CODE BOOK

Книга кодов

for / для MIKROLATEST®

ANAEROtest 23

Erba Lachema s.r.o.

Erba® Mannheim

Introduction

The codebook for ANAEROtest 23 is designed to identify bacteria using the MIKRO-LA-TEST ANAEROtest 23 test kit. The identification, ie interpretation of the result, can be done in three ways:

1. **manually** - by comparing the result with the data in the differentiation table
2. **using an identification program** (eg ErbaExpert)
3. **using this codebook**

All of the mentioned identification methods are based on the data of the identification matrix (frequency matrix, etc.), ie table containing taxa, tests and percentages of positive reactions of particular tests for each taxon. The taxon may be:

- genus - *Morganella sp*.
- species - *Morganella morganii*
- subspecies - *Morganella morganii ssp. morganii*

The codebook contains the identification results using the ANAEROtest 23 set ordered according to the value of the so-called profile code. Sorting results by the value of the profile code allows for easy and fast search. The identification quality evaluation for each profile code is based on the calculation of the probability of the result from the identification matrix data; the evaluation is detailed below.

Creating a profile code

For easier identification, it is advisable to convert the obtained results of each test (+/-) into a numerical profile (octal code). Tests are first divided into groups:

IND		GLU		MLT		FRU		GAL		LAC		MLZ		URE	
NIT	1	SUC	2	SAL	3	TRE	4	MAN	5	RHA	6	NAG	7	BGL	8
ESL		MNS		RAF		CEL		XYL		ARA		SOR			

Test groups are created in accordance with the test order in a vertical triples in the triplestrip.

The last group is incomplete and contains the remaining strip tests.

In each created group, there are assigned values 1, 2, and 4 to the particilar tests according to the following scheme:

IND	1	GLU	1	MLT	1	FRU	1	GAL	1	LAC	1	MLZ	1	URE	1
NIT	2	SUC	2	SAL	2	TRE	2	MAN	2	RHA	2	NAG	2	BGL	2
ESL	4	MNS	4	RAF	4	CEL	4	XYL	4	ARA	4	SOR	4		

When creating a profile code for a specific result, proceed as follows:

- positive tests are assigned 1, 2 or 4 values
- negative tests are always assigned a zero value
- the values assigned to the tests are added in each test group
- the resulting group of numbers represents the **profile code**

IND +	1	GLU +	1	MLT -	1	FRU +	1	GAL -	0	LAC -	0	MLZ -	0	URE +	1
NIT -	0	SUC +	2	SAL -	0	TRE +	2	MAN -	0	RHA +	2	NAG -	0	BGL +	0
ESL -	0	MNS +	4	RAF +	4	CEL -	0	XYL -	0	ARA -	0	SOR -	0		
1		**7**		**4**		**3**		**0**		**2**		**0**		**1**	

In this example, a profile **17430201** was created.

The results are organized in the codebook according to the value of the profile code in ascending order.

Using the codebook

The following information is provided for each profile code in the codebook:

- Profile code.
- Identification score and T-index values. They are listed for each taxon:

  Identification score – the taxon with the highest value is ranked first; the identification score indicates the probability of occurrence of a given result for that taxon, related to the probability of occurrence of the result for all other taxa.

  Identification score ≥ 99 … the strain is excellently distinguished
  Identification score ≥ 95 … the strain is very good distinguished
  Identification score ≥ 90 … the strain is distinguished
  Identification score < 90 … the strain is not sufficiently distinguished

  T-index – an indication of the extent to which the result corresponds to the most typical result for that taxon; an entirely typical result corresponds to a T-index value equal to 1. The value may lie in the range from 0 to 1 and is inversely proportional to the number of atypical tests.

  T-index ≥ 0,75 … typical strain
  T-index ≥ 0,50 … less typical strain
  T-index ≥ 0,25 … atypical strain
  T-index < 0,25 … completely atypical strain

- **List of atypical characters (tests)** - the list of all atypical characters (listed below) is valid only for the first taxon; the percentage of positive reactions is given for every atypical test.
- **List of additional tests** – if taxa are insufficiently differentiated, additional tests allow for more accurate identification. Additional tests show percentages of positive reactions.

If the profile code in the diagnostic list is not listed, it is a very atypical result, or it is a taxon that is not included in the identification matrix.

List of taxa used to create a codebook

Acidaminococcus fermentans
Actinomyces israelii
Actinomyces meyeri
Actinomyces naeslundii
Actinomyces odontolyticus
Actinomyces viscosus
Anaerococcus prevotii
Atopobium minutum
Atopobium parvulum
Bacteroides eggerthii
Bacteroides fragilis
Bacteroides ovatus
Bacteroides thetaiotaomicron
Bacteroides uniformis
Bacteroides vulgatus
Bifidobacterium adolescentis
Bifidobacterium breve
Bifidobacterium dentium
Bifidobacterium longum subsp. infantis
Bifidobacterium longum subsp. longum
Blautia hansenii
Blautia producta
Campylobacter gracilis
Campylobacter ureolyticus
Capnocytophaga ochracea
Clostridium barati
Clostridium bifermentans
Clostridium botulinum biovar A
Clostridium botulinum biovar B
Clostridium butyricum
Clostridium cadaveris
Clostridium chauvoei
Clostridium clostridioforme
Clostridium difficile
Clostridium glycolicum
Clostridium histolyticum
Clostridium innocuum
Clostridium limosum
Clostridium novyi biovar A
Clostridium paraputrificum
Clostridium perfringens
Clostridium ramosum
Clostridium septicum
Clostridium sordellii
Clostridium sporogenes
Clostridium tertium
Clostridium tetani
Collinsella aerofaciens
Eggerthella lenta
Eubacterium contortum
Eubacterium limosum
Eubacterium saburreum
Eubacterium tortuosum

Finegoldia magna
Fusobacterium mortiferum
Fusobacterium necrophorum subsp. necrophorum
Fusobacterium nucleatum subsp. nucleatum
Fusobacterium varium
Gemella morbillorum
Lactobacillus acidophilus
Lactobacillus catenaformis
Lactobacillus fermentum
Lactobacillus jensenii
Leptotrichia buccalis
Megasphaera elsdenii
Mitsuokella multacida
Odoribacter splanchnicus
Parabacteroides distasonis
Parvimonas micra
Peptococcus niger
Peptoniphilus asaccharolyticus
Peptostreptococcus anaerobius
Porphyromonas asaccharolytica
Prevotella bivia
Prevotella disiens
Prevotella intermedia
Prevotella melaninogenica
Prevotella oralis
Propionibacterium acnes
Propionibacterium granulosum
Propionibacterium propionicum
Pseudoflavonifractor capillosus
Pseudoramibacter alactolyticus
Veillonella parvula

List of tests used to create a codebook

Basic set of tests

IND indole production
NIT nitrate reduction
ESL aesculine hydrolysis
GLU acid from glucose
SUC acid from saccharose
MNS acid from mannose
MLT acid from maltose
SAL acid from salicine
RAF acid from raffinose
FRU acid from fructose
TRE acid from trehalose
CEL acid from cellobiose
GAL acid from galactose
MAN acid from mannitol
XYL acid from D-xylose
LAC acid from lactose
RHA acid from rhamnose
ARA acid from L-arabinose
MLZ acid from melezitose
NAG N-acetyl-glucosaminidase
SOR acid from sorbitol
URE urea hydrolysis
BGL ß-glucosidase

Additional (distinguishing) tests

CAT catalase production
COC coccoid cells
GRA Gram positive cells
SPO spores production

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
00000000	**Low discrimination**						**00000000**
					COC	SPO	
Clostridium histolyticum		13.81	1.00		1	90	
Clostridium limosum		13.81	1.00		1	99	
Finegoldia magna		13.81	1.00		99	1	
00000001	**Low discrimination**						**00000001**
					COC	GRA	
Campylobacter ureolyticus		70.71	1.00		1	1	
Anaerococcus prevotii		14.31	0.92		99	99	
Peptococcus niger		7.70	0.83		99	99	
00000002	**Low discrimination**						**00000002**
					GRA		
Peptococcus niger		31.06	0.87		99		
Acidaminococcus fermentans		25.29	0.84		1		
Parvimonas micra		22.99	0.84		99		
00000003	**Low discrimination**						**00000003**
					COC	GRA	
Anaerococcus prevotii		60.01	0.82		99	99	
Peptococcus niger		22.80	0.70		99	99	
Campylobacter ureolyticus		11.98	0.66		1	1	
00000010	**Low discrimination**						**00000010**
					COC	SPO	
Clostridium histolyticum		13.74	0.66	MLZ 1	1	90	
Clostridium limosum		13.74	0.66		1	99	
Finegoldia magna		13.74	0.66		99	1	
00000011	**Low discrimination**						**00000011**
					COC	GRA	
Campylobacter ureolyticus		68.04	0.66	MLZ 1	1	1	
Anaerococcus prevotii		13.77	0.58		99	99	
Peptococcus niger		7.41	0.49		99	99	
00000012	**Low discrimination**						**00000012**
					GRA		
Peptococcus niger		30.58	0.53	MLZ 1	99		
Acidaminococcus fermentans		24.90	0.50		1		
Parvimonas micra		22.64	0.50		99		
00000013	**Low discrimination**						**00000013**
					COC	GRA	
Anaerococcus prevotii		53.81	0.47	MLZ 1	99	99	
Peptococcus niger		20.44	0.36		99	99	
Campylobacter ureolyticus		10.74	0.32		1	1	
00000020	**Low discrimination**						**00000020**
					COC	SPO	
Peptococcus niger		48.11	0.84	NAG 10	99	1	
Atopobium minutum		7.50	0.88		1	1	
Clostridium histolyticum		6.22	0.66		1	90	
00000021	**Low discrimination**						**00000021**
					COC	GRA	
Peptococcus niger		47.07	0.67	NAG 10	99	99	
Campylobacter ureolyticus		39.29	0.66	URE 9	1	1	
Anaerococcus prevotii		7.95	0.58		99	99	
00000022	**Good identification**						**00000022**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoflavonifractor capillosus		94.43	1.00		1	1		
Peptococcus niger		3.12	0.71		99	99		
Atopobium minutum		1.84	0.85		1	99		
00000023	**Low discrimination**							**00000023**
					COC	GRA		
Pseudoflavonifractor capillosus		69.17	0.66	URE 1	1	1		
Peptococcus niger		22.37	0.54		99	99		
Anaerococcus prevotii		5.35	0.47		99	99		
00000030	**Low discrimination**							**00000030**
					COC	SPO		
Peptococcus niger		45.64	0.50	MLZ 1	99	1		
Atopobium minutum		7.11	0.54	NAG 10	1	1		
Clostridium histolyticum		5.90	0.32		1	90		
00000031	**Low discrimination**							**00000031**
					COC	GRA	SPO	
Clostridium novyi biovar A		35.38	0.56	GLU 99	1	99	99	
Peptococcus niger		30.61	0.32		99	99	1	
Campylobacter ureolyticus		25.55	0.32		1	1	1	
00000032	**Good identification**							**00000032**
					COC	GRA		
Pseudoflavonifractor capillosus		94.11	0.66	MLZ 1	1	1		
Peptococcus niger		3.11	0.37		99	99		
Atopobium minutum		1.83	0.51		1	99		
00000040	**Low discrimination**							**00000040**
					COC	SPO		
Clostridium histolyticum		13.21	0.66	SOR 1	1	90		
Clostridium limosum		13.21	0.66		1	99		
Finegoldia magna		13.21	0.66		99	1		
00000041	**Low discrimination**							**00000041**
					COC	GRA		
Campylobacter ureolyticus		68.69	0.66	SOR 1	1	1		
Anaerococcus prevotii		13.90	0.58		99	99		
Peptococcus niger		7.48	0.49		99	99		
00000042	**Low discrimination**							**00000042**
					GRA			
Peptococcus niger		30.49	0.53	SOR 1	99			
Acidaminococcus fermentans		24.83	0.50		1			
Parvimonas micra		22.57	0.50		99			
00000043	**Low discrimination**							**00000043**
					COC	GRA		
Anaerococcus prevotii		55.73	0.47	SOR 1	99	99		
Peptococcus niger		21.17	0.36		99	99		
Campylobacter ureolyticus		11.13	0.32		1	1		
00000050	**Low discrimination**							**00000050**
Clostridium novyi biovar A		23.82	0.63	GLU 99				
Clostridium histolyticum		10.10	0.32					
Clostridium limosum		10.10	0.32					
00000051	**Low discrimination**							**00000051**
					COC	GRA	SPO	
Clostridium novyi biovar A		72.28	0.63	GLU 99	1	99	99	
Campylobacter ureolyticus		19.57	0.32		1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Anaerococcus prevotii		3.96	0.24		99	99	1	
00000052	**Low discrimination**							**00000052**
					COC	GRA	SPO	
Clostridium novyi biovar A		51.06	0.50	GLU 99	1	99	99	
Peptococcus niger		15.23	0.19		99	99	1	
Acidaminococcus fermentans		12.40	0.15		99	1	1	
00000053	**Low discrimination**							**00000053**
					COC	SPO		
Clostridium novyi biovar A		89.08	0.50	GLU 99	1	99		
Anaerococcus prevotii		6.92	0.13		99	1		
Peptococcus niger		2.63	0.02		99	1		
00000060	**Low discrimination**							**00000060**
					COC	SPO		
Peptococcus niger		45.56	0.50	NAG 10	99	1		
Atopobium minutum		7.10	0.54	SOR 1	1	1		
Clostridium histolyticum		5.89	0.32		1	90		
00000062	**Good identification**							**00000062**
					COC	GRA		
Pseudoflavonifractor capillosus		94.21	0.66	SOR 1	1	1		
Peptococcus niger		3.11	0.37		99	99		
Atopobium minutum		1.83	0.51		1	99		
00000070	**Low discrimination**							**00000070**
					COC	SPO		
Clostridium novyi biovar A		78.28	0.53	GLU 99	1	99		
Peptococcus niger		10.37	0.15		99	1		
Atopobium minutum		1.62	0.20		1	1		
00000071	**Good identification**							**00000071**
					COC	GRA	SPO	
Clostridium novyi biovar A		97.65	0.53	GLU 99	1	99	99	
Peptococcus niger		1.28	-.02		99	99	1	
Campylobacter ureolyticus		1.07	-.03		1	1	1	
00000073	**Good identification**							**00000073**
					GRA	SPO		
Clostridium novyi biovar A		96.10	0.40	GLU 99	99	99		
Pseudoflavonifractor capillosus		3.90	-.03		1	1		
00000100	**Low discrimination**							**00000100**
					COC	SPO		
Clostridium histolyticum		13.81	0.66	LAC 1	1	90		
Clostridium limosum		13.81	0.66		1	99		
Finegoldia magna		13.81	0.66		99	1		
00000101	**Low discrimination**							**00000101**
					COC	GRA		
Campylobacter ureolyticus		70.71	0.66	LAC 1	1	1		
Anaerococcus prevotii		14.31	0.58		99	99		
Peptococcus niger		7.70	0.49		99	99		
00000102	**Low discrimination**							**00000102**
					GRA			
Peptococcus niger		31.05	0.53	LAC 1	99			
Acidaminococcus fermentans		25.29	0.50		1			
Parvimonas micra		22.99	0.50		99			
00000103	**Low discrimination**							**00000103**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Anaerococcus prevotii		60.05	0.47	LAC 1	99	99	
Peptococcus niger		22.81	0.36		99	99	
Campylobacter ureolyticus		11.99	0.32		1	1	
00000120	**Low discrimination**						**00000120**
					COC	SPO	
Peptococcus niger		48.06	0.50	LAC 1	99	1	
Atopobium minutum		7.49	0.54	NAG 10	1	1	
Clostridium histolyticum		6.22	0.32		1	90	
00000122	**Good identification**						**00000122**
					COC	GRA	
Pseudoflavonifractor capillosus		94.44	0.66	LAC 1	1	1	
Peptococcus niger		3.12	0.37		99	99	
Atopobium minutum		1.84	0.51		1	99	
00000200	**Low discrimination**						**00000200**
					COC	SPO	
Clostridium histolyticum		13.81	0.66	RHA 1	1	90	
Clostridium limosum		13.81	0.66		1	99	
Finegoldia magna		13.81	0.66		99	1	
00000201	**Low discrimination**						**00000201**
					COC	GRA	
Campylobacter ureolyticus		70.71	0.66	RHA 1	1	1	
Anaerococcus prevotii		14.31	0.58		99	99	
Peptococcus niger		7.70	0.49		99	99	
00000202	**Low discrimination**						**00000202**
					GRA		
Peptococcus niger		31.06	0.53	RHA 1	99		
Acidaminococcus fermentans		25.29	0.50		1		
Parvimonas micra		22.99	0.50		99		
00000203	**Low discrimination**						**00000203**
					COC	GRA	
Anaerococcus prevotii		60.05	0.47	RHA 1	99	99	
Peptococcus niger		22.81	0.36		99	99	
Campylobacter ureolyticus		11.99	0.32		1	1	
00000220	**Low discrimination**						**00000220**
					COC	SPO	
Peptococcus niger		48.12	0.50	RHA 1	99	1	
Atopobium minutum		7.50	0.54	NAG 10	1	1	
Clostridium histolyticum		6.22	0.32		1	90	
00000222	**Good identification**						**00000222**
					COC	GRA	
Pseudoflavonifractor capillosus		94.44	0.66	RHA 1	1	1	
Peptococcus niger		3.12	0.37		99	99	
Atopobium minutum		1.84	0.51		1	99	
00000400	**Low discrimination**						**00000400**
					COC	SPO	
Clostridium histolyticum		13.81	0.66	ARA 1	1	90	
Clostridium limosum		13.81	0.66		1	99	
Finegoldia magna		13.81	0.66		99	1	
00000401	**Low discrimination**						**00000401**
					COC	GRA	
Campylobacter ureolyticus		70.71	0.66	ARA 1	1	1	
Anaerococcus prevotii		14.31	0.58		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptococcus niger		7.70	0.49		99	99	
00000402	**Low discrimination**						**00000402**
					GRA		
Peptococcus niger		31.06	0.53	ARA 1	99		
Acidaminococcus fermentans		25.29	0.50		1		
Parvimonas micra		22.99	0.50		99		
00000403	**Low discrimination**						**00000403**
					COC	GRA	
Anaerococcus prevotii		60.05	0.47	ARA 1	99	99	
Peptococcus niger		22.81	0.36		99	99	
Campylobacter ureolyticus		11.99	0.32		1	1	
00000420	**Low discrimination**						**00000420**
					COC	SPO	
Peptococcus niger		48.12	0.50	ARA 1	99	1	
Atopobium minutum		7.50	0.54	NAG 10	1	1	
Clostridium histolyticum		6.22	0.32		1	90	
00000422	**Good identification**						**00000422**
					COC	GRA	
Pseudoflavonifractor capillosus		94.44	0.66	ARA 1	1	1	
Peptococcus niger		3.12	0.37		99	99	
Atopobium minutum		1.84	0.51		1	99	
00001000	**Low discrimination**						**00001000**
					SPO		
Atopobium minutum		13.60	0.88		1		
Clostridium histolyticum		11.29	0.66		90		
Clostridium limosum		11.29	0.66		99		
00001001	**Low discrimination**						**00001001**
					COC	GRA	
Campylobacter ureolyticus		69.16	0.66	GAL 1	1	1	
Anaerococcus prevotii		14.00	0.58		99	99	
Peptococcus niger		7.53	0.49		99	99	
00001002	**Low discrimination**						**00001002**
					COC	GRA	
Atopobium minutum		65.53	0.85		1	99	
Peptococcus niger		10.12	0.53		99	99	
Acidaminococcus fermentans		8.24	0.50		99	1	
00001003	**Low discrimination**						**00001003**
					COC		
Anaerococcus prevotii		51.46	0.47	GAL 1	99		
Peptococcus niger		19.55	0.36		99		
Atopobium minutum		12.93	0.51		1		
00001010	**Low discrimination**						**00001010**
					SPO		
Atopobium minutum		13.03	0.54	MLZ 1	1		
Clostridium histolyticum		10.81	0.32		90		
Clostridium limosum		10.81	0.32		99		
00001012	**Low discrimination**						**00001012**
					COC	GRA	
Atopobium minutum		62.24	0.51	MLZ 1	1	99	
Peptococcus niger		9.61	0.19		99	99	
Acidaminococcus fermentans		7.83	0.15		99	1	
00001013	**Low discrimination**						**00001013**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Clostridium novyi biovar A		53.98	0.37	GLU 99	1	99		
Anaerococcus prevotii		25.14	0.13	GAL 10	99	1		
Peptococcus niger		9.55	0.02		99	1		
00001020	**Low discrimination**							**00001020**
					COC	GRA		
Atopobium minutum		67.00	0.80		1	99		
Peptococcus niger		13.03	0.50		99	99		
Pseudoflavonifractor capillosus		7.74	0.50		1	1		
00001022	**Good identification**							**00001022**
					COC	GRA		
Pseudoflavonifractor capillosus		94.17	0.84	GAL 10	1	1		
Atopobium minutum		5.49	0.77		1	99		
Peptococcus niger		0.28	0.37		99	99		
00001023	**Acceptable identification**							**00001023**
					COC	GRA		
Pseudoflavonifractor capillosus		90.92	0.50	GAL 10	1	1		
Atopobium minutum		5.30	0.43	URE 1	1	99		
Peptococcus niger		2.67	0.19		99	99		
00001030	**Low discrimination**							**00001030**
					COC	SPO		
Atopobium minutum		57.80	0.46	MLZ 1	1	1		
Clostridium novyi biovar A		14.14	0.39		1	99		
Peptococcus niger		11.24	0.15		99	1		
00001031	**Low discrimination**							**00001031**
					COC	GRA	SPO	
Clostridium novyi biovar A		84.35	0.39	GLU 99	1	99	99	
Peptococcus niger		6.63	-.02	GAL 10	99	99	1	
Campylobacter ureolyticus		5.54	-.03		1	1	1	
00001032	**Acceptable identification**							**00001032**
					GRA	SPO		
Pseudoflavonifractor capillosus		93.89	0.50	GAL 10	1	1		
Atopobium minutum		5.47	0.43	MLZ 1	99	1		
Clostridium novyi biovar A		0.35	0.26		99	99		
00001040	**Low discrimination**							**00001040**
					SPO			
Atopobium minutum		12.78	0.54	SOR 1	1			
Clostridium histolyticum		10.61	0.32		90			
Clostridium limosum		10.61	0.32		99			
00001042	**Low discrimination**							**00001042**
					COC	GRA		
Atopobium minutum		63.21	0.51	SOR 1	1	99		
Peptococcus niger		9.77	0.19		99	99		
Acidaminococcus fermentans		7.95	0.15		99	1		
00001050	**Low discrimination**							**00001050**
					SPO			
Clostridium novyi biovar A		74.21	0.47	GLU 99	99			
Atopobium minutum		3.45	0.20	GAL 10	1			
Clostridium histolyticum		2.86	-.03		90			
00001051	**Good identification**							**00001051**
					GRA	SPO		
Clostridium novyi biovar A		97.60	0.47	GLU 99	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Campylobacter ureolyticus		2.40	-.03	GAL 10	1	1		
00001060	**Low discrimination**							**00001060**
					COC	SPO		
Atopobium minutum		60.37	0.46	SOR 1	1	1		
Peptococcus niger		11.74	0.15		99	1		
Clostridium novyi biovar A		9.85	0.36		1	99		
00001061	**Low discrimination**							**00001061**
					COC	GRA	SPO	
Clostridium novyi biovar A		78.23	0.36	GLU 99	1	99	99	
Peptococcus niger		9.23	-.02	GAL 10	99	99	1	
Campylobacter ureolyticus		7.70	-.03		1	1	1	
00001062	**Acceptable identification**							**00001062**
					COC	GRA		
Pseudoflavonifractor capillosus		94.00	0.50	GAL 10	1	1		
Atopobium minutum		5.48	0.43	SOR 1	1	99		
Peptococcus niger		0.28	0.03		99	99		
00001070	**Acceptable identification**							**00001070**
					SPO			
Clostridium novyi biovar A		94.17	0.36	GLU 99	99			
Atopobium minutum		5.83	0.11	GAL 10	1			
00001071	**Good identification**							**00001071**
Clostridium novyi biovar A		100.0	0.36	GLU 99				
00001100	**Low discrimination**							**00001100**
					SPO			
Atopobium minutum		13.38	0.54	LAC 1	1			
Clostridium histolyticum		11.11	0.32		90			
Clostridium limosum		11.11	0.32		99			
00001102	**Low discrimination**							**00001102**
					COC	GRA		
Atopobium minutum		65.54	0.51	LAC 1	1	99		
Peptococcus niger		10.12	0.19		99	99		
Acidaminococcus fermentans		8.24	0.15		99	1		
00001120	**Low discrimination**							**00001120**
					COC	GRA		
Atopobium minutum		65.24	0.46	LAC 1	1	99		
Peptococcus niger		12.69	0.15		99	99		
Pseudoflavonifractor capillosus		7.54	0.15		1	1		
00001122	**Acceptable identification**							**00001122**
					COC	GRA		
Pseudoflavonifractor capillosus		94.22	0.50	GAL 10	1	1		
Atopobium minutum		5.49	0.43	LAC 1	1	99		
Peptococcus niger		0.28	0.03		99	99		
00001200	**Low discrimination**							**00001200**
					SPO			
Atopobium minutum		13.60	0.54	RHA 1	1			
Clostridium histolyticum		11.29	0.32		90			
Clostridium limosum		11.29	0.32		99			
00001202	**Low discrimination**							**00001202**
					COC	GRA		
Atopobium minutum		65.77	0.51	RHA 1	1	99		
Peptococcus niger		10.16	0.19		99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Acidaminococcus fermentans		8.27	0.15		99	1	
00001220	**Low discrimination**						**00001220**
					COC	GRA	
Atopobium minutum		67.31	0.46	RHA 1	1	99	
Peptococcus niger		13.09	0.15		99	99	
Pseudoflavonifractor capillosus		7.78	0.15		1	1	
00001222	**Acceptable identification**						**00001222**
					COC	GRA	
Pseudoflavonifractor capillosus		94.22	0.50	GAL 10	1	1	
Atopobium minutum		5.49	0.43	RHA 1	1	99	
Peptococcus niger		0.28	0.03		99	99	
00001400	**Low discrimination**						**00001400**
					SPO		
Atopobium minutum		13.59	0.54	ARA 1	1		
Clostridium histolyticum		11.28	0.32		90		
Clostridium limosum		11.28	0.32		99		
00001402	**Low discrimination**						**00001402**
					COC	GRA	
Atopobium minutum		65.77	0.51	ARA 1	1	99	
Peptococcus niger		10.16	0.19		99	99	
Acidaminococcus fermentans		8.27	0.15		99	1	
00001420	**Low discrimination**						**00001420**
					COC	GRA	
Atopobium minutum		67.31	0.46	ARA 1	1	99	
Peptococcus niger		13.09	0.15		99	99	
Pseudoflavonifractor capillosus		7.78	0.15		1	1	
00001422	**Acceptable identification**						**00001422**
					COC	GRA	
Pseudoflavonifractor capillosus		94.22	0.50	GAL 10	1	1	
Atopobium minutum		5.49	0.43	ARA 1	1	99	
Peptococcus niger		0.28	0.03		99	99	
00002000	**Low discrimination**						**00002000**
					COC	SPO	
Clostridium histolyticum		13.80	0.66	MAN 1	1	90	
Clostridium limosum		13.80	0.66		1	99	
Finegoldia magna		13.80	0.66		99	1	
00002001	**Low discrimination**						**00002001**
					COC	GRA	
Campylobacter ureolyticus		70.70	0.66	MAN 1	1	1	
Anaerococcus prevotii		14.31	0.58		99	99	
Peptococcus niger		7.70	0.49		99	99	
00002002	**Low discrimination**						**00002002**
					GRA		
Peptococcus niger		31.02	0.53	MAN 1	99		
Acidaminococcus fermentans		25.26	0.50		1		
Parvimonas micra		22.96	0.50		99		
00002003	**Low discrimination**						**00002003**
					COC	GRA	
Anaerococcus prevotii		60.05	0.47	MAN 1	99	99	
Peptococcus niger		22.81	0.36		99	99	
Campylobacter ureolyticus		11.99	0.32		1	1	
00002020	**Low discrimination**						**00002020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Peptococcus niger		48.00	0.50	MAN 1	99	1	
Atopobium minutum		7.48	0.54	NAG 10	1	1	
Clostridium histolyticum		6.21	0.32		1	90	
00002022	**Good identification**						**00002022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.43	0.66	MAN 1	1	1	
Peptococcus niger		3.12	0.37		99	99	
Atopobium minutum		1.84	0.51		1	99	
00003000	**Low discrimination**						**00003000**
					SPO		
Atopobium minutum		13.59	0.54	MAN 1	1		
Clostridium histolyticum		11.28	0.32		90		
Clostridium limosum		11.28	0.32		99		
00003002	**Low discrimination**						**00003002**
					COC	GRA	
Atopobium minutum		65.77	0.51	MAN 1	1	99	
Peptococcus niger		10.16	0.19		99	99	
Acidaminococcus fermentans		8.27	0.15		99	1	
00003020	**Low discrimination**						**00003020**
					COC	GRA	
Atopobium minutum		67.31	0.46	MAN 1	1	99	
Peptococcus niger		13.09	0.15		99	99	
Pseudoflavonifractor capillosus		7.78	0.15		1	1	
00003022	**Acceptable identification**						**00003022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.22	0.50	GAL 10	1	1	
Atopobium minutum		5.49	0.43	MAN 1	1	99	
Peptococcus niger		0.28	0.03		99	99	
00004000	**Low discrimination**						**00004000**
					COC	SPO	
Clostridium histolyticum		13.80	0.66	XYL 1	1	90	
Clostridium limosum		13.80	0.66		1	99	
Finegoldia magna		13.80	0.66		99	1	
00004001	**Low discrimination**						**00004001**
					COC	GRA	
Campylobacter ureolyticus		70.71	0.66	XYL 1	1	1	
Anaerococcus prevotii		14.31	0.58		99	99	
Peptococcus niger		7.70	0.49		99	99	
00004002	**Low discrimination**						**00004002**
					GRA		
Peptococcus niger		31.06	0.53	XYL 1	99		
Acidaminococcus fermentans		25.29	0.50		1		
Parvimonas micra		22.99	0.50		99		
00004003	**Low discrimination**						**00004003**
					COC	GRA	
Anaerococcus prevotii		60.05	0.47	XYL 1	99	99	
Peptococcus niger		22.81	0.36		99	99	
Campylobacter ureolyticus		11.99	0.32		1	1	
00004020	**Low discrimination**						**00004020**
					COC	SPO	
Peptococcus niger		48.10	0.50	XYL 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		7.49	0.54	NAG 10	1	1	
Clostridium histolyticum		6.22	0.32		1	90	
00004022	**Good identification**						**00004022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.44	0.66	XYL 1	1	1	
Peptococcus niger		3.12	0.37		99	99	
Atopobium minutum		1.84	0.51		1	99	
00004040	**Low discrimination**						**00004040**
Clostridium glycolicum		21.32	0.49	GLU 99			
Clostridium histolyticum		10.40	0.32	FRU 90			
Clostridium limosum		10.40	0.32				
00005000	**Low discrimination**						**00005000**
					SPO		
Atopobium minutum		13.48	0.54	XYL 1	1		
Clostridium histolyticum		11.19	0.32		90		
Clostridium limosum		11.19	0.32		99		
00005002	**Low discrimination**						**00005002**
					COC	GRA	
Atopobium minutum		65.66	0.51	XYL 1	1	99	
Peptococcus niger		10.14	0.19		99	99	
Acidaminococcus fermentans		8.26	0.15		99	1	
00005020	**Low discrimination**						**00005020**
					COC	GRA	
Atopobium minutum		66.29	0.46	XYL 1	1	99	
Peptococcus niger		12.90	0.15		99	99	
Pseudoflavonifractor capillosus		7.66	0.15		1	1	
00005022	**Acceptable identification**						**00005022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.22	0.50	GAL 10	1	1	
Atopobium minutum		5.49	0.43	XYL 1	1	99	
Peptococcus niger		0.28	0.03		99	99	
00005100	**Low discrimination**						**00005100**
					SPO		
Actinomyces meyeri		75.44	0.39	GLU 93	1		
Atopobium minutum		3.40	0.20	SUC 93	1		
Clostridium histolyticum		2.82	-.03	FRU 95	90		
00010000	**Low discrimination**						**00010000**
					GRA	SPO	
Atopobium minutum		25.90	0.95		99	1	
Fusobacterium varium		13.36	0.77		1	1	
Clostridium histolyticum		7.76	0.66		99	90	
00010001	**Low discrimination**						**00010001**
					COC	GRA	
Campylobacter ureolyticus		66.13	0.66	FRU 1	1	1	
Anaerococcus prevotii		13.38	0.58		99	99	
Peptococcus niger		7.20	0.49		99	99	
00010002	**Low discrimination**						**00010002**
					COC	GRA	
Atopobium minutum		81.37	0.92		1	99	
Peptococcus niger		4.54	0.53		99	99	
Acidaminococcus fermentans		3.70	0.50		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
00010003	**Low discrimination**						**00010003**
					COC		
Anaerococcus prevotii		41.40	0.47	FRU 1	99		
Atopobium minutum		28.80	0.58		1		
Peptococcus niger		15.73	0.36		99		
00010010	**Low discrimination**						**00010010**
					GRA	SPO	
Atopobium minutum		24.28	0.61	MLZ 1	99	1	
Fusobacterium varium		12.52	0.42		1	1	
Clostridium histolyticum		7.28	0.32		99	90	
00010012	**Low discrimination**						**00010012**
					COC	GRA	
Atopobium minutum		79.34	0.58	MLZ 1	1	99	
Peptococcus niger		4.43	0.19		99	99	
Acidaminococcus fermentans		3.60	0.15		99	1	
00010013	**Low discrimination**						**00010013**
					COC	SPO	
Clostridium novyi biovar A		48.55	0.37	GLU 99	1	99	
Anaerococcus prevotii		22.61	0.13	FRU 10	99	1	
Atopobium minutum		15.73	0.24		1	1	
00010020	**Low discrimination**						**00010020**
					COC		
Atopobium minutum		78.00	0.87		1		
Peptococcus niger		5.48	0.50		99		
Pseudoramibacter alactolyticus		5.13	0.53		1		
00010021	**Low discrimination**						**00010021**
					COC	GRA	
Atopobium minutum		37.39	0.53	URE 1	1	99	
Peptococcus niger		25.72	0.32		99	99	
Campylobacter ureolyticus		21.46	0.32		1	1	
00010022	**Low discrimination**						**00010022**
					COC	GRA	
Pseudoflavonifractor capillosus		85.69	0.84	FRU 10	1	1	
Atopobium minutum		13.83	0.84		1	99	
Peptococcus niger		0.26	0.37		99	99	
00010023	**Low discrimination**						**00010023**
					COC	GRA	
Pseudoflavonifractor capillosus		83.00	0.50	FRU 10	1	1	
Atopobium minutum		13.40	0.50	URE 1	1	99	
Peptococcus niger		2.44	0.19		99	99	
00010030	**Low discrimination**						**00010030**
					COC	SPO	
Atopobium minutum		73.03	0.53	MLZ 1	1	1	
Clostridium novyi biovar A		6.45	0.39		1	99	
Peptococcus niger		5.13	0.15		99	1	
00010031	**Low discrimination**						**00010031**
					COC	SPO	
Clostridium novyi biovar A		79.45	0.39	GLU 99	1	99	
Atopobium minutum		9.08	0.19	FRU 10	1	1	
Peptococcus niger		6.25	-.02		99	1	
00010032	**Low discrimination**						**00010032**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		85.47	0.50	FRU 10	1	1	
Atopobium minutum		13.80	0.50	MLZ 1	99	1	
Clostridium novyi biovar A		0.32	0.26		99	99	
00010040	**Low discrimination**						**00010040**
					COC	GRA	
Pseudoramibacter alactolyticus		35.75	0.56	GLU 99	1	99	
Atopobium minutum		11.63	0.61		1	99	
Megasphaera elsdenii		10.04	0.57		99	1	
00010042	**Low discrimination**						**00010042**
					COC	GRA	
Atopobium minutum		52.80	0.58	SOR 1	1	99	
Pseudoramibacter alactolyticus		27.04	0.39		1	99	
Megasphaera elsdenii		7.59	0.41		99	1	
00010050	**Low discrimination**						**00010050**
					SPO		
Clostridium novyi biovar A		45.85	0.47	GLU 99	99		
Pseudoramibacter alactolyticus		18.12	0.21	FRU 10	1		
Atopobium minutum		5.90	0.27		1		
00010051	**Good identification**						**00010051**
					GRA	SPO	
Clostridium novyi biovar A		97.60	0.47	GLU 99	99	99	
Campylobacter ureolyticus		2.40	-.03	FRU 10	1	1	
00010060	**Low discrimination**						**00010060**
					COC		
Pseudoramibacter alactolyticus		54.87	0.43	GLU 99	1		
Atopobium minutum		33.73	0.53		1		
Peptococcus niger		2.37	0.15		99		
00010061	**Low discrimination**						**00010061**
					SPO		
Clostridium novyi biovar A		59.99	0.36	GLU 99	99		
Pseudoramibacter alactolyticus		16.74	0.09	FRU 10	1		
Atopobium minutum		10.29	0.19		1		
00010062	**Low discrimination**						**00010062**
					GRA		
Pseudoflavonifractor capillosus		82.55	0.50	FRU 10	1		
Atopobium minutum		13.33	0.50	SOR 1	99		
Pseudoramibacter alactolyticus		3.61	0.26		99		
00010070	**Low discrimination**						**00010070**
					SPO		
Clostridium novyi biovar A		68.94	0.36	GLU 99	99		
Pseudoramibacter alactolyticus		19.23	0.09	FRU 10	1		
Atopobium minutum		11.82	0.19		1		
00010071	**Good identification**						**00010071**
Clostridium novyi biovar A		100.0	0.36	GLU 99			
00010100	**Low discrimination**						**00010100**
					GRA		
Atopobium minutum		22.75	0.61	LAC 1	99		
Actinomyces meyeri		12.01	0.53		99		
Fusobacterium varium		11.73	0.42		1		
00010102	**Low discrimination**						**00010102**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		79.80	0.58	LAC 1	1	99	
Peptococcus niger		4.45	0.19		99	99	
Acidaminococcus fermentans		3.63	0.15		99	1	
00010120	**Low discrimination**						**00010120**
					COC		
Atopobium minutum		66.99	0.53	LAC 1	1		
Actinomyces meyeri		14.47	0.38		1		
Peptococcus niger		4.71	0.15		99		
00010122	**Low discrimination**						**00010122**
					COC	GRA	
Pseudoflavonifractor capillosus		85.63	0.50	FRU 10	1	1	
Atopobium minutum		13.83	0.50	LAC 1	1	99	
Peptococcus niger		0.26	0.03		99	99	
00010200	**Low discrimination**						**00010200**
					GRA	SPO	
Atopobium minutum		25.91	0.61	RHA 1	99	1	
Fusobacterium varium		13.36	0.42		1	1	
Clostridium histolyticum		7.77	0.32		99	90	
00010202	**Low discrimination**						**00010202**
					COC	GRA	
Atopobium minutum		81.50	0.58	RHA 1	1	99	
Peptococcus niger		4.55	0.19		99	99	
Acidaminococcus fermentans		3.70	0.15		99	1	
00010220	**Low discrimination**						**00010220**
					COC		
Atopobium minutum		78.33	0.53	RHA 1	1		
Peptococcus niger		5.50	0.15		99		
Pseudoramibacter alactolyticus		5.15	0.19		1		
00010222	**Low discrimination**						**00010222**
					COC	GRA	
Pseudoflavonifractor capillosus		85.75	0.50	FRU 10	1	1	
Atopobium minutum		13.84	0.50	RHA 1	1	99	
Peptococcus niger		0.26	0.03		99	99	
00010400	**Low discrimination**						**00010400**
					GRA	SPO	
Atopobium minutum		25.69	0.61	ARA 1	99	1	
Fusobacterium varium		13.25	0.42		1	1	
Clostridium histolyticum		7.70	0.32		99	90	
00010402	**Low discrimination**						**00010402**
					COC	GRA	
Atopobium minutum		81.39	0.58	ARA 1	1	99	
Peptococcus niger		4.54	0.19		99	99	
Acidaminococcus fermentans		3.70	0.15		99	1	
00010420	**Low discrimination**						**00010420**
					COC		
Atopobium minutum		77.51	0.53	ARA 1	1		
Peptococcus niger		5.45	0.15		99		
Pseudoramibacter alactolyticus		5.09	0.19		1		
00010422	**Low discrimination**						**00010422**
					COC	GRA	
Pseudoflavonifractor capillosus		85.75	0.50	FRU 10	1	1	
Atopobium minutum		13.84	0.50	ARA 1	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptococcus niger		0.26	0.03		99	99	
00010500	**Low discrimination**						**00010500**
					GRA		
Actinomyces meyeri		77.69	0.42	GLU 93	99		
Atopobium minutum		5.94	0.27	SUC 93	99		
Fusobacterium varium		3.07	0.08		1		
00011000	**Good identification**						**00011000**
					GRA	SPO	
Atopobium minutum		91.06	0.87		99	1	
Fusobacterium varium		1.42	0.42		1	1	
Clostridium histolyticum		0.83	0.32		99	90	
00011001	**Low discrimination**						**00011001**
					COC	GRA	
Atopobium minutum		52.34	0.53	URE 1	1	99	
Campylobacter ureolyticus		30.05	0.32		1	1	
Anaerococcus prevotii		6.08	0.24		99	99	
00011002	**Very good identification**						**00011002**
					COC	GRA	
Atopobium minutum		98.59	0.84		1	99	
Pseudoflavonifractor capillosus		0.69	0.33		1	1	
Peptococcus niger		0.17	0.19		99	99	
00011003	**Acceptable identification**						**00011003**
					COC		
Atopobium minutum		92.07	0.50	URE 1	1		
Anaerococcus prevotii		4.01	0.13		99		
Peptococcus niger		1.52	0.02		99		
00011010	**Low discrimination**						**00011010**
					GRA	SPO	
Atopobium minutum		87.94	0.53	MLZ 1	99	1	
Clostridium novyi biovar A		3.45	0.33		99	99	
Fusobacterium varium		1.37	0.08		1	1	
00011012	**Good identification**						**00011012**
					GRA	SPO	
Atopobium minutum		98.30	0.50	MLZ 1	99	1	
Clostridium novyi biovar A		1.02	0.20		99	99	
Pseudoflavonifractor capillosus		0.68	-.01		1	1	
00011020	**Very good identification**						**00011020**
					GRA		
Atopobium minutum		97.43	0.79		99		
Pseudoflavonifractor capillosus		1.36	0.33		1		
Actinomyces meyeri		0.24	0.31		99		
00011021	**Acceptable identification**						**00011021**
					COC	SPO	
Atopobium minutum		92.46	0.45	URE 1	1	1	
Clostridium novyi biovar A		2.72	0.23		1	99	
Peptococcus niger		1.93	-.02		99	1	
00011022	**Low discrimination**						**00011022**
					GRA		
Pseudoflavonifractor capillosus		67.33	0.67	FRU 10	1		
Atopobium minutum		32.61	0.76	GAL 10	99		
Lactobacillus fermentum		0.02	0.28		99		
00011030	**Good identification**						**00011030**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Atopobium minutum		95.85	0.45	MLZ 1	99	1	
Clostridium novyi biovar A		2.82	0.23		99	99	
Pseudoflavonifractor capillosus		1.33	-.01		1	1	
00011040	**Low discrimination**						**00011040**
					SPO		
Atopobium minutum		74.08	0.53	SOR 1	1		
Clostridium glycolicum		7.19	0.27		99		
Pseudoramibacter alactolyticus		6.90	0.21		1		
00011042	**Good identification**						**00011042**
					SPO		
Atopobium minutum		96.73	0.50	SOR 1	1		
Pseudoramibacter alactolyticus		1.50	0.05		1		
Clostridium novyi biovar A		0.67	0.17		99		
00011060	**Low discrimination**						**00011060**
					CAT		
Atopobium minutum		88.60	0.45	SOR 1	1		
Pseudoramibacter alactolyticus		4.37	0.09		1		
Propionibacterium acnes		4.07	0.15		95		
00011100	**Low discrimination**						**00011100**
Actinomyces meyeri		66.51	0.56	GLU 93			
Atopobium minutum		27.99	0.53	SUC 93			
Lactobacillus fermentum		1.85	0.40				
00011102	**Low discrimination**						**00011102**
Atopobium minutum		81.73	0.50	LAC 1			
Actinomyces meyeri		9.01	0.30				
Lactobacillus fermentum		8.11	0.40				
00011120	**Low discrimination**						**00011120**
Atopobium minutum		48.73	0.45	LAC 1			
Actinomyces meyeri		47.37	0.41				
Lactobacillus fermentum		3.22	0.32				
00011200	**Acceptable identification**						**00011200**
					GRA	SPO	
Atopobium minutum		91.45	0.53	RHA 1	99	1	
Fusobacterium varium		1.43	0.08		1	1	
Clostridium histolyticum		0.83	-.03		99	90	
00011202	**Good identification**						**00011202**
					GRA		
Atopobium minutum		99.31	0.50	RHA 1	99		
Pseudoflavonifractor capillosus		0.69	-.01		1		
00011220	**Good identification**						**00011220**
					GRA		
Atopobium minutum		98.63	0.45	RHA 1	99		
Pseudoflavonifractor capillosus		1.37	-.01		1		
00011400	**Low discrimination**						**00011400**
Atopobium minutum		78.34	0.53	ARA 1			
Actinomyces meyeri		11.64	0.35				
Lactobacillus fermentum		3.17	0.36				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
00011402	**Acceptable identification**						**00011402**
					GRA		
Atopobium minutum		93.06	0.50	ARA 1	99		
Lactobacillus fermentum		5.66	0.36		99		
Pseudoflavonifractor capillosus		0.65	-.01		1		
00011420	**Low discrimination**						**00011420**
Atopobium minutum		89.67	0.45	ARA 1			
Actinomyces meyeri		5.45	0.20				
Lactobacillus fermentum		3.63	0.28				
00011500	**Low discrimination**						**00011500**
Actinomyces meyeri		88.63	0.45	GLU 93			
Lactobacillus fermentum		9.86	0.40	SUC 93			
Atopobium minutum		1.51	0.19				
00011502	**Low discrimination**						**00011502**
Lactobacillus fermentum		72.48	0.40	GLU 99			
Actinomyces meyeri		20.14	0.20	SUC 87			
Atopobium minutum		7.38	0.16				
00012000	**Low discrimination**						**00012000**
					COC	GRA	
Pseudoramibacter alactolyticus		41.11	0.61	GLU 99	1	99	
Eubacterium limosum		26.14	0.60		1	99	
Megasphaera elsdenii		8.04	0.60		99	1	
00012002	**Low discrimination**						**00012002**
					COC	GRA	
Pseudoramibacter alactolyticus		42.58	0.45	GLU 99	1	99	
Atopobium minutum		38.60	0.58	BGL 10	1	99	
Megasphaera elsdenii		8.32	0.44		99	1	
00012010	**Low discrimination**						**00012010**
					SPO		
Clostridium difficile		88.90	0.53	GLU 99	99		
Pseudoramibacter alactolyticus		4.63	0.27		1		
Eubacterium limosum		2.95	0.26		1		
00012020	**Low discrimination**						**00012020**
Pseudoramibacter alactolyticus		71.09	0.48	GLU 99			
Atopobium minutum		20.29	0.53				
Eubacterium limosum		2.59	0.26				
00012022	**Low discrimination**						**00012022**
					GRA		
Pseudoflavonifractor capillosus		79.38	0.50	FRU 10	1		
Atopobium minutum		12.82	0.50	MAN 1	99		
Pseudoramibacter alactolyticus		7.48	0.32		99		
00012040	**Low discrimination**						**00012040**
					COC	GRA	
Pseudoramibacter alactolyticus		63.16	0.51	GLU 99	1	99	
Megasphaera elsdenii		32.92	0.57		99	1	
Eubacterium limosum		1.62	0.26		1	99	
00012060	**Good identification**						**00012060**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoramibacter alactolyticus		95.13	0.38	GLU 99	1	99	
Megasphaera elsdenii		2.84	0.23		99	1	
Atopobium minutum		1.10	0.19		1	99	
00013000	**Low discrimination**						**00013000**
Atopobium minutum		68.26	0.53	MAN 1			
Pseudoramibacter alactolyticus		13.69	0.27				
Eubacterium limosum		8.71	0.26				
00013002	**Good identification**						**00013002**
					GRA		
Atopobium minutum		95.52	0.50	MAN 1	99		
Pseudoramibacter alactolyticus		3.19	0.11		99		
Pseudoflavonifractor capillosus		0.66	-.01		1		
00013020	**Low discrimination**						**00013020**
					CAT		
Atopobium minutum		80.80	0.45	MAN 1	1		
Propionibacterium acnes		9.50	0.22		95		
Pseudoramibacter alactolyticus		8.58	0.14		1		
00014000	**Low discrimination**						**00014000**
					GRA	SPO	
Fusobacterium varium		35.34	0.53	XYL 4	1	1	
Atopobium minutum		16.61	0.61		99	1	
Clostridium histolyticum		4.98	0.32		99	90	
00014002	**Low discrimination**						**00014002**
					COC	GRA	
Atopobium minutum		78.88	0.58	XYL 1	1	99	
Peptococcus niger		4.40	0.19		99	99	
Acidaminococcus fermentans		3.58	0.15		99	1	
00014020	**Low discrimination**						**00014020**
					COC		
Atopobium minutum		69.48	0.53	XYL 1	1		
Actinomyces meyeri		7.29	0.33		1		
Peptococcus niger		4.88	0.15		99		
00014022	**Low discrimination**						**00014022**
					COC	GRA	
Pseudoflavonifractor capillosus		85.69	0.50	FRU 10	1	1	
Atopobium minutum		13.83	0.50	XYL 1	1	99	
Peptococcus niger		0.26	0.03		99	99	
00014040	**Very good identification**						**00014040**
					GRA	SPO	
Clostridium glycolicum		98.19	0.65	GLU 99	99	99	
Pseudoramibacter alactolyticus		0.55	0.21		99	1	
Fusobacterium varium		0.38	0.19		1	1	
00014100	**Good identification**						**00014100**
					GRA		
Actinomyces meyeri		94.66	0.58	GLU 93	99		
Fusobacterium varium		1.98	0.19	SUC 93	1		
Atopobium minutum		0.93	0.27		99		
00014120	**Good identification**						**00014120**
Actinomyces meyeri		97.65	0.43	GLU 93			
Atopobium minutum		2.35	0.19	SUC 93			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
00014400	**Low discrimination**					**00014400**
					GRA	
Actinomyces meyeri		52.86	0.37	GLU 93	99	
Fusobacterium varium		17.73	0.19	SUC 93	1	
Atopobium minutum		8.33	0.27		99	
00014500	**Good identification**					**00014500**
Actinomyces meyeri		100.0	0.47	GLU 93		
00015000	**Low discrimination**					**00015000**
					SPO	
Actinomyces meyeri		47.85	0.50	GLU 93	1	
Atopobium minutum		41.49	0.53	SUC 93	1	
Clostridium glycolicum		4.03	0.27		99	
00015002	**Acceptable identification**					**00015002**
Atopobium minutum		91.23	0.50	XYL 1		
Actinomyces meyeri		4.88	0.25			
Lactobacillus fermentum		3.26	0.32			
00015020	**Low discrimination**					**00015020**
Atopobium minutum		66.25	0.45	XYL 1		
Actinomyces meyeri		31.26	0.36			
Lactobacillus fermentum		1.58	0.24			
00015040	**Good identification**					**00015040**
					SPO	
Clostridium glycolicum		98.83	0.49	GLU 99	99	
Actinomyces meyeri		0.63	0.16	GAL 10	1	
Atopobium minutum		0.54	0.19		1	
00015100	**Very good identification**					**00015100**
Actinomyces meyeri		98.88	0.61	GLU 93		
Lactobacillus fermentum		0.83	0.36	SUC 93		
Atopobium minutum		0.22	0.19			
00015120	**Good identification**					**00015120**
Actinomyces meyeri		97.47	0.46	GLU 93		
Lactobacillus fermentum		2.01	0.28	SUC 93		
Atopobium minutum		0.52	0.11	NAG 12		
00015400	**Low discrimination**					**00015400**
Actinomyces meyeri		89.48	0.40	GLU 93		
Lactobacillus fermentum		7.39	0.32	SUC 93		
Atopobium minutum		3.14	0.19			
00015500	**Good identification**					**00015500**
Actinomyces meyeri		96.74	0.50	GLU 93		
Lactobacillus fermentum		3.26	0.36	SUC 93		
00015502	**Low discrimination**					**00015502**
Lactobacillus fermentum		52.12	0.36	GLU 99		
Actinomyces meyeri		47.88	0.25	SUC 87		
00015520	**Acceptable identification**					**00015520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		92.40	0.36	GLU 93			
Lactobacillus fermentum		7.61	0.28	SUC 93			
00020000	**Low discrimination**						**00020000**
					COC	SPO	
Clostridium histolyticum		13.13	0.66	TRE 1	1	90	
Clostridium limosum		13.13	0.66		1	99	
Finegoldia magna		13.13	0.66		99	1	
00020001	**Low discrimination**						**00020001**
					COC	GRA	
Campylobacter ureolyticus		70.41	0.66	TRE 1	1	1	
Anaerococcus prevotii		14.25	0.58		99	99	
Peptococcus niger		7.67	0.49		99	99	
00020002	**Low discrimination**						**00020002**
					GRA		
Peptococcus niger		27.90	0.53	TRE 1	99		
Acidaminococcus fermentans		22.72	0.50		1		
Parvimonas micra		20.65	0.50		99		
00020003	**Low discrimination**						**00020003**
					COC	GRA	
Anaerococcus prevotii		59.53	0.47	TRE 1	99	99	
Peptococcus niger		22.61	0.36		99	99	
Campylobacter ureolyticus		11.88	0.32		1	1	
00020020	**Low discrimination**						**00020020**
					COC	SPO	
Peptococcus niger		46.70	0.50	TRE 1	99	1	
Atopobium minutum		7.28	0.54	NAG 10	1	1	
Clostridium histolyticum		6.04	0.32		1	90	
00020022	**Good identification**						**00020022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.34	0.66	TRE 1	1	1	
Peptococcus niger		3.12	0.37		99	99	
Atopobium minutum		1.83	0.51		1	99	
00020040	**Low discrimination**						**00020040**
Clostridium botulinum biovar A		50.88	0.60	GLU 90			
Clostridium histolyticum		6.65	0.32				
Clostridium limosum		6.65	0.32				
00021000	**Low discrimination**						**00021000**
					SPO		
Atopobium minutum		13.04	0.54	TRE 1	1		
Clostridium histolyticum		10.83	0.32		90		
Clostridium limosum		10.83	0.32		99		
00021002	**Low discrimination**						**00021002**
					COC	GRA	
Atopobium minutum		63.35	0.51	TRE 1	1	99	
Peptococcus niger		9.79	0.19		99	99	
Acidaminococcus fermentans		7.97	0.15		99	1	
00021020	**Low discrimination**						**00021020**
					COC	GRA	
Atopobium minutum		67.02	0.46	TRE 1	1	99	
Peptococcus niger		13.04	0.15		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		7.74	0.15		1	1	
00021022	**Acceptable identification**						**00021022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.22	0.50	TRE 1	1	1	
Atopobium minutum		5.49	0.43	GAL 10	1	99	
Peptococcus niger		0.28	0.03		99	99	
00030000	**Low discrimination**						**00030000**
					GRA	SPO	
Atopobium minutum		19.98	0.61	TRE 1	99	1	
Clostridium sporogenes		13.81	0.49		99	99	
Fusobacterium varium		10.31	0.42		1	1	
00030002	**Low discrimination**						**00030002**
					COC	SPO	
Atopobium minutum		63.71	0.58	TRE 1	1	1	
Clostridium sporogenes		22.02	0.40		1	99	
Peptococcus niger		3.55	0.19		99	1	
00030020	**Low discrimination**						**00030020**
					COC	SPO	
Atopobium minutum		70.73	0.53	TRE 1	1	1	
Clostridium cadaveris		8.87	0.27		1	99	
Peptococcus niger		4.97	0.15		99	1	
00030022	**Low discrimination**						**00030022**
					COC	GRA	
Pseudoflavonifractor capillosus		85.62	0.50	FRU 10	1	1	
Atopobium minutum		13.82	0.50	TRE 1	1	99	
Peptococcus niger		0.26	0.03		99	99	
00030040	**Low discrimination**						**00030040**
					SPO		
Clostridium botulinum biovar A		62.68	0.39	GLU 90	99		
Pseudoramibacter alactolyticus		13.30	0.21	FRU 6	1		
Atopobium minutum		4.33	0.27		1		
00031000	**Low discrimination**						**00031000**
					GRA	SPO	
Atopobium minutum		87.93	0.53	TRE 1	99	1	
Clostridium sporogenes		1.84	0.14		99	99	
Fusobacterium varium		1.37	0.08		1	1	
00031002	**Good identification**						**00031002**
					GRA	SPO	
Atopobium minutum		98.29	0.50	TRE 1	99	1	
Clostridium sporogenes		1.03	0.06		99	99	
Pseudoflavonifractor capillosus		0.68	-.01		1	1	
00031020	**Good identification**						**00031020**
					GRA		
Atopobium minutum		98.63	0.45	TRE 1	99		
Pseudoflavonifractor capillosus		1.37	-.01		1		
00040000	**Low discrimination**						**00040000**
					COC	SPO	
Clostridium histolyticum		13.81	0.66	CEL 1	1	90	
Clostridium limosum		13.81	0.66		1	99	
Finegoldia magna		13.81	0.66		99	1	
00040001	**Low discrimination**						**00040001**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Campylobacter ureolyticus		70.71	0.66	CEL 1	1	1	
Anaerococcus prevotii		14.31	0.58		99	99	
Peptococcus niger		7.70	0.49		99	99	
00040002	**Low discrimination**						**00040002**
					GRA		
Peptococcus niger		31.06	0.53	CEL 1	99		
Acidaminococcus fermentans		25.29	0.50		1		
Parvimonas micra		22.99	0.50		99		
00040003	**Low discrimination**						**00040003**
					COC	GRA	
Anaerococcus prevotii		60.05	0.47	CEL 1	99	99	
Peptococcus niger		22.81	0.36		99	99	
Campylobacter ureolyticus		11.99	0.32		1	1	
00040020	**Low discrimination**						**00040020**
					COC	SPO	
Peptococcus niger		48.12	0.50	CEL 1	99	1	
Atopobium minutum		7.50	0.54	NAG 10	1	1	
Clostridium histolyticum		6.22	0.32		1	90	
00040022	**Good identification**						**00040022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.44	0.66	CEL 1	1	1	
Peptococcus niger		3.12	0.37		99	99	
Atopobium minutum		1.84	0.51		1	99	
00041000	**Low discrimination**						**00041000**
					SPO		
Atopobium minutum		13.60	0.54	CEL 1	1		
Clostridium histolyticum		11.29	0.32		90		
Clostridium limosum		11.29	0.32		99		
00041002	**Low discrimination**						**00041002**
					COC	GRA	
Atopobium minutum		65.77	0.51	CEL 1	1	99	
Peptococcus niger		10.16	0.19		99	99	
Acidaminococcus fermentans		8.27	0.15		99	1	
00041020	**Low discrimination**						**00041020**
					COC	GRA	
Atopobium minutum		67.31	0.46	CEL 1	1	99	
Peptococcus niger		13.09	0.15		99	99	
Pseudoflavonifractor capillosus		7.78	0.15		1	1	
00041022	**Acceptable identification**						**00041022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.22	0.50	CEL 1	1	1	
Atopobium minutum		5.49	0.43	GAL 10	1	99	
Peptococcus niger		0.28	0.03		99	99	
00050000	**Low discrimination**						**00050000**
					GRA	SPO	
Atopobium minutum		25.89	0.61	CEL 1	99	1	
Fusobacterium varium		13.35	0.42		1	1	
Clostridium histolyticum		7.76	0.32		99	90	
00050002	**Low discrimination**						**00050002**
					COC	GRA	
Atopobium minutum		81.50	0.58	CEL 1	1	99	
Peptococcus niger		4.55	0.19		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Acidaminococcus fermentans		3.70	0.15		99	1		
00050020	**Low discrimination**							**00050020**
					COC			
Atopobium minutum		78.33	0.53	CEL 1	1			
Peptococcus niger		5.50	0.15		99			
Pseudoramibacter alactolyticus		5.15	0.19		1			
00050022	**Low discrimination**							**00050022**
					COC	GRA		
Pseudoflavonifractor capillosus		85.75	0.50	FRU 10	1	1		
Atopobium minutum		13.84	0.50	CEL 1	1	99		
Peptococcus niger		0.26	0.03		99	99		
00051000	**Acceptable identification**							**00051000**
					GRA	SPO		
Atopobium minutum		90.94	0.53	CEL 1	99	1		
Fusobacterium varium		1.42	0.08		1	1		
Clostridium histolyticum		0.83	-.03		99	90		
00051002	**Good identification**							**00051002**
					GRA			
Atopobium minutum		99.31	0.50	CEL 1	99			
Pseudoflavonifractor capillosus		0.69	-.01		1			
00051020	**Good identification**							**00051020**
					GRA			
Atopobium minutum		98.63	0.45	CEL 1	99			
Pseudoflavonifractor capillosus		1.37	-.01		1			
00100000	**Low discrimination**							**00100000**
					COC	SPO		
Peptostreptococcus anaerobius		66.79	0.96		99	1		
Clostridium histolyticum		3.79	0.66		1	90		
Clostridium limosum		3.79	0.66		1	99		
00100001	**Low discrimination**							**00100001**
					COC	GRA		
Campylobacter ureolyticus		55.55	0.66	MLT 1	1	1		
Peptostreptococcus anaerobius		15.50	0.61		99	99		
Anaerococcus prevotii		11.24	0.58		99	99		
00100002	**Low discrimination**							**00100002**
					COC	SPO		
Clostridium sporogenes		29.34	0.66		1	99		
Peptostreptococcus anaerobius		21.32	0.61		99	1		
Peptococcus niger		14.85	0.53		99	1		
00100003	**Low discrimination**							**00100003**
					COC	GRA		
Anaerococcus prevotii		50.04	0.47	MLT 1	99	99		
Peptococcus niger		19.01	0.36		99	99		
Campylobacter ureolyticus		9.99	0.32		1	1		
00100010	**Low discrimination**							**00100010**
					COC	SPO		
Peptostreptococcus anaerobius		58.98	0.61	MLZ 1	99	1		
Clostridium novyi biovar A		11.83	0.66		1	99		
Clostridium histolyticum		3.35	0.32		1	90		
00100011	**Low discrimination**							**00100011**
					COC	GRA	SPO	
Clostridium novyi biovar A		76.10	0.66	GLU 99	1	99	99	

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Campylobacter ureolyticus		13.74	0.32		1	1	1	
Peptostreptococcus anaerobius		3.83	0.27		99	99	1	
00100012	**Low discrimination**							**00100012**
					COC	SPO		
Clostridium novyi biovar A		43.08	0.53	GLU 99	1	99		
Clostridium sporogenes		16.92	0.32		1	99		
Peptostreptococcus anaerobius		12.29	0.27		99	1		
00100013	**Acceptable identification**							**00100013**
					COC	SPO		
Clostridium novyi biovar A		91.86	0.53	GLU 99	1	99		
Anaerococcus prevotii		4.75	0.13		99	1		
Peptococcus niger		1.81	0.02		99	1		
00100020	**Low discrimination**							**00100020**
					COC			
Peptostreptococcus anaerobius		48.26	0.61	NAG 1	99			
Peptococcus niger		21.17	0.50		99			
Atopobium minutum		3.30	0.54		1			
00100021	**Low discrimination**							**00100021**
					COC	GRA	SPO	
Clostridium novyi biovar A		32.60	0.56	GLU 99	1	99	99	
Peptococcus niger		28.20	0.32		99	99	1	
Campylobacter ureolyticus		23.54	0.32		1	1	1	
00100022	**Good identification**							**00100022**
					COC	GRA		
Pseudoflavonifractor capillosus		93.19	0.66	MLT 1	1	1		
Peptococcus niger		3.08	0.37		99	99		
Atopobium minutum		1.81	0.51		1	99		
00100030	**Low discrimination**							**00100030**
					COC	SPO		
Clostridium novyi biovar A		71.30	0.56	GLU 99	1	99		
Peptostreptococcus anaerobius		14.36	0.27		99	1		
Peptococcus niger		6.30	0.15		99	1		
00100031	**Good identification**							**00100031**
					COC	GRA	SPO	
Clostridium novyi biovar A		98.23	0.56	GLU 99	1	99	99	
Peptococcus niger		0.86	-.02		99	99	1	
Campylobacter ureolyticus		0.72	-.03		1	1	1	
00100033	**Good identification**							**00100033**
					GRA	SPO		
Clostridium novyi biovar A		97.37	0.43	GLU 99	99	99		
Pseudoflavonifractor capillosus		2.63	-.03		1	1		
00100040	**Low discrimination**							**00100040**
					COC	SPO		
Clostridium botulinum biovar A		46.83	0.69	GLU 90	1	99		
Peptostreptococcus anaerobius		32.99	0.61		99	1		
Clostridium novyi biovar A		4.41	0.63		1	99		
00100041	**Low discrimination**							**00100041**
					GRA	SPO		
Clostridium novyi biovar A		62.48	0.63	GLU 99	99	99		
Campylobacter ureolyticus		16.92	0.32		1	1		
Clostridium botulinum biovar A		6.70	0.34		99	99		
00100042	**Low discrimination**							**00100042**

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Clostridium novyi biovar A		27.50	0.50	GLU 99				
Clostridium botulinum biovar A		16.71	0.34					
Clostridium sporogenes		16.20	0.32					
00100043	**Low discrimination**							**00100043**
					COC	SPO		
Clostridium novyi biovar A		87.79	0.50	GLU 99	1	99		
Anaerococcus prevotii		6.82	0.13		99	1		
Peptococcus niger		2.59	0.02		99	1		
00100050	**Low discrimination**							**00100050**
					COC	SPO		
Clostridium novyi biovar A		82.17	0.63	GLU 99	1	99		
Clostridium botulinum biovar A		8.81	0.34		1	99		
Peptostreptococcus anaerobius		6.21	0.27		99	1		
00100051	**Very good identification**							**00100051**
					GRA	SPO		
Clostridium novyi biovar A		99.54	0.63	GLU 99	99	99		
Campylobacter ureolyticus		0.27	-.03		1	1		
Clostridium botulinum biovar A		0.11	0.00		99	99		
00100052	**Good identification**							**00100052**
Clostridium novyi biovar A		98.39	0.50	GLU 99				
Clostridium botulinum biovar A		0.60	0.00					
Clostridium sporogenes		0.59	-.02					
00100053	**Good identification**							**00100053**
Clostridium novyi biovar A		100.0	0.50	GLU 99				
00100060	**Low discrimination**							**00100060**
					COC	SPO		
Clostridium novyi biovar A		48.94	0.53	GLU 99	1	99		
Clostridium botulinum biovar A		20.99	0.34		1	99		
Peptostreptococcus anaerobius		14.79	0.27		99	1		
00100061	**Good identification**							**00100061**
					COC	GRA	SPO	
Clostridium novyi biovar A		96.95	0.53	GLU 99	1	99	99	
Peptococcus niger		1.27	-.02		99	99	1	
Campylobacter ureolyticus		1.06	-.03		1	1	1	
00100063	**Good identification**							**00100063**
					GRA	SPO		
Clostridium novyi biovar A		96.10	0.40	GLU 99	99	99		
Pseudoflavonifractor capillosus		3.90	-.03		1	1		
00100070	**Good identification**							**00100070**
					COC	SPO		
Clostridium novyi biovar A		99.27	0.53	GLU 99	1	99		
Clostridium botulinum biovar A		0.43	0.00		1	99		
Peptostreptococcus anaerobius		0.30	-.07		99	1		
00100071	**Good identification**							**00100071**
Clostridium novyi biovar A		100.0	0.53	GLU 99				
00100072	**Good identification**							**00100072**
					GRA	SPO		
Clostridium novyi biovar A		96.10	0.40	GLU 99	99	99		

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Pseudoflavonifractor capillosus		3.90	-.03		1	1	
00100073	**Good identification**						**00100073**
Clostridium novyi biovar A		100.0	0.40	GLU 99			
00100100	**Low discrimination**						**00100100**
					COC	SPO	
Peptostreptococcus anaerobius		66.51	0.61	LAC 1	99	1	
Clostridium histolyticum		3.77	0.32		1	90	
Clostridium limosum		3.77	0.32		1	99	
00100200	**Low discrimination**						**00100200**
					COC	SPO	
Peptostreptococcus anaerobius		66.80	0.61	RHA 1	99	1	
Clostridium histolyticum		3.79	0.32		1	90	
Clostridium limosum		3.79	0.32		1	99	
00100400	**Low discrimination**						**00100400**
					COC	SPO	
Peptostreptococcus anaerobius		66.75	0.61	ARA 1	99	1	
Clostridium histolyticum		3.79	0.32		1	90	
Clostridium limosum		3.79	0.32		1	99	
00101000	**Low discrimination**						**00101000**
					COC	SPO	
Peptostreptococcus anaerobius		61.86	0.61	GAL 1	99	1	
Atopobium minutum		4.23	0.54		1	1	
Clostridium histolyticum		3.51	0.32		1	90	
00101002	**Low discrimination**						**00101002**
					COC	SPO	
Atopobium minutum		46.71	0.51	MLT 1	1	1	
Clostridium sporogenes		14.26	0.32		1	99	
Peptostreptococcus anaerobius		10.36	0.27		99	1	
00101003	**Low discrimination**						**00101003**
					COC	SPO	
Clostridium novyi biovar A		51.02	0.37	GLU 99	1	99	
Anaerococcus prevotii		23.76	0.13	GAL 10	99	1	
Peptococcus niger		9.03	0.02		99	1	
00101010	**Low discrimination**						**00101010**
					COC	SPO	
Clostridium novyi biovar A		58.45	0.50	GLU 99	1	99	
Peptostreptococcus anaerobius		26.49	0.27	GAL 10	99	1	
Atopobium minutum		1.81	0.20		1	1	
00101011	**Good identification**						**00101011**
					GRA	SPO	
Clostridium novyi biovar A		97.18	0.50	GLU 99	99	99	
Campylobacter ureolyticus		1.60	-.03	GAL 10	1	1	
Clostridium sordellii		0.78	-.03		99	99	
00101012	**Low discrimination**						**00101012**
					SPO		
Clostridium novyi biovar A		84.84	0.37	GLU 99	99		
Atopobium minutum		9.93	0.17	GAL 10	1		
Clostridium sporogenes		3.03	-.02		99		
00101013	**Good identification**						**00101013**
Clostridium novyi biovar A		100.0	0.37	GLU 99			

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00101020	**Low discrimination**							**00101020**
					COC	SPO		
Atopobium minutum		44.55	0.46	MLT 1	1	1		
Peptostreptococcus anaerobius		19.76	0.27		99	1		
Clostridium novyi biovar A		10.90	0.39		1	99		
00101021	**Low discrimination**							**00101021**
					COC	GRA	SPO	
Clostridium novyi biovar A		80.90	0.39	GLU 99	1	99	99	
Peptococcus niger		6.36	-.02	GAL 10	99	99	1	
Campylobacter ureolyticus		5.31	-.03		1	1	1	
00101022	**Acceptable identification**							**00101022**
					GRA	SPO		
Pseudoflavonifractor capillosus		93.81	0.50	MLT 1	1	1		
Atopobium minutum		5.47	0.43	GAL 10	99	1		
Clostridium novyi biovar A		0.35	0.26		99	99		
00101030	**Acceptable identification**							**00101030**
					COC	SPO		
Clostridium novyi biovar A		94.38	0.39	GLU 99	1	99		
Atopobium minutum		3.90	0.11	GAL 10	1	1		
Peptostreptococcus anaerobius		1.73	-.07		99	1		
00101031	**Good identification**							**00101031**
Clostridium novyi biovar A		100.0	0.39	GLU 99				
00101040	**Low discrimination**							**00101040**
					COC	SPO		
Clostridium novyi biovar A		32.61	0.47	GLU 99	1	99		
Clostridium botulinum biovar A		31.47	0.34	GAL 10	1	99		
Peptostreptococcus anaerobius		22.17	0.27		99	1		
00101041	**Acceptable identification**							**00101041**
					GRA	SPO		
Clostridium novyi biovar A		91.65	0.47	GLU 99	99	99		
Clostridium sordellii		4.58	0.07	GAL 10	99	99		
Campylobacter ureolyticus		2.26	-.03		1	1		
00101050	**Good identification**							**00101050**
					COC	SPO		
Clostridium novyi biovar A		98.37	0.47	GLU 99	1	99		
Clostridium botulinum biovar A		0.96	0.00	GAL 10	1	99		
Peptostreptococcus anaerobius		0.68	-.07		99	1		
00101051	**Good identification**							**00101051**
Clostridium novyi biovar A		100.0	0.47	GLU 99				
00101060	**Low discrimination**							**00101060**
					SPO			
Clostridium novyi biovar A		88.62	0.36	GLU 99	99			
Atopobium minutum		5.49	0.11	GAL 10	1			
Clostridium botulinum biovar A		3.46	0.00		99			
00101061	**Good identification**							**00101061**
Clostridium novyi biovar A		100.0	0.36	GLU 99				
00101070	**Good identification**							**00101070**
Clostridium novyi biovar A		100.0	0.36	GLU 99				

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00101071	**Good identification**							**00101071**
Clostridium novyi biovar A		100.0	0.36	GLU 99				
00102000	**Low discrimination**							**00102000**
					COC	SPO		
Peptostreptococcus anaerobius		65.95	0.61	MAN 1	99	1		
Clostridium histolyticum		3.74	0.32		1	90		
Clostridium limosum		3.74	0.32		1	99		
00104000	**Low discrimination**							**00104000**
					COC	SPO		
Peptostreptococcus anaerobius		66.35	0.61	XYL 1	99	1		
Clostridium histolyticum		3.76	0.32		1	90		
Clostridium limosum		3.76	0.32		1	99		
00104040	**Low discrimination**							**00104040**
					COC	SPO		
Clostridium glycolicum		80.57	0.50	GLU 99	1	99		
Clostridium botulinum biovar A		9.17	0.34	FRU 90	1	99		
Peptostreptococcus anaerobius		6.46	0.27		99	1		
00104100	**Low discrimination**							**00104100**
					COC	SPO		
Actinomyces meyeri		43.60	0.37	GLU 93	1	1		
Peptostreptococcus anaerobius		38.26	0.27	SUC 93	99	1		
Clostridium histolyticum		2.17	-.03	FRU 95	1	90		
00105100	**Low discrimination**							**00105100**
					COC			
Actinomyces meyeri		84.44	0.40	GLU 93	1			
Blautia hansenii		15.06	0.19	SUC 93	99			
Peptostreptococcus anaerobius		0.50	-.07	FRU 95	99			
00110000	**Low discrimination**							**00110000**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		24.57	0.61	FRU 1	99	99	1	
Clostridium sporogenes		17.90	0.61		1	99	99	
Prevotella disiens		16.55	0.57		1	1	1	
00110001	**Low discrimination**							**00110001**
					GRA	SPO		
Clostridium sordellii		47.83	0.44	IND 99	99	99		
Campylobacter ureolyticus		17.17	0.32	GLU 95	1	1		
Clostridium novyi biovar A		10.57	0.50		99	99		
00110002	**Low discrimination**							**00110002**
					COC	SPO	GRA	
Clostridium sporogenes		52.36	0.53		1	99	99	
Atopobium minutum		27.20	0.58		1	1	99	
Megasphaera elsdenii		8.80	0.47		99	1	1	
00110003	**Low discrimination**							**00110003**
					COC	SPO		
Clostridium novyi biovar A		32.54	0.37	GLU 99	1	99		
Clostridium sporogenes		20.30	0.19	FRU 10	1	99		
Anaerococcus prevotii		15.16	0.13		99	1		
00110010	**Low discrimination**							**00110010**
					COC	SPO		
Clostridium novyi biovar A		35.24	0.50	GLU 99	1	99		
Peptostreptococcus anaerobius		15.97	0.27	FRU 10	99	1		

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Clostridium sporogenes		11.64	0.27		1	99	
00110011	**Acceptable identification**						**00110011**
					GRA	SPO	
Clostridium novyi biovar A		93.75	0.50	GLU 99	99	99	
Clostridium sordellii		4.29	0.10	FRU 10	99	99	
Campylobacter ureolyticus		1.54	-.03		1	1	
00110012	**Low discrimination**						**00110012**
					SPO		
Clostridium novyi biovar A		48.11	0.37	GLU 99	99		
Clostridium sporogenes		30.01	0.19	FRU 10	99		
Atopobium minutum		15.59	0.24		1		
00110013	**Good identification**						**00110013**
Clostridium novyi biovar A		100.0	0.37	GLU 99			
00110020	**Low discrimination**						**00110020**
					COC	SPO	
Atopobium minutum		48.03	0.53	MLT 1	1	1	
Peptostreptococcus anaerobius		7.69	0.27		99	1	
Clostridium cadaveris		6.02	0.27		1	99	
00110021	**Low discrimination**						**00110021**
					SPO		
Clostridium novyi biovar A		68.50	0.39	GLU 99	99		
Clostridium sordellii		12.53	0.10	FRU 10	99		
Atopobium minutum		7.83	0.19		1		
00110022	**Low discrimination**						**00110022**
					GRA	SPO	
Pseudoflavonifractor capillosus		84.61	0.50	MLT 1	1	1	
Atopobium minutum		13.66	0.50	FRU 10	99	1	
Clostridium sporogenes		0.80	0.19		99	99	
00110030	**Low discrimination**						**00110030**
					COC	SPO	
Clostridium novyi biovar A		88.29	0.39	GLU 99	1	99	
Atopobium minutum		10.09	0.19	FRU 10	1	1	
Peptostreptococcus anaerobius		1.62	-.07		99	1	
00110031	**Good identification**						**00110031**
Clostridium novyi biovar A		100.0	0.39	GLU 99			
00110040	**Low discrimination**						**00110040**
					COC	GRA	
Megasphaera elsdenii		54.33	0.60	GLU 99	99	1	
Clostridium botulinum biovar A		20.11	0.48		1	99	
Clostridium glycolicum		13.24	0.44		1	99	
00110041	**Low discrimination**						**00110041**
					COC	GRA	
Clostridium novyi biovar A		63.17	0.47	GLU 99	1	99	
Clostridium sordellii		17.87	0.20	FRU 10	1	99	
Megasphaera elsdenii		10.51	0.26		99	1	
00110042	**Low discrimination**						**00110042**
					COC	GRA	
Megasphaera elsdenii		75.93	0.44	GLU 99	99	1	
Clostridium novyi biovar A		7.32	0.34	BGL 10	1	99	
Clostridium sporogenes		6.85	0.19		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
00110050	**Low discrimination**							**00110050**
					COC	GRA		
Clostridium novyi biovar A		77.33	0.47	GLU 99	1	99		
Megasphaera elsdenii		12.87	0.26	FRU 10	99	1		
Clostridium botulinum biovar A		4.76	0.14		1	99		
00110051	**Good identification**							**00110051**
Clostridium novyi biovar A		100.0	0.47	GLU 99				
00110060	**Low discrimination**							**00110060**
					COC	SPO	GRA	
Clostridium novyi biovar A		38.12	0.36	GLU 99	1	99	99	
Megasphaera elsdenii		25.38	0.26	FRU 10	99	1	1	
Pseudoramibacter alactolyticus		10.64	0.09		1	1	99	
00110061	**Good identification**							**00110061**
Clostridium novyi biovar A		100.0	0.36	GLU 99				
00110070	**Good identification**							**00110070**
Clostridium novyi biovar A		96.62	0.36	GLU 99				
Clostridium botulinum biovar B		3.38	-.03	FRU 10				
00110071	**Good identification**							**00110071**
Clostridium novyi biovar A		100.0	0.36	GLU 99				
00110100	**Low discrimination**							**00110100**
					COC	SPO		
Actinomyces meyeri		73.33	0.54	GLU 93	1	1		
Peptostreptococcus anaerobius		6.57	0.27	SUC 93	99	1		
Clostridium sporogenes		4.79	0.27		1	99		
00110120	**Low discrimination**							**00110120**
					GRA			
Actinomyces meyeri		73.09	0.39	GLU 93	99			
Prevotella bivia		23.40	0.15	SUC 93	1			
Atopobium minutum		3.03	0.19	NAG 12	99			
00110500	**Good identification**							**00110500**
					COC			
Actinomyces meyeri		99.20	0.43	GLU 93	1			
Lactobacillus fermentum		0.45	0.14	SUC 93	1			
Peptostreptococcus anaerobius		0.36	-.07		99			
00111000	**Low discrimination**							**00111000**
					GRA			
Prevotella intermedia		57.74	0.57	GLU 90	1			
Atopobium minutum		17.05	0.53		99			
Actinomyces meyeri		11.42	0.46		99			
00111002	**Low discrimination**							**00111002**
					SPO			
Atopobium minutum		73.93	0.50	MLT 1	1			
Lactobacillus fermentum		13.48	0.44		1			
Clostridium sporogenes		4.31	0.19		99			
00111020	**Low discrimination**							**00111020**
Atopobium minutum		63.87	0.45	MLT 1				
Actinomyces meyeri		17.51	0.32					

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Lactobacillus fermentum	7.77	0.36			
00111100	**Good identification**					**00111100**
					GRA	
	Actinomyces meyeri	91.39	0.57	GLU 93	99	
	Lactobacillus fermentum	6.77	0.48	SUC 93	99	
	Prevotella intermedia	1.17	0.22		1	
00111102	**Low discrimination**					**00111102**
	Lactobacillus fermentum	68.88	0.48	GLU 99		
	Actinomyces meyeri	28.78	0.31	SUC 87		
	Atopobium minutum	2.34	0.16			
00111120	**Low discrimination**					**00111120**
					GRA	
	Actinomyces meyeri	73.54	0.42	GLU 93	99	
	Lactobacillus fermentum	13.31	0.40	SUC 93	99	
	Prevotella melaninogenica	7.85	0.17	NAG 12	1	
00111122	**Low discrimination**					**00111122**
					GRA	
	Lactobacillus fermentum	69.91	0.40	GLU 99	99	
	Actinomyces meyeri	11.95	0.16	SUC 87	99	
	Prevotella melaninogenica	7.85	0.05		1	
00111400	**Low discrimination**					**00111400**
					GRA	
	Actinomyces meyeri	49.06	0.36	GLU 93	99	
	Lactobacillus fermentum	35.61	0.44	SUC 93	99	
	Prevotella intermedia	10.02	0.22		1	
00111402	**Acceptable identification**					**00111402**
	Lactobacillus fermentum	91.07	0.44	GLU 99		
	Atopobium minutum	5.05	0.16	SUC 87		
	Actinomyces meyeri	3.88	0.10			
00111420	**Low discrimination**					**00111420**
	Lactobacillus fermentum	60.73	0.36	GLU 99		
	Actinomyces meyeri	34.23	0.21	SUC 87		
	Atopobium minutum	5.05	0.11			
00111422	**Low discrimination**					**00111422**
					GRA	
	Lactobacillus fermentum	85.49	0.36	GLU 99	99	
	Pseudoflavonifractor capillosus	9.78	-.01	SUC 87	1	
	Atopobium minutum	4.74	0.08		99	
00111500	**Low discrimination**					**00111500**
	Actinomyces meyeri	77.16	0.46	GLU 93		
	Lactobacillus fermentum	22.84	0.48	SUC 93		
00111502	**Acceptable identification**					**00111502**
	Lactobacillus fermentum	90.54	0.48	GLU 99		
	Actinomyces meyeri	9.46	0.21	SUC 87		
00111522	**Good identification**					**00111522**
	Lactobacillus fermentum	95.90	0.40	GLU 99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		4.10	0.06	SUC 87				
00112000	**Low discrimination**							**00112000**
					COC	GRA		
Megasphaera elsdenii		86.33	0.63	GLU 99	99	1		
Pseudoramibacter alactolyticus		2.97	0.27		1	99		
Peptostreptococcus anaerobius		2.38	0.27		99	99		
00112002	**Low discrimination**							**00112002**
					COC	SPO	GRA	
Megasphaera elsdenii		88.63	0.47	GLU 99	99	1	1	
Clostridium sporogenes		5.33	0.19	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		3.05	0.11		1	1	99	
00112040	**Very good identification**							**00112040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.03	0.60	GLU 99	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.17		1	1	99	
Clostridium botulinum biovar A		0.37	0.14		1	99	99	
00112042	**Good identification**							**00112042**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	GLU 99	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
00114000	**Low discrimination**							**00114000**
					COC	SPO		
Clostridium glycolicum		37.57	0.44	GLU 99	1	99		
Actinomyces meyeri		34.43	0.48	SOR 95	1	1		
Peptostreptococcus anaerobius		6.36	0.27		99	1		
00114040	**Very good identification**							**00114040**
					COC	GRA		
Clostridium glycolicum		99.63	0.66	GLU 99	1	99		
Megasphaera elsdenii		0.22	0.26		99	1		
Clostridium botulinum biovar A		0.08	0.14		1	99		
00114100	**Very good identification**							**00114100**
					COC	SPO		
Actinomyces meyeri		99.68	0.59	GLU 93	1	1		
Clostridium glycolicum		0.28	0.10	SUC 93	1	99		
Peptostreptococcus anaerobius		0.05	-.07		99	1		
00114120	**Good identification**							**00114120**
Actinomyces meyeri		100.0	0.44	GLU 93				
00114400	**Good identification**							**00114400**
					COC	SPO		
Actinomyces meyeri		95.10	0.38	GLU 93	1	1		
Clostridium glycolicum		4.19	0.10	SUC 93	1	99		
Peptostreptococcus anaerobius		0.71	-.07		99	1		
00114500	**Good identification**							**00114500**
Actinomyces meyeri		100.0	0.48	GLU 93				
00115000	**Low discrimination**							**00115000**
					SPO			
Actinomyces meyeri		85.04	0.51	GLU 93	1			
Clostridium glycolicum		6.87	0.28	SUC 93	99			
Lactobacillus fermentum		4.67	0.41		1			
00115002	**Low discrimination**							**00115002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		60.33	0.41	GLU 99		
Actinomyces meyeri		33.99	0.25	SUC 87		
Atopobium minutum		5.69	0.16			
00115020	**Low discrimination**					**00115020**
Actinomyces meyeri		86.72	0.36	GLU 93		
Lactobacillus fermentum		11.64	0.32	SUC 93		
Atopobium minutum		1.65	0.11	NAG 12		
00115040	**Good identification**					**00115040**
					SPO	
Clostridium glycolicum		99.35	0.50	GLU 99	99	
Actinomyces meyeri		0.65	0.17	GAL 10	1	
00115100	**Very good identification**					**00115100**
Actinomyces meyeri		97.81	0.62	GLU 93		
Lactobacillus fermentum		2.19	0.44	SUC 93		
00115102	**Low discrimination**					**00115102**
Actinomyces meyeri		58.00	0.36	GLU 93		
Lactobacillus fermentum		42.00	0.44	SUC 93		
00115120	**Acceptable identification**					**00115120**
Actinomyces meyeri		94.81	0.47	GLU 93		
Lactobacillus fermentum		5.19	0.36	SUC 93		
00115122	**Low discrimination**					**00115122**
Lactobacillus fermentum		63.90	0.36	GLU 99		
Actinomyces meyeri		36.10	0.21	SUC 87		
00115400	**Low discrimination**					**00115400**
Actinomyces meyeri		81.99	0.41	GLU 93		
Lactobacillus fermentum		18.01	0.41	SUC 93		
00115402	**Low discrimination**					**00115402**
Lactobacillus fermentum		87.66	0.41	GLU 99		
Actinomyces meyeri		12.35	0.15	SUC 87		
00115500	**Acceptable identification**					**00115500**
Actinomyces meyeri		91.78	0.51	GLU 93		
Lactobacillus fermentum		8.22	0.44	SUC 93		
00115502	**Low discrimination**					**00115502**
Lactobacillus fermentum		74.33	0.44	GLU 99		
Actinomyces meyeri		25.67	0.25	SUC 87		
00115520	**Low discrimination**					**00115520**
Actinomyces meyeri		82.04	0.36	GLU 93		
Lactobacillus fermentum		17.96	0.36	SUC 93		
00115522	**Low discrimination**					**00115522**
Lactobacillus fermentum		87.62	0.36	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		12.38	0.11	SUC 87			
00120000	**Low discrimination**						**00120000**
					COC	SPO	
Clostridium sporogenes		34.65	0.66		1	99	
Clostridium botulinum biovar A		29.91	0.67		1	99	
Peptostreptococcus anaerobius		25.18	0.61		99	1	
00120002	**Good identification**						**00120002**
					COC	SPO	
Clostridium sporogenes		91.11	0.58		1	99	
Clostridium botulinum biovar A		2.38	0.33		1	99	
Peptostreptococcus anaerobius		2.01	0.27		99	1	
00120040	**Good identification**						**00120040**
					COC	SPO	
Clostridium botulinum biovar A		92.22	0.58	GLU 90	1	99	
Clostridium sporogenes		3.61	0.32		1	99	
Peptostreptococcus anaerobius		2.63	0.27		99	1	
00130000	**Low discrimination**						**00130000**
					COC	SPO	
Clostridium sporogenes		69.55	0.53		1	99	
Clostridium botulinum biovar A		21.72	0.47		1	99	
Peptostreptococcus anaerobius		2.89	0.27		99	1	
00130002	**Good identification**						**00130002**
					SPO		
Clostridium sporogenes		96.94	0.45		99		
Atopobium minutum		1.53	0.24		1		
Clostridium botulinum biovar A		0.92	0.13		99		
00130040	**Low discrimination**						**00130040**
					COC	GRA	
Clostridium botulinum biovar A		75.41	0.38	GLU 90	1	99	
Megasphaera elsdenii		8.23	0.26	FRU 6	99	1	
Clostridium sporogenes		8.17	0.19		1	99	
00140000	**Low discrimination**						**00140000**
					COC	SPO	
Peptostreptococcus anaerobius		66.80	0.61	CEL 1	99	1	
Clostridium histolyticum		3.79	0.32		1	90	
Clostridium limosum		3.79	0.32		1	99	
00200000	**Low discrimination**						**00200000**
					COC	SPO	
Clostridium histolyticum		13.80	0.66	SAL 1	1	90	
Clostridium limosum		13.80	0.66		1	99	
Finegoldia magna		13.80	0.66		99	1	
00200001	**Low discrimination**						**00200001**
					COC	GRA	
Campylobacter ureolyticus		70.49	0.66	SAL 1	1	1	
Anaerococcus prevotii		14.27	0.58		99	99	
Peptococcus niger		7.68	0.49		99	99	
00200002	**Low discrimination**						**00200002**
					GRA		
Peptococcus niger		31.02	0.53	SAL 1	99		
Acidaminococcus fermentans		25.26	0.50		1		
Parvimonas micra		22.96	0.50		99		
00200003	**Low discrimination**						**00200003**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Anaerococcus prevotii		59.51	0.47	SAL 1	99	99		
Peptococcus niger		22.61	0.36		99	99		
Campylobacter ureolyticus		11.88	0.32		1	1		
00200020	**Low discrimination**							**00200020**
					COC	SPO		
Peptococcus niger		47.86	0.50	SAL 1	99	1		
Atopobium minutum		7.46	0.54	NAG 10	1	1		
Clostridium histolyticum		6.19	0.32		1	90		
00200022	**Good identification**							**00200022**
					COC	GRA		
Pseudoflavonifractor capillosus		94.42	0.66	SAL 1	1	1		
Peptococcus niger		3.12	0.37		99	99		
Atopobium minutum		1.84	0.51		1	99		
00200031	**Low discrimination**							**00200031**
					COC	GRA	SPO	
Clostridium novyi biovar A		84.43	0.37	GLU 99	1	99	99	
Peptococcus niger		8.48	-.02	SAL 8	99	99	1	
Campylobacter ureolyticus		7.08	-.03		1	1	1	
00200050	**Low discrimination**							**00200050**
Clostridium novyi biovar A		73.39	0.45	GLU 99				
Clostridium histolyticum		3.62	-.03	SAL 8				
Clostridium limosum		3.62	-.03					
00200051	**Good identification**							**00200051**
					GRA	SPO		
Clostridium novyi biovar A		96.95	0.45	GLU 99	99	99		
Campylobacter ureolyticus		3.05	-.03	SAL 8	1	1		
00201000	**Low discrimination**							**00201000**
					SPO			
Atopobium minutum		13.54	0.54	SAL 1	1			
Clostridium histolyticum		11.24	0.32		90			
Clostridium limosum		11.24	0.32		99			
00201002	**Low discrimination**							**00201002**
					COC	GRA		
Atopobium minutum		65.45	0.51	SAL 1	1	99		
Peptococcus niger		10.11	0.19		99	99		
Acidaminococcus fermentans		8.23	0.15		99	1		
00201020	**Low discrimination**							**00201020**
					COC	GRA		
Atopobium minutum		66.36	0.46	SAL 1	1	99		
Peptococcus niger		12.91	0.15		99	99		
Pseudoflavonifractor capillosus		7.67	0.15		1	1		
00201022	**Acceptable identification**							**00201022**
					COC	GRA		
Pseudoflavonifractor capillosus		94.22	0.50	SAL 1	1	1		
Atopobium minutum		5.49	0.43	GAL 10	1	99		
Peptococcus niger		0.28	0.03		99	99		
00210000	**Low discrimination**							**00210000**
					GRA	SPO		
Atopobium minutum		25.71	0.61	SAL 1	99	1		
Fusobacterium varium		13.26	0.42		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium histolyticum		7.71	0.32		99	90	
00210002	**Low discrimination**						**00210002**
					COC	GRA	
Atopobium minutum		81.23	0.58	SAL 1	1	99	
Peptococcus niger		4.53	0.19		99	99	
Acidaminococcus fermentans		3.69	0.15		99	1	
00210020	**Low discrimination**						**00210020**
					COC		
Atopobium minutum		77.86	0.53	SAL 1	1		
Peptococcus niger		5.47	0.15		99		
Pseudoramibacter alactolyticus		5.12	0.19		1		
00210022	**Low discrimination**						**00210022**
					COC	GRA	
Pseudoflavonifractor capillosus		85.75	0.50	SAL 1	1	1	
Atopobium minutum		13.84	0.50	FRU 10	1	99	
Peptococcus niger		0.26	0.03		99	99	
00210100	**Low discrimination**						**00210100**
					GRA		
Fusobacterium mortiferum		29.92	0.37	GLU 99	1		
Atopobium minutum		16.37	0.27		99		
Actinomyces meyeri		8.64	0.19		99		
00211000	**Low discrimination**						**00211000**
					GRA		
Atopobium minutum		88.30	0.53	SAL 1	99		
Fusobacterium mortiferum		3.14	0.34		1		
Fusobacterium varium		1.38	0.08		1		
00211002	**Good identification**						**00211002**
					GRA		
Atopobium minutum		98.73	0.50	SAL 1	99		
Pseudoflavonifractor capillosus		0.69	-.01		1		
Fusobacterium mortiferum		0.59	0.17		1		
00211020	**Good identification**						**00211020**
					GRA		
Atopobium minutum		98.63	0.45	SAL 1	99		
Pseudoflavonifractor capillosus		1.37	-.01		1		
00211100	**Low discrimination**						**00211100**
					GRA		
Fusobacterium mortiferum		70.90	0.40	GLU 99	1		
Actinomyces meyeri		20.48	0.22		99		
Atopobium minutum		8.62	0.19		99		
00212010	**Good identification**						**00212010**
					SPO		
Clostridium difficile		99.81	0.43	GLU 99	99		
Pseudoramibacter alactolyticus		0.19	-.07		1		
00300000	**Low discrimination**						**00300000**
					COC	SPO	
Peptostreptococcus anaerobius		64.58	0.61	SAL 1	99	1	
Clostridium histolyticum		3.66	0.32		1	90	
Clostridium limosum		3.66	0.32		1	99	
00300010	**Low discrimination**						**00300010**
					COC	SPO	
Clostridium novyi biovar A		53.28	0.48	GLU 99	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptostreptococcus anaerobius		30.86	0.27	SAL 8	99	1		
Clostridium histolyticum		1.75	-.03		1	90		
00300011	**Good identification**							**00300011**
					COC	GRA	SPO	
Clostridium novyi biovar A		97.39	0.48	GLU 99	1	99	99	
Campylobacter ureolyticus		2.04	-.03	SAL 8	1	1	1	
Peptostreptococcus anaerobius		0.57	-.07		99	99	1	
00300021	**Low discrimination**							**00300021**
					COC	GRA	SPO	
Clostridium novyi biovar A		82.80	0.37	GLU 99	1	99	99	
Peptococcus niger		8.32	-.02	SAL 8	99	99	1	
Campylobacter ureolyticus		6.95	-.03		1	1	1	
00300030	**Good identification**							**00300030**
					COC	SPO		
Clostridium novyi biovar A		97.71	0.37	GLU 99	1	99		
Peptostreptococcus anaerobius		2.29	-.07	SAL 8	99	1		
00300031	**Good identification**							**00300031**
Clostridium novyi biovar A		100.0	0.37	GLU 99				
00300041	**Acceptable identification**							**00300041**
					GRA	SPO		
Clostridium novyi biovar A		95.00	0.45	GLU 99	99	99		
Campylobacter ureolyticus		2.99	-.03	SAL 8	1	1		
Clostridium botulinum biovar A		1.18	0.00		99	99		
00300050	**Good identification**							**00300050**
					COC	SPO		
Clostridium novyi biovar A		97.92	0.45	GLU 99	1	99		
Clostridium botulinum biovar A		1.22	0.00	SAL 8	1	99		
Peptostreptococcus anaerobius		0.86	-.07		99	1		
00300051	**Good identification**							**00300051**
Clostridium novyi biovar A		100.0	0.45	GLU 99				
00400000	**Low discrimination**							**00400000**
					COC	SPO		
Clostridium histolyticum		12.98	0.66	RAF 1	1	90		
Clostridium limosum		12.98	0.66		1	99		
Finegoldia magna		12.98	0.66		99	1		
00400001	**Low discrimination**							**00400001**
					COC	GRA		
Anaerococcus prevotii		84.64	0.84		99	99		
Campylobacter ureolyticus		12.67	0.66		1	1		
Peptococcus niger		1.38	0.49		99	99		
00400002	**Low discrimination**							**00400002**
					GRA			
Peptococcus niger		24.12	0.53	RAF 1	99			
Anaerococcus prevotii		23.03	0.57		99			
Acidaminococcus fermentans		19.64	0.50		1			
00400003	**Very good identification**							**00400003**
					COC	GRA		
Anaerococcus prevotii		98.02	0.73		99	99		
Peptococcus niger		1.13	0.36		99	99		
Campylobacter ureolyticus		0.59	0.32		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
00400011	**Low discrimination**						**00400011**
					COC	GRA	
Anaerococcus prevotii		84.07	0.50	MLZ 1	99	99	
Campylobacter ureolyticus		12.59	0.32		1	1	
Peptococcus niger		1.37	0.15		99	99	
00400013	**Good identification**						**00400013**
					COC	GRA	
Anaerococcus prevotii		97.71	0.39	MLZ 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.59	-.03		1	1	
00400020	**Low discrimination**						**00400020**
					COC	SPO	
Peptococcus niger		46.78	0.50	RAF 1	99	1	
Atopobium minutum		7.29	0.54	NAG 10	1	1	
Clostridium histolyticum		6.05	0.32		1	90	
00400021	**Low discrimination**						**00400021**
					COC	GRA	
Anaerococcus prevotii		74.05	0.50	NAG 1	99	99	
Peptococcus niger		13.28	0.32		99	99	
Campylobacter ureolyticus		11.09	0.32		1	1	
00400022	**Good identification**						**00400022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.20	0.66	RAF 1	1	1	
Peptococcus niger		3.11	0.37		99	99	
Atopobium minutum		1.83	0.51		1	99	
00400023	**Low discrimination**						**00400023**
					COC	GRA	
Anaerococcus prevotii		65.28	0.39	NAG 1	99	99	
Pseudoflavonifractor capillosus		25.57	0.32		1	1	
Peptococcus niger		8.27	0.19		99	99	
00400041	**Low discrimination**						**00400041**
					COC	GRA	
Anaerococcus prevotii		84.22	0.50	SOR 1	99	99	
Campylobacter ureolyticus		12.61	0.32		1	1	
Peptococcus niger		1.37	0.15		99	99	
00400043	**Good identification**						**00400043**
					COC	GRA	
Anaerococcus prevotii		97.90	0.39	SOR 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.59	-.03		1	1	
00400101	**Low discrimination**						**00400101**
					COC	GRA	
Anaerococcus prevotii		84.67	0.50	LAC 1	99	99	
Campylobacter ureolyticus		12.68	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00400103	**Good identification**						**00400103**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	LAC 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00400201	**Low discrimination**						**00400201**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Anaerococcus prevotii		84.67	0.50	RHA 1	99	99	
Campylobacter ureolyticus		12.68	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00400203	**Good identification**						**00400203**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	RHA 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00400401	**Low discrimination**						**00400401**
					COC	GRA	
Anaerococcus prevotii		84.67	0.50	ARA 1	99	99	
Campylobacter ureolyticus		12.68	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00400403	**Good identification**						**00400403**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	ARA 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00401000	**Low discrimination**						**00401000**
					SPO		
Atopobium minutum		12.92	0.54	RAF 1	1		
Clostridium histolyticum		10.73	0.32		90		
Clostridium limosum		10.73	0.32		99		
00401001	**Low discrimination**						**00401001**
					COC	GRA	
Anaerococcus prevotii		84.32	0.50	GAL 1	99	99	
Campylobacter ureolyticus		12.63	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00401002	**Low discrimination**						**00401002**
					COC		
Atopobium minutum		60.12	0.51	RAF 1	1		
Peptococcus niger		9.29	0.19		99		
Anaerococcus prevotii		8.87	0.23		99		
00401003	**Good identification**						**00401003**
					COC		
Anaerococcus prevotii		97.54	0.39	GAL 1	99		
Peptococcus niger		1.12	0.02		99		
Atopobium minutum		0.74	0.17		1		
00401020	**Low discrimination**						**00401020**
					COC	GRA	
Atopobium minutum		66.78	0.46	RAF 1	1	99	
Peptococcus niger		12.99	0.15		99	99	
Pseudoflavonifractor capillosus		7.71	0.15		1	1	
00401022	**Acceptable identification**						**00401022**
					COC	GRA	
Pseudoflavonifractor capillosus		94.22	0.50	RAF 1	1	1	
Atopobium minutum		5.49	0.43	GAL 10	1	99	
Peptococcus niger		0.28	0.03		99	99	
00402001	**Low discrimination**						**00402001**
					COC	GRA	
Anaerococcus prevotii		84.67	0.50	MAN 1	99	99	
Campylobacter ureolyticus		12.68	0.32		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptococcus niger		1.38	0.15		99	99	
00402003	**Good identification**						**00402003**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	MAN 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00404001	**Low discrimination**						**00404001**
					COC	GRA	
Anaerococcus prevotii		84.67	0.50	XYL 1	99	99	
Campylobacter ureolyticus		12.68	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00404003	**Good identification**						**00404003**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	XYL 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00410000	**Low discrimination**						**00410000**
					GRA	SPO	
Atopobium minutum		24.86	0.61	RAF 1	99	1	
Fusobacterium varium		12.82	0.42		1	1	
Clostridium histolyticum		7.45	0.32		99	90	
00410001	**Low discrimination**						**00410001**
					COC	GRA	
Anaerococcus prevotii		83.65	0.50	FRU 1	99	99	
Campylobacter ureolyticus		12.53	0.32		1	1	
Peptococcus niger		1.36	0.15		99	99	
00410002	**Low discrimination**						**00410002**
					COC		
Atopobium minutum		78.12	0.58	RAF 1	1		
Peptococcus niger		4.36	0.19		99		
Anaerococcus prevotii		4.16	0.23		99		
00410003	**Good identification**						**00410003**
					COC		
Anaerococcus prevotii		96.28	0.39	FRU 1	99		
Atopobium minutum		2.03	0.24		1		
Peptococcus niger		1.11	0.02		99		
00410020	**Low discrimination**						**00410020**
					COC		
Atopobium minutum		78.07	0.53	RAF 1	1		
Peptococcus niger		5.48	0.15		99		
Pseudoramibacter alactolyticus		5.13	0.19		1		
00410022	**Low discrimination**						**00410022**
					COC	GRA	
Pseudoflavonifractor capillosus		85.75	0.50	RAF 1	1	1	
Atopobium minutum		13.84	0.50	FRU 10	1	99	
Peptococcus niger		0.26	0.03		99	99	
00410100	**Low discrimination**						**00410100**
					GRA		
Fusobacterium mortiferum		29.24	0.37	GLU 99	1		
Atopobium minutum		16.00	0.27		99		
Actinomyces meyeri		8.45	0.19		99		
00411000	**Low discrimination**						**00411000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Atopobium minutum		79.95	0.53	RAF 1	99	
Lactobacillus fermentum		5.29	0.40		99	
Prevotella intermedia		4.10	0.25		1	
00411002	**Low discrimination**					**00411002**
					GRA	
Atopobium minutum		89.92	0.50	RAF 1	99	
Lactobacillus fermentum		8.92	0.40		99	
Pseudoflavonifractor capillosus		0.63	-.01		1	
00411020	**Acceptable identification**					**00411020**
					GRA	
Atopobium minutum		92.59	0.45	RAF 1	99	
Lactobacillus fermentum		6.12	0.32		99	
Pseudoflavonifractor capillosus		1.29	-.01		1	
00411100	**Low discrimination**					**00411100**
					GRA	
Lactobacillus fermentum		47.34	0.44	GLU 99	99	
Fusobacterium mortiferum		36.46	0.40	SUC 87	1	
Actinomyces meyeri		10.54	0.22		99	
00411102	**Low discrimination**					**00411102**
					GRA	
Lactobacillus fermentum		87.11	0.44	GLU 99	99	
Fusobacterium mortiferum		7.45	0.24	SUC 87	1	
Atopobium minutum		5.44	0.16		99	
00411120	**Low discrimination**					**00411120**
					GRA	
Lactobacillus fermentum		77.93	0.35	GLU 99	99	
Prevotella melaninogenica		7.68	-.01	SUC 87	1	
Atopobium minutum		7.30	0.11		99	
00411122	**Low discrimination**					**00411122**
					GRA	
Lactobacillus fermentum		83.94	0.35	GLU 99	99	
Pseudoflavonifractor capillosus		10.82	-.01	SUC 87	1	
Atopobium minutum		5.24	0.08		99	
00411400	**Low discrimination**					**00411400**
Lactobacillus fermentum		86.75	0.40	GLU 99		
Atopobium minutum		13.25	0.19	SUC 87		
00411402	**Acceptable identification**					**00411402**
Lactobacillus fermentum		90.76	0.40	GLU 99		
Atopobium minutum		9.24	0.16	SUC 87		
00411500	**Acceptable identification**					**00411500**
Lactobacillus fermentum		94.73	0.44	GLU 99		
Actinomyces meyeri		5.27	0.11	SUC 87		
00411502	**Good identification**					**00411502**
Lactobacillus fermentum		100.0	0.44	GLU 99		
00411520	**Good identification**					**00411520**
Lactobacillus fermentum		100.0	0.35	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
00411522	**Good identification**						**00411522**
Lactobacillus fermentum		100.0	0.35	GLU 99			
00415000	**Low discrimination**						**00415000**
Lactobacillus fermentum		61.38	0.36	GLU 99			
Actinomyces meyeri		18.41	0.16	SUC 87			
Atopobium minutum		15.97	0.19				
00415002	**Low discrimination**						**00415002**
Lactobacillus fermentum		85.22	0.36	GLU 99			
Atopobium minutum		14.78	0.16	SUC 87			
00415100	**Low discrimination**						**00415100**
					GRA		
Lactobacillus fermentum		54.52	0.40	GLU 99	99		
Actinomyces meyeri		40.10	0.27	SUC 87	99		
Fusobacterium mortiferum		5.38	0.21		1		
00415102	**Good identification**						**00415102**
Lactobacillus fermentum		100.0	0.40	GLU 99			
00415400	**Acceptable identification**						**00415400**
Lactobacillus fermentum		93.02	0.36	GLU 99			
Actinomyces meyeri		6.98	0.06	SUC 87			
00415402	**Good identification**						**00415402**
Lactobacillus fermentum		100.0	0.36	GLU 99			
00415500	**Low discrimination**						**00415500**
Lactobacillus fermentum		84.47	0.40	GLU 99			
Actinomyces meyeri		15.53	0.16	SUC 87			
00415502	**Good identification**						**00415502**
Lactobacillus fermentum		100.0	0.40	GLU 99			
00420001	**Low discrimination**						**00420001**
					COC	GRA	
Anaerococcus prevotii		84.63	0.50	TRE 1	99	99	
Campylobacter ureolyticus		12.67	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00420003	**Good identification**						**00420003**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	TRE 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00440001	**Low discrimination**						**00440001**
					COC	GRA	
Anaerococcus prevotii		84.67	0.50	CEL 1	99	99	
Campylobacter ureolyticus		12.68	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00440003	**Good identification**						**00440003**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	CEL 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00500000	**Low discrimination**						**00500000**
					COC	SPO	
Peptostreptococcus anaerobius		65.51	0.61	RAF 1	99	1	
Clostridium histolyticum		3.72	0.32		1	90	
Clostridium limosum		3.72	0.32		1	99	
00500001	**Low discrimination**						**00500001**
					COC	GRA	
Anaerococcus prevotii		80.74	0.50	MLT 1	99	99	
Campylobacter ureolyticus		12.09	0.32		1	1	
Peptostreptococcus anaerobius		3.37	0.27		99	99	
00500003	**Good identification**						**00500003**
					COC	GRA	
Anaerococcus prevotii		97.33	0.39	MLT 1	99	99	
Peptococcus niger		1.12	0.02		99	99	
Campylobacter ureolyticus		0.59	-.03		1	1	
00501100	**Very good identification**						**00501100**
					COC		
Blautia hansenii		99.98	0.66	GLU 99	99		
Lactobacillus fermentum		0.01	0.18		1		
Peptostreptococcus anaerobius		0.01	-.07		99		
00501102	**Good identification**						**00501102**
					COC		
Blautia hansenii		99.89	0.50	GLU 99	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		
00501120	**Good identification**						**00501120**
					COC	GRA	
Blautia hansenii		98.74	0.50	GLU 99	99	99	
Prevotella melaninogenica		1.27	0.17	NAG 10	1	1	
00501500	**Good identification**						**00501500**
					COC		
Blautia hansenii		99.93	0.53	GLU 99	99		
Lactobacillus fermentum		0.07	0.18		1		
00501502	**Good identification**						**00501502**
					COC		
Blautia hansenii		99.40	0.37	GLU 99	99		
Lactobacillus fermentum		0.61	0.18	BGL 10	1		
00501520	**Good identification**						**00501520**
Blautia hansenii		100.0	0.37	GLU 99			
00505100	**Good identification**						**00505100**
					COC		
Blautia hansenii		99.90	0.53	GLU 99	99		
Actinomyces meyeri		0.06	0.06		1		
Lactobacillus fermentum		0.04	0.14		1		
00505102	**Good identification**						**00505102**
					COC		
Blautia hansenii		99.64	0.37	GLU 99	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
00505120	**Good identification**						**00505120**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		100.0	0.37	GLU 99				
00505500	**Good identification**							**00505500**
					COC			
Blautia hansenii		99.78	0.40	GLU 99	99			
Lactobacillus fermentum		0.23	0.14		1			
00510000	**Low discrimination**							**00510000**
					COC	GRA	SPO	
Prevotella intermedia		89.07	0.60	GLU 90	1	1	1	
Peptostreptococcus anaerobius		2.81	0.27		99	99	1	
Clostridium sporogenes		2.05	0.27		1	99	99	
00511000	**Very good identification**							**00511000**
					GRA			
Prevotella intermedia		95.78	0.60	GLU 90	1			
Lactobacillus fermentum		3.74	0.48		99			
Atopobium minutum		0.19	0.19		99			
00511002	**Low discrimination**							**00511002**
					GRA			
Lactobacillus fermentum		77.36	0.48	GLU 99	99			
Prevotella intermedia		20.02	0.25	SUC 87	1			
Atopobium minutum		2.63	0.16		99			
00511020	**Low discrimination**							**00511020**
					GRA			
Lactobacillus fermentum		53.17	0.40	GLU 99	99			
Prevotella intermedia		41.27	0.25	SUC 87	1			
Prevotella melaninogenica		2.85	-.01		1			
00511022	**Acceptable identification**							**00511022**
					GRA			
Lactobacillus fermentum		90.58	0.40	GLU 99	99			
Pseudoflavonifractor capillosus		6.35	-.01	SUC 87	1			
Atopobium minutum		3.08	0.08		99			
00511100	**Low discrimination**							**00511100**
					COC	GRA		
Lactobacillus fermentum		46.73	0.52	GLU 99	1	99		
Blautia hansenii		39.96	0.32	SUC 87	99	99		
Prevotella intermedia		7.41	0.25		1	1		
00511102	**Acceptable identification**							**00511102**
					COC			
Lactobacillus fermentum		91.32	0.52	GLU 99	1			
Blautia hansenii		8.68	0.15	SUC 87	99			
00511122	**Low discrimination**							**00511122**
					COC	GRA		
Lactobacillus fermentum		60.56	0.44	GLU 99	1	99		
Prevotella melaninogenica		37.53	0.21	SUC 87	1	1		
Blautia hansenii		1.92	-.01		99	99		
00511400	**Low discrimination**							**00511400**
					GRA			
Lactobacillus fermentum		78.91	0.48	GLU 99	99			
Prevotella intermedia		20.42	0.25	SUC 87	1			
Actinomyces meyeri		0.67	0.02		99			
00511402	**Good identification**							**00511402**
Lactobacillus fermentum		100.0	0.48	GLU 99				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
00511420	**Good identification**						**00511420**
Lactobacillus fermentum		100.0	0.40	GLU 99			
00511422	**Good identification**						**00511422**
Lactobacillus fermentum		100.0	0.40	GLU 99			
00511500	**Low discrimination**						**00511500**
					COC		
Lactobacillus fermentum		85.34	0.52	GLU 99	1		
Blautia hansenii		12.88	0.19	SUC 87	99		
Actinomyces meyeri		1.79	0.12		1		
00511502	**Good identification**						**00511502**
					COC		
Lactobacillus fermentum		98.35	0.52	GLU 99	1		
Blautia hansenii		1.65	0.03	SUC 87	99		
00511520	**Acceptable identification**						**00511520**
					COC	GRA	
Lactobacillus fermentum		91.03	0.44	GLU 99	1	99	
Blautia hansenii		4.58	0.03	SUC 87	99	99	
Prevotella melaninogenica		2.99	-.01		1	1	
00511522	**Good identification**						**00511522**
Lactobacillus fermentum		98.49	0.44	GLU 99			
Bifidobacterium longum subsp. longum		1.51	0.05	SUC 87			
00515000	**Low discrimination**						**00515000**
					GRA		
Lactobacillus fermentum		64.38	0.44	GLU 99	99		
Prevotella intermedia		28.36	0.25	SUC 87	1		
Actinomyces meyeri		7.26	0.17		99		
00515002	**Good identification**						**00515002**
Lactobacillus fermentum		100.0	0.44	GLU 99			
00515020	**Good identification**						**00515020**
Lactobacillus fermentum		95.59	0.36	GLU 99			
Actinomyces meyeri		4.41	0.02	SUC 87			
00515022	**Good identification**						**00515022**
Lactobacillus fermentum		100.0	0.36	GLU 99			
00515100	**Low discrimination**						**00515100**
					COC		
Lactobacillus fermentum		65.22	0.48	GLU 99	1		
Actinomyces meyeri		18.03	0.27	SUC 87	1		
Blautia hansenii		16.76	0.19		99		
00515102	**Good identification**						**00515102**
					COC		
Lactobacillus fermentum		96.42	0.48	GLU 99	1		
Blautia hansenii		2.75	0.03	SUC 87	99		
Actinomyces meyeri		0.82	0.02		1		
00515120	**Low discrimination**						**00515120**
					COC		
Lactobacillus fermentum		79.70	0.40	GLU 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		9.01	0.13	SUC 87	1		
Blautia hansenii		6.83	0.03		99		
00515122	**Good identification**						**00515122**
Lactobacillus fermentum		100.0	0.40	GLU 99			
00515400	**Good identification**						**00515400**
Lactobacillus fermentum		97.26	0.44	GLU 99			
Actinomyces meyeri		2.74	0.07	SUC 87			
00515402	**Good identification**						**00515402**
Lactobacillus fermentum		100.0	0.44	GLU 99			
00515420	**Good identification**						**00515420**
Lactobacillus fermentum		100.0	0.36	GLU 99			
00515422	**Good identification**						**00515422**
Lactobacillus fermentum		100.0	0.36	GLU 99			
00515500	**Low discrimination**						**00515500**
					COC		
Lactobacillus fermentum		89.73	0.48	GLU 99	1		
Actinomyces meyeri		6.20	0.17	SUC 87	1		
Blautia hansenii		4.07	0.06		99		
00515502	**Good identification**						**00515502**
Lactobacillus fermentum		100.0	0.48	GLU 99			
00515520	**Acceptable identification**						**00515520**
Lactobacillus fermentum		90.37	0.40	GLU 99			
Bifidobacterium longum subsp. longum		7.08	0.14	SUC 87			
Actinomyces meyeri		2.56	0.02				
00515522	**Acceptable identification**						**00515522**
Lactobacillus fermentum		92.74	0.40	GLU 99			
Bifidobacterium longum subsp. longum		7.26	0.14	SUC 87			
00600001	**Low discrimination**						**00600001**
					COC	GRA	
Anaerococcus prevotii		84.62	0.50	SAL 1	99	99	
Campylobacter ureolyticus		12.67	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
00600003	**Good identification**						**00600003**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	SAL 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
00610000	**Low discrimination**						**00610000**
					GRA		
Fusobacterium mortiferum		28.08	0.36	GLU 99	1		
Atopobium minutum		18.47	0.27		99		
Fusobacterium varium		9.52	0.08		1		
00610100	**Good identification**						**00610100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		100.0	0.42	GLU 99				
00611000	**Low discrimination**							**00611000**
					GRA			
Fusobacterium mortiferum		87.25	0.39	GLU 99	1			
Atopobium minutum		12.75	0.19		99			
00611100	**Good identification**							**00611100**
Fusobacterium mortiferum		100.0	0.45	GLU 99				
00631100	**Good identification**							**00631100**
Fusobacterium mortiferum		100.0	0.36	GLU 99				
00651100	**Good identification**							**00651100**
Fusobacterium mortiferum		100.0	0.37	GLU 99				
01000000	**Low discrimination**							**01000000**
					COC	SPO		
Peptostreptococcus anaerobius		52.15	1.00		99	1		
Clostridium botulinum biovar A		15.02	0.95		1	99		
Atopobium minutum		9.93	1.00		1	1		
01000001	**Low discrimination**							**01000001**
					COC	GRA	SPO	
Clostridium novyi biovar A		70.81	1.00		1	99	99	
Campylobacter ureolyticus		12.78	0.66		1	1	1	
Peptostreptococcus anaerobius		6.54	0.66		99	99	1	
01000002	**Low discrimination**							**01000002**
					SPO			
Atopobium minutum		64.65	0.97		1			
Clostridium sporogenes		11.79	0.74		99			
Clostridium novyi biovar A		9.84	0.87		99			
01000003	**Low discrimination**							**01000003**
					COC	SPO		
Clostridium novyi biovar A		85.26	0.87		1	99		
Atopobium minutum		5.66	0.63		1	1		
Anaerococcus prevotii		4.41	0.47		99	1		
01000010	**Low discrimination**							**01000010**
					COC	SPO		
Clostridium novyi biovar A		85.60	1.00		1	99		
Peptostreptococcus anaerobius		7.90	0.66		99	1		
Clostridium botulinum biovar A		2.28	0.61		1	99		
01000011	**Excellent identification**							**01000011**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.59	1.00		1	99	99	
Campylobacter ureolyticus		0.18	0.32		1	1	1	
Peptostreptococcus anaerobius		0.09	0.32		99	99	1	
01000012	**Good identification**							**01000012**
					SPO			
Clostridium novyi biovar A		91.52	0.87		99			
Atopobium minutum		6.08	0.63		1			
Clostridium sporogenes		1.11	0.40		99			
01000013	**Very good identification**							**01000013**
					COC	SPO		
Clostridium novyi biovar A		99.83	0.87		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		0.07	0.29		1	1	
Anaerococcus prevotii		0.05	0.13		99	1	
01000020	**Low discrimination**						**01000020**
					SPO		
Atopobium minutum		49.58	0.92		1		
Clostridium novyi biovar A		21.37	0.90		99		
Clostridium cadaveris		12.28	0.71		99		
01000021	**Good identification**						**01000021**
					COC	SPO	
Clostridium novyi biovar A		94.86	0.90		1	99	
Atopobium minutum		2.22	0.58		1	1	
Peptococcus niger		0.83	0.32		99	1	
01000022	**Low discrimination**						**01000022**
					GRA	SPO	
Atopobium minutum		68.96	0.89		99	1	
Pseudoflavonifractor capillosus		21.05	0.66		1	1	
Clostridium novyi biovar A		7.87	0.77		99	99	
01000023	**Low discrimination**						**01000023**
					GRA	SPO	
Clostridium novyi biovar A		88.62	0.77		99	99	
Atopobium minutum		7.85	0.55		99	1	
Pseudoflavonifractor capillosus		2.40	0.32		1	1	
01000030	**Very good identification**						**01000030**
					SPO		
Clostridium novyi biovar A		96.42	0.90		99		
Atopobium minutum		2.26	0.58		1		
Clostridium cadaveris		0.56	0.37		99		
01000031	**Very good identification**						**01000031**
					SPO		
Clostridium novyi biovar A		99.95	0.90		99		
Atopobium minutum		0.02	0.24		1		
Clostridium cadaveris		0.01	0.03		99		
01000032	**Low discrimination**						**01000032**
					GRA	SPO	
Clostridium novyi biovar A		89.45	0.77		99	99	
Atopobium minutum		7.92	0.55		99	1	
Pseudoflavonifractor capillosus		2.42	0.32		1	1	
01000033	**Very good identification**						**01000033**
					GRA	SPO	
Clostridium novyi biovar A		99.88	0.77		99	99	
Atopobium minutum		0.09	0.21		99	1	
Pseudoflavonifractor capillosus		0.03	-.03		1	1	
01000040	**Low discrimination**						**01000040**
					COC	SPO	
Clostridium botulinum biovar A		46.34	0.86		1	99	
Clostridium novyi biovar A		39.29	0.97		1	99	
Peptostreptococcus anaerobius		5.44	0.66		99	1	
01000041	**Very good identification**						**01000041**
					GRA	SPO	
Clostridium novyi biovar A		98.06	0.97		99	99	
Clostridium botulinum biovar A		1.17	0.52		99	99	
Campylobacter ureolyticus		0.27	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01000042	**Low discrimination**						**01000042**
					SPO		
Clostridium novyi biovar A		78.54	0.84		99		
Atopobium minutum		7.82	0.63		1		
Clostridium botulinum biovar A		5.30	0.52		99		
01000043	**Very good identification**						**01000043**
					COC	SPO	
Clostridium novyi biovar A		99.61	0.84		1	99	
Atopobium minutum		0.10	0.29		1	1	
Anaerococcus prevotii		0.08	0.13		99	1	
01000050	**Very good identification**						**01000050**
					COC	SPO	
Clostridium novyi biovar A		98.45	0.97		1	99	
Clostridium botulinum biovar A		1.17	0.52		1	99	
Peptostreptococcus anaerobius		0.14	0.32		99	1	
01000051	**Excellent identification**						**01000051**
					GRA	SPO	
Clostridium novyi biovar A		99.98	0.97		99	99	
Clostridium botulinum biovar A		0.01	0.18		99	99	
Campylobacter ureolyticus		0.00	-.03		1	1	
01000052	**Very good identification**						**01000052**
					SPO		
Clostridium novyi biovar A		99.74	0.84		99		
Atopobium minutum		0.10	0.29		1		
Clostridium botulinum biovar A		0.07	0.18		99		
01000053	**Very good identification**						**01000053**
Clostridium novyi biovar A		100.0	0.84				
01000060	**Low discrimination**						**01000060**
					SPO		
Clostridium novyi biovar A		87.93	0.87		99		
Clostridium botulinum biovar A		4.19	0.52		99		
Atopobium minutum		3.09	0.58		1		
01000061	**Very good identification**						**01000061**
					SPO		
Clostridium novyi biovar A		99.85	0.87		99		
Clostridium botulinum biovar A		0.05	0.18		99		
Atopobium minutum		0.04	0.24		1		
01000062	**Low discrimination**						**01000062**
					GRA	SPO	
Clostridium novyi biovar A		83.33	0.74		99	99	
Atopobium minutum		11.07	0.55		99	1	
Pseudoflavonifractor capillosus		3.38	0.32		1	1	
01000063	**Very good identification**						**01000063**
					GRA	SPO	
Clostridium novyi biovar A		99.81	0.74		99	99	
Atopobium minutum		0.13	0.21		99	1	
Pseudoflavonifractor capillosus		0.04	-.03		1	1	
01000070	**Very good identification**						**01000070**
					SPO		
Clostridium novyi biovar A		99.87	0.87		99		
Clostridium botulinum biovar A		0.05	0.18		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium minutum		0.04	0.24		1			
01000071	**Very good identification**							**01000071**
Clostridium novyi biovar A		100.0	0.87					
01000072	**Very good identification**							**01000072**
					GRA	SPO		
Clostridium novyi biovar A		99.81	0.74		99	99		
Atopobium minutum		0.13	0.21		99	1		
Pseudoflavonifractor capillosus		0.04	-.03		1	1		
01000073	**Very good identification**							**01000073**
Clostridium novyi biovar A		100.0	0.74					
01000100	**Low discrimination**							**01000100**
					COC	SPO		
Peptostreptococcus anaerobius		51.10	0.66	LAC 1	99	1		
Clostridium botulinum biovar A		14.72	0.61		1	99		
Atopobium minutum		9.73	0.66		1	1		
01000101	**Low discrimination**							**01000101**
					COC	GRA	SPO	
Clostridium novyi biovar A		70.67	0.66	LAC 1	1	99	99	
Campylobacter ureolyticus		12.76	0.32		1	1	1	
Peptostreptococcus anaerobius		6.53	0.32		99	99	1	
01000102	**Low discrimination**							**01000102**
					SPO			
Atopobium minutum		64.27	0.63	LAC 1	1			
Clostridium sporogenes		11.72	0.40		99			
Clostridium novyi biovar A		9.78	0.53		99			
01000103	**Low discrimination**							**01000103**
					COC	SPO		
Clostridium novyi biovar A		85.80	0.53	LAC 1	1	99		
Atopobium minutum		5.70	0.29		1	1		
Anaerococcus prevotii		4.44	0.13		99	1		
01000110	**Low discrimination**							**01000110**
					COC	SPO		
Clostridium novyi biovar A		85.40	0.66	LAC 1	1	99		
Peptostreptococcus anaerobius		7.89	0.32		99	1		
Clostridium botulinum biovar A		2.27	0.27		1	99		
01000111	**Very good identification**							**01000111**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.73	0.66	LAC 1	1	99	99	
Campylobacter ureolyticus		0.18	-.03		1	1	1	
Peptostreptococcus anaerobius		0.09	-.03		99	99	1	
01000112	**Acceptable identification**							**01000112**
					SPO			
Clostridium novyi biovar A		92.27	0.53	LAC 1	99			
Atopobium minutum		6.13	0.29		1			
Clostridium sporogenes		1.12	0.06		99			
01000113	**Good identification**							**01000113**
Clostridium novyi biovar A		100.0	0.53	LAC 1				
01000120	**Low discrimination**							**01000120**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		40.36	0.58	LAC 1	99	1	
Clostridium novyi biovar A		17.40	0.56		99	99	
Prevotella bivia		15.36	0.32		1	1	
01000121	**Acceptable identification**						**01000121**
					GRA	SPO	
Clostridium novyi biovar A		94.53	0.56	LAC 1	99	99	
Atopobium minutum		2.22	0.24		99	1	
Prevotella bivia		0.84	-.03		1	1	
01000122	**Low discrimination**						**01000122**
					GRA	SPO	
Atopobium minutum		68.65	0.55	LAC 1	99	1	
Pseudoflavonifractor capillosus		20.96	0.32		1	1	
Clostridium novyi biovar A		7.83	0.43		99	99	
01000123	**Low discrimination**						**01000123**
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	LAC 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01000130	**Good identification**						**01000130**
					GRA	SPO	
Clostridium novyi biovar A		95.98	0.56	LAC 1	99	99	
Atopobium minutum		2.25	0.24		99	1	
Prevotella bivia		0.86	-.03		1	1	
01000131	**Good identification**						**01000131**
Clostridium novyi biovar A		100.0	0.56	LAC 1			
01000132	**Low discrimination**						**01000132**
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	LAC 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01000133	**Good identification**						**01000133**
Clostridium novyi biovar A		100.0	0.43	LAC 1			
01000140	**Low discrimination**						**01000140**
					COC	SPO	
Clostridium botulinum biovar A		46.27	0.52	LAC 1	1	99	
Clostridium novyi biovar A		39.23	0.63		1	99	
Peptostreptococcus anaerobius		5.43	0.32		99	1	
01000141	**Very good identification**						**01000141**
					GRA	SPO	
Clostridium novyi biovar A		98.42	0.63	LAC 1	99	99	
Clostridium botulinum biovar A		1.17	0.18		99	99	
Campylobacter ureolyticus		0.27	-.03		1	1	
01000142	**Low discrimination**						**01000142**
					SPO		
Clostridium novyi biovar A		79.29	0.50	LAC 1	99		
Atopobium minutum		7.90	0.29		1		
Clostridium botulinum biovar A		5.35	0.18		99		
01000143	**Good identification**						**01000143**
Clostridium novyi biovar A		100.0	0.50	LAC 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01000150	**Very good identification**							**01000150**
					COC	SPO		
Clostridium novyi biovar A		98.69	0.63	LAC 1	1	99		
Clostridium botulinum biovar A		1.18	0.18		1	99		
Peptostreptococcus anaerobius		0.14	-.03		99	1		
01000151	**Very good identification**							**01000151**
Clostridium novyi biovar A		100.0	0.63	LAC 1				
01000152	**Good identification**							**01000152**
Clostridium novyi biovar A		100.0	0.50	LAC 1				
01000153	**Good identification**							**01000153**
Clostridium novyi biovar A		100.0	0.50	LAC 1				
01000160	**Low discrimination**							**01000160**
					SPO			
Clostridium novyi biovar A		87.61	0.53	LAC 1	99			
Clostridium botulinum biovar A		4.18	0.18		99			
Atopobium minutum		3.08	0.24		1			
01000161	**Good identification**							**01000161**
Clostridium novyi biovar A		100.0	0.53	LAC 1				
01000162	**Low discrimination**							**01000162**
					GRA	SPO		
Clostridium novyi biovar A		83.83	0.40	LAC 1	99	99		
Atopobium minutum		11.13	0.21		99	1		
Pseudoflavonifractor capillosus		3.40	-.03		1	1		
01000163	**Good identification**							**01000163**
Clostridium novyi biovar A		100.0	0.40	LAC 1				
01000170	**Good identification**							**01000170**
Clostridium novyi biovar A		100.0	0.53	LAC 1				
01000171	**Good identification**							**01000171**
Clostridium novyi biovar A		100.0	0.53	LAC 1				
01000172	**Good identification**							**01000172**
Clostridium novyi biovar A		100.0	0.40	LAC 1				
01000173	**Good identification**							**01000173**
Clostridium novyi biovar A		100.0	0.40	LAC 1				
01000200	**Low discrimination**							**01000200**
					COC	SPO		
Peptostreptococcus anaerobius		52.15	0.66	RHA 1	99	1		
Clostridium botulinum biovar A		15.02	0.61		1	99		
Atopobium minutum		9.93	0.66		1	1		
01000201	**Low discrimination**							**01000201**
					COC	GRA	SPO	
Clostridium novyi biovar A		70.85	0.66	RHA 1	1	99	99	
Campylobacter ureolyticus		12.79	0.32		1	1	1	
Peptostreptococcus anaerobius		6.54	0.32		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01000202	**Low discrimination**							**01000202**
					SPO			
Atopobium minutum		64.69	0.63	RHA 1	1			
Clostridium sporogenes		11.79	0.40		99			
Clostridium novyi biovar A		9.84	0.53		99			
01000203	**Low discrimination**							**01000203**
					COC	SPO		
Clostridium novyi biovar A		85.80	0.53	RHA 1	1	99		
Atopobium minutum		5.70	0.29		1	1		
Anaerococcus prevotii		4.44	0.13		99	1		
01000210	**Low discrimination**							**01000210**
					COC	SPO		
Clostridium novyi biovar A		85.66	0.66	RHA 1	1	99		
Peptostreptococcus anaerobius		7.91	0.32		99	1		
Clostridium botulinum biovar A		2.28	0.27		1	99		
01000211	**Very good identification**							**01000211**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.73	0.66	RHA 1	1	99	99	
Campylobacter ureolyticus		0.18	-.03		1	1	1	
Peptostreptococcus anaerobius		0.09	-.03		99	99	1	
01000212	**Acceptable identification**							**01000212**
					SPO			
Clostridium novyi biovar A		92.27	0.53	RHA 1	99			
Atopobium minutum		6.13	0.29		1			
Clostridium sporogenes		1.12	0.06		99			
01000213	**Good identification**							**01000213**
Clostridium novyi biovar A		100.0	0.53	RHA 1				
01000220	**Low discrimination**							**01000220**
					SPO			
Atopobium minutum		49.63	0.58	RHA 1	1			
Clostridium novyi biovar A		21.39	0.56		99			
Clostridium cadaveris		12.30	0.37		99			
01000221	**Good identification**							**01000221**
					COC	SPO		
Clostridium novyi biovar A		95.33	0.56	RHA 1	1	99		
Atopobium minutum		2.23	0.24		1	1		
Peptococcus niger		0.83	-.02		99	1		
01000222	**Low discrimination**							**01000222**
					GRA	SPO		
Atopobium minutum		69.11	0.55	RHA 1	99	1		
Pseudoflavonifractor capillosus		21.10	0.32		1	1		
Clostridium novyi biovar A		7.89	0.43		99	99		
01000223	**Low discrimination**							**01000223**
					GRA	SPO		
Clostridium novyi biovar A		89.64	0.43	RHA 1	99	99		
Atopobium minutum		7.94	0.21		99	1		
Pseudoflavonifractor capillosus		2.42	-.03		1	1		
01000230	**Good identification**							**01000230**
					SPO			
Clostridium novyi biovar A		96.81	0.56	RHA 1	99			
Atopobium minutum		2.27	0.24		1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium cadaveris		0.56	0.03		99		
01000231	**Good identification**						**01000231**
Clostridium novyi biovar A		100.0	0.56	RHA 1			
01000232	**Low discrimination**						**01000232**
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	RHA 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01000233	**Good identification**						**01000233**
Clostridium novyi biovar A		100.0	0.43	RHA 1			
01000240	**Low discrimination**						**01000240**
					COC	SPO	
Clostridium botulinum biovar A		46.36	0.52	RHA 1	1	99	
Clostridium novyi biovar A		39.31	0.63		1	99	
Peptostreptococcus anaerobius		5.44	0.32		99	1	
01000241	**Very good identification**						**01000241**
					GRA	SPO	
Clostridium novyi biovar A		98.42	0.63	RHA 1	99	99	
Clostridium botulinum biovar A		1.17	0.18		99	99	
Campylobacter ureolyticus		0.27	-.03		1	1	
01000242	**Low discrimination**						**01000242**
					SPO		
Clostridium novyi biovar A		79.29	0.50	RHA 1	99		
Atopobium minutum		7.90	0.29		1		
Clostridium botulinum biovar A		5.35	0.18		99		
01000243	**Good identification**						**01000243**
Clostridium novyi biovar A		100.0	0.50	RHA 1			
01000250	**Very good identification**						**01000250**
					COC	SPO	
Clostridium novyi biovar A		98.69	0.63	RHA 1	1	99	
Clostridium botulinum biovar A		1.18	0.18		1	99	
Peptostreptococcus anaerobius		0.14	-.03		99	1	
01000251	**Very good identification**						**01000251**
Clostridium novyi biovar A		100.0	0.63	RHA 1			
01000252	**Good identification**						**01000252**
Clostridium novyi biovar A		100.0	0.50	RHA 1			
01000253	**Good identification**						**01000253**
Clostridium novyi biovar A		100.0	0.50	RHA 1			
01000260	**Low discrimination**						**01000260**
					SPO		
Clostridium novyi biovar A		88.64	0.53	RHA 1	99		
Clostridium botulinum biovar A		4.22	0.18		99		
Atopobium minutum		3.12	0.24		1		
01000261	**Good identification**						**01000261**
Clostridium novyi biovar A		100.0	0.53	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01000262	**Low discrimination**							**01000262**
					GRA	SPO		
Clostridium novyi biovar A		83.83	0.40	RHA 1	99	99		
Atopobium minutum		11.13	0.21		99	1		
Pseudoflavonifractor capillosus		3.40	-.03		1	1		
01000263	**Good identification**							**01000263**
Clostridium novyi biovar A		100.0	0.40	RHA 1				
01000270	**Good identification**							**01000270**
Clostridium novyi biovar A		100.0	0.53	RHA 1				
01000271	**Good identification**							**01000271**
Clostridium novyi biovar A		100.0	0.53	RHA 1				
01000272	**Good identification**							**01000272**
Clostridium novyi biovar A		100.0	0.40	RHA 1				
01000273	**Good identification**							**01000273**
Clostridium novyi biovar A		100.0	0.40	RHA 1				
01000400	**Low discrimination**							**01000400**
					COC	SPO		
Peptostreptococcus anaerobius		51.95	0.66	ARA 1	99	1		
Clostridium botulinum biovar A		14.96	0.61		1	99		
Atopobium minutum		9.89	0.66		1	1		
01000401	**Low discrimination**							**01000401**
					COC	GRA	SPO	
Clostridium novyi biovar A		70.82	0.66	ARA 1	1	99	99	
Campylobacter ureolyticus		12.78	0.32		1	1	1	
Peptostreptococcus anaerobius		6.54	0.32		99	99	1	
01000402	**Low discrimination**							**01000402**
					SPO			
Atopobium minutum		64.65	0.63	ARA 1	1			
Clostridium sporogenes		11.79	0.40		99			
Clostridium novyi biovar A		9.84	0.53		99			
01000403	**Low discrimination**							**01000403**
					COC	SPO		
Clostridium novyi biovar A		85.80	0.53	ARA 1	1	99		
Atopobium minutum		5.70	0.29		1	1		
Anaerococcus prevotii		4.44	0.13		99	1		
01000410	**Low discrimination**							**01000410**
					COC	SPO		
Clostridium novyi biovar A		85.62	0.66	ARA 1	1	99		
Peptostreptococcus anaerobius		7.91	0.32		99	1		
Clostridium botulinum biovar A		2.28	0.27		1	99		
01000411	**Very good identification**							**01000411**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.73	0.66	ARA 1	1	99	99	
Campylobacter ureolyticus		0.18	-.03		1	1	1	
Peptostreptococcus anaerobius		0.09	-.03		99	99	1	
01000412	**Acceptable identification**							**01000412**
					SPO			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		92.27	0.53	ARA 1	99		
Atopobium minutum		6.13	0.29		1		
Clostridium sporogenes		1.12	0.06		99		
01000413	**Good identification**						**01000413**
Clostridium novyi biovar A		100.0	0.53	ARA 1			
01000420	**Low discrimination**						**01000420**
					SPO		
Atopobium minutum		49.27	0.58	ARA 1	1		
Clostridium novyi biovar A		21.24	0.56		99		
Clostridium cadaveris		12.21	0.37		99		
01000421	**Good identification**						**01000421**
					COC	SPO	
Clostridium novyi biovar A		95.33	0.56	ARA 1	1	99	
Atopobium minutum		2.23	0.24		1	1	
Peptococcus niger		0.83	-.02		99	1	
01000422	**Low discrimination**						**01000422**
					GRA	SPO	
Atopobium minutum		69.11	0.55	ARA 1	99	1	
Pseudoflavonifractor capillosus		21.10	0.32		1	1	
Clostridium novyi biovar A		7.89	0.43		99	99	
01000423	**Low discrimination**						**01000423**
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	ARA 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01000430	**Good identification**						**01000430**
					SPO		
Clostridium novyi biovar A		96.81	0.56	ARA 1	99		
Atopobium minutum		2.27	0.24		1		
Clostridium cadaveris		0.56	0.03		99		
01000431	**Good identification**						**01000431**
Clostridium novyi biovar A		100.0	0.56	ARA 1			
01000432	**Low discrimination**						**01000432**
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	ARA 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01000433	**Good identification**						**01000433**
Clostridium novyi biovar A		100.0	0.43	ARA 1			
01000440	**Low discrimination**						**01000440**
					COC	SPO	
Clostridium botulinum biovar A		46.35	0.52	ARA 1	1	99	
Clostridium novyi biovar A		39.30	0.63		1	99	
Peptostreptococcus anaerobius		5.44	0.32		99	1	
01000441	**Very good identification**						**01000441**
					GRA	SPO	
Clostridium novyi biovar A		98.42	0.63	ARA 1	99	99	
Clostridium botulinum biovar A		1.17	0.18		99	99	
Campylobacter ureolyticus		0.27	-.03		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
01000442	**Low discrimination**						**01000442**
					SPO		
Clostridium novyi biovar A		79.29	0.50	ARA 1	99		
Atopobium minutum		7.90	0.29		1		
Clostridium botulinum biovar A		5.35	0.18		99		
01000443	**Good identification**						**01000443**
Clostridium novyi biovar A		100.0	0.50	ARA 1			
01000450	**Very good identification**						**01000450**
					COC	SPO	
Clostridium novyi biovar A		98.69	0.63	ARA 1	1	99	
Clostridium botulinum biovar A		1.18	0.18		1	99	
Peptostreptococcus anaerobius		0.14	-.03		99	1	
01000451	**Very good identification**						**01000451**
Clostridium novyi biovar A		100.0	0.63	ARA 1			
01000452	**Good identification**						**01000452**
Clostridium novyi biovar A		100.0	0.50	ARA 1			
01000453	**Good identification**						**01000453**
Clostridium novyi biovar A		100.0	0.50	ARA 1			
01000460	**Low discrimination**						**01000460**
					SPO		
Clostridium novyi biovar A		88.64	0.53	ARA 1	99		
Clostridium botulinum biovar A		4.22	0.18		99		
Atopobium minutum		3.12	0.24		1		
01000461	**Good identification**						**01000461**
Clostridium novyi biovar A		100.0	0.53	ARA 1			
01000462	**Low discrimination**						**01000462**
					GRA	SPO	
Clostridium novyi biovar A		83.83	0.40	ARA 1	99	99	
Atopobium minutum		11.13	0.21		99	1	
Pseudoflavonifractor capillosus		3.40	-.03		1	1	
01000463	**Good identification**						**01000463**
Clostridium novyi biovar A		100.0	0.40	ARA 1			
01000470	**Good identification**						**01000470**
Clostridium novyi biovar A		100.0	0.53	ARA 1			
01000471	**Good identification**						**01000471**
Clostridium novyi biovar A		100.0	0.53	ARA 1			
01000472	**Good identification**						**01000472**
Clostridium novyi biovar A		100.0	0.40	ARA 1			
01000473	**Good identification**						**01000473**
Clostridium novyi biovar A		100.0	0.40	ARA 1			
01001000	**Low discrimination**						**01001000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Atopobium minutum		67.30	0.92		1	1	
Clostridium novyi biovar A		12.89	0.84		1	99	
Peptostreptococcus anaerobius		10.71	0.66		99	1	
01001001	**Good identification**						**01001001**
					SPO		
Clostridium novyi biovar A		90.37	0.84	GAL 10	99		
Atopobium minutum		4.77	0.58		1		
Clostridium sordellii		1.54	0.37		99		
01001002	**Good identification**						**01001002**
					SPO		
Atopobium minutum		93.99	0.89		1		
Clostridium novyi biovar A		4.77	0.71		99		
Clostridium sporogenes		0.52	0.40		99		
01001003	**Low discrimination**						**01001003**
					COC	SPO	
Clostridium novyi biovar A		82.70	0.71	GAL 10	1	99	
Atopobium minutum		16.47	0.55		1	1	
Anaerococcus prevotii		0.39	0.13		99	1	
01001010	**Good identification**						**01001010**
					COC	SPO	
Clostridium novyi biovar A		93.61	0.84	GAL 10	1	99	
Atopobium minutum		4.94	0.58		1	1	
Peptostreptococcus anaerobius		0.79	0.32		99	1	
01001011	**Very good identification**						**01001011**
					SPO		
Clostridium novyi biovar A		99.91	0.84	GAL 10	99		
Atopobium minutum		0.05	0.24		1		
Clostridium sordellii		0.02	0.03		99		
01001012	**Low discrimination**						**01001012**
					SPO		
Clostridium novyi biovar A		83.28	0.71	GAL 10	99		
Atopobium minutum		16.59	0.55		1		
Clostridium sporogenes		0.09	0.06		99		
01001013	**Very good identification**						**01001013**
					SPO		
Clostridium novyi biovar A		99.80	0.71	GAL 10	99		
Atopobium minutum		0.20	0.21		1		
01001020	**Low discrimination**						**01001020**
					SPO		
Atopobium minutum		85.71	0.84		1		
Clostridium novyi biovar A		12.32	0.73		99		
Clostridium cadaveris		0.64	0.37		99		
01001021	**Good identification**						**01001021**
					COC	SPO	
Clostridium novyi biovar A		93.17	0.73	GAL 10	1	99	
Atopobium minutum		6.55	0.50		1	1	
Peptococcus niger		0.07	-.02		99	1	
01001022	**Low discrimination**						**01001022**
					GRA	SPO	
Atopobium minutum		87.64	0.81		99	1	
Pseudoflavonifractor capillosus		8.92	0.50		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		3.33	0.61		99	99	
01001023	**Low discrimination**						**01001023**
					GRA	SPO	
Clostridium novyi biovar A		77.36	0.61	GAL 10	99	99	
Atopobium minutum		20.55	0.47		99	1	
Pseudoflavonifractor capillosus		2.09	0.15		1	1	
01001030	**Good identification**						**01001030**
					SPO		
Clostridium novyi biovar A		93.36	0.73	GAL 10	99		
Atopobium minutum		6.56	0.50		1		
Clostridium cadaveris		0.05	0.03		99		
01001031	**Very good identification**						**01001031**
					SPO		
Clostridium novyi biovar A		99.93	0.73	GAL 10	99		
Atopobium minutum		0.07	0.15		1		
01001032	**Low discrimination**						**01001032**
					GRA	SPO	
Clostridium novyi biovar A		77.36	0.61	GAL 10	99	99	
Atopobium minutum		20.55	0.47		99	1	
Pseudoflavonifractor capillosus		2.09	0.15		1	1	
01001033	**Very good identification**						**01001033**
					SPO		
Clostridium novyi biovar A		99.73	0.61	GAL 10	99		
Atopobium minutum		0.27	0.12		1		
01001040	**Low discrimination**						**01001040**
					SPO		
Clostridium novyi biovar A		79.27	0.81	GAL 10	99		
Clostridium botulinum biovar A		8.50	0.52		99		
Atopobium minutum		6.27	0.58		1		
01001041	**Very good identification**						**01001041**
Clostridium novyi biovar A		99.62	0.81	GAL 10			
Clostridium botulinum biovar A		0.11	0.18				
Clostridium sordellii		0.11	0.13				
01001042	**Low discrimination**						**01001042**
					SPO		
Clostridium novyi biovar A		75.98	0.68	GAL 10	99		
Atopobium minutum		22.70	0.55		1		
Clostridium botulinum biovar A		0.47	0.18		99		
01001043	**Very good identification**						**01001043**
					SPO		
Clostridium novyi biovar A		99.70	0.68	GAL 10	99		
Atopobium minutum		0.30	0.21		1		
01001050	**Very good identification**						**01001050**
					SPO		
Clostridium novyi biovar A		99.75	0.81	GAL 10	99		
Clostridium botulinum biovar A		0.11	0.18		99		
Atopobium minutum		0.08	0.24		1		
01001051	**Very good identification**						**01001051**
Clostridium novyi biovar A		100.0	0.81	GAL 10			
01001052	**Very good identification**						**01001052**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium novyi biovar A		99.70	0.68	GAL 10	99		
Atopobium minutum		0.30	0.21		1		
01001053	**Very good identification**						**01001053**
Clostridium novyi biovar A		100.0	0.68	GAL 10			
01001060	**Low discrimination**						**01001060**
					SPO		
Clostridium novyi biovar A		89.42	0.70	GAL 10	99		
Atopobium minutum		9.43	0.50		1		
Clostridium botulinum biovar A		0.39	0.18		99		
01001061	**Very good identification**						**01001061**
					SPO		
Clostridium novyi biovar A		99.89	0.70	GAL 10	99		
Atopobium minutum		0.11	0.15		1		
01001062	**Low discrimination**						**01001062**
					GRA	SPO	
Clostridium novyi biovar A		69.41	0.58	GAL 10	99	99	
Atopobium minutum		27.65	0.47		99	1	
Pseudoflavonifractor capillosus		2.81	0.15		1	1	
01001063	**Very good identification**						**01001063**
					SPO		
Clostridium novyi biovar A		99.60	0.58	GAL 10	99		
Atopobium minutum		0.40	0.12		1		
01001070	**Very good identification**						**01001070**
					SPO		
Clostridium novyi biovar A		99.89	0.70	GAL 10	99		
Atopobium minutum		0.11	0.15		1		
01001071	**Very good identification**						**01001071**
Clostridium novyi biovar A		100.0	0.70	GAL 10			
01001072	**Very good identification**						**01001072**
					SPO		
Clostridium novyi biovar A		99.60	0.58	GAL 10	99		
Atopobium minutum		0.40	0.12		1		
01001073	**Very good identification**						**01001073**
Clostridium novyi biovar A		100.0	0.58	GAL 10			
01001100	**Low discrimination**						**01001100**
					SPO		
Atopobium minutum		38.58	0.58	LAC 1	1		
Actinomyces meyeri		34.76	0.53		1		
Clostridium novyi biovar A		7.39	0.50		99		
01001101	**Low discrimination**						**01001101**
					SPO		
Clostridium novyi biovar A		86.39	0.50	GAL 10	99		
Atopobium minutum		4.56	0.24	LAC 1	1		
Actinomyces meyeri		4.10	0.19		1		
01001102	**Low discrimination**						**01001102**
					SPO		
Atopobium minutum		84.53	0.55	LAC 1	1		
Lactobacillus fermentum		4.55	0.40		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		4.29	0.37		99	
01001103	**Low discrimination**					**01001103**
					SPO	
Clostridium novyi biovar A		83.39	0.37	GAL 10	99	
Atopobium minutum		16.61	0.21	LAC 1	1	
01001110	**Low discrimination**					**01001110**
					SPO	
Clostridium novyi biovar A		89.56	0.50	GAL 10	99	
Atopobium minutum		4.72	0.24	LAC 1	1	
Actinomyces meyeri		4.25	0.19		1	
01001111	**Good identification**					**01001111**
Clostridium novyi biovar A		100.0	0.50	GAL 10		
01001112	**Low discrimination**					**01001112**
					SPO	
Clostridium novyi biovar A		83.39	0.37	GAL 10	99	
Atopobium minutum		16.61	0.21	LAC 1	1	
01001113	**Good identification**					**01001113**
Clostridium novyi biovar A		100.0	0.37	GAL 10		
01001120	**Low discrimination**					**01001120**
					GRA	
Atopobium minutum		55.19	0.50	LAC 1	99	
Actinomyces meyeri		20.34	0.38		99	
Prevotella bivia		11.12	0.19		1	
01001121	**Acceptable identification**					**01001121**
					SPO	
Clostridium novyi biovar A		91.22	0.39	GAL 10	99	
Atopobium minutum		6.41	0.15	LAC 1	1	
Actinomyces meyeri		2.36	0.04		1	
01001122	**Low discrimination**					**01001122**
					GRA	
Atopobium minutum		82.04	0.47	LAC 1	99	
Pseudoflavonifractor capillosus		8.35	0.15		1	
Lactobacillus fermentum		4.41	0.32		99	
01001130	**Acceptable identification**					**01001130**
					SPO	
Clostridium novyi biovar A		91.22	0.39	GAL 10	99	
Atopobium minutum		6.41	0.15	LAC 1	1	
Actinomyces meyeri		2.36	0.04		1	
01001131	**Good identification**					**01001131**
Clostridium novyi biovar A		100.0	0.39	GAL 10		
01001140	**Low discrimination**					**01001140**
					SPO	
Clostridium novyi biovar A		75.31	0.47	GAL 10	99	
Clostridium botulinum biovar A		8.08	0.18	LAC 1	99	
Atopobium minutum		5.96	0.24		1	
01001141	**Good identification**					**01001141**
Clostridium novyi biovar A		100.0	0.47	GAL 10		
01001150	**Good identification**					**01001150**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01001151	**Good identification**						**01001151**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01001160	**Low discrimination**						**01001160**
					SPO		
Clostridium novyi biovar A		87.39	0.36	GAL 10	99		
Atopobium minutum		9.22	0.15	LAC 1	1		
Actinomyces meyeri		3.40	0.04		1		
01001161	**Good identification**						**01001161**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01001170	**Good identification**						**01001170**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01001171	**Good identification**						**01001171**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01001200	**Low discrimination**						**01001200**
					COC	SPO	
Atopobium minutum		67.38	0.58	RHA 1	1	1	
Clostridium novyi biovar A		12.91	0.50		1	99	
Peptostreptococcus anaerobius		10.73	0.32		99	1	
01001201	**Acceptable identification**						**01001201**
					SPO		
Clostridium novyi biovar A		91.36	0.50	GAL 10	99		
Atopobium minutum		4.82	0.24	RHA 1	1		
Clostridium sordellii		1.56	0.03		99		
01001202	**Acceptable identification**						**01001202**
					SPO		
Atopobium minutum		94.46	0.55	RHA 1	1		
Clostridium novyi biovar A		4.79	0.37		99		
Clostridium sporogenes		0.52	0.06		99		
01001203	**Low discrimination**						**01001203**
					SPO		
Clostridium novyi biovar A		83.39	0.37	GAL 10	99		
Atopobium minutum		16.61	0.21	RHA 1	1		
01001210	**Acceptable identification**						**01001210**
					COC	SPO	
Clostridium novyi biovar A		94.24	0.50	GAL 10	1	99	
Atopobium minutum		4.97	0.24	RHA 1	1	1	
Peptostreptococcus anaerobius		0.79	-.03		99	1	
01001211	**Good identification**						**01001211**
Clostridium novyi biovar A		100.0	0.50	GAL 10			
01001212	**Low discrimination**						**01001212**
					SPO		
Clostridium novyi biovar A		83.39	0.37	GAL 10	99		
Atopobium minutum		16.61	0.21	RHA 1	1		
01001213	**Good identification**						**01001213**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01001220	**Low discrimination**						**01001220**
					SPO		
Atopobium minutum		86.50	0.50	RHA 1	1		
Clostridium novyi biovar A		12.43	0.39		99		
Clostridium cadaveris		0.65	0.03		99		
01001221	**Acceptable identification**						**01001221**
					SPO		
Clostridium novyi biovar A		93.43	0.39	GAL 10	99		
Atopobium minutum		6.57	0.15	RHA 1	1		
01001222	**Low discrimination**						**01001222**
					GRA	SPO	
Atopobium minutum		87.74	0.47	RHA 1	99	1	
Pseudoflavonifractor capillosus		8.93	0.15		1	1	
Clostridium novyi biovar A		3.34	0.26		99	99	
01001230	**Acceptable identification**						**01001230**
					SPO		
Clostridium novyi biovar A		93.43	0.39	GAL 10	99		
Atopobium minutum		6.57	0.15	RHA 1	1		
01001231	**Good identification**						**01001231**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01001240	**Low discrimination**						**01001240**
					SPO		
Clostridium novyi biovar A		80.34	0.47	GAL 10	99		
Clostridium botulinum biovar A		8.62	0.18	RHA 1	99		
Atopobium minutum		6.35	0.24		1		
01001241	**Good identification**						**01001241**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01001250	**Good identification**						**01001250**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01001251	**Good identification**						**01001251**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01001260	**Acceptable identification**						**01001260**
					SPO		
Clostridium novyi biovar A		90.46	0.36	GAL 10	99		
Atopobium minutum		9.54	0.15	RHA 1	1		
01001261	**Good identification**						**01001261**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01001270	**Good identification**						**01001270**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01001271	**Good identification**						**01001271**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01001400	**Low discrimination**						**01001400**
					COC	SPO	
Atopobium minutum		62.30	0.58	ARA 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		11.93	0.50		1	99	
Peptostreptococcus anaerobius		9.92	0.32		99	1	
01001401	**Acceptable identification**						**01001401**
					SPO		
Clostridium novyi biovar A		91.36	0.50	GAL 10	99		
Atopobium minutum		4.82	0.24	ARA 1	1		
Clostridium sordellii		1.56	0.03		99		
01001402	**Acceptable identification**						**01001402**
					SPO		
Atopobium minutum		91.39	0.55	ARA 1	1		
Clostridium novyi biovar A		4.63	0.37		99		
Lactobacillus fermentum		3.01	0.36		1		
01001403	**Low discrimination**						**01001403**
					SPO		
Clostridium novyi biovar A		83.39	0.37	GAL 10	99		
Atopobium minutum		16.61	0.21	ARA 1	1		
01001410	**Acceptable identification**						**01001410**
					COC	SPO	
Clostridium novyi biovar A		94.24	0.50	GAL 10	1	99	
Atopobium minutum		4.97	0.24	ARA 1	1	1	
Peptostreptococcus anaerobius		0.79	-.03		99	1	
01001411	**Good identification**						**01001411**
Clostridium novyi biovar A		100.0	0.50	GAL 10			
01001412	**Low discrimination**						**01001412**
					SPO		
Clostridium novyi biovar A		83.39	0.37	GAL 10	99		
Atopobium minutum		16.61	0.21	ARA 1	1		
01001413	**Good identification**						**01001413**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01001420	**Low discrimination**						**01001420**
					SPO		
Atopobium minutum		82.13	0.50	ARA 1	1		
Clostridium novyi biovar A		11.80	0.39		99		
Actinomyces meyeri		1.89	0.18		1		
01001421	**Acceptable identification**						**01001421**
					SPO		
Clostridium novyi biovar A		93.43	0.39	GAL 10	99		
Atopobium minutum		6.57	0.15	ARA 1	1		
01001422	**Low discrimination**						**01001422**
					GRA	SPO	
Atopobium minutum		85.27	0.47	ARA 1	99	1	
Pseudoflavonifractor capillosus		8.68	0.15		1	1	
Clostridium novyi biovar A		3.24	0.26		99	99	
01001430	**Acceptable identification**						**01001430**
					SPO		
Clostridium novyi biovar A		93.43	0.39	GAL 10	99		
Atopobium minutum		6.57	0.15	ARA 1	1		
01001431	**Good identification**						**01001431**
Clostridium novyi biovar A		100.0	0.39	GAL 10			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01001440	**Low discrimination**							**01001440**
					SPO			
Clostridium novyi biovar A		80.34	0.47	GAL 10	99			
Clostridium botulinum biovar A		8.62	0.18	ARA 1	99			
Atopobium minutum		6.35	0.24		1			
01001441	**Good identification**							**01001441**
Clostridium novyi biovar A		100.0	0.47	GAL 10				
01001450	**Good identification**							**01001450**
Clostridium novyi biovar A		100.0	0.47	GAL 10				
01001451	**Good identification**							**01001451**
Clostridium novyi biovar A		100.0	0.47	GAL 10				
01001460	**Acceptable identification**							**01001460**
					SPO			
Clostridium novyi biovar A		90.46	0.36	GAL 10	99			
Atopobium minutum		9.54	0.15	ARA 1	1			
01001461	**Good identification**							**01001461**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01001470	**Good identification**							**01001470**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01001471	**Good identification**							**01001471**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01001500	**Low discrimination**							**01001500**
					COC			
Actinomyces meyeri		74.78	0.43	SUC 93	1			
Lactobacillus fermentum		11.90	0.40	FRU 95	1			
Blautia hansenii		8.79	0.19		99			
01001502	**Low discrimination**							**01001502**
Lactobacillus fermentum		68.29	0.40	SUC 87				
Actinomyces meyeri		13.27	0.17	FRU 99				
Atopobium minutum		12.83	0.21					
01002000	**Low discrimination**							**01002000**
					COC	GRA		
Peptostreptococcus anaerobius		26.76	0.66	MAN 1	99	99		
Megasphaera elsdenii		18.62	0.72		99	1		
Pseudoramibacter alactolyticus		18.28	0.61		1	99		
01002001	**Low discrimination**							**01002001**
					COC	GRA	SPO	
Clostridium novyi biovar A		63.31	0.66	MAN 1	1	99	99	
Campylobacter ureolyticus		11.43	0.32		1	1	1	
Peptostreptococcus anaerobius		5.85	0.32		99	99	1	
01002002	**Low discrimination**							**01002002**
					COC	GRA		
Atopobium minutum		36.07	0.63	MAN 1	1	99		
Megasphaera elsdenii		21.97	0.56		99	1		
Pseudoramibacter alactolyticus		21.57	0.45		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01002003	**Low discrimination**						**01002003**
					COC	SPO	
Clostridium novyi biovar A		80.28	0.53	MAN 1	1	99	
Atopobium minutum		5.33	0.29		1	1	
Anaerococcus prevotii		4.15	0.13		99	1	
01002010	**Low discrimination**						**01002010**
					COC	SPO	
Clostridium difficile		47.61	0.53	FRU 99	1	99	
Clostridium novyi biovar A		39.34	0.66		1	99	
Peptostreptococcus anaerobius		3.63	0.32		99	1	
01002011	**Very good identification**						**01002011**
					GRA	SPO	
Clostridium novyi biovar A		98.40	0.66	MAN 1	99	99	
Clostridium difficile		1.20	0.19		99	99	
Campylobacter ureolyticus		0.18	-.03		1	1	
01002012	**Low discrimination**						**01002012**
					SPO		
Clostridium novyi biovar A		81.09	0.53	MAN 1	99		
Clostridium difficile		5.62	0.19		99		
Atopobium minutum		5.38	0.29		1		
01002013	**Good identification**						**01002013**
Clostridium novyi biovar A		100.0	0.53	MAN 1			
01002020	**Low discrimination**						**01002020**
					SPO		
Pseudoramibacter alactolyticus		46.52	0.48	FRU 99	1		
Atopobium minutum		24.49	0.58		1		
Clostridium novyi biovar A		10.56	0.56		99		
01002021	**Acceptable identification**						**01002021**
					SPO		
Clostridium novyi biovar A		91.23	0.56	MAN 1	99		
Pseudoramibacter alactolyticus		4.06	0.14		1		
Atopobium minutum		2.14	0.24		1		
01002022	**Low discrimination**						**01002022**
					GRA		
Atopobium minutum		56.30	0.55	MAN 1	99		
Pseudoramibacter alactolyticus		17.82	0.32		99		
Pseudoflavonifractor capillosus		17.19	0.32		1		
01002023	**Low discrimination**						**01002023**
					SPO		
Clostridium novyi biovar A		87.44	0.43	MAN 1	99		
Atopobium minutum		7.74	0.21		1		
Pseudoramibacter alactolyticus		2.45	-.02		1		
01002030	**Low discrimination**						**01002030**
					SPO		
Clostridium novyi biovar A		88.58	0.56	MAN 1	99		
Clostridium difficile		4.33	0.19		99		
Pseudoramibacter alactolyticus		3.94	0.14		1		
01002031	**Good identification**						**01002031**
Clostridium novyi biovar A		100.0	0.56	MAN 1			
01002032	**Low discrimination**						**01002032**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Clostridium novyi biovar A		87.44	0.43	MAN 1	99			
Atopobium minutum		7.74	0.21		1			
Pseudoramibacter alactolyticus		2.45	-.02		1			
01002033	**Good identification**							**01002033**
Clostridium novyi biovar A		100.0	0.43	MAN 1				
01002040	**Low discrimination**							**01002040**
					COC	SPO	GRA	
Megasphaera elsdenii		56.70	0.69	FRU 95	99	1	1	
Pseudoramibacter alactolyticus		20.87	0.51		1	1	99	
Clostridium botulinum biovar A		10.52	0.52		1	99	99	
01002041	**Good identification**							**01002041**
					COC	SPO	GRA	
Clostridium novyi biovar A		90.59	0.63	MAN 1	1	99	99	
Megasphaera elsdenii		5.82	0.35		99	1	1	
Pseudoramibacter alactolyticus		2.14	0.17		1	1	99	
01002042	**Low discrimination**							**01002042**
					COC	SPO	GRA	
Megasphaera elsdenii		60.02	0.53	FRU 95	99	1	1	
Pseudoramibacter alactolyticus		22.10	0.35	BGL 10	1	1	99	
Clostridium novyi biovar A		14.99	0.50		1	99	99	
01002043	**Acceptable identification**							**01002043**
					COC	SPO	GRA	
Clostridium novyi biovar A		94.76	0.50	MAN 1	1	99	99	
Megasphaera elsdenii		3.83	0.19		99	1	1	
Pseudoramibacter alactolyticus		1.41	0.00		1	1	99	
01002050	**Low discrimination**							**01002050**
					COC	GRA		
Clostridium novyi biovar A		83.56	0.63	MAN 1	1	99		
Clostridium difficile		7.98	0.31		1	99		
Megasphaera elsdenii		5.37	0.35		99	1		
01002051	**Very good identification**							**01002051**
Clostridium novyi biovar A		99.90	0.63	MAN 1				
Clostridium difficile		0.10	-.03					
01002052	**Acceptable identification**							**01002052**
					COC	SPO	GRA	
Clostridium novyi biovar A		94.27	0.50	MAN 1	1	99	99	
Megasphaera elsdenii		3.81	0.19		99	1	1	
Pseudoramibacter alactolyticus		1.40	0.00		1	1	99	
01002053	**Good identification**							**01002053**
Clostridium novyi biovar A		100.0	0.50	MAN 1				
01002060	**Low discrimination**							**01002060**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		54.67	0.38	FRU 99	1	1	99	
Clostridium novyi biovar A		33.08	0.53		1	99	99	
Megasphaera elsdenii		8.50	0.35		99	1	1	
01002061	**Good identification**							**01002061**
					SPO			
Clostridium novyi biovar A		98.36	0.53	MAN 1	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		1.64	0.04		1			
01002063	**Good identification**							**01002063**
Clostridium novyi biovar A		100.0	0.40	MAN 1				
01002070	**Good identification**							**01002070**
					SPO			
Clostridium novyi biovar A		97.99	0.53	MAN 1	99			
Pseudoramibacter alactolyticus		1.64	0.04		1			
Clostridium difficile		0.38	-.03		99			
01002071	**Good identification**							**01002071**
Clostridium novyi biovar A		100.0	0.53	MAN 1				
01002072	**Good identification**							**01002072**
Clostridium novyi biovar A		100.0	0.40	MAN 1				
01002073	**Good identification**							**01002073**
Clostridium novyi biovar A		100.0	0.40	MAN 1				
01003000	**Low discrimination**							**01003000**
					COC	SPO		
Atopobium minutum		56.37	0.58	MAN 1	1	1		
Clostridium novyi biovar A		10.80	0.50		1	99		
Peptostreptococcus anaerobius		8.97	0.32		99	1		
01003001	**Acceptable identification**							**01003001**
					SPO			
Clostridium novyi biovar A		90.41	0.50	GAL 10	99			
Atopobium minutum		4.77	0.24	MAN 1	1			
Clostridium sordellii		1.54	0.03		99			
01003002	**Acceptable identification**							**01003002**
					COC	SPO	GRA	
Atopobium minutum		91.31	0.55	MAN 1	1	1	99	
Clostridium novyi biovar A		4.63	0.37		1	99	99	
Megasphaera elsdenii		1.69	0.22		99	1	1	
01003003	**Low discrimination**							**01003003**
					SPO			
Clostridium novyi biovar A		83.39	0.37	GAL 10	99			
Atopobium minutum		16.61	0.21	MAN 1	1			
01003010	**Low discrimination**							**01003010**
					SPO			
Clostridium novyi biovar A		84.55	0.50	GAL 10	99			
Clostridium difficile		9.30	0.19	MAN 1	99			
Atopobium minutum		4.46	0.24		1			
01003011	**Good identification**							**01003011**
Clostridium novyi biovar A		100.0	0.50	GAL 10				
01003012	**Low discrimination**							**01003012**
					SPO			
Clostridium novyi biovar A		83.39	0.37	GAL 10	99			
Atopobium minutum		16.61	0.21	MAN 1	1			
01003013	**Good identification**							**01003013**
Clostridium novyi biovar A		100.0	0.37	GAL 10				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
01003020	**Low discrimination**							**01003020**
					SPO	CAT		
Atopobium minutum		78.08	0.50	MAN 1	1	1		
Clostridium novyi biovar A		11.22	0.39		99	1		
Propionibacterium acnes		4.98	0.22		1	95		
01003021	**Acceptable identification**							**01003021**
					SPO			
Clostridium novyi biovar A		93.43	0.39	GAL 10	99			
Atopobium minutum		6.57	0.15	MAN 1	1			
01003022	**Low discrimination**							**01003022**
					GRA	SPO		
Atopobium minutum		87.00	0.47	MAN 1	99	1		
Pseudoflavonifractor capillosus		8.85	0.15		1	1		
Clostridium novyi biovar A		3.31	0.26		99	99		
01003030	**Acceptable identification**							**01003030**
					SPO			
Clostridium novyi biovar A		93.43	0.39	GAL 10	99			
Atopobium minutum		6.57	0.15	MAN 1	1			
01003031	**Good identification**							**01003031**
Clostridium novyi biovar A		100.0	0.39	GAL 10				
01003040	**Low discrimination**							**01003040**
					COC	SPO	GRA	
Clostridium novyi biovar A		49.13	0.47	GAL 10	1	99	99	
Megasphaera elsdenii		28.40	0.35	MAN 1	99	1	1	
Pseudoramibacter alactolyticus		10.46	0.17		1	1	99	
01003041	**Good identification**							**01003041**
Clostridium novyi biovar A		100.0	0.47	GAL 10				
01003050	**Good identification**							**01003050**
Clostridium novyi biovar A		99.14	0.47	GAL 10				
Clostridium difficile		0.86	-.03	MAN 1				
01003051	**Good identification**							**01003051**
Clostridium novyi biovar A		100.0	0.47	GAL 10				
01003060	**Low discrimination**							**01003060**
					SPO	CAT		
Clostridium novyi biovar A		71.34	0.36	GAL 10	99	1		
Pseudoramibacter alactolyticus		10.72	0.04	MAN 1	1	1		
Propionibacterium acnes		10.42	0.11		1	95		
01003061	**Good identification**							**01003061**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01003070	**Good identification**							**01003070**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01003071	**Good identification**							**01003071**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01004000	**Low discrimination**							**01004000**
					COC	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptostreptococcus anaerobius		43.91	0.66	XYL 1	99	1		
Clostridium glycolicum		14.51	0.61		1	99		
Clostridium botulinum biovar A		12.65	0.61		1	99		
01004001	**Low discrimination**							**01004001**
					COC	GRA	SPO	
Clostridium novyi biovar A		69.21	0.66	XYL 1	1	99	99	
Campylobacter ureolyticus		12.49	0.32		1	1	1	
Peptostreptococcus anaerobius		6.39	0.32		99	99	1	
01004002	**Low discrimination**							**01004002**
					SPO			
Atopobium minutum		63.38	0.63	XYL 1	1			
Clostridium sporogenes		11.56	0.40		99			
Clostridium novyi biovar A		9.64	0.53		99			
01004003	**Low discrimination**							**01004003**
					COC	SPO		
Clostridium novyi biovar A		85.80	0.53	XYL 1	1	99		
Atopobium minutum		5.70	0.29		1	1		
Anaerococcus prevotii		4.44	0.13		99	1		
01004010	**Low discrimination**							**01004010**
					COC	SPO		
Clostridium novyi biovar A		82.88	0.66	XYL 1	1	99		
Peptostreptococcus anaerobius		7.65	0.32		99	1		
Clostridium glycolicum		2.53	0.27		1	99		
01004011	**Very good identification**							**01004011**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.73	0.66	XYL 1	1	99	99	
Campylobacter ureolyticus		0.18	-.03		1	1	1	
Peptostreptococcus anaerobius		0.09	-.03		99	99	1	
01004012	**Acceptable identification**							**01004012**
					SPO			
Clostridium novyi biovar A		92.27	0.53	XYL 1	99			
Atopobium minutum		6.13	0.29		1			
Clostridium sporogenes		1.12	0.06		99			
01004013	**Good identification**							**01004013**
Clostridium novyi biovar A		100.0	0.53	XYL 1				
01004020	**Low discrimination**							**01004020**
					SPO			
Atopobium minutum		47.22	0.58	XYL 1	1			
Clostridium novyi biovar A		20.35	0.56		99			
Clostridium cadaveris		11.70	0.37		99			
01004021	**Good identification**							**01004021**
					COC	SPO		
Clostridium novyi biovar A		95.33	0.56	XYL 1	1	99		
Atopobium minutum		2.23	0.24		1	1		
Peptococcus niger		0.83	-.02		99	1		
01004022	**Low discrimination**							**01004022**
					GRA	SPO		
Atopobium minutum		69.02	0.55	XYL 1	99	1		
Pseudoflavonifractor capillosus		21.07	0.32		1	1		
Clostridium novyi biovar A		7.88	0.43		99	99		
01004023	**Low discrimination**							**01004023**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	XYL 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01004030	**Good identification**						**01004030**
					SPO		
Clostridium novyi biovar A		96.81	0.56	XYL 1	99		
Atopobium minutum		2.27	0.24		1		
Clostridium cadaveris		0.56	0.03		99		
01004031	**Good identification**						**01004031**
Clostridium novyi biovar A		100.0	0.56	XYL 1			
01004032	**Low discrimination**						**01004032**
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	XYL 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01004033	**Good identification**						**01004033**
Clostridium novyi biovar A		100.0	0.43	XYL 1			
01004040	**Very good identification**						**01004040**
Clostridium glycolicum		97.18	0.83	FRU 90			
Clostridium botulinum biovar A		1.33	0.52				
Clostridium novyi biovar A		1.13	0.63				
01004041	**Low discrimination**						**01004041**
Clostridium novyi biovar A		53.04	0.63	XYL 1			
Clostridium glycolicum		46.11	0.49				
Clostridium botulinum biovar A		0.63	0.18				
01004042	**Low discrimination**						**01004042**
					SPO		
Clostridium glycolicum		79.62	0.49	FRU 90	99		
Clostridium novyi biovar A		16.16	0.50	BGL 1	99		
Atopobium minutum		1.61	0.29		1		
01004043	**Good identification**						**01004043**
Clostridium novyi biovar A		95.26	0.50	XYL 1			
Clostridium glycolicum		4.74	0.15				
01004050	**Low discrimination**						**01004050**
Clostridium novyi biovar A		53.11	0.63	XYL 1			
Clostridium glycolicum		46.18	0.49				
Clostridium botulinum biovar A		0.63	0.18				
01004051	**Very good identification**						**01004051**
Clostridium novyi biovar A		99.13	0.63	XYL 1			
Clostridium glycolicum		0.87	0.15				
01004052	**Good identification**						**01004052**
Clostridium novyi biovar A		95.26	0.50	XYL 1			
Clostridium glycolicum		4.74	0.15				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01004053	**Good identification**						**01004053**
Clostridium novyi biovar A		100.0	0.50	XYL 1			
01004060	**Low discrimination**						**01004060**
Clostridium glycolicum		75.51	0.49	FRU 90			
Clostridium novyi biovar A		21.71	0.53	NAG 1			
Clostridium botulinum biovar A		1.04	0.18				
01004061	**Good identification**						**01004061**
Clostridium novyi biovar A		96.61	0.53	XYL 1			
Clostridium glycolicum		3.39	0.15				
01004062	**Low discrimination**						**01004062**
					SPO		
Clostridium novyi biovar A		71.84	0.40	XYL 1	99		
Clostridium glycolicum		14.30	0.15		99		
Atopobium minutum		9.54	0.21		1		
01004063	**Good identification**						**01004063**
Clostridium novyi biovar A		100.0	0.40	XYL 1			
01004070	**Good identification**						**01004070**
Clostridium novyi biovar A		96.61	0.53	XYL 1			
Clostridium glycolicum		3.39	0.15				
01004071	**Good identification**						**01004071**
Clostridium novyi biovar A		100.0	0.53	XYL 1			
01004072	**Good identification**						**01004072**
Clostridium novyi biovar A		100.0	0.40	XYL 1			
01004073	**Good identification**						**01004073**
Clostridium novyi biovar A		100.0	0.40	XYL 1			
01004100	**Low discrimination**						**01004100**
					COC	SPO	
Actinomyces meyeri		76.43	0.55	SUC 93	1	1	
Peptostreptococcus anaerobius		10.44	0.32	FRU 95	99	1	
Clostridium glycolicum		3.45	0.27		1	99	
01004120	**Low discrimination**						**01004120**
					SPO		
Actinomyces meyeri		87.65	0.40	SUC 93	1		
Atopobium minutum		5.57	0.24	FRU 95	1		
Clostridium novyi biovar A		2.40	0.21	NAG 12	99		
01004140	**Good identification**						**01004140**
					SPO		
Clostridium glycolicum		96.27	0.49	FRU 90	99		
Clostridium botulinum biovar A		1.32	0.18	LAC 1	99		
Actinomyces meyeri		1.13	0.21		1		
01004240	**Good identification**						**01004240**
Clostridium glycolicum		97.38	0.49	FRU 90			
Clostridium botulinum biovar A		1.33	0.18	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		1.13	0.29			
01004440	**Good identification**					**01004440**
Clostridium glycolicum		97.38	0.49	FRU 90		
Clostridium botulinum biovar A		1.33	0.18	ARA 1		
Clostridium novyi biovar A		1.13	0.29			
01004500	**Good identification**					**01004500**
					COC	
Actinomyces meyeri		99.45	0.45	SUC 93	1	
Peptostreptococcus anaerobius		0.55	-.03	FRU 95	99	
01005000	**Low discrimination**					**01005000**
					SPO	
Atopobium minutum		39.11	0.58	XYL 1	1	
Clostridium glycolicum		22.63	0.45		99	
Actinomyces meyeri		17.10	0.48		1	
01005001	**Low discrimination**					**01005001**
					SPO	
Clostridium novyi biovar A		87.10	0.50	GAL 10	99	
Atopobium minutum		4.59	0.24	XYL 1	1	
Clostridium glycolicum		2.66	0.11		99	
01005002	**Acceptable identification**					**01005002**
					SPO	
Atopobium minutum		90.33	0.55	XYL 1	1	
Clostridium novyi biovar A		4.58	0.37		99	
Actinomyces meyeri		1.83	0.22		1	
01005003	**Low discrimination**					**01005003**
					SPO	
Clostridium novyi biovar A		83.39	0.37	GAL 10	99	
Atopobium minutum		16.61	0.21	XYL 1	1	
01005010	**Low discrimination**					**01005010**
					SPO	
Clostridium novyi biovar A		89.71	0.50	GAL 10	99	
Atopobium minutum		4.73	0.24	XYL 1	1	
Clostridium glycolicum		2.74	0.11		99	
01005011	**Good identification**					**01005011**
Clostridium novyi biovar A		100.0	0.50	GAL 10		
01005012	**Low discrimination**					**01005012**
					SPO	
Clostridium novyi biovar A		83.39	0.37	GAL 10	99	
Atopobium minutum		16.61	0.21	XYL 1	1	
01005013	**Good identification**					**01005013**
Clostridium novyi biovar A		100.0	0.37	GAL 10		
01005020	**Low discrimination**					**01005020**
					SPO	
Atopobium minutum		72.37	0.50	XYL 1	1	
Actinomyces meyeri		12.94	0.33		1	
Clostridium novyi biovar A		10.40	0.39		99	
01005021	**Acceptable identification**					**01005021**
					SPO	
Clostridium novyi biovar A		93.43	0.39	GAL 10	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		6.57	0.15	XYL 1	1		
01005022	**Low discrimination**						**01005022**
					GRA	SPO	
Atopobium minutum		85.66	0.47	XYL 1	99	1	
Pseudoflavonifractor capillosus		8.72	0.15		1	1	
Clostridium novyi biovar A		3.26	0.26		99	99	
01005030	**Acceptable identification**						**01005030**
					SPO		
Clostridium novyi biovar A		93.43	0.39	GAL 10	99		
Atopobium minutum		6.57	0.15	XYL 1	1		
01005031	**Good identification**						**01005031**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01005040	**Very good identification**						**01005040**
Clostridium glycolicum		98.59	0.67	FRU 90			
Clostridium novyi biovar A		1.15	0.47	GAL 10			
Clostridium botulinum biovar A		0.12	0.18				
01005041	**Low discrimination**						**01005041**
Clostridium novyi biovar A		53.49	0.47	GAL 10			
Clostridium glycolicum		46.51	0.33	XYL 1			
01005050	**Low discrimination**						**01005050**
Clostridium novyi biovar A		53.49	0.47	GAL 10			
Clostridium glycolicum		46.51	0.33	XYL 1			
01005051	**Good identification**						**01005051**
Clostridium novyi biovar A		99.13	0.47	GAL 10			
Clostridium glycolicum		0.87	-.02	XYL 1			
01005061	**Good identification**						**01005061**
Clostridium novyi biovar A		96.61	0.36	GAL 10			
Clostridium glycolicum		3.39	-.02	XYL 1			
01005070	**Good identification**						**01005070**
Clostridium novyi biovar A		96.61	0.36	GAL 10			
Clostridium glycolicum		3.39	-.02	XYL 1			
01005071	**Good identification**						**01005071**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01005100	**Very good identification**						**01005100**
					COC		
Actinomyces meyeri		96.22	0.58	SUC 93	1		
Blautia hansenii		1.46	0.19	FRU 95	99		
Lactobacillus fermentum		1.16	0.36		1		
01005120	**Acceptable identification**						**01005120**
Actinomyces meyeri		94.71	0.43	SUC 93			
Lactobacillus fermentum		2.79	0.28	FRU 95			
Atopobium minutum		1.34	0.15	NAG 12			
01005400	**Low discrimination**						**01005400**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		74.25	0.37	SUC 93				
Lactobacillus fermentum		8.77	0.32	FRU 95				
Atopobium minutum		6.86	0.24					
01005500	**Acceptable identification**							**01005500**
					COC			
Actinomyces meyeri		94.44	0.48	SUC 93	1			
Lactobacillus fermentum		4.55	0.36	FRU 95	1			
Blautia hansenii		1.01	0.06		99			
01005502	**Low discrimination**							**01005502**
Lactobacillus fermentum		60.89	0.36	SUC 87				
Actinomyces meyeri		39.11	0.22	FRU 99				
01006040	**Low discrimination**							**01006040**
					COC	SPO	GRA	
Clostridium glycolicum		88.56	0.49	FRU 90	1	99	99	
Megasphaera elsdenii		6.54	0.35	MAN 1	99	1	1	
Pseudoramibacter alactolyticus		2.41	0.17		1	1	99	
01010000	**Low discrimination**							**01010000**
					GRA			
Pseudoramibacter alactolyticus		66.71	1.00		99			
Fusobacterium varium		9.37	0.86		1			
Atopobium minutum		9.24	0.99		99			
01010001	**Low discrimination**							**01010001**
					GRA	SPO		
Pseudoramibacter alactolyticus		39.22	0.66	URE 1	99	1		
Clostridium novyi biovar A		37.21	0.84		99	99		
Fusobacterium varium		5.51	0.51		1	1		
01010002	**Low discrimination**							**01010002**
					COC	GRA		
Pseudoramibacter alactolyticus		49.91	0.84	BGL 10	1	99		
Atopobium minutum		41.49	0.96		1	99		
Megasphaera elsdenii		5.25	0.78		99	1		
01010003	**Low discrimination**							**01010003**
					SPO			
Clostridium novyi biovar A		42.90	0.71	FRU 10	99			
Pseudoramibacter alactolyticus		28.47	0.50		1			
Atopobium minutum		23.66	0.62		1			
01010010	**Low discrimination**							**01010010**
					SPO			
Clostridium difficile		32.04	0.71	MAN 90	99			
Pseudoramibacter alactolyticus		27.90	0.66		1			
Clostridium novyi biovar A		26.47	0.84		99			
01010011	**Very good identification**							**01010011**
					SPO			
Clostridium novyi biovar A		97.17	0.84	FRU 10	99			
Clostridium difficile		1.19	0.36		99			
Pseudoramibacter alactolyticus		1.03	0.32		1			
01010012	**Low discrimination**							**01010012**
					SPO			
Clostridium novyi biovar A		41.88	0.71	FRU 10	99			
Pseudoramibacter alactolyticus		27.79	0.50		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium minutum		23.10	0.62		1			
01010013	**Very good identification**							**01010013**
					SPO			
Clostridium novyi biovar A		98.65	0.71	FRU 10	99			
Pseudoramibacter alactolyticus		0.66	0.15		1			
Atopobium minutum		0.55	0.28		1			
01010020	**Low discrimination**							**01010020**
					SPO			
Pseudoramibacter alactolyticus		72.57	0.87		1			
Atopobium minutum		18.99	0.91		1			
Clostridium cadaveris		5.10	0.71		99			
01010021	**Low discrimination**							**01010021**
					SPO			
Clostridium novyi biovar A		49.51	0.73	FRU 10	99			
Pseudoramibacter alactolyticus		36.83	0.53		1			
Atopobium minutum		9.64	0.57		1			
01010022	**Low discrimination**							**01010022**
					GRA			
Atopobium minutum		58.89	0.88		99			
Pseudoramibacter alactolyticus		37.51	0.71		99			
Pseudoflavonifractor capillosus		2.16	0.50		1			
01010023	**Low discrimination**							**01010023**
					SPO			
Clostridium novyi biovar A		44.71	0.61	FRU 10	99			
Atopobium minutum		32.88	0.54		1			
Pseudoramibacter alactolyticus		20.94	0.37		1			
01010030	**Low discrimination**							**01010030**
					SPO			
Clostridium novyi biovar A		48.46	0.73	FRU 10	99			
Pseudoramibacter alactolyticus		36.05	0.53		1			
Atopobium minutum		9.43	0.57		1			
01010031	**Very good identification**							**01010031**
					SPO			
Clostridium novyi biovar A		98.96	0.73	FRU 10	99			
Pseudoramibacter alactolyticus		0.74	0.19		1			
Atopobium minutum		0.20	0.23		1			
01010032	**Low discrimination**							**01010032**
					SPO			
Clostridium novyi biovar A		44.65	0.61	FRU 10	99			
Atopobium minutum		32.84	0.54		1			
Pseudoramibacter alactolyticus		20.92	0.37		1			
01010033	**Very good identification**							**01010033**
					SPO			
Clostridium novyi biovar A		98.80	0.61	FRU 10	99			
Atopobium minutum		0.73	0.20		1			
Pseudoramibacter alactolyticus		0.47	0.03		1			
01010040	**Low discrimination**							**01010040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		69.75	0.90		1	1	99	
Megasphaera elsdenii		19.58	0.91		99	1	1	
Clostridium glycolicum		6.61	0.78		1	99	99	
01010041	**Low discrimination**							**01010041**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO	GRA	
Clostridium novyi biovar A		63.99	0.81	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		25.29	0.56		1	1	99	
Megasphaera elsdenii		7.10	0.57		99	1	1	
01010042	**Low discrimination**							**01010042**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		73.15	0.73	BGL 10	1	1	99	
Megasphaera elsdenii		20.53	0.75		99	1	1	
Clostridium novyi biovar A		2.97	0.68		1	99	99	
01010043	**Low discrimination**							**01010043**
					COC	SPO	GRA	
Clostridium novyi biovar A		75.21	0.68	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		18.72	0.39		1	1	99	
Megasphaera elsdenii		5.25	0.41		99	1	1	
01010050	**Low discrimination**							**01010050**
					COC	SPO	GRA	
Clostridium novyi biovar A		60.53	0.81	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		23.93	0.56		1	1	99	
Megasphaera elsdenii		6.72	0.57		99	1	1	
01010051	**Very good identification**							**01010051**
					COC	SPO	GRA	
Clostridium novyi biovar A		99.35	0.81	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		0.40	0.21		1	1	99	
Megasphaera elsdenii		0.11	0.23		99	1	1	
01010052	**Low discrimination**							**01010052**
					COC	SPO	GRA	
Clostridium novyi biovar A		74.90	0.68	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		18.64	0.39		1	1	99	
Megasphaera elsdenii		5.23	0.41		99	1	1	
01010053	**Very good identification**							**01010053**
					COC	SPO	GRA	
Clostridium novyi biovar A		99.68	0.68	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		0.25	0.05		1	1	99	
Megasphaera elsdenii		0.07	0.06		99	1	1	
01010060	**Good identification**							**01010060**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		92.41	0.77		1	1	99	
Clostridium novyi biovar A		3.35	0.70		1	99	99	
Megasphaera elsdenii		1.49	0.57		99	1	1	
01010061	**Low discrimination**							**01010061**
					COC	SPO	GRA	
Clostridium novyi biovar A		77.42	0.70	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		21.60	0.43		1	1	99	
Megasphaera elsdenii		0.35	0.23		99	1	1	
01010062	**Low discrimination**							**01010062**
					SPO			
Pseudoramibacter alactolyticus		87.64	0.61	BGL 10	1			
Atopobium minutum		5.56	0.54		1			
Clostridium novyi biovar A		5.04	0.58		99			
01010063	**Low discrimination**							**01010063**
					SPO			
Clostridium novyi biovar A		84.06	0.58	FRU 10	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		14.77	0.26		1			
Atopobium minutum		0.94	0.20		1			
01010070	**Low discrimination**							**01010070**
					COC	SPO	GRA	
Clostridium novyi biovar A		77.17	0.70	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		21.53	0.43		1	1	99	
Megasphaera elsdenii		0.35	0.23		99	1	1	
01010071	**Very good identification**							**01010071**
					SPO			
Clostridium novyi biovar A		99.72	0.70	FRU 10	99			
Pseudoramibacter alactolyticus		0.28	0.09		1			
01010072	**Low discrimination**							**01010072**
					SPO			
Clostridium novyi biovar A		84.06	0.58	FRU 10	99			
Pseudoramibacter alactolyticus		14.77	0.26		1			
Atopobium minutum		0.94	0.20		1			
01010073	**Very good identification**							**01010073**
					SPO			
Clostridium novyi biovar A		99.82	0.58	FRU 10	99			
Pseudoramibacter alactolyticus		0.18	-.08		1			
01010100	**Low discrimination**							**01010100**
					GRA			
Pseudoramibacter alactolyticus		46.59	0.66	LAC 1	99			
Actinomyces meyeri		26.60	0.72		99			
Fusobacterium varium		6.55	0.51		1			
01010102	**Low discrimination**							**01010102**
Pseudoramibacter alactolyticus		44.39	0.50	LAC 1				
Atopobium minutum		36.89	0.62	BGL 10				
Actinomyces meyeri		7.05	0.46					
01010103	**Low discrimination**							**01010103**
					SPO			
Clostridium novyi biovar A		40.96	0.37	FRU 10	99			
Pseudoramibacter alactolyticus		27.18	0.15	LAC 1	1			
Atopobium minutum		22.59	0.28		1			
01010110	**Low discrimination**							**01010110**
					SPO			
Clostridium difficile		27.17	0.36	MAN 90	99			
Pseudoramibacter alactolyticus		23.67	0.32	LAC 1	1			
Clostridium novyi biovar A		22.45	0.50		99			
01010111	**Good identification**							**01010111**
					SPO			
Clostridium novyi biovar A		97.19	0.50	FRU 10	99			
Clostridium difficile		1.19	0.02	LAC 1	99			
Pseudoramibacter alactolyticus		1.04	-.03		1			
01010112	**Low discrimination**							**01010112**
					SPO			
Clostridium novyi biovar A		39.83	0.37	FRU 10	99			
Pseudoramibacter alactolyticus		26.43	0.15	LAC 1	1			
Atopobium minutum		21.97	0.28		1			
01010113	**Good identification**							**01010113**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.37	FRU 10				
01010120	**Low discrimination**							**01010120**
Pseudoramibacter alactolyticus		54.44	0.53	LAC 1				
Actinomyces meyeri		24.01	0.57					
Atopobium minutum		14.25	0.57					
01010121	**Low discrimination**							**01010121**
					SPO			
Clostridium novyi biovar A		43.12	0.39	FRU 10	99			
Pseudoramibacter alactolyticus		32.08	0.19	LAC 1	1			
Actinomyces meyeri		14.15	0.23		1			
01010122	**Low discrimination**							**01010122**
Atopobium minutum		55.69	0.54	LAC 1				
Pseudoramibacter alactolyticus		35.47	0.37					
Actinomyces meyeri		4.36	0.31					
01010130	**Low discrimination**							**01010130**
					SPO			
Clostridium novyi biovar A		42.23	0.39	FRU 10	99			
Pseudoramibacter alactolyticus		31.42	0.19	LAC 1	1			
Actinomyces meyeri		13.86	0.23		1			
01010131	**Good identification**							**01010131**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01010140	**Low discrimination**							**01010140**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		68.51	0.56	LAC 1	1	1	99	
Megasphaera elsdenii		19.23	0.57		99	1	1	
Clostridium glycolicum		6.49	0.43		1	99	99	
01010141	**Low discrimination**							**01010141**
					COC	SPO	GRA	
Clostridium novyi biovar A		64.12	0.47	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		25.34	0.21	LAC 1	1	1	99	
Megasphaera elsdenii		7.11	0.23		99	1	1	
01010142	**Low discrimination**							**01010142**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		72.75	0.39	LAC 1	1	1	99	
Megasphaera elsdenii		20.42	0.41	BGL 10	99	1	1	
Clostridium novyi biovar A		2.95	0.34		1	99	99	
01010150	**Low discrimination**							**01010150**
					COC	SPO	GRA	
Clostridium novyi biovar A		60.42	0.47	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		23.88	0.21	LAC 1	1	1	99	
Megasphaera elsdenii		6.70	0.23		99	1	1	
01010151	**Good identification**							**01010151**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01010160	**Acceptable identification**							**01010160**
					SPO			
Pseudoramibacter alactolyticus		90.97	0.43	LAC 1	1			
Clostridium novyi biovar A		3.29	0.36		99			
Actinomyces meyeri		1.62	0.23		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01010161	**Low discrimination**						**01010161**
					SPO		
Clostridium novyi biovar A		78.19	0.36	FRU 10	99		
Pseudoramibacter alactolyticus		21.81	0.09	LAC 1	1		
01010170	**Low discrimination**						**01010170**
					SPO		
Clostridium novyi biovar A		78.19	0.36	FRU 10	99		
Pseudoramibacter alactolyticus		21.81	0.09	LAC 1	1		
01010171	**Good identification**						**01010171**
Clostridium novyi biovar A		100.0	0.36	FRU 10			
01010200	**Low discrimination**						**01010200**
					GRA		
Pseudoramibacter alactolyticus		66.71	0.66	RHA 1	99		
Fusobacterium varium		9.37	0.51		1		
Atopobium minutum		9.24	0.65		99		
01010202	**Low discrimination**						**01010202**
					COC	GRA	
Pseudoramibacter alactolyticus		49.96	0.50	RHA 1	1	99	
Atopobium minutum		41.52	0.62	BGL 10	1	99	
Megasphaera elsdenii		5.26	0.44		99	1	
01010203	**Low discrimination**						**01010203**
					SPO		
Clostridium novyi biovar A		43.76	0.37	FRU 10	99		
Pseudoramibacter alactolyticus		29.04	0.15	RHA 1	1		
Atopobium minutum		24.14	0.28		1		
01010210	**Low discrimination**						**01010210**
					SPO		
Clostridium difficile		32.09	0.36	MAN 90	99		
Pseudoramibacter alactolyticus		27.95	0.32	RHA 1	1		
Clostridium novyi biovar A		26.52	0.50		99		
01010211	**Good identification**						**01010211**
					SPO		
Clostridium novyi biovar A		97.76	0.50	FRU 10	99		
Clostridium difficile		1.20	0.02	RHA 1	99		
Pseudoramibacter alactolyticus		1.04	-.03		1		
01010212	**Low discrimination**						**01010212**
					SPO		
Clostridium novyi biovar A		42.48	0.37	FRU 10	99		
Pseudoramibacter alactolyticus		28.19	0.15	RHA 1	1		
Atopobium minutum		23.43	0.28		1		
01010213	**Good identification**						**01010213**
Clostridium novyi biovar A		100.0	0.37	FRU 10			
01010220	**Low discrimination**						**01010220**
					SPO		
Pseudoramibacter alactolyticus		72.61	0.53	RHA 1	1		
Atopobium minutum		19.00	0.57		1		
Clostridium cadaveris		5.10	0.37		99		
01010221	**Low discrimination**						**01010221**
					SPO		
Clostridium novyi biovar A		50.23	0.39	FRU 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		37.37	0.19	RHA 1	1			
Atopobium minutum		9.78	0.23		1			
01010222	**Low discrimination**							**01010222**
					GRA			
Atopobium minutum		58.99	0.54	RHA 1	99			
Pseudoramibacter alactolyticus		37.57	0.37		99			
Pseudoflavonifractor capillosus		2.17	0.15		1			
01010230	**Low discrimination**							**01010230**
					SPO			
Clostridium novyi biovar A		49.02	0.39	FRU 10	99			
Pseudoramibacter alactolyticus		36.47	0.19	RHA 1	1			
Atopobium minutum		9.54	0.23		1			
01010231	**Good identification**							**01010231**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01010240	**Low discrimination**							**01010240**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		69.76	0.56	RHA 1	1	1	99	
Megasphaera elsdenii		19.58	0.57		99	1	1	
Clostridium glycolicum		6.61	0.43		1	99	99	
01010241	**Low discrimination**							**01010241**
					COC	SPO	GRA	
Clostridium novyi biovar A		64.50	0.47	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		25.49	0.21	RHA 1	1	1	99	
Megasphaera elsdenii		7.16	0.23		99	1	1	
01010242	**Low discrimination**							**01010242**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		73.26	0.39	RHA 1	1	1	99	
Megasphaera elsdenii		20.56	0.41	BGL 10	99	1	1	
Clostridium novyi biovar A		2.97	0.34		1	99	99	
01010250	**Low discrimination**							**01010250**
					COC	SPO	GRA	
Clostridium novyi biovar A		60.75	0.47	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		24.01	0.21	RHA 1	1	1	99	
Megasphaera elsdenii		6.74	0.23		99	1	1	
01010251	**Good identification**							**01010251**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01010260	**Acceptable identification**							**01010260**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		92.46	0.43	RHA 1	1	1	99	
Clostridium novyi biovar A		3.35	0.36		1	99	99	
Megasphaera elsdenii		1.49	0.23		99	1	1	
01010261	**Low discrimination**							**01010261**
					SPO			
Clostridium novyi biovar A		78.19	0.36	FRU 10	99			
Pseudoramibacter alactolyticus		21.81	0.09	RHA 1	1			
01010270	**Low discrimination**							**01010270**
					SPO			
Clostridium novyi biovar A		78.19	0.36	FRU 10	99			
Pseudoramibacter alactolyticus		21.81	0.09	RHA 1	1			
01010271	**Good identification**							**01010271**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.36	FRU 10			
01010400	**Low discrimination**						**01010400**
					GRA		
Pseudoramibacter alactolyticus		64.96	0.66	ARA 1	99		
Fusobacterium varium		9.13	0.51		1		
Atopobium minutum		9.00	0.65		99		
01010402	**Low discrimination**						**01010402**
					COC	GRA	
Pseudoramibacter alactolyticus		49.46	0.50	ARA 1	1	99	
Atopobium minutum		41.12	0.62	BGL 10	1	99	
Megasphaera elsdenii		5.21	0.44		99	1	
01010403	**Low discrimination**						**01010403**
					SPO		
Clostridium novyi biovar A		43.76	0.37	FRU 10	99		
Pseudoramibacter alactolyticus		29.04	0.15	ARA 1	1		
Atopobium minutum		24.14	0.28		1		
01010410	**Low discrimination**						**01010410**
					SPO		
Clostridium difficile		31.72	0.36	MAN 90	99		
Pseudoramibacter alactolyticus		27.63	0.32	ARA 1	1		
Clostridium novyi biovar A		26.21	0.50		99		
01010411	**Good identification**						**01010411**
					SPO		
Clostridium novyi biovar A		97.76	0.50	FRU 10	99		
Clostridium difficile		1.20	0.02	ARA 1	99		
Pseudoramibacter alactolyticus		1.04	-.03		1		
01010412	**Low discrimination**						**01010412**
					SPO		
Clostridium novyi biovar A		42.48	0.37	FRU 10	99		
Pseudoramibacter alactolyticus		28.19	0.15	ARA 1	1		
Atopobium minutum		23.43	0.28		1		
01010413	**Good identification**						**01010413**
Clostridium novyi biovar A		100.0	0.37	FRU 10			
01010420	**Low discrimination**						**01010420**
					SPO		
Pseudoramibacter alactolyticus		70.97	0.53	ARA 1	1		
Atopobium minutum		18.57	0.57		1		
Clostridium cadaveris		4.99	0.37		99		
01010421	**Low discrimination**						**01010421**
					SPO		
Clostridium novyi biovar A		49.72	0.39	FRU 10	99		
Pseudoramibacter alactolyticus		36.99	0.19	ARA 1	1		
Atopobium minutum		9.68	0.23		1		
01010422	**Low discrimination**						**01010422**
					GRA		
Atopobium minutum		58.41	0.54	ARA 1	99		
Pseudoramibacter alactolyticus		37.20	0.37		99		
Pseudoflavonifractor capillosus		2.15	0.15		1		
01010430	**Low discrimination**						**01010430**
					SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		48.54	0.39	FRU 10	99			
Pseudoramibacter alactolyticus		36.11	0.19	ARA 1	1			
Atopobium minutum		9.45	0.23		1			
01010431	**Good identification**							**01010431**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01010440	**Low discrimination**							**01010440**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		69.68	0.56	ARA 1	1	1	99	
Megasphaera elsdenii		19.56	0.57		99	1	1	
Clostridium glycolicum		6.60	0.43		1	99	99	
01010441	**Low discrimination**							**01010441**
					COC	SPO	GRA	
Clostridium novyi biovar A		64.50	0.47	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		25.49	0.21	ARA 1	1	1	99	
Megasphaera elsdenii		7.16	0.23		99	1	1	
01010442	**Low discrimination**							**01010442**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		73.26	0.39	ARA 1	1	1	99	
Megasphaera elsdenii		20.56	0.41	BGL 10	99	1	1	
Clostridium novyi biovar A		2.97	0.34		1	99	99	
01010450	**Low discrimination**							**01010450**
					COC	SPO	GRA	
Clostridium novyi biovar A		60.75	0.47	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		24.01	0.21	ARA 1	1	1	99	
Megasphaera elsdenii		6.74	0.23		99	1	1	
01010451	**Good identification**							**01010451**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01010460	**Acceptable identification**							**01010460**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		92.37	0.43	ARA 1	1	1	99	
Clostridium novyi biovar A		3.34	0.36		1	99	99	
Megasphaera elsdenii		1.48	0.23		99	1	1	
01010461	**Low discrimination**							**01010461**
					SPO			
Clostridium novyi biovar A		78.19	0.36	FRU 10	99			
Pseudoramibacter alactolyticus		21.81	0.09	ARA 1	1			
01010470	**Low discrimination**							**01010470**
					SPO			
Clostridium novyi biovar A		78.19	0.36	FRU 10	99			
Pseudoramibacter alactolyticus		21.81	0.09	ARA 1	1			
01010471	**Good identification**							**01010471**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01010500	**Low discrimination**							**01010500**
Actinomyces meyeri		88.98	0.62	SUC 93				
Pseudoramibacter alactolyticus		6.30	0.32					
Lactobacillus fermentum		1.12	0.40					
01010502	**Low discrimination**							**01010502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		52.13	0.36	SUC 93		
Lactobacillus fermentum		21.18	0.40	BGL 3		
Pseudoramibacter alactolyticus		13.26	0.15			
01010520	**Low discrimination**					**01010520**
Actinomyces meyeri		86.75	0.47	SUC 93		
Pseudoramibacter alactolyticus		7.95	0.19	NAG 12		
Lactobacillus fermentum		2.66	0.32			
01011000	**Low discrimination**					**01011000**
Atopobium minutum		70.86	0.91			
Pseudoramibacter alactolyticus		15.50	0.66			
Actinomyces meyeri		3.32	0.65			
01011001	**Low discrimination**					**01011001**
					SPO	
Clostridium novyi biovar A		56.76	0.67	FRU 10	99	
Atopobium minutum		24.86	0.57	GAL 10	1	
Clostridium sordellii		8.70	0.37		99	
01011002	**Good identification**					**01011002**
Atopobium minutum		91.89	0.88			
Pseudoramibacter alactolyticus		3.35	0.50			
Lactobacillus fermentum		3.28	0.70			
01011003	**Low discrimination**					**01011003**
					SPO	
Atopobium minutum		59.19	0.54	URE 1	1	
Clostridium novyi biovar A		35.77	0.54		99	
Pseudoramibacter alactolyticus		2.16	0.15		1	
01011010	**Low discrimination**					**01011010**
					SPO	
Clostridium novyi biovar A		58.14	0.67	FRU 10	99	
Atopobium minutum		25.47	0.57	GAL 10	1	
Clostridium difficile		6.40	0.36		99	
01011011	**Very good identification**					**01011011**
					SPO	
Clostridium novyi biovar A		99.20	0.67	FRU 10	99	
Atopobium minutum		0.44	0.23	GAL 10	1	
Clostridium sordellii		0.15	0.03		99	
01011012	**Low discrimination**					**01011012**
					SPO	
Atopobium minutum		59.15	0.54	MLZ 1	1	
Clostridium novyi biovar A		35.75	0.54		99	
Pseudoramibacter alactolyticus		2.16	0.15		1	
01011013	**Good identification**					**01011013**
					SPO	
Clostridium novyi biovar A		98.36	0.54	FRU 10	99	
Atopobium minutum		1.64	0.20	GAL 10	1	
01011020	**Low discrimination**					**01011020**
					CAT	
Atopobium minutum		76.21	0.83		1	
Propionibacterium acnes		9.36	0.60		95	
Pseudoramibacter alactolyticus		8.83	0.53		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01011021	**Low discrimination**						**01011021**
					SPO	CAT	
Clostridium novyi biovar A		56.84	0.57	FRU 10	99	1	
Atopobium minutum		33.20	0.49	GAL 10	1	1	
Propionibacterium acnes		4.08	0.26		1	95	
01011022	**Good identification**						**01011022**
Atopobium minutum		93.02	0.80				
Lactobacillus fermentum		3.32	0.62				
Pseudoramibacter alactolyticus		1.80	0.37				
01011023	**Low discrimination**						**01011023**
					SPO		
Atopobium minutum		65.77	0.46	URE 1	1		
Clostridium novyi biovar A		29.81	0.44		99		
Lactobacillus fermentum		2.35	0.28		1		
01011030	**Low discrimination**						**01011030**
					SPO	CAT	
Clostridium novyi biovar A		56.80	0.57	FRU 10	99	1	
Atopobium minutum		33.18	0.49	GAL 10	1	1	
Propionibacterium acnes		4.07	0.26		1	95	
01011031	**Very good identification**						**01011031**
					SPO		
Clostridium novyi biovar A		99.41	0.57	FRU 10	99		
Atopobium minutum		0.59	0.15	GAL 10	1		
01011032	**Low discrimination**						**01011032**
					SPO		
Atopobium minutum		65.77	0.46	MLZ 1	1		
Clostridium novyi biovar A		29.81	0.44		99		
Lactobacillus fermentum		2.35	0.28		1		
01011033	**Good identification**						**01011033**
					SPO		
Clostridium novyi biovar A		97.82	0.44	FRU 10	99		
Atopobium minutum		2.18	0.12	GAL 10	1		
01011040	**Low discrimination**						**01011040**
					SPO		
Clostridium glycolicum		36.78	0.61	GAL 10	99		
Pseudoramibacter alactolyticus		35.29	0.56	XYL 95	1		
Clostridium novyi biovar A		9.92	0.64		99		
01011041	**Good identification**						**01011041**
					SPO		
Clostridium novyi biovar A		90.88	0.64	FRU 10	99		
Clostridium glycolicum		3.40	0.27	GAL 10	99		
Pseudoramibacter alactolyticus		3.27	0.21		1		
01011042	**Low discrimination**						**01011042**
					SPO		
Atopobium minutum		37.24	0.54	SOR 1	1		
Pseudoramibacter alactolyticus		33.61	0.39		1		
Clostridium novyi biovar A		15.00	0.51		99		
01011043	**Acceptable identification**						**01011043**
					SPO		
Clostridium novyi biovar A		94.87	0.51	FRU 10	99		
Atopobium minutum		2.38	0.20	GAL 10	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Pseudoramibacter alactolyticus		2.15	0.05		1		
01011050	**Good identification**						**01011050**
					SPO		
Clostridium novyi biovar A		90.96	0.64	FRU 10	99		
Clostridium glycolicum		3.41	0.27	GAL 10	99		
Pseudoramibacter alactolyticus		3.27	0.21		1		
01011051	**Very good identification**						**01011051**
Clostridium novyi biovar A		100.0	0.64	FRU 10			
01011052	**Acceptable identification**						**01011052**
					SPO		
Clostridium novyi biovar A		94.87	0.51	FRU 10	99		
Atopobium minutum		2.38	0.20	GAL 10	1		
Pseudoramibacter alactolyticus		2.15	0.05		1		
01011053	**Good identification**						**01011053**
Clostridium novyi biovar A		100.0	0.51	FRU 10			
01011060	**Low discrimination**						**01011060**
					SPO	CAT	
Pseudoramibacter alactolyticus		36.07	0.43	GAL 1	1	1	
Propionibacterium acnes		33.58	0.49		1	95	
Clostridium novyi biovar A		14.36	0.54		99	1	
01011061	**Acceptable identification**						**01011061**
					SPO	CAT	
Clostridium novyi biovar A		94.53	0.54	FRU 10	99	1	
Pseudoramibacter alactolyticus		2.40	0.09	GAL 10	1	1	
Propionibacterium acnes		2.23	0.15		1	95	
01011062	**Low discrimination**						**01011062**
					SPO		
Atopobium minutum		53.31	0.46	SOR 1	1		
Pseudoramibacter alactolyticus		25.46	0.26		1		
Clostridium novyi biovar A		16.11	0.41		99		
01011063	**Good identification**						**01011063**
					SPO		
Clostridium novyi biovar A		95.29	0.41	FRU 10	99		
Atopobium minutum		3.19	0.12	GAL 10	1		
Pseudoramibacter alactolyticus		1.52	-.08		1		
01011070	**Acceptable identification**						**01011070**
					SPO	CAT	
Clostridium novyi biovar A		94.53	0.54	FRU 10	99	1	
Pseudoramibacter alactolyticus		2.40	0.09	GAL 10	1	1	
Propionibacterium acnes		2.23	0.15		1	95	
01011071	**Good identification**						**01011071**
Clostridium novyi biovar A		100.0	0.54	FRU 10			
01011072	**Good identification**						**01011072**
					SPO		
Clostridium novyi biovar A		95.29	0.41	FRU 10	99		
Atopobium minutum		3.19	0.12	GAL 10	1		
Pseudoramibacter alactolyticus		1.52	-.08		1		
01011073	**Good identification**						**01011073**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.41	FRU 10		
01011100	**Low discrimination**					**01011100**
					GRA	
Actinomyces meyeri		70.76	0.75	SUC 93	99	
Lactobacillus fermentum		14.67	0.74		99	
Fusobacterium mortiferum		9.50	0.69		1	
01011101	**Low discrimination**					**01011101**
					GRA	
Actinomyces meyeri		63.75	0.41	SUC 93	99	
Lactobacillus fermentum		13.22	0.40	URE 1	99	
Fusobacterium mortiferum		8.56	0.35		1	
01011102	**Low discrimination**					**01011102**
Lactobacillus fermentum		71.30	0.74	SUC 87		
Atopobium minutum		12.36	0.54			
Actinomyces meyeri		10.64	0.49			
01011103	**Low discrimination**					**01011103**
Lactobacillus fermentum		66.70	0.40	SUC 87		
Atopobium minutum		11.57	0.20	URE 1		
Actinomyces meyeri		9.95	0.15			
01011110	**Low discrimination**					**01011110**
					GRA	
Actinomyces meyeri		64.77	0.41	SUC 93	99	
Lactobacillus fermentum		13.43	0.40	MLZ 1	99	
Fusobacterium mortiferum		8.69	0.35		1	
01011112	**Low discrimination**					**01011112**
Lactobacillus fermentum		66.70	0.40	SUC 87		
Atopobium minutum		11.57	0.20	MLZ 1		
Actinomyces meyeri		9.95	0.15			
01011120	**Low discrimination**					**01011120**
Actinomyces meyeri		59.05	0.60	SUC 93		
Lactobacillus fermentum		29.93	0.66	NAG 12		
Atopobium minutum		7.79	0.49			
01011122	**Low discrimination**					**01011122**
Lactobacillus fermentum		80.37	0.66	SUC 87		
Atopobium minutum		13.94	0.46			
Actinomyces meyeri		4.91	0.34			
01011140	**Low discrimination**					**01011140**
					SPO	
Actinomyces meyeri		46.28	0.41	SUC 93	1	
Clostridium glycolicum		14.08	0.27	SOR 1	99	
Pseudoramibacter alactolyticus		13.51	0.21		1	
01011142	**Low discrimination**					**01011142**
Lactobacillus fermentum		60.07	0.40	SUC 87		
Atopobium minutum		10.42	0.20	SOR 1		
Pseudoramibacter alactolyticus		9.40	0.05			
01011200	**Low discrimination**					**01011200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Atopobium minutum		70.90	0.57	RHA 1		
Pseudoramibacter alactolyticus		15.51	0.32			
Actinomyces meyeri		3.32	0.31			
01011202	**Acceptable identification**					**01011202**
Atopobium minutum		92.08	0.54	RHA 1		
Pseudoramibacter alactolyticus		3.36	0.15			
Lactobacillus fermentum		3.29	0.36			
01011220	**Low discrimination**					**01011220**
					CAT	
Atopobium minutum		76.51	0.49	RHA 1	1	
Propionibacterium acnes		9.39	0.26		95	
Pseudoramibacter alactolyticus		8.86	0.19		1	
01011222	**Acceptable identification**					**01011222**
Atopobium minutum		93.30	0.46	RHA 1		
Lactobacillus fermentum		3.33	0.28			
Pseudoramibacter alactolyticus		1.80	0.03			
01011300	**Low discrimination**					**01011300**
					GRA	
Actinomyces meyeri		71.06	0.41	SUC 93	99	
Lactobacillus fermentum		14.73	0.40	RHA 1	99	
Fusobacterium mortiferum		9.54	0.35		1	
01011302	**Low discrimination**					**01011302**
Lactobacillus fermentum		71.71	0.40	SUC 87		
Atopobium minutum		12.44	0.20	RHA 1		
Actinomyces meyeri		10.70	0.15			
01011400	**Low discrimination**					**01011400**
Lactobacillus fermentum		47.96	0.70	SUC 87		
Actinomyces meyeri		23.59	0.54			
Atopobium minutum		20.35	0.57			
01011401	**Low discrimination**					**01011401**
					SPO	
Lactobacillus fermentum		31.98	0.36	SUC 87	1	
Clostridium novyi biovar A		30.99	0.33	URE 1	99	
Actinomyces meyeri		15.73	0.20		1	
01011402	**Low discrimination**					**01011402**
Lactobacillus fermentum		76.23	0.70	SUC 87		
Atopobium minutum		21.57	0.54			
Actinomyces meyeri		1.16	0.29			
01011403	**Low discrimination**					**01011403**
					SPO	
Lactobacillus fermentum		68.78	0.36	SUC 87	1	
Atopobium minutum		19.46	0.20	URE 1	1	
Clostridium novyi biovar A		11.76	0.20		99	
01011410	**Low discrimination**					**01011410**
					SPO	
Lactobacillus fermentum		32.42	0.36	SUC 87	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		31.41	0.33	MLZ 1	99	
Actinomyces meyeri		15.95	0.20		1	
01011412	**Low discrimination**					**01011412**
					SPO	
Lactobacillus fermentum		68.78	0.36	SUC 87	1	
Atopobium minutum		19.46	0.20	MLZ 1	1	
Clostridium novyi biovar A		11.76	0.20		99	
01011420	**Low discrimination**					**01011420**
Lactobacillus fermentum		57.24	0.62	SUC 87		
Atopobium minutum		24.30	0.49			
Actinomyces meyeri		11.52	0.40			
01011422	**Low discrimination**					**01011422**
Lactobacillus fermentum		76.96	0.62	SUC 87		
Atopobium minutum		21.78	0.46			
Actinomyces meyeri		0.48	0.14			
01011442	**Low discrimination**					**01011442**
Lactobacillus fermentum		58.01	0.36	SUC 87		
Atopobium minutum		16.41	0.20	SOR 1		
Pseudoramibacter alactolyticus		14.81	0.05			
01011500	**Low discrimination**					**01011500**
					GRA	
Actinomyces meyeri		54.43	0.65	SUC 93	99	
Lactobacillus fermentum		45.13	0.74		99	
Fusobacterium mortiferum		0.30	0.35		1	
01011502	**Very good identification**					**01011502**
Lactobacillus fermentum		96.18	0.74	SUC 87		
Actinomyces meyeri		3.59	0.39			
Atopobium minutum		0.17	0.20			
01011503	**Good identification**					**01011503**
Lactobacillus fermentum		96.41	0.40	SUC 87		
Actinomyces meyeri		3.60	0.05	URE 1		
01011512	**Good identification**					**01011512**
Lactobacillus fermentum		96.41	0.40	SUC 87		
Actinomyces meyeri		3.60	0.05	MLZ 1		
01011520	**Low discrimination**					**01011520**
Lactobacillus fermentum		66.76	0.66	SUC 87		
Actinomyces meyeri		32.93	0.50			
Atopobium minutum		0.18	0.15			
01011522	**Very good identification**					**01011522**
Lactobacillus fermentum		98.33	0.66	SUC 87		
Actinomyces meyeri		1.50	0.24			
Atopobium minutum		0.17	0.12			
01011542	**Good identification**					**01011542**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		96.41	0.40	SUC 87				
Actinomyces meyeri		3.60	0.05	SOR 1				
01011600	**Low discrimination**							**01011600**
Lactobacillus fermentum		49.77	0.36	SUC 87				
Actinomyces meyeri		24.48	0.20	RHA 1				
Atopobium minutum		21.13	0.23					
01011602	**Low discrimination**							**01011602**
Lactobacillus fermentum		77.95	0.36	SUC 87				
Atopobium minutum		22.05	0.20	RHA 1				
01011702	**Good identification**							**01011702**
Lactobacillus fermentum		96.41	0.40	SUC 87				
Actinomyces meyeri		3.60	0.05	RHA 1				
01012000	**Low discrimination**							**01012000**
					COC	GRA		
Pseudoramibacter alactolyticus		53.26	0.95		1	99		
Eubacterium limosum		33.87	0.94		1	99		
Megasphaera elsdenii		10.41	0.94		99	1		
01012001	**Low discrimination**							**01012001**
					COC	GRA		
Pseudoramibacter alactolyticus		52.72	0.61	URE 1	1	99		
Eubacterium limosum		33.53	0.60		1	99		
Megasphaera elsdenii		10.31	0.60		99	1		
01012002	**Low discrimination**							**01012002**
					COC	GRA		
Pseudoramibacter alactolyticus		78.51	0.79	BGL 10	1	99		
Megasphaera elsdenii		15.35	0.78		99	1		
Eubacterium limosum		4.54	0.60		1	99		
01012003	**Low discrimination**							**01012003**
					COC	GRA		
Pseudoramibacter alactolyticus		76.88	0.45	URE 1	1	99		
Megasphaera elsdenii		15.03	0.44	BGL 10	99	1		
Eubacterium limosum		4.45	0.26		1	99		
01012010	**Good identification**							**01012010**
					SPO			
Clostridium difficile		91.18	0.87		99			
Pseudoramibacter alactolyticus		4.75	0.61		1			
Eubacterium limosum		3.02	0.60		1			
01012011	**Low discrimination**							**01012011**
					SPO			
Clostridium difficile		84.23	0.53	URE 1	99			
Clostridium novyi biovar A		7.73	0.50		99			
Pseudoramibacter alactolyticus		4.39	0.27		1			
01012012	**Low discrimination**							**01012012**
					COC	SPO	GRA	
Clostridium difficile		57.34	0.53	BGL 1	1	99	99	
Pseudoramibacter alactolyticus		32.86	0.45		1	1	99	
Megasphaera elsdenii		6.42	0.44		99	1	1	
01012013	**Low discrimination**							**01012013**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		48.77	0.37	FRU 10	99			
Clostridium difficile		30.40	0.19	MAN 1	99			
Pseudoramibacter alactolyticus		17.43	0.11		1			
01012020	**Good identification**							**01012020**
					COC	GRA		
Pseudoramibacter alactolyticus		93.65	0.83		1	99		
Eubacterium limosum		3.41	0.60		1	99		
Megasphaera elsdenii		1.05	0.60		99	1		
01012021	**Acceptable identification**							**01012021**
					SPO			
Pseudoramibacter alactolyticus		91.53	0.48	URE 1	1			
Eubacterium limosum		3.33	0.26		1			
Clostridium novyi biovar A		2.31	0.39		99			
01012022	**Very good identification**							**01012022**
					COC	GRA		
Pseudoramibacter alactolyticus		95.52	0.66	BGL 10	1	99		
Atopobium minutum		2.81	0.54		1	99		
Megasphaera elsdenii		1.07	0.44		99	1		
01012030	**Low discrimination**							**01012030**
					SPO			
Clostridium difficile		50.18	0.53	NAG 1	99			
Pseudoramibacter alactolyticus		45.68	0.48		1			
Eubacterium limosum		1.66	0.26		1			
01012031	**Low discrimination**							**01012031**
					SPO			
Clostridium novyi biovar A		54.33	0.39	FRU 10	99			
Clostridium difficile		23.91	0.19	MAN 1	99			
Pseudoramibacter alactolyticus		21.76	0.14		1			
01012040	**Low discrimination**							**01012040**
					COC	GRA		
Pseudoramibacter alactolyticus		64.18	0.85		1	99		
Megasphaera elsdenii		33.46	0.91		99	1		
Eubacterium limosum		1.65	0.60		1	99		
01012041	**Low discrimination**							**01012041**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		62.31	0.51	URE 1	1	1	99	
Megasphaera elsdenii		32.48	0.57		99	1	1	
Clostridium novyi biovar A		2.96	0.47		1	99	99	
01012042	**Low discrimination**							**01012042**
					COC	GRA		
Pseudoramibacter alactolyticus		65.53	0.69	BGL 10	1	99		
Megasphaera elsdenii		34.16	0.75		99	1		
Eubacterium limosum		0.15	0.26		1	99		
01012050	**Low discrimination**							**01012050**
					COC	SPO	GRA	
Clostridium difficile		71.68	0.65	SOR 5	1	99	99	
Pseudoramibacter alactolyticus		17.74	0.51		1	1	99	
Megasphaera elsdenii		9.25	0.57		99	1	1	
01012051	**Low discrimination**							**01012051**
					SPO			
Clostridium novyi biovar A		45.80	0.47	FRU 10	99			
Clostridium difficile		39.38	0.31	MAN 1	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		9.75	0.17		1			
01012060	**Very good identification**							**01012060**
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		95.91	0.72		1	1	99	
Megasphaera elsdenii		2.86	0.57		99	1	1	
Propionibacterium acnes		0.94	0.45		1	95	99	
01012061	**Acceptable identification**							**01012061**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		90.34	0.38	URE 1	1	1	99	
Clostridium novyi biovar A		6.08	0.36		1	99	99	
Megasphaera elsdenii		2.70	0.23		99	1	1	
01012062	**Good identification**							**01012062**
					COC	GRA		
Pseudoramibacter alactolyticus		96.81	0.56	BGL 10	1	99		
Megasphaera elsdenii		2.89	0.41		99	1		
Atopobium minutum		0.12	0.20		1	99		
01012070	**Low discrimination**							**01012070**
					SPO			
Pseudoramibacter alactolyticus		74.73	0.38	MLZ 1	1			
Clostridium difficile		17.28	0.31		99			
Clostridium novyi biovar A		5.03	0.36		99			
01012071	**Low discrimination**							**01012071**
					SPO			
Clostridium novyi biovar A		84.39	0.36	FRU 10	99			
Pseudoramibacter alactolyticus		12.68	0.04	MAN 1	1			
Clostridium difficile		2.93	-.03		99			
01012100	**Low discrimination**							**01012100**
					COC	GRA		
Pseudoramibacter alactolyticus		52.92	0.61	LAC 1	1	99		
Eubacterium limosum		33.66	0.60		1	99		
Megasphaera elsdenii		10.35	0.60		99	1		
01012102	**Low discrimination**							**01012102**
					COC	GRA		
Pseudoramibacter alactolyticus		78.21	0.45	LAC 1	1	99		
Megasphaera elsdenii		15.29	0.44	BGL 10	99	1		
Eubacterium limosum		4.52	0.26		1	99		
01012110	**Acceptable identification**							**01012110**
					SPO			
Clostridium difficile		91.17	0.53	LAC 1	99			
Pseudoramibacter alactolyticus		4.75	0.27		1			
Eubacterium limosum		3.02	0.26		1			
01012120	**Acceptable identification**							**01012120**
					COC	GRA		
Pseudoramibacter alactolyticus		92.98	0.48	LAC 1	1	99		
Eubacterium limosum		3.39	0.26		1	99		
Megasphaera elsdenii		1.04	0.26		99	1		
01012140	**Low discrimination**							**01012140**
					COC	GRA		
Pseudoramibacter alactolyticus		64.15	0.51	LAC 1	1	99		
Megasphaera elsdenii		33.44	0.57		99	1		
Eubacterium limosum		1.65	0.26		1	99		
01012160	**Good identification**							**01012160**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		96.19	0.38	LAC 1	1	1	99	
Megasphaera elsdenii		2.87	0.23		99	1	1	
Propionibacterium acnes		0.95	0.11		1	95	99	
01012200	**Low discrimination**							**01012200**
					COC	GRA		
Pseudoramibacter alactolyticus		53.26	0.61	RHA 1	1	99		
Eubacterium limosum		33.87	0.60		1	99		
Megasphaera elsdenii		10.41	0.60		99	1		
01012202	**Low discrimination**							**01012202**
					COC	GRA		
Pseudoramibacter alactolyticus		78.59	0.45	RHA 1	1	99		
Megasphaera elsdenii		15.36	0.44	BGL 10	99	1		
Eubacterium limosum		4.54	0.26		1	99		
01012210	**Acceptable identification**							**01012210**
					SPO			
Clostridium difficile		91.21	0.53	RHA 1	99			
Pseudoramibacter alactolyticus		4.75	0.27		1			
Eubacterium limosum		3.02	0.26		1			
01012220	**Acceptable identification**							**01012220**
					COC	GRA		
Pseudoramibacter alactolyticus		93.70	0.48	RHA 1	1	99		
Eubacterium limosum		3.41	0.26		1	99		
Megasphaera elsdenii		1.05	0.26		99	1		
01012240	**Low discrimination**							**01012240**
					COC	GRA		
Pseudoramibacter alactolyticus		64.19	0.51	RHA 1	1	99		
Megasphaera elsdenii		33.46	0.57		99	1		
Eubacterium limosum		1.65	0.26		1	99		
01012260	**Good identification**							**01012260**
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		96.19	0.38	RHA 1	1	1	99	
Megasphaera elsdenii		2.87	0.23		99	1	1	
Propionibacterium acnes		0.95	0.11		1	95	99	
01012400	**Low discrimination**							**01012400**
					COC	GRA		
Pseudoramibacter alactolyticus		53.25	0.61	ARA 1	1	99		
Eubacterium limosum		33.86	0.60		1	99		
Megasphaera elsdenii		10.41	0.60		99	1		
01012402	**Low discrimination**							**01012402**
					COC	GRA		
Pseudoramibacter alactolyticus		78.59	0.45	ARA 1	1	99		
Megasphaera elsdenii		15.36	0.44	BGL 10	99	1		
Eubacterium limosum		4.54	0.26		1	99		
01012410	**Acceptable identification**							**01012410**
					SPO			
Clostridium difficile		91.21	0.53	ARA 1	99			
Pseudoramibacter alactolyticus		4.75	0.27		1			
Eubacterium limosum		3.02	0.26		1			
01012420	**Acceptable identification**							**01012420**
					COC	GRA		
Pseudoramibacter alactolyticus		93.65	0.48	ARA 1	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium limosum		3.41	0.26		1	99		
Megasphaera elsdenii		1.05	0.26		99	1		
01012440	**Low discrimination**							**01012440**
					COC	GRA		
Pseudoramibacter alactolyticus		64.19	0.51	ARA 1	1	99		
Megasphaera elsdenii		33.46	0.57		99	1		
Eubacterium limosum		1.65	0.26		1	99		
01012460	**Good identification**							**01012460**
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		96.19	0.38	ARA 1	1	1	99	
Megasphaera elsdenii		2.87	0.23		99	1	1	
Propionibacterium acnes		0.95	0.11		1	95	99	
01013000	**Low discrimination**							**01013000**
					COC	GRA		
Pseudoramibacter alactolyticus		50.63	0.61	GAL 1	1	99		
Eubacterium limosum		32.20	0.60		1	99		
Megasphaera elsdenii		9.90	0.60		99	1		
01013002	**Low discrimination**							**01013002**
					COC	GRA		
Pseudoramibacter alactolyticus		55.49	0.45	GAL 1	1	99		
Atopobium minutum		28.56	0.54	BGL 10	1	99		
Megasphaera elsdenii		10.85	0.44		99	1		
01013010	**Low discrimination**							**01013010**
					SPO			
Clostridium difficile		89.91	0.53	GAL 1	99			
Pseudoramibacter alactolyticus		4.68	0.27		1			
Eubacterium limosum		2.98	0.26		1			
01013020	**Low discrimination**							**01013020**
					CAT			
Propionibacterium acnes		47.44	0.56	IND 87	95			
Pseudoramibacter alactolyticus		42.83	0.48	NIT 92	1			
Atopobium minutum		6.94	0.49		1			
01013040	**Low discrimination**							**01013040**
					COC	GRA		
Pseudoramibacter alactolyticus		63.06	0.51	GAL 1	1	99		
Megasphaera elsdenii		32.88	0.57		99	1		
Eubacterium limosum		1.62	0.26		1	99		
01013060	**Low discrimination**							**01013060**
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		49.60	0.38	GAL 1	1	1	99	
Propionibacterium acnes		48.23	0.45		1	95	99	
Megasphaera elsdenii		1.48	0.23		99	1	1	
01013100	**Low discrimination**							**01013100**
Actinomyces meyeri		38.76	0.41	SUC 93				
Pseudoramibacter alactolyticus		24.38	0.27	MAN 1				
Eubacterium limosum		15.50	0.26					
01013102	**Low discrimination**							**01013102**
Lactobacillus fermentum		53.98	0.40	SUC 87				
Pseudoramibacter alactolyticus		18.19	0.11	MAN 1				
Atopobium minutum		9.36	0.20					

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
01013402	**Low discrimination**						**01013402**
Lactobacillus fermentum		51.54	0.36	SUC 87			
Pseudoramibacter alactolyticus		28.34	0.11	MAN 1			
Atopobium minutum		14.58	0.20				
01013502	**Good identification**						**01013502**
Lactobacillus fermentum		96.41	0.40	SUC 87			
Actinomyces meyeri		3.60	0.05	MAN 1			
01014000	**Low discrimination**						**01014000**
					GRA	SPO	
Clostridium glycolicum		51.25	0.78	SOR 95	99	99	
Pseudoramibacter alactolyticus		21.86	0.66		99	1	
Fusobacterium varium		12.67	0.62		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01014001	**Low discrimination**					**01014001**
					SPO	
Clostridium glycolicum		41.75	0.43	SOR 95	99	
Pseudoramibacter alactolyticus		17.81	0.32	URE 1	1	
Clostridium novyi biovar A		16.90	0.50		99	
01014002	**Low discrimination**					**01014002**
					SPO	
Pseudoramibacter alactolyticus		42.71	0.50	XYL 1	1	
Atopobium minutum		35.50	0.62	BGL 10	1	
Clostridium glycolicum		9.10	0.43		99	
01014003	**Low discrimination**					**01014003**
					SPO	
Clostridium novyi biovar A		39.80	0.37	FRU 10	99	
Pseudoramibacter alactolyticus		26.41	0.15	XYL 1	1	
Atopobium minutum		21.95	0.28		1	
01014010	**Low discrimination**					**01014010**
					SPO	
Clostridium difficile		52.12	0.49	MAN 90	99	
Clostridium glycolicum		20.43	0.43	XYL 5	99	
Pseudoramibacter alactolyticus		8.71	0.32		1	
01014011	**Acceptable identification**					**01014011**
Clostridium novyi biovar A		90.46	0.50	FRU 10		
Clostridium difficile		5.76	0.15	XYL 1		
Clostridium glycolicum		2.26	0.09			
01014012	**Low discrimination**					**01014012**
					SPO	
Clostridium novyi biovar A		34.80	0.37	FRU 10	99	
Pseudoramibacter alactolyticus		23.09	0.15	XYL 1	1	
Atopobium minutum		19.20	0.28		1	
01014013	**Good identification**					**01014013**
Clostridium novyi biovar A		100.0	0.37	FRU 10		
01014020	**Low discrimination**					**01014020**
Pseudoramibacter alactolyticus		56.99	0.53	XYL 1		
Atopobium minutum		14.91	0.57			
Actinomyces meyeri		12.20	0.52			
01014021	**Low discrimination**					**01014021**
					SPO	
Clostridium novyi biovar A		43.96	0.39	FRU 10	99	
Pseudoramibacter alactolyticus		32.71	0.19	XYL 1	1	
Atopobium minutum		8.56	0.23		1	
01014022	**Low discrimination**					**01014022**
Atopobium minutum		57.15	0.54	XYL 1		
Pseudoramibacter alactolyticus		36.40	0.37			
Actinomyces meyeri		2.17	0.26			
01014030	**Low discrimination**					**01014030**
					SPO	
Clostridium novyi biovar A		39.53	0.39	FRU 10	99	
Pseudoramibacter alactolyticus		29.41	0.19	XYL 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		10.07	0.15		99			
01014031	**Good identification**							**01014031**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01014040	**Excellent identification**							**01014040**
					COC	SPO	GRA	
Clostridium glycolicum		99.24	0.99		1	99	99	
Pseudoramibacter alactolyticus		0.56	0.56		1	1	99	
Megasphaera elsdenii		0.16	0.57		99	1	1	
01014041	**Very good identification**							**01014041**
					SPO			
Clostridium glycolicum		97.88	0.65	URE 1	99			
Clostridium novyi biovar A		1.39	0.47		99			
Pseudoramibacter alactolyticus		0.55	0.21		1			
01014042	**Good identification**							**01014042**
					COC	SPO	GRA	
Clostridium glycolicum		92.25	0.65	BGL 1	1	99	99	
Pseudoramibacter alactolyticus		5.69	0.39		1	1	99	
Megasphaera elsdenii		1.60	0.41		99	1	1	
01014050	**Very good identification**							**01014050**
Clostridium glycolicum		97.21	0.65	MLZ 1				
Clostridium novyi biovar A		1.38	0.47					
Clostridium difficile		0.69	0.27					
01014051	**Low discrimination**							**01014051**
Clostridium novyi biovar A		58.43	0.47	FRU 10				
Clostridium glycolicum		41.57	0.31	XYL 1				
01014060	**Good identification**							**01014060**
					SPO			
Clostridium glycolicum		90.38	0.65	NAG 1	99			
Pseudoramibacter alactolyticus		8.86	0.43		1			
Clostridium novyi biovar A		0.32	0.36		99			
01014071	**Good identification**							**01014071**
Clostridium novyi biovar A		97.21	0.36	FRU 10				
Clostridium glycolicum		2.79	-.03	XYL 1				
01014100	**Very good identification**							**01014100**
					SPO			
Actinomyces meyeri		95.67	0.77	SUC 93	1			
Clostridium glycolicum		2.05	0.43		99			
Pseudoramibacter alactolyticus		0.87	0.32		1			
01014101	**Good identification**							**01014101**
					SPO			
Actinomyces meyeri		95.27	0.43	SUC 93	1			
Clostridium glycolicum		2.04	0.09	URE 1	99			
Pseudoramibacter alactolyticus		0.87	-.03		1			
01014102	**Low discrimination**							**01014102**
Actinomyces meyeri		89.16	0.51	SUC 93				
Pseudoramibacter alactolyticus		2.92	0.15	BGL 3				
Lactobacillus fermentum		2.74	0.36					

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01014110	**Acceptable identification**							**01014110**
					SPO			
Actinomyces meyeri		90.57	0.43	SUC 93	1			
Clostridium difficile		4.94	0.15	MLZ 1	99			
Clostridium glycolicum		1.94	0.09		99			
01014120	**Very good identification**							**01014120**
Actinomyces meyeri		98.00	0.62	SUC 93				
Pseudoramibacter alactolyticus		1.16	0.19	NAG 12				
Atopobium minutum		0.30	0.23					
01014122	**Low discrimination**							**01014122**
Actinomyces meyeri		84.46	0.36	SUC 93				
Lactobacillus fermentum		6.34	0.28	NAG 12				
Atopobium minutum		5.62	0.20	BGL 3				
01014140	**Very good identification**							**01014140**
					SPO			
Clostridium glycolicum		96.85	0.65	LAC 1	99			
Actinomyces meyeri		2.41	0.43		1			
Pseudoramibacter alactolyticus		0.54	0.21		1			
01014200	**Low discrimination**							**01014200**
					GRA	SPO		
Clostridium glycolicum		51.31	0.43	RHA 1	99	99		
Pseudoramibacter alactolyticus		21.88	0.32	SOR 95	99	1		
Fusobacterium varium		12.68	0.28		1	1		
01014240	**Very good identification**							**01014240**
					COC	SPO	GRA	
Clostridium glycolicum		99.25	0.65	RHA 1	1	99	99	
Pseudoramibacter alactolyticus		0.56	0.21		1	1	99	
Megasphaera elsdenii		0.16	0.23		99	1	1	
01014300	**Good identification**							**01014300**
					SPO			
Actinomyces meyeri		96.06	0.43	SUC 93	1			
Clostridium glycolicum		2.05	0.09	RHA 1	99			
Pseudoramibacter alactolyticus		0.88	-.03		1			
01014400	**Low discrimination**							**01014400**
					SPO			
Actinomyces meyeri		60.84	0.56	SUC 93	1			
Clostridium glycolicum		20.81	0.43		99			
Pseudoramibacter alactolyticus		8.87	0.32		1			
01014420	**Low discrimination**							**01014420**
Actinomyces meyeri		76.16	0.41	SUC 93				
Pseudoramibacter alactolyticus		14.37	0.19	NAG 12				
Atopobium minutum		3.76	0.23					
01014440	**Very good identification**							**01014440**
					COC	SPO	GRA	
Clostridium glycolicum		99.10	0.65	ARA 1	1	99	99	
Pseudoramibacter alactolyticus		0.56	0.21		1	1	99	
Megasphaera elsdenii		0.16	0.23		99	1	1	
01014500	**Very good identification**							**01014500**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		99.46	0.67	SUC 93	1	
Lactobacillus fermentum		0.38	0.36		1	
Clostridium glycolicum		0.09	0.09		99	
01014502	**Low discrimination**					**01014502**
Actinomyces meyeri		89.05	0.41	SUC 93		
Lactobacillus fermentum		10.95	0.36	BGL 3		
01014520	**Good identification**					**01014520**
Actinomyces meyeri		99.08	0.52	SUC 93		
Lactobacillus fermentum		0.92	0.28	NAG 12		
01015000	**Low discrimination**					**01015000**
					SPO	
Actinomyces meyeri		51.17	0.70	SUC 93	1	
Clostridium glycolicum		32.08	0.61		99	
Lactobacillus fermentum		7.87	0.67		1	
01015001	**Low discrimination**					**01015001**
					SPO	
Actinomyces meyeri		44.99	0.35	SUC 93	1	
Clostridium glycolicum		28.20	0.27	URE 1	99	
Clostridium novyi biovar A		11.41	0.33		99	
01015002	**Low discrimination**					**01015002**
Lactobacillus fermentum		57.02	0.67	SUC 87		
Atopobium minutum		27.47	0.54			
Actinomyces meyeri		11.47	0.44			
01015010	**Low discrimination**					**01015010**
					SPO	
Actinomyces meyeri		42.93	0.35	SUC 93	1	
Clostridium glycolicum		26.91	0.27	MLZ 1	99	
Clostridium novyi biovar A		10.89	0.33		99	
01015020	**Low discrimination**					**01015020**
Actinomyces meyeri		56.41	0.55	SUC 93		
Lactobacillus fermentum		21.21	0.58	NAG 12		
Atopobium minutum		15.32	0.49			
01015022	**Low discrimination**					**01015022**
Lactobacillus fermentum		63.23	0.58	SUC 87		
Atopobium minutum		30.46	0.46			
Actinomyces meyeri		5.20	0.29			
01015040	**Very good identification**					**01015040**
					SPO	
Clostridium glycolicum		99.81	0.83	GAL 10	99	
Actinomyces meyeri		0.09	0.35		1	
Pseudoramibacter alactolyticus		0.05	0.21		1	
01015041	**Good identification**					**01015041**
Clostridium glycolicum		98.60	0.49	GAL 10		
Clostridium novyi biovar A		1.40	0.30	URE 1		
01015042	**Good identification**					**01015042**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium glycolicum		96.95	0.49	GAL 10	99	
Lactobacillus fermentum		1.25	0.32	BGL 1	1	
Atopobium minutum		0.60	0.20		1	
01015050	**Good identification**					**01015050**
Clostridium glycolicum		98.60	0.49	GAL 10		
Clostridium novyi biovar A		1.40	0.30	MLZ 1		
01015060	**Good identification**					**01015060**
					SPO	
Clostridium glycolicum		96.19	0.49	GAL 10	99	
Actinomyces meyeri		1.10	0.21	NAG 1	1	
Pseudoramibacter alactolyticus		0.86	0.09		1	
01015100	**Good identification**					**01015100**
					GRA	
Actinomyces meyeri		93.47	0.80	SUC 93	99	
Lactobacillus fermentum		5.86	0.70		99	
Fusobacterium mortiferum		0.49	0.50		1	
01015101	**Acceptable identification**					**01015101**
					GRA	
Actinomyces meyeri		93.64	0.46	SUC 93	99	
Lactobacillus fermentum		5.87	0.36	URE 1	99	
Fusobacterium mortiferum		0.49	0.16		1	
01015102	**Low discrimination**					**01015102**
					GRA	
Lactobacillus fermentum		66.43	0.70	SUC 87	99	
Actinomyces meyeri		32.76	0.54		99	
Fusobacterium mortiferum		0.61	0.34		1	
01015103	**Low discrimination**					**01015103**
Lactobacillus fermentum		66.98	0.36	SUC 87		
Actinomyces meyeri		33.03	0.20	URE 1		
01015110	**Acceptable identification**					**01015110**
					GRA	
Actinomyces meyeri		93.64	0.46	SUC 93	99	
Lactobacillus fermentum		5.87	0.36	MLZ 1	99	
Fusobacterium mortiferum		0.49	0.16		1	
01015112	**Low discrimination**					**01015112**
Lactobacillus fermentum		66.98	0.36	SUC 87		
Actinomyces meyeri		33.03	0.20	MLZ 1		
01015120	**Low discrimination**					**01015120**
Actinomyces meyeri		86.63	0.65	SUC 93		
Lactobacillus fermentum		13.28	0.62	NAG 12		
Atopobium minutum		0.06	0.15			
01015122	**Low discrimination**					**01015122**
Lactobacillus fermentum		83.01	0.62	SUC 87		
Actinomyces meyeri		16.74	0.39			
Atopobium minutum		0.25	0.12			
01015140	**Low discrimination**					**01015140**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium glycolicum		73.60	0.49	GAL 10	99	
Actinomyces meyeri		24.72	0.46	LAC 1	1	
Lactobacillus fermentum		1.55	0.36		1	
01015142	**Low discrimination**					**01015142**
					SPO	
Lactobacillus fermentum		50.69	0.36	SUC 87	1	
Actinomyces meyeri		25.00	0.20	SOR 1	1	
Clostridium glycolicum		24.31	0.15		99	
01015200	**Low discrimination**					**01015200**
					SPO	
Actinomyces meyeri		51.81	0.35	SUC 93	1	
Clostridium glycolicum		32.48	0.27	RHA 1	99	
Lactobacillus fermentum		7.97	0.32		1	
01015240	**Good identification**					**01015240**
Clostridium glycolicum		100.0	0.49	GAL 10		
01015300	**Acceptable identification**					**01015300**
					GRA	
Actinomyces meyeri		93.64	0.46	SUC 93	99	
Lactobacillus fermentum		5.87	0.36	RHA 1	99	
Fusobacterium mortiferum		0.49	0.16		1	
01015302	**Low discrimination**					**01015302**
Lactobacillus fermentum		66.98	0.36	SUC 87		
Actinomyces meyeri		33.03	0.20	RHA 1		
01015400	**Low discrimination**					**01015400**
					SPO	
Actinomyces meyeri		60.62	0.59	SUC 93	1	
Lactobacillus fermentum		37.29	0.67		1	
Clostridium glycolicum		1.54	0.27		99	
01015402	**Good identification**					**01015402**
Lactobacillus fermentum		94.77	0.67	SUC 87		
Actinomyces meyeri		4.77	0.33			
Atopobium minutum		0.46	0.20			
01015420	**Low discrimination**					**01015420**
Lactobacillus fermentum		59.80	0.58	SUC 87		
Actinomyces meyeri		39.77	0.45			
Atopobium minutum		0.44	0.15			
01015422	**Very good identification**					**01015422**
Lactobacillus fermentum		97.52	0.58	SUC 87		
Actinomyces meyeri		2.01	0.19			
Atopobium minutum		0.48	0.12			
01015440	**Good identification**					**01015440**
					SPO	
Clostridium glycolicum		96.72	0.49	GAL 10	99	
Actinomyces meyeri		2.03	0.25	ARA 1	1	
Lactobacillus fermentum		1.25	0.32		1	
01015500	**Low discrimination**					**01015500**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		79.93	0.70	SUC 93	99	
Lactobacillus fermentum		20.05	0.70		99	
Fusobacterium mortiferum		0.02	0.16		1	
01015501	**Low discrimination**					**01015501**
Actinomyces meyeri		79.94	0.35	SUC 93		
Lactobacillus fermentum		20.06	0.36	URE 1		
01015502	**Low discrimination**					**01015502**
Lactobacillus fermentum		89.03	0.70	SUC 87		
Actinomyces meyeri		10.97	0.44			
01015503	**Low discrimination**					**01015503**
Lactobacillus fermentum		89.03	0.36	SUC 87		
Actinomyces meyeri		10.98	0.10	URE 1		
01015510	**Low discrimination**					**01015510**
Actinomyces meyeri		79.94	0.35	SUC 93		
Lactobacillus fermentum		20.06	0.36	MLZ 1		
01015512	**Low discrimination**					**01015512**
Lactobacillus fermentum		89.03	0.36	SUC 87		
Actinomyces meyeri		10.98	0.10	MLZ 1		
01015520	**Low discrimination**					**01015520**
Actinomyces meyeri		61.99	0.55	SUC 93		
Lactobacillus fermentum		38.02	0.62	NAG 12		
01015522	**Very good identification**					**01015522**
Lactobacillus fermentum		95.20	0.62	SUC 87		
Actinomyces meyeri		4.80	0.29			
01015540	**Low discrimination**					**01015540**
					SPO	
Actinomyces meyeri		72.93	0.35	SUC 93	1	
Lactobacillus fermentum		18.30	0.36	SOR 1	1	
Clostridium glycolicum		8.78	0.15		99	
01015542	**Low discrimination**					**01015542**
Lactobacillus fermentum		89.03	0.36	SUC 87		
Actinomyces meyeri		10.98	0.10	SOR 1		
01015700	**Low discrimination**					**01015700**
Actinomyces meyeri		79.94	0.35	SUC 93		
Lactobacillus fermentum		20.06	0.36	RHA 1		
01015702	**Low discrimination**					**01015702**
Lactobacillus fermentum		89.03	0.36	SUC 87		
Actinomyces meyeri		10.98	0.10	RHA 1		
01016000	**Low discrimination**					**01016000**
					SPO	
Pseudoramibacter alactolyticus		47.57	0.61	XYL 1	1	
Eubacterium limosum		30.26	0.60		1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		9.84	0.53		99			
01016002	**Low discrimination**							**01016002**
					COC	GRA		
Pseudoramibacter alactolyticus		77.27	0.45	XYL 1	1	99		
Megasphaera elsdenii		15.11	0.44	BGL 10	99	1		
Eubacterium limosum		4.47	0.26		1	99		
01016010	**Very good identification**							**01016010**
					SPO			
Clostridium difficile		98.13	0.65	XYL 5	99			
Pseudoramibacter alactolyticus		0.98	0.27		1			
Eubacterium limosum		0.62	0.26		1			
01016020	**Acceptable identification**							**01016020**
					SPO			
Pseudoramibacter alactolyticus		92.25	0.48	XYL 1	1			
Eubacterium limosum		3.36	0.26		1			
Clostridium difficile		1.09	0.19		99			
01016040	**Low discrimination**							**01016040**
					COC	SPO	GRA	
Clostridium glycolicum		67.74	0.65	MAN 1	1	99	99	
Pseudoramibacter alactolyticus		20.27	0.51		1	1	99	
Megasphaera elsdenii		10.56	0.57		99	1	1	
01016050	**Low discrimination**							**01016050**
					SPO			
Clostridium difficile		81.08	0.43	XYL 5	99			
Clostridium glycolicum		12.88	0.31	SOR 5	99			
Pseudoramibacter alactolyticus		3.85	0.17		1			
01016060	**Low discrimination**							**01016060**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		81.07	0.38	XYL 1	1	1	99	
Clostridium glycolicum		15.51	0.31		1	99	99	
Megasphaera elsdenii		2.42	0.23		99	1	1	
01016100	**Low discrimination**							**01016100**
Actinomyces meyeri		49.45	0.43	SUC 93				
Pseudoramibacter alactolyticus		24.03	0.27	MAN 1				
Eubacterium limosum		15.28	0.26					
01017040	**Good identification**							**01017040**
					COC	SPO	GRA	
Clostridium glycolicum		95.92	0.49	GAL 10	1	99	99	
Pseudoramibacter alactolyticus		2.61	0.17	MAN 1	1	1	99	
Megasphaera elsdenii		1.36	0.23		99	1	1	
01017100	**Acceptable identification**							**01017100**
					GRA			
Actinomyces meyeri		93.35	0.46	SUC 93	99			
Lactobacillus fermentum		5.86	0.36	MAN 1	99			
Fusobacterium mortiferum		0.49	0.16		1			
01017102	**Low discrimination**							**01017102**
Lactobacillus fermentum		66.98	0.36	SUC 87				
Actinomyces meyeri		33.03	0.20	MAN 1				
01017500	**Low discrimination**							**01017500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		79.94	0.35	SUC 93			
Lactobacillus fermentum		20.06	0.36	MAN 1			
01017502	**Low discrimination**						**01017502**
Lactobacillus fermentum		89.03	0.36	SUC 87			
Actinomyces meyeri		10.98	0.10	MAN 1			
01020000	**Low discrimination**						**01020000**
					COC	SPO	
Clostridium botulinum biovar A		64.92	0.85		1	99	
Clostridium sporogenes		20.86	0.74		1	99	
Peptostreptococcus anaerobius		9.11	0.66		99	1	
01020001	**Low discrimination**						**01020001**
Clostridium novyi biovar A		44.48	0.66	TRE 1			
Clostridium botulinum biovar A		29.28	0.51				
Clostridium sporogenes		9.41	0.40				
01020002	**Low discrimination**						**01020002**
					SPO		
Clostridium sporogenes		75.92	0.66		99		
Atopobium minutum		12.62	0.63		1		
Clostridium botulinum biovar A		7.16	0.51		99		
01020003	**Low discrimination**						**01020003**
					SPO		
Clostridium novyi biovar A		62.87	0.53	TRE 1	99		
Clostridium sporogenes		25.12	0.32		99		
Atopobium minutum		4.17	0.29		1		
01020010	**Low discrimination**						**01020010**
Clostridium novyi biovar A		49.68	0.66	TRE 1			
Clostridium botulinum biovar A		32.70	0.51				
Clostridium sporogenes		10.51	0.40				
01020011	**Very good identification**						**01020011**
Clostridium novyi biovar A		98.86	0.66	TRE 1			
Clostridium botulinum biovar A		0.66	0.17				
Clostridium sporogenes		0.21	0.06				
01020012	**Low discrimination**						**01020012**
					SPO		
Clostridium novyi biovar A		66.28	0.53	TRE 1	99		
Clostridium sporogenes		26.48	0.32		99		
Atopobium minutum		4.40	0.29		1		
01020013	**Good identification**						**01020013**
Clostridium novyi biovar A		99.60	0.53	TRE 1			
Clostridium sporogenes		0.40	-.02				
01020020	**Low discrimination**						**01020020**
					SPO		
Clostridium cadaveris		28.68	0.49	IND 98	99		
Clostridium botulinum biovar A		25.21	0.51	TRE 5	99		
Atopobium minutum		22.22	0.58		1		
01020021	**Acceptable identification**						**01020021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		90.21	0.56	TRE 1			
Clostridium cadaveris		2.73	0.15				
Clostridium botulinum biovar A		2.40	0.17				
01020022	**Low discrimination**						**01020022**
					GRA	SPO	
Atopobium minutum		60.36	0.55	TRE 1	99	1	
Pseudoflavonifractor capillosus		18.42	0.32		1	1	
Clostridium sporogenes		11.00	0.32		99	99	
01020023	**Low discrimination**						**01020023**
					GRA	SPO	
Clostridium novyi biovar A		88.36	0.43	TRE 1	99	99	
Atopobium minutum		7.82	0.21		99	1	
Pseudoflavonifractor capillosus		2.39	-.03		1	1	
01020030	**Acceptable identification**						**01020030**
Clostridium novyi biovar A		91.53	0.56	TRE 1			
Clostridium cadaveris		2.77	0.15				
Clostridium botulinum biovar A		2.43	0.17				
01020031	**Good identification**						**01020031**
Clostridium novyi biovar A		100.0	0.56	TRE 1			
01020032	**Low discrimination**						**01020032**
					GRA	SPO	
Clostridium novyi biovar A		88.36	0.43	TRE 1	99	99	
Atopobium minutum		7.82	0.21		99	1	
Pseudoflavonifractor capillosus		2.39	-.03		1	1	
01020033	**Good identification**						**01020033**
Clostridium novyi biovar A		100.0	0.43	TRE 1			
01020040	**Good identification**						**01020040**
Clostridium botulinum biovar A		94.57	0.76				
Clostridium novyi biovar A		3.24	0.63				
Clostridium sporogenes		1.03	0.40				
01020041	**Low discrimination**						**01020041**
Clostridium novyi biovar A		76.79	0.63	TRE 1			
Clostridium botulinum biovar A		22.64	0.42				
Clostridium sporogenes		0.25	0.06				
01020042	**Low discrimination**						**01020042**
Clostridium botulinum biovar A		48.49	0.42	BGL 1			
Clostridium novyi biovar A		29.02	0.50				
Clostridium sporogenes		17.39	0.32				
01020043	**Good identification**						**01020043**
Clostridium novyi biovar A		97.76	0.50	TRE 1			
Clostridium botulinum biovar A		1.65	0.08				
Clostridium sporogenes		0.59	-.02				
01020050	**Low discrimination**						**01020050**
Clostridium novyi biovar A		76.82	0.63	TRE 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		22.65	0.42				
Clostridium sporogenes		0.25	0.06				
01020051	**Very good identification**						**01020051**
Clostridium novyi biovar A		99.70	0.63	TRE 1			
Clostridium botulinum biovar A		0.30	0.08				
01020052	**Good identification**						**01020052**
Clostridium novyi biovar A		97.76	0.50	TRE 1			
Clostridium botulinum biovar A		1.65	0.08				
Clostridium sporogenes		0.59	-.02				
01020053	**Good identification**						**01020053**
Clostridium novyi biovar A		100.0	0.50	TRE 1			
01020060	**Low discrimination**						**01020060**
Clostridium botulinum biovar A		50.94	0.42	NAG 1			
Clostridium novyi biovar A		43.19	0.53				
Clostridium cadaveris		1.96	0.15				
01020061	**Good identification**						**01020061**
Clostridium novyi biovar A		98.82	0.53	TRE 1			
Clostridium botulinum biovar A		1.18	0.08				
01020062	**Low discrimination**						**01020062**
					SPO		
Clostridium novyi biovar A		77.85	0.40	TRE 1	99		
Atopobium minutum		10.34	0.21		1		
Clostridium botulinum biovar A		5.26	0.08		99		
01020063	**Good identification**						**01020063**
Clostridium novyi biovar A		100.0	0.40	TRE 1			
01020070	**Good identification**						**01020070**
Clostridium novyi biovar A		98.82	0.53	TRE 1			
Clostridium botulinum biovar A		1.18	0.08				
01020071	**Good identification**						**01020071**
Clostridium novyi biovar A		100.0	0.53	TRE 1			
01020072	**Good identification**						**01020072**
Clostridium novyi biovar A		100.0	0.40	TRE 1			
01020073	**Good identification**						**01020073**
Clostridium novyi biovar A		100.0	0.40	TRE 1			
01020100	**Low discrimination**						**01020100**
					COC	SPO	
Clostridium botulinum biovar A		64.59	0.51	LAC 1	1	99	
Clostridium sporogenes		20.76	0.40		1	99	
Peptostreptococcus anaerobius		9.06	0.32		99	1	
01020140	**Good identification**						**01020140**
Clostridium botulinum biovar A		95.25	0.42	LAC 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		3.26	0.29				
Clostridium sporogenes		1.04	0.06				
01020200	**Low discrimination**						**01020200**
					COC	SPO	
Clostridium botulinum biovar A		64.98	0.51	RHA 1	1	99	
Clostridium sporogenes		20.88	0.40		1	99	
Peptostreptococcus anaerobius		9.12	0.32		99	1	
01020240	**Good identification**						**01020240**
Clostridium botulinum biovar A		95.25	0.42	RHA 1			
Clostridium novyi biovar A		3.26	0.29				
Clostridium sporogenes		1.04	0.06				
01020400	**Low discrimination**						**01020400**
					COC	SPO	
Clostridium botulinum biovar A		64.95	0.51	ARA 1	1	99	
Clostridium sporogenes		20.87	0.40		1	99	
Peptostreptococcus anaerobius		9.11	0.32		99	1	
01020440	**Good identification**						**01020440**
Clostridium botulinum biovar A		95.25	0.42	ARA 1			
Clostridium novyi biovar A		3.26	0.29				
Clostridium sporogenes		1.04	0.06				
01021000	**Low discrimination**						**01021000**
					SPO		
Clostridium botulinum biovar A		38.71	0.51	GAL 1	99		
Atopobium minutum		34.12	0.58		1		
Clostridium sporogenes		12.44	0.40		99		
01021001	**Low discrimination**						**01021001**
					SPO		
Clostridium novyi biovar A		85.21	0.50	TRE 1	99		
Clostridium botulinum biovar A		5.10	0.17	GAL 10	99		
Atopobium minutum		4.49	0.24		1		
01021002	**Low discrimination**						**01021002**
					SPO		
Atopobium minutum		79.83	0.55	TRE 1	1		
Clostridium sporogenes		14.56	0.32		99		
Clostridium novyi biovar A		4.05	0.37		99		
01021003	**Low discrimination**						**01021003**
					SPO		
Clostridium novyi biovar A		80.94	0.37	TRE 1	99		
Atopobium minutum		16.12	0.21	GAL 10	1		
Clostridium sporogenes		2.94	-.02		99		
01021010	**Low discrimination**						**01021010**
					SPO		
Clostridium novyi biovar A		87.71	0.50	TRE 1	99		
Clostridium botulinum biovar A		5.25	0.17	GAL 10	99		
Atopobium minutum		4.62	0.24		1		
01021011	**Good identification**						**01021011**
Clostridium novyi biovar A		100.0	0.50	TRE 1			
01021012	**Low discrimination**						**01021012**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		80.94	0.37	TRE 1	99		
Atopobium minutum		16.12	0.21	GAL 10	1		
Clostridium sporogenes		2.94	-.02		99		
01021013	**Good identification**						**01021013**
Clostridium novyi biovar A		100.0	0.37	TRE 1			
01021020	**Low discrimination**						**01021020**
					SPO		
Atopobium minutum		81.10	0.50	TRE 1	1		
Clostridium novyi biovar A		11.65	0.39		99		
Clostridium cadaveris		3.17	0.15		99		
01021021	**Acceptable identification**						**01021021**
					SPO		
Clostridium novyi biovar A		93.43	0.39	TRE 1	99		
Atopobium minutum		6.57	0.15	GAL 10	1		
01021022	**Low discrimination**						**01021022**
					GRA	SPO	
Atopobium minutum		87.31	0.47	TRE 1	99	1	
Pseudoflavonifractor capillosus		8.88	0.15		1	1	
Clostridium novyi biovar A		3.32	0.26		99	99	
01021030	**Acceptable identification**						**01021030**
					SPO		
Clostridium novyi biovar A		93.43	0.39	TRE 1	99		
Atopobium minutum		6.57	0.15	GAL 10	1		
01021031	**Good identification**						**01021031**
Clostridium novyi biovar A		100.0	0.39	TRE 1			
01021040	**Low discrimination**						**01021040**
					SPO		
Clostridium botulinum biovar A		69.47	0.42	GAL 1	99		
Clostridium novyi biovar A		26.18	0.47		99		
Atopobium minutum		2.07	0.24		1		
01021041	**Good identification**						**01021041**
Clostridium novyi biovar A		97.39	0.47	TRE 1			
Clostridium botulinum biovar A		2.61	0.08	GAL 10			
01021050	**Good identification**						**01021050**
Clostridium novyi biovar A		97.39	0.47	TRE 1			
Clostridium botulinum biovar A		2.61	0.08	GAL 10			
01021051	**Good identification**						**01021051**
Clostridium novyi biovar A		100.0	0.47	TRE 1			
01021060	**Low discrimination**						**01021060**
					SPO		
Clostridium novyi biovar A		82.46	0.36	TRE 1	99		
Clostridium botulinum biovar A		8.84	0.08	GAL 10	99		
Atopobium minutum		8.70	0.15		1		
01021061	**Good identification**						**01021061**
Clostridium novyi biovar A		100.0	0.36	TRE 1			
01021070	**Good identification**						**01021070**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.36	TRE 1				
01021071	**Good identification**							**01021071**
Clostridium novyi biovar A		100.0	0.36	TRE 1				
01022000	**Low discrimination**							**01022000**
					COC	SPO		
Clostridium botulinum biovar A		54.96	0.51	MAN 1	1	99		
Clostridium sporogenes		17.66	0.40		1	99		
Peptostreptococcus anaerobius		7.71	0.32		99	1		
01022040	**Low discrimination**							**01022040**
					COC	SPO	GRA	
Clostridium botulinum biovar A		73.92	0.42	MAN 1	1	99	99	
Megasphaera elsdenii		16.10	0.35		99	1	1	
Pseudoramibacter alactolyticus		5.93	0.17		1	1	99	
01024000	**Low discrimination**							**01024000**
					COC	SPO		
Clostridium botulinum biovar A		62.91	0.51	XYL 1	1	99		
Clostridium sporogenes		20.22	0.40		1	99		
Peptostreptococcus anaerobius		8.83	0.32		99	1		
01024040	**Low discrimination**							**01024040**
Clostridium glycolicum		73.75	0.49	FRU 90				
Clostridium botulinum biovar A		25.01	0.42	TRE 1				
Clostridium novyi biovar A		0.86	0.29					
01030000	**Low discrimination**							**01030000**
					SPO			
Pseudoramibacter alactolyticus		45.79	0.66	TRE 1	1			
Clostridium botulinum biovar A		16.42	0.65		99			
Clostridium sporogenes		14.59	0.61		99			
01030002	**Low discrimination**							**01030002**
					SPO			
Pseudoramibacter alactolyticus		33.48	0.50	TRE 1	1			
Clostridium sporogenes		32.01	0.53	BGL 10	99			
Atopobium minutum		27.83	0.62		1			
01030003	**Low discrimination**							**01030003**
					SPO			
Clostridium novyi biovar A		34.25	0.37	FRU 10	99			
Pseudoramibacter alactolyticus		22.73	0.15	TRE 1	1			
Clostridium sporogenes		21.73	0.19		99			
01030010	**Low discrimination**							**01030010**
					SPO			
Clostridium difficile		72.89	0.51	TRE 7	99			
Pseudoramibacter alactolyticus		8.52	0.32	MAN 90	1			
Clostridium novyi biovar A		8.08	0.50		99			
01030011	**Acceptable identification**							**01030011**
					SPO			
Clostridium novyi biovar A		90.77	0.50	FRU 10	99			
Clostridium difficile		8.27	0.17	TRE 1	99			
Pseudoramibacter alactolyticus		0.97	-.03		1			
01030012	**Low discrimination**							**01030012**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		28.02	0.37	FRU 10	99			
Pseudoramibacter alactolyticus		18.59	0.15	TRE 1	1			
Clostridium sporogenes		17.78	0.19		99			
01030013	**Good identification**							**01030013**
Clostridium novyi biovar A		100.0	0.37	FRU 10				
01030020	**Low discrimination**							**01030020**
					SPO			
Pseudoramibacter alactolyticus		58.09	0.53	TRE 1	1			
Clostridium cadaveris		21.26	0.49		99			
Atopobium minutum		15.20	0.57		1			
01030021	**Low discrimination**							**01030021**
					SPO			
Clostridium novyi biovar A		45.23	0.39	FRU 10	99			
Pseudoramibacter alactolyticus		33.65	0.19	TRE 1	1			
Clostridium cadaveris		12.32	0.15		99			
01030022	**Low discrimination**							**01030022**
					GRA			
Atopobium minutum		56.80	0.54	TRE 1	99			
Pseudoramibacter alactolyticus		36.17	0.37		99			
Pseudoflavonifractor capillosus		2.09	0.15		1			
01030030	**Low discrimination**							**01030030**
					SPO			
Clostridium novyi biovar A		37.46	0.39	FRU 10	99			
Pseudoramibacter alactolyticus		27.87	0.19	TRE 1	1			
Clostridium difficile		13.65	0.17		99			
01030031	**Good identification**							**01030031**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01030040	**Low discrimination**							**01030040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		53.65	0.56	TRE 1	1	1	99	
Clostridium botulinum biovar A		23.00	0.56		1	99	99	
Megasphaera elsdenii		15.06	0.57		99	1	1	
01030041	**Low discrimination**							**01030041**
					SPO			
Clostridium novyi biovar A		58.38	0.47	FRU 10	99			
Pseudoramibacter alactolyticus		23.07	0.21	TRE 1	1			
Clostridium botulinum biovar A		9.89	0.22		99			
01030042	**Low discrimination**							**01030042**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		69.39	0.39	TRE 1	1	1	99	
Megasphaera elsdenii		19.48	0.41	BGL 10	99	1	1	
Clostridium novyi biovar A		2.82	0.34		1	99	99	
01030050	**Low discrimination**							**01030050**
					SPO			
Clostridium novyi biovar A		40.35	0.47	FRU 10	99			
Clostridium difficile		28.73	0.29	TRE 1	99			
Pseudoramibacter alactolyticus		15.95	0.21		1			
01030051	**Good identification**							**01030051**
Clostridium novyi biovar A		99.29	0.47	FRU 10				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		0.71	-.05	TRE 1			
01030060	**Low discrimination**						**01030060**
					SPO		
Pseudoramibacter alactolyticus		88.91	0.43	TRE 1	1		
Clostridium novyi biovar A		3.22	0.36		99		
Clostridium botulinum biovar A		2.18	0.22		99		
01030061	**Low discrimination**						**01030061**
					SPO		
Clostridium novyi biovar A		78.19	0.36	FRU 10	99		
Pseudoramibacter alactolyticus		21.81	0.09	TRE 1	1		
01030070	**Low discrimination**						**01030070**
					SPO		
Clostridium novyi biovar A		70.79	0.36	FRU 10	99		
Pseudoramibacter alactolyticus		19.75	0.09	TRE 1	1		
Clostridium botulinum biovar B		7.43	0.05		99		
01030071	**Good identification**						**01030071**
Clostridium novyi biovar A		100.0	0.36	FRU 10			
01030100	**Low discrimination**						**01030100**
					GRA		
Fusobacterium mortiferum		45.27	0.57		1		
Pseudoramibacter alactolyticus		19.98	0.32		99		
Actinomyces meyeri		11.41	0.38		99		
01030102	**Low discrimination**						**01030102**
					GRA	SPO	
Fusobacterium mortiferum		42.61	0.41	BGL 10	1	1	
Pseudoramibacter alactolyticus		18.81	0.15		99	1	
Clostridium sporogenes		17.98	0.19		99	99	
01031000	**Low discrimination**						**01031000**
					GRA		
Atopobium minutum		49.83	0.57	TRE 1	99		
Fusobacterium mortiferum		15.87	0.54		1		
Pseudoramibacter alactolyticus		10.90	0.32		99		
01031002	**Low discrimination**						**01031002**
					GRA	CAT	
Atopobium minutum		81.56	0.54	TRE 1	99	1	
Propionibacterium granulosum		4.39	0.35		99	90	
Fusobacterium mortiferum		4.33	0.38		1	5	
01031020	**Low discrimination**						**01031020**
					CAT		
Atopobium minutum		68.57	0.49	TRE 1	1		
Propionibacterium acnes		8.42	0.26		95		
Pseudoramibacter alactolyticus		7.94	0.19		1		
01031022	**Low discrimination**						**01031022**
					CAT		
Atopobium minutum		89.91	0.46	TRE 1	1		
Propionibacterium granulosum		3.63	0.24		90		
Lactobacillus fermentum		3.21	0.28		1		
01031100	**Low discrimination**						**01031100**
					GRA		
Fusobacterium mortiferum		75.60	0.60		1		
Actinomyces meyeri		19.05	0.41		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		3.95	0.40		99			
01031102	**Low discrimination**							**01031102**
					GRA			
Fusobacterium mortiferum		61.66	0.44	BGL 10	1			
Lactobacillus fermentum		28.99	0.40		99			
Atopobium minutum		5.03	0.20		99			
01031400	**Low discrimination**							**01031400**
Lactobacillus fermentum		45.35	0.36	SUC 87				
Actinomyces meyeri		22.31	0.20	TRE 1				
Atopobium minutum		19.25	0.23					
01031402	**Low discrimination**							**01031402**
Lactobacillus fermentum		77.95	0.36	SUC 87				
Atopobium minutum		22.05	0.20	TRE 1				
01031502	**Good identification**							**01031502**
Lactobacillus fermentum		96.41	0.40	SUC 87				
Actinomyces meyeri		3.60	0.05	TRE 1				
01032000	**Low discrimination**							**01032000**
					SPO			
Pseudoramibacter alactolyticus		46.40	0.61	TRE 1	1			
Eubacterium limosum		29.51	0.60		1			
Clostridium difficile		13.73	0.56		99			
01032002	**Low discrimination**							**01032002**
					COC	GRA		
Pseudoramibacter alactolyticus		74.93	0.45	TRE 1	1	99		
Megasphaera elsdenii		14.65	0.44	BGL 10	99	1		
Eubacterium limosum		4.33	0.26		1	99		
01032010	**Very good identification**							**01032010**
					SPO			
Clostridium difficile		98.64	0.68	TRE 7	99			
Pseudoramibacter alactolyticus		0.69	0.27		1			
Eubacterium limosum		0.44	0.26		1			
01032020	**Acceptable identification**							**01032020**
					SPO			
Pseudoramibacter alactolyticus		90.86	0.48	TRE 1	1			
Eubacterium limosum		3.31	0.26		1			
Clostridium difficile		1.54	0.22		99			
01032040	**Low discrimination**							**01032040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		61.73	0.51	TRE 1	1	1	99	
Megasphaera elsdenii		32.18	0.57		99	1	1	
Clostridium difficile		3.85	0.34		1	99	99	
01032050	**Good identification**							**01032050**
					COC	SPO	GRA	
Clostridium difficile		95.05	0.46	TRE 7	1	99	99	
Pseudoramibacter alactolyticus		3.16	0.17	SOR 5	1	1	99	
Megasphaera elsdenii		1.65	0.23		99	1	1	
01032060	**Good identification**							**01032060**
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		95.86	0.38	TRE 1	1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		2.86	0.23		99	1	1	
Propionibacterium acnes		0.94	0.11		1	95	99	
01034000	**Low discrimination**							**01034000**
					GRA	SPO		
Clostridium glycolicum		43.56	0.43	TRE 1	99	99		
Pseudoramibacter alactolyticus		18.58	0.32	SOR 95	99	1		
Fusobacterium varium		10.77	0.28		1	1		
01034040	**Very good identification**							**01034040**
					SPO			
Clostridium glycolicum		99.01	0.65	TRE 1	99			
Pseudoramibacter alactolyticus		0.56	0.21		1			
Clostridium botulinum biovar A		0.24	0.22		99			
01034100	**Low discrimination**							**01034100**
					GRA	SPO		
Actinomyces meyeri		84.06	0.43	SUC 93	99	1		
Fusobacterium mortiferum		12.94	0.38	TRE 1	1	1		
Clostridium glycolicum		1.80	0.09		99	99		
01035000	**Low discrimination**							**01035000**
					GRA	SPO		
Actinomyces meyeri		45.25	0.35	SUC 93	99	1		
Clostridium glycolicum		28.37	0.27	TRE 1	99	99		
Fusobacterium mortiferum		11.94	0.35		1	1		
01035040	**Good identification**							**01035040**
Clostridium glycolicum		100.0	0.49	TRE 1				
01035100	**Low discrimination**							**01035100**
					GRA			
Actinomyces meyeri		82.20	0.46	SUC 93	99			
Fusobacterium mortiferum		12.65	0.41	TRE 1	1			
Lactobacillus fermentum		5.16	0.36		99			
01035102	**Low discrimination**							**01035102**
					GRA			
Lactobacillus fermentum		56.64	0.36	SUC 87	99			
Actinomyces meyeri		27.93	0.20	TRE 1	99			
Fusobacterium mortiferum		15.44	0.25		1			
01035500	**Low discrimination**							**01035500**
Actinomyces meyeri		79.94	0.35	SUC 93				
Lactobacillus fermentum		20.06	0.36	TRE 1				
01035502	**Low discrimination**							**01035502**
Lactobacillus fermentum		89.03	0.36	SUC 87				
Actinomyces meyeri		10.98	0.10	TRE 1				
01036010	**Good identification**							**01036010**
					SPO			
Clostridium difficile		99.87	0.46	TRE 7	99			
Pseudoramibacter alactolyticus		0.13	-.07	XYL 5	1			
01040000	**Low discrimination**							**01040000**
					COC	SPO		
Peptostreptococcus anaerobius		52.15	0.66	CEL 1	99	1		
Clostridium botulinum biovar A		15.02	0.61		1	99		
Atopobium minutum		9.93	0.66		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01040001	**Low discrimination**							**01040001**
					COC	GRA	SPO	
Clostridium novyi biovar A		70.85	0.66	CEL 1	1	99	99	
Campylobacter ureolyticus		12.79	0.32		1	1	1	
Peptostreptococcus anaerobius		6.54	0.32		99	99	1	
01040002	**Low discrimination**							**01040002**
					SPO			
Atopobium minutum		64.69	0.63	CEL 1	1			
Clostridium sporogenes		11.79	0.40		99			
Clostridium novyi biovar A		9.84	0.53		99			
01040003	**Low discrimination**							**01040003**
					COC	SPO		
Clostridium novyi biovar A		85.80	0.53	CEL 1	1	99		
Atopobium minutum		5.70	0.29		1	1		
Anaerococcus prevotii		4.44	0.13		99	1		
01040010	**Low discrimination**							**01040010**
					COC	SPO		
Clostridium novyi biovar A		85.24	0.66	CEL 1	1	99		
Peptostreptococcus anaerobius		7.87	0.32		99	1		
Clostridium botulinum biovar A		2.27	0.27		1	99		
01040011	**Very good identification**							**01040011**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.73	0.66	CEL 1	1	99	99	
Campylobacter ureolyticus		0.18	-.03		1	1	1	
Peptostreptococcus anaerobius		0.09	-.03		99	99	1	
01040012	**Acceptable identification**							**01040012**
					SPO			
Clostridium novyi biovar A		92.27	0.53	CEL 1	99			
Atopobium minutum		6.13	0.29		1			
Clostridium sporogenes		1.12	0.06		99			
01040013	**Good identification**							**01040013**
Clostridium novyi biovar A		100.0	0.53	CEL 1				
01040020	**Low discrimination**							**01040020**
					SPO			
Atopobium minutum		49.63	0.58	CEL 1	1			
Clostridium novyi biovar A		21.39	0.56		99			
Clostridium cadaveris		12.30	0.37		99			
01040021	**Good identification**							**01040021**
					COC	SPO		
Clostridium novyi biovar A		95.33	0.56	CEL 1	1	99		
Atopobium minutum		2.23	0.24		1	1		
Peptococcus niger		0.83	-.02		99	1		
01040022	**Low discrimination**							**01040022**
					GRA	SPO		
Atopobium minutum		69.11	0.55	CEL 1	99	1		
Pseudoflavonifractor capillosus		21.10	0.32		1	1		
Clostridium novyi biovar A		7.89	0.43		99	99		
01040023	**Low discrimination**							**01040023**
					GRA	SPO		
Clostridium novyi biovar A		89.64	0.43	CEL 1	99	99		
Atopobium minutum		7.94	0.21		99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01040030	**Good identification**						**01040030**
					SPO		
Clostridium novyi biovar A		96.81	0.56	CEL 1	99		
Atopobium minutum		2.27	0.24		1		
Clostridium cadaveris		0.56	0.03		99		
01040031	**Good identification**						**01040031**
Clostridium novyi biovar A		100.0	0.56	CEL 1			
01040032	**Low discrimination**						**01040032**
					GRA	SPO	
Clostridium novyi biovar A		89.64	0.43	CEL 1	99	99	
Atopobium minutum		7.94	0.21		99	1	
Pseudoflavonifractor capillosus		2.42	-.03		1	1	
01040033	**Good identification**						**01040033**
Clostridium novyi biovar A		100.0	0.43	CEL 1			
01040040	**Low discrimination**						**01040040**
					COC	SPO	
Clostridium botulinum biovar A		46.36	0.52	CEL 1	1	99	
Clostridium novyi biovar A		39.31	0.63		1	99	
Peptostreptococcus anaerobius		5.44	0.32		99	1	
01040041	**Very good identification**						**01040041**
					GRA	SPO	
Clostridium novyi biovar A		98.42	0.63	CEL 1	99	99	
Clostridium botulinum biovar A		1.17	0.18		99	99	
Campylobacter ureolyticus		0.27	-.03		1	1	
01040042	**Low discrimination**						**01040042**
					SPO		
Clostridium novyi biovar A		79.29	0.50	CEL 1	99		
Atopobium minutum		7.90	0.29		1		
Clostridium botulinum biovar A		5.35	0.18		99		
01040043	**Good identification**						**01040043**
Clostridium novyi biovar A		100.0	0.50	CEL 1			
01040050	**Very good identification**						**01040050**
					COC	SPO	
Clostridium novyi biovar A		98.69	0.63	CEL 1	1	99	
Clostridium botulinum biovar A		1.18	0.18		1	99	
Peptostreptococcus anaerobius		0.14	-.03		99	1	
01040051	**Very good identification**						**01040051**
Clostridium novyi biovar A		100.0	0.63	CEL 1			
01040052	**Good identification**						**01040052**
Clostridium novyi biovar A		100.0	0.50	CEL 1			
01040053	**Good identification**						**01040053**
Clostridium novyi biovar A		100.0	0.50	CEL 1			
01040060	**Low discrimination**						**01040060**
					SPO		
Clostridium novyi biovar A		88.64	0.53	CEL 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		4.22	0.18		99		
Atopobium minutum		3.12	0.24		1		
01040061	**Good identification**						**01040061**
Clostridium novyi biovar A		100.0	0.53	CEL 1			
01040062	**Low discrimination**						**01040062**
					GRA	SPO	
Clostridium novyi biovar A		83.83	0.40	CEL 1	99	99	
Atopobium minutum		11.13	0.21		99	1	
Pseudoflavonifractor capillosus		3.40	-.03		1	1	
01040063	**Good identification**						**01040063**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
01040070	**Good identification**						**01040070**
Clostridium novyi biovar A		100.0	0.53	CEL 1			
01040071	**Good identification**						**01040071**
Clostridium novyi biovar A		100.0	0.53	CEL 1			
01040072	**Good identification**						**01040072**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
01040073	**Good identification**						**01040073**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
01041000	**Low discrimination**						**01041000**
					COC	SPO	
Atopobium minutum		67.23	0.58	CEL 1	1	1	
Clostridium novyi biovar A		12.88	0.50		1	99	
Peptostreptococcus anaerobius		10.70	0.32		99	1	
01041001	**Acceptable identification**						**01041001**
					SPO		
Clostridium novyi biovar A		91.36	0.50	CEL 1	99		
Atopobium minutum		4.82	0.24	GAL 10	1		
Clostridium sordellii		1.56	0.03		99		
01041002	**Acceptable identification**						**01041002**
					SPO		
Atopobium minutum		94.46	0.55	CEL 1	1		
Clostridium novyi biovar A		4.79	0.37		99		
Clostridium sporogenes		0.52	0.06		99		
01041003	**Low discrimination**						**01041003**
					SPO		
Clostridium novyi biovar A		83.39	0.37	CEL 1	99		
Atopobium minutum		16.61	0.21	GAL 10	1		
01041010	**Acceptable identification**						**01041010**
					COC	SPO	
Clostridium novyi biovar A		94.24	0.50	CEL 1	1	99	
Atopobium minutum		4.97	0.24	GAL 10	1	1	
Peptostreptococcus anaerobius		0.79	-.03		99	1	
01041011	**Good identification**						**01041011**
Clostridium novyi biovar A		100.0	0.50	CEL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01041012	**Low discrimination**						**01041012**
					SPO		
Clostridium novyi biovar A		83.39	0.37	CEL 1	99		
Atopobium minutum		16.61	0.21	GAL 10	1		
01041013	**Good identification**						**01041013**
Clostridium novyi biovar A		100.0	0.37	CEL 1			
01041020	**Low discrimination**						**01041020**
					SPO		
Atopobium minutum		86.50	0.50	CEL 1	1		
Clostridium novyi biovar A		12.43	0.39		99		
Clostridium cadaveris		0.65	0.03		99		
01041021	**Acceptable identification**						**01041021**
					SPO		
Clostridium novyi biovar A		93.43	0.39	CEL 1	99		
Atopobium minutum		6.57	0.15	GAL 10	1		
01041022	**Low discrimination**						**01041022**
					GRA	SPO	
Atopobium minutum		87.74	0.47	CEL 1	99	1	
Pseudoflavonifractor capillosus		8.93	0.15		1	1	
Clostridium novyi biovar A		3.34	0.26		99	99	
01041030	**Acceptable identification**						**01041030**
					SPO		
Clostridium novyi biovar A		93.43	0.39	CEL 1	99		
Atopobium minutum		6.57	0.15	GAL 10	1		
01041031	**Good identification**						**01041031**
Clostridium novyi biovar A		100.0	0.39	CEL 1			
01041040	**Low discrimination**						**01041040**
					SPO		
Clostridium novyi biovar A		80.34	0.47	CEL 1	99		
Clostridium botulinum biovar A		8.62	0.18	GAL 10	99		
Atopobium minutum		6.35	0.24		1		
01041041	**Good identification**						**01041041**
Clostridium novyi biovar A		100.0	0.47	CEL 1			
01041050	**Good identification**						**01041050**
Clostridium novyi biovar A		100.0	0.47	CEL 1			
01041051	**Good identification**						**01041051**
Clostridium novyi biovar A		100.0	0.47	CEL 1			
01041060	**Acceptable identification**						**01041060**
					SPO		
Clostridium novyi biovar A		90.46	0.36	CEL 1	99		
Atopobium minutum		9.54	0.15	GAL 10	1		
01041061	**Good identification**						**01041061**
Clostridium novyi biovar A		100.0	0.36	CEL 1			
01041070	**Good identification**						**01041070**
Clostridium novyi biovar A		100.0	0.36	CEL 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
01041071	**Good identification**						**01041071**
	Clostridium novyi biovar A	100.0	0.36	CEL 1			
01044040	**Good identification**						**01044040**
	Clostridium glycolicum	97.38	0.49	FRU 90			
	Clostridium botulinum biovar A	1.33	0.18	CEL 1			
	Clostridium novyi biovar A	1.13	0.29				
01050000	**Low discrimination**						**01050000**
					GRA		
	Pseudoramibacter alactolyticus	65.81	0.66	CEL 1	99		
	Fusobacterium varium	9.25	0.51		1		
	Atopobium minutum	9.12	0.65		99		
01050002	**Low discrimination**						**01050002**
					COC	GRA	
	Pseudoramibacter alactolyticus	49.66	0.50	CEL 1	1	99	
	Atopobium minutum	41.28	0.62	BGL 10	1	99	
	Megasphaera elsdenii	5.23	0.44		99	1	
01050003	**Low discrimination**						**01050003**
					SPO		
	Clostridium novyi biovar A	43.76	0.37	FRU 10	99		
	Pseudoramibacter alactolyticus	29.04	0.15	CEL 1	1		
	Atopobium minutum	24.14	0.28		1		
01050010	**Low discrimination**						**01050010**
					SPO		
	Clostridium difficile	71.03	0.49	CEL 5	99		
	Pseudoramibacter alactolyticus	11.87	0.32	MAN 90	1		
	Clostridium novyi biovar A	11.26	0.50		99		
01050011	**Acceptable identification**						**01050011**
					SPO		
	Clostridium novyi biovar A	93.08	0.50	FRU 10	99		
	Clostridium difficile	5.93	0.15	CEL 1	99		
	Pseudoramibacter alactolyticus	0.99	-.03		1		
01050012	**Low discrimination**						**01050012**
					SPO		
	Clostridium novyi biovar A	37.79	0.37	FRU 10	99		
	Pseudoramibacter alactolyticus	25.08	0.15	CEL 1	1		
	Atopobium minutum	20.85	0.28		1		
01050013	**Good identification**						**01050013**
	Clostridium novyi biovar A	100.0	0.37	FRU 10			
01050020	**Low discrimination**						**01050020**
					SPO		
	Pseudoramibacter alactolyticus	72.54	0.53	CEL 1	1		
	Atopobium minutum	18.98	0.57		1		
	Clostridium cadaveris	5.10	0.37		99		
01050021	**Low discrimination**						**01050021**
					SPO		
	Clostridium novyi biovar A	50.23	0.39	FRU 10	99		
	Pseudoramibacter alactolyticus	37.37	0.19	CEL 1	1		
	Atopobium minutum	9.78	0.23		1		
01050022	**Low discrimination**						**01050022**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Atopobium minutum		58.99	0.54	CEL 1	99			
Pseudoramibacter alactolyticus		37.57	0.37		99			
Pseudoflavonifractor capillosus		2.17	0.15		1			
01050030	**Low discrimination**							**01050030**
					SPO			
Clostridium novyi biovar A		44.53	0.39	FRU 10	99			
Pseudoramibacter alactolyticus		33.13	0.19	CEL 1	1			
Clostridium difficile		11.35	0.15		99			
01050031	**Good identification**							**01050031**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01050040	**Low discrimination**							**01050040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		69.64	0.56	CEL 1	1	1	99	
Megasphaera elsdenii		19.55	0.57		99	1	1	
Clostridium glycolicum		6.60	0.43		1	99	99	
01050041	**Low discrimination**							**01050041**
					COC	SPO	GRA	
Clostridium novyi biovar A		64.50	0.47	FRU 10	1	99	99	
Pseudoramibacter alactolyticus		25.49	0.21	CEL 1	1	1	99	
Megasphaera elsdenii		7.16	0.23		99	1	1	
01050042	**Low discrimination**							**01050042**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		73.26	0.39	CEL 1	1	1	99	
Megasphaera elsdenii		20.56	0.41	BGL 10	99	1	1	
Clostridium novyi biovar A		2.97	0.34		1	99	99	
01050050	**Low discrimination**							**01050050**
					SPO			
Clostridium novyi biovar A		48.82	0.47	FRU 10	99			
Clostridium difficile		24.31	0.27	CEL 1	99			
Pseudoramibacter alactolyticus		19.30	0.21		1			
01050051	**Good identification**							**01050051**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01050060	**Acceptable identification**							**01050060**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		92.46	0.43	CEL 1	1	1	99	
Clostridium novyi biovar A		3.35	0.36		1	99	99	
Megasphaera elsdenii		1.49	0.23		99	1	1	
01050061	**Low discrimination**							**01050061**
					SPO			
Clostridium novyi biovar A		78.19	0.36	FRU 10	99			
Pseudoramibacter alactolyticus		21.81	0.09	CEL 1	1			
01050070	**Low discrimination**							**01050070**
					SPO			
Clostridium novyi biovar A		78.19	0.36	FRU 10	99			
Pseudoramibacter alactolyticus		21.81	0.09	CEL 1	1			
01050071	**Good identification**							**01050071**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01050100	**Low discrimination**							**01050100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		54.94	0.58		1	
Pseudoramibacter alactolyticus		21.73	0.32		99	
Actinomyces meyeri		12.40	0.38		99	
01050102	**Low discrimination**					**01050102**
					GRA	
Fusobacterium mortiferum		54.68	0.42	BGL 10	1	
Pseudoramibacter alactolyticus		21.63	0.15		99	
Atopobium minutum		17.98	0.28		99	
01051000	**Low discrimination**					**01051000**
					GRA	
Atopobium minutum		56.94	0.57	CEL 1	99	
Fusobacterium mortiferum		20.24	0.55		1	
Pseudoramibacter alactolyticus		12.45	0.32		99	
01051002	**Low discrimination**					**01051002**
					GRA	
Atopobium minutum		87.45	0.54	CEL 1	99	
Fusobacterium mortiferum		5.18	0.39		1	
Pseudoramibacter alactolyticus		3.19	0.15		99	
01051020	**Low discrimination**					**01051020**
					CAT	
Atopobium minutum		75.89	0.49	CEL 1	1	
Propionibacterium acnes		9.32	0.26		95	
Pseudoramibacter alactolyticus		8.79	0.19		1	
01051022	**Acceptable identification**					**01051022**
Atopobium minutum		93.30	0.46	CEL 1		
Lactobacillus fermentum		3.33	0.28			
Pseudoramibacter alactolyticus		1.80	0.03			
01051100	**Low discrimination**					**01051100**
					GRA	
Fusobacterium mortiferum		77.67	0.61		1	
Actinomyces meyeri		17.54	0.41		99	
Lactobacillus fermentum		3.64	0.40		99	
01051102	**Low discrimination**					**01051102**
					GRA	
Fusobacterium mortiferum		64.22	0.45	BGL 10	1	
Lactobacillus fermentum		27.06	0.40		99	
Atopobium minutum		4.69	0.20		99	
01051400	**Low discrimination**					**01051400**
Lactobacillus fermentum		46.30	0.36	SUC 87		
Actinomyces meyeri		22.77	0.20	CEL 1		
Atopobium minutum		19.65	0.23			
01051402	**Low discrimination**					**01051402**
Lactobacillus fermentum		77.95	0.36	SUC 87		
Atopobium minutum		22.05	0.20	CEL 1		
01051502	**Good identification**					**01051502**
Lactobacillus fermentum		96.41	0.40	SUC 87		
Actinomyces meyeri		3.60	0.05	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01052000	**Low discrimination**							**01052000**
					SPO			
Pseudoramibacter alactolyticus		48.90	0.61	CEL 1	1			
Eubacterium limosum		31.10	0.60		1			
Clostridium difficile		10.12	0.53		99			
01052002	**Low discrimination**							**01052002**
					COC	GRA		
Pseudoramibacter alactolyticus		77.58	0.45	CEL 1	1	99		
Megasphaera elsdenii		15.17	0.44	BGL 10	99	1		
Eubacterium limosum		4.49	0.26		1	99		
01052010	**Very good identification**							**01052010**
					SPO			
Clostridium difficile		98.18	0.65	CEL 5	99			
Pseudoramibacter alactolyticus		0.98	0.27		1			
Eubacterium limosum		0.62	0.26		1			
01052020	**Acceptable identification**							**01052020**
					SPO			
Pseudoramibacter alactolyticus		92.86	0.48	CEL 1	1			
Eubacterium limosum		3.38	0.26		1			
Clostridium difficile		1.10	0.19		99			
01052040	**Low discrimination**							**01052040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		62.77	0.51	CEL 1	1	1	99	
Megasphaera elsdenii		32.72	0.57		99	1	1	
Clostridium difficile		2.73	0.31		1	99	99	
01052050	**Acceptable identification**							**01052050**
					COC	SPO	GRA	
Clostridium difficile		93.07	0.43	CEL 5	1	99	99	
Pseudoramibacter alactolyticus		4.42	0.17	SOR 5	1	1	99	
Megasphaera elsdenii		2.30	0.23		99	1	1	
01052060	**Good identification**							**01052060**
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		95.96	0.38	CEL 1	1	1	99	
Megasphaera elsdenii		2.86	0.23		99	1	1	
Propionibacterium acnes		0.94	0.11		1	95	99	
01054000	**Low discrimination**							**01054000**
					GRA	SPO		
Clostridium glycolicum		49.85	0.43	CEL 1	99	99		
Pseudoramibacter alactolyticus		21.26	0.32	SOR 95	99	1		
Fusobacterium varium		12.32	0.28		1	1		
01054040	**Very good identification**							**01054040**
					COC	SPO	GRA	
Clostridium glycolicum		99.25	0.65	CEL 1	1	99	99	
Pseudoramibacter alactolyticus		0.56	0.21		1	1	99	
Megasphaera elsdenii		0.16	0.23		99	1	1	
01054100	**Low discrimination**							**01054100**
					GRA	SPO		
Actinomyces meyeri		82.82	0.43	SUC 93	99	1		
Fusobacterium mortiferum		14.22	0.39	CEL 1	1	1		
Clostridium glycolicum		1.77	0.09		99	99		
01055000	**Low discrimination**							**01055000**
					GRA	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		44.95	0.35	SUC 93	99	1	
Clostridium glycolicum		28.18	0.27	CEL 1	99	99	
Fusobacterium mortiferum		13.23	0.36		1	1	
01055040	**Good identification**						**01055040**
Clostridium glycolicum		100.0	0.49	CEL 1			
01055100	**Low discrimination**						**01055100**
					GRA		
Actinomyces meyeri		81.01	0.46	SUC 93	99		
Fusobacterium mortiferum		13.91	0.42	CEL 1	1		
Lactobacillus fermentum		5.08	0.36		99		
01055102	**Low discrimination**						**01055102**
					GRA		
Lactobacillus fermentum		55.64	0.36	SUC 87	99		
Actinomyces meyeri		27.44	0.20	CEL 1	99		
Fusobacterium mortiferum		16.93	0.26		1		
01055500	**Low discrimination**						**01055500**
Actinomyces meyeri		79.94	0.35	SUC 93			
Lactobacillus fermentum		20.06	0.36	CEL 1			
01055502	**Low discrimination**						**01055502**
Lactobacillus fermentum		89.03	0.36	SUC 87			
Actinomyces meyeri		10.98	0.10	CEL 1			
01056010	**Good identification**						**01056010**
					SPO		
Clostridium difficile		99.81	0.43	CEL 5	99		
Pseudoramibacter alactolyticus		0.19	-.07	XYL 5	1		
01060000	**Low discrimination**						**01060000**
					COC	SPO	
Clostridium botulinum biovar A		64.98	0.51	CEL 1	1	99	
Clostridium sporogenes		20.88	0.40		1	99	
Peptostreptococcus anaerobius		9.12	0.32		99	1	
01060040	**Good identification**						**01060040**
Clostridium botulinum biovar A		95.25	0.42	CEL 1			
Clostridium novyi biovar A		3.26	0.29				
Clostridium sporogenes		1.04	0.06				
01070100	**Good identification**						**01070100**
					GRA		
Fusobacterium mortiferum		97.94	0.49		1		
Pseudoramibacter alactolyticus		1.31	-.03		99		
Actinomyces meyeri		0.75	0.04		99		
01071000	**Low discrimination**						**01071000**
					GRA		
Fusobacterium mortiferum		87.20	0.46		1		
Atopobium minutum		8.29	0.23		99		
Pseudoramibacter alactolyticus		1.81	-.03		99		
01071100	**Good identification**						**01071100**
					GRA	SPO	
Fusobacterium mortiferum		98.80	0.52		1	1	
Actinomyces meyeri		0.75	0.07		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium septicum		0.45	-.05		99	99		
01071102	**Good identification**							**01071102**
Fusobacterium mortiferum		100.0	0.36	BGL 10				
01072010	**Good identification**							**01072010**
					SPO			
Clostridium difficile		99.87	0.46	TRE 7	99			
Pseudoramibacter alactolyticus		0.13	-.07	CEL 5	1			
01100000	**Low discrimination**							**01100000**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		43.04	0.97		99	99	1	
Prevotella disiens		23.72	0.91		1	1	1	
Clostridium botulinum biovar A		15.21	0.94		1	99	99	
01100001	**Low discrimination**							**01100001**
					COC	SPO		
Clostridium novyi biovar A		81.18	1.00		1	99		
Clostridium sordellii		7.84	0.66		1	99		
Peptostreptococcus anaerobius		5.00	0.63		99	1		
01100002	**Low discrimination**							**01100002**
					COC	SPO		
Clostridium sporogenes		55.13	0.79		1	99		
Clostridium novyi biovar A		24.77	0.87		1	99		
Peptostreptococcus anaerobius		8.64	0.63		99	1		
01100003	**Very good identification**							**01100003**
Clostridium novyi biovar A		96.44	0.87					
Clostridium sporogenes		2.17	0.45					
Clostridium sordellii		0.53	0.32					
01100010	**Low discrimination**							**01100010**
					COC	GRA	SPO	
Clostridium novyi biovar A		88.27	1.00		1	99	99	
Peptostreptococcus anaerobius		5.43	0.63		99	99	1	
Prevotella disiens		2.99	0.57		1	1	1	
01100011	**Excellent identification**							**01100011**
					COC	SPO		
Clostridium novyi biovar A		99.77	1.00		1	99		
Clostridium sordellii		0.10	0.32		1	99		
Peptostreptococcus anaerobius		0.06	0.29		99	1		
01100012	**Very good identification**							**01100012**
					COC	SPO		
Clostridium novyi biovar A		97.03	0.87		1	99		
Clostridium sporogenes		2.18	0.45		1	99		
Peptostreptococcus anaerobius		0.34	0.29		99	1		
01100013	**Very good identification**							**01100013**
Clostridium novyi biovar A		99.97	0.87					
Clostridium sporogenes		0.02	0.11					
Clostridium sordellii		0.01	-.03					
01100020	**Low discrimination**							**01100020**
					COC	GRA	SPO	
Clostridium novyi biovar A		60.57	0.90		1	99	99	
Peptostreptococcus anaerobius		14.91	0.63		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella disiens		8.22	0.57		1	1	1	
01100021	**Very good identification**							**01100021**
					COC	SPO		
Clostridium novyi biovar A		98.95	0.90		1	99		
Clostridium sordellii		0.39	0.32		1	99		
Peptostreptococcus anaerobius		0.25	0.29		99	1		
01100022	**Low discrimination**							**01100022**
					SPO			
Clostridium novyi biovar A		80.51	0.77		99			
Clostridium sporogenes		7.24	0.45		99			
Atopobium minutum		7.13	0.55		1			
01100023	**Very good identification**							**01100023**
					SPO			
Clostridium novyi biovar A		99.76	0.77		99			
Clostridium sporogenes		0.09	0.11		99			
Atopobium minutum		0.09	0.21		1			
01100030	**Very good identification**							**01100030**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.35	0.90		1	99	99	
Peptostreptococcus anaerobius		0.25	0.29		99	99	1	
Prevotella disiens		0.14	0.23		1	1	1	
01100031	**Very good identification**							**01100031**
					COC	SPO		
Clostridium novyi biovar A		99.99	0.90		1	99		
Clostridium sordellii		0.00	-.03		1	99		
Peptostreptococcus anaerobius		0.00	-.06		99	1		
01100032	**Very good identification**							**01100032**
					SPO			
Clostridium novyi biovar A		99.78	0.77		99			
Clostridium sporogenes		0.09	0.11		99			
Atopobium minutum		0.09	0.21		1			
01100033	**Very good identification**							**01100033**
Clostridium novyi biovar A		100.0	0.77					
01100040	**Low discrimination**							**01100040**
					COC	GRA		
Clostridium novyi biovar A		43.72	0.97		1	99		
Clostridium botulinum biovar A		42.19	0.85		1	99		
Megasphaera elsdenii		4.21	0.72		99	1		
01100041	**Very good identification**							**01100041**
Clostridium novyi biovar A		98.13	0.97					
Clostridium botulinum biovar A		0.96	0.51					
Clostridium sordellii		0.59	0.42					
01100042	**Low discrimination**							**01100042**
					COC	GRA		
Clostridium novyi biovar A		86.00	0.84		1	99		
Megasphaera elsdenii		5.22	0.56		99	1		
Clostridium botulinum biovar A		4.75	0.51		1	99		
01100043	**Very good identification**							**01100043**
					COC	GRA		
Clostridium novyi biovar A		99.81	0.84		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		0.06	0.22		99	1	
Clostridium botulinum biovar A		0.06	0.17		1	99	
01100050	**Very good identification**						**01100050**
					COC	GRA	
Clostridium novyi biovar A		98.67	0.97		1	99	
Clostridium botulinum biovar A		0.96	0.51		1	99	
Megasphaera elsdenii		0.10	0.38		99	1	
01100051	**Excellent identification**						**01100051**
Clostridium novyi biovar A		99.98	0.97				
Clostridium botulinum biovar A		0.01	0.17				
Clostridium sordellii		0.01	0.08				
01100052	**Very good identification**						**01100052**
					COC	GRA	
Clostridium novyi biovar A		99.84	0.84		1	99	
Megasphaera elsdenii		0.06	0.22		99	1	
Clostridium botulinum biovar A		0.06	0.17		1	99	
01100053	**Very good identification**						**01100053**
Clostridium novyi biovar A		100.0	0.84				
01100060	**Good identification**						**01100060**
					COC	GRA	
Clostridium novyi biovar A		94.51	0.87		1	99	
Clostridium botulinum biovar A		3.69	0.51		1	99	
Megasphaera elsdenii		0.37	0.38		99	1	
01100061	**Very good identification**						**01100061**
Clostridium novyi biovar A		99.93	0.87				
Clostridium botulinum biovar A		0.04	0.17				
Clostridium sordellii		0.02	0.08				
01100062	**Very good identification**						**01100062**
					COC	GRA	
Clostridium novyi biovar A		99.17	0.74		1	99	
Megasphaera elsdenii		0.24	0.22		99	1	
Clostridium botulinum biovar A		0.22	0.17		1	99	
01100063	**Very good identification**						**01100063**
Clostridium novyi biovar A		100.0	0.74				
01100070	**Very good identification**						**01100070**
Clostridium novyi biovar A		99.85	0.87				
Clostridium botulinum biovar B		0.10	0.21				
Clostridium botulinum biovar A		0.04	0.17				
01100071	**Very good identification**						**01100071**
Clostridium novyi biovar A		100.0	0.87				
01100072	**Very good identification**						**01100072**
Clostridium novyi biovar A		100.0	0.74				
01100073	**Very good identification**						**01100073**
Clostridium novyi biovar A		100.0	0.74				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01100100	**Low discrimination**							**01100100**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		41.01	0.63	LAC 1	99	99	1	
Prevotella disiens		22.60	0.57		1	1	1	
Clostridium botulinum biovar A		14.50	0.60		1	99	99	
01100101	**Low discrimination**							**01100101**
					COC	SPO		
Clostridium novyi biovar A		80.75	0.66	LAC 1	1	99		
Clostridium sordellii		7.79	0.32		1	99		
Peptostreptococcus anaerobius		4.97	0.29		99	1		
01100102	**Low discrimination**							**01100102**
					COC	SPO		
Clostridium sporogenes		53.39	0.45	LAC 1	1	99		
Clostridium novyi biovar A		23.99	0.53		1	99		
Peptostreptococcus anaerobius		8.37	0.29		99	1		
01100103	**Good identification**							**01100103**
Clostridium novyi biovar A		96.94	0.53	LAC 1				
Clostridium sporogenes		2.18	0.11					
Clostridium sordellii		0.54	-.03					
01100110	**Low discrimination**							**01100110**
					COC	GRA	SPO	
Clostridium novyi biovar A		87.78	0.66	LAC 1	1	99	99	
Peptostreptococcus anaerobius		5.40	0.29		99	99	1	
Prevotella disiens		2.98	0.23		1	1	1	
01100111	**Very good identification**							**01100111**
					COC	SPO		
Clostridium novyi biovar A		99.84	0.66	LAC 1	1	99		
Clostridium sordellii		0.10	-.03		1	99		
Peptostreptococcus anaerobius		0.06	-.06		99	1		
01100112	**Good identification**							**01100112**
					COC	SPO		
Clostridium novyi biovar A		97.47	0.53	LAC 1	1	99		
Clostridium sporogenes		2.19	0.11		1	99		
Peptostreptococcus anaerobius		0.34	-.06		99	1		
01100113	**Good identification**							**01100113**
Clostridium novyi biovar A		100.0	0.53	LAC 1				
01100120	**Very good identification**							**01100120**
					COC	GRA	SPO	
Prevotella bivia		97.88	0.66	MNS 99	1	1	1	
Clostridium novyi biovar A		1.12	0.56		1	99	99	
Peptostreptococcus anaerobius		0.28	0.29		99	99	1	
01100121	**Low discrimination**							**01100121**
					GRA	SPO		
Clostridium novyi biovar A		52.87	0.56	LAC 1	99	99		
Prevotella bivia		46.68	0.32		1	1		
Clostridium sordellii		0.21	-.03		99	99		
01100123	**Good identification**							**01100123**
					GRA	SPO		
Clostridium novyi biovar A		95.19	0.43	LAC 1	99	99		
Prevotella bivia		4.81	-.03		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01100130	**Low discrimination**							**01100130**
					COC	GRA	SPO	
Clostridium novyi biovar A		52.98	0.56	LAC 1	1	99	99	
Prevotella bivia		46.78	0.32		1	1	1	
Peptostreptococcus anaerobius		0.13	-.06		99	99	1	
01100131	**Good identification**							**01100131**
					GRA	SPO		
Clostridium novyi biovar A		99.12	0.56	LAC 1	99	99		
Prevotella bivia		0.88	-.03		1	1		
01100132	**Good identification**							**01100132**
					GRA	SPO		
Clostridium novyi biovar A		95.19	0.43	LAC 1	99	99		
Prevotella bivia		4.81	-.03		1	1		
01100133	**Good identification**							**01100133**
Clostridium novyi biovar A		100.0	0.43	LAC 1				
01100140	**Low discrimination**							**01100140**
					COC	GRA		
Clostridium novyi biovar A		43.55	0.63	LAC 1	1	99		
Clostridium botulinum biovar A		42.02	0.51		1	99		
Megasphaera elsdenii		4.20	0.38		99	1		
01100141	**Very good identification**							**01100141**
Clostridium novyi biovar A		98.26	0.63	LAC 1				
Clostridium botulinum biovar A		0.96	0.17					
Clostridium sordellii		0.59	0.08					
01100142	**Low discrimination**							**01100142**
					COC	GRA		
Clostridium novyi biovar A		86.59	0.50	LAC 1	1	99		
Megasphaera elsdenii		5.25	0.22		99	1		
Clostridium botulinum biovar A		4.78	0.17		1	99		
01100143	**Good identification**							**01100143**
Clostridium novyi biovar A		100.0	0.50	LAC 1				
01100150	**Very good identification**							**01100150**
					COC	GRA		
Clostridium novyi biovar A		98.85	0.63	LAC 1	1	99		
Clostridium botulinum biovar A		0.96	0.17		1	99		
Megasphaera elsdenii		0.10	0.04		99	1		
01100151	**Very good identification**							**01100151**
Clostridium novyi biovar A		100.0	0.63	LAC 1				
01100152	**Good identification**							**01100152**
Clostridium novyi biovar A		100.0	0.50	LAC 1				
01100153	**Good identification**							**01100153**
Clostridium novyi biovar A		100.0	0.50	LAC 1				
01100161	**Good identification**							**01100161**
					GRA	SPO		
Clostridium novyi biovar A		98.68	0.53	LAC 1	99	99		
Prevotella bivia		1.32	-.03		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01100162	**Acceptable identification**							**01100162**
					GRA	SPO		
Clostridium novyi biovar A		92.95	0.40	LAC 1	99	99		
Prevotella bivia		7.05	-.03		1	1		
01100163	**Good identification**							**01100163**
Clostridium novyi biovar A		100.0	0.40	LAC 1				
01100170	**Good identification**							**01100170**
					GRA	SPO		
Clostridium novyi biovar A		98.68	0.53	LAC 1	99	99		
Prevotella bivia		1.32	-.03		1	1		
01100171	**Good identification**							**01100171**
Clostridium novyi biovar A		100.0	0.53	LAC 1				
01100172	**Good identification**							**01100172**
Clostridium novyi biovar A		100.0	0.40	LAC 1				
01100173	**Good identification**							**01100173**
Clostridium novyi biovar A		100.0	0.40	LAC 1				
01100200	**Low discrimination**							**01100200**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		43.04	0.63	RHA 1	99	99	1	
Prevotella disiens		23.72	0.57		1	1	1	
Clostridium botulinum biovar A		15.21	0.60		1	99	99	
01100201	**Low discrimination**							**01100201**
					COC	SPO		
Clostridium novyi biovar A		81.27	0.66	RHA 1	1	99		
Clostridium sordellii		7.84	0.32		1	99		
Peptostreptococcus anaerobius		5.00	0.29		99	1		
01100202	**Low discrimination**							**01100202**
					COC	SPO		
Clostridium sporogenes		55.24	0.45	RHA 1	1	99		
Clostridium novyi biovar A		24.82	0.53		1	99		
Peptostreptococcus anaerobius		8.66	0.29		99	1		
01100203	**Good identification**							**01100203**
Clostridium novyi biovar A		96.94	0.53	RHA 1				
Clostridium sporogenes		2.18	0.11					
Clostridium sordellii		0.54	-.03					
01100210	**Low discrimination**							**01100210**
					COC	GRA	SPO	
Clostridium novyi biovar A		88.33	0.66	RHA 1	1	99	99	
Peptostreptococcus anaerobius		5.44	0.29		99	99	1	
Prevotella disiens		3.00	0.23		1	1	1	
01100211	**Very good identification**							**01100211**
					COC	SPO		
Clostridium novyi biovar A		99.84	0.66	RHA 1	1	99		
Clostridium sordellii		0.10	-.03		1	99		
Peptostreptococcus anaerobius		0.06	-.06		99	1		
01100212	**Good identification**							**01100212**
					COC	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		97.47	0.53	RHA 1	1	99		
Clostridium sporogenes		2.19	0.11		1	99		
Peptostreptococcus anaerobius		0.34	-.06		99	1		
01100213	**Good identification**							**01100213**
Clostridium novyi biovar A		100.0	0.53	RHA 1				
01100220	**Low discrimination**							**01100220**
					COC	GRA	SPO	
Clostridium novyi biovar A		60.69	0.56	RHA 1	1	99	99	
Peptostreptococcus anaerobius		14.94	0.29		99	99	1	
Prevotella disiens		8.24	0.23		1	1	1	
01100221	**Good identification**							**01100221**
					COC	SPO		
Clostridium novyi biovar A		99.37	0.56	RHA 1	1	99		
Clostridium sordellii		0.39	-.03		1	99		
Peptostreptococcus anaerobius		0.25	-.06		99	1		
01100222	**Low discrimination**							**01100222**
					SPO			
Clostridium novyi biovar A		82.00	0.43	RHA 1	99			
Clostridium sporogenes		7.37	0.11		99			
Atopobium minutum		7.26	0.21		1			
01100223	**Good identification**							**01100223**
Clostridium novyi biovar A		100.0	0.43	RHA 1				
01100230	**Good identification**							**01100230**
					COC	SPO		
Clostridium novyi biovar A		99.75	0.56	RHA 1	1	99		
Peptostreptococcus anaerobius		0.25	-.06		99	1		
01100231	**Good identification**							**01100231**
Clostridium novyi biovar A		100.0	0.56	RHA 1				
01100232	**Good identification**							**01100232**
Clostridium novyi biovar A		100.0	0.43	RHA 1				
01100233	**Good identification**							**01100233**
Clostridium novyi biovar A		100.0	0.43	RHA 1				
01100240	**Low discrimination**							**01100240**
					COC	GRA		
Clostridium novyi biovar A		43.75	0.63	RHA 1	1	99		
Clostridium botulinum biovar A		42.22	0.51		1	99		
Megasphaera elsdenii		4.22	0.38		99	1		
01100241	**Very good identification**							**01100241**
Clostridium novyi biovar A		98.26	0.63	RHA 1				
Clostridium botulinum biovar A		0.96	0.17					
Clostridium sordellii		0.59	0.08					
01100242	**Low discrimination**							**01100242**
					COC	GRA		
Clostridium novyi biovar A		86.59	0.50	RHA 1	1	99		
Megasphaera elsdenii		5.25	0.22		99	1		
Clostridium botulinum biovar A		4.78	0.17		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01100243	**Good identification**							**01100243**
Clostridium novyi biovar A		100.0	0.50	RHA 1				
01100250	**Very good identification**							**01100250**
					COC	GRA		
Clostridium novyi biovar A		98.85	0.63	RHA 1	1	99		
Clostridium botulinum biovar A		0.96	0.17		1	99		
Megasphaera elsdenii		0.10	0.04		99	1		
01100251	**Very good identification**							**01100251**
Clostridium novyi biovar A		100.0	0.63	RHA 1				
01100252	**Good identification**							**01100252**
Clostridium novyi biovar A		100.0	0.50	RHA 1				
01100253	**Good identification**							**01100253**
Clostridium novyi biovar A		100.0	0.50	RHA 1				
01100260	**Good identification**							**01100260**
					COC	GRA		
Clostridium novyi biovar A		95.25	0.53	RHA 1	1	99		
Clostridium botulinum biovar A		3.71	0.17		1	99		
Megasphaera elsdenii		0.37	0.04		99	1		
01100261	**Good identification**							**01100261**
Clostridium novyi biovar A		100.0	0.53	RHA 1				
01100262	**Good identification**							**01100262**
Clostridium novyi biovar A		100.0	0.40	RHA 1				
01100263	**Good identification**							**01100263**
Clostridium novyi biovar A		100.0	0.40	RHA 1				
01100270	**Good identification**							**01100270**
Clostridium novyi biovar A		100.0	0.53	RHA 1				
01100271	**Good identification**							**01100271**
Clostridium novyi biovar A		100.0	0.53	RHA 1				
01100272	**Good identification**							**01100272**
Clostridium novyi biovar A		100.0	0.40	RHA 1				
01100273	**Good identification**							**01100273**
Clostridium novyi biovar A		100.0	0.40	RHA 1				
01100400	**Low discrimination**							**01100400**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		42.16	0.63	ARA 1	99	99	1	
Prevotella disiens		23.23	0.57		1	1	1	
Clostridium botulinum biovar A		14.90	0.60		1	99	99	
01100401	**Low discrimination**							**01100401**
					COC	SPO		
Clostridium novyi biovar A		81.07	0.66	ARA 1	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sordellii		7.83	0.32		1	99		
Peptostreptococcus anaerobius		4.99	0.29		99	1		
01100402	**Low discrimination**							**01100402**
					COC	SPO		
Clostridium sporogenes		54.95	0.45	ARA 1	1	99		
Clostridium novyi biovar A		24.69	0.53		1	99		
Peptostreptococcus anaerobius		8.61	0.29		99	1		
01100403	**Good identification**							**01100403**
Clostridium novyi biovar A		96.94	0.53	ARA 1				
Clostridium sporogenes		2.18	0.11					
Clostridium sordellii		0.54	-.03					
01100410	**Low discrimination**							**01100410**
					COC	GRA	SPO	
Clostridium novyi biovar A		88.09	0.66	ARA 1	1	99	99	
Peptostreptococcus anaerobius		5.42	0.29		99	99	1	
Prevotella disiens		2.99	0.23		1	1	1	
01100411	**Very good identification**							**01100411**
					COC	SPO		
Clostridium novyi biovar A		99.84	0.66	ARA 1	1	99		
Clostridium sordellii		0.10	-.03		1	99		
Peptostreptococcus anaerobius		0.06	-.06		99	1		
01100412	**Good identification**							**01100412**
					COC	SPO		
Clostridium novyi biovar A		97.47	0.53	ARA 1	1	99		
Clostridium sporogenes		2.19	0.11		1	99		
Peptostreptococcus anaerobius		0.34	-.06		99	1		
01100413	**Good identification**							**01100413**
Clostridium novyi biovar A		100.0	0.53	ARA 1				
01100420	**Low discrimination**							**01100420**
					COC	GRA	SPO	
Clostridium novyi biovar A		56.08	0.56	ARA 1	1	99	99	
Peptostreptococcus anaerobius		13.81	0.29		99	99	1	
Prevotella disiens		7.61	0.23		1	1	1	
01100421	**Good identification**							**01100421**
					COC	SPO		
Clostridium novyi biovar A		99.37	0.56	ARA 1	1	99		
Clostridium sordellii		0.39	-.03		1	99		
Peptostreptococcus anaerobius		0.25	-.06		99	1		
01100422	**Low discrimination**							**01100422**
					SPO			
Clostridium novyi biovar A		82.00	0.43	ARA 1	99			
Clostridium sporogenes		7.37	0.11		99			
Atopobium minutum		7.26	0.21		1			
01100423	**Good identification**							**01100423**
Clostridium novyi biovar A		100.0	0.43	ARA 1				
01100430	**Good identification**							**01100430**
					COC	SPO		
Clostridium novyi biovar A		99.75	0.56	ARA 1	1	99		
Peptostreptococcus anaerobius		0.25	-.06		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01100431	**Good identification**						**01100431**
Clostridium novyi biovar A		100.0	0.56	ARA 1			
01100432	**Good identification**						**01100432**
Clostridium novyi biovar A		100.0	0.43	ARA 1			
01100433	**Good identification**						**01100433**
Clostridium novyi biovar A		100.0	0.43	ARA 1			
01100440	**Low discrimination**						**01100440**
					COC	GRA	
Clostridium novyi biovar A		43.66	0.63	ARA 1	1	99	
Clostridium botulinum biovar A		42.13	0.51		1	99	
Megasphaera elsdenii		4.21	0.38		99	1	
01100441	**Very good identification**						**01100441**
Clostridium novyi biovar A		98.26	0.63	ARA 1			
Clostridium botulinum biovar A		0.96	0.17				
Clostridium sordellii		0.59	0.08				
01100442	**Low discrimination**						**01100442**
					COC	GRA	
Clostridium novyi biovar A		86.59	0.50	ARA 1	1	99	
Megasphaera elsdenii		5.25	0.22		99	1	
Clostridium botulinum biovar A		4.78	0.17		1	99	
01100443	**Good identification**						**01100443**
Clostridium novyi biovar A		100.0	0.50	ARA 1			
01100450	**Very good identification**						**01100450**
					COC	GRA	
Clostridium novyi biovar A		98.85	0.63	ARA 1	1	99	
Clostridium botulinum biovar A		0.96	0.17		1	99	
Megasphaera elsdenii		0.10	0.04		99	1	
01100451	**Very good identification**						**01100451**
Clostridium novyi biovar A		100.0	0.63	ARA 1			
01100452	**Good identification**						**01100452**
Clostridium novyi biovar A		100.0	0.50	ARA 1			
01100453	**Good identification**						**01100453**
Clostridium novyi biovar A		100.0	0.50	ARA 1			
01100460	**Good identification**						**01100460**
					COC	GRA	
Clostridium novyi biovar A		95.25	0.53	ARA 1	1	99	
Clostridium botulinum biovar A		3.71	0.17		1	99	
Megasphaera elsdenii		0.37	0.04		99	1	
01100461	**Good identification**						**01100461**
Clostridium novyi biovar A		100.0	0.53	ARA 1			
01100462	**Good identification**						**01100462**
Clostridium novyi biovar A		100.0	0.40	ARA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01100463	**Good identification**							**01100463**
Clostridium novyi biovar A		100.0	0.40	ARA 1				
01100470	**Good identification**							**01100470**
Clostridium novyi biovar A		100.0	0.53	ARA 1				
01100471	**Good identification**							**01100471**
Clostridium novyi biovar A		100.0	0.53	ARA 1				
01100472	**Good identification**							**01100472**
Clostridium novyi biovar A		100.0	0.40	ARA 1				
01100473	**Good identification**							**01100473**
Clostridium novyi biovar A		100.0	0.40	ARA 1				
01100500	**Low discrimination**							**01100500**
					COC	GRA		
Actinomyces meyeri		37.52	0.41	SUC 93	1	99		
Peptostreptococcus anaerobius		23.52	0.29	FRU 95	99	99		
Prevotella disiens		12.96	0.23		1	1		
01101000	**Low discrimination**							**01101000**
					COC	GRA	SPO	
Clostridium novyi biovar A		41.39	0.84	GAL 10	1	99	99	
Peptostreptococcus anaerobius		22.93	0.63		99	99	1	
Prevotella disiens		12.64	0.57		1	1	1	
01101001	**Low discrimination**							**01101001**
					COC	SPO		
Clostridium novyi biovar A		85.64	0.84	GAL 10	1	99		
Clostridium sordellii		13.13	0.53		1	99		
Peptostreptococcus anaerobius		0.48	0.29		99	1		
01101002	**Low discrimination**							**01101002**
					SPO			
Clostridium novyi biovar A		66.51	0.71	GAL 10	99			
Clostridium sporogenes		13.46	0.45		99			
Atopobium minutum		13.25	0.55		1			
01101003	**Very good identification**							**01101003**
					SPO			
Clostridium novyi biovar A		98.70	0.71	GAL 10	99			
Clostridium sordellii		0.87	0.19		99			
Atopobium minutum		0.20	0.21		1			
01101010	**Very good identification**							**01101010**
					COC	GRA	SPO	
Clostridium novyi biovar A		98.60	0.84	GAL 10	1	99	99	
Peptostreptococcus anaerobius		0.55	0.29		99	99	1	
Prevotella disiens		0.30	0.23		1	1	1	
01101011	**Very good identification**							**01101011**
					COC	SPO		
Clostridium novyi biovar A		99.84	0.84	GAL 10	1	99		
Clostridium sordellii		0.16	0.19		1	99		
Peptostreptococcus anaerobius		0.01	-.06		99	1		
01101012	**Very good identification**							**01101012**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		99.56	0.71	GAL 10	99		
Clostridium sporogenes		0.20	0.11		99		
Atopobium minutum		0.20	0.21		1		
01101013	**Very good identification**						**01101013**
Clostridium novyi biovar A		100.0	0.71	GAL 10			
01101020	**Low discrimination**						**01101020**
					COC	SPO	
Clostridium novyi biovar A		82.35	0.73	GAL 10	1	99	
Atopobium minutum		5.79	0.50		1	1	
Gemella morbillorum		4.88	0.43		99	1	
01101021	**Very good identification**						**01101021**
					SPO		
Clostridium novyi biovar A		99.23	0.73	GAL 10	99		
Clostridium sordellii		0.62	0.19		99		
Atopobium minutum		0.07	0.15		1		
01101022	**Low discrimination**						**01101022**
					GRA	SPO	
Clostridium novyi biovar A		74.90	0.61	GAL 10	99	99	
Atopobium minutum		19.89	0.47		99	1	
Pseudoflavonifractor capillosus		2.02	0.15		1	1	
01101023	**Very good identification**						**01101023**
					SPO		
Clostridium novyi biovar A		99.73	0.61	GAL 10	99		
Atopobium minutum		0.27	0.12		1		
01101030	**Very good identification**						**01101030**
					COC	SPO	
Clostridium novyi biovar A		99.85	0.73	GAL 10	1	99	
Atopobium minutum		0.07	0.15		1	1	
Gemella morbillorum		0.06	0.09		99	1	
01101031	**Very good identification**						**01101031**
Clostridium novyi biovar A		100.0	0.73	GAL 10			
01101032	**Very good identification**						**01101032**
					SPO		
Clostridium novyi biovar A		99.73	0.61	GAL 10	99		
Atopobium minutum		0.27	0.12		1		
01101033	**Very good identification**						**01101033**
Clostridium novyi biovar A		100.0	0.61	GAL 10			
01101040	**Low discrimination**						**01101040**
Clostridium novyi biovar A		85.81	0.81	GAL 10			
Clostridium botulinum biovar A		7.53	0.51				
Clostridium glycolicum		4.25	0.45				
01101041	**Very good identification**						**01101041**
Clostridium novyi biovar A		98.90	0.81	GAL 10			
Clostridium sordellii		0.95	0.29				
Clostridium botulinum biovar A		0.09	0.17				
01101042	**Very good identification**						**01101042**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		98.05	0.68	GAL 10	1	99	
Megasphaera elsdenii		0.54	0.22		99	1	
Clostridium botulinum biovar A		0.49	0.17		1	99	
01101043	**Very good identification**						**01101043**
Clostridium novyi biovar A		99.95	0.68	GAL 10			
Clostridium sordellii		0.06	-.05				
01101050	**Very good identification**						**01101050**
Clostridium novyi biovar A		99.84	0.81	GAL 10			
Clostridium botulinum biovar A		0.09	0.17				
Clostridium glycolicum		0.05	0.11				
01101051	**Very good identification**						**01101051**
Clostridium novyi biovar A		99.99	0.81	GAL 10			
Clostridium sordellii		0.01	-.05				
01101052	**Very good identification**						**01101052**
Clostridium novyi biovar A		100.0	0.68	GAL 10			
01101053	**Very good identification**						**01101053**
Clostridium novyi biovar A		100.0	0.68	GAL 10			
01101060	**Very good identification**						**01101060**
Clostridium novyi biovar A		99.16	0.70	GAL 10			
Clostridium botulinum biovar A		0.35	0.17				
Clostridium glycolicum		0.20	0.11				
01101061	**Very good identification**						**01101061**
Clostridium novyi biovar A		99.96	0.70	GAL 10			
Clostridium sordellii		0.04	-.05				
01101062	**Very good identification**						**01101062**
					SPO		
Clostridium novyi biovar A		99.60	0.58	GAL 10	99		
Atopobium minutum		0.40	0.12		1		
01101063	**Very good identification**						**01101063**
Clostridium novyi biovar A		100.0	0.58	GAL 10			
01101070	**Very good identification**						**01101070**
Clostridium novyi biovar A		100.0	0.70	GAL 10			
01101071	**Very good identification**						**01101071**
Clostridium novyi biovar A		100.0	0.70	GAL 10			
01101072	**Very good identification**						**01101072**
Clostridium novyi biovar A		100.0	0.58	GAL 10			
01101073	**Very good identification**						**01101073**
Clostridium novyi biovar A		100.0	0.58	GAL 10			
01101100	**Low discrimination**						**01101100**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		89.99	0.66	RAF 99	99	1	
Actinomyces meyeri		6.16	0.54		1	1	
Clostridium novyi biovar A		1.16	0.50		1	99	
01101101	**Low discrimination**						**01101101**
					COC	SPO	
Clostridium novyi biovar A		36.42	0.50	GAL 10	1	99	
Blautia hansenii		28.52	0.32	LAC 1	99	1	
Clostridium chauvoei		27.11	0.37		1	99	
01101102	**Low discrimination**						**01101102**
					COC	SPO	
Blautia hansenii		89.80	0.50	RAF 99	99	1	
Lactobacillus fermentum		5.85	0.48	BGL 10	1	1	
Clostridium novyi biovar A		1.84	0.37		1	99	
01101103	**Low discrimination**						**01101103**
					COC	SPO	
Clostridium novyi biovar A		44.06	0.37	GAL 10	1	99	
Clostridium chauvoei		32.80	0.24	LAC 1	1	99	
Blautia hansenii		21.72	0.15		99	1	
01101110	**Low discrimination**						**01101110**
					COC	SPO	
Clostridium novyi biovar A		54.12	0.50	GAL 10	1	99	
Blautia hansenii		42.38	0.32	LAC 1	99	1	
Actinomyces meyeri		2.90	0.20		1	1	
01101111	**Good identification**						**01101111**
					COC	SPO	
Clostridium novyi biovar A		98.48	0.50	GAL 10	1	99	
Blautia hansenii		0.78	-.03	LAC 1	99	1	
Clostridium chauvoei		0.74	0.03		1	99	
01101112	**Low discrimination**						**01101112**
					COC	SPO	
Clostridium novyi biovar A		65.57	0.37	GAL 10	1	99	
Blautia hansenii		32.33	0.15	LAC 1	99	1	
Lactobacillus fermentum		2.11	0.14		1	1	
01101113	**Good identification**						**01101113**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01101120	**Low discrimination**						**01101120**
					COC	GRA	
Prevotella bivia		77.72	0.53	MNS 99	1	1	
Blautia hansenii		19.29	0.50		99	99	
Actinomyces meyeri		1.62	0.39		1	99	
01101131	**Good identification**						**01101131**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01101141	**Good identification**						**01101141**
					COC	SPO	
Clostridium novyi biovar A		96.83	0.47	GAL 10	1	99	
Blautia hansenii		1.15	-.03	LAC 1	99	1	
Clostridium chauvoei		1.09	0.03		1	99	
01101150	**Good identification**						**01101150**
					COC	SPO	
Clostridium novyi biovar A		98.83	0.47	GAL 10	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		1.17	-.03	LAC 1	99	1		
01101151	**Good identification**							**01101151**
Clostridium novyi biovar A		100.0	0.47	GAL 10				
01101161	**Good identification**							**01101161**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01101170	**Good identification**							**01101170**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01101171	**Good identification**							**01101171**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01101200	**Low discrimination**							**01101200**
					COC	GRA	SPO	
Clostridium novyi biovar A		41.48	0.50	GAL 10	1	99	99	
Peptostreptococcus anaerobius		22.98	0.29	RHA 1	99	99	1	
Prevotella disiens		12.67	0.23		1	1	1	
01101201	**Low discrimination**							**01101201**
					COC	SPO		
Clostridium novyi biovar A		86.29	0.50	GAL 10	1	99		
Clostridium sordellii		13.23	0.19	RHA 1	1	99		
Peptostreptococcus anaerobius		0.48	-.06		99	1		
01101202	**Low discrimination**							**01101202**
					SPO			
Clostridium novyi biovar A		69.77	0.37	GAL 10	99			
Clostridium sporogenes		14.12	0.11	RHA 1	99			
Atopobium minutum		13.90	0.21		1			
01101203	**Good identification**							**01101203**
Clostridium novyi biovar A		100.0	0.37	GAL 10				
01101210	**Good identification**							**01101210**
					COC	SPO		
Clostridium novyi biovar A		99.44	0.50	GAL 10	1	99		
Peptostreptococcus anaerobius		0.56	-.06	RHA 1	99	1		
01101211	**Good identification**							**01101211**
Clostridium novyi biovar A		100.0	0.50	GAL 10				
01101212	**Good identification**							**01101212**
Clostridium novyi biovar A		100.0	0.37	GAL 10				
01101213	**Good identification**							**01101213**
Clostridium novyi biovar A		100.0	0.37	GAL 10				
01101220	**Low discrimination**							**01101220**
					COC	SPO		
Clostridium novyi biovar A		86.81	0.39	GAL 10	1	99		
Atopobium minutum		6.10	0.15	RHA 1	1	1		
Gemella morbillorum		5.14	0.09		99	1		
01101221	**Good identification**							**01101221**
Clostridium novyi biovar A		100.0	0.39	GAL 10				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
01101230	**Good identification**						**01101230**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01101231	**Good identification**						**01101231**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01101240	**Low discrimination**						**01101240**
Clostridium novyi biovar A		86.62	0.47	GAL 10			
Clostridium botulinum biovar A		7.60	0.17	RHA 1			
Clostridium glycolicum		4.29	0.11				
01101241	**Good identification**						**01101241**
Clostridium novyi biovar A		99.05	0.47	GAL 10			
Clostridium sordellii		0.95	-.05	RHA 1			
01101250	**Good identification**						**01101250**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01101251	**Good identification**						**01101251**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01101260	**Good identification**						**01101260**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01101261	**Good identification**						**01101261**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01101270	**Good identification**						**01101270**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01101271	**Good identification**						**01101271**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01101400	**Low discrimination**						**01101400**
					COC	SPO	
Gemella morbillorum		32.35	0.43	MNS 99	99	1	
Clostridium novyi biovar A		22.07	0.50	ARA 10	1	99	
Peptostreptococcus anaerobius		12.23	0.29		99	1	
01101401	**Low discrimination**						**01101401**
					COC	SPO	
Clostridium novyi biovar A		85.20	0.50	GAL 10	1	99	
Clostridium sordellii		13.06	0.19	ARA 1	1	99	
Gemella morbillorum		1.26	0.09		99	1	
01101402	**Low discrimination**						**01101402**
					SPO		
Lactobacillus fermentum		53.98	0.44	SUC 87	1		
Clostridium novyi biovar A		27.69	0.37	FRU 99	99		
Clostridium sporogenes		5.60	0.11		99		
01101403	**Good identification**						**01101403**
Clostridium novyi biovar A		100.0	0.37	GAL 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01101410	**Good identification**						**01101410**
					COC	SPO	
Clostridium novyi biovar A		98.00	0.50	GAL 10	1	99	
Gemella morbillorum		1.45	0.09	ARA 1	99	1	
Peptostreptococcus anaerobius		0.55	-.06		99	1	
01101411	**Good identification**						**01101411**
Clostridium novyi biovar A		100.0	0.50	GAL 10			
01101412	**Good identification**						**01101412**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01101413	**Good identification**						**01101413**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01101420	**Low discrimination**						**01101420**
					COC	SPO	
Clostridium novyi biovar A		40.90	0.39	GAL 10	1	99	
Gemella morbillorum		26.64	0.27	ARA 1	99	1	
Lactobacillus fermentum		18.76	0.36		1	1	
01101421	**Good identification**						**01101421**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01101422	**Low discrimination**						**01101422**
					SPO		
Lactobacillus fermentum		67.26	0.36	SUC 87	1		
Clostridium novyi biovar A		25.87	0.26	FRU 99	99		
Atopobium minutum		6.87	0.12		1		
01101430	**Good identification**						**01101430**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01101431	**Good identification**						**01101431**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01101440	**Low discrimination**						**01101440**
Clostridium novyi biovar A		84.99	0.47	GAL 10			
Clostridium botulinum biovar A		7.46	0.17	ARA 1			
Clostridium glycolicum		4.21	0.11				
01101441	**Good identification**						**01101441**
Clostridium novyi biovar A		99.05	0.47	GAL 10			
Clostridium sordellii		0.95	-.05	ARA 1			
01101450	**Good identification**						**01101450**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01101451	**Good identification**						**01101451**
Clostridium novyi biovar A		100.0	0.47	GAL 10			
01101460	**Good identification**						**01101460**
Clostridium novyi biovar A		100.0	0.36	GAL 10			

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01101461	**Good identification**							**01101461**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01101470	**Good identification**							**01101470**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01101471	**Good identification**							**01101471**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01101500	**Low discrimination**							**01101500**
					COC			
Blautia hansenii		87.71	0.53	RAF 99	99			
Actinomyces meyeri		8.50	0.44		1			
Lactobacillus fermentum		3.60	0.48		1			
01101502	**Low discrimination**							**01101502**
					COC			
Blautia hansenii		71.62	0.37	RAF 99	99			
Lactobacillus fermentum		26.45	0.48	BGL 10	1			
Actinomyces meyeri		1.93	0.18		1			
01101520	**Low discrimination**							**01101520**
					COC	GRA		
Blautia hansenii		67.91	0.37	RAF 99	99	99		
Prevotella bivia		15.66	0.19	NAG 10	1	1		
Lactobacillus fermentum		8.36	0.40		1	99		
01101522	**Low discrimination**							**01101522**
					COC			
Lactobacillus fermentum		51.74	0.40	SUC 87	1			
Blautia hansenii		46.71	0.20	FRU 99	99			
Actinomyces meyeri		1.55	0.03		1			
01102000	**Low discrimination**							**01102000**
					COC	GRA		
Megasphaera elsdenii		40.24	0.75	FRU 95	99	1		
Peptostreptococcus anaerobius		25.71	0.63		99	99		
Prevotella disiens		14.17	0.57		1	1		
01102001	**Low discrimination**							**01102001**
					COC	GRA		
Clostridium novyi biovar A		75.38	0.66	MAN 1	1	99		
Clostridium sordellii		7.28	0.32		1	99		
Megasphaera elsdenii		7.26	0.41		99	1		
01102002	**Low discrimination**							**01102002**
					COC	GRA		
Megasphaera elsdenii		59.98	0.59	FRU 95	99	1		
Clostridium sporogenes		22.23	0.45	BGL 10	1	99		
Clostridium novyi biovar A		9.99	0.53		1	99		
01102003	**Acceptable identification**							**01102003**
					COC	GRA		
Clostridium novyi biovar A		91.56	0.53	MAN 1	1	99		
Megasphaera elsdenii		5.55	0.25		99	1		
Clostridium sporogenes		2.06	0.11		1	99		
01102010	**Low discrimination**							**01102010**
					COC	GRA	SPO	
Clostridium novyi biovar A		80.62	0.66	MAN 1	1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		7.77	0.41		99	1	1	
Peptostreptococcus anaerobius		4.96	0.29		99	99	1	
01102011	**Very good identification**							**01102011**
					COC	GRA		
Clostridium novyi biovar A		99.74	0.66	MAN 1	1	99		
Megasphaera elsdenii		0.10	0.07		99	1		
Clostridium sordellii		0.10	-.03		1	99		
01102012	**Acceptable identification**							**01102012**
					COC	GRA		
Clostridium novyi biovar A		92.03	0.53	MAN 1	1	99		
Megasphaera elsdenii		5.58	0.25		99	1		
Clostridium sporogenes		2.07	0.11		1	99		
01102013	**Good identification**							**01102013**
Clostridium novyi biovar A		100.0	0.53	MAN 1				
01102020	**Low discrimination**							**01102020**
					COC	GRA	SPO	
Clostridium novyi biovar A		48.22	0.56	MAN 1	1	99	99	
Megasphaera elsdenii		18.58	0.41		99	1	1	
Peptostreptococcus anaerobius		11.87	0.29		99	99	1	
01102021	**Good identification**							**01102021**
					COC	GRA		
Clostridium novyi biovar A		98.98	0.56	MAN 1	1	99		
Clostridium sordellii		0.39	-.03		1	99		
Megasphaera elsdenii		0.39	0.07		99	1		
01102022	**Low discrimination**							**01102022**
					COC	GRA		
Clostridium novyi biovar A		67.10	0.43	MAN 1	1	99		
Megasphaera elsdenii		16.28	0.25		99	1		
Clostridium sporogenes		6.03	0.11		1	99		
01102023	**Good identification**							**01102023**
Clostridium novyi biovar A		100.0	0.43	MAN 1				
01102030	**Good identification**							**01102030**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.37	0.56	MAN 1	1	99	99	
Megasphaera elsdenii		0.39	0.07		99	1	1	
Peptostreptococcus anaerobius		0.25	-.06		99	99	1	
01102031	**Good identification**							**01102031**
Clostridium novyi biovar A		100.0	0.56	MAN 1				
01102032	**Good identification**							**01102032**
Clostridium novyi biovar A		100.0	0.43	MAN 1				
01102033	**Good identification**							**01102033**
Clostridium novyi biovar A		100.0	0.43	MAN 1				
01102040	**Low discrimination**							**01102040**
					COC	GRA		
Megasphaera elsdenii		81.17	0.72	FRU 95	99	1		
Clostridium novyi biovar A		8.51	0.63		1	99		
Clostridium botulinum biovar A		8.21	0.51		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01102041	**Low discrimination**						**01102041**
					COC	GRA	
Clostridium novyi biovar A		89.84	0.63	MAN 1	1	99	
Megasphaera elsdenii		8.66	0.38		99	1	
Clostridium botulinum biovar A		0.88	0.17		1	99	
01102042	**Low discrimination**						**01102042**
					COC	GRA	
Megasphaera elsdenii		84.41	0.56	FRU 95	99	1	
Clostridium novyi biovar A		14.06	0.50	BGL 10	1	99	
Clostridium botulinum biovar A		0.78	0.17		1	99	
01102043	**Acceptable identification**						**01102043**
					COC	GRA	
Clostridium novyi biovar A		94.28	0.50	MAN 1	1	99	
Megasphaera elsdenii		5.72	0.22		99	1	
01102050	**Good identification**						**01102050**
					COC	GRA	
Clostridium novyi biovar A		90.25	0.63	MAN 1	1	99	
Megasphaera elsdenii		8.70	0.38		99	1	
Clostridium botulinum biovar A		0.88	0.17		1	99	
01102051	**Very good identification**						**01102051**
					COC	GRA	
Clostridium novyi biovar A		99.90	0.63	MAN 1	1	99	
Megasphaera elsdenii		0.10	0.04		99	1	
01102052	**Acceptable identification**						**01102052**
					COC	GRA	
Clostridium novyi biovar A		94.28	0.50	MAN 1	1	99	
Megasphaera elsdenii		5.72	0.22		99	1	
01102053	**Good identification**						**01102053**
Clostridium novyi biovar A		100.0	0.50	MAN 1			
01102060	**Low discrimination**						**01102060**
					COC	GRA	
Clostridium novyi biovar A		69.06	0.53	MAN 1	1	99	
Megasphaera elsdenii		26.61	0.38		99	1	
Clostridium botulinum biovar A		2.69	0.17		1	99	
01102061	**Good identification**						**01102061**
					COC	GRA	
Clostridium novyi biovar A		99.61	0.53	MAN 1	1	99	
Megasphaera elsdenii		0.39	0.04		99	1	
01102062	**Low discrimination**						**01102062**
					COC	GRA	
Clostridium novyi biovar A		80.47	0.40	MAN 1	1	99	
Megasphaera elsdenii		19.53	0.22		99	1	
01102063	**Good identification**						**01102063**
Clostridium novyi biovar A		100.0	0.40	MAN 1			
01102070	**Good identification**						**01102070**
					COC	GRA	
Clostridium novyi biovar A		99.61	0.53	MAN 1	1	99	
Megasphaera elsdenii		0.39	0.04		99	1	
01102071	**Good identification**						**01102071**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.53	MAN 1				
01102072	**Good identification**							**01102072**
Clostridium novyi biovar A		100.0	0.40	MAN 1				
01102073	**Good identification**							**01102073**
Clostridium novyi biovar A		100.0	0.40	MAN 1				
01102100	**Low discrimination**							**01102100**
					COC	GRA		
Megasphaera elsdenii		39.14	0.41	FRU 95	99	1		
Peptostreptococcus anaerobius		25.01	0.29	LAC 1	99	99		
Prevotella disiens		13.78	0.23		1	1		
01102140	**Low discrimination**							**01102140**
					COC	GRA		
Megasphaera elsdenii		81.46	0.38	FRU 95	99	1		
Clostridium novyi biovar A		8.54	0.29	LAC 1	1	99		
Clostridium botulinum biovar A		8.24	0.17		1	99		
01102200	**Low discrimination**							**01102200**
					COC	GRA		
Megasphaera elsdenii		40.43	0.41	FRU 95	99	1		
Peptostreptococcus anaerobius		25.83	0.29	RHA 1	99	99		
Prevotella disiens		14.24	0.23		1	1		
01102240	**Low discrimination**							**01102240**
					COC	GRA		
Megasphaera elsdenii		82.26	0.38	FRU 95	99	1		
Clostridium novyi biovar A		8.62	0.29	RHA 1	1	99		
Clostridium botulinum biovar A		8.32	0.17		1	99		
01102400	**Low discrimination**							**01102400**
					COC	GRA		
Megasphaera elsdenii		39.92	0.41	FRU 95	99	1		
Peptostreptococcus anaerobius		25.50	0.29	ARA 1	99	99		
Prevotella disiens		14.06	0.23		1	1		
01102440	**Low discrimination**							**01102440**
					COC	GRA		
Megasphaera elsdenii		82.26	0.38	FRU 95	99	1		
Clostridium novyi biovar A		8.62	0.29	ARA 1	1	99		
Clostridium botulinum biovar A		8.32	0.17		1	99		
01103000	**Low discrimination**							**01103000**
					COC	GRA	SPO	
Clostridium novyi biovar A		30.58	0.50	GAL 10	1	99	99	
Megasphaera elsdenii		26.52	0.41	MAN 1	99	1	1	
Peptostreptococcus anaerobius		16.94	0.29		99	99	1	
01103001	**Low discrimination**							**01103001**
					COC	GRA		
Clostridium novyi biovar A		85.64	0.50	GAL 10	1	99		
Clostridium sordellii		13.13	0.19	MAN 1	1	99		
Megasphaera elsdenii		0.75	0.07		99	1		
01103002	**Low discrimination**							**01103002**
					COC	GRA		
Clostridium novyi biovar A		50.53	0.37	GAL 10	1	99		
Megasphaera elsdenii		27.59	0.25	MAN 1	99	1		
Clostridium sporogenes		10.22	0.11		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01103003	**Good identification**							**01103003**
Clostridium novyi biovar A		100.0	0.37	GAL 10				
01103010	**Good identification**							**01103010**
					COC	GRA	SPO	
Clostridium novyi biovar A		98.59	0.50	GAL 10	1	99	99	
Megasphaera elsdenii		0.86	0.07	MAN 1	99	1	1	
Peptostreptococcus anaerobius		0.55	-.06		99	99	1	
01103011	**Good identification**							**01103011**
Clostridium novyi biovar A		100.0	0.50	GAL 10				
01103012	**Good identification**							**01103012**
Clostridium novyi biovar A		100.0	0.37	GAL 10				
01103013	**Good identification**							**01103013**
Clostridium novyi biovar A		100.0	0.37	GAL 10				
01103020	**Low discrimination**							**01103020**
					COC	SPO		
Clostridium novyi biovar A		84.25	0.39	GAL 10	1	99		
Atopobium minutum		5.92	0.15	MAN 1	1	1		
Gemella morbillorum		4.99	0.09		99	1		
01103021	**Good identification**							**01103021**
Clostridium novyi biovar A		100.0	0.39	GAL 10				
01103030	**Good identification**							**01103030**
Clostridium novyi biovar A		100.0	0.39	GAL 10				
01103031	**Good identification**							**01103031**
Clostridium novyi biovar A		100.0	0.39	GAL 10				
01103040	**Low discrimination**							**01103040**
					COC	GRA		
Clostridium novyi biovar A		49.68	0.47	GAL 10	1	99		
Megasphaera elsdenii		43.08	0.38	MAN 1	99	1		
Clostridium botulinum biovar A		4.36	0.17		1	99		
01103041	**Good identification**							**01103041**
					COC	GRA		
Clostridium novyi biovar A		98.20	0.47	GAL 10	1	99		
Clostridium sordellii		0.94	-.05	MAN 1	1	99		
Megasphaera elsdenii		0.86	0.04		99	1		
01103050	**Good identification**							**01103050**
					COC	GRA		
Clostridium novyi biovar A		99.13	0.47	GAL 10	1	99		
Megasphaera elsdenii		0.87	0.04	MAN 1	99	1		
01103051	**Good identification**							**01103051**
Clostridium novyi biovar A		100.0	0.47	GAL 10				
01103060	**Good identification**							**01103060**
					COC	GRA		
Clostridium novyi biovar A		96.62	0.36	GAL 10	1	99		
Megasphaera elsdenii		3.39	0.04	MAN 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01103061	**Good identification**							**01103061**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01103070	**Good identification**							**01103070**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01103071	**Good identification**							**01103071**
Clostridium novyi biovar A		100.0	0.36	GAL 10				
01104000	**Low discrimination**							**01104000**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		34.06	0.63	XYL 1	99	99	1	
Prevotella disiens		18.77	0.57		1	1	1	
Clostridium glycolicum		18.29	0.62		1	99	99	
01104001	**Low discrimination**							**01104001**
					COC	SPO		
Clostridium novyi biovar A		78.83	0.66	XYL 1	1	99		
Clostridium sordellii		7.61	0.32		1	99		
Peptostreptococcus anaerobius		4.85	0.29		99	1		
01104002	**Low discrimination**							**01104002**
					COC	SPO		
Clostridium sporogenes		52.16	0.45	XYL 1	1	99		
Clostridium novyi biovar A		23.44	0.53		1	99		
Peptostreptococcus anaerobius		8.18	0.29		99	1		
01104003	**Good identification**							**01104003**
Clostridium novyi biovar A		96.94	0.53	XYL 1				
Clostridium sporogenes		2.18	0.11					
Clostridium sordellii		0.54	-.03					
01104010	**Low discrimination**							**01104010**
					COC	GRA	SPO	
Clostridium novyi biovar A		85.45	0.66	XYL 1	1	99	99	
Peptostreptococcus anaerobius		5.26	0.29		99	99	1	
Prevotella disiens		2.90	0.23		1	1	1	
01104011	**Very good identification**							**01104011**
					COC	SPO		
Clostridium novyi biovar A		99.84	0.66	XYL 1	1	99		
Clostridium sordellii		0.10	-.03		1	99		
Peptostreptococcus anaerobius		0.06	-.06		99	1		
01104012	**Good identification**							**01104012**
					COC	SPO		
Clostridium novyi biovar A		97.47	0.53	XYL 1	1	99		
Clostridium sporogenes		2.19	0.11		1	99		
Peptostreptococcus anaerobius		0.34	-.06		99	1		
01104013	**Good identification**							**01104013**
Clostridium novyi biovar A		100.0	0.53	XYL 1				
01104020	**Low discrimination**							**01104020**
					COC	GRA	SPO	
Clostridium novyi biovar A		49.85	0.56	XYL 1	1	99	99	
Peptostreptococcus anaerobius		12.27	0.29		99	99	1	
Prevotella disiens		6.76	0.23		1	1	1	

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01104021	**Good identification**						**01104021**
					COC	SPO	
Clostridium novyi biovar A		99.37	0.56	XYL 1	1	99	
Clostridium sordellii		0.39	-.03		1	99	
Peptostreptococcus anaerobius		0.25	-.06		99	1	
01104022	**Low discrimination**						**01104022**
					SPO		
Clostridium novyi biovar A		80.79	0.43	XYL 1	99		
Clostridium sporogenes		7.27	0.11		99		
Atopobium minutum		7.15	0.21		1		
01104023	**Good identification**						**01104023**
Clostridium novyi biovar A		100.0	0.43	XYL 1			
01104030	**Good identification**						**01104030**
					COC	SPO	
Clostridium novyi biovar A		99.75	0.56	XYL 1	1	99	
Peptostreptococcus anaerobius		0.25	-.06		99	1	
01104031	**Good identification**						**01104031**
Clostridium novyi biovar A		100.0	0.56	XYL 1			
01104032	**Good identification**						**01104032**
Clostridium novyi biovar A		100.0	0.43	XYL 1			
01104033	**Good identification**						**01104033**
Clostridium novyi biovar A		100.0	0.43	XYL 1			
01104040	**Very good identification**						**01104040**
Clostridium glycolicum		97.65	0.84	FRU 90			
Clostridium novyi biovar A		1.05	0.63				
Clostridium botulinum biovar A		1.01	0.51				
01104041	**Low discrimination**						**01104041**
Clostridium novyi biovar A		51.03	0.63	XYL 1			
Clostridium glycolicum		48.07	0.50				
Clostridium botulinum biovar A		0.50	0.17				
01104042	**Low discrimination**						**01104042**
					COC	GRA	
Clostridium glycolicum		82.21	0.50	FRU 90	1	99	
Clostridium novyi biovar A		15.40	0.50	BGL 1	1	99	
Megasphaera elsdenii		0.93	0.22		99	1	
01104043	**Acceptable identification**						**01104043**
Clostridium novyi biovar A		94.88	0.50	XYL 1			
Clostridium glycolicum		5.12	0.15				
01104050	**Low discrimination**						**01104050**
Clostridium novyi biovar A		51.19	0.63	XYL 1			
Clostridium glycolicum		48.22	0.50				
Clostridium botulinum biovar A		0.50	0.17				
01104051	**Very good identification**						**01104051**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		99.06	0.63	XYL 1			
Clostridium glycolicum		0.94	0.15				
01104052	**Acceptable identification**						**01104052**
Clostridium novyi biovar A		94.88	0.50	XYL 1			
Clostridium glycolicum		5.12	0.15				
01104053	**Good identification**						**01104053**
Clostridium novyi biovar A		100.0	0.50	XYL 1			
01104060	**Low discrimination**						**01104060**
Clostridium glycolicum		78.21	0.50	FRU 90			
Clostridium novyi biovar A		20.76	0.53	NAG 1			
Clostridium botulinum biovar A		0.81	0.17				
01104061	**Good identification**						**01104061**
Clostridium novyi biovar A		96.33	0.53	XYL 1			
Clostridium glycolicum		3.67	0.15				
01104062	**Low discrimination**						**01104062**
Clostridium novyi biovar A		82.26	0.40	XYL 1			
Clostridium glycolicum		17.74	0.15				
01104063	**Good identification**						**01104063**
Clostridium novyi biovar A		100.0	0.40	XYL 1			
01104070	**Good identification**						**01104070**
Clostridium novyi biovar A		96.33	0.53	XYL 1			
Clostridium glycolicum		3.67	0.15				
01104071	**Good identification**						**01104071**
Clostridium novyi biovar A		100.0	0.53	XYL 1			
01104072	**Good identification**						**01104072**
Clostridium novyi biovar A		100.0	0.40	XYL 1			
01104073	**Good identification**						**01104073**
Clostridium novyi biovar A		100.0	0.40	XYL 1			
01104100	**Low discrimination**						**01104100**
					COC	GRA	
Actinomyces meyeri		79.55	0.56	SUC 93	1	99	
Peptostreptococcus anaerobius		6.42	0.29	FRU 95	99	99	
Prevotella disiens		3.54	0.23		1	1	
01104140	**Good identification**						**01104140**
					SPO		
Clostridium glycolicum		96.59	0.50	FRU 90	99		
Actinomyces meyeri		1.19	0.22	LAC 1	1		
Clostridium novyi biovar A		1.04	0.29		99		
01104240	**Good identification**						**01104240**
Clostridium glycolicum		97.74	0.50	FRU 90			
Clostridium novyi biovar A		1.05	0.29	RHA 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		1.01	0.17				
01104440	**Good identification**						**01104440**
Clostridium glycolicum		97.74	0.50	FRU 90			
Clostridium novyi biovar A		1.05	0.29	ARA 1			
Clostridium botulinum biovar A		1.01	0.17				
01104500	**Good identification**						**01104500**
					COC		
Actinomyces meyeri		98.28	0.46	SUC 93	1		
Blautia hansenii		1.40	0.06	FRU 95	99		
Peptostreptococcus anaerobius		0.32	-.06		99		
01105000	**Low discrimination**						**01105000**
					COC	SPO	
Clostridium glycolicum		33.04	0.45	FRU 90	1	99	
Actinomyces meyeri		25.99	0.49	GAL 10	1	1	
Gemella morbillorum		14.80	0.43	SOR 95	99	1	
01105001	**Low discrimination**						**01105001**
Clostridium novyi biovar A		81.12	0.50	GAL 10			
Clostridium sordellii		12.44	0.19	XYL 1			
Clostridium glycolicum		2.68	0.11				
01105002	**Low discrimination**						**01105002**
					SPO		
Lactobacillus fermentum		33.80	0.41	SUC 87	1		
Clostridium novyi biovar A		29.52	0.37	FRU 99	99		
Actinomyces meyeri		13.31	0.23		1		
01105003	**Good identification**						**01105003**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01105010	**Acceptable identification**						**01105010**
					SPO		
Clostridium novyi biovar A		92.64	0.50	GAL 10	99		
Clostridium glycolicum		3.06	0.11	XYL 1	99		
Actinomyces meyeri		2.41	0.14		1		
01105011	**Good identification**						**01105011**
Clostridium novyi biovar A		100.0	0.50	GAL 10			
01105012	**Good identification**						**01105012**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01105013	**Good identification**						**01105013**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
01105021	**Good identification**						**01105021**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01105030	**Good identification**						**01105030**
Clostridium novyi biovar A		100.0	0.39	GAL 10			
01105031	**Good identification**						**01105031**
Clostridium novyi biovar A		100.0	0.39	GAL 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01105040	**Very good identification**						**01105040**
Clostridium glycolicum		98.77	0.67	FRU 90			
Clostridium novyi biovar A		1.06	0.47	GAL 10			
Clostridium botulinum biovar A		0.09	0.17				
01105041	**Low discrimination**						**01105041**
Clostridium novyi biovar A		51.24	0.47	GAL 10			
Clostridium glycolicum		48.27	0.33	XYL 1			
Clostridium sordellii		0.49	-.05				
01105050	**Low discrimination**						**01105050**
Clostridium novyi biovar A		51.49	0.47	GAL 10			
Clostridium glycolicum		48.51	0.33	XYL 1			
01105051	**Good identification**						**01105051**
Clostridium novyi biovar A		99.06	0.47	GAL 10			
Clostridium glycolicum		0.94	-.01	XYL 1			
01105061	**Good identification**						**01105061**
Clostridium novyi biovar A		96.33	0.36	GAL 10			
Clostridium glycolicum		3.67	-.01	XYL 1			
01105070	**Good identification**						**01105070**
Clostridium novyi biovar A		96.33	0.36	GAL 10			
Clostridium glycolicum		3.67	-.01	XYL 1			
01105071	**Good identification**						**01105071**
Clostridium novyi biovar A		100.0	0.36	GAL 10			
01105100	**Low discrimination**						**01105100**
					COC		
Blautia hansenii		56.14	0.53	RAF 99	99		
Actinomyces meyeri		42.24	0.59		1		
Lactobacillus fermentum		1.35	0.44		1		
01105102	**Low discrimination**						**01105102**
					COC		
Blautia hansenii		70.11	0.37	RAF 99	99		
Lactobacillus fermentum		15.20	0.44	BGL 10	1		
Actinomyces meyeri		14.68	0.33		1		
01105120	**Low discrimination**						**01105120**
					COC	GRA	
Blautia hansenii		44.92	0.37	RAF 99	99	99	
Actinomyces meyeri		41.48	0.44	NAG 10	1	99	
Prevotella bivia		10.36	0.19		1	1	
01105400	**Low discrimination**						**01105400**
					COC		
Actinomyces meyeri		59.49	0.38	SUC 93	1		
Lactobacillus fermentum		18.68	0.41	FRU 95	1		
Gemella morbillorum		15.06	0.27		99		
01105402	**Acceptable identification**						**01105402**
Lactobacillus fermentum		91.03	0.41	SUC 87			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		8.97	0.12	FRU 99				
01105500	**Low discrimination**							**01105500**
					COC			
Actinomyces meyeri		48.40	0.49	SUC 93	1			
Blautia hansenii		45.40	0.40	FRU 95	99			
Lactobacillus fermentum		6.20	0.44		1			
01105502	**Low discrimination**							**01105502**
					COC			
Lactobacillus fermentum		48.66	0.44	SUC 87	1			
Blautia hansenii		39.60	0.24	FRU 99	99			
Actinomyces meyeri		11.75	0.23		1			
01105522	**Low discrimination**							**01105522**
					COC			
Lactobacillus fermentum		70.75	0.36	SUC 87	1			
Blautia hansenii		19.19	0.07	FRU 99	99			
Actinomyces meyeri		6.99	0.08		1			
01106000	**Low discrimination**							**01106000**
					COC	GRA		
Megasphaera elsdenii		34.87	0.41	FRU 95	99	1		
Peptostreptococcus anaerobius		22.28	0.29	XYL 1	99	99		
Prevotella disiens		12.28	0.23		1	1		
01106040	**Low discrimination**							**01106040**
					COC	GRA		
Clostridium glycolicum		88.94	0.50	FRU 90	1	99		
Megasphaera elsdenii		9.10	0.38	MAN 1	99	1		
Clostridium novyi biovar A		0.95	0.29		1	99		
01110000	**Low discrimination**							**01110000**
					COC	GRA	SPO	
Prevotella disiens		55.59	0.91		1	1	1	
Megasphaera elsdenii		30.30	0.97		99	1	1	
Clostridium sporogenes		3.44	0.74		1	99	99	
01110001	**Low discrimination**							**01110001**
					GRA	SPO		
Clostridium novyi biovar A		41.53	0.84	FRU 10	99	99		
Clostridium sordellii		36.08	0.66		99	99		
Prevotella disiens		12.68	0.57		1	1		
01110002	**Low discrimination**							**01110002**
					COC	GRA	SPO	
Megasphaera elsdenii		57.30	0.81	BGL 10	99	1	1	
Clostridium sporogenes		19.53	0.66		1	99	99	
Prevotella disiens		9.56	0.57		1	1	1	
01110003	**Low discrimination**							**01110003**
					COC	GRA		
Clostridium novyi biovar A		81.85	0.71	FRU 10	1	99		
Megasphaera elsdenii		8.58	0.47		99	1		
Clostridium sordellii		4.07	0.32		1	99		
01110010	**Low discrimination**							**01110010**
					COC	GRA	SPO	
Clostridium novyi biovar A		64.45	0.84	FRU 10	1	99	99	
Prevotella disiens		19.68	0.57		1	1	1	
Megasphaera elsdenii		10.73	0.63		99	1	1	
01110011	**Very good identification**							**01110011**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium novyi biovar A		98.60	0.84	FRU 10	99	99	
Clostridium sordellii		0.87	0.32		99	99	
Prevotella disiens		0.30	0.23		1	1	
01110012	**Low discrimination**						**01110012**
					COC	GRA	
Clostridium novyi biovar A		85.27	0.71	FRU 10	1	99	
Megasphaera elsdenii		8.93	0.47		99	1	
Clostridium sporogenes		3.05	0.32		1	99	
01110013	**Very good identification**						**01110013**
					COC	GRA	
Clostridium novyi biovar A		99.81	0.71	FRU 10	1	99	
Megasphaera elsdenii		0.11	0.12		99	1	
Clostridium sordellii		0.05	-.03		1	99	
01110020	**Low discrimination**						**01110020**
					GRA	SPO	
Prevotella disiens		26.72	0.57	NAG 1	1	1	
Clostridium novyi biovar A		21.88	0.73		99	99	
Pseudoramibacter alactolyticus		16.28	0.53		99	1	
01110021	**Good identification**						**01110021**
					GRA	SPO	
Clostridium novyi biovar A		93.39	0.73	FRU 10	99	99	
Clostridium sordellii		3.28	0.32		99	99	
Prevotella disiens		1.15	0.23		1	1	
01110022	**Low discrimination**						**01110022**
					SPO		
Clostridium novyi biovar A		34.04	0.61	FRU 10	99		
Atopobium minutum		25.04	0.54		1		
Pseudoramibacter alactolyticus		15.95	0.37		1		
01110023	**Very good identification**						**01110023**
					SPO		
Clostridium novyi biovar A		98.05	0.61	FRU 10	99		
Atopobium minutum		0.73	0.20		1		
Pseudoramibacter alactolyticus		0.46	0.03		1		
01110030	**Very good identification**						**01110030**
					GRA	SPO	
Clostridium novyi biovar A		96.49	0.73	FRU 10	99	99	
Prevotella disiens		1.19	0.23		1	1	
Pseudoramibacter alactolyticus		0.73	0.19		99	1	
01110031	**Very good identification**						**01110031**
Clostridium novyi biovar A		99.97	0.73	FRU 10			
Clostridium sordellii		0.04	-.03				
01110032	**Very good identification**						**01110032**
					SPO		
Clostridium novyi biovar A		98.25	0.61	FRU 10	99		
Atopobium minutum		0.73	0.20		1		
Pseudoramibacter alactolyticus		0.47	0.03		1		
01110033	**Very good identification**						**01110033**
Clostridium novyi biovar A		100.0	0.61	FRU 10			
01110040	**Low discrimination**						**01110040**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Megasphaera elsdenii		71.62	0.94		99	1		
Clostridium glycolicum		17.46	0.78		1	99		
Clostridium novyi biovar A		4.35	0.81		1	99		
01110041	**Low discrimination**							**01110041**
					COC	GRA		
Clostridium novyi biovar A		78.35	0.81	FRU 10	1	99		
Megasphaera elsdenii		13.04	0.60		99	1		
Clostridium sordellii		4.25	0.42		1	99		
01110042	**Low discrimination**							**01110042**
					COC	SPO	GRA	
Megasphaera elsdenii		86.62	0.78	BGL 10	99	1	1	
Clostridium novyi biovar A		8.35	0.68		1	99	99	
Pseudoramibacter alactolyticus		2.08	0.39		1	1	99	
01110043	**Low discrimination**							**01110043**
					COC	GRA		
Clostridium novyi biovar A		89.83	0.68	FRU 10	1	99		
Megasphaera elsdenii		9.41	0.44		99	1		
Clostridium sordellii		0.28	0.08		1	99		
01110050	**Low discrimination**							**01110050**
					COC	GRA		
Clostridium novyi biovar A		80.62	0.81	FRU 10	1	99		
Megasphaera elsdenii		13.42	0.60		99	1		
Clostridium glycolicum		3.27	0.44		1	99		
01110051	**Very good identification**							**01110051**
					COC	GRA		
Clostridium novyi biovar A		99.72	0.81	FRU 10	1	99		
Megasphaera elsdenii		0.17	0.26		99	1		
Clostridium sordellii		0.06	0.08		1	99		
01110052	**Low discrimination**							**01110052**
					COC	SPO	GRA	
Clostridium novyi biovar A		90.00	0.68	FRU 10	1	99	99	
Megasphaera elsdenii		9.43	0.44		99	1	1	
Pseudoramibacter alactolyticus		0.23	0.05		1	1	99	
01110053	**Very good identification**							**01110053**
					COC	GRA		
Clostridium novyi biovar A		99.89	0.68	FRU 10	1	99		
Megasphaera elsdenii		0.11	0.09		99	1		
01110060	**Low discrimination**							**01110060**
					COC	SPO	GRA	
Clostridium novyi biovar A		44.73	0.70	FRU 10	1	99	99	
Megasphaera elsdenii		29.78	0.60		99	1	1	
Pseudoramibacter alactolyticus		12.48	0.43		1	1	99	
01110061	**Very good identification**							**01110061**
					COC	SPO	GRA	
Clostridium novyi biovar A		98.61	0.70	FRU 10	1	99	99	
Megasphaera elsdenii		0.66	0.26		99	1	1	
Pseudoramibacter alactolyticus		0.28	0.09		1	1	99	
01110062	**Low discrimination**							**01110062**
					COC	SPO	GRA	
Clostridium novyi biovar A		61.67	0.58	FRU 10	1	99	99	
Megasphaera elsdenii		25.85	0.44		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		10.83	0.26		1	1	99	
01110063	**Very good identification**							**01110063**
					COC	SPO	GRA	
Clostridium novyi biovar A		99.40	0.58	FRU 10	1	99	99	
Megasphaera elsdenii		0.42	0.09		99	1	1	
Pseudoramibacter alactolyticus		0.18	-.08		1	1	99	
01110070	**Very good identification**							**01110070**
					COC	GRA		
Clostridium novyi biovar A		95.59	0.70	FRU 10	1	99		
Clostridium botulinum biovar B		3.34	0.31		1	99		
Megasphaera elsdenii		0.64	0.26		99	1		
01110071	**Very good identification**							**01110071**
Clostridium novyi biovar A		99.97	0.70	FRU 10				
Clostridium botulinum biovar B		0.04	-.03					
01110072	**Very good identification**							**01110072**
					COC	GRA		
Clostridium novyi biovar A		99.21	0.58	FRU 10	1	99		
Megasphaera elsdenii		0.42	0.09		99	1		
Clostridium botulinum biovar B		0.20	-.03		1	99		
01110073	**Very good identification**							**01110073**
Clostridium novyi biovar A		100.0	0.58	FRU 10				
01110100	**Low discrimination**							**01110100**
					COC	GRA		
Actinomyces meyeri		54.54	0.73	SUC 93	1	99		
Prevotella disiens		24.54	0.57		1	1		
Megasphaera elsdenii		13.38	0.63		99	1		
01110101	**Low discrimination**							**01110101**
					SPO			
Clostridium novyi biovar A		32.17	0.50	FRU 10	99			
Clostridium sordellii		27.95	0.32	LAC 1	99			
Actinomyces meyeri		21.83	0.39		1			
01110102	**Low discrimination**							**01110102**
					COC	SPO	GRA	
Actinomyces meyeri		34.95	0.47	SUC 93	1	1	99	
Megasphaera elsdenii		30.79	0.47	BGL 3	99	1	1	
Clostridium sporogenes		10.49	0.32		1	99	99	
01110103	**Low discrimination**							**01110103**
					COC	SPO	GRA	
Clostridium novyi biovar A		74.56	0.37	FRU 10	1	99	99	
Actinomyces meyeri		8.87	0.13	LAC 1	1	1	99	
Megasphaera elsdenii		7.81	0.12		99	1	1	
01110110	**Low discrimination**							**01110110**
					GRA	SPO		
Clostridium novyi biovar A		44.75	0.50	FRU 10	99	99		
Actinomyces meyeri		30.36	0.39	LAC 1	99	1		
Prevotella disiens		13.66	0.23		1	1		
01110111	**Good identification**							**01110111**
					SPO			
Clostridium novyi biovar A		98.46	0.50	FRU 10	99			
Clostridium sordellii		0.86	-.03	LAC 1	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		0.68	0.05		1			
01110112	**Low discrimination**							**01110112**
					COC	SPO	GRA	
Clostridium novyi biovar A		77.43	0.37	FRU 10	1	99	99	
Actinomyces meyeri		9.21	0.13	LAC 1	1	1	99	
Megasphaera elsdenii		8.11	0.12		99	1	1	
01110113	**Good identification**							**01110113**
Clostridium novyi biovar A		100.0	0.37	FRU 10				
01110120	**Low discrimination**							**01110120**
					GRA			
Prevotella bivia		67.54	0.50	MNS 99	1			
Actinomyces meyeri		28.31	0.58	FRU 10	99			
Prevotella disiens		0.94	0.23		1			
01110121	**Low discrimination**							**01110121**
					GRA	SPO		
Clostridium novyi biovar A		43.26	0.39	FRU 10	99	99		
Prevotella bivia		38.20	0.15	LAC 1	1	1		
Actinomyces meyeri		16.01	0.24		99	1		
01110130	**Low discrimination**							**01110130**
					GRA	SPO		
Clostridium novyi biovar A		44.39	0.39	FRU 10	99	99		
Prevotella bivia		39.19	0.15	LAC 1	1	1		
Actinomyces meyeri		16.43	0.24		99	1		
01110131	**Good identification**							**01110131**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01110140	**Low discrimination**							**01110140**
					COC	SPO	GRA	
Megasphaera elsdenii		68.54	0.60	LAC 1	99	1	1	
Clostridium glycolicum		16.71	0.44		1	99	99	
Actinomyces meyeri		4.23	0.39		1	1	99	
01110141	**Low discrimination**							**01110141**
					COC	GRA		
Clostridium novyi biovar A		78.64	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.09	0.26	LAC 1	99	1		
Clostridium sordellii		4.27	0.08		1	99		
01110142	**Low discrimination**							**01110142**
					COC	SPO	GRA	
Megasphaera elsdenii		84.89	0.44	LAC 1	99	1	1	
Clostridium novyi biovar A		8.18	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.04	0.05		1	1	99	
01110150	**Low discrimination**							**01110150**
					COC	GRA		
Clostridium novyi biovar A		80.97	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.48	0.26	LAC 1	99	1		
Clostridium glycolicum		3.29	0.10		1	99		
01110151	**Good identification**							**01110151**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01110161	**Good identification**							**01110161**

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Clostridium novyi biovar A		100.0	0.36	FRU 10				
01110170	**Good identification**							**01110170**
Clostridium novyi biovar A		96.62	0.36	FRU 10				
Clostridium botulinum biovar B		3.38	-.03	LAC 1				
01110171	**Good identification**							**01110171**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01110200	**Low discrimination**							**01110200**
					COC	GRA	SPO	
Prevotella disiens		55.52	0.57	RHA 1	1	1	1	
Megasphaera elsdenii		30.26	0.63		99	1	1	
Clostridium sporogenes		3.44	0.40		1	99	99	
01110201	**Low discrimination**							**01110201**
					GRA	SPO		
Clostridium novyi biovar A		41.72	0.50	FRU 10	99	99		
Clostridium sordellii		36.24	0.32	RHA 1	99	99		
Prevotella disiens		12.74	0.23		1	1		
01110202	**Low discrimination**							**01110202**
					COC	GRA	SPO	
Megasphaera elsdenii		56.61	0.47	RHA 1	99	1	1	
Clostridium sporogenes		19.29	0.32	BGL 10	1	99	99	
Prevotella disiens		9.44	0.23		1	1	1	
01110203	**Low discrimination**							**01110203**
					COC	GRA		
Clostridium novyi biovar A		84.02	0.37	FRU 10	1	99		
Megasphaera elsdenii		8.80	0.12	RHA 1	99	1		
Clostridium sordellii		4.18	-.03		1	99		
01110210	**Low discrimination**							**01110210**
					COC	GRA	SPO	
Clostridium novyi biovar A		64.92	0.50	FRU 10	1	99	99	
Prevotella disiens		19.82	0.23	RHA 1	1	1	1	
Megasphaera elsdenii		10.80	0.29		99	1	1	
01110211	**Good identification**							**01110211**
Clostridium novyi biovar A		99.13	0.50	FRU 10				
Clostridium sordellii		0.87	-.03	RHA 1				
01110212	**Low discrimination**							**01110212**
					COC	GRA		
Clostridium novyi biovar A		87.68	0.37	FRU 10	1	99		
Megasphaera elsdenii		9.19	0.12	RHA 1	99	1		
Clostridium sporogenes		3.13	-.02		1	99		
01110213	**Good identification**							**01110213**
Clostridium novyi biovar A		100.0	0.37	FRU 10				
01110221	**Good identification**							**01110221**
Clostridium novyi biovar A		96.61	0.39	FRU 10				
Clostridium sordellii		3.39	-.03	RHA 1				
01110230	**Good identification**							**01110230**
Clostridium novyi biovar A		100.0	0.39	FRU 10				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01110231	**Good identification**							**01110231**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01110240	**Low discrimination**							**01110240**
					COC	GRA		
Megasphaera elsdenii		71.68	0.60	RHA 1	99	1		
Clostridium glycolicum		17.47	0.44		1	99		
Clostridium novyi biovar A		4.35	0.47		1	99		
01110241	**Low discrimination**							**01110241**
					COC	GRA		
Clostridium novyi biovar A		79.29	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.19	0.26	RHA 1	99	1		
Clostridium sordellii		4.31	0.08		1	99		
01110242	**Low discrimination**							**01110242**
					COC	SPO	GRA	
Megasphaera elsdenii		87.13	0.44	RHA 1	99	1	1	
Clostridium novyi biovar A		8.40	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.09	0.05		1	1	99	
01110250	**Low discrimination**							**01110250**
					COC	GRA		
Clostridium novyi biovar A		81.65	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.59	0.26	RHA 1	99	1		
Clostridium glycolicum		3.31	0.10		1	99		
01110251	**Good identification**							**01110251**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01110260	**Low discrimination**							**01110260**
					COC	SPO	GRA	
Clostridium novyi biovar A		45.92	0.36	FRU 10	1	99	99	
Megasphaera elsdenii		30.56	0.26	RHA 1	99	1	1	
Pseudoramibacter alactolyticus		12.81	0.09		1	1	99	
01110261	**Good identification**							**01110261**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01110270	**Good identification**							**01110270**
Clostridium novyi biovar A		96.62	0.36	FRU 10				
Clostridium botulinum biovar B		3.38	-.03	RHA 1				
01110271	**Good identification**							**01110271**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01110300	**Low discrimination**							**01110300**
					COC	GRA		
Actinomyces meyeri		53.90	0.39	SUC 93	1	99		
Prevotella disiens		24.26	0.23	RHA 1	1	1		
Megasphaera elsdenii		13.22	0.29		99	1		
01110400	**Low discrimination**							**01110400**
					COC	GRA		
Prevotella disiens		48.94	0.57	ARA 1	1	1		
Megasphaera elsdenii		26.68	0.63		99	1		
Actinomyces meyeri		6.80	0.52		1	99		
01110401	**Low discrimination**							**01110401**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Clostridium novyi biovar A		40.44	0.50	FRU 10	99	99		
Clostridium sordellii		35.13	0.32	ARA 1	99	99		
Prevotella disiens		12.35	0.23		1	1		
01110402	**Low discrimination**							**01110402**
					COC	SPO	GRA	
Megasphaera elsdenii		49.68	0.47	ARA 1	99	1	1	
Clostridium sporogenes		16.93	0.32	BGL 10	1	99	99	
Lactobacillus fermentum		9.34	0.44		1	1	99	
01110403	**Low discrimination**							**01110403**
					COC	GRA		
Clostridium novyi biovar A		84.02	0.37	FRU 10	1	99		
Megasphaera elsdenii		8.80	0.12	ARA 1	99	1		
Clostridium sordellii		4.18	-.03		1	99		
01110410	**Low discrimination**							**01110410**
					COC	GRA	SPO	
Clostridium novyi biovar A		61.87	0.50	FRU 10	1	99	99	
Prevotella disiens		18.89	0.23	ARA 1	1	1	1	
Megasphaera elsdenii		10.30	0.29		99	1	1	
01110411	**Good identification**							**01110411**
Clostridium novyi biovar A		99.13	0.50	FRU 10				
Clostridium sordellii		0.87	-.03	ARA 1				
01110412	**Low discrimination**							**01110412**
					COC	GRA		
Clostridium novyi biovar A		87.68	0.37	FRU 10	1	99		
Megasphaera elsdenii		9.19	0.12	ARA 1	99	1		
Clostridium sporogenes		3.13	-.02		1	99		
01110413	**Good identification**							**01110413**
Clostridium novyi biovar A		100.0	0.37	FRU 10				
01110420	**Low discrimination**							**01110420**
					COC	GRA		
Actinomyces meyeri		26.84	0.37	SUC 93	1	99		
Gemella morbillorum		17.33	0.33	NAG 12	99	99		
Prevotella disiens		14.31	0.23		1	1		
01110421	**Acceptable identification**							**01110421**
					SPO			
Clostridium novyi biovar A		94.50	0.39	FRU 10	99			
Clostridium sordellii		3.32	-.03	ARA 1	99			
Actinomyces meyeri		2.19	0.03		1			
01110422	**Low discrimination**							**01110422**
					SPO			
Lactobacillus fermentum		45.08	0.36	SUC 87	1			
Clostridium novyi biovar A		17.34	0.26	GAL 99	99			
Atopobium minutum		12.75	0.20		1			
01110430	**Good identification**							**01110430**
					SPO			
Clostridium novyi biovar A		97.74	0.39	FRU 10	99			
Actinomyces meyeri		2.26	0.03	ARA 1	1			
01110431	**Good identification**							**01110431**

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Clostridium novyi biovar A		100.0	0.39	FRU 10				
01110440	**Low discrimination**							**01110440**
					COC	GRA		
Megasphaera elsdenii		71.33	0.60	ARA 1	99	1		
Clostridium glycolicum		17.39	0.44		1	99		
Clostridium novyi biovar A		4.33	0.47		1	99		
01110441	**Low discrimination**							**01110441**
					COC	GRA		
Clostridium novyi biovar A		79.29	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.19	0.26	ARA 1	99	1		
Clostridium sordellii		4.31	0.08		1	99		
01110442	**Low discrimination**							**01110442**
					COC	SPO	GRA	
Megasphaera elsdenii		87.13	0.44	ARA 1	99	1	1	
Clostridium novyi biovar A		8.40	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.09	0.05		1	1	99	
01110450	**Low discrimination**							**01110450**
					COC	GRA		
Clostridium novyi biovar A		81.65	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.59	0.26	ARA 1	99	1		
Clostridium glycolicum		3.31	0.10		1	99		
01110451	**Good identification**							**01110451**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01110460	**Low discrimination**							**01110460**
					COC	SPO	GRA	
Clostridium novyi biovar A		45.20	0.36	FRU 10	1	99	99	
Megasphaera elsdenii		30.09	0.26	ARA 1	99	1	1	
Pseudoramibacter alactolyticus		12.61	0.09		1	1	99	
01110461	**Good identification**							**01110461**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01110470	**Good identification**							**01110470**
Clostridium novyi biovar A		96.62	0.36	FRU 10				
Clostridium botulinum biovar B		3.38	-.03	ARA 1				
01110471	**Good identification**							**01110471**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01110500	**Good identification**							**01110500**
					GRA			
Actinomyces meyeri		93.63	0.63	SUC 93	99			
Lactobacillus fermentum		3.13	0.48		99			
Prevotella disiens		1.70	0.23		1			
01110502	**Low discrimination**							**01110502**
					COC	GRA		
Lactobacillus fermentum		50.78	0.48	SUC 87	1	99		
Actinomyces meyeri		46.98	0.37	GAL 99	1	99		
Megasphaera elsdenii		1.67	0.12		99	1		
01110520	**Low discrimination**							**01110520**
					GRA			
Actinomyces meyeri		84.88	0.48	SUC 93	99			

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Prevotella bivia		8.18	0.15	NAG 12	1		
Lactobacillus fermentum		6.94	0.40		99		
01110522	**Low discrimination**						**01110522**
Lactobacillus fermentum		72.54	0.40	SUC 87			
Actinomyces meyeri		27.46	0.22	GAL 99			
01111000	**Low discrimination**						**01111000**
					GRA		
Prevotella intermedia		37.24	0.73		1		
Lactobacillus fermentum		14.70	0.79		99		
Prevotella disiens		13.05	0.57		1		
01111001	**Low discrimination**						**01111001**
					GRA	SPO	
Clostridium sordellii		53.41	0.53	IND 99	99	99	
Clostridium novyi biovar A		38.71	0.67		99	99	
Prevotella intermedia		3.07	0.39		1	1	
01111002	**Low discrimination**						**01111002**
					SPO		
Lactobacillus fermentum		77.32	0.79	SUC 87	1		
Atopobium minutum		7.29	0.54		1		
Clostridium novyi biovar A		4.41	0.54		99		
01111003	**Low discrimination**						**01111003**
					SPO		
Clostridium novyi biovar A		77.26	0.54	FRU 10	99		
Lactobacillus fermentum		13.69	0.44	GAL 10	1		
Clostridium sordellii		6.10	0.19		99		
01111010	**Low discrimination**						**01111010**
					GRA	SPO	
Clostridium novyi biovar A		83.02	0.67	FRU 10	99	99	
Prevotella intermedia		6.58	0.39	GAL 10	1	1	
Lactobacillus fermentum		2.60	0.44		99	1	
01111011	**Very good identification**						**01111011**
					GRA	SPO	
Clostridium novyi biovar A		98.55	0.67	FRU 10	99	99	
Clostridium sordellii		1.37	0.19	GAL 10	99	99	
Prevotella intermedia		0.08	0.05		1	1	
01111012	**Low discrimination**						**01111012**
					SPO		
Clostridium novyi biovar A		82.28	0.54	FRU 10	99		
Lactobacillus fermentum		14.58	0.44	GAL 10	1		
Atopobium minutum		1.38	0.20		1		
01111013	**Good identification**						**01111013**
Clostridium novyi biovar A		100.0	0.54	FRU 10			
01111020	**Low discrimination**						**01111020**
					SPO		
Lactobacillus fermentum		49.12	0.70	SUC 87	1		
Actinomyces meyeri		14.86	0.51		1		
Clostridium novyi biovar A		11.90	0.57		99		
01111021	**Low discrimination**						**01111021**
					SPO		
Clostridium novyi biovar A		88.66	0.57	FRU 10	99		

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Clostridium sordellii		4.94	0.19	GAL 10	99			
Lactobacillus fermentum		3.70	0.36		1			
01111022	**Low discrimination**							**01111022**
					SPO			
Lactobacillus fermentum		85.90	0.70	SUC 87	1			
Atopobium minutum		8.10	0.46		1			
Clostridium novyi biovar A		3.67	0.44		99			
01111023	**Low discrimination**							**01111023**
					SPO			
Clostridium novyi biovar A		79.45	0.44	FRU 10	99			
Lactobacillus fermentum		18.78	0.36	GAL 10	1			
Atopobium minutum		1.77	0.12		1			
01111030	**Good identification**							**01111030**
					SPO			
Clostridium novyi biovar A		93.46	0.57	FRU 10	99			
Lactobacillus fermentum		3.90	0.36	GAL 10	1			
Actinomyces meyeri		1.18	0.17		1			
01111031	**Very good identification**							**01111031**
Clostridium novyi biovar A		100.0	0.57	FRU 10				
01111032	**Low discrimination**							**01111032**
					SPO			
Clostridium novyi biovar A		79.45	0.44	FRU 10	99			
Lactobacillus fermentum		18.78	0.36	GAL 10	1			
Atopobium minutum		1.77	0.12		1			
01111033	**Good identification**							**01111033**
Clostridium novyi biovar A		100.0	0.44	FRU 10				
01111040	**Low discrimination**							**01111040**
					COC	GRA		
Clostridium glycolicum		58.38	0.62	GAL 10	1	99		
Megasphaera elsdenii		21.77	0.60	XYL 95	99	1		
Clostridium novyi biovar A		14.54	0.64		1	99		
01111041	**Low discrimination**							**01111041**
Clostridium novyi biovar A		87.48	0.64	FRU 10				
Clostridium sordellii		7.54	0.29	GAL 10				
Clostridium glycolicum		3.55	0.28					
01111042	**Low discrimination**							**01111042**
					COC	SPO	GRA	
Clostridium novyi biovar A		39.93	0.51	FRU 10	1	99	99	
Megasphaera elsdenii		37.66	0.44	GAL 10	99	1	1	
Lactobacillus fermentum		10.62	0.44		1	1	99	
01111043	**Good identification**							**01111043**
					COC	GRA		
Clostridium novyi biovar A		98.58	0.51	FRU 10	1	99		
Megasphaera elsdenii		0.94	0.09	GAL 10	99	1		
Clostridium sordellii		0.49	-.05		1	99		
01111050	**Good identification**							**01111050**
					COC	GRA		
Clostridium novyi biovar A		94.47	0.64	FRU 10	1	99		
Clostridium glycolicum		3.83	0.28	GAL 10	1	99		

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Megasphaera elsdenii		1.43	0.26		99	1	
01111051	**Very good identification**						**01111051**
Clostridium novyi biovar A		99.91	0.64	FRU 10			
Clostridium sordellii		0.09	-.05	GAL 10			
01111052	**Good identification**						**01111052**
					COC	GRA	
Clostridium novyi biovar A		99.06	0.51	FRU 10	1	99	
Megasphaera elsdenii		0.94	0.09	GAL 10	99	1	
01111053	**Good identification**						**01111053**
Clostridium novyi biovar A		100.0	0.51	FRU 10			
01111060	**Low discrimination**						**01111060**
					SPO		
Clostridium novyi biovar A		71.06	0.54	FRU 10	99		
Clostridium glycolicum		11.53	0.28	GAL 10	99		
Lactobacillus fermentum		4.45	0.36		1		
01111061	**Good identification**						**01111061**
Clostridium novyi biovar A		99.65	0.54	FRU 10			
Clostridium sordellii		0.35	-.05	GAL 10			
01111062	**Low discrimination**						**01111062**
					SPO		
Clostridium novyi biovar A		66.40	0.41	FRU 10	99		
Lactobacillus fermentum		23.54	0.36	GAL 10	1		
Clostridium perfringens		4.25	0.11		84		
01111063	**Good identification**						**01111063**
Clostridium novyi biovar A		100.0	0.41	FRU 10			
01111070	**Good identification**						**01111070**
Clostridium novyi biovar A		99.68	0.54	FRU 10			
Clostridium botulinum biovar B		0.32	-.03	GAL 10			
01111071	**Good identification**						**01111071**
Clostridium novyi biovar A		100.0	0.54	FRU 10			
01111072	**Good identification**						**01111072**
Clostridium novyi biovar A		100.0	0.41	FRU 10			
01111073	**Good identification**						**01111073**
Clostridium novyi biovar A		100.0	0.41	FRU 10			
01111100	**Low discrimination**						**01111100**
					GRA		
Actinomyces meyeri		62.79	0.76	SUC 93	99		
Lactobacillus fermentum		34.63	0.82		99		
Fusobacterium mortiferum		0.92	0.54		1		
01111101	**Low discrimination**						**01111101**
					SPO		
Clostridium chauvoei		48.09	0.44	SUC 99	99		
Actinomyces meyeri		28.17	0.42	MNS 90	1		
Lactobacillus fermentum		15.54	0.48		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01111102	**Good identification**					**01111102**
					GRA	
Lactobacillus fermentum		94.08	0.82	SUC 87	99	
Actinomyces meyeri		5.28	0.50		99	
Fusobacterium mortiferum		0.28	0.37		1	
01111103	**Low discrimination**					**01111103**
					SPO	
Lactobacillus fermentum		61.08	0.48	SUC 87	1	
Clostridium chauvoei		33.36	0.31	URE 1	99	
Actinomyces meyeri		3.43	0.16		1	
01111110	**Low discrimination**					**01111110**
					SPO	
Actinomyces meyeri		59.11	0.42	SUC 93	1	
Lactobacillus fermentum		32.60	0.48	MLZ 1	1	
Clostridium novyi biovar A		6.45	0.33		99	
01111112	**Acceptable identification**					**01111112**
					SPO	
Lactobacillus fermentum		91.66	0.48	SUC 87	1	
Actinomyces meyeri		5.14	0.16	MLZ 1	1	
Clostridium novyi biovar A		3.20	0.20		99	
01111120	**Low discrimination**					**01111120**
					GRA	
Lactobacillus fermentum		46.46	0.74	SUC 87	99	
Actinomyces meyeri		34.46	0.61		99	
Prevotella bivia		9.67	0.37		1	
01111121	**Low discrimination**					**01111121**
					SPO	
Clostridium chauvoei		57.46	0.35	SUC 99	99	
Lactobacillus fermentum		18.57	0.40	MNS 90	1	
Actinomyces meyeri		13.77	0.27		1	
01111122	**Good identification**					**01111122**
Lactobacillus fermentum		92.49	0.74	SUC 87		
Clostridium perfringens		3.98	0.38			
Actinomyces meyeri		2.12	0.35			
01111123	**Low discrimination**					**01111123**
					SPO	
Lactobacillus fermentum		61.90	0.40	SUC 87	1	
Clostridium chauvoei		33.81	0.22	URE 1	99	
Clostridium perfringens		2.67	0.04		84	
01111130	**Low discrimination**					**01111130**
					GRA	
Lactobacillus fermentum		43.64	0.40	SUC 87	99	
Actinomyces meyeri		32.37	0.27	MLZ 1	99	
Prevotella bivia		9.09	0.03		1	
01111132	**Acceptable identification**					**01111132**
					SPO	
Lactobacillus fermentum		93.52	0.40	SUC 87	1	
Clostridium perfringens		4.03	0.04	MLZ 1	84	
Clostridium novyi biovar A		2.45	0.10		99	
01111140	**Low discrimination**					**01111140**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		48.23	0.42	SUC 93	1	
Lactobacillus fermentum		26.60	0.48	SOR 1	1	
Clostridium glycolicum		14.10	0.28		99	
01111142	**Low discrimination**					**01111142**
					SPO	
Lactobacillus fermentum		88.65	0.48	SUC 87	1	
Actinomyces meyeri		4.97	0.16	SOR 1	1	
Clostridium novyi biovar A		2.06	0.17		99	
01111162	**Low discrimination**					**01111162**
Lactobacillus fermentum		55.18	0.40	SUC 87		
Clostridium perfringens		44.82	0.26	SOR 1		
01111200	**Low discrimination**					**01111200**
					GRA	
Prevotella intermedia		36.47	0.39	RHA 1	1	
Lactobacillus fermentum		14.40	0.44		99	
Prevotella disiens		12.78	0.23		1	
01111202	**Low discrimination**					**01111202**
Lactobacillus fermentum		72.94	0.44	SUC 87		
Actinomyces odontolyticus		7.24	0.20	RHA 1		
Atopobium minutum		6.88	0.20			
01111220	**Low discrimination**					**01111220**
					SPO	
Lactobacillus fermentum		50.00	0.36	SUC 87	1	
Actinomyces meyeri		15.13	0.17	RHA 1	1	
Clostridium novyi biovar A		12.11	0.23		99	
01111222	**Low discrimination**					**01111222**
Lactobacillus fermentum		84.35	0.36	SUC 87		
Atopobium minutum		7.96	0.12	RHA 1		
Actinomyces odontolyticus		4.09	0.07			
01111300	**Low discrimination**					**01111300**
Actinomyces meyeri		62.50	0.42	SUC 93		
Lactobacillus fermentum		34.47	0.48	RHA 1		
Actinomyces odontolyticus		1.08	0.15			
01111302	**Acceptable identification**					**01111302**
Lactobacillus fermentum		93.00	0.48	SUC 87		
Actinomyces meyeri		5.22	0.16	RHA 1		
Actinomyces odontolyticus		1.79	0.12			
01111320	**Low discrimination**					**01111320**
					GRA	
Lactobacillus fermentum		46.66	0.40	SUC 87	99	
Actinomyces meyeri		34.61	0.27	RHA 1	99	
Prevotella bivia		9.72	0.03		1	
01111322	**Acceptable identification**					**01111322**
					GRA	
Lactobacillus fermentum		93.68	0.40	SUC 87	99	
Clostridium perfringens		4.04	0.04	RHA 1	99	
Prevotella oralis		2.28	-.03		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01111400	**Low discrimination**					**01111400**
					COC	
Lactobacillus fermentum		77.90	0.79	SUC 87	1	
Actinomyces meyeri		14.40	0.55		1	
Gemella morbillorum		4.10	0.45		99	
01111401	**Low discrimination**					**01111401**
					SPO	
Lactobacillus fermentum		49.22	0.44	SUC 87	1	
Clostridium sordellii		21.93	0.19	URE 1	99	
Clostridium novyi biovar A		15.90	0.33		99	
01111402	**Very good identification**					**01111402**
Lactobacillus fermentum		99.13	0.79	SUC 87		
Actinomyces meyeri		0.57	0.29			
Atopobium minutum		0.09	0.20			
01111403	**Acceptable identification**					**01111403**
					SPO	
Lactobacillus fermentum		94.61	0.44	SUC 87	1	
Clostridium novyi biovar A		5.39	0.20	URE 1	99	
01111410	**Low discrimination**					**01111410**
					SPO	
Lactobacillus fermentum		63.05	0.44	SUC 87	1	
Clostridium novyi biovar A		20.36	0.33	MLZ 1	99	
Actinomyces meyeri		11.66	0.21		1	
01111412	**Acceptable identification**					**01111412**
					SPO	
Lactobacillus fermentum		94.61	0.44	SUC 87	1	
Clostridium novyi biovar A		5.39	0.20	MLZ 1	99	
01111420	**Good identification**					**01111420**
					COC	
Lactobacillus fermentum		91.09	0.70	SUC 87	1	
Actinomyces meyeri		6.89	0.40		1	
Gemella morbillorum		1.60	0.29		99	
01111421	**Low discrimination**					**01111421**
					SPO	
Lactobacillus fermentum		75.88	0.36	SUC 87	1	
Clostridium novyi biovar A		18.38	0.23	URE 1	99	
Actinomyces meyeri		5.74	0.06		1	
01111422	**Very good identification**					**01111422**
Lactobacillus fermentum		99.63	0.70	SUC 87		
Actinomyces meyeri		0.23	0.15			
Atopobium minutum		0.10	0.12			
01111423	**Good identification**					**01111423**
					SPO	
Lactobacillus fermentum		95.90	0.36	SUC 87	1	
Clostridium novyi biovar A		4.10	0.10	URE 1	99	
01111430	**Low discrimination**					**01111430**
					SPO	
Lactobacillus fermentum		75.88	0.36	SUC 87	1	
Clostridium novyi biovar A		18.38	0.23	MLZ 1	99	
Actinomyces meyeri		5.74	0.06		1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01111432	**Good identification**							**01111432**
					SPO			
Lactobacillus fermentum		95.90	0.36	SUC 87	1			
Clostridium novyi biovar A		4.10	0.10	MLZ 1	99			
01111440	**Low discrimination**							**01111440**
					COC	SPO	GRA	
Lactobacillus fermentum		37.52	0.44	SUC 87	1	1	99	
Clostridium glycolicum		32.44	0.28	SOR 1	1	99	99	
Megasphaera elsdenii		12.10	0.26		99	1	1	
01111442	**Acceptable identification**							**01111442**
					COC	SPO	GRA	
Lactobacillus fermentum		93.13	0.44	SUC 87	1	1	99	
Clostridium novyi biovar A		3.54	0.17	SOR 1	1	99	99	
Megasphaera elsdenii		3.34	0.09		99	1	1	
01111460	**Low discrimination**							**01111460**
					SPO			
Lactobacillus fermentum		80.83	0.36	SUC 87	1			
Clostridium novyi biovar A		13.05	0.20	SOR 1	99			
Actinomyces meyeri		6.12	0.06		1			
01111462	**Good identification**							**01111462**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01111500	**Low discrimination**							**01111500**
					COC			
Lactobacillus fermentum		68.66	0.82	SUC 87	1			
Actinomyces meyeri		31.12	0.66		1			
Blautia hansenii		0.17	0.19		99			
01111501	**Low discrimination**							**01111501**
					SPO			
Lactobacillus fermentum		67.36	0.48	SUC 87	1			
Actinomyces meyeri		30.53	0.31	URE 1	1			
Clostridium chauvoei		2.11	0.09		99			
01111502	**Very good identification**							**01111502**
					COC			
Lactobacillus fermentum		98.59	0.82	SUC 87	1			
Actinomyces meyeri		1.38	0.40		1			
Blautia hansenii		0.03	0.03		99			
01111503	**Good identification**							**01111503**
Lactobacillus fermentum		98.62	0.48	SUC 87				
Actinomyces meyeri		1.38	0.06	URE 1				
01111510	**Low discrimination**							**01111510**
Lactobacillus fermentum		68.30	0.48	SUC 87				
Actinomyces meyeri		30.96	0.31	MLZ 1				
Bifidobacterium longum subsp. longum		0.74	0.07					
01111512	**Good identification**							**01111512**
Lactobacillus fermentum		97.57	0.48	SUC 87				
Actinomyces meyeri		1.37	0.06	MLZ 1				
Bifidobacterium longum subsp. longum		1.06	0.07					
01111520	**Low discrimination**							**01111520**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Lactobacillus fermentum		83.96	0.74	SUC 87	99	
Actinomyces meyeri		15.57	0.51		99	
Prevotella bivia		0.18	0.03		1	
01111521	**Low discrimination**					**01111521**
					SPO	
Lactobacillus fermentum		82.19	0.40	SUC 87	1	
Actinomyces meyeri		15.24	0.17	URE 1	1	
Clostridium chauvoei		2.57	0.01		99	
01111522	**Very good identification**					**01111522**
Lactobacillus fermentum		99.31	0.74	SUC 87		
Actinomyces meyeri		0.57	0.25			
Bifidobacterium longum subsp. longum		0.08	0.14			
01111523	**Good identification**					**01111523**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01111530	**Low discrimination**					**01111530**
Lactobacillus fermentum		76.00	0.40	SUC 87		
Actinomyces meyeri		14.09	0.17	MLZ 1		
Bifidobacterium longum subsp. longum		9.91	0.18			
01111532	**Low discrimination**					**01111532**
Lactobacillus fermentum		88.47	0.40	SUC 87		
Bifidobacterium longum subsp. longum		11.54	0.18	MLZ 1		
01111540	**Low discrimination**					**01111540**
Lactobacillus fermentum		68.81	0.48	SUC 87		
Actinomyces meyeri		31.19	0.31	SOR 1		
01111542	**Good identification**					**01111542**
Lactobacillus fermentum		98.62	0.48	SUC 87		
Actinomyces meyeri		1.38	0.06	SOR 1		
01111560	**Low discrimination**					**01111560**
Lactobacillus fermentum		84.36	0.40	SUC 87		
Actinomyces meyeri		15.64	0.17	SOR 1		
01111562	**Good identification**					**01111562**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01111600	**Low discrimination**					**01111600**
					COC	
Lactobacillus fermentum		79.17	0.44	SUC 87	1	
Actinomyces meyeri		14.64	0.21	RHA 1	1	
Gemella morbillorum		4.16	0.11		99	
01111602	**Good identification**					**01111602**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01111620	**Acceptable identification**					**01111620**
Lactobacillus fermentum		92.97	0.36	SUC 87		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		7.03	0.06	RHA 1				
01111622	**Good identification**							**01111622**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01111700	**Low discrimination**							**01111700**
Lactobacillus fermentum		68.81	0.48	SUC 87				
Actinomyces meyeri		31.19	0.31	RHA 1				
01111702	**Good identification**							**01111702**
Lactobacillus fermentum		98.62	0.48	SUC 87				
Actinomyces meyeri		1.38	0.06	RHA 1				
01111720	**Low discrimination**							**01111720**
Lactobacillus fermentum		84.36	0.40	SUC 87				
Actinomyces meyeri		15.64	0.17	RHA 1				
01111722	**Good identification**							**01111722**
Lactobacillus fermentum		100.0	0.40	SUC 87				
01112000	**Good identification**							**01112000**
					COC	GRA		
Megasphaera elsdenii		92.57	0.97		99	1		
Pseudoramibacter alactolyticus		3.19	0.61		1	99		
Eubacterium limosum		2.03	0.60		1	99		
01112001	**Low discrimination**							**01112001**
					COC	GRA		
Megasphaera elsdenii		83.86	0.63	URE 1	99	1		
Clostridium novyi biovar A		5.09	0.50		1	99		
Clostridium sordellii		4.42	0.32		1	99		
01112002	**Very good identification**							**01112002**
					COC	SPO	GRA	
Megasphaera elsdenii		95.83	0.81	BGL 10	99	1	1	
Pseudoramibacter alactolyticus		3.30	0.45		1	1	99	
Clostridium sporogenes		0.33	0.32		1	99	99	
01112003	**Low discrimination**							**01112003**
					COC	SPO	GRA	
Megasphaera elsdenii		87.79	0.47	URE 1	99	1	1	
Clostridium novyi biovar A		8.46	0.37	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		3.02	0.11		1	1	99	
01112010	**Low discrimination**							**01112010**
					COC	GRA		
Megasphaera elsdenii		55.55	0.63	MLZ 1	99	1		
Clostridium difficile		36.73	0.53		1	99		
Clostridium novyi biovar A		3.37	0.50		1	99		
01112011	**Low discrimination**							**01112011**
					COC	GRA		
Clostridium novyi biovar A		77.46	0.50	FRU 10	1	99		
Megasphaera elsdenii		12.89	0.29	MAN 1	99	1		
Clostridium difficile		8.52	0.19		1	99		
01112012	**Low discrimination**							**01112012**
					COC	GRA		
Megasphaera elsdenii		83.72	0.47	MLZ 1	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		8.07	0.37	BGL 10	1	99		
Clostridium difficile		5.03	0.19		1	99		
01112013	**Acceptable identification**							**01112013**
					COC	GRA		
Clostridium novyi biovar A		90.52	0.37	FRU 10	1	99		
Megasphaera elsdenii		9.48	0.12	MAN 1	99	1		
01112020	**Low discrimination**							**01112020**
					COC	GRA		
Megasphaera elsdenii		59.51	0.63	NAG 1	99	1		
Pseudoramibacter alactolyticus		35.81	0.48		1	99		
Eubacterium limosum		1.30	0.26		1	99		
01112021	**Low discrimination**							**01112021**
					COC	SPO	GRA	
Clostridium novyi biovar A		47.59	0.39	FRU 10	1	99	99	
Megasphaera elsdenii		31.68	0.29	MAN 1	99	1	1	
Pseudoramibacter alactolyticus		19.06	0.14		1	1	99	
01112022	**Low discrimination**							**01112022**
					COC	SPO	GRA	
Megasphaera elsdenii		60.62	0.47	NAG 1	99	1	1	
Pseudoramibacter alactolyticus		36.48	0.32	BGL 10	1	1	99	
Clostridium novyi biovar A		1.46	0.26		1	99	99	
01112030	**Low discrimination**							**01112030**
					COC	GRA		
Clostridium novyi biovar A		39.90	0.39	FRU 10	1	99		
Megasphaera elsdenii		26.56	0.29	MAN 1	99	1		
Clostridium difficile		17.56	0.19		1	99		
01112031	**Good identification**							**01112031**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01112040	**Very good identification**							**01112040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.32	0.94		99	1	1	
Pseudoramibacter alactolyticus		1.27	0.51		1	1	99	
Clostridium glycolicum		0.24	0.44		1	99	99	
01112041	**Good identification**							**01112041**
					COC	SPO	GRA	
Megasphaera elsdenii		92.65	0.60	URE 1	99	1	1	
Clostridium novyi biovar A		5.62	0.47		1	99	99	
Pseudoramibacter alactolyticus		1.20	0.17		1	1	99	
01112042	**Very good identification**							**01112042**
					COC	SPO	GRA	
Megasphaera elsdenii		98.60	0.78	BGL 10	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.35		1	1	99	
Clostridium novyi biovar A		0.10	0.34		1	99	99	
01112043	**Acceptable identification**							**01112043**
					COC	SPO	GRA	
Megasphaera elsdenii		90.15	0.44	URE 1	99	1	1	
Clostridium novyi biovar A		8.69	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		1.17	0.00		1	1	99	
01112050	**Low discrimination**							**01112050**
					COC	GRA		
Megasphaera elsdenii		88.55	0.60	MLZ 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		5.38	0.47		1	99		
Clostridium difficile		4.62	0.31		1	99		
01112051	**Low discrimination**							**01112051**
					COC	GRA		
Clostridium novyi biovar A		85.10	0.47	FRU 10	1	99		
Megasphaera elsdenii		14.16	0.26	MAN 1	99	1		
Clostridium difficile		0.74	-.03		1	99		
01112052	**Low discrimination**							**01112052**
					COC	SPO	GRA	
Megasphaera elsdenii		89.76	0.44	MLZ 1	99	1	1	
Clostridium novyi biovar A		8.65	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		1.16	0.00		1	1	99	
01112060	**Low discrimination**							**01112060**
					COC	SPO	GRA	
Megasphaera elsdenii		80.22	0.60	NAG 1	99	1	1	
Pseudoramibacter alactolyticus		18.10	0.38		1	1	99	
Clostridium novyi biovar A		1.22	0.36		1	99	99	
01112061	**Low discrimination**							**01112061**
					COC	SPO	GRA	
Clostridium novyi biovar A		55.07	0.36	FRU 10	1	99	99	
Megasphaera elsdenii		36.66	0.26	MAN 1	99	1	1	
Pseudoramibacter alactolyticus		8.27	0.04		1	1	99	
01112062	**Low discrimination**							**01112062**
					COC	SPO	GRA	
Megasphaera elsdenii		80.01	0.44	NAG 1	99	1	1	
Pseudoramibacter alactolyticus		18.06	0.22	BGL 10	1	1	99	
Clostridium novyi biovar A		1.93	0.23		1	99	99	
01112070	**Low discrimination**							**01112070**
					COC	SPO	GRA	
Clostridium novyi biovar A		53.03	0.36	FRU 10	1	99	99	
Megasphaera elsdenii		35.30	0.26	MAN 1	99	1	1	
Pseudoramibacter alactolyticus		7.97	0.04		1	1	99	
01112071	**Good identification**							**01112071**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01112100	**Low discrimination**							**01112100**
					COC	GRA		
Megasphaera elsdenii		85.98	0.63	LAC 1	99	1		
Propionibacterium propionicum		3.56	0.49		1	99		
Actinomyces meyeri		3.54	0.39		1	99		
01112102	**Low discrimination**							**01112102**
					COC	GRA		
Megasphaera elsdenii		62.09	0.47	LAC 1	99	1		
Propionibacterium propionicum		34.66	0.52	BGL 10	1	99		
Pseudoramibacter alactolyticus		2.14	0.11		1	99		
01112103	**Low discrimination**							**01112103**
					COC	GRA		
Propionibacterium propionicum		86.00	0.36	NIT 86	1	99		
Megasphaera elsdenii		14.01	0.12	SUC 99	99	1		
01112140	**Good identification**							**01112140**
					COC	GRA		
Megasphaera elsdenii		92.71	0.60	LAC 1	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		5.75	0.49		1	99		
Pseudoramibacter alactolyticus		1.20	0.17		1	99		
01112142	**Low discrimination**							**01112142**
					COC	GRA		
Megasphaera elsdenii		54.05	0.44	LAC 1	99	1		
Propionibacterium propionicum		45.26	0.52	BGL 10	1	99		
Pseudoramibacter alactolyticus		0.70	0.00		1	99		
01112143	**Acceptable identification**							**01112143**
					COC	GRA		
Propionibacterium propionicum		90.21	0.36	NIT 86	1	99		
Megasphaera elsdenii		9.79	0.09	SUC 99	99	1		
01112200	**Good identification**							**01112200**
					COC	GRA		
Megasphaera elsdenii		92.62	0.63	RHA 1	99	1		
Pseudoramibacter alactolyticus		3.19	0.27		1	99		
Eubacterium limosum		2.03	0.26		1	99		
01112202	**Good identification**							**01112202**
					COC	SPO	GRA	
Megasphaera elsdenii		96.35	0.47	RHA 1	99	1	1	
Pseudoramibacter alactolyticus		3.32	0.11	BGL 10	1	1	99	
Clostridium sporogenes		0.33	-.02		1	99	99	
01112240	**Very good identification**							**01112240**
					COC	SPO	GRA	
Megasphaera elsdenii		98.43	0.60	RHA 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.17		1	1	99	
Clostridium glycolicum		0.24	0.10		1	99	99	
01112242	**Good identification**							**01112242**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	RHA 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
01112400	**Good identification**							**01112400**
					COC	GRA		
Megasphaera elsdenii		92.23	0.63	ARA 1	99	1		
Pseudoramibacter alactolyticus		3.18	0.27		1	99		
Eubacterium limosum		2.02	0.26		1	99		
01112402	**Good identification**							**01112402**
					COC	SPO	GRA	
Megasphaera elsdenii		96.35	0.47	ARA 1	99	1	1	
Pseudoramibacter alactolyticus		3.32	0.11	BGL 10	1	1	99	
Clostridium sporogenes		0.33	-.02		1	99	99	
01112440	**Very good identification**							**01112440**
					COC	SPO	GRA	
Megasphaera elsdenii		98.43	0.60	ARA 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.17		1	1	99	
Clostridium glycolicum		0.24	0.10		1	99	99	
01112442	**Good identification**							**01112442**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	ARA 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
01113000	**Low discrimination**							**01113000**
					COC	GRA		
Megasphaera elsdenii		83.77	0.63	GAL 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		4.43	0.39		1	1		
Pseudoramibacter alactolyticus		2.89	0.27		1	99		
01113002	**Low discrimination**							**01113002**
					COC	GRA		
Megasphaera elsdenii		78.15	0.47	GAL 1	99	1		
Lactobacillus fermentum		14.69	0.44	BGL 10	1	99		
Pseudoramibacter alactolyticus		2.69	0.11		1	99		
01113022	**Low discrimination**							**01113022**
					COC	GRA		
Lactobacillus fermentum		69.55	0.36	SUC 87	1	99		
Megasphaera elsdenii		11.21	0.12	MAN 1	99	1		
Pseudoramibacter alactolyticus		6.75	-.02		1	99		
01113040	**Very good identification**							**01113040**
					COC	SPO	GRA	
Megasphaera elsdenii		95.40	0.60	GAL 1	99	1	1	
Clostridium glycolicum		2.58	0.28		1	99	99	
Pseudoramibacter alactolyticus		1.23	0.17		1	1	99	
01113042	**Good identification**							**01113042**
					COC	SPO	GRA	
Megasphaera elsdenii		96.89	0.44	GAL 1	99	1	1	
Pseudoramibacter alactolyticus		1.25	0.00	BGL 10	1	1	99	
Clostridium novyi biovar A		1.04	0.17		1	99	99	
01113100	**Low discrimination**							**01113100**
Actinomyces meyeri		41.30	0.42	SUC 93				
Propionibacterium propionicum		27.65	0.49	MAN 1				
Lactobacillus fermentum		22.78	0.48					
01113102	**Low discrimination**							**01113102**
Propionibacterium propionicum		62.57	0.52	NIT 86				
Lactobacillus fermentum		34.37	0.48	SUC 99				
Actinomyces meyeri		1.93	0.16					
01113103	**Good identification**							**01113103**
Propionibacterium propionicum		95.24	0.36	NIT 86				
Lactobacillus fermentum		4.76	0.14	SUC 99				
01113120	**Low discrimination**							**01113120**
					GRA			
Lactobacillus fermentum		45.88	0.40	SUC 87	99			
Actinomyces meyeri		34.03	0.27	MAN 1	99			
Prevotella bivia		9.55	0.03		1			
01113122	**Acceptable identification**							**01113122**
Lactobacillus fermentum		91.06	0.40	SUC 87				
Propionibacterium propionicum		5.02	0.18	MAN 1				
Clostridium perfringens		3.92	0.04					
01113140	**Low discrimination**							**01113140**
					COC	GRA		
Propionibacterium propionicum		84.30	0.49	NIT 86	1	99		
Megasphaera elsdenii		13.73	0.26	SUC 99	99	1		
Actinomyces meyeri		1.27	0.08		1	99		
01113142	**Good identification**							**01113142**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
	Propionibacterium propionicum	98.27	0.52	NIT 86	1	99	
	Megasphaera elsdenii	1.19	0.09	SUC 99	99	1	
	Lactobacillus fermentum	0.55	0.14		1	99	
01113143	**Good identification**						**01113143**
	Propionibacterium propionicum	100.0	0.36	NIT 86			
01113400	**Low discrimination**						**01113400**
					COC	GRA	
	Lactobacillus fermentum	56.71	0.44	SUC 87	1	99	
	Megasphaera elsdenii	27.43	0.29	MAN 1	99	1	
	Actinomyces meyeri	10.49	0.21		1	99	
01113402	**Acceptable identification**						**01113402**
					COC	GRA	
	Lactobacillus fermentum	94.90	0.44	SUC 87	1	99	
	Megasphaera elsdenii	5.10	0.12	MAN 1	99	1	
01113420	**Acceptable identification**						**01113420**
	Lactobacillus fermentum	92.97	0.36	SUC 87			
	Actinomyces meyeri	7.03	0.06	MAN 1			
01113422	**Good identification**						**01113422**
	Lactobacillus fermentum	100.0	0.36	SUC 87			
01113500	**Low discrimination**						**01113500**
	Lactobacillus fermentum	68.23	0.48	SUC 87			
	Actinomyces meyeri	30.93	0.31	MAN 1			
	Propionibacterium propionicum	0.84	0.15				
01113502	**Good identification**						**01113502**
	Lactobacillus fermentum	96.86	0.48	SUC 87			
	Propionibacterium propionicum	1.78	0.18	MAN 1			
	Actinomyces meyeri	1.36	0.06				
01113520	**Low discrimination**						**01113520**
	Lactobacillus fermentum	84.36	0.40	SUC 87			
	Actinomyces meyeri	15.64	0.17	MAN 1			
01113522	**Good identification**						**01113522**
	Lactobacillus fermentum	100.0	0.40	SUC 87			
01114000	**Low discrimination**						**01114000**
					GRA	SPO	
	Clostridium glycolicum	74.57	0.78	SOR 95	99	99	
	Actinomyces meyeri	9.17	0.68		99	1	
	Prevotella disiens	8.50	0.57		1	1	
01114001	**Low discrimination**						**01114001**
	Clostridium glycolicum	49.19	0.44	SOR 95			
	Clostridium novyi biovar A	18.37	0.50	URE 1			
	Clostridium sordellii	15.96	0.32				
01114002	**Low discrimination**						**01114002**
					COC	SPO	GRA

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium glycolicum		37.61	0.44	SOR 95	1	99	99	
Megasphaera elsdenii		25.72	0.47	BGL 1	99	1	1	
Actinomyces meyeri		14.16	0.42		1	1	99	
01114003	**Low discrimination**							**01114003**
					COC	GRA		
Clostridium novyi biovar A		71.37	0.37	FRU 10	1	99		
Clostridium glycolicum		10.94	0.10	XYL 1	1	99		
Megasphaera elsdenii		7.48	0.12		99	1		
01114010	**Low discrimination**							**01114010**
					SPO			
Clostridium glycolicum		57.72	0.44	MLZ 1	99			
Clostridium novyi biovar A		21.56	0.50	SOR 95	99			
Actinomyces meyeri		7.10	0.33		1			
01114011	**Good identification**							**01114011**
Clostridium novyi biovar A		96.54	0.50	FRU 10				
Clostridium glycolicum		2.61	0.10	XYL 1				
Clostridium sordellii		0.85	-.03					
01114012	**Low discrimination**							**01114012**
					COC	GRA		
Clostridium novyi biovar A		74.00	0.37	FRU 10	1	99		
Clostridium glycolicum		11.34	0.10	XYL 1	1	99		
Megasphaera elsdenii		7.75	0.12		99	1		
01114013	**Good identification**							**01114013**
Clostridium novyi biovar A		100.0	0.37	FRU 10				
01114020	**Low discrimination**							**01114020**
					COC	SPO		
Actinomyces meyeri		51.44	0.53	SUC 93	1	1		
Clostridium glycolicum		30.98	0.44	NAG 12	1	99		
Gemella morbillorum		4.28	0.33		99	1		
01114021	**Low discrimination**							**01114021**
					SPO			
Clostridium novyi biovar A		75.59	0.39	FRU 10	99			
Actinomyces meyeri		13.58	0.19	XYL 1	1			
Clostridium glycolicum		8.18	0.10		99			
01114030	**Low discrimination**							**01114030**
					SPO			
Clostridium novyi biovar A		77.65	0.39	FRU 10	99			
Actinomyces meyeri		13.95	0.19	XYL 1	1			
Clostridium glycolicum		8.40	0.10		99			
01114031	**Good identification**							**01114031**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01114040	**Excellent identification**							**01114040**
					COC	GRA		
Clostridium glycolicum		99.74	1.00		1	99		
Megasphaera elsdenii		0.22	0.60		99	1		
Clostridium novyi biovar A		0.01	0.47		1	99		
01114041	**Very good identification**							**01114041**
					COC	GRA		
Clostridium glycolicum		98.43	0.66	URE 1	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		1.29	0.47		1	99	
Megasphaera elsdenii		0.22	0.26		99	1	
01114042	**Very good identification**						**01114042**
					COC	GRA	
Clostridium glycolicum		97.35	0.66	BGL 1	1	99	
Megasphaera elsdenii		2.34	0.44		99	1	
Clostridium novyi biovar A		0.23	0.34		1	99	
01114050	**Very good identification**						**01114050**
					COC	GRA	
Clostridium glycolicum		98.47	0.66	MLZ 1	1	99	
Clostridium novyi biovar A		1.29	0.47		1	99	
Megasphaera elsdenii		0.22	0.26		99	1	
01114051	**Low discrimination**						**01114051**
Clostridium novyi biovar A		56.48	0.47	FRU 10			
Clostridium glycolicum		43.53	0.32	XYL 1			
01114060	**Very good identification**						**01114060**
					COC	GRA	
Clostridium glycolicum		99.26	0.66	NAG 1	1	99	
Clostridium novyi biovar A		0.33	0.36		1	99	
Megasphaera elsdenii		0.22	0.26		99	1	
01114071	**Good identification**						**01114071**
Clostridium novyi biovar A		96.98	0.36	FRU 10			
Clostridium glycolicum		3.02	-.03	XYL 1			
01114100	**Very good identification**						**01114100**
					SPO		
Actinomyces meyeri		97.27	0.78	SUC 93	1		
Clostridium glycolicum		2.00	0.44		99		
Lactobacillus fermentum		0.25	0.44		1		
01114101	**Good identification**						**01114101**
					SPO		
Actinomyces meyeri		96.63	0.44	SUC 93	1		
Clostridium glycolicum		1.98	0.10	URE 1	99		
Clostridium novyi biovar A		0.74	0.15		99		
01114102	**Acceptable identification**						**01114102**
					SPO		
Actinomyces meyeri		91.19	0.52	SUC 93	1		
Lactobacillus fermentum		7.45	0.44	BGL 3	1		
Clostridium glycolicum		0.61	0.10		99		
01114110	**Good identification**						**01114110**
					SPO		
Actinomyces meyeri		97.26	0.44	SUC 93	1		
Clostridium glycolicum		2.00	0.10	MLZ 1	99		
Clostridium novyi biovar A		0.75	0.15		99		
01114120	**Very good identification**						**01114120**
					GRA		
Actinomyces meyeri		98.03	0.63	SUC 93	99		
Prevotella bivia		1.22	0.15	NAG 12	1		
Lactobacillus fermentum		0.61	0.36		99		
01114122	**Low discrimination**						**01114122**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		83.35	0.37	SUC 93				
Lactobacillus fermentum		16.66	0.36	NAG 12				
01114140	**Very good identification**							**01114140**
					COC	SPO	GRA	
Clostridium glycolicum		97.26	0.66	LAC 1	1	99	99	
Actinomyces meyeri		2.52	0.44		1	1	99	
Megasphaera elsdenii		0.21	0.26		99	1	1	
01114200	**Low discrimination**							**01114200**
					GRA	SPO		
Clostridium glycolicum		73.84	0.44	RHA 1	99	99		
Actinomyces meyeri		9.08	0.33	SOR 95	99	1		
Prevotella disiens		8.42	0.23		1	1		
01114240	**Very good identification**							**01114240**
					COC	GRA		
Clostridium glycolicum		99.77	0.66	RHA 1	1	99		
Megasphaera elsdenii		0.22	0.26		99	1		
Clostridium novyi biovar A		0.01	0.12		1	99		
01114300	**Good identification**							**01114300**
					SPO			
Actinomyces meyeri		97.05	0.44	SUC 93	1			
Clostridium glycolicum		1.99	0.10	RHA 1	99			
Actinomyces odontolyticus		0.96	0.13		1			
01114400	**Low discrimination**							**01114400**
					COC	SPO		
Actinomyces meyeri		68.24	0.57	SUC 93	1	1		
Clostridium glycolicum		22.42	0.44		1	99		
Gemella morbillorum		3.10	0.33		99	1		
01114420	**Low discrimination**							**01114420**
					COC			
Actinomyces meyeri		89.13	0.42	SUC 93	1			
Lactobacillus fermentum		5.40	0.32	NAG 12	1			
Gemella morbillorum		3.30	0.17		99			
01114440	**Very good identification**							**01114440**
					COC	SPO	GRA	
Clostridium glycolicum		99.61	0.66	ARA 1	1	99	99	
Megasphaera elsdenii		0.22	0.26		99	1	1	
Actinomyces meyeri		0.16	0.23		1	1	99	
01114500	**Very good identification**							**01114500**
					SPO			
Actinomyces meyeri		98.92	0.68	SUC 93	1			
Lactobacillus fermentum		1.00	0.44		1			
Clostridium glycolicum		0.08	0.10		99			
01114502	**Low discrimination**							**01114502**
Actinomyces meyeri		75.36	0.42	SUC 93				
Lactobacillus fermentum		24.64	0.44	BGL 3				
01114520	**Good identification**							**01114520**
Actinomyces meyeri		97.59	0.53	SUC 93				
Lactobacillus fermentum		2.41	0.36	NAG 12				
01115000	**Low discrimination**							**01115000**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		48.05	0.71	SUC 93	1	
Clostridium glycolicum		28.94	0.62		99	
Lactobacillus fermentum		19.66	0.75		1	
01115001	**Low discrimination**					**01115001**
					SPO	
Actinomyces meyeri		38.44	0.36	SUC 93	1	
Clostridium glycolicum		23.15	0.28	URE 1	99	
Lactobacillus fermentum		15.73	0.41		1	
01115002	**Good identification**					**01115002**
					SPO	
Lactobacillus fermentum		91.21	0.75	SUC 87	1	
Actinomyces meyeri		6.90	0.45		1	
Clostridium glycolicum		1.36	0.28		99	
01115003	**Low discrimination**					**01115003**
					SPO	
Lactobacillus fermentum		85.28	0.41	SUC 87	1	
Clostridium novyi biovar A		8.28	0.20	URE 1	99	
Actinomyces meyeri		6.45	0.11		1	
01115010	**Low discrimination**					**01115010**
					SPO	
Actinomyces meyeri		43.65	0.36	SUC 93	1	
Clostridium glycolicum		26.29	0.28	MLZ 1	99	
Lactobacillus fermentum		17.86	0.41		1	
01115012	**Low discrimination**					**01115012**
					SPO	
Lactobacillus fermentum		85.28	0.41	SUC 87	1	
Clostridium novyi biovar A		8.28	0.20	MLZ 1	99	
Actinomyces meyeri		6.45	0.11		1	
01115020	**Low discrimination**					**01115020**
					SPO	
Lactobacillus fermentum		48.02	0.67	SUC 87	1	
Actinomyces meyeri		48.02	0.56		1	
Clostridium glycolicum		2.14	0.28		99	
01115022	**Very good identification**					**01115022**
Lactobacillus fermentum		96.78	0.67	SUC 87		
Actinomyces meyeri		2.99	0.30			
Atopobium minutum		0.16	0.12			
01115040	**Very good identification**					**01115040**
					SPO	
Clostridium glycolicum		99.84	0.84	GAL 10	99	
Actinomyces meyeri		0.09	0.36		1	
Lactobacillus fermentum		0.04	0.41		1	
01115041	**Good identification**					**01115041**
Clostridium glycolicum		98.51	0.50	GAL 10		
Clostridium novyi biovar A		1.29	0.30	URE 1		
Clostridium sordellii		0.11	-.05			
01115042	**Good identification**					**01115042**
					SPO	
Clostridium glycolicum		95.88	0.50	GAL 10	99	
Lactobacillus fermentum		3.43	0.41	BGL 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		0.26	0.11		1	
01115050	**Good identification**					**01115050**
					SPO	
Clostridium glycolicum		98.62	0.50	GAL 10	99	
Clostridium novyi biovar A		1.29	0.30	MLZ 1	99	
Actinomyces meyeri		0.09	0.02		1	
01115060	**Good identification**					**01115060**
					SPO	
Clostridium glycolicum		97.36	0.50	GAL 10	99	
Lactobacillus fermentum		1.16	0.32	NAG 1	1	
Actinomyces meyeri		1.16	0.22		1	
01115100	**Low discrimination**					**01115100**
					SPO	
Actinomyces meyeri		85.49	0.81	SUC 93	1	
Lactobacillus fermentum		14.27	0.78		1	
Clostridium glycolicum		0.13	0.28		99	
01115101	**Low discrimination**					**01115101**
					SPO	
Actinomyces meyeri		85.05	0.47	SUC 93	1	
Lactobacillus fermentum		14.19	0.44	URE 1	1	
Clostridium chauvoei		0.76	0.09		99	
01115102	**Low discrimination**					**01115102**
					COC	
Lactobacillus fermentum		84.30	0.78	SUC 87	1	
Actinomyces meyeri		15.63	0.55		1	
Blautia hansenii		0.04	0.03		99	
01115103	**Low discrimination**					**01115103**
Lactobacillus fermentum		84.36	0.44	SUC 87		
Actinomyces meyeri		15.64	0.21	URE 1		
01115110	**Low discrimination**					**01115110**
Actinomyces meyeri		85.70	0.47	SUC 93		
Lactobacillus fermentum		14.30	0.44	MLZ 1		
01115112	**Low discrimination**					**01115112**
Lactobacillus fermentum		84.36	0.44	SUC 87		
Actinomyces meyeri		15.64	0.21	MLZ 1		
01115120	**Low discrimination**					**01115120**
					GRA	
Actinomyces meyeri		70.86	0.66	SUC 93	99	
Lactobacillus fermentum		28.90	0.70	NAG 12	99	
Prevotella bivia		0.10	0.03		1	
01115122	**Good identification**					**01115122**
Lactobacillus fermentum		92.89	0.70	SUC 87		
Actinomyces meyeri		7.04	0.40			
Clostridium perfringens		0.07	0.04			
01115123	**Acceptable identification**					**01115123**
Lactobacillus fermentum		92.95	0.36	SUC 87		
Actinomyces meyeri		7.05	0.06	URE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01115132	**Acceptable identification**					**01115132**
Lactobacillus fermentum		92.95	0.36	SUC 87		
Actinomyces meyeri		7.05	0.06	MLZ 1		
01115140	**Low discrimination**					**01115140**
					SPO	
Clostridium glycolicum		71.03	0.50	GAL 10	99	
Actinomyces meyeri		24.83	0.47	LAC 1	1	
Lactobacillus fermentum		4.14	0.44		1	
01115142	**Low discrimination**					**01115142**
					SPO	
Lactobacillus fermentum		73.61	0.44	SUC 87	1	
Actinomyces meyeri		13.64	0.21	SOR 1	1	
Clostridium glycolicum		12.75	0.15		99	
01115162	**Acceptable identification**					**01115162**
Lactobacillus fermentum		92.95	0.36	SUC 87		
Actinomyces meyeri		7.05	0.06	SOR 1		
01115200	**Low discrimination**					**01115200**
					SPO	
Actinomyces meyeri		47.14	0.36	SUC 93	1	
Clostridium glycolicum		28.39	0.28	RHA 1	99	
Lactobacillus fermentum		19.28	0.41		1	
01115202	**Low discrimination**					**01115202**
Lactobacillus fermentum		86.30	0.41	SUC 87		
Actinomyces odontolyticus		7.18	0.15	RHA 1		
Actinomyces meyeri		6.52	0.11			
01115240	**Good identification**					**01115240**
					SPO	
Clostridium glycolicum		99.91	0.50	GAL 10	99	
Actinomyces meyeri		0.09	0.02	RHA 1	1	
01115300	**Low discrimination**					**01115300**
Actinomyces meyeri		85.38	0.47	SUC 93		
Lactobacillus fermentum		14.25	0.44	RHA 1		
Actinomyces odontolyticus		0.37	0.10			
01115302	**Low discrimination**					**01115302**
Lactobacillus fermentum		83.23	0.44	SUC 87		
Actinomyces meyeri		15.43	0.21	RHA 1		
Actinomyces odontolyticus		1.34	0.06			
01115322	**Acceptable identification**					**01115322**
Lactobacillus fermentum		92.95	0.36	SUC 87		
Actinomyces meyeri		7.05	0.06	RHA 1		
01115400	**Low discrimination**					**01115400**
					SPO	
Lactobacillus fermentum		61.11	0.75	SUC 87	1	
Actinomyces meyeri		37.35	0.60		1	
Clostridium glycolicum		0.91	0.28		99	
01115401	**Low discrimination**					**01115401**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		62.07	0.41	SUC 87		
Actinomyces meyeri		37.93	0.26	URE 1		
01115402	**Very good identification**					**01115402**
Lactobacillus fermentum		98.15	0.75	SUC 87		
Actinomyces meyeri		1.86	0.34			
01115403	**Good identification**					**01115403**
Lactobacillus fermentum		100.0	0.41	SUC 87		
01115410	**Low discrimination**					**01115410**
Lactobacillus fermentum		62.07	0.41	SUC 87		
Actinomyces meyeri		37.93	0.26	MLZ 1		
01115412	**Good identification**					**01115412**
Lactobacillus fermentum		100.0	0.41	SUC 87		
01115420	**Low discrimination**					**01115420**
					COC	
Lactobacillus fermentum		79.79	0.67	SUC 87	1	
Actinomyces meyeri		19.95	0.45		1	
Gemella morbillorum		0.27	0.12		99	
01115422	**Very good identification**					**01115422**
Lactobacillus fermentum		99.23	0.67	SUC 87		
Actinomyces meyeri		0.77	0.20			
01115440	**Acceptable identification**					**01115440**
					SPO	
Clostridium glycolicum		94.55	0.50	GAL 10	99	
Lactobacillus fermentum		3.38	0.41	ARA 1	1	
Actinomyces meyeri		2.07	0.26		1	
01115442	**Low discrimination**					**01115442**
					SPO	
Lactobacillus fermentum		77.97	0.41	SUC 87	1	
Clostridium glycolicum		22.03	0.15	SOR 1	99	
01115500	**Low discrimination**					**01115500**
					COC	
Actinomyces meyeri		59.95	0.71	SUC 93	1	
Lactobacillus fermentum		40.01	0.78		1	
Blautia hansenii		0.03	0.06		99	
01115501	**Low discrimination**					**01115501**
Actinomyces meyeri		59.97	0.36	SUC 93		
Lactobacillus fermentum		40.03	0.44	URE 1		
01115502	**Very good identification**					**01115502**
Lactobacillus fermentum		95.54	0.78	SUC 87		
Actinomyces meyeri		4.43	0.45			
Bifidobacterium longum subsp. longum		0.03	0.12			
01115503	**Good identification**					**01115503**
Lactobacillus fermentum		95.57	0.44	SUC 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		4.43	0.11	URE 1		
01115510	**Low discrimination**					**01115510**
Actinomyces meyeri		58.67	0.36	SUC 93		
Lactobacillus fermentum		39.16	0.44	MLZ 1		
Bifidobacterium longum subsp. longum		2.17	0.16			
01115512	**Acceptable identification**					**01115512**
Lactobacillus fermentum		90.76	0.44	SUC 87		
Bifidobacterium longum subsp. longum		5.04	0.16	MLZ 1		
Actinomyces meyeri		4.21	0.11			
01115520	**Low discrimination**					**01115520**
Lactobacillus fermentum		61.83	0.70	SUC 87		
Actinomyces meyeri		37.89	0.56			
Bifidobacterium longum subsp. longum		0.24	0.22			
01115521	**Low discrimination**					**01115521**
Lactobacillus fermentum		62.00	0.36	SUC 87		
Actinomyces meyeri		38.00	0.22	URE 1		
01115522	**Very good identification**					**01115522**
Lactobacillus fermentum		97.76	0.70	SUC 87		
Actinomyces meyeri		1.85	0.30			
Bifidobacterium longum subsp. longum		0.39	0.22			
01115523	**Good identification**					**01115523**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01115530	**Low discrimination**					**01115530**
Lactobacillus fermentum		43.88	0.36	SUC 87		
Bifidobacterium longum subsp. longum		29.23	0.26	MLZ 1		
Actinomyces meyeri		26.89	0.22			
01115532	**Low discrimination**					**01115532**
Lactobacillus fermentum		60.02	0.36	SUC 87		
Bifidobacterium longum subsp. longum		39.98	0.26	MLZ 1		
01115540	**Low discrimination**					**01115540**
					SPO	
Actinomyces meyeri		56.08	0.36	SUC 93	1	
Lactobacillus fermentum		37.43	0.44	SOR 1	1	
Clostridium glycolicum		6.48	0.15		99	
01115542	**Good identification**					**01115542**
Lactobacillus fermentum		95.57	0.44	SUC 87		
Actinomyces meyeri		4.43	0.11	SOR 1		
01115560	**Low discrimination**					**01115560**
Lactobacillus fermentum		62.00	0.36	SUC 87		
Actinomyces meyeri		38.00	0.22	SOR 1		
01115562	**Good identification**					**01115562**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		100.0	0.36	SUC 87				
01115600	**Low discrimination**							**01115600**
Lactobacillus fermentum		62.07	0.41	SUC 87				
Actinomyces meyeri		37.93	0.26	RHA 1				
01115602	**Good identification**							**01115602**
Lactobacillus fermentum		100.0	0.41	SUC 87				
01115700	**Low discrimination**							**01115700**
Actinomyces meyeri		59.97	0.36	SUC 93				
Lactobacillus fermentum		40.03	0.44	RHA 1				
01115702	**Good identification**							**01115702**
Lactobacillus fermentum		95.57	0.44	SUC 87				
Actinomyces meyeri		4.43	0.11	RHA 1				
01115720	**Low discrimination**							**01115720**
					GRA			
Lactobacillus fermentum		46.43	0.36	SUC 87	99			
Actinomyces meyeri		28.46	0.22	RHA 1	99			
Bacteroides vulgatus		25.11	0.10		1			
01115722	**Good identification**							**01115722**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01116000	**Low discrimination**							**01116000**
					COC	SPO	GRA	
Megasphaera elsdenii		78.72	0.63	XYL 1	99	1	1	
Clostridium glycolicum		12.79	0.44		1	99	99	
Pseudoramibacter alactolyticus		2.71	0.27		1	1	99	
01116002	**Acceptable identification**							**01116002**
					COC	SPO	GRA	
Megasphaera elsdenii		94.50	0.47	XYL 1	99	1	1	
Pseudoramibacter alactolyticus		3.26	0.11	BGL 10	1	1	99	
Clostridium glycolicum		1.40	0.10		1	99	99	
01116040	**Low discrimination**							**01116040**
					COC	SPO	GRA	
Clostridium glycolicum		82.04	0.66	MAN 1	1	99	99	
Megasphaera elsdenii		17.71	0.60		99	1	1	
Pseudoramibacter alactolyticus		0.23	0.17		1	1	99	
01116042	**Low discrimination**							**01116042**
					COC	SPO	GRA	
Megasphaera elsdenii		69.74	0.44	XYL 1	99	1	1	
Clostridium glycolicum		29.36	0.32	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		0.90	0.00		1	1	99	
01116100	**Low discrimination**							**01116100**
					COC	SPO	GRA	
Actinomyces meyeri		86.47	0.44	SUC 93	1	1	99	
Megasphaera elsdenii		10.93	0.29	MAN 1	99	1	1	
Clostridium glycolicum		1.78	0.10		1	99	99	
01117000	**Low discrimination**							**01117000**
					SPO			
Actinomyces meyeri		41.42	0.36	SUC 93	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium glycolicum		24.95	0.28	MAN 1	99			
Lactobacillus fermentum		16.94	0.41		1			
01117002	**Low discrimination**							**01117002**
					COC	GRA		
Lactobacillus fermentum		85.68	0.41	SUC 87	1	99		
Megasphaera elsdenii		7.84	0.12	MAN 1	99	1		
Actinomyces meyeri		6.48	0.11		1	99		
01117040	**Good identification**							**01117040**
					COC	SPO	GRA	
Clostridium glycolicum		97.99	0.50	GAL 10	1	99	99	
Megasphaera elsdenii		1.92	0.26	MAN 1	99	1	1	
Actinomyces meyeri		0.09	0.02		1	1	99	
01117100	**Low discrimination**							**01117100**
Actinomyces meyeri		85.45	0.47	SUC 93				
Lactobacillus fermentum		14.26	0.44	MAN 1				
Propionibacterium propionicum		0.30	0.15					
01117102	**Low discrimination**							**01117102**
Lactobacillus fermentum		82.19	0.44	SUC 87				
Actinomyces meyeri		15.23	0.21	MAN 1				
Propionibacterium propionicum		2.57	0.18					
01117122	**Acceptable identification**							**01117122**
Lactobacillus fermentum		92.95	0.36	SUC 87				
Actinomyces meyeri		7.05	0.06	MAN 1				
01117400	**Low discrimination**							**01117400**
Lactobacillus fermentum		62.07	0.41	SUC 87				
Actinomyces meyeri		37.93	0.26	MAN 1				
01117402	**Good identification**							**01117402**
Lactobacillus fermentum		100.0	0.41	SUC 87				
01117500	**Low discrimination**							**01117500**
Actinomyces meyeri		59.97	0.36	SUC 93				
Lactobacillus fermentum		40.03	0.44	MAN 1				
01117502	**Good identification**							**01117502**
Lactobacillus fermentum		95.57	0.44	SUC 87				
Actinomyces meyeri		4.43	0.11	MAN 1				
01117520	**Low discrimination**							**01117520**
Lactobacillus fermentum		62.00	0.36	SUC 87				
Actinomyces meyeri		38.00	0.22	MAN 1				
01117522	**Good identification**							**01117522**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01120000	**Low discrimination**							**01120000**
					COC	SPO		
Clostridium botulinum biovar A		51.75	0.84		1	99		
Clostridium sporogenes		37.75	0.79		1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptostreptococcus anaerobius		5.92	0.63		99	1	
01120001	**Low discrimination**						**01120001**
Clostridium novyi biovar A		46.99	0.66	TRE	1		
Clostridium botulinum biovar A		25.30	0.49				
Clostridium sporogenes		18.46	0.45				
01120002	**Good identification**						**01120002**
Clostridium sporogenes		93.95	0.71				
Clostridium botulinum biovar A		3.90	0.49				
Clostridium novyi biovar A		1.28	0.53				
01120003	**Low discrimination**						**01120003**
Clostridium novyi biovar A		56.13	0.53	TRE	1		
Clostridium sporogenes		41.64	0.37				
Clostridium botulinum biovar A		1.73	0.15				
01120010	**Low discrimination**						**01120010**
Clostridium novyi biovar A		49.24	0.66	TRE	1		
Clostridium botulinum biovar A		26.52	0.49				
Clostridium sporogenes		19.34	0.45				
01120011	**Very good identification**						**01120011**
Clostridium novyi biovar A		98.91	0.66	TRE	1		
Clostridium botulinum biovar A		0.54	0.15				
Clostridium sporogenes		0.39	0.11				
01120012	**Low discrimination**						**01120012**
Clostridium novyi biovar A		56.30	0.53	TRE	1		
Clostridium sporogenes		41.77	0.37				
Clostridium botulinum biovar A		1.74	0.15				
01120013	**Good identification**						**01120013**
Clostridium novyi biovar A		99.26	0.53	TRE	1		
Clostridium sporogenes		0.74	0.03				
01120020	**Low discrimination**						**01120020**
Clostridium botulinum biovar A		40.11	0.49	NAG	1		
Clostridium sporogenes		29.26	0.45				
Clostridium novyi biovar A		18.63	0.56				
01120021	**Good identification**						**01120021**
Clostridium novyi biovar A		95.78	0.56	TRE	1		
Clostridium botulinum biovar A		2.08	0.15				
Clostridium sporogenes		1.52	0.11				
01120022	**Low discrimination**						**01120022**
Clostridium sporogenes		70.31	0.37	NAG	1		
Clostridium novyi biovar A		23.69	0.43				
Clostridium botulinum biovar A		2.92	0.15				
01120023	**Good identification**						**01120023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		97.09	0.43	TRE 1		
Clostridium sporogenes		2.91	0.03			
01120030	**Good identification**					**01120030**
Clostridium novyi biovar A		96.14	0.56	TRE 1		
Clostridium botulinum biovar A		2.09	0.15			
Clostridium sporogenes		1.53	0.11			
01120031	**Good identification**					**01120031**
Clostridium novyi biovar A		100.0	0.56	TRE 1		
01120032	**Good identification**					**01120032**
Clostridium novyi biovar A		97.09	0.43	TRE 1		
Clostridium sporogenes		2.91	0.03			
01120033	**Good identification**					**01120033**
Clostridium novyi biovar A		100.0	0.43	TRE 1		
01120040	**Good identification**					**01120040**
Clostridium botulinum biovar A		91.61	0.75			
Clostridium novyi biovar A		3.84	0.63			
Clostridium sporogenes		2.26	0.45			
01120041	**Low discrimination**					**01120041**
Clostridium novyi biovar A		79.48	0.63	TRE 1		
Clostridium botulinum biovar A		19.18	0.41			
Clostridium sordellii		0.48	0.08			
01120042	**Low discrimination**					**01120042**
Clostridium botulinum biovar A		38.37	0.41	BGL 1		
Clostridium sporogenes		31.23	0.37			
Clostridium novyi biovar A		28.07	0.50			
01120043	**Good identification**					**01120043**
Clostridium novyi biovar A		97.56	0.50	TRE 1		
Clostridium botulinum biovar A		1.35	0.06			
Clostridium sporogenes		1.10	0.03			
01120050	**Low discrimination**					**01120050**
Clostridium novyi biovar A		71.66	0.63	TRE 1		
Clostridium botulinum biovar A		17.29	0.41			
Clostridium botulinum biovar B		10.49	0.34			
01120051	**Very good identification**					**01120051**
Clostridium novyi biovar A		99.61	0.63	TRE 1		
Clostridium botulinum biovar A		0.24	0.06			
Clostridium botulinum biovar B		0.15	0.00			
01120052	**Good identification**					**01120052**
Clostridium novyi biovar A		96.77	0.50	TRE 1		
Clostridium botulinum biovar A		1.34	0.06			
Clostridium sporogenes		1.09	0.03			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
01120053	**Good identification**						**01120053**
Clostridium novyi biovar A		100.0	0.50	TRE 1			
01120060	**Low discrimination**						**01120060**
Clostridium novyi biovar A		38.68	0.53	TRE 1			
Clostridium botulinum biovar A		37.32	0.41				
Clostridium botulinum biovar B		22.66	0.34				
01120061	**Good identification**						**01120061**
Clostridium novyi biovar A		98.46	0.53	TRE 1			
Clostridium botulinum biovar A		0.96	0.06				
Clostridium botulinum biovar B		0.58	0.00				
01120062	**Low discrimination**						**01120062**
Clostridium novyi biovar A		88.20	0.40	TRE 1			
Clostridium botulinum biovar A		4.87	0.06				
Clostridium sporogenes		3.97	0.03				
01120063	**Good identification**						**01120063**
Clostridium novyi biovar A		100.0	0.40	TRE 1			
01120070	**Low discrimination**						**01120070**
Clostridium novyi biovar A		77.03	0.53	TRE 1			
Clostridium botulinum biovar B		22.22	0.29				
Clostridium botulinum biovar A		0.75	0.06				
01120071	**Good identification**						**01120071**
Clostridium novyi biovar A		100.0	0.53	TRE 1			
01120072	**Good identification**						**01120072**
Clostridium novyi biovar A		100.0	0.40	TRE 1			
01120073	**Good identification**						**01120073**
Clostridium novyi biovar A		100.0	0.40	TRE 1			
01120100	**Low discrimination**						**01120100**
					COC	SPO	
Clostridium botulinum biovar A		51.43	0.49	LAC 1	1	99	
Clostridium sporogenes		37.52	0.45		1	99	
Peptostreptococcus anaerobius		5.88	0.29		99	1	
01120102	**Acceptable identification**						**01120102**
Clostridium sporogenes		94.35	0.37	LAC 1			
Clostridium botulinum biovar A		3.92	0.15				
Clostridium novyi biovar A		1.29	0.19				
01120140	**Acceptable identification**						**01120140**
Clostridium botulinum biovar A		92.01	0.41	LAC 1			
Clostridium novyi biovar A		3.85	0.29				
Clostridium sporogenes		2.27	0.11				
01120200	**Low discrimination**						**01120200**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		51.78	0.49	RHA 1	1	99	
Clostridium sporogenes		37.77	0.45		1	99	
Peptostreptococcus anaerobius		5.92	0.29		99	1	
01120202	**Acceptable identification**						**01120202**
Clostridium sporogenes		94.35	0.37	RHA 1			
Clostridium botulinum biovar A		3.92	0.15				
Clostridium novyi biovar A		1.29	0.19				
01120240	**Acceptable identification**						**01120240**
Clostridium botulinum biovar A		92.01	0.41	RHA 1			
Clostridium novyi biovar A		3.85	0.29				
Clostridium sporogenes		2.27	0.11				
01120400	**Low discrimination**						**01120400**
					COC	SPO	
Clostridium botulinum biovar A		51.63	0.49	ARA 1	1	99	
Clostridium sporogenes		37.66	0.45		1	99	
Peptostreptococcus anaerobius		5.90	0.29		99	1	
01120402	**Acceptable identification**						**01120402**
Clostridium sporogenes		94.35	0.37	ARA 1			
Clostridium botulinum biovar A		3.92	0.15				
Clostridium novyi biovar A		1.29	0.19				
01120440	**Acceptable identification**						**01120440**
Clostridium botulinum biovar A		92.01	0.41	ARA 1			
Clostridium novyi biovar A		3.85	0.29				
Clostridium sporogenes		2.27	0.11				
01121000	**Low discrimination**						**01121000**
Clostridium botulinum biovar A		46.23	0.49	GAL 1			
Clostridium sporogenes		33.72	0.45				
Clostridium novyi biovar A		9.54	0.50				
01121001	**Low discrimination**						**01121001**
Clostridium novyi biovar A		80.42	0.50	TRE 1			
Clostridium sordellii		12.33	0.19	GAL 10			
Clostridium botulinum biovar A		3.94	0.15				
01121002	**Low discrimination**						**01121002**
Clostridium sporogenes		81.57	0.37	GAL 1			
Clostridium novyi biovar A		12.22	0.37				
Clostridium botulinum biovar A		3.39	0.15				
01121003	**Acceptable identification**						**01121003**
Clostridium novyi biovar A		93.68	0.37	TRE 1			
Clostridium sporogenes		6.32	0.03	GAL 10			
01121010	**Acceptable identification**						**01121010**
Clostridium novyi biovar A		91.72	0.50	TRE 1			
Clostridium botulinum biovar A		4.49	0.15	GAL 10			
Clostridium sporogenes		3.28	0.11				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01121011	**Good identification**					**01121011**
Clostridium novyi biovar A		100.0	0.50	TRE 1		
01121012	**Acceptable identification**					**01121012**
Clostridium novyi biovar A		93.68	0.37	TRE 1		
Clostridium sporogenes		6.32	0.03	GAL 10		
01121013	**Good identification**					**01121013**
Clostridium novyi biovar A		100.0	0.37	TRE 1		
01121020	**Low discrimination**					**01121020**
Clostridium novyi biovar A		67.09	0.39	TRE 1		
Clostridium botulinum biovar A		13.14	0.15	GAL 10		
Clostridium sporogenes		9.58	0.11			
01121021	**Good identification**					**01121021**
Clostridium novyi biovar A		100.0	0.39	TRE 1		
01121030	**Good identification**					**01121030**
Clostridium novyi biovar A		100.0	0.39	TRE 1		
01121031	**Good identification**					**01121031**
Clostridium novyi biovar A		100.0	0.39	TRE 1		
01121040	**Low discrimination**					**01121040**
Clostridium botulinum biovar A		65.43	0.41	GAL 1		
Clostridium novyi biovar A		30.13	0.47			
Clostridium sporogenes		1.61	0.11			
01121041	**Good identification**					**01121041**
Clostridium novyi biovar A		96.95	0.47	TRE 1		
Clostridium botulinum biovar A		2.13	0.06	GAL 10		
Clostridium sordellii		0.93	-.05			
01121050	**Good identification**					**01121050**
Clostridium novyi biovar A		96.60	0.47	TRE 1		
Clostridium botulinum biovar A		2.12	0.06	GAL 10		
Clostridium botulinum biovar B		1.29	0.00			
01121051	**Good identification**					**01121051**
Clostridium novyi biovar A		100.0	0.47	TRE 1		
01121060	**Low discrimination**					**01121060**
Clostridium novyi biovar A		87.64	0.36	TRE 1		
Clostridium botulinum biovar A		7.69	0.06	GAL 10		
Clostridium botulinum biovar B		4.67	0.00			
01121061	**Good identification**					**01121061**
Clostridium novyi biovar A		100.0	0.36	TRE 1		
01121070	**Good identification**					**01121070**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.36	TRE 1			
01121071	**Good identification**						**01121071**
Clostridium novyi biovar A		100.0	0.36	TRE 1			
01122000	**Low discrimination**						**01122000**
					COC	GRA	
Clostridium botulinum biovar A		47.41	0.49	MAN 1	1	99	
Clostridium sporogenes		34.58	0.45		1	99	
Megasphaera elsdenii		8.48	0.41		99	1	
01122002	**Low discrimination**						**01122002**
					COC	GRA	
Clostridium sporogenes		87.59	0.37	MAN 1	1	99	
Megasphaera elsdenii		7.16	0.25		99	1	
Clostridium botulinum biovar A		3.64	0.15		1	99	
01122040	**Low discrimination**						**01122040**
					COC	GRA	
Clostridium botulinum biovar A		67.47	0.41	MAN 1	1	99	
Megasphaera elsdenii		26.94	0.38		99	1	
Clostridium novyi biovar A		2.83	0.29		1	99	
01124000	**Low discrimination**						**01124000**
					COC	SPO	
Clostridium botulinum biovar A		49.96	0.49	XYL 1	1	99	
Clostridium sporogenes		36.44	0.45		1	99	
Peptostreptococcus anaerobius		5.71	0.29		99	1	
01124002	**Acceptable identification**						**01124002**
Clostridium sporogenes		94.35	0.37	XYL 1			
Clostridium botulinum biovar A		3.92	0.15				
Clostridium novyi biovar A		1.29	0.19				
01124040	**Low discrimination**						**01124040**
Clostridium glycolicum		78.23	0.50	FRU 90			
Clostridium botulinum biovar A		20.04	0.41	TRE 1			
Clostridium novyi biovar A		0.84	0.29				
01130000	**Low discrimination**						**01130000**
					GRA	SPO	
Clostridium sporogenes		42.84	0.66		99	99	
Clostridium botulinum biovar A		21.24	0.63		99	99	
Prevotella disiens		20.97	0.57		1	1	
01130001	**Low discrimination**						**01130001**
Clostridium novyi biovar A		30.32	0.50	FRU 10			
Clostridium sordellii		26.34	0.32	TRE 1			
Clostridium sporogenes		18.91	0.32				
01130002	**Low discrimination**						**01130002**
					COC	GRA	
Clostridium sporogenes		87.48	0.58		1	99	
Megasphaera elsdenii		7.78	0.47		99	1	
Clostridium botulinum biovar A		1.31	0.29		1	99	
01130010	**Low discrimination**						**01130010**
Clostridium novyi biovar A		38.48	0.50	FRU 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		24.00	0.32	TRE 1			
Clostridium botulinum biovar A		11.90	0.29				
01130011	**Good identification**						**01130011**
Clostridium novyi biovar A		98.52	0.50	FRU 10			
Clostridium sordellii		0.87	-.03	TRE 1			
Clostridium sporogenes		0.62	-.02				
01130013	**Good identification**						**01130013**
Clostridium novyi biovar A		100.0	0.37	FRU 10			
01130021	**Acceptable identification**						**01130021**
Clostridium novyi biovar A		94.31	0.39	FRU 10			
Clostridium sordellii		3.31	-.03	TRE 1			
Clostridium sporogenes		2.38	-.02				
01130030	**Low discrimination**						**01130030**
					SPO	CAT	
Clostridium novyi biovar A		87.91	0.39	FRU 10	99	1	
Clostridium botulinum biovar B		6.15	0.05	TRE 1	99	1	
Propionibacterium granulosum		3.72	0.14		1	90	
01130031	**Good identification**						**01130031**
Clostridium novyi biovar A		100.0	0.39	FRU 10			
01130040	**Low discrimination**						**01130040**
					COC	GRA	
Megasphaera elsdenii		34.52	0.60	TRE 1	99	1	
Clostridium botulinum biovar A		28.75	0.54		1	99	
Clostridium botulinum biovar B		22.43	0.50		1	99	
01130041	**Low discrimination**						**01130041**
					COC	GRA	
Clostridium novyi biovar A		65.90	0.47	FRU 10	1	99	
Megasphaera elsdenii		10.97	0.26	TRE 1	99	1	
Clostridium botulinum biovar A		9.13	0.20		1	99	
01130042	**Low discrimination**						**01130042**
					COC	GRA	
Megasphaera elsdenii		69.07	0.44	TRE 1	99	1	
Clostridium sporogenes		11.77	0.24	BGL 10	1	99	
Clostridium novyi biovar A		6.66	0.34		1	99	
01130050	**Low discrimination**						**01130050**
					COC	GRA	
Clostridium botulinum biovar B		79.47	0.44	SUC 90	1	99	
Clostridium novyi biovar A		15.07	0.47	MNS 99	1	99	
Megasphaera elsdenii		2.51	0.26		99	1	
01130051	**Acceptable identification**						**01130051**
Clostridium novyi biovar A		94.94	0.47	FRU 10			
Clostridium botulinum biovar B		5.06	0.10	TRE 1			
01130060	**Low discrimination**						**01130060**
					COC	GRA	
Clostridium botulinum biovar B		88.65	0.44	SUC 90	1	99	
Clostridium novyi biovar A		4.20	0.36	MNS 99	1	99	
Megasphaera elsdenii		2.80	0.26		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01130061	**Low discrimination**						**01130061**
Clostridium novyi biovar A		82.44	0.36	FRU 10			
Clostridium botulinum biovar B		17.56	0.10	TRE 1			
01130070	**Acceptable identification**						**01130070**
Clostridium botulinum biovar B		91.22	0.39	SUC 90			
Clostridium novyi biovar A		8.78	0.36	MNS 99			
01130071	**Acceptable identification**						**01130071**
Clostridium novyi biovar A		90.51	0.36	FRU 10			
Clostridium botulinum biovar B		9.50	0.05	TRE 1			
01130100	**Low discrimination**						**01130100**
					SPO		
Actinomyces meyeri		27.98	0.39	SUC 93	1		
Clostridium sporogenes		25.73	0.32	TRE 1	99		
Clostridium botulinum biovar A		12.76	0.29		99		
01131000	**Low discrimination**						**01131000**
					GRA	SPO	
Prevotella intermedia		23.23	0.39	TRE 1	1	1	
Clostridium sporogenes		16.63	0.32		99	99	
Lactobacillus fermentum		9.17	0.44		99	1	
01131002	**Low discrimination**						**01131002**
					SPO	CAT	
Lactobacillus fermentum		45.87	0.44	SUC 87	1	1	
Clostridium sporogenes		27.72	0.24	TRE 1	99	1	
Propionibacterium granulosum		10.35	0.29		1	90	
01131100	**Low discrimination**						**01131100**
					GRA		
Actinomyces meyeri		44.77	0.42	SUC 93	99		
Lactobacillus fermentum		24.69	0.48	TRE 1	99		
Fusobacterium mortiferum		19.47	0.45		1		
01131102	**Low discrimination**						**01131102**
					GRA		
Lactobacillus fermentum		78.65	0.48	SUC 87	99		
Fusobacterium mortiferum		6.89	0.28	TRE 1	1		
Clostridium perfringens		6.42	0.17		99		
01131120	**Low discrimination**						**01131120**
					SPO		
Clostridium perfringens		50.75	0.49	SUC 91	84		
Clostridium septicum		42.92	0.41	MNS 99	99		
Lactobacillus fermentum		3.02	0.40		1		
01131122	**Low discrimination**						**01131122**
					SPO		
Clostridium perfringens		85.99	0.43	SUC 91	84		
Lactobacillus fermentum		10.86	0.40	MNS 99	1		
Clostridium septicum		3.15	0.12		99		
01131160	**Good identification**						**01131160**
Clostridium perfringens		95.71	0.36	SUC 91			
Clostridium septicum		4.29	0.07	MNS 99			
01131400	**Low discrimination**						**01131400**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC			
Lactobacillus fermentum		78.04	0.44	SUC 87	1			
Actinomyces meyeri		14.43	0.21	TRE 1	1			
Gemella morbillorum		4.11	0.11		99			
01131402	**Good identification**							**01131402**
Lactobacillus fermentum		100.0	0.44	SUC 87				
01131420	**Acceptable identification**							**01131420**
Lactobacillus fermentum		92.97	0.36	SUC 87				
Actinomyces meyeri		7.03	0.06	TRE 1				
01131422	**Good identification**							**01131422**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01131500	**Low discrimination**							**01131500**
Lactobacillus fermentum		68.81	0.48	SUC 87				
Actinomyces meyeri		31.19	0.31	TRE 1				
01131502	**Good identification**							**01131502**
Lactobacillus fermentum		98.62	0.48	SUC 87				
Actinomyces meyeri		1.38	0.06	TRE 1				
01131520	**Low discrimination**							**01131520**
Lactobacillus fermentum		66.71	0.40	SUC 87				
Actinomyces meyeri		12.37	0.17	TRE 1				
Clostridium perfringens		11.34	0.15					
01131522	**Acceptable identification**							**01131522**
Lactobacillus fermentum		92.59	0.40	SUC 87				
Clostridium perfringens		7.41	0.09	TRE 1				
01132000	**Low discrimination**							**01132000**
					COC	SPO	GRA	
Megasphaera elsdenii		86.37	0.63	TRE 1	99	1	1	
Clostridium sporogenes		3.27	0.32		1	99	99	
Pseudoramibacter alactolyticus		2.98	0.27		1	1	99	
01132002	**Low discrimination**							**01132002**
					COC	SPO	GRA	
Megasphaera elsdenii		84.41	0.47	TRE 1	99	1	1	
Clostridium sporogenes		9.59	0.24	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.91	0.11		1	1	99	
01132040	**Very good identification**							**01132040**
					COC	SPO	GRA	
Megasphaera elsdenii		96.82	0.60	TRE 1	99	1	1	
Pseudoramibacter alactolyticus		1.25	0.17		1	1	99	
Clostridium botulinum biovar A		0.81	0.20		1	99	99	
01132042	**Good identification**							**01132042**
					COC	GRA		
Megasphaera elsdenii		98.06	0.44	TRE 1	99	1		
Pseudoramibacter alactolyticus		1.27	0.00	BGL 10	1	99		
Propionibacterium propionicum		0.68	0.17		1	99		
01132100	**Low discrimination**							**01132100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Propionibacterium propionicum		75.05	0.48	NIT 86	1	99		
Megasphaera elsdenii		22.41	0.29	SUC 99	99	1		
Actinomyces meyeri		0.92	0.05		1	99		
01132102	**Good identification**							**01132102**
					COC	GRA		
Propionibacterium propionicum		97.84	0.51	NIT 86	1	99		
Megasphaera elsdenii		2.16	0.12	SUC 99	99	1		
01132140	**Low discrimination**							**01132140**
					COC	GRA		
Propionibacterium propionicum		83.40	0.48	NIT 86	1	99		
Megasphaera elsdenii		16.60	0.26	SUC 99	99	1		
01132142	**Good identification**							**01132142**
					COC	GRA		
Propionibacterium propionicum		98.55	0.51	NIT 86	1	99		
Megasphaera elsdenii		1.45	0.09	SUC 99	99	1		
01133000	**Low discrimination**							**01133000**
					COC	GRA	CAT	
Propionibacterium granulosum		54.85	0.37	SUC 93	1	99	90	
Megasphaera elsdenii		38.95	0.29	MNS 99	99	1	1	
Prevotella intermedia		2.06	0.05		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01133100	**Good identification**						**01133100**
Propionibacterium propionicum		97.22	0.48	NIT 86			
Actinomyces meyeri		1.79	0.08	SUC 99			
Lactobacillus fermentum		0.99	0.14				
01133102	**Good identification**						**01133102**
Propionibacterium propionicum		99.33	0.51	NIT 86			
Lactobacillus fermentum		0.67	0.14	SUC 99			
01133140	**Good identification**						**01133140**
Propionibacterium propionicum		100.0	0.48	NIT 86			
01133142	**Good identification**						**01133142**
Propionibacterium propionicum		100.0	0.51	NIT 86			
01134000	**Low discrimination**						**01134000**
					SPO		
Clostridium glycolicum		59.55	0.44	TRE 1	99		
Clostridium sporogenes		13.88	0.32	SOR 95	99		
Actinomyces meyeri		7.32	0.33		1		
01134040	**Very good identification**						**01134040**
					COC	GRA	
Clostridium glycolicum		99.44	0.66	TRE 1	1	99	
Megasphaera elsdenii		0.22	0.26		99	1	
Clostridium botulinum biovar A		0.18	0.20		1	99	
01134100	**Good identification**						**01134100**
					GRA	SPO	
Actinomyces meyeri		95.95	0.44	SUC 93	99	1	
Clostridium glycolicum		1.97	0.10	TRE 1	99	99	
Fusobacterium mortiferum		1.62	0.23		1	1	
01135000	**Low discrimination**						**01135000**
					SPO		
Actinomyces meyeri		47.45	0.36	SUC 93	1		
Clostridium glycolicum		28.58	0.28	TRE 1	99		
Lactobacillus fermentum		19.41	0.41		1		
01135002	**Acceptable identification**						**01135002**
Lactobacillus fermentum		92.97	0.41	SUC 87			
Actinomyces meyeri		7.03	0.11	TRE 1			
01135040	**Good identification**						**01135040**
					SPO		
Clostridium glycolicum		99.91	0.50	TRE 1	99		
Actinomyces meyeri		0.09	0.02	GAL 10	1		
01135100	**Low discrimination**						**01135100**
					COC		
Actinomyces meyeri		78.34	0.47	SUC 93	1		
Lactobacillus fermentum		13.07	0.44	TRE 1	1		
Atopobium parvulum		7.27	0.27		99		
01135102	**Low discrimination**						**01135102**
					COC		
Lactobacillus fermentum		77.91	0.44	SUC 87	1		
Actinomyces meyeri		14.44	0.21	TRE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		7.65	0.14		99			
01135122	**Low discrimination**							**01135122**
Lactobacillus fermentum		75.78	0.36	SUC 87				
Clostridium perfringens		10.32	0.09	TRE 1				
Actinomyces meyeri		5.75	0.06					
01135400	**Low discrimination**							**01135400**
Lactobacillus fermentum		62.07	0.41	SUC 87				
Actinomyces meyeri		37.93	0.26	TRE 1				
01135402	**Good identification**							**01135402**
Lactobacillus fermentum		100.0	0.41	SUC 87				
01135500	**Low discrimination**							**01135500**
Actinomyces meyeri		59.97	0.36	SUC 93				
Lactobacillus fermentum		40.03	0.44	TRE 1				
01135502	**Good identification**							**01135502**
Lactobacillus fermentum		95.57	0.44	SUC 87				
Actinomyces meyeri		4.43	0.11	TRE 1				
01135520	**Low discrimination**							**01135520**
Lactobacillus fermentum		58.46	0.36	SUC 87				
Actinomyces meyeri		35.83	0.22	TRE 1				
Bifidobacterium longum subsp. longum		5.72	0.12					
01135522	**Acceptable identification**							**01135522**
Lactobacillus fermentum		91.09	0.36	SUC 87				
Bifidobacterium longum subsp. longum		8.91	0.12	TRE 1				
01140000	**Low discrimination**							**01140000**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		43.05	0.63	CEL 1	99	99	1	
Prevotella disiens		23.72	0.57		1	1	1	
Clostridium botulinum biovar A		15.21	0.60		1	99	99	
01140001	**Low discrimination**							**01140001**
					COC	SPO		
Clostridium novyi biovar A		81.27	0.66	CEL 1	1	99		
Clostridium sordellii		7.84	0.32		1	99		
Peptostreptococcus anaerobius		5.00	0.29		99	1		
01140002	**Low discrimination**							**01140002**
					COC	SPO		
Clostridium sporogenes		55.24	0.45	CEL 1	1	99		
Clostridium novyi biovar A		24.82	0.53		1	99		
Peptostreptococcus anaerobius		8.66	0.29		99	1		
01140003	**Good identification**							**01140003**
Clostridium novyi biovar A		96.94	0.53	CEL 1				
Clostridium sporogenes		2.18	0.11					
Clostridium sordellii		0.54	-.03					
01140010	**Low discrimination**							**01140010**
					COC	GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		88.33	0.66	CEL 1	1	99	99	
Peptostreptococcus anaerobius		5.44	0.29		99	99	1	
Prevotella disiens		3.00	0.23		1	1	1	
01140011	**Very good identification**							**01140011**
					COC	SPO		
Clostridium novyi biovar A		99.84	0.66	CEL 1	1	99		
Clostridium sordellii		0.10	-.03		1	99		
Peptostreptococcus anaerobius		0.06	-.06		99	1		
01140012	**Good identification**							**01140012**
					COC	SPO		
Clostridium novyi biovar A		97.47	0.53	CEL 1	1	99		
Clostridium sporogenes		2.19	0.11		1	99		
Peptostreptococcus anaerobius		0.34	-.06		99	1		
01140013	**Good identification**							**01140013**
Clostridium novyi biovar A		100.0	0.53	CEL 1				
01140020	**Low discrimination**							**01140020**
					COC	GRA	SPO	
Clostridium novyi biovar A		60.69	0.56	CEL 1	1	99	99	
Peptostreptococcus anaerobius		14.94	0.29		99	99	1	
Prevotella disiens		8.24	0.23		1	1	1	
01140021	**Good identification**							**01140021**
					COC	SPO		
Clostridium novyi biovar A		99.37	0.56	CEL 1	1	99		
Clostridium sordellii		0.39	-.03		1	99		
Peptostreptococcus anaerobius		0.25	-.06		99	1		
01140022	**Low discrimination**							**01140022**
					SPO			
Clostridium novyi biovar A		82.00	0.43	CEL 1	99			
Clostridium sporogenes		7.37	0.11		99			
Atopobium minutum		7.26	0.21		1			
01140023	**Good identification**							**01140023**
Clostridium novyi biovar A		100.0	0.43	CEL 1				
01140030	**Good identification**							**01140030**
					COC	SPO		
Clostridium novyi biovar A		99.75	0.56	CEL 1	1	99		
Peptostreptococcus anaerobius		0.25	-.06		99	1		
01140031	**Good identification**							**01140031**
Clostridium novyi biovar A		100.0	0.56	CEL 1				
01140032	**Good identification**							**01140032**
Clostridium novyi biovar A		100.0	0.43	CEL 1				
01140033	**Good identification**							**01140033**
Clostridium novyi biovar A		100.0	0.43	CEL 1				
01140040	**Low discrimination**							**01140040**
					COC	GRA		
Clostridium novyi biovar A		43.75	0.63	CEL 1	1	99		
Clostridium botulinum biovar A		42.22	0.51		1	99		
Megasphaera elsdenii		4.22	0.38		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01140041	**Very good identification**						**01140041**
Clostridium novyi biovar A		98.26	0.63	CEL 1			
Clostridium botulinum biovar A		0.96	0.17				
Clostridium sordellii		0.59	0.08				
01140042	**Low discrimination**						**01140042**
					COC	GRA	
Clostridium novyi biovar A		86.59	0.50	CEL 1	1	99	
Megasphaera elsdenii		5.25	0.22		99	1	
Clostridium botulinum biovar A		4.78	0.17		1	99	
01140043	**Good identification**						**01140043**
Clostridium novyi biovar A		100.0	0.50	CEL 1			
01140050	**Very good identification**						**01140050**
					COC	GRA	
Clostridium novyi biovar A		98.85	0.63	CEL 1	1	99	
Clostridium botulinum biovar A		0.96	0.17		1	99	
Megasphaera elsdenii		0.10	0.04		99	1	
01140051	**Very good identification**						**01140051**
Clostridium novyi biovar A		100.0	0.63	CEL 1			
01140052	**Good identification**						**01140052**
Clostridium novyi biovar A		100.0	0.50	CEL 1			
01140053	**Good identification**						**01140053**
Clostridium novyi biovar A		100.0	0.50	CEL 1			
01140060	**Good identification**						**01140060**
					COC	GRA	
Clostridium novyi biovar A		95.25	0.53	CEL 1	1	99	
Clostridium botulinum biovar A		3.71	0.17		1	99	
Megasphaera elsdenii		0.37	0.04		99	1	
01140061	**Good identification**						**01140061**
Clostridium novyi biovar A		100.0	0.53	CEL 1			
01140062	**Good identification**						**01140062**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
01140063	**Good identification**						**01140063**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
01140070	**Good identification**						**01140070**
Clostridium novyi biovar A		100.0	0.53	CEL 1			
01140071	**Good identification**						**01140071**
Clostridium novyi biovar A		100.0	0.53	CEL 1			
01140072	**Good identification**						**01140072**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
01140073	**Good identification**						**01140073**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.40	CEL 1				
01141000	**Low discrimination**							**01141000**
					COC	GRA	SPO	
Clostridium novyi biovar A		41.57	0.50	CEL 1	1	99	99	
Peptostreptococcus anaerobius		23.03	0.29	GAL 10	99	99	1	
Prevotella disiens		12.69	0.23		1	1	1	
01141001	**Low discrimination**							**01141001**
					COC	SPO		
Clostridium novyi biovar A		86.29	0.50	CEL 1	1	99		
Clostridium sordellii		13.23	0.19	GAL 10	1	99		
Peptostreptococcus anaerobius		0.48	-.06		99	1		
01141002	**Low discrimination**							**01141002**
					SPO			
Clostridium novyi biovar A		69.77	0.37	CEL 1	99			
Clostridium sporogenes		14.12	0.11	GAL 10	99			
Atopobium minutum		13.90	0.21		1			
01141003	**Good identification**							**01141003**
Clostridium novyi biovar A		100.0	0.37	CEL 1				
01141010	**Good identification**							**01141010**
					COC	SPO		
Clostridium novyi biovar A		99.44	0.50	CEL 1	1	99		
Peptostreptococcus anaerobius		0.56	-.06	GAL 10	99	1		
01141011	**Good identification**							**01141011**
Clostridium novyi biovar A		100.0	0.50	CEL 1				
01141012	**Good identification**							**01141012**
Clostridium novyi biovar A		100.0	0.37	CEL 1				
01141013	**Good identification**							**01141013**
Clostridium novyi biovar A		100.0	0.37	CEL 1				
01141020	**Low discrimination**							**01141020**
					COC	SPO		
Clostridium novyi biovar A		86.81	0.39	CEL 1	1	99		
Atopobium minutum		6.10	0.15	GAL 10	1	1		
Gemella morbillorum		5.14	0.09		99	1		
01141021	**Good identification**							**01141021**
Clostridium novyi biovar A		100.0	0.39	CEL 1				
01141030	**Good identification**							**01141030**
Clostridium novyi biovar A		100.0	0.39	CEL 1				
01141031	**Good identification**							**01141031**
Clostridium novyi biovar A		100.0	0.39	CEL 1				
01141040	**Low discrimination**							**01141040**
Clostridium novyi biovar A		86.62	0.47	CEL 1				
Clostridium botulinum biovar A		7.60	0.17	GAL 10				
Clostridium glycolicum		4.29	0.11					

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
01141041	**Good identification**							**01141041**
Clostridium novyi biovar A		99.05	0.47	CEL 1				
Clostridium sordellii		0.95	-.05	GAL 10				
01141050	**Good identification**							**01141050**
Clostridium novyi biovar A		100.0	0.47	CEL 1				
01141051	**Good identification**							**01141051**
Clostridium novyi biovar A		100.0	0.47	CEL 1				
01141060	**Good identification**							**01141060**
Clostridium novyi biovar A		100.0	0.36	CEL 1				
01141061	**Good identification**							**01141061**
Clostridium novyi biovar A		100.0	0.36	CEL 1				
01141070	**Good identification**							**01141070**
Clostridium novyi biovar A		100.0	0.36	CEL 1				
01141071	**Good identification**							**01141071**
Clostridium novyi biovar A		100.0	0.36	CEL 1				
01142000	**Low discrimination**							**01142000**
					COC	GRA		
Megasphaera elsdenii		40.43	0.41	FRU 95	99	1		
Peptostreptococcus anaerobius		25.83	0.29	CEL 1	99	99		
Prevotella disiens		14.24	0.23		1	1		
01142040	**Low discrimination**							**01142040**
					COC	GRA		
Megasphaera elsdenii		82.26	0.38	FRU 95	99	1		
Clostridium novyi biovar A		8.62	0.29	CEL 1	1	99		
Clostridium botulinum biovar A		8.32	0.17		1	99		
01144040	**Good identification**							**01144040**
Clostridium glycolicum		97.74	0.50	FRU 90				
Clostridium novyi biovar A		1.05	0.29	CEL 1				
Clostridium botulinum biovar A		1.01	0.17					
01150000	**Low discrimination**							**01150000**
					COC	GRA	SPO	
Prevotella disiens		55.45	0.57	CEL 1	1	1	1	
Megasphaera elsdenii		30.22	0.63		99	1	1	
Clostridium sporogenes		3.43	0.40		1	99	99	
01150001	**Low discrimination**							**01150001**
					GRA	SPO		
Clostridium novyi biovar A		41.72	0.50	FRU 10	99	99		
Clostridium sordellii		36.24	0.32	CEL 1	99	99		
Prevotella disiens		12.74	0.23		1	1		
01150002	**Low discrimination**							**01150002**
					COC	GRA	SPO	
Megasphaera elsdenii		57.23	0.47	CEL 1	99	1	1	
Clostridium sporogenes		19.50	0.32	BGL 10	1	99	99	
Prevotella disiens		9.55	0.23		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01150003	**Low discrimination**							**01150003**
					COC	GRA		
Clostridium novyi biovar A		84.02	0.37	FRU 10	1	99		
Megasphaera elsdenii		8.80	0.12	CEL 1	99	1		
Clostridium sordellii		4.18	-.03		1	99		
01150010	**Low discrimination**							**01150010**
					COC	GRA	SPO	
Clostridium novyi biovar A		62.82	0.50	FRU 10	1	99	99	
Prevotella disiens		19.18	0.23	CEL 1	1	1	1	
Megasphaera elsdenii		10.45	0.29		99	1	1	
01150011	**Good identification**							**01150011**
Clostridium novyi biovar A		99.13	0.50	FRU 10				
Clostridium sordellii		0.87	-.03	CEL 1				
01150012	**Low discrimination**							**01150012**
					COC	GRA		
Clostridium novyi biovar A		87.68	0.37	FRU 10	1	99		
Megasphaera elsdenii		9.19	0.12	CEL 1	99	1		
Clostridium sporogenes		3.13	-.02		1	99		
01150013	**Good identification**							**01150013**
Clostridium novyi biovar A		100.0	0.37	FRU 10				
01150021	**Good identification**							**01150021**
Clostridium novyi biovar A		96.61	0.39	FRU 10				
Clostridium sordellii		3.39	-.03	CEL 1				
01150030	**Good identification**							**01150030**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01150031	**Good identification**							**01150031**
Clostridium novyi biovar A		100.0	0.39	FRU 10				
01150040	**Low discrimination**							**01150040**
					COC	GRA		
Megasphaera elsdenii		71.68	0.60	CEL 1	99	1		
Clostridium glycolicum		17.47	0.44		1	99		
Clostridium novyi biovar A		4.35	0.47		1	99		
01150041	**Low discrimination**							**01150041**
					COC	GRA		
Clostridium novyi biovar A		79.29	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.19	0.26	CEL 1	99	1		
Clostridium sordellii		4.31	0.08		1	99		
01150042	**Low discrimination**							**01150042**
					COC	SPO	GRA	
Megasphaera elsdenii		87.13	0.44	CEL 1	99	1	1	
Clostridium novyi biovar A		8.40	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.09	0.05		1	1	99	
01150050	**Low discrimination**							**01150050**
					COC	GRA		
Clostridium novyi biovar A		81.65	0.47	FRU 10	1	99		
Megasphaera elsdenii		13.59	0.26	CEL 1	99	1		
Clostridium glycolicum		3.31	0.10		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01150051	**Good identification**							**01150051**
Clostridium novyi biovar A		100.0	0.47	FRU 10				
01150060	**Low discrimination**							**01150060**
					COC	SPO	GRA	
Clostridium novyi biovar A		45.92	0.36	FRU 10	1	99	99	
Megasphaera elsdenii		30.56	0.26	CEL 1	99	1	1	
Pseudoramibacter alactolyticus		12.81	0.09		1	1	99	
01150061	**Good identification**							**01150061**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01150070	**Good identification**							**01150070**
Clostridium novyi biovar A		96.62	0.36	FRU 10				
Clostridium botulinum biovar B		3.38	-.03	CEL 1				
01150071	**Good identification**							**01150071**
Clostridium novyi biovar A		100.0	0.36	FRU 10				
01150100	**Low discrimination**							**01150100**
					GRA			
Actinomyces meyeri		43.72	0.39	SUC 93	99			
Fusobacterium mortiferum		21.22	0.43	CEL 1	1			
Prevotella disiens		19.68	0.23		1			
01151000	**Low discrimination**							**01151000**
					GRA			
Prevotella intermedia		33.58	0.39	CEL 1	1			
Lactobacillus fermentum		13.26	0.44		99			
Prevotella disiens		11.76	0.23		1			
01151002	**Low discrimination**							**01151002**
Lactobacillus fermentum		69.03	0.44	SUC 87				
Lactobacillus jensenii		7.49	0.16	CEL 1				
Atopobium minutum		6.51	0.20					
01151020	**Low discrimination**							**01151020**
					SPO			
Lactobacillus fermentum		42.11	0.36	SUC 87	1			
Actinomyces meyeri		12.74	0.17	CEL 1	1			
Clostridium septicum		10.43	0.07		99			
01151022	**Low discrimination**							**01151022**
Lactobacillus fermentum		79.49	0.36	SUC 87				
Atopobium minutum		7.50	0.12	CEL 1				
Lactobacillus jensenii		6.47	0.06					
01151100	**Low discrimination**							**01151100**
					GRA			
Actinomyces meyeri		48.05	0.42	SUC 93	99			
Lactobacillus fermentum		26.50	0.48	CEL 1	99			
Fusobacterium mortiferum		23.32	0.46		1			
01151102	**Low discrimination**							**01151102**
					GRA			
Lactobacillus fermentum		86.67	0.48	SUC 87	99			
Fusobacterium mortiferum		8.48	0.29	CEL 1	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		4.86	0.16		99			
01151120	**Low discrimination**							**01151120**
					SPO			
Clostridium septicum		83.61	0.41	MNS 99	99			
Lactobacillus fermentum		5.57	0.40		1			
Actinomyces meyeri		4.13	0.27		1			
01151122	**Low discrimination**							**01151122**
					GRA	SPO		
Lactobacillus fermentum		55.62	0.40	SUC 87	99	1		
Clostridium septicum		17.05	0.12	CEL 1	99	99		
Prevotella oralis		11.20	0.12		1	1		
01151400	**Low discrimination**							**01151400**
					COC			
Lactobacillus fermentum		79.17	0.44	SUC 87	1			
Actinomyces meyeri		14.64	0.21	CEL 1	1			
Gemella morbillorum		4.16	0.11		99			
01151402	**Good identification**							**01151402**
Lactobacillus fermentum		100.0	0.44	SUC 87				
01151420	**Acceptable identification**							**01151420**
Lactobacillus fermentum		92.97	0.36	SUC 87				
Actinomyces meyeri		7.03	0.06	CEL 1				
01151422	**Good identification**							**01151422**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01151500	**Low discrimination**							**01151500**
					GRA			
Lactobacillus fermentum		68.39	0.48	SUC 87	99			
Actinomyces meyeri		31.00	0.31	CEL 1	99			
Fusobacterium mortiferum		0.61	0.11		1			
01151502	**Good identification**							**01151502**
Lactobacillus fermentum		98.62	0.48	SUC 87				
Actinomyces meyeri		1.38	0.06	CEL 1				
01151520	**Low discrimination**							**01151520**
					SPO			
Lactobacillus fermentum		74.79	0.40	SUC 87	1			
Actinomyces meyeri		13.87	0.17	CEL 1	1			
Clostridium septicum		11.35	0.07		99			
01151522	**Good identification**							**01151522**
Lactobacillus fermentum		100.0	0.40	SUC 87				
01152000	**Good identification**							**01152000**
					COC	GRA		
Megasphaera elsdenii		92.13	0.63	CEL 1	99	1		
Pseudoramibacter alactolyticus		3.17	0.27		1	99		
Eubacterium limosum		2.02	0.26		1	99		
01152002	**Good identification**							**01152002**
					COC	SPO	GRA	
Megasphaera elsdenii		96.35	0.47	CEL 1	99	1	1	
Pseudoramibacter alactolyticus		3.32	0.11	BGL 10	1	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sporogenes		0.33	-.02		1	99	99	
01152040	**Very good identification**							**01152040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.37	0.60	CEL 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.17		1	1	99	
Clostridium glycolicum		0.24	0.10		1	99	99	
01152042	**Good identification**							**01152042**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	CEL 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
01154000	**Low discrimination**							**01154000**
					GRA	SPO		
Clostridium glycolicum		74.47	0.44	CEL 1	99	99		
Actinomyces meyeri		9.16	0.33	SOR 95	99	1		
Prevotella disiens		8.49	0.23		1	1		
01154040	**Very good identification**							**01154040**
					COC	GRA		
Clostridium glycolicum		99.77	0.66	CEL 1	1	99		
Megasphaera elsdenii		0.22	0.26		99	1		
Clostridium novyi biovar A		0.01	0.12		1	99		
01154100	**Good identification**							**01154100**
					GRA	SPO		
Actinomyces meyeri		96.21	0.44	SUC 93	99	1		
Clostridium glycolicum		1.98	0.10	CEL 1	99	99		
Fusobacterium mortiferum		1.81	0.23		1	1		
01155000	**Low discrimination**							**01155000**
					SPO			
Actinomyces meyeri		47.66	0.36	SUC 93	1			
Clostridium glycolicum		28.71	0.28	CEL 1	99			
Lactobacillus fermentum		19.50	0.41		1			
01155002	**Acceptable identification**							**01155002**
Lactobacillus fermentum		92.97	0.41	SUC 87				
Actinomyces meyeri		7.03	0.11	CEL 1				
01155040	**Good identification**							**01155040**
					SPO			
Clostridium glycolicum		99.91	0.50	CEL 1	99			
Actinomyces meyeri		0.09	0.02	GAL 10	1			
01155100	**Low discrimination**							**01155100**
					GRA			
Actinomyces meyeri		84.34	0.47	SUC 93	99			
Lactobacillus fermentum		14.07	0.44	CEL 1	99			
Fusobacterium mortiferum		1.59	0.26		1			
01155102	**Low discrimination**							**01155102**
Lactobacillus fermentum		84.36	0.44	SUC 87				
Actinomyces meyeri		15.64	0.21	CEL 1				
01155122	**Low discrimination**							**01155122**
					GRA			
Lactobacillus fermentum		89.43	0.36	SUC 87	99			
Actinomyces meyeri		6.78	0.06	CEL 1	99			
Prevotella oralis		3.79	-.03		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01155400	**Low discrimination**							**01155400**
Lactobacillus fermentum		62.07	0.41	SUC 87				
Actinomyces meyeri		37.93	0.26	CEL 1				
01155402	**Good identification**							**01155402**
Lactobacillus fermentum		100.0	0.41	SUC 87				
01155500	**Low discrimination**							**01155500**
Actinomyces meyeri		59.97	0.36	SUC 93				
Lactobacillus fermentum		40.03	0.44	CEL 1				
01155502	**Acceptable identification**							**01155502**
Lactobacillus fermentum		93.95	0.44	SUC 87				
Actinomyces meyeri		4.35	0.11	CEL 1				
Bifidobacterium adolescentis		1.70	0.09					
01155520	**Low discrimination**							**01155520**
Lactobacillus fermentum		62.00	0.36	SUC 87				
Actinomyces meyeri		38.00	0.22	CEL 1				
01155522	**Good identification**							**01155522**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01160000	**Low discrimination**							**01160000**
					COC	SPO		
Clostridium botulinum biovar A		51.78	0.49	CEL 1	1	99		
Clostridium sporogenes		37.77	0.45		1	99		
Peptostreptococcus anaerobius		5.92	0.29		99	1		
01160002	**Acceptable identification**							**01160002**
Clostridium sporogenes		94.35	0.37	CEL 1				
Clostridium botulinum biovar A		3.92	0.15					
Clostridium novyi biovar A		1.29	0.19					
01160040	**Acceptable identification**							**01160040**
Clostridium botulinum biovar A		92.01	0.41	CEL 1				
Clostridium novyi biovar A		3.85	0.29					
Clostridium sporogenes		2.27	0.11					
01171120	**Good identification**							**01171120**
					COC	SPO		
Clostridium septicum		98.00	0.49	MNS 99	1	99		
Clostridium perfringens		1.53	0.25		1	84		
Atopobium parvulum		0.47	0.16		99	1		
01200000	**Low discrimination**							**01200000**
					COC	SPO		
Peptostreptococcus anaerobius		35.11	0.66	SAL 1	99	1		
Clostridium novyi biovar A		33.07	0.82		1	99		
Clostridium botulinum biovar A		10.11	0.61		1	99		
01200001	**Very good identification**							**01200001**
					COC	GRA	SPO	
Clostridium novyi biovar A		95.34	0.82	SAL 8	1	99	99	
Campylobacter ureolyticus		2.00	0.32		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptostreptococcus anaerobius		1.02	0.32		99	99	1	
01200002	**Low discrimination**							**01200002**
					SPO			
Clostridium novyi biovar A		48.30	0.69	SAL 8	99			
Atopobium minutum		36.88	0.63		1			
Clostridium sporogenes		6.72	0.40		99			
01200003	**Very good identification**							**01200003**
					COC	SPO		
Clostridium novyi biovar A		98.11	0.69	SAL 8	1	99		
Atopobium minutum		0.76	0.29		1	1		
Anaerococcus prevotii		0.59	0.13		99	1		
01200010	**Very good identification**							**01200010**
					COC	SPO		
Clostridium novyi biovar A		97.59	0.82	SAL 8	1	99		
Peptostreptococcus anaerobius		1.05	0.32		99	1		
Clostridium difficile		0.43	0.27		1	99		
01200011	**Very good identification**							**01200011**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.97	0.82	SAL 8	1	99	99	
Campylobacter ureolyticus		0.02	-.03		1	1	1	
Peptostreptococcus anaerobius		0.01	-.03		99	99	1	
01200012	**Very good identification**							**01200012**
					SPO			
Clostridium novyi biovar A		99.04	0.69	SAL 8	99			
Atopobium minutum		0.76	0.29		1			
Clostridium sporogenes		0.14	0.06		99			
01200013	**Very good identification**							**01200013**
Clostridium novyi biovar A		100.0	0.69	SAL 8				
01200020	**Low discrimination**							**01200020**
					SPO			
Clostridium novyi biovar A		68.82	0.72	SAL 8	99			
Atopobium minutum		18.55	0.58		1			
Clostridium cadaveris		4.60	0.37		99			
01200021	**Very good identification**							**01200021**
					COC	SPO		
Clostridium novyi biovar A		99.41	0.72	SAL 8	1	99		
Atopobium minutum		0.27	0.24		1	1		
Peptococcus niger		0.10	-.02		99	1		
01200022	**Low discrimination**							**01200022**
					GRA	SPO		
Atopobium minutum		43.16	0.55	SAL 1	99	1		
Clostridium novyi biovar A		42.40	0.59		99	99		
Pseudoflavonifractor capillosus		13.18	0.32		1	1		
01200023	**Very good identification**							**01200023**
					GRA	SPO		
Clostridium novyi biovar A		98.68	0.59	SAL 8	99	99		
Atopobium minutum		1.02	0.21		99	1		
Pseudoflavonifractor capillosus		0.31	-.03		1	1		
01200030	**Very good identification**							**01200030**
					SPO			
Clostridium novyi biovar A		99.59	0.72	SAL 8	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		0.27	0.24		1		
Clostridium cadaveris		0.07	0.03		99		
01200031	**Very good identification**						**01200031**
Clostridium novyi biovar A		100.0	0.72	SAL 8			
01200032	**Very good identification**						**01200032**
					GRA	SPO	
Clostridium novyi biovar A		98.68	0.59	SAL 8	99	99	
Atopobium minutum		1.02	0.21		99	1	
Pseudoflavonifractor capillosus		0.31	-.03		1	1	
01200033	**Very good identification**						**01200033**
Clostridium novyi biovar A		100.0	0.59	SAL 8			
01200040	**Low discrimination**						**01200040**
					COC	SPO	
Clostridium novyi biovar A		84.34	0.79	SAL 8	1	99	
Clostridium botulinum biovar A		11.56	0.52		1	99	
Peptostreptococcus anaerobius		1.36	0.32		99	1	
01200041	**Very good identification**						**01200041**
					GRA	SPO	
Clostridium novyi biovar A		99.81	0.79	SAL 8	99	99	
Clostridium botulinum biovar A		0.14	0.18		99	99	
Campylobacter ureolyticus		0.03	-.03		1	1	
01200042	**Very good identification**						**01200042**
					SPO		
Clostridium novyi biovar A		97.05	0.66	SAL 8	99		
Atopobium minutum		1.12	0.29		1		
Clostridium botulinum biovar A		0.76	0.18		99		
01200043	**Very good identification**						**01200043**
Clostridium novyi biovar A		100.0	0.66	SAL 8			
01200050	**Very good identification**						**01200050**
Clostridium novyi biovar A		99.81	0.79	SAL 8			
Clostridium botulinum biovar A		0.14	0.18				
Clostridium difficile		0.04	0.05				
01200051	**Very good identification**						**01200051**
Clostridium novyi biovar A		100.0	0.79	SAL 8			
01200052	**Very good identification**						**01200052**
Clostridium novyi biovar A		100.0	0.66	SAL 8			
01200053	**Very good identification**						**01200053**
Clostridium novyi biovar A		100.0	0.66	SAL 8			
01200060	**Very good identification**						**01200060**
					SPO		
Clostridium novyi biovar A		98.36	0.69	SAL 8	99		
Clostridium botulinum biovar A		0.54	0.18		99		
Atopobium minutum		0.40	0.24		1		
01200061	**Very good identification**						**01200061**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.69	SAL 8				
01200062	**Good identification**							**01200062**
					GRA	SPO		
Clostridium novyi biovar A		97.81	0.56	SAL 8	99	99		
Atopobium minutum		1.51	0.21		99	1		
Pseudoflavonifractor capillosus		0.46	-.03		1	1		
01200063	**Good identification**							**01200063**
Clostridium novyi biovar A		100.0	0.56	SAL 8				
01200070	**Very good identification**							**01200070**
Clostridium novyi biovar A		100.0	0.69	SAL 8				
01200071	**Very good identification**							**01200071**
Clostridium novyi biovar A		100.0	0.69	SAL 8				
01200072	**Good identification**							**01200072**
Clostridium novyi biovar A		100.0	0.56	SAL 8				
01200073	**Good identification**							**01200073**
Clostridium novyi biovar A		100.0	0.56	SAL 8				
01200101	**Good identification**							**01200101**
					COC	GRA	SPO	
Clostridium novyi biovar A		96.93	0.48	SAL 8	1	99	99	
Campylobacter ureolyticus		2.03	-.03	LAC 1	1	1	1	
Peptostreptococcus anaerobius		1.04	-.03		99	99	1	
01200110	**Good identification**							**01200110**
					COC	SPO		
Clostridium novyi biovar A		98.94	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		1.06	-.03	LAC 1	99	1		
01200111	**Good identification**							**01200111**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01200120	**Low discrimination**							**01200120**
					GRA	SPO		
Clostridium novyi biovar A		66.21	0.37	SAL 8	99	99		
Atopobium minutum		17.84	0.24	LAC 1	99	1		
Prevotella bivia		6.79	-.03		1	1		
01200121	**Good identification**							**01200121**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200130	**Good identification**							**01200130**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200131	**Good identification**							**01200131**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200140	**Low discrimination**							**01200140**
					COC	SPO		
Clostridium novyi biovar A		86.72	0.45	SAL 8	1	99		
Clostridium botulinum biovar A		11.88	0.18	LAC 1	1	99		
Peptostreptococcus anaerobius		1.40	-.03		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01200141	**Good identification**							**01200141**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200150	**Good identification**							**01200150**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200151	**Good identification**							**01200151**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200201	**Good identification**							**01200201**
					COC	GRA	SPO	
Clostridium novyi biovar A		96.93	0.48	SAL 8	1	99	99	
Campylobacter ureolyticus		2.03	-.03	RHA 1	1	1	1	
Peptostreptococcus anaerobius		1.04	-.03		99	99	1	
01200210	**Good identification**							**01200210**
					COC	SPO		
Clostridium novyi biovar A		98.94	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		1.06	-.03	RHA 1	99	1		
01200211	**Good identification**							**01200211**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01200220	**Low discrimination**							**01200220**
					SPO			
Clostridium novyi biovar A		71.03	0.37	SAL 8	99			
Atopobium minutum		19.14	0.24	RHA 1	1			
Clostridium cadaveris		4.74	0.03		99			
01200221	**Good identification**							**01200221**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200230	**Good identification**							**01200230**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200231	**Good identification**							**01200231**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200240	**Low discrimination**							**01200240**
					COC	SPO		
Clostridium novyi biovar A		86.72	0.45	SAL 8	1	99		
Clostridium botulinum biovar A		11.88	0.18	RHA 1	1	99		
Peptostreptococcus anaerobius		1.40	-.03		99	1		
01200241	**Good identification**							**01200241**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200250	**Good identification**							**01200250**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200251	**Good identification**							**01200251**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200401	**Good identification**							**01200401**
					COC	GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		96.93	0.48	SAL 8	1	99	99	
Campylobacter ureolyticus		2.03	-.03	ARA 1	1	1	1	
Peptostreptococcus anaerobius		1.04	-.03		99	99	1	
01200410	**Good identification**							**01200410**
					COC	SPO		
Clostridium novyi biovar A		98.94	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		1.06	-.03	ARA 1	99	1		
01200411	**Good identification**							**01200411**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01200420	**Low discrimination**							**01200420**
					SPO			
Clostridium novyi biovar A		71.03	0.37	SAL 8	99			
Atopobium minutum		19.14	0.24	ARA 1	1			
Clostridium cadaveris		4.74	0.03		99			
01200421	**Good identification**							**01200421**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200430	**Good identification**							**01200430**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200431	**Good identification**							**01200431**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01200440	**Low discrimination**							**01200440**
					COC	SPO		
Clostridium novyi biovar A		86.72	0.45	SAL 8	1	99		
Clostridium botulinum biovar A		11.88	0.18	ARA 1	1	99		
Peptostreptococcus anaerobius		1.40	-.03		99	1		
01200441	**Good identification**							**01200441**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200450	**Good identification**							**01200450**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01200451	**Good identification**							**01200451**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01201000	**Low discrimination**							**01201000**
					COC	SPO		
Clostridium novyi biovar A		52.96	0.66	SAL 8	1	99		
Atopobium minutum		32.11	0.58	GAL 10	1	1		
Peptostreptococcus anaerobius		5.11	0.32		99	1		
01201001	**Very good identification**							**01201001**
					SPO			
Clostridium novyi biovar A		98.83	0.66	SAL 8	99			
Atopobium minutum		0.61	0.24	GAL 10	1			
Clostridium sordellii		0.20	0.03		99			
01201002	**Low discrimination**							**01201002**
					SPO			
Atopobium minutum		67.38	0.55	SAL 1	1			
Clostridium novyi biovar A		29.42	0.53		99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium tortuosum		2.45	0.19		1		
01201003	**Good identification**						**01201003**
					SPO		
Clostridium novyi biovar A		97.74	0.53	SAL 8	99		
Atopobium minutum		2.26	0.21	GAL 10	1		
01201010	**Very good identification**						**01201010**
					COC	SPO	
Clostridium novyi biovar A		99.21	0.66	SAL 8	1	99	
Atopobium minutum		0.61	0.24	GAL 10	1	1	
Peptostreptococcus anaerobius		0.10	-.03		99	1	
01201011	**Very good identification**						**01201011**
Clostridium novyi biovar A		100.0	0.66	SAL 8			
01201012	**Good identification**						**01201012**
					SPO		
Clostridium novyi biovar A		97.74	0.53	SAL 8	99		
Atopobium minutum		2.26	0.21	GAL 10	1		
01201013	**Good identification**						**01201013**
Clostridium novyi biovar A		100.0	0.53	SAL 8			
01201020	**Low discrimination**						**01201020**
					SPO		
Clostridium novyi biovar A		53.19	0.55	SAL 8	99		
Atopobium minutum		43.01	0.50	GAL 10	1		
Eubacterium tortuosum		3.13	0.19		1		
01201021	**Good identification**						**01201021**
					SPO		
Clostridium novyi biovar A		99.19	0.55	SAL 8	99		
Atopobium minutum		0.81	0.15	GAL 10	1		
01201022	**Low discrimination**						**01201022**
					GRA	SPO	
Atopobium minutum		69.04	0.47	SAL 1	99	1	
Clostridium novyi biovar A		22.61	0.42		99	99	
Pseudoflavonifractor capillosus		7.03	0.15		1	1	
01201023	**Good identification**						**01201023**
					SPO		
Clostridium novyi biovar A		97.01	0.42	SAL 8	99		
Atopobium minutum		2.99	0.12	GAL 10	1		
01201030	**Good identification**						**01201030**
					SPO		
Clostridium novyi biovar A		99.19	0.55	SAL 8	99		
Atopobium minutum		0.81	0.15	GAL 10	1		
01201031	**Good identification**						**01201031**
Clostridium novyi biovar A		100.0	0.55	SAL 8			
01201032	**Good identification**						**01201032**
					SPO		
Clostridium novyi biovar A		97.01	0.42	SAL 8	99		
Atopobium minutum		2.99	0.12	GAL 10	1		
01201033	**Good identification**						**01201033**
Clostridium novyi biovar A		100.0	0.42	SAL 8			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01201040	**Very good identification**					**01201040**
					SPO	
Clostridium novyi biovar A		97.12	0.63	SAL 8	99	
Clostridium botulinum biovar A		1.21	0.18	GAL 10	99	
Atopobium minutum		0.89	0.24		1	
01201041	**Very good identification**					**01201041**
Clostridium novyi biovar A		100.0	0.63	SAL 8		
01201042	**Good identification**					**01201042**
					SPO	
Clostridium novyi biovar A		96.65	0.50	SAL 8	99	
Atopobium minutum		3.35	0.21	GAL 10	1	
01201043	**Good identification**					**01201043**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
01201050	**Very good identification**					**01201050**
Clostridium novyi biovar A		100.0	0.63	SAL 8		
01201051	**Very good identification**					**01201051**
Clostridium novyi biovar A		100.0	0.63	SAL 8		
01201052	**Good identification**					**01201052**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
01201053	**Good identification**					**01201053**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
01201060	**Good identification**					**01201060**
					SPO	
Clostridium novyi biovar A		98.79	0.52	SAL 8	99	
Atopobium minutum		1.21	0.15	GAL 10	1	
01201061	**Good identification**					**01201061**
Clostridium novyi biovar A		100.0	0.52	SAL 8		
01201062	**Good identification**					**01201062**
					SPO	
Clostridium novyi biovar A		95.58	0.39	SAL 8	99	
Atopobium minutum		4.42	0.12	GAL 10	1	
01201063	**Good identification**					**01201063**
Clostridium novyi biovar A		100.0	0.39	SAL 8		
01201070	**Good identification**					**01201070**
Clostridium novyi biovar A		100.0	0.52	SAL 8		
01201071	**Good identification**					**01201071**
Clostridium novyi biovar A		100.0	0.52	SAL 8		
01201072	**Good identification**					**01201072**
Clostridium novyi biovar A		100.0	0.39	SAL 8		
01201073	**Good identification**					**01201073**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01201100	**Low discrimination**							**01201100**
					GRA	SPO		
Fusobacterium mortiferum		52.73	0.40	FRU 99	1	1		
Clostridium novyi biovar A		19.50	0.31		99	99		
Atopobium minutum		11.83	0.24		99	1		
01202001	**Acceptable identification**							**01202001**
					COC	GRA	SPO	
Clostridium novyi biovar A		94.82	0.48	SAL 8	1	99	99	
Campylobacter ureolyticus		1.99	-.03	MAN 1	1	1	1	
Peptostreptococcus anaerobius		1.02	-.03		99	99	1	
01202010	**Low discrimination**							**01202010**
					COC	SPO		
Clostridium difficile		79.29	0.43	FRU 99	1	99		
Clostridium novyi biovar A		20.20	0.48		1	99		
Peptostreptococcus anaerobius		0.22	-.03		99	1		
01202011	**Good identification**							**01202011**
Clostridium novyi biovar A		96.19	0.48	SAL 8				
Clostridium difficile		3.81	0.09	MAN 1				
01202020	**Low discrimination**							**01202020**
					SPO			
Clostridium novyi biovar A		50.45	0.37	SAL 8	99			
Pseudoramibacter alactolyticus		25.83	0.14	MAN 1	1			
Atopobium minutum		13.60	0.24		1			
01202021	**Good identification**							**01202021**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01202030	**Low discrimination**							**01202030**
Clostridium novyi biovar A		86.31	0.37	SAL 8				
Clostridium difficile		13.69	0.09	MAN 1				
01202031	**Good identification**							**01202031**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01202040	**Low discrimination**							**01202040**
					COC	SPO	GRA	
Clostridium novyi biovar A		44.90	0.45	SAL 8	1	99	99	
Megasphaera elsdenii		33.17	0.35	MAN 1	99	1	1	
Pseudoramibacter alactolyticus		12.21	0.17		1	1	99	
01202041	**Good identification**							**01202041**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01202050	**Low discrimination**							**01202050**
Clostridium novyi biovar A		76.34	0.45	SAL 8				
Clostridium difficile		23.66	0.21	MAN 1				
01202051	**Good identification**							**01202051**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01204001	**Good identification**							**01204001**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Clostridium novyi biovar A		96.93	0.48	SAL 8	1	99	99	
Campylobacter ureolyticus		2.03	-.03	XYL 1	1	1	1	
Peptostreptococcus anaerobius		1.04	-.03		99	99	1	
01204010	**Good identification**							**01204010**
					COC	SPO		
Clostridium novyi biovar A		96.74	0.48	SAL 8	1	99		
Clostridium difficile		2.22	0.05	XYL 1	1	99		
Peptostreptococcus anaerobius		1.04	-.03		99	1		
01204011	**Good identification**							**01204011**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01204020	**Low discrimination**							**01204020**
					SPO			
Clostridium novyi biovar A		71.03	0.37	SAL 8	99			
Atopobium minutum		19.14	0.24	XYL 1	1			
Clostridium cadaveris		4.74	0.03		99			
01204021	**Good identification**							**01204021**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01204030	**Good identification**							**01204030**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01204031	**Good identification**							**01204031**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01204040	**Low discrimination**							**01204040**
Clostridium glycolicum		89.66	0.49	SAL 1				
Clostridium novyi biovar A		8.97	0.45	FRU 90				
Clostridium botulinum biovar A		1.23	0.18					
01204041	**Acceptable identification**							**01204041**
Clostridium novyi biovar A		90.83	0.45	SAL 8				
Clostridium glycolicum		9.17	0.15	XYL 1				
01204050	**Acceptable identification**							**01204050**
Clostridium novyi biovar A		90.83	0.45	SAL 8				
Clostridium glycolicum		9.17	0.15	XYL 1				
01204051	**Good identification**							**01204051**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01210000	**Low discrimination**							**01210000**
					GRA			
Pseudoramibacter alactolyticus		58.71	0.66	SAL 1	99			
Fusobacterium varium		8.25	0.51		1			
Atopobium minutum		8.13	0.65		99			
01210001	**Low discrimination**							**01210001**
					GRA	SPO		
Clostridium novyi biovar A		82.58	0.66	SAL 8	99	99		
Pseudoramibacter alactolyticus		10.11	0.32	FRU 10	99	1		
Fusobacterium varium		1.42	0.17		1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01210002	**Low discrimination**					**01210002**
					SPO	
Pseudoramibacter alactolyticus		45.71	0.50	SAL 1	1	
Atopobium minutum		37.99	0.62	BGL 10	1	
Clostridium novyi biovar A		5.99	0.53		99	
01210003	**Low discrimination**					**01210003**
					SPO	
Clostridium novyi biovar A		86.64	0.53	SAL 8	99	
Pseudoramibacter alactolyticus		6.68	0.15	FRU 10	1	
Atopobium minutum		5.55	0.28		1	
01210010	**Low discrimination**					**01210010**
					SPO	
Clostridium difficile		76.74	0.61	MAN 90	99	
Clostridium novyi biovar A		19.55	0.66		99	
Pseudoramibacter alactolyticus		2.39	0.32		1	
01210011	**Very good identification**					**01210011**
					SPO	
Clostridium novyi biovar A		96.07	0.66	SAL 8	99	
Clostridium difficile		3.81	0.27	FRU 10	99	
Pseudoramibacter alactolyticus		0.12	-.03		1	
01210012	**Low discrimination**					**01210012**
					SPO	
Clostridium novyi biovar A		72.52	0.53	SAL 8	99	
Clostridium difficile		16.30	0.27	FRU 10	99	
Pseudoramibacter alactolyticus		5.59	0.15		1	
01210013	**Good identification**					**01210013**
Clostridium novyi biovar A		100.0	0.53	SAL 8		
01210020	**Low discrimination**					**01210020**
					SPO	
Pseudoramibacter alactolyticus		67.13	0.53	SAL 1	1	
Atopobium minutum		17.57	0.57		1	
Clostridium novyi biovar A		7.85	0.55		99	
01210021	**Low discrimination**					**01210021**
					SPO	
Clostridium novyi biovar A		89.68	0.55	SAL 8	99	
Pseudoramibacter alactolyticus		7.75	0.19	FRU 10	1	
Atopobium minutum		2.03	0.23		1	
01210022	**Low discrimination**					**01210022**
					SPO	
Atopobium minutum		55.49	0.54	SAL 1	1	
Pseudoramibacter alactolyticus		35.34	0.37		1	
Clostridium novyi biovar A		6.56	0.42		99	
01210023	**Low discrimination**					**01210023**
					SPO	
Clostridium novyi biovar A		87.73	0.42	SAL 8	99	
Atopobium minutum		7.50	0.20	FRU 10	1	
Pseudoramibacter alactolyticus		4.77	0.03		1	
01210030	**Low discrimination**					**01210030**
					SPO	
Clostridium novyi biovar A		78.51	0.55	SAL 8	99	
Clostridium difficile		12.45	0.27	FRU 10	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Pseudoramibacter alactolyticus	6.79	0.19		1			
01210031	**Good identification**							**01210031**
	Clostridium novyi biovar A	100.0	0.55	SAL 8				
01210032	**Low discrimination**							**01210032**
					SPO			
	Clostridium novyi biovar A	87.73	0.42	SAL 8	99			
	Atopobium minutum	7.50	0.20	FRU 10	1			
	Pseudoramibacter alactolyticus	4.77	0.03		1			
01210033	**Good identification**							**01210033**
	Clostridium novyi biovar A	100.0	0.42	SAL 8				
01210040	**Low discrimination**							**01210040**
					COC	SPO	GRA	
	Pseudoramibacter alactolyticus	60.80	0.56	SAL 1	1	1	99	
	Megasphaera elsdenii	17.07	0.57		99	1	1	
	Clostridium novyi biovar A	13.38	0.63		1	99	99	
01210041	**Good identification**							**01210041**
					COC	SPO	GRA	
	Clostridium novyi biovar A	93.99	0.63	SAL 8	1	99	99	
	Pseudoramibacter alactolyticus	4.32	0.21	FRU 10	1	1	99	
	Megasphaera elsdenii	1.21	0.23		99	1	1	
01210042	**Low discrimination**							**01210042**
					COC	SPO	GRA	
	Pseudoramibacter alactolyticus	59.65	0.39	SAL 1	1	1	99	
	Clostridium novyi biovar A	20.84	0.50	BGL 10	1	99	99	
	Megasphaera elsdenii	16.74	0.41		99	1	1	
01210043	**Good identification**							**01210043**
					COC	SPO	GRA	
	Clostridium novyi biovar A	96.43	0.50	SAL 8	1	99	99	
	Pseudoramibacter alactolyticus	2.79	0.05	FRU 10	1	1	99	
	Megasphaera elsdenii	0.78	0.06		99	1	1	
01210050	**Low discrimination**							**01210050**
					SPO			
	Clostridium novyi biovar A	72.79	0.63	SAL 8	99			
	Clostridium difficile	22.56	0.39	FRU 10	99			
	Pseudoramibacter alactolyticus	3.34	0.21		1			
01210051	**Very good identification**							**01210051**
	Clostridium novyi biovar A	99.69	0.63	SAL 8				
	Clostridium difficile	0.31	0.05	FRU 10				
01210052	**Acceptable identification**							**01210052**
					SPO			
	Clostridium novyi biovar A	94.81	0.50	SAL 8	99			
	Pseudoramibacter alactolyticus	2.74	0.05	FRU 10	1			
	Clostridium difficile	1.68	0.05		99			
01210053	**Good identification**							**01210053**
	Clostridium novyi biovar A	100.0	0.50	SAL 8				
01210060	**Low discrimination**							**01210060**
					COC	SPO	GRA	
	Pseudoramibacter alactolyticus	73.69	0.43	SAL 1	1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		22.97	0.52		1	99	99	
Megasphaera elsdenii		1.18	0.23		99	1	1	
01210061	**Good identification**							**01210061**
					SPO			
Clostridium novyi biovar A		96.86	0.52	SAL 8	99			
Pseudoramibacter alactolyticus		3.14	0.09	FRU 10	1			
01210063	**Good identification**							**01210063**
					SPO			
Clostridium novyi biovar A		98.00	0.39	SAL 8	99			
Pseudoramibacter alactolyticus		2.00	-.08	FRU 10	1			
01210070	**Good identification**							**01210070**
					SPO			
Clostridium novyi biovar A		95.70	0.52	SAL 8	99			
Pseudoramibacter alactolyticus		3.10	0.09	FRU 10	1			
Clostridium difficile		1.20	0.05		99			
01210071	**Good identification**							**01210071**
Clostridium novyi biovar A		100.0	0.52	SAL 8				
01210072	**Good identification**							**01210072**
					SPO			
Clostridium novyi biovar A		98.00	0.39	SAL 8	99			
Pseudoramibacter alactolyticus		2.00	-.08	FRU 10	1			
01210073	**Good identification**							**01210073**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01210100	**Low discrimination**							**01210100**
					GRA			
Fusobacterium mortiferum		86.97	0.71		1			
Pseudoramibacter alactolyticus		5.91	0.32		99			
Actinomyces meyeri		3.37	0.38		99			
01210101	**Low discrimination**							**01210101**
					GRA	SPO		
Fusobacterium mortiferum		60.19	0.37	URE 1	1	1		
Clostridium novyi biovar A		33.39	0.31		99	99		
Pseudoramibacter alactolyticus		4.09	-.03		99	1		
01210102	**Low discrimination**							**01210102**
					GRA			
Fusobacterium mortiferum		86.87	0.55	BGL 10	1			
Pseudoramibacter alactolyticus		5.90	0.15		99			
Atopobium minutum		4.90	0.28		99			
01210300	**Acceptable identification**							**01210300**
					GRA			
Fusobacterium mortiferum		90.36	0.37	RHA 1	1			
Pseudoramibacter alactolyticus		6.14	-.03		99			
Actinomyces meyeri		3.50	0.04		99			
01210500	**Low discrimination**							**01210500**
					GRA			
Fusobacterium mortiferum		49.32	0.37	ARA 1	1			
Actinomyces meyeri		47.33	0.28		99			
Pseudoramibacter alactolyticus		3.35	-.03		99			
01211000	**Low discrimination**							**01211000**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		56.55	0.68		1		
Atopobium minutum		27.32	0.57		99		
Pseudoramibacter alactolyticus		5.98	0.32		99		
01211001	**Low discrimination**						**01211001**
					GRA	SPO	
Clostridium novyi biovar A		84.38	0.49	SAL 8	99	99	
Fusobacterium mortiferum		8.89	0.34	FRU 10	1	1	
Atopobium minutum		4.29	0.23	GAL 10	99	1	
01211002	**Low discrimination**						**01211002**
					GRA	SPO	
Atopobium minutum		67.76	0.54	SAL 1	99	1	
Fusobacterium mortiferum		23.38	0.52		1	1	
Clostridium novyi biovar A		3.56	0.36		99	99	
01211003	**Low discrimination**						**01211003**
					GRA	SPO	
Clostridium novyi biovar A		79.46	0.36	SAL 8	99	99	
Atopobium minutum		15.27	0.20	FRU 10	99	1	
Fusobacterium mortiferum		5.27	0.17	GAL 10	1	1	
01211010	**Low discrimination**						**01211010**
					GRA	SPO	
Clostridium novyi biovar A		65.61	0.49	SAL 8	99	99	
Clostridium difficile		23.41	0.27	FRU 10	99	99	
Fusobacterium mortiferum		6.91	0.34	GAL 10	1	1	
01211011	**Good identification**						**01211011**
Clostridium novyi biovar A		100.0	0.49	SAL 8			
01211012	**Low discrimination**						**01211012**
					GRA	SPO	
Clostridium novyi biovar A		79.46	0.36	SAL 8	99	99	
Atopobium minutum		15.27	0.20	FRU 10	99	1	
Fusobacterium mortiferum		5.27	0.17	GAL 10	1	1	
01211013	**Good identification**						**01211013**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01211020	**Low discrimination**						**01211020**
					SPO	CAT	
Atopobium minutum		66.61	0.49	SAL 1	1	1	
Clostridium novyi biovar A		9.92	0.39		99	1	
Propionibacterium acnes		8.18	0.26		1	95	
01211021	**Acceptable identification**						**01211021**
					SPO		
Clostridium novyi biovar A		93.65	0.39	SAL 8	99		
Atopobium minutum		6.35	0.15	FRU 10	1		
01211022	**Low discrimination**						**01211022**
					SPO		
Atopobium minutum		89.52	0.46	SAL 1	1		
Clostridium novyi biovar A		3.53	0.26		99		
Lactobacillus fermentum		3.20	0.28		1		
01211030	**Acceptable identification**						**01211030**
					SPO		
Clostridium novyi biovar A		93.65	0.39	SAL 8	99		
Atopobium minutum		6.35	0.15	FRU 10	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01211031	**Good identification**						**01211031**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
01211040	**Low discrimination**						**01211040**
					SPO		
Clostridium novyi biovar A		45.43	0.46	SAL 8	99		
Clostridium glycolicum		19.56	0.27	FRU 10	99		
Pseudoramibacter alactolyticus		18.77	0.21	GAL 10	1		
01211041	**Good identification**						**01211041**
Clostridium novyi biovar A		100.0	0.46	SAL 8			
01211050	**Good identification**						**01211050**
Clostridium novyi biovar A		97.26	0.46	SAL 8			
Clostridium difficile		2.74	0.05	FRU 10			
01211051	**Good identification**						**01211051**
Clostridium novyi biovar A		100.0	0.46	SAL 8			
01211060	**Low discrimination**						**01211060**
					SPO	CAT	
Clostridium novyi biovar A		60.06	0.36	SAL 8	99	1	
Pseudoramibacter alactolyticus		17.52	0.09	FRU 10	1	1	
Propionibacterium acnes		16.31	0.15	GAL 10	1	95	
01211061	**Good identification**						**01211061**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01211070	**Good identification**						**01211070**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01211071	**Good identification**						**01211071**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01211100	**Very good identification**						**01211100**
					GRA		
Fusobacterium mortiferum		95.26	0.74		1		
Actinomyces meyeri		3.69	0.41		99		
Lactobacillus fermentum		0.77	0.40		99		
01211101	**Acceptable identification**						**01211101**
					GRA	SPO	
Fusobacterium mortiferum		92.60	0.40	URE 1	1	1	
Clostridium novyi biovar A		3.81	0.15		99	99	
Actinomyces meyeri		3.59	0.07		99	1	
01211102	**Good identification**						**01211102**
					GRA		
Fusobacterium mortiferum		91.27	0.58	BGL 10	1		
Lactobacillus fermentum		6.60	0.40		99		
Atopobium minutum		1.15	0.20		99		
01211110	**Acceptable identification**						**01211110**
					GRA	SPO	
Fusobacterium mortiferum		92.60	0.40	MLZ 1	1	1	
Clostridium novyi biovar A		3.81	0.15		99	99	
Actinomyces meyeri		3.59	0.07		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01211120	**Low discrimination**							**01211120**
					GRA	SPO		
Fusobacterium mortiferum		35.44	0.40	NAG 1	1	1		
Clostridium septicum		34.16	0.22		99	99		
Actinomyces meyeri		18.55	0.26		99	1		
01211140	**Acceptable identification**							**01211140**
					GRA	SPO		
Fusobacterium mortiferum		93.80	0.40	SOR 1	1	1		
Actinomyces meyeri		3.64	0.07		99	1		
Clostridium novyi biovar A		2.57	0.12		99	99		
01211300	**Good identification**							**01211300**
					GRA			
Fusobacterium mortiferum		96.27	0.40	RHA 1	1			
Actinomyces meyeri		3.73	0.07		99			
01211400	**Low discrimination**							**01211400**
					GRA			
Lactobacillus fermentum		33.65	0.36	SUC 87	99			
Fusobacterium mortiferum		29.57	0.34	SAL 1	1			
Actinomyces meyeri		16.55	0.20		99			
01211402	**Low discrimination**							**01211402**
					GRA			
Lactobacillus fermentum		72.43	0.36	SUC 87	99			
Atopobium minutum		20.49	0.20	SAL 1	99			
Fusobacterium mortiferum		7.07	0.17		1			
01211500	**Low discrimination**							**01211500**
					GRA			
Fusobacterium mortiferum		36.29	0.40	ARA 1	1			
Actinomyces meyeri		34.83	0.31		99			
Lactobacillus fermentum		28.88	0.40		99			
01211502	**Low discrimination**							**01211502**
					GRA			
Lactobacillus fermentum		84.97	0.40	SUC 87	99			
Fusobacterium mortiferum		11.86	0.24	SAL 1	1			
Actinomyces meyeri		3.17	0.05		99			
01212000	**Low discrimination**							**01212000**
					SPO			
Clostridium difficile		37.60	0.66		99			
Pseudoramibacter alactolyticus		33.91	0.61		1			
Eubacterium limosum		21.56	0.60		1			
01212002	**Low discrimination**							**01212002**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		72.74	0.45	SAL 1	1	1	99	
Megasphaera elsdenii		14.22	0.44	BGL 10	99	1	1	
Clostridium difficile		7.33	0.32		1	99	99	
01212010	**Very good identification**							**01212010**
					SPO			
Clostridium difficile		99.63	0.78		99			
Pseudoramibacter alactolyticus		0.19	0.27		1			
Eubacterium limosum		0.12	0.26		1			
01212011	**Good identification**							**01212011**
					SPO			
Clostridium difficile		97.07	0.43	URE 1	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		2.75	0.31		99			
Pseudoramibacter alactolyticus		0.18	-.07		1			
01212012	**Good identification**							**01212012**
					SPO			
Clostridium difficile		97.13	0.43	BGL 1	99			
Pseudoramibacter alactolyticus		1.99	0.11		1			
Clostridium novyi biovar A		0.49	0.19		99			
01212020	**Low discrimination**							**01212020**
					SPO			
Pseudoramibacter alactolyticus		88.44	0.48	SAL 1	1			
Clostridium difficile		5.61	0.32		99			
Eubacterium limosum		3.22	0.26		1			
01212030	**Good identification**							**01212030**
					SPO			
Clostridium difficile		96.18	0.43	NAG 1	99			
Pseudoramibacter alactolyticus		3.14	0.14		1			
Clostridium novyi biovar A		0.68	0.21		99			
01212040	**Low discrimination**							**01212040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		55.97	0.51	SAL 1	1	1	99	
Megasphaera elsdenii		29.18	0.57		99	1	1	
Clostridium difficile		13.07	0.44		1	99	99	
01212050	**Good identification**							**01212050**
					COC	SPO	GRA	
Clostridium difficile		98.32	0.56	SOR 5	1	99	99	
Pseudoramibacter alactolyticus		0.87	0.17		1	1	99	
Megasphaera elsdenii		0.45	0.23		99	1	1	
01212060	**Acceptable identification**							**01212060**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		94.44	0.38	SAL 1	1	1	99	
Megasphaera elsdenii		2.82	0.23		99	1	1	
Clostridium difficile		1.26	0.10		1	99	99	
01212110	**Good identification**							**01212110**
					SPO			
Clostridium difficile		99.81	0.43	LAC 1	99			
Pseudoramibacter alactolyticus		0.19	-.07		1			
01212210	**Good identification**							**01212210**
					SPO			
Clostridium difficile		99.81	0.43	RHA 1	99			
Pseudoramibacter alactolyticus		0.19	-.07		1			
01212410	**Good identification**							**01212410**
					SPO			
Clostridium difficile		99.81	0.43	ARA 1	99			
Pseudoramibacter alactolyticus		0.19	-.07		1			
01213010	**Good identification**							**01213010**
Clostridium difficile		99.29	0.43	GAL 1				
Clostridium novyi biovar A		0.31	0.15					
Clostridium innocuum		0.22	0.01					
01213100	**Acceptable identification**							**01213100**
					GRA	SPO		
Fusobacterium mortiferum		91.73	0.40	MAN 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		3.56	0.07		99	1		
Clostridium difficile		2.48	-.03		99	99		
01214000	**Low discrimination**							**01214000**
					GRA	SPO		
Clostridium glycolicum		42.98	0.43	SAL 1	99	99		
Pseudoramibacter alactolyticus		18.33	0.32	SOR 95	99	1		
Fusobacterium varium		10.62	0.28		1	1		
01214010	**Acceptable identification**							**01214010**
Clostridium difficile		93.00	0.39	MAN 90				
Clostridium novyi biovar A		4.55	0.31	XYL 5				
Clostridium glycolicum		1.31	0.09					
01214040	**Very good identification**							**01214040**
					COC	SPO	GRA	
Clostridium glycolicum		99.09	0.65	SAL 1	1	99	99	
Pseudoramibacter alactolyticus		0.56	0.21		1	1	99	
Megasphaera elsdenii		0.16	0.23		99	1	1	
01214100	**Low discrimination**							**01214100**
					GRA	SPO		
Fusobacterium mortiferum		49.12	0.52	XYL 7	1	1		
Actinomyces meyeri		49.12	0.43		99	1		
Clostridium glycolicum		1.05	0.09		99	99		
01214102	**Low discrimination**							**01214102**
					GRA			
Fusobacterium mortiferum		78.23	0.36	XYL 7	1			
Actinomyces meyeri		21.77	0.17	BGL 10	99			
01215000	**Low discrimination**							**01215000**
					GRA	SPO		
Fusobacterium mortiferum		46.76	0.49	XYL 7	1	1		
Actinomyces meyeri		27.27	0.35		99	1		
Clostridium glycolicum		17.10	0.27		99	99		
01215040	**Good identification**							**01215040**
					GRA	SPO		
Clostridium glycolicum		99.73	0.49	SAL 1	99	99		
Fusobacterium mortiferum		0.15	0.15	GAL 10	1	1		
Clostridium novyi biovar A		0.12	0.12		99	99		
01215100	**Low discrimination**							**01215100**
					GRA			
Fusobacterium mortiferum		48.48	0.55	XYL 7	1			
Actinomyces meyeri		48.48	0.46		99			
Lactobacillus fermentum		3.04	0.36		99			
01215102	**Low discrimination**							**01215102**
					GRA			
Fusobacterium mortiferum		54.26	0.39	XYL 7	1			
Lactobacillus fermentum		30.63	0.36	BGL 10	99			
Actinomyces meyeri		15.10	0.20		99			
01215500	**Low discrimination**							**01215500**
					GRA			
Actinomyces meyeri		77.44	0.35	SUC 93	99			
Lactobacillus fermentum		19.43	0.36	SAL 1	99			
Fusobacterium mortiferum		3.13	0.21		1			
01215502	**Low discrimination**							**01215502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		89.03	0.36	SUC 87			
Actinomyces meyeri		10.98	0.10	SAL 1			
01216000	**Low discrimination**						**01216000**
					SPO		
Clostridium difficile		75.30	0.44	XYL 5	99		
Pseudoramibacter alactolyticus		13.03	0.27		1		
Eubacterium limosum		8.29	0.26		1		
01216010	**Good identification**						**01216010**
					SPO		
Clostridium difficile		99.96	0.56	XYL 5	99		
Pseudoramibacter alactolyticus		0.04	-.07		1		
01220000	**Low discrimination**						**01220000**
					COC	SPO	
Clostridium botulinum biovar A		59.76	0.51	SAL 1	1	99	
Clostridium sporogenes		19.20	0.40		1	99	
Peptostreptococcus anaerobius		8.38	0.32		99	1	
01220001	**Low discrimination**						**01220001**
Clostridium novyi biovar A		88.28	0.48	SAL 8			
Clostridium botulinum biovar A		6.75	0.17	TRE 1			
Clostridium sporogenes		2.17	0.06				
01220010	**Low discrimination**						**01220010**
Clostridium novyi biovar A		87.37	0.48	SAL 8			
Clostridium botulinum biovar A		6.68	0.17	TRE 1			
Clostridium difficile		2.87	0.08				
01220011	**Good identification**						**01220011**
Clostridium novyi biovar A		100.0	0.48	SAL 8			
01220020	**Low discrimination**						**01220020**
Clostridium novyi biovar A		47.78	0.37	SAL 8			
Clostridium cadaveris		16.62	0.15	TRE 1			
Clostridium botulinum biovar A		14.61	0.17				
01220021	**Good identification**						**01220021**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01220030	**Good identification**						**01220030**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01220031	**Good identification**						**01220031**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01220040	**Low discrimination**						**01220040**
Clostridium botulinum biovar A		76.31	0.42	SAL 1			
Clostridium novyi biovar A		22.50	0.45				
Clostridium sporogenes		0.83	0.06				
01220041	**Good identification**						**01220041**
Clostridium novyi biovar A		96.69	0.45	SAL 8			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		3.31	0.08	TRE 1			
01220050	**Good identification**						**01220050**
Clostridium novyi biovar A		96.69	0.45	SAL 8			
Clostridium botulinum biovar A		3.31	0.08	TRE 1			
01220051	**Good identification**						**01220051**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01230000	**Low discrimination**						**01230000**
					GRA	SPO	
Fusobacterium mortiferum		40.91	0.56		1	1	
Pseudoramibacter alactolyticus		21.71	0.32		99	1	
Clostridium difficile		10.73	0.30		99	99	
01230002	**Low discrimination**						**01230002**
					GRA	SPO	
Fusobacterium mortiferum		38.05	0.40	BGL 10	1	1	
Pseudoramibacter alactolyticus		20.19	0.15		99	1	
Clostridium sporogenes		19.30	0.19		99	99	
01230010	**Good identification**						**01230010**
					GRA	SPO	
Clostridium difficile		95.58	0.42	TRE 7	99	99	
Clostridium novyi biovar A		3.27	0.31	MAN 90	99	99	
Fusobacterium mortiferum		0.75	0.22		1	1	
01230100	**Very good identification**						**01230100**
					GRA		
Fusobacterium mortiferum		99.53	0.62		1		
Pseudoramibacter alactolyticus		0.23	-.03		99		
Actinomyces meyeri		0.13	0.04		99		
01230102	**Good identification**						**01230102**
Fusobacterium mortiferum		100.0	0.46	BGL 10			
01231000	**Very good identification**						**01231000**
					GRA		
Fusobacterium mortiferum		97.07	0.59		1		
Atopobium minutum		1.59	0.23		99		
Pseudoramibacter alactolyticus		0.35	-.03		99		
01231002	**Low discrimination**						**01231002**
					GRA		
Fusobacterium mortiferum		89.62	0.43	BGL 10	1		
Atopobium minutum		8.78	0.20		99		
Lactobacillus jensenii		1.60	0.03		99		
01231100	**Very good identification**						**01231100**
					GRA	SPO	
Fusobacterium mortiferum		99.77	0.65		1	1	
Actinomyces meyeri		0.13	0.07		99	1	
Clostridium septicum		0.10	-.04		99	99	
01231102	**Good identification**						**01231102**
Fusobacterium mortiferum		100.0	0.49	BGL 10			
01232000	**Low discrimination**						**01232000**
					SPO		
Clostridium difficile		81.49	0.47	TRE 7	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		9.86	0.27		1			
Eubacterium limosum		6.27	0.26		1			
01232010	**Very good identification**							**01232010**
					SPO			
Clostridium difficile		99.98	0.58	TRE 7	99			
Pseudoramibacter alactolyticus		0.03	-.07		1			
01232050	**Good identification**							**01232050**
Clostridium difficile		100.0	0.36	TRE 7				
01234000	**Low discrimination**							**01234000**
					GRA	SPO		
Fusobacterium mortiferum		68.36	0.37	XYL 7	1	1		
Clostridium difficile		12.54	0.08		99	99		
Clostridium glycolicum		11.41	0.09		99	99		
01234100	**Good identification**							**01234100**
					GRA			
Fusobacterium mortiferum		96.73	0.43	XYL 7	1			
Actinomyces meyeri		3.27	0.09		99			
01235000	**Good identification**							**01235000**
Fusobacterium mortiferum		100.0	0.40	XYL 7				
01235100	**Good identification**							**01235100**
					GRA			
Fusobacterium mortiferum		96.73	0.46	XYL 7	1			
Actinomyces meyeri		3.27	0.12		99			
01236010	**Good identification**							**01236010**
Clostridium difficile		100.0	0.36	TRE 7				
01240001	**Good identification**							**01240001**
					COC	GRA	SPO	
Clostridium novyi biovar A		96.93	0.48	SAL 8	1	99	99	
Campylobacter ureolyticus		2.03	-.03	CEL 1	1	1	1	
Peptostreptococcus anaerobius		1.04	-.03		99	99	1	
01240010	**Good identification**							**01240010**
					COC	SPO		
Clostridium novyi biovar A		96.74	0.48	SAL 8	1	99		
Clostridium difficile		2.22	0.05	CEL 1	1	99		
Peptostreptococcus anaerobius		1.04	-.03		99	1		
01240011	**Good identification**							**01240011**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01240020	**Low discrimination**							**01240020**
					SPO			
Clostridium novyi biovar A		71.03	0.37	SAL 8	99			
Atopobium minutum		19.14	0.24	CEL 1	1			
Clostridium cadaveris		4.74	0.03		99			
01240021	**Good identification**							**01240021**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01240030	**Good identification**							**01240030**
Clostridium novyi biovar A		100.0	0.37	SAL 8				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01240031	**Good identification**						**01240031**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01240040	**Low discrimination**						**01240040**
					COC	SPO	
Clostridium novyi biovar A		86.72	0.45	SAL 8	1	99	
Clostridium botulinum biovar A		11.88	0.18	CEL 1	1	99	
Peptostreptococcus anaerobius		1.40	-.03		99	1	
01240041	**Good identification**						**01240041**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01240050	**Good identification**						**01240050**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01240051	**Good identification**						**01240051**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01243000	**Good identification**						**01243000**
Clostridium innocuum		100.0	0.37	MNS 99			
01243002	**Good identification**						**01243002**
Clostridium innocuum		100.0	0.37	MNS 99			
01250000	**Low discrimination**						**01250000**
					GRA	SPO	
Fusobacterium mortiferum		52.26	0.57		1	1	
Pseudoramibacter alactolyticus		24.85	0.32		99	1	
Clostridium difficile		8.59	0.28		99	99	
01250002	**Low discrimination**						**01250002**
					GRA		
Fusobacterium mortiferum		49.58	0.40	BGL 10	1		
Pseudoramibacter alactolyticus		23.57	0.15		99		
Atopobium minutum		19.59	0.28		99		
01250010	**Acceptable identification**						**01250010**
					GRA	SPO	
Clostridium difficile		93.68	0.39	CEL 5	99	99	
Clostridium novyi biovar A		4.58	0.31	MAN 90	99	99	
Fusobacterium mortiferum		1.18	0.23		1	1	
01250100	**Very good identification**						**01250100**
					GRA		
Fusobacterium mortiferum		99.68	0.63		1		
Pseudoramibacter alactolyticus		0.21	-.03		99		
Actinomyces meyeri		0.12	0.04		99		
01250102	**Good identification**						**01250102**
Fusobacterium mortiferum		100.0	0.47	BGL 10			
01251000	**Good identification**						**01251000**
					GRA	SPO	
Fusobacterium mortiferum		91.58	0.60		1	1	
Clostridium innocuum		4.39	0.25		99	99	
Lactobacillus jensenii		1.54	0.19		99	1	
01251002	**Low discrimination**						**01251002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		59.57	0.43	BGL 10	1	1	
Clostridium innocuum		25.72	0.25		99	99	
Lactobacillus jensenii		6.02	0.16		99	1	
01251100	**Very good identification**						**01251100**
					GRA		
Fusobacterium mortiferum		99.38	0.66		1		
Lactobacillus catenaformis		0.41	0.15		99		
Actinomyces meyeri		0.12	0.07		99		
01251102	**Acceptable identification**						**01251102**
					GRA		
Fusobacterium mortiferum		94.91	0.50	BGL 10	1		
Lactobacillus catenaformis		3.54	0.15		99		
Lactobacillus acidophilus		0.90	0.02		99		
01252000	**Low discrimination**						**01252000**
					SPO		
Clostridium difficile		63.40	0.44	CEL 5	99		
Clostridium innocuum		16.07	0.37		99		
Pseudoramibacter alactolyticus		10.97	0.27		1		
01252002	**Low discrimination**						**01252002**
					SPO		
Clostridium innocuum		88.45	0.37	MNS 99	99		
Pseudoramibacter alactolyticus		6.71	0.11	GAL 95	1		
Clostridium difficile		3.52	0.10		99		
01252010	**Good identification**						**01252010**
					SPO		
Clostridium difficile		99.75	0.56	CEL 5	99		
Clostridium innocuum		0.22	0.13		99		
Pseudoramibacter alactolyticus		0.04	-.07		1		
01253000	**Very good identification**						**01253000**
					GRA	SPO	
Clostridium innocuum		99.54	0.59	MNS 99	99	99	
Fusobacterium mortiferum		0.21	0.26		1	1	
Clostridium difficile		0.21	0.10		99	99	
01253002	**Very good identification**						**01253002**
Clostridium innocuum		100.0	0.59	MNS 99			
01253010	**Low discrimination**						**01253010**
Clostridium innocuum		80.28	0.35	MNS 99			
Clostridium difficile		19.73	0.21	MLZ 4			
01253012	**Good identification**						**01253012**
Clostridium innocuum		100.0	0.35	MNS 99			
01253400	**Good identification**						**01253400**
Clostridium innocuum		100.0	0.46	MNS 99			
01253402	**Good identification**						**01253402**
Clostridium innocuum		100.0	0.46	MNS 99			
01254000	**Low discrimination**						**01254000**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		73.24	0.38	XYL 7	1	1	
Clostridium glycolicum		10.96	0.09		99	99	
Clostridium difficile		8.42	0.06		99	99	
01254100	**Good identification**						**01254100**
					GRA		
Fusobacterium mortiferum		97.06	0.44	XYL 7	1		
Actinomyces meyeri		2.94	0.09		99		
01255000	**Good identification**						**01255000**
					GRA	SPO	
Fusobacterium mortiferum		96.76	0.41	XYL 7	1	1	
Clostridium innocuum		3.25	0.03		99	99	
01255100	**Good identification**						**01255100**
					GRA		
Fusobacterium mortiferum		97.06	0.47	XYL 7	1		
Actinomyces meyeri		2.94	0.12		99		
01257000	**Good identification**						**01257000**
Clostridium innocuum		100.0	0.37	MNS 99			
01257002	**Good identification**						**01257002**
Clostridium innocuum		100.0	0.37	MNS 99			
01270000	**Acceptable identification**						**01270000**
					GRA	SPO	
Fusobacterium mortiferum		94.56	0.48		1	1	
Clostridium difficile		3.92	0.08		99	99	
Pseudoramibacter alactolyticus		1.52	-.03		99	1	
01270100	**Good identification**						**01270100**
Fusobacterium mortiferum		100.0	0.54				
01270102	**Good identification**						**01270102**
Fusobacterium mortiferum		100.0	0.38	BGL 10			
01271000	**Low discrimination**						**01271000**
					GRA	SPO	
Fusobacterium mortiferum		81.59	0.51		1	1	
Lactobacillus jensenii		13.80	0.28		99	1	
Clostridium innocuum		4.60	0.17		99	99	
01271100	**Very good identification**						**01271100**
					GRA	SPO	
Fusobacterium mortiferum		99.07	0.57		1	1	
Clostridium septicum		0.93	0.05		99	99	
01271102	**Acceptable identification**						**01271102**
					GRA		
Fusobacterium mortiferum		90.39	0.41	BGL 10	1		
Lactobacillus acidophilus		9.61	0.11		99		
01271120	**Good identification**						**01271120**
					GRA	SPO	
Clostridium septicum		98.92	0.39	MNS 99	99	99	
Fusobacterium mortiferum		1.08	0.23	MLT 94	1	1	
01272010	**Good identification**						**01272010**
Clostridium difficile		99.00	0.36	TRE 7			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium innocuum		1.00	0.06	CEL 5				
01273000	**Good identification**							**01273000**
					GRA	SPO		
Clostridium innocuum		99.82	0.51	MNS 99	99	99		
Fusobacterium mortiferum		0.18	0.17		1	1		
01273002	**Good identification**							**01273002**
Clostridium innocuum		100.0	0.51	MNS 99				
01273400	**Good identification**							**01273400**
Clostridium innocuum		100.0	0.38	MNS 99				
01273402	**Good identification**							**01273402**
Clostridium innocuum		100.0	0.38	MNS 99				
01275100	**Good identification**							**01275100**
					COC	GRA		
Fusobacterium mortiferum		96.82	0.38	XYL 7	1	1		
Atopobium parvulum		3.18	0.01		99	99		
01300000	**Low discrimination**							**01300000**
					COC	GRA	SPO	
Clostridium novyi biovar A		36.14	0.82	SAL 8	1	99	99	
Peptostreptococcus anaerobius		25.58	0.63		99	99	1	
Prevotella disiens		14.10	0.57		1	1	1	
01300001	**Very good identification**							**01300001**
					COC	SPO		
Clostridium novyi biovar A		97.16	0.82	SAL 8	1	99		
Clostridium sordellii		1.09	0.32		1	99		
Peptostreptococcus anaerobius		0.70	0.29		99	1		
01300002	**Low discrimination**							**01300002**
					COC	SPO		
Clostridium novyi biovar A		73.23	0.69	SAL 8	1	99		
Clostridium sporogenes		18.93	0.45		1	99		
Peptostreptococcus anaerobius		2.97	0.29		99	1		
01300003	**Very good identification**							**01300003**
Clostridium novyi biovar A		99.64	0.69	SAL 8				
Clostridium sporogenes		0.26	0.11					
Clostridium sordellii		0.06	-.03					
01300010	**Very good identification**							**01300010**
					COC	GRA	SPO	
Clostridium novyi biovar A		98.25	0.82	SAL 8	1	99	99	
Peptostreptococcus anaerobius		0.70	0.29		99	99	1	
Prevotella disiens		0.39	0.23		1	1	1	
01300011	**Very good identification**							**01300011**
					COC	SPO		
Clostridium novyi biovar A		99.98	0.82	SAL 8	1	99		
Clostridium sordellii		0.01	-.03		1	99		
Peptostreptococcus anaerobius		0.01	-.06		99	1		
01300012	**Very good identification**							**01300012**
					COC	SPO		
Clostridium novyi biovar A		99.70	0.69	SAL 8	1	99		
Clostridium sporogenes		0.26	0.11		1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptostreptococcus anaerobius		0.04	-.06		99	1	
01300013	**Very good identification**						**01300013**
Clostridium novyi biovar A		100.0	0.69	SAL 8			
01300020	**Low discrimination**						**01300020**
					COC	SPO	
Clostridium novyi biovar A		88.01	0.72	SAL 8	1	99	
Clostridium bifermentans		5.90	0.39		1	99	
Peptostreptococcus anaerobius		2.52	0.29		99	1	
01300021	**Very good identification**						**01300021**
Clostridium novyi biovar A		99.86	0.72	SAL 8			
Clostridium bifermentans		0.07	0.05				
Clostridium sordellii		0.05	-.03				
01300022	**Very good identification**						**01300022**
					SPO		
Clostridium novyi biovar A		97.15	0.59	SAL 8	99		
Clostridium sporogenes		1.02	0.11		99		
Atopobium minutum		1.00	0.21		1		
01300023	**Very good identification**						**01300023**
Clostridium novyi biovar A		100.0	0.59	SAL 8			
01300030	**Very good identification**						**01300030**
					COC	SPO	
Clostridium novyi biovar A		99.90	0.72	SAL 8	1	99	
Clostridium bifermentans		0.07	0.05		1	99	
Peptostreptococcus anaerobius		0.03	-.06		99	1	
01300031	**Very good identification**						**01300031**
Clostridium novyi biovar A		100.0	0.72	SAL 8			
01300032	**Very good identification**						**01300032**
Clostridium novyi biovar A		100.0	0.59	SAL 8			
01300033	**Very good identification**						**01300033**
Clostridium novyi biovar A		100.0	0.59	SAL 8			
01300040	**Low discrimination**						**01300040**
Clostridium novyi biovar A		85.89	0.79	SAL 8			
Clostridium botulinum biovar A		9.63	0.51				
Clostridium bifermentans		1.41	0.36				
01300041	**Very good identification**						**01300041**
Clostridium novyi biovar A		99.78	0.79	SAL 8			
Clostridium botulinum biovar A		0.11	0.17				
Clostridium sordellii		0.07	0.08				
01300042	**Very good identification**						**01300042**
					COC	GRA	
Clostridium novyi biovar A		98.14	0.66	SAL 8	1	99	
Megasphaera elsdenii		0.69	0.22		99	1	
Clostridium botulinum biovar A		0.63	0.17		1	99	
01300043	**Very good identification**						**01300043**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.66	SAL 8				
01300050	**Very good identification**							**01300050**
Clostridium novyi biovar A		99.85	0.79	SAL 8				
Clostridium botulinum biovar A		0.11	0.17					
Clostridium bifermentans		0.02	0.02					
01300051	**Very good identification**							**01300051**
Clostridium novyi biovar A		100.0	0.79	SAL 8				
01300052	**Very good identification**							**01300052**
Clostridium novyi biovar A		100.0	0.66	SAL 8				
01300053	**Very good identification**							**01300053**
Clostridium novyi biovar A		100.0	0.66	SAL 8				
01300060	**Very good identification**							**01300060**
Clostridium novyi biovar A		99.05	0.69	SAL 8				
Clostridium botulinum biovar A		0.45	0.17					
Clostridium bifermentans		0.42	0.16					
01300061	**Very good identification**							**01300061**
Clostridium novyi biovar A		100.0	0.69	SAL 8				
01300062	**Good identification**							**01300062**
Clostridium novyi biovar A		100.0	0.56	SAL 8				
01300063	**Good identification**							**01300063**
Clostridium novyi biovar A		100.0	0.56	SAL 8				
01300070	**Very good identification**							**01300070**
Clostridium novyi biovar A		100.0	0.69	SAL 8				
01300071	**Very good identification**							**01300071**
Clostridium novyi biovar A		100.0	0.69	SAL 8				
01300072	**Good identification**							**01300072**
Clostridium novyi biovar A		100.0	0.56	SAL 8				
01300073	**Good identification**							**01300073**
Clostridium novyi biovar A		100.0	0.56	SAL 8				
01300100	**Low discrimination**							**01300100**
					COC	GRA	SPO	
Clostridium novyi biovar A		34.90	0.48	SAL 8	1	99	99	
Peptostreptococcus anaerobius		24.71	0.29	LAC 1	99	99	1	
Prevotella disiens		13.62	0.23		1	1	1	
01300101	**Good identification**							**01300101**
					COC	SPO		
Clostridium novyi biovar A		98.20	0.48	SAL 8	1	99		
Clostridium sordellii		1.10	-.03	LAC 1	1	99		
Peptostreptococcus anaerobius		0.70	-.06		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01300110	**Good identification**							**01300110**
					COC	SPO		
Clostridium novyi biovar A		99.29	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		0.71	-.06	LAC 1	99	1		
01300111	**Good identification**							**01300111**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01300121	**Acceptable identification**							**01300121**
					GRA	SPO		
Clostridium novyi biovar A		90.70	0.37	SAL 8	99	99		
Prevotella bivia		9.30	-.03	LAC 1	1	1		
01300130	**Acceptable identification**							**01300130**
					GRA	SPO		
Clostridium novyi biovar A		90.70	0.37	SAL 8	99	99		
Prevotella bivia		9.30	-.03	LAC 1	1	1		
01300131	**Good identification**							**01300131**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01300140	**Low discrimination**							**01300140**
Clostridium novyi biovar A		86.92	0.45	SAL 8				
Clostridium botulinum biovar A		9.74	0.17	LAC 1				
Clostridium bifermentans		1.43	0.02					
01300141	**Good identification**							**01300141**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300150	**Good identification**							**01300150**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300151	**Good identification**							**01300151**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300200	**Low discrimination**							**01300200**
					COC	GRA	SPO	
Clostridium novyi biovar A		36.01	0.48	SAL 8	1	99	99	
Peptostreptococcus anaerobius		25.49	0.29	RHA 1	99	99	1	
Prevotella disiens		14.05	0.23		1	1	1	
01300201	**Good identification**							**01300201**
					COC	SPO		
Clostridium novyi biovar A		98.20	0.48	SAL 8	1	99		
Clostridium sordellii		1.10	-.03	RHA 1	1	99		
Peptostreptococcus anaerobius		0.70	-.06		99	1		
01300210	**Good identification**							**01300210**
					COC	SPO		
Clostridium novyi biovar A		99.29	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		0.71	-.06	RHA 1	99	1		
01300211	**Good identification**							**01300211**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01300220	**Acceptable identification**							**01300220**
					COC	SPO		
Clostridium novyi biovar A		91.27	0.37	SAL 8	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		6.12	0.05	RHA 1	1	99		
Peptostreptococcus anaerobius		2.61	-.06		99	1		
01300221	**Good identification**							**01300221**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01300230	**Good identification**							**01300230**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01300231	**Good identification**							**01300231**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01300240	**Low discrimination**							**01300240**
Clostridium novyi biovar A		86.92	0.45	SAL 8				
Clostridium botulinum biovar A		9.74	0.17	RHA 1				
Clostridium bifermentans		1.43	0.02					
01300241	**Good identification**							**01300241**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300250	**Good identification**							**01300250**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300251	**Good identification**							**01300251**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300400	**Low discrimination**							**01300400**
					COC	GRA	SPO	
Clostridium novyi biovar A		35.79	0.48	SAL 8	1	99	99	
Peptostreptococcus anaerobius		25.33	0.29	ARA 1	99	99	1	
Prevotella disiens		13.96	0.23		1	1	1	
01300401	**Good identification**							**01300401**
					COC	SPO		
Clostridium novyi biovar A		98.20	0.48	SAL 8	1	99		
Clostridium sordellii		1.10	-.03	ARA 1	1	99		
Peptostreptococcus anaerobius		0.70	-.06		99	1		
01300410	**Good identification**							**01300410**
					COC	SPO		
Clostridium novyi biovar A		99.29	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		0.71	-.06	ARA 1	99	1		
01300411	**Good identification**							**01300411**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01300420	**Acceptable identification**							**01300420**
					COC	SPO		
Clostridium novyi biovar A		91.27	0.37	SAL 8	1	99		
Clostridium bifermentans		6.12	0.05	ARA 1	1	99		
Peptostreptococcus anaerobius		2.61	-.06		99	1		
01300421	**Good identification**							**01300421**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01300430	**Good identification**							**01300430**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01300431	**Good identification**							**01300431**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01300440	**Low discrimination**							**01300440**
Clostridium novyi biovar A		86.92	0.45	SAL 8				
Clostridium botulinum biovar A		9.74	0.17	ARA 1				
Clostridium bifermentans		1.43	0.02					
01300441	**Good identification**							**01300441**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300450	**Good identification**							**01300450**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01300451	**Good identification**							**01300451**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01301000	**Low discrimination**							**01301000**
					COC	GRA	SPO	
Clostridium novyi biovar A		84.19	0.66	SAL 8	1	99	99	
Peptostreptococcus anaerobius		5.42	0.29	GAL 10	99	99	1	
Prevotella disiens		2.99	0.23		1	1	1	
01301001	**Very good identification**							**01301001**
					COC	SPO		
Clostridium novyi biovar A		98.19	0.66	SAL 8	1	99		
Clostridium sordellii		1.75	0.19	GAL 10	1	99		
Peptostreptococcus anaerobius		0.06	-.06		99	1		
01301002	**Good identification**							**01301002**
					SPO			
Clostridium novyi biovar A		95.21	0.53	SAL 8	99			
Clostridium sporogenes		2.24	0.11	GAL 10	99			
Atopobium minutum		2.20	0.21		1			
01301003	**Good identification**							**01301003**
Clostridium novyi biovar A		100.0	0.53	SAL 8				
01301010	**Very good identification**							**01301010**
					COC	SPO		
Clostridium novyi biovar A		99.94	0.66	SAL 8	1	99		
Peptostreptococcus anaerobius		0.07	-.06	GAL 10	99	1		
01301011	**Very good identification**							**01301011**
Clostridium novyi biovar A		100.0	0.66	SAL 8				
01301012	**Good identification**							**01301012**
Clostridium novyi biovar A		100.0	0.53	SAL 8				
01301013	**Good identification**							**01301013**
Clostridium novyi biovar A		100.0	0.53	SAL 8				
01301020	**Good identification**							**01301020**
					COC	SPO		
Clostridium novyi biovar A		97.68	0.55	SAL 8	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		0.80	0.15	GAL 10	1	1	
Gemella morbillorum		0.67	0.09		99	1	
01301021	**Good identification**						**01301021**
Clostridium novyi biovar A		100.0	0.55	SAL 8			
01301022	**Good identification**						**01301022**
					SPO		
Clostridium novyi biovar A		97.01	0.42	SAL 8	99		
Atopobium minutum		2.99	0.12	GAL 10	1		
01301023	**Good identification**						**01301023**
Clostridium novyi biovar A		100.0	0.42	SAL 8			
01301030	**Good identification**						**01301030**
Clostridium novyi biovar A		100.0	0.55	SAL 8			
01301031	**Good identification**						**01301031**
Clostridium novyi biovar A		100.0	0.55	SAL 8			
01301032	**Good identification**						**01301032**
Clostridium novyi biovar A		100.0	0.42	SAL 8			
01301033	**Good identification**						**01301033**
Clostridium novyi biovar A		100.0	0.42	SAL 8			
01301040	**Very good identification**						**01301040**
Clostridium novyi biovar A		98.09	0.63	SAL 8			
Clostridium botulinum biovar A		1.00	0.17	GAL 10			
Clostridium glycolicum		0.57	0.11				
01301041	**Very good identification**						**01301041**
Clostridium novyi biovar A		99.89	0.63	SAL 8			
Clostridium sordellii		0.11	-.05	GAL 10			
01301042	**Good identification**						**01301042**
Clostridium novyi biovar A		100.0	0.50	SAL 8			
01301043	**Good identification**						**01301043**
Clostridium novyi biovar A		100.0	0.50	SAL 8			
01301050	**Very good identification**						**01301050**
Clostridium novyi biovar A		100.0	0.63	SAL 8			
01301051	**Very good identification**						**01301051**
Clostridium novyi biovar A		100.0	0.63	SAL 8			
01301052	**Good identification**						**01301052**
Clostridium novyi biovar A		100.0	0.50	SAL 8			
01301053	**Good identification**						**01301053**
Clostridium novyi biovar A		100.0	0.50	SAL 8			

Code		ID	T	tests	Differentiation tests			Code
Taxon		score	index	against				
01301060	**Good identification**							**01301060**
Clostridium novyi biovar A		100.0	0.52	SAL 8				
01301061	**Good identification**							**01301061**
Clostridium novyi biovar A		100.0	0.52	SAL 8				
01301062	**Good identification**							**01301062**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01301063	**Good identification**							**01301063**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01301070	**Good identification**							**01301070**
Clostridium novyi biovar A		100.0	0.52	SAL 8				
01301071	**Good identification**							**01301071**
Clostridium novyi biovar A		100.0	0.52	SAL 8				
01301072	**Good identification**							**01301072**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01301073	**Good identification**							**01301073**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01302000	**Low discrimination**							**01302000**
					COC	GRA	SPO	
Megasphaera elsdenii		28.62	0.41	SAL 1	99	1	1	
Clostridium novyi biovar A		25.83	0.48	FRU 95	1	99	99	
Peptostreptococcus anaerobius		18.29	0.29		99	99	1	
01302001	**Good identification**							**01302001**
					COC	GRA		
Clostridium novyi biovar A		97.13	0.48	SAL 8	1	99		
Clostridium sordellii		1.09	-.03	MAN 1	1	99		
Megasphaera elsdenii		1.09	0.07		99	1		
01302010	**Acceptable identification**							**01302010**
					COC	GRA		
Clostridium novyi biovar A		94.52	0.48	SAL 8	1	99		
Clostridium difficile		3.75	0.09	MAN 1	1	99		
Megasphaera elsdenii		1.06	0.07		99	1		
01302011	**Good identification**							**01302011**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01302020	**Low discrimination**							**01302020**
					COC	GRA		
Clostridium novyi biovar A		87.69	0.37	SAL 8	1	99		
Clostridium bifermentans		5.88	0.05	MAN 1	1	99		
Megasphaera elsdenii		3.93	0.07		99	1		
01302021	**Good identification**							**01302021**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01302030	**Good identification**							**01302030**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01302031	**Good identification**							**01302031**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01302040	**Low discrimination**							**01302040**
					COC	GRA		
Megasphaera elsdenii		49.30	0.38	SAL 1	99	1		
Clostridium novyi biovar A		44.50	0.45	FRU 95	1	99		
Clostridium botulinum biovar A		4.99	0.17		1	99		
01302041	**Good identification**							**01302041**
					COC	GRA		
Clostridium novyi biovar A		98.89	0.45	SAL 8	1	99		
Megasphaera elsdenii		1.11	0.04	MAN 1	99	1		
01302050	**Good identification**							**01302050**
					COC	GRA		
Clostridium novyi biovar A		98.89	0.45	SAL 8	1	99		
Megasphaera elsdenii		1.11	0.04	MAN 1	99	1		
01302051	**Good identification**							**01302051**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01304000	**Low discrimination**							**01304000**
					COC	GRA	SPO	
Clostridium novyi biovar A		31.29	0.48	SAL 8	1	99	99	
Peptostreptococcus anaerobius		22.15	0.29	XYL 1	99	99	1	
Prevotella disiens		12.21	0.23		1	1	1	
01304001	**Good identification**							**01304001**
					COC	SPO		
Clostridium novyi biovar A		98.20	0.48	SAL 8	1	99		
Clostridium sordellii		1.10	-.03	XYL 1	1	99		
Peptostreptococcus anaerobius		0.70	-.06		99	1		
01304010	**Good identification**							**01304010**
					COC	SPO		
Clostridium novyi biovar A		99.29	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		0.71	-.06	XYL 1	99	1		
01304011	**Good identification**							**01304011**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01304020	**Acceptable identification**							**01304020**
					COC	SPO		
Clostridium novyi biovar A		91.27	0.37	SAL 8	1	99		
Clostridium bifermentans		6.12	0.05	XYL 1	1	99		
Peptostreptococcus anaerobius		2.61	-.06		99	1		
01304021	**Good identification**							**01304021**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01304030	**Good identification**							**01304030**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01304031	**Good identification**							**01304031**
Clostridium novyi biovar A		100.0	0.37	SAL 8				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01304040	**Acceptable identification**							**01304040**
Clostridium glycolicum		90.40	0.50	SAL 1				
Clostridium novyi biovar A		8.35	0.45	FRU 90				
Clostridium botulinum biovar A		0.94	0.17					
01304041	**Acceptable identification**							**01304041**
Clostridium novyi biovar A		90.14	0.45	SAL 8				
Clostridium glycolicum		9.86	0.15	XYL 1				
01304050	**Acceptable identification**							**01304050**
Clostridium novyi biovar A		90.14	0.45	SAL 8				
Clostridium glycolicum		9.86	0.15	XYL 1				
01304051	**Good identification**							**01304051**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01310000	**Low discrimination**							**01310000**
					COC	GRA	SPO	
Prevotella disiens		47.29	0.57	SAL 1	1	1	1	
Megasphaera elsdenii		25.78	0.63		99	1	1	
Clostridium novyi biovar A		13.47	0.66		1	99	99	
01310001	**Low discrimination**							**01310001**
					GRA	SPO		
Clostridium novyi biovar A		85.88	0.66	SAL 8	99	99		
Clostridium sordellii		8.67	0.32	FRU 10	99	99		
Prevotella disiens		3.05	0.23		1	1		
01310002	**Low discrimination**							**01310002**
					COC	GRA		
Megasphaera elsdenii		39.12	0.47	SAL 1	99	1		
Clostridium novyi biovar A		32.47	0.53	BGL 10	1	99		
Clostridium sporogenes		13.33	0.32		1	99		
01310003	**Good identification**							**01310003**
					COC	GRA		
Clostridium novyi biovar A		97.84	0.53	SAL 8	1	99		
Megasphaera elsdenii		1.19	0.12	FRU 10	99	1		
Clostridium sordellii		0.57	-.03		1	99		
01310010	**Good identification**							**01310010**
					GRA	SPO		
Clostridium novyi biovar A		90.64	0.66	SAL 8	99	99		
Clostridium difficile		3.59	0.27	FRU 10	99	99		
Prevotella disiens		3.22	0.23		1	1		
01310011	**Very good identification**							**01310011**
Clostridium novyi biovar A		99.90	0.66	SAL 8				
Clostridium sordellii		0.10	-.03	FRU 10				
01310012	**Good identification**							**01310012**
					COC	GRA		
Clostridium novyi biovar A		98.39	0.53	SAL 8	1	99		
Megasphaera elsdenii		1.20	0.12	FRU 10	99	1		
Clostridium sporogenes		0.41	-.02		1	99		
01310013	**Good identification**							**01310013**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.53	SAL 8				
01310020	**Low discrimination**							**01310020**
					GRA	SPO		
Clostridium novyi biovar A		68.99	0.55	SAL 8	99	99		
Prevotella disiens		9.79	0.23	FRU 10	1	1		
Pseudoramibacter alactolyticus		5.96	0.19		99	1		
01310021	**Good identification**							**01310021**
Clostridium novyi biovar A		99.59	0.55	SAL 8				
Clostridium sordellii		0.41	-.03	FRU 10				
01310022	**Low discrimination**							**01310022**
					SPO			
Clostridium novyi biovar A		81.50	0.42	SAL 8	99			
Atopobium minutum		6.96	0.20	FRU 10	1			
Pseudoramibacter alactolyticus		4.43	0.03		1			
01310023	**Good identification**							**01310023**
Clostridium novyi biovar A		100.0	0.42	SAL 8				
01310030	**Good identification**							**01310030**
Clostridium novyi biovar A		100.0	0.55	SAL 8				
01310031	**Good identification**							**01310031**
Clostridium novyi biovar A		100.0	0.55	SAL 8				
01310032	**Good identification**							**01310032**
Clostridium novyi biovar A		100.0	0.42	SAL 8				
01310033	**Good identification**							**01310033**
Clostridium novyi biovar A		100.0	0.42	SAL 8				
01310040	**Low discrimination**							**01310040**
					COC	GRA		
Megasphaera elsdenii		53.71	0.60	SAL 1	99	1		
Clostridium novyi biovar A		28.07	0.63		1	99		
Clostridium glycolicum		13.09	0.44		1	99		
01310041	**Very good identification**							**01310041**
					COC	GRA		
Clostridium novyi biovar A		97.05	0.63	SAL 8	1	99		
Megasphaera elsdenii		1.88	0.26	FRU 10	99	1		
Clostridium sordellii		0.61	0.08		1	99		
01310042	**Low discrimination**							**01310042**
					COC	SPO	GRA	
Megasphaera elsdenii		53.16	0.44	SAL 1	99	1	1	
Clostridium novyi biovar A		44.12	0.50	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		1.28	0.05		1	1	99	
01310043	**Good identification**							**01310043**
					COC	GRA		
Clostridium novyi biovar A		98.80	0.50	SAL 8	1	99		
Megasphaera elsdenii		1.20	0.09	FRU 10	99	1		
01310050	**Very good identification**							**01310050**
					COC	GRA		
Clostridium novyi biovar A		97.16	0.63	SAL 8	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		1.88	0.26	FRU 10	99	1		
Clostridium glycolicum		0.46	0.10		1	99		
01310051	**Very good identification**							**01310051**
Clostridium novyi biovar A		100.0	0.63	SAL 8				
01310052	**Good identification**							**01310052**
					COC	GRA		
Clostridium novyi biovar A		98.80	0.50	SAL 8	1	99		
Megasphaera elsdenii		1.20	0.09	FRU 10	99	1		
01310053	**Good identification**							**01310053**
Clostridium novyi biovar A		100.0	0.50	SAL 8				
01310060	**Low discrimination**							**01310060**
					COC	SPO	GRA	
Clostridium novyi biovar A		87.96	0.52	SAL 8	1	99	99	
Megasphaera elsdenii		6.80	0.26	FRU 10	99	1	1	
Pseudoramibacter alactolyticus		2.85	0.09		1	1	99	
01310061	**Good identification**							**01310061**
Clostridium novyi biovar A		100.0	0.52	SAL 8				
01310062	**Acceptable identification**							**01310062**
					COC	SPO	GRA	
Clostridium novyi biovar A		93.54	0.39	SAL 8	1	99	99	
Megasphaera elsdenii		4.55	0.09	FRU 10	99	1	1	
Pseudoramibacter alactolyticus		1.91	-.08		1	1	99	
01310063	**Good identification**							**01310063**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01310070	**Good identification**							**01310070**
Clostridium novyi biovar A		99.60	0.52	SAL 8				
Clostridium botulinum biovar B		0.41	-.03	FRU 10				
01310071	**Good identification**							**01310071**
Clostridium novyi biovar A		100.0	0.52	SAL 8				
01310072	**Good identification**							**01310072**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01310073	**Good identification**							**01310073**
Clostridium novyi biovar A		100.0	0.39	SAL 8				
01310100	**Low discrimination**							**01310100**
					GRA			
Fusobacterium mortiferum		58.93	0.56	MLT 11	1			
Actinomyces meyeri		20.85	0.39		99			
Prevotella disiens		9.38	0.23		1			
01310102	**Low discrimination**							**01310102**
					COC	GRA		
Fusobacterium mortiferum		71.98	0.39	MLT 11	1	1		
Actinomyces meyeri		7.09	0.13	BGL 10	1	99		
Megasphaera elsdenii		6.25	0.12		99	1		
01311000	**Low discrimination**							**01311000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		27.23	0.52	MLT 11	1	1	
Clostridium novyi biovar A		21.12	0.49		99	99	
Prevotella intermedia		19.24	0.39		1	1	
01311001	**Low discrimination**						**01311001**
					GRA	SPO	
Clostridium novyi biovar A		84.57	0.49	SAL 8	99	99	
Clostridium sordellii		13.55	0.19	FRU 10	99	99	
Fusobacterium mortiferum		1.10	0.18	GAL 10	1	1	
01311002	**Low discrimination**						**01311002**
					GRA	SPO	
Lactobacillus fermentum		43.26	0.44	SUC 87	99	1	
Clostridium novyi biovar A		21.23	0.36	SAL 1	99	99	
Fusobacterium mortiferum		17.23	0.36		1	1	
01311003	**Good identification**						**01311003**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01311010	**Good identification**						**01311010**
					GRA	SPO	
Clostridium novyi biovar A		95.99	0.49	SAL 8	99	99	
Lactobacillus jensenii		1.87	0.11	FRU 10	99	1	
Fusobacterium mortiferum		1.25	0.18	GAL 10	1	1	
01311011	**Good identification**						**01311011**
Clostridium novyi biovar A		100.0	0.49	SAL 8			
01311012	**Acceptable identification**						**01311012**
					SPO		
Clostridium novyi biovar A		93.13	0.36	SAL 8	99		
Lactobacillus jensenii		6.87	0.08	FRU 10	1		
01311013	**Good identification**						**01311013**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01311020	**Low discrimination**						**01311020**
					SPO		
Clostridium novyi biovar A		42.64	0.39	SAL 8	99		
Lactobacillus fermentum		20.45	0.36	FRU 10	1		
Clostridium perfringens		10.31	0.19	GAL 10	84		
01311021	**Good identification**						**01311021**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
01311022	**Low discrimination**						**01311022**
					SPO		
Lactobacillus fermentum		54.67	0.36	SUC 87	1		
Clostridium novyi biovar A		20.12	0.26	SAL 1	99		
Clostridium perfringens		12.97	0.13		84		
01311030	**Good identification**						**01311030**
					SPO		
Clostridium novyi biovar A		98.09	0.39	SAL 8	99		
Lactobacillus jensenii		1.91	0.01	FRU 10	1		
01311031	**Good identification**						**01311031**
Clostridium novyi biovar A		100.0	0.39	SAL 8			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01311040	**Low discrimination**						**01311040**
					COC	GRA	
Clostridium novyi biovar A		59.75	0.46	SAL 8	1	99	
Clostridium glycolicum		27.87	0.28	FRU 10	1	99	
Megasphaera elsdenii		10.39	0.26	GAL 10	99	1	
01311041	**Good identification**						**01311041**
Clostridium novyi biovar A		99.01	0.46	SAL 8			
Clostridium sordellii		0.99	-.05	FRU 10			
01311050	**Good identification**						**01311050**
Clostridium novyi biovar A		100.0	0.46	SAL 8			
01311051	**Good identification**						**01311051**
Clostridium novyi biovar A		100.0	0.46	SAL 8			
01311060	**Acceptable identification**						**01311060**
Clostridium novyi biovar A		93.54	0.36	SAL 8			
Clostridium perfringens		6.46	0.07	FRU 10			
01311061	**Good identification**						**01311061**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01311070	**Good identification**						**01311070**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01311071	**Good identification**						**01311071**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
01311100	**Low discrimination**						**01311100**
					GRA		
Fusobacterium mortiferum		63.24	0.59	MLT 11	1		
Actinomyces meyeri		22.37	0.42		99		
Lactobacillus fermentum		12.34	0.48		99		
01311102	**Low discrimination**						**01311102**
					GRA		
Lactobacillus fermentum		60.40	0.48	SUC 87	99		
Fusobacterium mortiferum		34.39	0.42	SAL 1	1		
Actinomyces meyeri		3.39	0.16		99		
01311120	**Low discrimination**						**01311120**
					SPO		
Clostridium septicum		81.09	0.43	MNS 99	99		
Clostridium perfringens		9.68	0.34		84		
Lactobacillus fermentum		4.27	0.40		1		
01311400	**Low discrimination**						**01311400**
					COC		
Lactobacillus fermentum		75.34	0.44	SUC 87	1		
Actinomyces meyeri		13.93	0.21	SAL 1	1		
Gemella morbillorum		3.96	0.11		99		
01311402	**Good identification**						**01311402**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01311420	**Acceptable identification**						**01311420**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		92.97	0.36	SUC 87				
Actinomyces meyeri		7.03	0.06	SAL 1				
01311422	**Good identification**							**01311422**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01311500	**Low discrimination**							**01311500**
					GRA			
Lactobacillus fermentum		66.44	0.48	SUC 87	99			
Actinomyces meyeri		30.12	0.31	SAL 1	99			
Fusobacterium mortiferum		3.44	0.25		1			
01311502	**Good identification**							**01311502**
Lactobacillus fermentum		98.62	0.48	SUC 87				
Actinomyces meyeri		1.38	0.06	SAL 1				
01311520	**Low discrimination**							**01311520**
					SPO			
Lactobacillus fermentum		72.62	0.40	SUC 87	1			
Clostridium septicum		13.92	0.09	SAL 1	99			
Actinomyces meyeri		13.47	0.17		1			
01311522	**Good identification**							**01311522**
Lactobacillus fermentum		100.0	0.40	SUC 87				
01312000	**Low discrimination**							**01312000**
					COC	SPO	GRA	
Megasphaera elsdenii		89.10	0.63	SAL 1	99	1	1	
Clostridium difficile		3.40	0.32		1	99	99	
Pseudoramibacter alactolyticus		3.07	0.27		1	1	99	
01312002	**Good identification**							**01312002**
					COC	SPO	GRA	
Megasphaera elsdenii		95.26	0.47	SAL 1	99	1	1	
Pseudoramibacter alactolyticus		3.28	0.11	BGL 10	1	1	99	
Clostridium novyi biovar A		0.80	0.19		1	99	99	
01312010	**Acceptable identification**							**01312010**
					COC	GRA		
Clostridium difficile		92.22	0.43	MLT 1	1	99		
Megasphaera elsdenii		5.00	0.29		99	1		
Clostridium novyi biovar A		2.61	0.31		1	99		
01312040	**Very good identification**							**01312040**
					COC	SPO	GRA	
Megasphaera elsdenii		97.69	0.60	SAL 1	99	1	1	
Pseudoramibacter alactolyticus		1.26	0.17		1	1	99	
Clostridium novyi biovar A		0.52	0.28		1	99	99	
01312042	**Good identification**							**01312042**
					COC	SPO	GRA	
Megasphaera elsdenii		97.91	0.44	SAL 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.00	BGL 10	1	1	99	
Clostridium novyi biovar A		0.82	0.15		1	99	99	
01314000	**Low discrimination**							**01314000**
					GRA	SPO		
Clostridium glycolicum		70.92	0.44	SAL 1	99	99		
Actinomyces meyeri		8.72	0.33	SOR 95	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella disiens		8.09	0.23		1	1	
01314040	**Very good identification**						**01314040**
					COC	GRA	
Clostridium glycolicum		99.67	0.66	SAL 1	1	99	
Megasphaera elsdenii		0.22	0.26		99	1	
Clostridium novyi biovar A		0.11	0.28		1	99	
01314100	**Low discrimination**						**01314100**
					GRA	SPO	
Actinomyces meyeri		87.72	0.44	SUC 93	99	1	
Fusobacterium mortiferum		9.62	0.36	SAL 1	1	1	
Clostridium glycolicum		1.80	0.10		99	99	
01315000	**Low discrimination**						**01315000**
					SPO		
Actinomyces meyeri		42.94	0.36	SUC 93	1		
Clostridium glycolicum		25.86	0.28	SAL 1	99		
Lactobacillus fermentum		17.57	0.41		1		
01315002	**Low discrimination**						**01315002**
Lactobacillus fermentum		82.66	0.41	SUC 87			
Actinomyces odontolyticus		6.88	0.15	SAL 1			
Actinomyces meyeri		6.25	0.11				
01315040	**Good identification**						**01315040**
					SPO		
Clostridium glycolicum		99.80	0.50	SAL 1	99		
Clostridium novyi biovar A		0.11	0.12	GAL 10	99		
Actinomyces meyeri		0.09	0.02		1		
01315100	**Low discrimination**						**01315100**
					GRA		
Actinomyces meyeri		78.07	0.47	SUC 93	99		
Lactobacillus fermentum		13.03	0.44	SAL 1	99		
Fusobacterium mortiferum		8.56	0.40		1		
01315102	**Low discrimination**						**01315102**
					GRA		
Lactobacillus fermentum		78.47	0.44	SUC 87	99		
Actinomyces meyeri		14.54	0.21	SAL 1	99		
Fusobacterium mortiferum		5.73	0.23		1		
01315122	**Acceptable identification**						**01315122**
Lactobacillus fermentum		92.95	0.36	SUC 87			
Actinomyces meyeri		7.05	0.06	SAL 1			
01315400	**Low discrimination**						**01315400**
Lactobacillus fermentum		62.07	0.41	SUC 87			
Actinomyces meyeri		37.93	0.26	SAL 1			
01315402	**Good identification**						**01315402**
Lactobacillus fermentum		100.0	0.41	SUC 87			
01315500	**Low discrimination**						**01315500**
Actinomyces meyeri		59.97	0.36	SUC 93			
Lactobacillus fermentum		40.03	0.44	SAL 1			
01315502	**Acceptable identification**						**01315502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		91.35	0.44	SUC 87		
Bifidobacterium adolescentis		4.41	0.16	SAL 1		
Actinomyces meyeri		4.23	0.11			
01315520	**Low discrimination**					**01315520**
Lactobacillus fermentum		62.00	0.36	SUC 87		
Actinomyces meyeri		38.00	0.22	SAL 1		
01315522	**Good identification**					**01315522**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01320000	**Low discrimination**					**01320000**
Clostridium botulinum biovar A		47.34	0.49	SAL 1		
Clostridium sporogenes		34.53	0.45			
Clostridium novyi biovar A		7.65	0.48			
01320001	**Low discrimination**					**01320001**
Clostridium novyi biovar A		88.77	0.48	SAL 8		
Clostridium botulinum biovar A		5.55	0.15	TRE 1		
Clostridium sporogenes		4.05	0.11			
01320002	**Low discrimination**					**01320002**
Clostridium sporogenes		85.95	0.37	SAL 1		
Clostridium novyi biovar A		10.07	0.35			
Clostridium botulinum biovar A		3.57	0.15			
01320010	**Low discrimination**					**01320010**
Clostridium novyi biovar A		89.66	0.48	SAL 8		
Clostridium botulinum biovar A		5.61	0.15	TRE 1		
Clostridium sporogenes		4.09	0.11			
01320011	**Good identification**					**01320011**
Clostridium novyi biovar A		100.0	0.48	SAL 8		
01320020	**Low discrimination**					**01320020**
Clostridium novyi biovar A		65.43	0.37	SAL 8		
Clostridium botulinum biovar A		16.37	0.15	TRE 1		
Clostridium sporogenes		11.94	0.11			
01320021	**Good identification**					**01320021**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
01320030	**Good identification**					**01320030**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
01320031	**Good identification**					**01320031**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
01320040	**Low discrimination**					**01320040**
Clostridium botulinum biovar A		70.85	0.41	SAL 1		
Clostridium novyi biovar A		25.54	0.45			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sporogenes		1.75	0.11					
01320041	**Good identification**							**01320041**
Clostridium novyi biovar A		97.27	0.45	SAL 8				
Clostridium botulinum biovar A		2.73	0.06	TRE 1				
01320050	**Good identification**							**01320050**
Clostridium novyi biovar A		95.69	0.45	SAL 8				
Clostridium botulinum biovar A		2.68	0.06	TRE 1				
Clostridium botulinum biovar B		1.63	0.00					
01320051	**Good identification**							**01320051**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01330100	**Good identification**							**01330100**
					GRA	SPO		
Fusobacterium mortiferum		97.76	0.47	MLT 11	1	1		
Actinomyces meyeri		1.17	0.05		99	1		
Clostridium sporogenes		1.08	-.02		99	99		
01331000	**Low discrimination**							**01331000**
					GRA	SPO		
Fusobacterium mortiferum		65.64	0.43	MLT 11	1	1		
Lactobacillus jensenii		29.95	0.28		99	1		
Clostridium novyi biovar A		1.72	0.15		99	99		
01331100	**Low discrimination**							**01331100**
					COC	GRA	SPO	
Fusobacterium mortiferum		75.18	0.50	MLT 11	1	1	1	
Atopobium parvulum		11.45	0.24		99	99	1	
Clostridium septicum		9.41	0.17		1	99	99	
01331120	**Acceptable identification**							**01331120**
					COC	SPO		
Clostridium septicum		92.82	0.51	MNS 99	1	99		
Clostridium perfringens		6.86	0.38		1	84		
Atopobium parvulum		0.20	0.11		99	1		
01340000	**Low discrimination**							**01340000**
					COC	GRA	SPO	
Clostridium novyi biovar A		36.25	0.48	SAL 8	1	99	99	
Peptostreptococcus anaerobius		25.66	0.29	CEL 1	99	99	1	
Prevotella disiens		14.14	0.23		1	1	1	
01340001	**Good identification**							**01340001**
					COC	SPO		
Clostridium novyi biovar A		98.20	0.48	SAL 8	1	99		
Clostridium sordellii		1.10	-.03	CEL 1	1	99		
Peptostreptococcus anaerobius		0.70	-.06		99	1		
01340010	**Good identification**							**01340010**
					COC	SPO		
Clostridium novyi biovar A		99.29	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		0.71	-.06	CEL 1	99	1		
01340011	**Good identification**							**01340011**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01340020	**Acceptable identification**							**01340020**
					COC	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		91.27	0.37	SAL 8	1	99	
Clostridium bifermentans		6.12	0.05	CEL 1	1	99	
Peptostreptococcus anaerobius		2.61	-.06		99	1	
01340021	**Good identification**						**01340021**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01340030	**Good identification**						**01340030**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01340031	**Good identification**						**01340031**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01340040	**Low discrimination**						**01340040**
Clostridium novyi biovar A		86.92	0.45	SAL 8			
Clostridium botulinum biovar A		9.74	0.17	CEL 1			
Clostridium bifermentans		1.43	0.02				
01340041	**Good identification**						**01340041**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01340050	**Good identification**						**01340050**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01340051	**Good identification**						**01340051**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01350100	**Good identification**						**01350100**
					GRA		
Fusobacterium mortiferum		98.94	0.48	MLT 11	1		
Actinomyces meyeri		1.06	0.05		99		
01351000	**Low discrimination**						**01351000**
					GRA		
Lactobacillus jensenii		63.11	0.41	SUC 90	99		
Fusobacterium mortiferum		24.37	0.44	MNS 99	1		
Lactobacillus catenaformis		11.43	0.29		99		
01351002	**Low discrimination**						**01351002**
					GRA		
Lactobacillus jensenii		73.24	0.38	SUC 90	99		
Lactobacillus catenaformis		19.89	0.29	MNS 99	99		
Fusobacterium mortiferum		4.71	0.28		1		
01351100	**Low discrimination**						**01351100**
					GRA	SPO	
Fusobacterium mortiferum		67.57	0.51	MLT 11	1	1	
Lactobacillus catenaformis		20.37	0.32		99	1	
Clostridium septicum		8.00	0.17		99	99	
01351120	**Good identification**						**01351120**
					SPO		
Clostridium septicum		97.99	0.51	MNS 99	99		
Clostridium paraputrificum		0.89	0.14		99		
Lactobacillus catenaformis		0.63	0.21		1		
01357100	**Good identification**						**01357100**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.37	SUC 98			
01357102	**Good identification**						**01357102**
Actinomyces israelii		100.0	0.40	SUC 98			
01357500	**Good identification**						**01357500**
Actinomyces israelii		100.0	0.36	SUC 98			
01357502	**Low discrimination**						**01357502**
Actinomyces israelii		83.53	0.39	SUC 98			
Bifidobacterium adolescentis		16.47	0.20				
01370100	**Good identification**						**01370100**
Fusobacterium mortiferum		100.0	0.39	MLT 11			
01371000	**Good identification**						**01371000**
					GRA		
Lactobacillus jensenii		96.30	0.50	SUC 90	99		
Fusobacterium mortiferum		3.70	0.35	MNS 99	1		
01371002	**Good identification**						**01371002**
					GRA		
Lactobacillus jensenii		97.66	0.47	SUC 90	99		
Lactobacillus acidophilus		1.71	0.14	MNS 99	99		
Fusobacterium mortiferum		0.63	0.19		1		
01371010	**Good identification**						**01371010**
Lactobacillus jensenii		100.0	0.41	SUC 90			
01371012	**Good identification**						**01371012**
Lactobacillus jensenii		100.0	0.38	SUC 90			
01371020	**Low discrimination**						**01371020**
					SPO		
Lactobacillus jensenii		70.09	0.39	SUC 90	1		
Clostridium septicum		29.92	0.25	MNS 99	99		
01371022	**Acceptable identification**						**01371022**
					SPO		
Lactobacillus jensenii		92.32	0.36	SUC 90	1		
Lactobacillus acidophilus		6.47	0.14	MNS 99	1		
Clostridium septicum		1.21	-.04		99		
01371120	**Very good identification**						**01371120**
					COC	SPO	
Clostridium septicum		99.62	0.60	MNS 99	1	99	
Atopobium parvulum		0.16	0.18		99	1	
Clostridium perfringens		0.10	0.15		1	84	
01375100	**Low discrimination**						**01375100**
					COC	GRA	
Atopobium parvulum		85.33	0.35	SUC 96	99	99	
Actinomyces israelii		11.43	0.34	MNS 99	1	99	
Fusobacterium mortiferum		3.24	0.22		1	1	
01375102	**Low discrimination**						**01375102**
					COC		
Actinomyces israelii		53.24	0.37	SUC 98	1		
Atopobium parvulum		46.76	0.22		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01375502	**Low discrimination**						**01375502**
Actinomyces israelii		80.52	0.36	SUC 98			
Bifidobacterium adolescentis		19.48	0.19				
01377000	**Good identification**						**01377000**
Actinomyces israelii		100.0	0.37	SUC 98			
01377002	**Good identification**						**01377002**
Actinomyces israelii		100.0	0.40	SUC 98			
01377100	**Good identification**						**01377100**
					COC	SPO	
Actinomyces israelii		96.92	0.47	SUC 98	1	1	
Clostridium tertium		1.79	0.03		1	99	
Atopobium parvulum		1.29	0.01		99	1	
01377102	**Good identification**						**01377102**
					SPO		
Actinomyces israelii		96.44	0.50	SUC 98	1		
Clostridium tertium		3.56	0.11		99		
01377112	**Low discrimination**						**01377112**
					SPO		
Actinomyces israelii		69.51	0.36	SUC 98	1		
Clostridium tertium		30.49	0.15	MLZ 13	99		
01377140	**Good identification**						**01377140**
Actinomyces israelii		100.0	0.40	SUC 98			
01377142	**Good identification**						**01377142**
Actinomyces israelii		100.0	0.43	SUC 98			
01377400	**Good identification**						**01377400**
Actinomyces israelii		100.0	0.35	SUC 98			
01377402	**Good identification**						**01377402**
Actinomyces israelii		100.0	0.38	SUC 98			
01377500	**Good identification**						**01377500**
Actinomyces israelii		100.0	0.46	SUC 98			
01377502	**Good identification**						**01377502**
Actinomyces israelii		96.63	0.49	SUC 98			
Bifidobacterium adolescentis		3.38	0.17				
01377540	**Good identification**						**01377540**
Actinomyces israelii		100.0	0.38	SUC 98			
01377542	**Good identification**						**01377542**
Actinomyces israelii		100.0	0.41	SUC 98			
01400000	**Low discrimination**						**01400000**
					COC	SPO	
Peptostreptococcus anaerobius		51.74	0.66	RAF 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		14.90	0.61		1	99		
Atopobium minutum		9.85	0.66		1	1		
01400001	**Low discrimination**							**01400001**
					COC	GRA	SPO	
Anaerococcus prevotii		46.72	0.50	GLU 1	99	99	1	
Clostridium novyi biovar A		38.75	0.66		1	99	99	
Campylobacter ureolyticus		7.00	0.32		1	1	1	
01400002	**Low discrimination**							**01400002**
					SPO			
Atopobium minutum		63.51	0.63	RAF 1	1			
Clostridium sporogenes		11.58	0.40		99			
Clostridium novyi biovar A		9.66	0.53		99			
01400003	**Low discrimination**							**01400003**
					COC	SPO		
Anaerococcus prevotii		60.53	0.39	GLU 1	99	1		
Clostridium novyi biovar A		35.44	0.53		1	99		
Atopobium minutum		2.35	0.29		1	1		
01400010	**Low discrimination**							**01400010**
					COC	SPO		
Clostridium novyi biovar A		85.56	0.66	RAF 1	1	99		
Peptostreptococcus anaerobius		7.90	0.32		99	1		
Clostridium botulinum biovar A		2.28	0.27		1	99		
01400011	**Very good identification**							**01400011**
					COC	GRA	SPO	
Clostridium novyi biovar A		98.53	0.66	RAF 1	1	99	99	
Anaerococcus prevotii		1.20	0.15		99	99	1	
Campylobacter ureolyticus		0.18	-.03		1	1	1	
01400012	**Acceptable identification**							**01400012**
					SPO			
Clostridium novyi biovar A		92.27	0.53	RAF 1	99			
Atopobium minutum		6.13	0.29		1			
Clostridium sporogenes		1.12	0.06		99			
01400013	**Good identification**							**01400013**
					COC	SPO		
Clostridium novyi biovar A		98.30	0.53	RAF 1	1	99		
Anaerococcus prevotii		1.70	0.05		99	1		
01400020	**Low discrimination**							**01400020**
					SPO			
Atopobium minutum		49.57	0.58	RAF 1	1			
Clostridium novyi biovar A		21.37	0.56		99			
Clostridium cadaveris		12.28	0.37		99			
01400021	**Acceptable identification**							**01400021**
					COC	SPO		
Clostridium novyi biovar A		91.10	0.56	RAF 1	1	99		
Anaerococcus prevotii		4.44	0.15		99	1		
Atopobium minutum		2.14	0.24		1	1		
01400022	**Low discrimination**							**01400022**
					GRA	SPO		
Atopobium minutum		69.11	0.55	RAF 1	99	1		
Pseudoflavonifractor capillosus		21.10	0.32		1	1		
Clostridium novyi biovar A		7.89	0.43		99	99		
01400023	**Low discrimination**							**01400023**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				COC	SPO	
Clostridium novyi biovar A	84.42	0.43	RAF 1	1	99	
Atopobium minutum	7.47	0.21		1	1	
Anaerococcus prevotii	5.83	0.05		99	1	
01400030 Good identification						**01400030**
				SPO		
Clostridium novyi biovar A	96.81	0.56	RAF 1	99		
Atopobium minutum	2.27	0.24		1		
Clostridium cadaveris	0.56	0.03		99		
01400031 Good identification						**01400031**
Clostridium novyi biovar A	100.0	0.56	RAF 1			
01400032 Low discrimination						**01400032**
				GRA	SPO	
Clostridium novyi biovar A	89.64	0.43	RAF 1	99	99	
Atopobium minutum	7.94	0.21		99	1	
Pseudoflavonifractor capillosus	2.42	-.03		1	1	
01400033 Good identification						**01400033**
Clostridium novyi biovar A	100.0	0.43	RAF 1			
01400040 Low discrimination						**01400040**
				COC	SPO	
Clostridium botulinum biovar A	46.33	0.52	RAF 1	1	99	
Clostridium novyi biovar A	39.28	0.63		1	99	
Peptostreptococcus anaerobius	5.44	0.32		99	1	
01400041 Very good identification						**01400041**
				COC	SPO	
Clostridium novyi biovar A	96.68	0.63	RAF 1	1	99	
Anaerococcus prevotii	1.77	0.15		99	1	
Clostridium botulinum biovar A	1.15	0.18		1	99	
01400042 Low discrimination						**01400042**
				SPO		
Clostridium novyi biovar A	79.28	0.50	RAF 1	99		
Atopobium minutum	7.90	0.29		1		
Clostridium botulinum biovar A	5.35	0.18		99		
01400043 Good identification						**01400043**
				COC	SPO	
Clostridium novyi biovar A	97.48	0.50	RAF 1	1	99	
Anaerococcus prevotii	2.52	0.05		99	1	
01400050 Very good identification						**01400050**
				COC	SPO	
Clostridium novyi biovar A	98.69	0.63	RAF 1	1	99	
Clostridium botulinum biovar A	1.18	0.18		1	99	
Peptostreptococcus anaerobius	0.14	-.03		99	1	
01400051 Very good identification						**01400051**
Clostridium novyi biovar A	100.0	0.63	RAF 1			
01400052 Good identification						**01400052**
Clostridium novyi biovar A	100.0	0.50	RAF 1			
01400053 Good identification						**01400053**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.50	RAF 1			
01400060	**Low discrimination**						**01400060**
					SPO		
Clostridium novyi biovar A		88.64	0.53	RAF 1	99		
Clostridium botulinum biovar A		4.22	0.18		99		
Atopobium minutum		3.12	0.24		1		
01400061	**Good identification**						**01400061**
Clostridium novyi biovar A		100.0	0.53	RAF 1			
01400062	**Low discrimination**						**01400062**
					GRA	SPO	
Clostridium novyi biovar A		83.83	0.40	RAF 1	99	99	
Atopobium minutum		11.13	0.21		99	1	
Pseudoflavonifractor capillosus		3.40	-.03		1	1	
01400063	**Good identification**						**01400063**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
01400070	**Good identification**						**01400070**
Clostridium novyi biovar A		100.0	0.53	RAF 1			
01400071	**Good identification**						**01400071**
Clostridium novyi biovar A		100.0	0.53	RAF 1			
01400072	**Good identification**						**01400072**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
01400073	**Good identification**						**01400073**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
01401000	**Low discrimination**						**01401000**
					COC	SPO	
Atopobium minutum		59.02	0.58	RAF 1	1	1	
Clostridium novyi biovar A		11.31	0.50		1	99	
Peptostreptococcus anaerobius		9.40	0.32		99	1	
01401001	**Low discrimination**						**01401001**
					COC	SPO	
Clostridium novyi biovar A		82.50	0.50	RAF 1	1	99	
Anaerococcus prevotii		9.04	0.15	GAL 10	99	1	
Atopobium minutum		4.35	0.24		1	1	
01401002	**Low discrimination**						**01401002**
					SPO		
Atopobium minutum		87.67	0.55	RAF 1	1		
Lactobacillus fermentum		4.71	0.40		1		
Clostridium novyi biovar A		4.45	0.37		99		
01401003	**Low discrimination**						**01401003**
					COC	SPO	
Clostridium novyi biovar A		73.83	0.37	RAF 1	1	99	
Atopobium minutum		14.71	0.21	GAL 10	1	1	
Anaerococcus prevotii		11.46	0.05		99	1	
01401010	**Acceptable identification**						**01401010**
					COC	SPO	
Clostridium novyi biovar A		93.54	0.50	RAF 1	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		4.93	0.24	GAL 10	1	1	
Peptostreptococcus anaerobius		0.79	-.03		99	1	
01401011	**Good identification**						**01401011**
Clostridium novyi biovar A		100.0	0.50	RAF 1			
01401012	**Low discrimination**						**01401012**
					SPO		
Clostridium novyi biovar A		83.39	0.37	RAF 1	99		
Atopobium minutum		16.61	0.21	GAL 10	1		
01401013	**Good identification**						**01401013**
Clostridium novyi biovar A		100.0	0.37	RAF 1			
01401020	**Low discrimination**						**01401020**
					COC	SPO	
Atopobium minutum		80.52	0.50	RAF 1	1	1	
Clostridium novyi biovar A		11.57	0.39		1	99	
Blautia hansenii		4.03	0.15		99	1	
01401021	**Acceptable identification**						**01401021**
					SPO		
Clostridium novyi biovar A		93.43	0.39	RAF 1	99		
Atopobium minutum		6.57	0.15	GAL 10	1		
01401022	**Low discrimination**						**01401022**
					GRA		
Atopobium minutum		83.20	0.47	RAF 1	99		
Pseudoflavonifractor capillosus		8.47	0.15		1		
Lactobacillus fermentum		4.47	0.32		99		
01401030	**Acceptable identification**						**01401030**
					SPO		
Clostridium novyi biovar A		93.43	0.39	RAF 1	99		
Atopobium minutum		6.57	0.15	GAL 10	1		
01401031	**Good identification**						**01401031**
Clostridium novyi biovar A		100.0	0.39	RAF 1			
01401040	**Low discrimination**						**01401040**
					SPO		
Clostridium novyi biovar A		79.58	0.47	RAF 1	99		
Clostridium botulinum biovar A		8.53	0.18	GAL 10	99		
Atopobium minutum		6.29	0.24		1		
01401041	**Good identification**						**01401041**
Clostridium novyi biovar A		100.0	0.47	RAF 1			
01401050	**Good identification**						**01401050**
Clostridium novyi biovar A		100.0	0.47	RAF 1			
01401051	**Good identification**						**01401051**
Clostridium novyi biovar A		100.0	0.47	RAF 1			
01401060	**Acceptable identification**						**01401060**
					SPO		
Clostridium novyi biovar A		90.46	0.36	RAF 1	99		
Atopobium minutum		9.54	0.15	GAL 10	1		
01401061	**Good identification**						**01401061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.36	RAF 1			
01401070	**Good identification**						**01401070**
Clostridium novyi biovar A		100.0	0.36	RAF 1			
01401071	**Good identification**						**01401071**
Clostridium novyi biovar A		100.0	0.36	RAF 1			
01401100	**Very good identification**						**01401100**
					COC	GRA	
Blautia hansenii		99.16	0.66	MLT 99	99	99	
Lactobacillus fermentum		0.39	0.44		1	99	
Fusobacterium mortiferum		0.30	0.40		1	1	
01401102	**Good identification**						**01401102**
					COC		
Blautia hansenii		95.92	0.50	MLT 99	99		
Lactobacillus fermentum		3.40	0.44	BGL 10	1		
Atopobium minutum		0.39	0.21		1		
01401120	**Good identification**						**01401120**
					COC		
Blautia hansenii		98.45	0.50	MLT 99	99		
Lactobacillus fermentum		1.16	0.35	NAG 10	1		
Atopobium minutum		0.20	0.15		1		
01401400	**Low discrimination**						**01401400**
					COC		
Lactobacillus fermentum		47.19	0.40	SUC 87	1		
Blautia hansenii		34.85	0.19	FRU 99	99		
Atopobium minutum		13.30	0.24		1		
01401402	**Low discrimination**						**01401402**
					COC		
Lactobacillus fermentum		78.74	0.40	SUC 87	1		
Atopobium minutum		14.79	0.21	FRU 99	1		
Blautia hansenii		6.46	0.03		99		
01401500	**Good identification**						**01401500**
					COC		
Blautia hansenii		97.73	0.53	MLT 99	99		
Lactobacillus fermentum		2.18	0.44		1		
Actinomyces meyeri		0.09	0.09		1		
01401502	**Low discrimination**						**01401502**
					COC		
Blautia hansenii		83.28	0.37	MLT 99	99		
Lactobacillus fermentum		16.72	0.44	BGL 10	1		
01401520	**Acceptable identification**						**01401520**
					COC		
Blautia hansenii		93.73	0.37	MLT 99	99		
Lactobacillus fermentum		6.27	0.35	NAG 10	1		
01404040	**Good identification**						**01404040**
Clostridium glycolicum		97.38	0.49	RAF 1			
Clostridium botulinum biovar A		1.33	0.18	FRU 90			
Clostridium novyi biovar A		1.13	0.29				
01405002	**Low discrimination**						**01405002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Lactobacillus fermentum		68.51	0.36	SUC 87	1		
Atopobium minutum		21.92	0.21	FRU 99	1		
Blautia hansenii		9.57	0.03		99		
01405100	**Good identification**						**01405100**
					COC		
Blautia hansenii		97.93	0.53	MLT 99	99		
Lactobacillus fermentum		1.28	0.40		1		
Actinomyces meyeri		0.66	0.24		1		
01405102	**Low discrimination**						**01405102**
					COC		
Blautia hansenii		89.45	0.37	MLT 99	99		
Lactobacillus fermentum		10.55	0.40	BGL 10	1		
01405120	**Good identification**						**01405120**
					COC		
Blautia hansenii		95.46	0.37	MLT 99	99		
Lactobacillus fermentum		3.75	0.31	NAG 10	1		
Actinomyces meyeri		0.79	0.09		1		
01405400	**Low discrimination**						**01405400**
					COC		
Lactobacillus fermentum		78.47	0.36	SUC 87	1		
Blautia hansenii		17.42	0.06	FRU 99	99		
Actinomyces meyeri		4.11	0.03		1		
01405402	**Good identification**						**01405402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01405500	**Acceptable identification**						**01405500**
					COC		
Blautia hansenii		92.27	0.40	MLT 99	99		
Lactobacillus fermentum		6.85	0.40		1		
Actinomyces meyeri		0.88	0.14		1		
01410000	**Low discrimination**						**01410000**
					GRA		
Pseudoramibacter alactolyticus		63.31	0.66	RAF 1	99		
Fusobacterium varium		8.89	0.51		1		
Atopobium minutum		8.77	0.65		99		
01410002	**Low discrimination**						**01410002**
					COC	GRA	
Pseudoramibacter alactolyticus		47.87	0.50	RAF 1	1	99	
Atopobium minutum		39.79	0.62	BGL 10	1	99	
Megasphaera elsdenii		5.04	0.44		99	1	
01410003	**Low discrimination**						**01410003**
					SPO		
Clostridium novyi biovar A		40.28	0.37	RAF 1	99		
Pseudoramibacter alactolyticus		26.73	0.15	FRU 10	1		
Atopobium minutum		22.22	0.28		1		
01410010	**Low discrimination**						**01410010**
					SPO		
Clostridium difficile		31.10	0.36	RAF 1	99		
Pseudoramibacter alactolyticus		27.08	0.32	MAN 90	1		
Clostridium novyi biovar A		25.70	0.50		99		
01410011	**Good identification**						**01410011**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Clostridium novyi biovar A		97.76	0.50	RAF 1	99			
Clostridium difficile		1.20	0.02	FRU 10	99			
Pseudoramibacter alactolyticus		1.04	-.03		1			
01410012	**Low discrimination**							**01410012**
					SPO			
Clostridium novyi biovar A		40.76	0.37	RAF 1	99			
Pseudoramibacter alactolyticus		27.05	0.15	FRU 10	1			
Atopobium minutum		22.48	0.28		1			
01410013	**Good identification**							**01410013**
Clostridium novyi biovar A		100.0	0.37	RAF 1				
01410020	**Low discrimination**							**01410020**
					SPO			
Pseudoramibacter alactolyticus		72.00	0.53	RAF 1	1			
Atopobium minutum		18.84	0.57		1			
Clostridium cadaveris		5.06	0.37		99			
01410021	**Low discrimination**							**01410021**
					SPO			
Clostridium novyi biovar A		50.23	0.39	RAF 1	99			
Pseudoramibacter alactolyticus		37.37	0.19	FRU 10	1			
Atopobium minutum		9.78	0.23		1			
01410022	**Low discrimination**							**01410022**
					GRA			
Atopobium minutum		58.05	0.54	RAF 1	99			
Pseudoramibacter alactolyticus		36.97	0.37		99			
Pseudoflavonifractor capillosus		2.13	0.15		1			
01410030	**Low discrimination**							**01410030**
					SPO			
Clostridium novyi biovar A		48.08	0.39	RAF 1	99			
Pseudoramibacter alactolyticus		35.77	0.19	FRU 10	1			
Atopobium minutum		9.36	0.23		1			
01410031	**Good identification**							**01410031**
Clostridium novyi biovar A		100.0	0.39	RAF 1				
01410040	**Low discrimination**							**01410040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		69.61	0.56	RAF 1	1	1	99	
Megasphaera elsdenii		19.54	0.57		99	1	1	
Clostridium glycolicum		6.60	0.43		1	99	99	
01410041	**Low discrimination**							**01410041**
					COC	SPO	GRA	
Clostridium novyi biovar A		64.50	0.47	RAF 1	1	99	99	
Pseudoramibacter alactolyticus		25.49	0.21	FRU 10	1	1	99	
Megasphaera elsdenii		7.16	0.23		99	1	1	
01410042	**Low discrimination**							**01410042**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		73.12	0.39	RAF 1	1	1	99	
Megasphaera elsdenii		20.52	0.41	BGL 10	99	1	1	
Clostridium novyi biovar A		2.97	0.34		1	99	99	
01410050	**Low discrimination**							**01410050**
					COC	SPO	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		60.75	0.47	RAF 1	1	99	99	
Pseudoramibacter alactolyticus		24.01	0.21	FRU 10	1	1	99	
Megasphaera elsdenii		6.74	0.23		99	1	1	
01410051	**Good identification**							**01410051**
Clostridium novyi biovar A		100.0	0.47	RAF 1				
01410060	**Acceptable identification**							**01410060**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		92.46	0.43	RAF 1	1	1	99	
Clostridium novyi biovar A		3.35	0.36		1	99	99	
Megasphaera elsdenii		1.49	0.23		99	1	1	
01410061	**Low discrimination**							**01410061**
					SPO			
Clostridium novyi biovar A		78.19	0.36	RAF 1	99			
Pseudoramibacter alactolyticus		21.81	0.09	FRU 10	1			
01410070	**Low discrimination**							**01410070**
					SPO			
Clostridium novyi biovar A		78.19	0.36	RAF 1	99			
Pseudoramibacter alactolyticus		21.81	0.09	FRU 10	1			
01410071	**Good identification**							**01410071**
Clostridium novyi biovar A		100.0	0.36	RAF 1				
01410100	**Low discrimination**							**01410100**
					GRA			
Fusobacterium mortiferum		86.13	0.71		1			
Pseudoramibacter alactolyticus		5.85	0.32		99			
Actinomyces meyeri		3.34	0.38		99			
01410101	**Low discrimination**							**01410101**
					GRA	SPO		
Fusobacterium mortiferum		85.39	0.37	URE 1	1	1		
Pseudoramibacter alactolyticus		5.80	-.03		99	1		
Clostridium novyi biovar A		5.50	0.15		99	99		
01410102	**Low discrimination**							**01410102**
					GRA			
Fusobacterium mortiferum		75.79	0.55	BGL 10	1			
Lactobacillus fermentum		13.42	0.44		99			
Pseudoramibacter alactolyticus		5.15	0.15		99			
01410110	**Low discrimination**							**01410110**
					GRA	SPO		
Fusobacterium mortiferum		80.06	0.37	MLZ 1	1	1		
Clostridium difficile		6.24	0.02		99	99		
Pseudoramibacter alactolyticus		5.44	-.03		99	1		
01410122	**Low discrimination**							**01410122**
Lactobacillus fermentum		59.06	0.35	SUC 87				
Atopobium minutum		18.83	0.20	GAL 99				
Pseudoramibacter alactolyticus		12.00	0.03					
01410300	**Acceptable identification**							**01410300**
					GRA			
Fusobacterium mortiferum		90.36	0.37	RHA 1	1			
Pseudoramibacter alactolyticus		6.14	-.03		99			
Actinomyces meyeri		3.50	0.04		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01410402	**Low discrimination**						**01410402**
Lactobacillus fermentum		44.41	0.40	SUC 87			
Pseudoramibacter alactolyticus		27.79	0.15	GAL 99			
Atopobium minutum		23.10	0.28				
01410500	**Low discrimination**						**01410500**
					GRA		
Lactobacillus fermentum		48.99	0.44	SUC 87	99		
Fusobacterium mortiferum		25.16	0.37	GAL 99	1		
Actinomyces meyeri		24.14	0.28		99		
01410502	**Acceptable identification**						**01410502**
					GRA		
Lactobacillus fermentum		93.26	0.44	SUC 87	99		
Fusobacterium mortiferum		5.32	0.21	GAL 99	1		
Actinomyces meyeri		1.42	0.02		99		
01410520	**Low discrimination**						**01410520**
Lactobacillus fermentum		83.22	0.35	SUC 87			
Actinomyces meyeri		16.78	0.13	GAL 99			
01410522	**Good identification**						**01410522**
Lactobacillus fermentum		100.0	0.35	SUC 87			
01411000	**Low discrimination**						**01411000**
					GRA		
Lactobacillus fermentum		50.44	0.74	SUC 87	99		
Fusobacterium mortiferum		27.17	0.68		1		
Atopobium minutum		13.12	0.57		99		
01411001	**Low discrimination**						**01411001**
					GRA	SPO	
Lactobacillus fermentum		38.30	0.40	SUC 87	99	1	
Clostridium novyi biovar A		22.74	0.33	URE 1	99	99	
Fusobacterium mortiferum		20.63	0.34		1	1	
01411002	**Low discrimination**						**01411002**
					GRA		
Lactobacillus fermentum		80.23	0.74	SUC 87	99		
Atopobium minutum		13.91	0.54		99		
Fusobacterium mortiferum		4.80	0.52		1		
01411003	**Low discrimination**						**01411003**
					SPO		
Lactobacillus fermentum		74.74	0.40	SUC 87	1		
Atopobium minutum		12.96	0.20	URE 1	1		
Clostridium novyi biovar A		7.83	0.20		99		
01411010	**Low discrimination**						**01411010**
					GRA	SPO	
Lactobacillus fermentum		35.73	0.40	SUC 87	99	1	
Clostridium novyi biovar A		21.22	0.33	MLZ 1	99	99	
Fusobacterium mortiferum		19.24	0.34		1	1	
01411012	**Low discrimination**						**01411012**
					SPO		
Lactobacillus fermentum		71.87	0.40	SUC 87	1		
Atopobium minutum		12.46	0.20	MLZ 1	1		
Clostridium novyi biovar A		7.53	0.20		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01411020	**Low discrimination**						**01411020**
					CAT		
Lactobacillus fermentum		73.31	0.66	SUC 87	1		
Atopobium minutum		19.07	0.49		1		
Propionibacterium acnes		2.34	0.26		95		
01411022	**Low discrimination**						**01411022**
Lactobacillus fermentum		84.43	0.66	SUC 87			
Atopobium minutum		14.64	0.46				
Pseudoramibacter alactolyticus		0.28	0.03				
01411042	**Low discrimination**						**01411042**
Lactobacillus fermentum		66.51	0.40	SUC 87			
Atopobium minutum		11.53	0.20	SOR 1			
Pseudoramibacter alactolyticus		10.41	0.05				
01411100	**Low discrimination**						**01411100**
					GRA		
Lactobacillus fermentum		54.69	0.78	SUC 87	99		
Fusobacterium mortiferum		42.12	0.74		1		
Actinomyces meyeri		1.63	0.41		99		
01411101	**Low discrimination**						**01411101**
					GRA		
Lactobacillus fermentum		54.77	0.44	SUC 87	99		
Fusobacterium mortiferum		42.18	0.40	URE 1	1		
Actinomyces meyeri		1.64	0.07		99		
01411102	**Good identification**						**01411102**
					COC	GRA	
Lactobacillus fermentum		91.71	0.78	SUC 87	1	99	
Fusobacterium mortiferum		7.85	0.58		1	1	
Blautia hansenii		0.26	0.15		99	99	
01411103	**Acceptable identification**						**01411103**
					GRA		
Lactobacillus fermentum		92.12	0.44	SUC 87	99		
Fusobacterium mortiferum		7.88	0.24	URE 1	1		
01411110	**Low discrimination**						**01411110**
					GRA		
Lactobacillus fermentum		54.77	0.44	SUC 87	99		
Fusobacterium mortiferum		42.18	0.40	MLZ 1	1		
Actinomyces meyeri		1.64	0.07		99		
01411112	**Acceptable identification**						**01411112**
					GRA		
Lactobacillus fermentum		92.12	0.44	SUC 87	99		
Fusobacterium mortiferum		7.88	0.24	MLZ 1	1		
01411120	**Good identification**						**01411120**
					GRA		
Lactobacillus fermentum		94.81	0.70	SUC 87	99		
Fusobacterium mortiferum		2.21	0.40		1		
Actinomyces meyeri		1.16	0.26		99		
01411121	**Good identification**						**01411121**
Lactobacillus fermentum		100.0	0.35	SUC 87			
01411122	**Very good identification**						**01411122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Lactobacillus fermentum		99.37	0.70	SUC 87	99	
Fusobacterium mortiferum		0.26	0.24		1	
Prevotella melaninogenica		0.17	0.03		1	
01411123	**Good identification**					**01411123**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411130	**Good identification**					**01411130**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411132	**Good identification**					**01411132**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411140	**Low discrimination**					**01411140**
					GRA	
Lactobacillus fermentum		54.77	0.44	SUC 87	99	
Fusobacterium mortiferum		42.18	0.40	SOR 1	1	
Actinomyces meyeri		1.64	0.07		99	
01411142	**Acceptable identification**					**01411142**
					GRA	
Lactobacillus fermentum		92.12	0.44	SUC 87	99	
Fusobacterium mortiferum		7.88	0.24	SOR 1	1	
01411160	**Good identification**					**01411160**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411162	**Good identification**					**01411162**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411200	**Low discrimination**					**01411200**
					GRA	
Lactobacillus fermentum		50.64	0.40	SUC 87	99	
Fusobacterium mortiferum		27.27	0.34	RHA 1	1	
Atopobium minutum		13.17	0.23		99	
01411202	**Low discrimination**					**01411202**
					GRA	
Lactobacillus fermentum		81.09	0.40	SUC 87	99	
Atopobium minutum		14.06	0.20	RHA 1	99	
Fusobacterium mortiferum		4.85	0.17		1	
01411300	**Low discrimination**					**01411300**
					GRA	
Lactobacillus fermentum		54.77	0.44	SUC 87	99	
Fusobacterium mortiferum		42.18	0.40	RHA 1	1	
Actinomyces meyeri		1.64	0.07		99	
01411302	**Acceptable identification**					**01411302**
					GRA	
Lactobacillus fermentum		92.12	0.44	SUC 87	99	
Fusobacterium mortiferum		7.88	0.24	RHA 1	1	
01411320	**Good identification**					**01411320**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411322	**Low discrimination**					**01411322**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		89.07	0.35	SUC 87	99		
Parabacteroides distasonis		10.93	0.17	RHA 1	1		
01411400	**Very good identification**						**01411400**
					GRA		
Lactobacillus fermentum		98.78	0.74	SUC 87	99		
Fusobacterium mortiferum		0.54	0.34		1		
Actinomyces meyeri		0.30	0.20		99		
01411401	**Good identification**						**01411401**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01411402	**Very good identification**						**01411402**
					GRA		
Lactobacillus fermentum		99.77	0.74	SUC 87	99		
Atopobium minutum		0.18	0.20		99		
Fusobacterium mortiferum		0.06	0.17		1		
01411403	**Good identification**						**01411403**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01411410	**Good identification**						**01411410**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01411412	**Good identification**						**01411412**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01411420	**Very good identification**						**01411420**
Lactobacillus fermentum		99.61	0.66	SUC 87			
Atopobium minutum		0.26	0.15				
Actinomyces meyeri		0.12	0.05				
01411422	**Very good identification**						**01411422**
Lactobacillus fermentum		99.83	0.66	SUC 87			
Atopobium minutum		0.18	0.12				
01411440	**Good identification**						**01411440**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01411442	**Good identification**						**01411442**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01411500	**Very good identification**						**01411500**
					GRA		
Lactobacillus fermentum		98.06	0.78	SUC 87	99		
Fusobacterium mortiferum		0.76	0.40		1		
Actinomyces meyeri		0.73	0.31		99		
01411501	**Good identification**						**01411501**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01411502	**Very good identification**						**01411502**
					COC	GRA	
Lactobacillus fermentum		99.84	0.78	SUC 87	1	99	
Fusobacterium mortiferum		0.09	0.24		1	1	
Blautia hansenii		0.05	0.03		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01411503	**Good identification**					**01411503**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01411510	**Good identification**					**01411510**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01411512	**Good identification**					**01411512**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01411520	**Very good identification**					**01411520**
					COC	
Lactobacillus fermentum		99.50	0.70	SUC 87	1	
Actinomyces meyeri		0.30	0.16		1	
Blautia hansenii		0.15	0.03		99	
01411521	**Good identification**					**01411521**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411522	**Very good identification**					**01411522**
Lactobacillus fermentum		99.95	0.70	SUC 87		
Bifidobacterium longum subsp. longum		0.05	0.05			
01411523	**Good identification**					**01411523**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411530	**Acceptable identification**					**01411530**
Lactobacillus fermentum		92.74	0.35	SUC 87		
Bifidobacterium longum subsp. longum		7.26	0.09	MLZ 1		
01411532	**Acceptable identification**					**01411532**
Lactobacillus fermentum		92.74	0.35	SUC 87		
Bifidobacterium longum subsp. longum		7.26	0.09	MLZ 1		
01411540	**Good identification**					**01411540**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01411542	**Good identification**					**01411542**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01411560	**Good identification**					**01411560**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411562	**Good identification**					**01411562**
Lactobacillus fermentum		100.0	0.35	SUC 87		
01411600	**Good identification**					**01411600**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01411602	**Good identification**					**01411602**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01411700	**Good identification**					**01411700**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		100.0	0.44	SUC 87				
01411702	**Good identification**							**01411702**
Lactobacillus fermentum		100.0	0.44	SUC 87				
01411720	**Good identification**							**01411720**
Lactobacillus fermentum		100.0	0.35	SUC 87				
01411722	**Acceptable identification**							**01411722**
					GRA			
Lactobacillus fermentum		93.01	0.35	SUC 87	99			
Parabacteroides distasonis		7.00	0.13	RHA 1	1			
01412000	**Low discrimination**							**01412000**
					COC	GRA		
Pseudoramibacter alactolyticus		53.12	0.61	RAF 1	1	99		
Eubacterium limosum		33.78	0.60		1	99		
Megasphaera elsdenii		10.38	0.60		99	1		
01412002	**Low discrimination**							**01412002**
					COC	GRA		
Pseudoramibacter alactolyticus		77.97	0.45	RAF 1	1	99		
Megasphaera elsdenii		15.24	0.44	BGL 10	99	1		
Eubacterium limosum		4.51	0.26		1	99		
01412010	**Acceptable identification**							**01412010**
					SPO			
Clostridium difficile		90.98	0.53	RAF 1	99			
Pseudoramibacter alactolyticus		4.74	0.27		1			
Eubacterium limosum		3.02	0.26		1			
01412020	**Acceptable identification**							**01412020**
					COC	GRA		
Pseudoramibacter alactolyticus		93.21	0.48	RAF 1	1	99		
Eubacterium limosum		3.39	0.26		1	99		
Megasphaera elsdenii		1.04	0.26		99	1		
01412040	**Low discrimination**							**01412040**
					COC	GRA		
Pseudoramibacter alactolyticus		64.19	0.51	RAF 1	1	99		
Megasphaera elsdenii		33.46	0.57		99	1		
Eubacterium limosum		1.65	0.26		1	99		
01412060	**Good identification**							**01412060**
					COC	CAT	GRA	
Pseudoramibacter alactolyticus		96.19	0.38	RAF 1	1	1	99	
Megasphaera elsdenii		2.87	0.23		99	1	1	
Propionibacterium acnes		0.95	0.11		1	95	99	
01413002	**Low discrimination**							**01413002**
					CAT			
Lactobacillus fermentum		51.77	0.40	SUC 87	1			
Pseudoramibacter alactolyticus		17.45	0.11	MAN 1	1			
Propionibacterium granulosum		15.31	0.26		90			
01413100	**Low discrimination**							**01413100**
					GRA			
Lactobacillus fermentum		53.77	0.44	SUC 87	99			
Fusobacterium mortiferum		41.41	0.40	MAN 1	1			
Actinomyces meyeri		1.61	0.07		99			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
01413102	**Acceptable identification**							**01413102**
					GRA			
Lactobacillus fermentum		90.24	0.44	SUC 87	99			
Fusobacterium mortiferum		7.72	0.24	MAN 1	1			
Propionibacterium propionicum		2.03	0.15		99			
01413120	**Good identification**							**01413120**
Lactobacillus fermentum		100.0	0.35	SUC 87				
01413122	**Good identification**							**01413122**
Lactobacillus fermentum		100.0	0.35	SUC 87				
01413400	**Good identification**							**01413400**
Lactobacillus fermentum		97.03	0.40	SUC 87				
Pseudoramibacter alactolyticus		2.97	-.07	MAN 1				
01413402	**Good identification**							**01413402**
Lactobacillus fermentum		100.0	0.40	SUC 87				
01413500	**Good identification**							**01413500**
Lactobacillus fermentum		100.0	0.44	SUC 87				
01413502	**Good identification**							**01413502**
Lactobacillus fermentum		100.0	0.44	SUC 87				
01413520	**Good identification**							**01413520**
Lactobacillus fermentum		100.0	0.35	SUC 87				
01413522	**Good identification**							**01413522**
Lactobacillus fermentum		100.0	0.35	SUC 87				
01414000	**Low discrimination**							**01414000**
					GRA	SPO		
Clostridium glycolicum		45.43	0.43	RAF 1	99	99		
Pseudoramibacter alactolyticus		19.38	0.32	SOR 95	99	1		
Fusobacterium varium		11.23	0.28		1	1		
01414040	**Very good identification**							**01414040**
					COC	SPO	GRA	
Clostridium glycolicum		99.24	0.65	RAF 1	1	99	99	
Pseudoramibacter alactolyticus		0.56	0.21		1	1	99	
Megasphaera elsdenii		0.16	0.23		99	1	1	
01414100	**Low discrimination**							**01414100**
					GRA			
Fusobacterium mortiferum		45.68	0.52	XYL 7	1			
Actinomyces meyeri		45.68	0.43		99			
Lactobacillus fermentum		7.01	0.40		99			
01414102	**Low discrimination**							**01414102**
					GRA			
Lactobacillus fermentum		51.94	0.40	SUC 87	99			
Fusobacterium mortiferum		37.60	0.36	GAL 99	1			
Actinomyces meyeri		10.46	0.17		99			
01414400	**Low discrimination**							**01414400**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		47.84	0.36	SUC 87	1		
Actinomyces meyeri		31.78	0.22	GAL 99	1		
Clostridium glycolicum		10.87	0.09		99		
01414402	**Good identification**						**01414402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01414502	**Good identification**						**01414502**
Lactobacillus fermentum		95.21	0.40	SUC 87			
Actinomyces meyeri		4.80	0.07	GAL 99			
01415000	**Low discrimination**						**01415000**
					GRA		
Lactobacillus fermentum		87.58	0.70	SUC 87	99		
Fusobacterium mortiferum		6.05	0.49		1		
Actinomyces meyeri		3.53	0.35		99		
01415001	**Acceptable identification**						**01415001**
					GRA		
Lactobacillus fermentum		93.54	0.36	SUC 87	99		
Fusobacterium mortiferum		6.46	0.15	URE 1	1		
01415002	**Very good identification**						**01415002**
					GRA		
Lactobacillus fermentum		98.82	0.70	SUC 87	99		
Fusobacterium mortiferum		0.76	0.32		1		
Atopobium minutum		0.30	0.20		99		
01415003	**Good identification**						**01415003**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01415010	**Acceptable identification**						**01415010**
					GRA		
Lactobacillus fermentum		93.54	0.36	SUC 87	99		
Fusobacterium mortiferum		6.46	0.15	MLZ 1	1		
01415012	**Good identification**						**01415012**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01415020	**Very good identification**						**01415020**
Lactobacillus fermentum		97.75	0.62	SUC 87			
Actinomyces meyeri		1.61	0.21				
Atopobium minutum		0.44	0.15				
01415022	**Very good identification**						**01415022**
Lactobacillus fermentum		99.70	0.62	SUC 87			
Atopobium minutum		0.30	0.12				
01415040	**Good identification**						**01415040**
					GRA	SPO	
Clostridium glycolicum		97.80	0.49	RAF 1	99	99	
Lactobacillus fermentum		2.06	0.36	GAL 10	99	1	
Fusobacterium mortiferum		0.14	0.15		1	1	
01415042	**Low discrimination**						**01415042**
					SPO		
Lactobacillus fermentum		67.59	0.36	SUC 87	1		
Clostridium glycolicum		32.41	0.15	SOR 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01415100	**Low discrimination**						**01415100**
					GRA		
Lactobacillus fermentum		82.97	0.74	SUC 87	99		
Fusobacterium mortiferum		8.19	0.55		1		
Actinomyces meyeri		8.19	0.46		99		
01415101	**Low discrimination**						**01415101**
					GRA		
Lactobacillus fermentum		83.51	0.40	SUC 87	99		
Fusobacterium mortiferum		8.24	0.21	URE 1	1		
Actinomyces meyeri		8.24	0.12		99		
01415102	**Very good identification**						**01415102**
					GRA		
Lactobacillus fermentum		98.53	0.74	SUC 87	99		
Fusobacterium mortiferum		1.08	0.39		1		
Actinomyces meyeri		0.30	0.20		99		
01415103	**Good identification**						**01415103**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01415110	**Low discrimination**						**01415110**
					GRA		
Lactobacillus fermentum		83.51	0.40	SUC 87	99		
Fusobacterium mortiferum		8.24	0.21	MLZ 1	1		
Actinomyces meyeri		8.24	0.12		99		
01415112	**Good identification**						**01415112**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01415120	**Very good identification**						**01415120**
					GRA		
Lactobacillus fermentum		95.61	0.66	SUC 87	99		
Actinomyces meyeri		3.86	0.31		99		
Fusobacterium mortiferum		0.29	0.21		1		
01415122	**Very good identification**						**01415122**
					GRA		
Lactobacillus fermentum		99.25	0.66	SUC 87	99		
Parabacteroides distasonis		0.63	0.25		1		
Actinomyces meyeri		0.12	0.05		99		
01415140	**Low discrimination**						**01415140**
					GRA	SPO	
Lactobacillus fermentum		67.05	0.40	SUC 87	99	1	
Clostridium glycolicum		19.71	0.15	SOR 1	99	99	
Fusobacterium mortiferum		6.62	0.21		1	1	
01415142	**Good identification**						**01415142**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01415200	**Acceptable identification**						**01415200**
					GRA		
Lactobacillus fermentum		93.54	0.36	SUC 87	99		
Fusobacterium mortiferum		6.46	0.15	RHA 1	1		
01415202	**Good identification**						**01415202**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01415300	**Low discrimination**						**01415300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Lactobacillus fermentum		83.51	0.40	SUC 87	99	
Fusobacterium mortiferum		8.24	0.21	RHA 1	1	
Actinomyces meyeri		8.24	0.12		99	
01415302	**Good identification**					**01415302**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01415400	**Very good identification**					**01415400**
					GRA	
Lactobacillus fermentum		98.94	0.70	SUC 87	99	
Actinomyces meyeri		1.00	0.25		99	
Fusobacterium mortiferum		0.07	0.15		1	
01415401	**Good identification**					**01415401**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01415402	**Very good identification**					**01415402**
Lactobacillus fermentum		100.0	0.70	SUC 87		
01415403	**Good identification**					**01415403**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01415410	**Good identification**					**01415410**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01415412	**Good identification**					**01415412**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01415420	**Very good identification**					**01415420**
Lactobacillus fermentum		99.59	0.62	SUC 87		
Actinomyces meyeri		0.41	0.10			
01415422	**Very good identification**					**01415422**
Lactobacillus fermentum		100.0	0.62	SUC 87		
01415440	**Low discrimination**					**01415440**
					SPO	
Lactobacillus fermentum		67.59	0.36	SUC 87	1	
Clostridium glycolicum		32.41	0.15	SOR 1	99	
01415442	**Good identification**					**01415442**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01415500	**Very good identification**					**01415500**
					COC	
Lactobacillus fermentum		97.37	0.74	SUC 87	1	
Actinomyces meyeri		2.40	0.35		1	
Blautia hansenii		0.13	0.06		99	
01415501	**Good identification**					**01415501**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01415502	**Very good identification**					**01415502**
Lactobacillus fermentum		99.92	0.74	SUC 87		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		0.08	0.10			
01415503	**Good identification**					**01415503**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01415510	**Good identification**					**01415510**
Lactobacillus fermentum		96.78	0.40	SUC 87		
Bifidobacterium longum subsp. longum		3.23	0.07	MLZ 1		
01415512	**Good identification**					**01415512**
Lactobacillus fermentum		96.78	0.40	SUC 87		
Bifidobacterium longum subsp. longum		3.23	0.07	MLZ 1		
01415520	**Very good identification**					**01415520**
Lactobacillus fermentum		98.77	0.66	SUC 87		
Actinomyces meyeri		1.00	0.21			
Bifidobacterium longum subsp. longum		0.23	0.14			
01415522	**Very good identification**					**01415522**
					GRA	
Lactobacillus fermentum		99.38	0.66	SUC 87	99	
Parabacteroides distasonis		0.39	0.21		1	
Bifidobacterium longum subsp. longum		0.24	0.14		99	
01415540	**Good identification**					**01415540**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01415542	**Good identification**					**01415542**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01415600	**Good identification**					**01415600**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01415602	**Good identification**					**01415602**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01415700	**Good identification**					**01415700**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01415702	**Good identification**					**01415702**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01417000	**Low discrimination**					**01417000**
					GRA	
Lactobacillus fermentum		89.19	0.36	SUC 87	99	
Fusobacterium mortiferum		6.16	0.15	MAN 1	1	
Pseudoramibacter alactolyticus		4.65	-.07		99	
01417002	**Good identification**					**01417002**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01417100	**Low discrimination**					**01417100**
					GRA	
Lactobacillus fermentum		83.51	0.40	SUC 87	99	
Fusobacterium mortiferum		8.24	0.21	MAN 1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		8.24	0.12		99		
01417102	**Good identification**						**01417102**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01417400	**Good identification**						**01417400**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01417402	**Good identification**						**01417402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01417500	**Good identification**						**01417500**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01417502	**Good identification**						**01417502**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01420000	**Low discrimination**						**01420000**
					COC	SPO	
Clostridium botulinum biovar A		64.69	0.51	RAF 1	1	99	
Clostridium sporogenes		20.79	0.40		1	99	
Peptostreptococcus anaerobius		9.08	0.32		99	1	
01420040	**Good identification**						**01420040**
Clostridium botulinum biovar A		95.25	0.42	RAF 1			
Clostridium novyi biovar A		3.26	0.29				
Clostridium sporogenes		1.04	0.06				
01430000	**Low discrimination**						**01430000**
					GRA	CAT	
Fusobacterium mortiferum		38.51	0.56		1	5	
Pseudoramibacter alactolyticus		20.43	0.32		99	1	
Propionibacterium granulosum		17.06	0.47		99	90	
01430002	**Low discrimination**						**01430002**
					GRA	CAT	
Fusobacterium mortiferum		28.32	0.40	BGL 10	1	5	
Propionibacterium granulosum		28.22	0.36		99	90	
Pseudoramibacter alactolyticus		15.02	0.15		99	1	
01430020	**Low discrimination**						**01430020**
					SPO	CAT	
Propionibacterium granulosum		40.53	0.36	SUC 93	1	90	
Pseudoramibacter alactolyticus		34.27	0.19	MNS 99	1	1	
Clostridium cadaveris		12.54	0.15		99	1	
01430100	**Very good identification**						**01430100**
					GRA	CAT	
Fusobacterium mortiferum		99.45	0.62		1	5	
Pseudoramibacter alactolyticus		0.23	-.03		99	1	
Propionibacterium granulosum		0.19	0.13		99	90	
01430102	**Good identification**						**01430102**
Fusobacterium mortiferum		100.0	0.46	BGL 10			
01431000	**Low discrimination**						**01431000**
					GRA	CAT	
Fusobacterium mortiferum		72.34	0.59		1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		21.36	0.47		99	90	
Lactobacillus fermentum		4.54	0.40		99	1	
01431002	**Low discrimination**						**01431002**
					GRA	CAT	
Fusobacterium mortiferum		42.96	0.43	BGL 10	1	5	
Propionibacterium granulosum		28.55	0.36		99	90	
Lactobacillus fermentum		24.28	0.40		99	1	
01431020	**Low discrimination**						**01431020**
					GRA	CAT	
Propionibacterium granulosum		66.93	0.36	SUC 93	99	90	
Lactobacillus fermentum		18.98	0.32	MNS 99	99	1	
Fusobacterium mortiferum		9.16	0.25		1	5	
01431100	**Very good identification**						**01431100**
					GRA	CAT	
Fusobacterium mortiferum		95.45	0.65		1	5	
Lactobacillus fermentum		4.19	0.44		99	1	
Propionibacterium granulosum		0.12	0.13		99	90	
01431102	**Low discrimination**						**01431102**
					GRA		
Fusobacterium mortiferum		71.68	0.49	BGL 10	1		
Lactobacillus fermentum		28.32	0.44		99		
01431120	**Low discrimination**						**01431120**
					GRA		
Lactobacillus fermentum		57.15	0.35	SUC 87	99		
Fusobacterium mortiferum		39.45	0.31	TRE 1	1		
Clostridium perfringens		3.40	0.02		99		
01431122	**Low discrimination**						**01431122**
					GRA		
Lactobacillus fermentum		79.18	0.35	SUC 87	99		
Parabacteroides distasonis		14.75	0.20	TRE 1	1		
Fusobacterium mortiferum		6.07	0.15		1		
01431400	**Low discrimination**						**01431400**
					GRA	CAT	
Lactobacillus fermentum		82.76	0.40	SUC 87	99	1	
Fusobacterium mortiferum		13.31	0.25	TRE 1	1	5	
Propionibacterium granulosum		3.93	0.13		99	90	
01431402	**Good identification**						**01431402**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01431500	**Low discrimination**						**01431500**
					GRA		
Lactobacillus fermentum		81.30	0.44	SUC 87	99		
Fusobacterium mortiferum		18.70	0.31	TRE 1	1		
01431502	**Good identification**						**01431502**
					GRA		
Lactobacillus fermentum		97.51	0.44	SUC 87	99		
Fusobacterium mortiferum		2.49	0.15	TRE 1	1		
01431520	**Good identification**						**01431520**
Lactobacillus fermentum		100.0	0.35	SUC 87			
01431522	**Low discrimination**						**01431522**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		89.75	0.35	SUC 87	99		
Parabacteroides distasonis		10.25	0.16	TRE 1	1		
01432000	**Low discrimination**						**01432000**
					CAT		
Propionibacterium granulosum		35.10	0.45	SUC 93	90		
Pseudoramibacter alactolyticus		30.01	0.27	MNS 99	1		
Eubacterium limosum		19.08	0.26		1		
01433000	**Acceptable identification**						**01433000**
					GRA	CAT	
Propionibacterium granulosum		94.88	0.45	SUC 93	99	90	
Fusobacterium mortiferum		4.30	0.25	MNS 99	1	5	
Pseudoramibacter alactolyticus		0.82	-.07		99	1	
01434000	**Low discrimination**						**01434000**
					GRA	SPO	
Fusobacterium mortiferum		74.69	0.37	XYL 7	1	1	
Clostridium glycolicum		12.47	0.09		99	99	
Pseudoramibacter alactolyticus		5.32	-.03		99	1	
01434100	**Good identification**						**01434100**
					GRA		
Fusobacterium mortiferum		96.73	0.43	XYL 7	1		
Actinomyces meyeri		3.27	0.09		99		
01435000	**Low discrimination**						**01435000**
					GRA	CAT	
Fusobacterium mortiferum		65.38	0.40	XYL 7	1	5	
Lactobacillus fermentum		32.03	0.36		99	1	
Propionibacterium granulosum		2.59	0.13		99	90	
01435002	**Low discrimination**						**01435002**
					GRA		
Lactobacillus fermentum		81.52	0.36	SUC 87	99		
Fusobacterium mortiferum		18.49	0.23	TRE 1	1		
01435100	**Low discrimination**						**01435100**
					GRA		
Fusobacterium mortiferum		72.65	0.46	XYL 7	1		
Lactobacillus fermentum		24.89	0.40		99		
Actinomyces meyeri		2.46	0.12		99		
01435102	**Low discrimination**						**01435102**
					GRA		
Lactobacillus fermentum		75.51	0.40	SUC 87	99		
Fusobacterium mortiferum		24.49	0.30	TRE 1	1		
01435122	**Low discrimination**						**01435122**
					GRA		
Parabacteroides distasonis		74.06	0.36	MNS 98	1		
Lactobacillus fermentum		25.94	0.31		99		
01435322	**Good identification**						**01435322**
Parabacteroides distasonis		100.0	0.45	MNS 98			
01435332	**Good identification**						**01435332**
Parabacteroides distasonis		100.0	0.40	MNS 98			
01435400	**Good identification**						**01435400**
Lactobacillus fermentum		100.0	0.36	SUC 87			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01435402	**Good identification**					**01435402**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01435500	**Good identification**					**01435500**
					GRA	
Lactobacillus fermentum		97.14	0.40	SUC 87	99	
Fusobacterium mortiferum		2.86	0.12	TRE 1	1	
01435502	**Good identification**					**01435502**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01435722	**Good identification**					**01435722**
Parabacteroides distasonis		100.0	0.41	MNS 98		
01435732	**Good identification**					**01435732**
Parabacteroides distasonis		100.0	0.36	MNS 98		
01450000	**Low discrimination**					**01450000**
					GRA	
Fusobacterium mortiferum		58.12	0.57		1	
Pseudoramibacter alactolyticus		27.63	0.32		99	
Fusobacterium varium		3.88	0.17		1	
01450002	**Low discrimination**					**01450002**
					GRA	
Fusobacterium mortiferum		52.07	0.40	BGL 10	1	
Pseudoramibacter alactolyticus		24.75	0.15		99	
Atopobium minutum		20.58	0.28		99	
01450100	**Very good identification**					**01450100**
					GRA	
Fusobacterium mortiferum		99.68	0.63		1	
Pseudoramibacter alactolyticus		0.21	-.03		99	
Actinomyces meyeri		0.12	0.04		99	
01450102	**Good identification**					**01450102**
Fusobacterium mortiferum		100.0	0.47	BGL 10		
01451000	**Good identification**					**01451000**
					GRA	
Fusobacterium mortiferum		92.76	0.60		1	
Lactobacillus fermentum		5.22	0.40		99	
Atopobium minutum		1.36	0.23		99	
01451002	**Low discrimination**					**01451002**
					GRA	
Fusobacterium mortiferum		62.73	0.43	BGL 10	1	
Lactobacillus fermentum		31.77	0.40		99	
Atopobium minutum		5.51	0.20		99	
01451100	**Very good identification**					**01451100**
					GRA	
Fusobacterium mortiferum		96.01	0.66		1	
Lactobacillus fermentum		3.78	0.44		99	
Actinomyces meyeri		0.11	0.07		99	
01451102	**Low discrimination**					**01451102**
					GRA	
Fusobacterium mortiferum		73.85	0.50	BGL 10	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		26.15	0.44		99		
01451120	**Low discrimination**						**01451120**
					GRA		
Lactobacillus fermentum		53.51	0.35	SUC 87	99		
Fusobacterium mortiferum		41.21	0.32	CEL 1	1		
Prevotella melaninogenica		5.28	-.01		1		
01451122	**Acceptable identification**						**01451122**
					GRA		
Lactobacillus fermentum		92.12	0.35	SUC 87	99		
Fusobacterium mortiferum		7.88	0.16	CEL 1	1		
01451400	**Low discrimination**						**01451400**
					GRA		
Lactobacillus fermentum		84.78	0.40	SUC 87	99		
Fusobacterium mortiferum		15.22	0.26	CEL 1	1		
01451402	**Good identification**						**01451402**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01451500	**Low discrimination**						**01451500**
					GRA		
Lactobacillus fermentum		79.57	0.44	SUC 87	99		
Fusobacterium mortiferum		20.43	0.32	CEL 1	1		
01451502	**Good identification**						**01451502**
					GRA		
Lactobacillus fermentum		97.23	0.44	SUC 87	99		
Fusobacterium mortiferum		2.77	0.16	CEL 1	1		
01451520	**Good identification**						**01451520**
Lactobacillus fermentum		100.0	0.35	SUC 87			
01451522	**Good identification**						**01451522**
Lactobacillus fermentum		100.0	0.35	SUC 87			
01454000	**Low discrimination**						**01454000**
					GRA	SPO	
Fusobacterium mortiferum		79.98	0.38	XYL 7	1	1	
Clostridium glycolicum		11.97	0.09		99	99	
Pseudoramibacter alactolyticus		5.10	-.03		99	1	
01454100	**Good identification**						**01454100**
					GRA		
Fusobacterium mortiferum		97.06	0.44	XYL 7	1		
Actinomyces meyeri		2.94	0.09		99		
01455000	**Low discrimination**						**01455000**
					GRA		
Fusobacterium mortiferum		69.49	0.41	XYL 7	1		
Lactobacillus fermentum		30.51	0.36		99		
01455002	**Low discrimination**						**01455002**
					GRA		
Lactobacillus fermentum		79.80	0.36	SUC 87	99		
Fusobacterium mortiferum		20.20	0.24	CEL 1	1		
01455100	**Low discrimination**						**01455100**
					GRA		
Fusobacterium mortiferum		74.78	0.47	XYL 7	1		
Lactobacillus fermentum		22.96	0.40		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		2.27	0.12		99		
01455102	**Low discrimination**						**01455102**
					GRA		
Lactobacillus fermentum		73.43	0.40	SUC 87	99		
Fusobacterium mortiferum		26.58	0.31	CEL 1	1		
01455400	**Good identification**						**01455400**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01455402	**Good identification**						**01455402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01455500	**Good identification**						**01455500**
					GRA		
Lactobacillus fermentum		96.81	0.40	SUC 87	99		
Fusobacterium mortiferum		3.19	0.13	CEL 1	1		
01455502	**Good identification**						**01455502**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01470000	**Good identification**						**01470000**
					GRA	CAT	
Fusobacterium mortiferum		97.13	0.48		1	5	
Pseudoramibacter alactolyticus		1.56	-.03		99	1	
Propionibacterium granulosum		1.30	0.13		99	90	
01470100	**Good identification**						**01470100**
Fusobacterium mortiferum		100.0	0.54				
01470102	**Good identification**						**01470102**
Fusobacterium mortiferum		100.0	0.38	BGL 10			
01471000	**Good identification**						**01471000**
					GRA	CAT	
Fusobacterium mortiferum		99.11	0.51		1	5	
Propionibacterium granulosum		0.89	0.13		99	90	
01471100	**Very good identification**						**01471100**
Fusobacterium mortiferum		100.0	0.57				
01471102	**Good identification**						**01471102**
Fusobacterium mortiferum		100.0	0.41	BGL 10			
01475100	**Good identification**						**01475100**
Fusobacterium mortiferum		100.0	0.38	XYL 7			
01475322	**Good identification**						**01475322**
Parabacteroides distasonis		100.0	0.40	MNS 98			
01475332	**Good identification**						**01475332**
Parabacteroides distasonis		100.0	0.35	MNS 98			
01475722	**Good identification**						**01475722**
Parabacteroides distasonis		100.0	0.36	MNS 98			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01500000	**Low discrimination**							**01500000**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		42.35	0.63	RAF 1	99	99	1	
Prevotella disiens		23.34	0.57		1	1	1	
Clostridium botulinum biovar A		14.97	0.60		1	99	99	
01500001	**Low discrimination**							**01500001**
					COC	SPO		
Clostridium novyi biovar A		80.32	0.66	RAF 1	1	99		
Clostridium sordellii		7.75	0.32		1	99		
Peptostreptococcus anaerobius		4.94	0.29		99	1		
01500002	**Low discrimination**							**01500002**
					COC	SPO		
Clostridium sporogenes		54.34	0.45	RAF 1	1	99		
Clostridium novyi biovar A		24.42	0.53		1	99		
Peptostreptococcus anaerobius		8.52	0.29		99	1		
01500003	**Good identification**							**01500003**
					COC	SPO		
Clostridium novyi biovar A		95.35	0.53	RAF 1	1	99		
Clostridium sporogenes		2.14	0.11		1	99		
Anaerococcus prevotii		1.65	0.05		99	1		
01500010	**Low discrimination**							**01500010**
					COC	GRA	SPO	
Clostridium novyi biovar A		88.14	0.66	RAF 1	1	99	99	
Peptostreptococcus anaerobius		5.43	0.29		99	99	1	
Prevotella disiens		2.99	0.23		1	1	1	
01500011	**Very good identification**							**01500011**
					COC	SPO		
Clostridium novyi biovar A		99.84	0.66	RAF 1	1	99		
Clostridium sordellii		0.10	-.03		1	99		
Peptostreptococcus anaerobius		0.06	-.06		99	1		
01500012	**Good identification**							**01500012**
					COC	SPO		
Clostridium novyi biovar A		97.47	0.53	RAF 1	1	99		
Clostridium sporogenes		2.19	0.11		1	99		
Peptostreptococcus anaerobius		0.34	-.06		99	1		
01500013	**Good identification**							**01500013**
Clostridium novyi biovar A		100.0	0.53	RAF 1				
01500020	**Low discrimination**							**01500020**
					COC	GRA	SPO	
Clostridium novyi biovar A		59.09	0.56	RAF 1	1	99	99	
Peptostreptococcus anaerobius		14.55	0.29		99	99	1	
Prevotella disiens		8.02	0.23		1	1	1	
01500021	**Good identification**							**01500021**
					COC	SPO		
Clostridium novyi biovar A		99.37	0.56	RAF 1	1	99		
Clostridium sordellii		0.39	-.03		1	99		
Peptostreptococcus anaerobius		0.25	-.06		99	1		
01500022	**Low discrimination**							**01500022**
					SPO			
Clostridium novyi biovar A		80.39	0.43	RAF 1	99			
Clostridium sporogenes		7.23	0.11		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		7.12	0.21		1		
01500023	**Good identification**						**01500023**
Clostridium novyi biovar A		100.0	0.43	RAF 1			
01500030	**Good identification**						**01500030**
					COC	SPO	
Clostridium novyi biovar A		99.75	0.56	RAF 1	1	99	
Peptostreptococcus anaerobius		0.25	-.06		99	1	
01500031	**Good identification**						**01500031**
Clostridium novyi biovar A		100.0	0.56	RAF 1			
01500032	**Good identification**						**01500032**
Clostridium novyi biovar A		100.0	0.43	RAF 1			
01500033	**Good identification**						**01500033**
Clostridium novyi biovar A		100.0	0.43	RAF 1			
01500040	**Low discrimination**						**01500040**
					COC	GRA	
Clostridium novyi biovar A		43.68	0.63	RAF 1	1	99	
Clostridium botulinum biovar A		42.15	0.51		1	99	
Megasphaera elsdenii		4.21	0.38		99	1	
01500041	**Very good identification**						**01500041**
Clostridium novyi biovar A		98.26	0.63	RAF 1			
Clostridium botulinum biovar A		0.96	0.17				
Clostridium sordellii		0.59	0.08				
01500042	**Low discrimination**						**01500042**
					COC	GRA	
Clostridium novyi biovar A		86.59	0.50	RAF 1	1	99	
Megasphaera elsdenii		5.25	0.22		99	1	
Clostridium botulinum biovar A		4.78	0.17		1	99	
01500043	**Good identification**						**01500043**
Clostridium novyi biovar A		100.0	0.50	RAF 1			
01500050	**Very good identification**						**01500050**
					COC	GRA	
Clostridium novyi biovar A		98.85	0.63	RAF 1	1	99	
Clostridium botulinum biovar A		0.96	0.17		1	99	
Megasphaera elsdenii		0.10	0.04		99	1	
01500051	**Very good identification**						**01500051**
Clostridium novyi biovar A		100.0	0.63	RAF 1			
01500052	**Good identification**						**01500052**
Clostridium novyi biovar A		100.0	0.50	RAF 1			
01500053	**Good identification**						**01500053**
Clostridium novyi biovar A		100.0	0.50	RAF 1			
01500060	**Good identification**						**01500060**
					COC	GRA	
Clostridium novyi biovar A		95.25	0.53	RAF 1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		3.71	0.17		1	99		
Megasphaera elsdenii		0.37	0.04		99	1		
01500061	**Good identification**							**01500061**
Clostridium novyi biovar A		100.0	0.53	RAF 1				
01500062	**Good identification**							**01500062**
Clostridium novyi biovar A		100.0	0.40	RAF 1				
01500063	**Good identification**							**01500063**
Clostridium novyi biovar A		100.0	0.40	RAF 1				
01500070	**Good identification**							**01500070**
Clostridium novyi biovar A		100.0	0.53	RAF 1				
01500071	**Good identification**							**01500071**
Clostridium novyi biovar A		100.0	0.53	RAF 1				
01500072	**Good identification**							**01500072**
Clostridium novyi biovar A		100.0	0.40	RAF 1				
01500073	**Good identification**							**01500073**
Clostridium novyi biovar A		100.0	0.40	RAF 1				
01500100	**Very good identification**							**01500100**
					COC	GRA		
Blautia hansenii		98.25	0.66	GAL 99	99	99		
Peptostreptococcus anaerobius		0.70	0.29		99	99		
Prevotella disiens		0.39	0.23		1	1		
01500102	**Good identification**							**01500102**
					COC	SPO		
Blautia hansenii		99.23	0.50	GAL 99	99	1		
Clostridium sporogenes		0.41	0.11	BGL 10	1	99		
Clostridium novyi biovar A		0.19	0.19		1	99		
01500120	**Low discrimination**							**01500120**
					COC	GRA	SPO	
Blautia hansenii		80.93	0.50	GAL 99	99	99	1	
Prevotella bivia		18.66	0.32	NAG 10	1	1	1	
Clostridium novyi biovar A		0.21	0.21		1	99	99	
01500500	**Good identification**							**01500500**
					COC			
Blautia hansenii		99.83	0.53	GAL 99	99			
Lactobacillus fermentum		0.07	0.18		1			
Actinomyces meyeri		0.07	0.07		1			
01500502	**Good identification**							**01500502**
					COC			
Blautia hansenii		99.40	0.37	GAL 99	99			
Lactobacillus fermentum		0.61	0.18	BGL 10	1			
01500520	**Good identification**							**01500520**
					COC	GRA		
Blautia hansenii		98.70	0.37	GAL 99	99	99		
Prevotella bivia		1.30	-.03	NAG 10	1	1		
01501000	**Good identification**							**01501000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Blautia hansenii		94.78	0.66	LAC 99	99	99	1	
Prevotella intermedia		1.60	0.42		1	1	1	
Clostridium novyi biovar A		1.22	0.50		1	99	99	
01501001	**Low discrimination**							**01501001**
					COC	SPO		
Clostridium novyi biovar A		51.00	0.50	RAF 1	1	99		
Blautia hansenii		39.94	0.32	GAL 10	99	1		
Clostridium sordellii		7.82	0.19		1	99		
01501002	**Acceptable identification**							**01501002**
					COC	SPO		
Blautia hansenii		91.24	0.50	LAC 99	99	1		
Lactobacillus fermentum		5.95	0.48	BGL 10	1	1		
Clostridium novyi biovar A		1.87	0.37		1	99		
01501003	**Low discrimination**							**01501003**
					COC	SPO		
Clostridium novyi biovar A		65.57	0.37	RAF 1	1	99		
Blautia hansenii		32.33	0.15	GAL 10	99	1		
Lactobacillus fermentum		2.11	0.14		1	1		
01501010	**Low discrimination**							**01501010**
					COC	GRA	SPO	
Clostridium novyi biovar A		55.32	0.50	RAF 1	1	99	99	
Blautia hansenii		43.32	0.32	GAL 10	99	99	1	
Prevotella intermedia		0.73	0.08		1	1	1	
01501011	**Good identification**							**01501011**
					COC	SPO		
Clostridium novyi biovar A		99.22	0.50	RAF 1	1	99		
Blautia hansenii		0.79	-.03	GAL 10	99	1		
01501012	**Low discrimination**							**01501012**
					COC	SPO		
Clostridium novyi biovar A		65.57	0.37	RAF 1	1	99		
Blautia hansenii		32.33	0.15	GAL 10	99	1		
Lactobacillus fermentum		2.11	0.14		1	1		
01501013	**Good identification**							**01501013**
Clostridium novyi biovar A		100.0	0.37	RAF 1				
01501020	**Acceptable identification**							**01501020**
					COC	SPO		
Blautia hansenii		94.53	0.50	LAC 99	99	1		
Clostridium novyi biovar A		2.74	0.39	NAG 10	1	99		
Lactobacillus fermentum		2.05	0.40		1	1		
01501021	**Low discrimination**							**01501021**
					COC	SPO		
Clostridium novyi biovar A		74.18	0.39	RAF 1	1	99		
Blautia hansenii		25.82	0.15	GAL 10	99	1		
01501030	**Low discrimination**							**01501030**
					COC	SPO		
Clostridium novyi biovar A		74.18	0.39	RAF 1	1	99		
Blautia hansenii		25.82	0.15	GAL 10	99	1		
01501031	**Good identification**							**01501031**
Clostridium novyi biovar A		100.0	0.39	RAF 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01501041	**Good identification**						**01501041**
					COC	SPO	
Clostridium novyi biovar A		97.90	0.47	RAF 1	1	99	
Blautia hansenii		1.16	-.03	GAL 10	99	1	
Clostridium sordellii		0.94	-.05		1	99	
01501050	**Good identification**						**01501050**
					COC	SPO	
Clostridium novyi biovar A		98.83	0.47	RAF 1	1	99	
Blautia hansenii		1.17	-.03	GAL 10	99	1	
01501051	**Good identification**						**01501051**
Clostridium novyi biovar A		100.0	0.47	RAF 1			
01501060	**Low discrimination**						**01501060**
					COC	SPO	
Clostridium novyi biovar A		65.70	0.36	RAF 1	1	99	
Blautia hansenii		34.30	0.15	GAL 10	99	1	
01501061	**Good identification**						**01501061**
Clostridium novyi biovar A		100.0	0.36	RAF 1			
01501070	**Good identification**						**01501070**
Clostridium novyi biovar A		100.0	0.36	RAF 1			
01501071	**Good identification**						**01501071**
Clostridium novyi biovar A		100.0	0.36	RAF 1			
01501100	**Excellent identification**						**01501100**
					COC	GRA	
Blautia hansenii		99.99	1.00		99	99	
Lactobacillus fermentum		0.01	0.52		1	99	
Fusobacterium mortiferum		0.00	0.25		1	1	
01501101	**Very good identification**						**01501101**
					COC	SPO	
Blautia hansenii		99.97	0.66	URE 1	99	1	
Clostridium novyi biovar A		0.01	0.15		1	99	
Lactobacillus fermentum		0.01	0.18		1	1	
01501102	**Very good identification**						**01501102**
					COC		
Blautia hansenii		99.89	0.84	BGL 10	99		
Lactobacillus fermentum		0.11	0.52		1		
01501103	**Good identification**						**01501103**
					COC		
Blautia hansenii		99.89	0.50	URE 1	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		
01501110	**Very good identification**						**01501110**
					COC	SPO	
Blautia hansenii		99.98	0.66	MLZ 1	99	1	
Clostridium novyi biovar A		0.01	0.15		1	99	
Lactobacillus fermentum		0.01	0.18		1	1	
01501112	**Good identification**						**01501112**
					COC		
Blautia hansenii		99.89	0.50	MLZ 1	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01501120	**Very good identification**						**01501120**
					COC	GRA	
Blautia hansenii		99.81	0.84	NAG 10	99	99	
Prevotella melaninogenica		0.12	0.33		1	1	
Lactobacillus fermentum		0.04	0.44		1	99	
01501121	**Good identification**						**01501121**
					COC	GRA	
Blautia hansenii		99.88	0.50	NAG 10	99	99	
Prevotella melaninogenica		0.12	-.01	URE 1	1	1	
01501122	**Very good identification**						**01501122**
					COC	GRA	
Blautia hansenii		99.48	0.67	NAG 10	99	99	
Lactobacillus fermentum		0.32	0.44	BGL 10	1	99	
Prevotella melaninogenica		0.20	0.21		1	1	
01501130	**Good identification**						**01501130**
					COC	GRA	
Blautia hansenii		99.88	0.50	MLZ 1	99	99	
Prevotella melaninogenica		0.12	-.01	NAG 10	1	1	
01501140	**Very good identification**						**01501140**
					COC	SPO	
Blautia hansenii		99.98	0.66	SOR 1	99	1	
Lactobacillus fermentum		0.01	0.18		1	1	
Clostridium novyi biovar A		0.01	0.12		1	99	
01501142	**Good identification**						**01501142**
					COC		
Blautia hansenii		99.89	0.50	SOR 1	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		
01501160	**Good identification**						**01501160**
					COC	GRA	
Blautia hansenii		99.88	0.50	NAG 10	99	99	
Prevotella melaninogenica		0.12	-.01	SOR 1	1	1	
01501300	**Very good identification**						**01501300**
					COC		
Blautia hansenii		99.99	0.66	RHA 1	99		
Lactobacillus fermentum		0.01	0.18		1		
01501302	**Good identification**						**01501302**
					COC		
Blautia hansenii		99.89	0.50	RHA 1	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		
01501320	**Good identification**						**01501320**
					COC	GRA	
Blautia hansenii		99.88	0.50	RHA 1	99	99	
Prevotella melaninogenica		0.12	-.01	NAG 10	1	1	
01501400	**Good identification**						**01501400**
					COC		
Blautia hansenii		95.77	0.53	LAC 99	99		
Lactobacillus fermentum		3.93	0.48		1		
Gemella morbillorum		0.10	0.09		99		
01501402	**Low discrimination**						**01501402**
					COC		
Blautia hansenii		73.03	0.37	LAC 99	99		
Lactobacillus fermentum		26.97	0.48	BGL 10	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01501420	**Low discrimination**					**01501420**
					COC	
Blautia hansenii		89.04	0.37	LAC 99	99	
Lactobacillus fermentum		10.96	0.40	NAG 10	1	
01501422	**Low discrimination**					**01501422**
					COC	
Lactobacillus fermentum		52.56	0.40	SUC 87	1	
Blautia hansenii		47.44	0.20	FRU 99	99	
01501500	**Very good identification**					**01501500**
					COC	
Blautia hansenii		99.93	0.87		99	
Lactobacillus fermentum		0.07	0.52		1	
Bifidobacterium longum subsp. longum		0.00	0.13		1	
01501501	**Good identification**					**01501501**
					COC	
Blautia hansenii		99.93	0.53	URE 1	99	
Lactobacillus fermentum		0.07	0.18		1	
01501502	**Very good identification**					**01501502**
					COC	
Blautia hansenii		99.39	0.71	BGL 10	99	
Lactobacillus fermentum		0.61	0.52		1	
Bifidobacterium longum subsp. longum		0.01	0.13		1	
01501503	**Good identification**					**01501503**
					COC	
Blautia hansenii		99.40	0.37	URE 1	99	
Lactobacillus fermentum		0.61	0.18	BGL 10	1	
01501510	**Good identification**					**01501510**
					COC	
Blautia hansenii		99.77	0.53	MLZ 1	99	
Bifidobacterium longum subsp. longum		0.16	0.17		1	
Lactobacillus fermentum		0.07	0.18		1	
01501512	**Good identification**					**01501512**
					COC	
Blautia hansenii		97.99	0.37	MLZ 1	99	
Bifidobacterium longum subsp. longum		1.41	0.17	BGL 10	1	
Lactobacillus fermentum		0.60	0.18		1	
01501520	**Very good identification**					**01501520**
					COC	
Blautia hansenii		99.76	0.71	NAG 10	99	
Lactobacillus fermentum		0.20	0.44		1	
Bifidobacterium longum subsp. longum		0.03	0.23		1	
01501521	**Good identification**					**01501521**
Blautia hansenii		100.0	0.37	NAG 10		
01501522	**Good identification**					**01501522**
					COC	
Blautia hansenii		97.91	0.54	NAG 10	99	
Lactobacillus fermentum		1.79	0.44	BGL 10	1	
Bifidobacterium longum subsp. longum		0.30	0.23		1	
01501530	**Acceptable identification**					**01501530**
					COC	
Blautia hansenii		94.55	0.37	MLZ 1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		5.45	0.27	NAG 10	1		
01501540	**Good identification**						**01501540**
					COC		
Blautia hansenii		99.93	0.53	SOR 1	99		
Lactobacillus fermentum		0.07	0.18		1		
01501542	**Good identification**						**01501542**
					COC		
Blautia hansenii		99.40	0.37	SOR 1	99		
Lactobacillus fermentum		0.61	0.18	BGL 10	1		
01501560	**Good identification**						**01501560**
Blautia hansenii		100.0	0.37	NAG 10			
01501700	**Good identification**						**01501700**
					COC		
Blautia hansenii		99.93	0.53	RHA 1	99		
Lactobacillus fermentum		0.07	0.18		1		
01501702	**Good identification**						**01501702**
					COC		
Blautia hansenii		99.40	0.37	RHA 1	99		
Lactobacillus fermentum		0.61	0.18	BGL 10	1		
01501720	**Good identification**						**01501720**
Blautia hansenii		100.0	0.37	RHA 1			
01502000	**Low discrimination**						**01502000**
					COC	GRA	
Megasphaera elsdenii		40.04	0.41	RAF 1	99	1	
Peptostreptococcus anaerobius		25.58	0.29	FRU 95	99	99	
Prevotella disiens		14.10	0.23		1	1	
01502040	**Low discrimination**						**01502040**
					COC	GRA	
Megasphaera elsdenii		82.26	0.38	RAF 1	99	1	
Clostridium novyi biovar A		8.62	0.29	FRU 95	1	99	
Clostridium botulinum biovar A		8.32	0.17		1	99	
01503100	**Very good identification**						**01503100**
					COC		
Blautia hansenii		99.98	0.66	MAN 1	99		
Lactobacillus fermentum		0.01	0.18		1		
Propionibacterium propionicum		0.01	0.12		1		
01503102	**Good identification**						**01503102**
					COC		
Blautia hansenii		99.81	0.50	MAN 1	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		
Propionibacterium propionicum		0.08	0.15		1		
01503120	**Good identification**						**01503120**
					COC	GRA	
Blautia hansenii		99.88	0.50	MAN 1	99	99	
Prevotella melaninogenica		0.12	-.01	NAG 10	1	1	
01503500	**Good identification**						**01503500**
					COC		
Blautia hansenii		99.93	0.53	MAN 1	99		
Lactobacillus fermentum		0.07	0.18		1		
01503502	**Good identification**						**01503502**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Blautia hansenii		99.40	0.37	MAN 1	99		
Lactobacillus fermentum		0.61	0.18	BGL 10	1		
01503520	**Good identification**						**01503520**
Blautia hansenii		100.0	0.37	MAN 1			
01504040	**Good identification**						**01504040**
Clostridium glycolicum		97.74	0.50	RAF 1			
Clostridium novyi biovar A		1.05	0.29	FRU 90			
Clostridium botulinum biovar A		1.01	0.17				
01504100	**Good identification**						**01504100**
					COC		
Blautia hansenii		99.42	0.53	GAL 99	99		
Actinomyces meyeri		0.50	0.22		1		
Lactobacillus fermentum		0.04	0.14		1		
01504102	**Good identification**						**01504102**
					COC		
Blautia hansenii		99.64	0.37	GAL 99	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
01504120	**Good identification**						**01504120**
					COC	GRA	
Blautia hansenii		98.10	0.37	GAL 99	99	99	
Prevotella bivia		1.30	-.03	NAG 10	1	1	
Actinomyces meyeri		0.61	0.07		1	99	
01504500	**Good identification**						**01504500**
					COC		
Blautia hansenii		99.07	0.40	GAL 99	99		
Actinomyces meyeri		0.71	0.11		1		
Lactobacillus fermentum		0.22	0.14		1		
01505000	**Good identification**						**01505000**
					COC	SPO	
Blautia hansenii		96.94	0.53	LAC 99	99	1	
Lactobacillus fermentum		2.34	0.44		1	1	
Clostridium glycolicum		0.23	0.11		1	99	
01505002	**Low discrimination**						**01505002**
					COC		
Blautia hansenii		82.18	0.37	LAC 99	99		
Lactobacillus fermentum		17.82	0.44	BGL 10	1		
01505020	**Acceptable identification**						**01505020**
					COC		
Blautia hansenii		93.26	0.37	LAC 99	99		
Lactobacillus fermentum		6.74	0.36	NAG 10	1		
01505100	**Very good identification**						**01505100**
					COC		
Blautia hansenii		99.95	0.87		99		
Lactobacillus fermentum		0.04	0.48		1		
Actinomyces meyeri		0.01	0.25		1		
01505101	**Good identification**						**01505101**
					COC		
Blautia hansenii		99.96	0.53	URE 1	99		
Lactobacillus fermentum		0.04	0.14		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01505102	**Very good identification**						**01505102**
					COC		
Blautia hansenii		99.64	0.71	BGL 10	99		
Lactobacillus fermentum		0.36	0.48		1		
01505103	**Good identification**						**01505103**
					COC		
Blautia hansenii		99.64	0.37	URE 1	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
01505110	**Good identification**						**01505110**
					COC		
Blautia hansenii		99.96	0.53	MLZ 1	99		
Lactobacillus fermentum		0.04	0.14		1		
01505112	**Good identification**						**01505112**
					COC		
Blautia hansenii		99.64	0.37	MLZ 1	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
01505120	**Very good identification**						**01505120**
					COC		
Blautia hansenii		99.87	0.71	NAG 10	99		
Lactobacillus fermentum		0.12	0.40		1		
Actinomyces meyeri		0.01	0.10		1		
01505121	**Good identification**						**01505121**
Blautia hansenii		100.0	0.37	NAG 10			
01505122	**Good identification**						**01505122**
					COC		
Blautia hansenii		98.94	0.54	NAG 10	99		
Lactobacillus fermentum		1.06	0.40	BGL 10	1		
01505130	**Good identification**						**01505130**
Blautia hansenii		100.0	0.37	MLZ 1			
01505140	**Good identification**						**01505140**
					COC	SPO	
Blautia hansenii		99.91	0.53	SOR 1	99	1	
Clostridium glycolicum		0.05	-.01		1	99	
Lactobacillus fermentum		0.04	0.14		1	1	
01505142	**Good identification**						**01505142**
					COC		
Blautia hansenii		99.64	0.37	SOR 1	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
01505160	**Good identification**						**01505160**
Blautia hansenii		100.0	0.37	NAG 10			
01505300	**Good identification**						**01505300**
					COC		
Blautia hansenii		99.96	0.53	RHA 1	99		
Lactobacillus fermentum		0.04	0.14		1		
01505302	**Good identification**						**01505302**
					COC		
Blautia hansenii		99.64	0.37	RHA 1	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
01505320	**Good identification**						**01505320**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Blautia hansenii		100.0	0.37	RHA 1		
01505400	**Low discrimination**					**01505400**
					COC	
Blautia hansenii		87.78	0.40	LAC 99	99	
Lactobacillus fermentum		11.99	0.44		1	
Actinomyces meyeri		0.24	0.04		1	
01505402	**Low discrimination**					**01505402**
					COC	
Lactobacillus fermentum		55.13	0.44	SUC 87	1	
Blautia hansenii		44.87	0.24	FRU 99	99	
01505422	**Low discrimination**					**01505422**
					COC	
Lactobacillus fermentum		78.66	0.36	SUC 87	1	
Blautia hansenii		21.34	0.07	FRU 99	99	
01505500	**Very good identification**					**01505500**
					COC	
Blautia hansenii		99.75	0.74		99	
Lactobacillus fermentum		0.22	0.48		1	
Bifidobacterium longum subsp. longum		0.02	0.21		1	
01505501	**Good identification**					**01505501**
					COC	
Blautia hansenii		99.78	0.40	URE 1	99	
Lactobacillus fermentum		0.23	0.14		1	
01505502	**Very good identification**					**01505502**
					COC	
Blautia hansenii		97.88	0.58	BGL 10	99	
Lactobacillus fermentum		1.98	0.48		1	
Bifidobacterium longum subsp. longum		0.14	0.21		1	
01505510	**Good identification**					**01505510**
					COC	
Blautia hansenii		97.14	0.40	MLZ 1	99	
Bifidobacterium longum subsp. longum		2.65	0.25		1	
Lactobacillus fermentum		0.22	0.14		1	
01505520	**Very good identification**					**01505520**
					COC	
Blautia hansenii		98.76	0.58	NAG 10	99	
Lactobacillus fermentum		0.67	0.40		1	
Bifidobacterium longum subsp. longum		0.57	0.31		1	
01505522	**Low discrimination**					**01505522**
					COC	
Blautia hansenii		89.84	0.42	NAG 10	99	
Lactobacillus fermentum		5.46	0.40	BGL 10	1	
Bifidobacterium longum subsp. longum		4.70	0.31		1	
01505532	**Low discrimination**					**01505532**
					COC	
Bifidobacterium longum subsp. longum		89.82	0.35	SUC 99	1	
Blautia hansenii		10.18	0.07	MNS 87	99	
01505540	**Good identification**					**01505540**
					COC	
Blautia hansenii		99.78	0.40	SOR 1	99	
Lactobacillus fermentum		0.23	0.14		1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01505700	**Good identification**							**01505700**
					COC			
Blautia hansenii		99.78	0.40	RHA 1	99			
Lactobacillus fermentum		0.23	0.14		1			
01507100	**Good identification**							**01507100**
					COC			
Blautia hansenii		99.96	0.53	MAN 1	99			
Lactobacillus fermentum		0.04	0.14		1			
01507102	**Good identification**							**01507102**
					COC			
Blautia hansenii		99.64	0.37	MAN 1	99			
Lactobacillus fermentum		0.36	0.14	BGL 10	1			
01507120	**Good identification**							**01507120**
Blautia hansenii		100.0	0.37	MAN 1				
01507500	**Good identification**							**01507500**
					COC			
Blautia hansenii		99.78	0.40	MAN 1	99			
Lactobacillus fermentum		0.23	0.14		1			
01510000	**Low discrimination**							**01510000**
					COC			
Prevotella intermedia		70.17	0.76		1			
Prevotella disiens		16.39	0.57		1			
Megasphaera elsdenii		8.93	0.63		99			
01510001	**Low discrimination**							**01510001**
					GRA	SPO		
Prevotella intermedia		35.24	0.42	URE 1	1	1		
Clostridium novyi biovar A		26.96	0.50		99	99		
Clostridium sordellii		23.42	0.32		99	99		
01510002	**Low discrimination**							**01510002**
					COC	GRA	SPO	
Megasphaera elsdenii		35.36	0.47	RAF 1	99	1	1	
Prevotella intermedia		25.25	0.42	BGL 10	1	1	1	
Clostridium sporogenes		12.05	0.32		1	99	99	
01510003	**Low discrimination**							**01510003**
					COC	GRA	SPO	
Clostridium novyi biovar A		77.09	0.37	RAF 1	1	99	99	
Megasphaera elsdenii		8.08	0.12	FRU 10	99	1	1	
Prevotella intermedia		5.77	0.08		1	1	1	
01510010	**Low discrimination**							**01510010**
					GRA	SPO		
Prevotella intermedia		45.53	0.42	MLZ 1	1	1		
Clostridium novyi biovar A		34.84	0.50		99	99		
Prevotella disiens		10.64	0.23		1	1		
01510011	**Good identification**							**01510011**
					GRA	SPO		
Clostridium novyi biovar A		97.85	0.50	RAF 1	99	99		
Prevotella intermedia		1.29	0.08	FRU 10	1	1		
Clostridium sordellii		0.86	-.03		99	99		
01510012	**Low discrimination**							**01510012**
					COC	GRA	SPO	
Clostridium novyi biovar A		80.16	0.37	RAF 1	1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		8.40	0.12	FRU 10	99	1	1	
Prevotella intermedia		6.00	0.08		1	1	1	
01510013	**Good identification**							**01510013**
Clostridium novyi biovar A		100.0	0.37	RAF 1				
01510020	**Low discrimination**							**01510020**
					GRA	SPO		
Prevotella intermedia		49.14	0.42	NAG 1	1	1		
Prevotella disiens		11.48	0.23		1	1		
Clostridium novyi biovar A		9.40	0.39		99	99		
01510021	**Acceptable identification**							**01510021**
					GRA	SPO		
Clostridium novyi biovar A		91.92	0.39	RAF 1	99	99		
Prevotella intermedia		4.86	0.08	FRU 10	1	1		
Clostridium sordellii		3.23	-.03		99	99		
01510022	**Low discrimination**							**01510022**
					SPO			
Lactobacillus fermentum		56.60	0.40	SUC 87	1			
Clostridium novyi biovar A		13.35	0.26	GAL 99	99			
Atopobium minutum		9.82	0.20		1			
01510030	**Acceptable identification**							**01510030**
					GRA	SPO		
Clostridium novyi biovar A		94.98	0.39	RAF 1	99	99		
Prevotella intermedia		5.02	0.08	FRU 10	1	1		
01510031	**Good identification**							**01510031**
Clostridium novyi biovar A		100.0	0.39	RAF 1				
01510040	**Low discrimination**							**01510040**
					COC	GRA	SPO	
Megasphaera elsdenii		66.02	0.60	RAF 1	99	1	1	
Clostridium glycolicum		16.09	0.44		1	99	99	
Prevotella intermedia		7.86	0.42		1	1	1	
01510041	**Low discrimination**							**01510041**
					COC	GRA		
Clostridium novyi biovar A		78.06	0.47	RAF 1	1	99		
Megasphaera elsdenii		12.99	0.26	FRU 10	99	1		
Clostridium sordellii		4.24	0.08		1	99		
01510042	**Low discrimination**							**01510042**
					COC	SPO	GRA	
Megasphaera elsdenii		85.97	0.44	RAF 1	99	1	1	
Clostridium novyi biovar A		8.29	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.06	0.05		1	1	99	
01510050	**Low discrimination**							**01510050**
					COC	GRA		
Clostridium novyi biovar A		80.35	0.47	RAF 1	1	99		
Megasphaera elsdenii		13.37	0.26	FRU 10	99	1		
Clostridium glycolicum		3.26	0.10		1	99		
01510051	**Good identification**							**01510051**
Clostridium novyi biovar A		100.0	0.47	RAF 1				
01510060	**Low discrimination**							**01510060**
					COC	SPO	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		44.30	0.36	RAF 1	1	99	99	
Megasphaera elsdenii		29.49	0.26	FRU 10	99	1	1	
Pseudoramibacter alactolyticus		12.36	0.09		1	1	99	
01510061	**Good identification**							**01510061**
Clostridium novyi biovar A		100.0	0.36	RAF 1				
01510070	**Good identification**							**01510070**
Clostridium novyi biovar A		96.62	0.36	RAF 1				
Clostridium botulinum biovar B		3.38	-.03	FRU 10				
01510071	**Good identification**							**01510071**
Clostridium novyi biovar A		100.0	0.36	RAF 1				
01510100	**Low discrimination**							**01510100**
					GRA			
Fusobacterium mortiferum		31.25	0.56	MLT 11	1			
Prevotella intermedia		21.30	0.42		1			
Lactobacillus fermentum		14.92	0.52		99			
01510102	**Low discrimination**							**01510102**
					COC	GRA		
Lactobacillus fermentum		70.50	0.52	SUC 87	1	99		
Fusobacterium mortiferum		16.41	0.39	GAL 99	1	1		
Blautia hansenii		6.70	0.15		99	99		
01510120	**Low discrimination**							**01510120**
					GRA			
Lactobacillus fermentum		36.85	0.44	SUC 87	99			
Prevotella bivia		26.65	0.15	GAL 99	1			
Actinomyces meyeri		11.17	0.24		99			
01510122	**Acceptable identification**							**01510122**
					COC	GRA		
Lactobacillus fermentum		91.91	0.44	SUC 87	1	99		
Prevotella melaninogenica		5.18	0.03	GAL 99	1	1		
Blautia hansenii		2.91	-.01		99	99		
01510200	**Low discrimination**							**01510200**
					COC			
Prevotella intermedia		70.63	0.42	RHA 1	1			
Prevotella disiens		16.50	0.23		1			
Megasphaera elsdenii		8.99	0.29		99			
01510400	**Low discrimination**							**01510400**
					GRA			
Prevotella intermedia		52.66	0.42	ARA 1	1			
Lactobacillus fermentum		22.61	0.48		99			
Prevotella disiens		12.30	0.23		1			
01510402	**Acceptable identification**							**01510402**
					COC	GRA		
Lactobacillus fermentum		93.66	0.48	SUC 87	1	99		
Megasphaera elsdenii		3.09	0.12	GAL 99	99	1		
Prevotella intermedia		2.20	0.08		1	1		
01510420	**Acceptable identification**							**01510420**
					GRA			
Lactobacillus fermentum		90.79	0.40	SUC 87	99			
Prevotella intermedia		6.41	0.08	GAL 99	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		2.81	0.03		99		
01510422	**Good identification**						**01510422**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01510500	**Low discrimination**						**01510500**
					COC		
Lactobacillus fermentum		72.90	0.52	SUC 87	1		
Actinomyces meyeri		13.50	0.28	GAL 99	1		
Blautia hansenii		11.00	0.19		99		
01510502	**Good identification**						**01510502**
					COC		
Lactobacillus fermentum		97.80	0.52	SUC 87	1		
Blautia hansenii		1.64	0.03	GAL 99	99		
Actinomyces meyeri		0.56	0.03		1		
01510520	**Low discrimination**						**01510520**
					COC		
Lactobacillus fermentum		87.61	0.44	SUC 87	1		
Actinomyces meyeri		6.64	0.14	GAL 99	1		
Blautia hansenii		4.41	0.03		99		
01510522	**Good identification**						**01510522**
Lactobacillus fermentum		98.49	0.44	SUC 87			
Bifidobacterium longum subsp. longum		1.51	0.05	GAL 99			
01511000	**Low discrimination**						**01511000**
					GRA		
Prevotella intermedia		68.75	0.76		1		
Lactobacillus fermentum		29.52	0.82		99		
Fusobacterium mortiferum		0.66	0.52		1		
01511001	**Low discrimination**						**01511001**
					GRA	SPO	
Prevotella intermedia		60.71	0.42	URE 1	1	1	
Lactobacillus fermentum		26.07	0.48		99	1	
Clostridium sordellii		7.12	0.19		99	99	
01511002	**Very good identification**						**01511002**
					GRA		
Lactobacillus fermentum		97.11	0.82	SUC 87	99		
Prevotella intermedia		2.28	0.42		1		
Fusobacterium mortiferum		0.24	0.36		1		
01511003	**Acceptable identification**						**01511003**
					GRA	SPO	
Lactobacillus fermentum		94.48	0.48	SUC 87	99	1	
Clostridium novyi biovar A		3.30	0.20	URE 1	99	99	
Prevotella intermedia		2.22	0.08		1	1	
01511010	**Low discrimination**						**01511010**
					GRA	SPO	
Prevotella intermedia		64.78	0.42	MLZ 1	1	1	
Lactobacillus fermentum		27.82	0.48		99	1	
Clostridium novyi biovar A		5.51	0.33		99	99	
01511012	**Acceptable identification**						**01511012**
					GRA	SPO	
Lactobacillus fermentum		94.48	0.48	SUC 87	99	1	
Clostridium novyi biovar A		3.30	0.20	MLZ 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		2.22	0.08		1	1	
01511020	**Good identification**						**01511020**
					GRA		
Lactobacillus fermentum		91.73	0.74	SUC 87	99		
Prevotella intermedia		6.47	0.42		1		
Prevotella melaninogenica		0.45	0.15		1		
01511021	**Low discrimination**						**01511021**
					GRA	SPO	
Lactobacillus fermentum		82.03	0.40	SUC 87	99	1	
Clostridium novyi biovar A		12.18	0.23	URE 1	99	99	
Prevotella intermedia		5.79	0.08		1	1	
01511022	**Very good identification**						**01511022**
					GRA		
Lactobacillus fermentum		99.50	0.74	SUC 87	99		
Clostridium perfringens		0.11	0.11		99		
Prevotella melaninogenica		0.09	0.03		1		
01511023	**Good identification**						**01511023**
					SPO		
Lactobacillus fermentum		97.45	0.40	SUC 87	1		
Clostridium novyi biovar A		2.55	0.10	URE 1	99		
01511030	**Low discrimination**						**01511030**
					GRA	SPO	
Lactobacillus fermentum		82.03	0.40	SUC 87	99	1	
Clostridium novyi biovar A		12.18	0.23	MLZ 1	99	99	
Prevotella intermedia		5.79	0.08		1	1	
01511032	**Good identification**						**01511032**
					SPO		
Lactobacillus fermentum		97.45	0.40	SUC 87	1		
Clostridium novyi biovar A		2.55	0.10	MLZ 1	99		
01511040	**Low discrimination**						**01511040**
					GRA	SPO	
Prevotella intermedia		55.12	0.42	SOR 1	1	1	
Lactobacillus fermentum		23.67	0.48		99	1	
Clostridium glycolicum		12.54	0.28		99	99	
01511042	**Acceptable identification**						**01511042**
					GRA	SPO	
Lactobacillus fermentum		93.57	0.48	SUC 87	99	1	
Prevotella intermedia		2.20	0.08	SOR 1	1	1	
Clostridium novyi biovar A		2.18	0.17		99	99	
01511060	**Low discrimination**						**01511060**
					GRA	SPO	
Lactobacillus fermentum		82.35	0.40	SUC 87	99	1	
Clostridium novyi biovar A		8.15	0.20	SOR 1	99	99	
Prevotella intermedia		5.81	0.08		1	1	
01511062	**Good identification**						**01511062**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511100	**Low discrimination**						**01511100**
					COC	GRA	
Lactobacillus fermentum		52.26	0.86	SUC 87	1	99	
Blautia hansenii		44.68	0.66		99	99	
Fusobacterium mortiferum		1.66	0.59		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01511101	**Low discrimination**						**01511101**
					COC	GRA	
Lactobacillus fermentum		51.42	0.52	SUC 87	1	99	
Blautia hansenii		43.96	0.32	URE 1	99	99	
Fusobacterium mortiferum		1.63	0.25		1	1	
01511102	**Good identification**						**01511102**
					COC	GRA	
Lactobacillus fermentum		90.94	0.86	SUC 87	1	99	
Blautia hansenii		8.64	0.50		99	99	
Fusobacterium mortiferum		0.32	0.42		1	1	
01511103	**Acceptable identification**						**01511103**
					COC		
Lactobacillus fermentum		90.23	0.52	SUC 87	1		
Blautia hansenii		8.57	0.15	URE 1	99		
Actinomyces naeslundii		1.20	0.30		1		
01511110	**Low discrimination**						**01511110**
					COC	GRA	
Lactobacillus fermentum		52.29	0.52	SUC 87	1	99	
Blautia hansenii		44.71	0.32	MLZ 1	99	99	
Fusobacterium mortiferum		1.66	0.25		1	1	
01511112	**Acceptable identification**						**01511112**
					COC		
Lactobacillus fermentum		91.32	0.52	SUC 87	1		
Blautia hansenii		8.68	0.15	MLZ 1	99		
01511120	**Low discrimination**						**01511120**
					COC	GRA	
Lactobacillus fermentum		62.51	0.78	SUC 87	1	99	
Prevotella melaninogenica		18.49	0.50		1	1	
Blautia hansenii		17.82	0.50		99	99	
01511121	**Low discrimination**						**01511121**
					COC	GRA	
Lactobacillus fermentum		60.98	0.44	SUC 87	1	99	
Prevotella melaninogenica		18.04	0.15	URE 1	1	1	
Blautia hansenii		17.38	0.15		99	99	
01511122	**Good identification**						**01511122**
					COC	GRA	
Lactobacillus fermentum		90.98	0.78	SUC 87	1	99	
Prevotella melaninogenica		5.13	0.37		1	1	
Blautia hansenii		2.88	0.33		99	99	
01511123	**Low discrimination**						**01511123**
					GRA		
Lactobacillus fermentum		88.65	0.44	SUC 87	99		
Prevotella melaninogenica		5.00	0.03	URE 1	1		
Actinomyces naeslundii		3.54	0.30		99		
01511130	**Low discrimination**						**01511130**
					COC	GRA	
Lactobacillus fermentum		62.23	0.44	SUC 87	1	99	
Prevotella melaninogenica		18.41	0.15	MLZ 1	1	1	
Blautia hansenii		17.74	0.15		99	99	
01511132	**Low discrimination**						**01511132**
					COC	GRA	
Lactobacillus fermentum		87.71	0.44	SUC 87	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		4.94	0.03	MLZ 1	1	1	
Blautia hansenii		2.78	-.01		99	99	
01511140	**Low discrimination**						**01511140**
					COC	GRA	
Lactobacillus fermentum		52.15	0.52	SUC 87	1	99	
Blautia hansenii		44.59	0.32	SOR 1	99	99	
Fusobacterium mortiferum		1.66	0.25		1	1	
01511142	**Acceptable identification**						**01511142**
					COC		
Lactobacillus fermentum		90.70	0.52	SUC 87	1		
Blautia hansenii		8.62	0.15	SOR 1	99		
Propionibacterium propionicum		0.68	0.15		1		
01511160	**Low discrimination**						**01511160**
					COC	GRA	
Lactobacillus fermentum		56.11	0.44	SUC 87	1	99	
Prevotella melaninogenica		16.60	0.15	SOR 1	1	1	
Blautia hansenii		15.99	0.15		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01511162	**Low discrimination**						**01511162**
					GRA		
Lactobacillus fermentum		84.54	0.44	SUC 87	99		
Clostridium perfringens		8.02	0.14	SOR 1	99		
Prevotella melaninogenica		4.76	0.03		1		
01511200	**Low discrimination**						**01511200**
					GRA		
Prevotella intermedia		69.21	0.42	RHA 1	1		
Lactobacillus fermentum		29.72	0.48		99		
Fusobacterium mortiferum		0.66	0.18		1		
01511202	**Good identification**						**01511202**
					GRA		
Lactobacillus fermentum		97.70	0.48	SUC 87	99		
Prevotella intermedia		2.30	0.08	RHA 1	1		
01511220	**Acceptable identification**						**01511220**
					GRA		
Lactobacillus fermentum		93.41	0.40	SUC 87	99		
Prevotella intermedia		6.59	0.08	RHA 1	1		
01511222	**Good identification**						**01511222**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511300	**Low discrimination**						**01511300**
					COC	GRA	
Lactobacillus fermentum		52.29	0.52	SUC 87	1	99	
Blautia hansenii		44.71	0.32	RHA 1	99	99	
Fusobacterium mortiferum		1.66	0.25		1	1	
01511302	**Acceptable identification**						**01511302**
					COC		
Lactobacillus fermentum		91.32	0.52	SUC 87	1		
Blautia hansenii		8.68	0.15	RHA 1	99		
01511320	**Low discrimination**						**01511320**
					COC	GRA	
Lactobacillus fermentum		63.26	0.44	SUC 87	1	99	
Prevotella melaninogenica		18.71	0.15	RHA 1	1	1	
Blautia hansenii		18.03	0.15		99	99	
01511322	**Low discrimination**						**01511322**
					GRA		
Lactobacillus fermentum		71.19	0.44	SUC 87	99		
Prevotella oralis		12.22	0.15	RHA 1	1		
Parabacteroides distasonis		10.32	0.26		1		
01511400	**Very good identification**						**01511400**
					COC	GRA	
Lactobacillus fermentum		97.30	0.82	SUC 87	1	99	
Prevotella intermedia		2.29	0.42		1	1	
Blautia hansenii		0.24	0.19		99	99	
01511401	**Good identification**						**01511401**
					GRA		
Lactobacillus fermentum		97.70	0.48	SUC 87	99		
Prevotella intermedia		2.30	0.08	URE 1	1		
01511402	**Very good identification**						**01511402**
					COC	GRA	
Lactobacillus fermentum		99.95	0.82	SUC 87	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		0.03	0.03		99	99	
Prevotella intermedia		0.02	0.08		1	1	
01511403	**Good identification**						**01511403**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01511410	**Good identification**						**01511410**
					GRA		
Lactobacillus fermentum		97.70	0.48	SUC 87	99		
Prevotella intermedia		2.30	0.08	MLZ 1	1		
01511412	**Good identification**						**01511412**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01511420	**Very good identification**						**01511420**
					COC	GRA	
Lactobacillus fermentum		99.77	0.74	SUC 87	1	99	
Blautia hansenii		0.08	0.03		99	99	
Prevotella intermedia		0.07	0.08		1	1	
01511421	**Good identification**						**01511421**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511422	**Very good identification**						**01511422**
Lactobacillus fermentum		99.98	0.74	SUC 87			
Bifidobacterium longum subsp. longum		0.03	0.05				
01511423	**Good identification**						**01511423**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511430	**Good identification**						**01511430**
Lactobacillus fermentum		95.92	0.40	SUC 87			
Bifidobacterium longum subsp. longum		4.09	0.09	MLZ 1			
01511432	**Good identification**						**01511432**
Lactobacillus fermentum		95.92	0.40	SUC 87			
Bifidobacterium longum subsp. longum		4.09	0.09	MLZ 1			
01511440	**Good identification**						**01511440**
					GRA		
Lactobacillus fermentum		97.70	0.48	SUC 87	99		
Prevotella intermedia		2.30	0.08	SOR 1	1		
01511442	**Good identification**						**01511442**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01511460	**Good identification**						**01511460**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511462	**Good identification**						**01511462**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511500	**Low discrimination**						**01511500**
					COC		
Lactobacillus fermentum		86.55	0.86	SUC 87	1		
Blautia hansenii		13.06	0.53		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		0.24	0.31		1	
01511501	**Low discrimination**					**01511501**
					COC	
Lactobacillus fermentum		86.89	0.52	SUC 87	1	
Blautia hansenii		13.11	0.19	URE 1	99	
01511502	**Very good identification**					**01511502**
					COC	
Lactobacillus fermentum		98.21	0.86	SUC 87	1	
Blautia hansenii		1.65	0.37		99	
Bifidobacterium longum subsp. longum		0.13	0.29		1	
01511503	**Good identification**					**01511503**
					COC	
Lactobacillus fermentum		98.35	0.52	SUC 87	1	
Blautia hansenii		1.65	0.03	URE 1	99	
01511510	**Low discrimination**					**01511510**
					COC	
Lactobacillus fermentum		73.19	0.52	SUC 87	1	
Bifidobacterium longum subsp. longum		15.76	0.33	MLZ 1	1	
Blautia hansenii		11.05	0.19		99	
01511512	**Low discrimination**					**01511512**
					COC	
Lactobacillus fermentum		80.74	0.52	SUC 87	1	
Bifidobacterium longum subsp. longum		17.39	0.33	MLZ 1	1	
Blautia hansenii		1.35	0.03		99	
01511520	**Good identification**					**01511520**
					COC	
Lactobacillus fermentum		93.48	0.78	SUC 87	1	
Blautia hansenii		4.70	0.37		99	
Bifidobacterium longum subsp. longum		1.43	0.40		1	
01511521	**Acceptable identification**					**01511521**
					COC	
Lactobacillus fermentum		93.84	0.44	SUC 87	1	
Blautia hansenii		4.72	0.03	URE 1	99	
Bifidobacterium longum subsp. longum		1.44	0.05		1	
01511522	**Very good identification**					**01511522**
					COC	
Lactobacillus fermentum		97.85	0.78	SUC 87	1	
Bifidobacterium longum subsp. longum		1.50	0.40		1	
Blautia hansenii		0.55	0.20		99	
01511523	**Good identification**					**01511523**
Lactobacillus fermentum		98.49	0.44	SUC 87		
Bifidobacterium longum subsp. longum		1.51	0.05	URE 1		
01511530	**Low discrimination**					**01511530**
					COC	
Bifidobacterium longum subsp. longum		71.10	0.44	SUC 99	1	
Lactobacillus fermentum		27.52	0.44	MNS 87	1	
Blautia hansenii		1.38	0.03		99	
01511532	**Low discrimination**					**01511532**
					GRA	
Bifidobacterium longum subsp. longum		71.78	0.44	SUC 99	99	
Lactobacillus fermentum		27.78	0.44	MNS 87	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		0.44	0.10		1		
01511540	**Low discrimination**						**01511540**
					COC		
Lactobacillus fermentum		86.89	0.52	SUC 87	1		
Blautia hansenii		13.11	0.19	SOR 1	99		
01511542	**Good identification**						**01511542**
					COC		
Lactobacillus fermentum		98.35	0.52	SUC 87	1		
Blautia hansenii		1.65	0.03	SOR 1	99		
01511560	**Acceptable identification**						**01511560**
					COC		
Lactobacillus fermentum		93.84	0.44	SUC 87	1		
Blautia hansenii		4.72	0.03	SOR 1	99		
Bifidobacterium longum subsp. longum		1.44	0.05		1		
01511562	**Good identification**						**01511562**
Lactobacillus fermentum		98.49	0.44	SUC 87			
Bifidobacterium longum subsp. longum		1.51	0.05	SOR 1			
01511600	**Good identification**						**01511600**
					GRA		
Lactobacillus fermentum		97.70	0.48	SUC 87	99		
Prevotella intermedia		2.30	0.08	RHA 1	1		
01511602	**Good identification**						**01511602**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01511620	**Good identification**						**01511620**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511622	**Good identification**						**01511622**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01511700	**Low discrimination**						**01511700**
					COC		
Lactobacillus fermentum		86.89	0.52	SUC 87	1		
Blautia hansenii		13.11	0.19	RHA 1	99		
01511702	**Good identification**						**01511702**
					COC		
Lactobacillus fermentum		98.35	0.52	SUC 87	1		
Blautia hansenii		1.65	0.03	RHA 1	99		
01511720	**Low discrimination**						**01511720**
					COC	GRA	
Lactobacillus fermentum		83.75	0.44	SUC 87	1	99	
Bacteroides vulgatus		10.76	0.07	RHA 1	1	1	
Blautia hansenii		4.21	0.03		99	99	
01511722	**Acceptable identification**						**01511722**
					GRA		
Lactobacillus fermentum		90.56	0.44	SUC 87	99		
Parabacteroides distasonis		8.05	0.23	RHA 1	1		
Bifidobacterium longum subsp. longum		1.39	0.05		99		
01512000	**Low discrimination**						**01512000**
					COC	GRA	
Megasphaera elsdenii		85.79	0.63	RAF 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		6.81	0.42		1	1		
Pseudoramibacter alactolyticus		2.96	0.27		1	99		
01512002	**Acceptable identification**							**01512002**
					COC	CAT	GRA	
Megasphaera elsdenii		93.85	0.47	RAF 1	99	1	1	
Pseudoramibacter alactolyticus		3.23	0.11	BGL 10	1	1	99	
Propionibacterium granulosum		1.27	0.20		1	90	99	
01512040	**Very good identification**							**01512040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.27	0.60	RAF 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.17		1	1	99	
Clostridium glycolicum		0.24	0.10		1	99	99	
01512042	**Good identification**							**01512042**
					COC	GRA		
Megasphaera elsdenii		98.18	0.44	RAF 1	99	1		
Pseudoramibacter alactolyticus		1.27	0.00	BGL 10	1	99		
Propionibacterium propionicum		0.55	0.15		1	99		
01512100	**Low discrimination**							**01512100**
					COC	GRA		
Propionibacterium propionicum		66.50	0.46	NIT 86	1	99		
Megasphaera elsdenii		24.37	0.29	SUC 99	99	1		
Fusobacterium mortiferum		2.84	0.22		1	1		
01512102	**Good identification**							**01512102**
					COC	GRA		
Propionibacterium propionicum		96.09	0.49	NIT 86	1	99		
Megasphaera elsdenii		2.61	0.12	SUC 99	99	1		
Lactobacillus fermentum		1.31	0.18		1	99		
01512140	**Low discrimination**							**01512140**
					COC	GRA		
Propionibacterium propionicum		80.37	0.46	NIT 86	1	99		
Megasphaera elsdenii		19.63	0.26	SUC 99	99	1		
01512142	**Good identification**							**01512142**
					COC	GRA		
Propionibacterium propionicum		98.22	0.49	NIT 86	1	99		
Megasphaera elsdenii		1.78	0.09	SUC 99	99	1		
01513000	**Low discrimination**							**01513000**
					COC	GRA		
Prevotella intermedia		60.22	0.42	MAN 1	1	1		
Lactobacillus fermentum		25.86	0.48		1	99		
Megasphaera elsdenii		7.67	0.29		99	1		
01513002	**Low discrimination**							**01513002**
					COC	CAT	GRA	
Lactobacillus fermentum		89.06	0.48	SUC 87	1	1	99	
Propionibacterium granulosum		3.94	0.20	MAN 1	1	90	99	
Megasphaera elsdenii		2.93	0.12		99	1	1	
01513020	**Low discrimination**							**01513020**
					GRA	CAT		
Lactobacillus fermentum		81.10	0.40	SUC 87	99	1		
Propionibacterium granulosum		10.77	0.20	MAN 1	99	90		
Prevotella intermedia		5.72	0.08		1	1		
01513022	**Acceptable identification**							**01513022**
					CAT			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		94.09	0.40	SUC 87	1		
Propionibacterium granulosum		3.13	0.10	MAN 1	90		
Actinomyces naeslundii		2.79	0.24		25		
01513100	**Low discrimination**						**01513100**
					COC		
Lactobacillus fermentum		41.23	0.52	SUC 87	1		
Blautia hansenii		35.25	0.32	MAN 1	99		
Propionibacterium propionicum		20.44	0.46		1		
01513102	**Low discrimination**						**01513102**
					COC		
Lactobacillus fermentum		53.88	0.52	SUC 87	1		
Propionibacterium propionicum		40.08	0.49	MAN 1	1		
Blautia hansenii		5.12	0.15		99		
01513120	**Low discrimination**						**01513120**
					COC	GRA	
Lactobacillus fermentum		60.70	0.44	SUC 87	1	99	
Prevotella melaninogenica		17.95	0.15	MAN 1	1	1	
Blautia hansenii		17.30	0.15		99	99	
01513122	**Low discrimination**						**01513122**
					GRA		
Lactobacillus fermentum		86.04	0.44	SUC 87	99		
Prevotella melaninogenica		4.85	0.03	MAN 1	1		
Actinomyces naeslundii		4.45	0.32		99		
01513140	**Good identification**						**01513140**
					COC		
Propionibacterium propionicum		96.36	0.46	NIT 86	1		
Lactobacillus fermentum		1.96	0.18	SUC 99	1		
Blautia hansenii		1.68	-.03		99		
01513142	**Good identification**						**01513142**
Propionibacterium propionicum		98.66	0.49	NIT 86			
Lactobacillus fermentum		1.34	0.18	SUC 99			
01513400	**Good identification**						**01513400**
					GRA		
Lactobacillus fermentum		97.70	0.48	SUC 87	99		
Prevotella intermedia		2.30	0.08	MAN 1	1		
01513402	**Good identification**						**01513402**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01513420	**Good identification**						**01513420**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01513422	**Good identification**						**01513422**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01513500	**Low discrimination**						**01513500**
					COC		
Lactobacillus fermentum		86.51	0.52	SUC 87	1		
Blautia hansenii		13.05	0.19	MAN 1	99		
Propionibacterium propionicum		0.43	0.12		1		
01513502	**Good identification**						**01513502**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		97.63	0.52	SUC 87	1			
Blautia hansenii		1.64	0.03	MAN 1	99			
Propionibacterium propionicum		0.73	0.15		1			
01513520	**Acceptable identification**							**01513520**
					COC			
Lactobacillus fermentum		93.84	0.44	SUC 87	1			
Blautia hansenii		4.72	0.03	MAN 1	99			
Bifidobacterium longum subsp. longum		1.44	0.05		1			
01513522	**Good identification**							**01513522**
Lactobacillus fermentum		98.49	0.44	SUC 87				
Bifidobacterium longum subsp. longum		1.51	0.05	MAN 1				
01514000	**Low discrimination**							**01514000**
					GRA	SPO		
Clostridium glycolicum		50.76	0.44	RAF 1	99	99		
Prevotella intermedia		24.78	0.42	SOR 95	1	1		
Lactobacillus fermentum		6.25	0.44		99	1		
01514002	**Low discrimination**							**01514002**
					COC	SPO	GRA	
Lactobacillus fermentum		80.06	0.44	SUC 87	1	1	99	
Clostridium glycolicum		6.57	0.10	GAL 99	1	99	99	
Megasphaera elsdenii		4.49	0.12		99	1	1	
01514020	**Low discrimination**							**01514020**
					SPO			
Lactobacillus fermentum		56.34	0.36	SUC 87	1			
Actinomyces meyeri		23.02	0.19	GAL 99	1			
Clostridium glycolicum		13.87	0.10		99			
01514022	**Good identification**							**01514022**
Lactobacillus fermentum		100.0	0.36	SUC 87				
01514040	**Very good identification**							**01514040**
					COC	GRA	SPO	
Clostridium glycolicum		99.74	0.66	RAF 1	1	99	99	
Megasphaera elsdenii		0.22	0.26		99	1	1	
Prevotella intermedia		0.03	0.08		1	1	1	
01514100	**Low discrimination**							**01514100**
					GRA			
Actinomyces meyeri		60.48	0.44	SUC 93	99			
Lactobacillus fermentum		24.70	0.48	RAF 1	99			
Fusobacterium mortiferum		6.63	0.36		1			
01514102	**Low discrimination**							**01514102**
					GRA			
Lactobacillus fermentum		88.18	0.48	SUC 87	99			
Actinomyces meyeri		6.68	0.18	GAL 99	99			
Fusobacterium mortiferum		2.63	0.20		1			
01514122	**Good identification**							**01514122**
Lactobacillus fermentum		97.00	0.40	SUC 87				
Actinomyces meyeri		3.01	0.03	GAL 99				
01514400	**Low discrimination**							**01514400**
					SPO			
Lactobacillus fermentum		72.90	0.44	SUC 87	1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		18.20	0.23	GAL 99	1	
Clostridium glycolicum		5.98	0.10		99	
01514402	**Good identification**					**01514402**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01514420	**Acceptable identification**					**01514420**
Lactobacillus fermentum		90.73	0.36	SUC 87		
Actinomyces meyeri		9.27	0.08	GAL 99		
01514422	**Good identification**					**01514422**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01514500	**Low discrimination**					**01514500**
					COC	
Lactobacillus fermentum		60.33	0.48	SUC 87	1	
Actinomyces meyeri		36.93	0.33	GAL 99	1	
Blautia hansenii		2.74	0.06		99	
01514502	**Good identification**					**01514502**
Lactobacillus fermentum		98.14	0.48	SUC 87		
Actinomyces meyeri		1.86	0.07	GAL 99		
01514520	**Low discrimination**					**01514520**
Lactobacillus fermentum		75.26	0.40	SUC 87		
Actinomyces meyeri		18.85	0.19	GAL 99		
Bifidobacterium longum subsp. longum		5.89	0.14			
01514522	**Acceptable identification**					**01514522**
Lactobacillus fermentum		92.74	0.40	SUC 87		
Bifidobacterium longum subsp. longum		7.26	0.14	GAL 99		
01515000	**Good identification**					**01515000**
					GRA	
Lactobacillus fermentum		93.29	0.78	SUC 87	99	
Prevotella intermedia		3.74	0.42		1	
Actinomyces meyeri		1.41	0.36		99	
01515001	**Acceptable identification**					**01515001**
					GRA	
Lactobacillus fermentum		94.77	0.44	SUC 87	99	
Prevotella intermedia		3.80	0.08	URE 1	1	
Actinomyces meyeri		1.43	0.02		99	
01515002	**Very good identification**					**01515002**
					COC	
Lactobacillus fermentum		99.83	0.78	SUC 87	1	
Actinomyces meyeri		0.05	0.11		1	
Blautia hansenii		0.05	0.03		99	
01515003	**Good identification**					**01515003**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01515010	**Acceptable identification**					**01515010**
					GRA	
Lactobacillus fermentum		94.77	0.44	SUC 87	99	
Prevotella intermedia		3.80	0.08	MLZ 1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		1.43	0.02		99		
01515012	**Good identification**						**01515012**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01515020	**Very good identification**						**01515020**
					COC		
Lactobacillus fermentum		99.13	0.70	SUC 87	1		
Actinomyces meyeri		0.61	0.22		1		
Blautia hansenii		0.14	0.03		99		
01515021	**Good identification**						**01515021**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515022	**Very good identification**						**01515022**
Lactobacillus fermentum		100.0	0.70	SUC 87			
01515023	**Good identification**						**01515023**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515030	**Good identification**						**01515030**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515032	**Good identification**						**01515032**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515040	**Acceptable identification**						**01515040**
					GRA	SPO	
Clostridium glycolicum		94.20	0.50	RAF 1	99	99	
Lactobacillus fermentum		5.50	0.44	GAL 10	99	1	
Prevotella intermedia		0.22	0.08		1	1	
01515042	**Low discrimination**						**01515042**
					SPO		
Lactobacillus fermentum		85.24	0.44	SUC 87	1		
Clostridium glycolicum		14.76	0.15	SOR 1	99		
01515060	**Low discrimination**						**01515060**
					SPO		
Lactobacillus fermentum		65.81	0.36	SUC 87	1		
Clostridium glycolicum		34.19	0.15	SOR 1	99		
01515062	**Good identification**						**01515062**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515100	**Low discrimination**						**01515100**
					COC		
Lactobacillus fermentum		77.02	0.82	SUC 87	1		
Blautia hansenii		19.79	0.53		99		
Actinomyces meyeri		2.86	0.47		1		
01515101	**Low discrimination**						**01515101**
					COC		
Lactobacillus fermentum		77.28	0.48	SUC 87	1		
Blautia hansenii		19.86	0.19	URE 1	99		
Actinomyces meyeri		2.87	0.13		1		
01515102	**Very good identification**						**01515102**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		97.07	0.82	SUC 87	1		
Blautia hansenii		2.77	0.37		99		
Actinomyces meyeri		0.11	0.21		1		
01515103	**Good identification**						**01515103**
					COC		
Lactobacillus fermentum		97.23	0.48	SUC 87	1		
Blautia hansenii		2.78	0.03	URE 1	99		
01515110	**Low discrimination**						**01515110**
					COC		
Lactobacillus fermentum		76.62	0.48	SUC 87	1		
Blautia hansenii		19.69	0.19	MLZ 1	99		
Actinomyces meyeri		2.84	0.13		1		
01515112	**Good identification**						**01515112**
					COC		
Lactobacillus fermentum		95.18	0.48	SUC 87	1		
Blautia hansenii		2.72	0.03	MLZ 1	99		
Bifidobacterium longum subsp. longum		1.06	0.07		1		
01515120	**Good identification**						**01515120**
					COC		
Lactobacillus fermentum		90.29	0.74	SUC 87	1		
Blautia hansenii		7.73	0.37		99		
Actinomyces meyeri		1.37	0.32		1		
01515121	**Acceptable identification**						**01515121**
					COC		
Lactobacillus fermentum		92.11	0.40	SUC 87	1		
Blautia hansenii		7.89	0.03	URE 1	99		
01515122	**Very good identification**						**01515122**
					COC	GRA	
Lactobacillus fermentum		97.89	0.74	SUC 87	1	99	
Blautia hansenii		0.93	0.20		99	99	
Parabacteroides distasonis		0.73	0.34		1	1	
01515123	**Good identification**						**01515123**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01515130	**Low discrimination**						**01515130**
					COC		
Lactobacillus fermentum		82.04	0.40	SUC 87	1		
Bifidobacterium longum subsp. longum		10.94	0.18	MLZ 1	1		
Blautia hansenii		7.03	0.03		99		
01515132	**Low discrimination**						**01515132**
					GRA		
Lactobacillus fermentum		65.27	0.40	SUC 87	99		
Parabacteroides distasonis		26.03	0.30	MLZ 1	1		
Bifidobacterium longum subsp. longum		8.70	0.18		99		
01515140	**Low discrimination**						**01515140**
					COC	SPO	
Lactobacillus fermentum		71.42	0.48	SUC 87	1	1	
Blautia hansenii		18.35	0.19	SOR 1	99	1	
Clostridium glycolicum		7.58	0.15		1	99	
01515142	**Good identification**						**01515142**
					COC		
Lactobacillus fermentum		97.23	0.48	SUC 87	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		2.78	0.03	SOR 1	99		
01515160	**Acceptable identification**						**01515160**
					COC		
Lactobacillus fermentum		92.11	0.40	SUC 87	1		
Blautia hansenii		7.89	0.03	SOR 1	99		
01515162	**Good identification**						**01515162**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01515200	**Acceptable identification**						**01515200**
					GRA		
Lactobacillus fermentum		94.77	0.44	SUC 87	99		
Prevotella intermedia		3.80	0.08	RHA 1	1		
Actinomyces meyeri		1.43	0.02		99		
01515202	**Good identification**						**01515202**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01515220	**Good identification**						**01515220**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515222	**Good identification**						**01515222**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515300	**Low discrimination**						**01515300**
					COC		
Lactobacillus fermentum		77.28	0.48	SUC 87	1		
Blautia hansenii		19.86	0.19	RHA 1	99		
Actinomyces meyeri		2.87	0.13		1		
01515302	**Good identification**						**01515302**
					COC		
Lactobacillus fermentum		97.23	0.48	SUC 87	1		
Blautia hansenii		2.78	0.03	RHA 1	99		
01515320	**Low discrimination**						**01515320**
					COC	GRA	
Lactobacillus fermentum		52.85	0.40	SUC 87	1	99	
Bacteroides vulgatus		42.63	0.17	RHA 1	1	1	
Blautia hansenii		4.53	0.03		99	99	
01515322	**Low discrimination**						**01515322**
					GRA		
Parabacteroides distasonis		68.20	0.43	MNS 98	1		
Lactobacillus fermentum		30.69	0.40		99		
Prevotella oralis		1.11	-.01		1		
01515332	**Good identification**						**01515332**
Parabacteroides distasonis		100.0	0.38	MNS 98			
01515400	**Very good identification**						**01515400**
					COC		
Lactobacillus fermentum		99.51	0.78	SUC 87	1		
Actinomyces meyeri		0.38	0.26		1		
Blautia hansenii		0.07	0.06		99		
01515401	**Good identification**						**01515401**
Lactobacillus fermentum		100.0	0.44	SUC 87			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01515402	**Very good identification**					**01515402**
Lactobacillus fermentum		100.0	0.78	SUC 87		
01515403	**Good identification**					**01515403**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01515410	**Good identification**					**01515410**
Lactobacillus fermentum		98.22	0.44	SUC 87		
Bifidobacterium longum subsp. longum		1.78	0.07	MLZ 1		
01515412	**Good identification**					**01515412**
Lactobacillus fermentum		98.22	0.44	SUC 87		
Bifidobacterium longum subsp. longum		1.78	0.07	MLZ 1		
01515420	**Very good identification**					**01515420**
Lactobacillus fermentum		99.72	0.70	SUC 87		
Actinomyces meyeri		0.15	0.11			
Bifidobacterium longum subsp. longum		0.13	0.14			
01515421	**Good identification**					**01515421**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01515422	**Very good identification**					**01515422**
Lactobacillus fermentum		99.87	0.70	SUC 87		
Bifidobacterium longum subsp. longum		0.13	0.14			
01515423	**Good identification**					**01515423**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01515430	**Low discrimination**					**01515430**
Lactobacillus fermentum		82.13	0.36	SUC 87		
Bifidobacterium longum subsp. longum		17.87	0.18	MLZ 1		
01515432	**Low discrimination**					**01515432**
Lactobacillus fermentum		82.13	0.36	SUC 87		
Bifidobacterium longum subsp. longum		17.87	0.18	MLZ 1		
01515440	**Low discrimination**					**01515440**
					SPO	
Lactobacillus fermentum		85.24	0.44	SUC 87	1	
Clostridium glycolicum		14.76	0.15	SOR 1	99	
01515442	**Good identification**					**01515442**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01515460	**Good identification**					**01515460**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01515462	**Good identification**					**01515462**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01515500	**Good identification**					**01515500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Lactobacillus fermentum		94.24	0.82	SUC 87	1	
Blautia hansenii		4.27	0.40		99	
Actinomyces meyeri		0.87	0.36		1	
01515501	**Acceptable identification**					**01515501**
					COC	
Lactobacillus fermentum		94.82	0.48	SUC 87	1	
Blautia hansenii		4.30	0.06	URE 1	99	
Actinomyces meyeri		0.88	0.02		1	
01515502	**Very good identification**					**01515502**
					COC	
Lactobacillus fermentum		98.77	0.82	SUC 87	1	
Bifidobacterium longum subsp. longum		0.64	0.37		1	
Blautia hansenii		0.50	0.24		99	
01515503	**Good identification**					**01515503**
Lactobacillus fermentum		100.0	0.48	SUC 87		
01515510	**Low discrimination**					**01515510**
					COC	
Bifidobacterium longum subsp. longum		51.05	0.41	SUC 99	1	
Lactobacillus fermentum		46.41	0.48	MNS 87	1	
Blautia hansenii		2.10	0.06		99	
01515512	**Low discrimination**					**01515512**
Bifidobacterium longum subsp. longum		50.61	0.41	SUC 99		
Lactobacillus fermentum		46.01	0.48	MNS 87		
Bifidobacterium adolescentis		3.38	0.23			
01515520	**Good identification**					**01515520**
					COC	
Lactobacillus fermentum		90.15	0.74	SUC 87	1	
Bifidobacterium longum subsp. longum		7.06	0.48		1	
Blautia hansenii		1.36	0.24		99	
01515521	**Acceptable identification**					**01515521**
Lactobacillus fermentum		92.74	0.40	SUC 87		
Bifidobacterium longum subsp. longum		7.26	0.14	URE 1		
01515522	**Good identification**					**01515522**
					GRA	
Lactobacillus fermentum		92.20	0.74	SUC 87	99	
Bifidobacterium longum subsp. longum		7.22	0.48		99	
Parabacteroides distasonis		0.42	0.31		1	
01515523	**Acceptable identification**					**01515523**
Lactobacillus fermentum		92.74	0.40	SUC 87		
Bifidobacterium longum subsp. longum		7.26	0.14	URE 1		
01515530	**Acceptable identification**					**01515530**
Bifidobacterium longum subsp. longum		92.96	0.52	SUC 99		
Lactobacillus fermentum		7.04	0.40	MNS 87		
01515532	**Acceptable identification**					**01515532**
					GRA	
Bifidobacterium longum subsp. longum		91.38	0.52	SUC 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		6.92	0.40	MNS 87	99		
Parabacteroides distasonis		1.69	0.26		1		
01515540	**Acceptable identification**						**01515540**
					COC		
Lactobacillus fermentum		94.82	0.48	SUC 87	1		
Blautia hansenii		4.30	0.06	SOR 1	99		
Actinomyces meyeri		0.88	0.02		1		
01515542	**Good identification**						**01515542**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01515560	**Acceptable identification**						**01515560**
Lactobacillus fermentum		92.74	0.40	SUC 87			
Bifidobacterium longum subsp. longum		7.26	0.14	SOR 1			
01515562	**Acceptable identification**						**01515562**
Lactobacillus fermentum		92.74	0.40	SUC 87			
Bifidobacterium longum subsp. longum		7.26	0.14	SOR 1			
01515600	**Good identification**						**01515600**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01515602	**Good identification**						**01515602**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01515620	**Low discrimination**						**01515620**
					GRA		
Lactobacillus fermentum		84.99	0.36	SUC 87	99		
Bacteroides vulgatus		15.01	0.02	RHA 1	1		
01515622	**Good identification**						**01515622**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01515700	**Acceptable identification**						**01515700**
					COC	GRA	
Lactobacillus fermentum		91.68	0.48	SUC 87	1	99	
Blautia hansenii		4.16	0.06	RHA 1	99	99	
Bacteroides vulgatus		3.31	0.02		1	1	
01515702	**Good identification**						**01515702**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01515720	**Acceptable identification**						**01515720**
					GRA		
Bacteroides vulgatus		90.86	0.36	SUC 96	1		
Lactobacillus fermentum		8.48	0.40	MNS 99	99		
Bifidobacterium longum subsp. longum		0.66	0.14		99		
01515722	**Low discrimination**						**01515722**
					GRA		
Parabacteroides distasonis		53.44	0.39	MNS 98	1		
Lactobacillus fermentum		39.24	0.40		99		
Bacteroides vulgatus		4.25	0.02		1		
01517000	**Acceptable identification**						**01517000**
					GRA		
Lactobacillus fermentum		94.77	0.44	SUC 87	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella intermedia		3.80	0.08	MAN 1	1	
Actinomyces meyeri		1.43	0.02		99	
01517002	**Good identification**					**01517002**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01517020	**Good identification**					**01517020**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01517022	**Good identification**					**01517022**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01517100	**Low discrimination**					**01517100**
					COC	
Lactobacillus fermentum		76.77	0.48	SUC 87	1	
Blautia hansenii		19.73	0.19	MAN 1	99	
Actinomyces meyeri		2.85	0.13		1	
01517102	**Good identification**					**01517102**
					COC	
Lactobacillus fermentum		96.03	0.48	SUC 87	1	
Blautia hansenii		2.74	0.03	MAN 1	99	
Propionibacterium propionicum		1.23	0.15		1	
01517120	**Acceptable identification**					**01517120**
					COC	
Lactobacillus fermentum		92.11	0.40	SUC 87	1	
Blautia hansenii		7.89	0.03	MAN 1	99	
01517122	**Good identification**					**01517122**
Lactobacillus fermentum		100.0	0.40	SUC 87		
01517400	**Good identification**					**01517400**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01517402	**Good identification**					**01517402**
Lactobacillus fermentum		100.0	0.44	SUC 87		
01517420	**Good identification**					**01517420**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01517422	**Good identification**					**01517422**
Lactobacillus fermentum		100.0	0.36	SUC 87		
01517500	**Acceptable identification**					**01517500**
					COC	
Lactobacillus fermentum		94.82	0.48	SUC 87	1	
Blautia hansenii		4.30	0.06	MAN 1	99	
Actinomyces meyeri		0.88	0.02		1	
01517502	**Good identification**					**01517502**
Lactobacillus fermentum		95.31	0.48	SUC 87		
Bifidobacterium adolescentis		4.69	0.20	MAN 1		
01517520	**Acceptable identification**					**01517520**
Lactobacillus fermentum		92.74	0.40	SUC 87		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		7.26	0.14	MAN 1			
01517522	**Acceptable identification**						**01517522**
Lactobacillus fermentum		92.74	0.40	SUC 87			
Bifidobacterium longum subsp. longum		7.26	0.14	MAN 1			
01520000	**Low discrimination**						**01520000**
					COC	SPO	
Clostridium botulinum biovar A		51.67	0.49	RAF 1	1	99	
Clostridium sporogenes		37.69	0.45		1	99	
Peptostreptococcus anaerobius		5.91	0.29		99	1	
01520002	**Acceptable identification**						**01520002**
Clostridium sporogenes		94.35	0.37	RAF 1			
Clostridium botulinum biovar A		3.92	0.15				
Clostridium novyi biovar A		1.29	0.19				
01520040	**Acceptable identification**						**01520040**
Clostridium botulinum biovar A		92.01	0.41	RAF 1			
Clostridium novyi biovar A		3.85	0.29				
Clostridium sporogenes		2.27	0.11				
01521100	**Very good identification**						**01521100**
					COC	GRA	
Blautia hansenii		99.98	0.66	TRE 1	99	99	
Lactobacillus fermentum		0.01	0.18		1	99	
Fusobacterium mortiferum		0.01	0.16		1	1	
01521102	**Good identification**						**01521102**
					COC		
Blautia hansenii		99.89	0.50	TRE 1	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		
01521120	**Good identification**						**01521120**
					COC	GRA	
Blautia hansenii		99.81	0.50	TRE 1	99	99	
Prevotella melaninogenica		0.12	-.01	NAG 10	1	1	
Clostridium perfringens		0.07	0.02		1	99	
01521500	**Good identification**						**01521500**
					COC		
Blautia hansenii		99.93	0.53	TRE 1	99		
Lactobacillus fermentum		0.07	0.18		1		
01521502	**Good identification**						**01521502**
					COC		
Blautia hansenii		99.40	0.37	TRE 1	99		
Lactobacillus fermentum		0.61	0.18	BGL 10	1		
01521520	**Good identification**						**01521520**
					COC		
Blautia hansenii		99.16	0.37	TRE 1	99		
Bifidobacterium longum subsp. longum		0.84	0.13	NAG 10	1		
01525100	**Good identification**						**01525100**
					COC		
Blautia hansenii		99.96	0.53	TRE 1	99		
Lactobacillus fermentum		0.04	0.14		1		
01525102	**Good identification**						**01525102**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		99.64	0.37	TRE 1	99			
Lactobacillus fermentum		0.36	0.14	BGL 10	1			
01525120	**Good identification**							**01525120**
Blautia hansenii		100.0	0.37	TRE 1				
01525500	**Good identification**							**01525500**
					COC			
Blautia hansenii		99.38	0.40	TRE 1	99			
Bifidobacterium longum subsp. longum		0.40	0.11		1			
Lactobacillus fermentum		0.22	0.14		1			
01530000	**Low discrimination**							**01530000**
					GRA	SPO	CAT	
Prevotella intermedia		38.62	0.42	TRE 1	1	1	1	
Clostridium sporogenes		18.43	0.32		99	99	1	
Propionibacterium granulosum		11.68	0.41		99	1	90	
01530100	**Acceptable identification**							**01530100**
					GRA			
Fusobacterium mortiferum		92.94	0.47	MLT 11	1			
Prevotella intermedia		2.14	0.08		1			
Lactobacillus fermentum		1.50	0.18		99			
01531000	**Low discrimination**							**01531000**
					GRA	CAT		
Prevotella intermedia		49.29	0.42	TRE 1	1	1		
Lactobacillus fermentum		21.17	0.48		99	1		
Propionibacterium granulosum		14.91	0.41		99	90		
01531002	**Low discrimination**							**01531002**
					GRA	CAT		
Lactobacillus fermentum		77.98	0.48	SUC 87	99	1		
Propionibacterium granulosum		13.73	0.30	TRE 1	99	90		
Fusobacterium mortiferum		5.69	0.27		1	5		
01531020	**Low discrimination**							**01531020**
					CAT			
Lactobacillus fermentum		48.41	0.40	SUC 87	1			
Propionibacterium granulosum		25.57	0.30	TRE 1	90			
Clostridium perfringens		21.16	0.22		1			
01531022	**Low discrimination**							**01531022**
					CAT			
Lactobacillus fermentum		73.13	0.40	SUC 87	1			
Clostridium perfringens		15.04	0.16	TRE 1	1			
Propionibacterium granulosum		9.66	0.20		90			
01531100	**Low discrimination**							**01531100**
					COC	GRA		
Lactobacillus fermentum		35.01	0.52	SUC 87	1	99		
Fusobacterium mortiferum		32.85	0.50	TRE 1	1	1		
Blautia hansenii		29.93	0.32		99	99		
01531102	**Low discrimination**							**01531102**
					COC	GRA		
Lactobacillus fermentum		81.16	0.52	SUC 87	1	99		
Fusobacterium mortiferum		8.46	0.33	TRE 1	1	1		
Blautia hansenii		7.71	0.15		99	99		
01531120	**Low discrimination**							**01531120**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		52.90	0.36	SUC 91	99			
Lactobacillus fermentum		26.93	0.44	MNS 99	99			
Prevotella melaninogenica		7.97	0.15		1			
01531122	**Low discrimination**							**01531122**
					GRA			
Lactobacillus fermentum		43.29	0.44	SUC 87	99			
Clostridium perfringens		40.02	0.31	TRE 1	99			
Parabacteroides distasonis		9.53	0.29		1			
01531322	**Good identification**							**01531322**
Parabacteroides distasonis		100.0	0.38	MNS 98				
01531400	**Good identification**							**01531400**
					GRA			
Lactobacillus fermentum		97.70	0.48	SUC 87	99			
Prevotella intermedia		2.30	0.08	TRE 1	1			
01531402	**Good identification**							**01531402**
Lactobacillus fermentum		100.0	0.48	SUC 87				
01531420	**Good identification**							**01531420**
Lactobacillus fermentum		100.0	0.40	SUC 87				
01531422	**Good identification**							**01531422**
Lactobacillus fermentum		100.0	0.40	SUC 87				
01531500	**Low discrimination**							**01531500**
					COC			
Lactobacillus fermentum		83.89	0.52	SUC 87	1			
Blautia hansenii		12.66	0.19	TRE 1	99			
Bifidobacterium longum subsp. longum		2.65	0.19		1			
01531502	**Good identification**							**01531502**
					COC			
Lactobacillus fermentum		95.39	0.52	SUC 87	1			
Bifidobacterium longum subsp. longum		3.02	0.19	TRE 1	1			
Blautia hansenii		1.60	0.03		99			
01531520	**Low discrimination**							**01531520**
					COC			
Lactobacillus fermentum		68.99	0.44	SUC 87	1			
Bifidobacterium longum subsp. longum		26.17	0.29	TRE 1	1			
Blautia hansenii		3.47	0.03		99			
01531522	**Low discrimination**							**01531522**
					GRA			
Lactobacillus fermentum		66.04	0.44	SUC 87	99			
Bifidobacterium longum subsp. longum		25.05	0.29	TRE 1	99			
Parabacteroides distasonis		8.91	0.26		1			
01532000	**Low discrimination**							**01532000**
					COC	GRA	CAT	
Propionibacterium granulosum		60.88	0.39	SUC 93	1	99	90	
Megasphaera elsdenii		33.97	0.29	MNS 99	99	1	1	
Prevotella intermedia		2.70	0.08		1	1	1	
01532100	**Good identification**							**01532100**
					GRA			
Propionibacterium propionicum		98.47	0.45	NIT 86	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		1.53	0.13	SUC 99	1		
01532102	**Good identification**						**01532102**
Propionibacterium propionicum		100.0	0.48	NIT 86			
01532140	**Good identification**						**01532140**
Propionibacterium propionicum		100.0	0.45	NIT 86			
01532142	**Good identification**						**01532142**
Propionibacterium propionicum		100.0	0.48	NIT 86			
01533000	**Acceptable identification**						**01533000**
					GRA	CAT	
Propionibacterium granulosum		90.96	0.39	SUC 93	99	90	
Prevotella intermedia		4.03	0.08	MNS 99	1	1	
Actinomyces naeslundii		3.28	0.29		99	25	
01533100	**Low discrimination**						**01533100**
Propionibacterium propionicum		86.80	0.45	NIT 86			
Actinomyces naeslundii		7.16	0.37	SUC 99			
Lactobacillus fermentum		2.16	0.18				
01533101	**Low discrimination**						**01533101**
Actinomyces naeslundii		52.69	0.40	SUC 99			
Propionibacterium propionicum		47.31	0.29	MNS 88			
01533102	**Acceptable identification**						**01533102**
Propionibacterium propionicum		93.32	0.48	NIT 86			
Actinomyces naeslundii		5.13	0.37	SUC 99			
Lactobacillus fermentum		1.55	0.18				
01533120	**Low discrimination**						**01533120**
Actinomyces naeslundii		83.50	0.37	SUC 99			
Clostridium perfringens		16.50	0.02	MNS 88			
01533121	**Good identification**						**01533121**
Actinomyces naeslundii		100.0	0.40	SUC 99			
01533122	**Low discrimination**						**01533122**
Actinomyces naeslundii		84.48	0.37	SUC 99			
Propionibacterium propionicum		15.52	0.14	MNS 88			
01533123	**Good identification**						**01533123**
Actinomyces naeslundii		100.0	0.40	SUC 99			
01533140	**Good identification**						**01533140**
Propionibacterium propionicum		98.57	0.45	NIT 86			
Actinomyces naeslundii		1.44	0.24	SUC 99			
01533142	**Good identification**						**01533142**
Propionibacterium propionicum		99.04	0.48	NIT 86			
Actinomyces naeslundii		0.96	0.24	SUC 99			
01535000	**Low discrimination**						**01535000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Lactobacillus fermentum		87.78	0.44	SUC 87	99		
Fusobacterium mortiferum		7.38	0.24	TRE 1	1		
Prevotella intermedia		3.52	0.08		1		
01535002	**Good identification**						**01535002**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01535020	**Good identification**						**01535020**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01535022	**Acceptable identification**						**01535022**
					GRA		
Lactobacillus fermentum		94.73	0.36	SUC 87	99		
Parabacteroides distasonis		5.27	0.12	TRE 1	1		
01535100	**Low discrimination**						**01535100**
					COC	GRA	
Lactobacillus fermentum		70.71	0.48	SUC 87	1	99	
Blautia hansenii		18.17	0.19	TRE 1	99	99	
Fusobacterium mortiferum		8.50	0.31		1	1	
01535102	**Acceptable identification**						**01535102**
					COC	GRA	
Lactobacillus fermentum		94.94	0.48	SUC 87	1	99	
Blautia hansenii		2.71	0.03	TRE 1	99	99	
Fusobacterium mortiferum		1.27	0.14		1	1	
01535120	**Low discrimination**						**01535120**
					COC	GRA	
Lactobacillus fermentum		86.69	0.40	SUC 87	1	99	
Blautia hansenii		7.42	0.03	TRE 1	99	99	
Parabacteroides distasonis		2.96	0.12		1	1	
01535122	**Low discrimination**						**01535122**
					GRA		
Parabacteroides distasonis		77.14	0.46	MNS 98	1		
Lactobacillus fermentum		22.86	0.40		99		
01535132	**Good identification**						**01535132**
					GRA		
Parabacteroides distasonis		98.20	0.41	MNS 98	1		
Bifidobacterium longum subsp. longum		1.80	0.07		99		
01535322	**Good identification**						**01535322**
Parabacteroides distasonis		100.0	0.54	MNS 98			
01535332	**Good identification**						**01535332**
Parabacteroides distasonis		100.0	0.49	MNS 98			
01535400	**Good identification**						**01535400**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01535402	**Good identification**						**01535402**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01535420	**Good identification**						**01535420**
Lactobacillus fermentum		100.0	0.36	SUC 87			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
01535422	**Good identification**						**01535422**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01535500	**Low discrimination**						**01535500**
					COC		
Lactobacillus fermentum		82.23	0.48	SUC 87	1		
Bifidobacterium longum subsp. longum		13.28	0.27	TRE 1	1		
Blautia hansenii		3.73	0.06		99		
01535502	**Low discrimination**						**01535502**
Lactobacillus fermentum		83.23	0.48	SUC 87			
Bifidobacterium longum subsp. longum		13.44	0.27	TRE 1			
Bifidobacterium adolescentis		3.33	0.18				
01535520	**Low discrimination**						**01535520**
					GRA		
Bifidobacterium longum subsp. longum		64.87	0.37	SUC 99	99		
Lactobacillus fermentum		33.47	0.40	MNS 87	99		
Bacteroides vulgatus		1.66	-.04		1		
01535522	**Low discrimination**						**01535522**
					GRA		
Parabacteroides distasonis		41.31	0.42	MNS 98	1		
Bifidobacterium longum subsp. longum		38.71	0.37		99		
Lactobacillus fermentum		19.98	0.40		99		
01535530	**Good identification**						**01535530**
Bifidobacterium longum subsp. longum		100.0	0.41	SUC 99			
01535532	**Low discrimination**						**01535532**
					GRA		
Bifidobacterium longum subsp. longum		74.77	0.41	SUC 99	99		
Parabacteroides distasonis		25.23	0.38	MNS 87	1		
01535722	**Good identification**						**01535722**
Parabacteroides distasonis		100.0	0.50	MNS 98			
01535732	**Good identification**						**01535732**
					GRA		
Parabacteroides distasonis		99.01	0.46	MNS 98	1		
Bifidobacterium longum subsp. longum		0.99	0.07		99		
01541100	**Very good identification**						**01541100**
					COC	GRA	
Blautia hansenii		99.98	0.66	CEL 1	99	99	
Lactobacillus fermentum		0.01	0.18		1	99	
Fusobacterium mortiferum		0.01	0.16		1	1	
01541102	**Good identification**						**01541102**
					COC		
Blautia hansenii		99.89	0.50	CEL 1	99		
Lactobacillus fermentum		0.11	0.18	BGL 10	1		
01541120	**Good identification**						**01541120**
					COC	GRA	
Blautia hansenii		98.74	0.50	CEL 1	99	99	
Prevotella melaninogenica		1.27	0.17	NAG 10	1	1	
01541500	**Good identification**						**01541500**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Blautia hansenii		99.93	0.53	CEL 1	99	
Lactobacillus fermentum		0.07	0.18		1	
01541502	**Good identification**					**01541502**
					COC	
Blautia hansenii		99.40	0.37	CEL 1	99	
Lactobacillus fermentum		0.61	0.18	BGL 10	1	
01541520	**Good identification**					**01541520**
Blautia hansenii		100.0	0.37	CEL 1		
01545100	**Good identification**					**01545100**
					COC	
Blautia hansenii		99.96	0.53	CEL 1	99	
Lactobacillus fermentum		0.04	0.14		1	
01545102	**Good identification**					**01545102**
					COC	
Blautia hansenii		99.64	0.37	CEL 1	99	
Lactobacillus fermentum		0.36	0.14	BGL 10	1	
01545120	**Good identification**					**01545120**
Blautia hansenii		100.0	0.37	CEL 1		
01545500	**Good identification**					**01545500**
					COC	
Blautia hansenii		99.78	0.40	CEL 1	99	
Lactobacillus fermentum		0.23	0.14		1	
01550000	**Low discrimination**					**01550000**
Prevotella intermedia		61.77	0.42	CEL 1		
Prevotella disiens		14.43	0.23			
Fusobacterium mortiferum		12.95	0.41			
01550100	**Acceptable identification**					**01550100**
					GRA	
Fusobacterium mortiferum		94.50	0.48	MLT 11	1	
Prevotella intermedia		1.95	0.08		1	
Lactobacillus fermentum		1.37	0.18		99	
01551000	**Low discrimination**					**01551000**
					GRA	
Prevotella intermedia		57.15	0.42	CEL 1	1	
Lactobacillus fermentum		24.54	0.48		99	
Fusobacterium mortiferum		17.97	0.44		1	
01551002	**Acceptable identification**					**01551002**
					GRA	
Lactobacillus fermentum		90.51	0.48	SUC 87	99	
Fusobacterium mortiferum		7.36	0.28	CEL 1	1	
Prevotella intermedia		2.13	0.08		1	
01551020	**Low discrimination**					**01551020**
					GRA	
Lactobacillus fermentum		88.95	0.40	SUC 87	99	
Prevotella intermedia		6.28	0.08	CEL 1	1	
Prevotella melaninogenica		4.77	-.01		1	
01551022	**Low discrimination**					**01551022**
					GRA	
Lactobacillus fermentum		87.12	0.40	SUC 87	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		12.88	0.10	CEL 1	1		
01551100	**Low discrimination**						**01551100**
					COC	GRA	
Fusobacterium mortiferum		35.49	0.51	MLT 11	1	1	
Lactobacillus fermentum		33.89	0.52		1	99	
Blautia hansenii		28.98	0.32		99	99	
01551102	**Low discrimination**						**01551102**
					COC	GRA	
Lactobacillus fermentum		80.13	0.52	SUC 87	1	99	
Fusobacterium mortiferum		9.32	0.34	CEL 1	1	1	
Blautia hansenii		7.61	0.15		99	99	
01551400	**Good identification**						**01551400**
					GRA		
Lactobacillus fermentum		97.70	0.48	SUC 87	99		
Prevotella intermedia		2.30	0.08	CEL 1	1		
01551402	**Good identification**						**01551402**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01551420	**Good identification**						**01551420**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01551422	**Good identification**						**01551422**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01551500	**Low discrimination**						**01551500**
					COC	GRA	
Lactobacillus fermentum		86.10	0.52	SUC 87	1	99	
Blautia hansenii		12.99	0.19	CEL 1	99	99	
Fusobacterium mortiferum		0.91	0.16		1	1	
01551502	**Good identification**						**01551502**
					COC		
Lactobacillus fermentum		97.28	0.52	SUC 87	1		
Blautia hansenii		1.63	0.03	CEL 1	99		
Bifidobacterium adolescentis		1.09	0.13		1		
01551520	**Acceptable identification**						**01551520**
					COC	GRA	
Lactobacillus fermentum		91.03	0.44	SUC 87	1	99	
Blautia hansenii		4.58	0.03	CEL 1	99	99	
Prevotella melaninogenica		2.99	-.01		1	1	
01551522	**Good identification**						**01551522**
					GRA		
Lactobacillus fermentum		96.97	0.44	SUC 87	99		
Parabacteroides distasonis		1.55	0.10	CEL 1	1		
Bifidobacterium longum subsp. longum		1.49	0.05		99		
01553101	**Good identification**						**01553101**
Actinomyces naeslundii		100.0	0.37	SUC 99			
01553103	**Good identification**						**01553103**
Actinomyces naeslundii		100.0	0.37	SUC 99			
01553121	**Good identification**						**01553121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	SUC 99			
01553123	**Good identification**						**01553123**
Actinomyces naeslundii		100.0	0.37	SUC 99			
01555000	**Low discrimination**						**01555000**
					GRA		
Lactobacillus fermentum		87.03	0.44	SUC 87	99		
Fusobacterium mortiferum		8.17	0.25	CEL 1	1		
Prevotella intermedia		3.49	0.08		1		
01555002	**Good identification**						**01555002**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01555020	**Good identification**						**01555020**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01555022	**Good identification**						**01555022**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01555100	**Low discrimination**						**01555100**
					COC	GRA	
Lactobacillus fermentum		69.42	0.48	SUC 87	1	99	
Blautia hansenii		17.84	0.19	CEL 1	99	99	
Fusobacterium mortiferum		9.32	0.31		1	1	
01555102	**Acceptable identification**						**01555102**
					COC		
Lactobacillus fermentum		92.51	0.48	SUC 87	1		
Blautia hansenii		2.64	0.03	CEL 1	99		
Bifidobacterium adolescentis		1.77	0.13		1		
01555120	**Low discrimination**						**01555120**
					COC	GRA	
Lactobacillus fermentum		87.60	0.40	SUC 87	1	99	
Blautia hansenii		7.50	0.03	CEL 1	99	99	
Prevotella melaninogenica		4.90	-.01		1	1	
01555122	**Low discrimination**						**01555122**
					GRA		
Lactobacillus fermentum		57.52	0.40	SUC 87	99		
Parabacteroides distasonis		22.94	0.30	CEL 1	1		
Prevotella oralis		17.19	0.15		1		
01555322	**Acceptable identification**						**01555322**
Parabacteroides distasonis		90.33	0.38	MNS 98			
Prevotella oralis		9.67	0.09				
01555400	**Good identification**						**01555400**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01555402	**Good identification**						**01555402**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01555420	**Good identification**						**01555420**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01555422	**Good identification**						**01555422**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		100.0	0.36	SUC 87		
01555500	**Acceptable identification**					**01555500**
					COC	
Lactobacillus fermentum		93.93	0.48	SUC 87	1	
Blautia hansenii		4.26	0.06	CEL 1	99	
Actinomyces israelii		0.94	0.14		1	
01555502	**Low discrimination**					**01555502**
Lactobacillus fermentum		87.51	0.48	SUC 87		
Bifidobacterium adolescentis		11.18	0.27	CEL 1		
Actinomyces israelii		1.32	0.17			
01555520	**Low discrimination**					**01555520**
					GRA	
Lactobacillus fermentum		87.65	0.40	SUC 87	99	
Bifidobacterium longum subsp. longum		6.86	0.14	CEL 1	99	
Bacteroides vulgatus		5.49	-.02		1	
01555522	**Low discrimination**					**01555522**
					GRA	
Lactobacillus fermentum		75.60	0.40	SUC 87	99	
Parabacteroides distasonis		18.48	0.26	CEL 1	1	
Bifidobacterium longum subsp. longum		5.92	0.14		99	
01570100	**Good identification**					**01570100**
Fusobacterium mortiferum		100.0	0.39	MLT 11		
01571000	**Low discrimination**					**01571000**
					GRA	
Fusobacterium mortiferum		82.40	0.35	MLT 11	1	
Prevotella intermedia		8.86	0.08		1	
Actinomyces naeslundii		4.94	0.26		99	
01571100	**Low discrimination**					**01571100**
					GRA	
Fusobacterium mortiferum		88.26	0.42	MLT 11	1	
Actinomyces naeslundii		6.45	0.34		99	
Lactobacillus fermentum		2.85	0.18		99	
01571101	**Good identification**					**01571101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
01571103	**Good identification**					**01571103**
Actinomyces naeslundii		100.0	0.37	SUC 99		
01571121	**Good identification**					**01571121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
01571123	**Good identification**					**01571123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
01573100	**Low discrimination**					**01573100**
Actinomyces naeslundii		76.84	0.39	SUC 99		
Actinomyces israelii		23.17	0.21	MNS 88		
01573101	**Good identification**					**01573101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	SUC 99		
01573102	**Low discrimination**					**01573102**
Actinomyces naeslundii		62.86	0.39	SUC 99		
Actinomyces israelii		28.43	0.24	MNS 88		
Propionibacterium propionicum		8.71	0.14			
01573103	**Good identification**					**01573103**
Actinomyces naeslundii		100.0	0.42	SUC 99		
01573120	**Good identification**					**01573120**
Actinomyces naeslundii		100.0	0.39	SUC 99		
01573121	**Good identification**					**01573121**
Actinomyces naeslundii		100.0	0.42	SUC 99		
01573122	**Good identification**					**01573122**
Actinomyces naeslundii		100.0	0.39	SUC 99		
01573123	**Good identification**					**01573123**
Actinomyces naeslundii		100.0	0.42	SUC 99		
01575122	**Good identification**					**01575122**
Parabacteroides distasonis		100.0	0.41	MNS 98		
01575132	**Good identification**					**01575132**
Parabacteroides distasonis		100.0	0.37	MNS 98		
01575322	**Good identification**					**01575322**
Parabacteroides distasonis		100.0	0.49	MNS 98		
01575332	**Good identification**					**01575332**
Parabacteroides distasonis		100.0	0.45	MNS 98		
01575522	**Good identification**					**01575522**
Parabacteroides distasonis		100.0	0.38	MNS 98		
01575722	**Good identification**					**01575722**
Parabacteroides distasonis		100.0	0.46	MNS 98		
01575732	**Good identification**					**01575732**
Parabacteroides distasonis		97.58	0.41	MNS 98		
Clostridium clostridioforme		2.42	0.08			
01577100	**Good identification**					**01577100**
Actinomyces israelii		100.0	0.39	SUC 98		
01577102	**Good identification**					**01577102**
Actinomyces israelii		97.42	0.42	SUC 98		
Bifidobacterium adolescentis		2.58	0.08	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
01577500	**Good identification**							**01577500**
Actinomyces israelii		100.0	0.37	SUC 98				
01577502	**Low discrimination**							**01577502**
Actinomyces israelii		82.19	0.40	SUC 98				
Bifidobacterium adolescentis		17.81	0.23	SAL 92				
01600001	**Low discrimination**							**01600001**
					COC	GRA	SPO	
Clostridium novyi biovar A		85.34	0.48	SAL 8	1	99	99	
Anaerococcus prevotii		11.95	0.15	RAF 1	99	99	1	
Campylobacter ureolyticus		1.79	-.03		1	1	1	
01600010	**Good identification**							**01600010**
					COC	SPO		
Clostridium novyi biovar A		98.94	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		1.06	-.03	RAF 1	99	1		
01600011	**Good identification**							**01600011**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01600020	**Low discrimination**							**01600020**
					SPO			
Clostridium novyi biovar A		71.03	0.37	SAL 8	99			
Atopobium minutum		19.14	0.24	RAF 1	1			
Clostridium cadaveris		4.74	0.03		99			
01600021	**Good identification**							**01600021**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01600030	**Good identification**							**01600030**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01600031	**Good identification**							**01600031**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01600040	**Low discrimination**							**01600040**
					COC	SPO		
Clostridium novyi biovar A		86.72	0.45	SAL 8	1	99		
Clostridium botulinum biovar A		11.88	0.18	RAF 1	1	99		
Peptostreptococcus anaerobius		1.40	-.03		99	1		
01600041	**Good identification**							**01600041**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01600050	**Good identification**							**01600050**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01600051	**Good identification**							**01600051**
Clostridium novyi biovar A		100.0	0.45	SAL 8				
01600100	**Acceptable identification**							**01600100**
					COC	GRA	SPO	
Fusobacterium mortiferum		92.63	0.42	FRU 99	1	1	1	
Peptostreptococcus anaerobius		2.56	-.03		99	99	1	
Clostridium novyi biovar A		2.41	0.14		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01610000	**Low discrimination**						**01610000**
					GRA		
Fusobacterium mortiferum		88.15	0.70		1		
Pseudoramibacter alactolyticus		7.20	0.32		99		
Fusobacterium varium		1.01	0.17		1		
01610001	**Low discrimination**						**01610001**
					GRA	SPO	
Fusobacterium mortiferum		57.19	0.36	URE 1	1	1	
Clostridium novyi biovar A		38.14	0.31		99	99	
Pseudoramibacter alactolyticus		4.67	-.03		99	1	
01610002	**Low discrimination**						**01610002**
					GRA		
Fusobacterium mortiferum		85.56	0.54	BGL 10	1		
Pseudoramibacter alactolyticus		6.99	0.15		99		
Atopobium minutum		5.81	0.28		99		
01610100	**Very good identification**						**01610100**
					GRA		
Fusobacterium mortiferum		99.95	0.76		1		
Pseudoramibacter alactolyticus		0.04	-.03		99		
Actinomyces meyeri		0.02	0.04		99		
01610101	**Good identification**						**01610101**
Fusobacterium mortiferum		100.0	0.42	URE 1			
01610102	**Very good identification**						**01610102**
Fusobacterium mortiferum		100.0	0.60	BGL 10			
01610110	**Good identification**						**01610110**
Fusobacterium mortiferum		100.0	0.42	MLZ 1			
01610120	**Good identification**						**01610120**
Fusobacterium mortiferum		100.0	0.42	NAG 1			
01610140	**Good identification**						**01610140**
Fusobacterium mortiferum		100.0	0.42	SOR 1			
01610200	**Acceptable identification**						**01610200**
					GRA		
Fusobacterium mortiferum		92.45	0.36	RHA 1	1		
Pseudoramibacter alactolyticus		7.55	-.03		99		
01610300	**Good identification**						**01610300**
Fusobacterium mortiferum		100.0	0.42	RHA 1			
01610400	**Acceptable identification**						**01610400**
					GRA		
Fusobacterium mortiferum		92.45	0.36	ARA 1	1		
Pseudoramibacter alactolyticus		7.55	-.03		99		
01610500	**Good identification**						**01610500**
Fusobacterium mortiferum		100.0	0.42	ARA 1			
01611000	**Very good identification**						**01611000**
					GRA		
Fusobacterium mortiferum		98.63	0.73		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		0.95	0.40		99		
Atopobium minutum		0.25	0.23		99		
01611001	**Good identification**						**01611001**
					GRA	SPO	
Fusobacterium mortiferum		95.29	0.39	URE 1	1	1	
Clostridium novyi biovar A		4.71	0.15		99	99	
01611002	**Good identification**						**01611002**
					GRA		
Fusobacterium mortiferum		90.74	0.57	BGL 10	1		
Lactobacillus fermentum		7.89	0.40		99		
Atopobium minutum		1.37	0.20		99		
01611010	**Good identification**						**01611010**
					GRA	SPO	
Fusobacterium mortiferum		95.29	0.39	MLZ 1	1	1	
Clostridium novyi biovar A		4.71	0.15		99	99	
01611020	**Low discrimination**						**01611020**
					GRA		
Fusobacterium mortiferum		71.34	0.39	NAG 1	1		
Lactobacillus fermentum		22.75	0.32		99		
Atopobium minutum		5.92	0.15		99		
01611040	**Good identification**						**01611040**
					GRA	SPO	
Fusobacterium mortiferum		96.81	0.39	SOR 1	1	1	
Clostridium novyi biovar A		3.19	0.12		99	99	
01611100	**Very good identification**						**01611100**
					GRA		
Fusobacterium mortiferum		99.29	0.79		1		
Lactobacillus fermentum		0.67	0.44		99		
Actinomyces meyeri		0.02	0.07		99		
01611101	**Good identification**						**01611101**
Fusobacterium mortiferum		100.0	0.45	URE 1			
01611102	**Good identification**						**01611102**
					GRA		
Fusobacterium mortiferum		94.27	0.63	BGL 10	1		
Lactobacillus fermentum		5.73	0.44		99		
01611110	**Good identification**						**01611110**
Fusobacterium mortiferum		100.0	0.45	MLZ 1			
01611120	**Low discrimination**						**01611120**
					GRA		
Fusobacterium mortiferum		81.77	0.45	NAG 1	1		
Lactobacillus fermentum		18.23	0.35		99		
01611122	**Low discrimination**						**01611122**
					GRA		
Lactobacillus fermentum		66.74	0.35	SUC 87	99		
Fusobacterium mortiferum		33.26	0.29	SAL 1	1		
01611140	**Good identification**						**01611140**
Fusobacterium mortiferum		100.0	0.45	SOR 1			
01611200	**Good identification**						**01611200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.39	RHA 1			
01611300	**Good identification**						**01611300**
Fusobacterium mortiferum		100.0	0.45	RHA 1			
01611400	**Low discrimination**						**01611400**
					GRA		
Fusobacterium mortiferum		51.11	0.39	ARA 1	1		
Lactobacillus fermentum		48.89	0.40		99		
01611402	**Low discrimination**						**01611402**
					GRA		
Lactobacillus fermentum		89.59	0.40	SUC 87	99		
Fusobacterium mortiferum		10.41	0.22	SAL 1	1		
01611500	**Low discrimination**						**01611500**
					GRA		
Fusobacterium mortiferum		59.92	0.45	ARA 1	1		
Lactobacillus fermentum		40.08	0.44		99		
01611502	**Low discrimination**						**01611502**
					GRA		
Lactobacillus fermentum		85.76	0.44	SUC 87	99		
Fusobacterium mortiferum		14.25	0.29	SAL 1	1		
01611520	**Good identification**						**01611520**
					GRA		
Lactobacillus fermentum		95.67	0.35	SUC 87	99		
Fusobacterium mortiferum		4.34	0.11	SAL 1	1		
01611522	**Good identification**						**01611522**
Lactobacillus fermentum		100.0	0.35	SUC 87			
01612010	**Good identification**						**01612010**
					SPO		
Clostridium difficile		99.81	0.43	RAF 1	99		
Pseudoramibacter alactolyticus		0.19	-.07		1		
01612100	**Good identification**						**01612100**
					GRA	SPO	
Fusobacterium mortiferum		96.18	0.42	MAN 1	1	1	
Clostridium difficile		2.01	-.03		99	99	
Pseudoramibacter alactolyticus		1.81	-.07		99	1	
01613000	**Acceptable identification**						**01613000**
					GRA	SPO	
Fusobacterium mortiferum		94.18	0.39	MAN 1	1	1	
Clostridium difficile		3.06	-.03		99	99	
Pseudoramibacter alactolyticus		2.76	-.07		99	1	
01613100	**Good identification**						**01613100**
Fusobacterium mortiferum		100.0	0.45	MAN 1			
01614000	**Good identification**						**01614000**
					GRA	SPO	
Fusobacterium mortiferum		95.88	0.51	XYL 7	1	1	
Clostridium glycolicum		2.46	0.09		99	99	
Pseudoramibacter alactolyticus		1.05	-.03		99	1	
01614100	**Very good identification**						**01614100**
					GRA		
Fusobacterium mortiferum		99.48	0.57	XYL 7	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		0.52	0.09		99		
01614102	**Good identification**						**01614102**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
01615000	**Acceptable identification**						**01615000**
					GRA		
Fusobacterium mortiferum		92.99	0.54	XYL 7	1		
Lactobacillus fermentum		7.01	0.36		99		
01615002	**Low discrimination**						**01615002**
					GRA		
Fusobacterium mortiferum		59.58	0.37	XYL 7	1		
Lactobacillus fermentum		40.42	0.36	BGL 10	99		
01615100	**Good identification**						**01615100**
					GRA		
Fusobacterium mortiferum		94.53	0.60	XYL 7	1		
Lactobacillus fermentum		4.98	0.40		99		
Actinomyces meyeri		0.49	0.12		99		
01615102	**Low discrimination**						**01615102**
					GRA		
Fusobacterium mortiferum		67.82	0.44	XYL 7	1		
Lactobacillus fermentum		32.18	0.40	BGL 10	99		
01615400	**Low discrimination**						**01615400**
					GRA		
Lactobacillus fermentum		88.19	0.36	SUC 87	99		
Fusobacterium mortiferum		11.82	0.20	SAL 1	1		
01615402	**Good identification**						**01615402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01615500	**Low discrimination**						**01615500**
					GRA		
Lactobacillus fermentum		83.92	0.40	SUC 87	99		
Fusobacterium mortiferum		16.08	0.26	SAL 1	1		
01615502	**Good identification**						**01615502**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01621100	**Acceptable identification**						**01621100**
					COC	GRA	
Fusobacterium mortiferum		94.52	0.36	FRU 99	1	1	
Blautia hansenii		5.49	-.03		99	99	
01630000	**Very good identification**						**01630000**
					GRA	CAT	
Fusobacterium mortiferum		99.36	0.61		1	5	
Pseudoramibacter alactolyticus		0.27	-.03		99	1	
Propionibacterium granulosum		0.23	0.13		99	90	
01630002	**Good identification**						**01630002**
Fusobacterium mortiferum		100.0	0.45	BGL 10			
01630100	**Very good identification**						**01630100**
Fusobacterium mortiferum		100.0	0.67				
01630102	**Good identification**						**01630102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.51	BGL 10			
01631000	**Very good identification**						**01631000**
					GRA	CAT	
Fusobacterium mortiferum		99.85	0.64		1	5	
Propionibacterium granulosum		0.15	0.13		99	90	
01631002	**Good identification**						**01631002**
Fusobacterium mortiferum		100.0	0.48	BGL 10			
01631100	**Very good identification**						**01631100**
Fusobacterium mortiferum		100.0	0.70				
01631101	**Good identification**						**01631101**
					GRA		
Fusobacterium mortiferum		96.22	0.36	URE 1	1		
Actinomyces naeslundii		3.78	0.24		99		
01631102	**Good identification**						**01631102**
Fusobacterium mortiferum		100.0	0.54	BGL 10			
01631110	**Good identification**						**01631110**
Fusobacterium mortiferum		100.0	0.36	MLZ 1			
01631120	**Good identification**						**01631120**
					GRA	SPO	
Fusobacterium mortiferum		95.20	0.36	NAG 1	1	1	
Clostridium septicum		4.80	-.04		99	99	
01631140	**Good identification**						**01631140**
Fusobacterium mortiferum		100.0	0.36	SOR 1			
01631300	**Good identification**						**01631300**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
01631500	**Good identification**						**01631500**
Fusobacterium mortiferum		100.0	0.36	ARA 1			
01633100	**Good identification**						**01633100**
					GRA		
Fusobacterium mortiferum		95.16	0.36	MAN 1	1		
Actinomyces naeslundii		4.84	0.26		99		
01634000	**Good identification**						**01634000**
Fusobacterium mortiferum		100.0	0.42	XYL 7			
01634100	**Good identification**						**01634100**
Fusobacterium mortiferum		100.0	0.48	XYL 7			
01635000	**Good identification**						**01635000**
Fusobacterium mortiferum		100.0	0.45	XYL 7			
01635100	**Good identification**						**01635100**
Fusobacterium mortiferum		100.0	0.51	XYL 7			
01635122	**Good identification**						**01635122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.37	MNS 98			
01635322	**Good identification**						**01635322**
Parabacteroides distasonis		100.0	0.45	MNS 98			
01635332	**Good identification**						**01635332**
Parabacteroides distasonis		100.0	0.41	MNS 98			
01635722	**Good identification**						**01635722**
Parabacteroides distasonis		100.0	0.41	MNS 98			
01635732	**Good identification**						**01635732**
Parabacteroides distasonis		100.0	0.37	MNS 98			
01641100	**Good identification**						**01641100**
					COC	GRA	
Fusobacterium mortiferum		95.06	0.37	FRU 99	1	1	
Blautia hansenii		4.94	-.03		99	99	
01650000	**Very good identification**						**01650000**
					GRA		
Fusobacterium mortiferum		99.75	0.62		1		
Pseudoramibacter alactolyticus		0.25	-.03		99		
01650002	**Good identification**						**01650002**
Fusobacterium mortiferum		100.0	0.45	BGL 10			
01650100	**Very good identification**						**01650100**
Fusobacterium mortiferum		100.0	0.68				
01650102	**Good identification**						**01650102**
Fusobacterium mortiferum		100.0	0.52	BGL 10			
01651000	**Very good identification**						**01651000**
Fusobacterium mortiferum		100.0	0.65				
01651002	**Good identification**						**01651002**
Fusobacterium mortiferum		100.0	0.48	BGL 10			
01651100	**Very good identification**						**01651100**
Fusobacterium mortiferum		100.0	0.71				
01651101	**Good identification**						**01651101**
Fusobacterium mortiferum		100.0	0.37	URE 1			
01651102	**Good identification**						**01651102**
Fusobacterium mortiferum		100.0	0.55	BGL 10			
01651110	**Good identification**						**01651110**
Fusobacterium mortiferum		100.0	0.37	MLZ 1			
01651120	**Good identification**						**01651120**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		95.45	0.37	NAG 1	1	1	
Clostridium septicum		4.55	-.03		99	99	
01651140	**Good identification**						**01651140**
Fusobacterium mortiferum		100.0	0.37	SOR 1			
01651300	**Good identification**						**01651300**
Fusobacterium mortiferum		100.0	0.37	RHA 1			
01651500	**Good identification**						**01651500**
Fusobacterium mortiferum		100.0	0.37	ARA 1			
01653100	**Good identification**						**01653100**
					GRA		
Fusobacterium mortiferum		96.98	0.37	MAN 1	1		
Actinomyces naeslundii		3.02	0.23		99		
01654000	**Good identification**						**01654000**
Fusobacterium mortiferum		100.0	0.43	XYL 7			
01654100	**Good identification**						**01654100**
Fusobacterium mortiferum		100.0	0.49	XYL 7			
01655000	**Good identification**						**01655000**
Fusobacterium mortiferum		100.0	0.46	XYL 7			
01655100	**Good identification**						**01655100**
Fusobacterium mortiferum		100.0	0.52	XYL 7			
01655102	**Good identification**						**01655102**
Fusobacterium mortiferum		100.0	0.35	XYL 7			
01670000	**Good identification**						**01670000**
Fusobacterium mortiferum		100.0	0.53				
01670002	**Good identification**						**01670002**
Fusobacterium mortiferum		100.0	0.36	BGL 10			
01670100	**Very good identification**						**01670100**
Fusobacterium mortiferum		100.0	0.59				
01670102	**Good identification**						**01670102**
Fusobacterium mortiferum		100.0	0.43	BGL 10			
01671000	**Good identification**						**01671000**
Fusobacterium mortiferum		100.0	0.56				
01671002	**Good identification**						**01671002**
Fusobacterium mortiferum		100.0	0.39	BGL 10			
01671100	**Very good identification**						**01671100**
					GRA		
Fusobacterium mortiferum		99.90	0.62		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		0.11	0.23		99			
01671102	**Good identification**							**01671102**
					GRA			
Fusobacterium mortiferum		98.05	0.46	BGL 10	1			
Lactobacillus acidophilus		1.02	-.01		99			
Actinomyces naeslundii		0.93	0.23		99			
01674100	**Good identification**							**01674100**
Fusobacterium mortiferum		100.0	0.40	XYL 7				
01675000	**Good identification**							**01675000**
Fusobacterium mortiferum		100.0	0.37	XYL 7				
01675100	**Good identification**							**01675100**
Fusobacterium mortiferum		100.0	0.43	XYL 7				
01675322	**Good identification**							**01675322**
Parabacteroides distasonis		100.0	0.41	MNS 98				
01675332	**Good identification**							**01675332**
Parabacteroides distasonis		100.0	0.36	MNS 98				
01675722	**Good identification**							**01675722**
Parabacteroides distasonis		100.0	0.37	MNS 98				
01700000	**Low discrimination**							**01700000**
					COC	GRA	SPO	
Clostridium novyi biovar A		35.63	0.48	SAL 8	1	99	99	
Peptostreptococcus anaerobius		25.22	0.29	RAF 1	99	99	1	
Prevotella disiens		13.90	0.23		1	1	1	
01700001	**Good identification**							**01700001**
					COC	SPO		
Clostridium novyi biovar A		98.20	0.48	SAL 8	1	99		
Clostridium sordellii		1.10	-.03	RAF 1	1	99		
Peptostreptococcus anaerobius		0.70	-.06		99	1		
01700010	**Good identification**							**01700010**
					COC	SPO		
Clostridium novyi biovar A		99.29	0.48	SAL 8	1	99		
Peptostreptococcus anaerobius		0.71	-.06	RAF 1	99	1		
01700011	**Good identification**							**01700011**
Clostridium novyi biovar A		100.0	0.48	SAL 8				
01700020	**Acceptable identification**							**01700020**
					COC	SPO		
Clostridium novyi biovar A		91.27	0.37	SAL 8	1	99		
Clostridium bifermentans		6.12	0.05	RAF 1	1	99		
Peptostreptococcus anaerobius		2.61	-.06		99	1		
01700021	**Good identification**							**01700021**
Clostridium novyi biovar A		100.0	0.37	SAL 8				
01700030	**Good identification**							**01700030**
Clostridium novyi biovar A		100.0	0.37	SAL 8				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
01700031	**Good identification**						**01700031**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
01700040	**Low discrimination**						**01700040**
Clostridium novyi biovar A		86.92	0.45	SAL 8			
Clostridium botulinum biovar A		9.74	0.17	RAF 1			
Clostridium bifermentans		1.43	0.02				
01700041	**Good identification**						**01700041**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01700050	**Good identification**						**01700050**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01700051	**Good identification**						**01700051**
Clostridium novyi biovar A		100.0	0.45	SAL 8			
01701100	**Very good identification**						**01701100**
					COC	GRA	
Blautia hansenii		99.92	0.66	SAL 1	99	99	
Fusobacterium mortiferum		0.07	0.30		1	1	
Lactobacillus fermentum		0.01	0.18		1	99	
01701102	**Good identification**						**01701102**
					COC	GRA	
Blautia hansenii		99.82	0.50	SAL 1	99	99	
Lactobacillus fermentum		0.11	0.18	BGL 10	1	99	
Fusobacterium mortiferum		0.07	0.13		1	1	
01701120	**Good identification**						**01701120**
					COC	GRA	
Blautia hansenii		99.88	0.50	SAL 1	99	99	
Prevotella melaninogenica		0.12	-.01	NAG 10	1	1	
01701500	**Good identification**						**01701500**
					COC		
Blautia hansenii		99.93	0.53	SAL 1	99		
Lactobacillus fermentum		0.07	0.18		1		
01701502	**Good identification**						**01701502**
					COC		
Blautia hansenii		99.40	0.37	SAL 1	99		
Lactobacillus fermentum		0.61	0.18	BGL 10	1		
01701520	**Good identification**						**01701520**
Blautia hansenii		100.0	0.37	SAL 1			
01705100	**Good identification**						**01705100**
					COC		
Blautia hansenii		99.96	0.53	SAL 1	99		
Lactobacillus fermentum		0.04	0.14		1		
01705102	**Good identification**						**01705102**
					COC		
Blautia hansenii		99.64	0.37	SAL 1	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
01705120	**Good identification**						**01705120**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		100.0	0.37	SAL 1				
01705500	**Good identification**							**01705500**
					COC			
Blautia hansenii		99.78	0.40	SAL 1	99			
Lactobacillus fermentum		0.23	0.14		1			
01710000	**Low discrimination**							**01710000**
Fusobacterium mortiferum		45.26	0.54	MLT 11				
Prevotella intermedia		37.08	0.42					
Prevotella disiens		8.66	0.23					
01710002	**Low discrimination**							**01710002**
					COC	GRA	SPO	
Fusobacterium mortiferum		75.03	0.38	MLT 11	1	1	1	
Megasphaera elsdenii		7.83	0.12	BGL 10	99	1	1	
Clostridium novyi biovar A		6.49	0.19		1	99	99	
01710100	**Very good identification**							**01710100**
					GRA			
Fusobacterium mortiferum		99.01	0.61	MLT 11	1			
Prevotella intermedia		0.35	0.08		1			
Lactobacillus fermentum		0.25	0.18		99			
01710102	**Good identification**							**01710102**
					GRA			
Fusobacterium mortiferum		96.77	0.44	MLT 11	1			
Lactobacillus fermentum		2.16	0.18	BGL 10	99			
Actinomyces viscosus		1.07	-.02		99			
01711000	**Low discrimination**							**01711000**
					GRA			
Fusobacterium mortiferum		55.80	0.57	MLT 11	1			
Prevotella intermedia		30.47	0.42		1			
Lactobacillus fermentum		13.09	0.48		99			
01711002	**Low discrimination**							**01711002**
					GRA			
Lactobacillus fermentum		65.99	0.48	SUC 87	99			
Fusobacterium mortiferum		31.27	0.41	SAL 1	1			
Prevotella intermedia		1.55	0.08		1			
01711020	**Low discrimination**							**01711020**
					GRA			
Lactobacillus fermentum		76.17	0.40	SUC 87	99			
Fusobacterium mortiferum		9.84	0.23	SAL 1	1			
Prevotella intermedia		5.38	0.08		1			
01711022	**Low discrimination**							**01711022**
					GRA			
Lactobacillus fermentum		86.51	0.40	SUC 87	99			
Prevotella oralis		6.41	0.05	SAL 1	1			
Actinomyces naeslundii		4.69	0.28		99			
01711100	**Low discrimination**							**01711100**
					COC	GRA		
Fusobacterium mortiferum		76.13	0.64	MLT 11	1	1		
Lactobacillus fermentum		12.48	0.52		1	99		
Blautia hansenii		10.68	0.32		99	99		
01711102	**Low discrimination**							**01711102**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		54.90	0.52	SUC 87	1	99	
Fusobacterium mortiferum		37.20	0.47	SAL 1	1	1	
Blautia hansenii		5.22	0.15		99	99	
01711103	**Low discrimination**						**01711103**
					GRA		
Actinomyces naeslundii		73.60	0.39	SUC 99	99		
Lactobacillus fermentum		15.74	0.18	MNS 88	99		
Fusobacterium mortiferum		10.66	0.13		1		
01711120	**Low discrimination**						**01711120**
					COC	GRA	
Lactobacillus fermentum		43.87	0.44	SUC 87	1	99	
Prevotella melaninogenica		12.98	0.15	SAL 1	1	1	
Blautia hansenii		12.50	0.15		99	99	
01711121	**Good identification**						**01711121**
Actinomyces naeslundii		100.0	0.39	SUC 99			
01711122	**Low discrimination**						**01711122**
					GRA		
Lactobacillus fermentum		47.21	0.44	SUC 87	99		
Prevotella oralis		33.61	0.25	SAL 1	1		
Clostridium perfringens		5.88	0.16		99		
01711123	**Good identification**						**01711123**
Actinomyces naeslundii		100.0	0.39	SUC 99			
01711400	**Acceptable identification**						**01711400**
					GRA		
Lactobacillus fermentum		93.76	0.48	SUC 87	99		
Fusobacterium mortiferum		4.04	0.23	SAL 1	1		
Prevotella intermedia		2.21	0.08		1		
01711402	**Good identification**						**01711402**
Lactobacillus fermentum		100.0	0.48	SUC 87			
01711420	**Good identification**						**01711420**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01711422	**Good identification**						**01711422**
Lactobacillus fermentum		100.0	0.40	SUC 87			
01711500	**Low discrimination**						**01711500**
					COC	GRA	
Lactobacillus fermentum		82.48	0.52	SUC 87	1	99	
Blautia hansenii		12.45	0.19	SAL 1	99	99	
Fusobacterium mortiferum		5.08	0.30		1	1	
01711502	**Acceptable identification**						**01711502**
					COC		
Lactobacillus fermentum		94.92	0.52	SUC 87	1		
Bifidobacterium adolescentis		2.84	0.20	SAL 1	1		
Blautia hansenii		1.59	0.03		99		
01711520	**Acceptable identification**						**01711520**
					COC		
Lactobacillus fermentum		93.84	0.44	SUC 87	1		
Blautia hansenii		4.72	0.03	SAL 1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		1.44	0.05		1		
01711522	**Good identification**						**01711522**
					GRA		
Lactobacillus fermentum		95.47	0.44	SUC 87	99		
Parabacteroides distasonis		3.06	0.15	SAL 1	1		
Bifidobacterium longum subsp. longum		1.46	0.05		99		
01713001	**Good identification**						**01713001**
Actinomyces naeslundii		100.0	0.36	SUC 99			
01713003	**Good identification**						**01713003**
Actinomyces naeslundii		100.0	0.36	SUC 99			
01713021	**Good identification**						**01713021**
Actinomyces naeslundii		100.0	0.36	SUC 99			
01713023	**Good identification**						**01713023**
Actinomyces naeslundii		100.0	0.36	SUC 99			
01713101	**Good identification**						**01713101**
Actinomyces naeslundii		100.0	0.44	SUC 99			
01713102	**Low discrimination**						**01713102**
Actinomyces naeslundii		71.42	0.41	SUC 99			
Lactobacillus fermentum		11.80	0.18	MNS 88			
Propionibacterium propionicum		8.78	0.15				
01713103	**Good identification**						**01713103**
Actinomyces naeslundii		100.0	0.44	SUC 99			
01713120	**Good identification**						**01713120**
Actinomyces naeslundii		100.0	0.41	SUC 99			
01713121	**Good identification**						**01713121**
Actinomyces naeslundii		100.0	0.44	SUC 99			
01713122	**Good identification**						**01713122**
Actinomyces naeslundii		100.0	0.41	SUC 99			
01713123	**Good identification**						**01713123**
Actinomyces naeslundii		100.0	0.44	SUC 99			
01714000	**Low discrimination**						**01714000**
					GRA	SPO	
Fusobacterium mortiferum		74.90	0.35	MLT 11	1	1	
Clostridium glycolicum		16.87	0.10	XYL 7	99	99	
Prevotella intermedia		8.24	0.08		1	1	
01714100	**Acceptable identification**						**01714100**
					GRA		
Fusobacterium mortiferum		93.73	0.41	MLT 11	1		
Actinomyces meyeri		4.45	0.09	XYL 7	99		
Lactobacillus fermentum		1.82	0.14		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
01715000	**Low discrimination**						**01715000**
					GRA		
Lactobacillus fermentum		62.44	0.44	SUC 87	99		
Fusobacterium mortiferum		34.12	0.38	SAL 1	1		
Prevotella intermedia		2.50	0.08		1		
01715002	**Acceptable identification**						**01715002**
					GRA		
Lactobacillus fermentum		94.28	0.44	SUC 87	99		
Fusobacterium mortiferum		5.72	0.22	SAL 1	1		
01715020	**Good identification**						**01715020**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01715022	**Good identification**						**01715022**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01715100	**Low discrimination**						**01715100**
					COC	GRA	
Lactobacillus fermentum		47.94	0.48	SUC 87	1	99	
Fusobacterium mortiferum		37.46	0.44	SAL 1	1	1	
Blautia hansenii		12.32	0.19		99	99	
01715102	**Low discrimination**						**01715102**
					GRA		
Lactobacillus fermentum		84.59	0.48	SUC 87	99		
Fusobacterium mortiferum		7.35	0.28	SAL 1	1		
Bifidobacterium adolescentis		4.30	0.20		99		
01715120	**Acceptable identification**						**01715120**
					COC		
Lactobacillus fermentum		92.11	0.40	SUC 87	1		
Blautia hansenii		7.89	0.03	SAL 1	99		
01715122	**Low discrimination**						**01715122**
					GRA		
Lactobacillus fermentum		50.27	0.40	SUC 87	99		
Parabacteroides distasonis		40.33	0.35	SAL 1	1		
Prevotella oralis		7.53	0.10		1		
01715322	**Good identification**						**01715322**
					GRA		
Parabacteroides distasonis		95.90	0.43	MNS 98	1		
Prevotella oralis		2.56	0.04		1		
Eubacterium contortum		1.55	0.01		99		
01715332	**Good identification**						**01715332**
Parabacteroides distasonis		100.0	0.39	MNS 98			
01715400	**Good identification**						**01715400**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01715402	**Good identification**						**01715402**
Lactobacillus fermentum		100.0	0.44	SUC 87			
01715420	**Good identification**						**01715420**
Lactobacillus fermentum		100.0	0.36	SUC 87			
01715422	**Good identification**						**01715422**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		100.0	0.36	SUC 87			
01715500	**Acceptable identification**						**01715500**
					COC		
Lactobacillus fermentum		94.05	0.48	SUC 87	1		
Blautia hansenii		4.26	0.06	SAL 1	99		
Actinomyces meyeri		0.87	0.02		1		
01715502	**Low discrimination**						**01715502**
Lactobacillus fermentum		73.88	0.48	SUC 87			
Bifidobacterium adolescentis		25.16	0.34	SAL 1			
Actinomyces israelii		0.96	0.16				
01715512	**Good identification**						**01715512**
Bifidobacterium adolescentis		95.15	0.36	SUC 99			
Bifidobacterium longum subsp. longum		2.54	0.07	MNS 87			
Lactobacillus fermentum		2.31	0.14				
01715520	**Low discrimination**						**01715520**
					GRA		
Lactobacillus fermentum		72.68	0.40	SUC 87	99		
Bacteroides vulgatus		21.63	0.09	SAL 1	1		
Bifidobacterium longum subsp. longum		5.69	0.14		99		
01715522	**Low discrimination**						**01715522**
					GRA		
Lactobacillus fermentum		63.69	0.40	SUC 87	99		
Parabacteroides distasonis		31.32	0.31	SAL 1	1		
Bifidobacterium longum subsp. longum		4.99	0.14		99		
01715722	**Good identification**						**01715722**
					GRA		
Parabacteroides distasonis		96.17	0.40	MNS 98	1		
Eubacterium contortum		2.07	0.00		99		
Bacteroides vulgatus		1.76	-.09		1		
01730000	**Good identification**						**01730000**
					GRA	SPO	
Fusobacterium mortiferum		96.07	0.45	MLT 11	1	1	
Prevotella intermedia		2.66	0.08		1	1	
Clostridium sporogenes		1.27	-.02		99	99	
01730100	**Good identification**						**01730100**
Fusobacterium mortiferum		100.0	0.52	MLT 11			
01730102	**Good identification**						**01730102**
Fusobacterium mortiferum		100.0	0.35	MLT 11			
01731000	**Acceptable identification**						**01731000**
					GRA		
Fusobacterium mortiferum		94.88	0.48	MLT 11	1		
Actinomyces naeslundii		2.61	0.33		99		
Prevotella intermedia		1.75	0.08		1		
01731001	**Low discrimination**						**01731001**
					GRA		
Actinomyces naeslundii		80.34	0.36	SUC 99	99		
Fusobacterium mortiferum		19.66	0.14	MNS 88	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01731003	**Good identification**					**01731003**
Actinomyces naeslundii		100.0	0.36	SUC 99		
01731021	**Good identification**					**01731021**
Actinomyces naeslundii		100.0	0.36	SUC 99		
01731023	**Good identification**					**01731023**
Actinomyces naeslundii		100.0	0.36	SUC 99		
01731100	**Good identification**					**01731100**
					GRA	
Fusobacterium mortiferum		95.55	0.55	MLT 11	1	
Actinomyces naeslundii		3.21	0.41		99	
Lactobacillus fermentum		0.53	0.18		99	
01731101	**Low discrimination**					**01731101**
					GRA	
Actinomyces naeslundii		83.29	0.44	SUC 99	99	
Fusobacterium mortiferum		16.71	0.21	MNS 88	1	
01731102	**Low discrimination**					**01731102**
					GRA	
Fusobacterium mortiferum		73.97	0.38	MLT 11	1	
Actinomyces naeslundii		22.34	0.41	BGL 10	99	
Lactobacillus fermentum		3.69	0.18		99	
01731103	**Good identification**					**01731103**
Actinomyces naeslundii		100.0	0.44	SUC 99		
01731121	**Good identification**					**01731121**
Actinomyces naeslundii		100.0	0.44	SUC 99		
01731123	**Good identification**					**01731123**
Actinomyces naeslundii		100.0	0.44	SUC 99		
01731322	**Good identification**					**01731322**
Parabacteroides distasonis		97.35	0.38	MNS 98		
Prevotella oralis		2.65	-.01	XYL 90		
01733000	**Low discrimination**					**01733000**
					GRA	
Actinomyces naeslundii		84.10	0.38	SUC 99	99	
Fusobacterium mortiferum		15.90	0.14	MNS 88	1	
01733001	**Good identification**					**01733001**
Actinomyces naeslundii		100.0	0.41	SUC 99		
01733002	**Good identification**					**01733002**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01733003	**Good identification**					**01733003**
Actinomyces naeslundii		100.0	0.41	SUC 99		
01733020	**Good identification**					**01733020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	SUC 99		
01733021	**Good identification**					**01733021**
Actinomyces naeslundii		100.0	0.41	SUC 99		
01733022	**Good identification**					**01733022**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01733023	**Good identification**					**01733023**
Actinomyces naeslundii		100.0	0.41	SUC 99		
01733100	**Low discrimination**					**01733100**
					GRA	
Actinomyces naeslundii		79.83	0.46	SUC 99	99	
Fusobacterium mortiferum		12.38	0.21	MNS 88	1	
Actinomyces israelii		7.79	0.19		99	
01733101	**Good identification**					**01733101**
Actinomyces naeslundii		100.0	0.49	SUC 99		
01733102	**Low discrimination**					**01733102**
Actinomyces naeslundii		83.46	0.46	SUC 99		
Actinomyces israelii		12.21	0.22	MNS 88		
Propionibacterium propionicum		4.32	0.14			
01733103	**Good identification**					**01733103**
Actinomyces naeslundii		100.0	0.49	SUC 99		
01733120	**Good identification**					**01733120**
Actinomyces naeslundii		100.0	0.46	SUC 99		
01733121	**Good identification**					**01733121**
Actinomyces naeslundii		100.0	0.49	SUC 99		
01733122	**Good identification**					**01733122**
Actinomyces naeslundii		100.0	0.46	SUC 99		
01733123	**Good identification**					**01733123**
Actinomyces naeslundii		100.0	0.49	SUC 99		
01733141	**Good identification**					**01733141**
Actinomyces naeslundii		100.0	0.36	SUC 99		
01733143	**Good identification**					**01733143**
Actinomyces naeslundii		100.0	0.36	SUC 99		
01733161	**Good identification**					**01733161**
Actinomyces naeslundii		100.0	0.36	SUC 99		
01733163	**Good identification**					**01733163**
Actinomyces naeslundii		100.0	0.36	SUC 99		
01735100	**Low discrimination**					**01735100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		82.33	0.35	MLT 11	1	
Actinomyces israelii		14.11	0.25	XYL 7	99	
Lactobacillus fermentum		3.56	0.14		99	
01735122	**Good identification**					**01735122**
Parabacteroides distasonis		100.0	0.46	MNS 98		
01735132	**Good identification**					**01735132**
Parabacteroides distasonis		100.0	0.42	MNS 98		
01735322	**Good identification**					**01735322**
Parabacteroides distasonis		100.0	0.55	MNS 98		
01735332	**Good identification**					**01735332**
Parabacteroides distasonis		100.0	0.50	MNS 98		
01735522	**Good identification**					**01735522**
Parabacteroides distasonis		100.0	0.43	MNS 98		
01735532	**Good identification**					**01735532**
					GRA	
Parabacteroides distasonis		97.31	0.38	MNS 98	1	
Bifidobacterium longum subsp. longum		2.69	0.07		99	
01735722	**Good identification**					**01735722**
Parabacteroides distasonis		100.0	0.51	MNS 98		
01735732	**Good identification**					**01735732**
Parabacteroides distasonis		97.37	0.46	MNS 98		
Clostridium clostridioforme		2.63	0.14			
01737100	**Good identification**					**01737100**
Actinomyces israelii		100.0	0.38	SUC 98		
01737102	**Acceptable identification**					**01737102**
Actinomyces israelii		92.45	0.41	SUC 98		
Bifidobacterium adolescentis		7.55	0.16	CEL 93		
01737500	**Good identification**					**01737500**
Actinomyces israelii		100.0	0.36	SUC 98		
01737502	**Low discrimination**					**01737502**
Actinomyces israelii		59.96	0.39	SUC 98		
Bifidobacterium adolescentis		40.04	0.30	CEL 93		
01750000	**Good identification**					**01750000**
Fusobacterium mortiferum		97.58	0.46	MLT 11		
Prevotella intermedia		2.42	0.08			
01750100	**Good identification**					**01750100**
Fusobacterium mortiferum		100.0	0.52	MLT 11		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01750102	**Good identification**					**01750102**
Fusobacterium mortiferum		100.0	0.36	MLT 11		
01751000	**Acceptable identification**					**01751000**
					GRA	
Fusobacterium mortiferum		94.88	0.49	MLT 11	1	
Actinomyces naeslundii		1.60	0.30		99	
Prevotella intermedia		1.57	0.08		1	
01751100	**Good identification**					**01751100**
					GRA	
Fusobacterium mortiferum		96.78	0.56	MLT 11	1	
Actinomyces naeslundii		1.99	0.38		99	
Lactobacillus fermentum		0.48	0.18		99	
01751101	**Low discrimination**					**01751101**
					GRA	
Actinomyces naeslundii		75.31	0.41	SUC 99	99	
Fusobacterium mortiferum		24.69	0.21	MNS 88	1	
01751102	**Low discrimination**					**01751102**
					GRA	
Fusobacterium mortiferum		61.22	0.39	MLT 11	1	
Prevotella oralis		12.00	0.13	BGL 10	1	
Actinomyces naeslundii		11.31	0.38		99	
01751103	**Good identification**					**01751103**
Actinomyces naeslundii		100.0	0.41	SUC 99		
01751121	**Good identification**					**01751121**
Actinomyces naeslundii		100.0	0.41	SUC 99		
01751123	**Low discrimination**					**01751123**
					GRA	
Actinomyces naeslundii		88.05	0.41	SUC 99	99	
Prevotella oralis		11.95	0.01	MNS 88	1	
01753000	**Low discrimination**					**01753000**
					GRA	
Actinomyces naeslundii		59.95	0.35	SUC 99	99	
Actinomyces israelii		21.52	0.19	MNS 88	99	
Fusobacterium mortiferum		18.53	0.15		1	
01753001	**Good identification**					**01753001**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01753002	**Low discrimination**					**01753002**
Actinomyces naeslundii		65.00	0.35	SUC 99		
Actinomyces israelii		35.00	0.22	MNS 88		
01753003	**Good identification**					**01753003**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01753020	**Good identification**					**01753020**
Actinomyces naeslundii		100.0	0.35	SUC 99		
01753021	**Good identification**					**01753021**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	SUC 99			
01753022	**Good identification**						**01753022**
Actinomyces naeslundii		100.0	0.35	SUC 99			
01753023	**Good identification**						**01753023**
Actinomyces naeslundii		100.0	0.38	SUC 99			
01753100	**Low discrimination**						**01753100**
					GRA		
Actinomyces naeslundii		56.89	0.43	SUC 99	99		
Actinomyces israelii		28.70	0.29	MNS 88	99		
Fusobacterium mortiferum		14.41	0.21		1		
01753101	**Good identification**						**01753101**
Actinomyces naeslundii		100.0	0.46	SUC 99			
01753102	**Low discrimination**						**01753102**
Actinomyces naeslundii		53.98	0.43	SUC 99			
Actinomyces israelii		40.85	0.32	MNS 88			
Bifidobacterium adolescentis		5.17	0.10				
01753103	**Good identification**						**01753103**
Actinomyces naeslundii		100.0	0.46	SUC 99			
01753120	**Good identification**						**01753120**
Actinomyces naeslundii		100.0	0.43	SUC 99			
01753121	**Good identification**						**01753121**
Actinomyces naeslundii		100.0	0.46	SUC 99			
01753122	**Low discrimination**						**01753122**
					SPO	GRA	
Actinomyces naeslundii		78.58	0.43	SUC 99	1	99	
Clostridium ramosum		13.18	0.05	MNS 88	92	50	
Prevotella oralis		8.24	0.01		1	1	
01753123	**Good identification**						**01753123**
Actinomyces naeslundii		100.0	0.46	SUC 99			
01755100	**Low discrimination**						**01755100**
					GRA		
Fusobacterium mortiferum		63.25	0.36	MLT 11	1		
Actinomyces israelii		34.30	0.34	XYL 7	99		
Lactobacillus fermentum		2.45	0.14		99		
01755102	**Low discrimination**						**01755102**
					GRA		
Actinomyces israelii		57.52	0.37	SUC 98	99		
Bifidobacterium adolescentis		29.35	0.26		99		
Fusobacterium mortiferum		7.86	0.20		1		
01755322	**Low discrimination**						**01755322**
Parabacteroides distasonis		85.06	0.39	MNS 98			
Prevotella oralis		14.94	0.13				
01755502	**Low discrimination**						**01755502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		79.77	0.40	SUC 99		
Actinomyces israelii		19.11	0.36	MNS 87		
Lactobacillus fermentum		1.11	0.14			
01755512	**Good identification**					**01755512**
Bifidobacterium adolescentis		97.15	0.41	SUC 99		
Actinomyces israelii		2.85	0.21	MNS 87		
01757000	**Good identification**					**01757000**
Actinomyces israelii		100.0	0.37	SUC 98		
01757002	**Good identification**					**01757002**
Actinomyces israelii		100.0	0.40	SUC 98		
01757100	**Good identification**					**01757100**
Actinomyces israelii		100.0	0.47	SUC 98		
01757102	**Acceptable identification**					**01757102**
Actinomyces israelii		93.14	0.50	SUC 98		
Bifidobacterium adolescentis		6.86	0.24			
01757112	**Low discrimination**					**01757112**
Actinomyces israelii		62.40	0.36	SUC 98		
Bifidobacterium adolescentis		37.60	0.26	MLZ 13		
01757140	**Good identification**					**01757140**
Actinomyces israelii		100.0	0.40	SUC 98		
01757142	**Good identification**					**01757142**
Actinomyces israelii		97.71	0.43	SUC 98		
Bifidobacterium adolescentis		2.29	0.09			
01757400	**Good identification**					**01757400**
Actinomyces israelii		100.0	0.35	SUC 98		
01757402	**Good identification**					**01757402**
Actinomyces israelii		100.0	0.38	SUC 98		
01757500	**Good identification**					**01757500**
Actinomyces israelii		100.0	0.46	SUC 98		
01757502	**Low discrimination**					**01757502**
Actinomyces israelii		62.40	0.49	SUC 98		
Bifidobacterium adolescentis		37.60	0.38			
01757512	**Low discrimination**					**01757512**
Bifidobacterium adolescentis		83.13	0.40	SUC 99		
Actinomyces israelii		16.87	0.34	MNS 87		
01757540	**Good identification**					**01757540**
Actinomyces israelii		100.0	0.38	SUC 98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01757542	**Low discrimination**					**01757542**
Actinomyces israelii		83.93	0.41	SUC 98		
Bifidobacterium adolescentis		16.07	0.23			
01770000	**Good identification**					**01770000**
Fusobacterium mortiferum		100.0	0.37	MLT 11		
01770100	**Good identification**					**01770100**
Fusobacterium mortiferum		100.0	0.44	MLT 11		
01771000	**Low discrimination**					**01771000**
					GRA	
Fusobacterium mortiferum		78.69	0.40	MLT 11	1	
Lactobacillus jensenii		10.65	0.15		99	
Actinomyces naeslundii		8.61	0.35		99	
01771001	**Good identification**					**01771001**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01771003	**Good identification**					**01771003**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01771020	**Low discrimination**					**01771020**
Actinomyces naeslundii		76.38	0.35	SUC 99		
Lactobacillus jensenii		23.62	0.05	MNS 88		
01771021	**Good identification**					**01771021**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01771022	**Low discrimination**					**01771022**
Actinomyces naeslundii		67.63	0.35	SUC 99		
Lactobacillus acidophilus		18.43	0.01	MNS 88		
Lactobacillus jensenii		13.94	0.02			
01771023	**Good identification**					**01771023**
Actinomyces naeslundii		100.0	0.38	SUC 99		
01771100	**Low discrimination**					**01771100**
					GRA	
Fusobacterium mortiferum		84.88	0.47	MLT 11	1	
Actinomyces naeslundii		11.33	0.43		99	
Actinomyces israelii		3.79	0.26		99	
01771101	**Good identification**					**01771101**
					GRA	
Actinomyces naeslundii		95.20	0.46	SUC 99	99	
Fusobacterium mortiferum		4.80	0.12	MNS 88	1	
01771103	**Good identification**					**01771103**
Actinomyces naeslundii		100.0	0.46	SUC 99		
01771121	**Good identification**					**01771121**
Actinomyces naeslundii		100.0	0.46	SUC 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
01771123	**Good identification**					**01771123**
Actinomyces naeslundii		100.0	0.46	SUC 99		
01773000	**Low discrimination**					**01773000**
Actinomyces naeslundii		58.97	0.40	SUC 99		
Actinomyces israelii		41.03	0.28	MNS 88		
01773001	**Good identification**					**01773001**
Actinomyces naeslundii		100.0	0.43	SUC 99		
01773003	**Good identification**					**01773003**
Actinomyces naeslundii		100.0	0.43	SUC 99		
01773020	**Good identification**					**01773020**
Actinomyces naeslundii		100.0	0.40	SUC 99		
01773021	**Good identification**					**01773021**
Actinomyces naeslundii		100.0	0.43	SUC 99		
01773022	**Good identification**					**01773022**
Actinomyces naeslundii		100.0	0.40	SUC 99		
01773023	**Good identification**					**01773023**
Actinomyces naeslundii		100.0	0.43	SUC 99		
01773100	**Low discrimination**					**01773100**
					GRA	
Actinomyces naeslundii		49.58	0.48	SUC 99	99	
Actinomyces israelii		48.48	0.39	MNS 88	99	
Fusobacterium mortiferum		1.93	0.12		1	
01773101	**Good identification**					**01773101**
Actinomyces naeslundii		100.0	0.51	SUC 99		
01773102	**Low discrimination**					**01773102**
Actinomyces israelii		58.68	0.42	SUC 98		
Actinomyces naeslundii		40.01	0.48	XYL 92		
Bifidobacterium adolescentis		1.32	0.07			
01773103	**Good identification**					**01773103**
Actinomyces naeslundii		100.0	0.51	SUC 99		
01773120	**Low discrimination**					**01773120**
					SPO	
Actinomyces naeslundii		71.38	0.48	SUC 99	1	
Clostridium ramosum		24.16	0.15	MNS 88	92	
Actinomyces israelii		4.46	0.18		1	
01773121	**Good identification**					**01773121**
Actinomyces naeslundii		100.0	0.51	SUC 99		
01773123	**Good identification**					**01773123**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.51	SUC 99			
01773141	**Good identification**						**01773141**
Actinomyces naeslundii		100.0	0.38	SUC 99			
01773143	**Good identification**						**01773143**
Actinomyces naeslundii		100.0	0.38	SUC 99			
01773160	**Good identification**						**01773160**
Actinomyces naeslundii		100.0	0.35	SUC 99			
01773161	**Good identification**						**01773161**
Actinomyces naeslundii		100.0	0.38	SUC 99			
01773162	**Low discrimination**						**01773162**
Actinomyces naeslundii		82.89	0.35	SUC 99			
Actinomyces israelii		17.11	0.14	MNS 88			
01773163	**Good identification**						**01773163**
Actinomyces naeslundii		100.0	0.38	SUC 99			
01773500	**Good identification**						**01773500**
Actinomyces israelii		100.0	0.37	SUC 98			
01773502	**Low discrimination**						**01773502**
Actinomyces israelii		84.49	0.40	SUC 98			
Bifidobacterium adolescentis		15.51	0.21	XYL 92			
01775002	**Good identification**						**01775002**
Actinomyces israelii		100.0	0.37	SUC 98			
01775100	**Low discrimination**						**01775100**
					GRA	SPO	
Actinomyces israelii		82.23	0.44	SUC 98	99	1	
Fusobacterium mortiferum		12.04	0.27		1	1	
Clostridium butyricum		4.08	0.01		99	99	
01775102	**Low discrimination**						**01775102**
					SPO		
Actinomyces israelii		89.76	0.47	SUC 98	1		
Bifidobacterium adolescentis		8.12	0.23		1		
Clostridium butyricum		1.16	-.07		99		
01775122	**Acceptable identification**						**01775122**
					GRA		
Parabacteroides distasonis		91.34	0.42	MNS 98	1		
Actinomyces israelii		7.08	0.27		99		
Prevotella oralis		1.57	-.01		1		
01775132	**Good identification**						**01775132**
Parabacteroides distasonis		100.0	0.37	MNS 98			
01775140	**Good identification**						**01775140**
Actinomyces israelii		100.0	0.37	SUC 98			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
01775142	**Good identification**					**01775142**
Actinomyces israelii		97.21	0.40	SUC 98		
Bifidobacterium adolescentis		2.79	0.07			
01775322	**Good identification**					**01775322**
Parabacteroides distasonis		100.0	0.50	MNS 98		
01775332	**Good identification**					**01775332**
Parabacteroides distasonis		100.0	0.45	MNS 98		
01775402	**Good identification**					**01775402**
Actinomyces israelii		100.0	0.35	SUC 98		
01775500	**Low discrimination**					**01775500**
					SPO	
Actinomyces israelii		64.68	0.42	SUC 98	1	
Clostridium butyricum		35.32	0.17		99	
01775502	**Low discrimination**					**01775502**
					SPO	
Actinomyces israelii		53.16	0.45	SUC 98	1	
Bifidobacterium adolescentis		39.32	0.37		1	
Clostridium butyricum		7.53	0.10		99	
01775512	**Low discrimination**					**01775512**
Bifidobacterium adolescentis		84.00	0.38	SUC 99		
Actinomyces israelii		13.88	0.31	MNS 87		
Clostridium clostridioforme		2.12	0.08			
01775522	**Acceptable identification**					**01775522**
					GRA	
Parabacteroides distasonis		90.62	0.38	MNS 98	1	
Actinomyces israelii		9.38	0.25		99	
01775540	**Low discrimination**					**01775540**
					SPO	
Actinomyces israelii		89.12	0.35	SUC 98	1	
Clostridium butyricum		10.88	-.01		99	
01775542	**Low discrimination**					**01775542**
Actinomyces israelii		80.97	0.38	SUC 98		
Bifidobacterium adolescentis		19.03	0.21			
01775722	**Good identification**					**01775722**
Parabacteroides distasonis		98.05	0.46	MNS 98		
Clostridium clostridioforme		1.95	0.11			
01775732	**Low discrimination**					**01775732**
Parabacteroides distasonis		85.63	0.42	MNS 98		
Clostridium clostridioforme		14.37	0.23			
01777000	**Good identification**					**01777000**
Actinomyces israelii		100.0	0.47	SUC 98		
01777002	**Good identification**					**01777002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.50	SUC 98		
01777012	**Good identification**					**01777012**
Actinomyces israelii		100.0	0.35	SUC 98		
01777040	**Good identification**					**01777040**
Actinomyces israelii		100.0	0.40	SUC 98		
01777042	**Good identification**					**01777042**
Actinomyces israelii		100.0	0.43	SUC 98		
01777100	**Very good identification**					**01777100**
Actinomyces israelii		100.0	0.57	SUC 98		
01777102	**Very good identification**					**01777102**
Actinomyces israelii		98.71	0.60	SUC 98		
Bifidobacterium adolescentis		1.29	0.21			
01777110	**Good identification**					**01777110**
Actinomyces israelii		100.0	0.43	SUC 98		
01777112	**Acceptable identification**					**01777112**
Actinomyces israelii		90.35	0.46	SUC 98		
Bifidobacterium adolescentis		9.65	0.23	MLZ 13		
01777120	**Good identification**					**01777120**
Actinomyces israelii		100.0	0.36	SUC 98		
01777122	**Good identification**					**01777122**
Actinomyces israelii		100.0	0.39	SUC 98		
01777140	**Good identification**					**01777140**
Actinomyces israelii		100.0	0.50	SUC 98		
01777142	**Good identification**					**01777142**
Actinomyces israelii		100.0	0.53	SUC 98		
01777150	**Good identification**					**01777150**
Actinomyces israelii		100.0	0.36	SUC 98		
01777152	**Good identification**					**01777152**
Actinomyces israelii		96.72	0.39	SUC 98		
Bifidobacterium adolescentis		3.28	0.07	MLZ 13		
01777300	**Good identification**					**01777300**
Actinomyces israelii		100.0	0.38	SUC 98		
01777302	**Good identification**					**01777302**
Actinomyces israelii		100.0	0.41	SUC 98		
01777400	**Good identification**					**01777400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.45	SUC 98			
01777402	**Good identification**						**01777402**
Actinomyces israelii		100.0	0.48	SUC 98			
01777440	**Good identification**						**01777440**
Actinomyces israelii		100.0	0.38	SUC 98			
01777442	**Good identification**						**01777442**
Actinomyces israelii		100.0	0.41	SUC 98			
01777500	**Good identification**						**01777500**
					SPO		
Actinomyces israelii		98.94	0.55	SUC 98	1		
Clostridium butyricum		1.06	0.01		99		
01777502	**Good identification**						**01777502**
					SPO		
Actinomyces israelii		90.13	0.58	SUC 98	1		
Bifidobacterium adolescentis		9.63	0.35		1		
Clostridium butyricum		0.25	-.07		99		
01777510	**Good identification**						**01777510**
Actinomyces israelii		100.0	0.41	SUC 98			
01777512	**Low discrimination**						**01777512**
Actinomyces israelii		52.60	0.44	SUC 98			
Bifidobacterium adolescentis		45.95	0.37	MLZ 13			
Bifidobacterium dentium		1.46	-.07				
01777522	**Good identification**						**01777522**
Actinomyces israelii		100.0	0.38	SUC 98			
01777540	**Good identification**						**01777540**
Actinomyces israelii		100.0	0.48	SUC 98			
01777542	**Good identification**						**01777542**
Actinomyces israelii		96.72	0.51	SUC 98			
Bifidobacterium adolescentis		3.28	0.20				
01777552	**Low discrimination**						**01777552**
Actinomyces israelii		78.27	0.37	SUC 98			
Bifidobacterium adolescentis		21.73	0.21	MLZ 13			
01777700	**Good identification**						**01777700**
Actinomyces israelii		100.0	0.36	SUC 98			
01777702	**Good identification**						**01777702**
Actinomyces israelii		100.0	0.39	SUC 98			
02000000	**Low discrimination**						**02000000**
					COC	SPO	
Clostridium histolyticum		13.39	0.66	SUC 1	1	90	
Clostridium limosum		13.39	0.66		1	99	
Finegoldia magna		13.39	0.66		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
02000001	**Low discrimination**						**02000001**
					COC	GRA	
Campylobacter ureolyticus		70.53	0.66	SUC 1	1	1	
Anaerococcus prevotii		14.27	0.58		99	99	
Peptococcus niger		7.68	0.49		99	99	
02000002	**Low discrimination**						**02000002**
					COC	GRA	
Atopobium minutum		25.20	0.71	SUC 5	1	99	
Peptococcus niger		24.66	0.53		99	99	
Acidaminococcus fermentans		20.08	0.50		99	1	
02000003	**Low discrimination**						**02000003**
					COC	GRA	
Anaerococcus prevotii		58.92	0.47	SUC 1	99	99	
Peptococcus niger		22.38	0.36		99	99	
Campylobacter ureolyticus		11.76	0.32		1	1	
02000012	**Low discrimination**						**02000012**
					COC	GRA	
Atopobium minutum		25.14	0.37	SUC 5	1	99	
Peptococcus niger		24.60	0.19	MLZ 1	99	99	
Acidaminococcus fermentans		20.03	0.15		99	1	
02000020	**Low discrimination**						**02000020**
					COC	SPO	
Peptococcus niger		36.09	0.50	SUC 1	99	1	
Atopobium minutum		29.29	0.66	NAG 10	1	1	
Clostridium histolyticum		4.67	0.32		1	90	
02000022	**Low discrimination**						**02000022**
					COC	GRA	
Pseudoflavonifractor capillosus		87.67	0.66	SUC 1	1	1	
Atopobium minutum		8.88	0.63		1	99	
Peptococcus niger		2.90	0.37		99	99	
02000040	**Low discrimination**						**02000040**
Clostridium botulinum biovar A		23.16	0.51	GLU 90			
Clostridium histolyticum		10.06	0.32	SUC 7			
Clostridium limosum		10.06	0.32				
02000042	**Low discrimination**						**02000042**
					COC	GRA	
Atopobium minutum		23.02	0.37	SUC 5	1	99	
Peptococcus niger		22.52	0.19	SOR 1	99	99	
Acidaminococcus fermentans		18.34	0.15		99	1	
02000100	**Low discrimination**						**02000100**
					SPO		
Actinomyces meyeri		14.16	0.50	GLU 93	1		
Clostridium histolyticum		11.49	0.32	FRU 95	90		
Clostridium limosum		11.49	0.32		99		
02000102	**Low discrimination**						**02000102**
					COC	GRA	
Atopobium minutum		23.65	0.37	SUC 5	1	99	
Peptococcus niger		23.14	0.19	LAC 1	99	99	
Acidaminococcus fermentans		18.84	0.15		99	1	
02000120	**Low discrimination**						**02000120**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		44.01	0.35	GLU 93	1		
Peptococcus niger		20.47	0.15	FRU 95	99		
Atopobium minutum		16.61	0.32	NAG 12	1		
02000202	**Low discrimination**						**02000202**
					COC	GRA	
Atopobium minutum		25.46	0.37	SUC 5	1	99	
Peptococcus niger		24.91	0.19	RHA 1	99	99	
Acidaminococcus fermentans		20.28	0.15		99	1	
02000402	**Low discrimination**						**02000402**
					COC	GRA	
Atopobium minutum		25.34	0.37	SUC 5	1	99	
Peptococcus niger		24.79	0.19	ARA 1	99	99	
Acidaminococcus fermentans		20.18	0.15		99	1	
02000500	**Low discrimination**						**02000500**
					SPO		
Actinomyces meyeri		80.81	0.40	GLU 93	1		
Clostridium histolyticum		2.65	-.03	FRU 95	90		
Clostridium limosum		2.65	-.03		99		
02001000	**Low discrimination**						**02001000**
					SPO		
Atopobium minutum		43.35	0.66	SUC 5	1		
Clostridium histolyticum		6.91	0.32		90		
Clostridium limosum		6.91	0.32		99		
02001002	**Good identification**						**02001002**
					COC	GRA	
Atopobium minutum		90.58	0.63	SUC 5	1	99	
Peptococcus niger		2.69	0.19		99	99	
Acidaminococcus fermentans		2.19	0.15		99	1	
02001020	**Low discrimination**						**02001020**
					COC		
Atopobium minutum		88.95	0.58	SUC 5	1		
Peptococcus niger		3.32	0.15		99		
Actinomyces meyeri		2.68	0.28		1		
02001022	**Low discrimination**						**02001022**
					COC	GRA	
Pseudoflavonifractor capillosus		76.50	0.50	SUC 1	1	1	
Atopobium minutum		23.24	0.55	GAL 10	1	99	
Peptococcus niger		0.23	0.03		99	99	
02001100	**Acceptable identification**						**02001100**
					SPO		
Actinomyces meyeri		92.64	0.53	GLU 93	1		
Atopobium minutum		3.18	0.32	FRU 95	1		
Clostridium histolyticum		0.51	-.03		90		
02001120	**Acceptable identification**						**02001120**
					GRA		
Actinomyces meyeri		90.66	0.38	GLU 93	99		
Atopobium minutum		7.60	0.24	FRU 95	99		
Prevotella melaninogenica		1.73	0.01	NAG 12	1		
02001500	**Good identification**						**02001500**
Actinomyces meyeri		98.94	0.43	GLU 93			
Lactobacillus fermentum		1.06	0.20	FRU 95			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
02002002	**Low discrimination**						**02002002**
					COC	GRA	
Atopobium minutum		23.66	0.37	SUC 5	1	99	
Peptococcus niger		23.15	0.19	MAN 1	99	99	
Acidaminococcus fermentans		18.85	0.15		99	1	
02002040	**Low discrimination**						**02002040**
					COC	GRA	
Megasphaera elsdenii		70.21	0.38	GLU 99	99	1	
Clostridium botulinum biovar A		7.19	0.17	FRU 95	1	99	
Clostridium histolyticum		3.12	-.03		1	99	
02004002	**Low discrimination**						**02004002**
					COC	GRA	
Atopobium minutum		24.55	0.37	SUC 5	1	99	
Peptococcus niger		24.02	0.19	XYL 1	99	99	
Acidaminococcus fermentans		19.56	0.15		99	1	
02004100	**Good identification**						**02004100**
					SPO		
Actinomyces meyeri		97.03	0.55	GLU 93	1		
Clostridium histolyticum		0.41	-.03	FRU 95	90		
Clostridium limosum		0.41	-.03		99		
02004120	**Good identification**						**02004120**
Actinomyces meyeri		100.0	0.40	GLU 93			
02004500	**Good identification**						**02004500**
Actinomyces meyeri		100.0	0.45	GLU 93			
02005000	**Low discrimination**						**02005000**
					SPO		
Actinomyces meyeri		86.42	0.48	GLU 93	1		
Atopobium minutum		6.11	0.32	FRU 95	1		
Clostridium histolyticum		0.97	-.03		90		
02005100	**Very good identification**						**02005100**
Actinomyces meyeri		99.92	0.58	GLU 93			
Lactobacillus fermentum		0.08	0.16	FRU 95			
02005120	**Good identification**						**02005120**
Actinomyces meyeri		100.0	0.43	GLU 93			
02005400	**Good identification**						**02005400**
Actinomyces meyeri		99.21	0.37	GLU 93			
Lactobacillus fermentum		0.79	0.12	FRU 95			
02005500	**Good identification**						**02005500**
Actinomyces meyeri		99.68	0.48	GLU 93			
Lactobacillus fermentum		0.32	0.16	FRU 95			
02010000	**Low discrimination**						**02010000**
					COC	GRA	
Atopobium minutum		43.72	0.74	SUC 5	1	99	
Megasphaera elsdenii		16.28	0.63		99	1	
Actinomyces meyeri		14.72	0.62		1	99	
02010002	**Low discrimination**						**02010002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Atopobium minutum		89.08	0.71	SUC 5	1	99		
Megasphaera elsdenii		5.53	0.47		99	1		
Actinomyces meyeri		1.39	0.36		1	99		
02010003	**Low discrimination**							**02010003**
					COC			
Atopobium minutum		66.07	0.36	SUC 5	1			
Anaerococcus prevotii		18.23	0.13	URE 1	99			
Peptococcus niger		6.93	0.02		99			
02010010	**Low discrimination**							**02010010**
					COC	GRA		
Atopobium minutum		43.18	0.39	SUC 5	1	99		
Megasphaera elsdenii		16.08	0.29	MLZ 1	99	1		
Actinomyces meyeri		14.54	0.28		1	99		
02010012	**Acceptable identification**							**02010012**
					COC	GRA		
Atopobium minutum		92.79	0.36	SUC 5	1	99		
Megasphaera elsdenii		5.76	0.12	MLZ 1	99	1		
Actinomyces meyeri		1.45	0.02		1	99		
02010020	**Low discrimination**							**02010020**
					SPO			
Atopobium minutum		81.06	0.65	SUC 5	1			
Actinomyces meyeri		11.16	0.47		1			
Clostridium cadaveris		2.79	0.30		99			
02010022	**Low discrimination**							**02010022**
					GRA			
Pseudoflavonifractor capillosus		53.96	0.50	SUC 1	1			
Atopobium minutum		45.39	0.62	FRU 10	99			
Actinomyces meyeri		0.29	0.21		99			
02010040	**Low discrimination**							**02010040**
					COC	SPO	GRA	
Megasphaera elsdenii		88.04	0.60	GLU 99	99	1	1	
Atopobium minutum		3.58	0.39		1	1	99	
Clostridium botulinum biovar A		3.00	0.30		1	99	99	
02010042	**Low discrimination**							**02010042**
					COC	GRA		
Megasphaera elsdenii		78.62	0.44	GLU 99	99	1		
Atopobium minutum		19.19	0.36	BGL 10	1	99		
Pseudoramibacter alactolyticus		1.89	0.05		1	99		
02010100	**Very good identification**							**02010100**
					GRA			
Actinomyces meyeri		97.81	0.72	GLU 93	99			
Fusobacterium mortiferum		0.75	0.45		1			
Atopobium minutum		0.73	0.39		99			
02010101	**Good identification**							**02010101**
					GRA			
Actinomyces meyeri		96.62	0.38	GLU 93	99			
Campylobacter ureolyticus		2.64	-.03	URE 1	1			
Fusobacterium mortiferum		0.74	0.11		1			
02010102	**Low discrimination**							**02010102**
					GRA			
Actinomyces meyeri		82.91	0.46	GLU 93	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		13.41	0.36	BGL 3	99		
Fusobacterium mortiferum		2.29	0.29		1		
02010110	**Good identification**						**02010110**
					GRA		
Actinomyces meyeri		99.24	0.38	GLU 93	99		
Fusobacterium mortiferum		0.76	0.11	MLZ 1	1		
02010120	**Very good identification**						**02010120**
					GRA		
Actinomyces meyeri		98.08	0.57	GLU 93	99		
Atopobium minutum		1.80	0.31	NAG 12	99		
Fusobacterium mortiferum		0.06	0.11		1		
02010140	**Low discrimination**						**02010140**
					COC	GRA	
Actinomyces meyeri		83.89	0.38	GLU 93	1	99	
Megasphaera elsdenii		15.47	0.26	SOR 1	99	1	
Fusobacterium mortiferum		0.64	0.11		1	1	
02010200	**Low discrimination**						**02010200**
					COC	GRA	
Atopobium minutum		44.14	0.39	SUC 5	1	99	
Megasphaera elsdenii		16.44	0.29	RHA 1	99	1	
Actinomyces meyeri		14.86	0.28		1	99	
02010202	**Acceptable identification**						**02010202**
					COC	GRA	
Atopobium minutum		92.79	0.36	SUC 5	1	99	
Megasphaera elsdenii		5.76	0.12	RHA 1	99	1	
Actinomyces meyeri		1.45	0.02		1	99	
02010300	**Good identification**						**02010300**
					GRA		
Actinomyces meyeri		99.24	0.38	GLU 93	99		
Fusobacterium mortiferum		0.76	0.11	RHA 1	1		
02010400	**Low discrimination**						**02010400**
					COC	GRA	
Actinomyces meyeri		80.96	0.51	GLU 93	1	99	
Atopobium minutum		9.72	0.39		1	99	
Megasphaera elsdenii		3.62	0.29		99	1	
02010402	**Low discrimination**						**02010402**
					COC	GRA	
Atopobium minutum		67.81	0.36	SUC 5	1	99	
Actinomyces meyeri		26.21	0.26	ARA 1	1	99	
Megasphaera elsdenii		4.21	0.12		99	1	
02010420	**Low discrimination**						**02010420**
					SPO		
Actinomyces meyeri		76.72	0.37	GLU 93	1		
Atopobium minutum		22.51	0.31	NAG 12	1		
Clostridium cadaveris		0.77	-.04		99		
02010500	**Very good identification**						**02010500**
					GRA		
Actinomyces meyeri		99.88	0.62	GLU 93	99		
Lactobacillus fermentum		0.09	0.20		99		
Fusobacterium mortiferum		0.03	0.11		1		
02010502	**Good identification**						**02010502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		97.33	0.36	GLU 93		
Lactobacillus fermentum		2.67	0.20	BGL 3		
02010520	**Good identification**					**02010520**
Actinomyces meyeri		100.0	0.47	GLU 93		
02011000	**Low discrimination**					**02011000**
Actinomyces meyeri		57.75	0.65	GLU 93		
Atopobium minutum		38.12	0.65			
Lactobacillus fermentum		1.98	0.50			
02011002	**Low discrimination**					**02011002**
Atopobium minutum		86.53	0.62	SUC 5		
Lactobacillus fermentum		6.76	0.50			
Actinomyces meyeri		6.08	0.39			
02011020	**Low discrimination**					**02011020**
Atopobium minutum		59.63	0.57	SUC 5		
Actinomyces meyeri		36.95	0.50			
Lactobacillus fermentum		3.10	0.42			
02011022	**Low discrimination**					**02011022**
					GRA	
Atopobium minutum		66.53	0.54	SUC 5	99	
Pseudoflavonifractor capillosus		26.36	0.33		1	
Lactobacillus fermentum		5.19	0.42		99	
02011100	**Very good identification**					**02011100**
					GRA	
Actinomyces meyeri		97.72	0.75	GLU 93	99	
Lactobacillus fermentum		1.37	0.54		99	
Fusobacterium mortiferum		0.75	0.48		1	
02011101	**Good identification**					**02011101**
					SPO	
Actinomyces meyeri		96.05	0.41	GLU 93	1	
Clostridium chauvoei		1.87	0.09	URE 1	99	
Lactobacillus fermentum		1.35	0.20		1	
02011102	**Low discrimination**					**02011102**
Actinomyces meyeri		65.93	0.49	GLU 93		
Lactobacillus fermentum		29.88	0.54	BGL 3		
Atopobium minutum		2.37	0.28			
02011110	**Good identification**					**02011110**
					GRA	
Actinomyces meyeri		97.88	0.41	GLU 93	99	
Lactobacillus fermentum		1.37	0.20	MLZ 1	99	
Fusobacterium mortiferum		0.75	0.14		1	
02011120	**Very good identification**					**02011120**
					GRA	
Actinomyces meyeri		95.42	0.60	GLU 93	99	
Lactobacillus fermentum		3.27	0.46	NAG 12	99	
Prevotella melaninogenica		0.86	0.17		1	
02011122	**Low discrimination**					**02011122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		48.45	0.46	GLU 99		
Actinomyces meyeri		43.74	0.34			
Atopobium minutum		3.84	0.20			
02011140	**Good identification**					**02011140**
					GRA	
Actinomyces meyeri		97.88	0.41	GLU 93	99	
Lactobacillus fermentum		1.37	0.20	SOR 1	99	
Fusobacterium mortiferum		0.75	0.14		1	
02011300	**Good identification**					**02011300**
					GRA	
Actinomyces meyeri		97.88	0.41	GLU 93	99	
Lactobacillus fermentum		1.37	0.20	RHA 1	99	
Fusobacterium mortiferum		0.75	0.14		1	
02011400	**Low discrimination**					**02011400**
Actinomyces meyeri		85.90	0.54	GLU 93		
Lactobacillus fermentum		11.81	0.50			
Atopobium minutum		2.29	0.31			
02011402	**Low discrimination**					**02011402**
Lactobacillus fermentum		73.83	0.50	GLU 99		
Actinomyces meyeri		16.62	0.29			
Atopobium minutum		9.55	0.28			
02011420	**Low discrimination**					**02011420**
Actinomyces meyeri		71.37	0.40	GLU 93		
Lactobacillus fermentum		23.98	0.42	NAG 12		
Atopobium minutum		4.65	0.23			
02011422	**Low discrimination**					**02011422**
Lactobacillus fermentum		78.57	0.42	GLU 99		
Atopobium minutum		10.17	0.20			
Actinomyces meyeri		7.23	0.14			
02011500	**Good identification**					**02011500**
					GRA	
Actinomyces meyeri		94.66	0.65	GLU 93	99	
Lactobacillus fermentum		5.31	0.54		99	
Fusobacterium mortiferum		0.03	0.14		1	
02011502	**Low discrimination**					**02011502**
Lactobacillus fermentum		64.45	0.54	GLU 99		
Actinomyces meyeri		35.56	0.39			
02011520	**Low discrimination**					**02011520**
Actinomyces meyeri		87.95	0.50	GLU 93		
Lactobacillus fermentum		12.05	0.46	NAG 12		
02011522	**Low discrimination**					**02011522**
Lactobacillus fermentum		81.59	0.46	GLU 99		
Actinomyces meyeri		18.41	0.24			
02012000	**Good identification**					**02012000**
					COC GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		90.10	0.63	GLU 99	99	1	
Pseudoramibacter alactolyticus		3.10	0.27		1	99	
Atopobium minutum		2.44	0.39		1	99	
02012002	**Low discrimination**						**02012002**
					COC	GRA	
Megasphaera elsdenii		83.13	0.47	GLU 99	99	1	
Atopobium minutum		13.53	0.36	BGL 10	1	99	
Pseudoramibacter alactolyticus		2.86	0.11		1	99	
02012040	**Very good identification**						**02012040**
					COC	GRA	
Megasphaera elsdenii		98.73	0.60	GLU 99	99	1	
Pseudoramibacter alactolyticus		1.28	0.17		1	99	
02012042	**Good identification**						**02012042**
					COC	GRA	
Megasphaera elsdenii		98.73	0.44	GLU 99	99	1	
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99	
02012100	**Low discrimination**						**02012100**
					COC	GRA	
Actinomyces meyeri		76.62	0.38	GLU 93	1	99	
Megasphaera elsdenii		21.19	0.29	MAN 1	99	1	
Propionibacterium propionicum		0.88	0.15		1	99	
02013100	**Good identification**						**02013100**
					GRA		
Actinomyces meyeri		97.15	0.41	GLU 93	99		
Lactobacillus fermentum		1.36	0.20	MAN 1	99		
Fusobacterium mortiferum		0.75	0.14		1		
02014000	**Very good identification**						**02014000**
					GRA		
Actinomyces meyeri		96.49	0.67	GLU 93	99		
Atopobium minutum		1.49	0.39		99		
Fusobacterium varium		0.61	0.19		1		
02014002	**Low discrimination**						**02014002**
					COC	GRA	
Actinomyces meyeri		73.59	0.41	GLU 93	1	99	
Atopobium minutum		24.52	0.36	BGL 3	1	99	
Megasphaera elsdenii		1.52	0.12		99	1	
02014020	**Good identification**						**02014020**
					SPO		
Actinomyces meyeri		96.24	0.52	GLU 93	1		
Atopobium minutum		3.64	0.31	NAG 12	1		
Clostridium cadaveris		0.13	-.04		99		
02014100	**Very good identification**						**02014100**
					GRA		
Actinomyces meyeri		99.96	0.77	GLU 93	99		
Fusobacterium mortiferum		0.03	0.26		1		
Lactobacillus fermentum		0.01	0.16		99		
02014101	**Good identification**						**02014101**
Actinomyces meyeri		100.0	0.43	GLU 93			
02014102	**Good identification**						**02014102**
Actinomyces meyeri		99.79	0.51	GLU 93			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		0.21	0.16	BGL 3		
02014110	**Good identification**					**02014110**
Actinomyces meyeri		100.0	0.43	GLU 93		
02014120	**Very good identification**					**02014120**
Actinomyces meyeri		100.0	0.62	GLU 93		
02014122	**Good identification**					**02014122**
Actinomyces meyeri		100.0	0.36	GLU 93		
02014140	**Good identification**					**02014140**
					SPO	
Actinomyces meyeri		99.59	0.43	GLU 93	1	
Clostridium glycolicum		0.41	-.03	SOR 1	99	
02014300	**Good identification**					**02014300**
Actinomyces meyeri		100.0	0.43	GLU 93		
02014400	**Very good identification**					**02014400**
Actinomyces meyeri		99.94	0.56	GLU 93		
Lactobacillus fermentum		0.06	0.12			
02014420	**Good identification**					**02014420**
Actinomyces meyeri		100.0	0.41	GLU 93		
02014500	**Very good identification**					**02014500**
Actinomyces meyeri		99.97	0.67	GLU 93		
Lactobacillus fermentum		0.03	0.16			
02014502	**Good identification**					**02014502**
Actinomyces meyeri		99.18	0.41	GLU 93		
Lactobacillus fermentum		0.82	0.16	BGL 3		
02014520	**Good identification**					**02014520**
Actinomyces meyeri		100.0	0.52	GLU 93		
02015000	**Very good identification**					**02015000**
Actinomyces meyeri		98.59	0.70	GLU 93		
Lactobacillus fermentum		1.03	0.46			
Atopobium minutum		0.34	0.31			
02015001	**Good identification**					**02015001**
Actinomyces meyeri		98.97	0.35	GLU 93		
Lactobacillus fermentum		1.03	0.12	URE 1		
02015002	**Low discrimination**					**02015002**
Actinomyces meyeri		70.92	0.44	GLU 93		
Lactobacillus fermentum		23.84	0.46	BGL 3		
Atopobium minutum		5.25	0.28			
02015010	**Good identification**					**02015010**
Actinomyces meyeri		98.97	0.35	GLU 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		1.03	0.12	MLZ 1		
02015020	**Good identification**					**02015020**
Actinomyces meyeri		96.73	0.55	GLU 93		
Lactobacillus fermentum		2.46	0.38	NAG 12		
Atopobium minutum		0.81	0.23			
02015040	**Low discrimination**					**02015040**
					SPO	
Actinomyces meyeri		88.44	0.35	GLU 93	1	
Clostridium glycolicum		10.64	0.15	SOR 1	99	
Lactobacillus fermentum		0.92	0.12		1	
02015100	**Very good identification**					**02015100**
					GRA	
Actinomyces meyeri		99.55	0.80	GLU 93	99	
Lactobacillus fermentum		0.42	0.50		99	
Fusobacterium mortiferum		0.03	0.29		1	
02015101	**Good identification**					**02015101**
Actinomyces meyeri		99.58	0.46	GLU 93		
Lactobacillus fermentum		0.42	0.16	URE 1		
02015102	**Low discrimination**					**02015102**
					GRA	
Actinomyces meyeri		87.86	0.54	GLU 93	99	
Lactobacillus fermentum		12.05	0.50	BGL 3	99	
Fusobacterium mortiferum		0.09	0.13		1	
02015110	**Good identification**					**02015110**
Actinomyces meyeri		99.58	0.46	GLU 93		
Lactobacillus fermentum		0.42	0.16	MLZ 1		
02015120	**Very good identification**					**02015120**
Actinomyces meyeri		98.97	0.65	GLU 93		
Lactobacillus fermentum		1.03	0.42	NAG 12		
02015122	**Low discrimination**					**02015122**
Actinomyces meyeri		74.90	0.39	GLU 93		
Lactobacillus fermentum		25.10	0.42	NAG 12		
02015140	**Good identification**					**02015140**
Actinomyces meyeri		99.58	0.46	GLU 93		
Lactobacillus fermentum		0.42	0.16	SOR 1		
02015200	**Good identification**					**02015200**
Actinomyces meyeri		98.97	0.35	GLU 93		
Lactobacillus fermentum		1.03	0.12	RHA 1		
02015300	**Good identification**					**02015300**
Actinomyces meyeri		99.58	0.46	GLU 93		
Lactobacillus fermentum		0.42	0.16	RHA 1		
02015400	**Very good identification**					**02015400**
Actinomyces meyeri		96.01	0.59	GLU 93		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Lactobacillus fermentum	3.99	0.46			
02015402	**Low discrimination**					**02015402**
	Lactobacillus fermentum	57.35	0.46	GLU 99		
	Actinomyces meyeri	42.66	0.33			
02015420	**Acceptable identification**					**02015420**
	Actinomyces meyeri	90.77	0.45	GLU 93		
	Lactobacillus fermentum	9.23	0.38	NAG 12		
02015422	**Low discrimination**					**02015422**
	Lactobacillus fermentum	76.67	0.38	GLU 99		
	Actinomyces meyeri	23.33	0.19			
02015500	**Very good identification**					**02015500**
	Actinomyces meyeri	98.33	0.70	GLU 93		
	Lactobacillus fermentum	1.67	0.50			
02015501	**Good identification**					**02015501**
	Actinomyces meyeri	98.33	0.35	GLU 93		
	Lactobacillus fermentum	1.67	0.16	URE 1		
02015502	**Low discrimination**					**02015502**
	Actinomyces meyeri	64.58	0.44	GLU 93		
	Lactobacillus fermentum	35.42	0.50	BGL 3		
02015510	**Good identification**					**02015510**
	Actinomyces meyeri	98.33	0.35	GLU 93		
	Lactobacillus fermentum	1.67	0.16	MLZ 1		
02015520	**Good identification**					**02015520**
	Actinomyces meyeri	96.02	0.55	GLU 93		
	Lactobacillus fermentum	3.98	0.42	NAG 12		
02015522	**Low discrimination**					**02015522**
	Lactobacillus fermentum	57.27	0.42	GLU 99		
	Actinomyces meyeri	42.73	0.29			
02015540	**Good identification**					**02015540**
	Actinomyces meyeri	98.33	0.35	GLU 93		
	Lactobacillus fermentum	1.67	0.16	SOR 1		
02015700	**Good identification**					**02015700**
	Actinomyces meyeri	98.33	0.35	GLU 93		
	Lactobacillus fermentum	1.67	0.16	RHA 1		
02016100	**Good identification**					**02016100**
	Actinomyces meyeri	100.0	0.43	GLU 93		
02017000	**Good identification**					**02017000**
	Actinomyces meyeri	98.97	0.35	GLU 93		
	Lactobacillus fermentum	1.03	0.12	MAN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
02017100	**Good identification**							**02017100**
Actinomyces meyeri		99.58	0.46	GLU 93				
Lactobacillus fermentum		0.42	0.16	MAN 1				
02017500	**Good identification**							**02017500**
Actinomyces meyeri		98.33	0.35	GLU 93				
Lactobacillus fermentum		1.67	0.16	MAN 1				
02020000	**Low discrimination**							**02020000**
Clostridium botulinum biovar A		20.40	0.50	GLU 90				
Clostridium histolyticum		10.59	0.32	SUC 7				
Clostridium limosum		10.59	0.32					
02020002	**Low discrimination**							**02020002**
					COC	GRA		
Atopobium minutum		22.58	0.37	SUC 5	1	99		
Peptococcus niger		22.09	0.19	TRE 1	99	99		
Acidaminococcus fermentans		17.99	0.15		99	1		
02020040	**Low discrimination**							**02020040**
Clostridium botulinum biovar A		88.41	0.41	GLU 90				
Clostridium histolyticum		1.55	-.03	SUC 7				
Clostridium limosum		1.55	-.03					
02030000	**Low discrimination**							**02030000**
					COC	SPO	GRA	
Atopobium minutum		29.33	0.39	SUC 5	1	1	99	
Clostridium botulinum biovar A		20.55	0.29	TRE 1	1	99	99	
Megasphaera elsdenii		10.92	0.29		99	1	1	
02030002	**Low discrimination**							**02030002**
					SPO	CAT		
Atopobium minutum		80.40	0.36	SUC 5	1	1		
Propionibacterium granulosum		6.26	0.20	TRE 1	1	90		
Clostridium sporogenes		5.33	0.06		99	1		
02030100	**Low discrimination**							**02030100**
					GRA			
Actinomyces meyeri		81.49	0.38	GLU 93	99			
Fusobacterium mortiferum		18.51	0.36	TRE 1	1			
02031100	**Low discrimination**							**02031100**
					GRA			
Actinomyces meyeri		80.57	0.41	GLU 93	99			
Fusobacterium mortiferum		18.30	0.39	TRE 1	1			
Lactobacillus fermentum		1.13	0.20		99			
02034100	**Good identification**							**02034100**
					GRA			
Actinomyces meyeri		99.13	0.43	GLU 93	99			
Fusobacterium mortiferum		0.87	0.17	TRE 1	1			
02035000	**Good identification**							**02035000**
					GRA			
Actinomyces meyeri		97.51	0.35	GLU 93	99			
Fusobacterium mortiferum		1.47	0.14	TRE 1	1			
Lactobacillus fermentum		1.01	0.12		99			
02035100	**Good identification**							**02035100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces meyeri		98.71	0.46	GLU 93	99		
Fusobacterium mortiferum		0.87	0.20	TRE 1	1		
Lactobacillus fermentum		0.42	0.16		99		
02035500	**Good identification**						**02035500**
Actinomyces meyeri		98.33	0.35	GLU 93			
Lactobacillus fermentum		1.67	0.16	TRE 1			
02040002	**Low discrimination**						**02040002**
					COC	GRA	
Atopobium minutum		25.46	0.37	SUC 5	1	99	
Peptococcus niger		24.91	0.19	CEL 1	99	99	
Acidaminococcus fermentans		20.28	0.15		99	1	
02050000	**Low discrimination**						**02050000**
					COC	GRA	
Atopobium minutum		41.46	0.39	SUC 5	1	99	
Megasphaera elsdenii		15.44	0.29	CEL 1	99	1	
Actinomyces meyeri		13.96	0.28		1	99	
02050002	**Acceptable identification**						**02050002**
					COC	GRA	
Atopobium minutum		90.74	0.36	SUC 5	1	99	
Megasphaera elsdenii		5.63	0.12	CEL 1	99	1	
Fusobacterium mortiferum		2.21	0.14		1	1	
02050100	**Low discrimination**						**02050100**
					GRA		
Actinomyces meyeri		79.78	0.38	GLU 93	99		
Fusobacterium mortiferum		20.22	0.37	CEL 1	1		
02051100	**Low discrimination**						**02051100**
					GRA		
Actinomyces meyeri		78.90	0.41	GLU 93	99		
Fusobacterium mortiferum		20.00	0.40	CEL 1	1		
Lactobacillus fermentum		1.11	0.20		99		
02054100	**Good identification**						**02054100**
					GRA		
Actinomyces meyeri		99.03	0.43	GLU 93	99		
Fusobacterium mortiferum		0.97	0.18	CEL 1	1		
02055000	**Good identification**						**02055000**
					GRA		
Actinomyces meyeri		97.35	0.35	GLU 93	99		
Fusobacterium mortiferum		1.64	0.14	CEL 1	1		
Lactobacillus fermentum		1.01	0.12		99		
02055100	**Good identification**						**02055100**
					GRA		
Actinomyces meyeri		98.61	0.46	GLU 93	99		
Fusobacterium mortiferum		0.97	0.21	CEL 1	1		
Lactobacillus fermentum		0.42	0.16		99		
02055500	**Good identification**						**02055500**
Actinomyces meyeri		98.33	0.35	GLU 93			
Lactobacillus fermentum		1.67	0.16	CEL 1			
02100000	**Low discrimination**						**02100000**
					COC	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptostreptococcus anaerobius		53.46	0.61	SUC 1	99	1		
Clostridium botulinum biovar A		19.12	0.58		1	99		
Clostridium histolyticum		3.03	0.32		1	90		
02100040	**Low discrimination**							**02100040**
					COC	GRA	SPO	
Clostridium botulinum biovar A		74.41	0.49	GLU 90	1	99	99	
Megasphaera elsdenii		13.46	0.41	SUC 7	99	1	1	
Peptostreptococcus anaerobius		7.04	0.27		99	99	1	
02100100	**Low discrimination**							**02100100**
					COC	SPO		
Actinomyces meyeri		81.05	0.51	GLU 93	1	1		
Peptostreptococcus anaerobius		10.39	0.27	FRU 95	99	1		
Clostridium botulinum biovar A		3.72	0.24		1	99		
02100120	**Acceptable identification**							**02100120**
					GRA			
Actinomyces meyeri		92.89	0.36	GLU 93	99			
Prevotella bivia		3.87	-.03	FRU 95	1			
Prevotella melaninogenica		2.36	0.01	NAG 12	1			
02100500	**Good identification**							**02100500**
					COC			
Actinomyces meyeri		99.49	0.41	GLU 93	1			
Peptostreptococcus anaerobius		0.52	-.07	FRU 95	99			
02101000	**Low discrimination**							**02101000**
					COC	SPO		
Actinomyces meyeri		52.30	0.44	GLU 93	1	1		
Peptostreptococcus anaerobius		17.88	0.27	FRU 95	99	1		
Clostridium botulinum biovar A		6.40	0.24		1	99		
02101100	**Good identification**							**02101100**
					GRA			
Actinomyces meyeri		98.69	0.54	GLU 93	99			
Lactobacillus fermentum		0.71	0.28	FRU 95	99			
Prevotella melaninogenica		0.23	0.01		1			
02101101	**Acceptable identification**							**02101101**
					SPO			
Clostridium chauvoei		93.29	0.37	GLU 99	99			
Actinomyces meyeri		6.71	0.20	MNS 90	1			
02101500	**Good identification**							**02101500**
Actinomyces meyeri		97.22	0.44	GLU 93				
Lactobacillus fermentum		2.78	0.28	FRU 95				
02102000	**Low discrimination**							**02102000**
					COC	GRA	SPO	
Megasphaera elsdenii		61.26	0.44	GLU 99	99	1	1	
Peptostreptococcus anaerobius		21.35	0.27	FRU 95	99	99	1	
Clostridium botulinum biovar A		7.64	0.24		1	99	99	
02102040	**Acceptable identification**							**02102040**
					COC	GRA	SPO	
Megasphaera elsdenii		94.24	0.41	GLU 99	99	1	1	
Clostridium botulinum biovar A		5.26	0.15	FRU 95	1	99	99	
Peptostreptococcus anaerobius		0.50	-.07		99	99	1	
02104000	**Low discrimination**							**02104000**
					COC	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		66.76	0.46	GLU 93	1	1	
Peptostreptococcus anaerobius		17.64	0.27	FRU 95	99	1	
Clostridium botulinum biovar A		6.31	0.24		1	99	
02104100	**Good identification**						**02104100**
					COC		
Actinomyces meyeri		99.93	0.56	GLU 93	1		
Peptostreptococcus anaerobius		0.07	-.07	FRU 95	99		
02104120	**Good identification**						**02104120**
Actinomyces meyeri		100.0	0.41	GLU 93			
02104400	**Good identification**						**02104400**
					COC		
Actinomyces meyeri		98.94	0.35	GLU 93	1		
Peptostreptococcus anaerobius		1.06	-.07	FRU 95	99		
02104500	**Good identification**						**02104500**
Actinomyces meyeri		100.0	0.46	GLU 93			
02105000	**Good identification**						**02105000**
					COC		
Actinomyces meyeri		98.73	0.49	GLU 93	1		
Gemella morbillorum		0.57	0.09	FRU 95	99		
Lactobacillus fermentum		0.52	0.20		1		
02105100	**Very good identification**						**02105100**
Actinomyces meyeri		99.78	0.59	GLU 93			
Lactobacillus fermentum		0.22	0.24	FRU 95			
02105120	**Good identification**						**02105120**
					GRA		
Actinomyces meyeri		98.62	0.44	GLU 93	99		
Prevotella melaninogenica		0.86	0.01	FRU 95	1		
Lactobacillus fermentum		0.52	0.16	NAG 12	99		
02105400	**Good identification**						**02105400**
Actinomyces meyeri		97.92	0.38	GLU 93			
Lactobacillus fermentum		2.08	0.20	FRU 95			
02105500	**Good identification**						**02105500**
Actinomyces meyeri		99.14	0.49	GLU 93			
Lactobacillus fermentum		0.86	0.24	FRU 95			
02110000	**Low discrimination**						**02110000**
					COC	GRA	
Megasphaera elsdenii		49.98	0.66	GLU 99	99	1	
Actinomyces meyeri		33.97	0.63		1	99	
Prevotella intermedia		9.55	0.51		1	1	
02110002	**Low discrimination**						**02110002**
					COC	GRA	
Megasphaera elsdenii		72.54	0.50	GLU 99	99	1	
Actinomyces meyeri		13.72	0.37	BGL 10	1	99	
Atopobium minutum		7.87	0.36		1	99	
02110020	**Low discrimination**						**02110020**
					COC	GRA	
Actinomyces meyeri		81.03	0.48	GLU 93	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		8.83	0.32	NAG 12	99	1	
Atopobium minutum		5.27	0.31		1	99	
02110040	**Very good identification**						**02110040**
					COC	GRA	
Megasphaera elsdenii		95.49	0.63	GLU 99	99	1	
Clostridium botulinum biovar A		1.77	0.29		1	99	
Clostridium botulinum biovar B		1.25	0.24		1	99	
02110042	**Good identification**						**02110042**
					COC	GRA	
Megasphaera elsdenii		99.71	0.47	GLU 99	99	1	
Actinomyces meyeri		0.29	0.03	BGL 10	1	99	
02110100	**Very good identification**						**02110100**
					COC	GRA	
Actinomyces meyeri		99.37	0.73	GLU 93	1	99	
Megasphaera elsdenii		0.37	0.32		99	1	
Fusobacterium mortiferum		0.08	0.30		1	1	
02110101	**Good identification**						**02110101**
					SPO		
Actinomyces meyeri		97.48	0.39	GLU 93	1		
Clostridium chauvoei		2.52	0.09	URE 1	99		
02110102	**Good identification**						**02110102**
					COC	GRA	
Actinomyces meyeri		95.82	0.47	GLU 93	1	99	
Lactobacillus fermentum		1.75	0.28	BGL 3	1	99	
Megasphaera elsdenii		1.28	0.15		99	1	
02110110	**Good identification**						**02110110**
Actinomyces meyeri		100.0	0.39	GLU 93			
02110120	**Very good identification**						**02110120**
					GRA		
Actinomyces meyeri		98.68	0.58	GLU 93	99		
Prevotella melaninogenica		1.19	0.17	NAG 12	1		
Lactobacillus fermentum		0.14	0.20		99		
02110140	**Low discrimination**						**02110140**
					COC	GRA	
Actinomyces meyeri		79.66	0.39	GLU 93	1	99	
Megasphaera elsdenii		19.54	0.29	SOR 1	99	1	
Propionibacterium propionicum		0.81	0.15		1	99	
02110300	**Good identification**						**02110300**
Actinomyces meyeri		96.26	0.39	GLU 93			
Actinomyces odontolyticus		3.74	0.18	RHA 1			
02110400	**Acceptable identification**						**02110400**
					COC	GRA	
Actinomyces meyeri		91.92	0.52	GLU 93	1	99	
Megasphaera elsdenii		5.47	0.32		99	1	
Prevotella intermedia		1.04	0.17		1	1	
02110420	**Good identification**						**02110420**
Actinomyces meyeri		98.66	0.37	GLU 93			
Lactobacillus fermentum		1.34	0.16	NAG 12			
02110500	**Very good identification**						**02110500**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		99.78	0.63	GLU 93		
Lactobacillus fermentum		0.23	0.28			
02110502	**Acceptable identification**					**02110502**
Actinomyces meyeri		93.19	0.37	GLU 93		
Lactobacillus fermentum		6.81	0.28	BGL 3		
02110520	**Good identification**					**02110520**
Actinomyces meyeri		99.45	0.48	GLU 93		
Lactobacillus fermentum		0.55	0.20	NAG 12		
02111000	**Low discrimination**					**02111000**
					GRA	
Actinomyces meyeri		76.64	0.66	GLU 93	99	
Prevotella intermedia		14.37	0.51		1	
Lactobacillus fermentum		7.01	0.59		99	
02111002	**Low discrimination**					**02111002**
Lactobacillus fermentum		70.33	0.59	GLU 99		
Actinomyces meyeri		23.80	0.40			
Atopobium minutum		3.03	0.28			
02111020	**Low discrimination**					**02111020**
					GRA	
Actinomyces meyeri		77.08	0.51	GLU 93	99	
Lactobacillus fermentum		17.22	0.50	NAG 12	99	
Prevotella melaninogenica		2.48	0.17		1	
02111022	**Low discrimination**					**02111022**
Lactobacillus fermentum		80.61	0.50	GLU 99		
Actinomyces meyeri		11.16	0.25			
Atopobium minutum		3.48	0.20			
02111100	**Very good identification**					**02111100**
					GRA	
Actinomyces meyeri		96.15	0.76	GLU 93	99	
Lactobacillus fermentum		3.59	0.62		99	
Prevotella melaninogenica		0.11	0.17		1	
02111101	**Low discrimination**					**02111101**
					SPO	
Clostridium chauvoei		62.20	0.44	GLU 99	99	
Actinomyces meyeri		36.44	0.42	MNS 90	1	
Lactobacillus fermentum		1.36	0.28		1	
02111102	**Low discrimination**					**02111102**
					GRA	
Lactobacillus fermentum		54.29	0.62	GLU 99	99	
Actinomyces meyeri		45.04	0.50		99	
Prevotella melaninogenica		0.30	0.05		1	
02111110	**Good identification**					**02111110**
Actinomyces meyeri		96.41	0.42	GLU 93		
Lactobacillus fermentum		3.59	0.28	MLZ 1		
02111120	**Low discrimination**					**02111120**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		52.65	0.61	GLU 93	99		
Prevotella melaninogenica		41.87	0.51	NAG 12	1		
Lactobacillus fermentum		4.80	0.54		99		
02111121	**Low discrimination**						**02111121**
					GRA	SPO	
Clostridium chauvoei		68.87	0.35	GLU 99	99	99	
Actinomyces meyeri		16.51	0.27	MNS 90	99	1	
Prevotella melaninogenica		13.13	0.17		1	1	
02111122	**Low discrimination**						**02111122**
					GRA		
Prevotella melaninogenica		54.20	0.39	GLU 90	1		
Lactobacillus fermentum		32.61	0.54	MNS 90	99		
Actinomyces meyeri		11.07	0.35	RAF 90	99		
02111140	**Good identification**						**02111140**
Actinomyces meyeri		95.78	0.42	GLU 93			
Lactobacillus fermentum		3.57	0.28	SOR 1			
Propionibacterium propionicum		0.65	0.15				
02111300	**Acceptable identification**						**02111300**
Actinomyces meyeri		94.83	0.42	GLU 93			
Lactobacillus fermentum		3.54	0.28	RHA 1			
Actinomyces odontolyticus		1.64	0.15				
02111400	**Low discrimination**						**02111400**
					GRA		
Actinomyces meyeri		72.67	0.55	GLU 93	99		
Lactobacillus fermentum		26.57	0.59		99		
Prevotella intermedia		0.55	0.17		1		
02111402	**Good identification**						**02111402**
Lactobacillus fermentum		92.20	0.59	GLU 99			
Actinomyces meyeri		7.80	0.29				
02111420	**Low discrimination**						**02111420**
Actinomyces meyeri		52.81	0.40	GLU 93			
Lactobacillus fermentum		47.19	0.50	NAG 12			
02111422	**Good identification**						**02111422**
Lactobacillus fermentum		96.66	0.50	GLU 99			
Actinomyces meyeri		3.35	0.15				
02111500	**Low discrimination**						**02111500**
Actinomyces meyeri		87.02	0.66	GLU 93			
Lactobacillus fermentum		12.98	0.62				
02111502	**Low discrimination**						**02111502**
Lactobacillus fermentum		82.82	0.62	GLU 99			
Actinomyces meyeri		17.18	0.40				
02111520	**Low discrimination**						**02111520**
					GRA		
Actinomyces meyeri		71.39	0.51	GLU 93	99		
Lactobacillus fermentum		26.02	0.54	NAG 12	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		2.29	0.17		1		
02111522	**Low discrimination**						**02111522**
					GRA		
Lactobacillus fermentum		89.84	0.54	GLU 99	99		
Actinomyces meyeri		7.62	0.25		99		
Prevotella melaninogenica		1.51	0.05		1		
02112000	**Very good identification**						**02112000**
					COC	GRA	
Megasphaera elsdenii		98.98	0.66	GLU 99	99	1	
Actinomyces meyeri		0.68	0.28		1	99	
Prevotella intermedia		0.19	0.17		1	1	
02112002	**Good identification**						**02112002**
					COC	GRA	
Megasphaera elsdenii		99.44	0.50	GLU 99	99	1	
Propionibacterium propionicum		0.37	0.18	BGL 10	1	99	
Actinomyces meyeri		0.19	0.03		1	99	
02112040	**Very good identification**						**02112040**
					COC	GRA	
Megasphaera elsdenii		99.96	0.63	GLU 99	99	1	
Propionibacterium propionicum		0.04	0.15		1	99	
02112042	**Good identification**						**02112042**
					COC	GRA	
Megasphaera elsdenii		99.44	0.47	GLU 99	99	1	
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99	
02112100	**Low discrimination**						**02112100**
					COC	GRA	
Propionibacterium propionicum		42.33	0.49	NIT 86	1	99	
Actinomyces meyeri		42.16	0.39	GLU 99	1	99	
Megasphaera elsdenii		15.51	0.32		99	1	
02112102	**Good identification**						**02112102**
					COC	GRA	
Propionibacterium propionicum		95.45	0.52	NIT 86	1	99	
Megasphaera elsdenii		2.59	0.15	GLU 99	99	1	
Actinomyces meyeri		1.96	0.13		1	99	
02112103	**Good identification**						**02112103**
Propionibacterium propionicum		100.0	0.36	NIT 86			
02112140	**Low discrimination**						**02112140**
					COC	GRA	
Propionibacterium propionicum		79.72	0.49	NIT 86	1	99	
Megasphaera elsdenii		19.48	0.29	GLU 99	99	1	
Actinomyces meyeri		0.80	0.05		1	99	
02112142	**Good identification**						**02112142**
					COC	GRA	
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99	
Megasphaera elsdenii		1.78	0.12	GLU 99	99	1	
02112143	**Good identification**						**02112143**
Propionibacterium propionicum		100.0	0.36	NIT 86			
02113100	**Low discrimination**						**02113100**
Actinomyces meyeri		58.59	0.42	GLU 93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		39.22	0.49	MAN 1			
Lactobacillus fermentum		2.19	0.28				
02113102	**Acceptable identification**						**02113102**
Propionibacterium propionicum		93.64	0.52	NIT 86			
Lactobacillus fermentum		3.48	0.28	GLU 99			
Actinomyces meyeri		2.88	0.16				
02113103	**Good identification**						**02113103**
Propionibacterium propionicum		100.0	0.36	NIT 86			
02113140	**Good identification**						**02113140**
Propionibacterium propionicum		98.51	0.49	NIT 86			
Actinomyces meyeri		1.49	0.08	GLU 99			
02113142	**Good identification**						**02113142**
Propionibacterium propionicum		100.0	0.52	NIT 86			
02113143	**Good identification**						**02113143**
Propionibacterium propionicum		100.0	0.36	NIT 86			
02114000	**Very good identification**						**02114000**
					COC	GRA	
Actinomyces meyeri		98.77	0.68	GLU 93	1	99	
Megasphaera elsdenii		0.76	0.32		99	1	
Prevotella intermedia		0.14	0.17		1	1	
02114002	**Acceptable identification**						**02114002**
					COC	GRA	
Actinomyces meyeri		94.88	0.42	GLU 93	1	99	
Megasphaera elsdenii		2.61	0.15	BGL 3	99	1	
Lactobacillus fermentum		1.29	0.20		1	99	
02114020	**Good identification**						**02114020**
Actinomyces meyeri		99.90	0.53	GLU 93			
Lactobacillus fermentum		0.10	0.12	NAG 12			
02114100	**Very good identification**						**02114100**
Actinomyces meyeri		99.98	0.78	GLU 93			
Lactobacillus fermentum		0.02	0.24				
Actinomyces odontolyticus		0.01	0.08				
02114101	**Good identification**						**02114101**
Actinomyces meyeri		100.0	0.44	GLU 93			
02114102	**Good identification**						**02114102**
Actinomyces meyeri		99.45	0.52	GLU 93			
Lactobacillus fermentum		0.55	0.24	BGL 3			
02114110	**Good identification**						**02114110**
Actinomyces meyeri		100.0	0.44	GLU 93			
02114120	**Very good identification**						**02114120**
Actinomyces meyeri		99.96	0.63	GLU 93			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Lactobacillus fermentum	0.04	0.16	NAG 12		
02114122	**Good identification**					**02114122**
	Actinomyces meyeri	98.67	0.37	GLU 93		
	Lactobacillus fermentum	1.33	0.16	NAG 12		
02114140	**Good identification**					**02114140**
					SPO	
	Actinomyces meyeri	99.61	0.44	GLU 93	1	
	Clostridium glycolicum	0.39	-.03	SOR 1	99	
02114300	**Good identification**					**02114300**
	Actinomyces meyeri	99.02	0.44	GLU 93		
	Actinomyces odontolyticus	0.98	0.13	RHA 1		
02114400	**Very good identification**					**02114400**
	Actinomyces meyeri	99.83	0.57	GLU 93		
	Lactobacillus fermentum	0.17	0.20			
02114420	**Good identification**					**02114420**
	Actinomyces meyeri	99.59	0.42	GLU 93		
	Lactobacillus fermentum	0.41	0.12	NAG 12		
02114500	**Very good identification**					**02114500**
	Actinomyces meyeri	99.93	0.68	GLU 93		
	Lactobacillus fermentum	0.07	0.24			
02114502	**Good identification**					**02114502**
	Actinomyces meyeri	97.84	0.42	GLU 93		
	Lactobacillus fermentum	2.16	0.24	BGL 3		
02114520	**Good identification**					**02114520**
	Actinomyces meyeri	99.83	0.53	GLU 93		
	Lactobacillus fermentum	0.17	0.16	NAG 12		
02115000	**Very good identification**					**02115000**
					GRA	
	Actinomyces meyeri	97.15	0.71	GLU 93	99	
	Lactobacillus fermentum	2.69	0.55		99	
	Prevotella intermedia	0.10	0.17		1	
02115001	**Good identification**					**02115001**
	Actinomyces meyeri	97.31	0.36	GLU 93		
	Lactobacillus fermentum	2.69	0.20	URE 1		
02115002	**Low discrimination**					**02115002**
	Actinomyces meyeri	52.64	0.45	GLU 93		
	Lactobacillus fermentum	47.06	0.55	BGL 3		
	Actinomyces odontolyticus	0.30	0.10			
02115010	**Good identification**					**02115010**
	Actinomyces meyeri	97.31	0.36	GLU 93		
	Lactobacillus fermentum	2.69	0.20	MLZ 1		
02115020	**Acceptable identification**					**02115020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		93.67	0.56	GLU 93		
Lactobacillus fermentum		6.33	0.46	NAG 12		
02115022	**Low discrimination**					**02115022**
Lactobacillus fermentum		68.61	0.46	GLU 99		
Actinomyces meyeri		31.39	0.30			
02115040	**Low discrimination**					**02115040**
					SPO	
Actinomyces meyeri		87.47	0.36	GLU 93	1	
Clostridium glycolicum		10.11	0.15	SOR 1	99	
Lactobacillus fermentum		2.42	0.20		1	
02115100	**Very good identification**					**02115100**
					GRA	
Actinomyces meyeri		98.88	0.81	GLU 93	99	
Lactobacillus fermentum		1.12	0.58		99	
Fusobacterium mortiferum		0.00	0.13		1	
02115101	**Good identification**					**02115101**
					SPO	
Actinomyces meyeri		98.02	0.47	GLU 93	1	
Lactobacillus fermentum		1.11	0.24	URE 1	1	
Clostridium chauvoei		0.87	0.09		99	
02115102	**Low discrimination**					**02115102**
Actinomyces meyeri		73.28	0.55	GLU 93		
Lactobacillus fermentum		26.73	0.58	BGL 3		
02115110	**Good identification**					**02115110**
Actinomyces meyeri		98.89	0.47	GLU 93		
Lactobacillus fermentum		1.12	0.24	MLZ 1		
02115120	**Very good identification**					**02115120**
					GRA	
Actinomyces meyeri		96.93	0.66	GLU 93	99	
Lactobacillus fermentum		2.67	0.50	NAG 12	99	
Prevotella melaninogenica		0.40	0.17		1	
02115122	**Low discrimination**					**02115122**
					GRA	
Actinomyces meyeri		52.16	0.40	GLU 93	99	
Lactobacillus fermentum		46.51	0.50	NAG 12	99	
Prevotella melaninogenica		1.33	0.05	BGL 3	1	
02115140	**Good identification**					**02115140**
Actinomyces meyeri		98.89	0.47	GLU 93		
Lactobacillus fermentum		1.12	0.24	SOR 1		
02115200	**Acceptable identification**					**02115200**
Actinomyces meyeri		92.32	0.36	GLU 93		
Actinomyces odontolyticus		5.13	0.19	RHA 1		
Lactobacillus fermentum		2.55	0.20			
02115300	**Good identification**					**02115300**
Actinomyces meyeri		98.46	0.47	GLU 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		1.11	0.24	RHA 1		
Actinomyces odontolyticus		0.43	0.10			
02115400	**Good identification**					**02115400**
Actinomyces meyeri		90.04	0.60	GLU 93		
Lactobacillus fermentum		9.96	0.55			
02115402	**Low discrimination**					**02115402**
Lactobacillus fermentum		78.15	0.55	GLU 99		
Actinomyces meyeri		21.85	0.34			
02115420	**Low discrimination**					**02115420**
Actinomyces meyeri		78.72	0.45	GLU 93		
Lactobacillus fermentum		21.28	0.46	NAG 12		
02115422	**Low discrimination**					**02115422**
Lactobacillus fermentum		89.74	0.46	GLU 99		
Actinomyces meyeri		10.26	0.20			
02115500	**Very good identification**					**02115500**
Actinomyces meyeri		95.66	0.71	GLU 93		
Lactobacillus fermentum		4.32	0.58			
Bifidobacterium longum subsp. longum		0.02	0.12			
02115501	**Good identification**					**02115501**
Actinomyces meyeri		95.68	0.36	GLU 93		
Lactobacillus fermentum		4.32	0.24	URE 1		
02115502	**Low discrimination**					**02115502**
Lactobacillus fermentum		59.16	0.58	GLU 99		
Actinomyces meyeri		40.55	0.45			
Bifidobacterium longum subsp. longum		0.29	0.12			
02115510	**Acceptable identification**					**02115510**
Actinomyces meyeri		92.41	0.36	GLU 93		
Lactobacillus fermentum		4.17	0.24	MLZ 1		
Bifidobacterium longum subsp. longum		3.42	0.16			
02115520	**Low discrimination**					**02115520**
Actinomyces meyeri		89.55	0.56	GLU 93		
Lactobacillus fermentum		9.88	0.50	NAG 12		
Bifidobacterium longum subsp. longum		0.58	0.22			
02115522	**Low discrimination**					**02115522**
Lactobacillus fermentum		74.69	0.50	GLU 99		
Actinomyces meyeri		20.94	0.30			
Bifidobacterium longum subsp. longum		4.37	0.22			
02115540	**Good identification**					**02115540**
Actinomyces meyeri		95.68	0.36	GLU 93		
Lactobacillus fermentum		4.32	0.24	SOR 1		
02115700	**Good identification**					**02115700**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		95.68	0.36	GLU 93			
Lactobacillus fermentum		4.32	0.24	RHA 1			
02116100	**Good identification**						**02116100**
Actinomyces meyeri		99.48	0.44	GLU 93			
Propionibacterium propionicum		0.52	0.15	MAN 1			
02117000	**Good identification**						**02117000**
Actinomyces meyeri		97.31	0.36	GLU 93			
Lactobacillus fermentum		2.69	0.20	MAN 1			
02117100	**Good identification**						**02117100**
Actinomyces meyeri		98.55	0.47	GLU 93			
Lactobacillus fermentum		1.11	0.24	MAN 1			
Propionibacterium propionicum		0.34	0.15				
02117500	**Good identification**						**02117500**
Actinomyces meyeri		95.68	0.36	GLU 93			
Lactobacillus fermentum		4.32	0.24	MAN 1			
02120000	**Low discrimination**						**02120000**
					COC	SPO	
Clostridium botulinum biovar A		75.84	0.48	GLU 90	1	99	
Clostridium sporogenes		11.79	0.32	SUC 7	1	99	
Peptostreptococcus anaerobius		8.57	0.27		99	1	
02120040	**Low discrimination**						**02120040**
					COC	GRA	
Clostridium botulinum biovar A		86.83	0.39	GLU 90	1	99	
Clostridium botulinum biovar B		11.75	0.21	SUC 7	1	99	
Megasphaera elsdenii		0.64	0.07		99	1	
02130100	**Good identification**						**02130100**
					GRA		
Actinomyces meyeri		96.79	0.39	GLU 93	99		
Fusobacterium mortiferum		2.41	0.21	TRE 1	1		
Propionibacterium propionicum		0.80	0.14		99		
02131100	**Acceptable identification**						**02131100**
					GRA		
Actinomyces meyeri		90.83	0.42	GLU 93	99		
Lactobacillus fermentum		3.39	0.28	TRE 1	99		
Fusobacterium mortiferum		2.26	0.24		1		
02132100	**Good identification**						**02132100**
Propionibacterium propionicum		98.79	0.48	NIT 86			
Actinomyces meyeri		1.21	0.05	GLU 99			
02132102	**Good identification**						**02132102**
Propionibacterium propionicum		100.0	0.51	NIT 86			
02132140	**Good identification**						**02132140**
Propionibacterium propionicum		100.0	0.48	NIT 86			
02132142	**Good identification**						**02132142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.51	NIT 86		
02133100	**Good identification**					**02133100**
Propionibacterium propionicum		98.19	0.48	NIT 86		
Actinomyces meyeri		1.81	0.08	GLU 99		
02133102	**Good identification**					**02133102**
Propionibacterium propionicum		100.0	0.51	NIT 86		
02133140	**Good identification**					**02133140**
Propionibacterium propionicum		100.0	0.48	NIT 86		
02133142	**Good identification**					**02133142**
Propionibacterium propionicum		100.0	0.51	NIT 86		
02134100	**Good identification**					**02134100**
Actinomyces meyeri		100.0	0.44	GLU 93		
02135000	**Good identification**					**02135000**
Actinomyces meyeri		97.31	0.36	GLU 93		
Lactobacillus fermentum		2.69	0.20	TRE 1		
02135100	**Good identification**					**02135100**
					COC	
Actinomyces meyeri		96.73	0.47	GLU 93	1	
Atopobium parvulum		2.18	0.16	TRE 1	99	
Lactobacillus fermentum		1.09	0.24		1	
02135500	**Good identification**					**02135500**
Actinomyces meyeri		95.68	0.36	GLU 93		
Lactobacillus fermentum		4.32	0.24	TRE 1		
02150100	**Good identification**					**02150100**
					GRA	
Actinomyces meyeri		97.30	0.39	GLU 93	99	
Fusobacterium mortiferum		2.70	0.21	CEL 1	1	
02151100	**Acceptable identification**					**02151100**
					GRA	
Actinomyces meyeri		92.84	0.42	GLU 93	99	
Lactobacillus fermentum		3.46	0.28	CEL 1	99	
Fusobacterium mortiferum		2.58	0.24		1	
02154100	**Good identification**					**02154100**
Actinomyces meyeri		100.0	0.44	GLU 93		
02155000	**Good identification**					**02155000**
Actinomyces meyeri		97.31	0.36	GLU 93		
Lactobacillus fermentum		2.69	0.20	CEL 1		
02155100	**Good identification**					**02155100**
Actinomyces meyeri		98.89	0.47	GLU 93		
Lactobacillus fermentum		1.12	0.24	CEL 1		
02155500	**Good identification**					**02155500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		95.68	0.36	GLU 93			
Lactobacillus fermentum		4.32	0.24	CEL 1			
02200002	**Low discrimination**						**02200002**
					COC	GRA	
Atopobium minutum		25.46	0.37	SUC 5	1	99	
Peptococcus niger		24.91	0.19	SAL 1	99	99	
Acidaminococcus fermentans		20.28	0.15		99	1	
02210000	**Low discrimination**						**02210000**
					COC	GRA	
Atopobium minutum		32.08	0.39	SUC 5	1	99	
Fusobacterium mortiferum		27.33	0.44	SAL 1	1	1	
Megasphaera elsdenii		11.95	0.29		99	1	
02210002	**Low discrimination**						**02210002**
					COC	GRA	
Atopobium minutum		81.99	0.36	SUC 5	1	99	
Fusobacterium mortiferum		11.64	0.27	SAL 1	1	1	
Megasphaera elsdenii		5.09	0.12		99	1	
02210100	**Low discrimination**						**02210100**
					GRA		
Fusobacterium mortiferum		59.62	0.50	GLU 99	1		
Actinomyces meyeri		40.38	0.38		99		
02211000	**Low discrimination**						**02211000**
					GRA		
Fusobacterium mortiferum		59.44	0.47	GLU 99	1		
Actinomyces meyeri		23.49	0.31		99		
Atopobium minutum		15.51	0.31		99		
02211100	**Low discrimination**						**02211100**
					GRA		
Fusobacterium mortiferum		59.28	0.53	GLU 99	1		
Actinomyces meyeri		40.16	0.41		99		
Lactobacillus fermentum		0.56	0.20		99		
02211102	**Low discrimination**						**02211102**
					GRA		
Fusobacterium mortiferum		78.49	0.37	GLU 99	1		
Actinomyces meyeri		14.80	0.15	BGL 10	99		
Lactobacillus fermentum		6.71	0.20		99		
02214100	**Acceptable identification**						**02214100**
					GRA		
Actinomyces meyeri		94.59	0.43	GLU 93	99		
Fusobacterium mortiferum		5.41	0.31	SAL 1	1		
02215000	**Acceptable identification**						**02215000**
					GRA		
Actinomyces meyeri		90.21	0.35	GLU 93	99		
Fusobacterium mortiferum		8.85	0.27	SAL 1	1		
Lactobacillus fermentum		0.94	0.12		99		
02215100	**Acceptable identification**						**02215100**
					GRA		
Actinomyces meyeri		94.21	0.46	GLU 93	99		
Fusobacterium mortiferum		5.39	0.34	SAL 1	1		
Lactobacillus fermentum		0.40	0.16		99		
02215500	**Good identification**						**02215500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		98.33	0.35	GLU 93		
Lactobacillus fermentum		1.67	0.16	SAL 1		
02230100	**Good identification**					**02230100**
					GRA	
Fusobacterium mortiferum		97.76	0.41	GLU 99	1	
Actinomyces meyeri		2.24	0.04		99	
02231000	**Good identification**					**02231000**
Fusobacterium mortiferum		100.0	0.38	GLU 99		
02231100	**Good identification**					**02231100**
					GRA	
Fusobacterium mortiferum		97.76	0.44	GLU 99	1	
Actinomyces meyeri		2.24	0.07		99	
02250000	**Good identification**					**02250000**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
02250100	**Good identification**					**02250100**
					GRA	
Fusobacterium mortiferum		97.99	0.42	GLU 99	1	
Actinomyces meyeri		2.01	0.04		99	
02251000	**Low discrimination**					**02251000**
					GRA	
Fusobacterium mortiferum		87.78	0.39	GLU 99	1	
Lactobacillus catenaformis		9.87	0.12		99	
Lactobacillus jensenii		2.35	0.02		99	
02251100	**Acceptable identification**					**02251100**
					GRA	
Fusobacterium mortiferum		91.51	0.45	GLU 99	1	
Lactobacillus catenaformis		6.62	0.15		99	
Actinomyces meyeri		1.88	0.07		99	
02271100	**Good identification**					**02271100**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
02310100	**Low discrimination**					**02310100**
					GRA	
Actinomyces meyeri		83.29	0.39	GLU 93	99	
Fusobacterium mortiferum		13.48	0.34	SAL 1	1	
Actinomyces odontolyticus		3.24	0.18		99	
02311100	**Low discrimination**					**02311100**
					GRA	
Actinomyces meyeri		82.21	0.42	GLU 93	99	
Fusobacterium mortiferum		13.30	0.37	SAL 1	1	
Lactobacillus fermentum		3.07	0.28		99	
02314100	**Good identification**					**02314100**
					GRA	
Actinomyces meyeri		98.41	0.44	GLU 93	99	
Actinomyces odontolyticus		0.97	0.13	SAL 1	99	
Fusobacterium mortiferum		0.62	0.15		1	
02315000	**Acceptable identification**					**02315000**
Actinomyces meyeri		91.41	0.36	GLU 93		
Actinomyces odontolyticus		5.08	0.19	SAL 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		2.53	0.20				
02315100	**Good identification**						**02315100**
					GRA		
Actinomyces meyeri		97.85	0.47	GLU 93	99		
Lactobacillus fermentum		1.10	0.24	SAL 1	99		
Fusobacterium mortiferum		0.61	0.18		1		
02315500	**Good identification**						**02315500**
Actinomyces meyeri		95.68	0.36	GLU 93			
Lactobacillus fermentum		4.32	0.24	SAL 1			
02377100	**Good identification**						**02377100**
					SPO		
Actinomyces israelii		96.41	0.42	GLU 99	1		
Clostridium tertium		3.59	0.03		99		
02377102	**Acceptable identification**						**02377102**
					SPO		
Actinomyces israelii		93.07	0.45	GLU 99	1		
Clostridium tertium		6.93	0.11		99		
02377142	**Good identification**						**02377142**
Actinomyces israelii		100.0	0.38	GLU 99			
02377500	**Good identification**						**02377500**
Actinomyces israelii		100.0	0.40	GLU 99			
02377502	**Acceptable identification**						**02377502**
Actinomyces israelii		93.41	0.43	GLU 99			
Bifidobacterium adolescentis		6.59	0.17				
02377542	**Good identification**						**02377542**
Actinomyces israelii		100.0	0.36	GLU 99			
02400001	**Low discrimination**						**02400001**
					COC	GRA	
Anaerococcus prevotii		84.64	0.50	SUC 1	99	99	
Campylobacter ureolyticus		12.67	0.32		1	1	
Peptococcus niger		1.38	0.15		99	99	
02400002	**Low discrimination**						**02400002**
					COC		
Atopobium minutum		20.69	0.37	SUC 5	1		
Peptococcus niger		20.24	0.19	RAF 1	99		
Anaerococcus prevotii		19.32	0.23		99		
02400003	**Good identification**						**02400003**
					COC	GRA	
Anaerococcus prevotii		98.27	0.39	SUC 1	99	99	
Peptococcus niger		1.13	0.02		99	99	
Campylobacter ureolyticus		0.60	-.03		1	1	
02410000	**Low discrimination**						**02410000**
					COC	GRA	
Atopobium minutum		29.23	0.39	SUC 5	1	99	
Fusobacterium mortiferum		24.90	0.44	RAF 1	1	1	
Megasphaera elsdenii		10.89	0.29		99	1	
02410002	**Low discrimination**						**02410002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Atopobium minutum		77.21	0.36	SUC 5	1	99	
Fusobacterium mortiferum		10.96	0.27	RAF 1	1	1	
Megasphaera elsdenii		4.79	0.12		99	1	
02410100	**Low discrimination**						**02410100**
					GRA		
Fusobacterium mortiferum		58.80	0.50	GLU 99	1		
Actinomyces meyeri		39.83	0.38		99		
Lactobacillus fermentum		1.37	0.24		99		
02411000	**Low discrimination**						**02411000**
					GRA		
Lactobacillus fermentum		54.24	0.54	GLU 99	99		
Fusobacterium mortiferum		24.73	0.47		1		
Actinomyces meyeri		9.77	0.31		99		
02411002	**Low discrimination**						**02411002**
					GRA		
Lactobacillus fermentum		87.46	0.54	GLU 99	99		
Atopobium minutum		6.94	0.28		99		
Fusobacterium mortiferum		4.43	0.30		1		
02411020	**Low discrimination**						**02411020**
Lactobacillus fermentum		79.74	0.46	GLU 99			
Atopobium minutum		9.48	0.23				
Actinomyces meyeri		5.88	0.16				
02411022	**Acceptable identification**						**02411022**
					GRA		
Lactobacillus fermentum		90.03	0.46	GLU 99	99		
Atopobium minutum		7.14	0.20		99		
Pseudoflavonifractor capillosus		2.83	-.01		1		
02411100	**Low discrimination**						**02411100**
					GRA		
Lactobacillus fermentum		47.65	0.58	GLU 99	99		
Fusobacterium mortiferum		31.07	0.53		1		
Actinomyces meyeri		21.05	0.41		99		
02411102	**Good identification**						**02411102**
					GRA		
Lactobacillus fermentum		92.07	0.58	GLU 99	99		
Fusobacterium mortiferum		6.67	0.37		1		
Actinomyces meyeri		1.26	0.15		99		
02411122	**Low discrimination**						**02411122**
					GRA		
Lactobacillus fermentum		78.26	0.50	GLU 99	99		
Prevotella melaninogenica		21.74	0.21		1		
02411400	**Good identification**						**02411400**
					GRA		
Lactobacillus fermentum		95.27	0.54	GLU 99	99		
Actinomyces meyeri		4.29	0.20		99		
Fusobacterium mortiferum		0.44	0.13		1		
02411402	**Good identification**						**02411402**
Lactobacillus fermentum		100.0	0.54	GLU 99			
02411420	**Good identification**						**02411420**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		98.19	0.46	GLU 99		
Actinomyces meyeri		1.81	0.05			
02411422	**Good identification**					**02411422**
Lactobacillus fermentum		100.0	0.46	GLU 99		
02411500	**Low discrimination**					**02411500**
					GRA	
Lactobacillus fermentum		89.52	0.58	GLU 99	99	
Actinomyces meyeri		9.89	0.31		99	
Fusobacterium mortiferum		0.59	0.19		1	
02411502	**Very good identification**					**02411502**
Lactobacillus fermentum		99.66	0.58	GLU 99		
Actinomyces meyeri		0.34	0.05			
02411520	**Acceptable identification**					**02411520**
					GRA	
Lactobacillus fermentum		93.74	0.50	GLU 99	99	
Actinomyces meyeri		4.24	0.16		99	
Prevotella melaninogenica		1.38	-.01		1	
02411522	**Good identification**					**02411522**
Lactobacillus fermentum		99.32	0.50	GLU 99		
Bifidobacterium longum subsp. longum		0.68	0.05			
02414100	**Acceptable identification**					**02414100**
					GRA	
Actinomyces meyeri		93.67	0.43	GLU 93	99	
Fusobacterium mortiferum		5.36	0.31	RAF 1	1	
Lactobacillus fermentum		0.97	0.20		99	
02415000	**Low discrimination**					**02415000**
					GRA	
Lactobacillus fermentum		60.46	0.50	GLU 99	99	
Actinomyces meyeri		36.01	0.35		99	
Fusobacterium mortiferum		3.53	0.27		1	
02415002	**Good identification**					**02415002**
					GRA	
Lactobacillus fermentum		97.57	0.50	GLU 99	99	
Actinomyces meyeri		1.80	0.10		99	
Fusobacterium mortiferum		0.63	0.11		1	
02415020	**Low discrimination**					**02415020**
Lactobacillus fermentum		80.41	0.42	GLU 99		
Actinomyces meyeri		19.59	0.21			
02415022	**Good identification**					**02415022**
Lactobacillus fermentum		100.0	0.42	GLU 99		
02415100	**Low discrimination**					**02415100**
					GRA	
Actinomyces meyeri		57.40	0.46	GLU 93	99	
Lactobacillus fermentum		39.31	0.54	RAF 1	99	
Fusobacterium mortiferum		3.29	0.34		1	
02415102	**Acceptable identification**					**02415102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Lactobacillus fermentum		94.84	0.54	GLU 99	99		
Actinomyces meyeri		4.28	0.20		99		
Fusobacterium mortiferum		0.88	0.17		1		
02415120	**Low discrimination**						**02415120**
					GRA		
Lactobacillus fermentum		61.64	0.46	GLU 99	99		
Actinomyces meyeri		36.82	0.31		99		
Prevotella melaninogenica		1.55	-.01		1		
02415122	**Good identification**						**02415122**
Lactobacillus fermentum		98.19	0.46	GLU 99			
Actinomyces meyeri		1.81	0.05				
02415400	**Low discrimination**						**02415400**
Lactobacillus fermentum		87.04	0.50	GLU 99			
Actinomyces meyeri		12.96	0.25				
02415402	**Good identification**						**02415402**
Lactobacillus fermentum		100.0	0.50	GLU 99			
02415420	**Acceptable identification**						**02415420**
Lactobacillus fermentum		94.26	0.42	GLU 99			
Actinomyces meyeri		5.74	0.10				
02415422	**Good identification**						**02415422**
Lactobacillus fermentum		100.0	0.42	GLU 99			
02415500	**Low discrimination**						**02415500**
Lactobacillus fermentum		73.26	0.54	GLU 99			
Actinomyces meyeri		26.74	0.35				
02415502	**Good identification**						**02415502**
Lactobacillus fermentum		98.88	0.54	GLU 99			
Actinomyces meyeri		1.12	0.10				
02415520	**Low discrimination**						**02415520**
Lactobacillus fermentum		84.43	0.46	GLU 99			
Actinomyces meyeri		12.61	0.21				
Bifidobacterium longum subsp. longum		2.96	0.14				
02415522	**Good identification**						**02415522**
Lactobacillus fermentum		96.61	0.46	GLU 99			
Bifidobacterium longum subsp. longum		3.39	0.14				
02430100	**Good identification**						**02430100**
					GRA		
Fusobacterium mortiferum		97.76	0.41	GLU 99	1		
Actinomyces meyeri		2.24	0.04		99		
02431000	**Low discrimination**						**02431000**
					GRA	CAT	
Fusobacterium mortiferum		56.61	0.38	GLU 99	1	5	
Propionibacterium granulosum		39.19	0.32		99	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		4.20	0.20		99	1	
02431100	**Acceptable identification**						**02431100**
					GRA		
Fusobacterium mortiferum		93.04	0.44	GLU 99	1		
Lactobacillus fermentum		4.83	0.24		99		
Actinomyces meyeri		2.13	0.07		99		
02450000	**Good identification**						**02450000**
Fusobacterium mortiferum		100.0	0.36	GLU 99			
02450100	**Good identification**						**02450100**
					GRA		
Fusobacterium mortiferum		97.99	0.42	GLU 99	1		
Actinomyces meyeri		2.01	0.04		99		
02451000	**Acceptable identification**						**02451000**
					GRA		
Fusobacterium mortiferum		93.77	0.39	GLU 99	1		
Lactobacillus fermentum		6.23	0.20		99		
02451100	**Acceptable identification**						**02451100**
					GRA		
Fusobacterium mortiferum		93.72	0.45	GLU 99	1		
Lactobacillus fermentum		4.36	0.24		99		
Actinomyces meyeri		1.92	0.07		99		
02471100	**Good identification**						**02471100**
Fusobacterium mortiferum		100.0	0.36	GLU 99			
02501120	**Good identification**						**02501120**
					COC	GRA	
Prevotella melaninogenica		98.96	0.51	GLU 90	1	1	
Blautia hansenii		0.79	0.15	MNS 90	99	99	
Lactobacillus fermentum		0.19	0.24	FRU 90	1	99	
02501122	**Good identification**						**02501122**
					COC	GRA	
Prevotella melaninogenica		98.57	0.39	GLU 90	1	1	
Lactobacillus fermentum		0.98	0.24	MNS 90	1	99	
Blautia hansenii		0.46	-.01	FRU 90	99	99	
02505530	**Good identification**						**02505530**
Bifidobacterium longum subsp. longum		100.0	0.35	GLU 99			
02505532	**Good identification**						**02505532**
Bifidobacterium longum subsp. longum		100.0	0.35	GLU 99			
02510000	**Acceptable identification**						**02510000**
					COC	GRA	
Prevotella intermedia		92.95	0.54	GLU 90	1	1	
Megasphaera elsdenii		3.28	0.32		99	1	
Actinomyces meyeri		2.23	0.28		1	99	
02510100	**Low discrimination**						**02510100**
					GRA		
Actinomyces meyeri		72.46	0.39	GLU 93	99		
Fusobacterium mortiferum		11.73	0.34	RAF 1	1		
Prevotella intermedia		7.64	0.20		1		
02511000	**Low discrimination**						**02511000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Prevotella intermedia		63.28	0.54	GLU 90	1		
Lactobacillus fermentum		33.57	0.62		99		
Actinomyces meyeri		2.27	0.31		99		
02511002	**Very good identification**						**02511002**
					GRA		
Lactobacillus fermentum		97.58	0.62	GLU 99	99		
Prevotella intermedia		1.86	0.20		1		
Fusobacterium mortiferum		0.20	0.15		1		
02511020	**Low discrimination**						**02511020**
					GRA		
Lactobacillus fermentum		53.25	0.54	GLU 99	99		
Prevotella melaninogenica		42.24	0.33		1		
Prevotella intermedia		3.04	0.20		1		
02511022	**Low discrimination**						**02511022**
					GRA		
Lactobacillus fermentum		86.87	0.54	GLU 99	99		
Prevotella melaninogenica		13.13	0.21		1		
02511100	**Low discrimination**						**02511100**
					GRA		
Lactobacillus fermentum		72.99	0.66	GLU 99	99		
Actinomyces meyeri		12.12	0.42		99		
Prevotella melaninogenica		11.83	0.33		1		
02511102	**Very good identification**						**02511102**
					GRA		
Lactobacillus fermentum		95.86	0.66	GLU 99	99		
Prevotella melaninogenica		2.96	0.21		1		
Actinomyces meyeri		0.49	0.16		99		
02511120	**Very good identification**						**02511120**
					GRA		
Prevotella melaninogenica		97.79	0.67	GLU 90	1		
Lactobacillus fermentum		2.03	0.58	MNS 90	99		
Actinomyces meyeri		0.14	0.27		99		
02511122	**Low discrimination**						**02511122**
					GRA	CAT	
Prevotella melaninogenica		89.83	0.55	GLU 90	1	1	
Lactobacillus fermentum		9.80	0.58	MNS 90	99	1	
Capnocytophaga ochracea		0.17	0.12		1	90	
02511400	**Very good identification**						**02511400**
					GRA		
Lactobacillus fermentum		96.53	0.62	GLU 99	99		
Prevotella intermedia		1.84	0.20		1		
Actinomyces meyeri		1.63	0.21		99		
02511402	**Very good identification**						**02511402**
Lactobacillus fermentum		100.0	0.62	GLU 99			
02511420	**Good identification**						**02511420**
					GRA		
Lactobacillus fermentum		98.17	0.54	GLU 99	99		
Prevotella melaninogenica		0.79	-.01		1		
Actinomyces meyeri		0.68	0.06		99		
02511422	**Good identification**						**02511422**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		99.63	0.54	GLU 99		
Bifidobacterium longum subsp. longum		0.37	0.05			
02511500	**Good identification**					**02511500**
Lactobacillus fermentum		94.16	0.66	GLU 99		
Actinomyces meyeri		3.91	0.31			
Bifidobacterium longum subsp. longum		1.78	0.29			
02511502	**Very good identification**					**02511502**
Lactobacillus fermentum		97.95	0.66	GLU 99		
Bifidobacterium longum subsp. longum		1.85	0.29			
Actinomyces meyeri		0.13	0.06			
02511520	**Low discrimination**					**02511520**
					GRA	
Lactobacillus fermentum		57.81	0.58	GLU 99	99	
Prevotella melaninogenica		28.10	0.33		1	
Bifidobacterium longum subsp. longum		13.11	0.40		99	
02511522	**Low discrimination**					**02511522**
					GRA	
Lactobacillus fermentum		75.62	0.58	GLU 99	99	
Bifidobacterium longum subsp. longum		17.15	0.40		99	
Prevotella melaninogenica		7.00	0.21		1	
02511530	**Good identification**					**02511530**
					GRA	
Bifidobacterium longum subsp. longum		96.26	0.44	GLU 99	99	
Lactobacillus fermentum		2.52	0.24	MNS 87	99	
Prevotella melaninogenica		1.22	-.01		1	
02511532	**Good identification**					**02511532**
Bifidobacterium longum subsp. longum		97.45	0.44	GLU 99		
Lactobacillus fermentum		2.55	0.24	MNS 87		
02512100	**Good identification**					**02512100**
Propionibacterium propionicum		98.51	0.46	NIT 86		
Actinomyces meyeri		1.49	0.05	GLU 99		
02512102	**Good identification**					**02512102**
Propionibacterium propionicum		100.0	0.49	NIT 86		
02512140	**Good identification**					**02512140**
Propionibacterium propionicum		100.0	0.46	NIT 86		
02512142	**Good identification**					**02512142**
Propionibacterium propionicum		100.0	0.49	NIT 86		
02513100	**Low discrimination**					**02513100**
Propionibacterium propionicum		82.25	0.46	NIT 86		
Lactobacillus fermentum		11.21	0.32	GLU 99		
Actinomyces naeslundii		2.86	0.32			
02513102	**Low discrimination**					**02513102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		89.76	0.49	NIT 86			
Lactobacillus fermentum		8.16	0.32	GLU 99			
Actinomyces naeslundii		2.08	0.32				
02513140	**Good identification**						**02513140**
Propionibacterium propionicum		100.0	0.46	NIT 86			
02513142	**Good identification**						**02513142**
Propionibacterium propionicum		100.0	0.49	NIT 86			
02514100	**Good identification**						**02514100**
					GRA		
Actinomyces meyeri		96.72	0.44	GLU 93	99		
Lactobacillus fermentum		2.67	0.28	RAF 1	99		
Fusobacterium mortiferum		0.61	0.15		1		
02515000	**Low discrimination**						**02515000**
					GRA		
Lactobacillus fermentum		79.45	0.58	GLU 99	99		
Actinomyces meyeri		17.79	0.36		99		
Prevotella intermedia		2.58	0.20		1		
02515002	**Very good identification**						**02515002**
Lactobacillus fermentum		99.31	0.58	GLU 99			
Actinomyces meyeri		0.69	0.11				
02515020	**Acceptable identification**						**02515020**
					GRA		
Lactobacillus fermentum		90.48	0.50	GLU 99	99		
Actinomyces meyeri		8.29	0.22		99		
Prevotella melaninogenica		1.23	-.01		1		
02515022	**Good identification**						**02515022**
Lactobacillus fermentum		100.0	0.50	GLU 99			
02515100	**Low discrimination**						**02515100**
					GRA		
Lactobacillus fermentum		64.31	0.62	GLU 99	99		
Actinomyces meyeri		35.29	0.47		99		
Fusobacterium mortiferum		0.22	0.18		1		
02515102	**Very good identification**						**02515102**
Lactobacillus fermentum		98.20	0.62	GLU 99			
Actinomyces meyeri		1.67	0.21				
Bifidobacterium adolescentis		0.13	0.07				
02515120	**Low discrimination**						**02515120**
					GRA		
Lactobacillus fermentum		48.45	0.54	GLU 99	99		
Prevotella melaninogenica		40.11	0.33		1		
Actinomyces meyeri		10.88	0.32		99		
02515122	**Low discrimination**						**02515122**
					CAT	GRA	
Lactobacillus fermentum		83.16	0.54	GLU 99	1	99	
Prevotella melaninogenica		13.11	0.21		1	1	
Bacteroides fragilis		1.19	-.03		99	1	
02515400	**Good identification**						**02515400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		94.70	0.58	GLU 99		
Actinomyces meyeri		5.30	0.26			
02515402	**Very good identification**					**02515402**
Lactobacillus fermentum		100.0	0.58	GLU 99		
02515420	**Good identification**					**02515420**
Lactobacillus fermentum		95.97	0.50	GLU 99		
Actinomyces meyeri		2.20	0.11			
Bifidobacterium longum subsp. longum		1.83	0.14			
02515422	**Good identification**					**02515422**
Lactobacillus fermentum		98.13	0.50	GLU 99		
Bifidobacterium longum subsp. longum		1.87	0.14			
02515500	**Low discrimination**					**02515500**
Lactobacillus fermentum		81.05	0.62	GLU 99		
Actinomyces meyeri		11.12	0.36			
Bifidobacterium longum subsp. longum		7.82	0.37			
02515502	**Good identification**					**02515502**
Lactobacillus fermentum		90.11	0.62	GLU 99		
Bifidobacterium longum subsp. longum		8.70	0.37			
Bifidobacterium adolescentis		0.81	0.21			
02515510	**Acceptable identification**					**02515510**
Bifidobacterium longum subsp. longum		93.47	0.41	GLU 99		
Lactobacillus fermentum		5.74	0.28	MNS 87		
Actinomyces meyeri		0.79	0.02			
02515512	**Low discrimination**					**02515512**
Bifidobacterium longum subsp. longum		88.63	0.41	GLU 99		
Bifidobacterium adolescentis		5.92	0.23	MNS 87		
Lactobacillus fermentum		5.45	0.28			
02515520	**Low discrimination**					**02515520**
Bifidobacterium longum subsp. longum		51.12	0.48	GLU 99		
Lactobacillus fermentum		44.13	0.54	MNS 87		
Actinomyces meyeri		2.48	0.22			
02515522	**Low discrimination**					**02515522**
Bifidobacterium longum subsp. longum		53.67	0.48	GLU 99		
Lactobacillus fermentum		46.33	0.54	MNS 87		
02515530	**Good identification**					**02515530**
Bifidobacterium longum subsp. longum		99.49	0.52	GLU 99		
Lactobacillus fermentum		0.51	0.20	MNS 87		
02515532	**Good identification**					**02515532**
Bifidobacterium longum subsp. longum		99.49	0.52	GLU 99		
Lactobacillus fermentum		0.51	0.20	MNS 87		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
02532100	**Good identification**					**02532100**
Propionibacterium propionicum		100.0	0.45	NIT 86		
02532102	**Good identification**					**02532102**
Propionibacterium propionicum		100.0	0.48	NIT 86		
02532140	**Good identification**					**02532140**
Propionibacterium propionicum		100.0	0.45	NIT 86		
02532142	**Good identification**					**02532142**
Propionibacterium propionicum		100.0	0.48	NIT 86		
02533100	**Acceptable identification**					**02533100**
Propionibacterium propionicum		92.38	0.45	NIT 86		
Actinomyces naeslundii		7.62	0.37	GLU 99		
02533101	**Low discrimination**					**02533101**
Actinomyces naeslundii		52.69	0.40	GLU 99		
Propionibacterium propionicum		47.31	0.29	MNS 88		
02533102	**Acceptable identification**					**02533102**
Propionibacterium propionicum		94.79	0.48	NIT 86		
Actinomyces naeslundii		5.21	0.37	GLU 99		
02533120	**Low discrimination**					**02533120**
					GRA	
Actinomyces naeslundii		75.34	0.37	GLU 99	99	
Prevotella melaninogenica		24.66	-.01	MNS 88	1	
02533121	**Good identification**					**02533121**
Actinomyces naeslundii		100.0	0.40	GLU 99		
02533122	**Low discrimination**					**02533122**
Actinomyces naeslundii		84.48	0.37	GLU 99		
Propionibacterium propionicum		15.52	0.14	MNS 88		
02533123	**Good identification**					**02533123**
Actinomyces naeslundii		100.0	0.40	GLU 99		
02533140	**Good identification**					**02533140**
Propionibacterium propionicum		98.57	0.45	NIT 86		
Actinomyces naeslundii		1.44	0.24	GLU 99		
02533142	**Good identification**					**02533142**
Propionibacterium propionicum		99.04	0.48	NIT 86		
Actinomyces naeslundii		0.96	0.24	GLU 99		
02535520	**Good identification**					**02535520**
Bifidobacterium longum subsp. longum		96.63	0.37	GLU 99		
Lactobacillus fermentum		3.37	0.20	MNS 87		
02535522	**Acceptable identification**					**02535522**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium longum subsp. longum		92.77	0.37	GLU 99	99	
Parabacteroides distasonis		4.00	0.18	MNS 87	1	
Lactobacillus fermentum		3.24	0.20		99	
02535530	**Good identification**					**02535530**
Bifidobacterium longum subsp. longum		100.0	0.41	GLU 99		
02535532	**Good identification**					**02535532**
					GRA	
Bifidobacterium longum subsp. longum		98.66	0.41	GLU 99	99	
Parabacteroides distasonis		1.35	0.14	MNS 87	1	
02551120	**Good identification**					**02551120**
					GRA	
Prevotella melaninogenica		99.71	0.51	GLU 90	1	
Lactobacillus fermentum		0.19	0.24	MNS 90	99	
Actinomyces naeslundii		0.10	0.29	CEL 10	99	
02551122	**Good identification**					**02551122**
					GRA	
Prevotella melaninogenica		95.82	0.39	GLU 90	1	
Prevotella oralis		2.32	0.15	MNS 90	1	
Lactobacillus fermentum		0.95	0.24	CEL 10	99	
02553101	**Good identification**					**02553101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02553103	**Good identification**					**02553103**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02553121	**Good identification**					**02553121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02553123	**Good identification**					**02553123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02571101	**Good identification**					**02571101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02571103	**Good identification**					**02571103**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02571121	**Good identification**					**02571121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02571123	**Good identification**					**02571123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
02573100	**Low discrimination**					**02573100**
Actinomyces naeslundii		87.01	0.39	GLU 99		
Actinomyces israelii		12.99	0.15	MNS 88		
02573101	**Good identification**					**02573101**
Actinomyces naeslundii		100.0	0.42	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
02573102	**Low discrimination**					**02573102**
Actinomyces naeslundii		73.40	0.39	GLU 99		
Actinomyces israelii		16.43	0.18	MNS 88		
Propionibacterium propionicum		10.17	0.14			
02573103	**Good identification**					**02573103**
Actinomyces naeslundii		100.0	0.42	GLU 99		
02573120	**Good identification**					**02573120**
Actinomyces naeslundii		100.0	0.39	GLU 99		
02573121	**Good identification**					**02573121**
Actinomyces naeslundii		100.0	0.42	GLU 99		
02573122	**Good identification**					**02573122**
Actinomyces naeslundii		100.0	0.39	GLU 99		
02573123	**Good identification**					**02573123**
Actinomyces naeslundii		100.0	0.42	GLU 99		
02577102	**Acceptable identification**					**02577102**
Actinomyces israelii		94.92	0.36	GLU 99		
Bifidobacterium adolescentis		5.08	0.08	SAL 92		
02610000	**Good identification**					**02610000**
Fusobacterium mortiferum		100.0	0.49	GLU 99		
02610100	**Good identification**					**02610100**
					GRA	
Fusobacterium mortiferum		99.65	0.55	GLU 99	1	
Actinomyces meyeri		0.35	0.04		99	
02610102	**Good identification**					**02610102**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
02611000	**Good identification**					**02611000**
					GRA	
Fusobacterium mortiferum		98.87	0.52	GLU 99	1	
Lactobacillus fermentum		1.13	0.20		99	
02611002	**Acceptable identification**					**02611002**
					GRA	
Fusobacterium mortiferum		90.69	0.35	GLU 99	1	
Lactobacillus fermentum		9.31	0.20	BGL 10	99	
02611100	**Very good identification**					**02611100**
					GRA	
Fusobacterium mortiferum		98.86	0.58	GLU 99	1	
Lactobacillus fermentum		0.79	0.24		99	
Actinomyces meyeri		0.35	0.07		99	
02611102	**Acceptable identification**					**02611102**
					GRA	
Fusobacterium mortiferum		93.30	0.42	GLU 99	1	
Lactobacillus fermentum		6.70	0.24	BGL 10	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
02614100	**Acceptable identification**					**02614100**
					GRA	
Fusobacterium mortiferum		91.67	0.36	GLU 99	1	
Actinomyces meyeri		8.33	0.09	XYL 7	99	
02615100	**Low discrimination**					**02615100**
					GRA	
Fusobacterium mortiferum		86.72	0.39	GLU 99	1	
Actinomyces meyeri		7.88	0.12	XYL 7	99	
Lactobacillus fermentum		5.40	0.20		99	
02630000	**Good identification**					**02630000**
Fusobacterium mortiferum		100.0	0.40	GLU 99		
02630100	**Good identification**					**02630100**
Fusobacterium mortiferum		100.0	0.46	GLU 99		
02631000	**Good identification**					**02631000**
Fusobacterium mortiferum		100.0	0.43	GLU 99		
02631100	**Good identification**					**02631100**
Fusobacterium mortiferum		100.0	0.49	GLU 99		
02650000	**Good identification**					**02650000**
Fusobacterium mortiferum		100.0	0.40	GLU 99		
02650100	**Good identification**					**02650100**
Fusobacterium mortiferum		100.0	0.47	GLU 99		
02651000	**Good identification**					**02651000**
Fusobacterium mortiferum		100.0	0.43	GLU 99		
02651100	**Good identification**					**02651100**
Fusobacterium mortiferum		100.0	0.50	GLU 99		
02670100	**Good identification**					**02670100**
Fusobacterium mortiferum		100.0	0.38	GLU 99		
02671100	**Good identification**					**02671100**
					GRA	
Fusobacterium mortiferum		98.19	0.41	GLU 99	1	
Actinomyces naeslundii		1.81	0.23		99	
02710100	**Good identification**					**02710100**
					GRA	
Fusobacterium mortiferum		96.89	0.39	GLU 99	1	
Actinomyces meyeri		3.12	0.05	MLT 11	99	
02711000	**Low discrimination**					**02711000**
					GRA	
Fusobacterium mortiferum		53.39	0.36	GLU 99	1	
Prevotella intermedia		27.88	0.20	MLT 11	1	
Lactobacillus fermentum		14.79	0.28		99	
02711100	**Low discrimination**					**02711100**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		74.21	0.42	GLU 99	1	
Lactobacillus fermentum		14.37	0.32	MLT 11	99	
Actinomyces naeslundii		6.70	0.36		99	
02711101	**Good identification**					**02711101**
Actinomyces naeslundii		100.0	0.39	GLU 99		
02711103	**Good identification**					**02711103**
Actinomyces naeslundii		100.0	0.39	GLU 99		
02711121	**Low discrimination**					**02711121**
					GRA	
Actinomyces naeslundii		81.17	0.39	GLU 99	99	
Prevotella melaninogenica		18.83	-.01	MNS 88	1	
02711123	**Good identification**					**02711123**
Actinomyces naeslundii		100.0	0.39	GLU 99		
02713001	**Good identification**					**02713001**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02713003	**Good identification**					**02713003**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02713021	**Good identification**					**02713021**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02713023	**Good identification**					**02713023**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02713100	**Acceptable identification**					**02713100**
Actinomyces naeslundii		92.43	0.41	GLU 99		
Propionibacterium propionicum		7.57	0.12	MNS 88		
02713101	**Good identification**					**02713101**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02713102	**Low discrimination**					**02713102**
Actinomyces naeslundii		89.06	0.41	GLU 99		
Propionibacterium propionicum		10.95	0.15	MNS 88		
02713103	**Good identification**					**02713103**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02713120	**Low discrimination**					**02713120**
					GRA	
Actinomyces naeslundii		84.80	0.41	GLU 99	99	
Prevotella melaninogenica		15.20	-.01	MNS 88	1	
02713121	**Good identification**					**02713121**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02713122	**Good identification**					**02713122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	GLU 99		
02713123	**Good identification**					**02713123**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02715512	**Good identification**					**02715512**
Bifidobacterium adolescentis		97.40	0.36	GLU 99		
Bifidobacterium longum subsp. longum		2.60	0.07	MNS 87		
02731001	**Good identification**					**02731001**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02731003	**Good identification**					**02731003**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02731021	**Good identification**					**02731021**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02731023	**Good identification**					**02731023**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02731101	**Good identification**					**02731101**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02731102	**Low discrimination**					**02731102**
					GRA	
Actinomyces naeslundii		84.07	0.41	GLU 99	99	
Fusobacterium mortiferum		15.93	0.17	MNS 88	1	
02731103	**Good identification**					**02731103**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02731120	**Low discrimination**					**02731120**
					GRA	
Actinomyces naeslundii		68.84	0.41	GLU 99	99	
Clostridium perfringens		18.83	0.09	MNS 88	99	
Prevotella melaninogenica		12.34	-.01		1	
02731121	**Good identification**					**02731121**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02731122	**Low discrimination**					**02731122**
Actinomyces naeslundii		88.59	0.41	GLU 99		
Clostridium perfringens		11.41	0.03	MNS 88		
02731123	**Good identification**					**02731123**
Actinomyces naeslundii		100.0	0.44	GLU 99		
02733000	**Good identification**					**02733000**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02733001	**Good identification**					**02733001**
Actinomyces naeslundii		100.0	0.41	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
02733002	**Good identification**					**02733002**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02733003	**Good identification**					**02733003**
Actinomyces naeslundii		100.0	0.41	GLU 99		
02733020	**Good identification**					**02733020**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02733021	**Good identification**					**02733021**
Actinomyces naeslundii		100.0	0.41	GLU 99		
02733022	**Good identification**					**02733022**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02733023	**Good identification**					**02733023**
Actinomyces naeslundii		100.0	0.41	GLU 99		
02733100	**Good identification**					**02733100**
Actinomyces naeslundii		95.40	0.46	GLU 99		
Actinomyces israelii		4.61	0.14	MNS 88		
02733101	**Good identification**					**02733101**
Actinomyces naeslundii		100.0	0.49	GLU 99		
02733102	**Low discrimination**					**02733102**
Actinomyces naeslundii		88.95	0.46	GLU 99		
Actinomyces israelii		6.44	0.17	MNS 88		
Propionibacterium propionicum		4.61	0.14			
02733103	**Good identification**					**02733103**
Actinomyces naeslundii		100.0	0.49	GLU 99		
02733120	**Good identification**					**02733120**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02733121	**Good identification**					**02733121**
Actinomyces naeslundii		100.0	0.49	GLU 99		
02733122	**Good identification**					**02733122**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02733123	**Good identification**					**02733123**
Actinomyces naeslundii		100.0	0.49	GLU 99		
02733141	**Good identification**					**02733141**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02733143	**Good identification**					**02733143**
Actinomyces naeslundii		100.0	0.36	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
02733161	**Good identification**					**02733161**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02733163	**Good identification**					**02733163**
Actinomyces naeslundii		100.0	0.36	GLU 99		
02737102	**Low discrimination**					**02737102**
Actinomyces israelii		85.84	0.35	GLU 99		
Bifidobacterium adolescentis		14.16	0.16	CEL 93		
02751101	**Good identification**					**02751101**
Actinomyces naeslundii		100.0	0.41	GLU 99		
02751102	**Low discrimination**					**02751102**
					GRA	
Actinomyces naeslundii		41.20	0.38	GLU 99	99	
Lactobacillus acidophilus		26.30	0.10	MNS 88	99	
Fusobacterium mortiferum		12.76	0.18		1	
02751103	**Good identification**					**02751103**
Actinomyces naeslundii		100.0	0.41	GLU 99		
02751121	**Good identification**					**02751121**
Actinomyces naeslundii		100.0	0.41	GLU 99		
02751123	**Good identification**					**02751123**
Actinomyces naeslundii		100.0	0.41	GLU 99		
02753000	**Low discrimination**					**02753000**
Actinomyces naeslundii		84.91	0.35	GLU 99		
Actinomyces israelii		15.09	0.13	MNS 88		
02753001	**Good identification**					**02753001**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02753002	**Low discrimination**					**02753002**
Actinomyces naeslundii		78.96	0.35	GLU 99		
Actinomyces israelii		21.05	0.16	MNS 88		
02753003	**Good identification**					**02753003**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02753020	**Good identification**					**02753020**
Actinomyces naeslundii		100.0	0.35	GLU 99		
02753021	**Good identification**					**02753021**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02753022	**Good identification**					**02753022**
Actinomyces naeslundii		100.0	0.35	GLU 99		
02753023	**Good identification**					**02753023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	GLU 99		
02753100	**Low discrimination**					**02753100**
Actinomyces naeslundii		80.02	0.43	GLU 99		
Actinomyces israelii		19.98	0.24	MNS 88		
02753101	**Good identification**					**02753101**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02753102	**Low discrimination**					**02753102**
Actinomyces naeslundii		68.01	0.43	GLU 99		
Actinomyces israelii		25.48	0.27	MNS 88		
Bifidobacterium adolescentis		6.52	0.10			
02753103	**Good identification**					**02753103**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02753120	**Good identification**					**02753120**
Actinomyces naeslundii		100.0	0.43	GLU 99		
02753121	**Good identification**					**02753121**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02753122	**Low discrimination**					**02753122**
					SPO	
Actinomyces naeslundii		85.63	0.43	GLU 99	1	
Clostridium ramosum		14.37	0.05	MNS 88	92	
02753123	**Good identification**					**02753123**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02755502	**Low discrimination**					**02755502**
Bifidobacterium adolescentis		89.40	0.40	GLU 99		
Actinomyces israelii		10.60	0.30	MNS 87		
02755512	**Good identification**					**02755512**
Bifidobacterium adolescentis		95.65	0.41	GLU 99		
Clostridium clostridioforme		2.96	0.12	MNS 87		
Actinomyces israelii		1.39	0.16			
02757100	**Good identification**					**02757100**
Actinomyces israelii		100.0	0.42	GLU 99		
02757102	**Low discrimination**					**02757102**
Actinomyces israelii		87.04	0.45	GLU 99		
Bifidobacterium adolescentis		12.96	0.24			
02757142	**Good identification**					**02757142**
Actinomyces israelii		95.48	0.38	GLU 99		
Bifidobacterium adolescentis		4.52	0.09			
02757500	**Good identification**					**02757500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.40	GLU 99		
02757502	**Low discrimination**					**02757502**
Bifidobacterium adolescentis		54.91	0.38	GLU 99		
Actinomyces israelii		45.09	0.43	MNS 87		
02757512	**Acceptable identification**					**02757512**
Bifidobacterium adolescentis		90.88	0.40	GLU 99		
Actinomyces israelii		9.13	0.29	MNS 87		
02757542	**Low discrimination**					**02757542**
Actinomyces israelii		72.10	0.36	GLU 99		
Bifidobacterium adolescentis		27.90	0.23			
02771000	**Low discrimination**					**02771000**
					GRA	
Actinomyces naeslundii		65.65	0.35	GLU 99	99	
Fusobacterium mortiferum		34.35	0.19	MNS 88	1	
02771001	**Good identification**					**02771001**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02771002	**Low discrimination**					**02771002**
Actinomyces naeslundii		68.99	0.35	GLU 99		
Lactobacillus acidophilus		18.80	0.01	MNS 88		
Actinomyces israelii		12.21	0.13			
02771003	**Good identification**					**02771003**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02771020	**Good identification**					**02771020**
Actinomyces naeslundii		100.0	0.35	GLU 99		
02771021	**Good identification**					**02771021**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02771022	**Low discrimination**					**02771022**
Actinomyces naeslundii		78.59	0.35	GLU 99		
Lactobacillus acidophilus		21.41	0.01	MNS 88		
02771023	**Good identification**					**02771023**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02771100	**Low discrimination**					**02771100**
					GRA	
Actinomyces naeslundii		62.71	0.43	GLU 99	99	
Fusobacterium mortiferum		26.90	0.25	MNS 88	1	
Actinomyces israelii		10.40	0.21		99	
02771101	**Good identification**					**02771101**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02771103	**Good identification**					**02771103**
Actinomyces naeslundii		100.0	0.46	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
02771120	**Low discrimination**					**02771120**
					SPO	
Actinomyces naeslundii		89.61	0.43	GLU 99	1	
Clostridium ramosum		10.39	0.02	MNS 88	92	
02771121	**Good identification**					**02771121**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02771123	**Good identification**					**02771123**
Actinomyces naeslundii		100.0	0.46	GLU 99		
02773000	**Low discrimination**					**02773000**
Actinomyces naeslundii		74.39	0.40	GLU 99		
Actinomyces israelii		25.61	0.23	MNS 88		
02773001	**Good identification**					**02773001**
Actinomyces naeslundii		100.0	0.43	GLU 99		
02773002	**Low discrimination**					**02773002**
Actinomyces naeslundii		65.94	0.40	GLU 99		
Actinomyces israelii		34.06	0.26	MNS 88		
02773003	**Good identification**					**02773003**
Actinomyces naeslundii		100.0	0.43	GLU 99		
02773020	**Good identification**					**02773020**
Actinomyces naeslundii		100.0	0.40	GLU 99		
02773021	**Good identification**					**02773021**
Actinomyces naeslundii		100.0	0.43	GLU 99		
02773022	**Good identification**					**02773022**
Actinomyces naeslundii		100.0	0.40	GLU 99		
02773023	**Good identification**					**02773023**
Actinomyces naeslundii		100.0	0.43	GLU 99		
02773100	**Low discrimination**					**02773100**
Actinomyces naeslundii		67.39	0.48	GLU 99		
Actinomyces israelii		32.61	0.33	MNS 88		
02773101	**Good identification**					**02773101**
Actinomyces naeslundii		100.0	0.51	GLU 99		
02773102	**Low discrimination**					**02773102**
Actinomyces naeslundii		56.86	0.48	GLU 99		
Actinomyces israelii		41.27	0.36	MNS 88		
Bifidobacterium adolescentis		1.87	0.07			
02773103	**Good identification**					**02773103**
Actinomyces naeslundii		100.0	0.51	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
02773120	**Low discrimination**					**02773120**
					SPO	
Actinomyces naeslundii		74.71	0.48	GLU 99	1	
Clostridium ramosum		25.29	0.15	MNS 88	92	
02773121	**Good identification**					**02773121**
Actinomyces naeslundii		100.0	0.51	GLU 99		
02773123	**Good identification**					**02773123**
Actinomyces naeslundii		100.0	0.51	GLU 99		
02773141	**Good identification**					**02773141**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02773143	**Good identification**					**02773143**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02773160	**Good identification**					**02773160**
Actinomyces naeslundii		100.0	0.35	GLU 99		
02773161	**Good identification**					**02773161**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02773162	**Good identification**					**02773162**
Actinomyces naeslundii		100.0	0.35	GLU 99		
02773163	**Good identification**					**02773163**
Actinomyces naeslundii		100.0	0.38	GLU 99		
02775100	**Acceptable identification**					**02775100**
					SPO	
Actinomyces israelii		90.89	0.39	GLU 99	1	
Clostridium butyricum		9.12	0.01		99	
02775102	**Low discrimination**					**02775102**
					SPO	
Actinomyces israelii		82.73	0.42	GLU 99	1	
Bifidobacterium adolescentis		15.12	0.23		1	
Clostridium butyricum		2.15	-.07		99	
02775502	**Low discrimination**					**02775502**
					SPO	
Bifidobacterium adolescentis		51.23	0.37	GLU 99	1	
Actinomyces israelii		34.28	0.40	MNS 87	1	
Clostridium butyricum		9.81	0.10		99	
02775512	**Low discrimination**					**02775512**
Bifidobacterium adolescentis		70.31	0.38	GLU 99		
Clostridium clostridioforme		23.94	0.27	MNS 87		
Actinomyces israelii		5.75	0.26			
02775712	**Good identification**					**02775712**
Clostridium clostridioforme		100.0	0.42	GLU 88		
02775732	**Good identification**					**02775732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		98.25	0.42	GLU 88		
Parabacteroides distasonis		1.75	0.18	MNS 99		
02777000	**Good identification**					**02777000**
Actinomyces israelii		100.0	0.41	GLU 99		
02777002	**Good identification**					**02777002**
Actinomyces israelii		100.0	0.44	GLU 99		
02777042	**Good identification**					**02777042**
Actinomyces israelii		100.0	0.37	GLU 99		
02777100	**Good identification**					**02777100**
Actinomyces israelii		100.0	0.52	GLU 99		
02777102	**Good identification**					**02777102**
Actinomyces israelii		97.43	0.55	GLU 99		
Bifidobacterium adolescentis		2.57	0.21			
02777110	**Good identification**					**02777110**
Actinomyces israelii		100.0	0.37	GLU 99		
02777112	**Low discrimination**					**02777112**
Actinomyces israelii		82.25	0.41	GLU 99		
Bifidobacterium adolescentis		17.75	0.23	MLZ 13		
02777140	**Good identification**					**02777140**
Actinomyces israelii		100.0	0.45	GLU 99		
02777142	**Good identification**					**02777142**
Actinomyces israelii		100.0	0.48	GLU 99		
02777302	**Good identification**					**02777302**
Actinomyces israelii		100.0	0.35	GLU 99		
02777400	**Good identification**					**02777400**
Actinomyces israelii		100.0	0.40	GLU 99		
02777402	**Good identification**					**02777402**
Actinomyces israelii		100.0	0.43	GLU 99		
02777442	**Good identification**					**02777442**
Actinomyces israelii		100.0	0.36	GLU 99		
02777500	**Good identification**					**02777500**
					SPO	
Actinomyces israelii		97.88	0.50	GLU 99	1	
Clostridium butyricum		2.12	0.01		99	
02777502	**Low discrimination**					**02777502**
					SPO	
Actinomyces israelii		81.87	0.53	GLU 99	1	

Code Taxon	ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium adolescentis	17.67	0.35		1			
Clostridium butyricum	0.46	-.07		99			
02777510 Good identification							**02777510**
Actinomyces israelii	100.0	0.36	GLU 99				
02777512 Low discrimination							**02777512**
Bifidobacterium adolescentis	62.56	0.37	GLU 99				
Actinomyces israelii	35.45	0.39	MNS 87				
Bifidobacterium dentium	1.99	-.07					
02777540 Good identification							**02777540**
Actinomyces israelii	100.0	0.43	GLU 99				
02777542 Acceptable identification							**02777542**
Actinomyces israelii	93.58	0.46	GLU 99				
Bifidobacterium adolescentis	6.42	0.20					
03000000 Low discrimination							**03000000**
				COC	GRA	SPO	
Clostridium botulinum biovar A	37.03	0.76	SUC 7	1	99	99	
Megasphaera elsdenii	18.00	0.75		99	1	1	
Peptostreptococcus anaerobius	17.25	0.66		99	99	1	
03000001 Low discrimination							**03000001**
				GRA	SPO		
Clostridium novyi biovar A	56.52	0.66	SUC 1	99	99		
Clostridium botulinum biovar A	11.20	0.42		99	99		
Campylobacter ureolyticus	10.20	0.32		1	1		
03000002 Low discrimination							**03000002**
				COC	SPO	GRA	
Atopobium minutum	76.06	0.75	SUC 5	1	1	99	
Megasphaera elsdenii	13.34	0.59		99	1	1	
Clostridium sporogenes	2.66	0.40		1	99	99	
03000003 Low discrimination							**03000003**
				COC	SPO	GRA	
Clostridium novyi biovar A	65.92	0.53	SUC 1	1	99	99	
Atopobium minutum	22.80	0.41		1	1	99	
Megasphaera elsdenii	4.00	0.25		99	1	1	
03000010 Low discrimination							**03000010**
				COC	GRA		
Clostridium novyi biovar A	65.57	0.66	SUC 1	1	99		
Clostridium botulinum biovar A	12.99	0.42		1	99		
Megasphaera elsdenii	6.32	0.41		99	1		
03000011 Very good identification							**03000011**
				GRA	SPO		
Clostridium novyi biovar A	99.34	0.66	SUC 1	99	99		
Clostridium botulinum biovar A	0.20	0.08		99	99		
Campylobacter ureolyticus	0.18	-.03		1	1		
03000012 Low discrimination							**03000012**
				COC	SPO	GRA	
Clostridium novyi biovar A	69.67	0.53	SUC 1	1	99	99	
Atopobium minutum	24.10	0.41		1	1	99	
Megasphaera elsdenii	4.23	0.25		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03000013	**Good identification**							**03000013**
					SPO			
Clostridium novyi biovar A		99.65	0.53	SUC 1	99			
Atopobium minutum		0.35	0.07		1			
03000020	**Low discrimination**							**03000020**
					SPO			
Atopobium minutum		60.24	0.70	SUC 5	1			
Clostridium cadaveris		21.35	0.52		99			
Clostridium novyi biovar A		4.98	0.56		99			
03000021	**Low discrimination**							**03000021**
					SPO			
Clostridium novyi biovar A		82.95	0.56	SUC 1	99			
Atopobium minutum		10.13	0.36		1			
Clostridium cadaveris		3.59	0.18		99			
03000022	**Good identification**							**03000022**
					GRA	SPO		
Atopobium minutum		90.96	0.67	SUC 5	99	1		
Pseudoflavonifractor capillosus		5.33	0.32		1	1		
Clostridium novyi biovar A		1.99	0.43		99	99		
03000023	**Low discrimination**							**03000023**
					GRA	SPO		
Clostridium novyi biovar A		67.19	0.43	SUC 1	99	99		
Atopobium minutum		30.99	0.33		99	1		
Pseudoflavonifractor capillosus		1.82	-.03		1	1		
03000030	**Low discrimination**							**03000030**
					SPO			
Clostridium novyi biovar A		84.07	0.56	SUC 1	99			
Atopobium minutum		10.27	0.36		1			
Clostridium cadaveris		3.64	0.18		99			
03000031	**Good identification**							**03000031**
Clostridium novyi biovar A		100.0	0.56	SUC 1				
03000032	**Low discrimination**							**03000032**
					GRA	SPO		
Clostridium novyi biovar A		67.19	0.43	SUC 1	99	99		
Atopobium minutum		30.99	0.33		99	1		
Pseudoflavonifractor capillosus		1.82	-.03		1	1		
03000033	**Good identification**							**03000033**
Clostridium novyi biovar A		100.0	0.43	SUC 1				
03000040	**Low discrimination**							**03000040**
					COC	GRA		
Megasphaera elsdenii		48.24	0.72	FRU 95	99	1		
Clostridium botulinum biovar A		44.45	0.67		1	99		
Clostridium novyi biovar A		5.06	0.63		1	99		
03000041	**Low discrimination**							**03000041**
					COC	GRA		
Clostridium novyi biovar A		83.99	0.63	SUC 1	1	99		
Megasphaera elsdenii		8.09	0.38		99	1		
Clostridium botulinum biovar A		7.46	0.33		1	99		
03000042	**Low discrimination**							**03000042**
					COC	SPO	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		74.31	0.56	FRU 95	99	1	1	
Clostridium novyi biovar A		12.37	0.50	BGL 10	1	99	99	
Atopobium minutum		6.42	0.41		1	1	99	
03000043	**Acceptable identification**							**03000043**
					COC	SPO	GRA	
Clostridium novyi biovar A		93.37	0.50	SUC 1	1	99	99	
Megasphaera elsdenii		5.66	0.22		99	1	1	
Atopobium minutum		0.49	0.07		1	1	99	
03000050	**Low discrimination**							**03000050**
					COC	GRA		
Clostridium novyi biovar A		83.87	0.63	SUC 1	1	99		
Megasphaera elsdenii		8.08	0.38		99	1		
Clostridium botulinum biovar A		7.45	0.33		1	99		
03000051	**Very good identification**							**03000051**
					COC	GRA		
Clostridium novyi biovar A		99.81	0.63	SUC 1	1	99		
Megasphaera elsdenii		0.10	0.04		99	1		
Clostridium botulinum biovar A		0.09	-.01		1	99		
03000052	**Acceptable identification**							**03000052**
					COC	SPO	GRA	
Clostridium novyi biovar A		93.37	0.50	SUC 1	1	99	99	
Megasphaera elsdenii		5.66	0.22		99	1	1	
Atopobium minutum		0.49	0.07		1	1	99	
03000053	**Good identification**							**03000053**
Clostridium novyi biovar A		100.0	0.50	SUC 1				
03000060	**Low discrimination**							**03000060**
					COC	GRA		
Clostridium novyi biovar A		48.65	0.53	SUC 1	1	99		
Megasphaera elsdenii		18.75	0.38		99	1		
Clostridium botulinum biovar A		17.27	0.33		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03000061	**Good identification**							**03000061**
					COC	GRA		
Clostridium novyi biovar A		99.26	0.53	SUC 1	1	99		
Megasphaera elsdenii		0.39	0.04		99	1		
Clostridium botulinum biovar A		0.36	-.01		1	99		
03000062	**Low discrimination**							**03000062**
					COC	SPO	GRA	
Clostridium novyi biovar A		49.63	0.40	SUC 1	1	99	99	
Atopobium minutum		34.34	0.33		1	1	99	
Megasphaera elsdenii		12.04	0.22		99	1	1	
03000063	**Good identification**							**03000063**
Clostridium novyi biovar A		100.0	0.40	SUC 1				
03000070	**Good identification**							**03000070**
					COC	GRA		
Clostridium novyi biovar A		98.40	0.53	SUC 1	1	99		
Clostridium botulinum biovar B		0.86	0.03		1	99		
Megasphaera elsdenii		0.38	0.04		99	1		
03000071	**Good identification**							**03000071**
Clostridium novyi biovar A		100.0	0.53	SUC 1				
03000072	**Good identification**							**03000072**
Clostridium novyi biovar A		100.0	0.40	SUC 1				
03000073	**Good identification**							**03000073**
Clostridium novyi biovar A		100.0	0.40	SUC 1				
03000100	**Low discrimination**							**03000100**
					COC	SPO	GRA	
Actinomyces meyeri		88.96	0.69	FRU 95	1	1	99	
Clostridium botulinum biovar A		3.81	0.42		1	99	99	
Megasphaera elsdenii		1.85	0.41		99	1	1	
03000101	**Low discrimination**							**03000101**
					SPO			
Actinomyces meyeri		72.08	0.35	FRU 95	1			
Clostridium novyi biovar A		15.58	0.32	URE 1	99			
Clostridium botulinum biovar A		3.09	0.08		99			
03000102	**Low discrimination**							**03000102**
					COC	GRA		
Actinomyces meyeri		62.56	0.43	FRU 95	1	99		
Atopobium minutum		26.68	0.41	BGL 3	1	99		
Megasphaera elsdenii		4.68	0.25		99	1		
03000110	**Low discrimination**							**03000110**
					SPO			
Actinomyces meyeri		75.16	0.35	FRU 95	1			
Clostridium novyi biovar A		16.24	0.32	MLZ 1	99			
Clostridium botulinum biovar A		3.22	0.08		99			
03000120	**Acceptable identification**							**03000120**
					SPO			
Actinomyces meyeri		92.52	0.55	FRU 95	1			
Atopobium minutum		4.48	0.36	NAG 12	1			
Clostridium cadaveris		1.59	0.18		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03000140	**Low discrimination**							**03000140**
					COC	SPO	GRA	
Megasphaera elsdenii		35.83	0.38	FRU 95	99	1	1	
Clostridium botulinum biovar A		33.02	0.33	LAC 1	1	99	99	
Actinomyces meyeri		26.08	0.35		1	1	99	
03000200	**Low discrimination**							**03000200**
					COC	GRA	SPO	
Clostridium botulinum biovar A		37.05	0.42	SUC 7	1	99	99	
Megasphaera elsdenii		18.02	0.41	RHA 1	99	1	1	
Peptostreptococcus anaerobius		17.27	0.32		99	99	1	
03000202	**Low discrimination**							**03000202**
					COC	SPO	GRA	
Atopobium minutum		77.67	0.41	SUC 5	1	1	99	
Megasphaera elsdenii		13.62	0.25	RHA 1	99	1	1	
Clostridium sporogenes		2.72	0.06		1	99	99	
03000220	**Low discrimination**							**03000220**
					SPO			
Atopobium minutum		61.17	0.36	SUC 5	1			
Clostridium cadaveris		21.67	0.18	RHA 1	99			
Clostridium novyi biovar A		5.06	0.21		99			
03000240	**Low discrimination**							**03000240**
					COC	GRA		
Megasphaera elsdenii		48.66	0.38	FRU 95	99	1		
Clostridium botulinum biovar A		44.84	0.33	RHA 1	1	99		
Clostridium novyi biovar A		5.10	0.29		1	99		
03000300	**Low discrimination**							**03000300**
					COC	SPO	GRA	
Actinomyces meyeri		89.74	0.35	FRU 95	1	1	99	
Clostridium botulinum biovar A		3.84	0.08	RHA 1	1	99	99	
Megasphaera elsdenii		1.87	0.07		99	1	1	
03000400	**Low discrimination**							**03000400**
					COC	SPO	GRA	
Actinomyces meyeri		33.60	0.49	FRU 95	1	1	99	
Clostridium botulinum biovar A		23.02	0.42		1	99	99	
Megasphaera elsdenii		11.19	0.41		99	1	1	
03000402	**Low discrimination**							**03000402**
					COC	GRA		
Atopobium minutum		68.63	0.41	SUC 5	1	99		
Megasphaera elsdenii		12.03	0.25	ARA 1	99	1		
Actinomyces meyeri		10.06	0.23		1	99		
03000440	**Low discrimination**							**03000440**
					COC	GRA		
Megasphaera elsdenii		47.41	0.38	FRU 95	99	1		
Clostridium botulinum biovar A		43.69	0.33	ARA 1	1	99		
Clostridium novyi biovar A		4.97	0.29		1	99		
03000500	**Very good identification**							**03000500**
					SPO			
Actinomyces meyeri		99.37	0.59	FRU 95	1			
Clostridium botulinum biovar A		0.17	0.08		99			
Lactobacillus fermentum		0.12	0.20		1			
03000520	**Good identification**							**03000520**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		100.0	0.44	FRU 95		
03001000	**Low discrimination**					**03001000**
					COC	
Atopobium minutum		52.45	0.70	SUC 5	1	
Actinomyces meyeri		30.12	0.62		1	
Gemella morbillorum		4.49	0.47		99	
03001001	**Low discrimination**					**03001001**
					SPO	
Clostridium novyi biovar A		64.64	0.50	SUC 1	99	
Atopobium minutum		17.76	0.36	GAL 10	1	
Actinomyces meyeri		10.20	0.28		1	
03001002	**Good identification**					**03001002**
Atopobium minutum		91.15	0.67	SUC 5		
Lactobacillus fermentum		3.86	0.50			
Actinomyces meyeri		2.43	0.36			
03001010	**Low discrimination**					**03001010**
					SPO	
Clostridium novyi biovar A		66.39	0.50	SUC 1	99	
Atopobium minutum		18.24	0.36	GAL 10	1	
Actinomyces meyeri		10.48	0.28		1	
03001011	**Good identification**					**03001011**
Clostridium novyi biovar A		100.0	0.50	SUC 1		
03001013	**Good identification**					**03001013**
Clostridium novyi biovar A		100.0	0.37	SUC 1		
03001020	**Low discrimination**					**03001020**
					COC	
Atopobium minutum		74.15	0.62	SUC 5	1	
Actinomyces meyeri		17.42	0.47		1	
Gemella morbillorum		2.12	0.31		99	
03001021	**Low discrimination**					**03001021**
					SPO	
Clostridium novyi biovar A		68.85	0.39	SUC 1	99	
Atopobium minutum		25.22	0.28	GAL 10	1	
Actinomyces meyeri		5.93	0.13		1	
03001022	**Good identification**					**03001022**
					GRA	
Atopobium minutum		92.27	0.59	SUC 5	99	
Lactobacillus fermentum		3.91	0.42		99	
Pseudoflavonifractor capillosus		1.80	0.15		1	
03001030	**Low discrimination**					**03001030**
					SPO	
Clostridium novyi biovar A		68.85	0.39	SUC 1	99	
Atopobium minutum		25.22	0.28	GAL 10	1	
Actinomyces meyeri		5.93	0.13		1	
03001031	**Good identification**					**03001031**
Clostridium novyi biovar A		100.0	0.39	SUC 1		
03001040	**Low discrimination**					**03001040**
					COC GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		29.15	0.47	SUC 1	1	99	
Megasphaera elsdenii		25.27	0.38	GAL 10	99	1	
Clostridium botulinum biovar A		23.29	0.33		1	99	
03001041	**Good identification**						**03001041**
					COC	GRA	
Clostridium novyi biovar A		98.35	0.47	SUC 1	1	99	
Megasphaera elsdenii		0.86	0.04	GAL 10	99	1	
Clostridium botulinum biovar A		0.79	-.01		1	99	
03001050	**Good identification**						**03001050**
					COC	GRA	
Clostridium novyi biovar A		98.35	0.47	SUC 1	1	99	
Megasphaera elsdenii		0.86	0.04	GAL 10	99	1	
Clostridium botulinum biovar A		0.79	-.01		1	99	
03001051	**Good identification**						**03001051**
Clostridium novyi biovar A		100.0	0.47	SUC 1			
03001060	**Low discrimination**						**03001060**
					SPO		
Clostridium novyi biovar A		57.28	0.36	SUC 1	99		
Atopobium minutum		31.47	0.28	GAL 10	1		
Actinomyces meyeri		7.39	0.13		1		
03001061	**Good identification**						**03001061**
Clostridium novyi biovar A		100.0	0.36	SUC 1			
03001070	**Good identification**						**03001070**
Clostridium novyi biovar A		100.0	0.36	SUC 1			
03001071	**Good identification**						**03001071**
Clostridium novyi biovar A		100.0	0.36	SUC 1			
03001100	**Very good identification**						**03001100**
					GRA		
Actinomyces meyeri		96.43	0.72	FRU 95	99		
Lactobacillus fermentum		1.93	0.54		99		
Fusobacterium mortiferum		1.06	0.48		1		
03001101	**Low discrimination**						**03001101**
					SPO		
Clostridium chauvoei		52.98	0.37	MNS 90	99		
Actinomyces meyeri		44.91	0.38	MLT 99	1		
Lactobacillus fermentum		0.90	0.20		1		
03001102	**Low discrimination**						**03001102**
Actinomyces meyeri		56.11	0.46	FRU 95			
Lactobacillus fermentum		36.36	0.54	BGL 3			
Atopobium minutum		5.32	0.33				
03001110	**Acceptable identification**						**03001110**
					SPO		
Actinomyces meyeri		94.58	0.38	FRU 95	1		
Lactobacillus fermentum		1.90	0.20	MLZ 1	1		
Clostridium novyi biovar A		1.51	0.15		99		
03001120	**Good identification**						**03001120**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		93.15	0.58	FRU 95	99	
Lactobacillus fermentum		4.56	0.46	NAG 12	99	
Prevotella melaninogenica		1.21	0.17		1	
03001122	**Low discrimination**					**03001122**
Lactobacillus fermentum		54.71	0.46	FRU 99		
Actinomyces meyeri		34.54	0.32			
Atopobium minutum		8.00	0.25			
03001140	**Acceptable identification**					**03001140**
					GRA	
Actinomyces meyeri		94.39	0.38	FRU 95	99	
Lactobacillus fermentum		1.89	0.20	SOR 1	99	
Fusobacterium mortiferum		1.04	0.14		1	
03001200	**Low discrimination**					**03001200**
					COC	
Atopobium minutum		52.72	0.36	SUC 5	1	
Actinomyces meyeri		30.27	0.28	RHA 1	1	
Gemella morbillorum		4.51	0.13		99	
03001300	**Good identification**					**03001300**
					GRA	
Actinomyces meyeri		96.99	0.38	FRU 95	99	
Lactobacillus fermentum		1.94	0.20	RHA 1	99	
Fusobacterium mortiferum		1.07	0.14		1	
03001400	**Low discrimination**					**03001400**
Actinomyces meyeri		74.26	0.52	FRU 95		
Lactobacillus fermentum		14.60	0.50			
Atopobium minutum		5.23	0.36			
03001402	**Low discrimination**					**03001402**
Lactobacillus fermentum		71.63	0.50	FRU 99		
Atopobium minutum		17.10	0.33			
Actinomyces meyeri		11.27	0.26			
03001420	**Low discrimination**					**03001420**
Actinomyces meyeri		58.60	0.37	FRU 95		
Lactobacillus fermentum		28.16	0.42	NAG 12		
Atopobium minutum		10.08	0.28			
03001422	**Low discrimination**					**03001422**
Lactobacillus fermentum		76.74	0.42	FRU 99		
Atopobium minutum		18.32	0.25			
Actinomyces meyeri		4.94	0.11			
03001500	**Good identification**					**03001500**
					GRA	
Actinomyces meyeri		92.54	0.62	FRU 95	99	
Lactobacillus fermentum		7.42	0.54		99	
Fusobacterium mortiferum		0.04	0.14		1	
03001502	**Low discrimination**					**03001502**
Lactobacillus fermentum		72.16	0.54	FRU 99		
Actinomyces meyeri		27.84	0.36			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03001520	**Low discrimination**							**03001520**
Actinomyces meyeri		83.61	0.47	FRU 95				
Lactobacillus fermentum		16.39	0.46	NAG 12				
03001522	**Low discrimination**							**03001522**
Lactobacillus fermentum		86.37	0.46	FRU 99				
Actinomyces meyeri		13.63	0.21					
03002000	**Good identification**							**03002000**
					COC	GRA	SPO	
Megasphaera elsdenii		94.60	0.75	FRU 95	99	1	1	
Clostridium botulinum biovar A		1.97	0.42		1	99	99	
Peptostreptococcus anaerobius		0.92	0.32		99	99	1	
03002001	**Low discrimination**							**03002001**
					COC	GRA		
Megasphaera elsdenii		85.44	0.41	FRU 95	99	1		
Clostridium novyi biovar A		8.96	0.32	URE 1	1	99		
Clostridium botulinum biovar A		1.78	0.08		1	99		
03002002	**Good identification**							**03002002**
					COC	GRA		
Megasphaera elsdenii		93.35	0.59	FRU 95	99	1		
Atopobium minutum		5.38	0.41	BGL 10	1	99		
Pseudoramibacter alactolyticus		0.62	0.11		1	99		
03002010	**Low discrimination**							**03002010**
					COC	GRA		
Megasphaera elsdenii		78.22	0.41	FRU 95	99	1		
Clostridium difficile		9.93	0.19	MLZ 1	1	99		
Clostridium novyi biovar A		8.20	0.32		1	99		
03002020	**Low discrimination**							**03002020**
					COC	GRA		
Megasphaera elsdenii		61.35	0.41	FRU 95	99	1		
Atopobium minutum		19.44	0.36	NAG 1	1	99		
Pseudoramibacter alactolyticus		7.09	0.14		1	99		
03002040	**Very good identification**							**03002040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.70	0.72	FRU 95	99	1	1	
Clostridium botulinum biovar A		0.92	0.33		1	99	99	
Pseudoramibacter alactolyticus		0.25	0.17		1	1	99	
03002041	**Low discrimination**							**03002041**
					COC	GRA		
Megasphaera elsdenii		89.75	0.38	FRU 95	99	1		
Clostridium novyi biovar A		9.41	0.29	URE 1	1	99		
Clostridium botulinum biovar A		0.84	-.01		1	99		
03002042	**Good identification**							**03002042**
					COC	SPO	GRA	
Megasphaera elsdenii		99.42	0.56	FRU 95	99	1	1	
Pseudoramibacter alactolyticus		0.25	0.00	BGL 10	1	1	99	
Clostridium novyi biovar A		0.17	0.16		1	99	99	
03002050	**Low discrimination**							**03002050**
					COC	GRA		
Megasphaera elsdenii		88.96	0.38	FRU 95	99	1		
Clostridium novyi biovar A		9.33	0.29	MLZ 1	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		0.89	-.03		1	99		
03002060	**Acceptable identification**							**03002060**
					COC	SPO	GRA	
Megasphaera elsdenii		92.69	0.38	FRU 95	99	1	1	
Pseudoramibacter alactolyticus		4.02	0.04	NAG 1	1	1	99	
Clostridium novyi biovar A		2.43	0.18		1	99	99	
03002100	**Low discrimination**							**03002100**
					COC	SPO	GRA	
Megasphaera elsdenii		64.72	0.41	FRU 95	99	1	1	
Actinomyces meyeri		31.40	0.35	LAC 1	1	1	99	
Clostridium botulinum biovar A		1.34	0.08		1	99	99	
03002140	**Good identification**							**03002140**
					COC	SPO	GRA	
Megasphaera elsdenii		97.92	0.38	FRU 95	99	1	1	
Propionibacterium propionicum		1.17	0.15	LAC 1	1	1	99	
Clostridium botulinum biovar A		0.91	-.01		1	99	99	
03002200	**Good identification**							**03002200**
					COC	SPO	GRA	
Megasphaera elsdenii		95.54	0.41	FRU 95	99	1	1	
Clostridium botulinum biovar A		1.99	0.08	RHA 1	1	99	99	
Atopobium minutum		0.92	0.10		1	1	99	
03002240	**Good identification**							**03002240**
					COC	GRA		
Megasphaera elsdenii		99.08	0.38	FRU 95	99	1		
Clostridium botulinum biovar A		0.92	-.01	RHA 1	1	99		
03002400	**Acceptable identification**							**03002400**
					COC	SPO	GRA	
Megasphaera elsdenii		92.37	0.41	FRU 95	99	1	1	
Actinomyces meyeri		2.80	0.15	ARA 1	1	1	99	
Clostridium botulinum biovar A		1.92	0.08		1	99	99	
03002440	**Good identification**							**03002440**
					COC	GRA		
Megasphaera elsdenii		99.08	0.38	FRU 95	99	1		
Clostridium botulinum biovar A		0.92	-.01	ARA 1	1	99		
03003000	**Low discrimination**							**03003000**
					COC	GRA		
Megasphaera elsdenii		61.77	0.41	FRU 95	99	1		
Atopobium minutum		19.57	0.36	GAL 1	1	99		
Actinomyces meyeri		11.24	0.28		1	99		
03003040	**Good identification**							**03003040**
					COC	GRA		
Megasphaera elsdenii		97.95	0.38	FRU 95	99	1		
Clostridium novyi biovar A		1.14	0.12	GAL 1	1	99		
Clostridium botulinum biovar A		0.91	-.01		1	99		
03003100	**Acceptable identification**							**03003100**
					COC	GRA		
Actinomyces meyeri		94.71	0.38	FRU 95	1	99		
Lactobacillus fermentum		1.90	0.20	MAN 1	1	99		
Megasphaera elsdenii		1.32	0.07		99	1		
03004000	**Low discrimination**							**03004000**
					COC	SPO	GRA	
Actinomyces meyeri		78.80	0.64	FRU 95	1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		6.95	0.42		1	99	99	
Megasphaera elsdenii		3.38	0.41		99	1	1	
03004002	**Low discrimination**							**03004002**
					COC	GRA		
Actinomyces meyeri		46.58	0.38	FRU 95	1	99		
Atopobium minutum		40.94	0.41	BGL 3	1	99		
Megasphaera elsdenii		7.18	0.25		99	1		
03004020	**Low discrimination**							**03004020**
					SPO			
Actinomyces meyeri		85.08	0.49	FRU 95	1			
Atopobium minutum		8.48	0.36	NAG 12	1			
Clostridium cadaveris		3.01	0.18		99			
03004040	**Low discrimination**							**03004040**
					COC	GRA		
Clostridium glycolicum		78.88	0.49	SUC 1	1	99		
Megasphaera elsdenii		8.74	0.38	FRU 90	99	1		
Clostridium botulinum biovar A		8.05	0.33		1	99		
03004100	**Very good identification**							**03004100**
					GRA	SPO		
Actinomyces meyeri		99.89	0.74	FRU 95	99	1		
Fusobacterium mortiferum		0.04	0.26		1	1		
Clostridium botulinum biovar A		0.02	0.08		99	99		
03004101	**Good identification**							**03004101**
Actinomyces meyeri		100.0	0.40	FRU 95				
03004102	**Good identification**							**03004102**
Actinomyces meyeri		99.48	0.48	FRU 95				
Lactobacillus fermentum		0.30	0.16	BGL 3				
Atopobium minutum		0.22	0.07					
03004110	**Good identification**							**03004110**
Actinomyces meyeri		100.0	0.40	FRU 95				
03004120	**Very good identification**							**03004120**
Actinomyces meyeri		100.0	0.59	FRU 95				
03004140	**Acceptable identification**							**03004140**
					COC	SPO	GRA	
Actinomyces meyeri		92.74	0.40	FRU 95	1	1	99	
Clostridium glycolicum		5.98	0.15	SOR 1	1	99	99	
Megasphaera elsdenii		0.66	0.04		99	1	1	
03004300	**Good identification**							**03004300**
Actinomyces meyeri		100.0	0.40	FRU 95				
03004400	**Good identification**							**03004400**
					COC	SPO		
Actinomyces meyeri		98.03	0.54	FRU 95	1	1		
Gemella morbillorum		1.04	0.19		99	1		
Clostridium botulinum biovar A		0.35	0.08		1	99		
03004420	**Good identification**							**03004420**
					COC			
Actinomyces meyeri		99.15	0.39	FRU 95	1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Gemella morbillorum		0.85	0.03	NAG 12	99	
03004500	**Very good identification**					**03004500**
Actinomyces meyeri		99.96	0.64	FRU 95		
Lactobacillus fermentum		0.04	0.16			
03004502	**Good identification**					**03004502**
Actinomyces meyeri		98.83	0.38	FRU 95		
Lactobacillus fermentum		1.17	0.16	BGL 3		
03004520	**Good identification**					**03004520**
Actinomyces meyeri		100.0	0.49	FRU 95		
03005000	**Very good identification**					**03005000**
Actinomyces meyeri		96.53	0.67	FRU 95		
Lactobacillus fermentum		1.44	0.46			
Atopobium minutum		0.88	0.36			
03005002	**Low discrimination**					**03005002**
Actinomyces meyeri		59.67	0.41	FRU 95		
Lactobacillus fermentum		28.68	0.46	BGL 3		
Atopobium minutum		11.65	0.33			
03005020	**Acceptable identification**					**03005020**
Actinomyces meyeri		93.86	0.52	FRU 95		
Lactobacillus fermentum		3.41	0.38	NAG 12		
Atopobium minutum		2.08	0.28			
03005022	**Low discrimination**					**03005022**
Lactobacillus fermentum		44.30	0.38	FRU 99		
Actinomyces meyeri		37.70	0.26			
Atopobium minutum		18.00	0.25			
03005100	**Very good identification**					**03005100**
					GRA	
Actinomyces meyeri		99.36	0.77	FRU 95	99	
Lactobacillus fermentum		0.60	0.50		99	
Fusobacterium mortiferum		0.04	0.29		1	
03005101	**Good identification**					**03005101**
					SPO	
Actinomyces meyeri		98.79	0.43	FRU 95	1	
Clostridium chauvoei		0.61	0.03	URE 1	99	
Lactobacillus fermentum		0.60	0.16		1	
03005102	**Low discrimination**					**03005102**
					GRA	
Actinomyces meyeri		83.50	0.51	FRU 95	99	
Lactobacillus fermentum		16.37	0.50	BGL 3	99	
Fusobacterium mortiferum		0.13	0.13		1	
03005110	**Good identification**					**03005110**
Actinomyces meyeri		99.40	0.43	FRU 95		
Lactobacillus fermentum		0.60	0.16	MLZ 1		
03005120	**Very good identification**					**03005120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		98.54	0.62	FRU 95		
Lactobacillus fermentum		1.46	0.42	NAG 12		
03005122	**Low discrimination**					**03005122**
Actinomyces meyeri		67.60	0.37	FRU 95		
Lactobacillus fermentum		32.40	0.42	NAG 12		
03005140	**Good identification**					**03005140**
					SPO	
Actinomyces meyeri		98.93	0.43	FRU 95	1	
Lactobacillus fermentum		0.60	0.16	SOR 1	1	
Clostridium glycolicum		0.47	-.02		99	
03005300	**Good identification**					**03005300**
Actinomyces meyeri		99.40	0.43	FRU 95		
Lactobacillus fermentum		0.60	0.16	RHA 1		
03005400	**Good identification**					**03005400**
					COC	
Actinomyces meyeri		94.05	0.57	FRU 95	1	
Lactobacillus fermentum		5.59	0.46		1	
Gemella morbillorum		0.36	0.14		99	
03005402	**Low discrimination**					**03005402**
Lactobacillus fermentum		65.78	0.46	FRU 99		
Actinomyces meyeri		34.22	0.31			
03005420	**Low discrimination**					**03005420**
Actinomyces meyeri		87.31	0.42	FRU 95		
Lactobacillus fermentum		12.69	0.38	NAG 12		
03005422	**Low discrimination**					**03005422**
Lactobacillus fermentum		82.46	0.38	FRU 99		
Actinomyces meyeri		17.55	0.16			
03005500	**Very good identification**					**03005500**
Actinomyces meyeri		97.63	0.67	FRU 95		
Lactobacillus fermentum		2.37	0.50			
03005502	**Low discrimination**					**03005502**
Actinomyces meyeri		56.05	0.41	FRU 95		
Lactobacillus fermentum		43.95	0.50	BGL 3		
03005520	**Acceptable identification**					**03005520**
Actinomyces meyeri		94.30	0.52	FRU 95		
Lactobacillus fermentum		5.59	0.42	NAG 12		
Bifidobacterium longum subsp. longum		0.11	0.06			
03005522	**Low discrimination**					**03005522**
Lactobacillus fermentum		64.89	0.42	FRU 99		
Actinomyces meyeri		33.85	0.26			
Bifidobacterium longum subsp. longum		1.27	0.06			
03006000	**Low discrimination**					**03006000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO	GRA	
Megasphaera elsdenii		77.66	0.41	FRU 95	99	1	1	
Actinomyces meyeri		18.29	0.30	XYL 1	1	1	99	
Clostridium botulinum biovar A		1.61	0.08		1	99	99	
03006040	**Acceptable identification**							**03006040**
					COC	GRA		
Megasphaera elsdenii		90.87	0.38	FRU 95	99	1		
Clostridium glycolicum		8.28	0.15	XYL 1	1	99		
Clostridium botulinum biovar A		0.85	-.01		1	99		
03006100	**Good identification**							**03006100**
					COC	GRA		
Actinomyces meyeri		98.94	0.40	FRU 95	1	99		
Megasphaera elsdenii		1.06	0.07	MAN 1	99	1		
03007100	**Good identification**							**03007100**
Actinomyces meyeri		99.40	0.43	FRU 95				
Lactobacillus fermentum		0.60	0.16	MAN 1				
03010000	**Low discrimination**							**03010000**
					COC	GRA		
Megasphaera elsdenii		78.65	0.97		99	1		
Actinomyces meyeri		9.54	0.81		1	99		
Pseudoramibacter alactolyticus		5.03	0.66		1	99		
03010001	**Low discrimination**							**03010001**
					COC	GRA		
Megasphaera elsdenii		74.76	0.63	URE 1	99	1		
Actinomyces meyeri		9.07	0.47		1	99		
Pseudoramibacter alactolyticus		4.78	0.32		1	99		
03010002	**Low discrimination**							**03010002**
					COC	GRA		
Megasphaera elsdenii		71.41	0.81	BGL 10	99	1		
Atopobium minutum		19.79	0.75		1	99		
Pseudoramibacter alactolyticus		4.57	0.50		1	99		
03010003	**Low discrimination**							**03010003**
					COC	SPO	GRA	
Megasphaera elsdenii		67.04	0.47	URE 1	99	1	1	
Atopobium minutum		18.58	0.40	BGL 10	1	1	99	
Clostridium novyi biovar A		6.46	0.37		1	99	99	
03010010	**Low discrimination**							**03010010**
					COC	SPO	GRA	
Megasphaera elsdenii		70.04	0.63	MLZ 1	99	1	1	
Actinomyces meyeri		8.50	0.47		1	1	99	
Clostridium difficile		5.15	0.36		1	99	99	
03010011	**Low discrimination**							**03010011**
					COC	SPO	GRA	
Clostridium novyi biovar A		82.16	0.50	SUC 1	1	99	99	
Megasphaera elsdenii		13.67	0.29	FRU 10	99	1	1	
Actinomyces meyeri		1.66	0.13		1	1	99	
03010012	**Low discrimination**							**03010012**
					COC	SPO	GRA	
Megasphaera elsdenii		64.47	0.47	MLZ 1	99	1	1	
Atopobium minutum		17.86	0.40	BGL 10	1	1	99	
Clostridium novyi biovar A		6.22	0.37		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03010013	**Low discrimination**							**03010013**
					COC	SPO	GRA	
Clostridium novyi biovar A		88.20	0.37	SUC 1	1	99	99	
Megasphaera elsdenii		9.24	0.12	FRU 10	99	1	1	
Atopobium minutum		2.56	0.06		1	1	99	
03010020	**Low discrimination**							**03010020**
Actinomyces meyeri		27.19	0.66	NAG 12				
Atopobium minutum		25.30	0.69					
Pseudoramibacter alactolyticus		18.56	0.53					
03010022	**Low discrimination**							**03010022**
					COC	GRA		
Atopobium minutum		75.45	0.66	SUC 5	1	99		
Pseudoramibacter alactolyticus		9.22	0.37		1	99		
Megasphaera elsdenii		8.25	0.47		99	1		
03010031	**Good identification**							**03010031**
Clostridium novyi biovar A		100.0	0.39	SUC 1				
03010040	**Very good identification**							**03010040**
					COC	SPO	GRA	
Megasphaera elsdenii		96.79	0.94		99	1	1	
Pseudoramibacter alactolyticus		2.32	0.56		1	1	99	
Clostridium botulinum biovar A		0.30	0.47		1	99	99	
03010041	**Good identification**							**03010041**
					COC	SPO	GRA	
Megasphaera elsdenii		91.53	0.60	URE 1	99	1	1	
Clostridium novyi biovar A		5.56	0.47		1	99	99	
Pseudoramibacter alactolyticus		2.20	0.21		1	1	99	
03010042	**Very good identification**							**03010042**
					COC	GRA		
Megasphaera elsdenii		97.04	0.78	BGL 10	99	1		
Pseudoramibacter alactolyticus		2.33	0.39		1	99		
Atopobium minutum		0.41	0.40		1	99		
03010043	**Low discrimination**							**03010043**
					COC	SPO	GRA	
Megasphaera elsdenii		88.92	0.44	URE 1	99	1	1	
Clostridium novyi biovar A		8.57	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.13	0.05		1	1	99	
03010050	**Good identification**							**03010050**
					COC	SPO	GRA	
Megasphaera elsdenii		90.25	0.60	MLZ 1	99	1	1	
Clostridium novyi biovar A		5.48	0.47		1	99	99	
Pseudoramibacter alactolyticus		2.17	0.21		1	1	99	
03010051	**Low discrimination**							**03010051**
					COC	GRA		
Clostridium novyi biovar A		85.73	0.47	SUC 1	1	99		
Megasphaera elsdenii		14.27	0.26	FRU 10	99	1		
03010052	**Low discrimination**							**03010052**
					COC	SPO	GRA	
Megasphaera elsdenii		88.92	0.44	MLZ 1	99	1	1	
Clostridium novyi biovar A		8.57	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.13	0.05		1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03010060	**Low discrimination**							**03010060**
					COC	GRA		
Megasphaera elsdenii		66.17	0.60	NAG 1	99	1		
Pseudoramibacter alactolyticus		27.73	0.43		1	99		
Actinomyces meyeri		1.64	0.32		1	99		
03010061	**Low discrimination**							**03010061**
					COC	SPO	GRA	
Clostridium novyi biovar A		51.42	0.36	SUC 1	1	99	99	
Megasphaera elsdenii		34.23	0.26	FRU 10	99	1	1	
Pseudoramibacter alactolyticus		14.35	0.09		1	1	99	
03010062	**Low discrimination**							**03010062**
					COC	GRA		
Megasphaera elsdenii		62.95	0.44	NAG 1	99	1		
Pseudoramibacter alactolyticus		26.38	0.26	BGL 10	1	99		
Atopobium minutum		8.72	0.32		1	99		
03010070	**Low discrimination**							**03010070**
					COC	GRA		
Clostridium novyi biovar A		44.26	0.36	SUC 1	1	99		
Megasphaera elsdenii		29.46	0.26	FRU 10	99	1		
Clostridium botulinum biovar B		13.93	0.13		1	99		
03010071	**Good identification**							**03010071**
Clostridium novyi biovar A		100.0	0.36	SUC 1				
03010100	**Good identification**							**03010100**
					COC	GRA		
Actinomyces meyeri		92.38	0.91		1	99		
Fusobacterium mortiferum		5.29	0.79		1	1		
Megasphaera elsdenii		1.92	0.63		99	1		
03010101	**Good identification**							**03010101**
					COC	GRA		
Actinomyces meyeri		92.13	0.57	URE 1	1	99		
Fusobacterium mortiferum		5.27	0.45		1	1		
Megasphaera elsdenii		1.92	0.29		99	1		
03010102	**Low discrimination**							**03010102**
					COC	GRA		
Actinomyces meyeri		73.63	0.65	BGL 3	1	99		
Fusobacterium mortiferum		15.14	0.63		1	1		
Megasphaera elsdenii		5.51	0.47		99	1		
03010110	**Good identification**							**03010110**
					COC	GRA		
Actinomyces meyeri		92.19	0.57	MLZ 1	1	99		
Fusobacterium mortiferum		5.28	0.45		1	1		
Megasphaera elsdenii		1.92	0.29		99	1		
03010120	**Very good identification**							**03010120**
					GRA			
Actinomyces meyeri		98.54	0.76	NAG 12	99			
Fusobacterium mortiferum		0.42	0.45		1			
Lactobacillus fermentum		0.38	0.46		99			
03010121	**Good identification**							**03010121**
					GRA	SPO		
Actinomyces meyeri		99.05	0.42	NAG 12	99	1		
Clostridium chauvoei		0.53	0.01	URE 1	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		0.42	0.11		1	1		
03010122	**Low discrimination**							**03010122**
Actinomyces meyeri		83.15	0.51	NAG 12				
Lactobacillus fermentum		10.40	0.46	BGL 3				
Atopobium minutum		4.21	0.32					
03010130	**Good identification**							**03010130**
					GRA			
Actinomyces meyeri		99.58	0.42	MLZ 1	99			
Fusobacterium mortiferum		0.42	0.11	NAG 12	1			
03010140	**Low discrimination**							**03010140**
					COC	GRA		
Megasphaera elsdenii		55.50	0.60	LAC 1	99	1		
Actinomyces meyeri		40.39	0.57		1	99		
Fusobacterium mortiferum		2.31	0.45		1	1		
03010142	**Low discrimination**							**03010142**
					COC	GRA		
Megasphaera elsdenii		77.44	0.44	LAC 1	99	1		
Actinomyces meyeri		15.69	0.31	BGL 10	1	99		
Fusobacterium mortiferum		3.23	0.29		1	1		
03010160	**Low discrimination**							**03010160**
					COC	GRA		
Actinomyces meyeri		87.06	0.42	NAG 12	1	99		
Megasphaera elsdenii		8.86	0.26	SOR 1	99	1		
Pseudoramibacter alactolyticus		3.71	0.09		1	99		
03010200	**Low discrimination**							**03010200**
					COC	GRA		
Megasphaera elsdenii		78.15	0.63	RHA 1	99	1		
Actinomyces meyeri		9.48	0.47		1	99		
Pseudoramibacter alactolyticus		5.00	0.32		1	99		
03010202	**Low discrimination**							**03010202**
					COC	GRA		
Megasphaera elsdenii		69.84	0.47	RHA 1	99	1		
Atopobium minutum		19.35	0.40	BGL 10	1	99		
Pseudoramibacter alactolyticus		4.47	0.15		1	99		
03010240	**Very good identification**							**03010240**
					COC	SPO	GRA	
Megasphaera elsdenii		96.85	0.60	RHA 1	99	1	1	
Pseudoramibacter alactolyticus		2.32	0.21		1	1	99	
Clostridium botulinum biovar A		0.30	0.13		1	99	99	
03010242	**Good identification**							**03010242**
					COC	GRA		
Megasphaera elsdenii		97.26	0.44	RHA 1	99	1		
Pseudoramibacter alactolyticus		2.33	0.05	BGL 10	1	99		
Atopobium minutum		0.41	0.06		1	99		
03010300	**Good identification**							**03010300**
					COC	GRA		
Actinomyces meyeri		92.15	0.57	RHA 1	1	99		
Fusobacterium mortiferum		5.28	0.45		1	1		
Megasphaera elsdenii		1.92	0.29		99	1		
03010320	**Good identification**							**03010320**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		99.58	0.42	RHA 1	99			
Fusobacterium mortiferum		0.42	0.11	NAG 12	1			
03010400	**Low discrimination**							**03010400**
					COC	GRA		
Actinomyces meyeri		70.97	0.71		1	99		
Megasphaera elsdenii		23.64	0.63		99	1		
Pseudoramibacter alactolyticus		1.51	0.32		1	99		
03010401	**Low discrimination**							**03010401**
					COC	GRA		
Actinomyces meyeri		70.55	0.36	URE 1	1	99		
Megasphaera elsdenii		23.50	0.29		99	1		
Pseudoramibacter alactolyticus		1.50	-.03		1	99		
03010402	**Low discrimination**							**03010402**
					COC	GRA		
Megasphaera elsdenii		38.24	0.47	ARA 1	99	1		
Actinomyces meyeri		31.95	0.45	BGL 10	1	99		
Lactobacillus fermentum		16.03	0.50		1	99		
03010410	**Low discrimination**							**03010410**
					COC	SPO	GRA	
Actinomyces meyeri		69.36	0.36	MLZ 1	1	1	99	
Megasphaera elsdenii		23.10	0.29		99	1	1	
Clostridium difficile		1.70	0.02		1	99	99	
03010420	**Low discrimination**							**03010420**
Actinomyces meyeri		86.73	0.56	NAG 12				
Lactobacillus fermentum		3.29	0.42					
Atopobium minutum		3.26	0.35					
03010422	**Low discrimination**							**03010422**
Lactobacillus fermentum		38.03	0.42	GAL 99				
Actinomyces meyeri		31.01	0.30					
Atopobium minutum		25.14	0.32					
03010440	**Good identification**							**03010440**
					COC	GRA		
Megasphaera elsdenii		92.82	0.60	ARA 1	99	1		
Actinomyces meyeri		4.22	0.36		1	99		
Pseudoramibacter alactolyticus		2.23	0.21		1	99		
03010442	**Good identification**							**03010442**
					COC	GRA		
Megasphaera elsdenii		95.49	0.44	ARA 1	99	1		
Pseudoramibacter alactolyticus		2.29	0.05	BGL 10	1	99		
Actinomyces meyeri		1.21	0.11		1	99		
03010500	**Very good identification**							**03010500**
					GRA			
Actinomyces meyeri		99.05	0.81		99			
Lactobacillus fermentum		0.63	0.54		99			
Fusobacterium mortiferum		0.23	0.45		1			
03010501	**Good identification**							**03010501**
					GRA			
Actinomyces meyeri		99.14	0.47	URE 1	99			
Lactobacillus fermentum		0.63	0.20		99			
Fusobacterium mortiferum		0.23	0.11		1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
03010502	**Low discrimination**						**03010502**
					GRA		
Actinomyces meyeri		82.18	0.55	BGL 3	99		
Lactobacillus fermentum		16.82	0.54		99		
Fusobacterium mortiferum		0.68	0.29		1		
03010510	**Good identification**						**03010510**
					GRA		
Actinomyces meyeri		99.14	0.47	MLZ 1	99		
Lactobacillus fermentum		0.63	0.20		99		
Fusobacterium mortiferum		0.23	0.11		1		
03010520	**Very good identification**						**03010520**
					GRA		
Actinomyces meyeri		98.46	0.66	NAG 12	99		
Lactobacillus fermentum		1.52	0.46		99		
Fusobacterium mortiferum		0.02	0.11		1		
03010522	**Low discrimination**						**03010522**
Actinomyces meyeri		66.66	0.40	NAG 12			
Lactobacillus fermentum		33.34	0.46	BGL 3			
03010540	**Acceptable identification**						**03010540**
					COC	GRA	
Actinomyces meyeri		93.97	0.47	SOR 1	1	99	
Megasphaera elsdenii		5.22	0.26		99	1	
Lactobacillus fermentum		0.60	0.20		1	99	
03010600	**Low discrimination**						**03010600**
					COC	GRA	
Actinomyces meyeri		71.57	0.36	RHA 1	1	99	
Megasphaera elsdenii		23.84	0.29		99	1	
Pseudoramibacter alactolyticus		1.53	-.03		1	99	
03010700	**Good identification**						**03010700**
					GRA		
Actinomyces meyeri		99.14	0.47	RHA 1	99		
Lactobacillus fermentum		0.63	0.20		99		
Fusobacterium mortiferum		0.23	0.11		1		
03011000	**Low discrimination**						**03011000**
					GRA		
Actinomyces meyeri		65.75	0.84		99		
Lactobacillus fermentum		16.83	0.85		99		
Fusobacterium mortiferum		6.45	0.76		1		
03011001	**Low discrimination**						**03011001**
					GRA		
Actinomyces meyeri		63.84	0.50	URE 1	99		
Lactobacillus fermentum		16.35	0.50		99		
Fusobacterium mortiferum		6.27	0.42		1		
03011002	**Low discrimination**						**03011002**
Lactobacillus fermentum		70.41	0.85				
Atopobium minutum		15.51	0.66				
Actinomyces meyeri		8.51	0.58				
03011003	**Low discrimination**						**03011003**
Lactobacillus fermentum		69.28	0.50	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		15.26	0.32				
Actinomyces meyeri		8.37	0.24				
03011010	**Low discrimination**						**03011010**
					CAT		
Actinomyces meyeri		59.15	0.50	MLZ 1	1		
Lactobacillus fermentum		15.14	0.50		1		
Propionibacterium granulosum		8.24	0.41		90		
03011012	**Low discrimination**						**03011012**
					CAT		
Lactobacillus fermentum		63.72	0.50	MLZ 1	1		
Atopobium minutum		14.04	0.32		1		
Propionibacterium granulosum		8.67	0.31		90		
03011020	**Low discrimination**						**03011020**
Actinomyces meyeri		53.07	0.69	NAG 12			
Lactobacillus fermentum		33.21	0.76				
Atopobium minutum		10.98	0.61				
03011022	**Low discrimination**						**03011022**
Lactobacillus fermentum		78.05	0.76				
Atopobium minutum		17.19	0.58				
Actinomyces meyeri		3.86	0.43				
03011023	**Low discrimination**						**03011023**
Lactobacillus fermentum		77.59	0.42	URE 1			
Atopobium minutum		17.09	0.24				
Actinomyces meyeri		3.83	0.09				
03011032	**Low discrimination**						**03011032**
					CAT		
Lactobacillus fermentum		71.90	0.42	MLZ 1	1		
Atopobium minutum		15.84	0.24		1		
Propionibacterium granulosum		7.33	0.21		90		
03011040	**Low discrimination**						**03011040**
					COC	GRA	
Megasphaera elsdenii		68.64	0.60	GAL 1	99	1	
Actinomyces meyeri		18.73	0.50		1	99	
Lactobacillus fermentum		4.80	0.50		1	99	
03011042	**Low discrimination**						**03011042**
					COC	GRA	
Megasphaera elsdenii		52.32	0.44	GAL 1	99	1	
Lactobacillus fermentum		32.90	0.50	BGL 10	1	99	
Atopobium minutum		7.25	0.32		1	99	
03011062	**Low discrimination**						**03011062**
Lactobacillus fermentum		74.73	0.42	SOR 1			
Atopobium minutum		16.46	0.24				
Actinomyces meyeri		3.69	0.09				
03011100	**Low discrimination**						**03011100**
					GRA		
Actinomyces meyeri		86.05	0.94		99		
Lactobacillus fermentum		8.99	0.88		99		
Fusobacterium mortiferum		4.93	0.82		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03011101	**Low discrimination**					**03011101**
					SPO	
Actinomyces meyeri		76.51	0.60	URE 1	1	
Clostridium chauvoei		11.08	0.44		99	
Lactobacillus fermentum		7.99	0.54		1	
03011102	**Low discrimination**					**03011102**
					GRA	
Lactobacillus fermentum		73.60	0.88		99	
Actinomyces meyeri		21.80	0.68		99	
Fusobacterium mortiferum		4.48	0.66		1	
03011103	**Low discrimination**					**03011103**
					SPO	
Lactobacillus fermentum		62.22	0.54	URE 1	1	
Actinomyces meyeri		18.43	0.34		1	
Clostridium chauvoei		15.23	0.31		99	
03011110	**Low discrimination**					**03011110**
					GRA	
Actinomyces meyeri		85.96	0.60	MLZ 1	99	
Lactobacillus fermentum		8.98	0.54		99	
Fusobacterium mortiferum		4.92	0.48		1	
03011112	**Low discrimination**					**03011112**
					GRA	
Lactobacillus fermentum		73.69	0.54	MLZ 1	99	
Actinomyces meyeri		21.82	0.34		99	
Fusobacterium mortiferum		4.49	0.32		1	
03011120	**Low discrimination**					**03011120**
					GRA	
Actinomyces meyeri		78.91	0.79	NAG 12	99	
Lactobacillus fermentum		20.14	0.80		99	
Prevotella melaninogenica		0.48	0.33		1	
03011121	**Low discrimination**					**03011121**
					SPO	
Actinomyces meyeri		61.58	0.45	NAG 12	1	
Clostridium chauvoei		21.81	0.35	URE 1	99	
Lactobacillus fermentum		15.72	0.46		1	
03011122	**Low discrimination**					**03011122**
					GRA	
Lactobacillus fermentum		88.42	0.80		99	
Actinomyces meyeri		10.71	0.54		99	
Prevotella melaninogenica		0.41	0.21		1	
03011123	**Low discrimination**					**03011123**
					SPO	
Lactobacillus fermentum		72.36	0.46	URE 1	1	
Clostridium chauvoei		17.71	0.22		99	
Actinomyces meyeri		8.77	0.19		1	
03011130	**Low discrimination**					**03011130**
					GRA	
Actinomyces meyeri		79.01	0.45	MLZ 1	99	
Lactobacillus fermentum		20.17	0.46	NAG 12	99	
Prevotella melaninogenica		0.49	-.01		1	
03011132	**Low discrimination**					**03011132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		89.19	0.46	MLZ 1		
Actinomyces meyeri		10.81	0.19			
03011140	**Low discrimination**					**03011140**
					GRA	
Actinomyces meyeri		85.36	0.60	SOR 1	99	
Lactobacillus fermentum		8.91	0.54		99	
Fusobacterium mortiferum		4.89	0.48		1	
03011142	**Low discrimination**					**03011142**
					GRA	
Lactobacillus fermentum		72.72	0.54	SOR 1	99	
Actinomyces meyeri		21.54	0.34		99	
Fusobacterium mortiferum		4.43	0.32		1	
03011160	**Low discrimination**					**03011160**
Actinomyces meyeri		77.77	0.45	NAG 12		
Lactobacillus fermentum		19.85	0.46	SOR 1		
Clostridium perfringens		1.57	0.15			
03011162	**Low discrimination**					**03011162**
Lactobacillus fermentum		86.33	0.46	SOR 1		
Actinomyces meyeri		10.46	0.19			
Clostridium perfringens		3.21	0.09			
03011200	**Low discrimination**					**03011200**
					GRA	
Actinomyces meyeri		64.20	0.50	RHA 1	99	
Lactobacillus fermentum		16.44	0.50		99	
Fusobacterium mortiferum		6.30	0.42		1	
03011202	**Low discrimination**					**03011202**
Lactobacillus fermentum		67.43	0.50	RHA 1		
Atopobium minutum		14.85	0.32			
Actinomyces meyeri		8.15	0.24			
03011222	**Low discrimination**					**03011222**
Lactobacillus fermentum		76.07	0.42	RHA 1		
Atopobium minutum		16.76	0.24			
Actinomyces meyeri		3.76	0.09			
03011300	**Low discrimination**					**03011300**
					GRA	
Actinomyces meyeri		85.97	0.60	RHA 1	99	
Lactobacillus fermentum		8.98	0.54		99	
Fusobacterium mortiferum		4.92	0.48		1	
03011302	**Low discrimination**					**03011302**
					GRA	
Lactobacillus fermentum		73.22	0.54	RHA 1	99	
Actinomyces meyeri		21.69	0.34		99	
Fusobacterium mortiferum		4.46	0.32		1	
03011320	**Low discrimination**					**03011320**
					GRA	
Actinomyces meyeri		79.01	0.45	RHA 1	99	
Lactobacillus fermentum		20.17	0.46	NAG 12	99	
Prevotella melaninogenica		0.49	-.01		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03011322	**Low discrimination**					**03011322**
Lactobacillus fermentum		89.19	0.46	RHA 1		
Actinomyces meyeri		10.81	0.19			
03011400	**Low discrimination**					**03011400**
					COC	
Lactobacillus fermentum		50.24	0.85		1	
Actinomyces meyeri		49.06	0.74		1	
Gemella morbillorum		0.21	0.32		99	
03011401	**Low discrimination**					**03011401**
Lactobacillus fermentum		50.60	0.50	URE 1		
Actinomyces meyeri		49.41	0.39			
03011402	**Very good identification**					**03011402**
Lactobacillus fermentum		96.79	0.85			
Actinomyces meyeri		2.92	0.48			
Atopobium minutum		0.22	0.32			
03011403	**Good identification**					**03011403**
Lactobacillus fermentum		97.07	0.50	URE 1		
Actinomyces meyeri		2.93	0.14			
03011410	**Low discrimination**					**03011410**
Lactobacillus fermentum		50.60	0.50	MLZ 1		
Actinomyces meyeri		49.41	0.39			
03011412	**Good identification**					**03011412**
Lactobacillus fermentum		97.07	0.50	MLZ 1		
Actinomyces meyeri		2.93	0.14			
03011420	**Low discrimination**					**03011420**
Lactobacillus fermentum		71.20	0.76			
Actinomyces meyeri		28.44	0.59			
Atopobium minutum		0.24	0.27			
03011421	**Low discrimination**					**03011421**
Lactobacillus fermentum		71.46	0.42	URE 1		
Actinomyces meyeri		28.55	0.25			
03011422	**Very good identification**					**03011422**
Lactobacillus fermentum		98.56	0.76			
Actinomyces meyeri		1.22	0.33			
Atopobium minutum		0.22	0.24			
03011423	**Good identification**					**03011423**
Lactobacillus fermentum		100.0	0.42	URE 1		
03011430	**Low discrimination**					**03011430**
Lactobacillus fermentum		71.46	0.42	MLZ 1		
Actinomyces meyeri		28.55	0.25			
03011432	**Good identification**					**03011432**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		100.0	0.42	MLZ 1			
03011440	**Low discrimination**						**03011440**
					COC	GRA	
Lactobacillus fermentum		47.15	0.50	SOR 1	1	99	
Actinomyces meyeri		46.04	0.39		1	99	
Megasphaera elsdenii		6.82	0.26		99	1	
03011442	**Good identification**						**03011442**
					COC	GRA	
Lactobacillus fermentum		95.58	0.50	SOR 1	1	99	
Actinomyces meyeri		2.89	0.14		1	99	
Megasphaera elsdenii		1.54	0.09		99	1	
03011460	**Low discrimination**						**03011460**
Lactobacillus fermentum		71.46	0.42	SOR 1			
Actinomyces meyeri		28.55	0.25				
03011462	**Good identification**						**03011462**
Lactobacillus fermentum		100.0	0.42	SOR 1			
03011500	**Low discrimination**						**03011500**
					GRA		
Actinomyces meyeri		70.42	0.84		99		
Lactobacillus fermentum		29.42	0.88		99		
Fusobacterium mortiferum		0.16	0.48		1		
03011501	**Low discrimination**						**03011501**
					SPO		
Actinomyces meyeri		70.13	0.50	URE 1	1		
Lactobacillus fermentum		29.30	0.54		1		
Clostridium chauvoei		0.41	0.09		99		
03011502	**Good identification**						**03011502**
					GRA		
Lactobacillus fermentum		93.05	0.88		99		
Actinomyces meyeri		6.89	0.58		99		
Fusobacterium mortiferum		0.06	0.32		1		
03011503	**Acceptable identification**						**03011503**
Lactobacillus fermentum		93.11	0.54	URE 1			
Actinomyces meyeri		6.89	0.24				
03011510	**Low discrimination**						**03011510**
					GRA		
Actinomyces meyeri		70.32	0.50	MLZ 1	99		
Lactobacillus fermentum		29.37	0.54		99		
Fusobacterium mortiferum		0.16	0.14		1		
03011512	**Acceptable identification**						**03011512**
Lactobacillus fermentum		92.69	0.54	MLZ 1			
Actinomyces meyeri		6.86	0.24				
Bifidobacterium longum subsp. longum		0.45	0.07				
03011520	**Low discrimination**						**03011520**
Lactobacillus fermentum		50.50	0.80				
Actinomyces meyeri		49.46	0.69				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		0.02	0.14			
03011521	**Low discrimination**					**03011521**
					SPO	
Lactobacillus fermentum		50.17	0.46	URE 1	1	
Actinomyces meyeri		49.13	0.35		1	
Clostridium chauvoei		0.70	0.01		99	
03011522	**Very good identification**					**03011522**
Lactobacillus fermentum		97.03	0.80			
Actinomyces meyeri		2.94	0.43			
Bifidobacterium longum subsp. longum		0.03	0.14			
03011523	**Good identification**					**03011523**
Lactobacillus fermentum		97.06	0.46	URE 1		
Actinomyces meyeri		2.94	0.09			
03011530	**Low discrimination**					**03011530**
Lactobacillus fermentum		49.07	0.46	MLZ 1		
Actinomyces meyeri		48.06	0.35			
Bifidobacterium longum subsp. longum		2.87	0.18			
03011532	**Acceptable identification**					**03011532**
Lactobacillus fermentum		91.85	0.46	MLZ 1		
Bifidobacterium longum subsp. longum		5.37	0.18			
Actinomyces meyeri		2.78	0.09			
03011540	**Low discrimination**					**03011540**
					GRA	
Actinomyces meyeri		70.42	0.50	SOR 1	99	
Lactobacillus fermentum		29.42	0.54		99	
Fusobacterium mortiferum		0.16	0.14		1	
03011542	**Acceptable identification**					**03011542**
Lactobacillus fermentum		93.11	0.54	SOR 1		
Actinomyces meyeri		6.89	0.24			
03011560	**Low discrimination**					**03011560**
Lactobacillus fermentum		50.52	0.46	SOR 1		
Actinomyces meyeri		49.48	0.35			
03011562	**Good identification**					**03011562**
Lactobacillus fermentum		97.06	0.46	SOR 1		
Actinomyces meyeri		2.94	0.09			
03011600	**Low discrimination**					**03011600**
Lactobacillus fermentum		50.60	0.50	RHA 1		
Actinomyces meyeri		49.41	0.39			
03011602	**Good identification**					**03011602**
Lactobacillus fermentum		97.07	0.50	RHA 1		
Actinomyces meyeri		2.93	0.14			
03011620	**Low discrimination**					**03011620**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		71.46	0.42	RHA 1				
Actinomyces meyeri		28.55	0.25					
03011622	**Good identification**							**03011622**
Lactobacillus fermentum		100.0	0.42	RHA 1				
03011700	**Low discrimination**							**03011700**
					GRA			
Actinomyces meyeri		70.42	0.50	RHA 1	99			
Lactobacillus fermentum		29.42	0.54		99			
Fusobacterium mortiferum		0.16	0.14		1			
03011702	**Acceptable identification**							**03011702**
Lactobacillus fermentum		93.11	0.54	RHA 1				
Actinomyces meyeri		6.89	0.24					
03011720	**Low discrimination**							**03011720**
Lactobacillus fermentum		50.52	0.46	RHA 1				
Actinomyces meyeri		49.48	0.35					
03011722	**Good identification**							**03011722**
Lactobacillus fermentum		97.06	0.46	RHA 1				
Actinomyces meyeri		2.94	0.09					
03012000	**Good identification**							**03012000**
					COC	GRA		
Megasphaera elsdenii		94.24	0.97		99	1		
Pseudoramibacter alactolyticus		3.25	0.61		1	99		
Eubacterium limosum		2.06	0.60		1	99		
03012001	**Good identification**							**03012001**
					COC	GRA		
Megasphaera elsdenii		94.21	0.63	URE 1	99	1		
Pseudoramibacter alactolyticus		3.25	0.27		1	99		
Eubacterium limosum		2.06	0.26		1	99		
03012002	**Very good identification**							**03012002**
					COC	CAT	GRA	
Megasphaera elsdenii		95.87	0.81	BGL 10	99	1	1	
Pseudoramibacter alactolyticus		3.30	0.45		1	1	99	
Propionibacterium granulosum		0.31	0.44		1	90	99	
03012003	**Good identification**							**03012003**
					COC	CAT	GRA	
Megasphaera elsdenii		96.11	0.47	URE 1	99	1	1	
Pseudoramibacter alactolyticus		3.31	0.11	BGL 10	1	1	99	
Propionibacterium granulosum		0.31	0.09		1	90	99	
03012010	**Low discrimination**							**03012010**
					COC	SPO	GRA	
Megasphaera elsdenii		57.51	0.63	MLZ 1	99	1	1	
Clostridium difficile		38.02	0.53		1	99	99	
Pseudoramibacter alactolyticus		1.98	0.27		1	1	99	
03012012	**Low discrimination**							**03012012**
					COC	SPO	CAT	
Megasphaera elsdenii		87.69	0.47	MLZ 1	99	1	1	
Clostridium difficile		5.27	0.19	BGL 10	1	99	1	
Propionibacterium granulosum		3.77	0.29		1	1	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
03012020	**Low discrimination**							**03012020**
					COC	CAT	GRA	
Megasphaera elsdenii		58.65	0.63	NAG 1	99	1	1	
Pseudoramibacter alactolyticus		35.30	0.48		1	1	99	
Propionibacterium granulosum		2.06	0.44		1	90	99	
03012022	**Low discrimination**							**03012022**
					COC	GRA		
Megasphaera elsdenii		56.25	0.47	NAG 1	99	1		
Pseudoramibacter alactolyticus		33.86	0.32	BGL 10	1	99		
Atopobium minutum		5.20	0.32		1	99		
03012040	**Very good identification**							**03012040**
					COC	GRA		
Megasphaera elsdenii		98.67	0.94		99	1		
Pseudoramibacter alactolyticus		1.28	0.51		1	99		
Eubacterium limosum		0.03	0.26		1	99		
03012041	**Very good identification**							**03012041**
					COC	SPO	GRA	
Megasphaera elsdenii		98.67	0.60	URE 1	99	1	1	
Pseudoramibacter alactolyticus		1.28	0.17		1	1	99	
Clostridium novyi biovar A		0.06	0.12		1	99	99	
03012042	**Very good identification**							**03012042**
					COC	GRA		
Megasphaera elsdenii		98.71	0.78	BGL 10	99	1		
Pseudoramibacter alactolyticus		1.28	0.35		1	99		
Propionibacterium propionicum		0.01	0.18		1	99		
03012043	**Good identification**							**03012043**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	URE 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
03012050	**Good identification**							**03012050**
					COC	SPO	GRA	
Megasphaera elsdenii		93.83	0.60	MLZ 1	99	1	1	
Clostridium difficile		4.90	0.31		1	99	99	
Pseudoramibacter alactolyticus		1.21	0.17		1	1	99	
03012052	**Good identification**							**03012052**
					COC	SPO	GRA	
Megasphaera elsdenii		98.26	0.44	MLZ 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.00	BGL 10	1	1	99	
Clostridium difficile		0.47	-.03		1	99	99	
03012060	**Low discrimination**							**03012060**
					COC	CAT	GRA	
Megasphaera elsdenii		81.41	0.60	NAG 1	99	1	1	
Pseudoramibacter alactolyticus		18.37	0.38		1	1	99	
Propionibacterium acnes		0.18	0.11		1	95	99	
03012062	**Low discrimination**							**03012062**
					COC	GRA		
Megasphaera elsdenii		81.59	0.44	NAG 1	99	1		
Pseudoramibacter alactolyticus		18.41	0.22	BGL 10	1	99		
03012100	**Low discrimination**							**03012100**
					COC	GRA		
Megasphaera elsdenii		61.93	0.63	LAC 1	99	1		
Actinomyces meyeri		30.05	0.57		1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		2.56	0.49		1	99		
03012102	**Low discrimination**							**03012102**
					COC	GRA		
Megasphaera elsdenii		56.55	0.47	LAC 1	99	1		
Propionibacterium propionicum		31.57	0.52	BGL 10	1	99		
Actinomyces meyeri		7.64	0.31		1	99		
03012103	**Low discrimination**							**03012103**
					COC	GRA		
Propionibacterium propionicum		86.00	0.36	NIT 86	1	99		
Megasphaera elsdenii		14.01	0.12	MLT 99	99	1		
03012120	**Low discrimination**							**03012120**
					COC	GRA		
Actinomyces meyeri		79.34	0.42	MAN 1	1	99		
Megasphaera elsdenii		12.11	0.29	NAG 12	99	1		
Pseudoramibacter alactolyticus		7.29	0.14		1	99		
03012140	**Good identification**							**03012140**
					COC	GRA		
Megasphaera elsdenii		92.36	0.60	LAC 1	99	1		
Propionibacterium propionicum		5.73	0.49		1	99		
Pseudoramibacter alactolyticus		1.19	0.17		1	99		
03012142	**Low discrimination**							**03012142**
					COC	GRA		
Megasphaera elsdenii		54.05	0.44	LAC 1	99	1		
Propionibacterium propionicum		45.26	0.52	BGL 10	1	99		
Pseudoramibacter alactolyticus		0.70	0.00		1	99		
03012143	**Acceptable identification**							**03012143**
					COC	GRA		
Propionibacterium propionicum		90.21	0.36	NIT 86	1	99		
Megasphaera elsdenii		9.79	0.09	MLT 99	99	1		
03012200	**Good identification**							**03012200**
					COC	GRA		
Megasphaera elsdenii		94.01	0.63	RHA 1	99	1		
Pseudoramibacter alactolyticus		3.24	0.27		1	99		
Eubacterium limosum		2.06	0.26		1	99		
03012202	**Good identification**							**03012202**
					COC	CAT	GRA	
Megasphaera elsdenii		95.51	0.47	RHA 1	99	1	1	
Pseudoramibacter alactolyticus		3.29	0.11	BGL 10	1	1	99	
Propionibacterium granulosum		0.93	0.18		1	90	99	
03012240	**Very good identification**							**03012240**
					COC	GRA		
Megasphaera elsdenii		98.73	0.60	RHA 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.17		1	99		
03012242	**Good identification**							**03012242**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	RHA 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
03012400	**Good identification**							**03012400**
					COC	GRA		
Megasphaera elsdenii		91.68	0.63	ARA 1	99	1		
Pseudoramibacter alactolyticus		3.16	0.27		1	99		
Actinomyces meyeri		2.78	0.36		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03012402	**Acceptable identification**							**03012402**
					COC	GRA		
Megasphaera elsdenii		94.96	0.47	ARA 1	99	1		
Pseudoramibacter alactolyticus		3.27	0.11	BGL 10	1	99		
Actinomyces meyeri		0.80	0.11		1	99		
03012440	**Very good identification**							**03012440**
					COC	GRA		
Megasphaera elsdenii		98.68	0.60	ARA 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.17		1	99		
Actinomyces meyeri		0.05	0.02		1	99		
03012442	**Good identification**							**03012442**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	ARA 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
03012500	**Acceptable identification**							**03012500**
					COC	GRA		
Actinomyces meyeri		91.06	0.47	MAN 1	1	99		
Megasphaera elsdenii		7.58	0.29		99	1		
Lactobacillus fermentum		0.58	0.20		1	99		
03013000	**Low discrimination**							**03013000**
					COC	CAT	GRA	
Megasphaera elsdenii		68.41	0.63	GAL 1	99	1	1	
Actinomyces meyeri		12.45	0.50		1	1	99	
Propionibacterium granulosum		9.59	0.54		1	90	99	
03013002	**Low discrimination**							**03013002**
					COC	CAT	GRA	
Megasphaera elsdenii		51.66	0.47	GAL 1	99	1	1	
Lactobacillus fermentum		21.66	0.50	BGL 10	1	1	99	
Propionibacterium granulosum		16.30	0.44		1	90	99	
03013010	**Low discrimination**							**03013010**
					COC	SPO	CAT	
Propionibacterium granulosum		49.59	0.39	MNS 99	1	1	90	
Megasphaera elsdenii		26.20	0.29	MLZ 12	99	1	1	
Clostridium difficile		17.32	0.19		1	99	1	
03013020	**Low discrimination**							**03013020**
					CAT			
Propionibacterium granulosum		33.89	0.44	MNS 99	90			
Actinomyces meyeri		23.99	0.35		1			
Lactobacillus fermentum		15.01	0.42		1			
03013022	**Low discrimination**							**03013022**
					CAT			
Lactobacillus fermentum		51.29	0.42	MAN 1	1			
Propionibacterium granulosum		28.94	0.33		90			
Atopobium minutum		11.30	0.24		1			
03013040	**Very good identification**							**03013040**
					COC	GRA		
Megasphaera elsdenii		98.12	0.60	GAL 1	99	1		
Pseudoramibacter alactolyticus		1.27	0.17		1	99		
Actinomyces meyeri		0.27	0.16		1	99		
03013042	**Good identification**							**03013042**
					COC	GRA		
Megasphaera elsdenii		96.85	0.44	GAL 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoramibacter alactolyticus		1.25	0.00	BGL 10	1	99	
Propionibacterium propionicum		0.82	0.18		1	99	
03013100	**Low discrimination**						**03013100**
					GRA		
Actinomyces meyeri		80.94	0.60	MAN 1	99		
Lactobacillus fermentum		8.45	0.54		99		
Fusobacterium mortiferum		4.63	0.48		1		
03013102	**Low discrimination**						**03013102**
Lactobacillus fermentum		45.47	0.54	MAN 1			
Propionibacterium propionicum		37.10	0.52				
Actinomyces meyeri		13.47	0.34				
03013103	**Low discrimination**						**03013103**
Propionibacterium propionicum		81.56	0.36	NIT 86			
Actinomyces naeslundii		9.35	0.30	MLT 99			
Lactobacillus fermentum		9.09	0.20	URE 10			
03013120	**Low discrimination**						**03013120**
					GRA		
Actinomyces meyeri		78.20	0.45	MAN 1	99		
Lactobacillus fermentum		19.96	0.46	NAG 12	99		
Prevotella melaninogenica		0.48	-.01		1		
03013122	**Low discrimination**						**03013122**
Lactobacillus fermentum		85.71	0.46	MAN 1			
Actinomyces meyeri		10.39	0.19				
Propionibacterium propionicum		2.12	0.18				
03013140	**Low discrimination**						**03013140**
					COC	GRA	
Propionibacterium propionicum		73.03	0.49	NIT 86	1	99	
Actinomyces meyeri		12.98	0.26	MLT 99	1	99	
Megasphaera elsdenii		11.89	0.26		99	1	
03013142	**Good identification**						**03013142**
					COC	GRA	
Propionibacterium propionicum		97.61	0.52	NIT 86	1	99	
Lactobacillus fermentum		1.21	0.20	MLT 99	1	99	
Megasphaera elsdenii		1.18	0.09		99	1	
03013143	**Good identification**						**03013143**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03013400	**Low discrimination**						**03013400**
					COC	GRA	
Lactobacillus fermentum		44.82	0.50	MAN 1	1	99	
Actinomyces meyeri		43.77	0.39		1	99	
Megasphaera elsdenii		9.72	0.29		99	1	
03013402	**Acceptable identification**						**03013402**
					COC	GRA	
Lactobacillus fermentum		94.17	0.50	MAN 1	1	99	
Actinomyces meyeri		2.84	0.14		1	99	
Megasphaera elsdenii		2.27	0.12		99	1	
03013420	**Low discrimination**						**03013420**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		70.31	0.42	MAN 1	1			
Actinomyces meyeri		28.09	0.25		1			
Propionibacterium granulosum		1.60	0.09		90			
03013422	**Good identification**							**03013422**
Lactobacillus fermentum		100.0	0.42	MAN 1				
03013500	**Low discrimination**							**03013500**
Actinomyces meyeri		70.31	0.50	MAN 1				
Lactobacillus fermentum		29.37	0.54					
Propionibacterium propionicum		0.16	0.15					
03013502	**Acceptable identification**							**03013502**
Lactobacillus fermentum		92.40	0.54	MAN 1				
Actinomyces meyeri		6.84	0.24					
Propionibacterium propionicum		0.76	0.18					
03013520	**Low discrimination**							**03013520**
Lactobacillus fermentum		50.52	0.46	MAN 1				
Actinomyces meyeri		49.48	0.35					
03013522	**Good identification**							**03013522**
Lactobacillus fermentum		97.06	0.46	MAN 1				
Actinomyces meyeri		2.94	0.09					
03014000	**Good identification**							**03014000**
					COC	SPO	GRA	
Actinomyces meyeri		94.09	0.86		1	1	99	
Megasphaera elsdenii		4.04	0.63		99	1	1	
Clostridium glycolicum		0.61	0.43		1	99	99	
03014001	**Acceptable identification**							**03014001**
					COC	SPO	GRA	
Actinomyces meyeri		93.92	0.52	URE 1	1	1	99	
Megasphaera elsdenii		4.03	0.29		99	1	1	
Clostridium glycolicum		0.60	0.09		1	99	99	
03014002	**Low discrimination**							**03014002**
					COC	GRA		
Actinomyces meyeri		79.17	0.60	BGL 3	1	99		
Megasphaera elsdenii		12.20	0.47		99	1		
Atopobium minutum		3.38	0.40		1	99		
03014010	**Acceptable identification**							**03014010**
					COC	SPO	GRA	
Actinomyces meyeri		92.49	0.52	MLZ 1	1	1	99	
Megasphaera elsdenii		3.97	0.29		99	1	1	
Clostridium difficile		1.52	0.15		1	99	99	
03014020	**Very good identification**							**03014020**
Actinomyces meyeri		98.18	0.71	NAG 12				
Atopobium minutum		0.48	0.35					
Pseudoramibacter alactolyticus		0.35	0.19					
03014021	**Good identification**							**03014021**
Actinomyces meyeri		100.0	0.37	NAG 12				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
03014022	**Low discrimination**							**03014022**
Actinomyces meyeri		81.88	0.45	NAG 12				
Atopobium minutum		8.55	0.32	BGL 3				
Lactobacillus fermentum		7.60	0.38					
03014030	**Good identification**							**03014030**
Actinomyces meyeri		100.0	0.37	MLZ 1				
03014040	**Low discrimination**							**03014040**
					COC	SPO	GRA	
Clostridium glycolicum		75.54	0.65	SUC 1	1	99	99	
Megasphaera elsdenii		17.67	0.60		99	1	1	
Actinomyces meyeri		6.24	0.52		1	1	99	
03014042	**Low discrimination**							**03014042**
					COC	SPO	GRA	
Megasphaera elsdenii		65.84	0.44	XYL 1	99	1	1	
Clostridium glycolicum		25.59	0.31	BGL 10	1	99	99	
Actinomyces meyeri		6.47	0.26		1	1	99	
03014060	**Low discrimination**							**03014060**
					COC	SPO	GRA	
Actinomyces meyeri		45.58	0.37	NAG 12	1	1	99	
Clostridium glycolicum		40.86	0.31	SOR 1	1	99	99	
Megasphaera elsdenii		9.56	0.26		99	1	1	
03014100	**Excellent identification**							**03014100**
					GRA			
Actinomyces meyeri		99.72	0.96		99			
Fusobacterium mortiferum		0.22	0.60		1			
Lactobacillus fermentum		0.05	0.50		99			
03014101	**Very good identification**							**03014101**
					GRA			
Actinomyces meyeri		99.73	0.62	URE 1	99			
Fusobacterium mortiferum		0.22	0.26		1			
Lactobacillus fermentum		0.05	0.16		99			
03014102	**Very good identification**							**03014102**
					GRA			
Actinomyces meyeri		97.66	0.70	BGL 3	99			
Lactobacillus fermentum		1.51	0.50		99			
Fusobacterium mortiferum		0.78	0.44		1			
03014103	**Good identification**							**03014103**
Actinomyces meyeri		98.48	0.36	URE 1				
Lactobacillus fermentum		1.52	0.16	BGL 3				
03014110	**Very good identification**							**03014110**
					GRA			
Actinomyces meyeri		99.73	0.62	MLZ 1	99			
Fusobacterium mortiferum		0.22	0.26		1			
Lactobacillus fermentum		0.05	0.16		99			
03014112	**Good identification**							**03014112**
Actinomyces meyeri		98.48	0.36	MLZ 1				
Lactobacillus fermentum		1.52	0.16	BGL 3				
03014120	**Very good identification**							**03014120**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Actinomyces meyeri		99.87	0.81	NAG 12	99			
Lactobacillus fermentum		0.12	0.42		99			
Fusobacterium mortiferum		0.02	0.26		1			
03014121	**Good identification**							**03014121**
Actinomyces meyeri		100.0	0.47	NAG 12				
03014122	**Good identification**							**03014122**
Actinomyces meyeri		96.35	0.55	NAG 12				
Lactobacillus fermentum		3.65	0.42	BGL 3				
03014130	**Good identification**							**03014130**
Actinomyces meyeri		100.0	0.47	MLZ 1				
03014140	**Very good identification**							**03014140**
					COC	SPO	GRA	
Actinomyces meyeri		96.12	0.62	SOR 1	1	1	99	
Clostridium glycolicum		2.94	0.31		1	99	99	
Megasphaera elsdenii		0.69	0.26		99	1	1	
03014142	**Good identification**							**03014142**
					COC	GRA		
Actinomyces meyeri		95.13	0.36	SOR 1	1	99		
Megasphaera elsdenii		2.44	0.09	BGL 3	99	1		
Lactobacillus fermentum		1.47	0.16		1	99		
03014160	**Good identification**							**03014160**
					SPO			
Actinomyces meyeri		99.77	0.47	NAG 12	1			
Clostridium glycolicum		0.23	-.03	SOR 1	99			
03014200	**Acceptable identification**							**03014200**
					COC	GRA		
Actinomyces meyeri		93.22	0.52	RHA 1	1	99		
Megasphaera elsdenii		4.00	0.29		99	1		
Actinomyces odontolyticus		0.99	0.22		1	99		
03014220	**Good identification**							**03014220**
Actinomyces meyeri		98.75	0.37	RHA 1				
Actinomyces odontolyticus		1.25	0.08	NAG 12				
03014300	**Very good identification**							**03014300**
					GRA			
Actinomyces meyeri		99.65	0.62	RHA 1	99			
Fusobacterium mortiferum		0.22	0.26		1			
Actinomyces odontolyticus		0.08	0.13		99			
03014302	**Good identification**							**03014302**
Actinomyces meyeri		96.89	0.36	RHA 1				
Actinomyces odontolyticus		1.61	0.09	BGL 3				
Lactobacillus fermentum		1.50	0.16					
03014320	**Good identification**							**03014320**
Actinomyces meyeri		100.0	0.47	RHA 1				
03014400	**Very good identification**							**03014400**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		99.23	0.76		1	99		
Lactobacillus fermentum		0.47	0.46		1	99		
Megasphaera elsdenii		0.17	0.29		99	1		
03014401	**Good identification**							**03014401**
Actinomyces meyeri		99.53	0.41	URE 1				
Lactobacillus fermentum		0.47	0.12					
03014402	**Low discrimination**							**03014402**
					COC	GRA		
Actinomyces meyeri		86.23	0.50	BGL 3	1	99		
Lactobacillus fermentum		13.09	0.46		1	99		
Megasphaera elsdenii		0.54	0.12		99	1		
03014410	**Good identification**							**03014410**
Actinomyces meyeri		99.53	0.41	MLZ 1				
Lactobacillus fermentum		0.47	0.12					
03014420	**Very good identification**							**03014420**
					COC			
Actinomyces meyeri		98.81	0.61	NAG 12	1			
Lactobacillus fermentum		1.13	0.38		1			
Gemella morbillorum		0.06	0.04		99			
03014440	**Low discrimination**							**03014440**
					COC	SPO	GRA	
Actinomyces meyeri		62.18	0.41	SOR 1	1	1	99	
Clostridium glycolicum		30.41	0.31		1	99	99	
Megasphaera elsdenii		7.11	0.26		99	1	1	
03014500	**Very good identification**							**03014500**
					GRA			
Actinomyces meyeri		99.80	0.86		99			
Lactobacillus fermentum		0.19	0.50		99			
Fusobacterium mortiferum		0.01	0.26		1			
03014501	**Good identification**							**03014501**
Actinomyces meyeri		99.81	0.52	URE 1				
Lactobacillus fermentum		0.19	0.16					
03014502	**Good identification**							**03014502**
Actinomyces meyeri		94.17	0.60	BGL 3				
Lactobacillus fermentum		5.83	0.50					
03014510	**Good identification**							**03014510**
Actinomyces meyeri		99.81	0.52	MLZ 1				
Lactobacillus fermentum		0.19	0.16					
03014520	**Very good identification**							**03014520**
Actinomyces meyeri		99.53	0.71	NAG 12				
Lactobacillus fermentum		0.47	0.42					
03014521	**Good identification**							**03014521**
Actinomyces meyeri		100.0	0.37	NAG 12				
03014522	**Low discrimination**							**03014522**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		86.86	0.45	NAG 12		
Lactobacillus fermentum		13.15	0.42	BGL 3		
03014530	**Good identification**					**03014530**
Actinomyces meyeri		100.0	0.37	MLZ 1		
03014540	**Good identification**					**03014540**
					SPO	
Actinomyces meyeri		99.69	0.52	SOR 1	1	
Lactobacillus fermentum		0.19	0.16		1	
Clostridium glycolicum		0.12	-.03		99	
03014560	**Good identification**					**03014560**
Actinomyces meyeri		100.0	0.37	NAG 12		
03014600	**Good identification**					**03014600**
Actinomyces meyeri		99.53	0.41	RHA 1		
Lactobacillus fermentum		0.47	0.12			
03014700	**Good identification**					**03014700**
Actinomyces meyeri		99.81	0.52	RHA 1		
Lactobacillus fermentum		0.19	0.16			
03014720	**Good identification**					**03014720**
Actinomyces meyeri		100.0	0.37	RHA 1		
03015000	**Good identification**					**03015000**
					GRA	
Actinomyces meyeri		92.33	0.89		99	
Lactobacillus fermentum		7.15	0.81		99	
Fusobacterium mortiferum		0.35	0.57		1	
03015001	**Acceptable identification**					**03015001**
					GRA	
Actinomyces meyeri		92.48	0.55	URE 1	99	
Lactobacillus fermentum		7.16	0.46		99	
Fusobacterium mortiferum		0.35	0.23		1	
03015002	**Low discrimination**					**03015002**
					GRA	
Lactobacillus fermentum		70.96	0.81		99	
Actinomyces meyeri		28.33	0.63		99	
Fusobacterium mortiferum		0.39	0.40		1	
03015003	**Low discrimination**					**03015003**
Lactobacillus fermentum		71.47	0.46	URE 1		
Actinomyces meyeri		28.54	0.29			
03015010	**Acceptable identification**					**03015010**
					GRA	
Actinomyces meyeri		92.48	0.55	MLZ 1	99	
Lactobacillus fermentum		7.16	0.46		99	
Fusobacterium mortiferum		0.35	0.23		1	
03015012	**Low discrimination**					**03015012**
Lactobacillus fermentum		71.47	0.46	MLZ 1		
Actinomyces meyeri		28.54	0.29			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03015020	**Low discrimination**					**03015020**
Actinomyces meyeri		83.95	0.74	NAG 12		
Lactobacillus fermentum		15.90	0.72			
Atopobium minutum		0.09	0.27			
03015021	**Low discrimination**					**03015021**
Actinomyces meyeri		84.08	0.40	NAG 12		
Lactobacillus fermentum		15.92	0.38	URE 1		
03015022	**Low discrimination**					**03015022**
Lactobacillus fermentum		85.68	0.72			
Actinomyces meyeri		14.00	0.48			
Atopobium minutum		0.33	0.24			
03015023	**Low discrimination**					**03015023**
Lactobacillus fermentum		85.96	0.38	URE 1		
Actinomyces meyeri		14.04	0.14			
03015030	**Low discrimination**					**03015030**
Actinomyces meyeri		84.08	0.40	MLZ 1		
Lactobacillus fermentum		15.92	0.38	NAG 12		
03015032	**Low discrimination**					**03015032**
Lactobacillus fermentum		85.96	0.38	MLZ 1		
Actinomyces meyeri		14.04	0.14			
03015040	**Low discrimination**					**03015040**
					SPO	
Actinomyces meyeri		50.08	0.55	SOR 1	1	
Clostridium glycolicum		44.90	0.49		99	
Lactobacillus fermentum		3.88	0.46		1	
03015042	**Low discrimination**					**03015042**
					SPO	
Lactobacillus fermentum		64.79	0.46	SOR 1	1	
Actinomyces meyeri		25.87	0.29		1	
Clostridium glycolicum		7.58	0.15		99	
03015060	**Low discrimination**					**03015060**
					SPO	
Actinomyces meyeri		79.63	0.40	NAG 12	1	
Lactobacillus fermentum		15.08	0.38	SOR 1	1	
Clostridium glycolicum		5.29	0.15		99	
03015062	**Low discrimination**					**03015062**
Lactobacillus fermentum		85.96	0.38	SOR 1		
Actinomyces meyeri		14.04	0.14			
03015100	**Very good identification**					**03015100**
					GRA	
Actinomyces meyeri		96.73	0.99		99	
Lactobacillus fermentum		3.06	0.84		99	
Fusobacterium mortiferum		0.22	0.63		1	
03015101	**Very good identification**					**03015101**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		96.66	0.65	URE 1	99	
Lactobacillus fermentum		3.05	0.50		99	
Fusobacterium mortiferum		0.22	0.29		1	
03015102	**Low discrimination**					**03015102**
					GRA	
Lactobacillus fermentum		50.34	0.84		99	
Actinomyces meyeri		49.27	0.73		99	
Fusobacterium mortiferum		0.39	0.47		1	
03015103	**Low discrimination**					**03015103**
					GRA	
Lactobacillus fermentum		50.34	0.50	URE 1	99	
Actinomyces meyeri		49.27	0.39		99	
Fusobacterium mortiferum		0.39	0.13		1	
03015110	**Very good identification**					**03015110**
					GRA	
Actinomyces meyeri		96.69	0.65	MLZ 1	99	
Lactobacillus fermentum		3.06	0.50		99	
Fusobacterium mortiferum		0.22	0.29		1	
03015112	**Low discrimination**					**03015112**
					GRA	
Lactobacillus fermentum		50.34	0.50	MLZ 1	99	
Actinomyces meyeri		49.27	0.39		99	
Fusobacterium mortiferum		0.39	0.13		1	
03015120	**Good identification**					**03015120**
					GRA	
Actinomyces meyeri		92.81	0.84	NAG 12	99	
Lactobacillus fermentum		7.17	0.76		99	
Fusobacterium mortiferum		0.02	0.29		1	
03015121	**Acceptable identification**					**03015121**
					SPO	
Actinomyces meyeri		92.67	0.50	NAG 12	1	
Lactobacillus fermentum		7.16	0.42	URE 1	1	
Clostridium chauvoei		0.17	0.01		99	
03015122	**Low discrimination**					**03015122**
					GRA	
Lactobacillus fermentum		71.39	0.76		99	
Actinomyces meyeri		28.59	0.59		99	
Fusobacterium mortiferum		0.02	0.13		1	
03015123	**Low discrimination**					**03015123**
Lactobacillus fermentum		71.41	0.42	URE 1		
Actinomyces meyeri		28.60	0.24			
03015130	**Acceptable identification**					**03015130**
Actinomyces meyeri		92.83	0.50	MLZ 1		
Lactobacillus fermentum		7.17	0.42	NAG 12		
03015132	**Low discrimination**					**03015132**
Lactobacillus fermentum		71.41	0.42	MLZ 1		
Actinomyces meyeri		28.60	0.24			
03015140	**Very good identification**					**03015140**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		96.52	0.65	SOR 1	1	
Lactobacillus fermentum		3.05	0.50		1	
Clostridium glycolicum		0.22	0.15		99	
03015142	**Low discrimination**					**03015142**
					GRA	
Lactobacillus fermentum		50.34	0.50	SOR 1	99	
Actinomyces meyeri		49.27	0.39		99	
Fusobacterium mortiferum		0.39	0.13		1	
03015160	**Acceptable identification**					**03015160**
Actinomyces meyeri		92.83	0.50	NAG 12		
Lactobacillus fermentum		7.17	0.42	SOR 1		
03015162	**Low discrimination**					**03015162**
Lactobacillus fermentum		71.41	0.42	SOR 1		
Actinomyces meyeri		28.60	0.24			
03015200	**Acceptable identification**					**03015200**
Actinomyces meyeri		92.08	0.55	RHA 1		
Lactobacillus fermentum		7.13	0.46			
Actinomyces odontolyticus		0.43	0.19			
03015202	**Low discrimination**					**03015202**
Lactobacillus fermentum		69.61	0.46	RHA 1		
Actinomyces meyeri		27.79	0.29			
Actinomyces odontolyticus		2.60	0.15			
03015220	**Low discrimination**					**03015220**
Actinomyces meyeri		84.08	0.40	RHA 1		
Lactobacillus fermentum		15.92	0.38	NAG 12		
03015222	**Low discrimination**					**03015222**
Lactobacillus fermentum		85.96	0.38	RHA 1		
Actinomyces meyeri		14.04	0.14			
03015300	**Very good identification**					**03015300**
					GRA	
Actinomyces meyeri		96.69	0.65	RHA 1	99	
Lactobacillus fermentum		3.06	0.50		99	
Fusobacterium mortiferum		0.22	0.29		1	
03015302	**Low discrimination**					**03015302**
					GRA	
Lactobacillus fermentum		50.15	0.50	RHA 1	99	
Actinomyces meyeri		49.09	0.39		99	
Fusobacterium mortiferum		0.39	0.13		1	
03015320	**Acceptable identification**					**03015320**
Actinomyces meyeri		92.83	0.50	RHA 1		
Lactobacillus fermentum		7.17	0.42	NAG 12		
03015322	**Low discrimination**					**03015322**
					GRA	
Lactobacillus fermentum		70.47	0.42	RHA 1	99	
Actinomyces meyeri		28.22	0.24		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		1.31	0.09		1	
03015400	**Low discrimination**					**03015400**
					COC	
Actinomyces meyeri		76.32	0.79		1	
Lactobacillus fermentum		23.65	0.81		1	
Gemella morbillorum		0.02	0.16		99	
03015401	**Low discrimination**					**03015401**
Actinomyces meyeri		76.35	0.44	URE 1		
Lactobacillus fermentum		23.65	0.46			
03015402	**Good identification**					**03015402**
Lactobacillus fermentum		90.92	0.81			
Actinomyces meyeri		9.08	0.53			
03015403	**Acceptable identification**					**03015403**
Lactobacillus fermentum		90.92	0.46	URE 1		
Actinomyces meyeri		9.08	0.19			
03015410	**Low discrimination**					**03015410**
Actinomyces meyeri		76.35	0.44	MLZ 1		
Lactobacillus fermentum		23.65	0.46			
03015412	**Acceptable identification**					**03015412**
Lactobacillus fermentum		90.92	0.46	MLZ 1		
Actinomyces meyeri		9.08	0.19			
03015420	**Low discrimination**					**03015420**
Actinomyces meyeri		56.90	0.64	NAG 12		
Lactobacillus fermentum		43.10	0.72			
03015422	**Very good identification**					**03015422**
Lactobacillus fermentum		96.08	0.72			
Actinomyces meyeri		3.92	0.38			
03015423	**Good identification**					**03015423**
Lactobacillus fermentum		96.08	0.38	URE 1		
Actinomyces meyeri		3.92	0.04			
03015432	**Good identification**					**03015432**
Lactobacillus fermentum		96.08	0.38	MLZ 1		
Actinomyces meyeri		3.92	0.04			
03015440	**Low discrimination**					**03015440**
					SPO	
Actinomyces meyeri		74.29	0.44	SOR 1	1	
Lactobacillus fermentum		23.02	0.46		1	
Clostridium glycolicum		2.69	0.15		99	
03015442	**Acceptable identification**					**03015442**
Lactobacillus fermentum		90.92	0.46	SOR 1		
Actinomyces meyeri		9.08	0.19			
03015462	**Good identification**					**03015462**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		96.08	0.38	SOR 1		
Actinomyces meyeri		3.92	0.04			
03015500	**Low discrimination**					**03015500**
					GRA	
Actinomyces meyeri		88.77	0.89		99	
Lactobacillus fermentum		11.22	0.84		99	
Fusobacterium mortiferum		0.01	0.29		1	
03015501	**Low discrimination**					**03015501**
Actinomyces meyeri		88.78	0.55	URE 1		
Lactobacillus fermentum		11.22	0.50			
03015502	**Low discrimination**					**03015502**
					GRA	
Lactobacillus fermentum		80.33	0.84		99	
Actinomyces meyeri		19.66	0.63		99	
Fusobacterium mortiferum		0.01	0.13		1	
03015503	**Low discrimination**					**03015503**
Lactobacillus fermentum		80.34	0.50	URE 1		
Actinomyces meyeri		19.66	0.29			
03015510	**Low discrimination**					**03015510**
Actinomyces meyeri		88.53	0.55	MLZ 1		
Lactobacillus fermentum		11.19	0.50			
Bifidobacterium longum subsp. longum		0.28	0.16			
03015512	**Low discrimination**					**03015512**
Lactobacillus fermentum		78.77	0.50	MLZ 1		
Actinomyces meyeri		19.28	0.29			
Bifidobacterium longum subsp. longum		1.96	0.16			
03015520	**Low discrimination**					**03015520**
Actinomyces meyeri		76.37	0.74	NAG 12		
Lactobacillus fermentum		23.59	0.76			
Bifidobacterium longum subsp. longum		0.04	0.22			
03015521	**Low discrimination**					**03015521**
Actinomyces meyeri		76.40	0.40	NAG 12		
Lactobacillus fermentum		23.60	0.42	URE 1		
03015522	**Good identification**					**03015522**
Lactobacillus fermentum		90.75	0.76			
Actinomyces meyeri		9.09	0.48			
Bifidobacterium longum subsp. longum		0.16	0.22			
03015523	**Acceptable identification**					**03015523**
Lactobacillus fermentum		90.90	0.42	URE 1		
Actinomyces meyeri		9.10	0.14			
03015530	**Low discrimination**					**03015530**
Actinomyces meyeri		71.37	0.40	MLZ 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		22.05	0.42	NAG 12		
Bifidobacterium longum subsp. longum		6.58	0.26			
03015532	**Low discrimination**					**03015532**
Lactobacillus fermentum		71.50	0.42	MLZ 1		
Bifidobacterium longum subsp. longum		21.35	0.26			
Actinomyces meyeri		7.16	0.14			
03015540	**Low discrimination**					**03015540**
Actinomyces meyeri		88.78	0.55	SOR 1		
Lactobacillus fermentum		11.22	0.50			
03015542	**Low discrimination**					**03015542**
Lactobacillus fermentum		80.34	0.50	SOR 1		
Actinomyces meyeri		19.66	0.29			
03015560	**Low discrimination**					**03015560**
Actinomyces meyeri		76.40	0.40	NAG 12		
Lactobacillus fermentum		23.60	0.42	SOR 1		
03015562	**Acceptable identification**					**03015562**
Lactobacillus fermentum		90.90	0.42	SOR 1		
Actinomyces meyeri		9.10	0.14			
03015600	**Low discrimination**					**03015600**
Actinomyces meyeri		76.35	0.44	RHA 1		
Lactobacillus fermentum		23.65	0.46			
03015602	**Acceptable identification**					**03015602**
Lactobacillus fermentum		90.92	0.46	RHA 1		
Actinomyces meyeri		9.08	0.19			
03015622	**Good identification**					**03015622**
Lactobacillus fermentum		96.08	0.38	RHA 1		
Actinomyces meyeri		3.92	0.04			
03015700	**Low discrimination**					**03015700**
Actinomyces meyeri		88.78	0.55	RHA 1		
Lactobacillus fermentum		11.22	0.50			
03015702	**Low discrimination**					**03015702**
Lactobacillus fermentum		80.34	0.50	RHA 1		
Actinomyces meyeri		19.66	0.29			
03015720	**Low discrimination**					**03015720**
					GRA	
Actinomyces meyeri		75.35	0.40	RHA 1	99	
Lactobacillus fermentum		23.28	0.42	NAG 12	99	
Bacteroides vulgatus		1.37	-.01		1	
03015722	**Acceptable identification**					**03015722**
Lactobacillus fermentum		90.90	0.42	RHA 1		
Actinomyces meyeri		9.10	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03016000	**Low discrimination**							**03016000**
					COC	GRA		
Megasphaera elsdenii		76.67	0.63	XYL 1	99	1		
Actinomyces meyeri		18.05	0.52		1	99		
Pseudoramibacter alactolyticus		2.64	0.27		1	99		
03016002	**Acceptable identification**							**03016002**
					COC	GRA		
Megasphaera elsdenii		90.22	0.47	XYL 1	99	1		
Actinomyces meyeri		5.91	0.26	BGL 10	1	99		
Pseudoramibacter alactolyticus		3.11	0.11		1	99		
03016020	**Low discrimination**							**03016020**
					COC	GRA		
Actinomyces meyeri		66.01	0.37	MAN 1	1	99		
Megasphaera elsdenii		20.77	0.29	NAG 12	99	1		
Pseudoramibacter alactolyticus		12.50	0.14		1	99		
03016040	**Good identification**							**03016040**
					COC	SPO	GRA	
Megasphaera elsdenii		94.32	0.60	XYL 1	99	1	1	
Clostridium glycolicum		4.07	0.31		1	99	99	
Pseudoramibacter alactolyticus		1.22	0.17		1	1	99	
03016042	**Good identification**							**03016042**
					COC	SPO	GRA	
Megasphaera elsdenii		98.34	0.44	XYL 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.00	BGL 10	1	1	99	
Clostridium glycolicum		0.39	-.03		1	99	99	
03016100	**Very good identification**							**03016100**
					COC	GRA		
Actinomyces meyeri		98.60	0.62	MAN 1	1	99		
Megasphaera elsdenii		1.06	0.29		99	1		
Fusobacterium mortiferum		0.22	0.26		1	1		
03016102	**Acceptable identification**							**03016102**
					COC	GRA		
Actinomyces meyeri		92.98	0.36	MAN 1	1	99		
Megasphaera elsdenii		3.58	0.12	BGL 3	99	1		
Propionibacterium propionicum		2.00	0.18		1	99		
03016120	**Good identification**							**03016120**
Actinomyces meyeri		100.0	0.47	MAN 1				
03016400	**Low discrimination**							**03016400**
					COC	GRA		
Actinomyces meyeri		84.59	0.41	MAN 1	1	99		
Megasphaera elsdenii		14.52	0.29		99	1		
Pseudoramibacter alactolyticus		0.50	-.07		1	99		
03016500	**Good identification**							**03016500**
Actinomyces meyeri		99.81	0.52	MAN 1				
Lactobacillus fermentum		0.19	0.16					
03016520	**Good identification**							**03016520**
Actinomyces meyeri		100.0	0.37	MAN 1				
03017000	**Low discrimination**							**03017000**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		89.70	0.55	MAN 1	1	99	
Lactobacillus fermentum		6.95	0.46		1	99	
Megasphaera elsdenii		2.57	0.29		99	1	
03017002	**Low discrimination**						**03017002**
					COC	GRA	
Lactobacillus fermentum		68.81	0.46	MAN 1	1	99	
Actinomyces meyeri		27.48	0.29		1	99	
Megasphaera elsdenii		2.82	0.12		99	1	
03017020	**Low discrimination**						**03017020**
					CAT		
Actinomyces meyeri		83.56	0.40	MAN 1	1		
Lactobacillus fermentum		15.82	0.38	NAG 12	1		
Propionibacterium granulosum		0.61	0.09		90		
03017022	**Low discrimination**						**03017022**
Lactobacillus fermentum		85.96	0.38	MAN 1			
Actinomyces meyeri		14.04	0.14				
03017100	**Very good identification**						**03017100**
					GRA		
Actinomyces meyeri		96.70	0.65	MAN 1	99		
Lactobacillus fermentum		3.06	0.50		99		
Fusobacterium mortiferum		0.22	0.29		1		
03017102	**Low discrimination**						**03017102**
Lactobacillus fermentum		49.98	0.50	MAN 1			
Actinomyces meyeri		48.93	0.39				
Propionibacterium propionicum		0.70	0.18				
03017120	**Acceptable identification**						**03017120**
Actinomyces meyeri		92.83	0.50	MAN 1			
Lactobacillus fermentum		7.17	0.42	NAG 12			
03017122	**Low discrimination**						**03017122**
Lactobacillus fermentum		71.41	0.42	MAN 1			
Actinomyces meyeri		28.60	0.24				
03017400	**Low discrimination**						**03017400**
Actinomyces meyeri		76.35	0.44	MAN 1			
Lactobacillus fermentum		23.65	0.46				
03017402	**Acceptable identification**						**03017402**
Lactobacillus fermentum		90.92	0.46	MAN 1			
Actinomyces meyeri		9.08	0.19				
03017422	**Good identification**						**03017422**
Lactobacillus fermentum		96.08	0.38	MAN 1			
Actinomyces meyeri		3.92	0.04				
03017500	**Low discrimination**						**03017500**
Actinomyces meyeri		88.78	0.55	MAN 1			
Lactobacillus fermentum		11.22	0.50				
03017502	**Low discrimination**						**03017502**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		80.34	0.50	MAN 1				
Actinomyces meyeri		19.66	0.29					
03017520	**Low discrimination**							**03017520**
Actinomyces meyeri		76.40	0.40	MAN 1				
Lactobacillus fermentum		23.60	0.42	NAG 12				
03017522	**Acceptable identification**							**03017522**
Lactobacillus fermentum		90.90	0.42	MAN 1				
Actinomyces meyeri		9.10	0.14					
03020000	**Low discrimination**							**03020000**
					COC	GRA		
Clostridium botulinum biovar A		89.78	0.66	SUC 7	1	99		
Clostridium sporogenes		3.87	0.40		1	99		
Megasphaera elsdenii		1.76	0.41		99	1		
03020040	**Good identification**							**03020040**
					COC	GRA		
Clostridium botulinum biovar A		94.03	0.57	SUC 7	1	99		
Megasphaera elsdenii		4.12	0.38		99	1		
Clostridium botulinum biovar B		1.16	0.21		1	99		
03021100	**Low discrimination**							**03021100**
					GRA			
Actinomyces meyeri		73.89	0.38	FRU 95	99			
Fusobacterium mortiferum		24.01	0.39	TRE 1	1			
Lactobacillus fermentum		1.48	0.20		99			
03022000	**Low discrimination**							**03022000**
					COC	SPO	CAT	
Megasphaera elsdenii		60.87	0.41	FRU 95	99	1	1	
Clostridium botulinum biovar A		31.29	0.32	TRE 1	1	99	1	
Propionibacterium granulosum		4.91	0.28		1	1	90	
03022040	**Low discrimination**							**03022040**
					COC	GRA		
Megasphaera elsdenii		81.28	0.38	FRU 95	99	1		
Clostridium botulinum biovar A		18.72	0.23	TRE 1	1	99		
03024100	**Good identification**							**03024100**
					GRA			
Actinomyces meyeri		98.76	0.40	FRU 95	99			
Fusobacterium mortiferum		1.24	0.17	TRE 1	1			
03025100	**Good identification**							**03025100**
					GRA			
Actinomyces meyeri		98.17	0.43	FRU 95	99			
Fusobacterium mortiferum		1.24	0.20	TRE 1	1			
Lactobacillus fermentum		0.60	0.16		99			
03030000	**Low discrimination**							**03030000**
					COC	GRA	CAT	
Megasphaera elsdenii		42.51	0.63	TRE 1	99	1	1	
Propionibacterium granulosum		23.70	0.64		1	99	90	
Fusobacterium mortiferum		14.96	0.64		1	1	5	
03030002	**Low discrimination**							**03030002**
					COC	GRA	CAT	
Propionibacterium granulosum		40.97	0.54	MNS 99	1	99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		32.66	0.47		99	1	1	
Fusobacterium mortiferum		11.49	0.48		1	1	5	
03030010	**Low discrimination**							**03030010**
					COC	SPO	CAT	
Propionibacterium granulosum		76.11	0.49	MNS 99	1	1	90	
Megasphaera elsdenii		10.11	0.29	MLZ 12	99	1	1	
Clostridium difficile		5.54	0.17		1	99	1	
03030012	**Acceptable identification**							**03030012**
					COC	GRA	CAT	
Propionibacterium granulosum		91.22	0.39	MNS 99	1	99	90	
Megasphaera elsdenii		5.39	0.12	MLZ 12	99	1	1	
Fusobacterium mortiferum		1.90	0.13		1	1	5	
03030020	**Low discrimination**							**03030020**
					SPO	CAT		
Propionibacterium granulosum		60.83	0.54	MNS 99	1	90		
Clostridium cadaveris		13.44	0.30		99	1		
Actinomyces meyeri		7.22	0.32		1	1		
03030022	**Low discrimination**							**03030022**
					CAT			
Propionibacterium granulosum		71.55	0.43	MNS 99	90			
Atopobium minutum		21.07	0.32		1			
Pseudoramibacter alactolyticus		2.58	0.03		1			
03030030	**Good identification**							**03030030**
					SPO	CAT		
Propionibacterium granulosum		98.39	0.39	MNS 99	1	90		
Clostridium cadaveris		1.61	-.04	MLZ 12	99	1		
03030040	**Low discrimination**							**03030040**
					COC	GRA		
Megasphaera elsdenii		84.83	0.60	TRE 1	99	1		
Clostridium botulinum biovar A		6.50	0.36		1	99		
Clostridium botulinum biovar B		5.01	0.32		1	99		
03030042	**Acceptable identification**							**03030042**
					COC	CAT	GRA	
Megasphaera elsdenii		93.91	0.44	TRE 1	99	1	1	
Pseudoramibacter alactolyticus		2.25	0.05	BGL 10	1	1	99	
Propionibacterium granulosum		1.79	0.20		1	90	99	
03030100	**Low discrimination**							**03030100**
					COC	GRA		
Fusobacterium mortiferum		61.93	0.70		1	1		
Actinomyces meyeri		36.59	0.57		1	99		
Megasphaera elsdenii		0.76	0.29		99	1		
03030101	**Low discrimination**							**03030101**
					GRA			
Fusobacterium mortiferum		62.86	0.36	URE 1	1			
Actinomyces meyeri		37.14	0.23		99			
03030102	**Low discrimination**							**03030102**
					GRA	CAT		
Fusobacterium mortiferum		82.67	0.54	BGL 10	1	5		
Actinomyces meyeri		13.60	0.31		99	1		
Propionibacterium granulosum		1.28	0.20		99	90		
03030110	**Low discrimination**							**03030110**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		59.41	0.36	MLZ 1	1	5		
Actinomyces meyeri		35.10	0.23		99	1		
Propionibacterium granulosum		5.50	0.15		99	90		
03030120	**Low discrimination**							**03030120**
					GRA	CAT		
Actinomyces meyeri		86.85	0.42	TRE 1	99	1		
Fusobacterium mortiferum		10.89	0.36	NAG 12	1	5		
Propionibacterium granulosum		1.85	0.20		99	90		
03030140	**Low discrimination**							**03030140**
					COC	GRA		
Fusobacterium mortiferum		39.16	0.36	SOR 1	1	1		
Megasphaera elsdenii		31.79	0.26		99	1		
Actinomyces meyeri		23.13	0.23		1	99		
03030200	**Low discrimination**							**03030200**
					COC	GRA	CAT	
Propionibacterium granulosum		49.32	0.38	MNS 99	1	99	90	
Megasphaera elsdenii		28.89	0.29	RHA 3	99	1	1	
Fusobacterium mortiferum		10.17	0.30		1	1	5	
03030300	**Low discrimination**							**03030300**
					GRA			
Fusobacterium mortiferum		62.86	0.36	RHA 1	1			
Actinomyces meyeri		37.14	0.23		99			
03030400	**Low discrimination**							**03030400**
					COC	CAT	GRA	
Actinomyces meyeri		56.91	0.36	TRE 1	1	1	99	
Megasphaera elsdenii		18.96	0.29		99	1	1	
Propionibacterium granulosum		10.57	0.30		1	90	99	
03030500	**Acceptable identification**							**03030500**
					GRA			
Actinomyces meyeri		93.05	0.47	TRE 1	99			
Fusobacterium mortiferum		6.36	0.36		1			
Lactobacillus fermentum		0.59	0.20		99			
03031000	**Low discrimination**							**03031000**
					GRA	CAT		
Propionibacterium granulosum		41.48	0.64	MNS 99	99	90		
Fusobacterium mortiferum		39.28	0.67		1	5		
Actinomyces meyeri		13.54	0.50		99	1		
03031002	**Low discrimination**							**03031002**
					GRA	CAT		
Propionibacterium granulosum		53.27	0.54	MNS 99	99	90		
Fusobacterium mortiferum		22.42	0.51		1	5		
Lactobacillus fermentum		17.80	0.50		99	1		
03031010	**Acceptable identification**							**03031010**
					GRA	CAT		
Propionibacterium granulosum		90.50	0.49	MNS 99	99	90		
Fusobacterium mortiferum		6.35	0.33	MLZ 12	1	5		
Actinomyces meyeri		2.19	0.16		99	1		
03031012	**Acceptable identification**							**03031012**
					GRA	CAT		
Propionibacterium granulosum		94.70	0.39	MNS 99	99	90		
Fusobacterium mortiferum		2.95	0.16	MLZ 12	1	5		
Lactobacillus fermentum		2.34	0.16		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03031020	**Low discrimination**						**03031020**
					CAT		
Propionibacterium granulosum		72.16	0.54	MNS 99	90		
Actinomyces meyeri		12.85	0.35		1		
Lactobacillus fermentum		8.04	0.42		1		
03031022	**Low discrimination**						**03031022**
					CAT		
Propionibacterium granulosum		61.78	0.43	MNS 99	90		
Lactobacillus fermentum		27.53	0.42		1		
Atopobium minutum		6.06	0.24		1		
03031030	**Good identification**						**03031030**
Propionibacterium granulosum		100.0	0.39	MNS 99			
03031100	**Low discrimination**						**03031100**
					GRA		
Fusobacterium mortiferum		60.23	0.73		1		
Actinomyces meyeri		35.58	0.60		99		
Lactobacillus fermentum		3.72	0.54		99		
03031101	**Low discrimination**						**03031101**
					GRA	SPO	
Fusobacterium mortiferum		37.63	0.39	URE 1	1	1	
Clostridium chauvoei		35.43	0.27		99	99	
Actinomyces meyeri		22.23	0.26		99	1	
03031102	**Low discrimination**						**03031102**
					GRA		
Fusobacterium mortiferum		57.48	0.57	BGL 10	1		
Lactobacillus fermentum		31.92	0.54		99		
Actinomyces meyeri		9.45	0.34		99		
03031110	**Low discrimination**						**03031110**
					GRA		
Fusobacterium mortiferum		56.26	0.39	MLZ 1	1		
Actinomyces meyeri		33.24	0.26		99		
Eubacterium saburreum		3.56	0.17		85		
03031120	**Low discrimination**						**03031120**
Actinomyces meyeri		62.39	0.45	TRE 1			
Lactobacillus fermentum		15.93	0.46	NAG 12			
Clostridium perfringens		12.27	0.32				
03031122	**Low discrimination**						**03031122**
Lactobacillus fermentum		63.58	0.46	TRE 1			
Clostridium perfringens		23.05	0.26				
Actinomyces meyeri		7.70	0.19				
03031140	**Low discrimination**						**03031140**
					GRA		
Fusobacterium mortiferum		59.53	0.39	SOR 1	1		
Actinomyces meyeri		35.17	0.26		99		
Lactobacillus fermentum		3.67	0.20		99		
03031200	**Low discrimination**						**03031200**
					GRA	CAT	
Propionibacterium granulosum		69.29	0.38	MNS 99	99	90	
Fusobacterium mortiferum		21.43	0.33	RHA 3	1	5	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		7.39	0.16		99	1		
03031300	**Low discrimination**							**03031300**
					GRA			
Fusobacterium mortiferum		60.52	0.39	RHA 1	1			
Actinomyces meyeri		35.75	0.26		99			
Lactobacillus fermentum		3.73	0.20		99			
03031400	**Low discrimination**							**03031400**
					CAT			
Lactobacillus fermentum		45.21	0.50	TRE 1	1			
Actinomyces meyeri		44.15	0.39		1			
Propionibacterium granulosum		5.47	0.30		90			
03031402	**Acceptable identification**							**03031402**
					CAT			
Lactobacillus fermentum		93.18	0.50	TRE 1	1			
Propionibacterium granulosum		2.82	0.20		90			
Actinomyces meyeri		2.81	0.14		1			
03031420	**Low discrimination**							**03031420**
					CAT			
Lactobacillus fermentum		67.11	0.42	TRE 1	1			
Actinomyces meyeri		26.81	0.25		1			
Propionibacterium granulosum		6.09	0.20		90			
03031422	**Good identification**							**03031422**
					CAT			
Lactobacillus fermentum		97.78	0.42	TRE 1	1			
Propionibacterium granulosum		2.22	0.09		90			
03031500	**Low discrimination**							**03031500**
					GRA			
Actinomyces meyeri		67.29	0.50	TRE 1	99			
Lactobacillus fermentum		28.11	0.54		99			
Fusobacterium mortiferum		4.60	0.39		1			
03031502	**Acceptable identification**							**03031502**
					GRA			
Lactobacillus fermentum		91.56	0.54	TRE 1	99			
Actinomyces meyeri		6.78	0.24		99			
Fusobacterium mortiferum		1.67	0.23		1			
03031520	**Low discrimination**							**03031520**
Lactobacillus fermentum		50.52	0.46	TRE 1				
Actinomyces meyeri		49.48	0.35					
03031522	**Good identification**							**03031522**
Lactobacillus fermentum		97.06	0.46	TRE 1				
Actinomyces meyeri		2.94	0.09					
03032000	**Low discrimination**							**03032000**
					COC	CAT	GRA	
Megasphaera elsdenii		66.91	0.63	TRE 1	99	1	1	
Propionibacterium granulosum		28.14	0.62		1	90	99	
Pseudoramibacter alactolyticus		2.31	0.27		1	1	99	
03032002	**Low discrimination**							**03032002**
					COC	CAT	GRA	
Megasphaera elsdenii		50.21	0.47	TRE 1	99	1	1	
Propionibacterium granulosum		47.51	0.52	BGL 10	1	90	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		1.73	0.11		1	1	99	
03032010	**Low discrimination**							**03032010**
					COC	SPO	CAT	
Propionibacterium granulosum		48.79	0.47	MNS 99	1	1	90	
Clostridium difficile		42.33	0.34	MLZ 12	1	99	1	
Megasphaera elsdenii		8.59	0.29		99	1	1	
03032012	**Low discrimination**							**03032012**
					COC	SPO	CAT	
Propionibacterium granulosum		89.82	0.37	MNS 99	1	1	90	
Megasphaera elsdenii		7.03	0.12	MLZ 12	99	1	1	
Clostridium difficile		3.15	-.01		1	99	1	
03032020	**Low discrimination**							**03032020**
					COC	CAT	GRA	
Propionibacterium granulosum		86.44	0.52	MNS 99	1	90	99	
Megasphaera elsdenii		8.30	0.29		99	1	1	
Pseudoramibacter alactolyticus		5.00	0.14		1	1	99	
03032022	**Acceptable identification**							**03032022**
					COC	CAT	GRA	
Propionibacterium granulosum		93.60	0.41	MNS 99	1	90	99	
Megasphaera elsdenii		4.00	0.12		99	1	1	
Pseudoramibacter alactolyticus		2.41	-.02		1	1	99	
03032030	**Good identification**							**03032030**
					SPO	CAT		
Propionibacterium granulosum		96.61	0.37	MNS 99	1	90		
Clostridium difficile		3.39	-.01	MLZ 12	99	1		
03032040	**Very good identification**							**03032040**
					COC	CAT	GRA	
Megasphaera elsdenii		97.85	0.60	TRE 1	99	1	1	
Pseudoramibacter alactolyticus		1.26	0.17		1	1	99	
Propionibacterium granulosum		0.62	0.28		1	90	99	
03032042	**Good identification**							**03032042**
					COC	CAT	GRA	
Megasphaera elsdenii		96.70	0.44	TRE 1	99	1	1	
Propionibacterium granulosum		1.39	0.18	BGL 10	1	90	99	
Pseudoramibacter alactolyticus		1.25	0.00		1	1	99	
03032100	**Low discrimination**							**03032100**
					COC	GRA		
Propionibacterium propionicum		54.81	0.48	NIT 86	1	99		
Megasphaera elsdenii		16.36	0.29	MLT 99	99	1		
Fusobacterium mortiferum		13.44	0.36		1	1		
03032102	**Acceptable identification**							**03032102**
					COC	CAT	GRA	
Propionibacterium propionicum		94.23	0.51	NIT 86	1	1	99	
Megasphaera elsdenii		2.08	0.12	MLT 99	99	1	1	
Propionibacterium granulosum		1.97	0.18		1	90	99	
03032140	**Low discrimination**							**03032140**
					COC	GRA		
Propionibacterium propionicum		83.40	0.48	NIT 86	1	99		
Megasphaera elsdenii		16.60	0.26	MLT 99	99	1		
03032142	**Good identification**							**03032142**
					COC	GRA		
Propionibacterium propionicum		98.55	0.51	NIT 86	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		1.45	0.09	MLT 99	99	1		
03032200	**Low discrimination**							**03032200**
					COC	CAT	GRA	
Propionibacterium granulosum		55.46	0.36	MNS 99	1	90	99	
Megasphaera elsdenii		43.06	0.29	RHA 3	99	1	1	
Pseudoramibacter alactolyticus		1.48	-.07		1	1	99	
03033000	**Very good identification**							**03033000**
					COC	GRA	CAT	
Propionibacterium granulosum		95.72	0.62	MNS 99	1	99	90	
Megasphaera elsdenii		2.30	0.29		99	1	1	
Fusobacterium mortiferum		1.21	0.33		1	1	5	
03033002	**Good identification**							**03033002**
					COC	GRA	CAT	
Propionibacterium granulosum		97.18	0.52	MNS 99	1	99	90	
Megasphaera elsdenii		1.04	0.12		99	1	1	
Fusobacterium mortiferum		0.55	0.16		1	1	5	
03033010	**Good identification**							**03033010**
					SPO	CAT		
Propionibacterium granulosum		99.13	0.47	MNS 99	1	90		
Clostridium difficile		0.87	-.01	MLZ 12	99	1		
03033012	**Good identification**							**03033012**
Propionibacterium granulosum		100.0	0.37	MNS 99				
03033020	**Good identification**							**03033020**
Propionibacterium granulosum		99.66	0.52	MNS 99				
Actinomyces naeslundii		0.34	0.24					
03033022	**Good identification**							**03033022**
Propionibacterium granulosum		98.65	0.41	MNS 99				
Actinomyces naeslundii		1.35	0.24					
03033030	**Good identification**							**03033030**
Propionibacterium granulosum		100.0	0.37	MNS 99				
03033100	**Low discrimination**							**03033100**
					GRA			
Propionibacterium propionicum		56.70	0.48	NIT 86	99			
Fusobacterium mortiferum		20.86	0.39	MLT 99	1			
Actinomyces meyeri		12.32	0.26		99			
03033102	**Acceptable identification**							**03033102**
					GRA	CAT		
Propionibacterium propionicum		92.30	0.51	NIT 86	99	1		
Fusobacterium mortiferum		2.52	0.23	MLT 99	1	5		
Propionibacterium granulosum		1.93	0.18		99	90		
03033121	**Good identification**							**03033121**
Actinomyces naeslundii		100.0	0.35	MNS 88				
03033123	**Good identification**							**03033123**
Actinomyces naeslundii		100.0	0.35	MNS 88				
03033140	**Good identification**							**03033140**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		100.0	0.48	NIT 86			
03033142	**Good identification**						**03033142**
Propionibacterium propionicum		100.0	0.51	NIT 86			
03033200	**Good identification**						**03033200**
Propionibacterium granulosum		100.0	0.36	MNS 99			
03034000	**Low discrimination**						**03034000**
					COC	GRA	
Actinomyces meyeri		83.19	0.52	TRE 1	1	99	
Fusobacterium mortiferum		9.36	0.45		1	1	
Megasphaera elsdenii		3.57	0.29		99	1	
03034020	**Acceptable identification**						**03034020**
					SPO	CAT	
Actinomyces meyeri		94.92	0.37	TRE 1	1	1	
Propionibacterium granulosum		4.16	0.20	NAG 12	1	90	
Clostridium cadaveris		0.92	-.04		99	1	
03034100	**Good identification**						**03034100**
					GRA		
Actinomyces meyeri		93.80	0.62	TRE 1	99		
Fusobacterium mortiferum		6.16	0.51		1		
Lactobacillus fermentum		0.05	0.16		99		
03034102	**Low discrimination**						**03034102**
					GRA		
Actinomyces meyeri		79.92	0.36	TRE 1	99		
Fusobacterium mortiferum		18.85	0.35	BGL 3	1		
Lactobacillus fermentum		1.24	0.16		99		
03034120	**Good identification**						**03034120**
					GRA		
Actinomyces meyeri		99.52	0.47	TRE 1	99		
Fusobacterium mortiferum		0.48	0.17	NAG 12	1		
03034400	**Good identification**						**03034400**
Actinomyces meyeri		99.53	0.41	TRE 1			
Lactobacillus fermentum		0.47	0.12				
03034500	**Good identification**						**03034500**
					GRA		
Actinomyces meyeri		99.55	0.52	TRE 1	99		
Fusobacterium mortiferum		0.26	0.17		1		
Lactobacillus fermentum		0.19	0.16		99		
03034520	**Good identification**						**03034520**
Actinomyces meyeri		100.0	0.37	TRE 1			
03035000	**Low discrimination**						**03035000**
					GRA		
Actinomyces meyeri		82.92	0.55	TRE 1	99		
Fusobacterium mortiferum		9.33	0.48		1		
Lactobacillus fermentum		6.42	0.46		99		
03035002	**Low discrimination**						**03035002**
					GRA		
Lactobacillus fermentum		62.03	0.46	TRE 1	99		
Actinomyces meyeri		24.77	0.29		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		10.01	0.31		1	
03035020	**Low discrimination**					**03035020**
					CAT	
Actinomyces meyeri		81.51	0.40	TRE 1	1	
Lactobacillus fermentum		15.43	0.38	NAG 12	1	
Propionibacterium granulosum		2.38	0.20		90	
03035022	**Low discrimination**					**03035022**
					CAT	
Lactobacillus fermentum		83.20	0.38	TRE 1	1	
Actinomyces meyeri		13.59	0.14		1	
Propionibacterium granulosum		3.21	0.09		90	
03035100	**Good identification**					**03035100**
					GRA	
Actinomyces meyeri		90.95	0.65	TRE 1	99	
Fusobacterium mortiferum		5.97	0.54		1	
Lactobacillus fermentum		2.87	0.50		99	
03035102	**Low discrimination**					**03035102**
					GRA	
Lactobacillus fermentum		45.04	0.50	TRE 1	99	
Actinomyces meyeri		44.09	0.39		99	
Fusobacterium mortiferum		10.40	0.38		1	
03035120	**Acceptable identification**					**03035120**
					GRA	
Actinomyces meyeri		92.20	0.50	TRE 1	99	
Lactobacillus fermentum		7.12	0.42	NAG 12	99	
Fusobacterium mortiferum		0.45	0.20		1	
03035122	**Low discrimination**					**03035122**
					GRA	
Lactobacillus fermentum		70.00	0.42	TRE 1	99	
Actinomyces meyeri		28.03	0.24		99	
Parabacteroides distasonis		1.97	0.12		1	
03035400	**Low discrimination**					**03035400**
					GRA	
Actinomyces meyeri		76.08	0.44	TRE 1	99	
Lactobacillus fermentum		23.57	0.46		99	
Fusobacterium mortiferum		0.35	0.14		1	
03035402	**Acceptable identification**					**03035402**
Lactobacillus fermentum		90.92	0.46	TRE 1		
Actinomyces meyeri		9.08	0.19			
03035422	**Good identification**					**03035422**
Lactobacillus fermentum		96.08	0.38	TRE 1		
Actinomyces meyeri		3.92	0.04			
03035500	**Low discrimination**					**03035500**
					GRA	
Actinomyces meyeri		88.57	0.55	TRE 1	99	
Lactobacillus fermentum		11.19	0.50		99	
Fusobacterium mortiferum		0.24	0.20		1	
03035502	**Low discrimination**					**03035502**
Lactobacillus fermentum		80.34	0.50	TRE 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Actinomyces meyeri	19.66	0.29					
03035520	**Low discrimination**							**03035520**
	Actinomyces meyeri	75.62	0.40	TRE 1				
	Lactobacillus fermentum	23.36	0.42	NAG 12				
	Bifidobacterium longum subsp. longum	1.02	0.12					
03035522	**Low discrimination**							**03035522**
	Lactobacillus fermentum	87.42	0.42	TRE 1				
	Actinomyces meyeri	8.75	0.14					
	Bifidobacterium longum subsp. longum	3.83	0.12					
03040000	**Low discrimination**							**03040000**
					COC	GRA	SPO	
	Clostridium botulinum biovar A	36.61	0.42	SUC 7	1	99	99	
	Megasphaera elsdenii	17.80	0.41	CEL 1	99	1	1	
	Peptostreptococcus anaerobius	17.06	0.32		99	99	1	
03040002	**Low discrimination**							**03040002**
					COC	SPO	GRA	
	Atopobium minutum	76.88	0.41	SUC 5	1	1	99	
	Megasphaera elsdenii	13.48	0.25	CEL 1	99	1	1	
	Clostridium sporogenes	2.69	0.06		1	99	99	
03040020	**Low discrimination**							**03040020**
					SPO			
	Atopobium minutum	61.17	0.36	SUC 5	1			
	Clostridium cadaveris	21.67	0.18	CEL 1	99			
	Clostridium novyi biovar A	5.06	0.21		99			
03040040	**Low discrimination**							**03040040**
					COC	GRA		
	Megasphaera elsdenii	48.66	0.38	FRU 95	99	1		
	Clostridium botulinum biovar A	44.84	0.33	CEL 1	1	99		
	Clostridium novyi biovar A	5.10	0.29		1	99		
03040100	**Low discrimination**							**03040100**
					GRA	SPO		
	Actinomyces meyeri	68.22	0.35	FRU 95	99	1		
	Fusobacterium mortiferum	24.73	0.37	CEL 1	1	1		
	Clostridium botulinum biovar A	2.92	0.08		99	99		
03041000	**Low discrimination**							**03041000**
					GRA			
	Atopobium minutum	40.00	0.36	SUC 5	99			
	Actinomyces meyeri	22.97	0.28	CEL 1	99			
	Fusobacterium mortiferum	14.28	0.34		1			
03041100	**Low discrimination**							**03041100**
					GRA			
	Actinomyces meyeri	72.33	0.38	FRU 95	99			
	Fusobacterium mortiferum	26.22	0.40	CEL 1	1			
	Lactobacillus fermentum	1.45	0.20		99			
03042000	**Good identification**							**03042000**
					COC	SPO	GRA	
	Megasphaera elsdenii	95.54	0.41	FRU 95	99	1	1	
	Clostridium botulinum biovar A	1.99	0.08	CEL 1	1	99	99	
	Atopobium minutum	0.92	0.10		1	1	99	
03042040	**Good identification**							**03042040**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Megasphaera elsdenii		99.08	0.38	FRU 95	99	1	
Clostridium botulinum biovar A		0.92	-.01	CEL 1	1	99	
03044100	**Good identification**						**03044100**
					GRA		
Actinomyces meyeri		98.61	0.40	FRU 95	99		
Fusobacterium mortiferum		1.39	0.18	CEL 1	1		
03045100	**Good identification**						**03045100**
					GRA		
Actinomyces meyeri		98.03	0.43	FRU 95	99		
Fusobacterium mortiferum		1.38	0.21	CEL 1	1		
Lactobacillus fermentum		0.59	0.16		99		
03050000	**Low discrimination**						**03050000**
					COC	GRA	
Megasphaera elsdenii		60.52	0.63	CEL 1	99	1	
Fusobacterium mortiferum		23.77	0.65		1	1	
Actinomyces meyeri		7.34	0.47		1	99	
03050002	**Low discrimination**						**03050002**
					COC	GRA	
Megasphaera elsdenii		56.30	0.47	CEL 1	99	1	
Fusobacterium mortiferum		22.12	0.48	BGL 10	1	1	
Atopobium minutum		15.60	0.40		1	99	
03050040	**Very good identification**						**03050040**
					COC	GRA	
Megasphaera elsdenii		96.30	0.60	CEL 1	99	1	
Pseudoramibacter alactolyticus		2.31	0.21		1	99	
Fusobacterium mortiferum		0.57	0.31		1	1	
03050042	**Good identification**						**03050042**
					COC	GRA	
Megasphaera elsdenii		96.70	0.44	CEL 1	99	1	
Pseudoramibacter alactolyticus		2.32	0.05	BGL 10	1	99	
Fusobacterium mortiferum		0.58	0.14		1	1	
03050100	**Low discrimination**						**03050100**
					COC	GRA	
Fusobacterium mortiferum		64.83	0.71		1	1	
Actinomyces meyeri		34.32	0.57		1	99	
Megasphaera elsdenii		0.72	0.29		99	1	
03050101	**Low discrimination**						**03050101**
					GRA		
Fusobacterium mortiferum		65.39	0.37	URE 1	1		
Actinomyces meyeri		34.61	0.23		99		
03050102	**Low discrimination**						**03050102**
					COC	GRA	
Fusobacterium mortiferum		85.55	0.55	BGL 10	1	1	
Actinomyces meyeri		12.61	0.31		1	99	
Megasphaera elsdenii		0.94	0.12		99	1	
03050110	**Low discrimination**						**03050110**
					GRA		
Fusobacterium mortiferum		65.39	0.37	MLZ 1	1		
Actinomyces meyeri		34.61	0.23		99		
03050120	**Low discrimination**						**03050120**
					GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		87.73	0.42	CEL 1	99		
Fusobacterium mortiferum		12.28	0.37	NAG 12	1		
03050140	**Low discrimination**						**03050140**
					COC	GRA	
Fusobacterium mortiferum		44.31	0.37	SOR 1	1	1	
Megasphaera elsdenii		32.23	0.26		99	1	
Actinomyces meyeri		23.46	0.23		1	99	
03050300	**Low discrimination**						**03050300**
					GRA		
Fusobacterium mortiferum		65.39	0.37	RHA 1	1		
Actinomyces meyeri		34.61	0.23		99		
03050400	**Low discrimination**						**03050400**
					COC	GRA	
Actinomyces meyeri		65.45	0.36	CEL 1	1	99	
Megasphaera elsdenii		21.80	0.29		99	1	
Fusobacterium mortiferum		8.56	0.31		1	1	
03050500	**Acceptable identification**						**03050500**
					GRA		
Actinomyces meyeri		92.37	0.47	CEL 1	99		
Fusobacterium mortiferum		7.05	0.37		1		
Lactobacillus fermentum		0.59	0.20		99		
03051000	**Low discrimination**						**03051000**
					GRA		
Fusobacterium mortiferum		69.29	0.68		1		
Actinomyces meyeri		21.40	0.50		99		
Lactobacillus fermentum		5.48	0.50		99		
03051002	**Low discrimination**						**03051002**
					GRA		
Fusobacterium mortiferum		49.63	0.51	BGL 10	1		
Lactobacillus fermentum		35.32	0.50		99		
Atopobium minutum		7.78	0.32		99		
03051022	**Low discrimination**						**03051022**
Lactobacillus fermentum		73.72	0.42	CEL 1			
Atopobium minutum		16.24	0.24				
Actinomyces meyeri		3.64	0.09				
03051100	**Low discrimination**						**03051100**
					GRA		
Fusobacterium mortiferum		63.04	0.74		1		
Actinomyces meyeri		33.37	0.60		99		
Lactobacillus fermentum		3.49	0.54		99		
03051101	**Low discrimination**						**03051101**
					GRA	SPO	
Fusobacterium mortiferum		58.82	0.40	URE 1	1	1	
Actinomyces meyeri		31.14	0.26		99	1	
Clostridium chauvoei		4.51	0.09		99	99	
03051102	**Low discrimination**						**03051102**
					GRA		
Fusobacterium mortiferum		60.04	0.58	BGL 10	1		
Lactobacillus fermentum		29.87	0.54		99		
Actinomyces meyeri		8.85	0.34		99		
03051110	**Low discrimination**						**03051110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		63.10	0.40	MLZ 1	1	
Actinomyces meyeri		33.41	0.26		99	
Lactobacillus fermentum		3.49	0.20		99	
03051120	**Low discrimination**					**03051120**
					GRA	
Actinomyces meyeri		67.60	0.45	CEL 1	99	
Lactobacillus fermentum		17.26	0.46	NAG 12	99	
Fusobacterium mortiferum		9.46	0.40		1	
03051122	**Low discrimination**					**03051122**
					GRA	
Lactobacillus fermentum		75.29	0.46	CEL 1	99	
Actinomyces meyeri		9.12	0.19		99	
Fusobacterium mortiferum		4.59	0.24		1	
03051140	**Low discrimination**					**03051140**
					GRA	
Fusobacterium mortiferum		63.10	0.40	SOR 1	1	
Actinomyces meyeri		33.41	0.26		99	
Lactobacillus fermentum		3.49	0.20		99	
03051300	**Low discrimination**					**03051300**
					GRA	
Fusobacterium mortiferum		63.10	0.40	RHA 1	1	
Actinomyces meyeri		33.41	0.26		99	
Lactobacillus fermentum		3.49	0.20		99	
03051400	**Low discrimination**					**03051400**
					GRA	
Lactobacillus fermentum		47.52	0.50	CEL 1	99	
Actinomyces meyeri		46.41	0.39		99	
Fusobacterium mortiferum		6.07	0.34		1	
03051402	**Good identification**					**03051402**
					GRA	
Lactobacillus fermentum		95.75	0.50	CEL 1	99	
Actinomyces meyeri		2.89	0.14		99	
Fusobacterium mortiferum		1.36	0.17		1	
03051420	**Low discrimination**					**03051420**
Lactobacillus fermentum		71.46	0.42	CEL 1		
Actinomyces meyeri		28.55	0.25			
03051422	**Good identification**					**03051422**
Lactobacillus fermentum		100.0	0.42	CEL 1		
03051500	**Low discrimination**					**03051500**
					GRA	
Actinomyces meyeri		66.93	0.50	CEL 1	99	
Lactobacillus fermentum		27.96	0.54		99	
Fusobacterium mortiferum		5.11	0.40		1	
03051502	**Acceptable identification**					**03051502**
					GRA	
Lactobacillus fermentum		91.38	0.54	CEL 1	99	
Actinomyces meyeri		6.77	0.24		99	
Fusobacterium mortiferum		1.86	0.24		1	
03051520	**Low discrimination**					**03051520**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		50.52	0.46	CEL 1				
Actinomyces meyeri		49.48	0.35					
03051522	**Good identification**							**03051522**
Lactobacillus fermentum		97.06	0.46	CEL 1				
Actinomyces meyeri		2.94	0.09					
03052000	**Good identification**							**03052000**
					COC	GRA		
Megasphaera elsdenii		93.30	0.63	CEL 1	99	1		
Pseudoramibacter alactolyticus		3.21	0.27		1	99		
Eubacterium limosum		2.04	0.26		1	99		
03052002	**Acceptable identification**							**03052002**
					COC	SPO	GRA	
Megasphaera elsdenii		94.68	0.47	CEL 1	99	1	1	
Pseudoramibacter alactolyticus		3.26	0.11	BGL 10	1	1	99	
Clostridium innocuum		1.12	0.10		1	99	99	
03052040	**Very good identification**							**03052040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.67	0.60	CEL 1	99	1	1	
Pseudoramibacter alactolyticus		1.28	0.17		1	1	99	
Clostridium difficile		0.06	-.03		1	99	99	
03052042	**Good identification**							**03052042**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	CEL 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
03053100	**Low discrimination**							**03053100**
					GRA			
Fusobacterium mortiferum		60.06	0.40	MAN 1	1			
Actinomyces meyeri		31.80	0.26		99			
Lactobacillus fermentum		3.32	0.20		99			
03054000	**Low discrimination**							**03054000**
					COC	GRA		
Actinomyces meyeri		84.47	0.52	CEL 1	1	99		
Fusobacterium mortiferum		10.61	0.46		1	1		
Megasphaera elsdenii		3.62	0.29		99	1		
03054020	**Good identification**							**03054020**
					GRA			
Actinomyces meyeri		99.08	0.37	CEL 1	99			
Fusobacterium mortiferum		0.92	0.11	NAG 12	1			
03054100	**Good identification**							**03054100**
					GRA			
Actinomyces meyeri		93.13	0.62	CEL 1	99			
Fusobacterium mortiferum		6.82	0.52		1			
Lactobacillus fermentum		0.05	0.16		99			
03054102	**Low discrimination**							**03054102**
					GRA			
Actinomyces meyeri		78.21	0.36	CEL 1	99			
Fusobacterium mortiferum		20.58	0.35	BGL 3	1			
Lactobacillus fermentum		1.21	0.16		99			
03054120	**Good identification**							**03054120**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		99.46	0.47	CEL 1	99	
Fusobacterium mortiferum		0.54	0.18	NAG 12	1	
03054400	**Good identification**					**03054400**
					GRA	
Actinomyces meyeri		99.03	0.41	CEL 1	99	
Fusobacterium mortiferum		0.50	0.11		1	
Lactobacillus fermentum		0.47	0.12		99	
03054500	**Good identification**					**03054500**
					GRA	
Actinomyces meyeri		99.52	0.52	CEL 1	99	
Fusobacterium mortiferum		0.29	0.18		1	
Lactobacillus fermentum		0.19	0.16		99	
03054520	**Good identification**					**03054520**
Actinomyces meyeri		100.0	0.37	CEL 1		
03055000	**Low discrimination**					**03055000**
					GRA	
Actinomyces meyeri		83.12	0.55	CEL 1	99	
Fusobacterium mortiferum		10.44	0.49		1	
Lactobacillus fermentum		6.44	0.46		99	
03055002	**Low discrimination**					**03055002**
					GRA	
Lactobacillus fermentum		63.32	0.46	CEL 1	99	
Actinomyces meyeri		25.28	0.29		99	
Fusobacterium mortiferum		11.40	0.32		1	
03055020	**Low discrimination**					**03055020**
					GRA	
Actinomyces meyeri		83.43	0.40	CEL 1	99	
Lactobacillus fermentum		15.80	0.38	NAG 12	99	
Fusobacterium mortiferum		0.78	0.14		1	
03055022	**Low discrimination**					**03055022**
Lactobacillus fermentum		85.96	0.38	CEL 1		
Actinomyces meyeri		14.04	0.14			
03055100	**Good identification**					**03055100**
					GRA	
Actinomyces meyeri		90.51	0.65	CEL 1	99	
Fusobacterium mortiferum		6.63	0.55		1	
Lactobacillus fermentum		2.86	0.50		99	
03055102	**Low discrimination**					**03055102**
					GRA	
Lactobacillus fermentum		44.71	0.50	CEL 1	99	
Actinomyces meyeri		43.77	0.39		99	
Fusobacterium mortiferum		11.52	0.39		1	
03055120	**Acceptable identification**					**03055120**
					GRA	
Actinomyces meyeri		92.37	0.50	CEL 1	99	
Lactobacillus fermentum		7.13	0.42	NAG 12	99	
Fusobacterium mortiferum		0.50	0.21		1	
03055122	**Low discrimination**					**03055122**
Lactobacillus fermentum		71.41	0.42	CEL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Actinomyces meyeri	28.60	0.24					
03055400	**Low discrimination**							**03055400**
					GRA			
	Actinomyces meyeri	76.05	0.44	CEL 1	99			
	Lactobacillus fermentum	23.56	0.46		99			
	Fusobacterium mortiferum	0.39	0.14		1			
03055402	**Acceptable identification**							**03055402**
	Lactobacillus fermentum	90.92	0.46	CEL 1				
	Actinomyces meyeri	9.08	0.19					
03055422	**Good identification**							**03055422**
	Lactobacillus fermentum	96.08	0.38	CEL 1				
	Actinomyces meyeri	3.92	0.04					
03055500	**Low discrimination**							**03055500**
					GRA			
	Actinomyces meyeri	88.55	0.55	CEL 1	99			
	Lactobacillus fermentum	11.19	0.50		99			
	Fusobacterium mortiferum	0.26	0.21		1			
03055502	**Low discrimination**							**03055502**
	Lactobacillus fermentum	79.82	0.50	CEL 1				
	Actinomyces meyeri	19.53	0.29					
	Bifidobacterium adolescentis	0.65	0.09					
03055520	**Low discrimination**							**03055520**
	Actinomyces meyeri	76.40	0.40	CEL 1				
	Lactobacillus fermentum	23.60	0.42	NAG 12				
03055522	**Acceptable identification**							**03055522**
	Lactobacillus fermentum	90.90	0.42	CEL 1				
	Actinomyces meyeri	9.10	0.14					
03070000	**Low discrimination**							**03070000**
					COC	GRA	CAT	
	Fusobacterium mortiferum	85.56	0.56		1	1	5	
	Megasphaera elsdenii	7.37	0.29		99	1	1	
	Propionibacterium granulosum	4.11	0.30		1	99	90	
03070002	**Low discrimination**							**03070002**
					COC	GRA	CAT	
	Fusobacterium mortiferum	82.11	0.39	BGL 10	1	1	5	
	Propionibacterium granulosum	8.87	0.20		1	99	90	
	Megasphaera elsdenii	7.07	0.12		99	1	1	
03070100	**Very good identification**							**03070100**
					GRA			
	Fusobacterium mortiferum	98.24	0.62		1			
	Actinomyces meyeri	1.76	0.23		99			
03070102	**Good identification**							**03070102**
	Fusobacterium mortiferum	100.0	0.46	BGL 10				
03071000	**Good identification**							**03071000**
					GRA	CAT		
	Fusobacterium mortiferum	93.23	0.59		1	5		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		2.98	0.30		99	90		
Lactobacillus jensenii		2.56	0.22		99	1		
03071002	**Low discrimination**							**03071002**
					GRA	CAT		
Fusobacterium mortiferum		79.32	0.42	BGL 10	1	5		
Lactobacillus jensenii		13.06	0.19		99	1		
Propionibacterium granulosum		5.71	0.20		99	90		
03071100	**Very good identification**							**03071100**
					COC	GRA		
Fusobacterium mortiferum		97.11	0.65		1	1		
Actinomyces meyeri		1.74	0.26		1	99		
Atopobium parvulum		0.80	0.18		99	99		
03071102	**Acceptable identification**							**03071102**
					GRA			
Fusobacterium mortiferum		92.45	0.49	BGL 10	1			
Lactobacillus acidophilus		3.37	0.11		99			
Lactobacillus fermentum		1.56	0.20		99			
03073101	**Good identification**							**03073101**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03073103	**Good identification**							**03073103**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03073121	**Good identification**							**03073121**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03073123	**Good identification**							**03073123**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03074000	**Low discrimination**							**03074000**
					GRA			
Fusobacterium mortiferum		78.78	0.37	XYL 7	1			
Actinomyces meyeri		21.22	0.18		99			
03074100	**Low discrimination**							**03074100**
					GRA			
Fusobacterium mortiferum		68.41	0.43	XYL 7	1			
Actinomyces meyeri		31.59	0.28		99			
03075000	**Low discrimination**							**03075000**
					GRA			
Fusobacterium mortiferum		77.51	0.40	XYL 7	1			
Actinomyces meyeri		20.87	0.21		99			
Lactobacillus fermentum		1.62	0.12		99			
03075100	**Low discrimination**							**03075100**
					COC	GRA		
Fusobacterium mortiferum		59.52	0.46	XYL 7	1	1		
Actinomyces meyeri		27.48	0.31		1	99		
Atopobium parvulum		12.13	0.23		99	99		
03100000	**Low discrimination**							**03100000**
					COC	GRA	SPO	
Clostridium botulinum biovar A		34.56	0.75	SUC 7	1	99	99	
Megasphaera elsdenii		30.81	0.78		99	1	1	
Peptostreptococcus anaerobius		13.12	0.63		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03100001	**Low discrimination**							**03100001**
					COC	GRA		
Clostridium novyi biovar A		64.23	0.66	SUC 1	1	99		
Clostridium botulinum biovar A		10.41	0.41		1	99		
Megasphaera elsdenii		9.28	0.44		99	1		
03100002	**Low discrimination**							**03100002**
					COC	GRA		
Megasphaera elsdenii		61.73	0.62	FRU 95	99	1		
Clostridium sporogenes		15.25	0.45	BGL 10	1	99		
Clostridium novyi biovar A		6.85	0.53		1	99		
03100003	**Low discrimination**							**03100003**
					COC	GRA		
Clostridium novyi biovar A		88.08	0.53	SUC 1	1	99		
Megasphaera elsdenii		8.02	0.28		99	1		
Clostridium sporogenes		1.98	0.11		1	99		
03100010	**Low discrimination**							**03100010**
					COC	GRA		
Clostridium novyi biovar A		68.46	0.66	SUC 1	1	99		
Clostridium botulinum biovar A		11.10	0.41		1	99		
Megasphaera elsdenii		9.89	0.44		99	1		
03100011	**Very good identification**							**03100011**
					COC	GRA		
Clostridium novyi biovar A		99.53	0.66	SUC 1	1	99		
Clostridium botulinum biovar A		0.16	0.06		1	99		
Megasphaera elsdenii		0.15	0.10		99	1		
03100012	**Low discrimination**							**03100012**
					COC	GRA		
Clostridium novyi biovar A		88.51	0.53	SUC 1	1	99		
Megasphaera elsdenii		8.05	0.28		99	1		
Clostridium sporogenes		1.99	0.11		1	99		
03100013	**Good identification**							**03100013**
Clostridium novyi biovar A		100.0	0.53	SUC 1				
03100020	**Low discrimination**							**03100020**
					COC	SPO		
Gemella morbillorum		24.06	0.48	MNS 99	99	1		
Clostridium novyi biovar A		19.89	0.56	NAG 10	1	99		
Actinomyces meyeri		14.15	0.45		1	1		
03100021	**Good identification**							**03100021**
					COC	SPO		
Clostridium novyi biovar A		96.11	0.56	SUC 1	1	99		
Gemella morbillorum		1.18	0.14		99	1		
Actinomyces meyeri		0.69	0.11		1	1		
03100022	**Low discrimination**							**03100022**
					COC	SPO	GRA	
Clostridium novyi biovar A		43.24	0.43	SUC 1	1	99	99	
Atopobium minutum		19.94	0.33		1	1	99	
Megasphaera elsdenii		15.74	0.28		99	1	1	
03100023	**Good identification**							**03100023**
Clostridium novyi biovar A		100.0	0.43	SUC 1				
03100030	**Good identification**							**03100030**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium novyi biovar A		95.93	0.56	SUC 1	1	99	
Gemella morbillorum		1.17	0.14		99	1	
Actinomyces meyeri		0.69	0.11		1	1	
03100031	**Good identification**						**03100031**
Clostridium novyi biovar A		100.0	0.56	SUC 1			
03100032	**Good identification**						**03100032**
Clostridium novyi biovar A		100.0	0.43	SUC 1			
03100033	**Good identification**						**03100033**
Clostridium novyi biovar A		100.0	0.43	SUC 1			
03100040	**Low discrimination**						**03100040**
					COC	GRA	
Megasphaera elsdenii		60.43	0.75	FRU 95	99	1	
Clostridium botulinum biovar A		30.37	0.66		1	99	
Clostridium novyi biovar A		4.22	0.63		1	99	
03100041	**Low discrimination**						**03100041**
					COC	GRA	
Clostridium novyi biovar A		81.10	0.63	SUC 1	1	99	
Megasphaera elsdenii		11.72	0.41		99	1	
Clostridium botulinum biovar A		5.89	0.32		1	99	
03100042	**Low discrimination**						**03100042**
					COC	GRA	
Megasphaera elsdenii		85.67	0.59	FRU 95	99	1	
Clostridium novyi biovar A		9.51	0.50	BGL 10	1	99	
Clostridium botulinum biovar A		3.91	0.32		1	99	
03100043	**Acceptable identification**						**03100043**
					COC	GRA	
Clostridium novyi biovar A		91.66	0.50	SUC 1	1	99	
Megasphaera elsdenii		8.34	0.25		99	1	
03100050	**Low discrimination**						**03100050**
					COC	GRA	
Clostridium novyi biovar A		60.33	0.63	SUC 1	1	99	
Clostridium botulinum biovar B		26.51	0.42		1	99	
Megasphaera elsdenii		8.72	0.41		99	1	
03100051	**Very good identification**						**03100051**
					COC	GRA	
Clostridium novyi biovar A		99.41	0.63	SUC 1	1	99	
Clostridium botulinum biovar B		0.44	0.08		1	99	
Megasphaera elsdenii		0.15	0.07		99	1	
03100052	**Low discrimination**						**03100052**
					COC	GRA	
Clostridium novyi biovar A		89.59	0.50	SUC 1	1	99	
Megasphaera elsdenii		8.15	0.25		99	1	
Clostridium botulinum biovar B		2.25	0.08		1	99	
03100053	**Good identification**						**03100053**
Clostridium novyi biovar A		100.0	0.50	SUC 1			
03100060	**Low discrimination**						**03100060**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		47.94	0.42	MNS 99	1	99		
Clostridium novyi biovar A		27.28	0.53		1	99		
Megasphaera elsdenii		15.77	0.41		99	1		
03100061	**Good identification**							**03100061**
					COC	GRA		
Clostridium novyi biovar A		97.70	0.53	SUC 1	1	99		
Clostridium botulinum biovar B		1.73	0.08		1	99		
Megasphaera elsdenii		0.57	0.07		99	1		
03100062	**Low discrimination**							**03100062**
					COC	GRA		
Clostridium novyi biovar A		68.28	0.40	SUC 1	1	99		
Megasphaera elsdenii		24.85	0.25		99	1		
Clostridium botulinum biovar B		6.87	0.08		1	99		
03100063	**Good identification**							**03100063**
Clostridium novyi biovar A		100.0	0.40	SUC 1				
03100070	**Low discrimination**							**03100070**
					COC	GRA		
Clostridium novyi biovar A		53.44	0.53	SUC 1	1	99		
Clostridium botulinum biovar B		46.25	0.37		1	99		
Megasphaera elsdenii		0.31	0.07		99	1		
03100071	**Good identification**							**03100071**
Clostridium novyi biovar A		99.13	0.53	SUC 1				
Clostridium botulinum biovar B		0.87	0.03					
03100072	**Good identification**							**03100072**
Clostridium novyi biovar A		95.28	0.40	SUC 1				
Clostridium botulinum biovar B		4.72	0.03					
03100073	**Good identification**							**03100073**
Clostridium novyi biovar A		100.0	0.40	SUC 1				
03100100	**Good identification**							**03100100**
					COC	SPO	GRA	
Actinomyces meyeri		91.53	0.70	FRU 95	1	1	99	
Clostridium botulinum biovar A		2.84	0.41		1	99	99	
Megasphaera elsdenii		2.54	0.44		99	1	1	
03100101	**Low discrimination**							**03100101**
					SPO			
Clostridium chauvoei		55.06	0.37	MNS 90	99			
Actinomyces meyeri		35.09	0.36	GAL 99	1			
Clostridium novyi biovar A		6.72	0.32		99			
03100102	**Low discrimination**							**03100102**
					COC	GRA		
Actinomyces meyeri		80.40	0.44	FRU 95	1	99		
Megasphaera elsdenii		8.00	0.28	BGL 3	99	1		
Actinomyces odontolyticus		3.97	0.26		1	99		
03100110	**Low discrimination**							**03100110**
					SPO			
Actinomyces meyeri		79.23	0.36	FRU 95	1			
Clostridium novyi biovar A		15.18	0.32	MLZ 1	99			
Clostridium botulinum biovar A		2.46	0.06		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03100120	**Low discrimination**							**03100120**
					GRA			
Actinomyces meyeri		74.28	0.55	FRU 95	99			
Prevotella bivia		23.04	0.32	NAG 12	1			
Prevotella melaninogenica		1.28	0.17		1			
03100140	**Low discrimination**							**03100140**
					COC	SPO	GRA	
Megasphaera elsdenii		45.53	0.41	FRU 95	99	1	1	
Actinomyces meyeri		24.91	0.36	LAC 1	1	1	99	
Clostridium botulinum biovar A		22.88	0.32		1	99	99	
03100200	**Low discrimination**							**03100200**
					COC	SPO	GRA	
Clostridium botulinum biovar A		29.60	0.41	SUC 7	1	99	99	
Megasphaera elsdenii		26.39	0.44	RHA 1	99	1	1	
Actinomyces odontolyticus		14.59	0.43		1	1	99	
03100202	**Low discrimination**							**03100202**
					COC	SPO	GRA	
Actinomyces odontolyticus		66.49	0.39	NIT 99	1	1	99	
Megasphaera elsdenii		21.80	0.28	FRU 92	99	1	1	
Clostridium sporogenes		5.39	0.11		1	99	99	
03100240	**Low discrimination**							**03100240**
					COC	GRA		
Megasphaera elsdenii		60.62	0.41	FRU 95	99	1		
Clostridium botulinum biovar A		30.47	0.32	RHA 1	1	99		
Clostridium novyi biovar A		4.24	0.29		1	99		
03100300	**Low discrimination**							**03100300**
					SPO			
Actinomyces meyeri		64.55	0.36	FRU 95	1			
Actinomyces odontolyticus		30.90	0.34	RHA 1	1			
Clostridium botulinum biovar A		2.01	0.06		99			
03100400	**Low discrimination**							**03100400**
					COC	SPO		
Actinomyces meyeri		30.82	0.50	FRU 95	1	1		
Gemella morbillorum		28.58	0.48		99	1		
Clostridium botulinum biovar A		15.32	0.41		1	99		
03100440	**Low discrimination**							**03100440**
					COC	GRA		
Megasphaera elsdenii		58.58	0.41	FRU 95	99	1		
Clostridium botulinum biovar A		29.44	0.32	ARA 1	1	99		
Clostridium novyi biovar A		4.09	0.29		1	99		
03100500	**Very good identification**							**03100500**
					COC			
Actinomyces meyeri		99.17	0.60	FRU 95	1			
Lactobacillus fermentum		0.32	0.28		1			
Gemella morbillorum		0.23	0.14		99			
03100520	**Good identification**							**03100520**
					GRA			
Actinomyces meyeri		98.00	0.45	FRU 95	99			
Prevotella bivia		1.23	-.03	NAG 12	1			
Lactobacillus fermentum		0.77	0.20		99			
03101000	**Low discrimination**							**03101000**
					COC			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		45.29	0.63	FRU 95	1		
Gemella morbillorum		33.92	0.60		99		
Lactobacillus fermentum		5.92	0.59		1		
03101001	**Low discrimination**						**03101001**
					SPO		
Clostridium novyi biovar A		44.14	0.50	SUC 1	99		
Clostridium chauvoei		32.86	0.37	GAL 10	99		
Actinomyces meyeri		7.85	0.29		1		
03101002	**Low discrimination**						**03101002**
Lactobacillus fermentum		62.36	0.59	FRU 99			
Actinomyces meyeri		14.76	0.37				
Atopobium minutum		4.96	0.33				
03101003	**Low discrimination**						**03101003**
					SPO		
Clostridium novyi biovar A		52.43	0.37	SUC 1	99		
Clostridium chauvoei		39.03	0.24	GAL 10	99		
Lactobacillus fermentum		6.91	0.24		1		
03101010	**Low discrimination**						**03101010**
					COC	SPO	
Clostridium novyi biovar A		72.52	0.50	SUC 1	1	99	
Actinomyces meyeri		12.90	0.29	GAL 10	1	1	
Gemella morbillorum		9.66	0.26		99	1	
03101011	**Good identification**						**03101011**
Clostridium novyi biovar A		99.25	0.50	SUC 1			
Clostridium chauvoei		0.75	0.03	GAL 10			
03101012	**Low discrimination**						**03101012**
					SPO		
Clostridium novyi biovar A		85.99	0.37	SUC 1	99		
Lactobacillus fermentum		11.33	0.24	GAL 10	1		
Actinomyces meyeri		2.68	0.03		1		
03101013	**Good identification**						**03101013**
Clostridium novyi biovar A		100.0	0.37	SUC 1			
03101020	**Low discrimination**						**03101020**
					COC		
Actinomyces meyeri		46.28	0.48	FRU 95	1		
Gemella morbillorum		28.25	0.43	NAG 12	99		
Lactobacillus fermentum		14.79	0.50		1		
03101021	**Low discrimination**						**03101021**
					SPO		
Clostridium novyi biovar A		45.87	0.39	SUC 1	99		
Clostridium chauvoei		45.54	0.29	GAL 10	99		
Actinomyces meyeri		4.45	0.14		1		
03101022	**Low discrimination**						**03101022**
Lactobacillus fermentum		75.35	0.50	FRU 99			
Actinomyces meyeri		7.29	0.22				
Atopobium minutum		5.99	0.25				
03101030	**Low discrimination**						**03101030**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		84.23	0.39	SUC 1	1	99	
Actinomyces meyeri		8.17	0.14	GAL 10	1	1	
Gemella morbillorum		4.99	0.09		99	1	
03101031	**Good identification**						**03101031**
Clostridium novyi biovar A		100.0	0.39	SUC 1			
03101040	**Low discrimination**						**03101040**
					COC	GRA	
Megasphaera elsdenii		36.18	0.41	FRU 95	99	1	
Clostridium novyi biovar A		27.82	0.47	GAL 1	1	99	
Clostridium botulinum biovar A		18.18	0.32		1	99	
03101041	**Good identification**						**03101041**
					COC	GRA	
Clostridium novyi biovar A		96.71	0.47	SUC 1	1	99	
Megasphaera elsdenii		1.27	0.07	GAL 10	99	1	
Clostridium chauvoei		1.09	0.03		1	99	
03101050	**Acceptable identification**						**03101050**
					COC	GRA	
Clostridium novyi biovar A		94.96	0.47	SUC 1	1	99	
Clostridium botulinum biovar B		3.79	0.08	GAL 10	1	99	
Megasphaera elsdenii		1.25	0.07		99	1	
03101051	**Good identification**						**03101051**
Clostridium novyi biovar A		100.0	0.47	SUC 1			
03101060	**Low discrimination**						**03101060**
					SPO		
Clostridium novyi biovar A		66.97	0.36	SUC 1	99		
Clostridium botulinum biovar B		10.70	0.08	GAL 10	99		
Actinomyces meyeri		9.75	0.14		1		
03101061	**Good identification**						**03101061**
Clostridium novyi biovar A		100.0	0.36	SUC 1			
03101070	**Acceptable identification**						**03101070**
Clostridium novyi biovar A		92.70	0.36	SUC 1			
Clostridium botulinum biovar B		7.30	0.03	GAL 10			
03101071	**Good identification**						**03101071**
Clostridium novyi biovar A		100.0	0.36	SUC 1			
03101100	**Good identification**						**03101100**
					COC		
Actinomyces meyeri		92.42	0.73	FRU 95	1		
Lactobacillus fermentum		4.93	0.62		1		
Blautia hansenii		1.03	0.32		99		
03101101	**Very good identification**						**03101101**
					SPO		
Clostridium chauvoei		98.96	0.71	MNS 90	99		
Actinomyces meyeri		0.96	0.39		1		
Lactobacillus fermentum		0.05	0.28		1		
03101102	**Low discrimination**						**03101102**
					SPO		
Lactobacillus fermentum		60.12	0.62	FRU 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		34.87	0.47		1		
Clostridium chauvoei		2.10	0.24		99		
03101103	**Very good identification**						**03101103**
					SPO		
Clostridium chauvoei		99.50	0.58	MNS 90	99		
Lactobacillus fermentum		0.29	0.28		1		
Actinomyces meyeri		0.17	0.13		1		
03101110	**Low discrimination**						**03101110**
Actinomyces meyeri		88.83	0.39	FRU 95			
Lactobacillus fermentum		4.74	0.28	MLZ 1			
Eubacterium saburreum		3.24	0.22				
03101111	**Good identification**						**03101111**
					SPO		
Clostridium chauvoei		97.73	0.37	MNS 90	99		
Clostridium novyi biovar A		1.33	0.15	MLZ 1	99		
Actinomyces meyeri		0.94	0.05		1		
03101120	**Low discrimination**						**03101120**
					GRA		
Prevotella melaninogenica		48.18	0.51	MNS 90	1		
Actinomyces meyeri		42.37	0.58	RAF 90	99		
Lactobacillus fermentum		5.52	0.54	FRU 90	99		
03101121	**Very good identification**						**03101121**
					GRA	SPO	
Clostridium chauvoei		99.09	0.63	MNS 90	99	99	
Prevotella melaninogenica		0.45	0.17		1	1	
Actinomyces meyeri		0.39	0.24		99	1	
03101122	**Low discrimination**						**03101122**
					GRA		
Prevotella melaninogenica		55.16	0.39	MNS 90	1		
Lactobacillus fermentum		33.19	0.54	RAF 90	99		
Actinomyces meyeri		7.88	0.33	FRU 90	99		
03101123	**Good identification**						**03101123**
					GRA	SPO	
Clostridium chauvoei		99.10	0.50	MNS 90	99	99	
Prevotella melaninogenica		0.48	0.05		1	1	
Lactobacillus fermentum		0.29	0.20		99	1	
03101140	**Acceptable identification**						**03101140**
					COC	GRA	
Actinomyces meyeri		90.38	0.39	FRU 95	1	99	
Lactobacillus fermentum		4.82	0.28	SOR 1	1	99	
Megasphaera elsdenii		1.11	0.07		99	1	
03101141	**Good identification**						**03101141**
					SPO		
Clostridium chauvoei		98.16	0.37	MNS 90	99		
Actinomyces meyeri		0.95	0.05	SOR 1	1		
Clostridium novyi biovar A		0.89	0.12		99		
03101200	**Low discrimination**						**03101200**
					COC		
Actinomyces odontolyticus		55.92	0.40	NIT 99	1		
Actinomyces meyeri		20.75	0.29	FRU 92	1		
Gemella morbillorum		15.54	0.26		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03101202	**Acceptable identification**					**03101202**
Actinomyces odontolyticus		91.09	0.36	NIT 99		
Lactobacillus fermentum		7.21	0.24	FRU 92		
Actinomyces meyeri		1.71	0.03			
03101300	**Low discrimination**					**03101300**
Actinomyces meyeri		77.65	0.39	FRU 95		
Actinomyces odontolyticus		16.52	0.31	RHA 1		
Lactobacillus fermentum		4.14	0.28			
03101301	**Good identification**					**03101301**
					SPO	
Clostridium chauvoei		99.04	0.37	MNS 90	99	
Actinomyces meyeri		0.96	0.05	RHA 1	1	
03101400	**Low discrimination**					**03101400**
					COC	
Actinomyces meyeri		53.60	0.53	FRU 95	1	
Lactobacillus fermentum		28.03	0.59		1	
Gemella morbillorum		17.84	0.43		99	
03101402	**Good identification**					**03101402**
					COC	
Lactobacillus fermentum		93.85	0.59	FRU 99	1	
Actinomyces meyeri		5.55	0.27		1	
Gemella morbillorum		0.60	0.09		99	
03101420	**Low discrimination**					**03101420**
					COC	
Lactobacillus fermentum		50.14	0.50	FRU 99	1	
Actinomyces meyeri		39.23	0.38		1	
Gemella morbillorum		10.64	0.27		99	
03101422	**Good identification**					**03101422**
Lactobacillus fermentum		97.64	0.50	FRU 99		
Actinomyces meyeri		2.36	0.12			
03101500	**Low discrimination**					**03101500**
					COC	
Actinomyces meyeri		81.66	0.63	FRU 95	1	
Lactobacillus fermentum		17.42	0.62		1	
Blautia hansenii		0.64	0.19		99	
03101501	**Low discrimination**					**03101501**
					SPO	
Clostridium chauvoei		77.52	0.37	MNS 90	99	
Actinomyces meyeri		18.53	0.29	ARA 1	1	
Lactobacillus fermentum		3.95	0.28		1	
03101502	**Low discrimination**					**03101502**
Lactobacillus fermentum		86.24	0.62	FRU 99		
Actinomyces meyeri		12.51	0.37			
Bifidobacterium longum subsp. longum		0.91	0.21			
03101520	**Low discrimination**					**03101520**
Actinomyces meyeri		61.00	0.48	FRU 95		
Lactobacillus fermentum		31.80	0.54	NAG 12		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		4.00	0.32					
03101522	**Low discrimination**							**03101522**
Lactobacillus fermentum		83.20	0.54	FRU 99				
Bifidobacterium longum subsp. longum		10.47	0.32					
Actinomyces meyeri		4.94	0.22					
03101530	**Low discrimination**							**03101530**
Bifidobacterium longum subsp. longum		87.91	0.36	MNS 87				
Actinomyces meyeri		7.95	0.14	RAF 97				
Lactobacillus fermentum		4.14	0.20	FRU 90				
03101532	**Good identification**							**03101532**
Bifidobacterium longum subsp. longum		95.50	0.36	MNS 87				
Lactobacillus fermentum		4.50	0.20	RAF 97				
03102000	**Very good identification**							**03102000**
					COC	GRA	SPO	
Megasphaera elsdenii		97.77	0.78	FRU 95	99	1	1	
Clostridium botulinum biovar A		1.11	0.41		1	99	99	
Peptostreptococcus anaerobius		0.42	0.29		99	99	1	
03102001	**Acceptable identification**							**03102001**
					COC	GRA		
Megasphaera elsdenii		91.55	0.44	FRU 95	99	1		
Clostridium novyi biovar A		6.40	0.32	URE 1	1	99		
Clostridium botulinum biovar A		1.04	0.06		1	99		
03102002	**Very good identification**							**03102002**
					COC	GRA		
Megasphaera elsdenii		99.39	0.62	FRU 95	99	1		
Clostridium sporogenes		0.25	0.11	BGL 10	1	99		
Clostridium novyi biovar A		0.11	0.19		1	99		
03102010	**Acceptable identification**							**03102010**
					COC	GRA		
Megasphaera elsdenii		92.12	0.44	FRU 95	99	1		
Clostridium novyi biovar A		6.44	0.32	MLZ 1	1	99		
Clostridium botulinum biovar A		1.04	0.06		1	99		
03102020	**Acceptable identification**							**03102020**
					COC	GRA	SPO	
Megasphaera elsdenii		93.40	0.44	FRU 95	99	1	1	
Gemella morbillorum		1.98	0.14	NAG 1	99	99	1	
Clostridium novyi biovar A		1.63	0.21		1	99	99	
03102040	**Very good identification**							**03102040**
					COC	GRA		
Megasphaera elsdenii		99.35	0.75	FRU 95	99	1		
Clostridium botulinum biovar A		0.50	0.32		1	99		
Clostridium novyi biovar A		0.07	0.29		1	99		
03102041	**Acceptable identification**							**03102041**
					COC	GRA		
Megasphaera elsdenii		93.47	0.41	FRU 95	99	1		
Clostridium novyi biovar A		6.53	0.29	URE 1	1	99		
03102042	**Very good identification**							**03102042**
					COC	SPO	GRA	
Megasphaera elsdenii		99.78	0.59	FRU 95	99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		0.11	0.16	BGL 10	1	99	99	
Propionibacterium propionicum		0.11	0.18		1	1	99	
03102050	**Acceptable identification**							**03102050**
					COC	GRA		
Megasphaera elsdenii		90.86	0.41	FRU 95	99	1		
Clostridium novyi biovar A		6.35	0.29	MLZ 1	1	99		
Clostridium botulinum biovar B		2.79	0.08		1	99		
03102060	**Good identification**							**03102060**
					COC	GRA		
Megasphaera elsdenii		95.40	0.41	FRU 95	99	1		
Clostridium botulinum biovar B		2.93	0.08	NAG 1	1	99		
Clostridium novyi biovar A		1.67	0.18		1	99		
03102100	**Low discrimination**							**03102100**
					COC	GRA		
Megasphaera elsdenii		52.52	0.44	FRU 95	99	1		
Propionibacterium propionicum		27.51	0.49	LAC 1	1	99		
Actinomyces meyeri		19.16	0.36		1	99		
03102102	**Low discrimination**							**03102102**
					COC	GRA		
Propionibacterium propionicum		86.52	0.52	NIT 86	1	99		
Megasphaera elsdenii		12.24	0.28	FRU 99	99	1		
Actinomyces meyeri		1.24	0.10		1	99		
03102103	**Good identification**							**03102103**
Propionibacterium propionicum		100.0	0.36	NIT 86				
03102140	**Low discrimination**							**03102140**
					COC	GRA		
Megasphaera elsdenii		55.83	0.41	FRU 95	99	1		
Propionibacterium propionicum		43.86	0.49	LAC 1	1	99		
Actinomyces meyeri		0.31	0.02		1	99		
03102142	**Acceptable identification**							**03102142**
					COC	GRA		
Propionibacterium propionicum		91.38	0.52	NIT 86	1	99		
Megasphaera elsdenii		8.62	0.25	FRU 99	99	1		
03102143	**Good identification**							**03102143**
Propionibacterium propionicum		100.0	0.36	NIT 86				
03102200	**Good identification**							**03102200**
					COC	SPO	GRA	
Megasphaera elsdenii		97.92	0.44	FRU 95	99	1	1	
Clostridium botulinum biovar A		1.11	0.06	RHA 1	1	99	99	
Actinomyces odontolyticus		0.55	0.09		1	1	99	
03102240	**Good identification**							**03102240**
Megasphaera elsdenii		100.0	0.41	FRU 95				
03102400	**Acceptable identification**							**03102400**
					COC	GRA		
Megasphaera elsdenii		94.38	0.44	FRU 95	99	1		
Actinomyces meyeri		2.15	0.15	ARA 1	1	99		
Gemella morbillorum		2.00	0.14		99	99		
03102440	**Good identification**							**03102440**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		100.0	0.41	FRU 95				
03103000	**Low discrimination**							**03103000**
					COC	GRA		
Megasphaera elsdenii		77.29	0.44	FRU 95	99	1		
Actinomyces meyeri		10.57	0.29	GAL 1	1	99		
Gemella morbillorum		7.92	0.26		99	99		
03103040	**Good identification**							**03103040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.45	0.41	FRU 95	99	1	1	
Propionibacterium propionicum		0.78	0.15	GAL 1	1	1	99	
Clostridium novyi biovar A		0.76	0.12		1	99	99	
03103100	**Low discrimination**							**03103100**
Actinomyces meyeri		48.53	0.39	FRU 95				
Propionibacterium propionicum		46.45	0.49	MAN 1				
Lactobacillus fermentum		2.59	0.28					
03103101	**Low discrimination**							**03103101**
					SPO			
Clostridium chauvoei		88.83	0.37	MNS 90	99			
Propionibacterium propionicum		9.03	0.33	MAN 1	1			
Actinomyces naeslundii		1.28	0.29		1			
03103102	**Acceptable identification**							**03103102**
Propionibacterium propionicum		93.84	0.52	NIT 86				
Lactobacillus fermentum		3.48	0.28	FRU 99				
Actinomyces meyeri		2.02	0.13					
03103140	**Good identification**							**03103140**
					COC	GRA		
Propionibacterium propionicum		97.71	0.49	NIT 86	1	99		
Megasphaera elsdenii		1.26	0.07	FRU 99	99	1		
Actinomyces meyeri		1.03	0.05		1	99		
03103142	**Good identification**							**03103142**
Propionibacterium propionicum		100.0	0.52	NIT 86				
03103143	**Good identification**							**03103143**
Propionibacterium propionicum		100.0	0.36	NIT 86				
03104000	**Low discrimination**							**03104000**
					COC	SPO		
Actinomyces meyeri		76.35	0.65	FRU 95	1	1		
Gemella morbillorum		9.12	0.48		99	1		
Clostridium botulinum biovar A		4.89	0.41		1	99		
03104002	**Low discrimination**							**03104002**
					COC	GRA		
Actinomyces meyeri		65.27	0.39	FRU 95	1	99		
Megasphaera elsdenii		13.38	0.28	BGL 3	99	1		
Actinomyces odontolyticus		10.36	0.29		1	99		
03104020	**Low discrimination**							**03104020**
					COC			
Actinomyces meyeri		88.65	0.50	FRU 95	1			
Gemella morbillorum		8.63	0.32	NAG 12	99			
Actinomyces odontolyticus		0.85	0.19		1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
03104040	**Low discrimination**							**03104040**
					COC	GRA		
Clostridium glycolicum		77.08	0.50	SUC 1	1	99		
Megasphaera elsdenii		11.83	0.41	FRU 90	99	1		
Clostridium botulinum biovar A		5.94	0.32		1	99		
03104100	**Very good identification**							**03104100**
					COC			
Actinomyces meyeri		99.85	0.75	FRU 95	1			
Actinomyces odontolyticus		0.06	0.24		1			
Gemella morbillorum		0.03	0.14		99			
03104101	**Good identification**							**03104101**
					SPO			
Actinomyces meyeri		99.19	0.41	FRU 95	1			
Clostridium chauvoei		0.81	0.03	URE 1	99			
03104102	**Good identification**							**03104102**
Actinomyces meyeri		98.00	0.49	FRU 95				
Actinomyces odontolyticus		1.23	0.20	BGL 3				
Lactobacillus fermentum		0.77	0.24					
03104110	**Good identification**							**03104110**
Actinomyces meyeri		100.0	0.41	FRU 95				
03104120	**Very good identification**							**03104120**
					GRA			
Actinomyces meyeri		99.70	0.60	FRU 95	99			
Prevotella bivia		0.16	-.03	NAG 12	1			
Actinomyces odontolyticus		0.08	0.11		99			
03104140	**Acceptable identification**							**03104140**
					COC	SPO	GRA	
Actinomyces meyeri		93.33	0.41	FRU 95	1	1	99	
Clostridium glycolicum		5.79	0.15	SOR 1	1	99	99	
Megasphaera elsdenii		0.89	0.07		99	1	1	
03104200	**Low discrimination**							**03104200**
					COC			
Actinomyces odontolyticus		54.88	0.38	NIT 99	1			
Actinomyces meyeri		35.67	0.31	FRU 92	1			
Gemella morbillorum		4.26	0.14		99			
03104300	**Low discrimination**							**03104300**
Actinomyces meyeri		89.17	0.41	FRU 95				
Actinomyces odontolyticus		10.83	0.29	RHA 1				
03104400	**Acceptable identification**							**03104400**
					COC	SPO		
Actinomyces meyeri		94.22	0.55	FRU 95	1	1		
Gemella morbillorum		5.00	0.32		99	1		
Clostridium botulinum biovar A		0.24	0.06		1	99		
03104420	**Good identification**							**03104420**
					COC			
Actinomyces meyeri		95.32	0.40	FRU 95	1			
Gemella morbillorum		4.12	0.15	NAG 12	99			
Lactobacillus fermentum		0.56	0.12		1			
03104500	**Very good identification**							**03104500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		99.90	0.65	FRU 95		
Lactobacillus fermentum		0.10	0.24			
03104502	**Good identification**					**03104502**
Actinomyces meyeri		96.94	0.39	FRU 95		
Lactobacillus fermentum		3.06	0.24	BGL 3		
03104520	**Good identification**					**03104520**
Actinomyces meyeri		99.61	0.50	FRU 95		
Lactobacillus fermentum		0.24	0.16	NAG 12		
Bifidobacterium longum subsp. longum		0.15	0.06			
03105000	**Good identification**					**03105000**
					COC	
Actinomyces meyeri		91.88	0.68	FRU 95	1	
Gemella morbillorum		3.94	0.43		99	
Lactobacillus fermentum		3.63	0.55		1	
03105002	**Low discrimination**					**03105002**
Lactobacillus fermentum		54.11	0.55	FRU 99		
Actinomyces meyeri		42.32	0.42			
Actinomyces odontolyticus		2.98	0.26			
03105020	**Low discrimination**					**03105020**
					COC	
Actinomyces meyeri		88.04	0.53	FRU 95	1	
Lactobacillus fermentum		8.51	0.46	NAG 12	1	
Gemella morbillorum		3.08	0.27		99	
03105022	**Low discrimination**					**03105022**
Lactobacillus fermentum		74.25	0.46	FRU 99		
Actinomyces meyeri		23.75	0.27			
Actinomyces odontolyticus		2.00	0.12			
03105100	**Very good identification**					**03105100**
					COC	
Actinomyces meyeri		98.26	0.78	FRU 95	1	
Lactobacillus fermentum		1.59	0.58		1	
Blautia hansenii		0.10	0.19		99	
03105101	**Low discrimination**					**03105101**
					SPO	
Actinomyces meyeri		64.31	0.44	FRU 95	1	
Clostridium chauvoei		34.66	0.37	URE 1	99	
Lactobacillus fermentum		1.04	0.24		1	
03105102	**Low discrimination**					**03105102**
Actinomyces meyeri		65.33	0.52	FRU 95		
Lactobacillus fermentum		34.07	0.58	BGL 3		
Actinomyces odontolyticus		0.36	0.17			
03105110	**Good identification**					**03105110**
Actinomyces meyeri		97.44	0.44	FRU 95		
Lactobacillus fermentum		1.57	0.24	MLZ 1		
Eubacterium saburreum		0.99	0.17			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03105120	**Very good identification**					**03105120**
					GRA	
Actinomyces meyeri		95.53	0.63	FRU 95	99	
Lactobacillus fermentum		3.77	0.50	NAG 12	99	
Prevotella melaninogenica		0.57	0.17		1	
03105122	**Low discrimination**					**03105122**
					GRA	
Lactobacillus fermentum		54.96	0.50	FRU 99	99	
Actinomyces meyeri		43.11	0.38		99	
Prevotella melaninogenica		1.57	0.05		1	
03105140	**Good identification**					**03105140**
					SPO	
Actinomyces meyeri		97.97	0.44	FRU 95	1	
Lactobacillus fermentum		1.58	0.24	SOR 1	1	
Clostridium glycolicum		0.45	-.01		99	
03105300	**Acceptable identification**					**03105300**
Actinomyces meyeri		93.45	0.44	FRU 95		
Actinomyces odontolyticus		5.05	0.26	RHA 1		
Lactobacillus fermentum		1.51	0.24			
03105400	**Low discrimination**					**03105400**
					COC	
Actinomyces meyeri		84.94	0.58	FRU 95	1	
Lactobacillus fermentum		13.44	0.55		1	
Gemella morbillorum		1.62	0.27		99	
03105402	**Low discrimination**					**03105402**
Lactobacillus fermentum		83.65	0.55	FRU 99		
Actinomyces meyeri		16.35	0.32			
03105420	**Low discrimination**					**03105420**
					COC	
Actinomyces meyeri		71.11	0.43	FRU 95	1	
Lactobacillus fermentum		27.50	0.46	NAG 12	1	
Gemella morbillorum		1.10	0.11		99	
03105422	**Acceptable identification**					**03105422**
Lactobacillus fermentum		91.69	0.46	FRU 99		
Actinomyces meyeri		7.33	0.17			
Bifidobacterium longum subsp. longum		0.97	0.06			
03105500	**Good identification**					**03105500**
Actinomyces meyeri		93.57	0.68	FRU 95		
Lactobacillus fermentum		6.04	0.58			
Bifidobacterium longum subsp. longum		0.32	0.29			
03105502	**Low discrimination**					**03105502**
Lactobacillus fermentum		65.24	0.58	FRU 99		
Actinomyces meyeri		31.27	0.42			
Bifidobacterium longum subsp. longum		3.50	0.29			
03105520	**Low discrimination**					**03105520**
Actinomyces meyeri		79.42	0.53	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		12.53	0.50	NAG 12				
Bifidobacterium longum subsp. longum		8.05	0.40					
03105522	**Low discrimination**							**03105522**
Lactobacillus fermentum		54.38	0.50	FRU 99				
Bifidobacterium longum subsp. longum		34.96	0.40					
Actinomyces meyeri		10.66	0.27					
03105530	**Acceptable identification**							**03105530**
Bifidobacterium longum subsp. longum		93.66	0.44	MNS 87				
Actinomyces meyeri		5.48	0.19	RAF 97				
Lactobacillus fermentum		0.86	0.16	FRU 90				
03105532	**Good identification**							**03105532**
Bifidobacterium longum subsp. longum		99.09	0.44	MNS 87				
Lactobacillus fermentum		0.91	0.16	RAF 97				
03106000	**Low discrimination**							**03106000**
					COC	GRA		
Megasphaera elsdenii		82.39	0.44	FRU 95	99	1		
Actinomyces meyeri		14.58	0.31	XYL 1	1	99		
Gemella morbillorum		1.74	0.14		99	99		
03106040	**Acceptable identification**							**03106040**
					COC	GRA		
Megasphaera elsdenii		93.82	0.41	FRU 95	99	1		
Clostridium glycolicum		6.18	0.15	XYL 1	1	99		
03106100	**Good identification**							**03106100**
					COC	GRA		
Actinomyces meyeri		97.87	0.41	FRU 95	1	99		
Megasphaera elsdenii		1.40	0.10	MAN 1	99	1		
Propionibacterium propionicum		0.73	0.15		1	99		
03107100	**Good identification**							**03107100**
Actinomyces meyeri		97.93	0.44	FRU 95				
Lactobacillus fermentum		1.58	0.24	MAN 1				
Propionibacterium propionicum		0.49	0.15					
03110000	**Low discrimination**							**03110000**
					COC	GRA		
Megasphaera elsdenii		87.45	1.00		99	1		
Actinomyces meyeri		7.97	0.82		1	99		
Prevotella intermedia		1.52	0.68		1	1		
03110001	**Low discrimination**							**03110001**
					COC	SPO	GRA	
Megasphaera elsdenii		82.02	0.66	URE 1	99	1	1	
Actinomyces meyeri		7.48	0.48		1	1	99	
Clostridium novyi biovar A		3.32	0.50		1	99	99	
03110002	**Good identification**							**03110002**
					COC	GRA		
Megasphaera elsdenii		94.75	0.84	BGL 10	99	1		
Actinomyces meyeri		2.41	0.56		1	99		
Actinomyces odontolyticus		0.91	0.52		1	99		
03110003	**Low discrimination**							**03110003**
					COC	SPO	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		89.09	0.50	URE 1	99	1	1	
Clostridium novyi biovar A		5.73	0.37	BGL 10	1	99	99	
Actinomyces meyeri		2.26	0.22		1	1	99	
03110010	**Low discrimination**							**03110010**
					COC	SPO	GRA	
Megasphaera elsdenii		83.67	0.66	MLZ 1	99	1	1	
Actinomyces meyeri		7.63	0.48		1	1	99	
Clostridium novyi biovar A		3.39	0.50		1	99	99	
03110011	**Low discrimination**							**03110011**
					COC	SPO	GRA	
Clostridium novyi biovar A		78.05	0.50	SUC 1	1	99	99	
Megasphaera elsdenii		19.48	0.32	FRU 10	99	1	1	
Actinomyces meyeri		1.78	0.13		1	1	99	
03110012	**Low discrimination**							**03110012**
					COC	SPO	GRA	
Megasphaera elsdenii		88.58	0.50	MLZ 1	99	1	1	
Clostridium novyi biovar A		5.69	0.37	BGL 10	1	99	99	
Actinomyces meyeri		2.25	0.22		1	1	99	
03110013	**Low discrimination**							**03110013**
					COC	GRA		
Clostridium novyi biovar A		86.42	0.37	SUC 1	1	99		
Megasphaera elsdenii		13.58	0.15	FRU 10	99	1		
03110020	**Low discrimination**							**03110020**
					COC	GRA		
Actinomyces meyeri		48.69	0.67	NAG 12	1	99		
Megasphaera elsdenii		39.55	0.66		99	1		
Gemella morbillorum		5.32	0.50		99	99		
03110022	**Low discrimination**							**03110022**
					COC	GRA		
Megasphaera elsdenii		52.25	0.50	NAG 1	99	1		
Actinomyces meyeri		17.90	0.41	BGL 10	1	99		
Lactobacillus fermentum		14.60	0.50		1	99		
03110031	**Good identification**							**03110031**
Clostridium novyi biovar A		100.0	0.39	SUC 1				
03110040	**Very good identification**							**03110040**
					COC	GRA		
Megasphaera elsdenii		98.12	0.97		99	1		
Clostridium botulinum biovar B		1.29	0.58		1	99		
Clostridium botulinum biovar A		0.17	0.45		1	99		
03110041	**Good identification**							**03110041**
					COC	GRA		
Megasphaera elsdenii		94.30	0.63	URE 1	99	1		
Clostridium novyi biovar A		3.82	0.47		1	99		
Clostridium botulinum biovar B		1.24	0.24		1	99		
03110042	**Very good identification**							**03110042**
					COC	GRA		
Megasphaera elsdenii		99.69	0.81	BGL 10	99	1		
Clostridium botulinum biovar B		0.12	0.24		1	99		
Clostridium novyi biovar A		0.07	0.34		1	99		
03110043	**Acceptable identification**							**03110043**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		93.96	0.47	URE 1	99	1		
Clostridium novyi biovar A		6.04	0.34	BGL 10	1	99		
03110050	**Low discrimination**							**03110050**
					COC	GRA		
Megasphaera elsdenii		59.34	0.63	MLZ 1	99	1		
Clostridium botulinum biovar B		37.98	0.52		1	99		
Clostridium novyi biovar A		2.40	0.47		1	99		
03110051	**Low discrimination**							**03110051**
					COC	GRA		
Clostridium novyi biovar A		70.95	0.47	SUC 1	1	99		
Megasphaera elsdenii		17.71	0.29	FRU 10	99	1		
Clostridium botulinum biovar B		11.34	0.18		1	99		
03110052	**Low discrimination**							**03110052**
					COC	GRA		
Megasphaera elsdenii		89.09	0.47	MLZ 1	99	1		
Clostridium novyi biovar A		5.73	0.34	BGL 10	1	99		
Clostridium botulinum biovar B		5.18	0.18		1	99		
03110060	**Low discrimination**							**03110060**
					COC	SPO	GRA	
Megasphaera elsdenii		59.59	0.63	NAG 1	99	1	1	
Clostridium botulinum biovar B		38.14	0.52		1	99	99	
Actinomyces meyeri		1.11	0.33		1	1	99	
03110061	**Low discrimination**							**03110061**
					COC	GRA		
Clostridium novyi biovar A		37.91	0.36	SUC 1	1	99		
Megasphaera elsdenii		37.86	0.29	FRU 10	99	1		
Clostridium botulinum biovar B		24.23	0.18		1	99		
03110062	**Acceptable identification**							**03110062**
					COC	GRA		
Megasphaera elsdenii		92.04	0.47	NAG 1	99	1		
Clostridium botulinum biovar B		5.36	0.18	BGL 10	1	99		
Clostridium novyi biovar A		1.48	0.23		1	99		
03110070	**Acceptable identification**							**03110070**
					COC	GRA		
Clostridium botulinum biovar B		93.97	0.47	MNS 99	1	99		
Clostridium novyi biovar A		3.02	0.36		1	99		
Megasphaera elsdenii		3.01	0.29		99	1		
03110071	**Low discrimination**							**03110071**
Clostridium novyi biovar A		76.06	0.36	SUC 1				
Clostridium botulinum biovar B		23.94	0.13	FRU 10				
03110100	**Very good identification**							**03110100**
					COC	GRA		
Actinomyces meyeri		95.99	0.92		1	99		
Megasphaera elsdenii		2.66	0.66		99	1		
Fusobacterium mortiferum		0.60	0.64		1	1		
03110101	**Low discrimination**							**03110101**
					COC	SPO	GRA	
Actinomyces meyeri		80.46	0.58	URE 1	1	1	99	
Clostridium chauvoei		15.51	0.44		1	99	99	
Megasphaera elsdenii		2.23	0.32		99	1	1	
03110102	**Low discrimination**							**03110102**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Actinomyces meyeri		75.22	0.66	BGL 3	1	99		
Lactobacillus fermentum		10.24	0.62		1	99		
Megasphaera elsdenii		7.48	0.50		99	1		
03110110	**Very good identification**							**03110110**
					COC	GRA		
Actinomyces meyeri		96.01	0.58	MLZ 1	1	99		
Megasphaera elsdenii		2.66	0.32		99	1		
Fusobacterium mortiferum		0.60	0.30		1	1		
03110120	**Very good identification**							**03110120**
					GRA			
Actinomyces meyeri		97.43	0.77	NAG 12	99			
Lactobacillus fermentum		1.00	0.54		99			
Prevotella melaninogenica		0.80	0.33		1			
03110121	**Low discrimination**							**03110121**
					SPO			
Actinomyces meyeri		67.14	0.43	NAG 12	1			
Clostridium chauvoei		31.63	0.35	URE 1	99			
Lactobacillus fermentum		0.69	0.20		1			
03110122	**Low discrimination**							**03110122**
					GRA			
Actinomyces meyeri		68.87	0.51	NAG 12	99			
Lactobacillus fermentum		22.91	0.54	BGL 3	99			
Prevotella melaninogenica		3.46	0.21		1			
03110130	**Good identification**							**03110130**
					GRA			
Actinomyces meyeri		98.19	0.43	MLZ 1	99			
Lactobacillus fermentum		1.01	0.20	NAG 12	99			
Prevotella melaninogenica		0.80	-.01		1			
03110140	**Low discrimination**							**03110140**
					COC	GRA		
Megasphaera elsdenii		62.04	0.63	LAC 1	99	1		
Actinomyces meyeri		33.94	0.58		1	99		
Propionibacterium propionicum		2.57	0.49		1	99		
03110142	**Low discrimination**							**03110142**
					COC	GRA		
Megasphaera elsdenii		57.50	0.47	LAC 1	99	1		
Propionibacterium propionicum		32.10	0.52	BGL 10	1	99		
Actinomyces meyeri		8.76	0.32		1	99		
03110143	**Low discrimination**							**03110143**
					COC	GRA		
Propionibacterium propionicum		86.00	0.36	NIT 86	1	99		
Megasphaera elsdenii		14.01	0.12	MAN 99	99	1		
03110160	**Low discrimination**							**03110160**
					COC	SPO	GRA	
Actinomyces meyeri		78.56	0.43	NAG 12	1	1	99	
Megasphaera elsdenii		10.64	0.29	SOR 1	99	1	1	
Clostridium botulinum biovar B		6.81	0.18		1	99	99	
03110200	**Low discrimination**							**03110200**
					COC	GRA		
Megasphaera elsdenii		67.69	0.66	RHA 1	99	1		
Actinomyces odontolyticus		22.65	0.61		1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		6.17	0.48		1	99		
03110202	**Low discrimination**							**03110202**
					COC	GRA		
Actinomyces odontolyticus		63.92	0.57	NIT 99	1	99		
Megasphaera elsdenii		34.63	0.50		99	1		
Actinomyces meyeri		0.88	0.22		1	99		
03110220	**Low discrimination**							**03110220**
					COC	GRA		
Actinomyces odontolyticus		68.68	0.47	NIT 99	1	99		
Actinomyces meyeri		15.68	0.33	NAG 14	1	99		
Megasphaera elsdenii		12.73	0.32		99	1		
03110222	**Acceptable identification**							**03110222**
					COC	GRA		
Actinomyces odontolyticus		94.83	0.44	NIT 99	1	99		
Megasphaera elsdenii		3.19	0.15	NAG 14	99	1		
Actinomyces meyeri		1.09	0.07		1	99		
03110240	**Very good identification**							**03110240**
					COC	SPO	GRA	
Megasphaera elsdenii		97.72	0.63	RHA 1	99	1	1	
Clostridium botulinum biovar B		1.28	0.24		1	99	99	
Actinomyces odontolyticus		0.50	0.27		1	1	99	
03110242	**Good identification**							**03110242**
					COC	GRA		
Megasphaera elsdenii		97.28	0.47	RHA 1	99	1		
Actinomyces odontolyticus		2.72	0.23	BGL 10	1	99		
03110300	**Low discrimination**							**03110300**
					COC	GRA		
Actinomyces meyeri		75.26	0.58	RHA 1	1	99		
Actinomyces odontolyticus		21.81	0.52		1	99		
Megasphaera elsdenii		2.08	0.32		99	1		
03110302	**Low discrimination**							**03110302**
Actinomyces odontolyticus		81.60	0.49	NIT 99				
Actinomyces meyeri		14.21	0.32					
Lactobacillus fermentum		1.93	0.28					
03110320	**Low discrimination**							**03110320**
Actinomyces meyeri		73.30	0.43	RHA 1				
Actinomyces odontolyticus		25.35	0.39	NAG 12				
Lactobacillus fermentum		0.75	0.20					
03110322	**Low discrimination**							**03110322**
Actinomyces odontolyticus		83.72	0.35	NIT 99				
Actinomyces meyeri		12.21	0.17	NAG 14				
Lactobacillus fermentum		4.06	0.20					
03110400	**Low discrimination**							**03110400**
					COC	GRA		
Actinomyces meyeri		64.05	0.71		1	99		
Megasphaera elsdenii		28.38	0.66		99	1		
Gemella morbillorum		3.82	0.50		99	99		
03110401	**Low discrimination**							**03110401**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		63.39	0.37	URE 1	1	99		
Megasphaera elsdenii		28.09	0.32		99	1		
Gemella morbillorum		3.78	0.15		99	99		
03110402	**Low discrimination**							**03110402**
					COC	GRA		
Megasphaera elsdenii		40.12	0.50	ARA 1	99	1		
Lactobacillus fermentum		33.64	0.59	BGL 10	1	99		
Actinomyces meyeri		25.20	0.46		1	99		
03110410	**Low discrimination**							**03110410**
					COC	GRA		
Actinomyces meyeri		64.02	0.37	MLZ 1	1	99		
Megasphaera elsdenii		28.37	0.32		99	1		
Gemella morbillorum		3.82	0.15		99	99		
03110420	**Low discrimination**							**03110420**
					COC			
Actinomyces meyeri		84.47	0.57	NAG 12	1			
Lactobacillus fermentum		8.52	0.50		1			
Gemella morbillorum		4.11	0.33		99			
03110422	**Low discrimination**							**03110422**
					COC	GRA		
Lactobacillus fermentum		74.48	0.50	GAL 99	1	99		
Actinomyces meyeri		22.83	0.31		1	99		
Megasphaera elsdenii		2.69	0.15		99	1		
03110440	**Good identification**							**03110440**
					COC	SPO	GRA	
Megasphaera elsdenii		94.83	0.63	ARA 1	99	1	1	
Actinomyces meyeri		3.24	0.37		1	1	99	
Clostridium botulinum biovar B		1.25	0.24		1	99	99	
03110442	**Good identification**							**03110442**
					COC	GRA		
Megasphaera elsdenii		97.83	0.47	ARA 1	99	1		
Lactobacillus fermentum		1.24	0.24	BGL 10	1	99		
Actinomyces meyeri		0.93	0.11		1	99		
03110500	**Very good identification**							**03110500**
					COC	GRA		
Actinomyces meyeri		98.19	0.82		1	99		
Lactobacillus fermentum		1.65	0.62		1	99		
Megasphaera elsdenii		0.11	0.32		99	1		
03110501	**Good identification**							**03110501**
					SPO			
Actinomyces meyeri		97.60	0.48	URE 1	1			
Lactobacillus fermentum		1.64	0.28		1			
Clostridium chauvoei		0.76	0.09		99			
03110502	**Low discrimination**							**03110502**
					COC	GRA		
Actinomyces meyeri		64.43	0.56	BGL 3	1	99		
Lactobacillus fermentum		35.08	0.62		1	99		
Megasphaera elsdenii		0.26	0.15		99	1		
03110510	**Good identification**							**03110510**
Actinomyces meyeri		98.08	0.48	MLZ 1				
Lactobacillus fermentum		1.65	0.28					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		0.27	0.07				
03110520	**Very good identification**						**03110520**
Actinomyces meyeri		95.97	0.67	NAG 12			
Lactobacillus fermentum		3.95	0.54				
Bifidobacterium longum subsp. longum		0.05	0.14				
03110522	**Low discrimination**						**03110522**
Lactobacillus fermentum		56.73	0.54	GAL 99			
Actinomyces meyeri		42.63	0.41				
Bifidobacterium longum subsp. longum		0.65	0.14				
03110540	**Acceptable identification**						**03110540**
					COC	GRA	
Actinomyces meyeri		91.43	0.48	SOR 1	1	99	
Megasphaera elsdenii		6.75	0.29		99	1	
Lactobacillus fermentum		1.54	0.28		1	99	
03110600	**Low discrimination**						**03110600**
					COC	GRA	
Actinomyces meyeri		59.09	0.37	RHA 1	1	99	
Megasphaera elsdenii		26.18	0.32		99	1	
Actinomyces odontolyticus		8.76	0.27		1	99	
03110700	**Good identification**						**03110700**
Actinomyces meyeri		97.23	0.48	RHA 1			
Lactobacillus fermentum		1.64	0.28				
Actinomyces odontolyticus		1.14	0.18				
03111000	**Low discrimination**						**03111000**
					GRA		
Actinomyces meyeri		51.11	0.85		99		
Lactobacillus fermentum		34.81	0.93		99		
Prevotella intermedia		6.49	0.68		1		
03111001	**Low discrimination**						**03111001**
					SPO		
Actinomyces meyeri		39.12	0.51	URE 1	1		
Lactobacillus fermentum		26.64	0.59		1		
Clostridium chauvoei		20.10	0.44		99		
03111002	**Good identification**						**03111002**
					COC	GRA	
Lactobacillus fermentum		93.25	0.93		1	99	
Actinomyces meyeri		4.23	0.59		1	99	
Megasphaera elsdenii		1.12	0.50		99	1	
03111003	**Low discrimination**						**03111003**
					SPO		
Lactobacillus fermentum		81.42	0.59	URE 1	1		
Clostridium chauvoei		10.84	0.31		99		
Actinomyces meyeri		3.70	0.25		1		
03111010	**Low discrimination**						**03111010**
					GRA		
Actinomyces meyeri		48.78	0.51	MLZ 1	99		
Lactobacillus fermentum		33.22	0.59		99		
Prevotella intermedia		6.19	0.33		1		
03111012	**Good identification**						**03111012**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Lactobacillus fermentum		90.98	0.59	MLZ 1	1		
Actinomyces meyeri		4.13	0.25		1		
Propionibacterium granulosum		1.85	0.25		90		
03111020	**Low discrimination**						**03111020**
Lactobacillus fermentum		59.17	0.85				
Actinomyces meyeri		35.54	0.70				
Clostridium perfringens		1.82	0.46				
03111021	**Low discrimination**						**03111021**
					SPO		
Lactobacillus fermentum		39.79	0.50	URE 1	1		
Clostridium chauvoei		30.03	0.35		99		
Actinomyces meyeri		23.90	0.36		1		
03111022	**Very good identification**						**03111022**
Lactobacillus fermentum		95.95	0.85				
Actinomyces meyeri		1.78	0.44				
Clostridium perfringens		1.39	0.41				
03111023	**Low discrimination**						**03111023**
					SPO		
Lactobacillus fermentum		81.92	0.50	URE 1	1		
Clostridium chauvoei		10.91	0.22		99		
Actinomyces naeslundii		3.94	0.38		1		
03111030	**Low discrimination**						**03111030**
					CAT		
Lactobacillus fermentum		56.21	0.50	MLZ 1	1		
Actinomyces meyeri		33.76	0.36		1		
Propionibacterium granulosum		3.44	0.25		90		
03111032	**Acceptable identification**						**03111032**
					CAT		
Lactobacillus fermentum		94.81	0.50	MLZ 1	1		
Actinomyces meyeri		1.76	0.10		1		
Propionibacterium granulosum		1.45	0.15		90		
03111040	**Low discrimination**						**03111040**
					COC	GRA	
Megasphaera elsdenii		70.21	0.63	GAL 1	99	1	
Actinomyces meyeri		14.41	0.51		1	99	
Lactobacillus fermentum		9.81	0.59		1	99	
03111042	**Low discrimination**						**03111042**
					COC	GRA	
Lactobacillus fermentum		53.73	0.59	SOR 1	1	99	
Megasphaera elsdenii		42.72	0.47		99	1	
Actinomyces meyeri		2.44	0.25		1	99	
03111060	**Low discrimination**						**03111060**
Lactobacillus fermentum		38.41	0.50	SOR 1			
Actinomyces meyeri		23.07	0.36				
Clostridium perfringens		22.29	0.34				
03111062	**Low discrimination**						**03111062**
					COC	GRA	
Lactobacillus fermentum		76.01	0.50	SOR 1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		20.75	0.29		1	99	
Megasphaera elsdenii		1.83	0.12		99	1	
03111100	**Low discrimination**						**03111100**
					GRA		
Actinomyces meyeri		77.61	0.95		99		
Lactobacillus fermentum		21.56	0.96		99		
Fusobacterium mortiferum		0.49	0.67		1		
03111101	**Good identification**						**03111101**
					SPO		
Clostridium chauvoei		90.73	0.78	MNS 90	99		
Actinomyces meyeri		7.13	0.61		1		
Lactobacillus fermentum		1.98	0.62		1		
03111102	**Low discrimination**						**03111102**
Lactobacillus fermentum		89.23	0.96				
Actinomyces meyeri		9.93	0.69				
Propionibacterium propionicum		0.25	0.52				
03111103	**Low discrimination**						**03111103**
					SPO		
Clostridium chauvoei		87.26	0.65	MNS 90	99		
Lactobacillus fermentum		10.80	0.62		1		
Actinomyces meyeri		1.20	0.35		1		
03111110	**Low discrimination**						**03111110**
					GRA		
Actinomyces meyeri		77.28	0.61	MLZ 1	99		
Lactobacillus fermentum		21.47	0.62		99		
Fusobacterium mortiferum		0.49	0.33		1		
03111111	**Acceptable identification**						**03111111**
					SPO		
Clostridium chauvoei		90.61	0.44	MNS 90	99		
Actinomyces meyeri		7.12	0.27	MLZ 1	1		
Lactobacillus fermentum		1.98	0.28		1		
03111112	**Low discrimination**						**03111112**
Lactobacillus fermentum		88.76	0.62	MLZ 1			
Actinomyces meyeri		9.88	0.35				
Eubacterium saburreum		0.33	0.17				
03111120	**Low discrimination**						**03111120**
					GRA		
Actinomyces meyeri		43.15	0.80	NAG 12	99		
Lactobacillus fermentum		29.31	0.88		99		
Prevotella melaninogenica		23.25	0.67		1		
03111121	**Good identification**						**03111121**
					SPO		
Clostridium chauvoei		92.92	0.70	MNS 90	99		
Actinomyces meyeri		2.99	0.46		1		
Lactobacillus fermentum		2.03	0.54		1		
03111122	**Low discrimination**						**03111122**
					GRA		
Lactobacillus fermentum		78.85	0.88		99		
Prevotella melaninogenica		11.91	0.55		1		
Clostridium perfringens		5.13	0.56		99		

Code Taxon		ID score	T index	tests against		Differentiation tests		Code
03111123	**Low discrimination**							**03111123**
						GRA	SPO	
Clostridium chauvoei		85.72	0.57	MNS	90	99	99	
Lactobacillus fermentum		10.61	0.54			99	1	
Prevotella melaninogenica		1.60	0.21			1	1	
03111130	**Low discrimination**							**03111130**
						GRA		
Actinomyces meyeri		42.77	0.46	MLZ	1	99		
Lactobacillus fermentum		29.04	0.54	NAG	12	99		
Prevotella melaninogenica		23.04	0.33			1		
03111131	**Acceptable identification**							**03111131**
						SPO		
Clostridium chauvoei		92.52	0.35	MNS	90	99		
Actinomyces meyeri		2.98	0.12	MLZ	1	1		
Lactobacillus fermentum		2.02	0.20			1		
03111132	**Low discrimination**							**03111132**
						GRA		
Lactobacillus fermentum		75.96	0.54	MLZ	1	99		
Prevotella melaninogenica		11.48	0.21			1		
Clostridium perfringens		4.94	0.22			99		
03111140	**Low discrimination**							**03111140**
Actinomyces meyeri		73.64	0.61	SOR	1			
Lactobacillus fermentum		20.46	0.62					
Propionibacterium propionicum		3.71	0.49					
03111141	**Low discrimination**							**03111141**
						SPO		
Clostridium chauvoei		86.60	0.44	MNS	90	99		
Actinomyces meyeri		6.81	0.27	SOR	1	1		
Propionibacterium propionicum		3.77	0.33			1		
03111142	**Low discrimination**							**03111142**
Lactobacillus fermentum		70.95	0.62	SOR	1			
Propionibacterium propionicum		19.30	0.52					
Actinomyces meyeri		7.90	0.35					
03111160	**Low discrimination**							**03111160**
Clostridium perfringens		44.22	0.49	MNS	99			
Actinomyces meyeri		24.97	0.46					
Lactobacillus fermentum		16.96	0.54					
03111161	**Low discrimination**							**03111161**
						SPO		
Clostridium chauvoei		86.19	0.35	MNS	90	99		
Clostridium perfringens		4.91	0.15	SOR	1	84		
Actinomyces meyeri		2.77	0.12			1		
03111162	**Low discrimination**							**03111162**
						GRA		
Clostridium perfringens		49.54	0.43	MNS	99	99		
Lactobacillus fermentum		40.37	0.54			99		
Prevotella melaninogenica		6.10	0.21			1		
03111200	**Low discrimination**							**03111200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		45.50	0.58	NIT 99		
Actinomyces meyeri		27.90	0.51			
Lactobacillus fermentum		19.00	0.59			
03111202	**Low discrimination**					**03111202**
Actinomyces odontolyticus		58.02	0.54	NIT 99		
Lactobacillus fermentum		39.53	0.59			
Actinomyces meyeri		1.80	0.25			
03111220	**Low discrimination**					**03111220**
Actinomyces odontolyticus		41.02	0.44	NIT 99		
Lactobacillus fermentum		35.07	0.50	NAG 14		
Actinomyces meyeri		21.06	0.36			
03111222	**Low discrimination**					**03111222**
Lactobacillus fermentum		57.15	0.50	RHA 1		
Actinomyces odontolyticus		40.97	0.41			
Actinomyces meyeri		1.06	0.10			
03111300	**Low discrimination**					**03111300**
Actinomyces meyeri		70.63	0.61	RHA 1		
Lactobacillus fermentum		19.62	0.62			
Actinomyces odontolyticus		9.10	0.49			
03111301	**Acceptable identification**					**03111301**
					SPO	
Clostridium chauvoei		90.04	0.44	MNS 90	99	
Actinomyces meyeri		7.08	0.27	RHA 1	1	
Lactobacillus fermentum		1.97	0.28		1	
03111302	**Low discrimination**					**03111302**
Lactobacillus fermentum		71.39	0.62	RHA 1		
Actinomyces odontolyticus		20.28	0.46			
Actinomyces meyeri		7.95	0.35			
03111320	**Low discrimination**					**03111320**
					GRA	
Actinomyces meyeri		40.51	0.46	RHA 1	99	
Lactobacillus fermentum		27.51	0.54	NAG 12	99	
Prevotella melaninogenica		21.82	0.33		1	
03111321	**Acceptable identification**					**03111321**
					SPO	
Clostridium chauvoei		93.34	0.35	MNS 90	99	
Actinomyces meyeri		3.00	0.12	RHA 1	1	
Lactobacillus fermentum		2.04	0.20		1	
03111322	**Low discrimination**					**03111322**
					GRA	
Lactobacillus fermentum		69.33	0.54	RHA 1	99	
Prevotella melaninogenica		10.48	0.21		1	
Actinomyces odontolyticus		9.62	0.32		99	
03111400	**Low discrimination**					**03111400**
					COC	
Lactobacillus fermentum		72.56	0.93		1	
Actinomyces meyeri		26.63	0.75		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		0.57	0.45		99		
03111401	**Low discrimination**						**03111401**
					COC		
Lactobacillus fermentum		72.33	0.59	URE 1	1		
Actinomyces meyeri		26.55	0.40		1		
Gemella morbillorum		0.57	0.11		99		
03111402	**Very good identification**						**03111402**
					COC	GRA	
Lactobacillus fermentum		98.85	0.93		1	99	
Actinomyces meyeri		1.12	0.49		1	99	
Megasphaera elsdenii		0.01	0.15		99	1	
03111403	**Very good identification**						**03111403**
Lactobacillus fermentum		98.88	0.59	URE 1			
Actinomyces meyeri		1.12	0.15				
03111410	**Low discrimination**						**03111410**
					COC		
Lactobacillus fermentum		72.59	0.59	MLZ 1	1		
Actinomyces meyeri		26.65	0.40		1		
Gemella morbillorum		0.57	0.11		99		
03111412	**Very good identification**						**03111412**
Lactobacillus fermentum		98.62	0.59	MLZ 1			
Actinomyces meyeri		1.12	0.15				
Bifidobacterium longum subsp. longum		0.26	0.07				
03111420	**Low discrimination**						**03111420**
					COC		
Lactobacillus fermentum		86.70	0.85		1		
Actinomyces meyeri		13.02	0.60		1		
Gemella morbillorum		0.23	0.29		99		
03111421	**Low discrimination**						**03111421**
					SPO		
Lactobacillus fermentum		86.37	0.50	URE 1	1		
Actinomyces meyeri		12.97	0.26		1		
Clostridium chauvoei		0.66	0.01		99		
03111422	**Very good identification**						**03111422**
Lactobacillus fermentum		99.51	0.85				
Actinomyces meyeri		0.46	0.34				
Bifidobacterium longum subsp. longum		0.02	0.14				
03111423	**Good identification**						**03111423**
Lactobacillus fermentum		100.0	0.50	URE 1			
03111430	**Low discrimination**						**03111430**
Lactobacillus fermentum		84.61	0.50	MLZ 1			
Actinomyces meyeri		12.70	0.26				
Bifidobacterium longum subsp. longum		2.69	0.18				
03111432	**Good identification**						**03111432**
Lactobacillus fermentum		96.92	0.50	MLZ 1			
Bifidobacterium longum subsp. longum		3.08	0.18				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03111440	**Low discrimination**						**03111440**
					COC	GRA	
Lactobacillus fermentum		69.10	0.59	SOR 1	1	99	
Actinomyces meyeri		25.36	0.40		1	99	
Megasphaera elsdenii		4.99	0.29		99	1	
03111442	**Very good identification**						**03111442**
					COC	GRA	
Lactobacillus fermentum		98.10	0.59	SOR 1	1	99	
Actinomyces meyeri		1.11	0.15		1	99	
Megasphaera elsdenii		0.79	0.12		99	1	
03111460	**Low discrimination**						**03111460**
Lactobacillus fermentum		86.95	0.50	SOR 1			
Actinomyces meyeri		13.06	0.26				
03111462	**Good identification**						**03111462**
Lactobacillus fermentum		100.0	0.50	SOR 1			
03111500	**Low discrimination**						**03111500**
Lactobacillus fermentum		52.60	0.96				
Actinomyces meyeri		47.33	0.85				
Bifidobacterium longum subsp. longum		0.05	0.38				
03111501	**Low discrimination**						**03111501**
					SPO		
Lactobacillus fermentum		42.33	0.62	URE 1	1		
Actinomyces meyeri		38.09	0.51		1		
Clostridium chauvoei		19.58	0.44		99		
03111502	**Very good identification**						**03111502**
Lactobacillus fermentum		97.18	0.96				
Actinomyces meyeri		2.71	0.59				
Bifidobacterium longum subsp. longum		0.09	0.38				
03111503	**Good identification**						**03111503**
					SPO		
Lactobacillus fermentum		90.14	0.62	URE 1	1		
Clostridium chauvoei		7.36	0.31		99		
Actinomyces meyeri		2.51	0.25		1		
03111510	**Low discrimination**						**03111510**
Lactobacillus fermentum		48.53	0.62	MLZ 1			
Actinomyces meyeri		43.67	0.51				
Bifidobacterium longum subsp. longum		7.80	0.42				
03111512	**Low discrimination**						**03111512**
Lactobacillus fermentum		83.19	0.62	MLZ 1			
Bifidobacterium longum subsp. longum		13.37	0.42				
Actinomyces meyeri		2.32	0.25				
03111520	**Low discrimination**						**03111520**
Lactobacillus fermentum		71.99	0.88				
Actinomyces meyeri		26.50	0.70				
Bifidobacterium longum subsp. longum		0.82	0.48				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03111521	**Low discrimination**					**03111521**
					SPO	
Lactobacillus fermentum		54.05	0.54	URE 1	1	
Clostridium chauvoei		25.00	0.35		99	
Actinomyces meyeri		19.90	0.36		1	
03111522	**Very good identification**					**03111522**
Lactobacillus fermentum		97.51	0.88			
Bifidobacterium longum subsp. longum		1.12	0.48			
Actinomyces meyeri		1.11	0.44			
03111523	**Acceptable identification**					**03111523**
					SPO	
Lactobacillus fermentum		90.54	0.54	URE 1	1	
Clostridium chauvoei		7.39	0.22		99	
Bifidobacterium longum subsp. longum		1.04	0.14		1	
03111530	**Low discrimination**					**03111530**
Bifidobacterium longum subsp. longum		58.36	0.52	MNS 87		
Lactobacillus fermentum		30.26	0.54	RAF 97		
Actinomyces meyeri		11.14	0.36			
03111532	**Low discrimination**					**03111532**
Bifidobacterium longum subsp. longum		65.46	0.52	MNS 87		
Lactobacillus fermentum		33.94	0.54	RAF 97		
Actinomyces meyeri		0.39	0.10			
03111540	**Low discrimination**					**03111540**
Lactobacillus fermentum		52.59	0.62	SOR 1		
Actinomyces meyeri		47.32	0.51			
Propionibacterium propionicum		0.10	0.15			
03111542	**Very good identification**					**03111542**
Lactobacillus fermentum		96.90	0.62	SOR 1		
Actinomyces meyeri		2.70	0.25			
Propionibacterium propionicum		0.27	0.18			
03111560	**Low discrimination**					**03111560**
Lactobacillus fermentum		70.73	0.54	SOR 1		
Actinomyces meyeri		26.04	0.36			
Clostridium perfringens		1.86	0.15			
03111562	**Good identification**					**03111562**
Lactobacillus fermentum		96.02	0.54	SOR 1		
Clostridium perfringens		1.19	0.09			
Bifidobacterium longum subsp. longum		1.10	0.14			
03111600	**Low discrimination**					**03111600**
Lactobacillus fermentum		71.48	0.59	RHA 1		
Actinomyces meyeri		26.24	0.40			
Actinomyces odontolyticus		1.73	0.24			
03111602	**Very good identification**					**03111602**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		97.45	0.59	RHA 1				
Actinomyces odontolyticus		1.45	0.20					
Actinomyces meyeri		1.11	0.15					
03111620	**Low discrimination**							**03111620**
Lactobacillus fermentum		86.06	0.50	RHA 1				
Actinomyces meyeri		12.92	0.26					
Actinomyces odontolyticus		1.02	0.10					
03111622	**Good identification**							**03111622**
Lactobacillus fermentum		99.28	0.50	RHA 1				
Actinomyces odontolyticus		0.72	0.07					
03111700	**Low discrimination**							**03111700**
Lactobacillus fermentum		52.51	0.62	RHA 1				
Actinomyces meyeri		47.25	0.51					
Actinomyces odontolyticus		0.25	0.15					
03111702	**Very good identification**							**03111702**
Lactobacillus fermentum		97.02	0.62	RHA 1				
Actinomyces meyeri		2.70	0.25					
Actinomyces odontolyticus		0.28	0.12					
03111720	**Low discrimination**							**03111720**
					GRA			
Lactobacillus fermentum		70.88	0.54	RHA 1	99			
Actinomyces meyeri		26.09	0.36		99			
Bacteroides vulgatus		1.65	0.05		1			
03111722	**Good identification**							**03111722**
Lactobacillus fermentum		97.77	0.54	RHA 1				
Bifidobacterium longum subsp. longum		1.12	0.14					
Actinomyces meyeri		1.11	0.10					
03112000	**Excellent identification**							**03112000**
					COC	CAT	GRA	
Megasphaera elsdenii		99.75	1.00		99	1	1	
Actinomyces meyeri		0.09	0.48		1	1	99	
Propionibacterium granulosum		0.04	0.48		1	90	99	
03112001	**Very good identification**							**03112001**
					COC	GRA		
Megasphaera elsdenii		99.46	0.66	URE 1	99	1		
Propionibacterium propionicum		0.31	0.33		1	99		
Actinomyces meyeri		0.09	0.13		1	99		
03112002	**Very good identification**							**03112002**
					COC	CAT	GRA	
Megasphaera elsdenii		99.46	0.84	BGL 10	99	1	1	
Propionibacterium propionicum		0.37	0.52		1	1	99	
Propionibacterium granulosum		0.10	0.38		1	90	99	
03112003	**Good identification**							**03112003**
					COC	GRA		
Megasphaera elsdenii		96.03	0.50	URE 1	99	1		
Propionibacterium propionicum		3.97	0.36	BGL 10	1	99		
03112010	**Very good identification**							**03112010**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO	CAT	
Megasphaera elsdenii		98.81	0.66	MLZ 1	99	1	1	
Propionibacterium granulosum		0.57	0.33		1	1	90	
Clostridium difficile		0.44	0.19		1	99	1	
03112012	**Good identification**							**03112012**
					COC	CAT	GRA	
Megasphaera elsdenii		98.36	0.50	MLZ 1	99	1	1	
Propionibacterium granulosum		1.27	0.23	BGL 10	1	90	99	
Propionibacterium propionicum		0.37	0.18		1	1	99	
03112020	**Very good identification**							**03112020**
					COC	CAT	GRA	
Megasphaera elsdenii		97.14	0.66	NAG 1	99	1	1	
Actinomyces meyeri		1.21	0.33		1	1	99	
Propionibacterium granulosum		1.02	0.38		1	90	99	
03112022	**Good identification**							**03112022**
					COC	CAT	GRA	
Megasphaera elsdenii		96.37	0.50	NAG 1	99	1	1	
Propionibacterium granulosum		2.28	0.27	BGL 10	1	90	99	
Pseudoramibacter alactolyticus		0.39	-.02		1	1	99	
03112040	**Excellent identification**							**03112040**
					COC	SPO	GRA	
Megasphaera elsdenii		99.93	0.97		99	1	1	
Propionibacterium propionicum		0.04	0.49		1	1	99	
Clostridium botulinum biovar B		0.01	0.24		1	99	99	
03112041	**Very good identification**							**03112041**
					COC	SPO	GRA	
Megasphaera elsdenii		99.50	0.63	URE 1	99	1	1	
Propionibacterium propionicum		0.46	0.33		1	1	99	
Clostridium novyi biovar A		0.04	0.12		1	99	99	
03112042	**Very good identification**							**03112042**
					COC	GRA		
Megasphaera elsdenii		99.43	0.81	BGL 10	99	1		
Propionibacterium propionicum		0.56	0.52		1	99		
Pseudoramibacter alactolyticus		0.01	0.00		1	99		
03112043	**Acceptable identification**							**03112043**
					COC	GRA		
Megasphaera elsdenii		94.16	0.47	URE 1	99	1		
Propionibacterium propionicum		5.84	0.36	BGL 10	1	99		
03112050	**Very good identification**							**03112050**
					COC	SPO	GRA	
Megasphaera elsdenii		99.24	0.63	MLZ 1	99	1	1	
Clostridium botulinum biovar B		0.64	0.18		1	99	99	
Propionibacterium propionicum		0.04	0.15		1	1	99	
03112052	**Good identification**							**03112052**
					COC	GRA		
Megasphaera elsdenii		99.44	0.47	MLZ 1	99	1		
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99		
03112060	**Very good identification**							**03112060**
					COC	SPO	GRA	
Megasphaera elsdenii		99.17	0.63	NAG 1	99	1	1	
Clostridium botulinum biovar B		0.64	0.18		1	99	99	
Pseudoramibacter alactolyticus		0.15	0.04		1	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
03112062	**Good identification**						**03112062**
					COC	GRA	
Megasphaera elsdenii		99.44	0.47	NAG 1	99	1	
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99	
03112100	**Low discrimination**						**03112100**
					COC	GRA	
Propionibacterium propionicum		66.58	0.84	NIT 86	1	99	
Megasphaera elsdenii		24.40	0.66		99	1	
Actinomyces meyeri		8.90	0.58		1	99	
03112101	**Very good identification**						**03112101**
					COC	GRA	
Propionibacterium propionicum		95.34	0.67	NIT 86	1	99	
Megasphaera elsdenii		3.18	0.32	URE 10	99	1	
Actinomyces meyeri		1.16	0.24		1	99	
03112102	**Very good identification**						**03112102**
					COC	GRA	
Propionibacterium propionicum		97.04	0.87	NIT 86	1	99	
Megasphaera elsdenii		2.63	0.50		99	1	
Actinomyces meyeri		0.27	0.32		1	99	
03112103	**Very good identification**						**03112103**
					COC	GRA	
Propionibacterium propionicum		99.68	0.70	NIT 86	1	99	
Megasphaera elsdenii		0.25	0.15	URE 10	99	1	
Actinomyces naeslundii		0.08	0.27		1	99	
03112110	**Low discrimination**						**03112110**
					COC	GRA	
Propionibacterium propionicum		66.66	0.49	NIT 86	1	99	
Megasphaera elsdenii		24.43	0.32	MLZ 1	99	1	
Actinomyces meyeri		8.91	0.24		1	99	
03112112	**Good identification**						**03112112**
					COC	GRA	
Propionibacterium propionicum		97.36	0.52	NIT 86	1	99	
Megasphaera elsdenii		2.64	0.15	MLZ 1	99	1	
03112113	**Good identification**						**03112113**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03112120	**Low discrimination**						**03112120**
					COC	GRA	
Actinomyces meyeri		56.17	0.43	MAN 1	1	99	
Propionibacterium propionicum		31.13	0.49	NAG 12	1	99	
Megasphaera elsdenii		11.41	0.32		99	1	
03112122	**Acceptable identification**						**03112122**
					COC	GRA	
Propionibacterium propionicum		92.40	0.52	NIT 86	1	99	
Actinomyces meyeri		3.44	0.17	NAG 1	1	99	
Megasphaera elsdenii		2.51	0.15		99	1	
03112123	**Acceptable identification**						**03112123**
Propionibacterium propionicum		93.04	0.36	NIT 86			
Actinomyces naeslundii		6.96	0.27	NAG 1			
03112140	**Low discrimination**						**03112140**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		80.28	0.84	NIT 86	1	99		
Megasphaera elsdenii		19.61	0.63		99	1		
Actinomyces meyeri		0.11	0.24		1	99		
03112141	**Very good identification**							**03112141**
					COC	GRA		
Propionibacterium propionicum		97.83	0.67	NIT 86	1	99		
Megasphaera elsdenii		2.17	0.29	URE 10	99	1		
03112142	**Very good identification**							**03112142**
					COC	GRA		
Propionibacterium propionicum		98.22	0.87	NIT 86	1	99		
Megasphaera elsdenii		1.78	0.47		99	1		
03112143	**Very good identification**							**03112143**
					COC	GRA		
Propionibacterium propionicum		99.84	0.70	NIT 86	1	99		
Megasphaera elsdenii		0.16	0.12	URE 10	99	1		
03112150	**Low discrimination**							**03112150**
					COC	GRA		
Propionibacterium propionicum		80.37	0.49	NIT 86	1	99		
Megasphaera elsdenii		19.63	0.29	MLZ 1	99	1		
03112152	**Good identification**							**03112152**
					COC	GRA		
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99		
Megasphaera elsdenii		1.78	0.12	MLZ 1	99	1		
03112153	**Good identification**							**03112153**
Propionibacterium propionicum		100.0	0.36	NIT 86				
03112160	**Low discrimination**							**03112160**
					COC	GRA		
Propionibacterium propionicum		79.21	0.49	NIT 86	1	99		
Megasphaera elsdenii		19.35	0.29	NAG 1	99	1		
Actinomyces meyeri		1.44	0.09		1	99		
03112162	**Good identification**							**03112162**
					COC	GRA		
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99		
Megasphaera elsdenii		1.78	0.12	NAG 1	99	1		
03112163	**Good identification**							**03112163**
Propionibacterium propionicum		100.0	0.36	NIT 86				
03112200	**Very good identification**							**03112200**
					COC	CAT	GRA	
Megasphaera elsdenii		99.39	0.66	RHA 1	99	1	1	
Actinomyces odontolyticus		0.34	0.27		1	10	99	
Propionibacterium granulosum		0.13	0.22		1	90	99	
03112202	**Good identification**							**03112202**
					COC	GRA		
Megasphaera elsdenii		97.53	0.50	RHA 1	99	1		
Actinomyces odontolyticus		1.82	0.23	BGL 10	1	99		
Propionibacterium propionicum		0.37	0.18		1	99		
03112240	**Very good identification**							**03112240**
					COC	GRA		
Megasphaera elsdenii		99.96	0.63	RHA 1	99	1		
Propionibacterium propionicum		0.04	0.15		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03112242	**Good identification**						**03112242**
					COC	GRA	
Megasphaera elsdenii		99.44	0.47	RHA 1	99	1	
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99	
03112300	**Low discrimination**						**03112300**
					COC	GRA	
Propionibacterium propionicum		64.99	0.49	NIT 86	1	99	
Megasphaera elsdenii		23.81	0.32	RHA 1	99	1	
Actinomyces meyeri		8.69	0.24		1	99	
03112302	**Good identification**						**03112302**
					COC	GRA	
Propionibacterium propionicum		95.88	0.52	NIT 86	1	99	
Megasphaera elsdenii		2.60	0.15	RHA 1	99	1	
Actinomyces odontolyticus		1.52	0.15		1	99	
03112303	**Good identification**						**03112303**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03112340	**Low discrimination**						**03112340**
					COC	GRA	
Propionibacterium propionicum		80.37	0.49	NIT 86	1	99	
Megasphaera elsdenii		19.63	0.29	RHA 1	99	1	
03112342	**Good identification**						**03112342**
					COC	GRA	
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99	
Megasphaera elsdenii		1.78	0.12	RHA 1	99	1	
03112343	**Good identification**						**03112343**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03112400	**Very good identification**						**03112400**
					COC	GRA	
Megasphaera elsdenii		97.46	0.66	ARA 1	99	1	
Actinomyces meyeri		2.22	0.37		1	99	
Gemella morbillorum		0.13	0.15		99	99	
03112402	**Good identification**						**03112402**
					COC	GRA	
Megasphaera elsdenii		98.18	0.50	ARA 1	99	1	
Lactobacillus fermentum		0.83	0.24	BGL 10	1	99	
Actinomyces meyeri		0.62	0.11		1	99	
03112440	**Very good identification**						**03112440**
					COC	GRA	
Megasphaera elsdenii		99.92	0.63	ARA 1	99	1	
Propionibacterium propionicum		0.04	0.15		1	99	
Actinomyces meyeri		0.04	0.03		1	99	
03112442	**Good identification**						**03112442**
					COC	GRA	
Megasphaera elsdenii		99.44	0.47	ARA 1	99	1	
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99	
03112500	**Low discrimination**						**03112500**
					COC	GRA	
Actinomyces meyeri		69.94	0.48	MAN 1	1	99	
Propionibacterium propionicum		21.14	0.49		1	99	
Megasphaera elsdenii		7.75	0.32		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03112502	**Low discrimination**						**03112502**
Propionibacterium propionicum		88.30	0.52	NIT 86			
Actinomyces meyeri		6.02	0.22	ARA 1			
Lactobacillus fermentum		3.28	0.28				
03112503	**Good identification**						**03112503**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03112540	**Low discrimination**						**03112540**
					COC	GRA	
Propionibacterium propionicum		78.27	0.49	NIT 86	1	99	
Megasphaera elsdenii		19.12	0.29	ARA 1	99	1	
Actinomyces meyeri		2.62	0.13		1	99	
03112542	**Good identification**						**03112542**
					COC	GRA	
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99	
Megasphaera elsdenii		1.78	0.12	ARA 1	99	1	
03112543	**Good identification**						**03112543**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03113000	**Low discrimination**						**03113000**
					COC	GRA	
Megasphaera elsdenii		75.29	0.66	GAL 1	99	1	
Actinomyces meyeri		10.30	0.51		1	99	
Lactobacillus fermentum		7.01	0.59		1	99	
03113002	**Low discrimination**						**03113002**
					COC	GRA	
Megasphaera elsdenii		42.25	0.50	GAL 1	99	1	
Lactobacillus fermentum		35.42	0.59	BGL 10	1	99	
Propionibacterium propionicum		15.72	0.52		1	99	
03113003	**Low discrimination**						**03113003**
					COC	GRA	
Propionibacterium propionicum		48.08	0.36	NIT 86	1	99	
Actinomyces naeslundii		30.33	0.43	LAC 99	1	99	
Megasphaera elsdenii		11.74	0.15	URE 10	99	1	
03113020	**Low discrimination**						**03113020**
					CAT		
Lactobacillus fermentum		41.52	0.50	MAN 1	1		
Actinomyces meyeri		24.94	0.36		1		
Propionibacterium granulosum		14.04	0.38		90		
03113021	**Low discrimination**						**03113021**
					SPO		
Actinomyces naeslundii		79.69	0.43	MNS 88	1		
Lactobacillus fermentum		8.62	0.16	RAF 92	1		
Clostridium chauvoei		6.51	0.01		99		
03113022	**Low discrimination**						**03113022**
					CAT		
Lactobacillus fermentum		81.34	0.50	MAN 1	1		
Propionibacterium granulosum		6.87	0.27		90		
Actinomyces naeslundii		5.06	0.40		25		
03113023	**Acceptable identification**						**03113023**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		90.24	0.43	MNS 88			
Lactobacillus fermentum		9.76	0.16	RAF 92			
03113040	**Very good identification**						**03113040**
					COC	GRA	
Megasphaera elsdenii		95.60	0.63	GAL 1	99	1	
Propionibacterium propionicum		3.95	0.49		1	99	
Actinomyces meyeri		0.20	0.16		1	99	
03113042	**Low discrimination**						**03113042**
					COC	GRA	
Megasphaera elsdenii		63.47	0.47	GAL 1	99	1	
Propionibacterium propionicum		35.43	0.52	BGL 10	1	99	
Lactobacillus fermentum		0.81	0.24		1	99	
03113043	**Low discrimination**						**03113043**
					COC	GRA	
Propionibacterium propionicum		78.48	0.36	NIT 86	1	99	
Megasphaera elsdenii		12.78	0.12	LAC 99	99	1	
Actinomyces naeslundii		8.74	0.30	URE 10	1	99	
03113100	**Low discrimination**						**03113100**
Propionibacterium propionicum		79.12	0.84	NIT 86			
Actinomyces meyeri		15.86	0.61				
Lactobacillus fermentum		4.41	0.62				
03113101	**Low discrimination**						**03113101**
					SPO		
Propionibacterium propionicum		77.98	0.67	NIT 86	1		
Clostridium chauvoei		18.08	0.44	URE 10	99		
Actinomyces naeslundii		2.12	0.51		1		
03113102	**Very good identification**						**03113102**
Propionibacterium propionicum		95.87	0.87	NIT 86			
Lactobacillus fermentum		3.56	0.62				
Actinomyces meyeri		0.40	0.35				
03113103	**Very good identification**						**03113103**
					SPO		
Propionibacterium propionicum		95.35	0.70	NIT 86	1		
Clostridium chauvoei		2.60	0.31	URE 10	99		
Actinomyces naeslundii		1.73	0.51		1		
03113110	**Low discrimination**						**03113110**
Propionibacterium propionicum		78.95	0.49	NIT 86			
Actinomyces meyeri		15.83	0.27	MLZ 1			
Lactobacillus fermentum		4.40	0.28				
03113112	**Good identification**						**03113112**
Propionibacterium propionicum		95.22	0.52	NIT 86			
Lactobacillus fermentum		3.54	0.28	MLZ 1			
Actinomyces naeslundii		0.67	0.26				
03113113	**Acceptable identification**						**03113113**
Propionibacterium propionicum		91.37	0.36	NIT 86			
Actinomyces naeslundii		8.64	0.29	MLZ 1			
03113120	**Low discrimination**						**03113120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces meyeri		36.26	0.46	MAN 1	99		
Lactobacillus fermentum		24.62	0.54	NAG 12	99		
Prevotella melaninogenica		19.53	0.33		1		
03113121	**Low discrimination**						**03113121**
					SPO		
Clostridium chauvoei		64.34	0.35	MNS 90	99		
Actinomyces naeslundii		22.66	0.51	MAN 1	1		
Propionibacterium propionicum		8.41	0.33		1		
03113122	**Low discrimination**						**03113122**
					GRA		
Lactobacillus fermentum		45.74	0.54	MAN 1	99		
Propionibacterium propionicum		37.33	0.52		99		
Prevotella melaninogenica		6.91	0.21		1		
03113123	**Low discrimination**						**03113123**
					SPO		
Actinomyces naeslundii		47.17	0.51	MNS 88	1		
Propionibacterium propionicum		26.27	0.36	RAF 92	1		
Clostridium chauvoei		23.64	0.22		99		
03113140	**Very good identification**						**03113140**
					COC	GRA	
Propionibacterium propionicum		99.46	0.84	NIT 86	1	99	
Megasphaera elsdenii		0.25	0.29		99	1	
Actinomyces meyeri		0.20	0.27		1	99	
03113141	**Very good identification**						**03113141**
					SPO		
Propionibacterium propionicum		99.29	0.67	NIT 86	1		
Actinomyces naeslundii		0.48	0.38	URE 10	1		
Clostridium chauvoei		0.23	0.09		99		
03113142	**Very good identification**						**03113142**
Propionibacterium propionicum		99.92	0.87	NIT 86			
Lactobacillus fermentum		0.04	0.28				
Actinomyces naeslundii		0.02	0.35				
03113143	**Very good identification**						**03113143**
Propionibacterium propionicum		99.68	0.70	NIT 86			
Actinomyces naeslundii		0.32	0.38	URE 10			
03113150	**Good identification**						**03113150**
Propionibacterium propionicum		100.0	0.49	NIT 86			
03113152	**Good identification**						**03113152**
Propionibacterium propionicum		100.0	0.52	NIT 86			
03113153	**Good identification**						**03113153**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03113160	**Low discrimination**						**03113160**
Propionibacterium propionicum		87.39	0.49	NIT 86			
Clostridium perfringens		4.23	0.15	NAG 1			
Actinomyces naeslundii		3.08	0.35				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
03113162	**Good identification**						**03113162**
Propionibacterium propionicum		95.14	0.52	NIT 86			
Actinomyces naeslundii		2.23	0.35	NAG 1			
Clostridium perfringens		1.45	0.09				
03113163	**Low discrimination**						**03113163**
Propionibacterium propionicum		75.94	0.36	NIT 86			
Actinomyces naeslundii		24.06	0.38	NAG 1			
03113300	**Low discrimination**						**03113300**
Propionibacterium propionicum		78.00	0.49	NIT 86			
Actinomyces meyeri		15.64	0.27	RHA 1			
Lactobacillus fermentum		4.35	0.28				
03113302	**Good identification**						**03113302**
Propionibacterium propionicum		95.45	0.52	NIT 86			
Lactobacillus fermentum		3.54	0.28	RHA 1			
Actinomyces odontolyticus		1.01	0.12				
03113303	**Good identification**						**03113303**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03113340	**Good identification**						**03113340**
Propionibacterium propionicum		100.0	0.49	NIT 86			
03113342	**Good identification**						**03113342**
Propionibacterium propionicum		100.0	0.52	NIT 86			
03113343	**Good identification**						**03113343**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03113400	**Low discrimination**						**03113400**
					COC	GRA	
Lactobacillus fermentum		67.08	0.59	MAN 1	1	99	
Actinomyces meyeri		24.62	0.40		1	99	
Megasphaera elsdenii		7.27	0.32		99	1	
03113402	**Very good identification**						**03113402**
					COC	GRA	
Lactobacillus fermentum		97.29	0.59	MAN 1	1	99	
Megasphaera elsdenii		1.17	0.15		99	1	
Actinomyces meyeri		1.10	0.15		1	99	
03113420	**Low discrimination**						**03113420**
Lactobacillus fermentum		86.95	0.50	MAN 1			
Actinomyces meyeri		13.06	0.26				
03113422	**Good identification**						**03113422**
Lactobacillus fermentum		100.0	0.50	MAN 1			
03113500	**Low discrimination**						**03113500**
Lactobacillus fermentum		48.05	0.62	MAN 1			
Actinomyces meyeri		43.24	0.51				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Propionibacterium propionicum	8.71	0.49				
03113502	**Low discrimination**						**03113502**
	Lactobacillus fermentum	76.40	0.62	MAN 1			
	Propionibacterium propionicum	20.78	0.52				
	Actinomyces meyeri	2.13	0.25				
03113503	**Low discrimination**						**03113503**
	Propionibacterium propionicum	74.95	0.36	NIT 86			
	Lactobacillus fermentum	25.05	0.28	ARA 1			
03113520	**Low discrimination**						**03113520**
	Lactobacillus fermentum	71.78	0.54	MAN 1			
	Actinomyces meyeri	26.43	0.36				
	Bifidobacterium longum subsp. longum	0.82	0.14				
03113522	**Good identification**						**03113522**
	Lactobacillus fermentum	96.99	0.54	MAN 1			
	Bifidobacterium longum subsp. longum	1.11	0.14				
	Actinomyces meyeri	1.10	0.10				
03113540	**Acceptable identification**						**03113540**
	Propionibacterium propionicum	90.43	0.49	NIT 86			
	Lactobacillus fermentum	5.04	0.28	ARA 1			
	Actinomyces meyeri	4.53	0.16				
03113542	**Good identification**						**03113542**
	Propionibacterium propionicum	96.42	0.52	NIT 86			
	Lactobacillus fermentum	3.58	0.28	ARA 1			
03113543	**Good identification**						**03113543**
	Propionibacterium propionicum	100.0	0.36	NIT 86			
03114000	**Good identification**						**03114000**
					COC	GRA	
	Actinomyces meyeri	92.47	0.87		1	99	
	Megasphaera elsdenii	5.28	0.66		99	1	
	Gemella morbillorum	0.71	0.50		99	99	
03114001	**Acceptable identification**						**03114001**
					COC	GRA	
	Actinomyces meyeri	92.32	0.53	URE 1	1	99	
	Megasphaera elsdenii	5.27	0.32		99	1	
	Gemella morbillorum	0.71	0.15		99	99	
03114002	**Low discrimination**						**03114002**
					COC	GRA	
	Actinomyces meyeri	70.98	0.61	BGL 3	1	99	
	Megasphaera elsdenii	14.55	0.50		99	1	
	Lactobacillus fermentum	7.17	0.55		1	99	
03114010	**Acceptable identification**						**03114010**
					COC	GRA	
	Actinomyces meyeri	92.49	0.53	MLZ 1	1	99	
	Megasphaera elsdenii	5.28	0.32		99	1	
	Gemella morbillorum	0.71	0.15		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
03114020	**Very good identification**							**03114020**
					COC			
Actinomyces meyeri		97.61	0.72	NAG 12	1			
Lactobacillus fermentum		0.75	0.46		1			
Gemella morbillorum		0.61	0.33		99			
03114021	**Good identification**							**03114021**
Actinomyces meyeri		99.24	0.38	NAG 12				
Lactobacillus fermentum		0.76	0.12	URE 1				
03114022	**Low discrimination**							**03114022**
Actinomyces meyeri		72.64	0.46	NAG 12				
Lactobacillus fermentum		17.93	0.46	BGL 3				
Actinomyces odontolyticus		8.33	0.34					
03114030	**Good identification**							**03114030**
Actinomyces meyeri		99.24	0.38	MLZ 1				
Lactobacillus fermentum		0.76	0.12	NAG 12				
03114040	**Low discrimination**							**03114040**
					COC	SPO	GRA	
Clostridium glycolicum		70.61	0.66	SUC 1	1	99	99	
Megasphaera elsdenii		22.87	0.63		99	1	1	
Actinomyces meyeri		6.07	0.53		1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03114042	**Low discrimination**							**03114042**
					COC	SPO	GRA	
Megasphaera elsdenii		73.04	0.47	XYL 1	99	1	1	
Clostridium glycolicum		20.50	0.32	BGL 10	1	99	99	
Actinomyces meyeri		5.40	0.27		1	1	99	
03114060	**Low discrimination**							**03114060**
					COC	SPO	GRA	
Actinomyces meyeri		42.98	0.38	NAG 12	1	1	99	
Clostridium glycolicum		37.02	0.32	SOR 1	1	99	99	
Megasphaera elsdenii		11.99	0.29		99	1	1	
03114100	**Excellent identification**							**03114100**
Actinomyces meyeri		99.79	0.97					
Lactobacillus fermentum		0.13	0.58					
Actinomyces odontolyticus		0.04	0.42					
03114101	**Very good identification**							**03114101**
					SPO			
Actinomyces meyeri		99.74	0.63	URE 1	1			
Lactobacillus fermentum		0.13	0.24		1			
Clostridium chauvoei		0.10	0.09		99			
03114102	**Very good identification**							**03114102**
Actinomyces meyeri		95.21	0.71	BGL 3				
Lactobacillus fermentum		3.92	0.58					
Actinomyces odontolyticus		0.72	0.39					
03114103	**Good identification**							**03114103**
Actinomyces meyeri		96.05	0.37	URE 1				
Lactobacillus fermentum		3.96	0.24	BGL 3				
03114110	**Very good identification**							**03114110**
Actinomyces meyeri		99.84	0.63	MLZ 1				
Lactobacillus fermentum		0.13	0.24					
Actinomyces odontolyticus		0.04	0.08					
03114112	**Good identification**							**03114112**
Actinomyces meyeri		96.05	0.37	MLZ 1				
Lactobacillus fermentum		3.96	0.24	BGL 3				
03114120	**Very good identification**							**03114120**
Actinomyces meyeri		99.64	0.82	NAG 12				
Lactobacillus fermentum		0.31	0.50					
Actinomyces odontolyticus		0.05	0.29					
03114121	**Good identification**							**03114121**
					SPO			
Actinomyces meyeri		99.45	0.48	NAG 12	1			
Lactobacillus fermentum		0.31	0.16	URE 1	1			
Clostridium chauvoei		0.24	0.01		99			
03114122	**Good identification**							**03114122**
Actinomyces meyeri		90.11	0.56	NAG 12				
Lactobacillus fermentum		9.07	0.50	BGL 3				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Actinomyces odontolyticus	0.82	0.25					
03114130	**Good identification**							**03114130**
	Actinomyces meyeri	99.69	0.48	MLZ 1				
	Lactobacillus fermentum	0.31	0.16	NAG 12				
03114140	**Very good identification**							**03114140**
					COC	SPO	GRA	
	Actinomyces meyeri	96.07	0.63	SOR 1	1	1	99	
	Clostridium glycolicum	2.82	0.32		1	99	99	
	Megasphaera elsdenii	0.91	0.29		99	1	1	
03114142	**Acceptable identification**							**03114142**
					COC	GRA		
	Actinomyces meyeri	90.58	0.37	SOR 1	1	99		
	Lactobacillus fermentum	3.73	0.24	BGL 3	1	99		
	Megasphaera elsdenii	3.10	0.12		99	1		
03114160	**Good identification**							**03114160**
					SPO			
	Actinomyces meyeri	99.47	0.48	NAG 12	1			
	Lactobacillus fermentum	0.31	0.16	SOR 1	1			
	Clostridium glycolicum	0.22	-.03		99			
03114200	**Low discrimination**							**03114200**
					COC	GRA		
	Actinomyces meyeri	49.87	0.53	RHA 1	1	99		
	Actinomyces odontolyticus	46.44	0.56		1	99		
	Megasphaera elsdenii	2.85	0.32		99	1		
03114202	**Acceptable identification**							**03114202**
					COC	GRA		
	Actinomyces odontolyticus	93.39	0.52	NIT 99	1	99		
	Actinomyces meyeri	5.06	0.27		1	99		
	Megasphaera elsdenii	1.04	0.15		99	1		
03114220	**Low discrimination**							**03114220**
	Actinomyces odontolyticus	52.46	0.42	NIT 99				
	Actinomyces meyeri	47.19	0.38	NAG 14				
	Lactobacillus fermentum	0.36	0.12					
03114222	**Acceptable identification**							**03114222**
	Actinomyces odontolyticus	94.64	0.39	NIT 99				
	Actinomyces meyeri	4.30	0.12	NAG 14				
	Lactobacillus fermentum	1.06	0.12					
03114300	**Good identification**							**03114300**
	Actinomyces meyeri	93.04	0.63	RHA 1				
	Actinomyces odontolyticus	6.84	0.47					
	Lactobacillus fermentum	0.12	0.24					
03114302	**Low discrimination**							**03114302**
	Actinomyces odontolyticus	58.32	0.43	NIT 99				
	Actinomyces meyeri	40.03	0.37					
	Lactobacillus fermentum	1.65	0.24					
03114320	**Acceptable identification**							**03114320**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		91.67	0.48	RHA 1				
Actinomyces odontolyticus		8.05	0.34	NAG 12				
Lactobacillus fermentum		0.29	0.16					
03114400	**Very good identification**							**03114400**
					COC			
Actinomyces meyeri		98.17	0.76		1			
Lactobacillus fermentum		1.23	0.55		1			
Gemella morbillorum		0.34	0.33		99			
03114401	**Good identification**							**03114401**
Actinomyces meyeri		98.77	0.42	URE 1				
Lactobacillus fermentum		1.23	0.20					
03114402	**Low discrimination**							**03114402**
					COC	GRA		
Actinomyces meyeri		70.62	0.51	BGL 3	1	99		
Lactobacillus fermentum		28.52	0.55		1	99		
Megasphaera elsdenii		0.59	0.15		99	1		
03114410	**Good identification**							**03114410**
Actinomyces meyeri		98.77	0.42	MLZ 1				
Lactobacillus fermentum		1.23	0.20					
03114420	**Very good identification**							**03114420**
					COC			
Actinomyces meyeri		96.78	0.62	NAG 12	1			
Lactobacillus fermentum		2.95	0.46		1			
Gemella morbillorum		0.27	0.17		99			
03114422	**Low discrimination**							**03114422**
Actinomyces meyeri		50.33	0.36	NAG 12				
Lactobacillus fermentum		49.68	0.46	BGL 3				
03114440	**Low discrimination**							**03114440**
					COC	SPO	GRA	
Actinomyces meyeri		61.18	0.42	SOR 1	1	1	99	
Clostridium glycolicum		28.74	0.32		1	99	99	
Megasphaera elsdenii		9.31	0.29		99	1	1	
03114500	**Very good identification**							**03114500**
Actinomyces meyeri		99.49	0.87					
Lactobacillus fermentum		0.51	0.58					
Bifidobacterium longum subsp. longum		0.00	0.12					
03114501	**Good identification**							**03114501**
Actinomyces meyeri		99.49	0.53	URE 1				
Lactobacillus fermentum		0.51	0.24					
03114502	**Low discrimination**							**03114502**
Actinomyces meyeri		85.80	0.61	BGL 3				
Lactobacillus fermentum		14.13	0.58					
Bifidobacterium longum subsp. longum		0.07	0.12					
03114510	**Good identification**							**03114510**
Actinomyces meyeri		99.08	0.53	MLZ 1				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		0.51	0.24			
Bifidobacterium longum subsp. longum		0.41	0.16			
03114520	**Very good identification**					**03114520**
Actinomyces meyeri		98.70	0.72	NAG 12		
Lactobacillus fermentum		1.23	0.50			
Bifidobacterium longum subsp. longum		0.07	0.22			
03114521	**Good identification**					**03114521**
Actinomyces meyeri		98.77	0.38	NAG 12		
Lactobacillus fermentum		1.23	0.16	URE 1		
03114522	**Low discrimination**					**03114522**
Actinomyces meyeri		70.12	0.46	NAG 12		
Lactobacillus fermentum		28.23	0.50	BGL 3		
Bifidobacterium longum subsp. longum		1.65	0.22			
03114530	**Low discrimination**					**03114530**
Actinomyces meyeri		88.09	0.38	MLZ 1		
Bifidobacterium longum subsp. longum		10.81	0.26	NAG 12		
Lactobacillus fermentum		1.10	0.16			
03114540	**Good identification**					**03114540**
					SPO	
Actinomyces meyeri		99.38	0.53	SOR 1	1	
Lactobacillus fermentum		0.51	0.24		1	
Clostridium glycolicum		0.12	-.03		99	
03114560	**Good identification**					**03114560**
Actinomyces meyeri		98.77	0.38	NAG 12		
Lactobacillus fermentum		1.23	0.16	SOR 1		
03114600	**Good identification**					**03114600**
Actinomyces meyeri		95.23	0.42	RHA 1		
Actinomyces odontolyticus		3.58	0.22			
Lactobacillus fermentum		1.19	0.20			
03114700	**Good identification**					**03114700**
Actinomyces meyeri		99.20	0.53	RHA 1		
Lactobacillus fermentum		0.51	0.24			
Actinomyces odontolyticus		0.30	0.13			
03114720	**Good identification**					**03114720**
					GRA	
Actinomyces meyeri		96.47	0.38	RHA 1	99	
Bacteroides vulgatus		2.33	-.01	NAG 12	1	
Lactobacillus fermentum		1.20	0.16		99	
03115000	**Low discrimination**					**03115000**
					COC	
Actinomyces meyeri		82.44	0.90		1	
Lactobacillus fermentum		16.99	0.89		1	
Gemella morbillorum		0.23	0.45		99	
03115001	**Low discrimination**					**03115001**
					COC	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		82.40	0.56	URE	1	
Lactobacillus fermentum		16.98	0.55		1	
Gemella morbillorum		0.23	0.11		99	
03115002	**Low discrimination**					**03115002**
Lactobacillus fermentum		86.43	0.89			
Actinomyces meyeri		12.97	0.64			
Actinomyces odontolyticus		0.55	0.44			
03115003	**Low discrimination**					**03115003**
Lactobacillus fermentum		86.47	0.55	URE	1	
Actinomyces meyeri		12.98	0.30			
Actinomyces odontolyticus		0.55	0.10			
03115010	**Low discrimination**					**03115010**
Actinomyces meyeri		82.39	0.56	MLZ	1	
Lactobacillus fermentum		16.98	0.55			
Eubacterium saburreum		0.23	0.19			
03115012	**Low discrimination**					**03115012**
Lactobacillus fermentum		86.47	0.55	MLZ	1	
Actinomyces meyeri		12.98	0.30			
Actinomyces odontolyticus		0.55	0.10			
03115020	**Low discrimination**					**03115020**
Actinomyces meyeri		66.27	0.75	NAG	12	
Lactobacillus fermentum		33.38	0.81			
Actinomyces odontolyticus		0.17	0.34			
03115021	**Low discrimination**					**03115021**
					SPO	
Actinomyces meyeri		66.21	0.41	NAG	12 1	
Lactobacillus fermentum		33.35	0.46	URE	1 1	
Clostridium chauvoei		0.43	0.01		99	
03115022	**Good identification**					**03115022**
Lactobacillus fermentum		93.92	0.81			
Actinomyces meyeri		5.77	0.49			
Actinomyces odontolyticus		0.29	0.31			
03115023	**Acceptable identification**					**03115023**
Lactobacillus fermentum		94.22	0.46	URE	1	
Actinomyces meyeri		5.78	0.15			
03115030	**Low discrimination**					**03115030**
Actinomyces meyeri		66.50	0.41	MLZ	1	
Lactobacillus fermentum		33.50	0.46	NAG	12	
03115032	**Acceptable identification**					**03115032**
Lactobacillus fermentum		94.22	0.46	MLZ	1	
Actinomyces meyeri		5.78	0.15			
03115040	**Low discrimination**					**03115040**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		47.67	0.56	SOR 1	1	
Clostridium glycolicum		41.06	0.50		99	
Lactobacillus fermentum		9.82	0.55		1	
03115042	**Low discrimination**					**03115042**
					SPO	
Lactobacillus fermentum		82.48	0.55	SOR 1	1	
Actinomyces meyeri		12.38	0.30		1	
Clostridium glycolicum		3.48	0.15		99	
03115060	**Low discrimination**					**03115060**
					SPO	
Actinomyces meyeri		63.80	0.41	NAG 12	1	
Lactobacillus fermentum		32.14	0.46	SOR 1	1	
Clostridium glycolicum		4.07	0.15		99	
03115062	**Acceptable identification**					**03115062**
Lactobacillus fermentum		94.22	0.46	SOR 1		
Actinomyces meyeri		5.78	0.15			
03115100	**Good identification**					**03115100**
					GRA	
Actinomyces meyeri		92.21	1.00		99	
Lactobacillus fermentum		7.75	0.92		99	
Fusobacterium mortiferum		0.02	0.47		1	
03115101	**Low discrimination**					**03115101**
					SPO	
Actinomyces meyeri		86.92	0.66	URE 1	1	
Lactobacillus fermentum		7.31	0.58		1	
Clostridium chauvoei		5.75	0.44		99	
03115102	**Low discrimination**					**03115102**
Lactobacillus fermentum		72.99	0.92			
Actinomyces meyeri		26.86	0.74			
Actinomyces odontolyticus		0.09	0.36			
03115103	**Low discrimination**					**03115103**
					SPO	
Lactobacillus fermentum		66.36	0.58	URE 1	1	
Actinomyces meyeri		24.42	0.40		1	
Clostridium chauvoei		9.22	0.31		99	
03115110	**Good identification**					**03115110**
Actinomyces meyeri		92.03	0.66	MLZ 1		
Lactobacillus fermentum		7.74	0.58			
Eubacterium saburreum		0.15	0.26			
03115112	**Low discrimination**					**03115112**
Lactobacillus fermentum		71.29	0.58	MLZ 1		
Actinomyces meyeri		26.23	0.40			
Bifidobacterium adolescentis		1.64	0.25			
03115120	**Low discrimination**					**03115120**
					GRA	
Actinomyces meyeri		82.70	0.85	NAG 12	99	
Lactobacillus fermentum		16.99	0.84		99	
Prevotella melaninogenica		0.23	0.33		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03115121	**Low discrimination**					**03115121**
					SPO	
Actinomyces meyeri		72.99	0.51	NAG 12	1	
Lactobacillus fermentum		15.00	0.50	URE 1	1	
Clostridium chauvoei		11.81	0.35		99	
03115122	**Low discrimination**					**03115122**
					GRA	
Lactobacillus fermentum		86.41	0.84		99	
Actinomyces meyeri		13.01	0.59		99	
Prevotella melaninogenica		0.23	0.21		1	
03115123	**Low discrimination**					**03115123**
					SPO	
Lactobacillus fermentum		77.55	0.50	URE 1	1	
Actinomyces meyeri		11.67	0.25		1	
Clostridium chauvoei		10.78	0.22		99	
03115130	**Low discrimination**					**03115130**
Actinomyces meyeri		81.25	0.51	MLZ 1		
Lactobacillus fermentum		16.69	0.50	NAG 12		
Bifidobacterium longum subsp. longum		1.66	0.26			
03115132	**Low discrimination**					**03115132**
Lactobacillus fermentum		79.05	0.50	MLZ 1		
Actinomyces meyeri		11.90	0.25			
Bifidobacterium longum subsp. longum		7.87	0.26			
03115140	**Good identification**					**03115140**
					SPO	
Actinomyces meyeri		92.02	0.66	SOR 1	1	
Lactobacillus fermentum		7.74	0.58		1	
Clostridium glycolicum		0.20	0.15		99	
03115142	**Low discrimination**					**03115142**
Lactobacillus fermentum		72.73	0.58	SOR 1		
Actinomyces meyeri		26.76	0.40			
Propionibacterium propionicum		0.34	0.18			
03115160	**Low discrimination**					**03115160**
Actinomyces meyeri		82.14	0.51	NAG 12		
Lactobacillus fermentum		16.88	0.50	SOR 1		
Clostridium perfringens		0.76	0.15			
03115162	**Low discrimination**					**03115162**
Lactobacillus fermentum		84.27	0.50	SOR 1		
Actinomyces meyeri		12.69	0.25			
Clostridium perfringens		1.78	0.09			
03115200	**Low discrimination**					**03115200**
Actinomyces meyeri		61.63	0.56	RHA 1		
Actinomyces odontolyticus		25.51	0.53			
Lactobacillus fermentum		12.70	0.55			
03115202	**Low discrimination**					**03115202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		51.70	0.49	NIT 99		
Lactobacillus fermentum		42.00	0.55			
Actinomyces meyeri		6.30	0.30			
03115220	**Low discrimination**					**03115220**
Actinomyces meyeri		50.06	0.41	RHA 1		
Lactobacillus fermentum		25.21	0.46	NAG 12		
Actinomyces odontolyticus		24.73	0.39			
03115222	**Low discrimination**					**03115222**
Lactobacillus fermentum		60.15	0.46	RHA 1		
Actinomyces odontolyticus		36.16	0.36			
Actinomyces meyeri		3.69	0.15			
03115300	**Low discrimination**					**03115300**
Actinomyces meyeri		88.28	0.66	RHA 1		
Lactobacillus fermentum		7.42	0.58			
Actinomyces odontolyticus		2.88	0.44			
03115302	**Low discrimination**					**03115302**
Lactobacillus fermentum		61.00	0.58	RHA 1		
Actinomyces meyeri		22.45	0.40			
Actinomyces odontolyticus		14.53	0.40			
03115320	**Low discrimination**					**03115320**
Actinomyces meyeri		77.06	0.51	RHA 1		
Lactobacillus fermentum		15.83	0.50	NAG 12		
Actinomyces odontolyticus		3.01	0.31			
03115322	**Low discrimination**					**03115322**
Lactobacillus fermentum		76.04	0.50	RHA 1		
Actinomyces meyeri		11.45	0.25			
Actinomyces odontolyticus		8.85	0.27			
03115400	**Low discrimination**					**03115400**
					COC	
Actinomyces meyeri		54.78	0.79		1	
Lactobacillus fermentum		45.15	0.89		1	
Gemella morbillorum		0.07	0.29		99	
03115401	**Low discrimination**					**03115401**
Actinomyces meyeri		54.82	0.45	URE 1		
Lactobacillus fermentum		45.18	0.55			
03115402	**Very good identification**					**03115402**
Lactobacillus fermentum		96.37	0.89			
Actinomyces meyeri		3.62	0.54			
Bifidobacterium longum subsp. longum		0.01	0.12			
03115403	**Good identification**					**03115403**
Lactobacillus fermentum		96.38	0.55	URE 1		
Actinomyces meyeri		3.62	0.19			
03115410	**Low discrimination**					**03115410**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		54.48	0.45	MLZ 1		
Lactobacillus fermentum		44.91	0.55			
Bifidobacterium longum subsp. longum		0.61	0.16			
03115412	**Good identification**					**03115412**
Lactobacillus fermentum		95.14	0.55	MLZ 1		
Actinomyces meyeri		3.57	0.19			
Bifidobacterium longum subsp. longum		1.29	0.16			
03115420	**Low discrimination**					**03115420**
Lactobacillus fermentum		66.77	0.81			
Actinomyces meyeri		33.14	0.65			
Bifidobacterium longum subsp. longum		0.06	0.22			
03115421	**Low discrimination**					**03115421**
Lactobacillus fermentum		66.83	0.46	URE 1		
Actinomyces meyeri		33.17	0.30			
03115422	**Very good identification**					**03115422**
Lactobacillus fermentum		98.40	0.81			
Actinomyces meyeri		1.51	0.39			
Bifidobacterium longum subsp. longum		0.10	0.22			
03115423	**Good identification**					**03115423**
Lactobacillus fermentum		98.49	0.46	URE 1		
Actinomyces meyeri		1.51	0.05			
03115430	**Low discrimination**					**03115430**
Lactobacillus fermentum		60.29	0.46	MLZ 1		
Actinomyces meyeri		29.92	0.30			
Bifidobacterium longum subsp. longum		9.79	0.26			
03115432	**Low discrimination**					**03115432**
Lactobacillus fermentum		84.91	0.46	MLZ 1		
Bifidobacterium longum subsp. longum		13.79	0.26			
Actinomyces meyeri		1.30	0.05			
03115440	**Low discrimination**					**03115440**
					SPO	
Actinomyces meyeri		53.79	0.45	SOR 1	1	
Lactobacillus fermentum		44.34	0.55		1	
Clostridium glycolicum		1.87	0.15		99	
03115442	**Good identification**					**03115442**
Lactobacillus fermentum		96.38	0.55	SOR 1		
Actinomyces meyeri		3.62	0.19			
03115460	**Low discrimination**					**03115460**
Lactobacillus fermentum		66.83	0.46	SOR 1		
Actinomyces meyeri		33.17	0.30			
03115462	**Good identification**					**03115462**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		98.49	0.46	SOR 1		
Actinomyces meyeri		1.51	0.05			
03115500	**Low discrimination**					**03115500**
Actinomyces meyeri		74.75	0.90			
Lactobacillus fermentum		25.13	0.92			
Bifidobacterium longum subsp. longum		0.12	0.46			
03115501	**Low discrimination**					**03115501**
					SPO	
Actinomyces meyeri		74.60	0.56	URE 1	1	
Lactobacillus fermentum		25.08	0.58		1	
Clostridium chauvoei		0.20	0.09		99	
03115502	**Good identification**					**03115502**
Lactobacillus fermentum		91.06	0.92			
Actinomyces meyeri		8.38	0.64			
Bifidobacterium longum subsp. longum		0.44	0.46			
03115503	**Good identification**					**03115503**
Lactobacillus fermentum		91.17	0.58	URE 1		
Actinomyces meyeri		8.39	0.30			
Bifidobacterium longum subsp. longum		0.44	0.12			
03115510	**Low discrimination**					**03115510**
Actinomyces meyeri		62.02	0.56	MLZ 1		
Lactobacillus fermentum		20.85	0.58			
Bifidobacterium longum subsp. longum		17.12	0.50			
03115512	**Low discrimination**					**03115512**
Lactobacillus fermentum		48.37	0.58	MLZ 1		
Bifidobacterium longum subsp. longum		39.72	0.50			
Bifidobacterium adolescentis		7.46	0.39			
03115520	**Low discrimination**					**03115520**
Actinomyces meyeri		53.42	0.75	NAG 12		
Lactobacillus fermentum		43.91	0.84			
Bifidobacterium longum subsp. longum		2.57	0.56			
03115521	**Low discrimination**					**03115521**
Actinomyces meyeri		53.29	0.41	NAG 12		
Lactobacillus fermentum		43.80	0.50	URE 1		
Bifidobacterium longum subsp. longum		2.56	0.22			
03115522	**Good identification**					**03115522**
Lactobacillus fermentum		91.21	0.84			
Bifidobacterium longum subsp. longum		5.33	0.56			
Actinomyces meyeri		3.43	0.49			
03115523	**Acceptable identification**					**03115523**
Lactobacillus fermentum		91.23	0.50	URE 1		
Bifidobacterium longum subsp. longum		5.33	0.22			
Actinomyces meyeri		3.43	0.15			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03115530	**Low discrimination**					**03115530**
Bifidobacterium longum subsp. longum		81.63	0.60	MNS 87		
Actinomyces meyeri		10.08	0.41	RAF 97		
Lactobacillus fermentum		8.29	0.50			
03115532	**Good identification**					**03115532**
Bifidobacterium longum subsp. longum		90.47	0.60	MNS 87		
Lactobacillus fermentum		9.18	0.50	RAF 97		
Actinomyces meyeri		0.35	0.15			
03115540	**Low discrimination**					**03115540**
Actinomyces meyeri		74.75	0.56	SOR 1		
Lactobacillus fermentum		25.13	0.58			
Bifidobacterium longum subsp. longum		0.12	0.12			
03115542	**Low discrimination**					**03115542**
Lactobacillus fermentum		89.89	0.58	SOR 1		
Actinomyces meyeri		8.27	0.30			
Bifidobacterium adolescentis		1.40	0.22			
03115560	**Low discrimination**					**03115560**
Actinomyces meyeri		53.48	0.41	NAG 12		
Lactobacillus fermentum		43.95	0.50	SOR 1		
Bifidobacterium longum subsp. longum		2.57	0.22			
03115562	**Acceptable identification**					**03115562**
Lactobacillus fermentum		91.23	0.50	SOR 1		
Bifidobacterium longum subsp. longum		5.33	0.22			
Actinomyces meyeri		3.43	0.15			
03115600	**Low discrimination**					**03115600**
Actinomyces meyeri		54.32	0.45	RHA 1		
Lactobacillus fermentum		44.77	0.55			
Actinomyces odontolyticus		0.91	0.19			
03115602	**Good identification**					**03115602**
Lactobacillus fermentum		95.24	0.55	RHA 1		
Actinomyces meyeri		3.57	0.19			
Actinomyces odontolyticus		1.18	0.15			
03115620	**Low discrimination**					**03115620**
					GRA	
Lactobacillus fermentum		65.43	0.46	RHA 1	99	
Actinomyces meyeri		32.48	0.30		99	
Bacteroides vulgatus		2.09	-.01		1	
03115622	**Good identification**					**03115622**
Lactobacillus fermentum		98.49	0.46	RHA 1		
Actinomyces meyeri		1.51	0.05			
03115700	**Low discrimination**					**03115700**
Actinomyces meyeri		71.78	0.56	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		24.14	0.58					
Eubacterium contortum		3.71	0.30					
03115702	**Low discrimination**							**03115702**
Lactobacillus fermentum		88.78	0.58	RHA 1				
Actinomyces meyeri		8.17	0.30					
Eubacterium contortum		2.41	0.17					
03115722	**Low discrimination**							**03115722**
Lactobacillus fermentum		87.45	0.50	RHA 1				
Bifidobacterium longum subsp. longum		5.11	0.22					
Actinomyces meyeri		3.29	0.15					
03116000	**Low discrimination**							**03116000**
					COC	GRA		
Megasphaera elsdenii		84.62	0.66	XYL 1	99	1		
Actinomyces meyeri		14.98	0.53		1	99		
Gemella morbillorum		0.12	0.15		99	99		
03116002	**Acceptable identification**							**03116002**
					COC	GRA		
Megasphaera elsdenii		94.10	0.50	XYL 1	99	1		
Actinomyces meyeri		4.64	0.27	BGL 10	1	99		
Lactobacillus fermentum		0.47	0.20		1	99		
03116020	**Low discrimination**							**03116020**
					COC	GRA		
Actinomyces meyeri		70.12	0.38	MAN 1	1	99		
Megasphaera elsdenii		29.34	0.32	NAG 12	99	1		
Lactobacillus fermentum		0.54	0.12		1	99		
03116040	**Very good identification**							**03116040**
					COC	SPO	GRA	
Megasphaera elsdenii		96.69	0.63	XYL 1	99	1	1	
Clostridium glycolicum		3.02	0.32		1	99	99	
Actinomyces meyeri		0.26	0.18		1	1	99	
03116042	**Good identification**							**03116042**
					COC	SPO	GRA	
Megasphaera elsdenii		99.16	0.47	XYL 1	99	1	1	
Propionibacterium propionicum		0.56	0.18	BGL 10	1	1	99	
Clostridium glycolicum		0.28	-.03		1	99	99	
03116100	**Good identification**							**03116100**
					COC	GRA		
Actinomyces meyeri		94.80	0.63	MAN 1	1	99		
Propionibacterium propionicum		3.69	0.49		1	99		
Megasphaera elsdenii		1.35	0.32		99	1		
03116102	**Low discrimination**							**03116102**
					COC	GRA		
Propionibacterium propionicum		63.35	0.52	NIT 86	1	99		
Actinomyces meyeri		33.55	0.37	XYL 1	1	99		
Megasphaera elsdenii		1.72	0.15		99	1		
03116103	**Good identification**							**03116103**
Propionibacterium propionicum		95.41	0.36	NIT 86				
Actinomyces meyeri		4.59	0.03	XYL 1				
03116120	**Good identification**							**03116120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		99.40	0.48	MAN 1			
Lactobacillus fermentum		0.31	0.16	NAG 12			
Propionibacterium propionicum		0.29	0.15				
03116140	**Low discrimination**						**03116140**
					COC	GRA	
Propionibacterium propionicum		66.17	0.49	NIT 86	1	99	
Actinomyces meyeri		17.17	0.29	XYL 1	1	99	
Megasphaera elsdenii		16.16	0.29		99	1	
03116142	**Good identification**						**03116142**
					COC	GRA	
Propionibacterium propionicum		97.71	0.52	NIT 86	1	99	
Megasphaera elsdenii		1.77	0.12	XYL 1	99	1	
Actinomyces meyeri		0.52	0.03		1	99	
03116143	**Good identification**						**03116143**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03116400	**Low discrimination**						**03116400**
					COC	GRA	
Actinomyces meyeri		80.60	0.42	MAN 1	1	99	
Megasphaera elsdenii		18.40	0.32		99	1	
Lactobacillus fermentum		1.01	0.20		1	99	
03116500	**Good identification**						**03116500**
Actinomyces meyeri		99.34	0.53	MAN 1			
Lactobacillus fermentum		0.51	0.24				
Propionibacterium propionicum		0.16	0.15				
03116520	**Good identification**						**03116520**
Actinomyces meyeri		98.77	0.38	MAN 1			
Lactobacillus fermentum		1.23	0.16	NAG 12			
03117000	**Low discrimination**						**03117000**
					COC	GRA	
Actinomyces meyeri		79.80	0.56	MAN 1	1	99	
Lactobacillus fermentum		16.45	0.55		1	99	
Megasphaera elsdenii		3.04	0.32		99	1	
03117002	**Low discrimination**						**03117002**
					COC	GRA	
Lactobacillus fermentum		83.70	0.55	MAN 1	1	99	
Actinomyces meyeri		12.56	0.30		1	99	
Megasphaera elsdenii		1.72	0.15		99	1	
03117020	**Low discrimination**						**03117020**
Actinomyces meyeri		66.50	0.41	MAN 1			
Lactobacillus fermentum		33.50	0.46	NAG 12			
03117022	**Acceptable identification**						**03117022**
Lactobacillus fermentum		94.22	0.46	MAN 1			
Actinomyces meyeri		5.78	0.15				
03117100	**Low discrimination**						**03117100**
Actinomyces meyeri		89.96	0.66	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		7.56	0.58			
Propionibacterium propionicum		2.34	0.49			
03117102	**Low discrimination**					**03117102**
Lactobacillus fermentum		53.43	0.58	MAN 1		
Propionibacterium propionicum		24.75	0.52			
Actinomyces meyeri		19.66	0.40			
03117103	**Low discrimination**					**03117103**
Propionibacterium propionicum		78.83	0.36	NIT 86		
Lactobacillus fermentum		15.47	0.24	XYL 1		
Actinomyces meyeri		5.69	0.06	URE 10		
03117120	**Low discrimination**					**03117120**
					GRA	
Actinomyces meyeri		82.63	0.51	MAN 1	99	
Lactobacillus fermentum		16.98	0.50	NAG 12	99	
Prevotella melaninogenica		0.23	-.01		1	
03117122	**Low discrimination**					**03117122**
Lactobacillus fermentum		85.87	0.50	MAN 1		
Actinomyces meyeri		12.93	0.25			
Propionibacterium propionicum		1.21	0.18			
03117140	**Low discrimination**					**03117140**
Propionibacterium propionicum		69.31	0.49	NIT 86		
Actinomyces meyeri		26.97	0.32	XYL 1		
Lactobacillus fermentum		2.27	0.24			
03117142	**Good identification**					**03117142**
Propionibacterium propionicum		95.17	0.52	NIT 86		
Lactobacillus fermentum		2.08	0.24	XYL 1		
Actinomyces israelii		1.99	0.25			
03117143	**Good identification**					**03117143**
Propionibacterium propionicum		100.0	0.36	NIT 86		
03117400	**Low discrimination**					**03117400**
Actinomyces meyeri		54.68	0.45	MAN 1		
Lactobacillus fermentum		45.07	0.55			
Actinomyces israelii		0.25	0.17			
03117402	**Good identification**					**03117402**
Lactobacillus fermentum		95.62	0.55	MAN 1		
Actinomyces meyeri		3.59	0.19			
Actinomyces israelii		0.80	0.20			
03117420	**Low discrimination**					**03117420**
Lactobacillus fermentum		66.83	0.46	MAN 1		
Actinomyces meyeri		33.17	0.30			
03117422	**Good identification**					**03117422**
Lactobacillus fermentum		98.49	0.46	MAN 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		1.51	0.05				
03117500	**Low discrimination**						**03117500**
Actinomyces meyeri		74.43	0.56	MAN 1			
Lactobacillus fermentum		25.03	0.58				
Actinomyces israelii		0.34	0.27				
03117502	**Low discrimination**						**03117502**
Lactobacillus fermentum		81.62	0.58	MAN 1			
Bifidobacterium adolescentis		8.42	0.36				
Actinomyces meyeri		7.51	0.30				
03117512	**Low discrimination**						**03117512**
Bifidobacterium adolescentis		85.47	0.37	MNS 87			
Lactobacillus fermentum		6.85	0.24	RAF 92			
Bifidobacterium longum subsp. longum		5.62	0.16				
03117520	**Low discrimination**						**03117520**
Actinomyces meyeri		53.48	0.41	MAN 1			
Lactobacillus fermentum		43.95	0.50	NAG 12			
Bifidobacterium longum subsp. longum		2.57	0.22				
03117522	**Acceptable identification**						**03117522**
Lactobacillus fermentum		91.23	0.50	MAN 1			
Bifidobacterium longum subsp. longum		5.33	0.22				
Actinomyces meyeri		3.43	0.15				
03120000	**Low discrimination**						**03120000**
					COC	GRA	
Clostridium botulinum biovar A		84.97	0.64	SUC 7	1	99	
Clostridium sporogenes		8.32	0.45		1	99	
Megasphaera elsdenii		3.06	0.44		99	1	
03120002	**Low discrimination**						**03120002**
					COC	GRA	
Clostridium sporogenes		68.30	0.37	SUC 1	1	99	
Clostridium botulinum biovar A		21.14	0.30		1	99	
Megasphaera elsdenii		8.38	0.28		99	1	
03120040	**Low discrimination**						**03120040**
					COC	GRA	
Clostridium botulinum biovar B		57.73	0.56	MNS 99	1	99	
Clostridium botulinum biovar A		38.78	0.55		1	99	
Megasphaera elsdenii		3.12	0.41		99	1	
03120050	**Good identification**						**03120050**
Clostridium botulinum biovar B		97.80	0.50	MNS 99			
Clostridium botulinum biovar A		1.35	0.21				
Clostridium novyi biovar A		0.75	0.29				
03120060	**Good identification**						**03120060**
Clostridium botulinum biovar B		98.35	0.50	MNS 99			
Clostridium botulinum biovar A		1.36	0.21				
Clostridium novyi biovar A		0.19	0.18				
03120070	**Good identification**						**03120070**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar B		99.61	0.45	MNS 99			
Clostridium novyi biovar A		0.39	0.18				
03120100	**Low discrimination**						**03120100**
					SPO		
Actinomyces meyeri		51.79	0.36	FRU 95	1		
Clostridium botulinum biovar A		39.82	0.30	TRE 1	99		
Clostridium sporogenes		3.90	0.11		99		
03121100	**Low discrimination**						**03121100**
					SPO		
Actinomyces meyeri		76.24	0.39	FRU 95	1		
Clostridium chauvoei		8.86	0.21	TRE 1	99		
Lactobacillus fermentum		4.07	0.28		1		
03121101	**Good identification**						**03121101**
					SPO		
Clostridium chauvoei		99.78	0.55	MNS 90	99		
Actinomyces naeslundii		0.13	0.29	TRE 10	1		
Actinomyces meyeri		0.09	0.05		1		
03121103	**Good identification**						**03121103**
					SPO		
Clostridium chauvoei		99.26	0.42	MNS 90	99		
Actinomyces naeslundii		0.74	0.29	TRE 10	1		
03121121	**Good identification**						**03121121**
					SPO		
Clostridium chauvoei		99.61	0.47	MNS 90	99		
Actinomyces naeslundii		0.39	0.29	TRE 10	1		
03122000	**Low discrimination**						**03122000**
					COC	GRA	
Megasphaera elsdenii		74.59	0.44	FRU 95	99	1	
Clostridium botulinum biovar A		20.92	0.30	TRE 1	1	99	
Clostridium sporogenes		2.05	0.11		1	99	
03122040	**Low discrimination**						**03122040**
					COC	GRA	
Megasphaera elsdenii		75.80	0.41	FRU 95	99	1	
Clostridium botulinum biovar B		14.18	0.21	TRE 1	1	99	
Clostridium botulinum biovar A		9.53	0.21		1	99	
03122100	**Good identification**						**03122100**
					COC	GRA	
Propionibacterium propionicum		96.88	0.48	NIT 86	1	99	
Megasphaera elsdenii		2.28	0.10	FRU 99	99	1	
Actinomyces meyeri		0.83	0.02		1	99	
03122102	**Good identification**						**03122102**
Propionibacterium propionicum		100.0	0.51	NIT 86			
03122140	**Good identification**						**03122140**
					COC	GRA	
Propionibacterium propionicum		98.45	0.48	NIT 86	1	99	
Megasphaera elsdenii		1.55	0.07	FRU 99	99	1	
03122142	**Good identification**						**03122142**
Propionibacterium propionicum		100.0	0.51	NIT 86			
03123100	**Good identification**						**03123100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		96.36	0.48	NIT 86				
Actinomyces naeslundii		2.40	0.31	FRU 99				
Actinomyces meyeri		1.24	0.05					
03123102	**Good identification**							**03123102**
Propionibacterium propionicum		98.37	0.51	NIT 86				
Actinomyces naeslundii		1.63	0.31	FRU 99				
03123140	**Good identification**							**03123140**
Propionibacterium propionicum		100.0	0.48	NIT 86				
03123142	**Good identification**							**03123142**
Propionibacterium propionicum		100.0	0.51	NIT 86				
03124100	**Good identification**							**03124100**
Actinomyces meyeri		100.0	0.41	FRU 95				
03125100	**Acceptable identification**							**03125100**
					COC			
Actinomyces meyeri		94.65	0.44	FRU 95	1			
Atopobium parvulum		3.05	0.16	TRE 1	99			
Lactobacillus fermentum		1.53	0.24		1			
03130000	**Low discrimination**							**03130000**
					COC	SPO	CAT	
Megasphaera elsdenii		67.65	0.66	TRE 1	99	1	1	
Propionibacterium granulosum		11.30	0.58		1	1	90	
Clostridium botulinum biovar A		6.31	0.44		1	99	1	
03130002	**Low discrimination**							**03130002**
					COC	SPO	CAT	
Megasphaera elsdenii		64.88	0.50	TRE 1	99	1	1	
Propionibacterium granulosum		24.38	0.48	BGL 10	1	1	90	
Clostridium sporogenes		4.91	0.24		1	99	1	
03130010	**Low discrimination**							**03130010**
					COC	SPO	CAT	
Propionibacterium granulosum		46.83	0.43	MNS 99	1	1	90	
Clostridium botulinum biovar B		26.59	0.26	MLZ 12	1	99	1	
Megasphaera elsdenii		20.77	0.32		99	1	1	
03130020	**Low discrimination**							**03130020**
					SPO	CAT		
Propionibacterium granulosum		49.99	0.48	MNS 99	1	90		
Clostridium botulinum biovar B		15.48	0.26		99	1		
Actinomyces meyeri		14.89	0.33		1	1		
03130022	**Low discrimination**							**03130022**
					COC	CAT	GRA	
Propionibacterium granulosum		79.72	0.38	MNS 99	1	90	99	
Megasphaera elsdenii		8.57	0.15		99	1	1	
Clostridium perfringens		6.38	0.11		1	1	99	
03130040	**Low discrimination**							**03130040**
					COC	GRA		
Clostridium botulinum biovar B		78.80	0.66	MNS 99	1	99		
Megasphaera elsdenii		20.21	0.63		99	1		
Clostridium botulinum biovar A		0.85	0.35		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03130042	**Low discrimination**							**03130042**
					COC	SPO	CAT	
Megasphaera elsdenii		73.07	0.47	TRE 1	99	1	1	
Clostridium botulinum biovar B		25.90	0.32	BGL 10	1	99	1	
Propionibacterium granulosum		0.42	0.14		1	1	90	
03130050	**Very good identification**							**03130050**
					COC	GRA		
Clostridium botulinum biovar B		99.42	0.61	MNS 99	1	99		
Megasphaera elsdenii		0.52	0.29		99	1		
Clostridium novyi biovar A		0.02	0.12		1	99		
03130060	**Very good identification**							**03130060**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.40	0.61	MNS 99	1	99	1	
Megasphaera elsdenii		0.52	0.29		99	1	1	
Propionibacterium granulosum		0.03	0.14		1	1	90	
03130070	**Good identification**							**03130070**
Clostridium botulinum biovar B		100.0	0.55	MNS 99				
03130100	**Low discrimination**							**03130100**
					GRA			
Actinomyces meyeri		77.29	0.58	TRE 1	99			
Fusobacterium mortiferum		14.34	0.55		1			
Propionibacterium propionicum		4.78	0.48		99			
03130102	**Low discrimination**							**03130102**
					GRA			
Propionibacterium propionicum		59.56	0.51	NIT 86	99			
Actinomyces meyeri		19.86	0.32	MAN 99	99			
Fusobacterium mortiferum		13.24	0.38		1			
03130120	**Low discrimination**							**03130120**
					GRA			
Actinomyces meyeri		76.25	0.43	TRE 1	99			
Clostridium perfringens		19.95	0.32	NAG 12	99			
Fusobacterium mortiferum		1.05	0.21		1			
03130142	**Good identification**							**03130142**
					COC	SPO	GRA	
Propionibacterium propionicum		97.09	0.51	NIT 86	1	1	99	
Megasphaera elsdenii		2.15	0.12	MAN 99	99	1	1	
Clostridium botulinum biovar B		0.76	-.03		1	99	99	
03130400	**Low discrimination**							**03130400**
					COC	CAT	GRA	
Actinomyces meyeri		59.00	0.37	TRE 1	1	1	99	
Megasphaera elsdenii		26.14	0.32		99	1	1	
Propionibacterium granulosum		4.37	0.24		1	90	99	
03130500	**Good identification**							**03130500**
					GRA			
Actinomyces meyeri		97.39	0.48	TRE 1	99			
Lactobacillus fermentum		1.64	0.28		99			
Fusobacterium mortiferum		0.73	0.21		1			
03131000	**Low discrimination**							**03131000**
					CAT			
Propionibacterium granulosum		34.08	0.58	MNS 99	90			
Actinomyces meyeri		27.91	0.51		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		19.01	0.59		1		
03131002	**Low discrimination**						**03131002**
					GRA	CAT	
Lactobacillus fermentum		60.90	0.59	TRE 1	99	1	
Propionibacterium granulosum		27.29	0.48		99	90	
Fusobacterium mortiferum		3.16	0.35		1	5	
03131003	**Low discrimination**						**03131003**
					SPO		
Actinomyces naeslundii		51.40	0.43	MNS 88	1		
Clostridium chauvoei		24.44	0.14	RAF 92	99		
Lactobacillus fermentum		16.69	0.24		1		
03131010	**Low discrimination**						**03131010**
					CAT		
Propionibacterium granulosum		82.78	0.43	MNS 99	90		
Actinomyces meyeri		5.02	0.16	MLZ 12	1		
Lactobacillus jensenii		3.90	0.14		1		
03131020	**Low discrimination**						**03131020**
					CAT		
Clostridium perfringens		64.63	0.51	MNS 99	1		
Propionibacterium granulosum		15.36	0.48	LAC 88	90		
Lactobacillus fermentum		11.43	0.50		1		
03131021	**Low discrimination**						**03131021**
					SPO		
Actinomyces naeslundii		35.35	0.43	MNS 88	1		
Clostridium chauvoei		31.75	0.19	RAF 92	99		
Clostridium perfringens		21.64	0.17		84		
03131022	**Low discrimination**						**03131022**
					CAT		
Clostridium perfringens		64.99	0.45	MNS 99	1		
Lactobacillus fermentum		24.41	0.50	LAC 88	1		
Propionibacterium granulosum		8.21	0.38		90		
03131023	**Low discrimination**						**03131023**
					SPO		
Actinomyces naeslundii		64.32	0.43	MNS 88	1		
Clostridium perfringens		18.53	0.11	RAF 92	84		
Clostridium chauvoei		10.19	0.06		99		
03131060	**Low discrimination**						**03131060**
					CAT	SPO	
Clostridium perfringens		70.21	0.39	MNS 99	1	84	
Clostridium botulinum biovar B		27.13	0.26	LAC 88	1	99	
Propionibacterium granulosum		0.89	0.14		90	1	
03131100	**Low discrimination**						**03131100**
					GRA		
Actinomyces meyeri		55.98	0.61	TRE 1	99		
Lactobacillus fermentum		15.55	0.62		99		
Fusobacterium mortiferum		10.39	0.58		1		
03131101	**Very good identification**						**03131101**
					SPO		
Clostridium chauvoei		97.45	0.61	MNS 90	99		
Actinomyces naeslundii		1.04	0.51	TRE 10	1		
Actinomyces meyeri		0.70	0.27		1		
03131102	**Low discrimination**						**03131102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		58.94	0.62	TRE 1		
Propionibacterium propionicum		13.12	0.51			
Clostridium perfringens		7.27	0.34			
03131103	**Acceptable identification**					**03131103**
					SPO	
Clostridium chauvoei		90.86	0.48	MNS 90	99	
Actinomyces naeslundii		5.49	0.51	TRE 10	1	
Propionibacterium propionicum		2.50	0.35		1	
03131120	**Low discrimination**					**03131120**
Clostridium perfringens		86.93	0.66	MNS 99		
Actinomyces meyeri		5.04	0.46			
Lactobacillus fermentum		3.42	0.54			
03131121	**Acceptable identification**					**03131121**
					SPO	
Clostridium chauvoei		91.83	0.53	MNS 90	99	
Clostridium perfringens		4.64	0.32	TRE 10	84	
Actinomyces naeslundii		2.94	0.51		1	
03131122	**Low discrimination**					**03131122**
					GRA	
Clostridium perfringens		89.62	0.60	MNS 99	99	
Lactobacillus fermentum		7.49	0.54		99	
Prevotella melaninogenica		1.13	0.21		1	
03131123	**Low discrimination**					**03131123**
					SPO	
Clostridium chauvoei		75.34	0.40	MNS 90	99	
Actinomyces naeslundii		13.67	0.51	TRE 10	1	
Clostridium perfringens		10.14	0.26		84	
03131140	**Low discrimination**					**03131140**
Propionibacterium propionicum		55.78	0.48	NIT 86		
Clostridium perfringens		18.77	0.28	MAN 99		
Actinomyces meyeri		13.67	0.27			
03131142	**Low discrimination**					**03131142**
Propionibacterium propionicum		84.77	0.51	NIT 86		
Clostridium perfringens		8.95	0.22	MAN 99		
Lactobacillus fermentum		3.85	0.28			
03131160	**Good identification**					**03131160**
Clostridium perfringens		98.84	0.54	MNS 99		
Actinomyces naeslundii		0.39	0.35			
Actinomyces meyeri		0.30	0.12			
03131162	**Good identification**					**03131162**
Clostridium perfringens		98.45	0.48	MNS 99		
Actinomyces naeslundii		0.83	0.35			
Lactobacillus fermentum		0.44	0.20			
03131163	**Low discrimination**					**03131163**
					SPO	
Actinomyces naeslundii		47.25	0.38	MNS 88	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		37.85	0.14	RAF 92	84	
Clostridium chauvoei		14.91	0.06		99	
03131400	**Low discrimination**					**03131400**
					CAT	
Lactobacillus fermentum		71.55	0.59	TRE 1	1	
Actinomyces meyeri		26.26	0.40		1	
Propionibacterium granulosum		1.30	0.24		90	
03131402	**Very good identification**					**03131402**
					CAT	
Lactobacillus fermentum		98.44	0.59	TRE 1	1	
Actinomyces meyeri		1.12	0.15		1	
Propionibacterium granulosum		0.45	0.14		90	
03131420	**Low discrimination**					**03131420**
Lactobacillus fermentum		81.91	0.50	TRE 1		
Actinomyces meyeri		12.30	0.26			
Clostridium perfringens		4.68	0.17			
03131422	**Good identification**					**03131422**
Lactobacillus fermentum		97.38	0.50	TRE 1		
Clostridium perfringens		2.62	0.11			
03131500	**Low discrimination**					**03131500**
Lactobacillus fermentum		51.53	0.62	TRE 1		
Actinomyces meyeri		46.37	0.51			
Bifidobacterium longum subsp. longum		1.22	0.27			
03131502	**Good identification**					**03131502**
Lactobacillus fermentum		94.14	0.62	TRE 1		
Actinomyces meyeri		2.62	0.25			
Bifidobacterium longum subsp. longum		2.22	0.27			
03131520	**Low discrimination**					**03131520**
Lactobacillus fermentum		52.19	0.54	TRE 1		
Actinomyces meyeri		19.21	0.36			
Bifidobacterium longum subsp. longum		14.78	0.38			
03131522	**Low discrimination**					**03131522**
Lactobacillus fermentum		70.65	0.54	TRE 1		
Bifidobacterium longum subsp. longum		20.01	0.38			
Clostridium perfringens		8.54	0.26			
03131530	**Good identification**					**03131530**
Bifidobacterium longum subsp. longum		97.21	0.42	MNS 87		
Lactobacillus fermentum		2.04	0.20	RAF 97		
Actinomyces meyeri		0.75	0.02			
03131532	**Good identification**					**03131532**
Bifidobacterium longum subsp. longum		97.95	0.42	MNS 87		
Lactobacillus fermentum		2.05	0.20	RAF 97		
03132000	**Low discrimination**					**03132000**
					COC CAT GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		86.85	0.66	TRE 1	99	1	1	
Propionibacterium granulosum		10.94	0.56		1	90	99	
Propionibacterium propionicum		1.96	0.48		1	1	99	
03132002	**Low discrimination**							**03132002**
					COC	CAT	GRA	
Megasphaera elsdenii		62.98	0.50	TRE 1	99	1	1	
Propionibacterium propionicum		19.18	0.51	BGL 10	1	1	99	
Propionibacterium granulosum		17.85	0.46		1	90	99	
03132010	**Low discrimination**							**03132010**
					COC	SPO	CAT	
Propionibacterium granulosum		61.70	0.41	MNS 99	1	1	90	
Megasphaera elsdenii		36.28	0.32	MLZ 12	99	1	1	
Clostridium difficile		1.20	-.01		1	99	1	
03132020	**Low discrimination**							**03132020**
					COC	CAT	GRA	
Propionibacterium granulosum		75.30	0.46	MNS 99	1	90	99	
Megasphaera elsdenii		24.15	0.32		99	1	1	
Propionibacterium propionicum		0.55	0.14		1	1	99	
03132022	**Low discrimination**							**03132022**
					COC	CAT	GRA	
Propionibacterium granulosum		84.32	0.35	MNS 99	1	90	99	
Megasphaera elsdenii		12.02	0.15		99	1	1	
Propionibacterium propionicum		3.66	0.17		1	1	99	
03132040	**Good identification**							**03132040**
					COC	SPO	GRA	
Megasphaera elsdenii		92.98	0.63	TRE 1	99	1	1	
Clostridium botulinum biovar B		3.66	0.32		1	99	99	
Propionibacterium propionicum		3.15	0.48		1	1	99	
03132042	**Low discrimination**							**03132042**
					COC	CAT	GRA	
Megasphaera elsdenii		68.28	0.47	TRE 1	99	1	1	
Propionibacterium propionicum		31.18	0.51	BGL 10	1	1	99	
Propionibacterium granulosum		0.29	0.12		1	90	99	
03132100	**Very good identification**							**03132100**
					COC	GRA		
Propionibacterium propionicum		99.28	0.82	NIT 86	1	99		
Megasphaera elsdenii		0.45	0.32		99	1		
Actinomyces meyeri		0.16	0.24		1	99		
03132101	**Very good identification**							**03132101**
Propionibacterium propionicum		99.73	0.66	NIT 86				
Actinomyces naeslundii		0.27	0.32	URE 10				
03132102	**Very good identification**							**03132102**
					COC	GRA		
Propionibacterium propionicum		99.94	0.85	NIT 86	1	99		
Megasphaera elsdenii		0.03	0.15		99	1		
Actinomyces naeslundii		0.01	0.29		1	99		
03132103	**Very good identification**							**03132103**
Propionibacterium propionicum		99.82	0.69	NIT 86				
Actinomyces naeslundii		0.18	0.32	URE 10				
03132110	**Good identification**							**03132110**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		100.0	0.48	NIT 86				
03132112	**Good identification**							**03132112**
Propionibacterium propionicum		100.0	0.51	NIT 86				
03132120	**Acceptable identification**							**03132120**
Propionibacterium propionicum		94.69	0.48	NIT 86				
Actinomyces meyeri		2.11	0.09	NAG 1				
Actinomyces naeslundii		1.87	0.29					
03132122	**Good identification**							**03132122**
Propionibacterium propionicum		98.70	0.51	NIT 86				
Actinomyces naeslundii		1.30	0.29	NAG 1				
03132140	**Very good identification**							**03132140**
					COC	SPO	GRA	
Propionibacterium propionicum		99.69	0.82	NIT 86	1	1	99	
Megasphaera elsdenii		0.30	0.29		99	1	1	
Clostridium botulinum biovar B		0.01	-.03		1	99	99	
03132141	**Very good identification**							**03132141**
Propionibacterium propionicum		100.0	0.66	NIT 86				
03132142	**Very good identification**							**03132142**
					COC	GRA		
Propionibacterium propionicum		99.98	0.85	NIT 86	1	99		
Megasphaera elsdenii		0.02	0.12		99	1		
03132143	**Very good identification**							**03132143**
Propionibacterium propionicum		100.0	0.69	NIT 86				
03132150	**Good identification**							**03132150**
Propionibacterium propionicum		100.0	0.48	NIT 86				
03132152	**Good identification**							**03132152**
Propionibacterium propionicum		100.0	0.51	NIT 86				
03132160	**Good identification**							**03132160**
Propionibacterium propionicum		100.0	0.48	NIT 86				
03132162	**Good identification**							**03132162**
Propionibacterium propionicum		100.0	0.51	NIT 86				
03132300	**Good identification**							**03132300**
Propionibacterium propionicum		100.0	0.48	NIT 86				
03132302	**Good identification**							**03132302**
Propionibacterium propionicum		100.0	0.51	NIT 86				
03132340	**Good identification**							**03132340**
Propionibacterium propionicum		100.0	0.48	NIT 86				
03132342	**Good identification**							**03132342**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		100.0	0.51	NIT 86				
03132500	**Good identification**							**03132500**
Propionibacterium propionicum		96.08	0.48	NIT 86				
Actinomyces meyeri		3.92	0.13	ARA 1				
03132502	**Good identification**							**03132502**
Propionibacterium propionicum		100.0	0.51	NIT 86				
03132540	**Good identification**							**03132540**
Propionibacterium propionicum		100.0	0.48	NIT 86				
03132542	**Good identification**							**03132542**
Propionibacterium propionicum		100.0	0.51	NIT 86				
03133000	**Low discrimination**							**03133000**
					COC	CAT	GRA	
Propionibacterium granulosum		76.28	0.56	MNS 99	1	90	99	
Propionibacterium propionicum		13.66	0.48		1	1	99	
Megasphaera elsdenii		6.12	0.32		99	1	1	
03133001	**Low discrimination**							**03133001**
					CAT			
Actinomyces naeslundii		59.83	0.48	MNS 88	25			
Propionibacterium propionicum		26.64	0.32	RAF 92	1			
Propionibacterium granulosum		13.53	0.22		90			
03133002	**Low discrimination**							**03133002**
					CAT			
Propionibacterium propionicum		47.56	0.51	NIT 86	1			
Propionibacterium granulosum		44.27	0.46	LAC 99	90			
Actinomyces naeslundii		5.27	0.45		25			
03133003	**Low discrimination**							**03133003**
					CAT			
Actinomyces naeslundii		57.99	0.48	MNS 88	25			
Propionibacterium propionicum		38.74	0.35	RAF 92	1			
Propionibacterium granulosum		3.28	0.12		90			
03133010	**Good identification**							**03133010**
					CAT			
Propionibacterium granulosum		97.59	0.41	MNS 99	90			
Propionibacterium propionicum		1.29	0.14	MLZ 12	1			
Actinomyces naeslundii		1.12	0.23		25			
03133020	**Low discrimination**							**03133020**
					CAT			
Propionibacterium granulosum		83.43	0.46	MNS 99	90			
Actinomyces naeslundii		9.94	0.45		25			
Clostridium perfringens		4.70	0.17		1			
03133021	**Acceptable identification**							**03133021**
Actinomyces naeslundii		94.65	0.48	MNS 88				
Propionibacterium granulosum		5.35	0.12	RAF 92				
03133022	**Low discrimination**							**03133022**
					CAT			
Propionibacterium granulosum		60.03	0.35	MNS 99	90			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		28.60	0.45		25			
Clostridium perfringens		6.37	0.11		1			
03133023	**Good identification**							**03133023**
Actinomyces naeslundii		100.0	0.48	MNS 88				
03133040	**Low discrimination**							**03133040**
					COC	CAT	GRA	
Propionibacterium propionicum		71.63	0.48	NIT 86	1	1	99	
Megasphaera elsdenii		21.39	0.29	LAC 99	99	1	1	
Propionibacterium granulosum		4.04	0.22		1	90	99	
03133042	**Good identification**							**03133042**
					COC	GRA		
Propionibacterium propionicum		95.14	0.51	NIT 86	1	99		
Megasphaera elsdenii		2.10	0.12	LAC 99	99	1		
Actinomyces naeslundii		1.86	0.32		1	99		
03133100	**Very good identification**							**03133100**
Propionibacterium propionicum		99.03	0.82	NIT 86				
Actinomyces naeslundii		0.47	0.53					
Actinomyces meyeri		0.25	0.27					
03133101	**Good identification**							**03133101**
					SPO			
Propionibacterium propionicum		91.24	0.66	NIT 86	1			
Actinomyces naeslundii		5.89	0.56	URE 10	1			
Clostridium chauvoei		2.87	0.27		99			
03133102	**Very good identification**							**03133102**
Propionibacterium propionicum		99.60	0.85	NIT 86				
Actinomyces naeslundii		0.32	0.53					
Lactobacillus fermentum		0.05	0.28					
03133103	**Very good identification**							**03133103**
					SPO			
Propionibacterium propionicum		95.54	0.69	NIT 86	1			
Actinomyces naeslundii		4.11	0.56	URE 10	1			
Clostridium chauvoei		0.35	0.14		99			
03133110	**Good identification**							**03133110**
Propionibacterium propionicum		97.57	0.48	NIT 86				
Actinomyces naeslundii		2.43	0.31	MLZ 1				
03133112	**Good identification**							**03133112**
Propionibacterium propionicum		98.37	0.51	NIT 86				
Actinomyces naeslundii		1.63	0.31	MLZ 1				
03133120	**Low discrimination**							**03133120**
Propionibacterium propionicum		46.76	0.48	NIT 86				
Clostridium perfringens		26.98	0.32	NAG 1				
Actinomyces naeslundii		22.14	0.53					
03133121	**Low discrimination**							**03133121**
					SPO			
Actinomyces naeslundii		75.81	0.56	MNS 88	1			
Clostridium chauvoei		12.32	0.19	RAF 92	99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		11.86	0.32		1	
03133122	**Low discrimination**					**03133122**
Propionibacterium propionicum		66.15	0.51	NIT 86		
Actinomyces naeslundii		20.88	0.53	NAG 1		
Clostridium perfringens		11.97	0.26			
03133123	**Low discrimination**					**03133123**
					SPO	
Actinomyces naeslundii		79.15	0.56	MNS 88	1	
Propionibacterium propionicum		18.58	0.35	RAF 92	1	
Clostridium chauvoei		2.27	0.06		99	
03133140	**Very good identification**					**03133140**
Propionibacterium propionicum		99.91	0.82	NIT 86		
Actinomyces naeslundii		0.08	0.40			
Actinomyces israelii		0.01	0.13			
03133141	**Very good identification**					**03133141**
Propionibacterium propionicum		98.87	0.66	NIT 86		
Actinomyces naeslundii		1.13	0.43	URE 10		
03133142	**Very good identification**					**03133142**
Propionibacterium propionicum		99.94	0.85	NIT 86		
Actinomyces naeslundii		0.06	0.40			
Actinomyces israelii		0.01	0.16			
03133143	**Very good identification**					**03133143**
Propionibacterium propionicum		99.25	0.69	NIT 86		
Actinomyces naeslundii		0.75	0.43	URE 10		
03133150	**Good identification**					**03133150**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03133152	**Good identification**					**03133152**
Propionibacterium propionicum		100.0	0.51	NIT 86		
03133160	**Low discrimination**					**03133160**
Propionibacterium propionicum		83.79	0.48	NIT 86		
Clostridium perfringens		9.21	0.19	NAG 1		
Actinomyces naeslundii		7.00	0.40			
03133161	**Low discrimination**					**03133161**
Actinomyces naeslundii		53.00	0.43	MNS 88		
Propionibacterium propionicum		47.00	0.32	RAF 92		
03133162	**Acceptable identification**					**03133162**
Propionibacterium propionicum		91.73	0.51	NIT 86		
Actinomyces naeslundii		5.11	0.40	NAG 1		
Clostridium perfringens		3.16	0.14			
03133300	**Good identification**					**03133300**
Propionibacterium propionicum		100.0	0.48	NIT 86		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03133302	**Good identification**						**03133302**
Propionibacterium propionicum		100.0	0.51	NIT 86			
03133340	**Good identification**						**03133340**
Propionibacterium propionicum		100.0	0.48	NIT 86			
03133342	**Good identification**						**03133342**
Propionibacterium propionicum		100.0	0.51	NIT 86			
03133500	**Low discrimination**						**03133500**
Propionibacterium propionicum		87.19	0.48	NIT 86			
Lactobacillus fermentum		5.94	0.28	ARA 1			
Actinomyces meyeri		5.34	0.16				
03133502	**Acceptable identification**						**03133502**
Propionibacterium propionicum		91.73	0.51	NIT 86			
Lactobacillus fermentum		4.16	0.28	ARA 1			
Bifidobacterium adolescentis		2.49	0.19				
03133540	**Good identification**						**03133540**
Propionibacterium propionicum		100.0	0.48	NIT 86			
03133542	**Good identification**						**03133542**
Propionibacterium propionicum		99.32	0.51	NIT 86			
Actinomyces israelii		0.68	0.15	ARA 1			
03134000	**Acceptable identification**						**03134000**
					COC	GRA	
Actinomyces meyeri		90.17	0.53	TRE 1	1	99	
Megasphaera elsdenii		5.15	0.32		99	1	
Fusobacterium mortiferum		1.11	0.29		1	1	
03134020	**Good identification**						**03134020**
					CAT		
Actinomyces meyeri		97.55	0.38	TRE 1	1		
Propionibacterium granulosum		1.70	0.14	NAG 12	90		
Lactobacillus fermentum		0.75	0.12		1		
03134100	**Very good identification**						**03134100**
					COC	GRA	
Actinomyces meyeri		98.84	0.63	TRE 1	1	99	
Fusobacterium mortiferum		0.71	0.35		1	1	
Atopobium parvulum		0.25	0.16		99	99	
03134102	**Acceptable identification**						**03134102**
					GRA		
Actinomyces meyeri		91.15	0.37	TRE 1	99		
Lactobacillus fermentum		3.75	0.24	BGL 3	99		
Fusobacterium mortiferum		2.36	0.19		1		
03134120	**Good identification**						**03134120**
					COC		
Actinomyces meyeri		99.36	0.48	TRE 1	1		
Atopobium parvulum		0.33	0.03	NAG 12	99		
Lactobacillus fermentum		0.31	0.16		1		
03134400	**Good identification**						**03134400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		98.77	0.42	TRE 1		
Lactobacillus fermentum		1.23	0.20			
03134500	**Good identification**					**03134500**
Actinomyces meyeri		99.49	0.53	TRE 1		
Lactobacillus fermentum		0.51	0.24			
03134520	**Good identification**					**03134520**
Actinomyces meyeri		97.04	0.38	TRE 1		
Bifidobacterium longum subsp. longum		1.75	0.12	NAG 12		
Lactobacillus fermentum		1.21	0.16			
03135000	**Low discrimination**					**03135000**
					GRA	
Actinomyces meyeri		80.67	0.56	TRE 1	99	
Lactobacillus fermentum		16.62	0.55		99	
Fusobacterium mortiferum		1.00	0.32		1	
03135002	**Low discrimination**					**03135002**
					CAT	
Lactobacillus fermentum		84.50	0.55	TRE 1	1	
Actinomyces meyeri		12.68	0.30		1	
Propionibacterium granulosum		0.65	0.14		90	
03135020	**Low discrimination**					**03135020**
Actinomyces meyeri		63.57	0.41	TRE 1		
Lactobacillus fermentum		32.02	0.46	NAG 12		
Clostridium perfringens		3.12	0.17			
03135022	**Acceptable identification**					**03135022**
Lactobacillus fermentum		90.32	0.46	TRE 1		
Actinomyces meyeri		5.55	0.15			
Clostridium perfringens		4.14	0.11			
03135100	**Low discrimination**					**03135100**
					COC	
Actinomyces meyeri		79.25	0.66	TRE 1	1	
Atopobium parvulum		13.29	0.50		99	
Lactobacillus fermentum		6.66	0.58		1	
03135102	**Low discrimination**					**03135102**
					COC	
Lactobacillus fermentum		56.58	0.58	TRE 1	1	
Actinomyces meyeri		20.82	0.40		1	
Atopobium parvulum		19.92	0.38		99	
03135120	**Low discrimination**					**03135120**
					COC	
Actinomyces meyeri		65.80	0.51	TRE 1	1	
Atopobium parvulum		14.28	0.38	NAG 12	99	
Lactobacillus fermentum		13.52	0.50		1	
03135122	**Low discrimination**					**03135122**
					COC	
Lactobacillus fermentum		62.87	0.50	TRE 1	1	
Clostridium perfringens		12.94	0.26		1	
Atopobium parvulum		11.72	0.25		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03135400	**Low discrimination**					**03135400**
Actinomyces meyeri		54.73	0.45	TRE 1		
Lactobacillus fermentum		45.11	0.55			
Actinomyces israelii		0.17	0.14			
03135402	**Good identification**					**03135402**
Lactobacillus fermentum		95.87	0.55	TRE 1		
Actinomyces meyeri		3.60	0.19			
Actinomyces israelii		0.53	0.17			
03135420	**Low discrimination**					**03135420**
Lactobacillus fermentum		65.78	0.46	TRE 1		
Actinomyces meyeri		32.65	0.30			
Bifidobacterium longum subsp. longum		1.57	0.12			
03135422	**Good identification**					**03135422**
Lactobacillus fermentum		96.23	0.46	TRE 1		
Bifidobacterium longum subsp. longum		2.29	0.12			
Actinomyces meyeri		1.48	0.05			
03135500	**Low discrimination**					**03135500**
Actinomyces meyeri		72.12	0.56	TRE 1		
Lactobacillus fermentum		24.25	0.58			
Bifidobacterium longum subsp. longum		2.92	0.36			
03135502	**Low discrimination**					**03135502**
Lactobacillus fermentum		76.12	0.58	TRE 1		
Bifidobacterium longum subsp. longum		9.18	0.36			
Actinomyces meyeri		7.00	0.30			
03135510	**Low discrimination**					**03135510**
Bifidobacterium longum subsp. longum		83.64	0.39	MNS 87		
Actinomyces meyeri		12.24	0.21	RAF 97		
Lactobacillus fermentum		4.12	0.24			
03135512	**Low discrimination**					**03135512**
Bifidobacterium longum subsp. longum		64.52	0.39	MNS 87		
Bifidobacterium adolescentis		32.30	0.36	RAF 97		
Lactobacillus fermentum		3.18	0.24			
03135520	**Low discrimination**					**03135520**
Bifidobacterium longum subsp. longum		39.28	0.46	MNS 87		
Actinomyces meyeri		33.04	0.41	RAF 97		
Lactobacillus fermentum		27.15	0.50			
03135522	**Low discrimination**					**03135522**
Bifidobacterium longum subsp. longum		57.76	0.46	MNS 87		
Lactobacillus fermentum		39.93	0.50	RAF 97		
Actinomyces meyeri		1.50	0.15			
03135530	**Good identification**					**03135530**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		99.10	0.50	MNS 87		
Actinomyces meyeri		0.50	0.07	RAF 97		
Lactobacillus fermentum		0.41	0.16			
03135532	**Good identification**					**03135532**
					GRA	
Bifidobacterium longum subsp. longum		99.15	0.50	MNS 87	99	
Parabacteroides distasonis		0.44	0.14	RAF 97	1	
Lactobacillus fermentum		0.41	0.16		99	
03136100	**Low discrimination**					**03136100**
Propionibacterium propionicum		75.92	0.48	NIT 86		
Actinomyces meyeri		24.08	0.29	XYL 1		
03136102	**Good identification**					**03136102**
Propionibacterium propionicum		99.35	0.51	NIT 86		
Actinomyces meyeri		0.65	0.03	XYL 1		
03136140	**Good identification**					**03136140**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03136142	**Good identification**					**03136142**
Propionibacterium propionicum		100.0	0.51	NIT 86		
03137100	**Low discrimination**					**03137100**
Propionibacterium propionicum		54.25	0.48	NIT 86		
Actinomyces meyeri		25.81	0.32	XYL 1		
Actinomyces israelii		13.44	0.39			
03137102	**Low discrimination**					**03137102**
Propionibacterium propionicum		75.72	0.51	NIT 86		
Actinomyces israelii		18.76	0.42	XYL 1		
Bifidobacterium adolescentis		2.05	0.19			
03137140	**Acceptable identification**					**03137140**
Propionibacterium propionicum		91.21	0.48	NIT 86		
Actinomyces israelii		8.79	0.32	XYL 1		
03137142	**Acceptable identification**					**03137142**
Propionibacterium propionicum		91.21	0.51	NIT 86		
Actinomyces israelii		8.79	0.35	XYL 1		
03137500	**Low discrimination**					**03137500**
Actinomyces israelii		54.53	0.37	SAL 92		
Actinomyces meyeri		31.99	0.21	CEL 93		
Lactobacillus fermentum		10.76	0.24			
03137502	**Low discrimination**					**03137502**
Actinomyces israelii		48.15	0.40	SAL 92		
Bifidobacterium adolescentis		43.12	0.33	CEL 93		
Lactobacillus fermentum		6.33	0.24			
03140000	**Low discrimination**					**03140000**
					COC GRA SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		34.59	0.41	SUC 7	1	99	99	
Megasphaera elsdenii		30.84	0.44	CEL 1	99	1	1	
Peptostreptococcus anaerobius		13.13	0.29		99	99	1	
03140040	**Low discrimination**							**03140040**
					COC	GRA		
Megasphaera elsdenii		60.93	0.41	FRU 95	99	1		
Clostridium botulinum biovar A		30.62	0.32	CEL 1	1	99		
Clostridium novyi biovar A		4.26	0.29		1	99		
03140100	**Acceptable identification**							**03140100**
					GRA	SPO		
Actinomyces meyeri		90.07	0.36	FRU 95	99	1		
Fusobacterium mortiferum		3.58	0.21	CEL 1	1	1		
Clostridium botulinum biovar A		2.80	0.06		99	99		
03141100	**Low discrimination**							**03141100**
					GRA			
Actinomyces meyeri		87.81	0.39	FRU 95	99			
Lactobacillus fermentum		4.68	0.28	CEL 1	99			
Fusobacterium mortiferum		3.49	0.24		1			
03141101	**Good identification**							**03141101**
					SPO			
Clostridium chauvoei		98.09	0.37	MNS 90	99			
Actinomyces naeslundii		0.97	0.26	CEL 1	1			
Actinomyces meyeri		0.95	0.05		1			
03142000	**Good identification**							**03142000**
					COC	GRA	SPO	
Megasphaera elsdenii		98.46	0.44	FRU 95	99	1	1	
Clostridium botulinum biovar A		1.12	0.06	CEL 1	1	99	99	
Peptostreptococcus anaerobius		0.42	-.06		99	99	1	
03142040	**Good identification**							**03142040**
Megasphaera elsdenii		100.0	0.41	FRU 95				
03144100	**Good identification**							**03144100**
Actinomyces meyeri		100.0	0.41	FRU 95				
03145100	**Good identification**							**03145100**
Actinomyces meyeri		98.41	0.44	FRU 95				
Lactobacillus fermentum		1.59	0.24	CEL 1				
03150000	**Low discrimination**							**03150000**
					COC	GRA		
Megasphaera elsdenii		85.12	0.66	CEL 1	99	1		
Actinomyces meyeri		7.76	0.48		1	99		
Fusobacterium mortiferum		2.76	0.49		1	1		
03150002	**Acceptable identification**							**03150002**
					COC	GRA		
Megasphaera elsdenii		92.09	0.50	CEL 1	99	1		
Fusobacterium mortiferum		2.98	0.33	BGL 10	1	1		
Actinomyces meyeri		2.34	0.22		1	99		
03150040	**Very good identification**							**03150040**
					COC	GRA		
Megasphaera elsdenii		98.16	0.63	CEL 1	99	1		
Clostridium botulinum biovar B		1.29	0.24		1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		0.17	0.11		1	99	
03150042	**Good identification**						**03150042**
Megasphaera elsdenii		100.0	0.47	CEL 1			
03150100	**Low discrimination**						**03150100**
					COC	GRA	
Actinomyces meyeri		80.57	0.58	CEL 1	1	99	
Fusobacterium mortiferum		16.68	0.56		1	1	
Megasphaera elsdenii		2.23	0.32		99	1	
03150120	**Low discrimination**						**03150120**
					GRA		
Actinomyces meyeri		89.18	0.43	CEL 1	99		
Prevotella melaninogenica		8.01	0.17	NAG 12	1		
Fusobacterium mortiferum		1.37	0.21		1		
03150400	**Low discrimination**						**03150400**
					COC	GRA	
Actinomyces meyeri		64.17	0.37	CEL 1	1	99	
Megasphaera elsdenii		28.43	0.32		99	1	
Gemella morbillorum		3.83	0.15		99	99	
03150500	**Good identification**						**03150500**
					GRA		
Actinomyces meyeri		97.54	0.48	CEL 1	99		
Lactobacillus fermentum		1.64	0.28		99		
Fusobacterium mortiferum		0.82	0.21		1		
03151000	**Low discrimination**						**03151000**
					GRA		
Actinomyces meyeri		39.79	0.51	CEL 1	99		
Lactobacillus fermentum		27.10	0.59		99		
Fusobacterium mortiferum		14.12	0.52		1		
03151001	**Low discrimination**						**03151001**
Actinomyces naeslundii		34.68	0.40	MNS 88			
Actinomyces meyeri		24.20	0.16	RAF 92			
Lactobacillus fermentum		16.48	0.24				
03151002	**Low discrimination**						**03151002**
Lactobacillus fermentum		72.45	0.59	CEL 1			
Lactobacillus jensenii		10.58	0.33				
Lactobacillus catenaformis		6.06	0.29				
03151003	**Low discrimination**						**03151003**
Actinomyces naeslundii		67.78	0.40	MNS 88			
Lactobacillus fermentum		32.22	0.24	RAF 92			
03151020	**Low discrimination**						**03151020**
Lactobacillus fermentum		45.48	0.50	CEL 1			
Actinomyces meyeri		27.32	0.36				
Lactobacillus jensenii		7.47	0.26				
03151021	**Low discrimination**						**03151021**
					SPO		
Actinomyces naeslundii		72.82	0.40	MNS 88	1		
Lactobacillus fermentum		11.54	0.16	RAF 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		8.71	0.01		99		
03151022	**Low discrimination**						**03151022**
Lactobacillus fermentum		70.98	0.50	CEL 1			
Lactobacillus jensenii		7.77	0.23				
Lactobacillus catenaformis		4.46	0.18				
03151023	**Low discrimination**						**03151023**
Actinomyces naeslundii		86.32	0.40	MNS 88			
Lactobacillus fermentum		13.68	0.16	RAF 92			
03151100	**Low discrimination**						**03151100**
					GRA		
Actinomyces meyeri		65.24	0.61	CEL 1	99		
Lactobacillus fermentum		18.13	0.62		99		
Fusobacterium mortiferum		13.51	0.59		1		
03151101	**Low discrimination**						**03151101**
					SPO		
Clostridium chauvoei		83.55	0.44	MNS 90	99		
Actinomyces naeslundii		6.70	0.48	CEL 1	1		
Actinomyces meyeri		6.57	0.27		1		
03151102	**Low discrimination**						**03151102**
					GRA		
Lactobacillus fermentum		72.83	0.62	CEL 1	99		
Actinomyces meyeri		8.11	0.35		99		
Fusobacterium mortiferum		6.03	0.42		1		
03151120	**Low discrimination**						**03151120**
					GRA		
Prevotella melaninogenica		72.61	0.51	MNS 90	1		
Actinomyces meyeri		12.25	0.46	RAF 90	99		
Lactobacillus fermentum		8.32	0.54	CEL 10	99		
03151121	**Low discrimination**						**03151121**
					GRA	SPO	
Clostridium chauvoei		66.78	0.35	MNS 90	99	99	
Actinomyces naeslundii		16.06	0.48	CEL 1	99	1	
Prevotella melaninogenica		12.73	0.17		1	1	
03151122	**Low discrimination**						**03151122**
					GRA		
Prevotella melaninogenica		44.81	0.39	MNS 90	1		
Lactobacillus fermentum		26.96	0.54	RAF 90	99		
Prevotella oralis		9.33	0.31	CEL 10	1		
03151123	**Low discrimination**						**03151123**
					GRA	SPO	
Actinomyces naeslundii		50.61	0.48	MNS 88	99	1	
Clostridium chauvoei		37.15	0.22	RAF 92	99	99	
Prevotella melaninogenica		7.64	0.05		1	1	
03151400	**Low discrimination**						**03151400**
					COC		
Lactobacillus fermentum		72.34	0.59	CEL 1	1		
Actinomyces meyeri		26.55	0.40		1		
Gemella morbillorum		0.57	0.11		99		
03151402	**Very good identification**						**03151402**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		98.88	0.59	CEL 1				
Actinomyces meyeri		1.12	0.15					
03151420	**Low discrimination**							**03151420**
Lactobacillus fermentum		86.95	0.50	CEL 1				
Actinomyces meyeri		13.06	0.26					
03151422	**Good identification**							**03151422**
Lactobacillus fermentum		100.0	0.50	CEL 1				
03151500	**Low discrimination**							**03151500**
					GRA			
Lactobacillus fermentum		52.38	0.62	CEL 1	99			
Actinomyces meyeri		47.14	0.51		99			
Fusobacterium mortiferum		0.39	0.24		1			
03151502	**Good identification**							**03151502**
Lactobacillus fermentum		94.90	0.62	CEL 1				
Actinomyces meyeri		2.64	0.25					
Bifidobacterium adolescentis		2.23	0.29					
03151520	**Low discrimination**							**03151520**
					GRA			
Lactobacillus fermentum		67.87	0.54	CEL 1	99			
Actinomyces meyeri		24.98	0.36		99			
Prevotella melaninogenica		5.98	0.17		1			
03151522	**Acceptable identification**							**03151522**
					GRA	CAT		
Lactobacillus fermentum		94.91	0.54	CEL 1	99	1		
Prevotella melaninogenica		1.59	0.05		1	1		
Capnocytophaga ochracea		1.33	0.07		1	90		
03152000	**Very good identification**							**03152000**
					COC	CAT	GRA	
Megasphaera elsdenii		99.78	0.66	CEL 1	99	1	1	
Actinomyces meyeri		0.09	0.13		1	1	99	
Propionibacterium granulosum		0.04	0.14		1	90	99	
03152002	**Good identification**							**03152002**
					COC	GRA		
Megasphaera elsdenii		99.63	0.50	CEL 1	99	1		
Propionibacterium propionicum		0.38	0.18	BGL 10	1	99		
03152040	**Very good identification**							**03152040**
					COC	GRA		
Megasphaera elsdenii		99.96	0.63	CEL 1	99	1		
Propionibacterium propionicum		0.04	0.15		1	99		
03152042	**Good identification**							**03152042**
					COC	GRA		
Megasphaera elsdenii		99.44	0.47	CEL 1	99	1		
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99		
03152100	**Low discrimination**							**03152100**
					COC	GRA		
Propionibacterium propionicum		64.99	0.49	NIT 86	1	99		
Megasphaera elsdenii		23.81	0.32	CEL 1	99	1		
Actinomyces meyeri		8.69	0.24		1	99		
03152102	**Good identification**							**03152102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Propionibacterium propionicum		96.67	0.52	NIT 86	1	99	
Megasphaera elsdenii		2.62	0.15	CEL 1	99	1	
Actinomyces naeslundii		0.71	0.26		1	99	
03152103	**Acceptable identification**						**03152103**
Propionibacterium propionicum		90.98	0.36	NIT 86			
Actinomyces naeslundii		9.02	0.29	CEL 1			
03152140	**Low discrimination**						**03152140**
					COC	GRA	
Propionibacterium propionicum		80.37	0.49	NIT 86	1	99	
Megasphaera elsdenii		19.63	0.29	CEL 1	99	1	
03152142	**Good identification**						**03152142**
					COC	GRA	
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99	
Megasphaera elsdenii		1.78	0.12	CEL 1	99	1	
03152143	**Good identification**						**03152143**
Propionibacterium propionicum		100.0	0.36	NIT 86			
03153001	**Good identification**						**03153001**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03153002	**Low discrimination**						**03153002**
					COC	GRA	
Actinomyces naeslundii		45.05	0.42	MNS 88	1	99	
Megasphaera elsdenii		19.74	0.15	RAF 92	99	1	
Lactobacillus fermentum		16.55	0.24		1	99	
03153003	**Good identification**						**03153003**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03153020	**Low discrimination**						**03153020**
Actinomyces naeslundii		83.62	0.42	MNS 88			
Lactobacillus fermentum		10.24	0.16	RAF 92			
Actinomyces meyeri		6.15	0.02				
03153021	**Good identification**						**03153021**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03153022	**Low discrimination**						**03153022**
Actinomyces naeslundii		89.09	0.42	MNS 88			
Lactobacillus fermentum		10.91	0.16	RAF 92			
03153023	**Good identification**						**03153023**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03153100	**Low discrimination**						**03153100**
Propionibacterium propionicum		61.56	0.49	NIT 86			
Actinomyces naeslundii		16.28	0.50	CEL 1			
Actinomyces meyeri		12.34	0.27				
03153101	**Low discrimination**						**03153101**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		74.35	0.53	MNS 88	1	
Propionibacterium propionicum		20.82	0.33	RAF 92	1	
Clostridium chauvoei		4.83	0.09		99	
03153102	**Low discrimination**					**03153102**
Propionibacterium propionicum		77.35	0.52	NIT 86		
Actinomyces naeslundii		13.64	0.50	CEL 1		
Actinomyces israelii		4.82	0.33			
03153103	**Low discrimination**					**03153103**
Actinomyces naeslundii		70.42	0.53	MNS 88		
Propionibacterium propionicum		29.58	0.36	RAF 92		
03153120	**Low discrimination**					**03153120**
					GRA	
Actinomyces naeslundii		53.64	0.50	MNS 88	99	
Prevotella melaninogenica		32.85	0.17	RAF 92	1	
Actinomyces meyeri		5.54	0.12		99	
03153121	**Good identification**					**03153121**
					SPO	
Actinomyces naeslundii		97.88	0.53	MNS 88	1	
Clostridium chauvoei		2.12	0.01	RAF 92	99	
03153122	**Low discrimination**					**03153122**
					GRA	
Actinomyces naeslundii		73.77	0.50	MNS 88	99	
Prevotella melaninogenica		8.61	0.05	RAF 92	1	
Lactobacillus fermentum		5.18	0.20		99	
03153123	**Good identification**					**03153123**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03153140	**Acceptable identification**					**03153140**
Propionibacterium propionicum		93.38	0.49	NIT 86		
Actinomyces naeslundii		4.36	0.37	CEL 1		
Actinomyces israelii		2.26	0.23			
03153142	**Acceptable identification**					**03153142**
Propionibacterium propionicum		94.76	0.52	NIT 86		
Actinomyces naeslundii		2.95	0.37	CEL 1		
Actinomyces israelii		2.29	0.26			
03153143	**Low discrimination**					**03153143**
Propionibacterium propionicum		70.42	0.36	NIT 86		
Actinomyces naeslundii		29.58	0.40	CEL 1		
03153160	**Low discrimination**					**03153160**
Actinomyces naeslundii		82.21	0.37	MNS 88		
Propionibacterium propionicum		17.79	0.15	RAF 92		
03153161	**Good identification**					**03153161**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03153162	**Low discrimination**					**03153162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		75.49	0.37	MNS 88			
Propionibacterium propionicum		24.51	0.18	RAF 92			
03153163	**Good identification**						**03153163**
Actinomyces naeslundii		100.0	0.40	MNS 88			
03154000	**Acceptable identification**						**03154000**
					COC	GRA	
Actinomyces meyeri		91.52	0.53	CEL 1	1	99	
Megasphaera elsdenii		5.22	0.32		99	1	
Fusobacterium mortiferum		1.26	0.30		1	1	
03154020	**Good identification**						**03154020**
Actinomyces meyeri		99.24	0.38	CEL 1			
Lactobacillus fermentum		0.76	0.12	NAG 12			
03154100	**Very good identification**						**03154100**
					GRA		
Actinomyces meyeri		99.04	0.63	CEL 1	99		
Fusobacterium mortiferum		0.80	0.36		1		
Lactobacillus fermentum		0.13	0.24		99		
03154102	**Acceptable identification**						**03154102**
					GRA		
Actinomyces meyeri		93.46	0.37	CEL 1	99		
Lactobacillus fermentum		3.85	0.24	BGL 3	99		
Fusobacterium mortiferum		2.70	0.20		1		
03154120	**Good identification**						**03154120**
Actinomyces meyeri		99.69	0.48	CEL 1			
Lactobacillus fermentum		0.31	0.16	NAG 12			
03154400	**Good identification**						**03154400**
Actinomyces meyeri		98.77	0.42	CEL 1			
Lactobacillus fermentum		1.23	0.20				
03154500	**Good identification**						**03154500**
Actinomyces meyeri		99.49	0.53	CEL 1			
Lactobacillus fermentum		0.51	0.24				
03154520	**Good identification**						**03154520**
Actinomyces meyeri		98.77	0.38	CEL 1			
Lactobacillus fermentum		1.23	0.16	NAG 12			
03155000	**Low discrimination**						**03155000**
					GRA		
Actinomyces meyeri		81.38	0.56	CEL 1	99		
Lactobacillus fermentum		16.77	0.55		99		
Fusobacterium mortiferum		1.12	0.33		1		
03155002	**Low discrimination**						**03155002**
Lactobacillus fermentum		84.19	0.55	CEL 1			
Actinomyces meyeri		12.64	0.30				
Actinomyces israelii		2.01	0.28				
03155020	**Low discrimination**						**03155020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		66.50	0.41	CEL 1		
Lactobacillus fermentum		33.50	0.46	NAG 12		
03155022	**Acceptable identification**					**03155022**
Lactobacillus fermentum		94.22	0.46	CEL 1		
Actinomyces meyeri		5.78	0.15			
03155100	**Good identification**					**03155100**
					GRA	
Actinomyces meyeri		90.97	0.66	CEL 1	99	
Lactobacillus fermentum		7.65	0.58		99	
Fusobacterium mortiferum		0.73	0.39		1	
03155102	**Low discrimination**					**03155102**
Lactobacillus fermentum		67.32	0.58	CEL 1		
Actinomyces meyeri		24.77	0.40			
Actinomyces israelii		3.94	0.38			
03155120	**Low discrimination**					**03155120**
					GRA	
Actinomyces meyeri		80.32	0.51	CEL 1	99	
Lactobacillus fermentum		16.50	0.50	NAG 12	99	
Prevotella melaninogenica		2.48	0.17		1	
03155122	**Low discrimination**					**03155122**
					GRA	
Lactobacillus fermentum		76.52	0.50	CEL 1	99	
Actinomyces meyeri		11.52	0.25		99	
Prevotella oralis		5.57	0.15		1	
03155400	**Low discrimination**					**03155400**
Actinomyces meyeri		54.50	0.45	CEL 1		
Lactobacillus fermentum		44.92	0.55			
Actinomyces israelii		0.59	0.23			
03155402	**Acceptable identification**					**03155402**
Lactobacillus fermentum		94.21	0.55	CEL 1		
Actinomyces meyeri		3.54	0.19			
Actinomyces israelii		1.84	0.26			
03155420	**Low discrimination**					**03155420**
Lactobacillus fermentum		66.83	0.46	CEL 1		
Actinomyces meyeri		33.17	0.30			
03155422	**Good identification**					**03155422**
Lactobacillus fermentum		98.49	0.46	CEL 1		
Actinomyces meyeri		1.51	0.05			
03155500	**Low discrimination**					**03155500**
Actinomyces meyeri		74.10	0.56	CEL 1		
Lactobacillus fermentum		24.91	0.58			
Actinomyces israelii		0.80	0.34			
03155502	**Low discrimination**					**03155502**
Lactobacillus fermentum		70.78	0.58	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		18.97	0.43			
Actinomyces meyeri		6.51	0.30			
03155512	**Acceptable identification**					**03155512**
Bifidobacterium adolescentis		92.77	0.45	MNS 87		
Lactobacillus fermentum		2.86	0.24	RAF 92		
Bifidobacterium longum subsp. longum		2.35	0.16			
03155520	**Low discrimination**					**03155520**
Actinomyces meyeri		53.21	0.41	CEL 1		
Lactobacillus fermentum		43.73	0.50	NAG 12		
Bifidobacterium longum subsp. longum		2.56	0.22			
03155522	**Low discrimination**					**03155522**
Lactobacillus fermentum		89.82	0.50	CEL 1		
Bifidobacterium longum subsp. longum		5.25	0.22			
Actinomyces meyeri		3.38	0.15			
03157000	**Low discrimination**					**03157000**
Actinomyces israelii		60.40	0.38	SAL 92		
Actinomyces meyeri		32.84	0.21			
Lactobacillus fermentum		6.77	0.20			
03157002	**Acceptable identification**					**03157002**
Actinomyces israelii		93.05	0.41	SAL 92		
Lactobacillus fermentum		6.95	0.20			
03157100	**Low discrimination**					**03157100**
Actinomyces israelii		62.37	0.48	SAL 92		
Actinomyces meyeri		33.90	0.32			
Lactobacillus fermentum		2.85	0.24			
03157102	**Low discrimination**					**03157102**
Actinomyces israelii		86.61	0.51	SAL 92		
Bifidobacterium adolescentis		8.56	0.27			
Lactobacillus fermentum		2.64	0.24			
03157112	**Low discrimination**					**03157112**
					SPO	
Actinomyces israelii		54.02	0.37	SAL 92	1	
Bifidobacterium adolescentis		43.66	0.29	MLZ 13	1	
Clostridium tertium		2.33	-.01		99	
03157140	**Good identification**					**03157140**
Actinomyces israelii		96.50	0.41	SAL 92		
Propionibacterium propionicum		3.50	0.15			
03157142	**Acceptable identification**					**03157142**
Actinomyces israelii		93.66	0.44	SAL 92		
Propionibacterium propionicum		3.40	0.18			
Bifidobacterium adolescentis		2.94	0.12			
03157400	**Low discrimination**					**03157400**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		76.74	0.36	SAL 92			
Actinomyces meyeri		12.75	0.11				
Lactobacillus fermentum		10.51	0.20				
03157402	**Low discrimination**						**03157402**
Actinomyces israelii		88.97	0.39	SAL 92			
Lactobacillus fermentum		8.12	0.20				
Bifidobacterium adolescentis		2.91	0.07				
03157500	**Low discrimination**						**03157500**
Actinomyces israelii		81.02	0.46	SAL 92			
Actinomyces meyeri		13.46	0.21				
Lactobacillus fermentum		4.53	0.24				
03157502	**Low discrimination**						**03157502**
Actinomyces israelii		54.19	0.49	SAL 92			
Bifidobacterium adolescentis		43.80	0.42				
Lactobacillus fermentum		2.02	0.24				
03157512	**Low discrimination**						**03157512**
Bifidobacterium adolescentis		86.86	0.43	MNS 87			
Actinomyces israelii		13.14	0.35	RAF 92			
03157540	**Good identification**						**03157540**
Actinomyces israelii		100.0	0.39	SAL 92			
03157542	**Low discrimination**						**03157542**
Actinomyces israelii		79.56	0.42	SAL 92			
Bifidobacterium adolescentis		20.44	0.26				
03163101	**Good identification**						**03163101**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03163103	**Good identification**						**03163103**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03163121	**Good identification**						**03163121**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03163123	**Good identification**						**03163123**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03170100	**Low discrimination**						**03170100**
					COC	GRA	
Fusobacterium mortiferum		77.69	0.47	MLT 11	1	1	
Actinomyces meyeri		12.69	0.24		1	99	
Atopobium parvulum		7.79	0.18		99	99	
03171000	**Low discrimination**						**03171000**
					GRA		
Lactobacillus jensenii		71.96	0.44	MNS 99	99		
Fusobacterium mortiferum		17.05	0.43	SAL 95	1		
Actinomyces naeslundii		3.01	0.42		99		
03171001	**Low discrimination**						**03171001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		86.14	0.45	MNS 88			
Lactobacillus jensenii		13.86	0.10	RAF 92			
03171002	**Low discrimination**						**03171002**
					GRA		
Lactobacillus jensenii		83.83	0.41	MNS 99	99		
Actinomyces naeslundii		5.26	0.42	SAL 95	99		
Fusobacterium mortiferum		3.31	0.27		1		
03171003	**Acceptable identification**						**03171003**
Actinomyces naeslundii		90.31	0.45	MNS 88			
Lactobacillus jensenii		9.69	0.07	RAF 92			
03171010	**Good identification**						**03171010**
					CAT		
Lactobacillus jensenii		98.89	0.36	MNS 99	1		
Propionibacterium granulosum		1.12	0.09	SAL 95	90		
03171021	**Good identification**						**03171021**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03171023	**Good identification**						**03171023**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03171100	**Low discrimination**						**03171100**
					COC	GRA	
Atopobium parvulum		81.20	0.52	MNS 99	99	99	
Fusobacterium mortiferum		12.28	0.50		1	1	
Actinomyces naeslundii		2.65	0.50		1	99	
03171101	**Low discrimination**						**03171101**
					COC	SPO	
Actinomyces naeslundii		51.23	0.53	MNS 88	1	1	
Clostridium chauvoei		36.58	0.27	RAF 92	1	99	
Atopobium parvulum		10.59	0.18		99	1	
03171102	**Low discrimination**						**03171102**
					COC		
Atopobium parvulum		54.08	0.39	MNS 99	99		
Lactobacillus acidophilus		23.76	0.32		1		
Actinomyces naeslundii		9.98	0.50		1		
03171103	**Low discrimination**						**03171103**
					COC	SPO	
Actinomyces naeslundii		86.02	0.53	MNS 88	1	1	
Clostridium chauvoei		10.84	0.14	RAF 92	1	99	
Atopobium parvulum		3.14	0.05		99	1	
03171120	**Low discrimination**						**03171120**
					COC		
Clostridium perfringens		48.80	0.42	MNS 99	1		
Atopobium parvulum		35.91	0.39	CEL 4	99		
Actinomyces naeslundii		6.63	0.50		1		
03171121	**Low discrimination**						**03171121**
					SPO		
Actinomyces naeslundii		75.52	0.53	MNS 88	1		
Clostridium chauvoei		17.98	0.19	RAF 92	99		
Clostridium perfringens		3.74	0.08		84		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03171122	**Low discrimination**					**03171122**
Clostridium perfringens		42.01	0.37	MNS 99		
Lactobacillus acidophilus		28.86	0.32	CEL 4		
Actinomyces naeslundii		12.13	0.50			
03171123	**Acceptable identification**					**03171123**
					SPO	
Actinomyces naeslundii		93.87	0.53	MNS 88	1	
Clostridium chauvoei		3.94	0.06	RAF 92	99	
Clostridium perfringens		2.19	0.03		84	
03171141	**Good identification**					**03171141**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03171143	**Good identification**					**03171143**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03171161	**Good identification**					**03171161**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03171163	**Good identification**					**03171163**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03172100	**Good identification**					**03172100**
					GRA	
Propionibacterium propionicum		96.50	0.48	NIT 86	99	
Actinomyces naeslundii		2.52	0.31	CEL 1	99	
Fusobacterium mortiferum		0.98	0.12		1	
03172102	**Good identification**					**03172102**
Propionibacterium propionicum		98.29	0.51	NIT 86		
Actinomyces naeslundii		1.71	0.31	CEL 1		
03172140	**Good identification**					**03172140**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03172142	**Good identification**					**03172142**
Propionibacterium propionicum		100.0	0.51	NIT 86		
03173000	**Low discrimination**					**03173000**
					CAT	
Actinomyces naeslundii		57.13	0.47	MNS 88	25	
Actinomyces israelii		18.54	0.29	RAF 92	1	
Propionibacterium granulosum		14.62	0.22		90	
03173001	**Good identification**					**03173001**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03173002	**Low discrimination**					**03173002**
Actinomyces naeslundii		58.74	0.47	MNS 88		
Actinomyces israelii		28.60	0.32	RAF 92		
Lactobacillus jensenii		4.87	0.07			
03173003	**Good identification**					**03173003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MNS 88		
03173020	**Acceptable identification**					**03173020**
Actinomyces naeslundii		93.99	0.47	MNS 88		
Propionibacterium granulosum		6.01	0.12	RAF 92		
03173021	**Good identification**					**03173021**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03173022	**Good identification**					**03173022**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03173023	**Good identification**					**03173023**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03173041	**Good identification**					**03173041**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03173043	**Good identification**					**03173043**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03173061	**Good identification**					**03173061**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03173063	**Good identification**					**03173063**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03173100	**Low discrimination**					**03173100**
Propionibacterium propionicum		49.03	0.48	NIT 86		
Actinomyces naeslundii		30.77	0.55	CEL 1		
Actinomyces israelii		14.03	0.40			
03173101	**Low discrimination**					**03173101**
Actinomyces naeslundii		89.44	0.58	MNS 88		
Propionibacterium propionicum		10.56	0.32	RAF 92		
03173102	**Low discrimination**					**03173102**
Propionibacterium propionicum		56.41	0.51	NIT 86		
Actinomyces naeslundii		23.60	0.55	CEL 1		
Actinomyces israelii		16.15	0.43			
03173103	**Low discrimination**					**03173103**
Actinomyces naeslundii		84.96	0.58	MNS 88		
Propionibacterium propionicum		15.04	0.35	RAF 92		
03173111	**Good identification**					**03173111**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03173113	**Good identification**					**03173113**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03173120	**Low discrimination**					**03173120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		82.82	0.55	MNS 88	1	
Clostridium tertium		4.08	0.09	RAF 92	99	
Clostridium ramosum		3.85	0.07		92	
03173121	**Very good identification**					**03173121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03173122	**Low discrimination**					**03173122**
					SPO	
Actinomyces naeslundii		67.38	0.55	MNS 88	1	
Clostridium ramosum		12.52	0.17	RAF 92	92	
Clostridium tertium		9.97	0.17		99	
03173123	**Very good identification**					**03173123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03173131	**Good identification**					**03173131**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03173133	**Good identification**					**03173133**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03173140	**Low discrimination**					**03173140**
Propionibacterium propionicum		81.83	0.48	NIT 86		
Actinomyces israelii		9.11	0.33	CEL 1		
Actinomyces naeslundii		9.06	0.42			
03173141	**Low discrimination**					**03173141**
Actinomyces naeslundii		59.92	0.45	MNS 88		
Propionibacterium propionicum		40.08	0.32	RAF 92		
03173142	**Low discrimination**					**03173142**
Propionibacterium propionicum		84.38	0.51	NIT 86		
Actinomyces israelii		9.39	0.36	CEL 1		
Actinomyces naeslundii		6.23	0.42			
03173160	**Acceptable identification**					**03173160**
Actinomyces naeslundii		91.64	0.42	MNS 88		
Propionibacterium propionicum		8.36	0.14	RAF 92		
03173161	**Good identification**					**03173161**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03173162	**Low discrimination**					**03173162**
Actinomyces naeslundii		81.10	0.42	MNS 88		
Propionibacterium propionicum		11.10	0.17	RAF 92		
Actinomyces israelii		7.81	0.15			
03173163	**Good identification**					**03173163**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03173500	**Good identification**					**03173500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		95.87	0.38	SAL 92		
Propionibacterium propionicum		4.14	0.14	XYL 92		
03173502	**Low discrimination**					**03173502**
Actinomyces israelii		77.56	0.41	SAL 92		
Bifidobacterium adolescentis		19.09	0.24	XYL 92		
Propionibacterium propionicum		3.35	0.17			
03175002	**Low discrimination**					**03175002**
					COC	
Actinomyces israelii		74.23	0.38	SAL 92	1	
Atopobium parvulum		11.19	0.10		99	
Lactobacillus fermentum		8.35	0.20		1	
03175100	**Good identification**					**03175100**
					COC	
Atopobium parvulum		93.84	0.57	MNS 99	99	
Actinomyces israelii		2.96	0.45		1	
Actinomyces meyeri		2.42	0.32		1	
03175102	**Low discrimination**					**03175102**
					COC	
Atopobium parvulum		75.70	0.44	MNS 99	99	
Actinomyces israelii		20.28	0.48		1	
Bifidobacterium adolescentis		2.46	0.26		1	
03175120	**Good identification**					**03175120**
					COC	
Atopobium parvulum		95.89	0.44	MNS 99	99	
Actinomyces meyeri		1.91	0.17		1	
Actinomyces israelii		1.09	0.24		1	
03175140	**Low discrimination**					**03175140**
					COC	
Actinomyces israelii		54.82	0.38	SAL 92	1	
Atopobium parvulum		45.18	0.23		99	
03175142	**Low discrimination**					**03175142**
					COC	
Actinomyces israelii		88.07	0.41	SAL 92	1	
Atopobium parvulum		8.54	0.10		99	
Bifidobacterium adolescentis		3.40	0.10		1	
03175402	**Low discrimination**					**03175402**
Actinomyces israelii		87.91	0.36	SAL 92		
Lactobacillus fermentum		12.09	0.20			
03175500	**Low discrimination**					**03175500**
					COC	
Actinomyces israelii		57.10	0.43	SAL 92	1	
Atopobium parvulum		22.36	0.23		99	
Actinomyces meyeri		14.29	0.21		1	
03175502	**Low discrimination**					**03175502**
Actinomyces israelii		47.75	0.46	SAL 92		
Bifidobacterium adolescentis		47.37	0.40			
Lactobacillus fermentum		2.68	0.24			
03175512	**Low discrimination**					**03175512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		87.18	0.42	MNS 87			
Actinomyces israelii		10.74	0.32	RAF 92			
Clostridium clostridioforme		1.25	0.07				
03175540	**Good identification**						**03175540**
Actinomyces israelii		100.0	0.36	SAL 92			
03175542	**Low discrimination**						**03175542**
Actinomyces israelii		76.03	0.39	SAL 92			
Bifidobacterium adolescentis		23.97	0.25				
03177000	**Good identification**						**03177000**
Actinomyces israelii		100.0	0.47	SAL 92			
03177002	**Good identification**						**03177002**
Actinomyces israelii		100.0	0.50	SAL 92			
03177012	**Good identification**						**03177012**
Actinomyces israelii		100.0	0.36	SAL 92			
03177040	**Good identification**						**03177040**
Actinomyces israelii		100.0	0.40	SAL 92			
03177042	**Good identification**						**03177042**
Actinomyces israelii		100.0	0.43	SAL 92			
03177100	**Good identification**						**03177100**
					COC	SPO	
Actinomyces israelii		93.99	0.58	SAL 92	1	1	
Atopobium parvulum		5.32	0.23		99	1	
Clostridium tertium		0.41	0.03		1	99	
03177102	**Very good identification**						**03177102**
					SPO		
Actinomyces israelii		96.54	0.61	SAL 92	1		
Bifidobacterium adolescentis		1.69	0.24		1		
Clostridium tertium		0.84	0.11		99		
03177110	**Good identification**						**03177110**
					SPO		
Actinomyces israelii		95.10	0.44	SAL 92	1		
Clostridium tertium		4.90	0.07	MLZ 13	99		
03177112	**Low discrimination**						**03177112**
					SPO		
Actinomyces israelii		80.24	0.47	SAL 92	1		
Bifidobacterium adolescentis		11.49	0.26	MLZ 13	1		
Clostridium tertium		8.26	0.15		99		
03177120	**Low discrimination**						**03177120**
					COC	SPO	
Actinomyces israelii		73.55	0.37	SAL 92	1	1	
Clostridium tertium		14.96	0.11	NAG 6	1	99	
Atopobium parvulum		11.50	0.10		99	1	
03177122	**Low discrimination**						**03177122**
					SPO		
Actinomyces israelii		71.09	0.40	SAL 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		28.91	0.19	NAG 6	99	
03177140	**Good identification**					**03177140**
Actinomyces israelii		99.22	0.51	SAL 92		
Propionibacterium propionicum		0.78	0.14			
03177142	**Good identification**					**03177142**
Actinomyces israelii		98.67	0.54	SAL 92		
Propionibacterium propionicum		0.78	0.17			
Bifidobacterium adolescentis		0.55	0.09			
03177150	**Good identification**					**03177150**
Actinomyces israelii		100.0	0.37	SAL 92		
03177152	**Good identification**					**03177152**
Actinomyces israelii		95.65	0.40	SAL 92		
Bifidobacterium adolescentis		4.35	0.10	MLZ 13		
03177300	**Good identification**					**03177300**
Actinomyces israelii		100.0	0.39	SAL 92		
03177302	**Good identification**					**03177302**
Actinomyces israelii		100.0	0.42	SAL 92		
03177400	**Good identification**					**03177400**
Actinomyces israelii		100.0	0.46	SAL 92		
03177402	**Good identification**					**03177402**
Actinomyces israelii		100.0	0.49	SAL 92		
03177440	**Good identification**					**03177440**
Actinomyces israelii		100.0	0.39	SAL 92		
03177442	**Good identification**					**03177442**
Actinomyces israelii		100.0	0.42	SAL 92		
03177500	**Very good identification**					**03177500**
Actinomyces israelii		100.0	0.56	SAL 92		
03177502	**Low discrimination**					**03177502**
Actinomyces israelii		87.47	0.59	SAL 92		
Bifidobacterium adolescentis		12.53	0.39			
03177510	**Good identification**					**03177510**
Actinomyces israelii		100.0	0.42	SAL 92		
03177512	**Low discrimination**					**03177512**
Bifidobacterium adolescentis		53.95	0.40	MNS 87		
Actinomyces israelii		46.05	0.45	RAF 92		
03177520	**Good identification**					**03177520**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		100.0	0.36	SAL 92				
03177522	**Good identification**							**03177522**
Actinomyces israelii		100.0	0.39	SAL 92				
03177540	**Good identification**							**03177540**
Actinomyces israelii		100.0	0.49	SAL 92				
03177542	**Good identification**							**03177542**
Actinomyces israelii		95.65	0.52	SAL 92				
Bifidobacterium adolescentis		4.35	0.23					
03177550	**Good identification**							**03177550**
Actinomyces israelii		100.0	0.35	SAL 92				
03177552	**Low discrimination**							**03177552**
Actinomyces israelii		72.87	0.38	SAL 92				
Bifidobacterium adolescentis		27.13	0.25	MLZ 13				
03177700	**Good identification**							**03177700**
Actinomyces israelii		100.0	0.37	SAL 92				
03177702	**Good identification**							**03177702**
Actinomyces israelii		100.0	0.40	SAL 92				
03200000	**Low discrimination**							**03200000**
					COC	GRA	SPO	
Clostridium botulinum biovar A		26.68	0.42	SUC 7	1	99	99	
Megasphaera elsdenii		12.98	0.41	SAL 1	99	1	1	
Peptostreptococcus anaerobius		12.43	0.32		99	99	1	
03200001	**Acceptable identification**							**03200001**
					GRA	SPO		
Clostridium novyi biovar A		92.10	0.48	SUC 1	99	99		
Clostridium botulinum biovar A		2.12	0.08	SAL 8	99	99		
Campylobacter ureolyticus		1.93	-.03		1	1		
03200002	**Low discrimination**							**03200002**
					SPO			
Atopobium minutum		55.03	0.41	SUC 5	1			
Clostridium novyi biovar A		13.83	0.35	SAL 1	99			
Eubacterium tortuosum		12.69	0.19		1			
03200010	**Acceptable identification**							**03200010**
					COC	GRA		
Clostridium novyi biovar A		93.91	0.48	SUC 1	1	99		
Clostridium botulinum biovar A		2.16	0.08	SAL 8	1	99		
Megasphaera elsdenii		1.05	0.07		99	1		
03200011	**Good identification**							**03200011**
Clostridium novyi biovar A		100.0	0.48	SUC 1				
03200020	**Low discrimination**							**03200020**
					SPO			
Atopobium minutum		36.42	0.36	SUC 5	1			
Clostridium novyi biovar A		25.94	0.37	SAL 1	99			
Eubacterium tortuosum		16.79	0.19		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03200021	**Good identification**						**03200021**
Clostridium novyi biovar A		100.0	0.37	SUC 1			
03200030	**Good identification**						**03200030**
Clostridium novyi biovar A		100.0	0.37	SUC 1			
03200031	**Good identification**						**03200031**
Clostridium novyi biovar A		100.0	0.37	SUC 1			
03200040	**Low discrimination**						**03200040**
					COC	GRA	
Megasphaera elsdenii		34.90	0.38	SAL 1	99	1	
Clostridium botulinum biovar A		32.16	0.33	FRU 95	1	99	
Clostridium novyi biovar A		31.50	0.45		1	99	
03200041	**Good identification**						**03200041**
					COC	GRA	
Clostridium novyi biovar A		97.90	0.45	SUC 1	1	99	
Megasphaera elsdenii		1.10	0.04	SAL 8	99	1	
Clostridium botulinum biovar A		1.01	-.01		1	99	
03200050	**Good identification**						**03200050**
					COC	GRA	
Clostridium novyi biovar A		97.90	0.45	SUC 1	1	99	
Megasphaera elsdenii		1.10	0.04	SAL 8	99	1	
Clostridium botulinum biovar A		1.01	-.01		1	99	
03200051	**Good identification**						**03200051**
Clostridium novyi biovar A		100.0	0.45	SUC 1			
03200100	**Low discrimination**						**03200100**
					GRA	SPO	
Fusobacterium mortiferum		64.95	0.50	FRU 99	1	1	
Actinomyces meyeri		30.76	0.35		99	1	
Clostridium botulinum biovar A		1.32	0.08		99	99	
03201000	**Very good identification**						**03201000**
					GRA		
Eubacterium tortuosum		98.39	0.66	ESL 99	99		
Fusobacterium mortiferum		0.79	0.47		1		
Atopobium minutum		0.38	0.36		99		
03201002	**Good identification**						**03201002**
					GRA		
Eubacterium tortuosum		97.86	0.53	ESL 99	99		
Atopobium minutum		1.43	0.33		99		
Fusobacterium mortiferum		0.50	0.30		1		
03201020	**Good identification**						**03201020**
					SPO		
Eubacterium tortuosum		98.89	0.53	ESL 99	1		
Atopobium minutum		0.72	0.28		1		
Clostridium novyi biovar A		0.17	0.21		99		
03201022	**Good identification**						**03201022**
Eubacterium tortuosum		97.32	0.40	ESL 99			
Atopobium minutum		2.69	0.25				
03201100	**Low discrimination**						**03201100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Fusobacterium mortiferum		49.46	0.53	FRU 99	1			
Eubacterium tortuosum		26.65	0.32		99			
Actinomyces meyeri		23.43	0.38		99			
03201102	**Low discrimination**							**03201102**
					GRA			
Fusobacterium mortiferum		48.24	0.37	FRU 99	1			
Eubacterium tortuosum		41.28	0.19	BGL 10	99			
Actinomyces meyeri		6.36	0.12		99			
03202000	**Acceptable identification**							**03202000**
					COC	SPO	GRA	
Megasphaera elsdenii		93.33	0.41	SAL 1	99	1	1	
Clostridium botulinum biovar A		1.94	0.08	FRU 95	1	99	99	
Atopobium minutum		0.90	0.10		1	1	99	
03202040	**Good identification**							**03202040**
					COC	GRA		
Megasphaera elsdenii		98.19	0.38	SAL 1	99	1		
Clostridium botulinum biovar A		0.91	-.01	FRU 95	1	99		
Clostridium novyi biovar A		0.90	0.11		1	99		
03204100	**Acceptable identification**							**03204100**
					GRA			
Actinomyces meyeri		92.43	0.40	SAL 1	99			
Fusobacterium mortiferum		7.57	0.31	FRU 95	1			
03205100	**Acceptable identification**							**03205100**
					GRA			
Actinomyces meyeri		91.42	0.43	SAL 1	99			
Fusobacterium mortiferum		7.48	0.34	FRU 95	1			
Lactobacillus fermentum		0.55	0.16		99			
03210000	**Low discrimination**							**03210000**
					COC	GRA		
Fusobacterium mortiferum		64.24	0.78		1	1		
Megasphaera elsdenii		28.08	0.63		99	1		
Actinomyces meyeri		3.41	0.47		1	99		
03210001	**Low discrimination**							**03210001**
					COC	GRA	SPO	
Fusobacterium mortiferum		56.60	0.44	URE 1	1	1	1	
Megasphaera elsdenii		24.74	0.29		99	1	1	
Clostridium novyi biovar A		12.93	0.31		1	99	99	
03210002	**Low discrimination**							**03210002**
					COC	GRA		
Fusobacterium mortiferum		61.71	0.62	BGL 10	1	1		
Megasphaera elsdenii		26.98	0.47		99	1		
Atopobium minutum		7.47	0.40		1	99		
03210010	**Low discrimination**							**03210010**
					COC	GRA	SPO	
Fusobacterium mortiferum		37.55	0.44	MLZ 1	1	1	1	
Clostridium difficile		33.67	0.27		1	99	99	
Megasphaera elsdenii		16.41	0.29		99	1	1	
03210020	**Low discrimination**							**03210020**
					GRA			
Fusobacterium mortiferum		27.08	0.44	NAG 1	1			
Actinomyces meyeri		19.39	0.32		99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		18.04	0.35		99		
03210040	**Good identification**						**03210040**
					COC	GRA	
Megasphaera elsdenii		93.30	0.60	SAL 1	99	1	
Fusobacterium mortiferum		3.23	0.44		1	1	
Pseudoramibacter alactolyticus		2.24	0.21		1	99	
03210042	**Acceptable identification**						**03210042**
					COC	GRA	
Megasphaera elsdenii		93.35	0.44	SAL 1	99	1	
Fusobacterium mortiferum		3.24	0.27	BGL 10	1	1	
Pseudoramibacter alactolyticus		2.24	0.05		1	99	
03210100	**Good identification**						**03210100**
					COC	GRA	
Fusobacterium mortiferum		91.45	0.84		1	1	
Actinomyces meyeri		8.31	0.57		1	99	
Megasphaera elsdenii		0.17	0.29		99	1	
03210101	**Acceptable identification**						**03210101**
					GRA		
Fusobacterium mortiferum		91.67	0.50	URE 1	1		
Actinomyces meyeri		8.33	0.23		99		
03210102	**Very good identification**						**03210102**
					COC	GRA	
Fusobacterium mortiferum		97.02	0.68	BGL 10	1	1	
Actinomyces meyeri		2.46	0.31		1	99	
Megasphaera elsdenii		0.18	0.12		99	1	
03210110	**Acceptable identification**						**03210110**
					GRA		
Fusobacterium mortiferum		91.67	0.50	MLZ 1	1		
Actinomyces meyeri		8.33	0.23		99		
03210120	**Low discrimination**						**03210120**
					GRA		
Actinomyces meyeri		55.10	0.42	SAL 1	99		
Fusobacterium mortiferum		44.90	0.50	NAG 12	1		
03210140	**Low discrimination**						**03210140**
					COC	GRA	
Fusobacterium mortiferum		82.25	0.50	SOR 1	1	1	
Megasphaera elsdenii		10.27	0.26		99	1	
Actinomyces meyeri		7.48	0.23		1	99	
03210200	**Low discrimination**						**03210200**
					COC	GRA	
Fusobacterium mortiferum		50.91	0.44	RHA 1	1	1	
Megasphaera elsdenii		22.26	0.29		99	1	
Actinomyces odontolyticus		21.69	0.32		1	99	
03210300	**Low discrimination**						**03210300**
					GRA		
Fusobacterium mortiferum		87.07	0.50	RHA 1	1		
Actinomyces meyeri		7.92	0.23		99		
Actinomyces odontolyticus		5.02	0.23		99		
03210400	**Low discrimination**						**03210400**
					COC	GRA	
Actinomyces meyeri		46.32	0.36	SAL 1	1	99	
Fusobacterium mortiferum		35.29	0.44		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		15.43	0.29		99	1	
03210500	**Low discrimination**						**03210500**
					GRA		
Actinomyces meyeri		68.93	0.47	SAL 1	99		
Fusobacterium mortiferum		30.64	0.50		1		
Lactobacillus fermentum		0.44	0.20		99		
03211000	**Good identification**						**03211000**
					GRA		
Fusobacterium mortiferum		91.74	0.81		1		
Actinomyces meyeri		4.87	0.50		99		
Lactobacillus fermentum		1.25	0.50		99		
03211001	**Acceptable identification**						**03211001**
					GRA	SPO	
Fusobacterium mortiferum		90.02	0.47	URE 1	1	1	
Actinomyces meyeri		4.77	0.16		99	1	
Clostridium novyi biovar A		1.52	0.15		99	99	
03211002	**Low discrimination**						**03211002**
					GRA		
Fusobacterium mortiferum		83.28	0.65	BGL 10	1		
Lactobacillus fermentum		10.18	0.50		99		
Atopobium minutum		2.24	0.32		99		
03211010	**Low discrimination**						**03211010**
					GRA	SPO	
Fusobacterium mortiferum		89.95	0.47	MLZ 1	1	1	
Actinomyces meyeri		4.77	0.16		99	1	
Clostridium novyi biovar A		1.52	0.15		99	99	
03211020	**Low discrimination**						**03211020**
					GRA		
Fusobacterium mortiferum		38.40	0.47	NAG 1	1		
Actinomyces meyeri		27.49	0.35		99		
Lactobacillus fermentum		17.20	0.42		99		
03211022	**Low discrimination**						**03211022**
					GRA		
Lactobacillus fermentum		59.07	0.42	SAL 1	99		
Fusobacterium mortiferum		14.65	0.30		1		
Atopobium minutum		13.01	0.24		99		
03211040	**Low discrimination**						**03211040**
					COC	GRA	
Fusobacterium mortiferum		77.83	0.47	SOR 1	1	1	
Megasphaera elsdenii		15.12	0.26		99	1	
Actinomyces meyeri		4.13	0.16		1	99	
03211100	**Good identification**						**03211100**
					GRA		
Fusobacterium mortiferum		90.83	0.87		1		
Actinomyces meyeri		8.26	0.60		99		
Lactobacillus fermentum		0.86	0.54		99		
03211101	**Low discrimination**						**03211101**
					GRA		
Fusobacterium mortiferum		88.39	0.53	URE 1	1		
Actinomyces meyeri		8.04	0.26		99		
Actinomyces naeslundii		1.58	0.35		99		
03211102	**Low discrimination**						**03211102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		89.63	0.71	BGL 10	1	
Lactobacillus fermentum		7.66	0.54		99	
Actinomyces meyeri		2.27	0.34		99	
03211103	**Low discrimination**					**03211103**
					GRA	
Fusobacterium mortiferum		80.26	0.37	URE 1	1	
Actinomyces naeslundii		12.89	0.35	BGL 10	99	
Lactobacillus fermentum		6.86	0.20		99	
03211110	**Acceptable identification**					**03211110**
					GRA	
Fusobacterium mortiferum		90.88	0.53	MLZ 1	1	
Actinomyces meyeri		8.26	0.26		99	
Lactobacillus fermentum		0.86	0.20		99	
03211112	**Acceptable identification**					**03211112**
					GRA	
Fusobacterium mortiferum		92.13	0.37	MLZ 1	1	
Lactobacillus fermentum		7.87	0.20	BGL 10	99	
03211120	**Low discrimination**					**03211120**
					GRA	
Actinomyces meyeri		47.15	0.45	SAL 1	99	
Fusobacterium mortiferum		38.42	0.53	NAG 12	1	
Lactobacillus fermentum		12.04	0.46		99	
03211122	**Low discrimination**					**03211122**
					GRA	
Lactobacillus fermentum		60.82	0.46	SAL 1	99	
Fusobacterium mortiferum		21.57	0.37		1	
Actinomyces meyeri		7.37	0.19		99	
03211140	**Acceptable identification**					**03211140**
					GRA	
Fusobacterium mortiferum		90.88	0.53	SOR 1	1	
Actinomyces meyeri		8.26	0.26		99	
Lactobacillus fermentum		0.86	0.20		99	
03211142	**Acceptable identification**					**03211142**
					GRA	
Fusobacterium mortiferum		92.13	0.37	SOR 1	1	
Lactobacillus fermentum		7.87	0.20	BGL 10	99	
03211200	**Low discrimination**					**03211200**
					GRA	
Fusobacterium mortiferum		78.82	0.47	RHA 1	1	
Actinomyces odontolyticus		14.93	0.29		99	
Actinomyces meyeri		4.18	0.16		99	
03211300	**Low discrimination**					**03211300**
					GRA	
Fusobacterium mortiferum		88.81	0.53	RHA 1	1	
Actinomyces meyeri		8.07	0.26		99	
Actinomyces odontolyticus		2.28	0.20		99	
03211302	**Low discrimination**					**03211302**
					GRA	
Fusobacterium mortiferum		81.51	0.37	RHA 1	1	
Actinomyces odontolyticus		11.52	0.17	BGL 10	99	
Lactobacillus fermentum		6.96	0.20		99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03211400	**Low discrimination**							**03211400**
					GRA			
Lactobacillus fermentum		36.63	0.50	SAL 1	99			
Actinomyces meyeri		35.77	0.39		99			
Fusobacterium mortiferum		27.25	0.47		1			
03211402	**Low discrimination**							**03211402**
					GRA			
Lactobacillus fermentum		89.86	0.50	SAL 1	99			
Fusobacterium mortiferum		7.43	0.30		1			
Actinomyces meyeri		2.71	0.14		99			
03211420	**Low discrimination**							**03211420**
					GRA			
Lactobacillus fermentum		70.32	0.42	SAL 1	99			
Actinomyces meyeri		28.09	0.25		99			
Fusobacterium mortiferum		1.59	0.13		1			
03211422	**Good identification**							**03211422**
Lactobacillus fermentum		100.0	0.42	SAL 1				
03211500	**Low discrimination**							**03211500**
					GRA			
Actinomyces meyeri		53.70	0.50	SAL 1	99			
Fusobacterium mortiferum		23.87	0.53		1			
Lactobacillus fermentum		22.43	0.54		99			
03211502	**Low discrimination**							**03211502**
					GRA			
Lactobacillus fermentum		83.87	0.54	SAL 1	99			
Fusobacterium mortiferum		9.92	0.37		1			
Actinomyces meyeri		6.21	0.24		99			
03211520	**Low discrimination**							**03211520**
					GRA			
Lactobacillus fermentum		49.71	0.46	SAL 1	99			
Actinomyces meyeri		48.69	0.35		99			
Fusobacterium mortiferum		1.60	0.19		1			
03211522	**Good identification**							**03211522**
Lactobacillus fermentum		97.06	0.46	SAL 1				
Actinomyces meyeri		2.94	0.09					
03212000	**Low discrimination**							**03212000**
					COC	SPO	GRA	
Megasphaera elsdenii		89.12	0.63	SAL 1	99	1	1	
Clostridium difficile		3.40	0.32		1	99	99	
Pseudoramibacter alactolyticus		3.07	0.27		1	1	99	
03212002	**Acceptable identification**							**03212002**
					COC	GRA		
Megasphaera elsdenii		92.70	0.47	SAL 1	99	1		
Pseudoramibacter alactolyticus		3.19	0.11	BGL 10	1	99		
Fusobacterium mortiferum		2.14	0.27		1	1		
03212010	**Acceptable identification**							**03212010**
					COC	SPO	GRA	
Clostridium difficile		94.69	0.43	SUC 1	1	99	99	
Megasphaera elsdenii		5.13	0.29		99	1	1	
Pseudoramibacter alactolyticus		0.18	-.07		1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03212040	**Very good identification**							**03212040**
					COC	SPO	GRA	
Megasphaera elsdenii		98.43	0.60	SAL 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.17		1	1	99	
Clostridium difficile		0.30	0.10		1	99	99	
03212042	**Good identification**							**03212042**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	SAL 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
03212100	**Low discrimination**							**03212100**
					COC	GRA		
Fusobacterium mortiferum		76.95	0.50	MAN 1	1	1		
Megasphaera elsdenii		14.42	0.29		99	1		
Actinomyces meyeri		7.00	0.23		1	99		
03213000	**Low discrimination**							**03213000**
					COC	GRA	SPO	
Fusobacterium mortiferum		46.08	0.47	MAN 1	1	1	1	
Clostridium innocuum		33.11	0.32		1	99	99	
Megasphaera elsdenii		13.43	0.29		99	1	1	
03213100	**Low discrimination**							**03213100**
					GRA			
Fusobacterium mortiferum		88.60	0.53	MAN 1	1			
Actinomyces meyeri		8.05	0.26		99			
Actinomyces naeslundii		2.05	0.37		99			
03213101	**Low discrimination**							**03213101**
					GRA			
Actinomyces naeslundii		77.42	0.40	MNS 88	99			
Fusobacterium mortiferum		22.58	0.19	MLT 94	1			
03213102	**Low discrimination**							**03213102**
					GRA			
Fusobacterium mortiferum		73.37	0.37	MAN 1	1			
Actinomyces naeslundii		15.25	0.37	BGL 10	99			
Lactobacillus fermentum		6.27	0.20		99			
03213103	**Good identification**							**03213103**
Actinomyces naeslundii		100.0	0.40	MNS 88				
03213120	**Low discrimination**							**03213120**
					GRA			
Actinomyces naeslundii		50.65	0.37	MNS 88	99			
Actinomyces meyeri		27.19	0.11	MLT 94	99			
Fusobacterium mortiferum		22.16	0.19	RAF 92	1			
03213121	**Good identification**							**03213121**
Actinomyces naeslundii		100.0	0.40	MNS 88				
03213122	**Good identification**							**03213122**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03213123	**Good identification**							**03213123**
Actinomyces naeslundii		100.0	0.40	MNS 88				
03214000	**Low discrimination**							**03214000**
					COC	GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		55.55	0.52	SAL 1	1	99	
Fusobacterium mortiferum		40.62	0.59		1	1	
Megasphaera elsdenii		2.38	0.29		99	1	
03214002	**Low discrimination**						**03214002**
					GRA		
Fusobacterium mortiferum		64.51	0.42	XYL 7	1		
Actinomyces meyeri		24.56	0.26	BGL 10	99		
Actinomyces odontolyticus		5.16	0.18		99		
03214020	**Acceptable identification**						**03214020**
					GRA		
Actinomyces meyeri		93.74	0.37	SAL 1	99		
Fusobacterium mortiferum		5.08	0.24	NAG 12	1		
Actinomyces odontolyticus		1.19	0.08		99		
03214100	**Low discrimination**						**03214100**
					GRA		
Actinomyces meyeri		70.04	0.62	SAL 1	99		
Fusobacterium mortiferum		29.87	0.65		1		
Actinomyces odontolyticus		0.06	0.13		99		
03214102	**Low discrimination**						**03214102**
					GRA		
Fusobacterium mortiferum		59.76	0.49	XYL 7	1		
Actinomyces meyeri		39.00	0.36	BGL 10	99		
Actinomyces odontolyticus		0.65	0.09		99		
03214120	**Good identification**						**03214120**
					GRA		
Actinomyces meyeri		96.94	0.47	SAL 1	99		
Fusobacterium mortiferum		3.06	0.31	NAG 12	1		
03214400	**Good identification**						**03214400**
					GRA		
Actinomyces meyeri		96.69	0.41	SAL 1	99		
Fusobacterium mortiferum		2.86	0.24		1		
Lactobacillus fermentum		0.45	0.12		99		
03214500	**Good identification**						**03214500**
					GRA		
Actinomyces meyeri		98.12	0.52	SAL 1	99		
Fusobacterium mortiferum		1.69	0.31		1		
Lactobacillus fermentum		0.19	0.16		99		
03214520	**Good identification**						**03214520**
Actinomyces meyeri		100.0	0.37	SAL 1			
03215000	**Low discrimination**						**03215000**
					GRA		
Actinomyces meyeri		55.11	0.55	SAL 1	99		
Fusobacterium mortiferum		40.30	0.62		1		
Lactobacillus fermentum		4.27	0.46		99		
03215002	**Low discrimination**						**03215002**
					GRA		
Fusobacterium mortiferum		42.20	0.45	XYL 7	1		
Lactobacillus fermentum		40.23	0.46	BGL 10	99		
Actinomyces meyeri		16.07	0.29		99		
03215020	**Low discrimination**						**03215020**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		80.42	0.40	SAL 1	99	
Lactobacillus fermentum		15.23	0.38	NAG 12	99	
Fusobacterium mortiferum		4.36	0.27		1	
03215022	**Low discrimination**					**03215022**
					GRA	
Lactobacillus fermentum		83.67	0.38	SAL 1	99	
Actinomyces meyeri		13.67	0.14		99	
Fusobacterium mortiferum		2.66	0.11		1	
03215100	**Low discrimination**					**03215100**
					GRA	
Actinomyces meyeri		68.56	0.65	SAL 1	99	
Fusobacterium mortiferum		29.25	0.68		1	
Lactobacillus fermentum		2.17	0.50		99	
03215102	**Low discrimination**					**03215102**
					GRA	
Fusobacterium mortiferum		43.03	0.52	XYL 7	1	
Lactobacillus fermentum		28.69	0.50	BGL 10	99	
Actinomyces meyeri		28.08	0.39		99	
03215120	**Acceptable identification**					**03215120**
					GRA	
Actinomyces meyeri		90.19	0.50	SAL 1	99	
Lactobacillus fermentum		6.97	0.42	NAG 12	99	
Fusobacterium mortiferum		2.85	0.34		1	
03215122	**Low discrimination**					**03215122**
					GRA	
Lactobacillus fermentum		69.16	0.42	SAL 1	99	
Actinomyces meyeri		27.70	0.24		99	
Fusobacterium mortiferum		3.14	0.17		1	
03215400	**Low discrimination**					**03215400**
					GRA	
Actinomyces meyeri		74.66	0.44	SAL 1	99	
Lactobacillus fermentum		23.13	0.46		99	
Fusobacterium mortiferum		2.21	0.27		1	
03215402	**Acceptable identification**					**03215402**
					GRA	
Lactobacillus fermentum		90.06	0.46	SAL 1	99	
Actinomyces meyeri		8.99	0.19		99	
Fusobacterium mortiferum		0.95	0.11		1	
03215422	**Good identification**					**03215422**
Lactobacillus fermentum		96.08	0.38	SAL 1		
Actinomyces meyeri		3.92	0.04			
03215500	**Low discrimination**					**03215500**
					GRA	
Actinomyces meyeri		87.44	0.55	SAL 1	99	
Lactobacillus fermentum		11.05	0.50		99	
Fusobacterium mortiferum		1.51	0.34		1	
03215502	**Low discrimination**					**03215502**
Lactobacillus fermentum		78.03	0.50	SAL 1		
Actinomyces meyeri		19.10	0.29			
Bifidobacterium adolescentis		1.69	0.16			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03215520	**Low discrimination**							**03215520**
Actinomyces meyeri		76.40	0.40	SAL 1				
Lactobacillus fermentum		23.60	0.42	NAG 12				
03215522	**Acceptable identification**							**03215522**
Lactobacillus fermentum		90.90	0.42	SAL 1				
Actinomyces meyeri		9.10	0.14					
03220100	**Good identification**							**03220100**
Fusobacterium mortiferum		100.0	0.41	FRU 99				
03221100	**Good identification**							**03221100**
					GRA			
Fusobacterium mortiferum		96.69	0.44	FRU 99	1			
Eubacterium tortuosum		1.76	-.03		99			
Actinomyces meyeri		1.55	0.04		99			
03230000	**Very good identification**							**03230000**
					COC	GRA	CAT	
Fusobacterium mortiferum		97.14	0.69		1	1	5	
Megasphaera elsdenii		1.44	0.29		99	1	1	
Propionibacterium granulosum		0.80	0.30		1	99	90	
03230002	**Good identification**							**03230002**
					COC	GRA	CAT	
Fusobacterium mortiferum		96.39	0.53	BGL 10	1	1	5	
Propionibacterium granulosum		1.79	0.20		1	99	90	
Megasphaera elsdenii		1.43	0.12		99	1	1	
03230100	**Very good identification**							**03230100**
					GRA			
Fusobacterium mortiferum		99.69	0.75		1			
Actinomyces meyeri		0.31	0.23		99			
03230101	**Good identification**							**03230101**
Fusobacterium mortiferum		100.0	0.41	URE 1				
03230102	**Very good identification**							**03230102**
Fusobacterium mortiferum		100.0	0.59	BGL 10				
03230110	**Good identification**							**03230110**
Fusobacterium mortiferum		100.0	0.41	MLZ 1				
03230120	**Good identification**							**03230120**
					GRA			
Fusobacterium mortiferum		96.02	0.41	NAG 1	1			
Actinomyces meyeri		3.99	0.08		99			
03230140	**Good identification**							**03230140**
Fusobacterium mortiferum		100.0	0.41	SOR 1				
03230300	**Good identification**							**03230300**
Fusobacterium mortiferum		100.0	0.41	RHA 1				
03230500	**Acceptable identification**							**03230500**
					GRA			
Fusobacterium mortiferum		92.93	0.41	ARA 1	1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		7.07	0.13		99		
03231000	**Very good identification**						**03231000**
					GRA	CAT	
Fusobacterium mortiferum		98.67	0.72		1	5	
Propionibacterium granulosum		0.54	0.30		99	90	
Lactobacillus jensenii		0.47	0.22		99	1	
03231001	**Acceptable identification**						**03231001**
					GRA		
Fusobacterium mortiferum		91.32	0.38	URE 1	1		
Actinomyces naeslundii		8.68	0.32		99		
03231002	**Good identification**						**03231002**
					GRA	CAT	
Fusobacterium mortiferum		95.19	0.56	BGL 10	1	5	
Lactobacillus jensenii		2.69	0.19		99	1	
Propionibacterium granulosum		1.18	0.20		99	90	
03231010	**Low discrimination**						**03231010**
					GRA	CAT	
Fusobacterium mortiferum		81.32	0.38	MLZ 1	1	5	
Lactobacillus jensenii		12.65	0.14		99	1	
Propionibacterium granulosum		6.03	0.15		99	90	
03231020	**Low discrimination**						**03231020**
					GRA	CAT	
Fusobacterium mortiferum		72.45	0.38	NAG 1	1	5	
Propionibacterium granulosum		9.85	0.20		99	90	
Lactobacillus jensenii		8.45	0.12		99	1	
03231040	**Good identification**						**03231040**
Fusobacterium mortiferum		100.0	0.38	SOR 1			
03231100	**Very good identification**						**03231100**
					GRA		
Fusobacterium mortiferum		99.50	0.78		1		
Actinomyces meyeri		0.31	0.26		99		
Actinomyces naeslundii		0.08	0.37		99		
03231101	**Low discrimination**						**03231101**
					GRA		
Fusobacterium mortiferum		89.61	0.44	URE 1	1		
Actinomyces naeslundii		10.39	0.40		99		
03231102	**Very good identification**						**03231102**
					GRA		
Fusobacterium mortiferum		98.29	0.62	BGL 10	1		
Actinomyces naeslundii		0.69	0.37		99		
Lactobacillus acidophilus		0.62	0.11		99		
03231103	**Low discrimination**						**03231103**
					GRA		
Actinomyces naeslundii		51.06	0.40	MNS 88	99		
Fusobacterium mortiferum		48.94	0.28	MLT 94	1		
03231110	**Good identification**						**03231110**
Fusobacterium mortiferum		100.0	0.44	MLZ 1			
03231120	**Low discrimination**						**03231120**
					GRA		
Fusobacterium mortiferum		73.96	0.44	NAG 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		14.94	0.21		99			
Actinomyces naeslundii		5.72	0.37		99			
03231121	**Good identification**							**03231121**
Actinomyces naeslundii		100.0	0.40	MNS 88				
03231123	**Good identification**							**03231123**
Actinomyces naeslundii		100.0	0.40	MNS 88				
03231140	**Good identification**							**03231140**
					GRA			
Fusobacterium mortiferum		98.65	0.44	SOR 1	1			
Actinomyces naeslundii		1.35	0.24		99			
03231200	**Good identification**							**03231200**
Fusobacterium mortiferum		100.0	0.38	RHA 1				
03231300	**Good identification**							**03231300**
Fusobacterium mortiferum		100.0	0.44	RHA 1				
03231400	**Acceptable identification**							**03231400**
					GRA			
Fusobacterium mortiferum		91.76	0.38	ARA 1	1			
Lactobacillus fermentum		4.17	0.16		99			
Actinomyces meyeri		4.07	0.05		99			
03231500	**Acceptable identification**							**03231500**
					GRA			
Fusobacterium mortiferum		90.26	0.44	ARA 1	1			
Actinomyces meyeri		6.87	0.16		99			
Lactobacillus fermentum		2.87	0.20		99			
03232100	**Good identification**							**03232100**
					GRA			
Fusobacterium mortiferum		97.92	0.41	MAN 1	1			
Propionibacterium propionicum		2.08	0.14		99			
03233000	**Low discrimination**							**03233000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		42.05	0.38	MAN 1	1	1	5	
Clostridium innocuum		35.53	0.24		99	99	1	
Propionibacterium granulosum		17.25	0.28		99	1	90	
03233001	**Good identification**							**03233001**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03233003	**Good identification**							**03233003**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03233021	**Good identification**							**03233021**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03233023	**Good identification**							**03233023**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03233100	**Low discrimination**							**03233100**
					GRA			
Fusobacterium mortiferum		85.90	0.44	MAN 1	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		12.89	0.42		99		
Propionibacterium propionicum		1.22	0.14		99		
03233101	**Good identification**						**03233101**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03233102	**Low discrimination**						**03233102**
					GRA		
Actinomyces naeslundii		53.14	0.42	MNS 88	99		
Fusobacterium mortiferum		39.35	0.28	MLT 94	1		
Propionibacterium propionicum		7.52	0.17	RAF 92	99		
03233103	**Good identification**						**03233103**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03233120	**Good identification**						**03233120**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03233121	**Good identification**						**03233121**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03233122	**Good identification**						**03233122**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03233123	**Good identification**						**03233123**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03234000	**Good identification**						**03234000**
					GRA		
Fusobacterium mortiferum		95.58	0.50	XYL 7	1		
Actinomyces meyeri		4.42	0.18		99		
03234100	**Acceptable identification**						**03234100**
					GRA		
Fusobacterium mortiferum		92.65	0.56	XYL 7	1		
Actinomyces meyeri		7.35	0.28		99		
03234102	**Good identification**						**03234102**
					GRA		
Fusobacterium mortiferum		97.84	0.40	XYL 7	1		
Actinomyces meyeri		2.16	0.02	BGL 10	99		
03235000	**Good identification**						**03235000**
					GRA		
Fusobacterium mortiferum		95.25	0.53	XYL 7	1		
Actinomyces meyeri		4.41	0.21		99		
Lactobacillus fermentum		0.34	0.12		99		
03235002	**Good identification**						**03235002**
					GRA		
Fusobacterium mortiferum		96.88	0.36	XYL 7	1		
Lactobacillus fermentum		3.12	0.12	BGL 10	99		
03235100	**Good identification**						**03235100**
					COC	GRA	
Fusobacterium mortiferum		90.77	0.59	XYL 7	1	1	
Actinomyces meyeri		7.20	0.31		1	99	
Atopobium parvulum		1.80	0.18		99	99	
03235102	**Acceptable identification**						**03235102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Fusobacterium mortiferum		92.92	0.43	XYL 7	1	1	
Atopobium parvulum		2.93	0.06	BGL 10	99	99	
Lactobacillus fermentum		2.10	0.16		1	99	
03240000	**Low discrimination**						**03240000**
					GRA	SPO	
Fusobacterium mortiferum		49.17	0.36	FRU 99	1	1	
Eubacterium tortuosum		30.91	0.15		99	1	
Clostridium botulinum biovar A		6.98	0.08		99	99	
03240100	**Good identification**						**03240100**
Fusobacterium mortiferum		100.0	0.42	FRU 99			
03241000	**Good identification**						**03241000**
					GRA		
Eubacterium tortuosum		97.07	0.50	ESL 99	99		
Fusobacterium mortiferum		2.34	0.39	CEL 10	1		
Lactobacillus catenaformis		0.26	0.12		99		
03241002	**Good identification**						**03241002**
					SPO		
Eubacterium tortuosum		95.60	0.37	ESL 99	1		
Clostridium innocuum		1.48	0.10	CEL 10	99		
Lactobacillus catenaformis		1.47	0.12		1		
03241020	**Good identification**						**03241020**
Eubacterium tortuosum		99.62	0.37	ESL 99			
Lactobacillus catenaformis		0.38	0.02	CEL 10			
03241100	**Low discrimination**						**03241100**
					GRA		
Fusobacterium mortiferum		78.97	0.45	FRU 99	1		
Eubacterium tortuosum		14.18	0.15		99		
Lactobacillus catenaformis		5.71	0.15		99		
03243000	**Good identification**						**03243000**
					SPO		
Clostridium innocuum		96.40	0.44	MNS 99	99		
Eubacterium tortuosum		3.60	0.15	FRU 95	1		
03243002	**Good identification**						**03243002**
					SPO		
Clostridium innocuum		99.35	0.44	MNS 99	99		
Eubacterium tortuosum		0.65	0.03	FRU 95	1		
03250000	**Very good identification**						**03250000**
					COC	GRA	
Fusobacterium mortiferum		98.03	0.70		1	1	
Megasphaera elsdenii		1.30	0.29		99	1	
Lactobacillus catenaformis		0.17	0.12		1	99	
03250001	**Good identification**						**03250001**
Fusobacterium mortiferum		100.0	0.36	URE 1			
03250002	**Acceptable identification**						**03250002**
					GRA	SPO	
Fusobacterium mortiferum		94.78	0.53	BGL 10	1	1	
Clostridium innocuum		1.47	0.10		99	99	
Lactobacillus catenaformis		1.45	0.12		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03250010	**Low discrimination**						**03250010**
					GRA	SPO	
Fusobacterium mortiferum		87.60	0.36	MLZ 1	1	1	
Clostridium difficile		12.40	0.05		99	99	
03250020	**Low discrimination**						**03250020**
					GRA	SPO	
Fusobacterium mortiferum		89.00	0.36	NAG 1	1	1	
Clostridium barati		7.25	0.02		99	99	
Lactobacillus catenaformis		3.75	0.02		99	1	
03250040	**Low discrimination**						**03250040**
					COC		
Fusobacterium mortiferum		53.35	0.36	SOR 1	1		
Megasphaera elsdenii		46.65	0.26		99		
03250100	**Very good identification**						**03250100**
					GRA		
Fusobacterium mortiferum		99.58	0.76		1		
Actinomyces meyeri		0.27	0.23		99		
Lactobacillus catenaformis		0.11	0.15		99		
03250101	**Good identification**						**03250101**
Fusobacterium mortiferum		100.0	0.42	URE 1			
03250102	**Very good identification**						**03250102**
					GRA	SPO	
Fusobacterium mortiferum		97.86	0.60	BGL 10	1	1	
Lactobacillus catenaformis		0.97	0.15		99	1	
Clostridium barati		0.93	0.10		99	99	
03250110	**Good identification**						**03250110**
Fusobacterium mortiferum		100.0	0.42	MLZ 1			
03250120	**Low discrimination**						**03250120**
					GRA	SPO	
Fusobacterium mortiferum		85.54	0.42	NAG 1	1	1	
Clostridium barati		8.96	0.10		99	99	
Actinomyces meyeri		3.18	0.08		99	1	
03250140	**Good identification**						**03250140**
Fusobacterium mortiferum		100.0	0.42	SOR 1			
03250200	**Good identification**						**03250200**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
03250300	**Good identification**						**03250300**
Fusobacterium mortiferum		100.0	0.42	RHA 1			
03250400	**Good identification**						**03250400**
					GRA		
Fusobacterium mortiferum		96.17	0.36	ARA 1	1		
Actinomyces meyeri		3.83	0.02		99		
03250500	**Acceptable identification**						**03250500**
					GRA		
Fusobacterium mortiferum		93.62	0.42	ARA 1	1		
Actinomyces meyeri		6.38	0.13		99		
03251000	**Low discrimination**						**03251000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		85.39	0.73		1		
Lactobacillus catenaformis		9.61	0.47		99		
Lactobacillus jensenii		2.28	0.36		99		
03251001	**Low discrimination**						**03251001**
					GRA		
Fusobacterium mortiferum		83.51	0.39	URE 1	1		
Lactobacillus catenaformis		9.40	0.12		99		
Actinomyces naeslundii		4.86	0.29		99		
03251002	**Low discrimination**						**03251002**
					GRA	SPO	
Lactobacillus catenaformis		40.33	0.47	MNS 99	99	1	
Fusobacterium mortiferum		39.84	0.56	MLT 90	1	1	
Clostridium innocuum		7.81	0.32		99	99	
03251010	**Low discrimination**						**03251010**
					GRA		
Fusobacterium mortiferum		47.96	0.39	MLZ 1	1		
Lactobacillus jensenii		42.34	0.28		99		
Lactobacillus catenaformis		5.40	0.12		99		
03251020	**Low discrimination**						**03251020**
					GRA	SPO	
Lactobacillus catenaformis		49.80	0.36	MNS 99	99	1	
Clostridium barati		18.46	0.24	MLT 90	99	99	
Fusobacterium mortiferum		17.89	0.39		1	1	
03251040	**Low discrimination**						**03251040**
					GRA	SPO	
Fusobacterium mortiferum		85.53	0.39	SOR 1	1	1	
Lactobacillus catenaformis		9.62	0.12		99	1	
Clostridium barati		2.56	-.03		99	99	
03251100	**Good identification**						**03251100**
					GRA	SPO	
Fusobacterium mortiferum		92.50	0.79		1	1	
Lactobacillus catenaformis		6.69	0.50		99	1	
Clostridium barati		0.41	0.24		99	99	
03251101	**Low discrimination**						**03251101**
					GRA		
Fusobacterium mortiferum		87.47	0.45	URE 1	1		
Lactobacillus catenaformis		6.33	0.15		99		
Actinomyces naeslundii		6.21	0.37		99		
03251102	**Low discrimination**						**03251102**
					GRA		
Fusobacterium mortiferum		51.44	0.63	BGL 10	1		
Lactobacillus catenaformis		33.48	0.50		99		
Lactobacillus acidophilus		8.52	0.36		99		
03251110	**Acceptable identification**						**03251110**
					GRA		
Fusobacterium mortiferum		93.26	0.45	MLZ 1	1		
Lactobacillus catenaformis		6.74	0.15		99		
03251120	**Low discrimination**						**03251120**
					GRA	SPO	
Lactobacillus catenaformis		41.99	0.39	MNS 99	99	1	
Clostridium barati		31.13	0.32	MLT 90	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		23.46	0.45		1	1		
03251121	**Low discrimination**							**03251121**
					SPO			
Actinomyces naeslundii		69.27	0.37	MNS 88	1			
Lactobacillus catenaformis		17.65	0.05	MLT 94	1			
Clostridium barati		13.09	-.02	RAF 92	99			
03251122	**Low discrimination**							**03251122**
					SPO			
Clostridium barati		51.16	0.40	ESL 87	99			
Lactobacillus acidophilus		23.43	0.36	MNS 99	1			
Lactobacillus catenaformis		23.00	0.39		1			
03251123	**Low discrimination**							**03251123**
					SPO			
Actinomyces naeslundii		48.05	0.37	MNS 88	1			
Clostridium barati		27.23	0.06	MLT 94	99			
Lactobacillus acidophilus		12.47	0.02	RAF 92	1			
03251140	**Acceptable identification**							**03251140**
					GRA	SPO		
Fusobacterium mortiferum		90.03	0.45	SOR 1	1	1		
Lactobacillus catenaformis		6.51	0.15		99	1		
Clostridium barati		3.46	0.05		99	99		
03251200	**Low discrimination**							**03251200**
					GRA	SPO		
Fusobacterium mortiferum		85.53	0.39	RHA 1	1	1		
Lactobacillus catenaformis		9.62	0.12		99	1		
Clostridium barati		2.56	-.03		99	99		
03251300	**Acceptable identification**							**03251300**
					GRA	SPO		
Fusobacterium mortiferum		90.03	0.45	RHA 1	1	1		
Lactobacillus catenaformis		6.51	0.15		99	1		
Clostridium barati		3.46	0.05		99	99		
03251400	**Low discrimination**							**03251400**
					GRA	SPO		
Fusobacterium mortiferum		62.50	0.39	ARA 1	1	1		
Clostridium innocuum		23.77	0.19		99	99		
Lactobacillus catenaformis		7.03	0.12		99	1		
03251500	**Low discrimination**							**03251500**
					GRA			
Fusobacterium mortiferum		85.55	0.45	ARA 1	1			
Lactobacillus catenaformis		6.19	0.15		99			
Actinomyces meyeri		5.83	0.16		99			
03252000	**Low discrimination**							**03252000**
					COC	GRA	SPO	
Clostridium innocuum		86.55	0.44	MNS 99	1	99	99	
Megasphaera elsdenii		6.74	0.29	GAL 95	99	1	1	
Fusobacterium mortiferum		5.14	0.36		1	1	1	
03252002	**Good identification**							**03252002**
					COC	GRA	SPO	
Clostridium innocuum		98.50	0.44	MNS 99	1	99	99	
Megasphaera elsdenii		0.85	0.12	GAL 95	99	1	1	
Fusobacterium mortiferum		0.65	0.19		1	1	1	
03252100	**Acceptable identification**							**03252100**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		93.20	0.42	MAN 1	1	1	
Clostridium innocuum		6.80	0.10		99	99	
03253000	**Very good identification**						**03253000**
					GRA	SPO	
Clostridium innocuum		99.42	0.66	MNS 99	99	99	
Fusobacterium mortiferum		0.47	0.39		1	1	
Lactobacillus catenaformis		0.05	0.12		99	1	
03253002	**Very good identification**						**03253002**
					GRA	SPO	
Clostridium innocuum		99.82	0.66	MNS 99	99	99	
Lactobacillus catenaformis		0.05	0.12		99	1	
Fusobacterium mortiferum		0.05	0.22		1	1	
03253010	**Good identification**						**03253010**
Clostridium innocuum		100.0	0.42	MNS 99			
03253012	**Good identification**						**03253012**
Clostridium innocuum		100.0	0.42	MNS 99			
03253100	**Low discrimination**						**03253100**
					GRA	SPO	
Fusobacterium mortiferum		46.79	0.45	MAN 1	1	1	
Clostridium innocuum		43.22	0.32		99	99	
Actinomyces naeslundii		4.30	0.39		99	1	
03253101	**Good identification**						**03253101**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03253103	**Good identification**						**03253103**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03253121	**Good identification**						**03253121**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03253123	**Good identification**						**03253123**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03253400	**Good identification**						**03253400**
Clostridium innocuum		100.0	0.53	MNS 99			
03253402	**Good identification**						**03253402**
Clostridium innocuum		100.0	0.53	MNS 99			
03254000	**Good identification**						**03254000**
					GRA		
Fusobacterium mortiferum		96.02	0.50	XYL 7	1		
Actinomyces meyeri		3.98	0.18		99		
03254100	**Good identification**						**03254100**
					GRA		
Fusobacterium mortiferum		93.37	0.57	XYL 7	1		
Actinomyces meyeri		6.63	0.28		99		
03254102	**Good identification**						**03254102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		98.06	0.40	XYL 7	1		
Actinomyces meyeri		1.94	0.02	BGL 10	99		
03255000	**Acceptable identification**						**03255000**
					GRA	SPO	
Fusobacterium mortiferum		92.72	0.53	XYL 7	1	1	
Actinomyces meyeri		3.84	0.21		99	1	
Clostridium innocuum		1.41	0.10		99	99	
03255002	**Low discrimination**						**03255002**
					GRA	SPO	
Fusobacterium mortiferum		76.82	0.37	XYL 7	1	1	
Clostridium innocuum		10.53	0.10	BGL 10	99	99	
Lactobacillus catenaformis		10.44	0.12		99	1	
03255100	**Good identification**						**03255100**
					GRA		
Fusobacterium mortiferum		92.08	0.60	XYL 7	1		
Actinomyces meyeri		6.54	0.31		99		
Lactobacillus catenaformis		0.89	0.15		99		
03255102	**Low discrimination**						**03255102**
					GRA		
Fusobacterium mortiferum		81.95	0.43	XYL 7	1		
Lactobacillus catenaformis		7.16	0.15	BGL 10	99		
Actinomyces israelii		3.34	0.22		99		
03257000	**Good identification**						**03257000**
					GRA	SPO	
Clostridium innocuum		98.69	0.44	MNS 99	99	99	
Fusobacterium mortiferum		0.66	0.19	XYL 5	1	1	
Actinomyces israelii		0.65	0.22		99	1	
03257002	**Good identification**						**03257002**
					SPO		
Clostridium innocuum		99.02	0.44	MNS 99	99		
Actinomyces israelii		0.98	0.25	XYL 5	1		
03261100	**Good identification**						**03261100**
Fusobacterium mortiferum		100.0	0.36	FRU 99			
03263000	**Good identification**						**03263000**
Clostridium innocuum		100.0	0.36	MNS 99			
03263002	**Good identification**						**03263002**
Clostridium innocuum		100.0	0.36	MNS 99			
03270000	**Very good identification**						**03270000**
					GRA	SPO	
Fusobacterium mortiferum		99.39	0.61		1	1	
Lactobacillus jensenii		0.41	0.10		99	1	
Clostridium innocuum		0.20	0.02		99	99	
03270002	**Good identification**						**03270002**
					GRA	SPO	
Fusobacterium mortiferum		95.91	0.44	BGL 10	1	1	
Lactobacillus jensenii		2.34	0.07		99	1	
Clostridium innocuum		1.75	0.02		99	99	
03270100	**Very good identification**						**03270100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.67				
03270102	**Good identification**						**03270102**
					GRA		
Fusobacterium mortiferum		97.26	0.51	BGL 10	1		
Lactobacillus acidophilus		2.74	0.11		99		
03271000	**Low discrimination**						**03271000**
					GRA	SPO	
Fusobacterium mortiferum		76.83	0.64		1	1	
Lactobacillus jensenii		20.64	0.44		99	1	
Clostridium innocuum		1.97	0.24		99	99	
03271002	**Low discrimination**						**03271002**
					GRA		
Lactobacillus jensenii		48.79	0.41	MNS 99	99		
Fusobacterium mortiferum		30.27	0.47	MLT 95	1		
Lactobacillus acidophilus		11.41	0.27		99		
03271010	**Low discrimination**						**03271010**
					GRA	SPO	
Lactobacillus jensenii		88.91	0.36	MNS 99	99	1	
Fusobacterium mortiferum		10.03	0.30	MLT 95	1	1	
Clostridium innocuum		1.06	0.00		99	99	
03271100	**Very good identification**						**03271100**
					COC	GRA	
Fusobacterium mortiferum		98.45	0.70		1	1	
Atopobium parvulum		0.56	0.20		99	99	
Actinomyces naeslundii		0.31	0.39		1	99	
03271101	**Low discrimination**						**03271101**
					GRA		
Fusobacterium mortiferum		68.44	0.36	URE 1	1		
Actinomyces naeslundii		31.56	0.42		99		
03271102	**Low discrimination**						**03271102**
					GRA	SPO	
Lactobacillus acidophilus		62.71	0.45	MNS 99	99	1	
Fusobacterium mortiferum		33.77	0.54	MLT 94	1	1	
Clostridium barati		1.19	0.14		99	99	
03271103	**Low discrimination**						**03271103**
					GRA		
Actinomyces naeslundii		59.22	0.42	MNS 88	99		
Lactobacillus acidophilus		26.51	0.11	MLT 94	99		
Fusobacterium mortiferum		14.27	0.20	RAF 92	1		
03271110	**Good identification**						**03271110**
					GRA		
Fusobacterium mortiferum		96.30	0.36	MLZ 1	1		
Lactobacillus jensenii		3.70	0.02		99		
03271120	**Low discrimination**						**03271120**
					GRA	SPO	
Fusobacterium mortiferum		44.99	0.36	NAG 1	1	1	
Clostridium barati		17.38	0.14		99	99	
Actinomyces naeslundii		13.83	0.39		99	1	
03271121	**Good identification**						**03271121**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03271122	**Acceptable identification**						**03271122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Lactobacillus acidophilus		92.22	0.45	MNS 99	1		
Clostridium barati		5.23	0.22	MLT 94	99		
Actinomyces naeslundii		1.39	0.39		1		
03271123	**Low discrimination**						**03271123**
Actinomyces naeslundii		69.08	0.42	MNS 88			
Lactobacillus acidophilus		30.92	0.11	MLT 94			
03271140	**Acceptable identification**						**03271140**
					GRA		
Fusobacterium mortiferum		94.86	0.36	SOR 1	1		
Actinomyces naeslundii		5.15	0.26		99		
03271300	**Good identification**						**03271300**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
03271500	**Good identification**						**03271500**
Fusobacterium mortiferum		100.0	0.36	ARA 1			
03272000	**Good identification**						**03272000**
					GRA	SPO	
Clostridium innocuum		95.20	0.36	MNS 99	99	99	
Fusobacterium mortiferum		4.81	0.27	GAL 95	1	1	
03272002	**Good identification**						**03272002**
Clostridium innocuum		100.0	0.36	MNS 99			
03273000	**Very good identification**						**03273000**
					GRA	SPO	
Clostridium innocuum		99.27	0.58	MNS 99	99	99	
Fusobacterium mortiferum		0.40	0.30		1	1	
Actinomyces naeslundii		0.19	0.36		99	1	
03273002	**Very good identification**						**03273002**
					SPO		
Clostridium innocuum		99.56	0.58	MNS 99	99		
Actinomyces naeslundii		0.19	0.36		1		
Lactobacillus acidophilus		0.09	0.05		1		
03273021	**Good identification**						**03273021**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03273023	**Good identification**						**03273023**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03273101	**Good identification**						**03273101**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03273103	**Good identification**						**03273103**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03273120	**Low discrimination**						**03273120**
					SPO		
Actinomyces naeslundii		70.27	0.44	MNS 88	1		
Clostridium tertium		15.31	0.09	MLT 94	99		
Clostridium ramosum		14.43	0.07	RAF 92	92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03273121	**Good identification**						**03273121**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03273123	**Good identification**						**03273123**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03273400	**Good identification**						**03273400**
Clostridium innocuum		100.0	0.45	MNS 99			
03273402	**Good identification**						**03273402**
					SPO		
Clostridium innocuum		99.78	0.45	MNS 99	99		
Actinomyces israelii		0.23	0.15		1		
03274000	**Good identification**						**03274000**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
03274100	**Good identification**						**03274100**
Fusobacterium mortiferum		100.0	0.48	XYL 7			
03275000	**Acceptable identification**						**03275000**
					GRA		
Fusobacterium mortiferum		92.92	0.44	XYL 7	1		
Lactobacillus jensenii		3.35	0.10		99		
Actinomyces israelii		2.06	0.19		99		
03275100	**Low discrimination**						**03275100**
					COC	GRA	
Fusobacterium mortiferum		84.95	0.51	XYL 7	1	1	
Atopobium parvulum		11.82	0.25		99	99	
Actinomyces israelii		3.23	0.29		1	99	
03277000	**Acceptable identification**						**03277000**
					SPO		
Clostridium innocuum		93.38	0.36	MNS 99	99		
Actinomyces israelii		6.62	0.31	XYL 5	1		
03277002	**Acceptable identification**						**03277002**
					SPO		
Clostridium innocuum		90.38	0.36	MNS 99	99		
Actinomyces israelii		9.62	0.34	XYL 5	1		
03277100	**Low discrimination**						**03277100**
					GRA	SPO	
Actinomyces israelii		89.31	0.42	MLT 99	99	1	
Fusobacterium mortiferum		4.19	0.17		1	1	
Clostridium tertium		3.33	0.03		99	99	
03277102	**Acceptable identification**						**03277102**
					SPO		
Actinomyces israelii		91.06	0.45	MLT 99	1		
Clostridium tertium		6.78	0.11		99		
Clostridium innocuum		2.16	0.02		99		
03277142	**Good identification**						**03277142**
Actinomyces israelii		100.0	0.38	MLT 99			
03277500	**Good identification**						**03277500**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		100.0	0.40	MLT 99				
03277502	**Acceptable identification**							**03277502**
Actinomyces israelii		93.41	0.43	MLT 99				
Bifidobacterium adolescentis		6.59	0.17					
03277542	**Good identification**							**03277542**
Actinomyces israelii		100.0	0.36	MLT 99				
03300000	**Low discrimination**							**03300000**
					COC	GRA		
Clostridium botulinum biovar A		24.89	0.41	SUC 7	1	99		
Megasphaera elsdenii		22.19	0.44	SAL 1	99	1		
Clostridium novyi biovar A		13.35	0.48		1	99		
03300001	**Acceptable identification**							**03300001**
					COC	GRA		
Clostridium novyi biovar A		94.05	0.48	SUC 1	1	99		
Clostridium botulinum biovar A		1.77	0.06	SAL 8	1	99		
Megasphaera elsdenii		1.58	0.10		99	1		
03300002	**Low discrimination**							**03300002**
					COC	SPO	GRA	
Actinomyces odontolyticus		55.78	0.39	NIT 99	1	1	99	
Megasphaera elsdenii		18.29	0.28	FRU 92	99	1	1	
Clostridium novyi biovar A		17.48	0.35		1	99	99	
03300010	**Good identification**							**03300010**
					COC	GRA		
Clostridium novyi biovar A		95.05	0.48	SUC 1	1	99		
Clostridium botulinum biovar A		1.79	0.06	SAL 8	1	99		
Megasphaera elsdenii		1.60	0.10		99	1		
03300011	**Good identification**							**03300011**
Clostridium novyi biovar A		100.0	0.48	SUC 1				
03300020	**Low discrimination**							**03300020**
					COC	SPO		
Clostridium novyi biovar A		47.94	0.37	SUC 1	1	99		
Actinomyces odontolyticus		28.69	0.29	SAL 8	1	1		
Gemella morbillorum		6.74	0.14		99	1		
03300021	**Good identification**							**03300021**
Clostridium novyi biovar A		100.0	0.37	SUC 1				
03300030	**Good identification**							**03300030**
Clostridium novyi biovar A		100.0	0.37	SUC 1				
03300031	**Good identification**							**03300031**
Clostridium novyi biovar A		100.0	0.37	SUC 1				
03300040	**Low discrimination**							**03300040**
					COC	GRA		
Megasphaera elsdenii		45.63	0.41	SAL 1	99	1		
Clostridium novyi biovar A		27.46	0.45	FRU 95	1	99		
Clostridium botulinum biovar A		22.94	0.32		1	99		
03300041	**Good identification**							**03300041**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		98.35	0.45	SUC 1	1	99		
Megasphaera elsdenii		1.65	0.07	SAL 8	99	1		
03300050	**Acceptable identification**							**03300050**
					COC	GRA		
Clostridium novyi biovar A		93.65	0.45	SUC 1	1	99		
Clostridium botulinum biovar B		4.78	0.08	SAL 8	1	99		
Megasphaera elsdenii		1.57	0.07		99	1		
03300051	**Good identification**							**03300051**
Clostridium novyi biovar A		100.0	0.45	SUC 1				
03300100	**Low discrimination**							**03300100**
					GRA			
Actinomyces meyeri		55.64	0.36	SAL 1	99			
Actinomyces odontolyticus		26.63	0.34	FRU 95	99			
Fusobacterium mortiferum		12.88	0.34		1			
03300200	**Good identification**							**03300200**
					COC	SPO	GRA	
Actinomyces odontolyticus		97.12	0.48	NIT 99	1	1	99	
Clostridium botulinum biovar A		1.03	0.06	FRU 92	1	99	99	
Megasphaera elsdenii		0.91	0.10		99	1	1	
03300202	**Good identification**							**03300202**
Actinomyces odontolyticus		100.0	0.44	NIT 99				
03300300	**Good identification**							**03300300**
Actinomyces odontolyticus		98.92	0.39	NIT 99				
Actinomyces meyeri		1.08	0.02	FRU 92				
03300302	**Good identification**							**03300302**
Actinomyces odontolyticus		100.0	0.36	NIT 99				
03301002	**Low discrimination**							**03301002**
Actinomyces odontolyticus		61.60	0.36	NIT 99				
Eubacterium tortuosum		26.55	0.19	FRU 92				
Lactobacillus fermentum		4.88	0.24					
03301100	**Low discrimination**							**03301100**
					GRA			
Actinomyces meyeri		64.61	0.39	SAL 1	99			
Fusobacterium mortiferum		14.95	0.37	FRU 95	1			
Actinomyces odontolyticus		13.75	0.31		99			
03301101	**Good identification**							**03301101**
					SPO			
Clostridium chauvoei		96.53	0.37	MNS 90	99			
Actinomyces naeslundii		2.54	0.33	SAL 1	1			
Actinomyces meyeri		0.93	0.05		1			
03301200	**Good identification**							**03301200**
Actinomyces odontolyticus		99.23	0.45	NIT 99				
Eubacterium tortuosum		0.77	-.03	FRU 92				
03301202	**Good identification**							**03301202**
Actinomyces odontolyticus		100.0	0.41	NIT 99				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
03301300	**Good identification**						**03301300**
Actinomyces odontolyticus		97.61	0.36	NIT 99			
Actinomyces meyeri		2.39	0.05	FRU 92			
03302000	**Good identification**						**03302000**
					COC	GRA	
Megasphaera elsdenii		97.34	0.44	SAL 1	99	1	
Clostridium botulinum biovar A		1.10	0.06	FRU 95	1	99	
Clostridium novyi biovar A		0.59	0.14		1	99	
03302040	**Good identification**						**03302040**
					COC	GRA	
Megasphaera elsdenii		99.40	0.41	SAL 1	99	1	
Clostridium novyi biovar A		0.60	0.11	FRU 95	1	99	
03303100	**Low discrimination**						**03303100**
Actinomyces naeslundii		64.32	0.35	MNS 88			
Actinomyces meyeri		18.23	0.05	RAF 92			
Propionibacterium propionicum		17.45	0.15	FRU 95			
03303101	**Low discrimination**						**03303101**
					SPO		
Actinomyces naeslundii		83.49	0.38	MNS 88	1		
Clostridium chauvoei		16.51	0.03	RAF 92	99		
03303102	**Low discrimination**						**03303102**
Actinomyces naeslundii		71.08	0.35	MNS 88			
Propionibacterium propionicum		28.92	0.18	RAF 92			
03303103	**Good identification**						**03303103**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03303120	**Good identification**						**03303120**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03303121	**Good identification**						**03303121**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03303122	**Good identification**						**03303122**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03303123	**Good identification**						**03303123**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03304000	**Low discrimination**						**03304000**
					COC		
Actinomyces odontolyticus		54.22	0.38	NIT 99	1		
Actinomyces meyeri		35.24	0.31	FRU 92	1		
Gemella morbillorum		4.21	0.14		99		
03304100	**Low discrimination**						**03304100**
					GRA		
Actinomyces meyeri		88.46	0.41	SAL 1	99		
Actinomyces odontolyticus		10.75	0.29	FRU 95	99		
Fusobacterium mortiferum		0.79	0.15		1		
03304200	**Good identification**						**03304200**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces odontolyticus		100.0	0.42	NIT 99				
03304202	**Good identification**							**03304202**
Actinomyces odontolyticus		100.0	0.39	NIT 99				
03305100	**Acceptable identification**							**03305100**
Actinomyces meyeri		92.67	0.44	SAL 1				
Actinomyces odontolyticus		5.00	0.26	FRU 95				
Lactobacillus fermentum		1.50	0.24					
03305200	**Good identification**							**03305200**
Actinomyces odontolyticus		100.0	0.39	NIT 99				
03305202	**Good identification**							**03305202**
Actinomyces odontolyticus		100.0	0.36	NIT 99				
03310000	**Low discrimination**							**03310000**
					COC	GRA		
Megasphaera elsdenii		59.97	0.66	SAL 1	99	1		
Actinomyces odontolyticus		20.07	0.61		1	99		
Fusobacterium mortiferum		11.30	0.62		1	1		
03310002	**Low discrimination**							**03310002**
					COC	GRA		
Actinomyces odontolyticus		59.86	0.57	NIT 99	1	99		
Megasphaera elsdenii		32.42	0.50		99	1		
Fusobacterium mortiferum		6.11	0.46		1	1		
03310020	**Low discrimination**							**03310020**
					COC	GRA		
Actinomyces odontolyticus		66.51	0.47	NIT 99	1	99		
Actinomyces meyeri		15.18	0.33	NAG 14	1	99		
Megasphaera elsdenii		12.33	0.32		99	1		
03310022	**Acceptable identification**							**03310022**
					COC	GRA		
Actinomyces odontolyticus		94.26	0.44	NIT 99	1	99		
Megasphaera elsdenii		3.17	0.15	NAG 14	99	1		
Actinomyces meyeri		1.09	0.07		1	99		
03310040	**Very good identification**							**03310040**
					COC	SPO	GRA	
Megasphaera elsdenii		97.16	0.63	SAL 1	99	1	1	
Clostridium botulinum biovar B		1.28	0.24		1	99	99	
Actinomyces odontolyticus		0.49	0.27		1	1	99	
03310042	**Good identification**							**03310042**
					COC	SPO	GRA	
Megasphaera elsdenii		96.49	0.47	SAL 1	99	1	1	
Actinomyces odontolyticus		2.70	0.23	BGL 10	1	1	99	
Clostridium novyi biovar A		0.54	0.15		1	99	99	
03310100	**Low discrimination**							**03310100**
					GRA			
Fusobacterium mortiferum		47.67	0.69	MLT 11	1			
Actinomyces meyeri		39.54	0.58		99			
Actinomyces odontolyticus		11.46	0.52		99			
03310102	**Low discrimination**							**03310102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces odontolyticus		49.99	0.49	NIT 99	99		
Fusobacterium mortiferum		37.71	0.52		1		
Actinomyces meyeri		8.71	0.32		99		
03310120	**Low discrimination**						**03310120**
					GRA		
Actinomyces meyeri		66.79	0.43	SAL 1	99		
Actinomyces odontolyticus		23.10	0.39	NAG 12	99		
Fusobacterium mortiferum		5.97	0.34		1		
03310122	**Low discrimination**						**03310122**
Actinomyces odontolyticus		72.64	0.35	NIT 99			
Actinomyces meyeri		10.60	0.17	NAG 14			
Clostridium perfringens		5.68	0.11				
03310200	**Very good identification**						**03310200**
					COC	GRA	
Actinomyces odontolyticus		98.05	0.66	NIT 99	1	99	
Megasphaera elsdenii		1.52	0.32		99	1	
Fusobacterium mortiferum		0.29	0.28		1	1	
03310202	**Very good identification**						**03310202**
					COC	GRA	
Actinomyces odontolyticus		99.67	0.62	NIT 99	1	99	
Megasphaera elsdenii		0.28	0.15		99	1	
Fusobacterium mortiferum		0.05	0.12		1	1	
03310220	**Good identification**						**03310220**
Actinomyces odontolyticus		100.0	0.52	NIT 99			
03310222	**Good identification**						**03310222**
Actinomyces odontolyticus		100.0	0.49	NIT 99			
03310300	**Very good identification**						**03310300**
					GRA		
Actinomyces odontolyticus		96.19	0.57	NIT 99	99		
Fusobacterium mortiferum		2.08	0.34		1		
Actinomyces meyeri		1.73	0.24		99		
03310302	**Good identification**						**03310302**
					GRA		
Actinomyces odontolyticus		99.61	0.54	NIT 99	99		
Fusobacterium mortiferum		0.39	0.18		1		
03310320	**Good identification**						**03310320**
Actinomyces odontolyticus		98.52	0.44	NIT 99			
Actinomyces meyeri		1.48	0.09	NAG 14			
03310322	**Good identification**						**03310322**
Actinomyces odontolyticus		100.0	0.40	NIT 99			
03310400	**Low discrimination**						**03310400**
					COC	GRA	
Actinomyces meyeri		56.31	0.37	SAL 1	1	99	
Megasphaera elsdenii		24.95	0.32		99	1	
Actinomyces odontolyticus		8.35	0.27		1	99	
03310500	**Acceptable identification**						**03310500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces meyeri		92.83	0.48	SAL 1	99		
Fusobacterium mortiferum		4.52	0.34		1		
Lactobacillus fermentum		1.56	0.28		99		
03311000	**Low discrimination**						**03311000**
					GRA		
Fusobacterium mortiferum		35.18	0.65	MLT 11	1		
Actinomyces odontolyticus		27.77	0.58		99		
Actinomyces meyeri		17.02	0.51		99		
03311001	**Low discrimination**						**03311001**
					GRA		
Actinomyces naeslundii		37.73	0.47	MNS 88	99		
Fusobacterium mortiferum		20.35	0.31	RAF 92	1		
Actinomyces odontolyticus		16.06	0.24		99		
03311002	**Low discrimination**						**03311002**
					GRA		
Actinomyces odontolyticus		46.64	0.54	NIT 99	99		
Lactobacillus fermentum		31.78	0.59		99		
Fusobacterium mortiferum		10.71	0.49		1		
03311003	**Low discrimination**						**03311003**
Actinomyces naeslundii		66.73	0.47	MNS 88			
Actinomyces odontolyticus		17.41	0.20	RAF 92			
Lactobacillus fermentum		11.86	0.24				
03311020	**Low discrimination**						**03311020**
Actinomyces odontolyticus		29.14	0.44	NIT 99			
Lactobacillus fermentum		24.92	0.50	NAG 14			
Clostridium perfringens		18.97	0.36				
03311021	**Low discrimination**						**03311021**
Actinomyces naeslundii		79.75	0.47	MNS 88			
Actinomyces odontolyticus		5.53	0.10	RAF 92			
Lactobacillus fermentum		4.72	0.16				
03311022	**Low discrimination**						**03311022**
Lactobacillus fermentum		40.65	0.50	SAL 1			
Actinomyces odontolyticus		29.14	0.41				
Clostridium perfringens		14.56	0.31				
03311023	**Acceptable identification**						**03311023**
Actinomyces naeslundii		90.77	0.47	MNS 88			
Lactobacillus fermentum		5.38	0.16	RAF 92			
Actinomyces odontolyticus		3.86	0.07				
03311100	**Low discrimination**						**03311100**
					GRA		
Fusobacterium mortiferum		45.16	0.72	MLT 11	1		
Actinomyces meyeri		37.46	0.61		99		
Lactobacillus fermentum		10.41	0.62		99		
03311101	**Low discrimination**						**03311101**
					GRA	SPO	
Clostridium chauvoei		70.43	0.44	MNS 90	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		15.10	0.55	SAL 1	99	1	
Fusobacterium mortiferum		6.68	0.37		1	1	
03311102	**Low discrimination**						**03311102**
					GRA		
Lactobacillus fermentum		47.41	0.62	SAL 1	99		
Fusobacterium mortiferum		22.86	0.55		1		
Actinomyces odontolyticus		13.47	0.46		99		
03311103	**Low discrimination**						**03311103**
					SPO		
Actinomyces naeslundii		49.53	0.55	MNS 88	1		
Clostridium chauvoei		40.77	0.31	RAF 92	99		
Lactobacillus fermentum		5.05	0.28		1		
03311110	**Low discrimination**						**03311110**
					GRA		
Fusobacterium mortiferum		44.52	0.37	MLT 11	1		
Actinomyces meyeri		36.93	0.27	MLZ 1	99		
Lactobacillus fermentum		10.26	0.28		99		
03311120	**Low discrimination**						**03311120**
Clostridium perfringens		46.36	0.51	MNS 99			
Actinomyces meyeri		19.95	0.46				
Lactobacillus fermentum		13.55	0.54				
03311121	**Low discrimination**						**03311121**
					SPO		
Clostridium chauvoei		55.89	0.35	MNS 90	99		
Actinomyces naeslundii		35.95	0.55	SAL 1	1		
Clostridium perfringens		4.18	0.17		84		
03311122	**Low discrimination**						**03311122**
Clostridium perfringens		44.74	0.45	MNS 99			
Lactobacillus fermentum		27.78	0.54				
Actinomyces naeslundii		5.51	0.52				
03311123	**Low discrimination**						**03311123**
					SPO		
Actinomyces naeslundii		73.37	0.55	MNS 88	1		
Clostridium chauvoei		20.13	0.22	RAF 92	99		
Clostridium perfringens		4.01	0.11		84		
03311140	**Low discrimination**						**03311140**
					GRA		
Fusobacterium mortiferum		38.05	0.37	MLT 11	1		
Actinomyces meyeri		31.56	0.27	SOR 1	99		
Actinomyces naeslundii		10.13	0.39		99		
03311141	**Low discrimination**						**03311141**
					SPO		
Actinomyces naeslundii		78.93	0.42	MNS 88	1		
Clostridium chauvoei		21.08	0.09	RAF 92	99		
03311142	**Low discrimination**						**03311142**
					GRA		
Actinomyces naeslundii		29.43	0.39	MNS 88	99		
Lactobacillus fermentum		25.49	0.28	RAF 92	99		
Fusobacterium mortiferum		12.29	0.21		1		
03311143	**Good identification**						**03311143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	MNS 88		
03311160	**Acceptable identification**					**03311160**
Clostridium perfringens		90.56	0.39	MNS 99		
Actinomyces naeslundii		4.86	0.39			
Actinomyces meyeri		2.07	0.12			
03311161	**Low discrimination**					**03311161**
					SPO	
Actinomyces naeslundii		82.34	0.42	MNS 88	1	
Clostridium perfringens		10.33	0.04	RAF 92	84	
Clostridium chauvoei		7.33	0.01		99	
03311163	**Good identification**					**03311163**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03311200	**Very good identification**					**03311200**
					GRA	
Actinomyces odontolyticus		98.77	0.63	NIT 99	99	
Fusobacterium mortiferum		0.65	0.31		1	
Actinomyces meyeri		0.32	0.16		99	
03311202	**Very good identification**					**03311202**
					GRA	
Actinomyces odontolyticus		99.53	0.59	NIT 99	99	
Lactobacillus fermentum		0.35	0.24		99	
Fusobacterium mortiferum		0.12	0.15		1	
03311220	**Good identification**					**03311220**
Actinomyces odontolyticus		98.96	0.49	NIT 99		
Lactobacillus fermentum		0.44	0.16	NAG 14		
Clostridium perfringens		0.34	0.02			
03311222	**Good identification**					**03311222**
					GRA	
Actinomyces odontolyticus		98.73	0.46	NIT 99	99	
Lactobacillus fermentum		0.72	0.16	NAG 14	99	
Prevotella oralis		0.55	-.01		1	
03311300	**Low discrimination**					**03311300**
					GRA	
Actinomyces odontolyticus		81.76	0.54	NIT 99	99	
Eubacterium contortum		10.03	0.32		99	
Fusobacterium mortiferum		3.98	0.37		1	
03311302	**Acceptable identification**					**03311302**
Actinomyces odontolyticus		93.50	0.51	NIT 99		
Eubacterium contortum		3.30	0.19			
Lactobacillus fermentum		1.71	0.28			
03311320	**Low discrimination**					**03311320**
Actinomyces odontolyticus		77.18	0.41	NIT 99		
Eubacterium contortum		10.27	0.19	NAG 14		
Clostridium perfringens		6.07	0.17			
03311322	**Low discrimination**					**03311322**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		68.33	0.37	NIT 99	99		
Prevotella oralis		18.87	0.19	NAG 14	1		
Clostridium perfringens		4.13	0.11		99		
03311400	**Low discrimination**						**03311400**
					GRA		
Lactobacillus fermentum		69.83	0.59	SAL 1	99		
Actinomyces meyeri		25.63	0.40		99		
Fusobacterium mortiferum		2.14	0.31		1		
03311402	**Very good identification**						**03311402**
Lactobacillus fermentum		97.13	0.59	SAL 1			
Actinomyces odontolyticus		1.44	0.20				
Actinomyces meyeri		1.10	0.15				
03311420	**Low discrimination**						**03311420**
Lactobacillus fermentum		85.50	0.50	SAL 1			
Actinomyces meyeri		12.84	0.26				
Actinomyces odontolyticus		1.01	0.10				
03311422	**Good identification**						**03311422**
Lactobacillus fermentum		99.28	0.50	SAL 1			
Actinomyces odontolyticus		0.72	0.07				
03311500	**Low discrimination**						**03311500**
					GRA		
Lactobacillus fermentum		51.29	0.62	SAL 1	99		
Actinomyces meyeri		46.15	0.51		99		
Fusobacterium mortiferum		2.25	0.37		1		
03311502	**Good identification**						**03311502**
Lactobacillus fermentum		90.88	0.62	SAL 1			
Bifidobacterium adolescentis		5.70	0.36				
Actinomyces meyeri		2.53	0.25				
03311512	**Low discrimination**						**03311512**
Bifidobacterium adolescentis		86.74	0.38	MNS 87			
Lactobacillus fermentum		11.43	0.28	RAF 92			
Bifidobacterium longum subsp. longum		1.84	0.07	XYL 87			
03311520	**Low discrimination**						**03311520**
Lactobacillus fermentum		70.32	0.54	SAL 1			
Actinomyces meyeri		25.89	0.36				
Clostridium perfringens		2.43	0.17				
03311522	**Good identification**						**03311522**
Lactobacillus fermentum		96.24	0.54	SAL 1			
Clostridium perfringens		1.57	0.11				
Bifidobacterium longum subsp. longum		1.10	0.14				
03312000	**Very good identification**						**03312000**
					COC	GRA	
Megasphaera elsdenii		99.24	0.66	SAL 1	99	1	
Actinomyces odontolyticus		0.34	0.27		1	99	
Fusobacterium mortiferum		0.19	0.28		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03312002	**Good identification**						**03312002**
					COC	GRA	
Megasphaera elsdenii		97.38	0.50	SAL 1	99	1	
Actinomyces odontolyticus		1.82	0.23	BGL 10	1	99	
Propionibacterium propionicum		0.37	0.18		1	99	
03312040	**Very good identification**						**03312040**
					COC	GRA	
Megasphaera elsdenii		99.96	0.63	SAL 1	99	1	
Propionibacterium propionicum		0.04	0.15		1	99	
03312042	**Good identification**						**03312042**
					COC	GRA	
Megasphaera elsdenii		99.44	0.47	SAL 1	99	1	
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99	
03312100	**Low discrimination**						**03312100**
					COC	GRA	
Propionibacterium propionicum		57.82	0.49	NIT 86	1	99	
Megasphaera elsdenii		21.19	0.32	SAL 1	99	1	
Fusobacterium mortiferum		9.32	0.34		1	1	
03312102	**Acceptable identification**						**03312102**
					COC	GRA	
Propionibacterium propionicum		93.06	0.52	NIT 86	1	99	
Megasphaera elsdenii		2.53	0.15	SAL 1	99	1	
Actinomyces naeslundii		1.83	0.33		1	99	
03312103	**Low discrimination**						**03312103**
Propionibacterium propionicum		79.03	0.36	NIT 86			
Actinomyces naeslundii		20.97	0.36	SAL 1			
03312121	**Good identification**						**03312121**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03312123	**Good identification**						**03312123**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03312140	**Low discrimination**						**03312140**
					COC	GRA	
Propionibacterium propionicum		80.37	0.49	NIT 86	1	99	
Megasphaera elsdenii		19.63	0.29	SAL 1	99	1	
03312142	**Good identification**						**03312142**
					COC	GRA	
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99	
Megasphaera elsdenii		1.78	0.12	SAL 1	99	1	
03312143	**Good identification**						**03312143**
Propionibacterium propionicum		95.53	0.36	NIT 86			
Actinomyces naeslundii		4.47	0.24	SAL 1			
03313001	**Good identification**						**03313001**
Actinomyces naeslundii		100.0	0.52	MNS 88			
03313002	**Low discrimination**						**03313002**
					COC	GRA	
Actinomyces naeslundii		59.13	0.49	MNS 88	1	99	
Actinomyces odontolyticus		11.92	0.20	RAF 92	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		9.69	0.15		99	1	
03313003	**Good identification**						**03313003**
Actinomyces naeslundii		100.0	0.52	MNS 88			
03313020	**Low discrimination**						**03313020**
Actinomyces naeslundii		86.08	0.49	MNS 88			
Actinomyces odontolyticus		4.61	0.10	RAF 92			
Lactobacillus fermentum		3.94	0.16				
03313021	**Good identification**						**03313021**
Actinomyces naeslundii		100.0	0.52	MNS 88			
03313022	**Acceptable identification**						**03313022**
Actinomyces naeslundii		92.71	0.49	MNS 88			
Lactobacillus fermentum		4.24	0.16	RAF 92			
Actinomyces odontolyticus		3.04	0.07				
03313023	**Good identification**						**03313023**
Actinomyces naeslundii		100.0	0.52	MNS 88			
03313041	**Good identification**						**03313041**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03313042	**Low discrimination**						**03313042**
					COC	GRA	
Actinomyces naeslundii		46.66	0.36	MNS 88	1	99	
Megasphaera elsdenii		28.87	0.12	RAF 92	99	1	
Propionibacterium propionicum		16.12	0.18		1	99	
03313043	**Good identification**						**03313043**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03313060	**Good identification**						**03313060**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03313061	**Good identification**						**03313061**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03313062	**Good identification**						**03313062**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03313063	**Good identification**						**03313063**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03313100	**Low discrimination**						**03313100**
					GRA		
Propionibacterium propionicum		43.76	0.49	NIT 86	99		
Actinomyces naeslundii		30.96	0.57	SAL 1	99		
Fusobacterium mortiferum		10.58	0.37		1		
03313101	**Low discrimination**						**03313101**
					SPO		
Actinomyces naeslundii		88.58	0.60	MNS 88	1		
Propionibacterium propionicum		9.28	0.33	RAF 92	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		2.15	0.09		99	
03313102	**Low discrimination**					**03313102**
Propionibacterium propionicum		61.51	0.52	NIT 86		
Actinomyces naeslundii		29.01	0.57	SAL 1		
Actinomyces israelii		3.31	0.32			
03313103	**Low discrimination**					**03313103**
Actinomyces naeslundii		86.43	0.60	MNS 88		
Propionibacterium propionicum		13.57	0.36	RAF 92		
03313110	**Low discrimination**					**03313110**
Actinomyces naeslundii		67.23	0.35	MNS 88		
Propionibacterium propionicum		18.24	0.15	RAF 92		
Actinomyces israelii		14.54	0.15	MLZ 5		
03313111	**Good identification**					**03313111**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03313113	**Good identification**					**03313113**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03313120	**Low discrimination**					**03313120**
Actinomyces naeslundii		84.05	0.57	MNS 88		
Clostridium perfringens		7.55	0.17	RAF 92		
Actinomyces meyeri		3.25	0.12			
03313121	**Very good identification**					**03313121**
					SPO	
Actinomyces naeslundii		99.20	0.60	MNS 88	1	
Clostridium chauvoei		0.80	0.01	RAF 92	99	
03313122	**Low discrimination**					**03313122**
Actinomyces naeslundii		88.37	0.57	MNS 88		
Clostridium perfringens		3.74	0.11	RAF 92		
Lactobacillus fermentum		2.32	0.20			
03313123	**Very good identification**					**03313123**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03313130	**Good identification**					**03313130**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03313131	**Good identification**					**03313131**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03313132	**Low discrimination**					**03313132**
					SPO	
Actinomyces naeslundii		71.28	0.35	MNS 88	1	
Clostridium tertium		28.72	0.05	RAF 92	99	
03313133	**Good identification**					**03313133**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03313140	**Low discrimination**					**03313140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		87.28	0.49	NIT 86		
Actinomyces naeslundii		10.90	0.44	SAL 1		
Actinomyces israelii		1.83	0.22			
03313141	**Low discrimination**					**03313141**
Actinomyces naeslundii		62.76	0.47	MNS 88		
Propionibacterium propionicum		37.24	0.33	RAF 92		
03313142	**Acceptable identification**					**03313142**
Propionibacterium propionicum		90.57	0.52	NIT 86		
Actinomyces naeslundii		7.54	0.44	SAL 1		
Actinomyces israelii		1.90	0.25			
03313143	**Low discrimination**					**03313143**
Actinomyces naeslundii		52.91	0.47	MNS 88		
Propionibacterium propionicum		47.09	0.36	RAF 92		
03313160	**Low discrimination**					**03313160**
Actinomyces naeslundii		84.90	0.44	MNS 88		
Clostridium perfringens		8.23	0.04	RAF 92		
Propionibacterium propionicum		6.87	0.15			
03313161	**Good identification**					**03313161**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03313162	**Low discrimination**					**03313162**
Actinomyces naeslundii		89.18	0.44	MNS 88		
Propionibacterium propionicum		10.82	0.18	RAF 92		
03313163	**Good identification**					**03313163**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03313512	**Good identification**					**03313512**
Bifidobacterium adolescentis		97.22	0.36	MNS 87		
Actinomyces israelii		2.78	0.16	RAF 92		
03314000	**Low discrimination**					**03314000**
					GRA	
Actinomyces meyeri		47.95	0.53	SAL 1	99	
Actinomyces odontolyticus		44.65	0.56		99	
Fusobacterium mortiferum		3.84	0.43		1	
03314002	**Acceptable identification**					**03314002**
					GRA	
Actinomyces odontolyticus		92.05	0.52	NIT 99	99	
Actinomyces meyeri		4.99	0.27		99	
Fusobacterium mortiferum		1.44	0.27		1	
03314020	**Low discrimination**					**03314020**
Actinomyces odontolyticus		52.46	0.42	NIT 99		
Actinomyces meyeri		47.19	0.38	NAG 14		
Lactobacillus fermentum		0.36	0.12			
03314022	**Acceptable identification**					**03314022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		94.64	0.39	NIT 99		
Actinomyces meyeri		4.30	0.12	NAG 14		
Lactobacillus fermentum		1.06	0.12			
03314100	**Low discrimination**					**03314100**
					GRA	
Actinomyces meyeri		89.16	0.63	SAL 1	99	
Actinomyces odontolyticus		6.56	0.47		99	
Fusobacterium mortiferum		4.17	0.49		1	
03314102	**Low discrimination**					**03314102**
					GRA	
Actinomyces odontolyticus		54.65	0.43	NIT 99	99	
Actinomyces meyeri		37.51	0.37		99	
Fusobacterium mortiferum		6.30	0.33		1	
03314120	**Acceptable identification**					**03314120**
					GRA	
Actinomyces meyeri		91.38	0.48	SAL 1	99	
Actinomyces odontolyticus		8.02	0.34	NAG 12	99	
Fusobacterium mortiferum		0.32	0.15		1	
03314200	**Very good identification**					**03314200**
Actinomyces odontolyticus		99.44	0.61	NIT 99		
Actinomyces meyeri		0.56	0.18			
03314202	**Very good identification**					**03314202**
Actinomyces odontolyticus		100.0	0.57	NIT 99		
03314220	**Good identification**					**03314220**
Actinomyces odontolyticus		99.53	0.47	NIT 99		
Actinomyces meyeri		0.47	0.04	NAG 14		
03314222	**Good identification**					**03314222**
Actinomyces odontolyticus		100.0	0.43	NIT 99		
03314300	**Low discrimination**					**03314300**
Actinomyces odontolyticus		80.63	0.52	NIT 99		
Eubacterium contortum		13.39	0.32			
Actinomyces meyeri		5.71	0.29			
03314302	**Good identification**					**03314302**
Actinomyces odontolyticus		95.11	0.48	NIT 99		
Eubacterium contortum		4.55	0.19			
Actinomyces meyeri		0.34	0.03			
03314320	**Low discrimination**					**03314320**
Actinomyces odontolyticus		80.69	0.39	NIT 99		
Eubacterium contortum		14.53	0.19	NAG 14		
Actinomyces meyeri		4.78	0.14			
03314400	**Good identification**					**03314400**
Actinomyces meyeri		95.23	0.42	SAL 1		
Actinomyces odontolyticus		3.58	0.22			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Lactobacillus fermentum	1.19	0.20			
03314500	**Good identification**					**03314500**
	Actinomyces meyeri	99.01	0.53	SAL 1		
	Lactobacillus fermentum	0.50	0.24			
	Actinomyces odontolyticus	0.29	0.13			
03314520	**Good identification**					**03314520**
	Actinomyces meyeri	98.77	0.38	SAL 1		
	Lactobacillus fermentum	1.23	0.16	NAG 12		
03315000	**Low discrimination**					**03315000**
	Actinomyces meyeri	58.60	0.56	SAL 1		
	Actinomyces odontolyticus	24.25	0.53			
	Lactobacillus fermentum	12.08	0.55			
03315002	**Low discrimination**					**03315002**
	Actinomyces odontolyticus	50.35	0.49	NIT 99		
	Lactobacillus fermentum	40.90	0.55			
	Actinomyces meyeri	6.14	0.30			
03315020	**Low discrimination**					**03315020**
	Actinomyces meyeri	49.74	0.41	SAL 1		
	Lactobacillus fermentum	25.06	0.46	NAG 12		
	Actinomyces odontolyticus	24.58	0.39			
03315022	**Low discrimination**					**03315022**
	Lactobacillus fermentum	60.15	0.46	SAL 1		
	Actinomyces odontolyticus	36.16	0.36			
	Actinomyces meyeri	3.69	0.15			
03315100	**Low discrimination**					**03315100**
					GRA	
	Actinomyces meyeri	84.43	0.66	SAL 1	99	
	Lactobacillus fermentum	7.10	0.58		99	
	Fusobacterium mortiferum	3.95	0.52		1	
03315102	**Low discrimination**					**03315102**
	Lactobacillus fermentum	53.41	0.58	SAL 1		
	Actinomyces meyeri	19.65	0.40			
	Actinomyces odontolyticus	12.73	0.40			
03315112	**Low discrimination**					**03315112**
	Bifidobacterium adolescentis	84.58	0.38	MNS 87		
	Lactobacillus fermentum	6.55	0.24	RAF 92		
	Actinomyces israelii	4.90	0.23	ARA 87		
03315120	**Low discrimination**					**03315120**
	Actinomyces meyeri	77.67	0.51	SAL 1		
	Lactobacillus fermentum	15.96	0.50	NAG 12		
	Actinomyces odontolyticus	3.03	0.31			
03315122	**Low discrimination**					**03315122**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Lactobacillus fermentum	70.75	0.50	SAL 1		
	Actinomyces meyeri	10.65	0.25			
	Actinomyces odontolyticus	8.23	0.27			
03315200	**Good identification**					**03315200**
	Actinomyces odontolyticus	91.34	0.58	NIT 99		
	Eubacterium contortum	7.19	0.32			
	Actinomyces meyeri	1.15	0.21			
03315202	**Good identification**					**03315202**
	Actinomyces odontolyticus	97.38	0.54	NIT 99		
	Eubacterium contortum	2.21	0.19			
	Lactobacillus fermentum	0.41	0.20			
03315220	**Acceptable identification**					**03315220**
	Actinomyces odontolyticus	90.82	0.44	NIT 99		
	Eubacterium contortum	7.75	0.19	NAG 14		
	Actinomyces meyeri	0.96	0.07			
03315222	**Good identification**					**03315222**
	Actinomyces odontolyticus	96.79	0.40	NIT 99		
	Eubacterium contortum	2.38	0.06	NAG 14		
	Lactobacillus fermentum	0.84	0.12			
03315300	**Very good identification**					**03315300**
	Eubacterium contortum	95.42	0.66	ESL 99		
	Actinomyces odontolyticus	3.87	0.49			
	Actinomyces meyeri	0.62	0.32			
03315302	**Low discrimination**					**03315302**
	Eubacterium contortum	87.25	0.53	ESL 99		
	Actinomyces odontolyticus	12.29	0.45			
	Lactobacillus fermentum	0.27	0.24			
03315320	**Good identification**					**03315320**
	Eubacterium contortum	95.50	0.53	ESL 99		
	Actinomyces odontolyticus	3.57	0.36			
	Actinomyces meyeri	0.48	0.17			
03315322	**Low discrimination**					**03315322**
					GRA	
	Eubacterium contortum	86.33	0.40	ESL 99	99	
	Actinomyces odontolyticus	11.21	0.32		99	
	Parabacteroides distasonis	1.18	0.19		1	
03315400	**Low discrimination**					**03315400**
	Actinomyces meyeri	53.95	0.45	SAL 1		
	Lactobacillus fermentum	44.47	0.55			
	Actinomyces odontolyticus	0.90	0.19			
03315402	**Acceptable identification**					**03315402**
	Lactobacillus fermentum	92.71	0.55	SAL 1		
	Actinomyces meyeri	3.48	0.19			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	1.57	0.25			
03315420	**Low discrimination**					**03315420**
	Lactobacillus fermentum	66.83	0.46	SAL 1		
	Actinomyces meyeri	33.17	0.30			
03315422	**Good identification**					**03315422**
	Lactobacillus fermentum	98.49	0.46	SAL 1		
	Actinomyces meyeri	1.51	0.05			
03315500	**Low discrimination**					**03315500**
	Actinomyces meyeri	71.20	0.56	SAL 1		
	Lactobacillus fermentum	23.94	0.58			
	Eubacterium contortum	3.68	0.30			
03315502	**Low discrimination**					**03315502**
	Lactobacillus fermentum	53.12	0.58	SAL 1		
	Bifidobacterium adolescentis	37.97	0.50			
	Actinomyces meyeri	4.89	0.30			
03315512	**Good identification**					**03315512**
	Bifidobacterium adolescentis	97.26	0.52	MNS 87		
	Lactobacillus fermentum	1.13	0.24	RAF 92		
	Bifidobacterium longum subsp. longum	0.92	0.16			
03315520	**Low discrimination**					**03315520**
	Actinomyces meyeri	50.48	0.41	SAL 1		
	Lactobacillus fermentum	41.49	0.50	NAG 12		
	Eubacterium contortum	3.38	0.17			
03315522	**Low discrimination**					**03315522**
	Lactobacillus fermentum	87.72	0.50	SAL 1		
	Bifidobacterium longum subsp. longum	5.13	0.22			
	Actinomyces meyeri	3.30	0.15			
03315552	**Good identification**					**03315552**
	Bifidobacterium adolescentis	97.82	0.36	MNS 87		
	Actinomyces israelii	2.18	0.14	RAF 92		
03315700	**Very good identification**					**03315700**
	Eubacterium contortum	99.67	0.64	ESL 99		
	Actinomyces meyeri	0.20	0.21			
	Lactobacillus fermentum	0.07	0.24			
03315702	**Good identification**					**03315702**
	Eubacterium contortum	99.07	0.52	ESL 99		
	Lactobacillus fermentum	0.37	0.24			
	Bifidobacterium adolescentis	0.27	0.16			
03315720	**Acceptable identification**					**03315720**
					GRA	
	Eubacterium contortum	94.08	0.52	ESL 99	99	
	Bacteroides vulgatus	5.66	0.23		1	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		0.14	0.07		99	
03315722	**Good identification**					**03315722**
					GRA	
Eubacterium contortum		98.30	0.39	ESL 99	99	
Parabacteroides distasonis		1.00	0.16		1	
Lactobacillus fermentum		0.70	0.16		99	
03317000	**Low discrimination**					**03317000**
Actinomyces israelii		48.36	0.37	CEL 93		
Actinomyces meyeri		30.37	0.21			
Actinomyces odontolyticus		12.57	0.19			
03317002	**Low discrimination**					**03317002**
Actinomyces israelii		83.86	0.40	CEL 93		
Actinomyces odontolyticus		8.91	0.15			
Lactobacillus fermentum		7.24	0.20			
03317100	**Low discrimination**					**03317100**
Actinomyces israelii		56.87	0.47	CEL 93		
Actinomyces meyeri		35.71	0.32			
Lactobacillus fermentum		3.00	0.24			
03317102	**Low discrimination**					**03317102**
Actinomyces israelii		72.20	0.50	CEL 93		
Bifidobacterium adolescentis		21.98	0.35			
Lactobacillus fermentum		2.54	0.24			
03317112	**Low discrimination**					**03317112**
					SPO	
Bifidobacterium adolescentis		70.35	0.36	MNS 87	1	
Actinomyces israelii		28.25	0.36	RAF 92	1	
Clostridium tertium		1.41	-.01	ARA 87	99	
03317140	**Good identification**					**03317140**
Actinomyces israelii		95.98	0.40	CEL 93		
Propionibacterium propionicum		4.02	0.15			
03317142	**Low discrimination**					**03317142**
Actinomyces israelii		87.82	0.43	CEL 93		
Bifidobacterium adolescentis		8.50	0.19			
Propionibacterium propionicum		3.68	0.18			
03317400	**Low discrimination**					**03317400**
Actinomyces israelii		74.07	0.35	CEL 93		
Actinomyces meyeri		14.22	0.11			
Lactobacillus fermentum		11.72	0.20			
03317402	**Low discrimination**					**03317402**
Actinomyces israelii		82.91	0.38	CEL 93		
Lactobacillus fermentum		8.74	0.20			
Bifidobacterium adolescentis		8.34	0.15			
03317500	**Low discrimination**					**03317500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		77.27	0.45	CEL 93			
Actinomyces meyeri		14.83	0.21				
Lactobacillus fermentum		4.99	0.24				
03317502	**Low discrimination**						**03317502**
Bifidobacterium adolescentis		70.48	0.49	MNS 87			
Actinomyces israelii		28.30	0.48	RAF 92			
Lactobacillus fermentum		1.22	0.24				
03317512	**Good identification**						**03317512**
Bifidobacterium adolescentis		95.32	0.50	MNS 87			
Actinomyces israelii		4.68	0.34	RAF 92			
03317540	**Good identification**						**03317540**
Actinomyces israelii		100.0	0.38	CEL 93			
03317542	**Low discrimination**						**03317542**
Actinomyces israelii		55.82	0.41	CEL 93			
Bifidobacterium adolescentis		44.18	0.33				
03321102	**Low discrimination**						**03321102**
					COC	GRA	
Actinomyces naeslundii		69.62	0.35	MNS 88	1	99	
Atopobium parvulum		15.37	0.01	RAF 92	99	99	
Fusobacterium mortiferum		15.00	0.12	FRU 95	1	1	
03321103	**Low discrimination**						**03321103**
					SPO		
Actinomyces naeslundii		72.27	0.38	MNS 88	1		
Clostridium chauvoei		27.73	0.08	RAF 92	99		
03321121	**Low discrimination**						**03321121**
					SPO		
Actinomyces naeslundii		57.98	0.38	MNS 88	1		
Clostridium chauvoei		42.02	0.13	RAF 92	99		
03321123	**Low discrimination**						**03321123**
					SPO		
Actinomyces naeslundii		88.66	0.38	MNS 88	1		
Clostridium chauvoei		11.34	0.00	RAF 92	99		
03323001	**Good identification**						**03323001**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03323003	**Good identification**						**03323003**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03323021	**Good identification**						**03323021**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03323023	**Good identification**						**03323023**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03323100	**Low discrimination**						**03323100**
Actinomyces naeslundii		89.74	0.40	MNS 88			
Propionibacterium propionicum		10.26	0.14	RAF 92			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
03323101	**Good identification**						**03323101**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03323102	**Low discrimination**						**03323102**
Actinomyces naeslundii		85.36	0.40	MNS 88			
Propionibacterium propionicum		14.64	0.17	RAF 92			
03323103	**Good identification**						**03323103**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03323120	**Good identification**						**03323120**
Actinomyces naeslundii		100.0	0.40	MNS 88			
03323121	**Good identification**						**03323121**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03323122	**Good identification**						**03323122**
Actinomyces naeslundii		100.0	0.40	MNS 88			
03323123	**Good identification**						**03323123**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03330000	**Low discrimination**						**03330000**
					COC	GRA	
Fusobacterium mortiferum		75.31	0.53	MLT 11	1	1	
Megasphaera elsdenii		13.51	0.32		99	1	
Actinomyces odontolyticus		4.52	0.27		1	99	
03330002	**Low discrimination**						**03330002**
					COC	GRA	
Fusobacterium mortiferum		59.19	0.37	MLT 11	1	1	
Actinomyces odontolyticus		19.60	0.23	BGL 10	1	99	
Megasphaera elsdenii		10.62	0.15		99	1	
03330100	**Good identification**						**03330100**
					COC	GRA	
Fusobacterium mortiferum		94.89	0.60	MLT 11	1	1	
Actinomyces meyeri		2.66	0.24		1	99	
Atopobium parvulum		0.93	0.14		99	99	
03330102	**Low discrimination**						**03330102**
					GRA		
Fusobacterium mortiferum		87.76	0.43	MLT 11	1		
Actinomyces naeslundii		4.89	0.33	BGL 10	99		
Actinomyces odontolyticus		3.93	0.15		99		
03330103	**Good identification**						**03330103**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03330121	**Good identification**						**03330121**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03330123	**Good identification**						**03330123**
Actinomyces naeslundii		100.0	0.36	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03331000	**Low discrimination**					**03331000**
					GRA	
Fusobacterium mortiferum		52.24	0.56	MLT 11	1	
Lactobacillus jensenii		37.86	0.44		99	
Actinomyces naeslundii		4.24	0.49		99	
03331001	**Low discrimination**					**03331001**
					GRA	
Actinomyces naeslundii		87.48	0.52	MNS 88	99	
Fusobacterium mortiferum		7.26	0.22	RAF 92	1	
Lactobacillus jensenii		5.26	0.10		99	
03331002	**Low discrimination**					**03331002**
					GRA	
Lactobacillus jensenii		65.25	0.41	MNS 99	99	
Fusobacterium mortiferum		15.01	0.40	CEL 95	1	
Actinomyces naeslundii		10.96	0.49		99	
03331003	**Good identification**					**03331003**
Actinomyces naeslundii		96.15	0.52	MNS 88		
Lactobacillus jensenii		3.86	0.07	RAF 92		
03331010	**Acceptable identification**					**03331010**
					GRA	
Lactobacillus jensenii		93.39	0.36	MNS 99	99	
Fusobacterium mortiferum		3.91	0.22	CEL 95	1	
Actinomyces naeslundii		1.65	0.27		99	
03331020	**Low discrimination**					**03331020**
Clostridium perfringens		64.17	0.41	MNS 99		
Lactobacillus jensenii		22.35	0.34	LAC 88		
Actinomyces naeslundii		10.01	0.49			
03331021	**Good identification**					**03331021**
Actinomyces naeslundii		95.86	0.52	MNS 88		
Clostridium perfringens		4.14	0.07	RAF 92		
03331022	**Low discrimination**					**03331022**
Clostridium perfringens		51.97	0.35	MNS 99		
Lactobacillus jensenii		25.65	0.31	LAC 88		
Actinomyces naeslundii		17.23	0.49			
03331023	**Good identification**					**03331023**
Actinomyces naeslundii		98.01	0.52	MNS 88		
Clostridium perfringens		1.99	0.01	RAF 92		
03331040	**Low discrimination**					**03331040**
					GRA	
Actinomyces naeslundii		37.74	0.36	MNS 88	99	
Fusobacterium mortiferum		26.62	0.22	RAF 92	1	
Lactobacillus jensenii		19.29	0.10		99	
03331041	**Good identification**					**03331041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03331042	**Low discrimination**					**03331042**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		66.89	0.36	MNS 88			
Lactobacillus jensenii		22.80	0.07	RAF 92			
Lactobacillus acidophilus		10.32	-.02				
03331043	**Good identification**						**03331043**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03331061	**Good identification**						**03331061**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03331063	**Good identification**						**03331063**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03331100	**Low discrimination**						**03331100**
					COC	GRA	
Fusobacterium mortiferum		54.24	0.63	MLT 11	1	1	
Atopobium parvulum		35.00	0.48		99	99	
Actinomyces naeslundii		5.37	0.57		1	99	
03331101	**Low discrimination**						**03331101**
					GRA	SPO	
Actinomyces naeslundii		72.52	0.60	MNS 88	99	1	
Clostridium chauvoei		19.36	0.27	RAF 92	99	99	
Fusobacterium mortiferum		4.93	0.28		1	1	
03331102	**Low discrimination**						**03331102**
					COC	GRA	
Atopobium parvulum		24.65	0.35	MNS 99	99	99	
Fusobacterium mortiferum		24.05	0.46		1	1	
Actinomyces naeslundii		21.42	0.57		1	99	
03331103	**Good identification**						**03331103**
					COC	SPO	
Actinomyces naeslundii		94.12	0.60	MNS 88	1	1	
Clostridium chauvoei		4.44	0.14	RAF 92	1	99	
Atopobium parvulum		0.73	0.01		99	1	
03331111	**Good identification**						**03331111**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03331113	**Good identification**						**03331113**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03331120	**Low discrimination**						**03331120**
					COC		
Clostridium perfringens		86.57	0.56	MNS 99	1		
Atopobium parvulum		6.03	0.35		99		
Actinomyces naeslundii		5.24	0.57		1		
03331121	**Low discrimination**						**03331121**
					SPO		
Actinomyces naeslundii		82.79	0.60	MNS 88	1		
Clostridium perfringens		9.21	0.21	RAF 92	84		
Clostridium chauvoei		7.37	0.19		99		
03331122	**Low discrimination**						**03331122**
Clostridium perfringens		77.95	0.50	MNS 99			
Actinomyces naeslundii		10.03	0.57				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		8.93	0.32			
03331123	**Good identification**					**03331123**
					SPO	
Actinomyces naeslundii		93.63	0.60	MNS 88	1	
Clostridium perfringens		4.90	0.16	RAF 92	84	
Clostridium chauvoei		1.47	0.06		99	
03331131	**Good identification**					**03331131**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03331133	**Good identification**					**03331133**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03331140	**Low discrimination**					**03331140**
					GRA	
Actinomyces naeslundii		37.31	0.44	MNS 88	99	
Fusobacterium mortiferum		21.58	0.28	RAF 92	1	
Clostridium perfringens		20.57	0.17		99	
03331141	**Good identification**					**03331141**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03331142	**Low discrimination**					**03331142**
Actinomyces naeslundii		43.87	0.44	MNS 88		
Lactobacillus acidophilus		27.35	0.16	RAF 92		
Clostridium perfringens		11.38	0.12			
03331143	**Good identification**					**03331143**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03331160	**Acceptable identification**					**03331160**
					COC	
Clostridium perfringens		94.36	0.43	MNS 99	1	
Actinomyces naeslundii		5.29	0.44		1	
Atopobium parvulum		0.35	0.01		99	
03331161	**Low discrimination**					**03331161**
Actinomyces naeslundii		89.28	0.47	MNS 88		
Clostridium perfringens		10.72	0.09	RAF 92		
03331162	**Low discrimination**					**03331162**
Clostridium perfringens		83.79	0.38	MNS 99		
Actinomyces naeslundii		9.99	0.44			
Lactobacillus acidophilus		6.23	0.16			
03331163	**Acceptable identification**					**03331163**
Actinomyces naeslundii		94.65	0.47	MNS 88		
Clostridium perfringens		5.35	0.03	RAF 92		
03332100	**Low discrimination**					**03332100**
					GRA	
Propionibacterium propionicum		88.59	0.48	NIT 86	99	
Actinomyces naeslundii		6.20	0.38	SAL 1	99	
Fusobacterium mortiferum		5.21	0.25		1	
03332102	**Good identification**					**03332102**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		95.55	0.51	NIT 86		
Actinomyces naeslundii		4.45	0.38	SAL 1		
03332120	**Low discrimination**					**03332120**
Actinomyces naeslundii		87.38	0.38	MNS 88		
Propionibacterium propionicum		12.62	0.14	RAF 92		
03332121	**Good identification**					**03332121**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03332122	**Low discrimination**					**03332122**
Actinomyces naeslundii		82.19	0.38	MNS 88		
Propionibacterium propionicum		17.81	0.17	RAF 92		
03332123	**Good identification**					**03332123**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03332140	**Good identification**					**03332140**
Propionibacterium propionicum		98.78	0.48	NIT 86		
Actinomyces naeslundii		1.22	0.25	SAL 1		
03332142	**Good identification**					**03332142**
Propionibacterium propionicum		99.18	0.51	NIT 86		
Actinomyces naeslundii		0.82	0.25	SAL 1		
03333000	**Low discrimination**					**03333000**
					CAT	
Actinomyces naeslundii		75.28	0.54	MNS 88	25	
Actinomyces israelii		7.91	0.28	RAF 92	1	
Propionibacterium granulosum		7.20	0.22		90	
03333001	**Very good identification**					**03333001**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03333002	**Low discrimination**					**03333002**
Actinomyces naeslundii		80.77	0.54	MNS 88		
Actinomyces israelii		12.72	0.31	RAF 92		
Lactobacillus jensenii		2.50	0.07			
03333003	**Very good identification**					**03333003**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03333011	**Good identification**					**03333011**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03333013	**Good identification**					**03333013**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03333020	**Acceptable identification**					**03333020**
					CAT	
Actinomyces naeslundii		94.58	0.54	MNS 88	25	
Clostridium perfringens		3.16	0.07	RAF 92	1	
Propionibacterium granulosum		2.26	0.12		90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03333021	**Very good identification**					**03333021**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03333022	**Good identification**					**03333022**
Actinomyces naeslundii		98.46	0.54	MNS 88		
Clostridium perfringens		1.55	0.01	RAF 92		
03333023	**Very good identification**					**03333023**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03333031	**Good identification**					**03333031**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03333033	**Good identification**					**03333033**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03333040	**Low discrimination**					**03333040**
Actinomyces naeslundii		75.27	0.41	MNS 88		
Actinomyces israelii		17.42	0.21	RAF 92		
Propionibacterium propionicum		7.31	0.14			
03333041	**Good identification**					**03333041**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03333042	**Low discrimination**					**03333042**
Actinomyces naeslundii		66.99	0.41	MNS 88		
Actinomyces israelii		23.26	0.24	RAF 92		
Propionibacterium propionicum		9.75	0.17			
03333043	**Good identification**					**03333043**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03333060	**Good identification**					**03333060**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03333061	**Good identification**					**03333061**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03333062	**Good identification**					**03333062**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03333063	**Good identification**					**03333063**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03333100	**Low discrimination**					**03333100**
Actinomyces naeslundii		54.47	0.62	MNS 88		
Propionibacterium propionicum		32.45	0.48	RAF 92		
Actinomyces israelii		8.04	0.39			
03333101	**Very good identification**					**03333101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.77	0.65	MNS 88		
Propionibacterium propionicum		4.23	0.32	RAF 92		
03333102	**Low discrimination**					**03333102**
Actinomyces naeslundii		45.66	0.62	MNS 88		
Propionibacterium propionicum		40.81	0.51	RAF 92		
Actinomyces israelii		10.11	0.42			
03333103	**Good identification**					**03333103**
Actinomyces naeslundii		93.79	0.65	MNS 88		
Propionibacterium propionicum		6.21	0.35	RAF 92		
03333110	**Low discrimination**					**03333110**
					SPO	
Actinomyces naeslundii		57.44	0.40	MNS 88	1	
Actinomyces israelii		24.07	0.24	RAF 92	1	
Clostridium tertium		11.92	0.05	MLZ 5	99	
03333111	**Good identification**					**03333111**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03333112	**Low discrimination**					**03333112**
					SPO	
Actinomyces naeslundii		34.49	0.40	MNS 88	1	
Actinomyces israelii		21.68	0.27	RAF 92	1	
Clostridium tertium		21.47	0.13	MLZ 5	99	
03333113	**Good identification**					**03333113**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03333120	**Low discrimination**					**03333120**
					SPO	
Actinomyces naeslundii		87.48	0.62	MNS 88	1	
Clostridium perfringens		7.52	0.21	RAF 92	84	
Clostridium tertium		1.61	0.09		99	
03333121	**Very good identification**					**03333121**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03333122	**Low discrimination**					**03333122**
					SPO	
Actinomyces naeslundii		82.47	0.62	MNS 88	1	
Clostridium ramosum		5.73	0.17	RAF 92	92	
Clostridium tertium		4.56	0.17		99	
03333123	**Very good identification**					**03333123**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03333130	**Low discrimination**					**03333130**
					SPO	
Actinomyces naeslundii		61.63	0.40	MNS 88	1	
Clostridium tertium		38.37	0.13	RAF 92	99	
03333131	**Good identification**					**03333131**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03333133	**Good identification**					**03333133**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Actinomyces naeslundii	100.0	0.43	MNS 88			
03333140	**Low discrimination**						**03333140**
	Propionibacterium propionicum	71.81	0.48	NIT 86			
	Actinomyces naeslundii	21.27	0.49	SAL 1			
	Actinomyces israelii	6.92	0.32				
03333141	**Low discrimination**						**03333141**
	Actinomyces naeslundii	80.00	0.52	MNS 88			
	Propionibacterium propionicum	20.01	0.32	RAF 92			
03333142	**Low discrimination**						**03333142**
	Propionibacterium propionicum	77.29	0.51	NIT 86			
	Actinomyces naeslundii	15.26	0.49	SAL 1			
	Actinomyces israelii	7.45	0.35				
03333143	**Low discrimination**						**03333143**
	Actinomyces naeslundii	72.72	0.52	MNS 88			
	Propionibacterium propionicum	27.28	0.35	RAF 92			
03333160	**Low discrimination**						**03333160**
	Actinomyces naeslundii	88.74	0.49	MNS 88			
	Clostridium perfringens	8.23	0.09	RAF 92			
	Propionibacterium propionicum	3.03	0.14				
03333161	**Good identification**						**03333161**
	Actinomyces naeslundii	100.0	0.52	MNS 88			
03333162	**Low discrimination**						**03333162**
	Actinomyces naeslundii	88.81	0.49	MNS 88			
	Propionibacterium propionicum	4.54	0.17	RAF 92			
	Clostridium perfringens	3.88	0.03				
03333163	**Good identification**						**03333163**
	Actinomyces naeslundii	100.0	0.52	MNS 88			
03333500	**Low discrimination**						**03333500**
	Actinomyces israelii	88.23	0.37	CEL 93			
	Actinomyces naeslundii	7.38	0.28	XYL 92			
	Propionibacterium propionicum	4.40	0.14				
03333502	**Low discrimination**						**03333502**
	Actinomyces israelii	53.65	0.40	CEL 93			
	Bifidobacterium adolescentis	40.69	0.32	XYL 92			
	Actinomyces naeslundii	2.99	0.28				
03334100	**Low discrimination**						**03334100**
					COC	GRA	
	Fusobacterium mortiferum	49.56	0.40	MLT 11	1	1	
	Actinomyces meyeri	35.85	0.29	XYL 7	1	99	
	Atopobium parvulum	11.96	0.18		99	99	
03335000	**Low discrimination**						**03335000**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		38.43	0.37	MLT 11	1		
Actinomyces israelii		17.14	0.33	XYL 7	99		
Actinomyces meyeri		16.22	0.21		99		
03335002	**Low discrimination**						**03335002**
					GRA		
Actinomyces israelii		60.54	0.37	CEL 93	99		
Fusobacterium mortiferum		10.05	0.21		1		
Actinomyces odontolyticus		9.69	0.15		99		
03335100	**Low discrimination**						**03335100**
					COC	GRA	
Atopobium parvulum		86.03	0.53	MNS 99	99	99	
Fusobacterium mortiferum		5.40	0.43		1	1	
Actinomyces israelii		4.13	0.44		1	99	
03335102	**Low discrimination**						**03335102**
					COC		
Atopobium parvulum		60.98	0.40	MNS 99	99		
Actinomyces israelii		24.89	0.47		1		
Bifidobacterium adolescentis		9.30	0.33		1		
03335120	**Low discrimination**						**03335120**
					COC		
Atopobium parvulum		87.53	0.40	MNS 99	99		
Clostridium perfringens		6.83	0.21		1		
Actinomyces meyeri		3.07	0.17		1		
03335140	**Low discrimination**						**03335140**
					COC		
Actinomyces israelii		64.90	0.37	CEL 93	1		
Atopobium parvulum		35.10	0.18		99		
03335142	**Low discrimination**						**03335142**
					COC		
Actinomyces israelii		84.57	0.40	CEL 93	1		
Bifidobacterium adolescentis		10.05	0.18		1		
Atopobium parvulum		5.38	0.06		99		
03335402	**Low discrimination**						**03335402**
Actinomyces israelii		77.98	0.35	CEL 93			
Lactobacillus fermentum		12.39	0.20				
Bifidobacterium adolescentis		9.63	0.13				
03335500	**Low discrimination**						**03335500**
					COC		
Actinomyces israelii		58.18	0.42	CEL 93	1		
Actinomyces meyeri		16.82	0.21		1		
Atopobium parvulum		14.96	0.18		99		
03335502	**Low discrimination**						**03335502**
Bifidobacterium adolescentis		73.62	0.47	MNS 87			
Actinomyces israelii		24.09	0.45	RAF 92			
Lactobacillus fermentum		1.56	0.24				
03335512	**Acceptable identification**						**03335512**
Bifidobacterium adolescentis		94.76	0.49	MNS 87			
Actinomyces israelii		3.79	0.31	RAF 92			
Clostridium clostridioforme		1.11	0.13				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03335540	**Good identification**						**03335540**
Actinomyces israelii		100.0	0.35	CEL 93			
03335542	**Low discrimination**						**03335542**
Actinomyces israelii		50.73	0.38	CEL 93			
Bifidobacterium adolescentis		49.27	0.32				
03337000	**Good identification**						**03337000**
Actinomyces israelii		100.0	0.46	CEL 93			
03337002	**Good identification**						**03337002**
Actinomyces israelii		100.0	0.49	CEL 93			
03337012	**Good identification**						**03337012**
Actinomyces israelii		100.0	0.35	CEL 93			
03337040	**Good identification**						**03337040**
Actinomyces israelii		100.0	0.39	CEL 93			
03337042	**Good identification**						**03337042**
Actinomyces israelii		100.0	0.42	CEL 93			
03337100	**Very good identification**						**03337100**
					COC		
Actinomyces israelii		95.09	0.57	CEL 93	1		
Atopobium parvulum		3.53	0.18		99		
Actinomyces naeslundii		0.57	0.28		1		
03337102	**Good identification**						**03337102**
					SPO		
Actinomyces israelii		92.95	0.60	CEL 93	1		
Bifidobacterium adolescentis		5.02	0.32		1		
Clostridium tertium		0.93	0.11		99		
03337110	**Acceptable identification**						**03337110**
					SPO		
Actinomyces israelii		94.39	0.43	CEL 93	1		
Clostridium tertium		5.61	0.07	MLZ 13	99		
03337112	**Low discrimination**						**03337112**
					SPO		
Actinomyces israelii		64.09	0.46	CEL 93	1		
Bifidobacterium adolescentis		28.28	0.33	MLZ 13	1		
Clostridium tertium		7.62	0.15		99		
03337120	**Low discrimination**						**03337120**
					COC	SPO	
Actinomyces israelii		69.89	0.36	CEL 93	1	1	
Clostridium tertium		16.42	0.11	NAG 6	1	99	
Atopobium parvulum		7.17	0.06		99	1	
03337122	**Low discrimination**						**03337122**
					SPO		
Actinomyces israelii		65.28	0.39	CEL 93	1		
Clostridium tertium		30.67	0.19	NAG 6	99		
Actinomyces naeslundii		4.06	0.28		1		
03337140	**Good identification**						**03337140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		99.10	0.50	CEL 93		
Propionibacterium propionicum		0.90	0.14			
03337142	**Good identification**					**03337142**
Actinomyces israelii		97.44	0.53	CEL 93		
Bifidobacterium adolescentis		1.67	0.16			
Propionibacterium propionicum		0.89	0.17			
03337150	**Good identification**					**03337150**
Actinomyces israelii		100.0	0.36	CEL 93		
03337152	**Low discrimination**					**03337152**
Actinomyces israelii		87.70	0.39	CEL 93		
Bifidobacterium adolescentis		12.30	0.18	MLZ 13		
03337300	**Good identification**					**03337300**
Actinomyces israelii		100.0	0.37	CEL 93		
03337302	**Good identification**					**03337302**
Actinomyces israelii		100.0	0.40	CEL 93		
03337400	**Good identification**					**03337400**
Actinomyces israelii		100.0	0.45	CEL 93		
03337402	**Good identification**					**03337402**
Actinomyces israelii		98.25	0.48	CEL 93		
Bifidobacterium adolescentis		1.75	0.12			
03337440	**Good identification**					**03337440**
Actinomyces israelii		100.0	0.38	CEL 93		
03337442	**Good identification**					**03337442**
Actinomyces israelii		100.0	0.41	CEL 93		
03337500	**Good identification**					**03337500**
Actinomyces israelii		99.34	0.55	CEL 93		
Bifidobacterium adolescentis		0.66	0.12			
03337502	**Low discrimination**					**03337502**
Actinomyces israelii		69.38	0.58	CEL 93		
Bifidobacterium adolescentis		30.62	0.46			
03337510	**Acceptable identification**					**03337510**
Actinomyces israelii		94.82	0.41	CEL 93		
Bifidobacterium adolescentis		5.19	0.13	MLZ 13		
03337512	**Low discrimination**					**03337512**
Bifidobacterium adolescentis		78.31	0.47	MNS 87		
Actinomyces israelii		21.69	0.44	RAF 92		
03337522	**Acceptable identification**					**03337522**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		93.47	0.38	CEL 93		
Bifidobacterium adolescentis		6.53	0.12	NAG 6		
03337540	**Good identification**					**03337540**
Actinomyces israelii		100.0	0.48	CEL 93		
03337542	**Low discrimination**					**03337542**
Actinomyces israelii		87.70	0.51	CEL 93		
Bifidobacterium adolescentis		12.30	0.30			
03337700	**Good identification**					**03337700**
Actinomyces israelii		100.0	0.36	CEL 93		
03337702	**Acceptable identification**					**03337702**
Actinomyces israelii		94.41	0.39	CEL 93		
Bifidobacterium adolescentis		5.59	0.12	RHA 7		
03341101	**Low discrimination**					**03341101**
					SPO	
Actinomyces naeslundii		77.55	0.35	MNS 88	1	
Clostridium chauvoei		22.45	0.03	RAF 92	99	
03341103	**Good identification**					**03341103**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03341121	**Good identification**					**03341121**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03341123	**Good identification**					**03341123**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03343100	**Low discrimination**					**03343100**
Actinomyces naeslundii		87.22	0.37	MNS 88		
Actinomyces israelii		12.78	0.14	RAF 92		
03343101	**Good identification**					**03343101**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03343102	**Low discrimination**					**03343102**
Actinomyces naeslundii		70.41	0.37	MNS 88		
Actinomyces israelii		15.47	0.17	RAF 92		
Lactobacillus acidophilus		14.12	0.01	FRU 95		
03343103	**Good identification**					**03343103**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03343120	**Good identification**					**03343120**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03343121	**Good identification**					**03343121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03343122	**Low discrimination**					**03343122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		62.62	0.37	MNS 88	1		
Clostridium barati		14.12	-.01	RAF 92	99		
Lactobacillus acidophilus		12.56	0.01	FRU 95	1		
03343123	**Good identification**						**03343123**
Actinomyces naeslundii		100.0	0.40	MNS 88			
03350000	**Low discrimination**						**03350000**
					COC	GRA	
Fusobacterium mortiferum		69.95	0.54	MLT 11	1	1	
Megasphaera elsdenii		11.25	0.32		99	1	
Lactobacillus catenaformis		8.68	0.29		1	99	
03350100	**Low discrimination**						**03350100**
					GRA		
Fusobacterium mortiferum		88.05	0.60	MLT 11	1		
Lactobacillus catenaformis		7.02	0.32		99		
Actinomyces meyeri		2.21	0.24		99		
03350102	**Low discrimination**						**03350102**
					GRA	SPO	
Fusobacterium mortiferum		39.91	0.44	MLT 11	1	1	
Lactobacillus catenaformis		28.66	0.32	BGL 10	99	1	
Clostridium barati		15.01	0.22		99	99	
03351000	**Low discrimination**						**03351000**
					GRA		
Lactobacillus catenaformis		60.40	0.63	MNS 99	99		
Lactobacillus jensenii		30.32	0.58		99		
Fusobacterium mortiferum		7.37	0.57		1		
03351002	**Low discrimination**						**03351002**
Lactobacillus catenaformis		68.16	0.63	MNS 99			
Lactobacillus jensenii		22.81	0.55				
Lactobacillus acidophilus		3.94	0.39				
03351010	**Acceptable identification**						**03351010**
					GRA		
Lactobacillus jensenii		93.49	0.50	MNS 99	99		
Lactobacillus catenaformis		5.64	0.29		99		
Fusobacterium mortiferum		0.69	0.23		1		
03351012	**Acceptable identification**						**03351012**
Lactobacillus jensenii		90.46	0.47	MNS 99			
Lactobacillus catenaformis		8.19	0.29				
Lactobacillus acidophilus		0.47	0.05				
03351020	**Low discrimination**						**03351020**
					SPO		
Lactobacillus catenaformis		57.25	0.53	MNS 99	1		
Lactobacillus jensenii		28.74	0.47		1		
Clostridium barati		11.51	0.35		99		
03351021	**Low discrimination**						**03351021**
Actinomyces naeslundii		67.89	0.49	MNS 88			
Lactobacillus catenaformis		18.86	0.18	RAF 92			
Lactobacillus jensenii		9.47	0.13				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03351022	**Low discrimination**					**03351022**
					SPO	
Lactobacillus catenaformis		45.25	0.53	MNS 99	1	
Clostridium barati		27.30	0.44		99	
Lactobacillus jensenii		15.15	0.44		1	
03351023	**Low discrimination**					**03351023**
					SPO	
Actinomyces naeslundii		62.40	0.49	MNS 88	1	
Lactobacillus catenaformis		17.33	0.18	RAF 92	1	
Clostridium barati		10.46	0.09		99	
03351030	**Acceptable identification**					**03351030**
					SPO	
Lactobacillus jensenii		92.58	0.39	MNS 99	1	
Lactobacillus catenaformis		5.59	0.18		1	
Clostridium barati		1.12	0.01		99	
03351032	**Low discrimination**					**03351032**
					SPO	
Lactobacillus jensenii		84.92	0.36	MNS 99	1	
Lactobacillus catenaformis		7.69	0.18		1	
Clostridium barati		4.64	0.09		99	
03351041	**Good identification**					**03351041**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03351043	**Good identification**					**03351043**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03351061	**Good identification**					**03351061**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03351063	**Good identification**					**03351063**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03351100	**Low discrimination**					**03351100**
					GRA	
Lactobacillus catenaformis		75.23	0.66	MNS 99	99	
Fusobacterium mortiferum		14.29	0.63		1	
Collinsella aerofaciens		5.53	0.32		99	
03351101	**Low discrimination**					**03351101**
					GRA	
Actinomyces naeslundii		55.71	0.57	MNS 88	99	
Lactobacillus catenaformis		32.62	0.32	RAF 92	99	
Fusobacterium mortiferum		6.20	0.29		1	
03351102	**Low discrimination**					**03351102**
					SPO	
Lactobacillus catenaformis		62.59	0.66	MNS 99	1	
Lactobacillus acidophilus		27.74	0.57		1	
Clostridium barati		6.29	0.44		99	
03351103	**Low discrimination**					**03351103**
Actinomyces naeslundii		52.19	0.57	MNS 88		
Lactobacillus catenaformis		30.56	0.32	RAF 92		
Lactobacillus acidophilus		13.54	0.23			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03351111	**Good identification**					**03351111**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03351113	**Good identification**					**03351113**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03351120	**Low discrimination**					**03351120**
					SPO	
Lactobacillus catenaformis		57.28	0.56	MNS 99	1	
Clostridium barati		23.04	0.44		99	
Clostridium paraputrificum		9.43	0.32		99	
03351121	**Low discrimination**					**03351121**
					SPO	
Actinomyces naeslundii		80.57	0.57	MNS 88	1	
Lactobacillus catenaformis		11.79	0.21	RAF 92	1	
Clostridium barati		4.74	0.09		99	
03351122	**Low discrimination**					**03351122**
					SPO	
Lactobacillus acidophilus		41.82	0.57	MNS 99	1	
Clostridium barati		28.46	0.52		99	
Lactobacillus catenaformis		23.59	0.56		1	
03351123	**Low discrimination**					**03351123**
					SPO	
Actinomyces naeslundii		62.17	0.57	MNS 88	1	
Lactobacillus acidophilus		16.13	0.23	RAF 92	1	
Clostridium barati		10.98	0.18		99	
03351131	**Good identification**					**03351131**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03351133	**Good identification**					**03351133**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03351141	**Good identification**					**03351141**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03351142	**Low discrimination**					**03351142**
					SPO	
Lactobacillus acidophilus		70.28	0.41	MNS 99	1	
Lactobacillus catenaformis		12.96	0.32	SOR 11	1	
Clostridium barati		11.22	0.25		99	
03351143	**Low discrimination**					**03351143**
Actinomyces naeslundii		84.62	0.44	MNS 88		
Lactobacillus acidophilus		15.38	0.07	RAF 92		
03351161	**Good identification**					**03351161**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03351162	**Low discrimination**					**03351162**
					SPO	
Lactobacillus acidophilus		62.30	0.41	MNS 99	1	
Clostridium barati		29.83	0.34	SOR 11	99	
Lactobacillus catenaformis		2.87	0.21		1	

Code Taxon	ID score	T index	tests against	Differentiation tests	Code

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03351163	**Low discrimination**						**03351163**
					SPO		
Actinomyces naeslundii		78.82	0.44	MNS 88	1		
Lactobacillus acidophilus		14.32	0.07	RAF 92	1		
Clostridium barati		6.86	-.01		99		
03351302	**Low discrimination**						**03351302**
					SPO		
Lactobacillus acidophilus		61.67	0.38	MNS 99	1		
Lactobacillus catenaformis		18.67	0.32	RHA 7	1		
Clostridium barati		16.16	0.25		99		
03351322	**Low discrimination**						**03351322**
					GRA	SPO	
Lactobacillus acidophilus		43.22	0.38	MNS 99	99	1	
Clostridium barati		33.98	0.34	RHA 7	99	99	
Prevotella oralis		14.95	0.29		1	1	
03351502	**Low discrimination**						**03351502**
Bifidobacterium adolescentis		45.66	0.42	MNS 87			
Lactobacillus catenaformis		26.36	0.32	RAF 92			
Lactobacillus acidophilus		11.68	0.23	XYL 87			
03351512	**Good identification**						**03351512**
Bifidobacterium adolescentis		97.49	0.43	MNS 87			
Actinomyces israelii		2.51	0.22	RAF 92			
03352101	**Good identification**						**03352101**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03352102	**Low discrimination**						**03352102**
					SPO		
Actinomyces naeslundii		41.44	0.36	MNS 88	1		
Clostridium barati		20.50	0.04	RAF 92	99		
Propionibacterium propionicum		16.07	0.18	GAL 96	1		
03352103	**Good identification**						**03352103**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03352120	**Low discrimination**						**03352120**
					SPO		
Actinomyces naeslundii		66.91	0.36	MNS 88	1		
Clostridium barati		33.09	0.04	RAF 92	99		
03352121	**Good identification**						**03352121**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03352123	**Good identification**						**03352123**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03353001	**Good identification**						**03353001**
Actinomyces naeslundii		100.0	0.54	MNS 88			
03353003	**Good identification**						**03353003**
Actinomyces naeslundii		100.0	0.54	MNS 88			
03353020	**Low discrimination**						**03353020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		57.85	0.51	MNS 88	1	
Clostridium barati		21.49	0.17	RAF 92	99	
Lactobacillus catenaformis		12.42	0.18		1	
03353021	**Good identification**					**03353021**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03353023	**Good identification**					**03353023**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03353041	**Good identification**					**03353041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03353043	**Good identification**					**03353043**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03353060	**Low discrimination**					**03353060**
					SPO	
Actinomyces naeslundii		84.53	0.38	MNS 88	1	
Clostridium barati		15.47	-.01	RAF 92	99	
03353061	**Good identification**					**03353061**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03353062	**Low discrimination**					**03353062**
					SPO	
Actinomyces naeslundii		54.20	0.38	MNS 88	1	
Clostridium barati		29.77	0.07	RAF 92	99	
Lactobacillus acidophilus		9.82	0.01		1	
03353063	**Good identification**					**03353063**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03353100	**Low discrimination**					**03353100**
Actinomyces naeslundii		38.67	0.59	MNS 88		
Actinomyces israelii		29.52	0.48	RAF 92		
Lactobacillus catenaformis		17.50	0.32			
03353101	**Very good identification**					**03353101**
Actinomyces naeslundii		99.49	0.62	MNS 88		
Actinomyces israelii		0.51	0.14	RAF 92		
03353102	**Low discrimination**					**03353102**
Actinomyces israelii		25.72	0.51	XYL 92		
Lactobacillus acidophilus		23.47	0.35			
Actinomyces naeslundii		22.46	0.59			
03353103	**Very good identification**					**03353103**
Actinomyces naeslundii		98.55	0.62	MNS 88		
Actinomyces israelii		0.76	0.17	RAF 92		
Lactobacillus acidophilus		0.69	0.01			
03353111	**Good identification**					**03353111**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MNS 88		
03353113	**Good identification**					**03353113**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03353120	**Low discrimination**					**03353120**
					SPO	
Actinomyces naeslundii		49.01	0.59	MNS 88	1	
Clostridium barati		19.19	0.25	RAF 92	99	
Clostridium tertium		14.55	0.27		99	
03353121	**Very good identification**					**03353121**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03353123	**Very good identification**					**03353123**
					SPO	
Actinomyces naeslundii		97.95	0.62	MNS 88	1	
Clostridium barati		0.78	-.01	RAF 92	99	
Lactobacillus acidophilus		0.69	0.01		1	
03353131	**Good identification**					**03353131**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03353132	**Acceptable identification**					**03353132**
					SPO	
Clostridium tertium		94.32	0.39	MNS 99	99	
Actinomyces naeslundii		3.14	0.37	TRE 90	1	
Clostridium barati		0.71	-.01		99	
03353133	**Low discrimination**					**03353133**
					SPO	
Actinomyces naeslundii		83.15	0.40	MNS 88	1	
Clostridium tertium		16.85	0.05	RAF 92	99	
03353140	**Low discrimination**					**03353140**
					SPO	
Actinomyces israelii		59.92	0.41	XYL 92	1	
Actinomyces naeslundii		35.62	0.46		1	
Clostridium barati		2.29	-.01		99	
03353141	**Good identification**					**03353141**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03353142	**Low discrimination**					**03353142**
Actinomyces israelii		54.78	0.44	XYL 92		
Actinomyces naeslundii		21.71	0.46			
Lactobacillus acidophilus		15.89	0.19			
03353143	**Good identification**					**03353143**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03353160	**Low discrimination**					**03353160**
					SPO	
Actinomyces naeslundii		76.90	0.46	MNS 88	1	
Clostridium barati		14.84	0.07	RAF 92	99	
Actinomyces israelii		8.26	0.20		1	
03353161	**Good identification**					**03353161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	MNS 88		
03353162	**Low discrimination**					**03353162**
					SPO	
Actinomyces naeslundii		38.83	0.46	MNS 88	1	
Lactobacillus acidophilus		28.41	0.19	RAF 92	1	
Clostridium barati		22.48	0.15		99	
03353163	**Good identification**					**03353163**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03353400	**Low discrimination**					**03353400**
					SPO	
Actinomyces israelii		57.95	0.36	XYL 92	1	
Clostridium innocuum		42.05	0.19		99	
03353402	**Low discrimination**					**03353402**
					SPO	
Actinomyces israelii		67.40	0.39	XYL 92	1	
Clostridium innocuum		32.61	0.19		99	
03353500	**Good identification**					**03353500**
Actinomyces israelii		98.41	0.46	XYL 92		
Actinomyces naeslundii		1.59	0.25			
03353502	**Low discrimination**					**03353502**
Actinomyces israelii		58.60	0.49	XYL 92		
Bifidobacterium adolescentis		40.11	0.40			
Lactobacillus acidophilus		0.66	0.01			
03353512	**Low discrimination**					**03353512**
Bifidobacterium adolescentis		84.84	0.42	MNS 87		
Actinomyces israelii		15.16	0.35	RAF 92		
03353540	**Good identification**					**03353540**
Actinomyces israelii		100.0	0.39	XYL 92		
03353542	**Low discrimination**					**03353542**
Actinomyces israelii		82.13	0.42	XYL 92		
Bifidobacterium adolescentis		17.87	0.25			
03354100	**Low discrimination**					**03354100**
					GRA	
Fusobacterium mortiferum		57.72	0.41	MLT 11	1	
Actinomyces meyeri		37.41	0.29	XYL 7	99	
Actinomyces odontolyticus		2.75	0.13		99	
03355000	**Low discrimination**					**03355000**
					GRA	
Actinomyces israelii		30.19	0.43		99	
Lactobacillus catenaformis		23.51	0.29		99	
Fusobacterium mortiferum		21.39	0.38		1	
03355002	**Low discrimination**					**03355002**
Actinomyces israelii		52.71	0.46			
Lactobacillus catenaformis		27.37	0.29			
Lactobacillus jensenii		9.16	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03355040	**Good identification**					**03355040**
Actinomyces israelii		100.0	0.36			
03355042	**Good identification**					**03355042**
Actinomyces israelii		100.0	0.39			
03355100	**Low discrimination**					**03355100**
					GRA	
Actinomyces israelii		45.78	0.53		99	
Fusobacterium mortiferum		18.92	0.44		1	
Lactobacillus catenaformis		13.37	0.32		99	
03355102	**Low discrimination**					**03355102**
Actinomyces israelii		58.40	0.56			
Bifidobacterium adolescentis		19.70	0.42			
Lactobacillus catenaformis		11.37	0.32			
03355110	**Good identification**					**03355110**
Actinomyces israelii		95.99	0.39	MLZ 13		
Bifidobacterium adolescentis		4.01	0.09			
03355112	**Low discrimination**					**03355112**
					SPO	
Bifidobacterium adolescentis		72.98	0.43	MNS 87	1	
Actinomyces israelii		26.46	0.42	RAF 92	1	
Clostridium tertium		0.56	-.01	ARA 87	99	
03355140	**Good identification**					**03355140**
Actinomyces israelii		100.0	0.46			
03355142	**Low discrimination**					**03355142**
Actinomyces israelii		88.14	0.49			
Bifidobacterium adolescentis		9.45	0.26			
Lactobacillus acidophilus		2.42	0.07			
03355152	**Low discrimination**					**03355152**
Actinomyces israelii		53.28	0.35	MLZ 13		
Bifidobacterium adolescentis		46.72	0.28			
03355302	**Low discrimination**					**03355302**
Actinomyces israelii		53.23	0.37	RHA 7		
Eubacterium contortum		34.86	0.19			
Actinomyces odontolyticus		4.91	0.11			
03355400	**Low discrimination**					**03355400**
Actinomyces israelii		87.01	0.41			
Actinomyces meyeri		7.12	0.11			
Lactobacillus fermentum		5.87	0.20			
03355402	**Low discrimination**					**03355402**
Actinomyces israelii		86.47	0.44			
Bifidobacterium adolescentis		9.64	0.22			
Lactobacillus fermentum		3.89	0.20			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03355442	**Good identification**					**03355442**
Actinomyces israelii		100.0	0.37			
03355500	**Low discrimination**					**03355500**
					SPO	
Actinomyces israelii		83.08	0.52		1	
Actinomyces meyeri		6.80	0.21		1	
Clostridium butyricum		4.36	0.09		99	
03355502	**Low discrimination**					**03355502**
Bifidobacterium adolescentis		72.78	0.56	MNS 87		
Actinomyces israelii		26.38	0.55	RAF 92		
Lactobacillus fermentum		0.48	0.24			
03355510	**Low discrimination**					**03355510**
Actinomyces israelii		74.52	0.38	MLZ 13		
Bifidobacterium adolescentis		25.48	0.23			
03355512	**Very good identification**					**03355512**
Bifidobacterium adolescentis		95.60	0.57	MNS 87		
Actinomyces israelii		4.24	0.41	RAF 92		
Clostridium clostridioforme		0.16	0.07			
03355540	**Good identification**					**03355540**
Actinomyces israelii		100.0	0.45			
03355542	**Low discrimination**					**03355542**
Actinomyces israelii		53.28	0.48			
Bifidobacterium adolescentis		46.72	0.40			
03355552	**Low discrimination**					**03355552**
Bifidobacterium adolescentis		87.76	0.42	MNS 87		
Actinomyces israelii		12.24	0.34	RAF 92		
03355702	**Low discrimination**					**03355702**
Actinomyces israelii		46.59	0.35	RHA 7		
Eubacterium contortum		30.51	0.17			
Bifidobacterium adolescentis		17.25	0.22			
03357000	**Good identification**					**03357000**
					SPO	
Actinomyces israelii		98.48	0.56		1	
Clostridium innocuum		1.52	0.10		99	
03357002	**Very good identification**					**03357002**
					SPO	
Actinomyces israelii		98.98	0.59		1	
Clostridium innocuum		1.02	0.10		99	
03357010	**Good identification**					**03357010**
Actinomyces israelii		100.0	0.42	MLZ 13		
03357012	**Good identification**					**03357012**
					SPO	
Actinomyces israelii		96.99	0.45	MLZ 13	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		1.56	0.08		1	
Clostridium tertium		1.45	-.01		99	
03357020	**Good identification**					**03357020**
Actinomyces israelii		100.0	0.35	NAG 6		
03357022	**Acceptable identification**					**03357022**
					SPO	
Actinomyces israelii		94.42	0.38	NAG 6	1	
Clostridium tertium		5.58	0.03		99	
03357040	**Good identification**					**03357040**
Actinomyces israelii		100.0	0.49			
03357042	**Good identification**					**03357042**
Actinomyces israelii		100.0	0.52			
03357052	**Good identification**					**03357052**
Actinomyces israelii		100.0	0.38	MLZ 13		
03357100	**Very good identification**					**03357100**
					SPO	
Actinomyces israelii		98.36	0.66		1	
Clostridium tertium		1.53	0.21		99	
Actinomyces naeslundii		0.11	0.25		1	
03357102	**Good identification**					**03357102**
					SPO	
Actinomyces israelii		92.47	0.69		1	
Bifidobacterium adolescentis		4.50	0.40		1	
Clostridium tertium		2.88	0.29		99	
03357110	**Low discrimination**					**03357110**
					SPO	
Actinomyces israelii		83.95	0.52	MLZ 13	1	
Clostridium tertium		15.55	0.25		99	
Bifidobacterium adolescentis		0.51	0.08		1	
03357112	**Low discrimination**					**03357112**
					SPO	
Actinomyces israelii		56.54	0.55	MLZ 13	1	
Bifidobacterium adolescentis		22.52	0.42		1	
Clostridium tertium		20.94	0.33		99	
03357120	**Low discrimination**					**03357120**
					SPO	
Actinomyces israelii		57.15	0.46	NAG 6	1	
Clostridium tertium		41.82	0.29		99	
Actinomyces naeslundii		1.03	0.25		1	
03357122	**Low discrimination**					**03357122**
					SPO	
Clostridium tertium		58.49	0.37	MNS 99	99	
Actinomyces israelii		39.97	0.49	TRE 90	1	
Clostridium barati		0.56	-.01		99	
03357132	**Acceptable identification**					**03357132**
					SPO	
Clostridium tertium		94.25	0.41	MNS 99	99	
Actinomyces israelii		5.41	0.35	TRE 90	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Bifidobacterium adolescentis	0.34	0.08		1	
03357140	**Very good identification**					**03357140**
	Actinomyces israelii	100.0	0.59			
03357142	**Very good identification**					**03357142**
	Actinomyces israelii	98.48	0.62			
	Bifidobacterium adolescentis	1.52	0.25			
03357150	**Good identification**					**03357150**
	Actinomyces israelii	100.0	0.45	MLZ 13		
03357152	**Low discrimination**					**03357152**
					SPO	
	Actinomyces israelii	88.01	0.48	MLZ 13	1	
	Bifidobacterium adolescentis	11.14	0.26		1	
	Clostridium tertium	0.85	-.01		99	
03357160	**Good identification**					**03357160**
	Actinomyces israelii	100.0	0.39	NAG 6		
03357162	**Good identification**					**03357162**
					SPO	
	Actinomyces israelii	96.34	0.42	NAG 6	1	
	Clostridium tertium	3.66	0.03		99	
03357200	**Good identification**					**03357200**
	Actinomyces israelii	100.0	0.37	RHA 7		
03357202	**Good identification**					**03357202**
	Actinomyces israelii	100.0	0.40	RHA 7		
03357300	**Good identification**					**03357300**
	Actinomyces israelii	100.0	0.47	RHA 7		
03357302	**Good identification**					**03357302**
	Actinomyces israelii	100.0	0.50	RHA 7		
03357312	**Acceptable identification**					**03357312**
					SPO	
	Actinomyces israelii	90.65	0.36	RHA 7	1	
	Bifidobacterium adolescentis	4.85	0.08	MLZ 13	1	
	Clostridium tertium	4.51	-.01		99	
03357340	**Good identification**					**03357340**
	Actinomyces israelii	100.0	0.40	RHA 7		
03357342	**Good identification**					**03357342**
	Actinomyces israelii	100.0	0.43	RHA 7		
03357400	**Good identification**					**03357400**
	Actinomyces israelii	100.0	0.54			
03357402	**Very good identification**					**03357402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		98.42	0.57			
Bifidobacterium adolescentis		1.58	0.20			
03357410	**Good identification**					**03357410**
Actinomyces israelii		100.0	0.40	MLZ 13		
03357412	**Low discrimination**					**03357412**
Actinomyces israelii		88.37	0.43	MLZ 13		
Bifidobacterium adolescentis		11.63	0.22			
03357422	**Good identification**					**03357422**
Actinomyces israelii		100.0	0.37	NAG 6		
03357440	**Good identification**					**03357440**
Actinomyces israelii		100.0	0.47			
03357442	**Good identification**					**03357442**
Actinomyces israelii		100.0	0.50			
03357452	**Good identification**					**03357452**
Actinomyces israelii		100.0	0.36	MLZ 13		
03357500	**Very good identification**					**03357500**
					SPO	
Actinomyces israelii		99.30	0.65		1	
Bifidobacterium adolescentis		0.60	0.20		1	
Clostridium butyricum		0.10	-.08		99	
03357502	**Low discrimination**					**03357502**
Actinomyces israelii		71.52	0.68			
Bifidobacterium adolescentis		28.48	0.54			
03357510	**Good identification**					**03357510**
Actinomyces israelii		95.30	0.50	MLZ 13		
Bifidobacterium adolescentis		4.70	0.22			
03357512	**Low discrimination**					**03357512**
					SPO	
Bifidobacterium adolescentis		76.43	0.56	MNS 87	1	
Actinomyces israelii		23.46	0.53	RAF 92	1	
Clostridium tertium		0.11	-.01		99	
03357520	**Good identification**					**03357520**
Actinomyces israelii		100.0	0.44	NAG 6		
03357522	**Acceptable identification**					**03357522**
					SPO	
Actinomyces israelii		92.50	0.47	NAG 6	1	
Bifidobacterium adolescentis		5.83	0.20		1	
Clostridium tertium		1.67	0.03		99	
03357540	**Very good identification**					**03357540**
Actinomyces israelii		100.0	0.58			
03357542	**Low discrimination**					**03357542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		88.76	0.61			
Bifidobacterium adolescentis		11.24	0.39			
03357550	**Good identification**					**03357550**
Actinomyces israelii		100.0	0.43	MLZ 13		
03357552	**Low discrimination**					**03357552**
Bifidobacterium adolescentis		50.87	0.40	MNS 87		
Actinomyces israelii		49.13	0.46	RAF 92		
03357560	**Good identification**					**03357560**
Actinomyces israelii		100.0	0.37	NAG 6		
03357562	**Good identification**					**03357562**
Actinomyces israelii		100.0	0.40	NAG 6		
03357600	**Good identification**					**03357600**
Actinomyces israelii		100.0	0.35	RHA 7		
03357602	**Good identification**					**03357602**
Actinomyces israelii		100.0	0.38	RHA 7		
03357700	**Good identification**					**03357700**
Actinomyces israelii		100.0	0.45	RHA 7		
03357702	**Acceptable identification**					**03357702**
Actinomyces israelii		94.93	0.48	RHA 7		
Bifidobacterium adolescentis		5.07	0.20			
03357740	**Good identification**					**03357740**
Actinomyces israelii		100.0	0.38	RHA 7		
03357742	**Good identification**					**03357742**
Actinomyces israelii		100.0	0.41	RHA 7		
03361101	**Good identification**					**03361101**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03361103	**Good identification**					**03361103**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03361120	**Low discrimination**					**03361120**
					COC	
Actinomyces naeslundii		72.01	0.37	MNS 88	1	
Atopobium parvulum		27.99	0.07	RAF 92	99	
03361121	**Good identification**					**03361121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03361123	**Good identification**					**03361123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03363001	**Good identification**					**03363001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	MNS 88		
03363003	**Good identification**					**03363003**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03363021	**Good identification**					**03363021**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03363023	**Good identification**					**03363023**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03363100	**Low discrimination**					**03363100**
					SPO	
Actinomyces naeslundii		72.89	0.42	MNS 88	1	
Actinomyces israelii		20.70	0.24	RAF 92	1	
Clostridium tertium		6.42	0.01	FRU 95	99	
03363101	**Good identification**					**03363101**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03363102	**Low discrimination**					**03363102**
Actinomyces naeslundii		49.12	0.42	MNS 88		
Actinomyces israelii		20.92	0.27	RAF 92		
Lactobacillus acidophilus		16.99	0.10	FRU 95		
03363103	**Good identification**					**03363103**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03363120	**Low discrimination**					**03363120**
					SPO	
Actinomyces naeslundii		66.08	0.42	MNS 88	1	
Clostridium tertium		17.46	0.09	RAF 92	99	
Clostridium ramosum		16.46	0.07	FRU 95	92	
03363121	**Good identification**					**03363121**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03363122	**Low discrimination**					**03363122**
					SPO	
Actinomyces naeslundii		31.90	0.42	MNS 88	1	
Clostridium ramosum		31.78	0.17	RAF 92	92	
Clostridium tertium		25.29	0.17	FRU 95	99	
03363123	**Good identification**					**03363123**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03367100	**Good identification**					**03367100**
					SPO	
Actinomyces israelii		96.41	0.42	FRU 99	1	
Clostridium tertium		3.59	0.03		99	
03367102	**Acceptable identification**					**03367102**
					SPO	
Actinomyces israelii		93.07	0.45	FRU 99	1	
Clostridium tertium		6.93	0.11		99	
03367142	**Good identification**					**03367142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.38	FRU 99			
03367500	**Good identification**						**03367500**
Actinomyces israelii		100.0	0.40	FRU 99			
03367502	**Acceptable identification**						**03367502**
Actinomyces israelii		93.41	0.43	FRU 99			
Bifidobacterium adolescentis		6.59	0.17				
03367542	**Good identification**						**03367542**
Actinomyces israelii		100.0	0.36	FRU 99			
03370000	**Low discrimination**						**03370000**
					GRA		
Fusobacterium mortiferum		60.76	0.45	MLT 11	1		
Lactobacillus jensenii		38.02	0.32		99		
Actinomyces naeslundii		1.23	0.28		99		
03370100	**Acceptable identification**						**03370100**
					COC	GRA	
Fusobacterium mortiferum		91.49	0.51	MLT 11	1	1	
Atopobium parvulum		6.26	0.20		99	99	
Actinomyces naeslundii		2.25	0.36		1	99	
03370101	**Low discrimination**						**03370101**
					GRA		
Actinomyces naeslundii		78.51	0.39	MNS 88	99		
Fusobacterium mortiferum		21.49	0.17	RAF 92	1		
03370103	**Good identification**						**03370103**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03370120	**Low discrimination**						**03370120**
					COC	SPO	
Actinomyces naeslundii		41.73	0.36	MNS 88	1	1	
Clostridium barati		20.64	0.04	RAF 92	1	99	
Atopobium parvulum		20.49	0.07	GAL 96	99	1	
03370121	**Good identification**						**03370121**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03370123	**Good identification**						**03370123**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03371000	**Very good identification**						**03371000**
					GRA		
Lactobacillus jensenii		95.57	0.66	MNS 99	99		
Fusobacterium mortiferum		2.31	0.48		1		
Actinomyces naeslundii		0.75	0.51		99		
03371001	**Low discrimination**						**03371001**
					GRA		
Actinomyces naeslundii		53.11	0.54	MNS 88	99		
Lactobacillus jensenii		45.78	0.32	RAF 92	99		
Fusobacterium mortiferum		1.11	0.14		1		
03371002	**Low discrimination**						**03371002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		81.31	0.63	MNS 99		
Lactobacillus acidophilus		15.67	0.48			
Actinomyces naeslundii		0.95	0.51			
03371003	**Low discrimination**					**03371003**
Actinomyces naeslundii		59.33	0.54	MNS 88		
Lactobacillus jensenii		34.10	0.29	RAF 92		
Lactobacillus acidophilus		6.57	0.14			
03371010	**Very good identification**					**03371010**
Lactobacillus jensenii		99.68	0.58	MNS 99		
Actinomyces israelii		0.13	0.20			
Actinomyces naeslundii		0.12	0.29			
03371012	**Good identification**					**03371012**
Lactobacillus jensenii		98.71	0.55	MNS 99		
Lactobacillus acidophilus		0.58	0.14			
Actinomyces israelii		0.28	0.23			
03371020	**Acceptable identification**					**03371020**
					SPO	
Lactobacillus jensenii		93.94	0.56	MNS 99	1	
Actinomyces naeslundii		2.94	0.51		1	
Clostridium barati		1.09	0.17		99	
03371021	**Low discrimination**					**03371021**
Actinomyces naeslundii		82.27	0.54	MNS 88		
Lactobacillus jensenii		17.73	0.21	RAF 92		
03371022	**Low discrimination**					**03371022**
					SPO	
Lactobacillus jensenii		52.95	0.53	MNS 99	1	
Lactobacillus acidophilus		40.82	0.48		1	
Clostridium barati		2.77	0.25		99	
03371023	**Low discrimination**					**03371023**
Actinomyces naeslundii		79.72	0.54	MNS 88		
Lactobacillus jensenii		11.45	0.18	RAF 92		
Lactobacillus acidophilus		8.83	0.14			
03371030	**Good identification**					**03371030**
Lactobacillus jensenii		99.51	0.47	MNS 99		
Actinomyces naeslundii		0.49	0.29			
03371032	**Good identification**					**03371032**
Lactobacillus jensenii		97.02	0.44	MNS 99		
Lactobacillus acidophilus		2.27	0.14			
Actinomyces naeslundii		0.72	0.29			
03371041	**Good identification**					**03371041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03371043	**Acceptable identification**					**03371043**
Actinomyces naeslundii		92.80	0.41	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		7.20	-.02	RAF 92			
03371061	**Good identification**						**03371061**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03371063	**Acceptable identification**						**03371063**
Actinomyces naeslundii		92.80	0.41	MNS 88			
Lactobacillus acidophilus		7.20	-.02	RAF 92			
03371100	**Low discrimination**						**03371100**
					COC	GRA	
Atopobium parvulum		66.37	0.54	MNS 99	99	99	
Fusobacterium mortiferum		14.70	0.54		1	1	
Actinomyces naeslundii		5.78	0.59		1	99	
03371101	**Good identification**						**03371101**
					COC	GRA	
Actinomyces naeslundii		91.38	0.62	MNS 88	1	99	
Atopobium parvulum		7.06	0.20	RAF 92	99	99	
Fusobacterium mortiferum		1.56	0.20		1	1	
03371102	**Good identification**						**03371102**
					COC		
Lactobacillus acidophilus		92.52	0.66	MNS 99	1		
Atopobium parvulum		2.82	0.41		99		
Actinomyces naeslundii		1.39	0.59		1		
03371103	**Low discrimination**						**03371103**
					COC		
Actinomyces naeslundii		68.20	0.62	MNS 88	1		
Lactobacillus acidophilus		30.52	0.32	RAF 92	1		
Atopobium parvulum		0.93	0.07		99		
03371111	**Good identification**						**03371111**
Actinomyces naeslundii		100.0	0.40	MNS 88			
03371113	**Low discrimination**						**03371113**
					GRA		
Actinomyces naeslundii		75.81	0.40	MNS 88	99		
Leptotrichia buccalis		24.19	-.03	RAF 92	1		
03371120	**Low discrimination**						**03371120**
					COC		
Atopobium parvulum		31.71	0.41	MNS 99	99		
Actinomyces naeslundii		15.66	0.59		1		
Lactobacillus acidophilus		10.51	0.32		1		
03371121	**Very good identification**						**03371121**
					COC		
Actinomyces naeslundii		98.65	0.62	MNS 88	1		
Atopobium parvulum		1.35	0.07	RAF 92	99		
03371122	**Good identification**						**03371122**
					SPO		
Lactobacillus acidophilus		92.78	0.66	MNS 99	1		
Clostridium ramosum		2.48	0.38		92		
Clostridium barati		1.64	0.34		99		
03371123	**Low discrimination**						**03371123**
					SPO		
Actinomyces naeslundii		68.15	0.62	MNS 88	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		30.50	0.32	RAF 92	1	
Clostridium ramosum		0.82	0.04		92	
03371130	**Low discrimination**					**03371130**
					SPO	
Actinomyces naeslundii		33.12	0.37	MNS 88	1	
Clostridium tertium		30.19	0.13	RAF 92	99	
Lactobacillus jensenii		23.82	0.13	MLZ 5	1	
03371131	**Good identification**					**03371131**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03371133	**Good identification**					**03371133**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03371140	**Low discrimination**					**03371140**
					COC	
Actinomyces israelii		32.34	0.38	XYL 92	1	
Actinomyces naeslundii		28.96	0.46		1	
Atopobium parvulum		19.02	0.20		99	
03371141	**Good identification**					**03371141**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03371142	**Acceptable identification**					**03371142**
Lactobacillus acidophilus		93.87	0.50	MNS 99		
Actinomyces israelii		3.38	0.41	SOR 11		
Actinomyces naeslundii		2.02	0.46			
03371143	**Low discrimination**					**03371143**
Actinomyces naeslundii		76.13	0.49	MNS 88		
Lactobacillus acidophilus		23.87	0.16	RAF 92		
03371160	**Low discrimination**					**03371160**
Actinomyces naeslundii		38.17	0.46	MNS 88		
Clostridium perfringens		21.39	0.20	RAF 92		
Lactobacillus acidophilus		17.95	0.16			
03371161	**Good identification**					**03371161**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03371162	**Good identification**					**03371162**
					SPO	
Lactobacillus acidophilus		95.60	0.50	MNS 99	1	
Actinomyces naeslundii		2.05	0.46	SOR 11	1	
Clostridium barati		1.19	0.15		99	
03371163	**Low discrimination**					**03371163**
Actinomyces naeslundii		76.13	0.49	MNS 88		
Lactobacillus acidophilus		23.87	0.16	RAF 92		
03371302	**Good identification**					**03371302**
					SPO	
Lactobacillus acidophilus		97.17	0.47	MNS 99	1	
Actinomyces israelii		1.11	0.29	RHA 7	1	
Clostridium barati		0.66	0.07		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
03371322	**Low discrimination**						**03371322**
					SPO		
Lactobacillus acidophilus		66.59	0.47	MNS 99	1		
Clostridium ramosum		31.37	0.40	RHA 7	92		
Clostridium barati		1.36	0.15		99		
03371500	**Low discrimination**						**03371500**
					COC	SPO	
Actinomyces israelii		69.28	0.43	XYL 92	1	1	
Atopobium parvulum		19.37	0.20		99	1	
Clostridium butyricum		5.38	0.03		1	99	
03371512	**Low discrimination**						**03371512**
					GRA		
Bifidobacterium adolescentis		83.53	0.40	MNS 87	99		
Actinomyces israelii		12.16	0.32	RAF 92	99		
Leptotrichia buccalis		3.30	-.03	XYL 87	1		
03371540	**Good identification**						**03371540**
Actinomyces israelii		100.0	0.36	XYL 92			
03371542	**Low discrimination**						**03371542**
Actinomyces israelii		62.11	0.39	XYL 92			
Lactobacillus acidophilus		21.31	0.16				
Bifidobacterium adolescentis		16.58	0.23				
03372001	**Good identification**						**03372001**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03372003	**Good identification**						**03372003**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03372021	**Good identification**						**03372021**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03372023	**Good identification**						**03372023**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03372100	**Low discrimination**						**03372100**
					GRA		
Actinomyces naeslundii		55.85	0.40	MNS 88	99		
Actinomyces israelii		20.03	0.24	RAF 92	99		
Fusobacterium mortiferum		11.82	0.17	GAL 96	1		
03372101	**Good identification**						**03372101**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03372102	**Low discrimination**						**03372102**
Actinomyces naeslundii		40.45	0.40	MNS 88			
Actinomyces israelii		21.76	0.27	RAF 92			
Lactobacillus acidophilus		17.67	0.10	GAL 96			
03372103	**Good identification**						**03372103**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03372120	**Low discrimination**						**03372120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		60.67	0.40	MNS 88	1	
Clostridium tertium		20.25	0.09	RAF 92	99	
Clostridium ramosum		19.08	0.07	GAL 96	92	
03372121	**Good identification**					**03372121**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03372123	**Good identification**					**03372123**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03373000	**Low discrimination**					**03373000**
Actinomyces israelii		34.19	0.47	XYL 92		
Actinomyces naeslundii		32.48	0.56			
Lactobacillus jensenii		21.63	0.32			
03373001	**Very good identification**					**03373001**
Actinomyces naeslundii		99.30	0.59	MNS 88		
Actinomyces israelii		0.70	0.13	RAF 92		
03373002	**Low discrimination**					**03373002**
Actinomyces israelii		40.95	0.50	XYL 92		
Actinomyces naeslundii		25.93	0.56			
Lactobacillus acidophilus		11.57	0.26			
03373003	**Very good identification**					**03373003**
Actinomyces naeslundii		98.95	0.59	MNS 88		
Actinomyces israelii		1.05	0.16	RAF 92		
03373011	**Good identification**					**03373011**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03373012	**Low discrimination**					**03373012**
					SPO	
Actinomyces israelii		45.31	0.36	XYL 92	1	
Lactobacillus jensenii		28.43	0.21	MLZ 13	1	
Clostridium tertium		13.51	0.13		99	
03373013	**Good identification**					**03373013**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03373020	**Low discrimination**					**03373020**
Actinomyces naeslundii		75.35	0.56	MNS 88		
Lactobacillus jensenii		12.55	0.21	RAF 92		
Actinomyces israelii		5.06	0.27			
03373021	**Very good identification**					**03373021**
Actinomyces naeslundii		100.0	0.59	MNS 88		
03373022	**Low discrimination**					**03373022**
					SPO	
Actinomyces naeslundii		50.62	0.56	MNS 88	1	
Lactobacillus acidophilus		22.57	0.26	RAF 92	1	
Clostridium ramosum		7.55	0.17		92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03373023	**Very good identification**					**03373023**
Actinomyces naeslundii		100.0	0.59	MNS 88		
03373031	**Good identification**					**03373031**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03373033	**Good identification**					**03373033**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03373040	**Low discrimination**					**03373040**
Actinomyces israelii		69.88	0.40	XYL 92		
Actinomyces naeslundii		30.12	0.43			
03373041	**Good identification**					**03373041**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03373042	**Low discrimination**					**03373042**
Actinomyces israelii		72.61	0.43	XYL 92		
Actinomyces naeslundii		20.87	0.43			
Lactobacillus acidophilus		6.52	0.10			
03373043	**Good identification**					**03373043**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03373060	**Low discrimination**					**03373060**
Actinomyces naeslundii		87.10	0.43	MNS 88		
Actinomyces israelii		12.90	0.20	RAF 92		
03373061	**Good identification**					**03373061**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03373062	**Low discrimination**					**03373062**
Actinomyces naeslundii		65.17	0.43	MNS 88		
Lactobacillus acidophilus		20.36	0.10	RAF 92		
Actinomyces israelii		14.47	0.23			
03373063	**Good identification**					**03373063**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03373100	**Low discrimination**					**03373100**
					SPO	
Actinomyces israelii		48.46	0.58	XYL 92	1	
Actinomyces naeslundii		32.76	0.64		1	
Clostridium tertium		15.03	0.35		99	
03373101	**Very good identification**					**03373101**
					SPO	
Actinomyces naeslundii		98.71	0.67	MNS 88	1	
Actinomyces israelii		0.98	0.24	RAF 92	1	
Clostridium tertium		0.31	0.01		99	
03373102	**Low discrimination**					**03373102**
					SPO	
Actinomyces israelii		34.05	0.61	XYL 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		27.65	0.44		1	
Clostridium tertium		21.12	0.43		99	
03373103	**Very good identification**					**03373103**
Actinomyces naeslundii		96.49	0.67	MNS 88		
Actinomyces israelii		1.44	0.27	RAF 92		
Lactobacillus acidophilus		1.17	0.10			
03373110	**Low discrimination**					**03373110**
					SPO	
Clostridium tertium		74.88	0.39	MNS 99	99	
Actinomyces israelii		20.29	0.44		1	
Actinomyces naeslundii		4.83	0.42		1	
03373111	**Acceptable identification**					**03373111**
					SPO	
Actinomyces naeslundii		90.55	0.45	MNS 88	1	
Clostridium tertium		9.45	0.05	RAF 92	99	
03373112	**Low discrimination**					**03373112**
					SPO	
Clostridium tertium		84.31	0.47	MNS 99	99	
Actinomyces israelii		11.42	0.47		1	
Actinomyces naeslundii		1.81	0.42		1	
03373113	**Low discrimination**					**03373113**
					SPO	
Actinomyces naeslundii		76.16	0.45	MNS 88	1	
Clostridium tertium		23.85	0.13	RAF 92	99	
03373120	**Low discrimination**					**03373120**
					SPO	
Clostridium tertium		36.09	0.43	MNS 99	99	
Clostridium ramosum		34.01	0.41		92	
Actinomyces naeslundii		26.21	0.64		1	
03373121	**Very good identification**					**03373121**
					SPO	
Actinomyces naeslundii		98.23	0.67	MNS 88	1	
Clostridium tertium		0.91	0.09	RAF 92	99	
Clostridium ramosum		0.86	0.07		92	
03373122	**Low discrimination**					**03373122**
					SPO	
Clostridium ramosum		42.14	0.51	MNS 99	92	
Clostridium tertium		33.53	0.51		99	
Lactobacillus acidophilus		14.63	0.44		1	
03373123	**Good identification**					**03373123**
					SPO	
Actinomyces naeslundii		93.03	0.67	MNS 88	1	
Clostridium ramosum		3.25	0.17	RAF 92	92	
Clostridium tertium		2.59	0.17		99	
03373130	**Good identification**					**03373130**
					SPO	
Clostridium tertium		96.84	0.47	MNS 99	99	
Actinomyces naeslundii		2.08	0.42		1	
Actinomyces israelii		0.56	0.23		1	
03373131	**Low discrimination**					**03373131**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		76.16	0.45	MNS 88	1	
Clostridium tertium		23.85	0.13	RAF 92	99	
03373132	**Good identification**					**03373132**
					SPO	
Clostridium tertium		98.07	0.55	MNS 99	99	
Actinomyces naeslundii		0.70	0.42		1	
Clostridium ramosum		0.70	0.17		92	
03373133	**Low discrimination**					**03373133**
					SPO	
Actinomyces naeslundii		51.56	0.45	MNS 88	1	
Clostridium tertium		48.44	0.21	RAF 92	99	
03373140	**Low discrimination**					**03373140**
					SPO	
Actinomyces israelii		75.60	0.51	XYL 92	1	
Actinomyces naeslundii		23.19	0.51		1	
Clostridium tertium		0.61	0.01		99	
03373141	**Good identification**					**03373141**
Actinomyces naeslundii		97.85	0.54	MNS 88		
Actinomyces israelii		2.15	0.17	RAF 92		
03373142	**Low discrimination**					**03373142**
Actinomyces israelii		67.06	0.54	XYL 92		
Lactobacillus acidophilus		17.31	0.28			
Actinomyces naeslundii		13.71	0.51			
03373143	**Good identification**					**03373143**
Actinomyces naeslundii		96.81	0.54	MNS 88		
Actinomyces israelii		3.19	0.20	RAF 92		
03373150	**Low discrimination**					**03373150**
					SPO	
Actinomyces israelii		83.07	0.37	XYL 92	1	
Actinomyces naeslundii		8.97	0.29	MLZ 13	1	
Clostridium tertium		7.96	0.05		99	
03373152	**Low discrimination**					**03373152**
					SPO	
Actinomyces israelii		76.79	0.40	XYL 92	1	
Clostridium tertium		14.72	0.13	MLZ 13	99	
Actinomyces naeslundii		5.53	0.29		1	
03373160	**Low discrimination**					**03373160**
					SPO	
Actinomyces naeslundii		73.47	0.51	MNS 88	1	
Actinomyces israelii		15.29	0.30	RAF 92	1	
Clostridium tertium		5.79	0.09		99	
03373161	**Good identification**					**03373161**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03373163	**Good identification**					**03373163**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03373300	**Low discrimination**					**03373300**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		77.62	0.39	XYL 92	1	
Clostridium ramosum		12.11	0.09	RHA 7	92	
Actinomyces naeslundii		7.04	0.30		1	
03373302	**Low discrimination**					**03373302**
					SPO	
Actinomyces israelii		42.16	0.42	XYL 92	1	
Lactobacillus acidophilus		34.24	0.25	RHA 7	1	
Clostridium ramosum		17.54	0.19		92	
03373320	**Good identification**					**03373320**
					SPO	
Clostridium ramosum		98.22	0.43	MNS 99	92	
Clostridium tertium		0.79	0.09		99	
Actinomyces naeslundii		0.58	0.30		1	
03373322	**Good identification**					**03373322**
					SPO	
Clostridium ramosum		97.14	0.53	MNS 99	92	
Lactobacillus acidophilus		1.92	0.25		1	
Clostridium tertium		0.59	0.17		99	
03373400	**Acceptable identification**					**03373400**
					SPO	
Actinomyces israelii		93.65	0.46	XYL 92	1	
Clostridium innocuum		6.35	0.11		99	
03373402	**Good identification**					**03373402**
					SPO	
Actinomyces israelii		95.68	0.49	XYL 92	1	
Clostridium innocuum		4.32	0.11		99	
03373440	**Good identification**					**03373440**
Actinomyces israelii		100.0	0.39	XYL 92		
03373442	**Good identification**					**03373442**
Actinomyces israelii		100.0	0.42	XYL 92		
03373500	**Very good identification**					**03373500**
					SPO	
Actinomyces israelii		98.80	0.56	XYL 92	1	
Actinomyces naeslundii		0.82	0.30		1	
Clostridium tertium		0.38	0.01		99	
03373502	**Low discrimination**					**03373502**
Actinomyces israelii		87.37	0.59	XYL 92		
Bifidobacterium adolescentis		10.60	0.37			
Lactobacillus acidophilus		0.88	0.10			
03373510	**Good identification**					**03373510**
					SPO	
Actinomyces israelii		95.64	0.42	XYL 92	1	
Clostridium tertium		4.36	0.05	MLZ 13	99	
03373512	**Low discrimination**					**03373512**
					SPO	
Actinomyces israelii		48.00	0.45	XYL 92	1	
Bifidobacterium adolescentis		47.62	0.39	MLZ 13	1	
Clostridium tertium		4.37	0.13		99	
03373520	**Low discrimination**					**03373520**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		67.55	0.36	XYL 92	1	
Clostridium tertium		12.16	0.09	NAG 6	99	
Clostridium ramosum		11.46	0.07		92	
03373522	**Low discrimination**					**03373522**
					SPO	
Actinomyces israelii		48.63	0.39	XYL 92	1	
Clostridium ramosum		22.00	0.17	NAG 6	92	
Clostridium tertium		17.50	0.17		99	
03373540	**Good identification**					**03373540**
Actinomyces israelii		100.0	0.49	XYL 92		
03373542	**Good identification**					**03373542**
Actinomyces israelii		96.29	0.52	XYL 92		
Bifidobacterium adolescentis		3.71	0.22			
03373550	**Good identification**					**03373550**
Actinomyces israelii		100.0	0.35	XYL 92		
03373552	**Low discrimination**					**03373552**
Actinomyces israelii		76.03	0.38	XYL 92		
Bifidobacterium adolescentis		23.97	0.23	MLZ 13		
03373700	**Good identification**					**03373700**
Actinomyces israelii		100.0	0.37	XYL 92		
03373702	**Good identification**					**03373702**
Actinomyces israelii		100.0	0.40	XYL 92		
03375000	**Low discrimination**					**03375000**
					COC	
Actinomyces israelii		65.97	0.53		1	
Lactobacillus jensenii		20.57	0.32		1	
Atopobium parvulum		9.75	0.25		99	
03375002	**Low discrimination**					**03375002**
Actinomyces israelii		84.26	0.56			
Lactobacillus jensenii		11.68	0.29			
Lactobacillus acidophilus		2.25	0.14			
03375010	**Low discrimination**					**03375010**
Actinomyces israelii		58.98	0.39	MLZ 13		
Lactobacillus jensenii		41.02	0.24			
03375012	**Low discrimination**					**03375012**
Actinomyces israelii		76.39	0.42	MLZ 13		
Lactobacillus jensenii		23.61	0.21			
03375022	**Low discrimination**					**03375022**
Actinomyces israelii		50.99	0.35	NAG 6		
Lactobacillus jensenii		27.68	0.18			
Lactobacillus acidophilus		21.34	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03375040	**Good identification**						**03375040**
Actinomyces israelii		100.0	0.46				
03375042	**Good identification**						**03375042**
Actinomyces israelii		99.16	0.49				
Lactobacillus acidophilus		0.84	-.02				
03375100	**Low discrimination**						**03375100**
					COC	GRA	
Atopobium parvulum		77.59	0.59	MNS 99	99	99	
Actinomyces israelii		21.22	0.63		1	99	
Fusobacterium mortiferum		0.70	0.35		1	1	
03375102	**Low discrimination**						**03375102**
					COC		
Actinomyces israelii		63.30	0.66		1		
Atopobium parvulum		27.23	0.46		99		
Lactobacillus acidophilus		4.86	0.32		1		
03375110	**Low discrimination**						**03375110**
					COC	SPO	
Actinomyces israelii		78.58	0.49	MLZ 13	1	1	
Atopobium parvulum		19.43	0.25		99	1	
Clostridium tertium		1.99	0.07		1	99	
03375112	**Low discrimination**						**03375112**
					SPO		
Actinomyces israelii		62.97	0.52	MLZ 13	1		
Bifidobacterium adolescentis		30.79	0.40		1		
Clostridium tertium		3.20	0.15		99		
03375120	**Low discrimination**						**03375120**
					COC	SPO	
Atopobium parvulum		89.75	0.46	MNS 99	99	1	
Actinomyces israelii		8.88	0.43		1	1	
Clostridium tertium		0.89	0.11		1	99	
03375140	**Acceptable identification**						**03375140**
					COC	SPO	
Actinomyces israelii		90.96	0.56		1	1	
Atopobium parvulum		8.64	0.25		99	1	
Clostridium butyricum		0.40	-.05		1	99	
03375142	**Good identification**						**03375142**
Actinomyces israelii		94.83	0.59				
Lactobacillus acidophilus		2.31	0.16				
Bifidobacterium adolescentis		1.80	0.23				
03375150	**Good identification**						**03375150**
Actinomyces israelii		100.0	0.42	MLZ 13			
03375152	**Low discrimination**						**03375152**
Actinomyces israelii		86.55	0.45	MLZ 13			
Bifidobacterium adolescentis		13.45	0.25				
03375160	**Low discrimination**						**03375160**
					COC		
Actinomyces israelii		79.20	0.36	NAG 6	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		20.80	0.12		99		
03375162	**Low discrimination**						**03375162**
Actinomyces israelii		72.34	0.39	NAG 6			
Lactobacillus acidophilus		27.66	0.16				
03375202	**Good identification**						**03375202**
Actinomyces israelii		100.0	0.37	RHA 7			
03375300	**Low discrimination**						**03375300**
					COC		
Actinomyces israelii		67.08	0.44	RHA 7	1		
Atopobium parvulum		32.92	0.25		99		
03375302	**Low discrimination**						**03375302**
					COC		
Actinomyces israelii		87.19	0.47	RHA 7	1		
Lactobacillus acidophilus		6.70	0.12		1		
Atopobium parvulum		5.03	0.12		99		
03375340	**Good identification**						**03375340**
Actinomyces israelii		100.0	0.37	RHA 7			
03375342	**Good identification**						**03375342**
Actinomyces israelii		100.0	0.40	RHA 7			
03375400	**Good identification**						**03375400**
					SPO		
Actinomyces israelii		96.49	0.51		1		
Clostridium butyricum		3.51	0.06		99		
03375402	**Good identification**						**03375402**
					SPO		
Actinomyces israelii		97.16	0.54		1		
Bifidobacterium adolescentis		1.92	0.19		1		
Clostridium butyricum		0.92	-.02		99		
03375410	**Good identification**						**03375410**
Actinomyces israelii		100.0	0.37	MLZ 13			
03375412	**Low discrimination**						**03375412**
Actinomyces israelii		80.00	0.40	MLZ 13			
Bifidobacterium adolescentis		12.93	0.20				
Clostridium clostridioforme		7.07	0.13				
03375440	**Good identification**						**03375440**
Actinomyces israelii		100.0	0.44				
03375442	**Good identification**						**03375442**
Actinomyces israelii		100.0	0.47				
03375500	**Low discrimination**						**03375500**
					COC	SPO	
Actinomyces israelii		78.68	0.62		1	1	
Clostridium butyricum		17.19	0.29		1	99	
Atopobium parvulum		3.55	0.25		99	1	
03375502	**Low discrimination**						**03375502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		64.38	0.65		1		
Bifidobacterium adolescentis		31.47	0.53		1		
Clostridium butyricum		3.65	0.22		99		
03375510	**Low discrimination**						**03375510**
					SPO		
Actinomyces israelii		87.91	0.47	MLZ 13	1		
Clostridium butyricum		6.76	0.07		99		
Bifidobacterium adolescentis		5.33	0.20		1		
03375512	**Low discrimination**						**03375512**
Bifidobacterium adolescentis		78.52	0.54	MNS 87			
Actinomyces israelii		19.64	0.50	RAF 92			
Clostridium clostridioforme		1.45	0.22				
03375520	**Low discrimination**						**03375520**
					COC	SPO	
Actinomyces israelii		70.30	0.41	NAG 6	1	1	
Clostridium butyricum		20.92	0.11		1	99	
Atopobium parvulum		8.78	0.12		99	1	
03375522	**Low discrimination**						**03375522**
					SPO		
Actinomyces israelii		81.80	0.44	NAG 6	1		
Bifidobacterium adolescentis		6.33	0.19		1		
Clostridium butyricum		6.31	0.04		99		
03375540	**Good identification**						**03375540**
					SPO		
Actinomyces israelii		95.34	0.55		1		
Clostridium butyricum		4.66	0.11		99		
03375542	**Low discrimination**						**03375542**
					SPO		
Actinomyces israelii		85.61	0.58		1		
Bifidobacterium adolescentis		13.30	0.37		1		
Clostridium butyricum		1.08	0.04		99		
03375550	**Good identification**						**03375550**
Actinomyces israelii		100.0	0.40	MLZ 13			
03375552	**Low discrimination**						**03375552**
Bifidobacterium adolescentis		55.96	0.39	MNS 87			
Actinomyces israelii		44.04	0.43	RAF 92			
03375562	**Good identification**						**03375562**
Actinomyces israelii		100.0	0.37	NAG 6			
03375602	**Low discrimination**						**03375602**
Actinomyces israelii		77.97	0.35	RHA 7			
Clostridium clostridioforme		22.03	0.16				
03375700	**Acceptable identification**						**03375700**
					SPO		
Actinomyces israelii		91.76	0.42	RHA 7	1		
Clostridium butyricum		8.24	0.03		99		
03375702	**Low discrimination**						**03375702**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		75.45	0.45	RHA 7		
Clostridium clostridioforme		17.85	0.25			
Bifidobacterium adolescentis		4.95	0.19			
03375712	**Low discrimination**					**03375712**
Clostridium clostridioforme		82.43	0.36	MNS 99		
Actinomyces israelii		11.43	0.31	RAF 94		
Bifidobacterium adolescentis		6.14	0.20			
03375732	**Good identification**					**03375732**
Clostridium clostridioforme		96.33	0.36	MNS 99		
Parabacteroides distasonis		3.67	0.18	RAF 94		
03375740	**Good identification**					**03375740**
Actinomyces israelii		100.0	0.35	RHA 7		
03375742	**Good identification**					**03375742**
Actinomyces israelii		100.0	0.38	RHA 7		
03376100	**Good identification**					**03376100**
					SPO	
Actinomyces israelii		96.41	0.42	GAL 99	1	
Clostridium tertium		3.59	0.03		99	
03376102	**Acceptable identification**					**03376102**
					SPO	
Actinomyces israelii		93.07	0.45	GAL 99	1	
Clostridium tertium		6.93	0.11		99	
03376142	**Good identification**					**03376142**
Actinomyces israelii		100.0	0.38	GAL 99		
03376500	**Good identification**					**03376500**
Actinomyces israelii		100.0	0.40	GAL 99		
03376502	**Acceptable identification**					**03376502**
Actinomyces israelii		93.41	0.43	GAL 99		
Bifidobacterium adolescentis		6.59	0.17			
03376542	**Good identification**					**03376542**
Actinomyces israelii		100.0	0.36	GAL 99		
03377000	**Very good identification**					**03377000**
					SPO	
Actinomyces israelii		99.71	0.66		1	
Clostridium tertium		0.15	0.03		99	
Clostridium innocuum		0.14	0.02		99	
03377002	**Very good identification**					**03377002**
					SPO	
Actinomyces israelii		99.61	0.69		1	
Clostridium tertium		0.30	0.11		99	
Clostridium innocuum		0.10	0.02		99	
03377010	**Good identification**					**03377010**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		98.24	0.51	MLZ 13	1		
Clostridium tertium		1.76	0.07		99		
03377012	**Good identification**						**03377012**
					SPO		
Actinomyces israelii		96.54	0.55	MLZ 13	1		
Clostridium tertium		3.46	0.15		99		
03377020	**Acceptable identification**						**03377020**
					SPO		
Actinomyces israelii		93.39	0.45	NAG 6	1		
Clostridium tertium		6.61	0.11		99		
03377022	**Low discrimination**						**03377022**
					SPO		
Actinomyces israelii		87.61	0.48	NAG 6	1		
Clostridium tertium		12.39	0.19		99		
03377040	**Very good identification**						**03377040**
Actinomyces israelii		100.0	0.59				
03377042	**Very good identification**						**03377042**
Actinomyces israelii		100.0	0.62				
03377050	**Good identification**						**03377050**
Actinomyces israelii		100.0	0.44	MLZ 13			
03377052	**Good identification**						**03377052**
Actinomyces israelii		100.0	0.47	MLZ 13			
03377060	**Good identification**						**03377060**
Actinomyces israelii		100.0	0.38	NAG 6			
03377062	**Good identification**						**03377062**
Actinomyces israelii		100.0	0.41	NAG 6			
03377100	**Very good identification**						**03377100**
					COC	SPO	
Actinomyces israelii		95.72	0.76		1	1	
Clostridium tertium		3.57	0.37		1	99	
Atopobium parvulum		0.62	0.25		99	1	
03377101	**Low discrimination**						**03377101**
					SPO		
Actinomyces israelii		88.86	0.42	URE 1	1		
Actinomyces naeslundii		7.83	0.33		1		
Clostridium tertium		3.31	0.03		99		
03377102	**Good identification**						**03377102**
					SPO		
Actinomyces israelii		92.17	0.79		1		
Clostridium tertium		6.87	0.45		99		
Bifidobacterium adolescentis		0.80	0.37		1		
03377103	**Low discrimination**						**03377103**
					SPO		
Actinomyces israelii		88.24	0.45	URE 1	1		
Clostridium tertium		6.57	0.11		99		
Actinomyces naeslundii		5.19	0.33		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03377110	**Low discrimination**						**03377110**
					SPO		
Actinomyces israelii		69.29	0.62	MLZ 13	1		
Clostridium tertium		30.71	0.41		99		
03377112	**Low discrimination**						**03377112**
					SPO		
Actinomyces israelii		51.10	0.65	MLZ 13	1		
Clostridium tertium		45.29	0.50		99		
Bifidobacterium adolescentis		3.61	0.39		1		
03377120	**Low discrimination**						**03377120**
					COC	SPO	
Clostridium tertium		62.75	0.45	MNS 99	1	99	
Actinomyces israelii		35.84	0.55		1	1	
Atopobium parvulum		0.65	0.12		99	1	
03377122	**Low discrimination**						**03377122**
					SPO		
Clostridium tertium		76.94	0.53	MNS 99	99		
Actinomyces israelii		21.98	0.58		1		
Clostridium ramosum		0.71	0.17		92		
03377130	**Good identification**						**03377130**
					SPO		
Clostridium tertium		95.42	0.50	MNS 99	99		
Actinomyces israelii		4.58	0.41		1		
03377132	**Very good identification**						**03377132**
					SPO		
Clostridium tertium		97.66	0.58	MNS 99	99		
Actinomyces israelii		2.34	0.44		1		
03377140	**Very good identification**						**03377140**
					SPO		
Actinomyces israelii		99.90	0.69		1		
Clostridium tertium		0.10	0.03		99		
03377142	**Very good identification**						**03377142**
					SPO		
Actinomyces israelii		99.53	0.72		1		
Bifidobacterium adolescentis		0.27	0.22		1		
Clostridium tertium		0.19	0.11		99		
03377143	**Good identification**						**03377143**
Actinomyces israelii		100.0	0.38	URE 1			
03377150	**Good identification**						**03377150**
					SPO		
Actinomyces israelii		98.86	0.55	MLZ 13	1		
Clostridium tertium		1.14	0.07		99		
03377152	**Very good identification**						**03377152**
					SPO		
Actinomyces israelii		95.65	0.58	MLZ 13	1		
Clostridium tertium		2.20	0.15		99		
Bifidobacterium adolescentis		2.15	0.23		1		
03377160	**Good identification**						**03377160**
					SPO		
Actinomyces israelii		95.65	0.48	NAG 6	1		
Clostridium tertium		4.35	0.11		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03377162	**Acceptable identification**					**03377162**
					SPO	
Actinomyces israelii		91.66	0.51	NAG 6	1	
Clostridium tertium		8.34	0.19		99	
03377200	**Good identification**					**03377200**
Actinomyces israelii		100.0	0.46	RHA 7		
03377202	**Good identification**					**03377202**
Actinomyces israelii		100.0	0.49	RHA 7		
03377212	**Good identification**					**03377212**
Actinomyces israelii		100.0	0.35	RHA 7		
03377240	**Good identification**					**03377240**
Actinomyces israelii		100.0	0.39	RHA 7		
03377242	**Good identification**					**03377242**
Actinomyces israelii		100.0	0.42	RHA 7		
03377300	**Very good identification**					**03377300**
					SPO	
Actinomyces israelii		99.50	0.57	RHA 7	1	
Clostridium tertium		0.50	0.03		99	
03377302	**Very good identification**					**03377302**
					SPO	
Actinomyces israelii		99.01	0.60	RHA 7	1	
Clostridium tertium		0.99	0.11		99	
03377310	**Acceptable identification**					**03377310**
					SPO	
Actinomyces israelii		94.39	0.43	RHA 7	1	
Clostridium tertium		5.61	0.07	MLZ 13	99	
03377312	**Low discrimination**					**03377312**
					SPO	
Actinomyces israelii		89.37	0.46	RHA 7	1	
Clostridium tertium		10.63	0.15	MLZ 13	99	
03377320	**Low discrimination**					**03377320**
					SPO	
Actinomyces israelii		69.09	0.36	RHA 7	1	
Clostridium tertium		16.23	0.11	NAG 6	99	
Clostridium ramosum		14.69	0.09		92	
03377322	**Low discrimination**					**03377322**
					SPO	
Actinomyces israelii		49.10	0.39	RHA 7	1	
Clostridium ramosum		27.83	0.19	NAG 6	92	
Clostridium tertium		23.07	0.19		99	
03377340	**Good identification**					**03377340**
Actinomyces israelii		100.0	0.50	RHA 7		
03377342	**Good identification**					**03377342**
Actinomyces israelii		100.0	0.53	RHA 7		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03377350	**Good identification**					**03377350**
Actinomyces israelii		100.0	0.36	RHA 7		
03377352	**Good identification**					**03377352**
Actinomyces israelii		100.0	0.39	RHA 7		
03377400	**Very good identification**					**03377400**
Actinomyces israelii		100.0	0.64			
03377402	**Very good identification**					**03377402**
Actinomyces israelii		99.72	0.67			
Bifidobacterium adolescentis		0.28	0.17			
03377410	**Good identification**					**03377410**
Actinomyces israelii		100.0	0.50	MLZ 13		
03377412	**Good identification**					**03377412**
Actinomyces israelii		97.72	0.53	MLZ 13		
Bifidobacterium adolescentis		2.28	0.19			
03377420	**Good identification**					**03377420**
Actinomyces israelii		100.0	0.44	NAG 6		
03377422	**Good identification**					**03377422**
Actinomyces israelii		100.0	0.47	NAG 6		
03377440	**Very good identification**					**03377440**
Actinomyces israelii		100.0	0.57			
03377442	**Very good identification**					**03377442**
Actinomyces israelii		100.0	0.60			
03377450	**Good identification**					**03377450**
Actinomyces israelii		100.0	0.43	MLZ 13		
03377452	**Good identification**					**03377452**
Actinomyces israelii		100.0	0.46	MLZ 13		
03377460	**Good identification**					**03377460**
Actinomyces israelii		100.0	0.37	NAG 6		
03377462	**Good identification**					**03377462**
Actinomyces israelii		100.0	0.40	NAG 6		
03377500	**Very good identification**					**03377500**
					SPO	
Actinomyces israelii		99.42	0.74		1	
Clostridium butyricum		0.43	0.13		99	
Bifidobacterium adolescentis		0.11	0.17		1	
03377501	**Good identification**					**03377501**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.40	URE 1		
03377502	**Good identification**					**03377502**
					SPO	
Actinomyces israelii		93.23	0.77		1	
Bifidobacterium adolescentis		6.58	0.51		1	
Clostridium butyricum		0.10	0.06		99	
03377503	**Acceptable identification**					**03377503**
Actinomyces israelii		93.41	0.43	URE 1		
Bifidobacterium adolescentis		6.59	0.17			
03377510	**Very good identification**					**03377510**
					SPO	
Actinomyces israelii		98.60	0.60	MLZ 13	1	
Bifidobacterium adolescentis		0.86	0.19		1	
Clostridium tertium		0.54	0.07		99	
03377512	**Low discrimination**					**03377512**
Actinomyces israelii		62.52	0.63	MLZ 13		
Bifidobacterium adolescentis		36.10	0.53			
Bifidobacterium dentium		0.69	0.06			
03377520	**Good identification**					**03377520**
					SPO	
Actinomyces israelii		97.33	0.54	NAG 6	1	
Clostridium tertium		2.10	0.11		99	
Clostridium butyricum		0.57	-.05		99	
03377522	**Good identification**					**03377522**
					SPO	
Actinomyces israelii		94.84	0.57	NAG 6	1	
Clostridium tertium		4.10	0.19		99	
Bifidobacterium adolescentis		1.06	0.17		1	
03377530	**Low discrimination**					**03377530**
					SPO	
Actinomyces israelii		76.75	0.40	MLZ 13	1	
Clostridium tertium		19.73	0.15	NAG 6	99	
Bifidobacterium dentium		3.52	-.07		1	
03377532	**Low discrimination**					**03377532**
					SPO	
Actinomyces israelii		52.22	0.43	MLZ 13	1	
Clostridium tertium		26.86	0.24	NAG 6	99	
Bifidobacterium dentium		16.15	0.10		1	
03377540	**Very good identification**					**03377540**
					SPO	
Actinomyces israelii		99.90	0.67		1	
Clostridium butyricum		0.10	-.05		99	
03377542	**Very good identification**					**03377542**
Actinomyces israelii		97.81	0.70			
Bifidobacterium adolescentis		2.19	0.36			
03377543	**Good identification**					**03377543**
Actinomyces israelii		100.0	0.36	URE 1		
03377550	**Good identification**					**03377550**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Actinomyces israelii		99.20	0.53	MLZ 13	1	
Blautia producta		0.80	-.03		99	
03377552	**Low discrimination**					**03377552**
Actinomyces israelii		84.50	0.56	MLZ 13		
Bifidobacterium adolescentis		15.50	0.37			
03377560	**Good identification**					**03377560**
Actinomyces israelii		100.0	0.47	NAG 6		
03377562	**Good identification**					**03377562**
Actinomyces israelii		100.0	0.50	NAG 6		
03377572	**Good identification**					**03377572**
Actinomyces israelii		100.0	0.36	MLZ 13		
03377600	**Good identification**					**03377600**
Actinomyces israelii		100.0	0.45	RHA 7		
03377602	**Good identification**					**03377602**
Actinomyces israelii		100.0	0.48	RHA 7		
03377640	**Good identification**					**03377640**
Actinomyces israelii		100.0	0.38	RHA 7		
03377642	**Good identification**					**03377642**
Actinomyces israelii		100.0	0.41	RHA 7		
03377700	**Good identification**					**03377700**
Actinomyces israelii		100.0	0.55	RHA 7		
03377702	**Very good identification**					**03377702**
Actinomyces israelii		99.06	0.58	RHA 7		
Bifidobacterium adolescentis		0.94	0.17			
03377710	**Good identification**					**03377710**
Actinomyces israelii		100.0	0.41	RHA 7		
03377712	**Acceptable identification**					**03377712**
Actinomyces israelii		92.81	0.44	RHA 7		
Bifidobacterium adolescentis		7.19	0.19	MLZ 13		
03377722	**Good identification**					**03377722**
Actinomyces israelii		100.0	0.38	RHA 7		
03377740	**Good identification**					**03377740**
Actinomyces israelii		100.0	0.48	RHA 7		
03377742	**Good identification**					**03377742**
Actinomyces israelii		100.0	0.51	RHA 7		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03377752	**Good identification**							**03377752**
Actinomyces israelii		100.0	0.37	RHA 7				
03400000	**Low discrimination**							**03400000**
					COC	GRA	SPO	
Clostridium botulinum biovar A		33.92	0.42	SUC 7	1	99	99	
Megasphaera elsdenii		16.50	0.41	RAF 1	99	1	1	
Peptostreptococcus anaerobius		15.81	0.32		99	99	1	
03400002	**Low discrimination**							**03400002**
					COC	GRA		
Atopobium minutum		71.15	0.41	SUC 5	1	99		
Megasphaera elsdenii		12.48	0.25	RAF 1	99	1		
Fusobacterium mortiferum		5.48	0.27		1	1		
03400020	**Low discrimination**							**03400020**
					SPO			
Atopobium minutum		59.85	0.36	SUC 5	1			
Clostridium cadaveris		21.21	0.18	RAF 1	99			
Clostridium novyi biovar A		4.95	0.21		99			
03400040	**Low discrimination**							**03400040**
					COC	GRA		
Megasphaera elsdenii		48.66	0.38	RAF 1	99	1		
Clostridium botulinum biovar A		44.84	0.33	FRU 95	1	99		
Clostridium novyi biovar A		5.10	0.29		1	99		
03400100	**Low discrimination**							**03400100**
					GRA			
Fusobacterium mortiferum		64.32	0.50	FRU 99	1			
Actinomyces meyeri		30.47	0.35		99			
Lactobacillus fermentum		1.49	0.24		99			
03401000	**Low discrimination**							**03401000**
					GRA			
Lactobacillus fermentum		47.84	0.54	FRU 99	99			
Fusobacterium mortiferum		21.82	0.47		1			
Atopobium minutum		10.50	0.36		99			
03401002	**Low discrimination**							**03401002**
					GRA			
Lactobacillus fermentum		80.49	0.54	FRU 99	99			
Atopobium minutum		11.77	0.33		99			
Fusobacterium mortiferum		4.08	0.30		1			
03401020	**Low discrimination**							**03401020**
Lactobacillus fermentum		69.76	0.46	FRU 99				
Atopobium minutum		15.31	0.28					
Eubacterium tortuosum		7.06	0.11					
03401022	**Low discrimination**							**03401022**
Lactobacillus fermentum		85.90	0.46	FRU 99				
Atopobium minutum		12.57	0.25					
Eubacterium tortuosum		1.53	-.02					
03401100	**Low discrimination**							**03401100**
					COC	GRA		
Lactobacillus fermentum		42.57	0.58	FRU 99	1	99		
Fusobacterium mortiferum		27.76	0.53		1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		16.32	0.32		99	99		
03401102	**Low discrimination**							**03401102**
					COC	GRA		
Lactobacillus fermentum		88.92	0.58	FRU 99	1	99		
Fusobacterium mortiferum		6.44	0.37		1	1		
Blautia hansenii		3.79	0.15		99	99		
03401122	**Low discrimination**							**03401122**
					COC	GRA		
Lactobacillus fermentum		77.40	0.50	FRU 99	1	99		
Prevotella melaninogenica		21.50	0.21		1	1		
Blautia hansenii		1.10	-.01		99	99		
03401400	**Good identification**							**03401400**
					GRA			
Lactobacillus fermentum		96.52	0.54	FRU 99	99			
Actinomyces meyeri		3.04	0.18		99			
Fusobacterium mortiferum		0.45	0.13		1			
03401402	**Good identification**							**03401402**
Lactobacillus fermentum		100.0	0.54	FRU 99				
03401420	**Good identification**							**03401420**
Lactobacillus fermentum		98.73	0.46	FRU 99				
Actinomyces meyeri		1.27	0.03					
03401422	**Good identification**							**03401422**
Lactobacillus fermentum		100.0	0.46	FRU 99				
03401500	**Low discrimination**							**03401500**
					COC			
Lactobacillus fermentum		86.37	0.58	FRU 99	1			
Actinomyces meyeri		6.67	0.28		1			
Blautia hansenii		5.84	0.19		99			
03401502	**Very good identification**							**03401502**
					COC			
Lactobacillus fermentum		98.41	0.58	FRU 99	1			
Blautia hansenii		0.74	0.03		99			
Bifidobacterium longum subsp. longum		0.62	0.13		1			
03401520	**Low discrimination**							**03401520**
Lactobacillus fermentum		87.38	0.50	FRU 99				
Bifidobacterium longum subsp. longum		6.61	0.23					
Actinomyces meyeri		2.76	0.13					
03401522	**Acceptable identification**							**03401522**
Lactobacillus fermentum		92.97	0.50	FRU 99				
Bifidobacterium longum subsp. longum		7.03	0.23					
03402000	**Acceptable identification**							**03402000**
					COC	SPO	CAT	
Megasphaera elsdenii		92.51	0.41	RAF 1	99	1	1	
Propionibacterium granulosum		3.17	0.21	FRU 95	1	1	90	
Clostridium botulinum biovar A		1.92	0.08		1	99	1	
03402040	**Good identification**							**03402040**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		99.08	0.38	RAF 1	99	1	
Clostridium botulinum biovar A		0.92	-.01	FRU 95	1	99	
03404100	**Acceptable identification**						**03404100**
					GRA		
Actinomyces meyeri		91.18	0.40	RAF 1	99		
Fusobacterium mortiferum		7.46	0.31	FRU 95	1		
Lactobacillus fermentum		1.35	0.20		99		
03405000	**Low discrimination**						**03405000**
					GRA		
Lactobacillus fermentum		67.80	0.50	FRU 99	99		
Actinomyces meyeri		28.24	0.33		99		
Fusobacterium mortiferum		3.96	0.27		1		
03405002	**Good identification**						**03405002**
					GRA		
Lactobacillus fermentum		98.10	0.50	FRU 99	99		
Actinomyces meyeri		1.26	0.07		99		
Fusobacterium mortiferum		0.64	0.11		1		
03405020	**Low discrimination**						**03405020**
Lactobacillus fermentum		85.44	0.42	FRU 99			
Actinomyces meyeri		14.56	0.18				
03405022	**Good identification**						**03405022**
Lactobacillus fermentum		100.0	0.42	FRU 99			
03405100	**Low discrimination**						**03405100**
					COC		
Actinomyces meyeri		46.00	0.43	RAF 1	1		
Lactobacillus fermentum		45.05	0.54	FRU 95	1		
Blautia hansenii		5.19	0.19		99		
03405102	**Acceptable identification**						**03405102**
					COC		
Lactobacillus fermentum		94.91	0.54	FRU 99	1		
Actinomyces meyeri		3.00	0.17		1		
Blautia hansenii		1.22	0.03		99		
03405120	**Low discrimination**						**03405120**
					COC		
Lactobacillus fermentum		67.52	0.46	FRU 99	1		
Actinomyces meyeri		28.20	0.28		1		
Blautia hansenii		2.59	0.03		99		
03405122	**Good identification**						**03405122**
Lactobacillus fermentum		98.72	0.46	FRU 99			
Actinomyces meyeri		1.28	0.02				
03405400	**Acceptable identification**						**03405400**
Lactobacillus fermentum		90.57	0.50	FRU 99			
Actinomyces meyeri		9.43	0.22				
03405402	**Good identification**						**03405402**
Lactobacillus fermentum		100.0	0.50	FRU 99			
03405420	**Good identification**						**03405420**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		95.91	0.42	FRU 99				
Actinomyces meyeri		4.09	0.08					
03405422	**Good identification**							**03405422**
Lactobacillus fermentum		100.0	0.42	FRU 99				
03405500	**Low discrimination**							**03405500**
Lactobacillus fermentum		76.47	0.54	FRU 99				
Actinomyces meyeri		19.52	0.33					
Bifidobacterium longum subsp. longum		2.46	0.21					
03405502	**Good identification**							**03405502**
Lactobacillus fermentum		96.15	0.54	FRU 99				
Bifidobacterium longum subsp. longum		3.09	0.21					
Actinomyces meyeri		0.76	0.07					
03405520	**Low discrimination**							**03405520**
Lactobacillus fermentum		67.09	0.46	FRU 99				
Bifidobacterium longum subsp. longum		25.90	0.31					
Actinomyces meyeri		7.01	0.18					
03405522	**Low discrimination**							**03405522**
Lactobacillus fermentum		72.14	0.46	FRU 99				
Bifidobacterium longum subsp. longum		27.86	0.31					
03405530	**Good identification**							**03405530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87				
03405532	**Good identification**							**03405532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87				
03410000	**Low discrimination**							**03410000**
					COC	GRA	CAT	
Fusobacterium mortiferum		58.72	0.78		1	1	5	
Megasphaera elsdenii		25.67	0.63		99	1	1	
Propionibacterium granulosum		6.07	0.58		1	99	90	
03410001	**Low discrimination**							**03410001**
					COC	GRA	CAT	
Fusobacterium mortiferum		58.19	0.44	URE 1	1	1	5	
Megasphaera elsdenii		25.44	0.29		99	1	1	
Propionibacterium granulosum		6.02	0.24		1	99	90	
03410002	**Low discrimination**							**03410002**
					COC	GRA		
Fusobacterium mortiferum		46.80	0.62	BGL 10	1	1		
Megasphaera elsdenii		20.46	0.47		99	1		
Lactobacillus fermentum		14.00	0.54		1	99		
03410010	**Low discrimination**							**03410010**
					COC	GRA	CAT	
Propionibacterium granulosum		45.87	0.43	MNS 99	1	99	90	
Fusobacterium mortiferum		32.86	0.44	MLZ 12	1	1	5	
Megasphaera elsdenii		14.37	0.29		99	1	1	
03410020	**Low discrimination**							**03410020**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		35.29	0.47	MNS 99	99	90	
Lactobacillus fermentum		15.12	0.46		99	1	
Fusobacterium mortiferum		13.79	0.44		1	5	
03410022	**Low discrimination**						**03410022**
					CAT		
Lactobacillus fermentum		45.37	0.46	GAL 99	1		
Propionibacterium granulosum		26.48	0.37		90		
Atopobium minutum		18.38	0.32		1		
03410040	**Good identification**						**03410040**
					COC	GRA	
Megasphaera elsdenii		93.24	0.60	RAF 1	99	1	
Fusobacterium mortiferum		3.23	0.44		1	1	
Pseudoramibacter alactolyticus		2.24	0.21		1	99	
03410042	**Acceptable identification**						**03410042**
					COC	GRA	
Megasphaera elsdenii		92.48	0.44	RAF 1	99	1	
Fusobacterium mortiferum		3.21	0.27	BGL 10	1	1	
Pseudoramibacter alactolyticus		2.22	0.05		1	99	
03410100	**Low discrimination**						**03410100**
					GRA		
Fusobacterium mortiferum		89.54	0.84		1		
Actinomyces meyeri		8.14	0.57		99		
Lactobacillus fermentum		2.08	0.58		99		
03410101	**Low discrimination**						**03410101**
					GRA		
Fusobacterium mortiferum		89.76	0.50	URE 1	1		
Actinomyces meyeri		8.16	0.23		99		
Lactobacillus fermentum		2.09	0.24		99		
03410102	**Low discrimination**						**03410102**
					GRA		
Fusobacterium mortiferum		80.63	0.68	BGL 10	1		
Lactobacillus fermentum		16.86	0.58		99		
Actinomyces meyeri		2.04	0.31		99		
03410110	**Low discrimination**						**03410110**
					GRA		
Fusobacterium mortiferum		89.27	0.50	MLZ 1	1		
Actinomyces meyeri		8.12	0.23		99		
Lactobacillus fermentum		2.07	0.24		99		
03410120	**Low discrimination**						**03410120**
					GRA		
Actinomyces meyeri		39.50	0.42	RAF 1	99		
Fusobacterium mortiferum		32.19	0.50	NAG 12	1		
Lactobacillus fermentum		24.68	0.50		99		
03410122	**Low discrimination**						**03410122**
					GRA		
Lactobacillus fermentum		81.99	0.50	GAL 99	99		
Fusobacterium mortiferum		11.88	0.34		1		
Actinomyces meyeri		4.06	0.16		99		
03410140	**Low discrimination**						**03410140**
					COC	GRA	
Fusobacterium mortiferum		80.37	0.50	SOR 1	1	1	
Megasphaera elsdenii		10.04	0.26		99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		7.31	0.23		1	99		
03410200	**Low discrimination**							**03410200**
					COC	GRA	CAT	
Fusobacterium mortiferum		52.49	0.44	RHA 1	1	1	5	
Megasphaera elsdenii		22.95	0.29		99	1	1	
Propionibacterium granulosum		16.62	0.32		1	99	90	
03410300	**Low discrimination**							**03410300**
					GRA			
Fusobacterium mortiferum		89.76	0.50	RHA 1	1			
Actinomyces meyeri		8.16	0.23		99			
Lactobacillus fermentum		2.09	0.24		99			
03410400	**Low discrimination**							**03410400**
					GRA			
Lactobacillus fermentum		52.86	0.54	GAL 99	99			
Actinomyces meyeri		21.09	0.36		99			
Fusobacterium mortiferum		16.07	0.44		1			
03410402	**Acceptable identification**							**03410402**
					COC	GRA		
Lactobacillus fermentum		93.21	0.54	GAL 99	1	99		
Fusobacterium mortiferum		3.15	0.27		1	1		
Megasphaera elsdenii		1.38	0.12		99	1		
03410420	**Low discrimination**							**03410420**
					CAT			
Lactobacillus fermentum		84.26	0.46	GAL 99	1			
Actinomyces meyeri		13.75	0.22		1			
Propionibacterium granulosum		1.99	0.13		90			
03410422	**Good identification**							**03410422**
Lactobacillus fermentum		100.0	0.46	GAL 99				
03410500	**Low discrimination**							**03410500**
					GRA			
Lactobacillus fermentum		41.44	0.58	GAL 99	99			
Actinomyces meyeri		40.54	0.47		99			
Fusobacterium mortiferum		18.02	0.50		1			
03410502	**Good identification**							**03410502**
					GRA			
Lactobacillus fermentum		92.72	0.58	GAL 99	99			
Fusobacterium mortiferum		4.48	0.34		1			
Actinomyces meyeri		2.81	0.21		99			
03410520	**Low discrimination**							**03410520**
					GRA			
Lactobacillus fermentum		70.41	0.50	GAL 99	99			
Actinomyces meyeri		28.18	0.32		99			
Fusobacterium mortiferum		0.93	0.16		1			
03410522	**Good identification**							**03410522**
Lactobacillus fermentum		98.11	0.50	GAL 99				
Actinomyces meyeri		1.21	0.06					
Bifidobacterium longum subsp. longum		0.67	0.05					
03411000	**Low discrimination**							**03411000**
					GRA	CAT		
Lactobacillus fermentum		65.76	0.88		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		29.99	0.81		1	5	
Propionibacterium granulosum		2.07	0.58		99	90	
03411001	**Low discrimination**						**03411001**
					GRA	CAT	
Lactobacillus fermentum		65.00	0.54	URE 1	99	1	
Fusobacterium mortiferum		29.64	0.47		1	5	
Propionibacterium granulosum		2.04	0.24		99	90	
03411002	**Good identification**						**03411002**
					GRA	CAT	
Lactobacillus fermentum		94.25	0.88		99	1	
Fusobacterium mortiferum		4.78	0.65		1	5	
Propionibacterium granulosum		0.74	0.47		99	90	
03411003	**Acceptable identification**						**03411003**
					GRA		
Lactobacillus fermentum		92.58	0.54	URE 1	99		
Fusobacterium mortiferum		4.69	0.30		1		
Actinomyces naeslundii		2.00	0.36		99		
03411010	**Low discrimination**						**03411010**
					GRA	CAT	
Lactobacillus fermentum		52.36	0.54	MLZ 1	99	1	
Fusobacterium mortiferum		23.88	0.47		1	5	
Propionibacterium granulosum		22.22	0.43		99	90	
03411012	**Low discrimination**						**03411012**
					GRA	CAT	
Lactobacillus fermentum		86.45	0.54	MLZ 1	99	1	
Propionibacterium granulosum		9.17	0.33		99	90	
Fusobacterium mortiferum		4.38	0.30		1	5	
03411020	**Very good identification**						**03411020**
					GRA	CAT	
Lactobacillus fermentum		95.05	0.80		99	1	
Propionibacterium granulosum		2.24	0.47		99	90	
Fusobacterium mortiferum		1.31	0.47		1	5	
03411021	**Acceptable identification**						**03411021**
					CAT		
Lactobacillus fermentum		90.72	0.46	URE 1	1		
Actinomyces naeslundii		5.89	0.36		25		
Propionibacterium granulosum		2.14	0.13		90		
03411022	**Very good identification**						**03411022**
					GRA	CAT	
Lactobacillus fermentum		99.02	0.80		99	1	
Propionibacterium granulosum		0.58	0.37		99	90	
Fusobacterium mortiferum		0.15	0.30		1	5	
03411023	**Acceptable identification**						**03411023**
Lactobacillus fermentum		93.91	0.46	URE 1			
Actinomyces naeslundii		6.10	0.36				
03411030	**Low discrimination**						**03411030**
					GRA	CAT	
Lactobacillus fermentum		75.07	0.46	MLZ 1	99	1	
Propionibacterium granulosum		23.89	0.33		99	90	
Fusobacterium mortiferum		1.04	0.13		1	5	
03411032	**Acceptable identification**						**03411032**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Lactobacillus fermentum		92.63	0.46	MLZ 1	1		
Propionibacterium granulosum		7.37	0.22		90		
03411040	**Low discrimination**						**03411040**
					COC	GRA	
Lactobacillus fermentum		62.29	0.54	SOR 1	1	99	
Fusobacterium mortiferum		28.41	0.47		1	1	
Megasphaera elsdenii		5.52	0.26		99	1	
03411042	**Acceptable identification**						**03411042**
					COC	GRA	
Lactobacillus fermentum		93.60	0.54	SOR 1	1	99	
Fusobacterium mortiferum		4.74	0.30		1	1	
Megasphaera elsdenii		0.92	0.09		99	1	
03411060	**Good identification**						**03411060**
					GRA	CAT	
Lactobacillus fermentum		96.40	0.46	SOR 1	99	1	
Propionibacterium granulosum		2.27	0.13		99	90	
Fusobacterium mortiferum		1.33	0.13		1	5	
03411062	**Good identification**						**03411062**
Lactobacillus fermentum		100.0	0.46	SOR 1			
03411100	**Low discrimination**						**03411100**
					GRA		
Lactobacillus fermentum		58.40	0.92		99		
Fusobacterium mortiferum		38.08	0.87		1		
Actinomyces meyeri		3.46	0.60		99		
03411101	**Low discrimination**						**03411101**
					GRA		
Lactobacillus fermentum		56.89	0.58	URE 1	99		
Fusobacterium mortiferum		37.10	0.53		1		
Actinomyces meyeri		3.37	0.26		99		
03411102	**Good identification**						**03411102**
					GRA		
Lactobacillus fermentum		93.04	0.92		99		
Fusobacterium mortiferum		6.74	0.71		1		
Actinomyces meyeri		0.17	0.34		99		
03411103	**Good identification**						**03411103**
					GRA		
Lactobacillus fermentum		90.07	0.58	URE 1	99		
Fusobacterium mortiferum		6.53	0.37		1		
Actinomyces naeslundii		3.40	0.43		99		
03411110	**Low discrimination**						**03411110**
					GRA		
Lactobacillus fermentum		58.17	0.58	MLZ 1	99		
Fusobacterium mortiferum		37.93	0.53		1		
Actinomyces meyeri		3.45	0.26		99		
03411112	**Good identification**						**03411112**
					GRA		
Lactobacillus fermentum		93.24	0.58	MLZ 1	99		
Fusobacterium mortiferum		6.76	0.37		1		
03411120	**Low discrimination**						**03411120**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		84.88	0.84		99		
Prevotella melaninogenica		11.26	0.50		1		
Actinomyces meyeri		2.06	0.45		99		
03411121	**Low discrimination**						**03411121**
					GRA		
Lactobacillus fermentum		77.02	0.50	URE 1	99		
Prevotella melaninogenica		10.21	0.15		1		
Actinomyces naeslundii		8.72	0.43		99		
03411122	**Very good identification**						**03411122**
					GRA		
Lactobacillus fermentum		97.06	0.84		99		
Prevotella melaninogenica		2.45	0.37		1		
Fusobacterium mortiferum		0.21	0.37		1		
03411123	**Low discrimination**						**03411123**
					GRA		
Lactobacillus fermentum		87.84	0.50	URE 1	99		
Actinomyces naeslundii		9.95	0.43		99		
Prevotella melaninogenica		2.22	0.03		1		
03411130	**Low discrimination**						**03411130**
					GRA		
Lactobacillus fermentum		84.15	0.50	MLZ 1	99		
Prevotella melaninogenica		11.16	0.15		1		
Actinomyces meyeri		2.04	0.11		99		
03411132	**Good identification**						**03411132**
					GRA		
Lactobacillus fermentum		95.23	0.50	MLZ 1	99		
Prevotella melaninogenica		2.41	0.03		1		
Parabacteroides distasonis		1.25	0.14		1		
03411140	**Low discrimination**						**03411140**
					GRA		
Lactobacillus fermentum		58.20	0.58	SOR 1	99		
Fusobacterium mortiferum		37.96	0.53		1		
Actinomyces meyeri		3.45	0.26		99		
03411142	**Good identification**						**03411142**
					GRA		
Lactobacillus fermentum		92.57	0.58	SOR 1	99		
Fusobacterium mortiferum		6.71	0.37		1		
Actinomyces naeslundii		0.41	0.28		99		
03411160	**Low discrimination**						**03411160**
					GRA		
Lactobacillus fermentum		83.39	0.50	SOR 1	99		
Prevotella melaninogenica		11.06	0.15		1		
Actinomyces meyeri		2.02	0.11		99		
03411162	**Good identification**						**03411162**
					GRA		
Lactobacillus fermentum		96.29	0.50	SOR 1	99		
Prevotella melaninogenica		2.43	0.03		1		
Actinomyces naeslundii		1.28	0.28		99		
03411200	**Low discrimination**						**03411200**
					GRA	CAT	
Lactobacillus fermentum		63.23	0.54	RHA 1	99	1	
Fusobacterium mortiferum		28.83	0.47		1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		6.09	0.32		99	90	
03411202	**Acceptable identification**						**03411202**
					GRA	CAT	
Lactobacillus fermentum		93.05	0.54	RHA 1	99	1	
Fusobacterium mortiferum		4.71	0.30		1	5	
Propionibacterium granulosum		2.24	0.22		99	90	
03411220	**Acceptable identification**						**03411220**
					GRA	CAT	
Lactobacillus fermentum		92.08	0.46	RHA 1	99	1	
Propionibacterium granulosum		6.65	0.22		99	90	
Fusobacterium mortiferum		1.27	0.13		1	5	
03411222	**Good identification**						**03411222**
					CAT		
Lactobacillus fermentum		98.23	0.46	RHA 1	1		
Propionibacterium granulosum		1.77	0.11		90		
03411300	**Low discrimination**						**03411300**
					GRA		
Lactobacillus fermentum		58.43	0.58	RHA 1	99		
Fusobacterium mortiferum		38.11	0.53		1		
Actinomyces meyeri		3.46	0.26		99		
03411302	**Good identification**						**03411302**
					GRA		
Lactobacillus fermentum		93.24	0.58	RHA 1	99		
Fusobacterium mortiferum		6.76	0.37		1		
03411320	**Low discrimination**						**03411320**
					GRA		
Lactobacillus fermentum		84.99	0.50	RHA 1	99		
Prevotella melaninogenica		11.27	0.15		1		
Actinomyces meyeri		2.06	0.11		99		
03411322	**Acceptable identification**						**03411322**
					GRA		
Lactobacillus fermentum		90.29	0.50	RHA 1	99		
Parabacteroides distasonis		6.62	0.27		1		
Prevotella melaninogenica		2.28	0.03		1		
03411400	**Very good identification**						**03411400**
					GRA		
Lactobacillus fermentum		98.91	0.88		99		
Actinomyces meyeri		0.60	0.39		99		
Fusobacterium mortiferum		0.46	0.47		1		
03411401	**Good identification**						**03411401**
					GRA		
Lactobacillus fermentum		98.95	0.54	URE 1	99		
Actinomyces meyeri		0.60	0.05		99		
Fusobacterium mortiferum		0.46	0.13		1		
03411402	**Very good identification**						**03411402**
					GRA		
Lactobacillus fermentum		99.92	0.88		99		
Fusobacterium mortiferum		0.05	0.30		1		
Actinomyces meyeri		0.02	0.14		99		
03411403	**Good identification**						**03411403**
Lactobacillus fermentum		100.0	0.54	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03411410	**Good identification**					**03411410**
					GRA	
Lactobacillus fermentum		98.53	0.54	MLZ 1	99	
Actinomyces meyeri		0.60	0.05		99	
Fusobacterium mortiferum		0.45	0.13		1	
03411412	**Good identification**					**03411412**
Lactobacillus fermentum		100.0	0.54	MLZ 1		
03411420	**Very good identification**					**03411420**
					CAT	
Lactobacillus fermentum		99.70	0.80		1	
Actinomyces meyeri		0.25	0.25		1	
Propionibacterium granulosum		0.02	0.13		90	
03411421	**Good identification**					**03411421**
Lactobacillus fermentum		100.0	0.46	URE 1		
03411422	**Very good identification**					**03411422**
Lactobacillus fermentum		99.99	0.80			
Bifidobacterium longum subsp. longum		0.01	0.05			
03411423	**Good identification**					**03411423**
Lactobacillus fermentum		100.0	0.46	URE 1		
03411430	**Good identification**					**03411430**
Lactobacillus fermentum		98.13	0.46	MLZ 1		
Bifidobacterium longum subsp. longum		1.87	0.09			
03411432	**Good identification**					**03411432**
Lactobacillus fermentum		98.13	0.46	MLZ 1		
Bifidobacterium longum subsp. longum		1.87	0.09			
03411440	**Good identification**					**03411440**
					GRA	
Lactobacillus fermentum		98.95	0.54	SOR 1	99	
Actinomyces meyeri		0.60	0.05		99	
Fusobacterium mortiferum		0.46	0.13		1	
03411442	**Good identification**					**03411442**
Lactobacillus fermentum		100.0	0.54	SOR 1		
03411460	**Good identification**					**03411460**
Lactobacillus fermentum		100.0	0.46	SOR 1		
03411462	**Good identification**					**03411462**
Lactobacillus fermentum		100.0	0.46	SOR 1		
03411500	**Very good identification**					**03411500**
					GRA	
Lactobacillus fermentum		97.85	0.92		99	
Actinomyces meyeri		1.45	0.50		99	
Fusobacterium mortiferum		0.65	0.53		1	
03411501	**Very good identification**					**03411501**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		97.90	0.58	URE 1	99	
Actinomyces meyeri		1.45	0.16		99	
Fusobacterium mortiferum		0.65	0.19		1	
03411502	**Excellent identification**					**03411502**
					GRA	
Lactobacillus fermentum		99.82	0.92		99	
Fusobacterium mortiferum		0.07	0.37		1	
Bifidobacterium longum subsp. longum		0.06	0.29		99	
03411503	**Very good identification**					**03411503**
Lactobacillus fermentum		100.0	0.58	URE 1		
03411510	**Low discrimination**					**03411510**
Lactobacillus fermentum		89.45	0.58	MLZ 1		
Bifidobacterium longum subsp. longum		8.64	0.33			
Actinomyces meyeri		1.33	0.16			
03411512	**Good identification**					**03411512**
Lactobacillus fermentum		90.96	0.58	MLZ 1		
Bifidobacterium longum subsp. longum		8.78	0.33			
Bifidobacterium adolescentis		0.26	0.09			
03411520	**Very good identification**					**03411520**
Lactobacillus fermentum		98.57	0.84			
Bifidobacterium longum subsp. longum		0.68	0.40			
Actinomyces meyeri		0.60	0.35			
03411521	**Good identification**					**03411521**
Lactobacillus fermentum		99.32	0.50	URE 1		
Bifidobacterium longum subsp. longum		0.68	0.05			
03411522	**Very good identification**					**03411522**
Lactobacillus fermentum		99.25	0.84			
Bifidobacterium longum subsp. longum		0.68	0.40			
Actinomyces meyeri		0.02	0.09			
03411523	**Good identification**					**03411523**
Lactobacillus fermentum		99.32	0.50	URE 1		
Bifidobacterium longum subsp. longum		0.68	0.05			
03411530	**Low discrimination**					**03411530**
Bifidobacterium longum subsp. longum		53.67	0.44	MNS 87		
Lactobacillus fermentum		46.33	0.50	MLT 99		
03411532	**Low discrimination**					**03411532**
					GRA	
Bifidobacterium longum subsp. longum		53.47	0.44	MNS 87	99	
Lactobacillus fermentum		46.16	0.50	MLT 99	99	
Parabacteroides distasonis		0.37	0.11		1	
03411540	**Very good identification**					**03411540**
					GRA	
Lactobacillus fermentum		97.90	0.58	SOR 1	99	
Actinomyces meyeri		1.45	0.16		99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		0.65	0.19		1			
03411542	**Very good identification**							**03411542**
Lactobacillus fermentum		100.0	0.58	SOR 1				
03411560	**Good identification**							**03411560**
Lactobacillus fermentum		99.32	0.50	SOR 1				
Bifidobacterium longum subsp. longum		0.68	0.05					
03411562	**Good identification**							**03411562**
Lactobacillus fermentum		99.32	0.50	SOR 1				
Bifidobacterium longum subsp. longum		0.68	0.05					
03411600	**Good identification**							**03411600**
					GRA			
Lactobacillus fermentum		98.95	0.54	RHA 1	99			
Actinomyces meyeri		0.60	0.05		99			
Fusobacterium mortiferum		0.46	0.13		1			
03411602	**Good identification**							**03411602**
Lactobacillus fermentum		100.0	0.54	RHA 1				
03411620	**Good identification**							**03411620**
Lactobacillus fermentum		100.0	0.46	RHA 1				
03411622	**Good identification**							**03411622**
Lactobacillus fermentum		100.0	0.46	RHA 1				
03411700	**Very good identification**							**03411700**
					GRA			
Lactobacillus fermentum		97.90	0.58	RHA 1	99			
Actinomyces meyeri		1.45	0.16		99			
Fusobacterium mortiferum		0.65	0.19		1			
03411702	**Very good identification**							**03411702**
Lactobacillus fermentum		100.0	0.58	RHA 1				
03411720	**Good identification**							**03411720**
					GRA			
Lactobacillus fermentum		97.96	0.50	RHA 1	99			
Bacteroides vulgatus		1.37	-.04		1			
Bifidobacterium longum subsp. longum		0.67	0.05		99			
03411722	**Good identification**							**03411722**
					GRA			
Lactobacillus fermentum		95.07	0.50	RHA 1	99			
Parabacteroides distasonis		4.27	0.24		1			
Bifidobacterium longum subsp. longum		0.65	0.05		99			
03412000	**Low discrimination**							**03412000**
					COC	CAT	GRA	
Megasphaera elsdenii		79.21	0.63	RAF 1	99	1	1	
Propionibacterium granulosum		14.14	0.56		1	90	99	
Pseudoramibacter alactolyticus		2.73	0.27		1	1	99	
03412002	**Low discrimination**							**03412002**
					COC	CAT	GRA	
Megasphaera elsdenii		67.91	0.47	RAF 1	99	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		27.27	0.45	BGL 10	1	90	99	
Pseudoramibacter alactolyticus		2.34	0.11		1	1	99	
03412010	**Low discrimination**							**03412010**
					COC	SPO	CAT	
Propionibacterium granulosum		58.69	0.41	MNS 99	1	1	90	
Megasphaera elsdenii		24.36	0.29	MLZ 12	99	1	1	
Clostridium difficile		16.11	0.19		1	99	1	
03412020	**Low discrimination**							**03412020**
					COC	CAT	GRA	
Propionibacterium granulosum		73.38	0.45	MNS 99	1	90	99	
Megasphaera elsdenii		16.62	0.29		99	1	1	
Pseudoramibacter alactolyticus		10.00	0.14		1	1	99	
03412022	**Low discrimination**							**03412022**
					COC	CAT	GRA	
Propionibacterium granulosum		86.12	0.35	MNS 99	1	90	99	
Megasphaera elsdenii		8.67	0.12		99	1	1	
Pseudoramibacter alactolyticus		5.22	-.02		1	1	99	
03412040	**Very good identification**							**03412040**
					COC	CAT	GRA	
Megasphaera elsdenii		98.42	0.60	RAF 1	99	1	1	
Pseudoramibacter alactolyticus		1.27	0.17		1	1	99	
Propionibacterium granulosum		0.27	0.21		1	90	99	
03412042	**Good identification**							**03412042**
					COC	CAT	GRA	
Megasphaera elsdenii		97.60	0.44	RAF 1	99	1	1	
Pseudoramibacter alactolyticus		1.26	0.00	BGL 10	1	1	99	
Propionibacterium granulosum		0.59	0.11		1	90	99	
03412100	**Low discrimination**							**03412100**
					COC	GRA		
Fusobacterium mortiferum		53.98	0.50	MAN 1	1	1		
Propionibacterium propionicum		27.60	0.46		1	99		
Megasphaera elsdenii		10.11	0.29		99	1		
03412102	**Low discrimination**							**03412102**
					GRA			
Propionibacterium propionicum		82.43	0.49	NIT 86	99			
Fusobacterium mortiferum		11.94	0.34	MLT 99	1			
Lactobacillus fermentum		2.50	0.24		99			
03412140	**Low discrimination**							**03412140**
					COC	GRA		
Propionibacterium propionicum		79.11	0.46	NIT 86	1	99		
Megasphaera elsdenii		19.33	0.26	MLT 99	99	1		
Fusobacterium mortiferum		1.56	0.16		1	1		
03412142	**Good identification**							**03412142**
					COC	GRA		
Propionibacterium propionicum		98.22	0.49	NIT 86	1	99		
Megasphaera elsdenii		1.78	0.09	MLT 99	99	1		
03413000	**Low discrimination**							**03413000**
					GRA	CAT		
Propionibacterium granulosum		58.66	0.56	MNS 99	99	90		
Lactobacillus fermentum		24.99	0.54		99	1		
Fusobacterium mortiferum		11.39	0.47		1	5		
03413001	**Low discrimination**							**03413001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Actinomyces naeslundii		52.22	0.41	MNS 88	25		
Propionibacterium granulosum		29.49	0.21	MLT 94	90		
Lactobacillus fermentum		12.56	0.20		1		
03413002	**Low discrimination**						**03413002**
					GRA	CAT	
Lactobacillus fermentum		59.32	0.54	MAN 1	99	1	
Propionibacterium granulosum		34.82	0.45		99	90	
Fusobacterium mortiferum		3.01	0.30		1	5	
03413003	**Low discrimination**						**03413003**
					CAT		
Actinomyces naeslundii		72.37	0.41	MNS 88	25		
Lactobacillus fermentum		17.41	0.20	MLT 94	1		
Propionibacterium granulosum		10.22	0.11		90		
03413010	**Good identification**						**03413010**
					GRA	CAT	
Propionibacterium granulosum		95.61	0.41	MNS 99	99	90	
Lactobacillus fermentum		3.02	0.20	MLZ 12	99	1	
Fusobacterium mortiferum		1.38	0.13		1	5	
03413020	**Low discrimination**						**03413020**
					CAT		
Propionibacterium granulosum		61.60	0.45	MNS 99	90		
Lactobacillus fermentum		34.98	0.46		1		
Actinomyces naeslundii		2.94	0.38		25		
03413021	**Low discrimination**						**03413021**
Actinomyces naeslundii		87.63	0.41	MNS 88			
Propionibacterium granulosum		12.37	0.11	MLT 94			
03413022	**Low discrimination**						**03413022**
					CAT		
Lactobacillus fermentum		65.61	0.46	MAN 1	1		
Propionibacterium granulosum		28.88	0.35		90		
Actinomyces naeslundii		5.51	0.38		25		
03413023	**Good identification**						**03413023**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03413100	**Low discrimination**						**03413100**
					GRA		
Lactobacillus fermentum		50.07	0.58	MAN 1	99		
Fusobacterium mortiferum		32.66	0.53		1		
Propionibacterium propionicum		11.13	0.46		99		
03413101	**Low discrimination**						**03413101**
Actinomyces naeslundii		63.90	0.48	MNS 88			
Propionibacterium propionicum		21.54	0.30	MLT 94			
Lactobacillus fermentum		8.81	0.24				
03413102	**Low discrimination**						**03413102**
					GRA		
Lactobacillus fermentum		68.57	0.58	MAN 1	99		
Propionibacterium propionicum		22.86	0.49		99		
Fusobacterium mortiferum		4.97	0.37		1		
03413103	**Low discrimination**						**03413103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		60.84	0.48	MNS 88		
Propionibacterium propionicum		30.77	0.33	MLT 94		
Lactobacillus fermentum		8.39	0.24			
03413120	**Low discrimination**					**03413120**
					GRA	
Lactobacillus fermentum		74.59	0.50	MAN 1	99	
Actinomyces naeslundii		10.93	0.45		99	
Prevotella melaninogenica		9.89	0.15		1	
03413121	**Good identification**					**03413121**
Actinomyces naeslundii		95.61	0.48	MNS 88		
Lactobacillus fermentum		4.39	0.15	MLT 94		
03413122	**Low discrimination**					**03413122**
					GRA	
Lactobacillus fermentum		84.61	0.50	MAN 1	99	
Actinomyces naeslundii		12.40	0.45		99	
Prevotella melaninogenica		2.14	0.03		1	
03413123	**Good identification**					**03413123**
Actinomyces naeslundii		95.61	0.48	MNS 88		
Lactobacillus fermentum		4.39	0.15	MLT 94		
03413140	**Low discrimination**					**03413140**
Propionibacterium propionicum		89.78	0.46	NIT 86		
Lactobacillus fermentum		4.08	0.24	MLT 99		
Actinomyces naeslundii		3.48	0.32			
03413142	**Acceptable identification**					**03413142**
Propionibacterium propionicum		94.69	0.49	NIT 86		
Lactobacillus fermentum		2.87	0.24	MLT 99		
Actinomyces naeslundii		2.45	0.32			
03413161	**Good identification**					**03413161**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03413163	**Good identification**					**03413163**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03413400	**Good identification**					**03413400**
					CAT	
Lactobacillus fermentum		96.68	0.54	MAN 1	1	
Propionibacterium granulosum		2.29	0.21		90	
Actinomyces meyeri		0.58	0.05		1	
03413402	**Good identification**					**03413402**
					CAT	
Lactobacillus fermentum		99.41	0.54	MAN 1	1	
Propionibacterium granulosum		0.59	0.11		90	
03413420	**Good identification**					**03413420**
					CAT	
Lactobacillus fermentum		98.25	0.46	MAN 1	1	
Propionibacterium granulosum		1.75	0.11		90	
03413422	**Good identification**					**03413422**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		100.0	0.46	MAN 1		
03413500	**Very good identification**					**03413500**
					GRA	
Lactobacillus fermentum		97.69	0.58	MAN 1	99	
Actinomyces meyeri		1.45	0.16		99	
Fusobacterium mortiferum		0.64	0.19		1	
03413502	**Very good identification**					**03413502**
Lactobacillus fermentum		99.66	0.58	MAN 1		
Propionibacterium propionicum		0.34	0.15			
03413520	**Good identification**					**03413520**
Lactobacillus fermentum		99.32	0.50	MAN 1		
Bifidobacterium longum subsp. longum		0.68	0.05			
03413522	**Good identification**					**03413522**
Lactobacillus fermentum		99.32	0.50	MAN 1		
Bifidobacterium longum subsp. longum		0.68	0.05			
03414000	**Low discrimination**					**03414000**
					GRA	
Actinomyces meyeri		50.26	0.52	RAF 1	99	
Fusobacterium mortiferum		36.75	0.59		1	
Lactobacillus fermentum		9.53	0.50		99	
03414002	**Low discrimination**					**03414002**
					GRA	
Lactobacillus fermentum		61.08	0.50	GAL 99	99	
Fusobacterium mortiferum		26.18	0.42		1	
Actinomyces meyeri		9.97	0.26		99	
03414020	**Low discrimination**					**03414020**
					GRA	
Actinomyces meyeri		65.10	0.37	RAF 1	99	
Lactobacillus fermentum		30.17	0.42	NAG 12	99	
Fusobacterium mortiferum		3.53	0.24		1	
03414022	**Acceptable identification**					**03414022**
Lactobacillus fermentum		93.74	0.42	GAL 99		
Actinomyces meyeri		6.26	0.11			
03414100	**Low discrimination**					**03414100**
					GRA	
Actinomyces meyeri		66.50	0.62	RAF 1	99	
Fusobacterium mortiferum		28.36	0.65		1	
Lactobacillus fermentum		5.14	0.54		99	
03414102	**Low discrimination**					**03414102**
					GRA	
Lactobacillus fermentum		49.68	0.54	GAL 99	99	
Fusobacterium mortiferum		30.45	0.49		1	
Actinomyces meyeri		19.87	0.36		99	
03414120	**Low discrimination**					**03414120**
					GRA	
Actinomyces meyeri		81.93	0.47	RAF 1	99	
Lactobacillus fermentum		15.49	0.46	NAG 12	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		2.59	0.31		1	
03414122	**Low discrimination**					**03414122**
					GRA	
Lactobacillus fermentum		84.59	0.46	GAL 99	99	
Actinomyces meyeri		13.84	0.21		99	
Fusobacterium mortiferum		1.57	0.14		1	
03414400	**Low discrimination**					**03414400**
					GRA	
Actinomyces meyeri		55.93	0.41	RAF 1	99	
Lactobacillus fermentum		42.41	0.50		99	
Fusobacterium mortiferum		1.65	0.24		1	
03414402	**Good identification**					**03414402**
Lactobacillus fermentum		96.08	0.50	GAL 99		
Actinomyces meyeri		3.92	0.16			
03414420	**Low discrimination**					**03414420**
Lactobacillus fermentum		64.96	0.42	GAL 99		
Actinomyces meyeri		35.04	0.27			
03414422	**Good identification**					**03414422**
Lactobacillus fermentum		100.0	0.42	GAL 99		
03414500	**Low discrimination**					**03414500**
					GRA	
Actinomyces meyeri		75.39	0.52	RAF 1	99	
Lactobacillus fermentum		23.32	0.54		99	
Fusobacterium mortiferum		1.30	0.31		1	
03414502	**Acceptable identification**					**03414502**
					GRA	
Lactobacillus fermentum		90.40	0.54	GAL 99	99	
Actinomyces meyeri		9.04	0.26		99	
Fusobacterium mortiferum		0.56	0.14		1	
03414520	**Low discrimination**					**03414520**
Actinomyces meyeri		56.10	0.37	RAF 1		
Lactobacillus fermentum		42.41	0.46	NAG 12		
Bifidobacterium longum subsp. longum		1.49	0.14			
03414522	**Acceptable identification**					**03414522**
Lactobacillus fermentum		92.94	0.46	GAL 99		
Actinomyces meyeri		3.80	0.11			
Bifidobacterium longum subsp. longum		3.26	0.14			
03415000	**Low discrimination**					**03415000**
					GRA	
Lactobacillus fermentum		87.80	0.84		99	
Actinomyces meyeri		7.02	0.55		99	
Fusobacterium mortiferum		5.13	0.62		1	
03415001	**Low discrimination**					**03415001**
					GRA	
Lactobacillus fermentum		87.85	0.50	URE 1	99	
Actinomyces meyeri		7.02	0.21		99	
Fusobacterium mortiferum		5.13	0.27		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03415002	**Very good identification**					**03415002**
					GRA	
Lactobacillus fermentum		99.10	0.84		99	
Fusobacterium mortiferum		0.64	0.45		1	
Actinomyces meyeri		0.25	0.29		99	
03415003	**Good identification**					**03415003**
					GRA	
Lactobacillus fermentum		99.36	0.50	URE 1	99	
Fusobacterium mortiferum		0.65	0.11		1	
03415010	**Low discrimination**					**03415010**
					GRA	
Lactobacillus fermentum		87.29	0.50	MLZ 1	99	
Actinomyces meyeri		6.98	0.21		99	
Fusobacterium mortiferum		5.10	0.27		1	
03415012	**Good identification**					**03415012**
					GRA	
Lactobacillus fermentum		99.36	0.50	MLZ 1	99	
Fusobacterium mortiferum		0.65	0.11		1	
03415020	**Very good identification**					**03415020**
					GRA	
Lactobacillus fermentum		96.63	0.76		99	
Actinomyces meyeri		3.16	0.40		99	
Fusobacterium mortiferum		0.17	0.27		1	
03415021	**Good identification**					**03415021**
Lactobacillus fermentum		96.83	0.42	URE 1		
Actinomyces meyeri		3.17	0.06			
03415022	**Very good identification**					**03415022**
					GRA	
Lactobacillus fermentum		99.88	0.76		99	
Actinomyces meyeri		0.10	0.14		99	
Fusobacterium mortiferum		0.02	0.11		1	
03415023	**Good identification**					**03415023**
Lactobacillus fermentum		100.0	0.42	URE 1		
03415030	**Good identification**					**03415030**
Lactobacillus fermentum		96.83	0.42	MLZ 1		
Actinomyces meyeri		3.17	0.06			
03415032	**Good identification**					**03415032**
Lactobacillus fermentum		100.0	0.42	MLZ 1		
03415040	**Low discrimination**					**03415040**
					SPO	
Lactobacillus fermentum		82.64	0.50	SOR 1	1	
Actinomyces meyeri		6.61	0.21		1	
Clostridium glycolicum		5.92	0.15		99	
03415042	**Good identification**					**03415042**
					GRA	
Lactobacillus fermentum		99.36	0.50	SOR 1	99	
Fusobacterium mortiferum		0.65	0.11		1	
03415060	**Good identification**					**03415060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		96.83	0.42	SOR 1		
Actinomyces meyeri		3.17	0.06			
03415062	**Good identification**					**03415062**
Lactobacillus fermentum		100.0	0.42	SOR 1		
03415100	**Low discrimination**					**03415100**
					GRA	
Lactobacillus fermentum		78.15	0.88		99	
Actinomyces meyeri		15.31	0.65		99	
Fusobacterium mortiferum		6.53	0.68		1	
03415101	**Low discrimination**					**03415101**
					GRA	
Lactobacillus fermentum		78.15	0.54	URE 1	99	
Actinomyces meyeri		15.31	0.31		99	
Fusobacterium mortiferum		6.53	0.34		1	
03415102	**Very good identification**					**03415102**
					GRA	
Lactobacillus fermentum		98.49	0.88		99	
Fusobacterium mortiferum		0.92	0.52		1	
Actinomyces meyeri		0.60	0.39		99	
03415103	**Good identification**					**03415103**
					GRA	
Lactobacillus fermentum		98.49	0.54	URE 1	99	
Fusobacterium mortiferum		0.92	0.17		1	
Actinomyces meyeri		0.60	0.05		99	
03415110	**Low discrimination**					**03415110**
					GRA	
Lactobacillus fermentum		77.57	0.54	MLZ 1	99	
Actinomyces meyeri		15.20	0.31		99	
Fusobacterium mortiferum		6.48	0.34		1	
03415112	**Good identification**					**03415112**
					GRA	
Lactobacillus fermentum		97.54	0.54	MLZ 1	99	
Fusobacterium mortiferum		0.91	0.17		1	
Actinomyces meyeri		0.59	0.05		99	
03415120	**Good identification**					**03415120**
					GRA	
Lactobacillus fermentum		92.14	0.80		99	
Actinomyces meyeri		7.39	0.50		99	
Fusobacterium mortiferum		0.23	0.34		1	
03415121	**Acceptable identification**					**03415121**
Lactobacillus fermentum		92.58	0.46	URE 1		
Actinomyces meyeri		7.42	0.16			
03415122	**Very good identification**					**03415122**
					GRA	
Lactobacillus fermentum		99.21	0.80		99	
Parabacteroides distasonis		0.38	0.35		1	
Actinomyces meyeri		0.25	0.24		99	
03415123	**Good identification**					**03415123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		100.0	0.46	URE 1		
03415130	**Low discrimination**					**03415130**
Lactobacillus fermentum		87.73	0.46	MLZ 1		
Actinomyces meyeri		7.03	0.16			
Bifidobacterium longum subsp. longum		5.24	0.18			
03415132	**Low discrimination**					**03415132**
					GRA	
Lactobacillus fermentum		79.27	0.46	MLZ 1	99	
Parabacteroides distasonis		15.99	0.31		1	
Bifidobacterium longum subsp. longum		4.74	0.18		99	
03415140	**Low discrimination**					**03415140**
					GRA	
Lactobacillus fermentum		78.15	0.54	SOR 1	99	
Actinomyces meyeri		15.31	0.31		99	
Fusobacterium mortiferum		6.53	0.34		1	
03415142	**Good identification**					**03415142**
					GRA	
Lactobacillus fermentum		98.49	0.54	SOR 1	99	
Fusobacterium mortiferum		0.92	0.17		1	
Actinomyces meyeri		0.60	0.05		99	
03415160	**Acceptable identification**					**03415160**
Lactobacillus fermentum		92.58	0.46	SOR 1		
Actinomyces meyeri		7.42	0.16			
03415162	**Good identification**					**03415162**
Lactobacillus fermentum		100.0	0.46	SOR 1		
03415200	**Low discrimination**					**03415200**
					GRA	
Lactobacillus fermentum		87.85	0.50	RHA 1	99	
Actinomyces meyeri		7.02	0.21		99	
Fusobacterium mortiferum		5.13	0.27		1	
03415202	**Good identification**					**03415202**
					GRA	
Lactobacillus fermentum		99.36	0.50	RHA 1	99	
Fusobacterium mortiferum		0.65	0.11		1	
03415220	**Good identification**					**03415220**
Lactobacillus fermentum		96.83	0.42	RHA 1		
Actinomyces meyeri		3.17	0.06			
03415222	**Good identification**					**03415222**
					GRA	
Lactobacillus fermentum		98.18	0.42	RHA 1	99	
Parabacteroides distasonis		1.82	0.09		1	
03415300	**Low discrimination**					**03415300**
					GRA	
Lactobacillus fermentum		78.15	0.54	RHA 1	99	
Actinomyces meyeri		15.31	0.31		99	
Fusobacterium mortiferum		6.53	0.34		1	
03415302	**Good identification**					**03415302**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		98.12	0.54	RHA 1	99	
Fusobacterium mortiferum		0.91	0.17		1	
Actinomyces meyeri		0.60	0.05		99	
03415320	**Low discrimination**					**03415320**
					GRA	
Lactobacillus fermentum		84.81	0.46	RHA 1	99	
Bacteroides vulgatus		7.43	0.06		1	
Actinomyces meyeri		6.80	0.16		99	
03415322	**Low discrimination**					**03415322**
					GRA	
Parabacteroides distasonis		52.91	0.44	MNS 98	1	
Lactobacillus fermentum		47.09	0.46		99	
03415332	**Good identification**					**03415332**
Parabacteroides distasonis		100.0	0.39	MNS 98		
03415400	**Very good identification**					**03415400**
					GRA	
Lactobacillus fermentum		97.98	0.84		99	
Actinomyces meyeri		1.96	0.44		99	
Fusobacterium mortiferum		0.06	0.27		1	
03415401	**Good identification**					**03415401**
Lactobacillus fermentum		98.04	0.50	URE 1		
Actinomyces meyeri		1.96	0.10			
03415402	**Very good identification**					**03415402**
					GRA	
Lactobacillus fermentum		99.93	0.84		99	
Actinomyces meyeri		0.06	0.19		99	
Fusobacterium mortiferum		0.01	0.11		1	
03415403	**Good identification**					**03415403**
Lactobacillus fermentum		100.0	0.50	URE 1		
03415410	**Good identification**					**03415410**
Lactobacillus fermentum		97.27	0.50	MLZ 1		
Actinomyces meyeri		1.94	0.10			
Bifidobacterium longum subsp. longum		0.79	0.07			
03415412	**Good identification**					**03415412**
Lactobacillus fermentum		99.19	0.50	MLZ 1		
Bifidobacterium longum subsp. longum		0.81	0.07			
03415420	**Very good identification**					**03415420**
Lactobacillus fermentum		99.13	0.76			
Actinomyces meyeri		0.81	0.30			
Bifidobacterium longum subsp. longum		0.06	0.14			
03415421	**Good identification**					**03415421**
Lactobacillus fermentum		100.0	0.42	URE 1		
03415422	**Very good identification**					**03415422**
Lactobacillus fermentum		99.92	0.76			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		0.06	0.14			
Actinomyces meyeri		0.03	0.04			
03415423	**Good identification**					**03415423**
Lactobacillus fermentum		100.0	0.42	URE 1		
03415430	**Acceptable identification**					**03415430**
Lactobacillus fermentum		91.12	0.42	MLZ 1		
Bifidobacterium longum subsp. longum		8.89	0.18			
03415432	**Acceptable identification**					**03415432**
Lactobacillus fermentum		91.12	0.42	MLZ 1		
Bifidobacterium longum subsp. longum		8.89	0.18			
03415440	**Good identification**					**03415440**
Lactobacillus fermentum		98.04	0.50	SOR 1		
Actinomyces meyeri		1.96	0.10			
03415442	**Good identification**					**03415442**
Lactobacillus fermentum		100.0	0.50	SOR 1		
03415460	**Good identification**					**03415460**
Lactobacillus fermentum		100.0	0.42	SOR 1		
03415462	**Good identification**					**03415462**
Lactobacillus fermentum		100.0	0.42	SOR 1		
03415500	**Good identification**					**03415500**
Lactobacillus fermentum		94.99	0.88			
Actinomyces meyeri		4.65	0.55			
Bifidobacterium longum subsp. longum		0.28	0.37			
03415501	**Good identification**					**03415501**
Lactobacillus fermentum		95.33	0.54	URE 1		
Actinomyces meyeri		4.67	0.21			
03415502	**Very good identification**					**03415502**
Lactobacillus fermentum		99.52	0.88			
Bifidobacterium longum subsp. longum		0.29	0.37			
Actinomyces meyeri		0.15	0.29			
03415503	**Good identification**					**03415503**
Lactobacillus fermentum		100.0	0.54	URE 1		
03415510	**Low discrimination**					**03415510**
Lactobacillus fermentum		64.85	0.54	MLZ 1		
Bifidobacterium longum subsp. longum		31.98	0.41			
Actinomyces meyeri		3.18	0.21			
03415512	**Low discrimination**					**03415512**
Lactobacillus fermentum		65.53	0.54	MLZ 1		
Bifidobacterium longum subsp. longum		32.31	0.41			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		2.16	0.23			
03415520	**Good identification**					**03415520**
Lactobacillus fermentum		94.66	0.80			
Bifidobacterium longum subsp. longum		3.32	0.48			
Actinomyces meyeri		1.90	0.40			
03415521	**Acceptable identification**					**03415521**
Lactobacillus fermentum		94.77	0.46	URE 1		
Bifidobacterium longum subsp. longum		3.33	0.14			
Actinomyces meyeri		1.90	0.06			
03415522	**Very good identification**					**03415522**
					GRA	
Lactobacillus fermentum		96.34	0.80		99	
Bifidobacterium longum subsp. longum		3.38	0.48		99	
Parabacteroides distasonis		0.22	0.32		1	
03415523	**Good identification**					**03415523**
Lactobacillus fermentum		96.61	0.46	URE 1		
Bifidobacterium longum subsp. longum		3.39	0.14			
03415530	**Low discrimination**					**03415530**
Bifidobacterium longum subsp. longum		85.30	0.52	MNS 87		
Lactobacillus fermentum		14.42	0.46	MLT 99		
Actinomyces meyeri		0.29	0.06			
03415532	**Low discrimination**					**03415532**
					GRA	
Bifidobacterium longum subsp. longum		84.04	0.52	MNS 87	99	
Lactobacillus fermentum		14.20	0.46	MLT 99	99	
Parabacteroides distasonis		1.76	0.27		1	
03415540	**Good identification**					**03415540**
Lactobacillus fermentum		95.33	0.54	SOR 1		
Actinomyces meyeri		4.67	0.21			
03415542	**Good identification**					**03415542**
Lactobacillus fermentum		100.0	0.54	SOR 1		
03415560	**Acceptable identification**					**03415560**
Lactobacillus fermentum		94.77	0.46	SOR 1		
Bifidobacterium longum subsp. longum		3.33	0.14			
Actinomyces meyeri		1.90	0.06			
03415562	**Good identification**					**03415562**
Lactobacillus fermentum		96.61	0.46	SOR 1		
Bifidobacterium longum subsp. longum		3.39	0.14			
03415600	**Good identification**					**03415600**
Lactobacillus fermentum		98.04	0.50	RHA 1		
Actinomyces meyeri		1.96	0.10			
03415602	**Good identification**					**03415602**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		100.0	0.50	RHA 1			
03415620	**Good identification**						**03415620**
					GRA		
Lactobacillus fermentum		98.12	0.42	RHA 1	99		
Bacteroides vulgatus		1.88	-.09		1		
03415622	**Good identification**						**03415622**
Lactobacillus fermentum		100.0	0.42	RHA 1			
03415700	**Acceptable identification**						**03415700**
					GRA		
Lactobacillus fermentum		94.98	0.54	RHA 1	99		
Actinomyces meyeri		4.65	0.21		99		
Bacteroides vulgatus		0.37	-.09		1		
03415702	**Good identification**						**03415702**
Lactobacillus fermentum		100.0	0.54	RHA 1			
03415722	**Low discrimination**						**03415722**
					GRA		
Lactobacillus fermentum		57.62	0.46	RHA 1	99		
Parabacteroides distasonis		39.68	0.40		1		
Bifidobacterium longum subsp. longum		2.02	0.14		99		
03415732	**Low discrimination**						**03415732**
					GRA		
Parabacteroides distasonis		86.12	0.35	MNS 98	1		
Bifidobacterium longum subsp. longum		13.88	0.18		99		
03417000	**Low discrimination**						**03417000**
					GRA		
Lactobacillus fermentum		84.84	0.50	MAN 1	99		
Actinomyces meyeri		6.78	0.21		99		
Fusobacterium mortiferum		4.96	0.27		1		
03417002	**Good identification**						**03417002**
					GRA	CAT	
Lactobacillus fermentum		98.37	0.50	MAN 1	99	1	
Propionibacterium granulosum		0.99	0.11		99	90	
Fusobacterium mortiferum		0.64	0.11		1	5	
03417020	**Acceptable identification**						**03417020**
					CAT		
Lactobacillus fermentum		94.08	0.42	MAN 1	1		
Actinomyces meyeri		3.08	0.06		1		
Propionibacterium granulosum		2.85	0.11		90		
03417022	**Good identification**						**03417022**
Lactobacillus fermentum		100.0	0.42	MAN 1			
03417100	**Low discrimination**						**03417100**
					GRA		
Lactobacillus fermentum		77.92	0.54	MAN 1	99		
Actinomyces meyeri		15.27	0.31		99		
Fusobacterium mortiferum		6.51	0.34		1		
03417102	**Good identification**						**03417102**
					GRA		
Lactobacillus fermentum		97.94	0.54	MAN 1	99		
Fusobacterium mortiferum		0.91	0.17		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		0.59	0.05		99		
03417120	**Acceptable identification**						**03417120**
Lactobacillus fermentum		92.58	0.46	MAN 1			
Actinomyces meyeri		7.42	0.16				
03417122	**Good identification**						**03417122**
Lactobacillus fermentum		100.0	0.46	MAN 1			
03417400	**Good identification**						**03417400**
Lactobacillus fermentum		98.04	0.50	MAN 1			
Actinomyces meyeri		1.96	0.10				
03417402	**Good identification**						**03417402**
Lactobacillus fermentum		100.0	0.50	MAN 1			
03417420	**Good identification**						**03417420**
Lactobacillus fermentum		100.0	0.42	MAN 1			
03417422	**Good identification**						**03417422**
Lactobacillus fermentum		100.0	0.42	MAN 1			
03417500	**Good identification**						**03417500**
Lactobacillus fermentum		95.33	0.54	MAN 1			
Actinomyces meyeri		4.67	0.21				
03417502	**Good identification**						**03417502**
Lactobacillus fermentum		97.84	0.54	MAN 1			
Bifidobacterium adolescentis		2.16	0.20				
03417520	**Acceptable identification**						**03417520**
Lactobacillus fermentum		94.77	0.46	MAN 1			
Bifidobacterium longum subsp. longum		3.33	0.14				
Actinomyces meyeri		1.90	0.06				
03417522	**Good identification**						**03417522**
Lactobacillus fermentum		96.61	0.46	MAN 1			
Bifidobacterium longum subsp. longum		3.39	0.14				
03420100	**Good identification**						**03420100**
Fusobacterium mortiferum		100.0	0.41	FRU 99			
03421000	**Low discrimination**						**03421000**
					GRA	CAT	
Fusobacterium mortiferum		56.61	0.38	FRU 99	1	5	
Propionibacterium granulosum		39.19	0.32		99	90	
Lactobacillus fermentum		4.20	0.20		99	1	
03421100	**Acceptable identification**						**03421100**
					COC	GRA	
Fusobacterium mortiferum		90.83	0.44	FRU 99	1	1	
Lactobacillus fermentum		4.71	0.24		1	99	
Blautia hansenii		1.81	-.03		99	99	
03430000	**Low discrimination**						**03430000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	CAT	
Propionibacterium granulosum		50.39	0.66	MNS 99	1	99	90	
Fusobacterium mortiferum		48.52	0.69		1	1	5	
Megasphaera elsdenii		0.72	0.29		99	1	1	
03430002	**Low discrimination**							**03430002**
					COC	GRA	CAT	
Propionibacterium granulosum		69.43	0.56	MNS 99	1	99	90	
Fusobacterium mortiferum		29.71	0.53		1	1	5	
Megasphaera elsdenii		0.44	0.12		99	1	1	
03430010	**Acceptable identification**							**03430010**
					GRA	CAT		
Propionibacterium granulosum		93.34	0.51	MNS 99	99	90		
Fusobacterium mortiferum		6.66	0.35	MLZ 12	1	5		
03430012	**Good identification**							**03430012**
					GRA	CAT		
Propionibacterium granulosum		96.93	0.41	MNS 99	99	90		
Fusobacterium mortiferum		3.07	0.18	MLZ 12	1	5		
03430020	**Good identification**							**03430020**
					GRA	SPO	CAT	
Propionibacterium granulosum		96.09	0.56	MNS 99	99	1	90	
Fusobacterium mortiferum		3.74	0.35		1	1	5	
Clostridium cadaveris		0.17	-.04		99	99	1	
03430022	**Good identification**							**03430022**
					GRA	CAT		
Propionibacterium granulosum		98.30	0.45	MNS 99	99	90		
Fusobacterium mortiferum		1.70	0.18		1	5		
03430030	**Good identification**							**03430030**
Propionibacterium granulosum		100.0	0.41	MNS 99				
03430100	**Very good identification**							**03430100**
					GRA	CAT		
Fusobacterium mortiferum		99.17	0.75		1	5		
Propionibacterium granulosum		0.45	0.32		99	90		
Actinomyces meyeri		0.31	0.23		99	1		
03430101	**Good identification**							**03430101**
					GRA			
Fusobacterium mortiferum		97.70	0.41	URE 1	1			
Actinomyces naeslundii		2.30	0.25		99			
03430102	**Very good identification**							**03430102**
					GRA	CAT		
Fusobacterium mortiferum		98.12	0.59	BGL 10	1	5		
Propionibacterium granulosum		0.99	0.21		99	90		
Lactobacillus fermentum		0.69	0.24		99	1		
03430110	**Acceptable identification**							**03430110**
					GRA	CAT		
Fusobacterium mortiferum		94.28	0.41	MLZ 1	1	5		
Propionibacterium granulosum		5.72	0.17		99	90		
03430120	**Low discrimination**							**03430120**
					GRA	CAT		
Fusobacterium mortiferum		84.84	0.41	NAG 1	1	5		
Propionibacterium granulosum		9.44	0.21		99	90		
Actinomyces meyeri		3.52	0.08		99	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
03430140	**Good identification**						**03430140**
Fusobacterium mortiferum		100.0	0.41	SOR 1			
03430200	**Low discrimination**						**03430200**
					GRA	CAT	
Propionibacterium granulosum		76.08	0.40	MNS 99	99	90	
Fusobacterium mortiferum		23.92	0.35	RHA 3	1	5	
03430300	**Good identification**						**03430300**
Fusobacterium mortiferum		100.0	0.41	RHA 1			
03430500	**Low discrimination**						**03430500**
					GRA		
Fusobacterium mortiferum		86.67	0.41	ARA 1	1		
Lactobacillus fermentum		6.74	0.24		99		
Actinomyces meyeri		6.59	0.13		99		
03431000	**Low discrimination**						**03431000**
					GRA	CAT	
Fusobacterium mortiferum		56.47	0.72		1	5	
Propionibacterium granulosum		39.10	0.66		99	90	
Lactobacillus fermentum		4.19	0.54		99	1	
03431001	**Low discrimination**						**03431001**
					GRA	CAT	
Fusobacterium mortiferum		48.20	0.38	URE 1	1	5	
Propionibacterium granulosum		33.37	0.32		99	90	
Actinomyces naeslundii		14.86	0.41		99	25	
03431002	**Low discrimination**						**03431002**
					GRA	CAT	
Propionibacterium granulosum		48.02	0.56	MNS 99	99	90	
Fusobacterium mortiferum		30.83	0.56		1	5	
Lactobacillus fermentum		20.57	0.54		99	1	
03431003	**Low discrimination**						**03431003**
					GRA	CAT	
Actinomyces naeslundii		46.25	0.41	MNS 88	99	25	
Propionibacterium granulosum		25.97	0.21	MLT 94	99	90	
Fusobacterium mortiferum		16.67	0.21		1	5	
03431010	**Low discrimination**						**03431010**
					GRA	CAT	
Propionibacterium granulosum		89.69	0.51	MNS 99	99	90	
Fusobacterium mortiferum		9.60	0.38	MLZ 12	1	5	
Lactobacillus fermentum		0.71	0.20		99	1	
03431012	**Acceptable identification**						**03431012**
					GRA	CAT	
Propionibacterium granulosum		92.65	0.41	MNS 99	99	90	
Fusobacterium mortiferum		4.41	0.21	MLZ 12	1	5	
Lactobacillus fermentum		2.94	0.20		99	1	
03431020	**Low discrimination**						**03431020**
					GRA	CAT	
Propionibacterium granulosum		82.24	0.56	MNS 99	99	90	
Lactobacillus fermentum		11.74	0.46		99	1	
Fusobacterium mortiferum		4.80	0.38		1	5	
03431021	**Low discrimination**						**03431021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		64.04	0.41	MNS 88			
Propionibacterium granulosum		35.96	0.21	MLT 94			
03431022	**Low discrimination**						**03431022**
					CAT		
Propionibacterium granulosum		60.79	0.45	MNS 99	90		
Lactobacillus fermentum		34.72	0.46		1		
Actinomyces naeslundii		2.92	0.38		25		
03431023	**Low discrimination**						**03431023**
Actinomyces naeslundii		87.69	0.41	MNS 88			
Propionibacterium granulosum		12.31	0.11	MLT 94			
03431030	**Good identification**						**03431030**
Propionibacterium granulosum		100.0	0.41	MNS 99			
03431040	**Low discrimination**						**03431040**
					GRA	CAT	
Fusobacterium mortiferum		55.47	0.38	SOR 1	1	5	
Propionibacterium granulosum		38.41	0.32		99	90	
Lactobacillus fermentum		4.11	0.20		99	1	
03431100	**Good identification**						**03431100**
					GRA		
Fusobacterium mortiferum		94.28	0.78		1		
Lactobacillus fermentum		4.89	0.58		99		
Actinomyces meyeri		0.29	0.26		99		
03431101	**Low discrimination**						**03431101**
					GRA		
Fusobacterium mortiferum		70.03	0.44	URE 1	1		
Actinomyces naeslundii		26.34	0.48		99		
Lactobacillus fermentum		3.63	0.24		99		
03431102	**Low discrimination**						**03431102**
					GRA		
Fusobacterium mortiferum		66.78	0.62	BGL 10	1		
Lactobacillus fermentum		31.16	0.58		99		
Actinomyces naeslundii		1.52	0.45		99		
03431103	**Low discrimination**						**03431103**
					GRA		
Actinomyces naeslundii		69.77	0.48	MNS 88	99		
Fusobacterium mortiferum		20.61	0.28	MLT 94	1		
Lactobacillus fermentum		9.62	0.24		99		
03431110	**Low discrimination**						**03431110**
					GRA	CAT	
Fusobacterium mortiferum		89.04	0.44	MLZ 1	1	5	
Lactobacillus fermentum		4.62	0.24		99	1	
Propionibacterium granulosum		3.60	0.17		99	90	
03431120	**Low discrimination**						**03431120**
					GRA		
Lactobacillus fermentum		49.48	0.50	TRE 1	99		
Fusobacterium mortiferum		28.92	0.44		1		
Actinomyces naeslundii		7.25	0.45		99		
03431121	**Good identification**						**03431121**
Actinomyces naeslundii		95.61	0.48	MNS 88			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		4.39	0.15	MLT 94			
03431122	**Low discrimination**						**03431122**
					GRA		
Lactobacillus fermentum		71.37	0.50	TRE 1	99		
Actinomyces naeslundii		10.46	0.45		99		
Parabacteroides distasonis		7.95	0.30		1		
03431123	**Good identification**						**03431123**
Actinomyces naeslundii		95.61	0.48	MNS 88			
Lactobacillus fermentum		4.39	0.15	MLT 94			
03431140	**Acceptable identification**						**03431140**
					GRA		
Fusobacterium mortiferum		91.23	0.44	SOR 1	1		
Lactobacillus fermentum		4.73	0.24		99		
Actinomyces naeslundii		4.04	0.32		99		
03431141	**Good identification**						**03431141**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03431143	**Good identification**						**03431143**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03431161	**Good identification**						**03431161**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03431163	**Good identification**						**03431163**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03431200	**Low discrimination**						**03431200**
					GRA	CAT	
Propionibacterium granulosum		66.37	0.40	MNS 99	99	90	
Fusobacterium mortiferum		31.31	0.38	RHA 3	1	5	
Lactobacillus fermentum		2.32	0.20		99	1	
03431300	**Good identification**						**03431300**
					GRA		
Fusobacterium mortiferum		95.07	0.44	RHA 1	1		
Lactobacillus fermentum		4.93	0.24		99		
03431322	**Good identification**						**03431322**
					GRA		
Parabacteroides distasonis		97.07	0.38	MNS 98	1		
Lactobacillus fermentum		2.93	0.15	XYL 90	99		
03431400	**Low discrimination**						**03431400**
					GRA	CAT	
Lactobacillus fermentum		80.87	0.54	TRE 1	99	1	
Fusobacterium mortiferum		11.02	0.38		1	5	
Propionibacterium granulosum		7.63	0.32		99	90	
03431402	**Good identification**						**03431402**
					GRA	CAT	
Lactobacillus fermentum		96.27	0.54	TRE 1	99	1	
Propionibacterium granulosum		2.27	0.21		99	90	
Fusobacterium mortiferum		1.46	0.21		1	5	
03431420	**Acceptable identification**						**03431420**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		93.39	0.46	TRE 1	1			
Propionibacterium granulosum		6.61	0.21		90			
03431422	**Good identification**							**03431422**
					CAT			
Lactobacillus fermentum		98.26	0.46	TRE 1	1			
Propionibacterium granulosum		1.74	0.11		90			
03431500	**Low discrimination**							**03431500**
					GRA			
Lactobacillus fermentum		81.71	0.58	TRE 1	99			
Fusobacterium mortiferum		15.92	0.44		1			
Actinomyces meyeri		1.21	0.16		99			
03431502	**Very good identification**							**03431502**
					GRA			
Lactobacillus fermentum		96.54	0.58	TRE 1	99			
Fusobacterium mortiferum		2.09	0.28		1			
Bifidobacterium longum subsp. longum		1.37	0.19		99			
03431520	**Low discrimination**							**03431520**
Lactobacillus fermentum		85.47	0.50	TRE 1				
Bifidobacterium longum subsp. longum		14.54	0.29					
03431522	**Low discrimination**							**03431522**
					GRA			
Lactobacillus fermentum		80.76	0.50	TRE 1	99			
Bifidobacterium longum subsp. longum		13.73	0.29		99			
Parabacteroides distasonis		5.51	0.27		1			
03432000	**Very good identification**							**03432000**
					COC	GRA	CAT	
Propionibacterium granulosum		96.86	0.64	MNS 99	1	99	90	
Megasphaera elsdenii		1.83	0.29		99	1	1	
Fusobacterium mortiferum		1.25	0.35		1	1	5	
03432002	**Good identification**							**03432002**
					COC	GRA	CAT	
Propionibacterium granulosum		98.36	0.53	MNS 99	1	99	90	
Megasphaera elsdenii		0.83	0.12		99	1	1	
Fusobacterium mortiferum		0.56	0.18		1	1	5	
03432010	**Good identification**							**03432010**
					SPO	CAT		
Propionibacterium granulosum		99.32	0.49	MNS 99	1	90		
Clostridium difficile		0.68	-.01	MLZ 12	99	1		
03432012	**Good identification**							**03432012**
Propionibacterium granulosum		100.0	0.39	MNS 99				
03432020	**Good identification**							**03432020**
Propionibacterium granulosum		100.0	0.53	MNS 99				
03432022	**Good identification**							**03432022**
Propionibacterium granulosum		100.0	0.43	MNS 99				
03432030	**Good identification**							**03432030**
Propionibacterium granulosum		100.0	0.39	MNS 99				
03432100	**Low discrimination**							**03432100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Propionibacterium propionicum		50.62	0.45	NIT 86	99	1	
Fusobacterium mortiferum		36.14	0.41	MLT 99	1	5	
Propionibacterium granulosum		12.14	0.30		99	90	
03432102	**Acceptable identification**						**03432102**
					GRA	CAT	
Propionibacterium propionicum		90.31	0.48	NIT 86	99	1	
Fusobacterium mortiferum		4.78	0.25	MLT 99	1	5	
Propionibacterium granulosum		3.61	0.19		99	90	
03432140	**Good identification**						**03432140**
Propionibacterium propionicum		100.0	0.45	NIT 86			
03432142	**Good identification**						**03432142**
Propionibacterium propionicum		100.0	0.48	NIT 86			
03432200	**Good identification**						**03432200**
Propionibacterium granulosum		100.0	0.38	MNS 99			
03433000	**Very good identification**						**03433000**
					GRA	CAT	
Propionibacterium granulosum		97.23	0.64	MNS 99	99	90	
Fusobacterium mortiferum		1.88	0.38		1	5	
Actinomyces naeslundii		0.75	0.42		99	25	
03433001	**Low discrimination**						**03433001**
Actinomyces naeslundii		53.40	0.46	MNS 88			
Propionibacterium granulosum		46.60	0.30	MLT 94			
03433002	**Good identification**						**03433002**
					GRA	CAT	
Propionibacterium granulosum		95.44	0.53	MNS 99	99	90	
Actinomyces naeslundii		2.95	0.42		99	25	
Fusobacterium mortiferum		0.82	0.21		1	5	
03433003	**Low discrimination**						**03433003**
Actinomyces naeslundii		82.09	0.46	MNS 88			
Propionibacterium granulosum		17.91	0.19	MLT 94			
03433010	**Good identification**						**03433010**
Propionibacterium granulosum		100.0	0.49	MNS 99			
03433012	**Good identification**						**03433012**
Propionibacterium granulosum		100.0	0.39	MNS 99			
03433020	**Good identification**						**03433020**
Propionibacterium granulosum		97.01	0.53	MNS 99			
Actinomyces naeslundii		2.99	0.42				
03433021	**Low discrimination**						**03433021**
Actinomyces naeslundii		82.09	0.46	MNS 88			
Propionibacterium granulosum		17.91	0.19	MLT 94			
03433022	**Low discrimination**						**03433022**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Propionibacterium granulosum	89.01	0.43	MNS 99		
	Actinomyces naeslundii	10.99	0.42			
03433023	**Acceptable identification**					**03433023**
	Actinomyces naeslundii	94.83	0.46	MNS 88		
	Propionibacterium granulosum	5.17	0.09	MLT 94		
03433030	**Good identification**					**03433030**
	Propionibacterium granulosum	100.0	0.39	MNS 99		
03433100	**Low discrimination**					**03433100**
					GRA	
	Fusobacterium mortiferum	37.09	0.44	MAN 1	1	
	Propionibacterium propionicum	34.63	0.45		99	
	Actinomyces naeslundii	18.05	0.50		99	
03433101	**Low discrimination**					**03433101**
	Actinomyces naeslundii	87.56	0.53	MNS 88		
	Propionibacterium propionicum	12.44	0.29	MLT 94		
03433102	**Low discrimination**					**03433102**
					GRA	
	Propionibacterium propionicum	66.50	0.48	NIT 86	99	
	Actinomyces naeslundii	23.11	0.50	MLT 99	99	
	Fusobacterium mortiferum	5.28	0.28		1	
03433103	**Low discrimination**					**03433103**
	Actinomyces naeslundii	82.43	0.53	MNS 88		
	Propionibacterium propionicum	17.57	0.32	MLT 94		
03433120	**Low discrimination**					**03433120**
					CAT	
	Actinomyces naeslundii	86.92	0.50	MNS 88	25	
	Propionibacterium granulosum	10.00	0.19	MLT 94	90	
	Lactobacillus fermentum	3.09	0.15		1	
03433121	**Good identification**					**03433121**
	Actinomyces naeslundii	100.0	0.53	MNS 88		
03433122	**Acceptable identification**					**03433122**
	Actinomyces naeslundii	91.47	0.50	MNS 88		
	Lactobacillus fermentum	3.25	0.15	MLT 94		
	Propionibacterium propionicum	2.66	0.14			
03433123	**Good identification**					**03433123**
	Actinomyces naeslundii	100.0	0.53	MNS 88		
03433140	**Acceptable identification**					**03433140**
	Propionibacterium propionicum	91.58	0.45	NIT 86		
	Actinomyces naeslundii	8.42	0.37	MLT 99		
03433141	**Low discrimination**					**03433141**
	Actinomyces naeslundii	55.39	0.40	MNS 88		
	Propionibacterium propionicum	44.61	0.29	MLT 94		
03433142	**Acceptable identification**					**03433142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		94.22	0.48	NIT 86			
Actinomyces naeslundii		5.78	0.37	MLT 99			
03433160	**Good identification**						**03433160**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03433161	**Good identification**						**03433161**
Actinomyces naeslundii		100.0	0.40	MNS 88			
03433162	**Low discrimination**						**03433162**
Actinomyces naeslundii		85.86	0.37	MNS 88			
Propionibacterium propionicum		14.14	0.14	MLT 94			
03433163	**Good identification**						**03433163**
Actinomyces naeslundii		100.0	0.40	MNS 88			
03433200	**Good identification**						**03433200**
Propionibacterium granulosum		100.0	0.38	MNS 99			
03434000	**Low discrimination**						**03434000**
					GRA	CAT	
Fusobacterium mortiferum		83.73	0.50	XYL 7	1	5	
Propionibacterium granulosum		11.67	0.32		99	90	
Actinomyces meyeri		3.87	0.18		99	1	
03434100	**Acceptable identification**						**03434100**
					GRA		
Fusobacterium mortiferum		92.13	0.56	XYL 7	1		
Actinomyces meyeri		7.30	0.28		99		
Lactobacillus fermentum		0.57	0.20		99		
03434102	**Acceptable identification**						**03434102**
					GRA		
Fusobacterium mortiferum		92.83	0.40	XYL 7	1		
Lactobacillus fermentum		5.12	0.20	BGL 10	99		
Actinomyces meyeri		2.05	0.02		99		
03435000	**Low discrimination**						**03435000**
					GRA	CAT	
Fusobacterium mortiferum		58.21	0.53	XYL 7	1	5	
Lactobacillus fermentum		33.68	0.50		99	1	
Propionibacterium granulosum		5.41	0.32		99	90	
03435002	**Low discrimination**						**03435002**
					GRA	CAT	
Lactobacillus fermentum		81.16	0.50	TRE 1	99	1	
Fusobacterium mortiferum		15.58	0.36		1	5	
Propionibacterium granulosum		3.26	0.21		99	90	
03435020	**Low discrimination**						**03435020**
					GRA	CAT	
Lactobacillus fermentum		82.95	0.42	TRE 1	99	1	
Propionibacterium granulosum		9.99	0.21		99	90	
Fusobacterium mortiferum		4.34	0.19		1	5	
03435022	**Acceptable identification**						**03435022**
					GRA	CAT	
Lactobacillus fermentum		94.50	0.42	TRE 1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		2.85	0.11		99	90	
Parabacteroides distasonis		2.66	0.12		1	76	
03435100	**Low discrimination**						**03435100**
					GRA		
Fusobacterium mortiferum		67.32	0.59	XYL 7	1		
Lactobacillus fermentum		27.24	0.54		99		
Actinomyces meyeri		5.34	0.31		99		
03435102	**Low discrimination**						**03435102**
					GRA		
Lactobacillus fermentum		77.74	0.54	TRE 1	99		
Fusobacterium mortiferum		21.35	0.43		1		
Actinomyces meyeri		0.47	0.05		99		
03435120	**Low discrimination**						**03435120**
					GRA		
Lactobacillus fermentum		85.30	0.46	TRE 1	99		
Actinomyces meyeri		6.84	0.16		99		
Fusobacterium mortiferum		6.39	0.25		1		
03435122	**Low discrimination**						**03435122**
					GRA		
Parabacteroides distasonis		63.05	0.47	MNS 98	1		
Lactobacillus fermentum		36.95	0.46		99		
03435132	**Good identification**						**03435132**
					GRA		
Parabacteroides distasonis		98.40	0.42	MNS 98	1		
Bifidobacterium longum subsp. longum		1.60	0.07		99		
03435322	**Good identification**						**03435322**
Parabacteroides distasonis		100.0	0.55	MNS 98			
03435332	**Good identification**						**03435332**
Parabacteroides distasonis		100.0	0.50	MNS 98			
03435400	**Good identification**						**03435400**
					GRA		
Lactobacillus fermentum		96.39	0.50	TRE 1	99		
Actinomyces meyeri		1.93	0.10		99		
Fusobacterium mortiferum		1.68	0.19		1		
03435402	**Good identification**						**03435402**
Lactobacillus fermentum		100.0	0.50	TRE 1			
03435420	**Good identification**						**03435420**
Lactobacillus fermentum		100.0	0.42	TRE 1			
03435422	**Good identification**						**03435422**
Lactobacillus fermentum		100.0	0.42	TRE 1			
03435500	**Low discrimination**						**03435500**
Lactobacillus fermentum		87.23	0.54	TRE 1			
Bifidobacterium longum subsp. longum		6.32	0.27				
Actinomyces meyeri		4.27	0.21				
03435502	**Acceptable identification**						**03435502**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		91.71	0.54	TRE 1				
Bifidobacterium longum subsp. longum		6.64	0.27					
Bifidobacterium adolescentis		1.65	0.18					
03435520	**Low discrimination**							**03435520**
Lactobacillus fermentum		52.94	0.46	TRE 1				
Bifidobacterium longum subsp. longum		46.00	0.37					
Actinomyces meyeri		1.06	0.06					
03435522	**Low discrimination**							**03435522**
					GRA			
Parabacteroides distasonis		35.88	0.43	MNS 98	1			
Lactobacillus fermentum		34.31	0.46		99			
Bifidobacterium longum subsp. longum		29.81	0.37		99			
03435530	**Good identification**							**03435530**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87				
03435532	**Low discrimination**							**03435532**
					GRA			
Bifidobacterium longum subsp. longum		72.43	0.41	MNS 87	99			
Parabacteroides distasonis		27.57	0.38	MLT 99	1			
03435722	**Good identification**							**03435722**
Parabacteroides distasonis		100.0	0.51	MNS 98				
03435732	**Good identification**							**03435732**
					GRA			
Parabacteroides distasonis		99.12	0.47	MNS 98	1			
Bifidobacterium longum subsp. longum		0.88	0.07		99			
03440000	**Low discrimination**							**03440000**
					COC	GRA	SPO	
Fusobacterium mortiferum		74.47	0.36	FRU 99	1	1	1	
Clostridium botulinum biovar A		10.57	0.08		1	99	99	
Megasphaera elsdenii		5.14	0.07		99	1	1	
03440100	**Good identification**							**03440100**
Fusobacterium mortiferum		100.0	0.42	FRU 99				
03441000	**Low discrimination**							**03441000**
					GRA			
Fusobacterium mortiferum		82.91	0.39	FRU 99	1			
Eubacterium tortuosum		11.58	0.07		99			
Lactobacillus fermentum		5.51	0.20		99			
03441100	**Acceptable identification**							**03441100**
					COC	GRA		
Fusobacterium mortiferum		92.71	0.45	FRU 99	1	1		
Lactobacillus fermentum		4.31	0.24		1	99		
Blautia hansenii		1.65	-.03		99	99		
03450000	**Very good identification**							**03450000**
					COC	GRA	CAT	
Fusobacterium mortiferum		97.95	0.70		1	1	5	
Megasphaera elsdenii		1.30	0.29		99	1	1	
Propionibacterium granulosum		0.31	0.24		1	99	90	
03450001	**Good identification**							**03450001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.36	URE 1			
03450002	**Good identification**						**03450002**
					COC	GRA	
Fusobacterium mortiferum		96.80	0.53	BGL 10	1	1	
Megasphaera elsdenii		1.28	0.12		99	1	
Lactobacillus fermentum		0.88	0.20		1	99	
03450010	**Good identification**						**03450010**
					GRA	CAT	
Fusobacterium mortiferum		95.94	0.36	MLZ 1	1	5	
Propionibacterium granulosum		4.06	0.09		99	90	
03450020	**Acceptable identification**						**03450020**
					GRA	CAT	
Fusobacterium mortiferum		92.80	0.36	NAG 1	1	5	
Propionibacterium granulosum		7.20	0.13		99	90	
03450040	**Low discrimination**						**03450040**
					COC		
Fusobacterium mortiferum		53.35	0.36	SOR 1	1		
Megasphaera elsdenii		46.65	0.26		99		
03450100	**Very good identification**						**03450100**
					GRA		
Fusobacterium mortiferum		99.66	0.76		1		
Actinomyces meyeri		0.28	0.23		99		
Lactobacillus fermentum		0.07	0.24		99		
03450101	**Good identification**						**03450101**
Fusobacterium mortiferum		100.0	0.42	URE 1			
03450102	**Very good identification**						**03450102**
					GRA		
Fusobacterium mortiferum		99.23	0.60	BGL 10	1		
Lactobacillus fermentum		0.63	0.24		99		
Actinomyces viscosus		0.14	-.02		99		
03450110	**Good identification**						**03450110**
Fusobacterium mortiferum		100.0	0.42	MLZ 1			
03450120	**Acceptable identification**						**03450120**
					GRA		
Fusobacterium mortiferum		91.38	0.42	NAG 1	1		
Actinomyces meyeri		3.40	0.08		99		
Prevotella melaninogenica		3.10	-.01		1		
03450140	**Good identification**						**03450140**
Fusobacterium mortiferum		100.0	0.42	SOR 1			
03450200	**Good identification**						**03450200**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
03450300	**Good identification**						**03450300**
Fusobacterium mortiferum		100.0	0.42	RHA 1			
03450400	**Low discrimination**						**03450400**
					GRA		
Fusobacterium mortiferum		87.76	0.36	ARA 1	1		
Lactobacillus fermentum		8.75	0.20		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		3.49	0.02		99		
03450500	**Low discrimination**						**03450500**
					GRA		
Fusobacterium mortiferum		87.88	0.42	ARA 1	1		
Lactobacillus fermentum		6.13	0.24		99		
Actinomyces meyeri		5.99	0.13		99		
03451000	**Good identification**						**03451000**
					GRA	CAT	
Fusobacterium mortiferum		93.30	0.73		1	5	
Lactobacillus fermentum		6.20	0.54		99	1	
Propionibacterium granulosum		0.20	0.24		99	90	
03451001	**Low discrimination**						**03451001**
					GRA		
Fusobacterium mortiferum		79.67	0.39	URE 1	1		
Actinomyces naeslundii		15.03	0.38		99		
Lactobacillus fermentum		5.29	0.20		99		
03451002	**Low discrimination**						**03451002**
					GRA		
Fusobacterium mortiferum		61.95	0.56	BGL 10	1		
Lactobacillus fermentum		37.05	0.54		99		
Actinomyces naeslundii		0.71	0.35		99		
03451003	**Low discrimination**						**03451003**
					GRA		
Actinomyces naeslundii		51.52	0.38	MNS 88	99		
Fusobacterium mortiferum		30.34	0.22	MLT 94	1		
Lactobacillus fermentum		18.15	0.20		99		
03451010	**Acceptable identification**						**03451010**
					GRA	CAT	
Fusobacterium mortiferum		91.35	0.39	MLZ 1	1	5	
Lactobacillus fermentum		6.07	0.20		99	1	
Propionibacterium granulosum		2.58	0.09		99	90	
03451020	**Low discrimination**						**03451020**
					GRA		
Lactobacillus fermentum		64.06	0.46	CEL 1	99		
Fusobacterium mortiferum		29.21	0.39		1		
Actinomyces naeslundii		3.67	0.35		99		
03451021	**Good identification**						**03451021**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03451022	**Acceptable identification**						**03451022**
					GRA		
Lactobacillus fermentum		90.25	0.46	CEL 1	99		
Actinomyces naeslundii		5.18	0.35		99		
Fusobacterium mortiferum		4.57	0.22		1		
03451023	**Good identification**						**03451023**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03451040	**Acceptable identification**						**03451040**
					GRA		
Fusobacterium mortiferum		93.77	0.39	SOR 1	1		
Lactobacillus fermentum		6.23	0.20		99		
03451100	**Very good identification**						**03451100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		95.15	0.79		1	
Lactobacillus fermentum		4.42	0.58		99	
Actinomyces meyeri		0.26	0.26		99	
03451101	**Low discrimination**					**03451101**
					GRA	
Fusobacterium mortiferum		78.33	0.45	URE 1	1	
Actinomyces naeslundii		18.03	0.46		99	
Lactobacillus fermentum		3.64	0.24		99	
03451102	**Low discrimination**					**03451102**
					GRA	
Fusobacterium mortiferum		69.83	0.63	BGL 10	1	
Lactobacillus fermentum		29.20	0.58		99	
Actinomyces naeslundii		0.97	0.42		99	
03451103	**Low discrimination**					**03451103**
					GRA	
Actinomyces naeslundii		59.36	0.46	MNS 88	99	
Fusobacterium mortiferum		28.66	0.29	MLT 94	1	
Lactobacillus fermentum		11.98	0.24		99	
03451110	**Good identification**					**03451110**
					GRA	
Fusobacterium mortiferum		95.56	0.45	MLZ 1	1	
Lactobacillus fermentum		4.44	0.24		99	
03451121	**Low discrimination**					**03451121**
					GRA	
Actinomyces naeslundii		85.80	0.46	MNS 88	99	
Prevotella melaninogenica		8.42	-.01	MLT 94	1	
Lactobacillus fermentum		5.77	0.15		99	
03451122	**Low discrimination**					**03451122**
					GRA	
Lactobacillus fermentum		64.49	0.50	CEL 1	99	
Prevotella melaninogenica		17.92	0.21		1	
Actinomyces naeslundii		6.45	0.42		99	
03451123	**Acceptable identification**					**03451123**
Actinomyces naeslundii		93.70	0.46	MNS 88		
Lactobacillus fermentum		6.31	0.15	MLT 94		
03451140	**Acceptable identification**					**03451140**
					GRA	
Fusobacterium mortiferum		93.15	0.45	SOR 1	1	
Lactobacillus fermentum		4.33	0.24		99	
Actinomyces naeslundii		2.52	0.30		99	
03451200	**Acceptable identification**					**03451200**
					GRA	
Fusobacterium mortiferum		93.77	0.39	RHA 1	1	
Lactobacillus fermentum		6.23	0.20		99	
03451300	**Good identification**					**03451300**
					GRA	
Fusobacterium mortiferum		95.56	0.45	RHA 1	1	
Lactobacillus fermentum		4.44	0.24		99	
03451400	**Low discrimination**					**03451400**
					GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		86.35	0.54	CEL 1	99		
Fusobacterium mortiferum		13.13	0.39		1		
Actinomyces meyeri		0.52	0.05		99		
03451402	**Good identification**						**03451402**
					GRA		
Lactobacillus fermentum		98.34	0.54	CEL 1	99		
Fusobacterium mortiferum		1.66	0.22		1		
03451420	**Good identification**						**03451420**
Lactobacillus fermentum		100.0	0.46	CEL 1			
03451422	**Good identification**						**03451422**
Lactobacillus fermentum		100.0	0.46	CEL 1			
03451500	**Low discrimination**						**03451500**
					GRA		
Lactobacillus fermentum		81.16	0.58	CEL 1	99		
Fusobacterium mortiferum		17.64	0.45		1		
Actinomyces meyeri		1.20	0.16		99		
03451502	**Very good identification**						**03451502**
					GRA		
Lactobacillus fermentum		97.17	0.58	CEL 1	99		
Fusobacterium mortiferum		2.35	0.29		1		
Bifidobacterium adolescentis		0.49	0.13		99		
03451520	**Good identification**						**03451520**
					GRA		
Lactobacillus fermentum		97.89	0.50	CEL 1	99		
Prevotella melaninogenica		1.44	-.01		1		
Bifidobacterium longum subsp. longum		0.67	0.05		99		
03451522	**Good identification**						**03451522**
					GRA		
Lactobacillus fermentum		98.53	0.50	CEL 1	99		
Parabacteroides distasonis		0.80	0.11		1		
Bifidobacterium longum subsp. longum		0.68	0.05		99		
03452100	**Good identification**						**03452100**
					GRA		
Fusobacterium mortiferum		96.70	0.42	MAN 1	1		
Actinomyces naeslundii		1.80	0.24		99		
Propionibacterium propionicum		1.50	0.12		99		
03453000	**Low discrimination**						**03453000**
					GRA	CAT	
Fusobacterium mortiferum		68.18	0.39	MAN 1	1	5	
Actinomyces naeslundii		16.65	0.40		99	25	
Propionibacterium granulosum		10.64	0.21		99	90	
03453001	**Good identification**						**03453001**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03453002	**Low discrimination**						**03453002**
					GRA		
Actinomyces naeslundii		53.00	0.40	MNS 88	99		
Fusobacterium mortiferum		24.12	0.22	MLT 94	1		
Lactobacillus fermentum		14.42	0.20		99		
03453003	**Good identification**						**03453003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	MNS 88		
03453020	**Low discrimination**					**03453020**
Actinomyces naeslundii		86.23	0.40	MNS 88		
Propionibacterium granulosum		13.78	0.11	MLT 94		
03453021	**Good identification**					**03453021**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03453022	**Good identification**					**03453022**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03453023	**Good identification**					**03453023**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03453100	**Low discrimination**					**03453100**
					GRA	
Fusobacterium mortiferum		73.82	0.45	MAN 1	1	
Actinomyces naeslundii		21.99	0.47		99	
Lactobacillus fermentum		3.43	0.24		99	
03453101	**Good identification**					**03453101**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03453102	**Low discrimination**					**03453102**
					GRA	
Actinomyces naeslundii		63.25	0.47	MNS 88	99	
Fusobacterium mortiferum		23.60	0.29	MLT 94	1	
Lactobacillus fermentum		9.87	0.24		99	
03453103	**Good identification**					**03453103**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03453120	**Low discrimination**					**03453120**
					GRA	
Actinomyces naeslundii		88.66	0.47	MNS 88	99	
Prevotella melaninogenica		6.73	-.01	MLT 94	1	
Lactobacillus fermentum		4.61	0.15		99	
03453121	**Good identification**					**03453121**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03453122	**Good identification**					**03453122**
Actinomyces naeslundii		95.06	0.47	MNS 88		
Lactobacillus fermentum		4.94	0.15	MLT 94		
03453123	**Good identification**					**03453123**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03453141	**Good identification**					**03453141**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03453143	**Good identification**					**03453143**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03453161	**Good identification**					**03453161**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03453163	**Good identification**					**03453163**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03454000	**Good identification**					**03454000**
					GRA	
Fusobacterium mortiferum		95.30	0.50	XYL 7	1	
Actinomyces meyeri		3.95	0.18		99	
Lactobacillus fermentum		0.75	0.16		99	
03454100	**Good identification**					**03454100**
					GRA	
Fusobacterium mortiferum		92.89	0.57	XYL 7	1	
Actinomyces meyeri		6.60	0.28		99	
Lactobacillus fermentum		0.51	0.20		99	
03454102	**Acceptable identification**					**03454102**
					GRA	
Fusobacterium mortiferum		93.53	0.40	XYL 7	1	
Lactobacillus fermentum		4.62	0.20	BGL 10	99	
Actinomyces meyeri		1.85	0.02		99	
03455000	**Low discrimination**					**03455000**
					GRA	
Fusobacterium mortiferum		64.11	0.53	XYL 7	1	
Lactobacillus fermentum		33.24	0.50		99	
Actinomyces meyeri		2.66	0.21		99	
03455002	**Low discrimination**					**03455002**
					GRA	
Lactobacillus fermentum		82.35	0.50	CEL 1	99	
Fusobacterium mortiferum		17.65	0.37		1	
03455020	**Acceptable identification**					**03455020**
					GRA	
Lactobacillus fermentum		91.65	0.42	CEL 1	99	
Fusobacterium mortiferum		5.36	0.19		1	
Actinomyces meyeri		3.00	0.06		99	
03455022	**Good identification**					**03455022**
Lactobacillus fermentum		100.0	0.42	CEL 1		
03455100	**Low discrimination**					**03455100**
					GRA	
Fusobacterium mortiferum		69.75	0.60	XYL 7	1	
Lactobacillus fermentum		25.29	0.54		99	
Actinomyces meyeri		4.96	0.31		99	
03455102	**Low discrimination**					**03455102**
					GRA	
Lactobacillus fermentum		75.46	0.54	CEL 1	99	
Fusobacterium mortiferum		23.13	0.43		1	
Bifidobacterium adolescentis		0.65	0.13		99	
03455120	**Low discrimination**					**03455120**
					GRA	
Lactobacillus fermentum		84.12	0.46	CEL 1	99	
Fusobacterium mortiferum		7.03	0.26		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		6.74	0.16		99		
03455122	**Low discrimination**						**03455122**
					GRA	CAT	
Lactobacillus fermentum		80.93	0.46	CEL 1	99	1	
Parabacteroides distasonis		16.33	0.31		1	76	
Bacteroides fragilis		1.48	-.09		1	99	
03455322	**Good identification**						**03455322**
Parabacteroides distasonis		100.0	0.39	MNS 98			
03455400	**Good identification**						**03455400**
					GRA		
Lactobacillus fermentum		96.20	0.50	CEL 1	99		
Actinomyces meyeri		1.92	0.10		99		
Fusobacterium mortiferum		1.87	0.19		1		
03455402	**Good identification**						**03455402**
Lactobacillus fermentum		100.0	0.50	CEL 1			
03455420	**Good identification**						**03455420**
Lactobacillus fermentum		100.0	0.42	CEL 1			
03455422	**Good identification**						**03455422**
Lactobacillus fermentum		100.0	0.42	CEL 1			
03455500	**Acceptable identification**						**03455500**
					GRA		
Lactobacillus fermentum		92.86	0.54	CEL 1	99		
Actinomyces meyeri		4.55	0.21		99		
Fusobacterium mortiferum		2.59	0.26		1		
03455502	**Acceptable identification**						**03455502**
Lactobacillus fermentum		94.59	0.54	CEL 1			
Bifidobacterium adolescentis		5.42	0.27				
03455520	**Acceptable identification**						**03455520**
Lactobacillus fermentum		94.77	0.46	CEL 1			
Bifidobacterium longum subsp. longum		3.33	0.14				
Actinomyces meyeri		1.90	0.06				
03455522	**Low discrimination**						**03455522**
					GRA		
Lactobacillus fermentum		86.30	0.46	CEL 1	99		
Parabacteroides distasonis		10.67	0.27		1		
Bifidobacterium longum subsp. longum		3.03	0.14		99		
03455722	**Good identification**						**03455722**
Parabacteroides distasonis		100.0	0.35	MNS 98			
03461100	**Good identification**						**03461100**
					GRA		
Fusobacterium mortiferum		95.01	0.36	FRU 99	1		
Actinomyces naeslundii		4.99	0.26		99		
03470000	**Very good identification**						**03470000**
					GRA	CAT	
Fusobacterium mortiferum		96.95	0.61		1	5	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		3.05	0.32		99	90	
03470002	**Acceptable identification**						**03470002**
					GRA	CAT	
Fusobacterium mortiferum		93.39	0.44	BGL 10	1	5	
Propionibacterium granulosum		6.61	0.21		99	90	
03470100	**Very good identification**						**03470100**
					GRA		
Fusobacterium mortiferum		99.94	0.67		1		
Actinomyces naeslundii		0.06	0.24		99		
03470102	**Good identification**						**03470102**
					GRA		
Fusobacterium mortiferum		99.44	0.51	BGL 10	1		
Actinomyces naeslundii		0.56	0.24		99		
03471000	**Very good identification**						**03471000**
					GRA	CAT	
Fusobacterium mortiferum		96.95	0.64		1	5	
Propionibacterium granulosum		2.03	0.32		99	90	
Actinomyces naeslundii		0.80	0.40		99	25	
03471001	**Low discrimination**						**03471001**
					GRA		
Actinomyces naeslundii		55.08	0.43	MNS 88	99		
Fusobacterium mortiferum		44.92	0.30	MLT 94	1		
03471002	**Low discrimination**						**03471002**
					GRA	CAT	
Fusobacterium mortiferum		87.59	0.47	BGL 10	1	5	
Actinomyces naeslundii		6.51	0.40		99	25	
Propionibacterium granulosum		4.14	0.21		99	90	
03471003	**Acceptable identification**						**03471003**
					GRA		
Actinomyces naeslundii		91.69	0.43	MNS 88	99		
Fusobacterium mortiferum		8.31	0.13	MLT 94	1		
03471021	**Good identification**						**03471021**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03471022	**Low discrimination**						**03471022**
					GRA	CAT	
Actinomyces naeslundii		77.23	0.40	MNS 88	99	25	
Propionibacterium granulosum		12.27	0.11	MLT 94	99	90	
Fusobacterium mortiferum		10.50	0.13		1	5	
03471023	**Good identification**						**03471023**
Actinomyces naeslundii		100.0	0.43	MNS 88			
03471100	**Very good identification**						**03471100**
					GRA		
Fusobacterium mortiferum		98.85	0.70		1		
Actinomyces naeslundii		1.00	0.47		99		
Lactobacillus fermentum		0.16	0.24		99		
03471101	**Low discrimination**						**03471101**
					GRA		
Actinomyces naeslundii		59.93	0.50	MNS 88	99		
Fusobacterium mortiferum		40.07	0.36	MLT 94	1		
03471102	**Acceptable identification**						**03471102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		90.22	0.54	BGL 10	1	
Actinomyces naeslundii		8.18	0.47		99	
Lactobacillus fermentum		1.28	0.24		99	
03471103	**Acceptable identification**					**03471103**
					GRA	
Actinomyces naeslundii		93.09	0.50	MNS 88	99	
Fusobacterium mortiferum		6.92	0.20	MLT 94	1	
03471110	**Good identification**					**03471110**
					GRA	
Fusobacterium mortiferum		95.01	0.36	MLZ 1	1	
Actinomyces naeslundii		4.99	0.26		99	
03471120	**Low discrimination**					**03471120**
					GRA	
Fusobacterium mortiferum		47.07	0.36	NAG 1	1	
Actinomyces naeslundii		46.93	0.47		99	
Prevotella melaninogenica		3.56	-.01		1	
03471121	**Good identification**					**03471121**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03471122	**Low discrimination**					**03471122**
					GRA	
Actinomyces naeslundii		66.16	0.47	MNS 88	99	
Parabacteroides distasonis		20.42	0.26	MLT 94	1	
Fusobacterium mortiferum		7.37	0.20		1	
03471123	**Good identification**					**03471123**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03471140	**Low discrimination**					**03471140**
					GRA	
Fusobacterium mortiferum		85.04	0.36	SOR 1	1	
Actinomyces naeslundii		14.96	0.35		99	
03471141	**Good identification**					**03471141**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03471143	**Good identification**					**03471143**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03471161	**Good identification**					**03471161**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03471163	**Good identification**					**03471163**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03471300	**Good identification**					**03471300**
Fusobacterium mortiferum		100.0	0.36	RHA 1		
03471500	**Low discrimination**					**03471500**
					GRA	
Fusobacterium mortiferum		86.54	0.36	ARA 1	1	
Lactobacillus fermentum		13.46	0.24		99	
03473000	**Low discrimination**					**03473000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces naeslundii		38.20	0.45	MNS 88	99	25	
Propionibacterium granulosum		37.73	0.30	MLT 94	99	90	
Fusobacterium mortiferum		24.08	0.30		1	5	
03473001	**Good identification**						**03473001**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03473002	**Low discrimination**						**03473002**
					GRA	CAT	
Actinomyces naeslundii		75.93	0.45	MNS 88	99	25	
Propionibacterium granulosum		18.75	0.19	MLT 94	99	90	
Fusobacterium mortiferum		5.32	0.13		1	5	
03473003	**Good identification**						**03473003**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03473020	**Low discrimination**						**03473020**
Actinomyces naeslundii		80.20	0.45	MNS 88			
Propionibacterium granulosum		19.80	0.19	MLT 94			
03473021	**Good identification**						**03473021**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03473022	**Acceptable identification**						**03473022**
Actinomyces naeslundii		94.19	0.45	MNS 88			
Propionibacterium granulosum		5.81	0.09	MLT 94			
03473023	**Good identification**						**03473023**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03473100	**Low discrimination**						**03473100**
					GRA		
Actinomyces naeslundii		64.92	0.52	MNS 88	99		
Fusobacterium mortiferum		33.54	0.36	MLT 94	1		
Actinomyces israelii		1.53	0.15		99		
03473101	**Good identification**						**03473101**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03473102	**Low discrimination**						**03473102**
					GRA		
Actinomyces naeslundii		89.71	0.52	MNS 88	99		
Fusobacterium mortiferum		5.15	0.20	MLT 94	1		
Actinomyces israelii		3.18	0.18		99		
03473103	**Good identification**						**03473103**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03473120	**Good identification**						**03473120**
Actinomyces naeslundii		100.0	0.52	MNS 88			
03473121	**Good identification**						**03473121**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03473122	**Good identification**						**03473122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	MNS 88		
03473123	**Good identification**					**03473123**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03473140	**Good identification**					**03473140**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03473141	**Good identification**					**03473141**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03473142	**Low discrimination**					**03473142**
Actinomyces naeslundii		88.95	0.39	MNS 88		
Propionibacterium propionicum		11.05	0.14	MLT 94		
03473143	**Good identification**					**03473143**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03473160	**Good identification**					**03473160**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03473161	**Good identification**					**03473161**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03473162	**Good identification**					**03473162**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03473163	**Good identification**					**03473163**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03474000	**Good identification**					**03474000**
Fusobacterium mortiferum		100.0	0.41	XYL 7		
03474100	**Good identification**					**03474100**
Fusobacterium mortiferum		100.0	0.48	XYL 7		
03475000	**Good identification**					**03475000**
					GRA	
Fusobacterium mortiferum		98.28	0.44	XYL 7	1	
Lactobacillus fermentum		1.72	0.16		99	
03475100	**Good identification**					**03475100**
					GRA	
Fusobacterium mortiferum		97.59	0.51	XYL 7	1	
Actinomyces israelii		1.22	0.21		99	
Lactobacillus fermentum		1.20	0.20		99	
03475122	**Good identification**					**03475122**
Parabacteroides distasonis		100.0	0.42	MNS 98		
03475132	**Good identification**					**03475132**
Parabacteroides distasonis		100.0	0.37	MNS 98		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
03475322	**Good identification**							**03475322**
Parabacteroides distasonis		100.0	0.50	MNS 98				
03475332	**Good identification**							**03475332**
Parabacteroides distasonis		100.0	0.46	MNS 98				
03475522	**Good identification**							**03475522**
Parabacteroides distasonis		100.0	0.38	MNS 98				
03475722	**Good identification**							**03475722**
Parabacteroides distasonis		100.0	0.47	MNS 98				
03475732	**Good identification**							**03475732**
Parabacteroides distasonis		97.85	0.42	MNS 98				
Clostridium clostridioforme		2.15	0.08					
03477102	**Acceptable identification**							**03477102**
Actinomyces israelii		94.92	0.36	MLT 99				
Bifidobacterium adolescentis		5.08	0.08	SAL 92				
03500000	**Low discrimination**							**03500000**
					COC	GRA	SPO	
Clostridium botulinum biovar A		29.32	0.41	SUC 7	1	99	99	
Megasphaera elsdenii		26.14	0.44	RAF 1	99	1	1	
Prevotella intermedia		12.94	0.36		1	1	1	
03500040	**Low discrimination**							**03500040**
					COC	GRA		
Megasphaera elsdenii		60.65	0.41	RAF 1	99	1		
Clostridium botulinum biovar A		30.48	0.32	FRU 95	1	99		
Clostridium novyi biovar A		4.24	0.29		1	99		
03501000	**Low discrimination**							**03501000**
					COC	GRA		
Lactobacillus fermentum		66.30	0.62	FRU 99	1	99		
Blautia hansenii		13.82	0.32		99	99		
Prevotella intermedia		11.36	0.36		1	1		
03501002	**Very good identification**							**03501002**
					COC			
Lactobacillus fermentum		95.98	0.62	FRU 99	1			
Blautia hansenii		2.22	0.15		99			
Actinomyces viscosus		0.79	0.12		1			
03501020	**Low discrimination**							**03501020**
					COC	GRA		
Lactobacillus fermentum		52.33	0.54	FRU 99	1	99		
Prevotella melaninogenica		41.51	0.33		1	1		
Blautia hansenii		3.64	0.15		99	99		
03501022	**Low discrimination**							**03501022**
					GRA			
Lactobacillus fermentum		85.42	0.54	FRU 99	99			
Prevotella melaninogenica		12.91	0.21		1			
Actinomyces naeslundii		1.02	0.31		99			
03501100	**Good identification**							**03501100**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		90.55	0.66	SUC 1	99	99	
Lactobacillus fermentum		7.16	0.66		1	99	
Prevotella melaninogenica		1.16	0.33		1	1	
03501102	**Low discrimination**						**03501102**
					COC	GRA	
Blautia hansenii		57.09	0.50	SUC 1	99	99	
Lactobacillus fermentum		40.62	0.66	BGL 10	1	99	
Prevotella melaninogenica		1.25	0.21		1	1	
03501120	**Good identification**						**03501120**
					COC	GRA	
Prevotella melaninogenica		90.04	0.67	MNS 90	1	1	
Blautia hansenii		7.89	0.50	FRU 90	99	99	
Lactobacillus fermentum		1.87	0.58		1	99	
03501122	**Low discrimination**						**03501122**
					COC	GRA	
Prevotella melaninogenica		85.56	0.55	MNS 90	1	1	
Lactobacillus fermentum		9.33	0.58	FRU 90	1	99	
Blautia hansenii		4.37	0.33		99	99	
03501400	**Good identification**						**03501400**
					COC		
Lactobacillus fermentum		94.54	0.62	FRU 99	1		
Blautia hansenii		3.48	0.19		99		
Actinomyces meyeri		1.12	0.18		1		
03501402	**Very good identification**						**03501402**
					COC		
Lactobacillus fermentum		99.25	0.62	FRU 99	1		
Blautia hansenii		0.41	0.03		99		
Bifidobacterium longum subsp. longum		0.34	0.13		1		
03501420	**Acceptable identification**						**03501420**
					COC		
Lactobacillus fermentum		93.79	0.54	FRU 99	1		
Bifidobacterium longum subsp. longum		3.86	0.23		1		
Blautia hansenii		1.15	0.03		99		
03501422	**Good identification**						**03501422**
Lactobacillus fermentum		96.05	0.54	FRU 99			
Bifidobacterium longum subsp. longum		3.95	0.23				
03501500	**Low discrimination**						**03501500**
					COC		
Blautia hansenii		64.32	0.53	SUC 1	99		
Lactobacillus fermentum		28.81	0.66		1		
Bifidobacterium longum subsp. longum		5.99	0.47		1		
03501502	**Low discrimination**						**03501502**
					COC		
Lactobacillus fermentum		68.61	0.66	FRU 99	1		
Blautia hansenii		17.02	0.37		99		
Bifidobacterium longum subsp. longum		14.26	0.47		1		
03501510	**Acceptable identification**						**03501510**
					COC		
Bifidobacterium longum subsp. longum		91.56	0.51	MNS 87	1		
Blautia hansenii		5.83	0.19	FRU 90	99		
Lactobacillus fermentum		2.61	0.32		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03501512	**Good identification**							**03501512**
					COC			
Bifidobacterium longum subsp. longum		96.31	0.51	MNS 87	1			
Lactobacillus fermentum		2.75	0.32	FRU 90	1			
Blautia hansenii		0.68	0.03		99			
03501520	**Low discrimination**							**03501520**
					COC			
Bifidobacterium longum subsp. longum		52.66	0.57	MNS 87	1			
Lactobacillus fermentum		21.11	0.58	FRU 90	1			
Blautia hansenii		15.71	0.37		99			
03501522	**Low discrimination**							**03501522**
					GRA			
Bifidobacterium longum subsp. longum		67.92	0.57	MNS 87	99			
Lactobacillus fermentum		27.23	0.58	FRU 90	99			
Prevotella melaninogenica		2.52	0.21		1			
03501530	**Very good identification**							**03501530**
					COC			
Bifidobacterium longum subsp. longum		99.47	0.61	MNS 87	1			
Lactobacillus fermentum		0.24	0.24	FRU 90	1			
Blautia hansenii		0.18	0.03		99			
03501532	**Very good identification**							**03501532**
Bifidobacterium longum subsp. longum		99.76	0.61	MNS 87				
Lactobacillus fermentum		0.24	0.24	FRU 90				
03502000	**Good identification**							**03502000**
					COC	SPO	CAT	
Megasphaera elsdenii		96.67	0.44	RAF 1	99	1	1	
Clostridium botulinum biovar A		1.10	0.06	FRU 95	1	99	1	
Propionibacterium granulosum		0.99	0.16		1	1	90	
03502040	**Good identification**							**03502040**
					COC	GRA		
Megasphaera elsdenii		99.47	0.41	RAF 1	99	1		
Propionibacterium propionicum		0.53	0.12	FRU 95	1	99		
03502100	**Acceptable identification**							**03502100**
					COC	GRA		
Propionibacterium propionicum		94.62	0.46	NIT 86	1	99		
Megasphaera elsdenii		2.74	0.10	FRU 99	99	1		
Blautia hansenii		1.65	-.03		99	99		
03502102	**Good identification**							**03502102**
Propionibacterium propionicum		100.0	0.49	NIT 86				
03502140	**Good identification**							**03502140**
					COC	GRA		
Propionibacterium propionicum		98.11	0.46	NIT 86	1	99		
Megasphaera elsdenii		1.89	0.07	FRU 99	99	1		
03502142	**Good identification**							**03502142**
Propionibacterium propionicum		100.0	0.49	NIT 86				
03503001	**Good identification**							**03503001**
Actinomyces naeslundii		100.0	0.39	MNS 88				
03503003	**Good identification**							**03503003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	MNS 88		
03503020	**Low discrimination**					**03503020**
					GRA	
Actinomyces naeslundii		56.04	0.36	MNS 88	99	
Lactobacillus fermentum		24.52	0.20	FRU 95	99	
Prevotella melaninogenica		19.45	-.01		1	
03503021	**Good identification**					**03503021**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03503022	**Low discrimination**					**03503022**
Actinomyces naeslundii		69.56	0.36	MNS 88		
Lactobacillus fermentum		30.44	0.20	FRU 95		
03503023	**Good identification**					**03503023**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03503101	**Low discrimination**					**03503101**
					COC	
Actinomyces naeslundii		65.19	0.47	MNS 88	1	
Propionibacterium propionicum		26.66	0.30	FRU 95	1	
Blautia hansenii		4.18	-.03		99	
03503102	**Low discrimination**					**03503102**
					COC	
Propionibacterium propionicum		74.66	0.49	NIT 86	1	
Blautia hansenii		9.54	0.15	FRU 99	99	
Actinomyces naeslundii		9.02	0.44		1	
03503103	**Low discrimination**					**03503103**
Actinomyces naeslundii		61.98	0.47	MNS 88		
Propionibacterium propionicum		38.02	0.33	FRU 95		
03503121	**Acceptable identification**					**03503121**
					GRA	
Actinomyces naeslundii		92.48	0.47	MNS 88	99	
Prevotella melaninogenica		7.52	-.01	FRU 95	1	
03503123	**Good identification**					**03503123**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03503140	**Good identification**					**03503140**
					COC	
Propionibacterium propionicum		95.29	0.46	NIT 86	1	
Actinomyces naeslundii		3.05	0.31	FRU 99	1	
Blautia hansenii		1.66	-.03		99	
03503142	**Good identification**					**03503142**
Propionibacterium propionicum		97.91	0.49	NIT 86		
Actinomyces naeslundii		2.09	0.31	FRU 99		
03504100	**Low discrimination**					**03504100**
					COC	
Actinomyces meyeri		83.44	0.41	RAF 1	1	
Blautia hansenii		12.52	0.19	FRU 95	99	
Lactobacillus fermentum		3.29	0.28		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
03504530	**Good identification**						**03504530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03504532	**Good identification**						**03504532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03505000	**Low discrimination**						**03505000**
					COC		
Lactobacillus fermentum		81.22	0.58	FRU 99	1		
Actinomyces meyeri		12.71	0.34		1		
Blautia hansenii		5.09	0.19		99		
03505002	**Very good identification**						**03505002**
					COC		
Lactobacillus fermentum		98.83	0.58	FRU 99	1		
Blautia hansenii		0.69	0.03		99		
Actinomyces meyeri		0.48	0.08		1		
03505020	**Acceptable identification**						**03505020**
					COC		
Lactobacillus fermentum		91.03	0.50	FRU 99	1		
Actinomyces meyeri		5.83	0.19		1		
Blautia hansenii		1.90	0.03		99		
03505022	**Good identification**						**03505022**
Lactobacillus fermentum		100.0	0.50	FRU 99			
03505100	**Low discrimination**						**03505100**
					COC		
Blautia hansenii		73.04	0.53	SUC 1	99		
Lactobacillus fermentum		19.22	0.62		1		
Actinomyces meyeri		7.37	0.44		1		
03505102	**Low discrimination**						**03505102**
					COC		
Lactobacillus fermentum		69.14	0.62	FRU 99	1		
Blautia hansenii		29.20	0.37		99		
Actinomyces meyeri		0.82	0.18		1		
03505120	**Low discrimination**						**03505120**
					COC	GRA	
Blautia hansenii		37.48	0.37	SUC 1	99	99	
Lactobacillus fermentum		29.58	0.54	NAG 10	1	99	
Prevotella melaninogenica		24.49	0.33		1	1	
03505122	**Low discrimination**						**03505122**
					COC	GRA	
Lactobacillus fermentum		68.58	0.54	FRU 99	1	99	
Prevotella melaninogenica		10.81	0.21		1	1	
Blautia hansenii		9.65	0.20		99	99	
03505130	**Low discrimination**						**03505130**
					COC		
Bifidobacterium longum subsp. longum		87.52	0.35	MNS 87	1		
Blautia hansenii		5.11	0.03	FRU 90	99		
Lactobacillus fermentum		4.03	0.20	ARA 99	1		
03505132	**Good identification**						**03505132**
Bifidobacterium longum subsp. longum		95.59	0.35	MNS 87			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		4.41	0.20	FRU 90		
03505400	**Good identification**					**03505400**
Lactobacillus fermentum		93.66	0.58	FRU 99		
Actinomyces meyeri		3.67	0.23			
Bifidobacterium longum subsp. longum		1.64	0.21			
03505402	**Very good identification**					**03505402**
Lactobacillus fermentum		98.28	0.58	FRU 99		
Bifidobacterium longum subsp. longum		1.72	0.21			
03505420	**Low discrimination**					**03505420**
Lactobacillus fermentum		81.57	0.50	FRU 99		
Bifidobacterium longum subsp. longum		17.13	0.31			
Actinomyces meyeri		1.31	0.09			
03505422	**Low discrimination**					**03505422**
Lactobacillus fermentum		82.65	0.50	FRU 99		
Bifidobacterium longum subsp. longum		17.36	0.31			
03505430	**Good identification**					**03505430**
Bifidobacterium longum subsp. longum		97.25	0.35	MNS 87		
Lactobacillus fermentum		2.75	0.16	FRU 90		
03505432	**Good identification**					**03505432**
Bifidobacterium longum subsp. longum		97.25	0.35	MNS 87		
Lactobacillus fermentum		2.75	0.16	FRU 90		
03505500	**Low discrimination**					**03505500**
					COC	
Bifidobacterium longum subsp. longum		37.54	0.55	MNS 87	1	
Lactobacillus fermentum		35.36	0.62	FRU 90	1	
Blautia hansenii		23.71	0.40		99	
03505502	**Low discrimination**					**03505502**
					COC	
Bifidobacterium longum subsp. longum		49.43	0.55	MNS 87	1	
Lactobacillus fermentum		46.55	0.62	FRU 90	1	
Blautia hansenii		3.47	0.24		99	
03505510	**Very good identification**					**03505510**
					COC	
Bifidobacterium longum subsp. longum		99.08	0.59	MNS 87	1	
Lactobacillus fermentum		0.55	0.28	FRU 90	1	
Blautia hansenii		0.37	0.06		99	
03505512	**Very good identification**					**03505512**
Bifidobacterium longum subsp. longum		98.85	0.59	MNS 87		
Bifidobacterium adolescentis		0.60	0.23	FRU 90		
Lactobacillus fermentum		0.55	0.28			
03505520	**Good identification**					**03505520**
					COC	
Bifidobacterium longum subsp. longum		90.65	0.66	MNS 87	1	
Lactobacillus fermentum		7.11	0.54	FRU 90	1	
Blautia hansenii		1.59	0.24		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03505522	**Good identification**						**03505522**
					COC		
Bifidobacterium longum subsp. longum		92.56	0.66	MNS 87	1		
Lactobacillus fermentum		7.26	0.54	FRU 90	1		
Blautia hansenii		0.18	0.07		99		
03505530	**Very good identification**						**03505530**
Bifidobacterium longum subsp. longum		99.95	0.70	MNS 87			
Lactobacillus fermentum		0.05	0.20	FRU 90			
03505531	**Good identification**						**03505531**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03505532	**Very good identification**						**03505532**
Bifidobacterium longum subsp. longum		99.95	0.70	MNS 87			
Lactobacillus fermentum		0.05	0.20	FRU 90			
03505533	**Good identification**						**03505533**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03505570	**Good identification**						**03505570**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03505572	**Good identification**						**03505572**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03505730	**Good identification**						**03505730**
					GRA		
Bifidobacterium longum subsp. longum		98.24	0.35	MNS 87	99		
Bacteroides vulgatus		1.76	-.09	FRU 90	1		
03505732	**Good identification**						**03505732**
					GRA		
Bifidobacterium longum subsp. longum		98.02	0.35	MNS 87	99		
Parabacteroides distasonis		1.98	0.11	FRU 90	1		
03507530	**Good identification**						**03507530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03507532	**Good identification**						**03507532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
03510000	**Low discrimination**						**03510000**
					COC		
Prevotella intermedia		62.62	0.71		1		
Megasphaera elsdenii		24.28	0.66		99		
Fusobacterium mortiferum		4.58	0.62		1		
03510001	**Low discrimination**						**03510001**
					COC		
Prevotella intermedia		61.00	0.36	URE 1	1		
Megasphaera elsdenii		23.65	0.32		99		
Fusobacterium mortiferum		4.46	0.28		1		
03510002	**Low discrimination**						**03510002**
					COC	GRA	
Lactobacillus fermentum		40.66	0.62	GAL 99	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		29.72	0.50		99	1		
Actinomyces viscosus		10.36	0.38		1	99		
03510010	**Low discrimination**							**03510010**
					COC	GRA	CAT	
Prevotella intermedia		51.50	0.36	MLZ 1	1	1	1	
Megasphaera elsdenii		19.97	0.32		99	1	1	
Propionibacterium granulosum		19.10	0.37		1	99	90	
03510020	**Low discrimination**							**03510020**
					GRA	CAT		
Lactobacillus fermentum		40.39	0.54	GAL 99	99	1		
Prevotella intermedia		20.77	0.36		1	1		
Propionibacterium granulosum		14.12	0.41		99	90		
03510022	**Low discrimination**							**03510022**
					COC	CAT	GRA	
Lactobacillus fermentum		85.63	0.54	GAL 99	1	1	99	
Propionibacterium granulosum		7.48	0.31		1	90	99	
Megasphaera elsdenii		1.90	0.15		99	1	1	
03510040	**Good identification**							**03510040**
					COC	GRA	SPO	
Megasphaera elsdenii		94.00	0.63	RAF 1	99	1	1	
Prevotella intermedia		3.67	0.36		1	1	1	
Clostridium botulinum biovar B		1.23	0.24		1	99	99	
03510042	**Good identification**							**03510042**
					COC	GRA		
Megasphaera elsdenii		96.51	0.47	RAF 1	99	1		
Lactobacillus fermentum		2.00	0.28	BGL 10	1	99		
Actinomyces viscosus		0.51	0.04		1	99		
03510100	**Low discrimination**							**03510100**
					GRA			
Fusobacterium mortiferum		39.43	0.69	MLT 11	1			
Actinomyces meyeri		32.70	0.58		99			
Lactobacillus fermentum		22.23	0.66		99			
03510102	**Low discrimination**							**03510102**
					GRA			
Lactobacillus fermentum		60.63	0.66	GAL 99	99			
Actinomyces viscosus		16.84	0.42		99			
Fusobacterium mortiferum		11.95	0.52		1			
03510120	**Low discrimination**							**03510120**
					GRA			
Prevotella melaninogenica		72.03	0.50	MNS 90	1			
Lactobacillus fermentum		16.46	0.58	GAL 99	99			
Actinomyces meyeri		9.91	0.43		99			
03510122	**Low discrimination**							**03510122**
					GRA			
Lactobacillus fermentum		52.06	0.58	GAL 99	99			
Prevotella melaninogenica		43.39	0.37		1			
Actinomyces meyeri		0.97	0.17		99			
03510123	**Low discrimination**							**03510123**
					GRA			
Actinomyces naeslundii		53.60	0.40	MNS 88	99			
Lactobacillus fermentum		21.98	0.24	GAL 96	99			
Prevotella melaninogenica		18.32	0.03		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03510140	**Low discrimination**						**03510140**
					COC	GRA	
Propionibacterium propionicum		50.70	0.46	NIT 86	1	99	
Megasphaera elsdenii		18.58	0.29	MAN 99	99	1	
Fusobacterium mortiferum		12.26	0.34		1	1	
03510142	**Low discrimination**						**03510142**
					COC	GRA	
Propionibacterium propionicum		85.48	0.49	NIT 86	1	99	
Lactobacillus fermentum		7.77	0.32	MAN 99	1	99	
Megasphaera elsdenii		2.32	0.12		99	1	
03510200	**Low discrimination**						**03510200**
					COC	GRA	
Prevotella intermedia		49.10	0.36	RHA 1	1	1	
Actinomyces odontolyticus		19.51	0.35		1	99	
Megasphaera elsdenii		19.04	0.32		99	1	
03510400	**Low discrimination**						**03510400**
					GRA		
Lactobacillus fermentum		70.72	0.62	GAL 99	99		
Prevotella intermedia		12.12	0.36		1		
Actinomyces meyeri		10.61	0.37		99		
03510402	**Very good identification**						**03510402**
					COC	GRA	
Lactobacillus fermentum		98.26	0.62	GAL 99	1	99	
Megasphaera elsdenii		0.73	0.15		99	1	
Actinomyces meyeri		0.46	0.11		1	99	
03510420	**Acceptable identification**						**03510420**
					GRA		
Lactobacillus fermentum		93.43	0.54	GAL 99	99		
Actinomyces meyeri		5.73	0.23		99		
Prevotella intermedia		0.49	0.02		1		
03510422	**Good identification**						**03510422**
Lactobacillus fermentum		99.63	0.54	GAL 99			
Bifidobacterium longum subsp. longum		0.37	0.05				
03510500	**Low discrimination**						**03510500**
Lactobacillus fermentum		71.11	0.66	GAL 99			
Actinomyces meyeri		26.15	0.48				
Bifidobacterium longum subsp. longum		1.34	0.29				
03510502	**Very good identification**						**03510502**
Lactobacillus fermentum		96.43	0.66	GAL 99			
Bifidobacterium longum subsp. longum		1.82	0.29				
Actinomyces meyeri		1.10	0.22				
03510520	**Low discrimination**						**03510520**
Lactobacillus fermentum		70.35	0.58	GAL 99			
Bifidobacterium longum subsp. longum		15.95	0.40				
Actinomyces meyeri		10.58	0.33				
03510522	**Low discrimination**						**03510522**
					GRA		
Lactobacillus fermentum		80.45	0.58	GAL 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		18.24	0.40		99		
Prevotella melaninogenica		0.68	0.03		1		
03510530	**Good identification**						**03510530**
Bifidobacterium longum subsp. longum		97.45	0.44	MNS 87			
Lactobacillus fermentum		2.55	0.24	GAL 99			
03510532	**Good identification**						**03510532**
Bifidobacterium longum subsp. longum		97.45	0.44	MNS 87			
Lactobacillus fermentum		2.55	0.24	GAL 99			
03511000	**Low discrimination**						**03511000**
					GRA		
Lactobacillus fermentum		82.89	0.96		99		
Prevotella intermedia		14.21	0.71		1		
Fusobacterium mortiferum		1.56	0.65		1		
03511001	**Low discrimination**						**03511001**
					GRA		
Lactobacillus fermentum		75.66	0.62	URE 1	99		
Prevotella intermedia		12.97	0.36		1		
Actinomyces naeslundii		8.55	0.56		99		
03511002	**Excellent identification**						**03511002**
					GRA		
Lactobacillus fermentum		99.21	0.96		99		
Fusobacterium mortiferum		0.21	0.49		1		
Prevotella intermedia		0.17	0.36		1		
03511003	**Low discrimination**						**03511003**
Lactobacillus fermentum		85.55	0.62	URE 1			
Actinomyces naeslundii		9.67	0.56				
Actinomyces viscosus		4.45	0.26				
03511010	**Low discrimination**						**03511010**
					GRA	CAT	
Lactobacillus fermentum		78.64	0.62	MLZ 1	99	1	
Prevotella intermedia		13.48	0.36		1	1	
Propionibacterium granulosum		5.00	0.37		99	90	
03511012	**Very good identification**						**03511012**
					CAT		
Lactobacillus fermentum		97.54	0.62	MLZ 1	1		
Propionibacterium granulosum		1.55	0.27		90		
Actinomyces naeslundii		0.39	0.31		25		
03511020	**Good identification**						**03511020**
					GRA		
Lactobacillus fermentum		91.63	0.88		99		
Prevotella melaninogenica		6.61	0.50		1		
Prevotella intermedia		0.48	0.36		1		
03511021	**Low discrimination**						**03511021**
					GRA		
Lactobacillus fermentum		70.19	0.54	URE 1	99		
Actinomyces naeslundii		23.79	0.56		99		
Prevotella melaninogenica		5.06	0.15		1		
03511022	**Very good identification**						**03511022**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		98.04	0.88		99		
Prevotella melaninogenica		1.35	0.37		1		
Actinomyces naeslundii		0.22	0.53		99		
03511023	**Low discrimination**						**03511023**
					GRA		
Lactobacillus fermentum		73.93	0.54	URE 1	99		
Actinomyces naeslundii		25.06	0.56		99		
Prevotella melaninogenica		1.02	0.03		1		
03511030	**Low discrimination**						**03511030**
					GRA	CAT	
Lactobacillus fermentum		87.19	0.54	MLZ 1	99	1	
Prevotella melaninogenica		6.29	0.15		1	1	
Propionibacterium granulosum		4.16	0.27		99	90	
03511032	**Good identification**						**03511032**
					GRA		
Lactobacillus fermentum		95.79	0.54	MLZ 1	99		
Prevotella melaninogenica		1.32	0.03		1		
Actinomyces naeslundii		1.14	0.31		99		
03511040	**Low discrimination**						**03511040**
					COC	GRA	
Lactobacillus fermentum		79.19	0.62	SOR 1	1	99	
Prevotella intermedia		13.57	0.36		1	1	
Megasphaera elsdenii		3.51	0.29		99	1	
03511041	**Low discrimination**						**03511041**
					GRA		
Actinomyces naeslundii		62.76	0.43	MNS 88	99		
Lactobacillus fermentum		31.79	0.28		99		
Prevotella intermedia		5.45	0.02		1		
03511042	**Very good identification**						**03511042**
					COC	GRA	
Lactobacillus fermentum		97.52	0.62	SOR 1	1	99	
Actinomyces naeslundii		1.30	0.40		1	99	
Megasphaera elsdenii		0.48	0.12		99	1	
03511043	**Low discrimination**						**03511043**
Actinomyces naeslundii		66.37	0.43	MNS 88			
Lactobacillus fermentum		33.63	0.28				
03511060	**Low discrimination**						**03511060**
					GRA		
Lactobacillus fermentum		84.13	0.54	SOR 1	99		
Prevotella melaninogenica		6.07	0.15		1		
Clostridium perfringens		5.70	0.22		99		
03511061	**Low discrimination**						**03511061**
Actinomyces naeslundii		85.55	0.43	MNS 88			
Lactobacillus fermentum		14.45	0.20				
03511062	**Acceptable identification**						**03511062**
Lactobacillus fermentum		92.13	0.54	SOR 1			
Actinomyces naeslundii		3.67	0.40				
Clostridium perfringens		2.94	0.16				
03511063	**Low discrimination**						**03511063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		85.55	0.43	MNS 88		
Lactobacillus fermentum		14.45	0.20			
03511100	**Good identification**					**03511100**
					GRA	
Lactobacillus fermentum		93.55	1.00		99	
Fusobacterium mortiferum		2.51	0.72		1	
Actinomyces meyeri		2.09	0.61		99	
03511101	**Low discrimination**					**03511101**
					SPO	
Lactobacillus fermentum		64.31	0.66	URE 1	1	
Clostridium chauvoei		18.23	0.44		99	
Actinomyces naeslundii		12.67	0.64		1	
03511102	**Very good identification**					**03511102**
					GRA	
Lactobacillus fermentum		98.81	1.00		99	
Fusobacterium mortiferum		0.30	0.55		1	
Prevotella melaninogenica		0.28	0.37		1	
03511103	**Low discrimination**					**03511103**
Lactobacillus fermentum		75.65	0.66	URE 1		
Actinomyces naeslundii		14.91	0.64			
Actinomyces viscosus		4.29	0.30			
03511110	**Low discrimination**					**03511110**
					GRA	
Lactobacillus fermentum		89.79	0.66	MLZ 1	99	
Bifidobacterium longum subsp. longum		2.89	0.33		99	
Fusobacterium mortiferum		2.41	0.37		1	
03511111	**Low discrimination**					**03511111**
					SPO	
Actinomyces naeslundii		44.45	0.42	MNS 88	1	
Lactobacillus fermentum		43.28	0.32	MLZ 5	1	
Clostridium chauvoei		12.27	0.09		99	
03511112	**Good identification**					**03511112**
Lactobacillus fermentum		93.42	0.66	MLZ 1		
Bifidobacterium longum subsp. longum		3.01	0.33			
Bifidobacterium adolescentis		1.33	0.29			
03511113	**Low discrimination**					**03511113**
Actinomyces naeslundii		50.66	0.42	MNS 88		
Lactobacillus fermentum		49.34	0.32	MLZ 5		
03511120	**Low discrimination**					**03511120**
					GRA	
Prevotella melaninogenica		80.87	0.84	MNS 90	1	
Lactobacillus fermentum		18.48	0.92		99	
Clostridium perfringens		0.30	0.49		99	
03511121	**Low discrimination**					**03511121**
					GRA	
Prevotella melaninogenica		69.70	0.50	MNS 90	1	
Lactobacillus fermentum		15.93	0.58	URE 1	99	
Actinomyces naeslundii		9.42	0.64		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03511122	**Low discrimination**						**03511122**
					GRA	CAT	
Lactobacillus fermentum		53.39	0.92		99	1	
Prevotella melaninogenica		44.50	0.71		1	1	
Capnocytophaga ochracea		0.95	0.47		1	90	
03511123	**Low discrimination**						**03511123**
					GRA		
Lactobacillus fermentum		39.85	0.58	URE 1	99		
Prevotella melaninogenica		33.21	0.37		1		
Actinomyces naeslundii		23.56	0.64		99		
03511130	**Low discrimination**						**03511130**
					GRA		
Prevotella melaninogenica		75.01	0.50	MNS 90	1		
Lactobacillus fermentum		17.14	0.58	MLZ 1	99		
Bifidobacterium longum subsp. longum		6.62	0.44		99		
03511132	**Low discrimination**						**03511132**
					GRA		
Lactobacillus fermentum		40.57	0.58	MLZ 1	99		
Prevotella melaninogenica		33.81	0.37		1		
Bifidobacterium longum subsp. longum		15.66	0.44		99		
03511133	**Low discrimination**						**03511133**
					GRA		
Actinomyces naeslundii		58.12	0.42	MNS 88	99		
Lactobacillus fermentum		18.87	0.24	MLZ 5	99		
Prevotella melaninogenica		15.73	0.03		1		
03511140	**Low discrimination**						**03511140**
					GRA		
Lactobacillus fermentum		85.66	0.66	SOR 1	99		
Propionibacterium propionicum		6.35	0.46		99		
Fusobacterium mortiferum		2.30	0.37		1		
03511141	**Low discrimination**						**03511141**
Actinomyces naeslundii		62.13	0.51	MNS 88			
Lactobacillus fermentum		18.05	0.32				
Propionibacterium propionicum		14.71	0.30				
03511142	**Low discrimination**						**03511142**
Lactobacillus fermentum		86.81	0.66	SOR 1			
Propionibacterium propionicum		9.65	0.49				
Actinomyces naeslundii		2.01	0.48				
03511143	**Low discrimination**						**03511143**
Actinomyces naeslundii		60.77	0.51	MNS 88			
Propionibacterium propionicum		21.58	0.33				
Lactobacillus fermentum		17.65	0.32				
03511160	**Low discrimination**						**03511160**
					GRA		
Prevotella melaninogenica		75.64	0.50	MNS 90	1		
Lactobacillus fermentum		17.29	0.58	SOR 1	99		
Clostridium perfringens		5.26	0.37		99		
03511161	**Low discrimination**						**03511161**
					GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		63.40	0.51	MNS 88	99		
Prevotella melaninogenica		26.86	0.15		1		
Lactobacillus fermentum		6.14	0.24		99		
03511162	**Low discrimination**						**03511162**
					GRA	CAT	
Lactobacillus fermentum		44.38	0.58	SOR 1	99	1	
Prevotella melaninogenica		36.98	0.37		1	1	
Capnocytophaga ochracea		8.64	0.30		1	90	
03511163	**Low discrimination**						**03511163**
					GRA		
Actinomyces naeslundii		84.93	0.51	MNS 88	99		
Lactobacillus fermentum		8.22	0.24		99		
Prevotella melaninogenica		6.85	0.03		1		
03511200	**Low discrimination**						**03511200**
					GRA		
Lactobacillus fermentum		79.42	0.62	RHA 1	99		
Prevotella intermedia		13.61	0.36		1		
Actinomyces odontolyticus		3.61	0.32		99		
03511202	**Very good identification**						**03511202**
					CAT		
Lactobacillus fermentum		96.45	0.62	RHA 1	1		
Actinomyces odontolyticus		2.68	0.28		10		
Propionibacterium granulosum		0.35	0.16		90		
03511220	**Low discrimination**						**03511220**
					GRA		
Lactobacillus fermentum		89.77	0.54	RHA 1	99		
Prevotella melaninogenica		6.47	0.15		1		
Actinomyces odontolyticus		1.99	0.19		99		
03511222	**Acceptable identification**						**03511222**
					GRA		
Lactobacillus fermentum		94.52	0.54	RHA 1	99		
Prevotella oralis		2.90	0.13		1		
Prevotella melaninogenica		1.30	0.03		1		
03511300	**Good identification**						**03511300**
					GRA		
Lactobacillus fermentum		92.84	0.66	RHA 1	99		
Fusobacterium mortiferum		2.49	0.37		1		
Actinomyces meyeri		2.07	0.27		99		
03511302	**Very good identification**						**03511302**
					GRA		
Lactobacillus fermentum		97.99	0.66	RHA 1	99		
Actinomyces odontolyticus		0.53	0.20		99		
Prevotella oralis		0.51	0.11		1		
03511320	**Low discrimination**						**03511320**
					GRA		
Prevotella melaninogenica		80.12	0.50	MNS 90	1		
Lactobacillus fermentum		18.31	0.58	RHA 1	99		
Bacteroides vulgatus		0.64	0.11		1		
03511322	**Low discrimination**						**03511322**
					GRA		
Lactobacillus fermentum		44.46	0.58	RHA 1	99		
Prevotella melaninogenica		37.05	0.37		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		13.11	0.33		1	
03511400	**Excellent identification**					**03511400**
					GRA	
Lactobacillus fermentum		99.54	0.96		99	
Actinomyces meyeri		0.23	0.40		99	
Prevotella intermedia		0.17	0.36		1	
03511401	**Very good identification**					**03511401**
					GRA	
Lactobacillus fermentum		99.60	0.62	URE 1	99	
Actinomyces meyeri		0.23	0.06		99	
Prevotella intermedia		0.17	0.02		1	
03511402	**Excellent identification**					**03511402**
Lactobacillus fermentum		99.96	0.96			
Bifidobacterium longum subsp. longum		0.03	0.29			
Actinomyces meyeri		0.01	0.15			
03511403	**Very good identification**					**03511403**
Lactobacillus fermentum		100.0	0.62	URE 1		
03511410	**Good identification**					**03511410**
Lactobacillus fermentum		94.65	0.62	MLZ 1		
Bifidobacterium longum subsp. longum		4.97	0.33			
Actinomyces meyeri		0.22	0.06			
03511412	**Good identification**					**03511412**
Lactobacillus fermentum		94.87	0.62	MLZ 1		
Bifidobacterium longum subsp. longum		4.98	0.33			
Bifidobacterium adolescentis		0.15	0.09			
03511420	**Very good identification**					**03511420**
Lactobacillus fermentum		99.46	0.88			
Bifidobacterium longum subsp. longum		0.37	0.40			
Actinomyces meyeri		0.09	0.26			
03511421	**Good identification**					**03511421**
Lactobacillus fermentum		99.63	0.54	URE 1		
Bifidobacterium longum subsp. longum		0.37	0.05			
03511422	**Very good identification**					**03511422**
					GRA	
Lactobacillus fermentum		99.61	0.88		99	
Bifidobacterium longum subsp. longum		0.37	0.40		99	
Prevotella melaninogenica		0.01	0.03		1	
03511423	**Good identification**					**03511423**
Lactobacillus fermentum		99.63	0.54	URE 1		
Bifidobacterium longum subsp. longum		0.37	0.05			
03511430	**Low discrimination**					**03511430**
Lactobacillus fermentum		61.35	0.54	MLZ 1		
Bifidobacterium longum subsp. longum		38.65	0.44			
03511432	**Low discrimination**					**03511432**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		61.35	0.54	MLZ 1		
Bifidobacterium longum subsp. longum		38.65	0.44			
03511440	**Very good identification**					**03511440**
					GRA	
Lactobacillus fermentum		99.60	0.62	SOR 1	99	
Actinomyces meyeri		0.23	0.06		99	
Prevotella intermedia		0.17	0.02		1	
03511442	**Very good identification**					**03511442**
Lactobacillus fermentum		100.0	0.62	SOR 1		
03511460	**Good identification**					**03511460**
Lactobacillus fermentum		99.63	0.54	SOR 1		
Bifidobacterium longum subsp. longum		0.37	0.05			
03511462	**Good identification**					**03511462**
Lactobacillus fermentum		99.63	0.54	SOR 1		
Bifidobacterium longum subsp. longum		0.37	0.05			
03511500	**Very good identification**					**03511500**
Lactobacillus fermentum		97.55	1.00			
Bifidobacterium longum subsp. longum		1.84	0.63			
Actinomyces meyeri		0.54	0.51			
03511501	**Very good identification**					**03511501**
Lactobacillus fermentum		97.15	0.66	URE 1		
Bifidobacterium longum subsp. longum		1.84	0.29			
Actinomyces meyeri		0.54	0.16			
03511502	**Very good identification**					**03511502**
Lactobacillus fermentum		98.03	1.00			
Bifidobacterium longum subsp. longum		1.85	0.63			
Bifidobacterium adolescentis		0.08	0.41			
03511503	**Very good identification**					**03511503**
Lactobacillus fermentum		97.88	0.66	URE 1		
Bifidobacterium longum subsp. longum		1.85	0.29			
Actinomyces naeslundii		0.20	0.30			
03511510	**Low discrimination**					**03511510**
Bifidobacterium longum subsp. longum		75.99	0.67	MNS 87		
Lactobacillus fermentum		23.86	0.66			
Actinomyces meyeri		0.13	0.16			
03511512	**Low discrimination**					**03511512**
Bifidobacterium longum subsp. longum		74.39	0.67	MNS 87		
Lactobacillus fermentum		23.35	0.66			
Bifidobacterium adolescentis		2.23	0.43			
03511520	**Low discrimination**					**03511520**
					GRA	
Lactobacillus fermentum		78.47	0.92		99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		17.79	0.74		99		
Prevotella melaninogenica		3.47	0.50		1		
03511521	**Low discrimination**						**03511521**
					GRA		
Lactobacillus fermentum		78.00	0.58	URE 1	99		
Bifidobacterium longum subsp. longum		17.69	0.40		99		
Prevotella melaninogenica		3.45	0.15		1		
03511522	**Low discrimination**						**03511522**
					GRA		
Lactobacillus fermentum		80.73	0.92		99		
Bifidobacterium longum subsp. longum		18.31	0.74		99		
Prevotella melaninogenica		0.68	0.37		1		
03511523	**Low discrimination**						**03511523**
					GRA		
Lactobacillus fermentum		80.37	0.58	URE 1	99		
Bifidobacterium longum subsp. longum		18.23	0.40		99		
Prevotella melaninogenica		0.68	0.03		1		
03511530	**Very good identification**						**03511530**
					GRA		
Bifidobacterium longum subsp. longum		97.33	0.78	MNS 87	99		
Lactobacillus fermentum		2.55	0.58		99		
Prevotella melaninogenica		0.11	0.15		1		
03511531	**Good identification**						**03511531**
Bifidobacterium longum subsp. longum		97.45	0.44	MNS 87			
Lactobacillus fermentum		2.55	0.24	URE 1			
03511532	**Very good identification**						**03511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.31	0.78	MNS 87	99	1	
Lactobacillus fermentum		2.55	0.58		99	1	
Capnocytophaga ochracea		0.09	0.17		1	90	
03511533	**Good identification**						**03511533**
Bifidobacterium longum subsp. longum		97.45	0.44	MNS 87			
Lactobacillus fermentum		2.55	0.24	URE 1			
03511540	**Very good identification**						**03511540**
Lactobacillus fermentum		97.54	0.66	SOR 1			
Bifidobacterium longum subsp. longum		1.84	0.29				
Actinomyces meyeri		0.54	0.16				
03511542	**Very good identification**						**03511542**
Lactobacillus fermentum		97.00	0.66	SOR 1			
Bifidobacterium longum subsp. longum		1.83	0.29				
Bifidobacterium adolescentis		0.94	0.26				
03511560	**Low discrimination**						**03511560**
					GRA		
Lactobacillus fermentum		78.12	0.58	SOR 1	99		
Bifidobacterium longum subsp. longum		17.71	0.40		99		
Prevotella melaninogenica		3.45	0.15		1		
03511562	**Low discrimination**						**03511562**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		78.77	0.58	SOR 1	99	1	
Bifidobacterium longum subsp. longum		17.86	0.40		99	1	
Capnocytophaga ochracea		2.71	0.17		1	90	
03511570	**Good identification**						**03511570**
Bifidobacterium longum subsp. longum		97.45	0.44	MNS 87			
Lactobacillus fermentum		2.55	0.24	SOR 1			
03511572	**Good identification**						**03511572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.52	0.44	MNS 87	99	1	
Lactobacillus fermentum		2.53	0.24	SOR 1	99	1	
Capnocytophaga ochracea		0.95	0.01		1	90	
03511600	**Very good identification**						**03511600**
					GRA		
Lactobacillus fermentum		99.60	0.62	RHA 1	99		
Actinomyces meyeri		0.23	0.06		99		
Prevotella intermedia		0.17	0.02		1		
03511602	**Very good identification**						**03511602**
Lactobacillus fermentum		100.0	0.62	RHA 1			
03511620	**Good identification**						**03511620**
					GRA		
Lactobacillus fermentum		98.88	0.54	RHA 1	99		
Bacteroides vulgatus		0.75	-.04		1		
Bifidobacterium longum subsp. longum		0.37	0.05		99		
03511622	**Good identification**						**03511622**
Lactobacillus fermentum		99.63	0.54	RHA 1			
Bifidobacterium longum subsp. longum		0.37	0.05				
03511700	**Very good identification**						**03511700**
Lactobacillus fermentum		97.46	0.66	RHA 1			
Bifidobacterium longum subsp. longum		1.84	0.29				
Actinomyces meyeri		0.54	0.16				
03511702	**Very good identification**						**03511702**
Lactobacillus fermentum		98.07	0.66	RHA 1			
Bifidobacterium longum subsp. longum		1.85	0.29				
Bifidobacterium adolescentis		0.08	0.07				
03511720	**Low discrimination**						**03511720**
					GRA		
Lactobacillus fermentum		57.67	0.58	RHA 1	99		
Bacteroides vulgatus		26.57	0.31		1		
Bifidobacterium longum subsp. longum		13.08	0.40		99		
03511722	**Low discrimination**						**03511722**
					GRA		
Lactobacillus fermentum		76.98	0.58	RHA 1	99		
Bifidobacterium longum subsp. longum		17.46	0.40		99		
Parabacteroides distasonis		4.09	0.33		1		
03511730	**Good identification**						**03511730**
					GRA		
Bifidobacterium longum subsp. longum		96.32	0.44	MNS 87	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		2.52	0.24	RHA 1	99			
Bacteroides vulgatus		1.16	-.04		1			
03511732	**Acceptable identification**							**03511732**
					GRA			
Bifidobacterium longum subsp. longum		90.89	0.44	MNS 87	99			
Parabacteroides distasonis		6.73	0.28	RHA 1	1			
Lactobacillus fermentum		2.38	0.24		99			
03512000	**Good identification**							**03512000**
					COC	GRA	CAT	
Megasphaera elsdenii		90.63	0.66	RAF 1	99	1	1	
Propionibacterium granulosum		4.84	0.50		1	99	90	
Prevotella intermedia		2.36	0.36		1	1	1	
03512002	**Low discrimination**							**03512002**
					COC	CAT	GRA	
Megasphaera elsdenii		71.52	0.50	RAF 1	99	1	1	
Propionibacterium propionicum		17.74	0.49	BGL 10	1	1	99	
Propionibacterium granulosum		8.60	0.39		1	90	99	
03512020	**Low discrimination**							**03512020**
					COC	GRA		
Propionibacterium granulosum		52.71	0.39	MNS 99	1	99		
Megasphaera elsdenii		39.85	0.32		99	1		
Actinomyces naeslundii		3.65	0.34		1	99		
03512021	**Good identification**							**03512021**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03512023	**Good identification**							**03512023**
Actinomyces naeslundii		100.0	0.37	MNS 88				
03512040	**Very good identification**							**03512040**
					COC	CAT	GRA	
Megasphaera elsdenii		97.20	0.63	RAF 1	99	1	1	
Propionibacterium propionicum		2.68	0.46		1	1	99	
Propionibacterium granulosum		0.08	0.16		1	90	99	
03512042	**Low discrimination**							**03512042**
					COC	GRA		
Megasphaera elsdenii		72.88	0.47	RAF 1	99	1		
Propionibacterium propionicum		27.12	0.49	BGL 10	1	99		
03512100	**Very good identification**							**03512100**
					COC	GRA		
Propionibacterium propionicum		98.69	0.81	NIT 86	1	99		
Megasphaera elsdenii		0.55	0.32		99	1		
Fusobacterium mortiferum		0.24	0.34		1	1		
03512101	**Very good identification**							**03512101**
Propionibacterium propionicum		98.08	0.64	NIT 86				
Actinomyces naeslundii		1.92	0.45	URE 10				
03512102	**Very good identification**							**03512102**
Propionibacterium propionicum		99.73	0.84	NIT 86				
Actinomyces naeslundii		0.10	0.42					
Lactobacillus fermentum		0.09	0.32					
03512103	**Very good identification**							**03512103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		98.64	0.67	NIT 86			
Actinomyces naeslundii		1.29	0.45	URE 10			
Actinomyces viscosus		0.08	0.00				
03512110	**Good identification**						**03512110**
Propionibacterium propionicum		100.0	0.46	NIT 86			
03512112	**Good identification**						**03512112**
Propionibacterium propionicum		100.0	0.49	NIT 86			
03512120	**Low discrimination**						**03512120**
					GRA		
Propionibacterium propionicum		70.67	0.46	NIT 86	99		
Prevotella melaninogenica		14.05	0.15	NAG 1	1		
Actinomyces naeslundii		10.14	0.42		99		
03512121	**Low discrimination**						**03512121**
Actinomyces naeslundii		65.94	0.45	MNS 88			
Propionibacterium propionicum		34.06	0.30	GAL 96			
03512122	**Low discrimination**						**03512122**
Propionibacterium propionicum		86.87	0.49	NIT 86			
Actinomyces naeslundii		8.31	0.42	NAG 1			
Lactobacillus fermentum		2.63	0.24				
03512123	**Low discrimination**						**03512123**
Actinomyces naeslundii		56.35	0.45	MNS 88			
Propionibacterium propionicum		43.65	0.33	GAL 96			
03512140	**Very good identification**						**03512140**
					COC	GRA	
Propionibacterium propionicum		99.61	0.81	NIT 86	1	99	
Megasphaera elsdenii		0.37	0.29		99	1	
Actinomyces naeslundii		0.03	0.29		1	99	
03512141	**Very good identification**						**03512141**
Propionibacterium propionicum		99.66	0.64	NIT 86			
Actinomyces naeslundii		0.34	0.32	URE 10			
03512142	**Very good identification**						**03512142**
					COC	GRA	
Propionibacterium propionicum		99.96	0.84	NIT 86	1	99	
Megasphaera elsdenii		0.03	0.12		99	1	
Actinomyces naeslundii		0.02	0.29		1	99	
03512143	**Very good identification**						**03512143**
Propionibacterium propionicum		99.77	0.67	NIT 86			
Actinomyces naeslundii		0.23	0.32	URE 10			
03512150	**Good identification**						**03512150**
Propionibacterium propionicum		100.0	0.46	NIT 86			
03512152	**Good identification**						**03512152**
Propionibacterium propionicum		100.0	0.49	NIT 86			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
03512160	**Good identification**						**03512160**
Propionibacterium propionicum		97.53	0.46	NIT 86			
Actinomyces naeslundii		2.47	0.29	NAG 1			
03512162	**Good identification**						**03512162**
Propionibacterium propionicum		98.34	0.49	NIT 86			
Actinomyces naeslundii		1.66	0.29	NAG 1			
03512300	**Good identification**						**03512300**
Propionibacterium propionicum		100.0	0.46	NIT 86			
03512302	**Good identification**						**03512302**
Propionibacterium propionicum		100.0	0.49	NIT 86			
03512340	**Good identification**						**03512340**
Propionibacterium propionicum		100.0	0.46	NIT 86			
03512342	**Good identification**						**03512342**
Propionibacterium propionicum		100.0	0.49	NIT 86			
03512500	**Low discrimination**						**03512500**
Propionibacterium propionicum		84.29	0.46	NIT 86			
Lactobacillus fermentum		11.49	0.32	ARA 1			
Actinomyces meyeri		4.23	0.13				
03512502	**Acceptable identification**						**03512502**
Propionibacterium propionicum		91.67	0.49	NIT 86			
Lactobacillus fermentum		8.33	0.32	ARA 1			
03512540	**Good identification**						**03512540**
Propionibacterium propionicum		100.0	0.46	NIT 86			
03512542	**Good identification**						**03512542**
Propionibacterium propionicum		100.0	0.49	NIT 86			
03513000	**Low discrimination**						**03513000**
					GRA	CAT	
Lactobacillus fermentum		53.07	0.62	MAN 1	99	1	
Propionibacterium granulosum		18.66	0.50		99	90	
Prevotella intermedia		9.10	0.36		1	1	
03513001	**Low discrimination**						**03513001**
Actinomyces naeslundii		88.39	0.61	MNS 88			
Propionibacterium propionicum		5.41	0.30				
Lactobacillus fermentum		4.07	0.28				
03513002	**Low discrimination**						**03513002**
Lactobacillus fermentum		70.02	0.62	MAN 1			
Propionibacterium propionicum		12.70	0.49				
Actinomyces naeslundii		10.24	0.58				
03513003	**Low discrimination**						**03513003**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		87.88	0.61	MNS 88			
Propionibacterium propionicum		8.07	0.33				
Lactobacillus fermentum		4.05	0.28				
03513011	**Good identification**						**03513011**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03513013	**Good identification**						**03513013**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03513020	**Low discrimination**						**03513020**
					CAT		
Lactobacillus fermentum		55.96	0.54	MAN 1	1		
Actinomyces naeslundii		24.55	0.58		25		
Propionibacterium granulosum		14.76	0.39		90		
03513021	**Very good identification**						**03513021**
Actinomyces naeslundii		98.49	0.61	MNS 88			
Lactobacillus fermentum		1.51	0.20				
03513022	**Low discrimination**						**03513022**
					CAT		
Lactobacillus fermentum		65.63	0.54	MAN 1	1		
Actinomyces naeslundii		28.79	0.58		25		
Propionibacterium granulosum		4.33	0.29		90		
03513023	**Very good identification**						**03513023**
Actinomyces naeslundii		98.49	0.61	MNS 88			
Lactobacillus fermentum		1.51	0.20				
03513031	**Good identification**						**03513031**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03513032	**Low discrimination**						**03513032**
					CAT		
Actinomyces naeslundii		54.74	0.36	MNS 88	25		
Lactobacillus fermentum		23.95	0.20	MLZ 5	1		
Propionibacterium granulosum		21.31	0.14		90		
03513033	**Good identification**						**03513033**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03513040	**Low discrimination**						**03513040**
					COC	GRA	
Propionibacterium propionicum		58.58	0.46	NIT 86	1	99	
Megasphaera elsdenii		21.47	0.29	LAC 99	99	1	
Actinomyces naeslundii		12.50	0.45		1	99	
03513041	**Low discrimination**						**03513041**
Actinomyces naeslundii		74.23	0.48	MNS 88			
Propionibacterium propionicum		25.77	0.30				
03513042	**Low discrimination**						**03513042**
Propionibacterium propionicum		81.63	0.49	NIT 86			
Actinomyces naeslundii		11.61	0.45	LAC 99			
Lactobacillus fermentum		4.55	0.28				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03513043	**Low discrimination**					**03513043**
Actinomyces naeslundii		65.76	0.48	MNS 88		
Propionibacterium propionicum		34.24	0.33			
03513060	**Low discrimination**					**03513060**
Actinomyces naeslundii		84.90	0.45	MNS 88		
Lactobacillus fermentum		11.08	0.20			
Propionibacterium propionicum		4.02	0.12			
03513061	**Good identification**					**03513061**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03513062	**Low discrimination**					**03513062**
Actinomyces naeslundii		83.23	0.45	MNS 88		
Lactobacillus fermentum		10.86	0.20			
Propionibacterium propionicum		5.91	0.15			
03513063	**Good identification**					**03513063**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03513100	**Low discrimination**					**03513100**
Propionibacterium propionicum		84.70	0.81	NIT 86		
Lactobacillus fermentum		11.55	0.66			
Actinomyces naeslundii		2.95	0.66			
03513101	**Low discrimination**					**03513101**
Propionibacterium propionicum		67.33	0.64	NIT 86		
Actinomyces naeslundii		31.60	0.69	URE 10		
Lactobacillus fermentum		0.83	0.32			
03513102	**Low discrimination**					**03513102**
Propionibacterium propionicum		89.62	0.84	NIT 86		
Lactobacillus fermentum		8.14	0.66			
Actinomyces naeslundii		2.08	0.66			
03513103	**Low discrimination**					**03513103**
Propionibacterium propionicum		75.69	0.67	NIT 86		
Actinomyces naeslundii		23.68	0.69	URE 10		
Lactobacillus fermentum		0.63	0.32			
03513110	**Low discrimination**					**03513110**
Propionibacterium propionicum		75.90	0.46	NIT 86		
Actinomyces naeslundii		13.75	0.44	MLZ 1		
Lactobacillus fermentum		10.35	0.32			
03513111	**Low discrimination**					**03513111**
Actinomyces naeslundii		70.98	0.47	MNS 88		
Propionibacterium propionicum		29.02	0.30	MLZ 5		
03513112	**Low discrimination**					**03513112**
Propionibacterium propionicum		75.96	0.49	NIT 86		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		9.17	0.44	MLZ 1		
Bifidobacterium adolescentis		7.97	0.27			
03513113	**Low discrimination**					**03513113**
Actinomyces naeslundii		61.98	0.47	MNS 88		
Propionibacterium propionicum		38.02	0.33	MLZ 5		
03513120	**Low discrimination**					**03513120**
					GRA	
Prevotella melaninogenica		68.47	0.50	MNS 90	1	
Lactobacillus fermentum		15.65	0.58	MAN 1	99	
Actinomyces naeslundii		11.97	0.66		99	
03513121	**Good identification**					**03513121**
					GRA	
Actinomyces naeslundii		93.34	0.69	MNS 88	99	
Prevotella melaninogenica		3.59	0.15		1	
Propionibacterium propionicum		2.01	0.30		99	
03513122	**Low discrimination**					**03513122**
					GRA	
Lactobacillus fermentum		33.71	0.58	MAN 1	99	
Prevotella melaninogenica		28.09	0.37		1	
Actinomyces naeslundii		25.79	0.66		99	
03513123	**Very good identification**					**03513123**
Actinomyces naeslundii		95.38	0.69	MNS 88		
Propionibacterium propionicum		3.08	0.33			
Lactobacillus fermentum		0.84	0.24			
03513131	**Good identification**					**03513131**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03513132	**Low discrimination**					**03513132**
					GRA	
Actinomyces naeslundii		60.96	0.44	MNS 88	99	
Lactobacillus fermentum		15.29	0.24	MLZ 5	99	
Prevotella melaninogenica		12.75	0.03		1	
03513133	**Good identification**					**03513133**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03513140	**Very good identification**					**03513140**
Propionibacterium propionicum		99.24	0.81	NIT 86		
Actinomyces naeslundii		0.61	0.53			
Lactobacillus fermentum		0.14	0.32			
03513141	**Good identification**					**03513141**
Propionibacterium propionicum		92.35	0.64	NIT 86		
Actinomyces naeslundii		7.65	0.56	URE 10		
03513142	**Very good identification**					**03513142**
Propionibacterium propionicum		99.48	0.84	NIT 86		
Actinomyces naeslundii		0.41	0.53			
Lactobacillus fermentum		0.09	0.32			
03513143	**Good identification**					**03513143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		94.77	0.67	NIT 86		
Actinomyces naeslundii		5.23	0.56	URE 10		
03513150	**Good identification**					**03513150**
Propionibacterium propionicum		96.90	0.46	NIT 86		
Actinomyces naeslundii		3.10	0.31	MLZ 1		
03513152	**Good identification**					**03513152**
Propionibacterium propionicum		96.69	0.49	NIT 86		
Actinomyces naeslundii		2.06	0.31	MLZ 1		
Bifidobacterium adolescentis		1.25	0.12			
03513160	**Low discrimination**					**03513160**
					GRA	
Propionibacterium propionicum		53.61	0.46	NIT 86	99	
Actinomyces naeslundii		32.56	0.53	NAG 1	99	
Prevotella melaninogenica		10.66	0.15		1	
03513161	**Low discrimination**					**03513161**
Actinomyces naeslundii		89.13	0.56	MNS 88		
Propionibacterium propionicum		10.87	0.30			
03513162	**Low discrimination**					**03513162**
Propionibacterium propionicum		68.47	0.49	NIT 86		
Actinomyces naeslundii		27.73	0.53	NAG 1		
Lactobacillus fermentum		2.07	0.24			
03513163	**Low discrimination**					**03513163**
Actinomyces naeslundii		84.54	0.56	MNS 88		
Propionibacterium propionicum		15.46	0.33			
03513300	**Low discrimination**					**03513300**
Propionibacterium propionicum		85.39	0.46	NIT 86		
Lactobacillus fermentum		11.64	0.32	RHA 1		
Actinomyces naeslundii		2.97	0.32			
03513302	**Low discrimination**					**03513302**
Propionibacterium propionicum		89.76	0.49	NIT 86		
Lactobacillus fermentum		8.16	0.32	RHA 1		
Actinomyces naeslundii		2.08	0.32			
03513340	**Good identification**					**03513340**
Propionibacterium propionicum		100.0	0.46	NIT 86		
03513342	**Good identification**					**03513342**
Propionibacterium propionicum		100.0	0.49	NIT 86		
03513400	**Very good identification**					**03513400**
					CAT	
Lactobacillus fermentum		98.99	0.62	MAN 1	1	
Propionibacterium granulosum		0.35	0.16		90	
Actinomyces meyeri		0.23	0.06		1	
03513402	**Very good identification**					**03513402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		99.67	0.62	MAN 1		
Propionibacterium propionicum		0.18	0.15			
Actinomyces naeslundii		0.15	0.24			
03513420	**Good identification**					**03513420**
Lactobacillus fermentum		99.19	0.54	MAN 1		
Actinomyces naeslundii		0.44	0.24			
Bifidobacterium longum subsp. longum		0.37	0.05			
03513422	**Good identification**					**03513422**
Lactobacillus fermentum		99.19	0.54	MAN 1		
Actinomyces naeslundii		0.44	0.24			
Bifidobacterium longum subsp. longum		0.37	0.05			
03513500	**Good identification**					**03513500**
Lactobacillus fermentum		90.68	0.66	MAN 1		
Propionibacterium propionicum		6.72	0.46			
Bifidobacterium longum subsp. longum		1.71	0.29			
03513502	**Low discrimination**					**03513502**
Lactobacillus fermentum		83.41	0.66	MAN 1		
Propionibacterium propionicum		9.27	0.49			
Bifidobacterium adolescentis		5.32	0.40			
03513512	**Low discrimination**					**03513512**
Bifidobacterium adolescentis		64.26	0.41	MNS 87		
Bifidobacterium longum subsp. longum		26.50	0.33	XYL 87		
Lactobacillus fermentum		8.32	0.32			
03513520	**Low discrimination**					**03513520**
					GRA	
Lactobacillus fermentum		77.93	0.58	MAN 1	99	
Bifidobacterium longum subsp. longum		17.67	0.40		99	
Prevotella melaninogenica		3.44	0.15		1	
03513522	**Low discrimination**					**03513522**
					GRA	
Lactobacillus fermentum		80.04	0.58	MAN 1	99	
Bifidobacterium longum subsp. longum		18.15	0.40		99	
Prevotella melaninogenica		0.67	0.03		1	
03513530	**Good identification**					**03513530**
Bifidobacterium longum subsp. longum		97.45	0.44	MNS 87		
Lactobacillus fermentum		2.55	0.24	MAN 1		
03513532	**Good identification**					**03513532**
Bifidobacterium longum subsp. longum		96.87	0.44	MNS 87		
Lactobacillus fermentum		2.53	0.24	MAN 1		
Bifidobacterium adolescentis		0.59	0.07			
03513540	**Low discrimination**					**03513540**
Propionibacterium propionicum		88.00	0.46	NIT 86		
Lactobacillus fermentum		12.00	0.32	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03513542	**Low discrimination**					**03513542**
Propionibacterium propionicum		85.45	0.49	NIT 86		
Lactobacillus fermentum		7.77	0.32	ARA 1		
Bifidobacterium adolescentis		6.07	0.24			
03514000	**Low discrimination**					**03514000**
					GRA	
Actinomyces meyeri		54.90	0.53	RAF 1	99	
Lactobacillus fermentum		27.69	0.58		99	
Prevotella intermedia		8.08	0.36		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03514002	**Low discrimination**					**03514002**
Lactobacillus fermentum		89.53	0.58	GAL 99		
Actinomyces meyeri		5.49	0.27			
Actinomyces odontolyticus		1.61	0.21			
03514020	**Low discrimination**					**03514020**
Lactobacillus fermentum		54.52	0.50	GAL 99		
Actinomyces meyeri		44.22	0.38			
Actinomyces odontolyticus		0.78	0.11			
03514022	**Good identification**					**03514022**
Lactobacillus fermentum		96.72	0.50	GAL 99		
Actinomyces meyeri		2.43	0.12			
Actinomyces odontolyticus		0.85	0.08			
03514100	**Low discrimination**					**03514100**
					GRA	
Actinomyces meyeri		79.73	0.63	RAF 1	99	
Lactobacillus fermentum		16.40	0.62		99	
Fusobacterium mortiferum		3.73	0.49		1	
03514102	**Low discrimination**					**03514102**
					GRA	
Lactobacillus fermentum		84.24	0.62	GAL 99	99	
Actinomyces meyeri		12.67	0.37		99	
Fusobacterium mortiferum		2.13	0.33		1	
03514120	**Low discrimination**					**03514120**
					GRA	
Actinomyces meyeri		64.52	0.48	RAF 1	99	
Lactobacillus fermentum		32.44	0.54	NAG 12	99	
Prevotella melaninogenica		2.44	0.15		1	
03514122	**Low discrimination**					**03514122**
					GRA	
Lactobacillus fermentum		89.78	0.54	GAL 99	99	
Actinomyces meyeri		5.52	0.22		99	
Prevotella melaninogenica		1.29	0.03		1	
03514400	**Low discrimination**					**03514400**
					GRA	
Lactobacillus fermentum		66.73	0.58	GAL 99	99	
Actinomyces meyeri		33.08	0.42		99	
Prevotella intermedia		0.20	0.02		1	
03514402	**Very good identification**					**03514402**
Lactobacillus fermentum		98.49	0.58	GAL 99		
Actinomyces meyeri		1.51	0.16			
03514420	**Low discrimination**					**03514420**
Lactobacillus fermentum		81.84	0.50	GAL 99		
Actinomyces meyeri		16.60	0.27			
Bifidobacterium longum subsp. longum		1.56	0.14			
03514422	**Good identification**					**03514422**
Lactobacillus fermentum		97.53	0.50	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		1.86	0.14			
Actinomyces meyeri		0.61	0.02			
03514500	**Low discrimination**					**03514500**
Actinomyces meyeri		52.52	0.53	RAF 1		
Lactobacillus fermentum		43.21	0.62			
Bifidobacterium longum subsp. longum		4.17	0.37			
03514502	**Low discrimination**					**03514502**
Lactobacillus fermentum		87.48	0.62	GAL 99		
Bifidobacterium longum subsp. longum		8.44	0.37			
Actinomyces meyeri		3.29	0.27			
03514510	**Low discrimination**					**03514510**
Bifidobacterium longum subsp. longum		88.02	0.41	MNS 87		
Actinomyces meyeri		6.57	0.18	GAL 99		
Lactobacillus fermentum		5.41	0.28			
03514512	**Low discrimination**					**03514512**
Bifidobacterium longum subsp. longum		88.63	0.41	MNS 87		
Bifidobacterium adolescentis		5.92	0.23	GAL 99		
Lactobacillus fermentum		5.45	0.28			
03514520	**Low discrimination**					**03514520**
Bifidobacterium longum subsp. longum		42.92	0.48	MNS 87		
Lactobacillus fermentum		37.06	0.54	GAL 99		
Actinomyces meyeri		18.42	0.38			
03514522	**Low discrimination**					**03514522**
Bifidobacterium longum subsp. longum		53.29	0.48	MNS 87		
Lactobacillus fermentum		46.00	0.54	GAL 99		
Actinomyces meyeri		0.71	0.12			
03514530	**Good identification**					**03514530**
Bifidobacterium longum subsp. longum		99.24	0.52	MNS 87		
Lactobacillus fermentum		0.51	0.20	GAL 99		
Actinomyces meyeri		0.25	0.04			
03514532	**Good identification**					**03514532**
Bifidobacterium longum subsp. longum		99.49	0.52	MNS 87		
Lactobacillus fermentum		0.51	0.20	GAL 99		
03515000	**Very good identification**					**03515000**
					GRA	
Lactobacillus fermentum		96.54	0.92		99	
Actinomyces meyeri		2.90	0.56		99	
Prevotella intermedia		0.29	0.36		1	
03515001	**Very good identification**					**03515001**
					GRA	
Lactobacillus fermentum		96.58	0.58	URE 1	99	
Actinomyces meyeri		2.90	0.21		99	
Prevotella intermedia		0.29	0.02		1	
03515002	**Excellent identification**					**03515002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Lactobacillus fermentum		99.86	0.92		99	
Actinomyces meyeri		0.09	0.30		99	
Fusobacterium mortiferum		0.03	0.30		1	
03515003	**Very good identification**					**03515003**
Lactobacillus fermentum		100.0	0.58	URE 1		
03515010	**Very good identification**					**03515010**
Lactobacillus fermentum		95.92	0.58	MLZ 1		
Actinomyces meyeri		2.88	0.21			
Eubacterium saburreum		0.42	0.15			
03515012	**Very good identification**					**03515012**
Lactobacillus fermentum		99.46	0.58	MLZ 1		
Bifidobacterium longum subsp. longum		0.27	0.07			
Bifidobacterium adolescentis		0.27	0.09			
03515020	**Very good identification**					**03515020**
					GRA	
Lactobacillus fermentum		98.62	0.84		99	
Actinomyces meyeri		1.21	0.41		99	
Prevotella melaninogenica		0.12	0.15		1	
03515021	**Good identification**					**03515021**
Lactobacillus fermentum		98.79	0.50	URE 1		
Actinomyces meyeri		1.21	0.07			
03515022	**Very good identification**					**03515022**
Lactobacillus fermentum		99.86	0.84			
Actinomyces meyeri		0.04	0.15			
Bifidobacterium longum subsp. longum		0.02	0.14			
03515023	**Good identification**					**03515023**
Lactobacillus fermentum		100.0	0.50	URE 1		
03515030	**Good identification**					**03515030**
Lactobacillus fermentum		95.71	0.50	MLZ 1		
Bifidobacterium longum subsp. longum		3.11	0.18			
Actinomyces meyeri		1.18	0.07			
03515032	**Good identification**					**03515032**
Lactobacillus fermentum		96.85	0.50	MLZ 1		
Bifidobacterium longum subsp. longum		3.15	0.18			
03515040	**Good identification**					**03515040**
					SPO	
Lactobacillus fermentum		94.23	0.58	SOR 1	1	
Actinomyces meyeri		2.83	0.21		1	
Clostridium glycolicum		2.44	0.15		99	
03515042	**Very good identification**					**03515042**
Lactobacillus fermentum		100.0	0.58	SOR 1		
03515060	**Good identification**					**03515060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		98.79	0.50	SOR 1			
Actinomyces meyeri		1.21	0.07				
03515062	**Good identification**						**03515062**
Lactobacillus fermentum		100.0	0.50	SOR 1			
03515100	**Good identification**						**03515100**
					GRA		
Lactobacillus fermentum		92.68	0.96		99		
Actinomyces meyeri		6.83	0.66		99		
Fusobacterium mortiferum		0.32	0.52		1		
03515101	**Good identification**						**03515101**
					SPO		
Lactobacillus fermentum		92.14	0.62	URE 1	1		
Actinomyces meyeri		6.79	0.32		1		
Clostridium chauvoei		0.45	0.09		99		
03515102	**Excellent identification**						**03515102**
Lactobacillus fermentum		99.44	0.96				
Actinomyces meyeri		0.23	0.40				
Bifidobacterium adolescentis		0.13	0.41				
03515103	**Very good identification**						**03515103**
Lactobacillus fermentum		99.30	0.62	URE 1			
Actinomyces naeslundii		0.34	0.30				
Actinomyces meyeri		0.23	0.06				
03515110	**Low discrimination**						**03515110**
Lactobacillus fermentum		80.04	0.62	MLZ 1			
Bifidobacterium longum subsp. longum		13.16	0.41				
Actinomyces meyeri		5.90	0.32				
03515112	**Low discrimination**						**03515112**
Lactobacillus fermentum		74.99	0.62	MLZ 1			
Bifidobacterium longum subsp. longum		12.33	0.41				
Bifidobacterium adolescentis		12.17	0.43				
03515120	**Low discrimination**						**03515120**
					GRA		
Lactobacillus fermentum		89.07	0.88		99		
Prevotella melaninogenica		6.70	0.50		1		
Actinomyces meyeri		2.68	0.51		99		
03515121	**Low discrimination**						**03515121**
					GRA		
Lactobacillus fermentum		88.33	0.54	URE 1	99		
Prevotella melaninogenica		6.65	0.15		1		
Actinomyces meyeri		2.66	0.17		99		
03515122	**Good identification**						**03515122**
					CAT	GRA	
Lactobacillus fermentum		94.79	0.88		1	99	
Prevotella melaninogenica		1.36	0.37		1	1	
Bacteroides fragilis		1.35	0.32		99	1	
03515123	**Acceptable identification**						**03515123**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT	GRA	
Lactobacillus fermentum		94.12	0.54	URE 1	1	99	
Prevotella melaninogenica		1.35	0.03		1	1	
Bacteroides fragilis		1.34	-.03		99	1	
03515130	**Low discrimination**						**03515130**
					GRA		
Bifidobacterium longum subsp. longum		63.89	0.52	MNS 87	99		
Lactobacillus fermentum		32.40	0.54	ARA 99	99		
Prevotella melaninogenica		2.44	0.15		1		
03515132	**Low discrimination**						**03515132**
					GRA		
Bifidobacterium longum subsp. longum		59.28	0.52	MNS 87	99		
Lactobacillus fermentum		30.06	0.54	ARA 99	99		
Parabacteroides distasonis		7.17	0.40		1		
03515140	**Good identification**						**03515140**
					GRA		
Lactobacillus fermentum		92.51	0.62	SOR 1	99		
Actinomyces meyeri		6.81	0.32		99		
Fusobacterium mortiferum		0.32	0.18		1		
03515142	**Very good identification**						**03515142**
Lactobacillus fermentum		97.31	0.62	SOR 1			
Bifidobacterium adolescentis		1.60	0.26				
Actinomyces israelii		0.38	0.22				
03515160	**Low discrimination**						**03515160**
					GRA		
Lactobacillus fermentum		88.38	0.54	SOR 1	99		
Prevotella melaninogenica		6.65	0.15		1		
Actinomyces meyeri		2.66	0.17		99		
03515162	**Low discrimination**						**03515162**
					GRA	CAT	
Lactobacillus fermentum		88.68	0.54	SOR 1	99	1	
Capnocytophaga ochracea		7.35	0.20		1	90	
Prevotella melaninogenica		1.27	0.03		1	1	
03515200	**Good identification**						**03515200**
Lactobacillus fermentum		93.16	0.58	RHA 1			
Actinomyces odontolyticus		3.55	0.27				
Actinomyces meyeri		2.80	0.21				
03515202	**Very good identification**						**03515202**
Lactobacillus fermentum		97.72	0.58	RHA 1			
Actinomyces odontolyticus		2.28	0.23				
03515220	**Acceptable identification**						**03515220**
					GRA		
Lactobacillus fermentum		92.72	0.50	RHA 1	99		
Bacteroides vulgatus		4.42	0.06		1		
Actinomyces odontolyticus		1.72	0.13		99		
03515222	**Good identification**						**03515222**
					GRA		
Lactobacillus fermentum		96.18	0.50	RHA 1	99		
Parabacteroides distasonis		2.11	0.19		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		1.10	0.10		99	
03515300	**Low discrimination**					**03515300**
Lactobacillus fermentum		86.49	0.62	RHA 1		
Actinomyces meyeri		6.37	0.32			
Eubacterium contortum		5.37	0.27			
03515302	**Very good identification**					**03515302**
					GRA	
Lactobacillus fermentum		97.70	0.62	RHA 1	99	
Eubacterium contortum		1.07	0.14		99	
Parabacteroides distasonis		0.44	0.19		1	
03515320	**Low discrimination**					**03515320**
					GRA	
Bacteroides vulgatus		71.32	0.40	MNS 99	1	
Lactobacillus fermentum		24.65	0.54	ARA 93	99	
Prevotella melaninogenica		1.86	0.15		1	
03515322	**Low discrimination**					**03515322**
					GRA	
Parabacteroides distasonis		53.70	0.53	MNS 98	1	
Lactobacillus fermentum		40.44	0.54		99	
Prevotella oralis		2.51	0.17		1	
03515332	**Good identification**					**03515332**
					GRA	
Parabacteroides distasonis		95.97	0.48	MNS 98	1	
Bifidobacterium longum subsp. longum		2.67	0.18		99	
Lactobacillus fermentum		1.36	0.20		99	
03515400	**Excellent identification**					**03515400**
Lactobacillus fermentum		99.09	0.92			
Actinomyces meyeri		0.74	0.45			
Bifidobacterium longum subsp. longum		0.16	0.37			
03515401	**Very good identification**					**03515401**
Lactobacillus fermentum		99.26	0.58	URE 1		
Actinomyces meyeri		0.75	0.11			
03515402	**Excellent identification**					**03515402**
Lactobacillus fermentum		99.80	0.92			
Bifidobacterium longum subsp. longum		0.16	0.37			
Actinomyces meyeri		0.02	0.19			
03515403	**Very good identification**					**03515403**
Lactobacillus fermentum		100.0	0.58	URE 1		
03515410	**Low discrimination**					**03515410**
Lactobacillus fermentum		78.39	0.58	MLZ 1		
Bifidobacterium longum subsp. longum		21.02	0.41			
Actinomyces meyeri		0.59	0.11			
03515412	**Low discrimination**					**03515412**
Lactobacillus fermentum		77.76	0.58	MLZ 1		
Bifidobacterium longum subsp. longum		20.85	0.41			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		1.39	0.23			
03515420	**Very good identification**					**03515420**
Lactobacillus fermentum		97.76	0.84			
Bifidobacterium longum subsp. longum		1.87	0.48			
Actinomyces meyeri		0.30	0.30			
03515421	**Good identification**					**03515421**
Lactobacillus fermentum		98.13	0.50	URE 1		
Bifidobacterium longum subsp. longum		1.87	0.14			
03515422	**Very good identification**					**03515422**
Lactobacillus fermentum		98.12	0.84			
Bifidobacterium longum subsp. longum		1.87	0.48			
Actinomyces meyeri		0.01	0.05			
03515423	**Good identification**					**03515423**
Lactobacillus fermentum		98.13	0.50	URE 1		
Bifidobacterium longum subsp. longum		1.87	0.14			
03515430	**Low discrimination**					**03515430**
Bifidobacterium longum subsp. longum		76.29	0.52	MNS 87		
Lactobacillus fermentum		23.71	0.50	LAC 99		
03515432	**Low discrimination**					**03515432**
Bifidobacterium longum subsp. longum		76.29	0.52	MNS 87		
Lactobacillus fermentum		23.71	0.50	LAC 99		
03515440	**Very good identification**					**03515440**
Lactobacillus fermentum		99.26	0.58	SOR 1		
Actinomyces meyeri		0.75	0.11			
03515442	**Very good identification**					**03515442**
Lactobacillus fermentum		100.0	0.58	SOR 1		
03515460	**Good identification**					**03515460**
Lactobacillus fermentum		98.13	0.50	SOR 1		
Bifidobacterium longum subsp. longum		1.87	0.14			
03515462	**Good identification**					**03515462**
Lactobacillus fermentum		98.13	0.50	SOR 1		
Bifidobacterium longum subsp. longum		1.87	0.14			
03515500	**Low discrimination**					**03515500**
Lactobacillus fermentum		89.66	0.96			
Bifidobacterium longum subsp. longum		8.66	0.72			
Actinomyces meyeri		1.65	0.56			
03515501	**Low discrimination**					**03515501**
Lactobacillus fermentum		89.69	0.62	URE 1		
Bifidobacterium longum subsp. longum		8.66	0.37			
Actinomyces meyeri		1.65	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03515502	**Good identification**					**03515502**
Lactobacillus fermentum		90.40	0.96			
Bifidobacterium longum subsp. longum		8.73	0.72			
Bifidobacterium adolescentis		0.81	0.56			
03515503	**Good identification**					**03515503**
Lactobacillus fermentum		90.46	0.62	URE 1		
Bifidobacterium longum subsp. longum		8.73	0.37			
Bifidobacterium adolescentis		0.81	0.21			
03515510	**Good identification**					**03515510**
Bifidobacterium longum subsp. longum		94.05	0.76	MNS 87		
Lactobacillus fermentum		5.78	0.62			
Actinomyces meyeri		0.11	0.21			
03515511	**Acceptable identification**					**03515511**
Bifidobacterium longum subsp. longum		94.21	0.41	MNS 87		
Lactobacillus fermentum		5.79	0.28	URE 1		
03515512	**Low discrimination**					**03515512**
Bifidobacterium longum subsp. longum		88.63	0.76	MNS 87		
Bifidobacterium adolescentis		5.92	0.57			
Lactobacillus fermentum		5.45	0.62			
03515513	**Low discrimination**					**03515513**
Bifidobacterium longum subsp. longum		88.63	0.41	MNS 87		
Bifidobacterium adolescentis		5.92	0.23	URE 1		
Lactobacillus fermentum		5.45	0.28			
03515520	**Low discrimination**					**03515520**
					GRA	
Bifidobacterium longum subsp. longum		52.42	0.82	MNS 87	99	
Lactobacillus fermentum		45.25	0.88		99	
Bacteroides vulgatus		1.95	0.43		1	
03515521	**Low discrimination**					**03515521**
					GRA	
Bifidobacterium longum subsp. longum		52.44	0.48	MNS 87	99	
Lactobacillus fermentum		45.27	0.54	URE 1	99	
Bacteroides vulgatus		1.95	0.09		1	
03515522	**Low discrimination**					**03515522**
					GRA	
Bifidobacterium longum subsp. longum		53.54	0.82	MNS 87	99	
Lactobacillus fermentum		46.22	0.88		99	
Parabacteroides distasonis		0.13	0.41		1	
03515523	**Low discrimination**					**03515523**
Bifidobacterium longum subsp. longum		53.67	0.48	MNS 87		
Lactobacillus fermentum		46.33	0.54	URE 1		
03515530	**Very good identification**					**03515530**
					GRA	
Bifidobacterium longum subsp. longum		99.47	0.86	MNS 87	99	
Lactobacillus fermentum		0.51	0.54		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		0.02	0.09		1		
03515531	**Good identification**						**03515531**
Bifidobacterium longum subsp. longum		99.49	0.52	MNS 87			
Lactobacillus fermentum		0.51	0.20	URE 1			
03515532	**Very good identification**						**03515532**
					GRA		
Bifidobacterium longum subsp. longum		99.39	0.86	MNS 87	99		
Lactobacillus fermentum		0.51	0.54		99		
Parabacteroides distasonis		0.07	0.37		1		
03515533	**Good identification**						**03515533**
Bifidobacterium longum subsp. longum		99.49	0.52	MNS 87			
Lactobacillus fermentum		0.51	0.20	URE 1			
03515540	**Low discrimination**						**03515540**
Lactobacillus fermentum		89.52	0.62	SOR 1			
Bifidobacterium longum subsp. longum		8.64	0.37				
Actinomyces meyeri		1.65	0.21				
03515542	**Low discrimination**						**03515542**
Lactobacillus fermentum		82.67	0.62	SOR 1			
Bifidobacterium adolescentis		9.08	0.40				
Bifidobacterium longum subsp. longum		7.98	0.37				
03515550	**Acceptable identification**						**03515550**
Bifidobacterium longum subsp. longum		93.48	0.41	MNS 87			
Lactobacillus fermentum		5.75	0.28	SOR 1			
Bifidobacterium adolescentis		0.77	0.07				
03515552	**Low discrimination**						**03515552**
Bifidobacterium longum subsp. longum		53.23	0.41	MNS 87			
Bifidobacterium adolescentis		43.50	0.42	SOR 1			
Lactobacillus fermentum		3.27	0.28				
03515560	**Low discrimination**						**03515560**
					GRA		
Bifidobacterium longum subsp. longum		52.44	0.48	MNS 87	99		
Lactobacillus fermentum		45.27	0.54	SOR 1	99		
Bacteroides vulgatus		1.95	0.09		1		
03515562	**Low discrimination**						**03515562**
					GRA	CAT	
Bifidobacterium longum subsp. longum		53.31	0.48	MNS 87	99	1	
Lactobacillus fermentum		46.02	0.54	SOR 1	99	1	
Capnocytophaga ochracea		0.67	0.07		1	90	
03515570	**Good identification**						**03515570**
Bifidobacterium longum subsp. longum		99.49	0.52	MNS 87			
Lactobacillus fermentum		0.51	0.20	SOR 1			
03515572	**Good identification**						**03515572**
Bifidobacterium longum subsp. longum		99.29	0.52	MNS 87			
Lactobacillus fermentum		0.51	0.20	SOR 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		0.21	0.07			
03515600	**Very good identification**					**03515600**
					GRA	
Lactobacillus fermentum		99.05	0.58	RHA 1	99	
Actinomyces meyeri		0.74	0.11		99	
Bacteroides vulgatus		0.21	-.09		1	
03515602	**Very good identification**					**03515602**
Lactobacillus fermentum		100.0	0.58	RHA 1		
03515620	**Low discrimination**					**03515620**
					GRA	
Lactobacillus fermentum		60.52	0.50	RHA 1	99	
Bacteroides vulgatus		38.33	0.25		1	
Bifidobacterium longum subsp. longum		1.16	0.14		99	
03515622	**Good identification**					**03515622**
					GRA	
Lactobacillus fermentum		96.26	0.50	RHA 1	99	
Bifidobacterium longum subsp. longum		1.84	0.14		99	
Parabacteroides distasonis		1.29	0.15		1	
03515700	**Low discrimination**					**03515700**
					GRA	
Lactobacillus fermentum		77.21	0.62	RHA 1	99	
Bacteroides vulgatus		9.99	0.25		1	
Bifidobacterium longum subsp. longum		7.45	0.37		99	
03515702	**Low discrimination**					**03515702**
Lactobacillus fermentum		89.25	0.62	RHA 1		
Bifidobacterium longum subsp. longum		8.62	0.37			
Bifidobacterium adolescentis		0.80	0.21			
03515710	**Acceptable identification**					**03515710**
					GRA	
Bifidobacterium longum subsp. longum		93.51	0.41	MNS 87	99	
Lactobacillus fermentum		5.75	0.28	RHA 1	99	
Bacteroides vulgatus		0.74	-.09		1	
03515712	**Low discrimination**					**03515712**
Bifidobacterium longum subsp. longum		83.91	0.41	MNS 87		
Bifidobacterium adolescentis		5.60	0.23	RHA 1		
Lactobacillus fermentum		5.16	0.28			
03515720	**Good identification**					**03515720**
					GRA	
Bacteroides vulgatus		94.58	0.60	MNS 99	1	
Bifidobacterium longum subsp. longum		2.85	0.48		99	
Lactobacillus fermentum		2.46	0.54		99	
03515722	**Low discrimination**					**03515722**
					GRA	
Bifidobacterium longum subsp. longum		34.36	0.48	MNS 87	99	
Lactobacillus fermentum		29.66	0.54	RHA 1	99	
Parabacteroides distasonis		24.15	0.49		1	
03515730	**Low discrimination**					**03515730**
					GRA	
Bifidobacterium longum subsp. longum		83.04	0.52	MNS 87	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		16.35	0.25	RHA 1	1	
Lactobacillus fermentum		0.43	0.20		99	
03515732	**Low discrimination**					**03515732**
					GRA	
Bifidobacterium longum subsp. longum		80.45	0.52	MNS 87	99	
Parabacteroides distasonis		17.88	0.45	RHA 1	1	
Clostridium clostridioforme		1.10	0.18		50	
03516100	**Low discrimination**					**03516100**
Propionibacterium propionicum		67.23	0.46	NIT 86		
Actinomyces meyeri		26.17	0.29	XYL 1		
Lactobacillus fermentum		5.38	0.28			
03516102	**Acceptable identification**					**03516102**
Propionibacterium propionicum		94.22	0.49	NIT 86		
Lactobacillus fermentum		5.03	0.28	XYL 1		
Actinomyces meyeri		0.76	0.03			
03516140	**Good identification**					**03516140**
Propionibacterium propionicum		100.0	0.46	NIT 86		
03516142	**Good identification**					**03516142**
Propionibacterium propionicum		100.0	0.49	NIT 86		
03517000	**Good identification**					**03517000**
Lactobacillus fermentum		94.19	0.58	MAN 1		
Actinomyces meyeri		2.83	0.21			
Actinomyces israelii		1.47	0.28			
03517002	**Very good identification**					**03517002**
Lactobacillus fermentum		97.17	0.58	MAN 1		
Actinomyces israelii		2.28	0.31			
Propionibacterium propionicum		0.30	0.15			
03517020	**Good identification**					**03517020**
Lactobacillus fermentum		98.06	0.50	MAN 1		
Actinomyces meyeri		1.21	0.07			
Actinomyces naeslundii		0.74	0.24			
03517022	**Good identification**					**03517022**
Lactobacillus fermentum		99.25	0.50	MAN 1		
Actinomyces naeslundii		0.75	0.24			
03517100	**Low discrimination**					**03517100**
Lactobacillus fermentum		80.26	0.62	MAN 1		
Propionibacterium propionicum		10.13	0.46			
Actinomyces meyeri		5.91	0.32			
03517102	**Low discrimination**					**03517102**
Lactobacillus fermentum		73.41	0.62	MAN 1		
Propionibacterium propionicum		13.89	0.49			
Bifidobacterium adolescentis		7.98	0.40			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
03517112	**Low discrimination**						**03517112**
Bifidobacterium adolescentis		85.64	0.41	MNS 87			
Lactobacillus fermentum		6.51	0.28	ARA 87			
Actinomyces israelii		5.54	0.27				
03517120	**Low discrimination**						**03517120**
					GRA		
Lactobacillus fermentum		87.61	0.54	MAN 1	99		
Prevotella melaninogenica		6.59	0.15		1		
Actinomyces meyeri		2.64	0.17		99		
03517122	**Acceptable identification**						**03517122**
					CAT	GRA	
Lactobacillus fermentum		92.41	0.54	MAN 1	1	99	
Prevotella melaninogenica		1.33	0.03		1	1	
Bacteroides fragilis		1.32	-.03		99	1	
03517140	**Low discrimination**						**03517140**
Propionibacterium propionicum		83.45	0.46	NIT 86			
Actinomyces israelii		9.87	0.32	XYL 1			
Lactobacillus fermentum		6.68	0.28				
03517142	**Low discrimination**						**03517142**
Propionibacterium propionicum		80.48	0.49	NIT 86			
Actinomyces israelii		9.52	0.35	XYL 1			
Bifidobacterium adolescentis		5.71	0.24				
03517400	**Very good identification**						**03517400**
Lactobacillus fermentum		98.01	0.58	MAN 1			
Actinomyces israelii		1.26	0.27				
Actinomyces meyeri		0.74	0.11				
03517402	**Very good identification**						**03517402**
Lactobacillus fermentum		96.98	0.58	MAN 1			
Actinomyces israelii		1.86	0.30				
Bifidobacterium adolescentis		1.16	0.20				
03517420	**Good identification**						**03517420**
Lactobacillus fermentum		98.13	0.50	MAN 1			
Bifidobacterium longum subsp. longum		1.87	0.14				
03517422	**Good identification**						**03517422**
Lactobacillus fermentum		98.13	0.50	MAN 1			
Bifidobacterium longum subsp. longum		1.87	0.14				
03517500	**Low discrimination**						**03517500**
Lactobacillus fermentum		86.58	0.62	MAN 1			
Bifidobacterium longum subsp. longum		8.36	0.37				
Actinomyces israelii		2.72	0.37				
03517502	**Low discrimination**						**03517502**
Lactobacillus fermentum		53.39	0.62	MAN 1			
Bifidobacterium adolescentis		38.84	0.54				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		5.15	0.37			
03517510	**Low discrimination**					**03517510**
Bifidobacterium longum subsp. longum		87.36	0.41	MNS 87		
Lactobacillus fermentum		5.37	0.28	MAN 1		
Bifidobacterium adolescentis		4.77	0.21			
03517512	**Low discrimination**					**03517512**
Bifidobacterium adolescentis		83.05	0.56	MNS 87		
Bifidobacterium longum subsp. longum		15.35	0.41			
Lactobacillus fermentum		0.94	0.28			
03517520	**Low discrimination**					**03517520**
					GRA	
Bifidobacterium longum subsp. longum		52.30	0.48	MNS 87	99	
Lactobacillus fermentum		45.15	0.54	MAN 1	99	
Bacteroides vulgatus		1.95	0.09		1	
03517522	**Low discrimination**					**03517522**
Bifidobacterium longum subsp. longum		52.91	0.48	MNS 87		
Lactobacillus fermentum		45.68	0.54	MAN 1		
Bifidobacterium adolescentis		1.01	0.20			
03517530	**Good identification**					**03517530**
Bifidobacterium longum subsp. longum		99.49	0.52	MNS 87		
Lactobacillus fermentum		0.51	0.20	MAN 1		
03517532	**Good identification**					**03517532**
Bifidobacterium longum subsp. longum		98.16	0.52	MNS 87		
Bifidobacterium adolescentis		1.34	0.21	MAN 1		
Lactobacillus fermentum		0.50	0.20			
03517542	**Low discrimination**					**03517542**
Bifidobacterium adolescentis		74.78	0.39	MNS 87		
Actinomyces israelii		15.23	0.33	SOR 11		
Lactobacillus fermentum		8.40	0.28			
03517552	**Good identification**					**03517552**
Bifidobacterium adolescentis		96.15	0.40	MNS 87		
Actinomyces israelii		2.39	0.19	SOR 11		
Bifidobacterium longum subsp. longum		1.45	0.07			
03521001	**Good identification**					**03521001**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03521003	**Good identification**					**03521003**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03521020	**Low discrimination**					**03521020**
					CAT	
Actinomyces naeslundii		39.49	0.36	MNS 88	25	
Propionibacterium granulosum		18.12	0.15	FRU 95	90	
Lactobacillus fermentum		17.28	0.20		1	
03521021	**Good identification**					**03521021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	MNS 88			
03521022	**Low discrimination**						**03521022**
Actinomyces naeslundii		69.56	0.36	MNS 88			
Lactobacillus fermentum		30.44	0.20	FRU 95			
03521023	**Good identification**						**03521023**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03521101	**Low discrimination**						**03521101**
					COC	SPO	
Actinomyces naeslundii		57.66	0.47	MNS 88	1	1	
Clostridium chauvoei		38.65	0.21	FRU 95	1	99	
Blautia hansenii		3.70	-.03		99	1	
03521103	**Low discrimination**						**03521103**
					SPO		
Actinomyces naeslundii		89.42	0.47	MNS 88	1		
Clostridium chauvoei		10.58	0.08	FRU 95	99		
03521121	**Low discrimination**						**03521121**
					GRA	SPO	
Actinomyces naeslundii		76.64	0.47	MNS 88	99	1	
Clostridium chauvoei		17.13	0.13	FRU 95	99	99	
Prevotella melaninogenica		6.23	-.01		1	1	
03521123	**Good identification**						**03521123**
					SPO		
Actinomyces naeslundii		96.21	0.47	MNS 88	1		
Clostridium chauvoei		3.79	0.00	FRU 95	99		
03521500	**Low discrimination**						**03521500**
					COC		
Bifidobacterium longum subsp. longum		61.42	0.37	MNS 87	1		
Blautia hansenii		26.65	0.19	FRU 90	99		
Lactobacillus fermentum		11.94	0.32		1		
03521502	**Low discrimination**						**03521502**
					COC		
Bifidobacterium longum subsp. longum		80.48	0.37	MNS 87	1		
Lactobacillus fermentum		15.64	0.32	FRU 90	1		
Blautia hansenii		3.88	0.03		99		
03521510	**Good identification**						**03521510**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87			
03521512	**Good identification**						**03521512**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87			
03521520	**Good identification**						**03521520**
					COC		
Bifidobacterium longum subsp. longum		96.51	0.47	MNS 87	1		
Lactobacillus fermentum		1.56	0.24	FRU 90	1		
Blautia hansenii		1.16	0.03		99		
03521522	**Good identification**						**03521522**
Bifidobacterium longum subsp. longum		98.41	0.47	MNS 87			
Lactobacillus fermentum		1.59	0.24	FRU 90			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03521530	**Good identification**					**03521530**
Bifidobacterium longum subsp. longum		100.0	0.51	MNS 87		
03521532	**Good identification**					**03521532**
Bifidobacterium longum subsp. longum		100.0	0.51	MNS 87		
03522100	**Good identification**					**03522100**
Propionibacterium propionicum		98.24	0.45	NIT 86		
Actinomyces naeslundii		1.76	0.25	FRU 99		
03522102	**Good identification**					**03522102**
Propionibacterium propionicum		98.82	0.48	NIT 86		
Actinomyces naeslundii		1.18	0.25	FRU 99		
03522140	**Good identification**					**03522140**
Propionibacterium propionicum		100.0	0.45	NIT 86		
03522142	**Good identification**					**03522142**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03523000	**Low discrimination**					**03523000**
Actinomyces naeslundii		58.37	0.41	MNS 88		
Propionibacterium granulosum		41.63	0.24	FRU 95		
03523001	**Good identification**					**03523001**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03523002	**Low discrimination**					**03523002**
					CAT	
Actinomyces naeslundii		78.21	0.41	MNS 88	25	
Propionibacterium granulosum		13.95	0.13	FRU 95	90	
Propionibacterium propionicum		7.85	0.14		1	
03523003	**Good identification**					**03523003**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03523020	**Low discrimination**					**03523020**
Actinomyces naeslundii		84.87	0.41	MNS 88		
Propionibacterium granulosum		15.13	0.13	FRU 95		
03523021	**Good identification**					**03523021**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03523022	**Good identification**					**03523022**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03523023	**Good identification**					**03523023**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03523100	**Low discrimination**					**03523100**
					COC	
Propionibacterium propionicum		68.91	0.45	NIT 86	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		29.62	0.49	FRU 99	1	
Blautia hansenii		1.47	-.03		99	
03523101	**Low discrimination**					**03523101**
Actinomyces naeslundii		85.30	0.52	MNS 88		
Propionibacterium propionicum		14.70	0.29	FRU 95		
03523102	**Low discrimination**					**03523102**
Propionibacterium propionicum		77.73	0.48	NIT 86		
Actinomyces naeslundii		22.27	0.49	FRU 99		
03523103	**Low discrimination**					**03523103**
Actinomyces naeslundii		79.46	0.52	MNS 88		
Propionibacterium propionicum		20.54	0.32	FRU 95		
03523120	**Acceptable identification**					**03523120**
					GRA	
Actinomyces naeslundii		94.09	0.49	MNS 88	99	
Prevotella melaninogenica		5.91	-.01	FRU 95	1	
03523121	**Good identification**					**03523121**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03523122	**Good identification**					**03523122**
Actinomyces naeslundii		96.60	0.49	MNS 88		
Propionibacterium propionicum		3.41	0.14	FRU 95		
03523123	**Good identification**					**03523123**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03523140	**Acceptable identification**					**03523140**
Propionibacterium propionicum		92.95	0.45	NIT 86		
Actinomyces naeslundii		7.05	0.36	FRU 99		
03523141	**Low discrimination**					**03523141**
Actinomyces naeslundii		50.59	0.39	MNS 88		
Propionibacterium propionicum		49.41	0.29	FRU 95		
03523142	**Good identification**					**03523142**
Propionibacterium propionicum		95.19	0.48	NIT 86		
Actinomyces naeslundii		4.81	0.36	FRU 99		
03523160	**Good identification**					**03523160**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03523161	**Good identification**					**03523161**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03523162	**Low discrimination**					**03523162**
Actinomyces naeslundii		83.35	0.36	MNS 88		
Propionibacterium propionicum		16.65	0.14	FRU 95		
03523163	**Good identification**					**03523163**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	MNS 88			
03525500	**Acceptable identification**						**03525500**
					COC		
Bifidobacterium longum subsp. longum		94.02	0.45	MNS 87	1		
Lactobacillus fermentum		3.58	0.28	FRU 90	1		
Blautia hansenii		2.40	0.06		99		
03525502	**Acceptable identification**						**03525502**
Bifidobacterium longum subsp. longum		94.29	0.45	MNS 87			
Lactobacillus fermentum		3.59	0.28	FRU 90			
Bifidobacterium adolescentis		2.13	0.18				
03525510	**Good identification**						**03525510**
Bifidobacterium longum subsp. longum		100.0	0.49	MNS 87			
03525512	**Good identification**						**03525512**
Bifidobacterium longum subsp. longum		98.41	0.49	MNS 87			
Bifidobacterium adolescentis		1.59	0.20	FRU 90			
03525520	**Good identification**						**03525520**
Bifidobacterium longum subsp. longum		99.68	0.55	MNS 87			
Lactobacillus fermentum		0.32	0.20	FRU 90			
03525522	**Good identification**						**03525522**
					GRA		
Bifidobacterium longum subsp. longum		99.30	0.55	MNS 87	99		
Parabacteroides distasonis		0.39	0.18	FRU 90	1		
Lactobacillus fermentum		0.32	0.20		99		
03525530	**Very good identification**						**03525530**
Bifidobacterium longum subsp. longum		100.0	0.59	MNS 87			
03525532	**Very good identification**						**03525532**
					GRA		
Bifidobacterium longum subsp. longum		99.88	0.59	MNS 87	99		
Parabacteroides distasonis		0.12	0.14	FRU 90	1		
03530000	**Low discrimination**						**03530000**
					GRA	CAT	
Propionibacterium granulosum		68.60	0.60	MNS 99	99	90	
Fusobacterium mortiferum		18.17	0.53		1	5	
Prevotella intermedia		8.41	0.36		1	1	
03530002	**Low discrimination**						**03530002**
					GRA	CAT	
Propionibacterium granulosum		80.05	0.50	MNS 99	99	90	
Fusobacterium mortiferum		9.42	0.37		1	5	
Actinomyces viscosus		4.39	0.19		99	76	
03530003	**Low discrimination**						**03530003**
Actinomyces naeslundii		47.94	0.37	MNS 88			
Actinomyces viscosus		33.54	0.11	GAL 96			
Propionibacterium granulosum		18.52	0.15				
03530010	**Good identification**						**03530010**
					GRA	CAT	
Propionibacterium granulosum		97.21	0.45	MNS 99	99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		1.91	0.19	MLZ 12	1	5	
Prevotella intermedia		0.88	0.02		1	1	
03530020	**Good identification**						**03530020**
					GRA	CAT	
Propionibacterium granulosum		95.32	0.50	MNS 99	99	90	
Actinomyces naeslundii		1.66	0.34		99	25	
Fusobacterium mortiferum		1.02	0.19		1	5	
03530021	**Low discrimination**						**03530021**
Actinomyces naeslundii		72.13	0.37	MNS 88			
Propionibacterium granulosum		27.87	0.15	GAL 96			
03530022	**Acceptable identification**						**03530022**
					CAT		
Propionibacterium granulosum		90.23	0.39	MNS 99	90		
Actinomyces naeslundii		6.29	0.34		25		
Lactobacillus fermentum		3.48	0.20		1		
03530023	**Good identification**						**03530023**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03530100	**Low discrimination**						**03530100**
					GRA		
Fusobacterium mortiferum		83.33	0.60	MLT 11	1		
Propionibacterium propionicum		9.54	0.45		99		
Actinomyces meyeri		2.34	0.24		99		
03530101	**Low discrimination**						**03530101**
					GRA		
Actinomyces naeslundii		56.88	0.45	MNS 88	99		
Propionibacterium propionicum		24.03	0.29	GAL 96	99		
Fusobacterium mortiferum		19.09	0.25		1		
03530102	**Low discrimination**						**03530102**
					GRA		
Propionibacterium propionicum		46.98	0.48	NIT 86	99		
Fusobacterium mortiferum		30.40	0.43	MAN 99	1		
Actinomyces viscosus		10.80	0.23		99		
03530103	**Low discrimination**						**03530103**
Actinomyces naeslundii		48.29	0.45	MNS 88			
Propionibacterium propionicum		30.61	0.32	GAL 96			
Actinomyces viscosus		21.10	0.15				
03530121	**Good identification**						**03530121**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03530122	**Low discrimination**						**03530122**
Actinomyces naeslundii		47.40	0.42	MNS 88			
Clostridium perfringens		20.97	0.14	GAL 96			
Lactobacillus fermentum		15.02	0.24				
03530123	**Good identification**						**03530123**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03530140	**Low discrimination**						**03530140**
					GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		87.96	0.45	NIT 86	99		
Fusobacterium mortiferum		7.76	0.25	MAN 99	1		
Actinomyces naeslundii		2.72	0.29		99		
03530142	**Good identification**						**03530142**
Propionibacterium propionicum		97.98	0.48	NIT 86			
Actinomyces naeslundii		2.02	0.29	MAN 99			
03531000	**Low discrimination**						**03531000**
					GRA	CAT	
Propionibacterium granulosum		42.48	0.60	MNS 99	99	90	
Lactobacillus fermentum		30.38	0.62		99	1	
Fusobacterium mortiferum		16.88	0.56		1	5	
03531001	**Low discrimination**						**03531001**
					CAT		
Actinomyces naeslundii		87.42	0.61	MNS 88	25		
Propionibacterium granulosum		5.63	0.26		90		
Lactobacillus fermentum		4.03	0.28		1		
03531002	**Low discrimination**						**03531002**
					CAT		
Lactobacillus fermentum		63.47	0.62	TRE 1	1		
Propionibacterium granulosum		22.19	0.50		90		
Actinomyces naeslundii		9.28	0.58		25		
03531003	**Good identification**						**03531003**
Actinomyces naeslundii		92.59	0.61	MNS 88			
Lactobacillus fermentum		4.27	0.28				
Actinomyces viscosus		1.65	0.07				
03531010	**Low discrimination**						**03531010**
					CAT		
Propionibacterium granulosum		87.06	0.45	MNS 99	90		
Lactobacillus fermentum		4.61	0.28	MLZ 12	1		
Actinomyces naeslundii		3.51	0.36		25		
03531011	**Low discrimination**						**03531011**
Actinomyces naeslundii		85.70	0.39	MNS 88			
Propionibacterium granulosum		14.30	0.11	MLZ 5			
03531013	**Good identification**						**03531013**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03531020	**Low discrimination**						**03531020**
					CAT		
Propionibacterium granulosum		32.31	0.50	MNS 99	90		
Lactobacillus fermentum		30.81	0.54		1		
Clostridium perfringens		20.35	0.39		1		
03531021	**Very good identification**						**03531021**
					CAT		
Actinomyces naeslundii		96.01	0.61	MNS 88	25		
Propionibacterium granulosum		1.55	0.15		90		
Lactobacillus fermentum		1.47	0.20		1		
03531022	**Low discrimination**						**03531022**
Lactobacillus fermentum		49.27	0.54	TRE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		21.61	0.58			
Clostridium perfringens		15.31	0.33			
03531023	**Very good identification**					**03531023**
Actinomyces naeslundii		98.49	0.61	MNS 88		
Lactobacillus fermentum		1.51	0.20			
03531031	**Good identification**					**03531031**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03531033	**Good identification**					**03531033**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03531040	**Low discrimination**					**03531040**
					CAT	
Actinomyces naeslundii		42.71	0.45	MNS 88	25	
Propionibacterium granulosum		23.38	0.26		90	
Lactobacillus fermentum		16.72	0.28		1	
03531041	**Good identification**					**03531041**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03531042	**Low discrimination**					**03531042**
					CAT	
Actinomyces naeslundii		63.04	0.45	MNS 88	25	
Lactobacillus fermentum		24.68	0.28		1	
Propionibacterium granulosum		8.63	0.15		90	
03531043	**Good identification**					**03531043**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03531061	**Good identification**					**03531061**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03531062	**Low discrimination**					**03531062**
Actinomyces naeslundii		52.77	0.45	MNS 88		
Clostridium perfringens		40.35	0.21			
Lactobacillus fermentum		6.89	0.20			
03531063	**Good identification**					**03531063**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03531100	**Low discrimination**					**03531100**
					GRA	
Lactobacillus fermentum		45.37	0.66	TRE 1	99	
Fusobacterium mortiferum		36.04	0.63		1	
Actinomyces naeslundii		11.57	0.66		99	
03531101	**Low discrimination**					**03531101**
					SPO	
Actinomyces naeslundii		87.16	0.69	MNS 88	1	
Clostridium chauvoei		7.18	0.27		99	
Lactobacillus fermentum		2.30	0.32		1	
03531102	**Low discrimination**					**03531102**
Lactobacillus fermentum		67.45	0.66	TRE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		17.20	0.66				
Propionibacterium propionicum		6.13	0.48				
03531103	**Good identification**						**03531103**
Actinomyces naeslundii		92.52	0.69	MNS 88			
Propionibacterium propionicum		2.44	0.32				
Lactobacillus fermentum		2.44	0.32				
03531110	**Low discrimination**						**03531110**
Actinomyces naeslundii		29.61	0.44	MNS 88			
Lactobacillus fermentum		22.28	0.32	MLZ 5			
Bifidobacterium longum subsp. longum		17.74	0.23				
03531111	**Good identification**						**03531111**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03531112	**Low discrimination**						**03531112**
Actinomyces naeslundii		30.73	0.44	MNS 88			
Lactobacillus fermentum		23.12	0.32	MLZ 5			
Bifidobacterium adolescentis		21.75	0.26				
03531113	**Good identification**						**03531113**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03531120	**Low discrimination**						**03531120**
					GRA		
Prevotella melaninogenica		47.51	0.50	MNS 90	1		
Clostridium perfringens		32.23	0.54	TRE 1	99		
Lactobacillus fermentum		10.86	0.58		99		
03531121	**Good identification**						**03531121**
					GRA	SPO	
Actinomyces naeslundii		90.84	0.69	MNS 88	99	1	
Prevotella melaninogenica		3.50	0.15		1	1	
Clostridium chauvoei		2.49	0.19		99	99	
03531122	**Low discrimination**						**03531122**
					GRA		
Clostridium perfringens		32.86	0.48	MNS 99	99		
Lactobacillus fermentum		23.53	0.58		99		
Prevotella melaninogenica		19.61	0.37		1		
03531123	**Very good identification**						**03531123**
Actinomyces naeslundii		96.78	0.69	MNS 88			
Clostridium perfringens		1.19	0.14				
Lactobacillus fermentum		0.85	0.24				
03531131	**Good identification**						**03531131**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03531133	**Good identification**						**03531133**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03531140	**Low discrimination**						**03531140**
Propionibacterium propionicum		46.78	0.45	NIT 86			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		34.73	0.53	MAN 99		
Lactobacillus fermentum		7.79	0.32			
03531141	**Acceptable identification**					**03531141**
Actinomyces naeslundii		90.93	0.56	MNS 88		
Propionibacterium propionicum		9.07	0.29			
03531142	**Low discrimination**					**03531142**
Propionibacterium propionicum		60.07	0.48	NIT 86		
Actinomyces naeslundii		29.73	0.53	MAN 99		
Lactobacillus fermentum		6.67	0.32			
03531143	**Low discrimination**					**03531143**
Actinomyces naeslundii		86.98	0.56	MNS 88		
Propionibacterium propionicum		13.02	0.32			
03531160	**Low discrimination**					**03531160**
					GRA	
Clostridium perfringens		74.91	0.41	MNS 99	99	
Actinomyces naeslundii		17.89	0.53		99	
Prevotella melaninogenica		5.86	0.15		1	
03531161	**Good identification**					**03531161**
Actinomyces naeslundii		97.26	0.56	MNS 88		
Clostridium perfringens		2.74	0.07			
03531162	**Low discrimination**					**03531162**
Clostridium perfringens		62.99	0.36	MNS 99		
Actinomyces naeslundii		31.97	0.53			
Lactobacillus fermentum		2.39	0.24			
03531163	**Good identification**					**03531163**
Actinomyces naeslundii		98.69	0.56	MNS 88		
Clostridium perfringens		1.31	0.01			
03531322	**Low discrimination**					**03531322**
					GRA	
Parabacteroides distasonis		86.97	0.48	MNS 98	1	
Prevotella oralis		4.15	0.13	XYL 90	1	
Clostridium perfringens		3.11	0.14		99	
03531332	**Good identification**					**03531332**
Parabacteroides distasonis		100.0	0.43	MNS 98		
03531400	**Very good identification**					**03531400**
					CAT	
Lactobacillus fermentum		96.81	0.62	TRE 1	1	
Propionibacterium granulosum		1.37	0.26		90	
Bifidobacterium longum subsp. longum		0.75	0.19		1	
03531402	**Very good identification**					**03531402**
					CAT	
Lactobacillus fermentum		98.74	0.62	TRE 1	1	
Bifidobacterium longum subsp. longum		0.76	0.19		1	
Propionibacterium granulosum		0.35	0.15		90	
03531420	**Low discrimination**					**03531420**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Lactobacillus fermentum		89.75	0.54	TRE 1	1	
Bifidobacterium longum subsp. longum		8.30	0.29		1	
Propionibacterium granulosum		0.95	0.15		90	
03531422	**Acceptable identification**					**03531422**
Lactobacillus fermentum		91.16	0.54	TRE 1		
Bifidobacterium longum subsp. longum		8.43	0.29			
Actinomyces naeslundii		0.40	0.24			
03531500	**Low discrimination**					**03531500**
					GRA	
Lactobacillus fermentum		67.34	0.66	TRE 1	99	
Bifidobacterium longum subsp. longum		31.50	0.53		99	
Fusobacterium mortiferum		0.54	0.28		1	
03531502	**Low discrimination**					**03531502**
Lactobacillus fermentum		65.54	0.66	TRE 1		
Bifidobacterium longum subsp. longum		30.66	0.53			
Bifidobacterium adolescentis		3.41	0.38			
03531510	**Very good identification**					**03531510**
Bifidobacterium longum subsp. longum		98.75	0.57	MNS 87		
Lactobacillus fermentum		1.25	0.32			
03531512	**Good identification**					**03531512**
Bifidobacterium longum subsp. longum		91.53	0.57	MNS 87		
Bifidobacterium adolescentis		7.31	0.40			
Lactobacillus fermentum		1.16	0.32			
03531520	**Low discrimination**					**03531520**
					GRA	
Bifidobacterium longum subsp. longum		83.81	0.63	MNS 87	99	
Lactobacillus fermentum		14.93	0.58		99	
Prevotella melaninogenica		0.66	0.15		1	
03531522	**Low discrimination**					**03531522**
					GRA	
Bifidobacterium longum subsp. longum		83.44	0.63	MNS 87	99	
Lactobacillus fermentum		14.87	0.58		99	
Parabacteroides distasonis		1.20	0.36		1	
03531530	**Very good identification**					**03531530**
Bifidobacterium longum subsp. longum		99.89	0.67	MNS 87		
Lactobacillus fermentum		0.11	0.24			
03531532	**Very good identification**					**03531532**
					GRA	
Bifidobacterium longum subsp. longum		99.44	0.67	MNS 87	99	
Parabacteroides distasonis		0.45	0.31		1	
Lactobacillus fermentum		0.11	0.24		99	
03531722	**Low discrimination**					**03531722**
					GRA	
Parabacteroides distasonis		77.74	0.44	MNS 98	1	
Bifidobacterium longum subsp. longum		18.21	0.29	XYL 90	99	
Lactobacillus fermentum		3.25	0.24		99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
03531732	**Low discrimination**							**03531732**
					GRA			
Parabacteroides distasonis		54.68	0.40	MNS 98	1			
Bifidobacterium longum subsp. longum		40.51	0.33	XYL 90	99			
Clostridium clostridioforme		4.81	0.16		50			
03532000	**Low discrimination**							**03532000**
					COC	CAT	GRA	
Propionibacterium granulosum		85.24	0.58	MNS 99	1	90	99	
Propionibacterium propionicum		7.99	0.45		1	1	99	
Megasphaera elsdenii		5.37	0.32		99	1	1	
03532001	**Low discrimination**							**03532001**
					CAT			
Actinomyces naeslundii		45.04	0.42	MNS 88	25			
Propionibacterium propionicum		27.91	0.29	GAL 96	1			
Propionibacterium granulosum		27.05	0.24		90			
03532002	**Low discrimination**							**03532002**
					CAT			
Propionibacterium granulosum		61.14	0.48	MNS 99	90			
Propionibacterium propionicum		34.40	0.48		1			
Actinomyces naeslundii		2.74	0.39		25			
03532003	**Low discrimination**							**03532003**
					CAT			
Actinomyces naeslundii		48.09	0.42	MNS 88	25			
Propionibacterium propionicum		44.69	0.32	GAL 96	1			
Propionibacterium granulosum		7.22	0.13		90			
03532010	**Good identification**							**03532010**
Propionibacterium granulosum		100.0	0.43	MNS 99				
03532020	**Good identification**							**03532020**
Propionibacterium granulosum		95.71	0.48	MNS 99				
Actinomyces naeslundii		4.29	0.39					
03532021	**Low discrimination**							**03532021**
Actinomyces naeslundii		86.95	0.42	MNS 88				
Propionibacterium granulosum		13.05	0.13	GAL 96				
03532022	**Low discrimination**							**03532022**
					CAT			
Propionibacterium granulosum		83.19	0.37	MNS 99	90			
Actinomyces naeslundii		14.92	0.39		25			
Propionibacterium propionicum		1.89	0.14		1			
03532023	**Good identification**							**03532023**
Actinomyces naeslundii		100.0	0.42	MNS 88				
03532040	**Low discrimination**							**03532040**
					COC	CAT	GRA	
Propionibacterium propionicum		62.72	0.45	NIT 86	1	1	99	
Megasphaera elsdenii		28.09	0.29	LAC 99	99	1	1	
Propionibacterium granulosum		6.76	0.24		1	90	99	
03532042	**Acceptable identification**							**03532042**
					COC	CAT	GRA	
Propionibacterium propionicum		93.88	0.48	NIT 86	1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		3.12	0.12	LAC 99	99	1	1	
Propionibacterium granulosum		1.69	0.13		1	90	99	
03532100	**Very good identification**							**03532100**
					CAT			
Propionibacterium propionicum		99.46	0.79	NIT 86	1			
Actinomyces naeslundii		0.34	0.47		25			
Propionibacterium granulosum		0.11	0.24		90			
03532101	**Very good identification**							**03532101**
Propionibacterium propionicum		95.57	0.63	NIT 86				
Actinomyces naeslundii		4.43	0.50	URE 10				
03532102	**Very good identification**							**03532102**
					CAT			
Propionibacterium propionicum		99.75	0.82	NIT 86	1			
Actinomyces naeslundii		0.23	0.47		25			
Propionibacterium granulosum		0.02	0.13		90			
03532103	**Very good identification**							**03532103**
Propionibacterium propionicum		97.00	0.66	NIT 86				
Actinomyces naeslundii		3.00	0.50	URE 10				
03532110	**Good identification**							**03532110**
					CAT			
Propionibacterium propionicum		96.84	0.45	NIT 86	1			
Actinomyces naeslundii		1.73	0.25	MLZ 1	25			
Propionibacterium granulosum		1.42	0.09		90			
03532112	**Good identification**							**03532112**
Propionibacterium propionicum		98.82	0.48	NIT 86				
Actinomyces naeslundii		1.18	0.25	MLZ 1				
03532120	**Low discrimination**							**03532120**
					CAT			
Propionibacterium propionicum		73.14	0.45	NIT 86	1			
Actinomyces naeslundii		24.89	0.47	NAG 1	25			
Propionibacterium granulosum		1.97	0.13		90			
03532121	**Low discrimination**							**03532121**
Actinomyces naeslundii		82.12	0.50	MNS 88				
Propionibacterium propionicum		17.88	0.29	GAL 96				
03532122	**Low discrimination**							**03532122**
Propionibacterium propionicum		81.51	0.48	NIT 86				
Actinomyces naeslundii		18.49	0.47	NAG 1				
03532123	**Low discrimination**							**03532123**
Actinomyces naeslundii		75.38	0.50	MNS 88				
Propionibacterium propionicum		24.62	0.32	GAL 96				
03532140	**Very good identification**							**03532140**
Propionibacterium propionicum		99.94	0.79	NIT 86				
Actinomyces naeslundii		0.06	0.34					
03532141	**Very good identification**							**03532141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.19	0.63	NIT 86		
Actinomyces naeslundii		0.81	0.37	URE 10		
03532142	**Very good identification**					**03532142**
Propionibacterium propionicum		99.96	0.82	NIT 86		
Actinomyces naeslundii		0.04	0.34			
03532143	**Very good identification**					**03532143**
Propionibacterium propionicum		99.46	0.66	NIT 86		
Actinomyces naeslundii		0.54	0.37	URE 10		
03532150	**Good identification**					**03532150**
Propionibacterium propionicum		100.0	0.45	NIT 86		
03532152	**Good identification**					**03532152**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03532160	**Acceptable identification**					**03532160**
Propionibacterium propionicum		94.34	0.45	NIT 86		
Actinomyces naeslundii		5.67	0.34	NAG 1		
03532162	**Good identification**					**03532162**
Propionibacterium propionicum		96.15	0.48	NIT 86		
Actinomyces naeslundii		3.85	0.34	NAG 1		
03532300	**Good identification**					**03532300**
Propionibacterium propionicum		100.0	0.45	NIT 86		
03532302	**Good identification**					**03532302**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03532340	**Good identification**					**03532340**
Propionibacterium propionicum		100.0	0.45	NIT 86		
03532342	**Good identification**					**03532342**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03532500	**Good identification**					**03532500**
Propionibacterium propionicum		100.0	0.45	NIT 86		
03532502	**Good identification**					**03532502**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03532540	**Good identification**					**03532540**
Propionibacterium propionicum		100.0	0.45	NIT 86		
03532542	**Good identification**					**03532542**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03533000	**Low discrimination**					**03533000**
Propionibacterium granulosum		72.35	0.58	MNS 99	CAT 90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		19.47	0.63		25	
Propionibacterium propionicum		6.78	0.45		1	
03533001	**Very good identification**					**03533001**
					CAT	
Actinomyces naeslundii		95.16	0.66	MNS 88	25	
Propionibacterium propionicum		2.46	0.29		1	
Propionibacterium granulosum		2.38	0.24		90	
03533002	**Low discrimination**					**03533002**
					CAT	
Actinomyces naeslundii		39.95	0.63	MNS 88	25	
Propionibacterium granulosum		37.12	0.48		90	
Propionibacterium propionicum		20.88	0.48		1	
03533003	**Very good identification**					**03533003**
					CAT	
Actinomyces naeslundii		95.70	0.66	MNS 88	25	
Propionibacterium propionicum		3.71	0.32		1	
Propionibacterium granulosum		0.60	0.13		90	
03533010	**Acceptable identification**					**03533010**
Propionibacterium granulosum		90.59	0.43	MNS 99		
Actinomyces naeslundii		9.41	0.41	MLZ 12		
03533011	**Acceptable identification**					**03533011**
Actinomyces naeslundii		93.91	0.44	MNS 88		
Propionibacterium granulosum		6.09	0.09	MLZ 5		
03533013	**Good identification**					**03533013**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03533020	**Low discrimination**					**03533020**
					CAT	
Actinomyces naeslundii		51.31	0.63	MNS 88	25	
Propionibacterium granulosum		47.68	0.48		90	
Lactobacillus fermentum		0.61	0.20		1	
03533021	**Very good identification**					**03533021**
Actinomyces naeslundii		99.38	0.66	MNS 88		
Propionibacterium granulosum		0.62	0.13			
03533022	**Low discrimination**					**03533022**
					CAT	
Actinomyces naeslundii		80.04	0.63	MNS 88	25	
Propionibacterium granulosum		18.59	0.37		90	
Lactobacillus fermentum		0.95	0.20		1	
03533023	**Very good identification**					**03533023**
Actinomyces naeslundii		100.0	0.66	MNS 88		
03533031	**Good identification**					**03533031**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03533032	**Low discrimination**					**03533032**
Actinomyces naeslundii		62.43	0.41	MNS 88		
Propionibacterium granulosum		37.57	0.22	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03533033	**Good identification**					**03533033**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03533040	**Low discrimination**					**03533040**
					CAT	
Propionibacterium propionicum		61.96	0.45	NIT 86	1	
Actinomyces naeslundii		31.37	0.50	LAC 99	25	
Propionibacterium granulosum		6.67	0.24		90	
03533041	**Low discrimination**					**03533041**
Actinomyces naeslundii		87.24	0.53	MNS 88		
Propionibacterium propionicum		12.76	0.29			
03533042	**Low discrimination**					**03533042**
					CAT	
Propionibacterium propionicum		73.14	0.48	NIT 86	1	
Actinomyces naeslundii		24.69	0.50	LAC 99	25	
Propionibacterium granulosum		1.31	0.13		90	
03533043	**Low discrimination**					**03533043**
Actinomyces naeslundii		82.01	0.53	MNS 88		
Propionibacterium propionicum		18.00	0.32			
03533060	**Acceptable identification**					**03533060**
Actinomyces naeslundii		94.95	0.50	MNS 88		
Propionibacterium granulosum		5.05	0.13			
03533061	**Good identification**					**03533061**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03533062	**Good identification**					**03533062**
Actinomyces naeslundii		97.10	0.50	MNS 88		
Propionibacterium propionicum		2.91	0.14			
03533063	**Good identification**					**03533063**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03533100	**Good identification**					**03533100**
Propionibacterium propionicum		91.93	0.79	NIT 86		
Actinomyces naeslundii		7.58	0.71			
Lactobacillus fermentum		0.16	0.32			
03533101	**Low discrimination**					**03533101**
Actinomyces naeslundii		52.69	0.74	MNS 88		
Propionibacterium propionicum		47.31	0.63			
03533102	**Good identification**					**03533102**
Propionibacterium propionicum		94.49	0.82	NIT 86		
Actinomyces naeslundii		5.20	0.71			
Actinomyces israelii		0.12	0.33			
03533103	**Low discrimination**					**03533103**
Propionibacterium propionicum		57.39	0.66	NIT 86		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		42.61	0.74	URE 10		
03533110	**Low discrimination**					**03533110**
Propionibacterium propionicum		68.37	0.45	NIT 86		
Actinomyces naeslundii		29.39	0.49	MLZ 1		
Actinomyces israelii		1.24	0.16			
03533111	**Low discrimination**					**03533111**
Actinomyces naeslundii		85.30	0.52	MNS 88		
Propionibacterium propionicum		14.70	0.29	MLZ 5		
03533112	**Low discrimination**					**03533112**
Propionibacterium propionicum		71.93	0.48	NIT 86		
Actinomyces naeslundii		20.61	0.49	MLZ 1		
Bifidobacterium adolescentis		6.15	0.24			
03533113	**Low discrimination**					**03533113**
Actinomyces naeslundii		79.46	0.52	MNS 88		
Propionibacterium propionicum		20.54	0.32	MLZ 5		
03533120	**Low discrimination**					**03533120**
					GRA	
Actinomyces naeslundii		84.57	0.71	MNS 88	99	
Propionibacterium propionicum		10.36	0.45		99	
Prevotella melaninogenica		2.52	0.15		1	
03533121	**Very good identification**					**03533121**
Actinomyces naeslundii		99.10	0.74	MNS 88		
Propionibacterium propionicum		0.90	0.29			
03533122	**Low discrimination**					**03533122**
Actinomyces naeslundii		82.94	0.71	MNS 88		
Propionibacterium propionicum		15.24	0.48			
Clostridium perfringens		0.79	0.14			
03533123	**Very good identification**					**03533123**
Actinomyces naeslundii		98.66	0.74	MNS 88		
Propionibacterium propionicum		1.34	0.32			
03533130	**Good identification**					**03533130**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03533131	**Good identification**					**03533131**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03533132	**Good identification**					**03533132**
Actinomyces naeslundii		96.60	0.49	MNS 88		
Propionibacterium propionicum		3.41	0.14	MLZ 5		
03533133	**Good identification**					**03533133**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03533140	**Very good identification**					**03533140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		98.52	0.79	NIT 86		
Actinomyces naeslundii		1.43	0.58			
Actinomyces israelii		0.05	0.23			
03533141	**Low discrimination**					**03533141**
Propionibacterium propionicum		83.58	0.63	NIT 86		
Actinomyces naeslundii		16.42	0.61	URE 10		
03533142	**Very good identification**					**03533142**
Propionibacterium propionicum		98.98	0.82	NIT 86		
Actinomyces naeslundii		0.96	0.58			
Actinomyces israelii		0.05	0.26			
03533143	**Low discrimination**					**03533143**
Propionibacterium propionicum		88.42	0.66	NIT 86		
Actinomyces naeslundii		11.58	0.61	URE 10		
03533150	**Acceptable identification**					**03533150**
Propionibacterium propionicum		92.95	0.45	NIT 86		
Actinomyces naeslundii		7.05	0.36	MLZ 1		
03533151	**Low discrimination**					**03533151**
Actinomyces naeslundii		50.59	0.39	MNS 88		
Propionibacterium propionicum		49.41	0.29	MLZ 5		
03533152	**Acceptable identification**					**03533152**
Propionibacterium propionicum		94.24	0.48	NIT 86		
Actinomyces naeslundii		4.77	0.36	MLZ 1		
Bifidobacterium adolescentis		1.00	0.09			
03533160	**Low discrimination**					**03533160**
Actinomyces naeslundii		58.29	0.58	MNS 88		
Propionibacterium propionicum		40.45	0.45			
Clostridium perfringens		1.27	0.07			
03533161	**Very good identification**					**03533161**
Actinomyces naeslundii		95.11	0.61	MNS 88		
Propionibacterium propionicum		4.89	0.29			
03533162	**Low discrimination**					**03533162**
Propionibacterium propionicum		50.75	0.48	NIT 86		
Actinomyces naeslundii		48.75	0.58	NAG 1		
Clostridium perfringens		0.50	0.01			
03533163	**Good identification**					**03533163**
Actinomyces naeslundii		92.84	0.61	MNS 88		
Propionibacterium propionicum		7.16	0.32			
03533170	**Good identification**					**03533170**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03533171	**Good identification**					**03533171**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	MNS 88		
03533172	**Low discrimination**					**03533172**
Actinomyces naeslundii		83.35	0.36	MNS 88		
Propionibacterium propionicum		16.65	0.14	MLZ 5		
03533173	**Good identification**					**03533173**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03533300	**Acceptable identification**					**03533300**
Propionibacterium propionicum		92.38	0.45	NIT 86		
Actinomyces naeslundii		7.62	0.37	RHA 1		
03533301	**Low discrimination**					**03533301**
Actinomyces naeslundii		52.69	0.40	MNS 88		
Propionibacterium propionicum		47.31	0.29	RHA 1		
03533302	**Acceptable identification**					**03533302**
Propionibacterium propionicum		93.97	0.48	NIT 86		
Actinomyces naeslundii		5.17	0.37	RHA 1		
Actinomyces israelii		0.86	0.14			
03533320	**Good identification**					**03533320**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03533321	**Good identification**					**03533321**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03533322	**Low discrimination**					**03533322**
					GRA	
Actinomyces naeslundii		68.99	0.37	MNS 88	99	
Parabacteroides distasonis		18.34	0.14	RHA 1	1	
Propionibacterium propionicum		12.67	0.14		99	
03533323	**Good identification**					**03533323**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03533340	**Good identification**					**03533340**
Propionibacterium propionicum		98.57	0.45	NIT 86		
Actinomyces naeslundii		1.44	0.24	RHA 1		
03533342	**Good identification**					**03533342**
Propionibacterium propionicum		99.04	0.48	NIT 86		
Actinomyces naeslundii		0.96	0.24	RHA 1		
03533500	**Low discrimination**					**03533500**
Propionibacterium propionicum		70.10	0.45	NIT 86		
Lactobacillus fermentum		11.68	0.32	ARA 1		
Actinomyces israelii		6.97	0.29			
03533501	**Low discrimination**					**03533501**
Actinomyces naeslundii		52.69	0.40	MNS 88		
Propionibacterium propionicum		47.31	0.29	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03533502	**Low discrimination**					**03533502**
Propionibacterium propionicum		56.01	0.48	NIT 86		
Bifidobacterium adolescentis		26.21	0.37	ARA 1		
Lactobacillus fermentum		6.22	0.32			
03533512	**Low discrimination**					**03533512**
Bifidobacterium adolescentis		83.45	0.38	MNS 87		
Bifidobacterium longum subsp. longum		12.91	0.23	XYL 87		
Actinomyces israelii		2.17	0.18			
03533521	**Good identification**					**03533521**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03533523	**Good identification**					**03533523**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03533540	**Acceptable identification**					**03533540**
Propionibacterium propionicum		94.95	0.45	NIT 86		
Actinomyces israelii		3.67	0.22	ARA 1		
Actinomyces naeslundii		1.38	0.24			
03533542	**Acceptable identification**					**03533542**
Propionibacterium propionicum		90.40	0.48	NIT 86		
Bifidobacterium adolescentis		5.23	0.21	ARA 1		
Actinomyces israelii		3.50	0.25			
03534100	**Low discrimination**					**03534100**
					GRA	
Fusobacterium mortiferum		53.42	0.40	MLT 11	1	
Actinomyces meyeri		38.64	0.29	XYL 7	99	
Lactobacillus fermentum		7.95	0.28		99	
03534520	**Good identification**					**03534520**
Bifidobacterium longum subsp. longum		95.04	0.37	MNS 87		
Lactobacillus fermentum		3.32	0.20	GAL 99		
Actinomyces meyeri		1.65	0.04			
03534522	**Acceptable identification**					**03534522**
					GRA	
Bifidobacterium longum subsp. longum		92.77	0.37	MNS 87	99	
Parabacteroides distasonis		4.00	0.18	GAL 99	1	
Lactobacillus fermentum		3.24	0.20		99	
03534530	**Good identification**					**03534530**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87		
03534532	**Good identification**					**03534532**
					GRA	
Bifidobacterium longum subsp. longum		96.92	0.41	MNS 87	99	
Clostridium clostridioforme		1.76	0.10	GAL 99	50	
Parabacteroides distasonis		1.32	0.14		1	
03535000	**Low discrimination**					**03535000**
					GRA	
Lactobacillus fermentum		87.36	0.58	TRE 1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		6.22	0.37		1	
Actinomyces meyeri		2.62	0.21		99	
03535002	**Very good identification**					**03535002**
					GRA	
Lactobacillus fermentum		96.90	0.58	TRE 1	99	
Actinomyces israelii		1.51	0.28		99	
Fusobacterium mortiferum		0.77	0.21		1	
03535020	**Good identification**					**03535020**
					CAT	
Lactobacillus fermentum		95.31	0.50	TRE 1	1	
Propionibacterium granulosum		1.72	0.15		90	
Actinomyces meyeri		1.17	0.07		1	
03535022	**Good identification**					**03535022**
					GRA	
Lactobacillus fermentum		96.08	0.50	TRE 1	99	
Parabacteroides distasonis		3.19	0.22		1	
Actinomyces naeslundii		0.73	0.24		99	
03535100	**Low discrimination**					**03535100**
					GRA	
Lactobacillus fermentum		80.20	0.62	TRE 1	99	
Fusobacterium mortiferum		8.17	0.43		1	
Actinomyces meyeri		5.91	0.32		99	
03535102	**Low discrimination**					**03535102**
Lactobacillus fermentum		84.79	0.62	TRE 1		
Bifidobacterium adolescentis		7.51	0.38			
Actinomyces israelii		3.24	0.39			
03535112	**Low discrimination**					**03535112**
Bifidobacterium adolescentis		64.13	0.40	MNS 87		
Bifidobacterium longum subsp. longum		24.34	0.31	ARA 87		
Lactobacillus fermentum		5.98	0.28			
03535120	**Low discrimination**					**03535120**
					GRA	
Lactobacillus fermentum		66.47	0.54	TRE 1	99	
Bifidobacterium longum subsp. longum		19.25	0.37		99	
Prevotella melaninogenica		5.00	0.15		1	
03535122	**Low discrimination**					**03535122**
					GRA	
Parabacteroides distasonis		59.38	0.56	MNS 98	1	
Lactobacillus fermentum		29.45	0.54		99	
Bifidobacterium longum subsp. longum		8.53	0.37		99	
03535130	**Good identification**					**03535130**
					GRA	
Bifidobacterium longum subsp. longum		95.90	0.41	MNS 87	99	
Parabacteroides distasonis		2.13	0.17	ARA 99	1	
Lactobacillus fermentum		1.97	0.20		99	
03535132	**Low discrimination**					**03535132**
					GRA	
Parabacteroides distasonis		68.34	0.51	MNS 98	1	
Bifidobacterium longum subsp. longum		31.03	0.41		99	
Lactobacillus fermentum		0.64	0.20		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03535322	**Very good identification**					**03535322**
					GRA	
Parabacteroides distasonis		99.77	0.64	MNS 98	1	
Lactobacillus fermentum		0.17	0.20		99	
Prevotella oralis		0.07	-.03		1	
03535332	**Very good identification**					**03535332**
					GRA	
Parabacteroides distasonis		99.46	0.60	MNS 98	1	
Clostridium clostridioforme		0.38	0.13		50	
Bifidobacterium longum subsp. longum		0.15	0.07		99	
03535400	**Good identification**					**03535400**
Lactobacillus fermentum		94.75	0.58	TRE 1		
Bifidobacterium longum subsp. longum		3.73	0.27			
Actinomyces israelii		0.81	0.24			
03535402	**Good identification**					**03535402**
Lactobacillus fermentum		94.17	0.58	TRE 1		
Bifidobacterium longum subsp. longum		3.71	0.27			
Actinomyces israelii		1.20	0.27			
03535420	**Low discrimination**					**03535420**
Lactobacillus fermentum		67.91	0.50	TRE 1		
Bifidobacterium longum subsp. longum		32.09	0.37			
03535422	**Low discrimination**					**03535422**
					GRA	
Lactobacillus fermentum		66.99	0.50	TRE 1	99	
Bifidobacterium longum subsp. longum		31.65	0.37		99	
Parabacteroides distasonis		1.37	0.18		1	
03535430	**Good identification**					**03535430**
Bifidobacterium longum subsp. longum		98.76	0.41	MNS 87		
Lactobacillus fermentum		1.24	0.16	LAC 99		
03535432	**Good identification**					**03535432**
					GRA	
Bifidobacterium longum subsp. longum		95.93	0.41	MNS 87	99	
Clostridium clostridioforme		1.56	0.09	LAC 99	50	
Parabacteroides distasonis		1.31	0.14		1	
03535500	**Low discrimination**					**03535500**
Bifidobacterium longum subsp. longum		69.57	0.61	MNS 87		
Lactobacillus fermentum		29.12	0.62			
Actinomyces israelii		0.61	0.34			
03535502	**Low discrimination**					**03535502**
Bifidobacterium longum subsp. longum		59.44	0.61	MNS 87		
Lactobacillus fermentum		24.88	0.62			
Bifidobacterium adolescentis		14.75	0.53			
03535510	**Very good identification**					**03535510**
Bifidobacterium longum subsp. longum		99.50	0.65	MNS 87		
Lactobacillus fermentum		0.25	0.28			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Bifidobacterium adolescentis	0.18	0.20			
03535512	**Low discrimination**					**03535512**
	Bifidobacterium longum subsp. longum	84.43	0.65	MNS 87		
	Bifidobacterium adolescentis	15.03	0.54			
	Lactobacillus fermentum	0.21	0.28			
03535520	**Very good identification**					**03535520**
					GRA	
	Bifidobacterium longum subsp. longum	95.98	0.72	MNS 87	99	
	Lactobacillus fermentum	3.35	0.54		99	
	Bacteroides vulgatus	0.60	0.20		1	
03535521	**Good identification**					**03535521**
	Bifidobacterium longum subsp. longum	96.63	0.37	MNS 87		
	Lactobacillus fermentum	3.37	0.20	URE 1		
03535522	**Good identification**					**03535522**
					GRA	
	Bifidobacterium longum subsp. longum	92.68	0.72	MNS 87	99	
	Parabacteroides distasonis	4.00	0.52		1	
	Lactobacillus fermentum	3.23	0.54		99	
03535523	**Acceptable identification**					**03535523**
					GRA	
	Bifidobacterium longum subsp. longum	92.77	0.37	MNS 87	99	
	Parabacteroides distasonis	4.00	0.18	URE 1	1	
	Lactobacillus fermentum	3.24	0.20		99	
03535530	**Very good identification**					**03535530**
					GRA	
	Bifidobacterium longum subsp. longum	99.97	0.76	MNS 87	99	
	Lactobacillus fermentum	0.02	0.20		99	
	Parabacteroides distasonis	0.01	0.14		1	
03535531	**Good identification**					**03535531**
	Bifidobacterium longum subsp. longum	100.0	0.41	MNS 87		
03535532	**Very good identification**					**03535532**
					GRA	
	Bifidobacterium longum subsp. longum	98.54	0.76	MNS 87	99	
	Parabacteroides distasonis	1.34	0.48		1	
	Clostridium clostridioforme	0.05	0.18		50	
03535533	**Good identification**					**03535533**
					GRA	
	Bifidobacterium longum subsp. longum	98.66	0.41	MNS 87	99	
	Parabacteroides distasonis	1.35	0.14	URE 1	1	
03535542	**Low discrimination**					**03535542**
	Bifidobacterium adolescentis	61.23	0.37	MNS 87		
	Bifidobacterium longum subsp. longum	20.17	0.27	SOR 11		
	Actinomyces israelii	10.16	0.30			
03535552	**Low discrimination**					**03535552**
	Bifidobacterium adolescentis	67.60	0.39	MNS 87		
	Bifidobacterium longum subsp. longum	31.03	0.31	SOR 11		
	Actinomyces israelii	1.37	0.16			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03535560	**Good identification**					**03535560**
Bifidobacterium longum subsp. longum		96.63	0.37	MNS 87		
Lactobacillus fermentum		3.37	0.20	SOR 1		
03535562	**Acceptable identification**					**03535562**
					GRA	
Bifidobacterium longum subsp. longum		92.77	0.37	MNS 87	99	
Parabacteroides distasonis		4.00	0.18	SOR 1	1	
Lactobacillus fermentum		3.24	0.20		99	
03535570	**Good identification**					**03535570**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87		
03535572	**Good identification**					**03535572**
					GRA	
Bifidobacterium longum subsp. longum		98.66	0.41	MNS 87	99	
Parabacteroides distasonis		1.35	0.14	SOR 1	1	
03535720	**Low discrimination**					**03535720**
					GRA	
Bacteroides vulgatus		82.61	0.36	MNS 99	1	
Bifidobacterium longum subsp. longum		14.94	0.37	TRE 4	99	
Parabacteroides distasonis		1.93	0.26		1	
03535722	**Good identification**					**03535722**
					GRA	
Parabacteroides distasonis		90.57	0.61	MNS 98	1	
Bifidobacterium longum subsp. longum		7.07	0.37		99	
Clostridium clostridioforme		1.06	0.22		50	
03535730	**Acceptable identification**					**03535730**
					GRA	
Bifidobacterium longum subsp. longum		93.14	0.41	MNS 87	99	
Parabacteroides distasonis		3.81	0.22	RHA 1	1	
Bacteroides vulgatus		3.06	0.02		1	
03535732	**Low discrimination**					**03535732**
					GRA	
Parabacteroides distasonis		74.02	0.56	MNS 98	1	
Bifidobacterium longum subsp. longum		18.28	0.41		99	
Clostridium clostridioforme		7.31	0.33		50	
03536100	**Good identification**					**03536100**
Propionibacterium propionicum		98.61	0.45	NIT 86		
Actinomyces israelii		1.39	0.14	XYL 1		
03536102	**Good identification**					**03536102**
Propionibacterium propionicum		98.61	0.48	NIT 86		
Actinomyces israelii		1.39	0.17	XYL 1		
03536140	**Good identification**					**03536140**
Propionibacterium propionicum		100.0	0.45	NIT 86		
03536142	**Good identification**					**03536142**
Propionibacterium propionicum		100.0	0.48	NIT 86		
03537000	**Low discrimination**					**03537000**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		63.99	0.38	SAL 92	1	
Propionibacterium granulosum		19.72	0.24	CEL 93	90	
Lactobacillus fermentum		10.98	0.24		1	
03537002	**Low discrimination**					**03537002**
Actinomyces israelii		80.00	0.41	SAL 92		
Lactobacillus fermentum		9.15	0.24	CEL 93		
Actinomyces naeslundii		4.42	0.29			
03537100	**Low discrimination**					**03537100**
Actinomyces israelii		54.22	0.48	SAL 92		
Propionibacterium propionicum		38.78	0.45	CEL 93		
Lactobacillus fermentum		3.80	0.28			
03537101	**Low discrimination**					**03537101**
Actinomyces naeslundii		49.70	0.40	MNS 88		
Propionibacterium propionicum		44.63	0.29	XYL 1		
Actinomyces israelii		5.67	0.14			
03537102	**Low discrimination**					**03537102**
Actinomyces israelii		46.82	0.51	SAL 92		
Propionibacterium propionicum		33.49	0.48	CEL 93		
Bifidobacterium adolescentis		15.67	0.37			
03537112	**Low discrimination**					**03537112**
Bifidobacterium adolescentis		72.31	0.38	MNS 87		
Actinomyces israelii		26.41	0.37	ARA 87		
Propionibacterium propionicum		1.28	0.14			
03537121	**Good identification**					**03537121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03537123	**Good identification**					**03537123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03537140	**Low discrimination**					**03537140**
Propionibacterium propionicum		64.18	0.45	NIT 86		
Actinomyces israelii		34.89	0.41	XYL 1		
Actinomyces naeslundii		0.93	0.24			
03537142	**Low discrimination**					**03537142**
Propionibacterium propionicum		62.07	0.48	NIT 86		
Actinomyces israelii		33.74	0.44	XYL 1		
Bifidobacterium adolescentis		3.59	0.21			
03537400	**Low discrimination**					**03537400**
Actinomyces israelii		82.66	0.37	SAL 92		
Lactobacillus fermentum		17.34	0.24	CEL 93		
03537402	**Low discrimination**					**03537402**
Actinomyces israelii		79.97	0.40	SAL 92		
Lactobacillus fermentum		11.18	0.24	CEL 93		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Bifidobacterium adolescentis	8.85	0.17			
03537500	**Low discrimination**					**03537500**
	Actinomyces israelii	75.11	0.47	SAL 92		
	Bifidobacterium longum subsp. longum	15.35	0.27	CEL 93		
	Lactobacillus fermentum	6.43	0.28			
03537502	**Low discrimination**					**03537502**
	Bifidobacterium adolescentis	69.49	0.51	MNS 87		
	Actinomyces israelii	25.38	0.50			
	Bifidobacterium longum subsp. longum	3.46	0.27			
03537512	**Low discrimination**					**03537512**
	Bifidobacterium adolescentis	89.77	0.53	MNS 87		
	Bifidobacterium longum subsp. longum	6.22	0.31			
	Actinomyces israelii	4.01	0.36			
03537520	**Low discrimination**					**03537520**
	Bifidobacterium longum subsp. longum	89.85	0.37	MNS 87		
	Actinomyces israelii	7.02	0.26	MAN 1		
	Lactobacillus fermentum	3.13	0.20			
03537522	**Low discrimination**					**03537522**
	Bifidobacterium longum subsp. longum	80.27	0.37	MNS 87		
	Actinomyces israelii	9.40	0.29	MAN 1		
	Bifidobacterium adolescentis	4.07	0.17			
03537530	**Good identification**					**03537530**
	Bifidobacterium longum subsp. longum	100.0	0.41	MNS 87		
03537532	**Acceptable identification**					**03537532**
					GRA	
	Bifidobacterium longum subsp. longum	94.31	0.41	MNS 87	99	
	Bifidobacterium adolescentis	3.44	0.18	MAN 1	99	
	Parabacteroides distasonis	1.29	0.14		1	
03537540	**Good identification**					**03537540**
	Actinomyces israelii	100.0	0.40	SAL 92		
03537542	**Low discrimination**					**03537542**
	Actinomyces israelii	52.83	0.43	SAL 92		
	Bifidobacterium adolescentis	45.97	0.36	CEL 93		
	Propionibacterium propionicum	1.20	0.14			
03537552	**Low discrimination**					**03537552**
	Bifidobacterium adolescentis	87.68	0.37	MNS 87		
	Actinomyces israelii	12.32	0.29	SOR 11		
03541001	**Good identification**					**03541001**
	Actinomyces naeslundii	100.0	0.36	MNS 88		
03541003	**Good identification**					**03541003**
	Actinomyces naeslundii	100.0	0.36	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
03541021	**Good identification**						**03541021**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03541023	**Good identification**						**03541023**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03541101	**Low discrimination**						**03541101**
					COC	SPO	
Actinomyces naeslundii		84.52	0.44	MNS 88	1	1	
Blautia hansenii		7.94	-.03	FRU 95	99	1	
Clostridium chauvoei		7.54	0.03		1	99	
03541103	**Acceptable identification**						**03541103**
Actinomyces naeslundii		94.54	0.44	MNS 88			
Actinomyces viscosus		5.46	-.01	FRU 95			
03541120	**Good identification**						**03541120**
					COC	GRA	
Prevotella melaninogenica		98.52	0.51	MNS 90	1	1	
Blautia hansenii		0.79	0.15	FRU 90	99	99	
Actinomyces naeslundii		0.51	0.41	CEL 10	1	99	
03541122	**Acceptable identification**						**03541122**
					GRA		
Prevotella melaninogenica		93.45	0.39	MNS 90	1		
Actinomyces naeslundii		2.52	0.41	FRU 90	99		
Prevotella oralis		2.26	0.15	CEL 10	1		
03541123	**Low discrimination**						**03541123**
					GRA		
Actinomyces naeslundii		80.04	0.44	MNS 88	99		
Prevotella melaninogenica		19.96	0.05	FRU 95	1		
03543000	**Good identification**						**03543000**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03543001	**Good identification**						**03543001**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03543002	**Good identification**						**03543002**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03543003	**Good identification**						**03543003**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03543020	**Good identification**						**03543020**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03543021	**Good identification**						**03543021**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03543022	**Good identification**						**03543022**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03543023	**Good identification**						**03543023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MNS 88		
03543100	**Low discrimination**					**03543100**
					COC	
Actinomyces naeslundii		89.72	0.46	MNS 88	1	
Blautia hansenii		6.51	-.03	FRU 95	99	
Propionibacterium propionicum		3.77	0.12		1	
03543101	**Good identification**					**03543101**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03543102	**Acceptable identification**					**03543102**
Actinomyces naeslundii		94.07	0.46	MNS 88		
Propionibacterium propionicum		5.94	0.15	FRU 95		
03543103	**Good identification**					**03543103**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03543121	**Good identification**					**03543121**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03543122	**Low discrimination**					**03543122**
					GRA	
Actinomyces naeslundii		83.84	0.46	MNS 88	99	
Prevotella melaninogenica		16.16	0.05	FRU 95	1	
03543123	**Good identification**					**03543123**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03543141	**Good identification**					**03543141**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03543143	**Good identification**					**03543143**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03543161	**Good identification**					**03543161**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03543163	**Good identification**					**03543163**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03545530	**Good identification**					**03545530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87		
03545532	**Good identification**					**03545532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87		
03550000	**Low discrimination**					**03550000**
					COC	
Fusobacterium mortiferum		61.12	0.54	MLT 11	1	
Prevotella intermedia		25.35	0.36		1	
Megasphaera elsdenii		9.83	0.32		99	
03550100	**Good identification**					**03550100**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
				GRA	
Fusobacterium mortiferum	94.20	0.60	MLT 11	1	
Actinomyces meyeri	2.37	0.24		99	
Lactobacillus fermentum	1.61	0.32		99	
03550101 Low discrimination					**03550101**
				GRA	
Actinomyces naeslundii	64.58	0.42	MNS 88	99	
Fusobacterium mortiferum	35.42	0.26	GAL 96	1	
03550121 Low discrimination					**03550121**
				GRA	
Actinomyces naeslundii	86.93	0.42	MNS 88	99	
Prevotella melaninogenica	13.07	-.01	GAL 96	1	
03550123 Good identification					**03550123**
Actinomyces naeslundii	100.0	0.42	MNS 88		
03551000 Low discrimination					**03551000**
				GRA	
Lactobacillus fermentum	52.33	0.62	CEL 1	99	
Fusobacterium mortiferum	32.44	0.57		1	
Prevotella intermedia	8.97	0.36		1	
03551001 Low discrimination					**03551001**
				GRA	
Actinomyces naeslundii	89.22	0.58	MNS 88	99	
Lactobacillus fermentum	6.02	0.28		99	
Fusobacterium mortiferum	3.73	0.23		1	
03551002 Low discrimination					**03551002**
				GRA	
Lactobacillus fermentum	82.02	0.62	CEL 1	99	
Actinomyces naeslundii	8.19	0.55		99	
Fusobacterium mortiferum	5.65	0.41		1	
03551003 Low discrimination					**03551003**
Actinomyces naeslundii	86.22	0.58	MNS 88		
Actinomyces viscosus	7.97	0.16			
Lactobacillus fermentum	5.82	0.28			
03551011 Good identification					**03551011**
Actinomyces naeslundii	100.0	0.36	MNS 88		
03551013 Good identification					**03551013**
Actinomyces naeslundii	100.0	0.36	MNS 88		
03551020 Low discrimination					**03551020**
				GRA	
Lactobacillus fermentum	46.54	0.54	CEL 1	99	
Prevotella melaninogenica	36.92	0.33		1	
Actinomyces naeslundii	13.94	0.55		99	
03551021 Very good identification					**03551021**
				GRA	
Actinomyces naeslundii	96.13	0.58	MNS 88	99	
Lactobacillus fermentum	2.16	0.20		99	
Prevotella melaninogenica	1.71	-.01		1	
03551022 Low discrimination					**03551022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Lactobacillus fermentum		58.18	0.54	CEL 1	99	
Actinomyces naeslundii		17.43	0.55		99	
Prevotella oralis		14.78	0.28		1	
03551023	**Very good identification**					**03551023**
Actinomyces naeslundii		97.80	0.58	MNS 88		
Lactobacillus fermentum		2.20	0.20			
03551031	**Good identification**					**03551031**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03551033	**Good identification**					**03551033**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03551040	**Low discrimination**					**03551040**
					GRA	
Actinomyces naeslundii		49.34	0.42	MNS 88	99	
Lactobacillus fermentum		28.28	0.28		99	
Fusobacterium mortiferum		17.53	0.23		1	
03551041	**Good identification**					**03551041**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03551042	**Low discrimination**					**03551042**
Actinomyces naeslundii		63.56	0.42	MNS 88		
Lactobacillus fermentum		36.44	0.28			
03551043	**Good identification**					**03551043**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03551060	**Low discrimination**					**03551060**
					GRA	
Actinomyces naeslundii		74.48	0.42	MNS 88	99	
Lactobacillus fermentum		14.23	0.20		99	
Prevotella melaninogenica		11.29	-.01		1	
03551061	**Good identification**					**03551061**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03551062	**Low discrimination**					**03551062**
Actinomyces naeslundii		83.96	0.42	MNS 88		
Lactobacillus fermentum		16.04	0.20			
03551063	**Good identification**					**03551063**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03551100	**Low discrimination**					**03551100**
					GRA	
Lactobacillus fermentum		44.31	0.66	CEL 1	99	
Fusobacterium mortiferum		39.29	0.63		1	
Actinomyces naeslundii		7.72	0.63		99	
03551101	**Good identification**					**03551101**
					GRA	
Actinomyces naeslundii		91.73	0.66	MNS 88	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		3.55	0.32		99	
Fusobacterium mortiferum		3.14	0.29		1	
03551102	**Low discrimination**					**03551102**
					GRA	
Lactobacillus fermentum		69.22	0.66	CEL 1	99	
Actinomyces naeslundii		12.06	0.63		99	
Fusobacterium mortiferum		6.82	0.47		1	
03551103	**Good identification**					**03551103**
Actinomyces naeslundii		90.89	0.66	MNS 88		
Actinomyces viscosus		5.25	0.21			
Lactobacillus fermentum		3.51	0.32			
03551111	**Good identification**					**03551111**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03551113	**Good identification**					**03551113**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03551120	**Very good identification**					**03551120**
					GRA	
Prevotella melaninogenica		96.39	0.67	MNS 90	1	
Lactobacillus fermentum		2.00	0.58	CEL 10	99	
Actinomyces naeslundii		1.05	0.63		99	
03551121	**Low discrimination**					**03551121**
					GRA	
Actinomyces naeslundii		61.09	0.66	MNS 88	99	
Prevotella melaninogenica		37.90	0.33		1	
Lactobacillus fermentum		0.79	0.24		99	
03551122	**Low discrimination**					**03551122**
					GRA	
Prevotella melaninogenica		67.23	0.55	MNS 90	1	
Prevotella oralis		17.89	0.49	CEL 10	1	
Lactobacillus fermentum		7.33	0.58		99	
03551123	**Low discrimination**					**03551123**
					GRA	
Actinomyces naeslundii		85.61	0.66	MNS 88	99	
Prevotella melaninogenica		10.12	0.21		1	
Prevotella oralis		2.69	0.15		1	
03551131	**Low discrimination**					**03551131**
					GRA	
Actinomyces naeslundii		89.36	0.44	MNS 88	99	
Prevotella melaninogenica		10.64	-.01	MLZ 5	1	
03551133	**Good identification**					**03551133**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03551140	**Low discrimination**					**03551140**
					GRA	
Actinomyces naeslundii		56.96	0.50	MNS 88	99	
Lactobacillus fermentum		18.72	0.32		99	
Fusobacterium mortiferum		16.60	0.29		1	
03551141	**Good identification**					**03551141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	MNS 88		
03551142	**Low discrimination**					**03551142**
Actinomyces naeslundii		60.87	0.50	MNS 88		
Lactobacillus fermentum		20.00	0.32			
Bifidobacterium adolescentis		6.06	0.17			
03551143	**Good identification**					**03551143**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03551161	**Good identification**					**03551161**
					GRA	
Actinomyces naeslundii		96.57	0.53	MNS 88	99	
Prevotella melaninogenica		3.43	-.01		1	
03551163	**Good identification**					**03551163**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03551322	**Low discrimination**					**03551322**
Prevotella oralis		86.80	0.42	MNS 99		
Prevotella melaninogenica		7.69	0.21			
Parabacteroides distasonis		3.87	0.32			
03551400	**Very good identification**					**03551400**
					GRA	
Lactobacillus fermentum		98.83	0.62	CEL 1	99	
Fusobacterium mortiferum		0.62	0.23		1	
Actinomyces meyeri		0.23	0.06		99	
03551402	**Very good identification**					**03551402**
Lactobacillus fermentum		99.50	0.62	CEL 1		
Bifidobacterium adolescentis		0.27	0.13			
Actinomyces israelii		0.23	0.18			
03551420	**Good identification**					**03551420**
					GRA	
Lactobacillus fermentum		98.84	0.54	CEL 1	99	
Prevotella melaninogenica		0.79	-.01		1	
Bifidobacterium longum subsp. longum		0.37	0.05		99	
03551422	**Good identification**					**03551422**
Lactobacillus fermentum		99.63	0.54	CEL 1		
Bifidobacterium longum subsp. longum		0.37	0.05			
03551500	**Very good identification**					**03551500**
					GRA	
Lactobacillus fermentum		95.95	0.66	CEL 1	99	
Bifidobacterium longum subsp. longum		1.81	0.29		99	
Fusobacterium mortiferum		0.86	0.29		1	
03551502	**Low discrimination**					**03551502**
Lactobacillus fermentum		83.62	0.66	CEL 1		
Bifidobacterium adolescentis		13.86	0.47			
Bifidobacterium longum subsp. longum		1.58	0.29			
03551512	**Low discrimination**					**03551512**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		82.45	0.49	MNS 87			
Bifidobacterium longum subsp. longum		13.09	0.33	XYL 87			
Lactobacillus fermentum		4.11	0.32				
03551520	**Low discrimination**						**03551520**
					GRA		
Lactobacillus fermentum		57.75	0.58	CEL 1	99		
Prevotella melaninogenica		28.07	0.33		1		
Bifidobacterium longum subsp. longum		13.09	0.40		99		
03551522	**Low discrimination**						**03551522**
					GRA		
Lactobacillus fermentum		69.39	0.58	CEL 1	99		
Bifidobacterium longum subsp. longum		15.74	0.40		99		
Prevotella melaninogenica		6.43	0.21		1		
03551530	**Good identification**						**03551530**
					GRA		
Bifidobacterium longum subsp. longum		96.26	0.44	MNS 87	99		
Lactobacillus fermentum		2.52	0.24	CEL 1	99		
Prevotella melaninogenica		1.22	-.01		1		
03551532	**Acceptable identification**						**03551532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.80	0.44	MNS 87	99	1	
Lactobacillus fermentum		2.43	0.24	CEL 1	99	1	
Capnocytophaga ochracea		2.06	0.07		1	90	
03552001	**Good identification**						**03552001**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03552002	**Low discrimination**						**03552002**
					COC	GRA	
Actinomyces naeslundii		46.70	0.36	MNS 88	1	99	
Megasphaera elsdenii		42.70	0.15	GAL 96	99	1	
Propionibacterium propionicum		10.59	0.15		1	99	
03552003	**Good identification**						**03552003**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03552020	**Good identification**						**03552020**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03552021	**Good identification**						**03552021**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03552022	**Good identification**						**03552022**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03552023	**Good identification**						**03552023**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03552100	**Low discrimination**						**03552100**
					GRA		
Propionibacterium propionicum		78.70	0.46	NIT 86	99		
Actinomyces naeslundii		14.96	0.44	CEL 1	99		
Fusobacterium mortiferum		6.34	0.26		1		
03552101	**Low discrimination**						**03552101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		71.96	0.47	MNS 88		
Propionibacterium propionicum		28.04	0.30	GAL 96		
03552102	**Low discrimination**					**03552102**
Propionibacterium propionicum		88.23	0.49	NIT 86		
Actinomyces naeslundii		11.18	0.44	CEL 1		
Actinomyces viscosus		0.59	-.02			
03552103	**Low discrimination**					**03552103**
Actinomyces naeslundii		63.11	0.47	MNS 88		
Propionibacterium propionicum		36.89	0.33	GAL 96		
03552120	**Low discrimination**					**03552120**
					GRA	
Actinomyces naeslundii		85.52	0.44	MNS 88	99	
Prevotella melaninogenica		9.94	-.01	GAL 96	1	
Propionibacterium propionicum		4.54	0.12		99	
03552121	**Good identification**					**03552121**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03552122	**Acceptable identification**					**03552122**
Actinomyces naeslundii		92.62	0.44	MNS 88		
Propionibacterium propionicum		7.38	0.15	GAL 96		
03552123	**Good identification**					**03552123**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03552140	**Good identification**					**03552140**
Propionibacterium propionicum		96.75	0.46	NIT 86		
Actinomyces naeslundii		3.25	0.31	CEL 1		
03552142	**Good identification**					**03552142**
Propionibacterium propionicum		97.81	0.49	NIT 86		
Actinomyces naeslundii		2.19	0.31	CEL 1		
03553000	**Low discrimination**					**03553000**
Actinomyces naeslundii		85.27	0.60	MNS 88		
Actinomyces israelii		4.67	0.29			
Lactobacillus fermentum		4.44	0.28			
03553001	**Very good identification**					**03553001**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03553002	**Low discrimination**					**03553002**
Actinomyces naeslundii		87.43	0.60	MNS 88		
Actinomyces israelii		7.18	0.32			
Lactobacillus fermentum		4.56	0.28			
03553003	**Very good identification**					**03553003**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03553010	**Low discrimination**					**03553010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		86.54	0.38	MNS 88		
Actinomyces israelii		13.46	0.15	MLZ 5		
03553011	**Good identification**					**03553011**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03553012	**Low discrimination**					**03553012**
Actinomyces naeslundii		81.08	0.38	MNS 88		
Actinomyces israelii		18.92	0.18	MLZ 5		
03553013	**Good identification**					**03553013**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03553020	**Very good identification**					**03553020**
					GRA	
Actinomyces naeslundii		96.98	0.60	MNS 88	99	
Lactobacillus fermentum		1.69	0.20		99	
Prevotella melaninogenica		1.34	-.01		1	
03553021	**Very good identification**					**03553021**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03553022	**Very good identification**					**03553022**
Actinomyces naeslundii		98.29	0.60	MNS 88		
Lactobacillus fermentum		1.71	0.20			
03553023	**Very good identification**					**03553023**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03553030	**Good identification**					**03553030**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03553031	**Good identification**					**03553031**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03553032	**Good identification**					**03553032**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03553033	**Good identification**					**03553033**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03553040	**Low discrimination**					**03553040**
Actinomyces naeslundii		86.47	0.47	MNS 88		
Actinomyces israelii		10.44	0.22			
Propionibacterium propionicum		3.09	0.12			
03553041	**Good identification**					**03553041**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03553042	**Low discrimination**					**03553042**
Actinomyces naeslundii		81.00	0.47	MNS 88		
Actinomyces israelii		14.67	0.25			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		4.34	0.15			
03553043	**Good identification**					**03553043**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03553060	**Good identification**					**03553060**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03553061	**Good identification**					**03553061**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03553062	**Good identification**					**03553062**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03553063	**Good identification**					**03553063**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03553100	**Low discrimination**					**03553100**
Actinomyces naeslundii		73.67	0.68	MNS 88		
Propionibacterium propionicum		16.15	0.46			
Actinomyces israelii		5.67	0.40			
03553101	**Very good identification**					**03553101**
Actinomyces naeslundii		98.40	0.71	MNS 88		
Propionibacterium propionicum		1.60	0.30			
03553102	**Low discrimination**					**03553102**
Actinomyces naeslundii		65.06	0.68	MNS 88		
Propionibacterium propionicum		21.39	0.49			
Actinomyces israelii		7.51	0.43			
03553103	**Very good identification**					**03553103**
Actinomyces naeslundii		97.62	0.71	MNS 88		
Propionibacterium propionicum		2.38	0.33			
03553110	**Low discrimination**					**03553110**
Actinomyces naeslundii		79.33	0.46	MNS 88		
Actinomyces israelii		17.34	0.25	MLZ 5		
Propionibacterium propionicum		3.34	0.12			
03553111	**Good identification**					**03553111**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03553113	**Good identification**					**03553113**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03553120	**Low discrimination**					**03553120**
					GRA	
Actinomyces naeslundii		66.83	0.68	MNS 88	99	
Prevotella melaninogenica		32.03	0.33		1	
Lactobacillus fermentum		0.67	0.24		99	
03553121	**Very good identification**					**03553121**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.68	0.71	MNS 88	99	
Prevotella melaninogenica		0.32	-.01		1	
03553122	**Low discrimination**					**03553122**
					GRA	
Actinomyces naeslundii		87.21	0.68	MNS 88	99	
Prevotella melaninogenica		7.96	0.21		1	
Prevotella oralis		2.12	0.15		1	
03553123	**Very good identification**					**03553123**
Actinomyces naeslundii		100.0	0.71	MNS 88		
03553130	**Acceptable identification**					**03553130**
					GRA	
Actinomyces naeslundii		91.58	0.46	MNS 88	99	
Prevotella melaninogenica		8.43	-.01	MLZ 5	1	
03553131	**Good identification**					**03553131**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03553132	**Good identification**					**03553132**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03553133	**Good identification**					**03553133**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03553140	**Low discrimination**					**03553140**
Propionibacterium propionicum		51.50	0.46	NIT 86		
Actinomyces naeslundii		41.47	0.55	CEL 1		
Actinomyces israelii		7.04	0.33			
03553141	**Good identification**					**03553141**
Actinomyces naeslundii		91.58	0.58	MNS 88		
Propionibacterium propionicum		8.43	0.30			
03553142	**Low discrimination**					**03553142**
Propionibacterium propionicum		58.97	0.49	NIT 86		
Actinomyces naeslundii		31.65	0.55	CEL 1		
Actinomyces israelii		8.06	0.36			
03553143	**Low discrimination**					**03553143**
Actinomyces naeslundii		87.87	0.58	MNS 88		
Propionibacterium propionicum		12.13	0.33			
03553151	**Good identification**					**03553151**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03553153	**Good identification**					**03553153**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03553160	**Good identification**					**03553160**
					GRA	
Actinomyces naeslundii		96.16	0.55	MNS 88	99	
Prevotella melaninogenica		2.64	-.01		1	
Propionibacterium propionicum		1.21	0.12		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03553161	**Very good identification**					**03553161**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03553162	**Good identification**					**03553162**
Actinomyces naeslundii		96.61	0.55	MNS 88		
Propionibacterium propionicum		1.82	0.15			
Actinomyces israelii		1.57	0.15			
03553163	**Very good identification**					**03553163**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03553171	**Good identification**					**03553171**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03553173	**Good identification**					**03553173**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03553301	**Good identification**					**03553301**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03553303	**Good identification**					**03553303**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03553321	**Good identification**					**03553321**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03553323	**Good identification**					**03553323**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03553500	**Low discrimination**					**03553500**
Actinomyces israelii		57.67	0.38	SAL 92		
Lactobacillus fermentum		27.35	0.32	XYL 92		
Actinomyces naeslundii		9.25	0.34			
03553501	**Good identification**					**03553501**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03553502	**Low discrimination**					**03553502**
Bifidobacterium adolescentis		74.44	0.46	MNS 87		
Actinomyces israelii		17.53	0.41	XYL 87		
Lactobacillus fermentum		5.54	0.32			
03553503	**Low discrimination**					**03553503**
Actinomyces naeslundii		78.90	0.37	MNS 88		
Bifidobacterium adolescentis		21.10	0.11	ARA 1		
03553512	**Good identification**					**03553512**
Bifidobacterium adolescentis		97.20	0.47	MNS 87		
Actinomyces israelii		2.80	0.27	XYL 87		
03553521	**Good identification**					**03553521**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	MNS 88			
03553523	**Good identification**						**03553523**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03554100	**Low discrimination**						**03554100**
					GRA		
Fusobacterium mortiferum		56.13	0.41	MLT 11	1		
Actinomyces meyeri		36.38	0.29	XYL 7	99		
Lactobacillus fermentum		7.49	0.28		99		
03555000	**Low discrimination**						**03555000**
					GRA		
Lactobacillus fermentum		87.02	0.58	CEL 1	99		
Fusobacterium mortiferum		6.91	0.38		1		
Actinomyces israelii		3.19	0.35		99		
03555002	**Good identification**						**03555002**
					GRA		
Lactobacillus fermentum		93.59	0.58	CEL 1	99		
Actinomyces israelii		5.15	0.38		99		
Fusobacterium mortiferum		0.83	0.22		1		
03555020	**Good identification**						**03555020**
					GRA		
Lactobacillus fermentum		96.81	0.50	CEL 1	99		
Prevotella melaninogenica		1.32	-.01		1		
Actinomyces meyeri		1.19	0.07		99		
03555022	**Acceptable identification**						**03555022**
					GRA		
Lactobacillus fermentum		93.11	0.50	CEL 1	99		
Prevotella oralis		4.98	0.13		1		
Actinomyces israelii		0.98	0.17		99		
03555100	**Low discrimination**						**03555100**
					GRA		
Lactobacillus fermentum		77.77	0.62	CEL 1	99		
Fusobacterium mortiferum		8.84	0.44		1		
Actinomyces israelii		7.00	0.45		99		
03555102	**Low discrimination**						**03555102**
Lactobacillus fermentum		68.82	0.62	CEL 1			
Bifidobacterium adolescentis		19.43	0.47				
Actinomyces israelii		9.29	0.48				
03555112	**Acceptable identification**						**03555112**
Bifidobacterium adolescentis		91.53	0.49	MNS 87			
Actinomyces israelii		5.35	0.34	ARA 87			
Lactobacillus fermentum		2.68	0.28				
03555120	**Low discrimination**						**03555120**
					CAT	GRA	
Lactobacillus fermentum		50.19	0.54	CEL 1	1	99	
Prevotella melaninogenica		41.55	0.33		1	1	
Bacteroides fragilis		1.60	0.04		99	1	
03555122	**Low discrimination**						**03555122**
					CAT	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		36.29	0.54	CEL 1	1	99	
Bacteroides fragilis		21.93	0.25		99	1	
Prevotella oralis		18.63	0.33		1	1	
03555132	**Low discrimination**						**03555132**
					GRA	CAT	
Parabacteroides distasonis		62.70	0.36	MNS 98	1	76	
Capnocytophaga ochracea		10.12	0.10		1	90	
Bifidobacterium longum subsp. longum		9.73	0.18		99	1	
03555140	**Low discrimination**						**03555140**
Actinomyces israelii		77.59	0.38	SAL 92			
Lactobacillus fermentum		22.41	0.28				
03555142	**Low discrimination**						**03555142**
Actinomyces israelii		53.84	0.41	SAL 92			
Bifidobacterium adolescentis		35.80	0.32				
Lactobacillus fermentum		10.37	0.28				
03555322	**Low discrimination**						**03555322**
					GRA		
Parabacteroides distasonis		75.17	0.48	MNS 98	1		
Prevotella oralis		23.13	0.27		1		
Lactobacillus fermentum		1.06	0.20		99		
03555332	**Good identification**						**03555332**
Parabacteroides distasonis		98.53	0.44	MNS 98			
Clostridium clostridioforme		1.47	0.07				
03555400	**Very good identification**						**03555400**
Lactobacillus fermentum		96.38	0.58	CEL 1			
Actinomyces israelii		2.89	0.33				
Actinomyces meyeri		0.72	0.11				
03555402	**Good identification**						**03555402**
Lactobacillus fermentum		92.92	0.58	CEL 1			
Actinomyces israelii		4.18	0.36				
Bifidobacterium adolescentis		2.89	0.27				
03555420	**Good identification**						**03555420**
Lactobacillus fermentum		98.13	0.50	CEL 1			
Bifidobacterium longum subsp. longum		1.87	0.14				
03555422	**Good identification**						**03555422**
Lactobacillus fermentum		97.30	0.50	CEL 1			
Bifidobacterium longum subsp. longum		1.86	0.14				
Actinomyces israelii		0.84	0.16				
03555500	**Low discrimination**						**03555500**
Lactobacillus fermentum		82.65	0.62	CEL 1			
Bifidobacterium longum subsp. longum		7.98	0.37				
Actinomyces israelii		6.08	0.43				
03555502	**Low discrimination**						**03555502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		61.02	0.61	MNS 87		
Lactobacillus fermentum		32.30	0.62			
Actinomyces israelii		3.57	0.46			
03555510	**Low discrimination**					**03555510**
Bifidobacterium longum subsp. longum		78.72	0.41	MNS 87		
Bifidobacterium adolescentis		11.17	0.29	CEL 1		
Actinomyces israelii		5.27	0.29			
03555512	**Good identification**					**03555512**
Bifidobacterium adolescentis		92.21	0.63	MNS 87		
Bifidobacterium longum subsp. longum		6.56	0.41			
Actinomyces israelii		0.66	0.32			
03555520	**Low discrimination**					**03555520**
					GRA	
Bifidobacterium longum subsp. longum		48.00	0.48	MNS 87	99	
Lactobacillus fermentum		41.44	0.54	CEL 1	99	
Bacteroides vulgatus		9.31	0.21		1	
03555522	**Low discrimination**					**03555522**
					GRA	
Bifidobacterium longum subsp. longum		47.73	0.48	MNS 87	99	
Lactobacillus fermentum		41.20	0.54	CEL 1	99	
Parabacteroides distasonis		6.02	0.37		1	
03555530	**Good identification**					**03555530**
Bifidobacterium longum subsp. longum		99.49	0.52	MNS 87		
Lactobacillus fermentum		0.51	0.20	CEL 1		
03555532	**Acceptable identification**					**03555532**
					GRA	
Bifidobacterium longum subsp. longum		92.02	0.52	MNS 87	99	
Parabacteroides distasonis		3.67	0.32	CEL 1	1	
Bifidobacterium adolescentis		3.26	0.29		99	
03555540	**Low discrimination**					**03555540**
Actinomyces israelii		69.67	0.36	SAL 92		
Lactobacillus fermentum		24.59	0.28			
Bifidobacterium adolescentis		5.74	0.12			
03555542	**Low discrimination**					**03555542**
Bifidobacterium adolescentis		81.49	0.46	MNS 87		
Actinomyces israelii		14.98	0.39	SOR 11		
Lactobacillus fermentum		3.53	0.28			
03555552	**Good identification**					**03555552**
Bifidobacterium adolescentis		97.25	0.47	MNS 87		
Actinomyces israelii		2.19	0.25	SOR 11		
Bifidobacterium longum subsp. longum		0.57	0.07			
03555720	**Good identification**					**03555720**
					GRA	
Bacteroides vulgatus		98.72	0.38	MNS 99	1	
Bifidobacterium longum subsp. longum		0.57	0.14	CEL 5	99	
Lactobacillus fermentum		0.49	0.20		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03555722	**Low discrimination**					**03555722**
Parabacteroides distasonis		86.86	0.45	MNS 98		
Bacteroides vulgatus		4.05	0.04			
Clostridium clostridioforme		3.91	0.16			
03555732	**Low discrimination**					**03555732**
					GRA	
Parabacteroides distasonis		68.23	0.40	MNS 98	1	
Clostridium clostridioforme		26.01	0.27		50	
Bifidobacterium longum subsp. longum		5.76	0.18		99	
03557000	**Acceptable identification**					**03557000**
Actinomyces israelii		93.93	0.47	SAL 92		
Lactobacillus fermentum		4.56	0.24			
Actinomyces naeslundii		1.51	0.26			
03557002	**Acceptable identification**					**03557002**
Actinomyces israelii		94.76	0.50	SAL 92		
Lactobacillus fermentum		3.07	0.24			
Bifidobacterium adolescentis		1.16	0.11			
03557012	**Acceptable identification**					**03557012**
Actinomyces israelii		90.92	0.36	SAL 92		
Bifidobacterium adolescentis		9.08	0.13	MLZ 13		
03557040	**Good identification**					**03557040**
Actinomyces israelii		100.0	0.40	SAL 92		
03557042	**Good identification**					**03557042**
Actinomyces israelii		100.0	0.43	SAL 92		
03557100	**Very good identification**					**03557100**
Actinomyces israelii		96.31	0.58	SAL 92		
Lactobacillus fermentum		1.91	0.28			
Actinomyces naeslundii		1.10	0.34			
03557101	**Low discrimination**					**03557101**
Actinomyces naeslundii		62.88	0.37	MNS 88		
Actinomyces israelii		37.12	0.24	XYL 1		
03557102	**Low discrimination**					**03557102**
Actinomyces israelii		75.45	0.61	SAL 92		
Bifidobacterium adolescentis		22.79	0.46			
Lactobacillus fermentum		1.00	0.28			
03557103	**Low discrimination**					**03557103**
Actinomyces naeslundii		46.45	0.37	MNS 88		
Actinomyces israelii		41.13	0.27	XYL 1		
Bifidobacterium adolescentis		12.43	0.11			
03557110	**Good identification**					**03557110**
Actinomyces israelii		96.39	0.44	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		3.61	0.13	MLZ 13		
03557112	**Low discrimination**					**03557112**
Bifidobacterium adolescentis		71.19	0.47	MNS 87		
Actinomyces israelii		28.81	0.47	ARA 87		
03557120	**Low discrimination**					**03557120**
Actinomyces israelii		73.11	0.37	SAL 92		
Actinomyces naeslundii		13.07	0.34	NAG 6		
Lactobacillus fermentum		7.57	0.20			
03557121	**Good identification**					**03557121**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03557122	**Low discrimination**					**03557122**
Actinomyces israelii		76.16	0.40	SAL 92		
Actinomyces naeslundii		9.07	0.34	NAG 6		
Lactobacillus fermentum		5.26	0.20			
03557123	**Good identification**					**03557123**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03557140	**Good identification**					**03557140**
Actinomyces israelii		99.36	0.51	SAL 92		
Propionibacterium propionicum		0.64	0.12			
03557142	**Acceptable identification**					**03557142**
Actinomyces israelii		90.71	0.54	SAL 92		
Bifidobacterium adolescentis		8.71	0.30			
Propionibacterium propionicum		0.58	0.15			
03557150	**Good identification**					**03557150**
Actinomyces israelii		100.0	0.37	SAL 92		
03557152	**Low discrimination**					**03557152**
Actinomyces israelii		56.01	0.40	SAL 92		
Bifidobacterium adolescentis		43.99	0.32	MLZ 13		
03557300	**Good identification**					**03557300**
Actinomyces israelii		100.0	0.39	SAL 92		
03557302	**Good identification**					**03557302**
Actinomyces israelii		96.10	0.42	SAL 92		
Bifidobacterium adolescentis		3.90	0.11	RHA 7		
03557400	**Acceptable identification**					**03557400**
Actinomyces israelii		94.39	0.46	SAL 92		
Lactobacillus fermentum		5.61	0.24			
03557402	**Low discrimination**					**03557402**
Actinomyces israelii		87.76	0.49	SAL 92		
Bifidobacterium adolescentis		8.76	0.26			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Lactobacillus fermentum	3.48	0.24			
03557440	**Good identification**					**03557440**
	Actinomyces israelii	100.0	0.39	SAL 92		
03557442	**Good identification**					**03557442**
	Actinomyces israelii	96.92	0.42	SAL 92		
	Bifidobacterium adolescentis	3.08	0.10			
03557500	**Good identification**					**03557500**
	Actinomyces israelii	94.19	0.56	SAL 92		
	Bifidobacterium adolescentis	3.53	0.26			
	Lactobacillus fermentum	2.28	0.28			
03557502	**Low discrimination**					**03557502**
	Bifidobacterium adolescentis	70.86	0.60	MNS 87		
	Actinomyces israelii	28.68	0.59			
	Lactobacillus fermentum	0.46	0.28			
03557510	**Low discrimination**					**03557510**
	Actinomyces israelii	75.03	0.42	SAL 92		
	Bifidobacterium adolescentis	22.98	0.27	MLZ 13		
	Bifidobacterium longum subsp. longum	2.00	0.07			
03557512	**Very good identification**					**03557512**
	Bifidobacterium adolescentis	95.21	0.61	MNS 87		
	Actinomyces israelii	4.71	0.45			
	Bifidobacterium longum subsp. longum	0.08	0.07			
03557520	**Low discrimination**					**03557520**
	Actinomyces israelii	78.55	0.36	SAL 92		
	Bifidobacterium longum subsp. longum	11.51	0.14	NAG 6		
	Lactobacillus fermentum	9.94	0.20			
03557522	**Low discrimination**					**03557522**
	Actinomyces israelii	63.57	0.39	SAL 92		
	Bifidobacterium adolescentis	24.86	0.26	NAG 6		
	Bifidobacterium longum subsp. longum	6.21	0.14			
03557540	**Good identification**					**03557540**
	Actinomyces israelii	98.82	0.49	SAL 92		
	Bifidobacterium adolescentis	1.18	0.10			
03557542	**Low discrimination**					**03557542**
	Actinomyces israelii	56.01	0.52	SAL 92		
	Bifidobacterium adolescentis	43.99	0.44			
03557550	**Acceptable identification**					**03557550**
	Actinomyces israelii	91.13	0.35	SAL 92		
	Bifidobacterium adolescentis	8.87	0.12	MLZ 13		
03557552	**Low discrimination**					**03557552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		86.53	0.46	MNS 87			
Actinomyces israelii		13.47	0.38	SOR 11			
03557700	**Good identification**						**03557700**
Actinomyces israelii		100.0	0.37	SAL 92			
03557702	**Low discrimination**						**03557702**
Actinomyces israelii		75.10	0.40	SAL 92			
Bifidobacterium adolescentis		24.90	0.26	RHA 7			
03561000	**Low discrimination**						**03561000**
					GRA		
Actinomyces naeslundii		84.50	0.38	MNS 88	99		
Fusobacterium mortiferum		15.50	0.14	FRU 95	1		
03561001	**Good identification**						**03561001**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03561002	**Good identification**						**03561002**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03561003	**Good identification**						**03561003**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03561020	**Good identification**						**03561020**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03561021	**Good identification**						**03561021**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03561022	**Good identification**						**03561022**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03561023	**Good identification**						**03561023**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03561100	**Low discrimination**						**03561100**
					COC	GRA	
Actinomyces naeslundii		81.78	0.46	MNS 88	1	99	
Fusobacterium mortiferum		12.29	0.20	FRU 95	1	1	
Blautia hansenii		5.93	-.03		99	99	
03561101	**Good identification**						**03561101**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03561102	**Good identification**						**03561102**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03561103	**Good identification**						**03561103**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03561121	**Good identification**						**03561121**
Actinomyces naeslundii		100.0	0.49	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03561122	**Low discrimination**					**03561122**
					GRA	
Actinomyces naeslundii		83.84	0.46	MNS 88	99	
Prevotella melaninogenica		16.16	0.05	FRU 95	1	
03561123	**Good identification**					**03561123**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03561141	**Good identification**					**03561141**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03561143	**Good identification**					**03561143**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03561161	**Good identification**					**03561161**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03561163	**Good identification**					**03561163**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03563000	**Good identification**					**03563000**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03563001	**Good identification**					**03563001**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03563002	**Good identification**					**03563002**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03563003	**Good identification**					**03563003**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03563020	**Good identification**					**03563020**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03563021	**Good identification**					**03563021**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03563022	**Good identification**					**03563022**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03563023	**Good identification**					**03563023**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03563100	**Good identification**					**03563100**
Actinomyces naeslundii		97.22	0.51	MNS 88		
Actinomyces israelii		2.78	0.15	FRU 95		
03563101	**Good identification**					**03563101**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03563102	**Acceptable identification**					**03563102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		93.50	0.51	MNS 88			
Actinomyces israelii		4.02	0.18	FRU 95			
Propionibacterium propionicum		2.49	0.14				
03563103	**Good identification**						**03563103**
Actinomyces naeslundii		100.0	0.54	MNS 88			
03563120	**Good identification**						**03563120**
Actinomyces naeslundii		100.0	0.51	MNS 88			
03563121	**Good identification**						**03563121**
Actinomyces naeslundii		100.0	0.54	MNS 88			
03563122	**Good identification**						**03563122**
Actinomyces naeslundii		100.0	0.51	MNS 88			
03563123	**Good identification**						**03563123**
Actinomyces naeslundii		100.0	0.54	MNS 88			
03563140	**Good identification**						**03563140**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03563141	**Good identification**						**03563141**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03563142	**Low discrimination**						**03563142**
Actinomyces naeslundii		86.91	0.38	MNS 88			
Propionibacterium propionicum		13.10	0.14	FRU 95			
03563143	**Good identification**						**03563143**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03563160	**Good identification**						**03563160**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03563161	**Good identification**						**03563161**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03563162	**Good identification**						**03563162**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03563163	**Good identification**						**03563163**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03567102	**Acceptable identification**						**03567102**
Actinomyces israelii		94.92	0.36	SAL 92			
Bifidobacterium adolescentis		5.08	0.08	FRU 99			
03570000	**Low discrimination**						**03570000**
					GRA	CAT	
Fusobacterium mortiferum		83.76	0.45	MLT 11	1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		9.58	0.26		99	90	
Actinomyces naeslundii		5.48	0.36		99	25	
03570001	**Acceptable identification**						**03570001**
					GRA		
Actinomyces naeslundii		90.67	0.39	MNS 88	99		
Fusobacterium mortiferum		9.33	0.11	GAL 96	1		
03570003	**Low discrimination**						**03570003**
Actinomyces naeslundii		87.70	0.39	MNS 88			
Actinomyces viscosus		12.30	0.01	GAL 96			
03570020	**Low discrimination**						**03570020**
					GRA	CAT	
Actinomyces naeslundii		62.83	0.36	MNS 88	99	25	
Propionibacterium granulosum		27.47	0.15	GAL 96	99	90	
Fusobacterium mortiferum		9.70	0.11		1	5	
03570021	**Good identification**						**03570021**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03570022	**Good identification**						**03570022**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03570023	**Good identification**						**03570023**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03570100	**Acceptable identification**						**03570100**
					GRA		
Fusobacterium mortiferum		92.61	0.51	MLT 11	1		
Actinomyces naeslundii		7.39	0.44		99		
03570101	**Acceptable identification**						**03570101**
					GRA		
Actinomyces naeslundii		92.22	0.47	MNS 88	99		
Fusobacterium mortiferum		7.78	0.17	GAL 96	1		
03570102	**Low discrimination**						**03570102**
					GRA		
Fusobacterium mortiferum		48.83	0.35	MLT 11	1		
Actinomyces naeslundii		35.06	0.44	BGL 10	99		
Actinomyces viscosus		13.83	0.13		99		
03570103	**Acceptable identification**						**03570103**
Actinomyces naeslundii		91.94	0.47	MNS 88			
Actinomyces viscosus		8.06	0.05	GAL 96			
03570120	**Low discrimination**						**03570120**
					GRA		
Actinomyces naeslundii		80.46	0.44	MNS 88	99		
Fusobacterium mortiferum		10.19	0.17	GAL 96	1		
Prevotella melaninogenica		9.35	-.01		1		
03570121	**Good identification**						**03570121**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03570122	**Good identification**						**03570122**
Actinomyces naeslundii		100.0	0.44	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03570123	**Good identification**						**03570123**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03571000	**Low discrimination**						**03571000**
					GRA	CAT	
Actinomyces naeslundii		46.58	0.60	MNS 88	99	25	
Fusobacterium mortiferum		44.51	0.48		1	5	
Propionibacterium granulosum		3.40	0.26		99	90	
03571001	**Very good identification**						**03571001**
					GRA		
Actinomyces naeslundii		99.36	0.63	MNS 88	99		
Fusobacterium mortiferum		0.64	0.14		1		
03571002	**Low discrimination**						**03571002**
					GRA		
Actinomyces naeslundii		78.68	0.60	MNS 88	99		
Fusobacterium mortiferum		8.35	0.32		1		
Actinomyces israelii		4.29	0.29		99		
03571003	**Very good identification**						**03571003**
Actinomyces naeslundii		100.0	0.63	MNS 88			
03571010	**Low discrimination**						**03571010**
					GRA	CAT	
Actinomyces naeslundii		66.44	0.38	MNS 88	99	25	
Propionibacterium granulosum		12.55	0.11	MLZ 5	99	90	
Fusobacterium mortiferum		12.19	0.14		1	5	
03571011	**Good identification**						**03571011**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03571012	**Low discrimination**						**03571012**
Actinomyces naeslundii		86.59	0.38	MNS 88			
Actinomyces israelii		13.41	0.15	MLZ 5			
03571013	**Good identification**						**03571013**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03571020	**Good identification**						**03571020**
					CAT		
Actinomyces naeslundii		90.39	0.60	MNS 88	25		
Clostridium perfringens		4.28	0.15		1		
Propionibacterium granulosum		1.65	0.15		90		
03571021	**Very good identification**						**03571021**
Actinomyces naeslundii		100.0	0.63	MNS 88			
03571022	**Good identification**						**03571022**
					GRA		
Actinomyces naeslundii		92.83	0.60	MNS 88	99		
Prevotella oralis		2.59	0.08		1		
Clostridium perfringens		2.07	0.10		99		
03571023	**Very good identification**						**03571023**
Actinomyces naeslundii		100.0	0.63	MNS 88			
03571030	**Good identification**						**03571030**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
03571031	**Good identification**					**03571031**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03571032	**Good identification**					**03571032**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03571033	**Good identification**					**03571033**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03571040	**Low discrimination**					**03571040**
					GRA	
Actinomyces naeslundii		88.12	0.47	MNS 88	99	
Actinomyces israelii		7.06	0.19		99	
Fusobacterium mortiferum		4.82	0.14		1	
03571041	**Good identification**					**03571041**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03571042	**Low discrimination**					**03571042**
Actinomyces naeslundii		89.27	0.47	MNS 88		
Actinomyces israelii		10.73	0.22			
03571043	**Good identification**					**03571043**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03571060	**Good identification**					**03571060**
Actinomyces naeslundii		95.14	0.47	MNS 88		
Clostridium perfringens		4.86	0.03			
03571061	**Good identification**					**03571061**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03571062	**Good identification**					**03571062**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03571063	**Good identification**					**03571063**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03571100	**Low discrimination**					**03571100**
					GRA	
Actinomyces naeslundii		52.63	0.68	MNS 88	99	
Fusobacterium mortiferum		41.23	0.54		1	
Actinomyces israelii		2.69	0.37		99	
03571101	**Very good identification**					**03571101**
					GRA	
Actinomyces naeslundii		99.48	0.71	MNS 88	99	
Fusobacterium mortiferum		0.53	0.20		1	
03571102	**Low discrimination**					**03571102**
					GRA	
Actinomyces naeslundii		76.48	0.68	MNS 88	99	
Fusobacterium mortiferum		6.66	0.38		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		5.86	0.40		99	
03571103	**Very good identification**					**03571103**
Actinomyces naeslundii		99.78	0.71	MNS 88		
Actinomyces viscosus		0.22	0.01			
03571110	**Low discrimination**					**03571110**
					GRA	
Actinomyces naeslundii		77.19	0.46	MNS 88	99	
Fusobacterium mortiferum		11.61	0.20	MLZ 5	1	
Actinomyces israelii		11.20	0.22		99	
03571111	**Good identification**					**03571111**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03571113	**Good identification**					**03571113**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03571120	**Low discrimination**					**03571120**
					GRA	
Actinomyces naeslundii		61.00	0.68	MNS 88	99	
Prevotella melaninogenica		29.24	0.33		1	
Clostridium perfringens		7.44	0.30		99	
03571121	**Very good identification**					**03571121**
					GRA	
Actinomyces naeslundii		99.68	0.71	MNS 88	99	
Prevotella melaninogenica		0.32	-.01		1	
03571122	**Low discrimination**					**03571122**
					GRA	
Actinomyces naeslundii		69.09	0.68	MNS 88	99	
Prevotella oralis		10.61	0.28		1	
Prevotella melaninogenica		6.31	0.21		1	
03571123	**Very good identification**					**03571123**
Actinomyces naeslundii		100.0	0.71	MNS 88		
03571130	**Acceptable identification**					**03571130**
					GRA	
Actinomyces naeslundii		91.58	0.46	MNS 88	99	
Prevotella melaninogenica		8.43	-.01	MLZ 5	1	
03571131	**Good identification**					**03571131**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03571132	**Low discrimination**					**03571132**
					GRA	
Actinomyces naeslundii		58.32	0.46	MNS 88	99	
Parabacteroides distasonis		41.69	0.31	MLZ 5	1	
03571133	**Good identification**					**03571133**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03571140	**Low discrimination**					**03571140**
					GRA	
Actinomyces naeslundii		86.40	0.55	MNS 88	99	
Actinomyces israelii		9.73	0.30		99	
Fusobacterium mortiferum		3.87	0.20		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03571141	**Very good identification**					**03571141**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03571142	**Low discrimination**					**03571142**
Actinomyces naeslundii		80.25	0.55	MNS 88		
Actinomyces israelii		13.56	0.33			
Bifidobacterium adolescentis		2.75	0.14			
03571143	**Very good identification**					**03571143**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03571151	**Good identification**					**03571151**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03571153	**Good identification**					**03571153**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03571160	**Low discrimination**					**03571160**
					GRA	
Actinomyces naeslundii		86.28	0.55	MNS 88	99	
Clostridium perfringens		11.36	0.18		99	
Prevotella melaninogenica		2.37	-.01		1	
03571161	**Very good identification**					**03571161**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03571162	**Acceptable identification**					**03571162**
Actinomyces naeslundii		91.78	0.55	MNS 88		
Clostridium perfringens		5.68	0.12			
Lactobacillus acidophilus		2.53	0.04			
03571163	**Very good identification**					**03571163**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03571171	**Good identification**					**03571171**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03571173	**Good identification**					**03571173**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03571301	**Good identification**					**03571301**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03571303	**Good identification**					**03571303**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03571321	**Good identification**					**03571321**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03571322	**Low discrimination**					**03571322**
					SPO	
Parabacteroides distasonis		67.51	0.43	MNS 98	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		21.19	0.22	XYL 90	1		
Clostridium ramosum		7.09	0.14		92		
03571323	**Good identification**						**03571323**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03571332	**Good identification**						**03571332**
Parabacteroides distasonis		97.91	0.39	MNS 98			
Clostridium clostridioforme		2.09	0.04	XYL 90			
03571500	**Low discrimination**						**03571500**
Actinomyces israelii		39.09	0.35	SAL 92			
Lactobacillus fermentum		27.93	0.32	XYL 92			
Bifidobacterium longum subsp. longum		13.06	0.19				
03571501	**Good identification**						**03571501**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03571502	**Low discrimination**						**03571502**
Bifidobacterium adolescentis		73.70	0.44	MNS 87			
Actinomyces israelii		14.14	0.38	XYL 87			
Lactobacillus fermentum		6.74	0.32				
03571503	**Low discrimination**						**03571503**
Actinomyces naeslundii		82.11	0.37	MNS 88			
Bifidobacterium adolescentis		17.89	0.10	ARA 1			
03571512	**Acceptable identification**						**03571512**
Bifidobacterium adolescentis		91.25	0.46	MNS 87			
Bifidobacterium longum subsp. longum		5.43	0.23	XYL 87			
Actinomyces israelii		2.14	0.24				
03571521	**Good identification**						**03571521**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03571523	**Good identification**						**03571523**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03571722	**Acceptable identification**						**03571722**
Parabacteroides distasonis		93.94	0.40	MNS 98			
Clostridium clostridioforme		6.07	0.14	XYL 90			
03571732	**Low discrimination**						**03571732**
Parabacteroides distasonis		64.67	0.35	MNS 98			
Clostridium clostridioforme		35.33	0.25	XYL 90			
03572000	**Low discrimination**						**03572000**
					GRA	CAT	
Actinomyces naeslundii		56.84	0.41	MNS 88	99	25	
Propionibacterium granulosum		38.64	0.24	GAL 96	99	90	
Fusobacterium mortiferum		4.52	0.11		1	5	
03572001	**Good identification**						**03572001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	MNS 88		
03572002	**Low discrimination**					**03572002**
					CAT	
Actinomyces naeslundii		79.02	0.41	MNS 88	25	
Propionibacterium granulosum		13.43	0.13	GAL 96	90	
Propionibacterium propionicum		7.55	0.14		1	
03572003	**Good identification**					**03572003**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03572020	**Low discrimination**					**03572020**
Actinomyces naeslundii		85.48	0.41	MNS 88		
Propionibacterium granulosum		14.52	0.13	GAL 96		
03572021	**Good identification**					**03572021**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03572022	**Good identification**					**03572022**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03572023	**Good identification**					**03572023**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03572100	**Low discrimination**					**03572100**
					GRA	
Propionibacterium propionicum		66.81	0.45	NIT 86	99	
Actinomyces naeslundii		30.13	0.49	CEL 1	99	
Fusobacterium mortiferum		1.97	0.17		1	
03572101	**Low discrimination**					**03572101**
Actinomyces naeslundii		85.89	0.52	MNS 88		
Propionibacterium propionicum		14.11	0.29	GAL 96		
03572102	**Low discrimination**					**03572102**
Propionibacterium propionicum		75.93	0.48	NIT 86		
Actinomyces naeslundii		22.83	0.49	CEL 1		
Actinomyces israelii		1.24	0.18			
03572103	**Low discrimination**					**03572103**
Actinomyces naeslundii		80.24	0.52	MNS 88		
Propionibacterium propionicum		19.77	0.32	GAL 96		
03572120	**Good identification**					**03572120**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03572121	**Good identification**					**03572121**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03572122	**Good identification**					**03572122**
Actinomyces naeslundii		96.75	0.49	MNS 88		
Propionibacterium propionicum		3.25	0.14	GAL 96		
03572123	**Good identification**					**03572123**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	MNS 88		
03572140	**Acceptable identification**					**03572140**
Propionibacterium propionicum		92.63	0.45	NIT 86		
Actinomyces naeslundii		7.37	0.36	CEL 1		
03572141	**Low discrimination**					**03572141**
Actinomyces naeslundii		51.80	0.39	MNS 88		
Propionibacterium propionicum		48.20	0.29	GAL 96		
03572142	**Acceptable identification**					**03572142**
Propionibacterium propionicum		94.96	0.48	NIT 86		
Actinomyces naeslundii		5.04	0.36	CEL 1		
03572160	**Good identification**					**03572160**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03572161	**Good identification**					**03572161**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03572162	**Low discrimination**					**03572162**
Actinomyces naeslundii		84.01	0.36	MNS 88		
Propionibacterium propionicum		15.99	0.14	GAL 96		
03572163	**Good identification**					**03572163**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03573000	**Low discrimination**					**03573000**
					CAT	
Actinomyces naeslundii		87.76	0.65	MNS 88	25	
Actinomyces israelii		9.32	0.39		1	
Propionibacterium granulosum		2.49	0.24		90	
03573001	**Very good identification**					**03573001**
Actinomyces naeslundii		100.0	0.68	MNS 88		
03573002	**Low discrimination**					**03573002**
					CAT	
Actinomyces naeslundii		85.45	0.65	MNS 88	25	
Actinomyces israelii		13.61	0.42		1	
Propionibacterium granulosum		0.61	0.13		90	
03573003	**Very good identification**					**03573003**
Actinomyces naeslundii		100.0	0.68	MNS 88		
03573010	**Low discrimination**					**03573010**
					CAT	
Actinomyces naeslundii		72.74	0.43	MNS 88	25	
Actinomyces israelii		21.93	0.25	MLZ 5	1	
Propionibacterium granulosum		5.34	0.09		90	
03573011	**Good identification**					**03573011**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03573012	**Low discrimination**					**03573012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		68.87	0.43	MNS 88		
Actinomyces israelii		31.13	0.28	MLZ 5		
03573013	**Good identification**					**03573013**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03573020	**Very good identification**					**03573020**
					CAT	
Actinomyces naeslundii		98.63	0.65	MNS 88	25	
Propionibacterium granulosum		0.70	0.13		90	
Actinomyces israelii		0.67	0.19		1	
03573021	**Very good identification**					**03573021**
Actinomyces naeslundii		100.0	0.68	MNS 88		
03573022	**Very good identification**					**03573022**
Actinomyces naeslundii		98.99	0.65	MNS 88		
Actinomyces israelii		1.01	0.22			
03573023	**Very good identification**					**03573023**
Actinomyces naeslundii		100.0	0.68	MNS 88		
03573030	**Good identification**					**03573030**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03573031	**Good identification**					**03573031**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03573032	**Good identification**					**03573032**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03573033	**Good identification**					**03573033**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03573040	**Low discrimination**					**03573040**
Actinomyces naeslundii		81.04	0.52	MNS 88		
Actinomyces israelii		18.96	0.32			
03573041	**Good identification**					**03573041**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03573042	**Low discrimination**					**03573042**
Actinomyces naeslundii		72.81	0.52	MNS 88		
Actinomyces israelii		25.55	0.35			
Propionibacterium propionicum		1.64	0.14			
03573043	**Good identification**					**03573043**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03573060	**Good identification**					**03573060**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03573061	**Good identification**					**03573061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	MNS 88		
03573062	**Good identification**					**03573062**
Actinomyces naeslundii		97.81	0.52	MNS 88		
Actinomyces israelii		2.19	0.15			
03573063	**Good identification**					**03573063**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03573100	**Low discrimination**					**03573100**
Actinomyces naeslundii		80.28	0.73	MNS 88		
Actinomyces israelii		11.98	0.49			
Propionibacterium propionicum		7.42	0.45			
03573101	**Very good identification**					**03573101**
Actinomyces naeslundii		99.22	0.76	MNS 88		
Propionibacterium propionicum		0.68	0.29			
Actinomyces israelii		0.10	0.15			
03573102	**Low discrimination**					**03573102**
Actinomyces naeslundii		72.31	0.73	MNS 88		
Actinomyces israelii		16.18	0.52			
Propionibacterium propionicum		10.02	0.48			
03573103	**Very good identification**					**03573103**
Actinomyces naeslundii		98.84	0.76	MNS 88		
Propionibacterium propionicum		1.01	0.32			
Actinomyces israelii		0.15	0.18			
03573110	**Low discrimination**					**03573110**
Actinomyces naeslundii		70.25	0.51	MNS 88		
Actinomyces israelii		29.76	0.35	MLZ 5		
03573111	**Good identification**					**03573111**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03573112	**Low discrimination**					**03573112**
Actinomyces naeslundii		46.73	0.51	MNS 88		
Actinomyces israelii		29.69	0.38	MLZ 5		
Bifidobacterium adolescentis		22.34	0.30			
03573113	**Good identification**					**03573113**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03573120	**Very good identification**					**03573120**
					SPO	
Actinomyces naeslundii		97.64	0.73	MNS 88	1	
Clostridium ramosum		1.18	0.15		92	
Actinomyces israelii		0.93	0.29		1	
03573121	**Very good identification**					**03573121**
Actinomyces naeslundii		100.0	0.76	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03573122	**Good identification**					**03573122**
					SPO	
Actinomyces naeslundii		93.97	0.73	MNS 88	1	
Clostridium ramosum		4.56	0.25		92	
Actinomyces israelii		1.34	0.32		1	
03573123	**Very good identification**					**03573123**
Actinomyces naeslundii		100.0	0.76	MNS 88		
03573130	**Good identification**					**03573130**
Actinomyces naeslundii		97.37	0.51	MNS 88		
Actinomyces israelii		2.63	0.15	MLZ 5		
03573131	**Good identification**					**03573131**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03573132	**Good identification**					**03573132**
Actinomyces naeslundii		96.10	0.51	MNS 88		
Actinomyces israelii		3.90	0.18	MLZ 5		
03573133	**Good identification**					**03573133**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03573140	**Low discrimination**					**03573140**
Actinomyces naeslundii		53.99	0.60	MNS 88		
Propionibacterium propionicum		28.26	0.45			
Actinomyces israelii		17.75	0.42			
03573141	**Very good identification**					**03573141**
Actinomyces naeslundii		96.27	0.63	MNS 88		
Propionibacterium propionicum		3.73	0.29			
03573142	**Low discrimination**					**03573142**
Actinomyces naeslundii		43.61	0.60	MNS 88		
Propionibacterium propionicum		34.25	0.48			
Actinomyces israelii		21.51	0.45			
03573143	**Good identification**					**03573143**
Actinomyces naeslundii		94.50	0.63	MNS 88		
Propionibacterium propionicum		5.50	0.32			
03573150	**Low discrimination**					**03573150**
Actinomyces naeslundii		51.72	0.38	MNS 88		
Actinomyces israelii		48.28	0.28	MLZ 5		
03573151	**Good identification**					**03573151**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03573153	**Good identification**					**03573153**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03573160	**Very good identification**					**03573160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.94	0.60	MNS 88		
Actinomyces israelii		2.06	0.22			
03573161	**Very good identification**					**03573161**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03573162	**Very good identification**					**03573162**
Actinomyces naeslundii		96.21	0.60	MNS 88		
Actinomyces israelii		3.03	0.25			
Propionibacterium propionicum		0.76	0.14			
03573163	**Very good identification**					**03573163**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03573170	**Good identification**					**03573170**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03573171	**Good identification**					**03573171**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03573172	**Good identification**					**03573172**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03573173	**Good identification**					**03573173**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03573301	**Good identification**					**03573301**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03573303	**Good identification**					**03573303**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03573321	**Good identification**					**03573321**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03573323	**Good identification**					**03573323**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03573400	**Low discrimination**					**03573400**
Actinomyces israelii		89.58	0.38	SAL 92		
Actinomyces naeslundii		10.42	0.31	XYL 92		
03573402	**Acceptable identification**					**03573402**
Actinomyces israelii		90.26	0.41	SAL 92		
Actinomyces naeslundii		7.00	0.31	XYL 92		
Bifidobacterium adolescentis		2.74	0.08			
03573500	**Acceptable identification**					**03573500**
Actinomyces israelii		91.40	0.48	SAL 92		
Actinomyces naeslundii		7.56	0.39	XYL 92		
Bifidobacterium adolescentis		1.04	0.08			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03573501	**Acceptable identification**					**03573501**
Actinomyces naeslundii		92.47	0.42	MNS 88		
Actinomyces israelii		7.53	0.14	ARA 1		
03573502	**Low discrimination**					**03573502**
Actinomyces israelii		55.09	0.51	SAL 92		
Bifidobacterium adolescentis		41.45	0.43	XYL 92		
Actinomyces naeslundii		3.04	0.39			
03573503	**Low discrimination**					**03573503**
Actinomyces naeslundii		82.38	0.42	MNS 88		
Actinomyces israelii		10.06	0.17	ARA 1		
Bifidobacterium adolescentis		7.57	0.08			
03573512	**Low discrimination**					**03573512**
Bifidobacterium adolescentis		86.02	0.44	MNS 87		
Actinomyces israelii		13.98	0.37	XYL 87		
03573520	**Low discrimination**					**03573520**
Actinomyces naeslundii		56.45	0.39	MNS 88		
Actinomyces israelii		43.55	0.28	ARA 1		
03573521	**Good identification**					**03573521**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03573523	**Good identification**					**03573523**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03573540	**Good identification**					**03573540**
Actinomyces israelii		96.38	0.41	SAL 92		
Actinomyces naeslundii		3.62	0.26	XYL 92		
03573542	**Low discrimination**					**03573542**
Actinomyces israelii		77.89	0.44	SAL 92		
Bifidobacterium adolescentis		18.63	0.27	XYL 92		
Actinomyces naeslundii		1.95	0.26			
03575000	**Low discrimination**					**03575000**
					GRA	
Actinomyces israelii		78.78	0.44	SAL 92	99	
Fusobacterium mortiferum		13.55	0.29		1	
Lactobacillus fermentum		5.77	0.24		99	
03575002	**Acceptable identification**					**03575002**
Actinomyces israelii		91.52	0.47	SAL 92		
Lactobacillus fermentum		4.47	0.24			
Actinomyces naeslundii		1.47	0.26			
03575040	**Good identification**					**03575040**
Actinomyces israelii		100.0	0.37	SAL 92		
03575042	**Good identification**					**03575042**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	SAL 92			
03575100	**Low discrimination**						**03575100**
					COC	GRA	
Actinomyces israelii		80.01	0.55	SAL 92	1	99	
Fusobacterium mortiferum		8.03	0.35		1	1	
Atopobium parvulum		6.82	0.23		99	99	
03575101	**Low discrimination**						**03575101**
Actinomyces naeslundii		71.85	0.37	MNS 88			
Actinomyces israelii		28.15	0.21	XYL 1			
03575102	**Low discrimination**						**03575102**
Actinomyces israelii		69.94	0.58	SAL 92			
Bifidobacterium adolescentis		25.93	0.44				
Lactobacillus fermentum		1.39	0.28				
03575103	**Low discrimination**						**03575103**
Actinomyces naeslundii		55.38	0.37	MNS 88			
Actinomyces israelii		32.55	0.24	XYL 1			
Bifidobacterium adolescentis		12.07	0.10				
03575110	**Good identification**						**03575110**
Actinomyces israelii		95.61	0.41	SAL 92			
Bifidobacterium adolescentis		4.39	0.11	MLZ 13			
03575112	**Low discrimination**						**03575112**
					GRA		
Bifidobacterium adolescentis		74.26	0.46	MNS 87	99		
Actinomyces israelii		24.49	0.44	ARA 87	99		
Parabacteroides distasonis		0.64	0.13		1		
03575121	**Good identification**						**03575121**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03575122	**Low discrimination**						**03575122**
					GRA		
Parabacteroides distasonis		86.51	0.51	MNS 98	1		
Actinomyces israelii		7.74	0.37		99		
Prevotella oralis		2.61	0.13		1		
03575123	**Low discrimination**						**03575123**
					GRA		
Actinomyces naeslundii		70.46	0.37	MNS 88	99		
Parabacteroides distasonis		29.54	0.17	XYL 1	1		
03575132	**Acceptable identification**						**03575132**
					GRA		
Parabacteroides distasonis		94.78	0.47	MNS 98	1		
Actinomyces israelii		2.35	0.23		99		
Bifidobacterium adolescentis		1.13	0.11		99		
03575140	**Good identification**						**03575140**
Actinomyces israelii		100.0	0.48	SAL 92			
03575142	**Low discrimination**						**03575142**
Actinomyces israelii		89.46	0.51	SAL 92			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		10.54	0.29			
03575152	**Low discrimination**					**03575152**
Actinomyces israelii		50.92	0.37	SAL 92		
Bifidobacterium adolescentis		49.08	0.30	MLZ 13		
03575300	**Good identification**					**03575300**
Actinomyces israelii		100.0	0.35	SAL 92		
03575302	**Low discrimination**					**03575302**
					GRA	
Actinomyces israelii		70.54	0.39	SAL 92	99	
Parabacteroides distasonis		20.26	0.25	RHA 7	1	
Clostridium clostridioforme		5.69	0.10		50	
03575322	**Very good identification**					**03575322**
Parabacteroides distasonis		99.07	0.60	MNS 98		
Prevotella oralis		0.43	0.06			
Clostridium clostridioforme		0.28	0.10			
03575332	**Good identification**					**03575332**
Parabacteroides distasonis		97.66	0.55	MNS 98		
Clostridium clostridioforme		2.35	0.22			
03575400	**Acceptable identification**					**03575400**
					SPO	
Actinomyces israelii		90.02	0.43	SAL 92	1	
Lactobacillus fermentum		8.05	0.24		1	
Clostridium butyricum		1.92	-.07		99	
03575402	**Low discrimination**					**03575402**
Actinomyces israelii		83.26	0.46	SAL 92		
Bifidobacterium adolescentis		10.20	0.24			
Lactobacillus fermentum		4.97	0.24			
03575440	**Good identification**					**03575440**
Actinomyces israelii		100.0	0.36	SAL 92		
03575442	**Good identification**					**03575442**
Actinomyces israelii		96.25	0.39	SAL 92		
Bifidobacterium adolescentis		3.75	0.09			
03575500	**Low discrimination**					**03575500**
					SPO	
Actinomyces israelii		77.06	0.53	SAL 92	1	
Clostridium butyricum		9.87	0.17		99	
Bifidobacterium longum subsp. longum		6.72	0.27		1	
03575502	**Low discrimination**					**03575502**
Bifidobacterium adolescentis		72.73	0.58	MNS 87		
Actinomyces israelii		23.98	0.56			
Bifidobacterium longum subsp. longum		1.39	0.27			
03575510	**Low discrimination**					**03575510**
Actinomyces israelii		41.42	0.39	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		41.14	0.31	MLZ 13		
Bifidobacterium adolescentis		15.57	0.26			
03575512	**Good identification**					**03575512**
Bifidobacterium adolescentis		92.04	0.60	MNS 87		
Actinomyces israelii		3.71	0.42			
Bifidobacterium longum subsp. longum		2.46	0.31			
03575520	**Low discrimination**					**03575520**
Bifidobacterium longum subsp. longum		76.60	0.37	MNS 87		
Actinomyces israelii		14.02	0.33	CEL 1		
Lactobacillus fermentum		2.67	0.20			
03575522	**Low discrimination**					**03575522**
					GRA	
Parabacteroides distasonis		60.12	0.48	MNS 98	1	
Bifidobacterium longum subsp. longum		26.16	0.37		99	
Actinomyces israelii		7.18	0.36		99	
03575530	**Good identification**					**03575530**
Bifidobacterium longum subsp. longum		98.42	0.41	MNS 87		
Actinomyces israelii		1.58	0.19	CEL 1		
03575532	**Low discrimination**					**03575532**
					GRA	
Bifidobacterium longum subsp. longum		49.31	0.41	MNS 87	99	
Parabacteroides distasonis		35.84	0.43	CEL 1	1	
Clostridium clostridioforme		9.00	0.27		50	
03575540	**Good identification**					**03575540**
					SPO	
Actinomyces israelii		95.85	0.46	SAL 92	1	
Clostridium butyricum		2.75	-.01		99	
Bifidobacterium adolescentis		1.40	0.09		1	
03575542	**Low discrimination**					**03575542**
Actinomyces israelii		50.92	0.49	SAL 92		
Bifidobacterium adolescentis		49.08	0.43			
03575552	**Low discrimination**					**03575552**
Bifidobacterium adolescentis		88.74	0.44	MNS 87		
Actinomyces israelii		11.26	0.35	SOR 11		
03575712	**Acceptable identification**					**03575712**
Clostridium clostridioforme		90.22	0.42	MNS 99		
Bifidobacterium adolescentis		6.38	0.26	SAL 88		
Actinomyces israelii		1.92	0.23			
03575722	**Low discrimination**					**03575722**
Parabacteroides distasonis		89.46	0.56	MNS 98		
Clostridium clostridioforme		6.49	0.31			
Bacteroides thetaiotaomicron		2.57	0.14			
03575732	**Low discrimination**					**03575732**
Parabacteroides distasonis		60.39	0.51	MNS 98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		37.06	0.42			
Bacteroides thetaiotaomicron		1.25	0.07			
03576102	**Acceptable identification**					**03576102**
Actinomyces israelii		90.30	0.36	SAL 92		
Propionibacterium propionicum		4.86	0.14	GAL 99		
Bifidobacterium adolescentis		4.83	0.08			
03577000	**Very good identification**					**03577000**
Actinomyces israelii		99.18	0.57	SAL 92		
Actinomyces naeslundii		0.82	0.31			
03577002	**Very good identification**					**03577002**
Actinomyces israelii		99.24	0.60	SAL 92		
Actinomyces naeslundii		0.55	0.31			
Bifidobacterium adolescentis		0.22	0.08			
03577010	**Good identification**					**03577010**
Actinomyces israelii		100.0	0.43	SAL 92		
03577012	**Good identification**					**03577012**
Actinomyces israelii		98.26	0.46	SAL 92		
Bifidobacterium adolescentis		1.74	0.10	MLZ 13		
03577020	**Low discrimination**					**03577020**
Actinomyces israelii		88.53	0.37	SAL 92		
Actinomyces naeslundii		11.48	0.31	NAG 6		
03577022	**Acceptable identification**					**03577022**
Actinomyces israelii		92.05	0.40	SAL 92		
Actinomyces naeslundii		7.95	0.31	NAG 6		
03577040	**Good identification**					**03577040**
Actinomyces israelii		100.0	0.50	SAL 92		
03577042	**Good identification**					**03577042**
Actinomyces israelii		100.0	0.53	SAL 92		
03577050	**Good identification**					**03577050**
Actinomyces israelii		100.0	0.36	SAL 92		
03577052	**Good identification**					**03577052**
Actinomyces israelii		100.0	0.39	SAL 92		
03577100	**Very good identification**					**03577100**
Actinomyces israelii		99.34	0.68	SAL 92		
Actinomyces naeslundii		0.59	0.39			
Bifidobacterium adolescentis		0.08	0.08			
03577102	**Good identification**					**03577102**
Actinomyces israelii		94.52	0.71	SAL 92		
Bifidobacterium adolescentis		5.06	0.43			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.37	0.39			
03577103	**Low discrimination**					**03577103**
Actinomyces israelii		61.11	0.36	SAL 92		
Actinomyces naeslundii		35.62	0.42	URE 1		
Bifidobacterium adolescentis		3.27	0.08			
03577110	**Good identification**					**03577110**
Actinomyces israelii		99.34	0.53	SAL 92		
Bifidobacterium adolescentis		0.66	0.10	MLZ 13		
03577112	**Low discrimination**					**03577112**
Actinomyces israelii		69.55	0.57	SAL 92		
Bifidobacterium adolescentis		30.45	0.44	MLZ 13		
03577120	**Acceptable identification**					**03577120**
Actinomyces israelii		91.56	0.47	SAL 92		
Actinomyces naeslundii		8.44	0.39	NAG 6		
03577121	**Good identification**					**03577121**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03577122	**Acceptable identification**					**03577122**
					GRA	
Actinomyces israelii		91.75	0.50	SAL 92	99	
Actinomyces naeslundii		5.64	0.39	NAG 6	99	
Parabacteroides distasonis		1.83	0.17		1	
03577123	**Acceptable identification**					**03577123**
Actinomyces naeslundii		90.13	0.42	MNS 88		
Actinomyces israelii		9.87	0.16	XYL 1		
03577132	**Low discrimination**					**03577132**
					GRA	
Actinomyces israelii		87.64	0.36	SAL 92	99	
Parabacteroides distasonis		6.29	0.13	MLZ 13	1	
Bifidobacterium adolescentis		6.07	0.10	NAG 6	99	
03577140	**Very good identification**					**03577140**
Actinomyces israelii		99.73	0.61	SAL 92		
Actinomyces naeslundii		0.27	0.26			
03577142	**Very good identification**					**03577142**
Actinomyces israelii		98.02	0.64	SAL 92		
Bifidobacterium adolescentis		1.67	0.27			
Actinomyces naeslundii		0.18	0.26			
03577150	**Good identification**					**03577150**
Actinomyces israelii		100.0	0.46	SAL 92		
03577152	**Low discrimination**					**03577152**
Actinomyces israelii		87.78	0.49	SAL 92		
Bifidobacterium adolescentis		12.22	0.29	MLZ 13		
03577160	**Good identification**					**03577160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		95.98	0.40	SAL 92		
Actinomyces naeslundii		4.02	0.26	NAG 6		
03577162	**Good identification**					**03577162**
Actinomyces israelii		97.29	0.43	SAL 92		
Actinomyces naeslundii		2.71	0.26	NAG 6		
03577200	**Good identification**					**03577200**
Actinomyces israelii		100.0	0.38	SAL 92		
03577202	**Good identification**					**03577202**
Actinomyces israelii		100.0	0.41	SAL 92		
03577300	**Good identification**					**03577300**
Actinomyces israelii		100.0	0.48	SAL 92		
03577302	**Good identification**					**03577302**
Actinomyces israelii		99.29	0.51	SAL 92		
Bifidobacterium adolescentis		0.71	0.08	RHA 7		
03577312	**Acceptable identification**					**03577312**
Actinomyces israelii		94.45	0.37	SAL 92		
Bifidobacterium adolescentis		5.55	0.10	RHA 7		
03577340	**Good identification**					**03577340**
Actinomyces israelii		100.0	0.41	SAL 92		
03577342	**Good identification**					**03577342**
Actinomyces israelii		100.0	0.44	SAL 92		
03577400	**Good identification**					**03577400**
Actinomyces israelii		100.0	0.56	SAL 92		
03577402	**Very good identification**					**03577402**
Actinomyces israelii		98.26	0.59	SAL 92		
Bifidobacterium adolescentis		1.74	0.23			
03577410	**Good identification**					**03577410**
Actinomyces israelii		100.0	0.42	SAL 92		
03577412	**Low discrimination**					**03577412**
Actinomyces israelii		87.36	0.45	SAL 92		
Bifidobacterium adolescentis		12.64	0.24	MLZ 13		
03577420	**Good identification**					**03577420**
Actinomyces israelii		100.0	0.35	SAL 92		
03577422	**Good identification**					**03577422**
Actinomyces israelii		100.0	0.38	SAL 92		
03577440	**Good identification**					**03577440**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.49	SAL 92		
03577442	**Good identification**					**03577442**
Actinomyces israelii		99.44	0.52	SAL 92		
Bifidobacterium adolescentis		0.56	0.07			
03577452	**Good identification**					**03577452**
Actinomyces israelii		95.60	0.38	SAL 92		
Bifidobacterium adolescentis		4.40	0.09	MLZ 13		
03577500	**Very good identification**					**03577500**
					SPO	
Actinomyces israelii		99.09	0.66	SAL 92	1	
Bifidobacterium adolescentis		0.66	0.23		1	
Clostridium butyricum		0.25	0.01		99	
03577502	**Low discrimination**					**03577502**
					SPO	
Actinomyces israelii		69.52	0.69	SAL 92	1	
Bifidobacterium adolescentis		30.44	0.57		1	
Clostridium butyricum		0.05	-.07		99	
03577510	**Acceptable identification**					**03577510**
Actinomyces israelii		94.85	0.52	SAL 92		
Bifidobacterium adolescentis		5.15	0.24	MLZ 13		
03577512	**Low discrimination**					**03577512**
Bifidobacterium adolescentis		78.06	0.58	MNS 87		
Actinomyces israelii		21.80	0.55			
Bifidobacterium dentium		0.14	-.07			
03577520	**Good identification**					**03577520**
Actinomyces israelii		100.0	0.46	SAL 92		
03577522	**Acceptable identification**					**03577522**
					GRA	
Actinomyces israelii		92.23	0.49	SAL 92	99	
Bifidobacterium adolescentis		6.39	0.23	NAG 6	99	
Parabacteroides distasonis		1.38	0.14		1	
03577540	**Very good identification**					**03577540**
Actinomyces israelii		99.79	0.59	SAL 92		
Bifidobacterium adolescentis		0.21	0.07			
03577542	**Low discrimination**					**03577542**
Actinomyces israelii		87.78	0.62	SAL 92		
Bifidobacterium adolescentis		12.22	0.41			
03577550	**Good identification**					**03577550**
					COC	
Actinomyces israelii		95.97	0.45	SAL 92	1	
Blautia producta		2.38	-.03	MLZ 13	99	
Bifidobacterium adolescentis		1.66	0.09		1	
03577552	**Low discrimination**					**03577552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		53.24	0.43	MNS 87			
Actinomyces israelii		46.77	0.48	SOR 11			
03577560	**Good identification**						**03577560**
Actinomyces israelii		100.0	0.39	SAL 92			
03577562	**Good identification**						**03577562**
Actinomyces israelii		97.85	0.42	SAL 92			
Bifidobacterium adolescentis		2.16	0.07	NAG 6			
03577600	**Good identification**						**03577600**
Actinomyces israelii		100.0	0.37	SAL 92			
03577602	**Good identification**						**03577602**
Actinomyces israelii		100.0	0.40	SAL 92			
03577700	**Good identification**						**03577700**
Actinomyces israelii		100.0	0.47	SAL 92			
03577702	**Acceptable identification**						**03577702**
Actinomyces israelii		94.45	0.50	SAL 92			
Bifidobacterium adolescentis		5.55	0.23	RHA 7			
03577712	**Low discrimination**						**03577712**
Actinomyces israelii		63.92	0.36	SAL 92			
Bifidobacterium adolescentis		30.72	0.24	RHA 7			
Clostridium clostridioforme		5.37	0.08	MLZ 13			
03577740	**Good identification**						**03577740**
Actinomyces israelii		100.0	0.40	SAL 92			
03577742	**Good identification**						**03577742**
Actinomyces israelii		98.17	0.43	SAL 92			
Bifidobacterium adolescentis		1.83	0.07	RHA 7			
03600000	**Low discrimination**						**03600000**
					GRA	SPO	
Fusobacterium mortiferum		71.77	0.49	FRU 99	1	1	
Eubacterium tortuosum		23.24	0.24		99	1	
Clostridium botulinum biovar A		1.75	0.08		99	99	
03600100	**Good identification**						**03600100**
Fusobacterium mortiferum		100.0	0.55	FRU 99			
03600102	**Good identification**						**03600102**
Fusobacterium mortiferum		100.0	0.39	FRU 99			
03601000	**Very good identification**						**03601000**
					GRA		
Eubacterium tortuosum		95.48	0.58	ESL 99	99		
Fusobacterium mortiferum		4.47	0.52		1		
Lactobacillus fermentum		0.05	0.20		99		
03601002	**Good identification**						**03601002**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium tortuosum		96.85	0.45	ESL 99	99			
Fusobacterium mortiferum		2.85	0.35		1			
Lactobacillus fermentum		0.29	0.20		99			
03601020	**Good identification**							**03601020**
					GRA			
Eubacterium tortuosum		99.73	0.45	ESL 99	99			
Fusobacterium mortiferum		0.27	0.17		1			
03601100	**Good identification**							**03601100**
					GRA			
Fusobacterium mortiferum		90.11	0.58	FRU 99	1			
Eubacterium tortuosum		8.34	0.24		99			
Lactobacillus fermentum		0.72	0.24		99			
03601102	**Low discrimination**							**03601102**
					GRA			
Fusobacterium mortiferum		79.86	0.42	FRU 99	1			
Eubacterium tortuosum		11.74	0.11	BGL 10	99			
Lactobacillus fermentum		5.74	0.24		99			
03603101	**Good identification**							**03603101**
Actinomyces naeslundii		100.0	0.36	MNS 88				
03603103	**Good identification**							**03603103**
Actinomyces naeslundii		100.0	0.36	MNS 88				
03603121	**Good identification**							**03603121**
Actinomyces naeslundii		100.0	0.36	MNS 88				
03603123	**Good identification**							**03603123**
Actinomyces naeslundii		100.0	0.36	MNS 88				
03604100	**Acceptable identification**							**03604100**
					GRA			
Fusobacterium mortiferum		94.02	0.36	FRU 99	1			
Actinomyces meyeri		5.98	0.06	XYL 7	99			
03605100	**Low discrimination**							**03605100**
					GRA			
Fusobacterium mortiferum		88.82	0.39	FRU 99	1			
Actinomyces meyeri		5.65	0.09	XYL 7	99			
Lactobacillus fermentum		5.53	0.20		99			
03610000	**Very good identification**							**03610000**
					COC	GRA	CAT	
Fusobacterium mortiferum		99.64	0.83		1	1	5	
Megasphaera elsdenii		0.23	0.29		99	1	1	
Propionibacterium granulosum		0.05	0.24		1	99	90	
03610001	**Good identification**							**03610001**
Fusobacterium mortiferum		100.0	0.49	URE 1				
03610002	**Very good identification**							**03610002**
					COC	GRA		
Fusobacterium mortiferum		99.44	0.66	BGL 10	1	1		
Megasphaera elsdenii		0.23	0.12		99	1		
Lactobacillus fermentum		0.16	0.20		1	99		
03610010	**Good identification**							**03610010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Fusobacterium mortiferum		99.28	0.49	MLZ 1	1	5	
Propionibacterium granulosum		0.72	0.09		99	90	
03610020	**Good identification**						**03610020**
					GRA	CAT	
Fusobacterium mortiferum		98.69	0.49	NAG 1	1	5	
Propionibacterium granulosum		1.31	0.13		99	90	
03610040	**Low discrimination**						**03610040**
					COC		
Fusobacterium mortiferum		86.95	0.49	SOR 1	1		
Megasphaera elsdenii		13.05	0.26		99		
03610100	**Very good identification**						**03610100**
					GRA		
Fusobacterium mortiferum		99.94	0.89		1		
Actinomyces meyeri		0.05	0.23		99		
Lactobacillus fermentum		0.01	0.24		99		
03610101	**Good identification**						**03610101**
					GRA		
Fusobacterium mortiferum		99.34	0.55	URE 1	1		
Actinomyces naeslundii		0.66	0.29		99		
03610102	**Very good identification**						**03610102**
					GRA		
Fusobacterium mortiferum		99.83	0.73	BGL 10	1		
Lactobacillus fermentum		0.11	0.24		99		
Actinomyces naeslundii		0.04	0.26		99		
03610103	**Acceptable identification**						**03610103**
					GRA		
Fusobacterium mortiferum		94.39	0.39	URE 1	1		
Actinomyces naeslundii		5.61	0.29	BGL 10	99		
03610110	**Good identification**						**03610110**
Fusobacterium mortiferum		100.0	0.55	MLZ 1			
03610112	**Good identification**						**03610112**
Fusobacterium mortiferum		100.0	0.39	MLZ 1			
03610120	**Good identification**						**03610120**
					GRA		
Fusobacterium mortiferum		98.54	0.55	NAG 1	1		
Actinomyces meyeri		0.63	0.08		99		
Actinomyces naeslundii		0.43	0.26		99		
03610122	**Acceptable identification**						**03610122**
					GRA		
Fusobacterium mortiferum		92.98	0.39	NAG 1	1		
Actinomyces naeslundii		3.69	0.26	BGL 10	99		
Lactobacillus fermentum		3.34	0.15		99		
03610140	**Good identification**						**03610140**
Fusobacterium mortiferum		100.0	0.55	SOR 1			
03610142	**Good identification**						**03610142**
Fusobacterium mortiferum		100.0	0.39	SOR 1			
03610200	**Good identification**						**03610200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		99.33	0.49	RHA 1	1		
Actinomyces odontolyticus		0.67	0.06		99		
03610300	**Good identification**						**03610300**
Fusobacterium mortiferum		100.0	0.55	RHA 1			
03610302	**Good identification**						**03610302**
Fusobacterium mortiferum		100.0	0.39	RHA 1			
03610400	**Good identification**						**03610400**
					GRA		
Fusobacterium mortiferum		97.66	0.49	ARA 1	1		
Lactobacillus fermentum		1.67	0.20		99		
Actinomyces meyeri		0.67	0.02		99		
03610500	**Good identification**						**03610500**
					GRA		
Fusobacterium mortiferum		97.69	0.55	ARA 1	1		
Lactobacillus fermentum		1.17	0.24		99		
Actinomyces meyeri		1.14	0.13		99		
03610502	**Acceptable identification**						**03610502**
					GRA		
Fusobacterium mortiferum		90.28	0.39	ARA 1	1		
Lactobacillus fermentum		9.73	0.24	BGL 10	99		
03611000	**Very good identification**						**03611000**
					GRA		
Fusobacterium mortiferum		98.54	0.86		1		
Lactobacillus fermentum		1.12	0.54		99		
Eubacterium tortuosum		0.22	0.24		99		
03611001	**Acceptable identification**						**03611001**
					GRA		
Fusobacterium mortiferum		91.07	0.52	URE 1	1		
Actinomyces naeslundii		7.89	0.45		99		
Lactobacillus fermentum		1.04	0.20		99		
03611002	**Low discrimination**						**03611002**
					GRA		
Fusobacterium mortiferum		89.91	0.69	BGL 10	1		
Lactobacillus fermentum		9.23	0.54		99		
Actinomyces naeslundii		0.47	0.42		99		
03611003	**Low discrimination**						**03611003**
					GRA		
Fusobacterium mortiferum		53.12	0.35	URE 1	1		
Actinomyces naeslundii		41.43	0.45	BGL 10	99		
Lactobacillus fermentum		5.46	0.20		99		
03611010	**Good identification**						**03611010**
					GRA	CAT	
Fusobacterium mortiferum		98.40	0.52	MLZ 1	1	5	
Lactobacillus fermentum		1.12	0.20		99	1	
Propionibacterium granulosum		0.48	0.09		99	90	
03611012	**Acceptable identification**						**03611012**
					GRA		
Fusobacterium mortiferum		90.69	0.35	MLZ 1	1		
Lactobacillus fermentum		9.31	0.20	BGL 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03611020	**Low discrimination**					**03611020**
					GRA	
Fusobacterium mortiferum		67.51	0.52	NAG 1	1	
Lactobacillus fermentum		25.42	0.46		99	
Actinomyces naeslundii		3.90	0.42		99	
03611021	**Low discrimination**					**03611021**
					GRA	
Actinomyces naeslundii		89.56	0.45	MNS 88	99	
Fusobacterium mortiferum		10.44	0.17	MLT 94	1	
03611022	**Low discrimination**					**03611022**
					GRA	
Lactobacillus fermentum		68.20	0.46	SAL 1	99	
Fusobacterium mortiferum		20.12	0.35		1	
Actinomyces naeslundii		10.46	0.42		99	
03611023	**Good identification**					**03611023**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03611040	**Good identification**					**03611040**
					GRA	
Fusobacterium mortiferum		97.89	0.52	SOR 1	1	
Lactobacillus fermentum		1.12	0.20		99	
Actinomyces naeslundii		1.00	0.29		99	
03611042	**Low discrimination**					**03611042**
					GRA	
Fusobacterium mortiferum		83.72	0.35	SOR 1	1	
Lactobacillus fermentum		8.60	0.20	BGL 10	99	
Actinomyces naeslundii		7.68	0.29		99	
03611100	**Excellent identification**					**03611100**
					GRA	
Fusobacterium mortiferum		99.09	0.92		1	
Lactobacillus fermentum		0.79	0.58		99	
Actinomyces naeslundii		0.07	0.50		99	
03611101	**Low discrimination**					**03611101**
					GRA	
Fusobacterium mortiferum		89.79	0.58	URE 1	1	
Actinomyces naeslundii		9.49	0.53		99	
Lactobacillus fermentum		0.72	0.24		99	
03611102	**Good identification**					**03611102**
					GRA	
Fusobacterium mortiferum		92.73	0.76	BGL 10	1	
Lactobacillus fermentum		6.66	0.58		99	
Actinomyces naeslundii		0.59	0.50		99	
03611103	**Low discrimination**					**03611103**
					GRA	
Fusobacterium mortiferum		49.43	0.42	URE 1	1	
Actinomyces naeslundii		47.02	0.53	BGL 10	99	
Lactobacillus fermentum		3.55	0.24		99	
03611110	**Very good identification**					**03611110**
					GRA	
Fusobacterium mortiferum		98.85	0.58	MLZ 1	1	
Lactobacillus fermentum		0.79	0.24		99	
Actinomyces naeslundii		0.37	0.28		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03611112	**Low discrimination**					**03611112**
					GRA	
Fusobacterium mortiferum		89.08	0.42	MLZ 1	1	
Lactobacillus fermentum		6.40	0.24	BGL 10	99	
Actinomyces naeslundii		2.97	0.28		99	
03611120	**Low discrimination**					**03611120**
					GRA	
Fusobacterium mortiferum		72.56	0.58	NAG 1	1	
Lactobacillus fermentum		19.11	0.50		99	
Actinomyces naeslundii		5.11	0.50		99	
03611121	**Low discrimination**					**03611121**
					GRA	
Actinomyces naeslundii		89.23	0.53	MNS 88	99	
Fusobacterium mortiferum		8.53	0.24	MLT 94	1	
Lactobacillus fermentum		2.25	0.15		99	
03611122	**Low discrimination**					**03611122**
					GRA	
Lactobacillus fermentum		55.86	0.50	SAL 1	99	
Fusobacterium mortiferum		23.57	0.42		1	
Actinomyces naeslundii		14.95	0.50		99	
03611123	**Good identification**					**03611123**
Actinomyces naeslundii		97.55	0.53	MNS 88		
Lactobacillus fermentum		2.46	0.15	MLT 94		
03611140	**Very good identification**					**03611140**
					GRA	
Fusobacterium mortiferum		98.00	0.58	SOR 1	1	
Actinomyces naeslundii		1.22	0.37		99	
Lactobacillus fermentum		0.78	0.24		99	
03611141	**Low discrimination**					**03611141**
					GRA	
Actinomyces naeslundii		64.87	0.40	MNS 88	99	
Fusobacterium mortiferum		35.13	0.24	MLT 94	1	
03611142	**Low discrimination**					**03611142**
					GRA	
Fusobacterium mortiferum		84.48	0.42	SOR 1	1	
Actinomyces naeslundii		9.45	0.37	BGL 10	99	
Lactobacillus fermentum		6.07	0.24		99	
03611143	**Good identification**					**03611143**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03611160	**Low discrimination**					**03611160**
					GRA	
Actinomyces naeslundii		49.35	0.37	MNS 88	99	
Fusobacterium mortiferum		40.09	0.24	MLT 94	1	
Lactobacillus fermentum		10.56	0.15		99	
03611161	**Good identification**					**03611161**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03611162	**Low discrimination**					**03611162**
Actinomyces naeslundii		82.38	0.37	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		17.62	0.15	MLT 94		
03611163	**Good identification**					**03611163**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03611200	**Good identification**					**03611200**
					GRA	
Fusobacterium mortiferum		98.87	0.52	RHA 1	1	
Lactobacillus fermentum		1.13	0.20		99	
03611202	**Acceptable identification**					**03611202**
					GRA	
Fusobacterium mortiferum		90.69	0.35	RHA 1	1	
Lactobacillus fermentum		9.31	0.20	BGL 10	99	
03611300	**Very good identification**					**03611300**
					GRA	
Fusobacterium mortiferum		99.21	0.58	RHA 1	1	
Lactobacillus fermentum		0.79	0.24		99	
03611302	**Acceptable identification**					**03611302**
					GRA	
Fusobacterium mortiferum		93.30	0.42	RHA 1	1	
Lactobacillus fermentum		6.70	0.24	BGL 10	99	
03611400	**Low discrimination**					**03611400**
					GRA	
Lactobacillus fermentum		52.88	0.54	SAL 1	99	
Fusobacterium mortiferum		46.81	0.52		1	
Actinomyces meyeri		0.32	0.05		99	
03611402	**Acceptable identification**					**03611402**
					GRA	
Lactobacillus fermentum		91.05	0.54	SAL 1	99	
Fusobacterium mortiferum		8.96	0.35		1	
03611420	**Good identification**					**03611420**
					GRA	
Lactobacillus fermentum		97.39	0.46	SAL 1	99	
Fusobacterium mortiferum		2.61	0.17		1	
03611422	**Good identification**					**03611422**
Lactobacillus fermentum		100.0	0.46	SAL 1		
03611500	**Low discrimination**					**03611500**
					GRA	
Fusobacterium mortiferum		55.51	0.58	ARA 1	1	
Lactobacillus fermentum		43.85	0.58		99	
Actinomyces meyeri		0.65	0.16		99	
03611502	**Low discrimination**					**03611502**
					GRA	
Lactobacillus fermentum		86.65	0.58	SAL 1	99	
Fusobacterium mortiferum		12.19	0.42		1	
Bifidobacterium adolescentis		1.16	0.20		99	
03611520	**Good identification**					**03611520**
					GRA	
Lactobacillus fermentum		95.67	0.50	SAL 1	99	
Fusobacterium mortiferum		3.67	0.24		1	
Bifidobacterium longum subsp. longum		0.66	0.05		99	
03611522	**Good identification**					**03611522**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Lactobacillus fermentum		97.74	0.50	SAL 1	99			
Parabacteroides distasonis		1.59	0.16		1			
Bifidobacterium longum subsp. longum		0.67	0.05		99			
03612000	**Low discrimination**							**03612000**
					COC	GRA	CAT	
Fusobacterium mortiferum		77.59	0.49	MAN 1	1	1	5	
Megasphaera elsdenii		17.48	0.29		99	1	1	
Propionibacterium granulosum		3.12	0.21		1	99	90	
03612100	**Good identification**							**03612100**
					GRA			
Fusobacterium mortiferum		98.89	0.55	MAN 1	1			
Actinomyces naeslundii		0.85	0.31		99			
Propionibacterium propionicum		0.26	0.12		99			
03612102	**Low discrimination**							**03612102**
					GRA			
Fusobacterium mortiferum		89.86	0.39	MAN 1	1			
Actinomyces naeslundii		6.91	0.31	BGL 10	99			
Propionibacterium propionicum		3.23	0.15		99			
03613000	**Low discrimination**							**03613000**
					GRA	CAT		
Fusobacterium mortiferum		86.93	0.52	MAN 1	1	5		
Actinomyces naeslundii		9.75	0.47		99	25		
Propionibacterium granulosum		2.33	0.21		99	90		
03613001	**Acceptable identification**							**03613001**
					GRA			
Actinomyces naeslundii		94.34	0.50	MNS 88	99			
Fusobacterium mortiferum		5.67	0.17	MLT 94	1			
03613002	**Low discrimination**							**03613002**
					GRA			
Actinomyces naeslundii		46.46	0.47	MNS 88	99			
Fusobacterium mortiferum		46.04	0.35	MLT 94	1			
Lactobacillus fermentum		4.73	0.20		99			
03613003	**Good identification**							**03613003**
Actinomyces naeslundii		100.0	0.50	MNS 88				
03613020	**Low discrimination**							**03613020**
					GRA	CAT		
Actinomyces naeslundii		86.97	0.47	MNS 88	99	25		
Fusobacterium mortiferum		7.83	0.17	MLT 94	1	5		
Propionibacterium granulosum		5.20	0.11		99	90		
03613021	**Good identification**							**03613021**
Actinomyces naeslundii		100.0	0.50	MNS 88				
03613022	**Good identification**							**03613022**
Actinomyces naeslundii		100.0	0.47	MNS 88				
03613023	**Good identification**							**03613023**
Actinomyces naeslundii		100.0	0.50	MNS 88				
03613041	**Good identification**							**03613041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	MNS 88		
03613043	**Good identification**					**03613043**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03613061	**Good identification**					**03613061**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03613063	**Good identification**					**03613063**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03613100	**Low discrimination**					**03613100**
					GRA	
Fusobacterium mortiferum		87.22	0.58	MAN 1	1	
Actinomyces naeslundii		11.93	0.55		99	
Lactobacillus fermentum		0.70	0.24		99	
03613101	**Very good identification**					**03613101**
					GRA	
Actinomyces naeslundii		95.31	0.58	MNS 88	99	
Fusobacterium mortiferum		4.69	0.24	MLT 94	1	
03613102	**Low discrimination**					**03613102**
					GRA	
Actinomyces naeslundii		52.91	0.55	MNS 88	99	
Fusobacterium mortiferum		42.98	0.42	MLT 94	1	
Lactobacillus fermentum		3.09	0.24		99	
03613103	**Very good identification**					**03613103**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03613111	**Good identification**					**03613111**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03613113	**Good identification**					**03613113**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03613120	**Acceptable identification**					**03613120**
					GRA	
Actinomyces naeslundii		91.47	0.55	MNS 88	99	
Fusobacterium mortiferum		6.75	0.24	MLT 94	1	
Lactobacillus fermentum		1.78	0.15		99	
03613121	**Very good identification**					**03613121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03613122	**Good identification**					**03613122**
Actinomyces naeslundii		98.09	0.55	MNS 88		
Lactobacillus fermentum		1.91	0.15	MLT 94		
03613123	**Very good identification**					**03613123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03613131	**Good identification**					**03613131**
Actinomyces naeslundii		100.0	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03613133	**Good identification**					**03613133**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03613140	**Low discrimination**					**03613140**
					GRA	
Actinomyces naeslundii		67.03	0.42	MNS 88	99	
Fusobacterium mortiferum		28.05	0.24	MLT 94	1	
Propionibacterium propionicum		4.92	0.12		99	
03613141	**Good identification**					**03613141**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03613142	**Acceptable identification**					**03613142**
Actinomyces naeslundii		90.07	0.42	MNS 88		
Propionibacterium propionicum		9.93	0.15	MLT 94		
03613143	**Good identification**					**03613143**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03613160	**Good identification**					**03613160**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03613161	**Good identification**					**03613161**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03613162	**Good identification**					**03613162**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03613163	**Good identification**					**03613163**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03614000	**Very good identification**					**03614000**
					GRA	
Fusobacterium mortiferum		99.16	0.64	XYL 7	1	
Actinomyces meyeri		0.71	0.18		99	
Lactobacillus fermentum		0.13	0.16		99	
03614002	**Good identification**					**03614002**
					GRA	
Fusobacterium mortiferum		98.80	0.47	XYL 7	1	
Lactobacillus fermentum		1.20	0.16	BGL 10	99	
03614100	**Very good identification**					**03614100**
					GRA	
Fusobacterium mortiferum		98.70	0.70	XYL 7	1	
Actinomyces meyeri		1.20	0.28		99	
Lactobacillus fermentum		0.09	0.20		99	
03614101	**Good identification**					**03614101**
Fusobacterium mortiferum		100.0	0.36	XYL 7		
03614102	**Good identification**					**03614102**
					GRA	
Fusobacterium mortiferum		98.83	0.54	XYL 7	1	
Lactobacillus fermentum		0.84	0.20	BGL 10	99	
Actinomyces meyeri		0.34	0.02		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03614110	**Good identification**					**03614110**
Fusobacterium mortiferum		100.0	0.36	XYL 7		
03614120	**Low discrimination**					**03614120**
					GRA	
Fusobacterium mortiferum		85.86	0.36	XYL 7	1	
Actinomyces meyeri		14.14	0.13	NAG 1	99	
03614140	**Good identification**					**03614140**
Fusobacterium mortiferum		100.0	0.36	XYL 7		
03614300	**Good identification**					**03614300**
Fusobacterium mortiferum		100.0	0.36	XYL 7		
03614500	**Low discrimination**					**03614500**
					GRA	
Fusobacterium mortiferum		71.67	0.36	XYL 7	1	
Actinomyces meyeri		21.64	0.18	ARA 1	99	
Lactobacillus fermentum		6.69	0.20		99	
03615000	**Good identification**					**03615000**
					GRA	
Fusobacterium mortiferum		91.23	0.67	XYL 7	1	
Lactobacillus fermentum		8.12	0.50		99	
Actinomyces meyeri		0.65	0.21		99	
03615002	**Low discrimination**					**03615002**
					GRA	
Fusobacterium mortiferum		55.52	0.50	XYL 7	1	
Lactobacillus fermentum		44.49	0.50	BGL 10	99	
03615020	**Low discrimination**					**03615020**
					GRA	
Lactobacillus fermentum		72.83	0.42	SAL 1	99	
Fusobacterium mortiferum		24.79	0.32		1	
Actinomyces meyeri		2.38	0.06		99	
03615022	**Good identification**					**03615022**
					GRA	
Lactobacillus fermentum		96.36	0.42	SAL 1	99	
Fusobacterium mortiferum		3.64	0.16		1	
03615100	**Good identification**					**03615100**
					GRA	
Fusobacterium mortiferum		93.07	0.73	XYL 7	1	
Lactobacillus fermentum		5.79	0.54		99	
Actinomyces meyeri		1.14	0.31		99	
03615101	**Acceptable identification**					**03615101**
					GRA	
Fusobacterium mortiferum		94.14	0.39	XYL 7	1	
Lactobacillus fermentum		5.86	0.20	URE 1	99	
03615102	**Low discrimination**					**03615102**
					GRA	
Fusobacterium mortiferum		63.43	0.57	XYL 7	1	
Lactobacillus fermentum		35.54	0.54	BGL 10	99	
Bifidobacterium adolescentis		0.81	0.20		99	
03615110	**Acceptable identification**					**03615110**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		94.14	0.39	XYL 7	1	
Lactobacillus fermentum		5.86	0.20	MLZ 1	99	
03615120	**Low discrimination**					**03615120**
					GRA	
Lactobacillus fermentum		63.82	0.46	SAL 1	99	
Fusobacterium mortiferum		31.06	0.39		1	
Actinomyces meyeri		5.12	0.16		99	
03615122	**Low discrimination**					**03615122**
					GRA	
Lactobacillus fermentum		67.73	0.46	SAL 1	99	
Parabacteroides distasonis		27.48	0.36		1	
Fusobacterium mortiferum		3.66	0.22		1	
03615140	**Acceptable identification**					**03615140**
					GRA	
Fusobacterium mortiferum		94.14	0.39	XYL 7	1	
Lactobacillus fermentum		5.86	0.20	SOR 1	99	
03615322	**Good identification**					**03615322**
					GRA	
Parabacteroides distasonis		98.59	0.44	MNS 98	1	
Eubacterium contortum		1.41	0.01		99	
03615332	**Good identification**					**03615332**
Parabacteroides distasonis		100.0	0.40	MNS 98		
03615400	**Low discrimination**					**03615400**
					GRA	
Lactobacillus fermentum		88.23	0.50	SAL 1	99	
Fusobacterium mortiferum		10.01	0.32		1	
Actinomyces meyeri		1.76	0.10		99	
03615402	**Good identification**					**03615402**
					GRA	
Lactobacillus fermentum		98.76	0.50	SAL 1	99	
Fusobacterium mortiferum		1.25	0.16		1	
03615420	**Good identification**					**03615420**
Lactobacillus fermentum		100.0	0.42	SAL 1		
03615422	**Good identification**					**03615422**
Lactobacillus fermentum		100.0	0.42	SAL 1		
03615500	**Low discrimination**					**03615500**
					GRA	
Lactobacillus fermentum		82.56	0.54	SAL 1	99	
Fusobacterium mortiferum		13.40	0.39		1	
Actinomyces meyeri		4.04	0.21		99	
03615502	**Low discrimination**					**03615502**
					GRA	
Lactobacillus fermentum		85.42	0.54	SAL 1	99	
Bifidobacterium adolescentis		13.04	0.34		99	
Fusobacterium mortiferum		1.54	0.22		1	
03615512	**Acceptable identification**					**03615512**
Bifidobacterium adolescentis		92.52	0.36	MNS 87		
Lactobacillus fermentum		5.01	0.20	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		2.47	0.07			
03615520	**Acceptable identification**					**03615520**
					GRA	
Lactobacillus fermentum		91.96	0.46	SAL 1	99	
Bifidobacterium longum subsp. longum		3.23	0.14		99	
Bacteroides vulgatus		2.97	-.01		1	
03615522	**Low discrimination**					**03615522**
					GRA	
Lactobacillus fermentum		77.89	0.46	SAL 1	99	
Parabacteroides distasonis		19.37	0.32		1	
Bifidobacterium longum subsp. longum		2.73	0.14		99	
03615722	**Good identification**					**03615722**
					GRA	
Parabacteroides distasonis		98.13	0.40	MNS 98	1	
Eubacterium contortum		1.87	0.00		99	
03615732	**Good identification**					**03615732**
Parabacteroides distasonis		100.0	0.36	MNS 98		
03616100	**Good identification**					**03616100**
Fusobacterium mortiferum		100.0	0.36	MAN 1		
03617100	**Low discrimination**					**03617100**
					GRA	
Fusobacterium mortiferum		87.36	0.39	MAN 1	1	
Actinomyces israelii		7.20	0.23	XYL 7	99	
Lactobacillus fermentum		5.44	0.20		99	
03620000	**Good identification**					**03620000**
Fusobacterium mortiferum		100.0	0.40	FRU 99		
03620100	**Good identification**					**03620100**
Fusobacterium mortiferum		100.0	0.46	FRU 99		
03621000	**Low discrimination**					**03621000**
					GRA	
Fusobacterium mortiferum		57.39	0.43	FRU 99	1	
Eubacterium tortuosum		41.48	0.24		99	
Actinomyces naeslundii		1.13	0.25		99	
03621100	**Good identification**					**03621100**
					GRA	
Fusobacterium mortiferum		97.65	0.49	FRU 99	1	
Actinomyces naeslundii		2.35	0.33		99	
03621101	**Low discrimination**					**03621101**
					GRA	
Actinomyces naeslundii		78.16	0.36	MNS 88	99	
Fusobacterium mortiferum		21.84	0.15	MLT 94	1	
03621103	**Good identification**					**03621103**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03621121	**Good identification**					**03621121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03621123	**Good identification**					**03621123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.36	MNS 88			
03623100	**Low discrimination**						**03623100**
					GRA		
Actinomyces naeslundii		82.24	0.38	MNS 88	99		
Fusobacterium mortiferum		17.76	0.15	MLT 94	1		
03623101	**Good identification**						**03623101**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03623102	**Good identification**						**03623102**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03623103	**Good identification**						**03623103**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03623120	**Good identification**						**03623120**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03623121	**Good identification**						**03623121**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03623122	**Good identification**						**03623122**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03623123	**Good identification**						**03623123**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03630000	**Very good identification**						**03630000**
					GRA	CAT	
Fusobacterium mortiferum		99.44	0.74		1	5	
Propionibacterium granulosum		0.54	0.32		99	90	
Actinomyces naeslundii		0.02	0.23		99	25	
03630001	**Good identification**						**03630001**
					GRA		
Fusobacterium mortiferum		96.60	0.40	URE 1	1		
Actinomyces naeslundii		3.40	0.26		99		
03630002	**Very good identification**						**03630002**
					GRA	CAT	
Fusobacterium mortiferum		98.59	0.57	BGL 10	1	5	
Propionibacterium granulosum		1.20	0.21		99	90	
Actinomyces naeslundii		0.21	0.23		99	25	
03630010	**Acceptable identification**						**03630010**
					GRA	CAT	
Fusobacterium mortiferum		93.20	0.40	MLZ 1	1	5	
Propionibacterium granulosum		6.80	0.17		99	90	
03630020	**Low discrimination**						**03630020**
					GRA	CAT	
Fusobacterium mortiferum		86.41	0.40	NAG 1	1	5	
Propionibacterium granulosum		11.56	0.21		99	90	
Actinomyces naeslundii		2.03	0.23		99	25	
03630040	**Good identification**						**03630040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.40	SOR 1			
03630100	**Very good identification**						**03630100**
					GRA		
Fusobacterium mortiferum		99.97	0.80		1		
Actinomyces naeslundii		0.03	0.31		99		
03630101	**Good identification**						**03630101**
					GRA		
Fusobacterium mortiferum		95.88	0.46	URE 1	1		
Actinomyces naeslundii		4.12	0.34		99		
03630102	**Very good identification**						**03630102**
					GRA		
Fusobacterium mortiferum		99.74	0.64	BGL 10	1		
Actinomyces naeslundii		0.26	0.31		99		
03630110	**Good identification**						**03630110**
Fusobacterium mortiferum		100.0	0.46	MLZ 1			
03630120	**Good identification**						**03630120**
					GRA		
Fusobacterium mortiferum		97.22	0.46	NAG 1	1		
Actinomyces naeslundii		2.78	0.31		99		
03630140	**Good identification**						**03630140**
Fusobacterium mortiferum		100.0	0.46	SOR 1			
03630200	**Good identification**						**03630200**
Fusobacterium mortiferum		100.0	0.40	RHA 1			
03630300	**Good identification**						**03630300**
Fusobacterium mortiferum		100.0	0.46	RHA 1			
03630400	**Good identification**						**03630400**
Fusobacterium mortiferum		100.0	0.40	ARA 1			
03630500	**Good identification**						**03630500**
Fusobacterium mortiferum		100.0	0.46	ARA 1			
03631000	**Very good identification**						**03631000**
					GRA	CAT	
Fusobacterium mortiferum		99.23	0.77		1	5	
Actinomyces naeslundii		0.38	0.47		99	25	
Propionibacterium granulosum		0.36	0.32		99	90	
03631001	**Low discrimination**						**03631001**
					GRA		
Fusobacterium mortiferum		63.98	0.43	URE 1	1		
Actinomyces naeslundii		36.03	0.50		99		
03631002	**Very good identification**						**03631002**
					GRA	CAT	
Fusobacterium mortiferum		95.63	0.60	BGL 10	1	5	
Actinomyces naeslundii		3.26	0.47		99	25	
Propionibacterium granulosum		0.78	0.21		99	90	
03631003	**Low discrimination**						**03631003**
					GRA		
Actinomyces naeslundii		83.52	0.50	MNS 88	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		16.48	0.26	MLT 94	1		
03631010	**Acceptable identification**						**03631010**
					GRA	CAT	
Fusobacterium mortiferum		93.60	0.43	MLZ 1	1	5	
Propionibacterium granulosum		4.55	0.17		99	90	
Actinomyces naeslundii		1.85	0.25		99	25	
03631020	**Low discrimination**						**03631020**
					GRA	CAT	
Fusobacterium mortiferum		68.28	0.43	NAG 1	1	5	
Actinomyces naeslundii		25.63	0.47		99	25	
Propionibacterium granulosum		6.09	0.21		99	90	
03631021	**Good identification**						**03631021**
Actinomyces naeslundii		100.0	0.50	MNS 88			
03631022	**Low discrimination**						**03631022**
					GRA	CAT	
Actinomyces naeslundii		73.78	0.47	MNS 88	99	25	
Fusobacterium mortiferum		21.84	0.26	MLT 94	1	5	
Propionibacterium granulosum		4.38	0.11		99	90	
03631023	**Good identification**						**03631023**
Actinomyces naeslundii		100.0	0.50	MNS 88			
03631040	**Acceptable identification**						**03631040**
					GRA		
Fusobacterium mortiferum		93.79	0.43	SOR 1	1		
Actinomyces naeslundii		6.21	0.34		99		
03631041	**Good identification**						**03631041**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03631043	**Good identification**						**03631043**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03631061	**Good identification**						**03631061**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03631063	**Good identification**						**03631063**
Actinomyces naeslundii		100.0	0.37	MNS 88			
03631100	**Very good identification**						**03631100**
					GRA		
Fusobacterium mortiferum		99.51	0.83		1		
Actinomyces naeslundii		0.46	0.55		99		
Lactobacillus fermentum		0.03	0.24		99		
03631101	**Low discrimination**						**03631101**
					GRA		
Fusobacterium mortiferum		59.28	0.49	URE 1	1		
Actinomyces naeslundii		40.72	0.58		99		
03631102	**Very good identification**						**03631102**
					GRA		
Fusobacterium mortiferum		95.72	0.67	BGL 10	1		
Actinomyces naeslundii		3.99	0.55		99		
Lactobacillus fermentum		0.23	0.24		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03631103	**Low discrimination**					**03631103**
					GRA	
Actinomyces naeslundii		86.08	0.58	MNS 88	99	
Fusobacterium mortiferum		13.92	0.33	MLT 94	1	
03631110	**Good identification**					**03631110**
					GRA	
Fusobacterium mortiferum		97.65	0.49	MLZ 1	1	
Actinomyces naeslundii		2.35	0.33		99	
03631111	**Low discrimination**					**03631111**
					GRA	
Actinomyces naeslundii		78.16	0.36	MNS 88	99	
Fusobacterium mortiferum		21.84	0.15	MLT 94	1	
03631113	**Good identification**					**03631113**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03631120	**Low discrimination**					**03631120**
					GRA	
Fusobacterium mortiferum		67.27	0.49	NAG 1	1	
Actinomyces naeslundii		30.80	0.55		99	
Clostridium perfringens		1.33	0.09		99	
03631121	**Very good identification**					**03631121**
					GRA	
Actinomyces naeslundii		98.55	0.58	MNS 88	99	
Fusobacterium mortiferum		1.45	0.15	MLT 94	1	
03631122	**Low discrimination**					**03631122**
					GRA	
Actinomyces naeslundii		65.38	0.55	MNS 88	99	
Fusobacterium mortiferum		15.87	0.33	MLT 94	1	
Parabacteroides distasonis		15.18	0.31		1	
03631123	**Very good identification**					**03631123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03631131	**Good identification**					**03631131**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03631133	**Good identification**					**03631133**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03631140	**Acceptable identification**					**03631140**
					GRA	
Fusobacterium mortiferum		92.52	0.49	SOR 1	1	
Actinomyces naeslundii		7.48	0.42		99	
03631141	**Acceptable identification**					**03631141**
					GRA	
Actinomyces naeslundii		92.31	0.45	MNS 88	99	
Fusobacterium mortiferum		7.69	0.15	MLT 94	1	
03631143	**Good identification**					**03631143**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03631160	**Low discrimination**					**03631160**
					GRA	
Actinomyces naeslundii		88.89	0.42	MNS 88	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		11.11	0.15	MLT 94	1		
03631161	**Good identification**						**03631161**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03631162	**Good identification**						**03631162**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03631163	**Good identification**						**03631163**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03631200	**Good identification**						**03631200**
Fusobacterium mortiferum		100.0	0.43	RHA 1			
03631300	**Good identification**						**03631300**
Fusobacterium mortiferum		100.0	0.49	RHA 1			
03631322	**Good identification**						**03631322**
Parabacteroides distasonis		100.0	0.39	MNS 98			
03631400	**Good identification**						**03631400**
					GRA		
Fusobacterium mortiferum		96.32	0.43	ARA 1	1		
Lactobacillus fermentum		3.68	0.20		99		
03631500	**Good identification**						**03631500**
					GRA		
Fusobacterium mortiferum		97.40	0.49	ARA 1	1		
Lactobacillus fermentum		2.60	0.24		99		
03631722	**Good identification**						**03631722**
Parabacteroides distasonis		100.0	0.35	MNS 98			
03632000	**Low discrimination**						**03632000**
					GRA	CAT	
Fusobacterium mortiferum		69.01	0.40	MAN 1	1	5	
Propionibacterium granulosum		27.85	0.30		99	90	
Actinomyces naeslundii		3.14	0.28		99	25	
03632100	**Acceptable identification**						**03632100**
					GRA		
Fusobacterium mortiferum		94.74	0.46	MAN 1	1		
Actinomyces naeslundii		5.26	0.36		99		
03632101	**Low discrimination**						**03632101**
					GRA		
Actinomyces naeslundii		89.19	0.39	MNS 88	99		
Fusobacterium mortiferum		10.81	0.12	MLT 94	1		
03632103	**Good identification**						**03632103**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03632120	**Low discrimination**						**03632120**
					GRA		
Actinomyces naeslundii		84.62	0.36	MNS 88	99		
Fusobacterium mortiferum		15.38	0.12	MLT 94	1		
03632121	**Good identification**						**03632121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	MNS 88			
03632122	**Good identification**						**03632122**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03632123	**Good identification**						**03632123**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03633000	**Low discrimination**						**03633000**
					GRA	CAT	
Fusobacterium mortiferum		50.06	0.43	MAN 1	1	5	
Actinomyces naeslundii		36.48	0.52		99	25	
Propionibacterium granulosum		13.47	0.30		99	90	
03633001	**Good identification**						**03633001**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03633002	**Low discrimination**						**03633002**
					GRA	CAT	
Actinomyces naeslundii		80.34	0.52	MNS 88	99	25	
Fusobacterium mortiferum		12.25	0.26	MLT 94	1	5	
Propionibacterium granulosum		7.42	0.19		99	90	
03633003	**Good identification**						**03633003**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03633020	**Acceptable identification**						**03633020**
Actinomyces naeslundii		91.55	0.52	MNS 88			
Propionibacterium granulosum		8.45	0.19	MLT 94			
03633021	**Good identification**						**03633021**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03633022	**Good identification**						**03633022**
Actinomyces naeslundii		97.75	0.52	MNS 88			
Propionibacterium granulosum		2.26	0.09	MLT 94			
03633023	**Good identification**						**03633023**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03633040	**Good identification**						**03633040**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03633041	**Good identification**						**03633041**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03633042	**Good identification**						**03633042**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03633043	**Good identification**						**03633043**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03633060	**Good identification**						**03633060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	MNS 88		
03633061	**Good identification**					**03633061**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03633062	**Good identification**					**03633062**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03633063	**Good identification**					**03633063**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03633100	**Low discrimination**					**03633100**
					GRA	
Fusobacterium mortiferum		52.75	0.49	MAN 1	1	
Actinomyces naeslundii		46.89	0.60		99	
Actinomyces israelii		0.36	0.14		99	
03633101	**Very good identification**					**03633101**
					GRA	
Actinomyces naeslundii		99.25	0.63	MNS 88	99	
Fusobacterium mortiferum		0.75	0.15	MLT 94	1	
03633102	**Low discrimination**					**03633102**
					GRA	
Actinomyces naeslundii		87.36	0.60	MNS 88	99	
Fusobacterium mortiferum		10.92	0.33	MLT 94	1	
Actinomyces israelii		1.00	0.17		99	
03633103	**Very good identification**					**03633103**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03633110	**Low discrimination**					**03633110**
					GRA	
Actinomyces naeslundii		82.24	0.38	MNS 88	99	
Fusobacterium mortiferum		17.76	0.15	MLT 94	1	
03633111	**Good identification**					**03633111**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03633112	**Good identification**					**03633112**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03633113	**Good identification**					**03633113**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03633120	**Very good identification**					**03633120**
					GRA	
Actinomyces naeslundii		98.88	0.60	MNS 88	99	
Fusobacterium mortiferum		1.12	0.15	MLT 94	1	
03633121	**Very good identification**					**03633121**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03633122	**Very good identification**					**03633122**
Actinomyces naeslundii		100.0	0.60	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03633123	**Very good identification**					**03633123**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03633130	**Good identification**					**03633130**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03633131	**Good identification**					**03633131**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03633132	**Good identification**					**03633132**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03633133	**Good identification**					**03633133**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03633140	**Acceptable identification**					**03633140**
					GRA	
Actinomyces naeslundii		93.95	0.47	MNS 88	99	
Fusobacterium mortiferum		6.05	0.15	MLT 94	1	
03633141	**Good identification**					**03633141**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03633142	**Good identification**					**03633142**
Actinomyces naeslundii		95.56	0.47	MNS 88		
Propionibacterium propionicum		4.44	0.14	MLT 94		
03633143	**Good identification**					**03633143**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03633160	**Good identification**					**03633160**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03633161	**Good identification**					**03633161**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03633162	**Good identification**					**03633162**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03633163	**Good identification**					**03633163**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03634000	**Good identification**					**03634000**
Fusobacterium mortiferum		100.0	0.55	XYL 7		
03634002	**Good identification**					**03634002**
Fusobacterium mortiferum		100.0	0.38	XYL 7		
03634100	**Very good identification**					**03634100**
Fusobacterium mortiferum		100.0	0.61	XYL 7		
03634102	**Good identification**					**03634102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.45	XYL 7		
03635000	**Very good identification**					**03635000**
					GRA	
Fusobacterium mortiferum		99.70	0.58	XYL 7	1	
Lactobacillus fermentum		0.30	0.16		99	
03635002	**Good identification**					**03635002**
					GRA	
Fusobacterium mortiferum		97.36	0.41	XYL 7	1	
Lactobacillus fermentum		2.64	0.16	BGL 10	99	
03635100	**Very good identification**					**03635100**
					GRA	
Fusobacterium mortiferum		99.61	0.64	XYL 7	1	
Lactobacillus fermentum		0.21	0.20		99	
Actinomyces israelii		0.18	0.19		99	
03635102	**Acceptable identification**					**03635102**
					GRA	
Fusobacterium mortiferum		92.24	0.48	XYL 7	1	
Bifidobacterium adolescentis		2.63	0.17	BGL 10	99	
Actinomyces israelii		2.30	0.23		99	
03635122	**Good identification**					**03635122**
Parabacteroides distasonis		99.13	0.47	MNS 98		
Fusobacterium mortiferum		0.87	0.13			
03635132	**Good identification**					**03635132**
Parabacteroides distasonis		100.0	0.43	MNS 98		
03635322	**Good identification**					**03635322**
Parabacteroides distasonis		100.0	0.55	MNS 98		
03635332	**Good identification**					**03635332**
Parabacteroides distasonis		100.0	0.51	MNS 98		
03635522	**Good identification**					**03635522**
Parabacteroides distasonis		100.0	0.44	MNS 98		
03635532	**Good identification**					**03635532**
					GRA	
Parabacteroides distasonis		97.61	0.39	MNS 98	1	
Bifidobacterium longum subsp. longum		2.39	0.07		99	
03635722	**Good identification**					**03635722**
Parabacteroides distasonis		100.0	0.52	MNS 98		
03635732	**Good identification**					**03635732**
Parabacteroides distasonis		97.66	0.47	MNS 98		
Clostridium clostridioforme		2.34	0.14			
03637102	**Low discrimination**					**03637102**
Actinomyces israelii		76.16	0.35	MLT 99		
Bifidobacterium adolescentis		12.56	0.16	CEL 93		
Actinomyces naeslundii		5.84	0.26			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03640000	**Acceptable identification**					**03640000**
					GRA	
Fusobacterium mortiferum		90.26	0.40	FRU 99	1	
Eubacterium tortuosum		9.74	0.07		99	
03640100	**Good identification**					**03640100**
Fusobacterium mortiferum		100.0	0.47	FRU 99		
03641000	**Low discrimination**					**03641000**
					GRA	
Eubacterium tortuosum		87.69	0.41	ESL 99	99	
Fusobacterium mortiferum		12.31	0.43	CEL 10	1	
03641100	**Good identification**					**03641100**
					GRA	
Fusobacterium mortiferum		95.64	0.50	FRU 99	1	
Eubacterium tortuosum		2.95	0.07		99	
Actinomyces naeslundii		1.41	0.30		99	
03643101	**Good identification**					**03643101**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03643103	**Good identification**					**03643103**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03643121	**Good identification**					**03643121**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03643123	**Good identification**					**03643123**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03650000	**Very good identification**					**03650000**
Fusobacterium mortiferum		100.0	0.75			
03650001	**Good identification**					**03650001**
					GRA	
Fusobacterium mortiferum		97.89	0.40	URE 1	1	
Actinomyces naeslundii		2.11	0.24		99	
03650002	**Very good identification**					**03650002**
Fusobacterium mortiferum		100.0	0.58	BGL 10		
03650010	**Good identification**					**03650010**
Fusobacterium mortiferum		100.0	0.40	MLZ 1		
03650020	**Good identification**					**03650020**
Fusobacterium mortiferum		100.0	0.40	NAG 1		
03650040	**Good identification**					**03650040**
Fusobacterium mortiferum		100.0	0.40	SOR 1		
03650100	**Very good identification**					**03650100**
					GRA	
Fusobacterium mortiferum		99.98	0.81		1	
Actinomyces naeslundii		0.02	0.28		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03650101	**Good identification**					**03650101**
					GRA	
Fusobacterium mortiferum		97.44	0.47	URE 1	1	
Actinomyces naeslundii		2.56	0.31		99	
03650102	**Very good identification**					**03650102**
					GRA	
Fusobacterium mortiferum		99.84	0.65	BGL 10	1	
Actinomyces naeslundii		0.16	0.28		99	
03650110	**Good identification**					**03650110**
Fusobacterium mortiferum		100.0	0.47	MLZ 1		
03650120	**Good identification**					**03650120**
					GRA	
Fusobacterium mortiferum		98.28	0.47	NAG 1	1	
Actinomyces naeslundii		1.72	0.28		99	
03650140	**Good identification**					**03650140**
Fusobacterium mortiferum		100.0	0.47	SOR 1		
03650200	**Good identification**					**03650200**
Fusobacterium mortiferum		100.0	0.40	RHA 1		
03650300	**Good identification**					**03650300**
Fusobacterium mortiferum		100.0	0.47	RHA 1		
03650400	**Good identification**					**03650400**
Fusobacterium mortiferum		100.0	0.40	ARA 1		
03650500	**Good identification**					**03650500**
Fusobacterium mortiferum		100.0	0.47	ARA 1		
03651000	**Very good identification**					**03651000**
					GRA	
Fusobacterium mortiferum		99.59	0.78		1	
Actinomyces naeslundii		0.23	0.44		99	
Eubacterium tortuosum		0.07	0.07		99	
03651001	**Low discrimination**					**03651001**
					GRA	
Fusobacterium mortiferum		74.37	0.43	URE 1	1	
Actinomyces naeslundii		25.63	0.47		99	
03651002	**Very good identification**					**03651002**
					GRA	
Fusobacterium mortiferum		96.74	0.61	BGL 10	1	
Actinomyces naeslundii		2.02	0.44		99	
Lactobacillus catenaformis		0.51	0.12		99	
03651003	**Low discrimination**					**03651003**
					GRA	
Actinomyces naeslundii		75.62	0.47	MNS 88	99	
Fusobacterium mortiferum		24.38	0.27	MLT 94	1	
03651010	**Good identification**					**03651010**
Fusobacterium mortiferum		100.0	0.43	MLZ 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03651020	**Low discrimination**					**03651020**
					GRA	
Fusobacterium mortiferum		79.56	0.43	NAG 1	1	
Actinomyces naeslundii		18.28	0.44		99	
Lactobacillus catenaformis		1.15	0.02		99	
03651021	**Good identification**					**03651021**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03651022	**Low discrimination**					**03651022**
					GRA	
Actinomyces naeslundii		54.28	0.44	MNS 88	99	
Fusobacterium mortiferum		26.26	0.27	MLT 94	1	
Lactobacillus acidophilus		8.58	0.06		99	
03651023	**Good identification**					**03651023**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03651040	**Good identification**					**03651040**
					GRA	
Fusobacterium mortiferum		96.10	0.43	SOR 1	1	
Actinomyces naeslundii		3.90	0.31		99	
03651100	**Very good identification**					**03651100**
					GRA	
Fusobacterium mortiferum		99.66	0.84		1	
Actinomyces naeslundii		0.28	0.52		99	
Lactobacillus catenaformis		0.04	0.15		99	
03651101	**Low discrimination**					**03651101**
					GRA	
Fusobacterium mortiferum		70.41	0.50	URE 1	1	
Actinomyces naeslundii		29.59	0.55		99	
03651102	**Very good identification**					**03651102**
					GRA	
Fusobacterium mortiferum		95.25	0.68	BGL 10	1	
Actinomyces naeslundii		2.43	0.52		99	
Lactobacillus acidophilus		1.55	0.24		99	
03651103	**Low discrimination**					**03651103**
					GRA	
Actinomyces naeslundii		79.09	0.55	MNS 88	99	
Fusobacterium mortiferum		20.91	0.33	MLT 94	1	
03651110	**Good identification**					**03651110**
					GRA	
Fusobacterium mortiferum		98.55	0.50	MLZ 1	1	
Actinomyces naeslundii		1.45	0.30		99	
03651120	**Low discrimination**					**03651120**
					GRA	
Fusobacterium mortiferum		75.67	0.50	NAG 1	1	
Actinomyces naeslundii		21.20	0.52		99	
Prevotella melaninogenica		0.88	-.01		1	
03651121	**Good identification**					**03651121**
					GRA	
Actinomyces naeslundii		97.65	0.55	MNS 88	99	
Fusobacterium mortiferum		2.35	0.16	MLT 94	1	
03651122	**Low discrimination**					**03651122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		38.74	0.52	MNS 88	99		
Lactobacillus acidophilus		24.73	0.24	MLT 94	99		
Fusobacterium mortiferum		15.36	0.33		1		
03651123	**Good identification**						**03651123**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03651140	**Good identification**						**03651140**
					GRA		
Fusobacterium mortiferum		95.29	0.50	SOR 1	1		
Actinomyces naeslundii		4.71	0.39		99		
03651141	**Low discrimination**						**03651141**
					GRA		
Actinomyces naeslundii		88.02	0.42	MNS 88	99		
Fusobacterium mortiferum		11.98	0.16	MLT 94	1		
03651143	**Good identification**						**03651143**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03651160	**Low discrimination**						**03651160**
					GRA		
Actinomyces naeslundii		83.04	0.39	MNS 88	99		
Fusobacterium mortiferum		16.96	0.16	MLT 94	1		
03651161	**Good identification**						**03651161**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03651162	**Low discrimination**						**03651162**
Actinomyces naeslundii		69.11	0.39	MNS 88			
Lactobacillus acidophilus		30.89	0.09	MLT 94			
03651163	**Good identification**						**03651163**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03651200	**Good identification**						**03651200**
Fusobacterium mortiferum		100.0	0.43	RHA 1			
03651300	**Good identification**						**03651300**
Fusobacterium mortiferum		100.0	0.50	RHA 1			
03651400	**Good identification**						**03651400**
					GRA		
Fusobacterium mortiferum		96.69	0.43	ARA 1	1		
Lactobacillus fermentum		3.31	0.20		99		
03651500	**Good identification**						**03651500**
					GRA		
Fusobacterium mortiferum		97.66	0.50	ARA 1	1		
Lactobacillus fermentum		2.34	0.24		99		
03652000	**Low discrimination**						**03652000**
					GRA	SPO	
Fusobacterium mortiferum		89.64	0.40	MAN 1	1	1	
Clostridium innocuum		7.86	0.10		99	99	
Actinomyces naeslundii		2.50	0.25		99	1	
03652100	**Good identification**						**03652100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		96.71	0.47	MAN 1	1	
Actinomyces naeslundii		3.29	0.33		99	
03652101	**Low discrimination**					**03652101**
					GRA	
Actinomyces naeslundii		83.47	0.36	MNS 88	99	
Fusobacterium mortiferum		16.53	0.13	MLT 94	1	
03652103	**Good identification**					**03652103**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03652121	**Good identification**					**03652121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03652123	**Good identification**					**03652123**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03653001	**Good identification**					**03653001**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03653003	**Good identification**					**03653003**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03653020	**Good identification**					**03653020**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03653021	**Good identification**					**03653021**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03653022	**Good identification**					**03653022**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03653023	**Good identification**					**03653023**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03653040	**Good identification**					**03653040**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03653041	**Good identification**					**03653041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03653042	**Good identification**					**03653042**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03653043	**Good identification**					**03653043**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03653060	**Good identification**					**03653060**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03653061	**Good identification**					**03653061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	MNS 88		
03653062	**Good identification**					**03653062**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03653063	**Good identification**					**03653063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03653100	**Low discrimination**					**03653100**
					GRA	
Fusobacterium mortiferum		63.88	0.50	MAN 1	1	
Actinomyces naeslundii		34.75	0.57		99	
Actinomyces israelii		1.37	0.24		99	
03653101	**Very good identification**					**03653101**
					GRA	
Actinomyces naeslundii		98.78	0.60	MNS 88	99	
Fusobacterium mortiferum		1.22	0.16	MLT 94	1	
03653102	**Low discrimination**					**03653102**
					GRA	
Actinomyces naeslundii		77.16	0.57	MNS 88	99	
Fusobacterium mortiferum		15.76	0.33	MLT 94	1	
Actinomyces israelii		4.57	0.27		99	
03653103	**Very good identification**					**03653103**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03653111	**Good identification**					**03653111**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03653113	**Good identification**					**03653113**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03653120	**Very good identification**					**03653120**
					GRA	
Actinomyces naeslundii		98.18	0.57	MNS 88	99	
Fusobacterium mortiferum		1.82	0.16	MLT 94	1	
03653121	**Very good identification**					**03653121**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03653122	**Very good identification**					**03653122**
					SPO	
Actinomyces naeslundii		95.80	0.57	MNS 88	1	
Clostridium ramosum		2.54	0.05	MLT 94	92	
Lactobacillus acidophilus		1.66	0.02		1	
03653123	**Very good identification**					**03653123**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03653131	**Good identification**					**03653131**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03653133	**Good identification**					**03653133**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03653140	**Low discrimination**					**03653140**
					GRA	
Actinomyces naeslundii		83.87	0.44	MNS 88	99	
Fusobacterium mortiferum		8.82	0.16	MLT 94	1	
Actinomyces israelii		7.31	0.17		99	
03653141	**Good identification**					**03653141**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03653142	**Low discrimination**					**03653142**
Actinomyces naeslundii		88.45	0.44	MNS 88		
Actinomyces israelii		11.55	0.20	MLT 94		
03653143	**Good identification**					**03653143**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03653160	**Good identification**					**03653160**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03653161	**Good identification**					**03653161**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03653162	**Good identification**					**03653162**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03653163	**Good identification**					**03653163**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03654000	**Good identification**					**03654000**
Fusobacterium mortiferum		100.0	0.55	XYL 7		
03654002	**Good identification**					**03654002**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
03654100	**Very good identification**					**03654100**
Fusobacterium mortiferum		100.0	0.62	XYL 7		
03654102	**Good identification**					**03654102**
Fusobacterium mortiferum		100.0	0.45	XYL 7		
03655000	**Very good identification**					**03655000**
					GRA	
Fusobacterium mortiferum		99.39	0.58	XYL 7	1	
Actinomyces israelii		0.34	0.19		99	
Lactobacillus fermentum		0.27	0.16		99	
03655002	**Acceptable identification**					**03655002**
					GRA	
Fusobacterium mortiferum		93.43	0.42	XYL 7	1	
Actinomyces israelii		4.31	0.22	BGL 10	99	
Lactobacillus fermentum		2.27	0.16		99	
03655100	**Very good identification**					**03655100**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		99.23	0.65	XYL 7	1	
Actinomyces israelii		0.58	0.29		99	
Lactobacillus fermentum		0.19	0.20		99	
03655102	**Low discrimination**					**03655102**
					GRA	
Fusobacterium mortiferum		84.93	0.48	XYL 7	1	
Bifidobacterium adolescentis		6.92	0.26	BGL 10	99	
Actinomyces israelii		6.71	0.32		99	
03655322	**Good identification**					**03655322**
Parabacteroides distasonis		98.22	0.40	MNS 98		
Prevotella oralis		1.78	-.03			
03655502	**Low discrimination**					**03655502**
Bifidobacterium adolescentis		85.60	0.40	MNS 87		
Actinomyces israelii		10.15	0.30	MLT 99		
Lactobacillus fermentum		2.67	0.20			
03655512	**Good identification**					**03655512**
Bifidobacterium adolescentis		98.57	0.41	MNS 87		
Actinomyces israelii		1.43	0.16	MLT 99		
03655722	**Good identification**					**03655722**
Parabacteroides distasonis		100.0	0.36	MNS 98		
03657100	**Low discrimination**					**03657100**
					GRA	
Actinomyces israelii		75.37	0.42	MLT 99	99	
Fusobacterium mortiferum		22.95	0.31		1	
Actinomyces naeslundii		1.68	0.23		99	
03657102	**Low discrimination**					**03657102**
					GRA	
Actinomyces israelii		84.30	0.45	MLT 99	99	
Bifidobacterium adolescentis		12.55	0.24		99	
Fusobacterium mortiferum		1.90	0.14		1	
03657142	**Good identification**					**03657142**
Actinomyces israelii		95.48	0.38	MLT 99		
Bifidobacterium adolescentis		4.52	0.09			
03657500	**Good identification**					**03657500**
Actinomyces israelii		100.0	0.40	MLT 99		
03657502	**Low discrimination**					**03657502**
Bifidobacterium adolescentis		54.91	0.38	MNS 87		
Actinomyces israelii		45.09	0.43	MLT 99		
03657512	**Acceptable identification**					**03657512**
Bifidobacterium adolescentis		90.88	0.40	MNS 87		
Actinomyces israelii		9.13	0.29	MLT 99		
03657542	**Low discrimination**					**03657542**
Actinomyces israelii		72.10	0.36	MLT 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Bifidobacterium adolescentis	27.90	0.23			
03660100	**Good identification**					**03660100**
	Fusobacterium mortiferum	100.0	0.38	FRU 99		
03661100	**Acceptable identification**					**03661100**
					GRA	
	Fusobacterium mortiferum	91.25	0.41	FRU 99	1	
	Actinomyces naeslundii	8.75	0.35		99	
03661101	**Good identification**					**03661101**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03661103	**Good identification**					**03661103**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03661121	**Good identification**					**03661121**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03661123	**Good identification**					**03661123**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03663001	**Good identification**					**03663001**
	Actinomyces naeslundii	100.0	0.35	MNS 88		
03663003	**Good identification**					**03663003**
	Actinomyces naeslundii	100.0	0.35	MNS 88		
03663021	**Good identification**					**03663021**
	Actinomyces naeslundii	100.0	0.35	MNS 88		
03663023	**Good identification**					**03663023**
	Actinomyces naeslundii	100.0	0.35	MNS 88		
03663100	**Good identification**					**03663100**
	Actinomyces naeslundii	100.0	0.40	MNS 88		
03663101	**Good identification**					**03663101**
	Actinomyces naeslundii	100.0	0.43	MNS 88		
03663102	**Good identification**					**03663102**
	Actinomyces naeslundii	100.0	0.40	MNS 88		
03663103	**Good identification**					**03663103**
	Actinomyces naeslundii	100.0	0.43	MNS 88		
03663120	**Good identification**					**03663120**
	Actinomyces naeslundii	100.0	0.40	MNS 88		
03663121	**Good identification**					**03663121**
	Actinomyces naeslundii	100.0	0.43	MNS 88		
03663122	**Good identification**					**03663122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MNS 88		
03663123	**Good identification**					**03663123**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03670000	**Very good identification**					**03670000**
					GRA	
Fusobacterium mortiferum		99.91	0.66		1	
Actinomyces naeslundii		0.09	0.25		99	
03670002	**Good identification**					**03670002**
					GRA	
Fusobacterium mortiferum		99.16	0.49	BGL 10	1	
Actinomyces naeslundii		0.84	0.25		99	
03670100	**Very good identification**					**03670100**
					GRA	
Fusobacterium mortiferum		99.89	0.72		1	
Actinomyces naeslundii		0.12	0.33		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03670101	**Low discrimination**					**03670101**
					GRA	
Fusobacterium mortiferum		85.42	0.38	URE 1	1	
Actinomyces naeslundii		14.58	0.36		99	
03670102	**Good identification**					**03670102**
					GRA	
Fusobacterium mortiferum		98.71	0.56	BGL 10	1	
Actinomyces naeslundii		1.02	0.33		99	
Lactobacillus acidophilus		0.27	-.01		99	
03670103	**Low discrimination**					**03670103**
					GRA	
Actinomyces naeslundii		60.58	0.36	MNS 88	99	
Fusobacterium mortiferum		39.42	0.21	MLT 94	1	
03670110	**Good identification**					**03670110**
Fusobacterium mortiferum		100.0	0.38	MLZ 1		
03670120	**Low discrimination**					**03670120**
					GRA	
Fusobacterium mortiferum		89.78	0.38	NAG 1	1	
Actinomyces naeslundii		10.22	0.33		99	
03670121	**Good identification**					**03670121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03670123	**Good identification**					**03670123**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03670140	**Good identification**					**03670140**
Fusobacterium mortiferum		100.0	0.38	SOR 1		
03670300	**Good identification**					**03670300**
Fusobacterium mortiferum		100.0	0.38	RHA 1		
03670500	**Good identification**					**03670500**
Fusobacterium mortiferum		100.0	0.38	ARA 1		
03671000	**Very good identification**					**03671000**
					GRA	
Fusobacterium mortiferum		98.38	0.69		1	
Actinomyces naeslundii		1.48	0.49		99	
Lactobacillus jensenii		0.14	0.10		99	
03671001	**Low discrimination**					**03671001**
					GRA	
Actinomyces naeslundii		69.13	0.52	MNS 88	99	
Fusobacterium mortiferum		30.87	0.34	MLT 94	1	
03671002	**Low discrimination**					**03671002**
					GRA	
Fusobacterium mortiferum		84.40	0.52	BGL 10	1	
Actinomyces naeslundii		11.45	0.49		99	
Lactobacillus acidophilus		3.12	0.15		99	
03671003	**Good identification**					**03671003**
					GRA	
Actinomyces naeslundii		95.27	0.52	MNS 88	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		4.73	0.18	MLT 94	1	
03671020	**Low discrimination**					**03671020**
					GRA	
Actinomyces naeslundii		59.89	0.49	MNS 88	99	
Fusobacterium mortiferum		40.11	0.34	MLT 94	1	
03671021	**Good identification**					**03671021**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03671022	**Low discrimination**					**03671022**
					GRA	
Actinomyces naeslundii		74.25	0.49	MNS 88	99	
Lactobacillus acidophilus		20.23	0.15	MLT 94	99	
Fusobacterium mortiferum		5.53	0.18		1	
03671023	**Good identification**					**03671023**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03671041	**Good identification**					**03671041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03671042	**Low discrimination**					**03671042**
					GRA	
Actinomyces naeslundii		62.01	0.36	MNS 88	99	
Fusobacterium mortiferum		26.15	0.18	MLT 94	1	
Lactobacillus acidophilus		11.83	-.01		99	
03671043	**Good identification**					**03671043**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03671060	**Good identification**					**03671060**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03671061	**Good identification**					**03671061**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03671062	**Low discrimination**					**03671062**
Actinomyces naeslundii		83.98	0.36	MNS 88		
Lactobacillus acidophilus		16.02	-.01	MLT 94		
03671063	**Good identification**					**03671063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03671100	**Very good identification**					**03671100**
					GRA	
Fusobacterium mortiferum		98.13	0.75		1	
Actinomyces naeslundii		1.81	0.57		99	
Actinomyces israelii		0.05	0.21		99	
03671101	**Low discrimination**					**03671101**
					GRA	
Actinomyces naeslundii		73.20	0.60	MNS 88	99	
Fusobacterium mortiferum		26.80	0.41	MLT 94	1	
03671102	**Low discrimination**					**03671102**
					GRA	
Fusobacterium mortiferum		73.73	0.59	BGL 10	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		13.44	0.33		99		
Actinomyces naeslundii		12.21	0.57		99		
03671103	**Very good identification**						**03671103**
					GRA		
Actinomyces naeslundii		95.41	0.60	MNS 88	99		
Fusobacterium mortiferum		3.88	0.24	MLT 94	1		
Lactobacillus acidophilus		0.71	-.01		99		
03671110	**Acceptable identification**						**03671110**
					GRA		
Fusobacterium mortiferum		91.25	0.41	MLZ 1	1		
Actinomyces naeslundii		8.75	0.35		99		
03671111	**Good identification**						**03671111**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03671113	**Good identification**						**03671113**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03671120	**Low discrimination**						**03671120**
					SPO	GRA	
Actinomyces naeslundii		63.35	0.57	MNS 88	1	99	
Fusobacterium mortiferum		34.79	0.41	MLT 94	1	1	
Clostridium ramosum		1.16	0.02		92	50	
03671121	**Very good identification**						**03671121**
Actinomyces naeslundii		100.0	0.60	MNS 88			
03671123	**Very good identification**						**03671123**
Actinomyces naeslundii		99.26	0.60	MNS 88			
Lactobacillus acidophilus		0.74	-.01	MLT 94			
03671131	**Good identification**						**03671131**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03671133	**Good identification**						**03671133**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03671140	**Low discrimination**						**03671140**
					GRA		
Fusobacterium mortiferum		74.63	0.41	SOR 1	1		
Actinomyces naeslundii		23.98	0.44		99		
Actinomyces israelii		1.39	0.14		99		
03671141	**Good identification**						**03671141**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03671142	**Low discrimination**						**03671142**
					GRA		
Actinomyces naeslundii		45.38	0.44	MNS 88	99		
Lactobacillus acidophilus		34.99	0.18	MLT 94	99		
Fusobacterium mortiferum		15.69	0.24		1		
03671143	**Good identification**						**03671143**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03671160	**Good identification**						**03671160**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	MNS 88			
03671161	**Good identification**						**03671161**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03671162	**Low discrimination**						**03671162**
Actinomyces naeslundii		56.47	0.44	MNS 88			
Lactobacillus acidophilus		43.54	0.18	MLT 94			
03671163	**Good identification**						**03671163**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03671300	**Good identification**						**03671300**
					GRA		
Fusobacterium mortiferum		98.19	0.41	RHA 1	1		
Actinomyces naeslundii		1.81	0.23		99		
03671500	**Acceptable identification**						**03671500**
					GRA		
Fusobacterium mortiferum		94.57	0.41	ARA 1	1		
Actinomyces israelii		3.70	0.19		99		
Actinomyces naeslundii		1.74	0.23		99		
03672100	**Low discrimination**						**03672100**
					GRA		
Fusobacterium mortiferum		81.90	0.38	MAN 1	1		
Actinomyces naeslundii		18.10	0.38		99		
03672101	**Good identification**						**03672101**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03672102	**Low discrimination**						**03672102**
					GRA		
Actinomyces naeslundii		66.54	0.38	MNS 88	99		
Fusobacterium mortiferum		33.46	0.21	MLT 94	1		
03672103	**Good identification**						**03672103**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03672120	**Good identification**						**03672120**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03672121	**Good identification**						**03672121**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03672122	**Good identification**						**03672122**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03672123	**Good identification**						**03672123**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03673000	**Low discrimination**						**03673000**
					GRA	SPO	
Actinomyces naeslundii		54.05	0.54	MNS 88	99	1	
Clostridium innocuum		24.36	0.24	MLT 94	99	99	
Fusobacterium mortiferum		18.65	0.34		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03673001	**Very good identification**					**03673001**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03673002	**Low discrimination**					**03673002**
					SPO	
Actinomyces naeslundii		63.67	0.54	MNS 88	1	
Clostridium innocuum		28.69	0.24	MLT 94	99	
Actinomyces israelii		5.20	0.26		1	
03673003	**Very good identification**					**03673003**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03673011	**Good identification**					**03673011**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03673013	**Good identification**					**03673013**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03673020	**Good identification**					**03673020**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03673021	**Very good identification**					**03673021**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03673022	**Good identification**					**03673022**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03673023	**Very good identification**					**03673023**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03673031	**Good identification**					**03673031**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03673033	**Good identification**					**03673033**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03673040	**Low discrimination**					**03673040**
Actinomyces naeslundii		89.28	0.41	MNS 88		
Actinomyces israelii		10.72	0.16	MLT 94		
03673041	**Good identification**					**03673041**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03673042	**Low discrimination**					**03673042**
Actinomyces naeslundii		84.74	0.41	MNS 88		
Actinomyces israelii		15.27	0.19	MLT 94		
03673043	**Good identification**					**03673043**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03673060	**Good identification**					**03673060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MNS 88		
03673061	**Good identification**					**03673061**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03673062	**Good identification**					**03673062**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03673063	**Good identification**					**03673063**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03673100	**Low discrimination**					**03673100**
					GRA	
Actinomyces naeslundii		73.56	0.62	MNS 88	99	
Fusobacterium mortiferum		20.81	0.41	MLT 94	1	
Actinomyces israelii		5.63	0.33		99	
03673101	**Very good identification**					**03673101**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03673102	**Low discrimination**					**03673102**
					GRA	
Actinomyces naeslundii		84.65	0.62	MNS 88	99	
Actinomyces israelii		9.72	0.36	MLT 94	99	
Fusobacterium mortiferum		2.66	0.24		1	
03673103	**Very good identification**					**03673103**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03673110	**Low discrimination**					**03673110**
Actinomyces naeslundii		82.14	0.40	MNS 88		
Actinomyces israelii		17.86	0.19	MLT 94		
03673111	**Good identification**					**03673111**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03673112	**Low discrimination**					**03673112**
Actinomyces naeslundii		69.11	0.40	MNS 88		
Actinomyces israelii		22.54	0.22	MLT 94		
Bifidobacterium adolescentis		8.36	0.09	MLZ 5		
03673113	**Good identification**					**03673113**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03673120	**Good identification**					**03673120**
					SPO	
Actinomyces naeslundii		94.92	0.62	MNS 88	1	
Clostridium ramosum		5.08	0.15	MLT 94	92	
03673121	**Very good identification**					**03673121**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03673122	**Low discrimination**					**03673122**
					SPO	
Actinomyces naeslundii		79.91	0.62	MNS 88	1	
Clostridium ramosum		17.12	0.25	MLT 94	92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		2.39	0.11		1	
03673123	**Very good identification**					**03673123**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03673130	**Good identification**					**03673130**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03673131	**Good identification**					**03673131**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03673132	**Good identification**					**03673132**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03673133	**Good identification**					**03673133**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03673140	**Low discrimination**					**03673140**
Actinomyces naeslundii		85.56	0.49	MNS 88		
Actinomyces israelii		14.44	0.26	MLT 94		
03673141	**Good identification**					**03673141**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03673142	**Low discrimination**					**03673142**
Actinomyces naeslundii		79.80	0.49	MNS 88		
Actinomyces israelii		20.20	0.29	MLT 94		
03673143	**Good identification**					**03673143**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03673160	**Good identification**					**03673160**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03673161	**Good identification**					**03673161**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03673162	**Good identification**					**03673162**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03673163	**Good identification**					**03673163**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03674000	**Good identification**					**03674000**
Fusobacterium mortiferum		100.0	0.46	XYL 7		
03674100	**Good identification**					**03674100**
Fusobacterium mortiferum		100.0	0.53	XYL 7		
03674102	**Good identification**					**03674102**
Fusobacterium mortiferum		100.0	0.36	XYL 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03675000	**Good identification**						**03675000**
					GRA		
Fusobacterium mortiferum		95.88	0.49	XYL 7	1		
Actinomyces israelii		4.12	0.28		99		
03675100	**Acceptable identification**						**03675100**
					GRA	SPO	
Fusobacterium mortiferum		92.29	0.56	XYL 7	1	1	
Actinomyces israelii		6.80	0.39		99	1	
Clostridium butyricum		0.68	0.01		99	99	
03675102	**Low discrimination**						**03675102**
					GRA		
Fusobacterium mortiferum		44.45	0.39	XYL 7	1		
Actinomyces israelii		44.24	0.42	BGL 10	99		
Bifidobacterium adolescentis		8.08	0.23		99		
03675122	**Acceptable identification**						**03675122**
					GRA		
Parabacteroides distasonis		94.43	0.43	MNS 98	1		
Actinomyces israelii		3.21	0.21		99		
Lactobacillus acidophilus		1.24	-.01		99		
03675132	**Good identification**						**03675132**
Parabacteroides distasonis		100.0	0.38	MNS 98			
03675322	**Good identification**						**03675322**
Parabacteroides distasonis		100.0	0.51	MNS 98			
03675332	**Good identification**						**03675332**
Parabacteroides distasonis		100.0	0.46	MNS 98			
03675502	**Low discrimination**						**03675502**
					SPO		
Bifidobacterium adolescentis		53.75	0.37	MNS 87	1		
Actinomyces israelii		35.97	0.40	MLT 99	1		
Clostridium butyricum		10.29	0.10		99		
03675512	**Acceptable identification**						**03675512**
Bifidobacterium adolescentis		90.33	0.38	MNS 87			
Actinomyces israelii		7.39	0.26	MLT 99			
Clostridium clostridioforme		2.28	0.08				
03675522	**Good identification**						**03675522**
					GRA		
Parabacteroides distasonis		95.66	0.39	MNS 98	1		
Actinomyces israelii		4.35	0.20		99		
03675722	**Good identification**						**03675722**
Parabacteroides distasonis		98.27	0.47	MNS 98			
Clostridium clostridioforme		1.73	0.11				
03675732	**Low discrimination**						**03675732**
Parabacteroides distasonis		87.05	0.43	MNS 98			
Clostridium clostridioforme		12.95	0.23				
03677000	**Acceptable identification**						**03677000**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		92.65	0.41	MLT 99	99	1	
Fusobacterium mortiferum		3.84	0.15		1	1	
Clostridium innocuum		3.51	0.02		99	99	
03677002	**Good identification**						**03677002**
					SPO		
Actinomyces israelii		97.54	0.44	MLT 99	1		
Clostridium innocuum		2.46	0.02		99		
03677042	**Good identification**						**03677042**
Actinomyces israelii		100.0	0.37	MLT 99			
03677100	**Good identification**						**03677100**
					GRA		
Actinomyces israelii		96.56	0.52	MLT 99	99		
Fusobacterium mortiferum		2.34	0.22		1		
Actinomyces naeslundii		1.11	0.28		99		
03677102	**Good identification**						**03677102**
Actinomyces israelii		96.71	0.55	MLT 99			
Bifidobacterium adolescentis		2.55	0.21				
Actinomyces naeslundii		0.74	0.28				
03677110	**Good identification**						**03677110**
Actinomyces israelii		100.0	0.37	MLT 99			
03677112	**Low discrimination**						**03677112**
Actinomyces israelii		82.25	0.41	MLT 99			
Bifidobacterium adolescentis		17.75	0.23	MLZ 13			
03677140	**Good identification**						**03677140**
Actinomyces israelii		100.0	0.45	MLT 99			
03677142	**Good identification**						**03677142**
Actinomyces israelii		100.0	0.48	MLT 99			
03677302	**Good identification**						**03677302**
Actinomyces israelii		100.0	0.35	MLT 99			
03677400	**Good identification**						**03677400**
Actinomyces israelii		100.0	0.40	MLT 99			
03677402	**Good identification**						**03677402**
Actinomyces israelii		100.0	0.43	MLT 99			
03677442	**Good identification**						**03677442**
Actinomyces israelii		100.0	0.36	MLT 99			
03677500	**Good identification**						**03677500**
					SPO		
Actinomyces israelii		97.88	0.50	MLT 99	1		
Clostridium butyricum		2.12	0.01		99		
03677502	**Low discrimination**						**03677502**
					SPO		
Actinomyces israelii		81.87	0.53	MLT 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		17.67	0.35		1		
Clostridium butyricum		0.46	-.07		99		
03677510	**Good identification**						**03677510**
Actinomyces israelii		100.0	0.36	MLT 99			
03677512	**Low discrimination**						**03677512**
Bifidobacterium adolescentis		62.56	0.37	MNS 87			
Actinomyces israelii		35.45	0.39	MLT 99			
Bifidobacterium dentium		1.99	-.07				
03677540	**Good identification**						**03677540**
Actinomyces israelii		100.0	0.43	MLT 99			
03677542	**Acceptable identification**						**03677542**
Actinomyces israelii		93.58	0.46	MLT 99			
Bifidobacterium adolescentis		6.42	0.20				
03700100	**Low discrimination**						**03700100**
					COC	GRA	
Fusobacterium mortiferum		89.13	0.39	MLT 11	1	1	
Blautia hansenii		3.31	-.03	FRU 99	99	99	
Actinomyces odontolyticus		2.94	0.08		1	99	
03701001	**Good identification**						**03701001**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03701002	**Low discrimination**						**03701002**
Actinomyces naeslundii		31.90	0.41	MNS 88			
Eubacterium tortuosum		25.52	0.11	FRU 95			
Lactobacillus fermentum		22.93	0.28				
03701003	**Good identification**						**03701003**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03701020	**Low discrimination**						**03701020**
Actinomyces naeslundii		44.86	0.41	MNS 88			
Eubacterium tortuosum		35.88	0.11	FRU 95			
Lactobacillus fermentum		10.75	0.20				
03701021	**Good identification**						**03701021**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03701022	**Low discrimination**						**03701022**
Actinomyces naeslundii		72.43	0.41	MNS 88			
Lactobacillus fermentum		17.35	0.20	FRU 95			
Eubacterium tortuosum		10.22	-.02				
03701023	**Good identification**						**03701023**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03701101	**Acceptable identification**						**03701101**
					COC	SPO	
Actinomyces naeslundii		93.59	0.51	MNS 88	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		3.29	-.03	FRU 95	99	1	
Clostridium chauvoei		3.12	0.03		1	99	
03701102	**Low discrimination**						**03701102**
					COC		
Actinomyces naeslundii		42.46	0.48	MNS 88	1		
Blautia hansenii		24.59	0.15	FRU 95	99		
Lactobacillus fermentum		17.50	0.32		1		
03701103	**Good identification**						**03701103**
Actinomyces naeslundii		97.89	0.51	MNS 88			
Actinomyces viscosus		2.11	-.01	FRU 95			
03701121	**Good identification**						**03701121**
					GRA		
Actinomyces naeslundii		95.74	0.51	MNS 88	99		
Prevotella melaninogenica		4.26	-.01	FRU 95	1		
03701123	**Good identification**						**03701123**
Actinomyces naeslundii		100.0	0.51	MNS 88			
03701140	**Low discrimination**						**03701140**
					COC		
Actinomyces naeslundii		77.02	0.35	MNS 88	1		
Blautia hansenii		22.98	-.03	FRU 95	99		
03701141	**Good identification**						**03701141**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03701142	**Good identification**						**03701142**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03701143	**Good identification**						**03701143**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03701160	**Low discrimination**						**03701160**
					GRA		
Actinomyces naeslundii		72.55	0.35	MNS 88	99		
Prevotella melaninogenica		27.45	-.01	FRU 95	1		
03701161	**Good identification**						**03701161**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03701162	**Good identification**						**03701162**
Actinomyces naeslundii		100.0	0.35	MNS 88			
03701163	**Good identification**						**03701163**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03703000	**Good identification**						**03703000**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03703001	**Good identification**						**03703001**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03703002	**Good identification**						**03703002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MNS 88		
03703003	**Good identification**					**03703003**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03703020	**Good identification**					**03703020**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03703021	**Good identification**					**03703021**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03703022	**Good identification**					**03703022**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03703023	**Good identification**					**03703023**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03703041	**Good identification**					**03703041**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03703043	**Good identification**					**03703043**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03703061	**Good identification**					**03703061**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03703063	**Good identification**					**03703063**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03703100	**Good identification**					**03703100**
					COC	
Actinomyces naeslundii		95.89	0.53	MNS 88	1	
Blautia hansenii		2.60	-.03	FRU 95	99	
Propionibacterium propionicum		1.51	0.12		1	
03703101	**Very good identification**					**03703101**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03703102	**Good identification**					**03703102**
Actinomyces naeslundii		97.70	0.53	MNS 88		
Propionibacterium propionicum		2.30	0.15	FRU 95		
03703103	**Very good identification**					**03703103**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03703120	**Good identification**					**03703120**
					GRA	
Actinomyces naeslundii		96.68	0.53	MNS 88	99	
Prevotella melaninogenica		3.33	-.01	FRU 95	1	
03703121	**Very good identification**					**03703121**
Actinomyces naeslundii		100.0	0.56	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03703122	**Good identification**					**03703122**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03703123	**Very good identification**					**03703123**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03703140	**Acceptable identification**					**03703140**
Actinomyces naeslundii		91.82	0.40	MNS 88		
Propionibacterium propionicum		8.18	0.12	FRU 95		
03703141	**Good identification**					**03703141**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03703142	**Low discrimination**					**03703142**
Actinomyces naeslundii		88.21	0.40	MNS 88		
Propionibacterium propionicum		11.79	0.15	FRU 95		
03703143	**Good identification**					**03703143**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03703160	**Good identification**					**03703160**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03703161	**Good identification**					**03703161**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03703162	**Good identification**					**03703162**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03703163	**Good identification**					**03703163**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03705512	**Low discrimination**					**03705512**
Bifidobacterium adolescentis		77.30	0.36	MNS 87		
Bifidobacterium longum subsp. longum		22.70	0.25	FRU 99		
03705530	**Good identification**					**03705530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87		
03705532	**Good identification**					**03705532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87		
03710000	**Low discrimination**					**03710000**
					GRA	
Fusobacterium mortiferum		87.69	0.67	MLT 11	1	
Prevotella intermedia		6.24	0.36		1	
Actinomyces odontolyticus		2.48	0.35		99	
03710002	**Low discrimination**					**03710002**
					GRA	
Fusobacterium mortiferum		65.67	0.51	MLT 11	1	
Actinomyces viscosus		16.63	0.28	BGL 10	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		10.25	0.31		99	
03710021	**Good identification**					**03710021**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03710022	**Low discrimination**					**03710022**
Actinomyces naeslundii		46.38	0.39	MNS 88		
Actinomyces odontolyticus		28.32	0.18	GAL 96		
Lactobacillus fermentum		14.04	0.20			
03710023	**Good identification**					**03710023**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03710100	**Very good identification**					**03710100**
					GRA	
Fusobacterium mortiferum		98.28	0.74	MLT 11	1	
Actinomyces naeslundii		0.55	0.47		99	
Actinomyces meyeri		0.42	0.24		99	
03710101	**Low discrimination**					**03710101**
					GRA	
Fusobacterium mortiferum		54.43	0.39	MLT 11	1	
Actinomyces naeslundii		45.57	0.50	URE 1	99	
03710102	**Low discrimination**					**03710102**
					GRA	
Fusobacterium mortiferum		77.15	0.57	MLT 11	1	
Actinomyces viscosus		14.89	0.32	BGL 10	99	
Actinomyces naeslundii		3.92	0.47		99	
03710103	**Low discrimination**					**03710103**
					GRA	
Actinomyces naeslundii		50.56	0.50	MNS 88	99	
Actinomyces viscosus		42.73	0.24	GAL 96	99	
Fusobacterium mortiferum		6.71	0.23		1	
03710110	**Good identification**					**03710110**
					GRA	
Fusobacterium mortiferum		97.15	0.39	MLT 11	1	
Actinomyces naeslundii		2.85	0.25	MLZ 1	99	
03710120	**Low discrimination**					**03710120**
					GRA	
Fusobacterium mortiferum		44.69	0.39	MLT 11	1	
Actinomyces naeslundii		24.95	0.47	NAG 1	99	
Prevotella melaninogenica		18.94	0.15		1	
03710121	**Good identification**					**03710121**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03710122	**Low discrimination**					**03710122**
					GRA	
Actinomyces naeslundii		52.52	0.47	MNS 88	99	
Prevotella oralis		11.15	0.09	GAL 96	1	
Fusobacterium mortiferum		10.45	0.23		1	
03710123	**Good identification**					**03710123**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03710140	**Low discrimination**					**03710140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		89.28	0.39	MLT 11	1	
Actinomyces naeslundii		8.80	0.34	SOR 1	99	
Propionibacterium propionicum		1.92	0.12		99	
03710141	**Good identification**					**03710141**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03710143	**Good identification**					**03710143**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03710161	**Good identification**					**03710161**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03710163	**Good identification**					**03710163**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03710200	**Low discrimination**					**03710200**
					GRA	
Actinomyces odontolyticus		83.54	0.40	NIT 99	99	
Fusobacterium mortiferum		15.37	0.33	RAF 3	1	
Prevotella intermedia		1.09	0.02		1	
03710202	**Good identification**					**03710202**
					GRA	
Actinomyces odontolyticus		96.77	0.36	NIT 99	99	
Fusobacterium mortiferum		3.23	0.17	RAF 3	1	
03710300	**Low discrimination**					**03710300**
					GRA	
Fusobacterium mortiferum		57.61	0.39	MLT 11	1	
Actinomyces odontolyticus		42.39	0.31	RHA 1	99	
03710500	**Low discrimination**					**03710500**
					GRA	
Fusobacterium mortiferum		71.57	0.39	MLT 11	1	
Lactobacillus fermentum		20.79	0.32	ARA 1	99	
Actinomyces meyeri		7.64	0.13		99	
03711000	**Low discrimination**					**03711000**
					GRA	
Fusobacterium mortiferum		70.64	0.70	MLT 11	1	
Lactobacillus fermentum		19.57	0.62		99	
Actinomyces naeslundii		5.23	0.62		99	
03711001	**Low discrimination**					**03711001**
					GRA	
Actinomyces naeslundii		89.24	0.66	MNS 88	99	
Fusobacterium mortiferum		8.13	0.36		1	
Lactobacillus fermentum		2.25	0.28		99	
03711002	**Low discrimination**					**03711002**
					GRA	
Lactobacillus fermentum		57.30	0.62	SAL 1	99	
Fusobacterium mortiferum		22.99	0.54		1	
Actinomyces naeslundii		15.30	0.62		99	
03711003	**Good identification**					**03711003**
Actinomyces naeslundii		93.47	0.66	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		3.23	0.16			
Lactobacillus fermentum		2.36	0.28			
03711010	**Low discrimination**					**03711010**
					GRA	
Fusobacterium mortiferum		58.49	0.36	MLT 11	1	
Actinomyces naeslundii		22.54	0.41	MLZ 1	99	
Lactobacillus fermentum		16.20	0.28		99	
03711011	**Good identification**					**03711011**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03711012	**Low discrimination**					**03711012**
					GRA	
Actinomyces naeslundii		47.58	0.41	MNS 88	99	
Lactobacillus fermentum		34.20	0.28	MLZ 5	99	
Fusobacterium mortiferum		13.72	0.20		1	
03711013	**Good identification**					**03711013**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03711020	**Low discrimination**					**03711020**
					GRA	
Lactobacillus fermentum		47.41	0.54	SAL 1	99	
Actinomyces naeslundii		37.98	0.62		99	
Fusobacterium mortiferum		5.19	0.36		1	
03711021	**Very good identification**					**03711021**
Actinomyces naeslundii		99.17	0.66	MNS 88		
Lactobacillus fermentum		0.83	0.20			
03711022	**Low discrimination**					**03711022**
					GRA	
Lactobacillus fermentum		49.60	0.54	SAL 1	99	
Actinomyces naeslundii		39.73	0.62		99	
Prevotella oralis		6.32	0.23		1	
03711023	**Very good identification**					**03711023**
Actinomyces naeslundii		99.17	0.66	MNS 88		
Lactobacillus fermentum		0.83	0.20			
03711030	**Low discrimination**					**03711030**
Actinomyces naeslundii		80.67	0.41	MNS 88		
Lactobacillus fermentum		19.33	0.20	MLZ 5		
03711031	**Good identification**					**03711031**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03711032	**Low discrimination**					**03711032**
Actinomyces naeslundii		80.67	0.41	MNS 88		
Lactobacillus fermentum		19.33	0.20	MLZ 5		
03711033	**Good identification**					**03711033**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03711040	**Low discrimination**					**03711040**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		49.38	0.50	MNS 88	99	
Fusobacterium mortiferum		38.22	0.36		1	
Lactobacillus fermentum		10.59	0.28		99	
03711041	**Good identification**					**03711041**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03711042	**Low discrimination**					**03711042**
					GRA	
Actinomyces naeslundii		76.90	0.50	MNS 88	99	
Lactobacillus fermentum		16.48	0.28		99	
Fusobacterium mortiferum		6.61	0.20		1	
03711043	**Good identification**					**03711043**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03711060	**Low discrimination**					**03711060**
Actinomyces naeslundii		83.95	0.50	MNS 88		
Clostridium perfringens		10.05	0.12			
Lactobacillus fermentum		6.00	0.20			
03711061	**Good identification**					**03711061**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03711062	**Low discrimination**					**03711062**
Actinomyces naeslundii		88.67	0.50	MNS 88		
Lactobacillus fermentum		6.34	0.20			
Clostridium perfringens		5.00	0.06			
03711063	**Good identification**					**03711063**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03711100	**Low discrimination**					**03711100**
					GRA	
Fusobacterium mortiferum		77.13	0.77	MLT 11	1	
Lactobacillus fermentum		14.94	0.66		99	
Actinomyces naeslundii		6.96	0.70		99	
03711101	**Good identification**					**03711101**
					GRA	
Actinomyces naeslundii		91.48	0.73	MNS 88	99	
Fusobacterium mortiferum		6.83	0.42		1	
Lactobacillus fermentum		1.32	0.32		99	
03711102	**Low discrimination**					**03711102**
					GRA	
Lactobacillus fermentum		44.10	0.66	SAL 1	99	
Fusobacterium mortiferum		25.30	0.60		1	
Actinomyces naeslundii		20.54	0.70		99	
03711103	**Very good identification**					**03711103**
Actinomyces naeslundii		95.66	0.73	MNS 88		
Actinomyces viscosus		2.06	0.21			
Lactobacillus fermentum		1.38	0.32			
03711110	**Low discrimination**					**03711110**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		59.55	0.42	MLT 11	1	
Actinomyces naeslundii		27.99	0.48	MLZ 1	99	
Lactobacillus fermentum		11.53	0.32		99	
03711111	**Good identification**					**03711111**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03711112	**Low discrimination**					**03711112**
Bifidobacterium adolescentis		65.75	0.42	MNS 87		
Actinomyces naeslundii		19.97	0.48	XYL 87		
Lactobacillus fermentum		8.23	0.32	ARA 87		
03711113	**Good identification**					**03711113**
Actinomyces naeslundii		97.83	0.51	MNS 88		
Bifidobacterium adolescentis		2.17	0.08	MLZ 5		
03711120	**Low discrimination**					**03711120**
					GRA	
Prevotella melaninogenica		58.75	0.50	MNS 90	1	
Actinomyces naeslundii		18.76	0.70	SAL 1	99	
Lactobacillus fermentum		13.43	0.58		99	
03711121	**Very good identification**					**03711121**
					GRA	
Actinomyces naeslundii		97.16	0.73	MNS 88	99	
Prevotella melaninogenica		2.05	0.15		1	
Lactobacillus fermentum		0.47	0.24		99	
03711122	**Low discrimination**					**03711122**
					GRA	
Actinomyces naeslundii		27.29	0.70	MNS 88	99	
Prevotella oralis		23.89	0.44		1	
Lactobacillus fermentum		19.53	0.58		99	
03711123	**Very good identification**					**03711123**
					GRA	
Actinomyces naeslundii		98.36	0.73	MNS 88	99	
Prevotella oralis		0.58	0.09		1	
Lactobacillus fermentum		0.47	0.24		99	
03711130	**Low discrimination**					**03711130**
					GRA	
Actinomyces naeslundii		54.17	0.48	MNS 88	99	
Prevotella melaninogenica		32.55	0.15	MLZ 5	1	
Lactobacillus fermentum		7.44	0.24		99	
03711131	**Good identification**					**03711131**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03711132	**Low discrimination**					**03711132**
					GRA	
Actinomyces naeslundii		55.89	0.48	MNS 88	99	
Parabacteroides distasonis		12.80	0.24	MLZ 5	1	
Prevotella oralis		9.39	0.09		1	
03711133	**Good identification**					**03711133**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03711140	**Low discrimination**					**03711140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		55.72	0.57	MNS 88	99	
Fusobacterium mortiferum		35.35	0.42		1	
Lactobacillus fermentum		6.85	0.32		99	
03711141	**Very good identification**					**03711141**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03711142	**Low discrimination**					**03711142**
Actinomyces naeslundii		74.42	0.57	MNS 88		
Lactobacillus fermentum		9.15	0.32			
Bifidobacterium adolescentis		7.39	0.25			
03711143	**Very good identification**					**03711143**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03711150	**Good identification**					**03711150**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03711151	**Good identification**					**03711151**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03711153	**Good identification**					**03711153**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03711160	**Low discrimination**					**03711160**
					GRA	
Actinomyces naeslundii		65.41	0.57	MNS 88	99	
Clostridium perfringens		20.19	0.26		99	
Prevotella melaninogenica		11.72	0.15		1	
03711161	**Very good identification**					**03711161**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03711162	**Low discrimination**					**03711162**
					GRA	
Actinomyces naeslundii		77.73	0.57	MNS 88	99	
Clostridium perfringens		11.29	0.21		99	
Prevotella oralis		3.90	0.09		1	
03711163	**Very good identification**					**03711163**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03711170	**Good identification**					**03711170**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03711171	**Good identification**					**03711171**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03711172	**Good identification**					**03711172**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03711173	**Good identification**					**03711173**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03711200	**Low discrimination**					**03711200**
					GRA	
Actinomyces odontolyticus		63.34	0.37	NIT 99	99	
Fusobacterium mortiferum		26.21	0.36	RAF 3	1	
Lactobacillus fermentum		7.26	0.28		99	
03711300	**Low discrimination**					**03711300**
					GRA	
Fusobacterium mortiferum		43.30	0.42	MLT 11	1	
Eubacterium contortum		30.25	0.27	RHA 1	99	
Actinomyces odontolyticus		14.16	0.28		99	
03711301	**Good identification**					**03711301**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03711303	**Good identification**					**03711303**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03711321	**Good identification**					**03711321**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03711322	**Low discrimination**					**03711322**
					GRA	
Prevotella oralis		79.04	0.37	MNS 99	1	
Parabacteroides distasonis		14.16	0.37		1	
Actinomyces naeslundii		2.13	0.36		99	
03711323	**Low discrimination**					**03711323**
					GRA	
Actinomyces naeslundii		79.99	0.39	MNS 88	99	
Prevotella oralis		20.01	0.03	RHA 1	1	
03711400	**Very good identification**					**03711400**
					GRA	
Lactobacillus fermentum		95.74	0.62	SAL 1	99	
Fusobacterium mortiferum		3.49	0.36		1	
Actinomyces naeslundii		0.26	0.28		99	
03711402	**Very good identification**					**03711402**
					GRA	
Lactobacillus fermentum		98.42	0.62	SAL 1	99	
Bifidobacterium adolescentis		0.72	0.20		99	
Fusobacterium mortiferum		0.40	0.20		1	
03711420	**Good identification**					**03711420**
Lactobacillus fermentum		98.83	0.54	SAL 1		
Actinomyces naeslundii		0.80	0.28			
Bifidobacterium longum subsp. longum		0.37	0.05			
03711422	**Good identification**					**03711422**
Lactobacillus fermentum		98.83	0.54	SAL 1		
Actinomyces naeslundii		0.80	0.28			
Bifidobacterium longum subsp. longum		0.37	0.05			
03711500	**Good identification**					**03711500**
					GRA	
Lactobacillus fermentum		91.82	0.66	SAL 1	99	
Fusobacterium mortiferum		4.79	0.42		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		1.74	0.29		99	
03711502	**Low discrimination**					**03711502**
Lactobacillus fermentum		67.70	0.66	SAL 1		
Bifidobacterium adolescentis		29.93	0.54			
Bifidobacterium longum subsp. longum		1.28	0.29			
03711512	**Acceptable identification**					**03711512**
Bifidobacterium adolescentis		92.74	0.56	MNS 87		
Bifidobacterium longum subsp. longum		5.52	0.33	XYL 87		
Lactobacillus fermentum		1.73	0.32			
03711520	**Low discrimination**					**03711520**
					GRA	
Lactobacillus fermentum		76.68	0.58	SAL 1	99	
Bifidobacterium longum subsp. longum		17.39	0.40		99	
Prevotella melaninogenica		3.39	0.15		1	
03711521	**Low discrimination**					**03711521**
Actinomyces naeslundii		63.08	0.39	MNS 88		
Lactobacillus fermentum		30.09	0.24	ARA 1		
Bifidobacterium longum subsp. longum		6.83	0.05			
03711522	**Low discrimination**					**03711522**
					GRA	
Lactobacillus fermentum		76.84	0.58	SAL 1	99	
Bifidobacterium longum subsp. longum		17.43	0.40		99	
Parabacteroides distasonis		1.47	0.25		1	
03711523	**Low discrimination**					**03711523**
Actinomyces naeslundii		63.08	0.39	MNS 88		
Lactobacillus fermentum		30.09	0.24	ARA 1		
Bifidobacterium longum subsp. longum		6.83	0.05			
03711530	**Good identification**					**03711530**
Bifidobacterium longum subsp. longum		97.45	0.44	MNS 87		
Lactobacillus fermentum		2.55	0.24	SAL 1		
03711532	**Acceptable identification**					**03711532**
					GRA	
Bifidobacterium longum subsp. longum		91.30	0.44	MNS 87	99	
Bifidobacterium adolescentis		3.87	0.22	SAL 1	99	
Parabacteroides distasonis		2.44	0.21		1	
03711542	**Low discrimination**					**03711542**
Bifidobacterium adolescentis		80.99	0.39	MNS 87		
Lactobacillus fermentum		14.97	0.32	XYL 87		
Actinomyces israelii		2.81	0.20	SOR 11		
03711552	**Good identification**					**03711552**
Bifidobacterium adolescentis		100.0	0.40	MNS 87		
03712001	**Good identification**					**03712001**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03712002	**Low discrimination**					**03712002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Actinomyces naeslundii		64.44	0.44	MNS 88	1	99	
Megasphaera elsdenii		22.03	0.15	GAL 96	99	1	
Fusobacterium mortiferum		8.06	0.17		1	1	
03712003	**Good identification**						**03712003**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03712020	**Good identification**						**03712020**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03712021	**Good identification**						**03712021**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03712022	**Good identification**						**03712022**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03712023	**Good identification**						**03712023**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03712100	**Low discrimination**						**03712100**
					GRA		
Propionibacterium propionicum		50.56	0.46	NIT 86	99		
Actinomyces naeslundii		25.71	0.52	SAL 1	99		
Fusobacterium mortiferum		23.73	0.39		1		
03712101	**Low discrimination**						**03712101**
Actinomyces naeslundii		87.29	0.55	MNS 88			
Propionibacterium propionicum		12.72	0.30	GAL 96			
03712102	**Low discrimination**						**03712102**
					GRA		
Propionibacterium propionicum		72.44	0.49	NIT 86	99		
Actinomyces naeslundii		24.56	0.52	SAL 1	99		
Fusobacterium mortiferum		2.52	0.23		1		
03712103	**Low discrimination**						**03712103**
Actinomyces naeslundii		82.07	0.55	MNS 88			
Propionibacterium propionicum		17.93	0.33	GAL 96			
03712120	**Good identification**						**03712120**
Actinomyces naeslundii		98.05	0.52	MNS 88			
Propionibacterium propionicum		1.95	0.12	GAL 96			
03712121	**Good identification**						**03712121**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03712122	**Good identification**						**03712122**
Actinomyces naeslundii		97.11	0.52	MNS 88			
Propionibacterium propionicum		2.89	0.15	GAL 96			
03712123	**Good identification**						**03712123**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03712140	**Acceptable identification**						**03712140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		91.77	0.46	NIT 86		
Actinomyces naeslundii		8.23	0.39	SAL 1		
03712141	**Low discrimination**					**03712141**
Actinomyces naeslundii		54.78	0.42	MNS 88		
Propionibacterium propionicum		45.22	0.30	GAL 96		
03712142	**Acceptable identification**					**03712142**
Propionibacterium propionicum		94.36	0.49	NIT 86		
Actinomyces naeslundii		5.64	0.39	SAL 1		
03712160	**Low discrimination**					**03712160**
Actinomyces naeslundii		89.89	0.39	MNS 88		
Propionibacterium propionicum		10.12	0.12	GAL 96		
03712161	**Good identification**					**03712161**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03712162	**Low discrimination**					**03712162**
Actinomyces naeslundii		85.56	0.39	MNS 88		
Propionibacterium propionicum		14.45	0.15	GAL 96		
03712163	**Good identification**					**03712163**
Actinomyces naeslundii		100.0	0.42	MNS 88		
03713000	**Low discrimination**					**03713000**
					GRA	
Actinomyces naeslundii		89.28	0.67	MNS 88	99	
Fusobacterium mortiferum		6.28	0.36		1	
Lactobacillus fermentum		1.74	0.28		99	
03713001	**Very good identification**					**03713001**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03713002	**Good identification**					**03713002**
Actinomyces naeslundii		94.57	0.67	MNS 88		
Actinomyces israelii		2.52	0.31			
Lactobacillus fermentum		1.84	0.28			
03713003	**Very good identification**					**03713003**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03713010	**Good identification**					**03713010**
Actinomyces naeslundii		95.21	0.46	MNS 88		
Actinomyces israelii		4.79	0.14	MLZ 5		
03713011	**Good identification**					**03713011**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03713012	**Acceptable identification**					**03713012**
Actinomyces naeslundii		92.98	0.46	MNS 88		
Actinomyces israelii		7.02	0.17	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03713013	**Good identification**					**03713013**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03713020	**Very good identification**					**03713020**
Actinomyces naeslundii		99.36	0.67	MNS 88		
Lactobacillus fermentum		0.65	0.20			
03713021	**Very good identification**					**03713021**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03713022	**Very good identification**					**03713022**
Actinomyces naeslundii		99.36	0.67	MNS 88		
Lactobacillus fermentum		0.65	0.20			
03713023	**Very good identification**					**03713023**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03713030	**Good identification**					**03713030**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03713031	**Good identification**					**03713031**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03713032	**Good identification**					**03713032**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03713033	**Good identification**					**03713033**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03713040	**Good identification**					**03713040**
Actinomyces naeslundii		95.02	0.55	MNS 88		
Actinomyces israelii		3.71	0.21			
Propionibacterium propionicum		1.27	0.12			
03713041	**Very good identification**					**03713041**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03713042	**Acceptable identification**					**03713042**
Actinomyces naeslundii		92.71	0.55	MNS 88		
Actinomyces israelii		5.43	0.24			
Propionibacterium propionicum		1.86	0.15			
03713043	**Very good identification**					**03713043**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03713051	**Good identification**					**03713051**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03713053	**Good identification**					**03713053**
Actinomyces naeslundii		100.0	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03713060	**Good identification**					**03713060**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03713061	**Very good identification**					**03713061**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03713062	**Good identification**					**03713062**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03713063	**Very good identification**					**03713063**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03713071	**Good identification**					**03713071**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03713073	**Good identification**					**03713073**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03713100	**Low discrimination**					**03713100**
					GRA	
Actinomyces naeslundii		85.06	0.75	MNS 88	99	
Propionibacterium propionicum		6.97	0.46		99	
Fusobacterium mortiferum		4.91	0.42		1	
03713101	**Very good identification**					**03713101**
Actinomyces naeslundii		99.40	0.78	MNS 88		
Propionibacterium propionicum		0.60	0.30			
03713102	**Low discrimination**					**03713102**
Actinomyces naeslundii		80.75	0.75	MNS 88		
Propionibacterium propionicum		9.93	0.49			
Bifidobacterium adolescentis		4.83	0.39			
03713103	**Very good identification**					**03713103**
Actinomyces naeslundii		99.10	0.78	MNS 88		
Propionibacterium propionicum		0.90	0.33			
03713110	**Acceptable identification**					**03713110**
Actinomyces naeslundii		92.04	0.53	MNS 88		
Actinomyces israelii		6.51	0.24	MLZ 5		
Propionibacterium propionicum		1.45	0.12			
03713111	**Very good identification**					**03713111**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03713112	**Low discrimination**					**03713112**
Bifidobacterium adolescentis		55.13	0.40	MNS 87		
Actinomyces naeslundii		39.72	0.53	XYL 87		
Actinomyces israelii		4.21	0.27	ARA 87		
03713113	**Very good identification**					**03713113**
Actinomyces naeslundii		100.0	0.56	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03713120	**Very good identification**					**03713120**
					GRA	
Actinomyces naeslundii		97.66	0.75	MNS 88	99	
Prevotella melaninogenica		1.59	0.15		1	
Lactobacillus fermentum		0.36	0.24		99	
03713121	**Very good identification**					**03713121**
Actinomyces naeslundii		100.0	0.78	MNS 88		
03713122	**Very good identification**					**03713122**
					GRA	
Actinomyces naeslundii		98.16	0.75	MNS 88	99	
Prevotella oralis		0.45	0.09		1	
Lactobacillus fermentum		0.37	0.24		99	
03713123	**Very good identification**					**03713123**
Actinomyces naeslundii		100.0	0.78	MNS 88		
03713130	**Good identification**					**03713130**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03713131	**Very good identification**					**03713131**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03713132	**Good identification**					**03713132**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03713133	**Very good identification**					**03713133**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03713140	**Low discrimination**					**03713140**
Actinomyces naeslundii		65.82	0.62	MNS 88		
Propionibacterium propionicum		30.56	0.46			
Actinomyces israelii		3.61	0.32			
03713141	**Very good identification**					**03713141**
Actinomyces naeslundii		96.68	0.65	MNS 88		
Propionibacterium propionicum		3.33	0.30			
03713142	**Low discrimination**					**03713142**
Actinomyces naeslundii		54.92	0.62	MNS 88		
Propionibacterium propionicum		38.26	0.49			
Actinomyces israelii		4.52	0.35			
03713143	**Very good identification**					**03713143**
Actinomyces naeslundii		95.09	0.65	MNS 88		
Propionibacterium propionicum		4.91	0.33			
03713150	**Low discrimination**					**03713150**
Actinomyces naeslundii		80.32	0.40	MNS 88		
Actinomyces israelii		12.52	0.17	MLZ 5		
Propionibacterium propionicum		7.16	0.12			
03713151	**Good identification**					**03713151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	MNS 88		
03713152	**Low discrimination**					**03713152**
Actinomyces naeslundii		42.74	0.40	MNS 88		
Bifidobacterium adolescentis		41.55	0.25	MLZ 5		
Actinomyces israelii		9.99	0.20			
03713153	**Good identification**					**03713153**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03713160	**Very good identification**					**03713160**
Actinomyces naeslundii		99.53	0.62	MNS 88		
Propionibacterium propionicum		0.47	0.12			
03713161	**Very good identification**					**03713161**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03713162	**Very good identification**					**03713162**
Actinomyces naeslundii		98.79	0.62	MNS 88		
Propionibacterium propionicum		0.70	0.15			
Actinomyces israelii		0.52	0.14			
03713163	**Very good identification**					**03713163**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03713170	**Good identification**					**03713170**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03713171	**Good identification**					**03713171**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03713172	**Good identification**					**03713172**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03713173	**Good identification**					**03713173**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03713201	**Good identification**					**03713201**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03713203	**Good identification**					**03713203**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03713221	**Good identification**					**03713221**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03713223	**Good identification**					**03713223**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03713300	**Low discrimination**					**03713300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		78.89	0.41	MNS 88			
Actinomyces israelii		14.65	0.19	RHA 1			
Propionibacterium propionicum		6.46	0.12				
03713301	**Good identification**						**03713301**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03713302	**Low discrimination**						**03713302**
Actinomyces naeslundii		71.36	0.41	MNS 88			
Actinomyces israelii		19.87	0.22	RHA 1			
Propionibacterium propionicum		8.77	0.15				
03713303	**Good identification**						**03713303**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03713320	**Good identification**						**03713320**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03713321	**Good identification**						**03713321**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03713322	**Low discrimination**						**03713322**
					SPO	GRA	
Actinomyces naeslundii		67.20	0.41	MNS 88	1	99	
Clostridium ramosum		19.81	0.07	RHA 1	92	50	
Prevotella oralis		12.99	0.03		1	1	
03713323	**Good identification**						**03713323**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03713401	**Good identification**						**03713401**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03713403	**Good identification**						**03713403**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03713421	**Good identification**						**03713421**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03713423	**Good identification**						**03713423**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03713500	**Low discrimination**						**03713500**
Actinomyces israelii		43.82	0.37	CEL 93			
Lactobacillus fermentum		24.01	0.32	XYL 92			
Actinomyces naeslundii		21.71	0.41				
03713501	**Good identification**						**03713501**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03713502	**Low discrimination**						**03713502**
Bifidobacterium adolescentis		88.28	0.53	MNS 87			
Actinomyces israelii		6.75	0.40	XYL 87			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		2.47	0.32			
03713503	**Low discrimination**					**03713503**
Actinomyces naeslundii		78.95	0.44	MNS 88		
Bifidobacterium adolescentis		21.05	0.19	ARA 1		
03713512	**Good identification**					**03713512**
Bifidobacterium adolescentis		99.07	0.54	MNS 87		
Actinomyces israelii		0.93	0.26	XYL 87		
03713520	**Low discrimination**					**03713520**
Actinomyces naeslundii		63.25	0.41	MNS 88		
Lactobacillus fermentum		23.32	0.24	ARA 1		
Actinomyces israelii		8.15	0.17			
03713521	**Good identification**					**03713521**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03713522	**Low discrimination**					**03713522**
Actinomyces naeslundii		48.89	0.41	MNS 88		
Bifidobacterium adolescentis		19.55	0.19	ARA 1		
Lactobacillus fermentum		18.02	0.24			
03713523	**Good identification**					**03713523**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03713542	**Low discrimination**					**03713542**
Bifidobacterium adolescentis		76.82	0.37	MNS 87		
Actinomyces israelii		18.48	0.33	XYL 87		
Actinomyces naeslundii		2.77	0.28	SOR 11		
03713552	**Good identification**					**03713552**
Bifidobacterium adolescentis		97.14	0.39	MNS 87		
Actinomyces israelii		2.86	0.19	XYL 87		
03714000	**Low discrimination**					**03714000**
					GRA	
Fusobacterium mortiferum		77.38	0.48	MLT 11	1	
Actinomyces odontolyticus		14.33	0.30	XYL 7	99	
Actinomyces meyeri		5.02	0.18		99	
03714100	**Low discrimination**					**03714100**
					GRA	
Fusobacterium mortiferum		86.26	0.54	MLT 11	1	
Actinomyces meyeri		9.60	0.29	XYL 7	99	
Actinomyces odontolyticus		2.16	0.21		99	
03714102	**Low discrimination**					**03714102**
					GRA	
Fusobacterium mortiferum		64.26	0.38	MLT 11	1	
Lactobacillus fermentum		13.24	0.28	XYL 7	99	
Bifidobacterium adolescentis		9.97	0.20	BGL 10	99	
03714512	**Acceptable identification**					**03714512**
Bifidobacterium adolescentis		93.05	0.36	MNS 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		4.47	0.10	GAL 99		
Bifidobacterium longum subsp. longum		2.48	0.07			
03715000	**Low discrimination**					**03715000**
					GRA	
Lactobacillus fermentum		63.69	0.58	SAL 1	99	
Fusobacterium mortiferum		29.47	0.51		1	
Actinomyces odontolyticus		2.43	0.27		99	
03715002	**Low discrimination**					**03715002**
					GRA	
Lactobacillus fermentum		87.77	0.58	SAL 1	99	
Fusobacterium mortiferum		4.51	0.35		1	
Actinomyces israelii		4.18	0.37		99	
03715020	**Acceptable identification**					**03715020**
					GRA	
Lactobacillus fermentum		94.46	0.50	SAL 1	99	
Actinomyces odontolyticus		1.76	0.13		99	
Fusobacterium mortiferum		1.33	0.17		1	
03715022	**Acceptable identification**					**03715022**
					GRA	
Lactobacillus fermentum		92.76	0.50	SAL 1	99	
Prevotella oralis		2.49	0.08		1	
Actinomyces naeslundii		1.28	0.28		99	
03715100	**Low discrimination**					**03715100**
					GRA	
Lactobacillus fermentum		52.40	0.62	SAL 1	99	
Fusobacterium mortiferum		34.67	0.57		1	
Actinomyces israelii		4.08	0.44		99	
03715101	**Low discrimination**					**03715101**
					GRA	
Actinomyces naeslundii		41.72	0.39	MNS 88	99	
Lactobacillus fermentum		35.07	0.28	XYL 1	99	
Fusobacterium mortiferum		23.21	0.23		1	
03715102	**Low discrimination**					**03715102**
Lactobacillus fermentum		48.12	0.62	SAL 1		
Bifidobacterium adolescentis		36.22	0.54			
Actinomyces israelii		5.62	0.47			
03715103	**Low discrimination**					**03715103**
Actinomyces naeslundii		40.43	0.39	MNS 88		
Lactobacillus fermentum		33.99	0.28	XYL 1		
Bifidobacterium adolescentis		25.58	0.20			
03715112	**Good identification**					**03715112**
Bifidobacterium adolescentis		96.92	0.56	MNS 87		
Actinomyces israelii		1.84	0.33	ARA 87		
Lactobacillus fermentum		1.06	0.28			
03715120	**Low discrimination**					**03715120**
					GRA	
Lactobacillus fermentum		73.05	0.54	SAL 1	99	
Bacteroides vulgatus		5.87	0.14		1	
Prevotella melaninogenica		5.50	0.15		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03715121	**Low discrimination**					**03715121**
Actinomyces naeslundii		78.12	0.39	MNS 88		
Lactobacillus fermentum		21.89	0.20	XYL 1		
03715122	**Low discrimination**					**03715122**
					GRA	
Lactobacillus fermentum		41.50	0.54	SAL 1	99	
Bifidobacterium longum subsp. infantis		22.82	0.30		99	
Parabacteroides distasonis		19.90	0.45		1	
03715123	**Low discrimination**					**03715123**
Actinomyces naeslundii		63.75	0.39	MNS 88		
Lactobacillus fermentum		17.86	0.20	XYL 1		
Bifidobacterium longum subsp. infantis		9.82	-.04			
03715132	**Low discrimination**					**03715132**
					GRA	
Parabacteroides distasonis		80.27	0.41	MNS 98	1	
Bifidobacterium adolescentis		8.67	0.22		99	
Bifidobacterium longum subsp. longum		6.19	0.18		99	
03715140	**Low discrimination**					**03715140**
					GRA	
Actinomyces israelii		62.46	0.37	CEL 93	99	
Lactobacillus fermentum		20.84	0.28		99	
Fusobacterium mortiferum		13.79	0.23		1	
03715142	**Low discrimination**					**03715142**
Bifidobacterium adolescentis		62.04	0.39	MNS 87		
Actinomyces israelii		30.28	0.40	ARA 87		
Lactobacillus fermentum		6.74	0.28	SOR 11		
03715152	**Acceptable identification**					**03715152**
Bifidobacterium adolescentis		94.37	0.40	MNS 87		
Actinomyces israelii		5.63	0.26	ARA 87		
03715300	**Very good identification**					**03715300**
Eubacterium contortum		99.36	0.61	ESL 99		
Actinomyces odontolyticus		0.23	0.23			
Lactobacillus fermentum		0.16	0.28			
03715302	**Good identification**					**03715302**
Eubacterium contortum		96.27	0.48	ESL 99		
Lactobacillus fermentum		0.90	0.28			
Actinomyces israelii		0.78	0.28			
03715320	**Low discrimination**					**03715320**
					GRA	
Eubacterium contortum		81.04	0.48	ESL 99	99	
Bacteroides vulgatus		18.02	0.30		1	
Parabacteroides distasonis		0.36	0.19		1	
03715322	**Low discrimination**					**03715322**
					GRA	
Parabacteroides distasonis		66.91	0.54	MNS 98	1	
Eubacterium contortum		26.69	0.35		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		5.13	0.22		1	
03715332	**Good identification**					**03715332**
					GRA	
Parabacteroides distasonis		97.68	0.49	MNS 98	1	
Clostridium clostridioforme		1.59	0.13		50	
Eubacterium contortum		0.73	0.01		99	
03715400	**Very good identification**					**03715400**
Lactobacillus fermentum		96.32	0.58	SAL 1		
Actinomyces israelii		2.50	0.32			
Actinomyces meyeri		0.72	0.11			
03715402	**Low discrimination**					**03715402**
Lactobacillus fermentum		89.13	0.58	SAL 1		
Bifidobacterium adolescentis		7.40	0.34			
Actinomyces israelii		3.47	0.35			
03715412	**Low discrimination**					**03715412**
Bifidobacterium adolescentis		81.52	0.36	MNS 87		
Lactobacillus fermentum		8.12	0.24	LAC 99		
Actinomyces israelii		4.68	0.21			
03715420	**Good identification**					**03715420**
					GRA	
Lactobacillus fermentum		96.46	0.50	SAL 1	99	
Bifidobacterium longum subsp. longum		1.84	0.14		99	
Bacteroides vulgatus		1.70	-.01		1	
03715422	**Good identification**					**03715422**
Lactobacillus fermentum		97.41	0.50	SAL 1		
Bifidobacterium longum subsp. longum		1.86	0.14			
Actinomyces israelii		0.73	0.15			
03715500	**Low discrimination**					**03715500**
Lactobacillus fermentum		77.35	0.62	SAL 1		
Bifidobacterium longum subsp. longum		7.47	0.37			
Actinomyces israelii		4.93	0.42			
03715502	**Low discrimination**					**03715502**
Bifidobacterium adolescentis		80.72	0.69	MNS 87		
Lactobacillus fermentum		16.02	0.62			
Bifidobacterium longum subsp. longum		1.55	0.37			
03715510	**Low discrimination**					**03715510**
Bifidobacterium longum subsp. longum		66.77	0.41	MNS 87		
Bifidobacterium adolescentis		25.26	0.36	SAL 1		
Lactobacillus fermentum		4.10	0.28			
03715512	**Very good identification**					**03715512**
Bifidobacterium adolescentis		96.89	0.70	MNS 87		
Bifidobacterium longum subsp. longum		2.59	0.41			
Actinomyces israelii		0.23	0.31			
03715513	**Good identification**					**03715513**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		97.40	0.36	MNS 87		
Bifidobacterium longum subsp. longum		2.60	0.07	URE 1		
03715520	**Low discrimination**					**03715520**
					GRA	
Bifidobacterium longum subsp. longum		35.40	0.48	MNS 87	99	
Bacteroides vulgatus		32.62	0.33	SAL 1	1	
Lactobacillus fermentum		30.56	0.54		99	
03715522	**Low discrimination**					**03715522**
					GRA	
Bifidobacterium longum subsp. longum		43.71	0.48	MNS 87	99	
Lactobacillus fermentum		37.73	0.54	SAL 1	99	
Parabacteroides distasonis		11.09	0.42		1	
03715530	**Good identification**					**03715530**
					GRA	
Bifidobacterium longum subsp. longum		98.95	0.52	MNS 87	99	
Bacteroides vulgatus		0.54	-.01	SAL 1	1	
Lactobacillus fermentum		0.51	0.20		99	
03715532	**Low discrimination**					**03715532**
					GRA	
Bifidobacterium longum subsp. longum		83.78	0.52	MNS 87	99	
Bifidobacterium adolescentis		7.93	0.36	SAL 1	99	
Parabacteroides distasonis		6.72	0.37		1	
03715540	**Low discrimination**					**03715540**
Actinomyces israelii		60.18	0.35	CEL 93		
Lactobacillus fermentum		24.54	0.28			
Bifidobacterium adolescentis		15.28	0.19			
03715542	**Acceptable identification**					**03715542**
Bifidobacterium adolescentis		92.95	0.53	MNS 87		
Actinomyces israelii		5.55	0.38	SOR 11		
Lactobacillus fermentum		1.51	0.28			
03715552	**Good identification**					**03715552**
Bifidobacterium adolescentis		99.06	0.54	MNS 87		
Actinomyces israelii		0.72	0.24	SOR 11		
Bifidobacterium longum subsp. longum		0.22	0.07			
03715700	**Very good identification**					**03715700**
					GRA	
Eubacterium contortum		99.07	0.60	ESL 99	99	
Bacteroides vulgatus		0.64	0.15		1	
Lactobacillus fermentum		0.20	0.28		99	
03715702	**Acceptable identification**					**03715702**
Eubacterium contortum		91.20	0.47	ESL 99		
Bifidobacterium adolescentis		5.23	0.34			
Clostridium clostridioforme		1.48	0.22			
03715720	**Low discrimination**					**03715720**
					GRA	
Bacteroides vulgatus		78.12	0.49	MNS 99	1	
Eubacterium contortum		21.63	0.47		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		0.10	0.14		99	
03715722	**Low discrimination**					**03715722**
					GRA	
Parabacteroides distasonis		58.27	0.50	MNS 98	1	
Eubacterium contortum		31.02	0.34		99	
Bacteroides vulgatus		6.41	0.15		1	
03715732	**Low discrimination**					**03715732**
					GRA	
Parabacteroides distasonis		68.17	0.45	MNS 98	1	
Clostridium clostridioforme		28.29	0.33		50	
Bifidobacterium longum subsp. longum		2.86	0.18		99	
03717000	**Low discrimination**					**03717000**
Actinomyces israelii		88.37	0.46	CEL 93		
Lactobacillus fermentum		4.96	0.24			
Actinomyces naeslundii		4.38	0.33			
03717001	**Good identification**					**03717001**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03717002	**Acceptable identification**					**03717002**
Actinomyces israelii		90.25	0.49	CEL 93		
Bifidobacterium adolescentis		3.39	0.19			
Lactobacillus fermentum		3.38	0.24			
03717003	**Low discrimination**					**03717003**
Actinomyces naeslundii		83.06	0.36	MNS 88		
Actinomyces israelii		16.94	0.15	XYL 1		
03717012	**Low discrimination**					**03717012**
Actinomyces israelii		76.48	0.35	CEL 93		
Bifidobacterium adolescentis		23.52	0.20	MLZ 13		
03717021	**Good identification**					**03717021**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03717023	**Good identification**					**03717023**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03717040	**Good identification**					**03717040**
Actinomyces israelii		100.0	0.39	CEL 93		
03717042	**Good identification**					**03717042**
Actinomyces israelii		100.0	0.42	CEL 93		
03717100	**Good identification**					**03717100**
Actinomyces israelii		91.72	0.57	CEL 93		
Actinomyces naeslundii		3.23	0.41			
Lactobacillus fermentum		2.10	0.28			
03717101	**Low discrimination**					**03717101**
Actinomyces naeslundii		83.96	0.44	MNS 88		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		16.04	0.23	XYL 1		
03717102	**Low discrimination**					**03717102**
Actinomyces israelii		50.70	0.60	CEL 93		
Bifidobacterium adolescentis		47.19	0.53			
Actinomyces naeslundii		1.19	0.41			
03717103	**Low discrimination**					**03717103**
Actinomyces naeslundii		64.38	0.44	MNS 88		
Actinomyces israelii		18.45	0.26	XYL 1		
Bifidobacterium adolescentis		17.17	0.19			
03717110	**Low discrimination**					**03717110**
Actinomyces israelii		89.66	0.43	CEL 93		
Bifidobacterium adolescentis		10.34	0.20	MLZ 13		
03717112	**Low discrimination**					**03717112**
Bifidobacterium adolescentis		88.39	0.54	MNS 87		
Actinomyces israelii		11.61	0.46	ARA 87		
03717120	**Low discrimination**					**03717120**
Actinomyces israelii		59.81	0.36	CEL 93		
Actinomyces naeslundii		33.04	0.41	NAG 6		
Lactobacillus fermentum		7.15	0.20			
03717121	**Good identification**					**03717121**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03717122	**Low discrimination**					**03717122**
Actinomyces israelii		59.62	0.39	CEL 93		
Actinomyces naeslundii		21.95	0.41	NAG 6		
Bifidobacterium adolescentis		8.78	0.19			
03717123	**Good identification**					**03717123**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03717140	**Good identification**					**03717140**
Actinomyces israelii		97.71	0.50	CEL 93		
Actinomyces naeslundii		1.56	0.28			
Propionibacterium propionicum		0.73	0.12			
03717142	**Low discrimination**					**03717142**
Actinomyces israelii		76.11	0.53	CEL 93		
Bifidobacterium adolescentis		22.51	0.37			
Actinomyces naeslundii		0.81	0.28			
03717150	**Good identification**					**03717150**
Actinomyces israelii		100.0	0.36	CEL 93		
03717152	**Low discrimination**					**03717152**
Bifidobacterium adolescentis		70.76	0.39	MNS 87		
Actinomyces israelii		29.25	0.39	ARA 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03717302	**Low discrimination**					**03717302**
Actinomyces israelii		74.37	0.40	CEL 93		
Eubacterium contortum		16.35	0.14	RHA 7		
Bifidobacterium adolescentis		9.29	0.19			
03717400	**Acceptable identification**					**03717400**
Actinomyces israelii		93.58	0.45	CEL 93		
Lactobacillus fermentum		6.42	0.24			
03717402	**Low discrimination**					**03717402**
Actinomyces israelii		73.89	0.48	CEL 93		
Bifidobacterium adolescentis		22.73	0.33			
Lactobacillus fermentum		3.38	0.24			
03717440	**Good identification**					**03717440**
Actinomyces israelii		100.0	0.38	CEL 93		
03717442	**Acceptable identification**					**03717442**
Actinomyces israelii		91.10	0.41	CEL 93		
Bifidobacterium adolescentis		8.91	0.17			
03717500	**Low discrimination**					**03717500**
Actinomyces israelii		87.46	0.55	CEL 93		
Bifidobacterium adolescentis		10.09	0.33			
Lactobacillus fermentum		2.45	0.28			
03717502	**Low discrimination**					**03717502**
Bifidobacterium adolescentis		88.20	0.67	MNS 87		
Actinomyces israelii		11.59	0.58			
Lactobacillus fermentum		0.22	0.28			
03717510	**Low discrimination**					**03717510**
Actinomyces israelii		50.65	0.41	CEL 93		
Bifidobacterium adolescentis		47.79	0.34	MLZ 13		
Bifidobacterium longum subsp. longum		1.56	0.07			
03717512	**Very good identification**					**03717512**
Bifidobacterium adolescentis		98.39	0.69	MNS 87		
Actinomyces israelii		1.58	0.44			
Bifidobacterium longum subsp. longum		0.03	0.07			
03717540	**Good identification**					**03717540**
Actinomyces israelii		96.46	0.48	CEL 93		
Bifidobacterium adolescentis		3.54	0.17			
03717542	**Low discrimination**					**03717542**
Bifidobacterium adolescentis		70.76	0.51	MNS 87		
Actinomyces israelii		29.25	0.51	SOR 11		
03717552	**Good identification**					**03717552**
Bifidobacterium adolescentis		95.19	0.53	MNS 87		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		4.81	0.37	SOR 11			
03721000	**Low discrimination**						**03721000**
					GRA		
Actinomyces naeslundii		71.47	0.46	MNS 88	99		
Fusobacterium mortiferum		28.54	0.27	FRU 95	1		
03721001	**Good identification**						**03721001**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03721002	**Good identification**						**03721002**
					GRA		
Actinomyces naeslundii		95.75	0.46	MNS 88	99		
Fusobacterium mortiferum		4.25	0.11	FRU 95	1		
03721003	**Good identification**						**03721003**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03721020	**Good identification**						**03721020**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03721021	**Good identification**						**03721021**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03721022	**Good identification**						**03721022**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03721023	**Good identification**						**03721023**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03721041	**Good identification**						**03721041**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03721043	**Good identification**						**03721043**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03721061	**Good identification**						**03721061**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03721063	**Good identification**						**03721063**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03721100	**Low discrimination**						**03721100**
					COC	GRA	
Actinomyces naeslundii		73.83	0.53	MNS 88	1	99	
Fusobacterium mortiferum		24.17	0.33	FRU 95	1	1	
Blautia hansenii		2.00	-.03		99	99	
03721101	**Very good identification**						**03721101**
Actinomyces naeslundii		100.0	0.56	MNS 88			
03721102	**Good identification**						**03721102**
					GRA		
Actinomyces naeslundii		96.49	0.53	MNS 88	99		
Fusobacterium mortiferum		3.51	0.17	FRU 95	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03721103	**Very good identification**					**03721103**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03721120	**Acceptable identification**					**03721120**
					GRA	
Actinomyces naeslundii		92.01	0.53	MNS 88	99	
Clostridium perfringens		4.83	0.09	FRU 95	99	
Prevotella melaninogenica		3.17	-.01		1	
03721121	**Very good identification**					**03721121**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03721122	**Good identification**					**03721122**
Actinomyces naeslundii		97.59	0.53	MNS 88		
Clostridium perfringens		2.41	0.03	FRU 95		
03721123	**Very good identification**					**03721123**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03721140	**Good identification**					**03721140**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03721141	**Good identification**					**03721141**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03721142	**Good identification**					**03721142**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03721143	**Good identification**					**03721143**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03721160	**Good identification**					**03721160**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03721161	**Good identification**					**03721161**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03721162	**Good identification**					**03721162**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03721163	**Good identification**					**03721163**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03722101	**Good identification**					**03722101**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03722103	**Good identification**					**03722103**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03722121	**Good identification**					**03722121**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03722123	**Good identification**					**03722123**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03723000	**Good identification**					**03723000**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03723001	**Good identification**					**03723001**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03723002	**Good identification**					**03723002**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03723003	**Good identification**					**03723003**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03723020	**Good identification**					**03723020**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03723021	**Good identification**					**03723021**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03723022	**Good identification**					**03723022**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03723023	**Good identification**					**03723023**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03723040	**Good identification**					**03723040**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03723041	**Good identification**					**03723041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03723042	**Good identification**					**03723042**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03723043	**Good identification**					**03723043**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03723060	**Good identification**					**03723060**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03723061	**Good identification**					**03723061**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03723062	**Good identification**					**03723062**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03723063	**Good identification**					**03723063**
Actinomyces naeslundii		100.0	0.41	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03723100	**Very good identification**					**03723100**
Actinomyces naeslundii		99.08	0.58	MNS 88		
Actinomyces israelii		0.92	0.14	FRU 95		
03723101	**Very good identification**					**03723101**
Actinomyces naeslundii		100.0	0.61	MNS 88		
03723102	**Very good identification**					**03723102**
Actinomyces naeslundii		97.67	0.58	MNS 88		
Actinomyces israelii		1.36	0.17	FRU 95		
Propionibacterium propionicum		0.97	0.14			
03723103	**Very good identification**					**03723103**
Actinomyces naeslundii		100.0	0.61	MNS 88		
03723110	**Good identification**					**03723110**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03723111	**Good identification**					**03723111**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03723112	**Good identification**					**03723112**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03723113	**Good identification**					**03723113**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03723120	**Very good identification**					**03723120**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03723121	**Very good identification**					**03723121**
Actinomyces naeslundii		100.0	0.61	MNS 88		
03723122	**Very good identification**					**03723122**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03723123	**Very good identification**					**03723123**
Actinomyces naeslundii		100.0	0.61	MNS 88		
03723130	**Good identification**					**03723130**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03723131	**Good identification**					**03723131**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03723132	**Good identification**					**03723132**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03723133	**Good identification**					**03723133**
Actinomyces naeslundii		100.0	0.39	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03723140	**Good identification**						**03723140**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03723141	**Good identification**						**03723141**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03723142	**Acceptable identification**						**03723142**
Actinomyces naeslundii		94.67	0.45	MNS 88			
Propionibacterium propionicum		5.33	0.14	FRU 95			
03723143	**Good identification**						**03723143**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03723160	**Good identification**						**03723160**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03723161	**Good identification**						**03723161**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03723162	**Good identification**						**03723162**
Actinomyces naeslundii		100.0	0.45	MNS 88			
03723163	**Good identification**						**03723163**
Actinomyces naeslundii		100.0	0.48	MNS 88			
03727102	**Low discrimination**						**03727102**
Actinomyces israelii		81.42	0.35	FRU 99			
Bifidobacterium adolescentis		13.43	0.16	CEL 93			
Actinomyces naeslundii		5.15	0.24				
03730000	**Very good identification**						**03730000**
					GRA	CAT	
Fusobacterium mortiferum		95.05	0.58	MLT 11	1	5	
Actinomyces naeslundii		2.86	0.44		99	25	
Propionibacterium granulosum		1.87	0.26		99	90	
03730001	**Low discrimination**						**03730001**
					GRA		
Actinomyces naeslundii		81.69	0.47	MNS 88	99		
Fusobacterium mortiferum		18.31	0.24	GAL 96	1		
03730002	**Low discrimination**						**03730002**
					GRA		
Fusobacterium mortiferum		72.55	0.42	MLT 11	1		
Actinomyces naeslundii		19.62	0.44	BGL 10	99		
Actinomyces viscosus		4.63	0.09		99		
03730003	**Good identification**						**03730003**
Actinomyces naeslundii		95.02	0.47	MNS 88			
Actinomyces viscosus		4.98	0.01	GAL 96			
03730020	**Low discrimination**						**03730020**
					GRA	CAT	
Actinomyces naeslundii		66.68	0.44	MNS 88	99	25	
Fusobacterium mortiferum		22.42	0.24	GAL 96	1	5	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		10.90	0.15		99	90	
03730021	**Good identification**						**03730021**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03730022	**Good identification**						**03730022**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03730023	**Good identification**						**03730023**
Actinomyces naeslundii		100.0	0.47	MNS 88			
03730100	**Very good identification**						**03730100**
					GRA		
Fusobacterium mortiferum		96.47	0.65	MLT 11	1		
Actinomyces naeslundii		3.54	0.52		99		
03730101	**Low discrimination**						**03730101**
					GRA		
Actinomyces naeslundii		84.48	0.55	MNS 88	99		
Fusobacterium mortiferum		15.52	0.30	GAL 96	1		
03730102	**Low discrimination**						**03730102**
					GRA		
Fusobacterium mortiferum		72.13	0.48	MLT 11	1		
Actinomyces naeslundii		23.79	0.52	BGL 10	99		
Actinomyces viscosus		3.51	0.13		99		
03730103	**Acceptable identification**						**03730103**
					GRA		
Actinomyces naeslundii		94.95	0.55	MNS 88	99		
Actinomyces viscosus		3.11	0.05	GAL 96	99		
Fusobacterium mortiferum		1.94	0.14		1		
03730120	**Low discrimination**						**03730120**
					GRA		
Actinomyces naeslundii		74.52	0.52	MNS 88	99		
Fusobacterium mortiferum		20.54	0.30	GAL 96	1		
Clostridium perfringens		4.94	0.09		99		
03730121	**Good identification**						**03730121**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03730122	**Acceptable identification**						**03730122**
					GRA		
Actinomyces naeslundii		94.18	0.52	MNS 88	99		
Clostridium perfringens		2.94	0.03	GAL 96	99		
Fusobacterium mortiferum		2.88	0.14		1		
03730123	**Good identification**						**03730123**
Actinomyces naeslundii		100.0	0.55	MNS 88			
03730141	**Good identification**						**03730141**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03730142	**Low discrimination**						**03730142**
					GRA		
Actinomyces naeslundii		76.24	0.39	MNS 88	99		
Fusobacterium mortiferum		13.23	0.14	GAL 96	1		
Propionibacterium propionicum		10.53	0.14		99		

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
03730143	**Good identification**						**03730143**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03730160	**Good identification**						**03730160**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03730161	**Good identification**						**03730161**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03730162	**Good identification**						**03730162**
Actinomyces naeslundii		100.0	0.39	MNS 88			
03730163	**Good identification**						**03730163**
Actinomyces naeslundii		100.0	0.42	MNS 88			
03731000	**Low discrimination**						**03731000**
					GRA	CAT	
Fusobacterium mortiferum		65.97	0.61	MLT 11	1	5	
Actinomyces naeslundii		31.71	0.67		99	25	
Propionibacterium granulosum		0.86	0.26		99	90	
03731001	**Very good identification**						**03731001**
					GRA		
Actinomyces naeslundii		98.62	0.70	MNS 88	99		
Fusobacterium mortiferum		1.38	0.27		1		
03731002	**Low discrimination**						**03731002**
					GRA		
Actinomyces naeslundii		77.52	0.67	MNS 88	99		
Fusobacterium mortiferum		17.92	0.45		1		
Lactobacillus fermentum		1.51	0.28		99		
03731003	**Very good identification**						**03731003**
					GRA		
Actinomyces naeslundii		99.85	0.70	MNS 88	99		
Fusobacterium mortiferum		0.16	0.11		1		
03731010	**Low discrimination**						**03731010**
					GRA	CAT	
Actinomyces naeslundii		65.81	0.46	MNS 88	99	25	
Fusobacterium mortiferum		26.28	0.27	MLZ 5	1	5	
Propionibacterium granulosum		4.65	0.11		99	90	
03731011	**Good identification**						**03731011**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03731012	**Acceptable identification**						**03731012**
					GRA		
Actinomyces naeslundii		91.37	0.46	MNS 88	99		
Actinomyces israelii		4.58	0.14	MLZ 5	99		
Fusobacterium mortiferum		4.06	0.11		1		
03731013	**Good identification**						**03731013**
Actinomyces naeslundii		100.0	0.49	MNS 88			
03731020	**Low discrimination**						**03731020**
					GRA		
Actinomyces naeslundii		87.68	0.67	MNS 88	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		9.31	0.28		99	
Fusobacterium mortiferum		1.84	0.27		1	
03731021	**Very good identification**					**03731021**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03731022	**Good identification**					**03731022**
Actinomyces naeslundii		93.66	0.67	MNS 88		
Clostridium perfringens		4.68	0.23			
Lactobacillus fermentum		0.61	0.20			
03731023	**Very good identification**					**03731023**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03731030	**Good identification**					**03731030**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03731031	**Good identification**					**03731031**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03731032	**Good identification**					**03731032**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03731033	**Good identification**					**03731033**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03731040	**Low discrimination**					**03731040**
					GRA	
Actinomyces naeslundii		87.34	0.55	MNS 88	99	
Fusobacterium mortiferum		10.40	0.27		1	
Actinomyces israelii		2.27	0.18		99	
03731041	**Very good identification**					**03731041**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03731042	**Good identification**					**03731042**
					GRA	
Actinomyces naeslundii		95.05	0.55	MNS 88	99	
Actinomyces israelii		3.70	0.21		99	
Fusobacterium mortiferum		1.26	0.11		1	
03731043	**Very good identification**					**03731043**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03731051	**Good identification**					**03731051**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731053	**Good identification**					**03731053**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731060	**Low discrimination**					**03731060**
Actinomyces naeslundii		89.72	0.55	MNS 88		
Clostridium perfringens		10.28	0.16			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03731061	**Very good identification**					**03731061**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03731062	**Acceptable identification**					**03731062**
Actinomyces naeslundii		94.88	0.55	MNS 88		
Clostridium perfringens		5.12	0.11			
03731063	**Very good identification**					**03731063**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03731071	**Good identification**					**03731071**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731073	**Good identification**					**03731073**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731100	**Low discrimination**					**03731100**
					GRA	
Fusobacterium mortiferum		62.08	0.68	MLT 11	1	
Actinomyces naeslundii		36.40	0.75		99	
Actinomyces israelii		0.60	0.35		99	
03731101	**Very good identification**					**03731101**
					GRA	
Actinomyces naeslundii		98.86	0.78	MNS 88	99	
Fusobacterium mortiferum		1.14	0.33		1	
03731102	**Low discrimination**					**03731102**
					GRA	
Actinomyces naeslundii		76.86	0.75	MNS 88	99	
Fusobacterium mortiferum		14.57	0.51		1	
Bifidobacterium adolescentis		3.74	0.37		99	
03731103	**Very good identification**					**03731103**
					GRA	
Actinomyces naeslundii		99.79	0.78	MNS 88	99	
Fusobacterium mortiferum		0.13	0.17		1	
Actinomyces viscosus		0.08	0.01		99	
03731110	**Low discrimination**					**03731110**
					GRA	
Actinomyces naeslundii		72.76	0.53	MNS 88	99	
Fusobacterium mortiferum		23.82	0.33	MLZ 5	1	
Actinomyces israelii		3.42	0.21		99	
03731111	**Very good identification**					**03731111**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03731112	**Low discrimination**					**03731112**
Bifidobacterium adolescentis		50.13	0.39	MNS 87		
Actinomyces naeslundii		44.32	0.53	XYL 87		
Actinomyces israelii		3.12	0.24	ARA 87		
03731113	**Very good identification**					**03731113**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03731120	**Low discrimination**					**03731120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		75.99	0.75	MNS 88	99	
Clostridium perfringens		20.79	0.43		99	
Fusobacterium mortiferum		1.31	0.33		1	
03731121	**Very good identification**					**03731121**
Actinomyces naeslundii		99.82	0.78	MNS 88		
Clostridium perfringens		0.18	0.09			
03731122	**Low discrimination**					**03731122**
					GRA	
Actinomyces naeslundii		80.46	0.75	MNS 88	99	
Clostridium perfringens		10.36	0.38		99	
Parabacteroides distasonis		4.23	0.40		1	
03731123	**Very good identification**					**03731123**
Actinomyces naeslundii		99.91	0.78	MNS 88		
Clostridium perfringens		0.09	0.03			
03731130	**Good identification**					**03731130**
Actinomyces naeslundii		95.01	0.53	MNS 88		
Clostridium perfringens		4.99	0.09	MLZ 5		
03731131	**Very good identification**					**03731131**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03731132	**Low discrimination**					**03731132**
					GRA	
Actinomyces naeslundii		64.00	0.53	MNS 88	99	
Parabacteroides distasonis		34.41	0.36	MLZ 5	1	
Clostridium perfringens		1.58	0.03		99	
03731133	**Very good identification**					**03731133**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03731140	**Low discrimination**					**03731140**
					GRA	
Actinomyces naeslundii		87.47	0.62	MNS 88	99	
Fusobacterium mortiferum		8.54	0.33		1	
Actinomyces israelii		3.19	0.28		99	
03731141	**Very good identification**					**03731141**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03731142	**Low discrimination**					**03731142**
Actinomyces naeslundii		89.63	0.62	MNS 88		
Actinomyces israelii		4.90	0.31			
Bifidobacterium adolescentis		3.06	0.22			
03731143	**Very good identification**					**03731143**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03731150	**Acceptable identification**					**03731150**
Actinomyces naeslundii		90.62	0.40	MNS 88		
Actinomyces israelii		9.38	0.14	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03731151	**Good identification**					**03731151**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03731152	**Low discrimination**					**03731152**
Actinomyces naeslundii		51.35	0.40	MNS 88		
Bifidobacterium adolescentis		40.68	0.23	MLZ 5		
Actinomyces israelii		7.97	0.17			
03731153	**Good identification**					**03731153**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03731160	**Low discrimination**					**03731160**
Actinomyces naeslundii		77.21	0.62	MNS 88		
Clostridium perfringens		22.79	0.31			
03731161	**Very good identification**					**03731161**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03731162	**Low discrimination**					**03731162**
Actinomyces naeslundii		87.01	0.62	MNS 88		
Clostridium perfringens		12.09	0.25			
Lactobacillus acidophilus		0.90	0.04			
03731163	**Very good identification**					**03731163**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03731170	**Good identification**					**03731170**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03731171	**Good identification**					**03731171**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03731172	**Good identification**					**03731172**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03731173	**Good identification**					**03731173**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03731201	**Good identification**					**03731201**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731203	**Good identification**					**03731203**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731221	**Good identification**					**03731221**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731223	**Good identification**					**03731223**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731301	**Good identification**					**03731301**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	MNS 88		
03731302	**Low discrimination**					**03731302**
					GRA	
Actinomyces naeslundii		60.91	0.41	MNS 88	99	
Fusobacterium mortiferum		11.54	0.17	RHA 1	1	
Actinomyces israelii		11.26	0.19		99	
03731303	**Good identification**					**03731303**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03731320	**Low discrimination**					**03731320**
					SPO	
Actinomyces naeslundii		61.16	0.41	MNS 88	1	
Clostridium perfringens		16.73	0.09	RHA 1	84	
Clostridium ramosum		12.46	0.04		92	
03731321	**Good identification**					**03731321**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03731322	**Low discrimination**					**03731322**
					GRA	
Parabacteroides distasonis		82.66	0.48	MNS 98	1	
Prevotella oralis		6.47	0.17	XYL 90	1	
Actinomyces naeslundii		5.30	0.41		99	
03731323	**Acceptable identification**					**03731323**
					GRA	
Actinomyces naeslundii		90.49	0.44	MNS 88	99	
Parabacteroides distasonis		9.51	0.14	RHA 1	1	
03731332	**Good identification**					**03731332**
Parabacteroides distasonis		97.73	0.44	MNS 98		
Clostridium clostridioforme		2.27	0.10	XYL 90		
03731401	**Good identification**					**03731401**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731403	**Good identification**					**03731403**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731421	**Good identification**					**03731421**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731423	**Good identification**					**03731423**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03731501	**Good identification**					**03731501**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03731502	**Low discrimination**					**03731502**
Bifidobacterium adolescentis		86.78	0.51	MNS 87		
Actinomyces israelii		5.41	0.37	XYL 87		
Lactobacillus fermentum		2.97	0.32			
03731503	**Low discrimination**					**03731503**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		82.15	0.44	MNS 88			
Bifidobacterium adolescentis		17.85	0.17	ARA 1			
03731512	**Good identification**						**03731512**
Bifidobacterium adolescentis		96.10	0.53	MNS 87			
Bifidobacterium longum subsp. longum		2.15	0.23	XYL 87			
Clostridium clostridioforme		1.02	0.16				
03731521	**Good identification**						**03731521**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03731522	**Low discrimination**						**03731522**
					GRA		
Parabacteroides distasonis		43.68	0.37	MNS 98	1		
Bifidobacterium longum subsp. longum		28.34	0.29	XYL 90	99		
Actinomyces naeslundii		13.70	0.41		99		
03731523	**Good identification**						**03731523**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03731542	**Low discrimination**						**03731542**
Bifidobacterium adolescentis		80.63	0.36	MNS 87			
Actinomyces israelii		15.80	0.30	XYL 87			
Actinomyces naeslundii		3.57	0.28	SOR 11			
03731552	**Good identification**						**03731552**
Bifidobacterium adolescentis		97.66	0.37	MNS 87			
Actinomyces israelii		2.34	0.16	XYL 87			
03731722	**Acceptable identification**						**03731722**
Parabacteroides distasonis		93.43	0.45	MNS 98			
Clostridium clostridioforme		6.57	0.19	XYL 90			
03731732	**Low discrimination**						**03731732**
Parabacteroides distasonis		62.71	0.40	MNS 98			
Clostridium clostridioforme		37.29	0.31	XYL 90			
03732000	**Low discrimination**						**03732000**
					GRA	CAT	
Actinomyces naeslundii		70.06	0.49	MNS 88	99	25	
Propionibacterium granulosum		17.81	0.24	GAL 96	99	90	
Fusobacterium mortiferum		12.14	0.24		1	5	
03732001	**Good identification**						**03732001**
Actinomyces naeslundii		100.0	0.52	MNS 88			
03732002	**Acceptable identification**						**03732002**
					CAT		
Actinomyces naeslundii		90.97	0.49	MNS 88	25		
Propionibacterium granulosum		5.78	0.13	GAL 96	90		
Propionibacterium propionicum		3.25	0.14		1		
03732003	**Good identification**						**03732003**
Actinomyces naeslundii		100.0	0.52	MNS 88			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03732020	**Acceptable identification**					**03732020**
Actinomyces naeslundii		94.03	0.49	MNS 88		
Propionibacterium granulosum		5.97	0.13	GAL 96		
03732021	**Good identification**					**03732021**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03732022	**Good identification**					**03732022**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03732023	**Good identification**					**03732023**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03732040	**Good identification**					**03732040**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03732041	**Good identification**					**03732041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03732042	**Low discrimination**					**03732042**
Actinomyces naeslundii		83.16	0.36	MNS 88		
Propionibacterium propionicum		16.84	0.14	GAL 96		
03732043	**Good identification**					**03732043**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03732060	**Good identification**					**03732060**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03732061	**Good identification**					**03732061**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03732062	**Good identification**					**03732062**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03732063	**Good identification**					**03732063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03732100	**Low discrimination**					**03732100**
					GRA	
Actinomyces naeslundii		50.44	0.57	MNS 88	99	
Propionibacterium propionicum		41.81	0.45	GAL 96	99	
Fusobacterium mortiferum		7.16	0.30		1	
03732101	**Good identification**					**03732101**
Actinomyces naeslundii		94.22	0.60	MNS 88		
Propionibacterium propionicum		5.79	0.29	GAL 96		
03732102	**Low discrimination**					**03732102**
Propionibacterium propionicum		54.61	0.48	NIT 86		
Actinomyces naeslundii		43.92	0.57	SAL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	0.77	0.17			
03732103	**Good identification**					**03732103**
	Actinomyces naeslundii	91.57	0.60	MNS 88		
	Propionibacterium propionicum	8.43	0.32	GAL 96		
03732111	**Good identification**					**03732111**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03732113	**Good identification**					**03732113**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03732120	**Very good identification**					**03732120**
	Actinomyces naeslundii	100.0	0.57	MNS 88		
03732121	**Very good identification**					**03732121**
	Actinomyces naeslundii	100.0	0.60	MNS 88		
03732122	**Very good identification**					**03732122**
	Actinomyces naeslundii	98.76	0.57	MNS 88		
	Propionibacterium propionicum	1.24	0.14	GAL 96		
03732123	**Very good identification**					**03732123**
	Actinomyces naeslundii	100.0	0.60	MNS 88		
03732131	**Good identification**					**03732131**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03732133	**Good identification**					**03732133**
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03732140	**Low discrimination**					**03732140**
	Propionibacterium propionicum	82.45	0.45	NIT 86		
	Actinomyces naeslundii	17.55	0.44	SAL 1		
03732141	**Low discrimination**					**03732141**
	Actinomyces naeslundii	74.19	0.47	MNS 88		
	Propionibacterium propionicum	25.81	0.29	GAL 96		
03732142	**Low discrimination**					**03732142**
	Propionibacterium propionicum	87.57	0.48	NIT 86		
	Actinomyces naeslundii	12.43	0.44	SAL 1		
03732143	**Low discrimination**					**03732143**
	Actinomyces naeslundii	65.71	0.47	MNS 88		
	Propionibacterium propionicum	34.29	0.32	GAL 96		
03732160	**Good identification**					**03732160**
	Actinomyces naeslundii	100.0	0.44	MNS 88		
03732161	**Good identification**					**03732161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	MNS 88		
03732162	**Acceptable identification**					**03732162**
Actinomyces naeslundii		93.36	0.44	MNS 88		
Propionibacterium propionicum		6.64	0.14	GAL 96		
03732163	**Good identification**					**03732163**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03733000	**Good identification**					**03733000**
					GRA	
Actinomyces naeslundii		94.72	0.72	MNS 88	99	
Actinomyces israelii		3.25	0.38		99	
Fusobacterium mortiferum		1.03	0.27		1	
03733001	**Very good identification**					**03733001**
Actinomyces naeslundii		100.0	0.75	MNS 88		
03733002	**Good identification**					**03733002**
					CAT	
Actinomyces naeslundii		94.62	0.72	MNS 88	25	
Actinomyces israelii		4.88	0.41		1	
Propionibacterium granulosum		0.25	0.13		90	
03733003	**Very good identification**					**03733003**
Actinomyces naeslundii		100.0	0.75	MNS 88		
03733010	**Low discrimination**					**03733010**
					CAT	
Actinomyces naeslundii		88.89	0.50	MNS 88	25	
Actinomyces israelii		8.67	0.24	MLZ 5	1	
Propionibacterium granulosum		2.44	0.09		90	
03733011	**Good identification**					**03733011**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03733012	**Low discrimination**					**03733012**
Actinomyces naeslundii		87.24	0.50	MNS 88		
Actinomyces israelii		12.76	0.27	MLZ 5		
03733013	**Good identification**					**03733013**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03733020	**Very good identification**					**03733020**
					CAT	
Actinomyces naeslundii		99.52	0.72	MNS 88	25	
Propionibacterium granulosum		0.26	0.13		90	
Actinomyces israelii		0.22	0.18		1	
03733021	**Very good identification**					**03733021**
Actinomyces naeslundii		100.0	0.75	MNS 88		
03733022	**Very good identification**					**03733022**
Actinomyces naeslundii		99.67	0.72	MNS 88		
Actinomyces israelii		0.33	0.21			
03733023	**Very good identification**					**03733023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.75	MNS 88		
03733030	**Good identification**					**03733030**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03733031	**Good identification**					**03733031**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03733032	**Good identification**					**03733032**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03733033	**Good identification**					**03733033**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03733040	**Good identification**					**03733040**
Actinomyces naeslundii		92.96	0.59	MNS 88		
Actinomyces israelii		7.04	0.31			
03733041	**Very good identification**					**03733041**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03733042	**Low discrimination**					**03733042**
Actinomyces naeslundii		89.13	0.59	MNS 88		
Actinomyces israelii		10.12	0.34			
Propionibacterium propionicum		0.75	0.14			
03733043	**Very good identification**					**03733043**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03733050	**Low discrimination**					**03733050**
Actinomyces naeslundii		82.31	0.38	MNS 88		
Actinomyces israelii		17.69	0.17	MLZ 5		
03733051	**Good identification**					**03733051**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733052	**Low discrimination**					**03733052**
Actinomyces naeslundii		75.62	0.38	MNS 88		
Actinomyces israelii		24.38	0.20	MLZ 5		
03733053	**Good identification**					**03733053**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733060	**Very good identification**					**03733060**
Actinomyces naeslundii		100.0	0.59	MNS 88		
03733061	**Very good identification**					**03733061**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03733062	**Very good identification**					**03733062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.28	0.59	MNS 88		
Actinomyces israelii		0.72	0.14			
03733063	**Very good identification**					**03733063**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03733070	**Good identification**					**03733070**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03733071	**Good identification**					**03733071**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733072	**Good identification**					**03733072**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03733073	**Good identification**					**03733073**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733100	**Good identification**					**03733100**
Actinomyces naeslundii		91.60	0.80	MNS 88		
Actinomyces israelii		4.42	0.48			
Propionibacterium propionicum		3.16	0.45			
03733101	**Very good identification**					**03733101**
Actinomyces naeslundii		99.71	0.83	MNS 88		
Propionibacterium propionicum		0.26	0.29			
Actinomyces israelii		0.03	0.14			
03733102	**Low discrimination**					**03733102**
Actinomyces naeslundii		87.28	0.80	MNS 88		
Actinomyces israelii		6.32	0.51			
Propionibacterium propionicum		4.52	0.48			
03733103	**Very good identification**					**03733103**
Actinomyces naeslundii		99.57	0.83	MNS 88		
Propionibacterium propionicum		0.38	0.32			
Actinomyces israelii		0.05	0.17			
03733110	**Low discrimination**					**03733110**
Actinomyces naeslundii		87.94	0.58	MNS 88		
Actinomyces israelii		12.06	0.34	MLZ 5		
03733111	**Very good identification**					**03733111**
Actinomyces naeslundii		100.0	0.61	MNS 88		
03733112	**Low discrimination**					**03733112**
Actinomyces naeslundii		59.09	0.58	MNS 88		
Bifidobacterium adolescentis		28.17	0.37	MLZ 5		
Actinomyces israelii		12.15	0.37			
03733113	**Very good identification**					**03733113**
Actinomyces naeslundii		100.0	0.61	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03733120	**Very good identification**					**03733120**
					SPO	
Actinomyces naeslundii		99.10	0.80	MNS 88	1	
Clostridium ramosum		0.45	0.15		92	
Actinomyces israelii		0.31	0.28		1	
03733121	**Very good identification**					**03733121**
Actinomyces naeslundii		100.0	0.83	MNS 88		
03733122	**Very good identification**					**03733122**
					SPO	
Actinomyces naeslundii		97.66	0.80	MNS 88	1	
Clostridium ramosum		1.77	0.25		92	
Actinomyces israelii		0.45	0.31		1	
03733123	**Very good identification**					**03733123**
Actinomyces naeslundii		100.0	0.83	MNS 88		
03733130	**Very good identification**					**03733130**
Actinomyces naeslundii		99.13	0.58	MNS 88		
Actinomyces israelii		0.87	0.14	MLZ 5		
03733131	**Very good identification**					**03733131**
Actinomyces naeslundii		100.0	0.61	MNS 88		
03733132	**Very good identification**					**03733132**
Actinomyces naeslundii		98.70	0.58	MNS 88		
Actinomyces israelii		1.30	0.17	MLZ 5		
03733133	**Very good identification**					**03733133**
Actinomyces naeslundii		100.0	0.61	MNS 88		
03733140	**Low discrimination**					**03733140**
Actinomyces naeslundii		76.80	0.67	MNS 88		
Propionibacterium propionicum		15.03	0.45			
Actinomyces israelii		8.17	0.41			
03733141	**Very good identification**					**03733141**
Actinomyces naeslundii		98.57	0.70	MNS 88		
Propionibacterium propionicum		1.43	0.29			
03733142	**Low discrimination**					**03733142**
Actinomyces naeslundii		68.14	0.67	MNS 88		
Propionibacterium propionicum		20.00	0.48			
Actinomyces israelii		10.88	0.44			
03733143	**Very good identification**					**03733143**
Actinomyces naeslundii		97.87	0.70	MNS 88		
Propionibacterium propionicum		2.13	0.32			
03733150	**Low discrimination**					**03733150**
Actinomyces naeslundii		76.80	0.45	MNS 88		
Actinomyces israelii		23.20	0.27	MLZ 5		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03733151	**Good identification**					**03733151**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03733152	**Low discrimination**					**03733152**
Actinomyces naeslundii		54.25	0.45	MNS 88		
Actinomyces israelii		24.58	0.30	MLZ 5		
Bifidobacterium adolescentis		18.11	0.22			
03733153	**Good identification**					**03733153**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03733160	**Very good identification**					**03733160**
Actinomyces naeslundii		99.33	0.67	MNS 88		
Actinomyces israelii		0.68	0.21			
03733161	**Very good identification**					**03733161**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03733162	**Very good identification**					**03733162**
Actinomyces naeslundii		98.70	0.67	MNS 88		
Actinomyces israelii		1.01	0.24			
Propionibacterium propionicum		0.29	0.14			
03733163	**Very good identification**					**03733163**
Actinomyces naeslundii		100.0	0.70	MNS 88		
03733170	**Good identification**					**03733170**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03733171	**Good identification**					**03733171**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03733172	**Good identification**					**03733172**
Actinomyces naeslundii		100.0	0.45	MNS 88		
03733173	**Good identification**					**03733173**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03733200	**Low discrimination**					**03733200**
Actinomyces naeslundii		79.62	0.38	MNS 88		
Actinomyces israelii		20.38	0.19	RHA 1		
03733201	**Good identification**					**03733201**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733202	**Low discrimination**					**03733202**
Actinomyces naeslundii		72.25	0.38	MNS 88		
Actinomyces israelii		27.75	0.22	RHA 1		
03733203	**Good identification**					**03733203**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733220	**Good identification**					**03733220**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03733221	**Good identification**					**03733221**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733222	**Good identification**					**03733222**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03733223	**Good identification**					**03733223**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733300	**Low discrimination**					**03733300**
Actinomyces naeslundii		73.54	0.46	MNS 88		
Actinomyces israelii		26.46	0.29	RHA 1		
03733301	**Good identification**					**03733301**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03733302	**Low discrimination**					**03733302**
Actinomyces naeslundii		62.83	0.46	MNS 88		
Actinomyces israelii		33.91	0.32	RHA 1		
Propionibacterium propionicum		3.26	0.14			
03733303	**Good identification**					**03733303**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03733320	**Low discrimination**					**03733320**
					SPO	
Actinomyces naeslundii		62.70	0.46	MNS 88	1	
Clostridium ramosum		37.30	0.17	RHA 1	92	
03733321	**Good identification**					**03733321**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03733323	**Good identification**					**03733323**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03733341	**Good identification**					**03733341**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03733343	**Good identification**					**03733343**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03733361	**Good identification**					**03733361**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03733363	**Good identification**					**03733363**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03733400	**Low discrimination**					**03733400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		73.57	0.37	CEL 93		
Actinomyces naeslundii		26.44	0.38	XYL 92		
03733401	**Good identification**					**03733401**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733402	**Low discrimination**					**03733402**
Actinomyces israelii		75.01	0.40	CEL 93		
Actinomyces naeslundii		17.97	0.38	XYL 92		
Bifidobacterium adolescentis		7.03	0.16			
03733403	**Good identification**					**03733403**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733420	**Low discrimination**					**03733420**
Actinomyces naeslundii		84.92	0.38	MNS 88		
Actinomyces israelii		15.09	0.16	ARA 1		
03733421	**Good identification**					**03733421**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733422	**Low discrimination**					**03733422**
Actinomyces naeslundii		78.96	0.38	MNS 88		
Actinomyces israelii		21.04	0.19	ARA 1		
03733423	**Good identification**					**03733423**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03733500	**Low discrimination**					**03733500**
Actinomyces israelii		77.47	0.47	CEL 93		
Actinomyces naeslundii		19.81	0.46	XYL 92		
Bifidobacterium adolescentis		2.72	0.16			
03733501	**Good identification**					**03733501**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03733502	**Low discrimination**					**03733502**
Bifidobacterium adolescentis		66.28	0.50	MNS 87		
Actinomyces israelii		28.59	0.50	XYL 87		
Actinomyces naeslundii		4.87	0.46			
03733503	**Low discrimination**					**03733503**
Actinomyces naeslundii		88.41	0.49	MNS 88		
Bifidobacterium adolescentis		8.10	0.16	ARA 1		
Actinomyces israelii		3.49	0.16			
03733512	**Acceptable identification**					**03733512**
Bifidobacterium adolescentis		94.71	0.51	MNS 87		
Actinomyces israelii		5.00	0.36	XYL 87		
Actinomyces naeslundii		0.30	0.24			
03733520	**Low discrimination**					**03733520**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		80.02	0.46	MNS 88		
Actinomyces israelii		19.98	0.26	ARA 1		
03733521	**Good identification**					**03733521**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03733522	**Low discrimination**					**03733522**
Actinomyces naeslundii		66.14	0.46	MNS 88		
Actinomyces israelii		24.77	0.29	ARA 1		
Bifidobacterium adolescentis		9.09	0.16			
03733523	**Good identification**					**03733523**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03733540	**Low discrimination**					**03733540**
Actinomyces israelii		89.60	0.40	CEL 93		
Actinomyces naeslundii		10.40	0.33	XYL 92		
03733541	**Good identification**					**03733541**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03733542	**Low discrimination**					**03733542**
Actinomyces israelii		54.44	0.43	CEL 93		
Bifidobacterium adolescentis		40.11	0.34	XYL 92		
Actinomyces naeslundii		4.21	0.33			
03733543	**Good identification**					**03733543**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03733552	**Low discrimination**					**03733552**
Bifidobacterium adolescentis		85.77	0.36	MNS 87		
Actinomyces israelii		14.23	0.29	XYL 87		
03733561	**Good identification**					**03733561**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03733563	**Good identification**					**03733563**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03734000	**Good identification**					**03734000**
Fusobacterium mortiferum		100.0	0.39	MLT 11		
03734100	**Good identification**					**03734100**
					GRA	
Fusobacterium mortiferum		97.80	0.45	MLT 11	1	
Actinomyces israelii		2.20	0.19	XYL 7	99	
03734712	**Good identification**					**03734712**
Clostridium clostridioforme		100.0	0.40	MNS 99		
03734732	**Good identification**					**03734732**
Clostridium clostridioforme		95.66	0.40	MNS 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		4.34	0.22				
03735000	**Low discrimination**						**03735000**
					GRA		
Fusobacterium mortiferum		49.95	0.42	MLT 11	1		
Actinomyces israelii		43.18	0.43	XYL 7	99		
Lactobacillus fermentum		3.65	0.24		99		
03735001	**Good identification**						**03735001**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03735002	**Low discrimination**						**03735002**
					GRA		
Actinomyces israelii		80.78	0.46	CEL 93	99		
Fusobacterium mortiferum		6.92	0.26		1		
Lactobacillus fermentum		4.55	0.24		99		
03735003	**Good identification**						**03735003**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03735021	**Good identification**						**03735021**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03735023	**Good identification**						**03735023**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03735040	**Good identification**						**03735040**
Actinomyces israelii		100.0	0.36	CEL 93			
03735042	**Good identification**						**03735042**
Actinomyces israelii		100.0	0.39	CEL 93			
03735100	**Low discrimination**						**03735100**
					COC	GRA	
Actinomyces israelii		54.08	0.54	CEL 93	1	99	
Fusobacterium mortiferum		36.50	0.48		1	1	
Atopobium parvulum		3.02	0.18		99	99	
03735101	**Low discrimination**						**03735101**
					GRA		
Actinomyces naeslundii		82.48	0.44	MNS 88	99		
Actinomyces israelii		10.46	0.19	XYL 1	99		
Fusobacterium mortiferum		7.06	0.14		1		
03735102	**Low discrimination**						**03735102**
					GRA		
Bifidobacterium adolescentis		50.16	0.51	MNS 87	99		
Actinomyces israelii		43.91	0.57	ARA 87	99		
Fusobacterium mortiferum		2.20	0.32		1		
03735103	**Low discrimination**						**03735103**
Actinomyces naeslundii		71.05	0.44	MNS 88			
Bifidobacterium adolescentis		15.44	0.17	XYL 1			
Actinomyces israelii		13.52	0.23				
03735110	**Low discrimination**						**03735110**
					GRA		
Actinomyces israelii		84.23	0.40	CEL 93	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		11.93	0.19	MLZ 13	99	
Fusobacterium mortiferum		3.84	0.14		1	
03735112	**Low discrimination**					**03735112**
Bifidobacterium adolescentis		89.26	0.53	MNS 87		
Actinomyces israelii		9.55	0.43	ARA 87		
Clostridium clostridioforme		0.61	0.13			
03735121	**Good identification**					**03735121**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03735122	**Good identification**					**03735122**
					GRA	
Parabacteroides distasonis		92.23	0.57	MNS 98	1	
Actinomyces israelii		3.55	0.36		99	
Actinomyces naeslundii		1.97	0.41		99	
03735123	**Low discrimination**					**03735123**
					GRA	
Actinomyces naeslundii		76.03	0.44	MNS 88	99	
Parabacteroides distasonis		23.97	0.22	XYL 1	1	
03735132	**Good identification**					**03735132**
					GRA	
Parabacteroides distasonis		96.02	0.52	MNS 98	1	
Bifidobacterium adolescentis		1.52	0.19		99	
Actinomyces israelii		1.03	0.22		99	
03735140	**Good identification**					**03735140**
					GRA	
Actinomyces israelii		96.00	0.47	CEL 93	99	
Actinomyces naeslundii		2.31	0.28		99	
Fusobacterium mortiferum		1.68	0.14		1	
03735142	**Low discrimination**					**03735142**
Actinomyces israelii		72.51	0.50	CEL 93		
Bifidobacterium adolescentis		26.32	0.36			
Actinomyces naeslundii		1.17	0.28			
03735152	**Low discrimination**					**03735152**
Bifidobacterium adolescentis		74.81	0.37	MNS 87		
Actinomyces israelii		25.19	0.36	ARA 87		
03735302	**Low discrimination**					**03735302**
					GRA	
Actinomyces israelii		42.45	0.37	CEL 93	99	
Parabacteroides distasonis		28.33	0.31	RHA 7	1	
Eubacterium contortum		14.05	0.14		99	
03735322	**Very good identification**					**03735322**
Parabacteroides distasonis		99.40	0.65	MNS 98		
Clostridium clostridioforme		0.31	0.16			
Prevotella oralis		0.11	0.01			
03735332	**Very good identification**					**03735332**
Parabacteroides distasonis		97.45	0.60	MNS 98		
Clostridium clostridioforme		2.55	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03735400	**Low discrimination**					**03735400**
					SPO	
Actinomyces israelii		88.65	0.42	CEL 93	1	
Lactobacillus fermentum		9.16	0.24		1	
Clostridium butyricum		2.19	-.07		99	
03735402	**Low discrimination**					**03735402**
Actinomyces israelii		66.92	0.45	CEL 93		
Bifidobacterium adolescentis		25.26	0.31			
Lactobacillus fermentum		4.61	0.24			
03735442	**Low discrimination**					**03735442**
Actinomyces israelii		89.29	0.38	CEL 93		
Bifidobacterium adolescentis		10.71	0.16			
03735500	**Low discrimination**					**03735500**
					SPO	
Actinomyces israelii		69.41	0.52	CEL 93	1	
Clostridium butyricum		10.28	0.17		99	
Bifidobacterium adolescentis		9.83	0.31		1	
03735502	**Low discrimination**					**03735502**
Bifidobacterium adolescentis		88.73	0.66	MNS 87		
Actinomyces israelii		9.50	0.55			
Bifidobacterium longum subsp. longum		0.64	0.27			
03735512	**Very good identification**					**03735512**
Bifidobacterium adolescentis		96.18	0.67	MNS 87		
Clostridium clostridioforme		1.54	0.33			
Actinomyces israelii		1.26	0.41			
03735520	**Low discrimination**					**03735520**
					GRA	
Bifidobacterium longum subsp. longum		70.15	0.37	MNS 87	99	
Actinomyces israelii		11.12	0.32	SAL 1	99	
Bacteroides vulgatus		10.78	0.09		1	
03735522	**Low discrimination**					**03735522**
					GRA	
Parabacteroides distasonis		72.28	0.53	MNS 98	1	
Bifidobacterium longum subsp. longum		15.63	0.37		99	
Bifidobacterium adolescentis		5.49	0.31		99	
03735530	**Good identification**					**03735530**
					GRA	
Bifidobacterium longum subsp. longum		97.21	0.41	MNS 87	99	
Parabacteroides distasonis		1.44	0.14	SAL 1	1	
Actinomyces israelii		1.35	0.18		99	
03735532	**Low discrimination**					**03735532**
					GRA	
Parabacteroides distasonis		46.64	0.48	MNS 98	1	
Bifidobacterium longum subsp. longum		31.90	0.41		99	
Clostridium clostridioforme		12.75	0.33		50	
03735540	**Acceptable identification**					**03735540**
					SPO	
Actinomyces israelii		92.76	0.45	CEL 93	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		4.17	0.16		1	
Clostridium butyricum		3.07	-.01		99	
03735542	**Low discrimination**					**03735542**
Bifidobacterium adolescentis		74.81	0.50	MNS 87		
Actinomyces israelii		25.19	0.48	SOR 11		
03735552	**Good identification**					**03735552**
Bifidobacterium adolescentis		96.05	0.51	MNS 87		
Actinomyces israelii		3.95	0.34	SOR 11		
03735612	**Good identification**					**03735612**
Clostridium clostridioforme		100.0	0.39	MNS 99		
03735632	**Good identification**					**03735632**
Clostridium clostridioforme		95.18	0.39	MNS 99		
Parabacteroides distasonis		4.82	0.22			
03735702	**Low discrimination**					**03735702**
Clostridium clostridioforme		55.89	0.37	MNS 99		
Bifidobacterium adolescentis		17.93	0.31			
Actinomyces israelii		14.30	0.36			
03735712	**Acceptable identification**					**03735712**
					GRA	
Clostridium clostridioforme		90.13	0.48	MNS 99	50	
Bifidobacterium adolescentis		7.76	0.33		99	
Parabacteroides distasonis		1.36	0.22		1	
03735722	**Good identification**					**03735722**
					GRA	
Parabacteroides distasonis		92.28	0.61	MNS 98	1	
Clostridium clostridioforme		7.29	0.37		50	
Actinomyces israelii		0.12	0.15		99	
03735732	**Low discrimination**					**03735732**
					GRA	
Parabacteroides distasonis		59.86	0.57	MNS 98	1	
Clostridium clostridioforme		40.00	0.48		50	
Bifidobacterium longum subsp. longum		0.14	0.07		99	
03736102	**Low discrimination**					**03736102**
Actinomyces israelii		78.30	0.35	CEL 93		
Bifidobacterium adolescentis		12.92	0.16	GAL 99		
Propionibacterium propionicum		4.87	0.14			
03737000	**Very good identification**					**03737000**
Actinomyces israelii		97.51	0.56	CEL 93		
Actinomyces naeslundii		2.49	0.38			
03737001	**Low discrimination**					**03737001**
Actinomyces naeslundii		79.15	0.41	MNS 88		
Actinomyces israelii		20.85	0.22	XYL 1		
03737002	**Very good identification**					**03737002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		97.68	0.59	CEL 93		
Actinomyces naeslundii		1.67	0.38			
Bifidobacterium adolescentis		0.65	0.16			
03737003	**Low discrimination**					**03737003**
Actinomyces naeslundii		71.68	0.41	MNS 88		
Actinomyces israelii		28.32	0.25	XYL 1		
03737010	**Good identification**					**03737010**
Actinomyces israelii		100.0	0.42	CEL 93		
03737012	**Acceptable identification**					**03737012**
Actinomyces israelii		94.83	0.45	CEL 93		
Bifidobacterium adolescentis		5.17	0.17	MLZ 13		
03737020	**Low discrimination**					**03737020**
Actinomyces israelii		71.40	0.36	CEL 93		
Actinomyces naeslundii		28.60	0.38	NAG 6		
03737021	**Good identification**					**03737021**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03737022	**Low discrimination**					**03737022**
Actinomyces israelii		78.93	0.39	CEL 93		
Actinomyces naeslundii		21.08	0.38	NAG 6		
03737023	**Good identification**					**03737023**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03737040	**Good identification**					**03737040**
Actinomyces israelii		98.85	0.49	CEL 93		
Actinomyces naeslundii		1.15	0.25			
03737042	**Good identification**					**03737042**
Actinomyces israelii		99.23	0.52	CEL 93		
Actinomyces naeslundii		0.77	0.25			
03737050	**Good identification**					**03737050**
Actinomyces israelii		100.0	0.35	CEL 93		
03737052	**Good identification**					**03737052**
Actinomyces israelii		100.0	0.38	CEL 93		
03737100	**Very good identification**					**03737100**
Actinomyces israelii		97.86	0.67	CEL 93		
Actinomyces naeslundii		1.78	0.46			
Bifidobacterium adolescentis		0.25	0.16			
03737101	**Low discrimination**					**03737101**
Actinomyces naeslundii		72.98	0.49	MNS 88		
Actinomyces israelii		27.02	0.32	XYL 1		
03737102	**Low discrimination**					**03737102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		84.91	0.70	CEL 93		
Bifidobacterium adolescentis		14.01	0.50			
Actinomyces naeslundii		1.03	0.46			
03737103	**Low discrimination**					**03737103**
Actinomyces naeslundii		60.72	0.49	MNS 88		
Actinomyces israelii		33.72	0.35	XYL 1		
Bifidobacterium adolescentis		5.56	0.16			
03737110	**Good identification**					**03737110**
Actinomyces israelii		97.39	0.52	CEL 93		
Bifidobacterium adolescentis		1.99	0.17	MLZ 13		
Actinomyces naeslundii		0.62	0.24			
03737112	**Low discrimination**					**03737112**
Bifidobacterium adolescentis		57.33	0.51	MNS 87		
Actinomyces israelii		42.49	0.55	ARA 87		
Actinomyces naeslundii		0.18	0.24			
03737120	**Low discrimination**					**03737120**
Actinomyces israelii		77.82	0.46	CEL 93		
Actinomyces naeslundii		22.18	0.46	NAG 6		
03737121	**Good identification**					**03737121**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03737122	**Low discrimination**					**03737122**
					GRA	
Actinomyces israelii		79.22	0.49	CEL 93	99	
Actinomyces naeslundii		15.05	0.46	NAG 6	99	
Parabacteroides distasonis		3.67	0.22		1	
03737123	**Good identification**					**03737123**
Actinomyces naeslundii		96.58	0.49	MNS 88		
Actinomyces israelii		3.42	0.15	XYL 1		
03737140	**Very good identification**					**03737140**
Actinomyces israelii		99.18	0.60	CEL 93		
Actinomyces naeslundii		0.82	0.33			
03737141	**Low discrimination**					**03737141**
Actinomyces naeslundii		55.07	0.36	MNS 88		
Actinomyces israelii		44.93	0.25	XYL 1		
03737142	**Good identification**					**03737142**
Actinomyces israelii		94.38	0.63	CEL 93		
Bifidobacterium adolescentis		4.95	0.34			
Actinomyces naeslundii		0.52	0.33			
03737150	**Good identification**					**03737150**
Actinomyces israelii		100.0	0.45	CEL 93		
03737152	**Low discrimination**					**03737152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		69.99	0.48	CEL 93		
Bifidobacterium adolescentis		30.01	0.36	MLZ 13		
03737160	**Low discrimination**					**03737160**
Actinomyces israelii		88.55	0.39	CEL 93		
Actinomyces naeslundii		11.45	0.33	NAG 6		
03737161	**Good identification**					**03737161**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03737162	**Acceptable identification**					**03737162**
Actinomyces israelii		92.06	0.42	CEL 93		
Actinomyces naeslundii		7.94	0.33	NAG 6		
03737163	**Good identification**					**03737163**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03737200	**Good identification**					**03737200**
Actinomyces israelii		100.0	0.37	CEL 93		
03737202	**Good identification**					**03737202**
Actinomyces israelii		100.0	0.40	CEL 93		
03737300	**Good identification**					**03737300**
Actinomyces israelii		100.0	0.47	CEL 93		
03737302	**Good identification**					**03737302**
Actinomyces israelii		97.83	0.50	CEL 93		
Bifidobacterium adolescentis		2.17	0.16	RHA 7		
03737312	**Low discrimination**					**03737312**
Actinomyces israelii		84.67	0.36	CEL 93		
Bifidobacterium adolescentis		15.33	0.17	RHA 7		
03737340	**Good identification**					**03737340**
Actinomyces israelii		100.0	0.40	CEL 93		
03737342	**Good identification**					**03737342**
Actinomyces israelii		100.0	0.43	CEL 93		
03737400	**Good identification**					**03737400**
Actinomyces israelii		100.0	0.55	CEL 93		
03737402	**Good identification**					**03737402**
Actinomyces israelii		94.83	0.58	CEL 93		
Bifidobacterium adolescentis		5.17	0.30			
03737410	**Good identification**					**03737410**
Actinomyces israelii		100.0	0.41	CEL 93		
03737412	**Low discrimination**					**03737412**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		69.16	0.44	CEL 93		
Bifidobacterium adolescentis		30.84	0.31	MLZ 13		
03737422	**Good identification**					**03737422**
Actinomyces israelii		100.0	0.37	CEL 93		
03737440	**Good identification**					**03737440**
Actinomyces israelii		100.0	0.48	CEL 93		
03737442	**Good identification**					**03737442**
Actinomyces israelii		98.30	0.51	CEL 93		
Bifidobacterium adolescentis		1.70	0.14			
03737452	**Low discrimination**					**03737452**
Actinomyces israelii		87.59	0.37	CEL 93		
Bifidobacterium adolescentis		12.41	0.16	MLZ 13		
03737500	**Very good identification**					**03737500**
					SPO	
Actinomyces israelii		97.72	0.65	CEL 93	1	
Bifidobacterium adolescentis		2.00	0.30		1	
Clostridium butyricum		0.28	0.01		99	
03737502	**Low discrimination**					**03737502**
					SPO	
Bifidobacterium adolescentis		57.41	0.64	MNS 87	1	
Actinomyces israelii		42.56	0.68		1	
Clostridium butyricum		0.03	-.07		99	
03737510	**Low discrimination**					**03737510**
Actinomyces israelii		85.68	0.51	CEL 93		
Bifidobacterium adolescentis		14.32	0.31	MLZ 13		
03737512	**Good identification**					**03737512**
Bifidobacterium adolescentis		91.63	0.66	MNS 87		
Actinomyces israelii		8.30	0.54			
Bifidobacterium dentium		0.06	-.07			
03737520	**Good identification**					**03737520**
Actinomyces israelii		100.0	0.45	CEL 93		
03737522	**Low discrimination**					**03737522**
					GRA	
Actinomyces israelii		80.12	0.48	CEL 93	99	
Bifidobacterium adolescentis		17.10	0.30	NAG 6	99	
Parabacteroides distasonis		2.78	0.19		1	
03737540	**Very good identification**					**03737540**
Actinomyces israelii		99.36	0.58	CEL 93		
Bifidobacterium adolescentis		0.65	0.14			
03737542	**Low discrimination**					**03737542**
Actinomyces israelii		69.99	0.61	CEL 93		
Bifidobacterium adolescentis		30.01	0.48			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03737550	**Acceptable identification**					**03737550**
					COC	
Actinomyces israelii		92.44	0.44	CEL 93	1	
Bifidobacterium adolescentis		4.91	0.16	MLZ 13	1	
Blautia producta		2.65	-.03		99	
03737552	**Low discrimination**					**03737552**
Bifidobacterium adolescentis		77.81	0.50	MNS 87		
Actinomyces israelii		22.19	0.47	SOR 11		
03737560	**Good identification**					**03737560**
Actinomyces israelii		100.0	0.38	CEL 93		
03737562	**Acceptable identification**					**03737562**
Actinomyces israelii		93.65	0.41	CEL 93		
Bifidobacterium adolescentis		6.35	0.14	NAG 6		
03737600	**Good identification**					**03737600**
Actinomyces israelii		100.0	0.36	CEL 93		
03737602	**Good identification**					**03737602**
Actinomyces israelii		100.0	0.39	CEL 93		
03737700	**Good identification**					**03737700**
Actinomyces israelii		100.0	0.46	CEL 93		
03737702	**Low discrimination**					**03737702**
Actinomyces israelii		84.67	0.49	CEL 93		
Bifidobacterium adolescentis		15.33	0.30	RHA 7		
03737740	**Good identification**					**03737740**
Actinomyces israelii		100.0	0.39	CEL 93		
03737742	**Acceptable identification**					**03737742**
Actinomyces israelii		94.56	0.42	CEL 93		
Bifidobacterium adolescentis		5.44	0.14	RHA 7		
03741000	**Low discrimination**					**03741000**
					GRA	
Actinomyces naeslundii		49.20	0.43	MNS 88	99	
Fusobacterium mortiferum		32.10	0.28	FRU 95	1	
Eubacterium tortuosum		18.69	0.07		99	
03741001	**Good identification**					**03741001**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03741002	**Low discrimination**					**03741002**
					GRA	
Actinomyces naeslundii		87.76	0.43	MNS 88	99	
Fusobacterium mortiferum		6.36	0.12	FRU 95	1	
Eubacterium tortuosum		5.88	-.06		99	
03741003	**Good identification**					**03741003**
Actinomyces naeslundii		100.0	0.46	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03741020	**Acceptable identification**						**03741020**
Actinomyces naeslundii		93.72	0.43	MNS 88			
Eubacterium tortuosum		6.28	-.06	FRU 95			
03741021	**Good identification**						**03741021**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03741022	**Acceptable identification**						**03741022**
					GRA		
Actinomyces naeslundii		92.45	0.43	MNS 88	99		
Prevotella oralis		7.55	-.02	FRU 95	1		
03741023	**Good identification**						**03741023**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03741100	**Low discrimination**						**03741100**
					COC	GRA	
Actinomyces naeslundii		63.51	0.50	MNS 88	1	99	
Fusobacterium mortiferum		33.97	0.34	FRU 95	1	1	
Blautia hansenii		2.52	-.03		99	99	
03741101	**Good identification**						**03741101**
Actinomyces naeslundii		100.0	0.53	MNS 88			
03741102	**Low discrimination**						**03741102**
					GRA		
Actinomyces naeslundii		78.50	0.50	MNS 88	99		
Lactobacillus acidophilus		9.62	0.10	FRU 95	99		
Fusobacterium mortiferum		4.67	0.18		1		
03741103	**Good identification**						**03741103**
Actinomyces naeslundii		100.0	0.53	MNS 88			
03741120	**Low discrimination**						**03741120**
					GRA		
Actinomyces naeslundii		63.38	0.50	MNS 88	99		
Prevotella melaninogenica		35.12	0.17	FRU 95	1		
Prevotella oralis		1.50	-.03		1		
03741121	**Good identification**						**03741121**
Actinomyces naeslundii		100.0	0.53	MNS 88			
03741122	**Low discrimination**						**03741122**
					GRA		
Actinomyces naeslundii		59.62	0.50	MNS 88	99		
Prevotella oralis		26.79	0.19	FRU 95	1		
Lactobacillus acidophilus		7.30	0.10		99		
03741123	**Good identification**						**03741123**
Actinomyces naeslundii		100.0	0.53	MNS 88			
03741140	**Good identification**						**03741140**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03741141	**Good identification**						**03741141**
Actinomyces naeslundii		100.0	0.41	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03741142	**Good identification**					**03741142**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03741143	**Good identification**					**03741143**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03741160	**Good identification**					**03741160**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03741161	**Good identification**					**03741161**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03741162	**Good identification**					**03741162**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03741163	**Good identification**					**03741163**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03743000	**Good identification**					**03743000**
Actinomyces naeslundii		96.70	0.48	MNS 88		
Actinomyces israelii		3.30	0.13	FRU 95		
03743001	**Good identification**					**03743001**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03743002	**Good identification**					**03743002**
Actinomyces naeslundii		95.13	0.48	MNS 88		
Actinomyces israelii		4.87	0.16	FRU 95		
03743003	**Good identification**					**03743003**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03743020	**Good identification**					**03743020**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03743021	**Good identification**					**03743021**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03743022	**Good identification**					**03743022**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03743023	**Good identification**					**03743023**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03743041	**Good identification**					**03743041**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03743043	**Good identification**					**03743043**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03743061	**Good identification**					**03743061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
03743063	**Good identification**					**03743063**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03743100	**Good identification**					**03743100**
Actinomyces naeslundii		95.43	0.55	MNS 88		
Actinomyces israelii		4.57	0.24	FRU 95		
03743101	**Very good identification**					**03743101**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03743102	**Acceptable identification**					**03743102**
Actinomyces naeslundii		91.72	0.55	MNS 88		
Actinomyces israelii		6.59	0.27	FRU 95		
Bifidobacterium adolescentis		1.69	0.10			
03743103	**Very good identification**					**03743103**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03743111	**Good identification**					**03743111**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03743113	**Good identification**					**03743113**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03743120	**Good identification**					**03743120**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03743121	**Very good identification**					**03743121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03743122	**Good identification**					**03743122**
					SPO	
Actinomyces naeslundii		96.88	0.55	MNS 88	1	
Clostridium ramosum		3.12	0.05	FRU 95	92	
03743123	**Very good identification**					**03743123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03743131	**Good identification**					**03743131**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03743133	**Good identification**					**03743133**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03743140	**Acceptable identification**					**03743140**
Actinomyces naeslundii		90.45	0.43	MNS 88		
Actinomyces israelii		9.55	0.17	FRU 95		
03743141	**Good identification**					**03743141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MNS 88		
03743142	**Low discrimination**					**03743142**
Actinomyces naeslundii		86.33	0.43	MNS 88		
Actinomyces israelii		13.68	0.20	FRU 95		
03743143	**Good identification**					**03743143**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03743160	**Good identification**					**03743160**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03743161	**Good identification**					**03743161**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03743162	**Good identification**					**03743162**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03743163	**Good identification**					**03743163**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03745502	**Low discrimination**					**03745502**
Bifidobacterium adolescentis		89.40	0.40	MNS 87		
Actinomyces israelii		10.60	0.30	FRU 99		
03745512	**Good identification**					**03745512**
Bifidobacterium adolescentis		98.57	0.41	MNS 87		
Actinomyces israelii		1.43	0.16	FRU 99		
03747100	**Good identification**					**03747100**
Actinomyces israelii		100.0	0.42	FRU 99		
03747102	**Low discrimination**					**03747102**
Actinomyces israelii		87.04	0.45	FRU 99		
Bifidobacterium adolescentis		12.96	0.24			
03747142	**Good identification**					**03747142**
Actinomyces israelii		95.48	0.38	FRU 99		
Bifidobacterium adolescentis		4.52	0.09			
03747500	**Good identification**					**03747500**
Actinomyces israelii		100.0	0.40	FRU 99		
03747502	**Low discrimination**					**03747502**
Bifidobacterium adolescentis		54.91	0.38	MNS 87		
Actinomyces israelii		45.09	0.43	FRU 99		
03747512	**Acceptable identification**					**03747512**
Bifidobacterium adolescentis		90.88	0.40	MNS 87		
Actinomyces israelii		9.13	0.29	FRU 99		
03747542	**Low discrimination**					**03747542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		72.10	0.36	FRU 99		
Bifidobacterium adolescentis		27.90	0.23			
03750000	**Very good identification**					**03750000**
					GRA	
Fusobacterium mortiferum		97.99	0.59	MLT 11	1	
Actinomyces naeslundii		1.80	0.41		99	
Prevotella intermedia		0.21	0.02		1	
03750001	**Low discrimination**					**03750001**
					GRA	
Actinomyces naeslundii		73.19	0.44	MNS 88	99	
Fusobacterium mortiferum		26.81	0.25	GAL 96	1	
03750002	**Low discrimination**					**03750002**
					GRA	
Fusobacterium mortiferum		73.13	0.43	MLT 11	1	
Actinomyces viscosus		14.77	0.18	BGL 10	99	
Actinomyces naeslundii		12.10	0.41		99	
03750003	**Low discrimination**					**03750003**
Actinomyces naeslundii		78.66	0.44	MNS 88		
Actinomyces viscosus		21.34	0.10	GAL 96		
03750020	**Low discrimination**					**03750020**
					GRA	
Actinomyces naeslundii		64.54	0.41	MNS 88	99	
Fusobacterium mortiferum		35.46	0.25	GAL 96	1	
03750021	**Good identification**					**03750021**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03750022	**Acceptable identification**					**03750022**
					GRA	
Actinomyces naeslundii		90.65	0.41	MNS 88	99	
Prevotella oralis		9.35	-.02	GAL 96	1	
03750023	**Good identification**					**03750023**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03750100	**Very good identification**					**03750100**
					GRA	
Fusobacterium mortiferum		97.81	0.65	MLT 11	1	
Actinomyces naeslundii		2.19	0.49		99	
03750101	**Low discrimination**					**03750101**
					GRA	
Actinomyces naeslundii		76.91	0.52	MNS 88	99	
Fusobacterium mortiferum		23.10	0.31	GAL 96	1	
03750102	**Low discrimination**					**03750102**
					GRA	
Fusobacterium mortiferum		70.90	0.49	MLT 11	1	
Actinomyces naeslundii		14.31	0.49	BGL 10	99	
Actinomyces viscosus		10.91	0.23		99	
03750103	**Low discrimination**					**03750103**
					GRA	
Actinomyces naeslundii		83.14	0.52	MNS 88	99	
Actinomyces viscosus		14.09	0.14	GAL 96	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		2.77	0.15		1	
03750120	**Low discrimination**					**03750120**
					GRA	
Actinomyces naeslundii		64.77	0.49	MNS 88	99	
Fusobacterium mortiferum		29.18	0.31	GAL 96	1	
Prevotella melaninogenica		4.12	-.01		1	
03750121	**Good identification**					**03750121**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03750122	**Low discrimination**					**03750122**
					GRA	
Actinomyces naeslundii		55.51	0.49	MNS 88	99	
Prevotella oralis		31.51	0.19	GAL 96	1	
Lactobacillus acidophilus		8.59	0.10		99	
03750123	**Good identification**					**03750123**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03750141	**Good identification**					**03750141**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03750142	**Low discrimination**					**03750142**
					GRA	
Actinomyces naeslundii		77.91	0.36	MNS 88	99	
Fusobacterium mortiferum		22.09	0.15	GAL 96	1	
03750143	**Good identification**					**03750143**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03750160	**Good identification**					**03750160**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03750161	**Good identification**					**03750161**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03750162	**Good identification**					**03750162**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03750163	**Good identification**					**03750163**
Actinomyces naeslundii		100.0	0.39	MNS 88		
03751000	**Low discrimination**					**03751000**
					GRA	
Fusobacterium mortiferum		72.16	0.62	MLT 11	1	
Actinomyces naeslundii		21.22	0.65		99	
Lactobacillus catenaformis		3.08	0.29		99	
03751001	**Very good identification**					**03751001**
					GRA	
Actinomyces naeslundii		97.76	0.68	MNS 88	99	
Fusobacterium mortiferum		2.24	0.28		1	
03751002	**Low discrimination**					**03751002**
					GRA	
Actinomyces naeslundii		52.18	0.65	MNS 88	99	
Fusobacterium mortiferum		19.71	0.46		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		8.24	0.26		99	
03751003	**Very good identification**					**03751003**
					GRA	
Actinomyces naeslundii		99.06	0.68	MNS 88	99	
Actinomyces viscosus		0.69	0.06		99	
Fusobacterium mortiferum		0.25	0.12		1	
03751010	**Low discrimination**					**03751010**
					GRA	
Actinomyces naeslundii		43.74	0.43	MNS 88	99	
Fusobacterium mortiferum		28.54	0.28	MLZ 5	1	
Lactobacillus jensenii		20.16	0.15		99	
03751011	**Good identification**					**03751011**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03751012	**Low discrimination**					**03751012**
Actinomyces naeslundii		55.85	0.43	MNS 88		
Lactobacillus jensenii		17.16	0.12	MLZ 5		
Actinomyces israelii		14.48	0.23			
03751013	**Good identification**					**03751013**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03751020	**Low discrimination**					**03751020**
					GRA	
Actinomyces naeslundii		87.50	0.65	MNS 88	99	
Lactobacillus catenaformis		3.17	0.18		99	
Fusobacterium mortiferum		3.01	0.28		1	
03751021	**Very good identification**					**03751021**
Actinomyces naeslundii		100.0	0.68	MNS 88		
03751022	**Low discrimination**					**03751022**
					GRA	
Actinomyces naeslundii		59.78	0.65	MNS 88	99	
Prevotella oralis		25.43	0.33		1	
Lactobacillus acidophilus		9.44	0.26		99	
03751023	**Very good identification**					**03751023**
					GRA	
Actinomyces naeslundii		99.71	0.68	MNS 88	99	
Prevotella oralis		0.29	-.02		1	
03751030	**Low discrimination**					**03751030**
Actinomyces naeslundii		89.67	0.43	MNS 88		
Lactobacillus jensenii		10.33	0.05	MLZ 5		
03751031	**Good identification**					**03751031**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03751032	**Low discrimination**					**03751032**
					GRA	
Actinomyces naeslundii		86.32	0.43	MNS 88	99	
Prevotella oralis		7.05	-.02	MLZ 5	1	
Lactobacillus jensenii		6.63	0.02		99	
03751033	**Good identification**					**03751033**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MNS 88		
03751040	**Low discrimination**					**03751040**
					GRA	
Actinomyces naeslundii		75.26	0.52	MNS 88	99	
Fusobacterium mortiferum		14.65	0.28		1	
Actinomyces israelii		10.10	0.28		99	
03751041	**Good identification**					**03751041**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03751042	**Low discrimination**					**03751042**
Actinomyces naeslundii		74.99	0.52	MNS 88		
Actinomyces israelii		15.09	0.31			
Lactobacillus acidophilus		8.30	0.11			
03751043	**Good identification**					**03751043**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03751060	**Good identification**					**03751060**
Actinomyces naeslundii		100.0	0.52	MNS 88		
03751061	**Good identification**					**03751061**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03751062	**Low discrimination**					**03751062**
					GRA	
Actinomyces naeslundii		88.11	0.52	MNS 88	99	
Lactobacillus acidophilus		9.75	0.11		99	
Prevotella oralis		2.15	-.02		1	
03751063	**Good identification**					**03751063**
Actinomyces naeslundii		100.0	0.55	MNS 88		
03751100	**Low discrimination**					**03751100**
					GRA	
Fusobacterium mortiferum		69.80	0.68	MLT 11	1	
Actinomyces naeslundii		25.04	0.72		99	
Actinomyces israelii		2.14	0.45		99	
03751101	**Very good identification**					**03751101**
					GRA	
Actinomyces naeslundii		98.16	0.75	MNS 88	99	
Fusobacterium mortiferum		1.84	0.34		1	
03751102	**Low discrimination**					**03751102**
					GRA	
Actinomyces naeslundii		38.99	0.72	MNS 88	99	
Lactobacillus acidophilus		24.89	0.45		99	
Fusobacterium mortiferum		12.08	0.52		1	
03751103	**Very good identification**					**03751103**
Actinomyces naeslundii		98.63	0.75	MNS 88		
Actinomyces viscosus		0.43	0.11			
Lactobacillus acidophilus		0.42	0.10			
03751110	**Low discrimination**					**03751110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		53.68	0.50	MNS 88	99	
Fusobacterium mortiferum		28.72	0.34	MLZ 5	1	
Actinomyces israelii		13.04	0.31		99	
03751111	**Good identification**					**03751111**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03751112	**Low discrimination**					**03751112**
Bifidobacterium adolescentis		73.36	0.47	MNS 87		
Actinomyces naeslundii		13.90	0.50	XYL 87		
Actinomyces israelii		5.06	0.34	ARA 87		
03751113	**Good identification**					**03751113**
Actinomyces naeslundii		96.57	0.53	MNS 88		
Bifidobacterium adolescentis		3.43	0.13	MLZ 5		
03751120	**Low discrimination**					**03751120**
					GRA	
Actinomyces naeslundii		66.33	0.72	MNS 88	99	
Prevotella melaninogenica		17.41	0.33		1	
Prevotella oralis		8.17	0.31		1	
03751121	**Very good identification**					**03751121**
					GRA	
Actinomyces naeslundii		99.74	0.75	MNS 88	99	
Prevotella melaninogenica		0.18	-.01		1	
Prevotella oralis		0.08	-.03		1	
03751122	**Low discrimination**					**03751122**
					GRA	
Prevotella oralis		55.67	0.53	MNS 99	1	
Actinomyces naeslundii		23.78	0.72		99	
Lactobacillus acidophilus		15.18	0.45		99	
03751123	**Very good identification**					**03751123**
					GRA	
Actinomyces naeslundii		98.03	0.75	MNS 88	99	
Prevotella oralis		1.55	0.19		1	
Lactobacillus acidophilus		0.42	0.10		99	
03751130	**Low discrimination**					**03751130**
					GRA	
Actinomyces naeslundii		88.47	0.50	MNS 88	99	
Bifidobacterium breve		4.98	0.15	MLZ 5	99	
Prevotella melaninogenica		4.46	-.01		1	
03751131	**Good identification**					**03751131**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03751132	**Low discrimination**					**03751132**
					GRA	
Actinomyces naeslundii		43.28	0.50	MNS 88	99	
Bifidobacterium breve		24.62	0.32	MLZ 5	99	
Prevotella oralis		19.45	0.19		1	
03751133	**Good identification**					**03751133**
Actinomyces naeslundii		100.0	0.53	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03751140	**Low discrimination**					**03751140**
					GRA	
Actinomyces naeslundii		73.67	0.59	MNS 88	99	
Actinomyces israelii		13.89	0.38		99	
Fusobacterium mortiferum		11.75	0.34		1	
03751141	**Very good identification**					**03751141**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03751142	**Low discrimination**					**03751142**
Actinomyces naeslundii		49.77	0.59	MNS 88		
Lactobacillus acidophilus		22.25	0.29			
Actinomyces israelii		14.07	0.41			
03751143	**Very good identification**					**03751143**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03751150	**Low discrimination**					**03751150**
Actinomyces naeslundii		58.33	0.38	MNS 88		
Actinomyces israelii		31.22	0.24	MLZ 5		
Bifidobacterium breve		10.46	0.10			
03751151	**Good identification**					**03751151**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03751153	**Good identification**					**03751153**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03751160	**Good identification**					**03751160**
					GRA	
Actinomyces naeslundii		93.41	0.59	MNS 88	99	
Bifidobacterium breve		1.57	0.15		99	
Prevotella melaninogenica		1.40	-.01		1	
03751161	**Very good identification**					**03751161**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03751162	**Low discrimination**					**03751162**
Actinomyces naeslundii		56.18	0.59	MNS 88		
Lactobacillus acidophilus		25.11	0.29			
Bifidobacterium breve		9.53	0.32			
03751163	**Very good identification**					**03751163**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03751170	**Low discrimination**					**03751170**
Actinomyces naeslundii		75.83	0.38	MNS 88		
Bifidobacterium breve		24.17	0.15	MLZ 5		
03751171	**Good identification**					**03751171**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03751173	**Good identification**					**03751173**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	MNS 88			
03751301	**Good identification**						**03751301**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03751303	**Good identification**						**03751303**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03751321	**Good identification**						**03751321**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03751322	**Low discrimination**						**03751322**
					SPO	GRA	
Prevotella oralis		85.98	0.47	MNS 99	1	1	
Clostridium ramosum		6.53	0.28		92	50	
Lactobacillus acidophilus		4.12	0.25		1	99	
03751323	**Low discrimination**						**03751323**
					GRA		
Actinomyces naeslundii		59.92	0.41	MNS 88	99		
Prevotella oralis		40.08	0.13	RHA 1	1		
03751402	**Low discrimination**						**03751402**
Actinomyces israelii		46.43	0.36	XYL 92			
Bifidobacterium adolescentis		27.19	0.26				
Lactobacillus fermentum		17.74	0.28				
03751500	**Low discrimination**						**03751500**
					GRA		
Actinomyces israelii		47.75	0.43	XYL 92	99		
Fusobacterium mortiferum		19.21	0.34		1		
Lactobacillus fermentum		11.16	0.32		99		
03751501	**Good identification**						**03751501**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03751502	**Good identification**						**03751502**
Bifidobacterium adolescentis		91.31	0.60	MNS 87			
Actinomyces israelii		6.30	0.46	XYL 87			
Lactobacillus fermentum		0.98	0.32				
03751512	**Very good identification**						**03751512**
Bifidobacterium adolescentis		99.01	0.61	MNS 87			
Actinomyces israelii		0.84	0.32	XYL 87			
Clostridium clostridioforme		0.15	0.10				
03751520	**Low discrimination**						**03751520**
Actinomyces naeslundii		40.16	0.38	MNS 88			
Lactobacillus fermentum		21.68	0.24	ARA 1			
Actinomyces israelii		17.76	0.23				
03751521	**Good identification**						**03751521**
Actinomyces naeslundii		100.0	0.41	MNS 88			
03751523	**Good identification**						**03751523**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.41	MNS 88		
03751540	**Low discrimination**					**03751540**
	Actinomyces israelii	88.09	0.36	XYL 92		
	Bifidobacterium adolescentis	6.15	0.10			
	Actinomyces naeslundii	5.77	0.25			
03751542	**Low discrimination**					**03751542**
	Bifidobacterium adolescentis	81.53	0.44	MNS 87		
	Actinomyces israelii	17.70	0.39	XYL 87		
	Actinomyces naeslundii	0.77	0.25	SOR 11		
03751552	**Good identification**					**03751552**
	Bifidobacterium adolescentis	97.41	0.46	MNS 87		
	Actinomyces israelii	2.59	0.25	XYL 87		
03752000	**Low discrimination**					**03752000**
					GRA	
	Actinomyces naeslundii	75.41	0.46	MNS 88	99	
	Fusobacterium mortiferum	21.35	0.25	GAL 96	1	
	Actinomyces israelii	3.25	0.13		99	
03752001	**Good identification**					**03752001**
	Actinomyces naeslundii	100.0	0.49	MNS 88		
03752002	**Acceptable identification**					**03752002**
	Actinomyces naeslundii	93.93	0.46	MNS 88		
	Actinomyces israelii	6.07	0.16	GAL 96		
03752003	**Good identification**					**03752003**
	Actinomyces naeslundii	100.0	0.49	MNS 88		
03752020	**Good identification**					**03752020**
	Actinomyces naeslundii	100.0	0.46	MNS 88		
03752021	**Good identification**					**03752021**
	Actinomyces naeslundii	100.0	0.49	MNS 88		
03752022	**Good identification**					**03752022**
	Actinomyces naeslundii	100.0	0.46	MNS 88		
03752023	**Good identification**					**03752023**
	Actinomyces naeslundii	100.0	0.49	MNS 88		
03752041	**Good identification**					**03752041**
	Actinomyces naeslundii	100.0	0.36	MNS 88		
03752043	**Good identification**					**03752043**
	Actinomyces naeslundii	100.0	0.36	MNS 88		
03752061	**Good identification**					**03752061**
	Actinomyces naeslundii	100.0	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03752063	**Good identification**					**03752063**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03752100	**Low discrimination**					**03752100**
					GRA	
Actinomyces naeslundii		76.48	0.54	MNS 88	99	
Fusobacterium mortiferum		17.75	0.31	GAL 96	1	
Actinomyces israelii		4.63	0.24		99	
03752101	**Very good identification**					**03752101**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03752102	**Low discrimination**					**03752102**
					GRA	
Actinomyces naeslundii		86.04	0.54	MNS 88	99	
Actinomyces israelii		7.81	0.27	GAL 96	99	
Fusobacterium mortiferum		2.22	0.15		1	
03752103	**Very good identification**					**03752103**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03752120	**Good identification**					**03752120**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03752121	**Very good identification**					**03752121**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03752122	**Good identification**					**03752122**
					SPO	
Actinomyces naeslundii		96.09	0.54	MNS 88	1	
Clostridium ramosum		3.91	0.05	GAL 96	92	
03752123	**Very good identification**					**03752123**
Actinomyces naeslundii		100.0	0.57	MNS 88		
03752140	**Low discrimination**					**03752140**
Actinomyces naeslundii		82.08	0.41	MNS 88		
Actinomyces israelii		10.95	0.17	GAL 96		
Propionibacterium propionicum		6.97	0.12			
03752141	**Good identification**					**03752141**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03752142	**Low discrimination**					**03752142**
Actinomyces naeslundii		75.33	0.41	MNS 88		
Actinomyces israelii		15.07	0.20	GAL 96		
Propionibacterium propionicum		9.60	0.15			
03752143	**Good identification**					**03752143**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03752160	**Good identification**					**03752160**
Actinomyces naeslundii		100.0	0.41	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03752161	**Good identification**					**03752161**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03752162	**Good identification**					**03752162**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03752163	**Good identification**					**03752163**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03753000	**Low discrimination**					**03753000**
					GRA	
Actinomyces naeslundii		83.66	0.70	MNS 88	99	
Actinomyces israelii		14.87	0.47		99	
Fusobacterium mortiferum		1.48	0.28		1	
03753001	**Very good identification**					**03753001**
Actinomyces naeslundii		99.88	0.73	MNS 88		
Actinomyces israelii		0.12	0.13			
03753002	**Low discrimination**					**03753002**
Actinomyces naeslundii		78.40	0.70	MNS 88		
Actinomyces israelii		20.90	0.50			
Lactobacillus acidophilus		0.34	0.05			
03753003	**Very good identification**					**03753003**
Actinomyces naeslundii		99.82	0.73	MNS 88		
Actinomyces israelii		0.18	0.16			
03753010	**Low discrimination**					**03753010**
Actinomyces naeslundii		66.47	0.48	MNS 88		
Actinomyces israelii		33.53	0.33	MLZ 5		
03753011	**Good identification**					**03753011**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03753012	**Low discrimination**					**03753012**
Actinomyces naeslundii		54.93	0.48	MNS 88		
Actinomyces israelii		41.56	0.36	MLZ 5		
Bifidobacterium adolescentis		3.51	0.12			
03753013	**Good identification**					**03753013**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03753020	**Very good identification**					**03753020**
Actinomyces naeslundii		98.88	0.70	MNS 88		
Actinomyces israelii		1.12	0.27			
03753021	**Very good identification**					**03753021**
Actinomyces naeslundii		100.0	0.73	MNS 88		
03753022	**Very good identification**					**03753022**
					SPO	
Actinomyces naeslundii		97.25	0.70	MNS 88	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		1.65	0.30		1	
Clostridium ramosum		0.47	0.05		92	
03753023	**Very good identification**					**03753023**
Actinomyces naeslundii		100.0	0.73	MNS 88		
03753030	**Good identification**					**03753030**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03753031	**Good identification**					**03753031**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03753032	**Good identification**					**03753032**
Actinomyces naeslundii		95.39	0.48	MNS 88		
Actinomyces israelii		4.61	0.16	MLZ 5		
03753033	**Good identification**					**03753033**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03753040	**Low discrimination**					**03753040**
Actinomyces naeslundii		71.86	0.57	MNS 88		
Actinomyces israelii		28.14	0.40			
03753041	**Very good identification**					**03753041**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03753042	**Low discrimination**					**03753042**
Actinomyces naeslundii		63.00	0.57	MNS 88		
Actinomyces israelii		37.00	0.43			
03753043	**Very good identification**					**03753043**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03753051	**Good identification**					**03753051**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03753053	**Good identification**					**03753053**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03753060	**Very good identification**					**03753060**
Actinomyces naeslundii		97.56	0.57	MNS 88		
Actinomyces israelii		2.44	0.20			
03753061	**Very good identification**					**03753061**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03753062	**Very good identification**					**03753062**
Actinomyces naeslundii		96.39	0.57	MNS 88		
Actinomyces israelii		3.61	0.23			
03753063	**Very good identification**					**03753063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.60	MNS 88		
03753071	**Good identification**					**03753071**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03753073	**Good identification**					**03753073**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03753100	**Low discrimination**					**03753100**
					GRA	
Actinomyces naeslundii		78.95	0.77	MNS 88	99	
Actinomyces israelii		19.72	0.58		99	
Fusobacterium mortiferum		1.15	0.34		1	
03753101	**Very good identification**					**03753101**
Actinomyces naeslundii		99.83	0.80	MNS 88		
Actinomyces israelii		0.17	0.24			
03753102	**Low discrimination**					**03753102**
Actinomyces naeslundii		66.61	0.77	MNS 88		
Actinomyces israelii		24.95	0.61			
Bifidobacterium adolescentis		6.38	0.44			
03753103	**Very good identification**					**03753103**
Actinomyces naeslundii		99.68	0.80	MNS 88		
Actinomyces israelii		0.25	0.27			
Bifidobacterium adolescentis		0.06	0.10			
03753110	**Low discrimination**					**03753110**
Actinomyces naeslundii		57.22	0.55	MNS 88		
Actinomyces israelii		40.57	0.44	MLZ 5		
Bifidobacterium adolescentis		1.29	0.12			
03753111	**Very good identification**					**03753111**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03753112	**Low discrimination**					**03753112**
Bifidobacterium adolescentis		49.96	0.46	MNS 87		
Actinomyces israelii		23.88	0.47	XYL 87		
Actinomyces naeslundii		22.46	0.55	ARA 87		
03753113	**Very good identification**					**03753113**
Actinomyces naeslundii		98.52	0.58	MNS 88		
Bifidobacterium adolescentis		1.48	0.12	MLZ 5		
03753120	**Good identification**					**03753120**
					SPO	
Actinomyces naeslundii		94.21	0.77	MNS 88	1	
Clostridium ramosum		3.95	0.28		92	
Actinomyces israelii		1.50	0.37		1	
03753121	**Very good identification**					**03753121**
Actinomyces naeslundii		100.0	0.80	MNS 88		
03753122	**Low discrimination**					**03753122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		80.69	0.77	MNS 88	1	
Clostridium ramosum		13.54	0.39		92	
Actinomyces israelii		1.93	0.40		1	
03753123	**Very good identification**					**03753123**
					SPO	
Actinomyces naeslundii		99.89	0.80	MNS 88	1	
Clostridium ramosum		0.11	0.05		92	
03753130	**Acceptable identification**					**03753130**
Actinomyces naeslundii		93.09	0.55	MNS 88		
Actinomyces israelii		4.21	0.23	MLZ 5		
Bifidobacterium breve		2.70	0.15			
03753131	**Very good identification**					**03753131**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03753132	**Low discrimination**					**03753132**
Actinomyces naeslundii		69.35	0.55	MNS 88		
Bifidobacterium breve		20.32	0.32	MLZ 5		
Actinomyces israelii		4.71	0.26			
03753133	**Very good identification**					**03753133**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03753140	**Low discrimination**					**03753140**
Actinomyces naeslundii		64.16	0.64	MNS 88		
Actinomyces israelii		35.31	0.51			
Bifidobacterium breve		0.31	0.10			
03753141	**Very good identification**					**03753141**
Actinomyces naeslundii		99.63	0.67	MNS 88		
Actinomyces israelii		0.37	0.17			
03753142	**Low discrimination**					**03753142**
Actinomyces naeslundii		51.04	0.64	MNS 88		
Actinomyces israelii		42.13	0.54			
Bifidobacterium adolescentis		3.43	0.29			
03753143	**Very good identification**					**03753143**
Actinomyces naeslundii		99.45	0.67	MNS 88		
Actinomyces israelii		0.55	0.20			
03753150	**Low discrimination**					**03753150**
Actinomyces israelii		58.85	0.37	XYL 92		
Actinomyces naeslundii		37.67	0.43	MLZ 13		
Bifidobacterium breve		3.48	0.10			
03753151	**Good identification**					**03753151**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03753152	**Low discrimination**					**03753152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		40.16	0.40	XYL 92		
Bifidobacterium adolescentis		26.70	0.30	MLZ 13		
Actinomyces naeslundii		17.14	0.43			
03753153	**Good identification**					**03753153**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03753160	**Very good identification**					**03753160**
Actinomyces naeslundii		95.81	0.64	MNS 88		
Actinomyces israelii		3.37	0.30			
Bifidobacterium breve		0.83	0.15			
03753161	**Very good identification**					**03753161**
Actinomyces naeslundii		100.0	0.67	MNS 88		
03753162	**Low discrimination**					**03753162**
Actinomyces naeslundii		86.07	0.64	MNS 88		
Bifidobacterium breve		7.52	0.32			
Actinomyces israelii		4.54	0.33			
03753163	**Very good identification**					**03753163**
Actinomyces naeslundii		100.0	0.67	MNS 88		
03753170	**Low discrimination**					**03753170**
Actinomyces naeslundii		79.12	0.43	MNS 88		
Bifidobacterium breve		12.99	0.15	MLZ 5		
Actinomyces israelii		7.89	0.16			
03753171	**Good identification**					**03753171**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03753173	**Good identification**					**03753173**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03753201	**Good identification**					**03753201**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03753203	**Good identification**					**03753203**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03753220	**Good identification**					**03753220**
Actinomyces naeslundii		100.0	0.35	MNS 88		
03753221	**Good identification**					**03753221**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03753222	**Low discrimination**					**03753222**
					SPO	
Actinomyces naeslundii		61.24	0.35	MNS 88	1	
Clostridium ramosum		38.76	0.07	RHA 1	92	
03753223	**Good identification**					**03753223**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03753300	**Low discrimination**						**03753300**
Actinomyces israelii		65.04	0.39	XYL 92			
Actinomyces naeslundii		34.96	0.43	RHA 7			
03753301	**Good identification**						**03753301**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03753302	**Low discrimination**						**03753302**
					SPO		
Actinomyces israelii		65.86	0.42	XYL 92	1		
Actinomyces naeslundii		23.59	0.43	RHA 7	1		
Clostridium ramosum		5.25	0.07		92		
03753303	**Good identification**						**03753303**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03753321	**Good identification**						**03753321**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03753322	**Acceptable identification**						**03753322**
					SPO	GRA	
Clostridium ramosum		92.35	0.41	MNS 99	92	50	
Actinomyces naeslundii		4.20	0.43	TRE 94	1	99	
Prevotella oralis		2.17	0.13		1	1	
03753323	**Low discrimination**						**03753323**
					SPO		
Actinomyces naeslundii		87.09	0.46	MNS 88	1		
Clostridium ramosum		12.91	0.07	RHA 1	92		
03753400	**Acceptable identification**						**03753400**
Actinomyces israelii		93.50	0.46	XYL 92			
Actinomyces naeslundii		6.50	0.35				
03753401	**Good identification**						**03753401**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03753402	**Low discrimination**						**03753402**
Actinomyces israelii		88.43	0.49	XYL 92			
Bifidobacterium adolescentis		7.48	0.24				
Actinomyces naeslundii		4.10	0.35				
03753403	**Low discrimination**						**03753403**
Actinomyces naeslundii		87.31	0.38	MNS 88			
Actinomyces israelii		12.69	0.15	ARA 1			
03753420	**Low discrimination**						**03753420**
Actinomyces naeslundii		52.12	0.35	MNS 88			
Actinomyces israelii		47.88	0.26	ARA 1			
03753421	**Good identification**						**03753421**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03753423	**Good identification**						**03753423**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.38	MNS 88		
03753440	**Good identification**					**03753440**
	Actinomyces israelii	96.94	0.39	XYL 92		
	Actinomyces naeslundii	3.06	0.22			
03753442	**Good identification**					**03753442**
	Actinomyces israelii	95.43	0.42	XYL 92		
	Bifidobacterium adolescentis	2.56	0.09			
	Actinomyces naeslundii	2.01	0.22			
03753500	**Good identification**					**03753500**
	Actinomyces israelii	92.50	0.56	XYL 92		
	Actinomyces naeslundii	4.57	0.43			
	Bifidobacterium adolescentis	2.93	0.24			
03753501	**Low discrimination**					**03753501**
	Actinomyces naeslundii	88.01	0.46	MNS 88		
	Actinomyces israelii	11.99	0.22	ARA 1		
03753502	**Low discrimination**					**03753502**
	Bifidobacterium adolescentis	66.95	0.58	MNS 87		
	Actinomyces israelii	32.00	0.59	XYL 87		
	Actinomyces naeslundii	1.06	0.43			
03753503	**Low discrimination**					**03753503**
	Actinomyces naeslundii	61.28	0.46	MNS 88		
	Bifidobacterium adolescentis	26.20	0.24	ARA 1		
	Actinomyces israelii	12.52	0.25			
03753510	**Low discrimination**					**03753510**
	Actinomyces israelii	79.41	0.42	XYL 92		
	Bifidobacterium adolescentis	20.59	0.26	MLZ 13		
03753512	**Good identification**					**03753512**
	Bifidobacterium adolescentis	94.48	0.60	MNS 87		
	Actinomyces israelii	5.52	0.45	XYL 87		
03753520	**Low discrimination**					**03753520**
	Actinomyces israelii	56.35	0.36	XYL 92		
	Actinomyces naeslundii	43.65	0.43	NAG 6		
03753521	**Good identification**					**03753521**
	Actinomyces naeslundii	100.0	0.46	MNS 88		
03753522	**Low discrimination**					**03753522**
	Actinomyces israelii	51.70	0.39	XYL 92		
	Actinomyces naeslundii	26.70	0.43	NAG 6		
	Bifidobacterium adolescentis	17.12	0.24			
03753523	**Good identification**					**03753523**
	Actinomyces naeslundii	100.0	0.46	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03753540	**Good identification**					**03753540**
Actinomyces israelii		96.85	0.49	XYL 92		
Actinomyces naeslundii		2.17	0.30			
Bifidobacterium adolescentis		0.98	0.09			
03753542	**Low discrimination**					**03753542**
Actinomyces israelii		59.53	0.52	XYL 92		
Bifidobacterium adolescentis		39.58	0.43			
Actinomyces naeslundii		0.89	0.30			
03753550	**Acceptable identification**					**03753550**
Actinomyces israelii		92.39	0.35	XYL 92		
Bifidobacterium adolescentis		7.61	0.10	MLZ 13		
03753552	**Low discrimination**					**03753552**
Bifidobacterium adolescentis		84.47	0.44	MNS 87		
Actinomyces israelii		15.53	0.38	XYL 87		
03753700	**Good identification**					**03753700**
Actinomyces israelii		100.0	0.37	XYL 92		
03753702	**Low discrimination**					**03753702**
Actinomyces israelii		78.08	0.40	XYL 92		
Bifidobacterium adolescentis		21.92	0.24	RHA 7		
03754000	**Good identification**					**03754000**
					GRA	
Fusobacterium mortiferum		96.02	0.40	MLT 11	1	
Actinomyces israelii		3.98	0.19	XYL 7	99	
03754100	**Acceptable identification**					**03754100**
					GRA	
Fusobacterium mortiferum		93.37	0.46	MLT 11	1	
Actinomyces israelii		6.63	0.29	XYL 7	99	
03754502	**Low discrimination**					**03754502**
Bifidobacterium adolescentis		87.81	0.40	MNS 87		
Actinomyces israelii		10.41	0.30	GAL 99		
Clostridium clostridioforme		1.78	0.08			
03754512	**Acceptable identification**					**03754512**
Bifidobacterium adolescentis		91.73	0.41	MNS 87		
Clostridium clostridioforme		6.94	0.19	GAL 99		
Actinomyces israelii		1.33	0.16			
03755000	**Low discrimination**					**03755000**
					GRA	
Actinomyces israelii		71.23	0.53		99	
Fusobacterium mortiferum		26.04	0.43		1	
Lactobacillus fermentum		1.71	0.24		99	
03755002	**Acceptable identification**					**03755002**
					GRA	
Actinomyces israelii		91.38	0.56		99	
Bifidobacterium adolescentis		3.81	0.26		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		2.48	0.27		1		
03755010	**Good identification**						**03755010**
Actinomyces israelii		100.0	0.39	MLZ 13			
03755012	**Low discrimination**						**03755012**
Actinomyces israelii		74.59	0.42	MLZ 13			
Bifidobacterium adolescentis		25.41	0.27				
03755022	**Low discrimination**						**03755022**
					GRA		
Actinomyces israelii		49.54	0.35	NAG 6	99		
Prevotella oralis		38.86	0.17		1		
Actinomyces naeslundii		7.47	0.30		99		
03755040	**Good identification**						**03755040**
Actinomyces israelii		100.0	0.46				
03755042	**Good identification**						**03755042**
Actinomyces israelii		98.69	0.49				
Bifidobacterium adolescentis		1.31	0.10				
03755100	**Low discrimination**						**03755100**
					GRA	SPO	
Actinomyces israelii		78.56	0.63		99	1	
Fusobacterium mortiferum		16.75	0.49		1	1	
Clostridium butyricum		1.89	0.14		99	99	
03755101	**Low discrimination**						**03755101**
					GRA		
Actinomyces naeslundii		55.69	0.41	MNS 88	99		
Actinomyces israelii		36.52	0.29	XYL 1	99		
Fusobacterium mortiferum		7.79	0.15		1		
03755102	**Low discrimination**						**03755102**
					GRA		
Bifidobacterium adolescentis		49.54	0.60	MNS 87	99		
Actinomyces israelii		48.05	0.66	ARA 87	99		
Fusobacterium mortiferum		0.76	0.33		1		
03755110	**Low discrimination**						**03755110**
					GRA		
Actinomyces israelii		86.91	0.49	MLZ 13	99		
Bifidobacterium adolescentis		11.10	0.27		99		
Fusobacterium mortiferum		1.25	0.15		1		
03755112	**Low discrimination**						**03755112**
Bifidobacterium adolescentis		89.32	0.61	MNS 87			
Actinomyces israelii		10.59	0.52	ARA 87			
Clostridium clostridioforme		0.09	0.07				
03755120	**Low discrimination**						**03755120**
					GRA		
Actinomyces israelii		63.11	0.43	NAG 6	99		
Prevotella oralis		15.31	0.16		1		
Actinomyces naeslundii		10.15	0.38		99		
03755121	**Good identification**						**03755121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	MNS 88			
03755122	**Low discrimination**						**03755122**
					GRA		
Prevotella oralis		57.39	0.37	MNS 99	1		
Actinomyces israelii		18.68	0.46	XYL 11	99		
Parabacteroides distasonis		16.24	0.41		1		
03755123	**Low discrimination**						**03755123**
					GRA		
Actinomyces naeslundii		83.83	0.41	MNS 88	99		
Prevotella oralis		16.17	0.03	XYL 1	1		
03755132	**Low discrimination**						**03755132**
					GRA		
Parabacteroides distasonis		53.97	0.36	MNS 98	1		
Bifidobacterium adolescentis		22.99	0.27		99		
Actinomyces israelii		17.23	0.31		99		
03755140	**Good identification**						**03755140**
					GRA	SPO	
Actinomyces israelii		97.99	0.56		99	1	
Fusobacterium mortiferum		0.54	0.15		1	1	
Clostridium butyricum		0.53	-.04		99	99	
03755142	**Low discrimination**						**03755142**
Actinomyces israelii		75.15	0.59				
Bifidobacterium adolescentis		24.62	0.44				
Actinomyces naeslundii		0.23	0.25				
03755150	**Good identification**						**03755150**
Actinomyces israelii		96.10	0.42	MLZ 13			
Bifidobacterium adolescentis		3.90	0.12				
03755152	**Low discrimination**						**03755152**
Bifidobacterium adolescentis		72.83	0.46	MNS 87			
Actinomyces israelii		27.17	0.45	ARA 87			
03755160	**Acceptable identification**						**03755160**
Actinomyces israelii		93.20	0.36	NAG 6			
Actinomyces naeslundii		6.80	0.25				
03755162	**Low discrimination**						**03755162**
					GRA		
Actinomyces israelii		84.72	0.39	NAG 6	99		
Prevotella oralis		6.76	0.03		1		
Bifidobacterium adolescentis		4.39	0.10		99		
03755202	**Good identification**						**03755202**
Actinomyces israelii		100.0	0.37	RHA 7			
03755300	**Low discrimination**						**03755300**
					GRA		
Actinomyces israelii		54.78	0.44	RHA 7	99		
Eubacterium contortum		43.65	0.27		99		
Fusobacterium mortiferum		1.57	0.15		1		
03755302	**Low discrimination**						**03755302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		74.65	0.47	RHA 7		
Bifidobacterium adolescentis		10.33	0.26			
Eubacterium contortum		7.00	0.14			
03755322	**Low discrimination**					**03755322**
					GRA	
Parabacteroides distasonis		64.38	0.49	MNS 98	1	
Prevotella oralis		32.51	0.31		1	
Actinomyces israelii		1.86	0.26		99	
03755332	**Acceptable identification**					**03755332**
Parabacteroides distasonis		90.06	0.44	MNS 98		
Clostridium clostridioforme		9.09	0.22			
Prevotella oralis		0.85	-.03			
03755340	**Good identification**					**03755340**
Actinomyces israelii		100.0	0.37	RHA 7		
03755342	**Good identification**					**03755342**
Actinomyces israelii		95.79	0.40	RHA 7		
Bifidobacterium adolescentis		4.21	0.10			
03755400	**Acceptable identification**					**03755400**
					SPO	
Actinomyces israelii		93.16	0.51		1	
Clostridium butyricum		4.11	0.07		99	
Lactobacillus fermentum		2.73	0.24		1	
03755402	**Low discrimination**					**03755402**
Actinomyces israelii		72.58	0.54			
Bifidobacterium adolescentis		24.73	0.40			
Lactobacillus fermentum		1.42	0.24			
03755410	**Good identification**					**03755410**
Actinomyces israelii		95.95	0.37	MLZ 13		
Bifidobacterium adolescentis		4.05	0.07			
03755412	**Low discrimination**					**03755412**
Bifidobacterium adolescentis		70.10	0.41	MNS 87		
Actinomyces israelii		25.15	0.40	LAC 99		
Clostridium clostridioforme		4.75	0.18			
03755440	**Good identification**					**03755440**
Actinomyces israelii		100.0	0.44			
03755442	**Acceptable identification**					**03755442**
Actinomyces israelii		90.23	0.47			
Bifidobacterium adolescentis		9.77	0.24			
03755500	**Low discrimination**					**03755500**
					SPO	
Actinomyces israelii		71.06	0.62		1	
Clostridium butyricum		18.82	0.31		99	
Bifidobacterium adolescentis		9.08	0.40		1	
03755502	**Low discrimination**					**03755502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium adolescentis		88.63	0.74	MNS 87	1		
Actinomyces israelii		10.51	0.65		1		
Clostridium butyricum		0.72	0.24		99		
03755503	**Low discrimination**						**03755503**
Bifidobacterium adolescentis		89.40	0.40	MNS 87			
Actinomyces israelii		10.60	0.30	URE 1			
03755510	**Low discrimination**						**03755510**
					SPO		
Bifidobacterium adolescentis		48.58	0.41	MNS 87	1		
Actinomyces israelii		46.48	0.47	BGL 99	1		
Clostridium butyricum		4.34	0.09		99		
03755512	**Very good identification**						**03755512**
Bifidobacterium adolescentis		98.30	0.76	MNS 87			
Actinomyces israelii		1.43	0.50				
Clostridium clostridioforme		0.23	0.27				
03755513	**Good identification**						**03755513**
Bifidobacterium adolescentis		98.57	0.41	MNS 87			
Actinomyces israelii		1.43	0.16	URE 1			
03755520	**Low discrimination**						**03755520**
					GRA	SPO	
Actinomyces israelii		54.29	0.41	NAG 6	99	1	
Clostridium butyricum		19.59	0.12		99	99	
Bacteroides vulgatus		18.82	0.11		1	1	
03755522	**Low discrimination**						**03755522**
					GRA		
Bifidobacterium adolescentis		39.88	0.40	MNS 87	99		
Actinomyces israelii		29.88	0.44	NAG 1	99		
Parabacteroides distasonis		19.46	0.37		1		
03755532	**Low discrimination**						**03755532**
					GRA		
Bifidobacterium adolescentis		63.14	0.41	MNS 87	99		
Clostridium clostridioforme		14.33	0.27	NAG 1	50		
Parabacteroides distasonis		13.57	0.33		1		
03755540	**Acceptable identification**						**03755540**
					SPO		
Actinomyces israelii		90.92	0.55		1		
Clostridium butyricum		5.39	0.12		99		
Bifidobacterium adolescentis		3.69	0.24		1		
03755542	**Low discrimination**						**03755542**
					SPO		
Bifidobacterium adolescentis		72.52	0.59	MNS 87	1		
Actinomyces israelii		27.06	0.58	SOR 11	1		
Clostridium butyricum		0.42	0.05		99		
03755550	**Low discrimination**						**03755550**
Actinomyces israelii		75.07	0.40	MLZ 13			
Bifidobacterium adolescentis		24.93	0.26				
03755552	**Very good identification**						**03755552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		95.64	0.60	MNS 87		
Actinomyces israelii		4.36	0.43	SOR 11		
03755562	**Low discrimination**					**03755562**
Actinomyces israelii		70.22	0.37	NAG 6		
Bifidobacterium adolescentis		29.78	0.24			
03755602	**Low discrimination**					**03755602**
Actinomyces israelii		62.35	0.35	RHA 7		
Clostridium clostridioforme		37.65	0.22			
03755700	**Low discrimination**					**03755700**
					SPO	
Actinomyces israelii		51.75	0.42	RHA 7	1	
Eubacterium contortum		41.24	0.26		1	
Clostridium butyricum		5.63	0.05		99	
03755702	**Low discrimination**					**03755702**
Bifidobacterium adolescentis		40.77	0.40	MNS 87		
Actinomyces israelii		36.03	0.45	RHA 1		
Clostridium clostridioforme		18.20	0.31			
03755712	**Low discrimination**					**03755712**
Clostridium clostridioforme		60.03	0.42	MNS 99		
Bifidobacterium adolescentis		36.07	0.41	TRE 88		
Actinomyces israelii		3.90	0.31			
03755722	**Low discrimination**					**03755722**
Parabacteroides distasonis		68.73	0.45	MNS 98		
Clostridium clostridioforme		20.97	0.31			
Bacteroides ovatus		2.83	0.06			
03755732	**Low discrimination**					**03755732**
Clostridium clostridioforme		70.76	0.42	MNS 99		
Parabacteroides distasonis		27.42	0.41	TRE 88		
Bacteroides ovatus		1.40	0.03			
03755740	**Good identification**					**03755740**
Actinomyces israelii		100.0	0.35	RHA 7		
03755742	**Low discrimination**					**03755742**
Actinomyces israelii		73.55	0.38	RHA 7		
Bifidobacterium adolescentis		26.45	0.24			
03756100	**Good identification**					**03756100**
					GRA	
Actinomyces israelii		97.55	0.42	GAL 99	99	
Fusobacterium mortiferum		2.45	0.12		1	
03756102	**Low discrimination**					**03756102**
Actinomyces israelii		87.04	0.45	GAL 99		
Bifidobacterium adolescentis		12.96	0.24			
03756142	**Good identification**					**03756142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		95.48	0.38	GAL 99		
Bifidobacterium adolescentis		4.52	0.09			
03756500	**Good identification**					**03756500**
Actinomyces israelii		100.0	0.40	GAL 99		
03756502	**Low discrimination**					**03756502**
Bifidobacterium adolescentis		54.91	0.38	MNS 87		
Actinomyces israelii		45.09	0.43	GAL 99		
03756512	**Acceptable identification**					**03756512**
Bifidobacterium adolescentis		90.88	0.40	MNS 87		
Actinomyces israelii		9.13	0.29	GAL 99		
03756542	**Low discrimination**					**03756542**
Actinomyces israelii		72.10	0.36	GAL 99		
Bifidobacterium adolescentis		27.90	0.23			
03757000	**Very good identification**					**03757000**
Actinomyces israelii		99.51	0.66			
Actinomyces naeslundii		0.49	0.35			
03757002	**Very good identification**					**03757002**
Actinomyces israelii		99.08	0.69			
Bifidobacterium adolescentis		0.60	0.24			
Actinomyces naeslundii		0.33	0.35			
03757010	**Good identification**					**03757010**
Actinomyces israelii		100.0	0.51	MLZ 13		
03757012	**Good identification**					**03757012**
Actinomyces israelii		95.31	0.55	MLZ 13		
Bifidobacterium adolescentis		4.69	0.26			
03757020	**Acceptable identification**					**03757020**
Actinomyces israelii		92.81	0.45	NAG 6		
Actinomyces naeslundii		7.19	0.35			
03757021	**Good identification**					**03757021**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03757022	**Good identification**					**03757022**
Actinomyces israelii		95.09	0.48	NAG 6		
Actinomyces naeslundii		4.91	0.35			
03757023	**Low discrimination**					**03757023**
Actinomyces naeslundii		88.46	0.38	MNS 88		
Actinomyces israelii		11.54	0.14	XYL 1		
03757040	**Very good identification**					**03757040**
Actinomyces israelii		99.78	0.59			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.22	0.22			
03757042	**Very good identification**					**03757042**
Actinomyces israelii		99.66	0.62			
Bifidobacterium adolescentis		0.19	0.09			
Actinomyces naeslundii		0.15	0.22			
03757050	**Good identification**					**03757050**
Actinomyces israelii		100.0	0.44	MLZ 13		
03757052	**Good identification**					**03757052**
Actinomyces israelii		98.46	0.47	MLZ 13		
Bifidobacterium adolescentis		1.54	0.10			
03757060	**Good identification**					**03757060**
Actinomyces israelii		96.61	0.38	NAG 6		
Actinomyces naeslundii		3.40	0.22			
03757062	**Good identification**					**03757062**
Actinomyces israelii		97.71	0.41	NAG 6		
Actinomyces naeslundii		2.29	0.22			
03757100	**Very good identification**					**03757100**
Actinomyces israelii		99.34	0.76			
Actinomyces naeslundii		0.35	0.43			
Bifidobacterium adolescentis		0.22	0.24			
03757101	**Low discrimination**					**03757101**
Actinomyces israelii		65.69	0.42	URE 1		
Actinomyces naeslundii		34.31	0.46			
03757102	**Low discrimination**					**03757102**
Actinomyces israelii		86.87	0.79			
Bifidobacterium adolescentis		12.93	0.58			
Actinomyces naeslundii		0.20	0.43			
03757103	**Low discrimination**					**03757103**
Actinomyces israelii		66.80	0.45	URE 1		
Actinomyces naeslundii		23.26	0.46			
Bifidobacterium adolescentis		9.94	0.24			
03757110	**Very good identification**					**03757110**
Actinomyces israelii		98.19	0.62	MLZ 13		
Bifidobacterium adolescentis		1.81	0.26			
03757112	**Low discrimination**					**03757112**
					SPO	
Bifidobacterium adolescentis		54.88	0.60	MNS 87	1	
Actinomyces israelii		45.07	0.65	ARA 87	1	
Clostridium tertium		0.05	-.01		99	
03757120	**Acceptable identification**					**03757120**
Actinomyces israelii		94.78	0.55	NAG 6		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	5.22	0.43			
03757121	**Low discrimination**					**03757121**
	Actinomyces naeslundii	89.11	0.46	MNS 88		
	Actinomyces israelii	10.89	0.21	XYL 1		
03757122	**Good identification**					**03757122**
	Actinomyces israelii	92.95	0.58	NAG 6		
	Actinomyces naeslundii	3.42	0.43			
	Bifidobacterium adolescentis	2.19	0.24			
03757123	**Low discrimination**					**03757123**
	Actinomyces naeslundii	84.51	0.46	MNS 88		
	Actinomyces israelii	15.49	0.24	XYL 1		
03757130	**Good identification**					**03757130**
					SPO	
	Actinomyces israelii	97.72	0.41	MLZ 13	1	
	Clostridium tertium	2.29	-.01	NAG 6	99	
03757132	**Low discrimination**					**03757132**
					SPO	
	Actinomyces israelii	80.68	0.44	MLZ 13	1	
	Bifidobacterium adolescentis	15.55	0.26	NAG 6	1	
	Clostridium tertium	3.77	0.07		99	
03757140	**Very good identification**					**03757140**
	Actinomyces israelii	99.77	0.69			
	Actinomyces naeslundii	0.16	0.30			
	Bifidobacterium adolescentis	0.07	0.09			
03757142	**Very good identification**					**03757142**
	Actinomyces israelii	95.39	0.72			
	Bifidobacterium adolescentis	4.51	0.43			
	Actinomyces naeslundii	0.10	0.30			
03757143	**Low discrimination**					**03757143**
	Actinomyces israelii	82.97	0.38	URE 1		
	Actinomyces naeslundii	13.11	0.33			
	Bifidobacterium adolescentis	3.93	0.09			
03757150	**Good identification**					**03757150**
	Actinomyces israelii	99.42	0.55	MLZ 13		
	Bifidobacterium adolescentis	0.58	0.10			
03757152	**Low discrimination**					**03757152**
	Actinomyces israelii	72.10	0.58	MLZ 13		
	Bifidobacterium adolescentis	27.90	0.44			
03757160	**Good identification**					**03757160**
	Actinomyces israelii	97.56	0.48	NAG 6		
	Actinomyces naeslundii	2.44	0.30			
03757162	**Good identification**					**03757162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		97.64	0.51	NAG 6			
Actinomyces naeslundii		1.63	0.30				
Bifidobacterium adolescentis		0.73	0.09				
03757172	**Acceptable identification**						**03757172**
Actinomyces israelii		94.23	0.37	MLZ 13			
Bifidobacterium adolescentis		5.77	0.10	NAG 6			
03757200	**Good identification**						**03757200**
Actinomyces israelii		100.0	0.46	RHA 7			
03757202	**Good identification**						**03757202**
Actinomyces israelii		100.0	0.49	RHA 7			
03757212	**Good identification**						**03757212**
Actinomyces israelii		100.0	0.35	RHA 7			
03757240	**Good identification**						**03757240**
Actinomyces israelii		100.0	0.39	RHA 7			
03757242	**Good identification**						**03757242**
Actinomyces israelii		100.0	0.42	RHA 7			
03757300	**Very good identification**						**03757300**
Actinomyces israelii		100.0	0.57	RHA 7			
03757302	**Very good identification**						**03757302**
Actinomyces israelii		98.04	0.60	RHA 7			
Bifidobacterium adolescentis		1.96	0.24				
03757310	**Good identification**						**03757310**
Actinomyces israelii		100.0	0.43	RHA 7			
03757312	**Low discrimination**						**03757312**
Actinomyces israelii		85.96	0.46	RHA 7			
Bifidobacterium adolescentis		14.05	0.26	MLZ 13			
03757320	**Good identification**						**03757320**
Actinomyces israelii		100.0	0.36	RHA 7			
03757322	**Low discrimination**						**03757322**
					SPO	GRA	
Actinomyces israelii		83.23	0.39	RHA 7	1	99	
Clostridium ramosum		9.03	0.07	NAG 6	92	50	
Parabacteroides distasonis		5.14	0.15		1	1	
03757340	**Good identification**						**03757340**
Actinomyces israelii		100.0	0.50	RHA 7			
03757342	**Good identification**						**03757342**
Actinomyces israelii		99.37	0.53	RHA 7			
Bifidobacterium adolescentis		0.63	0.09				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03757350	**Good identification**					**03757350**
Actinomyces israelii		100.0	0.36	RHA 7		
03757352	**Good identification**					**03757352**
Actinomyces israelii		95.06	0.39	RHA 7		
Bifidobacterium adolescentis		4.94	0.10	MLZ 13		
03757400	**Very good identification**					**03757400**
Actinomyces israelii		100.0	0.64			
03757402	**Very good identification**					**03757402**
Actinomyces israelii		95.31	0.67			
Bifidobacterium adolescentis		4.69	0.38			
03757410	**Good identification**					**03757410**
Actinomyces israelii		100.0	0.50	MLZ 13		
03757412	**Low discrimination**					**03757412**
Actinomyces israelii		71.31	0.53	MLZ 13		
Bifidobacterium adolescentis		28.69	0.40			
03757420	**Good identification**					**03757420**
Actinomyces israelii		100.0	0.44	NAG 6		
03757422	**Good identification**					**03757422**
Actinomyces israelii		100.0	0.47	NAG 6		
03757440	**Very good identification**					**03757440**
Actinomyces israelii		100.0	0.57			
03757442	**Very good identification**					**03757442**
Actinomyces israelii		98.46	0.60			
Bifidobacterium adolescentis		1.54	0.23			
03757450	**Good identification**					**03757450**
Actinomyces israelii		100.0	0.43	MLZ 13		
03757452	**Low discrimination**					**03757452**
Actinomyces israelii		88.66	0.46	MLZ 13		
Bifidobacterium adolescentis		11.34	0.24			
03757460	**Good identification**					**03757460**
Actinomyces israelii		100.0	0.37	NAG 6		
03757462	**Good identification**					**03757462**
Actinomyces israelii		100.0	0.40	NAG 6		
03757500	**Very good identification**					**03757500**
					SPO	
Actinomyces israelii		97.69	0.74		1	
Bifidobacterium adolescentis		1.80	0.38		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		0.51	0.14		99	
03757501	**Good identification**					**03757501**
Actinomyces israelii		100.0	0.40	URE 1		
03757502	**Low discrimination**					**03757502**
					SPO	
Bifidobacterium adolescentis		54.87	0.73	MNS 87	1	
Actinomyces israelii		45.07	0.77		1	
Clostridium butyricum		0.06	0.07		99	
03757503	**Low discrimination**					**03757503**
Bifidobacterium adolescentis		54.91	0.38	MNS 87		
Actinomyces israelii		45.09	0.43	URE 1		
03757510	**Low discrimination**					**03757510**
					SPO	
Actinomyces israelii		86.75	0.60	MLZ 13	1	
Bifidobacterium adolescentis		13.09	0.40		1	
Clostridium butyricum		0.16	-.08		99	
03757512	**Good identification**					**03757512**
Bifidobacterium adolescentis		90.86	0.74	MNS 87		
Actinomyces israelii		9.12	0.63			
Bifidobacterium dentium		0.02	-.07			
03757513	**Acceptable identification**					**03757513**
Bifidobacterium adolescentis		90.88	0.40	MNS 87		
Actinomyces israelii		9.13	0.29	URE 1		
03757520	**Good identification**					**03757520**
					SPO	
Actinomyces israelii		99.30	0.54	NAG 6	1	
Clostridium butyricum		0.70	-.04		99	
03757522	**Low discrimination**					**03757522**
Actinomyces israelii		83.85	0.57	NAG 6		
Bifidobacterium adolescentis		16.16	0.38			
03757530	**Good identification**					**03757530**
Actinomyces israelii		100.0	0.40	MLZ 13		
03757532	**Low discrimination**					**03757532**
Bifidobacterium adolescentis		59.80	0.40	MNS 87		
Actinomyces israelii		37.95	0.43	NAG 1		
Bifidobacterium dentium		2.25	-.03			
03757540	**Very good identification**					**03757540**
					SPO	
Actinomyces israelii		99.30	0.67		1	
Bifidobacterium adolescentis		0.58	0.23		1	
Clostridium butyricum		0.12	-.04		99	
03757542	**Low discrimination**					**03757542**
Actinomyces israelii		72.10	0.70			
Bifidobacterium adolescentis		27.90	0.57			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03757543	**Low discrimination**					**03757543**
Actinomyces israelii		72.10	0.36	URE 1		
Bifidobacterium adolescentis		27.90	0.23			
03757550	**Acceptable identification**					**03757550**
					COC	
Actinomyces israelii		94.69	0.53	MLZ 13	1	
Bifidobacterium adolescentis		4.54	0.24		1	
Blautia producta		0.77	-.03		99	
03757552	**Low discrimination**					**03757552**
Bifidobacterium adolescentis		75.99	0.59	MNS 87		
Actinomyces israelii		24.01	0.56	SOR 11		
03757560	**Good identification**					**03757560**
Actinomyces israelii		100.0	0.47	NAG 6		
03757562	**Acceptable identification**					**03757562**
Actinomyces israelii		94.23	0.50	NAG 6		
Bifidobacterium adolescentis		5.77	0.23			
03757572	**Low discrimination**					**03757572**
Actinomyces israelii		66.63	0.36	MLZ 13		
Bifidobacterium adolescentis		33.37	0.24	NAG 6		
03757600	**Good identification**					**03757600**
Actinomyces israelii		100.0	0.45	RHA 7		
03757602	**Good identification**					**03757602**
Actinomyces israelii		100.0	0.48	RHA 7		
03757640	**Good identification**					**03757640**
Actinomyces israelii		100.0	0.38	RHA 7		
03757642	**Good identification**					**03757642**
Actinomyces israelii		100.0	0.41	RHA 7		
03757700	**Good identification**					**03757700**
Actinomyces israelii		100.0	0.55	RHA 7		
03757702	**Low discrimination**					**03757702**
Actinomyces israelii		85.96	0.58	RHA 7		
Bifidobacterium adolescentis		14.05	0.38			
03757710	**Good identification**					**03757710**
Actinomyces israelii		100.0	0.41	RHA 7		
03757712	**Low discrimination**					**03757712**
Bifidobacterium adolescentis		56.54	0.40	MNS 87		
Actinomyces israelii		42.30	0.44	RHA 1		
Clostridium clostridioforme		1.16	0.08			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03757722	**Acceptable identification**					**03757722**
					GRA	
Actinomyces israelii		90.16	0.38	RHA 7	99	
Bacteroides ovatus		5.67	-.03	NAG 6	1	
Parabacteroides distasonis		4.17	0.11		1	
03757740	**Good identification**					**03757740**
Actinomyces israelii		100.0	0.48	RHA 7		
03757742	**Good identification**					**03757742**
Actinomyces israelii		95.06	0.51	RHA 7		
Bifidobacterium adolescentis		4.94	0.23			
03757752	**Low discrimination**					**03757752**
Actinomyces israelii		70.19	0.37	RHA 7		
Bifidobacterium adolescentis		29.81	0.24	MLZ 13		
03761000	**Acceptable identification**					**03761000**
					GRA	
Actinomyces naeslundii		90.88	0.48	MNS 88	99	
Fusobacterium mortiferum		9.13	0.19	FRU 95	1	
03761001	**Good identification**					**03761001**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03761002	**Acceptable identification**					**03761002**
Actinomyces naeslundii		92.06	0.48	MNS 88		
Lactobacillus acidophilus		4.81	0.01	FRU 95		
Actinomyces israelii		3.13	0.13			
03761003	**Good identification**					**03761003**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03761020	**Good identification**					**03761020**
Actinomyces naeslundii		100.0	0.48	MNS 88		
03761021	**Good identification**					**03761021**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03761022	**Good identification**					**03761022**
Actinomyces naeslundii		95.03	0.48	MNS 88		
Lactobacillus acidophilus		4.97	0.01	FRU 95		
03761023	**Good identification**					**03761023**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03761041	**Good identification**					**03761041**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03761043	**Good identification**					**03761043**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03761061	**Good identification**					**03761061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
03761063	**Good identification**					**03761063**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03761100	**Low discrimination**					**03761100**
					GRA	
Actinomyces naeslundii		89.76	0.55	MNS 88	99	
Fusobacterium mortiferum		7.39	0.25	FRU 95	1	
Actinomyces israelii		2.86	0.21		99	
03761101	**Very good identification**					**03761101**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03761102	**Low discrimination**					**03761102**
Actinomyces naeslundii		78.49	0.55	MNS 88		
Lactobacillus acidophilus		16.58	0.19	FRU 95		
Actinomyces israelii		3.75	0.24			
03761103	**Very good identification**					**03761103**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03761111	**Good identification**					**03761111**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03761113	**Good identification**					**03761113**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03761120	**Good identification**					**03761120**
					SPO	
Actinomyces naeslundii		97.82	0.55	MNS 88	1	
Clostridium ramosum		2.18	0.02	FRU 95	92	
03761121	**Very good identification**					**03761121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03761122	**Low discrimination**					**03761122**
					SPO	
Actinomyces naeslundii		76.04	0.55	MNS 88	1	
Lactobacillus acidophilus		16.07	0.19	FRU 95	1	
Clostridium ramosum		6.77	0.12		92	
03761123	**Very good identification**					**03761123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03761131	**Good identification**					**03761131**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03761133	**Good identification**					**03761133**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03761140	**Acceptable identification**					**03761140**
Actinomyces naeslundii		93.45	0.43	MNS 88		
Actinomyces israelii		6.55	0.14	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03761141	**Good identification**					**03761141**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03761142	**Low discrimination**					**03761142**
Actinomyces naeslundii		79.80	0.43	MNS 88		
Lactobacillus acidophilus		11.81	0.04	FRU 95		
Actinomyces israelii		8.39	0.17			
03761143	**Good identification**					**03761143**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03761160	**Good identification**					**03761160**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03761161	**Good identification**					**03761161**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03761162	**Low discrimination**					**03761162**
Actinomyces naeslundii		87.11	0.43	MNS 88		
Lactobacillus acidophilus		12.89	0.04	FRU 95		
03761163	**Good identification**					**03761163**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03762100	**Good identification**					**03762100**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03762101	**Good identification**					**03762101**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03762102	**Good identification**					**03762102**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03762103	**Good identification**					**03762103**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03762120	**Good identification**					**03762120**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03762121	**Good identification**					**03762121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03762122	**Good identification**					**03762122**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03762123	**Good identification**					**03762123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03763000	**Acceptable identification**					**03763000**
Actinomyces naeslundii		93.80	0.53	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		6.20	0.23	FRU 95		
03763001	**Good identification**					**03763001**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03763002	**Acceptable identification**					**03763002**
Actinomyces naeslundii		90.98	0.53	MNS 88		
Actinomyces israelii		9.02	0.26	FRU 95		
03763003	**Good identification**					**03763003**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03763020	**Good identification**					**03763020**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03763021	**Good identification**					**03763021**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03763022	**Good identification**					**03763022**
Actinomyces naeslundii		100.0	0.53	MNS 88		
03763023	**Good identification**					**03763023**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03763040	**Low discrimination**					**03763040**
Actinomyces naeslundii		87.29	0.40	MNS 88		
Actinomyces israelii		12.71	0.16	FRU 95		
03763041	**Good identification**					**03763041**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03763042	**Low discrimination**					**03763042**
Actinomyces naeslundii		82.07	0.40	MNS 88		
Actinomyces israelii		17.93	0.19	FRU 95		
03763043	**Good identification**					**03763043**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03763060	**Good identification**					**03763060**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03763061	**Good identification**					**03763061**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03763062	**Good identification**					**03763062**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03763063	**Good identification**					**03763063**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03763100	**Good identification**					**03763100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.50	0.60	MNS 88		
Actinomyces israelii		8.50	0.33	FRU 95		
03763101	**Very good identification**					**03763101**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03763102	**Low discrimination**					**03763102**
Actinomyces naeslundii		87.29	0.60	MNS 88		
Actinomyces israelii		12.16	0.36	FRU 95		
Bifidobacterium adolescentis		0.55	0.07			
03763103	**Very good identification**					**03763103**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03763110	**Low discrimination**					**03763110**
Actinomyces naeslundii		79.13	0.38	MNS 88		
Actinomyces israelii		20.87	0.19	FRU 95		
03763111	**Good identification**					**03763111**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03763112	**Low discrimination**					**03763112**
Actinomyces naeslundii		64.84	0.38	MNS 88		
Actinomyces israelii		25.65	0.22	FRU 95		
Bifidobacterium adolescentis		9.51	0.09	MLZ 5		
03763113	**Good identification**					**03763113**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03763120	**Good identification**					**03763120**
					SPO	
Actinomyces naeslundii		93.90	0.60	MNS 88	1	
Clostridium ramosum		6.10	0.15	FRU 95	92	
03763121	**Very good identification**					**03763121**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03763122	**Low discrimination**					**03763122**
					SPO	
Actinomyces naeslundii		78.82	0.60	MNS 88	1	
Clostridium ramosum		20.48	0.25	FRU 95	92	
Actinomyces israelii		0.70	0.16		1	
03763123	**Very good identification**					**03763123**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03763130	**Good identification**					**03763130**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03763131	**Good identification**					**03763131**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03763132	**Good identification**					**03763132**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03763133	**Good identification**					**03763133**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03763140	**Low discrimination**					**03763140**
Actinomyces naeslundii		83.01	0.47	MNS 88		
Actinomyces israelii		16.99	0.26	FRU 95		
03763141	**Good identification**					**03763141**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03763142	**Low discrimination**					**03763142**
Actinomyces naeslundii		76.51	0.47	MNS 88		
Actinomyces israelii		23.49	0.29	FRU 95		
03763143	**Good identification**					**03763143**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03763160	**Good identification**					**03763160**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03763161	**Good identification**					**03763161**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03763162	**Good identification**					**03763162**
Actinomyces naeslundii		100.0	0.47	MNS 88		
03763163	**Good identification**					**03763163**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03765100	**Acceptable identification**					**03765100**
					SPO	
Actinomyces israelii		90.89	0.39	FRU 99	1	
Clostridium butyricum		9.12	0.01		99	
03765102	**Low discrimination**					**03765102**
					SPO	
Actinomyces israelii		82.73	0.42	FRU 99	1	
Bifidobacterium adolescentis		15.12	0.23		1	
Clostridium butyricum		2.15	-.07		99	
03765502	**Low discrimination**					**03765502**
					SPO	
Bifidobacterium adolescentis		53.75	0.37	MNS 87	1	
Actinomyces israelii		35.97	0.40	FRU 99	1	
Clostridium butyricum		10.29	0.10		99	
03765512	**Acceptable identification**					**03765512**
Bifidobacterium adolescentis		90.33	0.38	MNS 87		
Actinomyces israelii		7.39	0.26	FRU 99		
Clostridium clostridioforme		2.28	0.08			
03767000	**Good identification**					**03767000**
Actinomyces israelii		100.0	0.41	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03767002	**Good identification**					**03767002**
Actinomyces israelii		100.0	0.44	FRU 99		
03767042	**Good identification**					**03767042**
Actinomyces israelii		100.0	0.37	FRU 99		
03767100	**Good identification**					**03767100**
Actinomyces israelii		99.06	0.52	FRU 99		
Actinomyces naeslundii		0.94	0.26			
03767102	**Good identification**					**03767102**
Actinomyces israelii		96.84	0.55	FRU 99		
Bifidobacterium adolescentis		2.56	0.21			
Actinomyces naeslundii		0.61	0.26			
03767110	**Good identification**					**03767110**
Actinomyces israelii		100.0	0.37	FRU 99		
03767112	**Low discrimination**					**03767112**
Actinomyces israelii		82.25	0.41	FRU 99		
Bifidobacterium adolescentis		17.75	0.23	MLZ 13		
03767140	**Good identification**					**03767140**
Actinomyces israelii		100.0	0.45	FRU 99		
03767142	**Good identification**					**03767142**
Actinomyces israelii		100.0	0.48	FRU 99		
03767302	**Good identification**					**03767302**
Actinomyces israelii		100.0	0.35	FRU 99		
03767400	**Good identification**					**03767400**
Actinomyces israelii		100.0	0.40	FRU 99		
03767402	**Good identification**					**03767402**
Actinomyces israelii		100.0	0.43	FRU 99		
03767442	**Good identification**					**03767442**
Actinomyces israelii		100.0	0.36	FRU 99		
03767500	**Good identification**					**03767500**
					SPO	
Actinomyces israelii		97.88	0.50	FRU 99	1	
Clostridium butyricum		2.12	0.01		99	
03767502	**Low discrimination**					**03767502**
					SPO	
Actinomyces israelii		81.88	0.53	FRU 99	1	
Bifidobacterium adolescentis		17.67	0.35		1	
Clostridium butyricum		0.46	-.07		99	
03767510	**Good identification**					**03767510**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	FRU 99		
03767512	**Low discrimination**					**03767512**
Bifidobacterium adolescentis		62.56	0.37	MNS 87		
Actinomyces israelii		35.45	0.39	FRU 99		
Bifidobacterium dentium		1.99	-.07			
03767540	**Good identification**					**03767540**
Actinomyces israelii		100.0	0.43	FRU 99		
03767542	**Acceptable identification**					**03767542**
Actinomyces israelii		93.58	0.46	FRU 99		
Bifidobacterium adolescentis		6.42	0.20			
03770000	**Low discrimination**					**03770000**
					GRA	
Fusobacterium mortiferum		89.33	0.50	MLT 11	1	
Actinomyces naeslundii		10.67	0.46		99	
03770001	**Acceptable identification**					**03770001**
					GRA	
Actinomyces naeslundii		94.66	0.49	MNS 88	99	
Fusobacterium mortiferum		5.34	0.16	GAL 96	1	
03770002	**Low discrimination**					**03770002**
					GRA	
Actinomyces naeslundii		47.93	0.46	MNS 88	99	
Fusobacterium mortiferum		44.58	0.34	GAL 96	1	
Lactobacillus acidophilus		3.17	0.01		99	
03770003	**Good identification**					**03770003**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03770020	**Acceptable identification**					**03770020**
					GRA	
Actinomyces naeslundii		92.20	0.46	MNS 88	99	
Fusobacterium mortiferum		7.80	0.16	GAL 96	1	
03770021	**Good identification**					**03770021**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03770022	**Acceptable identification**					**03770022**
Actinomyces naeslundii		93.80	0.46	MNS 88		
Lactobacillus acidophilus		6.20	0.01	GAL 96		
03770023	**Good identification**					**03770023**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03770041	**Good identification**					**03770041**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03770043	**Good identification**					**03770043**
Actinomyces naeslundii		100.0	0.36	MNS 88		
03770061	**Good identification**					**03770061**
Actinomyces naeslundii		100.0	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
03770063	**Good identification**						**03770063**
Actinomyces naeslundii		100.0	0.36	MNS 88			
03770100	**Low discrimination**						**03770100**
					GRA		
Fusobacterium mortiferum		86.84	0.56	MLT 11	1		
Actinomyces naeslundii		12.65	0.54		99		
Actinomyces israelii		0.51	0.21		99		
03770101	**Very good identification**						**03770101**
					GRA		
Actinomyces naeslundii		95.58	0.57	MNS 88	99		
Fusobacterium mortiferum		4.42	0.22	GAL 96	1		
03770102	**Low discrimination**						**03770102**
					GRA		
Actinomyces naeslundii		46.77	0.54	MNS 88	99		
Fusobacterium mortiferum		35.66	0.40	GAL 96	1		
Lactobacillus acidophilus		12.48	0.19		99		
03770103	**Very good identification**						**03770103**
Actinomyces naeslundii		100.0	0.57	MNS 88			
03770120	**Acceptable identification**						**03770120**
					SPO	GRA	
Actinomyces naeslundii		91.12	0.54	MNS 88	1	99	
Fusobacterium mortiferum		6.32	0.22	GAL 96	1	1	
Clostridium ramosum		2.56	0.02		92	50	
03770121	**Very good identification**						**03770121**
Actinomyces naeslundii		100.0	0.57	MNS 88			
03770122	**Low discrimination**						**03770122**
					SPO		
Actinomyces naeslundii		71.53	0.54	MNS 88	1		
Lactobacillus acidophilus		19.09	0.19	GAL 96	1		
Clostridium ramosum		8.04	0.12		92		
03770123	**Very good identification**						**03770123**
Actinomyces naeslundii		100.0	0.57	MNS 88			
03770140	**Low discrimination**						**03770140**
					GRA		
Actinomyces naeslundii		67.51	0.41	MNS 88	99		
Fusobacterium mortiferum		26.52	0.22	GAL 96	1		
Actinomyces israelii		5.98	0.14		99		
03770141	**Good identification**						**03770141**
Actinomyces naeslundii		100.0	0.44	MNS 88			
03770142	**Low discrimination**						**03770142**
Actinomyces naeslundii		75.77	0.41	MNS 88			
Lactobacillus acidophilus		14.16	0.04	GAL 96			
Actinomyces israelii		10.07	0.17				
03770143	**Good identification**						**03770143**
Actinomyces naeslundii		100.0	0.44	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03770160	**Good identification**					**03770160**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03770161	**Good identification**					**03770161**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03770162	**Low discrimination**					**03770162**
Actinomyces naeslundii		84.25	0.41	MNS 88		
Lactobacillus acidophilus		15.75	0.04	GAL 96		
03770163	**Good identification**					**03770163**
Actinomyces naeslundii		100.0	0.44	MNS 88		
03771000	**Low discrimination**					**03771000**
					GRA	
Actinomyces naeslundii		56.94	0.70	MNS 88	99	
Fusobacterium mortiferum		29.79	0.53		1	
Actinomyces israelii		6.72	0.44		99	
03771001	**Very good identification**					**03771001**
					GRA	
Actinomyces naeslundii		99.65	0.73	MNS 88	99	
Fusobacterium mortiferum		0.35	0.19		1	
03771002	**Low discrimination**					**03771002**
Actinomyces naeslundii		63.10	0.70	MNS 88		
Lactobacillus acidophilus		17.19	0.35			
Actinomyces israelii		11.17	0.47			
03771003	**Very good identification**					**03771003**
Actinomyces naeslundii		99.70	0.73	MNS 88		
Lactobacillus acidophilus		0.18	0.01			
Actinomyces israelii		0.12	0.13			
03771010	**Low discrimination**					**03771010**
Actinomyces naeslundii		46.57	0.48	MNS 88		
Lactobacillus jensenii		33.16	0.24	MLZ 5		
Actinomyces israelii		15.60	0.30			
03771011	**Good identification**					**03771011**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03771012	**Low discrimination**					**03771012**
Actinomyces naeslundii		47.15	0.48	MNS 88		
Actinomyces israelii		23.69	0.33	MLZ 5		
Lactobacillus jensenii		22.38	0.21			
03771013	**Good identification**					**03771013**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03771020	**Very good identification**					**03771020**
Actinomyces naeslundii		95.21	0.70	MNS 88		
Lactobacillus jensenii		2.68	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.72	0.24			
03771021	**Very good identification**					**03771021**
Actinomyces naeslundii		100.0	0.73	MNS 88		
03771022	**Low discrimination**					**03771022**
Actinomyces naeslundii		75.20	0.70	MNS 88		
Lactobacillus acidophilus		20.49	0.35			
Lactobacillus jensenii		1.41	0.18			
03771023	**Very good identification**					**03771023**
Actinomyces naeslundii		99.82	0.73	MNS 88		
Lactobacillus acidophilus		0.18	0.01			
03771030	**Low discrimination**					**03771030**
Actinomyces naeslundii		84.89	0.48	MNS 88		
Lactobacillus jensenii		15.11	0.13	MLZ 5		
03771031	**Good identification**					**03771031**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03771032	**Low discrimination**					**03771032**
Actinomyces naeslundii		85.40	0.48	MNS 88		
Lactobacillus jensenii		10.14	0.10	MLZ 5		
Lactobacillus acidophilus		4.47	0.01			
03771033	**Good identification**					**03771033**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03771040	**Low discrimination**					**03771040**
					GRA	
Actinomyces naeslundii		77.53	0.57	MNS 88	99	
Actinomyces israelii		20.15	0.37		99	
Fusobacterium mortiferum		2.32	0.19		1	
03771041	**Very good identification**					**03771041**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03771042	**Low discrimination**					**03771042**
Actinomyces naeslundii		63.26	0.57	MNS 88		
Actinomyces israelii		24.67	0.40			
Lactobacillus acidophilus		12.07	0.20			
03771043	**Very good identification**					**03771043**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03771051	**Good identification**					**03771051**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771053	**Good identification**					**03771053**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771060	**Very good identification**					**03771060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.37	0.57	MNS 88		
Actinomyces israelii		1.63	0.17			
03771061	**Very good identification**					**03771061**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03771062	**Low discrimination**					**03771062**
Actinomyces naeslundii		82.26	0.57	MNS 88		
Lactobacillus acidophilus		15.70	0.20			
Actinomyces israelii		2.05	0.20			
03771063	**Very good identification**					**03771063**
Actinomyces naeslundii		100.0	0.60	MNS 88		
03771071	**Good identification**					**03771071**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771073	**Good identification**					**03771073**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771100	**Low discrimination**					**03771100**
					GRA	
Actinomyces naeslundii		61.59	0.77	MNS 88	99	
Fusobacterium mortiferum		26.41	0.59		1	
Actinomyces israelii		10.21	0.55		99	
03771101	**Very good identification**					**03771101**
					GRA	
Actinomyces naeslundii		99.60	0.80	MNS 88	99	
Fusobacterium mortiferum		0.29	0.25		1	
Actinomyces israelii		0.11	0.21		99	
03771102	**Low discrimination**					**03771102**
Lactobacillus acidophilus		44.12	0.54	MNS 99		
Actinomyces naeslundii		40.08	0.77			
Actinomyces israelii		9.97	0.58			
03771103	**Very good identification**					**03771103**
Actinomyces naeslundii		99.05	0.80	MNS 88		
Lactobacillus acidophilus		0.73	0.19			
Actinomyces israelii		0.17	0.24			
03771110	**Low discrimination**					**03771110**
					GRA	
Actinomyces naeslundii		63.09	0.55	MNS 88	99	
Actinomyces israelii		29.69	0.41	MLZ 5	99	
Fusobacterium mortiferum		5.19	0.25		1	
03771111	**Very good identification**					**03771111**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03771112	**Low discrimination**					**03771112**
					GRA	
Bifidobacterium adolescentis		34.69	0.44	MNS 87	99	
Leptotrichia buccalis		26.28	0.24	XYL 87	1	
Actinomyces naeslundii		19.13	0.55	ARA 87	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03771113	**Very good identification**					**03771113**
					GRA	
Actinomyces naeslundii		97.90	0.58	MNS 88	99	
Bifidobacterium adolescentis		1.20	0.10	MLZ 5	99	
Leptotrichia buccalis		0.91	-.11		1	
03771120	**Low discrimination**					**03771120**
					SPO	
Actinomyces naeslundii		86.06	0.77	MNS 88	1	
Clostridium ramosum		9.98	0.36		92	
Lactobacillus acidophilus		0.96	0.19		1	
03771121	**Very good identification**					**03771121**
					SPO	
Actinomyces naeslundii		99.92	0.80	MNS 88	1	
Clostridium ramosum		0.08	0.02		92	
03771122	**Low discrimination**					**03771122**
					SPO	
Lactobacillus acidophilus		40.83	0.54	MNS 99	1	
Actinomyces naeslundii		37.09	0.77		1	
Clostridium ramosum		17.20	0.46		92	
03771123	**Very good identification**					**03771123**
					SPO	
Actinomyces naeslundii		98.91	0.80	MNS 88	1	
Lactobacillus acidophilus		0.73	0.19		1	
Clostridium ramosum		0.31	0.12		92	
03771130	**Acceptable identification**					**03771130**
					SPO	
Actinomyces naeslundii		93.32	0.55	MNS 88	1	
Actinomyces israelii		2.80	0.20	MLZ 5	1	
Clostridium ramosum		2.08	0.02		92	
03771131	**Very good identification**					**03771131**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03771132	**Low discrimination**					**03771132**
					GRA	
Actinomyces naeslundii		51.08	0.55	MNS 88	99	
Parabacteroides distasonis		11.16	0.31	MLZ 5	1	
Lactobacillus acidophilus		10.79	0.19		99	
03771133	**Very good identification**					**03771133**
Actinomyces naeslundii		100.0	0.58	MNS 88		
03771140	**Low discrimination**					**03771140**
					GRA	
Actinomyces naeslundii		71.55	0.64	MNS 88	99	
Actinomyces israelii		26.14	0.48		99	
Fusobacterium mortiferum		1.76	0.25		1	
03771141	**Very good identification**					**03771141**
Actinomyces naeslundii		99.76	0.67	MNS 88		
Actinomyces israelii		0.25	0.14			
03771142	**Low discrimination**					**03771142**
Actinomyces naeslundii		41.56	0.64	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		32.04	0.38			
Actinomyces israelii		22.77	0.51			
03771143	**Very good identification**					**03771143**
Actinomyces naeslundii		99.12	0.67	MNS 88		
Lactobacillus acidophilus		0.52	0.04			
Actinomyces israelii		0.37	0.17			
03771151	**Good identification**					**03771151**
Actinomyces naeslundii		100.0	0.46	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03771152	**Low discrimination**					**03771152**
Actinomyces israelii		33.28	0.37	XYL 92		
Bifidobacterium adolescentis		27.16	0.29	MLZ 13		
Actinomyces naeslundii		21.39	0.43			
03771153	**Good identification**					**03771153**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03771160	**Good identification**					**03771160**
Actinomyces naeslundii		94.99	0.64	MNS 88		
Actinomyces israelii		2.22	0.27			
Clostridium perfringens		0.88	0.07			
03771161	**Very good identification**					**03771161**
Actinomyces naeslundii		100.0	0.67	MNS 88		
03771162	**Low discrimination**					**03771162**
Actinomyces naeslundii		52.64	0.64	MNS 88		
Lactobacillus acidophilus		40.59	0.38			
Bifidobacterium breve		3.07	0.29			
03771163	**Very good identification**					**03771163**
Actinomyces naeslundii		99.48	0.67	MNS 88		
Lactobacillus acidophilus		0.52	0.04			
03771170	**Acceptable identification**					**03771170**
Actinomyces naeslundii		90.13	0.43	MNS 88		
Bifidobacterium breve		9.87	0.12	MLZ 5		
03771171	**Good identification**					**03771171**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03771173	**Good identification**					**03771173**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03771200	**Low discrimination**					**03771200**
					GRA	
Actinomyces naeslundii		41.63	0.35	MNS 88	99	
Actinomyces israelii		36.59	0.25	RHA 1	99	
Fusobacterium mortiferum		21.78	0.19		1	
03771201	**Good identification**					**03771201**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771203	**Good identification**					**03771203**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771220	**Low discrimination**					**03771220**
					SPO	
Actinomyces naeslundii		69.57	0.35	MNS 88	1	
Clostridium ramosum		30.43	0.04	RHA 1	92	
03771221	**Good identification**					**03771221**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	MNS 88			
03771223	**Good identification**						**03771223**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03771300	**Low discrimination**						**03771300**
					GRA		
Actinomyces israelii		42.59	0.35	XYL 92	99		
Actinomyces naeslundii		34.47	0.43	RHA 7	99		
Fusobacterium mortiferum		14.79	0.25		1		
03771301	**Good identification**						**03771301**
Actinomyces naeslundii		100.0	0.46	MNS 88			
03771303	**Acceptable identification**						**03771303**
Actinomyces naeslundii		94.77	0.46	MNS 88			
Lactobacillus acidophilus		5.23	0.00	RHA 1			
03771320	**Acceptable identification**						**03771320**
					SPO	GRA	
Clostridium ramosum		91.59	0.38	MNS 99	92	50	
Actinomyces naeslundii		6.02	0.43		1	99	
Prevotella oralis		1.04	0.04		1	1	
03771321	**Acceptable identification**						**03771321**
					SPO		
Actinomyces naeslundii		90.71	0.46	MNS 88	1		
Clostridium ramosum		9.30	0.04	RHA 1	92		
03771322	**Low discrimination**						**03771322**
					SPO	GRA	
Clostridium ramosum		76.18	0.48	MNS 99	92	50	
Lactobacillus acidophilus		10.27	0.34		1	99	
Parabacteroides distasonis		7.94	0.44		1	1	
03771323	**Low discrimination**						**03771323**
					SPO		
Actinomyces naeslundii		66.32	0.46	MNS 88	1		
Clostridium ramosum		27.19	0.14	RHA 1	92		
Lactobacillus acidophilus		3.66	0.00		1		
03771332	**Low discrimination**						**03771332**
					SPO		
Parabacteroides distasonis		74.13	0.39	MNS 98	1		
Clostridium ramosum		13.35	0.14	XYL 90	92		
Clostridium clostridioforme		10.72	0.19		66		
03771400	**Low discrimination**						**03771400**
					SPO		
Actinomyces israelii		81.65	0.43	XYL 92	1		
Actinomyces naeslundii		8.55	0.35		1		
Clostridium butyricum		5.34	0.02		99		
03771401	**Good identification**						**03771401**
Actinomyces naeslundii		100.0	0.38	MNS 88			
03771402	**Low discrimination**						**03771402**
Actinomyces israelii		81.75	0.46	XYL 92			
Bifidobacterium adolescentis		8.48	0.23				

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		5.70	0.35			
03771403	**Good identification**					**03771403**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771420	**Low discrimination**					**03771420**
Actinomyces naeslundii		62.12	0.35	MNS 88		
Actinomyces israelii		37.88	0.22	ARA 1		
03771421	**Good identification**					**03771421**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771422	**Low discrimination**					**03771422**
Actinomyces naeslundii		41.94	0.35	MNS 88		
Actinomyces israelii		38.37	0.25	ARA 1		
Lactobacillus acidophilus		11.43	0.01			
03771423	**Good identification**					**03771423**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03771440	**Good identification**					**03771440**
Actinomyces israelii		95.47	0.36	XYL 92		
Actinomyces naeslundii		4.53	0.22			
03771442	**Acceptable identification**					**03771442**
Actinomyces israelii		93.93	0.39	XYL 92		
Bifidobacterium adolescentis		3.10	0.07			
Actinomyces naeslundii		2.97	0.22			
03771500	**Low discrimination**					**03771500**
					SPO	
Actinomyces israelii		65.03	0.53	XYL 92	1	
Clostridium butyricum		25.52	0.25		99	
Actinomyces naeslundii		4.84	0.43		1	
03771501	**Acceptable identification**					**03771501**
Actinomyces naeslundii		91.71	0.46	MNS 88		
Actinomyces israelii		8.29	0.19	ARA 1		
03771502	**Low discrimination**					**03771502**
					SPO	
Bifidobacterium adolescentis		67.84	0.57	MNS 87	1	
Actinomyces israelii		26.42	0.56	XYL 87	1	
Clostridium butyricum		2.69	0.18		99	
03771503	**Low discrimination**					**03771503**
Actinomyces naeslundii		67.39	0.46	MNS 88		
Bifidobacterium adolescentis		23.47	0.23	ARA 1		
Actinomyces israelii		9.14	0.22			
03771510	**Low discrimination**					**03771510**
					SPO	
Actinomyces israelii		68.66	0.39	XYL 92	1	
Bifidobacterium adolescentis		21.85	0.24	MLZ 13	1	
Clostridium butyricum		9.49	0.03		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03771512	**Good identification**					**03771512**
Bifidobacterium adolescentis		93.71	0.58	MNS 87		
Actinomyces israelii		4.46	0.42	XYL 87		
Clostridium clostridioforme		1.56	0.25			
03771520	**Low discrimination**					**03771520**
					SPO	
Actinomyces naeslundii		41.13	0.43	MNS 88	1	
Actinomyces israelii		35.25	0.33	ARA 1	1	
Clostridium butyricum		18.85	0.07		99	
03771521	**Good identification**					**03771521**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03771523	**Good identification**					**03771523**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03771540	**Low discrimination**					**03771540**
					SPO	
Actinomyces israelii		88.19	0.46	XYL 92	1	
Clostridium butyricum		7.74	0.07		99	
Actinomyces naeslundii		2.98	0.30		1	
03771542	**Low discrimination**					**03771542**
					SPO	
Actinomyces israelii		53.23	0.49	XYL 92	1	
Bifidobacterium adolescentis		43.44	0.41		1	
Clostridium butyricum		1.21	0.00		99	
03771552	**Low discrimination**					**03771552**
Bifidobacterium adolescentis		86.97	0.43	MNS 87		
Actinomyces israelii		13.03	0.35	XYL 87		
03771712	**Low discrimination**					**03771712**
Clostridium clostridioforme		89.94	0.40	MNS 99		
Bifidobacterium adolescentis		7.42	0.24	XYL 91		
Actinomyces israelii		2.63	0.23			
03771722	**Low discrimination**					**03771722**
					SPO	
Parabacteroides distasonis		60.92	0.40	MNS 98	1	
Clostridium clostridioforme		26.63	0.28	XYL 90	66	
Clostridium ramosum		9.64	0.14		92	
03771732	**Low discrimination**					**03771732**
Clostridium clostridioforme		78.71	0.40	MNS 99		
Parabacteroides distasonis		21.29	0.36	XYL 91		
03772000	**Low discrimination**					**03772000**
					GRA	
Actinomyces naeslundii		88.73	0.51	MNS 88	99	
Actinomyces israelii		7.41	0.23	GAL 96	99	
Fusobacterium mortiferum		3.86	0.16		1	
03772001	**Good identification**					**03772001**
Actinomyces naeslundii		100.0	0.54	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03772002	**Low discrimination**					**03772002**
Actinomyces naeslundii		88.87	0.51	MNS 88		
Actinomyces israelii		11.13	0.26	GAL 96		
03772003	**Good identification**					**03772003**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03772020	**Good identification**					**03772020**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03772021	**Good identification**					**03772021**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03772022	**Good identification**					**03772022**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03772023	**Good identification**					**03772023**
Actinomyces naeslundii		100.0	0.54	MNS 88		
03772040	**Low discrimination**					**03772040**
Actinomyces naeslundii		84.47	0.38	MNS 88		
Actinomyces israelii		15.54	0.16	GAL 96		
03772041	**Good identification**					**03772041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03772042	**Low discrimination**					**03772042**
Actinomyces naeslundii		78.37	0.38	MNS 88		
Actinomyces israelii		21.63	0.19	GAL 96		
03772043	**Good identification**					**03772043**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03772060	**Good identification**					**03772060**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03772061	**Good identification**					**03772061**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03772062	**Good identification**					**03772062**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03772063	**Good identification**					**03772063**
Actinomyces naeslundii		100.0	0.41	MNS 88		
03772100	**Low discrimination**					**03772100**
					GRA	
Actinomyces naeslundii		86.73	0.59	MNS 88	99	
Actinomyces israelii		10.18	0.33	GAL 96	99	
Fusobacterium mortiferum		3.10	0.22		1	
03772101	**Very good identification**					**03772101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.62	MNS 88		
03772102	**Low discrimination**					**03772102**
Actinomyces naeslundii		83.79	0.59	MNS 88		
Actinomyces israelii		14.75	0.36	GAL 96		
Propionibacterium propionicum		0.79	0.14			
03772103	**Very good identification**					**03772103**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03772110	**Low discrimination**					**03772110**
Actinomyces naeslundii		75.02	0.37	MNS 88		
Actinomyces israelii		24.99	0.19	GAL 96		
03772111	**Good identification**					**03772111**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03772112	**Low discrimination**					**03772112**
Actinomyces naeslundii		59.35	0.37	MNS 88		
Actinomyces israelii		29.65	0.22	GAL 96		
Bifidobacterium adolescentis		10.99	0.09	MLZ 5		
03772113	**Good identification**					**03772113**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03772120	**Good identification**					**03772120**
					SPO	
Actinomyces naeslundii		92.42	0.59	MNS 88	1	
Clostridium ramosum		7.58	0.15	GAL 96	92	
03772121	**Very good identification**					**03772121**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03772122	**Low discrimination**					**03772122**
					SPO	
Actinomyces naeslundii		74.66	0.59	MNS 88	1	
Clostridium ramosum		24.51	0.25	GAL 96	92	
Actinomyces israelii		0.84	0.16		1	
03772123	**Very good identification**					**03772123**
Actinomyces naeslundii		100.0	0.62	MNS 88		
03772130	**Good identification**					**03772130**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03772131	**Good identification**					**03772131**
Actinomyces naeslundii		100.0	0.40	MNS 88		
03772132	**Good identification**					**03772132**
Actinomyces naeslundii		100.0	0.37	MNS 88		
03772133	**Good identification**					**03772133**
Actinomyces naeslundii		100.0	0.40	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03772140	**Low discrimination**					**03772140**
Actinomyces naeslundii		79.46	0.46	MNS 88		
Actinomyces israelii		20.54	0.26	GAL 96		
03772141	**Good identification**					**03772141**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03772142	**Low discrimination**					**03772142**
Actinomyces naeslundii		69.37	0.46	MNS 88		
Actinomyces israelii		26.90	0.29	GAL 96		
Propionibacterium propionicum		3.72	0.14			
03772143	**Good identification**					**03772143**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03772160	**Good identification**					**03772160**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03772161	**Good identification**					**03772161**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03772162	**Good identification**					**03772162**
Actinomyces naeslundii		100.0	0.46	MNS 88		
03772163	**Good identification**					**03772163**
Actinomyces naeslundii		100.0	0.49	MNS 88		
03773000	**Low discrimination**					**03773000**
					GRA	
Actinomyces naeslundii		74.24	0.74	MNS 88	99	
Actinomyces israelii		25.56	0.57		99	
Fusobacterium mortiferum		0.20	0.19		1	
03773001	**Very good identification**					**03773001**
Actinomyces naeslundii		99.77	0.77	MNS 88		
Actinomyces israelii		0.23	0.23			
03773002	**Low discrimination**					**03773002**
Actinomyces naeslundii		65.58	0.74	MNS 88		
Actinomyces israelii		33.87	0.60			
Lactobacillus acidophilus		0.48	0.14			
03773003	**Very good identification**					**03773003**
Actinomyces naeslundii		99.65	0.77	MNS 88		
Actinomyces israelii		0.35	0.26			
03773010	**Low discrimination**					**03773010**
Actinomyces naeslundii		50.57	0.53	MNS 88		
Actinomyces israelii		49.43	0.43	MLZ 5		
03773011	**Good identification**					**03773011**
Actinomyces naeslundii		100.0	0.56	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03773012	**Low discrimination**					**03773012**
Actinomyces israelii		58.93	0.46	XYL 92		
Actinomyces naeslundii		40.19	0.53	MLZ 13		
Bifidobacterium adolescentis		0.88	0.09			
03773013	**Good identification**					**03773013**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03773020	**Very good identification**					**03773020**
					SPO	
Actinomyces naeslundii		96.93	0.74	MNS 88	1	
Actinomyces israelii		2.13	0.37		1	
Clostridium ramosum		0.94	0.15		92	
03773021	**Very good identification**					**03773021**
Actinomyces naeslundii		100.0	0.77	MNS 88		
03773022	**Good identification**					**03773022**
					SPO	
Actinomyces naeslundii		92.65	0.74	MNS 88	1	
Clostridium ramosum		3.61	0.25		92	
Actinomyces israelii		3.06	0.40		1	
03773023	**Very good identification**					**03773023**
Actinomyces naeslundii		100.0	0.77	MNS 88		
03773030	**Acceptable identification**					**03773030**
Actinomyces naeslundii		94.13	0.53	MNS 88		
Actinomyces israelii		5.87	0.23	MLZ 5		
03773031	**Good identification**					**03773031**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03773032	**Acceptable identification**					**03773032**
Actinomyces naeslundii		91.44	0.53	MNS 88		
Actinomyces israelii		8.56	0.26	MLZ 5		
03773033	**Good identification**					**03773033**
Actinomyces naeslundii		100.0	0.56	MNS 88		
03773040	**Low discrimination**					**03773040**
Actinomyces naeslundii		56.86	0.62	MNS 88		
Actinomyces israelii		43.14	0.50			
03773041	**Very good identification**					**03773041**
Actinomyces naeslundii		99.49	0.65	MNS 88		
Actinomyces israelii		0.51	0.16			
03773042	**Low discrimination**					**03773042**
Actinomyces israelii		53.11	0.53	XYL 92		
Actinomyces naeslundii		46.65	0.62			
Lactobacillus acidophilus		0.24	-.02			
03773043	**Very good identification**					**03773043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.24	0.65	MNS 88		
Actinomyces israelii		0.76	0.19			
03773050	**Low discrimination**					**03773050**
Actinomyces israelii		68.30	0.36	XYL 92		
Actinomyces naeslundii		31.70	0.40	MLZ 13		
03773051	**Good identification**					**03773051**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773052	**Low discrimination**					**03773052**
Actinomyces israelii		76.37	0.39	XYL 92		
Actinomyces naeslundii		23.63	0.40	MLZ 13		
03773053	**Good identification**					**03773053**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773060	**Very good identification**					**03773060**
Actinomyces naeslundii		95.38	0.62	MNS 88		
Actinomyces israelii		4.62	0.30			
03773061	**Very good identification**					**03773061**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03773062	**Good identification**					**03773062**
Actinomyces naeslundii		92.78	0.62	MNS 88		
Actinomyces israelii		6.74	0.33			
Lactobacillus acidophilus		0.48	-.02			
03773063	**Very good identification**					**03773063**
Actinomyces naeslundii		100.0	0.65	MNS 88		
03773070	**Low discrimination**					**03773070**
Actinomyces naeslundii		87.91	0.40	MNS 88		
Actinomyces israelii		12.09	0.16	MLZ 5		
03773071	**Good identification**					**03773071**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773072	**Low discrimination**					**03773072**
Actinomyces naeslundii		82.90	0.40	MNS 88		
Actinomyces israelii		17.10	0.19	MLZ 5		
03773073	**Good identification**					**03773073**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773100	**Low discrimination**					**03773100**
					SPO	
Actinomyces naeslundii		67.08	0.82	MNS 88	1	
Actinomyces israelii		32.47	0.68		1	
Clostridium ramosum		0.23	0.15		92	
03773101	**Very good identification**					**03773101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.68	0.85	MNS 88		
Actinomyces israelii		0.33	0.33			
03773102	**Low discrimination**					**03773102**
Actinomyces naeslundii		55.34	0.82	MNS 88		
Actinomyces israelii		40.17	0.71			
Bifidobacterium adolescentis		1.82	0.41			
03773103	**Very good identification**					**03773103**
Actinomyces naeslundii		99.49	0.85	MNS 88		
Actinomyces israelii		0.49	0.36			
Bifidobacterium adolescentis		0.02	0.07			
03773110	**Low discrimination**					**03773110**
					SPO	
Actinomyces israelii		57.36	0.53	XYL 92	1	
Actinomyces naeslundii		41.75	0.60	MLZ 13	1	
Clostridium tertium		0.57	0.05		99	
03773111	**Very good identification**					**03773111**
Actinomyces naeslundii		99.08	0.63	MNS 88		
Actinomyces israelii		0.92	0.19	MLZ 5		
03773112	**Low discrimination**					**03773112**
Actinomyces israelii		52.45	0.57	XYL 92		
Actinomyces naeslundii		25.45	0.60	MLZ 13		
Bifidobacterium adolescentis		19.45	0.43			
03773113	**Very good identification**					**03773113**
Actinomyces naeslundii		98.13	0.63	MNS 88		
Actinomyces israelii		1.36	0.22	MLZ 5		
Bifidobacterium adolescentis		0.51	0.09			
03773120	**Low discrimination**					**03773120**
					SPO	
Actinomyces naeslundii		72.93	0.82	MNS 88	1	
Clostridium ramosum		24.69	0.49		92	
Actinomyces israelii		2.25	0.47		1	
03773121	**Very good identification**					**03773121**
					SPO	
Actinomyces naeslundii		99.77	0.85	MNS 88	1	
Clostridium ramosum		0.23	0.15		92	
03773122	**Low discrimination**					**03773122**
					SPO	
Clostridium ramosum		55.50	0.59	MNS 99	92	
Actinomyces naeslundii		40.99	0.82		1	
Actinomyces israelii		1.90	0.50		1	
03773123	**Very good identification**					**03773123**
					SPO	
Actinomyces naeslundii		99.07	0.85	MNS 88	1	
Clostridium ramosum		0.90	0.25		92	
Actinomyces israelii		0.03	0.16		1	
03773130	**Low discrimination**					**03773130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		83.09	0.60	MNS 88	1	
Actinomyces israelii		7.29	0.33	MLZ 5	1	
Clostridium ramosum		5.40	0.15		92	
03773131	**Very good identification**					**03773131**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03773132	**Low discrimination**					**03773132**
					SPO	
Actinomyces naeslundii		61.64	0.60	MNS 88	1	
Clostridium ramosum		16.02	0.25	MLZ 5	92	
Actinomyces israelii		8.11	0.36		1	
03773133	**Very good identification**					**03773133**
Actinomyces naeslundii		100.0	0.63	MNS 88		
03773140	**Low discrimination**					**03773140**
Actinomyces israelii		51.61	0.61	XYL 92		
Actinomyces naeslundii		48.39	0.69			
03773141	**Very good identification**					**03773141**
Actinomyces naeslundii		99.29	0.72	MNS 88		
Actinomyces israelii		0.71	0.26			
03773142	**Low discrimination**					**03773142**
Actinomyces israelii		60.08	0.64	XYL 92		
Actinomyces naeslundii		37.56	0.69			
Bifidobacterium adolescentis		0.87	0.26			
03773143	**Very good identification**					**03773143**
Actinomyces naeslundii		98.93	0.72	MNS 88		
Actinomyces israelii		1.07	0.29			
03773150	**Low discrimination**					**03773150**
Actinomyces israelii		75.17	0.46	XYL 92		
Actinomyces naeslundii		24.83	0.47	MLZ 13		
03773151	**Good identification**					**03773151**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03773152	**Low discrimination**					**03773152**
Actinomyces israelii		70.99	0.49	XYL 92		
Actinomyces naeslundii		15.63	0.47	MLZ 13		
Bifidobacterium adolescentis		8.37	0.27			
03773153	**Good identification**					**03773153**
Actinomyces naeslundii		97.03	0.50	MNS 88		
Actinomyces israelii		2.97	0.15	MLZ 5		
03773160	**Good identification**					**03773160**
					SPO	
Actinomyces naeslundii		91.71	0.69	MNS 88	1	
Actinomyces israelii		6.24	0.40		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		1.78	0.15		92	
03773161	**Very good identification**					**03773161**
Actinomyces naeslundii		100.0	0.72	MNS 88		
03773162	**Low discrimination**					**03773162**
					SPO	
Actinomyces naeslundii		81.27	0.69	MNS 88	1	
Actinomyces israelii		8.30	0.43		1	
Clostridium ramosum		6.30	0.25		92	
03773163	**Very good identification**					**03773163**
Actinomyces naeslundii		100.0	0.72	MNS 88		
03773170	**Low discrimination**					**03773170**
Actinomyces naeslundii		80.02	0.47	MNS 88		
Actinomyces israelii		15.47	0.26	MLZ 5		
Bifidobacterium breve		4.51	0.12			
03773171	**Good identification**					**03773171**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03773172	**Low discrimination**					**03773172**
Actinomyces naeslundii		53.76	0.47	MNS 88		
Bifidobacterium breve		30.66	0.29	MLZ 5		
Actinomyces israelii		15.59	0.29			
03773173	**Good identification**					**03773173**
Actinomyces naeslundii		100.0	0.50	MNS 88		
03773200	**Low discrimination**					**03773200**
Actinomyces israelii		71.96	0.38	XYL 92		
Actinomyces naeslundii		28.04	0.40	RHA 7		
03773201	**Good identification**					**03773201**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773202	**Low discrimination**					**03773202**
Actinomyces israelii		79.38	0.41	XYL 92		
Actinomyces naeslundii		20.62	0.40	RHA 7		
03773203	**Good identification**					**03773203**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773221	**Good identification**					**03773221**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773223	**Good identification**					**03773223**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773300	**Low discrimination**					**03773300**
					SPO	
Actinomyces israelii		71.34	0.48	XYL 92	1	
Actinomyces naeslundii		19.78	0.48	RHA 7	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		8.88	0.17		92	
03773301	**Good identification**					**03773301**
Actinomyces naeslundii		97.63	0.51	MNS 88		
Actinomyces israelii		2.37	0.14	RHA 1		
03773302	**Low discrimination**					**03773302**
					SPO	
Actinomyces israelii		63.94	0.51	XYL 92	1	
Clostridium ramosum		21.22	0.27	RHA 7	92	
Actinomyces naeslundii		11.82	0.48		1	
03773303	**Good identification**					**03773303**
Actinomyces naeslundii		96.49	0.51	MNS 88		
Actinomyces israelii		3.51	0.17	RHA 1		
03773312	**Low discrimination**					**03773312**
Actinomyces israelii		89.70	0.37	XYL 92		
Actinomyces naeslundii		5.84	0.26	RHA 7		
Bifidobacterium adolescentis		4.46	0.09	MLZ 13		
03773320	**Good identification**					**03773320**
					SPO	
Clostridium ramosum		97.31	0.51	MNS 99	92	
Actinomyces naeslundii		2.19	0.48		1	
Actinomyces israelii		0.50	0.28		1	
03773321	**Low discrimination**					**03773321**
					SPO	
Actinomyces naeslundii		76.97	0.51	MNS 88	1	
Clostridium ramosum		23.03	0.17	RHA 1	92	
03773322	**Very good identification**					**03773322**
					SPO	
Clostridium ramosum		99.11	0.61	MNS 99	92	
Actinomyces naeslundii		0.56	0.48		1	
Actinomyces israelii		0.19	0.31		1	
03773340	**Low discrimination**					**03773340**
Actinomyces israelii		88.82	0.41	XYL 92		
Actinomyces naeslundii		11.18	0.35	RHA 7		
03773341	**Good identification**					**03773341**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03773342	**Acceptable identification**					**03773342**
Actinomyces israelii		92.26	0.44	XYL 92		
Actinomyces naeslundii		7.74	0.35	RHA 7		
03773343	**Good identification**					**03773343**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03773361	**Good identification**					**03773361**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03773363	**Good identification**					**03773363**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
03773400	**Good identification**					**03773400**
Actinomyces israelii		96.54	0.56	XYL 92		
Actinomyces naeslundii		3.46	0.40			
03773401	**Low discrimination**					**03773401**
Actinomyces naeslundii		84.19	0.43	MNS 88		
Actinomyces israelii		15.81	0.22	ARA 1		
03773402	**Very good identification**					**03773402**
Actinomyces israelii		96.26	0.59	XYL 92		
Actinomyces naeslundii		2.30	0.40			
Bifidobacterium adolescentis		1.44	0.21			
03773403	**Low discrimination**					**03773403**
Actinomyces naeslundii		78.02	0.43	MNS 88		
Actinomyces israelii		21.98	0.25	ARA 1		
03773410	**Good identification**					**03773410**
Actinomyces israelii		100.0	0.42	XYL 92		
03773412	**Low discrimination**					**03773412**
Actinomyces israelii		89.08	0.45	XYL 92		
Bifidobacterium adolescentis		10.92	0.23	MLZ 13		
03773420	**Low discrimination**					**03773420**
Actinomyces israelii		64.03	0.35	XYL 92		
Actinomyces naeslundii		35.97	0.40	NAG 6		
03773421	**Good identification**					**03773421**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773422	**Low discrimination**					**03773422**
Actinomyces israelii		72.76	0.38	XYL 92		
Actinomyces naeslundii		27.25	0.40	NAG 6		
03773423	**Good identification**					**03773423**
Actinomyces naeslundii		100.0	0.43	MNS 88		
03773440	**Good identification**					**03773440**
Actinomyces israelii		98.40	0.49	XYL 92		
Actinomyces naeslundii		1.60	0.27			
03773442	**Good identification**					**03773442**
Actinomyces israelii		98.93	0.52	XYL 92		
Actinomyces naeslundii		1.07	0.27			
03773452	**Good identification**					**03773452**
Actinomyces israelii		96.25	0.38	XYL 92		
Bifidobacterium adolescentis		3.75	0.07	MLZ 13		
03773500	**Very good identification**					**03773500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		96.26	0.66	XYL 92	1	
Actinomyces naeslundii		2.46	0.48		1	
Clostridium butyricum		0.74	0.09		99	
03773501	**Low discrimination**					**03773501**
Actinomyces naeslundii		79.12	0.51	MNS 88		
Actinomyces israelii		20.89	0.32	ARA 1		
03773502	**Low discrimination**					**03773502**
Actinomyces israelii		71.95	0.69	XYL 92		
Bifidobacterium adolescentis		26.68	0.55			
Actinomyces naeslundii		1.22	0.48			
03773503	**Low discrimination**					**03773503**
Actinomyces naeslundii		64.82	0.51	MNS 88		
Actinomyces israelii		25.67	0.35	ARA 1		
Bifidobacterium adolescentis		9.52	0.21			
03773510	**Acceptable identification**					**03773510**
Actinomyces israelii		94.79	0.52	XYL 92		
Bifidobacterium adolescentis		4.36	0.23	MLZ 13		
Actinomyces naeslundii		0.85	0.26			
03773512	**Low discrimination**					**03773512**
Bifidobacterium adolescentis		74.97	0.57	MNS 87		
Actinomyces israelii		24.72	0.55	XYL 87		
Bifidobacterium dentium		0.16	-.07			
03773520	**Low discrimination**					**03773520**
					SPO	
Actinomyces israelii		65.15	0.46	XYL 92	1	
Actinomyces naeslundii		26.04	0.48	NAG 6	1	
Clostridium ramosum		8.81	0.15		92	
03773521	**Good identification**					**03773521**
Actinomyces naeslundii		100.0	0.51	MNS 88		
03773522	**Low discrimination**					**03773522**
					SPO	
Actinomyces israelii		59.32	0.49	XYL 92	1	
Clostridium ramosum		21.40	0.25	NAG 6	92	
Actinomyces naeslundii		15.80	0.48		1	
03773523	**Good identification**					**03773523**
Actinomyces naeslundii		97.54	0.51	MNS 88		
Actinomyces israelii		2.46	0.15	ARA 1		
03773540	**Very good identification**					**03773540**
Actinomyces israelii		98.86	0.59	XYL 92		
Actinomyces naeslundii		1.14	0.35			
03773541	**Low discrimination**					**03773541**
Actinomyces naeslundii		63.22	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		36.78	0.25	ARA 1		
03773542	**Low discrimination**					**03773542**
Actinomyces israelii		88.85	0.62	XYL 92		
Bifidobacterium adolescentis		10.47	0.40			
Actinomyces naeslundii		0.69	0.35			
03773543	**Low discrimination**					**03773543**
Actinomyces naeslundii		53.40	0.38	MNS 88		
Actinomyces israelii		46.60	0.28	ARA 1		
03773550	**Good identification**					**03773550**
					COC	
Actinomyces israelii		96.21	0.45	XYL 92	1	
Blautia producta		2.38	-.03	MLZ 13	99	
Bifidobacterium adolescentis		1.41	0.07		1	
03773552	**Low discrimination**					**03773552**
Actinomyces israelii		50.92	0.48	XYL 92		
Bifidobacterium adolescentis		49.08	0.41	MLZ 13		
03773560	**Low discrimination**					**03773560**
Actinomyces israelii		84.65	0.39	XYL 92		
Actinomyces naeslundii		15.35	0.35	NAG 6		
03773561	**Good identification**					**03773561**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03773562	**Low discrimination**					**03773562**
Actinomyces israelii		89.21	0.42	XYL 92		
Actinomyces naeslundii		10.79	0.35	NAG 6		
03773563	**Good identification**					**03773563**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03773600	**Good identification**					**03773600**
Actinomyces israelii		100.0	0.37	XYL 92		
03773602	**Good identification**					**03773602**
Actinomyces israelii		100.0	0.40	XYL 92		
03773700	**Good identification**					**03773700**
Actinomyces israelii		100.0	0.47	XYL 92		
03773702	**Good identification**					**03773702**
Actinomyces israelii		95.26	0.50	XYL 92		
Bifidobacterium adolescentis		4.74	0.21	RHA 7		
03773712	**Low discrimination**					**03773712**
Actinomyces israelii		68.13	0.36	XYL 92		
Bifidobacterium adolescentis		27.73	0.23	RHA 7		
Clostridium clostridioforme		4.15	0.06	MLZ 13		
03773740	**Good identification**					**03773740**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	XYL 92			
03773742	**Good identification**						**03773742**
Actinomyces israelii		100.0	0.43	XYL 92			
03774100	**Low discrimination**						**03774100**
					GRA	SPO	
Fusobacterium mortiferum		50.39	0.37	MLT 11	1	1	
Actinomyces israelii		45.09	0.39	XYL 7	99	1	
Clostridium butyricum		4.52	0.01		99	99	
03774102	**Low discrimination**						**03774102**
					GRA		
Actinomyces israelii		77.43	0.42	GAL 99	99		
Bifidobacterium adolescentis		14.15	0.23		99		
Fusobacterium mortiferum		6.41	0.21		1		
03774502	**Low discrimination**						**03774502**
Bifidobacterium adolescentis		47.99	0.37	MNS 87			
Actinomyces israelii		32.11	0.40	GAL 99			
Clostridium clostridioforme		10.71	0.23				
03774512	**Low discrimination**						**03774512**
Bifidobacterium adolescentis		52.25	0.38	MNS 87			
Clostridium clostridioforme		43.48	0.34	GAL 99			
Actinomyces israelii		4.27	0.26				
03774612	**Good identification**						**03774612**
Clostridium clostridioforme		100.0	0.40	MNS 99			
03774632	**Good identification**						**03774632**
Clostridium clostridioforme		100.0	0.40	MNS 99			
03774702	**Good identification**						**03774702**
Clostridium clostridioforme		97.02	0.37	MNS 99			
Actinomyces israelii		2.99	0.21				
03774712	**Good identification**						**03774712**
Clostridium clostridioforme		100.0	0.49	MNS 99			
03774722	**Acceptable identification**						**03774722**
Clostridium clostridioforme		94.19	0.37	MNS 99			
Parabacteroides distasonis		5.81	0.22				
03774732	**Good identification**						**03774732**
Clostridium clostridioforme		99.28	0.49	MNS 99			
Parabacteroides distasonis		0.72	0.18				
03775000	**Good identification**						**03775000**
					GRA	SPO	
Actinomyces israelii		94.95	0.63		99	1	
Fusobacterium mortiferum		2.76	0.34		1	1	
Clostridium butyricum		1.59	0.11		99	99	
03775001	**Low discrimination**						**03775001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		52.51	0.38	MNS 88		
Actinomyces israelii		47.49	0.28	XYL 1		
03775002	**Very good identification**					**03775002**
Actinomyces israelii		98.02	0.66			
Bifidobacterium adolescentis		0.72	0.23			
Actinomyces naeslundii		0.49	0.35			
03775010	**Good identification**					**03775010**
Actinomyces israelii		100.0	0.48	MLZ 13		
03775012	**Acceptable identification**					**03775012**
Actinomyces israelii		92.63	0.51	MLZ 13		
Bifidobacterium adolescentis		5.59	0.24			
Clostridium clostridioforme		1.78	0.13			
03775020	**Low discrimination**					**03775020**
					SPO	
Actinomyces israelii		87.76	0.42	NAG 6	1	
Actinomyces naeslundii		10.24	0.35		1	
Clostridium butyricum		2.00	-.07		99	
03775021	**Good identification**					**03775021**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03775022	**Low discrimination**					**03775022**
					GRA	
Actinomyces israelii		87.61	0.45	NAG 6	99	
Actinomyces naeslundii		6.81	0.35		99	
Parabacteroides distasonis		3.73	0.18		1	
03775023	**Good identification**					**03775023**
Actinomyces naeslundii		100.0	0.38	MNS 88		
03775040	**Good identification**					**03775040**
					SPO	
Actinomyces israelii		99.29	0.56		1	
Clostridium butyricum		0.37	-.07		99	
Actinomyces naeslundii		0.34	0.22		1	
03775042	**Very good identification**					**03775042**
Actinomyces israelii		99.54	0.59			
Bifidobacterium adolescentis		0.23	0.07			
Actinomyces naeslundii		0.22	0.22			
03775050	**Good identification**					**03775050**
Actinomyces israelii		100.0	0.41	MLZ 13		
03775052	**Good identification**					**03775052**
Actinomyces israelii		98.12	0.44	MLZ 13		
Bifidobacterium adolescentis		1.88	0.09			
03775060	**Acceptable identification**					**03775060**
Actinomyces israelii		94.97	0.35	NAG 6		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		5.03	0.22				
03775062	**Good identification**						**03775062**
Actinomyces israelii		96.59	0.38	NAG 6			
Actinomyces naeslundii		3.41	0.22				
03775100	**Low discrimination**						**03775100**
					GRA	SPO	
Actinomyces israelii		88.10	0.73		99	1	
Clostridium butyricum		8.84	0.35		99	99	
Fusobacterium mortiferum		1.49	0.40		1	1	
03775101	**Low discrimination**						**03775101**
					SPO		
Actinomyces israelii		52.99	0.39	URE 1	1		
Actinomyces naeslundii		41.69	0.46		1		
Clostridium butyricum		5.31	0.01		99		
03775102	**Low discrimination**						**03775102**
					SPO		
Actinomyces israelii		81.99	0.76		1		
Bifidobacterium adolescentis		14.98	0.57		1		
Clostridium butyricum		2.13	0.28		99		
03775103	**Low discrimination**						**03775103**
Actinomyces israelii		57.70	0.42	URE 1			
Actinomyces naeslundii		30.26	0.46				
Bifidobacterium adolescentis		10.54	0.23				
03775110	**Good identification**						**03775110**
					SPO		
Actinomyces israelii		94.52	0.59	MLZ 13	1		
Clostridium butyricum		3.34	0.13		99		
Bifidobacterium adolescentis		2.14	0.24		1		
03775112	**Low discrimination**						**03775112**
Bifidobacterium adolescentis		59.21	0.58	MNS 87			
Actinomyces israelii		39.62	0.62	ARA 87			
Clostridium clostridioforme		0.64	0.22				
03775120	**Low discrimination**						**03775120**
					SPO		
Actinomyces israelii		78.59	0.52	NAG 6	1		
Clostridium butyricum		10.74	0.17		99		
Actinomyces naeslundii		6.52	0.43		1		
03775121	**Acceptable identification**						**03775121**
Actinomyces naeslundii		92.50	0.46	MNS 88			
Actinomyces israelii		7.50	0.18	XYL 1			
03775122	**Low discrimination**						**03775122**
					GRA		
Parabacteroides distasonis		45.39	0.52	MNS 98	1		
Actinomyces israelii		43.12	0.55		99		
Lactobacillus acidophilus		2.63	0.19		99		
03775123	**Low discrimination**						**03775123**
					GRA		
Actinomyces naeslundii		80.01	0.46	MNS 88	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		10.25	0.18	XYL 1	1		
Actinomyces israelii		9.74	0.21		99		
03775130	**Acceptable identification**						**03775130**
					GRA	SPO	
Actinomyces israelii		90.45	0.38	MLZ 13	99	1	
Parabacteroides distasonis		5.20	0.13	NAG 6	1	1	
Clostridium butyricum		4.35	-.05		99	99	
03775132	**Low discrimination**						**03775132**
					GRA		
Parabacteroides distasonis		71.82	0.47	MNS 98	1		
Actinomyces israelii		18.93	0.41		99		
Clostridium clostridioforme		4.78	0.22		50		
03775140	**Very good identification**						**03775140**
					SPO		
Actinomyces israelii		97.49	0.66		1		
Clostridium butyricum		2.19	0.17		99		
Actinomyces naeslundii		0.23	0.30		1		
03775142	**Good identification**						**03775142**
					SPO		
Actinomyces israelii		93.75	0.69		1		
Bifidobacterium adolescentis		5.44	0.41		1		
Clostridium butyricum		0.55	0.10		99		
03775150	**Good identification**						**03775150**
					SPO		
Actinomyces israelii		98.51	0.52	MLZ 13	1		
Clostridium butyricum		0.78	-.05		99		
Bifidobacterium adolescentis		0.71	0.09		1		
03775152	**Low discrimination**						**03775152**
Actinomyces israelii		67.80	0.55	MLZ 13			
Bifidobacterium adolescentis		32.20	0.43				
03775160	**Acceptable identification**						**03775160**
					SPO		
Actinomyces israelii		93.61	0.45	NAG 6	1		
Actinomyces naeslundii		3.53	0.30		1		
Clostridium butyricum		2.86	-.02		99		
03775162	**Acceptable identification**						**03775162**
					GRA		
Actinomyces israelii		92.51	0.48	NAG 6	99		
Parabacteroides distasonis		2.53	0.18		1		
Actinomyces naeslundii		2.32	0.30		99		
03775200	**Good identification**						**03775200**
Actinomyces israelii		100.0	0.43	RHA 7			
03775202	**Acceptable identification**						**03775202**
Actinomyces israelii		94.21	0.46	RHA 7			
Clostridium clostridioforme		5.79	0.16				
03775240	**Good identification**						**03775240**
Actinomyces israelii		100.0	0.36	RHA 7			
03775242	**Good identification**						**03775242**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.39	RHA 7		
03775300	**Good identification**					**03775300**
					SPO	
Actinomyces israelii		96.04	0.54	RHA 7	1	
Clostridium butyricum		3.96	0.09		99	
03775302	**Low discrimination**					**03775302**
					GRA	
Actinomyces israelii		89.48	0.57	RHA 7	99	
Clostridium clostridioforme		4.60	0.25		50	
Parabacteroides distasonis		2.42	0.26		1	
03775310	**Good identification**					**03775310**
Actinomyces israelii		100.0	0.40	RHA 7		
03775312	**Low discrimination**					**03775312**
Clostridium clostridioforme		54.72	0.36	MNS 99		
Actinomyces israelii		34.89	0.43	ARA 94		
Bifidobacterium adolescentis		7.00	0.24			
03775322	**Good identification**					**03775322**
					GRA	
Parabacteroides distasonis		93.98	0.60	MNS 98	1	
Actinomyces israelii		2.24	0.36		99	
Clostridium clostridioforme		1.81	0.25		50	
03775332	**Low discrimination**					**03775332**
					GRA	
Parabacteroides distasonis		85.38	0.56	MNS 98	1	
Clostridium clostridioforme		13.88	0.36		50	
Actinomyces israelii		0.57	0.22		99	
03775340	**Good identification**					**03775340**
Actinomyces israelii		100.0	0.47	RHA 7		
03775342	**Good identification**					**03775342**
Actinomyces israelii		99.23	0.50	RHA 7		
Bifidobacterium adolescentis		0.77	0.07			
03775352	**Acceptable identification**					**03775352**
Actinomyces israelii		94.01	0.36	RHA 7		
Bifidobacterium adolescentis		5.99	0.09	MLZ 13		
03775400	**Low discrimination**					**03775400**
					SPO	
Actinomyces israelii		84.47	0.61		1	
Clostridium butyricum		15.53	0.27		99	
03775402	**Low discrimination**					**03775402**
					SPO	
Actinomyces israelii		89.28	0.64		1	
Bifidobacterium adolescentis		5.39	0.37		1	
Clostridium butyricum		4.26	0.20		99	
03775410	**Acceptable identification**					**03775410**
					SPO	
Actinomyces israelii		93.92	0.47	MLZ 13	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		6.08	0.06		99	
03775412	**Low discrimination**					**03775412**
Actinomyces israelii		53.22	0.50	MLZ 13		
Bifidobacterium adolescentis		26.29	0.38			
Clostridium clostridioforme		19.60	0.33			
03775420	**Low discrimination**					**03775420**
					SPO	
Actinomyces israelii		79.97	0.41	NAG 6	1	
Clostridium butyricum		20.03	0.09		99	
03775422	**Low discrimination**					**03775422**
					SPO	
Actinomyces israelii		77.76	0.44	NAG 6	1	
Clostridium clostridioforme		14.72	0.22		66	
Clostridium butyricum		5.05	0.02		99	
03775440	**Good identification**					**03775440**
					SPO	
Actinomyces israelii		96.05	0.54		1	
Clostridium butyricum		3.95	0.09		99	
03775442	**Very good identification**					**03775442**
					SPO	
Actinomyces israelii		97.10	0.57		1	
Bifidobacterium adolescentis		1.86	0.21		1	
Clostridium butyricum		1.04	0.02		99	
03775450	**Good identification**					**03775450**
Actinomyces israelii		100.0	0.40	MLZ 13		
03775452	**Low discrimination**					**03775452**
Actinomyces israelii		86.43	0.43	MLZ 13		
Bifidobacterium adolescentis		13.57	0.23			
03775462	**Good identification**					**03775462**
Actinomyces israelii		100.0	0.37	NAG 6		
03775500	**Low discrimination**					**03775500**
					SPO	
Clostridium butyricum		51.89	0.51	ESL 88	99	
Actinomyces israelii		47.04	0.71	MNS 99	1	
Bifidobacterium adolescentis		1.07	0.37		1	
03775502	**Low discrimination**					**03775502**
					SPO	
Bifidobacterium adolescentis		53.54	0.71	MNS 87	1	
Actinomyces israelii		35.83	0.74		1	
Clostridium butyricum		10.25	0.44		99	
03775503	**Low discrimination**					**03775503**
					SPO	
Bifidobacterium adolescentis		53.75	0.37	MNS 87	1	
Actinomyces israelii		35.97	0.40	URE 1	1	
Clostridium butyricum		10.29	0.10		99	
03775510	**Low discrimination**					**03775510**
					SPO	
Actinomyces israelii		63.35	0.57	MLZ 13	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		24.62	0.29		99		
Bifidobacterium adolescentis		11.73	0.38		1		
03775512	**Low discrimination**						**03775512**
Bifidobacterium adolescentis		89.63	0.73	MNS 87			
Actinomyces israelii		7.33	0.60				
Clostridium clostridioforme		2.26	0.42				
03775513	**Acceptable identification**						**03775513**
Bifidobacterium adolescentis		90.33	0.38	MNS 87			
Actinomyces israelii		7.39	0.26	URE 1			
Clostridium clostridioforme		2.28	0.08				
03775522	**Low discrimination**						**03775522**
					GRA	SPO	
Actinomyces israelii		38.69	0.54	NAG 6	99	1	
Parabacteroides distasonis		30.51	0.48		1	1	
Clostridium butyricum		15.08	0.26		99	99	
03775530	**Low discrimination**						**03775530**
					SPO		
Actinomyces israelii		58.82	0.37	MLZ 13	1		
Clostridium butyricum		31.14	0.11	NAG 6	99		
Clostridium clostridioforme		4.30	0.08		66		
03775532	**Low discrimination**						**03775532**
					GRA		
Clostridium clostridioforme		44.23	0.42	MNS 99	50		
Parabacteroides distasonis		26.02	0.44	RHA 88	1		
Bifidobacterium adolescentis		17.72	0.38		99		
03775540	**Low discrimination**						**03775540**
					SPO		
Actinomyces israelii		79.75	0.64		1		
Clostridium butyricum		19.67	0.33		99		
Bifidobacterium adolescentis		0.57	0.21		1		
03775542	**Low discrimination**						**03775542**
					SPO		
Actinomyces israelii		64.98	0.67		1		
Bifidobacterium adolescentis		30.86	0.56		1		
Clostridium butyricum		4.16	0.26		99		
03775550	**Low discrimination**						**03775550**
					SPO		
Actinomyces israelii		86.36	0.50	MLZ 13	1		
Clostridium butyricum		7.50	0.11		99		
Bifidobacterium adolescentis		5.08	0.23		1		
03775552	**Low discrimination**						**03775552**
					SPO		
Bifidobacterium adolescentis		79.04	0.57	MNS 87	1		
Actinomyces israelii		20.34	0.53	SOR 11	1		
Clostridium butyricum		0.46	0.04		99		
03775560	**Low discrimination**						**03775560**
					SPO		
Actinomyces israelii		74.85	0.44	NAG 6	1		
Clostridium butyricum		25.15	0.15		99		
03775562	**Low discrimination**						**03775562**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		84.55	0.47	NAG 6	1	
Clostridium butyricum		7.37	0.08		99	
Bifidobacterium adolescentis		6.36	0.21		1	
03775600	**Acceptable identification**					**03775600**
					SPO	
Actinomyces israelii		92.98	0.42	RHA 7	1	
Clostridium butyricum		7.02	0.02		99	
03775602	**Low discrimination**					**03775602**
					SPO	
Clostridium clostridioforme		53.58	0.37	MNS 99	66	
Actinomyces israelii		45.53	0.45		1	
Clostridium butyricum		0.89	-.06		99	
03775612	**Good identification**					**03775612**
Clostridium clostridioforme		97.29	0.48	MNS 99		
Actinomyces israelii		2.71	0.31			
03775622	**Low discrimination**					**03775622**
					GRA	
Clostridium clostridioforme		89.04	0.37	MNS 99	50	
Parabacteroides distasonis		6.13	0.22		1	
Actinomyces israelii		4.83	0.24		99	
03775632	**Good identification**					**03775632**
Clostridium clostridioforme		99.19	0.48	MNS 99		
Parabacteroides distasonis		0.81	0.18			
03775642	**Good identification**					**03775642**
Actinomyces israelii		100.0	0.38	RHA 7		
03775700	**Low discrimination**					**03775700**
					SPO	
Actinomyces israelii		68.11	0.52	RHA 7	1	
Clostridium butyricum		30.87	0.25		99	
Clostridium clostridioforme		1.02	0.11		66	
03775702	**Low discrimination**					**03775702**
Actinomyces israelii		43.04	0.55	RHA 7		
Clostridium clostridioforme		42.40	0.46			
Bifidobacterium adolescentis		8.63	0.37			
03775710	**Low discrimination**					**03775710**
					SPO	
Actinomyces israelii		61.93	0.38	RHA 7	1	
Clostridium clostridioforme		28.18	0.23	MLZ 13	66	
Clostridium butyricum		9.89	0.03		99	
03775712	**Good identification**					**03775712**
Clostridium clostridioforme		91.60	0.57	MNS 99		
Bifidobacterium adolescentis		5.00	0.38			
Actinomyces israelii		3.05	0.41			
03775722	**Low discrimination**					**03775722**
Parabacteroides distasonis		61.57	0.57	MNS 98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		30.23	0.46			
Bacteroides ovatus		5.63	0.23			
03775732	**Low discrimination**					**03775732**
Clostridium clostridioforme		78.68	0.57	MNS 99		
Parabacteroides distasonis		18.94	0.52			
Bacteroides ovatus		2.15	0.20			
03775740	**Acceptable identification**					**03775740**
					SPO	
Actinomyces israelii		90.80	0.45	RHA 7	1	
Clostridium butyricum		9.20	0.07		99	
03775742	**Low discrimination**					**03775742**
					SPO	
Actinomyces israelii		89.64	0.48	RHA 7	1	
Bifidobacterium adolescentis		5.71	0.21		1	
Clostridium butyricum		2.36	0.00		99	
03776000	**Good identification**					**03776000**
Actinomyces israelii		100.0	0.41	GAL 99		
03776002	**Good identification**					**03776002**
Actinomyces israelii		100.0	0.44	GAL 99		
03776042	**Good identification**					**03776042**
Actinomyces israelii		100.0	0.37	GAL 99		
03776100	**Good identification**					**03776100**
Actinomyces israelii		99.26	0.52	GAL 99		
Actinomyces naeslundii		0.74	0.24			
03776102	**Good identification**					**03776102**
Actinomyces israelii		96.96	0.55	GAL 99		
Bifidobacterium adolescentis		2.56	0.21			
Actinomyces naeslundii		0.48	0.24			
03776110	**Good identification**					**03776110**
Actinomyces israelii		100.0	0.37	GAL 99		
03776112	**Low discrimination**					**03776112**
Actinomyces israelii		82.25	0.41	GAL 99		
Bifidobacterium adolescentis		17.75	0.23	MLZ 13		
03776140	**Good identification**					**03776140**
Actinomyces israelii		100.0	0.45	GAL 99		
03776142	**Good identification**					**03776142**
Actinomyces israelii		100.0	0.48	GAL 99		
03776302	**Good identification**					**03776302**
Actinomyces israelii		100.0	0.35	GAL 99		
03776400	**Good identification**					**03776400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.40	GAL 99		
03776402	**Good identification**					**03776402**
Actinomyces israelii		100.0	0.43	GAL 99		
03776442	**Good identification**					**03776442**
Actinomyces israelii		100.0	0.36	GAL 99		
03776500	**Good identification**					**03776500**
					SPO	
Actinomyces israelii		97.88	0.50	GAL 99	1	
Clostridium butyricum		2.12	0.01		99	
03776502	**Low discrimination**					**03776502**
					SPO	
Actinomyces israelii		81.88	0.53	GAL 99	1	
Bifidobacterium adolescentis		17.67	0.35		1	
Clostridium butyricum		0.46	-.07		99	
03776510	**Good identification**					**03776510**
Actinomyces israelii		100.0	0.36	GAL 99		
03776512	**Low discrimination**					**03776512**
Bifidobacterium adolescentis		62.56	0.37	MNS 87		
Actinomyces israelii		35.45	0.39	GAL 99		
Bifidobacterium dentium		1.99	-.07			
03776540	**Good identification**					**03776540**
Actinomyces israelii		100.0	0.43	GAL 99		
03776542	**Acceptable identification**					**03776542**
Actinomyces israelii		93.58	0.46	GAL 99		
Bifidobacterium adolescentis		6.42	0.20			
03777000	**Very good identification**					**03777000**
					SPO	
Actinomyces israelii		99.71	0.75		1	
Actinomyces naeslundii		0.25	0.40		1	
Clostridium butyricum		0.03	-.05		99	
03777001	**Low discrimination**					**03777001**
Actinomyces israelii		72.53	0.41	URE 1		
Actinomyces naeslundii		27.47	0.43			
03777002	**Very good identification**					**03777002**
Actinomyces israelii		99.72	0.78			
Actinomyces naeslundii		0.17	0.40			
Bifidobacterium adolescentis		0.11	0.21			
03777003	**Low discrimination**					**03777003**
Actinomyces israelii		79.84	0.44	URE 1		
Actinomyces naeslundii		20.16	0.43			
03777010	**Very good identification**					**03777010**
Actinomyces israelii		100.0	0.61	MLZ 13		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
03777012	**Very good identification**					**03777012**
Actinomyces israelii		99.14	0.64	MLZ 13		
Bifidobacterium adolescentis		0.86	0.23			
03777020	**Good identification**					**03777020**
Actinomyces israelii		96.16	0.55	NAG 6		
Actinomyces naeslundii		3.84	0.40			
03777021	**Low discrimination**					**03777021**
Actinomyces naeslundii		85.58	0.43	MNS 88		
Actinomyces israelii		14.42	0.21	XYL 1		
03777022	**Very good identification**					**03777022**
Actinomyces israelii		97.41	0.58	NAG 6		
Actinomyces naeslundii		2.60	0.40			
03777023	**Low discrimination**					**03777023**
Actinomyces naeslundii		79.82	0.43	MNS 88		
Actinomyces israelii		20.18	0.24	XYL 1		
03777030	**Good identification**					**03777030**
Actinomyces israelii		100.0	0.41	MLZ 13		
03777032	**Good identification**					**03777032**
Actinomyces israelii		100.0	0.44	MLZ 13		
03777040	**Very good identification**					**03777040**
Actinomyces israelii		99.88	0.68			
Actinomyces naeslundii		0.12	0.27			
03777042	**Very good identification**					**03777042**
Actinomyces israelii		99.92	0.71			
Actinomyces naeslundii		0.08	0.27			
03777043	**Low discrimination**					**03777043**
Actinomyces israelii		89.72	0.37	URE 1		
Actinomyces naeslundii		10.28	0.30			
03777050	**Good identification**					**03777050**
Actinomyces israelii		100.0	0.54	MLZ 13		
03777052	**Very good identification**					**03777052**
Actinomyces israelii		99.72	0.57	MLZ 13		
Bifidobacterium adolescentis		0.28	0.07			
03777060	**Good identification**					**03777060**
Actinomyces israelii		98.22	0.48	NAG 6		
Actinomyces naeslundii		1.78	0.27			
03777062	**Good identification**					**03777062**
Actinomyces israelii		98.81	0.51	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		1.20	0.27			
03777072	**Good identification**					**03777072**
Actinomyces israelii		100.0	0.37	MLZ 13		
03777100	**Very good identification**					**03777100**
					SPO	
Actinomyces israelii		99.57	0.86		1	
Clostridium butyricum		0.20	0.18		99	
Actinomyces naeslundii		0.18	0.48		1	
03777101	**Low discrimination**					**03777101**
Actinomyces israelii		78.77	0.52	URE 1		
Actinomyces naeslundii		21.23	0.51			
03777102	**Very good identification**					**03777102**
Actinomyces israelii		97.25	0.89			
Bifidobacterium adolescentis		2.57	0.55			
Actinomyces naeslundii		0.12	0.48			
03777103	**Low discrimination**					**03777103**
Actinomyces israelii		82.92	0.55	URE 1		
Actinomyces naeslundii		14.90	0.51			
Bifidobacterium adolescentis		2.19	0.21			
03777110	**Very good identification**					**03777110**
					SPO	
Actinomyces israelii		99.42	0.72	MLZ 13	1	
Bifidobacterium adolescentis		0.33	0.23		1	
Clostridium tertium		0.12	0.07		99	
03777111	**Acceptable identification**					**03777111**
Actinomyces israelii		91.33	0.37	MLZ 13		
Actinomyces naeslundii		8.67	0.29	URE 1		
03777112	**Low discrimination**					**03777112**
					SPO	
Actinomyces israelii		82.03	0.75	MLZ 13	1	
Bifidobacterium adolescentis		17.70	0.57		1	
Clostridium tertium		0.20	0.15		99	
03777113	**Low discrimination**					**03777113**
Actinomyces israelii		78.18	0.41	MLZ 13		
Bifidobacterium adolescentis		16.87	0.23	URE 1		
Actinomyces naeslundii		4.95	0.29			
03777120	**Very good identification**					**03777120**
					SPO	
Actinomyces israelii		95.65	0.65	NAG 6	1	
Actinomyces naeslundii		2.72	0.48		1	
Clostridium ramosum		0.92	0.15		92	
03777121	**Low discrimination**					**03777121**
Actinomyces naeslundii		80.85	0.51	MNS 88		
Actinomyces israelii		19.15	0.31	XYL 1		
03777122	**Good identification**					**03777122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		94.27	0.68	NAG 6	1	
Clostridium ramosum		2.42	0.25		92	
Actinomyces naeslundii		1.79	0.48		1	
03777123	**Low discrimination**					**03777123**
Actinomyces naeslundii		73.79	0.51	MNS 88		
Actinomyces israelii		26.21	0.34	XYL 1		
03777130	**Acceptable identification**					**03777130**
					SPO	
Actinomyces israelii		93.81	0.51	MLZ 13	1	
Clostridium tertium		5.25	0.15	NAG 6	99	
Actinomyces naeslundii		0.94	0.26		1	
03777132	**Low discrimination**					**03777132**
					SPO	
Actinomyces israelii		85.30	0.54	MLZ 13	1	
Clostridium tertium		9.54	0.24	NAG 6	99	
Bifidobacterium adolescentis		2.91	0.23		1	
03777140	**Very good identification**					**03777140**
					SPO	
Actinomyces israelii		99.87	0.79		1	
Actinomyces naeslundii		0.08	0.35		1	
Clostridium butyricum		0.04	0.00		99	
03777141	**Low discrimination**					**03777141**
Actinomyces israelii		89.10	0.45	URE 1		
Actinomyces naeslundii		10.90	0.38			
03777142	**Very good identification**					**03777142**
Actinomyces israelii		99.10	0.82			
Bifidobacterium adolescentis		0.83	0.40			
Actinomyces naeslundii		0.05	0.35			
03777143	**Acceptable identification**					**03777143**
Actinomyces israelii		92.46	0.48	URE 1		
Actinomyces naeslundii		7.54	0.38			
03777150	**Very good identification**					**03777150**
					COC	
Actinomyces israelii		99.72	0.65	MLZ 13	1	
Blautia producta		0.18	-.03		99	
Bifidobacterium adolescentis		0.10	0.07		1	
03777152	**Good identification**					**03777152**
Actinomyces israelii		93.58	0.68	MLZ 13		
Bifidobacterium adolescentis		6.42	0.41			
03777160	**Very good identification**					**03777160**
Actinomyces israelii		98.73	0.58	NAG 6		
Actinomyces naeslundii		1.27	0.35			
03777161	**Low discrimination**					**03777161**
Actinomyces naeslundii		65.71	0.38	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	34.29	0.24	XYL 1		
03777162	**Very good identification**					**03777162**
	Actinomyces israelii	99.15	0.61	NAG 6		
	Actinomyces naeslundii	0.85	0.35			
03777163	**Low discrimination**					**03777163**
	Actinomyces naeslundii	56.09	0.38	MNS 88		
	Actinomyces israelii	43.91	0.27	XYL 1		
03777170	**Good identification**					**03777170**
	Actinomyces israelii	100.0	0.44	MLZ 13		
03777172	**Good identification**					**03777172**
	Actinomyces israelii	98.93	0.47	MLZ 13		
	Bifidobacterium adolescentis	1.07	0.07	NAG 6		
03777200	**Very good identification**					**03777200**
	Actinomyces israelii	100.0	0.56	RHA 7		
03777202	**Very good identification**					**03777202**
	Actinomyces israelii	100.0	0.59	RHA 7		
03777210	**Good identification**					**03777210**
	Actinomyces israelii	100.0	0.42	RHA 7		
03777212	**Good identification**					**03777212**
	Actinomyces israelii	100.0	0.45	RHA 7		
03777220	**Good identification**					**03777220**
	Actinomyces israelii	100.0	0.36	RHA 7		
03777222	**Good identification**					**03777222**
	Actinomyces israelii	100.0	0.39	RHA 7		
03777240	**Good identification**					**03777240**
	Actinomyces israelii	100.0	0.49	RHA 7		
03777242	**Good identification**					**03777242**
	Actinomyces israelii	100.0	0.52	RHA 7		
03777250	**Good identification**					**03777250**
	Actinomyces israelii	100.0	0.35	RHA 7		
03777252	**Good identification**					**03777252**
	Actinomyces israelii	100.0	0.38	RHA 7		
03777300	**Very good identification**					**03777300**
					SPO	
	Actinomyces israelii	99.92	0.67	RHA 7	1	
	Clostridium butyricum	0.08	-.08		99	
03777302	**Very good identification**					**03777302**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		99.65	0.70	RHA 7			
Bifidobacterium adolescentis		0.35	0.21				
03777303	**Good identification**						**03777303**
Actinomyces israelii		100.0	0.35	RHA 7			
03777310	**Good identification**						**03777310**
Actinomyces israelii		100.0	0.52	RHA 7			
03777312	**Good identification**						**03777312**
Actinomyces israelii		97.19	0.55	RHA 7			
Bifidobacterium adolescentis		2.81	0.23	MLZ 13			
03777320	**Low discrimination**						**03777320**
					SPO		
Actinomyces israelii		85.51	0.46	RHA 7	1		
Clostridium ramosum		14.49	0.17	NAG 6	92		
03777322	**Low discrimination**						**03777322**
					SPO	GRA	
Actinomyces israelii		65.50	0.49	RHA 7	1	99	
Clostridium ramosum		29.61	0.27	NAG 6	92	50	
Parabacteroides distasonis		4.90	0.26		1	1	
03777340	**Very good identification**						**03777340**
Actinomyces israelii		100.0	0.60	RHA 7			
03777342	**Very good identification**						**03777342**
Actinomyces israelii		100.0	0.63	RHA 7			
03777350	**Good identification**						**03777350**
					COC		
Actinomyces israelii		97.71	0.45	RHA 7	1		
Blautia producta		2.29	-.03	MLZ 13	99		
03777352	**Good identification**						**03777352**
Actinomyces israelii		99.09	0.48	RHA 7			
Bifidobacterium adolescentis		0.91	0.07	MLZ 13			
03777360	**Good identification**						**03777360**
Actinomyces israelii		100.0	0.39	RHA 7			
03777362	**Good identification**						**03777362**
Actinomyces israelii		100.0	0.42	RHA 7			
03777400	**Very good identification**						**03777400**
					SPO		
Actinomyces israelii		99.64	0.74		1		
Clostridium butyricum		0.36	0.11		99		
03777401	**Good identification**						**03777401**
Actinomyces israelii		100.0	0.40	URE 1			
03777402	**Very good identification**						**03777402**
					SPO		
Actinomyces israelii		99.04	0.77		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		0.86	0.35		1	
Clostridium butyricum		0.09	0.04		99	
03777403	**Good identification**					**03777403**
Actinomyces israelii		100.0	0.43	URE 1		
03777410	**Very good identification**					**03777410**
Actinomyces israelii		100.0	0.60	MLZ 13		
03777412	**Good identification**					**03777412**
Actinomyces israelii		93.15	0.63	MLZ 13		
Bifidobacterium adolescentis		6.64	0.37			
Bifidobacterium dentium		0.21	-.07			
03777420	**Good identification**					**03777420**
					SPO	
Actinomyces israelii		99.51	0.54	NAG 6	1	
Clostridium butyricum		0.49	-.07		99	
03777422	**Very good identification**					**03777422**
Actinomyces israelii		100.0	0.57	NAG 6		
03777430	**Good identification**					**03777430**
Actinomyces israelii		100.0	0.39	MLZ 13		
03777432	**Acceptable identification**					**03777432**
Actinomyces israelii		94.06	0.42	MLZ 13		
Bifidobacterium dentium		5.94	-.03	NAG 6		
03777440	**Very good identification**					**03777440**
					SPO	
Actinomyces israelii		99.92	0.67		1	
Clostridium butyricum		0.08	-.07		99	
03777442	**Very good identification**					**03777442**
Actinomyces israelii		99.72	0.70			
Bifidobacterium adolescentis		0.28	0.20			
03777443	**Good identification**					**03777443**
Actinomyces israelii		100.0	0.36	URE 1		
03777450	**Good identification**					**03777450**
					COC	
Actinomyces israelii		99.15	0.53	MLZ 13	1	
Blautia producta		0.85	-.03		99	
03777452	**Good identification**					**03777452**
Actinomyces israelii		97.78	0.56	MLZ 13		
Bifidobacterium adolescentis		2.22	0.21			
03777460	**Good identification**					**03777460**
Actinomyces israelii		100.0	0.47	NAG 6		
03777462	**Good identification**					**03777462**
Actinomyces israelii		100.0	0.50	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
03777472	**Good identification**					**03777472**
Actinomyces israelii		100.0	0.35	MLZ 13		
03777500	**Very good identification**					**03777500**
					SPO	
Actinomyces israelii		97.57	0.84		1	
Clostridium butyricum		2.11	0.35		99	
Bifidobacterium adolescentis		0.32	0.35		1	
03777501	**Good identification**					**03777501**
					SPO	
Actinomyces israelii		97.88	0.50	URE 1	1	
Clostridium butyricum		2.12	0.01		99	
03777502	**Low discrimination**					**03777502**
					SPO	
Actinomyces israelii		81.87	0.87		1	
Bifidobacterium adolescentis		17.67	0.70		1	
Clostridium butyricum		0.46	0.28		99	
03777503	**Low discrimination**					**03777503**
					SPO	
Actinomyces israelii		81.88	0.53	URE 1	1	
Bifidobacterium adolescentis		17.67	0.35		1	
Clostridium butyricum		0.46	-.07		99	
03777510	**Very good identification**					**03777510**
Actinomyces israelii		95.83	0.70	MLZ 13		
Bifidobacterium adolescentis		2.56	0.37			
Bifidobacterium dentium		0.80	0.10			
03777511	**Good identification**					**03777511**
Actinomyces israelii		100.0	0.36	MLZ 13		
03777512	**Low discrimination**					**03777512**
Bifidobacterium adolescentis		62.51	0.71	MNS 87		
Actinomyces israelii		35.42	0.73			
Bifidobacterium dentium		1.99	0.27			
03777513	**Low discrimination**					**03777513**
Bifidobacterium adolescentis		62.56	0.37	MNS 87		
Actinomyces israelii		35.45	0.39	URE 1		
Bifidobacterium dentium		1.99	-.07			
03777520	**Very good identification**					**03777520**
					SPO	
Actinomyces israelii		97.14	0.64	NAG 6	1	
Clostridium butyricum		2.86	0.17		99	
03777522	**Very good identification**					**03777522**
					SPO	
Actinomyces israelii		95.64	0.67	NAG 6	1	
Bifidobacterium adolescentis		3.27	0.35		1	
Clostridium butyricum		0.73	0.10		99	
03777530	**Low discrimination**					**03777530**
					SPO	
Actinomyces israelii		80.51	0.50	MLZ 13	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		18.66	0.15	NAG 6	1		
Clostridium butyricum		0.84	-.05		99		
03777540	**Very good identification**						**03777540**
					COC	SPO	
Actinomyces israelii		98.38	0.77		1	1	
Blautia producta		1.05	0.24		99	1	
Clostridium butyricum		0.48	0.17		1	99	
03777541	**Good identification**						**03777541**
Actinomyces israelii		100.0	0.43	URE 1			
03777542	**Good identification**						**03777542**
					SPO		
Actinomyces israelii		93.40	0.80		1		
Bifidobacterium adolescentis		6.41	0.54		1		
Clostridium butyricum		0.12	0.10		99		
03777543	**Acceptable identification**						**03777543**
Actinomyces israelii		93.58	0.46	URE 1			
Bifidobacterium adolescentis		6.42	0.20				
03777550	**Low discrimination**						**03777550**
					COC		
Actinomyces israelii		81.74	0.63	MLZ 13	1		
Blautia producta		17.43	0.32		99		
Bifidobacterium adolescentis		0.70	0.21		1		
03777552	**Low discrimination**						**03777552**
					COC		
Actinomyces israelii		63.36	0.66	MLZ 13	1		
Bifidobacterium adolescentis		35.54	0.56		1		
Blautia producta		1.00	0.15		99		
03777560	**Very good identification**						**03777560**
					COC	SPO	
Actinomyces israelii		97.55	0.57	NAG 6	1	1	
Blautia producta		1.80	0.07		99	1	
Clostridium butyricum		0.64	-.02		1	99	
03777562	**Very good identification**						**03777562**
Actinomyces israelii		98.93	0.60	NAG 6			
Bifidobacterium adolescentis		1.07	0.20				
03777570	**Low discrimination**						**03777570**
					COC		
Actinomyces israelii		72.93	0.43	MLZ 13	1		
Blautia producta		27.07	0.15	NAG 6	99		
03777572	**Low discrimination**						**03777572**
Actinomyces israelii		86.44	0.46	MLZ 13			
Bifidobacterium adolescentis		7.67	0.21	NAG 6			
Bifidobacterium dentium		3.51	-.03				
03777600	**Good identification**						**03777600**
Actinomyces israelii		100.0	0.55	RHA 7			
03777602	**Very good identification**						**03777602**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.58	RHA 7		
03777610	**Good identification**					**03777610**
Actinomyces israelii		100.0	0.41	RHA 7		
03777612	**Acceptable identification**					**03777612**
Actinomyces israelii		93.99	0.44	RHA 7		
Clostridium clostridioforme		6.01	0.14	MLZ 13		
03777622	**Good identification**					**03777622**
Actinomyces israelii		100.0	0.37	RHA 7		
03777640	**Good identification**					**03777640**
Actinomyces israelii		100.0	0.48	RHA 7		
03777642	**Good identification**					**03777642**
Actinomyces israelii		100.0	0.51	RHA 7		
03777652	**Good identification**					**03777652**
Actinomyces israelii		100.0	0.37	RHA 7		
03777700	**Very good identification**					**03777700**
					SPO	
Actinomyces israelii		99.12	0.65	RHA 7	1	
Clostridium butyricum		0.88	0.09		99	
03777702	**Very good identification**					**03777702**
					SPO	
Actinomyces israelii		96.80	0.68	RHA 7	1	
Bifidobacterium adolescentis		2.80	0.35		1	
Clostridium butyricum		0.22	0.02		99	
03777710	**Good identification**					**03777710**
					COC	
Actinomyces israelii		98.90	0.51	RHA 7	1	
Blautia producta		1.10	-.03	MLZ 13	99	
03777712	**Low discrimination**					**03777712**
Actinomyces israelii		77.05	0.54	RHA 7		
Bifidobacterium adolescentis		18.25	0.37	MLZ 13		
Clostridium clostridioforme		4.12	0.23			
03777720	**Good identification**					**03777720**
Actinomyces israelii		100.0	0.45	RHA 7		
03777722	**Low discrimination**					**03777722**
					GRA	
Actinomyces israelii		79.83	0.48	RHA 7	99	
Bacteroides ovatus		13.50	0.15	NAG 6	1	
Parabacteroides distasonis		4.47	0.22		1	
03777740	**Low discrimination**					**03777740**
					COC	
Actinomyces israelii		87.63	0.58	RHA 7	1	
Blautia producta		12.37	0.24		99	
03777742	**Very good identification**					**03777742**
					COC	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		98.07	0.61	RHA 7	1		
Blautia producta		1.03	0.07		99		
Bifidobacterium adolescentis		0.90	0.20		1		
03777752	**Low discrimination**						**03777752**
					COC		
Actinomyces israelii		77.81	0.47	RHA 7	1		
Blautia producta		16.33	0.15	MLZ 13	99		
Bifidobacterium adolescentis		5.86	0.21		1		
03777760	**Low discrimination**						**03777760**
					COC		
Actinomyces israelii		80.28	0.38	RHA 7	1		
Blautia producta		19.72	0.07	NAG 6	99		
03777762	**Good identification**						**03777762**
Actinomyces israelii		100.0	0.41	RHA 7			
04000000	**Low discrimination**						**04000000**
Parvimonas micra		39.19	0.84	MNS 10			
Anaerococcus prevotii		18.38	0.84				
Peptococcus niger		9.28	0.74				
04000001	**Very good identification**						**04000001**
					COC	GRA	
Anaerococcus prevotii		97.41	1.00		99	99	
Campylobacter ureolyticus		1.62	0.66		1	1	
Peptococcus niger		0.54	0.57		99	99	
04000002	**Low discrimination**						**04000002**
					COC		
Anaerococcus prevotii		32.29	0.73	URE 90	99		
Parvimonas micra		30.60	0.67		99		
Atopobium minutum		20.58	0.84		1		
04000003	**Very good identification**						**04000003**
Anaerococcus prevotii		99.35	0.90				
Peptococcus niger		0.39	0.44				
Parvimonas micra		0.11	0.33				
04000010	**Low discrimination**						**04000010**
Parvimonas micra		39.12	0.50	MNS 10			
Anaerococcus prevotii		18.34	0.50	MLZ 1			
Peptococcus niger		9.26	0.40				
04000011	**Very good identification**						**04000011**
					COC	GRA	
Anaerococcus prevotii		97.33	0.66	MLZ 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04000012	**Low discrimination**						**04000012**
					COC		
Anaerococcus prevotii		32.28	0.39	MLZ 1	99		
Parvimonas micra		30.59	0.33	URE 90	99		
Atopobium minutum		20.58	0.50		1		
04000013	**Good identification**						**04000013**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Anaerococcus prevotii		99.30	0.56	MLZ 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04000020	**Low discrimination**						**04000020**
					COC		
Atopobium minutum		42.01	0.79		1		
Peptococcus niger		29.57	0.58		99		
Parvimonas micra		11.36	0.50		99		
04000021	**Good identification**						**04000021**
					COC	GRA	
Anaerococcus prevotii		91.61	0.66	NAG 1	99	99	
Peptococcus niger		5.59	0.41		99	99	
Campylobacter ureolyticus		1.52	0.32		1	1	
04000022	**Low discrimination**						**04000022**
					COC	GRA	
Pseudoflavonifractor capillosus		58.87	0.66	MNS 1	1	1	
Atopobium minutum		31.94	0.76		1	99	
Peptococcus niger		5.95	0.45		99	99	
04000023	**Low discrimination**						**04000023**
					COC	GRA	
Anaerococcus prevotii		89.93	0.56	NAG 1	99	99	
Pseudoflavonifractor capillosus		3.91	0.32		1	1	
Peptococcus niger		3.87	0.28		99	99	
04000030	**Low discrimination**						**04000030**
					COC		
Atopobium minutum		41.70	0.44	MLZ 1	1		
Peptococcus niger		29.35	0.24		99		
Parvimonas micra		11.27	0.15		99		
04000040	**Low discrimination**						**04000040**
					COC	SPO	
Parvimonas micra		35.58	0.50	MNS 10	99	1	
Anaerococcus prevotii		16.69	0.50	SOR 1	99	1	
Clostridium botulinum biovar A		9.01	0.51		1	99	
04000041	**Very good identification**						**04000041**
					COC	GRA	
Anaerococcus prevotii		97.30	0.66	SOR 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04000042	**Low discrimination**						**04000042**
					COC		
Anaerococcus prevotii		32.07	0.39	SOR 1	99		
Parvimonas micra		30.40	0.33	URE 90	99		
Atopobium minutum		20.45	0.50		1		
04000043	**Good identification**						**04000043**
Anaerococcus prevotii		99.32	0.56	SOR 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04000060	**Low discrimination**						**04000060**
					COC		
Atopobium minutum		40.36	0.44	SOR 1	1		
Peptococcus niger		28.41	0.24		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parvimonas micra		10.91	0.15		99		
04000100	**Low discrimination**						**04000100**
Parvimonas micra		39.18	0.50	MNS 10			
Anaerococcus prevotii		18.37	0.50	LAC 1			
Peptococcus niger		9.28	0.40				
04000101	**Very good identification**						**04000101**
					COC	GRA	
Anaerococcus prevotii		97.42	0.66	LAC 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04000102	**Low discrimination**						**04000102**
					COC		
Anaerococcus prevotii		32.34	0.39	LAC 1	99		
Parvimonas micra		30.65	0.33	URE 90	99		
Atopobium minutum		20.62	0.50		1		
04000103	**Good identification**						**04000103**
Anaerococcus prevotii		99.37	0.56	LAC 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04000200	**Low discrimination**						**04000200**
Parvimonas micra		39.19	0.50	MNS 10			
Anaerococcus prevotii		18.38	0.50	RHA 1			
Peptococcus niger		9.28	0.40				
04000201	**Very good identification**						**04000201**
					COC	GRA	
Anaerococcus prevotii		97.42	0.66	RHA 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04000202	**Low discrimination**						**04000202**
					COC		
Anaerococcus prevotii		32.34	0.39	RHA 1	99		
Parvimonas micra		30.65	0.33	URE 90	99		
Atopobium minutum		20.62	0.50		1		
04000203	**Good identification**						**04000203**
Anaerococcus prevotii		99.37	0.56	RHA 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04000220	**Low discrimination**						**04000220**
					COC		
Atopobium minutum		42.16	0.44	RHA 1	1		
Peptococcus niger		29.68	0.24		99		
Parvimonas micra		11.39	0.15		99		
04000400	**Low discrimination**						**04000400**
Parvimonas micra		38.90	0.50	MNS 10			
Anaerococcus prevotii		18.24	0.50	ARA 1			
Peptococcus niger		9.21	0.40				
04000401	**Very good identification**						**04000401**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Anaerococcus prevotii		97.41	0.66	ARA 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04000402	**Low discrimination**						**04000402**
					COC		
Anaerococcus prevotii		32.32	0.39	ARA 1	99		
Parvimonas micra		30.63	0.33	URE 90	99		
Atopobium minutum		20.61	0.50		1		
04000403	**Good identification**						**04000403**
Anaerococcus prevotii		99.37	0.56	ARA 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04000420	**Low discrimination**						**04000420**
					COC		
Atopobium minutum		41.18	0.44	ARA 1	1		
Peptococcus niger		28.99	0.24		99		
Parvimonas micra		11.13	0.15		99		
04001000	**Low discrimination**						**04001000**
					COC		
Atopobium minutum		58.84	0.79		1		
Parvimonas micra		15.90	0.50		99		
Anaerococcus prevotii		7.46	0.50		99		
04001001	**Very good identification**						**04001001**
					COC	GRA	
Anaerococcus prevotii		96.57	0.66	GAL 1	99	99	
Campylobacter ureolyticus		1.61	0.32		1	1	
Atopobium minutum		0.86	0.44		1	99	
04001002	**Low discrimination**						**04001002**
					COC		
Atopobium minutum		89.35	0.76		1		
Anaerococcus prevotii		4.25	0.39		99		
Parvimonas micra		4.03	0.33		99		
04001003	**Good identification**						**04001003**
					COC		
Anaerococcus prevotii		97.16	0.56	GAL 1	99		
Atopobium minutum		2.29	0.41		1		
Peptococcus niger		0.38	0.10		99		
04001010	**Low discrimination**						**04001010**
					COC		
Atopobium minutum		58.75	0.44	MLZ 1	1		
Parvimonas micra		15.88	0.15		99		
Anaerococcus prevotii		7.45	0.15		99		
04001012	**Acceptable identification**						**04001012**
					COC		
Atopobium minutum		90.13	0.41	MLZ 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04001020	**Good identification**						**04001020**
					COC		
Atopobium minutum		94.80	0.70		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptococcus niger		2.02	0.24		99		
Gemella morbillorum		0.86	0.31		99		
04001022	**Low discrimination**						**04001022**
					COC	GRA	
Atopobium minutum		61.63	0.67		1	99	
Pseudoflavonifractor capillosus		37.86	0.50		1	1	
Peptococcus niger		0.35	0.11		99	99	
04001030	**Good identification**						**04001030**
					COC		
Atopobium minutum		97.91	0.36	MLZ 1	1		
Peptococcus niger		2.09	-.11		99		
04001040	**Low discrimination**						**04001040**
					COC		
Atopobium minutum		56.60	0.44	SOR 1	1		
Parvimonas micra		15.30	0.15		99		
Anaerococcus prevotii		7.17	0.15		99		
04001042	**Acceptable identification**						**04001042**
					COC		
Atopobium minutum		90.13	0.41	SOR 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04001060	**Good identification**						**04001060**
					COC		
Atopobium minutum		97.91	0.36	SOR 1	1		
Peptococcus niger		2.09	-.11		99		
04001100	**Low discrimination**						**04001100**
					COC		
Atopobium minutum		59.16	0.44	LAC 1	1		
Parvimonas micra		15.99	0.15		99		
Anaerococcus prevotii		7.50	0.15		99		
04001102	**Acceptable identification**						**04001102**
					COC		
Atopobium minutum		90.13	0.41	LAC 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04001200	**Low discrimination**						**04001200**
					COC		
Atopobium minutum		59.16	0.44	RHA 1	1		
Parvimonas micra		15.99	0.15		99		
Anaerococcus prevotii		7.50	0.15		99		
04001202	**Acceptable identification**						**04001202**
					COC		
Atopobium minutum		90.13	0.41	RHA 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04001220	**Good identification**						**04001220**
					COC		
Atopobium minutum		97.91	0.36	RHA 1	1		
Peptococcus niger		2.09	-.11		99		
04001400	**Low discrimination**						**04001400**
					COC		
Atopobium minutum		50.96	0.44	ARA 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		15.24	0.31		99		
Parvimonas micra		13.77	0.15		99		
04001402	**Acceptable identification**						**04001402**
					COC		
Atopobium minutum		90.13	0.41	ARA 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04001420	**Low discrimination**						**04001420**
					COC		
Atopobium minutum		89.20	0.36	ARA 1	1		
Gemella morbillorum		8.89	0.14		99		
Peptococcus niger		1.90	-.11		99		
04002000	**Low discrimination**						**04002000**
Parvimonas micra		38.80	0.50	MNS 10			
Anaerococcus prevotii		18.19	0.50	MAN 1			
Peptococcus niger		9.19	0.40				
04002001	**Very good identification**						**04002001**
					COC	GRA	
Anaerococcus prevotii		97.41	0.66	MAN 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04002002	**Low discrimination**						**04002002**
					COC		
Anaerococcus prevotii		32.32	0.39	MAN 1	99		
Parvimonas micra		30.63	0.33	URE 90	99		
Atopobium minutum		20.61	0.50		1		
04002003	**Good identification**						**04002003**
Anaerococcus prevotii		99.37	0.56	MAN 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04002020	**Low discrimination**						**04002020**
					COC		
Atopobium minutum		42.05	0.44	MAN 1	1		
Peptococcus niger		29.60	0.24		99		
Parvimonas micra		11.36	0.15		99		
04003000	**Low discrimination**						**04003000**
					COC		
Atopobium minutum		58.93	0.44	MAN 1	1		
Parvimonas micra		15.93	0.15		99		
Anaerococcus prevotii		7.47	0.15		99		
04003002	**Acceptable identification**						**04003002**
					COC		
Atopobium minutum		90.13	0.41	MAN 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04003020	**Good identification**						**04003020**
					COC		
Atopobium minutum		97.91	0.36	MAN 1	1		
Peptococcus niger		2.09	-.11		99		
04004000	**Low discrimination**						**04004000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parvimonas micra		38.67	0.50	MNS 10			
Anaerococcus prevotii		18.14	0.50	XYL 1			
Peptococcus niger		9.16	0.40				
04004001	**Very good identification**						**04004001**
					COC	GRA	
Anaerococcus prevotii		97.41	0.66	XYL 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04004002	**Low discrimination**						**04004002**
					COC		
Anaerococcus prevotii		32.30	0.39	XYL 1	99		
Parvimonas micra		30.62	0.33	URE 90	99		
Atopobium minutum		20.60	0.50		1		
04004003	**Good identification**						**04004003**
Anaerococcus prevotii		99.37	0.56	XYL 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04004020	**Low discrimination**						**04004020**
					COC		
Atopobium minutum		41.09	0.44	XYL 1	1		
Peptococcus niger		28.92	0.24		99		
Parvimonas micra		11.11	0.15		99		
04005000	**Low discrimination**						**04005000**
					COC		
Atopobium minutum		50.82	0.44	XYL 1	1		
Gemella morbillorum		15.20	0.31		99		
Parvimonas micra		13.73	0.15		99		
04005002	**Acceptable identification**						**04005002**
					COC		
Atopobium minutum		90.13	0.41	XYL 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04005020	**Low discrimination**						**04005020**
					COC		
Atopobium minutum		89.20	0.36	XYL 1	1		
Gemella morbillorum		8.89	0.14		99		
Peptococcus niger		1.90	-.11		99		
04010000	**Low discrimination**						**04010000**
					COC	GRA	
Fusobacterium varium		78.00	0.84		1	1	
Atopobium minutum		16.59	0.86		1	99	
Parvimonas micra		1.62	0.50		99	99	
04010001	**Low discrimination**						**04010001**
					COC	GRA	
Anaerococcus prevotii		85.58	0.66	FRU 1	99	99	
Fusobacterium varium		9.87	0.50		1	1	
Atopobium minutum		2.10	0.52		1	99	
04010002	**Low discrimination**						**04010002**
					COC	GRA	
Atopobium minutum		89.54	0.83		1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium varium		6.38	0.50		1	1	
Anaerococcus prevotii		1.54	0.39		99	99	
04010003	**Acceptable identification**						**04010003**
					COC	GRA	
Anaerococcus prevotii		92.96	0.56	FRU 1	99	99	
Atopobium minutum		6.08	0.49		1	99	
Fusobacterium varium		0.43	0.15		1	1	
04010010	**Low discrimination**						**04010010**
					GRA	SPO	
Fusobacterium varium		66.99	0.50	MLZ 1	1	1	
Atopobium minutum		14.25	0.52		99	1	
Clostridium difficile		13.98	0.41		99	99	
04010012	**Low discrimination**						**04010012**
					COC	GRA	
Atopobium minutum		88.45	0.49	MLZ 1	1	99	
Fusobacterium varium		6.30	0.15		1	1	
Anaerococcus prevotii		1.52	0.05		99	99	
04010020	**Low discrimination**						**04010020**
					GRA	SPO	
Atopobium minutum		81.01	0.78		99	1	
Fusobacterium varium		11.54	0.50		1	1	
Clostridium cadaveris		4.61	0.46		99	99	
04010022	**Low discrimination**						**04010022**
					GRA		
Atopobium minutum		81.38	0.75		99		
Pseudoflavonifractor capillosus		18.05	0.50		1		
Fusobacterium varium		0.18	0.15		1		
04010023	**Low discrimination**						**04010023**
					COC	GRA	
Atopobium minutum		58.65	0.41	URE 1	1	99	
Anaerococcus prevotii		27.18	0.21		99	99	
Pseudoflavonifractor capillosus		13.01	0.15		1	1	
04010030	**Low discrimination**						**04010030**
					GRA	SPO	
Atopobium minutum		79.32	0.44	MLZ 1	99	1	
Fusobacterium varium		11.30	0.15		1	1	
Clostridium cadaveris		4.51	0.12		99	99	
04010032	**Low discrimination**						**04010032**
					GRA		
Atopobium minutum		81.85	0.41	MLZ 1	99		
Pseudoflavonifractor capillosus		18.16	0.15		1		
04010040	**Low discrimination**						**04010040**
					GRA	SPO	
Fusobacterium varium		74.06	0.50	SOR 1	1	1	
Atopobium minutum		15.75	0.52		99	1	
Clostridium botulinum biovar A		2.46	0.30		99	99	
04010042	**Low discrimination**						**04010042**
					GRA		
Atopobium minutum		88.16	0.49	SOR 1	99		
Fusobacterium varium		6.28	0.15		1		
Pseudoramibacter alactolyticus		1.62	0.05		99		
04010060	**Low discrimination**						**04010060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Atopobium minutum		69.73	0.44	SOR 1	99	1	
Propionibacterium acnes		10.22	0.22		99	95	
Fusobacterium varium		9.94	0.15		1	1	
04010062	**Low discrimination**						**04010062**
					GRA		
Atopobium minutum		81.20	0.41	SOR 1	99		
Pseudoflavonifractor capillosus		18.01	0.15		1		
Pseudoramibacter alactolyticus		0.79	-.08		99		
04010100	**Low discrimination**						**04010100**
					COC	GRA	
Fusobacterium varium		77.00	0.50	LAC 1	1	1	
Atopobium minutum		16.38	0.52		1	99	
Parvimonas micra		1.60	0.15		99	99	
04010102	**Low discrimination**						**04010102**
					COC	GRA	
Atopobium minutum		89.15	0.49	LAC 1	1	99	
Fusobacterium varium		6.35	0.15		1	1	
Anaerococcus prevotii		1.53	0.05		99	99	
04010120	**Low discrimination**						**04010120**
					GRA		
Atopobium minutum		66.42	0.44	LAC 1	99		
Prevotella bivia		18.37	0.15		1		
Fusobacterium varium		9.46	0.15		1		
04010122	**Low discrimination**						**04010122**
					GRA		
Atopobium minutum		81.85	0.41	LAC 1	99		
Pseudoflavonifractor capillosus		18.16	0.15		1		
04010200	**Low discrimination**						**04010200**
					COC	GRA	
Fusobacterium varium		78.04	0.50	RHA 1	1	1	
Atopobium minutum		16.60	0.52		1	99	
Parvimonas micra		1.62	0.15		99	99	
04010202	**Acceptable identification**						**04010202**
					COC	GRA	
Atopobium minutum		90.03	0.49	RHA 1	1	99	
Fusobacterium varium		6.41	0.15		1	1	
Anaerococcus prevotii		1.55	0.05		99	99	
04010220	**Low discrimination**						**04010220**
					GRA	SPO	
Atopobium minutum		81.89	0.44	RHA 1	99	1	
Fusobacterium varium		11.67	0.15		1	1	
Clostridium cadaveris		4.66	0.12		99	99	
04010222	**Low discrimination**						**04010222**
					GRA		
Atopobium minutum		81.85	0.41	RHA 1	99		
Pseudoflavonifractor capillosus		18.16	0.15		1		
04010400	**Low discrimination**						**04010400**
					COC	GRA	
Fusobacterium varium		75.29	0.50	ARA 1	1	1	
Atopobium minutum		16.01	0.52		1	99	
Gemella morbillorum		3.93	0.37		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04010402	**Low discrimination**						**04010402**
					COC	GRA	
Atopobium minutum		89.73	0.49	ARA 1	1	99	
Fusobacterium varium		6.39	0.15		1	1	
Anaerococcus prevotii		1.54	0.05		99	99	
04010420	**Low discrimination**						**04010420**
					COC	GRA	
Atopobium minutum		77.19	0.44	ARA 1	1	99	
Fusobacterium varium		11.00	0.15		1	1	
Gemella morbillorum		6.31	0.20		99	99	
04010422	**Low discrimination**						**04010422**
					GRA		
Atopobium minutum		81.85	0.41	ARA 1	99		
Pseudoflavonifractor capillosus		18.16	0.15		1		
04011000	**Low discrimination**						**04011000**
					COC	GRA	
Atopobium minutum		83.37	0.78		1	99	
Fusobacterium varium		11.88	0.50		1	1	
Gemella morbillorum		3.00	0.48		99	99	
04011002	**Very good identification**						**04011002**
					GRA	CAT	
Atopobium minutum		99.38	0.75		99	1	
Fusobacterium varium		0.22	0.15		1	1	
Propionibacterium granulosum		0.11	0.26		99	90	
04011003	**Low discrimination**						**04011003**
					COC		
Atopobium minutum		68.33	0.41	URE 1	1		
Anaerococcus prevotii		31.67	0.21		99		
04011010	**Low discrimination**						**04011010**
					GRA	CAT	
Atopobium minutum		79.71	0.44	MLZ 1	99	1	
Fusobacterium varium		11.36	0.15		1	1	
Propionibacterium granulosum		3.14	0.22		99	90	
04011012	**Good identification**						**04011012**
					CAT		
Atopobium minutum		98.55	0.41	MLZ 1	1		
Propionibacterium granulosum		1.46	0.12		90		
04011020	**Very good identification**						**04011020**
					COC	CAT	
Atopobium minutum		95.98	0.70		1	1	
Propionibacterium acnes		1.94	0.34		1	95	
Gemella morbillorum		1.15	0.32		99	1	
04011021	**Acceptable identification**						**04011021**
					COC	CAT	
Atopobium minutum		94.55	0.36	URE 1	1	1	
Anaerococcus prevotii		3.54	-.03		99	40	
Propionibacterium acnes		1.91	-.01		1	95	
04011022	**Good identification**						**04011022**
					GRA	CAT	
Atopobium minutum		92.96	0.67		99	1	
Pseudoflavonifractor capillosus		6.87	0.33		1	1	
Propionibacterium granulosum		0.08	0.16		99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04011030	**Good identification**						**04011030**
					CAT		
Atopobium minutum		95.26	0.36	MLZ 1	1		
Propionibacterium granulosum		2.81	0.12		90		
Propionibacterium acnes		1.93	-.01		95		
04011040	**Low discrimination**						**04011040**
					COC	GRA	
Atopobium minutum		84.36	0.44	SOR 1	1	99	
Fusobacterium varium		12.02	0.15		1	1	
Gemella morbillorum		3.04	0.14		99	99	
04011042	**Good identification**						**04011042**
Atopobium minutum		100.0	0.41	SOR 1			
04011060	**Low discrimination**						**04011060**
					CAT		
Atopobium minutum		69.45	0.36	SOR 1	1		
Propionibacterium acnes		30.55	0.22		95		
04011100	**Low discrimination**						**04011100**
					GRA		
Atopobium minutum		60.86	0.44	LAC 1	99		
Fusobacterium mortiferum		17.93	0.40		1		
Fusobacterium varium		8.67	0.15		1		
04011102	**Low discrimination**						**04011102**
					GRA		
Atopobium minutum		85.74	0.41	LAC 1	99		
Lactobacillus fermentum		10.05	0.32		99		
Fusobacterium mortiferum		4.21	0.24		1		
04011120	**Low discrimination**						**04011120**
					GRA	SPO	
Atopobium minutum		62.71	0.36	LAC 1	99	1	
Clostridium septicum		13.35	0.12		99	99	
Prevotella bivia		9.18	0.03		1	1	
04011200	**Low discrimination**						**04011200**
					COC	GRA	
Atopobium minutum		83.73	0.44	RHA 1	1	99	
Fusobacterium varium		11.93	0.15		1	1	
Gemella morbillorum		3.02	0.14		99	99	
04011202	**Good identification**						**04011202**
Atopobium minutum		100.0	0.41	RHA 1			
04011220	**Good identification**						**04011220**
					CAT		
Atopobium minutum		98.02	0.36	RHA 1	1		
Propionibacterium acnes		1.98	-.01		95		
04011400	**Low discrimination**						**04011400**
					COC	GRA	
Atopobium minutum		62.76	0.44	ARA 1	1	99	
Gemella morbillorum		24.86	0.32		99	99	
Fusobacterium varium		8.94	0.15		1	1	
04011402	**Acceptable identification**						**04011402**
Atopobium minutum		93.30	0.41	ARA 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		6.70	0.28				
04011420	**Low discrimination**						**04011420**
					COC		
Atopobium minutum		83.32	0.36	ARA 1	1		
Gemella morbillorum		11.00	0.16		99		
Lactobacillus fermentum		3.99	0.20		1		
04012000	**Low discrimination**						**04012000**
					GRA	SPO	
Eubacterium limosum		73.32	0.63	GLU 99	99	1	
Fusobacterium varium		15.40	0.50		1	1	
Clostridium difficile		5.92	0.46		99	99	
04012002	**Low discrimination**						**04012002**
					GRA		
Atopobium minutum		64.44	0.49	MAN 1	99		
Eubacterium limosum		21.86	0.29		99		
Fusobacterium varium		4.59	0.15		1		
04012010	**Very good identification**						**04012010**
					GRA	SPO	
Clostridium difficile		96.76	0.58	GLU 99	99	99	
Eubacterium limosum		2.48	0.29		99	1	
Fusobacterium varium		0.52	0.15		1	1	
04012020	**Low discrimination**						**04012020**
					CAT		
Atopobium minutum		39.95	0.44	MAN 1	1		
Eubacterium limosum		27.10	0.29		1		
Propionibacterium acnes		15.01	0.29		95		
04012022	**Low discrimination**						**04012022**
					GRA		
Atopobium minutum		78.53	0.41	MAN 1	99		
Pseudoflavonifractor capillosus		17.42	0.15		1		
Pseudoramibacter alactolyticus		1.64	-.02		99		
04012050	**Good identification**						**04012050**
					SPO		
Clostridium difficile		99.51	0.36	GLU 99	99		
Eubacterium limosum		0.49	-.06	SOR 5	1		
04013000	**Low discrimination**						**04013000**
					CAT		
Atopobium minutum		46.65	0.44	MAN 1	1		
Eubacterium limosum		31.64	0.29		1		
Propionibacterium granulosum		10.16	0.35		90		
04013002	**Acceptable identification**						**04013002**
					CAT		
Atopobium minutum		91.58	0.41	MAN 1	1		
Propionibacterium granulosum		7.48	0.24		90		
Eubacterium limosum		0.94	-.06		1		
04014000	**Good identification**						**04014000**
					COC	GRA	
Fusobacterium varium		92.50	0.60	XYL 4	1	1	
Atopobium minutum		4.77	0.52		1	99	
Gemella morbillorum		1.17	0.37		99	99	
04014002	**Low discrimination**						**04014002**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		74.79	0.49	XYL 1	1	99	
Fusobacterium varium		21.98	0.26		1	1	
Anaerococcus prevotii		1.28	0.05		99	99	
04014020	**Low discrimination**						**04014020**
					COC	GRA	
Atopobium minutum		57.45	0.44	XYL 1	1	99	
Fusobacterium varium		33.76	0.26		1	1	
Gemella morbillorum		4.70	0.20		99	99	
04014022	**Low discrimination**						**04014022**
					GRA		
Atopobium minutum		81.25	0.41	XYL 1	99		
Pseudoflavonifractor capillosus		18.02	0.15		1		
Fusobacterium varium		0.72	-.08		1		
04015000	**Low discrimination**						**04015000**
					COC	GRA	
Atopobium minutum		48.32	0.44	XYL 1	1	99	
Fusobacterium varium		28.40	0.26		1	1	
Gemella morbillorum		19.14	0.32		99	99	
04015002	**Good identification**						**04015002**
					GRA		
Atopobium minutum		95.14	0.41	XYL 1	99		
Lactobacillus fermentum		4.01	0.24		99		
Fusobacterium varium		0.85	-.08		1		
04015020	**Low discrimination**						**04015020**
					COC		
Atopobium minutum		83.46	0.36	XYL 1	1		
Gemella morbillorum		11.02	0.16		99		
Lactobacillus fermentum		2.35	0.16		1		
04016010	**Good identification**						**04016010**
					GRA	SPO	
Clostridium difficile		99.09	0.36	GLU 99	99	99	
Eubacterium limosum		0.49	-.06	XYL 5	99	1	
Fusobacterium varium		0.42	-.08		1	1	
04020000	**Low discrimination**						**04020000**
Parvimonas micra		36.11	0.50	MNS 10			
Anaerococcus prevotii		16.93	0.50	TRE 1			
Peptococcus niger		8.55	0.40				
04020001	**Very good identification**						**04020001**
					COC	GRA	
Anaerococcus prevotii		97.37	0.66	TRE 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04020002	**Low discrimination**						**04020002**
					COC		
Anaerococcus prevotii		31.73	0.39	TRE 1	99		
Parvimonas micra		30.07	0.33	URE 90	99		
Atopobium minutum		20.23	0.50		1		
04020003	**Good identification**						**04020003**
Anaerococcus prevotii		99.37	0.56	TRE 1			
Peptococcus niger		0.39	0.10				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parvimonas micra		0.11	-.01				
04020020	**Low discrimination**						**04020020**
					COC	SPO	
Atopobium minutum		37.73	0.44	TRE 1	1	1	
Peptococcus niger		26.56	0.24		99	1	
Clostridium cadaveris		10.32	0.24		1	99	
04020040	**Low discrimination**						**04020040**
					COC	SPO	
Clostridium botulinum biovar A		70.73	0.41	GLU 90	1	99	
Parvimonas micra		11.29	0.15	MNS 7	99	1	
Anaerococcus prevotii		5.29	0.15		99	1	
04021000	**Low discrimination**						**04021000**
					COC		
Atopobium minutum		56.95	0.44	TRE 1	1		
Parvimonas micra		15.39	0.15		99		
Anaerococcus prevotii		7.22	0.15		99		
04021002	**Acceptable identification**						**04021002**
					COC		
Atopobium minutum		90.13	0.41	TRE 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04021020	**Good identification**						**04021020**
					COC		
Atopobium minutum		97.91	0.36	TRE 1	1		
Peptococcus niger		2.09	-.11		99		
04030000	**Low discrimination**						**04030000**
					GRA	CAT	
Fusobacterium varium		72.53	0.50	TRE 1	1	1	
Atopobium minutum		15.43	0.52		99	1	
Propionibacterium granulosum		4.45	0.45		99	90	
04030002	**Low discrimination**						**04030002**
					GRA	CAT	
Atopobium minutum		81.21	0.49	TRE 1	99	1	
Propionibacterium granulosum		8.79	0.35		99	90	
Fusobacterium varium		5.79	0.15		1	1	
04030020	**Low discrimination**						**04030020**
					SPO	CAT	
Atopobium minutum		59.63	0.44	TRE 1	1	1	
Clostridium cadaveris		17.66	0.24		99	1	
Propionibacterium granulosum		12.91	0.35		1	90	
04030022	**Low discrimination**						**04030022**
					GRA	CAT	
Atopobium minutum		76.74	0.41	TRE 1	99	1	
Pseudoflavonifractor capillosus		17.02	0.15		1	1	
Propionibacterium granulosum		6.23	0.24		99	90	
04031000	**Low discrimination**						**04031000**
					GRA	CAT	
Atopobium minutum		47.86	0.44	TRE 1	99	1	
Propionibacterium granulosum		41.46	0.45		99	90	
Fusobacterium varium		6.82	0.15		1	1	
04031002	**Low discrimination**						**04031002**
					CAT		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		75.48	0.41	TRE 1	1		
Propionibacterium granulosum		24.52	0.35		90		
04031020	**Low discrimination**						**04031020**
					CAT		
Atopobium minutum		59.88	0.36	TRE 1	1		
Propionibacterium granulosum		38.90	0.35		90		
Propionibacterium acnes		1.21	-.01		95		
04032010	**Good identification**						**04032010**
					SPO	CAT	
Clostridium difficile		95.41	0.38	GLU 99	99	1	
Propionibacterium granulosum		4.26	0.28	TRE 7	1	90	
Eubacterium limosum		0.33	-.06		1	1	
04033000	**Good identification**						**04033000**
					CAT		
Propionibacterium granulosum		97.47	0.43	GLU 99	90		
Atopobium minutum		1.51	0.10	SUC 93	1		
Eubacterium limosum		1.02	-.06		1		
04040000	**Low discrimination**						**04040000**
Parvimonas micra		39.19	0.50	MNS 10			
Anaerococcus prevotii		18.38	0.50	CEL 1			
Peptococcus niger		9.28	0.40				
04040001	**Very good identification**						**04040001**
					COC	GRA	
Anaerococcus prevotii		97.42	0.66	CEL 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04040002	**Low discrimination**						**04040002**
					COC		
Anaerococcus prevotii		32.34	0.39	CEL 1	99		
Parvimonas micra		30.65	0.33	URE 90	99		
Atopobium minutum		20.62	0.50		1		
04040003	**Good identification**						**04040003**
Anaerococcus prevotii		99.37	0.56	CEL 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				
04040020	**Low discrimination**						**04040020**
					COC		
Atopobium minutum		42.16	0.44	CEL 1	1		
Peptococcus niger		29.68	0.24		99		
Parvimonas micra		11.39	0.15		99		
04041000	**Low discrimination**						**04041000**
					COC		
Atopobium minutum		59.16	0.44	CEL 1	1		
Parvimonas micra		15.99	0.15		99		
Anaerococcus prevotii		7.50	0.15		99		
04041002	**Acceptable identification**						**04041002**
					COC		
Atopobium minutum		90.13	0.41	CEL 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04041020	**Good identification**						**04041020**
					COC		
Atopobium minutum		97.91	0.36	CEL 1	1		
Peptococcus niger		2.09	-.11		99		
04050000	**Low discrimination**						**04050000**
					COC	GRA	
Fusobacterium varium		77.84	0.50	CEL 1	1	1	
Atopobium minutum		16.56	0.52		1	99	
Parvimonas micra		1.62	0.15		99	99	
04050002	**Acceptable identification**						**04050002**
					COC	GRA	
Atopobium minutum		90.03	0.49	CEL 1	1	99	
Fusobacterium varium		6.41	0.15		1	1	
Anaerococcus prevotii		1.55	0.05		99	99	
04050020	**Low discrimination**						**04050020**
					GRA	SPO	
Atopobium minutum		81.89	0.44	CEL 1	99	1	
Fusobacterium varium		11.67	0.15		1	1	
Clostridium cadaveris		4.66	0.12		99	99	
04050022	**Low discrimination**						**04050022**
					GRA		
Atopobium minutum		81.85	0.41	CEL 1	99		
Pseudoflavonifractor capillosus		18.16	0.15		1		
04051000	**Low discrimination**						**04051000**
					GRA		
Atopobium minutum		81.47	0.44	CEL 1	99		
Fusobacterium varium		11.61	0.15		1		
Fusobacterium mortiferum		3.43	0.26		1		
04051002	**Good identification**						**04051002**
Atopobium minutum		100.0	0.41	CEL 1			
04051020	**Good identification**						**04051020**
					CAT		
Atopobium minutum		98.02	0.36	CEL 1	1		
Propionibacterium acnes		1.98	-.01		95		
04052010	**Good identification**						**04052010**
					SPO		
Clostridium difficile		99.51	0.36	GLU 99	99		
Eubacterium limosum		0.49	-.06	CEL 5	1		
04100000	**Low discrimination**						**04100000**
					COC	GRA	
Peptostreptococcus anaerobius		46.36	0.72	MNS 4	99	99	
Prevotella disiens		23.98	0.66		1	1	
Gemella morbillorum		6.14	0.64		99	99	
04100001	**Good identification**						**04100001**
					COC	GRA	
Anaerococcus prevotii		93.93	0.66	MLT 1	99	99	
Peptostreptococcus anaerobius		1.80	0.38		99	99	
Campylobacter ureolyticus		1.56	0.32		1	1	
04100002	**Low discrimination**						**04100002**
Anaerococcus prevotii		21.79	0.39	MLT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parvimonas micra		20.65	0.33	URE 90				
Peptostreptococcus anaerobius		15.02	0.38					
04100003	**Good identification**							**04100003**
Anaerococcus prevotii		99.16	0.56	MLT 1				
Peptococcus niger		0.39	0.10					
Parvimonas micra		0.11	-.01					
04100010	**Low discrimination**							**04100010**
					COC	GRA		
Peptostreptococcus anaerobius		45.40	0.38	MNS 4	99	99		
Prevotella disiens		23.48	0.32	MLZ 1	1	1		
Gemella morbillorum		6.01	0.30		99	99		
04100020	**Low discrimination**							**04100020**
					COC	GRA		
Gemella morbillorum		24.30	0.48	GLU 99	99	99		
Prevotella bivia		17.74	0.32	NAG 10	1	1		
Peptostreptococcus anaerobius		16.69	0.38		99	99		
04100040	**Low discrimination**							**04100040**
					COC	GRA	SPO	
Clostridium botulinum biovar A		48.26	0.49	GLU 90	1	99	99	
Peptostreptococcus anaerobius		18.82	0.38	MNS 7	99	99	1	
Prevotella disiens		9.73	0.32		1	1	1	
04100120	**Very good identification**							**04100120**
					COC	GRA	SPO	
Prevotella bivia		99.96	0.66	GLU 99	1	1	1	
Gemella morbillorum		0.01	0.14		99	99	1	
Clostridium bifermentans		0.01	0.05		1	99	99	
04100200	**Low discrimination**							**04100200**
					COC	GRA		
Peptostreptococcus anaerobius		46.43	0.38	MNS 4	99	99		
Prevotella disiens		24.01	0.32	RHA 1	1	1		
Gemella morbillorum		6.15	0.30		99	99		
04100400	**Low discrimination**							**04100400**
					COC	GRA		
Gemella morbillorum		41.87	0.48	GLU 99	99	99		
Peptostreptococcus anaerobius		28.76	0.38	ARA 10	99	99		
Prevotella disiens		14.87	0.32		1	1		
04101000	**Low discrimination**							**04101000**
					COC	GRA		
Gemella morbillorum		71.79	0.60	GLU 99	99	99		
Peptostreptococcus anaerobius		10.18	0.38		99	99		
Prevotella disiens		5.26	0.32		1	1		
04101002	**Low discrimination**							**04101002**
					COC			
Atopobium minutum		70.88	0.41	MLT 1	1			
Gemella morbillorum		16.38	0.26		99			
Anaerococcus prevotii		3.37	0.05		99			
04101020	**Low discrimination**							**04101020**
					COC	GRA		
Gemella morbillorum		67.94	0.43	GLU 99	99	99		
Prevotella bivia		16.25	0.19	NAG 10	1	1		
Atopobium minutum		13.36	0.36		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04101120	**Good identification**						**04101120**
					GRA		
Prevotella bivia		99.57	0.53	GLU 99	1		
Prevotella melaninogenica		0.32	0.17		1		
Clostridium perfringens		0.05	0.10		99		
04101400	**Good identification**						**04101400**
					COC	GRA	
Gemella morbillorum		97.15	0.43	GLU 99	99	99	
Peptostreptococcus anaerobius		1.25	0.04	ARA 10	99	99	
Prevotella disiens		0.65	-.03		1	1	
04102000	**Low discrimination**						**04102000**
					COC	GRA	
Peptostreptococcus anaerobius		46.27	0.38	MNS 4	99	99	
Prevotella disiens		23.92	0.32	MAN 1	1	1	
Gemella morbillorum		6.12	0.30		99	99	
04104000	**Low discrimination**						**04104000**
					COC	GRA	
Gemella morbillorum		41.84	0.48	GLU 99	99	99	
Peptostreptococcus anaerobius		28.74	0.38	XYL 10	99	99	
Prevotella disiens		14.86	0.32		1	1	
04105000	**Good identification**						**04105000**
					COC	GRA	
Gemella morbillorum		97.15	0.43	GLU 99	99	99	
Peptostreptococcus anaerobius		1.25	0.04	XYL 10	99	99	
Prevotella disiens		0.65	-.03		1	1	
04110000	**Low discrimination**						**04110000**
					COC	GRA	
Prevotella disiens		48.14	0.66	GLU 99	1	1	
Prevotella intermedia		27.32	0.69		1	1	
Gemella morbillorum		15.06	0.66		99	99	
04110002	**Low discrimination**						**04110002**
					GRA		
Atopobium minutum		42.17	0.49	MLT 1	99		
Prevotella disiens		25.54	0.32		1		
Prevotella intermedia		14.50	0.35		1		
04110020	**Low discrimination**						**04110020**
					COC	GRA	
Gemella morbillorum		52.35	0.50	GLU 99	99	99	
Prevotella disiens		15.21	0.32	NAG 10	1	1	
Atopobium minutum		12.56	0.44		1	99	
04110022	**Low discrimination**						**04110022**
					COC	GRA	
Atopobium minutum		75.42	0.41	MLT 1	1	99	
Pseudoflavonifractor capillosus		16.73	0.15		1	1	
Gemella morbillorum		4.76	0.15		99	99	
04110040	**Low discrimination**						**04110040**
					GRA	SPO	
Clostridium botulinum biovar B		49.08	0.41	GLU 99	99	99	
Prevotella disiens		19.95	0.32	SUC 90	1	1	
Prevotella intermedia		11.32	0.35		1	1	
04110050	**Good identification**						**04110050**
					GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		98.15	0.36	GLU 99	99	99		
Prevotella disiens		0.82	-.03	SUC 90	1	1		
Clostridium novyi biovar A		0.56	0.12		99	99		
04110060	**Good identification**							**04110060**
					COC	GRA	SPO	
Clostridium botulinum biovar B		95.36	0.36	GLU 99	1	99	99	
Gemella morbillorum		2.74	0.15	SUC 90	99	99	1	
Prevotella disiens		0.80	-.03		1	1	1	
04110120	**Good identification**							**04110120**
					COC	GRA		
Prevotella bivia		99.54	0.50	GLU 99	1	1		
Gemella morbillorum		0.15	0.15	FRU 10	99	99		
Clostridium perfringens		0.08	0.10		1	99		
04110400	**Low discrimination**							**04110400**
					COC	GRA		
Gemella morbillorum		66.27	0.50	GLU 99	99	99		
Prevotella disiens		19.26	0.32	ARA 10	1	1		
Prevotella intermedia		10.93	0.35		1	1		
04111000	**Low discrimination**							**04111000**
					COC	GRA		
Prevotella intermedia		74.80	0.69	GLU 90	1	1		
Gemella morbillorum		22.20	0.61		99	99		
Prevotella disiens		1.33	0.32		1	1		
04111001	**Low discrimination**							**04111001**
					COC	GRA	SPO	
Prevotella intermedia		71.02	0.35	GLU 90	1	1	1	
Gemella morbillorum		21.08	0.27	URE 1	99	99	1	
Clostridium sordellii		4.31	0.07		1	99	99	
04111002	**Low discrimination**							**04111002**
					COC	GRA		
Prevotella intermedia		39.36	0.35	GLU 90	1	1		
Atopobium minutum		38.17	0.41	BGL 1	1	99		
Gemella morbillorum		11.68	0.27		99	99		
04111010	**Low discrimination**							**04111010**
					COC	GRA	CAT	
Prevotella intermedia		73.80	0.35	GLU 90	1	1	1	
Gemella morbillorum		21.90	0.27	MLZ 1	99	99	1	
Propionibacterium granulosum		1.90	0.16		1	99	90	
04111020	**Low discrimination**							**04111020**
					COC	GRA		
Gemella morbillorum		60.23	0.45	GLU 99	99	99		
Prevotella intermedia		18.45	0.35	NAG 10	1	1		
Atopobium minutum		8.94	0.36		1	99		
04111040	**Low discrimination**							**04111040**
					COC	GRA	SPO	
Prevotella intermedia		72.83	0.35	GLU 90	1	1	1	
Gemella morbillorum		21.62	0.27	SOR 1	99	99	1	
Clostridium botulinum biovar B		3.19	0.07		1	99	99	
04111100	**Low discrimination**							**04111100**
					COC	GRA		
Prevotella intermedia		48.45	0.35	GLU 90	1	1		
Lactobacillus fermentum		16.52	0.40	LAC 1	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		14.38	0.27		99	99		
04111102	**Low discrimination**							**04111102**
					GRA			
Lactobacillus fermentum		84.02	0.40	GLU 99	99			
Clostridium perfringens		11.08	0.13	SUC 87	99			
Prevotella intermedia		2.49	0.01		1			
04111120	**Low discrimination**							**04111120**
					GRA			
Prevotella bivia		51.22	0.37	GLU 99	1			
Clostridium perfringens		22.52	0.44	FRU 10	99			
Prevotella melaninogenica		13.19	0.33		1			
04111122	**Low discrimination**							**04111122**
					GRA			
Clostridium perfringens		71.42	0.38	GLU 99	99			
Prevotella melaninogenica		16.94	0.21	SUC 91	1			
Lactobacillus fermentum		5.58	0.32		99			
04111200	**Low discrimination**							**04111200**
					COC	GRA		
Prevotella intermedia		75.23	0.35	GLU 90	1	1		
Gemella morbillorum		22.33	0.27	RHA 1	99	99		
Prevotella disiens		1.34	-.03		1	1		
04111400	**Low discrimination**							**04111400**
					COC	GRA		
Gemella morbillorum		72.31	0.45	GLU 99	99	99		
Prevotella intermedia		22.15	0.35	ARA 10	1	1		
Lactobacillus fermentum		4.63	0.36		1	99		
04111402	**Low discrimination**							**04111402**
					COC	GRA		
Lactobacillus fermentum		79.81	0.36	GLU 99	1	99		
Gemella morbillorum		12.59	0.11	SUC 87	99	99		
Prevotella intermedia		3.86	0.01		1	1		
04111500	**Low discrimination**							**04111500**
					COC			
Lactobacillus fermentum		81.39	0.40	GLU 99	1			
Actinomyces meyeri		8.33	0.12	SUC 87	1			
Gemella morbillorum		7.87	0.11		99			
04111502	**Good identification**							**04111502**
Lactobacillus fermentum		100.0	0.40	GLU 99				
04113000	**Low discrimination**							**04113000**
					COC	GRA	CAT	
Prevotella intermedia		67.49	0.35	GLU 90	1	1	1	
Gemella morbillorum		20.03	0.27	MAN 1	99	99	1	
Propionibacterium granulosum		9.61	0.29		1	99	90	
04114000	**Low discrimination**							**04114000**
					COC	GRA		
Gemella morbillorum		61.30	0.50	GLU 99	99	99		
Prevotella disiens		17.82	0.32	XYL 10	1	1		
Prevotella intermedia		10.11	0.35		1	1		
04115000	**Low discrimination**							**04115000**
					COC	GRA		
Gemella morbillorum		72.61	0.45	GLU 99	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		22.24	0.35	XYL 10	1	1		
Lactobacillus fermentum		2.73	0.32		1	99		
04115102	**Good identification**							**04115102**
Lactobacillus fermentum		95.98	0.36	GLU 99				
Actinomyces meyeri		4.02	0.02	SUC 87				
04115500	**Low discrimination**							**04115500**
Lactobacillus fermentum		74.72	0.36	GLU 99				
Actinomyces meyeri		25.28	0.17	SUC 87				
04115502	**Good identification**							**04115502**
Lactobacillus fermentum		100.0	0.36	GLU 99				
04120000	**Low discrimination**							**04120000**
					COC	GRA	SPO	
Clostridium botulinum biovar A		47.20	0.48	GLU 90	1	99	99	
Peptostreptococcus anaerobius		22.00	0.38	MNS 7	99	99	1	
Prevotella disiens		11.37	0.32		1	1	1	
04120040	**Low discrimination**							**04120040**
					COC	SPO		
Clostridium botulinum biovar B		58.96	0.39	GLU 99	1	99		
Clostridium botulinum biovar A		39.61	0.39	SUC 90	1	99		
Peptostreptococcus anaerobius		0.62	0.04		99	1		
04130040	**Good identification**							**04130040**
					GRA	SPO		
Clostridium botulinum biovar B		98.73	0.50	GLU 99	99	99		
Clostridium botulinum biovar A		1.06	0.19	SUC 90	99	99		
Prevotella disiens		0.14	-.03		1	1		
04130050	**Good identification**							**04130050**
Clostridium botulinum biovar B		100.0	0.44	GLU 99				
04130060	**Good identification**							**04130060**
Clostridium botulinum biovar B		100.0	0.44	GLU 99				
04130070	**Good identification**							**04130070**
Clostridium botulinum biovar B		100.0	0.39	GLU 99				
04131000	**Low discrimination**							**04131000**
					COC	GRA	CAT	
Prevotella intermedia		51.66	0.35	GLU 90	1	1	1	
Propionibacterium granulosum		29.25	0.39	TRE 1	1	99	90	
Gemella morbillorum		15.33	0.27		99	99	1	
04131120	**Low discrimination**							**04131120**
					GRA	SPO		
Clostridium perfringens		53.57	0.49	GLU 99	99	84		
Clostridium septicum		45.30	0.41	SUC 91	99	99		
Prevotella bivia		0.66	0.03		1	1		
04131122	**Good identification**							**04131122**
Clostridium perfringens		96.46	0.43	GLU 99				
Clostridium septicum		3.54	0.12	SUC 91				
04131160	**Good identification**							**04131160**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		95.71	0.36	GLU 99			
Clostridium septicum		4.29	0.07	SUC 91			
04132000	**Acceptable identification**						**04132000**
					GRA	CAT	
Propionibacterium granulosum		91.86	0.37	GLU 99	99	90	
Prevotella disiens		3.83	-.03	SUC 93	1	1	
Prevotella intermedia		2.17	0.01		1	1	
04133000	**Good identification**						**04133000**
					GRA	CAT	
Propionibacterium granulosum		97.69	0.37	GLU 99	99	90	
Prevotella intermedia		2.31	0.01	SUC 93	1	1	
04140000	**Low discrimination**						**04140000**
					COC	GRA	
Peptostreptococcus anaerobius		46.43	0.38	MNS 4	99	99	
Prevotella disiens		24.01	0.32	CEL 1	1	1	
Gemella morbillorum		6.15	0.30		99	99	
04151000	**Low discrimination**						**04151000**
					COC	GRA	
Prevotella intermedia		69.86	0.35	GLU 90	1	1	
Gemella morbillorum		20.74	0.27	CEL 1	99	99	
Lactobacillus jensenii		7.13	0.19		1	99	
04151120	**Acceptable identification**						**04151120**
					GRA	SPO	
Clostridium septicum		92.74	0.41	GLU 99	99	99	
Prevotella melaninogenica		3.64	0.17		1	1	
Clostridium perfringens		2.33	0.20		99	84	
04171120	**Good identification**						**04171120**
					COC	SPO	
Clostridium septicum		98.00	0.49	GLU 99	1	99	
Clostridium perfringens		1.53	0.25		1	84	
Atopobium parvulum		0.47	0.16		99	1	
04200000	**Low discrimination**						**04200000**
Parvimonas micra		38.64	0.50	MNS 10			
Anaerococcus prevotii		18.12	0.50	SAL 1			
Peptococcus niger		9.15	0.40				
04200001	**Very good identification**						**04200001**
					COC	GRA	
Anaerococcus prevotii		97.40	0.66	SAL 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
04200002	**Low discrimination**						**04200002**
					COC		
Anaerococcus prevotii		32.30	0.39	SAL 1	99		
Parvimonas micra		30.62	0.33	URE 90	99		
Atopobium minutum		20.60	0.50		1		
04200003	**Good identification**						**04200003**
Anaerococcus prevotii		99.37	0.56	SAL 1			
Peptococcus niger		0.39	0.10				
Parvimonas micra		0.11	-.01				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04200020	**Low discrimination**						**04200020**
					COC		
Atopobium minutum		41.08	0.44	SAL 1	1		
Peptococcus niger		28.92	0.24		99		
Parvimonas micra		11.10	0.15		99		
04201000	**Low discrimination**						**04201000**
					COC		
Atopobium minutum		58.78	0.44	SAL 1	1		
Parvimonas micra		15.89	0.15		99		
Anaerococcus prevotii		7.45	0.15		99		
04201002	**Acceptable identification**						**04201002**
					COC		
Atopobium minutum		90.13	0.41	SAL 1	1		
Anaerococcus prevotii		4.28	0.05		99		
Parvimonas micra		4.06	-.01		99		
04201020	**Good identification**						**04201020**
					COC		
Atopobium minutum		97.91	0.36	SAL 1	1		
Peptococcus niger		2.09	-.11		99		
04210000	**Low discrimination**						**04210000**
					COC	GRA	
Fusobacterium varium		76.43	0.50	SAL 1	1	1	
Atopobium minutum		16.26	0.52		1	99	
Parvimonas micra		1.59	0.15		99	99	
04210002	**Low discrimination**						**04210002**
					COC	GRA	
Atopobium minutum		89.30	0.49	SAL 1	1	99	
Fusobacterium varium		6.36	0.15		1	1	
Anaerococcus prevotii		1.53	0.05		99	99	
04210020	**Low discrimination**						**04210020**
					GRA	SPO	
Atopobium minutum		81.89	0.44	SAL 1	99	1	
Fusobacterium varium		11.67	0.15		1	1	
Clostridium cadaveris		4.66	0.12		99	99	
04210022	**Low discrimination**						**04210022**
					GRA		
Atopobium minutum		81.85	0.41	SAL 1	99		
Pseudoflavonifractor capillosus		18.16	0.15		1		
04210100	**Low discrimination**						**04210100**
					GRA		
Fusobacterium mortiferum		68.81	0.42	GLU 99	1		
Fusobacterium varium		25.72	0.15		1		
Atopobium minutum		5.47	0.18		99		
04211000	**Low discrimination**						**04211000**
					GRA		
Atopobium minutum		69.91	0.44	SAL 1	99		
Fusobacterium mortiferum		17.13	0.39		1		
Fusobacterium varium		9.96	0.15		1		
04211002	**Good identification**						**04211002**
					GRA		
Atopobium minutum		96.08	0.41	SAL 1	99		
Fusobacterium mortiferum		3.92	0.22		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04211020	**Good identification**						**04211020**
					CAT		
Atopobium minutum		98.02	0.36	SAL 1	1		
Propionibacterium acnes		1.98	-.01		95		
04211100	**Good identification**						**04211100**
					GRA		
Fusobacterium mortiferum		98.26	0.45	GLU 99	1		
Atopobium minutum		1.74	0.10		99		
04212000	**Low discrimination**						**04212000**
					GRA	SPO	
Clostridium difficile		63.79	0.37	GLU 99	99	99	
Eubacterium limosum		28.27	0.29		99	1	
Fusobacterium varium		5.94	0.15		1	1	
04212010	**Good identification**						**04212010**
					SPO		
Clostridium difficile		99.91	0.48	GLU 99	99		
Eubacterium limosum		0.09	-.06		1		
04231100	**Good identification**						**04231100**
					GRA	SPO	
Fusobacterium mortiferum		95.20	0.36	GLU 99	1	1	
Clostridium septicum		4.80	-.04		99	99	
04243000	**Good identification**						**04243000**
Clostridium innocuum		100.0	0.37	GLU 99			
04243002	**Good identification**						**04243002**
Clostridium innocuum		100.0	0.37	GLU 99			
04251100	**Low discrimination**						**04251100**
					GRA	SPO	
Fusobacterium mortiferum		79.44	0.37	GLU 99	1	1	
Lactobacillus catenaformis		16.77	0.15		99	1	
Clostridium septicum		3.79	-.03		99	99	
04252000	**Acceptable identification**						**04252000**
					SPO		
Clostridium innocuum		92.26	0.37	GLU 99	99		
Clostridium difficile		7.14	0.15	GAL 95	99		
Eubacterium limosum		0.61	-.06		1		
04252002	**Good identification**						**04252002**
Clostridium innocuum		100.0	0.37	GLU 99			
04253000	**Very good identification**						**04253000**
Clostridium innocuum		100.0	0.59	GLU 99			
04253002	**Very good identification**						**04253002**
Clostridium innocuum		100.0	0.59	GLU 99			
04253010	**Good identification**						**04253010**
Clostridium innocuum		100.0	0.35	GLU 99			
04253012	**Good identification**						**04253012**
Clostridium innocuum		100.0	0.35	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
04253400	**Good identification**							**04253400**
Clostridium innocuum		100.0	0.46	GLU 99				
04253402	**Good identification**							**04253402**
Clostridium innocuum		100.0	0.46	GLU 99				
04257000	**Good identification**							**04257000**
Clostridium innocuum		100.0	0.37	GLU 99				
04257002	**Good identification**							**04257002**
Clostridium innocuum		100.0	0.37	GLU 99				
04271120	**Good identification**							**04271120**
Clostridium septicum		100.0	0.39	GLU 99				
04273000	**Good identification**							**04273000**
Clostridium innocuum		100.0	0.51	GLU 99				
04273002	**Good identification**							**04273002**
Clostridium innocuum		100.0	0.51	GLU 99				
04273400	**Good identification**							**04273400**
Clostridium innocuum		100.0	0.38	GLU 99				
04273402	**Good identification**							**04273402**
Clostridium innocuum		100.0	0.38	GLU 99				
04300000	**Low discrimination**							**04300000**
					COC	GRA	SPO	
Clostridium bifermentans		36.04	0.43	IND 90	1	99	99	
Peptostreptococcus anaerobius		31.16	0.38	GLU 92	99	99	1	
Prevotella disiens		16.11	0.32	SAL 10	1	1	1	
04311000	**Low discrimination**							**04311000**
					COC	GRA		
Prevotella intermedia		67.78	0.35	GLU 90	1	1		
Gemella morbillorum		20.12	0.27	SAL 1	99	99		
Lactobacillus jensenii		6.92	0.19		1	99		
04311120	**Low discrimination**							**04311120**
					GRA	SPO		
Clostridium septicum		88.25	0.43	GLU 99	99	99		
Clostridium perfringens		10.54	0.34		99	84		
Prevotella bivia		0.97	0.03		1	1		
04331120	**Acceptable identification**							**04331120**
					COC	SPO		
Clostridium septicum		92.93	0.51	GLU 99	1	99		
Clostridium perfringens		6.87	0.38		1	84		
Atopobium parvulum		0.20	0.11		99	1		
04351000	**Low discrimination**							**04351000**
					GRA			
Lactobacillus jensenii		83.76	0.41	GLU 99	99			
Lactobacillus catenaformis		15.17	0.29	SUC 90	99			
Fusobacterium mortiferum		0.63	0.15		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04351002	**Low discrimination**						**04351002**
Lactobacillus jensenii		76.87	0.38	GLU 99			
Lactobacillus catenaformis		20.88	0.29	SUC 90			
Lactobacillus acidophilus		1.21	0.05				
04351120	**Good identification**						**04351120**
					SPO		
Clostridium septicum		98.07	0.51	GLU 99	99		
Clostridium paraputrificum		0.89	0.14		99		
Lactobacillus catenaformis		0.63	0.21		1		
04371000	**Good identification**						**04371000**
Lactobacillus jensenii		100.0	0.50	GLU 99			
04371002	**Good identification**						**04371002**
Lactobacillus jensenii		98.28	0.47	GLU 99			
Lactobacillus acidophilus		1.72	0.14	SUC 90			
04371010	**Good identification**						**04371010**
Lactobacillus jensenii		100.0	0.41	GLU 99			
04371012	**Good identification**						**04371012**
Lactobacillus jensenii		100.0	0.38	GLU 99			
04371020	**Low discrimination**						**04371020**
					SPO		
Lactobacillus jensenii		70.09	0.39	GLU 99	1		
Clostridium septicum		29.92	0.25	SUC 90	99		
04371022	**Acceptable identification**						**04371022**
					SPO		
Lactobacillus jensenii		92.32	0.36	GLU 99	1		
Lactobacillus acidophilus		6.47	0.14	SUC 90	1		
Clostridium septicum		1.21	-.04		99		
04371120	**Very good identification**						**04371120**
					COC	SPO	
Clostridium septicum		99.64	0.60	GLU 99	1	99	
Atopobium parvulum		0.16	0.18		99	1	
Clostridium perfringens		0.10	0.15		1	84	
04375100	**Good identification**						**04375100**
Atopobium parvulum		100.0	0.35	GLU 99			
04400000	**Low discrimination**						**04400000**
Anaerococcus prevotii		88.14	0.76	URE 90			
Parvimonas micra		5.70	0.50				
Peptococcus niger		1.35	0.40				
04400001	**Excellent identification**						**04400001**
					COC	GRA	
Anaerococcus prevotii		99.92	0.92		99	99	
Campylobacter ureolyticus		0.05	0.32		1	1	
Peptococcus niger		0.02	0.23		99	99	
04400002	**Good identification**						**04400002**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Anaerococcus prevotii		94.04	0.65	URE 90	99		
Parvimonas micra		2.70	0.33		99		
Atopobium minutum		1.82	0.50		1		
04400003	**Very good identification**						**04400003**
					COC		
Anaerococcus prevotii		99.98	0.82		99		
Peptococcus niger		0.01	0.10		99		
Atopobium minutum		0.00	0.15		1		
04400010	**Low discrimination**						**04400010**
Anaerococcus prevotii		88.31	0.41	MLZ 1			
Parvimonas micra		5.71	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04400011	**Very good identification**						**04400011**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	MLZ 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04400013	**Good identification**						**04400013**
Anaerococcus prevotii		100.0	0.47	MLZ 1			
04400020	**Low discrimination**						**04400020**
					COC		
Anaerococcus prevotii		65.07	0.41	NAG 1	99		
Atopobium minutum		15.56	0.44	URE 90	1		
Peptococcus niger		10.95	0.24		99		
04400021	**Very good identification**						**04400021**
					COC	GRA	
Anaerococcus prevotii		99.74	0.58	NAG 1	99	99	
Peptococcus niger		0.18	0.07		99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
04400023	**Good identification**						**04400023**
					COC	GRA	
Anaerococcus prevotii		99.67	0.47	NAG 1	99	99	
Pseudoflavonifractor capillosus		0.13	-.03		1	1	
Peptococcus niger		0.13	-.06		99	99	
04400040	**Low discrimination**						**04400040**
					COC	SPO	
Anaerococcus prevotii		87.06	0.41	SOR 1	99	1	
Parvimonas micra		5.63	0.15	URE 90	99	1	
Clostridium botulinum biovar A		1.43	0.17		1	99	
04400041	**Very good identification**						**04400041**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	SOR 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04400043	**Good identification**						**04400043**
Anaerococcus prevotii		100.0	0.47	SOR 1			
04400100	**Low discrimination**						**04400100**
Anaerococcus prevotii		88.31	0.41	LAC 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parvimonas micra		5.71	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04400101	**Very good identification**						**04400101**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	LAC 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04400103	**Good identification**						**04400103**
Anaerococcus prevotii		100.0	0.47	LAC 1			
04400200	**Low discrimination**						**04400200**
Anaerococcus prevotii		88.31	0.41	RHA 1			
Parvimonas micra		5.71	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04400201	**Very good identification**						**04400201**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	RHA 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04400203	**Good identification**						**04400203**
Anaerococcus prevotii		100.0	0.47	RHA 1			
04400400	**Low discrimination**						**04400400**
Anaerococcus prevotii		88.21	0.41	ARA 1			
Parvimonas micra		5.70	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04400401	**Very good identification**						**04400401**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	ARA 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04400403	**Good identification**						**04400403**
Anaerococcus prevotii		100.0	0.47	ARA 1			
04401000	**Low discrimination**						**04401000**
					COC		
Anaerococcus prevotii		72.67	0.41	GAL 1	99		
Atopobium minutum		17.37	0.44	URE 90	1		
Parvimonas micra		4.70	0.15		99		
04401001	**Very good identification**						**04401001**
					COC	GRA	
Anaerococcus prevotii		99.91	0.58	GAL 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Atopobium minutum		0.03	0.10		1	99	
04401003	**Good identification**						**04401003**
					COC		
Anaerococcus prevotii		99.93	0.47	GAL 1	99		
Atopobium minutum		0.07	0.07		1		
04401020	**Low discrimination**						**04401020**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		87.10	0.36	RAF 1	1		
Anaerococcus prevotii		11.04	0.07		99		
Peptococcus niger		1.86	-.11		99		
04402000	**Low discrimination**						**04402000**
Anaerococcus prevotii		88.20	0.41	MAN 1			
Parvimonas micra		5.70	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04402001	**Very good identification**						**04402001**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	MAN 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04402003	**Good identification**						**04402003**
Anaerococcus prevotii		100.0	0.47	MAN 1			
04404000	**Low discrimination**						**04404000**
Anaerococcus prevotii		88.11	0.41	XYL 1			
Parvimonas micra		5.69	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04404001	**Very good identification**						**04404001**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	XYL 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04404003	**Good identification**						**04404003**
Anaerococcus prevotii		100.0	0.47	XYL 1			
04410000	**Low discrimination**						**04410000**
					COC	GRA	
Fusobacterium varium		58.74	0.50	RAF 1	1	1	
Anaerococcus prevotii		18.87	0.41		99	99	
Atopobium minutum		12.49	0.52		1	99	
04410001	**Very good identification**						**04410001**
					COC	GRA	
Anaerococcus prevotii		99.49	0.58	FRU 1	99	99	
Fusobacterium varium		0.35	0.15		1	1	
Atopobium minutum		0.07	0.18		1	99	
04410002	**Low discrimination**						**04410002**
					COC	GRA	
Atopobium minutum		58.13	0.49	RAF 1	1	99	
Anaerococcus prevotii		32.93	0.31		99	99	
Fusobacterium varium		4.14	0.15		1	1	
04410003	**Good identification**						**04410003**
					COC		
Anaerococcus prevotii		99.80	0.47	FRU 1	99		
Atopobium minutum		0.20	0.15		1		
04410020	**Low discrimination**						**04410020**
					GRA	CAT	
Atopobium minutum		73.03	0.44	RAF 1	99	1	
Fusobacterium varium		10.41	0.15		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		6.71	0.28		99	90	
04410022	**Low discrimination**						**04410022**
					GRA	CAT	
Atopobium minutum		78.53	0.41	RAF 1	99	1	
Pseudoflavonifractor capillosus		17.42	0.15		1	1	
Propionibacterium granulosum		2.71	0.18		99	90	
04410100	**Low discrimination**						**04410100**
					COC	GRA	
Fusobacterium mortiferum		62.48	0.42	GLU 99	1	1	
Fusobacterium varium		23.35	0.15		1	1	
Anaerococcus prevotii		7.50	0.07		99	99	
04411000	**Low discrimination**						**04411000**
					GRA	CAT	
Prevotella intermedia		35.00	0.38	GLU 90	1	1	
Atopobium minutum		33.94	0.44	MLT 99	99	1	
Propionibacterium granulosum		12.47	0.38		99	90	
04411002	**Low discrimination**						**04411002**
					CAT		
Atopobium minutum		75.26	0.41	RAF 1	1		
Propionibacterium granulosum		10.37	0.28		90		
Lactobacillus fermentum		8.82	0.32		1		
04411020	**Low discrimination**						**04411020**
					CAT		
Atopobium minutum		71.16	0.36	RAF 1	1		
Propionibacterium granulosum		19.61	0.28		90		
Lactobacillus fermentum		5.56	0.24		1		
04411100	**Low discrimination**						**04411100**
					GRA		
Fusobacterium mortiferum		79.44	0.45	GLU 99	1		
Lactobacillus fermentum		17.71	0.35		99		
Prevotella intermedia		1.45	0.04		1		
04411102	**Low discrimination**						**04411102**
					GRA		
Lactobacillus fermentum		64.47	0.35	GLU 99	99		
Fusobacterium mortiferum		32.13	0.29	SUC 87	1		
Atopobium minutum		3.41	0.07		99		
04411500	**Good identification**						**04411500**
					GRA		
Lactobacillus fermentum		95.67	0.35	GLU 99	99		
Fusobacterium mortiferum		4.34	0.11	SUC 87	1		
04411502	**Good identification**						**04411502**
Lactobacillus fermentum		100.0	0.35	GLU 99			
04413000	**Acceptable identification**						**04413000**
					GRA	CAT	
Propionibacterium granulosum		91.02	0.36	GLU 99	99	90	
Prevotella intermedia		3.42	0.04	SUC 93	1	1	
Atopobium minutum		3.32	0.10		99	1	
04420000	**Low discrimination**						**04420000**
Anaerococcus prevotii		87.07	0.41	TRE 1			
Parvimonas micra		5.63	0.15	URE 90			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptococcus niger		1.33	0.06				
04420001	**Very good identification**						**04420001**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	TRE 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04420003	**Good identification**						**04420003**
Anaerococcus prevotii		100.0	0.47	TRE 1			
04430000	**Low discrimination**						**04430000**
					GRA	CAT	
Propionibacterium granulosum		79.88	0.47	GLU 99	99	90	
Fusobacterium varium		10.32	0.15	SUC 93	1	1	
Fusobacterium mortiferum		3.54	0.27		1	5	
04430002	**Low discrimination**						**04430002**
					COC	CAT	
Propionibacterium granulosum		89.70	0.36	GLU 99	1	90	
Atopobium minutum		6.58	0.15	SUC 93	1	1	
Anaerococcus prevotii		3.72	-.03		99	40	
04430020	**Good identification**						**04430020**
					CAT		
Propionibacterium granulosum		96.47	0.36	GLU 99	90		
Atopobium minutum		3.54	0.10	SUC 93	1		
04431000	**Acceptable identification**						**04431000**
					GRA	CAT	
Propionibacterium granulosum		92.17	0.47	GLU 99	99	90	
Fusobacterium mortiferum		6.12	0.30	SUC 93	1	5	
Prevotella intermedia		0.87	0.04		1	1	
04431002	**Acceptable identification**						**04431002**
					GRA	CAT	
Propionibacterium granulosum		94.88	0.36	GLU 99	99	90	
Fusobacterium mortiferum		2.80	0.13	SUC 93	1	5	
Atopobium minutum		2.32	0.07		99	1	
04431020	**Good identification**						**04431020**
Propionibacterium granulosum		100.0	0.36	GLU 99			
04431100	**Acceptable identification**						**04431100**
					GRA	CAT	
Fusobacterium mortiferum		93.88	0.36	GLU 99	1	5	
Propionibacterium granulosum		6.12	0.13		99	90	
04432000	**Good identification**						**04432000**
					CAT		
Propionibacterium granulosum		99.18	0.45	GLU 99	90		
Eubacterium limosum		0.82	-.06	SUC 93	1		
04433000	**Good identification**						**04433000**
Propionibacterium granulosum		100.0	0.45	GLU 99			
04435322	**Good identification**						**04435322**
Parabacteroides distasonis		100.0	0.39	GLU 99			
04435722	**Good identification**						**04435722**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.36	GLU 99			
04440000	**Low discrimination**						**04440000**
Anaerococcus prevotii		88.31	0.41	CEL 1			
Parvimonas micra		5.71	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04440001	**Very good identification**						**04440001**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	CEL 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04440003	**Good identification**						**04440003**
Anaerococcus prevotii		100.0	0.47	CEL 1			
04451100	**Good identification**						**04451100**
Fusobacterium mortiferum		100.0	0.37	GLU 99			
04500000	**Low discrimination**						**04500000**
					COC	GRA	
Anaerococcus prevotii		43.30	0.41	MLT 1	99	99	
Peptostreptococcus anaerobius		22.39	0.38	URE 90	99	99	
Prevotella disiens		11.58	0.32		1	1	
04500001	**Very good identification**						**04500001**
					COC	GRA	
Anaerococcus prevotii		99.82	0.58	MLT 1	99	99	
Peptostreptococcus anaerobius		0.06	0.04		99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
04500003	**Good identification**						**04500003**
Anaerococcus prevotii		100.0	0.47	MLT 1			
04501000	**Low discrimination**						**04501000**
					COC	GRA	
Prevotella intermedia		79.28	0.38	GLU 90	1	1	
Gemella morbillorum		12.84	0.26	FRU 99	99	99	
Anaerococcus prevotii		3.52	0.07		99	99	
04510000	**Very good identification**						**04510000**
					COC	GRA	
Prevotella intermedia		98.04	0.72	GLU 90	1	1	
Prevotella disiens		1.16	0.32		1	1	
Gemella morbillorum		0.36	0.32		99	99	
04510001	**Low discrimination**						**04510001**
					COC	GRA	
Prevotella intermedia		70.86	0.38	GLU 90	1	1	
Anaerococcus prevotii		28.31	0.24	URE 1	99	99	
Prevotella disiens		0.84	-.03		1	1	
04510002	**Acceptable identification**						**04510002**
					GRA	CAT	
Prevotella intermedia		92.37	0.38	GLU 90	1	1	
Propionibacterium granulosum		3.70	0.22	BGL 1	99	90	
Atopobium minutum		1.81	0.15		99	1	
04510010	**Good identification**						**04510010**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		96.74	0.38	GLU 90	1	1		
Propionibacterium granulosum		2.11	0.18	MLZ 1	99	90		
Prevotella disiens		1.15	-.03		1	1		
04510020	**Acceptable identification**							**04510020**
					COC	GRA	CAT	
Prevotella intermedia		90.70	0.38	GLU 90	1	1	1	
Gemella morbillorum		3.70	0.15	NAG 1	99	99	1	
Propionibacterium granulosum		3.63	0.22		1	99	90	
04510040	**Good identification**							**04510040**
					GRA	SPO		
Prevotella intermedia		96.06	0.38	GLU 90	1	1		
Clostridium botulinum biovar B		2.80	0.07	SOR 1	99	99		
Prevotella disiens		1.14	-.03		1	1		
04510100	**Acceptable identification**							**04510100**
Prevotella intermedia		94.56	0.38	GLU 90				
Fusobacterium mortiferum		4.32	0.26	LAC 1				
Prevotella disiens		1.12	-.03					
04510200	**Good identification**							**04510200**
Prevotella intermedia		98.83	0.38	GLU 90				
Prevotella disiens		1.17	-.03	RHA 1				
04510400	**Acceptable identification**							**04510400**
					COC	GRA		
Prevotella intermedia		94.99	0.38	GLU 90	1	1		
Gemella morbillorum		3.88	0.15	ARA 1	99	99		
Prevotella disiens		1.13	-.03		1	1		
04511000	**Very good identification**							**04511000**
					COC	GRA		
Prevotella intermedia		99.36	0.72	GLU 90	1	1		
Lactobacillus fermentum		0.23	0.40		1	99		
Gemella morbillorum		0.20	0.27		99	99		
04511001	**Good identification**							**04511001**
Prevotella intermedia		100.0	0.38	GLU 90				
04511002	**Low discrimination**							**04511002**
					GRA	CAT		
Prevotella intermedia		78.50	0.38	GLU 90	1	1		
Lactobacillus fermentum		17.85	0.40	BGL 1	99	1		
Propionibacterium granulosum		3.14	0.22		99	90		
04511010	**Good identification**							**04511010**
					GRA	CAT		
Prevotella intermedia		97.86	0.38	GLU 90	1	1		
Propionibacterium granulosum		2.14	0.18	MLZ 1	99	90		
04511020	**Low discrimination**							**04511020**
					GRA			
Prevotella intermedia		76.86	0.38	GLU 90	1			
Prevotella melaninogenica		8.43	0.15	NAG 1	1			
Lactobacillus fermentum		5.83	0.32		99			
04511040	**Good identification**							**04511040**
Prevotella intermedia		100.0	0.38	GLU 90				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04511100	**Low discrimination**						**04511100**
					GRA		
Prevotella intermedia		63.66	0.38	GLU 90	1		
Lactobacillus fermentum		23.61	0.44	LAC 1	99		
Prevotella melaninogenica		6.98	0.15		1		
04511102	**Low discrimination**						**04511102**
					GRA		
Lactobacillus fermentum		87.68	0.44	GLU 99	99		
Prevotella melaninogenica		4.94	0.03	SUC 87	1		
Prevotella intermedia		2.39	0.04		1		
04511120	**Acceptable identification**						**04511120**
					GRA		
Prevotella melaninogenica		94.83	0.50	GLU 90	1		
Clostridium perfringens		3.43	0.32	SUC 99	99		
Lactobacillus fermentum		1.08	0.35		99		
04511122	**Low discrimination**						**04511122**
					GRA		
Prevotella melaninogenica		81.63	0.37	GLU 90	1		
Clostridium perfringens		7.29	0.26	SUC 99	99		
Prevotella oralis		5.86	0.21		1		
04511200	**Good identification**						**04511200**
Prevotella intermedia		100.0	0.38	GLU 90			
04511400	**Low discrimination**						**04511400**
					COC	GRA	
Prevotella intermedia		80.04	0.38	GLU 90	1	1	
Lactobacillus fermentum		18.20	0.40	ARA 1	1	99	
Gemella morbillorum		1.76	0.11		99	99	
04511402	**Good identification**						**04511402**
					GRA		
Lactobacillus fermentum		95.75	0.40	GLU 99	99		
Prevotella intermedia		4.25	0.04	SUC 87	1		
04511500	**Acceptable identification**						**04511500**
					GRA		
Lactobacillus fermentum		94.98	0.44	GLU 99	99		
Prevotella intermedia		2.59	0.04	SUC 87	1		
Bifidobacterium longum subsp. longum		2.44	0.09		99		
04511502	**Good identification**						**04511502**
Lactobacillus fermentum		97.50	0.44	GLU 99			
Bifidobacterium longum subsp. longum		2.50	0.09	SUC 87			
04511520	**Low discrimination**						**04511520**
					GRA		
Lactobacillus fermentum		45.56	0.35	GLU 99	99		
Prevotella melaninogenica		40.42	0.15	SUC 87	1		
Bifidobacterium longum subsp. longum		14.02	0.20		99		
04511522	**Low discrimination**						**04511522**
					GRA		
Lactobacillus fermentum		67.72	0.35	GLU 99	99		
Bifidobacterium longum subsp. longum		20.84	0.20	SUC 87	99		
Prevotella melaninogenica		11.45	0.03		1		
04512000	**Low discrimination**						**04512000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Prevotella intermedia		87.77	0.38	GLU 90	1	1	
Propionibacterium granulosum		10.60	0.30	MAN 1	99	90	
Prevotella disiens		1.04	-.03		1	1	
04513000	**Low discrimination**						**04513000**
					GRA	CAT	
Prevotella intermedia		89.22	0.38	GLU 90	1	1	
Propionibacterium granulosum		10.78	0.30	MAN 1	99	90	
04514000	**Acceptable identification**						**04514000**
					COC	GRA	
Prevotella intermedia		94.48	0.38	GLU 90	1	1	
Gemella morbillorum		3.86	0.15	XYL 1	99	99	
Prevotella disiens		1.12	-.03		1	1	
04515000	**Low discrimination**						**04515000**
					COC	GRA	
Prevotella intermedia		86.54	0.38	GLU 90	1	1	
Lactobacillus fermentum		11.56	0.36	XYL 1	1	99	
Gemella morbillorum		1.90	0.11		99	99	
04515002	**Acceptable identification**						**04515002**
					GRA		
Lactobacillus fermentum		92.97	0.36	GLU 99	99		
Prevotella intermedia		7.03	0.04	SUC 87	1		
04515100	**Good identification**						**04515100**
					GRA		
Lactobacillus fermentum		95.57	0.40	GLU 99	99		
Prevotella intermedia		4.43	0.04	SUC 87	1		
04515102	**Good identification**						**04515102**
Lactobacillus fermentum		100.0	0.40	GLU 99			
04515322	**Good identification**						**04515322**
Parabacteroides distasonis		96.82	0.37	GLU 99			
Prevotella oralis		3.18	-.01				
04515400	**Acceptable identification**						**04515400**
					GRA		
Lactobacillus fermentum		92.97	0.36	GLU 99	99		
Prevotella intermedia		7.03	0.04	SUC 87	1		
04515402	**Good identification**						**04515402**
Lactobacillus fermentum		100.0	0.36	GLU 99			
04515500	**Low discrimination**						**04515500**
Lactobacillus fermentum		88.42	0.40	GLU 99			
Bifidobacterium longum subsp. longum		11.58	0.17	SUC 87			
04515502	**Low discrimination**						**04515502**
Lactobacillus fermentum		88.42	0.40	GLU 99			
Bifidobacterium longum subsp. longum		11.58	0.17	SUC 87			
04515720	**Good identification**						**04515720**
Bacteroides vulgatus		100.0	0.36	GLU 99			
04530000	**Low discrimination**						**04530000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Prevotella intermedia		66.75	0.38	GLU 90	1	1	
Propionibacterium granulosum		32.07	0.41	TRE 1	99	90	
Prevotella disiens		0.79	-.03		1	1	
04531000	**Low discrimination**						**04531000**
					GRA	CAT	
Prevotella intermedia		67.15	0.38	GLU 90	1	1	
Propionibacterium granulosum		32.26	0.41	TRE 1	99	90	
Fusobacterium mortiferum		0.59	0.14		1	5	
04531120	**Low discrimination**						**04531120**
					GRA	SPO	
Clostridium perfringens		83.65	0.36	GLU 99	99	84	
Prevotella melaninogenica		12.60	0.15	SUC 91	1	1	
Clostridium septicum		3.75	0.07		99	99	
04532000	**Good identification**						**04532000**
					GRA	CAT	
Propionibacterium granulosum		97.29	0.39	GLU 99	99	90	
Prevotella intermedia		2.71	0.04	SUC 93	1	1	
04533000	**Good identification**						**04533000**
					GRA	CAT	
Propionibacterium granulosum		97.29	0.39	GLU 99	99	90	
Prevotella intermedia		2.71	0.04	SUC 93	1	1	
04535122	**Good identification**						**04535122**
Parabacteroides distasonis		100.0	0.41	GLU 99			
04535132	**Good identification**						**04535132**
Parabacteroides distasonis		100.0	0.36	GLU 99			
04535322	**Good identification**						**04535322**
Parabacteroides distasonis		100.0	0.49	GLU 99			
04535332	**Good identification**						**04535332**
Parabacteroides distasonis		100.0	0.44	GLU 99			
04535522	**Low discrimination**						**04535522**
					GRA		
Parabacteroides distasonis		88.65	0.37	GLU 99	1		
Bifidobacterium longum subsp. longum		11.35	0.17		99		
04535722	**Good identification**						**04535722**
Parabacteroides distasonis		98.73	0.45	GLU 99			
Clostridium clostridioforme		1.27	0.07				
04535732	**Acceptable identification**						**04535732**
Parabacteroides distasonis		90.19	0.40	GLU 99			
Clostridium clostridioforme		9.81	0.18				
04550000	**Good identification**						**04550000**
Prevotella intermedia		98.19	0.38	GLU 90			
Prevotella disiens		1.17	-.03	CEL 1			
Fusobacterium mortiferum		0.64	0.12				
04551000	**Good identification**						**04551000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		99.03	0.38	GLU 90			
Fusobacterium mortiferum		0.97	0.15	CEL 1			
04575122	**Good identification**						**04575122**
Parabacteroides distasonis		100.0	0.36	GLU 99			
04575322	**Good identification**						**04575322**
Parabacteroides distasonis		100.0	0.44	GLU 99			
04575332	**Good identification**						**04575332**
Parabacteroides distasonis		97.42	0.39	GLU 99			
Clostridium clostridioforme		2.58	0.07				
04575722	**Acceptable identification**						**04575722**
Parabacteroides distasonis		92.60	0.40	GLU 99			
Clostridium clostridioforme		7.40	0.16				
04575732	**Low discrimination**						**04575732**
Parabacteroides distasonis		59.65	0.36	GLU 99			
Clostridium clostridioforme		40.35	0.27				
04600000	**Low discrimination**						**04600000**
Anaerococcus prevotii		88.11	0.41	SAL 1			
Parvimonas micra		5.69	0.15	URE 90			
Peptococcus niger		1.35	0.06				
04600001	**Very good identification**						**04600001**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	SAL 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
04600003	**Good identification**						**04600003**
Anaerococcus prevotii		100.0	0.47	SAL 1			
04610000	**Low discrimination**						**04610000**
					COC	GRA	
Fusobacterium mortiferum		58.07	0.41	GLU 99	1	1	
Fusobacterium varium		26.09	0.15		1	1	
Anaerococcus prevotii		8.38	0.07		99	99	
04610100	**Good identification**						**04610100**
Fusobacterium mortiferum		100.0	0.47	GLU 99			
04611000	**Good identification**						**04611000**
					GRA		
Fusobacterium mortiferum		95.87	0.44	GLU 99	1		
Prevotella intermedia		2.10	0.04		1		
Atopobium minutum		2.04	0.10		99		
04611100	**Good identification**						**04611100**
Fusobacterium mortiferum		100.0	0.50	GLU 99			
04630100	**Good identification**						**04630100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.38	GLU 99		
04631100	**Good identification**					**04631100**
Fusobacterium mortiferum		100.0	0.41	GLU 99		
04635322	**Good identification**					**04635322**
Parabacteroides distasonis		100.0	0.40	GLU 99		
04635332	**Good identification**					**04635332**
Parabacteroides distasonis		100.0	0.35	GLU 99		
04635722	**Good identification**					**04635722**
Parabacteroides distasonis		100.0	0.36	GLU 99		
04650100	**Good identification**					**04650100**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
04651000	**Good identification**					**04651000**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
04651100	**Good identification**					**04651100**
Fusobacterium mortiferum		100.0	0.42	GLU 99		
04675322	**Good identification**					**04675322**
Parabacteroides distasonis		100.0	0.35	GLU 99		
04710000	**Good identification**					**04710000**
Prevotella intermedia		95.25	0.38	GLU 90		
Fusobacterium mortiferum		3.62	0.25	SAL 1		
Prevotella disiens		1.13	-.03			
04711000	**Acceptable identification**					**04711000**
Prevotella intermedia		94.61	0.38	GLU 90		
Fusobacterium mortiferum		5.40	0.28	SAL 1		
04715322	**Acceptable identification**					**04715322**
Parabacteroides distasonis		94.89	0.38	GLU 99		
Prevotella oralis		5.11	0.04			
04735122	**Good identification**					**04735122**
Parabacteroides distasonis		100.0	0.41	GLU 99		
04735132	**Good identification**					**04735132**
Parabacteroides distasonis		100.0	0.37	GLU 99		
04735322	**Good identification**					**04735322**
Parabacteroides distasonis		100.0	0.49	GLU 99		
04735332	**Good identification**					**04735332**
Parabacteroides distasonis		97.20	0.45	GLU 99		
Clostridium clostridioforme		2.80	0.13			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
04735522	**Good identification**						**04735522**
Parabacteroides distasonis		96.56	0.37	GLU 99			
Clostridium clostridioforme		3.44	0.07				
04735722	**Acceptable identification**						**04735722**
Parabacteroides distasonis		91.99	0.46	GLU 99			
Clostridium clostridioforme		8.01	0.22				
04735732	**Low discrimination**						**04735732**
Parabacteroides distasonis		57.59	0.41	GLU 99			
Clostridium clostridioforme		42.41	0.33				
04775122	**Good identification**						**04775122**
Parabacteroides distasonis		96.64	0.37	GLU 99			
Prevotella oralis		3.36	-.01				
04775322	**Good identification**						**04775322**
Parabacteroides distasonis		97.93	0.45	GLU 99			
Clostridium clostridioforme		2.07	0.10				
04775332	**Low discrimination**						**04775332**
Parabacteroides distasonis		84.81	0.40	GLU 99			
Clostridium clostridioforme		15.19	0.22				
04775712	**Good identification**						**04775712**
Clostridium clostridioforme		100.0	0.42	GLU 88			
04775722	**Low discrimination**						**04775722**
Parabacteroides distasonis		64.89	0.41	GLU 99			
Clostridium clostridioforme		35.11	0.31				
04775732	**Low discrimination**						**04775732**
Clostridium clostridioforme		82.07	0.42	GLU 88			
Parabacteroides distasonis		17.93	0.36	SUC 99			
05000000	**Low discrimination**						**05000000**
					COC		
Atopobium minutum		23.16	0.91		1		
Gemella morbillorum		22.67	0.86		99		
Peptostreptococcus anaerobius		17.97	0.76		99		
05000001	**Low discrimination**						**05000001**
					COC	SPO	
Anaerococcus prevotii		76.28	0.66	GLU 1	99	1	
Clostridium novyi biovar A		7.03	0.66		1	99	
Atopobium minutum		3.45	0.56		1	1	
05000002	**Good identification**						**05000002**
					COC		
Atopobium minutum		92.30	0.88		1		
Gemella morbillorum		1.37	0.52		99		
Peptostreptococcus anaerobius		1.09	0.42		99		
05000003	**Low discrimination**						**05000003**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Anaerococcus prevotii		83.49	0.56	GLU 1	99	1	
Atopobium minutum		10.07	0.53		1	1	
Clostridium novyi biovar A		5.43	0.53		1	99	
05000010	**Low discrimination**						**05000010**
					COC	SPO	
Clostridium novyi biovar A		29.66	0.66	MNS 1	1	99	
Atopobium minutum		14.56	0.56		1	1	
Gemella morbillorum		14.25	0.52		99	1	
05000011	**Low discrimination**						**05000011**
					COC	SPO	
Clostridium novyi biovar A		88.13	0.66	MNS 1	1	99	
Anaerococcus prevotii		9.66	0.32		99	1	
Atopobium minutum		0.44	0.22		1	1	
05000012	**Low discrimination**						**05000012**
					COC	SPO	
Atopobium minutum		61.57	0.53	MLZ 1	1	1	
Clostridium novyi biovar A		33.21	0.53		1	99	
Gemella morbillorum		0.91	0.17		99	1	
05000013	**Low discrimination**						**05000013**
					COC	SPO	
Clostridium novyi biovar A		85.18	0.53	MNS 1	1	99	
Anaerococcus prevotii		13.23	0.21		99	1	
Atopobium minutum		1.60	0.19		1	1	
05000020	**Low discrimination**						**05000020**
					COC	SPO	
Atopobium minutum		48.65	0.82		1	1	
Clostridium cadaveris		28.49	0.68		1	99	
Gemella morbillorum		15.87	0.69		99	1	
05000021	**Low discrimination**						**05000021**
					COC	SPO	
Clostridium novyi biovar A		35.73	0.56	MNS 1	1	99	
Atopobium minutum		23.38	0.48		1	1	
Anaerococcus prevotii		15.66	0.32		99	1	
05000022	**Very good identification**						**05000022**
					GRA	SPO	
Atopobium minutum		96.83	0.79		99	1	
Pseudoflavonifractor capillosus		1.06	0.32		1	1	
Clostridium cadaveris		0.86	0.34		99	99	
05000023	**Low discrimination**						**05000023**
					COC	SPO	
Atopobium minutum		59.13	0.45	URE 1	1	1	
Clostridium novyi biovar A		23.92	0.43		1	99	
Anaerococcus prevotii		14.85	0.21		99	1	
05000030	**Low discrimination**						**05000030**
					SPO		
Clostridium novyi biovar A		42.68	0.56	MNS 1	99		
Atopobium minutum		27.92	0.48		1		
Clostridium cadaveris		16.35	0.34		99		
05000031	**Good identification**						**05000031**
					COC	SPO	
Clostridium novyi biovar A		98.32	0.56	MNS 1	1	99	
Atopobium minutum		0.65	0.14		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Anaerococcus prevotii		0.44	-.03		99	1	
05000032	**Low discrimination**						**05000032**
					GRA	SPO	
Atopobium minutum		69.97	0.45	MLZ 1	99	1	
Clostridium novyi biovar A		28.30	0.43		99	99	
Pseudoflavonifractor capillosus		0.77	-.03		1	1	
05000033	**Good identification**						**05000033**
					SPO		
Clostridium novyi biovar A		97.56	0.43	MNS 1	99		
Atopobium minutum		2.44	0.11		1		
05000040	**Low discrimination**						**05000040**
					SPO		
Clostridium botulinum biovar A		65.53	0.67	MNS 7	99		
Clostridium novyi biovar A		7.46	0.63		99		
Atopobium minutum		5.49	0.56		1		
05000041	**Low discrimination**						**05000041**
					COC	SPO	
Clostridium novyi biovar A		77.35	0.63	MNS 1	1	99	
Anaerococcus prevotii		12.72	0.32		99	1	
Clostridium botulinum biovar A		6.87	0.33		1	99	
05000042	**Low discrimination**						**05000042**
					SPO		
Atopobium minutum		60.77	0.53	SOR 1	1		
Clostridium novyi biovar A		21.85	0.50		99		
Clostridium botulinum biovar A		10.99	0.33		99		
05000043	**Low discrimination**						**05000043**
					COC	SPO	
Clostridium novyi biovar A		78.99	0.50	MNS 1	1	99	
Anaerococcus prevotii		18.39	0.21		99	1	
Atopobium minutum		2.22	0.19		1	1	
05000050	**Low discrimination**						**05000050**
Clostridium novyi biovar A		84.01	0.63	MNS 1			
Clostridium botulinum biovar A		7.46	0.33				
Clostridium botulinum biovar B		4.10	0.26				
05000051	**Very good identification**						**05000051**
					COC	SPO	
Clostridium novyi biovar A		99.75	0.63	MNS 1	1	99	
Anaerococcus prevotii		0.17	-.03		99	1	
Clostridium botulinum biovar A		0.09	-.01		1	99	
05000052	**Good identification**						**05000052**
					SPO		
Clostridium novyi biovar A		96.79	0.50	MNS 1	99		
Atopobium minutum		2.72	0.19		1		
Clostridium botulinum biovar A		0.49	-.01		99		
05000053	**Good identification**						**05000053**
Clostridium novyi biovar A		100.0	0.50	MNS 1			
05000060	**Low discrimination**						**05000060**
					SPO		
Clostridium novyi biovar A		26.55	0.53	MNS 1	99		
Atopobium minutum		26.06	0.48		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium cadaveris		15.26	0.34		99		
05000061	**Good identification**						**05000061**
					COC	SPO	
Clostridium novyi biovar A		97.16	0.53	MNS 1	1	99	
Atopobium minutum		0.96	0.14		1	1	
Anaerococcus prevotii		0.65	-.03		99	1	
05000062	**Low discrimination**						**05000062**
					GRA	SPO	
Atopobium minutum		76.62	0.45	SOR 1	99	1	
Clostridium novyi biovar A		20.66	0.40		99	99	
Pseudoflavonifractor capillosus		0.84	-.03		1	1	
05000063	**Good identification**						**05000063**
					SPO		
Clostridium novyi biovar A		96.39	0.40	MNS 1	99		
Atopobium minutum		3.61	0.11		1		
05000070	**Low discrimination**						**05000070**
					SPO		
Clostridium novyi biovar A		89.39	0.53	MNS 1	99		
Clostridium botulinum biovar B		8.60	0.21		99		
Atopobium minutum		0.89	0.14		1		
05000071	**Good identification**						**05000071**
Clostridium novyi biovar A		100.0	0.53	MNS 1			
05000072	**Good identification**						**05000072**
					SPO		
Clostridium novyi biovar A		96.39	0.40	MNS 1	99		
Atopobium minutum		3.61	0.11		1		
05000073	**Good identification**						**05000073**
Clostridium novyi biovar A		100.0	0.40	MNS 1			
05000100	**Low discrimination**						**05000100**
					COC		
Atopobium minutum		20.80	0.56	LAC 1	1		
Gemella morbillorum		20.36	0.52		99		
Peptostreptococcus anaerobius		16.14	0.42		99		
05000102	**Acceptable identification**						**05000102**
					COC		
Atopobium minutum		91.55	0.53	LAC 1	1		
Gemella morbillorum		1.36	0.17		99		
Peptostreptococcus anaerobius		1.08	0.08		99		
05000120	**Very good identification**						**05000120**
					GRA	SPO	
Prevotella bivia		98.50	0.66	MLT 99	1	1	
Atopobium minutum		0.74	0.48		99	1	
Clostridium cadaveris		0.43	0.34		99	99	
05000200	**Low discrimination**						**05000200**
					COC		
Atopobium minutum		23.16	0.56	RHA 1	1		
Gemella morbillorum		22.67	0.52		99		
Peptostreptococcus anaerobius		17.98	0.42		99		
05000202	**Acceptable identification**						**05000202**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		92.63	0.53	RHA 1	1		
Gemella morbillorum		1.37	0.17		99		
Peptostreptococcus anaerobius		1.09	0.08		99		
05000220	**Low discrimination**						**05000220**
					COC	SPO	
Atopobium minutum		48.86	0.48	RHA 1	1	1	
Clostridium cadaveris		28.61	0.34		1	99	
Gemella morbillorum		15.94	0.35		99	1	
05000222	**Good identification**						**05000222**
					GRA	SPO	
Atopobium minutum		97.59	0.45	RHA 1	99	1	
Pseudoflavonifractor capillosus		1.07	-.03		1	1	
Clostridium cadaveris		0.87	0.00		99	99	
05000400	**Low discrimination**						**05000400**
					COC		
Gemella morbillorum		76.34	0.69	ARA 10	99		
Atopobium minutum		7.09	0.56		1		
Peptostreptococcus anaerobius		5.50	0.42		99		
05000402	**Low discrimination**						**05000402**
					COC		
Atopobium minutum		81.44	0.53	ARA 1	1		
Gemella morbillorum		13.29	0.35		99		
Peptostreptococcus anaerobius		0.96	0.08		99		
05000410	**Low discrimination**						**05000410**
					COC	SPO	
Gemella morbillorum		65.63	0.35	ARA 10	99	1	
Clostridium novyi biovar A		12.42	0.32	MLZ 1	1	99	
Atopobium minutum		6.10	0.22		1	1	
05000420	**Low discrimination**						**05000420**
					COC	SPO	
Gemella morbillorum		67.59	0.53	ARA 10	99	1	
Atopobium minutum		18.83	0.48	NAG 10	1	1	
Clostridium cadaveris		11.03	0.34		1	99	
05000422	**Acceptable identification**						**05000422**
					COC	GRA	
Atopobium minutum		93.09	0.45	ARA 1	1	99	
Gemella morbillorum		5.06	0.19		99	99	
Pseudoflavonifractor capillosus		1.02	-.03		1	1	
05000500	**Low discrimination**						**05000500**
					COC		
Gemella morbillorum		74.44	0.35	LAC 1	99		
Atopobium minutum		6.91	0.22	ARA 10	1		
Peptostreptococcus anaerobius		5.37	0.08		99		
05000600	**Low discrimination**						**05000600**
					COC		
Gemella morbillorum		77.82	0.35	RHA 1	99		
Atopobium minutum		7.23	0.22	ARA 10	1		
Peptostreptococcus anaerobius		5.61	0.08		99		
05001000	**Low discrimination**						**05001000**
					COC		
Gemella morbillorum		59.39	0.81		99		
Atopobium minutum		37.56	0.82		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptostreptococcus anaerobius		0.88	0.42		99		
05001001	**Low discrimination**						**05001001**
					COC	SPO	
Gemella morbillorum		38.99	0.47	URE 1	99	1	
Atopobium minutum		24.66	0.48		1	1	
Clostridium novyi biovar A		16.75	0.50		1	99	
05001002	**Very good identification**						**05001002**
					COC	SPO	
Atopobium minutum		97.21	0.79		1	1	
Gemella morbillorum		2.33	0.47		99	1	
Clostridium novyi biovar A		0.18	0.37		1	99	
05001003	**Low discrimination**						**05001003**
					COC	SPO	
Atopobium minutum		68.73	0.45	URE 1	1	1	
Anaerococcus prevotii		17.27	0.21		99	1	
Clostridium novyi biovar A		12.36	0.37		1	99	
05001010	**Low discrimination**						**05001010**
					COC	SPO	
Gemella morbillorum		47.05	0.47	MLZ 1	99	1	
Atopobium minutum		29.76	0.48		1	1	
Clostridium novyi biovar A		20.21	0.50		1	99	
05001011	**Good identification**						**05001011**
					COC	SPO	
Clostridium novyi biovar A		95.39	0.50	MNS 1	1	99	
Gemella morbillorum		2.24	0.13	GAL 10	99	1	
Atopobium minutum		1.42	0.14		1	1	
05001012	**Low discrimination**						**05001012**
					COC	SPO	
Atopobium minutum		82.53	0.45	MLZ 1	1	1	
Clostridium novyi biovar A		14.84	0.37		1	99	
Gemella morbillorum		1.98	0.13		99	1	
05001013	**Acceptable identification**						**05001013**
					SPO		
Clostridium novyi biovar A		94.68	0.37	MNS 1	99		
Atopobium minutum		5.32	0.11	GAL 10	1		
05001020	**Low discrimination**						**05001020**
					COC	SPO	
Atopobium minutum		63.59	0.74		1	1	
Gemella morbillorum		33.51	0.65		99	1	
Clostridium cadaveris		1.13	0.34		1	99	
05001021	**Low discrimination**						**05001021**
					COC	SPO	
Atopobium minutum		47.95	0.40	URE 1	1	1	
Gemella morbillorum		25.27	0.31		99	1	
Clostridium novyi biovar A		24.43	0.39		1	99	
05001022	**Very good identification**						**05001022**
					COC	GRA	
Atopobium minutum		98.59	0.71		1	99	
Gemella morbillorum		0.79	0.31		99	99	
Pseudoflavonifractor capillosus		0.36	0.15		1	1	
05001023	**Low discrimination**						**05001023**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		88.12	0.37	URE 1	1		
Clostridium novyi biovar A		11.88	0.26		99		
05001030	**Low discrimination**						**05001030**
					COC	SPO	
Atopobium minutum		48.05	0.40	MLZ 1	1	1	
Gemella morbillorum		25.32	0.31		99	1	
Clostridium novyi biovar A		24.48	0.39		1	99	
05001031	**Good identification**						**05001031**
					SPO		
Clostridium novyi biovar A		98.06	0.39	MNS 1	99		
Atopobium minutum		1.94	0.06	GAL 10	1		
05001032	**Low discrimination**						**05001032**
					SPO		
Atopobium minutum		88.12	0.37	MLZ 1	1		
Clostridium novyi biovar A		11.88	0.26		99		
05001040	**Low discrimination**						**05001040**
					COC	SPO	
Gemella morbillorum		45.28	0.47	SOR 1	99	1	
Atopobium minutum		28.64	0.48		1	1	
Clostridium novyi biovar A		12.97	0.47		1	99	
05001041	**Acceptable identification**						**05001041**
					COC	SPO	
Clostridium novyi biovar A		92.54	0.47	MNS 1	1	99	
Gemella morbillorum		3.26	0.13	GAL 10	99	1	
Atopobium minutum		2.07	0.14		1	1	
05001042	**Low discrimination**						**05001042**
					COC	SPO	
Atopobium minutum		87.01	0.45	SOR 1	1	1	
Clostridium novyi biovar A		10.43	0.34		1	99	
Gemella morbillorum		2.08	0.13		99	1	
05001050	**Acceptable identification**						**05001050**
					COC	SPO	
Clostridium novyi biovar A		93.84	0.47	MNS 1	1	99	
Gemella morbillorum		3.31	0.13	GAL 10	99	1	
Atopobium minutum		2.09	0.14		1	1	
05001051	**Good identification**						**05001051**
Clostridium novyi biovar A		100.0	0.47	MNS 1			
05001060	**Low discrimination**						**05001060**
					COC	SPO	
Atopobium minutum		46.86	0.40	SOR 1	1	1	
Gemella morbillorum		24.70	0.31		99	1	
Clostridium novyi biovar A		15.92	0.36		1	99	
05001061	**Good identification**						**05001061**
					SPO		
Clostridium novyi biovar A		97.11	0.36	MNS 1	99		
Atopobium minutum		2.89	0.06	GAL 10	1		
05001062	**Acceptable identification**						**05001062**
					SPO		
Atopobium minutum		91.75	0.37	SOR 1	1		
Clostridium novyi biovar A		8.25	0.23		99		
05001070	**Good identification**						**05001070**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium novyi biovar A		97.11	0.36	MNS 1	99		
Atopobium minutum		2.89	0.06	GAL 10	1		
05001071	**Good identification**						**05001071**
Clostridium novyi biovar A		100.0	0.36	MNS 1			
05001100	**Low discrimination**						**05001100**
					COC	GRA	
Gemella morbillorum		50.54	0.47	LAC 1	99	99	
Atopobium minutum		31.96	0.48		1	99	
Prevotella bivia		7.61	0.19		1	1	
05001102	**Low discrimination**						**05001102**
					GRA		
Atopobium minutum		89.76	0.45	LAC 1	99		
Lactobacillus fermentum		5.70	0.32		99		
Fusobacterium mortiferum		2.39	0.24		1		
05001120	**Good identification**						**05001120**
					COC	GRA	
Prevotella bivia		97.52	0.53	MLT 99	1	1	
Atopobium minutum		1.38	0.40		1	99	
Gemella morbillorum		0.73	0.31		99	99	
05001200	**Low discrimination**						**05001200**
					COC		
Gemella morbillorum		59.54	0.47	RHA 1	99		
Atopobium minutum		37.66	0.48		1		
Peptostreptococcus anaerobius		0.89	0.08		99		
05001202	**Good identification**						**05001202**
					COC		
Atopobium minutum		97.66	0.45	RHA 1	1		
Gemella morbillorum		2.34	0.13		99		
05001220	**Low discrimination**						**05001220**
					COC	SPO	
Atopobium minutum		64.28	0.40	RHA 1	1	1	
Gemella morbillorum		33.88	0.31		99	1	
Clostridium cadaveris		1.14	0.00		1	99	
05001222	**Good identification**						**05001222**
Atopobium minutum		100.0	0.37	RHA 1			
05001400	**Good identification**						**05001400**
					COC		
Gemella morbillorum		94.07	0.65	ARA 10	99		
Atopobium minutum		5.41	0.48		1		
Lactobacillus fermentum		0.14	0.28		1		
05001402	**Low discrimination**						**05001402**
					COC		
Atopobium minutum		76.78	0.45	ARA 1	1		
Gemella morbillorum		20.23	0.31		99		
Lactobacillus fermentum		2.99	0.28		1		
05001420	**Low discrimination**						**05001420**
					COC		
Gemella morbillorum		84.44	0.48	ARA 10	99		
Atopobium minutum		14.57	0.40	NAG 10	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		0.38	0.20		1		
05001422	**Low discrimination**						**05001422**
					COC		
Atopobium minutum		88.75	0.37	ARA 1	1		
Gemella morbillorum		7.80	0.14		99		
Lactobacillus fermentum		3.46	0.20		1		
05002000	**Low discrimination**						**05002000**
					SPO		
Eubacterium limosum		68.54	0.63	FRU 99	1		
Atopobium minutum		5.65	0.56		1		
Clostridium difficile		5.54	0.46		99		
05002002	**Low discrimination**						**05002002**
					COC	GRA	
Atopobium minutum		74.83	0.53	MAN 1	1	99	
Eubacterium limosum		13.76	0.29		1	99	
Megasphaera elsdenii		1.63	0.22		99	1	
05002010	**Very good identification**						**05002010**
					SPO		
Clostridium difficile		96.14	0.58	FRU 99	99		
Eubacterium limosum		2.46	0.29		1		
Clostridium novyi biovar A		0.41	0.32		99		
05002020	**Low discrimination**						**05002020**
					SPO		
Atopobium minutum		36.42	0.48	MAN 1	1		
Clostridium cadaveris		21.33	0.34		99		
Eubacterium limosum		13.39	0.29		1		
05002022	**Acceptable identification**						**05002022**
					CAT		
Atopobium minutum		94.99	0.45	MAN 1	1		
Pseudoramibacter alactolyticus		1.08	-.02		1		
Propionibacterium granulosum		1.05	0.14		90		
05002050	**Acceptable identification**						**05002050**
					SPO		
Clostridium difficile		93.97	0.36	FRU 99	99		
Clostridium novyi biovar A		5.12	0.29	SOR 5	99		
Eubacterium limosum		0.46	-.06		1		
05003000	**Low discrimination**						**05003000**
					COC		
Gemella morbillorum		49.72	0.47	MAN 1	99		
Atopobium minutum		31.45	0.48		1		
Eubacterium limosum		11.56	0.29		1		
05003002	**Acceptable identification**						**05003002**
					COC	CAT	
Atopobium minutum		92.39	0.45	MAN 1	1	1	
Propionibacterium granulosum		4.09	0.24		1	90	
Gemella morbillorum		2.21	0.13		99	1	
05003020	**Low discrimination**						**05003020**
					COC	CAT	
Atopobium minutum		44.34	0.40	MAN 1	1	1	
Propionibacterium acnes		27.09	0.29		1	95	
Gemella morbillorum		23.37	0.31		99	1	
05003022	**Good identification**						**05003022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Atopobium minutum		96.79	0.37	MAN 1	1		
Propionibacterium granulosum		3.21	0.14		90		
05004000	**Low discrimination**						**05004000**
					COC		
Gemella morbillorum		74.54	0.69	XYL 10	99		
Atopobium minutum		6.92	0.56		1		
Peptostreptococcus anaerobius		5.37	0.42		99		
05004002	**Low discrimination**						**05004002**
					COC		
Atopobium minutum		81.08	0.53	XYL 1	1		
Gemella morbillorum		13.23	0.35		99		
Peptostreptococcus anaerobius		0.95	0.08		99		
05004010	**Low discrimination**						**05004010**
					COC	SPO	
Gemella morbillorum		56.67	0.35	XYL 10	99	1	
Clostridium difficile		14.59	0.19	MLZ 1	1	99	
Clostridium novyi biovar A		10.72	0.32		1	99	
05004020	**Low discrimination**						**05004020**
					COC	SPO	
Gemella morbillorum		67.46	0.53	XYL 10	99	1	
Atopobium minutum		18.80	0.48	NAG 10	1	1	
Clostridium cadaveris		11.01	0.34		1	99	
05004022	**Acceptable identification**						**05004022**
					COC	GRA	
Atopobium minutum		93.09	0.45	XYL 1	1	99	
Gemella morbillorum		5.06	0.19		99	99	
Pseudoflavonifractor capillosus		1.02	-.03		1	1	
05004040	**Low discrimination**						**05004040**
					COC	SPO	
Clostridium glycolicum		80.73	0.49	MNS 1	1	99	
Clostridium botulinum biovar A		8.24	0.33	FRU 90	1	99	
Gemella morbillorum		7.43	0.35		99	1	
05004100	**Low discrimination**						**05004100**
					COC		
Gemella morbillorum		66.12	0.35	XYL 10	99		
Actinomyces meyeri		8.46	0.21	LAC 1	1		
Atopobium minutum		6.14	0.22		1		
05004200	**Low discrimination**						**05004200**
					COC		
Gemella morbillorum		75.87	0.35	XYL 10	99		
Atopobium minutum		7.05	0.22	RHA 1	1		
Peptostreptococcus anaerobius		5.47	0.08		99		
05004400	**Good identification**						**05004400**
					COC		
Gemella morbillorum		97.19	0.53	XYL 10	99		
Atopobium minutum		0.82	0.22	ARA 10	1		
Peptostreptococcus anaerobius		0.64	0.08		99		
05004420	**Good identification**						**05004420**
					COC	SPO	
Gemella morbillorum		96.14	0.37	XYL 10	99	1	
Atopobium minutum		2.44	0.14	ARA 10	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium cadaveris		1.43	0.00	NAG 10	1	99	
05005000	**Good identification**						**05005000**
					COC		
Gemella morbillorum		93.86	0.65	XYL 10	99		
Atopobium minutum		5.40	0.48		1		
Peptostreptococcus anaerobius		0.13	0.08		99		
05005002	**Low discrimination**						**05005002**
					COC		
Atopobium minutum		77.74	0.45	XYL 1	1		
Gemella morbillorum		20.49	0.31		99		
Lactobacillus fermentum		1.78	0.24		1		
05005020	**Low discrimination**						**05005020**
					COC	SPO	
Gemella morbillorum		84.74	0.48	XYL 10	99	1	
Atopobium minutum		14.62	0.40	NAG 10	1	1	
Clostridium cadaveris		0.26	0.00		1	99	
05005022	**Acceptable identification**						**05005022**
					COC		
Atopobium minutum		90.03	0.37	XYL 1	1		
Gemella morbillorum		7.91	0.14		99		
Lactobacillus fermentum		2.06	0.16		1		
05005400	**Good identification**						**05005400**
					COC		
Gemella morbillorum		98.51	0.48	XYL 10	99		
Lactobacillus fermentum		0.78	0.24	ARA 10	1		
Atopobium minutum		0.52	0.14		1		
05006010	**Good identification**						**05006010**
					COC	SPO	
Clostridium difficile		99.08	0.36	FRU 99	1	99	
Eubacterium limosum		0.49	-.06	XYL 5	1	1	
Gemella morbillorum		0.43	0.01		99	1	
05010000	**Low discrimination**						**05010000**
					COC	GRA	
Fusobacterium varium		61.82	0.93		1	1	
Eubacterium limosum		14.31	0.84		1	99	
Gemella morbillorum		8.67	0.87		99	99	
05010001	**Low discrimination**						**05010001**
					COC	GRA	
Fusobacterium varium		59.49	0.58	URE 1	1	1	
Eubacterium limosum		13.77	0.49		1	99	
Gemella morbillorum		8.34	0.53		99	99	
05010002	**Low discrimination**						**05010002**
					GRA		
Atopobium minutum		78.14	0.87		99		
Fusobacterium varium		10.95	0.58		1		
Pseudoramibacter alactolyticus		3.37	0.50		99		
05010003	**Low discrimination**						**05010003**
					COC	GRA	
Atopobium minutum		69.83	0.53	URE 1	1	99	
Fusobacterium varium		9.78	0.24		1	1	
Anaerococcus prevotii		6.34	0.21		99	99	
05010010	**Low discrimination**						**05010010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium difficile		78.32	0.76	MAN 90	99	99	
Fusobacterium varium		12.69	0.58		1	1	
Eubacterium limosum		2.94	0.49		99	1	
05010011	**Low discrimination**						**05010011**
					GRA	SPO	
Clostridium difficile		58.81	0.41	MAN 90	99	99	
Clostridium novyi biovar A		25.03	0.50	URE 1	99	99	
Fusobacterium varium		9.53	0.24		1	1	
05010012	**Low discrimination**						**05010012**
					SPO	CAT	
Atopobium minutum		40.50	0.53	MLZ 1	1	1	
Clostridium difficile		35.01	0.41		99	1	
Propionibacterium granulosum		11.59	0.46		1	90	
05010013	**Low discrimination**						**05010013**
					SPO		
Clostridium novyi biovar A		74.93	0.37	MNS 1	99		
Atopobium minutum		11.66	0.19	FRU 10	1		
Clostridium difficile		10.08	0.07		99		
05010020	**Low discrimination**						**05010020**
					COC	SPO	
Atopobium minutum		33.14	0.82		1	1	
Clostridium cadaveris		21.02	0.68		1	99	
Gemella morbillorum		14.32	0.71		99	1	
05010021	**Low discrimination**						**05010021**
					COC	SPO	
Atopobium minutum		31.25	0.48	URE 1	1	1	
Clostridium cadaveris		19.83	0.34		1	99	
Gemella morbillorum		13.50	0.37		99	1	
05010022	**Good identification**						**05010022**
					CAT		
Atopobium minutum		93.58	0.79		1		
Pseudoramibacter alactolyticus		2.14	0.37		1		
Propionibacterium granulosum		1.49	0.50		90		
05010023	**Acceptable identification**						**05010023**
					SPO		
Atopobium minutum		90.13	0.45	URE 1	1		
Clostridium novyi biovar A		4.39	0.26		99		
Pseudoramibacter alactolyticus		2.06	0.03		1		
05010030	**Low discrimination**						**05010030**
					SPO		
Clostridium difficile		31.71	0.41	MAN 90	99		
Atopobium minutum		18.34	0.48	NAG 1	1		
Clostridium cadaveris		11.64	0.34		99		
05010031	**Low discrimination**						**05010031**
					SPO		
Clostridium novyi biovar A		79.28	0.39	MNS 1	99		
Clostridium difficile		7.53	0.07	FRU 10	99		
Atopobium minutum		4.35	0.14		1		
05010032	**Low discrimination**						**05010032**
					SPO	CAT	
Atopobium minutum		74.94	0.45	MLZ 1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		16.08	0.35		1	90	
Clostridium novyi biovar A		3.65	0.26		99	1	
05010040	**Low discrimination**						**05010040**
					GRA		
Fusobacterium varium		36.15	0.58	SOR 1	1		
Pseudoramibacter alactolyticus		25.02	0.56		99		
Eubacterium limosum		8.37	0.49		99		
05010041	**Low discrimination**						**05010041**
					GRA	SPO	
Clostridium novyi biovar A		38.49	0.47	MNS 1	99	99	
Fusobacterium varium		21.98	0.24	FRU 10	1	1	
Pseudoramibacter alactolyticus		15.21	0.21		99	1	
05010042	**Low discrimination**						**05010042**
					COC	GRA	
Pseudoramibacter alactolyticus		39.85	0.39	MNS 1	1	99	
Atopobium minutum		37.37	0.53	BGL 10	1	99	
Megasphaera elsdenii		11.19	0.41		99	1	
05010050	**Low discrimination**						**05010050**
Clostridium difficile		81.06	0.54	MAN 90			
Clostridium botulinum biovar B		7.76	0.36	SOR 5			
Clostridium novyi biovar A		4.42	0.47				
05010051	**Low discrimination**						**05010051**
Clostridium novyi biovar A		83.11	0.47	MNS 1			
Clostridium difficile		15.41	0.19	FRU 10			
Clostridium botulinum biovar B		1.48	0.02				
05010060	**Low discrimination**						**05010060**
					CAT		
Propionibacterium acnes		55.00	0.56	IND 87	95		
Pseudoramibacter alactolyticus		21.85	0.43	NIT 92	1		
Atopobium minutum		6.45	0.48		1		
05010061	**Low discrimination**						**05010061**
					SPO	CAT	
Clostridium novyi biovar A		45.00	0.36	MNS 1	99	1	
Propionibacterium acnes		31.60	0.22	FRU 10	1	95	
Pseudoramibacter alactolyticus		12.55	0.09		1	1	
05010062	**Low discrimination**						**05010062**
					CAT		
Atopobium minutum		56.17	0.45	SOR 1	1		
Pseudoramibacter alactolyticus		31.71	0.26		1		
Propionibacterium acnes		7.26	0.22		95		
05010071	**Good identification**						**05010071**
Clostridium novyi biovar A		96.62	0.36	MNS 1			
Clostridium botulinum biovar B		3.38	-.03	FRU 10			
05010100	**Low discrimination**						**05010100**
					GRA		
Fusobacterium varium		48.76	0.58	LAC 1	1		
Fusobacterium mortiferum		20.07	0.71		1		
Eubacterium limosum		11.29	0.49		99		
05010102	**Low discrimination**						**05010102**

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					GRA			
Atopobium minutum		51.60	0.53	LAC 1	99			
Fusobacterium mortiferum		32.73	0.55		1			
Fusobacterium varium		7.23	0.24		1			
05010110	**Low discrimination**							**05010110**
					GRA	SPO		
Clostridium difficile		74.40	0.41	MAN 90	99	99		
Fusobacterium varium		12.06	0.24	LAC 1	1	1		
Fusobacterium mortiferum		4.96	0.37		1	1		
05010120	**Low discrimination**							**05010120**
					GRA	SPO		
Prevotella bivia		83.17	0.50	MLT 99	1	1		
Atopobium minutum		5.17	0.48	FRU 10	99	1		
Clostridium cadaveris		3.28	0.34		99	99		
05010122	**Low discrimination**							**05010122**
					GRA			
Atopobium minutum		73.77	0.45	LAC 1	99			
Prevotella bivia		17.97	0.15		1			
Lactobacillus fermentum		1.69	0.24		99			
05010200	**Low discrimination**							**05010200**
					COC	GRA		
Fusobacterium varium		61.35	0.58	RHA 1	1	1		
Eubacterium limosum		14.20	0.49		1	99		
Gemella morbillorum		8.60	0.53		99	99		
05010202	**Low discrimination**							**05010202**
					GRA	CAT		
Atopobium minutum		75.75	0.53	RHA 1	99	1		
Fusobacterium varium		10.61	0.24		1	1		
Propionibacterium granulosum		4.92	0.35		99	90		
05010210	**Low discrimination**							**05010210**
					GRA	SPO	CAT	
Clostridium difficile		76.78	0.41	MAN 90	99	99	1	
Fusobacterium varium		12.44	0.24	RHA 1	1	1	1	
Propionibacterium granulosum		3.15	0.30		99	1	90	
05010220	**Low discrimination**							**05010220**
					COC	SPO		
Atopobium minutum		32.30	0.48	RHA 1	1	1		
Clostridium cadaveris		20.49	0.34		1	99		
Gemella morbillorum		13.95	0.37		99	1		
05010222	**Acceptable identification**							**05010222**
					CAT			
Atopobium minutum		91.44	0.45	RHA 1	1			
Propionibacterium granulosum		4.45	0.24		90			
Pseudoramibacter alactolyticus		2.09	0.03		1			
05010400	**Low discrimination**							**05010400**
					COC	GRA		
Gemella morbillorum		51.04	0.71	ARA 10	99	99		
Fusobacterium varium		33.10	0.58		1	1		
Eubacterium limosum		7.66	0.49		1	99		
05010401	**Low discrimination**							**05010401**
					COC	GRA		
Gemella morbillorum		50.50	0.37	ARA 10	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium varium		32.75	0.24	URE 1	1	1		
Eubacterium limosum		7.58	0.15		1	99		
05010402	**Low discrimination**							**05010402**
					COC	GRA		
Atopobium minutum		67.25	0.53	ARA 1	1	99		
Gemella morbillorum		14.53	0.37		99	99		
Fusobacterium varium		9.42	0.24		1	1		
05010410	**Low discrimination**							**05010410**
					COC	GRA	SPO	
Clostridium difficile		66.58	0.41	MAN 90	1	99	99	
Gemella morbillorum		16.64	0.37	ARA 1	99	99	1	
Fusobacterium varium		10.79	0.24		1	1	1	
05010420	**Low discrimination**							**05010420**
					COC	SPO		
Gemella morbillorum		64.73	0.54	ARA 10	99	1		
Atopobium minutum		13.62	0.48	NAG 10	1	1		
Clostridium cadaveris		8.64	0.34		1	99		
05010422	**Low discrimination**							**05010422**
					COC			
Atopobium minutum		87.69	0.45	ARA 1	1			
Gemella morbillorum		6.31	0.20		99			
Pseudoramibacter alactolyticus		2.00	0.03		1			
05010440	**Low discrimination**							**05010440**
					COC	GRA		
Gemella morbillorum		37.17	0.37	ARA 10	99	99		
Fusobacterium varium		24.10	0.24	SOR 1	1	1		
Pseudoramibacter alactolyticus		16.68	0.21		1	99		
05010500	**Low discrimination**							**05010500**
					COC	GRA		
Gemella morbillorum		38.83	0.37	LAC 1	99	99		
Fusobacterium varium		25.18	0.24	ARA 10	1	1		
Fusobacterium mortiferum		10.37	0.37		1	1		
05010600	**Low discrimination**							**05010600**
					COC	GRA		
Gemella morbillorum		51.07	0.37	RHA 1	99	99		
Fusobacterium varium		33.12	0.24	ARA 10	1	1		
Eubacterium limosum		7.67	0.15		1	99		
05011000	**Low discrimination**							**05011000**
					COC	GRA		
Gemella morbillorum		55.24	0.83		99	99		
Atopobium minutum		26.39	0.82		1	99		
Fusobacterium varium		7.39	0.58		1	1		
05011001	**Low discrimination**							**05011001**
					COC	GRA		
Gemella morbillorum		53.34	0.48	URE 1	99	99		
Atopobium minutum		25.48	0.48		1	99		
Fusobacterium varium		7.14	0.24		1	1		
05011002	**Low discrimination**							**05011002**
					CAT			
Atopobium minutum		86.36	0.79		1			
Propionibacterium granulosum		5.49	0.61		90			
Lactobacillus fermentum		3.64	0.62		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05011003	**Low discrimination**							**05011003**
					CAT			
Atopobium minutum		85.36	0.45	URE 1	1			
Propionibacterium granulosum		5.43	0.26		90			
Lactobacillus fermentum		3.60	0.28		1			
05011010	**Low discrimination**							**05011010**
					COC	SPO	CAT	
Propionibacterium granulosum		29.68	0.56	SUC 93	1	1	90	
Gemella morbillorum		27.14	0.48	MLZ 12	99	1	1	
Clostridium difficile		22.41	0.41		1	99	1	
05011012	**Low discrimination**							**05011012**
					CAT			
Atopobium minutum		50.19	0.45	MLZ 1	1			
Propionibacterium granulosum		43.08	0.46		90			
Lactobacillus fermentum		2.12	0.28		1			
05011020	**Low discrimination**							**05011020**
					COC	CAT		
Propionibacterium acnes		38.25	0.68	IND 87	1	95		
Atopobium minutum		32.50	0.74	NIT 92	1	1		
Gemella morbillorum		22.68	0.66		99	1		
05011022	**Low discrimination**							**05011022**
					CAT			
Atopobium minutum		88.98	0.71		1			
Propionibacterium granulosum		4.24	0.50		90			
Lactobacillus fermentum		3.75	0.54		1			
05011023	**Low discrimination**							**05011023**
					CAT			
Atopobium minutum		88.97	0.37	URE 1	1			
Propionibacterium granulosum		4.24	0.16		90			
Lactobacillus fermentum		3.75	0.20		1			
05011030	**Low discrimination**							**05011030**
					CAT			
Propionibacterium granulosum		36.13	0.46	SUC 93	90			
Propionibacterium acnes		24.76	0.34	MLZ 12	95			
Atopobium minutum		21.04	0.40		1			
05011032	**Low discrimination**							**05011032**
					CAT			
Atopobium minutum		58.14	0.37	MLZ 1	1			
Propionibacterium granulosum		37.43	0.35		90			
Lactobacillus fermentum		2.45	0.20		1			
05011040	**Low discrimination**							**05011040**
					COC	GRA		
Gemella morbillorum		47.35	0.48	SOR 1	99	99		
Atopobium minutum		22.62	0.48		1	99		
Fusobacterium varium		6.34	0.24		1	1		
05011042	**Low discrimination**							**05011042**
					CAT			
Atopobium minutum		82.94	0.45	SOR 1	1			
Propionibacterium granulosum		5.27	0.26		90			
Lactobacillus fermentum		3.50	0.28		1			
05011060	**Good identification**							**05011060**
					COC	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		92.28	0.56	IND 87	1	95	
Atopobium minutum		3.61	0.40	NIT 92	1	1	
Gemella morbillorum		2.52	0.32		99	1	
05011062	**Low discrimination**						**05011062**
					CAT		
Atopobium minutum		63.98	0.37	SOR 1	1		
Propionibacterium acnes		24.79	0.22		95		
Clostridium perfringens		4.39	0.11		1		
05011100	**Low discrimination**						**05011100**
					COC	GRA	
Fusobacterium mortiferum		65.56	0.74		1	1	
Lactobacillus fermentum		17.39	0.66		1	99	
Gemella morbillorum		8.01	0.48		99	99	
05011101	**Low discrimination**						**05011101**
					GRA	SPO	
Fusobacterium mortiferum		46.38	0.40	URE 1	1	1	
Clostridium chauvoei		31.15	0.26		99	99	
Lactobacillus fermentum		12.30	0.32		99	1	
05011102	**Low discrimination**						**05011102**
					GRA		
Lactobacillus fermentum		62.97	0.66	SUC 87	99		
Fusobacterium mortiferum		26.38	0.58		1		
Atopobium minutum		9.24	0.45		99		
05011110	**Low discrimination**						**05011110**
					GRA	CAT	
Fusobacterium mortiferum		58.16	0.40	MLZ 1	1	5	
Lactobacillus fermentum		15.43	0.32		99	1	
Propionibacterium granulosum		7.78	0.22		99	90	
05011120	**Low discrimination**						**05011120**
					GRA	SPO	
Clostridium septicum		37.27	0.46	MLT 94	99	99	
Prevotella bivia		25.63	0.37		1	1	
Lactobacillus fermentum		13.68	0.58		99	1	
05011122	**Low discrimination**						**05011122**
Lactobacillus fermentum		60.80	0.58	SUC 87			
Clostridium perfringens		23.57	0.38				
Atopobium minutum		8.92	0.37				
05011140	**Low discrimination**						**05011140**
					COC	GRA	
Fusobacterium mortiferum		65.48	0.40	SOR 1	1	1	
Lactobacillus fermentum		17.37	0.32		1	99	
Gemella morbillorum		8.00	0.14		99	99	
05011200	**Low discrimination**						**05011200**
					COC	CAT	
Gemella morbillorum		50.64	0.48	RHA 1	99	1	
Atopobium minutum		24.19	0.48		1	1	
Propionibacterium granulosum		12.56	0.45		1	90	
05011202	**Low discrimination**						**05011202**
					CAT		
Atopobium minutum		78.07	0.45	RHA 1	1		
Propionibacterium granulosum		15.20	0.35		90		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		3.29	0.28		1		
05011222	**Low discrimination**						**05011222**
					CAT		
Atopobium minutum		82.92	0.37	RHA 1	1		
Propionibacterium granulosum		12.11	0.24		90		
Lactobacillus fermentum		3.50	0.20		1		
05011300	**Low discrimination**						**05011300**
					COC	GRA	
Fusobacterium mortiferum		66.01	0.40	RHA 1	1	1	
Lactobacillus fermentum		17.51	0.32		1	99	
Gemella morbillorum		8.07	0.14		99	99	
05011400	**Low discrimination**						**05011400**
					COC		
Gemella morbillorum		83.69	0.66	ARA 10	99		
Lactobacillus fermentum		10.12	0.62		1		
Atopobium minutum		3.63	0.48		1		
05011402	**Low discrimination**						**05011402**
					COC		
Lactobacillus fermentum		74.56	0.62	SUC 87	1		
Atopobium minutum		17.85	0.45		1		
Gemella morbillorum		6.23	0.32		99		
05011420	**Low discrimination**						**05011420**
					COC	CAT	
Gemella morbillorum		59.84	0.50	ARA 10	99	1	
Lactobacillus fermentum		21.71	0.54	NAG 10	1	1	
Propionibacterium acnes		9.17	0.34		1	95	
05011422	**Low discrimination**						**05011422**
					COC		
Lactobacillus fermentum		77.95	0.54	SUC 87	1		
Atopobium minutum		18.66	0.37		1		
Gemella morbillorum		2.17	0.16		99		
05011500	**Low discrimination**						**05011500**
					COC	GRA	
Lactobacillus fermentum		88.49	0.66	SUC 87	1	99	
Gemella morbillorum		4.53	0.32		99	99	
Fusobacterium mortiferum		3.37	0.40		1	1	
05011502	**Very good identification**						**05011502**
					GRA		
Lactobacillus fermentum		99.32	0.66	SUC 87	99		
Fusobacterium mortiferum		0.42	0.24		1		
Atopobium minutum		0.15	0.11		99		
05011520	**Low discrimination**						**05011520**
					GRA	SPO	
Lactobacillus fermentum		59.24	0.58	SUC 87	99	1	
Odoribacter splanchnicus		35.33	0.32		1	1	
Clostridium septicum		1.63	0.12		99	99	
05011522	**Very good identification**						**05011522**
					GRA		
Lactobacillus fermentum		99.02	0.58	SUC 87	99		
Odoribacter splanchnicus		0.60	-.03		1		
Clostridium perfringens		0.39	0.04		99		
05012000	**Good identification**						**05012000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Eubacterium limosum		90.40	0.97		1		
Clostridium difficile		7.30	0.80		99		
Pseudoramibacter alactolyticus		0.96	0.61		1		
05012001	**Good identification**						**05012001**
					SPO		
Eubacterium limosum		90.38	0.63	URE 1	1		
Clostridium difficile		7.30	0.46		99		
Pseudoramibacter alactolyticus		0.96	0.27		1		
05012002	**Low discrimination**						**05012002**
					SPO		
Eubacterium limosum		73.53	0.63	BGL 1	1		
Pseudoramibacter alactolyticus		8.56	0.45		1		
Clostridium difficile		5.94	0.46		99		
05012010	**Very good identification**						**05012010**
					SPO	CAT	
Clostridium difficile		97.34	0.92		99	1	
Eubacterium limosum		2.49	0.63		1	1	
Propionibacterium granulosum		0.11	0.54		1	90	
05012011	**Very good identification**						**05012011**
					SPO	CAT	
Clostridium difficile		97.33	0.58	URE 1	99	1	
Eubacterium limosum		2.49	0.29		1	1	
Propionibacterium granulosum		0.11	0.20		1	90	
05012012	**Good identification**						**05012012**
					SPO	CAT	
Clostridium difficile		94.50	0.58	BGL 1	99	1	
Propionibacterium granulosum		2.62	0.44		1	90	
Eubacterium limosum		2.42	0.29		1	1	
05012020	**Low discrimination**						**05012020**
					CAT		
Eubacterium limosum		50.97	0.63	NAG 1	1		
Propionibacterium acnes		28.23	0.63		95		
Pseudoramibacter alactolyticus		9.43	0.48		1		
05012030	**Good identification**						**05012030**
					SPO	CAT	
Clostridium difficile		93.14	0.58	NAG 1	99	1	
Propionibacterium granulosum		2.59	0.44		1	90	
Eubacterium limosum		2.39	0.29		1	1	
05012040	**Low discrimination**						**05012040**
					SPO		
Eubacterium limosum		54.49	0.63	SOR 1	1		
Clostridium difficile		22.94	0.58		99		
Pseudoramibacter alactolyticus		14.28	0.51		1		
05012050	**Very good identification**						**05012050**
					SPO		
Clostridium difficile		99.28	0.70	SOR 5	99		
Eubacterium limosum		0.49	0.29		1		
Pseudoramibacter alactolyticus		0.13	0.17		1		
05012051	**Good identification**						**05012051**
					SPO		
Clostridium difficile		98.92	0.36	SOR 5	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		0.60	0.12	URE 1	99			
Eubacterium limosum		0.49	-.06		1			
05012052	**Good identification**							**05012052**
					COC	SPO	GRA	
Clostridium difficile		96.92	0.36	SOR 5	1	99	99	
Pseudoramibacter alactolyticus		1.37	0.00	BGL 1	1	1	99	
Megasphaera elsdenii		0.72	0.06		99	1	1	
05012060	**Low discrimination**							**05012060**
					CAT			
Propionibacterium acnes		65.68	0.52	IND 87	95			
Pseudoramibacter alactolyticus		24.98	0.38	NIT 92	1			
Eubacterium limosum		5.46	0.29		1			
05012070	**Acceptable identification**							**05012070**
					SPO	CAT		
Clostridium difficile		91.15	0.36	NAG 1	99	1		
Propionibacterium acnes		5.39	0.18	SOR 5	1	95		
Pseudoramibacter alactolyticus		2.05	0.04		1	1		
05012100	**Good identification**							**05012100**
					SPO			
Eubacterium limosum		90.15	0.63	LAC 1	1			
Clostridium difficile		7.28	0.46		99			
Pseudoramibacter alactolyticus		0.96	0.27		1			
05012110	**Very good identification**							**05012110**
					SPO	CAT		
Clostridium difficile		97.37	0.58	LAC 1	99	1		
Eubacterium limosum		2.49	0.29		1	1		
Propionibacterium granulosum		0.11	0.20		1	90		
05012150	**Good identification**							**05012150**
					SPO			
Clostridium difficile		99.51	0.36	LAC 1	99			
Eubacterium limosum		0.49	-.06	SOR 5	1			
05012200	**Low discrimination**							**05012200**
					SPO			
Eubacterium limosum		89.86	0.63	RHA 1	1			
Clostridium difficile		7.26	0.46		99			
Pseudoramibacter alactolyticus		0.95	0.27		1			
05012210	**Very good identification**							**05012210**
					SPO	CAT		
Clostridium difficile		97.15	0.58	RHA 1	99	1		
Eubacterium limosum		2.49	0.29		1	1		
Propionibacterium granulosum		0.33	0.28		1	90		
05012250	**Good identification**							**05012250**
					SPO			
Clostridium difficile		99.51	0.36	RHA 1	99			
Eubacterium limosum		0.49	-.06	SOR 5	1			
05012400	**Low discrimination**							**05012400**
					COC	SPO		
Eubacterium limosum		89.60	0.63	ARA 1	1	1		
Clostridium difficile		7.24	0.46		1	99		
Gemella morbillorum		0.98	0.37		99	1		
05012410	**Very good identification**							**05012410**
					SPO	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		97.35	0.58	ARA 1	99	1	
Eubacterium limosum		2.49	0.29		1	1	
Propionibacterium granulosum		0.11	0.20		1	90	
05012450	**Good identification**						**05012450**
					SPO		
Clostridium difficile		99.51	0.36	ARA 1	99		
Eubacterium limosum		0.49	-.06	SOR 5	1		
05013000	**Low discrimination**						**05013000**
					SPO	CAT	
Eubacterium limosum		66.10	0.63	GAL 1	1	1	
Propionibacterium granulosum		21.23	0.69		1	90	
Clostridium difficile		5.34	0.46		99	1	
05013002	**Low discrimination**						**05013002**
					CAT		
Propionibacterium granulosum		70.46	0.58	SUC 93	90		
Atopobium minutum		14.84	0.45		1		
Eubacterium limosum		8.87	0.29		1		
05013010	**Low discrimination**						**05013010**
					SPO	CAT	
Clostridium difficile		87.78	0.58	GAL 1	99	1	
Propionibacterium granulosum		9.74	0.54		1	90	
Eubacterium limosum		2.25	0.29		1	1	
05013012	**Low discrimination**						**05013012**
					SPO	CAT	
Propionibacterium granulosum		71.42	0.44	SUC 93	1	90	
Clostridium difficile		25.99	0.24	MLZ 12	99	1	
Atopobium minutum		1.11	0.11		1	1	
05013020	**Low discrimination**						**05013020**
					CAT		
Propionibacterium acnes		83.71	0.63	IND 87	95		
Propionibacterium granulosum		12.13	0.58	NIT 92	90		
Eubacterium limosum		1.53	0.29		1		
05013022	**Low discrimination**						**05013022**
					CAT		
Propionibacterium granulosum		63.02	0.48	SUC 93	90		
Atopobium minutum		17.69	0.37		1		
Propionibacterium acnes		17.57	0.29		95		
05013030	**Low discrimination**						**05013030**
					SPO	CAT	
Propionibacterium granulosum		52.85	0.44	SUC 93	1	90	
Propionibacterium acnes		27.01	0.29	MLZ 12	1	95	
Clostridium difficile		19.24	0.24		99	1	
05013050	**Good identification**						**05013050**
					SPO	CAT	
Clostridium difficile		97.45	0.36	GAL 1	99	1	
Propionibacterium granulosum		2.08	0.20	SOR 5	1	90	
Eubacterium limosum		0.48	-.06		1	1	
05013060	**Good identification**						**05013060**
					CAT		
Propionibacterium acnes		98.81	0.52	IND 87	95		
Propionibacterium granulosum		0.66	0.24	NIT 92	90		
Pseudoramibacter alactolyticus		0.38	0.04		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05014000	**Low discrimination**							**05014000**
					COC	GRA		
Fusobacterium varium		65.65	0.69	XYL 4	1	1		
Gemella morbillorum		24.54	0.71		99	99		
Eubacterium limosum		3.68	0.49		1	99		
05014002	**Low discrimination**							**05014002**
					COC	GRA		
Atopobium minutum		51.11	0.53	XYL 1	1	99		
Fusobacterium varium		29.53	0.35		1	1		
Gemella morbillorum		11.04	0.37		99	99		
05014010	**Low discrimination**							**05014010**
					COC	GRA	SPO	
Clostridium difficile		83.61	0.54	MAN 90	1	99	99	
Fusobacterium varium		10.73	0.35	XYL 5	1	1	1	
Gemella morbillorum		4.01	0.37		99	99	1	
05014020	**Low discrimination**							**05014020**
					COC	GRA		
Gemella morbillorum		57.42	0.54	XYL 10	99	99		
Fusobacterium varium		13.96	0.35	NAG 10	1	1		
Atopobium minutum		12.08	0.48		1	99		
05014022	**Low discrimination**							**05014022**
					COC			
Atopobium minutum		86.80	0.45	XYL 1	1			
Gemella morbillorum		6.25	0.20		99			
Pseudoramibacter alactolyticus		1.98	0.03		1			
05014040	**Good identification**							**05014040**
					COC	GRA	SPO	
Clostridium glycolicum		94.27	0.65	MNS 1	1	99	99	
Fusobacterium varium		3.15	0.35		1	1	1	
Gemella morbillorum		1.18	0.37		99	99	1	
05014102	**Low discrimination**							**05014102**
					GRA			
Fusobacterium mortiferum		51.54	0.36	XYL 7	1			
Lactobacillus fermentum		14.55	0.28	BGL 10	99			
Actinomyces meyeri		14.35	0.17		99			
05014400	**Low discrimination**							**05014400**
					COC	GRA		
Gemella morbillorum		77.35	0.54	XYL 10	99	99		
Fusobacterium varium		18.81	0.35	ARA 10	1	1		
Eubacterium limosum		1.06	0.15		1	99		
05014420	**Acceptable identification**							**05014420**
					COC	GRA		
Gemella morbillorum		92.15	0.38	XYL 10	99	99		
Fusobacterium varium		2.04	0.01	ARA 10	1	1		
Atopobium minutum		1.76	0.14	NAG 10	1	99		
05015000	**Low discrimination**							**05015000**
					COC	GRA		
Gemella morbillorum		81.60	0.66	XYL 10	99	99		
Lactobacillus fermentum		5.80	0.58		1	99		
Fusobacterium varium		4.10	0.35		1	1		
05015002	**Low discrimination**							**05015002**
					COC			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		61.48	0.58	SUC 87	1		
Atopobium minutum		25.06	0.45		1		
Gemella morbillorum		8.75	0.32		99		
05015020	**Low discrimination**						**05015020**
					COC	CAT	
Gemella morbillorum		64.95	0.50	XYL 10	99	1	
Lactobacillus fermentum		13.84	0.50	NAG 10	1	1	
Propionibacterium acnes		9.96	0.34		1	95	
05015022	**Low discrimination**						**05015022**
					COC		
Lactobacillus fermentum		67.49	0.50	SUC 87	1		
Atopobium minutum		27.51	0.37		1		
Gemella morbillorum		3.20	0.16		99		
05015040	**Acceptable identification**						**05015040**
					COC	SPO	
Clostridium glycolicum		93.28	0.49	MNS 1	1	99	
Gemella morbillorum		5.65	0.32	GAL 10	99	1	
Lactobacillus fermentum		0.40	0.24		1	1	
05015100	**Low discrimination**						**05015100**
					GRA		
Lactobacillus fermentum		48.51	0.62	SUC 87	99		
Fusobacterium mortiferum		23.44	0.55		1		
Actinomyces meyeri		23.43	0.46		99		
05015102	**Good identification**						**05015102**
					GRA		
Lactobacillus fermentum		93.36	0.62	SUC 87	99		
Fusobacterium mortiferum		5.01	0.39		1		
Actinomyces meyeri		1.40	0.20		99		
05015120	**Low discrimination**						**05015120**
					SPO		
Lactobacillus fermentum		74.47	0.54	SUC 87	1		
Actinomyces meyeri		14.72	0.31		1		
Clostridium septicum		3.49	0.12		99		
05015122	**Good identification**						**05015122**
					GRA		
Lactobacillus fermentum		97.27	0.54	SUC 87	99		
Parabacteroides distasonis		1.49	0.20		1		
Clostridium perfringens		0.65	0.04		99		
05015400	**Low discrimination**						**05015400**
					COC		
Gemella morbillorum		59.48	0.50	XYL 10	99		
Lactobacillus fermentum		38.01	0.58	ARA 10	1		
Actinomyces meyeri		1.87	0.25		1		
05015402	**Very good identification**						**05015402**
					COC		
Lactobacillus fermentum		98.05	0.58	SUC 87	1		
Gemella morbillorum		1.55	0.16		99		
Atopobium minutum		0.40	0.11		1		
05015420	**Low discrimination**						**05015420**
					COC		
Lactobacillus fermentum		64.30	0.50	SUC 87	1		
Gemella morbillorum		33.54	0.34		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		1.30	0.10		1		
05015422	**Good identification**						**05015422**
Lactobacillus fermentum		100.0	0.50	SUC 87			
05015500	**Low discrimination**						**05015500**
					COC		
Lactobacillus fermentum		88.08	0.62	SUC 87	1		
Actinomyces meyeri		10.64	0.35		1		
Gemella morbillorum		0.85	0.16		99		
05015502	**Very good identification**						**05015502**
Lactobacillus fermentum		99.63	0.62	SUC 87			
Actinomyces meyeri		0.37	0.10				
05015520	**Acceptable identification**						**05015520**
					GRA		
Lactobacillus fermentum		93.89	0.54	SUC 87	99		
Actinomyces meyeri		4.64	0.21		99		
Odoribacter splanchnicus		0.96	-.03		1		
05015522	**Good identification**						**05015522**
					GRA		
Lactobacillus fermentum		99.07	0.54	SUC 87	99		
Parabacteroides distasonis		0.93	0.16		1		
05016000	**Low discrimination**						**05016000**
					GRA	SPO	
Eubacterium limosum		67.61	0.63	XYL 1	99	1	
Clostridium difficile		28.46	0.58		99	99	
Fusobacterium varium		1.98	0.35		1	1	
05016010	**Very good identification**						**05016010**
					SPO	CAT	
Clostridium difficile		99.47	0.70	XYL 5	99	1	
Eubacterium limosum		0.49	0.29		1	1	
Propionibacterium granulosum		0.02	0.20		1	90	
05016011	**Good identification**						**05016011**
					SPO		
Clostridium difficile		99.51	0.36	XYL 5	99		
Eubacterium limosum		0.49	-.06	URE 1	1		
05016012	**Good identification**						**05016012**
					SPO	CAT	
Clostridium difficile		98.99	0.36	XYL 5	99	1	
Propionibacterium granulosum		0.53	0.09	BGL 1	1	90	
Eubacterium limosum		0.49	-.06		1	1	
05016030	**Good identification**						**05016030**
					SPO	CAT	
Clostridium difficile		98.99	0.36	XYL 5	99	1	
Propionibacterium granulosum		0.53	0.09	NAG 1	1	90	
Eubacterium limosum		0.49	-.06		1	1	
05016040	**Low discrimination**						**05016040**
					SPO		
Clostridium difficile		48.65	0.36	XYL 5	99		
Eubacterium limosum		22.18	0.29	SOR 5	1		
Clostridium glycolicum		19.43	0.31		99		
05016050	**Good identification**						**05016050**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium difficile		99.82	0.48	XYL 5	99		
Eubacterium limosum		0.09	-.06	SOR 5	1		
Clostridium glycolicum		0.08	-.03		99		
05016110	**Good identification**						**05016110**
					SPO		
Clostridium difficile		99.51	0.36	XYL 5	99		
Eubacterium limosum		0.49	-.06	LAC 1	1		
05016210	**Good identification**						**05016210**
					SPO		
Clostridium difficile		99.51	0.36	XYL 5	99		
Eubacterium limosum		0.49	-.06	RHA 1	1		
05016410	**Good identification**						**05016410**
					SPO		
Clostridium difficile		99.51	0.36	XYL 5	99		
Eubacterium limosum		0.49	-.06	ARA 1	1		
05017010	**Good identification**						**05017010**
					SPO	CAT	
Clostridium difficile		97.45	0.36	GAL 1	99	1	
Propionibacterium granulosum		2.08	0.20	XYL 5	1	90	
Eubacterium limosum		0.48	-.06		1	1	
05020000	**Low discrimination**						**05020000**
					COC	SPO	
Clostridium botulinum biovar A		68.90	0.66	MNS 7	1	99	
Atopobium minutum		6.90	0.56		1	1	
Gemella morbillorum		6.75	0.52		99	1	
05020002	**Low discrimination**						**05020002**
					SPO		
Atopobium minutum		67.00	0.53	TRE 1	1		
Clostridium sporogenes		14.44	0.32		99		
Clostridium botulinum biovar A		10.14	0.32		99		
05020020	**Low discrimination**						**05020020**
					COC	SPO	
Clostridium cadaveris		61.77	0.46	IND 98	1	99	
Atopobium minutum		20.24	0.48	TRE 5	1	1	
Gemella morbillorum		6.61	0.35		99	1	
05020022	**Low discrimination**						**05020022**
					SPO	CAT	
Atopobium minutum		89.88	0.45	TRE 1	1	1	
Clostridium cadaveris		4.16	0.12		99	1	
Propionibacterium granulosum		3.96	0.24		1	90	
05020040	**Low discrimination**						**05020040**
Clostridium botulinum biovar A		86.46	0.57	MNS 7			
Clostridium botulinum biovar B		11.70	0.39				
Clostridium novyi biovar A		0.40	0.29				
05020400	**Low discrimination**						**05020400**
					COC	SPO	
Gemella morbillorum		44.84	0.35	TRE 1	99	1	
Clostridium botulinum biovar A		41.61	0.32	ARA 10	1	99	
Atopobium minutum		4.17	0.22		1	1	
05021000	**Low discrimination**						**05021000**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	CAT		
Gemella morbillorum		45.90	0.47	TRE 1	99	1		
Atopobium minutum		29.03	0.48		1	1		
Propionibacterium granulosum		13.63	0.45		1	90		
05021002	**Low discrimination**							**05021002**
					COC	CAT		
Atopobium minutum		82.88	0.45	TRE 1	1	1		
Propionibacterium granulosum		14.60	0.35		1	90		
Gemella morbillorum		1.99	0.13		99	1		
05021020	**Low discrimination**							**05021020**
					COC	CAT		
Atopobium minutum		50.44	0.40	TRE 1	1	1		
Gemella morbillorum		26.58	0.31		99	1		
Propionibacterium granulosum		17.76	0.35		1	90		
05021022	**Low discrimination**							**05021022**
					CAT			
Atopobium minutum		88.33	0.37	TRE 1	1			
Propionibacterium granulosum		11.67	0.24		90			
05022010	**Good identification**							**05022010**
					SPO	CAT		
Clostridium difficile		95.41	0.38	FRU 99	99	1		
Propionibacterium granulosum		4.26	0.28	TRE 7	1	90		
Eubacterium limosum		0.33	-.06		1	1		
05023000	**Acceptable identification**							**05023000**
					COC	CAT		
Propionibacterium granulosum		92.24	0.43	SUC 93	1	90		
Gemella morbillorum		4.16	0.13	FRU 99	99	1		
Atopobium minutum		2.63	0.14		1	1		
05024000	**Low discrimination**							**05024000**
					COC	SPO		
Gemella morbillorum		44.18	0.35	TRE 1	99	1		
Clostridium botulinum biovar A		41.01	0.32	XYL 10	1	99		
Atopobium minutum		4.10	0.22		1	1		
05030000	**Low discrimination**							**05030000**
					GRA	CAT		
Propionibacterium granulosum		49.84	0.79	SUC 93	99	90		
Fusobacterium varium		27.44	0.58		1	1		
Eubacterium limosum		6.35	0.49		99	1		
05030001	**Low discrimination**							**05030001**
					GRA	CAT		
Propionibacterium granulosum		49.38	0.45	SUC 93	99	90		
Fusobacterium varium		27.19	0.24	URE 1	1	1		
Eubacterium limosum		6.29	0.15		99	1		
05030002	**Low discrimination**							**05030002**
					GRA	CAT		
Propionibacterium granulosum		81.84	0.69	SUC 93	99	90		
Atopobium minutum		13.00	0.53		99	1		
Fusobacterium varium		1.82	0.24		1	1		
05030010	**Low discrimination**							**05030010**
					GRA	SPO	CAT	
Clostridium difficile		63.57	0.56	TRE 7	99	99	1	
Propionibacterium granulosum		33.90	0.64	MAN 90	99	1	90	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Fusobacterium varium	1.38	0.24		1	1	1	
05030012	**Acceptable identification**							**05030012**
					SPO	CAT		
	Propionibacterium granulosum	91.95	0.54	SUC 93	1	90		
	Clostridium difficile	6.97	0.22	MLZ 12	99	1		
	Atopobium minutum	1.08	0.19		1	1		
05030020	**Low discrimination**							**05030020**
					SPO	CAT		
	Propionibacterium granulosum	68.75	0.69	SUC 93	1	90		
	Clostridium cadaveris	18.05	0.46		99	1		
	Atopobium minutum	5.46	0.48		1	1		
05030022	**Low discrimination**							**05030022**
					SPO	CAT		
	Propionibacterium granulosum	81.31	0.58	SUC 93	1	90		
	Atopobium minutum	17.22	0.45		1	1		
	Clostridium cadaveris	0.86	0.12		99	1		
05030030	**Low discrimination**							**05030030**
					SPO	CAT		
	Propionibacterium granulosum	89.81	0.54	SUC 93	1	90		
	Clostridium difficile	6.81	0.22	MLZ 12	99	1		
	Clostridium cadaveris	1.75	0.12		99	1		
05030032	**Good identification**							**05030032**
					CAT			
	Propionibacterium granulosum	98.46	0.44	SUC 93	90			
	Atopobium minutum	1.55	0.11	MLZ 12	1			
05030040	**Low discrimination**							**05030040**
					SPO	CAT		
	Clostridium botulinum biovar B	72.52	0.50	SUC 90	99	1		
	Clostridium botulinum biovar A	8.55	0.36	MLT 99	99	1		
	Propionibacterium granulosum	7.03	0.45		1	90		
05030050	**Low discrimination**							**05030050**
					SPO	CAT		
	Clostridium botulinum biovar B	77.32	0.44	SUC 90	99	1		
	Clostridium difficile	20.27	0.34	MLT 99	99	1		
	Propionibacterium granulosum	2.07	0.30		1	90		
05030060	**Low discrimination**							**05030060**
					SPO	CAT		
	Clostridium botulinum biovar B	89.56	0.44	SUC 90	99	1		
	Propionibacterium granulosum	4.40	0.35	MLT 99	1	90		
	Propionibacterium acnes	2.98	0.22		1	95		
05030070	**Good identification**							**05030070**
					SPO	CAT		
	Clostridium botulinum biovar B	98.14	0.39	SUC 90	99	1		
	Propionibacterium granulosum	1.34	0.20	MLT 99	1	90		
	Clostridium difficile	0.53	0.00		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05030100	**Low discrimination**							**05030100**
					GRA	CAT		
Fusobacterium mortiferum		77.24	0.62		1	5		
Propionibacterium granulosum		11.52	0.45		99	90		
Fusobacterium varium		6.34	0.24		1	1		
05030102	**Low discrimination**							**05030102**
					GRA	CAT		
Fusobacterium mortiferum		72.00	0.46	BGL 10	1	5		
Propionibacterium granulosum		24.16	0.35		99	90		
Atopobium minutum		3.84	0.19		99	1		
05030200	**Low discrimination**							**05030200**
					GRA	CAT		
Propionibacterium granulosum		75.45	0.53	SUC 93	99	90		
Fusobacterium varium		13.57	0.24	RHA 3	1	1		
Eubacterium limosum		3.14	0.15		99	1		
05030202	**Good identification**							**05030202**
					CAT			
Propionibacterium granulosum		95.07	0.43	SUC 93	90			
Atopobium minutum		4.93	0.19	RHA 3	1			
05030210	**Low discrimination**							**05030210**
					SPO	CAT		
Propionibacterium granulosum		62.01	0.38	SUC 93	1	90		
Clostridium difficile		37.99	0.22	RHA 3	99	1		
05030220	**Low discrimination**							**05030220**
					SPO	CAT		
Propionibacterium granulosum		88.26	0.43	SUC 93	1	90		
Clostridium cadaveris		7.57	0.12	RHA 3	99	1		
Atopobium minutum		2.29	0.14		1	1		
05030400	**Low discrimination**							**05030400**
					COC	GRA	CAT	
Propionibacterium granulosum		36.13	0.45	SUC 93	1	99	90	
Gemella morbillorum		30.68	0.37	ARA 1	99	99	1	
Fusobacterium varium		19.89	0.24		1	1	1	
05031000	**Low discrimination**							**05031000**
					COC	GRA	CAT	
Propionibacterium granulosum		89.64	0.79	SUC 93	1	99	90	
Fusobacterium mortiferum		3.90	0.59		1	1	5	
Gemella morbillorum		3.73	0.48		99	99	1	
05031001	**Acceptable identification**							**05031001**
					COC	GRA	CAT	
Propionibacterium granulosum		90.50	0.45	SUC 93	1	99	90	
Fusobacterium mortiferum		3.94	0.25	URE 1	1	1	5	
Gemella morbillorum		3.76	0.14		99	99	1	
05031002	**Good identification**							**05031002**
					GRA	CAT		
Propionibacterium granulosum		92.73	0.69	SUC 93	99	90		
Atopobium minutum		4.91	0.45		99	1		
Fusobacterium mortiferum		1.79	0.43		1	5		
05031010	**Very good identification**							**05031010**
					GRA	SPO	CAT	
Propionibacterium granulosum		97.24	0.64	SUC 93	99	1	90	
Clostridium difficile		1.84	0.22	MLZ 12	99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		0.31	0.25		1	1	5	
05031012	**Good identification**							**05031012**
					CAT			
Propionibacterium granulosum		99.18	0.54	SUC 93	90			
Lactobacillus jensenii		0.44	0.07	MLZ 12	1			
Atopobium minutum		0.39	0.11		1			
05031020	**Low discrimination**							**05031020**
					CAT			
Propionibacterium granulosum		89.53	0.69	SUC 93	90			
Propionibacterium acnes		2.79	0.34		95			
Atopobium minutum		2.37	0.40		1			
05031022	**Low discrimination**							**05031022**
					CAT			
Propionibacterium granulosum		88.98	0.58	SUC 93	90			
Atopobium minutum		6.28	0.37		1			
Clostridium perfringens		4.20	0.28		1			
05031030	**Good identification**							**05031030**
					CAT			
Propionibacterium granulosum		99.58	0.54	SUC 93	90			
Propionibacterium acnes		0.23	-.01	MLZ 12	95			
Atopobium minutum		0.20	0.05		1			
05031032	**Good identification**							**05031032**
Propionibacterium granulosum		100.0	0.44	SUC 93				
05031040	**Low discrimination**							**05031040**
					GRA	SPO	CAT	
Propionibacterium granulosum		81.87	0.45	SUC 93	99	1	90	
Clostridium botulinum biovar B		8.54	0.15	SOR 1	99	99	1	
Fusobacterium mortiferum		3.56	0.25		1	1	5	
05031100	**Low discrimination**							**05031100**
					GRA	SPO	CAT	
Fusobacterium mortiferum		85.58	0.65		1	1	5	
Propionibacterium granulosum		8.51	0.45		99	1	90	
Clostridium septicum		2.09	0.20		99	99	1	
05031102	**Low discrimination**							**05031102**
					GRA	CAT		
Fusobacterium mortiferum		71.52	0.49	BGL 10	1	5		
Propionibacterium granulosum		16.00	0.35		99	90		
Lactobacillus fermentum		5.77	0.32		99	1		
05031120	**Low discrimination**							**05031120**
					CAT	SPO		
Clostridium septicum		82.87	0.54	MLT 94	1	99		
Clostridium perfringens		15.51	0.49		1	84		
Propionibacterium granulosum		0.85	0.35		90	1		
05031122	**Low discrimination**							**05031122**
					CAT	SPO		
Clostridium perfringens		77.73	0.43	SUC 91	1	84		
Clostridium septicum		18.01	0.25	MLT 99	1	99		
Propionibacterium granulosum		2.27	0.24		90	1		
05031160	**Low discrimination**							**05031160**
Clostridium perfringens		77.92	0.36	SUC 91				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium septicum		22.08	0.20	MLT 99				
05031200	**Good identification**							**05031200**
					COC	GRA	CAT	
Propionibacterium granulosum		96.69	0.53	SUC 93	1	99	90	
Fusobacterium mortiferum		1.38	0.25	RHA 3	1	1	5	
Gemella morbillorum		1.31	0.14		99	99	1	
05031202	**Good identification**							**05031202**
					CAT			
Propionibacterium granulosum		98.30	0.43	SUC 93	90			
Atopobium minutum		1.70	0.11	RHA 3	1			
05031210	**Good identification**							**05031210**
Propionibacterium granulosum		100.0	0.38	SUC 93				
05031220	**Good identification**							**05031220**
					CAT			
Propionibacterium granulosum		98.15	0.43	SUC 93	90			
Propionibacterium acnes		1.00	-.01	RHA 3	95			
Atopobium minutum		0.85	0.05		1			
05031400	**Low discrimination**							**05031400**
					COC	CAT		
Propionibacterium granulosum		63.46	0.45	SUC 93	1	90		
Gemella morbillorum		29.01	0.32	ARA 1	99	1		
Lactobacillus fermentum		3.51	0.28		1	1		
05032000	**Low discrimination**							**05032000**
					SPO	CAT		
Eubacterium limosum		38.63	0.63	TRE 1	1	1		
Propionibacterium granulosum		37.22	0.77		1	90		
Clostridium difficile		23.26	0.61		99	1		
05032002	**Good identification**							**05032002**
					SPO	CAT		
Propionibacterium granulosum		93.01	0.67	SUC 93	1	90		
Eubacterium limosum		3.90	0.29		1	1		
Clostridium difficile		2.35	0.27		99	1		
05032010	**Very good identification**							**05032010**
					SPO	CAT		
Clostridium difficile		95.41	0.73	TRE 7	99	1		
Propionibacterium granulosum		4.26	0.62		1	90		
Eubacterium limosum		0.33	0.29		1	1		
05032011	**Good identification**							**05032011**
					SPO	CAT		
Clostridium difficile		95.41	0.38	TRE 7	99	1		
Propionibacterium granulosum		4.26	0.28	URE 1	1	90		
Eubacterium limosum		0.33	-.06		1	1		
05032012	**Low discrimination**							**05032012**
					SPO	CAT		
Propionibacterium granulosum		52.43	0.52	SUC 93	1	90		
Clostridium difficile		47.40	0.38	MLZ 12	99	1		
Eubacterium limosum		0.16	-.06		1	1		
05032020	**Good identification**							**05032020**
					SPO	CAT		
Propionibacterium granulosum		90.64	0.67	SUC 93	1	90		
Eubacterium limosum		3.80	0.29		1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		2.29	0.27		99	1	
05032022	**Very good identification**						**05032022**
					CAT		
Propionibacterium granulosum		99.21	0.56	SUC 93	90		
Pseudoramibacter alactolyticus		0.34	-.02		1		
Atopobium minutum		0.28	0.11		1		
05032030	**Low discrimination**						**05032030**
					SPO	CAT	
Propionibacterium granulosum		52.43	0.52	SUC 93	1	90	
Clostridium difficile		47.40	0.38	MLZ 12	99	1	
Eubacterium limosum		0.16	-.06		1	1	
05032032	**Good identification**						**05032032**
					SPO	CAT	
Propionibacterium granulosum		96.48	0.42	SUC 93	1	90	
Clostridium difficile		3.52	0.04	MLZ 12	99	1	
05032040	**Low discrimination**						**05032040**
					SPO	CAT	
Clostridium difficile		55.54	0.39	TRE 7	99	1	
Eubacterium limosum		17.71	0.29	SOR 5	1	1	
Propionibacterium granulosum		17.06	0.43		1	90	
05032050	**Good identification**						**05032050**
					SPO	CAT	
Clostridium difficile		98.67	0.51	TRE 7	99	1	
Propionibacterium granulosum		0.85	0.28	SOR 5	1	90	
Clostridium botulinum biovar B		0.42	0.10		99	1	
05032110	**Good identification**						**05032110**
					SPO	CAT	
Clostridium difficile		95.41	0.38	TRE 7	99	1	
Propionibacterium granulosum		4.26	0.28	LAC 1	1	90	
Eubacterium limosum		0.33	-.06		1	1	
05032200	**Low discrimination**						**05032200**
					SPO	CAT	
Propionibacterium granulosum		64.65	0.51	SUC 93	1	90	
Eubacterium limosum		21.92	0.29	RHA 3	1	1	
Clostridium difficile		13.20	0.27		99	1	
05032202	**Good identification**						**05032202**
					CAT		
Propionibacterium granulosum		98.65	0.41	SUC 93	90		
Eubacterium limosum		1.35	-.06	RHA 3	1		
05032210	**Low discrimination**						**05032210**
					SPO	CAT	
Clostridium difficile		87.70	0.38	TRE 7	99	1	
Propionibacterium granulosum		12.00	0.36	RHA 1	1	90	
Eubacterium limosum		0.30	-.06		1	1	
05032220	**Good identification**						**05032220**
					CAT		
Propionibacterium granulosum		98.65	0.41	SUC 93	90		
Eubacterium limosum		1.35	-.06	RHA 3	1		
05032410	**Good identification**						**05032410**
					SPO	CAT	
Clostridium difficile		95.41	0.38	TRE 7	99	1	
Propionibacterium granulosum		4.26	0.28	ARA 1	1	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		0.33	-.06		1	1	
05033000	**Very good identification**						**05033000**
					SPO	CAT	
Propionibacterium granulosum		98.10	0.77	SUC 93	1	90	
Eubacterium limosum		1.03	0.29		1	1	
Clostridium difficile		0.62	0.27		99	1	
05033001	**Good identification**						**05033001**
					CAT		
Propionibacterium granulosum		98.96	0.43	SUC 93	90		
Eubacterium limosum		1.04	-.06	URE 1	1		
05033002	**Very good identification**						**05033002**
					SPO	CAT	
Propionibacterium granulosum		99.46	0.67	SUC 93	1	90	
Clostridium innocuum		0.43	0.17		99	1	
Atopobium minutum		0.07	0.11		1	1	
05033010	**Low discrimination**						**05033010**
					SPO	CAT	
Propionibacterium granulosum		81.51	0.62	SUC 93	1	90	
Clostridium difficile		18.42	0.38	MLZ 12	99	1	
Eubacterium limosum		0.06	-.06		1	1	
05033012	**Good identification**						**05033012**
					SPO	CAT	
Propionibacterium granulosum		99.10	0.52	SUC 93	1	90	
Clostridium difficile		0.91	0.04	MLZ 12	99	1	
05033020	**Very good identification**						**05033020**
					CAT		
Propionibacterium granulosum		97.66	0.67	SUC 93	90		
Propionibacterium acnes		2.27	0.29		95		
Eubacterium limosum		0.04	-.06		1		
05033022	**Very good identification**						**05033022**
Propionibacterium granulosum		100.0	0.56	SUC 93			
05033030	**Good identification**						**05033030**
					SPO	CAT	
Propionibacterium granulosum		99.10	0.52	SUC 93	1	90	
Clostridium difficile		0.91	0.04	MLZ 12	99	1	
05033032	**Good identification**						**05033032**
Propionibacterium granulosum		100.0	0.42	SUC 93			
05033040	**Good identification**						**05033040**
					SPO	CAT	
Propionibacterium granulosum		95.84	0.43	SUC 93	1	90	
Clostridium difficile		3.15	0.05	SOR 1	99	1	
Eubacterium limosum		1.01	-.06		1	1	
05033100	**Low discrimination**						**05033100**
					GRA	CAT	
Propionibacterium granulosum		87.33	0.43	SUC 93	99	90	
Fusobacterium mortiferum		11.76	0.31	LAC 1	1	5	
Eubacterium limosum		0.92	-.06		99	1	
05033200	**Good identification**						**05033200**
					CAT		
Propionibacterium granulosum		99.66	0.51	SUC 93	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium limosum		0.34	-.06	RHA 3	1			
05033202	**Good identification**							**05033202**
Propionibacterium granulosum		100.0	0.41	SUC 93				
05033210	**Acceptable identification**							**05033210**
					SPO	CAT		
Propionibacterium granulosum		93.13	0.36	SUC 93	1	90		
Clostridium difficile		6.87	0.04	RHA 3	99	1		
05033220	**Good identification**							**05033220**
Propionibacterium granulosum		100.0	0.41	SUC 93				
05033400	**Good identification**							**05033400**
					SPO	CAT		
Propionibacterium granulosum		97.14	0.43	SUC 93	1	90		
Clostridium innocuum		1.84	0.04	ARA 1	99	1		
Eubacterium limosum		1.02	-.06		1	1		
05034100	**Low discrimination**							**05034100**
					GRA			
Fusobacterium mortiferum		89.58	0.43	XYL 7	1			
Fusobacterium varium		4.07	0.01		1			
Actinomyces meyeri		3.03	0.09		99			
05035000	**Low discrimination**							**05035000**
					COC	GRA	CAT	
Propionibacterium granulosum		53.85	0.45	SUC 93	1	99	90	
Gemella morbillorum		24.62	0.32	XYL 1	99	99	1	
Fusobacterium mortiferum		17.47	0.40		1	1	5	
05035100	**Low discrimination**							**05035100**
					COC	GRA		
Fusobacterium mortiferum		68.14	0.46	XYL 7	1	1		
Atopobium parvulum		23.88	0.27		99	99		
Lactobacillus fermentum		4.77	0.28		1	99		
05035322	**Good identification**							**05035322**
Parabacteroides distasonis		100.0	0.39	RAF 99				
05035722	**Good identification**							**05035722**
Parabacteroides distasonis		100.0	0.36	RAF 99				
05036000	**Low discrimination**							**05036000**
					SPO	CAT		
Clostridium difficile		60.90	0.39	TRE 7	99	1		
Eubacterium limosum		19.41	0.29	XYL 5	1	1		
Propionibacterium granulosum		18.70	0.43		1	90		
05036010	**Good identification**							**05036010**
					SPO	CAT		
Clostridium difficile		99.09	0.51	TRE 7	99	1		
Propionibacterium granulosum		0.85	0.28	XYL 5	1	90		
Eubacterium limosum		0.07	-.06		1	1		
05037000	**Good identification**							**05037000**
					SPO	CAT		
Propionibacterium granulosum		95.84	0.43	SUC 93	1	90		
Clostridium difficile		3.15	0.05	XYL 1	99	1		
Eubacterium limosum		1.01	-.06		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05040000	**Low discrimination**						**05040000**
					COC		
Atopobium minutum		23.12	0.56	CEL 1	1		
Gemella morbillorum		22.63	0.52		99		
Peptostreptococcus anaerobius		17.94	0.42		99		
05040002	**Acceptable identification**						**05040002**
					COC		
Atopobium minutum		92.63	0.53	CEL 1	1		
Gemella morbillorum		1.37	0.17		99		
Peptostreptococcus anaerobius		1.09	0.08		99		
05040020	**Low discrimination**						**05040020**
					COC	SPO	
Atopobium minutum		48.86	0.48	CEL 1	1	1	
Clostridium cadaveris		28.61	0.34		1	99	
Gemella morbillorum		15.94	0.35		99	1	
05040022	**Good identification**						**05040022**
					GRA	SPO	
Atopobium minutum		97.59	0.45	CEL 1	99	1	
Pseudoflavonifractor capillosus		1.07	-.03		1	1	
Clostridium cadaveris		0.87	0.00		99	99	
05040400	**Low discrimination**						**05040400**
					COC		
Gemella morbillorum		77.82	0.35	CEL 1	99		
Atopobium minutum		7.23	0.22	ARA 10	1		
Peptostreptococcus anaerobius		5.61	0.08		99		
05041000	**Low discrimination**						**05041000**
					COC		
Gemella morbillorum		59.01	0.47	CEL 1	99		
Atopobium minutum		37.33	0.48		1		
Peptostreptococcus anaerobius		0.88	0.08		99		
05041002	**Good identification**						**05041002**
					COC	SPO	
Atopobium minutum		96.85	0.45	CEL 1	1	1	
Gemella morbillorum		2.32	0.13		99	1	
Clostridium innocuum		0.83	0.03		1	99	
05041020	**Low discrimination**						**05041020**
					COC	SPO	
Atopobium minutum		64.28	0.40	CEL 1	1	1	
Gemella morbillorum		33.88	0.31		99	1	
Clostridium cadaveris		1.14	0.00		1	99	
05041022	**Good identification**						**05041022**
Atopobium minutum		100.0	0.37	CEL 1			
05042010	**Good identification**						**05042010**
					SPO		
Clostridium difficile		99.51	0.36	FRU 99	99		
Eubacterium limosum		0.49	-.06	CEL 5	1		
05043000	**Acceptable identification**						**05043000**
					COC	SPO	
Clostridium innocuum		94.99	0.37	SAL 99	1	99	
Gemella morbillorum		2.69	0.13	FRU 95	99	1	
Atopobium minutum		1.70	0.14		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05043002	**Good identification**						**05043002**
					SPO		
Clostridium innocuum		98.82	0.37	SAL 99	99		
Atopobium minutum		1.18	0.11	FRU 95	1		
05044000	**Low discrimination**						**05044000**
					COC		
Gemella morbillorum		75.87	0.35	CEL 1	99		
Atopobium minutum		7.05	0.22	XYL 10	1		
Peptostreptococcus anaerobius		5.47	0.08		99		
05050000	**Low discrimination**						**05050000**
					COC	GRA	
Fusobacterium varium		57.86	0.58	CEL 1	1	1	
Eubacterium limosum		13.39	0.49		1	99	
Gemella morbillorum		8.11	0.53		99	99	
05050002	**Low discrimination**						**05050002**
					GRA		
Atopobium minutum		72.73	0.53	CEL 1	99		
Fusobacterium varium		10.19	0.24		1		
Fusobacterium mortiferum		6.59	0.40		1		
05050010	**Acceptable identification**						**05050010**
					GRA	SPO	
Clostridium difficile		94.83	0.54	CEL 5	99	99	
Fusobacterium varium		2.95	0.24	MAN 90	1	1	
Eubacterium limosum		0.68	0.15		99	1	
05050020	**Low discrimination**						**05050020**
					COC	SPO	
Atopobium minutum		32.89	0.48	CEL 1	1	1	
Clostridium cadaveris		20.87	0.34		1	99	
Gemella morbillorum		14.21	0.37		99	1	
05050022	**Acceptable identification**						**05050022**
					CAT		
Atopobium minutum		94.27	0.45	CEL 1	1		
Pseudoramibacter alactolyticus		2.15	0.03		1		
Propionibacterium granulosum		1.50	0.16		90		
05050100	**Low discrimination**						**05050100**
					GRA		
Fusobacterium mortiferum		89.06	0.63		1		
Fusobacterium varium		6.56	0.24		1		
Eubacterium limosum		1.52	0.15		99		
05050102	**Good identification**						**05050102**
					GRA		
Fusobacterium mortiferum		95.44	0.47	BGL 10	1		
Atopobium minutum		4.56	0.19		99		
05050400	**Low discrimination**						**05050400**
					COC	GRA	
Gemella morbillorum		49.53	0.37	CEL 1	99	99	
Fusobacterium varium		32.12	0.24	ARA 10	1	1	
Eubacterium limosum		7.44	0.15		1	99	
05051000	**Low discrimination**						**05051000**
					COC	GRA	
Fusobacterium mortiferum		37.53	0.60		1	1	
Gemella morbillorum		32.11	0.48		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		15.34	0.48		1	99	
05051002	**Low discrimination**						**05051002**
					GRA		
Atopobium minutum		51.03	0.45	CEL 1	99		
Fusobacterium mortiferum		20.81	0.43		1		
Lactobacillus jensenii		10.96	0.29		99		
05051022	**Low discrimination**						**05051022**
					CAT		
Atopobium minutum		75.69	0.37	CEL 1	1		
Lactobacillus jensenii		12.19	0.18		1		
Propionibacterium granulosum		3.61	0.16		90		
05051100	**Very good identification**						**05051100**
					GRA	SPO	
Fusobacterium mortiferum		95.97	0.66		1	1	
Clostridium septicum		2.22	0.21		99	99	
Lactobacillus fermentum		0.77	0.32		99	1	
05051102	**Low discrimination**						**05051102**
					GRA		
Fusobacterium mortiferum		87.98	0.50	BGL 10	1		
Lactobacillus fermentum		6.37	0.32		99		
Lactobacillus catenaformis		3.28	0.15		99		
05051120	**Good identification**						**05051120**
					GRA	SPO	
Clostridium septicum		98.65	0.55	MLT 94	99	99	
Fusobacterium mortiferum		0.44	0.32		1	1	
Clostridium perfringens		0.39	0.20		99	84	
05052000	**Low discrimination**						**05052000**
					SPO		
Eubacterium limosum		68.39	0.63	CEL 1	1		
Clostridium difficile		28.79	0.58		99		
Clostridium innocuum		1.06	0.37		99		
05052002	**Low discrimination**						**05052002**
					SPO		
Clostridium innocuum		47.58	0.37	SAL 99	99		
Eubacterium limosum		30.99	0.29	GAL 95	1		
Clostridium difficile		13.05	0.24		99		
05052010	**Very good identification**						**05052010**
					SPO		
Clostridium difficile		99.45	0.70	CEL 5	99		
Eubacterium limosum		0.49	0.29		1		
Clostridium innocuum		0.03	0.13		99		
05052011	**Good identification**						**05052011**
					SPO		
Clostridium difficile		99.51	0.36	CEL 5	99		
Eubacterium limosum		0.49	-.06	URE 1	1		
05052012	**Good identification**						**05052012**
					SPO	CAT	
Clostridium difficile		96.03	0.36	CEL 5	99	1	
Clostridium innocuum		2.99	0.13	BGL 1	99	1	
Propionibacterium granulosum		0.51	0.09		1	90	
05052030	**Good identification**						**05052030**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		98.99	0.36	CEL 5	99	1	
Propionibacterium granulosum		0.53	0.09	NAG 1	1	90	
Eubacterium limosum		0.49	-.06		1	1	
05052040	**Low discrimination**						**05052040**
					SPO		
Clostridium difficile		60.80	0.36	CEL 5	99		
Eubacterium limosum		27.72	0.29	SOR 5	1		
Pseudoramibacter alactolyticus		7.26	0.17		1		
05052050	**Good identification**						**05052050**
					SPO		
Clostridium difficile		99.91	0.48	CEL 5	99		
Eubacterium limosum		0.09	-.06	SOR 5	1		
05052110	**Good identification**						**05052110**
					SPO		
Clostridium difficile		99.51	0.36	CEL 5	99		
Eubacterium limosum		0.49	-.06	LAC 1	1		
05052210	**Good identification**						**05052210**
					SPO		
Clostridium difficile		99.51	0.36	CEL 5	99		
Eubacterium limosum		0.49	-.06	RHA 1	1		
05052410	**Good identification**						**05052410**
					SPO		
Clostridium difficile		98.97	0.36	CEL 5	99		
Clostridium innocuum		0.54	0.00	ARA 1	99		
Eubacterium limosum		0.49	-.06		1		
05053000	**Good identification**						**05053000**
					SPO		
Clostridium innocuum		93.91	0.59	SAL 99	99		
Eubacterium limosum		3.22	0.29		1		
Clostridium difficile		1.36	0.24		99		
05053002	**Very good identification**						**05053002**
					SPO	CAT	
Clostridium innocuum		99.63	0.59	SAL 99	99	1	
Propionibacterium granulosum		0.27	0.24		1	90	
Atopobium minutum		0.06	0.11		1	1	
05053010	**Low discrimination**						**05053010**
					SPO	CAT	
Clostridium difficile		61.82	0.36	CEL 5	99	1	
Clostridium innocuum		36.56	0.35	GAL 1	99	1	
Propionibacterium granulosum		1.32	0.20		1	90	
05053012	**Good identification**						**05053012**
					SPO	CAT	
Clostridium innocuum		97.46	0.35	SAL 99	99	1	
Clostridium difficile		1.67	0.02	MLZ 4	99	1	
Propionibacterium granulosum		0.88	0.09		1	90	
05053400	**Good identification**						**05053400**
					SPO		
Clostridium innocuum		99.80	0.46	SAL 99	99		
Eubacterium limosum		0.20	-.06		1		
05053402	**Good identification**						**05053402**
Clostridium innocuum		100.0	0.46	SAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05054100	**Acceptable identification**							**05054100**
					GRA			
Fusobacterium mortiferum		92.06	0.44	XYL 7	1			
Fusobacterium varium		3.75	0.01		1			
Actinomyces meyeri		2.79	0.09		99			
05055100	**Acceptable identification**							**05055100**
					GRA			
Fusobacterium mortiferum		91.49	0.47	XYL 7	1			
Lactobacillus fermentum		5.74	0.28		99			
Actinomyces meyeri		2.77	0.12		99			
05056000	**Low discrimination**							**05056000**
					SPO			
Clostridium difficile		65.97	0.36	CEL 5	99			
Eubacterium limosum		30.07	0.29	XYL 5	1			
Clostridium innocuum		2.43	0.15		99			
05056010	**Good identification**							**05056010**
					SPO			
Clostridium difficile		99.91	0.48	CEL 5	99			
Eubacterium limosum		0.09	-.06	XYL 5	1			
05057000	**Good identification**							**05057000**
					SPO			
Clostridium innocuum		97.94	0.37	SAL 99	99			
Clostridium difficile		1.41	0.02	XYL 5	99			
Eubacterium limosum		0.64	-.06		1			
05057002	**Good identification**							**05057002**
Clostridium innocuum		100.0	0.37	SAL 99				
05070000	**Low discrimination**							**05070000**
					GRA	CAT		
Propionibacterium granulosum		31.66	0.45	SUC 93	99	90		
Fusobacterium mortiferum		30.33	0.48	CEL 1	1	5		
Fusobacterium varium		17.44	0.24		1	1		
05070100	**Good identification**							**05070100**
					GRA	SPO	CAT	
Fusobacterium mortiferum		99.20	0.54		1	1	5	
Propionibacterium granulosum		0.45	0.11		99	1	90	
Clostridium septicum		0.35	-.05		99	99	1	
05070102	**Good identification**							**05070102**
Fusobacterium mortiferum		100.0	0.38	BGL 10				
05071000	**Low discrimination**							**05071000**
					GRA	CAT		
Fusobacterium mortiferum		37.28	0.51		1	5		
Lactobacillus jensenii		32.86	0.40		99	1		
Propionibacterium granulosum		25.95	0.45		99	90		
05071002	**Low discrimination**							**05071002**
					GRA	CAT		
Lactobacillus jensenii		62.62	0.37	SUC 90	99	1		
Propionibacterium granulosum		18.54	0.35	MLT 95	99	90		
Fusobacterium mortiferum		11.84	0.34	SAL 95	1	5		
05071100	**Low discrimination**							**05071100**
					COC	GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		74.71	0.57		1	1	1	
Clostridium septicum		17.35	0.29		1	99	99	
Atopobium parvulum		7.43	0.29		99	99	1	
05071102	**Low discrimination**							**05071102**
					COC	GRA		
Fusobacterium mortiferum		75.20	0.41	BGL 10	1	1		
Atopobium parvulum		11.88	0.16		99	99		
Lactobacillus acidophilus		8.00	0.11		1	99		
05071120	**Very good identification**							**05071120**
					COC	SPO		
Clostridium septicum		99.64	0.63	MLT 94	1	99		
Clostridium perfringens		0.25	0.25		1	84		
Atopobium parvulum		0.08	0.16		99	1		
05072000	**Low discrimination**							**05072000**
					SPO	CAT		
Clostridium difficile		55.40	0.39	TRE 7	99	1		
Eubacterium limosum		17.66	0.29	CEL 5	1	1		
Propionibacterium granulosum		17.01	0.43		1	90		
05072010	**Good identification**							**05072010**
					SPO	CAT		
Clostridium difficile		98.94	0.51	TRE 7	99	1		
Propionibacterium granulosum		0.85	0.28	CEL 5	1	90		
Clostridium innocuum		0.15	0.06		99	1		
05073000	**Acceptable identification**							**05073000**
					SPO	CAT		
Clostridium innocuum		90.79	0.51	SAL 99	99	1		
Propionibacterium granulosum		8.53	0.43		1	90		
Clostridium difficile		0.28	0.05		99	1		
05073002	**Good identification**							**05073002**
					SPO	CAT		
Clostridium innocuum		97.60	0.51	SAL 99	99	1		
Propionibacterium granulosum		2.29	0.32		1	90		
Lactobacillus jensenii		0.10	0.03		1	1		
05073400	**Good identification**							**05073400**
					SPO	CAT		
Clostridium innocuum		99.47	0.38	SAL 99	99	1		
Propionibacterium granulosum		0.54	0.09		1	90		
05073402	**Good identification**							**05073402**
Clostridium innocuum		100.0	0.38	SAL 99				
05100000	**Low discrimination**							**05100000**
					COC	GRA		
Prevotella disiens		75.63	1.00		1	1		
Gemella morbillorum		19.36	0.99		99	99		
Peptostreptococcus anaerobius		1.81	0.73		99	99		
05100001	**Low discrimination**							**05100001**
					COC	GRA	SPO	
Prevotella disiens		68.35	0.66	URE 1	1	1	1	
Gemella morbillorum		17.49	0.64		99	99	1	
Clostridium novyi biovar A		6.43	0.66		1	99	99	
05100002	**Low discrimination**							**05100002**
					COC	GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella disiens		71.14	0.66	BGL 1	1	1	1	
Gemella morbillorum		18.21	0.64		99	99	1	
Clostridium sporogenes		2.63	0.45		1	99	99	
05100003	**Low discrimination**							**05100003**
					COC	GRA	SPO	
Clostridium novyi biovar A		49.46	0.53	MNS 1	1	99	99	
Prevotella disiens		30.11	0.32		1	1	1	
Gemella morbillorum		7.71	0.30		99	99	1	
05100010	**Low discrimination**							**05100010**
					COC	GRA	SPO	
Prevotella disiens		70.49	0.66	MLZ 1	1	1	1	
Gemella morbillorum		18.04	0.64		99	99	1	
Clostridium novyi biovar A		6.63	0.66		1	99	99	
05100011	**Low discrimination**							**05100011**
					COC	GRA	SPO	
Clostridium novyi biovar A		87.20	0.66	MNS 1	1	99	99	
Prevotella disiens		9.37	0.32		1	1	1	
Gemella morbillorum		2.40	0.30		99	99	1	
05100012	**Low discrimination**							**05100012**
					COC	GRA	SPO	
Clostridium novyi biovar A		54.23	0.53	MNS 1	1	99	99	
Prevotella disiens		33.01	0.32		1	1	1	
Gemella morbillorum		8.45	0.30		99	99	1	
05100013	**Good identification**							**05100013**
					GRA	SPO		
Clostridium novyi biovar A		99.39	0.53	MNS 1	99	99		
Prevotella disiens		0.61	-.03		1	1		
05100020	**Low discrimination**							**05100020**
					COC	GRA		
Gemella morbillorum		45.78	0.82	NAG 10	99	99		
Prevotella bivia		33.41	0.66		1	1		
Prevotella disiens		16.26	0.66		1	1		
05100021	**Low discrimination**							**05100021**
					COC	GRA	SPO	
Gemella morbillorum		33.14	0.48	NAG 10	99	99	1	
Clostridium novyi biovar A		27.39	0.56	URE 1	1	99	99	
Prevotella bivia		24.19	0.32		1	1	1	
05100022	**Low discrimination**							**05100022**
					COC	GRA		
Gemella morbillorum		37.11	0.48	NAG 10	99	99		
Prevotella bivia		27.09	0.32	BGL 1	1	1		
Atopobium minutum		13.38	0.45		1	99		
05100023	**Low discrimination**							**05100023**
					COC	GRA	SPO	
Clostridium novyi biovar A		85.52	0.43	MNS 1	1	99	99	
Gemella morbillorum		5.92	0.14		99	99	1	
Prevotella bivia		4.32	-.03		1	1	1	
05100030	**Low discrimination**							**05100030**
					COC	GRA	SPO	
Gemella morbillorum		32.75	0.48	MLZ 1	99	99	1	
Clostridium novyi biovar A		27.07	0.56	NAG 10	1	99	99	
Prevotella bivia		23.90	0.32		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05100031	**Good identification**							**05100031**
					COC	GRA	SPO	
Clostridium novyi biovar A		97.52	0.56	MNS 1	1	99	99	
Gemella morbillorum		1.19	0.14		99	99	1	
Prevotella bivia		0.87	-.03		1	1	1	
05100032	**Low discrimination**							**05100032**
					COC	GRA	SPO	
Clostridium novyi biovar A		85.52	0.43	MNS 1	1	99	99	
Gemella morbillorum		5.92	0.14		99	99	1	
Prevotella bivia		4.32	-.03		1	1	1	
05100033	**Good identification**							**05100033**
Clostridium novyi biovar A		100.0	0.43	MNS 1				
05100040	**Low discrimination**							**05100040**
					GRA	SPO		
Prevotella disiens		39.73	0.66	SOR 1	1	1		
Clostridium botulinum biovar B		24.43	0.65		99	99		
Clostridium botulinum biovar A		17.91	0.66		99	99		
05100041	**Low discrimination**							**05100041**
					GRA	SPO		
Clostridium novyi biovar A		70.36	0.63	MNS 1	99	99		
Prevotella disiens		11.34	0.32		1	1		
Clostridium botulinum biovar B		6.97	0.31		99	99		
05100042	**Low discrimination**							**05100042**
					GRA	SPO		
Clostridium novyi biovar A		29.82	0.50	MNS 1	99	99		
Prevotella disiens		27.23	0.32		1	1		
Clostridium botulinum biovar B		16.75	0.31		99	99		
05100043	**Good identification**							**05100043**
					GRA	SPO		
Clostridium novyi biovar A		98.53	0.50	MNS 1	99	99		
Prevotella disiens		0.91	-.03		1	1		
Clostridium botulinum biovar B		0.56	-.03		99	99		
05100050	**Low discrimination**							**05100050**
					GRA	SPO		
Clostridium botulinum biovar B		78.86	0.60	SUC 90	99	99		
Clostridium novyi biovar A		16.32	0.63		99	99		
Prevotella disiens		2.63	0.32		1	1		
05100051	**Very good identification**							**05100051**
					GRA	SPO		
Clostridium novyi biovar A		95.20	0.63	MNS 1	99	99		
Clostridium botulinum biovar B		4.65	0.26		99	99		
Prevotella disiens		0.16	-.03		1	1		
05100052	**Low discrimination**							**05100052**
					GRA	SPO		
Clostridium novyi biovar A		77.77	0.50	MNS 1	99	99		
Clostridium botulinum biovar B		21.51	0.26		99	99		
Prevotella disiens		0.72	-.03		1	1		
05100053	**Good identification**							**05100053**
Clostridium novyi biovar A		100.0	0.50	MNS 1				
05100060	**Low discrimination**							**05100060**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Clostridium botulinum biovar B		77.76	0.60	SUC 90	1	99	99	
Gemella morbillorum		7.30	0.48		99	99	1	
Prevotella bivia		5.33	0.32		1	1	1	
05100061	**Low discrimination**							**05100061**
					COC	SPO		
Clostridium novyi biovar A		80.81	0.53	MNS 1	1	99		
Clostridium botulinum biovar B		15.78	0.26		1	99		
Gemella morbillorum		1.48	0.14		99	1		
05100063	**Good identification**							**05100063**
Clostridium novyi biovar A		100.0	0.40	MNS 1				
05100070	**Acceptable identification**							**05100070**
					COC	SPO		
Clostridium botulinum biovar B		90.13	0.55	SUC 90	1	99		
Clostridium novyi biovar A		9.47	0.53		1	99		
Gemella morbillorum		0.17	0.14		99	1		
05100071	**Acceptable identification**							**05100071**
Clostridium novyi biovar A		91.23	0.53	MNS 1				
Clostridium botulinum biovar B		8.77	0.21					
05100072	**Low discrimination**							**05100072**
Clostridium novyi biovar A		64.73	0.40	MNS 1				
Clostridium botulinum biovar B		35.28	0.21					
05100073	**Good identification**							**05100073**
Clostridium novyi biovar A		100.0	0.40	MNS 1				
05100100	**Low discrimination**							**05100100**
					COC	GRA		
Prevotella bivia		60.84	0.66	NAG 99	1	1		
Prevotella disiens		29.61	0.66		1	1		
Gemella morbillorum		7.58	0.64		99	99		
05100120	**Excellent identification**							**05100120**
					COC	GRA		
Prevotella bivia		99.98	1.00		1	1		
Gemella morbillorum		0.01	0.48		99	99		
Prevotella disiens		0.01	0.32		1	1		
05100121	**Very good identification**							**05100121**
					COC	GRA	SPO	
Prevotella bivia		99.96	0.66	URE 1	1	1	1	
Gemella morbillorum		0.01	0.14		99	99	1	
Clostridium novyi biovar A		0.01	0.21		1	99	99	
05100122	**Very good identification**							**05100122**
					COC	GRA		
Prevotella bivia		99.97	0.66	BGL 1	1	1		
Gemella morbillorum		0.01	0.14		99	99		
Atopobium minutum		0.01	0.11		1	99		
05100130	**Very good identification**							**05100130**
					COC	GRA	SPO	
Prevotella bivia		99.97	0.66	MLZ 1	1	1	1	
Gemella morbillorum		0.01	0.14		99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		0.01	0.21		1	99	99	
05100160	**Very good identification**							**05100160**
					GRA	SPO		
Prevotella bivia		99.82	0.66	SOR 1	1	1		
Clostridium botulinum biovar B		0.15	0.26		99	99		
Clostridium novyi biovar A		0.01	0.18		99	99		
05100200	**Low discrimination**							**05100200**
					COC	GRA		
Prevotella disiens		75.64	0.66	RHA 1	1	1		
Gemella morbillorum		19.36	0.64		99	99		
Peptostreptococcus anaerobius		1.81	0.39		99	99		
05100220	**Low discrimination**							**05100220**
					COC	GRA		
Gemella morbillorum		45.82	0.48	RHA 1	99	99		
Prevotella bivia		33.44	0.32	NAG 10	1	1		
Prevotella disiens		16.28	0.32		1	1		
05100320	**Very good identification**							**05100320**
					COC	GRA		
Prevotella bivia		99.98	0.66	RHA 1	1	1		
Gemella morbillorum		0.01	0.14		99	99		
Prevotella disiens		0.01	-.03		1	1		
05100400	**Low discrimination**							**05100400**
					COC	GRA	SPO	
Gemella morbillorum		72.53	0.82	ARA 10	99	99	1	
Prevotella disiens		25.76	0.66		1	1	1	
Clostridium bifermentans		0.61	0.43		1	99	99	
05100401	**Low discrimination**							**05100401**
					COC	GRA	SPO	
Gemella morbillorum		70.04	0.48	ARA 10	99	99	1	
Prevotella disiens		24.88	0.32	URE 1	1	1	1	
Clostridium novyi biovar A		2.34	0.32		1	99	99	
05100402	**Low discrimination**							**05100402**
					COC	GRA	SPO	
Gemella morbillorum		71.05	0.48	ARA 10	99	99	1	
Prevotella disiens		25.24	0.32	BGL 1	1	1	1	
Clostridium sporogenes		0.93	0.11		1	99	99	
05100410	**Low discrimination**							**05100410**
					COC	GRA	SPO	
Gemella morbillorum		70.88	0.48	ARA 10	99	99	1	
Prevotella disiens		25.18	0.32	MLZ 1	1	1	1	
Clostridium novyi biovar A		2.37	0.32		1	99	99	
05100420	**Good identification**							**05100420**
					COC	GRA		
Gemella morbillorum		90.29	0.66	ARA 10	99	99		
Prevotella bivia		5.99	0.32	NAG 10	1	1		
Prevotella disiens		2.92	0.32		1	1		
05100440	**Low discrimination**							**05100440**
					COC	GRA	SPO	
Gemella morbillorum		55.52	0.48	ARA 10	99	99	1	
Prevotella disiens		19.72	0.32	SOR 1	1	1	1	
Clostridium botulinum biovar B		12.13	0.31		1	99	99	
05100500	**Low discrimination**							**05100500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Gemella morbillorum		47.38	0.48	LAC 1	99	99		
Prevotella bivia		34.58	0.32	ARA 10	1	1		
Prevotella disiens		16.83	0.32		1	1		
05100520	**Very good identification**							**05100520**
					COC	GRA		
Prevotella bivia		99.84	0.66	ARA 1	1	1		
Gemella morbillorum		0.15	0.32		99	99		
Prevotella disiens		0.01	-.03		1	1		
05100600	**Low discrimination**							**05100600**
					COC	GRA		
Gemella morbillorum		72.60	0.48	RHA 1	99	99		
Prevotella disiens		25.79	0.32	ARA 10	1	1		
Peptostreptococcus anaerobius		0.62	0.05		99	99		
05101000	**Good identification**							**05101000**
					COC	GRA		
Gemella morbillorum		92.24	0.94		99	99		
Prevotella disiens		6.76	0.66		1	1		
Prevotella intermedia		0.35	0.52		1	1		
05101001	**Low discrimination**							**05101001**
					COC	GRA	SPO	
Gemella morbillorum		82.48	0.60	URE 1	99	99	1	
Clostridium novyi biovar A		6.25	0.50		1	99	99	
Prevotella disiens		6.05	0.32		1	1	1	
05101002	**Low discrimination**							**05101002**
					COC	GRA		
Gemella morbillorum		84.54	0.60	BGL 1	99	99		
Atopobium minutum		6.29	0.45		1	99		
Prevotella disiens		6.20	0.32		1	1		
05101003	**Low discrimination**							**05101003**
					COC	SPO		
Clostridium novyi biovar A		50.75	0.37	MNS 1	1	99		
Gemella morbillorum		38.32	0.26	GAL 10	99	1		
Clostridium chauvoei		3.44	0.07		1	99		
05101010	**Low discrimination**							**05101010**
					COC	GRA	SPO	
Gemella morbillorum		86.23	0.60	MLZ 1	99	99	1	
Clostridium novyi biovar A		6.54	0.50		1	99	99	
Prevotella disiens		6.32	0.32		1	1	1	
05101011	**Low discrimination**							**05101011**
					COC	GRA	SPO	
Clostridium novyi biovar A		87.00	0.50	MNS 1	1	99	99	
Gemella morbillorum		11.59	0.26	GAL 10	99	99	1	
Prevotella disiens		0.85	-.03		1	1	1	
05101012	**Low discrimination**							**05101012**
					COC	SPO		
Clostridium novyi biovar A		53.57	0.37	MNS 1	1	99		
Gemella morbillorum		40.45	0.26	GAL 10	99	1		
Atopobium minutum		3.01	0.11		1	1		
05101013	**Good identification**							**05101013**
Clostridium novyi biovar A		100.0	0.37	MNS 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05101020	**Low discrimination**							**05101020**
					COC	GRA		
Gemella morbillorum		79.68	0.78	NAG 10	99	99		
Prevotella bivia		19.06	0.53		1	1		
Prevotella disiens		0.53	0.32		1	1		
05101021	**Low discrimination**							**05101021**
					COC	GRA	SPO	
Gemella morbillorum		69.47	0.43	NAG 10	99	99	1	
Prevotella bivia		16.62	0.19	URE 1	1	1	1	
Clostridium novyi biovar A		11.85	0.39		1	99	99	
05101022	**Low discrimination**							**05101022**
					COC	GRA		
Gemella morbillorum		62.35	0.43	NAG 10	99	99		
Prevotella bivia		14.91	0.19	BGL 1	1	1		
Atopobium minutum		13.92	0.37		1	99		
05101030	**Low discrimination**							**05101030**
					COC	GRA	SPO	
Gemella morbillorum		70.43	0.43	MLZ 1	99	99	1	
Prevotella bivia		16.85	0.19	NAG 10	1	1	1	
Clostridium novyi biovar A		12.01	0.39		1	99	99	
05101031	**Acceptable identification**							**05101031**
					COC	SPO		
Clostridium novyi biovar A		94.41	0.39	MNS 1	1	99		
Gemella morbillorum		5.59	0.09	GAL 10	99	1		
05101040	**Low discrimination**							**05101040**
					COC	GRA	SPO	
Gemella morbillorum		82.06	0.60	SOR 1	99	99	1	
Prevotella disiens		6.01	0.32		1	1	1	
Clostridium novyi biovar A		4.15	0.47		1	99	99	
05101041	**Low discrimination**							**05101041**
					COC	SPO		
Clostridium novyi biovar A		79.17	0.47	MNS 1	1	99		
Gemella morbillorum		15.82	0.26	GAL 10	99	1		
Clostridium sordellii		3.13	0.06		1	99		
05101050	**Low discrimination**							**05101050**
					COC	SPO		
Clostridium novyi biovar A		60.47	0.47	MNS 1	1	99		
Clostridium botulinum biovar B		26.57	0.26	GAL 10	1	99		
Gemella morbillorum		12.09	0.26		99	1		
05101051	**Good identification**							**05101051**
Clostridium novyi biovar A		100.0	0.47	MNS 1				
05101060	**Low discrimination**							**05101060**
					COC	GRA	SPO	
Gemella morbillorum		62.62	0.43	NAG 10	99	99	1	
Prevotella bivia		14.98	0.19	SOR 1	1	1	1	
Clostridium botulinum biovar B		12.51	0.26		1	99	99	
05101061	**Acceptable identification**							**05101061**
					COC	SPO		
Clostridium novyi biovar A		91.84	0.36	MNS 1	1	99		
Gemella morbillorum		8.16	0.09	GAL 10	99	1		
05101070	**Low discrimination**							**05101070**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Clostridium novyi biovar A		51.17	0.36	MNS 1	1	99		
Clostridium botulinum biovar B		44.29	0.21	GAL 10	1	99		
Gemella morbillorum		4.55	0.09		99	1		
05101071	**Good identification**							**05101071**
Clostridium novyi biovar A		100.0	0.36	MNS 1				
05101100	**Low discrimination**							**05101100**
					COC	GRA		
Prevotella bivia		69.15	0.53	NAG 99	1	1		
Gemella morbillorum		26.28	0.60		99	99		
Prevotella disiens		1.93	0.32		1	1		
05101101	**Acceptable identification**							**05101101**
					COC	GRA	SPO	
Clostridium chauvoei		92.82	0.54	SUC 99	1	99	99	
Prevotella bivia		4.86	0.19		1	1	1	
Gemella morbillorum		1.85	0.26		99	99	1	
05101103	**Good identification**							**05101103**
Clostridium chauvoei		100.0	0.41	SUC 99				
05101120	**Very good identification**							**05101120**
					COC	GRA		
Prevotella bivia		99.85	0.87		1	1		
Clostridium perfringens		0.05	0.44		1	99		
Gemella morbillorum		0.04	0.43		99	99		
05101121	**Acceptable identification**							**05101121**
					GRA	SPO		
Prevotella bivia		93.82	0.53	URE 1	1	1		
Clostridium chauvoei		6.04	0.45		99	99		
Clostridium perfringens		0.05	0.10		99	84		
05101122	**Good identification**							**05101122**
					GRA			
Prevotella bivia		96.92	0.53	BGL 1	1			
Clostridium perfringens		2.23	0.38		99			
Prevotella melaninogenica		0.53	0.21		1			
05101130	**Good identification**							**05101130**
					COC	GRA		
Prevotella bivia		99.85	0.53	MLZ 1	1	1		
Clostridium perfringens		0.05	0.10		1	99		
Gemella morbillorum		0.04	0.09		99	99		
05101160	**Good identification**							**05101160**
					COC	GRA		
Prevotella bivia		98.98	0.53	SOR 1	1	1		
Clostridium perfringens		0.92	0.32		1	99		
Gemella morbillorum		0.04	0.09		99	99		
05101200	**Good identification**							**05101200**
					COC	GRA		
Gemella morbillorum		92.29	0.60	RHA 1	99	99		
Prevotella disiens		6.76	0.32		1	1		
Prevotella intermedia		0.35	0.17		1	1		
05101220	**Low discrimination**							**05101220**
					COC	GRA		

Code / Taxon		ID score	T index	tests against		Differentiation tests			Code
Gemella morbillorum		80.05	0.43	RHA	1	99	99		
Prevotella bivia		19.15	0.19	NAG	10	1	1		
Prevotella disiens		0.53	-.03			1	1		
05101320	**Good identification**								**05101320**
						COC	GRA		
Prevotella bivia		99.85	0.53	RHA	1	1	1		
Clostridium perfringens		0.05	0.10			1	99		
Gemella morbillorum		0.04	0.09			99	99		
05101400	**Very good identification**								**05101400**
						COC	GRA		
Gemella morbillorum		99.17	0.78	ARA	10	99	99		
Prevotella disiens		0.66	0.32			1	1		
Lactobacillus fermentum		0.08	0.36			1	99		
05101401	**Good identification**								**05101401**
						COC	GRA	SPO	
Gemella morbillorum		98.24	0.43	ARA	10	99	99	1	
Clostridium novyi biovar A		0.68	0.15	URE	1	1	99	99	
Prevotella disiens		0.66	-.03			1	1	1	
05101402	**Acceptable identification**								**05101402**
						COC			
Gemella morbillorum		91.60	0.43	ARA	10	99			
Lactobacillus fermentum		7.17	0.36	BGL	1	1			
Atopobium minutum		0.62	0.11			1			
05101410	**Good identification**								**05101410**
						COC	GRA	SPO	
Gemella morbillorum		98.66	0.43	ARA	10	99	99	1	
Clostridium novyi biovar A		0.68	0.15	MLZ	1	1	99	99	
Prevotella disiens		0.66	-.03			1	1	1	
05101420	**Very good identification**								**05101420**
						COC	GRA		
Gemella morbillorum		97.56	0.61	ARA	10	99	99		
Prevotella bivia		2.12	0.19	NAG	10	1	1		
Lactobacillus fermentum		0.23	0.28			1	99		
05101440	**Good identification**								**05101440**
						COC	GRA	SPO	
Gemella morbillorum		98.49	0.43	ARA	10	99	99	1	
Prevotella disiens		0.66	-.03	SOR	1	1	1	1	
Clostridium novyi biovar A		0.45	0.12			1	99	99	
05101500	**Low discrimination**								**05101500**
						COC	GRA		
Gemella morbillorum		67.71	0.43	LAC	1	99	99		
Prevotella bivia		16.19	0.19	ARA	10	1	1		
Lactobacillus fermentum		8.64	0.40			1	99		
05101502	**Low discrimination**								**05101502**
						COC			
Lactobacillus fermentum		86.12	0.40	SUC	87	1			
Blautia hansenii		7.07	0.03	FRU	99	99			
Gemella morbillorum		6.81	0.09			99			
05101520	**Good identification**								**05101520**
						COC	GRA		
Prevotella bivia		99.16	0.53	ARA	1	1	1		
Gemella morbillorum		0.47	0.27			99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		0.18	0.32		1	99		
05101600	**Good identification**							**05101600**
					COC	GRA		
Gemella morbillorum		99.34	0.43	RHA 1	99	99		
Prevotella disiens		0.66	-.03	ARA 10	1	1		
05102000	**Low discrimination**							**05102000**
					COC	GRA		
Prevotella disiens		74.74	0.66	MAN 1	1	1		
Gemella morbillorum		19.13	0.64		99	99		
Peptostreptococcus anaerobius		1.78	0.39		99	99		
05102020	**Low discrimination**							**05102020**
					COC	GRA		
Gemella morbillorum		45.56	0.48	MAN 1	99	99		
Prevotella bivia		33.25	0.32	NAG 10	1	1		
Prevotella disiens		16.18	0.32		1	1		
05102120	**Very good identification**							**05102120**
					COC	GRA		
Prevotella bivia		99.98	0.66	MAN 1	1	1		
Gemella morbillorum		0.01	0.14		99	99		
Prevotella disiens		0.01	-.03		1	1		
05102400	**Low discrimination**							**05102400**
					COC	GRA	SPO	
Gemella morbillorum		72.32	0.48	MAN 1	99	99	1	
Prevotella disiens		25.69	0.32	ARA 10	1	1	1	
Clostridium bifermentans		0.61	0.09		1	99	99	
05103000	**Good identification**							**05103000**
					COC	GRA	CAT	
Gemella morbillorum		91.70	0.60	MAN 1	99	99	1	
Prevotella disiens		6.72	0.32		1	1	1	
Propionibacterium granulosum		0.54	0.29		1	99	90	
05103020	**Low discrimination**							**05103020**
					COC	GRA	CAT	
Gemella morbillorum		79.21	0.43	MAN 1	99	99	1	
Prevotella bivia		18.94	0.19	NAG 10	1	1	1	
Propionibacterium granulosum		1.06	0.18		1	99	90	
05103120	**Good identification**							**05103120**
					COC	GRA		
Prevotella bivia		99.85	0.53	MAN 1	1	1		
Clostridium perfringens		0.05	0.10		1	99		
Gemella morbillorum		0.04	0.09		99	99		
05103400	**Good identification**							**05103400**
					COC	GRA		
Gemella morbillorum		99.34	0.43	MAN 1	99	99		
Prevotella disiens		0.66	-.03	ARA 10	1	1		
05104000	**Low discrimination**							**05104000**
					COC	GRA	SPO	
Gemella morbillorum		72.47	0.82	XYL 10	99	99	1	
Prevotella disiens		25.74	0.66		1	1	1	
Clostridium bifermentans		0.61	0.43		1	99	99	
05104001	**Low discrimination**							**05104001**
					COC	GRA	SPO	
Gemella morbillorum		70.04	0.48	XYL 10	99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella disiens		24.88	0.32	URE 1	1	1	1	
Clostridium novyi biovar A		2.34	0.32		1	99	99	
05104002	**Low discrimination**							**05104002**
					COC	GRA	SPO	
Gemella morbillorum		71.05	0.48	XYL 10	99	99	1	
Prevotella disiens		25.24	0.32	BGL 1	1	1	1	
Clostridium sporogenes		0.93	0.11		1	99	99	
05104010	**Low discrimination**							**05104010**
					COC	GRA	SPO	
Gemella morbillorum		70.88	0.48	XYL 10	99	99	1	
Prevotella disiens		25.18	0.32	MLZ 1	1	1	1	
Clostridium novyi biovar A		2.37	0.32		1	99	99	
05104020	**Good identification**							**05104020**
					COC	GRA		
Gemella morbillorum		90.29	0.66	XYL 10	99	99		
Prevotella bivia		5.99	0.32	NAG 10	1	1		
Prevotella disiens		2.92	0.32		1	1		
05104040	**Low discrimination**							**05104040**
					COC	GRA	SPO	
Clostridium glycolicum		53.55	0.50	MNS 1	1	99	99	
Gemella morbillorum		25.79	0.48	FRU 90	99	99	1	
Prevotella disiens		9.16	0.32		1	1	1	
05104100	**Low discrimination**							**05104100**
					COC	GRA		
Gemella morbillorum		46.89	0.48	XYL 10	99	99		
Prevotella bivia		34.22	0.32	LAC 1	1	1		
Prevotella disiens		16.65	0.32		1	1		
05104120	**Very good identification**							**05104120**
					COC	GRA		
Prevotella bivia		99.84	0.66	XYL 1	1	1		
Gemella morbillorum		0.15	0.32		99	99		
Actinomyces meyeri		0.01	0.07		1	99		
05104200	**Low discrimination**							**05104200**
					COC	GRA		
Gemella morbillorum		72.60	0.48	XYL 10	99	99		
Prevotella disiens		25.79	0.32	RHA 1	1	1		
Peptostreptococcus anaerobius		0.62	0.05		99	99		
05104400	**Very good identification**							**05104400**
					COC	GRA	SPO	
Gemella morbillorum		96.68	0.66	XYL 10	99	99	1	
Prevotella disiens		3.12	0.32	ARA 10	1	1	1	
Clostridium bifermentans		0.07	0.09		1	99	99	
05104420	**Good identification**							**05104420**
					COC	GRA		
Gemella morbillorum		99.11	0.50	XYL 10	99	99		
Prevotella bivia		0.60	-.03	ARA 10	1	1		
Prevotella disiens		0.29	-.03	NAG 10	1	1		
05105000	**Very good identification**							**05105000**
					COC	GRA		
Gemella morbillorum		99.16	0.78	XYL 10	99	99		
Prevotella disiens		0.66	0.32		1	1		
Lactobacillus fermentum		0.05	0.32		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05105001	**Good identification**							**05105001**
					COC	GRA	SPO	
Gemella morbillorum		98.24	0.43	XYL 10	99	99	1	
Clostridium novyi biovar A		0.68	0.15	URE 1	1	99	99	
Prevotella disiens		0.66	-.03		1	1	1	
05105002	**Acceptable identification**							**05105002**
					COC			
Gemella morbillorum		94.40	0.43	XYL 10	99			
Lactobacillus fermentum		4.34	0.32	BGL 1	1			
Atopobium minutum		0.64	0.11		1			
05105010	**Good identification**							**05105010**
					COC	GRA	SPO	
Gemella morbillorum		98.66	0.43	XYL 10	99	99	1	
Clostridium novyi biovar A		0.68	0.15	MLZ 1	1	99	99	
Prevotella disiens		0.66	-.03		1	1	1	
05105020	**Very good identification**							**05105020**
					COC	GRA		
Gemella morbillorum		97.65	0.61	XYL 10	99	99		
Prevotella bivia		2.12	0.19	NAG 10	1	1		
Lactobacillus fermentum		0.14	0.24		1	99		
05105040	**Low discrimination**							**05105040**
					COC	GRA	SPO	
Gemella morbillorum		69.26	0.43	XYL 10	99	99	1	
Clostridium glycolicum		29.68	0.33	SOR 1	1	99	99	
Prevotella disiens		0.46	-.03		1	1	1	
05105100	**Low discrimination**							**05105100**
					COC	GRA		
Gemella morbillorum		67.29	0.43	XYL 10	99	99		
Prevotella bivia		16.10	0.19	LAC 1	1	1		
Blautia hansenii		6.35	0.19		99	99		
05105102	**Low discrimination**							**05105102**
					COC			
Lactobacillus fermentum		78.47	0.36	SUC 87	1			
Blautia hansenii		10.96	0.03	FRU 99	99			
Gemella morbillorum		10.57	0.09		99			
05105120	**Good identification**							**05105120**
					COC	GRA		
Prevotella bivia		99.24	0.53	XYL 1	1	1		
Gemella morbillorum		0.47	0.27		99	99		
Lactobacillus fermentum		0.11	0.28		1	99		
05105200	**Good identification**							**05105200**
					COC	GRA		
Gemella morbillorum		99.34	0.43	XYL 10	99	99		
Prevotella disiens		0.66	-.03	RHA 1	1	1		
05105400	**Very good identification**							**05105400**
					COC	GRA		
Gemella morbillorum		99.48	0.61	XYL 10	99	99		
Lactobacillus fermentum		0.42	0.32	ARA 10	1	99		
Prevotella disiens		0.06	-.03		1	1		
05105420	**Good identification**							**05105420**
					COC			
Gemella morbillorum		98.76	0.45	XYL 10	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		1.24	0.24	ARA 10	1			
05105502	**Acceptable identification**							**05105502**
Lactobacillus fermentum		93.22	0.36	SUC 87				
Bifidobacterium longum subsp. longum		6.78	0.09	FRU 99				
05105722	**Good identification**							**05105722**
Bacteroides eggerthii		100.0	0.39	IND 99				
05106000	**Low discrimination**							**05106000**
					COC	GRA	SPO	
Gemella morbillorum		72.32	0.48	MAN 1	99	99	1	
Prevotella disiens		25.69	0.32	XYL 10	1	1	1	
Clostridium bifermentans		0.61	0.09		1	99	99	
05107000	**Good identification**							**05107000**
					COC	GRA		
Gemella morbillorum		99.34	0.43	MAN 1	99	99		
Prevotella disiens		0.66	-.03	XYL 10	1	1		
05110000	**Low discrimination**							**05110000**
					COC	GRA		
Prevotella disiens		72.68	1.00		1	1		
Gemella morbillorum		22.74	1.00		99	99		
Prevotella intermedia		3.75	0.86		1	1		
05110001	**Low discrimination**							**05110001**
					COC	GRA		
Prevotella disiens		70.24	0.66	URE 1	1	1		
Gemella morbillorum		21.97	0.66		99	99		
Prevotella intermedia		3.62	0.52		1	1		
05110002	**Low discrimination**							**05110002**
					COC	GRA		
Prevotella disiens		68.57	0.66	BGL 1	1	1		
Gemella morbillorum		21.45	0.66		99	99		
Prevotella intermedia		3.54	0.52		1	1		
05110010	**Low discrimination**							**05110010**
					COC	GRA		
Prevotella disiens		69.63	0.66	MLZ 1	1	1		
Gemella morbillorum		21.78	0.66		99	99		
Prevotella intermedia		3.59	0.52		1	1		
05110011	**Low discrimination**							**05110011**
					COC	GRA	SPO	
Clostridium novyi biovar A		41.58	0.50	MNS 1	1	99	99	
Prevotella disiens		40.20	0.32	FRU 10	1	1	1	
Gemella morbillorum		12.58	0.32		99	99	1	
05110013	**Acceptable identification**							**05110013**
					GRA	SPO		
Clostridium novyi biovar A		94.76	0.37	MNS 1	99	99		
Prevotella disiens		5.24	-.03	FRU 10	1	1		
05110020	**Low discrimination**							**05110020**
					COC	GRA		
Gemella morbillorum		70.94	0.84	NAG 10	99	99		
Prevotella disiens		20.62	0.66		1	1		
Prevotella bivia		4.71	0.50		1	1		
05110021	**Low discrimination**							**05110021**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Gemella morbillorum		67.07	0.50	NAG 10	99	99	1	
Prevotella disiens		19.49	0.32	URE 1	1	1	1	
Clostridium novyi biovar A		5.04	0.39		1	99	99	
05110022	**Low discrimination**							**05110022**
					COC	GRA		
Gemella morbillorum		51.84	0.50	NAG 10	99	99		
Prevotella disiens		15.07	0.32	BGL 1	1	1		
Atopobium minutum		14.12	0.45		1	99		
05110030	**Low discrimination**							**05110030**
					COC	GRA	SPO	
Gemella morbillorum		57.00	0.50	MLZ 1	99	99	1	
Prevotella disiens		16.57	0.32	NAG 10	1	1	1	
Clostridium botulinum biovar B		9.89	0.31		1	99	99	
05110031	**Low discrimination**							**05110031**
					COC	GRA	SPO	
Clostridium novyi biovar A		83.56	0.39	MNS 1	1	99	99	
Gemella morbillorum		11.23	0.15	FRU 10	99	99	1	
Prevotella disiens		3.26	-.03		1	1	1	
05110040	**Low discrimination**							**05110040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		61.23	0.76	SUC 90	1	99	99	
Prevotella disiens		24.89	0.66		1	1	1	
Gemella morbillorum		7.79	0.66		99	99	1	
05110041	**Low discrimination**							**05110041**
					GRA	SPO		
Clostridium botulinum biovar B		50.78	0.41	SUC 90	99	99		
Prevotella disiens		20.64	0.32	URE 1	1	1		
Clostridium novyi biovar A		14.23	0.47		99	99		
05110042	**Low discrimination**							**05110042**
					COC	GRA	SPO	
Clostridium botulinum biovar B		45.88	0.41	SUC 90	1	99	99	
Megasphaera elsdenii		23.54	0.44	BGL 1	99	1	1	
Prevotella disiens		18.65	0.32		1	1	1	
05110050	**Very good identification**							**05110050**
					GRA	SPO		
Clostridium botulinum biovar B		98.06	0.70	SUC 90	99	99		
Prevotella disiens		0.82	0.32		1	1		
Clostridium novyi biovar A		0.56	0.47		99	99		
05110051	**Low discrimination**							**05110051**
					GRA	SPO		
Clostridium botulinum biovar B		63.40	0.36	SUC 90	99	99		
Clostridium novyi biovar A		36.08	0.47	URE 1	99	99		
Prevotella disiens		0.53	-.03		1	1		
05110052	**Low discrimination**							**05110052**
					COC	GRA		
Clostridium botulinum biovar B		89.34	0.36	SUC 90	1	99		
Clostridium novyi biovar A		8.97	0.34	BGL 1	1	99		
Megasphaera elsdenii		0.94	0.09		99	1		
05110060	**Very good identification**							**05110060**
					COC	GRA	SPO	
Clostridium botulinum biovar B		95.69	0.70	SUC 90	1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		2.75	0.50		99	99	1	
Prevotella disiens		0.80	0.32		1	1	1	
05110061	**Low discrimination**							**05110061**
					COC	SPO		
Clostridium botulinum biovar B		84.80	0.36	SUC 90	1	99		
Clostridium novyi biovar A		12.06	0.36	URE 1	1	99		
Gemella morbillorum		2.43	0.15		99	1		
05110062	**Acceptable identification**							**05110062**
					COC	SPO		
Clostridium botulinum biovar B		90.25	0.36	SUC 90	1	99		
Gemella morbillorum		2.59	0.15	BGL 1	99	1		
Clostridium novyi biovar A		2.27	0.23		1	99		
05110070	**Very good identification**							**05110070**
					COC	SPO		
Clostridium botulinum biovar B		99.63	0.65	SUC 90	1	99		
Clostridium novyi biovar A		0.29	0.36		1	99		
Gemella morbillorum		0.06	0.15		99	1		
05110100	**Low discrimination**							**05110100**
					COC	GRA		
Prevotella disiens		61.21	0.66	LAC 1	1	1		
Gemella morbillorum		19.15	0.66		99	99		
Prevotella bivia		13.97	0.50		1	1		
05110120	**Very good identification**							**05110120**
					COC	GRA		
Prevotella bivia		99.66	0.84	FRU 10	1	1		
Gemella morbillorum		0.15	0.50		99	99		
Clostridium perfringens		0.08	0.44		1	99		
05110121	**Good identification**							**05110121**
					COC	GRA	SPO	
Prevotella bivia		99.44	0.50	FRU 10	1	1	1	
Clostridium chauvoei		0.24	0.18	URE 1	1	99	99	
Gemella morbillorum		0.15	0.15		99	99	1	
05110122	**Good identification**							**05110122**
					GRA			
Prevotella bivia		95.77	0.50	FRU 10	1			
Clostridium perfringens		3.50	0.38	BGL 1	99			
Lactobacillus fermentum		0.27	0.32		99			
05110130	**Good identification**							**05110130**
					COC	GRA		
Prevotella bivia		99.66	0.50	FRU 10	1	1		
Gemella morbillorum		0.15	0.15	MLZ 1	99	99		
Clostridium perfringens		0.08	0.10		1	99		
05110140	**Low discrimination**							**05110140**
					COC	GRA	SPO	
Clostridium botulinum biovar B		57.50	0.41	SUC 90	1	99	99	
Prevotella disiens		23.37	0.32	LAC 1	1	1	1	
Gemella morbillorum		7.31	0.32		99	99	1	
05110150	**Good identification**							**05110150**
					GRA	SPO		
Clostridium botulinum biovar B		98.61	0.36	SUC 90	99	99		
Prevotella disiens		0.82	-.03	LAC 1	1	1		
Clostridium novyi biovar A		0.57	0.12		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05110160	**Acceptable identification**							**05110160**
					GRA	SPO		
Prevotella bivia		93.39	0.50	FRU 10	1	1		
Clostridium botulinum biovar B		5.01	0.36	SOR 1	99	99		
Clostridium perfringens		1.38	0.32		99	84		
05110200	**Low discrimination**							**05110200**
					COC	GRA		
Prevotella disiens		72.56	0.66	RHA 1	1	1		
Gemella morbillorum		22.70	0.66		99	99		
Prevotella intermedia		3.74	0.52		1	1		
05110220	**Low discrimination**							**05110220**
					COC	GRA		
Gemella morbillorum		70.37	0.50	RHA 1	99	99		
Prevotella disiens		20.45	0.32	NAG 10	1	1		
Prevotella bivia		4.67	0.15		1	1		
05110240	**Low discrimination**							**05110240**
					COC	GRA	SPO	
Clostridium botulinum biovar B		61.38	0.41	SUC 90	1	99	99	
Prevotella disiens		24.95	0.32	RHA 1	1	1	1	
Gemella morbillorum		7.80	0.32		99	99	1	
05110250	**Good identification**							**05110250**
					GRA	SPO		
Clostridium botulinum biovar B		98.61	0.36	SUC 90	99	99		
Prevotella disiens		0.82	-.03	RHA 1	1	1		
Clostridium novyi biovar A		0.57	0.12		99	99		
05110260	**Good identification**							**05110260**
					COC	GRA	SPO	
Clostridium botulinum biovar B		96.43	0.36	SUC 90	1	99	99	
Gemella morbillorum		2.77	0.15	RHA 1	99	99	1	
Prevotella disiens		0.80	-.03		1	1	1	
05110320	**Good identification**							**05110320**
					COC	GRA		
Prevotella bivia		99.68	0.50	FRU 10	1	1		
Gemella morbillorum		0.15	0.15	RHA 1	99	99		
Clostridium perfringens		0.08	0.10		1	99		
05110400	**Low discrimination**							**05110400**
					COC	GRA		
Gemella morbillorum		76.37	0.84	ARA 10	99	99		
Prevotella disiens		22.19	0.66		1	1		
Prevotella intermedia		1.15	0.52		1	1		
05110401	**Low discrimination**							**05110401**
					COC	GRA		
Gemella morbillorum		75.79	0.50	ARA 10	99	99		
Prevotella disiens		22.02	0.32	URE 1	1	1		
Prevotella intermedia		1.14	0.17		1	1		
05110402	**Low discrimination**							**05110402**
					COC	GRA		
Gemella morbillorum		73.34	0.50	ARA 10	99	99		
Prevotella disiens		21.31	0.32	BGL 1	1	1		
Lactobacillus fermentum		2.53	0.36		1	99		
05110410	**Low discrimination**							**05110410**
					COC	GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		75.55	0.50	ARA 10	99	99		
Prevotella disiens		21.96	0.32	MLZ 1	1	1		
Prevotella intermedia		1.13	0.17		1	1		
05110420	**Very good identification**							**05110420**
					COC	GRA		
Gemella morbillorum		96.34	0.67	ARA 10	99	99		
Prevotella disiens		2.55	0.32	NAG 10	1	1		
Prevotella bivia		0.58	0.15		1	1		
05110440	**Low discrimination**							**05110440**
					COC	GRA	SPO	
Gemella morbillorum		48.22	0.50	ARA 10	99	99	1	
Clostridium botulinum biovar B		34.48	0.41	SOR 1	1	99	99	
Prevotella disiens		14.01	0.32		1	1	1	
05110450	**Good identification**							**05110450**
					COC	GRA	SPO	
Clostridium botulinum biovar B		95.90	0.36	SUC 90	1	99	99	
Gemella morbillorum		2.75	0.15	ARA 1	99	99	1	
Prevotella disiens		0.80	-.03		1	1	1	
05110460	**Low discrimination**							**05110460**
					COC	GRA	SPO	
Clostridium botulinum biovar B		75.54	0.36	SUC 90	1	99	99	
Gemella morbillorum		23.83	0.33	ARA 1	99	99	1	
Prevotella disiens		0.63	-.03		1	1	1	
05110500	**Low discrimination**							**05110500**
					COC	GRA		
Gemella morbillorum		67.34	0.50	LAC 1	99	99		
Prevotella disiens		19.57	0.32	ARA 10	1	1		
Prevotella bivia		4.47	0.15		1	1		
05110502	**Low discrimination**							**05110502**
					COC	GRA		
Lactobacillus fermentum		79.38	0.40	SUC 87	1	99		
Gemella morbillorum		14.26	0.15	GAL 99	99	99		
Prevotella disiens		4.14	-.03		1	1		
05110520	**Good identification**							**05110520**
					COC	GRA		
Prevotella bivia		97.80	0.50	FRU 10	1	1		
Gemella morbillorum		1.65	0.33	ARA 1	99	99		
Lactobacillus fermentum		0.28	0.32		1	99		
05110600	**Low discrimination**							**05110600**
					COC	GRA		
Gemella morbillorum		76.59	0.50	RHA 1	99	99		
Prevotella disiens		22.26	0.32	ARA 10	1	1		
Prevotella intermedia		1.15	0.17		1	1		
05111000	**Low discrimination**							**05111000**
					COC	GRA		
Gemella morbillorum		72.31	0.95		99	99		
Prevotella intermedia		22.15	0.86		1	1		
Prevotella disiens		4.34	0.66		1	1		
05111001	**Low discrimination**							**05111001**
					COC	GRA		
Gemella morbillorum		69.55	0.61	URE 1	99	99		
Prevotella intermedia		21.30	0.52		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella disiens		4.17	0.32		1	1		
05111002	**Low discrimination**							**05111002**
					COC	GRA		
Gemella morbillorum		42.91	0.61	BGL 1	99	99		
Lactobacillus fermentum		30.21	0.70		1	99		
Prevotella intermedia		13.14	0.52		1	1		
05111010	**Low discrimination**							**05111010**
					COC	GRA	CAT	
Gemella morbillorum		67.69	0.61	MLZ 1	99	99	1	
Prevotella intermedia		20.73	0.52		1	1	1	
Propionibacterium granulosum		5.87	0.50		1	99	90	
05111012	**Low discrimination**							**05111012**
					COC	CAT		
Propionibacterium granulosum		44.79	0.40	SUC 93	1	90		
Gemella morbillorum		20.87	0.27	MLZ 12	99	1		
Lactobacillus fermentum		14.70	0.36		1	1		
05111020	**Low discrimination**							**05111020**
					COC	GRA		
Gemella morbillorum		81.16	0.79	NAG 10	99	99		
Clostridium perfringens		10.47	0.63		1	99		
Prevotella intermedia		2.26	0.52		1	1		
05111021	**Low discrimination**							**05111021**
					COC	SPO		
Gemella morbillorum		78.19	0.45	NAG 10	99	1		
Clostridium perfringens		10.09	0.29	URE 1	1	84		
Clostridium chauvoei		2.30	0.18		1	99		
05111022	**Low discrimination**							**05111022**
					COC			
Clostridium perfringens		61.37	0.58	SUC 91	1			
Lactobacillus fermentum		21.57	0.62	LAC 88	1			
Gemella morbillorum		10.21	0.45		99			
05111030	**Low discrimination**							**05111030**
					COC	CAT		
Gemella morbillorum		69.28	0.45	MLZ 1	99	1		
Propionibacterium granulosum		13.52	0.40	NAG 10	1	90		
Clostridium perfringens		8.94	0.29		1	1		
05111040	**Low discrimination**							**05111040**
					COC	GRA	SPO	
Gemella morbillorum		63.72	0.61	SOR 1	99	99	1	
Prevotella intermedia		19.52	0.52		1	1	1	
Clostridium botulinum biovar B		9.40	0.41		1	99	99	
05111050	**Low discrimination**							**05111050**
					COC	SPO		
Clostridium botulinum biovar B		79.04	0.36	SUC 90	1	99		
Gemella morbillorum		10.99	0.27	GAL 1	99	1		
Clostridium novyi biovar A		5.00	0.30		1	99		
05111060	**Low discrimination**							**05111060**
					COC	SPO		
Clostridium perfringens		57.10	0.51	SUC 91	1	84		
Gemella morbillorum		23.47	0.45	LAC 88	99	1		
Clostridium botulinum biovar B		15.35	0.36		1	99		
05111062	**Good identification**							**05111062**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Clostridium perfringens		96.49	0.46	SUC 91	1		
Lactobacillus fermentum		1.80	0.28	LAC 88	1		
Gemella morbillorum		0.85	0.11		99		
05111100	**Low discrimination**						**05111100**
					COC		
Lactobacillus fermentum		33.76	0.74	SUC 87	1		
Gemella morbillorum		29.39	0.61		99		
Clostridium perfringens		9.46	0.52		1		
05111101	**Very good identification**						**05111101**
					COC	SPO	
Clostridium chauvoei		96.52	0.60	SUC 99	1	99	
Lactobacillus fermentum		1.16	0.40		1	1	
Gemella morbillorum		1.01	0.27		99	1	
05111102	**Low discrimination**						**05111102**
					GRA		
Lactobacillus fermentum		85.06	0.74	SUC 87	99		
Clostridium perfringens		11.22	0.47		99		
Fusobacterium mortiferum		1.47	0.42		1		
05111103	**Acceptable identification**						**05111103**
					SPO		
Clostridium chauvoei		92.46	0.47	SUC 99	99		
Lactobacillus fermentum		6.27	0.40		1		
Clostridium perfringens		0.83	0.13		84		
05111110	**Low discrimination**						**05111110**
					COC		
Lactobacillus fermentum		33.21	0.40	SUC 87	1		
Gemella morbillorum		28.91	0.27	MLZ 1	99		
Clostridium perfringens		9.31	0.18		1		
05111112	**Low discrimination**						**05111112**
Lactobacillus fermentum		88.35	0.40	SUC 87			
Clostridium perfringens		11.65	0.13	MLZ 1			
05111120	**Low discrimination**						**05111120**
					GRA	SPO	
Prevotella bivia		58.31	0.71	FRU 10	1	1	
Clostridium perfringens		25.63	0.78		99	84	
Clostridium septicum		13.42	0.67		99	99	
05111121	**Low discrimination**						**05111121**
					GRA	SPO	
Clostridium chauvoei		44.06	0.52	SUC 99	99	99	
Prevotella bivia		32.60	0.37		1	1	
Clostridium perfringens		14.33	0.44		99	84	
05111122	**Low discrimination**						**05111122**
					GRA		
Clostridium perfringens		84.34	0.73	SUC 91	99		
Lactobacillus fermentum		6.59	0.66		99		
Prevotella bivia		4.12	0.37		1		
05111123	**Low discrimination**						**05111123**
					SPO		
Clostridium chauvoei		49.11	0.39	SUC 99	99		
Clostridium perfringens		42.61	0.38		84		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		3.33	0.32		1		
05111130	**Low discrimination**						**05111130**
					GRA	SPO	
Prevotella bivia		58.34	0.37	FRU 10	1	1	
Clostridium perfringens		25.65	0.44	MLZ 1	99	84	
Clostridium septicum		13.43	0.33		99	99	
05111132	**Low discrimination**						**05111132**
					GRA		
Clostridium perfringens		84.42	0.38	SUC 91	99		
Lactobacillus fermentum		6.60	0.32	MLZ 1	99		
Prevotella bivia		4.12	0.03		1		
05111140	**Low discrimination**						**05111140**
					COC		
Clostridium perfringens		65.30	0.40	SUC 91	1		
Lactobacillus fermentum		12.36	0.40	NAG 97	1		
Gemella morbillorum		10.76	0.27		99		
05111160	**Low discrimination**						**05111160**
					GRA	SPO	
Clostridium perfringens		86.65	0.66	SUC 91	99	84	
Prevotella bivia		10.45	0.37		1	1	
Clostridium septicum		2.41	0.33		99	99	
05111162	**Very good identification**						**05111162**
					GRA		
Clostridium perfringens		99.03	0.60	SUC 91	99		
Lactobacillus fermentum		0.41	0.32		99		
Prevotella bivia		0.26	0.03		1		
05111200	**Low discrimination**						**05111200**
					COC	GRA	
Gemella morbillorum		71.66	0.61	RHA 1	99	99	
Prevotella intermedia		21.95	0.52		1	1	
Prevotella disiens		4.30	0.32		1	1	
05111220	**Low discrimination**						**05111220**
					COC	CAT	
Gemella morbillorum		79.35	0.45	RHA 1	99	1	
Clostridium perfringens		10.24	0.29	NAG 10	1	1	
Propionibacterium granulosum		3.51	0.29		1	90	
05111300	**Low discrimination**						**05111300**
					COC		
Lactobacillus fermentum		34.06	0.40	SUC 87	1		
Gemella morbillorum		29.65	0.27	RHA 1	99		
Clostridium perfringens		9.55	0.18		1		
05111302	**Low discrimination**						**05111302**
					GRA		
Lactobacillus fermentum		79.45	0.40	SUC 87	99		
Clostridium perfringens		10.48	0.13	RHA 1	99		
Prevotella oralis		10.08	0.09		1		
05111320	**Low discrimination**						**05111320**
					GRA	SPO	
Prevotella bivia		58.05	0.37	FRU 10	1	1	
Clostridium perfringens		25.52	0.44	RHA 1	99	84	
Clostridium septicum		13.36	0.33		99	99	
05111322	**Low discrimination**						**05111322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Clostridium perfringens		56.69	0.38	SUC 91	99		
Prevotella oralis		32.05	0.31	RHA 1	1		
Lactobacillus fermentum		4.43	0.32		99		
05111400	**Good identification**						**05111400**
					COC	GRA	
Gemella morbillorum		91.04	0.79	ARA 10	99	99	
Lactobacillus fermentum		5.83	0.70		1	99	
Prevotella intermedia		2.54	0.52		1	1	
05111401	**Acceptable identification**						**05111401**
					COC	GRA	
Gemella morbillorum		90.84	0.45	ARA 10	99	99	
Lactobacillus fermentum		5.82	0.36	URE 1	1	99	
Prevotella intermedia		2.53	0.17		1	1	
05111402	**Low discrimination**						**05111402**
					COC	GRA	
Lactobacillus fermentum		85.75	0.70	SUC 87	1	99	
Gemella morbillorum		13.53	0.45		99	99	
Prevotella intermedia		0.38	0.17		1	1	
05111403	**Low discrimination**						**05111403**
					COC		
Lactobacillus fermentum		86.37	0.36	SUC 87	1		
Gemella morbillorum		13.63	0.11	URE 1	99		
05111410	**Acceptable identification**						**05111410**
					COC	GRA	
Gemella morbillorum		90.48	0.45	ARA 10	99	99	
Lactobacillus fermentum		5.79	0.36	MLZ 1	1	99	
Prevotella intermedia		2.52	0.17		1	1	
05111412	**Low discrimination**						**05111412**
					COC		
Lactobacillus fermentum		86.37	0.36	SUC 87	1		
Gemella morbillorum		13.63	0.11	MLZ 1	99		
05111420	**Low discrimination**						**05111420**
					COC		
Gemella morbillorum		82.54	0.63	ARA 10	99		
Lactobacillus fermentum		15.85	0.62	NAG 10	1		
Clostridium perfringens		0.97	0.29		1		
05111422	**Good identification**						**05111422**
					COC		
Lactobacillus fermentum		92.33	0.62	SUC 87	1		
Gemella morbillorum		4.86	0.29		99		
Clostridium perfringens		2.65	0.24		1		
05111440	**Acceptable identification**						**05111440**
					COC	GRA	
Gemella morbillorum		90.03	0.45	ARA 10	99	99	
Lactobacillus fermentum		5.76	0.36	SOR 1	1	99	
Prevotella intermedia		2.51	0.17		1	1	
05111442	**Low discrimination**						**05111442**
					COC		
Lactobacillus fermentum		86.37	0.36	SUC 87	1		
Gemella morbillorum		13.63	0.11	SOR 1	99		
05111500	**Low discrimination**						**05111500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Lactobacillus fermentum		89.13	0.74	SUC 87	1		
Gemella morbillorum		8.62	0.45		99		
Actinomyces meyeri		1.22	0.31		1		
05111501	**Low discrimination**						**05111501**
					COC	SPO	
Lactobacillus fermentum		51.53	0.40	SUC 87	1	1	
Clostridium chauvoei		43.49	0.26	URE 1	1	99	
Gemella morbillorum		4.98	0.11		99	1	
05111502	**Very good identification**						**05111502**
Lactobacillus fermentum		99.60	0.74	SUC 87			
Clostridium perfringens		0.13	0.13				
Bifidobacterium longum subsp. longum		0.13	0.18				
05111503	**Low discrimination**						**05111503**
					SPO		
Lactobacillus fermentum		87.04	0.40	SUC 87	1		
Clostridium chauvoei		12.96	0.13	URE 1	99		
05111510	**Low discrimination**						**05111510**
					COC		
Lactobacillus fermentum		76.05	0.40	SUC 87	1		
Bifidobacterium longum subsp. longum		16.59	0.22	MLZ 1	1		
Gemella morbillorum		7.36	0.11		99		
05111512	**Low discrimination**						**05111512**
Lactobacillus fermentum		82.09	0.40	SUC 87			
Bifidobacterium longum subsp. longum		17.91	0.22	MLZ 1			
05111520	**Low discrimination**						**05111520**
					GRA		
Lactobacillus fermentum		47.28	0.66	SUC 87	99		
Prevotella bivia		29.53	0.37		1		
Clostridium perfringens		12.98	0.44		99		
05111522	**Low discrimination**						**05111522**
Lactobacillus fermentum		86.24	0.66	SUC 87			
Clostridium perfringens		11.14	0.38				
Bifidobacterium longum subsp. longum		1.34	0.28				
05111540	**Low discrimination**						**05111540**
					COC		
Lactobacillus fermentum		86.95	0.40	SUC 87	1		
Gemella morbillorum		8.41	0.11	SOR 1	99		
Clostridium perfringens		4.64	0.06		1		
05111542	**Good identification**						**05111542**
Lactobacillus fermentum		97.55	0.40	SUC 87			
Clostridium perfringens		2.45	0.00	SOR 1			
05111600	**Acceptable identification**						**05111600**
					COC	GRA	
Gemella morbillorum		91.13	0.45	RHA 1	99	99	
Lactobacillus fermentum		5.83	0.36	ARA 10	1	99	
Prevotella intermedia		2.54	0.17		1	1	
05111602	**Low discrimination**						**05111602**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC			
Lactobacillus fermentum		86.37	0.36	SUC 87	1			
Gemella morbillorum		13.63	0.11	RHA 1	99			
05111700	**Acceptable identification**							**05111700**
					COC			
Lactobacillus fermentum		91.18	0.40	SUC 87	1			
Gemella morbillorum		8.82	0.11	RHA 1	99			
05111702	**Good identification**							**05111702**
Lactobacillus fermentum		100.0	0.40	SUC 87				
05112000	**Low discrimination**							**05112000**
					COC	GRA		
Prevotella disiens		44.72	0.66	MAN 1	1	1		
Eubacterium limosum		25.04	0.63		1	99		
Gemella morbillorum		13.99	0.66		99	99		
05112002	**Low discrimination**							**05112002**
					COC	GRA	CAT	
Propionibacterium granulosum		33.50	0.53	SUC 93	1	99	90	
Megasphaera elsdenii		31.72	0.47		99	1	1	
Prevotella disiens		16.75	0.32		1	1	1	
05112010	**Low discrimination**							**05112010**
					GRA	SPO	CAT	
Clostridium difficile		87.18	0.58	MLT 1	99	99	1	
Propionibacterium granulosum		4.35	0.48		99	1	90	
Prevotella disiens		3.99	0.32		1	1	1	
05112012	**Low discrimination**							**05112012**
					COC	SPO	CAT	
Propionibacterium granulosum		51.60	0.38	SUC 93	1	1	90	
Clostridium difficile		41.80	0.24	MLZ 12	1	99	1	
Megasphaera elsdenii		3.62	0.12		99	1	1	
05112020	**Low discrimination**							**05112020**
					COC	GRA	CAT	
Gemella morbillorum		43.05	0.50	MAN 1	99	99	1	
Propionibacterium granulosum		25.02	0.53	NAG 10	1	99	90	
Prevotella disiens		12.51	0.32		1	1	1	
05112022	**Low discrimination**							**05112022**
					COC	GRA	CAT	
Propionibacterium granulosum		84.66	0.42	SUC 93	1	99	90	
Gemella morbillorum		5.89	0.15		99	99	1	
Megasphaera elsdenii		3.24	0.12		99	1	1	
05112030	**Low discrimination**							**05112030**
					COC	SPO	CAT	
Propionibacterium granulosum		49.57	0.38	SUC 93	1	1	90	
Clostridium difficile		40.16	0.24	MLZ 12	1	99	1	
Gemella morbillorum		6.32	0.15		99	1	1	
05112040	**Low discrimination**							**05112040**
					COC	GRA	SPO	
Megasphaera elsdenii		69.77	0.60	MNS 1	99	1	1	
Clostridium botulinum biovar B		15.11	0.41		1	99	99	
Prevotella disiens		6.14	0.32		1	1	1	
05112042	**Acceptable identification**							**05112042**
					COC	SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		93.74	0.44	MNS 1	99	1	1	
Clostridium botulinum biovar B		1.85	0.07	BGL 10	1	99	1	
Propionibacterium granulosum		1.50	0.18		1	1	90	
05112050	**Low discrimination**							**05112050**
					COC	GRA		
Clostridium botulinum biovar B		48.25	0.36	SUC 90	1	99		
Clostridium difficile		45.83	0.36	MAN 1	1	99		
Megasphaera elsdenii		4.57	0.26		99	1		
05112060	**Low discrimination**							**05112060**
					COC	SPO	CAT	
Clostridium botulinum biovar B		81.26	0.36	SUC 90	1	99	1	
Megasphaera elsdenii		7.70	0.26	MAN 1	99	1	1	
Propionibacterium acnes		4.57	0.18		1	1	95	
05112102	**Acceptable identification**							**05112102**
					COC	CAT	GRA	
Propionibacterium propionicum		92.75	0.49	NIT 86	1	1	99	
Propionibacterium granulosum		2.66	0.18	SUC 99	1	90	99	
Megasphaera elsdenii		2.52	0.12		99	1	1	
05112120	**Good identification**							**05112120**
					COC	GRA	CAT	
Prevotella bivia		99.55	0.50	FRU 10	1	1	1	
Gemella morbillorum		0.15	0.15	MAN 1	99	99	1	
Propionibacterium granulosum		0.09	0.18		1	99	90	
05112140	**Low discrimination**							**05112140**
					COC	SPO	GRA	
Propionibacterium propionicum		75.15	0.46	NIT 86	1	1	99	
Megasphaera elsdenii		18.36	0.26	SUC 99	99	1	1	
Clostridium botulinum biovar B		3.98	0.07		1	99	99	
05112142	**Good identification**							**05112142**
					COC	GRA		
Propionibacterium propionicum		98.22	0.49	NIT 86	1	99		
Megasphaera elsdenii		1.78	0.09	SUC 99	99	1		
05112400	**Low discrimination**							**05112400**
					COC	GRA		
Gemella morbillorum		64.24	0.50	MAN 1	99	99		
Prevotella disiens		18.67	0.32	ARA 10	1	1		
Eubacterium limosum		10.45	0.29		1	99		
05113000	**Low discrimination**							**05113000**
					COC	GRA	CAT	
Gemella morbillorum		52.57	0.61	MAN 1	99	99	1	
Propionibacterium granulosum		25.21	0.63		1	99	90	
Prevotella intermedia		16.10	0.52		1	1	1	
05113002	**Low discrimination**							**05113002**
					COC	CAT		
Propionibacterium granulosum		83.68	0.53	SUC 93	1	90		
Gemella morbillorum		7.05	0.27		99	1		
Lactobacillus fermentum		4.96	0.36		1	1		
05113010	**Low discrimination**							**05113010**
					COC	SPO	CAT	
Propionibacterium granulosum		70.49	0.48	SUC 93	1	1	90	
Clostridium difficile		14.28	0.24	MLZ 12	1	99	1	
Gemella morbillorum		10.89	0.27		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05113012	**Good identification**							**05113012**
Propionibacterium granulosum		100.0	0.38	SUC 93				
05113020	**Low discrimination**							**05113020**
					COC	CAT		
Propionibacterium granulosum		43.82	0.53	SUC 93	1	90		
Gemella morbillorum		40.61	0.45		99	1		
Propionibacterium acnes		6.80	0.29		1	95		
05113022	**Low discrimination**							**05113022**
					CAT			
Propionibacterium granulosum		74.54	0.42	SUC 93	90			
Clostridium perfringens		16.78	0.24		1			
Lactobacillus fermentum		5.90	0.28		1			
05113030	**Acceptable identification**							**05113030**
					COC	CAT		
Propionibacterium granulosum		93.58	0.38	SUC 93	1	90		
Gemella morbillorum		6.42	0.11	MLZ 12	99	1		
05113100	**Low discrimination**							**05113100**
					COC			
Propionibacterium propionicum		41.40	0.46	NIT 86	1			
Lactobacillus fermentum		17.06	0.40	SUC 99	1			
Gemella morbillorum		14.84	0.27		99			
05113102	**Low discrimination**							**05113102**
Propionibacterium propionicum		73.84	0.49	NIT 86				
Lactobacillus fermentum		20.28	0.40	SUC 99				
Clostridium perfringens		2.67	0.13					
05113120	**Low discrimination**							**05113120**
					GRA	SPO		
Prevotella bivia		58.04	0.37	FRU 10	1	1		
Clostridium perfringens		25.52	0.44	MAN 1	99	84		
Clostridium septicum		13.36	0.33		99	99		
05113122	**Low discrimination**							**05113122**
					GRA			
Clostridium perfringens		82.51	0.38	SUC 91	99			
Lactobacillus fermentum		6.45	0.32	MAN 1	99			
Prevotella bivia		4.03	0.03		1			
05113140	**Good identification**							**05113140**
Propionibacterium propionicum		97.85	0.46	NIT 86				
Clostridium perfringens		2.15	0.06	SUC 99				
05113142	**Good identification**							**05113142**
Propionibacterium propionicum		99.32	0.49	NIT 86				
Clostridium perfringens		0.69	0.00	SUC 99				
05113200	**Low discrimination**							**05113200**
					COC	GRA	CAT	
Propionibacterium granulosum		51.20	0.37	SUC 93	1	99	90	
Gemella morbillorum		34.86	0.27	RHA 3	99	99	1	
Prevotella intermedia		10.68	0.17		1	1	1	
05113400	**Low discrimination**							**05113400**
					COC	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		87.42	0.45	MAN 1	99	1		
Lactobacillus fermentum		5.60	0.36	ARA 10	1	1		
Propionibacterium granulosum		3.81	0.29		1	90		
05113402	**Low discrimination**							**05113402**
					COC	CAT		
Lactobacillus fermentum		75.30	0.36	SUC 87	1	1		
Propionibacterium granulosum		12.82	0.18	MAN 1	1	90		
Gemella morbillorum		11.88	0.11		99	1		
05113500	**Low discrimination**							**05113500**
					COC			
Lactobacillus fermentum		89.19	0.40	SUC 87	1			
Gemella morbillorum		8.63	0.11	MAN 1	99			
Propionibacterium propionicum		2.19	0.12		1			
05113502	**Good identification**							**05113502**
Lactobacillus fermentum		96.45	0.40	SUC 87				
Propionibacterium propionicum		3.55	0.15	MAN 1				
05114000	**Low discrimination**							**05114000**
					COC	GRA		
Gemella morbillorum		75.64	0.84	XYL 10	99	99		
Prevotella disiens		21.98	0.66		1	1		
Prevotella intermedia		1.13	0.52		1	1		
05114001	**Low discrimination**							**05114001**
					COC	GRA		
Gemella morbillorum		75.05	0.50	XYL 10	99	99		
Prevotella disiens		21.81	0.32	URE 1	1	1		
Prevotella intermedia		1.13	0.17		1	1		
05114002	**Low discrimination**							**05114002**
					COC	GRA		
Gemella morbillorum		73.13	0.50	XYL 10	99	99		
Prevotella disiens		21.25	0.32	BGL 1	1	1		
Lactobacillus fermentum		1.48	0.32		1	99		
05114010	**Low discrimination**							**05114010**
					COC	GRA	SPO	
Gemella morbillorum		73.18	0.50	XYL 10	99	99	1	
Prevotella disiens		21.27	0.32	MLZ 1	1	1	1	
Clostridium difficile		2.72	0.19		1	99	99	
05114020	**Very good identification**							**05114020**
					COC	GRA		
Gemella morbillorum		96.17	0.67	XYL 10	99	99		
Prevotella disiens		2.54	0.32	NAG 10	1	1		
Prevotella bivia		0.58	0.15		1	1		
05114040	**Low discrimination**							**05114040**
					COC	SPO		
Clostridium glycolicum		88.07	0.66	MNS 1	1	99		
Gemella morbillorum		5.76	0.50		99	1		
Clostridium botulinum biovar B		4.12	0.41		1	99		
05114050	**Low discrimination**							**05114050**
					COC	SPO		
Clostridium botulinum biovar B		67.26	0.36	SUC 90	1	99		
Clostridium glycolicum		29.50	0.32	XYL 1	1	99		
Gemella morbillorum		1.93	0.15		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05114060	**Low discrimination**						**05114060**
					COC	SPO	
Clostridium botulinum biovar B		56.74	0.36	SUC 90	1	99	
Clostridium glycolicum		24.88	0.32	XYL 1	1	99	
Gemella morbillorum		17.90	0.33		99	1	
05114100	**Low discrimination**						**05114100**
					COC	GRA	
Gemella morbillorum		53.85	0.50	XYL 10	99	99	
Actinomyces meyeri		21.32	0.44	LAC 1	1	99	
Prevotella disiens		15.65	0.32		1	1	
05114102	**Low discrimination**						**05114102**
					COC		
Lactobacillus fermentum		52.32	0.36	SUC 87	1		
Actinomyces meyeri		19.40	0.18	GAL 99	1		
Gemella morbillorum		16.00	0.15		99		
05114120	**Good identification**						**05114120**
					COC	GRA	
Prevotella bivia		97.23	0.50	FRU 10	1	1	
Gemella morbillorum		1.65	0.33	XYL 1	99	99	
Actinomyces meyeri		0.80	0.29		1	99	
05114200	**Low discrimination**						**05114200**
					COC	GRA	
Gemella morbillorum		75.84	0.50	XYL 10	99	99	
Prevotella disiens		22.04	0.32	RHA 1	1	1	
Prevotella intermedia		1.14	0.17		1	1	
05114400	**Very good identification**						**05114400**
					COC	GRA	
Gemella morbillorum		96.80	0.67	XYL 10	99	99	
Prevotella disiens		2.56	0.32	ARA 10	1	1	
Actinomyces meyeri		0.22	0.23		1	99	
05114420	**Good identification**						**05114420**
					COC		
Gemella morbillorum		98.94	0.51	XYL 10	99		
Lactobacillus fermentum		0.55	0.24	ARA 10	1		
Actinomyces meyeri		0.27	0.08	NAG 10	1		
05114502	**Acceptable identification**						**05114502**
Lactobacillus fermentum		91.52	0.36	SUC 87			
Actinomyces meyeri		8.48	0.07	GAL 99			
05115000	**Good identification**						**05115000**
					COC	GRA	
Gemella morbillorum		92.88	0.79	XYL 10	99	99	
Lactobacillus fermentum		3.49	0.67		1	99	
Prevotella intermedia		2.59	0.52		1	1	
05115001	**Acceptable identification**						**05115001**
					COC	GRA	
Gemella morbillorum		92.83	0.45	XYL 10	99	99	
Lactobacillus fermentum		3.49	0.32	URE 1	1	99	
Prevotella intermedia		2.59	0.17		1	1	
05115002	**Low discrimination**						**05115002**
					COC	GRA	
Lactobacillus fermentum		77.72	0.67	SUC 87	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		20.88	0.45		99	99	
Prevotella intermedia		0.58	0.17		1	1	
05115010	**Acceptable identification**						**05115010**
					COC	GRA	
Gemella morbillorum		92.46	0.45	XYL 10	99	99	
Lactobacillus fermentum		3.48	0.32	MLZ 1	1	99	
Prevotella intermedia		2.58	0.17		1	1	
05115020	**Low discrimination**						**05115020**
					COC		
Gemella morbillorum		88.09	0.63	XYL 10	99		
Lactobacillus fermentum		9.94	0.58	NAG 10	1		
Clostridium perfringens		1.03	0.29		1		
05115022	**Low discrimination**						**05115022**
					COC		
Lactobacillus fermentum		87.33	0.58	SUC 87	1		
Gemella morbillorum		7.82	0.29		99		
Clostridium perfringens		4.27	0.24		1		
05115040	**Low discrimination**						**05115040**
					COC	SPO	
Clostridium glycolicum		74.37	0.50	MNS 1	1	99	
Gemella morbillorum		23.57	0.45	GAL 10	99	1	
Lactobacillus fermentum		0.89	0.32		1	1	
05115100	**Low discrimination**						**05115100**
					COC		
Lactobacillus fermentum		71.54	0.70	SUC 87	1		
Actinomyces meyeri		12.99	0.47		1		
Gemella morbillorum		11.78	0.45		99		
05115102	**Very good identification**						**05115102**
					COC		
Lactobacillus fermentum		98.49	0.70	SUC 87	1		
Actinomyces meyeri		0.55	0.21		1		
Atopobium parvulum		0.29	0.14		99		
05115103	**Low discrimination**						**05115103**
					SPO		
Lactobacillus fermentum		79.77	0.36	SUC 87	1		
Clostridium chauvoei		20.23	0.13	URE 1	99		
05115110	**Low discrimination**						**05115110**
					COC		
Lactobacillus fermentum		74.28	0.36	SUC 87	1		
Actinomyces meyeri		13.49	0.13	MLZ 1	1		
Gemella morbillorum		12.23	0.11		99		
05115112	**Good identification**						**05115112**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05115120	**Low discrimination**						**05115120**
					GRA		
Prevotella bivia		36.00	0.37	FRU 10	1		
Lactobacillus fermentum		33.85	0.62	XYL 1	99		
Clostridium perfringens		15.83	0.44		99		
05115122	**Low discrimination**						**05115122**
					GRA		
Lactobacillus fermentum		76.60	0.62	SUC 87	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		16.85	0.38		99		
Prevotella oralis		2.75	0.22		1		
05115140	**Low discrimination**						**05115140**
					SPO		
Lactobacillus fermentum		51.11	0.36	SUC 87	1		
Clostridium glycolicum		26.55	0.15	SOR 1	99		
Actinomyces meyeri		9.28	0.13		1		
05115142	**Good identification**						**05115142**
Lactobacillus fermentum		95.90	0.36	SUC 87			
Clostridium perfringens		4.10	0.00	SOR 1			
05115200	**Acceptable identification**						**05115200**
					COC	GRA	
Gemella morbillorum		93.14	0.45	XYL 10	99	99	
Lactobacillus fermentum		3.50	0.32	RHA 1	1	99	
Prevotella intermedia		2.59	0.17		1	1	
05115300	**Low discrimination**						**05115300**
					COC		
Lactobacillus fermentum		74.28	0.36	SUC 87	1		
Actinomyces meyeri		13.49	0.13	RHA 1	1		
Gemella morbillorum		12.23	0.11		99		
05115302	**Good identification**						**05115302**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05115322	**Low discrimination**						**05115322**
					GRA		
Parabacteroides distasonis		66.25	0.37	RAF 99	1		
Prevotella oralis		18.73	0.15		1		
Lactobacillus fermentum		12.31	0.28		99		
05115400	**Low discrimination**						**05115400**
					COC		
Gemella morbillorum		74.20	0.63	XYL 10	99		
Lactobacillus fermentum		25.11	0.67	ARA 10	1		
Actinomyces meyeri		0.47	0.26		1		
05115402	**Very good identification**						**05115402**
					COC		
Lactobacillus fermentum		97.10	0.67	SUC 87	1		
Gemella morbillorum		2.90	0.29		99		
05115420	**Low discrimination**						**05115420**
					COC		
Lactobacillus fermentum		50.11	0.58	SUC 87	1		
Gemella morbillorum		49.36	0.46		99		
Actinomyces meyeri		0.38	0.11		1		
05115422	**Very good identification**						**05115422**
					COC		
Lactobacillus fermentum		99.02	0.58	SUC 87	1		
Gemella morbillorum		0.99	0.12		99		
05115500	**Good identification**						**05115500**
					COC		
Lactobacillus fermentum		93.38	0.70	SUC 87	1		
Actinomyces meyeri		4.24	0.36		1		
Gemella morbillorum		1.71	0.29		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05115501	**Good identification**					**05115501**
Lactobacillus fermentum		95.66	0.36	SUC 87		
Actinomyces meyeri		4.34	0.02	URE 1		
05115502	**Very good identification**					**05115502**
Lactobacillus fermentum		99.04	0.70	SUC 87		
Bifidobacterium longum subsp. longum		0.66	0.26			
Bifidobacterium adolescentis		0.17	0.17			
05115503	**Good identification**					**05115503**
Lactobacillus fermentum		100.0	0.36	SUC 87		
05115520	**Low discrimination**					**05115520**
					GRA	
Lactobacillus fermentum		76.79	0.62	SUC 87	99	
Bacteroides vulgatus		13.84	0.28		1	
Bifidobacterium longum subsp. longum		6.09	0.36		99	
05115522	**Good identification**					**05115522**
					GRA	
Lactobacillus fermentum		91.37	0.62	SUC 87	99	
Bifidobacterium longum subsp. longum		7.25	0.36		99	
Parabacteroides distasonis		1.02	0.26		1	
05115530	**Acceptable identification**					**05115530**
					GRA	
Bifidobacterium longum subsp. longum		91.89	0.40	SUC 99	99	
Lactobacillus fermentum		6.87	0.28	RAF 97	99	
Bacteroides vulgatus		1.24	-.06		1	
05115532	**Low discrimination**					**05115532**
					GRA	
Bifidobacterium longum subsp. longum		89.36	0.40	SUC 99	99	
Lactobacillus fermentum		6.68	0.28	RAF 97	99	
Parabacteroides distasonis		3.96	0.21		1	
05115540	**Good identification**					**05115540**
Lactobacillus fermentum		95.66	0.36	SUC 87		
Actinomyces meyeri		4.34	0.02	SOR 1		
05115542	**Good identification**					**05115542**
Lactobacillus fermentum		100.0	0.36	SUC 87		
05115700	**Low discrimination**					**05115700**
					GRA	
Lactobacillus fermentum		63.05	0.36	SUC 87	99	
Bacteroides vulgatus		34.09	0.10	RHA 1	1	
Actinomyces meyeri		2.86	0.02		99	
05115702	**Good identification**					**05115702**
Lactobacillus fermentum		100.0	0.36	SUC 87		
05115720	**Good identification**					**05115720**
					GRA	
Bacteroides vulgatus		99.38	0.44	SUC 96	1	
Lactobacillus fermentum		0.62	0.28	RAF 97	99	
05116000	**Low discrimination**					**05116000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Gemella morbillorum		61.54	0.50	MAN 1	99	99		
Prevotella disiens		17.88	0.32	XYL 10	1	1		
Eubacterium limosum		10.01	0.29		1	99		
05116010	**Acceptable identification**							**05116010**
					COC	SPO	CAT	
Clostridium difficile		94.93	0.36	MLT 1	1	99	1	
Gemella morbillorum		2.87	0.15	XYL 5	99	1	1	
Propionibacterium granulosum		0.91	0.14		1	1	90	
05117000	**Low discrimination**							**05117000**
					COC	CAT		
Gemella morbillorum		89.26	0.45	MAN 1	99	1		
Propionibacterium granulosum		3.89	0.29	XYL 10	1	90		
Lactobacillus fermentum		3.36	0.32		1	1		
05117100	**Low discrimination**							**05117100**
					COC			
Lactobacillus fermentum		72.05	0.36	SUC 87	1			
Actinomyces meyeri		13.08	0.13	MAN 1	1			
Gemella morbillorum		11.86	0.11		99			
05117102	**Acceptable identification**							**05117102**
Lactobacillus fermentum		94.11	0.36	SUC 87				
Propionibacterium propionicum		5.89	0.15	MAN 1				
05117500	**Good identification**							**05117500**
Lactobacillus fermentum		95.66	0.36	SUC 87				
Actinomyces meyeri		4.34	0.02	MAN 1				
05117502	**Low discrimination**							**05117502**
Lactobacillus fermentum		87.72	0.36	SUC 87				
Bifidobacterium adolescentis		12.28	0.16	MAN 1				
05120000	**Low discrimination**							**05120000**
					COC	GRA	SPO	
Prevotella disiens		57.41	0.66	TRE 1	1	1	1	
Clostridium botulinum biovar A		21.66	0.64		1	99	99	
Gemella morbillorum		14.70	0.64		99	99	1	
05120002	**Low discrimination**							**05120002**
					GRA	SPO		
Clostridium sporogenes		40.01	0.37	MNS 1	99	99		
Prevotella disiens		32.83	0.32		1	1		
Clostridium botulinum biovar A		12.39	0.30		99	99		
05120020	**Low discrimination**							**05120020**
					COC	GRA		
Gemella morbillorum		36.42	0.48	TRE 1	99	99		
Prevotella bivia		26.58	0.32	NAG 10	1	1		
Prevotella disiens		12.94	0.32		1	1		
05120040	**Good identification**							**05120040**
					GRA	SPO		
Clostridium botulinum biovar B		93.53	0.73	SUC 90	99	99		
Clostridium botulinum biovar A		5.71	0.55		99	99		
Prevotella disiens		0.51	0.32		1	1		
05120041	**Acceptable identification**							**05120041**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		90.87	0.39	SUC 90				
Clostridium botulinum biovar A		5.55	0.21	URE 1				
Clostridium novyi biovar A		3.09	0.29					
05120042	**Acceptable identification**							**05120042**
Clostridium botulinum biovar B		92.66	0.39	SUC 90				
Clostridium botulinum biovar A		5.66	0.21	BGL 1				
Clostridium sporogenes		0.62	0.03					
05120050	**Very good identification**							**05120050**
Clostridium botulinum biovar B		99.79	0.68	SUC 90				
Clostridium botulinum biovar A		0.13	0.21					
Clostridium novyi biovar A		0.07	0.29					
05120060	**Very good identification**							**05120060**
					COC	SPO		
Clostridium botulinum biovar B		99.79	0.68	SUC 90	1	99		
Clostridium botulinum biovar A		0.13	0.21		1	99		
Gemella morbillorum		0.03	0.14		99	1		
05120070	**Very good identification**							**05120070**
Clostridium botulinum biovar B		99.97	0.63	SUC 90				
Clostridium novyi biovar A		0.04	0.18					
05120120	**Very good identification**							**05120120**
					COC	GRA		
Prevotella bivia		99.95	0.66	TRE 1	1	1		
Clostridium perfringens		0.02	0.15		1	99		
Gemella morbillorum		0.01	0.14		99	99		
05120140	**Acceptable identification**							**05120140**
					GRA	SPO		
Clostridium botulinum biovar B		92.78	0.39	SUC 90	99	99		
Clostridium botulinum biovar A		5.67	0.21	LAC 1	99	99		
Prevotella bivia		1.04	-.03		1	1		
05120240	**Acceptable identification**							**05120240**
					GRA	SPO		
Clostridium botulinum biovar B		93.76	0.39	SUC 90	99	99		
Clostridium botulinum biovar A		5.73	0.21	RHA 1	99	99		
Prevotella disiens		0.51	-.03		1	1		
05120400	**Low discrimination**							**05120400**
					COC	GRA	SPO	
Gemella morbillorum		65.54	0.48	TRE 1	99	99	1	
Prevotella disiens		23.28	0.32	ARA 10	1	1	1	
Clostridium botulinum biovar A		8.78	0.30		1	99	99	
05120440	**Acceptable identification**							**05120440**
					COC	SPO		
Clostridium botulinum biovar B		92.43	0.39	SUC 90	1	99		
Clostridium botulinum biovar A		5.65	0.21	ARA 1	1	99		
Gemella morbillorum		1.42	0.14		99	1		
05121000	**Low discrimination**							**05121000**
					COC	GRA	SPO	
Gemella morbillorum		87.77	0.60	TRE 1	99	99	1	
Prevotella disiens		6.43	0.32		1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		2.43	0.30		1	99	99	
05121020	**Low discrimination**							**05121020**
					COC	GRA		
Gemella morbillorum		61.75	0.43	TRE 1	99	99		
Clostridium perfringens		18.09	0.34	NAG 10	1	99		
Prevotella bivia		14.77	0.19		1	1		
05121040	**Low discrimination**							**05121040**
					COC	SPO		
Clostridium botulinum biovar B		87.34	0.39	SUC 90	1	99		
Gemella morbillorum		6.52	0.26	GAL 1	99	1		
Clostridium botulinum biovar A		5.33	0.21		1	99		
05121101	**Good identification**							**05121101**
Clostridium chauvoei		100.0	0.37	SUC 99				
05121120	**Low discrimination**							**05121120**
					GRA	SPO		
Prevotella bivia		85.57	0.53	TRE 1	1	1		
Clostridium perfringens		7.76	0.49		99	84		
Clostridium septicum		6.57	0.41		99	99		
05121122	**Low discrimination**							**05121122**
					GRA	SPO		
Clostridium perfringens		78.54	0.43	SUC 91	99	84		
Prevotella bivia		18.58	0.19	FRU 99	1	1		
Clostridium septicum		2.88	0.12		99	99		
05121160	**Low discrimination**							**05121160**
					GRA	SPO		
Clostridium perfringens		60.82	0.36	SUC 91	99	84		
Prevotella bivia		35.55	0.19	FRU 99	1	1		
Clostridium septicum		2.73	0.07		99	99		
05121400	**Good identification**							**05121400**
					COC	GRA		
Gemella morbillorum		99.34	0.43	TRE 1	99	99		
Prevotella disiens		0.66	-.03	ARA 10	1	1		
05122040	**Acceptable identification**							**05122040**
					GRA	SPO		
Clostridium botulinum biovar B		93.47	0.39	SUC 90	99	99		
Clostridium botulinum biovar A		5.71	0.21	MAN 1	99	99		
Prevotella disiens		0.51	-.03		1	1		
05123000	**Low discrimination**							**05123000**
					COC	GRA	CAT	
Propionibacterium granulosum		62.10	0.37	SUC 93	1	99	90	
Gemella morbillorum		35.31	0.26	FRU 99	99	99	1	
Prevotella disiens		2.59	-.03		1	1	1	
05124000	**Low discrimination**							**05124000**
					COC	GRA	SPO	
Gemella morbillorum		65.54	0.48	TRE 1	99	99	1	
Prevotella disiens		23.28	0.32	XYL 10	1	1	1	
Clostridium botulinum biovar A		8.78	0.30		1	99	99	
05124040	**Low discrimination**							**05124040**
Clostridium botulinum biovar B		89.77	0.39	SUC 90				
Clostridium botulinum biovar A		5.48	0.21	XYL 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium glycolicum		2.87	0.15					
05125000	**Good identification**							**05125000**
					COC	GRA		
Gemella morbillorum		99.34	0.43	TRE 1	99	99		
Prevotella disiens		0.66	-.03	XYL 10	1	1		
05130000	**Low discrimination**							**05130000**
					COC	GRA	CAT	
Prevotella disiens		55.47	0.66	TRE 1	1	1	1	
Propionibacterium granulosum		17.82	0.73		1	99	90	
Gemella morbillorum		17.35	0.66		99	99	1	
05130002	**Low discrimination**							**05130002**
					COC	GRA	CAT	
Propionibacterium granulosum		81.82	0.63	SUC 93	1	99	90	
Prevotella disiens		10.29	0.32		1	1	1	
Gemella morbillorum		3.22	0.32		99	99	1	
05130010	**Low discrimination**							**05130010**
					GRA	SPO	CAT	
Propionibacterium granulosum		45.36	0.58	SUC 93	99	1	90	
Clostridium botulinum biovar B		38.02	0.44	MLZ 12	99	99	1	
Prevotella disiens		10.46	0.32		1	1	1	
05130012	**Good identification**							**05130012**
					GRA	SPO	CAT	
Propionibacterium granulosum		95.86	0.48	SUC 93	99	1	90	
Clostridium botulinum biovar B		3.25	0.10	MLZ 12	99	99	1	
Prevotella disiens		0.89	-.03		1	1	1	
05130020	**Low discrimination**							**05130020**
					COC	SPO	CAT	
Propionibacterium granulosum		46.55	0.63	SUC 93	1	1	90	
Clostridium botulinum biovar B		21.28	0.44		1	99	1	
Gemella morbillorum		20.14	0.50		99	1	1	
05130022	**Low discrimination**							**05130022**
					CAT	SPO		
Propionibacterium granulosum		86.89	0.52	SUC 93	90	1		
Clostridium perfringens		9.13	0.28		1	84		
Clostridium botulinum biovar B		1.61	0.10		1	99		
05130030	**Low discrimination**							**05130030**
					COC	SPO	CAT	
Clostridium botulinum biovar B		61.33	0.39	SUC 90	1	99	1	
Propionibacterium granulosum		37.14	0.48	SOR 99	1	1	90	
Gemella morbillorum		1.19	0.15		99	1	1	
05130032	**Acceptable identification**							**05130032**
					SPO	CAT		
Propionibacterium granulosum		93.75	0.38	SUC 93	1	90		
Clostridium botulinum biovar B		6.25	0.05	MLZ 12	99	1		
05130040	**Very good identification**							**05130040**
					GRA	SPO		
Clostridium botulinum biovar B		99.65	0.84	SUC 90	99	99		
Prevotella disiens		0.14	0.32		1	1		
Clostridium botulinum biovar A		0.10	0.35		99	99		
05130041	**Good identification**							**05130041**
					GRA	SPO		
Clostridium botulinum biovar B		99.67	0.50	SUC 90	99	99		

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Prevotella disiens		0.14	-.03	URE 1	1	1		
Clostridium botulinum biovar A		0.10	0.01		99	99		
05130042	**Good identification**							**05130042**
					COC	SPO	CAT	
Clostridium botulinum biovar B		98.53	0.50	SUC 90	1	99	1	
Propionibacterium granulosum		1.07	0.29	BGL 1	1	1	90	
Megasphaera elsdenii		0.17	0.09		99	1	1	
05130050	**Very good identification**							**05130050**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.78	SUC 90	99	1		
Propionibacterium granulosum		0.01	0.24		1	90		
Clostridium novyi biovar A		0.00	0.12		99	1		
05130051	**Good identification**							**05130051**
Clostridium botulinum biovar B		99.81	0.44	SUC 90				
Clostridium novyi biovar A		0.19	0.12	URE 1				
05130052	**Good identification**							**05130052**
					SPO	CAT		
Clostridium botulinum biovar B		99.70	0.44	SUC 90	99	1		
Propionibacterium granulosum		0.30	0.14	BGL 1	1	90		
05130060	**Very good identification**							**05130060**
					CAT	SPO		
Clostridium botulinum biovar B		99.94	0.78	SUC 90	1	99		
Clostridium perfringens		0.02	0.22		1	84		
Propionibacterium granulosum		0.02	0.29		90	1		
05130061	**Good identification**							**05130061**
Clostridium botulinum biovar B		100.0	0.44	SUC 90				
05130062	**Good identification**							**05130062**
					CAT	SPO		
Clostridium botulinum biovar B		98.38	0.44	SUC 90	1	99		
Clostridium perfringens		1.08	0.16	BGL 1	1	84		
Propionibacterium granulosum		0.54	0.18		90	1		
05130070	**Very good identification**							**05130070**
					SPO	CAT		
Clostridium botulinum biovar B		99.99	0.73	SUC 90	99	1		
Propionibacterium granulosum		0.01	0.14		1	90		
05130071	**Good identification**							**05130071**
Clostridium botulinum biovar B		100.0	0.39	SUC 90				
05130072	**Good identification**							**05130072**
Clostridium botulinum biovar B		100.0	0.39	SUC 90				
05130120	**Low discrimination**							**05130120**
					GRA	SPO		
Prevotella bivia		78.48	0.50	FRU 10	1	1		
Clostridium perfringens		11.31	0.49	TRE 1	99	84		
Clostridium septicum		9.56	0.41		99	99		
05130122	**Low discrimination**							**05130122**
					GRA	SPO		
Clostridium perfringens		83.42	0.43	SUC 91	99	84		
Prevotella bivia		12.43	0.15	GAL 99	1	1		

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Clostridium septicum		3.06	0.12		99	99		
05130140	**Good identification**							**05130140**
					GRA	SPO		
Clostridium botulinum biovar B		99.43	0.50	SUC 90	99	99		
Clostridium perfringens		0.26	0.11	LAC 1	99	84		
Prevotella disiens		0.14	-.03		1	1		
05130150	**Good identification**							**05130150**
Clostridium botulinum biovar B		100.0	0.44	SUC 90				
05130160	**Low discrimination**							**05130160**
					GRA	SPO		
Clostridium botulinum biovar B		80.58	0.44	SUC 90	99	99		
Clostridium perfringens		13.75	0.36	LAC 1	99	84		
Prevotella bivia		5.06	0.15		1	1		
05130170	**Good identification**							**05130170**
Clostridium botulinum biovar B		99.65	0.39	SUC 90				
Clostridium perfringens		0.35	0.02	LAC 1				
05130202	**Good identification**							**05130202**
					GRA	CAT		
Propionibacterium granulosum		96.06	0.37	SUC 93	99	90		
Prevotella disiens		3.95	-.03	RHA 3	1	1		
05130220	**Low discrimination**							**05130220**
					COC	SPO	CAT	
Propionibacterium granulosum		75.09	0.37	SUC 93	1	1	90	
Clostridium botulinum biovar B		11.21	0.10	RHA 3	1	99	1	
Gemella morbillorum		10.61	0.15		99	1	1	
05130240	**Good identification**							**05130240**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		99.63	0.50	SUC 90	99	99	1	
Prevotella disiens		0.14	-.03	RHA 1	1	1	1	
Propionibacterium granulosum		0.13	0.13		99	1	90	
05130250	**Good identification**							**05130250**
Clostridium botulinum biovar B		100.0	0.44	SUC 90				
05130260	**Good identification**							**05130260**
Clostridium botulinum biovar B		100.0	0.44	SUC 90				
05130270	**Good identification**							**05130270**
Clostridium botulinum biovar B		100.0	0.39	SUC 90				
05130400	**Low discrimination**							**05130400**
					COC	GRA	CAT	
Gemella morbillorum		69.95	0.50	TRE 1	99	99	1	
Prevotella disiens		20.33	0.32	ARA 10	1	1	1	
Propionibacterium granulosum		6.53	0.39		1	99	90	
05130440	**Good identification**							**05130440**
					COC	GRA	SPO	
Clostridium botulinum biovar B		99.30	0.50	SUC 90	1	99	99	
Gemella morbillorum		0.47	0.15	ARA 1	99	99	1	
Prevotella disiens		0.14	-.03		1	1	1	
05130450	**Good identification**							**05130450**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		100.0	0.44	SUC 90				
05130460	**Good identification**							**05130460**
Clostridium botulinum biovar B		100.0	0.44	SUC 90				
05130470	**Good identification**							**05130470**
Clostridium botulinum biovar B		100.0	0.39	SUC 90				
05131000	**Low discrimination**							**05131000**
					COC	GRA	CAT	
Propionibacterium granulosum		55.22	0.73	SUC 93	1	99	90	
Gemella morbillorum		28.95	0.61		99	99	1	
Prevotella intermedia		8.87	0.52		1	1	1	
05131001	**Low discrimination**							**05131001**
					COC	GRA	CAT	
Propionibacterium granulosum		53.15	0.39	SUC 93	1	99	90	
Gemella morbillorum		27.86	0.27	URE 1	99	99	1	
Prevotella intermedia		8.53	0.17		1	1	1	
05131002	**Low discrimination**							**05131002**
					CAT			
Propionibacterium granulosum		82.71	0.63	SUC 93	90			
Clostridium perfringens		6.66	0.37		1			
Lactobacillus jensenii		6.28	0.37		1			
05131010	**Low discrimination**							**05131010**
					COC	CAT		
Propionibacterium granulosum		88.16	0.58	SUC 93	1	90		
Lactobacillus jensenii		6.14	0.32	MLZ 12	1	1		
Gemella morbillorum		3.42	0.27		99	1		
05131012	**Low discrimination**							**05131012**
					CAT			
Propionibacterium granulosum		83.93	0.48	SUC 93	90			
Lactobacillus jensenii		15.58	0.29	MLZ 12	1			
Clostridium perfringens		0.50	0.02		1			
05131020	**Low discrimination**							**05131020**
					CAT	SPO		
Clostridium perfringens		77.19	0.68	SUC 91	1	84		
Propionibacterium granulosum		13.96	0.63	LAC 88	90	1		
Clostridium septicum		4.84	0.41		1	99		
05131022	**Good identification**							**05131022**
					CAT			
Clostridium perfringens		90.18	0.62	SUC 91	1			
Propionibacterium granulosum		8.67	0.52	LAC 88	90			
Lactobacillus jensenii		0.66	0.27		1			
05131030	**Low discrimination**							**05131030**
					CAT			
Propionibacterium granulosum		64.99	0.48	SUC 93	90			
Clostridium perfringens		26.62	0.34	MLZ 12	1			
Lactobacillus jensenii		4.52	0.21		1			
05131032	**Low discrimination**							**05131032**
					CAT			
Propionibacterium granulosum		51.11	0.38	SUC 93	90			
Clostridium perfringens		39.40	0.28	MLZ 12	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		9.49	0.18		1		
05131040	**Low discrimination**						**05131040**
					CAT	SPO	
Clostridium botulinum biovar B		89.93	0.50	SUC 90	1	99	
Propionibacterium granulosum		3.91	0.39	GAL 1	90	1	
Clostridium perfringens		3.16	0.30		1	84	
05131050	**Good identification**						**05131050**
					SPO	CAT	
Clostridium botulinum biovar B		98.81	0.44	SUC 90	99	1	
Propionibacterium granulosum		1.19	0.24	GAL 1	1	90	
05131060	**Low discrimination**						**05131060**
					CAT	SPO	
Clostridium perfringens		69.00	0.56	SUC 91	1	84	
Clostridium botulinum biovar B		29.96	0.44	LAC 88	1	99	
Propionibacterium granulosum		0.66	0.29		90	1	
05131062	**Good identification**						**05131062**
					CAT	SPO	
Clostridium perfringens		98.58	0.50	SUC 91	1	84	
Clostridium botulinum biovar B		0.92	0.10	LAC 88	1	99	
Propionibacterium granulosum		0.50	0.18		90	1	
05131070	**Acceptable identification**						**05131070**
					CAT	SPO	
Clostridium botulinum biovar B		94.94	0.39	SUC 90	1	99	
Clostridium perfringens		4.49	0.22	GAL 1	1	84	
Propionibacterium granulosum		0.58	0.14		90	1	
05131100	**Low discrimination**						**05131100**
					COC	SPO	
Atopobium parvulum		41.18	0.56	SUC 96	99	1	
Clostridium perfringens		40.60	0.57		1	84	
Clostridium septicum		11.21	0.41		1	99	
05131101	**Low discrimination**						**05131101**
					COC	SPO	
Clostridium chauvoei		87.56	0.44	SUC 99	1	99	
Atopobium parvulum		4.98	0.22	TRE 10	99	1	
Clostridium perfringens		4.91	0.23		1	84	
05131102	**Low discrimination**						**05131102**
					COC		
Clostridium perfringens		67.73	0.51	SUC 91	1		
Atopobium parvulum		25.76	0.44	NAG 97	99		
Lactobacillus fermentum		2.79	0.40		1		
05131120	**Low discrimination**						**05131120**
					GRA	SPO	
Clostridium perfringens		53.63	0.83	SUC 91	99	84	
Clostridium septicum		45.36	0.75		99	99	
Prevotella bivia		0.66	0.37		1	1	
05131121	**Low discrimination**						**05131121**
Clostridium perfringens		48.73	0.49	SUC 91			
Clostridium septicum		41.21	0.41	URE 1			
Clostridium chauvoei		8.96	0.35				
05131122	**Very good identification**						**05131122**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		96.11	0.77	SUC 91	1	84		
Clostridium septicum		3.53	0.46		1	99		
Atopobium parvulum		0.20	0.31		99	1		
05131123	**Low discrimination**							**05131123**
Clostridium perfringens		89.62	0.43	SUC 91				
Clostridium chauvoei		6.18	0.22	URE 1				
Clostridium septicum		3.29	0.12					
05131130	**Low discrimination**							**05131130**
					GRA	SPO		
Clostridium perfringens		53.56	0.49	SUC 91	99	84		
Clostridium septicum		45.30	0.41	MLZ 1	99	99		
Prevotella bivia		0.66	0.03		1	1		
05131132	**Good identification**							**05131132**
					GRA	SPO		
Clostridium perfringens		95.35	0.43	SUC 91	99	84		
Clostridium septicum		3.50	0.12	MLZ 1	99	99		
Parabacteroides distasonis		1.15	0.20		1	1		
05131140	**Low discrimination**							**05131140**
					COC	SPO		
Clostridium perfringens		89.89	0.45	SUC 91	1	84		
Atopobium parvulum		4.84	0.22	NAG 97	99	1		
Clostridium botulinum biovar B		3.53	0.15		1	99		
05131142	**Good identification**							**05131142**
					COC			
Clostridium perfringens		97.40	0.39	SUC 91	1			
Atopobium parvulum		1.97	0.09	NAG 97	99			
Propionibacterium propionicum		0.64	0.14		1			
05131160	**Very good identification**							**05131160**
Clostridium perfringens		95.57	0.71	SUC 91				
Clostridium septicum		4.29	0.41					
Clostridium botulinum biovar B		0.06	0.10					
05131161	**Acceptable identification**							**05131161**
Clostridium perfringens		94.82	0.36	SUC 91				
Clostridium septicum		4.25	0.07	URE 1				
Clostridium chauvoei		0.93	0.01					
05131162	**Very good identification**							**05131162**
Clostridium perfringens		99.81	0.65	SUC 91				
Clostridium septicum		0.19	0.12					
05131170	**Acceptable identification**							**05131170**
Clostridium perfringens		93.11	0.36	SUC 91				
Clostridium septicum		4.18	0.07	MLZ 1				
Clostridium botulinum biovar B		2.72	0.05					
05131200	**Low discrimination**							**05131200**
					COC	GRA	CAT	
Propionibacterium granulosum		79.54	0.47	SUC 93	1	99	90	
Gemella morbillorum		13.62	0.27	RHA 3	99	99	1	
Prevotella intermedia		4.17	0.17		1	1	1	

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05131202	**Good identification**						**05131202**
					CAT		
Propionibacterium granulosum		95.14	0.37	SUC 93	90		
Clostridium perfringens		2.50	0.02	RHA 3	1		
Lactobacillus jensenii		2.36	0.03		1		
05131320	**Low discrimination**						**05131320**
					GRA	SPO	
Clostridium perfringens		53.66	0.49	SUC 91	99	84	
Clostridium septicum		45.38	0.41	RHA 1	99	99	
Prevotella bivia		0.66	0.03		1	1	
05131322	**Low discrimination**						**05131322**
					GRA	SPO	
Clostridium perfringens		89.01	0.43	SUC 91	99	84	
Parabacteroides distasonis		6.00	0.32	RHA 1	1	1	
Clostridium septicum		3.26	0.12		99	99	
05131360	**Good identification**						**05131360**
Clostridium perfringens		95.71	0.36	SUC 91			
Clostridium septicum		4.29	0.07	RHA 1			
05131400	**Low discrimination**						**05131400**
					COC	CAT	
Gemella morbillorum		77.94	0.45	TRE 1	99	1	
Propionibacterium granulosum		13.51	0.39	ARA 10	1	90	
Lactobacillus fermentum		4.99	0.36		1	1	
05131402	**Low discrimination**						**05131402**
					COC	CAT	
Lactobacillus fermentum		51.52	0.36	SUC 87	1	1	
Propionibacterium granulosum		34.89	0.29	TRE 1	1	90	
Gemella morbillorum		8.13	0.11		99	1	
05131500	**Low discrimination**						**05131500**
					COC		
Lactobacillus fermentum		42.24	0.40	SUC 87	1		
Atopobium parvulum		22.30	0.22	TRE 1	99		
Clostridium perfringens		21.99	0.23		1		
05131502	**Low discrimination**						**05131502**
					COC		
Lactobacillus fermentum		72.99	0.40	SUC 87	1		
Clostridium perfringens		17.88	0.17	TRE 1	1		
Atopobium parvulum		6.80	0.09		99		
05131520	**Low discrimination**						**05131520**
					SPO		
Clostridium perfringens		52.88	0.49	SUC 91	84		
Clostridium septicum		44.72	0.41	ARA 1	99		
Lactobacillus fermentum		1.05	0.32		1		
05131522	**Acceptable identification**						**05131522**
					SPO		
Clostridium perfringens		90.20	0.43	SUC 91	84		
Lactobacillus fermentum		3.80	0.32	ARA 1	1		
Clostridium septicum		3.31	0.12		99		
05131560	**Good identification**						**05131560**
Clostridium perfringens		95.71	0.36	SUC 91			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium septicum		4.29	0.07	ARA 1				
05132000	**Good identification**							**05132000**
					GRA	CAT		
Propionibacterium granulosum		90.41	0.71	SUC 93	99	90		
Prevotella disiens		3.77	0.32		1	1		
Eubacterium limosum		2.11	0.29		99	1		
05132001	**Acceptable identification**							**05132001**
					GRA	CAT		
Propionibacterium granulosum		93.90	0.37	SUC 93	99	90		
Prevotella disiens		3.91	-.03	URE 1	1	1		
Eubacterium limosum		2.19	-.06		99	1		
05132002	**Very good identification**							**05132002**
					COC	GRA	CAT	
Propionibacterium granulosum		99.33	0.61	SUC 93	1	99	90	
Megasphaera elsdenii		0.32	0.12		99	1	1	
Prevotella disiens		0.17	-.03		1	1	1	
05132010	**Low discrimination**							**05132010**
					SPO	CAT		
Propionibacterium granulosum		65.83	0.56	SUC 93	1	90		
Clostridium difficile		33.12	0.38	MLZ 12	99	1		
Clostridium botulinum biovar B		0.74	0.10		99	1		
05132012	**Good identification**							**05132012**
					SPO	CAT		
Propionibacterium granulosum		98.01	0.46	SUC 93	1	90		
Clostridium difficile		1.99	0.04	MLZ 12	99	1		
05132020	**Very good identification**							**05132020**
					COC	SPO	CAT	
Propionibacterium granulosum		98.57	0.61	SUC 93	1	1	90	
Clostridium botulinum biovar B		0.60	0.10		1	99	1	
Gemella morbillorum		0.57	0.15		99	1	1	
05132022	**Good identification**							**05132022**
Propionibacterium granulosum		100.0	0.50	SUC 93				
05132030	**Good identification**							**05132030**
					SPO	CAT		
Propionibacterium granulosum		95.93	0.46	SUC 93	1	90		
Clostridium botulinum biovar B		2.12	0.05	MLZ 12	99	1		
Clostridium difficile		1.95	0.04		99	1		
05132032	**Good identification**							**05132032**
Propionibacterium granulosum		100.0	0.36	SUC 93				
05132040	**Acceptable identification**							**05132040**
					COC	SPO	CAT	
Clostridium botulinum biovar B		94.89	0.50	SUC 90	1	99	1	
Propionibacterium granulosum		3.12	0.37	MAN 1	1	1	90	
Megasphaera elsdenii		1.48	0.26		99	1	1	
05132050	**Good identification**							**05132050**
					SPO	CAT		
Clostridium botulinum biovar B		96.81	0.44	SUC 90	99	1		
Clostridium difficile		2.31	0.17	MAN 1	99	1		
Propionibacterium granulosum		0.88	0.22		1	90		
05132060	**Good identification**							**05132060**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	CAT		
Clostridium botulinum biovar B		98.36	0.44	SUC 90	99	1		
Propionibacterium granulosum		1.64	0.27	MAN 1	1	90		
05132070	**Good identification**							**05132070**
					SPO	CAT		
Clostridium botulinum biovar B		99.54	0.39	SUC 90	99	1		
Propionibacterium granulosum		0.46	0.12	MAN 1	1	90		
05132100	**Low discrimination**							**05132100**
					GRA	CAT		
Propionibacterium propionicum		59.27	0.45	NIT 86	99	1		
Propionibacterium granulosum		37.38	0.37	SUC 99	99	90		
Prevotella disiens		1.56	-.03		1	1		
05132102	**Acceptable identification**							**05132102**
					CAT			
Propionibacterium propionicum		90.49	0.48	NIT 86	1			
Propionibacterium granulosum		9.51	0.27	SUC 99	90			
05132140	**Low discrimination**							**05132140**
					SPO			
Propionibacterium propionicum		83.75	0.45	NIT 86	1			
Clostridium botulinum biovar B		16.25	0.15	SUC 99	99			
05132142	**Good identification**							**05132142**
Propionibacterium propionicum		100.0	0.48	NIT 86				
05132200	**Good identification**							**05132200**
					GRA	CAT		
Propionibacterium granulosum		97.92	0.45	SUC 93	99	90		
Prevotella disiens		1.33	-.03	RHA 3	1	1		
Eubacterium limosum		0.75	-.06		99	1		
05132400	**Low discrimination**							**05132400**
					COC	GRA	CAT	
Propionibacterium granulosum		82.75	0.37	SUC 93	1	99	90	
Gemella morbillorum		11.87	0.15	ARA 1	99	99	1	
Prevotella disiens		3.45	-.03		1	1	1	
05133000	**Very good identification**							**05133000**
					COC	GRA	CAT	
Propionibacterium granulosum		98.90	0.71	SUC 93	1	99	90	
Gemella morbillorum		0.69	0.27		99	99	1	
Prevotella intermedia		0.21	0.17		1	1	1	
05133001	**Good identification**							**05133001**
Propionibacterium granulosum		95.79	0.37	SUC 93				
Actinomyces naeslundii		4.21	0.28	URE 1				
05133002	**Very good identification**							**05133002**
					CAT			
Propionibacterium granulosum		99.58	0.61	SUC 93	90			
Actinomyces naeslundii		0.12	0.25		25			
Clostridium perfringens		0.11	0.02		1			
05133010	**Very good identification**							**05133010**
					SPO	CAT		
Propionibacterium granulosum		99.49	0.56	SUC 93	1	90		
Clostridium difficile		0.51	0.04	MLZ 12	99	1		
05133012	**Good identification**							**05133012**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.46	SUC 93			
05133020	**Good identification**						**05133020**
					CAT	SPO	
Propionibacterium granulosum		92.34	0.61	SUC 93	90	1	
Clostridium perfringens		6.84	0.34		1	84	
Clostridium septicum		0.43	0.07		1	99	
05133022	**Low discrimination**						**05133022**
					CAT		
Propionibacterium granulosum		87.41	0.50	SUC 93	90		
Clostridium perfringens		12.18	0.28		1		
Actinomyces naeslundii		0.41	0.25		25		
05133030	**Good identification**						**05133030**
Propionibacterium granulosum		100.0	0.46	SUC 93			
05133032	**Good identification**						**05133032**
Propionibacterium granulosum		100.0	0.36	SUC 93			
05133040	**Low discrimination**						**05133040**
					SPO	CAT	
Propionibacterium granulosum		76.48	0.37	SUC 93	1	90	
Clostridium botulinum biovar B		23.52	0.15	SOR 1	99	1	
05133100	**Low discrimination**						**05133100**
					COC	CAT	
Propionibacterium propionicum		43.35	0.45	NIT 86	1	1	
Propionibacterium granulosum		27.34	0.37	SUC 99	1	90	
Atopobium parvulum		11.52	0.22		99	1	
05133102	**Low discrimination**						**05133102**
					CAT		
Propionibacterium propionicum		79.77	0.48	NIT 86	1		
Propionibacterium granulosum		8.38	0.27	SUC 99	90		
Clostridium perfringens		6.56	0.17		1		
05133120	**Low discrimination**						**05133120**
					CAT	SPO	
Clostridium perfringens		52.96	0.49	SUC 91	1	84	
Clostridium septicum		44.78	0.41	MAN 1	1	99	
Propionibacterium granulosum		0.99	0.27		90	1	
05133122	**Acceptable identification**						**05133122**
					SPO		
Clostridium perfringens		94.03	0.43	SUC 91	84		
Clostridium septicum		3.45	0.12	MAN 1	99		
Actinomyces naeslundii		1.24	0.33		1		
05133123	**Low discrimination**						**05133123**
Actinomyces naeslundii		66.21	0.36	SUC 99			
Clostridium perfringens		33.79	0.09	RAF 92			
05133140	**Good identification**						**05133140**
Propionibacterium propionicum		95.25	0.45	NIT 86			
Clostridium perfringens		4.75	0.11	SUC 99			
05133142	**Good identification**						**05133142**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		98.46	0.48	NIT 86				
Clostridium perfringens		1.54	0.05	SUC 99				
05133160	**Good identification**							**05133160**
Clostridium perfringens		95.71	0.36	SUC 91				
Clostridium septicum		4.29	0.07	MAN 1				
05133200	**Good identification**							**05133200**
Propionibacterium granulosum		100.0	0.45	SUC 93				
05133400	**Acceptable identification**							**05133400**
					COC	CAT		
Propionibacterium granulosum		92.83	0.37	SUC 93	1	90		
Gemella morbillorum		7.17	0.11	ARA 1	99	1		
05134000	**Low discrimination**							**05134000**
					COC	GRA	CAT	
Gemella morbillorum		69.06	0.50	TRE 1	99	99	1	
Prevotella disiens		20.07	0.32	XYL 10	1	1	1	
Propionibacterium granulosum		6.45	0.39		1	99	90	
05134040	**Acceptable identification**							**05134040**
					COC	SPO		
Clostridium botulinum biovar B		92.67	0.50	SUC 90	1	99		
Clostridium glycolicum		6.67	0.32	XYL 1	1	99		
Gemella morbillorum		0.44	0.15		99	1		
05134050	**Good identification**							**05134050**
Clostridium botulinum biovar B		99.85	0.44	SUC 90				
Clostridium glycolicum		0.15	-.03	XYL 1				
05134060	**Good identification**							**05134060**
Clostridium botulinum biovar B		99.85	0.44	SUC 90				
Clostridium glycolicum		0.15	-.03	XYL 1				
05134070	**Good identification**							**05134070**
Clostridium botulinum biovar B		100.0	0.39	SUC 90				
05135000	**Low discrimination**							**05135000**
					COC	CAT		
Gemella morbillorum		72.61	0.45	TRE 1	99	1		
Propionibacterium granulosum		12.59	0.39	XYL 10	1	90		
Atopobium parvulum		7.43	0.27		99	1		
05135100	**Very good identification**							**05135100**
					COC			
Atopobium parvulum		98.18	0.61	SUC 96	99			
Lactobacillus fermentum		0.59	0.36		1			
Clostridium perfringens		0.53	0.23		1			
05135102	**Good identification**							**05135102**
					COC			
Atopobium parvulum		95.16	0.48	SUC 96	99			
Lactobacillus fermentum		3.26	0.36		1			
Clostridium perfringens		1.36	0.17		1			
05135120	**Low discrimination**							**05135120**
					COC	SPO		
Atopobium parvulum		35.25	0.48	SUC 96	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		34.63	0.49		1	84		
Clostridium septicum		29.29	0.41		1	99		
05135122	**Low discrimination**							**05135122**
					COC	GRA		
Clostridium perfringens		60.58	0.43	SUC 91	1	99		
Atopobium parvulum		23.12	0.35	XYL 1	99	99		
Parabacteroides distasonis		12.24	0.41		1	1		
05135132	**Low discrimination**							**05135132**
					COC	GRA		
Parabacteroides distasonis		88.63	0.36	RAF 99	1	1		
Clostridium perfringens		8.23	0.09		1	99		
Atopobium parvulum		3.14	0.01		99	99		
05135160	**Acceptable identification**							**05135160**
					COC	SPO		
Clostridium perfringens		91.01	0.36	SUC 91	1	84		
Atopobium parvulum		4.91	0.14	XYL 1	99	1		
Clostridium septicum		4.08	0.07		1	99		
05135322	**Good identification**							**05135322**
					COC	GRA		
Parabacteroides distasonis		97.75	0.49	RAF 99	1	1		
Clostridium perfringens		1.63	0.09		1	99		
Atopobium parvulum		0.62	0.01		99	99		
05135332	**Good identification**							**05135332**
Parabacteroides distasonis		100.0	0.44	RAF 99				
05135502	**Low discrimination**							**05135502**
					COC			
Lactobacillus fermentum		63.60	0.36	SUC 87	1			
Atopobium parvulum		18.73	0.14	TRE 1	99			
Bifidobacterium longum subsp. longum		10.41	0.15		1			
05135522	**Low discrimination**							**05135522**
					GRA			
Parabacteroides distasonis		58.68	0.37	RAF 99	1			
Bifidobacterium longum subsp. longum		23.00	0.26		99			
Lactobacillus fermentum		11.71	0.28		99			
05135722	**Good identification**							**05135722**
Parabacteroides distasonis		100.0	0.45	RAF 99				
05135732	**Good identification**							**05135732**
Parabacteroides distasonis		95.16	0.40	RAF 99				
Clostridium clostridioforme		4.85	0.13					
05136000	**Low discrimination**							**05136000**
					COC	SPO	CAT	
Propionibacterium granulosum		78.03	0.37	SUC 93	1	1	90	
Gemella morbillorum		11.19	0.15	XYL 1	99	1	1	
Clostridium difficile		5.71	0.05		1	99	1	
05137000	**Acceptable identification**							**05137000**
					COC	CAT		
Propionibacterium granulosum		92.83	0.37	SUC 93	1	90		
Gemella morbillorum		7.17	0.11	XYL 1	99	1		
05140000	**Low discrimination**							**05140000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Prevotella disiens		75.64	0.66	CEL 1	1	1		
Gemella morbillorum		19.36	0.64		99	99		
Peptostreptococcus anaerobius		1.81	0.39		99	99		
05140020	**Low discrimination**							**05140020**
					COC	GRA		
Gemella morbillorum		45.82	0.48	CEL 1	99	99		
Prevotella bivia		33.44	0.32	NAG 10	1	1		
Prevotella disiens		16.28	0.32		1	1		
05140120	**Very good identification**							**05140120**
					COC	GRA	SPO	
Prevotella bivia		99.97	0.66	CEL 1	1	1	1	
Gemella morbillorum		0.01	0.14		99	99	1	
Clostridium septicum		0.01	0.07		1	99	99	
05140400	**Low discrimination**							**05140400**
					COC	GRA		
Gemella morbillorum		72.60	0.48	CEL 1	99	99		
Prevotella disiens		25.79	0.32	ARA 10	1	1		
Peptostreptococcus anaerobius		0.62	0.05		99	99		
05141000	**Good identification**							**05141000**
					COC	GRA		
Gemella morbillorum		91.93	0.60	CEL 1	99	99		
Prevotella disiens		6.74	0.32		1	1		
Lactobacillus jensenii		0.39	0.19		1	99		
05141020	**Low discrimination**							**05141020**
					COC	GRA	SPO	
Gemella morbillorum		77.84	0.43	CEL 1	99	99	1	
Prevotella bivia		18.62	0.19	NAG 10	1	1	1	
Clostridium septicum		1.51	0.07		1	99	99	
05141120	**Acceptable identification**							**05141120**
					GRA	SPO		
Prevotella bivia		92.03	0.53	CEL 1	1	1		
Clostridium septicum		7.45	0.41		99	99		
Prevotella melaninogenica		0.29	0.17		1	1		
05141400	**Good identification**							**05141400**
					COC	GRA		
Gemella morbillorum		99.34	0.43	CEL 1	99	99		
Prevotella disiens		0.66	-.03	ARA 10	1	1		
05144000	**Low discrimination**							**05144000**
					COC	GRA		
Gemella morbillorum		72.60	0.48	CEL 1	99	99		
Prevotella disiens		25.79	0.32	XYL 10	1	1		
Peptostreptococcus anaerobius		0.62	0.05		99	99		
05145000	**Good identification**							**05145000**
					COC	GRA		
Gemella morbillorum		99.34	0.43	CEL 1	99	99		
Prevotella disiens		0.66	-.03	XYL 10	1	1		
05145722	**Good identification**							**05145722**
Bacteroides eggerthii		100.0	0.50	IND 99				
05150000	**Low discrimination**							**05150000**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella disiens		72.50	0.66	CEL 1	1	1		
Gemella morbillorum		22.68	0.66		99	99		
Prevotella intermedia		3.74	0.52		1	1		
05150020	**Low discrimination**							**05150020**
					COC	GRA		
Gemella morbillorum		70.39	0.50	CEL 1	99	99		
Prevotella disiens		20.46	0.32	NAG 10	1	1		
Prevotella bivia		4.67	0.15		1	1		
05150040	**Low discrimination**							**05150040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		61.38	0.41	SUC 90	1	99	99	
Prevotella disiens		24.95	0.32	CEL 1	1	1	1	
Gemella morbillorum		7.80	0.32		99	99	1	
05150050	**Good identification**							**05150050**
					GRA	SPO		
Clostridium botulinum biovar B		98.07	0.36	SUC 90	99	99		
Prevotella disiens		0.82	-.03	CEL 1	1	1		
Clostridium novyi biovar A		0.56	0.12		99	99		
05150060	**Good identification**							**05150060**
					COC	GRA	SPO	
Clostridium botulinum biovar B		96.43	0.36	SUC 90	1	99	99	
Gemella morbillorum		2.77	0.15	CEL 1	99	99	1	
Prevotella disiens		0.80	-.03		1	1	1	
05150120	**Low discrimination**							**05150120**
					GRA	SPO		
Prevotella bivia		88.14	0.50	FRU 10	1	1		
Clostridium septicum		11.34	0.41	CEL 1	99	99		
Clostridium perfringens		0.29	0.20		99	84		
05150400	**Low discrimination**							**05150400**
					COC	GRA		
Gemella morbillorum		76.59	0.50	CEL 1	99	99		
Prevotella disiens		22.26	0.32	ARA 10	1	1		
Prevotella intermedia		1.15	0.17		1	1		
05151000	**Low discrimination**							**05151000**
					COC	GRA		
Gemella morbillorum		56.64	0.61	CEL 1	99	99		
Lactobacillus jensenii		19.48	0.54		1	99		
Prevotella intermedia		17.35	0.52		1	1		
05151002	**Low discrimination**							**05151002**
					COC			
Lactobacillus jensenii		84.89	0.51	SUC 90	1			
Lactobacillus catenaformis		4.43	0.29	SAL 95	1			
Gemella morbillorum		3.74	0.27		99			
05151010	**Low discrimination**							**05151010**
					COC	GRA		
Lactobacillus jensenii		88.65	0.45	SUC 90	1	99		
Gemella morbillorum		7.81	0.27	SAL 95	99	99		
Prevotella intermedia		2.39	0.17		1	1		
05151012	**Good identification**							**05151012**
Lactobacillus jensenii		100.0	0.42	SUC 90				
05151020	**Low discrimination**							**05151020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium septicum		39.06	0.41	LAC 99	1	99	
Gemella morbillorum		24.90	0.45		99	1	
Lactobacillus jensenii		19.27	0.43		1	1	
05151022	**Low discrimination**						**05151022**
					GRA		
Lactobacillus jensenii		50.79	0.40	SUC 90	99		
Clostridium perfringens		24.66	0.34	SAL 95	99		
Prevotella oralis		13.09	0.26		1		
05151030	**Low discrimination**						**05151030**
					COC	SPO	
Lactobacillus jensenii		89.17	0.35	SUC 90	1	1	
Clostridium septicum		5.48	0.07	SAL 95	1	99	
Gemella morbillorum		3.49	0.11		99	1	
05151100	**Low discrimination**						**05151100**
					GRA	SPO	
Clostridium septicum		55.34	0.41	NAG 99	99	99	
Fusobacterium mortiferum		18.88	0.51		1	1	
Lactobacillus catenaformis		5.69	0.32		99	1	
05151120	**Very good identification**						**05151120**
					GRA	SPO	
Clostridium septicum		95.74	0.75		99	99	
Clostridium perfringens		2.41	0.55		99	84	
Prevotella bivia		1.33	0.37		1	1	
05151121	**Acceptable identification**						**05151121**
Clostridium septicum		93.90	0.41	URE 1			
Clostridium perfringens		2.36	0.20				
Clostridium chauvoei		1.76	0.18				
05151122	**Low discrimination**						**05151122**
					GRA	SPO	
Clostridium septicum		42.52	0.46	BGL 2	99	99	
Prevotella oralis		27.94	0.47		1	1	
Clostridium perfringens		24.64	0.49		99	84	
05151130	**Good identification**						**05151130**
					GRA	SPO	
Clostridium septicum		95.77	0.41	MLZ 1	99	99	
Clostridium perfringens		2.41	0.20		99	84	
Prevotella bivia		1.33	0.03		1	1	
05151160	**Low discrimination**						**05151160**
					GRA	SPO	
Clostridium septicum		66.93	0.41	SOR 1	99	99	
Clostridium perfringens		31.72	0.42		99	84	
Prevotella bivia		0.93	0.03		1	1	
05151162	**Low discrimination**						**05151162**
					GRA	SPO	
Clostridium perfringens		84.11	0.37	SUC 91	99	84	
Clostridium septicum		7.70	0.12	CEL 4	99	99	
Prevotella oralis		5.06	0.12		1	1	
05151320	**Acceptable identification**						**05151320**
					GRA	SPO	
Clostridium septicum		93.24	0.41	RHA 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella oralis		2.79	0.18		1	1		
Clostridium perfringens		2.34	0.20		99	84		
05151322	**Acceptable identification**							**05151322**
					GRA	SPO		
Prevotella oralis		93.04	0.40	SUC 92	1	1		
Clostridium septicum		3.34	0.12	RAF 92	99	99		
Clostridium perfringens		1.93	0.15		99	84		
05151400	**Low discrimination**							**05151400**
					COC			
Gemella morbillorum		88.61	0.45	CEL 1	99			
Lactobacillus fermentum		5.67	0.36	ARA 10	1			
Lactobacillus jensenii		2.77	0.19		1			
05151402	**Low discrimination**							**05151402**
					COC			
Lactobacillus fermentum		67.41	0.36	SUC 87	1			
Lactobacillus jensenii		21.95	0.16	CEL 1	1			
Gemella morbillorum		10.64	0.11		99			
05151500	**Low discrimination**							**05151500**
					COC	SPO		
Lactobacillus fermentum		76.91	0.40	SUC 87	1	1		
Clostridium septicum		11.67	0.07	CEL 1	1	99		
Gemella morbillorum		7.44	0.11		99	1		
05151502	**Good identification**							**05151502**
Lactobacillus fermentum		96.91	0.40	SUC 87				
Bifidobacterium adolescentis		3.09	0.09	CEL 1				
05151520	**Acceptable identification**							**05151520**
					SPO			
Clostridium septicum		93.92	0.41	ARA 1	99			
Clostridium perfringens		2.36	0.20		84			
Lactobacillus fermentum		2.08	0.32		1			
05152010	**Good identification**							**05152010**
					GRA	SPO	CAT	
Clostridium difficile		97.73	0.36	MLT 1	99	99	1	
Propionibacterium granulosum		0.94	0.14	CEL 5	99	1	90	
Prevotella disiens		0.86	-.03		1	1	1	
05153120	**Good identification**							**05153120**
					GRA	SPO		
Clostridium septicum		95.48	0.41	MAN 1	99	99		
Clostridium perfringens		2.40	0.20		99	84		
Prevotella bivia		1.32	0.03		1	1		
05154000	**Low discrimination**							**05154000**
					COC	GRA		
Gemella morbillorum		75.49	0.50	CEL 1	99	99		
Prevotella disiens		21.94	0.32	XYL 10	1	1		
Prevotella intermedia		1.13	0.17		1	1		
05155000	**Low discrimination**							**05155000**
					COC			
Gemella morbillorum		89.11	0.45	CEL 1	99			
Lactobacillus fermentum		3.35	0.32	XYL 10	1			
Lactobacillus jensenii		2.79	0.19		1			
05155102	**Low discrimination**							**05155102**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Lactobacillus fermentum		47.75	0.36	SUC 87	1	99	
Atopobium parvulum		32.82	0.20	CEL 1	99	99	
Prevotella oralis		10.55	0.09		1	1	
05155120	**Acceptable identification**						**05155120**
					COC	SPO	
Clostridium septicum		91.65	0.41	XYL 1	1	99	
Atopobium parvulum		2.46	0.20		99	1	
Clostridium perfringens		2.30	0.20		1	84	
05155500	**Good identification**						**05155500**
Lactobacillus fermentum		95.66	0.36	SUC 87			
Actinomyces meyeri		4.34	0.02	CEL 1			
05155502	**Low discrimination**						**05155502**
Lactobacillus fermentum		70.83	0.36	SUC 87			
Bifidobacterium adolescentis		25.77	0.23	CEL 1			
Actinomyces israelii		3.40	0.14				
05160040	**Acceptable identification**						**05160040**
					GRA	SPO	
Clostridium botulinum biovar B		93.76	0.39	SUC 90	99	99	
Clostridium botulinum biovar A		5.73	0.21	CEL 1	99	99	
Prevotella disiens		0.51	-.03		1	1	
05161120	**Acceptable identification**						**05161120**
					GRA	SPO	
Clostridium septicum		94.17	0.49	FRU 99	99	99	
Prevotella bivia		3.92	0.19		1	1	
Clostridium perfringens		1.47	0.25		99	84	
05170040	**Good identification**						**05170040**
					GRA	SPO	
Clostridium botulinum biovar B		99.77	0.50	SUC 90	99	99	
Prevotella disiens		0.14	-.03	CEL 1	1	1	
Clostridium botulinum biovar A		0.10	0.01		99	99	
05170050	**Good identification**						**05170050**
Clostridium botulinum biovar B		100.0	0.44	SUC 90			
05170060	**Good identification**						**05170060**
Clostridium botulinum biovar B		100.0	0.44	SUC 90			
05170070	**Good identification**						**05170070**
Clostridium botulinum biovar B		100.0	0.39	SUC 90			
05170120	**Good identification**						**05170120**
					GRA	SPO	
Clostridium septicum		95.55	0.49	GAL 99	99	99	
Prevotella bivia		2.50	0.15		1	1	
Clostridium perfringens		1.49	0.25		99	84	
05171000	**Good identification**						**05171000**
					COC	CAT	
Lactobacillus jensenii		94.16	0.62	SUC 90	1	1	
Propionibacterium granulosum		1.76	0.39	SAL 95	1	90	
Atopobium parvulum		1.31	0.29		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05171002	**Very good identification**						**05171002**
					COC	CAT	
Lactobacillus jensenii		98.23	0.59	SUC 90	1	1	
Propionibacterium granulosum		0.69	0.29	SAL 95	1	90	
Atopobium parvulum		0.36	0.16		99	1	
05171010	**Good identification**						**05171010**
					CAT		
Lactobacillus jensenii		99.24	0.54	SUC 90	1		
Propionibacterium granulosum		0.76	0.24	SAL 95	90		
05171012	**Good identification**						**05171012**
					CAT		
Lactobacillus jensenii		99.71	0.51	SUC 90	1		
Propionibacterium granulosum		0.29	0.14	SAL 95	90		
05171020	**Low discrimination**						**05171020**
					SPO		
Clostridium septicum		58.14	0.49	LAC 99	99		
Lactobacillus jensenii		28.68	0.52		1		
Clostridium perfringens		12.21	0.44		84		
05171022	**Low discrimination**						**05171022**
					SPO		
Lactobacillus jensenii		71.79	0.49	SUC 90	1		
Clostridium perfringens		21.58	0.39	SAL 95	84		
Clostridium septicum		4.46	0.20		99		
05171030	**Acceptable identification**						**05171030**
					SPO		
Lactobacillus jensenii		92.42	0.43	SUC 90	1		
Clostridium septicum		5.68	0.15	SAL 95	99		
Clostridium perfringens		1.19	0.10		84		
05171032	**Good identification**						**05171032**
Lactobacillus jensenii		99.10	0.40	SUC 90			
Clostridium perfringens		0.90	0.05	SAL 95			
05171100	**Low discrimination**						**05171100**
					COC	SPO	
Atopobium parvulum		71.04	0.63	SUC 96	99	1	
Clostridium septicum		26.25	0.49		1	99	
Clostridium perfringens		1.25	0.33		1	84	
05171102	**Low discrimination**						**05171102**
					COC		
Atopobium parvulum		80.75	0.50	SUC 96	99		
Lactobacillus acidophilus		8.60	0.32		1		
Clostridium perfringens		3.79	0.28		1		
05171120	**Very good identification**						**05171120**
					COC	SPO	
Clostridium septicum		97.99	0.83		1	99	
Clostridium perfringens		1.53	0.59		1	84	
Atopobium parvulum		0.47	0.50		99	1	
05171121	**Good identification**						**05171121**
					COC	SPO	
Clostridium septicum		97.71	0.49	URE 1	1	99	
Clostridium perfringens		1.52	0.25		1	84	
Atopobium parvulum		0.47	0.16		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05171122	**Low discrimination**							**05171122**
					COC	SPO		
Clostridium septicum		69.21	0.55	BGL 2	1	99		
Clostridium perfringens		24.83	0.54		1	84		
Atopobium parvulum		2.89	0.37		99	1		
05171130	**Good identification**							**05171130**
					COC	SPO		
Clostridium septicum		97.84	0.49	MLZ 1	1	99		
Clostridium perfringens		1.52	0.25		1	84		
Atopobium parvulum		0.47	0.16		99	1		
05171160	**Low discrimination**							**05171160**
					COC	SPO		
Clostridium septicum		77.03	0.49	SOR 1	1	99		
Clostridium perfringens		22.60	0.47		1	84		
Atopobium parvulum		0.37	0.16		99	1		
05171162	**Low discrimination**							**05171162**
					SPO			
Clostridium perfringens		83.37	0.41	SUC 91	84			
Clostridium septicum		12.32	0.20	CEL 4	99			
Lactobacillus acidophilus		3.80	0.16		1			
05171320	**Good identification**							**05171320**
					COC	SPO		
Clostridium septicum		97.94	0.49	RHA 1	1	99		
Clostridium perfringens		1.52	0.25		1	84		
Atopobium parvulum		0.47	0.16		99	1		
05171520	**Good identification**							**05171520**
					COC	SPO		
Clostridium septicum		98.00	0.49	ARA 1	1	99		
Clostridium perfringens		1.53	0.25		1	84		
Atopobium parvulum		0.47	0.16		99	1		
05172000	**Low discrimination**							**05172000**
					GRA	SPO	CAT	
Propionibacterium granulosum		87.86	0.37	SUC 93	99	1	90	
Clostridium difficile		6.43	0.05	CEL 1	99	99	1	
Prevotella disiens		3.66	-.03		1	1	1	
05173000	**Low discrimination**							**05173000**
					SPO	CAT		
Propionibacterium granulosum		50.12	0.37	SUC 93	1	90		
Lactobacillus jensenii		35.94	0.28	CEL 1	1	1		
Clostridium innocuum		11.99	0.17		99	1		
05173101	**Good identification**							**05173101**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05173103	**Good identification**							**05173103**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05173120	**Good identification**							**05173120**
					COC	SPO		
Clostridium septicum		97.33	0.49	MAN 1	1	99		
Clostridium perfringens		1.51	0.25		1	84		
Atopobium parvulum		0.47	0.16		99	1		
05173123	**Good identification**							**05173123**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		100.0	0.38	SUC 99				
05175100	**Very good identification**							**05175100**
					COC	GRA	SPO	
Atopobium parvulum		99.71	0.67	SUC 96	99	99	1	
Clostridium septicum		0.20	0.15		1	99	99	
Fusobacterium mortiferum		0.05	0.22		1	1	1	
05175102	**Good identification**							**05175102**
					COC			
Atopobium parvulum		99.68	0.54	SUC 96	99			
Actinomyces israelii		0.32	0.26		1			
05175120	**Low discrimination**							**05175120**
					COC	SPO		
Clostridium septicum		52.57	0.49	XYL 1	1	99		
Atopobium parvulum		46.62	0.54		99	1		
Clostridium perfringens		0.82	0.25		1	84		
05175122	**Low discrimination**							**05175122**
					COC	GRA	SPO	
Atopobium parvulum		75.66	0.42	SUC 96	99	99	1	
Clostridium septicum		9.87	0.20		1	99	99	
Parabacteroides distasonis		9.25	0.36		1	1	1	
05175322	**Acceptable identification**							**05175322**
					COC	GRA		
Parabacteroides distasonis		94.91	0.44	RAF 99	1	1		
Atopobium parvulum		2.62	0.07		99	99		
Prevotella oralis		2.48	0.04		1	1		
05175332	**Good identification**							**05175332**
Parabacteroides distasonis		100.0	0.39	RAF 99				
05175722	**Good identification**							**05175722**
Parabacteroides distasonis		96.39	0.40	RAF 99				
Clostridium clostridioforme		3.61	0.10					
05175732	**Low discrimination**							**05175732**
Parabacteroides distasonis		75.95	0.36	RAF 99				
Clostridium clostridioforme		24.05	0.22					
05177102	**Low discrimination**							**05177102**
					COC	SPO		
Actinomyces israelii		58.06	0.39	SUC 98	1	1		
Atopobium parvulum		31.89	0.20	SAL 92	99	1		
Clostridium tertium		10.05	0.11		1	99		
05177502	**Low discrimination**							**05177502**
Actinomyces israelii		83.77	0.37	SUC 98				
Bifidobacterium adolescentis		16.23	0.19	SAL 92				
05200000	**Low discrimination**							**05200000**
					COC	SPO		
Clostridium bifermentans		35.89	0.55	IND 90	1	99		
Atopobium minutum		14.87	0.56	SAL 10	1	1		
Gemella morbillorum		14.55	0.52		99	1		
05200002	**Low discrimination**							**05200002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Atopobium minutum		86.27	0.53	SAL 1	1		
Clostridium novyi biovar A		4.05	0.35		99		
Clostridium bifermentans		3.16	0.21		99		
05200010	**Low discrimination**						**05200010**
Clostridium novyi biovar A		45.40	0.48	MNS 1			
Clostridium difficile		38.44	0.32	SAL 8			
Clostridium bifermentans		6.25	0.21				
05200011	**Good identification**						**05200011**
					COC	SPO	
Clostridium novyi biovar A		97.92	0.48	MNS 1	1	99	
Anaerococcus prevotii		1.25	-.03	SAL 8	99	1	
Clostridium difficile		0.84	-.02		1	99	
05200020	**Low discrimination**						**05200020**
					SPO		
Atopobium minutum		38.69	0.48	SAL 1	1		
Clostridium cadaveris		22.66	0.34		99		
Clostridium bifermentans		17.88	0.34		99		
05200021	**Low discrimination**						**05200021**
					COC	SPO	
Clostridium novyi biovar A		81.21	0.37	MNS 1	1	99	
Atopobium minutum		6.17	0.14	SAL 8	1	1	
Anaerococcus prevotii		4.14	-.03		99	1	
05200022	**Acceptable identification**						**05200022**
					GRA	SPO	
Atopobium minutum		93.73	0.45	SAL 1	99	1	
Clostridium novyi biovar A		3.30	0.25		99	99	
Pseudoflavonifractor capillosus		1.03	-.03		1	1	
05200030	**Low discrimination**						**05200030**
					SPO		
Clostridium novyi biovar A		82.33	0.37	MNS 1	99		
Atopobium minutum		6.26	0.14	SAL 8	1		
Clostridium cadaveris		3.67	0.00		99		
05200031	**Good identification**						**05200031**
Clostridium novyi biovar A		100.0	0.37	MNS 1			
05200041	**Good identification**						**05200041**
					COC	SPO	
Clostridium novyi biovar A		97.14	0.45	MNS 1	1	99	
Anaerococcus prevotii		1.86	-.03	SAL 8	99	1	
Clostridium botulinum biovar A		1.00	-.01		1	99	
05200050	**Acceptable identification**						**05200050**
Clostridium novyi biovar A		92.84	0.45	MNS 1			
Clostridium difficile		6.21	0.10	SAL 8			
Clostridium botulinum biovar A		0.96	-.01				
05200051	**Good identification**						**05200051**
Clostridium novyi biovar A		100.0	0.45	MNS 1			
05200100	**Low discrimination**						**05200100**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		49.82	0.42	FRU 99	1	1	
Clostridium bifermentans		17.63	0.21		99	99	
Atopobium minutum		7.31	0.22		99	1	
05200400	**Low discrimination**						**05200400**
					COC	SPO	
Gemella morbillorum		66.43	0.35	SAL 1	99	1	
Clostridium bifermentans		14.89	0.21	ARA 10	1	99	
Atopobium minutum		6.17	0.22		1	1	
05201000	**Low discrimination**						**05201000**
					COC	GRA	
Gemella morbillorum		54.28	0.47	SAL 1	99	99	
Atopobium minutum		34.33	0.48		1	99	
Fusobacterium mortiferum		4.56	0.39		1	1	
05201002	**Acceptable identification**						**05201002**
					COC	GRA	
Atopobium minutum		93.43	0.45	SAL 1	1	99	
Gemella morbillorum		2.24	0.13		99	99	
Fusobacterium mortiferum		2.07	0.22		1	1	
05201020	**Low discrimination**						**05201020**
					COC	SPO	
Atopobium minutum		61.96	0.40	SAL 1	1	1	
Gemella morbillorum		32.65	0.31		99	1	
Clostridium novyi biovar A		2.74	0.21		1	99	
05201022	**Good identification**						**05201022**
Atopobium minutum		100.0	0.37	SAL 1			
05201100	**Acceptable identification**						**05201100**
					COC	GRA	
Fusobacterium mortiferum		92.24	0.45	FRU 99	1	1	
Gemella morbillorum		4.75	0.13		99	99	
Atopobium minutum		3.01	0.14		1	99	
05202000	**Low discrimination**						**05202000**
					SPO		
Clostridium difficile		58.04	0.37	FRU 99	99		
Eubacterium limosum		25.72	0.29		1		
Clostridium bifermentans		5.12	0.21		99		
05202010	**Good identification**						**05202010**
					SPO		
Clostridium difficile		99.78	0.48	FRU 99	99		
Clostridium novyi biovar A		0.13	0.14		99		
Eubacterium limosum		0.09	-.06		1		
05203000	**Acceptable identification**						**05203000**
					COC	SPO	
Clostridium innocuum		93.67	0.37	FRU 95	1	99	
Gemella morbillorum		2.65	0.13	CEL 99	99	1	
Atopobium minutum		1.68	0.14		1	1	
05203002	**Good identification**						**05203002**
					SPO		
Clostridium innocuum		98.82	0.37	FRU 95	99		
Atopobium minutum		1.18	0.11	CEL 99	1		
05204000	**Low discrimination**						**05204000**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		63.55	0.35	SAL 1	99	1	
Clostridium bifermentans		14.25	0.21	XYL 10	1	99	
Atopobium minutum		5.90	0.22		1	1	
05210000	**Low discrimination**						**05210000**
					GRA	SPO	
Fusobacterium varium		43.15	0.58	SAL 1	1	1	
Clostridium difficile		15.38	0.54		99	99	
Fusobacterium mortiferum		14.78	0.70		1	1	
05210002	**Low discrimination**						**05210002**
					GRA		
Atopobium minutum		53.75	0.53	SAL 1	99		
Fusobacterium mortiferum		28.37	0.54		1		
Fusobacterium varium		7.53	0.24		1		
05210010	**Very good identification**						**05210010**
					GRA	SPO	
Clostridium difficile		98.71	0.66	MAN 90	99	99	
Fusobacterium varium		0.57	0.24		1	1	
Fusobacterium mortiferum		0.20	0.36		1	1	
05210020	**Low discrimination**						**05210020**
					COC	SPO	
Atopobium minutum		30.84	0.48	SAL 1	1	1	
Clostridium cadaveris		19.56	0.34		1	99	
Gemella morbillorum		13.32	0.37		99	1	
05210022	**Acceptable identification**						**05210022**
					GRA		
Atopobium minutum		92.87	0.45	SAL 1	99		
Pseudoramibacter alactolyticus		2.12	0.03		99		
Fusobacterium mortiferum		1.49	0.19		1		
05210050	**Good identification**						**05210050**
Clostridium difficile		98.02	0.44	MAN 90			
Clostridium novyi biovar A		1.65	0.28	SOR 5			
Clostridium botulinum biovar B		0.34	0.02				
05210100	**Very good identification**						**05210100**
					GRA	SPO	
Fusobacterium mortiferum		97.57	0.76		1	1	
Fusobacterium varium		1.23	0.24		1	1	
Clostridium difficile		0.44	0.20		99	99	
05210101	**Good identification**						**05210101**
Fusobacterium mortiferum		100.0	0.42	URE 1			
05210102	**Very good identification**						**05210102**
					GRA		
Fusobacterium mortiferum		99.19	0.60	BGL 10	1		
Atopobium minutum		0.81	0.19		99		
05210120	**Low discrimination**						**05210120**
					GRA	SPO	
Fusobacterium mortiferum		38.74	0.42	NAG 1	1	1	
Clostridium septicum		28.85	0.22		99	99	
Prevotella bivia		28.11	0.15		1	1	
05210140	**Good identification**						**05210140**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		97.71	0.42	SOR 1	1	1		
Clostridium difficile		2.29	-.02		99	99		
05210300	**Good identification**							**05210300**
Fusobacterium mortiferum		100.0	0.42	RHA 1				
05210400	**Low discrimination**							**05210400**
					COC	GRA	SPO	
Gemella morbillorum		41.65	0.37	SAL 1	99	99	1	
Fusobacterium varium		27.01	0.24	ARA 10	1	1	1	
Clostridium difficile		9.63	0.20		1	99	99	
05210500	**Good identification**							**05210500**
					COC	GRA		
Fusobacterium mortiferum		98.09	0.42	ARA 1	1	1		
Gemella morbillorum		1.91	0.03		99	99		
05211000	**Low discrimination**							**05211000**
					COC	GRA		
Fusobacterium mortiferum		77.38	0.73		1	1		
Gemella morbillorum		11.37	0.48		99	99		
Atopobium minutum		5.43	0.48		1	99		
05211001	**Low discrimination**							**05211001**
					COC	GRA		
Fusobacterium mortiferum		78.96	0.39	URE 1	1	1		
Gemella morbillorum		11.60	0.14		99	99		
Atopobium minutum		5.54	0.14		1	99		
05211002	**Low discrimination**							**05211002**
					GRA			
Fusobacterium mortiferum		60.48	0.57	BGL 10	1			
Atopobium minutum		25.47	0.45		99			
Lactobacillus jensenii		5.47	0.29		99			
05211022	**Low discrimination**							**05211022**
Atopobium minutum		67.41	0.37	SAL 1				
Lactobacillus jensenii		10.86	0.18					
Clostridium perfringens		6.07	0.13					
05211040	**Low discrimination**							**05211040**
					COC	GRA		
Fusobacterium mortiferum		77.73	0.39	SOR 1	1	1		
Gemella morbillorum		11.42	0.14		99	99		
Atopobium minutum		5.46	0.14		1	99		
05211100	**Very good identification**							**05211100**
					GRA	SPO		
Fusobacterium mortiferum		99.15	0.79		1	1		
Clostridium septicum		0.50	0.22		99	99		
Lactobacillus fermentum		0.14	0.32		99	1		
05211101	**Good identification**							**05211101**
Fusobacterium mortiferum		100.0	0.45	URE 1				
05211102	**Very good identification**							**05211102**
					GRA			
Fusobacterium mortiferum		97.50	0.63	BGL 10	1			
Lactobacillus fermentum		1.21	0.32		99			
Lactobacillus catenaformis		0.62	0.15		99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
05211110	**Good identification**						**05211110**
					GRA	SPO	
Fusobacterium mortiferum		98.56	0.45	MLZ 1	1	1	
Clostridium difficile		1.45	-.02		99	99	
05211120	**Very good identification**						**05211120**
					GRA	SPO	
Clostridium septicum		95.88	0.57	MLT 94	99	99	
Fusobacterium mortiferum		1.95	0.45		1	1	
Clostridium perfringens		1.81	0.34		99	84	
05211140	**Good identification**						**05211140**
Fusobacterium mortiferum		100.0	0.45	SOR 1			
05211200	**Low discrimination**						**05211200**
					COC	GRA	
Fusobacterium mortiferum		79.77	0.39	RHA 1	1	1	
Gemella morbillorum		11.72	0.14		99	99	
Atopobium minutum		5.60	0.14		1	99	
05211300	**Good identification**						**05211300**
Fusobacterium mortiferum		100.0	0.45	RHA 1			
05211500	**Low discrimination**						**05211500**
					GRA		
Fusobacterium mortiferum		87.98	0.45	ARA 1	1		
Lactobacillus fermentum		12.02	0.32		99		
05212000	**Low discrimination**						**05212000**
					SPO		
Clostridium difficile		68.38	0.71		99		
Eubacterium limosum		30.31	0.63		1		
Clostridium innocuum		0.47	0.37		99		
05212001	**Low discrimination**						**05212001**
					SPO		
Clostridium difficile		68.74	0.37	URE 1	99		
Eubacterium limosum		30.47	0.29		1		
Clostridium innocuum		0.47	0.03		99		
05212002	**Low discrimination**						**05212002**
					SPO		
Clostridium difficile		44.34	0.37	BGL 1	99		
Clostridium innocuum		30.17	0.37		99		
Eubacterium limosum		19.65	0.29		1		
05212010	**Very good identification**						**05212010**
					SPO		
Clostridium difficile		99.90	0.82		99		
Eubacterium limosum		0.09	0.29		1		
Clostridium innocuum		0.01	0.13		99		
05212011	**Good identification**						**05212011**
					SPO		
Clostridium difficile		99.91	0.48	URE 1	99		
Eubacterium limosum		0.09	-.06		1		
05212012	**Good identification**						**05212012**
					SPO	CAT	
Clostridium difficile		99.23	0.48	BGL 1	99	1	
Clostridium innocuum		0.58	0.13		99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		0.10	0.09		1	90	
05212020	**Low discrimination**						**05212020**
					SPO	CAT	
Clostridium difficile		54.45	0.37	NAG 1	99	1	
Eubacterium limosum		24.14	0.29		1	1	
Propionibacterium acnes		13.37	0.29		1	95	
05212030	**Good identification**						**05212030**
					SPO	CAT	
Clostridium difficile		99.81	0.48	NAG 1	99	1	
Propionibacterium granulosum		0.10	0.09		1	90	
Eubacterium limosum		0.09	-.06		1	1	
05212040	**Low discrimination**						**05212040**
					SPO		
Clostridium difficile		89.26	0.49	SOR 5	99		
Eubacterium limosum		7.59	0.29		1		
Pseudoramibacter alactolyticus		1.99	0.17		1		
05212050	**Very good identification**						**05212050**
					SPO		
Clostridium difficile		99.98	0.61	SOR 5	99		
Eubacterium limosum		0.02	-.06		1		
05212100	**Low discrimination**						**05212100**
					GRA	SPO	
Clostridium difficile		58.69	0.37	LAC 1	99	99	
Eubacterium limosum		26.01	0.29		99	1	
Fusobacterium mortiferum		14.62	0.42		1	1	
05212110	**Good identification**						**05212110**
					SPO		
Clostridium difficile		99.91	0.48	LAC 1	99		
Eubacterium limosum		0.09	-.06		1		
05212200	**Low discrimination**						**05212200**
					SPO		
Clostridium difficile		68.53	0.37	RHA 1	99		
Eubacterium limosum		30.38	0.29		1		
Clostridium innocuum		0.47	0.03		99		
05212210	**Good identification**						**05212210**
					SPO		
Clostridium difficile		99.91	0.48	RHA 1	99		
Eubacterium limosum		0.09	-.06		1		
05212400	**Low discrimination**						**05212400**
					SPO		
Clostridium difficile		63.58	0.37	ARA 1	99		
Eubacterium limosum		28.18	0.29		1		
Clostridium innocuum		7.64	0.24		99		
05212410	**Good identification**						**05212410**
					SPO		
Clostridium difficile		99.81	0.48	ARA 1	99		
Clostridium innocuum		0.10	0.00		99		
Eubacterium limosum		0.09	-.06		1		
05213000	**Low discrimination**						**05213000**
					SPO		
Clostridium innocuum		87.87	0.59	CEL 99	99		
Clostridium difficile		6.80	0.37		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium limosum		3.01	0.29		1			
05213002	**Very good identification**							**05213002**
					GRA	SPO	CAT	
Clostridium innocuum		99.42	0.59	CEL 99	99	99	1	
Propionibacterium granulosum		0.27	0.24		99	1	90	
Fusobacterium mortiferum		0.14	0.22		1	1	5	
05213010	**Low discrimination**							**05213010**
					SPO	CAT		
Clostridium difficile		89.67	0.48	GAL 1	99	1		
Clostridium innocuum		9.89	0.35		99	1		
Propionibacterium granulosum		0.36	0.20		1	90		
05213012	**Acceptable identification**							**05213012**
					SPO	CAT		
Clostridium innocuum		90.86	0.35	CEL 99	99	1		
Clostridium difficile		8.32	0.14	MLZ 4	99	1		
Propionibacterium granulosum		0.82	0.09		1	90		
05213100	**Low discrimination**							**05213100**
					GRA	SPO		
Fusobacterium mortiferum		72.02	0.45	MAN 1	1	1		
Clostridium innocuum		25.17	0.25		99	99		
Clostridium difficile		1.95	0.02		99	99		
05213400	**Good identification**							**05213400**
					SPO			
Clostridium innocuum		99.37	0.46	CEL 99	99			
Clostridium difficile		0.44	0.02		99			
Eubacterium limosum		0.20	-.06		1			
05213402	**Good identification**							**05213402**
Clostridium innocuum		100.0	0.46	CEL 99				
05214010	**Good identification**							**05214010**
					GRA	SPO		
Clostridium difficile		99.09	0.44	MAN 90	99	99		
Fusobacterium varium		0.46	0.01	XYL 5	1	1		
Fusobacterium mortiferum		0.28	0.17		1	1		
05214100	**Very good identification**							**05214100**
					GRA			
Fusobacterium mortiferum		98.24	0.57	XYL 7	1			
Fusobacterium varium		0.69	0.01		1			
Actinomyces meyeri		0.51	0.09		99			
05214102	**Good identification**							**05214102**
Fusobacterium mortiferum		100.0	0.41	XYL 7				
05215000	**Low discrimination**							**05215000**
					COC	GRA		
Fusobacterium mortiferum		79.16	0.54	XYL 7	1	1		
Gemella morbillorum		17.17	0.32		99	99		
Lactobacillus fermentum		1.22	0.24		1	99		
05215002	**Low discrimination**							**05215002**
					GRA			
Fusobacterium mortiferum		80.19	0.37	XYL 7	1			
Lactobacillus fermentum		11.12	0.24	BGL 10	99			
Atopobium minutum		4.53	0.11		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05215100	**Very good identification**							**05215100**
					GRA			
Fusobacterium mortiferum		98.43	0.60	XYL 7	1			
Lactobacillus fermentum		1.06	0.28		99			
Actinomyces meyeri		0.51	0.12		99			
05215102	**Acceptable identification**							**05215102**
					GRA			
Fusobacterium mortiferum		91.16	0.44	XYL 7	1			
Lactobacillus fermentum		8.84	0.28	BGL 10	99			
05216000	**Acceptable identification**							**05216000**
					SPO			
Clostridium difficile		91.09	0.49	XYL 5	99			
Eubacterium limosum		7.75	0.29		1			
Clostridium innocuum		0.63	0.15		99			
05216010	**Very good identification**							**05216010**
					SPO			
Clostridium difficile		99.98	0.61	XYL 5	99			
Eubacterium limosum		0.02	-.06		1			
05216050	**Good identification**							**05216050**
Clostridium difficile		100.0	0.39	XYL 5				
05217000	**Acceptable identification**							**05217000**
					GRA	SPO		
Clostridium innocuum		90.77	0.37	CEL 99	99	99		
Clostridium difficile		7.02	0.15	XYL 5	99	99		
Fusobacterium mortiferum		1.61	0.20		1	1		
05217002	**Good identification**							**05217002**
Clostridium innocuum		100.0	0.37	CEL 99				
05221100	**Good identification**							**05221100**
					GRA	SPO		
Fusobacterium mortiferum		95.20	0.36	FRU 99	1	1		
Clostridium septicum		4.80	-.04		99	99		
05230000	**Low discrimination**							**05230000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		62.14	0.61		1	1	5	
Clostridium difficile		16.30	0.35		99	99	1	
Propionibacterium granulosum		11.14	0.45		99	1	90	
05230002	**Low discrimination**							**05230002**
					GRA	CAT		
Fusobacterium mortiferum		66.44	0.45	BGL 10	1	5		
Propionibacterium granulosum		26.80	0.35		99	90		
Atopobium minutum		4.26	0.19		99	1		
05230010	**Good identification**							**05230010**
					GRA	SPO	CAT	
Clostridium difficile		97.37	0.47	TRE 7	99	99	1	
Propionibacterium granulosum		1.86	0.30	MAN 90	99	1	90	
Fusobacterium mortiferum		0.77	0.27		1	1	5	
05230100	**Very good identification**							**05230100**
					GRA	SPO	CAT	
Fusobacterium mortiferum		99.73	0.67		1	1	5	
Clostridium difficile		0.11	0.01		99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.08	0.11		99	1	90	
05230102	**Good identification**							**05230102**
Fusobacterium mortiferum		100.0	0.51	BGL 10				
05231000	**Low discrimination**							**05231000**
					GRA	CAT		
Fusobacterium mortiferum		77.47	0.64		1	5		
Lactobacillus jensenii		11.73	0.40		99	1		
Propionibacterium granulosum		9.26	0.45		99	90		
05231002	**Low discrimination**							**05231002**
					GRA	CAT		
Fusobacterium mortiferum		43.89	0.48	BGL 10	1	5		
Lactobacillus jensenii		39.86	0.37		99	1		
Propionibacterium granulosum		11.80	0.35		99	90		
05231100	**Good identification**							**05231100**
					COC	GRA	SPO	
Fusobacterium mortiferum		94.07	0.70		1	1	1	
Clostridium septicum		4.74	0.31		1	99	99	
Atopobium parvulum		0.91	0.24		99	99	1	
05231101	**Good identification**							**05231101**
					GRA	SPO		
Fusobacterium mortiferum		95.20	0.36	URE 1	1	1		
Clostridium septicum		4.80	-.04		99	99		
05231102	**Acceptable identification**							**05231102**
					COC	GRA		
Fusobacterium mortiferum		94.83	0.54	BGL 10	1	1		
Lactobacillus acidophilus		1.73	0.11		1	99		
Atopobium parvulum		1.46	0.11		99	99		
05231110	**Good identification**							**05231110**
					GRA	SPO		
Fusobacterium mortiferum		95.20	0.36	MLZ 1	1	1		
Clostridium septicum		4.80	-.04		99	99		
05231120	**Very good identification**							**05231120**
					GRA	SPO		
Clostridium septicum		98.61	0.65	MLT 94	99	99		
Clostridium perfringens		1.15	0.38		99	84		
Fusobacterium mortiferum		0.20	0.36		1	1		
05231122	**Low discrimination**							**05231122**
					SPO			
Clostridium septicum		76.45	0.36	MLT 94	99			
Clostridium perfringens		20.62	0.33	BGL 2	84			
Lactobacillus acidophilus		1.52	0.11		1			
05231140	**Acceptable identification**							**05231140**
					GRA	SPO		
Fusobacterium mortiferum		92.22	0.36	SOR 1	1	1		
Clostridium septicum		4.65	-.04		99	99		
Clostridium perfringens		3.14	0.00		99	84		
05231300	**Good identification**							**05231300**
					GRA	SPO		
Fusobacterium mortiferum		95.20	0.36	RHA 1	1	1		
Clostridium septicum		4.80	-.04		99	99		
05231500	**Good identification**							**05231500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Fusobacterium mortiferum		95.20	0.36	ARA 1	1	1		
Clostridium septicum		4.80	-.04		99	99		
05232000	**Low discrimination**							**05232000**
					SPO	CAT		
Clostridium difficile		86.67	0.51	TRE 7	99	1		
Eubacterium limosum		5.16	0.29		1	1		
Propionibacterium granulosum		4.97	0.43		1	90		
05232010	**Very good identification**							**05232010**
					SPO	CAT		
Clostridium difficile		99.80	0.63	TRE 7	99	1		
Propionibacterium granulosum		0.16	0.28		1	90		
Clostridium innocuum		0.03	0.06		99	1		
05232050	**Good identification**							**05232050**
Clostridium difficile		100.0	0.41	TRE 7				
05233000	**Low discrimination**							**05233000**
					SPO	CAT		
Clostridium innocuum		88.99	0.51	CEL 99	99	1		
Propionibacterium granulosum		8.37	0.43		1	90		
Clostridium difficile		1.47	0.17		99	1		
05233002	**Good identification**							**05233002**
					GRA	SPO	CAT	
Clostridium innocuum		97.49	0.51	CEL 99	99	99	1	
Propionibacterium granulosum		2.29	0.32		99	1	90	
Fusobacterium mortiferum		0.11	0.13		1	1	5	
05233100	**Low discrimination**							**05233100**
					GRA	SPO		
Fusobacterium mortiferum		66.66	0.36	MAN 1	1	1		
Clostridium innocuum		27.41	0.17		99	99		
Clostridium septicum		3.36	-.04		99	99		
05233400	**Good identification**							**05233400**
					SPO	CAT		
Clostridium innocuum		99.47	0.38	CEL 99	99	1		
Propionibacterium granulosum		0.54	0.09		1	90		
05233402	**Good identification**							**05233402**
Clostridium innocuum		100.0	0.38	CEL 99				
05234000	**Low discrimination**							**05234000**
					GRA	SPO		
Fusobacterium mortiferum		77.97	0.42	XYL 7	1	1		
Clostridium difficile		14.30	0.13		99	99		
Fusobacterium varium		4.26	0.01		1	1		
05234100	**Good identification**							**05234100**
Fusobacterium mortiferum		100.0	0.48	XYL 7				
05235000	**Good identification**							**05235000**
					GRA	CAT		
Fusobacterium mortiferum		96.49	0.45	XYL 7	1	5		
Lactobacillus jensenii		1.96	0.06		99	1		
Propionibacterium granulosum		1.55	0.11		99	90		
05235100	**Low discrimination**							**05235100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Fusobacterium mortiferum		80.25	0.51	XYL 7	1	1	1	
Atopobium parvulum		19.21	0.29		99	99	1	
Clostridium septicum		0.54	-.04		1	99	99	
05235322	**Good identification**							**05235322**
Parabacteroides distasonis		100.0	0.40	RAF 99				
05235332	**Good identification**							**05235332**
Parabacteroides distasonis		100.0	0.35	RAF 99				
05235722	**Good identification**							**05235722**
Parabacteroides distasonis		100.0	0.36	RAF 99				
05236010	**Good identification**							**05236010**
Clostridium difficile		100.0	0.41	TRE 7				
05241000	**Low discrimination**							**05241000**
					GRA	SPO		
Clostridium innocuum		82.12	0.37	FRU 95	99	99		
Fusobacterium mortiferum		6.44	0.31	MAN 99	1	1		
Lactobacillus jensenii		5.53	0.19		99	1		
05241002	**Acceptable identification**							**05241002**
					SPO			
Clostridium innocuum		91.63	0.37	FRU 95	99			
Lactobacillus jensenii		4.12	0.16	MAN 99	1			
Lactobacillus catenaformis		2.36	0.12		1			
05241100	**Low discrimination**							**05241100**
					GRA	SPO		
Fusobacterium mortiferum		76.11	0.37	FRU 99	1	1		
Lactobacillus catenaformis		16.07	0.15		99	1		
Clostridium innocuum		4.20	0.03		99	99		
05242000	**Good identification**							**05242000**
					SPO			
Clostridium innocuum		98.42	0.49	FRU 95	99			
Clostridium difficile		1.46	0.15	GAL 95	99			
Eubacterium limosum		0.12	-.06		1			
05242002	**Good identification**							**05242002**
Clostridium innocuum		100.0	0.49	FRU 95				
05242400	**Good identification**							**05242400**
Clostridium innocuum		100.0	0.36	FRU 95				
05242402	**Good identification**							**05242402**
Clostridium innocuum		100.0	0.36	FRU 95				
05243000	**Very good identification**							**05243000**
Clostridium innocuum		100.0	0.71	FRU 95				
05243001	**Good identification**							**05243001**
Clostridium innocuum		100.0	0.37	FRU 95				
05243002	**Very good identification**							**05243002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium innocuum		100.0	0.71	FRU 95		
05243003	**Good identification**					**05243003**
Clostridium innocuum		100.0	0.37	FRU 95		
05243010	**Good identification**					**05243010**
Clostridium innocuum		100.0	0.48	FRU 95		
05243012	**Good identification**					**05243012**
Clostridium innocuum		100.0	0.48	FRU 95		
05243020	**Good identification**					**05243020**
Clostridium innocuum		100.0	0.37	FRU 95		
05243022	**Good identification**					**05243022**
Clostridium innocuum		100.0	0.37	FRU 95		
05243040	**Good identification**					**05243040**
Clostridium innocuum		100.0	0.37	FRU 95		
05243042	**Good identification**					**05243042**
Clostridium innocuum		100.0	0.37	FRU 95		
05243100	**Good identification**					**05243100**
Clostridium innocuum		100.0	0.37	FRU 95		
05243102	**Good identification**					**05243102**
Clostridium innocuum		100.0	0.37	FRU 95		
05243200	**Good identification**					**05243200**
Clostridium innocuum		100.0	0.37	FRU 95		
05243202	**Good identification**					**05243202**
Clostridium innocuum		100.0	0.37	FRU 95		
05243400	**Very good identification**					**05243400**
Clostridium innocuum		100.0	0.58	FRU 95		
05243402	**Very good identification**					**05243402**
Clostridium innocuum		100.0	0.58	FRU 95		
05247000	**Good identification**					**05247000**
Clostridium innocuum		100.0	0.49	FRU 95		
05247002	**Good identification**					**05247002**
Clostridium innocuum		100.0	0.49	FRU 95		
05247400	**Good identification**					**05247400**
Clostridium innocuum		100.0	0.36	FRU 95		
05247402	**Good identification**					**05247402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.36	FRU 95			
05250000	**Low discrimination**						**05250000**
					GRA	SPO	
Fusobacterium mortiferum		65.91	0.62		1	1	
Clostridium innocuum		12.73	0.37		99	99	
Clostridium difficile		10.83	0.32		99	99	
05250002	**Low discrimination**						**05250002**
					GRA	SPO	
Clostridium innocuum		58.83	0.37	GAL 95	99	99	
Fusobacterium mortiferum		33.84	0.45	MAN 99	1	1	
Lactobacillus jensenii		2.64	0.16		99	1	
05250010	**Good identification**						**05250010**
					GRA	SPO	
Clostridium difficile		97.27	0.44	CEL 5	99	99	
Fusobacterium mortiferum		1.23	0.28	MAN 90	1	1	
Clostridium innocuum		0.98	0.13		99	99	
05250100	**Very good identification**						**05250100**
					GRA	SPO	
Fusobacterium mortiferum		99.35	0.68		1	1	
Lactobacillus catenaformis		0.32	0.15		99	1	
Clostridium barati		0.10	0.02		99	99	
05250102	**Acceptable identification**						**05250102**
					GRA	SPO	
Fusobacterium mortiferum		93.33	0.52	BGL 10	1	1	
Lactobacillus catenaformis		2.69	0.15		99	1	
Clostridium barati		2.59	0.10		99	99	
05251000	**Low discrimination**						**05251000**
					GRA	SPO	
Clostridium innocuum		52.58	0.59	MAN 99	99	99	
Fusobacterium mortiferum		21.49	0.65		1	1	
Lactobacillus jensenii		18.46	0.54		99	1	
05251002	**Low discrimination**						**05251002**
					SPO		
Clostridium innocuum		69.87	0.59	MAN 99	99		
Lactobacillus jensenii		16.36	0.51		1		
Lactobacillus catenaformis		9.38	0.47		1		
05251010	**Low discrimination**						**05251010**
					GRA	SPO	
Lactobacillus jensenii		70.34	0.45	SUC 90	99	1	
Clostridium innocuum		25.04	0.35	MLT 95	99	99	
Fusobacterium mortiferum		2.48	0.31		1	1	
05251012	**Low discrimination**						**05251012**
					SPO		
Lactobacillus jensenii		64.21	0.42	SUC 90	1		
Clostridium innocuum		34.29	0.35	MLT 95	99		
Lactobacillus catenaformis		1.12	0.12		1		
05251020	**Low discrimination**						**05251020**
					SPO		
Lactobacillus jensenii		45.23	0.43	SUC 90	1		
Clostridium septicum		23.41	0.31	MLT 95	99		
Lactobacillus catenaformis		17.29	0.36		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05251022	**Low discrimination**						**05251022**
					SPO		
Lactobacillus jensenii		40.27	0.40	SUC 90	1		
Clostridium barati		25.68	0.32	MLT 95	99		
Lactobacillus catenaformis		23.09	0.36		1		
05251030	**Good identification**						**05251030**
					SPO		
Lactobacillus jensenii		96.00	0.35	SUC 90	1		
Clostridium septicum		1.51	-.03	MLT 95	99		
Clostridium innocuum		1.38	0.01		99		
05251100	**Low discrimination**						**05251100**
					GRA	SPO	
Fusobacterium mortiferum		77.64	0.71		1	1	
Lactobacillus catenaformis		16.39	0.50		99	1	
Clostridium septicum		3.70	0.31		99	99	
05251101	**Low discrimination**						**05251101**
					GRA	SPO	
Fusobacterium mortiferum		79.45	0.37	URE 1	1	1	
Lactobacillus catenaformis		16.77	0.15		99	1	
Clostridium septicum		3.79	-.03		99	99	
05251102	**Low discrimination**						**05251102**
					GRA		
Lactobacillus catenaformis		49.19	0.50	SUC 99	99		
Fusobacterium mortiferum		25.89	0.55	MLT 90	1		
Lactobacillus acidophilus		12.53	0.36		99		
05251110	**Low discrimination**						**05251110**
					GRA		
Fusobacterium mortiferum		70.17	0.37	MLZ 1	1		
Lactobacillus catenaformis		14.81	0.15		99		
Lactobacillus jensenii		8.61	0.11		99		
05251120	**Very good identification**						**05251120**
					SPO		
Clostridium septicum		97.69	0.65	MLT 94	99		
Lactobacillus catenaformis		1.09	0.39		1		
Clostridium barati		0.81	0.32		99		
05251122	**Low discrimination**						**05251122**
					SPO		
Clostridium barati		35.59	0.40	ESL 87	99		
Clostridium septicum		29.19	0.36	SUC 99	99		
Lactobacillus acidophilus		16.30	0.36		1		
05251140	**Low discrimination**						**05251140**
					GRA	SPO	
Fusobacterium mortiferum		72.94	0.37	SOR 1	1	1	
Lactobacillus catenaformis		15.40	0.15		99	1	
Clostridium barati		8.19	0.05		99	99	
05251300	**Low discrimination**						**05251300**
					GRA	SPO	
Fusobacterium mortiferum		72.94	0.37	RHA 1	1	1	
Lactobacillus catenaformis		15.40	0.15		99	1	
Clostridium barati		8.19	0.05		99	99	
05251400	**Good identification**						**05251400**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		95.13	0.46	MAN 99	99	99	
Fusobacterium mortiferum		2.23	0.31		1	1	
Lactobacillus jensenii		1.91	0.19		99	1	
05251402	**Good identification**						**05251402**
					SPO		
Clostridium innocuum		97.69	0.46	MAN 99	99		
Lactobacillus jensenii		1.31	0.16		1		
Lactobacillus catenaformis		0.75	0.12		1		
05251500	**Low discrimination**						**05251500**
					GRA	SPO	
Fusobacterium mortiferum		69.26	0.37	ARA 1	1	1	
Lactobacillus catenaformis		14.62	0.15		99	1	
Clostridium innocuum		12.82	0.12		99	99	
05252000	**Good identification**						**05252000**
					SPO		
Clostridium innocuum		92.21	0.71	GAL 95	99		
Clostridium difficile		7.13	0.49		99		
Eubacterium limosum		0.61	0.29		1		
05252001	**Acceptable identification**						**05252001**
					SPO		
Clostridium innocuum		92.26	0.37	GAL 95	99		
Clostridium difficile		7.14	0.15	URE 1	99		
Eubacterium limosum		0.61	-.06		1		
05252002	**Very good identification**						**05252002**
					GRA	SPO	
Clostridium innocuum		99.91	0.71	GAL 95	99	99	
Clostridium difficile		0.08	0.15		99	99	
Fusobacterium mortiferum		0.01	0.11		1	1	
05252003	**Good identification**						**05252003**
Clostridium innocuum		100.0	0.37	GAL 95			
05252010	**Good identification**						**05252010**
					SPO		
Clostridium difficile		90.05	0.61	CEL 5	99		
Clostridium innocuum		9.94	0.48		99		
Eubacterium limosum		0.02	-.06		1		
05252012	**Acceptable identification**						**05252012**
Clostridium innocuum		91.61	0.48	GAL 95			
Clostridium difficile		8.39	0.26	MLZ 4			
05252020	**Acceptable identification**						**05252020**
					SPO		
Clostridium innocuum		92.26	0.37	GAL 95	99		
Clostridium difficile		7.14	0.15	NAG 1	99		
Eubacterium limosum		0.61	-.06		1		
05252022	**Good identification**						**05252022**
Clostridium innocuum		100.0	0.37	GAL 95			
05252040	**Low discrimination**						**05252040**
					SPO		
Clostridium innocuum		70.94	0.37	GAL 95	99		
Clostridium difficile		28.59	0.27	SOR 1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		0.47	-.06		1		
05252042	**Good identification**						**05252042**
Clostridium innocuum		100.0	0.37	GAL 95			
05252050	**Good identification**						**05252050**
Clostridium difficile		97.93	0.39	CEL 5			
Clostridium innocuum		2.07	0.13	SOR 5			
05252100	**Low discrimination**						**05252100**
					GRA	SPO	
Clostridium innocuum		82.92	0.37	GAL 95	99	99	
Fusobacterium mortiferum		10.12	0.34	LAC 1	1	1	
Clostridium difficile		6.41	0.15		99	99	
05252102	**Good identification**						**05252102**
					GRA	SPO	
Clostridium innocuum		98.66	0.37	GAL 95	99	99	
Fusobacterium mortiferum		1.34	0.18	LAC 1	1	1	
05252200	**Acceptable identification**						**05252200**
					SPO		
Clostridium innocuum		92.26	0.37	GAL 95	99		
Clostridium difficile		7.14	0.15	RHA 1	99		
Eubacterium limosum		0.61	-.06		1		
05252202	**Good identification**						**05252202**
Clostridium innocuum		100.0	0.37	GAL 95			
05252400	**Very good identification**						**05252400**
					SPO		
Clostridium innocuum		99.52	0.58	GAL 95	99		
Clostridium difficile		0.44	0.15		99		
Eubacterium limosum		0.04	-.06		1		
05252402	**Very good identification**						**05252402**
Clostridium innocuum		100.0	0.58	GAL 95			
05253000	**Excellent identification**						**05253000**
					GRA	SPO	
Clostridium innocuum		99.99	0.93		99	99	
Fusobacterium mortiferum		0.00	0.31		1	1	
Lactobacillus jensenii		0.00	0.19		99	1	
05253001	**Very good identification**						**05253001**
Clostridium innocuum		100.0	0.59	URE 1			
05253002	**Excellent identification**						**05253002**
					GRA	SPO	
Clostridium innocuum		100.0	0.93		99	99	
Lactobacillus jensenii		0.00	0.16		99	1	
Fusobacterium mortiferum		0.00	0.14		1	1	
05253003	**Very good identification**						**05253003**
Clostridium innocuum		100.0	0.59	URE 1			
05253010	**Very good identification**						**05253010**
					SPO		
Clostridium innocuum		99.49	0.69	MLZ 4	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		0.48	0.26		99		
Lactobacillus jensenii		0.03	0.11		1		
05253011	**Good identification**						**05253011**
Clostridium innocuum		100.0	0.35	MLZ 4			
05253012	**Very good identification**						**05253012**
					SPO		
Clostridium innocuum		99.98	0.69	MLZ 4	99		
Lactobacillus jensenii		0.02	0.08		1		
05253013	**Good identification**						**05253013**
Clostridium innocuum		100.0	0.35	MLZ 4			
05253020	**Very good identification**						**05253020**
					SPO		
Clostridium innocuum		99.72	0.59	NAG 1	99		
Clostridium barati		0.11	0.05		99		
Lactobacillus jensenii		0.09	0.09		1		
05253022	**Very good identification**						**05253022**
					SPO		
Clostridium innocuum		99.58	0.59	NAG 1	99		
Clostridium barati		0.32	0.14		99		
Lactobacillus jensenii		0.06	0.06		1		
05253030	**Good identification**						**05253030**
					SPO		
Clostridium innocuum		99.30	0.35	MLZ 4	99		
Lactobacillus jensenii		0.70	0.01	NAG 1	1		
05253032	**Good identification**						**05253032**
Clostridium innocuum		100.0	0.35	MLZ 4			
05253040	**Very good identification**						**05253040**
Clostridium innocuum		100.0	0.59	SOR 1			
05253042	**Very good identification**						**05253042**
Clostridium innocuum		100.0	0.59	SOR 1			
05253050	**Good identification**						**05253050**
Clostridium innocuum		97.55	0.35	MLZ 4			
Clostridium difficile		2.45	0.05	SOR 1			
05253052	**Good identification**						**05253052**
Clostridium innocuum		100.0	0.35	MLZ 4			
05253100	**Very good identification**						**05253100**
					GRA	SPO	
Clostridium innocuum		98.70	0.59	LAC 1	99	99	
Fusobacterium mortiferum		0.95	0.37		1	1	
Lactobacillus catenaformis		0.20	0.15		99	1	
05253102	**Very good identification**						**05253102**
					SPO		
Clostridium innocuum		99.10	0.59	LAC 1	99		
Clostridium barati		0.32	0.14		99		
Lactobacillus acidophilus		0.27	0.15		1		

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
05253110	**Good identification**					**05253110**
Clostridium innocuum		100.0	0.35	LAC 1		
05253112	**Good identification**					**05253112**
Clostridium innocuum		100.0	0.35	LAC 1		
05253200	**Very good identification**					**05253200**
Clostridium innocuum		100.0	0.59	RHA 1		
05253202	**Very good identification**					**05253202**
Clostridium innocuum		100.0	0.59	RHA 1		
05253210	**Good identification**					**05253210**
Clostridium innocuum		100.0	0.35	RHA 1		
05253212	**Good identification**					**05253212**
Clostridium innocuum		100.0	0.35	RHA 1		
05253400	**Very good identification**					**05253400**
Clostridium innocuum		100.0	0.80			
05253401	**Good identification**					**05253401**
Clostridium innocuum		100.0	0.46	URE 1		
05253402	**Very good identification**					**05253402**
Clostridium innocuum		100.0	0.80			
05253403	**Good identification**					**05253403**
Clostridium innocuum		100.0	0.46	URE 1		
05253410	**Very good identification**					**05253410**
Clostridium innocuum		100.0	0.57	MLZ 4		
05253412	**Very good identification**					**05253412**
Clostridium innocuum		100.0	0.57	MLZ 4		
05253420	**Good identification**					**05253420**
Clostridium innocuum		100.0	0.46	NAG 1		
05253422	**Good identification**					**05253422**
Clostridium innocuum		100.0	0.46	NAG 1		
05253440	**Good identification**					**05253440**
Clostridium innocuum		100.0	0.46	SOR 1		
05253442	**Good identification**					**05253442**
Clostridium innocuum		100.0	0.46	SOR 1		
05253500	**Good identification**					**05253500**
Clostridium innocuum		100.0	0.46	LAC 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05253502	**Good identification**						**05253502**
Clostridium innocuum		100.0	0.46	LAC 1			
05253600	**Good identification**						**05253600**
Clostridium innocuum		100.0	0.46	RHA 1			
05253602	**Good identification**						**05253602**
Clostridium innocuum		100.0	0.46	RHA 1			
05254000	**Low discrimination**						**05254000**
					GRA	SPO	
Fusobacterium mortiferum		75.92	0.43	XYL 7	1	1	
Clostridium innocuum		10.25	0.15		99	99	
Clostridium difficile		8.72	0.11		99	99	
05254100	**Good identification**						**05254100**
Fusobacterium mortiferum		100.0	0.49	XYL 7			
05255000	**Low discrimination**						**05255000**
					GRA	SPO	
Clostridium innocuum		59.61	0.37	MAN 99	99	99	
Fusobacterium mortiferum		34.84	0.46	XYL 5	1	1	
Lactobacillus jensenii		4.02	0.19		99	1	
05255002	**Low discrimination**						**05255002**
					GRA	SPO	
Clostridium innocuum		88.06	0.37	MAN 99	99	99	
Fusobacterium mortiferum		5.72	0.29	XYL 5	1	1	
Lactobacillus jensenii		3.96	0.16		99	1	
05255100	**Good identification**						**05255100**
					GRA	SPO	
Fusobacterium mortiferum		95.49	0.52	XYL 7	1	1	
Lactobacillus catenaformis		2.71	0.15		99	1	
Clostridium innocuum		0.71	0.03		99	99	
05255102	**Low discrimination**						**05255102**
					GRA	SPO	
Fusobacterium mortiferum		69.75	0.35	XYL 7	1	1	
Lactobacillus catenaformis		17.78	0.15	BGL 10	99	1	
Clostridium innocuum		4.65	0.03		99	99	
05256000	**Acceptable identification**						**05256000**
					SPO		
Clostridium innocuum		92.71	0.49	GAL 95	99		
Clostridium difficile		7.17	0.27	XYL 5	99		
Eubacterium limosum		0.12	-.06		1		
05256002	**Good identification**						**05256002**
Clostridium innocuum		100.0	0.49	GAL 95			
05256010	**Acceptable identification**						**05256010**
Clostridium difficile		90.06	0.39	CEL 5			
Clostridium innocuum		9.94	0.26	XYL 5			
05256400	**Good identification**						**05256400**
Clostridium innocuum		100.0	0.36	GAL 95			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
05256402	**Good identification**						**05256402**
Clostridium innocuum		100.0	0.36	GAL 95			
05257000	**Very good identification**						**05257000**
					GRA	SPO	
Clostridium innocuum		99.99	0.71	XYL 5	99	99	
Fusobacterium mortiferum		0.01	0.11		1	1	
05257001	**Good identification**						**05257001**
Clostridium innocuum		100.0	0.37	XYL 5			
05257002	**Very good identification**						**05257002**
Clostridium innocuum		100.0	0.71	XYL 5			
05257003	**Good identification**						**05257003**
Clostridium innocuum		100.0	0.37	XYL 5			
05257010	**Good identification**						**05257010**
Clostridium innocuum		99.52	0.48	XYL 5			
Clostridium difficile		0.48	0.05	MLZ 4			
05257012	**Good identification**						**05257012**
Clostridium innocuum		100.0	0.48	XYL 5			
05257020	**Good identification**						**05257020**
Clostridium innocuum		100.0	0.37	XYL 5			
05257022	**Good identification**						**05257022**
Clostridium innocuum		100.0	0.37	XYL 5			
05257040	**Good identification**						**05257040**
Clostridium innocuum		100.0	0.37	XYL 5			
05257042	**Good identification**						**05257042**
Clostridium innocuum		100.0	0.37	XYL 5			
05257100	**Good identification**						**05257100**
					GRA	SPO	
Clostridium innocuum		98.64	0.37	XYL 5	99	99	
Fusobacterium mortiferum		1.36	0.18	LAC 1	1	1	
05257102	**Good identification**						**05257102**
Clostridium innocuum		100.0	0.37	XYL 5			
05257200	**Good identification**						**05257200**
Clostridium innocuum		100.0	0.37	XYL 5			
05257202	**Good identification**						**05257202**
Clostridium innocuum		100.0	0.37	XYL 5			
05257400	**Very good identification**						**05257400**
Clostridium innocuum		100.0	0.58	XYL 5			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05257402	**Very good identification**						**05257402**
Clostridium innocuum		100.0	0.58	XYL 5			
05261120	**Good identification**						**05261120**
Clostridium septicum		100.0	0.39	MLT 94			
05262000	**Good identification**						**05262000**
Clostridium innocuum		99.68	0.41	FRU 95			
Clostridium difficile		0.32	-.05	GAL 95			
05262002	**Good identification**						**05262002**
Clostridium innocuum		100.0	0.41	FRU 95			
05263000	**Very good identification**						**05263000**
Clostridium innocuum		100.0	0.63	FRU 95			
05263002	**Very good identification**						**05263002**
Clostridium innocuum		100.0	0.63	FRU 95			
05263010	**Good identification**						**05263010**
Clostridium innocuum		100.0	0.40	FRU 95			
05263012	**Good identification**						**05263012**
Clostridium innocuum		100.0	0.40	FRU 95			
05263400	**Good identification**						**05263400**
Clostridium innocuum		100.0	0.50	FRU 95			
05263402	**Good identification**						**05263402**
Clostridium innocuum		100.0	0.50	FRU 95			
05267000	**Good identification**						**05267000**
Clostridium innocuum		100.0	0.41	FRU 95			
05267002	**Good identification**						**05267002**
Clostridium innocuum		100.0	0.41	FRU 95			
05270000	**Low discrimination**						**05270000**
					GRA	SPO	
Fusobacterium mortiferum		71.19	0.53		1	1	
Clostridium innocuum		16.17	0.29		99	99	
Lactobacillus jensenii		9.31	0.28		99	1	
05270100	**Very good identification**						**05270100**
					GRA	SPO	
Fusobacterium mortiferum		99.28	0.59		1	1	
Clostridium septicum		0.72	0.05		99	99	
05270102	**Acceptable identification**						**05270102**
					GRA		
Fusobacterium mortiferum		92.41	0.43	BGL 10	1		
Lactobacillus acidophilus		7.59	0.11		99		
05270120	**Good identification**						**05270120**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Clostridium septicum		98.61	0.39	MLT 94	99	99		
Fusobacterium mortiferum		1.39	0.25	GAL 99	1	1		
05271000	**Low discrimination**							**05271000**
					GRA	SPO		
Lactobacillus jensenii		68.83	0.62	SUC 90	99	1		
Clostridium innocuum		22.96	0.51	MLT 95	99	99		
Fusobacterium mortiferum		7.98	0.56		1	1		
05271002	**Low discrimination**							**05271002**
					SPO			
Lactobacillus jensenii		64.75	0.59	SUC 90	1			
Clostridium innocuum		32.40	0.51	MLT 95	99			
Lactobacillus acidophilus		1.38	0.27		1			
05271010	**Good identification**							**05271010**
					GRA	SPO		
Lactobacillus jensenii		95.68	0.54	SUC 90	99	1		
Clostridium innocuum		3.99	0.27	MLT 95	99	99		
Fusobacterium mortiferum		0.34	0.22		1	1		
05271012	**Acceptable identification**							**05271012**
					SPO			
Lactobacillus jensenii		94.11	0.51	SUC 90	1			
Clostridium innocuum		5.89	0.27	MLT 95	99			
05271020	**Low discrimination**							**05271020**
					SPO			
Lactobacillus jensenii		64.88	0.52	SUC 90	1			
Clostridium septicum		33.59	0.39	MLT 95	99			
Clostridium innocuum		0.88	0.17		99			
05271022	**Low discrimination**							**05271022**
					SPO			
Lactobacillus jensenii		87.61	0.49	SUC 90	1			
Lactobacillus acidophilus		7.45	0.27	MLT 95	1			
Clostridium innocuum		1.77	0.17		99			
05271030	**Good identification**							**05271030**
					SPO			
Lactobacillus jensenii		98.46	0.43	SUC 90	1			
Clostridium septicum		1.54	0.05	MLT 95	99			
05271032	**Good identification**							**05271032**
Lactobacillus jensenii		100.0	0.40	SUC 90				
05271100	**Low discrimination**							**05271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		61.99	0.62		1	1	1	
Clostridium septicum		29.66	0.39		1	99	99	
Atopobium parvulum		4.21	0.31		99	99	1	
05271102	**Low discrimination**							**05271102**
					GRA			
Lactobacillus acidophilus		76.14	0.45	SUC 99	99			
Fusobacterium mortiferum		14.04	0.46	MLT 94	1			
Lactobacillus jensenii		3.15	0.25		99			
05271120	**Very good identification**							**05271120**
					COC	GRA	SPO	
Clostridium septicum		99.87	0.73	MLT 94	1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		0.03	0.18		99	99	1	
Fusobacterium mortiferum		0.02	0.28		1	1	1	
05271121	**Good identification**							**05271121**
					SPO			
Clostridium septicum		99.79	0.39	MLT 94	99			
Actinomyces naeslundii		0.21	0.22	URE 1	1			
05271122	**Low discrimination**							**05271122**
					SPO			
Clostridium septicum		59.52	0.44	MLT 94	99			
Lactobacillus acidophilus		37.09	0.45	BGL 2	1			
Clostridium barati		2.10	0.22		99			
05271130	**Good identification**							**05271130**
					SPO			
Clostridium septicum		99.35	0.39	MLT 94	99			
Lactobacillus jensenii		0.65	0.09	MLZ 1	1			
05271160	**Good identification**							**05271160**
Clostridium septicum		99.71	0.39	MLT 94				
Clostridium perfringens		0.29	0.02	SOR 1				
05271320	**Good identification**							**05271320**
Clostridium septicum		100.0	0.39	MLT 94				
05271400	**Low discrimination**							**05271400**
					GRA	SPO		
Clostridium innocuum		83.93	0.38	MAN 99	99	99		
Lactobacillus jensenii		14.40	0.28		99	1		
Fusobacterium mortiferum		1.67	0.22		1	1		
05271402	**Low discrimination**							**05271402**
					SPO			
Clostridium innocuum		89.74	0.38	MAN 99	99			
Lactobacillus jensenii		10.27	0.25		1			
05271520	**Good identification**							**05271520**
Clostridium septicum		100.0	0.39	MLT 94				
05272000	**Very good identification**							**05272000**
					GRA	SPO		
Clostridium innocuum		98.29	0.63	GAL 95	99	99		
Clostridium difficile		1.63	0.30		99	99		
Fusobacterium mortiferum		0.04	0.19		1	1		
05272002	**Very good identification**							**05272002**
Clostridium innocuum		99.98	0.63	GAL 95				
Clostridium difficile		0.02	-.05					
05272010	**Low discrimination**							**05272010**
Clostridium difficile		66.00	0.41	TRE 7				
Clostridium innocuum		34.00	0.40	CEL 5				
05272012	**Good identification**							**05272012**
Clostridium innocuum		98.08	0.40	GAL 95				
Clostridium difficile		1.92	0.07	MLZ 4				
05272400	**Good identification**							**05272400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		99.91	0.50	GAL 95			
Clostridium difficile		0.10	-.05				
05272402	**Good identification**						**05272402**
Clostridium innocuum		100.0	0.50	GAL 95			
05273000	**Very good identification**						**05273000**
					GRA	SPO	
Clostridium innocuum		99.96	0.85		99	99	
Lactobacillus jensenii		0.03	0.28		99	1	
Fusobacterium mortiferum		0.00	0.22		1	1	
05273001	**Good identification**						**05273001**
Clostridium innocuum		100.0	0.51	URE 1			
05273002	**Very good identification**						**05273002**
					SPO		
Clostridium innocuum		99.98	0.85		99		
Lactobacillus jensenii		0.02	0.25		1		
Lactobacillus acidophilus		0.00	0.05		1		
05273003	**Good identification**						**05273003**
Clostridium innocuum		100.0	0.51	URE 1			
05273010	**Very good identification**						**05273010**
					SPO		
Clostridium innocuum		99.65	0.62	MLZ 4	99		
Lactobacillus jensenii		0.24	0.19		1		
Clostridium difficile		0.10	0.07		99		
05273012	**Very good identification**						**05273012**
					SPO		
Clostridium innocuum		99.84	0.62	MLZ 4	99		
Lactobacillus jensenii		0.16	0.16		1		
05273020	**Good identification**						**05273020**
					SPO		
Clostridium innocuum		98.86	0.51	NAG 1	99		
Lactobacillus jensenii		0.75	0.17		1		
Clostridium septicum		0.39	0.05		99		
05273022	**Good identification**						**05273022**
					SPO		
Clostridium innocuum		99.28	0.51	NAG 1	99		
Lactobacillus jensenii		0.50	0.14		1		
Lactobacillus acidophilus		0.22	0.05		1		
05273040	**Good identification**						**05273040**
Clostridium innocuum		100.0	0.51	SOR 1			
05273042	**Good identification**						**05273042**
Clostridium innocuum		100.0	0.51	SOR 1			
05273100	**Good identification**						**05273100**
					GRA	SPO	
Clostridium innocuum		98.60	0.51	LAC 1	99	99	
Fusobacterium mortiferum		0.81	0.28		1	1	
Clostridium septicum		0.39	0.05		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05273102	**Good identification**					**05273102**
					SPO	
Clostridium innocuum		97.01	0.51	LAC 1	99	
Lactobacillus acidophilus		2.49	0.24		1	
Clostridium tertium		0.30	0.09		99	
05273120	**Good identification**					**05273120**
Clostridium septicum		95.76	0.39	MLT 94		
Clostridium innocuum		2.49	0.17	MAN 1		
Clostridium tertium		0.77	0.09			
05273200	**Good identification**					**05273200**
Clostridium innocuum		100.0	0.51	RHA 1		
05273202	**Good identification**					**05273202**
Clostridium innocuum		100.0	0.51	RHA 1		
05273400	**Very good identification**					**05273400**
Clostridium innocuum		100.0	0.72			
05273401	**Good identification**					**05273401**
Clostridium innocuum		100.0	0.38	URE 1		
05273402	**Very good identification**					**05273402**
Clostridium innocuum		100.0	0.72			
05273403	**Good identification**					**05273403**
Clostridium innocuum		100.0	0.38	URE 1		
05273410	**Good identification**					**05273410**
Clostridium innocuum		100.0	0.49	MLZ 4		
05273412	**Good identification**					**05273412**
Clostridium innocuum		100.0	0.49	MLZ 4		
05273420	**Good identification**					**05273420**
Clostridium innocuum		100.0	0.38	NAG 1		
05273422	**Good identification**					**05273422**
Clostridium innocuum		100.0	0.38	NAG 1		
05273440	**Good identification**					**05273440**
Clostridium innocuum		100.0	0.38	SOR 1		
05273442	**Good identification**					**05273442**
Clostridium innocuum		100.0	0.38	SOR 1		
05273500	**Good identification**					**05273500**
Clostridium innocuum		100.0	0.38	LAC 1		
05273502	**Good identification**					**05273502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium innocuum		100.0	0.38	LAC 1		
05273600	**Good identification**					**05273600**
Clostridium innocuum		100.0	0.38	RHA 1		

Code		ID	T	tests	Differentiation tests			Code
Taxon		score	index	against				
05273602	**Good identification**							**05273602**
Clostridium innocuum		100.0	0.38	RHA 1				
05274100	**Good identification**							**05274100**
					COC	GRA		
Fusobacterium mortiferum		97.53	0.40	XYL 7	1	1		
Atopobium parvulum		2.48	0.01		99	99		
05275100	**Low discrimination**							**05275100**
					COC	GRA	SPO	
Atopobium parvulum		61.16	0.35	SUC 96	99	99	1	
Fusobacterium mortiferum		36.50	0.43	MLT 99	1	1	1	
Clostridium septicum		2.34	0.05		1	99	99	
05275120	**Good identification**							**05275120**
					COC	SPO		
Clostridium septicum		95.56	0.39	MLT 94	1	99		
Atopobium parvulum		4.44	0.22	XYL 1	99	1		
05275322	**Good identification**							**05275322**
Parabacteroides distasonis		100.0	0.35	RAF 99				
05276000	**Good identification**							**05276000**
Clostridium innocuum		98.37	0.41	GAL 95				
Clostridium difficile		1.63	0.08	XYL 5				
05276002	**Good identification**							**05276002**
Clostridium innocuum		100.0	0.41	GAL 95				
05277000	**Very good identification**							**05277000**
Clostridium innocuum		100.0	0.63	XYL 5				
05277002	**Very good identification**							**05277002**
Clostridium innocuum		100.0	0.63	XYL 5				
05277010	**Good identification**							**05277010**
Clostridium innocuum		100.0	0.40	XYL 5				
05277012	**Good identification**							**05277012**
Clostridium innocuum		100.0	0.40	XYL 5				
05277400	**Good identification**							**05277400**
Clostridium innocuum		100.0	0.50	XYL 5				
05277402	**Good identification**							**05277402**
Clostridium innocuum		100.0	0.50	XYL 5				
05300000	**Low discrimination**							**05300000**
					COC	GRA	SPO	
Prevotella disiens		63.87	0.66	SAL 1	1	1	1	
Clostridium bifermentans		16.60	0.61		1	99	99	
Gemella morbillorum		16.35	0.64		99	99	1	
05300011	**Good identification**							**05300011**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		98.77	0.48	MNS 1	99	99		
Prevotella disiens		1.23	-.03	SAL 8	1	1		
05300020	**Low discrimination**							**05300020**
					COC	GRA	SPO	
Gemella morbillorum		36.02	0.48	SAL 1	99	99	1	
Prevotella bivia		26.29	0.32	NAG 10	1	1	1	
Clostridium bifermentans		21.01	0.41		1	99	99	
05300021	**Low discrimination**							**05300021**
					COC	GRA	SPO	
Clostridium novyi biovar A		72.73	0.37	MNS 1	1	99	99	
Gemella morbillorum		10.22	0.14	SAL 8	99	99	1	
Prevotella bivia		7.46	-.03		1	1	1	
05300030	**Low discrimination**							**05300030**
					COC	GRA	SPO	
Clostridium novyi biovar A		72.73	0.37	MNS 1	1	99	99	
Gemella morbillorum		10.22	0.14	SAL 8	99	99	1	
Prevotella bivia		7.46	-.03		1	1	1	
05300031	**Good identification**							**05300031**
Clostridium novyi biovar A		100.0	0.37	MNS 1				
05300040	**Low discrimination**							**05300040**
					GRA	SPO		
Clostridium bifermentans		27.05	0.37	IND 90	99	99		
Prevotella disiens		25.23	0.32	SAL 10	1	1		
Clostridium botulinum biovar B		15.52	0.31	SOR 4	99	99		
05300041	**Good identification**							**05300041**
					GRA	SPO		
Clostridium novyi biovar A		95.21	0.45	MNS 1	99	99		
Clostridium bifermentans		1.91	0.03	SAL 8	99	99		
Prevotella disiens		1.78	-.03		1	1		
05300050	**Low discrimination**							**05300050**
Clostridium novyi biovar A		62.49	0.45	MNS 1				
Clostridium botulinum biovar B		35.08	0.26	SAL 8				
Clostridium bifermentans		1.25	0.03					
05300051	**Good identification**							**05300051**
Clostridium novyi biovar A		100.0	0.45	MNS 1				
05300120	**Very good identification**							**05300120**
					COC	GRA	SPO	
Prevotella bivia		99.96	0.66	SAL 1	1	1	1	
Clostridium septicum		0.02	0.09		1	99	99	
Gemella morbillorum		0.01	0.14		99	99	1	
05300400	**Low discrimination**							**05300400**
					COC	GRA	SPO	
Gemella morbillorum		68.29	0.48	SAL 1	99	99	1	
Prevotella disiens		24.26	0.32	ARA 10	1	1	1	
Clostridium bifermentans		6.30	0.27		1	99	99	
05301000	**Low discrimination**							**05301000**
					COC	GRA	SPO	
Gemella morbillorum		89.86	0.60	SAL 1	99	99	1	
Prevotella disiens		6.59	0.32		1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		1.71	0.27		1	99	99	
05301020	**Low discrimination**							**05301020**
					COC	GRA		
Gemella morbillorum		74.16	0.43	SAL 1	99	99		
Prevotella bivia		17.74	0.19	NAG 10	1	1		
Clostridium perfringens		2.92	0.19		1	99		
05301120	**Low discrimination**							**05301120**
					GRA	SPO		
Prevotella bivia		89.67	0.53	SAL 1	1	1		
Clostridium septicum		9.17	0.43		99	99		
Clostridium perfringens		1.10	0.34		99	84		
05301400	**Good identification**							**05301400**
					COC	GRA		
Gemella morbillorum		99.34	0.43	SAL 1	99	99		
Prevotella disiens		0.66	-.03	ARA 10	1	1		
05304000	**Low discrimination**							**05304000**
					COC	GRA	SPO	
Gemella morbillorum		68.29	0.48	SAL 1	99	99	1	
Prevotella disiens		24.26	0.32	XYL 10	1	1	1	
Clostridium bifermentans		6.30	0.27		1	99	99	
05305000	**Good identification**							**05305000**
					COC	GRA		
Gemella morbillorum		99.34	0.43	SAL 1	99	99		
Prevotella disiens		0.66	-.03	XYL 10	1	1		
05310000	**Low discrimination**							**05310000**
					COC	GRA		
Prevotella disiens		71.04	0.66	SAL 1	1	1		
Gemella morbillorum		22.22	0.66		99	99		
Prevotella intermedia		3.67	0.52		1	1		
05310020	**Low discrimination**							**05310020**
					COC	GRA		
Gemella morbillorum		67.96	0.50	SAL 1	99	99		
Prevotella disiens		19.75	0.32	NAG 10	1	1		
Prevotella bivia		4.51	0.15		1	1		
05310040	**Low discrimination**							**05310040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		59.48	0.41	SUC 90	1	99	99	
Prevotella disiens		24.17	0.32	SAL 1	1	1	1	
Gemella morbillorum		7.56	0.32		99	99	1	
05310050	**Acceptable identification**							**05310050**
Clostridium botulinum biovar B		91.95	0.36	SUC 90				
Clostridium novyi biovar A		4.55	0.28	SAL 1				
Clostridium difficile		2.74	0.10					
05310060	**Acceptable identification**							**05310060**
					COC	SPO		
Clostridium botulinum biovar B		94.45	0.36	SUC 90	1	99		
Gemella morbillorum		2.71	0.15	SAL 1	99	1		
Clostridium novyi biovar A		1.17	0.18		1	99		
05310100	**Low discrimination**							**05310100**
					COC	GRA		
Fusobacterium mortiferum		69.83	0.61	MLT 11	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella disiens		18.13	0.32		1	1	
Gemella morbillorum		5.67	0.32		99	99	
05310102	**Acceptable identification**						**05310102**
					GRA		
Fusobacterium mortiferum		94.75	0.44	MLT 11	1		
Prevotella disiens		2.24	-.03	BGL 10	1		
Actinomyces viscosus		1.60	0.01		99		
05310120	**Low discrimination**						**05310120**
					GRA	SPO	
Prevotella bivia		84.35	0.50	SAL 1	1	1	
Clostridium septicum		13.70	0.43	FRU 10	99	99	
Clostridium perfringens		1.64	0.34		99	84	
05310400	**Low discrimination**						**05310400**
					COC	GRA	
Gemella morbillorum		76.08	0.50	SAL 1	99	99	
Prevotella disiens		22.11	0.32	ARA 10	1	1	
Prevotella intermedia		1.14	0.17		1	1	
05311000	**Low discrimination**						**05311000**
					COC	GRA	
Gemella morbillorum		52.62	0.61	SAL 1	99	99	
Lactobacillus jensenii		18.10	0.54		1	99	
Prevotella intermedia		16.12	0.52		1	1	
05311002	**Low discrimination**						**05311002**
					GRA		
Lactobacillus jensenii		79.59	0.51	SUC 90	99		
Fusobacterium mortiferum		5.73	0.41	CEL 95	1		
Lactobacillus catenaformis		4.15	0.29		99		
05311010	**Low discrimination**						**05311010**
					COC	GRA	
Lactobacillus jensenii		86.96	0.45	SUC 90	1	99	
Gemella morbillorum		7.66	0.27	CEL 95	99	99	
Prevotella intermedia		2.35	0.17		1	1	
05311012	**Good identification**						**05311012**
Lactobacillus jensenii		100.0	0.42	SUC 90			
05311020	**Low discrimination**						**05311020**
					COC	SPO	
Clostridium perfringens		44.99	0.53	SUC 91	1	84	
Clostridium septicum		27.92	0.43	LAC 88	1	99	
Gemella morbillorum		14.09	0.45		99	1	
05311022	**Low discrimination**						**05311022**
					GRA		
Clostridium perfringens		67.89	0.48	SUC 91	99		
Lactobacillus jensenii		23.30	0.40	LAC 88	99		
Prevotella oralis		3.01	0.21		1		
05311030	**Low discrimination**						**05311030**
					SPO		
Lactobacillus jensenii		80.53	0.35	SUC 90	1		
Clostridium perfringens		10.07	0.19	CEL 95	84		
Clostridium septicum		6.25	0.09		99		
05311060	**Acceptable identification**						**05311060**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		93.18	0.41	SUC 91	1	84	
Clostridium septicum		3.07	0.09	LAC 88	1	99	
Gemella morbillorum		1.55	0.11		99	1	
05311062	**Good identification**						**05311062**
Clostridium perfringens		98.21	0.35	SUC 91			
Lactobacillus jensenii		1.79	0.06	LAC 88			
05311100	**Low discrimination**						**05311100**
					GRA	SPO	
Fusobacterium mortiferum		48.89	0.64	MLT 11	1	1	
Clostridium septicum		31.08	0.43		99	99	
Clostridium perfringens		11.36	0.42		99	84	
05311102	**Low discrimination**						**05311102**
					GRA		
Fusobacterium mortiferum		29.02	0.47	MLT 11	1		
Clostridium perfringens		28.56	0.36	BGL 10	99		
Lactobacillus catenaformis		13.52	0.32		99		
05311120	**Low discrimination**						**05311120**
					GRA	SPO	
Clostridium septicum		88.34	0.77		99	99	
Clostridium perfringens		10.55	0.68		99	84	
Prevotella bivia		0.97	0.37		1	1	
05311121	**Low discrimination**						**05311121**
Clostridium septicum		86.73	0.43	URE 1			
Clostridium perfringens		10.36	0.34				
Clostridium chauvoei		1.29	0.18				
05311122	**Low discrimination**						**05311122**
					GRA	SPO	
Clostridium perfringens		67.34	0.62	SUC 91	99	84	
Clostridium septicum		24.46	0.48		99	99	
Prevotella oralis		6.38	0.41		1	1	
05311130	**Low discrimination**						**05311130**
					GRA	SPO	
Clostridium septicum		88.37	0.43	MLZ 1	99	99	
Clostridium perfringens		10.55	0.34		99	84	
Prevotella bivia		0.97	0.03		1	1	
05311160	**Low discrimination**						**05311160**
					GRA	SPO	
Clostridium perfringens		68.96	0.56	SUC 91	99	84	
Clostridium septicum		30.63	0.43		99	99	
Prevotella bivia		0.34	0.03		1	1	
05311162	**Good identification**						**05311162**
					GRA	SPO	
Clostridium perfringens		97.04	0.50	SUC 91	99	84	
Clostridium septicum		1.87	0.14		99	99	
Prevotella oralis		0.49	0.07		1	1	
05311320	**Low discrimination**						**05311320**
					GRA	SPO	
Clostridium septicum		87.55	0.43	RHA 1	99	99	
Clostridium perfringens		10.45	0.34		99	84	
Prevotella oralis		1.04	0.13		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05311322	**Low discrimination**							**05311322**
					GRA	SPO		
Prevotella oralis		71.93	0.35	SUC 92	1	1		
Clostridium perfringens		17.89	0.28	RAF 92	99	84		
Clostridium septicum		6.50	0.14		99	99		
05311400	**Low discrimination**							**05311400**
					COC			
Gemella morbillorum		87.56	0.45	SAL 1	99			
Lactobacillus fermentum		5.61	0.36	ARA 10	1			
Lactobacillus jensenii		2.74	0.19		1			
05311402	**Low discrimination**							**05311402**
					COC			
Lactobacillus fermentum		67.41	0.36	SUC 87	1			
Lactobacillus jensenii		21.95	0.16	SAL 1	1			
Gemella morbillorum		10.64	0.11		99			
05311500	**Low discrimination**							**05311500**
					GRA	SPO		
Lactobacillus fermentum		60.24	0.40	SUC 87	99	1		
Fusobacterium mortiferum		18.16	0.30	SAL 1	1	1		
Clostridium septicum		11.55	0.09		99	99		
05311502	**Low discrimination**							**05311502**
					GRA			
Lactobacillus fermentum		86.84	0.40	SUC 87	99			
Bifidobacterium adolescentis		7.39	0.16	SAL 1	99			
Fusobacterium mortiferum		2.91	0.13		1			
05311520	**Low discrimination**							**05311520**
					SPO			
Clostridium septicum		87.11	0.43	ARA 1	99			
Clostridium perfringens		10.40	0.34		84			
Lactobacillus fermentum		1.53	0.32		1			
05312000	**Low discrimination**							**05312000**
					GRA	SPO		
Clostridium difficile		36.25	0.37	MLT 1	99	99		
Prevotella disiens		28.69	0.32		1	1		
Eubacterium limosum		16.07	0.29		99	1		
05312010	**Good identification**							**05312010**
					GRA	SPO	CAT	
Clostridium difficile		99.57	0.48	MLT 1	99	99	1	
Propionibacterium granulosum		0.18	0.14		99	1	90	
Prevotella disiens		0.16	-.03		1	1	1	
05313101	**Good identification**							**05313101**
Actinomyces naeslundii		100.0	0.41	SUC 99				
05313102	**Low discrimination**							**05313102**
Actinomyces naeslundii		57.80	0.38	SUC 99				
Propionibacterium propionicum		11.14	0.15	RAF 92				
Lactobacillus acidophilus		10.92	0.01					
05313103	**Good identification**							**05313103**
Actinomyces naeslundii		100.0	0.41	SUC 99				
05313120	**Low discrimination**							**05313120**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium septicum		87.69	0.43	MAN 1	99	99	
Clostridium perfringens		10.47	0.34		99	84	
Prevotella bivia		0.96	0.03		1	1	
05313121	**Low discrimination**						**05313121**
					SPO		
Actinomyces naeslundii		59.89	0.41	SUC 99	1		
Clostridium septicum		40.11	0.09	RAF 92	99		
05313123	**Good identification**						**05313123**
Actinomyces naeslundii		100.0	0.41	SUC 99			
05314000	**Low discrimination**						**05314000**
					COC	GRA	
Gemella morbillorum		73.49	0.50	SAL 1	99	99	
Prevotella disiens		21.36	0.32	XYL 10	1	1	
Fusobacterium mortiferum		2.65	0.35		1	1	
05314100	**Low discrimination**						**05314100**
					COC	GRA	
Fusobacterium mortiferum		83.18	0.41	MLT 11	1	1	
Gemella morbillorum		9.98	0.15	XYL 7	99	99	
Actinomyces meyeri		3.95	0.09		1	99	
05315000	**Low discrimination**						**05315000**
					COC	GRA	
Gemella morbillorum		82.95	0.45	SAL 1	99	99	
Fusobacterium mortiferum		8.34	0.38	XYL 10	1	1	
Lactobacillus fermentum		3.12	0.32		1	99	
05315100	**Low discrimination**						**05315100**
					COC	GRA	
Fusobacterium mortiferum		48.86	0.44	MLT 11	1	1	
Atopobium parvulum		28.26	0.29	XYL 7	99	99	
Lactobacillus fermentum		12.77	0.36		1	99	
05315102	**Low discrimination**						**05315102**
					COC	GRA	
Lactobacillus fermentum		42.45	0.36	SUC 87	1	99	
Fusobacterium mortiferum		18.04	0.28	SAL 1	1	1	
Atopobium parvulum		16.57	0.16		99	99	
05315120	**Low discrimination**						**05315120**
					COC	SPO	
Clostridium septicum		86.23	0.43	XYL 1	1	99	
Clostridium perfringens		10.30	0.34		1	84	
Atopobium parvulum		1.04	0.16		99	1	
05315322	**Low discrimination**						**05315322**
					GRA		
Parabacteroides distasonis		66.48	0.38	RAF 99	1		
Prevotella oralis		30.84	0.20		1		
Eubacterium contortum		2.68	0.03		99		
05315500	**Good identification**						**05315500**
Lactobacillus fermentum		95.66	0.36	SUC 87			
Actinomyces meyeri		4.34	0.02	SAL 1			
05315502	**Low discrimination**						**05315502**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		49.71	0.36	SUC 87				
Bifidobacterium adolescentis		48.22	0.30	SAL 1				
Actinomyces israelii		2.07	0.13					
05320040	**Acceptable identification**							**05320040**
Clostridium botulinum biovar B		92.99	0.39	SUC 90				
Clostridium botulinum biovar A		5.68	0.21	SAL 1				
Clostridium bifermentans		0.55	0.03					
05321120	**Acceptable identification**							**05321120**
					GRA	SPO		
Clostridium septicum		90.17	0.51	FRU 99	99	99		
Clostridium perfringens		6.66	0.38		99	84		
Prevotella bivia		2.97	0.19		1	1		
05330040	**Good identification**							**05330040**
					GRA	SPO		
Clostridium botulinum biovar B		99.70	0.50	SUC 90	99	99		
Prevotella disiens		0.14	-.03	SAL 1	1	1		
Clostridium botulinum biovar A		0.10	0.01		99	99		
05330050	**Good identification**							**05330050**
Clostridium botulinum biovar B		100.0	0.44	SUC 90				
05330060	**Good identification**							**05330060**
Clostridium botulinum biovar B		99.42	0.44	SUC 90				
Clostridium perfringens		0.58	0.11	SAL 1				
05330070	**Good identification**							**05330070**
Clostridium botulinum biovar B		100.0	0.39	SUC 90				
05330100	**Low discrimination**							**05330100**
					COC	GRA	SPO	
Fusobacterium mortiferum		80.31	0.52	MLT 11	1	1	1	
Atopobium parvulum		9.46	0.24		99	99	1	
Clostridium septicum		7.77	0.17		1	99	99	
05330102	**Low discrimination**							**05330102**
					COC	GRA		
Fusobacterium mortiferum		78.14	0.35	MLT 11	1	1		
Atopobium parvulum		14.61	0.11	BGL 10	99	99		
Clostridium perfringens		7.25	0.07		1	99		
05330120	**Acceptable identification**							**05330120**
					GRA	SPO		
Clostridium septicum		91.09	0.51	GAL 99	99	99		
Clostridium perfringens		6.73	0.38		99	84		
Prevotella bivia		1.89	0.15		1	1		
05331000	**Good identification**							**05331000**
					GRA			
Lactobacillus jensenii		90.10	0.62	SUC 90	99			
Fusobacterium mortiferum		3.87	0.48	CEL 95	1			
Clostridium perfringens		1.78	0.32		99			
05331002	**Very good identification**							**05331002**
					GRA			
Lactobacillus jensenii		96.65	0.59	SUC 90	99			
Clostridium perfringens		1.35	0.26	CEL 95	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.69	0.32		1		
05331010	**Good identification**						**05331010**
					GRA	CAT	
Lactobacillus jensenii		99.11	0.54	SUC 90	99	1	
Propionibacterium granulosum		0.76	0.24	CEL 95	99	90	
Fusobacterium mortiferum		0.13	0.14		1	5	
05331012	**Good identification**						**05331012**
					CAT		
Lactobacillus jensenii		99.71	0.51	SUC 90	1		
Propionibacterium granulosum		0.29	0.14	CEL 95	90		
05331020	**Low discrimination**						**05331020**
					SPO		
Clostridium septicum		41.65	0.51	LAC 99	99		
Clostridium perfringens		41.55	0.58		84		
Lactobacillus jensenii		16.26	0.52		1		
05331022	**Low discrimination**						**05331022**
					SPO		
Clostridium perfringens		61.99	0.52	SUC 91	84		
Lactobacillus jensenii		34.37	0.49	LAC 88	1		
Clostridium septicum		2.69	0.22		99		
05331030	**Low discrimination**						**05331030**
					SPO		
Lactobacillus jensenii		86.01	0.43	SUC 90	1		
Clostridium septicum		6.68	0.17	CEL 95	99		
Clostridium perfringens		6.66	0.24		84		
05331032	**Acceptable identification**						**05331032**
Lactobacillus jensenii		94.82	0.40	SUC 90			
Clostridium perfringens		5.18	0.18	CEL 95			
05331060	**Acceptable identification**						**05331060**
					SPO		
Clostridium perfringens		91.62	0.45	SUC 91	84		
Clostridium septicum		4.87	0.17	LAC 88	99		
Lactobacillus jensenii		1.90	0.17		1		
05331062	**Good identification**						**05331062**
Clostridium perfringens		96.69	0.40	SUC 91			
Lactobacillus jensenii		2.84	0.14	LAC 88			
Lactobacillus acidophilus		0.47	-.02				
05331100	**Low discrimination**						**05331100**
					COC	SPO	
Atopobium parvulum		46.47	0.59	SUC 96	99	1	
Clostridium septicum		38.18	0.51		1	99	
Clostridium perfringens		8.64	0.47		1	84	
05331102	**Low discrimination**						**05331102**
					COC		
Atopobium parvulum		51.90	0.46	SUC 96	99		
Clostridium perfringens		25.74	0.41		1		
Lactobacillus acidophilus		9.73	0.32		1		
05331103	**Low discrimination**						**05331103**
					COC		
Actinomyces naeslundii		60.77	0.41	SUC 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		26.22	0.11	RAF 92	99			
Clostridium perfringens		13.00	0.07		1			
05331120	**Good identification**							**05331120**
					COC	SPO		
Clostridium septicum		92.92	0.85		1	99		
Clostridium perfringens		6.87	0.73		1	84		
Atopobium parvulum		0.20	0.46		99	1		
05331121	**Acceptable identification**							**05331121**
					SPO			
Clostridium septicum		92.45	0.51	URE 1	99			
Clostridium perfringens		6.83	0.38		84			
Actinomyces naeslundii		0.47	0.41		1			
05331122	**Low discrimination**							**05331122**
					SPO			
Clostridium perfringens		61.87	0.67	SUC 91	84			
Clostridium septicum		36.31	0.56		99			
Lactobacillus acidophilus		0.72	0.32		1			
05331130	**Acceptable identification**							**05331130**
					COC	SPO		
Clostridium septicum		92.82	0.51	MLZ 1	1	99		
Clostridium perfringens		6.86	0.38		1	84		
Atopobium parvulum		0.20	0.11		99	1		
05331160	**Low discrimination**							**05331160**
					COC	SPO		
Clostridium perfringens		58.16	0.60	SUC 91	1	84		
Clostridium septicum		41.73	0.51		1	99		
Atopobium parvulum		0.09	0.11		99	1		
05331162	**Good identification**							**05331162**
					SPO			
Clostridium perfringens		96.19	0.55	SUC 91	84			
Clostridium septicum		2.99	0.22		99			
Lactobacillus acidophilus		0.73	0.16		1			
05331320	**Acceptable identification**							**05331320**
					COC	SPO		
Clostridium septicum		92.93	0.51	RHA 1	1	99		
Clostridium perfringens		6.87	0.38		1	84		
Atopobium parvulum		0.20	0.11		99	1		
05331520	**Acceptable identification**							**05331520**
					COC	SPO		
Clostridium septicum		92.93	0.51	ARA 1	1	99		
Clostridium perfringens		6.87	0.38		1	84		
Atopobium parvulum		0.20	0.11		99	1		
05332000	**Low discrimination**							**05332000**
					GRA	SPO	CAT	
Propionibacterium granulosum		67.66	0.37	SUC 93	99	1	90	
Clostridium difficile		26.55	0.17	SAL 1	99	99	1	
Prevotella disiens		2.82	-.03		1	1	1	
05333000	**Low discrimination**							**05333000**
					SPO	CAT		
Propionibacterium granulosum		47.83	0.37	SUC 93	1	90		
Lactobacillus jensenii		34.29	0.28	SAL 1	1	1		
Clostridium innocuum		11.44	0.17		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05333001	**Good identification**							**05333001**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05333003	**Good identification**							**05333003**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05333021	**Good identification**							**05333021**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05333023	**Good identification**							**05333023**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05333101	**Good identification**							**05333101**
Actinomyces naeslundii		100.0	0.46	SUC 99				
05333102	**Low discrimination**							**05333102**
					COC			
Actinomyces naeslundii		43.40	0.43	SUC 99	1			
Atopobium parvulum		14.47	0.11	RAF 92	99			
Lactobacillus acidophilus		14.14	0.10		1			
05333103	**Good identification**							**05333103**
Actinomyces naeslundii		100.0	0.46	SUC 99				
05333120	**Acceptable identification**							**05333120**
					SPO			
Clostridium septicum		92.10	0.51	MAN 1	99			
Clostridium perfringens		6.81	0.38		84			
Actinomyces naeslundii		0.60	0.43		1			
05333123	**Good identification**							**05333123**
Actinomyces naeslundii		100.0	0.46	SUC 99				
05335100	**Very good identification**							**05335100**
					COC	GRA	SPO	
Atopobium parvulum		98.89	0.63	SUC 96	99	99	1	
Fusobacterium mortiferum		0.52	0.35		1	1	1	
Clostridium septicum		0.44	0.17		1	99	99	
05335102	**Good identification**							**05335102**
					COC	GRA		
Atopobium parvulum		98.68	0.50	SUC 96	99	99		
Actinomyces israelii		0.49	0.25		1	99		
Fusobacterium mortiferum		0.32	0.19		1	1		
05335120	**Low discrimination**							**05335120**
					COC	SPO		
Clostridium septicum		67.90	0.51	XYL 1	1	99		
Atopobium parvulum		27.08	0.50		99	1		
Clostridium perfringens		5.02	0.38		1	84		
05335122	**Low discrimination**							**05335122**
					COC	GRA		
Atopobium parvulum		44.71	0.37	SUC 96	99	99		
Clostridium perfringens		22.09	0.33		1	99		
Parabacteroides distasonis		19.35	0.41		1	1		
05335132	**Good identification**							**05335132**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Parabacteroides distasonis		95.85	0.37	RAF 99	1	1	
Atopobium parvulum		4.16	0.03		99	99	
05335322	**Good identification**						**05335322**
					COC	GRA	
Parabacteroides distasonis		98.59	0.49	RAF 99	1	1	
Atopobium parvulum		0.77	0.03		99	99	
Prevotella oralis		0.64	-.01		1	1	
05335332	**Good identification**						**05335332**
Parabacteroides distasonis		98.67	0.45	RAF 99			
Clostridium clostridioforme		1.33	0.07				
05335522	**Good identification**						**05335522**
					COC	GRA	
Parabacteroides distasonis		96.33	0.37	RAF 99	1	1	
Atopobium parvulum		3.67	0.03		99	99	
05335722	**Good identification**						**05335722**
Parabacteroides distasonis		96.09	0.46	RAF 99			
Clostridium clostridioforme		3.92	0.16				
05335732	**Low discrimination**						**05335732**
Parabacteroides distasonis		74.37	0.41	RAF 99			
Clostridium clostridioforme		25.63	0.27				
05337102	**Low discrimination**						**05337102**
					COC	SPO	
Actinomyces israelii		61.22	0.37	SUC 98	1	1	
Atopobium parvulum		22.07	0.16	CEL 93	99	1	
Clostridium tertium		12.25	0.11		1	99	
05337502	**Low discrimination**						**05337502**
Actinomyces israelii		62.62	0.36	SUC 98			
Bifidobacterium adolescentis		37.38	0.26	CEL 93			
05341000	**Low discrimination**						**05341000**
					COC		
Lactobacillus jensenii		75.14	0.41	SUC 90	1		
Lactobacillus catenaformis		13.61	0.29	FRU 99	1		
Gemella morbillorum		9.41	0.26		99		
05341002	**Low discrimination**						**05341002**
Lactobacillus jensenii		76.17	0.38	SUC 90			
Lactobacillus catenaformis		20.69	0.29	FRU 99			
Lactobacillus acidophilus		1.20	0.05				
05341120	**Good identification**						**05341120**
					GRA	SPO	
Clostridium septicum		95.16	0.51	FRU 99	99	99	
Prevotella bivia		2.97	0.19		1	1	
Clostridium paraputrificum		0.87	0.14		99	99	
05343000	**Good identification**						**05343000**
					SPO		
Clostridium innocuum		98.72	0.37	MLT 1	99		
Lactobacillus jensenii		1.28	0.07	FRU 95	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05343002	**Good identification**						**05343002**
					SPO		
Clostridium innocuum		99.15	0.37	MLT 1	99		
Lactobacillus jensenii		0.86	0.04	FRU 95	1		
05350000	**Low discrimination**						**05350000**
					GRA		
Lactobacillus jensenii		33.75	0.41	SUC 90	99		
Prevotella disiens		30.66	0.32	GAL 99	1		
Fusobacterium mortiferum		16.87	0.46		1		
05350002	**Low discrimination**						**05350002**
					GRA		
Lactobacillus jensenii		68.15	0.38	SUC 90	99		
Lactobacillus catenaformis		18.51	0.29	GAL 99	99		
Fusobacterium mortiferum		5.68	0.30		1		
05350100	**Low discrimination**						**05350100**
					GRA	SPO	
Fusobacterium mortiferum		71.49	0.52	MLT 11	1	1	
Lactobacillus catenaformis		16.65	0.32		99	1	
Clostridium septicum		6.54	0.17		99	99	
05350120	**Good identification**						**05350120**
					GRA	SPO	
Clostridium septicum		95.12	0.51	GAL 99	99	99	
Prevotella bivia		1.87	0.15		1	1	
Clostridium barati		1.28	0.22		99	99	
05351000	**Low discrimination**						**05351000**
					GRA		
Lactobacillus jensenii		83.55	0.76	SUC 90	99		
Lactobacillus catenaformis		15.13	0.63		99		
Fusobacterium mortiferum		0.63	0.49		1		
05351001	**Low discrimination**						**05351001**
Lactobacillus jensenii		83.28	0.41	SUC 90			
Lactobacillus catenaformis		15.08	0.29	URE 1			
Actinomyces naeslundii		1.01	0.30				
05351002	**Low discrimination**						**05351002**
Lactobacillus jensenii		76.63	0.73	SUC 90			
Lactobacillus catenaformis		20.81	0.63				
Lactobacillus acidophilus		1.20	0.39				
05351003	**Low discrimination**						**05351003**
Lactobacillus jensenii		75.81	0.38	SUC 90			
Lactobacillus catenaformis		20.59	0.29	URE 1			
Actinomyces naeslundii		1.37	0.30				
05351010	**Very good identification**						**05351010**
					GRA		
Lactobacillus jensenii		99.41	0.67	SUC 90	99		
Lactobacillus catenaformis		0.55	0.29		99		
Fusobacterium mortiferum		0.02	0.15		1		
05351012	**Very good identification**						**05351012**
Lactobacillus jensenii		99.07	0.64	SUC 90			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		0.82	0.29			
Lactobacillus acidophilus		0.05	0.05			
05351020	**Low discrimination**					**05351020**
					SPO	
Lactobacillus jensenii		60.25	0.65	SUC 90	1	
Clostridium septicum		25.72	0.51		99	
Lactobacillus catenaformis		10.91	0.53		1	
05351022	**Low discrimination**					**05351022**
					SPO	
Lactobacillus jensenii		64.73	0.62	SUC 90	1	
Lactobacillus catenaformis		17.58	0.53		1	
Clostridium barati		10.61	0.44		99	
05351030	**Very good identification**					**05351030**
					SPO	
Lactobacillus jensenii		98.09	0.57	SUC 90	1	
Clostridium septicum		1.27	0.17		99	
Lactobacillus catenaformis		0.54	0.18		1	
05351032	**Good identification**					**05351032**
					SPO	
Lactobacillus jensenii		98.51	0.54	SUC 90	1	
Lactobacillus catenaformis		0.81	0.18		1	
Clostridium barati		0.49	0.09		99	
05351040	**Low discrimination**					**05351040**
					SPO	
Lactobacillus jensenii		82.32	0.41	SUC 90	1	
Lactobacillus catenaformis		14.91	0.29	SOR 1	1	
Clostridium barati		2.15	0.09		99	
05351042	**Low discrimination**					**05351042**
Lactobacillus jensenii		63.24	0.38	SUC 90		
Lactobacillus catenaformis		17.18	0.29	SOR 1		
Lactobacillus acidophilus		12.15	0.23			
05351100	**Low discrimination**					**05351100**
					SPO	
Lactobacillus catenaformis		61.57	0.66	SUC 99	1	
Clostridium septicum		24.19	0.51		99	
Collinsella aerofaciens		4.53	0.32		1	
05351102	**Low discrimination**					**05351102**
					SPO	
Lactobacillus catenaformis		62.12	0.66	SUC 99	1	
Lactobacillus acidophilus		27.53	0.57		1	
Clostridium barati		6.25	0.44		99	
05351120	**Very good identification**					**05351120**
					SPO	
Clostridium septicum		97.97	0.86		99	
Clostridium paraputrificum		0.89	0.48		99	
Lactobacillus catenaformis		0.63	0.56		1	
05351121	**Good identification**					**05351121**
					SPO	
Clostridium septicum		97.76	0.51	URE 1	99	
Clostridium paraputrificum		0.89	0.14		99	
Lactobacillus catenaformis		0.63	0.21		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05351122	**Low discrimination**						**05351122**
					SPO		
Clostridium septicum		35.88	0.57	BGL 2	99		
Lactobacillus acidophilus		20.04	0.57		1		
Clostridium barati		13.64	0.52		99		
05351130	**Good identification**						**05351130**
					SPO		
Clostridium septicum		97.32	0.51	MLZ 1	99		
Clostridium paraputrificum		0.89	0.14		99		
Lactobacillus jensenii		0.77	0.23		1		
05351142	**Low discrimination**						**05351142**
					SPO		
Lactobacillus acidophilus		73.98	0.41	SUC 99	1		
Lactobacillus catenaformis		13.64	0.32	SOR 11	1		
Clostridium barati		11.81	0.25		99		
05351160	**Low discrimination**						**05351160**
Clostridium septicum		88.69	0.51	SOR 1			
Clostridium paraputrificum		6.02	0.29				
Clostridium perfringens		2.63	0.32				
05351162	**Low discrimination**						**05351162**
					SPO		
Lactobacillus acidophilus		49.35	0.41	SUC 99	1		
Clostridium barati		23.63	0.34	SOR 11	99		
Clostridium paraputrificum		10.30	0.22		99		
05351200	**Low discrimination**						**05351200**
					SPO		
Lactobacillus jensenii		82.32	0.41	SUC 90	1		
Lactobacillus catenaformis		14.91	0.29	RHA 1	1		
Clostridium barati		2.15	0.09		99		
05351202	**Low discrimination**						**05351202**
					SPO		
Lactobacillus jensenii		65.81	0.38	SUC 90	1		
Lactobacillus catenaformis		17.87	0.29	RHA 1	1		
Clostridium barati		7.74	0.17		99		
05351302	**Low discrimination**						**05351302**
					SPO		
Lactobacillus acidophilus		57.83	0.38	SUC 99	1		
Lactobacillus catenaformis		17.51	0.32	RHA 7	1		
Clostridium barati		15.16	0.25		99		
05351320	**Acceptable identification**						**05351320**
					GRA	SPO	
Clostridium septicum		94.73	0.51	RHA 1	99	99	
Clostridium barati		2.11	0.25		99	99	
Prevotella oralis		1.26	0.23		1	1	
05351322	**Low discrimination**						**05351322**
					GRA	SPO	
Prevotella oralis		56.73	0.45	SUC 92	1	1	
Lactobacillus acidophilus		19.05	0.38	RAF 92	99	1	
Clostridium barati		14.98	0.34		99	99	
05351400	**Low discrimination**						**05351400**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		81.16	0.41	SUC 90	1	
Lactobacillus catenaformis		14.70	0.29	ARA 1	1	
Clostridium innocuum		2.15	0.12		99	
05351402	**Low discrimination**					**05351402**
					SPO	
Lactobacillus jensenii		74.59	0.38	SUC 90	1	
Lactobacillus catenaformis		20.26	0.29	ARA 1	1	
Clostridium innocuum		2.96	0.12		99	
05351520	**Good identification**					**05351520**
					SPO	
Clostridium septicum		98.07	0.51	ARA 1	99	
Clostridium paraputrificum		0.89	0.14		99	
Lactobacillus catenaformis		0.63	0.21		1	
05352000	**Low discrimination**					**05352000**
					SPO	
Clostridium innocuum		89.68	0.37	MLT 1	99	
Clostridium difficile		6.94	0.15	GAL 95	99	
Lactobacillus jensenii		1.16	0.07		1	
05352002	**Good identification**					**05352002**
					SPO	
Clostridium innocuum		99.15	0.37	MLT 1	99	
Lactobacillus jensenii		0.86	0.04	GAL 95	1	
05353000	**Good identification**					**05353000**
					SPO	
Clostridium innocuum		92.31	0.59	MLT 1	99	
Lactobacillus jensenii		6.22	0.41		1	
Lactobacillus catenaformis		1.13	0.29		1	
05353002	**Good identification**					**05353002**
					SPO	
Clostridium innocuum		93.66	0.59	MLT 1	99	
Lactobacillus jensenii		4.21	0.38		1	
Lactobacillus catenaformis		1.14	0.29		1	
05353010	**Low discrimination**					**05353010**
					SPO	
Clostridium innocuum		64.98	0.35	MLT 1	99	
Lactobacillus jensenii		35.03	0.33	MLZ 4	1	
05353012	**Low discrimination**					**05353012**
					SPO	
Clostridium innocuum		73.56	0.35	MLT 1	99	
Lactobacillus jensenii		26.44	0.30	MLZ 4	1	
05353021	**Good identification**					**05353021**
Actinomyces naeslundii		100.0	0.35	SUC 99		
05353023	**Good identification**					**05353023**
Actinomyces naeslundii		100.0	0.35	SUC 99		
05353101	**Good identification**					**05353101**
Actinomyces naeslundii		100.0	0.43	SUC 99		
05353102	**Low discrimination**					**05353102**
					SPO	
Lactobacillus acidophilus		40.53	0.35	SUC 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		17.55	0.32	MAN 5	1		
Clostridium barati		15.19	0.25		99		
05353103	**Acceptable identification**						**05353103**
Actinomyces naeslundii		91.33	0.43	SUC 99			
Lactobacillus acidophilus		8.67	0.01	RAF 92			
05353120	**Acceptable identification**						**05353120**
Clostridium septicum		93.44	0.51	MAN 1			
Clostridium barati		2.08	0.25				
Clostridium tertium		1.58	0.27				
05353123	**Low discrimination**						**05353123**
					SPO		
Actinomyces naeslundii		77.96	0.43	SUC 99	1		
Clostridium barati		8.33	-.01	RAF 92	99		
Lactobacillus acidophilus		7.40	0.01		1		
05353132	**Good identification**						**05353132**
					SPO		
Clostridium tertium		98.03	0.39	SUC 99	99		
Clostridium barati		0.74	-.01	TRE 90	99		
Lactobacillus acidophilus		0.65	0.01		1		
05353400	**Good identification**						**05353400**
					SPO		
Clostridium innocuum		99.44	0.46	MLT 1	99		
Lactobacillus jensenii		0.38	0.07		1		
Actinomyces israelii		0.18	0.14		1		
05353402	**Good identification**						**05353402**
					SPO		
Clostridium innocuum		99.48	0.46	MLT 1	99		
Actinomyces israelii		0.27	0.17		1		
Lactobacillus jensenii		0.26	0.04		1		
05355000	**Low discrimination**						**05355000**
					GRA		
Lactobacillus jensenii		78.16	0.41	SUC 90	99		
Lactobacillus catenaformis		14.15	0.29	XYL 1	99		
Fusobacterium mortiferum		4.41	0.30		1		
05355002	**Low discrimination**						**05355002**
Lactobacillus jensenii		74.17	0.38	SUC 90			
Lactobacillus catenaformis		20.14	0.29	XYL 1			
Actinomyces israelii		1.94	0.24				
05355100	**Low discrimination**						**05355100**
					COC	GRA	
Atopobium parvulum		47.05	0.35	SUC 96	99	99	
Lactobacillus catenaformis		23.97	0.32	TRE 99	1	99	
Fusobacterium mortiferum		11.62	0.36		1	1	
05355120	**Good identification**						**05355120**
					COC	SPO	
Clostridium septicum		96.86	0.51	XYL 1	1	99	
Clostridium paraputrificum		0.88	0.14		1	99	
Atopobium parvulum		0.86	0.22		99	1	
05355122	**Low discrimination**						**05355122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Prevotella oralis		56.24	0.35	SUC 92	1	1	
Clostridium septicum		15.74	0.22	RAF 92	99	99	
Lactobacillus acidophilus		8.79	0.23	XYL 11	99	1	
05355502	**Low discrimination**						**05355502**
					SPO		
Bifidobacterium adolescentis		74.58	0.36	SUC 99	1		
Actinomyces israelii		19.98	0.32	RAF 92	1		
Clostridium butyricum		5.44	0.02		99		
05355512	**Acceptable identification**						**05355512**
Bifidobacterium adolescentis		94.55	0.37	SUC 99			
Actinomyces israelii		3.10	0.18	RAF 92			
Clostridium clostridioforme		2.35	0.07				
05357000	**Acceptable identification**						**05357000**
					SPO		
Clostridium innocuum		91.14	0.37	MLT 1	99		
Actinomyces israelii		7.68	0.33	XYL 5	1		
Lactobacillus jensenii		1.18	0.07		1		
05357002	**Low discrimination**						**05357002**
					SPO		
Clostridium innocuum		88.10	0.37	MLT 1	99		
Actinomyces israelii		11.14	0.37	XYL 5	1		
Lactobacillus jensenii		0.76	0.04		1		
05357100	**Low discrimination**						**05357100**
					SPO		
Actinomyces israelii		73.42	0.44	SUC 98	1		
Clostridium tertium		22.88	0.21		99		
Clostridium innocuum		2.20	0.03		99		
05357102	**Low discrimination**						**05357102**
					SPO		
Actinomyces israelii		57.97	0.47	SUC 98	1		
Clostridium tertium		36.13	0.29		99		
Bifidobacterium adolescentis		3.82	0.20		1		
05357122	**Acceptable identification**						**05357122**
					GRA	SPO	
Clostridium tertium		94.08	0.37	SUC 99	99	99	
Actinomyces israelii		3.21	0.26	TRE 90	99	1	
Prevotella oralis		0.99	0.01		1	1	
05357132	**Good identification**						**05357132**
Clostridium tertium		100.0	0.41	SUC 99			
05357140	**Good identification**						**05357140**
Actinomyces israelii		100.0	0.37	SUC 98			
05357142	**Good identification**						**05357142**
Actinomyces israelii		100.0	0.40	SUC 98			
05357500	**Good identification**						**05357500**
					SPO		
Actinomyces israelii		97.98	0.42	SUC 98	1		
Clostridium butyricum		2.02	-.08		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05357502	**Low discrimination**						**05357502**
Actinomyces israelii		64.98	0.45	SUC 98			
Bifidobacterium adolescentis		35.02	0.34				
05357512	**Low discrimination**						**05357512**
					SPO		
Bifidobacterium adolescentis		80.15	0.36	SUC 99	1		
Actinomyces israelii		18.18	0.31	RAF 92	1		
Clostridium tertium		1.67	-.01		99		
05357540	**Good identification**						**05357540**
Actinomyces israelii		100.0	0.35	SUC 98			
05357542	**Low discrimination**						**05357542**
Actinomyces israelii		85.38	0.38	SUC 98			
Bifidobacterium adolescentis		14.62	0.19				
05361000	**Good identification**						**05361000**
Lactobacillus jensenii		100.0	0.50	SUC 90			
05361002	**Good identification**						**05361002**
Lactobacillus jensenii		98.28	0.47	SUC 90			
Lactobacillus acidophilus		1.72	0.14	FRU 99			
05361010	**Good identification**						**05361010**
Lactobacillus jensenii		100.0	0.41	SUC 90			
05361012	**Good identification**						**05361012**
Lactobacillus jensenii		100.0	0.38	SUC 90			
05361020	**Low discrimination**						**05361020**
					SPO		
Lactobacillus jensenii		70.09	0.39	SUC 90	1		
Clostridium septicum		29.92	0.25	FRU 99	99		
05361022	**Acceptable identification**						**05361022**
					SPO		
Lactobacillus jensenii		92.32	0.36	SUC 90	1		
Lactobacillus acidophilus		6.47	0.14	FRU 99	1		
Clostridium septicum		1.21	-.04		99		
05361120	**Very good identification**						**05361120**
					COC	SPO	
Clostridium septicum		99.64	0.60	FRU 99	1	99	
Atopobium parvulum		0.16	0.18		99	1	
Clostridium perfringens		0.10	0.15		1	84	
05365100	**Good identification**						**05365100**
Atopobium parvulum		100.0	0.35	SUC 96			
05370000	**Acceptable identification**						**05370000**
					GRA		
Lactobacillus jensenii		94.98	0.50	SUC 90	99		
Fusobacterium mortiferum		4.73	0.37	GAL 99	1		
Prevotella disiens		0.29	-.03		1		
05370002	**Good identification**						**05370002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Lactobacillus jensenii		97.48	0.47	SUC 90	99			
Lactobacillus acidophilus		1.71	0.14	GAL 99	99			
Fusobacterium mortiferum		0.81	0.21		1			
05370010	**Good identification**							**05370010**
Lactobacillus jensenii		100.0	0.41	SUC 90				
05370012	**Good identification**							**05370012**
Lactobacillus jensenii		100.0	0.38	SUC 90				
05370020	**Low discrimination**							**05370020**
					SPO			
Lactobacillus jensenii		70.09	0.39	SUC 90	1			
Clostridium septicum		29.92	0.25	GAL 99	99			
05370022	**Acceptable identification**							**05370022**
					SPO			
Lactobacillus jensenii		90.26	0.36	SUC 90	1			
Lactobacillus acidophilus		6.33	0.14	GAL 99	1			
Clostridium barati		2.23	0.04		99			
05370100	**Low discrimination**							**05370100**
					COC	GRA	SPO	
Fusobacterium mortiferum		35.33	0.44	MLT 11	1	1	1	
Clostridium septicum		32.47	0.25		1	99	99	
Atopobium parvulum		29.12	0.31		99	99	1	
05370120	**Very good identification**							**05370120**
					COC	SPO		
Clostridium septicum		99.60	0.60	GAL 99	1	99		
Atopobium parvulum		0.16	0.18		99	1		
Clostridium perfringens		0.10	0.15		1	84		
05371000	**Very good identification**							**05371000**
					COC	SPO		
Lactobacillus jensenii		99.61	0.84	SUC 90	1	1		
Clostridium septicum		0.11	0.25		1	99		
Atopobium parvulum		0.10	0.31		99	1		
05371001	**Good identification**							**05371001**
Lactobacillus jensenii		99.23	0.50	SUC 90				
Actinomyces naeslundii		0.78	0.35	URE 1				
05371002	**Very good identification**							**05371002**
Lactobacillus jensenii		98.07	0.81	SUC 90				
Lactobacillus acidophilus		1.72	0.48					
Lactobacillus catenaformis		0.09	0.29					
05371003	**Good identification**							**05371003**
Lactobacillus jensenii		97.16	0.47	SUC 90				
Lactobacillus acidophilus		1.70	0.14	URE 1				
Actinomyces naeslundii		1.14	0.35					
05371010	**Very good identification**							**05371010**
Lactobacillus jensenii		100.0	0.76	SUC 90				
05371011	**Good identification**							**05371011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		100.0	0.41	SUC 90		
05371012	**Very good identification**					**05371012**
					GRA	
Lactobacillus jensenii		99.92	0.73	SUC 90	99	
Lactobacillus acidophilus		0.05	0.14		99	
Leptotrichia buccalis		0.02	-.03		1	
05371013	**Good identification**					**05371013**
Lactobacillus jensenii		100.0	0.38	SUC 90		
05371020	**Low discrimination**					**05371020**
					SPO	
Lactobacillus jensenii		69.65	0.73	SUC 90	1	
Clostridium septicum		29.73	0.60		99	
Clostridium perfringens		0.39	0.34		84	
05371021	**Low discrimination**					**05371021**
					SPO	
Lactobacillus jensenii		68.58	0.39	SUC 90	1	
Clostridium septicum		29.27	0.25	URE 1	99	
Actinomyces naeslundii		2.14	0.35		1	
05371022	**Good identification**					**05371022**
					SPO	
Lactobacillus jensenii		91.39	0.70	SUC 90	1	
Lactobacillus acidophilus		6.41	0.48		1	
Clostridium septicum		1.19	0.31		99	
05371023	**Low discrimination**					**05371023**
Lactobacillus jensenii		88.49	0.36	SUC 90		
Lactobacillus acidophilus		6.20	0.14	URE 1		
Actinomyces naeslundii		4.15	0.35			
05371030	**Very good identification**					**05371030**
					SPO	
Lactobacillus jensenii		98.72	0.65	SUC 90	1	
Clostridium septicum		1.28	0.25		99	
05371032	**Very good identification**					**05371032**
					SPO	
Lactobacillus jensenii		99.75	0.62	SUC 90	1	
Lactobacillus acidophilus		0.21	0.14		1	
Clostridium septicum		0.04	-.04		99	
05371040	**Good identification**					**05371040**
Lactobacillus jensenii		99.86	0.50	SUC 90		
Lactobacillus acidophilus		0.14	-.02	SOR 1		
05371042	**Low discrimination**					**05371042**
					SPO	
Lactobacillus jensenii		82.11	0.47	SUC 90	1	
Lactobacillus acidophilus		17.61	0.32	SOR 1	1	
Clostridium barati		0.28	-.01		99	
05371050	**Good identification**					**05371050**
Lactobacillus jensenii		100.0	0.41	SUC 90		
05371052	**Good identification**					**05371052**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		99.36	0.38	SUC 90			
Lactobacillus acidophilus		0.65	-.02	SOR 1			
05371060	**Low discrimination**						**05371060**
					SPO		
Lactobacillus jensenii		64.63	0.39	SUC 90	1		
Clostridium septicum		27.59	0.25	SOR 1	99		
Clostridium perfringens		6.83	0.22		84		
05371062	**Low discrimination**						**05371062**
Lactobacillus jensenii		50.35	0.36	SUC 90			
Lactobacillus acidophilus		43.18	0.32	SOR 1			
Clostridium perfringens		3.76	0.16				
05371100	**Low discrimination**						**05371100**
					COC	SPO	
Clostridium septicum		49.07	0.60	NAG 99	1	99	
Atopobium parvulum		44.00	0.65		99	1	
Lactobacillus jensenii		4.64	0.50		1	1	
05371102	**Low discrimination**						**05371102**
					COC		
Lactobacillus acidophilus		82.59	0.66	SUC 99	1		
Atopobium parvulum		10.38	0.52		99		
Lactobacillus jensenii		4.14	0.47		1		
05371110	**Low discrimination**						**05371110**
					COC	SPO	
Lactobacillus jensenii		61.08	0.41	SUC 90	1	1	
Clostridium septicum		19.55	0.25	LAC 1	1	99	
Atopobium parvulum		17.53	0.31		99	1	
05371120	**Excellent identification**						**05371120**
					COC	SPO	
Clostridium septicum		99.61	0.94		1	99	
Atopobium parvulum		0.16	0.52		99	1	
Clostridium perfringens		0.10	0.49		1	84	
05371121	**Very good identification**						**05371121**
					COC	SPO	
Clostridium septicum		99.43	0.60	URE 1	1	99	
Actinomyces naeslundii		0.21	0.43		1	1	
Atopobium parvulum		0.16	0.18		99	1	
05371122	**Low discrimination**						**05371122**
					SPO		
Clostridium septicum		58.11	0.65	BGL 2	99		
Lactobacillus acidophilus		36.21	0.66		1		
Clostridium perfringens		1.30	0.43		84		
05371130	**Very good identification**						**05371130**
					COC	SPO	
Clostridium septicum		98.80	0.60	MLZ 1	1	99	
Lactobacillus jensenii		0.78	0.31		1	1	
Atopobium parvulum		0.16	0.18		99	1	
05371142	**Good identification**						**05371142**
					COC		
Lactobacillus acidophilus		97.57	0.50	SUC 99	1		
Atopobium parvulum		1.00	0.18	SOR 11	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Lactobacillus jensenii	0.40	0.12		1		
05371160	**Very good identification**						**05371160**
	Clostridium septicum	97.21	0.60	SOR 1			
	Clostridium perfringens	1.78	0.37				
	Clostridium paraputrificum	0.59	0.19				
05371162	**Low discrimination**						**05371162**
					SPO		
	Lactobacillus acidophilus	81.82	0.50	SUC 99	1		
	Clostridium septicum	10.73	0.31	SOR 11	99		
	Clostridium perfringens	4.54	0.31		84		
05371200	**Good identification**						**05371200**
	Lactobacillus jensenii	100.0	0.50	SUC 90			
05371202	**Low discrimination**						**05371202**
					SPO		
	Lactobacillus jensenii	88.19	0.47	SUC 90	1		
	Lactobacillus acidophilus	11.52	0.29	RHA 1	1		
	Clostridium barati	0.30	-.01		99		
05371210	**Good identification**						**05371210**
	Lactobacillus jensenii	100.0	0.41	SUC 90			
05371212	**Good identification**						**05371212**
	Lactobacillus jensenii	100.0	0.38	SUC 90			
05371220	**Low discrimination**						**05371220**
					SPO		
	Lactobacillus jensenii	69.64	0.39	SUC 90	1		
	Clostridium septicum	29.73	0.25	RHA 1	99		
	Clostridium barati	0.63	-.01		99		
05371222	**Low discrimination**						**05371222**
					SPO		
	Lactobacillus jensenii	61.48	0.36	SUC 90	1		
	Lactobacillus acidophilus	32.11	0.29	RHA 1	1		
	Clostridium barati	2.51	0.07		99		
05371302	**Good identification**						**05371302**
					COC		
	Lactobacillus acidophilus	96.12	0.47	SUC 99	1		
	Atopobium parvulum	1.62	0.18	RHA 7	99		
	Lactobacillus jensenii	0.65	0.12		1		
05371320	**Very good identification**						**05371320**
					COC	SPO	
	Clostridium septicum	98.36	0.60	RHA 1	1	99	
	Clostridium ramosum	1.10	0.30		1	92	
	Atopobium parvulum	0.16	0.18		99	1	
05371322	**Low discrimination**						**05371322**
					SPO		
	Lactobacillus acidophilus	55.71	0.47	SUC 99	1		
	Clostridium ramosum	26.25	0.40	RHA 7	92		
	Clostridium septicum	12.00	0.31		99		
05371400	**Good identification**						**05371400**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		99.69	0.50	SUC 90	1		
Clostridium innocuum		0.31	0.04	ARA 1	99		
05371402	**Good identification**						**05371402**
					SPO		
Lactobacillus jensenii		97.61	0.47	SUC 90	1		
Lactobacillus acidophilus		1.71	0.14	ARA 1	1		
Clostridium innocuum		0.45	0.04		99		
05371410	**Good identification**						**05371410**
Lactobacillus jensenii		100.0	0.41	SUC 90			
05371412	**Good identification**						**05371412**
Lactobacillus jensenii		100.0	0.38	SUC 90			
05371420	**Low discrimination**						**05371420**
					SPO		
Lactobacillus jensenii		70.09	0.39	SUC 90	1		
Clostridium septicum		29.92	0.25	ARA 1	99		
05371422	**Acceptable identification**						**05371422**
					SPO		
Lactobacillus jensenii		92.32	0.36	SUC 90	1		
Lactobacillus acidophilus		6.47	0.14	ARA 1	1		
Clostridium septicum		1.21	-.04		99		
05371520	**Very good identification**						**05371520**
					COC	SPO	
Clostridium septicum		99.64	0.60	ARA 1	1	99	
Atopobium parvulum		0.16	0.18		99	1	
Clostridium perfringens		0.10	0.15		1	84	
05373000	**Low discrimination**						**05373000**
					SPO		
Clostridium innocuum		63.04	0.51	MLT 1	99		
Lactobacillus jensenii		36.27	0.50		1		
Actinomyces naeslundii		0.37	0.37		1		
05373002	**Low discrimination**						**05373002**
					SPO		
Clostridium innocuum		69.69	0.51	MLT 1	99		
Lactobacillus jensenii		26.73	0.47		1		
Lactobacillus acidophilus		2.44	0.26		1		
05373010	**Low discrimination**						**05373010**
					SPO		
Lactobacillus jensenii		81.51	0.41	SUC 90	1		
Clostridium innocuum		17.71	0.27	MAN 1	99		
Clostridium tertium		0.78	0.05		99		
05373012	**Low discrimination**						**05373012**
					SPO		
Lactobacillus jensenii		72.66	0.38	SUC 90	1		
Clostridium innocuum		23.68	0.27	MAN 1	99		
Clostridium tertium		3.14	0.13		99		
05373020	**Low discrimination**						**05373020**
					SPO		
Lactobacillus jensenii		62.95	0.39	SUC 90	1		
Clostridium septicum		26.87	0.25	MAN 1	99		
Clostridium innocuum		4.42	0.17		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05373021	**Low discrimination**					**05373021**
Actinomyces naeslundii		85.72	0.40	SUC 99		
Lactobacillus jensenii		14.28	0.05	RAF 92		
05373022	**Low discrimination**					**05373022**
					SPO	
Lactobacillus jensenii		55.42	0.36	SUC 90	1	
Lactobacillus acidophilus		20.24	0.26	MAN 1	1	
Clostridium ramosum		6.77	0.17		92	
05373023	**Acceptable identification**					**05373023**
Actinomyces naeslundii		90.01	0.40	SUC 99		
Lactobacillus jensenii		9.99	0.02	RAF 92		
05373101	**Good identification**					**05373101**
					SPO	
Actinomyces naeslundii		96.00	0.48	SUC 99	1	
Clostridium tertium		4.01	0.01	RAF 92	99	
05373102	**Low discrimination**					**05373102**
					SPO	
Lactobacillus acidophilus		50.83	0.44	SUC 99	1	
Clostridium tertium		38.83	0.43	MAN 5	99	
Actinomyces israelii		3.13	0.39		1	
05373103	**Low discrimination**					**05373103**
					SPO	
Actinomyces naeslundii		77.58	0.48	SUC 99	1	
Lactobacillus acidophilus		12.71	0.10	RAF 92	1	
Clostridium tertium		9.71	0.09		99	
05373110	**Good identification**					**05373110**
					SPO	
Clostridium tertium		97.19	0.39	SUC 99	99	
Actinomyces israelii		1.32	0.21		1	
Lactobacillus jensenii		1.03	0.07		1	
05373112	**Good identification**					**05373112**
					SPO	
Clostridium tertium		97.71	0.47	SUC 99	99	
Lactobacillus acidophilus		0.73	0.10		1	
Actinomyces israelii		0.66	0.24		1	
05373120	**Good identification**					**05373120**
Clostridium septicum		90.40	0.60	MAN 1		
Clostridium tertium		4.58	0.43			
Clostridium ramosum		4.31	0.41			
05373122	**Low discrimination**					**05373122**
					SPO	
Clostridium ramosum		43.82	0.51	SUC 99	92	
Clostridium tertium		34.87	0.51		99	
Lactobacillus acidophilus		15.21	0.44		1	
05373123	**Low discrimination**					**05373123**
					SPO	
Actinomyces naeslundii		48.51	0.48	SUC 99	1	
Clostridium ramosum		22.89	0.17	RAF 92	92	
Clostridium tertium		18.21	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05373130	**Low discrimination**						**05373130**
Clostridium tertium		89.35	0.47	SUC 99			
Clostridium septicum		10.03	0.25				
Clostridium ramosum		0.48	0.07				
05373132	**Good identification**						**05373132**
					SPO		
Clostridium tertium		98.92	0.55	SUC 99	99		
Clostridium ramosum		0.71	0.17		92		
Lactobacillus acidophilus		0.25	0.10		1		
05373320	**Low discrimination**						**05373320**
Clostridium ramosum		85.63	0.43	SUC 99			
Clostridium septicum		13.68	0.25				
Clostridium tertium		0.69	0.09				
05373322	**Good identification**						**05373322**
					SPO		
Clostridium ramosum		97.35	0.53	SUC 99	92		
Lactobacillus acidophilus		1.92	0.25		1		
Clostridium tertium		0.59	0.17		99		
05373400	**Good identification**						**05373400**
					SPO		
Clostridium innocuum		95.05	0.38	MLT 1	99		
Lactobacillus jensenii		3.13	0.15		1		
Actinomyces israelii		1.82	0.24		1		
05373402	**Good identification**						**05373402**
					SPO		
Clostridium innocuum		95.18	0.38	MLT 1	99		
Actinomyces israelii		2.73	0.27		1		
Lactobacillus jensenii		2.09	0.12		1		
05373502	**Low discrimination**						**05373502**
Actinomyces israelii		62.27	0.37	SUC 98			
Lactobacillus acidophilus		12.50	0.10	XYL 92			
Bifidobacterium adolescentis		10.22	0.17				
05374100	**Good identification**						**05374100**
					COC	GRA	
Atopobium parvulum		95.31	0.35	SUC 96	99	99	
Fusobacterium mortiferum		4.69	0.24	GAL 99	1	1	
05375000	**Low discrimination**						**05375000**
					COC		
Lactobacillus jensenii		83.48	0.50	SUC 90	1		
Atopobium parvulum		14.84	0.35	XYL 1	99		
Actinomyces israelii		1.22	0.30		1		
05375002	**Acceptable identification**						**05375002**
					COC		
Lactobacillus jensenii		91.13	0.47	SUC 90	1		
Atopobium parvulum		4.29	0.22	XYL 1	99		
Actinomyces israelii		2.99	0.33		1		
05375010	**Good identification**						**05375010**
					COC		
Lactobacillus jensenii		98.82	0.41	SUC 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.65	0.16	XYL 1	1		
Atopobium parvulum		0.53	0.01		99		
05375012	**Good identification**						**05375012**
Lactobacillus jensenii		98.55	0.38	SUC 90			
Actinomyces israelii		1.45	0.19	XYL 1			
05375020	**Low discrimination**						**05375020**
					COC	SPO	
Lactobacillus jensenii		64.42	0.39	SUC 90	1	1	
Clostridium septicum		27.50	0.25	XYL 1	1	99	
Atopobium parvulum		8.08	0.22		99	1	
05375022	**Low discrimination**						**05375022**
					COC		
Lactobacillus jensenii		89.58	0.36	SUC 90	1		
Lactobacillus acidophilus		6.28	0.14	XYL 1	1		
Atopobium parvulum		2.97	0.09		99		
05375100	**Very good identification**						**05375100**
					COC	SPO	
Atopobium parvulum		98.80	0.69	SUC 96	99	1	
Clostridium septicum		0.60	0.25		1	99	
Actinomyces israelii		0.33	0.41		1	1	
05375101	**Good identification**						**05375101**
Atopobium parvulum		100.0	0.35	SUC 96			
05375102	**Good identification**						**05375102**
					COC		
Atopobium parvulum		92.39	0.57	SUC 96	99		
Lactobacillus acidophilus		4.00	0.32		1		
Actinomyces israelii		2.60	0.44		1		
05375110	**Acceptable identification**						**05375110**
					COC	SPO	
Atopobium parvulum		91.51	0.35	SUC 96	99	1	
Actinomyces israelii		4.48	0.27	MLZ 1	1	1	
Clostridium tertium		2.28	0.07		1	99	
05375120	**Low discrimination**						**05375120**
					COC	SPO	
Clostridium septicum		77.09	0.60	XYL 1	1	99	
Atopobium parvulum		22.66	0.57		99	1	
Clostridium perfringens		0.08	0.15		1	84	
05375122	**Low discrimination**						**05375122**
					COC	SPO	
Atopobium parvulum		54.08	0.44	SUC 96	99	1	
Clostridium septicum		21.28	0.31		1	99	
Lactobacillus acidophilus		13.26	0.32		1	1	
05375140	**Low discrimination**						**05375140**
					COC	SPO	
Atopobium parvulum		87.81	0.35	SUC 96	99	1	
Actinomyces israelii		11.20	0.34	SOR 1	1	1	
Clostridium butyricum		1.00	-.05		1	99	
05375142	**Low discrimination**						**05375142**
					COC		
Actinomyces israelii		41.48	0.37	SUC 98	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		38.26	0.22		99		
Lactobacillus acidophilus		20.26	0.16		1		
05375300	**Good identification**						**05375300**
					COC		
Atopobium parvulum		97.59	0.35	SUC 96	99		
Actinomyces israelii		2.41	0.22	RHA 1	1		
05375322	**Low discrimination**						**05375322**
					GRA		
Parabacteroides distasonis		84.01	0.45	RAF 99	1		
Lactobacillus acidophilus		5.18	0.12		99		
Prevotella oralis		3.60	0.09		1		
05375332	**Acceptable identification**						**05375332**
Parabacteroides distasonis		92.26	0.40	RAF 99			
Clostridium clostridioforme		7.74	0.16				
05375542	**Low discrimination**						**05375542**
					SPO		
Actinomyces israelii		68.32	0.35	SUC 98	1		
Clostridium butyricum		17.32	0.04		99		
Bifidobacterium adolescentis		14.36	0.17		1		
05375712	**Good identification**						**05375712**
Clostridium clostridioforme		100.0	0.36	SUC 99			
05375722	**Low discrimination**						**05375722**
Parabacteroides distasonis		79.79	0.41	RAF 99			
Clostridium clostridioforme		20.21	0.25				
05375732	**Low discrimination**						**05375732**
Clostridium clostridioforme		68.18	0.36	SUC 99			
Parabacteroides distasonis		31.82	0.36	RAF 94			
05377002	**Low discrimination**						**05377002**
					SPO		
Actinomyces israelii		53.95	0.46	SUC 98	1		
Clostridium innocuum		39.86	0.29		99		
Clostridium tertium		3.25	0.11		99		
05377040	**Good identification**						**05377040**
Actinomyces israelii		100.0	0.36	SUC 98			
05377042	**Good identification**						**05377042**
Actinomyces israelii		100.0	0.39	SUC 98			
05377100	**Low discrimination**						**05377100**
					COC	SPO	
Actinomyces israelii		43.64	0.54	SUC 98	1	1	
Clostridium tertium		32.53	0.37		1	99	
Atopobium parvulum		23.49	0.35		99	1	
05377102	**Low discrimination**						**05377102**
					COC	SPO	
Clostridium tertium		57.78	0.45	SUC 99	1	99	
Actinomyces israelii		38.76	0.57		1	1	
Atopobium parvulum		2.45	0.22		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05377110	**Low discrimination**						**05377110**
					COC	SPO	
Clostridium tertium		89.54	0.41	SUC 99	1	99	
Actinomyces israelii		10.10	0.40		1	1	
Atopobium parvulum		0.37	0.01		99	1	
05377112	**Acceptable identification**						**05377112**
					SPO		
Clostridium tertium		94.18	0.50	SUC 99	99		
Actinomyces israelii		5.31	0.43		1		
Bifidobacterium adolescentis		0.51	0.19		1		
05377120	**Low discrimination**						**05377120**
					COC	SPO	
Clostridium tertium		81.81	0.45	SUC 99	1	99	
Clostridium septicum		11.82	0.25		1	99	
Atopobium parvulum		3.47	0.22		99	1	
05377122	**Good identification**						**05377122**
					SPO		
Clostridium tertium		97.08	0.53	SUC 99	99		
Actinomyces israelii		1.39	0.36		1		
Clostridium ramosum		0.89	0.17		92		
05377130	**Good identification**						**05377130**
					SPO		
Clostridium tertium		99.76	0.50	SUC 99	99		
Actinomyces israelii		0.24	0.19		1		
05377132	**Very good identification**						**05377132**
					SPO		
Clostridium tertium		99.88	0.58	SUC 99	99		
Actinomyces israelii		0.12	0.22		1		
05377140	**Good identification**						**05377140**
					COC	SPO	
Actinomyces israelii		96.77	0.47	SUC 98	1	1	
Clostridium tertium		1.87	0.03		1	99	
Atopobium parvulum		1.35	0.01		99	1	
05377142	**Good identification**						**05377142**
					SPO		
Actinomyces israelii		96.27	0.50	SUC 98	1		
Clostridium tertium		3.73	0.11		99		
05377152	**Low discrimination**						**05377152**
					SPO		
Actinomyces israelii		68.46	0.36	SUC 98	1		
Clostridium tertium		31.54	0.15	MLZ 13	99		
05377302	**Low discrimination**						**05377302**
					SPO		
Actinomyces israelii		83.33	0.37	SUC 98	1		
Clostridium tertium		16.67	0.11	RHA 7	99		
05377400	**Low discrimination**						**05377400**
					SPO		
Actinomyces israelii		80.71	0.42	SUC 98	1		
Clostridium innocuum		19.29	0.16		99		
05377402	**Low discrimination**						**05377402**
					SPO		
Actinomyces israelii		86.26	0.45	SUC 98	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		13.74	0.16		99		
05377442	**Good identification**						**05377442**
Actinomyces israelii		100.0	0.38	SUC 98			
05377500	**Acceptable identification**						**05377500**
					SPO		
Actinomyces israelii		90.78	0.52	SUC 98	1		
Clostridium butyricum		7.78	0.13		99		
Clostridium tertium		0.84	0.03		99		
05377502	**Low discrimination**						**05377502**
					SPO		
Actinomyces israelii		88.02	0.55	SUC 98	1		
Bifidobacterium adolescentis		8.41	0.31		1		
Clostridium butyricum		1.96	0.06		99		
05377510	**Acceptable identification**						**05377510**
					SPO		
Actinomyces israelii		90.13	0.38	SUC 98	1		
Clostridium tertium		9.87	0.07	MLZ 13	99		
05377512	**Low discrimination**						**05377512**
Actinomyces israelii		45.00	0.41	SUC 98			
Bifidobacterium adolescentis		35.15	0.33	MLZ 13			
Bifidobacterium dentium		10.00	0.06				
05377540	**Good identification**						**05377540**
					SPO		
Actinomyces israelii		98.12	0.45	SUC 98	1		
Clostridium butyricum		1.88	-.05		99		
05377542	**Good identification**						**05377542**
Actinomyces israelii		97.05	0.48	SUC 98			
Bifidobacterium adolescentis		2.95	0.16				
05377702	**Good identification**						**05377702**
Actinomyces israelii		100.0	0.36	SUC 98			
05400000	**Low discrimination**						**05400000**
					COC		
Atopobium minutum		19.15	0.56	RAF 1	1		
Gemella morbillorum		18.74	0.52		99		
Anaerococcus prevotii		15.68	0.41		99		
05400001	**Very good identification**						**05400001**
					COC	SPO	
Anaerococcus prevotii		99.11	0.58	GLU 1	99	1	
Clostridium novyi biovar A		0.28	0.32		1	99	
Atopobium minutum		0.14	0.22		1	1	
05400002	**Low discrimination**						**05400002**
					COC	CAT	
Atopobium minutum		71.07	0.53	RAF 1	1	1	
Anaerococcus prevotii		21.82	0.31		99	40	
Propionibacterium granulosum		1.77	0.28		1	90	
05400003	**Good identification**						**05400003**
					COC	SPO	
Anaerococcus prevotii		99.44	0.47	GLU 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		0.36	0.19		1	1	
Clostridium novyi biovar A		0.20	0.19		1	99	
05400020	**Low discrimination**						**05400020**
					COC	SPO	
Atopobium minutum		47.14	0.48	RAF 1	1	1	
Clostridium cadaveris		27.60	0.34		1	99	
Gemella morbillorum		15.38	0.35		99	1	
05400022	**Acceptable identification**						**05400022**
					GRA	CAT	
Atopobium minutum		94.99	0.45	RAF 1	99	1	
Propionibacterium granulosum		1.77	0.18		99	90	
Pseudoflavonifractor capillosus		1.04	-.03		1	1	
05400100	**Low discrimination**						**05400100**
					COC	GRA	
Fusobacterium mortiferum		55.38	0.42	FRU 99	1	1	
Atopobium minutum		8.12	0.22		1	99	
Gemella morbillorum		7.95	0.17		99	99	
05400400	**Low discrimination**						**05400400**
					COC		
Gemella morbillorum		73.47	0.35	RAF 1	99		
Atopobium minutum		6.82	0.22	ARA 10	1		
Anaerococcus prevotii		5.59	0.07		99		
05401000	**Low discrimination**						**05401000**
					COC	CAT	
Gemella morbillorum		50.90	0.47	RAF 1	99	1	
Atopobium minutum		32.19	0.48		1	1	
Propionibacterium granulosum		6.41	0.38		1	90	
05401002	**Low discrimination**						**05401002**
					CAT		
Atopobium minutum		83.78	0.45	RAF 1	1		
Propionibacterium granulosum		6.26	0.28		90		
Lactobacillus fermentum		5.32	0.32		1		
05401020	**Low discrimination**						**05401020**
					COC	CAT	
Atopobium minutum		57.22	0.40	RAF 1	1	1	
Gemella morbillorum		30.16	0.31		99	1	
Propionibacterium granulosum		8.55	0.28		1	90	
05401022	**Low discrimination**						**05401022**
					CAT		
Atopobium minutum		89.32	0.37	RAF 1	1		
Lactobacillus fermentum		5.68	0.24		1		
Propionibacterium granulosum		5.01	0.18		90		
05401102	**Low discrimination**						**05401102**
					COC	GRA	
Lactobacillus fermentum		40.80	0.35	SUC 87	1	99	
Blautia hansenii		34.89	0.15	FRU 99	99	99	
Fusobacterium mortiferum		20.33	0.29		1	1	
05401502	**Low discrimination**						**05401502**
					COC		
Lactobacillus fermentum		86.89	0.35	SUC 87	1		
Blautia hansenii		13.11	0.03	FRU 99	99		
05403000	**Low discrimination**						**05403000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	CAT		
Propionibacterium granulosum		83.46	0.36	SUC 93	1	90		
Gemella morbillorum		8.87	0.13	FRU 99	99	1		
Atopobium minutum		5.61	0.14		1	1		
05404000	**Low discrimination**							**05404000**
					COC			
Gemella morbillorum		70.69	0.35	RAF 1	99			
Atopobium minutum		6.57	0.22	XYL 10	1			
Anaerococcus prevotii		5.38	0.07		99			
05410000	**Low discrimination**							**05410000**
					GRA	CAT		
Fusobacterium varium		34.24	0.58	RAF 1	1	1		
Propionibacterium granulosum		26.38	0.73		99	90		
Fusobacterium mortiferum		11.73	0.70		1	5		
05410002	**Low discrimination**							**05410002**
					GRA	CAT		
Propionibacterium granulosum		59.29	0.62	SUC 93	99	90		
Atopobium minutum		22.20	0.53		99	1		
Fusobacterium mortiferum		11.71	0.54		1	5		
05410010	**Low discrimination**							**05410010**
					GRA	SPO	CAT	
Propionibacterium granulosum		55.55	0.58	SUC 93	99	1	90	
Clostridium difficile		32.96	0.41	MLZ 12	99	99	1	
Fusobacterium varium		5.34	0.24		1	1	1	
05410012	**Acceptable identification**							**05410012**
					SPO	CAT		
Propionibacterium granulosum		93.78	0.48	SUC 93	1	90		
Atopobium minutum		2.60	0.19	MLZ 12	1	1		
Clostridium difficile		2.25	0.07		99	1		
05410020	**Low discrimination**							**05410020**
					SPO	CAT		
Propionibacterium granulosum		63.04	0.62	SUC 93	1	90		
Atopobium minutum		11.80	0.48		1	1		
Clostridium cadaveris		7.49	0.34		99	1		
05410022	**Low discrimination**							**05410022**
					CAT			
Propionibacterium granulosum		64.89	0.52	SUC 93	90			
Atopobium minutum		32.39	0.45		1			
Lactobacillus fermentum		0.74	0.24		1			
05410030	**Acceptable identification**							**05410030**
					SPO	CAT		
Propionibacterium granulosum		94.51	0.48	SUC 93	1	90		
Clostridium difficile		2.27	0.07	MLZ 12	99	1		
Atopobium minutum		1.31	0.14		1	1		
05410032	**Good identification**							**05410032**
					CAT			
Propionibacterium granulosum		96.43	0.37	SUC 93	90			
Atopobium minutum		3.57	0.11	MLZ 12	1			
05410100	**Very good identification**							**05410100**
					GRA	CAT		
Fusobacterium mortiferum		96.52	0.76		1	5		
Fusobacterium varium		1.22	0.24		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.94	0.38		99	90		
05410101	**Good identification**							**05410101**
Fusobacterium mortiferum		100.0	0.42	URE 1				
05410102	**Good identification**							**05410102**
					GRA	CAT		
Fusobacterium mortiferum		94.29	0.60	BGL 10	1	5		
Lactobacillus fermentum		2.87	0.35		99	1		
Propionibacterium granulosum		2.07	0.28		99	90		
05410110	**Low discrimination**							**05410110**
					GRA	SPO	CAT	
Fusobacterium mortiferum		82.68	0.42	MLZ 1	1	1	5	
Propionibacterium granulosum		10.87	0.24		99	1	90	
Clostridium difficile		6.45	0.07		99	99	1	
05410120	**Low discrimination**							**05410120**
					GRA	CAT		
Fusobacterium mortiferum		45.13	0.42	NAG 1	1	5		
Prevotella bivia		32.74	0.15		1	1		
Propionibacterium granulosum		10.88	0.28		99	90		
05410140	**Good identification**							**05410140**
Fusobacterium mortiferum		100.0	0.42	SOR 1				
05410200	**Low discrimination**							**05410200**
					GRA	CAT		
Propionibacterium granulosum		52.46	0.47	SUC 93	99	90		
Fusobacterium varium		22.24	0.24	RHA 3	1	1		
Fusobacterium mortiferum		7.62	0.36		1	5		
05410202	**Low discrimination**							**05410202**
					GRA	CAT		
Propionibacterium granulosum		84.26	0.37	SUC 93	99	90		
Atopobium minutum		10.30	0.19	RHA 3	99	1		
Fusobacterium mortiferum		5.44	0.19		1	5		
05410220	**Low discrimination**							**05410220**
					SPO	CAT		
Propionibacterium granulosum		86.93	0.37	SUC 93	1	90		
Atopobium minutum		5.32	0.14	RHA 3	1	1		
Clostridium cadaveris		3.37	0.00		99	1		
05410300	**Good identification**							**05410300**
					GRA	CAT		
Fusobacterium mortiferum		97.11	0.42	RHA 1	1	5		
Propionibacterium granulosum		2.90	0.13		99	90		
05410400	**Low discrimination**							**05410400**
					COC	GRA	CAT	
Gemella morbillorum		34.47	0.37	RAF 1	99	99	1	
Fusobacterium varium		22.35	0.24	ARA 10	1	1	1	
Propionibacterium granulosum		17.22	0.38		1	99	90	
05410500	**Low discrimination**							**05410500**
					COC	GRA		
Fusobacterium mortiferum		73.86	0.42	ARA 1	1	1		
Lactobacillus fermentum		24.70	0.35		1	99		
Gemella morbillorum		1.44	0.03		99	99		
05410502	**Low discrimination**							**05410502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Lactobacillus fermentum		75.06	0.35	SUC 87	99		
Fusobacterium mortiferum		24.94	0.26	GAL 99	1		
05411000	**Low discrimination**						**05411000**
					GRA	CAT	
Propionibacterium granulosum		43.53	0.73	SUC 93	99	90	
Fusobacterium mortiferum		29.03	0.73		1	5	
Prevotella intermedia		11.10	0.55		1	1	
05411001	**Low discrimination**						**05411001**
					GRA	CAT	
Propionibacterium granulosum		43.87	0.38	SUC 93	99	90	
Fusobacterium mortiferum		29.26	0.39	URE 1	1	5	
Prevotella intermedia		11.19	0.20		1	1	
05411002	**Low discrimination**						**05411002**
					GRA	CAT	
Propionibacterium granulosum		43.75	0.62	SUC 93	99	90	
Lactobacillus fermentum		37.21	0.66		99	1	
Fusobacterium mortiferum		12.97	0.57		1	5	
05411010	**Good identification**						**05411010**
					GRA	CAT	
Propionibacterium granulosum		90.85	0.58	SUC 93	99	90	
Fusobacterium mortiferum		4.49	0.39	MLZ 12	1	5	
Prevotella intermedia		1.72	0.20		1	1	
05411012	**Acceptable identification**						**05411012**
					GRA	CAT	
Propionibacterium granulosum		91.39	0.48	SUC 93	99	90	
Lactobacillus fermentum		5.76	0.32	MLZ 12	99	1	
Fusobacterium mortiferum		2.01	0.22		1	5	
05411020	**Low discrimination**						**05411020**
					CAT		
Propionibacterium granulosum		66.29	0.62	SUC 93	90		
Lactobacillus fermentum		18.79	0.58		1		
Propionibacterium acnes		4.87	0.34		95		
05411022	**Low discrimination**						**05411022**
					CAT		
Lactobacillus fermentum		48.65	0.58	SUC 87	1		
Propionibacterium granulosum		42.90	0.52		90		
Atopobium minutum		7.14	0.37		1		
05411030	**Good identification**						**05411030**
					CAT		
Propionibacterium granulosum		96.99	0.48	SUC 93	90		
Lactobacillus fermentum		2.04	0.24	MLZ 12	1		
Propionibacterium acnes		0.53	-.01		95		
05411032	**Acceptable identification**						**05411032**
					CAT		
Propionibacterium granulosum		92.25	0.37	SUC 93	90		
Lactobacillus fermentum		7.75	0.24	MLZ 12	1		
05411040	**Low discrimination**						**05411040**
					GRA	CAT	
Propionibacterium granulosum		43.87	0.38	SUC 93	99	90	
Fusobacterium mortiferum		29.26	0.39	SOR 1	1	5	
Prevotella intermedia		11.19	0.20		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05411100	**Low discrimination**						**05411100**
					GRA	CAT	
Fusobacterium mortiferum		81.12	0.79		1	5	
Lactobacillus fermentum		18.09	0.70		99	1	
Propionibacterium granulosum		0.53	0.38		99	90	
05411101	**Low discrimination**						**05411101**
					GRA		
Fusobacterium mortiferum		80.94	0.45	URE 1	1		
Lactobacillus fermentum		18.05	0.35		99		
Actinomyces naeslundii		1.01	0.24		99		
05411102	**Low discrimination**						**05411102**
					GRA	CAT	
Lactobacillus fermentum		66.33	0.70	SUC 87	99	1	
Fusobacterium mortiferum		33.06	0.63		1	5	
Propionibacterium granulosum		0.48	0.28		99	90	
05411103	**Low discrimination**						**05411103**
					GRA		
Lactobacillus fermentum		64.33	0.35	SUC 87	99		
Fusobacterium mortiferum		32.06	0.29	URE 1	1		
Actinomyces naeslundii		3.61	0.24		99		
05411110	**Low discrimination**						**05411110**
					GRA	CAT	
Fusobacterium mortiferum		76.30	0.45	MLZ 1	1	5	
Lactobacillus fermentum		17.01	0.35		99	1	
Propionibacterium granulosum		6.69	0.24		99	90	
05411112	**Low discrimination**						**05411112**
					GRA	CAT	
Lactobacillus fermentum		62.63	0.35	SUC 87	99	1	
Fusobacterium mortiferum		31.21	0.29	MLZ 1	1	5	
Propionibacterium granulosum		6.15	0.13		99	90	
05411120	**Low discrimination**						**05411120**
					GRA		
Lactobacillus fermentum		65.39	0.61	SUC 87	99		
Prevotella melaninogenica		15.83	0.32		1		
Fusobacterium mortiferum		8.89	0.45		1		
05411122	**Low discrimination**						**05411122**
					GRA		
Lactobacillus fermentum		82.71	0.61	SUC 87	99		
Parabacteroides distasonis		5.02	0.38		1		
Prevotella melaninogenica		3.81	0.19		1		
05411140	**Low discrimination**						**05411140**
					GRA		
Fusobacterium mortiferum		81.77	0.45	SOR 1	1		
Lactobacillus fermentum		18.23	0.35		99		
05411142	**Low discrimination**						**05411142**
					GRA		
Lactobacillus fermentum		66.74	0.35	SUC 87	99		
Fusobacterium mortiferum		33.26	0.29	SOR 1	1		
05411200	**Low discrimination**						**05411200**
					GRA	CAT	
Propionibacterium granulosum		70.53	0.47	SUC 93	99	90	
Fusobacterium mortiferum		15.36	0.39	RHA 3	1	5	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		5.88	0.20		1	1	
05411202	**Low discrimination**						**05411202**
					GRA	CAT	
Propionibacterium granulosum		70.65	0.37	SUC 93	99	90	
Lactobacillus fermentum		19.63	0.32	RHA 3	99	1	
Fusobacterium mortiferum		6.84	0.22		1	5	
05411220	**Low discrimination**						**05411220**
					CAT		
Propionibacterium granulosum		87.95	0.37	SUC 93	90		
Lactobacillus fermentum		8.14	0.24	RHA 3	1		
Propionibacterium acnes		2.11	-.01		95		
05411300	**Low discrimination**						**05411300**
					GRA	CAT	
Fusobacterium mortiferum		80.46	0.45	RHA 1	1	5	
Lactobacillus fermentum		17.94	0.35		99	1	
Propionibacterium granulosum		1.60	0.13		99	90	
05411302	**Low discrimination**						**05411302**
					GRA		
Lactobacillus fermentum		64.14	0.35	SUC 87	99		
Fusobacterium mortiferum		31.96	0.29	RHA 1	1		
Parabacteroides distasonis		3.90	0.12		1		
05411322	**Low discrimination**						**05411322**
					GRA		
Parabacteroides distasonis		87.64	0.46	XYL 90	1		
Prevotella oralis		7.51	0.15		1		
Lactobacillus fermentum		4.86	0.27		99		
05411332	**Good identification**						**05411332**
Parabacteroides distasonis		100.0	0.41	XYL 90			
05411400	**Low discrimination**						**05411400**
					COC	CAT	
Lactobacillus fermentum		87.36	0.66	SUC 87	1	1	
Gemella morbillorum		4.47	0.32		99	1	
Propionibacterium granulosum		4.15	0.38		1	90	
05411402	**Very good identification**						**05411402**
					GRA	CAT	
Lactobacillus fermentum		98.34	0.66	SUC 87	99	1	
Propionibacterium granulosum		1.17	0.28		99	90	
Fusobacterium mortiferum		0.35	0.22		1	5	
05411420	**Good identification**						**05411420**
					COC	CAT	
Lactobacillus fermentum		94.56	0.58	SUC 87	1	1	
Propionibacterium granulosum		3.37	0.28		1	90	
Gemella morbillorum		1.61	0.16		99	1	
05411422	**Very good identification**						**05411422**
					CAT		
Lactobacillus fermentum		99.12	0.58	SUC 87	1		
Propionibacterium granulosum		0.88	0.18		90		
05411500	**Very good identification**						**05411500**
					GRA		
Lactobacillus fermentum		95.60	0.70	SUC 87	99		
Fusobacterium mortiferum		4.33	0.45		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		0.07	0.09		99	
05411501	**Good identification**					**05411501**
					GRA	
Lactobacillus fermentum		95.67	0.35	SUC 87	99	
Fusobacterium mortiferum		4.34	0.11	URE 1	1	
05411502	**Very good identification**					**05411502**
					GRA	
Lactobacillus fermentum		99.42	0.70	SUC 87	99	
Fusobacterium mortiferum		0.50	0.29		1	
Bifidobacterium longum subsp. longum		0.08	0.09		99	
05411503	**Good identification**					**05411503**
Lactobacillus fermentum		100.0	0.35	SUC 87		
05411510	**Low discrimination**					**05411510**
					GRA	
Lactobacillus fermentum		85.01	0.35	SUC 87	99	
Bifidobacterium longum subsp. longum		11.14	0.13	MLZ 1	99	
Fusobacterium mortiferum		3.85	0.11		1	
05411512	**Low discrimination**					**05411512**
Lactobacillus fermentum		88.42	0.35	SUC 87		
Bifidobacterium longum subsp. longum		11.58	0.13	MLZ 1		
05411520	**Very good identification**					**05411520**
					GRA	
Lactobacillus fermentum		98.34	0.61	SUC 87	99	
Bifidobacterium longum subsp. longum		0.92	0.20		99	
Odoribacter splanchnicus		0.36	-.03		1	
05411522	**Very good identification**					**05411522**
					GRA	
Lactobacillus fermentum		95.55	0.61	SUC 87	99	
Parabacteroides distasonis		3.56	0.34		1	
Bifidobacterium longum subsp. longum		0.89	0.20		99	
05411540	**Good identification**					**05411540**
					GRA	
Lactobacillus fermentum		95.67	0.35	SUC 87	99	
Fusobacterium mortiferum		4.34	0.11	SOR 1	1	
05411542	**Good identification**					**05411542**
Lactobacillus fermentum		100.0	0.35	SUC 87		
05411700	**Good identification**					**05411700**
					GRA	
Lactobacillus fermentum		95.67	0.35	SUC 87	99	
Fusobacterium mortiferum		4.34	0.11	RHA 1	1	
05411702	**Good identification**					**05411702**
Lactobacillus fermentum		100.0	0.35	SUC 87		
05411722	**Acceptable identification**					**05411722**
					GRA	
Parabacteroides distasonis		91.70	0.42	XYL 90	1	
Lactobacillus fermentum		8.30	0.27		99	
05411732	**Good identification**					**05411732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.38	XYL 90			
05412000	**Low discrimination**						**05412000**
					SPO	CAT	
Eubacterium limosum		65.99	0.63	RAF 1	1	1	
Propionibacterium granulosum		26.97	0.71		1	90	
Clostridium difficile		5.33	0.46		99	1	
05412002	**Low discrimination**						**05412002**
					CAT		
Propionibacterium granulosum		88.63	0.60	SUC 93	90		
Eubacterium limosum		8.76	0.29		1		
Pseudoramibacter alactolyticus		1.02	0.11		1		
05412010	**Low discrimination**						**05412010**
					SPO	CAT	
Clostridium difficile		85.68	0.58	RAF 1	99	1	
Propionibacterium granulosum		12.11	0.56		1	90	
Eubacterium limosum		2.19	0.29		1	1	
05412012	**Low discrimination**						**05412012**
					SPO	CAT	
Propionibacterium granulosum		77.32	0.45	SUC 93	1	90	
Clostridium difficile		22.11	0.24	MLZ 12	99	1	
Eubacterium limosum		0.57	-.06		1	1	
05412020	**Low discrimination**						**05412020**
					CAT		
Propionibacterium granulosum		84.38	0.60	SUC 93	90		
Eubacterium limosum		8.34	0.29		1		
Propionibacterium acnes		4.62	0.29		95		
05412022	**Good identification**						**05412022**
					CAT		
Propionibacterium granulosum		98.15	0.50	SUC 93	90		
Pseudoramibacter alactolyticus		0.80	-.02		1		
Atopobium minutum		0.66	0.11		1		
05412030	**Low discrimination**						**05412030**
					SPO	CAT	
Propionibacterium granulosum		77.32	0.45	SUC 93	1	90	
Clostridium difficile		22.11	0.24	MLZ 12	99	1	
Eubacterium limosum		0.57	-.06		1	1	
05412032	**Good identification**						**05412032**
Propionibacterium granulosum		100.0	0.35	SUC 93			
05412050	**Good identification**						**05412050**
					SPO	CAT	
Clostridium difficile		96.90	0.36	RAF 1	99	1	
Propionibacterium granulosum		2.63	0.22	SOR 5	1	90	
Eubacterium limosum		0.48	-.06		1	1	
05412200	**Low discrimination**						**05412200**
					SPO	CAT	
Propionibacterium granulosum		53.42	0.45	SUC 93	1	90	
Eubacterium limosum		42.68	0.29	RHA 3	1	1	
Clostridium difficile		3.45	0.12		99	1	
05413000	**Very good identification**						**05413000**
					GRA	CAT	
Propionibacterium granulosum		95.78	0.71	SUC 93	99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		2.37	0.29		99	1	
Fusobacterium mortiferum		0.86	0.39		1	5	
05413001	**Good identification**						**05413001**
					CAT		
Propionibacterium granulosum		97.59	0.36	SUC 93	90		
Eubacterium limosum		2.41	-.06	URE 1	1		
05413002	**Very good identification**						**05413002**
					GRA	CAT	
Propionibacterium granulosum		98.23	0.60	SUC 93	99	90	
Lactobacillus fermentum		1.12	0.32		99	1	
Fusobacterium mortiferum		0.39	0.22		1	5	
05413010	**Acceptable identification**						**05413010**
					SPO	CAT	
Propionibacterium granulosum		93.17	0.56	SUC 93	1	90	
Clostridium difficile		6.66	0.24	MLZ 12	99	1	
Eubacterium limosum		0.17	-.06		1	1	
05413012	**Good identification**						**05413012**
Propionibacterium granulosum		100.0	0.45	SUC 93			
05413020	**Good identification**						**05413020**
					CAT		
Propionibacterium granulosum		94.31	0.60	SUC 93	90		
Propionibacterium acnes		5.16	0.29		95		
Lactobacillus fermentum		0.36	0.24		1		
05413022	**Good identification**						**05413022**
					CAT		
Propionibacterium granulosum		98.50	0.50	SUC 93	90		
Lactobacillus fermentum		1.50	0.24		1		
05413030	**Good identification**						**05413030**
Propionibacterium granulosum		100.0	0.45	SUC 93			
05413032	**Good identification**						**05413032**
Propionibacterium granulosum		100.0	0.35	SUC 93			
05413040	**Good identification**						**05413040**
					CAT		
Propionibacterium granulosum		97.59	0.36	SUC 93	90		
Eubacterium limosum		2.41	-.06	SOR 1	1		
05413100	**Low discrimination**						**05413100**
					GRA	CAT	
Fusobacterium mortiferum		57.61	0.45	MAN 1	1	5	
Propionibacterium granulosum		27.93	0.36		99	90	
Lactobacillus fermentum		12.84	0.35		99	1	
05413102	**Low discrimination**						**05413102**
					GRA	CAT	
Lactobacillus fermentum		46.30	0.35	SUC 87	99	1	
Propionibacterium granulosum		25.17	0.26	MAN 1	99	90	
Fusobacterium mortiferum		23.08	0.29		1	5	
05413200	**Good identification**						**05413200**
					CAT		
Propionibacterium granulosum		99.20	0.45	SUC 93	90		
Eubacterium limosum		0.80	-.06	RHA 3	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05413400	**Low discrimination**						**05413400**
					CAT		
Propionibacterium granulosum		76.54	0.36	SUC 93	90		
Lactobacillus fermentum		21.57	0.32	ARA 1	1		
Eubacterium limosum		1.89	-.06		1		
05413500	**Good identification**						**05413500**
					GRA		
Lactobacillus fermentum		95.67	0.35	SUC 87	99		
Fusobacterium mortiferum		4.34	0.11	MAN 1	1		
05413502	**Good identification**						**05413502**
Lactobacillus fermentum		100.0	0.35	SUC 87			
05414100	**Very good identification**						**05414100**
					GRA		
Fusobacterium mortiferum		96.05	0.57	XYL 7	1		
Lactobacillus fermentum		2.53	0.31		99		
Fusobacterium varium		0.67	0.01		1		
05414102	**Low discrimination**						**05414102**
					GRA		
Fusobacterium mortiferum		80.83	0.41	XYL 7	1		
Lactobacillus fermentum		19.17	0.31	BGL 10	99		
05415000	**Low discrimination**						**05415000**
					COC	GRA	
Lactobacillus fermentum		62.55	0.62	SUC 87	1	99	
Fusobacterium mortiferum		25.14	0.54		1	1	
Gemella morbillorum		5.45	0.32		99	99	
05415002	**Good identification**						**05415002**
					GRA	CAT	
Lactobacillus fermentum		93.69	0.62	SUC 87	99	1	
Fusobacterium mortiferum		4.18	0.37		1	5	
Propionibacterium granulosum		1.89	0.28		99	90	
05415020	**Acceptable identification**						**05415020**
					COC	CAT	
Lactobacillus fermentum		90.08	0.54	SUC 87	1	1	
Propionibacterium granulosum		5.47	0.28		1	90	
Gemella morbillorum		2.62	0.16		99	1	
05415022	**Good identification**						**05415022**
					GRA	CAT	
Lactobacillus fermentum		97.04	0.54	SUC 87	99	1	
Parabacteroides distasonis		1.49	0.20		1	76	
Propionibacterium granulosum		1.47	0.18		99	90	
05415100	**Low discrimination**						**05415100**
					GRA		
Lactobacillus fermentum		63.38	0.66	SUC 87	99		
Fusobacterium mortiferum		36.43	0.60		1		
Actinomyces meyeri		0.19	0.12		99		
05415102	**Good identification**						**05415102**
					GRA		
Lactobacillus fermentum		93.72	0.66	SUC 87	99		
Fusobacterium mortiferum		5.99	0.44		1		
Parabacteroides distasonis		0.29	0.20		1		
05415120	**Very good identification**						**05415120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Lactobacillus fermentum		96.23	0.57	SUC 87	99		
Fusobacterium mortiferum		1.68	0.26		1		
Parabacteroides distasonis		0.91	0.20		1		
05415122	**Low discrimination**						**05415122**
					GRA	CAT	
Lactobacillus fermentum		51.36	0.57	SUC 87	99	1	
Parabacteroides distasonis		47.80	0.54		1	76	
Bacteroides fragilis		0.45	-.03		1	99	
05415132	**Good identification**						**05415132**
					GRA		
Parabacteroides distasonis		98.02	0.49		1		
Lactobacillus fermentum		1.98	0.23		99		
05415322	**Very good identification**						**05415322**
					GRA		
Parabacteroides distasonis		99.52	0.62		1		
Lactobacillus fermentum		0.36	0.23		99		
Prevotella oralis		0.12	-.01		1		
05415332	**Very good identification**						**05415332**
Parabacteroides distasonis		100.0	0.58				
05415400	**Very good identification**						**05415400**
					COC	GRA	
Lactobacillus fermentum		98.64	0.62	SUC 87	1	99	
Gemella morbillorum		0.96	0.16		99	99	
Fusobacterium mortiferum		0.40	0.20		1	1	
05415402	**Very good identification**						**05415402**
Lactobacillus fermentum		100.0	0.62	SUC 87			
05415420	**Good identification**						**05415420**
Lactobacillus fermentum		100.0	0.54	SUC 87			
05415422	**Good identification**						**05415422**
					GRA		
Lactobacillus fermentum		99.07	0.54	SUC 87	99		
Parabacteroides distasonis		0.93	0.16		1		
05415500	**Very good identification**						**05415500**
					GRA		
Lactobacillus fermentum		99.03	0.66	SUC 87	99		
Fusobacterium mortiferum		0.58	0.26		1		
Bifidobacterium longum subsp. longum		0.39	0.17		99		
05415502	**Very good identification**						**05415502**
					GRA		
Lactobacillus fermentum		99.41	0.66	SUC 87	99		
Bifidobacterium longum subsp. longum		0.40	0.17		99		
Parabacteroides distasonis		0.19	0.16		1		
05415520	**Low discrimination**						**05415520**
					GRA		
Lactobacillus fermentum		86.09	0.57	SUC 87	99		
Bacteroides vulgatus		9.32	0.20		1		
Bifidobacterium longum subsp. longum		4.10	0.28		99		
05415522	**Low discrimination**						**05415522**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Lactobacillus fermentum		61.81	0.57	SUC 87	99			
Parabacteroides distasonis		35.25	0.50		1			
Bifidobacterium longum subsp. longum		2.94	0.28		99			
05415532	**Low discrimination**							**05415532**
					GRA			
Parabacteroides distasonis		77.10	0.46		1			
Bifidobacterium longum subsp. longum		20.36	0.32		99			
Lactobacillus fermentum		2.54	0.23		99			
05415720	**Good identification**							**05415720**
					GRA			
Bacteroides vulgatus		97.27	0.36	SUC 96	1			
Parabacteroides distasonis		1.73	0.24	MLT 99	1			
Lactobacillus fermentum		1.01	0.23		99			
05415722	**Very good identification**							**05415722**
					GRA			
Parabacteroides distasonis		98.85	0.59		1			
Lactobacillus fermentum		0.58	0.23		99			
Bacteroides vulgatus		0.57	0.02		1			
05415732	**Good identification**							**05415732**
Parabacteroides distasonis		100.0	0.54					
05416010	**Good identification**							**05416010**
					SPO	CAT		
Clostridium difficile		96.90	0.36	RAF 1	99	1		
Propionibacterium granulosum		2.63	0.22	XYL 5	1	90		
Eubacterium limosum		0.48	-.06		1	1		
05417000	**Low discrimination**							**05417000**
					GRA	CAT		
Propionibacterium granulosum		79.57	0.36	SUC 93	99	90		
Lactobacillus fermentum		13.17	0.28	XYL 1	99	1		
Fusobacterium mortiferum		5.29	0.20		1	5		
05420000	**Low discrimination**							**05420000**
					SPO	CAT		
Propionibacterium granulosum		55.59	0.47	SUC 93	1	90		
Clostridium botulinum biovar A		28.16	0.32	FRU 99	99	1		
Atopobium minutum		2.82	0.22		1	1		
05420002	**Low discrimination**							**05420002**
					COC	CAT		
Propionibacterium granulosum		82.93	0.36	SUC 93	1	90		
Atopobium minutum		11.21	0.19	FRU 99	1	1		
Anaerococcus prevotii		3.44	-.03		99	40		
05420020	**Low discrimination**							**05420020**
					SPO	CAT		
Propionibacterium granulosum		77.17	0.36	SUC 93	1	90		
Clostridium cadaveris		15.92	0.12	FRU 99	99	1		
Atopobium minutum		5.22	0.14		1	1		
05421000	**Acceptable identification**							**05421000**
					COC	GRA	CAT	
Propionibacterium granulosum		90.09	0.47	SUC 93	1	99	90	
Fusobacterium mortiferum		5.98	0.30	FRU 99	1	1	5	
Gemella morbillorum		2.41	0.13		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05421002	**Acceptable identification**							**05421002**
					GRA	CAT		
Propionibacterium granulosum		93.06	0.36	SUC 93	99	90		
Atopobium minutum		4.19	0.11	FRU 99	99	1		
Fusobacterium mortiferum		2.75	0.13		1	5		
05421020	**Good identification**							**05421020**
					CAT			
Propionibacterium granulosum		97.80	0.36	SUC 93	90			
Atopobium minutum		2.20	0.06	FRU 99	1			
05421100	**Low discrimination**							**05421100**
					COC	GRA	CAT	
Fusobacterium mortiferum		89.03	0.36	FRU 99	1	1	5	
Propionibacterium granulosum		5.81	0.13		1	99	90	
Blautia hansenii		5.17	-.03		99	99	1	
05422000	**Good identification**							**05422000**
					CAT			
Propionibacterium granulosum		99.18	0.45	SUC 93	90			
Eubacterium limosum		0.82	-.06	FRU 99	1			
05423000	**Good identification**							**05423000**
Propionibacterium granulosum		100.0	0.45	SUC 93				
05425322	**Good identification**							**05425322**
Parabacteroides distasonis		100.0	0.39	FRU 99				
05425722	**Good identification**							**05425722**
Parabacteroides distasonis		100.0	0.36	FRU 99				
05430000	**Good identification**							**05430000**
					GRA	CAT		
Propionibacterium granulosum		94.99	0.81	SUC 93	99	90		
Fusobacterium mortiferum		4.20	0.61		1	5		
Fusobacterium varium		0.42	0.24		1	1		
05430001	**Good identification**							**05430001**
					GRA	CAT		
Propionibacterium granulosum		95.76	0.47	SUC 93	99	90		
Fusobacterium mortiferum		4.24	0.27	URE 1	1	5		
05430002	**Very good identification**							**05430002**
					GRA	CAT		
Propionibacterium granulosum		97.95	0.71	SUC 93	99	90		
Fusobacterium mortiferum		1.93	0.45		1	5		
Atopobium minutum		0.12	0.19		99	1		
05430003	**Good identification**							**05430003**
Propionibacterium granulosum		100.0	0.36	SUC 93				
05430010	**Very good identification**							**05430010**
					GRA	SPO	CAT	
Propionibacterium granulosum		98.22	0.66	SUC 93	99	1	90	
Clostridium difficile		1.46	0.22	MLZ 12	99	99	1	
Fusobacterium mortiferum		0.32	0.27		1	1	5	
05430012	**Good identification**							**05430012**
Propionibacterium granulosum		100.0	0.56	SUC 93				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05430020	**Very good identification**							**05430020**
					GRA	SPO	CAT	
Propionibacterium granulosum		99.50	0.71	SUC 93	99	1	90	
Clostridium cadaveris		0.21	0.12		99	99	1	
Fusobacterium mortiferum		0.18	0.27		1	1	5	
05430021	**Good identification**							**05430021**
Propionibacterium granulosum		100.0	0.36	SUC 93				
05430022	**Very good identification**							**05430022**
					CAT			
Propionibacterium granulosum		99.83	0.60	SUC 93	90			
Atopobium minutum		0.17	0.11		1			
05430030	**Good identification**							**05430030**
Propionibacterium granulosum		100.0	0.56	SUC 93				
05430032	**Good identification**							**05430032**
Propionibacterium granulosum		100.0	0.45	SUC 93				
05430040	**Low discrimination**							**05430040**
					GRA	SPO	CAT	
Propionibacterium granulosum		88.04	0.47	SUC 93	99	1	90	
Clostridium botulinum biovar B		7.21	0.15	SOR 1	99	99	1	
Fusobacterium mortiferum		3.90	0.27		1	1	5	
05430042	**Good identification**							**05430042**
Propionibacterium granulosum		100.0	0.36	SUC 93				
05430060	**Low discrimination**							**05430060**
					SPO	CAT		
Propionibacterium granulosum		86.10	0.36	SUC 93	1	90		
Clostridium botulinum biovar B		13.90	0.10	SOR 1	99	1		
05430100	**Good identification**							**05430100**
					GRA	CAT		
Fusobacterium mortiferum		91.09	0.67		1	5		
Propionibacterium granulosum		8.91	0.47		99	90		
05430102	**Low discrimination**							**05430102**
					GRA	CAT		
Fusobacterium mortiferum		81.96	0.51	BGL 10	1	5		
Propionibacterium granulosum		18.04	0.36		99	90		
05430120	**Low discrimination**							**05430120**
					GRA	CAT		
Propionibacterium granulosum		69.41	0.36	SUC 93	99	90		
Fusobacterium mortiferum		28.68	0.33	LAC 1	1	5		
Clostridium perfringens		1.91	0.02		99	1		
05430200	**Good identification**							**05430200**
					GRA	CAT		
Propionibacterium granulosum		98.58	0.55	SUC 93	99	90		
Fusobacterium mortiferum		1.43	0.27	RHA 3	1	5		
05430202	**Good identification**							**05430202**
Propionibacterium granulosum		100.0	0.45	SUC 93				
05430210	**Good identification**							**05430210**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.40	SUC 93				
05430220	**Good identification**							**05430220**
Propionibacterium granulosum		100.0	0.45	SUC 93				
05430400	**Good identification**							**05430400**
					COC	GRA	CAT	
Propionibacterium granulosum		95.15	0.47	SUC 93	1	99	90	
Fusobacterium mortiferum		4.21	0.27	ARA 1	1	1	5	
Gemella morbillorum		0.64	0.03		99	99	1	
05430402	**Good identification**							**05430402**
Propionibacterium granulosum		100.0	0.36	SUC 93				
05430420	**Good identification**							**05430420**
Propionibacterium granulosum		100.0	0.36	SUC 93				
05431000	**Good identification**							**05431000**
					GRA	CAT		
Propionibacterium granulosum		93.59	0.81	SUC 93	99	90		
Fusobacterium mortiferum		6.21	0.64		1	5		
Prevotella intermedia		0.08	0.20		1	1		
05431001	**Acceptable identification**							**05431001**
					GRA	CAT		
Propionibacterium granulosum		93.77	0.47	SUC 93	99	90		
Fusobacterium mortiferum		6.23	0.30	URE 1	1	5		
05431002	**Very good identification**							**05431002**
					GRA	CAT		
Propionibacterium granulosum		96.83	0.71	SUC 93	99	90		
Fusobacterium mortiferum		2.86	0.48		1	5		
Lactobacillus fermentum		0.28	0.32		99	1		
05431003	**Good identification**							**05431003**
					GRA	CAT		
Propionibacterium granulosum		97.13	0.36	SUC 93	99	90		
Fusobacterium mortiferum		2.87	0.13	URE 1	1	5		
05431010	**Very good identification**							**05431010**
					GRA	CAT		
Propionibacterium granulosum		99.51	0.66	SUC 93	99	90		
Fusobacterium mortiferum		0.49	0.30	MLZ 12	1	5		
05431012	**Good identification**							**05431012**
					GRA	CAT		
Propionibacterium granulosum		99.78	0.56	SUC 93	99	90		
Fusobacterium mortiferum		0.22	0.13	MLZ 12	1	5		
05431020	**Very good identification**							**05431020**
					GRA	CAT		
Propionibacterium granulosum		99.22	0.71	SUC 93	99	90		
Clostridium perfringens		0.37	0.22		99	1		
Fusobacterium mortiferum		0.27	0.30		1	5		
05431021	**Good identification**							**05431021**
Propionibacterium granulosum		100.0	0.36	SUC 93				
05431022	**Very good identification**							**05431022**
					CAT			
Propionibacterium granulosum		98.64	0.60	SUC 93	90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.70	0.16		1		
Lactobacillus fermentum		0.38	0.24		1		
05431030	**Good identification**						**05431030**
Propionibacterium granulosum		100.0	0.56	SUC 93			
05431032	**Good identification**						**05431032**
					GRA		
Propionibacterium granulosum		99.32	0.45	SUC 93	99		
Parabacteroides distasonis		0.68	0.10	MLZ 12	1		
05431040	**Acceptable identification**						**05431040**
					GRA	CAT	
Propionibacterium granulosum		93.77	0.47	SUC 93	99	90	
Fusobacterium mortiferum		6.23	0.30	SOR 1	1	5	
05431042	**Good identification**						**05431042**
					GRA	CAT	
Propionibacterium granulosum		97.13	0.36	SUC 93	99	90	
Fusobacterium mortiferum		2.87	0.13	SOR 1	1	5	
05431060	**Acceptable identification**						**05431060**
					CAT		
Propionibacterium granulosum		93.39	0.36	SUC 93	90		
Clostridium perfringens		6.61	0.09	SOR 1	1		
05431100	**Good identification**						**05431100**
					GRA	CAT	
Fusobacterium mortiferum		93.05	0.70		1	5	
Propionibacterium granulosum		6.07	0.47		99	90	
Lactobacillus fermentum		0.70	0.35		99	1	
05431101	**Low discrimination**						**05431101**
					GRA	CAT	
Fusobacterium mortiferum		87.22	0.36	URE 1	1	5	
Actinomyces naeslundii		7.09	0.29		99	25	
Propionibacterium granulosum		5.69	0.13		99	90	
05431102	**Low discrimination**						**05431102**
					GRA	CAT	
Fusobacterium mortiferum		81.20	0.54	BGL 10	1	5	
Propionibacterium granulosum		11.91	0.36		99	90	
Lactobacillus fermentum		5.51	0.35		99	1	
05431110	**Low discrimination**						**05431110**
					GRA	CAT	
Fusobacterium mortiferum		53.19	0.36	MLZ 1	1	5	
Propionibacterium granulosum		46.81	0.32		99	90	
05431120	**Low discrimination**						**05431120**
					CAT	SPO	
Clostridium perfringens		50.61	0.36	SUC 91	1	84	
Propionibacterium granulosum		18.57	0.36	MLT 99	90	1	
Clostridium septicum		14.34	0.20		1	99	
05431122	**Low discrimination**						**05431122**
					GRA	CAT	
Parabacteroides distasonis		69.82	0.49	XYL 90	1	76	
Clostridium perfringens		21.21	0.31		99	1	
Propionibacterium granulosum		4.13	0.26		99	90	
05431132	**Good identification**						**05431132**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		98.52	0.44	XYL 90	1		
Propionibacterium granulosum		1.48	0.11		99		
05431140	**Acceptable identification**						**05431140**
					GRA	CAT	
Fusobacterium mortiferum		93.88	0.36	SOR 1	1	5	
Propionibacterium granulosum		6.12	0.13		99	90	
05431200	**Good identification**						**05431200**
					GRA	CAT	
Propionibacterium granulosum		97.88	0.55	SUC 93	99	90	
Fusobacterium mortiferum		2.12	0.30	RHA 3	1	5	
05431202	**Good identification**						**05431202**
					GRA	CAT	
Propionibacterium granulosum		99.05	0.45	SUC 93	99	90	
Fusobacterium mortiferum		0.95	0.13	RHA 3	1	5	
05431210	**Good identification**						**05431210**
Propionibacterium granulosum		100.0	0.40	SUC 93			
05431220	**Good identification**						**05431220**
Propionibacterium granulosum		100.0	0.45	SUC 93			
05431300	**Low discrimination**						**05431300**
					GRA	CAT	
Fusobacterium mortiferum		83.36	0.36	RHA 1	1	5	
Propionibacterium granulosum		16.64	0.21		99	90	
05431322	**Very good identification**						**05431322**
Parabacteroides distasonis		100.0	0.57	XYL 90			
05431332	**Good identification**						**05431332**
Parabacteroides distasonis		100.0	0.53	XYL 90			
05431400	**Low discrimination**						**05431400**
					GRA	CAT	
Propionibacterium granulosum		87.93	0.47	SUC 93	99	90	
Lactobacillus fermentum		6.23	0.32	ARA 1	99	1	
Fusobacterium mortiferum		5.84	0.30		1	5	
05431402	**Low discrimination**						**05431402**
					GRA	CAT	
Propionibacterium granulosum		76.16	0.36	SUC 93	99	90	
Lactobacillus fermentum		21.59	0.32	ARA 1	99	1	
Fusobacterium mortiferum		2.25	0.13		1	5	
05431420	**Acceptable identification**						**05431420**
					CAT		
Propionibacterium granulosum		91.37	0.36	SUC 93	90		
Lactobacillus fermentum		8.63	0.24	ARA 1	1		
05431500	**Low discrimination**						**05431500**
					GRA	CAT	
Fusobacterium mortiferum		55.20	0.36	ARA 1	1	5	
Lactobacillus fermentum		41.20	0.35		99	1	
Propionibacterium granulosum		3.60	0.13		99	90	
05431502	**Low discrimination**						**05431502**
					GRA		
Lactobacillus fermentum		82.96	0.35	SUC 87	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		12.35	0.20	TRE 1	1		
Parabacteroides distasonis		4.69	0.11		1		
05431522	**Acceptable identification**						**05431522**
					GRA		
Parabacteroides distasonis		93.17	0.45	XYL 90	1		
Lactobacillus fermentum		5.55	0.27		99		
Bifidobacterium longum subsp. longum		1.28	0.09		99		
05431532	**Good identification**						**05431532**
					GRA		
Parabacteroides distasonis		95.83	0.41	XYL 90	1		
Bifidobacterium longum subsp. longum		4.17	0.13		99		
05431722	**Good identification**						**05431722**
Parabacteroides distasonis		100.0	0.53	XYL 90			
05431732	**Good identification**						**05431732**
Parabacteroides distasonis		100.0	0.49	XYL 90			
05432000	**Very good identification**						**05432000**
					SPO	CAT	
Propionibacterium granulosum		98.63	0.79	SUC 93	1	90	
Eubacterium limosum		0.81	0.29		1	1	
Clostridium difficile		0.49	0.27		99	1	
05432001	**Good identification**						**05432001**
					CAT		
Propionibacterium granulosum		99.18	0.45	SUC 93	90		
Eubacterium limosum		0.82	-.06	URE 1	1		
05432002	**Very good identification**						**05432002**
					CAT		
Propionibacterium granulosum		99.97	0.68	SUC 93	90		
Eubacterium limosum		0.03	-.06		1		
05432010	**Low discrimination**						**05432010**
					SPO	CAT	
Propionibacterium granulosum		84.88	0.64	SUC 93	1	90	
Clostridium difficile		15.07	0.38	MLZ 12	99	1	
Eubacterium limosum		0.05	-.06		1	1	
05432012	**Good identification**						**05432012**
					SPO	CAT	
Propionibacterium granulosum		99.29	0.54	SUC 93	1	90	
Clostridium difficile		0.71	0.04	MLZ 12	99	1	
05432020	**Very good identification**						**05432020**
					CAT		
Propionibacterium granulosum		99.97	0.68	SUC 93	90		
Eubacterium limosum		0.03	-.06		1		
05432022	**Very good identification**						**05432022**
Propionibacterium granulosum		100.0	0.58	SUC 93			
05432030	**Good identification**						**05432030**
					SPO	CAT	
Propionibacterium granulosum		99.29	0.54	SUC 93	1	90	
Clostridium difficile		0.71	0.04	MLZ 12	99	1	
05432032	**Good identification**						**05432032**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.43	SUC 93			
05432040	**Good identification**						**05432040**
					SPO	CAT	
Propionibacterium granulosum		96.70	0.45	SUC 93	1	90	
Clostridium difficile		2.50	0.05	SOR 1	99	1	
Eubacterium limosum		0.80	-.06		1	1	
05432100	**Low discrimination**						**05432100**
					GRA	CAT	
Propionibacterium granulosum		87.33	0.45	SUC 93	99	90	
Fusobacterium mortiferum		11.96	0.33	LAC 1	1	5	
Eubacterium limosum		0.72	-.06		99	1	
05432200	**Good identification**						**05432200**
					CAT		
Propionibacterium granulosum		99.73	0.53	SUC 93	90		
Eubacterium limosum		0.27	-.06	RHA 3	1		
05432202	**Good identification**						**05432202**
Propionibacterium granulosum		100.0	0.43	SUC 93			
05432210	**Acceptable identification**						**05432210**
					SPO	CAT	
Propionibacterium granulosum		94.52	0.38	SUC 93	1	90	
Clostridium difficile		5.48	0.04	RHA 3	99	1	
05432220	**Good identification**						**05432220**
Propionibacterium granulosum		100.0	0.43	SUC 93			
05432400	**Good identification**						**05432400**
					CAT		
Propionibacterium granulosum		99.18	0.45	SUC 93	90		
Eubacterium limosum		0.82	-.06	ARA 1	1		
05433000	**Very good identification**						**05433000**
					GRA	CAT	
Propionibacterium granulosum		99.90	0.79	SUC 93	99	90	
Fusobacterium mortiferum		0.09	0.30		1	5	
Actinomyces naeslundii		0.01	0.23		99	25	
05433001	**Good identification**						**05433001**
Propionibacterium granulosum		98.87	0.45	SUC 93			
Actinomyces naeslundii		1.13	0.26	URE 1			
05433002	**Very good identification**						**05433002**
					GRA	CAT	
Propionibacterium granulosum		99.93	0.68	SUC 93	99	90	
Fusobacterium mortiferum		0.04	0.13		1	5	
Actinomyces naeslundii		0.03	0.23		99	25	
05433010	**Very good identification**						**05433010**
					SPO	CAT	
Propionibacterium granulosum		99.82	0.64	SUC 93	1	90	
Clostridium difficile		0.18	0.04	MLZ 12	99	1	
05433012	**Good identification**						**05433012**
Propionibacterium granulosum		100.0	0.54	SUC 93			
05433020	**Very good identification**						**05433020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		99.97	0.68	SUC 93			
Actinomyces naeslundii		0.03	0.23				
05433022	**Very good identification**						**05433022**
Propionibacterium granulosum		99.88	0.58	SUC 93			
Actinomyces naeslundii		0.12	0.23				
05433030	**Good identification**						**05433030**
Propionibacterium granulosum		100.0	0.54	SUC 93			
05433032	**Good identification**						**05433032**
Propionibacterium granulosum		100.0	0.43	SUC 93			
05433040	**Good identification**						**05433040**
Propionibacterium granulosum		100.0	0.45	SUC 93			
05433100	**Low discrimination**						**05433100**
					GRA	CAT	
Propionibacterium granulosum		81.50	0.45	SUC 93	99	90	
Fusobacterium mortiferum		16.74	0.36	LAC 1	1	5	
Actinomyces naeslundii		1.76	0.31		99	25	
05433200	**Good identification**						**05433200**
Propionibacterium granulosum		100.0	0.53	SUC 93			
05433202	**Good identification**						**05433202**
Propionibacterium granulosum		100.0	0.43	SUC 93			
05433210	**Good identification**						**05433210**
Propionibacterium granulosum		100.0	0.38	SUC 93			
05433220	**Good identification**						**05433220**
Propionibacterium granulosum		100.0	0.43	SUC 93			
05433400	**Good identification**						**05433400**
Propionibacterium granulosum		100.0	0.45	SUC 93			
05434000	**Low discrimination**						**05434000**
					GRA	CAT	
Propionibacterium granulosum		73.82	0.47	SUC 93	99	90	
Fusobacterium mortiferum		24.35	0.42	XYL 1	1	5	
Fusobacterium varium		1.33	0.01		1	1	
05434002	**Low discrimination**						**05434002**
					GRA	CAT	
Propionibacterium granulosum		87.22	0.36	SUC 93	99	90	
Fusobacterium mortiferum		12.79	0.25	XYL 1	1	5	
05434020	**Good identification**						**05434020**
Propionibacterium granulosum		100.0	0.36	SUC 93			
05434100	**Good identification**						**05434100**
					GRA	CAT	
Fusobacterium mortiferum		98.71	0.48	XYL 7	1	5	
Propionibacterium granulosum		1.30	0.13		99	90	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
05434322	**Good identification**						**05434322**
Parabacteroides distasonis		100.0	0.39	GAL 99			
05434722	**Good identification**						**05434722**
Parabacteroides distasonis		100.0	0.36	GAL 99			
05435000	**Low discrimination**						**05435000**
					GRA	CAT	
Propionibacterium granulosum		65.09	0.47	SUC 93	99	90	
Fusobacterium mortiferum		32.20	0.45	XYL 1	1	5	
Lactobacillus fermentum		2.71	0.28		99	1	
05435002	**Low discrimination**						**05435002**
					GRA	CAT	
Propionibacterium granulosum		72.13	0.36	SUC 93	99	90	
Fusobacterium mortiferum		15.86	0.28	XYL 1	1	5	
Lactobacillus fermentum		12.01	0.28		99	1	
05435020	**Acceptable identification**						**05435020**
					CAT		
Propionibacterium granulosum		94.74	0.36	SUC 93	90		
Lactobacillus fermentum		5.26	0.20	XYL 1	1		
05435100	**Acceptable identification**						**05435100**
					GRA	CAT	
Fusobacterium mortiferum		93.67	0.51	XYL 7	1	5	
Lactobacillus fermentum		5.51	0.31		99	1	
Propionibacterium granulosum		0.82	0.13		99	90	
05435122	**Very good identification**						**05435122**
					GRA		
Parabacteroides distasonis		99.76	0.65		1		
Lactobacillus fermentum		0.24	0.23		99		
05435132	**Very good identification**						**05435132**
Parabacteroides distasonis		100.0	0.61				
05435222	**Good identification**						**05435222**
Parabacteroides distasonis		100.0	0.39	LAC 99			
05435302	**Good identification**						**05435302**
Parabacteroides distasonis		100.0	0.39	NAG 99			
05435320	**Good identification**						**05435320**
Parabacteroides distasonis		97.68	0.39	BGL 99			
Bacteroides vulgatus		2.32	-.07				
05435322	**Very good identification**						**05435322**
Parabacteroides distasonis		100.0	0.73				
05435323	**Good identification**						**05435323**
Parabacteroides distasonis		100.0	0.39	URE 1			
05435332	**Very good identification**						**05435332**
Parabacteroides distasonis		100.0	0.69				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
05435362	**Good identification**						**05435362**
Parabacteroides distasonis		100.0	0.39	SOR 1			
05435522	**Very good identification**						**05435522**
					GRA		
Parabacteroides distasonis		99.16	0.62		1		
Bifidobacterium longum subsp. longum		0.46	0.17		99		
Lactobacillus fermentum		0.39	0.23		99		
05435532	**Very good identification**						**05435532**
					GRA		
Parabacteroides distasonis		98.57	0.57		1		
Bifidobacterium longum subsp. longum		1.43	0.21		99		
05435622	**Good identification**						**05435622**
Parabacteroides distasonis		100.0	0.36	LAC 99			
05435702	**Good identification**						**05435702**
Parabacteroides distasonis		100.0	0.36	NAG 99			
05435720	**Low discrimination**						**05435720**
Parabacteroides distasonis		65.98	0.36	BGL 99			
Bacteroides vulgatus		34.02	0.12				
05435722	**Very good identification**						**05435722**
Parabacteroides distasonis		100.0	0.70				
05435723	**Good identification**						**05435723**
Parabacteroides distasonis		100.0	0.36	URE 1			
05435732	**Very good identification**						**05435732**
Parabacteroides distasonis		100.0	0.65				
05435762	**Good identification**						**05435762**
Parabacteroides distasonis		100.0	0.36	SOR 1			
05436000	**Good identification**						**05436000**
					SPO	CAT	
Propionibacterium granulosum		96.70	0.45	SUC 93	1	90	
Clostridium difficile		2.50	0.05	XYL 1	99	1	
Eubacterium limosum		0.80	-.06		1	1	
05437000	**Good identification**						**05437000**
Propionibacterium granulosum		100.0	0.45	SUC 93			
05437322	**Good identification**						**05437322**
Parabacteroides distasonis		100.0	0.39	MAN 1			
05437722	**Good identification**						**05437722**
Parabacteroides distasonis		100.0	0.36	MAN 1			
05441100	**Good identification**						**05441100**
					COC	GRA	
Fusobacterium mortiferum		95.06	0.37	FRU 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		4.94	-.03		99	99	
05450000	**Low discrimination**						**05450000**
					GRA	CAT	
Fusobacterium mortiferum		81.18	0.62		1	5	
Fusobacterium varium		7.18	0.24		1	1	
Propionibacterium granulosum		5.53	0.38		99	90	
05450002	**Low discrimination**						**05450002**
					GRA	CAT	
Fusobacterium mortiferum		82.59	0.45	BGL 10	1	5	
Propionibacterium granulosum		12.67	0.28		99	90	
Atopobium minutum		4.74	0.19		99	1	
05450100	**Very good identification**						**05450100**
Fusobacterium mortiferum		100.0	0.68				
05450102	**Good identification**						**05450102**
Fusobacterium mortiferum		100.0	0.52	BGL 10			
05451000	**Good identification**						**05451000**
					GRA	CAT	
Fusobacterium mortiferum		93.17	0.65		1	5	
Propionibacterium granulosum		4.23	0.38		99	90	
Prevotella intermedia		1.08	0.20		1	1	
05451002	**Low discrimination**						**05451002**
					GRA	CAT	
Fusobacterium mortiferum		83.20	0.48	BGL 10	1	5	
Propionibacterium granulosum		8.51	0.28		99	90	
Lactobacillus fermentum		7.24	0.32		99	1	
05451100	**Very good identification**						**05451100**
					GRA		
Fusobacterium mortiferum		99.30	0.71		1		
Lactobacillus fermentum		0.67	0.35		99		
Actinomyces naeslundii		0.03	0.23		99		
05451101	**Good identification**						**05451101**
					GRA		
Fusobacterium mortiferum		95.26	0.37	URE 1	1		
Actinomyces naeslundii		4.74	0.26		99		
05451102	**Acceptable identification**						**05451102**
					GRA		
Fusobacterium mortiferum		92.81	0.55	BGL 10	1		
Lactobacillus fermentum		5.64	0.35		99		
Prevotella oralis		1.27	0.09		1		
05451110	**Good identification**						**05451110**
Fusobacterium mortiferum		100.0	0.37	MLZ 1			
05451140	**Good identification**						**05451140**
Fusobacterium mortiferum		100.0	0.37	SOR 1			
05451300	**Good identification**						**05451300**
Fusobacterium mortiferum		100.0	0.37	RHA 1			
05451322	**Low discrimination**						**05451322**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		63.94	0.41	XYL 90				
Prevotella oralis		36.06	0.24					
05451332	**Good identification**							**05451332**
Parabacteroides distasonis		100.0	0.37	XYL 90				
05451500	**Low discrimination**							**05451500**
					GRA			
Fusobacterium mortiferum		59.92	0.37	ARA 1	1			
Lactobacillus fermentum		40.08	0.35		99			
05451502	**Low discrimination**							**05451502**
					GRA			
Lactobacillus fermentum		85.76	0.35	SUC 87	99			
Fusobacterium mortiferum		14.25	0.21	CEL 1	1			
05451722	**Good identification**							**05451722**
Parabacteroides distasonis		100.0	0.38	XYL 90				
05452010	**Good identification**							**05452010**
					SPO	CAT		
Clostridium difficile		96.90	0.36	RAF 1	99	1		
Propionibacterium granulosum		2.63	0.22	CEL 5	1	90		
Eubacterium limosum		0.48	-.06		1	1		
05453000	**Low discrimination**							**05453000**
					GRA	SPO	CAT	
Propionibacterium granulosum		49.01	0.36	SUC 93	99	1	90	
Clostridium innocuum		35.34	0.25	CEL 1	99	99	1	
Fusobacterium mortiferum		14.44	0.31		1	1	5	
05453100	**Acceptable identification**							**05453100**
					GRA			
Fusobacterium mortiferum		93.95	0.37	MAN 1	1			
Actinomyces naeslundii		6.05	0.28		99			
05454000	**Acceptable identification**							**05454000**
					COC	GRA		
Fusobacterium mortiferum		93.70	0.43	XYL 7	1	1		
Fusobacterium varium		4.59	0.01		1	1		
Gemella morbillorum		1.72	0.03		99	99		
05454100	**Good identification**							**05454100**
Fusobacterium mortiferum		100.0	0.49	XYL 7				
05455000	**Acceptable identification**							**05455000**
					GRA			
Fusobacterium mortiferum		92.99	0.46	XYL 7	1			
Lactobacillus fermentum		7.01	0.28		99			
05455100	**Acceptable identification**							**05455100**
					GRA			
Fusobacterium mortiferum		94.99	0.52	XYL 7	1			
Lactobacillus fermentum		5.01	0.31		99			
05455102	**Low discrimination**							**05455102**
					GRA			
Fusobacterium mortiferum		64.36	0.35	XYL 7	1			
Lactobacillus fermentum		30.54	0.31	BGL 10	99			
Parabacteroides distasonis		5.10	0.15		1			
05455122	**Acceptable identification**							**05455122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		92.44	0.49		1		
Prevotella oralis		5.01	0.15		1		
Lactobacillus fermentum		1.86	0.23		99		
05455132	**Good identification**						**05455132**
Parabacteroides distasonis		100.0	0.45				
05455322	**Very good identification**						**05455322**
Parabacteroides distasonis		99.23	0.58				
Prevotella oralis		0.77	0.09				
05455332	**Good identification**						**05455332**
Parabacteroides distasonis		100.0	0.53				
05455522	**Good identification**						**05455522**
					GRA		
Parabacteroides distasonis		96.82	0.46		1		
Lactobacillus fermentum		3.18	0.23		99		
05455532	**Good identification**						**05455532**
Parabacteroides distasonis		100.0	0.41				
05455722	**Good identification**						**05455722**
Parabacteroides distasonis		100.0	0.54				
05455732	**Good identification**						**05455732**
Parabacteroides distasonis		100.0	0.49				
05470000	**Low discrimination**						**05470000**
					GRA	CAT	
Fusobacterium mortiferum		59.36	0.53		1	5	
Propionibacterium granulosum		40.64	0.47		99	90	
05470002	**Low discrimination**						**05470002**
					GRA	CAT	
Propionibacterium granulosum		60.64	0.36	SUC 93	99	90	
Fusobacterium mortiferum		39.37	0.36	CEL 1	1	5	
05470020	**Acceptable identification**						**05470020**
					GRA	CAT	
Propionibacterium granulosum		94.43	0.36	SUC 93	99	90	
Fusobacterium mortiferum		5.57	0.19	CEL 1	1	5	
05470100	**Very good identification**						**05470100**
					GRA	CAT	
Fusobacterium mortiferum		99.71	0.59		1	5	
Propionibacterium granulosum		0.30	0.13		99	90	
05470102	**Good identification**						**05470102**
Fusobacterium mortiferum		100.0	0.43	BGL 10			
05471000	**Low discrimination**						**05471000**
					GRA	CAT	
Fusobacterium mortiferum		68.45	0.56		1	5	
Propionibacterium granulosum		31.24	0.47		99	90	
Lactobacillus jensenii		0.31	0.06		99	1	
05471002	**Low discrimination**						**05471002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Propionibacterium granulosum		49.98	0.36	SUC 93	99	90		
Fusobacterium mortiferum		48.68	0.39	CEL 1	1	5		
Lactobacillus jensenii		1.34	0.03		99	1		
05471020	**Low discrimination**							**05471020**
					GRA	SPO	CAT	
Propionibacterium granulosum		89.83	0.36	SUC 93	99	1	90	
Fusobacterium mortiferum		7.95	0.22	CEL 1	1	1	5	
Clostridium septicum		2.22	-.05		99	99	1	
05471100	**Very good identification**							**05471100**
					GRA	CAT		
Fusobacterium mortiferum		99.47	0.62		1	5		
Actinomyces naeslundii		0.22	0.28		99	25		
Propionibacterium granulosum		0.20	0.13		99	90		
05471102	**Good identification**							**05471102**
					GRA			
Fusobacterium mortiferum		96.14	0.46	BGL 10	1			
Actinomyces naeslundii		1.88	0.28		99			
Lactobacillus acidophilus		1.00	-.01		99			
05471122	**Low discrimination**							**05471122**
					GRA			
Parabacteroides distasonis		87.65	0.44	XYL 90	1			
Prevotella oralis		4.85	0.10		1			
Clostridium perfringens		2.06	0.07		99			
05471132	**Good identification**							**05471132**
Parabacteroides distasonis		100.0	0.40	XYL 90				
05471322	**Good identification**							**05471322**
Parabacteroides distasonis		99.22	0.53	XYL 90				
Prevotella oralis		0.78	0.04					
05471332	**Good identification**							**05471332**
Parabacteroides distasonis		100.0	0.48	XYL 90				
05471522	**Good identification**							**05471522**
Parabacteroides distasonis		100.0	0.41	XYL 90				
05471532	**Good identification**							**05471532**
Parabacteroides distasonis		100.0	0.36	XYL 90				
05471722	**Good identification**							**05471722**
Parabacteroides distasonis		100.0	0.49	XYL 90				
05471732	**Good identification**							**05471732**
Parabacteroides distasonis		100.0	0.44	XYL 90				
05472000	**Acceptable identification**							**05472000**
					GRA	SPO	CAT	
Propionibacterium granulosum		94.91	0.45	SUC 93	99	1	90	
Clostridium difficile		2.45	0.05	CEL 1	99	99	1	
Fusobacterium mortiferum		1.86	0.19		1	1	5	
05473000	**Low discrimination**							**05473000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Propionibacterium granulosum		88.98	0.45	SUC 93	99	1	90	
Clostridium innocuum		7.51	0.17	CEL 1	99	99	1	
Fusobacterium mortiferum		2.61	0.22		1	1	5	
05473101	**Good identification**							**05473101**
Actinomyces naeslundii		100.0	0.36	SUC 99				
05473103	**Good identification**							**05473103**
Actinomyces naeslundii		100.0	0.36	SUC 99				
05473121	**Good identification**							**05473121**
Actinomyces naeslundii		100.0	0.36	SUC 99				
05473123	**Good identification**							**05473123**
Actinomyces naeslundii		100.0	0.36	SUC 99				
05474100	**Good identification**							**05474100**
Fusobacterium mortiferum		100.0	0.40	XYL 7				
05475000	**Acceptable identification**							**05475000**
					GRA	CAT		
Fusobacterium mortiferum		94.23	0.37	XYL 7	1	5		
Propionibacterium granulosum		5.77	0.13		99	90		
05475100	**Good identification**							**05475100**
Fusobacterium mortiferum		100.0	0.43	XYL 7				
05475122	**Very good identification**							**05475122**
Parabacteroides distasonis		100.0	0.61					
05475132	**Good identification**							**05475132**
Parabacteroides distasonis		100.0	0.56					
05475322	**Very good identification**							**05475322**
Parabacteroides distasonis		100.0	0.69					
05475332	**Very good identification**							**05475332**
Parabacteroides distasonis		100.0	0.64					
05475522	**Very good identification**							**05475522**
Parabacteroides distasonis		100.0	0.57					
05475532	**Good identification**							**05475532**
Parabacteroides distasonis		100.0	0.52					
05475722	**Very good identification**							**05475722**
Parabacteroides distasonis		99.89	0.65					
Bacteroides thetaiotaomicron		0.11	-.02					
05475732	**Very good identification**							**05475732**
Parabacteroides distasonis		99.82	0.61					

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.18	0.08				
05500000	**Low discrimination**						**05500000**
					COC	GRA	
Prevotella disiens		71.38	0.66	RAF 1	1	1	
Gemella morbillorum		18.27	0.64		99	99	
Prevotella intermedia		5.52	0.55		1	1	
05500020	**Low discrimination**						**05500020**
					COC	GRA	
Gemella morbillorum		45.01	0.48	RAF 1	99	99	
Prevotella bivia		32.84	0.32	NAG 10	1	1	
Prevotella disiens		15.99	0.32		1	1	
05500120	**Very good identification**						**05500120**
					COC	GRA	
Prevotella bivia		99.89	0.66	RAF 1	1	1	
Prevotella melaninogenica		0.05	0.15		1	1	
Blautia hansenii		0.04	0.15		99	99	
05500400	**Low discrimination**						**05500400**
					COC	GRA	
Gemella morbillorum		71.18	0.48	RAF 1	99	99	
Prevotella disiens		25.28	0.32	ARA 10	1	1	
Prevotella intermedia		1.96	0.20		1	1	
05501000	**Low discrimination**						**05501000**
					COC	GRA	
Gemella morbillorum		57.92	0.60	RAF 1	99	99	
Prevotella intermedia		32.52	0.55		1	1	
Prevotella disiens		4.25	0.32		1	1	
05501002	**Low discrimination**						**05501002**
					COC		
Lactobacillus fermentum		34.50	0.40	SUC 87	1		
Gemella morbillorum		24.57	0.26	FRU 99	99		
Blautia hansenii		16.05	0.15		99		
05501020	**Low discrimination**						**05501020**
					COC	GRA	
Gemella morbillorum		65.95	0.43	RAF 1	99	99	
Prevotella bivia		15.77	0.19	NAG 10	1	1	
Prevotella melaninogenica		4.06	0.15		1	1	
05501100	**Very good identification**						**05501100**
					COC	GRA	
Blautia hansenii		98.69	0.66	MNS 1	99	99	
Prevotella bivia		0.45	0.19		1	1	
Lactobacillus fermentum		0.39	0.44		1	99	
05501102	**Good identification**						**05501102**
					COC	GRA	
Blautia hansenii		96.06	0.50	MNS 1	99	99	
Lactobacillus fermentum		3.40	0.44	BGL 10	1	99	
Prevotella melaninogenica		0.19	0.03		1	1	
05501120	**Low discrimination**						**05501120**
					COC	GRA	
Prevotella bivia		65.84	0.53	RAF 1	1	1	
Prevotella melaninogenica		16.96	0.50		1	1	
Blautia hansenii		16.34	0.50		99	99	
05501122	**Low discrimination**						**05501122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Prevotella melaninogenica		50.17	0.37	SUC 99	1	1	
Blautia hansenii		28.20	0.33	FRU 90	99	99	
Prevotella bivia		10.33	0.19		1	1	
05501400	**Low discrimination**						**05501400**
					COC		
Gemella morbillorum		78.15	0.43	RAF 1	99		
Lactobacillus fermentum		9.98	0.40	ARA 10	1		
Blautia hansenii		7.37	0.19		99		
05501402	**Low discrimination**						**05501402**
					COC		
Lactobacillus fermentum		86.12	0.40	SUC 87	1		
Blautia hansenii		7.07	0.03	FRU 99	99		
Gemella morbillorum		6.81	0.09		99		
05501500	**Good identification**						**05501500**
					COC		
Blautia hansenii		97.12	0.53	MNS 1	99		
Lactobacillus fermentum		2.17	0.44		1		
Bifidobacterium longum subsp. longum		0.61	0.27		1		
05501502	**Low discrimination**						**05501502**
					COC		
Blautia hansenii		79.53	0.37	MNS 1	99		
Lactobacillus fermentum		15.97	0.44	BGL 10	1		
Bifidobacterium longum subsp. longum		4.51	0.27		1		
05501520	**Low discrimination**						**05501520**
					COC	GRA	
Blautia hansenii		63.15	0.37	MNS 1	99	99	
Prevotella bivia		14.56	0.19	NAG 10	1	1	
Bifidobacterium longum subsp. longum		14.31	0.37		1	99	
05501522	**Low discrimination**						**05501522**
					COC		
Bifidobacterium longum subsp. longum		54.48	0.37	SUC 99	1		
Blautia hansenii		26.71	0.20	FRU 90	99		
Lactobacillus fermentum		16.09	0.35		1		
05501530	**Good identification**						**05501530**
					COC		
Bifidobacterium longum subsp. longum		97.45	0.41	SUC 99	1		
Blautia hansenii		2.55	0.03	FRU 90	99		
05501532	**Good identification**						**05501532**
Bifidobacterium longum subsp. longum		100.0	0.41	SUC 99			
05504000	**Low discrimination**						**05504000**
					COC	GRA	
Gemella morbillorum		71.18	0.48	RAF 1	99	99	
Prevotella disiens		25.28	0.32	XYL 10	1	1	
Prevotella intermedia		1.96	0.20		1	1	
05505000	**Low discrimination**						**05505000**
					COC		
Gemella morbillorum		81.50	0.43	RAF 1	99		
Blautia hansenii		7.68	0.19	XYL 10	99		
Lactobacillus fermentum		6.11	0.36		1		
05505002	**Low discrimination**						**05505002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Lactobacillus fermentum		78.47	0.36	SUC 87	1		
Blautia hansenii		10.96	0.03	FRU 99	99		
Gemella morbillorum		10.57	0.09		99		
05505100	**Good identification**						**05505100**
					COC		
Blautia hansenii		98.60	0.53	MNS 1	99		
Lactobacillus fermentum		1.29	0.40		1		
Gemella morbillorum		0.11	0.09		99		
05505102	**Low discrimination**						**05505102**
					COC		
Blautia hansenii		89.45	0.37	MNS 1	99		
Lactobacillus fermentum		10.55	0.40	BGL 10	1		
05505120	**Low discrimination**						**05505120**
					COC	GRA	
Blautia hansenii		74.25	0.37	MNS 1	99	99	
Prevotella bivia		17.12	0.19	NAG 10	1	1	
Prevotella melaninogenica		4.41	0.15		1	1	
05505322	**Good identification**						**05505322**
Parabacteroides distasonis		96.82	0.37	FRU 99			
Prevotella oralis		3.18	-.01				
05505402	**Good identification**						**05505402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05505500	**Low discrimination**						**05505500**
					COC		
Blautia hansenii		84.65	0.40	MNS 1	99		
Bifidobacterium longum subsp. longum		9.06	0.35		1		
Lactobacillus fermentum		6.29	0.40		1		
05505510	**Acceptable identification**						**05505510**
					COC		
Bifidobacterium longum subsp. longum		94.75	0.39	SUC 99	1		
Blautia hansenii		5.25	0.06	FRU 90	99		
05505512	**Good identification**						**05505512**
Bifidobacterium longum subsp. longum		100.0	0.39	SUC 99			
05505520	**Low discrimination**						**05505520**
					COC	GRA	
Bifidobacterium longum subsp. longum		65.62	0.46	SUC 99	1	99	
Blautia hansenii		17.03	0.24	FRU 90	99	99	
Bacteroides vulgatus		13.56	0.20		1	1	
05505522	**Low discrimination**						**05505522**
					GRA		
Bifidobacterium longum subsp. longum		88.84	0.46	SUC 99	99		
Lactobacillus fermentum		5.14	0.31	FRU 90	99		
Parabacteroides distasonis		3.46	0.26		1		
05505530	**Good identification**						**05505530**
Bifidobacterium longum subsp. longum		100.0	0.50	SUC 99			
05505532	**Good identification**						**05505532**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		98.78	0.50	SUC 99	99			
Parabacteroides distasonis		1.22	0.21	FRU 90	1			
05505720	**Good identification**							**05505720**
					GRA			
Bacteroides vulgatus		99.46	0.36	SUC 96	1			
Bifidobacterium longum subsp. longum		0.54	0.11	FRU 99	99			
05510000	**Low discrimination**							**05510000**
					COC	GRA		
Prevotella intermedia		83.83	0.89		1	1		
Prevotella disiens		10.94	0.66		1	1		
Gemella morbillorum		3.42	0.66		99	99		
05510001	**Low discrimination**							**05510001**
					COC	GRA		
Prevotella intermedia		83.50	0.55	URE 1	1	1		
Prevotella disiens		10.90	0.32		1	1		
Gemella morbillorum		3.41	0.32		99	99		
05510002	**Low discrimination**							**05510002**
					GRA	CAT		
Prevotella intermedia		58.85	0.55	BGL 1	1	1		
Propionibacterium granulosum		25.91	0.56		99	90		
Prevotella disiens		7.68	0.32		1	1		
05510010	**Low discrimination**							**05510010**
					GRA	CAT		
Prevotella intermedia		70.43	0.55	MLZ 1	1	1		
Propionibacterium granulosum		16.92	0.52		99	90		
Prevotella disiens		9.19	0.32		1	1		
05510012	**Low discrimination**							**05510012**
					GRA	CAT		
Propionibacterium granulosum		84.02	0.42	SUC 93	99	90		
Prevotella intermedia		14.13	0.20	MLZ 12	1	1		
Prevotella disiens		1.85	-.03		1	1		
05510020	**Low discrimination**							**05510020**
					COC	GRA	CAT	
Prevotella intermedia		48.11	0.55	NAG 1	1	1	1	
Gemella morbillorum		21.61	0.50		99	99	1	
Propionibacterium granulosum		21.18	0.56		1	99	90	
05510022	**Low discrimination**							**05510022**
					GRA	CAT		
Propionibacterium granulosum		78.89	0.46	SUC 93	99	90		
Prevotella intermedia		7.24	0.20		1	1		
Lactobacillus fermentum		6.03	0.32		99	1		
05510030	**Low discrimination**							**05510030**
					COC	GRA	CAT	
Propionibacterium granulosum		78.20	0.42	SUC 93	1	99	90	
Prevotella intermedia		13.15	0.20	MLZ 12	1	1	1	
Gemella morbillorum		5.91	0.15		99	99	1	
05510040	**Low discrimination**							**05510040**
					GRA	SPO		
Prevotella intermedia		65.09	0.55	SOR 1	1	1		
Clostridium botulinum biovar B		20.90	0.41		99	99		
Prevotella disiens		8.50	0.32		1	1		
05510050	**Acceptable identification**							**05510050**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Clostridium botulinum biovar B		91.47	0.36	SUC 90	99	99	1	
Prevotella intermedia		5.84	0.20	RAF 1	1	1	1	
Propionibacterium granulosum		1.40	0.18		99	1	90	
05510060	**Low discrimination**							**05510060**
					COC	GRA	SPO	
Clostridium botulinum biovar B		88.02	0.36	SUC 90	1	99	99	
Prevotella intermedia		5.62	0.20	RAF 1	1	1	1	
Gemella morbillorum		2.53	0.15		99	99	1	
05510100	**Low discrimination**							**05510100**
Prevotella intermedia		56.59	0.55	LAC 1				
Fusobacterium mortiferum		28.44	0.61					
Prevotella disiens		7.39	0.32					
05510102	**Low discrimination**							**05510102**
					GRA			
Fusobacterium mortiferum		32.71	0.44	MLT 11	1			
Actinomyces viscosus		32.62	0.32	BGL 10	99			
Lactobacillus fermentum		24.14	0.44		99			
05510120	**Acceptable identification**							**05510120**
					GRA			
Prevotella bivia		93.81	0.50	RAF 1	1			
Prevotella melaninogenica		3.49	0.32	FRU 10	1			
Clostridium perfringens		1.39	0.32		99			
05510200	**Low discrimination**							**05510200**
					GRA	CAT		
Prevotella intermedia		81.43	0.55	RHA 1	1	1		
Prevotella disiens		10.63	0.32		1	1		
Propionibacterium granulosum		4.44	0.41		99	90		
05510400	**Low discrimination**							**05510400**
					COC	GRA		
Prevotella intermedia		61.54	0.55	ARA 1	1	1		
Gemella morbillorum		27.64	0.50		99	99		
Prevotella disiens		8.03	0.32		1	1		
05510402	**Low discrimination**							**05510402**
					COC	GRA		
Lactobacillus fermentum		55.32	0.40	SUC 87	1	99		
Prevotella intermedia		22.12	0.20	GAL 99	1	1		
Gemella morbillorum		9.94	0.15		99	99		
05510500	**Low discrimination**							**05510500**
					GRA			
Lactobacillus fermentum		65.10	0.44	SUC 87	99			
Prevotella intermedia		15.96	0.20	GAL 99	1			
Fusobacterium mortiferum		8.02	0.26		1			
05510502	**Good identification**							**05510502**
Lactobacillus fermentum		97.50	0.44	SUC 87				
Bifidobacterium longum subsp. longum		2.50	0.09	GAL 99				
05510522	**Low discrimination**							**05510522**
Lactobacillus fermentum		76.47	0.35	SUC 87				
Bifidobacterium longum subsp. longum		23.53	0.20	GAL 99				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
05511000	**Good identification**						**05511000**
					COC	GRA	
Prevotella intermedia		93.47	0.89		1	1	
Lactobacillus fermentum		2.36	0.74		1	99	
Gemella morbillorum		2.06	0.61		99	99	
05511001	**Acceptable identification**						**05511001**
					COC	GRA	
Prevotella intermedia		93.05	0.55	URE 1	1	1	
Lactobacillus fermentum		2.35	0.40		1	99	
Gemella morbillorum		2.05	0.27		99	99	
05511002	**Low discrimination**						**05511002**
					GRA	CAT	
Lactobacillus fermentum		61.94	0.74	SUC 87	99	1	
Prevotella intermedia		24.77	0.55		1	1	
Propionibacterium granulosum		10.91	0.56		99	90	
05511003	**Low discrimination**						**05511003**
					GRA		
Lactobacillus fermentum		50.07	0.40	SUC 87	99		
Prevotella intermedia		20.02	0.20	URE 1	1		
Actinomyces viscosus		12.69	0.16		99		
05511010	**Low discrimination**						**05511010**
					GRA	CAT	
Prevotella intermedia		77.40	0.55	MLZ 1	1	1	
Propionibacterium granulosum		18.59	0.52		99	90	
Lactobacillus fermentum		1.96	0.40		99	1	
05511012	**Low discrimination**						**05511012**
					GRA	CAT	
Propionibacterium granulosum		62.94	0.42	SUC 93	99	90	
Lactobacillus fermentum		26.48	0.40	MLZ 12	99	1	
Prevotella intermedia		10.59	0.20		1	1	
05511020	**Low discrimination**						**05511020**
					GRA		
Prevotella intermedia		30.92	0.55	NAG 1	1		
Lactobacillus fermentum		25.77	0.66		99		
Clostridium perfringens		18.19	0.51		99		
05511022	**Low discrimination**						**05511022**
					CAT		
Lactobacillus fermentum		61.20	0.66	SUC 87	1		
Clostridium perfringens		20.33	0.46		1		
Propionibacterium granulosum		8.08	0.46		90		
05511030	**Low discrimination**						**05511030**
					GRA	CAT	
Propionibacterium granulosum		68.19	0.42	SUC 93	99	90	
Prevotella intermedia		11.47	0.20	MLZ 12	1	1	
Lactobacillus fermentum		9.56	0.32		99	1	
05511040	**Acceptable identification**						**05511040**
					COC	GRA	
Prevotella intermedia		92.91	0.55	SOR 1	1	1	
Lactobacillus fermentum		2.35	0.40		1	99	
Gemella morbillorum		2.04	0.27		99	99	
05511042	**Low discrimination**						**05511042**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		60.96	0.40	SUC 87	99	1	
Prevotella intermedia		24.38	0.20	SOR 1	1	1	
Propionibacterium granulosum		10.73	0.22		99	90	
05511060	**Low discrimination**						**05511060**
					GRA		
Clostridium perfringens		79.95	0.39	SUC 91	99		
Prevotella intermedia		7.20	0.20	LAC 88	1		
Lactobacillus fermentum		6.01	0.32		99		
05511100	**Low discrimination**						**05511100**
					GRA		
Lactobacillus fermentum		65.33	0.78	SUC 87	99		
Prevotella intermedia		16.01	0.55		1		
Fusobacterium mortiferum		12.07	0.64		1		
05511101	**Low discrimination**						**05511101**
					SPO		
Lactobacillus fermentum		43.00	0.44	SUC 87	1		
Clostridium chauvoei		22.24	0.26	URE 1	99		
Actinomyces naeslundii		12.60	0.45		1		
05511102	**Good identification**						**05511102**
					GRA		
Lactobacillus fermentum		92.50	0.78	SUC 87	99		
Prevotella oralis		1.95	0.33		1		
Fusobacterium mortiferum		1.90	0.47		1		
05511103	**Low discrimination**						**05511103**
Lactobacillus fermentum		58.23	0.44	SUC 87			
Actinomyces naeslundii		17.07	0.45	URE 1			
Actinomyces viscosus		16.08	0.20				
05511110	**Low discrimination**						**05511110**
					GRA		
Lactobacillus fermentum		61.81	0.44	SUC 87	99		
Prevotella intermedia		15.15	0.20	MLZ 1	1		
Fusobacterium mortiferum		11.42	0.30		1		
05511112	**Low discrimination**						**05511112**
					GRA		
Lactobacillus fermentum		89.29	0.44	SUC 87	99		
Bifidobacterium longum subsp. longum		3.90	0.13	MLZ 1	99		
Prevotella oralis		1.88	-.01		1		
05511120	**Low discrimination**						**05511120**
					GRA		
Prevotella melaninogenica		62.70	0.66	SUC 99	1		
Clostridium perfringens		24.92	0.66		99		
Lactobacillus fermentum		7.85	0.70		99		
05511122	**Low discrimination**						**05511122**
					GRA		
Prevotella melaninogenica		28.49	0.54	SUC 99	1		
Clostridium perfringens		27.97	0.60		99		
Prevotella oralis		22.50	0.55		1		
05511132	**Low discrimination**						**05511132**
					GRA		
Parabacteroides distasonis		38.56	0.42	XYL 90	1		
Prevotella melaninogenica		15.33	0.19		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		15.05	0.26		99		
05511140	**Low discrimination**						**05511140**
					GRA		
Lactobacillus fermentum		46.86	0.44	SUC 87	99		
Clostridium perfringens		28.91	0.28	SOR 1	99		
Prevotella intermedia		11.49	0.20		1		
05511142	**Low discrimination**						**05511142**
Lactobacillus fermentum		72.39	0.44	SUC 87			
Clostridium perfringens		21.02	0.22	SOR 1			
Actinomyces naeslundii		2.50	0.29				
05511160	**Low discrimination**						**05511160**
					GRA		
Clostridium perfringens		86.15	0.54	SUC 91	99		
Prevotella melaninogenica		11.50	0.32		1		
Lactobacillus fermentum		1.44	0.35		99		
05511162	**Low discrimination**						**05511162**
					GRA		
Clostridium perfringens		86.88	0.48	SUC 91	99		
Prevotella melaninogenica		4.69	0.19		1		
Prevotella oralis		3.71	0.21		1		
05511200	**Acceptable identification**						**05511200**
					GRA	CAT	
Prevotella intermedia		90.39	0.55	RHA 1	1	1	
Propionibacterium granulosum		4.92	0.41		99	90	
Lactobacillus fermentum		2.28	0.40		99	1	
05511202	**Low discrimination**						**05511202**
					GRA	CAT	
Lactobacillus fermentum		49.21	0.40	SUC 87	99	1	
Propionibacterium granulosum		26.53	0.31	RHA 1	99	90	
Prevotella intermedia		19.68	0.20		1	1	
05511300	**Low discrimination**						**05511300**
					GRA		
Lactobacillus fermentum		64.00	0.44	SUC 87	99		
Prevotella intermedia		15.69	0.20	RHA 1	1		
Fusobacterium mortiferum		11.83	0.30		1		
05511302	**Low discrimination**						**05511302**
					GRA		
Lactobacillus fermentum		49.95	0.44	SUC 87	99		
Prevotella oralis		44.67	0.27	RHA 1	1		
Parabacteroides distasonis		3.59	0.21		1		
05511322	**Low discrimination**						**05511322**
Prevotella oralis		66.80	0.49	SUC 92			
Parabacteroides distasonis		27.94	0.55				
Prevotella melaninogenica		1.99	0.19				
05511332	**Good identification**						**05511332**
Parabacteroides distasonis		95.71	0.51	XYL 90			
Prevotella oralis		4.29	0.15				
05511400	**Low discrimination**						**05511400**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		66.40	0.74	SUC 87	1	99	
Prevotella intermedia		26.55	0.55		1	1	
Gemella morbillorum		6.42	0.45		99	99	
05511401	**Low discrimination**						**05511401**
					COC	GRA	
Lactobacillus fermentum		66.82	0.40	SUC 87	1	99	
Prevotella intermedia		26.72	0.20	URE 1	1	1	
Gemella morbillorum		6.46	0.11		99	99	
05511402	**Very good identification**						**05511402**
					GRA	CAT	
Lactobacillus fermentum		99.28	0.74	SUC 87	99	1	
Prevotella intermedia		0.40	0.20		1	1	
Propionibacterium granulosum		0.18	0.22		99	90	
05511403	**Good identification**						**05511403**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05511410	**Low discrimination**						**05511410**
					COC	GRA	
Lactobacillus fermentum		60.10	0.40	SUC 87	1	99	
Prevotella intermedia		24.03	0.20	MLZ 1	1	1	
Gemella morbillorum		5.81	0.11		99	99	
05511412	**Acceptable identification**						**05511412**
Lactobacillus fermentum		93.35	0.40	SUC 87			
Bifidobacterium longum subsp. longum		6.65	0.13	MLZ 1			
05511420	**Good identification**						**05511420**
					COC	GRA	
Lactobacillus fermentum		94.05	0.66	SUC 87	1	99	
Gemella morbillorum		3.03	0.29		99	99	
Prevotella intermedia		1.14	0.20		1	1	
05511422	**Very good identification**						**05511422**
Lactobacillus fermentum		98.91	0.66	SUC 87			
Bifidobacterium longum subsp. longum		0.50	0.20				
Clostridium perfringens		0.33	0.11				
05511440	**Low discrimination**						**05511440**
					COC	GRA	
Lactobacillus fermentum		66.82	0.40	SUC 87	1	99	
Prevotella intermedia		26.72	0.20	SOR 1	1	1	
Gemella morbillorum		6.46	0.11		99	99	
05511442	**Good identification**						**05511442**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05511500	**Very good identification**						**05511500**
					COC		
Lactobacillus fermentum		96.55	0.78	SUC 87	1		
Bifidobacterium longum subsp. longum		2.48	0.43		1		
Blautia hansenii		0.44	0.19		99		
05511501	**Good identification**						**05511501**
Lactobacillus fermentum		97.50	0.44	SUC 87			
Bifidobacterium longum subsp. longum		2.50	0.09	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05511502	**Very good identification**					**05511502**
Lactobacillus fermentum		97.28	0.78	SUC 87		
Bifidobacterium longum subsp. longum		2.50	0.43			
Bifidobacterium adolescentis		0.10	0.21			
05511503	**Good identification**					**05511503**
Lactobacillus fermentum		97.50	0.44	SUC 87		
Bifidobacterium longum subsp. longum		2.50	0.09	URE 1		
05511510	**Low discrimination**					**05511510**
Bifidobacterium longum subsp. longum		81.21	0.47	SUC 99		
Lactobacillus fermentum		18.79	0.44			
05511512	**Low discrimination**					**05511512**
Bifidobacterium longum subsp. longum		79.29	0.47	SUC 99		
Lactobacillus fermentum		18.34	0.44			
Bifidobacterium adolescentis		2.37	0.23			
05511520	**Low discrimination**					**05511520**
					GRA	
Lactobacillus fermentum		67.99	0.70	SUC 87	99	
Bifidobacterium longum subsp. longum		20.92	0.54		99	
Prevotella melaninogenica		5.48	0.32		1	
05511521	**Low discrimination**					**05511521**
					GRA	
Lactobacillus fermentum		72.02	0.35	SUC 87	99	
Bifidobacterium longum subsp. longum		22.17	0.20	URE 1	99	
Prevotella melaninogenica		5.81	-.03		1	
05511522	**Low discrimination**					**05511522**
					GRA	
Lactobacillus fermentum		71.43	0.70	SUC 87	99	
Bifidobacterium longum subsp. longum		21.98	0.54		99	
Parabacteroides distasonis		3.14	0.43		1	
05511523	**Low discrimination**					**05511523**
Lactobacillus fermentum		76.47	0.35	SUC 87		
Bifidobacterium longum subsp. longum		23.53	0.20	URE 1		
05511530	**Very good identification**					**05511530**
					GRA	
Bifidobacterium longum subsp. longum		97.96	0.58	SUC 99	99	
Lactobacillus fermentum		1.89	0.35		99	
Prevotella melaninogenica		0.15	-.03		1	
05511532	**Good identification**					**05511532**
					GRA	
Bifidobacterium longum subsp. longum		93.84	0.58	SUC 99	99	
Parabacteroides distasonis		4.24	0.39		1	
Lactobacillus fermentum		1.81	0.35		99	
05511540	**Good identification**					**05511540**
Lactobacillus fermentum		97.50	0.44	SUC 87		
Bifidobacterium longum subsp. longum		2.50	0.09	SOR 1		
05511542	**Good identification**					**05511542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		97.50	0.44	SUC 87			
Bifidobacterium longum subsp. longum		2.50	0.09	SOR 1			
05511560	**Low discrimination**						**05511560**
Lactobacillus fermentum		50.18	0.35	SUC 87			
Clostridium perfringens		30.33	0.19	SOR 1			
Bifidobacterium longum subsp. longum		15.44	0.20				
05511562	**Low discrimination**						**05511562**
Lactobacillus fermentum		60.43	0.35	SUC 87			
Bifidobacterium longum subsp. longum		18.60	0.20	SOR 1			
Clostridium perfringens		17.19	0.14				
05511600	**Low discrimination**						**05511600**
					COC	GRA	
Lactobacillus fermentum		66.82	0.40	SUC 87	1	99	
Prevotella intermedia		26.72	0.20	RHA 1	1	1	
Gemella morbillorum		6.46	0.11		99	99	
05511602	**Good identification**						**05511602**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05511700	**Low discrimination**						**05511700**
					GRA		
Lactobacillus fermentum		86.65	0.44	SUC 87	99		
Bacteroides vulgatus		11.13	0.07	RHA 1	1		
Bifidobacterium longum subsp. longum		2.22	0.09		99		
05511702	**Acceptable identification**						**05511702**
					GRA		
Lactobacillus fermentum		93.49	0.44	SUC 87	99		
Parabacteroides distasonis		4.11	0.17	RHA 1	1		
Bifidobacterium longum subsp. longum		2.40	0.09		99		
05511720	**Good identification**						**05511720**
					GRA		
Bacteroides vulgatus		96.17	0.41	SUC 96	1		
Lactobacillus fermentum		2.52	0.35	XYL 98	99		
Bifidobacterium longum subsp. longum		0.78	0.20		99		
05511722	**Low discrimination**						**05511722**
					GRA		
Parabacteroides distasonis		85.54	0.52	XYL 90	1		
Lactobacillus fermentum		6.55	0.35		99		
Prevotella oralis		3.37	0.15		1		
05511732	**Acceptable identification**						**05511732**
					GRA		
Parabacteroides distasonis		93.07	0.47	XYL 90	1		
Bifidobacterium longum subsp. longum		6.93	0.24		99		
05512000	**Low discrimination**						**05512000**
					GRA	CAT	
Propionibacterium granulosum		51.02	0.65	SUC 93	99	90	
Prevotella intermedia		38.40	0.55		1	1	
Prevotella disiens		5.01	0.32		1	1	
05512002	**Good identification**						**05512002**
					COC	GRA	CAT

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		95.60	0.54	SUC 93	1	99	90	
Prevotella intermedia		2.91	0.20		1	1	1	
Megasphaera elsdenii		0.72	0.12		99	1	1	
05512010	**Low discrimination**							**05512010**
					GRA	SPO	CAT	
Propionibacterium granulosum		81.55	0.50	SUC 93	99	1	90	
Clostridium difficile		12.98	0.24	MLZ 12	99	99	1	
Prevotella intermedia		4.55	0.20		1	1	1	
05512012	**Good identification**							**05512012**
Propionibacterium granulosum		100.0	0.39	SUC 93				
05512020	**Good identification**							**05512020**
					COC	GRA	CAT	
Propionibacterium granulosum		95.22	0.54	SUC 93	1	99	90	
Prevotella intermedia		2.90	0.20		1	1	1	
Gemella morbillorum		1.30	0.15		99	99	1	
05512022	**Good identification**							**05512022**
Propionibacterium granulosum		100.0	0.44	SUC 93				
05512030	**Good identification**							**05512030**
Propionibacterium granulosum		100.0	0.39	SUC 93				
05512100	**Low discrimination**							**05512100**
					GRA	CAT		
Propionibacterium propionicum		56.63	0.43	NIT 86	99	1		
Propionibacterium granulosum		18.59	0.30	SUC 99	99	90		
Prevotella intermedia		13.99	0.20		1	1		
05512102	**Acceptable identification**							**05512102**
					CAT			
Propionibacterium propionicum		93.87	0.46	NIT 86	1			
Propionibacterium granulosum		5.14	0.20	SUC 99	90			
Actinomyces naeslundii		1.00	0.23		25			
05512140	**Good identification**							**05512140**
Propionibacterium propionicum		100.0	0.43	NIT 86				
05512142	**Good identification**							**05512142**
Propionibacterium propionicum		100.0	0.46	NIT 86				
05512200	**Low discrimination**							**05512200**
					GRA	CAT		
Propionibacterium granulosum		77.17	0.39	SUC 93	99	90		
Prevotella intermedia		18.97	0.20	RHA 3	1	1		
Prevotella disiens		2.48	-.03		1	1		
05513000	**Low discrimination**							**05513000**
					GRA	CAT		
Propionibacterium granulosum		55.67	0.65	SUC 93	99	90		
Prevotella intermedia		41.90	0.55		1	1		
Lactobacillus fermentum		1.06	0.40		99	1		
05513002	**Low discrimination**							**05513002**
					GRA	CAT		
Propionibacterium granulosum		88.90	0.54	SUC 93	99	90		
Lactobacillus fermentum		6.76	0.40		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		2.70	0.20		1	1	
05513003	**Low discrimination**						**05513003**
Actinomyces naeslundii		71.06	0.42	SUC 99			
Propionibacterium granulosum		28.94	0.20				
05513010	**Acceptable identification**						**05513010**
					GRA	CAT	
Propionibacterium granulosum		94.72	0.50	SUC 93	99	90	
Prevotella intermedia		5.28	0.20	MLZ 12	1	1	
05513012	**Good identification**						**05513012**
Propionibacterium granulosum		100.0	0.39	SUC 93			
05513020	**Acceptable identification**						**05513020**
					GRA	CAT	
Propionibacterium granulosum		90.84	0.54	SUC 93	99	90	
Prevotella intermedia		2.76	0.20		1	1	
Lactobacillus fermentum		2.30	0.32		99	1	
05513021	**Low discrimination**						**05513021**
Actinomyces naeslundii		71.06	0.42	SUC 99			
Propionibacterium granulosum		28.94	0.20				
05513022	**Low discrimination**						**05513022**
					CAT		
Propionibacterium granulosum		82.38	0.44	SUC 93	90		
Lactobacillus fermentum		8.35	0.32		1		
Actinomyces naeslundii		5.45	0.39		25		
05513023	**Acceptable identification**						**05513023**
Actinomyces naeslundii		90.76	0.42	SUC 99			
Propionibacterium granulosum		9.24	0.10				
05513030	**Good identification**						**05513030**
Propionibacterium granulosum		100.0	0.39	SUC 93			
05513100	**Low discrimination**						**05513100**
Lactobacillus fermentum		31.13	0.44	SUC 87			
Propionibacterium propionicum		30.88	0.43	MAN 1			
Actinomyces naeslundii		11.81	0.46				
05513101	**Low discrimination**						**05513101**
Actinomyces naeslundii		83.77	0.49	SUC 99			
Propionibacterium propionicum		16.23	0.27				
05513102	**Low discrimination**						**05513102**
Propionibacterium propionicum		49.50	0.46	NIT 86			
Lactobacillus fermentum		33.27	0.44	SUC 99			
Actinomyces naeslundii		12.62	0.46				
05513103	**Low discrimination**						**05513103**
Actinomyces naeslundii		77.49	0.49	SUC 99			
Propionibacterium propionicum		22.51	0.30				
05513121	**Good identification**						**05513121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		95.49	0.49	SUC 99	99		
Prevotella melaninogenica		4.51	-.03		1		
05513123	**Good identification**						**05513123**
Actinomyces naeslundii		100.0	0.49	SUC 99			
05513140	**Acceptable identification**						**05513140**
Propionibacterium propionicum		93.68	0.43	NIT 86			
Actinomyces naeslundii		6.32	0.34	SUC 99			
05513142	**Good identification**						**05513142**
Propionibacterium propionicum		95.69	0.46	NIT 86			
Actinomyces naeslundii		4.31	0.34	SUC 99			
05513161	**Good identification**						**05513161**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05513163	**Good identification**						**05513163**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05513200	**Low discrimination**						**05513200**
					GRA	CAT	
Propionibacterium granulosum		80.27	0.39	SUC 93	99	90	
Prevotella intermedia		19.73	0.20	RHA 3	1	1	
05513400	**Low discrimination**						**05513400**
					GRA	CAT	
Lactobacillus fermentum		49.31	0.40	SUC 87	99	1	
Propionibacterium granulosum		26.20	0.30	MAN 1	99	90	
Prevotella intermedia		19.72	0.20		1	1	
05513402	**Low discrimination**						**05513402**
					CAT		
Lactobacillus fermentum		88.28	0.40	SUC 87	1		
Propionibacterium granulosum		11.73	0.20	MAN 1	90		
05513500	**Good identification**						**05513500**
Lactobacillus fermentum		97.50	0.44	SUC 87			
Bifidobacterium longum subsp. longum		2.50	0.09	MAN 1			
05513502	**Low discrimination**						**05513502**
Lactobacillus fermentum		88.71	0.44	SUC 87			
Bifidobacterium adolescentis		7.69	0.20	MAN 1			
Bifidobacterium longum subsp. longum		2.28	0.09				
05513520	**Low discrimination**						**05513520**
					GRA		
Lactobacillus fermentum		72.02	0.35	SUC 87	99		
Bifidobacterium longum subsp. longum		22.17	0.20	MAN 1	99		
Prevotella melaninogenica		5.81	-.03		1		
05513522	**Low discrimination**						**05513522**
Lactobacillus fermentum		76.47	0.35	SUC 87			
Bifidobacterium longum subsp. longum		23.53	0.20	MAN 1			
05514000	**Low discrimination**						**05514000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Prevotella intermedia		61.19	0.55	XYL 1	1	1	
Gemella morbillorum		27.48	0.50		99	99	
Prevotella disiens		7.99	0.32		1	1	
05514002	**Low discrimination**						**05514002**
					COC	GRA	
Lactobacillus fermentum		40.05	0.36	SUC 87	1	99	
Prevotella intermedia		27.27	0.20	GAL 99	1	1	
Gemella morbillorum		12.25	0.15		99	99	
05514100	**Low discrimination**						**05514100**
					GRA		
Fusobacterium mortiferum		47.42	0.41	MLT 11	1		
Lactobacillus fermentum		30.34	0.40	XYL 7	99		
Prevotella intermedia		12.66	0.20		1		
05514102	**Low discrimination**						**05514102**
					GRA		
Lactobacillus fermentum		85.20	0.40	SUC 87	99		
Fusobacterium mortiferum		14.80	0.25	GAL 99	1		
05514322	**Good identification**						**05514322**
Parabacteroides distasonis		96.82	0.37	GAL 99			
Prevotella oralis		3.18	-.01				
05514402	**Good identification**						**05514402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05514500	**Low discrimination**						**05514500**
Lactobacillus fermentum		88.42	0.40	SUC 87			
Bifidobacterium longum subsp. longum		11.58	0.17	GAL 99			
05514502	**Low discrimination**						**05514502**
Lactobacillus fermentum		88.42	0.40	SUC 87			
Bifidobacterium longum subsp. longum		11.58	0.17	GAL 99			
05514720	**Good identification**						**05514720**
Bacteroides vulgatus		100.0	0.36	SUC 96			
05515000	**Low discrimination**						**05515000**
					COC	GRA	
Lactobacillus fermentum		53.32	0.70	SUC 87	1	99	
Prevotella intermedia		36.30	0.55		1	1	
Gemella morbillorum		8.78	0.45		99	99	
05515001	**Low discrimination**						**05515001**
					COC	GRA	
Lactobacillus fermentum		54.19	0.36	SUC 87	1	99	
Prevotella intermedia		36.89	0.20	URE 1	1	1	
Gemella morbillorum		8.92	0.11		99	99	
05515002	**Very good identification**						**05515002**
					GRA	CAT	
Lactobacillus fermentum		98.63	0.70	SUC 87	99	1	
Prevotella intermedia		0.68	0.20		1	1	
Propionibacterium granulosum		0.30	0.22		99	90	
05515003	**Good identification**						**05515003**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		100.0	0.36	SUC 87			
05515010	**Low discrimination**						**05515010**
					COC	GRA	
Lactobacillus fermentum		49.77	0.36	SUC 87	1	99	
Prevotella intermedia		33.89	0.20	MLZ 1	1	1	
Gemella morbillorum		8.20	0.11		99	99	
05515012	**Good identification**						**05515012**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05515020	**Good identification**						**05515020**
					COC	GRA	
Lactobacillus fermentum		90.51	0.62	SUC 87	1	99	
Gemella morbillorum		4.97	0.29		99	99	
Prevotella intermedia		1.87	0.20		1	1	
05515022	**Good identification**						**05515022**
					GRA		
Lactobacillus fermentum		94.59	0.62	SUC 87	99		
Prevotella oralis		2.49	0.19		1		
Parabacteroides distasonis		1.72	0.29		1		
05515040	**Low discrimination**						**05515040**
					GRA	SPO	
Lactobacillus fermentum		42.28	0.36	SUC 87	99	1	
Prevotella intermedia		28.79	0.20	SOR 1	1	1	
Clostridium glycolicum		21.97	0.15		99	99	
05515042	**Good identification**						**05515042**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05515100	**Very good identification**						**05515100**
					COC	GRA	
Lactobacillus fermentum		96.05	0.74	SUC 87	1	99	
Fusobacterium mortiferum		2.28	0.44		1	1	
Blautia hansenii		0.75	0.19		99	99	
05515101	**Good identification**						**05515101**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05515102	**Very good identification**						**05515102**
					GRA		
Lactobacillus fermentum		98.38	0.74	SUC 87	99		
Prevotella oralis		0.44	0.18		1		
Parabacteroides distasonis		0.36	0.29		1		
05515103	**Good identification**						**05515103**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05515110	**Low discrimination**						**05515110**
Lactobacillus fermentum		81.76	0.40	SUC 87			
Bifidobacterium longum subsp. longum		18.24	0.21	MLZ 1			
05515112	**Low discrimination**						**05515112**
Lactobacillus fermentum		60.95	0.40	SUC 87			
Bifidobacterium longum subsp. longum		13.60	0.21	MLZ 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		13.43	0.23				
05515120	**Low discrimination**						**05515120**
					GRA		
Lactobacillus fermentum		65.43	0.66	SUC 87	99		
Bacteroides vulgatus		17.59	0.35		1		
Prevotella melaninogenica		8.99	0.32		1		
05515122	**Low discrimination**						**05515122**
					GRA	CAT	
Parabacteroides distasonis		40.38	0.63		1	76	
Lactobacillus fermentum		36.71	0.66		99	1	
Bacteroides fragilis		10.50	0.32		1	99	
05515132	**Good identification**						**05515132**
					GRA		
Parabacteroides distasonis		93.07	0.59		1		
Bifidobacterium longum subsp. longum		4.25	0.32		99		
Lactobacillus fermentum		1.59	0.31		99		
05515140	**Good identification**						**05515140**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05515142	**Good identification**						**05515142**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05515200	**Low discrimination**						**05515200**
					COC	GRA	
Lactobacillus fermentum		54.19	0.36	SUC 87	1	99	
Prevotella intermedia		36.89	0.20	RHA 1	1	1	
Gemella morbillorum		8.92	0.11		99	99	
05515202	**Good identification**						**05515202**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05515222	**Low discrimination**						**05515222**
					GRA		
Parabacteroides distasonis		71.77	0.37	LAC 99	1		
Prevotella oralis		14.89	0.13		1		
Lactobacillus fermentum		13.33	0.28		99		
05515300	**Low discrimination**						**05515300**
					GRA		
Lactobacillus fermentum		55.35	0.40	SUC 87	99		
Bacteroides vulgatus		44.65	0.17	RHA 1	1		
05515302	**Low discrimination**						**05515302**
					GRA		
Parabacteroides distasonis		48.07	0.37	NAG 99	1		
Lactobacillus fermentum		43.71	0.40		99		
Prevotella oralis		8.23	0.12		1		
05515320	**Good identification**						**05515320**
					GRA		
Bacteroides vulgatus		97.96	0.51	SUC 96	1		
Parabacteroides distasonis		1.35	0.37	ARA 93	1		
Lactobacillus fermentum		0.41	0.31		99		
05515322	**Very good identification**						**05515322**
Parabacteroides distasonis		95.77	0.72				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		3.15	0.33				
Bacteroides vulgatus		0.71	0.17				
05515323	**Good identification**						**05515323**
Parabacteroides distasonis		96.82	0.37	URE 1			
Prevotella oralis		3.18	-.01				
05515332	**Very good identification**						**05515332**
Parabacteroides distasonis		99.94	0.67				
Prevotella oralis		0.06	-.01				
05515362	**Good identification**						**05515362**
Parabacteroides distasonis		96.82	0.37	SOR 1			
Prevotella oralis		3.18	-.01				
05515400	**Very good identification**						**05515400**
					COC	GRA	
Lactobacillus fermentum		97.34	0.70	SUC 87	1	99	
Gemella morbillorum		1.78	0.29		99	99	
Prevotella intermedia		0.67	0.20		1	1	
05515401	**Good identification**						**05515401**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05515402	**Very good identification**						**05515402**
Lactobacillus fermentum		99.79	0.70	SUC 87			
Bifidobacterium longum subsp. longum		0.22	0.17				
05515403	**Good identification**						**05515403**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05515410	**Low discrimination**						**05515410**
Lactobacillus fermentum		73.32	0.36	SUC 87			
Bifidobacterium longum subsp. longum		26.68	0.21	MLZ 1			
05515412	**Low discrimination**						**05515412**
Lactobacillus fermentum		73.32	0.36	SUC 87			
Bifidobacterium longum subsp. longum		26.68	0.21	MLZ 1			
05515420	**Good identification**						**05515420**
					GRA		
Lactobacillus fermentum		91.67	0.62	SUC 87	99		
Bacteroides vulgatus		5.40	0.20		1		
Bifidobacterium longum subsp. longum		2.38	0.28		99		
05515422	**Very good identification**						**05515422**
					GRA		
Lactobacillus fermentum		96.43	0.62	SUC 87	99		
Bifidobacterium longum subsp. longum		2.50	0.28		99		
Parabacteroides distasonis		1.07	0.26		1		
05515440	**Good identification**						**05515440**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05515442	**Good identification**						**05515442**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		100.0	0.36	SUC 87		
05515500	**Low discrimination**					**05515500**
					GRA	
Lactobacillus fermentum		87.36	0.74	SUC 87	99	
Bifidobacterium longum subsp. longum		11.45	0.52		99	
Bacteroides vulgatus		1.05	0.20		1	
05515501	**Low discrimination**					**05515501**
Lactobacillus fermentum		88.42	0.40	SUC 87		
Bifidobacterium longum subsp. longum		11.58	0.17	URE 1		
05515502	**Low discrimination**					**05515502**
Lactobacillus fermentum		87.30	0.74	SUC 87		
Bifidobacterium longum subsp. longum		11.44	0.52			
Bifidobacterium adolescentis		1.06	0.36			
05515503	**Low discrimination**					**05515503**
Lactobacillus fermentum		88.42	0.40	SUC 87		
Bifidobacterium longum subsp. longum		11.58	0.17	URE 1		
05515510	**Good identification**					**05515510**
Bifidobacterium longum subsp. longum		95.67	0.56	SUC 99		
Lactobacillus fermentum		4.33	0.40			
05515512	**Low discrimination**					**05515512**
Bifidobacterium longum subsp. longum		89.48	0.56	SUC 99		
Bifidobacterium adolescentis		5.98	0.37			
Lactobacillus fermentum		4.05	0.40			
05515520	**Low discrimination**					**05515520**
					GRA	
Bacteroides vulgatus		58.06	0.54	SUC 96	1	
Bifidobacterium longum subsp. longum		25.55	0.62	RHA 90	99	
Lactobacillus fermentum		16.25	0.66		99	
05515522	**Low discrimination**					**05515522**
					GRA	
Bifidobacterium longum subsp. longum		47.78	0.62	SUC 99	99	
Lactobacillus fermentum		30.39	0.66		99	
Parabacteroides distasonis		20.49	0.60		1	
05515530	**Very good identification**					**05515530**
					GRA	
Bifidobacterium longum subsp. longum		98.17	0.66	SUC 99	99	
Bacteroides vulgatus		1.32	0.20		1	
Lactobacillus fermentum		0.37	0.31		99	
05515532	**Low discrimination**					**05515532**
					GRA	
Bifidobacterium longum subsp. longum		87.77	0.66	SUC 99	99	
Parabacteroides distasonis		11.90	0.55		1	
Lactobacillus fermentum		0.33	0.31		99	
05515540	**Low discrimination**					**05515540**
Lactobacillus fermentum		88.42	0.40	SUC 87		
Bifidobacterium longum subsp. longum		11.58	0.17	SOR 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
05515542	**Low discrimination**					**05515542**
Lactobacillus fermentum		78.12	0.40	SUC 87		
Bifidobacterium adolescentis		11.65	0.20	SOR 1		
Bifidobacterium longum subsp. longum		10.23	0.17			
05515600	**Low discrimination**					**05515600**
					GRA	
Lactobacillus fermentum		84.99	0.36	SUC 87	99	
Bacteroides vulgatus		15.01	0.02	RHA 1	1	
05515602	**Good identification**					**05515602**
Lactobacillus fermentum		100.0	0.36	SUC 87		
05515620	**Good identification**					**05515620**
					GRA	
Bacteroides vulgatus		98.13	0.36	SUC 96	1	
Lactobacillus fermentum		1.87	0.28	LAC 99	99	
05515700	**Acceptable identification**					**05515700**
					GRA	
Bacteroides vulgatus		90.45	0.36	SUC 96	1	
Lactobacillus fermentum		8.44	0.40	NAG 99	99	
Bifidobacterium longum subsp. longum		1.11	0.17		99	
05515702	**Low discrimination**					**05515702**
					GRA	
Lactobacillus fermentum		51.21	0.40	SUC 87	99	
Parabacteroides distasonis		34.52	0.34	RHA 1	1	
Bifidobacterium longum subsp. longum		6.71	0.17		99	
05515720	**Very good identification**					**05515720**
					GRA	
Bacteroides vulgatus		99.86	0.70	SUC 96	1	
Parabacteroides distasonis		0.06	0.34		1	
Bifidobacterium longum subsp. longum		0.05	0.28		99	
05515721	**Good identification**					**05515721**
Bacteroides vulgatus		100.0	0.36	SUC 96		
05515722	**Low discrimination**					**05515722**
					GRA	
Parabacteroides distasonis		85.11	0.68		1	
Bacteroides vulgatus		13.67	0.36		1	
Bifidobacterium longum subsp. longum		0.67	0.28		99	
05515730	**Low discrimination**					**05515730**
					GRA	
Bacteroides vulgatus		89.52	0.36	SUC 96	1	
Bifidobacterium longum subsp. longum		7.45	0.32	MLZ 1	99	
Parabacteroides distasonis		3.03	0.29		1	
05515732	**Very good identification**					**05515732**
					GRA	
Parabacteroides distasonis		96.82	0.63		1	
Bifidobacterium longum subsp. longum		2.40	0.32		99	
Clostridium clostridioforme		0.49	0.18		50	
05515760	**Good identification**					**05515760**
Bacteroides vulgatus		100.0	0.36	SUC 96		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05517000	**Low discrimination**						**05517000**
					GRA	CAT	
Lactobacillus fermentum		36.36	0.36	SUC 87	99	1	
Propionibacterium granulosum		32.90	0.30	MAN 1	99	90	
Prevotella intermedia		24.76	0.20		1	1	
05517002	**Low discrimination**						**05517002**
					CAT		
Lactobacillus fermentum		81.56	0.36	SUC 87	1		
Propionibacterium granulosum		18.45	0.20	MAN 1	90		
05517100	**Good identification**						**05517100**
Lactobacillus fermentum		96.29	0.40	SUC 87			
Actinomyces israelii		3.71	0.16	MAN 1			
05517102	**Low discrimination**						**05517102**
Lactobacillus fermentum		81.25	0.40	SUC 87			
Bifidobacterium adolescentis		11.99	0.20	MAN 1			
Actinomyces israelii		4.69	0.19				
05517322	**Good identification**						**05517322**
Parabacteroides distasonis		96.82	0.37	MAN 1			
Prevotella oralis		3.18	-.01				
05517400	**Good identification**						**05517400**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05517402	**Good identification**						**05517402**
Lactobacillus fermentum		100.0	0.36	SUC 87			
05517500	**Low discrimination**						**05517500**
Lactobacillus fermentum		86.02	0.40	SUC 87			
Bifidobacterium longum subsp. longum		11.27	0.17	MAN 1			
Actinomyces israelii		2.71	0.15				
05517502	**Low discrimination**						**05517502**
Lactobacillus fermentum		46.18	0.40	SUC 87			
Bifidobacterium adolescentis		45.59	0.34	MAN 1			
Bifidobacterium longum subsp. longum		6.05	0.17				
05517512	**Low discrimination**						**05517512**
Bifidobacterium adolescentis		84.40	0.36	SUC 99			
Bifidobacterium longum subsp. longum		15.60	0.21				
05517720	**Good identification**						**05517720**
Bacteroides vulgatus		100.0	0.36	SUC 96			
05520040	**Acceptable identification**						**05520040**
					GRA	SPO	
Clostridium botulinum biovar B		93.76	0.39	SUC 90	99	99	
Clostridium botulinum biovar A		5.73	0.21	RAF 1	99	99	
Prevotella disiens		0.51	-.03		1	1	
05521000	**Low discrimination**						**05521000**
					COC	GRA	CAT

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		62.92	0.41	SUC 93	1	99	90	
Gemella morbillorum		21.21	0.26	FRU 99	99	99	1	
Prevotella intermedia		11.91	0.20		1	1	1	
05521120	**Low discrimination**							**05521120**
					GRA			
Clostridium perfringens		51.67	0.36	SUC 91	99			
Prevotella bivia		30.21	0.19	FRU 99	1			
Prevotella melaninogenica		7.78	0.15		1			
05522000	**Good identification**							**05522000**
					GRA	CAT		
Propionibacterium granulosum		96.83	0.39	SUC 93	99	90		
Prevotella disiens		3.17	-.03	FRU 99	1	1		
05523000	**Good identification**							**05523000**
Propionibacterium granulosum		100.0	0.39	SUC 93				
05525122	**Good identification**							**05525122**
Parabacteroides distasonis		100.0	0.41	FRU 99				
05525132	**Good identification**							**05525132**
Parabacteroides distasonis		100.0	0.36	FRU 99				
05525322	**Good identification**							**05525322**
Parabacteroides distasonis		100.0	0.49	FRU 99				
05525332	**Good identification**							**05525332**
Parabacteroides distasonis		100.0	0.44	FRU 99				
05525520	**Good identification**							**05525520**
					GRA			
Bifidobacterium longum subsp. longum		96.67	0.35	SUC 99	99			
Bacteroides vulgatus		3.33	-.04	FRU 90	1			
05525522	**Low discrimination**							**05525522**
					GRA			
Bifidobacterium longum subsp. longum		58.47	0.35	SUC 99	99			
Parabacteroides distasonis		41.53	0.37	FRU 90	1			
05525530	**Good identification**							**05525530**
Bifidobacterium longum subsp. longum		100.0	0.39	SUC 99				
05525532	**Low discrimination**							**05525532**
					GRA			
Bifidobacterium longum subsp. longum		81.66	0.39	SUC 99	99			
Parabacteroides distasonis		18.34	0.32	FRU 90	1			
05525722	**Good identification**							**05525722**
Parabacteroides distasonis		100.0	0.45	FRU 99				
05525732	**Good identification**							**05525732**
Parabacteroides distasonis		100.0	0.40	FRU 99				
05530000	**Low discrimination**							**05530000**
					GRA	CAT		
Propionibacterium granulosum		80.88	0.75	SUC 93	99	90		
Prevotella intermedia		15.31	0.55		1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella disiens		2.00	0.32		1	1		
05530001	**Low discrimination**							**05530001**
					GRA	CAT		
Propionibacterium granulosum		81.56	0.41	SUC 93	99	90		
Prevotella intermedia		15.44	0.20	URE 1	1	1		
Prevotella disiens		2.02	-.03		1	1		
05530002	**Very good identification**							**05530002**
					GRA	CAT		
Propionibacterium granulosum		98.44	0.65	SUC 93	99	90		
Prevotella intermedia		0.75	0.20		1	1		
Fusobacterium mortiferum		0.53	0.29		1	5		
05530010	**Very good identification**							**05530010**
					GRA	SPO	CAT	
Propionibacterium granulosum		97.71	0.60	SUC 93	99	1	90	
Prevotella intermedia		1.37	0.20	MLZ 12	1	1	1	
Clostridium botulinum biovar B		0.65	0.10		99	99	1	
05530012	**Good identification**							**05530012**
Propionibacterium granulosum		100.0	0.50	SUC 93				
05530020	**Very good identification**							**05530020**
					GRA	CAT		
Propionibacterium granulosum		97.60	0.65	SUC 93	99	90		
Clostridium perfringens		0.82	0.22		99	1		
Prevotella intermedia		0.75	0.20		1	1		
05530022	**Good identification**							**05530022**
					CAT			
Propionibacterium granulosum		98.45	0.54	SUC 93	90			
Clostridium perfringens		1.55	0.16		1			
05530030	**Good identification**							**05530030**
					SPO	CAT		
Propionibacterium granulosum		98.71	0.50	SUC 93	1	90		
Clostridium botulinum biovar B		1.29	0.05	MLZ 12	99	1		
05530032	**Good identification**							**05530032**
Propionibacterium granulosum		100.0	0.39	SUC 93				
05530040	**Acceptable identification**							**05530040**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		93.55	0.50	SUC 90	99	99	1	
Propionibacterium granulosum		5.18	0.41	RAF 1	99	1	90	
Prevotella intermedia		0.98	0.20		1	1	1	
05530050	**Good identification**							**05530050**
					SPO	CAT		
Clostridium botulinum biovar B		98.49	0.44	SUC 90	99	1		
Propionibacterium granulosum		1.51	0.26	RAF 1	1	90		
05530060	**Good identification**							**05530060**
					CAT	SPO		
Clostridium botulinum biovar B		96.85	0.44	SUC 90	1	99		
Propionibacterium granulosum		2.72	0.30	RAF 1	90	1		
Clostridium perfringens		0.43	0.09		1	84		
05530070	**Good identification**							**05530070**
					SPO	CAT		
Clostridium botulinum biovar B		99.23	0.39	SUC 90	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.77	0.16	RAF 1	1	90		
05530100	**Low discrimination**							**05530100**
					GRA	CAT		
Fusobacterium mortiferum		68.84	0.52	MLT 11	1	5		
Propionibacterium granulosum		24.47	0.41		99	90		
Prevotella intermedia		4.63	0.20		1	1		
05530102	**Low discrimination**							**05530102**
					GRA	CAT		
Fusobacterium mortiferum		46.28	0.35	MLT 11	1	5		
Propionibacterium granulosum		37.02	0.30	BGL 10	99	90		
Actinomyces viscosus		11.63	0.12		99	76		
05530120	**Low discrimination**							**05530120**
					GRA	CAT		
Clostridium perfringens		61.80	0.36	SUC 91	99	1		
Prevotella bivia		22.75	0.15	GAL 99	1	1		
Propionibacterium granulosum		10.19	0.30		99	90		
05530200	**Acceptable identification**							**05530200**
					GRA	CAT		
Propionibacterium granulosum		93.12	0.49	SUC 93	99	90		
Prevotella intermedia		5.76	0.20	RHA 3	1	1		
Prevotella disiens		0.75	-.03		1	1		
05530202	**Good identification**							**05530202**
Propionibacterium granulosum		100.0	0.39	SUC 93				
05530220	**Good identification**							**05530220**
Propionibacterium granulosum		100.0	0.39	SUC 93				
05530400	**Low discrimination**							**05530400**
					COC	GRA	CAT	
Propionibacterium granulosum		76.27	0.41	SUC 93	1	99	90	
Prevotella intermedia		14.43	0.20	ARA 1	1	1	1	
Gemella morbillorum		6.48	0.15		99	99	1	
05531000	**Low discrimination**							**05531000**
					GRA	CAT		
Propionibacterium granulosum		81.67	0.75	SUC 93	99	90		
Prevotella intermedia		15.46	0.55		1	1		
Fusobacterium mortiferum		1.49	0.48		1	5		
05531001	**Low discrimination**							**05531001**
					GRA	CAT		
Propionibacterium granulosum		73.43	0.41	SUC 93	99	90		
Prevotella intermedia		13.90	0.20	URE 1	1	1		
Actinomyces naeslundii		11.33	0.42		99	25		
05531002	**Good identification**							**05531002**
					CAT			
Propionibacterium granulosum		94.99	0.65	SUC 93	90			
Lactobacillus fermentum		1.82	0.40		1			
Clostridium perfringens		1.14	0.24		1			
05531010	**Very good identification**							**05531010**
					GRA	CAT		
Propionibacterium granulosum		98.49	0.60	SUC 93	99	90		
Prevotella intermedia		1.38	0.20	MLZ 12	1	1		
Fusobacterium mortiferum		0.13	0.14		1	5		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05531012	**Good identification**							**05531012**
Propionibacterium granulosum		100.0	0.50	SUC 93				
05531020	**Low discrimination**							**05531020**
					GRA	CAT		
Propionibacterium granulosum		54.00	0.65	SUC 93	99	90		
Clostridium perfringens		44.68	0.56		99	1		
Prevotella intermedia		0.41	0.20		1	1		
05531022	**Low discrimination**							**05531022**
					CAT			
Clostridium perfringens		59.14	0.50	SUC 91	1			
Propionibacterium granulosum		37.97	0.54	LAC 88	90			
Lactobacillus fermentum		0.97	0.32		1			
05531023	**Low discrimination**							**05531023**
					CAT			
Actinomyces naeslundii		49.13	0.42	SUC 99	25			
Clostridium perfringens		30.98	0.16		1			
Propionibacterium granulosum		19.89	0.20		90			
05531030	**Acceptable identification**							**05531030**
					CAT			
Propionibacterium granulosum		94.23	0.50	SUC 93	90			
Clostridium perfringens		5.78	0.22	MLZ 12	1			
05531032	**Low discrimination**							**05531032**
					GRA	CAT		
Propionibacterium granulosum		85.54	0.39	SUC 93	99	90		
Clostridium perfringens		9.87	0.16	MLZ 12	99	1		
Parabacteroides distasonis		4.59	0.20		1	76		
05531040	**Low discrimination**							**05531040**
					GRA	SPO	CAT	
Propionibacterium granulosum		65.42	0.41	SUC 93	99	1	90	
Prevotella intermedia		12.38	0.20	SOR 1	1	1	1	
Clostridium botulinum biovar B		11.93	0.15		99	99	1	
05531060	**Acceptable identification**							**05531060**
					CAT	SPO		
Clostridium perfringens		91.59	0.43	SUC 91	1	84		
Propionibacterium granulosum		5.87	0.30	LAC 88	90	1		
Clostridium botulinum biovar B		2.11	0.10		1	99		
05531062	**Good identification**							**05531062**
					CAT			
Clostridium perfringens		95.79	0.38	SUC 91	1			
Propionibacterium granulosum		3.26	0.20	LAC 88	90			
Actinomyces naeslundii		0.95	0.26		25			
05531100	**Low discrimination**							**05531100**
					GRA	CAT		
Clostridium perfringens		40.47	0.45	SUC 91	99	1		
Fusobacterium mortiferum		36.76	0.55	NAG 97	1	5		
Propionibacterium granulosum		8.71	0.41		99	90		
05531101	**Low discrimination**							**05531101**
					SPO			
Actinomyces naeslundii		76.64	0.49	SUC 99	1			
Clostridium perfringens		8.18	0.11		84			
Clostridium chauvoei		7.74	0.09		99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
05531102	**Low discrimination**						**05531102**
					GRA		
Clostridium perfringens		51.44	0.39	SUC 91	99		
Lactobacillus fermentum		18.17	0.44	NAG 97	99		
Fusobacterium mortiferum		11.03	0.38		1		
05531103	**Acceptable identification**						**05531103**
Actinomyces naeslundii		92.02	0.49	SUC 99			
Clostridium perfringens		4.62	0.05				
Actinomyces viscosus		3.36	0.01				
05531120	**Good identification**						**05531120**
					GRA	SPO	
Clostridium perfringens		93.77	0.71	SUC 91	99	84	
Clostridium septicum		4.21	0.41		99	99	
Prevotella melaninogenica		1.28	0.32		1	1	
05531121	**Low discrimination**						**05531121**
					SPO		
Clostridium perfringens		73.64	0.36	SUC 91	84		
Actinomyces naeslundii		21.33	0.49	URE 1	1		
Clostridium septicum		3.30	0.07		99		
05531122	**Low discrimination**						**05531122**
					GRA		
Clostridium perfringens		86.06	0.65	SUC 91	99		
Parabacteroides distasonis		10.15	0.58		1		
Prevotella oralis		2.38	0.35		1		
05531130	**Low discrimination**						**05531130**
					GRA	SPO	
Clostridium perfringens		87.47	0.36	SUC 91	99	84	
Clostridium septicum		3.92	0.07	MLZ 1	99	99	
Parabacteroides distasonis		2.61	0.20		1	1	
05531132	**Low discrimination**						**05531132**
					GRA		
Parabacteroides distasonis		85.14	0.54	XYL 90	1		
Clostridium perfringens		13.55	0.31		99		
Bifidobacterium longum subsp. longum		0.64	0.13		99		
05531141	**Good identification**						**05531141**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05531143	**Good identification**						**05531143**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05531160	**Very good identification**						**05531160**
					SPO		
Clostridium perfringens		99.51	0.58	SUC 91	84		
Clostridium septicum		0.24	0.07		99		
Actinomyces naeslundii		0.18	0.34		1		
05531162	**Good identification**						**05531162**
					GRA		
Clostridium perfringens		98.86	0.53	SUC 91	99		
Parabacteroides distasonis		0.62	0.24		1		
Actinomyces naeslundii		0.38	0.34		99		
05531200	**Acceptable identification**						**05531200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Propionibacterium granulosum		93.65	0.49	SUC 93	99	90	
Prevotella intermedia		5.79	0.20	RHA 3	1	1	
Fusobacterium mortiferum		0.56	0.14		1	5	
05531202	**Good identification**						**05531202**
Propionibacterium granulosum		100.0	0.39	SUC 93			
05531220	**Low discrimination**						**05531220**
					CAT		
Propionibacterium granulosum		78.73	0.39	SUC 93	90		
Clostridium perfringens		21.27	0.22	RHA 3	1		
05531320	**Low discrimination**						**05531320**
					GRA	SPO	
Clostridium perfringens		78.47	0.36	SUC 91	99	84	
Parabacteroides distasonis		13.06	0.32	RHA 1	1	1	
Clostridium septicum		3.52	0.07		99	99	
05531322	**Good identification**						**05531322**
					GRA		
Parabacteroides distasonis		94.16	0.67	XYL 90	1		
Prevotella oralis		3.16	0.29		1		
Clostridium perfringens		2.69	0.31		99		
05531332	**Very good identification**						**05531332**
Parabacteroides distasonis		100.0	0.62	XYL 90			
05531400	**Low discrimination**						**05531400**
					GRA	CAT	
Propionibacterium granulosum		57.92	0.41	SUC 93	99	90	
Lactobacillus fermentum		27.41	0.40	ARA 1	99	1	
Prevotella intermedia		10.96	0.20		1	1	
05531402	**Low discrimination**						**05531402**
					CAT		
Lactobacillus fermentum		65.44	0.40	SUC 87	1		
Propionibacterium granulosum		34.57	0.30	TRE 1	90		
05531500	**Low discrimination**						**05531500**
Lactobacillus fermentum		57.12	0.44	SUC 87			
Bifidobacterium longum subsp. longum		36.26	0.33	TRE 1			
Clostridium perfringens		3.47	0.11				
05531502	**Low discrimination**						**05531502**
Lactobacillus fermentum		55.53	0.44	SUC 87			
Bifidobacterium longum subsp. longum		35.25	0.33	TRE 1			
Bifidobacterium adolescentis		3.92	0.18				
05531510	**Good identification**						**05531510**
Bifidobacterium longum subsp. longum		100.0	0.37	SUC 99			
05531512	**Low discrimination**						**05531512**
					GRA		
Bifidobacterium longum subsp. longum		89.84	0.37	SUC 99	99		
Bifidobacterium adolescentis		7.17	0.20		99		
Parabacteroides distasonis		2.99	0.16		1		
05531520	**Low discrimination**						**05531520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		50.01	0.43	SUC 99			
Clostridium perfringens		38.69	0.36				
Lactobacillus fermentum		6.57	0.35				
05531522	**Low discrimination**						**05531522**
					GRA		
Parabacteroides distasonis		63.37	0.55	XYL 90	1		
Bifidobacterium longum subsp. longum		24.33	0.43		99		
Clostridium perfringens		8.86	0.31		99		
05531530	**Good identification**						**05531530**
					GRA		
Bifidobacterium longum subsp. longum		98.73	0.47	SUC 99	99		
Parabacteroides distasonis		0.82	0.16		1		
Clostridium perfringens		0.45	0.02		99		
05531532	**Low discrimination**						**05531532**
					GRA		
Bifidobacterium longum subsp. longum		54.84	0.47	SUC 99	99		
Parabacteroides distasonis		45.16	0.50		1		
05531722	**Very good identification**						**05531722**
					GRA		
Parabacteroides distasonis		99.79	0.63	XYL 90	1		
Bifidobacterium longum subsp. longum		0.13	0.09		99		
Clostridium clostridioforme		0.09	0.05		50		
05531732	**Very good identification**						**05531732**
					GRA		
Parabacteroides distasonis		98.89	0.58	XYL 90	1		
Clostridium clostridioforme		0.71	0.16		50		
Bifidobacterium longum subsp. longum		0.40	0.13		99		
05532000	**Very good identification**						**05532000**
					GRA	CAT	
Propionibacterium granulosum		99.68	0.73	SUC 93	99	90	
Prevotella intermedia		0.25	0.20		1	1	
Prevotella disiens		0.03	-.03		1	1	
05532001	**Good identification**						**05532001**
Propionibacterium granulosum		98.37	0.39	SUC 93			
Actinomyces naeslundii		1.63	0.23	URE 1			
05532002	**Very good identification**						**05532002**
Propionibacterium granulosum		100.0	0.62	SUC 93			
05532010	**Very good identification**						**05532010**
					SPO	CAT	
Propionibacterium granulosum		99.60	0.58	SUC 93	1	90	
Clostridium difficile		0.40	0.04	MLZ 12	99	1	
05532012	**Good identification**						**05532012**
Propionibacterium granulosum		100.0	0.48	SUC 93			
05532020	**Very good identification**						**05532020**
Propionibacterium granulosum		100.0	0.62	SUC 93			
05532022	**Good identification**						**05532022**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.52	SUC 93			
05532030	**Good identification**						**05532030**
Propionibacterium granulosum		100.0	0.48	SUC 93			
05532032	**Good identification**						**05532032**
Propionibacterium granulosum		100.0	0.37	SUC 93			
05532040	**Low discrimination**						**05532040**
					SPO	CAT	
Propionibacterium granulosum		80.54	0.39	SUC 93	1	90	
Clostridium botulinum biovar B		19.47	0.15	SOR 1	99	1	
05532100	**Low discrimination**						**05532100**
					GRA	CAT	
Propionibacterium granulosum		52.64	0.39	SUC 93	99	90	
Propionibacterium propionicum		43.72	0.42	LAC 1	99	1	
Fusobacterium mortiferum		1.98	0.18		1	5	
05532102	**Low discrimination**						**05532102**
					CAT		
Propionibacterium propionicum		81.58	0.45	NIT 86	1		
Propionibacterium granulosum		16.37	0.28	SUC 99	90		
Actinomyces naeslundii		2.06	0.28		25		
05532140	**Good identification**						**05532140**
Propionibacterium propionicum		100.0	0.42	NIT 86			
05532142	**Good identification**						**05532142**
Propionibacterium propionicum		100.0	0.45	NIT 86			
05532200	**Good identification**						**05532200**
Propionibacterium granulosum		100.0	0.47	SUC 93			
05532202	**Good identification**						**05532202**
Propionibacterium granulosum		100.0	0.37	SUC 93			
05532220	**Good identification**						**05532220**
Propionibacterium granulosum		100.0	0.37	SUC 93			
05532400	**Good identification**						**05532400**
Propionibacterium granulosum		100.0	0.39	SUC 93			
05533000	**Very good identification**						**05533000**
					GRA	CAT	
Propionibacterium granulosum		99.46	0.73	SUC 93	99	90	
Actinomyces naeslundii		0.27	0.44		99	25	
Prevotella intermedia		0.25	0.20		1	1	
05533001	**Low discrimination**						**05533001**
Propionibacterium granulosum		71.57	0.39	SUC 93			
Actinomyces naeslundii		28.43	0.47	URE 1			
05533002	**Very good identification**						**05533002**
Propionibacterium granulosum		98.94	0.62	SUC 93			
Actinomyces naeslundii		1.06	0.44				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05533003	**Low discrimination**					**05533003**
Actinomyces naeslundii		61.37	0.47	SUC 99		
Propionibacterium granulosum		38.63	0.28			
05533010	**Very good identification**					**05533010**
Propionibacterium granulosum		100.0	0.58	SUC 93		
05533012	**Good identification**					**05533012**
Propionibacterium granulosum		100.0	0.48	SUC 93		
05533020	**Very good identification**					**05533020**
					CAT	
Propionibacterium granulosum		97.87	0.62	SUC 93	90	
Clostridium perfringens		1.08	0.22		1	
Actinomyces naeslundii		1.05	0.44		25	
05533021	**Low discrimination**					**05533021**
Actinomyces naeslundii		61.37	0.47	SUC 99		
Propionibacterium granulosum		38.63	0.28			
05533022	**Acceptable identification**					**05533022**
					CAT	
Propionibacterium granulosum		94.02	0.52	SUC 93	90	
Actinomyces naeslundii		4.02	0.44		25	
Clostridium perfringens		1.96	0.16		1	
05533023	**Low discrimination**					**05533023**
Actinomyces naeslundii		86.41	0.47	SUC 99		
Propionibacterium granulosum		13.60	0.18			
05533030	**Good identification**					**05533030**
Propionibacterium granulosum		100.0	0.48	SUC 93		
05533032	**Good identification**					**05533032**
Propionibacterium granulosum		100.0	0.37	SUC 93		
05533040	**Good identification**					**05533040**
Propionibacterium granulosum		95.54	0.39	SUC 93		
Actinomyces naeslundii		4.46	0.31	SOR 1		
05533100	**Low discrimination**					**05533100**
					CAT	
Propionibacterium granulosum		37.00	0.39	SUC 93	90	
Propionibacterium propionicum		30.73	0.42	LAC 1	1	
Actinomyces naeslundii		27.88	0.51		25	
05533101	**Acceptable identification**					**05533101**
Actinomyces naeslundii		92.45	0.54	SUC 99		
Propionibacterium propionicum		7.55	0.26			
05533102	**Low discrimination**					**05533102**
					CAT	
Propionibacterium propionicum		54.67	0.45	NIT 86	1	
Actinomyces naeslundii		33.07	0.51	SUC 99	25	
Propionibacterium granulosum		10.97	0.28		90	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
05533103	**Low discrimination**					**05533103**
	Actinomyces naeslundii	89.09	0.54	SUC 99		
	Propionibacterium propionicum	10.91	0.29			
05533120	**Low discrimination**					**05533120**
					CAT	
	Clostridium perfringens	64.20	0.36	SUC 91	1	
	Actinomyces naeslundii	24.06	0.51	MAN 1	25	
	Propionibacterium granulosum	7.98	0.28		90	
05533121	**Good identification**					**05533121**
	Actinomyces naeslundii	98.24	0.54	SUC 99		
	Clostridium perfringens	1.77	0.02			
05533123	**Good identification**					**05533123**
	Actinomyces naeslundii	100.0	0.54	SUC 99		
05533140	**Low discrimination**					**05533140**
	Propionibacterium propionicum	86.20	0.42	NIT 86		
	Actinomyces naeslundii	13.80	0.38	SUC 99		
05533141	**Low discrimination**					**05533141**
	Actinomyces naeslundii	68.37	0.41	SUC 99		
	Propionibacterium propionicum	31.63	0.26			
05533142	**Acceptable identification**					**05533142**
	Propionibacterium propionicum	90.36	0.45	NIT 86		
	Actinomyces naeslundii	9.65	0.38	SUC 99		
05533143	**Low discrimination**					**05533143**
	Actinomyces naeslundii	59.04	0.41	SUC 99		
	Propionibacterium propionicum	40.97	0.29			
05533161	**Good identification**					**05533161**
	Actinomyces naeslundii	100.0	0.41	SUC 99		
05533163	**Good identification**					**05533163**
	Actinomyces naeslundii	100.0	0.41	SUC 99		
05533200	**Good identification**					**05533200**
	Propionibacterium granulosum	100.0	0.47	SUC 93		
05533202	**Good identification**					**05533202**
	Propionibacterium granulosum	100.0	0.37	SUC 93		
05533220	**Good identification**					**05533220**
	Propionibacterium granulosum	100.0	0.37	SUC 93		
05533400	**Good identification**					**05533400**
	Propionibacterium granulosum	100.0	0.39	SUC 93		
05534000	**Low discrimination**					**05534000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Propionibacterium granulosum		71.96	0.41	SUC 93	99	90	
Prevotella intermedia		13.62	0.20	XYL 1	1	1	
Fusobacterium mortiferum		6.53	0.26		1	5	
05534122	**Good identification**						**05534122**
Parabacteroides distasonis		100.0	0.41	GAL 99			
05534132	**Good identification**						**05534132**
Parabacteroides distasonis		100.0	0.36	GAL 99			
05534322	**Good identification**						**05534322**
Parabacteroides distasonis		100.0	0.49	GAL 99			
05534332	**Good identification**						**05534332**
Parabacteroides distasonis		98.97	0.44	GAL 99			
Clostridium clostridioforme		1.03	0.04				
05534522	**Low discrimination**						**05534522**
					GRA		
Parabacteroides distasonis		88.65	0.37	GAL 99	1		
Bifidobacterium longum subsp. longum		11.35	0.17		99		
05534722	**Good identification**						**05534722**
Parabacteroides distasonis		96.95	0.45	GAL 99			
Clostridium clostridioforme		3.05	0.14				
05534732	**Low discrimination**						**05534732**
Parabacteroides distasonis		79.00	0.40	GAL 99			
Clostridium clostridioforme		21.01	0.25				
05535000	**Low discrimination**						**05535000**
					GRA	CAT	
Propionibacterium granulosum		60.64	0.41	SUC 93	99	90	
Lactobacillus fermentum		16.86	0.36	XYL 1	99	1	
Prevotella intermedia		11.48	0.20		1	1	
05535002	**Low discrimination**						**05535002**
					GRA	CAT	
Lactobacillus fermentum		51.18	0.36	SUC 87	99	1	
Propionibacterium granulosum		46.03	0.30	TRE 1	99	90	
Fusobacterium mortiferum		2.78	0.13		1	5	
05535022	**Low discrimination**						**05535022**
					GRA		
Parabacteroides distasonis		73.87	0.41	LAC 99	1		
Clostridium perfringens		9.49	0.16		99		
Lactobacillus fermentum		9.04	0.28		99		
05535032	**Good identification**						**05535032**
Parabacteroides distasonis		100.0	0.36	LAC 99			
05535102	**Low discrimination**						**05535102**
					COC	GRA	
Parabacteroides distasonis		52.25	0.41	NAG 99	1	1	
Lactobacillus fermentum		31.29	0.40		1	99	
Atopobium parvulum		5.65	0.14		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05535112	**Low discrimination**					**05535112**
					GRA	
Parabacteroides distasonis		81.61	0.36	NAG 99	1	
Bifidobacterium adolescentis		13.33	0.20		99	
Bifidobacterium longum subsp. longum		5.06	0.11		99	
05535120	**Low discrimination**					**05535120**
					GRA	
Clostridium perfringens		53.49	0.36	SUC 91	99	
Parabacteroides distasonis		26.71	0.41	XYL 1	1	
Bacteroides vulgatus		5.91	0.11		1	
05535122	**Very good identification**					**05535122**
					GRA	
Parabacteroides distasonis		98.39	0.75		1	
Clostridium perfringens		0.94	0.31		99	
Prevotella oralis		0.32	0.19		1	
05535123	**Good identification**					**05535123**
Parabacteroides distasonis		100.0	0.41	URE 1		
05535130	**Low discrimination**					**05535130**
					GRA	
Parabacteroides distasonis		77.78	0.36	BGL 99	1	
Bifidobacterium longum subsp. longum		19.30	0.21		99	
Clostridium perfringens		2.92	0.02		99	
05535132	**Very good identification**					**05535132**
					GRA	
Parabacteroides distasonis		99.75	0.70		1	
Bifidobacterium longum subsp. longum		0.25	0.21		99	
05535133	**Good identification**					**05535133**
Parabacteroides distasonis		100.0	0.36	URE 1		
05535162	**Low discrimination**					**05535162**
					GRA	
Parabacteroides distasonis		84.78	0.41	SOR 1	1	
Clostridium perfringens		15.22	0.18		99	
05535172	**Good identification**					**05535172**
Parabacteroides distasonis		100.0	0.36	SOR 1		
05535222	**Good identification**					**05535222**
Parabacteroides distasonis		100.0	0.49	LAC 99		
05535232	**Good identification**					**05535232**
Parabacteroides distasonis		100.0	0.44	LAC 99		
05535302	**Good identification**					**05535302**
Parabacteroides distasonis		100.0	0.49	NAG 99		
05535312	**Good identification**					**05535312**
Parabacteroides distasonis		96.97	0.44	NAG 99		
Clostridium clostridioforme		3.03	0.13			
05535320	**Low discrimination**					**05535320**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		59.85	0.49	BGL 99	1	
Bacteroides vulgatus		39.75	0.27		1	
Clostridium perfringens		0.40	0.02		99	
05535322	**Very good identification**					**05535322**
Parabacteroides distasonis		99.95	0.83			
Prevotella oralis		0.05	0.13			
Bacteroides vulgatus		0.01	-.07			
05535323	**Good identification**					**05535323**
Parabacteroides distasonis		100.0	0.49	URE 1		
05535330	**Good identification**					**05535330**
Parabacteroides distasonis		98.77	0.44	BGL 99		
Bacteroides vulgatus		1.23	-.07			
05535332	**Very good identification**					**05535332**
Parabacteroides distasonis		99.97	0.78			
Clostridium clostridioforme		0.03	0.13			
05535333	**Good identification**					**05535333**
Parabacteroides distasonis		100.0	0.44	URE 1		
05535362	**Good identification**					**05535362**
Parabacteroides distasonis		100.0	0.49	SOR 1		
05535372	**Good identification**					**05535372**
Parabacteroides distasonis		100.0	0.44	SOR 1		
05535400	**Acceptable identification**					**05535400**
Lactobacillus fermentum		94.93	0.36	SUC 87		
Bifidobacterium longum subsp. longum		5.07	0.07	TRE 1		
05535402	**Acceptable identification**					**05535402**
Lactobacillus fermentum		94.93	0.36	SUC 87		
Bifidobacterium longum subsp. longum		5.07	0.07	TRE 1		
05535422	**Low discrimination**					**05535422**
					GRA	
Parabacteroides distasonis		75.32	0.37	LAC 99	1	
Lactobacillus fermentum		15.04	0.28		99	
Bifidobacterium longum subsp. longum		9.64	0.17		99	
05535500	**Low discrimination**					**05535500**
					GRA	
Bifidobacterium longum subsp. longum		75.55	0.41	SUC 99	99	
Lactobacillus fermentum		23.30	0.40		99	
Bacteroides vulgatus		1.16	-.04		1	
05535502	**Low discrimination**					**05535502**
					GRA	
Bifidobacterium longum subsp. longum		52.80	0.41	SUC 99	99	
Parabacteroides distasonis		16.67	0.37		1	
Lactobacillus fermentum		16.28	0.40		99	
05535510	**Good identification**					**05535510**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		100.0	0.45	SUC 99		
05535512	**Low discrimination**					**05535512**
					GRA	
Bifidobacterium longum subsp. longum		76.31	0.45	SUC 99	99	
Bifidobacterium adolescentis		13.59	0.34		99	
Parabacteroides distasonis		7.62	0.32		1	
05535520	**Low discrimination**					**05535520**
					GRA	
Bifidobacterium longum subsp. longum		67.30	0.52	SUC 99	99	
Bacteroides vulgatus		25.49	0.30		1	
Parabacteroides distasonis		5.31	0.37		1	
05535522	**Low discrimination**					**05535522**
					GRA	
Parabacteroides distasonis		88.31	0.71		1	
Bifidobacterium longum subsp. longum		11.30	0.52		99	
Lactobacillus fermentum		0.29	0.31		99	
05535523	**Low discrimination**					**05535523**
					GRA	
Parabacteroides distasonis		88.65	0.37	URE 1	1	
Bifidobacterium longum subsp. longum		11.35	0.17		99	
05535530	**Good identification**					**05535530**
					GRA	
Bifidobacterium longum subsp. longum		97.35	0.56	SUC 99	99	
Parabacteroides distasonis		2.43	0.32		1	
Bacteroides vulgatus		0.22	-.04		1	
05535532	**Low discrimination**					**05535532**
					GRA	
Parabacteroides distasonis		71.02	0.66		1	
Bifidobacterium longum subsp. longum		28.74	0.56		99	
Clostridium clostridioforme		0.23	0.18		50	
05535562	**Low discrimination**					**05535562**
					GRA	
Parabacteroides distasonis		88.65	0.37	SOR 1	1	
Bifidobacterium longum subsp. longum		11.35	0.17		99	
05535622	**Good identification**					**05535622**
Parabacteroides distasonis		97.26	0.45	LAC 99		
Clostridium clostridioforme		2.74	0.13			
05535632	**Low discrimination**					**05535632**
Parabacteroides distasonis		80.76	0.40	LAC 99		
Clostridium clostridioforme		19.24	0.24			
05535702	**Acceptable identification**					**05535702**
					GRA	
Parabacteroides distasonis		90.50	0.45	NAG 99	1	
Clostridium clostridioforme		8.53	0.22		50	
Bifidobacterium longum subsp. longum		0.97	0.07		99	
05535712	**Low discrimination**					**05535712**
					GRA	
Parabacteroides distasonis		54.60	0.40	NAG 99	1	
Clostridium clostridioforme		43.56	0.33		50	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		1.84	0.11		99	
05535720	**Acceptable identification**					**05535720**
					GRA	
Bacteroides vulgatus		93.25	0.47	SUC 96	1	
Parabacteroides distasonis		6.48	0.45	TRE 4	1	
Bifidobacterium longum subsp. longum		0.28	0.17		99	
05535722	**Very good identification**					**05535722**
					CAT	
Parabacteroides distasonis		99.60	0.79		76	
Bacteroides vulgatus		0.15	0.12		3	
Bacteroides thetaiotaomicron		0.12	0.13		89	
05535723	**Good identification**					**05535723**
Parabacteroides distasonis		100.0	0.45	URE 1		
05535730	**Low discrimination**					**05535730**
					GRA	
Parabacteroides distasonis		71.18	0.40	BGL 99	1	
Bacteroides vulgatus		19.22	0.12		1	
Bifidobacterium longum subsp. longum		9.60	0.21		99	
05535732	**Very good identification**					**05535732**
					GRA	
Parabacteroides distasonis		98.98	0.75		1	
Clostridium clostridioforme		0.80	0.33		50	
Bifidobacterium longum subsp. longum		0.14	0.21		99	
05535733	**Good identification**					**05535733**
Parabacteroides distasonis		100.0	0.40	URE 1		
05535762	**Good identification**					**05535762**
Parabacteroides distasonis		100.0	0.45	SOR 1		
05535772	**Good identification**					**05535772**
Parabacteroides distasonis		100.0	0.40	SOR 1		
05536000	**Good identification**					**05536000**
Propionibacterium granulosum		100.0	0.39	SUC 93		
05537000	**Good identification**					**05537000**
					CAT	
Propionibacterium granulosum		97.87	0.39	SUC 93	90	
Actinomyces israelii		2.13	0.16	XYL 1	1	
05537122	**Good identification**					**05537122**
Parabacteroides distasonis		100.0	0.41	MAN 1		
05537132	**Good identification**					**05537132**
Parabacteroides distasonis		100.0	0.36	MAN 1		
05537322	**Good identification**					**05537322**
Parabacteroides distasonis		100.0	0.49	MAN 1		
05537332	**Good identification**					**05537332**
Parabacteroides distasonis		100.0	0.44	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05537522	**Low discrimination**					**05537522**
					GRA	
Parabacteroides distasonis		88.65	0.37	MAN 1	1	
Bifidobacterium longum subsp. longum		11.35	0.17		99	
05537722	**Good identification**					**05537722**
Parabacteroides distasonis		100.0	0.45	MAN 1		
05537732	**Good identification**					**05537732**
Parabacteroides distasonis		100.0	0.40	MAN 1		
05550000	**Low discrimination**					**05550000**
Prevotella intermedia		79.31	0.55	CEL 1		
Prevotella disiens		10.35	0.32			
Fusobacterium mortiferum		5.69	0.46			
05550100	**Acceptable identification**					**05550100**
Fusobacterium mortiferum		93.62	0.52	MLT 11		
Prevotella intermedia		5.64	0.20			
Prevotella disiens		0.74	-.03			
05550102	**Low discrimination**					**05550102**
					GRA	
Fusobacterium mortiferum		51.00	0.36	MLT 11	1	
Actinomyces viscosus		40.56	0.22	BGL 10	99	
Prevotella oralis		8.44	0.09		1	
05551000	**Low discrimination**					**05551000**
					GRA	
Prevotella intermedia		84.24	0.55	CEL 1	1	
Fusobacterium mortiferum		9.07	0.49		1	
Lactobacillus fermentum		2.13	0.40		99	
05551002	**Low discrimination**					**05551002**
					GRA	
Lactobacillus fermentum		29.67	0.40	SUC 87	99	
Prevotella oralis		22.86	0.22	CEL 1	1	
Fusobacterium mortiferum		14.06	0.33		1	
05551003	**Low discrimination**					**05551003**
Actinomyces naeslundii		76.78	0.39	SUC 99		
Actinomyces viscosus		23.22	0.06			
05551021	**Good identification**					**05551021**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05551022	**Acceptable identification**					**05551022**
					GRA	
Prevotella oralis		92.87	0.44	SUC 92	1	
Clostridium perfringens		2.90	0.22	LAC 94	99	
Lactobacillus fermentum		2.12	0.32		99	
05551023	**Low discrimination**					**05551023**
					GRA	
Actinomyces naeslundii		60.10	0.39	SUC 99	99	
Prevotella oralis		39.90	0.10		1	
05551100	**Low discrimination**					**05551100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		71.40	0.56	MLT 11	1		
Lactobacillus fermentum		11.71	0.44		99		
Prevotella oralis		4.56	0.21		1		
05551101	**Low discrimination**						**05551101**
					GRA		
Actinomyces naeslundii		86.32	0.47	SUC 99	99		
Fusobacterium mortiferum		13.68	0.21		1		
05551102	**Low discrimination**						**05551102**
					GRA		
Prevotella oralis		75.35	0.43	SUC 92	1		
Lactobacillus fermentum		10.19	0.44	NAG 95	99		
Fusobacterium mortiferum		6.90	0.39		1		
05551103	**Low discrimination**						**05551103**
					GRA		
Actinomyces naeslundii		72.40	0.47	SUC 99	99		
Prevotella oralis		13.93	0.09		1		
Actinomyces viscosus		13.68	0.10		99		
05551120	**Low discrimination**						**05551120**
					GRA	SPO	
Prevotella melaninogenica		57.32	0.50	SUC 99	1	1	
Clostridium septicum		18.02	0.41	CEL 10	99	99	
Prevotella oralis		14.48	0.43		1	1	
05551121	**Low discrimination**						**05551121**
					GRA	SPO	
Actinomyces naeslundii		43.36	0.47	SUC 99	99	1	
Prevotella melaninogenica		33.01	0.15		1	1	
Clostridium septicum		10.38	0.07		99	99	
05551122	**Good identification**						**05551122**
					GRA		
Prevotella oralis		93.32	0.65	SUC 92	1		
Prevotella melaninogenica		3.70	0.37		1		
Clostridium perfringens		1.36	0.37		99		
05551222	**Good identification**						**05551222**
Prevotella oralis		99.01	0.38	SUC 92			
Parabacteroides distasonis		1.00	0.16	LAC 94			
05551302	**Good identification**						**05551302**
Prevotella oralis		98.80	0.37	SUC 92			
Parabacteroides distasonis		1.20	0.16	NAG 95			
05551320	**Low discrimination**						**05551320**
					GRA	SPO	
Prevotella oralis		85.87	0.37	SUC 92	1	1	
Prevotella melaninogenica		8.01	0.15	BGL 95	1	1	
Clostridium septicum		2.52	0.07		99	99	
05551322	**Good identification**						**05551322**
Prevotella oralis		93.92	0.58	SUC 92			
Parabacteroides distasonis		5.97	0.51				
Prevotella melaninogenica		0.09	0.03				
05551332	**Low discrimination**						**05551332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		77.20	0.46	XYL 90			
Prevotella oralis		22.80	0.24				
05551400	**Low discrimination**						**05551400**
					COC	GRA	
Lactobacillus fermentum		64.95	0.40	SUC 87	1	99	
Prevotella intermedia		25.97	0.20	CEL 1	1	1	
Gemella morbillorum		6.28	0.11		99	99	
05551402	**Good identification**						**05551402**
Lactobacillus fermentum		100.0	0.40	SUC 87			
05551500	**Acceptable identification**						**05551500**
					GRA		
Lactobacillus fermentum		91.98	0.44	SUC 87	99		
Fusobacterium mortiferum		5.67	0.21	CEL 1	1		
Bifidobacterium longum subsp. longum		2.36	0.09		99		
05551502	**Low discrimination**						**05551502**
					GRA		
Lactobacillus fermentum		75.45	0.44	SUC 87	99		
Bifidobacterium adolescentis		16.98	0.27	CEL 1	99		
Prevotella oralis		5.64	0.09		1		
05551520	**Low discrimination**						**05551520**
					GRA		
Lactobacillus fermentum		32.75	0.35	SUC 87	99		
Prevotella melaninogenica		29.06	0.15	CEL 1	1		
Bifidobacterium longum subsp. longum		10.08	0.20		99		
05551722	**Low discrimination**						**05551722**
Parabacteroides distasonis		79.40	0.47	XYL 90			
Prevotella oralis		20.60	0.24				
05551732	**Good identification**						**05551732**
Parabacteroides distasonis		97.31	0.42	XYL 90			
Clostridium clostridioforme		2.70	0.10				
05553001	**Good identification**						**05553001**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05553002	**Low discrimination**						**05553002**
Actinomyces naeslundii		68.45	0.41	SUC 99			
Propionibacterium granulosum		31.55	0.20				
05553003	**Good identification**						**05553003**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05553020	**Low discrimination**						**05553020**
Actinomyces naeslundii		68.45	0.41	SUC 99			
Propionibacterium granulosum		31.55	0.20				
05553021	**Good identification**						**05553021**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05553022	**Low discrimination**						**05553022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces naeslundii		61.42	0.41	SUC 99	99	25	
Prevotella oralis		31.51	0.10		1	1	
Propionibacterium granulosum		7.08	0.10		99	90	
05553023	**Good identification**						**05553023**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05553100	**Low discrimination**						**05553100**
					GRA		
Actinomyces naeslundii		85.15	0.49	SUC 99	99		
Fusobacterium mortiferum		10.43	0.21		1		
Actinomyces israelii		4.42	0.17		99		
05553101	**Good identification**						**05553101**
Actinomyces naeslundii		100.0	0.52	SUC 99			
05553102	**Low discrimination**						**05553102**
					GRA		
Actinomyces naeslundii		76.27	0.49	SUC 99	99		
Prevotella oralis		11.34	0.09		1		
Actinomyces israelii		5.94	0.20		99		
05553103	**Good identification**						**05553103**
Actinomyces naeslundii		100.0	0.52	SUC 99			
05553120	**Low discrimination**						**05553120**
					GRA	SPO	
Actinomyces naeslundii		49.76	0.49	SUC 99	99	1	
Prevotella melaninogenica		29.27	0.15		1	1	
Clostridium septicum		9.20	0.07		99	99	
05553121	**Good identification**						**05553121**
Actinomyces naeslundii		100.0	0.52	SUC 99			
05553123	**Good identification**						**05553123**
Actinomyces naeslundii		100.0	0.52	SUC 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
05553140	**Good identification**					**05553140**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05553141	**Good identification**					**05553141**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05553142	**Low discrimination**					**05553142**
Actinomyces naeslundii		74.58	0.36	SUC 99		
Actinomyces israelii		12.80	0.13			
Propionibacterium propionicum		12.63	0.12			
05553143	**Good identification**					**05553143**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05553160	**Good identification**					**05553160**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05553161	**Good identification**					**05553161**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05553162	**Good identification**					**05553162**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05553163	**Good identification**					**05553163**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05555000	**Low discrimination**					**05555000**
					GRA	
Lactobacillus fermentum		41.81	0.36	SUC 87	99	
Prevotella intermedia		28.46	0.20	CEL 1	1	
Fusobacterium mortiferum		22.85	0.30		1	
05555002	**Low discrimination**					**05555002**
					GRA	
Lactobacillus fermentum		78.24	0.36	SUC 87	99	
Prevotella oralis		12.69	0.07	CEL 1	1	
Fusobacterium mortiferum		4.75	0.14		1	
05555100	**Low discrimination**					**05555100**
					GRA	
Lactobacillus fermentum		51.19	0.40	SUC 87	99	
Fusobacterium mortiferum		40.00	0.36	CEL 1	1	
Actinomyces israelii		4.62	0.23		99	
05555122	**Low discrimination**					**05555122**
					CAT	
Prevotella oralis		55.04	0.49	SUC 92	1	
Parabacteroides distasonis		36.36	0.59	XYL 11	76	
Bacteroides fragilis		7.53	0.25		99	
05555132	**Good identification**					**05555132**
					CAT	
Parabacteroides distasonis		96.87	0.54		76	
Prevotella oralis		2.75	0.15		1	
Bacteroides fragilis		0.38	-.09		99	
05555322	**Low discrimination**					**05555322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		82.13	0.67			
Prevotella oralis		17.76	0.43			
Bacteroides vulgatus		0.06	-.05			
05555332	**Very good identification**					**05555332**
Parabacteroides distasonis		99.48	0.62			
Prevotella oralis		0.40	0.09			
Clostridium clostridioforme		0.12	0.07			
05555400	**Good identification**					**05555400**
Lactobacillus fermentum		100.0	0.36	SUC 87		
05555402	**Acceptable identification**					**05555402**
Lactobacillus fermentum		91.96	0.36	SUC 87		
Actinomyces israelii		4.15	0.14	CEL 1		
Bifidobacterium adolescentis		3.89	0.07			
05555500	**Low discrimination**					**05555500**
Lactobacillus fermentum		74.69	0.40	SUC 87		
Bifidobacterium longum subsp. longum		9.79	0.17	CEL 1		
Actinomyces israelii		5.51	0.21			
05555502	**Low discrimination**					**05555502**
Bifidobacterium adolescentis		65.30	0.41	SUC 99		
Lactobacillus fermentum		25.47	0.40			
Bifidobacterium longum subsp. longum		3.34	0.17			
05555512	**Low discrimination**					**05555512**
Bifidobacterium adolescentis		89.43	0.43	SUC 99		
Bifidobacterium longum subsp. longum		6.37	0.21			
Clostridium clostridioforme		2.34	0.12			
05555522	**Acceptable identification**					**05555522**
					GRA	
Parabacteroides distasonis		90.31	0.55		1	
Bifidobacterium longum subsp. longum		3.95	0.28		99	
Lactobacillus fermentum		2.51	0.31		99	
05555532	**Low discrimination**					**05555532**
					GRA	
Parabacteroides distasonis		86.51	0.51		1	
Bifidobacterium longum subsp. longum		11.97	0.32		99	
Clostridium clostridioforme		1.10	0.12		50	
05555720	**Good identification**					**05555720**
Bacteroides vulgatus		99.35	0.48	SUC 96		
Parabacteroides distasonis		0.65	0.29	CEL 5		
05555722	**Very good identification**					**05555722**
Parabacteroides distasonis		97.60	0.63			
Bacteroides vulgatus		1.53	0.14			
Clostridium clostridioforme		0.36	0.16			
05555732	**Very good identification**					**05555732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		96.98	0.59				
Clostridium clostridioforme		3.02	0.27				
05557100	**Good identification**						**05557100**
Actinomyces israelii		100.0	0.35	SUC 98			
05557102	**Low discrimination**						**05557102**
Actinomyces israelii		70.99	0.39	SUC 98			
Bifidobacterium adolescentis		29.01	0.26	SAL 92			
05557502	**Low discrimination**						**05557502**
Bifidobacterium adolescentis		76.98	0.40	SUC 99			
Actinomyces israelii		23.03	0.37				
05557512	**Good identification**						**05557512**
Bifidobacterium adolescentis		96.47	0.41	SUC 99			
Actinomyces israelii		3.53	0.23				
05565122	**Good identification**						**05565122**
Parabacteroides distasonis		100.0	0.36	FRU 99			
05565322	**Good identification**						**05565322**
Parabacteroides distasonis		100.0	0.44	FRU 99			
05565332	**Good identification**						**05565332**
Parabacteroides distasonis		100.0	0.39	FRU 99			
05565722	**Good identification**						**05565722**
Parabacteroides distasonis		100.0	0.40	FRU 99			
05565732	**Good identification**						**05565732**
Parabacteroides distasonis		95.23	0.36	FRU 99			
Clostridium clostridioforme		4.77	0.08				
05570000	**Low discrimination**						**05570000**
					GRA	CAT	
Propionibacterium granulosum		61.89	0.41	SUC 93	99	90	
Fusobacterium mortiferum		24.87	0.37	CEL 1	1	5	
Prevotella intermedia		11.71	0.20		1	1	
05570100	**Good identification**						**05570100**
					GRA		
Fusobacterium mortiferum		98.30	0.44	MLT 11	1		
Actinomyces naeslundii		1.70	0.25		99		
05571000	**Low discrimination**						**05571000**
					GRA	CAT	
Propionibacterium granulosum		42.02	0.41	SUC 93	99	90	
Fusobacterium mortiferum		25.33	0.40	CEL 1	1	5	
Lactobacillus jensenii		17.86	0.28		99	1	
05571001	**Good identification**						**05571001**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05571003	**Good identification**						**05571003**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	SUC 99			
05571021	**Good identification**						**05571021**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05571023	**Good identification**						**05571023**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05571100	**Low discrimination**						**05571100**
					COC	GRA	
Fusobacterium mortiferum		49.89	0.47	MLT 11	1	1	
Atopobium parvulum		20.68	0.29		99	99	
Actinomyces naeslundii		13.77	0.49		1	99	
05571101	**Good identification**						**05571101**
					GRA		
Actinomyces naeslundii		97.62	0.52	SUC 99	99		
Fusobacterium mortiferum		2.38	0.12		1		
05571102	**Low discrimination**						**05571102**
					GRA		
Actinomyces naeslundii		27.63	0.49	SUC 99	99		
Prevotella oralis		25.95	0.22		1		
Lactobacillus acidophilus		14.71	0.19		99		
05571103	**Good identification**						**05571103**
Actinomyces naeslundii		100.0	0.52	SUC 99			
05571120	**Low discrimination**						**05571120**
					SPO		
Clostridium septicum		74.18	0.49	RAF 1	99		
Clostridium perfringens		21.77	0.47		84		
Actinomyces naeslundii		1.35	0.49		1		
05571121	**Low discrimination**						**05571121**
					SPO		
Actinomyces naeslundii		67.64	0.52	SUC 99	1		
Clostridium septicum		25.02	0.15		99		
Clostridium perfringens		7.34	0.13		84		
05571122	**Low discrimination**						**05571122**
					GRA		
Prevotella oralis		44.59	0.44	SUC 92	1		
Parabacteroides distasonis		28.87	0.54	TRE 6	1		
Clostridium perfringens		18.95	0.41		99		
05571123	**Low discrimination**						**05571123**
					GRA		
Actinomyces naeslundii		80.06	0.52	SUC 99	99		
Prevotella oralis		9.62	0.10		1		
Parabacteroides distasonis		6.23	0.20		1		
05571132	**Good identification**						**05571132**
					GRA		
Parabacteroides distasonis		95.26	0.49	XYL 90	1		
Prevotella oralis		2.76	0.10		1		
Clostridium perfringens		1.17	0.07		99		
05571140	**Low discrimination**						**05571140**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		57.30	0.36	SUC 99	99	
Clostridium perfringens		30.82	0.09		99	
Fusobacterium mortiferum		11.88	0.12		1	
05571141	**Good identification**					**05571141**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05571142	**Low discrimination**					**05571142**
Actinomyces naeslundii		61.50	0.36	SUC 99		
Lactobacillus acidophilus		22.92	0.04			
Clostridium perfringens		15.57	0.03			
05571143	**Good identification**					**05571143**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05571161	**Low discrimination**					**05571161**
Actinomyces naeslundii		89.52	0.39	SUC 99		
Clostridium perfringens		10.49	0.00			
05571163	**Good identification**					**05571163**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05571322	**Low discrimination**					**05571322**
					SPO	
Parabacteroides distasonis		81.13	0.62	XYL 90	1	
Prevotella oralis		17.90	0.38		1	
Clostridium ramosum		0.70	0.14		92	
05571332	**Very good identification**					**05571332**
Parabacteroides distasonis		99.42	0.57	XYL 90		
Prevotella oralis		0.41	0.04			
Clostridium clostridioforme		0.17	0.04			
05571522	**Good identification**					**05571522**
					GRA	
Parabacteroides distasonis		95.83	0.50	XYL 90	1	
Prevotella oralis		2.44	0.10		1	
Clostridium perfringens		1.04	0.07		99	
05571532	**Good identification**					**05571532**
					GRA	
Parabacteroides distasonis		96.06	0.46	XYL 90	1	
Bifidobacterium longum subsp. longum		2.19	0.13		99	
Clostridium clostridioforme		1.75	0.10		50	
05571722	**Very good identification**					**05571722**
Parabacteroides distasonis		98.94	0.58	XYL 90		
Clostridium clostridioforme		0.52	0.14			
Prevotella oralis		0.36	0.04			
05571732	**Good identification**					**05571732**
Parabacteroides distasonis		95.73	0.54	XYL 90		
Clostridium clostridioforme		4.27	0.25			
05572000	**Good identification**					**05572000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		100.0	0.39	SUC 93		
05573000	**Low discrimination**					**05573000**
					CAT	
Propionibacterium granulosum		72.67	0.39	SUC 93	90	
Actinomyces naeslundii		25.51	0.46	CEL 1	25	
Actinomyces israelii		1.83	0.17		1	
05573001	**Good identification**					**05573001**
Actinomyces naeslundii		100.0	0.49	SUC 99		
05573002	**Low discrimination**					**05573002**
					CAT	
Actinomyces naeslundii		54.96	0.46	SUC 99	25	
Propionibacterium granulosum		39.14	0.28		90	
Actinomyces israelii		5.90	0.20		1	
05573003	**Good identification**					**05573003**
Actinomyces naeslundii		100.0	0.49	SUC 99		
05573020	**Low discrimination**					**05573020**
Actinomyces naeslundii		58.40	0.46	SUC 99		
Propionibacterium granulosum		41.60	0.28			
05573021	**Good identification**					**05573021**
Actinomyces naeslundii		100.0	0.49	SUC 99		
05573022	**Low discrimination**					**05573022**
Actinomyces naeslundii		84.89	0.46	SUC 99		
Propionibacterium granulosum		15.11	0.18			
05573023	**Good identification**					**05573023**
Actinomyces naeslundii		100.0	0.49	SUC 99		
05573041	**Good identification**					**05573041**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05573043	**Good identification**					**05573043**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05573061	**Good identification**					**05573061**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05573063	**Good identification**					**05573063**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05573100	**Low discrimination**					**05573100**
					GRA	
Actinomyces naeslundii		89.33	0.53	SUC 99	99	
Actinomyces israelii		8.99	0.27		99	
Fusobacterium mortiferum		1.68	0.12		1	
05573101	**Very good identification**					**05573101**
Actinomyces naeslundii		100.0	0.56	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05573102	**Low discrimination**						**05573102**
Actinomyces naeslundii		85.49	0.53	SUC 99			
Actinomyces israelii		12.91	0.30				
Bifidobacterium adolescentis		1.61	0.08				
05573103	**Very good identification**						**05573103**
Actinomyces naeslundii		100.0	0.56	SUC 99			
05573120	**Low discrimination**						**05573120**
					SPO		
Actinomyces naeslundii		65.22	0.53	SUC 99	1		
Clostridium septicum		18.64	0.15		99		
Clostridium ramosum		10.67	0.15		92		
05573121	**Very good identification**						**05573121**
Actinomyces naeslundii		100.0	0.56	SUC 99			
05573122	**Low discrimination**						**05573122**
					SPO	GRA	
Actinomyces naeslundii		54.14	0.53	SUC 99	1	99	
Clostridium ramosum		35.44	0.25		92	50	
Prevotella oralis		5.03	0.10		1	1	
05573123	**Very good identification**						**05573123**
Actinomyces naeslundii		100.0	0.56	SUC 99			
05573140	**Low discrimination**						**05573140**
Actinomyces naeslundii		81.85	0.41	SUC 99			
Actinomyces israelii		18.15	0.20				
05573141	**Good identification**						**05573141**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05573142	**Low discrimination**						**05573142**
Actinomyces naeslundii		75.04	0.41	SUC 99			
Actinomyces israelii		24.96	0.23				
05573143	**Good identification**						**05573143**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05573160	**Acceptable identification**						**05573160**
Actinomyces naeslundii		91.70	0.41	SUC 99			
Clostridium perfringens		8.30	0.00				
05573161	**Good identification**						**05573161**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05573162	**Good identification**						**05573162**
Actinomyces naeslundii		100.0	0.41	SUC 99			
05573163	**Good identification**						**05573163**
Actinomyces naeslundii		100.0	0.44	SUC 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
05574122	**Good identification**							**05574122**
Parabacteroides distasonis		100.0	0.36	GAL 99				
05574322	**Good identification**							**05574322**
Parabacteroides distasonis		100.0	0.44	GAL 99				
05574332	**Acceptable identification**							**05574332**
Parabacteroides distasonis		93.93	0.39	GAL 99				
Clostridium clostridioforme		6.08	0.13					
05574722	**Low discrimination**							**05574722**
Parabacteroides distasonis		83.65	0.40	GAL 99				
Clostridium clostridioforme		16.35	0.23					
05575022	**Low discrimination**							**05575022**
Parabacteroides distasonis		88.18	0.36	LAC 99				
Prevotella oralis		11.82	0.08					
05575102	**Low discrimination**							**05575102**
					COC	GRA		
Parabacteroides distasonis		40.13	0.36	NAG 99	1	1		
Actinomyces israelii		22.72	0.35		1	99		
Atopobium parvulum		18.80	0.20		99	99		
05575120	**Low discrimination**							**05575120**
					COC	GRA	SPO	
Parabacteroides distasonis		44.21	0.36	BGL 99	1	1	1	
Clostridium septicum		23.35	0.15		1	99	99	
Atopobium parvulum		20.71	0.20		99	99	1	
05575122	**Very good identification**							**05575122**
					COC	GRA		
Parabacteroides distasonis		97.69	0.70		1	1		
Prevotella oralis		2.07	0.29		1	1		
Atopobium parvulum		0.08	0.07		99	99		
05575123	**Good identification**							**05575123**
Parabacteroides distasonis		100.0	0.36	URE 1				
05575132	**Very good identification**							**05575132**
Parabacteroides distasonis		99.92	0.65					
Clostridium clostridioforme		0.08	0.07					
05575162	**Good identification**							**05575162**
Parabacteroides distasonis		100.0	0.36	SOR 1				
05575222	**Good identification**							**05575222**
Parabacteroides distasonis		98.12	0.44	LAC 99				
Prevotella oralis		1.88	0.02					
05575232	**Acceptable identification**							**05575232**
Parabacteroides distasonis		94.52	0.39	LAC 99				
Clostridium clostridioforme		5.48	0.13					
05575302	**Acceptable identification**							**05575302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		94.97	0.44	NAG 99		
Clostridium clostridioforme		2.18	0.10			
Prevotella oralis		1.50	0.01			
05575312	**Low discrimination**					**05575312**
Parabacteroides distasonis		83.75	0.39	NAG 99		
Clostridium clostridioforme		16.25	0.22			
05575320	**Acceptable identification**					**05575320**
Parabacteroides distasonis		92.53	0.44	BGL 99		
Bacteroides vulgatus		6.01	0.05			
Prevotella oralis		1.46	0.01			
05575322	**Very good identification**					**05575322**
Parabacteroides distasonis		99.68	0.78			
Prevotella oralis		0.30	0.22			
Clostridium clostridioforme		0.02	0.10			
05575323	**Good identification**					**05575323**
Parabacteroides distasonis		100.0	0.44	URE 1		
05575330	**Good identification**					**05575330**
Parabacteroides distasonis		100.0	0.39	BGL 99		
05575332	**Very good identification**					**05575332**
Parabacteroides distasonis		99.80	0.74			
Clostridium clostridioforme		0.20	0.22			
05575333	**Good identification**					**05575333**
Parabacteroides distasonis		100.0	0.39	URE 1		
05575362	**Good identification**					**05575362**
Parabacteroides distasonis		100.0	0.44	SOR 1		
05575372	**Good identification**					**05575372**
Parabacteroides distasonis		100.0	0.39	SOR 1		
05575502	**Low discrimination**					**05575502**
					GRA	
Bifidobacterium adolescentis		54.80	0.38	SUC 99	99	
Parabacteroides distasonis		17.67	0.32		1	
Actinomyces israelii		13.36	0.34		99	
05575512	**Low discrimination**					**05575512**
					GRA	
Bifidobacterium adolescentis		67.26	0.40	SUC 99	99	
Clostridium clostridioforme		19.39	0.27		50	
Parabacteroides distasonis		9.56	0.28		1	
05575522	**Very good identification**					**05575522**
					GRA	
Parabacteroides distasonis		99.37	0.66		1	
Clostridium clostridioforme		0.24	0.16		50	
Bifidobacterium longum subsp. longum		0.24	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05575532	**Very good identification**					**05575532**
					GRA	
Parabacteroides distasonis		97.27	0.62		1	
Clostridium clostridioforme		1.99	0.27		50	
Bifidobacterium longum subsp. longum		0.74	0.21		99	
05575622	**Low discrimination**					**05575622**
Parabacteroides distasonis		85.10	0.40	LAC 99		
Clostridium clostridioforme		14.90	0.22			
05575702	**Low discrimination**					**05575702**
					GRA	
Parabacteroides distasonis		62.30	0.40	NAG 99	1	
Clostridium clostridioforme		36.52	0.31		50	
Actinomyces israelii		1.18	0.15		99	
05575712	**Low discrimination**					**05575712**
Clostridium clostridioforme		83.22	0.42	SUC 99		
Parabacteroides distasonis		16.78	0.36	SAL 88		
05575722	**Very good identification**					**05575722**
Parabacteroides distasonis		98.84	0.75			
Clostridium clostridioforme		0.59	0.31			
Bacteroides thetaiotaomicron		0.47	0.19			
05575723	**Good identification**					**05575723**
Parabacteroides distasonis		100.0	0.40	URE 1		
05575730	**Good identification**					**05575730**
Parabacteroides distasonis		95.23	0.36	BGL 99		
Clostridium clostridioforme		4.77	0.08			
05575732	**Good identification**					**05575732**
Parabacteroides distasonis		94.80	0.70			
Clostridium clostridioforme		4.75	0.42			
Bacteroides thetaiotaomicron		0.32	0.12			
05575733	**Good identification**					**05575733**
Parabacteroides distasonis		95.23	0.36	URE 1		
Clostridium clostridioforme		4.77	0.08			
05575762	**Good identification**					**05575762**
Parabacteroides distasonis		100.0	0.40	SOR 1		
05575772	**Good identification**					**05575772**
Parabacteroides distasonis		95.23	0.36	SOR 1		
Clostridium clostridioforme		4.77	0.08			
05577000	**Good identification**					**05577000**
Actinomyces israelii		100.0	0.35	SUC 98		
05577002	**Good identification**					**05577002**
Actinomyces israelii		100.0	0.38	SUC 98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05577100	**Good identification**					**05577100**
Actinomyces israelii		100.0	0.45	SUC 98		
05577102	**Acceptable identification**					**05577102**
Actinomyces israelii		93.25	0.48	SUC 98		
Bifidobacterium adolescentis		6.75	0.23	SAL 92		
05577122	**Low discrimination**					**05577122**
					GRA	
Parabacteroides distasonis		83.00	0.36	MAN 1	1	
Actinomyces israelii		17.00	0.28		99	
05577140	**Good identification**					**05577140**
Actinomyces israelii		100.0	0.38	SUC 98		
05577142	**Good identification**					**05577142**
Actinomyces israelii		97.75	0.41	SUC 98		
Bifidobacterium adolescentis		2.25	0.07	SAL 92		
05577322	**Good identification**					**05577322**
Parabacteroides distasonis		100.0	0.44	MAN 1		
05577332	**Good identification**					**05577332**
Parabacteroides distasonis		100.0	0.39	MAN 1		
05577402	**Good identification**					**05577402**
Actinomyces israelii		100.0	0.37	SUC 98		
05577500	**Good identification**					**05577500**
					SPO	
Actinomyces israelii		95.21	0.44	SUC 98	1	
Clostridium butyricum		4.79	0.01	SAL 92	99	
05577502	**Low discrimination**					**05577502**
					SPO	
Actinomyces israelii		62.29	0.47	SUC 98	1	
Bifidobacterium adolescentis		36.90	0.37	SAL 92	1	
Clostridium butyricum		0.81	-.07		99	
05577512	**Low discrimination**					**05577512**
Bifidobacterium adolescentis		81.09	0.38	SUC 99		
Actinomyces israelii		16.73	0.33			
Bifidobacterium dentium		2.18	-.07			
05577540	**Good identification**					**05577540**
Actinomyces israelii		100.0	0.37	SUC 98		
05577542	**Low discrimination**					**05577542**
Actinomyces israelii		84.16	0.40	SUC 98		
Bifidobacterium adolescentis		15.85	0.21	SAL 92		
05577722	**Good identification**					**05577722**
Parabacteroides distasonis		96.49	0.40	MAN 1		
Bacteroides ovatus		3.51	0.00			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05577732	**Acceptable identification**						**05577732**
Parabacteroides distasonis		91.31	0.36	MAN 1			
Clostridium clostridioforme		4.57	0.08				
Bacteroides ovatus		4.12	-.03				
05600000	**Low discrimination**						**05600000**
					GRA	SPO	
Fusobacterium mortiferum		43.94	0.41	FRU 99	1	1	
Clostridium bifermentans		18.69	0.21		99	99	
Atopobium minutum		7.74	0.22		99	1	
05600100	**Good identification**						**05600100**
Fusobacterium mortiferum		100.0	0.47	FRU 99			
05601000	**Acceptable identification**						**05601000**
					COC	GRA	
Fusobacterium mortiferum		90.82	0.44	FRU 99	1	1	
Gemella morbillorum		5.62	0.13		99	99	
Atopobium minutum		3.56	0.14		1	99	
05601100	**Good identification**						**05601100**
					COC	GRA	
Fusobacterium mortiferum		99.12	0.50	FRU 99	1	1	
Blautia hansenii		0.89	-.03		99	99	
05610000	**Very good identification**						**05610000**
					GRA	CAT	
Fusobacterium mortiferum		95.75	0.75		1	5	
Fusobacterium varium		1.46	0.24		1	1	
Propionibacterium granulosum		1.12	0.38		99	90	
05610001	**Good identification**						**05610001**
Fusobacterium mortiferum		100.0	0.41	URE 1			
05610002	**Very good identification**						**05610002**
					GRA	CAT	
Fusobacterium mortiferum		96.51	0.58	BGL 10	1	5	
Propionibacterium granulosum		2.54	0.28		99	90	
Atopobium minutum		0.95	0.19		99	1	
05610020	**Low discrimination**						**05610020**
					GRA	CAT	
Fusobacterium mortiferum		70.27	0.41	NAG 1	1	5	
Propionibacterium granulosum		20.36	0.28		99	90	
Atopobium minutum		3.81	0.14		99	1	
05610040	**Good identification**						**05610040**
					GRA	SPO	
Fusobacterium mortiferum		97.26	0.41	SOR 1	1	1	
Clostridium difficile		2.75	-.02		99	99	
05610100	**Very good identification**						**05610100**
Fusobacterium mortiferum		100.0	0.81				
05610101	**Good identification**						**05610101**
Fusobacterium mortiferum		100.0	0.47	URE 1			
05610102	**Very good identification**						**05610102**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		100.0	0.65	BGL 10				
05610110	**Good identification**							**05610110**
					GRA	SPO		
Fusobacterium mortiferum		98.88	0.47	MLZ 1	1	1		
Clostridium difficile		1.12	-.02		99	99		
05610120	**Good identification**							**05610120**
Fusobacterium mortiferum		100.0	0.47	NAG 1				
05610140	**Good identification**							**05610140**
Fusobacterium mortiferum		100.0	0.47	SOR 1				
05610200	**Good identification**							**05610200**
					GRA	CAT		
Fusobacterium mortiferum		96.54	0.41	RHA 1	1	5		
Propionibacterium granulosum		3.46	0.13		99	90		
05610300	**Good identification**							**05610300**
Fusobacterium mortiferum		100.0	0.47	RHA 1				
05610400	**Good identification**							**05610400**
					COC	GRA		
Fusobacterium mortiferum		97.71	0.41	ARA 1	1	1		
Gemella morbillorum		2.29	0.03		99	99		
05610500	**Good identification**							**05610500**
Fusobacterium mortiferum		100.0	0.47	ARA 1				
05611000	**Very good identification**							**05611000**
					GRA	CAT		
Fusobacterium mortiferum		98.75	0.78		1	5		
Propionibacterium granulosum		0.77	0.38		99	90		
Prevotella intermedia		0.20	0.20		1	1		
05611001	**Good identification**							**05611001**
					GRA			
Fusobacterium mortiferum		98.16	0.44	URE 1	1			
Actinomyces naeslundii		1.84	0.26		99			
05611002	**Very good identification**							**05611002**
					GRA	CAT		
Fusobacterium mortiferum		96.54	0.62	BGL 10	1	5		
Propionibacterium granulosum		1.70	0.28		99	90		
Lactobacillus fermentum		1.44	0.32		99	1		
05611010	**Low discrimination**							**05611010**
					GRA	SPO	CAT	
Fusobacterium mortiferum		89.05	0.44	MLZ 1	1	1	5	
Propionibacterium granulosum		9.38	0.24		99	1	90	
Clostridium difficile		1.57	-.02		99	99	1	
05611020	**Low discrimination**							**05611020**
					GRA	CAT		
Fusobacterium mortiferum		76.01	0.44	NAG 1	1	5		
Propionibacterium granulosum		14.68	0.28		99	90		
Lactobacillus fermentum		4.16	0.24		99	1		
05611040	**Good identification**							**05611040**
Fusobacterium mortiferum		100.0	0.44	SOR 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05611100	**Very good identification**						**05611100**
					GRA		
Fusobacterium mortiferum		99.87	0.84		1		
Lactobacillus fermentum		0.12	0.35		99		
Actinomyces naeslundii		0.02	0.30		99		
05611101	**Good identification**						**05611101**
					GRA		
Fusobacterium mortiferum		97.77	0.50	URE 1	1		
Actinomyces naeslundii		2.23	0.33		99		
05611102	**Very good identification**						**05611102**
					GRA		
Fusobacterium mortiferum		98.67	0.68	BGL 10	1		
Lactobacillus fermentum		1.03	0.35		99		
Actinomyces naeslundii		0.14	0.30		99		
05611110	**Good identification**						**05611110**
Fusobacterium mortiferum		100.0	0.50	MLZ 1			
05611120	**Low discrimination**						**05611120**
					GRA	SPO	
Fusobacterium mortiferum		70.29	0.50	NAG 1	1	1	
Clostridium septicum		17.98	0.22		99	99	
Clostridium perfringens		6.41	0.21		99	84	
05611122	**Low discrimination**						**05611122**
Parabacteroides distasonis		34.80	0.38	XYL 90			
Prevotella oralis		34.23	0.25				
Fusobacterium mortiferum		15.51	0.34				
05611140	**Good identification**						**05611140**
Fusobacterium mortiferum		100.0	0.50	SOR 1			
05611200	**Good identification**						**05611200**
					GRA	CAT	
Fusobacterium mortiferum		97.67	0.44	RHA 1	1	5	
Propionibacterium granulosum		2.33	0.13		99	90	
05611300	**Good identification**						**05611300**
Fusobacterium mortiferum		100.0	0.50	RHA 1			
05611322	**Low discrimination**						**05611322**
Parabacteroides distasonis		87.68	0.46	XYL 90			
Prevotella oralis		12.32	0.19				
05611332	**Good identification**						**05611332**
Parabacteroides distasonis		100.0	0.42	XYL 90			
05611400	**Low discrimination**						**05611400**
					GRA		
Fusobacterium mortiferum		85.89	0.44	ARA 1	1		
Lactobacillus fermentum		14.11	0.32		99		
05611500	**Low discrimination**						**05611500**
					GRA		
Fusobacterium mortiferum		89.70	0.50	ARA 1	1		
Lactobacillus fermentum		10.30	0.35		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05611502	**Low discrimination**							**05611502**
					GRA			
Lactobacillus fermentum		50.83	0.35	SUC 87	99			
Fusobacterium mortiferum		49.17	0.34	SAL 1	1			
05611722	**Good identification**							**05611722**
Parabacteroides distasonis		100.0	0.43	XYL 90				
05611732	**Good identification**							**05611732**
Parabacteroides distasonis		100.0	0.38	XYL 90				
05612000	**Low discrimination**							**05612000**
					GRA	SPO		
Clostridium difficile		54.25	0.37	RAF 1	99	99		
Eubacterium limosum		24.05	0.29		99	1		
Fusobacterium mortiferum		11.24	0.41		1	1		
05612010	**Good identification**							**05612010**
					SPO	CAT		
Clostridium difficile		99.41	0.48	RAF 1	99	1		
Propionibacterium granulosum		0.50	0.22		1	90		
Eubacterium limosum		0.09	-.06		1	1		
05612100	**Good identification**							**05612100**
					GRA	SPO		
Fusobacterium mortiferum		97.07	0.47	MAN 1	1	1		
Clostridium difficile		2.03	0.02		99	99		
Eubacterium limosum		0.90	-.06		99	1		
05613000	**Low discrimination**							**05613000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		48.22	0.44	MAN 1	1	1	5	
Propionibacterium granulosum		28.10	0.36		99	1	90	
Clostridium innocuum		20.26	0.25		99	99	1	
05613100	**Good identification**							**05613100**
					GRA			
Fusobacterium mortiferum		97.13	0.50	MAN 1	1			
Actinomyces naeslundii		2.87	0.35		99			
05613101	**Low discrimination**							**05613101**
					GRA			
Actinomyces naeslundii		81.46	0.38	SUC 99	99			
Fusobacterium mortiferum		18.55	0.16	MLT 94	1			
05613103	**Good identification**							**05613103**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05613120	**Low discrimination**							**05613120**
					GRA			
Actinomyces naeslundii		74.54	0.35	SUC 99	99			
Fusobacterium mortiferum		25.46	0.16	MLT 94	1			
05613121	**Good identification**							**05613121**
Actinomyces naeslundii		100.0	0.38	SUC 99				
05613122	**Good identification**							**05613122**
Actinomyces naeslundii		100.0	0.35	SUC 99				
05613123	**Good identification**							**05613123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	SUC 99			
05614000	**Good identification**						**05614000**
					GRA	SPO	
Fusobacterium mortiferum		98.49	0.56	XYL 7	1	1	
Fusobacterium varium		0.83	0.01		1	1	
Clostridium difficile		0.37	-.02		99	99	
05614002	**Good identification**						**05614002**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
05614100	**Very good identification**						**05614100**
Fusobacterium mortiferum		100.0	0.62	XYL 7			
05614102	**Good identification**						**05614102**
Fusobacterium mortiferum		100.0	0.46	XYL 7			
05615000	**Very good identification**						**05615000**
					GRA		
Fusobacterium mortiferum		98.72	0.59	XYL 7	1		
Lactobacillus fermentum		1.28	0.28		99		
05615002	**Low discrimination**						**05615002**
					GRA		
Fusobacterium mortiferum		89.56	0.42	XYL 7	1		
Lactobacillus fermentum		10.44	0.28	BGL 10	99		
05615100	**Very good identification**						**05615100**
					GRA		
Fusobacterium mortiferum		99.10	0.65	XYL 7	1		
Lactobacillus fermentum		0.90	0.31		99		
05615102	**Acceptable identification**						**05615102**
					GRA		
Fusobacterium mortiferum		90.18	0.49	XYL 7	1		
Lactobacillus fermentum		7.35	0.31	BGL 10	99		
Parabacteroides distasonis		2.47	0.20		1		
05615122	**Good identification**						**05615122**
					GRA		
Parabacteroides distasonis		97.03	0.55		1		
Prevotella oralis		1.31	0.10		1		
Lactobacillus fermentum		0.97	0.23		99		
05615132	**Good identification**						**05615132**
Parabacteroides distasonis		100.0	0.50				
05615322	**Very good identification**						**05615322**
Parabacteroides distasonis		99.81	0.63				
Prevotella oralis		0.19	0.04				
05615332	**Very good identification**						**05615332**
Parabacteroides distasonis		100.0	0.58				
05615522	**Good identification**						**05615522**
					GRA		
Parabacteroides distasonis		98.39	0.51		1		
Lactobacillus fermentum		1.61	0.23		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05615532	**Good identification**							**05615532**
Parabacteroides distasonis		100.0	0.46					
05615722	**Very good identification**							**05615722**
Parabacteroides distasonis		99.87	0.59					
Bacteroides vulgatus		0.13	-.09					
05615732	**Good identification**							**05615732**
Parabacteroides distasonis		100.0	0.55					
05620100	**Good identification**							**05620100**
Fusobacterium mortiferum		100.0	0.38	FRU 99				
05621100	**Good identification**							**05621100**
Fusobacterium mortiferum		100.0	0.41	FRU 99				
05625322	**Good identification**							**05625322**
Parabacteroides distasonis		100.0	0.40	FRU 99				
05625332	**Good identification**							**05625332**
Parabacteroides distasonis		100.0	0.35	FRU 99				
05625722	**Good identification**							**05625722**
Parabacteroides distasonis		100.0	0.36	FRU 99				
05630000	**Low discrimination**							**05630000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		89.37	0.66		1	1	5	
Propionibacterium granulosum		10.51	0.47		99	1	90	
Clostridium difficile		0.12	0.01		99	99	1	
05630002	**Low discrimination**							**05630002**
					GRA	CAT		
Fusobacterium mortiferum		79.08	0.50	BGL 10	1	5		
Propionibacterium granulosum		20.92	0.36		99	90		
05630020	**Low discrimination**							**05630020**
					GRA	CAT		
Propionibacterium granulosum		74.42	0.36	SUC 93	99	90		
Fusobacterium mortiferum		25.58	0.32	SAL 1	1	5		
05630100	**Very good identification**							**05630100**
					GRA	CAT		
Fusobacterium mortiferum		99.95	0.72		1	5		
Propionibacterium granulosum		0.05	0.13		99	90		
05630101	**Good identification**							**05630101**
Fusobacterium mortiferum		100.0	0.38	URE 1				
05630102	**Good identification**							**05630102**
Fusobacterium mortiferum		100.0	0.56	BGL 10				
05630110	**Good identification**							**05630110**
Fusobacterium mortiferum		100.0	0.38	MLZ 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05630120	**Good identification**						**05630120**
					GRA	SPO	
Fusobacterium mortiferum		96.25	0.38	NAG 1	1	1	
Clostridium septicum		3.75	-.04		99	99	
05630140	**Good identification**						**05630140**
Fusobacterium mortiferum		100.0	0.38	SOR 1			
05630300	**Good identification**						**05630300**
Fusobacterium mortiferum		100.0	0.38	RHA 1			
05630500	**Good identification**						**05630500**
Fusobacterium mortiferum		100.0	0.38	ARA 1			
05631000	**Good identification**						**05631000**
					GRA	CAT	
Fusobacterium mortiferum		92.60	0.69		1	5	
Propionibacterium granulosum		7.26	0.47		99	90	
Actinomyces naeslundii		0.08	0.28		99	25	
05631002	**Low discrimination**						**05631002**
					GRA	CAT	
Fusobacterium mortiferum		84.14	0.53	BGL 10	1	5	
Propionibacterium granulosum		14.84	0.36		99	90	
Actinomyces naeslundii		0.62	0.28		99	25	
05631020	**Low discrimination**						**05631020**
					GRA	CAT	
Propionibacterium granulosum		59.47	0.36	SUC 93	99	90	
Fusobacterium mortiferum		30.66	0.35	SAL 1	1	5	
Clostridium perfringens		5.53	0.11		99	1	
05631100	**Very good identification**						**05631100**
					GRA	CAT	
Fusobacterium mortiferum		99.82	0.75		1	5	
Actinomyces naeslundii		0.10	0.35		99	25	
Propionibacterium granulosum		0.03	0.13		99	90	
05631101	**Low discrimination**						**05631101**
					GRA		
Fusobacterium mortiferum		87.07	0.41	URE 1	1		
Actinomyces naeslundii		12.93	0.38		99		
05631102	**Very good identification**						**05631102**
					GRA		
Fusobacterium mortiferum		98.59	0.59	BGL 10	1		
Actinomyces naeslundii		0.89	0.35		99		
Parabacteroides distasonis		0.34	0.15		1		
05631103	**Low discrimination**						**05631103**
					GRA		
Actinomyces naeslundii		57.21	0.38	SUC 99	99		
Fusobacterium mortiferum		42.80	0.25	MLT 94	1		
05631110	**Good identification**						**05631110**
Fusobacterium mortiferum		100.0	0.41	MLZ 1			
05631121	**Low discrimination**						**05631121**
					SPO		
Actinomyces naeslundii		85.13	0.38	SUC 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium septicum		14.87	-.04	MLT 94	99			
05631122	**Low discrimination**							**05631122**
					GRA			
Parabacteroides distasonis		86.35	0.50	XYL 90	1			
Clostridium perfringens		6.05	0.20		99			
Fusobacterium mortiferum		2.52	0.25		1			
05631123	**Low discrimination**							**05631123**
					GRA			
Actinomyces naeslundii		79.45	0.38	SUC 99	99			
Parabacteroides distasonis		20.55	0.15	MLT 94	1			
05631132	**Good identification**							**05631132**
Parabacteroides distasonis		100.0	0.45	XYL 90				
05631140	**Good identification**							**05631140**
					GRA			
Fusobacterium mortiferum		98.28	0.41	SOR 1	1			
Actinomyces naeslundii		1.72	0.22		99			
05631300	**Good identification**							**05631300**
Fusobacterium mortiferum		100.0	0.41	RHA 1				
05631322	**Very good identification**							**05631322**
Parabacteroides distasonis		99.80	0.58	XYL 90				
Prevotella oralis		0.20	-.01					
05631332	**Good identification**							**05631332**
Parabacteroides distasonis		100.0	0.53	XYL 90				
05631500	**Good identification**							**05631500**
Fusobacterium mortiferum		100.0	0.41	ARA 1				
05631522	**Good identification**							**05631522**
Parabacteroides distasonis		100.0	0.46	XYL 90				
05631532	**Good identification**							**05631532**
Parabacteroides distasonis		100.0	0.41	XYL 90				
05631722	**Good identification**							**05631722**
Parabacteroides distasonis		100.0	0.54	XYL 90				
05631732	**Good identification**							**05631732**
Parabacteroides distasonis		100.0	0.49	XYL 90				
05632000	**Low discrimination**							**05632000**
					GRA	SPO	CAT	
Propionibacterium granulosum		79.33	0.45	SUC 93	99	1	90	
Clostridium difficile		10.99	0.17	SAL 1	99	99	1	
Fusobacterium mortiferum		9.04	0.32		1	1	5	
05632100	**Good identification**							**05632100**
					GRA	CAT		
Fusobacterium mortiferum		96.34	0.38	MAN 1	1	5		
Propionibacterium granulosum		3.66	0.10		99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
05633000	**Low discrimination**							**05633000**
					GRA	SPO	CAT	
Propionibacterium granulosum		77.99	0.45	SUC 93	99	1	90	
Fusobacterium mortiferum		13.33	0.35	SAL 1	1	1	5	
Clostridium innocuum		6.59	0.17		99	99	1	
05633001	**Low discrimination**							**05633001**
Actinomyces naeslundii		79.99	0.36	SUC 99				
Propionibacterium granulosum		20.01	0.10	MLT 94				
05633003	**Good identification**							**05633003**
Actinomyces naeslundii		100.0	0.36	SUC 99				
05633021	**Good identification**							**05633021**
Actinomyces naeslundii		100.0	0.36	SUC 99				
05633023	**Good identification**							**05633023**
Actinomyces naeslundii		100.0	0.36	SUC 99				
05633100	**Low discrimination**							**05633100**
					GRA	CAT		
Fusobacterium mortiferum		82.13	0.41	MAN 1	1	5		
Actinomyces naeslundii		15.79	0.40		99	25		
Propionibacterium granulosum		2.08	0.10		99	90		
05633101	**Good identification**							**05633101**
Actinomyces naeslundii		100.0	0.43	SUC 99				
05633102	**Low discrimination**							**05633102**
					GRA			
Actinomyces naeslundii		63.37	0.40	SUC 99	99			
Fusobacterium mortiferum		36.63	0.25	MLT 94	1			
05633103	**Good identification**							**05633103**
Actinomyces naeslundii		100.0	0.43	SUC 99				
05633120	**Low discrimination**							**05633120**
					SPO			
Actinomyces naeslundii		88.10	0.40	SUC 99	1			
Clostridium septicum		11.90	-.04	MLT 94	99			
05633121	**Good identification**							**05633121**
Actinomyces naeslundii		100.0	0.43	SUC 99				
05633122	**Low discrimination**							**05633122**
					GRA			
Actinomyces naeslundii		83.34	0.40	SUC 99	99			
Parabacteroides distasonis		16.66	0.15	MLT 94	1			
05633123	**Good identification**							**05633123**
Actinomyces naeslundii		100.0	0.43	SUC 99				
05634000	**Good identification**							**05634000**
					GRA	CAT		
Fusobacterium mortiferum		98.45	0.47	XYL 7	1	5		
Propionibacterium granulosum		1.55	0.13		99	90		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
05634100	**Good identification**						**05634100**
Fusobacterium mortiferum		100.0	0.53	XYL 7			
05634102	**Good identification**						**05634102**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
05634322	**Good identification**						**05634322**
Parabacteroides distasonis		100.0	0.40	GAL 99			
05634332	**Good identification**						**05634332**
Parabacteroides distasonis		100.0	0.35	GAL 99			
05634722	**Good identification**						**05634722**
Parabacteroides distasonis		100.0	0.36	GAL 99			
05635000	**Good identification**						**05635000**
					GRA	CAT	
Fusobacterium mortiferum		98.96	0.50	XYL 7	1	5	
Propionibacterium granulosum		1.04	0.13		99	90	
05635100	**Good identification**						**05635100**
Fusobacterium mortiferum		100.0	0.56	XYL 7			
05635102	**Low discrimination**						**05635102**
Fusobacterium mortiferum		70.54	0.40	XYL 7			
Parabacteroides distasonis		29.46	0.32	BGL 10			
05635122	**Very good identification**						**05635122**
Parabacteroides distasonis		100.0	0.66				
05635132	**Very good identification**						**05635132**
Parabacteroides distasonis		100.0	0.61				
05635222	**Good identification**						**05635222**
Parabacteroides distasonis		100.0	0.40	LAC 99			
05635232	**Good identification**						**05635232**
Parabacteroides distasonis		100.0	0.35	LAC 99			
05635302	**Good identification**						**05635302**
Parabacteroides distasonis		100.0	0.40	NAG 99			
05635312	**Good identification**						**05635312**
Parabacteroides distasonis		100.0	0.35	NAG 99			
05635320	**Good identification**						**05635320**
Parabacteroides distasonis		100.0	0.40	BGL 99			
05635322	**Very good identification**						**05635322**
Parabacteroides distasonis		100.0	0.74				
05635323	**Good identification**						**05635323**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		100.0	0.40	URE 1		
05635330	**Good identification**					**05635330**
Parabacteroides distasonis		100.0	0.35	BGL 99		
05635332	**Very good identification**					**05635332**
Parabacteroides distasonis		100.0	0.69			
05635333	**Good identification**					**05635333**
Parabacteroides distasonis		100.0	0.35	URE 1		
05635362	**Good identification**					**05635362**
Parabacteroides distasonis		100.0	0.40	SOR 1		
05635372	**Good identification**					**05635372**
Parabacteroides distasonis		100.0	0.35	SOR 1		
05635522	**Very good identification**					**05635522**
Parabacteroides distasonis		100.0	0.62			
05635532	**Very good identification**					**05635532**
Parabacteroides distasonis		100.0	0.58			
05635622	**Good identification**					**05635622**
Parabacteroides distasonis		100.0	0.36	LAC 99		
05635702	**Good identification**					**05635702**
Parabacteroides distasonis		100.0	0.36	NAG 99		
05635720	**Low discrimination**					**05635720**
Parabacteroides distasonis		89.37	0.36	BGL 99		
Bacteroides vulgatus		10.63	0.02			
05635722	**Very good identification**					**05635722**
Parabacteroides distasonis		100.0	0.70			
05635723	**Good identification**					**05635723**
Parabacteroides distasonis		100.0	0.36	URE 1		
05635732	**Very good identification**					**05635732**
Parabacteroides distasonis		99.81	0.66			
Clostridium clostridioforme		0.20	0.14			
05635762	**Good identification**					**05635762**
Parabacteroides distasonis		100.0	0.36	SOR 1		
05637322	**Good identification**					**05637322**
Parabacteroides distasonis		100.0	0.40	MAN 1		
05637332	**Good identification**					**05637332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.35	MAN 1			
05637722	**Good identification**						**05637722**
Parabacteroides distasonis		100.0	0.36	MAN 1			
05640100	**Good identification**						**05640100**
Fusobacterium mortiferum		100.0	0.39	FRU 99			
05641000	**Acceptable identification**						**05641000**
					GRA	SPO	
Fusobacterium mortiferum		93.78	0.36	FRU 99	1	1	
Clostridium innocuum		6.22	0.03		99	99	
05641100	**Good identification**						**05641100**
Fusobacterium mortiferum		100.0	0.42	FRU 99			
05643000	**Good identification**						**05643000**
Clostridium innocuum		100.0	0.37	RAF 1			
05643002	**Good identification**						**05643002**
Clostridium innocuum		100.0	0.37	RAF 1			
05650000	**Very good identification**						**05650000**
					GRA	SPO	
Fusobacterium mortiferum		99.81	0.67		1	1	
Clostridium innocuum		0.10	0.03		99	99	
Clostridium difficile		0.09	-.02		99	99	
05650002	**Good identification**						**05650002**
					GRA	SPO	
Fusobacterium mortiferum		99.10	0.50	BGL 10	1	1	
Clostridium innocuum		0.90	0.03		99	99	
05650100	**Very good identification**						**05650100**
Fusobacterium mortiferum		100.0	0.73				
05650101	**Good identification**						**05650101**
Fusobacterium mortiferum		100.0	0.39	URE 1			
05650102	**Very good identification**						**05650102**
Fusobacterium mortiferum		100.0	0.57	BGL 10			
05650110	**Good identification**						**05650110**
Fusobacterium mortiferum		100.0	0.39	MLZ 1			
05650120	**Good identification**						**05650120**
					GRA	SPO	
Fusobacterium mortiferum		96.45	0.39	NAG 1	1	1	
Clostridium septicum		3.55	-.03		99	99	
05650140	**Good identification**						**05650140**
Fusobacterium mortiferum		100.0	0.39	SOR 1			
05650300	**Good identification**						**05650300**
Fusobacterium mortiferum		100.0	0.39	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05650500	**Good identification**						**05650500**
Fusobacterium mortiferum		100.0	0.39	ARA 1			
05651000	**Very good identification**						**05651000**
					GRA	SPO	
Fusobacterium mortiferum		98.10	0.70		1	1	
Clostridium innocuum		1.25	0.25		99	99	
Lactobacillus jensenii		0.44	0.19		99	1	
05651001	**Acceptable identification**						**05651001**
					GRA		
Fusobacterium mortiferum		93.07	0.36	URE 1	1		
Actinomyces naeslundii		6.93	0.28		99		
05651002	**Low discrimination**						**05651002**
					GRA	SPO	
Fusobacterium mortiferum		85.40	0.53	BGL 10	1	1	
Clostridium innocuum		9.79	0.25		99	99	
Lactobacillus jensenii		2.29	0.16		99	1	
05651010	**Low discrimination**						**05651010**
					GRA	SPO	
Fusobacterium mortiferum		83.33	0.36	MLZ 1	1	1	
Lactobacillus jensenii		12.29	0.11		99	1	
Clostridium innocuum		4.38	0.01		99	99	
05651020	**Low discrimination**						**05651020**
					GRA	SPO	
Fusobacterium mortiferum		79.37	0.36	NAG 1	1	1	
Lactobacillus jensenii		8.78	0.09		99	1	
Clostridium septicum		4.55	-.03		99	99	
05651040	**Good identification**						**05651040**
Fusobacterium mortiferum		100.0	0.36	SOR 1			
05651100	**Very good identification**						**05651100**
					GRA		
Fusobacterium mortiferum		99.81	0.76		1		
Lactobacillus catenaformis		0.11	0.15		99		
Actinomyces naeslundii		0.06	0.33		99		
05651101	**Acceptable identification**						**05651101**
					GRA		
Fusobacterium mortiferum		91.67	0.42	URE 1	1		
Actinomyces naeslundii		8.33	0.36		99		
05651102	**Good identification**						**05651102**
					GRA		
Fusobacterium mortiferum		92.84	0.60	BGL 10	1		
Lactobacillus acidophilus		4.41	0.24		99		
Prevotella oralis		1.16	0.13		1		
05651110	**Good identification**						**05651110**
Fusobacterium mortiferum		100.0	0.42	MLZ 1			
05651121	**Low discrimination**						**05651121**
					SPO		
Actinomyces naeslundii		78.74	0.36	SUC 99	1		
Clostridium septicum		21.26	-.03	MLT 94	99		
05651123	**Low discrimination**						**05651123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		77.52	0.36	SUC 99	99		
Prevotella oralis		22.48	0.01	MLT 94	1		
05651140	**Good identification**						**05651140**
Fusobacterium mortiferum		100.0	0.42	SOR 1			
05651200	**Good identification**						**05651200**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
05651300	**Good identification**						**05651300**
Fusobacterium mortiferum		100.0	0.42	RHA 1			
05651322	**Low discrimination**						**05651322**
					GRA		
Parabacteroides distasonis		51.08	0.42	XYL 90	1		
Prevotella oralis		47.26	0.29		1		
Lactobacillus acidophilus		1.66	0.05		99		
05651332	**Good identification**						**05651332**
Parabacteroides distasonis		100.0	0.37	XYL 90			
05651400	**Low discrimination**						**05651400**
					GRA	SPO	
Fusobacterium mortiferum		81.80	0.36	ARA 1	1	1	
Clostridium innocuum		18.20	0.12		99	99	
05651500	**Good identification**						**05651500**
Fusobacterium mortiferum		100.0	0.42	ARA 1			
05651722	**Good identification**						**05651722**
Parabacteroides distasonis		100.0	0.38	XYL 90			
05652000	**Low discrimination**						**05652000**
					GRA	SPO	
Clostridium innocuum		84.36	0.37	RAF 1	99	99	
Fusobacterium mortiferum		8.56	0.33	GAL 95	1	1	
Clostridium difficile		6.52	0.15		99	99	
05652002	**Good identification**						**05652002**
					GRA	SPO	
Clostridium innocuum		98.89	0.37	RAF 1	99	99	
Fusobacterium mortiferum		1.12	0.16	GAL 95	1	1	
05652100	**Good identification**						**05652100**
					GRA	SPO	
Fusobacterium mortiferum		95.91	0.39	MAN 1	1	1	
Clostridium innocuum		4.09	0.03		99	99	
05653000	**Very good identification**						**05653000**
					GRA	SPO	
Clostridium innocuum		99.13	0.59	RAF 1	99	99	
Fusobacterium mortiferum		0.79	0.36		1	1	
Actinomyces naeslundii		0.08	0.30		99	1	
05653002	**Very good identification**						**05653002**
					GRA	SPO	
Clostridium innocuum		99.83	0.59	RAF 1	99	99	
Fusobacterium mortiferum		0.09	0.19		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.08	0.30		99	1	
05653010	**Good identification**						**05653010**
Clostridium innocuum		100.0	0.35	RAF 1			
05653012	**Good identification**						**05653012**
Clostridium innocuum		100.0	0.35	RAF 1			
05653100	**Low discrimination**						**05653100**
					GRA	SPO	
Fusobacterium mortiferum		60.32	0.42	MAN 1	1	1	
Clostridium innocuum		32.58	0.25		99	99	
Actinomyces naeslundii		7.10	0.37		99	1	
05653101	**Good identification**						**05653101**
Actinomyces naeslundii		100.0	0.41	SUC 99			
05653103	**Good identification**						**05653103**
Actinomyces naeslundii		100.0	0.41	SUC 99			
05653120	**Low discrimination**						**05653120**
					SPO		
Actinomyces naeslundii		82.74	0.37	SUC 99	1		
Clostridium septicum		17.26	-.03	MLT 94	99		
05653121	**Good identification**						**05653121**
Actinomyces naeslundii		100.0	0.41	SUC 99			
05653122	**Low discrimination**						**05653122**
					SPO		
Actinomyces naeslundii		55.06	0.37	SUC 99	1		
Clostridium ramosum		19.74	0.05	MLT 94	92		
Lactobacillus acidophilus		12.87	0.02		1		
05653123	**Good identification**						**05653123**
Actinomyces naeslundii		100.0	0.41	SUC 99			
05653400	**Good identification**						**05653400**
Clostridium innocuum		100.0	0.46	RAF 1			
05653402	**Good identification**						**05653402**
Clostridium innocuum		100.0	0.46	RAF 1			
05654000	**Good identification**						**05654000**
Fusobacterium mortiferum		100.0	0.47	XYL 7			
05654100	**Good identification**						**05654100**
Fusobacterium mortiferum		100.0	0.54	XYL 7			
05654102	**Good identification**						**05654102**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
05655000	**Good identification**						**05655000**
					GRA	SPO	
Fusobacterium mortiferum		99.12	0.50	XYL 7	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		0.88	0.03		99	99	
05655100	**Very good identification**						**05655100**
Fusobacterium mortiferum		100.0	0.57	XYL 7			
05655102	**Acceptable identification**						**05655102**
Fusobacterium mortiferum		93.92	0.40	XYL 7			
Parabacteroides distasonis		4.15	0.16	BGL 10			
Prevotella oralis		1.93	-.03				
05655122	**Acceptable identification**						**05655122**
Parabacteroides distasonis		91.83	0.50				
Prevotella oralis		8.17	0.19				
05655132	**Good identification**						**05655132**
Parabacteroides distasonis		100.0	0.45				
05655322	**Very good identification**						**05655322**
Parabacteroides distasonis		98.75	0.58				
Prevotella oralis		1.26	0.13				
05655332	**Good identification**						**05655332**
Parabacteroides distasonis		100.0	0.54				
05655522	**Good identification**						**05655522**
Parabacteroides distasonis		100.0	0.46				
05655532	**Good identification**						**05655532**
Parabacteroides distasonis		100.0	0.42				
05655722	**Good identification**						**05655722**
Parabacteroides distasonis		100.0	0.55				
05655732	**Good identification**						**05655732**
Parabacteroides distasonis		99.25	0.50				
Clostridium clostridioforme		0.75	0.08				
05657000	**Good identification**						**05657000**
					GRA	SPO	
Clostridium innocuum		98.87	0.37	RAF 1	99	99	
Fusobacterium mortiferum		1.13	0.16	XYL 5	1	1	
05657002	**Good identification**						**05657002**
Clostridium innocuum		100.0	0.37	RAF 1			
05665322	**Good identification**						**05665322**
Parabacteroides distasonis		100.0	0.35	FRU 99			
05670000	**Very good identification**						**05670000**
					GRA	CAT	
Fusobacterium mortiferum		99.65	0.58		1	5	
Propionibacterium granulosum		0.36	0.13		99	90	
05670002	**Good identification**						**05670002**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.41	BGL 10			
05670100	**Very good identification**						**05670100**
Fusobacterium mortiferum		100.0	0.64				
05670102	**Good identification**						**05670102**
					GRA		
Fusobacterium mortiferum		99.20	0.48	BGL 10	1		
Lactobacillus acidophilus		0.80	-.01		99		
05671000	**Good identification**						**05671000**
					GRA	SPO	
Fusobacterium mortiferum		93.85	0.61		1	1	
Lactobacillus jensenii		4.21	0.28		99	1	
Clostridium innocuum		1.41	0.17		99	99	
05671002	**Low discrimination**						**05671002**
					GRA	SPO	
Fusobacterium mortiferum		64.88	0.44	BGL 10	1	1	
Lactobacillus jensenii		17.48	0.25		99	1	
Clostridium innocuum		8.74	0.17		99	99	
05671100	**Very good identification**						**05671100**
					GRA	SPO	
Fusobacterium mortiferum		99.30	0.67		1	1	
Actinomyces naeslundii		0.40	0.37		99	1	
Clostridium septicum		0.25	0.05		99	99	
05671102	**Low discrimination**						**05671102**
					GRA		
Fusobacterium mortiferum		63.56	0.51	BGL 10	1		
Lactobacillus acidophilus		33.81	0.33		99		
Actinomyces naeslundii		2.28	0.37		99		
05671103	**Low discrimination**						**05671103**
					GRA		
Actinomyces naeslundii		77.63	0.41	SUC 99	99		
Fusobacterium mortiferum		14.60	0.16	MLT 94	1		
Lactobacillus acidophilus		7.77	-.01		99		
05671120	**Acceptable identification**						**05671120**
					GRA	SPO	
Clostridium septicum		93.55	0.39	MLT 94	99	99	
Fusobacterium mortiferum		3.83	0.33	RAF 1	1	1	
Actinomyces naeslundii		1.51	0.37		99	1	
05671121	**Low discrimination**						**05671121**
					SPO		
Actinomyces naeslundii		70.56	0.41	SUC 99	1		
Clostridium septicum		29.45	0.05	MLT 94	99		
05671122	**Low discrimination**						**05671122**
					GRA		
Parabacteroides distasonis		44.05	0.45	XYL 90	1		
Lactobacillus acidophilus		41.96	0.33		99		
Prevotella oralis		4.00	0.14		1		
05671123	**Low discrimination**						**05671123**
					GRA		
Actinomyces naeslundii		82.98	0.41	SUC 99	99		
Parabacteroides distasonis		8.72	0.11	MLT 94	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		8.31	-.01		99		
05671132	**Good identification**						**05671132**
					GRA		
Parabacteroides distasonis		98.24	0.40	XYL 90	1		
Lactobacillus acidophilus		1.76	-.01		99		
05671322	**Acceptable identification**						**05671322**
					SPO	GRA	
Parabacteroides distasonis		93.90	0.53	XYL 90	1	1	
Clostridium ramosum		2.64	0.14		92	50	
Lactobacillus acidophilus		2.24	0.14		1	99	
05671332	**Good identification**						**05671332**
Parabacteroides distasonis		100.0	0.48	XYL 90			
05671522	**Good identification**						**05671522**
					GRA		
Parabacteroides distasonis		98.45	0.41	XYL 90	1		
Lactobacillus acidophilus		1.55	-.01		99		
05671532	**Good identification**						**05671532**
Parabacteroides distasonis		100.0	0.37	XYL 90			
05671722	**Good identification**						**05671722**
Parabacteroides distasonis		100.0	0.49	XYL 90			
05671732	**Good identification**						**05671732**
Parabacteroides distasonis		98.93	0.45	XYL 90			
Clostridium clostridioforme		1.07	0.06				
05673000	**Good identification**						**05673000**
					GRA	SPO	
Clostridium innocuum		98.79	0.51	RAF 1	99	99	
Fusobacterium mortiferum		0.67	0.27		1	1	
Actinomyces naeslundii		0.42	0.35		99	1	
05673002	**Good identification**						**05673002**
					SPO		
Clostridium innocuum		99.58	0.51	RAF 1	99		
Actinomyces naeslundii		0.43	0.35		1		
05673021	**Good identification**						**05673021**
Actinomyces naeslundii		100.0	0.38	SUC 99			
05673023	**Good identification**						**05673023**
Actinomyces naeslundii		100.0	0.38	SUC 99			
05673101	**Good identification**						**05673101**
Actinomyces naeslundii		100.0	0.45	SUC 99			
05673102	**Low discrimination**						**05673102**
					SPO		
Actinomyces naeslundii		40.70	0.42	SUC 99	1		
Clostridium innocuum		33.83	0.17	MLT 94	99		
Lactobacillus acidophilus		16.40	0.11		1		
05673103	**Good identification**						**05673103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.45	SUC 99			
05673120	**Low discrimination**						**05673120**
					SPO		
Actinomyces naeslundii		48.89	0.42	SUC 99	1		
Clostridium ramosum		35.35	0.15	MLT 94	92		
Clostridium septicum		15.77	0.05		99		
05673121	**Good identification**						**05673121**
Actinomyces naeslundii		100.0	0.45	SUC 99			
05673123	**Good identification**						**05673123**
Actinomyces naeslundii		100.0	0.45	SUC 99			
05673400	**Good identification**						**05673400**
Clostridium innocuum		100.0	0.38	RAF 1			
05673402	**Good identification**						**05673402**
Clostridium innocuum		100.0	0.38	RAF 1			
05674000	**Good identification**						**05674000**
Fusobacterium mortiferum		100.0	0.38	XYL 7			
05674100	**Good identification**						**05674100**
Fusobacterium mortiferum		100.0	0.45	XYL 7			
05674322	**Good identification**						**05674322**
Parabacteroides distasonis		100.0	0.35	GAL 99			
05675000	**Good identification**						**05675000**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
05675100	**Good identification**						**05675100**
					GRA	SPO	
Fusobacterium mortiferum		96.06	0.48	XYL 7	1	1	
Clostridium butyricum		2.07	0.01		99	99	
Actinomyces israelii		1.03	0.16		99	1	
05675122	**Very good identification**						**05675122**
					GRA		
Parabacteroides distasonis		99.77	0.61		1		
Prevotella oralis		0.12	-.01		1		
Lactobacillus acidophilus		0.11	-.01		99		
05675132	**Very good identification**						**05675132**
Parabacteroides distasonis		100.0	0.57				
05675222	**Good identification**						**05675222**
Parabacteroides distasonis		100.0	0.35	LAC 99			
05675302	**Good identification**						**05675302**
Parabacteroides distasonis		100.0	0.35	NAG 99			
05675320	**Good identification**						**05675320**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		100.0	0.35	BGL 99		
05675322	**Very good identification**					**05675322**
Parabacteroides distasonis		100.0	0.69			
05675323	**Good identification**					**05675323**
Parabacteroides distasonis		100.0	0.35	URE 1		
05675332	**Very good identification**					**05675332**
Parabacteroides distasonis		100.0	0.65			
05675362	**Good identification**					**05675362**
Parabacteroides distasonis		100.0	0.35	SOR 1		
05675522	**Very good identification**					**05675522**
Parabacteroides distasonis		100.0	0.58			
05675532	**Good identification**					**05675532**
Parabacteroides distasonis		99.51	0.53			
Clostridium clostridioforme		0.49	0.08			
05675722	**Very good identification**					**05675722**
Parabacteroides distasonis		99.86	0.66			
Clostridium clostridioforme		0.14	0.11			
05675732	**Very good identification**					**05675732**
Parabacteroides distasonis		98.80	0.61			
Clostridium clostridioforme		1.20	0.23			
05677322	**Good identification**					**05677322**
Parabacteroides distasonis		100.0	0.35	MAN 1		
05703101	**Good identification**					**05703101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05703103	**Good identification**					**05703103**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05703121	**Good identification**					**05703121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05703123	**Good identification**					**05703123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05705322	**Acceptable identification**					**05705322**
Parabacteroides distasonis		94.89	0.38	FRU 99		
Prevotella oralis		5.11	0.04			
05710000	**Low discrimination**					**05710000**
Prevotella intermedia		62.15	0.55	SAL 1		
Fusobacterium mortiferum		25.99	0.59			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella disiens		8.11	0.32			
05710002	**Low discrimination**					**05710002**
					GRA	
Fusobacterium mortiferum		65.76	0.43	MLT 11	1	
Prevotella intermedia		14.30	0.20	BGL 10	1	
Actinomyces viscosus		11.79	0.18		99	
05710100	**Very good identification**					**05710100**
Fusobacterium mortiferum		98.72	0.66	MLT 11		
Prevotella intermedia		1.02	0.20			
Prevotella disiens		0.13	-.03			
05710102	**Low discrimination**					**05710102**
					GRA	
Fusobacterium mortiferum		85.65	0.49	MLT 11	1	
Actinomyces viscosus		11.70	0.22	BGL 10	99	
Prevotella oralis		1.22	0.04		1	
05711000	**Low discrimination**					**05711000**
					GRA	
Prevotella intermedia		58.29	0.55	SAL 1	1	
Fusobacterium mortiferum		36.56	0.62		1	
Lactobacillus fermentum		1.47	0.40		99	
05711001	**Low discrimination**					**05711001**
					GRA	
Actinomyces naeslundii		47.79	0.46	SUC 99	99	
Prevotella intermedia		32.08	0.20		1	
Fusobacterium mortiferum		20.12	0.28		1	
05711002	**Low discrimination**					**05711002**
					GRA	
Fusobacterium mortiferum		49.01	0.46	MLT 11	1	
Lactobacillus fermentum		17.76	0.40	BGL 10	99	
Prevotella intermedia		7.10	0.20		1	
05711003	**Low discrimination**					**05711003**
					GRA	
Actinomyces naeslundii		86.22	0.46	SUC 99	99	
Actinomyces viscosus		9.75	0.06		99	
Fusobacterium mortiferum		4.03	0.12		1	
05711020	**Low discrimination**					**05711020**
					GRA	
Clostridium perfringens		71.25	0.41	SUC 91	99	
Prevotella intermedia		4.89	0.20	LAC 88	1	
Actinomyces naeslundii		4.86	0.43		99	
05711021	**Acceptable identification**					**05711021**
Actinomyces naeslundii		91.01	0.46	SUC 99		
Clostridium perfringens		8.99	0.07			
05711022	**Low discrimination**					**05711022**
					GRA	
Prevotella oralis		66.61	0.39	SUC 92	1	
Clostridium perfringens		24.88	0.35	LAC 94	99	
Actinomyces naeslundii		3.60	0.43		99	
05711023	**Low discrimination**					**05711023**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		85.40	0.46	SUC 99	99		
Prevotella oralis		10.63	0.05		1		
Clostridium perfringens		3.97	0.01		99		
05711100	**Good identification**						**05711100**
					GRA		
Fusobacterium mortiferum		91.59	0.69	MLT 11	1		
Lactobacillus fermentum		2.58	0.44		99		
Clostridium perfringens		2.09	0.30		99		
05711101	**Low discrimination**						**05711101**
					GRA		
Actinomyces naeslundii		74.34	0.54	SUC 99	99		
Fusobacterium mortiferum		25.66	0.34		1		
05711102	**Low discrimination**						**05711102**
					GRA		
Fusobacterium mortiferum		33.93	0.52	MLT 11	1		
Prevotella oralis		31.90	0.38	BGL 10	1		
Bifidobacterium longum subsp. infantis		12.77	0.27		99		
05711103	**Low discrimination**						**05711103**
					GRA		
Actinomyces naeslundii		86.25	0.54	SUC 99	99		
Actinomyces viscosus		6.09	0.10		99		
Fusobacterium mortiferum		3.31	0.18		1		
05711120	**Low discrimination**						**05711120**
					GRA	SPO	
Clostridium perfringens		56.87	0.56	SUC 91	99	84	
Clostridium septicum		25.26	0.43		99	99	
Prevotella oralis		8.06	0.38		1	1	
05711121	**Low discrimination**						**05711121**
					SPO		
Actinomyces naeslundii		69.96	0.54	SUC 99	1		
Clostridium perfringens		17.80	0.21		84		
Clostridium septicum		7.91	0.09		99		
05711122	**Low discrimination**						**05711122**
					GRA		
Prevotella oralis		78.40	0.60	SUC 92	1		
Clostridium perfringens		13.71	0.50		99		
Bifidobacterium longum subsp. infantis		2.94	0.32		99		
05711123	**Low discrimination**						**05711123**
					GRA		
Actinomyces naeslundii		53.90	0.54	SUC 99	99		
Prevotella oralis		36.92	0.25		1		
Clostridium perfringens		6.45	0.16		99		
05711132	**Low discrimination**						**05711132**
					GRA		
Parabacteroides distasonis		60.57	0.43	XYL 90	1		
Prevotella oralis		31.21	0.25		1		
Clostridium perfringens		5.46	0.16		99		
05711141	**Good identification**						**05711141**
Actinomyces naeslundii		100.0	0.41	SUC 99			
05711142	**Low discrimination**						**05711142**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		42.57	0.38	SUC 99	99	
Clostridium perfringens		25.27	0.12		99	
Fusobacterium mortiferum		13.88	0.18		1	
05711143	**Good identification**					**05711143**
Actinomyces naeslundii		100.0	0.41	SUC 99		
05711160	**Acceptable identification**					**05711160**
					SPO	
Clostridium perfringens		94.25	0.43	SUC 91	84	
Actinomyces naeslundii		2.31	0.38		1	
Clostridium septicum		2.22	0.09		99	
05711161	**Low discrimination**					**05711161**
Actinomyces naeslundii		78.45	0.41	SUC 99		
Clostridium perfringens		21.55	0.09			
05711162	**Low discrimination**					**05711162**
					GRA	
Clostridium perfringens		72.58	0.38	SUC 91	99	
Prevotella oralis		22.02	0.25		1	
Actinomyces naeslundii		3.78	0.38		99	
05711163	**Low discrimination**					**05711163**
Actinomyces naeslundii		88.56	0.41	SUC 99		
Clostridium perfringens		11.45	0.04			
05711322	**Low discrimination**					**05711322**
					GRA	
Prevotella oralis		79.38	0.53	SUC 92	1	
Parabacteroides distasonis		20.23	0.56		1	
Clostridium perfringens		0.33	0.16		99	
05711332	**Acceptable identification**					**05711332**
Parabacteroides distasonis		93.14	0.51	XYL 90		
Prevotella oralis		6.86	0.19			
05711400	**Low discrimination**					**05711400**
					GRA	
Lactobacillus fermentum		57.23	0.40	SUC 87	99	
Prevotella intermedia		22.88	0.20	SAL 1	1	
Fusobacterium mortiferum		14.35	0.28		1	
05711402	**Good identification**					**05711402**
					GRA	
Lactobacillus fermentum		97.29	0.40	SUC 87	99	
Fusobacterium mortiferum		2.71	0.12	SAL 1	1	
05711500	**Low discrimination**					**05711500**
					GRA	
Lactobacillus fermentum		72.24	0.44	SUC 87	99	
Fusobacterium mortiferum		25.91	0.34	SAL 1	1	
Bifidobacterium longum subsp. longum		1.85	0.09		99	
05711502	**Low discrimination**					**05711502**
					GRA	
Lactobacillus fermentum		57.68	0.44	SUC 87	99	
Bifidobacterium adolescentis		34.61	0.34	SAL 1	99	
Fusobacterium mortiferum		2.30	0.18		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05711512	**Acceptable identification**						**05711512**
Bifidobacterium adolescentis		94.38	0.36	SUC 99			
Bifidobacterium longum subsp. longum		5.62	0.13	XYL 87			
05711522	**Low discrimination**						**05711522**
					GRA		
Parabacteroides distasonis		54.76	0.44	XYL 90	1		
Prevotella oralis		24.79	0.25		1		
Lactobacillus fermentum		11.61	0.35		99		
05711532	**Low discrimination**						**05711532**
					GRA		
Parabacteroides distasonis		82.90	0.39	XYL 90	1		
Bifidobacterium longum subsp. longum		17.10	0.24		99		
05711722	**Acceptable identification**						**05711722**
Parabacteroides distasonis		93.02	0.52	XYL 90			
Prevotella oralis		6.02	0.19				
Bacteroides vulgatus		0.63	-.03				
05711732	**Good identification**						**05711732**
Parabacteroides distasonis		97.07	0.48	XYL 90			
Clostridium clostridioforme		2.93	0.16				
05712101	**Good identification**						**05712101**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05712103	**Good identification**						**05712103**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05712121	**Good identification**						**05712121**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05712123	**Good identification**						**05712123**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05713000	**Low discrimination**						**05713000**
					GRA	CAT	
Actinomyces naeslundii		39.47	0.48	SUC 99	99	25	
Propionibacterium granulosum		27.21	0.30		99	90	
Prevotella intermedia		20.48	0.20		1	1	
05713001	**Good identification**						**05713001**
Actinomyces naeslundii		100.0	0.51	SUC 99			
05713002	**Low discrimination**						**05713002**
					GRA	CAT	
Actinomyces naeslundii		82.75	0.48	SUC 99	99	25	
Propionibacterium granulosum		14.26	0.20		99	90	
Fusobacterium mortiferum		2.99	0.12		1	5	
05713003	**Good identification**						**05713003**
Actinomyces naeslundii		100.0	0.51	SUC 99			
05713020	**Low discrimination**						**05713020**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		80.09	0.48	SUC 99	25		
Propionibacterium granulosum		13.80	0.20		90		
Clostridium perfringens		6.11	0.07		1		
05713021	**Good identification**						**05713021**
Actinomyces naeslundii		100.0	0.51	SUC 99			
05713022	**Low discrimination**						**05713022**
					GRA	CAT	
Actinomyces naeslundii		85.10	0.48	SUC 99	99	25	
Prevotella oralis		8.18	0.05		1	1	
Propionibacterium granulosum		3.67	0.10		99	90	
05713023	**Good identification**						**05713023**
Actinomyces naeslundii		100.0	0.51	SUC 99			
05713040	**Good identification**						**05713040**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05713041	**Good identification**						**05713041**
Actinomyces naeslundii		100.0	0.38	SUC 99			
05713042	**Good identification**						**05713042**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05713043	**Good identification**						**05713043**
Actinomyces naeslundii		100.0	0.38	SUC 99			
05713060	**Good identification**						**05713060**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05713061	**Good identification**						**05713061**
Actinomyces naeslundii		100.0	0.38	SUC 99			
05713062	**Good identification**						**05713062**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05713063	**Good identification**						**05713063**
Actinomyces naeslundii		100.0	0.38	SUC 99			
05713100	**Low discrimination**						**05713100**
					GRA		
Actinomyces naeslundii		77.91	0.56	SUC 99	99		
Fusobacterium mortiferum		20.78	0.34		1		
Actinomyces israelii		1.31	0.16		99		
05713101	**Very good identification**						**05713101**
Actinomyces naeslundii		100.0	0.59	SUC 99			
05713102	**Low discrimination**						**05713102**
					GRA		
Actinomyces naeslundii		86.24	0.56	SUC 99	99		
Bifidobacterium adolescentis		4.70	0.19		99		
Fusobacterium mortiferum		2.56	0.18		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05713103	**Very good identification**					**05713103**
Actinomyces naeslundii		100.0	0.59	SUC 99		
05713111	**Good identification**					**05713111**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05713113	**Good identification**					**05713113**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05713120	**Low discrimination**					**05713120**
					SPO	
Actinomyces naeslundii		75.08	0.56	SUC 99	1	
Clostridium perfringens		14.77	0.21		84	
Clostridium septicum		6.56	0.09		99	
05713121	**Very good identification**					**05713121**
Actinomyces naeslundii		100.0	0.59	SUC 99		
05713122	**Low discrimination**					**05713122**
					GRA	
Actinomyces naeslundii		59.12	0.56	SUC 99	99	
Prevotella oralis		31.30	0.25		1	
Clostridium perfringens		5.47	0.16		99	
05713123	**Very good identification**					**05713123**
Actinomyces naeslundii		100.0	0.59	SUC 99		
05713131	**Good identification**					**05713131**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05713133	**Good identification**					**05713133**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05713140	**Good identification**					**05713140**
Actinomyces naeslundii		100.0	0.43	SUC 99		
05713141	**Good identification**					**05713141**
Actinomyces naeslundii		100.0	0.46	SUC 99		
05713142	**Acceptable identification**					**05713142**
Actinomyces naeslundii		94.05	0.43	SUC 99		
Propionibacterium propionicum		5.95	0.12			
05713143	**Good identification**					**05713143**
Actinomyces naeslundii		100.0	0.46	SUC 99		
05713160	**Low discrimination**					**05713160**
Actinomyces naeslundii		82.49	0.43	SUC 99		
Clostridium perfringens		17.51	0.09			
05713161	**Good identification**					**05713161**
Actinomyces naeslundii		100.0	0.46	SUC 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
05713162	**Acceptable identification**						**05713162**
Actinomyces naeslundii		90.92	0.43	SUC 99			
Clostridium perfringens		9.08	0.04				
05713163	**Good identification**						**05713163**
Actinomyces naeslundii		100.0	0.46	SUC 99			
05714000	**Low discrimination**						**05714000**
					COC	GRA	
Fusobacterium mortiferum		66.36	0.40	MLT 11	1	1	
Prevotella intermedia		21.30	0.20	XYL 7	1	1	
Gemella morbillorum		9.57	0.15		99	99	
05714100	**Good identification**						**05714100**
Fusobacterium mortiferum		100.0	0.46	MLT 11			
05714322	**Acceptable identification**						**05714322**
Parabacteroides distasonis		94.89	0.38	GAL 99			
Prevotella oralis		5.11	0.04				
05715000	**Low discrimination**						**05715000**
					GRA		
Fusobacterium mortiferum		63.29	0.43	MLT 11	1		
Lactobacillus fermentum		19.89	0.36	XYL 7	99		
Prevotella intermedia		13.54	0.20		1		
05715002	**Low discrimination**						**05715002**
					GRA		
Lactobacillus fermentum		67.45	0.36	SUC 87	99		
Fusobacterium mortiferum		23.85	0.27	SAL 1	1		
Prevotella oralis		5.48	0.02		1		
05715100	**Low discrimination**						**05715100**
					GRA		
Fusobacterium mortiferum		77.47	0.49	MLT 11	1		
Lactobacillus fermentum		17.02	0.40	XYL 7	99		
Bifidobacterium longum subsp. infantis		3.48	0.09		99		
05715112	**Low discrimination**						**05715112**
					GRA		
Bifidobacterium adolescentis		85.38	0.36	SUC 99	99		
Parabacteroides distasonis		14.62	0.25	ARA 87	1		
05715122	**Low discrimination**						**05715122**
					GRA		
Parabacteroides distasonis		67.24	0.64		1		
Prevotella oralis		25.37	0.44		1		
Bifidobacterium longum subsp. infantis		6.29	0.30		99		
05715132	**Very good identification**						**05715132**
					GRA		
Parabacteroides distasonis		99.12	0.59		1		
Prevotella oralis		0.70	0.10		1		
Bifidobacterium longum subsp. infantis		0.17	-.04		99		
05715222	**Low discrimination**						**05715222**
Parabacteroides distasonis		74.60	0.38	LAC 99			
Prevotella oralis		25.40	0.17				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
05715302	**Low discrimination**					**05715302**
					GRA	
Parabacteroides distasonis		71.24	0.38	NAG 99	1	
Prevotella oralis		20.00	0.16		1	
Eubacterium contortum		8.76	0.11		99	
05715320	**Acceptable identification**					**05715320**
Bacteroides vulgatus		92.27	0.41	SUC 96		
Parabacteroides distasonis		5.51	0.38	ARA 93		
Prevotella oralis		1.55	0.16			
05715322	**Good identification**					**05715322**
Parabacteroides distasonis		94.69	0.72			
Prevotella oralis		5.10	0.38			
Bacteroides vulgatus		0.16	0.06			
05715323	**Acceptable identification**					**05715323**
Parabacteroides distasonis		94.89	0.38	URE 1		
Prevotella oralis		5.11	0.04			
05715332	**Very good identification**					**05715332**
Parabacteroides distasonis		99.77	0.68			
Clostridium clostridioforme		0.13	0.13			
Prevotella oralis		0.10	0.04			
05715362	**Acceptable identification**					**05715362**
Parabacteroides distasonis		94.89	0.38	SOR 1		
Prevotella oralis		5.11	0.04			
05715400	**Good identification**					**05715400**
Lactobacillus fermentum		100.0	0.36	SUC 87		
05715402	**Low discrimination**					**05715402**
Lactobacillus fermentum		89.87	0.36	SUC 87		
Bifidobacterium adolescentis		10.13	0.14	SAL 1		
05715500	**Low discrimination**					**05715500**
					GRA	
Lactobacillus fermentum		60.49	0.40	SUC 87	99	
Bacteroides vulgatus		18.01	0.09	SAL 1	1	
Bifidobacterium longum subsp. longum		7.93	0.17		99	
05715502	**Low discrimination**					**05715502**
					GRA	
Bifidobacterium adolescentis		81.91	0.48	SUC 99	99	
Lactobacillus fermentum		11.98	0.40		99	
Parabacteroides distasonis		2.92	0.26		1	
05715512	**Acceptable identification**					**05715512**
Bifidobacterium adolescentis		93.99	0.50	SUC 99		
Bifidobacterium longum subsp. longum		2.51	0.21			
Clostridium clostridioforme		2.02	0.18			
05715520	**Good identification**					**05715520**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		96.40	0.43	SUC 96	1	
Bifidobacterium longum subsp. longum		1.71	0.28	RHA 90	99	
Lactobacillus fermentum		1.09	0.31		99	
05715522	**Good identification**					**05715522**
					GRA	
Parabacteroides distasonis		94.32	0.60		1	
Bifidobacterium longum subsp. longum		2.05	0.28		99	
Lactobacillus fermentum		1.31	0.31		99	
05715532	**Acceptable identification**					**05715532**
					GRA	
Parabacteroides distasonis		91.82	0.56		1	
Bifidobacterium longum subsp. longum		6.31	0.32		99	
Clostridium clostridioforme		1.27	0.18		50	
05715720	**Very good identification**					**05715720**
					GRA	
Bacteroides vulgatus		99.68	0.60	SUC 96	1	
Parabacteroides distasonis		0.27	0.34		1	
Eubacterium contortum		0.05	0.10		99	
05715722	**Very good identification**					**05715722**
Parabacteroides distasonis		95.97	0.69			
Bacteroides vulgatus		3.56	0.26			
Clostridium clostridioforme		0.38	0.22			
05715732	**Very good identification**					**05715732**
Parabacteroides distasonis		96.66	0.64			
Clostridium clostridioforme		3.28	0.33			
Bacteroides vulgatus		0.07	-.09			
05717322	**Acceptable identification**					**05717322**
Parabacteroides distasonis		94.89	0.38	MAN 1		
Prevotella oralis		5.11	0.04			
05717502	**Acceptable identification**					**05717502**
Bifidobacterium adolescentis		91.15	0.47	SUC 99		
Actinomyces israelii		8.85	0.36			
05717512	**Good identification**					**05717512**
Bifidobacterium adolescentis		98.83	0.48	SUC 99		
Actinomyces israelii		1.17	0.22			
05721101	**Good identification**					**05721101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05721103	**Good identification**					**05721103**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05721121	**Good identification**					**05721121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05721123	**Good identification**					**05721123**
Actinomyces naeslundii		100.0	0.37	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05723100	**Good identification**						**05723100**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05723101	**Good identification**						**05723101**
Actinomyces naeslundii		100.0	0.42	SUC 99			
05723102	**Good identification**						**05723102**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05723103	**Good identification**						**05723103**
Actinomyces naeslundii		100.0	0.42	SUC 99			
05723120	**Good identification**						**05723120**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05723121	**Good identification**						**05723121**
Actinomyces naeslundii		100.0	0.42	SUC 99			
05723122	**Good identification**						**05723122**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05723123	**Good identification**						**05723123**
Actinomyces naeslundii		100.0	0.42	SUC 99			
05725122	**Good identification**						**05725122**
Parabacteroides distasonis		100.0	0.41	FRU 99			
05725132	**Good identification**						**05725132**
Parabacteroides distasonis		100.0	0.37	FRU 99			
05725322	**Good identification**						**05725322**
Parabacteroides distasonis		100.0	0.49	FRU 99			
05725332	**Good identification**						**05725332**
Parabacteroides distasonis		100.0	0.45	FRU 99			
05725522	**Good identification**						**05725522**
Parabacteroides distasonis		100.0	0.37	FRU 99			
05725722	**Good identification**						**05725722**
Parabacteroides distasonis		100.0	0.46	FRU 99			
05725732	**Acceptable identification**						**05725732**
Parabacteroides distasonis		94.83	0.41	FRU 99			
Clostridium clostridioforme		5.17	0.14				
05730000	**Low discrimination**						**05730000**
					GRA	CAT	
Fusobacterium mortiferum		65.56	0.50	MLT 11	1	5	
Propionibacterium granulosum		28.02	0.41		99	90	
Prevotella intermedia		5.30	0.20		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05730100	**Very good identification**						**05730100**
					GRA		
Fusobacterium mortiferum		99.00	0.57	MLT 11	1		
Actinomyces naeslundii		0.78	0.32		99		
Clostridium perfringens		0.21	0.00		99		
05730101	**Low discrimination**						**05730101**
					GRA		
Actinomyces naeslundii		54.06	0.35	SUC 99	99		
Fusobacterium mortiferum		45.94	0.22	GAL 96	1		
05730102	**Acceptable identification**						**05730102**
					GRA		
Fusobacterium mortiferum		90.44	0.40	MLT 11	1		
Actinomyces naeslundii		6.45	0.32	BGL 10	99		
Actinomyces viscosus		3.11	0.03		99		
05730103	**Good identification**						**05730103**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05730121	**Good identification**						**05730121**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05730123	**Good identification**						**05730123**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05731000	**Low discrimination**						**05731000**
					GRA	CAT	
Fusobacterium mortiferum		62.15	0.53	MLT 11	1	5	
Propionibacterium granulosum		17.71	0.41		99	90	
Lactobacillus jensenii		7.53	0.28		99	1	
05731001	**Acceptable identification**						**05731001**
					GRA		
Actinomyces naeslundii		93.92	0.51	SUC 99	99		
Fusobacterium mortiferum		6.09	0.19		1		
05731002	**Low discrimination**						**05731002**
					GRA		
Fusobacterium mortiferum		28.01	0.37	MLT 11	1		
Actinomyces naeslundii		26.20	0.48	BGL 10	99		
Lactobacillus jensenii		20.35	0.25		99		
05731003	**Good identification**						**05731003**
Actinomyces naeslundii		100.0	0.51	SUC 99			
05731020	**Low discrimination**						**05731020**
					SPO		
Clostridium perfringens		83.11	0.45	SUC 91	84		
Actinomyces naeslundii		5.92	0.48	LAC 88	1		
Clostridium septicum		4.42	0.17		99		
05731021	**Acceptable identification**						**05731021**
Actinomyces naeslundii		91.37	0.51	SUC 99			
Clostridium perfringens		8.63	0.11				
05731022	**Low discrimination**						**05731022**
					GRA		
Clostridium perfringens		74.40	0.40	SUC 91	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		11.27	0.48	LAC 88	99	
Prevotella oralis		6.85	0.19		1	
05731023	**Good identification**					**05731023**
Actinomyces naeslundii		95.74	0.51	SUC 99		
Clostridium perfringens		4.26	0.06			
05731040	**Low discrimination**					**05731040**
					GRA	
Actinomyces naeslundii		49.53	0.35	SUC 99	99	
Fusobacterium mortiferum		27.28	0.19		1	
Clostridium perfringens		23.20	0.07		99	
05731041	**Good identification**					**05731041**
Actinomyces naeslundii		100.0	0.38	SUC 99		
05731042	**Low discrimination**					**05731042**
Actinomyces naeslundii		81.94	0.35	SUC 99		
Clostridium perfringens		18.06	0.02			
05731043	**Good identification**					**05731043**
Actinomyces naeslundii		100.0	0.38	SUC 99		
05731061	**Good identification**					**05731061**
Actinomyces naeslundii		100.0	0.38	SUC 99		
05731063	**Good identification**					**05731063**
Actinomyces naeslundii		100.0	0.38	SUC 99		
05731100	**Low discrimination**					**05731100**
					GRA	
Fusobacterium mortiferum		74.43	0.60	MLT 11	1	
Clostridium perfringens		10.55	0.34		99	
Actinomyces naeslundii		9.44	0.56		99	
05731101	**Good identification**					**05731101**
					GRA	
Actinomyces naeslundii		94.28	0.59	SUC 99	99	
Fusobacterium mortiferum		5.01	0.25		1	
Clostridium perfringens		0.71	0.00		99	
05731102	**Low discrimination**					**05731102**
					GRA	
Actinomyces naeslundii		33.26	0.56	SUC 99	99	
Fusobacterium mortiferum		29.15	0.43		1	
Clostridium perfringens		17.50	0.29		99	
05731103	**Very good identification**					**05731103**
Actinomyces naeslundii		100.0	0.59	SUC 99		
05731111	**Good identification**					**05731111**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05731113	**Good identification**					**05731113**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05731120	**Low discrimination**					**05731120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium perfringens		56.90	0.60	SUC 91	84	
Clostridium septicum		40.83	0.51		99	
Actinomyces naeslundii		1.57	0.56		1	
05731121	**Low discrimination**					**05731121**
					SPO	
Actinomyces naeslundii		70.51	0.59	SUC 99	1	
Clostridium perfringens		17.17	0.26		84	
Clostridium septicum		12.32	0.17		99	
05731122	**Low discrimination**					**05731122**
					GRA	
Clostridium perfringens		55.09	0.55	SUC 91	99	
Parabacteroides distasonis		28.14	0.59		1	
Prevotella oralis		10.83	0.39		1	
05731123	**Low discrimination**					**05731123**
					GRA	
Actinomyces naeslundii		83.64	0.59	SUC 99	99	
Clostridium perfringens		9.58	0.20		99	
Parabacteroides distasonis		4.90	0.25		1	
05731131	**Good identification**					**05731131**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05731132	**Acceptable identification**					**05731132**
					GRA	
Parabacteroides distasonis		94.77	0.54	XYL 90	1	
Clostridium perfringens		3.48	0.20		99	
Actinomyces naeslundii		1.07	0.34		99	
05731133	**Low discrimination**					**05731133**
					GRA	
Actinomyces naeslundii		62.55	0.37	SUC 99	99	
Parabacteroides distasonis		37.45	0.20	MLZ 5	1	
05731141	**Good identification**					**05731141**
Actinomyces naeslundii		100.0	0.46	SUC 99		
05731142	**Low discrimination**					**05731142**
Actinomyces naeslundii		58.57	0.43	SUC 99		
Clostridium perfringens		33.27	0.16			
Lactobacillus acidophilus		8.16	0.04			
05731143	**Good identification**					**05731143**
Actinomyces naeslundii		100.0	0.46	SUC 99		
05731160	**Acceptable identification**					**05731160**
					SPO	
Clostridium perfringens		94.01	0.48	SUC 91	84	
Clostridium septicum		3.58	0.17		99	
Actinomyces naeslundii		2.41	0.43		1	
05731161	**Low discrimination**					**05731161**
Actinomyces naeslundii		79.19	0.46	SUC 99		
Clostridium perfringens		20.81	0.14			
05731162	**Acceptable identification**					**05731162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Clostridium perfringens		90.95	0.42	SUC 91	99		
Actinomyces naeslundii		4.95	0.43		99		
Parabacteroides distasonis		2.46	0.25		1		
05731163	**Low discrimination**						**05731163**
Actinomyces naeslundii		88.99	0.46	SUC 99			
Clostridium perfringens		11.01	0.08				
05731322	**Good identification**						**05731322**
					GRA		
Parabacteroides distasonis		93.77	0.67	XYL 90	1		
Prevotella oralis		5.16	0.33		1		
Clostridium perfringens		0.62	0.20		99		
05731332	**Very good identification**						**05731332**
Parabacteroides distasonis		99.71	0.63	XYL 90			
Clostridium clostridioforme		0.19	0.10				
Prevotella oralis		0.10	-.01				
05731522	**Good identification**						**05731522**
					GRA		
Parabacteroides distasonis		95.74	0.55	XYL 90	1		
Clostridium perfringens		3.09	0.20		99		
Prevotella oralis		0.61	0.05		1		
05731532	**Good identification**						**05731532**
					GRA		
Parabacteroides distasonis		96.98	0.51	XYL 90	1		
Clostridium clostridioforme		1.93	0.16		50		
Bifidobacterium longum subsp. longum		1.10	0.13		99		
05731722	**Very good identification**						**05731722**
Parabacteroides distasonis		99.34	0.63	XYL 90			
Clostridium clostridioforme		0.57	0.19				
Prevotella oralis		0.09	-.01				
05731732	**Very good identification**						**05731732**
Parabacteroides distasonis		95.37	0.59	XYL 90			
Clostridium clostridioforme		4.63	0.31				
05732000	**Acceptable identification**						**05732000**
					GRA	CAT	
Propionibacterium granulosum		93.42	0.39	SUC 93	99	90	
Actinomyces naeslundii		3.66	0.29	SAL 1	99	25	
Fusobacterium mortiferum		2.93	0.16		1	5	
05732100	**Low discrimination**						**05732100**
					GRA		
Actinomyces naeslundii		60.36	0.37	SUC 99	99		
Fusobacterium mortiferum		39.64	0.22	GAL 96	1		
05732101	**Good identification**						**05732101**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05732102	**Good identification**						**05732102**
Actinomyces naeslundii		100.0	0.37	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05732103	**Good identification**						**05732103**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05732120	**Good identification**						**05732120**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05732121	**Good identification**						**05732121**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05732122	**Good identification**						**05732122**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05732123	**Good identification**						**05732123**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05733000	**Low discrimination**						**05733000**
					GRA	CAT	
Propionibacterium granulosum		49.81	0.39	SUC 93	99	90	
Actinomyces naeslundii		46.76	0.53	SAL 1	99	25	
Fusobacterium mortiferum		2.34	0.19		1	5	
05733001	**Good identification**						**05733001**
Actinomyces naeslundii		100.0	0.56	SUC 99			
05733002	**Low discrimination**						**05733002**
					CAT		
Actinomyces naeslundii		76.86	0.53	SUC 99	25		
Propionibacterium granulosum		20.47	0.28		90		
Actinomyces israelii		2.67	0.19		1		
05733003	**Good identification**						**05733003**
Actinomyces naeslundii		100.0	0.56	SUC 99			
05733020	**Low discrimination**						**05733020**
					CAT		
Actinomyces naeslundii		74.67	0.53	SUC 99	25		
Propionibacterium granulosum		19.88	0.28		90		
Clostridium perfringens		5.45	0.11		1		
05733021	**Good identification**						**05733021**
Actinomyces naeslundii		100.0	0.56	SUC 99			
05733022	**Acceptable identification**						**05733022**
					CAT		
Actinomyces naeslundii		90.83	0.53	SUC 99	25		
Propionibacterium granulosum		6.05	0.18		90		
Clostridium perfringens		3.12	0.06		1		
05733023	**Good identification**						**05733023**
Actinomyces naeslundii		100.0	0.56	SUC 99			
05733040	**Good identification**						**05733040**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05733041	**Good identification**						**05733041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	SUC 99		
05733042	**Good identification**					**05733042**
Actinomyces naeslundii		100.0	0.40	SUC 99		
05733043	**Good identification**					**05733043**
Actinomyces naeslundii		100.0	0.43	SUC 99		
05733060	**Good identification**					**05733060**
Actinomyces naeslundii		100.0	0.40	SUC 99		
05733061	**Good identification**					**05733061**
Actinomyces naeslundii		100.0	0.43	SUC 99		
05733062	**Good identification**					**05733062**
Actinomyces naeslundii		100.0	0.40	SUC 99		
05733063	**Good identification**					**05733063**
Actinomyces naeslundii		100.0	0.43	SUC 99		
05733100	**Good identification**					**05733100**
					GRA	
Actinomyces naeslundii		92.64	0.61	SUC 99	99	
Fusobacterium mortiferum		3.80	0.25		1	
Actinomyces israelii		3.02	0.26		99	
05733101	**Very good identification**					**05733101**
Actinomyces naeslundii		100.0	0.64	SUC 99		
05733102	**Good identification**					**05733102**
Actinomyces naeslundii		93.67	0.61	SUC 99		
Actinomyces israelii		4.58	0.29			
Bifidobacterium adolescentis		1.76	0.16			
05733103	**Very good identification**					**05733103**
Actinomyces naeslundii		100.0	0.64	SUC 99		
05733110	**Good identification**					**05733110**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05733111	**Good identification**					**05733111**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05733112	**Low discrimination**					**05733112**
Actinomyces naeslundii		63.54	0.39	SUC 99		
Bifidobacterium adolescentis		27.64	0.17	MLZ 5		
Actinomyces israelii		8.81	0.15			
05733113	**Good identification**					**05733113**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05733120	**Low discrimination**					**05733120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		72.24	0.61	SUC 99	1	
Clostridium perfringens		13.59	0.26		84	
Clostridium septicum		9.75	0.17		99	
05733121	**Very good identification**					**05733121**
Actinomyces naeslundii		100.0	0.64	SUC 99		
05733122	**Low discrimination**					**05733122**
					SPO	
Actinomyces naeslundii		71.64	0.61	SUC 99	1	
Clostridium ramosum		17.53	0.25		92	
Clostridium perfringens		6.34	0.20		84	
05733123	**Very good identification**					**05733123**
Actinomyces naeslundii		100.0	0.64	SUC 99		
05733130	**Good identification**					**05733130**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05733131	**Good identification**					**05733131**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05733132	**Low discrimination**					**05733132**
					GRA	
Actinomyces naeslundii		68.37	0.39	SUC 99	99	
Parabacteroides distasonis		31.63	0.20	MLZ 5	1	
05733133	**Good identification**					**05733133**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05733140	**Acceptable identification**					**05733140**
Actinomyces naeslundii		93.30	0.48	SUC 99		
Actinomyces israelii		6.70	0.19			
05733141	**Good identification**					**05733141**
Actinomyces naeslundii		100.0	0.51	SUC 99		
05733142	**Acceptable identification**					**05733142**
Actinomyces naeslundii		90.28	0.48	SUC 99		
Actinomyces israelii		9.72	0.22			
05733143	**Good identification**					**05733143**
Actinomyces naeslundii		100.0	0.51	SUC 99		
05733160	**Low discrimination**					**05733160**
Actinomyces naeslundii		83.12	0.48	SUC 99		
Clostridium perfringens		16.88	0.14			
05733161	**Good identification**					**05733161**
Actinomyces naeslundii		100.0	0.51	SUC 99		
05733162	**Acceptable identification**					**05733162**
Actinomyces naeslundii		91.28	0.48	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		8.72	0.08				
05733163	**Good identification**						**05733163**
Actinomyces naeslundii		100.0	0.51	SUC 99			
05734100	**Good identification**						**05734100**
Fusobacterium mortiferum		100.0	0.37	MLT 11			
05734122	**Good identification**						**05734122**
Parabacteroides distasonis		100.0	0.41	GAL 99			
05734132	**Good identification**						**05734132**
Parabacteroides distasonis		97.20	0.37	GAL 99			
Clostridium clostridioforme		2.80	0.04				
05734322	**Good identification**						**05734322**
Parabacteroides distasonis		99.18	0.49	GAL 99			
Clostridium clostridioforme		0.83	0.08				
05734332	**Acceptable identification**						**05734332**
Parabacteroides distasonis		93.42	0.45	GAL 99			
Clostridium clostridioforme		6.58	0.19				
05734522	**Acceptable identification**						**05734522**
Parabacteroides distasonis		91.99	0.37	GAL 99			
Clostridium clostridioforme		8.01	0.14				
05734712	**Good identification**						**05734712**
Clostridium clostridioforme		100.0	0.40	SUC 99			
05734722	**Low discrimination**						**05734722**
Parabacteroides distasonis		82.46	0.46	GAL 99			
Clostridium clostridioforme		17.54	0.28				
05734732	**Low discrimination**						**05734732**
Clostridium clostridioforme		64.29	0.40	SUC 99			
Parabacteroides distasonis		35.71	0.41				
05735022	**Acceptable identification**						**05735022**
					GRA		
Parabacteroides distasonis		94.07	0.41	LAC 99	1		
Prevotella oralis		3.14	0.03		1		
Clostridium perfringens		2.79	0.06		99		
05735032	**Good identification**						**05735032**
Parabacteroides distasonis		100.0	0.37	LAC 99			
05735100	**Low discrimination**						**05735100**
					COC	GRA	
Fusobacterium mortiferum		43.65	0.40	MLT 11	1	1	
Atopobium parvulum		43.55	0.29	XYL 7	99	99	
Actinomyces israelii		9.43	0.31		1	99	
05735102	**Low discrimination**						**05735102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		53.18	0.41	NAG 99	1		
Bifidobacterium adolescentis		20.02	0.31		99		
Actinomyces israelii		12.95	0.34		99		
05735112	**Low discrimination**						**05735112**
					GRA		
Parabacteroides distasonis		49.79	0.37	NAG 99	1		
Bifidobacterium adolescentis		42.54	0.33		99		
Clostridium clostridioforme		4.30	0.13		50		
05735120	**Low discrimination**						**05735120**
					GRA	SPO	
Parabacteroides distasonis		49.88	0.41	BGL 99	1	1	
Clostridium perfringens		23.06	0.26		99	84	
Clostridium septicum		16.54	0.17		99	99	
05735122	**Very good identification**						**05735122**
					GRA		
Parabacteroides distasonis		99.20	0.75		1		
Prevotella oralis		0.52	0.24		1		
Clostridium perfringens		0.22	0.20		99		
05735123	**Good identification**						**05735123**
					GRA		
Parabacteroides distasonis		98.12	0.41	URE 1	1		
Actinomyces naeslundii		1.88	0.25		99		
05735130	**Good identification**						**05735130**
Parabacteroides distasonis		100.0	0.37	BGL 99			
05735132	**Very good identification**						**05735132**
Parabacteroides distasonis		99.91	0.71				
Clostridium clostridioforme		0.09	0.13				
05735133	**Good identification**						**05735133**
Parabacteroides distasonis		100.0	0.37	URE 1			
05735162	**Good identification**						**05735162**
					GRA		
Parabacteroides distasonis		96.02	0.41	SOR 1	1		
Clostridium perfringens		3.98	0.08		99		
05735172	**Good identification**						**05735172**
Parabacteroides distasonis		100.0	0.37	SOR 1			
05735222	**Good identification**						**05735222**
Parabacteroides distasonis		98.79	0.49	LAC 99			
Clostridium clostridioforme		0.74	0.07				
Prevotella oralis		0.47	-.03				
05735232	**Acceptable identification**						**05735232**
Parabacteroides distasonis		94.06	0.45	LAC 99			
Clostridium clostridioforme		5.94	0.18				
05735302	**Good identification**						**05735302**
					GRA		
Parabacteroides distasonis		96.99	0.49	NAG 99	1		
Clostridium clostridioforme		2.42	0.16		50		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.59	0.15		99	
05735312	**Low discrimination**					**05735312**
Parabacteroides distasonis		82.56	0.45	NAG 99		
Clostridium clostridioforme		17.44	0.27			
05735320	**Low discrimination**					**05735320**
Parabacteroides distasonis		86.71	0.49	BGL 99		
Bacteroides vulgatus		13.29	0.17			
05735322	**Very good identification**					**05735322**
Parabacteroides distasonis		99.90	0.83			
Prevotella oralis		0.08	0.17			
Clostridium clostridioforme		0.03	0.16			
05735323	**Good identification**					**05735323**
Parabacteroides distasonis		100.0	0.49	URE 1		
05735330	**Good identification**					**05735330**
Parabacteroides distasonis		100.0	0.45	BGL 99		
05735332	**Very good identification**					**05735332**
Parabacteroides distasonis		99.79	0.79			
Clostridium clostridioforme		0.21	0.27			
05735333	**Good identification**					**05735333**
Parabacteroides distasonis		100.0	0.45	URE 1		
05735362	**Good identification**					**05735362**
Parabacteroides distasonis		100.0	0.49	SOR 1		
05735372	**Good identification**					**05735372**
Parabacteroides distasonis		100.0	0.45	SOR 1		
05735422	**Acceptable identification**					**05735422**
Parabacteroides distasonis		92.77	0.37	LAC 99		
Clostridium clostridioforme		7.24	0.13			
05735502	**Low discrimination**					**05735502**
					GRA	
Bifidobacterium adolescentis		68.78	0.45	SUC 99	99	
Parabacteroides distasonis		16.73	0.37		1	
Actinomyces israelii		5.44	0.33		99	
05735512	**Low discrimination**					**05735512**
					GRA	
Bifidobacterium adolescentis		73.33	0.47	SUC 99	99	
Clostridium clostridioforme		17.36	0.33		50	
Parabacteroides distasonis		7.86	0.33		1	
05735522	**Very good identification**					**05735522**
					GRA	
Parabacteroides distasonis		99.58	0.72		1	
Clostridium clostridioforme		0.26	0.22		50	
Bifidobacterium longum subsp. longum		0.12	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05735523	**Good identification**					**05735523**
Parabacteroides distasonis		100.0	0.37	URE 1		
05735532	**Very good identification**					**05735532**
					GRA	
Parabacteroides distasonis		97.37	0.67		1	
Clostridium clostridioforme		2.17	0.33		50	
Bifidobacterium longum subsp. longum		0.37	0.21		99	
05735562	**Good identification**					**05735562**
Parabacteroides distasonis		100.0	0.37	SOR 1		
05735612	**Good identification**					**05735612**
Clostridium clostridioforme		100.0	0.39	SUC 99		
05735622	**Low discrimination**					**05735622**
Parabacteroides distasonis		83.99	0.46	LAC 99		
Clostridium clostridioforme		16.01	0.28			
05735632	**Low discrimination**					**05735632**
Clostridium clostridioforme		61.73	0.39	SUC 99		
Parabacteroides distasonis		38.27	0.41			
05735702	**Low discrimination**					**05735702**
					GRA	
Parabacteroides distasonis		60.23	0.46	NAG 99	1	
Clostridium clostridioforme		38.44	0.37		50	
Bifidobacterium adolescentis		0.83	0.11		99	
05735712	**Low discrimination**					**05735712**
					GRA	
Clostridium clostridioforme		83.96	0.48	SUC 99	50	
Parabacteroides distasonis		15.55	0.41		1	
Bifidobacterium adolescentis		0.49	0.13		99	
05735720	**Low discrimination**					**05735720**
Bacteroides vulgatus		76.86	0.36	SUC 96		
Parabacteroides distasonis		23.14	0.46	TRE 4		
05735722	**Very good identification**					**05735722**
Parabacteroides distasonis		99.31	0.80			
Clostridium clostridioforme		0.64	0.37			
Bacteroides vulgatus		0.03	0.02			
05735723	**Good identification**					**05735723**
Parabacteroides distasonis		100.0	0.46	URE 1		
05735730	**Low discrimination**					**05735730**
Parabacteroides distasonis		89.54	0.41	BGL 99		
Bacteroides vulgatus		5.58	0.02			
Clostridium clostridioforme		4.88	0.14			
05735732	**Good identification**					**05735732**
Parabacteroides distasonis		94.83	0.75			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		5.17	0.48			
05735733	**Acceptable identification**					**05735733**
Parabacteroides distasonis		94.83	0.41	URE 1		
Clostridium clostridioforme		5.17	0.14			
05735762	**Good identification**					**05735762**
Parabacteroides distasonis		100.0	0.46	SOR 1		
05735772	**Acceptable identification**					**05735772**
Parabacteroides distasonis		94.83	0.41	SOR 1		
Clostridium clostridioforme		5.17	0.14			
05737002	**Good identification**					**05737002**
Actinomyces israelii		100.0	0.37	SUC 98		
05737100	**Good identification**					**05737100**
Actinomyces israelii		97.37	0.44	SUC 98		
Actinomyces naeslundii		2.63	0.27	CEL 93		
05737102	**Low discrimination**					**05737102**
Actinomyces israelii		80.57	0.47	SUC 98		
Bifidobacterium adolescentis		17.98	0.30	CEL 93		
Actinomyces naeslundii		1.45	0.27			
05737122	**Low discrimination**					**05737122**
					GRA	
Parabacteroides distasonis		89.85	0.41	MAN 1	1	
Actinomyces israelii		7.92	0.27		99	
Actinomyces naeslundii		2.23	0.27		99	
05737132	**Good identification**					**05737132**
Parabacteroides distasonis		100.0	0.37	MAN 1		
05737140	**Good identification**					**05737140**
Actinomyces israelii		100.0	0.37	SUC 98		
05737142	**Acceptable identification**					**05737142**
Actinomyces israelii		93.38	0.40	SUC 98		
Bifidobacterium adolescentis		6.62	0.14	CEL 93		
05737322	**Good identification**					**05737322**
Parabacteroides distasonis		100.0	0.49	MAN 1		
05737332	**Good identification**					**05737332**
Parabacteroides distasonis		100.0	0.45	MAN 1		
05737402	**Acceptable identification**					**05737402**
Actinomyces israelii		93.13	0.35	SUC 98		
Bifidobacterium adolescentis		6.87	0.10	CEL 93		
05737500	**Acceptable identification**					**05737500**
					SPO	
Actinomyces israelii		92.10	0.43	SUC 98	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		5.35	0.01	CEL 93	99	
Bifidobacterium adolescentis		2.55	0.10		1	
05737502	**Low discrimination**					**05737502**
					SPO	
Bifidobacterium adolescentis		64.26	0.44	SUC 99	1	
Actinomyces israelii		35.21	0.46		1	
Clostridium butyricum		0.53	-.07		99	
05737512	**Acceptable identification**					**05737512**
Bifidobacterium adolescentis		92.84	0.45	SUC 99		
Actinomyces israelii		6.22	0.32			
Bifidobacterium dentium		0.94	-.07			
05737522	**Low discrimination**					**05737522**
					GRA	
Parabacteroides distasonis		86.84	0.37	MAN 1	1	
Actinomyces israelii		10.21	0.25		99	
Bifidobacterium adolescentis		2.95	0.10		99	
05737540	**Good identification**					**05737540**
Actinomyces israelii		100.0	0.36	SUC 98		
05737542	**Low discrimination**					**05737542**
Actinomyces israelii		63.29	0.39	SUC 98		
Bifidobacterium adolescentis		36.71	0.28	CEL 93		
05737722	**Good identification**					**05737722**
Parabacteroides distasonis		100.0	0.46	MAN 1		
05737732	**Acceptable identification**					**05737732**
Parabacteroides distasonis		94.83	0.41	MAN 1		
Clostridium clostridioforme		5.17	0.14			
05743100	**Good identification**					**05743100**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05743101	**Good identification**					**05743101**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05743102	**Good identification**					**05743102**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05743103	**Good identification**					**05743103**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05743120	**Good identification**					**05743120**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05743121	**Good identification**					**05743121**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05743122	**Low discrimination**					**05743122**
					SPO GRA	
Actinomyces naeslundii		58.60	0.36	SUC 99	1 99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		25.47	0.05	FRU 95	92	50	
Prevotella oralis		15.92	0.01		1	1	
05743123	**Good identification**						**05743123**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05750000	**Acceptable identification**						**05750000**
					GRA		
Fusobacterium mortiferum		91.55	0.51	MLT 11	1		
Prevotella intermedia		6.63	0.20		1		
Lactobacillus jensenii		0.95	0.07		99		
05750100	**Very good identification**						**05750100**
					GRA		
Fusobacterium mortiferum		99.52	0.57	MLT 11	1		
Actinomyces naeslundii		0.48	0.29		99		
05750102	**Low discrimination**						**05750102**
					GRA		
Fusobacterium mortiferum		71.67	0.41	MLT 11	1		
Prevotella oralis		10.86	0.13	BGL 10	1		
Actinomyces viscosus		7.81	0.12		99		
05751000	**Low discrimination**						**05751000**
					GRA		
Fusobacterium mortiferum		51.76	0.54	MLT 11	1		
Lactobacillus jensenii		35.56	0.41		99		
Lactobacillus catenaformis		6.44	0.29		99		
05751001	**Low discrimination**						**05751001**
					GRA		
Actinomyces naeslundii		84.84	0.48	SUC 99	99		
Fusobacterium mortiferum		8.98	0.20		1		
Lactobacillus jensenii		6.17	0.07		99		
05751002	**Low discrimination**						**05751002**
					GRA		
Lactobacillus jensenii		42.61	0.38	SUC 90	99		
Prevotella oralis		15.39	0.27	RAF 1	1		
Lactobacillus acidophilus		12.62	0.26		99		
05751003	**Good identification**						**05751003**
Actinomyces naeslundii		95.37	0.48	SUC 99			
Lactobacillus jensenii		4.63	0.04				
05751021	**Good identification**						**05751021**
Actinomyces naeslundii		100.0	0.48	SUC 99			
05751022	**Low discrimination**						**05751022**
					GRA		
Prevotella oralis		89.28	0.49	SUC 92	1		
Lactobacillus acidophilus		3.85	0.26	LAC 94	99		
Lactobacillus jensenii		3.25	0.28		99		
05751023	**Low discrimination**						**05751023**
					GRA		
Actinomyces naeslundii		75.03	0.48	SUC 99	99		
Prevotella oralis		24.98	0.14		1		
05751041	**Good identification**						**05751041**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.35	SUC 99			
05751043	**Good identification**						**05751043**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05751061	**Good identification**						**05751061**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05751063	**Good identification**						**05751063**
Actinomyces naeslundii		100.0	0.35	SUC 99			
05751100	**Low discrimination**						**05751100**
					GRA		
Fusobacterium mortiferum		78.31	0.60	MLT 11	1		
Lactobacillus catenaformis		6.26	0.32		99		
Actinomyces naeslundii		6.08	0.53		99		
05751101	**Acceptable identification**						**05751101**
					GRA		
Actinomyces naeslundii		92.01	0.56	SUC 99	99		
Fusobacterium mortiferum		7.99	0.26		1		
05751102	**Low discrimination**						**05751102**
					GRA		
Prevotella oralis		52.75	0.47	SUC 92	1		
Lactobacillus acidophilus		31.74	0.45	NAG 95	99		
Fusobacterium mortiferum		5.28	0.44		1		
05751103	**Low discrimination**						**05751103**
					GRA		
Actinomyces naeslundii		85.57	0.56	SUC 99	99		
Prevotella oralis		8.25	0.13		1		
Lactobacillus acidophilus		4.97	0.10		99		
05751120	**Low discrimination**						**05751120**
					GRA	SPO	
Clostridium septicum		66.32	0.51	RAF 1	99	99	
Prevotella oralis		23.68	0.47		1	1	
Clostridium paraputrificum		4.50	0.29		99	99	
05751121	**Low discrimination**						**05751121**
					GRA	SPO	
Actinomyces naeslundii		72.21	0.56	SUC 99	99	1	
Clostridium septicum		19.50	0.17		99	99	
Prevotella oralis		6.96	0.13		1	1	
05751122	**Good identification**						**05751122**
					GRA		
Prevotella oralis		94.61	0.69	SUC 92	1		
Lactobacillus acidophilus		3.00	0.45		99		
Parabacteroides distasonis		0.52	0.43		1		
05751140	**Low discrimination**						**05751140**
					GRA		
Actinomyces naeslundii		50.19	0.40	SUC 99	99		
Fusobacterium mortiferum		37.03	0.26		1		
Bifidobacterium breve		6.39	0.10		99		
05751141	**Good identification**						**05751141**
Actinomyces naeslundii		100.0	0.43	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05751143	**Good identification**						**05751143**
Actinomyces naeslundii		100.0	0.43	SUC 99			
05751161	**Good identification**						**05751161**
Actinomyces naeslundii		100.0	0.43	SUC 99			
05751163	**Acceptable identification**						**05751163**
					GRA		
Actinomyces naeslundii		90.51	0.43	SUC 99	99		
Prevotella oralis		9.49	0.01		1		
05751222	**Good identification**						**05751222**
					GRA		
Prevotella oralis		98.65	0.42	SUC 92	1		
Lactobacillus acidophilus		0.75	0.07	LAC 94	99		
Parabacteroides distasonis		0.60	0.17		1		
05751302	**Low discrimination**						**05751302**
					GRA		
Prevotella oralis		89.84	0.41	SUC 92	1		
Lactobacillus acidophilus		9.49	0.25	NAG 95	99		
Parabacteroides distasonis		0.67	0.17		1		
05751320	**Low discrimination**						**05751320**
					SPO	GRA	
Prevotella oralis		89.30	0.41	SUC 92	1	1	
Clostridium septicum		5.89	0.17	BGL 95	99	99	
Clostridium ramosum		3.74	0.18		92	50	
05751322	**Good identification**						**05751322**
					SPO		
Prevotella oralis		94.94	0.63	SUC 92	1		
Parabacteroides distasonis		3.67	0.51		1		
Clostridium ramosum		0.84	0.28		92		
05751332	**Low discrimination**						**05751332**
Parabacteroides distasonis		67.02	0.47	XYL 90			
Prevotella oralis		32.49	0.29				
Clostridium clostridioforme		0.49	0.04				
05751502	**Low discrimination**						**05751502**
					GRA		
Bifidobacterium adolescentis		80.46	0.40	SUC 99	99		
Prevotella oralis		8.38	0.13	XYL 87	1		
Lactobacillus acidophilus		5.04	0.10		99		
05751512	**Good identification**						**05751512**
Bifidobacterium adolescentis		97.81	0.41	SUC 99			
Clostridium clostridioforme		2.20	0.10	XYL 87			
05751722	**Low discrimination**						**05751722**
Parabacteroides distasonis		69.06	0.48	XYL 90			
Prevotella oralis		29.41	0.29				
Clostridium clostridioforme		1.53	0.14				
05751732	**Low discrimination**						**05751732**
Parabacteroides distasonis		84.21	0.43	XYL 90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		15.79	0.25				
05752101	**Good identification**						**05752101**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05752103	**Good identification**						**05752103**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05752121	**Good identification**						**05752121**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05752123	**Good identification**						**05752123**
Actinomyces naeslundii		100.0	0.37	SUC 99			
05753000	**Low discrimination**						**05753000**
					SPO		
Actinomyces naeslundii		47.80	0.50	SUC 99	1		
Clostridium innocuum		39.88	0.25		99		
Actinomyces israelii		5.73	0.25		1		
05753001	**Good identification**						**05753001**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05753002	**Low discrimination**						**05753002**
					SPO		
Actinomyces naeslundii		47.41	0.50	SUC 99	1		
Clostridium innocuum		39.55	0.25		99		
Actinomyces israelii		8.52	0.28		1		
05753003	**Good identification**						**05753003**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05753020	**Good identification**						**05753020**
Actinomyces naeslundii		100.0	0.50	SUC 99			
05753021	**Good identification**						**05753021**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05753022	**Low discrimination**						**05753022**
					SPO	GRA	
Actinomyces naeslundii		72.45	0.50	SUC 99	1	99	
Prevotella oralis		18.64	0.14		1	1	
Clostridium ramosum		4.72	0.05		92	50	
05753023	**Good identification**						**05753023**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05753040	**Low discrimination**						**05753040**
Actinomyces naeslundii		79.11	0.37	SUC 99			
Actinomyces israelii		20.89	0.18				
05753041	**Good identification**						**05753041**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05753042	**Low discrimination**						**05753042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		71.62	0.37	SUC 99		
Actinomyces israelii		28.38	0.21			
05753043	**Good identification**					**05753043**
Actinomyces naeslundii		100.0	0.40	SUC 99		
05753060	**Good identification**					**05753060**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05753061	**Good identification**					**05753061**
Actinomyces naeslundii		100.0	0.40	SUC 99		
05753062	**Good identification**					**05753062**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05753063	**Good identification**					**05753063**
Actinomyces naeslundii		100.0	0.40	SUC 99		
05753100	**Low discrimination**					**05753100**
					GRA	
Actinomyces naeslundii		80.19	0.58	SUC 99	99	
Actinomyces israelii		13.51	0.35		99	
Fusobacterium mortiferum		5.38	0.26		1	
05753101	**Very good identification**					**05753101**
Actinomyces naeslundii		100.0	0.61	SUC 99		
05753102	**Low discrimination**					**05753102**
Actinomyces naeslundii		55.92	0.58	SUC 99		
Actinomyces israelii		14.13	0.39			
Lactobacillus acidophilus		13.07	0.23			
05753103	**Very good identification**					**05753103**
Actinomyces naeslundii		100.0	0.61	SUC 99		
05753110	**Low discrimination**					**05753110**
Actinomyces naeslundii		58.88	0.36	SUC 99		
Actinomyces israelii		28.16	0.21	MLZ 5		
Bifidobacterium breve		12.96	0.10			
05753111	**Good identification**					**05753111**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05753113	**Good identification**					**05753113**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05753120	**Low discrimination**					**05753120**
					SPO	
Actinomyces naeslundii		52.77	0.58	SUC 99	1	
Clostridium ramosum		29.88	0.28		92	
Clostridium septicum		11.01	0.17		99	
05753121	**Very good identification**					**05753121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.61	SUC 99			
05753122	**Low discrimination**						**05753122**
					SPO	GRA	
Clostridium ramosum		43.87	0.39	SUC 99	92	50	
Prevotella oralis		27.42	0.35	TRE 94	1	1	
Actinomyces naeslundii		19.37	0.58		1	99	
05753123	**Very good identification**						**05753123**
					SPO	GRA	
Actinomyces naeslundii		97.58	0.61	SUC 99	1	99	
Clostridium ramosum		1.49	0.05		92	50	
Prevotella oralis		0.93	0.01		1	1	
05753130	**Low discrimination**						**05753130**
Actinomyces naeslundii		71.88	0.36	SUC 99			
Bifidobacterium breve		28.12	0.15	MLZ 5			
05753131	**Good identification**						**05753131**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05753133	**Good identification**						**05753133**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05753140	**Low discrimination**						**05753140**
Actinomyces naeslundii		69.60	0.45	SUC 99			
Actinomyces israelii		25.83	0.28				
Bifidobacterium breve		4.57	0.10				
05753141	**Good identification**						**05753141**
Actinomyces naeslundii		100.0	0.48	SUC 99			
05753142	**Low discrimination**						**05753142**
Actinomyces naeslundii		40.89	0.45	SUC 99			
Bifidobacterium breve		27.14	0.27				
Actinomyces israelii		22.77	0.31				
05753143	**Good identification**						**05753143**
Actinomyces naeslundii		100.0	0.48	SUC 99			
05753160	**Low discrimination**						**05753160**
Actinomyces naeslundii		89.55	0.45	SUC 99			
Bifidobacterium breve		10.45	0.15				
05753161	**Good identification**						**05753161**
Actinomyces naeslundii		100.0	0.48	SUC 99			
05753163	**Good identification**						**05753163**
Actinomyces naeslundii		100.0	0.48	SUC 99			
05753322	**Low discrimination**						**05753322**
					SPO		
Clostridium ramosum		82.02	0.41	SUC 99	92		
Prevotella oralis		16.58	0.29	TRE 94	1		
Parabacteroides distasonis		0.64	0.17		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05753502	**Low discrimination**						**05753502**
Bifidobacterium adolescentis		72.96	0.38	SUC 99			
Actinomyces israelii		25.78	0.37	XYL 87			
Actinomyces naeslundii		1.26	0.24				
05753512	**Good identification**						**05753512**
Bifidobacterium adolescentis		95.86	0.40	SUC 99			
Actinomyces israelii		4.14	0.23	XYL 87			
05754100	**Good identification**						**05754100**
Fusobacterium mortiferum		100.0	0.38	MLT 11			
05755100	**Low discrimination**						**05755100**
					GRA	SPO	
Fusobacterium mortiferum		46.93	0.41	MLT 11	1	1	
Actinomyces israelii		32.10	0.41	XYL 7	99	1	
Clostridium butyricum		15.47	0.14		99	99	
05755112	**Low discrimination**						**05755112**
					GRA		
Bifidobacterium adolescentis		86.50	0.41	SUC 99	99		
Actinomyces israelii		7.58	0.30	ARA 87	99		
Parabacteroides distasonis		3.75	0.21		1		
05755122	**Low discrimination**						**05755122**
					GRA		
Prevotella oralis		71.00	0.53	SUC 92	1		
Parabacteroides distasonis		28.58	0.59	XYL 11	1		
Lactobacillus acidophilus		0.18	0.10		99		
05755132	**Good identification**						**05755132**
Parabacteroides distasonis		95.03	0.55				
Prevotella oralis		4.43	0.19				
Clostridium clostridioforme		0.32	0.07				
05755142	**Low discrimination**						**05755142**
					GRA		
Actinomyces israelii		67.14	0.37	SUC 98	99		
Bifidobacterium adolescentis		29.76	0.24		99		
Prevotella oralis		3.10	-.03		1		
05755322	**Low discrimination**						**05755322**
Parabacteroides distasonis		73.76	0.68				
Prevotella oralis		26.17	0.47				
Clostridium clostridioforme		0.07	0.10				
05755332	**Very good identification**						**05755332**
Parabacteroides distasonis		98.53	0.63				
Clostridium clostridioforme		0.81	0.22				
Prevotella oralis		0.66	0.13				
05755502	**Low discrimination**						**05755502**
					SPO		
Bifidobacterium adolescentis		81.42	0.54	SUC 99	1		
Clostridium butyricum		9.81	0.24		99		
Actinomyces israelii		7.14	0.42		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05755512	**Acceptable identification**						**05755512**
Bifidobacterium adolescentis		94.90	0.56	SUC 99			
Clostridium clostridioforme		3.22	0.27				
Actinomyces israelii		1.02	0.28				
05755522	**Acceptable identification**						**05755522**
					GRA	SPO	
Parabacteroides distasonis		91.94	0.56		1	1	
Prevotella oralis		3.76	0.19		1	1	
Clostridium butyricum		1.08	0.05		99	99	
05755532	**Low discrimination**						**05755532**
					GRA		
Parabacteroides distasonis		89.94	0.51		1		
Clostridium clostridioforme		7.75	0.27		50		
Bifidobacterium adolescentis		2.31	0.21		99		
05755542	**Low discrimination**						**05755542**
					SPO		
Bifidobacterium adolescentis		73.50	0.39	SUC 99	1		
Actinomyces israelii		20.27	0.35	SOR 11	1		
Clostridium butyricum		6.23	0.05		99		
05755552	**Good identification**						**05755552**
Bifidobacterium adolescentis		96.74	0.40	SUC 99			
Actinomyces israelii		3.26	0.21	SOR 11			
05755712	**Acceptable identification**						**05755712**
					GRA		
Clostridium clostridioforme		91.86	0.42	SUC 99	50		
Parabacteroides distasonis		4.41	0.25	TRE 88	1		
Bifidobacterium adolescentis		3.73	0.21		99		
05755720	**Good identification**						**05755720**
Bacteroides vulgatus		97.26	0.38	SUC 96			
Parabacteroides distasonis		2.74	0.30	CEL 5			
05755722	**Very good identification**						**05755722**
Parabacteroides distasonis		96.05	0.64				
Clostridium clostridioforme		2.39	0.31				
Prevotella oralis		0.56	0.13				
05755732	**Low discrimination**						**05755732**
Parabacteroides distasonis		82.32	0.59				
Clostridium clostridioforme		17.34	0.42				
Bacteroides ovatus		0.34	0.03				
05757000	**Good identification**						**05757000**
					SPO		
Actinomyces israelii		96.91	0.43	SUC 98	1		
Clostridium innocuum		3.09	0.03		99		
05757002	**Good identification**						**05757002**
					SPO		
Actinomyces israelii		97.92	0.46	SUC 98	1		
Clostridium innocuum		2.08	0.03		99		
05757040	**Good identification**						**05757040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	SUC 98		
05757042	**Good identification**					**05757042**
Actinomyces israelii		100.0	0.39	SUC 98		
05757100	**Good identification**					**05757100**
					SPO	
Actinomyces israelii		98.56	0.54	SUC 98	1	
Clostridium butyricum		0.93	-.02		99	
Actinomyces naeslundii		0.51	0.24		1	
05757102	**Low discrimination**					**05757102**
Actinomyces israelii		82.78	0.57	SUC 98		
Bifidobacterium adolescentis		16.67	0.38			
Actinomyces naeslundii		0.29	0.24			
05757110	**Good identification**					**05757110**
Actinomyces israelii		100.0	0.40	SUC 98		
05757112	**Low discrimination**					**05757112**
					SPO	
Bifidobacterium adolescentis		61.76	0.40	SUC 99	1	
Actinomyces israelii		37.49	0.43	ARA 87	1	
Clostridium tertium		0.75	-.01		99	
05757122	**Low discrimination**					**05757122**
					GRA	
Actinomyces israelii		38.76	0.36	SUC 98	99	
Prevotella oralis		36.57	0.19	NAG 6	1	
Parabacteroides distasonis		14.72	0.25		1	
05757140	**Good identification**					**05757140**
Actinomyces israelii		100.0	0.47	SUC 98		
05757142	**Acceptable identification**					**05757142**
Actinomyces israelii		93.98	0.50	SUC 98		
Bifidobacterium adolescentis		6.02	0.23			
05757152	**Low discrimination**					**05757152**
Actinomyces israelii		65.64	0.36	SUC 98		
Bifidobacterium adolescentis		34.36	0.24	MLZ 13		
05757302	**Good identification**					**05757302**
Actinomyces israelii		100.0	0.37	SUC 98		
05757400	**Good identification**					**05757400**
Actinomyces israelii		100.0	0.42	SUC 98		
05757402	**Acceptable identification**					**05757402**
Actinomyces israelii		93.76	0.45	SUC 98		
Bifidobacterium adolescentis		6.24	0.18			
05757442	**Good identification**					**05757442**
Actinomyces israelii		100.0	0.38	SUC 98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05757500	**Low discrimination**					**05757500**
					SPO	
Actinomyces israelii		88.58	0.52	SUC 98	1	
Clostridium butyricum		9.21	0.14		99	
Bifidobacterium adolescentis		2.21	0.18		1	
05757502	**Low discrimination**					**05757502**
					SPO	
Bifidobacterium adolescentis		61.60	0.53	SUC 99	1	
Actinomyces israelii		37.39	0.55		1	
Clostridium butyricum		1.01	0.07		99	
05757510	**Low discrimination**					**05757510**
					SPO	
Actinomyces israelii		80.60	0.38	SUC 98	1	
Bifidobacterium adolescentis		16.45	0.20	MLZ 13	1	
Clostridium butyricum		2.95	-.08		99	
05757512	**Acceptable identification**					**05757512**
Bifidobacterium adolescentis		92.82	0.54	SUC 99		
Actinomyces israelii		6.89	0.41			
Bifidobacterium dentium		0.29	-.07			
05757540	**Good identification**					**05757540**
					SPO	
Actinomyces israelii		97.73	0.45	SUC 98	1	
Clostridium butyricum		2.27	-.04		99	
05757542	**Low discrimination**					**05757542**
Actinomyces israelii		65.64	0.48	SUC 98		
Bifidobacterium adolescentis		34.36	0.37			
05757552	**Low discrimination**					**05757552**
Bifidobacterium adolescentis		81.07	0.39	SUC 99		
Actinomyces israelii		18.93	0.34	SOR 11		
05757702	**Low discrimination**					**05757702**
Actinomyces israelii		81.90	0.36	SUC 98		
Bifidobacterium adolescentis		18.11	0.18	RHA 7		
05761100	**Low discrimination**					**05761100**
					GRA	
Actinomyces naeslundii		72.43	0.36	SUC 99	99	
Fusobacterium mortiferum		27.57	0.17	FRU 95	1	
05761101	**Good identification**					**05761101**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05761103	**Good identification**					**05761103**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05761121	**Good identification**					**05761121**
Actinomyces naeslundii		100.0	0.39	SUC 99		
05761123	**Good identification**					**05761123**
Actinomyces naeslundii		100.0	0.39	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05763001	**Good identification**					**05763001**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05763003	**Good identification**					**05763003**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05763021	**Good identification**					**05763021**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05763023	**Good identification**					**05763023**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05763100	**Good identification**					**05763100**
Actinomyces naeslundii		100.0	0.41	SUC 99		
05763101	**Good identification**					**05763101**
Actinomyces naeslundii		100.0	0.44	SUC 99		
05763102	**Acceptable identification**					**05763102**
Actinomyces naeslundii		91.41	0.41	SUC 99		
Actinomyces israelii		8.59	0.14	FRU 95		
05763103	**Good identification**					**05763103**
Actinomyces naeslundii		100.0	0.44	SUC 99		
05763120	**Low discrimination**					**05763120**
					SPO	
Actinomyces naeslundii		53.28	0.41	SUC 99	1	
Clostridium ramosum		46.72	0.15	FRU 95	92	
05763121	**Good identification**					**05763121**
Actinomyces naeslundii		100.0	0.44	SUC 99		
05763123	**Good identification**					**05763123**
Actinomyces naeslundii		100.0	0.44	SUC 99		
05765122	**Good identification**					**05765122**
Parabacteroides distasonis		96.64	0.37	FRU 99		
Prevotella oralis		3.36	-.01			
05765322	**Good identification**					**05765322**
Parabacteroides distasonis		100.0	0.45	FRU 99		
05765332	**Good identification**					**05765332**
Parabacteroides distasonis		100.0	0.40	FRU 99		
05765722	**Good identification**					**05765722**
Parabacteroides distasonis		96.15	0.41	FRU 99		
Clostridium clostridioforme		3.85	0.11			
05765732	**Low discrimination**					**05765732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		74.68	0.36	FRU 99		
Clostridium clostridioforme		25.33	0.23			
05770000	**Low discrimination**					**05770000**
					GRA	
Fusobacterium mortiferum		88.46	0.42	MLT 11	1	
Lactobacillus jensenii		9.25	0.15		99	
Actinomyces naeslundii		2.29	0.27		99	
05770100	**Good identification**					**05770100**
					GRA	
Fusobacterium mortiferum		96.95	0.48	MLT 11	1	
Actinomyces naeslundii		3.06	0.34		99	
05770101	**Low discrimination**					**05770101**
					GRA	
Actinomyces naeslundii		82.40	0.37	SUC 99	99	
Fusobacterium mortiferum		17.61	0.14	GAL 96	1	
05770103	**Good identification**					**05770103**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05770121	**Good identification**					**05770121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05770123	**Good identification**					**05770123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
05771000	**Low discrimination**					**05771000**
					GRA	
Lactobacillus jensenii		82.52	0.50	SUC 90	99	
Fusobacterium mortiferum		11.96	0.45	RAF 1	1	
Actinomyces naeslundii		4.94	0.50		99	
05771001	**Low discrimination**					**05771001**
					GRA	
Actinomyces naeslundii		88.60	0.53	SUC 99	99	
Lactobacillus jensenii		9.96	0.15		99	
Fusobacterium mortiferum		1.44	0.11		1	
05771002	**Low discrimination**					**05771002**
Lactobacillus jensenii		68.36	0.47	SUC 90		
Lactobacillus acidophilus		22.59	0.35	RAF 1		
Actinomyces naeslundii		6.14	0.50			
05771003	**Acceptable identification**					**05771003**
Actinomyces naeslundii		90.93	0.53	SUC 99		
Lactobacillus jensenii		6.82	0.12			
Lactobacillus acidophilus		2.25	0.01			
05771010	**Good identification**					**05771010**
					GRA	
Lactobacillus jensenii		98.63	0.41	SUC 90	99	
Actinomyces naeslundii		0.93	0.28	RAF 1	99	
Fusobacterium mortiferum		0.43	0.11		1	
05771012	**Good identification**					**05771012**
Lactobacillus jensenii		96.92	0.38	SUC 90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		1.38	0.28	RAF 1			
Lactobacillus acidophilus		0.97	0.01				
05771020	**Low discrimination**						**05771020**
					SPO		
Lactobacillus jensenii		55.00	0.39	SUC 90	1		
Clostridium septicum		23.48	0.25	RAF 1	99		
Actinomyces naeslundii		13.18	0.50		1		
05771021	**Good identification**						**05771021**
Actinomyces naeslundii		97.27	0.53	SUC 99			
Lactobacillus jensenii		2.73	0.05				
05771022	**Low discrimination**						**05771022**
					GRA		
Lactobacillus acidophilus		38.27	0.35	SUC 99	99		
Lactobacillus jensenii		28.95	0.36	LAC 92	99		
Prevotella oralis		16.91	0.28		1		
05771023	**Good identification**						**05771023**
Actinomyces naeslundii		95.83	0.53	SUC 99			
Lactobacillus acidophilus		2.37	0.01				
Lactobacillus jensenii		1.80	0.02				
05771040	**Low discrimination**						**05771040**
Actinomyces naeslundii		44.06	0.37	SUC 99			
Lactobacillus jensenii		42.11	0.15				
Actinomyces israelii		7.73	0.15				
05771041	**Good identification**						**05771041**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05771043	**Good identification**						**05771043**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05771060	**Low discrimination**						**05771060**
Actinomyces naeslundii		58.31	0.37	SUC 99			
Clostridium perfringens		27.76	0.09				
Lactobacillus jensenii		13.93	0.05				
05771061	**Good identification**						**05771061**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05771063	**Good identification**						**05771063**
Actinomyces naeslundii		100.0	0.40	SUC 99			
05771100	**Low discrimination**						**05771100**
					GRA	SPO	
Fusobacterium mortiferum		42.68	0.51	MLT 11	1	1	
Actinomyces naeslundii		21.52	0.58		99	1	
Clostridium septicum		13.47	0.25		99	99	
05771101	**Very good identification**						**05771101**
					GRA		
Actinomyces naeslundii		98.68	0.61	SUC 99	99		
Fusobacterium mortiferum		1.32	0.17		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05771102	**Low discrimination**						**05771102**
					GRA		
Lactobacillus acidophilus		86.52	0.54	SUC 99	99		
Actinomyces naeslundii		5.82	0.58		99		
Prevotella oralis		2.74	0.27		1		
05771103	**Good identification**						**05771103**
Actinomyces naeslundii		90.90	0.61	SUC 99			
Lactobacillus acidophilus		9.10	0.19				
05771110	**Low discrimination**						**05771110**
					GRA		
Actinomyces naeslundii		44.23	0.36	SUC 99	99		
Fusobacterium mortiferum		16.84	0.17	MLZ 5	1		
Lactobacillus jensenii		16.60	0.07		99		
05771111	**Good identification**						**05771111**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05771113	**Low discrimination**						**05771113**
					GRA		
Actinomyces naeslundii		88.90	0.39	SUC 99	99		
Leptotrichia buccalis		11.10	-.11	MLZ 5	1		
05771120	**Good identification**						**05771120**
Clostridium septicum		92.70	0.60	RAF 1			
Clostridium ramosum		2.34	0.36				
Clostridium perfringens		1.70	0.37				
05771121	**Low discrimination**						**05771121**
					SPO		
Actinomyces naeslundii		69.66	0.61	SUC 99	1		
Clostridium septicum		29.07	0.25		99		
Clostridium ramosum		0.73	0.02		92		
05771122	**Low discrimination**						**05771122**
					SPO	GRA	
Lactobacillus acidophilus		40.44	0.54	SUC 99	1	99	
Prevotella oralis		24.35	0.49		1	1	
Clostridium ramosum		17.04	0.46		92	50	
05771123	**Low discrimination**						**05771123**
					GRA		
Actinomyces naeslundii		80.99	0.61	SUC 99	99		
Lactobacillus acidophilus		8.11	0.19		99		
Prevotella oralis		4.88	0.14		1		
05771131	**Good identification**						**05771131**
Actinomyces naeslundii		100.0	0.39	SUC 99			
05771132	**Low discrimination**						**05771132**
					GRA		
Parabacteroides distasonis		75.24	0.50	XYL 90	1		
Lactobacillus acidophilus		5.94	0.19		99		
Bifidobacterium breve		5.49	0.29		99		
05771133	**Low discrimination**						**05771133**
					GRA		
Actinomyces naeslundii		80.43	0.39	SUC 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		19.57	0.16	MLZ 5	1		
05771140	**Low discrimination**						**05771140**
					GRA		
Actinomyces naeslundii		68.26	0.45	SUC 99	99		
Actinomyces israelii		16.82	0.25		99		
Fusobacterium mortiferum		7.75	0.17		1		
05771141	**Good identification**						**05771141**
Actinomyces naeslundii		100.0	0.48	SUC 99			
05771142	**Low discrimination**						**05771142**
Lactobacillus acidophilus		84.83	0.38	SUC 99			
Actinomyces naeslundii		8.15	0.45	SOR 11			
Bifidobacterium breve		3.61	0.24				
05771143	**Acceptable identification**						**05771143**
Actinomyces naeslundii		93.45	0.48	SUC 99			
Lactobacillus acidophilus		6.55	0.04				
05771161	**Good identification**						**05771161**
Actinomyces naeslundii		100.0	0.48	SUC 99			
05771162	**Low discrimination**						**05771162**
Lactobacillus acidophilus		74.42	0.38	SUC 99			
Actinomyces naeslundii		7.15	0.45	SOR 11			
Bifidobacterium breve		5.62	0.29				
05771163	**Acceptable identification**						**05771163**
Actinomyces naeslundii		93.45	0.48	SUC 99			
Lactobacillus acidophilus		6.55	0.04				
05771320	**Low discrimination**						**05771320**
					SPO	GRA	
Clostridium ramosum		68.20	0.38	SUC 99	92	50	
Clostridium septicum		20.57	0.25		99	99	
Prevotella oralis		6.63	0.20		1	1	
05771322	**Low discrimination**						**05771322**
					SPO		
Parabacteroides distasonis		44.35	0.63	XYL 90	1		
Clostridium ramosum		34.75	0.48		92		
Prevotella oralis		16.06	0.42		1		
05771332	**Very good identification**						**05771332**
					SPO		
Parabacteroides distasonis		96.59	0.58	XYL 90	1		
Clostridium ramosum		1.42	0.14		92		
Clostridium clostridioforme		1.14	0.19		66		
05771502	**Low discrimination**						**05771502**
					SPO		
Bifidobacterium adolescentis		43.07	0.37	SUC 99	1		
Clostridium butyricum		25.24	0.18	XYL 87	99		
Lactobacillus acidophilus		13.56	0.19		1		
05771512	**Low discrimination**						**05771512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		74.38	0.38	SUC 99		
Clostridium clostridioforme		18.36	0.25	XYL 87		
Actinomyces israelii		2.62	0.20			
05771522	**Low discrimination**					**05771522**
					GRA	
Parabacteroides distasonis		84.69	0.51	XYL 90	1	
Lactobacillus acidophilus		5.88	0.19		99	
Prevotella oralis		3.54	0.14		1	
05771532	**Low discrimination**					**05771532**
Parabacteroides distasonis		89.01	0.46	XYL 90		
Clostridium clostridioforme		10.99	0.25			
05771712	**Good identification**					**05771712**
Clostridium clostridioforme		96.76	0.40	SUC 99		
Parabacteroides distasonis		3.24	0.20	XYL 91		
05771722	**Good identification**					**05771722**
					SPO	
Parabacteroides distasonis		94.67	0.59	XYL 90	1	
Clostridium clostridioforme		3.38	0.28		66	
Clostridium ramosum		1.22	0.14		92	
05771732	**Low discrimination**					**05771732**
Parabacteroides distasonis		76.81	0.54	XYL 90		
Clostridium clostridioforme		23.19	0.40			
05772100	**Low discrimination**					**05772100**
					GRA	
Actinomyces naeslundii		85.83	0.39	SUC 99	99	
Fusobacterium mortiferum		14.17	0.14	GAL 96	1	
05772101	**Good identification**					**05772101**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05772102	**Low discrimination**					**05772102**
Actinomyces naeslundii		89.39	0.39	SUC 99		
Actinomyces israelii		10.61	0.14	GAL 96		
05772103	**Good identification**					**05772103**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05772121	**Good identification**					**05772121**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05772123	**Good identification**					**05772123**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05773000	**Low discrimination**					**05773000**
					SPO	
Actinomyces naeslundii		67.44	0.55	SUC 99	1	
Actinomyces israelii		15.67	0.35		1	
Clostridium innocuum		10.18	0.17		99	
05773001	**Very good identification**					**05773001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.58	SUC 99		
05773002	**Low discrimination**					**05773002**
					SPO	
Actinomyces naeslundii		60.35	0.55	SUC 99	1	
Actinomyces israelii		21.03	0.38		1	
Clostridium innocuum		9.11	0.17		99	
05773003	**Very good identification**					**05773003**
Actinomyces naeslundii		100.0	0.58	SUC 99		
05773011	**Good identification**					**05773011**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05773013	**Good identification**					**05773013**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05773020	**Low discrimination**					**05773020**
					SPO	
Actinomyces naeslundii		85.62	0.55	SUC 99	1	
Clostridium ramosum		11.25	0.15		92	
Lactobacillus jensenii		1.86	0.05		1	
05773021	**Very good identification**					**05773021**
Actinomyces naeslundii		100.0	0.58	SUC 99		
05773022	**Low discrimination**					**05773022**
					SPO	
Actinomyces naeslundii		60.17	0.55	SUC 99	1	
Clostridium ramosum		31.62	0.25		92	
Lactobacillus acidophilus		6.00	0.14		1	
05773023	**Very good identification**					**05773023**
Actinomyces naeslundii		100.0	0.58	SUC 99		
05773031	**Good identification**					**05773031**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05773033	**Good identification**					**05773033**
Actinomyces naeslundii		100.0	0.36	SUC 99		
05773040	**Low discrimination**					**05773040**
Actinomyces naeslundii		66.15	0.42	SUC 99		
Actinomyces israelii		33.86	0.28			
05773041	**Good identification**					**05773041**
Actinomyces naeslundii		100.0	0.45	SUC 99		
05773042	**Low discrimination**					**05773042**
Actinomyces naeslundii		54.42	0.42	SUC 99		
Actinomyces israelii		41.78	0.31			
Lactobacillus acidophilus		3.80	-.02			
05773043	**Good identification**					**05773043**
Actinomyces naeslundii		100.0	0.45	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
05773060	**Good identification**					**05773060**
Actinomyces naeslundii		100.0	0.42	SUC 99		
05773061	**Good identification**					**05773061**
Actinomyces naeslundii		100.0	0.45	SUC 99		
05773062	**Acceptable identification**					**05773062**
Actinomyces naeslundii		93.47	0.42	SUC 99		
Lactobacillus acidophilus		6.53	-.02			
05773063	**Good identification**					**05773063**
Actinomyces naeslundii		100.0	0.45	SUC 99		
05773100	**Low discrimination**					**05773100**
					SPO	
Actinomyces naeslundii		71.79	0.63	SUC 99	1	
Actinomyces israelii		23.44	0.45		1	
Clostridium ramosum		3.31	0.15		92	
05773101	**Very good identification**					**05773101**
Actinomyces naeslundii		100.0	0.66	SUC 99		
05773102	**Low discrimination**					**05773102**
Actinomyces naeslundii		46.22	0.63	SUC 99		
Actinomyces israelii		22.63	0.48			
Lactobacillus acidophilus		18.62	0.32			
05773103	**Very good identification**					**05773103**
Actinomyces naeslundii		99.67	0.66	SUC 99		
Actinomyces israelii		0.33	0.14			
05773110	**Low discrimination**					**05773110**
					SPO	
Actinomyces naeslundii		47.38	0.41	SUC 99	1	
Actinomyces israelii		43.91	0.31	MLZ 5	1	
Clostridium tertium		8.71	0.05		99	
05773111	**Good identification**					**05773111**
Actinomyces naeslundii		100.0	0.44	SUC 99		
05773113	**Good identification**					**05773113**
Actinomyces naeslundii		100.0	0.44	SUC 99		
05773120	**Low discrimination**					**05773120**
					SPO	
Clostridium ramosum		76.90	0.49	SUC 99	92	
Actinomyces naeslundii		16.83	0.63		1	
Clostridium septicum		5.43	0.25		99	
05773121	**Very good identification**					**05773121**
					SPO	
Actinomyces naeslundii		97.02	0.66	SUC 99	1	
Clostridium ramosum		2.99	0.15		92	
05773122	**Good identification**					**05773122**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		91.83	0.59	SUC 99	92		
Actinomyces naeslundii		5.02	0.63		1		
Lactobacillus acidophilus		2.02	0.32		1		
05773123	**Low discrimination**						**05773123**
					SPO		
Actinomyces naeslundii		89.04	0.66	SUC 99	1		
Clostridium ramosum		10.96	0.25		92		
05773130	**Low discrimination**						**05773130**
					SPO		
Actinomyces naeslundii		39.02	0.41	SUC 99	1		
Clostridium ramosum		34.22	0.15	MLZ 5	92		
Clostridium tertium		21.52	0.13		99		
05773131	**Good identification**						**05773131**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05773133	**Good identification**						**05773133**
Actinomyces naeslundii		100.0	0.44	SUC 99			
05773140	**Low discrimination**						**05773140**
Actinomyces naeslundii		58.16	0.50	SUC 99			
Actinomyces israelii		41.84	0.38				
05773141	**Good identification**						**05773141**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05773142	**Low discrimination**						**05773142**
Actinomyces israelii		41.68	0.41	SUC 98			
Actinomyces naeslundii		38.62	0.50	XYL 92			
Lactobacillus acidophilus		10.90	0.16				
05773143	**Good identification**						**05773143**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05773160	**Low discrimination**						**05773160**
					SPO		
Actinomyces naeslundii		74.21	0.50	SUC 99	1		
Clostridium ramosum		19.41	0.15		92		
Actinomyces israelii		3.41	0.18		1		
05773161	**Good identification**						**05773161**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05773163	**Good identification**						**05773163**
Actinomyces naeslundii		100.0	0.53	SUC 99			
05773320	**Good identification**						**05773320**
					SPO		
Clostridium ramosum		99.83	0.51	SUC 99	92		
Actinomyces naeslundii		0.17	0.29		1		
05773322	**Very good identification**						**05773322**
					SPO	GRA	
Clostridium ramosum		99.53	0.61	SUC 99	92	50	
Parabacteroides distasonis		0.23	0.28		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		0.13	0.12		1 99	
05773402	**Acceptable identification**					**05773402**
					SPO	
Actinomyces israelii		91.45	0.37	SUC 98	1	
Clostridium innocuum		8.55	0.04	XYL 92	99	
05773500	**Low discrimination**					**05773500**
					SPO	
Actinomyces israelii		83.91	0.44	SUC 98	1	
Clostridium butyricum		12.92	0.09	XYL 92	99	
Actinomyces naeslundii		3.17	0.29		1	
05773502	**Low discrimination**					**05773502**
					SPO	
Actinomyces israelii		63.83	0.47	SUC 98	1	
Bifidobacterium adolescentis		32.02	0.35	XYL 92	1	
Clostridium butyricum		2.55	0.02		99	
05773512	**Low discrimination**					**05773512**
Bifidobacterium adolescentis		78.40	0.37	SUC 99		
Actinomyces israelii		19.11	0.33	XYL 87		
Bifidobacterium dentium		2.49	-.07			
05773540	**Good identification**					**05773540**
Actinomyces israelii		100.0	0.37	SUC 98		
05773542	**Low discrimination**					**05773542**
Actinomyces israelii		86.25	0.40	SUC 98		
Bifidobacterium adolescentis		13.75	0.20	XYL 92		
05774122	**Good identification**					**05774122**
Parabacteroides distasonis		96.64	0.37	GAL 99		
Prevotella oralis		3.36	-.01			
05774322	**Good identification**					**05774322**
Parabacteroides distasonis		95.08	0.45	GAL 99		
Clostridium clostridioforme		4.92	0.17			
05774332	**Low discrimination**					**05774332**
Parabacteroides distasonis		69.55	0.40	GAL 99		
Clostridium clostridioforme		30.45	0.28			
05774612	**Good identification**					**05774612**
Clostridium clostridioforme		100.0	0.40	SUC 99		
05774632	**Good identification**					**05774632**
Clostridium clostridioforme		100.0	0.40	SUC 99		
05774702	**Good identification**					**05774702**
Clostridium clostridioforme		100.0	0.37	SUC 99		
05774712	**Good identification**					**05774712**
Clostridium clostridioforme		100.0	0.49	SUC 99		
05774722	**Low discrimination**					**05774722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		56.95	0.37	SUC 99			
Parabacteroides distasonis		43.05	0.41				
05774732	**Acceptable identification**						**05774732**
Clostridium clostridioforme		91.80	0.49	SUC 99			
Parabacteroides distasonis		8.20	0.36				
05775000	**Low discrimination**						**05775000**
					GRA	SPO	
Actinomyces israelii		57.12	0.40	SUC 98	99	1	
Clostridium butyricum		19.11	0.11		99	99	
Fusobacterium mortiferum		11.37	0.26		1	1	
05775002	**Low discrimination**						**05775002**
					SPO		
Actinomyces israelii		83.63	0.43	SUC 98	1		
Clostridium butyricum		7.25	0.04		99		
Lactobacillus jensenii		6.85	0.12		1		
05775022	**Low discrimination**						**05775022**
					GRA		
Parabacteroides distasonis		72.08	0.37	LAC 99	1		
Prevotella oralis		15.85	0.13		1		
Actinomyces israelii		6.91	0.23		99		
05775042	**Good identification**						**05775042**
Actinomyces israelii		100.0	0.36	SUC 98			
05775102	**Low discrimination**						**05775102**
					SPO		
Actinomyces israelii		46.26	0.54	SUC 98	1		
Clostridium butyricum		24.07	0.28		99		
Bifidobacterium adolescentis		11.44	0.37		1		
05775110	**Low discrimination**						**05775110**
					COC	SPO	
Actinomyces israelii		56.76	0.36	SUC 98	1	1	
Clostridium butyricum		40.13	0.13	MLZ 13	1	99	
Atopobium parvulum		3.11	0.01		99	1	
05775112	**Low discrimination**						**05775112**
					GRA		
Bifidobacterium adolescentis		48.59	0.38	SUC 99	99		
Actinomyces israelii		24.03	0.39	ARA 87	99		
Parabacteroides distasonis		14.41	0.32		1		
05775122	**Very good identification**						**05775122**
					GRA		
Parabacteroides distasonis		95.19	0.71		1		
Prevotella oralis		3.31	0.33		1		
Lactobacillus acidophilus		0.45	0.19		99		
05775123	**Acceptable identification**						**05775123**
					GRA		
Parabacteroides distasonis		92.42	0.37	URE 1	1		
Actinomyces naeslundii		4.36	0.27		99		
Prevotella oralis		3.22	-.01		1		
05775132	**Very good identification**						**05775132**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		99.29	0.66		1		
Clostridium clostridioforme		0.54	0.22		50		
Actinomyces israelii		0.11	0.19		99		
05775140	**Low discrimination**						**05775140**
					COC	SPO	
Actinomyces israelii		68.03	0.44	SUC 98	1	1	
Clostridium butyricum		30.53	0.17		1	99	
Atopobium parvulum		1.43	0.01		99	1	
05775142	**Low discrimination**						**05775142**
					SPO		
Actinomyces israelii		81.99	0.47	SUC 98	1		
Clostridium butyricum		9.54	0.10		99		
Bifidobacterium adolescentis		6.44	0.21		1		
05775162	**Low discrimination**						**05775162**
					GRA		
Parabacteroides distasonis		80.53	0.37	SOR 1	1		
Actinomyces israelii		12.01	0.26		99		
Lactobacillus acidophilus		4.66	0.04		99		
05775222	**Acceptable identification**						**05775222**
Parabacteroides distasonis		92.78	0.45	LAC 99			
Clostridium clostridioforme		4.30	0.16				
Prevotella oralis		2.92	0.06				
05775232	**Low discrimination**						**05775232**
Parabacteroides distasonis		71.82	0.40	LAC 99			
Clostridium clostridioforme		28.18	0.27				
05775302	**Low discrimination**						**05775302**
					GRA		
Parabacteroides distasonis		72.68	0.45	NAG 99	1		
Clostridium clostridioforme		11.28	0.25		50		
Actinomyces israelii		10.96	0.34		99		
05775312	**Low discrimination**						**05775312**
					GRA		
Clostridium clostridioforme		55.77	0.36	SUC 99	50		
Parabacteroides distasonis		42.46	0.40	ARA 94	1		
Actinomyces israelii		1.78	0.20		99		
05775320	**Acceptable identification**						**05775320**
					SPO		
Parabacteroides distasonis		94.10	0.45	BGL 99	1		
Prevotella oralis		2.44	0.05		1		
Clostridium ramosum		2.05	0.04		92		
05775322	**Very good identification**						**05775322**
Parabacteroides distasonis		99.23	0.79				
Prevotella oralis		0.49	0.27				
Clostridium clostridioforme		0.16	0.25				
05775323	**Good identification**						**05775323**
Parabacteroides distasonis		100.0	0.45	URE 1			
05775330	**Good identification**						**05775330**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.40	BGL 99			
05775332	**Very good identification**						**05775332**
Parabacteroides distasonis		98.67	0.74				
Clostridium clostridioforme		1.31	0.36				
Bacteroides ovatus		0.02	-.06				
05775333	**Good identification**						**05775333**
Parabacteroides distasonis		100.0	0.40	URE 1			
05775362	**Good identification**						**05775362**
Parabacteroides distasonis		100.0	0.45	SOR 1			
05775372	**Good identification**						**05775372**
Parabacteroides distasonis		100.0	0.40	SOR 1			
05775402	**Low discrimination**						**05775402**
					SPO		
Actinomyces israelii		43.91	0.42	SUC 98	1		
Clostridium butyricum		41.89	0.20		99		
Clostridium clostridioforme		10.62	0.22		66		
05775500	**Good identification**						**05775500**
					SPO		
Clostridium butyricum		95.50	0.51	ESL 88	99		
Actinomyces israelii		4.33	0.49	SUC 99	1		
Bifidobacterium adolescentis		0.13	0.17		1		
05775502	**Low discrimination**						**05775502**
					SPO		
Clostridium butyricum		63.09	0.44	ESL 88	99		
Bifidobacterium adolescentis		22.28	0.51	SUC 99	1		
Actinomyces israelii		11.02	0.52		1		
05775512	**Low discrimination**						**05775512**
					SPO		
Bifidobacterium adolescentis		63.03	0.53	SUC 99	1		
Clostridium clostridioforme		23.51	0.42		66		
Clostridium butyricum		7.69	0.22		99		
05775522	**Good identification**						**05775522**
					GRA	SPO	
Parabacteroides distasonis		93.87	0.67		1	1	
Clostridium butyricum		3.79	0.26		99	99	
Clostridium clostridioforme		1.54	0.31		50	66	
05775532	**Low discrimination**						**05775532**
					GRA	SPO	
Parabacteroides distasonis		86.81	0.62		1	1	
Clostridium clostridioforme		12.04	0.42		50	66	
Clostridium butyricum		0.34	0.04		99	99	
05775552	**Low discrimination**						**05775552**
					SPO		
Bifidobacterium adolescentis		76.54	0.37	SUC 99	1		
Actinomyces israelii		14.56	0.31	SOR 11	1		
Clostridium butyricum		6.57	0.04		99		
05775602	**Acceptable identification**						**05775602**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		94.42	0.37	SUC 99	66		
Actinomyces israelii		4.01	0.23		1		
Clostridium butyricum		1.57	-.06		99		
05775612	**Good identification**						**05775612**
Clostridium clostridioforme		100.0	0.48	SUC 99			
05775622	**Low discrimination**						**05775622**
Clostridium clostridioforme		54.24	0.37	SUC 99			
Parabacteroides distasonis		45.76	0.41				
05775632	**Acceptable identification**						**05775632**
Clostridium clostridioforme		90.93	0.48	SUC 99			
Parabacteroides distasonis		9.07	0.36				
05775702	**Low discrimination**						**05775702**
					GRA	SPO	
Clostridium clostridioforme		69.65	0.46	SUC 99	50	66	
Parabacteroides distasonis		17.55	0.41		1	1	
Clostridium butyricum		8.31	0.18		99	99	
05775712	**Very good identification**						**05775712**
					GRA		
Clostridium clostridioforme		96.48	0.57	SUC 99	50		
Parabacteroides distasonis		2.87	0.36		1		
Bifidobacterium adolescentis		0.36	0.18		99		
05775720	**Low discrimination**						**05775720**
					GRA	SPO	
Parabacteroides distasonis		68.00	0.41	BGL 99	1	1	
Bacteroides vulgatus		22.08	0.14		1	1	
Clostridium butyricum		7.20	0.07		99	99	
05775722	**Very good identification**						**05775722**
Parabacteroides distasonis		95.35	0.75				
Clostridium clostridioforme		3.82	0.46				
Bacteroides ovatus		0.71	0.23				
05775723	**Good identification**						**05775723**
Parabacteroides distasonis		96.15	0.41	URE 1			
Clostridium clostridioforme		3.85	0.11				
05775730	**Low discrimination**						**05775730**
Parabacteroides distasonis		74.68	0.36	BGL 99			
Clostridium clostridioforme		25.33	0.23				
05775732	**Low discrimination**						**05775732**
Parabacteroides distasonis		74.13	0.71				
Clostridium clostridioforme		25.14	0.57				
Bacteroides ovatus		0.69	0.20				
05775733	**Low discrimination**						**05775733**
Parabacteroides distasonis		74.68	0.36	URE 1			
Clostridium clostridioforme		25.33	0.23				
05775762	**Acceptable identification**						**05775762**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		93.38	0.41	SOR 1		
Clostridium clostridioforme		3.74	0.11			
Bacteroides ovatus		2.88	-.01			
05775772	**Low discrimination**					**05775772**
Parabacteroides distasonis		72.61	0.36	SOR 1		
Clostridium clostridioforme		24.62	0.23			
Bacteroides ovatus		2.77	-.04			
05777000	**Good identification**					**05777000**
					SPO	
Actinomyces israelii		99.35	0.53	SUC 98	1	
Clostridium butyricum		0.65	-.05		99	
05777002	**Very good identification**					**05777002**
Actinomyces israelii		100.0	0.56	SUC 98		
05777010	**Good identification**					**05777010**
Actinomyces israelii		100.0	0.39	SUC 98		
05777012	**Good identification**					**05777012**
Actinomyces israelii		100.0	0.42	SUC 98		
05777022	**Good identification**					**05777022**
Actinomyces israelii		100.0	0.36	SUC 98		
05777040	**Good identification**					**05777040**
Actinomyces israelii		100.0	0.46	SUC 98		
05777042	**Good identification**					**05777042**
Actinomyces israelii		100.0	0.49	SUC 98		
05777052	**Good identification**					**05777052**
Actinomyces israelii		100.0	0.35	SUC 98		
05777100	**Very good identification**					**05777100**
					SPO	
Actinomyces israelii		95.65	0.64	SUC 98	1	
Clostridium butyricum		3.76	0.18		99	
Actinomyces naeslundii		0.26	0.29		1	
05777102	**Very good identification**					**05777102**
					SPO	
Actinomyces israelii		95.09	0.67	SUC 98	1	
Bifidobacterium adolescentis		3.39	0.35		1	
Clostridium butyricum		0.97	0.11		99	
05777110	**Good identification**					**05777110**
					SPO	
Actinomyces israelii		96.37	0.49	SUC 98	1	
Clostridium tertium		2.30	0.07	MLZ 13	99	
Clostridium butyricum		1.34	-.04		99	
05777112	**Low discrimination**					**05777112**
					SPO	
Actinomyces israelii		74.14	0.52	SUC 98	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		21.64	0.37	MLZ 13	1		
Clostridium tertium		3.53	0.15		99		
05777120	**Low discrimination**						**05777120**
					SPO		
Actinomyces israelii		72.33	0.43	SUC 98	1		
Clostridium ramosum		13.93	0.15	NAG 6	92		
Clostridium tertium		6.81	0.11		99		
05777122	**Low discrimination**						**05777122**
					SPO	GRA	
Actinomyces israelii		45.05	0.46	SUC 98	1	99	
Clostridium ramosum		23.14	0.25	NAG 6	92	50	
Parabacteroides distasonis		20.72	0.37		1	1	
05777140	**Very good identification**						**05777140**
					SPO		
Actinomyces israelii		99.13	0.56	SUC 98	1		
Clostridium butyricum		0.87	0.00		99		
05777142	**Very good identification**						**05777142**
					SPO		
Actinomyces israelii		98.66	0.59	SUC 98	1		
Bifidobacterium adolescentis		1.12	0.20		1		
Clostridium butyricum		0.23	-.07		99		
05777150	**Good identification**						**05777150**
					COC		
Actinomyces israelii		96.59	0.42	SUC 98	1		
Blautia producta		3.41	-.03	MLZ 13	99		
05777152	**Acceptable identification**						**05777152**
Actinomyces israelii		91.51	0.45	SUC 98			
Bifidobacterium adolescentis		8.49	0.21	MLZ 13			
05777160	**Good identification**						**05777160**
Actinomyces israelii		100.0	0.36	SUC 98			
05777162	**Good identification**						**05777162**
Actinomyces israelii		100.0	0.39	SUC 98			
05777202	**Good identification**						**05777202**
Actinomyces israelii		100.0	0.37	SUC 98			
05777300	**Good identification**						**05777300**
					SPO		
Actinomyces israelii		98.41	0.44	SUC 98	1		
Clostridium butyricum		1.59	-.08	RHA 7	99		
05777302	**Good identification**						**05777302**
					GRA		
Actinomyces israelii		98.83	0.47	SUC 98	99		
Parabacteroides distasonis		1.17	0.11	RHA 7	1		
05777322	**Low discrimination**						**05777322**
					SPO	GRA	
Parabacteroides distasonis		64.60	0.45	MAN 1	1	1	
Clostridium ramosum		31.88	0.27		92	50	
Actinomyces israelii		3.52	0.27		1	99	
05777332	**Good identification**						**05777332**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		100.0	0.40	MAN 1		
05777340	**Good identification**					**05777340**
Actinomyces israelii		100.0	0.37	SUC 98		
05777342	**Good identification**					**05777342**
Actinomyces israelii		100.0	0.40	SUC 98		
05777400	**Acceptable identification**					**05777400**
					SPO	
Actinomyces israelii		93.27	0.52	SUC 98	1	
Clostridium butyricum		6.73	0.11		99	
05777402	**Good identification**					**05777402**
					SPO	
Actinomyces israelii		97.04	0.55	SUC 98	1	
Clostridium butyricum		1.82	0.04		99	
Bifidobacterium adolescentis		1.15	0.15		1	
05777410	**Good identification**					**05777410**
Actinomyces israelii		100.0	0.38	SUC 98		
05777412	**Low discrimination**					**05777412**
Actinomyces israelii		87.58	0.41	SUC 98		
Bifidobacterium adolescentis		8.45	0.17	MLZ 13		
Bifidobacterium dentium		3.97	-.07			
05777440	**Good identification**					**05777440**
					SPO	
Actinomyces israelii		98.41	0.45	SUC 98	1	
Clostridium butyricum		1.59	-.07		99	
05777442	**Good identification**					**05777442**
Actinomyces israelii		100.0	0.48	SUC 98		
05777500	**Low discrimination**					**05777500**
					SPO	
Actinomyces israelii		69.57	0.62	SUC 98	1	
Clostridium butyricum		30.12	0.35		99	
Bifidobacterium adolescentis		0.31	0.15		1	
05777502	**Low discrimination**					**05777502**
					SPO	
Actinomyces israelii		71.13	0.65	SUC 98	1	
Bifidobacterium adolescentis		20.77	0.50		1	
Clostridium butyricum		7.98	0.28		99	
05777510	**Low discrimination**					**05777510**
					SPO	
Actinomyces israelii		72.89	0.48	SUC 98	1	
Bifidobacterium dentium		12.14	0.10	MLZ 13	1	
Clostridium butyricum		11.12	0.13		99	
05777512	**Low discrimination**					**05777512**
Bifidobacterium adolescentis		52.36	0.51	SUC 99		
Bifidobacterium dentium		24.61	0.27			
Actinomyces israelii		21.93	0.51			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
05777520	**Low discrimination**						**05777520**
					SPO		
Actinomyces israelii		62.90	0.42	SUC 98	1		
Clostridium butyricum		37.10	0.17	NAG 6	99		
05777522	**Low discrimination**						**05777522**
					GRA	SPO	
Actinomyces israelii		62.86	0.45	SUC 98	99	1	
Parabacteroides distasonis		21.66	0.33	NAG 6	1	1	
Clostridium butyricum		9.61	0.10		99	99	
05777540	**Low discrimination**						**05777540**
					COC	SPO	
Actinomyces israelii		76.37	0.55	SUC 98	1	1	
Blautia producta		16.24	0.24		99	1	
Clostridium butyricum		7.39	0.17		1	99	
05777542	**Low discrimination**						**05777542**
					SPO		
Actinomyces israelii		88.21	0.58	SUC 98	1		
Bifidobacterium adolescentis		8.18	0.34		1		
Clostridium butyricum		2.21	0.10		99		
05777552	**Low discrimination**						**05777552**
					COC		
Actinomyces israelii		47.52	0.44	SUC 98	1		
Bifidobacterium adolescentis		36.07	0.36	MLZ 13	1		
Blautia producta		15.03	0.15		99		
05777562	**Good identification**						**05777562**
Actinomyces israelii		100.0	0.38	SUC 98			
05777602	**Good identification**						**05777602**
Actinomyces israelii		100.0	0.35	SUC 98			
05777700	**Low discrimination**						**05777700**
					SPO		
Actinomyces israelii		84.90	0.43	SUC 98	1		
Clostridium butyricum		15.10	0.09	RHA 7	99		
05777702	**Low discrimination**						**05777702**
					SPO		
Actinomyces israelii		89.25	0.46	SUC 98	1		
Clostridium butyricum		4.12	0.02	RHA 7	99		
Bifidobacterium adolescentis		3.50	0.15		1		
05777722	**Low discrimination**						**05777722**
					GRA		
Parabacteroides distasonis		73.57	0.41	MAN 1	1		
Bacteroides ovatus		18.13	0.15		1		
Actinomyces israelii		5.36	0.25		99		
05777732	**Low discrimination**						**05777732**
Parabacteroides distasonis		57.84	0.36	MAN 1			
Clostridium clostridioforme		19.61	0.23				
Bacteroides ovatus		17.65	0.12				
05777742	**Low discrimination**						**05777742**
					COC		
Actinomyces israelii		82.70	0.39	SUC 98	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Blautia producta		17.30	0.07	RHA 7	99		
06000000	**Low discrimination**						**06000000**
					COC		
Parvimonas micra		30.62	0.50	SUC 1	99		
Atopobium minutum		17.89	0.65	MNS 10	1		
Anaerococcus prevotii		14.36	0.50		99		
06000001	**Very good identification**						**06000001**
					COC	GRA	
Anaerococcus prevotii		97.25	0.66	SUC 1	99	99	
Campylobacter ureolyticus		1.62	0.32		1	1	
Peptococcus niger		0.54	0.23		99	99	
06000002	**Low discrimination**						**06000002**
					COC		
Atopobium minutum		57.27	0.62	SUC 5	1		
Anaerococcus prevotii		17.24	0.39		99		
Parvimonas micra		16.34	0.33		99		
06000003	**Good identification**						**06000003**
					COC		
Anaerococcus prevotii		99.07	0.56	SUC 1	99		
Peptococcus niger		0.39	0.10		99		
Atopobium minutum		0.37	0.28		1		
06000020	**Low discrimination**						**06000020**
					COC		
Atopobium minutum		69.50	0.57	SUC 5	1		
Peptococcus niger		9.39	0.24		99		
Gemella morbillorum		7.41	0.35		99		
06000022	**Low discrimination**						**06000022**
					COC	GRA	
Atopobium minutum		70.98	0.54	SUC 5	1	99	
Pseudoflavonifractor capillosus		25.10	0.32		1	1	
Peptococcus niger		2.54	0.11		99	99	
06000400	**Low discrimination**						**06000400**
					COC		
Gemella morbillorum		40.24	0.35	GLU 99	99		
Parvimonas micra		19.58	0.15	ARA 10	99		
Atopobium minutum		11.44	0.31		1		
06001000	**Low discrimination**						**06001000**
					COC		
Atopobium minutum		60.31	0.57	SUC 5	1		
Gemella morbillorum		31.16	0.47		99		
Parvimonas micra		3.13	0.15		99		
06001002	**Good identification**						**06001002**
					COC		
Atopobium minutum		96.95	0.54	SUC 5	1		
Anaerococcus prevotii		0.88	0.05		99		
Parvimonas micra		0.84	-.01		99		
06001020	**Low discrimination**						**06001020**
					COC	CAT	
Atopobium minutum		84.64	0.48	SUC 5	1	1	
Gemella morbillorum		14.58	0.31		99	1	
Propionibacterium granulosum		0.44	0.11		1	90	
06001022	**Low discrimination**						**06001022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Atopobium minutum		89.45	0.45	SUC 5	99		
Pseudoflavonifractor capillosus		10.55	0.15		1		
06004000	**Low discrimination**						**06004000**
					COC		
Gemella morbillorum		39.57	0.35	GLU 99	99		
Parvimonas micra		19.25	0.15	XYL 10	99		
Atopobium minutum		11.25	0.31		1		
06010000	**Low discrimination**						**06010000**
					COC	GRA	
Atopobium minutum		36.87	0.64	SUC 5	1	99	
Fusobacterium varium		33.27	0.50		1	1	
Gemella morbillorum		15.62	0.53		99	99	
06010002	**Low discrimination**						**06010002**
					GRA	CAT	
Atopobium minutum		88.69	0.61	SUC 5	99	1	
Propionibacterium granulosum		8.16	0.46		99	90	
Fusobacterium varium		1.21	0.15		1	1	
06010010	**Low discrimination**						**06010010**
					GRA	CAT	
Propionibacterium granulosum		56.85	0.41	GLU 99	99	90	
Atopobium minutum		17.16	0.30	MLZ 12	99	1	
Fusobacterium varium		15.49	0.15		1	1	
06010020	**Low discrimination**						**06010020**
					COC	CAT	
Atopobium minutum		69.37	0.56	SUC 5	1	1	
Propionibacterium granulosum		12.77	0.46		1	90	
Gemella morbillorum		9.79	0.37		99	1	
06010022	**Low discrimination**						**06010022**
					GRA	CAT	
Atopobium minutum		89.79	0.53	SUC 5	99	1	
Propionibacterium granulosum		6.20	0.35		99	90	
Pseudoflavonifractor capillosus		3.82	0.15		1	1	
06010100	**Low discrimination**						**06010100**
					GRA		
Fusobacterium mortiferum		48.97	0.50	GLU 99	1		
Actinomyces meyeri		33.17	0.38		99		
Atopobium minutum		6.95	0.30		99		
06010400	**Low discrimination**						**06010400**
					COC	GRA	
Gemella morbillorum		65.57	0.37	GLU 99	99	99	
Atopobium minutum		14.08	0.30	ARA 10	1	99	
Fusobacterium varium		12.70	0.15		1	1	
06011000	**Low discrimination**						**06011000**
					COC	CAT	
Atopobium minutum		36.16	0.56	SUC 5	1	1	
Propionibacterium granulosum		26.62	0.56		1	90	
Gemella morbillorum		24.74	0.48		99	1	
06011002	**Low discrimination**						**06011002**
					CAT		
Atopobium minutum		71.22	0.53	SUC 5	1		
Propionibacterium granulosum		19.66	0.46		90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		6.57	0.42		1			
06011010	**Low discrimination**							**06011010**
					COC	CAT		
Propionibacterium granulosum		84.51	0.41	GLU 99	1	90		
Atopobium minutum		8.50	0.22	MLZ 12	1	1		
Gemella morbillorum		5.82	0.14		99	1		
06011020	**Low discrimination**							**06011020**
					COC	CAT		
Atopobium minutum		52.74	0.48	SUC 5	1	1		
Propionibacterium granulosum		29.12	0.46		1	90		
Gemella morbillorum		12.03	0.32		99	1		
06011022	**Low discrimination**							**06011022**
					CAT			
Atopobium minutum		75.44	0.45	SUC 5	1			
Propionibacterium granulosum		15.62	0.35		90			
Lactobacillus fermentum		6.96	0.34		1			
06011100	**Low discrimination**							**06011100**
					GRA			
Fusobacterium mortiferum		48.11	0.53	GLU 99	1			
Actinomyces meyeri		32.59	0.41		99			
Lactobacillus fermentum		15.07	0.46		99			
06011102	**Low discrimination**							**06011102**
					GRA			
Lactobacillus fermentum		66.35	0.46	GLU 99	99			
Fusobacterium mortiferum		23.53	0.37		1			
Atopobium minutum		4.45	0.19		99			
06011402	**Low discrimination**							**06011402**
					CAT			
Lactobacillus fermentum		87.74	0.42	GLU 99	1			
Atopobium minutum		9.61	0.19		1			
Propionibacterium granulosum		2.65	0.11		90			
06011500	**Low discrimination**							**06011500**
					GRA			
Lactobacillus fermentum		63.58	0.46	GLU 99	99			
Actinomyces meyeri		34.37	0.31		99			
Fusobacterium mortiferum		2.05	0.19		1			
06011502	**Good identification**							**06011502**
Lactobacillus fermentum		98.36	0.46	GLU 99				
Actinomyces meyeri		1.64	0.05					
06011520	**Low discrimination**							**06011520**
					GRA			
Lactobacillus fermentum		77.32	0.38	GLU 99	99			
Actinomyces meyeri		17.10	0.16		99			
Prevotella melaninogenica		5.58	-.01		1			
06011522	**Good identification**							**06011522**
Lactobacillus fermentum		100.0	0.38	GLU 99				
06012000	**Low discrimination**							**06012000**
					COC	CAT	GRA	
Propionibacterium granulosum		68.74	0.54	GLU 99	1	90	99	
Eubacterium limosum		16.11	0.29		1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		4.95	0.29		99	1	1	
06012002	**Low discrimination**							**06012002**
					COC	CAT	GRA	
Propionibacterium granulosum		84.25	0.44	GLU 99	1	90	99	
Atopobium minutum		12.26	0.27		1	1	99	
Megasphaera elsdenii		2.70	0.12		99	1	1	
06012010	**Low discrimination**							**06012010**
					SPO	CAT		
Propionibacterium granulosum		58.98	0.39	GLU 99	1	90		
Clostridium difficile		39.99	0.24	MLZ 12	99	1		
Eubacterium limosum		1.02	-.06		1	1		
06012020	**Acceptable identification**							**06012020**
					COC	CAT		
Propionibacterium granulosum		91.53	0.44	GLU 99	1	90		
Atopobium minutum		6.66	0.22		1	1		
Gemella morbillorum		0.94	0.03		99	1		
06013000	**Good identification**							**06013000**
					COC	CAT		
Propionibacterium granulosum		96.57	0.54	GLU 99	1	90		
Atopobium minutum		1.76	0.22		1	1		
Gemella morbillorum		1.20	0.14		99	1		
06013002	**Good identification**							**06013002**
					CAT			
Propionibacterium granulosum		95.38	0.44	GLU 99	90			
Atopobium minutum		4.63	0.19		1			
06013010	**Good identification**							**06013010**
Propionibacterium granulosum		100.0	0.39	GLU 99				
06013020	**Good identification**							**06013020**
					CAT			
Propionibacterium granulosum		97.63	0.44	GLU 99	90			
Atopobium minutum		2.37	0.14		1			
06014000	**Low discrimination**							**06014000**
					COC	GRA		
Gemella morbillorum		38.28	0.37	GLU 99	99	99		
Fusobacterium varium		30.59	0.26	XYL 10	1	1		
Actinomyces meyeri		19.04	0.32		1	99		
06014100	**Acceptable identification**							**06014100**
					COC	GRA		
Actinomyces meyeri		93.78	0.43	GLU 93	1	99		
Fusobacterium mortiferum		5.37	0.31	MNS 1	1	1		
Gemella morbillorum		0.48	0.03		99	99		
06015000	**Low discrimination**							**06015000**
					COC			
Actinomyces meyeri		42.73	0.35	GLU 93	1			
Gemella morbillorum		30.84	0.32	MNS 1	99			
Lactobacillus fermentum		14.66	0.38		1			
06015002	**Low discrimination**							**06015002**
Lactobacillus fermentum		73.54	0.38	GLU 99				
Atopobium minutum		13.71	0.19					
Actinomyces meyeri		6.63	0.10					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06015100	**Low discrimination**						**06015100**
					GRA		
Actinomyces meyeri		83.53	0.46	GLU 93	99		
Lactobacillus fermentum		11.69	0.42	MNS 1	99		
Fusobacterium mortiferum		4.78	0.34		1		
06015102	**Low discrimination**						**06015102**
					GRA		
Lactobacillus fermentum		78.96	0.42	GLU 99	99		
Actinomyces meyeri		17.45	0.20		99		
Fusobacterium mortiferum		3.59	0.17		1		
06015400	**Low discrimination**						**06015400**
					COC		
Lactobacillus fermentum		50.95	0.38	GLU 99	1		
Actinomyces meyeri		37.13	0.25		1		
Gemella morbillorum		11.91	0.16		99		
06015402	**Good identification**						**06015402**
Lactobacillus fermentum		100.0	0.38	GLU 99			
06015500	**Low discrimination**						**06015500**
Actinomyces meyeri		64.12	0.35	GLU 93			
Lactobacillus fermentum		35.88	0.42	MNS 1			
06015502	**Acceptable identification**						**06015502**
Lactobacillus fermentum		94.76	0.42	GLU 99			
Actinomyces meyeri		5.24	0.10				
06030000	**Very good identification**						**06030000**
					GRA	CAT	
Propionibacterium granulosum		95.46	0.64	GLU 99	99	90	
Atopobium minutum		1.31	0.30		99	1	
Fusobacterium mortiferum		1.18	0.35		1	5	
06030002	**Good identification**						**06030002**
					GRA	CAT	
Propionibacterium granulosum		95.96	0.54	GLU 99	99	90	
Atopobium minutum		3.51	0.27		99	1	
Fusobacterium mortiferum		0.53	0.18		1	5	
06030010	**Good identification**						**06030010**
Propionibacterium granulosum		100.0	0.49	GLU 99			
06030012	**Good identification**						**06030012**
Propionibacterium granulosum		100.0	0.39	GLU 99			
06030020	**Good identification**						**06030020**
					SPO	CAT	
Propionibacterium granulosum		97.22	0.54	GLU 99	1	90	
Atopobium minutum		1.78	0.22		1	1	
Clostridium cadaveris		0.75	0.05		99	1	
06030022	**Good identification**						**06030022**
					CAT		
Propionibacterium granulosum		95.35	0.43	GLU 99	90		
Atopobium minutum		4.65	0.19		1		
06030030	**Good identification**						**06030030**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.39	GLU 99				
06030100	**Low discrimination**							**06030100**
					GRA	CAT		
Fusobacterium mortiferum		72.85	0.41	GLU 99	1	5		
Propionibacterium granulosum		25.48	0.30		99	90		
Actinomyces meyeri		1.67	0.04		99	1		
06030200	**Good identification**							**06030200**
Propionibacterium granulosum		100.0	0.38	GLU 99				
06031000	**Very good identification**							**06031000**
					GRA	CAT		
Propionibacterium granulosum		97.44	0.64	GLU 99	99	90		
Fusobacterium mortiferum		1.81	0.38		1	5		
Atopobium minutum		0.45	0.22		99	1		
06031002	**Good identification**							**06031002**
					GRA	CAT		
Propionibacterium granulosum		98.00	0.54	GLU 99	99	90		
Atopobium minutum		1.20	0.19		99	1		
Fusobacterium mortiferum		0.81	0.21		1	5		
06031010	**Good identification**							**06031010**
Propionibacterium granulosum		100.0	0.49	GLU 99				
06031012	**Good identification**							**06031012**
Propionibacterium granulosum		100.0	0.39	GLU 99				
06031020	**Good identification**							**06031020**
					CAT			
Propionibacterium granulosum		97.54	0.54	GLU 99	90			
Clostridium perfringens		1.86	0.17		1			
Atopobium minutum		0.60	0.14		1			
06031022	**Good identification**							**06031022**
					CAT			
Propionibacterium granulosum		95.04	0.43	GLU 99	90			
Clostridium perfringens		3.42	0.11		1			
Atopobium minutum		1.55	0.11		1			
06031030	**Good identification**							**06031030**
Propionibacterium granulosum		100.0	0.39	GLU 99				
06031100	**Low discrimination**							**06031100**
					COC	GRA	CAT	
Fusobacterium mortiferum		74.56	0.44	GLU 99	1	1	5	
Propionibacterium granulosum		17.38	0.30		1	99	90	
Atopobium parvulum		4.49	0.12		99	99	1	
06031200	**Good identification**							**06031200**
Propionibacterium granulosum		100.0	0.38	GLU 99				
06032000	**Very good identification**							**06032000**
					CAT			
Propionibacterium granulosum		99.92	0.62	GLU 99	90			
Eubacterium limosum		0.08	-.06		1			
06032002	**Good identification**							**06032002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.52	GLU 99			
06032010	**Good identification**						**06032010**
					SPO	CAT	
Propionibacterium granulosum		98.33	0.47	GLU 99	1	90	
Clostridium difficile		1.67	0.04	MLZ 12	99	1	
06032012	**Good identification**						**06032012**
Propionibacterium granulosum		100.0	0.37	GLU 99			
06032020	**Good identification**						**06032020**
Propionibacterium granulosum		100.0	0.52	GLU 99			
06032022	**Good identification**						**06032022**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06032030	**Good identification**						**06032030**
Propionibacterium granulosum		100.0	0.37	GLU 99			
06032200	**Good identification**						**06032200**
Propionibacterium granulosum		100.0	0.36	GLU 99			
06033000	**Very good identification**						**06033000**
Propionibacterium granulosum		100.0	0.62	GLU 99			
06033002	**Good identification**						**06033002**
Propionibacterium granulosum		100.0	0.52	GLU 99			
06033010	**Good identification**						**06033010**
Propionibacterium granulosum		100.0	0.47	GLU 99			
06033012	**Good identification**						**06033012**
Propionibacterium granulosum		100.0	0.37	GLU 99			
06033020	**Good identification**						**06033020**
Propionibacterium granulosum		100.0	0.52	GLU 99			
06033022	**Good identification**						**06033022**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06033030	**Good identification**						**06033030**
Propionibacterium granulosum		100.0	0.37	GLU 99			
06033200	**Good identification**						**06033200**
Propionibacterium granulosum		100.0	0.36	GLU 99			
06050000	**Low discrimination**						**06050000**
					GRA		
Fusobacterium mortiferum		28.14	0.36	GLU 99	1		
Atopobium minutum		27.95	0.30		99		
Fusobacterium varium		25.22	0.15		1		
06050100	**Good identification**						**06050100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		97.99	0.42	GLU 99	1		
Actinomyces meyeri		2.01	0.04		99		
06051000	**Low discrimination**						**06051000**
					GRA	CAT	
Fusobacterium mortiferum		55.99	0.39	GLU 99	1	5	
Atopobium minutum		12.36	0.22		99	1	
Propionibacterium granulosum		9.10	0.22		99	90	
06051100	**Acceptable identification**						**06051100**
					GRA		
Fusobacterium mortiferum		91.51	0.45	GLU 99	1		
Lactobacillus catenaformis		6.62	0.15		99		
Actinomyces meyeri		1.88	0.07		99		
06071100	**Low discrimination**						**06071100**
					COC	GRA	
Fusobacterium mortiferum		70.37	0.36	GLU 99	1	1	
Atopobium parvulum		29.63	0.18		99	99	
06100000	**Low discrimination**						**06100000**
					COC	SPO	
Gemella morbillorum		82.24	0.64	GLU 99	99	1	
Peptostreptococcus anaerobius		6.28	0.38		99	1	
Clostridium botulinum biovar A		4.06	0.39		1	99	
06100020	**Acceptable identification**						**06100020**
					COC	SPO	
Gemella morbillorum		94.14	0.48	GLU 99	99	1	
Atopobium minutum		1.57	0.22	NAG 10	1	1	
Clostridium botulinum biovar B		0.92	0.08		1	99	
06100040	**Low discrimination**						**06100040**
					COC	SPO	
Clostridium botulinum biovar B		89.11	0.47	GLU 99	1	99	
Clostridium botulinum biovar A		6.01	0.30		1	99	
Gemella morbillorum		4.12	0.30		99	1	
06100050	**Good identification**						**06100050**
Clostridium botulinum biovar B		100.0	0.42	GLU 99			
06100060	**Good identification**						**06100060**
					COC	SPO	
Clostridium botulinum biovar B		98.97	0.42	GLU 99	1	99	
Gemella morbillorum		1.03	0.14		99	1	
06100070	**Good identification**						**06100070**
Clostridium botulinum biovar B		100.0	0.37	GLU 99			
06100400	**Good identification**						**06100400**
					COC	SPO	
Gemella morbillorum		98.52	0.48	GLU 99	99	1	
Peptostreptococcus anaerobius		0.68	0.04	ARA 10	99	1	
Clostridium botulinum biovar A		0.44	0.05		1	99	
06101000	**Very good identification**						**06101000**
					COC	GRA	
Gemella morbillorum		97.12	0.60	GLU 99	99	99	
Prevotella intermedia		1.99	0.30		1	1	
Atopobium minutum		0.34	0.22		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
06101020	**Acceptable identification**							**06101020**
					COC	GRA		
Gemella morbillorum		90.76	0.43	GLU 99	99	99		
Prevotella melaninogenica		6.83	0.17	NAG 10	1	1		
Clostridium perfringens		1.46	0.12		1	99		
06101101	**Good identification**							**06101101**
Clostridium chauvoei		100.0	0.54	GLU 99				
06101103	**Good identification**							**06101103**
Clostridium chauvoei		100.0	0.41	GLU 99				
06101120	**Good identification**							**06101120**
					GRA			
Prevotella melaninogenica		95.06	0.51	GLU 90	1			
Prevotella bivia		3.05	0.19	RAF 90	1			
Clostridium perfringens		1.51	0.27	FRU 90	99			
06101121	**Good identification**							**06101121**
					GRA	SPO		
Clostridium chauvoei		95.29	0.45	GLU 99	99	99		
Prevotella melaninogenica		4.71	0.17		1	1		
06101122	**Good identification**							**06101122**
					GRA			
Prevotella melaninogenica		96.23	0.39	GLU 90	1			
Clostridium perfringens		3.77	0.22	RAF 90	99			
06101400	**Good identification**							**06101400**
					COC			
Gemella morbillorum		99.47	0.43	GLU 99	99			
Lactobacillus fermentum		0.53	0.16	ARA 10	1			
06104000	**Good identification**							**06104000**
					COC			
Gemella morbillorum		97.91	0.48	GLU 99	99			
Peptostreptococcus anaerobius		0.68	0.04	XYL 10	99			
Actinomyces meyeri		0.62	0.11		1			
06105000	**Good identification**							**06105000**
					COC			
Gemella morbillorum		97.96	0.43	GLU 99	99			
Actinomyces meyeri		1.74	0.14	XYL 10	1			
Lactobacillus fermentum		0.30	0.12		1			
06110000	**Low discrimination**							**06110000**
					COC	GRA	CAT	
Gemella morbillorum		49.84	0.66	GLU 99	99	99	1	
Prevotella intermedia		44.54	0.64		1	1	1	
Propionibacterium granulosum		2.29	0.50		1	99	90	
06110002	**Low discrimination**							**06110002**
					COC	GRA	CAT	
Propionibacterium granulosum		32.11	0.40	GLU 99	1	99	90	
Gemella morbillorum		28.24	0.32		99	99	1	
Prevotella intermedia		25.23	0.30		1	1	1	
06110020	**Low discrimination**							**06110020**
					COC	GRA	CAT	
Gemella morbillorum		79.71	0.50	GLU 99	99	99	1	
Propionibacterium granulosum		8.24	0.40	NAG 10	1	99	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		6.48	0.30		1	1	1	
06110040	**Very good identification**							**06110040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		96.46	0.58	GLU 99	1	99	99	
Gemella morbillorum		1.36	0.32		99	99	1	
Prevotella intermedia		1.22	0.30		1	1	1	
06110050	**Good identification**							**06110050**
Clostridium botulinum biovar B		100.0	0.52	GLU 99				
06110060	**Good identification**							**06110060**
					COC	SPO		
Clostridium botulinum biovar B		99.68	0.52	GLU 99	1	99		
Gemella morbillorum		0.32	0.15		99	1		
06110070	**Good identification**							**06110070**
Clostridium botulinum biovar B		100.0	0.47	GLU 99				
06110100	**Low discrimination**							**06110100**
					COC	GRA		
Actinomyces meyeri		47.87	0.39	GLU 93	1	99		
Gemella morbillorum		21.34	0.32	MNS 1	99	99		
Prevotella intermedia		19.07	0.30		1	1		
06110400	**Acceptable identification**							**06110400**
					COC	GRA		
Gemella morbillorum		90.63	0.50	GLU 99	99	99		
Prevotella intermedia		7.36	0.30	ARA 10	1	1		
Actinomyces meyeri		1.16	0.18		1	99		
06111000	**Low discrimination**							**06111000**
					COC	GRA	CAT	
Prevotella intermedia		58.79	0.64	GLU 90	1	1	1	
Gemella morbillorum		35.43	0.61		99	99	1	
Propionibacterium granulosum		3.02	0.50		1	99	90	
06111002	**Low discrimination**							**06111002**
					GRA	CAT		
Lactobacillus fermentum		46.54	0.50	GLU 99	99	1		
Propionibacterium granulosum		20.86	0.40		99	90		
Prevotella intermedia		16.39	0.30		1	1		
06111020	**Low discrimination**							**06111020**
					COC	GRA		
Clostridium perfringens		38.65	0.46	GLU 99	1	99		
Gemella morbillorum		29.62	0.45	LAC 88	99	99		
Prevotella melaninogenica		16.42	0.33		1	1		
06111022	**Low discrimination**							**06111022**
					GRA			
Clostridium perfringens		66.02	0.41	GLU 99	99			
Lactobacillus fermentum		15.35	0.42	LAC 88	99			
Prevotella melaninogenica		11.35	0.21		1			
06111100	**Low discrimination**							**06111100**
					GRA			
Lactobacillus fermentum		27.24	0.54	GLU 99	99			
Actinomyces meyeri		22.14	0.42		99			
Prevotella melaninogenica		21.61	0.33		1			
06111101	**Very good identification**							**06111101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium chauvoei		99.71	0.60	GLU 99	99		
Lactobacillus fermentum		0.08	0.20		1		
Actinomyces meyeri		0.07	0.08		1		
06111102	**Low discrimination**						**06111102**
					GRA		
Lactobacillus fermentum		69.45	0.54	GLU 99	99		
Clostridium perfringens		13.84	0.30		99		
Prevotella melaninogenica		10.49	0.21		1		
06111103	**Good identification**						**06111103**
					SPO		
Clostridium chauvoei		99.07	0.47	GLU 99	99		
Lactobacillus fermentum		0.45	0.20		1		
Actinomyces naeslundii		0.28	0.26		1		
06111120	**Low discrimination**						**06111120**
					GRA		
Prevotella melaninogenica		84.65	0.67	GLU 90	1		
Clostridium perfringens		14.76	0.61	RAF 90	99		
Lactobacillus fermentum		0.36	0.46		99		
06111121	**Low discrimination**						**06111121**
					GRA	SPO	
Clostridium chauvoei		81.61	0.52	GLU 99	99	99	
Prevotella melaninogenica		15.55	0.33		1	1	
Clostridium perfringens		2.71	0.27		99	84	
06111122	**Low discrimination**						**06111122**
					GRA		
Prevotella melaninogenica		68.39	0.55	GLU 90	1		
Clostridium perfringens		29.46	0.56	RAF 90	99		
Lactobacillus fermentum		1.52	0.46		99		
06111123	**Low discrimination**						**06111123**
					GRA	SPO	
Clostridium chauvoei		76.76	0.39	GLU 99	99	99	
Prevotella melaninogenica		15.79	0.21		1	1	
Clostridium perfringens		6.80	0.22		99	84	
06111160	**Low discrimination**						**06111160**
					GRA		
Clostridium perfringens		76.68	0.49	GLU 99	99		
Prevotella melaninogenica		23.33	0.33		1		
06111162	**Low discrimination**						**06111162**
					GRA		
Clostridium perfringens		89.04	0.43	GLU 99	99		
Prevotella melaninogenica		10.96	0.21		1		
06111400	**Low discrimination**						**06111400**
					COC	GRA	
Gemella morbillorum		61.63	0.45	GLU 99	99	99	
Lactobacillus fermentum		26.40	0.50	ARA 10	1	99	
Prevotella intermedia		9.30	0.30		1	1	
06111402	**Good identification**						**06111402**
					COC		
Lactobacillus fermentum		97.70	0.50	GLU 99	1		
Gemella morbillorum		2.30	0.11		99		
06111420	**Low discrimination**						**06111420**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC			
Lactobacillus fermentum		51.45	0.42	GLU 99	1			
Gemella morbillorum		40.04	0.29		99			
Clostridium perfringens		4.75	0.12		1			
06111422	**Good identification**							**06111422**
Lactobacillus fermentum		95.84	0.42	GLU 99				
Clostridium perfringens		4.16	0.07					
06111500	**Low discrimination**							**06111500**
Lactobacillus fermentum		80.06	0.54	GLU 99				
Actinomyces meyeri		16.27	0.31					
Bifidobacterium longum subsp. longum		1.53	0.18					
06111502	**Good identification**							**06111502**
Lactobacillus fermentum		97.52	0.54	GLU 99				
Bifidobacterium longum subsp. longum		1.87	0.18					
Actinomyces meyeri		0.61	0.06					
06111522	**Low discrimination**							**06111522**
					GRA			
Lactobacillus fermentum		52.81	0.46	GLU 99	99			
Prevotella melaninogenica		23.94	0.21		1			
Bifidobacterium longum subsp. longum		12.13	0.28		99			
06112000	**Low discrimination**							**06112000**
					COC	GRA	CAT	
Propionibacterium granulosum		55.20	0.48	GLU 99	1	99	90	
Gemella morbillorum		16.09	0.32		99	99	1	
Prevotella intermedia		14.38	0.30		1	1	1	
06112002	**Low discrimination**							**06112002**
					COC	CAT	GRA	
Propionibacterium granulosum		88.23	0.38	GLU 99	1	90	99	
Megasphaera elsdenii		9.43	0.15		99	1	1	
Propionibacterium propionicum		2.34	0.15		1	1	99	
06112020	**Low discrimination**							**06112020**
					COC	CAT		
Propionibacterium granulosum		88.53	0.38	GLU 99	1	90		
Gemella morbillorum		11.47	0.15		99	1		
06112100	**Good identification**							**06112100**
					CAT			
Propionibacterium propionicum		96.32	0.46	NIT 86	1			
Propionibacterium granulosum		2.22	0.14	GLU 99	90			
Actinomyces meyeri		1.45	0.05		1			
06112102	**Good identification**							**06112102**
Propionibacterium propionicum		100.0	0.49	NIT 86				
06112140	**Good identification**							**06112140**
Propionibacterium propionicum		100.0	0.46	NIT 86				
06112142	**Good identification**							**06112142**
Propionibacterium propionicum		100.0	0.49	NIT 86				
06113000	**Low discrimination**							**06113000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	CAT	
Propionibacterium granulosum		69.97	0.48	GLU 99	1	99	90	
Prevotella intermedia		18.22	0.30		1	1	1	
Gemella morbillorum		10.98	0.27		99	99	1	
06113002	**Acceptable identification**							**06113002**
					CAT			
Propionibacterium granulosum		94.66	0.38	GLU 99	90			
Lactobacillus fermentum		2.83	0.16		1			
Propionibacterium propionicum		2.51	0.15		1			
06113020	**Low discrimination**							**06113020**
					COC	CAT		
Propionibacterium granulosum		83.37	0.38	GLU 99	1	90		
Clostridium perfringens		7.59	0.12		1	1		
Gemella morbillorum		5.82	0.11		99	1		
06113100	**Low discrimination**							**06113100**
					CAT			
Propionibacterium propionicum		88.53	0.46	NIT 86	1			
Lactobacillus fermentum		2.47	0.20	GLU 99	1			
Propionibacterium granulosum		2.04	0.14		90			
06113102	**Good identification**							**06113102**
Propionibacterium propionicum		96.77	0.49	NIT 86				
Lactobacillus fermentum		1.80	0.20	GLU 99				
Actinomyces naeslundii		1.43	0.28					
06113140	**Good identification**							**06113140**
Propionibacterium propionicum		100.0	0.46	NIT 86				
06113142	**Good identification**							**06113142**
Propionibacterium propionicum		100.0	0.49	NIT 86				
06114000	**Low discrimination**							**06114000**
					COC	GRA		
Gemella morbillorum		84.03	0.50	GLU 99	99	99		
Actinomyces meyeri		8.32	0.33	XYL 10	1	99		
Prevotella intermedia		6.83	0.30		1	1		
06114100	**Good identification**							**06114100**
					COC	GRA		
Actinomyces meyeri		96.39	0.44	GLU 93	1	99		
Gemella morbillorum		2.46	0.15	MNS 1	99	99		
Fusobacterium mortiferum		0.61	0.15		1	1		
06115000	**Low discrimination**							**06115000**
					COC			
Gemella morbillorum		59.21	0.45	GLU 99	99			
Actinomyces meyeri		16.33	0.36	XYL 10	1			
Lactobacillus fermentum		14.90	0.46		1			
06115002	**Acceptable identification**							**06115002**
					COC			
Lactobacillus fermentum		93.11	0.46	GLU 99	1			
Gemella morbillorum		3.74	0.11		99			
Actinomyces meyeri		3.16	0.11		1			
06115022	**Acceptable identification**							**06115022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		93.11	0.38	GLU 99		
Clostridium perfringens		6.89	0.07			
06115100	**Low discrimination**					**06115100**
					COC	
Actinomyces meyeri		70.52	0.47	GLU 93	1	
Lactobacillus fermentum		26.25	0.50	MNS 1	1	
Atopobium parvulum		1.59	0.16		99	
06115102	**Acceptable identification**					**06115102**
					COC	
Lactobacillus fermentum		91.43	0.50	GLU 99	1	
Actinomyces meyeri		7.60	0.21		1	
Atopobium parvulum		0.98	0.03		99	
06115122	**Low discrimination**					**06115122**
					GRA	
Lactobacillus fermentum		43.36	0.42	GLU 99	99	
Prevotella melaninogenica		33.46	0.21		1	
Clostridium perfringens		14.41	0.22		99	
06115400	**Low discrimination**					**06115400**
					COC	
Lactobacillus fermentum		58.29	0.46	GLU 99	1	
Gemella morbillorum		25.74	0.29		99	
Actinomyces meyeri		15.97	0.26		1	
06115402	**Good identification**					**06115402**
Lactobacillus fermentum		100.0	0.46	GLU 99		
06115420	**Low discrimination**					**06115420**
					COC	
Lactobacillus fermentum		79.41	0.38	GLU 99	1	
Gemella morbillorum		11.69	0.12		99	
Actinomyces meyeri		8.90	0.11		1	
06115422	**Good identification**					**06115422**
Lactobacillus fermentum		100.0	0.38	GLU 99		
06115500	**Low discrimination**					**06115500**
Lactobacillus fermentum		56.52	0.50	GLU 99		
Actinomyces meyeri		37.96	0.36			
Bifidobacterium longum subsp. longum		5.53	0.26			
06115502	**Low discrimination**					**06115502**
Lactobacillus fermentum		87.40	0.50	GLU 99		
Bifidobacterium longum subsp. longum		8.55	0.26			
Bifidobacterium adolescentis		2.24	0.17			
06115520	**Low discrimination**					**06115520**
					GRA	
Bifidobacterium longum subsp. longum		37.43	0.36	GLU 99	99	
Lactobacillus fermentum		31.89	0.42	RAF 97	99	
Bacteroides vulgatus		20.62	0.17		1	
06115522	**Low discrimination**					**06115522**
Bifidobacterium longum subsp. longum		53.99	0.36	GLU 99		
Lactobacillus fermentum		46.01	0.42	RAF 97		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
06115530	**Good identification**							**06115530**
Bifidobacterium longum subsp. longum		100.0	0.40	GLU 99				
06115532	**Good identification**							**06115532**
Bifidobacterium longum subsp. longum		100.0	0.40	GLU 99				
06120040	**Good identification**							**06120040**
Clostridium botulinum biovar B		99.44	0.56	GLU 99				
Clostridium botulinum biovar A		0.56	0.20					
06120050	**Good identification**							**06120050**
Clostridium botulinum biovar B		100.0	0.50	GLU 99				
06120060	**Good identification**							**06120060**
Clostridium botulinum biovar B		100.0	0.50	GLU 99				
06120070	**Good identification**							**06120070**
Clostridium botulinum biovar B		100.0	0.45	GLU 99				
06121101	**Good identification**							**06121101**
Clostridium chauvoei		100.0	0.37	GLU 99				
06130000	**Low discrimination**							**06130000**
					COC	SPO	CAT	
Propionibacterium granulosum		76.55	0.58	GLU 99	1	1	90	
Clostridium botulinum biovar B		12.03	0.32		1	99	1	
Gemella morbillorum		5.61	0.32		99	1	1	
06130002	**Good identification**							**06130002**
					SPO	CAT		
Propionibacterium granulosum		99.37	0.48	GLU 99	1	90		
Clostridium botulinum biovar B		0.63	-.03		99	1		
06130010	**Low discrimination**							**06130010**
					SPO	CAT		
Propionibacterium granulosum		63.78	0.43	GLU 99	1	90		
Clostridium botulinum biovar B		36.22	0.26	MLZ 12	99	1		
06130020	**Low discrimination**							**06130020**
					CAT	SPO		
Propionibacterium granulosum		72.21	0.48	GLU 99	90	1		
Clostridium botulinum biovar B		22.37	0.26		1	99		
Clostridium perfringens		3.07	0.17		1	84		
06130022	**Acceptable identification**							**06130022**
					CAT			
Propionibacterium granulosum		92.59	0.38	GLU 99	90			
Clostridium perfringens		7.41	0.11		1			
06130040	**Very good identification**							**06130040**
					SPO	CAT		
Clostridium botulinum biovar B		99.93	0.66	GLU 99	99	1		
Propionibacterium granulosum		0.07	0.24		1	90		
Clostridium botulinum biovar A		0.01	-.01		99	1		
06130050	**Very good identification**							**06130050**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.61	GLU 99	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.02	0.09		1	90		
06130060	**Very good identification**							**06130060**
					CAT	SPO		
Clostridium botulinum biovar B		99.94	0.61	GLU 99	1	99		
Propionibacterium granulosum		0.03	0.14		90	1		
Clostridium perfringens		0.03	0.05		1	84		
06130070	**Good identification**							**06130070**
Clostridium botulinum biovar B		100.0	0.55	GLU 99				
06131000	**Low discrimination**							**06131000**
					COC	GRA	CAT	
Propionibacterium granulosum		85.39	0.58	GLU 99	1	99	90	
Prevotella intermedia		5.59	0.30		1	1	1	
Gemella morbillorum		3.37	0.27		99	99	1	
06131002	**Low discrimination**							**06131002**
					CAT			
Propionibacterium granulosum		89.27	0.48	GLU 99	90			
Clostridium perfringens		5.47	0.20		1			
Lactobacillus jensenii		4.59	0.19		1			
06131010	**Good identification**							**06131010**
					CAT			
Propionibacterium granulosum		95.50	0.43	GLU 99	90			
Lactobacillus jensenii		4.50	0.14	MLZ 12	1			
06131020	**Low discrimination**							**06131020**
					CAT			
Clostridium perfringens		80.08	0.51	GLU 99	1			
Propionibacterium granulosum		19.03	0.48	LAC 88	90			
Lactobacillus jensenii		0.37	0.12		1			
06131022	**Low discrimination**							**06131022**
					CAT			
Clostridium perfringens		88.28	0.45	GLU 99	1			
Propionibacterium granulosum		11.15	0.38	LAC 88	90			
Lactobacillus jensenii		0.57	0.09		1			
06131060	**Low discrimination**							**06131060**
					CAT	SPO		
Clostridium perfringens		71.48	0.39	GLU 99	1	84		
Clostridium botulinum biovar B		27.62	0.26	LAC 88	1	99		
Propionibacterium granulosum		0.90	0.14		90	1		
06131100	**Low discrimination**							**06131100**
					COC	GRA		
Atopobium parvulum		67.60	0.46	GLU 99	99	99		
Clostridium perfringens		28.08	0.40		1	99		
Fusobacterium mortiferum		1.40	0.28		1	1		
06131101	**Good identification**							**06131101**
					COC	SPO		
Clostridium chauvoei		97.26	0.44	GLU 99	1	99		
Atopobium parvulum		1.34	0.12	TRE 10	99	1		
Actinomyces naeslundii		0.85	0.31		1	1		
06131120	**Very good identification**							**06131120**
					COC	GRA		
Clostridium perfringens		95.60	0.66	GLU 99	1	99		
Prevotella melaninogenica		2.98	0.33		1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		1.26	0.33		99	99	
06131121	**Low discrimination**						**06131121**
					SPO		
Clostridium chauvoei		62.55	0.35	GLU 99	99		
Clostridium perfringens		34.74	0.32	TRE 10	84		
Actinomyces naeslundii		1.63	0.31		1		
06131122	**Very good identification**						**06131122**
					COC	GRA	
Clostridium perfringens		98.16	0.60	GLU 99	1	99	
Prevotella melaninogenica		1.24	0.21		1	1	
Atopobium parvulum		0.48	0.20		99	99	
06131160	**Good identification**						**06131160**
					GRA		
Clostridium perfringens		99.84	0.54	GLU 99	99		
Prevotella melaninogenica		0.17	-.01		1		
06131162	**Good identification**						**06131162**
Clostridium perfringens		100.0	0.48	GLU 99			
06132000	**Good identification**						**06132000**
					SPO	CAT	
Propionibacterium granulosum		99.79	0.56	GLU 99	1	90	
Clostridium botulinum biovar B		0.21	-.03		99	1	
06132002	**Good identification**						**06132002**
					CAT		
Propionibacterium granulosum		99.28	0.46	GLU 99	90		
Propionibacterium propionicum		0.72	0.14		1		
06132010	**Good identification**						**06132010**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06132020	**Good identification**						**06132020**
Propionibacterium granulosum		100.0	0.46	GLU 99			
06132022	**Good identification**						**06132022**
Propionibacterium granulosum		100.0	0.35	GLU 99			
06132100	**Acceptable identification**						**06132100**
					CAT		
Propionibacterium propionicum		92.20	0.45	NIT 86	1		
Propionibacterium granulosum		7.80	0.22	GLU 99	90		
06132102	**Good identification**						**06132102**
					CAT		
Propionibacterium propionicum		98.61	0.48	NIT 86	1		
Propionibacterium granulosum		1.39	0.12	GLU 99	90		
06132140	**Good identification**						**06132140**
					SPO		
Propionibacterium propionicum		98.27	0.45	NIT 86	1		
Clostridium botulinum biovar B		1.73	-.03	GLU 99	99		
06132142	**Good identification**						**06132142**
Propionibacterium propionicum		100.0	0.48	NIT 86			
06133000	**Good identification**						**06133000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		99.78	0.56	GLU 99			
Actinomyces naeslundii		0.22	0.25				
06133002	**Good identification**						**06133002**
					CAT		
Propionibacterium granulosum		98.42	0.46	GLU 99	90		
Actinomyces naeslundii		0.87	0.25		25		
Propionibacterium propionicum		0.71	0.14		1		
06133010	**Good identification**						**06133010**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06133020	**Acceptable identification**						**06133020**
					CAT		
Propionibacterium granulosum		93.88	0.46	GLU 99	90		
Clostridium perfringens		5.29	0.17		1		
Actinomyces naeslundii		0.83	0.25		25		
06133022	**Low discrimination**						**06133022**
					CAT		
Propionibacterium granulosum		87.62	0.35	GLU 99	90		
Clostridium perfringens		9.29	0.11		1		
Actinomyces naeslundii		3.09	0.25		25		
06133100	**Low discrimination**						**06133100**
					COC	CAT	
Propionibacterium propionicum		81.41	0.45	NIT 86	1	1	
Propionibacterium granulosum		6.89	0.22	GLU 99	1	90	
Atopobium parvulum		5.25	0.12		99	1	
06133102	**Acceptable identification**						**06133102**
					CAT		
Propionibacterium propionicum		94.56	0.48	NIT 86	1		
Actinomyces naeslundii		3.32	0.33	GLU 99	25		
Propionibacterium granulosum		1.33	0.12		90		
06133121	**Low discrimination**						**06133121**
					SPO		
Actinomyces naeslundii		83.37	0.36	GLU 99	1		
Clostridium chauvoei		16.63	0.01	RAF 92	99		
06133123	**Good identification**						**06133123**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06133140	**Good identification**						**06133140**
Propionibacterium propionicum		100.0	0.45	NIT 86			
06133142	**Good identification**						**06133142**
Propionibacterium propionicum		100.0	0.48	NIT 86			
06135100	**Good identification**						**06135100**
					COC		
Atopobium parvulum		99.16	0.50	GLU 99	99		
Actinomyces meyeri		0.45	0.13		1		
Clostridium perfringens		0.22	0.06		1		
06135102	**Good identification**						**06135102**
					COC		
Atopobium parvulum		98.47	0.38	GLU 99	99		
Lactobacillus fermentum		0.94	0.16		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.59	0.00		1		
06135120	**Low discrimination**						**06135120**
					COC	GRA	
Atopobium parvulum		70.09	0.38	GLU 99	99	99	
Clostridium perfringens		29.01	0.32		1	99	
Prevotella melaninogenica		0.91	-.01		1	1	
06151000	**Low discrimination**						**06151000**
					GRA		
Lactobacillus jensenii		42.90	0.36	GLU 99	99		
Prevotella intermedia		23.00	0.30	SAL 95	1		
Lactobacillus catenaformis		16.40	0.29		99		
06151120	**Acceptable identification**						**06151120**
					GRA		
Prevotella melaninogenica		92.43	0.51	GLU 90	1		
Clostridium perfringens		6.04	0.38	RAF 90	99		
Lactobacillus catenaformis		0.61	0.21	CEL 10	99		
06151122	**Low discrimination**						**06151122**
					GRA		
Prevotella melaninogenica		65.87	0.39	GLU 90	1		
Prevotella oralis		13.72	0.31	RAF 90	1		
Clostridium perfringens		10.64	0.32	CEL 10	99		
06171000	**Acceptable identification**						**06171000**
					COC	CAT	
Lactobacillus jensenii		92.94	0.44	GLU 99	1	1	
Atopobium parvulum		3.43	0.18	SAL 95	99	1	
Propionibacterium granulosum		2.56	0.24		1	90	
06171002	**Acceptable identification**						**06171002**
					CAT		
Lactobacillus jensenii		94.18	0.41	GLU 99	1		
Lactobacillus acidophilus		3.48	0.14	SAL 95	1		
Propionibacterium granulosum		0.97	0.14		90		
06171010	**Good identification**						**06171010**
					CAT		
Lactobacillus jensenii		98.89	0.36	GLU 99	1		
Propionibacterium granulosum		1.12	0.09	SAL 95	90		
06171100	**Good identification**						**06171100**
					COC	GRA	
Atopobium parvulum		98.47	0.52	GLU 99	99	99	
Clostridium perfringens		0.73	0.16		1	99	
Fusobacterium mortiferum		0.29	0.20		1	1	
06171102	**Low discrimination**						**06171102**
					COC		
Atopobium parvulum		67.42	0.39	GLU 99	99		
Lactobacillus acidophilus		29.62	0.32		1		
Clostridium perfringens		1.33	0.11		1		
06171120	**Low discrimination**						**06171120**
					COC	GRA	
Clostridium perfringens		52.82	0.42	GLU 99	1	99	
Atopobium parvulum		38.87	0.39	CEL 4	99	99	
Prevotella melaninogenica		4.39	0.17		1	1	
06171122	**Low discrimination**						**06171122**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		47.78	0.37	GLU 99	1	
Lactobacillus acidophilus		32.83	0.32	CEL 4	1	
Atopobium parvulum		13.18	0.26		99	
06173101	**Good identification**					**06173101**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06173102	**Low discrimination**					**06173102**
					SPO	
Actinomyces naeslundii		29.74	0.35	GLU 99	1	
Lactobacillus acidophilus		25.91	0.10	RAF 92	1	
Clostridium tertium		19.79	0.09		99	
06173103	**Good identification**					**06173103**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06173120	**Low discrimination**					**06173120**
					SPO	
Actinomyces naeslundii		31.34	0.35	GLU 99	1	
Clostridium tertium		20.86	0.09	RAF 92	99	
Clostridium ramosum		19.66	0.07		92	
06173121	**Good identification**					**06173121**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06173123	**Good identification**					**06173123**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06175100	**Very good identification**					**06175100**
					COC	
Atopobium parvulum		99.92	0.57	GLU 99	99	
Actinomyces israelii		0.08	0.17		1	
06175102	**Good identification**					**06175102**
					COC	
Atopobium parvulum		99.34	0.44	GLU 99	99	
Actinomyces israelii		0.66	0.20		1	
06175120	**Good identification**					**06175120**
					COC	
Atopobium parvulum		99.27	0.44	GLU 99	99	
Clostridium perfringens		0.73	0.08		1	
06175712	**Good identification**					**06175712**
Clostridium clostridioforme		100.0	0.41	GLU 88		
06175732	**Good identification**					**06175732**
Clostridium clostridioforme		99.06	0.41	GLU 88		
Parabacteroides distasonis		0.94	0.12	SAL 88		
06210000	**Low discrimination**					**06210000**
					GRA	
Fusobacterium mortiferum		69.52	0.49	GLU 99	1	
Atopobium minutum		11.86	0.30		99	
Fusobacterium varium		10.70	0.15		1	
06210100	**Good identification**					**06210100**
					GRA	
Fusobacterium mortiferum		99.65	0.55	GLU 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		0.35	0.04		99		
06210102	**Good identification**						**06210102**
Fusobacterium mortiferum		100.0	0.39	GLU 99			
06211000	**Low discrimination**						**06211000**
					GRA	CAT	
Fusobacterium mortiferum		88.11	0.52	GLU 99	1	5	
Atopobium minutum		3.34	0.22		99	1	
Propionibacterium granulosum		2.46	0.22		99	90	
06211002	**Low discrimination**						**06211002**
					GRA		
Fusobacterium mortiferum		62.20	0.35	GLU 99	1		
Atopobium minutum		14.15	0.19	BGL 10	99		
Lactobacillus catenaformis		10.81	0.12		99		
06211100	**Very good identification**						**06211100**
					GRA		
Fusobacterium mortiferum		98.43	0.58	GLU 99	1		
Lactobacillus catenaformis		1.22	0.15		99		
Actinomyces meyeri		0.35	0.07		99		
06211102	**Low discrimination**						**06211102**
					GRA		
Fusobacterium mortiferum		86.14	0.42	GLU 99	1		
Lactobacillus catenaformis		9.63	0.15	BGL 10	99		
Lactobacillus acidophilus		2.45	0.02		99		
06214100	**Acceptable identification**						**06214100**
					GRA		
Fusobacterium mortiferum		91.67	0.36	GLU 99	1		
Actinomyces meyeri		8.33	0.09	XYL 7	99		
06215100	**Acceptable identification**						**06215100**
					GRA		
Fusobacterium mortiferum		91.67	0.39	GLU 99	1		
Actinomyces meyeri		8.33	0.12	XYL 7	99		
06230000	**Low discrimination**						**06230000**
					GRA	CAT	
Fusobacterium mortiferum		70.40	0.40	GLU 99	1	5	
Propionibacterium granulosum		29.60	0.30		99	90	
06230100	**Good identification**						**06230100**
Fusobacterium mortiferum		100.0	0.46	GLU 99			
06231000	**Low discrimination**						**06231000**
					GRA	CAT	
Fusobacterium mortiferum		65.76	0.43	GLU 99	1	5	
Propionibacterium granulosum		18.43	0.30		99	90	
Lactobacillus jensenii		15.81	0.22		99	1	
06231100	**Good identification**						**06231100**
					COC	GRA	
Fusobacterium mortiferum		96.05	0.49	GLU 99	1	1	
Atopobium parvulum		3.95	0.14		99	99	
06243000	**Good identification**						**06243000**
Clostridium innocuum		100.0	0.44	GLU 99			
06243002	**Good identification**						**06243002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.44	GLU 99			
06250000	**Low discrimination**						**06250000**
					GRA	SPO	
Fusobacterium mortiferum		83.66	0.40	GLU 99	1	1	
Clostridium innocuum		7.33	0.10		99	99	
Lactobacillus catenaformis		7.27	0.12		99	1	
06250100	**Acceptable identification**						**06250100**
					GRA	SPO	
Fusobacterium mortiferum		93.12	0.47	GLU 99	1	1	
Lactobacillus catenaformis		5.20	0.15		99	1	
Clostridium barati		1.68	0.02		99	99	
06251000	**Low discrimination**						**06251000**
					SPO		
Lactobacillus catenaformis		61.10	0.47	GLU 99	1		
Lactobacillus jensenii		14.53	0.36	MLT 90	1		
Clostridium innocuum		11.83	0.32		99		
06251002	**Low discrimination**						**06251002**
					SPO		
Lactobacillus catenaformis		66.79	0.47	GLU 99	1		
Clostridium innocuum		12.93	0.32	MLT 90	99		
Lactobacillus jensenii		10.59	0.33		1		
06251020	**Low discrimination**						**06251020**
					SPO		
Lactobacillus catenaformis		62.17	0.36	GLU 99	1		
Clostridium barati		23.05	0.24	MLT 90	99		
Lactobacillus jensenii		14.79	0.26		1		
06251100	**Low discrimination**						**06251100**
					GRA	SPO	
Lactobacillus catenaformis		74.42	0.50	GLU 99	99	1	
Fusobacterium mortiferum		20.18	0.50	MLT 90	1	1	
Clostridium barati		4.60	0.24		99	99	
06251102	**Low discrimination**						**06251102**
					SPO		
Lactobacillus catenaformis		68.02	0.50	GLU 99	1		
Lactobacillus acidophilus		17.32	0.36	MLT 90	1		
Clostridium barati		12.61	0.32		99		
06251120	**Low discrimination**						**06251120**
					SPO		
Lactobacillus catenaformis		55.97	0.39	GLU 99	1		
Clostridium barati		41.49	0.32	MLT 90	99		
Clostridium paraputrificum		0.84	-.03		99		
06251122	**Low discrimination**						**06251122**
					SPO		
Clostridium barati		52.22	0.40	ESL 87	99		
Lactobacillus acidophilus		23.92	0.36	GLU 99	1		
Lactobacillus catenaformis		23.48	0.39		1		
06252000	**Good identification**						**06252000**
Clostridium innocuum		100.0	0.44	GLU 99			
06252002	**Good identification**						**06252002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.44	GLU 99			
06253000	**Very good identification**						**06253000**
					SPO		
Clostridium innocuum		99.92	0.66	GLU 99	99		
Lactobacillus catenaformis		0.05	0.12		1		
Lactobacillus jensenii		0.01	0.02		1		
06253002	**Very good identification**						**06253002**
					SPO		
Clostridium innocuum		99.91	0.66	GLU 99	99		
Lactobacillus catenaformis		0.05	0.12		1		
Clostridium barati		0.04	0.05		99		
06253010	**Good identification**						**06253010**
Clostridium innocuum		100.0	0.42	GLU 99			
06253012	**Good identification**						**06253012**
Clostridium innocuum		100.0	0.42	GLU 99			
06253400	**Good identification**						**06253400**
Clostridium innocuum		100.0	0.53	GLU 99			
06253402	**Good identification**						**06253402**
Clostridium innocuum		100.0	0.53	GLU 99			
06257000	**Good identification**						**06257000**
Clostridium innocuum		100.0	0.44	GLU 99			
06257002	**Good identification**						**06257002**
Clostridium innocuum		100.0	0.44	GLU 99			
06263000	**Good identification**						**06263000**
Clostridium innocuum		100.0	0.36	GLU 99			
06263002	**Good identification**						**06263002**
Clostridium innocuum		100.0	0.36	GLU 99			
06270100	**Good identification**						**06270100**
Fusobacterium mortiferum		100.0	0.38	GLU 99			
06271000	**Low discrimination**						**06271000**
					GRA	SPO	
Lactobacillus jensenii		84.30	0.44	GLU 99	99	1	
Clostridium innocuum		8.04	0.24	MLT 95	99	99	
Fusobacterium mortiferum		6.15	0.34		1	1	
06271002	**Low discrimination**						**06271002**
					SPO		
Lactobacillus jensenii		70.07	0.41	GLU 99	1		
Lactobacillus acidophilus		16.38	0.27	MLT 95	1		
Clostridium innocuum		10.02	0.24		99		
06271010	**Good identification**						**06271010**
					SPO		
Lactobacillus jensenii		98.82	0.36	GLU 99	1		
Clostridium innocuum		1.18	0.00	MLT 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06271100	**Low discrimination**						**06271100**
					COC	GRA	
Fusobacterium mortiferum		60.82	0.41	GLU 99	1	1	
Atopobium parvulum		17.49	0.20		99	99	
Lactobacillus catenaformis		7.58	0.15		1	99	
06271102	**Good identification**						**06271102**
					SPO		
Lactobacillus acidophilus		95.25	0.45	GLU 99	1		
Clostridium barati		1.80	0.14	MLT 94	99		
Lactobacillus catenaformis		1.13	0.15		1		
06271122	**Acceptable identification**						**06271122**
					SPO		
Lactobacillus acidophilus		93.99	0.45	GLU 99	1		
Clostridium barati		5.33	0.22	MLT 94	99		
Clostridium ramosum		0.40	0.04		92		
06272000	**Good identification**						**06272000**
Clostridium innocuum		100.0	0.36	GLU 99			
06272002	**Good identification**						**06272002**
Clostridium innocuum		100.0	0.36	GLU 99			
06273000	**Very good identification**						**06273000**
					SPO		
Clostridium innocuum		99.89	0.58	GLU 99	99		
Lactobacillus jensenii		0.11	0.10		1		
06273002	**Very good identification**						**06273002**
					SPO		
Clostridium innocuum		99.84	0.58	GLU 99	99		
Lactobacillus acidophilus		0.09	0.05		1		
Lactobacillus jensenii		0.07	0.07		1		
06273400	**Good identification**						**06273400**
Clostridium innocuum		100.0	0.45	GLU 99			
06273402	**Good identification**						**06273402**
Clostridium innocuum		100.0	0.45	GLU 99			
06277000	**Good identification**						**06277000**
Clostridium innocuum		100.0	0.36	GLU 99			
06277002	**Good identification**						**06277002**
Clostridium innocuum		100.0	0.36	GLU 99			
06310100	**Good identification**						**06310100**
					GRA		
Fusobacterium mortiferum		96.89	0.39	GLU 99	1		
Actinomyces meyeri		3.12	0.05	MLT 11	99		
06311000	**Low discrimination**						**06311000**
					GRA		
Lactobacillus jensenii		38.23	0.36	GLU 99	99		
Prevotella intermedia		20.50	0.30	CEL 95	1		
Lactobacillus catenaformis		14.61	0.29		99		
06311020	**Low discrimination**						**06311020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		72.79	0.36	GLU 99		
Lactobacillus jensenii		15.70	0.26	LAC 88		
Lactobacillus catenaformis		6.00	0.18			
06311100	**Low discrimination**					**06311100**
					GRA	
Fusobacterium mortiferum		37.15	0.42	GLU 99	1	
Lactobacillus catenaformis		33.60	0.32	MLT 11	99	
Clostridium perfringens		15.41	0.25		99	
06311120	**Low discrimination**					**06311120**
					GRA	
Clostridium perfringens		78.44	0.51	GLU 99	99	
Prevotella melaninogenica		18.18	0.33		1	
Lactobacillus catenaformis		1.32	0.21		99	
06311122	**Low discrimination**					**06311122**
					GRA	
Clostridium perfringens		72.63	0.45	GLU 99	99	
Prevotella oralis		7.83	0.25		1	
Prevotella melaninogenica		6.81	0.21		1	
06311123	**Low discrimination**					**06311123**
					SPO	
Actinomyces naeslundii		48.19	0.36	GLU 99	1	
Clostridium perfringens		35.59	0.11	RAF 92	84	
Clostridium chauvoei		16.23	0.05		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06311160	**Good identification**						**06311160**
					GRA		
Clostridium perfringens		98.79	0.39	GLU 99	99		
Prevotella melaninogenica		1.21	-.01		1		
06313100	**Low discrimination**						**06313100**
Actinomyces naeslundii		88.62	0.38	GLU 99			
Propionibacterium propionicum		11.38	0.12	RAF 92			
06313101	**Good identification**						**06313101**
Actinomyces naeslundii		100.0	0.41	GLU 99			
06313102	**Low discrimination**						**06313102**
Actinomyces naeslundii		72.38	0.38	GLU 99			
Propionibacterium propionicum		13.95	0.15	RAF 92			
Lactobacillus acidophilus		13.67	0.01				
06313103	**Good identification**						**06313103**
Actinomyces naeslundii		100.0	0.41	GLU 99			
06313121	**Good identification**						**06313121**
Actinomyces naeslundii		100.0	0.41	GLU 99			
06313122	**Low discrimination**						**06313122**
					SPO		
Actinomyces naeslundii		46.89	0.38	GLU 99	1		
Clostridium perfringens		26.75	0.11	RAF 92	84		
Clostridium barati		9.96	-.01		99		
06313123	**Good identification**						**06313123**
Actinomyces naeslundii		100.0	0.41	GLU 99			
06331000	**Acceptable identification**						**06331000**
					GRA	CAT	
Lactobacillus jensenii		90.41	0.44	GLU 99	99	1	
Propionibacterium granulosum		2.49	0.24	CEL 95	99	90	
Fusobacterium mortiferum		2.45	0.27		1	5	
06331002	**Acceptable identification**						**06331002**
Lactobacillus jensenii		92.12	0.41	GLU 99			
Lactobacillus acidophilus		3.41	0.14	CEL 95			
Clostridium perfringens		1.44	0.09				
06331010	**Good identification**						**06331010**
					CAT		
Lactobacillus jensenii		98.89	0.36	GLU 99	1		
Propionibacterium granulosum		1.12	0.09	CEL 95	90		
06331020	**Low discrimination**						**06331020**
Clostridium perfringens		72.74	0.41	GLU 99			
Lactobacillus jensenii		25.34	0.34	LAC 88			
Actinomyces naeslundii		0.84	0.30				
06331022	**Low discrimination**						**06331022**
Clostridium perfringens		62.85	0.35	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		31.02	0.31	LAC 88			
Lactobacillus acidophilus		4.59	0.14				
06331100	**Low discrimination**						**06331100**
					COC	GRA	
Atopobium parvulum		88.89	0.48	GLU 99	99	99	
Clostridium perfringens		6.96	0.30		1	99	
Fusobacterium mortiferum		2.70	0.33		1	1	
06331101	**Low discrimination**						**06331101**
					COC	SPO	
Actinomyces naeslundii		52.07	0.41	GLU 99	1	1	
Atopobium parvulum		30.87	0.14	RAF 92	99	1	
Clostridium chauvoei		17.07	0.09		1	99	
06331102	**Low discrimination**						**06331102**
					COC		
Atopobium parvulum		47.98	0.35	GLU 99	99		
Lactobacillus acidophilus		37.11	0.32		1		
Clostridium perfringens		10.02	0.24		1		
06331103	**Acceptable identification**						**06331103**
					COC		
Actinomyces naeslundii		90.53	0.41	GLU 99	1		
Atopobium parvulum		9.47	0.01	RAF 92	99		
06331120	**Acceptable identification**						**06331120**
					COC	SPO	
Clostridium perfringens		91.83	0.56	GLU 99	1	84	
Atopobium parvulum		6.40	0.35		99	1	
Clostridium septicum		1.24	0.17		1	99	
06331122	**Low discrimination**						**06331122**
					COC		
Clostridium perfringens		86.28	0.50	GLU 99	1		
Lactobacillus acidophilus		9.88	0.32		1		
Atopobium parvulum		2.25	0.22		99		
06331123	**Low discrimination**						**06331123**
Actinomyces naeslundii		58.60	0.41	GLU 99			
Clostridium perfringens		41.40	0.16	RAF 92			
06331160	**Good identification**						**06331160**
					COC		
Clostridium perfringens		99.22	0.43	GLU 99	1		
Actinomyces naeslundii		0.41	0.25		1		
Atopobium parvulum		0.37	0.01		99		
06331162	**Acceptable identification**						**06331162**
Clostridium perfringens		92.32	0.38	GLU 99			
Lactobacillus acidophilus		6.86	0.16				
Actinomyces naeslundii		0.82	0.25				
06333001	**Good identification**						**06333001**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06333003	**Good identification**						**06333003**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06333021	**Good identification**						**06333021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	GLU 99			
06333023	**Good identification**						**06333023**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06333100	**Low discrimination**						**06333100**
					COC	SPO	
Actinomyces naeslundii		64.89	0.43	GLU 99	1	1	
Atopobium parvulum		29.72	0.14	RAF 92	99	1	
Clostridium tertium		5.38	0.01		1	99	
06333101	**Good identification**						**06333101**
Actinomyces naeslundii		100.0	0.46	GLU 99			
06333102	**Low discrimination**						**06333102**
					SPO		
Actinomyces naeslundii		55.23	0.43	GLU 99	1		
Lactobacillus acidophilus		17.99	0.10	RAF 92	1		
Clostridium tertium		13.74	0.09		99		
06333103	**Good identification**						**06333103**
Actinomyces naeslundii		100.0	0.46	GLU 99			
06333121	**Good identification**						**06333121**
Actinomyces naeslundii		100.0	0.46	GLU 99			
06333122	**Low discrimination**						**06333122**
					SPO		
Actinomyces naeslundii		28.12	0.43	GLU 99	1		
Clostridium ramosum		26.38	0.17	RAF 92	92		
Clostridium tertium		20.99	0.17		99		
06333123	**Good identification**						**06333123**
Actinomyces naeslundii		100.0	0.46	GLU 99			
06334712	**Good identification**						**06334712**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06334732	**Good identification**						**06334732**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06335100	**Good identification**						**06335100**
					COC	GRA	
Atopobium parvulum		99.76	0.53	GLU 99	99	99	
Fusobacterium mortiferum		0.12	0.14		1	1	
Actinomyces israelii		0.12	0.16		1	99	
06335102	**Good identification**						**06335102**
					COC		
Atopobium parvulum		98.00	0.40	GLU 99	99		
Bifidobacterium adolescentis		1.01	0.13		1		
Actinomyces israelii		0.99	0.19		1		
06335120	**Acceptable identification**						**06335120**
					COC		
Atopobium parvulum		92.76	0.40	GLU 99	99		
Clostridium perfringens		7.24	0.21		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06335612	**Good identification**					**06335612**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06335632	**Good identification**					**06335632**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06335702	**Good identification**					**06335702**
Clostridium clostridioforme		100.0	0.36	GLU 88		
06335712	**Good identification**					**06335712**
Clostridium clostridioforme		100.0	0.47	GLU 88		
06335722	**Acceptable identification**					**06335722**
Clostridium clostridioforme		93.16	0.36	GLU 88		
Parabacteroides distasonis		6.84	0.22	RAF 94		
06335732	**Good identification**					**06335732**
Clostridium clostridioforme		99.14	0.47	GLU 88		
Parabacteroides distasonis		0.86	0.17	RAF 94		
06351000	**Low discrimination**					**06351000**
					SPO	
Lactobacillus catenaformis		65.65	0.63	GLU 99	1	
Lactobacillus jensenii		32.96	0.58		1	
Clostridium barati		1.10	0.27		99	
06351002	**Low discrimination**					**06351002**
Lactobacillus catenaformis		69.28	0.63	GLU 99		
Lactobacillus jensenii		23.19	0.55			
Lactobacillus acidophilus		4.01	0.39			
06351010	**Acceptable identification**					**06351010**
Lactobacillus jensenii		94.31	0.50	GLU 99		
Lactobacillus catenaformis		5.69	0.29			
06351012	**Acceptable identification**					**06351012**
Lactobacillus jensenii		90.88	0.47	GLU 99		
Lactobacillus catenaformis		8.23	0.29			
Lactobacillus acidophilus		0.48	0.05			
06351020	**Low discrimination**					**06351020**
					SPO	
Lactobacillus catenaformis		58.31	0.53	GLU 99	1	
Lactobacillus jensenii		29.28	0.47		1	
Clostridium barati		11.73	0.35		99	
06351022	**Low discrimination**					**06351022**
					SPO	
Lactobacillus catenaformis		45.77	0.53	GLU 99	1	
Clostridium barati		27.61	0.44		99	
Lactobacillus jensenii		15.32	0.44		1	
06351030	**Acceptable identification**					**06351030**
					SPO	
Lactobacillus jensenii		93.24	0.39	GLU 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		5.63	0.18		1		
Clostridium barati		1.13	0.01		99		
06351032	**Low discrimination**						**06351032**
					SPO		
Lactobacillus jensenii		85.76	0.36	GLU 99	1		
Lactobacillus catenaformis		7.76	0.18		1		
Clostridium barati		4.68	0.09		99		
06351100	**Low discrimination**						**06351100**
					SPO		
Lactobacillus catenaformis		89.08	0.66	GLU 99	1		
Collinsella aerofaciens		6.55	0.32		1		
Clostridium barati		2.99	0.35		99		
06351102	**Low discrimination**						**06351102**
					SPO		
Lactobacillus catenaformis		64.20	0.66	GLU 99	1		
Lactobacillus acidophilus		28.45	0.57		1		
Clostridium barati		6.46	0.44		99		
06351120	**Low discrimination**						**06351120**
					SPO		
Lactobacillus catenaformis		59.16	0.56	GLU 99	1		
Clostridium barati		23.79	0.44		99		
Clostridium paraputrificum		9.74	0.32		99		
06351122	**Low discrimination**						**06351122**
					SPO		
Lactobacillus acidophilus		42.28	0.57	GLU 99	1		
Clostridium barati		28.77	0.52		99		
Lactobacillus catenaformis		23.85	0.56		1		
06351142	**Low discrimination**						**06351142**
					SPO		
Lactobacillus acidophilus		74.41	0.41	GLU 99	1		
Lactobacillus catenaformis		13.72	0.32	SOR 11	1		
Clostridium barati		11.88	0.25		99		
06351162	**Low discrimination**						**06351162**
					SPO		
Lactobacillus acidophilus		63.77	0.41	GLU 99	1		
Clostridium barati		30.53	0.34	SOR 11	99		
Lactobacillus catenaformis		2.94	0.21		1		
06351302	**Low discrimination**						**06351302**
					SPO		
Lactobacillus acidophilus		63.17	0.38	GLU 99	1		
Lactobacillus catenaformis		19.13	0.32	RHA 7	1		
Clostridium barati		16.56	0.25		99		
06351322	**Low discrimination**						**06351322**
					GRA	SPO	
Lactobacillus acidophilus		43.22	0.38	GLU 99	99	1	
Clostridium barati		33.98	0.34	RHA 7	99	99	
Prevotella oralis		14.95	0.29		1	1	
06353001	**Good identification**						**06353001**
Actinomyces naeslundii		100.0	0.35	GLU 99			
06353003	**Good identification**						**06353003**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	GLU 99		
06353021	**Good identification**					**06353021**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06353023	**Good identification**					**06353023**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06353101	**Good identification**					**06353101**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06353102	**Low discrimination**					**06353102**
					SPO	
Lactobacillus acidophilus		45.66	0.35	GLU 99	1	
Lactobacillus catenaformis		19.77	0.32	MAN 5	1	
Clostridium barati		17.11	0.25		99	
06353103	**Acceptable identification**					**06353103**
Actinomyces naeslundii		91.33	0.43	GLU 99		
Lactobacillus acidophilus		8.67	0.01	RAF 92		
06353121	**Good identification**					**06353121**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06353123	**Low discrimination**					**06353123**
					SPO	
Actinomyces naeslundii		77.96	0.43	GLU 99	1	
Clostridium barati		8.33	-.01	RAF 92	99	
Lactobacillus acidophilus		7.40	0.01		1	
06353132	**Good identification**					**06353132**
					SPO	
Clostridium tertium		98.03	0.39	GLU 99	99	
Clostridium barati		0.74	-.01	TRE 90	99	
Lactobacillus acidophilus		0.65	0.01		1	
06355502	**Low discrimination**					**06355502**
Bifidobacterium adolescentis		77.18	0.36	GLU 99		
Actinomyces israelii		10.24	0.27	RAF 92		
Clostridium clostridioforme		6.96	0.15			
06355512	**Low discrimination**					**06355512**
Bifidobacterium adolescentis		74.85	0.37	GLU 99		
Clostridium clostridioforme		25.15	0.26	RAF 92		
06355712	**Good identification**					**06355712**
Clostridium clostridioforme		100.0	0.41	GLU 88		
06355732	**Good identification**					**06355732**
Clostridium clostridioforme		100.0	0.41	GLU 88		
06357100	**Low discrimination**					**06357100**
					SPO	
Actinomyces israelii		61.37	0.39	GLU 99	1	
Clostridium tertium		38.63	0.21		99	
06357122	**Good identification**					**06357122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium tertium		96.61	0.37	GLU 99	99	
Actinomyces israelii		1.63	0.21	TRE 90	1	
Clostridium barati		0.93	-.01		99	
06357132	**Good identification**					**06357132**
Clostridium tertium		100.0	0.41	GLU 99		
06357500	**Good identification**					**06357500**
					SPO	
Actinomyces israelii		96.01	0.37	GLU 99	1	
Clostridium butyricum		3.99	-.08		99	
06357512	**Low discrimination**					**06357512**
					SPO	
Bifidobacterium adolescentis		88.26	0.36	GLU 99	1	
Actinomyces israelii		9.91	0.26	RAF 92	1	
Clostridium tertium		1.83	-.01		99	
06371000	**Very good identification**					**06371000**
					COC	
Lactobacillus jensenii		98.76	0.66	GLU 99	1	
Lactobacillus catenaformis		0.66	0.29		1	
Atopobium parvulum		0.26	0.20		99	
06371002	**Low discrimination**					**06371002**
Lactobacillus jensenii		82.73	0.63	GLU 99		
Lactobacillus acidophilus		15.95	0.48			
Lactobacillus catenaformis		0.83	0.29			
06371010	**Very good identification**					**06371010**
Lactobacillus jensenii		100.0	0.58	GLU 99		
06371012	**Good identification**					**06371012**
					GRA	
Lactobacillus jensenii		99.17	0.55	GLU 99	99	
Lactobacillus acidophilus		0.58	0.14		99	
Leptotrichia buccalis		0.25	-.03		1	
06371020	**Good identification**					**06371020**
					SPO	
Lactobacillus jensenii		96.72	0.56	GLU 99	1	
Clostridium barati		1.12	0.17		99	
Lactobacillus catenaformis		0.65	0.18		1	
06371022	**Low discrimination**					**06371022**
					SPO	
Lactobacillus jensenii		54.24	0.53	GLU 99	1	
Lactobacillus acidophilus		41.82	0.48		1	
Clostridium barati		2.83	0.25		99	
06371030	**Good identification**					**06371030**
Lactobacillus jensenii		100.0	0.47	GLU 99		
06371032	**Good identification**					**06371032**
Lactobacillus jensenii		97.72	0.44	GLU 99		
Lactobacillus acidophilus		2.28	0.14			
06371100	**Low discrimination**					**06371100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Atopobium parvulum		85.77	0.54	GLU 99	99		
Lactobacillus acidophilus		5.02	0.32		1		
Lactobacillus jensenii		3.40	0.32		1		
06371102	**Very good identification**						**06371102**
					COC		
Lactobacillus acidophilus		95.21	0.66	GLU 99	1		
Atopobium parvulum		2.90	0.41		99		
Lactobacillus catenaformis		0.65	0.32		1		
06371120	**Low discrimination**						**06371120**
					COC	SPO	
Atopobium parvulum		37.49	0.41	GLU 99	99	1	
Lactobacillus acidophilus		12.43	0.32		1	1	
Clostridium septicum		9.87	0.25		1	99	
06371121	**Low discrimination**						**06371121**
					COC		
Actinomyces naeslundii		84.45	0.43	GLU 99	1		
Atopobium parvulum		15.55	0.07	RAF 92	99		
06371122	**Good identification**						**06371122**
					SPO		
Lactobacillus acidophilus		94.06	0.66	GLU 99	1		
Clostridium ramosum		2.52	0.38		92		
Clostridium barati		1.66	0.34		99		
06371142	**Good identification**						**06371142**
					COC	SPO	
Lactobacillus acidophilus		99.10	0.50	GLU 99	1	1	
Clostridium barati		0.41	0.07	SOR 11	1	99	
Atopobium parvulum		0.25	0.07		99	1	
06371162	**Good identification**						**06371162**
					SPO		
Lactobacillus acidophilus		97.68	0.50	GLU 99	1		
Clostridium barati		1.22	0.15	SOR 11	99		
Clostridium perfringens		0.55	0.14		84		
06371302	**Good identification**						**06371302**
					SPO		
Lactobacillus acidophilus		98.25	0.47	GLU 99	1		
Clostridium barati		0.67	0.07	RHA 7	99		
Clostridium ramosum		0.47	0.06		92		
06371322	**Low discrimination**						**06371322**
					SPO		
Lactobacillus acidophilus		66.65	0.47	GLU 99	1		
Clostridium ramosum		31.40	0.40	RHA 7	92		
Clostridium barati		1.36	0.15		99		
06371712	**Good identification**						**06371712**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06371732	**Good identification**						**06371732**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06373001	**Good identification**						**06373001**
Actinomyces naeslundii		100.0	0.40	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06373003	**Good identification**						**06373003**
Actinomyces naeslundii		100.0	0.40	GLU 99			
06373021	**Good identification**						**06373021**
Actinomyces naeslundii		100.0	0.40	GLU 99			
06373023	**Good identification**						**06373023**
Actinomyces naeslundii		100.0	0.40	GLU 99			
06373101	**Good identification**						**06373101**
					SPO		
Actinomyces naeslundii		96.00	0.48	GLU 99	1		
Clostridium tertium		4.01	0.01	RAF 92	99		
06373102	**Low discrimination**						**06373102**
					SPO		
Lactobacillus acidophilus		53.19	0.44	GLU 99	1		
Clostridium tertium		40.63	0.43	MAN 5	99		
Actinomyces naeslundii		2.19	0.45		1		
06373103	**Low discrimination**						**06373103**
					SPO		
Actinomyces naeslundii		77.58	0.48	GLU 99	1		
Lactobacillus acidophilus		12.71	0.10	RAF 92	1		
Clostridium tertium		9.71	0.09		99		
06373110	**Good identification**						**06373110**
					SPO		
Clostridium tertium		98.87	0.39	GLU 99	99		
Actinomyces israelii		0.66	0.16		1		
Actinomyces naeslundii		0.47	0.23		1		
06373112	**Good identification**						**06373112**
					GRA	SPO	
Clostridium tertium		98.26	0.47	GLU 99	99	99	
Lactobacillus acidophilus		0.73	0.10		99	1	
Leptotrichia buccalis		0.52	-.03		1	1	
06373120	**Low discrimination**						**06373120**
					SPO		
Clostridium tertium		49.26	0.43	GLU 99	99		
Clostridium ramosum		46.43	0.41		92		
Actinomyces naeslundii		2.65	0.45		1		
06373121	**Low discrimination**						**06373121**
					SPO		
Actinomyces naeslundii		80.44	0.48	GLU 99	1		
Clostridium tertium		10.07	0.09	RAF 92	99		
Clostridium ramosum		9.49	0.07		92		
06373122	**Low discrimination**						**06373122**
					SPO		
Clostridium ramosum		46.14	0.51	GLU 99	92		
Clostridium tertium		36.71	0.51		99		
Lactobacillus acidophilus		16.02	0.44		1		
06373123	**Low discrimination**						**06373123**
					SPO		
Actinomyces naeslundii		49.72	0.48	GLU 99	1		
Clostridium ramosum		23.46	0.17	RAF 92	92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		18.67	0.17		99		
06373130	**Good identification**						**06373130**
					SPO		
Clostridium tertium		99.31	0.47	GLU 99	99		
Clostridium ramosum		0.53	0.07		92		
Actinomyces naeslundii		0.16	0.23		1		
06373132	**Good identification**						**06373132**
					SPO		
Clostridium tertium		99.00	0.55	GLU 99	99		
Clostridium ramosum		0.71	0.17		92		
Lactobacillus acidophilus		0.25	0.10		1		
06373320	**Good identification**						**06373320**
Clostridium ramosum		99.20	0.43	GLU 99			
Clostridium tertium		0.80	0.09				
06373322	**Good identification**						**06373322**
					SPO		
Clostridium ramosum		97.42	0.53	GLU 99	92		
Lactobacillus acidophilus		1.92	0.25		1		
Clostridium tertium		0.59	0.17		99		
06374612	**Good identification**						**06374612**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06374632	**Good identification**						**06374632**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06374702	**Good identification**						**06374702**
Clostridium clostridioforme		100.0	0.36	GLU 88			
06374712	**Good identification**						**06374712**
Clostridium clostridioforme		100.0	0.48	GLU 88			
06374722	**Good identification**						**06374722**
Clostridium clostridioforme		100.0	0.36	GLU 88			
06374732	**Good identification**						**06374732**
Clostridium clostridioforme		100.0	0.48	GLU 88			
06375100	**Very good identification**						**06375100**
					COC	SPO	
Atopobium parvulum		98.74	0.59	GLU 99	99	1	
Actinomyces israelii		0.67	0.36		1	1	
Clostridium butyricum		0.54	0.13		1	99	
06375102	**Low discrimination**						**06375102**
					COC		
Atopobium parvulum		78.67	0.46	GLU 99	99		
Lactobacillus acidophilus		14.04	0.32		1		
Actinomyces israelii		4.52	0.39		1		
06375120	**Good identification**						**06375120**
					COC	SPO	
Atopobium parvulum		98.52	0.46	GLU 99	99	1	
Clostridium tertium		0.98	0.11		1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.26	-.05		1	99	
06375312	**Good identification**						**06375312**
Clostridium clostridioforme		100.0	0.35	GLU 88			
06375332	**Good identification**						**06375332**
Clostridium clostridioforme		96.55	0.35	GLU 88			
Parabacteroides distasonis		3.45	0.16	RAF 94			
06375512	**Low discrimination**						**06375512**
Clostridium clostridioforme		76.03	0.41	GLU 88			
Bifidobacterium adolescentis		20.57	0.34	RAF 94			
Actinomyces israelii		1.88	0.23	RHA 88			
06375532	**Good identification**						**06375532**
Clostridium clostridioforme		100.0	0.41	GLU 88			
06375602	**Good identification**						**06375602**
Clostridium clostridioforme		100.0	0.36	GLU 88			
06375612	**Good identification**						**06375612**
Clostridium clostridioforme		100.0	0.47	GLU 88			
06375622	**Good identification**						**06375622**
Clostridium clostridioforme		100.0	0.36	GLU 88			
06375632	**Good identification**						**06375632**
Clostridium clostridioforme		100.0	0.47	GLU 88			
06375702	**Good identification**						**06375702**
					SPO		
Clostridium clostridioforme		98.52	0.45	GLU 88	66		
Actinomyces israelii		0.76	0.18	RAF 94	1		
Clostridium butyricum		0.72	-.04		99		
06375712	**Good identification**						**06375712**
Clostridium clostridioforme		100.0	0.56	GLU 88			
06375722	**Good identification**						**06375722**
Clostridium clostridioforme		98.83	0.45	GLU 88			
Parabacteroides distasonis		1.17	0.17	RAF 94			
06375732	**Good identification**						**06375732**
Clostridium clostridioforme		99.86	0.56	GLU 88			
Parabacteroides distasonis		0.14	0.13	RAF 94			
06377000	**Low discrimination**						**06377000**
					SPO		
Actinomyces israelii		89.37	0.38	GLU 99	1		
Clostridium tertium		5.44	0.03		99		
Clostridium innocuum		5.20	0.02		99		
06377002	**Low discrimination**						**06377002**
					SPO		
Actinomyces israelii		86.17	0.41	GLU 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		10.49	0.11		99		
Clostridium innocuum		3.34	0.02		99		
06377100	**Low discrimination**						**06377100**
					COC	SPO	
Clostridium tertium		54.07	0.37	GLU 99	1	99	
Actinomyces israelii		35.90	0.48		1	1	
Atopobium parvulum		9.46	0.25		99	1	
06377102	**Low discrimination**						**06377102**
					COC	SPO	
Clostridium tertium		73.55	0.45	GLU 99	1	99	
Actinomyces israelii		24.42	0.51		1	1	
Atopobium parvulum		0.76	0.12		99	1	
06377110	**Acceptable identification**						**06377110**
					SPO		
Clostridium tertium		94.71	0.41	GLU 99	99		
Actinomyces israelii		5.29	0.34		1		
06377112	**Good identification**						**06377112**
					SPO		
Clostridium tertium		96.78	0.50	GLU 99	99		
Actinomyces israelii		2.70	0.37		1		
Bifidobacterium adolescentis		0.52	0.19		1		
06377120	**Good identification**						**06377120**
					COC	SPO	
Clostridium tertium		96.96	0.45	GLU 99	1	99	
Actinomyces israelii		1.37	0.28		1	1	
Atopobium parvulum		1.00	0.12		99	1	
06377122	**Good identification**						**06377122**
					SPO		
Clostridium tertium		98.09	0.53	GLU 99	99		
Clostridium ramosum		0.90	0.17		92		
Actinomyces israelii		0.69	0.31		1		
06377130	**Good identification**						**06377130**
					SPO		
Clostridium tertium		99.88	0.50	GLU 99	99		
Actinomyces israelii		0.12	0.14		1		
06377132	**Very good identification**						**06377132**
					SPO		
Clostridium tertium		99.94	0.58	GLU 99	99		
Actinomyces israelii		0.06	0.17		1		
06377140	**Good identification**						**06377140**
					SPO		
Actinomyces israelii		96.24	0.41	GLU 99	1		
Clostridium tertium		3.76	0.03		99		
06377142	**Acceptable identification**						**06377142**
					SPO		
Actinomyces israelii		92.74	0.44	GLU 99	1		
Clostridium tertium		7.26	0.11		99		
06377400	**Good identification**						**06377400**
Actinomyces israelii		100.0	0.37	GLU 99			
06377402	**Good identification**						**06377402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	GLU 99			
06377500	**Low discrimination**						**06377500**
					SPO		
Actinomyces israelii		83.91	0.47	GLU 99	1		
Clostridium butyricum		14.53	0.13		99		
Clostridium tertium		1.56	0.03		99		
06377502	**Low discrimination**						**06377502**
					SPO		
Actinomyces israelii		78.43	0.50	GLU 99	1		
Bifidobacterium adolescentis		15.13	0.31		1		
Clostridium butyricum		3.52	0.06		99		
06377540	**Good identification**						**06377540**
					SPO		
Actinomyces israelii		96.27	0.40	GLU 99	1		
Clostridium butyricum		3.73	-.05		99		
06377542	**Acceptable identification**						**06377542**
Actinomyces israelii		94.22	0.43	GLU 99			
Bifidobacterium adolescentis		5.78	0.16				
06400000	**Low discrimination**						**06400000**
					COC		
Anaerococcus prevotii		83.89	0.41	SUC 1	99		
Parvimonas micra		5.42	0.15	URE 90	99		
Atopobium minutum		3.17	0.31		1		
06400001	**Very good identification**						**06400001**
					COC	GRA	
Anaerococcus prevotii		99.93	0.58	SUC 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Peptococcus niger		0.02	-.11		99	99	
06400003	**Good identification**						**06400003**
Anaerococcus prevotii		100.0	0.47	SUC 1			
06410000	**Low discrimination**						**06410000**
					GRA	CAT	
Propionibacterium granulosum		72.56	0.58	GLU 99	99	90	
Fusobacterium mortiferum		13.76	0.49		1	5	
Prevotella intermedia		7.55	0.33		1	1	
06410002	**Low discrimination**						**06410002**
					GRA	CAT	
Propionibacterium granulosum		84.75	0.47	GLU 99	99	90	
Atopobium minutum		7.31	0.27		99	1	
Fusobacterium mortiferum		7.14	0.32		1	5	
06410010	**Good identification**						**06410010**
					GRA	CAT	
Propionibacterium granulosum		98.62	0.43	GLU 99	99	90	
Fusobacterium mortiferum		1.39	0.14	MLZ 12	1	5	
06410020	**Acceptable identification**						**06410020**
					GRA	CAT	
Propionibacterium granulosum		94.62	0.47	GLU 99	99	90	
Atopobium minutum		4.08	0.22		99	1	
Fusobacterium mortiferum		0.73	0.14		1	5	
06410022	**Low discrimination**						**06410022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Propionibacterium granulosum		89.69	0.37	GLU 99	90		
Atopobium minutum		10.31	0.19		1		
06410100	**Good identification**						**06410100**
					GRA	CAT	
Fusobacterium mortiferum		97.06	0.55	GLU 99	1	5	
Propionibacterium granulosum		2.22	0.24		99	90	
Lactobacillus fermentum		0.39	0.15		99	1	
06410102	**Acceptable identification**						**06410102**
					GRA	CAT	
Fusobacterium mortiferum		91.97	0.39	GLU 99	1	5	
Propionibacterium granulosum		4.73	0.13	BGL 10	99	90	
Lactobacillus fermentum		3.30	0.15		99	1	
06411000	**Low discrimination**						**06411000**
					GRA	CAT	
Propionibacterium granulosum		66.06	0.58	GLU 99	99	90	
Fusobacterium mortiferum		18.79	0.52		1	5	
Lactobacillus fermentum		7.08	0.46		99	1	
06411002	**Low discrimination**						**06411002**
					GRA	CAT	
Propionibacterium granulosum		63.15	0.47	GLU 99	99	90	
Lactobacillus fermentum		27.05	0.46		99	1	
Fusobacterium mortiferum		7.98	0.35		1	5	
06411010	**Good identification**						**06411010**
					GRA	CAT	
Propionibacterium granulosum		97.94	0.43	GLU 99	99	90	
Fusobacterium mortiferum		2.06	0.17	MLZ 12	1	5	
06411020	**Low discrimination**						**06411020**
					GRA	CAT	
Propionibacterium granulosum		79.28	0.47	GLU 99	99	90	
Lactobacillus fermentum		11.32	0.38		99	1	
Prevotella melaninogenica		7.35	0.15		1	1	
06411022	**Low discrimination**						**06411022**
					GRA	CAT	
Propionibacterium granulosum		59.52	0.37	GLU 99	99	90	
Lactobacillus fermentum		34.00	0.38		99	1	
Prevotella melaninogenica		4.20	0.03		1	1	
06411100	**Low discrimination**						**06411100**
					GRA		
Fusobacterium mortiferum		75.93	0.58	GLU 99	1		
Lactobacillus fermentum		19.99	0.50		99		
Prevotella melaninogenica		2.65	0.15		1		
06411102	**Low discrimination**						**06411102**
					GRA		
Lactobacillus fermentum		68.41	0.50	GLU 99	99		
Fusobacterium mortiferum		28.87	0.42		1		
Prevotella melaninogenica		1.73	0.03		1		
06411120	**Good identification**						**06411120**
					GRA		
Prevotella melaninogenica		96.80	0.50	GLU 90	1		
Lactobacillus fermentum		2.46	0.41	MLT 99	99		
Clostridium perfringens		0.36	0.15		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06411122	**Low discrimination**						**06411122**
					GRA		
Prevotella melaninogenica		86.53	0.37	GLU 90	1		
Lactobacillus fermentum		11.54	0.41	MLT 99	99		
Clostridium perfringens		0.79	0.09		99		
06411400	**Low discrimination**						**06411400**
					GRA	CAT	
Lactobacillus fermentum		89.20	0.46	GLU 99	99	1	
Propionibacterium granulosum		8.41	0.24		99	90	
Fusobacterium mortiferum		2.39	0.17		1	5	
06411402	**Good identification**						**06411402**
					CAT		
Lactobacillus fermentum		97.70	0.46	GLU 99	1		
Propionibacterium granulosum		2.30	0.13		90		
06411420	**Acceptable identification**						**06411420**
					CAT		
Lactobacillus fermentum		93.39	0.38	GLU 99	1		
Propionibacterium granulosum		6.61	0.13		90		
06411422	**Good identification**						**06411422**
Lactobacillus fermentum		100.0	0.38	GLU 99			
06411500	**Good identification**						**06411500**
					GRA		
Lactobacillus fermentum		95.25	0.50	GLU 99	99		
Fusobacterium mortiferum		3.65	0.24		1		
Bifidobacterium longum subsp. longum		1.10	0.09		99		
06411502	**Good identification**						**06411502**
Lactobacillus fermentum		98.86	0.50	GLU 99			
Bifidobacterium longum subsp. longum		1.14	0.09				
06411520	**Low discrimination**						**06411520**
					GRA		
Lactobacillus fermentum		65.12	0.41	GLU 99	99		
Prevotella melaninogenica		25.90	0.15		1		
Bifidobacterium longum subsp. longum		8.98	0.20		99		
06411522	**Low discrimination**						**06411522**
					GRA		
Lactobacillus fermentum		80.91	0.41	GLU 99	99		
Bifidobacterium longum subsp. longum		11.16	0.20		99		
Prevotella melaninogenica		6.13	0.03		1		
06412000	**Good identification**						**06412000**
					GRA	CAT	
Propionibacterium granulosum		99.56	0.56	GLU 99	99	90	
Fusobacterium mortiferum		0.25	0.14		1	5	
Eubacterium limosum		0.19	-.06		99	1	
06412002	**Good identification**						**06412002**
Propionibacterium granulosum		100.0	0.45	GLU 99			
06412010	**Good identification**						**06412010**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06412020	**Good identification**						**06412020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.45	GLU 99			
06412022	**Good identification**						**06412022**
Propionibacterium granulosum		100.0	0.35	GLU 99			
06413000	**Good identification**						**06413000**
					GRA	CAT	
Propionibacterium granulosum		99.62	0.56	GLU 99	99	90	
Fusobacterium mortiferum		0.38	0.17		1	5	
06413002	**Good identification**						**06413002**
Propionibacterium granulosum		100.0	0.45	GLU 99			
06413010	**Good identification**						**06413010**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06413020	**Good identification**						**06413020**
Propionibacterium granulosum		100.0	0.45	GLU 99			
06413022	**Good identification**						**06413022**
Propionibacterium granulosum		100.0	0.35	GLU 99			
06414100	**Acceptable identification**						**06414100**
					GRA		
Fusobacterium mortiferum		91.67	0.36	GLU 99	1		
Actinomyces meyeri		8.33	0.09	XYL 7	99		
06415000	**Low discrimination**						**06415000**
					GRA	CAT	
Lactobacillus fermentum		66.63	0.42	GLU 99	99	1	
Fusobacterium mortiferum		22.68	0.32		1	5	
Propionibacterium granulosum		10.70	0.24		99	90	
06415002	**Acceptable identification**						**06415002**
					GRA	CAT	
Lactobacillus fermentum		92.77	0.42	GLU 99	99	1	
Propionibacterium granulosum		3.72	0.13		99	90	
Fusobacterium mortiferum		3.51	0.16		1	5	
06415100	**Low discrimination**						**06415100**
					GRA		
Lactobacillus fermentum		65.32	0.46	GLU 99	99		
Fusobacterium mortiferum		31.79	0.39		1		
Actinomyces meyeri		2.89	0.12		99		
06415102	**Acceptable identification**						**06415102**
					GRA		
Lactobacillus fermentum		94.87	0.46	GLU 99	99		
Fusobacterium mortiferum		5.13	0.22		1		
06415120	**Low discrimination**						**06415120**
					GRA		
Lactobacillus fermentum		59.62	0.37	GLU 99	99		
Prevotella melaninogenica		40.38	0.15		1		
06415122	**Low discrimination**						**06415122**
					GRA		
Lactobacillus fermentum		55.14	0.37	GLU 99	99		
Parabacteroides distasonis		30.67	0.30		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		7.11	0.03		1		
06415322	**Good identification**						**06415322**
Parabacteroides distasonis		100.0	0.38	GLU 99			
06415400	**Good identification**						**06415400**
Lactobacillus fermentum		100.0	0.42	GLU 99			
06415402	**Good identification**						**06415402**
Lactobacillus fermentum		100.0	0.42	GLU 99			
06415500	**Acceptable identification**						**06415500**
Lactobacillus fermentum		94.45	0.46	GLU 99			
Bifidobacterium longum subsp. longum		5.55	0.17				
06415502	**Acceptable identification**						**06415502**
Lactobacillus fermentum		94.45	0.46	GLU 99			
Bifidobacterium longum subsp. longum		5.55	0.17				
06415520	**Low discrimination**						**06415520**
					GRA		
Lactobacillus fermentum		47.78	0.37	GLU 99	99		
Bifidobacterium longum subsp. longum		33.67	0.28		99		
Bacteroides vulgatus		18.55	0.09		1		
06415522	**Low discrimination**						**06415522**
					GRA		
Lactobacillus fermentum		48.88	0.37	GLU 99	99		
Bifidobacterium longum subsp. longum		34.45	0.28		99		
Parabacteroides distasonis		16.67	0.27		1		
06430000	**Very good identification**						**06430000**
					GRA	CAT	
Propionibacterium granulosum		98.15	0.66	GLU 99	99	90	
Fusobacterium mortiferum		1.85	0.40		1	5	
06430002	**Good identification**						**06430002**
					GRA	CAT	
Propionibacterium granulosum		99.17	0.56	GLU 99	99	90	
Fusobacterium mortiferum		0.83	0.23		1	5	
06430010	**Good identification**						**06430010**
Propionibacterium granulosum		100.0	0.51	GLU 99			
06430012	**Good identification**						**06430012**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06430020	**Good identification**						**06430020**
Propionibacterium granulosum		100.0	0.56	GLU 99			
06430022	**Good identification**						**06430022**
Propionibacterium granulosum		100.0	0.45	GLU 99			
06430030	**Good identification**						**06430030**
Propionibacterium granulosum		100.0	0.41	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06430100	**Low discrimination**						**06430100**
					GRA	CAT	
Fusobacterium mortiferum		81.35	0.46	GLU 99	1	5	
Propionibacterium granulosum		18.65	0.32		99	90	
06430200	**Good identification**						**06430200**
Propionibacterium granulosum		100.0	0.40	GLU 99			
06431000	**Very good identification**						**06431000**
					GRA	CAT	
Propionibacterium granulosum		97.25	0.66	GLU 99	99	90	
Fusobacterium mortiferum		2.75	0.43		1	5	
06431002	**Good identification**						**06431002**
					GRA	CAT	
Propionibacterium granulosum		98.76	0.56	GLU 99	99	90	
Fusobacterium mortiferum		1.24	0.26		1	5	
06431010	**Good identification**						**06431010**
Propionibacterium granulosum		100.0	0.51	GLU 99			
06431012	**Good identification**						**06431012**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06431020	**Good identification**						**06431020**
					CAT		
Propionibacterium granulosum		99.72	0.56	GLU 99	90		
Clostridium perfringens		0.29	0.05		1		
06431022	**Good identification**						**06431022**
Propionibacterium granulosum		100.0	0.45	GLU 99			
06431030	**Good identification**						**06431030**
Propionibacterium granulosum		100.0	0.41	GLU 99			
06431100	**Low discrimination**						**06431100**
					GRA	CAT	
Fusobacterium mortiferum		85.37	0.49	GLU 99	1	5	
Propionibacterium granulosum		13.05	0.32		99	90	
Actinomyces naeslundii		0.82	0.26		99	25	
06431200	**Good identification**						**06431200**
Propionibacterium granulosum		100.0	0.40	GLU 99			
06432000	**Very good identification**						**06432000**
Propionibacterium granulosum		100.0	0.64	GLU 99			
06432002	**Good identification**						**06432002**
Propionibacterium granulosum		100.0	0.53	GLU 99			
06432010	**Good identification**						**06432010**
Propionibacterium granulosum		100.0	0.49	GLU 99			
06432012	**Good identification**						**06432012**
Propionibacterium granulosum		100.0	0.39	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06432020	**Good identification**					**06432020**
Propionibacterium granulosum		100.0	0.53	GLU 99		
06432022	**Good identification**					**06432022**
Propionibacterium granulosum		100.0	0.43	GLU 99		
06432030	**Good identification**					**06432030**
Propionibacterium granulosum		100.0	0.39	GLU 99		
06432200	**Good identification**					**06432200**
Propionibacterium granulosum		100.0	0.38	GLU 99		
06433000	**Very good identification**					**06433000**
Propionibacterium granulosum		99.94	0.64	GLU 99		
Actinomyces naeslundii		0.06	0.23			
06433002	**Good identification**					**06433002**
Propionibacterium granulosum		99.77	0.53	GLU 99		
Actinomyces naeslundii		0.23	0.23			
06433010	**Good identification**					**06433010**
Propionibacterium granulosum		100.0	0.49	GLU 99		
06433012	**Good identification**					**06433012**
Propionibacterium granulosum		100.0	0.39	GLU 99		
06433020	**Good identification**					**06433020**
Propionibacterium granulosum		99.77	0.53	GLU 99		
Actinomyces naeslundii		0.23	0.23			
06433022	**Good identification**					**06433022**
Propionibacterium granulosum		99.09	0.43	GLU 99		
Actinomyces naeslundii		0.91	0.23			
06433030	**Good identification**					**06433030**
Propionibacterium granulosum		100.0	0.39	GLU 99		
06433200	**Good identification**					**06433200**
Propionibacterium granulosum		100.0	0.38	GLU 99		
06435122	**Good identification**					**06435122**
Parabacteroides distasonis		100.0	0.41	GLU 99		
06435132	**Good identification**					**06435132**
Parabacteroides distasonis		100.0	0.37	GLU 99		
06435322	**Good identification**					**06435322**
Parabacteroides distasonis		100.0	0.50	GLU 99		
06435332	**Good identification**					**06435332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.45	GLU 99			
06435522	**Low discrimination**						**06435522**
					GRA		
Parabacteroides distasonis		89.81	0.38	GLU 99	1		
Bifidobacterium longum subsp. longum		10.19	0.17		99		
06435722	**Good identification**						**06435722**
Parabacteroides distasonis		97.89	0.46	GLU 99			
Clostridium clostridioforme		1.12	0.07				
Bacteroides thetaiotaomicron		1.00	-.04				
06435732	**Acceptable identification**						**06435732**
Parabacteroides distasonis		91.21	0.41	GLU 99			
Clostridium clostridioforme		8.79	0.18				
06450000	**Low discrimination**						**06450000**
					GRA	CAT	
Fusobacterium mortiferum		86.22	0.40	GLU 99	1	5	
Propionibacterium granulosum		13.78	0.24		99	90	
06450100	**Good identification**						**06450100**
Fusobacterium mortiferum		100.0	0.47	GLU 99			
06451000	**Acceptable identification**						**06451000**
					GRA	CAT	
Fusobacterium mortiferum		90.37	0.43	GLU 99	1	5	
Propionibacterium granulosum		9.63	0.24		99	90	
06451100	**Good identification**						**06451100**
					GRA		
Fusobacterium mortiferum		97.52	0.50	GLU 99	1		
Prevotella melaninogenica		1.14	-.01		1		
Lactobacillus fermentum		0.78	0.15		99		
06470100	**Good identification**						**06470100**
Fusobacterium mortiferum		100.0	0.38	GLU 99			
06471100	**Good identification**						**06471100**
					GRA		
Fusobacterium mortiferum		96.34	0.41	GLU 99	1		
Actinomyces naeslundii		3.67	0.28		99		
06473101	**Good identification**						**06473101**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06473103	**Good identification**						**06473103**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06473121	**Good identification**						**06473121**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06473123	**Good identification**						**06473123**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06475122	**Good identification**						**06475122**
Parabacteroides distasonis		100.0	0.37	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06475322	**Good identification**					**06475322**
Parabacteroides distasonis		100.0	0.45	GLU 99		
06475332	**Good identification**					**06475332**
Parabacteroides distasonis		97.71	0.40	GLU 99		
Clostridium clostridioforme		2.29	0.07			
06475722	**Acceptable identification**					**06475722**
Parabacteroides distasonis		90.00	0.41	GLU 99		
Clostridium clostridioforme		6.38	0.16			
Bacteroides thetaiotaomicron		3.62	0.01			
06475732	**Low discrimination**					**06475732**
Parabacteroides distasonis		61.41	0.37	GLU 99		
Clostridium clostridioforme		36.81	0.27			
Bacteroides thetaiotaomicron		1.78	-.06			
06501120	**Very good identification**					**06501120**
					GRA	
Prevotella melaninogenica		99.96	0.67	GLU 90	1	
Clostridium perfringens		0.03	0.15	FRU 90	99	
Lactobacillus fermentum		0.01	0.15		99	
06501122	**Good identification**					**06501122**
					GRA	
Prevotella melaninogenica		99.81	0.55	GLU 90	1	
Clostridium perfringens		0.08	0.09	FRU 90	99	
Prevotella oralis		0.08	0.05		1	
06501522	**Low discrimination**					**06501522**
					GRA	
Bifidobacterium longum subsp. longum		63.73	0.37	GLU 99	99	
Prevotella melaninogenica		35.00	0.21	FRU 90	1	
Lactobacillus fermentum		1.27	0.15		99	
06501530	**Good identification**					**06501530**
					GRA	
Bifidobacterium longum subsp. longum		98.32	0.41	GLU 99	99	
Prevotella melaninogenica		1.68	-.01	FRU 90	1	
06501532	**Good identification**					**06501532**
Bifidobacterium longum subsp. longum		100.0	0.41	GLU 99		
06505500	**Good identification**					**06505500**
Bifidobacterium longum subsp. longum		95.52	0.35	GLU 99		
Lactobacillus fermentum		4.48	0.20	FRU 90		
06505502	**Good identification**					**06505502**
Bifidobacterium longum subsp. longum		95.52	0.35	GLU 99		
Lactobacillus fermentum		4.48	0.20	FRU 90		
06505510	**Good identification**					**06505510**
Bifidobacterium longum subsp. longum		100.0	0.39	GLU 99		
06505512	**Good identification**					**06505512**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		100.0	0.39	GLU 99				
06505520	**Acceptable identification**							**06505520**
					GRA			
Bifidobacterium longum subsp. longum		94.36	0.46	GLU 99	99			
Bacteroides vulgatus		4.73	0.09	FRU 90	1			
Prevotella melaninogenica		0.92	-.01		1			
06505522	**Good identification**							**06505522**
Bifidobacterium longum subsp. longum		100.0	0.46	GLU 99				
06505530	**Good identification**							**06505530**
Bifidobacterium longum subsp. longum		100.0	0.50	GLU 99				
06505532	**Good identification**							**06505532**
Bifidobacterium longum subsp. longum		100.0	0.50	GLU 99				
06510000	**Good identification**							**06510000**
					COC	GRA	CAT	
Prevotella intermedia		94.88	0.67	GLU 90	1	1	1	
Propionibacterium granulosum		4.14	0.52		1	99	90	
Gemella morbillorum		0.72	0.32		99	99	1	
06510002	**Low discrimination**							**06510002**
					GRA	CAT		
Propionibacterium granulosum		39.87	0.41	GLU 99	99	90		
Prevotella intermedia		36.92	0.33		1	1		
Actinomyces viscosus		21.14	0.27		99	76		
06510020	**Low discrimination**							**06510020**
					GRA	CAT		
Propionibacterium granulosum		45.27	0.41	GLU 99	99	90		
Prevotella intermedia		41.92	0.33		1	1		
Prevotella melaninogenica		9.34	0.15		1	1		
06510120	**Good identification**							**06510120**
					GRA			
Prevotella melaninogenica		99.56	0.50	GLU 90	1			
Clostridium perfringens		0.37	0.15	GAL 99	99			
Lactobacillus fermentum		0.08	0.15		99			
06510122	**Good identification**							**06510122**
					GRA			
Prevotella melaninogenica		97.90	0.37	GLU 90	1			
Clostridium perfringens		0.89	0.09	GAL 99	99			
Prevotella oralis		0.82	0.05		1			
06511000	**Good identification**							**06511000**
					GRA	CAT		
Prevotella intermedia		92.13	0.67	GLU 90	1	1		
Propionibacterium granulosum		4.02	0.52		99	90		
Lactobacillus fermentum		2.88	0.54		99	1		
06511002	**Low discrimination**							**06511002**
					GRA	CAT		
Lactobacillus fermentum		54.39	0.54	GLU 99	99	1		
Propionibacterium granulosum		19.01	0.41		99	90		
Prevotella intermedia		17.61	0.33		1	1		
06511020	**Low discrimination**							**06511020**
					GRA	CAT		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		83.63	0.50	GLU 90	1	1	
Clostridium perfringens		4.17	0.34	LAC 99	99	1	
Propionibacterium granulosum		4.10	0.41		99	90	
06511022	**Low discrimination**						**06511022**
					GRA		
Prevotella melaninogenica		66.78	0.37	GLU 90	1		
Lactobacillus fermentum		16.37	0.46	LAC 99	99		
Clostridium perfringens		8.22	0.29		99		
06511023	**Low discrimination**						**06511023**
					GRA		
Actinomyces naeslundii		64.64	0.37	GLU 99	99		
Prevotella melaninogenica		35.36	0.03		1		
06511100	**Low discrimination**						**06511100**
					GRA		
Prevotella melaninogenica		75.10	0.50	GLU 90	1		
Lactobacillus fermentum		17.16	0.58	NAG 99	99		
Prevotella intermedia		3.41	0.33		1		
06511102	**Low discrimination**						**06511102**
					GRA		
Lactobacillus fermentum		47.81	0.58	GLU 99	99		
Prevotella melaninogenica		39.85	0.37		1		
Actinomyces viscosus		5.92	0.28		99		
06511103	**Low discrimination**						**06511103**
					SPO		
Actinomyces naeslundii		37.37	0.45	GLU 99	1		
Actinomyces viscosus		35.20	0.20		1		
Clostridium chauvoei		11.64	0.13		99		
06511120	**Very good identification**						**06511120**
					GRA		
Prevotella melaninogenica		99.53	0.84	GLU 90	1		
Clostridium perfringens		0.37	0.49		99		
Lactobacillus fermentum		0.08	0.50		99		
06511121	**Good identification**						**06511121**
					GRA	SPO	
Prevotella melaninogenica		98.01	0.50	GLU 90	1	1	
Actinomyces naeslundii		0.98	0.45	URE 1	99	1	
Clostridium chauvoei		0.58	0.18		99	99	
06511122	**Very good identification**						**06511122**
					GRA		
Prevotella melaninogenica		97.63	0.71	GLU 90	1		
Clostridium perfringens		0.89	0.43		99		
Prevotella oralis		0.81	0.39		1		
06511123	**Acceptable identification**						**06511123**
					GRA		
Prevotella melaninogenica		92.64	0.37	GLU 90	1		
Actinomyces naeslundii		4.87	0.45	URE 1	99		
Clostridium perfringens		0.84	0.09		99		
06511130	**Good identification**						**06511130**
					GRA		
Prevotella melaninogenica		98.97	0.50	GLU 90	1		
Bifidobacterium longum subsp. longum		0.59	0.24	MLZ 1	99		
Clostridium perfringens		0.37	0.15		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06511132	**Acceptable identification**						**06511132**
					GRA	CAT	
Prevotella melaninogenica		92.31	0.37	GLU 90	1	1	
Bifidobacterium longum subsp. longum		2.89	0.24	MLZ 1	99	1	
Capnocytophaga ochracea		1.96	0.12		1	90	
06511160	**Acceptable identification**						**06511160**
					GRA		
Prevotella melaninogenica		93.33	0.50	GLU 90	1		
Clostridium perfringens		6.49	0.37	SOR 1	99		
Actinomyces naeslundii		0.11	0.29		99		
06511162	**Low discrimination**						**06511162**
					GRA	CAT	
Prevotella melaninogenica		82.53	0.37	GLU 90	1	1	
Clostridium perfringens		14.19	0.31	SOR 1	99	1	
Capnocytophaga ochracea		1.75	0.12		1	90	
06511320	**Good identification**						**06511320**
					GRA		
Prevotella melaninogenica		98.43	0.50	GLU 90	1		
Bacteroides vulgatus		0.78	0.11	RHA 1	1		
Clostridium perfringens		0.36	0.15		99		
06511322	**Low discrimination**						**06511322**
Prevotella melaninogenica		70.50	0.37	GLU 90			
Prevotella oralis		24.95	0.33	RHA 1			
Parabacteroides distasonis		3.63	0.31				
06511400	**Low discrimination**						**06511400**
					COC	GRA	
Lactobacillus fermentum		73.61	0.54	GLU 99	1	99	
Prevotella intermedia		23.83	0.33		1	1	
Gemella morbillorum		1.06	0.11		99	99	
06511402	**Good identification**						**06511402**
Lactobacillus fermentum		99.38	0.54	GLU 99			
Bifidobacterium longum subsp. longum		0.62	0.09				
06511420	**Low discrimination**						**06511420**
					GRA		
Lactobacillus fermentum		77.44	0.46	GLU 99	99		
Prevotella melaninogenica		16.75	0.15		1		
Bifidobacterium longum subsp. longum		5.81	0.20		99		
06511422	**Low discrimination**						**06511422**
					GRA		
Lactobacillus fermentum		89.59	0.46	GLU 99	99		
Bifidobacterium longum subsp. longum		6.72	0.20		99		
Prevotella melaninogenica		3.69	0.03		1		
06511500	**Low discrimination**						**06511500**
					GRA		
Lactobacillus fermentum		70.25	0.58	GLU 99	99		
Bifidobacterium longum subsp. longum		26.65	0.43		99		
Prevotella melaninogenica		3.11	0.15		1		
06511502	**Low discrimination**						**06511502**
Lactobacillus fermentum		71.09	0.58	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		26.97	0.43				
Bifidobacterium adolescentis		1.13	0.21				
06511510	**Good identification**						**06511510**
Bifidobacterium longum subsp. longum		98.46	0.47	GLU 99			
Lactobacillus fermentum		1.54	0.24				
06511512	**Good identification**						**06511512**
Bifidobacterium longum subsp. longum		95.24	0.47	GLU 99			
Bifidobacterium adolescentis		2.85	0.23				
Lactobacillus fermentum		1.49	0.24				
06511520	**Low discrimination**						**06511520**
					GRA		
Prevotella melaninogenica		69.49	0.50	GLU 90	1		
Bifidobacterium longum subsp. longum		24.10	0.54	ARA 1	99		
Lactobacillus fermentum		5.29	0.50		99		
06511522	**Low discrimination**						**06511522**
					GRA		
Bifidobacterium longum subsp. longum		55.47	0.54	GLU 99	99		
Prevotella melaninogenica		30.46	0.37		1		
Lactobacillus fermentum		12.18	0.50		99		
06511530	**Very good identification**						**06511530**
					GRA		
Bifidobacterium longum subsp. longum		98.19	0.58	GLU 99	99		
Prevotella melaninogenica		1.68	0.15		1		
Lactobacillus fermentum		0.13	0.15		99		
06511532	**Very good identification**						**06511532**
					GRA		
Bifidobacterium longum subsp. longum		99.14	0.58	GLU 99	99		
Prevotella melaninogenica		0.32	0.03		1		
Parabacteroides distasonis		0.18	0.15		1		
06512000	**Low discrimination**						**06512000**
					GRA	CAT	
Propionibacterium granulosum		76.52	0.50	GLU 99	99	90	
Prevotella intermedia		23.48	0.33		1	1	
06512002	**Good identification**						**06512002**
					CAT		
Propionibacterium granulosum		98.63	0.39	GLU 99	90		
Propionibacterium propionicum		1.37	0.12		1		
06512020	**Good identification**						**06512020**
Propionibacterium granulosum		100.0	0.39	GLU 99			
06512100	**Acceptable identification**						**06512100**
					CAT		
Propionibacterium propionicum		94.34	0.43	NIT 86	1		
Propionibacterium granulosum		4.16	0.16	GLU 99	90		
Actinomyces naeslundii		1.50	0.23		25		
06512102	**Good identification**						**06512102**
Propionibacterium propionicum		98.95	0.46	NIT 86			
Actinomyces naeslundii		1.05	0.23	GLU 99			
06512140	**Good identification**						**06512140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		100.0	0.43	NIT 86			
06512142	**Good identification**						**06512142**
Propionibacterium propionicum		100.0	0.46	NIT 86			
06513000	**Low discrimination**						**06513000**
					GRA	CAT	
Propionibacterium granulosum		74.22	0.50	GLU 99	99	90	
Prevotella intermedia		22.78	0.33		1	1	
Actinomyces naeslundii		2.29	0.39		99	25	
06513001	**Low discrimination**						**06513001**
Actinomyces naeslundii		82.06	0.42	GLU 99			
Propionibacterium granulosum		17.94	0.16				
06513002	**Low discrimination**						**06513002**
					CAT		
Propionibacterium granulosum		85.08	0.39	GLU 99	90		
Actinomyces naeslundii		10.48	0.39		25		
Lactobacillus fermentum		3.26	0.20		1		
06513003	**Good identification**						**06513003**
Actinomyces naeslundii		100.0	0.42	GLU 99			
06513020	**Low discrimination**						**06513020**
					GRA	CAT	
Propionibacterium granulosum		71.60	0.39	GLU 99	99	90	
Prevotella melaninogenica		19.58	0.15		1	1	
Actinomyces naeslundii		8.82	0.39		99	25	
06513021	**Good identification**						**06513021**
Actinomyces naeslundii		100.0	0.42	GLU 99			
06513023	**Good identification**						**06513023**
Actinomyces naeslundii		100.0	0.42	GLU 99			
06513100	**Low discrimination**						**06513100**
					GRA		
Propionibacterium propionicum		55.78	0.43	NIT 86	99		
Actinomyces naeslundii		21.33	0.46	GLU 99	99		
Prevotella melaninogenica		16.63	0.15		1		
06513101	**Low discrimination**						**06513101**
Actinomyces naeslundii		83.77	0.49	GLU 99			
Propionibacterium propionicum		16.23	0.27				
06513102	**Low discrimination**						**06513102**
Propionibacterium propionicum		74.72	0.46	NIT 86			
Actinomyces naeslundii		19.05	0.46	GLU 99			
Lactobacillus fermentum		3.40	0.24				
06513103	**Low discrimination**						**06513103**
Actinomyces naeslundii		77.49	0.49	GLU 99			
Propionibacterium propionicum		22.51	0.30				
06513120	**Good identification**						**06513120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Prevotella melaninogenica		98.29	0.50	GLU 90	1		
Actinomyces naeslundii		1.27	0.46	MAN 1	99		
Clostridium perfringens		0.36	0.15		99		
06513121	**Low discrimination**						**06513121**
					GRA		
Actinomyces naeslundii		65.80	0.49	GLU 99	99		
Prevotella melaninogenica		34.21	0.15		1		
06513122	**Acceptable identification**						**06513122**
					GRA		
Prevotella melaninogenica		91.56	0.37	GLU 90	1		
Actinomyces naeslundii		6.23	0.46	MAN 1	99		
Clostridium perfringens		0.84	0.09		99		
06513123	**Acceptable identification**						**06513123**
					GRA		
Actinomyces naeslundii		90.99	0.49	GLU 99	99		
Prevotella melaninogenica		9.01	0.03		1		
06513140	**Acceptable identification**						**06513140**
Propionibacterium propionicum		93.68	0.43	NIT 86			
Actinomyces naeslundii		6.32	0.34	GLU 99			
06513142	**Good identification**						**06513142**
Propionibacterium propionicum		95.69	0.46	NIT 86			
Actinomyces naeslundii		4.31	0.34	GLU 99			
06513161	**Good identification**						**06513161**
Actinomyces naeslundii		100.0	0.37	GLU 99			
06513163	**Good identification**						**06513163**
Actinomyces naeslundii		100.0	0.37	GLU 99			
06515000	**Low discrimination**						**06515000**
					COC	GRA	
Lactobacillus fermentum		61.70	0.50	GLU 99	1	99	
Prevotella intermedia		34.01	0.33		1	1	
Gemella morbillorum		1.52	0.11		99	99	
06515002	**Good identification**						**06515002**
Lactobacillus fermentum		100.0	0.50	GLU 99			
06515020	**Low discrimination**						**06515020**
					GRA		
Lactobacillus fermentum		73.08	0.42	GLU 99	99		
Prevotella melaninogenica		26.92	0.15		1		
06515022	**Low discrimination**						**06515022**
					CAT	GRA	
Lactobacillus fermentum		84.38	0.42	GLU 99	1	99	
Prevotella melaninogenica		5.92	0.03		1	1	
Bacteroides fragilis		5.88	-.03		99	1	
06515100	**Low discrimination**						**06515100**
					GRA		
Lactobacillus fermentum		88.38	0.54	GLU 99	99		
Prevotella melaninogenica		6.65	0.15		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		1.77	0.23		1		
06515102	**Acceptable identification**						**06515102**
Lactobacillus fermentum		91.72	0.54	GLU 99			
Bifidobacterium adolescentis		2.47	0.21				
Bifidobacterium longum subsp. longum		1.80	0.17				
06515120	**Low discrimination**						**06515120**
					GRA		
Prevotella melaninogenica		89.76	0.50	GLU 90	1		
Lactobacillus fermentum		4.02	0.46	XYL 1	99		
Bacteroides vulgatus		3.87	0.24		1		
06515122	**Low discrimination**						**06515122**
					CAT	GRA	
Prevotella melaninogenica		38.45	0.37	GLU 90	1	1	
Bacteroides fragilis		38.20	0.32	XYL 1	99	1	
Lactobacillus fermentum		9.03	0.46		1	99	
06515320	**Good identification**						**06515320**
Bacteroides vulgatus		97.25	0.40	GLU 99			
Prevotella melaninogenica		2.53	0.15	ARA 93			
Parabacteroides distasonis		0.22	0.14				
06515322	**Low discrimination**						**06515322**
Parabacteroides distasonis		84.03	0.48	GLU 99			
Prevotella oralis		7.93	0.17				
Bacteroides vulgatus		3.74	0.06				
06515332	**Acceptable identification**						**06515332**
Parabacteroides distasonis		93.86	0.43	GLU 99			
Clostridium clostridioforme		6.14	0.17				
06515400	**Good identification**						**06515400**
Lactobacillus fermentum		96.91	0.50	GLU 99			
Bifidobacterium longum subsp. longum		3.10	0.17				
06515402	**Good identification**						**06515402**
Lactobacillus fermentum		96.91	0.50	GLU 99			
Bifidobacterium longum subsp. longum		3.10	0.17				
06515420	**Low discrimination**						**06515420**
					GRA		
Lactobacillus fermentum		62.72	0.42	GLU 99	99		
Bifidobacterium longum subsp. longum		24.04	0.28		99		
Bacteroides vulgatus		13.24	0.09		1		
06515422	**Low discrimination**						**06515422**
Lactobacillus fermentum		72.29	0.42	GLU 99			
Bifidobacterium longum subsp. longum		27.71	0.28				
06515500	**Low discrimination**						**06515500**
					GRA		
Bifidobacterium longum subsp. longum		64.92	0.52	GLU 99	99		
Lactobacillus fermentum		33.50	0.54		99		
Bacteroides vulgatus		1.45	0.09		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06515502	**Low discrimination**					**06515502**
Bifidobacterium longum subsp. longum		61.93	0.52	GLU 99		
Lactobacillus fermentum		31.96	0.54			
Bifidobacterium adolescentis		5.76	0.36			
06515510	**Good identification**					**06515510**
Bifidobacterium longum subsp. longum		99.70	0.56	GLU 99		
Lactobacillus fermentum		0.31	0.20			
06515512	**Acceptable identification**					**06515512**
Bifidobacterium longum subsp. longum		92.18	0.56	GLU 99		
Bifidobacterium adolescentis		6.16	0.37			
Clostridium clostridioforme		1.38	0.23			
06515520	**Low discrimination**					**06515520**
					GRA	
Bifidobacterium longum subsp. longum		62.36	0.62	GLU 99	99	
Bacteroides vulgatus		34.35	0.43		1	
Lactobacillus fermentum		2.68	0.46		99	
06515522	**Good identification**					**06515522**
					GRA	
Bifidobacterium longum subsp. longum		93.24	0.62	GLU 99	99	
Lactobacillus fermentum		4.01	0.46		99	
Parabacteroides distasonis		1.62	0.36		1	
06515530	**Very good identification**					**06515530**
					GRA	
Bifidobacterium longum subsp. longum		99.67	0.66	GLU 99	99	
Bacteroides vulgatus		0.33	0.09		1	
06515532	**Very good identification**					**06515532**
					GRA	
Bifidobacterium longum subsp. longum		99.09	0.66	GLU 99	99	
Parabacteroides distasonis		0.54	0.31		1	
Clostridium clostridioforme		0.37	0.23		50	
06515712	**Acceptable identification**					**06515712**
Clostridium clostridioforme		91.59	0.38	GLU 88		
Bifidobacterium longum subsp. longum		8.41	0.21	SAL 88		
06515720	**Very good identification**					**06515720**
					GRA	
Bacteroides vulgatus		99.79	0.60	GLU 99	1	
Bifidobacterium longum subsp. longum		0.20	0.28		99	
Parabacteroides distasonis		0.01	0.10		1	
06515722	**Low discrimination**					**06515722**
Parabacteroides distasonis		42.18	0.44	GLU 99		
Bacteroides vulgatus		40.64	0.25			
Clostridium clostridioforme		8.34	0.26			
06515732	**Low discrimination**					**06515732**
					GRA	
Clostridium clostridioforme		50.60	0.38	GLU 88	50	
Parabacteroides distasonis		30.26	0.39	SAL 88	1	
Bifidobacterium longum subsp. longum		18.59	0.32	TRE 88	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06517512	**Low discrimination**						**06517512**
Bifidobacterium adolescentis		84.40	0.36	GLU 99			
Bifidobacterium longum subsp. longum		15.60	0.21				
06525520	**Good identification**						**06525520**
Bifidobacterium longum subsp. longum		100.0	0.35	GLU 99			
06525522	**Good identification**						**06525522**
					GRA		
Bifidobacterium longum subsp. longum		97.21	0.35	GLU 99	99		
Parabacteroides distasonis		2.79	0.13	FRU 90	1		
06525530	**Good identification**						**06525530**
Bifidobacterium longum subsp. longum		100.0	0.39	GLU 99			
06525532	**Good identification**						**06525532**
Bifidobacterium longum subsp. longum		100.0	0.39	GLU 99			
06530000	**Good identification**						**06530000**
					GRA	CAT	
Propionibacterium granulosum		92.28	0.60	GLU 99	99	90	
Prevotella intermedia		7.12	0.33		1	1	
Fusobacterium mortiferum		0.48	0.24		1	5	
06530002	**Good identification**						**06530002**
Propionibacterium granulosum		98.69	0.50	GLU 99			
Actinomyces viscosus		1.31	0.08				
06530010	**Good identification**						**06530010**
Propionibacterium granulosum		100.0	0.45	GLU 99			
06530020	**Good identification**						**06530020**
					CAT		
Propionibacterium granulosum		99.37	0.50	GLU 99	90		
Clostridium perfringens		0.63	0.05		1		
06530022	**Good identification**						**06530022**
Propionibacterium granulosum		100.0	0.39	GLU 99			
06531000	**Good identification**						**06531000**
					GRA	CAT	
Propionibacterium granulosum		90.91	0.60	GLU 99	99	90	
Prevotella intermedia		7.02	0.33		1	1	
Fusobacterium mortiferum		0.71	0.27		1	5	
06531001	**Low discrimination**						**06531001**
Actinomyces naeslundii		53.49	0.42	GLU 99			
Propionibacterium granulosum		46.51	0.26				
06531002	**Acceptable identification**						**06531002**
					CAT		
Propionibacterium granulosum		94.22	0.50	GLU 99	90		
Actinomyces naeslundii		2.92	0.39		25		
Lactobacillus fermentum		0.91	0.20		1		
06531003	**Low discrimination**						**06531003**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		82.15	0.42	GLU 99			
Propionibacterium granulosum		17.86	0.15				
06531010	**Good identification**						**06531010**
Propionibacterium granulosum		100.0	0.45	GLU 99			
06531020	**Low discrimination**						**06531020**
					GRA	CAT	
Propionibacterium granulosum		57.82	0.50	GLU 99	99	90	
Clostridium perfringens		36.41	0.39		99	1	
Prevotella melaninogenica		3.98	0.15		1	1	
06531021	**Low discrimination**						**06531021**
					CAT		
Actinomyces naeslundii		73.84	0.42	GLU 99	25		
Propionibacterium granulosum		16.05	0.15		90		
Clostridium perfringens		10.11	0.05		1		
06531023	**Good identification**						**06531023**
Actinomyces naeslundii		100.0	0.42	GLU 99			
06531101	**Acceptable identification**						**06531101**
					SPO		
Actinomyces naeslundii		90.82	0.49	GLU 99	1		
Clostridium chauvoei		9.18	0.09		99		
06531102	**Low discrimination**						**06531102**
Actinomyces naeslundii		40.00	0.46	GLU 99			
Clostridium perfringens		30.48	0.22				
Lactobacillus fermentum		7.13	0.24				
06531103	**Good identification**						**06531103**
Actinomyces naeslundii		96.48	0.49	GLU 99			
Actinomyces viscosus		3.52	0.01				
06531120	**Low discrimination**						**06531120**
					GRA		
Prevotella melaninogenica		59.00	0.50	GLU 90	1		
Clostridium perfringens		40.02	0.54	TRE 1	99		
Actinomyces naeslundii		0.76	0.46		99		
06531121	**Low discrimination**						**06531121**
					GRA		
Actinomyces naeslundii		52.46	0.49	GLU 99	99		
Prevotella melaninogenica		27.27	0.15		1		
Clostridium perfringens		18.50	0.19		99		
06531122	**Low discrimination**						**06531122**
					GRA		
Clostridium perfringens		58.12	0.48	GLU 99	99		
Prevotella melaninogenica		34.68	0.37		1		
Parabacteroides distasonis		2.71	0.34		1		
06531123	**Low discrimination**						**06531123**
					GRA		
Actinomyces naeslundii		79.05	0.49	GLU 99	99		
Clostridium perfringens		13.12	0.14		99		
Prevotella melaninogenica		7.83	0.03		1		
06531141	**Good identification**						**06531141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GLU 99		
06531143	**Good identification**					**06531143**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06531160	**Acceptable identification**					**06531160**
					GRA	
Clostridium perfringens		91.25	0.41	GLU 99	99	
Prevotella melaninogenica		7.13	0.15		1	
Actinomyces naeslundii		1.61	0.34		99	
06531161	**Low discrimination**					**06531161**
Actinomyces naeslundii		72.43	0.37	GLU 99		
Clostridium perfringens		27.57	0.07			
06531162	**Acceptable identification**					**06531162**
					GRA	
Clostridium perfringens		93.52	0.36	GLU 99	99	
Actinomyces naeslundii		3.52	0.34		99	
Prevotella melaninogenica		2.96	0.03		1	
06531163	**Low discrimination**					**06531163**
Actinomyces naeslundii		84.81	0.37	GLU 99		
Clostridium perfringens		15.19	0.01			
06531322	**Low discrimination**					**06531322**
					GRA	
Parabacteroides distasonis		82.54	0.43	GLU 99	1	
Prevotella oralis		7.95	0.13	XYL 90	1	
Clostridium perfringens		5.96	0.14		99	
06531332	**Acceptable identification**					**06531332**
Parabacteroides distasonis		91.43	0.38	GLU 99		
Clostridium clostridioforme		8.57	0.15	XYL 90		
06531510	**Good identification**					**06531510**
Bifidobacterium longum subsp. longum		100.0	0.37	GLU 99		
06531512	**Low discrimination**					**06531512**
Bifidobacterium longum subsp. longum		82.62	0.37	GLU 99		
Clostridium clostridioforme		10.78	0.20			
Bifidobacterium adolescentis		6.60	0.20			
06531520	**Low discrimination**					**06531520**
					GRA	
Bifidobacterium longum subsp. longum		83.03	0.43	GLU 99	99	
Prevotella melaninogenica		9.67	0.15		1	
Clostridium perfringens		6.56	0.19		99	
06531522	**Low discrimination**					**06531522**
					GRA	
Bifidobacterium longum subsp. longum		84.34	0.43	GLU 99	99	
Parabacteroides distasonis		8.88	0.31		1	
Clostridium perfringens		3.14	0.14		99	
06531530	**Good identification**					**06531530**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		100.0	0.47	GLU 99		
06531532	**Acceptable identification**					**06531532**
					GRA	
Bifidobacterium longum subsp. longum		93.82	0.47	GLU 99	99	
Parabacteroides distasonis		3.12	0.26		1	
Clostridium clostridioforme		3.06	0.20		50	
06531712	**Good identification**					**06531712**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06531722	**Low discrimination**					**06531722**
					GRA	
Parabacteroides distasonis		76.03	0.39	GLU 99	1	
Clostridium clostridioforme		21.54	0.24	XYL 90	50	
Bifidobacterium longum subsp. longum		2.43	0.09		99	
06531732	**Low discrimination**					**06531732**
					GRA	
Clostridium clostridioforme		68.52	0.35	GLU 88	50	
Parabacteroides distasonis		28.59	0.34	SAL 88	1	
Bifidobacterium longum subsp. longum		2.89	0.13	XYL 91	99	
06532000	**Very good identification**					**06532000**
Propionibacterium granulosum		100.0	0.58	GLU 99		
06532002	**Good identification**					**06532002**
Propionibacterium granulosum		100.0	0.48	GLU 99		
06532010	**Good identification**					**06532010**
Propionibacterium granulosum		100.0	0.43	GLU 99		
06532020	**Good identification**					**06532020**
Propionibacterium granulosum		100.0	0.48	GLU 99		
06532022	**Good identification**					**06532022**
Propionibacterium granulosum		100.0	0.37	GLU 99		
06532100	**Low discrimination**					**06532100**
					CAT	
Propionibacterium propionicum		83.38	0.42	NIT 86	1	
Propionibacterium granulosum		13.47	0.24	GLU 99	90	
Actinomyces naeslundii		3.15	0.28		25	
06532102	**Good identification**					**06532102**
					CAT	
Propionibacterium propionicum		95.05	0.45	NIT 86	1	
Propionibacterium granulosum		2.56	0.13	GLU 99	90	
Actinomyces naeslundii		2.40	0.28		25	
06532140	**Good identification**					**06532140**
Propionibacterium propionicum		100.0	0.42	NIT 86		
06532142	**Good identification**					**06532142**
Propionibacterium propionicum		100.0	0.45	NIT 86		
06533000	**Very good identification**					**06533000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		98.05	0.58	GLU 99		
Actinomyces naeslundii		1.95	0.44			
06533001	**Low discrimination**					**06533001**
Actinomyces naeslundii		74.75	0.47	GLU 99		
Propionibacterium granulosum		25.26	0.24			
06533002	**Acceptable identification**					**06533002**
Propionibacterium granulosum		92.62	0.48	GLU 99		
Actinomyces naeslundii		7.38	0.44			
06533003	**Acceptable identification**					**06533003**
Actinomyces naeslundii		92.21	0.47	GLU 99		
Propionibacterium granulosum		7.79	0.13			
06533010	**Good identification**					**06533010**
Propionibacterium granulosum		100.0	0.43	GLU 99		
06533020	**Acceptable identification**					**06533020**
					CAT	
Propionibacterium granulosum		91.90	0.48	GLU 99	90	
Actinomyces naeslundii		7.33	0.44		25	
Clostridium perfringens		0.78	0.05		1	
06533021	**Acceptable identification**					**06533021**
Actinomyces naeslundii		92.21	0.47	GLU 99		
Propionibacterium granulosum		7.79	0.13			
06533022	**Low discrimination**					**06533022**
Propionibacterium granulosum		75.82	0.37	GLU 99		
Actinomyces naeslundii		24.18	0.44			
06533023	**Good identification**					**06533023**
Actinomyces naeslundii		100.0	0.47	GLU 99		
06533100	**Low discrimination**					**06533100**
					CAT	
Propionibacterium propionicum		48.33	0.42	NIT 86	1	
Actinomyces naeslundii		43.86	0.51	GLU 99	25	
Propionibacterium granulosum		7.81	0.24		90	
06533101	**Acceptable identification**					**06533101**
Actinomyces naeslundii		92.45	0.54	GLU 99		
Propionibacterium propionicum		7.55	0.26			
06533102	**Low discrimination**					**06533102**
					CAT	
Propionibacterium propionicum		61.28	0.45	NIT 86	1	
Actinomyces naeslundii		37.07	0.51	GLU 99	25	
Propionibacterium granulosum		1.65	0.13		90	
06533103	**Low discrimination**					**06533103**
Actinomyces naeslundii		89.09	0.54	GLU 99		
Propionibacterium propionicum		10.91	0.29			
06533120	**Low discrimination**					**06533120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		58.18	0.51	GLU 99	99	
Prevotella melaninogenica		23.37	0.15		1	
Clostridium perfringens		15.86	0.19		99	
06533121	**Good identification**					**06533121**
Actinomyces naeslundii		100.0	0.54	GLU 99		
06533122	**Low discrimination**					**06533122**
					GRA	
Actinomyces naeslundii		83.01	0.51	GLU 99	99	
Clostridium perfringens		10.64	0.14		99	
Prevotella melaninogenica		6.35	0.03		1	
06533123	**Good identification**					**06533123**
Actinomyces naeslundii		100.0	0.54	GLU 99		
06533140	**Low discrimination**					**06533140**
Propionibacterium propionicum		86.20	0.42	NIT 86		
Actinomyces naeslundii		13.80	0.38	GLU 99		
06533141	**Low discrimination**					**06533141**
Actinomyces naeslundii		68.37	0.41	GLU 99		
Propionibacterium propionicum		31.63	0.26			
06533142	**Acceptable identification**					**06533142**
Propionibacterium propionicum		90.36	0.45	NIT 86		
Actinomyces naeslundii		9.65	0.38	GLU 99		
06533143	**Low discrimination**					**06533143**
Actinomyces naeslundii		59.04	0.41	GLU 99		
Propionibacterium propionicum		40.97	0.29			
06533160	**Low discrimination**					**06533160**
Actinomyces naeslundii		77.27	0.38	GLU 99		
Clostridium perfringens		22.73	0.07			
06533161	**Good identification**					**06533161**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06533162	**Low discrimination**					**06533162**
Actinomyces naeslundii		87.84	0.38	GLU 99		
Clostridium perfringens		12.16	0.01			
06533163	**Good identification**					**06533163**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06534612	**Good identification**					**06534612**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06534632	**Good identification**					**06534632**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06534712	**Good identification**					**06534712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.44	GLU 88		
06534732	**Good identification**					**06534732**
Clostridium clostridioforme		98.89	0.44	GLU 88		
Parabacteroides distasonis		1.11	0.17	SAL 88		
06535122	**Acceptable identification**					**06535122**
					GRA	
Parabacteroides distasonis		92.03	0.51	GLU 99	1	
Clostridium perfringens		2.22	0.14		99	
Bifidobacterium longum subsp. longum		1.81	0.17		99	
06535132	**Acceptable identification**					**06535132**
					GRA	
Parabacteroides distasonis		90.49	0.46	GLU 99	1	
Bifidobacterium longum subsp. longum		5.61	0.21		99	
Clostridium clostridioforme		3.90	0.17		50	
06535322	**Very good identification**					**06535322**
Parabacteroides distasonis		98.64	0.59	GLU 99		
Clostridium clostridioforme		1.23	0.21			
Prevotella oralis		0.13	-.03			
06535332	**Acceptable identification**					**06535332**
Parabacteroides distasonis		90.47	0.54	GLU 99		
Clostridium clostridioforme		9.53	0.32			
06535500	**Good identification**					**06535500**
Bifidobacterium longum subsp. longum		97.96	0.41	GLU 99		
Lactobacillus fermentum		2.04	0.20			
06535502	**Low discrimination**					**06535502**
Bifidobacterium longum subsp. longum		69.15	0.41	GLU 99		
Bifidobacterium adolescentis		17.16	0.33			
Clostridium clostridioforme		11.37	0.26			
06535510	**Good identification**					**06535510**
Bifidobacterium longum subsp. longum		100.0	0.45	GLU 99		
06535512	**Low discrimination**					**06535512**
Bifidobacterium longum subsp. longum		61.81	0.45	GLU 99		
Clostridium clostridioforme		27.19	0.38			
Bifidobacterium adolescentis		11.01	0.34			
06535520	**Acceptable identification**					**06535520**
					GRA	
Bifidobacterium longum subsp. longum		91.32	0.52	GLU 99	99	
Bacteroides vulgatus		8.38	0.20		1	
Parabacteroides distasonis		0.29	0.13		1	
06535522	**Low discrimination**					**06535522**
					GRA	
Bifidobacterium longum subsp. longum		73.53	0.52	GLU 99	99	
Parabacteroides distasonis		23.21	0.47		1	
Clostridium clostridioforme		3.02	0.26		50	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06535530	**Good identification**					**06535530**
Bifidobacterium longum subsp. longum		100.0	0.56	GLU 99		
06535532	**Low discrimination**					**06535532**
					GRA	
Bifidobacterium longum subsp. longum		82.66	0.56	GLU 99	99	
Clostridium clostridioforme		9.09	0.38		50	
Parabacteroides distasonis		8.25	0.43		1	
06535612	**Good identification**					**06535612**
Clostridium clostridioforme		100.0	0.43	GLU 88		
06535632	**Good identification**					**06535632**
Clostridium clostridioforme		98.76	0.43	GLU 88		
Parabacteroides distasonis		1.24	0.17	SAL 88		
06535702	**Good identification**					**06535702**
					GRA	
Clostridium clostridioforme		96.14	0.41	GLU 88	50	
Parabacteroides distasonis		3.05	0.21	SAL 88	1	
Bifidobacterium longum subsp. longum		0.81	0.07		99	
06535712	**Good identification**					**06535712**
					GRA	
Clostridium clostridioforme		99.32	0.52	GLU 88	50	
Parabacteroides distasonis		0.37	0.17	SAL 88	1	
Bifidobacterium longum subsp. longum		0.31	0.11		99	
06535720	**Good identification**					**06535720**
					GRA	
Bacteroides vulgatus		97.33	0.36	GLU 99	1	
Bifidobacterium longum subsp. longum		1.19	0.17	TRE 4	99	
Parabacteroides distasonis		1.13	0.21		1	
06535722	**Low discrimination**					**06535722**
Parabacteroides distasonis		72.32	0.55	GLU 99		
Clostridium clostridioforme		23.01	0.41			
Bacteroides thetaiotaomicron		3.26	0.16			
06535732	**Low discrimination**					**06535732**
					GRA	
Clostridium clostridioforme		71.63	0.52	GLU 88	50	
Parabacteroides distasonis		26.61	0.51	SAL 88	1	
Bifidobacterium longum subsp. longum		0.90	0.21		99	
06541120	**Good identification**					**06541120**
Prevotella melaninogenica		100.0	0.51	GLU 90		
06541122	**Good identification**					**06541122**
Prevotella melaninogenica		97.64	0.39	GLU 90		
Prevotella oralis		2.36	0.15	FRU 90		
06551001	**Good identification**					**06551001**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06551003	**Low discrimination**					**06551003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		76.78	0.39	GLU 99		
Actinomyces viscosus		23.22	0.06			
06551021	**Low discrimination**					**06551021**
					GRA	
Actinomyces naeslundii		80.60	0.39	GLU 99	99	
Prevotella melaninogenica		19.40	-.01		1	
06551023	**Good identification**					**06551023**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06551101	**Acceptable identification**					**06551101**
					GRA	
Actinomyces naeslundii		92.20	0.47	GLU 99	99	
Prevotella melaninogenica		7.80	-.01		1	
06551103	**Low discrimination**					**06551103**
Actinomyces naeslundii		84.11	0.47	GLU 99		
Actinomyces viscosus		15.89	0.10			
06551120	**Very good identification**					**06551120**
					GRA	
Prevotella melaninogenica		99.50	0.67	GLU 90	1	
Prevotella oralis		0.27	0.27	CEL 10	1	
Clostridium perfringens		0.14	0.25		99	
06551122	**Low discrimination**					**06551122**
					CAT	
Prevotella melaninogenica		78.06	0.55	GLU 90	1	
Prevotella oralis		20.77	0.49	CEL 10	1	
Capnocytophaga ochracea		0.34	0.18		90	
06551322	**Acceptable identification**					**06551322**
Prevotella oralis		90.04	0.42	GLU 99		
Prevotella melaninogenica		7.97	0.21			
Parabacteroides distasonis		1.99	0.27			
06553000	**Low discrimination**					**06553000**
Actinomyces naeslundii		80.17	0.41	GLU 99		
Propionibacterium granulosum		19.83	0.16			
06553001	**Good identification**					**06553001**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06553002	**Good identification**					**06553002**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06553003	**Good identification**					**06553003**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06553020	**Low discrimination**					**06553020**
					GRA	
Actinomyces naeslundii		84.31	0.41	GLU 99	99	
Prevotella melaninogenica		15.69	-.01		1	
06553021	**Good identification**					**06553021**
Actinomyces naeslundii		100.0	0.44	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06553022	**Good identification**					**06553022**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06553023	**Good identification**					**06553023**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06553100	**Acceptable identification**					**06553100**
					GRA	
Actinomyces naeslundii		93.86	0.49	GLU 99	99	
Prevotella melaninogenica		6.14	-.01		1	
06553101	**Good identification**					**06553101**
Actinomyces naeslundii		100.0	0.52	GLU 99		
06553102	**Low discrimination**					**06553102**
Actinomyces naeslundii		89.04	0.49	GLU 99		
Bifidobacterium adolescentis		4.87	0.11			
Actinomyces israelii		3.43	0.15			
06553103	**Good identification**					**06553103**
Actinomyces naeslundii		100.0	0.52	GLU 99		
06553121	**Good identification**					**06553121**
					GRA	
Actinomyces naeslundii		95.82	0.52	GLU 99	99	
Prevotella melaninogenica		4.18	-.01		1	
06553123	**Good identification**					**06553123**
Actinomyces naeslundii		100.0	0.52	GLU 99		
06553140	**Good identification**					**06553140**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06553141	**Good identification**					**06553141**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06553142	**Low discrimination**					**06553142**
Actinomyces naeslundii		85.52	0.36	GLU 99		
Propionibacterium propionicum		14.48	0.12			
06553143	**Good identification**					**06553143**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06553160	**Low discrimination**					**06553160**
					GRA	
Actinomyces naeslundii		72.97	0.36	GLU 99	99	
Prevotella melaninogenica		27.03	-.01		1	
06553161	**Good identification**					**06553161**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06553162	**Good identification**					**06553162**
Actinomyces naeslundii		100.0	0.36	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06553163	**Good identification**					**06553163**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06554712	**Good identification**					**06554712**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06554732	**Good identification**					**06554732**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06555322	**Low discrimination**					**06555322**
Parabacteroides distasonis		59.28	0.43	GLU 99		
Prevotella oralis		36.85	0.27			
Clostridium clostridioforme		2.85	0.15			
06555332	**Low discrimination**					**06555332**
Parabacteroides distasonis		71.08	0.39	GLU 99		
Clostridium clostridioforme		28.92	0.26			
06555502	**Low discrimination**					**06555502**
Bifidobacterium adolescentis		83.77	0.41	GLU 99		
Clostridium clostridioforme		7.95	0.21			
Bifidobacterium longum subsp. longum		4.28	0.17			
06555512	**Low discrimination**					**06555512**
Bifidobacterium adolescentis		70.18	0.43	GLU 99		
Clostridium clostridioforme		24.83	0.32			
Bifidobacterium longum subsp. longum		5.00	0.21			
06555612	**Good identification**					**06555612**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06555632	**Good identification**					**06555632**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06555702	**Good identification**					**06555702**
Clostridium clostridioforme		98.57	0.35	GLU 88		
Bifidobacterium adolescentis		1.43	0.07	SAL 88		
06555712	**Good identification**					**06555712**
Clostridium clostridioforme		99.61	0.47	GLU 88		
Bifidobacterium adolescentis		0.39	0.09	SAL 88		
06555720	**Good identification**					**06555720**
Bacteroides vulgatus		100.0	0.38	GLU 99		
06555722	**Low discrimination**					**06555722**
Clostridium clostridioforme		51.55	0.35	GLU 88		
Parabacteroides distasonis		41.94	0.39	SAL 88		
Bacteroides vulgatus		3.95	0.04	TRE 88		
06555732	**Acceptable identification**					**06555732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		90.88	0.47	GLU 88			
Parabacteroides distasonis		8.74	0.35	SAL 88			
Bacteroides thetaiotaomicron		0.39	-.04	TRE 88			
06557502	**Low discrimination**						**06557502**
Bifidobacterium adolescentis		87.10	0.40	GLU 99			
Actinomyces israelii		12.90	0.32				
06557512	**Good identification**						**06557512**
Bifidobacterium adolescentis		98.22	0.41	GLU 99			
Actinomyces israelii		1.78	0.18				
06570712	**Good identification**						**06570712**
Clostridium clostridioforme		100.0	0.36	GLU 88			
06570732	**Good identification**						**06570732**
Clostridium clostridioforme		100.0	0.36	GLU 88			
06571000	**Low discrimination**						**06571000**
					CAT		
Actinomyces naeslundii		39.68	0.41	GLU 99	25		
Propionibacterium granulosum		39.04	0.26		90		
Lactobacillus jensenii		11.24	0.10		1		
06571001	**Good identification**						**06571001**
Actinomyces naeslundii		100.0	0.44	GLU 99			
06571002	**Low discrimination**						**06571002**
					CAT		
Actinomyces naeslundii		63.84	0.41	GLU 99	25		
Propionibacterium granulosum		15.70	0.15		90		
Lactobacillus jensenii		12.05	0.07		1		
06571003	**Good identification**						**06571003**
Actinomyces naeslundii		100.0	0.44	GLU 99			
06571020	**Low discrimination**						**06571020**
					CAT		
Actinomyces naeslundii		48.29	0.41	GLU 99	25		
Clostridium perfringens		30.85	0.15		1		
Propionibacterium granulosum		11.88	0.15		90		
06571021	**Good identification**						**06571021**
Actinomyces naeslundii		100.0	0.44	GLU 99			
06571022	**Low discrimination**						**06571022**
					GRA		
Actinomyces naeslundii		55.28	0.41	GLU 99	99		
Prevotella oralis		20.82	0.08		1		
Clostridium perfringens		16.62	0.10		99		
06571023	**Good identification**						**06571023**
Actinomyces naeslundii		100.0	0.44	GLU 99			
06571100	**Low discrimination**						**06571100**
					COC	GRA	
Actinomyces naeslundii		59.26	0.49	GLU 99	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		21.57	0.18		99	99	
Fusobacterium mortiferum		12.29	0.25		1	1	
06571101	**Good identification**						**06571101**
Actinomyces naeslundii		100.0	0.52	GLU 99			
06571102	**Low discrimination**						**06571102**
					GRA		
Actinomyces naeslundii		56.08	0.49	GLU 99	99		
Lactobacillus acidophilus		29.85	0.19		99		
Prevotella oralis		6.12	0.06		1		
06571103	**Good identification**						**06571103**
Actinomyces naeslundii		100.0	0.52	GLU 99			
06571121	**Good identification**						**06571121**
					GRA		
Actinomyces naeslundii		95.82	0.52	GLU 99	99		
Prevotella melaninogenica		4.18	-.01		1		
06571123	**Good identification**						**06571123**
Actinomyces naeslundii		100.0	0.52	GLU 99			
06571140	**Good identification**						**06571140**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06571141	**Good identification**						**06571141**
Actinomyces naeslundii		100.0	0.39	GLU 99			
06571142	**Low discrimination**						**06571142**
Actinomyces naeslundii		72.85	0.36	GLU 99			
Lactobacillus acidophilus		27.15	0.04				
06571143	**Good identification**						**06571143**
Actinomyces naeslundii		100.0	0.39	GLU 99			
06571161	**Good identification**						**06571161**
Actinomyces naeslundii		100.0	0.39	GLU 99			
06571162	**Low discrimination**						**06571162**
Actinomyces naeslundii		45.27	0.36	GLU 99			
Clostridium perfringens		37.85	0.12				
Lactobacillus acidophilus		16.87	0.04				
06571163	**Good identification**						**06571163**
Actinomyces naeslundii		100.0	0.39	GLU 99			
06571322	**Low discrimination**						**06571322**
					SPO		
Parabacteroides distasonis		51.44	0.38	GLU 99	1		
Prevotella oralis		32.63	0.22	XYL 90	1		
Clostridium ramosum		10.92	0.14		92		
06571612	**Good identification**						**06571612**
Clostridium clostridioforme		100.0	0.35	GLU 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06571632	**Good identification**					**06571632**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06571712	**Good identification**					**06571712**
Clostridium clostridioforme		100.0	0.44	GLU 88		
06571732	**Acceptable identification**					**06571732**
Clostridium clostridioforme		93.71	0.44	GLU 88		
Parabacteroides distasonis		6.29	0.30	SAL 88		
06573000	**Low discrimination**					**06573000**
Actinomyces naeslundii		72.34	0.46	GLU 99		
Propionibacterium granulosum		27.66	0.24			
06573001	**Good identification**					**06573001**
Actinomyces naeslundii		100.0	0.49	GLU 99		
06573002	**Low discrimination**					**06573002**
					CAT	
Actinomyces naeslundii		87.05	0.46	GLU 99	25	
Propionibacterium granulosum		8.32	0.13		90	
Actinomyces israelii		4.63	0.15		1	
06573003	**Good identification**					**06573003**
Actinomyces naeslundii		100.0	0.49	GLU 99		
06573020	**Acceptable identification**					**06573020**
Actinomyces naeslundii		91.28	0.46	GLU 99		
Propionibacterium granulosum		8.72	0.13			
06573021	**Good identification**					**06573021**
Actinomyces naeslundii		100.0	0.49	GLU 99		
06573022	**Good identification**					**06573022**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06573023	**Good identification**					**06573023**
Actinomyces naeslundii		100.0	0.49	GLU 99		
06573041	**Good identification**					**06573041**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06573043	**Good identification**					**06573043**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06573061	**Good identification**					**06573061**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06573063	**Good identification**					**06573063**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06573100	**Good identification**					**06573100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		95.26	0.53	GLU 99			
Actinomyces israelii		4.75	0.22				
06573101	**Very good identification**						**06573101**
Actinomyces naeslundii		100.0	0.56	GLU 99			
06573102	**Acceptable identification**						**06573102**
Actinomyces naeslundii		91.45	0.53	GLU 99			
Actinomyces israelii		6.83	0.25				
Bifidobacterium adolescentis		1.72	0.08				
06573103	**Very good identification**						**06573103**
Actinomyces naeslundii		100.0	0.56	GLU 99			
06573120	**Low discrimination**						**06573120**
					SPO	GRA	
Actinomyces naeslundii		83.52	0.53	GLU 99	1	99	
Clostridium ramosum		13.67	0.15		92	50	
Prevotella melaninogenica		2.81	-.01		1	1	
06573121	**Very good identification**						**06573121**
Actinomyces naeslundii		100.0	0.56	GLU 99			
06573122	**Low discrimination**						**06573122**
					SPO		
Actinomyces naeslundii		60.44	0.53	GLU 99	1		
Clostridium ramosum		39.56	0.25		92		
06573123	**Very good identification**						**06573123**
Actinomyces naeslundii		100.0	0.56	GLU 99			
06573140	**Acceptable identification**						**06573140**
Actinomyces naeslundii		90.11	0.41	GLU 99			
Actinomyces israelii		9.89	0.15				
06573141	**Good identification**						**06573141**
Actinomyces naeslundii		100.0	0.44	GLU 99			
06573142	**Low discrimination**						**06573142**
Actinomyces naeslundii		85.86	0.41	GLU 99			
Actinomyces israelii		14.14	0.18				
06573143	**Good identification**						**06573143**
Actinomyces naeslundii		100.0	0.44	GLU 99			
06573160	**Good identification**						**06573160**
Actinomyces naeslundii		100.0	0.41	GLU 99			
06573161	**Good identification**						**06573161**
Actinomyces naeslundii		100.0	0.44	GLU 99			
06573162	**Good identification**						**06573162**
Actinomyces naeslundii		100.0	0.41	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06573163	**Good identification**					**06573163**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06574512	**Good identification**					**06574512**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06574532	**Good identification**					**06574532**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06574612	**Good identification**					**06574612**
Clostridium clostridioforme		100.0	0.44	GLU 88		
06574632	**Good identification**					**06574632**
Clostridium clostridioforme		100.0	0.44	GLU 88		
06574702	**Good identification**					**06574702**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06574712	**Good identification**					**06574712**
Clostridium clostridioforme		100.0	0.53	GLU 88		
06574722	**Good identification**					**06574722**
Clostridium clostridioforme		98.49	0.42	GLU 88		
Parabacteroides distasonis		1.51	0.17	SAL 88		
06574732	**Good identification**					**06574732**
Clostridium clostridioforme		99.82	0.53	GLU 88		
Parabacteroides distasonis		0.18	0.12	SAL 88		
06575122	**Acceptable identification**					**06575122**
Parabacteroides distasonis		90.58	0.46	GLU 99		
Prevotella oralis		5.52	0.13			
Clostridium clostridioforme		2.87	0.15			
06575132	**Low discrimination**					**06575132**
Parabacteroides distasonis		78.87	0.42	GLU 99		
Clostridium clostridioforme		21.13	0.26			
06575312	**Good identification**					**06575312**
Clostridium clostridioforme		98.48	0.41	GLU 88		
Parabacteroides distasonis		1.52	0.16	SAL 88		
06575322	**Acceptable identification**					**06575322**
Parabacteroides distasonis		92.07	0.54	GLU 99		
Clostridium clostridioforme		7.13	0.30			
Prevotella oralis		0.80	0.06			
06575332	**Low discrimination**					**06575332**
Parabacteroides distasonis		60.43	0.50	GLU 99		
Clostridium clostridioforme		39.57	0.41			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06575412	**Good identification**					**06575412**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06575432	**Good identification**					**06575432**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06575502	**Low discrimination**					**06575502**
					SPO	
Clostridium clostridioforme		44.50	0.35	GLU 88	66	
Bifidobacterium adolescentis		42.63	0.38	SAL 88	1	
Clostridium butyricum		6.91	0.10	RHA 88	99	
06575512	**Low discrimination**					**06575512**
Clostridium clostridioforme		78.89	0.47	GLU 88		
Bifidobacterium adolescentis		20.27	0.40	SAL 88		
Bifidobacterium longum subsp. longum		0.54	0.11	RHA 88		
06575522	**Low discrimination**					**06575522**
					GRA	
Parabacteroides distasonis		52.39	0.43	GLU 99	1	
Clostridium clostridioforme		42.40	0.35		50	
Bifidobacterium longum subsp. longum		3.11	0.17		99	
06575532	**Low discrimination**					**06575532**
					GRA	
Clostridium clostridioforme		84.91	0.47	GLU 88	50	
Parabacteroides distasonis		12.40	0.38	SAL 88	1	
Bifidobacterium longum subsp. longum		2.33	0.21	RHA 88	99	
06575602	**Good identification**					**06575602**
Clostridium clostridioforme		100.0	0.41	GLU 88		
06575612	**Good identification**					**06575612**
Clostridium clostridioforme		100.0	0.52	GLU 88		
06575622	**Good identification**					**06575622**
Clostridium clostridioforme		98.32	0.41	GLU 88		
Parabacteroides distasonis		1.68	0.17	SAL 88		
06575632	**Good identification**					**06575632**
Clostridium clostridioforme		99.80	0.52	GLU 88		
Parabacteroides distasonis		0.20	0.12	SAL 88		
06575702	**Good identification**					**06575702**
Clostridium clostridioforme		99.49	0.50	GLU 88		
Parabacteroides distasonis		0.51	0.17	SAL 88		
06575712	**Very good identification**					**06575712**
Clostridium clostridioforme		99.94	0.61	GLU 88		
Parabacteroides distasonis		0.06	0.12	SAL 88		
06575722	**Low discrimination**					**06575722**
Clostridium clostridioforme		62.15	0.50	GLU 88		
Parabacteroides distasonis		31.41	0.51	SAL 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		5.58	0.22				
06575732	**Good identification**						**06575732**
Clostridium clostridioforme		93.51	0.61	GLU 88			
Parabacteroides distasonis		5.59	0.46	SAL 88			
Bacteroides thetaiotaomicron		0.72	0.15				
06577100	**Good identification**						**06577100**
Actinomyces israelii		100.0	0.40	GLU 99			
06577102	**Low discrimination**						**06577102**
Actinomyces israelii		87.23	0.43	GLU 99			
Bifidobacterium adolescentis		12.77	0.23	SAL 92			
06577142	**Good identification**						**06577142**
Actinomyces israelii		95.56	0.36	GLU 99			
Bifidobacterium adolescentis		4.44	0.07	SAL 92			
06577500	**Acceptable identification**						**06577500**
					SPO		
Actinomyces israelii		90.78	0.39	GLU 99	1		
Clostridium butyricum		9.22	0.01	SAL 92	99		
06577502	**Low discrimination**						**06577502**
					SPO		
Bifidobacterium adolescentis		53.84	0.37	GLU 99	1		
Actinomyces israelii		44.98	0.42		1		
Clostridium butyricum		1.19	-.07		99		
06577512	**Low discrimination**						**06577512**
Bifidobacterium adolescentis		84.96	0.38	GLU 99			
Actinomyces israelii		8.68	0.28				
Clostridium clostridioforme		4.08	0.12				
06610000	**Good identification**						**06610000**
					GRA	CAT	
Fusobacterium mortiferum		97.33	0.54	GLU 99	1	5	
Propionibacterium granulosum		2.67	0.24		99	90	
06610002	**Acceptable identification**						**06610002**
					GRA	CAT	
Fusobacterium mortiferum		94.18	0.37	GLU 99	1	5	
Propionibacterium granulosum		5.82	0.13	BGL 10	99	90	
06610100	**Very good identification**						**06610100**
Fusobacterium mortiferum		100.0	0.60	GLU 99			
06610102	**Good identification**						**06610102**
Fusobacterium mortiferum		100.0	0.44	GLU 99			
06611000	**Very good identification**						**06611000**
					GRA	CAT	
Fusobacterium mortiferum		97.99	0.57	GLU 99	1	5	
Propionibacterium granulosum		1.79	0.24		99	90	
Actinomyces naeslundii		0.22	0.23		99	25	
06611002	**Acceptable identification**						**06611002**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		94.25	0.40	GLU 99	1	5	
Propionibacterium granulosum		3.88	0.13	BGL 10	99	90	
Actinomyces naeslundii		1.87	0.23		99	25	
06611100	**Very good identification**						**06611100**
					GRA		
Fusobacterium mortiferum		99.60	0.63	GLU 99	1		
Actinomyces naeslundii		0.27	0.30		99		
Lactobacillus fermentum		0.14	0.15		99		
06611102	**Good identification**						**06611102**
					GRA		
Fusobacterium mortiferum		95.72	0.47	GLU 99	1		
Actinomyces naeslundii		2.32	0.30	BGL 10	99		
Lactobacillus fermentum		1.18	0.15		99		
06613101	**Good identification**						**06613101**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06613102	**Low discrimination**						**06613102**
					GRA		
Actinomyces naeslundii		82.30	0.35	GLU 99	99		
Fusobacterium mortiferum		17.70	0.12	MLT 94	1		
06613103	**Good identification**						**06613103**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06613120	**Good identification**						**06613120**
Actinomyces naeslundii		100.0	0.35	GLU 99			
06613121	**Good identification**						**06613121**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06613122	**Good identification**						**06613122**
Actinomyces naeslundii		100.0	0.35	GLU 99			
06613123	**Good identification**						**06613123**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06614100	**Good identification**						**06614100**
Fusobacterium mortiferum		100.0	0.41	GLU 99			
06615000	**Good identification**						**06615000**
Fusobacterium mortiferum		100.0	0.37	GLU 99			
06615100	**Good identification**						**06615100**
Fusobacterium mortiferum		100.0	0.44	GLU 99			
06615322	**Good identification**						**06615322**
Parabacteroides distasonis		100.0	0.39	GLU 99			
06615722	**Good identification**						**06615722**
Parabacteroides distasonis		95.43	0.35	GLU 99			
Clostridium clostridioforme		4.57	0.07				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06630000	**Low discrimination**						**06630000**
					GRA	CAT	
Fusobacterium mortiferum		78.39	0.45	GLU 99	1	5	
Propionibacterium granulosum		21.61	0.32		99	90	
06630100	**Good identification**						**06630100**
Fusobacterium mortiferum		100.0	0.51	GLU 99			
06631000	**Low discrimination**						**06631000**
					GRA	CAT	
Fusobacterium mortiferum		83.47	0.48	GLU 99	1	5	
Propionibacterium granulosum		15.34	0.32		99	90	
Actinomyces naeslundii		1.20	0.28		99	25	
06631100	**Good identification**						**06631100**
					GRA		
Fusobacterium mortiferum		98.28	0.54	GLU 99	1		
Actinomyces naeslundii		1.72	0.35		99		
06631101	**Low discrimination**						**06631101**
					GRA		
Actinomyces naeslundii		72.18	0.38	GLU 99	99		
Fusobacterium mortiferum		27.82	0.20	MLT 94	1		
06631102	**Low discrimination**						**06631102**
					GRA		
Fusobacterium mortiferum		84.14	0.38	GLU 99	1		
Actinomyces naeslundii		13.23	0.35	BGL 10	99		
Lactobacillus acidophilus		2.63	-.01		99		
06631103	**Good identification**						**06631103**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06631120	**Low discrimination**						**06631120**
					GRA		
Actinomyces naeslundii		46.25	0.35	GLU 99	99		
Clostridium perfringens		27.03	0.09	MLT 94	99		
Fusobacterium mortiferum		26.73	0.20		1		
06631121	**Good identification**						**06631121**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06631123	**Good identification**						**06631123**
Actinomyces naeslundii		100.0	0.38	GLU 99			
06633001	**Good identification**						**06633001**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06633003	**Good identification**						**06633003**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06633021	**Good identification**						**06633021**
Actinomyces naeslundii		100.0	0.36	GLU 99			
06633023	**Good identification**						**06633023**
Actinomyces naeslundii		100.0	0.36	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06633100	**Low discrimination**					**06633100**
					GRA	
Actinomyces naeslundii		77.06	0.40	GLU 99	99	
Fusobacterium mortiferum		22.95	0.20	MLT 94	1	
06633101	**Good identification**					**06633101**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06633102	**Good identification**					**06633102**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06633103	**Good identification**					**06633103**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06633120	**Good identification**					**06633120**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06633121	**Good identification**					**06633121**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06633122	**Good identification**					**06633122**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06633123	**Good identification**					**06633123**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06635122	**Good identification**					**06635122**
Parabacteroides distasonis		100.0	0.42	GLU 99		
06635132	**Good identification**					**06635132**
Parabacteroides distasonis		100.0	0.37	GLU 99		
06635322	**Good identification**					**06635322**
Parabacteroides distasonis		100.0	0.50	GLU 99		
06635332	**Good identification**					**06635332**
Parabacteroides distasonis		97.51	0.46	GLU 99		
Clostridium clostridioforme		2.49	0.13			
06635522	**Good identification**					**06635522**
Parabacteroides distasonis		96.94	0.38	GLU 99		
Clostridium clostridioforme		3.06	0.07			
06635722	**Acceptable identification**					**06635722**
Parabacteroides distasonis		92.84	0.47	GLU 99		
Clostridium clostridioforme		7.16	0.22			
06635732	**Low discrimination**					**06635732**
Parabacteroides distasonis		60.51	0.42	GLU 99		
Clostridium clostridioforme		39.49	0.33			
06650000	**Good identification**					**06650000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.45	GLU 99		
06650100	**Good identification**					**06650100**
Fusobacterium mortiferum		100.0	0.52	GLU 99		
06650102	**Good identification**					**06650102**
Fusobacterium mortiferum		100.0	0.35	GLU 99		
06651000	**Good identification**					**06651000**
					GRA	
Fusobacterium mortiferum		95.63	0.48	GLU 99	1	
Lactobacillus catenaformis		2.86	0.12		99	
Actinomyces naeslundii		0.84	0.25		99	
06651100	**Good identification**					**06651100**
					GRA	
Fusobacterium mortiferum		97.10	0.55	GLU 99	1	
Lactobacillus catenaformis		1.86	0.15		99	
Actinomyces naeslundii		1.04	0.33		99	
06651101	**Low discrimination**					**06651101**
					GRA	
Actinomyces naeslundii		61.36	0.36	GLU 99	99	
Fusobacterium mortiferum		38.64	0.21	MLT 94	1	
06651102	**Low discrimination**					**06651102**
					GRA	
Fusobacterium mortiferum		46.39	0.38	GLU 99	1	
Lactobacillus acidophilus		38.47	0.24	BGL 10	99	
Lactobacillus catenaformis		8.01	0.15		99	
06651103	**Good identification**					**06651103**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06651121	**Good identification**					**06651121**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06651123	**Good identification**					**06651123**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06653100	**Low discrimination**					**06653100**
					GRA	
Actinomyces naeslundii		67.27	0.37	GLU 99	99	
Fusobacterium mortiferum		32.74	0.21	MLT 94	1	
06653101	**Good identification**					**06653101**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06653102	**Low discrimination**					**06653102**
Actinomyces naeslundii		81.06	0.37	GLU 99		
Lactobacillus acidophilus		18.94	0.02	MLT 94		
06653103	**Good identification**					**06653103**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06653120	**Good identification**					**06653120**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GLU 99		
06653121	**Good identification**					**06653121**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06653122	**Low discrimination**					**06653122**
					SPO	
Actinomyces naeslundii		62.81	0.37	GLU 99	1	
Clostridium ramosum		22.51	0.05	MLT 94	92	
Lactobacillus acidophilus		14.68	0.02		1	
06653123	**Good identification**					**06653123**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06655100	**Good identification**					**06655100**
Fusobacterium mortiferum		100.0	0.35	GLU 99		
06670000	**Good identification**					**06670000**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
06670100	**Good identification**					**06670100**
Fusobacterium mortiferum		100.0	0.43	GLU 99		
06671000	**Low discrimination**					**06671000**
					GRA	
Fusobacterium mortiferum		88.63	0.39	GLU 99	1	
Lactobacillus jensenii		6.32	0.10		99	
Actinomyces naeslundii		5.05	0.30		99	
06671100	**Acceptable identification**					**06671100**
					GRA	
Fusobacterium mortiferum		92.60	0.46	GLU 99	1	
Actinomyces naeslundii		6.44	0.37		99	
Lactobacillus acidophilus		0.97	-.01		99	
06671101	**Acceptable identification**					**06671101**
					GRA	
Actinomyces naeslundii		91.17	0.41	GLU 99	99	
Fusobacterium mortiferum		8.83	0.12	MLT 94	1	
06671103	**Acceptable identification**					**06671103**
Actinomyces naeslundii		90.90	0.41	GLU 99		
Lactobacillus acidophilus		9.10	-.01	MLT 94		
06671120	**Low discrimination**					**06671120**
					SPO	
Actinomyces naeslundii		64.80	0.37	GLU 99	1	
Clostridium ramosum		16.06	0.02	MLT 94	92	
Lactobacillus acidophilus		9.73	-.01		1	
06671121	**Good identification**					**06671121**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06671123	**Acceptable identification**					**06671123**
Actinomyces naeslundii		90.90	0.41	GLU 99		
Lactobacillus acidophilus		9.10	-.01	MLT 94		
06673001	**Good identification**					**06673001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	GLU 99		
06673003	**Good identification**					**06673003**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06673021	**Good identification**					**06673021**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06673023	**Good identification**					**06673023**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06673100	**Acceptable identification**					**06673100**
					GRA	
Actinomyces naeslundii		93.04	0.42	GLU 99	99	
Fusobacterium mortiferum		6.97	0.12	MLT 94	1	
06673101	**Good identification**					**06673101**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06673102	**Low discrimination**					**06673102**
Actinomyces naeslundii		71.28	0.42	GLU 99		
Lactobacillus acidophilus		28.72	0.11	MLT 94		
06673103	**Good identification**					**06673103**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06673120	**Low discrimination**					**06673120**
					SPO	
Actinomyces naeslundii		58.03	0.42	GLU 99	1	
Clostridium ramosum		41.97	0.15	MLT 94	92	
06673121	**Good identification**					**06673121**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06673123	**Good identification**					**06673123**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06675122	**Good identification**					**06675122**
					GRA	
Parabacteroides distasonis		97.42	0.37	GLU 99	1	
Lactobacillus acidophilus		2.58	-.01		99	
06675322	**Good identification**					**06675322**
Parabacteroides distasonis		98.16	0.46	GLU 99		
Clostridium clostridioforme		1.84	0.10			
06675332	**Low discrimination**					**06675332**
Parabacteroides distasonis		86.30	0.41	GLU 99		
Clostridium clostridioforme		13.70	0.22			
06675712	**Good identification**					**06675712**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06675722	**Low discrimination**					**06675722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		67.58	0.42	GLU 99		
Clostridium clostridioforme		32.42	0.31			
06675732	**Low discrimination**					**06675732**
Clostridium clostridioforme		80.23	0.42	GLU 88		
Parabacteroides distasonis		19.77	0.37	MLT 99		
06703101	**Good identification**					**06703101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06703103	**Good identification**					**06703103**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06703121	**Good identification**					**06703121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06703123	**Good identification**					**06703123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06710100	**Good identification**					**06710100**
					GRA	
Fusobacterium mortiferum		97.91	0.44	GLU 99	1	
Actinomyces naeslundii		2.09	0.27	MLT 11	99	
06711001	**Good identification**					**06711001**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06711002	**Low discrimination**					**06711002**
					GRA	
Actinomyces naeslundii		44.42	0.43	GLU 99	99	
Actinomyces viscosus		22.60	0.14		99	
Fusobacterium mortiferum		17.66	0.25		1	
06711003	**Low discrimination**					**06711003**
Actinomyces naeslundii		89.84	0.46	GLU 99		
Actinomyces viscosus		10.16	0.06			
06711021	**Good identification**					**06711021**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06711023	**Good identification**					**06711023**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06711100	**Low discrimination**					**06711100**
					GRA	
Fusobacterium mortiferum		60.84	0.47	GLU 99	1	
Actinomyces naeslundii		20.74	0.51	MLT 11	99	
Prevotella melaninogenica		8.85	0.15		1	
06711101	**Good identification**					**06711101**
					GRA	
Actinomyces naeslundii		98.06	0.54	GLU 99	99	
Fusobacterium mortiferum		1.94	0.13		1	
06711103	**Acceptable identification**					**06711103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.16	0.54	GLU 99		
Actinomyces viscosus		6.51	0.10			
Bifidobacterium longum subsp. infantis		1.33	-.07			
06711120	**Low discrimination**					**06711120**
					GRA	
Prevotella melaninogenica		87.71	0.50	GLU 90	1	
Clostridium perfringens		8.01	0.39	SAL 1	99	
Actinomyces naeslundii		2.08	0.51		99	
06711121	**Low discrimination**					**06711121**
					GRA	
Actinomyces naeslundii		76.30	0.54	GLU 99	99	
Prevotella melaninogenica		21.72	0.15		1	
Clostridium perfringens		1.98	0.04		99	
06711122	**Low discrimination**					**06711122**
					GRA	
Prevotella oralis		44.10	0.44	GLU 99	1	
Prevotella melaninogenica		30.05	0.37		1	
Bifidobacterium longum subsp. infantis		14.23	0.32		99	
06711123	**Low discrimination**					**06711123**
					GRA	
Actinomyces naeslundii		86.25	0.54	GLU 99	99	
Prevotella oralis		6.86	0.09		1	
Prevotella melaninogenica		4.68	0.03		1	
06711140	**Low discrimination**					**06711140**
					GRA	
Actinomyces naeslundii		77.11	0.38	GLU 99	99	
Fusobacterium mortiferum		12.95	0.13		1	
Clostridium perfringens		9.94	0.00		99	
06711141	**Good identification**					**06711141**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06711142	**Low discrimination**					**06711142**
Actinomyces naeslundii		89.07	0.38	GLU 99		
Bifidobacterium longum subsp. infantis		10.93	-.07			
06711143	**Good identification**					**06711143**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06711161	**Good identification**					**06711161**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06711163	**Good identification**					**06711163**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06711322	**Low discrimination**					**06711322**
Prevotella oralis		89.90	0.37	GLU 99		
Parabacteroides distasonis		7.97	0.32			
Prevotella melaninogenica		1.44	0.03			
06711512	**Low discrimination**					**06711512**
Bifidobacterium adolescentis		79.88	0.36	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		15.36	0.20	XYL 87		
Bifidobacterium longum subsp. longum		4.76	0.13			
06711712	**Good identification**					**06711712**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06711732	**Acceptable identification**					**06711732**
Clostridium clostridioforme		90.97	0.35	GLU 88		
Parabacteroides distasonis		9.03	0.24	TRE 88		
06712101	**Good identification**					**06712101**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06712103	**Good identification**					**06712103**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06712121	**Good identification**					**06712121**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06712123	**Good identification**					**06712123**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06713000	**Acceptable identification**					**06713000**
Actinomyces naeslundii		91.53	0.48	GLU 99		
Propionibacterium granulosum		8.47	0.16			
06713001	**Good identification**					**06713001**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06713002	**Good identification**					**06713002**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06713003	**Good identification**					**06713003**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06713020	**Good identification**					**06713020**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06713021	**Good identification**					**06713021**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06713022	**Good identification**					**06713022**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06713023	**Good identification**					**06713023**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06713040	**Good identification**					**06713040**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06713041	**Good identification**					**06713041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	GLU 99		
06713042	**Good identification**					**06713042**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06713043	**Good identification**					**06713043**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06713060	**Good identification**					**06713060**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06713061	**Good identification**					**06713061**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06713062	**Good identification**					**06713062**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06713063	**Good identification**					**06713063**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06713100	**Good identification**					**06713100**
					GRA	
Actinomyces naeslundii		98.50	0.56	GLU 99	99	
Fusobacterium mortiferum		1.50	0.13		1	
06713101	**Very good identification**					**06713101**
Actinomyces naeslundii		100.0	0.59	GLU 99		
06713102	**Acceptable identification**					**06713102**
Actinomyces naeslundii		91.80	0.56	GLU 99		
Bifidobacterium adolescentis		5.01	0.19			
Actinomyces israelii		1.15	0.14			
06713103	**Very good identification**					**06713103**
Actinomyces naeslundii		100.0	0.59	GLU 99		
06713111	**Good identification**					**06713111**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06713113	**Good identification**					**06713113**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06713120	**Low discrimination**					**06713120**
					GRA	
Actinomyces naeslundii		80.64	0.56	GLU 99	99	
Prevotella melaninogenica		17.74	0.15		1	
Clostridium perfringens		1.62	0.04		99	
06713121	**Very good identification**					**06713121**
Actinomyces naeslundii		100.0	0.59	GLU 99		
06713122	**Low discrimination**					**06713122**
					GRA	
Actinomyces naeslundii		86.69	0.56	GLU 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		5.33	0.09		1	
Prevotella melaninogenica		3.63	0.03		1	
06713123	**Very good identification**					**06713123**
Actinomyces naeslundii		100.0	0.59	GLU 99		
06713131	**Good identification**					**06713131**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06713133	**Good identification**					**06713133**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06713140	**Good identification**					**06713140**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06713141	**Good identification**					**06713141**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06713142	**Acceptable identification**					**06713142**
Actinomyces naeslundii		94.05	0.43	GLU 99		
Propionibacterium propionicum		5.95	0.12			
06713143	**Good identification**					**06713143**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06713160	**Good identification**					**06713160**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06713161	**Good identification**					**06713161**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06713162	**Good identification**					**06713162**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06713163	**Good identification**					**06713163**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06714612	**Good identification**					**06714612**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06714632	**Good identification**					**06714632**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06714712	**Good identification**					**06714712**
Clostridium clostridioforme		100.0	0.44	GLU 88		
06714732	**Good identification**					**06714732**
Clostridium clostridioforme		100.0	0.44	GLU 88		
06715112	**Low discrimination**					**06715112**
Bifidobacterium adolescentis		88.96	0.36	GLU 99		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		11.04	0.17	ARA 87		
06715132	**Low discrimination**					**06715132**
					GRA	
Parabacteroides distasonis		81.66	0.36	GLU 99	1	
Clostridium clostridioforme		14.80	0.17		50	
Bifidobacterium longum subsp. infantis		3.55	-.04		99	
06715322	**Low discrimination**					**06715322**
Parabacteroides distasonis		81.25	0.48	GLU 99		
Prevotella oralis		12.59	0.22			
Clostridium clostridioforme		4.25	0.21			
06715332	**Low discrimination**					**06715332**
Parabacteroides distasonis		69.30	0.44	GLU 99		
Clostridium clostridioforme		30.70	0.32			
06715502	**Low discrimination**					**06715502**
Bifidobacterium adolescentis		89.77	0.48	GLU 99		
Clostridium clostridioforme		7.00	0.26			
Bifidobacterium longum subsp. longum		1.72	0.17			
06715512	**Low discrimination**					**06715512**
Bifidobacterium adolescentis		75.91	0.50	GLU 99		
Clostridium clostridioforme		22.06	0.38			
Bifidobacterium longum subsp. longum		2.03	0.21			
06715522	**Low discrimination**					**06715522**
					GRA	
Parabacteroides distasonis		44.08	0.37	GLU 99	1	
Clostridium clostridioforme		24.13	0.26		50	
Bifidobacterium longum subsp. longum		23.72	0.28		99	
06715532	**Low discrimination**					**06715532**
					GRA	
Clostridium clostridioforme		61.80	0.38	GLU 88	50	
Bifidobacterium longum subsp. longum		22.71	0.32	TRE 88	99	
Parabacteroides distasonis		13.34	0.32	RHA 88	1	
06715612	**Good identification**					**06715612**
Clostridium clostridioforme		100.0	0.43	GLU 88		
06715632	**Good identification**					**06715632**
Clostridium clostridioforme		100.0	0.43	GLU 88		
06715702	**Acceptable identification**					**06715702**
Clostridium clostridioforme		94.71	0.41	GLU 88		
Eubacterium contortum		2.91	0.10	TRE 88		
Bifidobacterium adolescentis		1.67	0.14			
06715712	**Good identification**					**06715712**
Clostridium clostridioforme		99.53	0.52	GLU 88		
Bifidobacterium adolescentis		0.47	0.16	TRE 88		
06715720	**Good identification**					**06715720**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		99.71	0.49	GLU 99	1	
Eubacterium contortum		0.19	0.10		99	
Clostridium clostridioforme		0.06	0.07		50	
06715722	**Low discrimination**					**06715722**
Clostridium clostridioforme		52.26	0.41	GLU 88		
Parabacteroides distasonis		39.05	0.45	TRE 88		
Bacteroides vulgatus		8.68	0.15			
06715732	**Acceptable identification**					**06715732**
Clostridium clostridioforme		91.88	0.52	GLU 88		
Parabacteroides distasonis		8.12	0.40	TRE 88		
06717502	**Good identification**					**06717502**
Bifidobacterium adolescentis		95.42	0.47	GLU 99		
Actinomyces israelii		4.59	0.31			
06717512	**Good identification**					**06717512**
Bifidobacterium adolescentis		99.42	0.48	GLU 99		
Actinomyces israelii		0.58	0.17			
06721101	**Good identification**					**06721101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06721103	**Good identification**					**06721103**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06721121	**Good identification**					**06721121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06721123	**Good identification**					**06721123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06723100	**Good identification**					**06723100**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06723101	**Good identification**					**06723101**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06723102	**Good identification**					**06723102**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06723103	**Good identification**					**06723103**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06723120	**Good identification**					**06723120**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06723121	**Good identification**					**06723121**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06723122	**Good identification**					**06723122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	GLU 99			
06723123	**Good identification**						**06723123**
Actinomyces naeslundii		100.0	0.42	GLU 99			
06730100	**Low discrimination**						**06730100**
					GRA		
Fusobacterium mortiferum		87.84	0.35	GLU 99	1		
Actinomyces naeslundii		12.16	0.32	MLT 11	99		
06730101	**Good identification**						**06730101**
Actinomyces naeslundii		100.0	0.35	GLU 99			
06730103	**Good identification**						**06730103**
Actinomyces naeslundii		100.0	0.35	GLU 99			
06730121	**Good identification**						**06730121**
Actinomyces naeslundii		100.0	0.35	GLU 99			
06730123	**Good identification**						**06730123**
Actinomyces naeslundii		100.0	0.35	GLU 99			
06730712	**Good identification**						**06730712**
Clostridium clostridioforme		100.0	0.42	GLU 88			
06730732	**Good identification**						**06730732**
Clostridium clostridioforme		100.0	0.42	GLU 88			
06731000	**Low discrimination**						**06731000**
					GRA	CAT	
Actinomyces naeslundii		48.34	0.48	GLU 99	99	25	
Fusobacterium mortiferum		26.62	0.32		1	5	
Propionibacterium granulosum		17.78	0.26		99	90	
06731001	**Good identification**						**06731001**
Actinomyces naeslundii		100.0	0.51	GLU 99			
06731002	**Low discrimination**						**06731002**
					CAT		
Actinomyces naeslundii		78.56	0.48	GLU 99	25		
Propionibacterium granulosum		7.22	0.15		90		
Lactobacillus jensenii		5.55	0.07		1		
06731003	**Good identification**						**06731003**
Actinomyces naeslundii		100.0	0.51	GLU 99			
06731021	**Good identification**						**06731021**
Actinomyces naeslundii		100.0	0.51	GLU 99			
06731022	**Low discrimination**						**06731022**
					GRA		
Actinomyces naeslundii		55.73	0.48	GLU 99	99		
Clostridium perfringens		37.59	0.23		99		
Prevotella oralis		3.94	0.03		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06731023	**Good identification**					**06731023**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06731040	**Good identification**					**06731040**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06731041	**Good identification**					**06731041**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06731042	**Good identification**					**06731042**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06731043	**Good identification**					**06731043**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06731061	**Good identification**					**06731061**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06731062	**Low discrimination**					**06731062**
Actinomyces naeslundii		57.87	0.35	GLU 99		
Clostridium perfringens		42.13	0.11			
06731063	**Good identification**					**06731063**
Actinomyces naeslundii		100.0	0.38	GLU 99		
06731100	**Low discrimination**					**06731100**
					GRA	
Actinomyces naeslundii		60.87	0.56	GLU 99	99	
Fusobacterium mortiferum		27.48	0.38		1	
Clostridium perfringens		6.95	0.17		99	
06731101	**Very good identification**					**06731101**
Actinomyces naeslundii		100.0	0.59	GLU 99		
06731102	**Low discrimination**					**06731102**
Actinomyces naeslundii		71.18	0.56	GLU 99		
Lactobacillus acidophilus		14.16	0.19			
Clostridium perfringens		3.83	0.12			
06731103	**Very good identification**					**06731103**
Actinomyces naeslundii		100.0	0.59	GLU 99		
06731111	**Good identification**					**06731111**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06731113	**Good identification**					**06731113**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06731120	**Low discrimination**					**06731120**
					GRA	
Clostridium perfringens		74.33	0.43	GLU 99	99	
Actinomyces naeslundii		20.13	0.56		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		4.43	0.15		1	
06731121	**Very good identification**					**06731121**
Actinomyces naeslundii		97.57	0.59	GLU 99		
Clostridium perfringens		2.43	0.09			
06731122	**Low discrimination**					**06731122**
					GRA	
Clostridium perfringens		45.47	0.38	GLU 99	99	
Actinomyces naeslundii		26.16	0.56		99	
Prevotella oralis		10.17	0.23		1	
06731123	**Very good identification**					**06731123**
Actinomyces naeslundii		98.84	0.59	GLU 99		
Clostridium perfringens		1.16	0.03			
06731131	**Good identification**					**06731131**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06731133	**Good identification**					**06731133**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06731140	**Low discrimination**					**06731140**
Actinomyces naeslundii		89.03	0.43	GLU 99		
Clostridium perfringens		10.97	0.05			
06731141	**Good identification**					**06731141**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06731142	**Low discrimination**					**06731142**
Actinomyces naeslundii		87.77	0.43	GLU 99		
Lactobacillus acidophilus		12.23	0.04			
06731143	**Good identification**					**06731143**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06731161	**Good identification**					**06731161**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06731163	**Good identification**					**06731163**
Actinomyces naeslundii		100.0	0.46	GLU 99		
06731322	**Low discrimination**					**06731322**
					SPO	
Parabacteroides distasonis		73.02	0.43	GLU 99	1	
Prevotella oralis		11.54	0.17	XYL 90	1	
Clostridium ramosum		7.70	0.14		92	
06731332	**Low discrimination**					**06731332**
Parabacteroides distasonis		61.18	0.39	GLU 99		
Clostridium clostridioforme		38.82	0.30	XYL 90		
06731512	**Low discrimination**					**06731512**
Clostridium clostridioforme		67.90	0.35	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		32.10	0.33	XYL 91		
06731532	**Low discrimination**					**06731532**
					GRA	
Clostridium clostridioforme		83.83	0.35	GLU 88	50	
Parabacteroides distasonis		12.63	0.27	XYL 91	1	
Bifidobacterium longum subsp. longum		3.54	0.13	RHA 88	99	
06731612	**Good identification**					**06731612**
Clostridium clostridioforme		100.0	0.41	GLU 88		
06731632	**Good identification**					**06731632**
Clostridium clostridioforme		100.0	0.41	GLU 88		
06731702	**Good identification**					**06731702**
Clostridium clostridioforme		100.0	0.39	GLU 88		
06731712	**Good identification**					**06731712**
Clostridium clostridioforme		100.0	0.50	GLU 88		
06731722	**Low discrimination**					**06731722**
Clostridium clostridioforme		65.72	0.39	GLU 88		
Parabacteroides distasonis		34.28	0.40	XYL 91		
06731732	**Acceptable identification**					**06731732**
Clostridium clostridioforme		94.19	0.50	GLU 88		
Parabacteroides distasonis		5.81	0.35	XYL 91		
06732100	**Good identification**					**06732100**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06732101	**Good identification**					**06732101**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06732102	**Good identification**					**06732102**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06732103	**Good identification**					**06732103**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06732120	**Good identification**					**06732120**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06732121	**Good identification**					**06732121**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06732122	**Good identification**					**06732122**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06732123	**Good identification**					**06732123**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06733000	**Low discrimination**					**06733000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		87.49	0.53	GLU 99		
Propionibacterium granulosum		12.51	0.24			
06733001	**Good identification**					**06733001**
Actinomyces naeslundii		100.0	0.56	GLU 99		
06733002	**Acceptable identification**					**06733002**
					CAT	
Actinomyces naeslundii		94.97	0.53	GLU 99	25	
Propionibacterium granulosum		3.39	0.13		90	
Actinomyces israelii		1.63	0.14		1	
06733003	**Good identification**					**06733003**
Actinomyces naeslundii		100.0	0.56	GLU 99		
06733020	**Good identification**					**06733020**
Actinomyces naeslundii		96.55	0.53	GLU 99		
Propionibacterium granulosum		3.45	0.13			
06733021	**Good identification**					**06733021**
Actinomyces naeslundii		100.0	0.56	GLU 99		
06733022	**Good identification**					**06733022**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06733023	**Good identification**					**06733023**
Actinomyces naeslundii		100.0	0.56	GLU 99		
06733040	**Good identification**					**06733040**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06733041	**Good identification**					**06733041**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06733042	**Good identification**					**06733042**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06733043	**Good identification**					**06733043**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06733060	**Good identification**					**06733060**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06733061	**Good identification**					**06733061**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06733062	**Good identification**					**06733062**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06733063	**Good identification**					**06733063**
Actinomyces naeslundii		100.0	0.43	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06733100	**Very good identification**					**06733100**
Actinomyces naeslundii		98.41	0.61	GLU 99		
Actinomyces israelii		1.59	0.21			
06733101	**Very good identification**					**06733101**
Actinomyces naeslundii		100.0	0.64	GLU 99		
06733102	**Very good identification**					**06733102**
Actinomyces naeslundii		95.89	0.61	GLU 99		
Actinomyces israelii		2.32	0.24			
Bifidobacterium adolescentis		1.80	0.16			
06733103	**Very good identification**					**06733103**
Actinomyces naeslundii		100.0	0.64	GLU 99		
06733110	**Good identification**					**06733110**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06733111	**Good identification**					**06733111**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06733112	**Low discrimination**					**06733112**
Actinomyces naeslundii		69.69	0.39	GLU 99		
Bifidobacterium adolescentis		30.32	0.17	MLZ 5		
06733113	**Good identification**					**06733113**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06733120	**Good identification**					**06733120**
					SPO	
Actinomyces naeslundii		92.56	0.61	GLU 99	1	
Clostridium ramosum		5.66	0.15		92	
Clostridium perfringens		1.78	0.09		84	
06733121	**Very good identification**					**06733121**
Actinomyces naeslundii		100.0	0.64	GLU 99		
06733122	**Low discrimination**					**06733122**
					SPO	
Actinomyces naeslundii		79.76	0.61	GLU 99	1	
Clostridium ramosum		19.52	0.25		92	
Clostridium perfringens		0.72	0.03		84	
06733123	**Very good identification**					**06733123**
Actinomyces naeslundii		100.0	0.64	GLU 99		
06733130	**Good identification**					**06733130**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06733131	**Good identification**					**06733131**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06733132	**Good identification**					**06733132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	GLU 99		
06733133	**Good identification**					**06733133**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06733140	**Good identification**					**06733140**
Actinomyces naeslundii		96.57	0.48	GLU 99		
Actinomyces israelii		3.43	0.14			
06733141	**Good identification**					**06733141**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06733142	**Acceptable identification**					**06733142**
Actinomyces naeslundii		94.94	0.48	GLU 99		
Actinomyces israelii		5.06	0.17			
06733143	**Good identification**					**06733143**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06733160	**Good identification**					**06733160**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06733161	**Good identification**					**06733161**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06733162	**Good identification**					**06733162**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06733163	**Good identification**					**06733163**
Actinomyces naeslundii		100.0	0.51	GLU 99		
06734312	**Good identification**					**06734312**
Clostridium clostridioforme		100.0	0.39	GLU 88		
06734332	**Good identification**					**06734332**
Clostridium clostridioforme		95.92	0.39	GLU 88		
Parabacteroides distasonis		4.08	0.21	ARA 94		
06734412	**Good identification**					**06734412**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06734432	**Good identification**					**06734432**
Clostridium clostridioforme		100.0	0.35	GLU 88		
06734512	**Good identification**					**06734512**
Clostridium clostridioforme		99.06	0.44	GLU 88		
Bifidobacterium adolescentis		0.94	0.13	RHA 88		
06734532	**Good identification**					**06734532**
Clostridium clostridioforme		100.0	0.44	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06734602	**Good identification**					**06734602**
Clostridium clostridioforme		100.0	0.39	GLU 88		
06734612	**Good identification**					**06734612**
Clostridium clostridioforme		100.0	0.50	GLU 88		
06734622	**Good identification**					**06734622**
Clostridium clostridioforme		100.0	0.39	GLU 88		
06734632	**Good identification**					**06734632**
Clostridium clostridioforme		100.0	0.50	GLU 88		
06734702	**Good identification**					**06734702**
Clostridium clostridioforme		100.0	0.48	GLU 88		
06734712	**Very good identification**					**06734712**
Clostridium clostridioforme		100.0	0.59	GLU 88		
06734722	**Good identification**					**06734722**
Clostridium clostridioforme		98.61	0.48	GLU 88		
Parabacteroides distasonis		1.39	0.22			
06734732	**Very good identification**					**06734732**
Clostridium clostridioforme		99.83	0.59	GLU 88		
Parabacteroides distasonis		0.17	0.17			
06735122	**Acceptable identification**					**06735122**
Parabacteroides distasonis		94.31	0.51	GLU 99		
Clostridium clostridioforme		3.25	0.21			
Prevotella oralis		1.43	0.08			
06735132	**Low discrimination**					**06735132**
Parabacteroides distasonis		77.42	0.47	GLU 99		
Clostridium clostridioforme		22.58	0.32			
06735212	**Good identification**					**06735212**
Clostridium clostridioforme		100.0	0.38	GLU 88		
06735232	**Good identification**					**06735232**
Clostridium clostridioforme		95.47	0.38	GLU 88		
Parabacteroides distasonis		4.53	0.21	ARA 94		
06735302	**Low discrimination**					**06735302**
Clostridium clostridioforme		89.30	0.36	GLU 88		
Parabacteroides distasonis		10.70	0.25	ARA 94		
06735312	**Good identification**					**06735312**
Clostridium clostridioforme		98.60	0.47	GLU 88		
Parabacteroides distasonis		1.40	0.21	ARA 94		
06735322	**Good identification**					**06735322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		92.04	0.60	GLU 99		
Clostridium clostridioforme		7.76	0.36			
Prevotella oralis		0.20	0.01			
06735332	**Low discrimination**					**06735332**
Parabacteroides distasonis		58.38	0.55	GLU 99		
Clostridium clostridioforme		41.62	0.47			
06735412	**Good identification**					**06735412**
Clostridium clostridioforme		98.95	0.43	GLU 88		
Bifidobacterium adolescentis		1.05	0.13	RHA 88		
06735432	**Good identification**					**06735432**
Clostridium clostridioforme		100.0	0.43	GLU 88		
06735502	**Low discrimination**					**06735502**
					SPO	
Bifidobacterium adolescentis		50.65	0.45	GLU 99	1	
Clostridium clostridioforme		43.43	0.41		66	
Clostridium butyricum		3.08	0.10		99	
06735512	**Low discrimination**					**06735512**
Clostridium clostridioforme		75.90	0.52	GLU 88		
Bifidobacterium adolescentis		23.75	0.47	RHA 88		
Bifidobacterium longum subsp. longum		0.24	0.11			
06735522	**Low discrimination**					**06735522**
					GRA	
Parabacteroides distasonis		52.05	0.48	GLU 99	1	
Clostridium clostridioforme		45.87	0.41		50	
Bifidobacterium longum subsp. longum		1.54	0.17		99	
06735532	**Low discrimination**					**06735532**
					GRA	
Clostridium clostridioforme		86.97	0.52	GLU 88	50	
Parabacteroides distasonis		11.67	0.43	RHA 88	1	
Bifidobacterium longum subsp. longum		1.09	0.21		99	
06735602	**Good identification**					**06735602**
Clostridium clostridioforme		100.0	0.47	GLU 88		
06735612	**Very good identification**					**06735612**
Clostridium clostridioforme		100.0	0.58	GLU 88		
06735622	**Good identification**					**06735622**
Clostridium clostridioforme		98.45	0.47	GLU 88		
Parabacteroides distasonis		1.55	0.22			
06735632	**Very good identification**					**06735632**
Clostridium clostridioforme		99.82	0.58	GLU 88		
Parabacteroides distasonis		0.19	0.17			
06735702	**Good identification**					**06735702**
					GRA	
Clostridium clostridioforme		99.37	0.56	GLU 88	50	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		0.47	0.22		1	
Bifidobacterium adolescentis		0.16	0.11		99	
06735712	**Very good identification**					**06735712**
					GRA	
Clostridium clostridioforme		99.90	0.67	GLU 88	50	
Parabacteroides distasonis		0.06	0.17		1	
Bifidobacterium adolescentis		0.04	0.13		99	
06735722	**Low discrimination**					**06735722**
Clostridium clostridioforme		68.09	0.56	GLU 88		
Parabacteroides distasonis		31.61	0.56			
Bacteroides thetaiotaomicron		0.23	0.03			
06735732	**Good identification**					**06735732**
Clostridium clostridioforme		94.77	0.67	GLU 88		
Parabacteroides distasonis		5.20	0.51			
Bacteroides thetaiotaomicron		0.03	-.04			
06737100	**Acceptable identification**					**06737100**
Actinomyces israelii		94.83	0.39	GLU 99		
Actinomyces naeslundii		5.17	0.27	CEL 93		
06737102	**Low discrimination**					**06737102**
Actinomyces israelii		67.24	0.42	GLU 99		
Bifidobacterium adolescentis		30.32	0.30	CEL 93		
Actinomyces naeslundii		2.44	0.27			
06737142	**Low discrimination**					**06737142**
Actinomyces israelii		87.47	0.35	GLU 99		
Bifidobacterium adolescentis		12.53	0.14	CEL 93		
06737500	**Low discrimination**					**06737500**
					SPO	
Actinomyces israelii		85.23	0.38	GLU 99	1	
Clostridium butyricum		10.00	0.01	CEL 93	99	
Bifidobacterium adolescentis		4.76	0.10		1	
06737502	**Low discrimination**					**06737502**
Bifidobacterium adolescentis		77.52	0.44	GLU 99		
Actinomyces israelii		21.02	0.41			
Clostridium clostridioforme		0.82	0.07			
06737512	**Acceptable identification**					**06737512**
Bifidobacterium adolescentis		92.36	0.45	GLU 99		
Clostridium clostridioforme		3.65	0.18			
Actinomyces israelii		3.06	0.26			
06743100	**Good identification**					**06743100**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06743101	**Good identification**					**06743101**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06743102	**Good identification**					**06743102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	GLU 99		
06743103	**Good identification**					**06743103**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06743120	**Good identification**					**06743120**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06743121	**Good identification**					**06743121**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06743122	**Low discrimination**					**06743122**
					SPO	
Actinomyces naeslundii		69.70	0.36	GLU 99	1	
Clostridium ramosum		30.30	0.05	FRU 95	92	
06743123	**Good identification**					**06743123**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06750100	**Acceptable identification**					**06750100**
					GRA	
Fusobacterium mortiferum		92.19	0.36	GLU 99	1	
Actinomyces naeslundii		7.81	0.29	MLT 11	99	
06750712	**Good identification**					**06750712**
Clostridium clostridioforme		100.0	0.36	GLU 88		
06750732	**Good identification**					**06750732**
Clostridium clostridioforme		100.0	0.36	GLU 88		
06751001	**Good identification**					**06751001**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06751003	**Good identification**					**06751003**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06751020	**Low discrimination**					**06751020**
					GRA	
Actinomyces naeslundii		43.78	0.45	GLU 99	99	
Lactobacillus catenaformis		21.41	0.18		99	
Prevotella oralis		13.23	0.11		1	
06751021	**Good identification**					**06751021**
Actinomyces naeslundii		100.0	0.48	GLU 99		
06751023	**Good identification**					**06751023**
					GRA	
Actinomyces naeslundii		96.28	0.48	GLU 99	99	
Prevotella oralis		3.72	-.02		1	
06751041	**Good identification**					**06751041**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06751043	**Good identification**					**06751043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	GLU 99		
06751061	**Good identification**					**06751061**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06751063	**Good identification**					**06751063**
Actinomyces naeslundii		100.0	0.35	GLU 99		
06751101	**Good identification**					**06751101**
Actinomyces naeslundii		100.0	0.56	GLU 99		
06751102	**Low discrimination**					**06751102**
					GRA	
Lactobacillus acidophilus		66.01	0.45	GLU 99	99	
Prevotella oralis		12.74	0.31		1	
Lactobacillus catenaformis		7.90	0.32		99	
06751103	**Acceptable identification**					**06751103**
Actinomyces naeslundii		92.30	0.56	GLU 99		
Lactobacillus acidophilus		5.36	0.10			
Actinomyces viscosus		1.31	0.00			
06751121	**Good identification**					**06751121**
					GRA	
Actinomyces naeslundii		96.61	0.56	GLU 99	99	
Prevotella melaninogenica		2.31	-.01		1	
Prevotella oralis		1.08	-.03		1	
06751122	**Low discrimination**					**06751122**
					GRA	
Prevotella oralis		71.95	0.53	GLU 99	1	
Lactobacillus acidophilus		19.62	0.45		99	
Actinomyces naeslundii		2.28	0.53		99	
06751123	**Low discrimination**					**06751123**
					GRA	
Actinomyces naeslundii		78.69	0.56	GLU 99	99	
Prevotella oralis		16.75	0.19		1	
Lactobacillus acidophilus		4.57	0.10		99	
06751140	**Low discrimination**					**06751140**
Actinomyces naeslundii		88.70	0.40	GLU 99		
Bifidobacterium breve		11.30	0.10			
06751141	**Good identification**					**06751141**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06751143	**Good identification**					**06751143**
Actinomyces naeslundii		100.0	0.43	GLU 99		
06751160	**Low discrimination**					**06751160**
					GRA	
Actinomyces naeslundii		55.86	0.40	GLU 99	99	
Bifidobacterium breve		12.65	0.15		99	
Prevotella melaninogenica		11.33	-.01		1	
06751161	**Good identification**					**06751161**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.43	GLU 99			
06751163	**Good identification**						**06751163**
Actinomyces naeslundii		100.0	0.43	GLU 99			
06751322	**Low discrimination**						**06751322**
					SPO	GRA	
Prevotella oralis		87.48	0.47	GLU 99	1	1	
Clostridium ramosum		6.64	0.28		92	50	
Lactobacillus acidophilus		4.19	0.25		1	99	
06751502	**Low discrimination**						**06751502**
Bifidobacterium adolescentis		82.90	0.40	GLU 99			
Clostridium clostridioforme		6.74	0.18	XYL 87			
Lactobacillus acidophilus		5.20	0.10				
06751512	**Low discrimination**						**06751512**
Bifidobacterium adolescentis		76.75	0.41	GLU 99			
Clostridium clostridioforme		23.25	0.29	XYL 87			
06751612	**Good identification**						**06751612**
Clostridium clostridioforme		100.0	0.35	GLU 88			
06751632	**Good identification**						**06751632**
Clostridium clostridioforme		100.0	0.35	GLU 88			
06751712	**Good identification**						**06751712**
Clostridium clostridioforme		99.55	0.44	GLU 88			
Bifidobacterium adolescentis		0.45	0.07	TRE 88			
06751732	**Good identification**						**06751732**
Clostridium clostridioforme		98.43	0.44	GLU 88			
Parabacteroides distasonis		1.57	0.19	TRE 88			
06752101	**Good identification**						**06752101**
Actinomyces naeslundii		100.0	0.37	GLU 99			
06752103	**Good identification**						**06752103**
Actinomyces naeslundii		100.0	0.37	GLU 99			
06752121	**Good identification**						**06752121**
Actinomyces naeslundii		100.0	0.37	GLU 99			
06752123	**Good identification**						**06752123**
Actinomyces naeslundii		100.0	0.37	GLU 99			
06753000	**Acceptable identification**						**06753000**
Actinomyces naeslundii		94.40	0.50	GLU 99			
Actinomyces israelii		5.60	0.20				
06753001	**Good identification**						**06753001**
Actinomyces naeslundii		100.0	0.53	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06753002	**Low discrimination**					**06753002**
Actinomyces naeslundii		87.20	0.50	GLU 99		
Actinomyces israelii		7.76	0.23			
Lactobacillus acidophilus		5.04	0.05			
06753003	**Good identification**					**06753003**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06753020	**Good identification**					**06753020**
Actinomyces naeslundii		100.0	0.50	GLU 99		
06753021	**Good identification**					**06753021**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06753022	**Low discrimination**					**06753022**
					SPO	
Actinomyces naeslundii		86.74	0.50	GLU 99	1	
Clostridium ramosum		5.65	0.05		92	
Lactobacillus acidophilus		5.02	0.05		1	
06753023	**Good identification**					**06753023**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06753040	**Good identification**					**06753040**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06753041	**Good identification**					**06753041**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06753042	**Low discrimination**					**06753042**
Actinomyces naeslundii		83.61	0.37	GLU 99		
Actinomyces israelii		16.39	0.16			
06753043	**Good identification**					**06753043**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06753060	**Good identification**					**06753060**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06753061	**Good identification**					**06753061**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06753062	**Good identification**					**06753062**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06753063	**Good identification**					**06753063**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06753100	**Good identification**					**06753100**
Actinomyces naeslundii		91.33	0.58	GLU 99		
Actinomyces israelii		7.61	0.30			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		1.06	0.10			
06753101	**Very good identification**					**06753101**
Actinomyces naeslundii		100.0	0.61	GLU 99		
06753102	**Low discrimination**					**06753102**
Actinomyces naeslundii		62.70	0.58	GLU 99		
Lactobacillus acidophilus		14.65	0.23			
Actinomyces israelii		7.84	0.33			
06753103	**Very good identification**					**06753103**
Actinomyces naeslundii		100.0	0.61	GLU 99		
06753110	**Low discrimination**					**06753110**
Actinomyces naeslundii		68.65	0.36	GLU 99		
Actinomyces israelii		16.25	0.16	MLZ 5		
Bifidobacterium breve		15.11	0.10			
06753111	**Good identification**					**06753111**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06753113	**Good identification**					**06753113**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06753120	**Low discrimination**					**06753120**
					SPO	
Actinomyces naeslundii		61.96	0.58	GLU 99	1	
Clostridium ramosum		35.08	0.28		92	
Bifidobacterium breve		1.28	0.15		1	
06753121	**Very good identification**					**06753121**
Actinomyces naeslundii		100.0	0.61	GLU 99		
06753122	**Low discrimination**					**06753122**
					SPO	
Clostridium ramosum		58.14	0.39	GLU 99	92	
Actinomyces naeslundii		25.67	0.58	TRE 94	1	
Lactobacillus acidophilus		6.00	0.23		1	
06753123	**Very good identification**					**06753123**
					SPO	
Actinomyces naeslundii		98.50	0.61	GLU 99	1	
Clostridium ramosum		1.50	0.05		92	
06753130	**Low discrimination**					**06753130**
Actinomyces naeslundii		71.88	0.36	GLU 99		
Bifidobacterium breve		28.12	0.15	MLZ 5		
06753131	**Good identification**					**06753131**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06753133	**Good identification**					**06753133**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06753140	**Low discrimination**					**06753140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		80.04	0.45	GLU 99			
Actinomyces israelii		14.71	0.23				
Bifidobacterium breve		5.25	0.10				
06753141	**Good identification**						**06753141**
Actinomyces naeslundii		100.0	0.48	GLU 99			
06753142	**Low discrimination**						**06753142**
Actinomyces naeslundii		46.21	0.45	GLU 99			
Bifidobacterium breve		30.67	0.27				
Actinomyces israelii		12.73	0.26				
06753143	**Good identification**						**06753143**
Actinomyces naeslundii		100.0	0.48	GLU 99			
06753160	**Low discrimination**						**06753160**
Actinomyces naeslundii		89.55	0.45	GLU 99			
Bifidobacterium breve		10.45	0.15				
06753161	**Good identification**						**06753161**
Actinomyces naeslundii		100.0	0.48	GLU 99			
06753163	**Good identification**						**06753163**
Actinomyces naeslundii		100.0	0.48	GLU 99			
06753322	**Good identification**						**06753322**
					SPO	GRA	
Clostridium ramosum		96.83	0.41	GLU 99	92	50	
Prevotella oralis		2.27	0.13	TRE 94	1	1	
Lactobacillus acidophilus		0.57	0.03		1	99	
06753502	**Low discrimination**						**06753502**
Bifidobacterium adolescentis		83.89	0.38	GLU 99			
Actinomyces israelii		14.67	0.32	XYL 87			
Actinomyces naeslundii		1.45	0.24				
06753512	**Good identification**						**06753512**
Bifidobacterium adolescentis		97.91	0.40	GLU 99			
Actinomyces israelii		2.09	0.18	XYL 87			
06754512	**Acceptable identification**						**06754512**
Clostridium clostridioforme		93.79	0.38	GLU 88			
Bifidobacterium adolescentis		6.21	0.21	TRE 88			
06754532	**Good identification**						**06754532**
Clostridium clostridioforme		100.0	0.38	GLU 88			
06754612	**Good identification**						**06754612**
Clostridium clostridioforme		100.0	0.44	GLU 88			
06754632	**Good identification**						**06754632**
Clostridium clostridioforme		100.0	0.44	GLU 88			
06754702	**Good identification**						**06754702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.42	GLU 88			
06754712	**Good identification**						**06754712**
Clostridium clostridioforme		100.0	0.53	GLU 88			
06754722	**Good identification**						**06754722**
Clostridium clostridioforme		100.0	0.42	GLU 88			
06754732	**Good identification**						**06754732**
Clostridium clostridioforme		100.0	0.53	GLU 88			
06755100	**Low discrimination**						**06755100**
					GRA	SPO	
Actinomyces israelii		46.67	0.36	GLU 99	99	1	
Clostridium butyricum		45.44	0.14		99	99	
Fusobacterium mortiferum		7.89	0.20		1	1	
06755102	**Low discrimination**						**06755102**
					GRA		
Bifidobacterium adolescentis		58.56	0.40	GLU 99	99		
Actinomyces israelii		20.78	0.39	ARA 87	99		
Prevotella oralis		8.67	0.16		1		
06755112	**Low discrimination**						**06755112**
Bifidobacterium adolescentis		80.71	0.41	GLU 99			
Clostridium clostridioforme		15.78	0.26	ARA 87			
Actinomyces israelii		3.50	0.24				
06755122	**Low discrimination**						**06755122**
					GRA		
Prevotella oralis		83.21	0.37	GLU 99	1		
Parabacteroides distasonis		11.65	0.36	XYL 11	1		
Lactobacillus acidophilus		1.85	0.10		99		
06755312	**Good identification**						**06755312**
Clostridium clostridioforme		99.30	0.41	GLU 88			
Bifidobacterium adolescentis		0.70	0.07	TRE 88			
06755332	**Low discrimination**						**06755332**
Clostridium clostridioforme		72.99	0.41	GLU 88			
Parabacteroides distasonis		26.51	0.39	TRE 88			
Prevotella oralis		0.51	-.03	ARA 94			
06755412	**Acceptable identification**						**06755412**
Clostridium clostridioforme		93.12	0.38	GLU 88			
Bifidobacterium adolescentis		6.88	0.21	TRE 88			
06755432	**Good identification**						**06755432**
Clostridium clostridioforme		100.0	0.38	GLU 88			
06755502	**Low discrimination**						**06755502**
					SPO		
Bifidobacterium adolescentis		77.72	0.54	GLU 99	1		
Clostridium clostridioforme		9.55	0.35		66		
Clostridium butyricum		9.36	0.24		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06755512	**Low discrimination**						**06755512**
Bifidobacterium adolescentis		68.11	0.56	GLU 99			
Clostridium clostridioforme		31.18	0.47				
Actinomyces israelii		0.36	0.23				
06755522	**Low discrimination**						**06755522**
					GRA	SPO	
Clostridium clostridioforme		63.63	0.35	GLU 88	50	66	
Parabacteroides distasonis		18.70	0.32	TRE 88	1	1	
Clostridium butyricum		5.43	0.05	RHA 88	99	99	
06755532	**Acceptable identification**						**06755532**
					GRA		
Clostridium clostridioforme		94.63	0.47	GLU 88	50		
Parabacteroides distasonis		3.29	0.27	TRE 88	1		
Bifidobacterium adolescentis		2.09	0.21	RHA 88	99		
06755542	**Low discrimination**						**06755542**
					SPO		
Bifidobacterium adolescentis		81.88	0.39	GLU 99	1		
Actinomyces israelii		11.18	0.30	SOR 11	1		
Clostridium butyricum		6.94	0.05		99		
06755552	**Acceptable identification**						**06755552**
Bifidobacterium adolescentis		94.87	0.40	GLU 99			
Clostridium clostridioforme		3.55	0.12	SOR 11			
Actinomyces israelii		1.58	0.16				
06755602	**Good identification**						**06755602**
Clostridium clostridioforme		100.0	0.41	GLU 88			
06755612	**Good identification**						**06755612**
Clostridium clostridioforme		100.0	0.52	GLU 88			
06755622	**Good identification**						**06755622**
Clostridium clostridioforme		100.0	0.41	GLU 88			
06755632	**Good identification**						**06755632**
Clostridium clostridioforme		100.0	0.52	GLU 88			
06755702	**Good identification**						**06755702**
					SPO		
Clostridium clostridioforme		98.14	0.50	GLU 88	66		
Bifidobacterium adolescentis		1.10	0.20	TRE 88	1		
Clostridium butyricum		0.41	-.02		99		
06755712	**Very good identification**						**06755712**
Clostridium clostridioforme		99.70	0.61	GLU 88			
Bifidobacterium adolescentis		0.30	0.21	TRE 88			
06755722	**Low discrimination**						**06755722**
Clostridium clostridioforme		87.35	0.50	GLU 88			
Parabacteroides distasonis		10.50	0.40	TRE 88			
Bacteroides ovatus		0.87	0.06				
06755732	**Very good identification**						**06755732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		98.46	0.61	GLU 88		
Parabacteroides distasonis		1.40	0.35	TRE 88		
Bacteroides ovatus		0.14	0.03			
06757000	**Good identification**					**06757000**
Actinomyces israelii		100.0	0.38	GLU 99		
06757002	**Good identification**					**06757002**
Actinomyces israelii		100.0	0.41	GLU 99		
06757100	**Good identification**					**06757100**
					SPO	
Actinomyces israelii		97.12	0.48	GLU 99	1	
Clostridium butyricum		1.86	-.02		99	
Actinomyces naeslundii		1.02	0.24		1	
06757102	**Low discrimination**					**06757102**
Actinomyces israelii		70.73	0.51	GLU 99		
Bifidobacterium adolescentis		28.78	0.38			
Actinomyces naeslundii		0.50	0.24			
06757112	**Low discrimination**					**06757112**
					SPO	
Bifidobacterium adolescentis		76.19	0.40	GLU 99	1	
Actinomyces israelii		22.89	0.37	ARA 87	1	
Clostridium tertium		0.92	-.01		99	
06757140	**Good identification**					**06757140**
Actinomyces israelii		100.0	0.41	GLU 99		
06757142	**Low discrimination**					**06757142**
Actinomyces israelii		88.55	0.44	GLU 99		
Bifidobacterium adolescentis		11.45	0.23			
06757400	**Good identification**					**06757400**
Actinomyces israelii		100.0	0.37	GLU 99		
06757402	**Low discrimination**					**06757402**
Actinomyces israelii		88.15	0.40	GLU 99		
Bifidobacterium adolescentis		11.85	0.18			
06757500	**Low discrimination**					**06757500**
					SPO	
Actinomyces israelii		79.34	0.47	GLU 99	1	
Clostridium butyricum		16.66	0.14		99	
Bifidobacterium adolescentis		4.00	0.18		1	
06757502	**Low discrimination**					**06757502**
					SPO	
Bifidobacterium adolescentis		75.94	0.53	GLU 99	1	
Actinomyces israelii		22.82	0.50		1	
Clostridium butyricum		1.24	0.07		99	
06757512	**Good identification**					**06757512**
Bifidobacterium adolescentis		95.64	0.54	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		3.51	0.36			
Clostridium clostridioforme		0.54	0.12			
06757540	**Good identification**					**06757540**
					SPO	
Actinomyces israelii		95.52	0.40	GLU 99	1	
Clostridium butyricum		4.49	-.04		99	
06757542	**Low discrimination**					**06757542**
Bifidobacterium adolescentis		51.40	0.37	GLU 99		
Actinomyces israelii		48.60	0.43	SOR 11		
06757552	**Low discrimination**					**06757552**
Bifidobacterium adolescentis		89.64	0.39	GLU 99		
Actinomyces israelii		10.36	0.29	SOR 11		
06761100	**Good identification**					**06761100**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06761101	**Good identification**					**06761101**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06761103	**Good identification**					**06761103**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06761120	**Low discrimination**					**06761120**
					SPO	
Actinomyces naeslundii		76.90	0.36	GLU 99	1	
Clostridium ramosum		23.11	0.02	FRU 95	92	
06761121	**Good identification**					**06761121**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06761123	**Good identification**					**06761123**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06763001	**Good identification**					**06763001**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06763003	**Good identification**					**06763003**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06763021	**Good identification**					**06763021**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06763023	**Good identification**					**06763023**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06763100	**Good identification**					**06763100**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06763101	**Good identification**					**06763101**
Actinomyces naeslundii		100.0	0.44	GLU 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
06763102	**Good identification**					**06763102**
Actinomyces naeslundii		100.0	0.41	GLU 99		
06763103	**Good identification**					**06763103**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06763120	**Low discrimination**					**06763120**
					SPO	
Actinomyces naeslundii		53.28	0.41	GLU 99	1	
Clostridium ramosum		46.72	0.15	FRU 95	92	
06763121	**Good identification**					**06763121**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06763123	**Good identification**					**06763123**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06765712	**Good identification**					**06765712**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06765732	**Good identification**					**06765732**
Clostridium clostridioforme		99.13	0.42	GLU 88		
Parabacteroides distasonis		0.88	0.13	FRU 99		
06770101	**Good identification**					**06770101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06770103	**Good identification**					**06770103**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06770121	**Good identification**					**06770121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06770123	**Good identification**					**06770123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06770512	**Good identification**					**06770512**
Clostridium clostridioforme		100.0	0.36	GLU 88		
06770532	**Good identification**					**06770532**
Clostridium clostridioforme		100.0	0.36	GLU 88		
06770612	**Good identification**					**06770612**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06770632	**Good identification**					**06770632**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06770702	**Good identification**					**06770702**
Clostridium clostridioforme		100.0	0.40	GLU 88		
06770712	**Good identification**					**06770712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.51	GLU 88		
06770722	**Good identification**					**06770722**
Clostridium clostridioforme		100.0	0.40	GLU 88		
06770732	**Good identification**					**06770732**
Clostridium clostridioforme		100.0	0.51	GLU 88		
06771001	**Good identification**					**06771001**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06771002	**Low discrimination**					**06771002**
Lactobacillus acidophilus		63.98	0.35	GLU 99		
Lactobacillus jensenii		17.60	0.29	LAC 92		
Actinomyces naeslundii		17.39	0.50			
06771003	**Good identification**					**06771003**
Actinomyces naeslundii		97.58	0.53	GLU 99		
Lactobacillus acidophilus		2.42	0.01			
06771020	**Low discrimination**					**06771020**
Actinomyces naeslundii		66.37	0.50	GLU 99		
Lactobacillus jensenii		25.19	0.21			
Clostridium perfringens		2.99	0.05			
06771021	**Good identification**					**06771021**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06771022	**Low discrimination**					**06771022**
Lactobacillus acidophilus		69.40	0.35	GLU 99		
Actinomyces naeslundii		18.87	0.50	LAC 92		
Lactobacillus jensenii		4.77	0.18			
06771023	**Good identification**					**06771023**
Actinomyces naeslundii		97.58	0.53	GLU 99		
Lactobacillus acidophilus		2.42	0.01			
06771040	**Good identification**					**06771040**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06771041	**Good identification**					**06771041**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06771043	**Good identification**					**06771043**
Actinomyces naeslundii		100.0	0.40	GLU 99		
06771060	**Good identification**					**06771060**
Actinomyces naeslundii		100.0	0.37	GLU 99		
06771061	**Good identification**					**06771061**
Actinomyces naeslundii		100.0	0.40	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06771063	**Good identification**						**06771063**
Actinomyces naeslundii		100.0	0.40	GLU 99			
06771100	**Low discrimination**						**06771100**
					COC		
Actinomyces naeslundii		64.49	0.58	GLU 99	1		
Lactobacillus acidophilus		9.68	0.19		1		
Atopobium parvulum		8.78	0.20		99		
06771101	**Very good identification**						**06771101**
Actinomyces naeslundii		100.0	0.61	GLU 99			
06771102	**Acceptable identification**						**06771102**
Lactobacillus acidophilus		91.08	0.54	GLU 99			
Actinomyces naeslundii		6.13	0.58				
Actinomyces israelii		0.51	0.30				
06771103	**Good identification**						**06771103**
Actinomyces naeslundii		90.90	0.61	GLU 99			
Lactobacillus acidophilus		9.10	0.19				
06771110	**Low discrimination**						**06771110**
					GRA		
Actinomyces naeslundii		84.22	0.36	GLU 99	99		
Leptotrichia buccalis		15.78	-.11	MLZ 5	1		
06771111	**Good identification**						**06771111**
Actinomyces naeslundii		100.0	0.39	GLU 99			
06771113	**Low discrimination**						**06771113**
					GRA		
Actinomyces naeslundii		88.90	0.39	GLU 99	99		
Leptotrichia buccalis		11.10	-.11	MLZ 5	1		
06771120	**Low discrimination**						**06771120**
					SPO		
Clostridium ramosum		52.43	0.36	GLU 99	92		
Actinomyces naeslundii		33.49	0.58		1		
Lactobacillus acidophilus		5.03	0.19		1		
06771121	**Very good identification**						**06771121**
					SPO		
Actinomyces naeslundii		98.96	0.61	GLU 99	1		
Clostridium ramosum		1.04	0.02		92		
06771122	**Low discrimination**						**06771122**
					SPO	GRA	
Lactobacillus acidophilus		63.08	0.54	GLU 99	1	99	
Clostridium ramosum		26.58	0.46		92	50	
Prevotella oralis		4.41	0.33		1	1	
06771123	**Low discrimination**						**06771123**
					SPO		
Actinomyces naeslundii		87.01	0.61	GLU 99	1		
Lactobacillus acidophilus		8.71	0.19		1		
Clostridium ramosum		3.67	0.12		92		
06771130	**Low discrimination**						**06771130**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		64.05	0.36	GLU 99	1		
Clostridium ramosum		19.25	0.02	MLZ 5	92		
Bifidobacterium breve		16.71	0.12		1		
06771131	**Good identification**						**06771131**
Actinomyces naeslundii		100.0	0.39	GLU 99			
06771133	**Good identification**						**06771133**
Actinomyces naeslundii		100.0	0.39	GLU 99			
06771140	**Low discrimination**						**06771140**
Actinomyces naeslundii		81.49	0.45	GLU 99			
Actinomyces israelii		9.94	0.20				
Lactobacillus acidophilus		8.57	0.04				
06771141	**Good identification**						**06771141**
Actinomyces naeslundii		100.0	0.48	GLU 99			
06771142	**Low discrimination**						**06771142**
Lactobacillus acidophilus		86.14	0.38	GLU 99			
Actinomyces naeslundii		8.28	0.45	SOR 11			
Bifidobacterium breve		3.66	0.24				
06771143	**Acceptable identification**						**06771143**
Actinomyces naeslundii		93.45	0.48	GLU 99			
Lactobacillus acidophilus		6.55	0.04				
06771160	**Low discrimination**						**06771160**
Actinomyces naeslundii		71.54	0.45	GLU 99			
Clostridium perfringens		8.96	0.07				
Lactobacillus acidophilus		7.52	0.04				
06771161	**Good identification**						**06771161**
Actinomyces naeslundii		100.0	0.48	GLU 99			
06771162	**Low discrimination**						**06771162**
Lactobacillus acidophilus		82.14	0.38	GLU 99			
Actinomyces naeslundii		7.89	0.45	SOR 11			
Bifidobacterium breve		6.21	0.29				
06771163	**Acceptable identification**						**06771163**
Actinomyces naeslundii		93.45	0.48	GLU 99			
Lactobacillus acidophilus		6.55	0.04				
06771312	**Good identification**						**06771312**
					GRA		
Clostridium clostridioforme		97.73	0.39	GLU 88	50		
Lactobacillus acidophilus		1.21	0.00	XYL 91	99		
Leptotrichia buccalis		1.06	-.11	ARA 94	1		
06771320	**Good identification**						**06771320**
					SPO	GRA	
Clostridium ramosum		97.89	0.38	GLU 99	92	50	
Prevotella oralis		1.11	0.04		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		0.53	0.00		1	99	
06771322	**Low discrimination**						**06771322**
					SPO	GRA	
Clostridium ramosum		79.04	0.48	GLU 99	92	50	
Lactobacillus acidophilus		10.65	0.34		1	99	
Prevotella oralis		4.24	0.26		1	1	
06771332	**Low discrimination**						**06771332**
					SPO		
Clostridium clostridioforme		73.63	0.39	GLU 88	66		
Parabacteroides distasonis		18.66	0.34	XYL 91	1		
Clostridium ramosum		6.79	0.14	ARA 94	92		
06771412	**Good identification**						**06771412**
Clostridium clostridioforme		100.0	0.35	GLU 88			
06771432	**Good identification**						**06771432**
Clostridium clostridioforme		100.0	0.35	GLU 88			
06771502	**Low discrimination**						**06771502**
					SPO		
Bifidobacterium adolescentis		33.80	0.37	GLU 99	1		
Clostridium clostridioforme		30.22	0.33	XYL 87	66		
Clostridium butyricum		19.81	0.18		99		
06771512	**Low discrimination**						**06771512**
					SPO		
Clostridium clostridioforme		75.88	0.44	GLU 88	66		
Bifidobacterium adolescentis		22.77	0.38	XYL 91	1		
Clostridium butyricum		0.57	-.04	RHA 88	99		
06771532	**Good identification**						**06771532**
Clostridium clostridioforme		97.63	0.44	GLU 88			
Parabacteroides distasonis		2.37	0.22	XYL 91			
06771602	**Good identification**						**06771602**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06771612	**Good identification**						**06771612**
Clostridium clostridioforme		100.0	0.50	GLU 88			
06771622	**Good identification**						**06771622**
Clostridium clostridioforme		100.0	0.39	GLU 88			
06771632	**Good identification**						**06771632**
Clostridium clostridioforme		100.0	0.50	GLU 88			
06771702	**Good identification**						**06771702**
					SPO		
Clostridium clostridioforme		99.37	0.48	GLU 88	66		
Lactobacillus acidophilus		0.36	0.00	XYL 91	1		
Clostridium butyricum		0.28	-.08		99		
06771712	**Very good identification**						**06771712**
Clostridium clostridioforme		100.0	0.59	GLU 88			
06771722	**Low discrimination**						**06771722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium clostridioforme		89.74	0.48	GLU 88	66	
Parabacteroides distasonis		7.53	0.35	XYL 91	1	
Clostridium ramosum		2.41	0.14		92	
06771732	**Very good identification**					**06771732**
Clostridium clostridioforme		99.02	0.59	GLU 88		
Parabacteroides distasonis		0.98	0.30	XYL 91		
06772100	**Good identification**					**06772100**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06772101	**Good identification**					**06772101**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06772102	**Good identification**					**06772102**
Actinomyces naeslundii		100.0	0.39	GLU 99		
06772103	**Good identification**					**06772103**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06772121	**Good identification**					**06772121**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06772123	**Good identification**					**06772123**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06773000	**Low discrimination**					**06773000**
Actinomyces naeslundii		89.69	0.55	GLU 99		
Actinomyces israelii		10.31	0.30			
06773001	**Very good identification**					**06773001**
Actinomyces naeslundii		100.0	0.58	GLU 99		
06773002	**Low discrimination**					**06773002**
Actinomyces naeslundii		78.61	0.55	GLU 99		
Actinomyces israelii		13.56	0.33			
Lactobacillus acidophilus		7.84	0.14			
06773003	**Very good identification**					**06773003**
Actinomyces naeslundii		100.0	0.58	GLU 99		
06773011	**Good identification**					**06773011**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06773013	**Good identification**					**06773013**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06773020	**Low discrimination**					**06773020**
					SPO	
Actinomyces naeslundii		88.39	0.55	GLU 99	1	
Clostridium ramosum		11.61	0.15		92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06773021	**Very good identification**					**06773021**
Actinomyces naeslundii		100.0	0.58	GLU 99		
06773022	**Low discrimination**					**06773022**
					SPO	
Actinomyces naeslundii		61.53	0.55	GLU 99	1	
Clostridium ramosum		32.33	0.25		92	
Lactobacillus acidophilus		6.14	0.14		1	
06773023	**Very good identification**					**06773023**
Actinomyces naeslundii		100.0	0.58	GLU 99		
06773031	**Good identification**					**06773031**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06773033	**Good identification**					**06773033**
Actinomyces naeslundii		100.0	0.36	GLU 99		
06773040	**Low discrimination**					**06773040**
Actinomyces naeslundii		79.79	0.42	GLU 99		
Actinomyces israelii		20.21	0.23			
06773041	**Good identification**					**06773041**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06773042	**Low discrimination**					**06773042**
Actinomyces naeslundii		68.97	0.42	GLU 99		
Actinomyces israelii		26.21	0.26			
Lactobacillus acidophilus		4.82	-.02			
06773043	**Good identification**					**06773043**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06773060	**Good identification**					**06773060**
Actinomyces naeslundii		100.0	0.42	GLU 99		
06773061	**Good identification**					**06773061**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06773062	**Acceptable identification**					**06773062**
Actinomyces naeslundii		93.47	0.42	GLU 99		
Lactobacillus acidophilus		6.53	-.02			
06773063	**Good identification**					**06773063**
Actinomyces naeslundii		100.0	0.45	GLU 99		
06773100	**Low discrimination**					**06773100**
					SPO	
Actinomyces naeslundii		82.12	0.63	GLU 99	1	
Actinomyces israelii		13.27	0.40		1	
Clostridium ramosum		3.79	0.15		92	
06773101	**Very good identification**					**06773101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.66	GLU 99		
06773102	**Low discrimination**					**06773102**
Actinomyces naeslundii		52.18	0.63	GLU 99		
Lactobacillus acidophilus		21.03	0.32			
Actinomyces israelii		12.65	0.43			
06773103	**Very good identification**					**06773103**
Actinomyces naeslundii		100.0	0.66	GLU 99		
06773110	**Low discrimination**					**06773110**
					SPO	
Actinomyces naeslundii		60.88	0.41	GLU 99	1	
Actinomyces israelii		27.93	0.26	MLZ 5	1	
Clostridium tertium		11.19	0.05		99	
06773111	**Good identification**					**06773111**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06773112	**Low discrimination**					**06773112**
Actinomyces naeslundii		26.33	0.41	GLU 99		
Bifidobacterium breve		20.13	0.24	MLZ 5		
Bifidobacterium adolescentis		18.36	0.23			
06773113	**Good identification**					**06773113**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06773120	**Low discrimination**					**06773120**
					SPO	
Clostridium ramosum		81.55	0.49	GLU 99	92	
Actinomyces naeslundii		17.85	0.63		1	
Clostridium tertium		0.29	0.09		99	
06773121	**Very good identification**					**06773121**
					SPO	
Actinomyces naeslundii		97.02	0.66	GLU 99	1	
Clostridium ramosum		2.99	0.15		92	
06773122	**Good identification**					**06773122**
					SPO	
Clostridium ramosum		92.21	0.59	GLU 99	92	
Actinomyces naeslundii		5.05	0.63		1	
Lactobacillus acidophilus		2.03	0.32		1	
06773123	**Low discrimination**					**06773123**
					SPO	
Actinomyces naeslundii		89.04	0.66	GLU 99	1	
Clostridium ramosum		10.96	0.25		92	
06773130	**Low discrimination**					**06773130**
					SPO	
Actinomyces naeslundii		39.02	0.41	GLU 99	1	
Clostridium ramosum		34.22	0.15	MLZ 5	92	
Clostridium tertium		21.52	0.13		99	
06773131	**Good identification**					**06773131**
Actinomyces naeslundii		100.0	0.44	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06773133	**Good identification**					**06773133**
Actinomyces naeslundii		100.0	0.44	GLU 99		
06773140	**Low discrimination**					**06773140**
Actinomyces naeslundii		73.74	0.50	GLU 99		
Actinomyces israelii		26.26	0.33			
06773141	**Good identification**					**06773141**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06773142	**Low discrimination**					**06773142**
Actinomyces naeslundii		48.92	0.50	GLU 99		
Actinomyces israelii		26.13	0.36			
Lactobacillus acidophilus		13.81	0.16			
06773143	**Good identification**					**06773143**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06773160	**Low discrimination**					**06773160**
					SPO	
Actinomyces naeslundii		76.83	0.50	GLU 99	1	
Clostridium ramosum		20.10	0.15		92	
Bifidobacterium breve		3.08	0.12		1	
06773161	**Good identification**					**06773161**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06773163	**Good identification**					**06773163**
Actinomyces naeslundii		100.0	0.53	GLU 99		
06773320	**Good identification**					**06773320**
					SPO	
Clostridium ramosum		99.83	0.51	GLU 99	92	
Actinomyces naeslundii		0.17	0.29		1	
06773322	**Very good identification**					**06773322**
					SPO	
Clostridium ramosum		99.83	0.61	GLU 99	92	
Lactobacillus acidophilus		0.13	0.12		1	
Actinomyces naeslundii		0.04	0.29		1	
06773500	**Low discrimination**					**06773500**
					SPO	
Actinomyces israelii		72.07	0.39	GLU 99	1	
Clostridium butyricum		22.42	0.09	XYL 92	99	
Actinomyces naeslundii		5.51	0.29		1	
06773502	**Low discrimination**					**06773502**
					SPO	
Bifidobacterium adolescentis		47.25	0.35	GLU 99	1	
Actinomyces israelii		46.62	0.42	XYL 87	1	
Clostridium butyricum		3.76	0.02		99	
06773512	**Low discrimination**					**06773512**
Bifidobacterium adolescentis		83.78	0.37	GLU 99		
Actinomyces israelii		10.11	0.28	XYL 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		3.45	0.10			
06774212	**Good identification**					**06774212**
Clostridium clostridioforme		100.0	0.39	GLU 88		
06774232	**Good identification**					**06774232**
Clostridium clostridioforme		100.0	0.39	GLU 88		
06774302	**Good identification**					**06774302**
Clostridium clostridioforme		100.0	0.36	GLU 88		
06774312	**Good identification**					**06774312**
Clostridium clostridioforme		100.0	0.48	GLU 88		
06774322	**Acceptable identification**					**06774322**
Clostridium clostridioforme		94.53	0.36	GLU 88		
Parabacteroides distasonis		5.47	0.21	ARA 94		
06774332	**Good identification**					**06774332**
Clostridium clostridioforme		99.32	0.48	GLU 88		
Parabacteroides distasonis		0.68	0.16	ARA 94		
06774412	**Good identification**					**06774412**
Clostridium clostridioforme		100.0	0.44	GLU 88		
06774432	**Good identification**					**06774432**
Clostridium clostridioforme		100.0	0.44	GLU 88		
06774502	**Acceptable identification**					**06774502**
					SPO	
Clostridium clostridioforme		92.09	0.42	GLU 88	66	
Clostridium butyricum		5.85	0.10	RHA 88	99	
Bifidobacterium adolescentis		2.07	0.17		1	
06774512	**Good identification**					**06774512**
Clostridium clostridioforme		99.40	0.53	GLU 88		
Bifidobacterium adolescentis		0.60	0.18	RHA 88		
06774522	**Good identification**					**06774522**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06774532	**Good identification**					**06774532**
Clostridium clostridioforme		100.0	0.53	GLU 88		
06774602	**Good identification**					**06774602**
Clostridium clostridioforme		100.0	0.48	GLU 88		
06774612	**Very good identification**					**06774612**
Clostridium clostridioforme		100.0	0.59	GLU 88		
06774622	**Good identification**					**06774622**
Clostridium clostridioforme		100.0	0.48	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06774632	**Very good identification**					**06774632**
Clostridium clostridioforme		100.0	0.59	GLU 88		
06774702	**Very good identification**					**06774702**
Clostridium clostridioforme		100.0	0.57	GLU 88		
06774712	**Very good identification**					**06774712**
Clostridium clostridioforme		100.0	0.68	GLU 88		
06774722	**Very good identification**					**06774722**
Clostridium clostridioforme		99.77	0.57	GLU 88		
Parabacteroides distasonis		0.23	0.17			
06774732	**Very good identification**					**06774732**
Clostridium clostridioforme		99.97	0.68	GLU 88		
Parabacteroides distasonis		0.03	0.13			
06775000	**Low discrimination**					**06775000**
					SPO	
Actinomyces israelii		59.67	0.35	GLU 99	1	
Clostridium butyricum		40.33	0.11		99	
06775002	**Low discrimination**					**06775002**
					SPO	
Actinomyces israelii		63.52	0.38	GLU 99	1	
Clostridium clostridioforme		21.88	0.21		66	
Clostridium butyricum		11.13	0.04		99	
06775112	**Low discrimination**					**06775112**
Clostridium clostridioforme		61.40	0.41	GLU 88		
Bifidobacterium adolescentis		28.54	0.38	RHA 88		
Actinomyces israelii		6.99	0.34	ARA 94		
06775122	**Low discrimination**					**06775122**
					GRA	
Parabacteroides distasonis		62.35	0.47	GLU 99	1	
Clostridium clostridioforme		13.36	0.30		50	
Lactobacillus acidophilus		7.29	0.19		99	
06775132	**Low discrimination**					**06775132**
					GRA	
Clostridium clostridioforme		64.16	0.41	GLU 88	50	
Parabacteroides distasonis		35.38	0.42	RHA 88	1	
Actinomyces israelii		0.47	0.14	ARA 94	99	
06775140	**Low discrimination**					**06775140**
					SPO	
Actinomyces israelii		52.44	0.38	GLU 99	1	
Clostridium butyricum		47.56	0.17		99	
06775142	**Low discrimination**					**06775142**
					SPO	
Actinomyces israelii		69.26	0.41	GLU 99	1	
Clostridium butyricum		16.28	0.10		99	
Bifidobacterium adolescentis		10.99	0.21		1	
06775202	**Good identification**					**06775202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		97.11	0.36	GLU 88		
Actinomyces israelii		2.89	0.19	ARA 94		
06775212	**Good identification**					**06775212**
Clostridium clostridioforme		100.0	0.47	GLU 88		
06775222	**Acceptable identification**					**06775222**
Clostridium clostridioforme		93.94	0.36	GLU 88		
Parabacteroides distasonis		6.06	0.21	ARA 94		
06775232	**Good identification**					**06775232**
Clostridium clostridioforme		99.24	0.47	GLU 88		
Parabacteroides distasonis		0.76	0.16	ARA 94		
06775302	**Acceptable identification**					**06775302**
					GRA	
Clostridium clostridioforme		92.95	0.45	GLU 88	50	
Actinomyces israelii		3.31	0.29	ARA 94	99	
Parabacteroides distasonis		1.79	0.21		1	
06775312	**Good identification**					**06775312**
					GRA	
Clostridium clostridioforme		99.66	0.56	GLU 88	50	
Parabacteroides distasonis		0.23	0.16	ARA 94	1	
Actinomyces israelii		0.12	0.15		99	
06775322	**Low discrimination**					**06775322**
					SPO	
Parabacteroides distasonis		63.84	0.55	GLU 99	1	
Clostridium clostridioforme		33.45	0.45		66	
Clostridium ramosum		1.39	0.14		92	
06775332	**Low discrimination**					**06775332**
Clostridium clostridioforme		81.53	0.56	GLU 88		
Parabacteroides distasonis		18.39	0.50	ARA 94		
Bacteroides ovatus		0.08	-.06			
06775402	**Low discrimination**					**06775402**
					SPO	
Clostridium clostridioforme		68.07	0.41	GLU 88	66	
Clostridium butyricum		19.90	0.20	RHA 88	99	
Actinomyces israelii		10.32	0.37		1	
06775412	**Good identification**					**06775412**
Clostridium clostridioforme		98.52	0.52	GLU 88		
Bifidobacterium adolescentis		0.66	0.18	RHA 88		
Actinomyces israelii		0.49	0.22			
06775422	**Good identification**					**06775422**
					SPO	
Clostridium clostridioforme		96.61	0.41	GLU 88	66	
Clostridium butyricum		2.46	0.02	RHA 88	99	
Actinomyces israelii		0.94	0.16		1	
06775432	**Good identification**					**06775432**
Clostridium clostridioforme		100.0	0.52	GLU 88		
06775500	**Good identification**					**06775500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium butyricum		97.50	0.51	ESL 88	99		
Actinomyces israelii		2.19	0.44	GLU 99	1		
Clostridium clostridioforme		0.18	0.16		66		
06775502	**Low discrimination**						**06775502**
					SPO		
Clostridium butyricum		52.17	0.44	ESL 88	99		
Clostridium clostridioforme		24.90	0.50	GLU 99	66		
Bifidobacterium adolescentis		18.43	0.51		1		
06775512	**Low discrimination**						**06775512**
					SPO		
Clostridium clostridioforme		81.33	0.61	GLU 88	66		
Bifidobacterium adolescentis		16.15	0.53	RHA 88	1		
Clostridium butyricum		1.97	0.22		99		
06775522	**Low discrimination**						**06775522**
					GRA	SPO	
Clostridium clostridioforme		71.99	0.50	GLU 88	50	66	
Parabacteroides distasonis		13.14	0.43	RHA 88	1	1	
Clostridium butyricum		13.12	0.26		99	99	
06775532	**Very good identification**						**06775532**
					GRA	SPO	
Clostridium clostridioforme		97.29	0.61	GLU 88	50	66	
Parabacteroides distasonis		2.10	0.39	RHA 88	1	1	
Clostridium butyricum		0.21	0.04		99	99	
06775552	**Low discrimination**						**06775552**
Bifidobacterium adolescentis		62.83	0.37	GLU 99			
Clostridium clostridioforme		25.86	0.27	SOR 11			
Actinomyces israelii		5.92	0.26				
06775602	**Good identification**						**06775602**
					SPO		
Clostridium clostridioforme		99.72	0.56	GLU 88	66		
Actinomyces israelii		0.16	0.17		1		
Clostridium butyricum		0.12	-.06		99		
06775612	**Very good identification**						**06775612**
Clostridium clostridioforme		100.0	0.67	GLU 88			
06775622	**Good identification**						**06775622**
Clostridium clostridioforme		99.75	0.56	GLU 88			
Parabacteroides distasonis		0.25	0.17				
06775632	**Very good identification**						**06775632**
Clostridium clostridioforme		99.97	0.67	GLU 88			
Parabacteroides distasonis		0.03	0.13				
06775702	**Very good identification**						**06775702**
					SPO		
Clostridium clostridioforme		98.77	0.65	GLU 88	66		
Clostridium butyricum		0.87	0.18		99		
Actinomyces israelii		0.18	0.28		1		
06775710	**Good identification**						**06775710**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		97.47	0.42	GLU 88		
Clostridium butyricum		2.53	0.03	BGL 99		
06775712	**Very good identification**					**06775712**
Clostridium clostridioforme		99.95	0.76	GLU 88		
Bifidobacterium adolescentis		0.03	0.18			
Actinomyces israelii		0.01	0.14			
06775713	**Good identification**					**06775713**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06775722	**Good identification**					**06775722**
Clostridium clostridioforme		91.63	0.65	GLU 88		
Parabacteroides distasonis		6.84	0.51			
Bacteroides ovatus		1.26	0.23			
06775730	**Good identification**					**06775730**
Clostridium clostridioforme		99.13	0.42	GLU 88		
Parabacteroides distasonis		0.88	0.13	BGL 99		
06775732	**Very good identification**					**06775732**
Clostridium clostridioforme		98.91	0.76	GLU 88		
Parabacteroides distasonis		0.87	0.47			
Bacteroides ovatus		0.20	0.20			
06775733	**Good identification**					**06775733**
Clostridium clostridioforme		99.13	0.42	GLU 88		
Parabacteroides distasonis		0.88	0.13	URE 1		
06775752	**Good identification**					**06775752**
Clostridium clostridioforme		100.0	0.42	GLU 88		
06775772	**Good identification**					**06775772**
Clostridium clostridioforme		98.31	0.42	GLU 88		
Parabacteroides distasonis		0.87	0.13	SOR 1		
Bacteroides ovatus		0.82	-.04			
06777000	**Good identification**					**06777000**
					SPO	
Actinomyces israelii		98.69	0.48	GLU 99	1	
Clostridium butyricum		1.31	-.05		99	
06777002	**Good identification**					**06777002**
Actinomyces israelii		100.0	0.51	GLU 99		
06777012	**Good identification**					**06777012**
Actinomyces israelii		100.0	0.37	GLU 99		
06777040	**Good identification**					**06777040**
Actinomyces israelii		100.0	0.41	GLU 99		
06777042	**Good identification**					**06777042**
Actinomyces israelii		100.0	0.44	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
06777100	**Good identification**					**06777100**
					SPO	
Actinomyces israelii		91.83	0.58	GLU 99	1	
Clostridium butyricum		7.30	0.18		99	
Actinomyces naeslundii		0.50	0.29		1	
06777102	**Good identification**					**06777102**
					SPO	
Actinomyces israelii		90.54	0.61	GLU 99	1	
Bifidobacterium adolescentis		6.53	0.35		1	
Clostridium butyricum		1.87	0.11		99	
06777110	**Acceptable identification**					**06777110**
					SPO	
Actinomyces israelii		92.93	0.44	GLU 99	1	
Clostridium tertium		4.47	0.07	MLZ 13	99	
Clostridium butyricum		2.60	-.04		99	
06777112	**Low discrimination**					**06777112**
					SPO	
Actinomyces israelii		58.13	0.47	GLU 99	1	
Bifidobacterium adolescentis		34.28	0.37	MLZ 13	1	
Clostridium tertium		5.59	0.15		99	
06777120	**Low discrimination**					**06777120**
					SPO	
Actinomyces israelii		56.41	0.38	GLU 99	1	
Clostridium ramosum		21.95	0.15	NAG 6	92	
Clostridium tertium		10.72	0.11		99	
06777140	**Good identification**					**06777140**
					SPO	
Actinomyces israelii		98.25	0.51	GLU 99	1	
Clostridium butyricum		1.75	0.00		99	
06777142	**Good identification**					**06777142**
					SPO	
Actinomyces israelii		97.32	0.54	GLU 99	1	
Bifidobacterium adolescentis		2.23	0.20		1	
Clostridium butyricum		0.45	-.07		99	
06777150	**Acceptable identification**					**06777150**
					COC	
Actinomyces israelii		93.35	0.37	GLU 99	1	
Blautia producta		6.65	-.03	MLZ 13	99	
06777152	**Low discrimination**					**06777152**
Actinomyces israelii		84.21	0.40	GLU 99		
Bifidobacterium adolescentis		15.79	0.21	MLZ 13		
06777300	**Good identification**					**06777300**
					SPO	
Actinomyces israelii		96.84	0.39	GLU 99	1	
Clostridium butyricum		3.16	-.08	RHA 7	99	
06777302	**Good identification**					**06777302**
Actinomyces israelii		95.23	0.42	GLU 99		
Clostridium clostridioforme		4.77	0.10	RHA 7		
06777342	**Good identification**					**06777342**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.35	GLU 99			
06777400	**Low discrimination**						**06777400**
					SPO		
Actinomyces israelii		87.28	0.46	GLU 99	1		
Clostridium butyricum		12.72	0.11		99		
06777402	**Acceptable identification**						**06777402**
					SPO		
Actinomyces israelii		93.16	0.49	GLU 99	1		
Clostridium butyricum		3.52	0.04		99		
Bifidobacterium adolescentis		2.22	0.15		1		
06777412	**Low discrimination**						**06777412**
Actinomyces israelii		60.79	0.35	GLU 99			
Clostridium clostridioforme		21.79	0.18	MLZ 13			
Bifidobacterium adolescentis		11.85	0.17				
06777440	**Good identification**						**06777440**
					SPO		
Actinomyces israelii		96.84	0.39	GLU 99	1		
Clostridium butyricum		3.16	-.07		99		
06777442	**Good identification**						**06777442**
Actinomyces israelii		100.0	0.42	GLU 99			
06777500	**Low discrimination**						**06777500**
					SPO		
Actinomyces israelii		53.09	0.57	GLU 99	1		
Clostridium butyricum		46.44	0.35		99		
Bifidobacterium adolescentis		0.47	0.15		1		
06777502	**Low discrimination**						**06777502**
					SPO		
Actinomyces israelii		54.65	0.60	GLU 99	1		
Bifidobacterium adolescentis		32.23	0.50		1		
Clostridium butyricum		12.39	0.28		99		
06777510	**Low discrimination**						**06777510**
					SPO		
Actinomyces israelii		57.09	0.43	GLU 99	1		
Bifidobacterium dentium		19.21	0.10	MLZ 13	1		
Clostridium butyricum		17.59	0.13		99		
06777512	**Low discrimination**						**06777512**
Bifidobacterium adolescentis		56.94	0.51	GLU 99			
Bifidobacterium dentium		26.77	0.27				
Actinomyces israelii		11.80	0.46				
06777522	**Low discrimination**						**06777522**
					SPO		
Actinomyces israelii		60.54	0.39	GLU 99	1		
Clostridium butyricum		18.70	0.10	NAG 6	99		
Clostridium clostridioforme		9.33	0.16		66		
06777540	**Low discrimination**						**06777540**
					COC	SPO	
Actinomyces israelii		61.53	0.50	GLU 99	1	1	
Blautia producta		26.43	0.24		99	1	
Clostridium butyricum		12.04	0.17		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
06777542	**Low discrimination**						**06777542**
					SPO		
Actinomyces israelii		78.74	0.53	GLU 99	1		
Bifidobacterium adolescentis		14.76	0.34		1		
Clostridium butyricum		3.99	0.10		99		
06777552	**Low discrimination**						**06777552**
					COC		
Bifidobacterium adolescentis		47.46	0.36	GLU 99	1		
Actinomyces israelii		30.95	0.39	SOR 11	1		
Blautia producta		19.77	0.15		99		
06777700	**Low discrimination**						**06777700**
					SPO		
Actinomyces israelii		73.56	0.38	GLU 99	1		
Clostridium butyricum		26.44	0.09	RHA 7	99		
06777702	**Low discrimination**						**06777702**
					SPO		
Actinomyces israelii		46.93	0.41	GLU 99	1		
Clostridium clostridioforme		44.98	0.31	RHA 7	66		
Clostridium butyricum		4.37	0.02		99		
06777712	**Acceptable identification**						**06777712**
Clostridium clostridioforme		93.74	0.42	GLU 88			
Actinomyces israelii		3.21	0.26	MAN 1			
Bifidobacterium adolescentis		2.08	0.17				
06777732	**Acceptable identification**						**06777732**
					GRA		
Clostridium clostridioforme		91.41	0.42	GLU 88	50		
Bacteroides ovatus		6.09	0.12	MAN 1	1		
Bifidobacterium dentium		1.69	-.03		99		
07000000	**Low discrimination**						**07000000**
					COC	SPO	
Gemella morbillorum		89.66	0.86		99	1	
Atopobium minutum		4.82	0.69		1	1	
Clostridium botulinum biovar A		2.78	0.57		1	99	
07000001	**Low discrimination**						**07000001**
					COC		
Gemella morbillorum		73.24	0.52	URE 1	99		
Anaerococcus prevotii		16.71	0.32		99		
Atopobium minutum		3.94	0.35		1		
07000002	**Low discrimination**						**07000002**
					COC	CAT	
Atopobium minutum		73.49	0.66	SUC 5	1	1	
Gemella morbillorum		20.71	0.52		99	1	
Propionibacterium granulosum		3.67	0.46		1	90	
07000010	**Low discrimination**						**07000010**
					COC	CAT	
Gemella morbillorum		81.08	0.52	MLZ 1	99	1	
Propionibacterium granulosum		7.83	0.41		1	90	
Atopobium minutum		4.36	0.35		1	1	
07000020	**Low discrimination**						**07000020**
					COC	SPO	
Gemella morbillorum		75.75	0.69	NAG 10	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		12.22	0.61		1	1	
Clostridium cadaveris		10.23	0.49		1	99	
07000021	**Low discrimination**						**07000021**
					COC	SPO	
Gemella morbillorum		72.43	0.35	NAG 10	99	1	
Atopobium minutum		11.69	0.26	URE 1	1	1	
Clostridium cadaveris		9.79	0.15		1	99	
07000022	**Low discrimination**						**07000022**
					COC	CAT	
Atopobium minutum		87.25	0.58	SUC 5	1	1	
Gemella morbillorum		8.20	0.35		99	1	
Propionibacterium granulosum		3.27	0.35		1	90	
07000030	**Low discrimination**						**07000030**
					COC	CAT	
Gemella morbillorum		62.96	0.35	MLZ 1	99	1	
Propionibacterium granulosum		13.68	0.31	NAG 10	1	90	
Atopobium minutum		10.16	0.26		1	1	
07000040	**Low discrimination**						**07000040**
					COC	SPO	
Clostridium botulinum biovar B		36.53	0.47	MLT 99	1	99	
Gemella morbillorum		29.51	0.52		99	1	
Clostridium botulinum biovar A		27.09	0.48		1	99	
07000050	**Acceptable identification**						**07000050**
					COC	SPO	
Clostridium botulinum biovar B		94.61	0.42	MLT 99	1	99	
Clostridium novyi biovar A		2.18	0.29		1	99	
Gemella morbillorum		1.57	0.17		99	1	
07000060	**Low discrimination**						**07000060**
					COC	SPO	
Clostridium botulinum biovar B		79.22	0.42	MLT 99	1	99	
Gemella morbillorum		14.44	0.35		99	1	
Atopobium minutum		2.33	0.26		1	1	
07000070	**Good identification**						**07000070**
					COC	SPO	
Clostridium botulinum biovar B		98.48	0.37	MLT 99	1	99	
Clostridium novyi biovar A		1.15	0.18		1	99	
Gemella morbillorum		0.37	0.01		99	1	
07000100	**Low discrimination**						**07000100**
					COC	GRA	
Gemella morbillorum		70.49	0.52	LAC 1	99	99	
Fusobacterium mortiferum		14.48	0.50		1	1	
Actinomyces meyeri		6.86	0.35		1	99	
07000200	**Low discrimination**						**07000200**
					COC	SPO	
Gemella morbillorum		88.82	0.52	RHA 1	99	1	
Atopobium minutum		4.78	0.35		1	1	
Clostridium botulinum biovar A		2.76	0.23		1	99	
07000220	**Low discrimination**						**07000220**
					COC	SPO	
Gemella morbillorum		74.31	0.35	RHA 1	99	1	
Atopobium minutum		11.99	0.26	NAG 10	1	1	
Clostridium cadaveris		10.04	0.15		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07000400	**Very good identification**						**07000400**
					COC	SPO	
Gemella morbillorum		98.95	0.69	ARA 10	99	1	
Atopobium minutum		0.48	0.35		1	1	
Clostridium botulinum biovar A		0.28	0.23		1	99	
07000401	**Good identification**						**07000401**
Gemella morbillorum		97.97	0.35	ARA 10			
Anaerococcus prevotii		2.03	-.03	URE 1			
07000402	**Low discrimination**						**07000402**
					COC	CAT	
Gemella morbillorum		74.70	0.35	ARA 10	99	1	
Atopobium minutum		24.10	0.32	BGL 1	1	1	
Propionibacterium granulosum		1.20	0.11		1	90	
07000410	**Good identification**						**07000410**
Gemella morbillorum		100.0	0.35	ARA 10			
07000420	**Good identification**						**07000420**
					COC	SPO	
Gemella morbillorum		97.24	0.53	ARA 10	99	1	
Atopobium minutum		1.43	0.26	NAG 10	1	1	
Clostridium cadaveris		1.19	0.15		1	99	
07000440	**Low discrimination**						**07000440**
					COC	SPO	
Gemella morbillorum		82.77	0.35	ARA 10	99	1	
Clostridium botulinum biovar B		9.32	0.13	SOR 1	1	99	
Clostridium botulinum biovar A		6.91	0.14		1	99	
07000500	**Low discrimination**						**07000500**
					COC	GRA	
Gemella morbillorum		80.80	0.35	LAC 1	99	99	
Actinomyces meyeri		17.69	0.25	ARA 10	1	99	
Fusobacterium mortiferum		1.51	0.16		1	1	
07000600	**Good identification**						**07000600**
Gemella morbillorum		100.0	0.35	RHA 1			
07001000	**Very good identification**						**07001000**
					COC	CAT	
Gemella morbillorum		95.06	0.81		99	1	
Atopobium minutum		3.16	0.61		1	1	
Propionibacterium granulosum		1.26	0.56		1	90	
07001001	**Acceptable identification**						**07001001**
					COC	SPO	
Gemella morbillorum		92.02	0.47	URE 1	99	1	
Atopobium minutum		3.06	0.26		1	1	
Clostridium chauvoei		2.68	0.19		1	99	
07001002	**Low discrimination**						**07001002**
					COC	CAT	
Atopobium minutum		59.94	0.58	SUC 5	1	1	
Gemella morbillorum		27.29	0.47		99	1	
Propionibacterium granulosum		8.97	0.46		1	90	
07001010	**Low discrimination**						**07001010**
					COC	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		82.00	0.47	MLZ 1	99	1		
Propionibacterium granulosum		14.71	0.41		1	90		
Atopobium minutum		2.73	0.26		1	1		
07001020	**Low discrimination**							**07001020**
					COC	CAT		
Gemella morbillorum		87.86	0.65	NAG 10	99	1		
Atopobium minutum		8.77	0.52		1	1		
Propionibacterium granulosum		2.63	0.46		1	90		
07001022	**Low discrimination**							**07001022**
					COC	CAT		
Atopobium minutum		75.73	0.49	SUC 5	1	1		
Gemella morbillorum		11.49	0.31		99	1		
Propionibacterium granulosum		8.50	0.35		1	90		
07001040	**Acceptable identification**							**07001040**
					COC	SPO		
Gemella morbillorum		91.33	0.47	SOR 1	99	1		
Atopobium minutum		3.04	0.26		1	1		
Clostridium botulinum biovar B		2.12	0.13		1	99		
07001100	**Low discrimination**							**07001100**
					COC	GRA		
Gemella morbillorum		47.51	0.47	LAC 1	99	99		
Fusobacterium mortiferum		27.19	0.53		1	1		
Actinomyces meyeri		12.88	0.38		1	99		
07001101	**Good identification**							**07001101**
					COC	GRA	SPO	
Clostridium chauvoei		99.36	0.54	MLT 99	1	99	99	
Gemella morbillorum		0.35	0.13		99	99	1	
Fusobacterium mortiferum		0.20	0.19		1	1	1	
07001102	**Low discrimination**							**07001102**
					GRA			
Lactobacillus fermentum		61.40	0.46	FRU 99	99			
Fusobacterium mortiferum		21.78	0.37		1			
Atopobium minutum		7.60	0.23		99			
07001103	**Good identification**							**07001103**
Clostridium chauvoei		100.0	0.41	MLT 99				
07001121	**Good identification**							**07001121**
					GRA	SPO		
Clostridium chauvoei		99.55	0.45	MLT 99	99	99		
Prevotella melaninogenica		0.45	-.01		1	1		
07001200	**Acceptable identification**							**07001200**
					COC	CAT		
Gemella morbillorum		92.90	0.47	RHA 1	99	1		
Propionibacterium granulosum		3.78	0.30		1	90		
Atopobium minutum		3.09	0.26		1	1		
07001400	**Very good identification**							**07001400**
					COC			
Gemella morbillorum		98.43	0.65	ARA 10	99			
Lactobacillus fermentum		0.98	0.42		1			
Atopobium minutum		0.30	0.26		1			
07001420	**Good identification**							**07001420**
					COC			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		95.82	0.48	ARA 10	99			
Lactobacillus fermentum		2.87	0.34	NAG 10	1			
Atopobium minutum		0.87	0.18		1			
07001500	**Low discrimination**							**07001500**
					COC			
Lactobacillus fermentum		49.26	0.46	FRU 99	1			
Gemella morbillorum		30.53	0.31		99			
Actinomyces meyeri		18.62	0.28		1			
07001502	**Good identification**							**07001502**
Lactobacillus fermentum		98.85	0.46	FRU 99				
Actinomyces meyeri		1.16	0.02					
07001520	**Low discrimination**							**07001520**
					COC			
Lactobacillus fermentum		64.19	0.38	FRU 99	1			
Gemella morbillorum		13.26	0.14		99			
Actinomyces meyeri		9.93	0.13		1			
07001522	**Acceptable identification**							**07001522**
Lactobacillus fermentum		92.89	0.38	FRU 99				
Bifidobacterium longum subsp. longum		7.11	0.12					
07002000	**Low discrimination**							**07002000**
					COC	GRA	CAT	
Gemella morbillorum		50.67	0.52	MAN 1	99	99	1	
Propionibacterium granulosum		27.07	0.54		1	99	90	
Megasphaera elsdenii		10.16	0.41		99	1	1	
07002002	**Low discrimination**							**07002002**
					COC	CAT	GRA	
Propionibacterium granulosum		65.78	0.44	FRU 99	1	90	99	
Atopobium minutum		17.65	0.32		1	1	99	
Megasphaera elsdenii		10.97	0.25		99	1	1	
07002010	**Low discrimination**							**07002010**
					COC	SPO	CAT	
Propionibacterium granulosum		53.71	0.39	FRU 99	1	1	90	
Clostridium difficile		36.42	0.24	MLZ 12	1	99	1	
Gemella morbillorum		7.45	0.17		99	1	1	
07002020	**Low discrimination**							**07002020**
					COC	CAT		
Propionibacterium granulosum		47.55	0.44	FRU 99	1	90		
Gemella morbillorum		39.55	0.35		99	1		
Atopobium minutum		6.38	0.26		1	1		
07002040	**Low discrimination**							**07002040**
					COC	GRA	SPO	
Megasphaera elsdenii		77.63	0.38	MNS 1	99	1	1	
Clostridium botulinum biovar B		7.26	0.13	FRU 95	1	99	99	
Gemella morbillorum		5.86	0.17		99	99	1	
07002400	**Acceptable identification**							**07002400**
					COC	GRA	CAT	
Gemella morbillorum		92.75	0.35	MAN 1	99	99	1	
Propionibacterium granulosum		4.51	0.20	ARA 10	1	99	90	
Megasphaera elsdenii		1.69	0.07		99	1	1	
07003000	**Low discrimination**							**07003000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	CAT	
Gemella morbillorum		48.94	0.47	MAN 1	99	1	
Propionibacterium granulosum		48.56	0.54		1	90	
Atopobium minutum		1.63	0.26		1	1	
07003002	**Low discrimination**						**07003002**
					COC	CAT	
Propionibacterium granulosum		85.65	0.44	FRU 99	1	90	
Atopobium minutum		7.66	0.23		1	1	
Gemella morbillorum		3.49	0.13		99	1	
07003010	**Acceptable identification**						**07003010**
					COC	CAT	
Propionibacterium granulosum		93.05	0.39	FRU 99	1	90	
Gemella morbillorum		6.95	0.13	MLZ 12	99	1	
07003020	**Low discrimination**						**07003020**
					COC	CAT	
Propionibacterium granulosum		67.00	0.44	FRU 99	1	90	
Gemella morbillorum		30.01	0.31		99	1	
Atopobium minutum		3.00	0.18		1	1	
07004000	**Very good identification**						**07004000**
					COC		
Gemella morbillorum		98.49	0.69	XYL 10	99		
Atopobium minutum		0.48	0.35		1		
Actinomyces meyeri		0.42	0.30		1		
07004001	**Good identification**						**07004001**
Gemella morbillorum		97.97	0.35	XYL 10			
Anaerococcus prevotii		2.03	-.03	URE 1			
07004002	**Low discrimination**						**07004002**
					COC	CAT	
Gemella morbillorum		73.97	0.35	XYL 10	99	1	
Atopobium minutum		23.87	0.32	BGL 1	1	1	
Propionibacterium granulosum		1.19	0.11		1	90	
07004010	**Good identification**						**07004010**
Gemella morbillorum		100.0	0.35	XYL 10			
07004020	**Good identification**						**07004020**
					COC	SPO	
Gemella morbillorum		96.74	0.53	XYL 10	99	1	
Atopobium minutum		1.42	0.26	NAG 10	1	1	
Clostridium cadaveris		1.19	0.15		1	99	
07004040	**Low discrimination**						**07004040**
					COC	SPO	
Gemella morbillorum		75.88	0.35	XYL 10	99	1	
Clostridium botulinum biovar B		8.54	0.13	SOR 1	1	99	
Clostridium glycolicum		8.32	0.15		1	99	
07004100	**Low discrimination**						**07004100**
					COC	GRA	
Actinomyces meyeri		59.88	0.40	MNS 1	1	99	
Gemella morbillorum		35.22	0.35	FRU 95	99	99	
Fusobacterium mortiferum		4.90	0.31		1	1	
07004200	**Good identification**						**07004200**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		100.0	0.35	XYL 10				
07004400	**Good identification**							**07004400**
					COC			
Gemella morbillorum		99.04	0.53	XYL 10	99			
Actinomyces meyeri		0.96	0.19	ARA 10	1			
07004420	**Good identification**							**07004420**
					COC			
Gemella morbillorum		98.83	0.37	XYL 10	99			
Actinomyces meyeri		1.17	0.05	ARA 10	1			
07005000	**Very good identification**							**07005000**
					COC			
Gemella morbillorum		97.68	0.65	XYL 10	99			
Actinomyces meyeri		1.17	0.33		1			
Lactobacillus fermentum		0.57	0.38		1			
07005020	**Good identification**							**07005020**
					COC			
Gemella morbillorum		95.78	0.48	XYL 10	99			
Lactobacillus fermentum		1.69	0.30	NAG 10	1			
Actinomyces meyeri		1.41	0.18		1			
07005100	**Low discrimination**							**07005100**
					COC			
Actinomyces meyeri		66.98	0.43	MNS 1	1			
Gemella morbillorum		14.14	0.31	FRU 95	99			
Lactobacillus fermentum		13.40	0.42		1			
07005102	**Low discrimination**							**07005102**
					GRA			
Lactobacillus fermentum		83.33	0.42	FRU 99	99			
Actinomyces meyeri		12.88	0.17		99			
Fusobacterium mortiferum		3.79	0.17		1			
07005400	**Acceptable identification**							**07005400**
					COC			
Gemella morbillorum		92.62	0.48	XYL 10	99			
Lactobacillus fermentum		4.89	0.38	ARA 10	1			
Actinomyces meyeri		2.49	0.22		1			
07005402	**Low discrimination**							**07005402**
					COC			
Lactobacillus fermentum		83.94	0.38	FRU 99	1			
Gemella morbillorum		16.06	0.14		99			
07005502	**Acceptable identification**							**07005502**
Lactobacillus fermentum		93.35	0.42	FRU 99				
Actinomyces meyeri		3.61	0.07					
Bifidobacterium longum subsp. longum		3.04	0.09					
07006000	**Acceptable identification**							**07006000**
					COC	GRA	CAT	
Gemella morbillorum		92.75	0.35	MAN 1	99	99	1	
Propionibacterium granulosum		4.51	0.20	XYL 10	1	99	90	
Megasphaera elsdenii		1.69	0.07		99	1	1	
07010000	**Low discrimination**							**07010000**
					COC	GRA	CAT	
Gemella morbillorum		54.58	0.87		99	99	1	
Propionibacterium granulosum		31.63	0.90		1	99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		3.93	0.78		1	1	5	
07010001	**Low discrimination**							**07010001**
					COC	GRA	CAT	
Gemella morbillorum		54.50	0.53	URE 1	99	99	1	
Propionibacterium granulosum		31.58	0.56		1	99	90	
Fusobacterium mortiferum		3.93	0.44		1	1	5	
07010002	**Low discrimination**							**07010002**
					COC	CAT		
Propionibacterium granulosum		74.13	0.80		1	90		
Atopobium minutum		13.85	0.65		1	1		
Gemella morbillorum		5.17	0.53		99	1		
07010003	**Low discrimination**							**07010003**
					COC	CAT		
Propionibacterium granulosum		74.49	0.46	URE 1	1	90		
Atopobium minutum		13.92	0.31		1	1		
Gemella morbillorum		5.19	0.19		99	1		
07010010	**Low discrimination**							**07010010**
					COC	SPO	CAT	
Propionibacterium granulosum		82.00	0.75	MLZ 12	1	1	90	
Gemella morbillorum		10.48	0.53		99	1	1	
Clostridium difficile		4.66	0.41		1	99	1	
07010011	**Low discrimination**							**07010011**
					COC	SPO	CAT	
Propionibacterium granulosum		82.72	0.41	MLZ 12	1	1	90	
Gemella morbillorum		10.58	0.19	URE 1	99	1	1	
Clostridium difficile		4.70	0.07		1	99	1	
07010012	**Very good identification**							**07010012**
					COC	CAT		
Propionibacterium granulosum		97.35	0.65	MLZ 12	1	90		
Atopobium minutum		1.35	0.31		1	1		
Gemella morbillorum		0.50	0.19		99	1		
07010020	**Low discrimination**							**07010020**
					COC	CAT		
Propionibacterium granulosum		50.62	0.80		1	90		
Gemella morbillorum		38.83	0.71		99	1		
Atopobium minutum		4.73	0.60		1	1		
07010021	**Low discrimination**							**07010021**
					COC	CAT		
Propionibacterium granulosum		51.22	0.46	URE 1	1	90		
Gemella morbillorum		39.29	0.37		99	1		
Atopobium minutum		4.79	0.26		1	1		
07010022	**Low discrimination**							**07010022**
					COC	CAT		
Propionibacterium granulosum		77.30	0.69		1	90		
Atopobium minutum		19.26	0.57		1	1		
Gemella morbillorum		2.40	0.37		99	1		
07010023	**Low discrimination**							**07010023**
					COC	CAT		
Propionibacterium granulosum		78.12	0.35	URE 1	1	90		
Atopobium minutum		19.46	0.23		1	1		
Gemella morbillorum		2.42	0.03		99	1		
07010030	**Good identification**							**07010030**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	CAT		
Propionibacterium granulosum		92.77	0.65	MLZ 12	1	90		
Gemella morbillorum		5.27	0.37		99	1		
Atopobium minutum		0.64	0.26		1	1		
07010032	**Good identification**							**07010032**
					COC	CAT		
Propionibacterium granulosum		97.97	0.55	MLZ 12	1	90		
Atopobium minutum		1.81	0.23		1	1		
Gemella morbillorum		0.23	0.03		99	1		
07010040	**Low discrimination**							**07010040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		50.43	0.58	MLT 99	1	99	99	
Megasphaera elsdenii		25.87	0.60		99	1	1	
Gemella morbillorum		12.45	0.53		99	99	1	
07010042	**Low discrimination**							**07010042**
					COC	SPO	CAT	
Megasphaera elsdenii		49.41	0.44	MNS 1	99	1	1	
Propionibacterium granulosum		30.99	0.46	BGL 10	1	1	90	
Clostridium botulinum biovar B		8.76	0.24		1	99	1	
07010050	**Acceptable identification**							**07010050**
					SPO	CAT		
Clostridium botulinum biovar B		93.67	0.52	MLT 99	99	1		
Propionibacterium granulosum		3.71	0.41		1	90		
Clostridium difficile		1.10	0.19		99	1		
07010060	**Low discrimination**							**07010060**
					COC	SPO	CAT	
Clostridium botulinum biovar B		85.67	0.52	MLT 99	1	99	1	
Propionibacterium granulosum		6.22	0.46		1	1	90	
Gemella morbillorum		4.77	0.37		99	1	1	
07010062	**Low discrimination**							**07010062**
					SPO	CAT		
Propionibacterium granulosum		51.86	0.35	SOR 1	1	90		
Clostridium botulinum biovar B		28.87	0.18		99	1		
Atopobium minutum		12.92	0.23		1	1		
07010070	**Good identification**							**07010070**
					COC	SPO	CAT	
Clostridium botulinum biovar B		97.92	0.47	MLT 99	1	99	1	
Propionibacterium granulosum		1.97	0.31		1	1	90	
Gemella morbillorum		0.11	0.03		99	1	1	
07010100	**Low discrimination**							**07010100**
					COC	GRA		
Fusobacterium mortiferum		83.25	0.84		1	1		
Actinomyces meyeri		7.57	0.57		1	99		
Gemella morbillorum		5.00	0.53		99	99		
07010101	**Low discrimination**							**07010101**
					GRA	SPO		
Fusobacterium mortiferum		73.34	0.50	URE 1	1	1		
Clostridium chauvoei		13.04	0.26		99	99		
Actinomyces meyeri		6.67	0.23		99	1		
07010102	**Low discrimination**							**07010102**
					GRA	CAT		
Fusobacterium mortiferum		85.34	0.68	BGL 10	1	5		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		6.69	0.46		99	90		
Lactobacillus fermentum		3.65	0.46		99	1		
07010110	**Low discrimination**							**07010110**
					GRA	CAT		
Fusobacterium mortiferum		60.69	0.50	MLZ 1	1	5		
Propionibacterium granulosum		28.53	0.41		99	90		
Actinomyces meyeri		5.52	0.23		99	1		
07010120	**Low discrimination**							**07010120**
					GRA	CAT		
Actinomyces meyeri		26.61	0.42	MNS 1	99	1		
Fusobacterium mortiferum		21.69	0.50	NAG 12	1	5		
Propionibacterium granulosum		18.69	0.46		99	90		
07010122	**Low discrimination**							**07010122**
					GRA	CAT		
Propionibacterium granulosum		31.11	0.35	LAC 1	99	90		
Lactobacillus fermentum		22.62	0.38		99	1		
Fusobacterium mortiferum		16.04	0.34		1	5		
07010140	**Low discrimination**							**07010140**
					COC	GRA	SPO	
Fusobacterium mortiferum		64.13	0.50	SOR 1	1	1	1	
Clostridium botulinum biovar B		15.61	0.24		1	99	99	
Megasphaera elsdenii		8.01	0.26		99	1	1	
07010200	**Low discrimination**							**07010200**
					COC	GRA	CAT	
Propionibacterium granulosum		58.61	0.64	RHA 3	1	99	90	
Gemella morbillorum		33.03	0.53		99	99	1	
Fusobacterium mortiferum		2.38	0.44		1	1	5	
07010202	**Low discrimination**							**07010202**
					COC	CAT		
Propionibacterium granulosum		89.94	0.54	RHA 3	1	90		
Atopobium minutum		5.49	0.31		1	1		
Gemella morbillorum		2.05	0.19		99	1		
07010210	**Acceptable identification**							**07010210**
					COC	SPO	CAT	
Propionibacterium granulosum		94.31	0.49	RHA 3	1	1	90	
Gemella morbillorum		3.94	0.19	MLZ 12	99	1	1	
Clostridium difficile		1.75	0.07		1	99	1	
07010212	**Good identification**							**07010212**
Propionibacterium granulosum		100.0	0.39	RHA 3				
07010220	**Low discrimination**							**07010220**
					COC	CAT		
Propionibacterium granulosum		76.28	0.54	RHA 3	1	90		
Gemella morbillorum		19.11	0.37		99	1		
Atopobium minutum		2.33	0.26		1	1		
07010222	**Acceptable identification**							**07010222**
					COC	CAT		
Propionibacterium granulosum		91.62	0.44	RHA 3	1	90		
Atopobium minutum		7.46	0.23		1	1		
Gemella morbillorum		0.93	0.03		99	1		
07010230	**Good identification**							**07010230**
					COC	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		98.18	0.39	RHA 3	1	90		
Gemella morbillorum		1.82	0.03	MLZ 12	99	1		
07010300	**Low discrimination**							**07010300**
					GRA	CAT		
Fusobacterium mortiferum		79.52	0.50	RHA 1	1	5		
Propionibacterium granulosum		8.48	0.30		99	90		
Actinomyces meyeri		7.23	0.23		99	1		
07010400	**Good identification**							**07010400**
					COC	CAT		
Gemella morbillorum		91.90	0.71	ARA 10	99	1		
Propionibacterium granulosum		4.84	0.56		1	90		
Actinomyces meyeri		0.79	0.36		1	1		
07010401	**Acceptable identification**							**07010401**
					COC	CAT		
Gemella morbillorum		94.23	0.37	ARA 10	99	1		
Propionibacterium granulosum		4.96	0.22	URE 1	1	90		
Actinomyces meyeri		0.81	0.02		1	1		
07010402	**Low discrimination**							**07010402**
					COC	CAT		
Propionibacterium granulosum		41.88	0.46	ARA 1	1	90		
Gemella morbillorum		32.12	0.37		99	1		
Lactobacillus fermentum		14.00	0.42		1	1		
07010410	**Low discrimination**							**07010410**
					COC	SPO	CAT	
Gemella morbillorum		56.81	0.37	ARA 10	99	1	1	
Propionibacterium granulosum		40.40	0.41	MLZ 1	1	1	90	
Clostridium difficile		2.30	0.07		1	99	1	
07010420	**Low discrimination**							**07010420**
					COC	CAT		
Gemella morbillorum		85.89	0.54	ARA 10	99	1		
Propionibacterium granulosum		10.18	0.46	NAG 10	1	90		
Lactobacillus fermentum		1.13	0.34		1	1		
07010422	**Low discrimination**							**07010422**
					COC	CAT		
Propionibacterium granulosum		49.13	0.35	ARA 1	1	90		
Lactobacillus fermentum		21.89	0.34		1	1		
Gemella morbillorum		16.75	0.20		99	1		
07010440	**Low discrimination**							**07010440**
					COC	GRA	SPO	
Gemella morbillorum		61.78	0.37	ARA 10	99	99	1	
Clostridium botulinum biovar B		22.76	0.24	SOR 1	1	99	99	
Megasphaera elsdenii		11.68	0.26		99	1	1	
07010500	**Low discrimination**							**07010500**
					COC	GRA		
Actinomyces meyeri		50.95	0.47	MNS 1	1	99		
Fusobacterium mortiferum		22.65	0.50		1	1		
Gemella morbillorum		14.97	0.37		99	99		
07010502	**Low discrimination**							**07010502**
					GRA			
Lactobacillus fermentum		70.56	0.46	GAL 99	99			
Fusobacterium mortiferum		16.68	0.34		1			
Actinomyces meyeri		10.45	0.21		99			

Code Taxon	ID score	T index	tests against	Differentiation tests	Code

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07010522	**Acceptable identification**							**07010522**
Lactobacillus fermentum		94.29	0.38	GAL 99				
Actinomyces meyeri		5.71	0.06					
07010600	**Low discrimination**							**07010600**
					COC	CAT		
Gemella morbillorum		85.48	0.37	RHA 1	99	1		
Propionibacterium granulosum		13.79	0.30	ARA 10	1	90		
Actinomyces meyeri		0.74	0.02		1	1		
07011000	**Low discrimination**							**07011000**
					COC	GRA	CAT	
Propionibacterium granulosum		44.57	0.90		1	99	90	
Gemella morbillorum		41.42	0.83		99	99	1	
Fusobacterium mortiferum		8.31	0.81		1	1	5	
07011001	**Low discrimination**							**07011001**
					COC	GRA	CAT	
Propionibacterium granulosum		43.52	0.56	URE 1	1	99	90	
Gemella morbillorum		40.44	0.48		99	99	1	
Fusobacterium mortiferum		8.12	0.47		1	1	5	
07011002	**Low discrimination**							**07011002**
					GRA	CAT		
Propionibacterium granulosum		65.75	0.80		99	90		
Lactobacillus fermentum		21.97	0.76		99	1		
Fusobacterium mortiferum		5.45	0.65		1	5		
07011003	**Low discrimination**							**07011003**
					GRA	CAT		
Propionibacterium granulosum		63.46	0.46	URE 1	99	90		
Lactobacillus fermentum		21.21	0.42		99	1		
Fusobacterium mortiferum		5.26	0.30		1	5		
07011010	**Good identification**							**07011010**
					COC	GRA	CAT	
Propionibacterium granulosum		91.44	0.75	MLZ 12	1	99	90	
Gemella morbillorum		6.29	0.48		99	99	1	
Fusobacterium mortiferum		1.26	0.47		1	1	5	
07011011	**Acceptable identification**							**07011011**
					COC	GRA	CAT	
Propionibacterium granulosum		92.37	0.41	MLZ 12	1	99	90	
Gemella morbillorum		6.36	0.14	URE 1	99	99	1	
Fusobacterium mortiferum		1.28	0.13		1	1	5	
07011012	**Very good identification**							**07011012**
					GRA	CAT		
Propionibacterium granulosum		96.04	0.65	MLZ 12	99	90		
Lactobacillus fermentum		2.38	0.42		99	1		
Fusobacterium mortiferum		0.59	0.30		1	5		
07011020	**Low discrimination**							**07011020**
					COC	CAT		
Propionibacterium granulosum		62.14	0.80		1	90		
Gemella morbillorum		25.66	0.66		99	1		
Lactobacillus fermentum		6.92	0.68		1	1		
07011021	**Low discrimination**							**07011021**
					COC	CAT		
Propionibacterium granulosum		59.09	0.46	URE 1	1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		24.40	0.32		99	1		
Lactobacillus fermentum		6.58	0.34		1	1		
07011022	**Low discrimination**							**07011022**
					CAT			
Propionibacterium granulosum		62.06	0.69		90			
Lactobacillus fermentum		27.65	0.68		1			
Atopobium minutum		5.15	0.49		1			
07011023	**Low discrimination**							**07011023**
					CAT			
Propionibacterium granulosum		57.95	0.35	URE 1	90			
Lactobacillus fermentum		25.82	0.34		1			
Actinomyces naeslundii		5.23	0.32		25			
07011030	**Very good identification**							**07011030**
					COC	CAT		
Propionibacterium granulosum		95.64	0.65	MLZ 12	1	90		
Gemella morbillorum		2.93	0.32		99	1		
Lactobacillus fermentum		0.79	0.34		1	1		
07011032	**Good identification**							**07011032**
					CAT			
Propionibacterium granulosum		95.56	0.55	MLZ 12	90			
Lactobacillus fermentum		3.15	0.34		1			
Atopobium minutum		0.59	0.15		1			
07011040	**Low discrimination**							**07011040**
					COC	GRA	CAT	
Propionibacterium granulosum		42.52	0.56	SOR 1	1	99	90	
Gemella morbillorum		39.51	0.48		99	99	1	
Fusobacterium mortiferum		7.93	0.47		1	1	5	
07011042	**Low discrimination**							**07011042**
					GRA	CAT		
Propionibacterium granulosum		64.87	0.46	SOR 1	99	90		
Lactobacillus fermentum		21.68	0.42		99	1		
Fusobacterium mortiferum		5.38	0.30		1	5		
07011050	**Low discrimination**							**07011050**
					COC	SPO	CAT	
Propionibacterium granulosum		74.75	0.41	MLZ 12	1	1	90	
Clostridium botulinum biovar B		19.07	0.18	SOR 1	1	99	1	
Gemella morbillorum		5.15	0.14		99	1	1	
07011060	**Low discrimination**							**07011060**
					COC	CAT		
Propionibacterium granulosum		41.04	0.46	SOR 1	1	90		
Clostridium perfringens		23.87	0.34		1	1		
Gemella morbillorum		16.95	0.32		99	1		
07011100	**Low discrimination**							**07011100**
					GRA			
Fusobacterium mortiferum		68.67	0.87		1			
Lactobacillus fermentum		21.51	0.80		99			
Actinomyces meyeri		6.24	0.60		99			
07011101	**Low discrimination**							**07011101**
					GRA	SPO		
Clostridium chauvoei		88.73	0.60	MLT 99	99	99		
Fusobacterium mortiferum		7.56	0.53		1	1		
Lactobacillus fermentum		2.37	0.46		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07011102	**Low discrimination**						**07011102**
					GRA	CAT	
Lactobacillus fermentum		71.75	0.80		99	1	
Fusobacterium mortiferum		25.45	0.71		1	5	
Propionibacterium granulosum		1.33	0.46		99	90	
07011103	**Low discrimination**						**07011103**
					GRA	SPO	
Clostridium chauvoei		81.47	0.47	MLT 99	99	99	
Lactobacillus fermentum		12.32	0.46		99	1	
Fusobacterium mortiferum		4.37	0.37		1	1	
07011110	**Low discrimination**						**07011110**
					GRA	CAT	
Fusobacterium mortiferum		57.25	0.53	MLZ 1	1	5	
Propionibacterium granulosum		17.94	0.41		99	90	
Lactobacillus fermentum		17.93	0.46		99	1	
07011112	**Low discrimination**						**07011112**
					GRA	CAT	
Lactobacillus fermentum		62.32	0.46	MLZ 1	99	1	
Fusobacterium mortiferum		22.10	0.37		1	5	
Propionibacterium granulosum		15.58	0.31		99	90	
07011120	**Low discrimination**						**07011120**
					GRA		
Clostridium perfringens		38.84	0.61	MLT 99	99		
Lactobacillus fermentum		31.21	0.72		99		
Prevotella melaninogenica		20.25	0.50		1		
07011121	**Acceptable identification**						**07011121**
					SPO		
Clostridium chauvoei		91.42	0.52	MLT 99	99		
Clostridium perfringens		3.04	0.27		84		
Lactobacillus fermentum		2.44	0.38		1		
07011122	**Low discrimination**						**07011122**
					GRA		
Lactobacillus fermentum		56.94	0.72		99		
Clostridium perfringens		33.34	0.56		99		
Prevotella melaninogenica		7.04	0.37		1		
07011123	**Low discrimination**						**07011123**
					SPO		
Clostridium chauvoei		76.22	0.39	MLT 99	99		
Lactobacillus fermentum		11.53	0.38		1		
Clostridium perfringens		6.75	0.22		84		
07011132	**Low discrimination**						**07011132**
					CAT		
Lactobacillus fermentum		52.72	0.38	MLZ 1	1		
Clostridium perfringens		30.88	0.22		1		
Propionibacterium granulosum		9.89	0.21		90		
07011140	**Low discrimination**						**07011140**
					GRA		
Fusobacterium mortiferum		65.25	0.53	SOR 1	1		
Lactobacillus fermentum		20.44	0.46		99		
Actinomyces meyeri		5.93	0.26		99		
07011142	**Low discrimination**						**07011142**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		65.90	0.46	SOR 1	99			
Fusobacterium mortiferum		23.37	0.37		1			
Clostridium perfringens		7.50	0.18		99			
07011160	**Acceptable identification**							**07011160**
					GRA			
Clostridium perfringens		92.29	0.49	MLT 99	99			
Lactobacillus fermentum		3.93	0.38		99			
Prevotella melaninogenica		2.55	0.15		1			
07011162	**Acceptable identification**							**07011162**
					GRA			
Clostridium perfringens		90.46	0.43	MLT 99	99			
Lactobacillus fermentum		8.19	0.38		99			
Prevotella melaninogenica		1.01	0.03		1			
07011200	**Low discrimination**							**07011200**
					COC	GRA	CAT	
Propionibacterium granulosum		71.14	0.64	RHA 3	1	99	90	
Gemella morbillorum		21.59	0.48		99	99	1	
Fusobacterium mortiferum		4.33	0.47		1	1	5	
07011202	**Low discrimination**							**07011202**
					GRA	CAT		
Propionibacterium granulosum		85.56	0.54	RHA 3	99	90		
Lactobacillus fermentum		9.34	0.42		99	1		
Fusobacterium mortiferum		2.32	0.30		1	5		
07011210	**Good identification**							**07011210**
					COC	GRA	CAT	
Propionibacterium granulosum		97.37	0.49	RHA 3	1	99	90	
Gemella morbillorum		2.19	0.14	MLZ 12	99	99	1	
Fusobacterium mortiferum		0.44	0.13		1	1	5	
07011212	**Good identification**							**07011212**
Propionibacterium granulosum		100.0	0.39	RHA 3				
07011220	**Low discrimination**							**07011220**
					COC	CAT		
Propionibacterium granulosum		83.59	0.54	RHA 3	1	90		
Gemella morbillorum		11.28	0.32		99	1		
Lactobacillus fermentum		3.04	0.34		1	1		
07011222	**Low discrimination**							**07011222**
					CAT			
Propionibacterium granulosum		83.92	0.44	RHA 3	90			
Lactobacillus fermentum		12.21	0.34		1			
Atopobium minutum		2.28	0.15		1			
07011230	**Good identification**							**07011230**
Propionibacterium granulosum		100.0	0.39	RHA 3				
07011300	**Low discrimination**							**07011300**
					GRA			
Fusobacterium mortiferum		66.63	0.53	RHA 1	1			
Lactobacillus fermentum		20.87	0.46		99			
Actinomyces meyeri		6.06	0.26		99			
07011302	**Low discrimination**							**07011302**
					GRA	CAT		
Lactobacillus fermentum		70.85	0.46	RHA 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		25.13	0.37		1	5	
Propionibacterium granulosum		4.02	0.20		99	90	
07011322	**Low discrimination**						**07011322**
					GRA		
Lactobacillus fermentum		43.37	0.38	RHA 1	99		
Clostridium perfringens		25.40	0.22		99		
Prevotella oralis		16.33	0.15		1		
07011400	**Low discrimination**						**07011400**
					COC	CAT	
Gemella morbillorum		51.22	0.66	ARA 10	99	1	
Lactobacillus fermentum		41.45	0.76		1	1	
Propionibacterium granulosum		5.01	0.56		1	90	
07011402	**Very good identification**						**07011402**
					COC	CAT	
Lactobacillus fermentum		95.42	0.76		1	1	
Propionibacterium granulosum		2.88	0.46		1	90	
Gemella morbillorum		1.19	0.32		99	1	
07011403	**Good identification**						**07011403**
					CAT		
Lactobacillus fermentum		97.07	0.42	URE 1	1		
Propionibacterium granulosum		2.93	0.11		90		
07011410	**Low discrimination**						**07011410**
					COC	CAT	
Propionibacterium granulosum		41.64	0.41	ARA 1	1	90	
Gemella morbillorum		31.52	0.32	MLZ 12	99	1	
Lactobacillus fermentum		25.51	0.42		1	1	
07011412	**Low discrimination**						**07011412**
					CAT		
Lactobacillus fermentum		71.02	0.42	MLZ 1	1		
Propionibacterium granulosum		28.98	0.31		90		
07011420	**Low discrimination**						**07011420**
					COC	CAT	
Lactobacillus fermentum		65.72	0.68		1	1	
Gemella morbillorum		27.07	0.50		99	1	
Propionibacterium granulosum		5.96	0.46		1	90	
07011422	**Very good identification**						**07011422**
					COC	CAT	
Lactobacillus fermentum		97.09	0.68		1	1	
Propionibacterium granulosum		2.20	0.35		1	90	
Gemella morbillorum		0.40	0.16		99	1	
07011442	**Good identification**						**07011442**
					CAT		
Lactobacillus fermentum		97.07	0.42	SOR 1	1		
Propionibacterium granulosum		2.93	0.11		90		
07011500	**Low discrimination**						**07011500**
					GRA		
Lactobacillus fermentum		89.78	0.80		99		
Actinomyces meyeri		6.51	0.50		99		
Fusobacterium mortiferum		2.90	0.53		1		
07011501	**Low discrimination**						**07011501**
					SPO		
Lactobacillus fermentum		67.42	0.46	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		25.51	0.26		99	
Actinomyces meyeri		4.89	0.16		1	
07011502	**Very good identification**					**07011502**
					GRA	
Lactobacillus fermentum		99.35	0.80		99	
Fusobacterium mortiferum		0.36	0.37		1	
Actinomyces meyeri		0.22	0.24		99	
07011503	**Acceptable identification**					**07011503**
					SPO	
Lactobacillus fermentum		93.74	0.46	URE 1	1	
Clostridium chauvoei		6.26	0.13		99	
07011510	**Low discrimination**					**07011510**
Lactobacillus fermentum		83.16	0.46	MLZ 1		
Bifidobacterium longum subsp. longum		8.13	0.22			
Actinomyces meyeri		6.03	0.16			
07011512	**Acceptable identification**					**07011512**
Lactobacillus fermentum		91.09	0.46	MLZ 1		
Bifidobacterium longum subsp. longum		8.91	0.22			
07011520	**Good identification**					**07011520**
Lactobacillus fermentum		94.27	0.72			
Actinomyces meyeri		2.80	0.35			
Clostridium perfringens		1.19	0.27			
07011521	**Low discrimination**					**07011521**
					SPO	
Lactobacillus fermentum		72.55	0.38	URE 1	1	
Clostridium chauvoei		27.45	0.18		99	
07011522	**Very good identification**					**07011522**
Lactobacillus fermentum		98.44	0.72			
Bifidobacterium longum subsp. longum		0.69	0.28			
Clostridium perfringens		0.58	0.22			
07011523	**Acceptable identification**					**07011523**
					SPO	
Lactobacillus fermentum		93.74	0.38	URE 1	1	
Clostridium chauvoei		6.26	0.05		99	
07011540	**Acceptable identification**					**07011540**
					GRA	
Lactobacillus fermentum		90.52	0.46	SOR 1	99	
Actinomyces meyeri		6.57	0.16		99	
Fusobacterium mortiferum		2.92	0.19		1	
07011542	**Good identification**					**07011542**
Lactobacillus fermentum		100.0	0.46	SOR 1		
07011560	**Low discrimination**					**07011560**
Lactobacillus fermentum		80.84	0.38	SOR 1		
Clostridium perfringens		19.16	0.15			
07011562	**Low discrimination**					**07011562**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		89.96	0.38	SOR 1				
Clostridium perfringens		10.04	0.09					
07011602	**Acceptable identification**							**07011602**
					CAT			
Lactobacillus fermentum		91.53	0.42	RHA 1	1			
Propionibacterium granulosum		8.47	0.20		90			
07011700	**Acceptable identification**							**07011700**
					GRA			
Lactobacillus fermentum		90.52	0.46	RHA 1	99			
Actinomyces meyeri		6.57	0.16		99			
Fusobacterium mortiferum		2.92	0.19		1			
07011702	**Good identification**							**07011702**
Lactobacillus fermentum		100.0	0.46	RHA 1				
07011720	**Low discrimination**							**07011720**
					GRA			
Lactobacillus fermentum		82.70	0.38	RHA 1	99			
Bacteroides vulgatus		17.30	0.05		1			
07011722	**Acceptable identification**							**07011722**
					GRA			
Lactobacillus fermentum		90.18	0.38	RHA 1	99			
Parabacteroides distasonis		9.82	0.18		1			
07012000	**Low discrimination**							**07012000**
					COC	CAT	GRA	
Propionibacterium granulosum		73.76	0.88		1	90	99	
Eubacterium limosum		17.29	0.63		1	1	99	
Megasphaera elsdenii		5.31	0.63		99	1	1	
07012001	**Low discrimination**							**07012001**
					COC	CAT	GRA	
Propionibacterium granulosum		74.02	0.54	URE 1	1	90	99	
Eubacterium limosum		17.35	0.29		1	1	99	
Megasphaera elsdenii		5.33	0.29		99	1	1	
07012002	**Very good identification**							**07012002**
					COC	CAT	GRA	
Propionibacterium granulosum		95.42	0.78		1	90	99	
Megasphaera elsdenii		3.06	0.47		99	1	1	
Eubacterium limosum		0.90	0.29		1	1	99	
07012003	**Good identification**							**07012003**
					COC	CAT	GRA	
Propionibacterium granulosum		96.02	0.44	URE 1	1	90	99	
Megasphaera elsdenii		3.08	0.12		99	1	1	
Eubacterium limosum		0.91	-.06		1	1	99	
07012010	**Low discrimination**							**07012010**
					SPO	CAT		
Propionibacterium granulosum		58.73	0.73	MLZ 12	1	90		
Clostridium difficile		39.82	0.58		99	1		
Eubacterium limosum		1.02	0.29		1	1		
07012011	**Low discrimination**							**07012011**
					SPO	CAT		
Propionibacterium granulosum		58.98	0.39	MLZ 12	1	90		
Clostridium difficile		39.99	0.24	URE 1	99	1		
Eubacterium limosum		1.02	-.06		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07012012	**Very good identification**							**07012012**
					COC	SPO	CAT	
Propionibacterium granulosum		97.04	0.63	MLZ 12	1	1	90	
Clostridium difficile		2.66	0.24		1	99	1	
Megasphaera elsdenii		0.23	0.12		99	1	1	
07012020	**Very good identification**							**07012020**
					COC	CAT		
Propionibacterium granulosum		96.82	0.78		1	90		
Gemella morbillorum		0.99	0.37		99	1		
Eubacterium limosum		0.92	0.29		1	1		
07012021	**Good identification**							**07012021**
					COC	CAT		
Propionibacterium granulosum		98.06	0.44	URE 1	1	90		
Gemella morbillorum		1.01	0.03		99	1		
Eubacterium limosum		0.93	-.06		1	1		
07012022	**Very good identification**							**07012022**
					COC	CAT	GRA	
Propionibacterium granulosum		99.38	0.67		1	90	99	
Atopobium minutum		0.33	0.23		1	1	99	
Megasphaera elsdenii		0.13	0.12		99	1	1	
07012030	**Very good identification**							**07012030**
					COC	SPO	CAT	
Propionibacterium granulosum		97.20	0.63	MLZ 12	1	1	90	
Clostridium difficile		2.66	0.24		1	99	1	
Gemella morbillorum		0.07	0.03		99	1	1	
07012032	**Good identification**							**07012032**
Propionibacterium granulosum		100.0	0.53	MLZ 12				
07012040	**Low discrimination**							**07012040**
					COC	CAT	GRA	
Megasphaera elsdenii		75.86	0.60	MNS 1	99	1	1	
Propionibacterium granulosum		15.95	0.54		1	90	99	
Eubacterium limosum		3.74	0.29		1	1	99	
07012042	**Low discrimination**							**07012042**
					COC	CAT	GRA	
Megasphaera elsdenii		66.84	0.44	MNS 1	99	1	1	
Propionibacterium granulosum		31.63	0.44	BGL 10	1	90	99	
Pseudoramibacter alactolyticus		0.86	0.00		1	1	99	
07012050	**Low discrimination**							**07012050**
					COC	SPO	CAT	
Clostridium difficile		67.41	0.36	SUC 1	1	99	1	
Propionibacterium granulosum		19.08	0.39	SOR 5	1	1	90	
Megasphaera elsdenii		6.72	0.26		99	1	1	
07012060	**Low discrimination**							**07012060**
					COC	SPO	CAT	
Propionibacterium granulosum		64.36	0.44	SOR 1	1	1	90	
Megasphaera elsdenii		12.37	0.26		99	1	1	
Clostridium botulinum biovar B		11.87	0.18		1	99	1	
07012100	**Low discrimination**							**07012100**
					GRA	CAT		
Propionibacterium granulosum		50.84	0.54	LAC 1	99	90		
Fusobacterium mortiferum		19.55	0.50		1	5		

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Eubacterium limosum		11.92	0.29		99	1		
07012102	**Low discrimination**							**07012102**
					GRA	CAT		
Propionibacterium propionicum		49.32	0.49	NIT 86	99	1		
Propionibacterium granulosum		41.81	0.44	MLT 99	99	90		
Fusobacterium mortiferum		7.15	0.34		1	5		
07012110	**Low discrimination**							**07012110**
					GRA	SPO	CAT	
Propionibacterium granulosum		57.52	0.39	LAC 1	99	1	90	
Clostridium difficile		39.00	0.24	MLZ 12	99	99	1	
Fusobacterium mortiferum		1.64	0.16		1	1	5	
07012120	**Acceptable identification**							**07012120**
					GRA	CAT		
Propionibacterium granulosum		94.14	0.44	LAC 1	99	90		
Actinomyces meyeri		1.80	0.08		99	1		
Fusobacterium mortiferum		1.46	0.16		1	5		
07012140	**Low discrimination**							**07012140**
					COC	CAT	GRA	
Propionibacterium propionicum		75.33	0.46	NIT 86	1	1	99	
Megasphaera elsdenii		18.40	0.26	MLT 99	99	1	1	
Propionibacterium granulosum		3.87	0.20		1	90	99	
07012142	**Good identification**							**07012142**
					COC	CAT	GRA	
Propionibacterium propionicum		97.40	0.49	NIT 86	1	1	99	
Megasphaera elsdenii		1.76	0.09	MLT 99	99	1	1	
Propionibacterium granulosum		0.83	0.09		1	90	99	
07012200	**Low discrimination**							**07012200**
					COC	CAT	GRA	
Propionibacterium granulosum		89.72	0.62	RHA 3	1	90	99	
Eubacterium limosum		6.87	0.29		1	1	99	
Megasphaera elsdenii		2.11	0.29		99	1	1	
07012202	**Good identification**							**07012202**
					COC	CAT	GRA	
Propionibacterium granulosum		98.66	0.52	RHA 3	1	90	99	
Megasphaera elsdenii		1.03	0.12		99	1	1	
Eubacterium limosum		0.31	-.06		1	1	99	
07012210	**Low discrimination**							**07012210**
					SPO	CAT		
Propionibacterium granulosum		81.49	0.47	RHA 3	1	90		
Clostridium difficile		18.05	0.24	MLZ 12	99	1		
Eubacterium limosum		0.46	-.06		1	1		
07012212	**Good identification**							**07012212**
Propionibacterium granulosum		100.0	0.37	RHA 3				
07012220	**Good identification**							**07012220**
					COC	CAT		
Propionibacterium granulosum		99.36	0.52	RHA 3	1	90		
Gemella morbillorum		0.33	0.03		99	1		
Eubacterium limosum		0.31	-.06		1	1		
07012222	**Good identification**							**07012222**
Propionibacterium granulosum		100.0	0.42	RHA 3				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07012230	**Good identification**							**07012230**
Propionibacterium granulosum		100.0	0.37	RHA 3				
07012400	**Low discrimination**							**07012400**
					COC	CAT		
Propionibacterium granulosum		63.12	0.54	ARA 1	1	90		
Gemella morbillorum		16.04	0.37		99	1		
Eubacterium limosum		14.80	0.29		1	1		
07012402	**Good identification**							**07012402**
					COC	GRA	CAT	
Propionibacterium granulosum		95.08	0.44	ARA 1	1	99	90	
Megasphaera elsdenii		3.05	0.12		99	1	1	
Gemella morbillorum		0.98	0.03		99	99	1	
07012410	**Low discrimination**							**07012410**
					COC	SPO	CAT	
Propionibacterium granulosum		58.34	0.39	ARA 1	1	1	90	
Clostridium difficile		39.55	0.24	MLZ 12	1	99	1	
Gemella morbillorum		1.10	0.03		99	1	1	
07012420	**Low discrimination**							**07012420**
					COC	CAT		
Propionibacterium granulosum		89.09	0.44	ARA 1	1	90		
Gemella morbillorum		10.06	0.20		99	1		
Eubacterium limosum		0.84	-.06		1	1		
07013000	**Very good identification**							**07013000**
					COC	GRA	CAT	
Propionibacterium granulosum		97.86	0.88		1	99	90	
Gemella morbillorum		1.22	0.48		99	99	1	
Fusobacterium mortiferum		0.24	0.47		1	1	5	
07013001	**Good identification**							**07013001**
					COC	CAT		
Propionibacterium granulosum		96.92	0.54	URE 1	1	90		
Actinomyces naeslundii		1.41	0.37		1	25		
Gemella morbillorum		1.21	0.14		99	1		
07013002	**Very good identification**							**07013002**
					SPO	CAT		
Propionibacterium granulosum		98.52	0.78		1	90		
Clostridium innocuum		0.71	0.32		99	1		
Lactobacillus fermentum		0.44	0.42		1	1		
07013003	**Acceptable identification**							**07013003**
Propionibacterium granulosum		94.51	0.44	URE 1				
Actinomyces naeslundii		5.49	0.37					
07013010	**Very good identification**							**07013010**
					COC	SPO	CAT	
Propionibacterium granulosum		99.14	0.73	MLZ 12	1	1	90	
Clostridium difficile		0.68	0.24		1	99	1	
Gemella morbillorum		0.09	0.14		99	1	1	
07013011	**Good identification**							**07013011**
Propionibacterium granulosum		100.0	0.39	MLZ 12				
07013012	**Very good identification**							**07013012**
					SPO	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		99.78	0.63	MLZ 12	1	90		
Clostridium innocuum		0.22	0.08		99	1		
07013020	**Very good identification**							**07013020**
					COC	CAT		
Propionibacterium granulosum		98.65	0.78		1	90		
Gemella morbillorum		0.55	0.32		99	1		
Propionibacterium acnes		0.52	0.29		1	95		
07013021	**Acceptable identification**							**07013021**
Propionibacterium granulosum		94.51	0.44	URE 1				
Actinomyces naeslundii		5.49	0.37					
07013022	**Very good identification**							**07013022**
					CAT			
Propionibacterium granulosum		99.07	0.67		90			
Lactobacillus fermentum		0.59	0.34		1			
Actinomyces naeslundii		0.16	0.34		25			
07013030	**Very good identification**							**07013030**
Propionibacterium granulosum		100.0	0.63	MLZ 12				
07013032	**Good identification**							**07013032**
Propionibacterium granulosum		100.0	0.53	MLZ 12				
07013040	**Acceptable identification**							**07013040**
					COC	GRA	CAT	
Propionibacterium granulosum		93.69	0.54	SOR 1	1	99	90	
Megasphaera elsdenii		4.50	0.26		99	1	1	
Gemella morbillorum		1.17	0.14		99	99	1	
07013042	**Good identification**							**07013042**
					COC	CAT	GRA	
Propionibacterium granulosum		96.78	0.44	SOR 1	1	90	99	
Megasphaera elsdenii		2.07	0.09		99	1	1	
Propionibacterium propionicum		1.15	0.15		1	1	99	
07013050	**Good identification**							**07013050**
					SPO	CAT		
Propionibacterium granulosum		96.56	0.39	MLZ 12	1	90		
Clostridium difficile		3.45	0.02	SOR 1	99	1		
07013060	**Low discrimination**							**07013060**
Propionibacterium granulosum		89.77	0.44	SOR 1				
Propionibacterium acnes		10.23	0.18					
07013100	**Low discrimination**							**07013100**
					GRA	CAT		
Propionibacterium granulosum		48.75	0.54	LAC 1	99	90		
Fusobacterium mortiferum		28.12	0.53		1	5		
Propionibacterium propionicum		9.59	0.46		99	1		
07013102	**Low discrimination**							**07013102**
					CAT			
Propionibacterium propionicum		36.09	0.49	NIT 86	1			
Propionibacterium granulosum		30.59	0.44	MLT 99	90			
Lactobacillus fermentum		22.11	0.46		1			
07013103	**Low discrimination**							**07013103**
					SPO			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		46.59	0.45	MLT 94	1			
Propionibacterium propionicum		36.95	0.33	RAF 92	1			
Clostridium chauvoei		13.61	0.13		99			
07013110	**Acceptable identification**							**07013110**
					GRA	CAT		
Propionibacterium granulosum		94.58	0.39	LAC 1	99	90		
Fusobacterium mortiferum		4.04	0.19	MLZ 12	1	5		
Propionibacterium propionicum		1.38	0.12		99	1		
07013120	**Low discrimination**							**07013120**
					CAT			
Propionibacterium granulosum		53.56	0.44	LAC 1	90			
Clostridium perfringens		16.06	0.27		1			
Lactobacillus fermentum		12.90	0.38		1			
07013121	**Low discrimination**							**07013121**
					SPO	CAT		
Actinomyces naeslundii		62.01	0.45	MLT 94	1	25		
Clostridium chauvoei		34.20	0.18	RAF 92	99	1		
Propionibacterium granulosum		3.79	0.09		1	90		
07013123	**Acceptable identification**							**07013123**
					SPO			
Actinomyces naeslundii		91.13	0.45	MLT 94	1			
Clostridium chauvoei		8.87	0.05	RAF 92	99			
07013140	**Acceptable identification**							**07013140**
					GRA	CAT		
Propionibacterium propionicum		90.44	0.46	NIT 86	99	1		
Propionibacterium granulosum		4.65	0.20	MLT 99	99	90		
Fusobacterium mortiferum		2.68	0.19		1	5		
07013142	**Good identification**							**07013142**
					CAT			
Propionibacterium propionicum		97.56	0.49	NIT 86	1			
Actinomyces naeslundii		1.61	0.29	MLT 99	25			
Propionibacterium granulosum		0.84	0.09		90			
07013200	**Very good identification**							**07013200**
					COC	GRA	CAT	
Propionibacterium granulosum		99.44	0.62	RHA 3	1	99	90	
Gemella morbillorum		0.40	0.14		99	99	1	
Fusobacterium mortiferum		0.08	0.13		1	1	5	
07013202	**Good identification**							**07013202**
Propionibacterium granulosum		100.0	0.52	RHA 3				
07013210	**Good identification**							**07013210**
Propionibacterium granulosum		100.0	0.47	RHA 3				
07013212	**Good identification**							**07013212**
Propionibacterium granulosum		100.0	0.37	RHA 3				
07013220	**Good identification**							**07013220**
Propionibacterium granulosum		100.0	0.52	RHA 3				
07013222	**Good identification**							**07013222**
Propionibacterium granulosum		100.0	0.42	RHA 3				

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07013230	**Good identification**							**07013230**
Propionibacterium granulosum		100.0	0.37	RHA 3				
07013400	**Low discrimination**							**07013400**
					COC	CAT		
Propionibacterium granulosum		77.69	0.54	ARA 1	1	90		
Gemella morbillorum		10.63	0.32		99	1		
Lactobacillus fermentum		8.61	0.42		1	1		
07013402	**Low discrimination**							**07013402**
					SPO	CAT		
Propionibacterium granulosum		63.76	0.44	ARA 1	1	90		
Lactobacillus fermentum		28.25	0.42		1	1		
Clostridium innocuum		7.99	0.19		99	1		
07013410	**Good identification**							**07013410**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07013420	**Low discrimination**							**07013420**
					COC	CAT		
Propionibacterium granulosum		82.75	0.44	ARA 1	1	90		
Lactobacillus fermentum		12.22	0.34		1	1		
Gemella morbillorum		5.03	0.16		99	1		
07013500	**Low discrimination**							**07013500**
					CAT			
Lactobacillus fermentum		85.35	0.46	MAN 1	1			
Actinomyces meyeri		6.19	0.16		1			
Propionibacterium granulosum		4.77	0.20		90			
07013502	**Good identification**							**07013502**
					CAT			
Lactobacillus fermentum		97.04	0.46	MAN 1	1			
Propionibacterium propionicum		1.60	0.15		1			
Propionibacterium granulosum		1.36	0.09		90			
07013520	**Good identification**							**07013520**
					CAT			
Lactobacillus fermentum		95.98	0.38	MAN 1	1			
Propionibacterium granulosum		4.02	0.09		90			
07013522	**Good identification**							**07013522**
Lactobacillus fermentum		100.0	0.38	MAN 1				
07014000	**Low discrimination**							**07014000**
					COC	CAT		
Gemella morbillorum		82.78	0.71	XYL 10	99	1		
Actinomyces meyeri		5.53	0.52		1	1		
Propionibacterium granulosum		4.36	0.56		1	90		
07014001	**Low discrimination**							**07014001**
					COC	CAT		
Gemella morbillorum		83.67	0.37	XYL 10	99	1		
Actinomyces meyeri		5.58	0.18	URE 1	1	1		
Propionibacterium granulosum		4.41	0.22		1	90		
07014002	**Low discrimination**							**07014002**
					COC	GRA	CAT	
Propionibacterium granulosum		36.19	0.46	XYL 1	1	99	90	
Gemella morbillorum		27.75	0.37		99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		14.90	0.42		1	1	5	
07014010	**Low discrimination**							**07014010**
					COC	SPO	CAT	
Gemella morbillorum		48.44	0.37	XYL 10	99	1	1	
Propionibacterium granulosum		34.45	0.41	MLZ 1	1	1	90	
Clostridium difficile		10.20	0.19		1	99	1	
07014020	**Low discrimination**							**07014020**
					COC	CAT		
Gemella morbillorum		80.84	0.54	XYL 10	99	1		
Propionibacterium granulosum		9.58	0.46	NAG 10	1	90		
Actinomyces meyeri		6.62	0.37		1	1		
07014022	**Low discrimination**							**07014022**
					COC	CAT		
Propionibacterium granulosum		51.62	0.35	XYL 1	1	90		
Gemella morbillorum		17.60	0.20		99	1		
Lactobacillus fermentum		13.51	0.30		1	1		
07014040	**Low discrimination**							**07014040**
					COC	SPO		
Gemella morbillorum		39.06	0.37	XYL 10	99	1		
Clostridium glycolicum		31.55	0.31	SOR 1	1	99		
Clostridium botulinum biovar B		14.39	0.24		1	99		
07014100	**Low discrimination**							**07014100**
					COC	GRA		
Actinomyces meyeri		67.42	0.62	MNS 1	1	99		
Fusobacterium mortiferum		28.76	0.65		1	1		
Gemella morbillorum		2.55	0.37		99	99		
07014102	**Low discrimination**							**07014102**
					GRA			
Fusobacterium mortiferum		49.89	0.49	XYL 7	1			
Actinomyces meyeri		32.56	0.36	BGL 10	99			
Lactobacillus fermentum		16.63	0.42		99			
07014120	**Acceptable identification**							**07014120**
					GRA			
Actinomyces meyeri		90.52	0.47	MNS 1	99			
Lactobacillus fermentum		3.50	0.34	NAG 12	99			
Fusobacterium mortiferum		2.86	0.31		1			
07014200	**Low discrimination**							**07014200**
					COC	CAT		
Gemella morbillorum		76.70	0.37	XYL 10	99	1		
Propionibacterium granulosum		12.37	0.30	RHA 1	1	90		
Actinomyces meyeri		5.12	0.18		1	1		
07014400	**Low discrimination**							**07014400**
					COC			
Gemella morbillorum		84.38	0.54	XYL 10	99			
Actinomyces meyeri		12.67	0.41	ARA 10	1			
Lactobacillus fermentum		1.96	0.38		1			
07014402	**Low discrimination**							**07014402**
					COC			
Lactobacillus fermentum		59.34	0.38	GAL 99	1			
Gemella morbillorum		25.76	0.20		99			
Actinomyces meyeri		11.85	0.16		1			
07014420	**Low discrimination**							**07014420**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Gemella morbillorum		79.06	0.38	XYL 10	99		
Actinomyces meyeri		14.57	0.27	ARA 10	1		
Lactobacillus fermentum		5.52	0.30	NAG 10	1		
07014500	**Acceptable identification**						**07014500**
					GRA		
Actinomyces meyeri		91.14	0.52	MNS 1	99		
Lactobacillus fermentum		5.76	0.42		99		
Fusobacterium mortiferum		1.57	0.31		1		
07014502	**Low discrimination**						**07014502**
					GRA		
Lactobacillus fermentum		65.80	0.42	GAL 99	99		
Actinomyces meyeri		32.21	0.26		99		
Fusobacterium mortiferum		1.99	0.14		1		
07014520	**Low discrimination**						**07014520**
					COC		
Actinomyces meyeri		85.60	0.37	MNS 1	1		
Lactobacillus fermentum		13.22	0.34	NAG 12	1		
Gemella morbillorum		1.17	0.04		99		
07015000	**Low discrimination**						**07015000**
					COC		
Gemella morbillorum		52.67	0.66	XYL 10	99		
Lactobacillus fermentum		25.03	0.72		1		
Actinomyces meyeri		9.79	0.55		1		
07015002	**Low discrimination**						**07015002**
					GRA	CAT	
Lactobacillus fermentum		89.30	0.72		99	1	
Propionibacterium granulosum		4.60	0.46		99	90	
Fusobacterium mortiferum		2.84	0.45		1	5	
07015003	**Acceptable identification**						**07015003**
					GRA	CAT	
Lactobacillus fermentum		92.32	0.38	URE 1	99	1	
Propionibacterium granulosum		4.75	0.11		99	90	
Fusobacterium mortiferum		2.93	0.11		1	5	
07015010	**Low discrimination**						**07015010**
					COC	CAT	
Propionibacterium granulosum		42.36	0.41	XYL 1	1	90	
Gemella morbillorum		32.07	0.32	MLZ 12	99	1	
Lactobacillus fermentum		15.24	0.38		1	1	
07015012	**Low discrimination**						**07015012**
					GRA	CAT	
Lactobacillus fermentum		57.92	0.38	MLZ 1	99	1	
Propionibacterium granulosum		40.24	0.31		99	90	
Fusobacterium mortiferum		1.84	0.11		1	5	
07015020	**Low discrimination**						**07015020**
					COC		
Lactobacillus fermentum		49.14	0.64		1		
Gemella morbillorum		34.46	0.50		99		
Actinomyces meyeri		7.86	0.40		1		
07015022	**Good identification**						**07015022**
					COC	CAT	
Lactobacillus fermentum		94.60	0.64		1	1	

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Propionibacterium granulosum		3.65	0.35		1	90	
Gemella morbillorum		0.67	0.16		99	1	
07015042	**Acceptable identification**						**07015042**
					GRA	CAT	
Lactobacillus fermentum		92.32	0.38	SOR 1	99	1	
Propionibacterium granulosum		4.75	0.11		99	90	
Fusobacterium mortiferum		2.93	0.11		1	5	
07015100	**Low discrimination**						**07015100**
					GRA		
Lactobacillus fermentum		41.94	0.76		99		
Actinomyces meyeri		40.22	0.65		99		
Fusobacterium mortiferum		17.16	0.68		1		
07015101	**Low discrimination**						**07015101**
					SPO		
Lactobacillus fermentum		33.20	0.42	URE 1	1		
Actinomyces meyeri		31.84	0.31		1		
Clostridium chauvoei		21.39	0.26		99		
07015102	**Good identification**						**07015102**
					GRA		
Lactobacillus fermentum		92.96	0.76		99		
Fusobacterium mortiferum		4.23	0.52		1		
Actinomyces meyeri		2.76	0.39		99		
07015103	**Low discrimination**						**07015103**
					GRA	SPO	
Lactobacillus fermentum		84.12	0.42	URE 1	99	1	
Clostridium chauvoei		9.56	0.13		99	99	
Fusobacterium mortiferum		3.82	0.17		1	1	
07015110	**Low discrimination**						**07015110**
					GRA		
Lactobacillus fermentum		42.23	0.42	MLZ 1	99		
Actinomyces meyeri		40.50	0.31		99		
Fusobacterium mortiferum		17.28	0.34		1		
07015112	**Acceptable identification**						**07015112**
					GRA		
Lactobacillus fermentum		93.01	0.42	MLZ 1	99		
Fusobacterium mortiferum		4.23	0.17		1		
Actinomyces meyeri		2.76	0.05		99		
07015120	**Low discrimination**						**07015120**
Lactobacillus fermentum		69.20	0.68				
Actinomyces meyeri		27.15	0.50				
Clostridium perfringens		1.48	0.27				
07015122	**Very good identification**						**07015122**
Lactobacillus fermentum		96.16	0.68				
Actinomyces meyeri		1.17	0.24				
Clostridium perfringens		0.97	0.22				
07015140	**Low discrimination**						**07015140**
					GRA		
Lactobacillus fermentum		42.23	0.42	SOR 1	99		
Actinomyces meyeri		40.50	0.31		99		
Fusobacterium mortiferum		17.28	0.34		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07015142	**Acceptable identification**						**07015142**
					GRA		
Lactobacillus fermentum		93.01	0.42	SOR 1	99		
Fusobacterium mortiferum		4.23	0.17		1		
Actinomyces meyeri		2.76	0.05		99		
07015202	**Low discrimination**						**07015202**
					GRA	CAT	
Lactobacillus fermentum		84.08	0.38	RHA 1	99	1	
Propionibacterium granulosum		13.25	0.20		99	90	
Fusobacterium mortiferum		2.67	0.11		1	5	
07015300	**Low discrimination**						**07015300**
					GRA		
Lactobacillus fermentum		42.23	0.42	RHA 1	99		
Actinomyces meyeri		40.50	0.31		99		
Fusobacterium mortiferum		17.28	0.34		1		
07015302	**Acceptable identification**						**07015302**
					GRA		
Lactobacillus fermentum		93.01	0.42	RHA 1	99		
Fusobacterium mortiferum		4.23	0.17		1		
Actinomyces meyeri		2.76	0.05		99		
07015322	**Low discrimination**						**07015322**
					GRA		
Parabacteroides distasonis		71.61	0.38	RAF 99	1		
Lactobacillus fermentum		26.30	0.34		99		
Prevotella oralis		2.08	-.01		1		
07015400	**Low discrimination**						**07015400**
					COC		
Lactobacillus fermentum		74.82	0.72		1		
Gemella morbillorum		17.49	0.50		99		
Actinomyces meyeri		7.32	0.44		1		
07015401	**Low discrimination**						**07015401**
					COC		
Lactobacillus fermentum		75.10	0.38	URE 1	1		
Gemella morbillorum		17.56	0.16		99		
Actinomyces meyeri		7.35	0.10		1		
07015402	**Very good identification**						**07015402**
					COC		
Lactobacillus fermentum		99.38	0.72		1		
Actinomyces meyeri		0.30	0.19		1		
Gemella morbillorum		0.24	0.16		99		
07015403	**Good identification**						**07015403**
Lactobacillus fermentum		100.0	0.38	URE 1			
07015410	**Low discrimination**						**07015410**
					COC		
Lactobacillus fermentum		75.10	0.38	MLZ 1	1		
Gemella morbillorum		17.56	0.16		99		
Actinomyces meyeri		7.35	0.10		1		
07015412	**Good identification**						**07015412**
Lactobacillus fermentum		100.0	0.38	MLZ 1			
07015420	**Low discrimination**						**07015420**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Lactobacillus fermentum		89.33	0.64		1	
Gemella morbillorum		6.96	0.34		99	
Actinomyces meyeri		3.57	0.30		1	
07015422	**Very good identification**					**07015422**
Lactobacillus fermentum		99.88	0.64			
Actinomyces meyeri		0.12	0.04			
07015440	**Low discrimination**					**07015440**
					COC	
Lactobacillus fermentum		75.10	0.38	SOR 1	1	
Gemella morbillorum		17.56	0.16		99	
Actinomyces meyeri		7.35	0.10		1	
07015442	**Good identification**					**07015442**
Lactobacillus fermentum		100.0	0.38	SOR 1		
07015500	**Low discrimination**					**07015500**
					GRA	
Lactobacillus fermentum		80.11	0.76		99	
Actinomyces meyeri		19.21	0.55		99	
Fusobacterium mortiferum		0.33	0.34		1	
07015501	**Low discrimination**					**07015501**
Lactobacillus fermentum		80.66	0.42	URE 1		
Actinomyces meyeri		19.34	0.21			
07015502	**Very good identification**					**07015502**
Lactobacillus fermentum		98.85	0.76			
Actinomyces meyeri		0.73	0.29			
Bifidobacterium longum subsp. longum		0.29	0.26			
07015503	**Good identification**					**07015503**
Lactobacillus fermentum		100.0	0.42	URE 1		
07015510	**Low discrimination**					**07015510**
Lactobacillus fermentum		57.49	0.42	MLZ 1		
Bifidobacterium longum subsp. longum		28.72	0.30			
Actinomyces meyeri		13.79	0.21			
07015512	**Low discrimination**					**07015512**
Lactobacillus fermentum		62.76	0.42	MLZ 1		
Bifidobacterium longum subsp. longum		31.35	0.30			
Bifidobacterium adolescentis		5.89	0.19			
07015520	**Low discrimination**					**07015520**
Lactobacillus fermentum		86.71	0.68			
Actinomyces meyeri		8.51	0.40			
Bifidobacterium longum subsp. longum		3.08	0.36			
07015522	**Very good identification**					**07015522**
					GRA	
Lactobacillus fermentum		95.77	0.68		99	
Bifidobacterium longum subsp. longum		3.41	0.36		99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		0.54	0.27		1			
07015530	**Low discrimination**							**07015530**
Bifidobacterium longum subsp. longum		84.52	0.40	MLT 99				
Lactobacillus fermentum		14.10	0.34	RAF 97				
Actinomyces meyeri		1.38	0.06					
07015532	**Low discrimination**							**07015532**
					GRA			
Bifidobacterium longum subsp. longum		82.18	0.40	MLT 99	99			
Lactobacillus fermentum		13.71	0.34	RAF 97	99			
Parabacteroides distasonis		4.11	0.22		1			
07015540	**Low discrimination**							**07015540**
Lactobacillus fermentum		80.66	0.42	SOR 1				
Actinomyces meyeri		19.34	0.21					
07015542	**Good identification**							**07015542**
Lactobacillus fermentum		100.0	0.42	SOR 1				
07015600	**Low discrimination**							**07015600**
					COC			
Lactobacillus fermentum		75.10	0.38	RHA 1	1			
Gemella morbillorum		17.56	0.16		99			
Actinomyces meyeri		7.35	0.10		1			
07015602	**Good identification**							**07015602**
Lactobacillus fermentum		100.0	0.38	RHA 1				
07015700	**Low discrimination**							**07015700**
					GRA			
Lactobacillus fermentum		77.01	0.42	RHA 1	99			
Actinomyces meyeri		18.47	0.21		99			
Bacteroides vulgatus		4.53	-.01		1			
07015702	**Good identification**							**07015702**
Lactobacillus fermentum		100.0	0.42	RHA 1				
07016000	**Low discrimination**							**07016000**
					COC	CAT		
Propionibacterium granulosum		58.89	0.54	XYL 1	1	90		
Gemella morbillorum		14.97	0.37		99	1		
Eubacterium limosum		13.80	0.29		1	1		
07016002	**Good identification**							**07016002**
					COC	GRA	CAT	
Propionibacterium granulosum		95.08	0.44	XYL 1	1	99	90	
Megasphaera elsdenii		3.05	0.12		99	1	1	
Gemella morbillorum		0.98	0.03		99	99	1	
07016010	**Low discrimination**							**07016010**
					COC	SPO	CAT	
Clostridium difficile		77.32	0.36	SUC 1	1	99	1	
Propionibacterium granulosum		21.89	0.39	XYL 5	1	1	90	
Gemella morbillorum		0.41	0.03		99	1	1	
07016020	**Low discrimination**							**07016020**
					COC	CAT		
Propionibacterium granulosum		88.36	0.44	XYL 1	1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		9.98	0.20		99	1		
Eubacterium limosum		0.84	-.06		1	1		
07017000	**Low discrimination**							**07017000**
					COC	CAT		
Propionibacterium granulosum		79.51	0.54	XYL 1	1	90		
Gemella morbillorum		10.88	0.32		99	1		
Lactobacillus fermentum		5.17	0.38		1	1		
07017002	**Low discrimination**							**07017002**
					SPO	CAT		
Propionibacterium granulosum		76.58	0.44	XYL 1	1	90		
Lactobacillus fermentum		19.92	0.38		1	1		
Clostridium innocuum		2.86	0.10		99	1		
07017010	**Good identification**							**07017010**
					SPO	CAT		
Propionibacterium granulosum		96.56	0.39	XYL 1	1	90		
Clostridium difficile		3.45	0.02	MLZ 12	99	1		
07017020	**Low discrimination**							**07017020**
					COC	CAT		
Propionibacterium granulosum		86.10	0.44	XYL 1	1	90		
Lactobacillus fermentum		7.47	0.30		1	1		
Gemella morbillorum		5.24	0.16		99	1		
07017100	**Low discrimination**							**07017100**
					GRA			
Lactobacillus fermentum		40.29	0.42	MAN 1	99			
Actinomyces meyeri		38.64	0.31		99			
Fusobacterium mortiferum		16.48	0.34		1			
07017102	**Low discrimination**							**07017102**
					GRA			
Lactobacillus fermentum		88.73	0.42	MAN 1	99			
Fusobacterium mortiferum		4.03	0.17		1			
Actinomyces meyeri		2.63	0.05		99			
07017400	**Low discrimination**							**07017400**
					COC	CAT		
Lactobacillus fermentum		67.26	0.38	MAN 1	1	1		
Gemella morbillorum		15.72	0.16		99	1		
Propionibacterium granulosum		10.44	0.20		1	90		
07017402	**Good identification**							**07017402**
					CAT			
Lactobacillus fermentum		96.26	0.38	MAN 1	1			
Propionibacterium granulosum		3.74	0.09		90			
07017500	**Low discrimination**							**07017500**
Lactobacillus fermentum		80.66	0.42	MAN 1				
Actinomyces meyeri		19.34	0.21					
07017502	**Acceptable identification**							**07017502**
Lactobacillus fermentum		94.09	0.42	MAN 1				
Bifidobacterium adolescentis		5.91	0.16					
07020000	**Low discrimination**							**07020000**
					COC	SPO	CAT	
Propionibacterium granulosum		52.67	0.64	FRU 99	1	1	90	
Gemella morbillorum		24.79	0.52		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		19.05	0.47		1	99	1	
07020002	**Acceptable identification**							**07020002**
					COC	CAT		
Propionibacterium granulosum		90.14	0.54	FRU 99	1	90		
Atopobium minutum		6.08	0.32		1	1		
Gemella morbillorum		1.71	0.17		99	1		
07020010	**Low discrimination**							**07020010**
					COC	SPO	CAT	
Propionibacterium granulosum		88.86	0.49	FRU 99	1	1	90	
Clostridium botulinum biovar B		5.67	0.16	MLZ 12	1	99	1	
Gemella morbillorum		3.10	0.17		99	1	1	
07020012	**Good identification**							**07020012**
Propionibacterium granulosum		100.0	0.39	FRU 99				
07020020	**Low discrimination**							**07020020**
					COC	SPO	CAT	
Propionibacterium granulosum		69.47	0.54	FRU 99	1	1	90	
Gemella morbillorum		14.53	0.35		99	1	1	
Clostridium cadaveris		10.23	0.27		1	99	1	
07020022	**Acceptable identification**							**07020022**
					COC	CAT		
Propionibacterium granulosum		91.04	0.43	FRU 99	1	90		
Atopobium minutum		8.19	0.23		1	1		
Gemella morbillorum		0.77	0.01		99	1		
07020030	**Low discrimination**							**07020030**
					COC	SPO	CAT	
Propionibacterium granulosum		87.63	0.39	FRU 99	1	1	90	
Clostridium botulinum biovar B		11.01	0.11	MLZ 12	1	99	1	
Gemella morbillorum		1.36	0.01		99	1	1	
07020040	**Acceptable identification**							**07020040**
					SPO	CAT		
Clostridium botulinum biovar B		93.40	0.56	MLT 99	99	1		
Clostridium botulinum biovar A		5.77	0.38		99	1		
Propionibacterium granulosum		0.54	0.30		1	90		
07020050	**Good identification**							**07020050**
					SPO	CAT		
Clostridium botulinum biovar B		99.71	0.50	MLT 99	99	1		
Propionibacterium granulosum		0.16	0.15		1	90		
Clostridium botulinum biovar A		0.13	0.04		99	1		
07020060	**Good identification**							**07020060**
					SPO	CAT		
Clostridium botulinum biovar B		99.52	0.50	MLT 99	99	1		
Propionibacterium granulosum		0.29	0.20		1	90		
Clostridium botulinum biovar A		0.13	0.04		99	1		
07020070	**Good identification**							**07020070**
Clostridium botulinum biovar B		100.0	0.45	MLT 99				
07020100	**Low discrimination**							**07020100**
					COC	GRA	CAT	
Fusobacterium mortiferum		60.95	0.41	FRU 99	1	1	5	
Propionibacterium granulosum		21.31	0.30		1	99	90	
Gemella morbillorum		10.03	0.17		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07020200	**Low discrimination**							**07020200**
					COC	SPO	CAT	
Propionibacterium granulosum		78.63	0.38	FRU 99	1	1	90	
Gemella morbillorum		12.09	0.17	RHA 3	99	1	1	
Clostridium botulinum biovar A		9.29	0.13		1	99	1	
07020400	**Low discrimination**							**07020400**
					COC	SPO	CAT	
Gemella morbillorum		79.18	0.35	TRE 1	99	1	1	
Propionibacterium granulosum		15.29	0.30	ARA 10	1	1	90	
Clostridium botulinum biovar A		5.53	0.13		1	99	1	
07021000	**Low discrimination**							**07021000**
					COC	GRA	CAT	
Propionibacterium granulosum		77.88	0.64	FRU 99	1	99	90	
Gemella morbillorum		19.74	0.47		99	99	1	
Fusobacterium mortiferum		1.45	0.38		1	1	5	
07021002	**Good identification**							**07021002**
					COC	CAT		
Propionibacterium granulosum		96.06	0.54	FRU 99	1	90		
Atopobium minutum		2.16	0.23		1	1		
Gemella morbillorum		0.98	0.13		99	1		
07021010	**Good identification**							**07021010**
					COC	CAT		
Propionibacterium granulosum		98.16	0.49	FRU 99	1	90		
Gemella morbillorum		1.84	0.13	MLZ 12	99	1		
07021012	**Good identification**							**07021012**
Propionibacterium granulosum		100.0	0.39	FRU 99				
07021020	**Low discrimination**							**07021020**
					COC	CAT		
Propionibacterium granulosum		87.49	0.54	FRU 99	1	90		
Gemella morbillorum		9.85	0.31		99	1		
Clostridium perfringens		1.67	0.17		1	1		
07021022	**Acceptable identification**							**07021022**
					CAT			
Propionibacterium granulosum		93.81	0.43	FRU 99	90			
Clostridium perfringens		3.37	0.11		1			
Atopobium minutum		2.81	0.15		1			
07021030	**Good identification**							**07021030**
Propionibacterium granulosum		100.0	0.39	FRU 99				
07021100	**Low discrimination**							**07021100**
					COC	GRA	CAT	
Fusobacterium mortiferum		70.81	0.44	FRU 99	1	1	5	
Propionibacterium granulosum		16.51	0.30		1	99	90	
Atopobium parvulum		4.26	0.12		99	99	1	
07021101	**Good identification**							**07021101**
					SPO			
Clostridium chauvoei		99.33	0.37	MLT 99	99			
Actinomyces naeslundii		0.67	0.23	TRE 10	1			
07021200	**Acceptable identification**							**07021200**
					COC	CAT		
Propionibacterium granulosum		92.36	0.38	FRU 99	1	90		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		7.64	0.13	RHA 3	99	1		
07022000	**Very good identification**							**07022000**
					COC	SPO	CAT	
Propionibacterium granulosum		98.70	0.62	FRU 99	1	1	90	
Gemella morbillorum		0.62	0.17		99	1	1	
Clostridium botulinum biovar A		0.48	0.13		1	99	1	
07022002	**Good identification**							**07022002**
Propionibacterium granulosum		100.0	0.52	FRU 99				
07022010	**Good identification**							**07022010**
					SPO	CAT		
Propionibacterium granulosum		98.33	0.47	FRU 99	1	90		
Clostridium difficile		1.67	0.04	MLZ 12	99	1		
07022012	**Good identification**							**07022012**
Propionibacterium granulosum		100.0	0.37	FRU 99				
07022020	**Good identification**							**07022020**
					COC	CAT		
Propionibacterium granulosum		99.72	0.52	FRU 99	1	90		
Gemella morbillorum		0.28	0.01		99	1		
07022022	**Good identification**							**07022022**
Propionibacterium granulosum		100.0	0.41	FRU 99				
07022030	**Good identification**							**07022030**
Propionibacterium granulosum		100.0	0.37	FRU 99				
07022200	**Good identification**							**07022200**
Propionibacterium granulosum		100.0	0.36	FRU 99				
07023000	**Very good identification**							**07023000**
					COC	SPO	CAT	
Propionibacterium granulosum		99.55	0.62	FRU 99	1	1	90	
Gemella morbillorum		0.34	0.13		99	1	1	
Clostridium innocuum		0.11	0.02		1	99	1	
07023002	**Good identification**							**07023002**
					SPO	CAT		
Propionibacterium granulosum		99.56	0.52	FRU 99	1	90		
Clostridium innocuum		0.44	0.02		99	1		
07023010	**Good identification**							**07023010**
Propionibacterium granulosum		100.0	0.47	FRU 99				
07023012	**Good identification**							**07023012**
Propionibacterium granulosum		100.0	0.37	FRU 99				
07023020	**Good identification**							**07023020**
Propionibacterium granulosum		100.0	0.52	FRU 99				
07023022	**Good identification**							**07023022**
Propionibacterium granulosum		100.0	0.41	FRU 99				
07023030	**Good identification**							**07023030**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.37	FRU 99				
07023200	**Good identification**							**07023200**
Propionibacterium granulosum		100.0	0.36	FRU 99				
07024000	**Low discrimination**							**07024000**
					COC	SPO	CAT	
Gemella morbillorum		78.07	0.35	TRE 1	99	1	1	
Propionibacterium granulosum		15.08	0.30	XYL 10	1	1	90	
Clostridium botulinum biovar A		5.45	0.13		1	99	1	
07030000	**Very good identification**							**07030000**
					COC	GRA	CAT	
Propionibacterium granulosum		98.02	0.98		1	99	90	
Fusobacterium mortiferum		1.21	0.69		1	1	5	
Gemella morbillorum		0.57	0.53		99	99	1	
07030001	**Very good identification**							**07030001**
					COC	GRA	CAT	
Propionibacterium granulosum		98.15	0.64	URE 1	1	99	90	
Fusobacterium mortiferum		1.22	0.35		1	1	5	
Gemella morbillorum		0.57	0.19		99	99	1	
07030002	**Very good identification**							**07030002**
					GRA	CAT		
Propionibacterium granulosum		99.36	0.88		99	90		
Fusobacterium mortiferum		0.55	0.53		1	5		
Atopobium minutum		0.06	0.31		99	1		
07030003	**Good identification**							**07030003**
					GRA	CAT		
Propionibacterium granulosum		99.45	0.54	URE 1	99	90		
Fusobacterium mortiferum		0.55	0.18		1	5		
07030010	**Very good identification**							**07030010**
					SPO	CAT		
Propionibacterium granulosum		99.47	0.83	MLZ 12	1	90		
Clostridium botulinum biovar B		0.26	0.26		99	1		
Clostridium difficile		0.14	0.22		99	1		
07030011	**Good identification**							**07030011**
Propionibacterium granulosum		100.0	0.49	MLZ 12				
07030012	**Very good identification**							**07030012**
					GRA	CAT		
Propionibacterium granulosum		99.96	0.73	MLZ 12	99	90		
Fusobacterium mortiferum		0.04	0.18		1	5		
07030013	**Good identification**							**07030013**
Propionibacterium granulosum		100.0	0.39	MLZ 12				
07030020	**Very good identification**							**07030020**
					COC	SPO	CAT	
Propionibacterium granulosum		99.35	0.88		1	1	90	
Gemella morbillorum		0.26	0.37		99	1	1	
Clostridium cadaveris		0.15	0.27		1	99	1	
07030021	**Good identification**							**07030021**
					COC	CAT		
Propionibacterium granulosum		99.74	0.54	URE 1	1	90		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		0.26	0.03		99	1		
07030022	**Very good identification**							**07030022**
					CAT			
Propionibacterium granulosum		99.85	0.78		90			
Atopobium minutum		0.08	0.23		1			
Clostridium perfringens		0.04	0.11		1			
07030023	**Good identification**							**07030023**
Propionibacterium granulosum		100.0	0.43	URE 1				
07030030	**Very good identification**							**07030030**
					COC	SPO	CAT	
Propionibacterium granulosum		99.48	0.73	MLZ 12	1	1	90	
Clostridium botulinum biovar B		0.51	0.21		1	99	1	
Gemella morbillorum		0.02	0.03		99	1	1	
07030031	**Good identification**							**07030031**
Propionibacterium granulosum		100.0	0.39	MLZ 12				
07030032	**Very good identification**							**07030032**
Propionibacterium granulosum		100.0	0.63	MLZ 12				
07030040	**Low discrimination**							**07030040**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		87.08	0.66	MLT 99	99	99	1	
Propionibacterium granulosum		12.46	0.64		99	1	90	
Fusobacterium mortiferum		0.15	0.35		1	1	5	
07030042	**Low discrimination**							**07030042**
					GRA	SPO	CAT	
Propionibacterium granulosum		77.32	0.54	SOR 1	99	1	90	
Clostridium botulinum biovar B		21.84	0.32		99	99	1	
Fusobacterium mortiferum		0.43	0.18		1	1	5	
07030050	**Very good identification**							**07030050**
					SPO	CAT		
Clostridium botulinum biovar B		96.16	0.61	MLT 99	99	1		
Propionibacterium granulosum		3.81	0.49		1	90		
Clostridium difficile		0.03	0.00		99	1		
07030060	**Good identification**							**07030060**
					CAT	SPO		
Clostridium botulinum biovar B		93.19	0.61	MLT 99	1	99		
Propionibacterium granulosum		6.77	0.54		90	1		
Clostridium perfringens		0.03	0.05		1	84		
07030062	**Low discrimination**							**07030062**
					SPO	CAT		
Propionibacterium granulosum		64.24	0.43	SOR 1	1	90		
Clostridium botulinum biovar B		35.76	0.26		99	1		
07030070	**Good identification**							**07030070**
					SPO	CAT		
Clostridium botulinum biovar B		98.03	0.55	MLT 99	99	1		
Propionibacterium granulosum		1.97	0.39		1	90		
07030100	**Low discrimination**							**07030100**
					GRA	CAT		
Fusobacterium mortiferum		73.73	0.75		1	5		
Propionibacterium granulosum		25.78	0.64		99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		0.23	0.23		99	1		
07030101	**Low discrimination**							**07030101**
					GRA	SPO	CAT	
Fusobacterium mortiferum		70.63	0.41	URE 1	1	1	5	
Propionibacterium granulosum		24.70	0.30		99	1	90	
Clostridium chauvoei		4.67	0.09		99	99	1	
07030102	**Low discrimination**							**07030102**
					GRA	CAT		
Fusobacterium mortiferum		55.85	0.59	BGL 10	1	5		
Propionibacterium granulosum		43.95	0.54		99	90		
Propionibacterium propionicum		0.11	0.14		99	1		
07030110	**Low discrimination**							**07030110**
					GRA	CAT		
Propionibacterium granulosum		82.52	0.49	LAC 1	99	90		
Fusobacterium mortiferum		17.48	0.41	MLZ 12	1	5		
07030112	**Acceptable identification**							**07030112**
					GRA	CAT		
Propionibacterium granulosum		91.40	0.39	LAC 1	99	90		
Fusobacterium mortiferum		8.60	0.25	MLZ 12	1	5		
07030120	**Low discrimination**							**07030120**
					GRA	CAT		
Propionibacterium granulosum		79.18	0.54	LAC 1	99	90		
Clostridium perfringens		11.09	0.32		99	1		
Fusobacterium mortiferum		9.15	0.41		1	5		
07030122	**Low discrimination**							**07030122**
					GRA	CAT		
Propionibacterium granulosum		76.05	0.43	LAC 1	99	90		
Clostridium perfringens		20.05	0.26		99	1		
Fusobacterium mortiferum		3.91	0.25		1	5		
07030130	**Good identification**							**07030130**
Propionibacterium granulosum		100.0	0.39	LAC 1				
07030200	**Very good identification**							**07030200**
					COC	GRA	CAT	
Propionibacterium granulosum		99.39	0.72	RHA 3	1	99	90	
Fusobacterium mortiferum		0.40	0.35		1	1	5	
Gemella morbillorum		0.19	0.19		99	99	1	
07030201	**Good identification**							**07030201**
Propionibacterium granulosum		100.0	0.38	RHA 3				
07030202	**Very good identification**							**07030202**
					GRA	CAT		
Propionibacterium granulosum		99.82	0.62	RHA 3	99	90		
Fusobacterium mortiferum		0.18	0.18		1	5		
07030210	**Very good identification**							**07030210**
Propionibacterium granulosum		100.0	0.58	RHA 3				
07030212	**Good identification**							**07030212**
Propionibacterium granulosum		100.0	0.47	RHA 3				
07030220	**Very good identification**							**07030220**
					COC	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		99.92	0.62	RHA 3	1	90		
Gemella morbillorum		0.08	0.03		99	1		
07030222	**Good identification**							**07030222**
Propionibacterium granulosum		100.0	0.52	RHA 3				
07030230	**Good identification**							**07030230**
Propionibacterium granulosum		100.0	0.47	RHA 3				
07030232	**Good identification**							**07030232**
Propionibacterium granulosum		100.0	0.37	RHA 3				
07030300	**Low discrimination**							**07030300**
					GRA	CAT		
Propionibacterium granulosum		51.71	0.38	LAC 1	99	90		
Fusobacterium mortiferum		48.29	0.41	RHA 3	1	5		
07030400	**Good identification**							**07030400**
					COC	GRA	CAT	
Propionibacterium granulosum		92.80	0.64	ARA 1	1	99	90	
Gemella morbillorum		5.93	0.37		99	99	1	
Fusobacterium mortiferum		1.15	0.35		1	1	5	
07030402	**Good identification**							**07030402**
					COC	GRA	CAT	
Propionibacterium granulosum		99.20	0.54	ARA 1	1	99	90	
Fusobacterium mortiferum		0.55	0.18		1	1	5	
Gemella morbillorum		0.26	0.03		99	99	1	
07030410	**Good identification**							**07030410**
					COC	CAT		
Propionibacterium granulosum		99.53	0.49	ARA 1	1	90		
Gemella morbillorum		0.47	0.03	MLZ 12	99	1		
07030412	**Good identification**							**07030412**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07030420	**Good identification**							**07030420**
					COC	CAT		
Propionibacterium granulosum		97.24	0.54	ARA 1	1	90		
Gemella morbillorum		2.76	0.20		99	1		
07030422	**Good identification**							**07030422**
Propionibacterium granulosum		100.0	0.43	ARA 1				
07030430	**Good identification**							**07030430**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07030500	**Low discrimination**							**07030500**
					GRA	CAT		
Fusobacterium mortiferum		69.05	0.41	ARA 1	1	5		
Propionibacterium granulosum		24.15	0.30		99	90		
Actinomyces meyeri		5.25	0.13		99	1		
07030600	**Good identification**							**07030600**
					COC	CAT		
Propionibacterium granulosum		97.96	0.38	RHA 3	1	90		
Gemella morbillorum		2.05	0.03	ARA 1	99	1		
07031000	**Very good identification**							**07031000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	CAT	
Propionibacterium granulosum		97.77	0.98		1	99	90	
Fusobacterium mortiferum		1.82	0.72		1	1	5	
Gemella morbillorum		0.31	0.48		99	99	1	
07031001	**Very good identification**							**07031001**
					GRA	CAT		
Propionibacterium granulosum		97.35	0.64	URE 1	99	90		
Fusobacterium mortiferum		1.81	0.38		1	5		
Actinomyces naeslundii		0.36	0.37		99	25		
07031002	**Very good identification**							**07031002**
					GRA	CAT		
Propionibacterium granulosum		98.88	0.88		99	90		
Fusobacterium mortiferum		0.82	0.56		1	5		
Lactobacillus jensenii		0.12	0.31		99	1		
07031003	**Good identification**							**07031003**
					GRA	CAT		
Propionibacterium granulosum		97.77	0.54	URE 1	99	90		
Actinomyces naeslundii		1.43	0.37		99	25		
Fusobacterium mortiferum		0.81	0.21		1	5		
07031010	**Very good identification**							**07031010**
					GRA	CAT		
Propionibacterium granulosum		99.73	0.83	MLZ 12	99	90		
Fusobacterium mortiferum		0.14	0.38		1	5		
Lactobacillus jensenii		0.11	0.26		99	1		
07031011	**Good identification**							**07031011**
Propionibacterium granulosum		100.0	0.49	MLZ 12				
07031012	**Very good identification**							**07031012**
					GRA	CAT		
Propionibacterium granulosum		99.64	0.73	MLZ 12	99	90		
Lactobacillus jensenii		0.30	0.23		99	1		
Fusobacterium mortiferum		0.06	0.21		1	5		
07031013	**Good identification**							**07031013**
Propionibacterium granulosum		100.0	0.39	MLZ 12				
07031020	**Very good identification**							**07031020**
					COC	CAT		
Propionibacterium granulosum		97.82	0.88		1	90		
Clostridium perfringens		1.87	0.51		1	1		
Gemella morbillorum		0.14	0.32		99	1		
07031021	**Good identification**							**07031021**
					CAT			
Propionibacterium granulosum		96.50	0.54	URE 1	90			
Clostridium perfringens		1.84	0.17		1			
Actinomyces naeslundii		1.41	0.37		25			
07031022	**Very good identification**							**07031022**
					CAT			
Propionibacterium granulosum		96.19	0.78		90			
Clostridium perfringens		3.46	0.45		1			
Lactobacillus fermentum		0.14	0.34		1			
07031023	**Acceptable identification**							**07031023**
					CAT			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		91.38	0.43	URE 1	90			
Actinomyces naeslundii		5.34	0.37		25			
Clostridium perfringens		3.28	0.11		1			
07031030	**Very good identification**							**07031030**
					CAT			
Propionibacterium granulosum		99.75	0.73	MLZ 12	90			
Clostridium perfringens		0.14	0.17		1			
Lactobacillus jensenii		0.11	0.16		1			
07031031	**Good identification**							**07031031**
Propionibacterium granulosum		100.0	0.39	MLZ 12				
07031032	**Very good identification**							**07031032**
					CAT			
Propionibacterium granulosum		99.44	0.63	MLZ 12	90			
Lactobacillus jensenii		0.30	0.13		1			
Clostridium perfringens		0.27	0.11		1			
07031040	**Good identification**							**07031040**
					GRA	SPO	CAT	
Propionibacterium granulosum		91.32	0.64	SOR 1	99	1	90	
Clostridium botulinum biovar B		6.45	0.32		99	99	1	
Fusobacterium mortiferum		1.70	0.38		1	1	5	
07031042	**Good identification**							**07031042**
					GRA	CAT		
Propionibacterium granulosum		98.39	0.54	SOR 1	99	90		
Fusobacterium mortiferum		0.81	0.21		1	5		
Clostridium perfringens		0.52	0.07		99	1		
07031050	**Low discrimination**							**07031050**
					SPO	CAT		
Propionibacterium granulosum		79.68	0.49	MLZ 12	1	90		
Clostridium botulinum biovar B		20.33	0.26	SOR 1	99	1		
07031052	**Good identification**							**07031052**
Propionibacterium granulosum		100.0	0.39	MLZ 12				
07031060	**Low discrimination**							**07031060**
					CAT	SPO		
Propionibacterium granulosum		66.70	0.54	SOR 1	90	1		
Clostridium perfringens		24.02	0.39		1	84		
Clostridium botulinum biovar B		9.28	0.26		1	99		
07031062	**Low discrimination**							**07031062**
					CAT			
Propionibacterium granulosum		59.60	0.43	SOR 1	90			
Clostridium perfringens		40.40	0.33		1			
07031070	**Low discrimination**							**07031070**
					CAT	SPO		
Propionibacterium granulosum		65.39	0.39	MLZ 12	90	1		
Clostridium botulinum biovar B		32.86	0.21	SOR 1	1	99		
Clostridium perfringens		1.74	0.05		1	84		
07031100	**Low discrimination**							**07031100**
					COC	GRA	CAT	
Fusobacterium mortiferum		74.92	0.78		1	1	5	
Propionibacterium granulosum		17.47	0.64		1	99	90	
Atopobium parvulum		4.51	0.46		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07031101	**Low discrimination**							**07031101**
					GRA	SPO		
Clostridium chauvoei		73.69	0.44	MLT 99	99	99		
Fusobacterium mortiferum		16.88	0.44	TRE 10	1	1		
Actinomyces naeslundii		4.05	0.45		99	1		
07031102	**Low discrimination**							**07031102**
					GRA	CAT		
Fusobacterium mortiferum		54.25	0.62	BGL 10	1	5		
Propionibacterium granulosum		28.46	0.54		99	90		
Clostridium perfringens		5.74	0.34		99	1		
07031110	**Low discrimination**							**07031110**
					COC	GRA	CAT	
Propionibacterium granulosum		74.36	0.49	LAC 1	1	99	90	
Fusobacterium mortiferum		23.63	0.44	MLZ 12	1	1	5	
Atopobium parvulum		1.42	0.12		99	99	1	
07031112	**Low discrimination**							**07031112**
					GRA	CAT		
Propionibacterium granulosum		86.49	0.39	LAC 1	99	90		
Fusobacterium mortiferum		12.21	0.28	MLZ 12	1	5		
Clostridium perfringens		1.29	0.00		99	1		
07031120	**Low discrimination**							**07031120**
					COC	CAT		
Clostridium perfringens		89.85	0.66	MLT 99	1	1		
Propionibacterium granulosum		6.48	0.54		1	90		
Atopobium parvulum		1.18	0.33		99	1		
07031121	**Low discrimination**							**07031121**
					SPO			
Clostridium chauvoei		56.81	0.35	MLT 99	99			
Clostridium perfringens		31.55	0.32	TRE 10	84			
Actinomyces naeslundii		9.36	0.45		1			
07031122	**Good identification**							**07031122**
					CAT			
Clostridium perfringens		93.83	0.60	MLT 99	1			
Propionibacterium granulosum		3.60	0.43		90			
Lactobacillus fermentum		0.87	0.38		1			
07031140	**Low discrimination**							**07031140**
					GRA	CAT		
Fusobacterium mortiferum		55.27	0.44	SOR 1	1	5		
Clostridium perfringens		26.05	0.28		99	1		
Propionibacterium granulosum		12.89	0.30		99	90		
07031160	**Good identification**							**07031160**
					CAT			
Clostridium perfringens		99.44	0.54	MLT 99	1			
Propionibacterium granulosum		0.38	0.20		90			
Actinomyces naeslundii		0.18	0.29		25			
07031162	**Good identification**							**07031162**
					CAT			
Clostridium perfringens		99.41	0.48	MLT 99	1			
Actinomyces naeslundii		0.39	0.29		25			
Propionibacterium granulosum		0.20	0.09		90			
07031200	**Very good identification**							**07031200**
					COC	GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		99.30	0.72	RHA 3	1	99	90	
Fusobacterium mortiferum		0.60	0.38		1	1	5	
Gemella morbillorum		0.10	0.14		99	99	1	
07031201	**Good identification**							**07031201**
Propionibacterium granulosum		100.0	0.38	RHA 3				
07031202	**Very good identification**							**07031202**
					GRA	CAT		
Propionibacterium granulosum		99.73	0.62	RHA 3	99	90		
Fusobacterium mortiferum		0.27	0.21		1	5		
07031210	**Very good identification**							**07031210**
Propionibacterium granulosum		100.0	0.58	RHA 3				
07031212	**Good identification**							**07031212**
Propionibacterium granulosum		100.0	0.47	RHA 3				
07031220	**Very good identification**							**07031220**
					CAT			
Propionibacterium granulosum		99.38	0.62	RHA 3	90			
Clostridium perfringens		0.62	0.17		1			
07031222	**Good identification**							**07031222**
					CAT			
Propionibacterium granulosum		98.84	0.52	RHA 3	90			
Clostridium perfringens		1.16	0.11		1			
07031230	**Good identification**							**07031230**
Propionibacterium granulosum		100.0	0.47	RHA 3				
07031232	**Good identification**							**07031232**
Propionibacterium granulosum		100.0	0.37	RHA 3				
07031240	**Good identification**							**07031240**
					SPO	CAT		
Propionibacterium granulosum		97.75	0.38	RHA 3	1	90		
Clostridium botulinum biovar B		2.26	-.03	SOR 1	99	1		
07031300	**Low discrimination**							**07031300**
					COC	GRA	CAT	
Fusobacterium mortiferum		55.59	0.44	RHA 1	1	1	5	
Propionibacterium granulosum		39.68	0.38		1	99	90	
Atopobium parvulum		3.34	0.12		99	99	1	
07031400	**Good identification**							**07031400**
					COC	CAT		
Propionibacterium granulosum		92.45	0.64	ARA 1	1	90		
Gemella morbillorum		3.18	0.32		99	1		
Lactobacillus fermentum		2.58	0.42		1	1		
07031402	**Low discrimination**							**07031402**
					GRA	CAT		
Propionibacterium granulosum		89.31	0.54	ARA 1	99	90		
Lactobacillus fermentum		9.95	0.42		99	1		
Fusobacterium mortiferum		0.74	0.21		1	5		
07031410	**Good identification**							**07031410**
Propionibacterium granulosum		100.0	0.49	ARA 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07031412	**Good identification**							**07031412**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07031420	**Acceptable identification**							**07031420**
					CAT			
Propionibacterium granulosum		93.33	0.54	ARA 1	90			
Lactobacillus fermentum		3.47	0.34		1			
Clostridium perfringens		1.78	0.17		1			
07031422	**Low discrimination**							**07031422**
					CAT			
Propionibacterium granulosum		84.43	0.43	ARA 1	90			
Lactobacillus fermentum		12.54	0.34		1			
Clostridium perfringens		3.03	0.11		1			
07031430	**Good identification**							**07031430**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07031500	**Low discrimination**							**07031500**
					GRA	CAT		
Lactobacillus fermentum		42.66	0.46	TRE 1	99	1		
Fusobacterium mortiferum		40.68	0.44		1	5		
Propionibacterium granulosum		9.48	0.30		99	90		
07031502	**Low discrimination**							**07031502**
					GRA	CAT		
Lactobacillus fermentum		84.24	0.46	TRE 1	99	1		
Fusobacterium mortiferum		8.92	0.28		1	5		
Propionibacterium granulosum		4.68	0.20		99	90		
07031600	**Good identification**							**07031600**
Propionibacterium granulosum		100.0	0.38	RHA 3				
07032000	**Excellent identification**							**07032000**
					SPO	CAT		
Propionibacterium granulosum		99.82	0.96		1	90		
Eubacterium limosum		0.08	0.29		1	1		
Clostridium difficile		0.05	0.27		99	1		
07032001	**Very good identification**							**07032001**
					CAT			
Propionibacterium granulosum		99.92	0.62	URE 1	90			
Eubacterium limosum		0.08	-.06		1			
07032002	**Very good identification**							**07032002**
					COC	GRA	CAT	
Propionibacterium granulosum		99.97	0.86		1	99	90	
Megasphaera elsdenii		0.01	0.12		99	1	1	
Fusobacterium mortiferum		0.01	0.18		1	1	5	
07032003	**Good identification**							**07032003**
Propionibacterium granulosum		100.0	0.52	URE 1				
07032010	**Very good identification**							**07032010**
					SPO	CAT		
Propionibacterium granulosum		98.32	0.81	MLZ 12	1	90		
Clostridium difficile		1.67	0.38		99	1		
Eubacterium limosum		0.01	-.06		1	1		
07032011	**Good identification**							**07032011**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	CAT		
Propionibacterium granulosum		98.33	0.47	MLZ 12	1	90		
Clostridium difficile		1.67	0.04	URE 1	99	1		
07032012	**Very good identification**							**07032012**
					SPO	CAT		
Propionibacterium granulosum		99.93	0.71	MLZ 12	1	90		
Clostridium difficile		0.07	0.04		99	1		
07032013	**Good identification**							**07032013**
Propionibacterium granulosum		100.0	0.37	MLZ 12				
07032020	**Very good identification**							**07032020**
					COC	CAT		
Propionibacterium granulosum		99.99	0.86		1	90		
Gemella morbillorum		0.00	0.03		99	1		
Eubacterium limosum		0.00	-.06		1	1		
07032021	**Good identification**							**07032021**
Propionibacterium granulosum		100.0	0.52	URE 1				
07032022	**Very good identification**							**07032022**
Propionibacterium granulosum		100.0	0.76					
07032023	**Good identification**							**07032023**
Propionibacterium granulosum		100.0	0.41	URE 1				
07032030	**Very good identification**							**07032030**
					SPO	CAT		
Propionibacterium granulosum		99.93	0.71	MLZ 12	1	90		
Clostridium difficile		0.07	0.04		99	1		
07032031	**Good identification**							**07032031**
Propionibacterium granulosum		100.0	0.37	MLZ 12				
07032032	**Very good identification**							**07032032**
Propionibacterium granulosum		100.0	0.61	MLZ 12				
07032040	**Low discrimination**							**07032040**
					COC	SPO	CAT	
Propionibacterium granulosum		89.86	0.62	SOR 1	1	1	90	
Clostridium botulinum biovar B		8.41	0.32		1	99	1	
Megasphaera elsdenii		1.44	0.26		99	1	1	
07032042	**Good identification**							**07032042**
					COC	SPO	CAT	
Propionibacterium granulosum		98.61	0.52	SOR 1	1	1	90	
Megasphaera elsdenii		0.70	0.09		99	1	1	
Clostridium botulinum biovar B		0.37	-.03		1	99	1	
07032050	**Low discrimination**							**07032050**
					SPO	CAT		
Propionibacterium granulosum		70.09	0.47	MLZ 12	1	90		
Clostridium botulinum biovar B		23.70	0.26	SOR 1	99	1		
Clostridium difficile		6.21	0.17		99	1		
07032052	**Good identification**							**07032052**
Propionibacterium granulosum		100.0	0.37	MLZ 12				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07032060	**Low discrimination**						**07032060**
					SPO	CAT	
Propionibacterium granulosum		84.43	0.52	SOR 1	1	90	
Clostridium botulinum biovar B		15.57	0.26		99	1	
07032062	**Good identification**						**07032062**
Propionibacterium granulosum		100.0	0.41	SOR 1			
07032070	**Low discrimination**						**07032070**
					SPO	CAT	
Propionibacterium granulosum		60.02	0.37	MLZ 12	1	90	
Clostridium botulinum biovar B		39.98	0.21	SOR 1	99	1	
07032100	**Good identification**						**07032100**
					GRA	CAT	
Propionibacterium granulosum		91.45	0.62	LAC 1	99	90	
Propionibacterium propionicum		4.90	0.45		99	1	
Fusobacterium mortiferum		3.50	0.41		1	5	
07032102	**Low discrimination**						**07032102**
					GRA	CAT	
Propionibacterium granulosum		74.53	0.52	LAC 1	99	90	
Propionibacterium propionicum		23.98	0.48		99	1	
Fusobacterium mortiferum		1.27	0.25		1	5	
07032110	**Good identification**						**07032110**
					SPO	CAT	
Propionibacterium granulosum		98.33	0.47	LAC 1	1	90	
Clostridium difficile		1.67	0.04	MLZ 12	99	1	
07032112	**Good identification**						**07032112**
					CAT		
Propionibacterium granulosum		97.67	0.37	LAC 1	90		
Propionibacterium propionicum		2.33	0.14	MLZ 12	1		
07032120	**Good identification**						**07032120**
Propionibacterium granulosum		99.70	0.52	LAC 1			
Actinomyces naeslundii		0.30	0.23				
07032122	**Good identification**						**07032122**
					CAT		
Propionibacterium granulosum		97.57	0.41	LAC 1	90		
Propionibacterium propionicum		1.27	0.14		1		
Actinomyces naeslundii		1.16	0.23		25		
07032130	**Good identification**						**07032130**
Propionibacterium granulosum		100.0	0.37	LAC 1			
07032140	**Low discrimination**						**07032140**
					SPO	CAT	
Propionibacterium propionicum		82.92	0.45	NIT 86	1	1	
Propionibacterium granulosum		15.62	0.28	MLT 99	1	90	
Clostridium botulinum biovar B		1.46	-.03		99	1	
07032142	**Good identification**						**07032142**
					CAT		
Propionibacterium propionicum		96.96	0.48	NIT 86	1		
Propionibacterium granulosum		3.04	0.18	MLT 99	90		
07032200	**Very good identification**						**07032200**
					CAT		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		99.97	0.70	RHA 3	90		
Eubacterium limosum		0.03	-.06		1		
07032201	**Good identification**						**07032201**
Propionibacterium granulosum		100.0	0.36	RHA 3			
07032202	**Very good identification**						**07032202**
Propionibacterium granulosum		100.0	0.60	RHA 3			
07032210	**Good identification**						**07032210**
					SPO	CAT	
Propionibacterium granulosum		99.45	0.56	RHA 3	1	90	
Clostridium difficile		0.55	0.04	MLZ 12	99	1	
07032212	**Good identification**						**07032212**
Propionibacterium granulosum		100.0	0.45	RHA 3			
07032220	**Very good identification**						**07032220**
Propionibacterium granulosum		100.0	0.60	RHA 3			
07032222	**Good identification**						**07032222**
Propionibacterium granulosum		100.0	0.50	RHA 3			
07032230	**Good identification**						**07032230**
Propionibacterium granulosum		100.0	0.45	RHA 3			
07032240	**Good identification**						**07032240**
					SPO	CAT	
Propionibacterium granulosum		97.03	0.36	RHA 3	1	90	
Clostridium botulinum biovar B		2.97	-.03	SOR 1	99	1	
07032300	**Good identification**						**07032300**
Propionibacterium granulosum		100.0	0.36	LAC 1			
07032400	**Very good identification**						**07032400**
					COC	CAT	
Propionibacterium granulosum		99.84	0.62	ARA 1	1	90	
Gemella morbillorum		0.09	0.03		99	1	
Eubacterium limosum		0.08	-.06		1	1	
07032402	**Good identification**						**07032402**
Propionibacterium granulosum		100.0	0.52	ARA 1			
07032410	**Good identification**						**07032410**
					SPO	CAT	
Propionibacterium granulosum		98.33	0.47	ARA 1	1	90	
Clostridium difficile		1.67	0.04	MLZ 12	99	1	
07032412	**Good identification**						**07032412**
Propionibacterium granulosum		100.0	0.37	ARA 1			
07032420	**Good identification**						**07032420**
Propionibacterium granulosum		100.0	0.52	ARA 1			
07032422	**Good identification**						**07032422**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.41	ARA 1				
07032430	**Good identification**							**07032430**
Propionibacterium granulosum		100.0	0.37	ARA 1				
07032600	**Good identification**							**07032600**
Propionibacterium granulosum		100.0	0.36	RHA 3				
07033000	**Excellent identification**							**07033000**
					GRA	SPO	CAT	
Propionibacterium granulosum		99.94	0.96		99	1	90	
Fusobacterium mortiferum		0.03	0.38		1	1	5	
Clostridium innocuum		0.02	0.24		99	99	1	
07033001	**Very good identification**							**07033001**
Propionibacterium granulosum		99.07	0.62	URE 1				
Actinomyces naeslundii		0.93	0.42					
07033002	**Very good identification**							**07033002**
					SPO	CAT		
Propionibacterium granulosum		99.88	0.86		1	90		
Clostridium innocuum		0.08	0.24		99	1		
Actinomyces naeslundii		0.03	0.39		1	25		
07033003	**Good identification**							**07033003**
Propionibacterium granulosum		96.38	0.52	URE 1				
Actinomyces naeslundii		3.62	0.42					
07033010	**Very good identification**							**07033010**
					SPO	CAT		
Propionibacterium granulosum		99.98	0.81	MLZ 12	1	90		
Clostridium difficile		0.02	0.04		99	1		
Clostridium innocuum		0.01	0.00		99	1		
07033011	**Good identification**							**07033011**
Propionibacterium granulosum		100.0	0.47	MLZ 12				
07033012	**Very good identification**							**07033012**
					SPO	CAT		
Propionibacterium granulosum		99.97	0.71	MLZ 12	1	90		
Clostridium innocuum		0.03	0.00		99	1		
07033013	**Good identification**							**07033013**
Propionibacterium granulosum		100.0	0.37	MLZ 12				
07033020	**Very good identification**							**07033020**
					CAT			
Propionibacterium granulosum		99.95	0.86		90			
Clostridium perfringens		0.03	0.17		1			
Actinomyces naeslundii		0.03	0.39		25			
07033021	**Good identification**							**07033021**
Propionibacterium granulosum		96.38	0.52	URE 1				
Actinomyces naeslundii		3.62	0.42					
07033022	**Very good identification**							**07033022**
					CAT			
Propionibacterium granulosum		99.85	0.76		90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.10	0.39		25		
Clostridium perfringens		0.05	0.11		1		
07033023	**Low discrimination**						**07033023**
Propionibacterium granulosum		86.93	0.41	URE 1			
Actinomyces naeslundii		13.07	0.42				
07033030	**Very good identification**						**07033030**
Propionibacterium granulosum		100.0	0.71	MLZ 12			
07033031	**Good identification**						**07033031**
Propionibacterium granulosum		100.0	0.37	MLZ 12			
07033032	**Very good identification**						**07033032**
Propionibacterium granulosum		100.0	0.61	MLZ 12			
07033040	**Very good identification**						**07033040**
					SPO	CAT	
Propionibacterium granulosum		99.80	0.62	SOR 1	1	90	
Actinomyces naeslundii		0.11	0.26		1	25	
Clostridium botulinum biovar B		0.09	-.03		99	1	
07033042	**Good identification**						**07033042**
					CAT		
Propionibacterium granulosum		99.24	0.52	SOR 1	90		
Actinomyces naeslundii		0.44	0.26		25		
Propionibacterium propionicum		0.32	0.14		1		
07033050	**Good identification**						**07033050**
Propionibacterium granulosum		100.0	0.47	MLZ 12			
07033052	**Good identification**						**07033052**
Propionibacterium granulosum		100.0	0.37	MLZ 12			
07033060	**Good identification**						**07033060**
					CAT		
Propionibacterium granulosum		99.08	0.52	SOR 1	90		
Clostridium perfringens		0.48	0.05		1		
Actinomyces naeslundii		0.44	0.26		25		
07033062	**Good identification**						**07033062**
Propionibacterium granulosum		98.26	0.41	SOR 1			
Actinomyces naeslundii		1.74	0.26				
07033070	**Good identification**						**07033070**
Propionibacterium granulosum		100.0	0.37	MLZ 12			
07033100	**Low discrimination**						**07033100**
					GRA	CAT	
Propionibacterium granulosum		88.20	0.62	LAC 1	99	90	
Fusobacterium mortiferum		5.07	0.44		1	5	
Propionibacterium propionicum		4.73	0.45		99	1	
07033101	**Low discrimination**						**07033101**
					CAT		
Actinomyces naeslundii		58.98	0.50	MLT 94	25		
Propionibacterium granulosum		22.29	0.28	RAF 92	90		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		13.14	0.29		1		
07033102	**Low discrimination**						**07033102**
					CAT		
Propionibacterium granulosum		70.36	0.52	LAC 1	90		
Propionibacterium propionicum		22.64	0.48		1		
Actinomyces naeslundii		5.02	0.47		25		
07033103	**Low discrimination**						**07033103**
					CAT		
Actinomyces naeslundii		69.99	0.50	MLT 94	25		
Propionibacterium propionicum		23.40	0.32	RAF 92	1		
Propionibacterium granulosum		6.61	0.18		90		
07033110	**Good identification**						**07033110**
Propionibacterium granulosum		99.32	0.47	LAC 1			
Actinomyces naeslundii		0.68	0.25	MLZ 12			
07033112	**Good identification**						**07033112**
					CAT		
Propionibacterium granulosum		95.12	0.37	LAC 1	90		
Actinomyces naeslundii		2.62	0.25	MLZ 12	25		
Propionibacterium propionicum		2.27	0.14		1		
07033120	**Low discrimination**						**07033120**
					CAT		
Propionibacterium granulosum		79.56	0.52	LAC 1	90		
Clostridium perfringens		14.77	0.32		1		
Actinomyces naeslundii		5.67	0.47		25		
07033121	**Low discrimination**						**07033121**
					SPO	CAT	
Actinomyces naeslundii		88.81	0.50	MLT 94	1	25	
Propionibacterium granulosum		8.39	0.18	RAF 92	1	90	
Clostridium chauvoei		2.80	0.01		99	1	
07033122	**Low discrimination**						**07033122**
					CAT		
Propionibacterium granulosum		60.70	0.41	LAC 1	90		
Clostridium perfringens		21.21	0.26		1		
Actinomyces naeslundii		17.31	0.47		25		
07033123	**Good identification**						**07033123**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07033130	**Good identification**						**07033130**
Propionibacterium granulosum		97.32	0.37	LAC 1			
Actinomyces naeslundii		2.68	0.25	MLZ 12			
07033140	**Low discrimination**						**07033140**
					CAT		
Propionibacterium propionicum		80.19	0.45	NIT 86	1		
Propionibacterium granulosum		15.11	0.28	MLT 99	90		
Actinomyces naeslundii		4.70	0.34		25		
07033142	**Acceptable identification**						**07033142**
					CAT		
Propionibacterium propionicum		93.42	0.48	NIT 86	1		
Actinomyces naeslundii		3.65	0.34	MLT 99	25		
Propionibacterium granulosum		2.93	0.18		90		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07033161	**Good identification**						**07033161**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07033163	**Good identification**						**07033163**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07033200	**Very good identification**						**07033200**
Propionibacterium granulosum		100.0	0.70	RHA 3			
07033201	**Good identification**						**07033201**
Propionibacterium granulosum		100.0	0.36	RHA 3			
07033202	**Very good identification**						**07033202**
Propionibacterium granulosum		100.0	0.60	RHA 3			
07033210	**Good identification**						**07033210**
Propionibacterium granulosum		100.0	0.56	RHA 3			
07033212	**Good identification**						**07033212**
Propionibacterium granulosum		100.0	0.45	RHA 3			
07033220	**Very good identification**						**07033220**
Propionibacterium granulosum		100.0	0.60	RHA 3			
07033222	**Good identification**						**07033222**
Propionibacterium granulosum		100.0	0.50	RHA 3			
07033230	**Good identification**						**07033230**
Propionibacterium granulosum		100.0	0.45	RHA 3			
07033240	**Good identification**						**07033240**
Propionibacterium granulosum		100.0	0.36	RHA 3			
07033300	**Good identification**						**07033300**
Propionibacterium granulosum		100.0	0.36	LAC 1			
07033400	**Very good identification**						**07033400**
					SPO	CAT	
Propionibacterium granulosum		99.63	0.62	ARA 1	1	90	
Clostridium innocuum		0.37	0.11		99	1	
07033402	**Good identification**						**07033402**
					SPO	CAT	
Propionibacterium granulosum		98.55	0.52	ARA 1	1	90	
Clostridium innocuum		1.45	0.11		99	1	
07033410	**Good identification**						**07033410**
Propionibacterium granulosum		100.0	0.47	ARA 1			
07033412	**Good identification**						**07033412**
Propionibacterium granulosum		100.0	0.37	ARA 1			
07033420	**Good identification**						**07033420**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.52	ARA 1				
07033422	**Good identification**							**07033422**
Propionibacterium granulosum		100.0	0.41	ARA 1				
07033430	**Good identification**							**07033430**
Propionibacterium granulosum		100.0	0.37	ARA 1				
07033600	**Good identification**							**07033600**
Propionibacterium granulosum		100.0	0.36	RHA 3				
07034000	**Low discrimination**							**07034000**
					COC	GRA	CAT	
Propionibacterium granulosum		86.00	0.64	XYL 1	1	99	90	
Fusobacterium mortiferum		7.93	0.50		1	1	5	
Gemella morbillorum		5.50	0.37		99	99	1	
07034002	**Good identification**							**07034002**
					COC	GRA	CAT	
Propionibacterium granulosum		95.82	0.54	XYL 1	1	99	90	
Fusobacterium mortiferum		3.93	0.33		1	1	5	
Gemella morbillorum		0.25	0.03		99	99	1	
07034010	**Good identification**							**07034010**
					GRA	SPO	CAT	
Propionibacterium granulosum		98.14	0.49	XYL 1	99	1	90	
Clostridium difficile		0.73	0.00	MLZ 12	99	99	1	
Fusobacterium mortiferum		0.67	0.15		1	1	5	
07034012	**Good identification**							**07034012**
Propionibacterium granulosum		100.0	0.39	XYL 1				
07034020	**Good identification**							**07034020**
					COC	GRA	CAT	
Propionibacterium granulosum		96.67	0.54	XYL 1	1	99	90	
Gemella morbillorum		2.75	0.20		99	99	1	
Fusobacterium mortiferum		0.36	0.15		1	1	5	
07034022	**Good identification**							**07034022**
Propionibacterium granulosum		100.0	0.43	XYL 1				
07034030	**Good identification**							**07034030**
Propionibacterium granulosum		100.0	0.39	XYL 1				
07034100	**Low discrimination**							**07034100**
					GRA	CAT		
Fusobacterium mortiferum		86.83	0.56	XYL 7	1	5		
Actinomyces meyeri		6.88	0.28		99	1		
Propionibacterium granulosum		4.08	0.30		99	90		
07034102	**Low discrimination**							**07034102**
					COC	GRA	CAT	
Fusobacterium mortiferum		85.96	0.40	XYL 7	1	1	5	
Propionibacterium granulosum		9.08	0.20	BGL 10	1	99	90	
Atopobium parvulum		3.07	0.03		99	99	1	
07034200	**Good identification**							**07034200**
					COC	GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		95.15	0.38	XYL 1	1	99	90	
Fusobacterium mortiferum		2.87	0.15	RHA 3	1	1	5	
Gemella morbillorum		1.99	0.03		99	99	1	
07035000	**Low discrimination**							**07035000**
					COC	GRA	CAT	
Propionibacterium granulosum		83.31	0.64	XYL 1	1	99	90	
Fusobacterium mortiferum		11.53	0.53		1	1	5	
Gemella morbillorum		2.87	0.32		99	99	1	
07035002	**Low discrimination**							**07035002**
					GRA	CAT		
Propionibacterium granulosum		88.47	0.54	XYL 1	99	90		
Lactobacillus fermentum		5.79	0.38		99	1		
Fusobacterium mortiferum		5.44	0.36		1	5		
07035010	**Good identification**							**07035010**
					GRA	CAT		
Propionibacterium granulosum		98.99	0.49	XYL 1	99	90		
Fusobacterium mortiferum		1.01	0.19	MLZ 12	1	5		
07035012	**Good identification**							**07035012**
Propionibacterium granulosum		100.0	0.39	XYL 1				
07035020	**Acceptable identification**							**07035020**
					CAT			
Propionibacterium granulosum		93.57	0.54	XYL 1	90			
Lactobacillus fermentum		2.04	0.30		1			
Clostridium perfringens		1.79	0.17		1			
07035022	**Low discrimination**							**07035022**
					CAT			
Propionibacterium granulosum		89.03	0.43	XYL 1	90			
Lactobacillus fermentum		7.77	0.30		1			
Clostridium perfringens		3.20	0.11		1			
07035030	**Good identification**							**07035030**
Propionibacterium granulosum		100.0	0.39	XYL 1				
07035100	**Low discrimination**							**07035100**
					COC	GRA		
Atopobium parvulum		55.37	0.50	MLT 99	99	99		
Fusobacterium mortiferum		37.30	0.59		1	1		
Lactobacillus fermentum		3.08	0.42		1	99		
07035102	**Low discrimination**							**07035102**
					COC	GRA		
Atopobium parvulum		56.02	0.38	MLT 99	99	99		
Fusobacterium mortiferum		23.76	0.43		1	1		
Lactobacillus fermentum		17.68	0.42		1	99		
07035120	**Low discrimination**							**07035120**
					COC			
Atopobium parvulum		61.40	0.38	MLT 99	99			
Clostridium perfringens		25.41	0.32		1			
Lactobacillus fermentum		6.46	0.34		1			
07035122	**Low discrimination**							**07035122**
					COC	GRA		
Parabacteroides distasonis		47.33	0.41	RAF 99	1	1		
Clostridium perfringens		21.20	0.26		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		19.21	0.25		99	99		
07035132	**Good identification**							**07035132**
Parabacteroides distasonis		100.0	0.37	RAF 99				
07035200	**Good identification**							**07035200**
					GRA	CAT		
Propionibacterium granulosum		95.68	0.38	XYL 1	99	90		
Fusobacterium mortiferum		4.32	0.19	RHA 3	1	5		
07035322	**Good identification**							**07035322**
Parabacteroides distasonis		100.0	0.50	RAF 99				
07035332	**Good identification**							**07035332**
Parabacteroides distasonis		100.0	0.45	RAF 99				
07035400	**Low discrimination**							**07035400**
					COC	CAT		
Lactobacillus fermentum		49.15	0.38	TRE 1	1	1		
Propionibacterium granulosum		30.35	0.30		1	90		
Gemella morbillorum		11.49	0.16		99	1		
07035402	**Low discrimination**							**07035402**
					CAT			
Lactobacillus fermentum		86.63	0.38	TRE 1	1			
Propionibacterium granulosum		13.37	0.20		90			
07035500	**Low discrimination**							**07035500**
					COC			
Lactobacillus fermentum		61.85	0.42	TRE 1	1			
Actinomyces meyeri		14.83	0.21		1			
Atopobium parvulum		11.22	0.16		99			
07035502	**Low discrimination**							**07035502**
Lactobacillus fermentum		86.47	0.42	TRE 1				
Bifidobacterium longum subsp. longum		6.34	0.15					
Bifidobacterium adolescentis		4.42	0.14					
07035522	**Low discrimination**							**07035522**
					GRA			
Parabacteroides distasonis		57.41	0.38	RAF 99	1			
Lactobacillus fermentum		22.66	0.34		99			
Bifidobacterium longum subsp. longum		19.94	0.26		99			
07035722	**Good identification**							**07035722**
Parabacteroides distasonis		100.0	0.46	RAF 99				
07035732	**Good identification**							**07035732**
Parabacteroides distasonis		95.68	0.41	RAF 99				
Clostridium clostridioforme		4.32	0.13					
07036000	**Very good identification**							**07036000**
					GRA	SPO	CAT	
Propionibacterium granulosum		99.47	0.62	XYL 1	99	1	90	
Clostridium difficile		0.25	0.05		99	99	1	
Fusobacterium mortiferum		0.12	0.15		1	1	5	
07036002	**Good identification**							**07036002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.52	XYL 1				
07036010	**Acceptable identification**							**07036010**
					SPO	CAT		
Propionibacterium granulosum		91.86	0.47	XYL 1	1	90		
Clostridium difficile		8.14	0.17	MLZ 12	99	1		
07036012	**Good identification**							**07036012**
Propionibacterium granulosum		100.0	0.37	XYL 1				
07036020	**Good identification**							**07036020**
Propionibacterium granulosum		100.0	0.52	XYL 1				
07036022	**Good identification**							**07036022**
Propionibacterium granulosum		100.0	0.41	XYL 1				
07036030	**Good identification**							**07036030**
Propionibacterium granulosum		100.0	0.37	XYL 1				
07036200	**Good identification**							**07036200**
Propionibacterium granulosum		100.0	0.36	XYL 1				
07037000	**Very good identification**							**07037000**
					GRA	SPO	CAT	
Propionibacterium granulosum		99.71	0.62	XYL 1	99	1	90	
Fusobacterium mortiferum		0.19	0.19		1	1	5	
Clostridium innocuum		0.11	0.02		99	99	1	
07037002	**Good identification**							**07037002**
					SPO	CAT		
Propionibacterium granulosum		99.56	0.52	XYL 1	1	90		
Clostridium innocuum		0.44	0.02		99	1		
07037010	**Good identification**							**07037010**
Propionibacterium granulosum		100.0	0.47	XYL 1				
07037012	**Good identification**							**07037012**
Propionibacterium granulosum		100.0	0.37	XYL 1				
07037020	**Good identification**							**07037020**
Propionibacterium granulosum		100.0	0.52	XYL 1				
07037022	**Good identification**							**07037022**
Propionibacterium granulosum		100.0	0.41	XYL 1				
07037030	**Good identification**							**07037030**
Propionibacterium granulosum		100.0	0.37	XYL 1				
07037200	**Good identification**							**07037200**
Propionibacterium granulosum		100.0	0.36	XYL 1				
07040000	**Low discrimination**							**07040000**
					COC	SPO		
Gemella morbillorum		87.68	0.52	CEL 1	99	1		
Atopobium minutum		4.72	0.35		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		2.72	0.23		1	99		
07040020	**Low discrimination**							**07040020**
					COC	SPO		
Gemella morbillorum		76.19	0.35	CEL 1	99	1		
Atopobium minutum		12.29	0.26	NAG 10	1	1		
Clostridium cadaveris		10.29	0.15		1	99		
07040100	**Low discrimination**							**07040100**
					COC	GRA		
Fusobacterium mortiferum		87.15	0.42	FRU 99	1	1		
Gemella morbillorum		12.85	0.17		99	99		
07040400	**Good identification**							**07040400**
Gemella morbillorum		100.0	0.35	CEL 1				
07041000	**Low discrimination**							**07041000**
					COC	GRA		
Gemella morbillorum		86.34	0.47	CEL 1	99	99		
Fusobacterium mortiferum		7.06	0.39		1	1		
Atopobium minutum		2.87	0.26		1	99		
07041100	**Low discrimination**							**07041100**
					COC	GRA		
Fusobacterium mortiferum		87.75	0.45	FRU 99	1	1		
Lactobacillus catenaformis		6.35	0.15		1	99		
Gemella morbillorum		4.65	0.13		99	99		
07043000	**Good identification**							**07043000**
					COC	SPO	CAT	
Clostridium innocuum		97.86	0.44	SAL 99	1	99	1	
Gemella morbillorum		1.08	0.13	FRU 95	99	1	1	
Propionibacterium granulosum		1.07	0.20		1	1	90	
07043002	**Good identification**							**07043002**
					SPO	CAT		
Clostridium innocuum		99.73	0.44	SAL 99	99	1		
Propionibacterium granulosum		0.27	0.09	FRU 95	1	90		
07044000	**Good identification**							**07044000**
					COC	GRA		
Gemella morbillorum		98.05	0.35	CEL 1	99	99		
Fusobacterium mortiferum		1.95	0.16	XYL 10	1	1		
07050000	**Low discrimination**							**07050000**
					COC	GRA	CAT	
Fusobacterium mortiferum		57.24	0.70		1	1	5	
Gemella morbillorum		24.08	0.53		99	99	1	
Propionibacterium granulosum		13.95	0.56		1	99	90	
07050001	**Low discrimination**							**07050001**
					COC	GRA	CAT	
Fusobacterium mortiferum		60.08	0.36	URE 1	1	1	5	
Gemella morbillorum		25.27	0.19		99	99	1	
Propionibacterium granulosum		14.65	0.22		1	99	90	
07050002	**Low discrimination**							**07050002**
					GRA	CAT		
Fusobacterium mortiferum		57.04	0.53	BGL 10	1	5		
Propionibacterium granulosum		31.28	0.46		99	90		
Atopobium minutum		5.85	0.31		99	1		
07050010	**Low discrimination**							**07050010**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Propionibacterium granulosum		57.17	0.41	CEL 1	99	1	90	
Fusobacterium mortiferum		17.38	0.36	MLZ 12	1	1	5	
Clostridium difficile		16.93	0.19		99	99	1	
07050020	**Low discrimination**							**07050020**
					COC	GRA	CAT	
Propionibacterium granulosum		46.57	0.46	CEL 1	1	99	90	
Gemella morbillorum		35.72	0.37		99	99	1	
Fusobacterium mortiferum		7.72	0.36		1	1	5	
07050022	**Low discrimination**							**07050022**
					SPO	CAT		
Propionibacterium granulosum		64.77	0.35	CEL 1	1	90		
Atopobium minutum		16.14	0.23		1	1		
Clostridium barati		10.50	0.10		99	1		
07050100	**Very good identification**							**07050100**
					COC	GRA		
Fusobacterium mortiferum		99.30	0.76		1	1		
Actinomyces meyeri		0.27	0.23		1	99		
Gemella morbillorum		0.18	0.19		99	99		
07050101	**Good identification**							**07050101**
Fusobacterium mortiferum		100.0	0.42	URE 1				
07050102	**Very good identification**							**07050102**
					GRA	SPO		
Fusobacterium mortiferum		97.43	0.60	BGL 10	1	1		
Lactobacillus catenaformis		0.96	0.15		99	1		
Clostridium barati		0.93	0.10		99	99		
07050110	**Good identification**							**07050110**
Fusobacterium mortiferum		100.0	0.42	MLZ 1				
07050120	**Low discrimination**							**07050120**
					GRA	SPO		
Fusobacterium mortiferum		78.53	0.42	NAG 1	1	1		
Clostridium barati		8.23	0.10		99	99		
Actinomyces meyeri		2.92	0.08		99	1		
07050140	**Good identification**							**07050140**
Fusobacterium mortiferum		100.0	0.42	SOR 1				
07050200	**Low discrimination**							**07050200**
					COC	GRA	CAT	
Fusobacterium mortiferum		46.15	0.36	RHA 1	1	1	5	
Propionibacterium granulosum		34.44	0.30		1	99	90	
Gemella morbillorum		19.41	0.19		99	99	1	
07050300	**Good identification**							**07050300**
Fusobacterium mortiferum		100.0	0.42	RHA 1				
07050400	**Low discrimination**							**07050400**
					COC	GRA	CAT	
Gemella morbillorum		78.28	0.37	CEL 1	99	99	1	
Fusobacterium mortiferum		16.92	0.36	ARA 10	1	1	5	
Propionibacterium granulosum		4.12	0.22		1	99	90	
07050500	**Acceptable identification**							**07050500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Fusobacterium mortiferum		91.89	0.42	ARA 1	1	1		
Actinomyces meyeri		6.27	0.13		1	99		
Gemella morbillorum		1.84	0.03		99	99		
07051000	**Low discrimination**							**07051000**
					COC	GRA	CAT	
Fusobacterium mortiferum		62.05	0.73		1	1	5	
Propionibacterium granulosum		10.08	0.56		1	99	90	
Gemella morbillorum		9.37	0.48		99	99	1	
07051001	**Low discrimination**							**07051001**
					COC	GRA	CAT	
Fusobacterium mortiferum		59.32	0.39	URE 1	1	1	5	
Propionibacterium granulosum		9.64	0.22		1	99	90	
Gemella morbillorum		8.96	0.14		99	99	1	
07051002	**Low discrimination**							**07051002**
					GRA			
Lactobacillus catenaformis		27.34	0.47	MLT 90	99			
Fusobacterium mortiferum		27.00	0.56	SAL 99	1			
Lactobacillus jensenii		22.58	0.45		99			
07051010	**Low discrimination**							**07051010**
					GRA	CAT		
Lactobacillus jensenii		56.46	0.40	MLT 95	99	1		
Propionibacterium granulosum		26.93	0.41	SAL 95	99	90		
Fusobacterium mortiferum		12.28	0.39		1	5		
07051012	**Low discrimination**							**07051012**
					CAT			
Lactobacillus jensenii		78.06	0.37	MLT 95	1			
Propionibacterium granulosum		13.96	0.31	SAL 95	90			
Lactobacillus catenaformis		2.86	0.12		1			
07051020	**Low discrimination**							**07051020**
					CAT			
Propionibacterium granulosum		26.47	0.46	CEL 1	90			
Lactobacillus jensenii		22.70	0.38		1			
Lactobacillus catenaformis		18.32	0.36		1			
07051040	**Low discrimination**							**07051040**
					COC	GRA	CAT	
Fusobacterium mortiferum		62.69	0.39	SOR 1	1	1	5	
Propionibacterium granulosum		10.19	0.22		1	99	90	
Gemella morbillorum		9.47	0.14		99	99	1	
07051100	**Good identification**							**07051100**
					GRA			
Fusobacterium mortiferum		91.35	0.79		1			
Lactobacillus catenaformis		6.61	0.50		99			
Lactobacillus fermentum		0.87	0.46		99			
07051101	**Low discrimination**							**07051101**
					GRA	SPO		
Fusobacterium mortiferum		63.51	0.45	URE 1	1	1		
Clostridium chauvoei		22.58	0.26		99	99		
Actinomyces naeslundii		9.32	0.42		99	1		
07051102	**Low discrimination**							**07051102**
					GRA			
Fusobacterium mortiferum		48.89	0.63	BGL 10	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus catenaformis		31.82	0.50		99			
Lactobacillus acidophilus		8.10	0.36		99			
07051103	**Low discrimination**							**07051103**
					GRA			
Actinomyces naeslundii		34.55	0.42	MLT 94	99			
Fusobacterium mortiferum		26.16	0.29	RAF 92	1			
Lactobacillus catenaformis		17.02	0.15		99			
07051110	**Acceptable identification**							**07051110**
					GRA			
Fusobacterium mortiferum		91.56	0.45	MLZ 1	1			
Lactobacillus catenaformis		6.62	0.15		99			
Lactobacillus jensenii		1.82	0.06		99			
07051121	**Low discrimination**							**07051121**
					GRA	SPO		
Actinomyces naeslundii		43.73	0.42	MLT 94	99	1		
Clostridium chauvoei		35.32	0.18	RAF 92	99	99		
Prevotella melaninogenica		6.73	-.01		1	1		
07051122	**Low discrimination**							**07051122**
					SPO			
Clostridium barati		38.62	0.40	ESL 87	99			
Lactobacillus acidophilus		17.69	0.36	SAL 99	1			
Lactobacillus catenaformis		17.37	0.39		1			
07051123	**Low discrimination**							**07051123**
					SPO			
Actinomyces naeslundii		60.05	0.42	MLT 94	1			
Clostridium barati		16.45	0.06	RAF 92	99			
Clostridium chauvoei		8.56	0.05		99			
07051140	**Low discrimination**							**07051140**
					GRA	SPO		
Fusobacterium mortiferum		88.65	0.45	SOR 1	1	1		
Lactobacillus catenaformis		6.41	0.15		99	1		
Clostridium barati		3.41	0.05		99	99		
07051200	**Low discrimination**							**07051200**
					COC	GRA	CAT	
Fusobacterium mortiferum		51.81	0.39	RHA 1	1	1	5	
Propionibacterium granulosum		25.78	0.30		1	99	90	
Gemella morbillorum		7.82	0.14		99	99	1	
07051300	**Acceptable identification**							**07051300**
					GRA	SPO		
Fusobacterium mortiferum		90.03	0.45	RHA 1	1	1		
Lactobacillus catenaformis		6.51	0.15		99	1		
Clostridium barati		3.46	0.05		99	99		
07051402	**Low discrimination**							**07051402**
					SPO			
Lactobacillus fermentum		64.57	0.42	CEL 1	1			
Clostridium innocuum		18.27	0.19		99			
Lactobacillus catenaformis		5.40	0.12		1			
07051500	**Low discrimination**							**07051500**
					GRA			
Fusobacterium mortiferum		48.07	0.45	ARA 1	1			
Lactobacillus fermentum		45.18	0.46		99			
Lactobacillus catenaformis		3.48	0.15		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07051502	**Low discrimination**							**07051502**
					GRA			
Lactobacillus fermentum		80.43	0.46	CEL 1	99			
Fusobacterium mortiferum		9.51	0.29		1			
Lactobacillus catenaformis		6.19	0.15		99			
07051520	**Low discrimination**							**07051520**
					GRA			
Lactobacillus fermentum		81.67	0.38	CEL 1	99			
Prevotella melaninogenica		5.89	-.01		1			
Lactobacillus catenaformis		4.71	0.05		99			
07051522	**Low discrimination**							**07051522**
					SPO			
Lactobacillus fermentum		80.33	0.38	CEL 1	1			
Clostridium barati		10.31	0.06		99			
Lactobacillus acidophilus		4.72	0.02		1			
07052000	**Low discrimination**							**07052000**
					SPO	CAT		
Propionibacterium granulosum		41.48	0.54	CEL 1	1	90		
Clostridium innocuum		38.38	0.44		99	1		
Eubacterium limosum		9.72	0.29		1	1		
07052002	**Low discrimination**							**07052002**
					COC	SPO	CAT	
Clostridium innocuum		77.64	0.44	SAL 99	1	99	1	
Propionibacterium granulosum		20.98	0.44	GAL 95	1	1	90	
Megasphaera elsdenii		0.67	0.12		99	1	1	
07052010	**Low discrimination**							**07052010**
					SPO	CAT		
Clostridium difficile		73.10	0.36	SUC 1	99	1		
Propionibacterium granulosum		20.69	0.39	CEL 5	1	90		
Clostridium innocuum		5.85	0.20		99	1		
07052020	**Acceptable identification**							**07052020**
					COC	SPO	CAT	
Propionibacterium granulosum		94.60	0.44	CEL 1	1	1	90	
Clostridium innocuum		3.54	0.10		1	99	1	
Gemella morbillorum		0.97	0.03		99	1	1	
07052100	**Low discrimination**							**07052100**
					GRA	SPO	CAT	
Fusobacterium mortiferum		84.34	0.42	MAN 1	1	1	5	
Propionibacterium granulosum		6.65	0.20		99	1	90	
Clostridium innocuum		6.15	0.10		99	99	1	
07053000	**Good identification**							**07053000**
					GRA	SPO	CAT	
Clostridium innocuum		93.95	0.66	SAL 99	99	99	1	
Propionibacterium granulosum		5.34	0.54		99	1	90	
Fusobacterium mortiferum		0.44	0.39		1	1	5	
07053002	**Very good identification**							**07053002**
					SPO	CAT		
Clostridium innocuum		98.34	0.66	SAL 99	99	1		
Propionibacterium granulosum		1.40	0.44		1	90		
Actinomyces naeslundii		0.07	0.36		1	25		
07053010	**Low discrimination**							**07053010**
					SPO	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		83.47	0.42	SAL 99	99	1	
Propionibacterium granulosum		15.54	0.39	MLZ 4	1	90	
Clostridium difficile		0.55	0.02		99	1	
07053012	**Good identification**						**07053012**
					SPO	CAT	
Clostridium innocuum		95.24	0.42	SAL 99	99	1	
Propionibacterium granulosum		4.43	0.29	MLZ 4	1	90	
Lactobacillus jensenii		0.33	0.03		1	1	
07053020	**Low discrimination**						**07053020**
					SPO	CAT	
Propionibacterium granulosum		55.19	0.44	CEL 1	1	90	
Clostridium innocuum		39.20	0.32		99	1	
Actinomyces naeslundii		2.83	0.36		1	25	
07053021	**Low discrimination**						**07053021**
Actinomyces naeslundii		88.40	0.39	MLT 94			
Propionibacterium granulosum		11.60	0.09	RAF 92			
07053023	**Good identification**						**07053023**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07053100	**Low discrimination**						**07053100**
					GRA	SPO	
Fusobacterium mortiferum		43.72	0.45	MAN 1	1	1	
Clostridium innocuum		40.38	0.32		99	99	
Actinomyces naeslundii		8.30	0.44		99	1	
07053101	**Good identification**						**07053101**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07053103	**Good identification**						**07053103**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07053120	**Low discrimination**						**07053120**
					GRA	SPO	
Actinomyces naeslundii		50.60	0.44	MLT 94	99	1	
Clostridium barati		30.74	0.14	RAF 92	99	99	
Prevotella melaninogenica		6.02	-.01		1	1	
07053121	**Good identification**						**07053121**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07053123	**Good identification**						**07053123**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07053400	**Good identification**						**07053400**
					SPO	CAT	
Clostridium innocuum		99.68	0.53	SAL 99	99	1	
Propionibacterium granulosum		0.33	0.20		1	90	
07053402	**Good identification**						**07053402**
					SPO	CAT	
Clostridium innocuum		99.92	0.53	SAL 99	99	1	
Propionibacterium granulosum		0.08	0.09		1	90	
07054000	**Low discrimination**						**07054000**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		58.42	0.50	XYL 7	1	1		
Gemella morbillorum		36.27	0.37		99	99		
Actinomyces meyeri		2.42	0.18		1	99		
07054100	**Good identification**							**07054100**
					COC	GRA		
Fusobacterium mortiferum		93.13	0.57	XYL 7	1	1		
Actinomyces meyeri		6.62	0.28		1	99		
Gemella morbillorum		0.25	0.03		99	99		
07054102	**Good identification**							**07054102**
					GRA			
Fusobacterium mortiferum		98.06	0.40	XYL 7	1			
Actinomyces meyeri		1.94	0.02	BGL 10	99			
07055000	**Low discrimination**							**07055000**
					COC	GRA		
Fusobacterium mortiferum		69.39	0.53	XYL 7	1	1		
Gemella morbillorum		15.47	0.32		99	99		
Lactobacillus fermentum		7.35	0.38		1	99		
07055002	**Low discrimination**							**07055002**
					GRA	SPO		
Fusobacterium mortiferum		41.88	0.37	XYL 7	1	1		
Lactobacillus fermentum		39.93	0.38	BGL 10	99	1		
Clostridium innocuum		5.74	0.10		99	99		
07055100	**Low discrimination**							**07055100**
					GRA			
Fusobacterium mortiferum		84.31	0.60	XYL 7	1			
Lactobacillus fermentum		6.25	0.42		99			
Actinomyces meyeri		5.99	0.31		99			
07055102	**Low discrimination**							**07055102**
					GRA			
Fusobacterium mortiferum		53.24	0.43	XYL 7	1			
Lactobacillus fermentum		35.49	0.42	BGL 10	99			
Lactobacillus catenaformis		4.65	0.15		99			
07055400	**Low discrimination**							**07055400**
					COC			
Lactobacillus fermentum		70.08	0.38	CEL 1	1			
Gemella morbillorum		16.38	0.16		99			
Actinomyces meyeri		6.85	0.10		1			
07055402	**Good identification**							**07055402**
Lactobacillus fermentum		100.0	0.38	CEL 1				
07055500	**Low discrimination**							**07055500**
					GRA			
Lactobacillus fermentum		72.67	0.42	CEL 1	99			
Actinomyces meyeri		17.42	0.21		99			
Fusobacterium mortiferum		9.91	0.26		1			
07055502	**Low discrimination**							**07055502**
Lactobacillus fermentum		85.98	0.42	CEL 1				
Bifidobacterium adolescentis		14.02	0.23					
07057000	**Good identification**							**07057000**
					GRA	SPO	CAT	
Clostridium innocuum		98.27	0.44	SAL 99	99	99	1	

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Propionibacterium granulosum		1.07	0.20	XYL 5	99	1	90	
Fusobacterium mortiferum		0.66	0.19		1	1	5	
07057002	**Good identification**							**07057002**
					SPO	CAT		
Clostridium innocuum		99.52	0.44	SAL 99	99	1		
Propionibacterium granulosum		0.27	0.09	XYL 5	1	90		
Actinomyces israelii		0.21	0.13		1	1		
07061100	**Low discrimination**							**07061100**
					COC	GRA		
Fusobacterium mortiferum		70.37	0.36	FRU 99	1	1		
Atopobium parvulum		29.63	0.18		99	99		
07063000	**Acceptable identification**							**07063000**
					SPO	CAT		
Clostridium innocuum		91.47	0.36	SAL 99	99	1		
Propionibacterium granulosum		8.53	0.28	FRU 95	1	90		
07063002	**Good identification**							**07063002**
					SPO	CAT		
Clostridium innocuum		97.72	0.36	SAL 99	99	1		
Propionibacterium granulosum		2.28	0.18	FRU 95	1	90		
07070000	**Low discrimination**							**07070000**
					GRA	CAT		
Propionibacterium granulosum		70.20	0.64	CEL 1	99	90		
Fusobacterium mortiferum		28.68	0.61		1	5		
Lactobacillus jensenii		0.61	0.22		99	1		
07070002	**Low discrimination**							**07070002**
					GRA	CAT		
Propionibacterium granulosum		82.78	0.54	CEL 1	99	90		
Fusobacterium mortiferum		15.03	0.44		1	5		
Lactobacillus jensenii		1.91	0.19		99	1		
07070010	**Acceptable identification**							**07070010**
					GRA	CAT		
Propionibacterium granulosum		94.44	0.49	CEL 1	99	90		
Fusobacterium mortiferum		2.86	0.27	MLZ 12	1	5		
Lactobacillus jensenii		2.00	0.14		99	1		
07070012	**Acceptable identification**							**07070012**
					CAT			
Propionibacterium granulosum		94.65	0.39	CEL 1	90			
Lactobacillus jensenii		5.35	0.11	MLZ 12	1			
07070020	**Good identification**							**07070020**
					GRA	CAT		
Propionibacterium granulosum		97.30	0.54	CEL 1	99	90		
Fusobacterium mortiferum		1.61	0.27		1	5		
Lactobacillus jensenii		0.84	0.12		99	1		
07070022	**Good identification**							**07070022**
					CAT			
Propionibacterium granulosum		97.74	0.43	CEL 1	90			
Lactobacillus jensenii		2.26	0.09		1			
07070030	**Good identification**							**07070030**
					CAT			
Propionibacterium granulosum		97.93	0.39	CEL 1	90			
Lactobacillus jensenii		2.07	0.04	MLZ 12	1			
07070100	**Very good identification**							**07070100**

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					COC	GRA	CAT	
Fusobacterium mortiferum		98.33	0.67		1	1	5	
Propionibacterium granulosum		1.04	0.30		1	99	90	
Atopobium parvulum		0.63	0.18		99	99	1	
07070102	**Acceptable identification**							**07070102**
					GRA	CAT		
Fusobacterium mortiferum		94.15	0.51	BGL 10	1	5		
Lactobacillus acidophilus		2.65	0.11		99	1		
Propionibacterium granulosum		2.25	0.20		99	90		
07070200	**Low discrimination**							**07070200**
					GRA	CAT		
Propionibacterium granulosum		88.23	0.38	CEL 1	99	90		
Fusobacterium mortiferum		11.77	0.27	RHA 3	1	5		
07071000	**Low discrimination**							**07071000**
					GRA	CAT		
Propionibacterium granulosum		39.96	0.64	CEL 1	99	90		
Lactobacillus jensenii		34.28	0.56		99	1		
Fusobacterium mortiferum		24.49	0.64		1	5		
07071002	**Low discrimination**							**07071002**
					GRA	CAT		
Lactobacillus jensenii		60.85	0.53	MLT 95	99	1		
Propionibacterium granulosum		26.60	0.54	SAL 95	99	90		
Fusobacterium mortiferum		7.24	0.47		1	5		
07071010	**Low discrimination**							**07071010**
					GRA	CAT		
Lactobacillus jensenii		66.63	0.48	MLT 95	99	1		
Propionibacterium granulosum		31.78	0.49	SAL 95	99	90		
Fusobacterium mortiferum		1.44	0.30		1	5		
07071012	**Low discrimination**							**07071012**
					GRA	CAT		
Lactobacillus jensenii		84.33	0.45	MLT 95	99	1		
Propionibacterium granulosum		15.08	0.39	SAL 95	99	90		
Fusobacterium mortiferum		0.30	0.13		1	5		
07071020	**Low discrimination**							**07071020**
					CAT			
Propionibacterium granulosum		50.24	0.54	CEL 1	90			
Lactobacillus jensenii		43.09	0.46		1			
Clostridium perfringens		3.96	0.27		1			
07071021	**Low discrimination**							**07071021**
					CAT			
Actinomyces naeslundii		50.78	0.39	MLT 94	25			
Propionibacterium granulosum		26.50	0.20	RAF 92	90			
Lactobacillus jensenii		22.73	0.12		1			
07071022	**Low discrimination**							**07071022**
					CAT			
Lactobacillus jensenii		57.08	0.43	MLT 95	1			
Propionibacterium granulosum		24.95	0.43	SAL 95	90			
Lactobacillus acidophilus		10.24	0.27		1			
07071023	**Low discrimination**							**07071023**
					CAT			
Actinomyces naeslundii		69.99	0.39	MLT 94	25			
Lactobacillus jensenii		20.88	0.09	RAF 92	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		9.13	0.09		90			
07071030	**Low discrimination**							**07071030**
					CAT			
Lactobacillus jensenii		67.71	0.38	MLT 95	1			
Propionibacterium granulosum		32.29	0.39	SAL 95	90			
07071100	**Low discrimination**							**07071100**
					COC	GRA	CAT	
Fusobacterium mortiferum		68.96	0.70		1	1	5	
Atopobium parvulum		29.04	0.52		99	99	1	
Propionibacterium granulosum		0.49	0.30		1	99	90	
07071101	**Low discrimination**							**07071101**
					COC	GRA		
Fusobacterium mortiferum		39.89	0.36	URE 1	1	1		
Actinomyces naeslundii		38.04	0.47		1	99		
Atopobium parvulum		16.80	0.18		99	99		
07071102	**Low discrimination**							**07071102**
					COC	GRA		
Lactobacillus acidophilus		50.11	0.45	MLT 94	1	99		
Fusobacterium mortiferum		26.98	0.54	SAL 99	1	1		
Atopobium parvulum		18.05	0.39		99	99		
07071103	**Low discrimination**							**07071103**
					GRA			
Actinomyces naeslundii		70.88	0.47	MLT 94	99			
Lactobacillus acidophilus		15.34	0.11	RAF 92	99			
Fusobacterium mortiferum		8.26	0.20		1			
07071110	**Low discrimination**							**07071110**
					COC	GRA		
Fusobacterium mortiferum		58.26	0.36	MLZ 1	1	1		
Atopobium parvulum		24.53	0.18		99	99		
Lactobacillus jensenii		11.65	0.14		1	99		
07071120	**Low discrimination**							**07071120**
					COC	SPO		
Clostridium perfringens		41.48	0.42	MLT 99	1	84		
Atopobium parvulum		30.53	0.39	CEL 4	99	1		
Clostridium septicum		16.83	0.29		1	99		
07071121	**Low discrimination**							**07071121**
					COC			
Actinomyces naeslundii		78.55	0.47	MLT 94	1			
Clostridium perfringens		8.32	0.08	RAF 92	1			
Atopobium parvulum		6.12	0.05		99			
07071122	**Low discrimination**							**07071122**
					COC			
Lactobacillus acidophilus		70.27	0.45	MLT 94	1			
Clostridium perfringens		16.18	0.37	SAL 99	1			
Atopobium parvulum		4.47	0.26		99			
07071123	**Low discrimination**							**07071123**
Actinomyces naeslundii		78.97	0.47	MLT 94				
Lactobacillus acidophilus		17.09	0.11	RAF 92				
Clostridium perfringens		3.94	0.03					
07071140	**Low discrimination**							**07071140**
					COC	GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		62.81	0.36	SOR 1	1	1		
Atopobium parvulum		26.45	0.18		99	99		
Actinomyces naeslundii		7.05	0.31		1	99		
07071200	**Low discrimination**							**07071200**
					GRA	CAT		
Propionibacterium granulosum		67.56	0.38	CEL 1	99	90		
Lactobacillus jensenii		18.93	0.22	RHA 3	99	1		
Fusobacterium mortiferum		13.52	0.30		1	5		
07071300	**Low discrimination**							**07071300**
					COC	GRA		
Fusobacterium mortiferum		70.37	0.36	RHA 1	1	1		
Atopobium parvulum		29.63	0.18		99	99		
07071500	**Low discrimination**							**07071500**
					COC	GRA		
Fusobacterium mortiferum		70.37	0.36	ARA 1	1	1		
Atopobium parvulum		29.63	0.18		99	99		
07072000	**Low discrimination**							**07072000**
					GRA	SPO	CAT	
Propionibacterium granulosum		89.52	0.62	CEL 1	99	1	90	
Clostridium innocuum		9.70	0.36		99	99	1	
Fusobacterium mortiferum		0.49	0.27		1	1	5	
07072002	**Low discrimination**							**07072002**
					SPO	CAT		
Propionibacterium granulosum		69.76	0.52	CEL 1	1	90		
Clostridium innocuum		30.24	0.36		99	1		
07072010	**Low discrimination**							**07072010**
					SPO	CAT		
Propionibacterium granulosum		89.15	0.47	CEL 1	1	90		
Clostridium difficile		7.90	0.17	MLZ 12	99	1		
Clostridium innocuum		2.95	0.13		99	1		
07072012	**Low discrimination**							**07072012**
					SPO	CAT		
Propionibacterium granulosum		88.30	0.37	CEL 1	1	90		
Clostridium innocuum		11.70	0.13	MLZ 12	99	1		
07072020	**Good identification**							**07072020**
					SPO	CAT		
Propionibacterium granulosum		99.56	0.52	CEL 1	1	90		
Clostridium innocuum		0.44	0.02		99	1		
07072022	**Good identification**							**07072022**
					SPO	CAT		
Propionibacterium granulosum		98.28	0.41	CEL 1	1	90		
Clostridium innocuum		1.72	0.02		99	1		
07072030	**Good identification**							**07072030**
Propionibacterium granulosum		100.0	0.37	CEL 1				
07072200	**Good identification**							**07072200**
					SPO	CAT		
Propionibacterium granulosum		96.58	0.36	CEL 1	1	90		
Clostridium innocuum		3.42	0.02	RHA 3	99	1		
07073000	**Low discrimination**							**07073000**
					SPO	CAT		
Clostridium innocuum		66.70	0.58	SAL 99	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		32.39	0.62		1	90	
Lactobacillus jensenii		0.37	0.22		1	1	
07073002	**Low discrimination**						**07073002**
					SPO	CAT	
Clostridium innocuum		88.46	0.58	SAL 99	99	1	
Propionibacterium granulosum		10.74	0.52		1	90	
Actinomyces naeslundii		0.36	0.41		1	25	
07073010	**Low discrimination**						**07073010**
					SPO	CAT	
Propionibacterium granulosum		60.34	0.47	CEL 1	1	90	
Clostridium innocuum		37.97	0.34	MLZ 12	99	1	
Lactobacillus jensenii		1.69	0.14		1	1	
07073020	**Low discrimination**						**07073020**
					SPO	CAT	
Propionibacterium granulosum		88.66	0.52	CEL 1	1	90	
Clostridium innocuum		7.38	0.24		99	1	
Actinomyces naeslundii		2.94	0.41		1	25	
07073021	**Low discrimination**						**07073021**
Actinomyces naeslundii		83.14	0.44	MLT 94			
Propionibacterium granulosum		16.86	0.18	RAF 92			
07073022	**Low discrimination**						**07073022**
					SPO	CAT	
Propionibacterium granulosum		65.58	0.41	CEL 1	1	90	
Clostridium innocuum		21.82	0.24		99	1	
Actinomyces naeslundii		8.71	0.41		1	25	
07073023	**Good identification**						**07073023**
Actinomyces naeslundii		100.0	0.44	MLT 94			
07073030	**Acceptable identification**						**07073030**
					SPO	CAT	
Propionibacterium granulosum		94.92	0.37	CEL 1	1	90	
Lactobacillus jensenii		2.67	0.04	MLZ 12	1	1	
Clostridium innocuum		2.41	0.00		99	1	
07073100	**Low discrimination**						**07073100**
					GRA	SPO	
Actinomyces naeslundii		28.62	0.49	MLT 94	99	1	
Clostridium innocuum		25.19	0.24	RAF 92	99	99	
Fusobacterium mortiferum		23.19	0.36		1	1	
07073101	**Good identification**						**07073101**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07073102	**Low discrimination**						**07073102**
					SPO		
Actinomyces naeslundii		31.40	0.49	MLT 94	1		
Clostridium innocuum		27.64	0.24	RAF 92	99		
Lactobacillus acidophilus		27.36	0.24		1		
07073103	**Good identification**						**07073103**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07073120	**Low discrimination**						**07073120**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		67.26	0.49	MLT 94	1	25	
Propionibacterium granulosum		7.19	0.18	RAF 92	1	90	
Clostridium tertium		7.08	0.09		99	1	
07073121	**Good identification**						**07073121**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07073122	**Low discrimination**						**07073122**
					SPO		
Actinomyces naeslundii		36.98	0.49	MLT 94	1		
Lactobacillus acidophilus		32.22	0.24	RAF 92	1		
Clostridium ramosum		14.68	0.17		92		
07073123	**Good identification**						**07073123**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07073140	**Good identification**						**07073140**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07073141	**Good identification**						**07073141**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07073142	**Low discrimination**						**07073142**
Actinomyces naeslundii		55.40	0.36	MLT 94			
Lactobacillus acidophilus		33.81	0.08	RAF 92			
Propionibacterium propionicum		10.79	0.14				
07073143	**Good identification**						**07073143**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07073160	**Good identification**						**07073160**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07073161	**Good identification**						**07073161**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07073162	**Low discrimination**						**07073162**
Actinomyces naeslundii		62.10	0.36	MLT 94			
Lactobacillus acidophilus		37.90	0.08	RAF 92			
07073163	**Good identification**						**07073163**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07073200	**Low discrimination**						**07073200**
					SPO	CAT	
Propionibacterium granulosum		59.79	0.36	CEL 1	1	90	
Clostridium innocuum		40.21	0.24	RHA 3	99	1	
07073400	**Good identification**						**07073400**
					SPO	CAT	
Clostridium innocuum		97.30	0.45	SAL 99	99	1	
Propionibacterium granulosum		2.71	0.28		1	90	
07073402	**Good identification**						**07073402**
					SPO	CAT	
Clostridium innocuum		99.31	0.45	SAL 99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.69	0.18		1	90		
07074000	**Low discrimination**							**07074000**
					COC	GRA	CAT	
Fusobacterium mortiferum		74.10	0.41	XYL 7	1	1	5	
Propionibacterium granulosum		24.34	0.30		1	99	90	
Gemella morbillorum		1.56	0.03		99	99	1	
07074100	**Low discrimination**							**07074100**
					COC	GRA		
Fusobacterium mortiferum		86.40	0.48	XYL 7	1	1		
Atopobium parvulum		13.60	0.23		99	99		
07075000	**Low discrimination**							**07075000**
					COC	GRA	CAT	
Fusobacterium mortiferum		59.89	0.44	XYL 7	1	1	5	
Atopobium parvulum		14.67	0.23		99	99	1	
Propionibacterium granulosum		13.12	0.30		1	99	90	
07075100	**Good identification**							**07075100**
					COC	GRA		
Atopobium parvulum		91.16	0.57	MLT 99	99	99		
Fusobacterium mortiferum		8.77	0.51		1	1		
Actinomyces israelii		0.07	0.17		1	99		
07075102	**Acceptable identification**							**07075102**
					COC	GRA		
Atopobium parvulum		92.39	0.44	MLT 99	99	99		
Fusobacterium mortiferum		5.60	0.34		1	1		
Lactobacillus acidophilus		1.40	0.11		1	99		
07075120	**Good identification**							**07075120**
					COC	GRA		
Atopobium parvulum		98.44	0.44	MLT 99	99	99		
Clostridium perfringens		0.73	0.08		1	99		
Fusobacterium mortiferum		0.54	0.17		1	1		
07075322	**Good identification**							**07075322**
Parabacteroides distasonis		100.0	0.45	RAF 99				
07075332	**Good identification**							**07075332**
Parabacteroides distasonis		100.0	0.40	RAF 99				
07075722	**Good identification**							**07075722**
Parabacteroides distasonis		96.79	0.41	RAF 99				
Clostridium clostridioforme		3.21	0.10					
07075732	**Low discrimination**							**07075732**
Parabacteroides distasonis		78.09	0.37	RAF 99				
Clostridium clostridioforme		21.91	0.22					
07077000	**Acceptable identification**							**07077000**
					SPO	CAT		
Clostridium innocuum		90.23	0.36	SAL 99	99	1		
Propionibacterium granulosum		8.41	0.28	XYL 5	1	90		
Actinomyces israelii		1.36	0.20		1	1		
07077002	**Good identification**							**07077002**
					SPO	CAT		
Clostridium innocuum		95.61	0.36	SAL 99	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		2.23	0.18	XYL 5	1	90		
Actinomyces israelii		2.17	0.23		1	1		
07100000	**Very good identification**							**07100000**
					COC	GRA	SPO	
Gemella morbillorum		95.20	0.99		99	99	1	
Prevotella disiens		3.76	0.66		1	1	1	
Clostridium botulinum biovar A		0.43	0.55		1	99	99	
07100001	**Good identification**							**07100001**
					COC	GRA	SPO	
Gemella morbillorum		94.47	0.64	URE 1	99	99	1	
Prevotella disiens		3.73	0.32		1	1	1	
Clostridium botulinum biovar A		0.42	0.21		1	99	99	
07100002	**Good identification**							**07100002**
					COC	GRA	CAT	
Gemella morbillorum		92.42	0.64	BGL 1	99	99	1	
Prevotella disiens		3.65	0.32		1	1	1	
Propionibacterium granulosum		1.30	0.40		1	99	90	
07100010	**Low discrimination**							**07100010**
					COC	GRA	SPO	
Gemella morbillorum		85.74	0.64	MLZ 1	99	99	1	
Clostridium botulinum biovar B		9.23	0.42		1	99	99	
Prevotella disiens		3.38	0.32		1	1	1	
07100020	**Very good identification**							**07100020**
					COC	GRA	SPO	
Gemella morbillorum		97.56	0.82	NAG 10	99	99	1	
Clostridium botulinum biovar B		0.95	0.42		1	99	99	
Prevotella bivia		0.72	0.32		1	1	1	
07100021	**Good identification**							**07100021**
					COC	SPO		
Gemella morbillorum		96.40	0.48	NAG 10	99	1		
Clostridium botulinum biovar B		0.94	0.08	URE 1	1	99		
Clostridium novyi biovar A		0.81	0.21		1	99		
07100022	**Acceptable identification**							**07100022**
					COC	CAT		
Gemella morbillorum		92.99	0.48	NAG 10	99	1		
Propionibacterium granulosum		2.94	0.29	BGL 1	1	90		
Atopobium minutum		1.77	0.23		1	1		
07100030	**Low discrimination**							**07100030**
					COC	SPO	CAT	
Gemella morbillorum		66.07	0.48	MLZ 1	99	1	1	
Clostridium botulinum biovar B		31.52	0.37	NAG 10	1	99	1	
Propionibacterium granulosum		1.14	0.25		1	1	90	
07100040	**Good identification**							**07100040**
					COC	SPO		
Clostridium botulinum biovar B		94.68	0.82		1	99		
Gemella morbillorum		4.38	0.64		99	1		
Clostridium botulinum biovar A		0.58	0.47		1	99		
07100041	**Acceptable identification**							**07100041**
					COC	SPO		
Clostridium botulinum biovar B		93.71	0.47	URE 1	1	99		
Gemella morbillorum		4.33	0.30		99	1		
Clostridium novyi biovar A		1.06	0.29		1	99		

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07100042	**Acceptable identification**							**07100042**
					COC	GRA	SPO	
Clostridium botulinum biovar B		93.09	0.47	BGL 1	1	99	99	
Gemella morbillorum		4.30	0.30		99	99	1	
Megasphaera elsdenii		1.68	0.25		99	1	1	
07100050	**Very good identification**							**07100050**
					COC	SPO		
Clostridium botulinum biovar B		99.86	0.76		1	99		
Gemella morbillorum		0.10	0.30		99	1		
Clostridium novyi biovar A		0.02	0.29		1	99		
07100051	**Good identification**							**07100051**
Clostridium botulinum biovar B		97.75	0.42	URE 1				
Clostridium novyi biovar A		2.25	0.29					
07100052	**Good identification**							**07100052**
Clostridium botulinum biovar B		99.60	0.42	BGL 1				
Clostridium novyi biovar A		0.40	0.16					
07100060	**Very good identification**							**07100060**
					COC	SPO		
Clostridium botulinum biovar B		98.93	0.76		1	99		
Gemella morbillorum		1.03	0.48		99	1		
Clostridium novyi biovar A		0.01	0.18		1	99		
07100061	**Good identification**							**07100061**
					COC	SPO		
Clostridium botulinum biovar B		98.41	0.42	URE 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
Clostridium novyi biovar A		0.57	0.18		1	99		
07100062	**Good identification**							**07100062**
					COC	SPO		
Clostridium botulinum biovar B		98.97	0.42	BGL 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
07100070	**Very good identification**							**07100070**
					COC	SPO		
Clostridium botulinum biovar B		99.97	0.71		1	99		
Gemella morbillorum		0.02	0.14		99	1		
Clostridium novyi biovar A		0.01	0.18		1	99		
07100071	**Good identification**							**07100071**
Clostridium botulinum biovar B		98.85	0.37	URE 1				
Clostridium novyi biovar A		1.15	0.18					
07100072	**Good identification**							**07100072**
Clostridium botulinum biovar B		100.0	0.37	BGL 1				
07100100	**Low discrimination**							**07100100**
					COC	GRA		
Gemella morbillorum		86.23	0.64	LAC 1	99	99		
Prevotella bivia		6.99	0.32		1	1		
Prevotella disiens		3.40	0.32		1	1		
07100101	**Good identification**							**07100101**
					COC	GRA	SPO	
Clostridium chauvoei		95.96	0.54	GAL 99	1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		3.54	0.30		99	99	1	
Prevotella bivia		0.29	-.03		1	1	1	
07100103	**Good identification**							**07100103**
					SPO			
Clostridium chauvoei		96.28	0.41	GAL 99	99			
Actinomyces viscosus		3.72	0.15		1			
07100120	**Very good identification**							**07100120**
					COC	GRA		
Prevotella bivia		98.00	0.66	SUC 1	1	1		
Gemella morbillorum		1.36	0.48		99	99		
Prevotella melaninogenica		0.50	0.33		1	1		
07100121	**Low discrimination**							**07100121**
					COC	GRA	SPO	
Clostridium chauvoei		52.46	0.45	GAL 99	1	99	99	
Prevotella bivia		46.66	0.32		1	1	1	
Gemella morbillorum		0.65	0.14		99	99	1	
07100140	**Acceptable identification**							**07100140**
					COC	SPO		
Clostridium botulinum biovar B		94.22	0.47	LAC 1	1	99		
Gemella morbillorum		4.36	0.30		99	1		
Clostridium botulinum biovar A		0.58	0.12		1	99		
07100150	**Good identification**							**07100150**
Clostridium botulinum biovar B		100.0	0.42	LAC 1				
07100160	**Low discrimination**							**07100160**
					GRA	SPO		
Clostridium botulinum biovar B		56.17	0.42	LAC 1	99	99		
Prevotella bivia		42.33	0.32		1	1		
Clostridium perfringens		0.70	0.15		99	84		
07100170	**Good identification**							**07100170**
					GRA	SPO		
Clostridium botulinum biovar B		98.48	0.37	LAC 1	99	99		
Prevotella bivia		1.52	-.03		1	1		
07100200	**Good identification**							**07100200**
					COC	GRA	SPO	
Gemella morbillorum		94.86	0.64	RHA 1	99	99	1	
Prevotella disiens		3.74	0.32		1	1	1	
Clostridium botulinum biovar A		0.43	0.21		1	99	99	
07100220	**Good identification**							**07100220**
					COC	GRA	SPO	
Gemella morbillorum		97.19	0.48	RHA 1	99	99	1	
Clostridium botulinum biovar B		0.95	0.08	NAG 10	1	99	99	
Prevotella bivia		0.72	-.03		1	1	1	
07100240	**Acceptable identification**							**07100240**
					COC	SPO		
Clostridium botulinum biovar B		94.71	0.47	RHA 1	1	99		
Gemella morbillorum		4.38	0.30		99	1		
Clostridium botulinum biovar A		0.58	0.12		1	99		
07100250	**Good identification**							**07100250**
Clostridium botulinum biovar B		100.0	0.42	RHA 1				
07100260	**Good identification**							**07100260**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Clostridium botulinum biovar B		98.97	0.42	RHA 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
07100270	**Good identification**							**07100270**
Clostridium botulinum biovar B		100.0	0.37	RHA 1				
07100400	**Very good identification**							**07100400**
					COC	GRA	SPO	
Gemella morbillorum		99.53	0.82	ARA 10	99	99	1	
Prevotella disiens		0.36	0.32		1	1	1	
Clostridium botulinum biovar A		0.04	0.21		1	99	99	
07100401	**Good identification**							**07100401**
					COC	GRA		
Gemella morbillorum		99.64	0.48	ARA 10	99	99		
Prevotella disiens		0.36	-.03	URE 1	1	1		
07100402	**Good identification**							**07100402**
					COC	GRA		
Gemella morbillorum		99.36	0.48	ARA 10	99	99		
Prevotella disiens		0.36	-.03	BGL 1	1	1		
Lactobacillus fermentum		0.28	0.16		1	99		
07100410	**Good identification**							**07100410**
					COC	GRA	SPO	
Gemella morbillorum		98.68	0.48	ARA 10	99	99	1	
Clostridium botulinum biovar B		0.97	0.08	MLZ 1	1	99	99	
Prevotella disiens		0.35	-.03		1	1	1	
07100420	**Very good identification**							**07100420**
					COC	GRA	SPO	
Gemella morbillorum		99.81	0.66	ARA 10	99	99	1	
Clostridium botulinum biovar B		0.09	0.08	NAG 10	1	99	99	
Prevotella bivia		0.07	-.03		1	1	1	
07100440	**Low discrimination**							**07100440**
					COC	SPO		
Clostridium botulinum biovar B		65.87	0.47	ARA 1	1	99		
Gemella morbillorum		33.50	0.48		99	1		
Clostridium botulinum biovar A		0.40	0.12		1	99		
07100450	**Good identification**							**07100450**
					COC	SPO		
Clostridium botulinum biovar B		98.97	0.42	ARA 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
07100460	**Low discrimination**							**07100460**
					COC	SPO		
Clostridium botulinum biovar B		89.71	0.42	ARA 1	1	99		
Gemella morbillorum		10.29	0.32		99	1		
07100470	**Good identification**							**07100470**
Clostridium botulinum biovar B		100.0	0.37	ARA 1				
07100500	**Acceptable identification**							**07100500**
					COC	GRA		
Gemella morbillorum		94.41	0.48	LAC 1	99	99		
Actinomyces meyeri		4.11	0.26	ARA 10	1	99		
Prevotella bivia		0.70	-.03		1	1		
07100600	**Good identification**							**07100600**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Gemella morbillorum		99.64	0.48	RHA 1	99	99		
Prevotella disiens		0.36	-.03	ARA 10	1	1		
07101000	**Excellent identification**							**07101000**
					COC	GRA	CAT	
Gemella morbillorum		99.52	0.94		99	99	1	
Prevotella intermedia		0.19	0.46		1	1	1	
Propionibacterium granulosum		0.11	0.50		1	99	90	
07101001	**Low discrimination**							**07101001**
					COC	GRA	SPO	
Gemella morbillorum		66.03	0.60	URE 1	99	99	1	
Clostridium chauvoei		33.54	0.54		1	99	99	
Prevotella intermedia		0.12	0.12		1	1	1	
07101002	**Good identification**							**07101002**
					COC	CAT		
Gemella morbillorum		90.74	0.60	BGL 1	99	1		
Lactobacillus fermentum		5.28	0.50		1	1		
Propionibacterium granulosum		2.37	0.40		1	90		
07101003	**Low discrimination**							**07101003**
					COC	SPO		
Clostridium chauvoei		87.37	0.41	LAC 99	1	99		
Gemella morbillorum		9.85	0.26		99	1		
Actinomyces viscosus		1.17	0.07		1	1		
07101010	**Very good identification**							**07101010**
					COC	SPO	CAT	
Gemella morbillorum		97.96	0.60	MLZ 1	99	1	1	
Propionibacterium granulosum		1.39	0.35		1	1	90	
Clostridium botulinum biovar B		0.20	0.08		1	99	1	
07101020	**Very good identification**							**07101020**
					COC	GRA		
Gemella morbillorum		97.04	0.78	NAG 10	99	99		
Clostridium perfringens		1.56	0.46		1	99		
Prevotella melaninogenica		0.66	0.33		1	1		
07101021	**Low discrimination**							**07101021**
					COC	SPO		
Clostridium chauvoei		59.61	0.45	LAC 99	1	99		
Gemella morbillorum		39.12	0.43		99	1		
Clostridium perfringens		0.63	0.12		1	84		
07101022	**Low discrimination**							**07101022**
					COC			
Gemella morbillorum		46.81	0.43	NAG 10	99			
Clostridium perfringens		35.13	0.41	BGL 1	1			
Lactobacillus fermentum		8.17	0.42		1			
07101030	**Acceptable identification**							**07101030**
					COC	CAT		
Gemella morbillorum		93.99	0.43	MLZ 1	99	1		
Propionibacterium granulosum		3.01	0.25	NAG 10	1	90		
Clostridium perfringens		1.51	0.12		1	1		
07101040	**Low discrimination**							**07101040**
					COC	SPO		
Gemella morbillorum		70.69	0.60	SOR 1	99	1		
Clostridium botulinum biovar B		28.68	0.47		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		0.18	0.12		1	99	
07101050	**Acceptable identification**						**07101050**
					COC	SPO	
Clostridium botulinum biovar B		94.96	0.42	GAL 1	1	99	
Gemella morbillorum		4.80	0.26		99	1	
Clostridium novyi biovar A		0.24	0.12		1	99	
07101060	**Low discrimination**						**07101060**
					COC	SPO	
Clostridium botulinum biovar B		57.85	0.42	GAL 1	1	99	
Gemella morbillorum		32.16	0.43		99	1	
Clostridium perfringens		9.77	0.34		1	84	
07101070	**Good identification**						**07101070**
					COC	SPO	
Clostridium botulinum biovar B		98.87	0.37	GAL 1	1	99	
Gemella morbillorum		1.13	0.09		99	1	
07101100	**Low discrimination**						**07101100**
					COC	SPO	
Gemella morbillorum		54.92	0.60	LAC 1	99	1	
Clostridium chauvoei		27.89	0.54		1	99	
Lactobacillus fermentum		5.21	0.54		1	1	
07101101	**Very good identification**						**07101101**
					COC	SPO	
Clostridium chauvoei		99.97	0.88		1	99	
Gemella morbillorum		0.02	0.26		99	1	
Actinomyces naeslundii		0.01	0.39		1	1	
07101102	**Low discrimination**						**07101102**
					SPO		
Lactobacillus fermentum		39.43	0.54	FRU 99	1		
Clostridium chauvoei		37.23	0.41		99		
Clostridium perfringens		7.86	0.30		84		
07101103	**Very good identification**						**07101103**
					SPO		
Clostridium chauvoei		99.93	0.75		99		
Actinomyces naeslundii		0.04	0.39		1		
Actinomyces viscosus		0.02	0.12		1		
07101111	**Good identification**						**07101111**
Clostridium chauvoei		100.0	0.54	MLZ 1			
07101113	**Good identification**						**07101113**
Clostridium chauvoei		100.0	0.41	MLZ 1			
07101120	**Low discrimination**						**07101120**
					GRA		
Prevotella melaninogenica		63.67	0.67	RAF 90	1		
Prevotella bivia		22.47	0.53	FRU 90	1		
Clostridium perfringens		11.10	0.61		99		
07101121	**Very good identification**						**07101121**
					GRA	SPO	
Clostridium chauvoei		99.29	0.80		99	99	
Prevotella melaninogenica		0.45	0.33		1	1	
Prevotella bivia		0.16	0.19		1	1	
07101122	**Low discrimination**						**07101122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Prevotella melaninogenica		66.49	0.55	RAF 90	1		
Clostridium perfringens		28.64	0.56	FRU 90	99		
Lactobacillus fermentum		1.48	0.46		99		
07101123	**Very good identification**						**07101123**
					GRA	SPO	
Clostridium chauvoei		99.21	0.67		99	99	
Prevotella melaninogenica		0.48	0.21		1	1	
Clostridium perfringens		0.21	0.22		99	84	
07101131	**Good identification**						**07101131**
					GRA	SPO	
Clostridium chauvoei		99.55	0.45	MLZ 1	99	99	
Prevotella melaninogenica		0.45	-.01		1	1	
07101141	**Good identification**						**07101141**
					SPO		
Clostridium chauvoei		99.89	0.54	SOR 1	99		
Actinomyces naeslundii		0.11	0.26		1		
07101143	**Good identification**						**07101143**
					SPO		
Clostridium chauvoei		99.40	0.41	SOR 1	99		
Actinomyces naeslundii		0.60	0.26		1		
07101160	**Low discrimination**						**07101160**
					GRA		
Clostridium perfringens		69.80	0.49	FRU 99	99		
Prevotella melaninogenica		21.23	0.33		1		
Prevotella bivia		7.50	0.19		1		
07101161	**Good identification**						**07101161**
					GRA	SPO	
Clostridium chauvoei		97.80	0.45	SOR 1	99	99	
Clostridium perfringens		1.44	0.15		99	84	
Prevotella melaninogenica		0.44	-.01		1	1	
07101162	**Low discrimination**						**07101162**
					GRA		
Clostridium perfringens		88.79	0.43	FRU 99	99		
Prevotella melaninogenica		10.93	0.21		1		
Actinomyces naeslundii		0.28	0.23		99		
07101200	**Very good identification**						**07101200**
					COC	CAT	
Gemella morbillorum		99.01	0.60	RHA 1	99	1	
Actinomyces odontolyticus		0.36	0.23		1	10	
Propionibacterium granulosum		0.32	0.24		1	90	
07101220	**Good identification**						**07101220**
					COC	CAT	
Gemella morbillorum		96.57	0.43	RHA 1	99	1	
Clostridium perfringens		1.56	0.12	NAG 10	1	1	
Propionibacterium granulosum		0.70	0.14		1	90	
07101301	**Good identification**						**07101301**
Clostridium chauvoei		100.0	0.54	RHA 1			
07101303	**Good identification**						**07101303**
Clostridium chauvoei		100.0	0.41	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07101321	**Good identification**						**07101321**
					GRA	SPO	
Clostridium chauvoei		99.55	0.45	RHA 1	99	99	
Prevotella melaninogenica		0.45	-.01		1	1	
07101400	**Very good identification**						**07101400**
					COC		
Gemella morbillorum		99.41	0.78	ARA 10	99		
Lactobacillus fermentum		0.53	0.50		1		
Actinomyces meyeri		0.03	0.18		1		
07101401	**Good identification**						**07101401**
					COC	SPO	
Gemella morbillorum		95.11	0.43	ARA 10	99	1	
Clostridium chauvoei		4.39	0.19	URE 1	1	99	
Lactobacillus fermentum		0.50	0.16		1	1	
07101402	**Low discrimination**						**07101402**
					COC		
Gemella morbillorum		65.63	0.43	ARA 10	99		
Lactobacillus fermentum		34.37	0.50	BGL 1	1		
07101410	**Good identification**						**07101410**
					COC		
Gemella morbillorum		99.17	0.43	ARA 10	99		
Lactobacillus fermentum		0.52	0.16	MLZ 1	1		
Bifidobacterium longum subsp. longum		0.31	0.05		1		
07101420	**Very good identification**						**07101420**
					COC		
Gemella morbillorum		98.14	0.61	ARA 10	99		
Lactobacillus fermentum		1.56	0.42	NAG 10	1		
Clostridium perfringens		0.14	0.12		1		
07101422	**Low discrimination**						**07101422**
					COC		
Lactobacillus fermentum		58.08	0.42	FRU 99	1		
Gemella morbillorum		36.98	0.27		99		
Clostridium perfringens		2.52	0.07		1		
07101440	**Good identification**						**07101440**
					COC	SPO	
Gemella morbillorum		95.95	0.43	ARA 10	99	1	
Clostridium botulinum biovar B		3.54	0.13	SOR 1	1	99	
Lactobacillus fermentum		0.51	0.16		1	1	
07101500	**Low discrimination**						**07101500**
					COC		
Gemella morbillorum		45.20	0.43	LAC 1	99		
Lactobacillus fermentum		38.62	0.54	ARA 10	1		
Bifidobacterium longum subsp. longum		8.13	0.35		1		
07101501	**Good identification**						**07101501**
					COC	SPO	
Clostridium chauvoei		99.59	0.54	ARA 1	1	99	
Gemella morbillorum		0.22	0.09		99	1	
Lactobacillus fermentum		0.19	0.20		1	1	
07101502	**Low discrimination**						**07101502**
					COC		
Lactobacillus fermentum		80.88	0.54	FRU 99	1		
Bifidobacterium longum subsp. longum		17.03	0.35		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		0.96	0.09		99			
07101503	**Good identification**							**07101503**
					SPO			
Clostridium chauvoei		98.94	0.41	ARA 1	99			
Lactobacillus fermentum		1.06	0.20		1			
07101510	**Acceptable identification**							**07101510**
					COC			
Bifidobacterium longum subsp. longum		94.24	0.39	RAF 97	1			
Gemella morbillorum		3.11	0.09	FRU 90	99			
Lactobacillus fermentum		2.66	0.20		1			
07101512	**Good identification**							**07101512**
Bifidobacterium longum subsp. longum		97.26	0.39	RAF 97				
Lactobacillus fermentum		2.74	0.20	FRU 90				
07101520	**Low discrimination**							**07101520**
					GRA			
Bifidobacterium longum subsp. longum		32.97	0.46	RAF 97	99			
Prevotella melaninogenica		31.05	0.33	FRU 90	1			
Lactobacillus fermentum		13.05	0.46		99			
07101521	**Good identification**							**07101521**
					GRA	SPO		
Clostridium chauvoei		99.08	0.45	ARA 1	99	99		
Bifidobacterium longum subsp. longum		0.47	0.12		99	1		
Prevotella melaninogenica		0.45	-.01		1	1		
07101522	**Low discrimination**							**07101522**
					GRA			
Bifidobacterium longum subsp. longum		60.30	0.46	RAF 97	99			
Lactobacillus fermentum		23.86	0.46	FRU 90	99			
Prevotella melaninogenica		10.82	0.21		1			
07101530	**Good identification**							**07101530**
					GRA			
Bifidobacterium longum subsp. longum		99.44	0.50	RAF 97	99			
Prevotella melaninogenica		0.56	-.01	FRU 90	1			
07101532	**Good identification**							**07101532**
Bifidobacterium longum subsp. longum		100.0	0.50	RAF 97				
07101600	**Good identification**							**07101600**
					COC			
Gemella morbillorum		99.47	0.43	RHA 1	99			
Lactobacillus fermentum		0.53	0.16	ARA 10	1			
07102000	**Low discrimination**							**07102000**
					COC	GRA	CAT	
Gemella morbillorum		87.34	0.64	MAN 1	99	99	1	
Megasphaera elsdenii		4.64	0.44		99	1	1	
Propionibacterium granulosum		3.70	0.48		1	99	90	
07102002	**Low discrimination**							**07102002**
					COC	GRA	CAT	
Propionibacterium granulosum		38.84	0.38	FRU 99	1	99	90	
Gemella morbillorum		37.04	0.30		99	99	1	
Megasphaera elsdenii		21.63	0.28		99	1	1	
07102020	**Low discrimination**							**07102020**
					COC	SPO	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		89.22	0.48	MAN 1	99	1	1	
Propionibacterium granulosum		8.50	0.38	NAG 10	1	1	90	
Clostridium botulinum biovar B		0.87	0.08		1	99	1	
07102040	**Low discrimination**							**07102040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		82.03	0.47	MAN 1	1	99	99	
Megasphaera elsdenii		13.29	0.41		99	1	1	
Gemella morbillorum		3.79	0.30		99	99	1	
07102050	**Good identification**							**07102050**
					COC	GRA		
Clostridium botulinum biovar B		99.67	0.42	MAN 1	1	99		
Megasphaera elsdenii		0.33	0.07		99	1		
07102060	**Good identification**							**07102060**
					COC	GRA	SPO	
Clostridium botulinum biovar B		98.64	0.42	MAN 1	1	99	99	
Gemella morbillorum		1.03	0.14		99	99	1	
Megasphaera elsdenii		0.33	0.07		99	1	1	
07102070	**Good identification**							**07102070**
Clostridium botulinum biovar B		100.0	0.37	MAN 1				
07102100	**Low discrimination**							**07102100**
					COC	GRA		
Propionibacterium propionicum		59.77	0.46	NIT 86	1	99		
Gemella morbillorum		32.57	0.30	FRU 99	99	99		
Prevotella bivia		2.64	-.03		1	1		
07102102	**Good identification**							**07102102**
Propionibacterium propionicum		100.0	0.49	NIT 86				
07102140	**Low discrimination**							**07102140**
					COC	SPO	GRA	
Propionibacterium propionicum		87.85	0.46	NIT 86	1	1	99	
Clostridium botulinum biovar B		10.46	0.13	FRU 99	1	99	99	
Megasphaera elsdenii		1.69	0.07		99	1	1	
07102142	**Good identification**							**07102142**
Propionibacterium propionicum		100.0	0.49	NIT 86				
07102400	**Good identification**							**07102400**
					COC	GRA	CAT	
Gemella morbillorum		98.79	0.48	MAN 1	99	99	1	
Megasphaera elsdenii		0.48	0.10	ARA 10	99	1	1	
Propionibacterium granulosum		0.38	0.14		1	99	90	
07103000	**Good identification**							**07103000**
					COC	GRA	CAT	
Gemella morbillorum		92.20	0.60	MAN 1	99	99	1	
Propionibacterium granulosum		7.25	0.48		1	99	90	
Prevotella intermedia		0.17	0.12		1	1	1	
07103002	**Low discrimination**							**07103002**
					COC	CAT		
Propionibacterium granulosum		60.94	0.38	FRU 99	1	90		
Gemella morbillorum		31.30	0.26		99	1		
Actinomyces naeslundii		4.33	0.33		1	25		
07103003	**Low discrimination**							**07103003**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		69.83	0.36	RAF 92	1		
Clostridium chauvoei		30.17	0.07	FRU 95	99		
07103020	**Low discrimination**						**07103020**
					COC	CAT	
Gemella morbillorum		82.47	0.43	MAN 1	99	1	
Propionibacterium granulosum		14.60	0.38	NAG 10	1	90	
Clostridium perfringens		1.33	0.12		1	1	
07103021	**Low discrimination**						**07103021**
					COC	SPO	
Actinomyces naeslundii		42.53	0.36	RAF 92	1	1	
Clostridium chauvoei		34.70	0.11	FRU 95	1	99	
Gemella morbillorum		22.77	0.09		99	1	
07103023	**Good identification**						**07103023**
Actinomyces naeslundii		100.0	0.36	RAF 92			
07103100	**Low discrimination**						**07103100**
					COC	SPO	
Propionibacterium propionicum		60.29	0.46	NIT 86	1	1	
Gemella morbillorum		17.69	0.26	FRU 99	99	1	
Clostridium chauvoei		8.99	0.19		1	99	
07103101	**Good identification**						**07103101**
					SPO		
Clostridium chauvoei		98.11	0.54	MAN 1	99		
Actinomyces naeslundii		1.15	0.44		1		
Propionibacterium propionicum		0.74	0.30		1		
07103102	**Low discrimination**						**07103102**
Propionibacterium propionicum		89.84	0.49	NIT 86			
Actinomyces naeslundii		6.92	0.41	FRU 99			
Lactobacillus fermentum		1.67	0.20				
07103103	**Low discrimination**						**07103103**
					SPO		
Clostridium chauvoei		88.45	0.41	MAN 1	99		
Actinomyces naeslundii		5.89	0.44		1		
Propionibacterium propionicum		5.66	0.33		1		
07103121	**Good identification**						**07103121**
					GRA	SPO	
Clostridium chauvoei		96.18	0.45	MAN 1	99	99	
Actinomyces naeslundii		3.39	0.44		99	1	
Prevotella melaninogenica		0.43	-.01		1	1	
07103140	**Good identification**						**07103140**
Propionibacterium propionicum		98.00	0.46	NIT 86			
Actinomyces naeslundii		2.00	0.28	FRU 99			
07103142	**Good identification**						**07103142**
Propionibacterium propionicum		98.66	0.49	NIT 86			
Actinomyces naeslundii		1.34	0.28	FRU 99			
07103400	**Good identification**						**07103400**
					COC	CAT	
Gemella morbillorum		98.77	0.43	MAN 1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.71	0.14	ARA 10	1	90		
Lactobacillus fermentum		0.52	0.16		1	1		
07104000	**Very good identification**							**07104000**
					COC	GRA		
Gemella morbillorum		99.45	0.82	XYL 10	99	99		
Prevotella disiens		0.36	0.32		1	1		
Actinomyces meyeri		0.09	0.31		1	99		
07104001	**Good identification**							**07104001**
					COC	GRA		
Gemella morbillorum		99.64	0.48	XYL 10	99	99		
Prevotella disiens		0.36	-.03	URE 1	1	1		
07104002	**Good identification**							**07104002**
					COC	GRA		
Gemella morbillorum		98.82	0.48	XYL 10	99	99		
Actinomyces odontolyticus		0.40	0.12	BGL 1	1	99		
Prevotella disiens		0.36	-.03		1	1		
07104010	**Good identification**							**07104010**
					COC	GRA	SPO	
Gemella morbillorum		98.68	0.48	XYL 10	99	99	1	
Clostridium botulinum biovar B		0.97	0.08	MLZ 1	1	99	99	
Prevotella disiens		0.35	-.03		1	1	1	
07104020	**Very good identification**							**07104020**
					COC	SPO		
Gemella morbillorum		99.71	0.66	XYL 10	99	1		
Actinomyces meyeri		0.11	0.16	NAG 10	1	1		
Clostridium botulinum biovar B		0.09	0.08		1	99		
07104040	**Low discrimination**							**07104040**
					COC	SPO		
Clostridium botulinum biovar B		65.41	0.47	XYL 1	1	99		
Gemella morbillorum		33.27	0.48		99	1		
Clostridium glycolicum		0.70	0.15		1	99		
07104050	**Good identification**							**07104050**
					COC	SPO		
Clostridium botulinum biovar B		98.97	0.42	XYL 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
07104060	**Low discrimination**							**07104060**
					COC	SPO		
Clostridium botulinum biovar B		89.71	0.42	XYL 1	1	99		
Gemella morbillorum		10.29	0.32		99	1		
07104070	**Good identification**							**07104070**
Clostridium botulinum biovar B		100.0	0.37	XYL 1				
07104100	**Low discrimination**							**07104100**
					COC	GRA		
Gemella morbillorum		73.68	0.48	XYL 10	99	99		
Actinomyces meyeri		24.93	0.41	LAC 1	1	99		
Prevotella bivia		0.54	-.03		1	1		
07104200	**Good identification**							**07104200**
					COC	GRA		
Gemella morbillorum		98.38	0.48	XYL 10	99	99		
Actinomyces odontolyticus		1.27	0.20	RHA 1	1	99		
Prevotella disiens		0.35	-.03		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07104400	**Very good identification**						**07104400**
					COC	GRA	
Gemella morbillorum		99.76	0.66	XYL 10	99	99	
Actinomyces meyeri		0.19	0.20	ARA 10	1	99	
Prevotella disiens		0.03	-.03		1	1	
07104420	**Good identification**						**07104420**
					COC		
Gemella morbillorum		99.77	0.50	XYL 10	99		
Actinomyces meyeri		0.24	0.06	ARA 10	1		
07105000	**Very good identification**						**07105000**
					COC		
Gemella morbillorum		99.41	0.78	XYL 10	99		
Lactobacillus fermentum		0.31	0.46		1		
Actinomyces meyeri		0.24	0.34		1		
07105001	**Good identification**						**07105001**
					COC	SPO	
Gemella morbillorum		95.30	0.43	XYL 10	99	1	
Clostridium chauvoei		4.40	0.19	URE 1	1	99	
Lactobacillus fermentum		0.30	0.12		1	1	
07105002	**Low discrimination**						**07105002**
					COC		
Gemella morbillorum		75.77	0.43	XYL 10	99		
Lactobacillus fermentum		23.30	0.46	BGL 1	1		
Actinomyces meyeri		0.55	0.08		1		
07105010	**Good identification**						**07105010**
					COC		
Gemella morbillorum		99.69	0.43	XYL 10	99		
Lactobacillus fermentum		0.31	0.12	MLZ 1	1		
07105020	**Very good identification**						**07105020**
					COC		
Gemella morbillorum		98.59	0.61	XYL 10	99		
Lactobacillus fermentum		0.92	0.38	NAG 10	1		
Actinomyces meyeri		0.29	0.19		1		
07105040	**Good identification**						**07105040**
					COC	SPO	
Gemella morbillorum		95.76	0.43	XYL 10	99	1	
Clostridium botulinum biovar B		3.53	0.13	SOR 1	1	99	
Clostridium glycolicum		0.41	-.01		1	99	
07105100	**Low discrimination**						**07105100**
					COC		
Gemella morbillorum		39.29	0.43	XYL 10	99		
Actinomyces meyeri		37.03	0.44	LAC 1	1		
Lactobacillus fermentum		19.71	0.50		1		
07105101	**Good identification**						**07105101**
					COC	SPO	
Clostridium chauvoei		99.46	0.54	XYL 1	1	99	
Gemella morbillorum		0.22	0.09		99	1	
Actinomyces meyeri		0.21	0.10		1	1	
07105102	**Low discrimination**						**07105102**
					COC		
Lactobacillus fermentum		89.61	0.50	FRU 99	1		
Actinomyces meyeri		5.21	0.18		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		1.80	0.09		99		
07105103	**Good identification**						**07105103**
					SPO		
Clostridium chauvoei		99.38	0.41	XYL 1	99		
Lactobacillus fermentum		0.62	0.16		1		
07105121	**Good identification**						**07105121**
					GRA	SPO	
Clostridium chauvoei		99.55	0.45	XYL 1	99	99	
Prevotella melaninogenica		0.45	-.01		1	1	
07105122	**Low discrimination**						**07105122**
					GRA		
Lactobacillus fermentum		41.21	0.42	FRU 99	99		
Prevotella melaninogenica		31.81	0.21		1		
Clostridium perfringens		13.70	0.22		99		
07105200	**Good identification**						**07105200**
					COC		
Gemella morbillorum		98.13	0.43	XYL 10	99		
Actinomyces odontolyticus		1.56	0.17	RHA 1	1		
Lactobacillus fermentum		0.31	0.12		1		
07105400	**Very good identification**						**07105400**
					COC		
Gemella morbillorum		96.73	0.61	XYL 10	99		
Lactobacillus fermentum		2.70	0.46	ARA 10	1		
Actinomyces meyeri		0.52	0.23		1		
07105402	**Low discrimination**						**07105402**
					COC		
Lactobacillus fermentum		72.51	0.46	FRU 99	1		
Gemella morbillorum		26.20	0.27		99		
Bifidobacterium longum subsp. longum		1.29	0.09		1		
07105420	**Acceptable identification**						**07105420**
					COC		
Gemella morbillorum		90.23	0.45	XYL 10	99		
Lactobacillus fermentum		7.57	0.38	ARA 10	1		
Bifidobacterium longum subsp. longum		1.61	0.20	NAG 10	1		
07105422	**Low discrimination**						**07105422**
					COC		
Lactobacillus fermentum		75.01	0.38	FRU 99	1		
Bifidobacterium longum subsp. longum		15.96	0.20		1		
Gemella morbillorum		9.03	0.11		99		
07105500	**Low discrimination**						**07105500**
Bifidobacterium longum subsp. longum		38.88	0.44	RAF 97			
Lactobacillus fermentum		36.14	0.50	FRU 90			
Actinomyces meyeri		16.97	0.34				
07105502	**Low discrimination**						**07105502**
Bifidobacterium longum subsp. longum		50.84	0.44	RAF 97			
Lactobacillus fermentum		47.26	0.50	FRU 90			
Bifidobacterium adolescentis		1.21	0.17				
07105510	**Good identification**						**07105510**
Bifidobacterium longum subsp. longum		99.45	0.48	RAF 97			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Lactobacillus fermentum	0.55	0.16	FRU 90				
07105512	**Good identification**							**07105512**
	Bifidobacterium longum subsp. longum	97.79	0.48	RAF 97				
	Bifidobacterium adolescentis	1.67	0.19	FRU 90				
	Lactobacillus fermentum	0.54	0.16					
07105520	**Low discrimination**							**07105520**
					GRA			
	Bifidobacterium longum subsp. longum	86.86	0.54	RAF 97	99			
	Lactobacillus fermentum	6.73	0.42	FRU 90	99			
	Bacteroides vulgatus	4.35	0.17		1			
07105522	**Acceptable identification**							**07105522**
	Bifidobacterium longum subsp. longum	92.81	0.54	RAF 97				
	Lactobacillus fermentum	7.19	0.42	FRU 90				
07105530	**Very good identification**							**07105530**
	Bifidobacterium longum subsp. longum	100.0	0.58	RAF 97				
07105532	**Very good identification**							**07105532**
	Bifidobacterium longum subsp. longum	100.0	0.58	RAF 97				
07106000	**Good identification**							**07106000**
					COC	GRA	CAT	
	Gemella morbillorum	98.79	0.48	MAN 1	99	99	1	
	Megasphaera elsdenii	0.48	0.10	XYL 10	99	1	1	
	Propionibacterium granulosum	0.38	0.14		1	99	90	
07107000	**Good identification**							**07107000**
					COC	CAT		
	Gemella morbillorum	98.99	0.43	MAN 1	99	1		
	Propionibacterium granulosum	0.71	0.14	XYL 10	1	90		
	Lactobacillus fermentum	0.31	0.12		1	1		
07110000	**Low discrimination**							**07110000**
					COC	GRA	CAT	
	Gemella morbillorum	84.87	1.00		99	99	1	
	Prevotella intermedia	6.89	0.80		1	1	1	
	Propionibacterium granulosum	3.90	0.84		1	99	90	
07110001	**Low discrimination**							**07110001**
					COC	GRA	CAT	
	Gemella morbillorum	84.37	0.66	URE 1	99	99	1	
	Prevotella intermedia	6.85	0.46		1	1	1	
	Propionibacterium granulosum	3.88	0.50		1	99	90	
07110002	**Low discrimination**							**07110002**
					COC	GRA	CAT	
	Propionibacterium granulosum	46.23	0.74		1	99	90	
	Gemella morbillorum	40.65	0.66		99	99	1	
	Megasphaera elsdenii	3.73	0.50		99	1	1	
07110010	**Low discrimination**							**07110010**
					COC	SPO	CAT	
	Gemella morbillorum	47.65	0.66	MLZ 1	99	1	1	
	Propionibacterium granulosum	29.56	0.69		1	1	90	
	Clostridium botulinum biovar B	16.78	0.52		1	99	1	
07110012	**Good identification**							**07110012**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO	CAT	
Propionibacterium granulosum		90.20	0.59	MLZ 12	1	1	90	
Gemella morbillorum		5.88	0.32		99	1	1	
Clostridium botulinum biovar B		2.07	0.18		1	99	1	
07110020	**Low discrimination**							**07110020**
					COC	SPO	CAT	
Gemella morbillorum		86.36	0.84	NAG 10	99	1	1	
Propionibacterium granulosum		8.93	0.74		1	1	90	
Clostridium botulinum biovar B		2.77	0.52		1	99	1	
07110021	**Low discrimination**							**07110021**
					COC	SPO	CAT	
Gemella morbillorum		85.32	0.50	NAG 10	99	1	1	
Propionibacterium granulosum		8.82	0.40	URE 1	1	1	90	
Clostridium botulinum biovar B		2.73	0.18		1	99	1	
07110022	**Low discrimination**							**07110022**
					COC	CAT		
Propionibacterium granulosum		62.98	0.64		1	90		
Gemella morbillorum		24.62	0.50		99	1		
Clostridium perfringens		8.14	0.41		1	1		
07110030	**Low discrimination**							**07110030**
					COC	SPO	CAT	
Clostridium botulinum biovar B		39.28	0.47	SOR 99	1	99	1	
Propionibacterium granulosum		35.12	0.59		1	1	90	
Gemella morbillorum		25.16	0.50		99	1	1	
07110032	**Acceptable identification**							**07110032**
					COC	SPO	CAT	
Propionibacterium granulosum		92.28	0.49	MLZ 12	1	1	90	
Clostridium botulinum biovar B		4.17	0.13		1	99	1	
Gemella morbillorum		2.67	0.15		99	1	1	
07110040	**Very good identification**							**07110040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		97.63	0.92		1	99	99	
Gemella morbillorum		1.38	0.66		99	99	1	
Megasphaera elsdenii		0.76	0.63		99	1	1	
07110041	**Very good identification**							**07110041**
					COC	GRA	SPO	
Clostridium botulinum biovar B		97.61	0.58	URE 1	1	99	99	
Gemella morbillorum		1.38	0.32		99	99	1	
Megasphaera elsdenii		0.76	0.29		99	1	1	
07110042	**Low discrimination**							**07110042**
					COC	SPO	CAT	
Clostridium botulinum biovar B		89.35	0.58	BGL 1	1	99	1	
Megasphaera elsdenii		7.64	0.47		99	1	1	
Propionibacterium granulosum		1.44	0.40		1	1	90	
07110050	**Very good identification**							**07110050**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.93	0.87		1	99	1	
Gemella morbillorum		0.03	0.32		99	1	1	
Propionibacterium granulosum		0.02	0.35		1	1	90	
07110051	**Good identification**							**07110051**
Clostridium botulinum biovar B		99.94	0.52	URE 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		0.06	0.12					
07110052	**Good identification**							**07110052**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.38	0.52	BGL 1	1	99	1	
Propionibacterium granulosum		0.44	0.25		1	1	90	
Megasphaera elsdenii		0.17	0.12		99	1	1	
07110060	**Very good identification**							**07110060**
					COC	SPO		
Clostridium botulinum biovar B		99.59	0.87		1	99		
Gemella morbillorum		0.32	0.50		99	1		
Clostridium perfringens		0.04	0.34		1	84		
07110061	**Good identification**							**07110061**
					COC	SPO		
Clostridium botulinum biovar B		99.68	0.52	URE 1	1	99		
Gemella morbillorum		0.32	0.15		99	1		
07110062	**Good identification**							**07110062**
					CAT	SPO		
Clostridium botulinum biovar B		96.81	0.52	BGL 1	1	99		
Clostridium perfringens		1.92	0.29		1	84		
Propionibacterium granulosum		0.79	0.29		90	1		
07110070	**Very good identification**							**07110070**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.98	0.81		1	99	1	
Propionibacterium granulosum		0.01	0.25		1	1	90	
Gemella morbillorum		0.01	0.15		99	1	1	
07110071	**Good identification**							**07110071**
Clostridium botulinum biovar B		100.0	0.47	URE 1				
07110072	**Good identification**							**07110072**
					SPO	CAT		
Clostridium botulinum biovar B		99.78	0.47	BGL 1	99	1		
Propionibacterium granulosum		0.23	0.15		1	90		
07110100	**Low discrimination**							**07110100**
					COC	GRA		
Gemella morbillorum		51.64	0.66	LAC 1	99	99		
Fusobacterium mortiferum		18.74	0.69		1	1		
Actinomyces meyeri		15.55	0.58		1	99		
07110101	**Very good identification**							**07110101**
					COC	GRA	SPO	
Clostridium chauvoei		95.94	0.60	GAL 99	1	99	99	
Gemella morbillorum		1.86	0.32		99	99	1	
Fusobacterium mortiferum		0.67	0.34		1	1	1	
07110102	**Low discrimination**							**07110102**
					GRA			
Actinomyces viscosus		36.52	0.45	NIT 99	99			
Lactobacillus fermentum		17.54	0.54	RAF 94	99			
Fusobacterium mortiferum		16.92	0.52		1			
07110103	**Low discrimination**							**07110103**
					SPO			
Clostridium chauvoei		73.37	0.47	GAL 99	99			
Actinomyces viscosus		23.11	0.37		1			
Propionibacterium propionicum		2.01	0.33		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07110120	**Low discrimination**							**07110120**
					GRA			
Prevotella bivia		40.76	0.50	SUC 1	1			
Clostridium perfringens		31.97	0.61	FRU 10	99			
Prevotella melaninogenica		16.67	0.50		1			
07110121	**Acceptable identification**							**07110121**
					GRA	SPO		
Clostridium chauvoei		90.26	0.52	GAL 99	99	99		
Prevotella bivia		3.82	0.15		1	1		
Clostridium perfringens		3.00	0.27		99	84		
07110122	**Low discrimination**							**07110122**
					GRA			
Clostridium perfringens		74.57	0.56	GAL 99	99			
Prevotella melaninogenica		15.74	0.37		1			
Lactobacillus fermentum		3.86	0.46		99			
07110123	**Low discrimination**							**07110123**
					SPO			
Clostridium chauvoei		86.83	0.39	GAL 99	99			
Clostridium perfringens		7.69	0.22		84			
Actinomyces naeslundii		3.03	0.37		1			
07110140	**Good identification**							**07110140**
					COC	SPO		
Clostridium botulinum biovar B		94.34	0.58	LAC 1	1	99		
Propionibacterium propionicum		2.00	0.46		1	1		
Gemella morbillorum		1.33	0.32		99	1		
07110142	**Low discrimination**							**07110142**
					SPO			
Propionibacterium propionicum		67.14	0.49	NIT 86	1			
Clostridium botulinum biovar B		21.31	0.24	MAN 99	99			
Clostridium perfringens		4.68	0.18		84			
07110150	**Good identification**							**07110150**
					SPO			
Clostridium botulinum biovar B		99.96	0.52	LAC 1	99			
Propionibacterium propionicum		0.04	0.12		1			
07110160	**Low discrimination**							**07110160**
					GRA	SPO		
Clostridium botulinum biovar B		74.41	0.52	LAC 1	99	99		
Clostridium perfringens		23.04	0.49		99	84		
Prevotella bivia		1.56	0.15		1	1		
07110162	**Acceptable identification**							**07110162**
					GRA	SPO		
Clostridium perfringens		92.17	0.43	GAL 99	99	84		
Clostridium botulinum biovar B		6.39	0.18		99	99		
Prevotella melaninogenica		1.03	0.03		1	1		
07110170	**Good identification**							**07110170**
Clostridium botulinum biovar B		99.37	0.47	LAC 1				
Clostridium perfringens		0.63	0.15					
07110200	**Low discrimination**							**07110200**
					COC	GRA	CAT	
Gemella morbillorum		76.93	0.66	RHA 1	99	99	1	
Propionibacterium granulosum		10.82	0.58		1	99	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		6.25	0.46		1	1	1	
07110202	**Low discrimination**							**07110202**
					COC	CAT		
Propionibacterium granulosum		54.09	0.48	RHA 3	1	90		
Actinomyces odontolyticus		25.75	0.40		1	10		
Gemella morbillorum		15.54	0.32		99	1		
07110210	**Low discrimination**							**07110210**
					COC	SPO	CAT	
Propionibacterium granulosum		55.99	0.43	RHA 3	1	1	90	
Gemella morbillorum		29.48	0.32	MLZ 12	99	1	1	
Clostridium botulinum biovar B		10.38	0.18		1	99	1	
07110220	**Low discrimination**							**07110220**
					COC	CAT		
Gemella morbillorum		71.13	0.50	RHA 1	99	1		
Propionibacterium granulosum		22.51	0.48	NAG 10	1	90		
Actinomyces odontolyticus		2.85	0.30		1	10		
07110222	**Low discrimination**							**07110222**
					COC	CAT		
Propionibacterium granulosum		67.57	0.38	RHA 3	1	90		
Actinomyces odontolyticus		20.95	0.27		1	10		
Gemella morbillorum		8.63	0.15		99	1		
07110240	**Very good identification**							**07110240**
					COC	GRA	SPO	
Clostridium botulinum biovar B		97.48	0.58	RHA 1	1	99	99	
Gemella morbillorum		1.38	0.32		99	99	1	
Megasphaera elsdenii		0.76	0.29		99	1	1	
07110250	**Good identification**							**07110250**
					SPO	CAT		
Clostridium botulinum biovar B		99.95	0.52	RHA 1	99	1		
Propionibacterium granulosum		0.06	0.09		1	90		
07110260	**Good identification**							**07110260**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.58	0.52	RHA 1	1	99	1	
Gemella morbillorum		0.32	0.15		99	1	1	
Propionibacterium granulosum		0.10	0.14		1	1	90	
07110270	**Good identification**							**07110270**
Clostridium botulinum biovar B		100.0	0.47	RHA 1				
07110400	**Very good identification**							**07110400**
					COC	GRA	CAT	
Gemella morbillorum		98.03	0.84	ARA 10	99	99	1	
Prevotella intermedia		0.72	0.46		1	1	1	
Propionibacterium granulosum		0.41	0.50		1	99	90	
07110401	**Good identification**							**07110401**
					COC	GRA	CAT	
Gemella morbillorum		98.18	0.50	ARA 10	99	99	1	
Prevotella intermedia		0.73	0.12	URE 1	1	1	1	
Propionibacterium granulosum		0.41	0.16		1	99	90	
07110402	**Low discrimination**							**07110402**
					COC	CAT		
Gemella morbillorum		73.32	0.50	ARA 10	99	1		
Lactobacillus fermentum		16.91	0.50	BGL 1	1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		7.58	0.40		1	90		
07110410	**Acceptable identification**							**07110410**
					COC	SPO	CAT	
Gemella morbillorum		90.68	0.50	ARA 10	99	1	1	
Propionibacterium granulosum		5.11	0.35	MLZ 1	1	1	90	
Clostridium botulinum biovar B		2.90	0.18		1	99	1	
07110420	**Very good identification**							**07110420**
					COC	CAT		
Gemella morbillorum		97.75	0.67	ARA 10	99	1		
Propionibacterium granulosum		0.92	0.40	NAG 10	1	90		
Lactobacillus fermentum		0.68	0.42		1	1		
07110440	**Low discrimination**							**07110440**
					COC	GRA	SPO	
Clostridium botulinum biovar B		85.76	0.58	ARA 1	1	99	99	
Gemella morbillorum		13.33	0.50		99	99	1	
Megasphaera elsdenii		0.67	0.29		99	1	1	
07110450	**Good identification**							**07110450**
					COC	SPO		
Clostridium botulinum biovar B		99.68	0.52	ARA 1	1	99		
Gemella morbillorum		0.32	0.15		99	1		
07110460	**Good identification**							**07110460**
					COC	SPO		
Clostridium botulinum biovar B		96.61	0.52	ARA 1	1	99		
Gemella morbillorum		3.39	0.33		99	1		
07110470	**Good identification**							**07110470**
Clostridium botulinum biovar B		100.0	0.47	ARA 1				
07110500	**Low discrimination**							**07110500**
					COC			
Gemella morbillorum		47.36	0.50	LAC 1	99			
Actinomyces meyeri		32.08	0.48	ARA 10	1			
Lactobacillus fermentum		17.83	0.54		1			
07110502	**Low discrimination**							**07110502**
					COC			
Lactobacillus fermentum		87.88	0.54	GAL 99	1			
Actinomyces meyeri		4.89	0.22		1			
Gemella morbillorum		2.36	0.15		99			
07110520	**Low discrimination**							**07110520**
					COC			
Lactobacillus fermentum		24.93	0.46	GAL 99	1			
Gemella morbillorum		22.08	0.33		99			
Actinomyces meyeri		18.35	0.33		1			
07110522	**Low discrimination**							**07110522**
Lactobacillus fermentum		66.48	0.46	GAL 99				
Bifidobacterium longum subsp. longum		15.27	0.28					
Clostridium perfringens		12.98	0.22					
07110600	**Good identification**							**07110600**
					COC	GRA	CAT	
Gemella morbillorum		97.12	0.50	RHA 1	99	99	1	
Propionibacterium granulosum		1.24	0.24	ARA 10	1	99	90	
Prevotella intermedia		0.72	0.12		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07111000	**Low discrimination**							**07111000**
					COC	GRA	CAT	
Gemella morbillorum		77.05	0.95		99	99	1	
Prevotella intermedia		11.62	0.80		1	1	1	
Propionibacterium granulosum		6.57	0.84		1	99	90	
07111001	**Low discrimination**							**07111001**
					COC	GRA	SPO	
Gemella morbillorum		43.79	0.61	URE 1	99	99	1	
Clostridium chauvoei		42.46	0.60		1	99	99	
Prevotella intermedia		6.61	0.46		1	1	1	
07111002	**Low discrimination**							**07111002**
					COC	CAT		
Lactobacillus fermentum		55.78	0.85		1	1		
Propionibacterium granulosum		24.99	0.74		1	90		
Gemella morbillorum		11.84	0.61		99	1		
07111003	**Low discrimination**							**07111003**
					SPO	CAT		
Clostridium chauvoei		58.86	0.47	LAC 99	99	1		
Lactobacillus fermentum		16.37	0.50		1	1		
Propionibacterium granulosum		7.34	0.40		1	90		
07111010	**Low discrimination**							**07111010**
					COC	GRA	CAT	
Propionibacterium granulosum		47.52	0.69	MLZ 12	1	99	90	
Gemella morbillorum		41.26	0.61		99	99	1	
Prevotella intermedia		6.22	0.46		1	1	1	
07111011	**Low discrimination**							**07111011**
					COC	SPO	CAT	
Propionibacterium granulosum		33.27	0.35	MLZ 12	1	1	90	
Gemella morbillorum		28.89	0.27	URE 1	99	1	1	
Clostridium chauvoei		28.01	0.26		1	99	1	
07111012	**Low discrimination**							**07111012**
					CAT			
Propionibacterium granulosum		74.38	0.59	MLZ 12	90			
Lactobacillus fermentum		12.30	0.50		1			
Lactobacillus jensenii		9.35	0.37		1			
07111020	**Low discrimination**							**07111020**
					COC	CAT		
Clostridium perfringens		48.08	0.81	LAC 88	1	1		
Gemella morbillorum		36.85	0.79		99	1		
Propionibacterium granulosum		7.08	0.74		1	90		
07111021	**Low discrimination**							**07111021**
					COC	SPO		
Clostridium chauvoei		50.42	0.52	LAC 99	1	99		
Clostridium perfringens		22.61	0.46		1	84		
Gemella morbillorum		17.33	0.45		99	1		
07111022	**Low discrimination**							**07111022**
					CAT			
Clostridium perfringens		73.65	0.75	LAC 88	1			
Lactobacillus fermentum		17.13	0.76		1			
Propionibacterium granulosum		5.76	0.64		90			
07111023	**Low discrimination**							**07111023**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		41.12	0.41	LAC 88	84			
Clostridium chauvoei		34.38	0.39	URE 1	99			
Actinomyces naeslundii		10.12	0.53		1			
07111030	**Low discrimination**							**07111030**
					COC	CAT		
Propionibacterium granulosum		49.25	0.59	MLZ 12	1	90		
Clostridium perfringens		24.79	0.46		1	1		
Gemella morbillorum		19.00	0.45		99	1		
07111032	**Low discrimination**							**07111032**
					CAT			
Propionibacterium granulosum		42.75	0.49	MLZ 12	90			
Clostridium perfringens		40.52	0.41		1			
Lactobacillus fermentum		9.42	0.42		1			
07111040	**Low discrimination**							**07111040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		48.86	0.58	GAL 1	1	99	99	
Gemella morbillorum		36.80	0.61		99	99	1	
Prevotella intermedia		5.55	0.46		1	1	1	
07111042	**Low discrimination**							**07111042**
					CAT			
Lactobacillus fermentum		34.13	0.50	SOR 1	1			
Clostridium perfringens		28.53	0.37		1			
Propionibacterium granulosum		15.30	0.40		90			
07111043	**Low discrimination**							**07111043**
					SPO			
Actinomyces naeslundii		53.15	0.40	RAF 92	1			
Clostridium chauvoei		31.01	0.13		99			
Lactobacillus fermentum		8.63	0.16		1			
07111050	**Good identification**							**07111050**
					COC	SPO	CAT	
Clostridium botulinum biovar B		96.30	0.52	GAL 1	1	99	1	
Propionibacterium granulosum		1.71	0.35		1	1	90	
Gemella morbillorum		1.49	0.27		99	1	1	
07111060	**Low discrimination**							**07111060**
					COC	SPO		
Clostridium perfringens		77.10	0.68	LAC 88	1	84		
Clostridium botulinum biovar B		18.44	0.52		1	99		
Gemella morbillorum		3.13	0.45		99	1		
07111062	**Very good identification**							**07111062**
					SPO			
Clostridium perfringens		97.54	0.63	LAC 88	84			
Lactobacillus fermentum		1.20	0.42		1			
Clostridium botulinum biovar B		0.50	0.18		99			
07111070	**Acceptable identification**							**07111070**
					CAT	SPO		
Clostridium botulinum biovar B		91.06	0.47	GAL 1	1	99		
Clostridium perfringens		7.81	0.34		1	84		
Propionibacterium granulosum		0.82	0.25		90	1		
07111100	**Low discrimination**							**07111100**
					GRA			
Lactobacillus fermentum		48.66	0.88		99			
Clostridium perfringens		20.60	0.70		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		6.40	0.72		1		
07111101	**Excellent identification**						**07111101**
					SPO		
Clostridium chauvoei		99.79	0.94		99		
Lactobacillus fermentum		0.08	0.54		1		
Actinomyces naeslundii		0.05	0.61		1		
07111102	**Low discrimination**						**07111102**
					SPO		
Lactobacillus fermentum		77.12	0.88		1		
Clostridium perfringens		15.37	0.64		84		
Clostridium chauvoei		1.72	0.47		99		
07111103	**Very good identification**						**07111103**
					SPO		
Clostridium chauvoei		98.94	0.81		99		
Lactobacillus fermentum		0.45	0.54		1		
Actinomyces naeslundii		0.28	0.61		1		
07111110	**Low discrimination**						**07111110**
					CAT		
Lactobacillus fermentum		44.15	0.54	MLZ 1	1		
Clostridium perfringens		18.69	0.35		1		
Propionibacterium granulosum		6.61	0.35		90		
07111111	**Very good identification**						**07111111**
					SPO		
Clostridium chauvoei		99.63	0.60	MLZ 1	99		
Actinomyces naeslundii		0.26	0.39		1		
Lactobacillus fermentum		0.08	0.20		1		
07111112	**Low discrimination**						**07111112**
Lactobacillus fermentum		69.18	0.54	MLZ 1			
Clostridium perfringens		13.78	0.30				
Bifidobacterium adolescentis		2.81	0.25				
07111113	**Good identification**						**07111113**
					SPO		
Clostridium chauvoei		97.92	0.47	MLZ 1	99		
Actinomyces naeslundii		1.44	0.39		1		
Lactobacillus fermentum		0.45	0.20		1		
07111120	**Low discrimination**						**07111120**
					GRA		
Clostridium perfringens		64.29	0.95		99		
Prevotella melaninogenica		33.53	0.84		1		
Lactobacillus fermentum		1.57	0.80		99		
07111121	**Good identification**						**07111121**
					GRA	SPO	
Clostridium chauvoei		94.96	0.86		99	99	
Clostridium perfringens		3.16	0.61		99	84	
Prevotella melaninogenica		1.65	0.50		1	1	
07111122	**Low discrimination**						**07111122**
					GRA		
Clostridium perfringens		77.80	0.90		99		
Prevotella melaninogenica		16.42	0.71		1		
Lactobacillus fermentum		4.03	0.80		99		
07111123	**Low discrimination**						**07111123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium chauvoei		89.12	0.73		99	99	
Clostridium perfringens		7.90	0.56		99	84	
Prevotella melaninogenica		1.67	0.37		1	1	
07111130	**Low discrimination**						**07111130**
					GRA		
Clostridium perfringens		63.67	0.61	MLZ 1	99		
Prevotella melaninogenica		33.20	0.50		1		
Lactobacillus fermentum		1.55	0.46		99		
07111131	**Acceptable identification**						**07111131**
					GRA	SPO	
Clostridium chauvoei		94.49	0.52	MLZ 1	99	99	
Clostridium perfringens		3.14	0.27		99	84	
Prevotella melaninogenica		1.64	0.15		1	1	
07111132	**Low discrimination**						**07111132**
					GRA		
Clostridium perfringens		73.78	0.56	MLZ 1	99		
Prevotella melaninogenica		15.57	0.37		1		
Lactobacillus fermentum		3.82	0.46		99		
07111133	**Low discrimination**						**07111133**
					SPO		
Clostridium chauvoei		86.85	0.39	MLZ 1	99		
Clostridium perfringens		7.70	0.22		84		
Actinomyces naeslundii		3.83	0.39		1		
07111140	**Low discrimination**						**07111140**
Clostridium perfringens		78.16	0.57	NAG 97			
Lactobacillus fermentum		9.79	0.54				
Propionibacterium propionicum		3.55	0.46				
07111141	**Very good identification**						**07111141**
					SPO		
Clostridium chauvoei		98.12	0.60	SOR 1	99		
Actinomyces naeslundii		0.85	0.48		1		
Clostridium perfringens		0.63	0.23		84		
07111142	**Low discrimination**						**07111142**
Clostridium perfringens		68.38	0.52	NAG 97			
Lactobacillus fermentum		18.20	0.54				
Propionibacterium propionicum		9.90	0.49				
07111143	**Acceptable identification**						**07111143**
					SPO		
Clostridium chauvoei		90.87	0.47	SOR 1	99		
Actinomyces naeslundii		4.48	0.48		1		
Propionibacterium propionicum		2.49	0.33		1		
07111160	**Very good identification**						**07111160**
					GRA		
Clostridium perfringens		97.07	0.83		99		
Prevotella melaninogenica		2.69	0.50		1		
Lactobacillus fermentum		0.13	0.46		99		
07111161	**Low discrimination**						**07111161**
					SPO		
Clostridium chauvoei		59.88	0.52	SOR 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		37.52	0.49		84			
Actinomyces naeslundii		1.56	0.48		1			
07111162	**Very good identification**							**07111162**
					GRA			
Clostridium perfringens		98.21	0.78		99			
Prevotella melaninogenica		1.10	0.37		1			
Lactobacillus fermentum		0.27	0.46		99			
07111163	**Low discrimination**							**07111163**
					SPO			
Clostridium perfringens		58.89	0.43	URE 1	84			
Clostridium chauvoei		35.25	0.39		99			
Actinomyces naeslundii		5.21	0.48		1			
07111170	**Good identification**							**07111170**
					GRA	SPO		
Clostridium perfringens		95.54	0.49	MLZ 1	99	84		
Prevotella melaninogenica		2.64	0.15		1	1		
Clostridium botulinum biovar B		1.54	0.13		99	99		
07111172	**Good identification**							**07111172**
					GRA	CAT		
Clostridium perfringens		95.87	0.43	MLZ 1	99	1		
Capnocytophaga ochracea		2.76	0.14		1	90		
Prevotella melaninogenica		1.07	0.03		1	1		
07111200	**Low discrimination**							**07111200**
					COC	GRA	CAT	
Gemella morbillorum		66.37	0.61	RHA 1	99	99	1	
Propionibacterium granulosum		17.34	0.58		1	99	90	
Prevotella intermedia		10.01	0.46		1	1	1	
07111202	**Low discrimination**							**07111202**
					CAT			
Propionibacterium granulosum		43.37	0.48	RHA 3	90			
Lactobacillus fermentum		31.61	0.50		1			
Actinomyces odontolyticus		13.77	0.37		10			
07111210	**Low discrimination**							**07111210**
					COC	GRA	CAT	
Propionibacterium granulosum		73.27	0.43	RHA 3	1	99	90	
Gemella morbillorum		20.78	0.27	MLZ 12	99	99	1	
Prevotella intermedia		3.13	0.12		1	1	1	
07111220	**Low discrimination**							**07111220**
					COC	CAT		
Clostridium perfringens		41.23	0.46	LAC 88	1	1		
Gemella morbillorum		31.59	0.45	RHA 1	99	1		
Propionibacterium granulosum		18.57	0.48		1	90		
07111222	**Low discrimination**							**07111222**
					GRA	CAT		
Clostridium perfringens		53.93	0.41	LAC 88	99	1		
Prevotella oralis		16.23	0.29	RHA 1	1	1		
Propionibacterium granulosum		12.91	0.38		99	90		
07111300	**Low discrimination**							**07111300**
Lactobacillus fermentum		45.01	0.54	RHA 1				
Clostridium perfringens		19.06	0.35					
Actinomyces odontolyticus		6.19	0.32					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07111301	**Very good identification**						**07111301**
					SPO		
Clostridium chauvoei		99.84	0.60	RHA 1	99		
Lactobacillus fermentum		0.08	0.20		1		
Actinomyces naeslundii		0.05	0.26		1		
07111302	**Low discrimination**						**07111302**
					GRA		
Lactobacillus fermentum		62.86	0.54	RHA 1	99		
Prevotella oralis		13.70	0.27		1		
Clostridium perfringens		12.52	0.30		99		
07111303	**Good identification**						**07111303**
					SPO		
Clostridium chauvoei		99.07	0.47	RHA 1	99		
Lactobacillus fermentum		0.45	0.20		1		
Actinomyces naeslundii		0.28	0.26		1		
07111320	**Low discrimination**						**07111320**
					GRA		
Clostridium perfringens		63.09	0.61	RHA 1	99		
Prevotella melaninogenica		32.90	0.50		1		
Lactobacillus fermentum		1.54	0.46		99		
07111321	**Good identification**						**07111321**
					GRA	SPO	
Clostridium chauvoei		95.05	0.52	RHA 1	99	99	
Clostridium perfringens		3.16	0.27		99	84	
Prevotella melaninogenica		1.65	0.15		1	1	
07111322	**Low discrimination**						**07111322**
					GRA		
Clostridium perfringens		51.89	0.56	RHA 1	99		
Prevotella oralis		33.36	0.49		1		
Prevotella melaninogenica		10.95	0.37		1		
07111323	**Low discrimination**						**07111323**
					GRA	SPO	
Clostridium chauvoei		85.27	0.39	RHA 1	99	99	
Clostridium perfringens		7.56	0.22		99	84	
Prevotella oralis		4.86	0.15		1	1	
07111360	**Good identification**						**07111360**
					GRA		
Clostridium perfringens		97.31	0.49	RHA 1	99		
Prevotella melaninogenica		2.69	0.15		1		
07111362	**Good identification**						**07111362**
					GRA		
Clostridium perfringens		95.67	0.43	RHA 1	99		
Prevotella oralis		3.26	0.15		1		
Prevotella melaninogenica		1.07	0.03		1		
07111400	**Low discrimination**						**07111400**
					COC	GRA	
Gemella morbillorum		68.69	0.79	ARA 10	99	99	
Lactobacillus fermentum		29.43	0.85		1	99	
Prevotella intermedia		0.94	0.46		1	1	
07111401	**Low discrimination**						**07111401**
					COC	SPO	
Gemella morbillorum		64.82	0.45	ARA 10	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		27.77	0.50	URE 1	1	1	
Clostridium chauvoei		5.71	0.26		1	99	
07111402	**Very good identification**						**07111402**
					COC	CAT	
Lactobacillus fermentum		97.09	0.85		1	1	
Gemella morbillorum		2.29	0.45		99	1	
Propionibacterium granulosum		0.44	0.40		1	90	
07111403	**Acceptable identification**						**07111403**
					COC	SPO	
Lactobacillus fermentum		94.35	0.50	URE 1	1	1	
Clostridium chauvoei		3.43	0.13		1	99	
Gemella morbillorum		2.23	0.11		99	1	
07111410	**Low discrimination**						**07111410**
					COC	CAT	
Gemella morbillorum		63.32	0.45	ARA 10	99	1	
Lactobacillus fermentum		27.13	0.50	MLZ 1	1	1	
Propionibacterium granulosum		6.63	0.35		1	90	
07111412	**Low discrimination**						**07111412**
					CAT		
Lactobacillus fermentum		87.29	0.50	MLZ 1	1		
Propionibacterium granulosum		5.33	0.25		90		
Bifidobacterium longum subsp. longum		4.64	0.22		1		
07111420	**Low discrimination**						**07111420**
					COC		
Lactobacillus fermentum		52.71	0.76		1		
Gemella morbillorum		41.01	0.63		99		
Clostridium perfringens		4.87	0.46		1		
07111421	**Low discrimination**						**07111421**
					COC	SPO	
Lactobacillus fermentum		48.17	0.42	URE 1	1	1	
Gemella morbillorum		37.48	0.29		99	1	
Clostridium chauvoei		9.91	0.18		1	99	
07111422	**Good identification**						**07111422**
					COC		
Lactobacillus fermentum		94.37	0.76		1		
Clostridium perfringens		4.10	0.41		1		
Gemella morbillorum		0.74	0.29		99		
07111423	**Acceptable identification**						**07111423**
					SPO		
Lactobacillus fermentum		92.62	0.42	URE 1	1		
Clostridium perfringens		4.02	0.07		84		
Clostridium chauvoei		3.36	0.05		99		
07111430	**Low discrimination**						**07111430**
					COC		
Lactobacillus fermentum		37.15	0.42	MLZ 1	1		
Gemella morbillorum		28.90	0.29		99		
Bifidobacterium longum subsp. longum		23.71	0.32		1		
07111432	**Low discrimination**						**07111432**
					CAT		
Lactobacillus fermentum		57.89	0.42	MLZ 1	1		
Bifidobacterium longum subsp. longum		36.95	0.32		1		
Propionibacterium granulosum		2.65	0.15		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07111440	**Low discrimination**						**07111440**
					COC	SPO	
Gemella morbillorum		63.17	0.45	ARA 10	99	1	
Lactobacillus fermentum		27.06	0.50	SOR 1	1	1	
Clostridium botulinum biovar B		7.63	0.24		1	99	
07111442	**Good identification**						**07111442**
					COC		
Lactobacillus fermentum		96.90	0.50	SOR 1	1		
Gemella morbillorum		2.29	0.11		99		
Clostridium perfringens		0.82	0.03		1		
07111462	**Low discrimination**						**07111462**
Lactobacillus fermentum		54.97	0.42	SOR 1			
Clostridium perfringens		45.03	0.29				
07111500	**Good identification**						**07111500**
Lactobacillus fermentum		93.54	0.88				
Actinomyces meyeri		2.55	0.51				
Bifidobacterium longum subsp. longum		1.79	0.52				
07111501	**Good identification**						**07111501**
					SPO		
Clostridium chauvoei		92.10	0.60	ARA 1	99		
Lactobacillus fermentum		7.38	0.54		1		
Actinomyces meyeri		0.20	0.16		1		
07111502	**Very good identification**						**07111502**
Lactobacillus fermentum		97.49	0.88				
Bifidobacterium longum subsp. longum		1.87	0.52				
Bifidobacterium adolescentis		0.22	0.37				
07111503	**Low discrimination**						**07111503**
					SPO		
Clostridium chauvoei		68.15	0.47	ARA 1	99		
Lactobacillus fermentum		30.93	0.54		1		
Bifidobacterium longum subsp. longum		0.59	0.18		1		
07111510	**Low discrimination**						**07111510**
Bifidobacterium longum subsp. longum		75.52	0.56	RAF 97			
Lactobacillus fermentum		23.40	0.54				
Actinomyces meyeri		0.64	0.16				
07111512	**Low discrimination**						**07111512**
Bifidobacterium longum subsp. longum		71.64	0.56	RAF 97			
Lactobacillus fermentum		22.20	0.54				
Bifidobacterium adolescentis		6.03	0.39				
07111520	**Low discrimination**						**07111520**
Lactobacillus fermentum		52.94	0.80				
Clostridium perfringens		21.96	0.61				
Bifidobacterium longum subsp. longum		12.16	0.62				
07111521	**Low discrimination**						**07111521**
					SPO		
Clostridium chauvoei		86.85	0.52	ARA 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		6.96	0.46		1		
Clostridium perfringens		2.89	0.27		84		
07111522	**Low discrimination**						**07111522**
Lactobacillus fermentum		67.32	0.80				
Bifidobacterium longum subsp. longum		15.47	0.62				
Clostridium perfringens		13.14	0.56				
07111523	**Low discrimination**						**07111523**
					SPO		
Clostridium chauvoei		59.50	0.39	ARA 1	99		
Lactobacillus fermentum		27.00	0.46		1		
Bifidobacterium longum subsp. longum		6.20	0.28		1		
07111530	**Very good identification**						**07111530**
Bifidobacterium longum subsp. longum		95.89	0.66	RAF 97			
Lactobacillus fermentum		2.48	0.46				
Clostridium perfringens		1.03	0.27				
07111532	**Very good identification**						**07111532**
Bifidobacterium longum subsp. longum		96.44	0.66	RAF 97			
Lactobacillus fermentum		2.49	0.46				
Clostridium perfringens		0.49	0.22				
07111540	**Low discrimination**						**07111540**
Lactobacillus fermentum		87.32	0.54	SOR 1			
Clostridium perfringens		7.04	0.23				
Actinomyces meyeri		2.38	0.16				
07111542	**Acceptable identification**						**07111542**
Lactobacillus fermentum		91.22	0.54	SOR 1			
Clostridium perfringens		3.46	0.18				
Bifidobacterium adolescentis		2.51	0.22				
07111560	**Low discrimination**						**07111560**
Clostridium perfringens		84.13	0.49	ARA 1			
Lactobacillus fermentum		10.75	0.46				
Bifidobacterium longum subsp. longum		2.47	0.28				
07111562	**Low discrimination**						**07111562**
Clostridium perfringens		71.89	0.43	ARA 1			
Lactobacillus fermentum		19.53	0.46				
Bifidobacterium longum subsp. longum		4.49	0.28				
07111600	**Low discrimination**						**07111600**
					COC	CAT	
Gemella morbillorum		67.86	0.45	RHA 1	99	1	
Lactobacillus fermentum		29.07	0.50	ARA 10	1	1	
Propionibacterium granulosum		1.61	0.24		1	90	
07111602	**Good identification**						**07111602**
					COC	CAT	
Lactobacillus fermentum		96.39	0.50	RHA 1	1	1	
Gemella morbillorum		2.27	0.11		99	1	
Propionibacterium granulosum		1.34	0.14		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07111620	**Low discrimination**							**07111620**
					COC	GRA		
Lactobacillus fermentum		49.37	0.42	RHA 1	1	99		
Gemella morbillorum		38.41	0.29		99	99		
Bacteroides vulgatus		5.62	0.05		1	1		
07111622	**Good identification**							**07111622**
Lactobacillus fermentum		95.84	0.42	RHA 1				
Clostridium perfringens		4.16	0.07					
07111700	**Acceptable identification**							**07111700**
					GRA			
Lactobacillus fermentum		91.88	0.54	RHA 1	99			
Actinomyces meyeri		2.51	0.16		99			
Bacteroides vulgatus		2.14	0.05		1			
07111702	**Good identification**							**07111702**
Lactobacillus fermentum		98.12	0.54	RHA 1				
Bifidobacterium longum subsp. longum		1.88	0.18					
07111720	**Low discrimination**							**07111720**
					GRA			
Bacteroides vulgatus		78.67	0.39	RAF 97	1			
Lactobacillus fermentum		11.40	0.46	XYL 98	99			
Clostridium perfringens		4.73	0.27		99			
07111722	**Low discrimination**							**07111722**
Lactobacillus fermentum		55.39	0.46	RHA 1				
Bifidobacterium longum subsp. longum		12.73	0.28					
Clostridium perfringens		10.81	0.22					
07112000	**Low discrimination**							**07112000**
					COC	GRA	CAT	
Propionibacterium granulosum		63.36	0.82		1	99	90	
Gemella morbillorum		18.47	0.66		99	99	1	
Megasphaera elsdenii		15.24	0.66		99	1	1	
07112001	**Low discrimination**							**07112001**
					COC	GRA	CAT	
Propionibacterium granulosum		60.97	0.48	URE 1	1	99	90	
Gemella morbillorum		17.77	0.32		99	99	1	
Megasphaera elsdenii		14.67	0.32		99	1	1	
07112002	**Low discrimination**							**07112002**
					COC	CAT	GRA	
Propionibacterium granulosum		87.08	0.72		1	90	99	
Megasphaera elsdenii		9.31	0.50		99	1	1	
Propionibacterium propionicum		2.31	0.49		1	1	99	
07112003	**Low discrimination**							**07112003**
					COC	CAT	GRA	
Propionibacterium granulosum		66.58	0.38	URE 1	1	90	99	
Propionibacterium propionicum		19.42	0.33		1	1	99	
Megasphaera elsdenii		7.12	0.15		99	1	1	
07112010	**Good identification**							**07112010**
					COC	GRA	CAT	
Propionibacterium granulosum		93.86	0.67	MLZ 12	1	99	90	
Gemella morbillorum		2.03	0.32		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		1.67	0.32		99	1	1	
07112012	**Very good identification**							**07112012**
					COC	CAT	GRA	
Propionibacterium granulosum		99.02	0.57	MLZ 12	1	90	99	
Megasphaera elsdenii		0.78	0.15		99	1	1	
Propionibacterium propionicum		0.20	0.15		1	1	99	
07112020	**Low discrimination**							**07112020**
					COC	GRA	CAT	
Propionibacterium granulosum		87.21	0.72		1	99	90	
Gemella morbillorum		11.30	0.50		99	99	1	
Megasphaera elsdenii		0.85	0.32		99	1	1	
07112021	**Low discrimination**							**07112021**
					COC	CAT		
Propionibacterium granulosum		82.38	0.38	URE 1	1	90		
Gemella morbillorum		10.67	0.15		99	1		
Actinomyces naeslundii		6.95	0.34		1	25		
07112022	**Very good identification**							**07112022**
					COC	GRA	CAT	
Propionibacterium granulosum		98.56	0.61		1	99	90	
Gemella morbillorum		0.52	0.15		99	99	1	
Megasphaera elsdenii		0.43	0.15		99	1	1	
07112030	**Very good identification**							**07112030**
					COC	SPO	CAT	
Propionibacterium granulosum		97.60	0.57	MLZ 12	1	1	90	
Clostridium botulinum biovar B		1.46	0.13		1	99	1	
Gemella morbillorum		0.94	0.15		99	1	1	
07112032	**Good identification**							**07112032**
Propionibacterium granulosum		100.0	0.47	MLZ 12				
07112040	**Low discrimination**							**07112040**
					COC	SPO	CAT	
Clostridium botulinum biovar B		53.90	0.58	MAN 1	1	99	1	
Megasphaera elsdenii		41.48	0.63		99	1	1	
Propionibacterium granulosum		2.61	0.48		1	1	90	
07112042	**Low discrimination**							**07112042**
					COC	CAT	GRA	
Megasphaera elsdenii		61.27	0.47	MNS 1	99	1	1	
Propionibacterium propionicum		22.80	0.49	BGL 10	1	1	99	
Propionibacterium granulosum		8.68	0.38		1	90	99	
07112050	**Good identification**							**07112050**
					COC	SPO	CAT	
Clostridium botulinum biovar B		97.02	0.52	MAN 1	1	99	1	
Megasphaera elsdenii		1.53	0.29		99	1	1	
Propionibacterium granulosum		1.30	0.33		1	1	90	
07112060	**Good identification**							**07112060**
					COC	SPO	CAT	
Clostridium botulinum biovar B		95.79	0.52	MAN 1	1	99	1	
Propionibacterium granulosum		2.36	0.38		1	1	90	
Megasphaera elsdenii		1.51	0.29		99	1	1	
07112070	**Good identification**							**07112070**
					SPO	CAT		
Clostridium botulinum biovar B		99.32	0.47	MAN 1	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.68	0.23		1	90		
07112100	**Very good identification**							**07112100**
					COC	CAT		
Propionibacterium propionicum		95.97	0.81	NIT 86	1	1		
Propionibacterium granulosum		2.22	0.48		1	90		
Gemella morbillorum		0.65	0.32		99	1		
07112101	**Very good identification**							**07112101**
					SPO			
Propionibacterium propionicum		95.59	0.64	NIT 86	1			
Clostridium chauvoei		3.02	0.26	URE 10	99			
Actinomyces naeslundii		1.19	0.42		1			
07112102	**Very good identification**							**07112102**
					CAT			
Propionibacterium propionicum		99.44	0.84	NIT 86	1			
Propionibacterium granulosum		0.38	0.38		90			
Actinomyces naeslundii		0.06	0.39		25			
07112103	**Very good identification**							**07112103**
					SPO			
Propionibacterium propionicum		98.70	0.67	NIT 86	1			
Actinomyces naeslundii		0.82	0.42	URE 10	1			
Clostridium chauvoei		0.37	0.13		99			
07112110	**Low discrimination**							**07112110**
					CAT			
Propionibacterium propionicum		76.24	0.46	NIT 86	1			
Propionibacterium granulosum		23.76	0.33	MLZ 1	90			
07112112	**Good identification**							**07112112**
					CAT			
Propionibacterium propionicum		95.06	0.49	NIT 86	1			
Propionibacterium granulosum		4.94	0.23	MLZ 1	90			
07112120	**Low discrimination**							**07112120**
					GRA	CAT		
Propionibacterium propionicum		35.33	0.46	NIT 86	99	1		
Propionibacterium granulosum		20.19	0.38	NAG 1	99	90		
Prevotella bivia		17.17	0.15		1	1		
07112121	**Low discrimination**							**07112121**
					SPO			
Actinomyces naeslundii		37.67	0.42	RAF 92	1			
Clostridium chauvoei		31.83	0.18	GAL 96	99			
Propionibacterium propionicum		30.51	0.30		1			
07112122	**Low discrimination**							**07112122**
					CAT			
Propionibacterium propionicum		76.86	0.49	NIT 86	1			
Clostridium perfringens		9.19	0.22	NAG 1	1			
Propionibacterium granulosum		7.32	0.27		90			
07112140	**Very good identification**							**07112140**
					COC	SPO	GRA	
Propionibacterium propionicum		99.12	0.81	NIT 86	1	1	99	
Clostridium botulinum biovar B		0.48	0.24		1	99	99	
Megasphaera elsdenii		0.37	0.29		99	1	1	
07112141	**Very good identification**							**07112141**
Propionibacterium propionicum		99.78	0.64	NIT 86				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		0.22	0.29	URE 10				
07112142	**Very good identification**							**07112142**
					COC	GRA		
Propionibacterium propionicum		99.96	0.84	NIT 86	1	99		
Megasphaera elsdenii		0.03	0.12		99	1		
Actinomyces naeslundii		0.01	0.26		1	99		
07112143	**Very good identification**							**07112143**
Propionibacterium propionicum		99.85	0.67	NIT 86				
Actinomyces naeslundii		0.15	0.29	URE 10				
07112150	**Low discrimination**							**07112150**
					SPO			
Propionibacterium propionicum		81.00	0.46	NIT 86	1			
Clostridium botulinum biovar B		19.00	0.18	MLZ 1	99			
07112152	**Good identification**							**07112152**
Propionibacterium propionicum		100.0	0.49	NIT 86				
07112160	**Low discrimination**							**07112160**
					SPO			
Propionibacterium propionicum		75.57	0.46	NIT 86	1			
Clostridium botulinum biovar B		17.72	0.18	NAG 1	99			
Clostridium perfringens		5.49	0.15		84			
07112162	**Good identification**							**07112162**
Propionibacterium propionicum		96.76	0.49	NIT 86				
Clostridium perfringens		2.20	0.09	NAG 1				
Actinomyces naeslundii		1.04	0.26					
07112200	**Low discrimination**							**07112200**
					COC	GRA	CAT	
Propionibacterium granulosum		84.01	0.56	RHA 3	1	99	90	
Gemella morbillorum		8.00	0.32		99	99	1	
Megasphaera elsdenii		6.60	0.32		99	1	1	
07112202	**Good identification**							**07112202**
					COC	CAT	GRA	
Propionibacterium granulosum		95.24	0.46	RHA 3	1	90	99	
Megasphaera elsdenii		3.33	0.15		99	1	1	
Propionibacterium propionicum		0.83	0.15		1	1	99	
07112210	**Good identification**							**07112210**
Propionibacterium granulosum		100.0	0.41	RHA 3				
07112220	**Good identification**							**07112220**
					COC	CAT		
Propionibacterium granulosum		95.94	0.46	RHA 3	1	90		
Gemella morbillorum		4.06	0.15		99	1		
07112222	**Good identification**							**07112222**
Propionibacterium granulosum		100.0	0.36	RHA 3				
07112300	**Acceptable identification**							**07112300**
					CAT			
Propionibacterium propionicum		93.40	0.46	NIT 86	1			
Propionibacterium granulosum		6.60	0.22	RHA 1	90			
07112302	**Good identification**							**07112302**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					CAT			
Propionibacterium propionicum		98.84	0.49	NIT 86	1			
Propionibacterium granulosum		1.16	0.12	RHA 1	90			
07112340	**Good identification**							**07112340**
Propionibacterium propionicum		100.0	0.46	NIT 86				
07112342	**Good identification**							**07112342**
Propionibacterium propionicum		100.0	0.49	NIT 86				
07112400	**Low discrimination**							**07112400**
					COC	GRA	CAT	
Gemella morbillorum		71.21	0.50	MAN 1	99	99	1	
Propionibacterium granulosum		22.21	0.48	ARA 10	1	99	90	
Megasphaera elsdenii		5.34	0.32		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07112402	**Low discrimination**							**07112402**
					COC	GRA	CAT	
Propionibacterium granulosum		77.35	0.38	ARA 1	1	99	90	
Gemella morbillorum		10.02	0.15		99	99	1	
Megasphaera elsdenii		8.27	0.15		99	1	1	
07112500	**Low discrimination**							**07112500**
					COC			
Propionibacterium propionicum		85.10	0.46	NIT 86	1			
Gemella morbillorum		6.30	0.15	ARA 1	99			
Actinomyces meyeri		4.27	0.13		1			
07112502	**Good identification**							**07112502**
Propionibacterium propionicum		98.18	0.49	NIT 86				
Lactobacillus fermentum		1.82	0.20	ARA 1				
07112540	**Good identification**							**07112540**
Propionibacterium propionicum		100.0	0.46	NIT 86				
07112542	**Good identification**							**07112542**
Propionibacterium propionicum		100.0	0.49	NIT 86				
07113000	**Low discrimination**							**07113000**
					COC	GRA	CAT	
Propionibacterium granulosum		83.30	0.82		1	99	90	
Gemella morbillorum		13.07	0.61		99	99	1	
Prevotella intermedia		1.97	0.46		1	1	1	
07113001	**Low discrimination**							**07113001**
					COC	CAT		
Propionibacterium granulosum		52.77	0.48	URE 1	1	90		
Actinomyces naeslundii		26.71	0.58		1	25		
Gemella morbillorum		8.28	0.27		99	1		
07113002	**Good identification**							**07113002**
					CAT			
Propionibacterium granulosum		92.50	0.72		90			
Lactobacillus fermentum		2.76	0.50		1			
Propionibacterium propionicum		2.45	0.49		1			
07113003	**Low discrimination**							**07113003**
					CAT			
Actinomyces naeslundii		58.63	0.58	RAF 92	25			
Propionibacterium granulosum		28.95	0.38		90			
Propionibacterium propionicum		8.45	0.33		1			
07113010	**Very good identification**							**07113010**
					COC	GRA	CAT	
Propionibacterium granulosum		98.41	0.67	MLZ 12	1	99	90	
Gemella morbillorum		1.14	0.27		99	99	1	
Prevotella intermedia		0.17	0.12		1	1	1	
07113012	**Very good identification**							**07113012**
					CAT			
Propionibacterium granulosum		98.90	0.57	MLZ 12	90			
Actinomyces naeslundii		0.52	0.33		25			
Lactobacillus fermentum		0.22	0.16		1			
07113020	**Low discrimination**							**07113020**
					COC	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		84.07	0.72		1	90	
Clostridium perfringens		7.65	0.46		1	1	
Gemella morbillorum		5.86	0.45		99	1	
07113021	**Low discrimination**						**07113021**
					SPO	CAT	
Actinomyces naeslundii		59.76	0.58	RAF 92	1	25	
Propionibacterium granulosum		29.51	0.38		1	90	
Clostridium chauvoei		5.99	0.18		99	1	
07113022	**Low discrimination**						**07113022**
					CAT		
Propionibacterium granulosum		78.53	0.61		90		
Clostridium perfringens		13.45	0.41		1		
Actinomyces naeslundii		4.28	0.55		25		
07113023	**Low discrimination**						**07113023**
					CAT		
Actinomyces naeslundii		86.04	0.58	RAF 92	25		
Propionibacterium granulosum		10.62	0.27		90		
Clostridium perfringens		1.82	0.07		1		
07113030	**Very good identification**						**07113030**
					CAT		
Propionibacterium granulosum		98.31	0.57	MLZ 12	90		
Clostridium perfringens		0.66	0.12		1		
Actinomyces naeslundii		0.52	0.33		25		
07113032	**Good identification**						**07113032**
					CAT		
Propionibacterium granulosum		96.74	0.47	MLZ 12	90		
Actinomyces naeslundii		2.04	0.33		25		
Clostridium perfringens		1.23	0.07		1		
07113033	**Low discrimination**						**07113033**
Actinomyces naeslundii		75.76	0.36	RAF 92			
Propionibacterium granulosum		24.24	0.12	MLZ 5			
07113040	**Low discrimination**						**07113040**
					SPO	CAT	
Propionibacterium granulosum		48.37	0.48	SOR 1	1	90	
Propionibacterium propionicum		21.17	0.46		1	1	
Clostridium botulinum biovar B		10.08	0.24		99	1	
07113041	**Low discrimination**						**07113041**
					CAT		
Actinomyces naeslundii		60.34	0.45	RAF 92	25		
Propionibacterium propionicum		32.84	0.30		1		
Propionibacterium granulosum		6.82	0.14		90		
07113042	**Low discrimination**						**07113042**
					CAT		
Propionibacterium propionicum		65.81	0.49	NIT 86	1		
Propionibacterium granulosum		25.06	0.38	LAC 99	90		
Actinomyces naeslundii		5.97	0.42		25		
07113043	**Low discrimination**						**07113043**
Actinomyces naeslundii		55.05	0.45	RAF 92			
Propionibacterium propionicum		44.95	0.33				
07113061	**Good identification**						**07113061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	RAF 92		
07113063	**Good identification**					**07113063**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07113100	**Good identification**					**07113100**
					CAT	
Propionibacterium propionicum		90.75	0.81	NIT 86	1	
Lactobacillus fermentum		2.53	0.54		1	
Propionibacterium granulosum		2.10	0.48		90	
07113101	**Low discrimination**					**07113101**
					SPO	
Clostridium chauvoei		70.57	0.60	MAN 1	99	
Propionibacterium propionicum		22.55	0.64		1	
Actinomyces naeslundii		6.75	0.65		1	
07113102	**Very good identification**					**07113102**
Propionibacterium propionicum		95.88	0.84	NIT 86		
Lactobacillus fermentum		1.78	0.54			
Actinomyces naeslundii		1.42	0.62			
07113103	**Low discrimination**					**07113103**
					SPO	
Propionibacterium propionicum		63.69	0.67	NIT 86	1	
Clostridium chauvoei		23.45	0.47	URE 10	99	
Actinomyces naeslundii		12.71	0.65		1	
07113110	**Low discrimination**					**07113110**
					CAT	
Propionibacterium propionicum		68.18	0.46	NIT 86	1	
Propionibacterium granulosum		21.24	0.33	MLZ 1	90	
Actinomyces naeslundii		7.88	0.41		25	
07113112	**Low discrimination**					**07113112**
Propionibacterium propionicum		82.74	0.49	NIT 86		
Actinomyces naeslundii		6.37	0.41	MLZ 1		
Bifidobacterium adolescentis		5.05	0.23			
07113113	**Low discrimination**					**07113113**
					SPO	
Actinomyces naeslundii		43.18	0.44	RAF 92	1	
Propionibacterium propionicum		41.53	0.33	MLZ 5	1	
Clostridium chauvoei		15.29	0.13		99	
07113120	**Low discrimination**					**07113120**
					GRA	
Clostridium perfringens		60.46	0.61	MAN 1	99	
Prevotella melaninogenica		31.53	0.50		1	
Actinomyces naeslundii		3.52	0.62		99	
07113121	**Low discrimination**					**07113121**
					SPO	
Clostridium chauvoei		74.23	0.52	MAN 1	99	
Actinomyces naeslundii		21.30	0.65		1	
Clostridium perfringens		2.47	0.27		84	
07113122	**Low discrimination**					**07113122**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		66.57	0.56	MAN 1	99	
Prevotella melaninogenica		14.05	0.37		1	
Actinomyces naeslundii		8.23	0.62		99	
07113123	**Low discrimination**					**07113123**
					SPO	
Actinomyces naeslundii		57.75	0.65	RAF 92	1	
Clostridium chauvoei		35.52	0.39		99	
Clostridium perfringens		3.15	0.22		84	
07113131	**Low discrimination**					**07113131**
					SPO	
Actinomyces naeslundii		59.92	0.44	RAF 92	1	
Clostridium chauvoei		40.08	0.18	MLZ 5	99	
07113133	**Low discrimination**					**07113133**
					SPO	
Actinomyces naeslundii		89.44	0.44	RAF 92	1	
Clostridium chauvoei		10.56	0.05	MLZ 5	99	
07113140	**Very good identification**					**07113140**
Propionibacterium propionicum		99.34	0.81	NIT 86		
Actinomyces naeslundii		0.39	0.50			
Clostridium perfringens		0.22	0.23			
07113141	**Good identification**					**07113141**
					SPO	
Propionibacterium propionicum		92.21	0.64	NIT 86	1	
Actinomyces naeslundii		4.87	0.53	URE 10	1	
Clostridium chauvoei		2.92	0.26		99	
07113142	**Very good identification**					**07113142**
Propionibacterium propionicum		99.64	0.84	NIT 86		
Actinomyces naeslundii		0.26	0.50			
Clostridium perfringens		0.07	0.18			
07113143	**Very good identification**					**07113143**
					SPO	
Propionibacterium propionicum		96.25	0.67	NIT 86	1	
Actinomyces naeslundii		3.39	0.53	URE 10	1	
Clostridium chauvoei		0.36	0.13		99	
07113150	**Good identification**					**07113150**
Propionibacterium propionicum		98.00	0.46	NIT 86		
Actinomyces naeslundii		2.00	0.28	MLZ 1		
07113152	**Good identification**					**07113152**
Propionibacterium propionicum		97.93	0.49	NIT 86		
Actinomyces naeslundii		1.33	0.28	MLZ 1		
Bifidobacterium adolescentis		0.74	0.08			
07113160	**Low discrimination**					**07113160**
Clostridium perfringens		81.93	0.49	MAN 1		
Propionibacterium propionicum		11.40	0.46			
Actinomyces naeslundii		4.41	0.50			
07113161	**Low discrimination**					**07113161**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		65.98	0.53	RAF 92	1			
Clostridium chauvoei		13.16	0.18		99			
Propionibacterium propionicum		12.62	0.30		1			
07113162	**Low discrimination**							**07113162**
Clostridium perfringens		63.73	0.43	MAN 1				
Propionibacterium propionicum		28.26	0.49					
Actinomyces naeslundii		7.30	0.50					
07113163	**Low discrimination**							**07113163**
Actinomyces naeslundii		72.42	0.53	RAF 92				
Propionibacterium propionicum		20.77	0.33					
Clostridium perfringens		4.26	0.09					
07113200	**Good identification**							**07113200**
					COC	GRA	CAT	
Propionibacterium granulosum		93.93	0.56	RHA 3	1	99	90	
Gemella morbillorum		4.81	0.27		99	99	1	
Prevotella intermedia		0.73	0.12		1	1	1	
07113202	**Good identification**							**07113202**
					CAT			
Propionibacterium granulosum		98.19	0.46	RHA 3	90			
Lactobacillus fermentum		0.96	0.16		1			
Propionibacterium propionicum		0.85	0.15		1			
07113210	**Good identification**							**07113210**
Propionibacterium granulosum		100.0	0.41	RHA 3				
07113220	**Good identification**							**07113220**
					COC	CAT		
Propionibacterium granulosum		95.01	0.46	RHA 3	1	90		
Clostridium perfringens		2.82	0.12		1	1		
Gemella morbillorum		2.16	0.11		99	1		
07113222	**Acceptable identification**							**07113222**
					CAT			
Propionibacterium granulosum		94.70	0.36	RHA 3	90			
Clostridium perfringens		5.30	0.07		1			
07113300	**Low discrimination**							**07113300**
					CAT			
Propionibacterium propionicum		88.30	0.46	NIT 86	1			
Propionibacterium granulosum		6.24	0.22	RHA 1	90			
Lactobacillus fermentum		2.46	0.20		1			
07113302	**Good identification**							**07113302**
Propionibacterium propionicum		95.68	0.49	NIT 86				
Lactobacillus fermentum		1.78	0.20	RHA 1				
Actinomyces naeslundii		1.41	0.28					
07113340	**Good identification**							**07113340**
Propionibacterium propionicum		100.0	0.46	NIT 86				
07113342	**Good identification**							**07113342**
Propionibacterium propionicum		100.0	0.49	NIT 86				
07113400	**Low discrimination**							**07113400**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	CAT	
Gemella morbillorum		49.29	0.45	MAN 1	99	1	
Propionibacterium granulosum		28.55	0.48	ARA 10	1	90	
Lactobacillus fermentum		21.12	0.50		1	1	
07113402	**Low discrimination**						**07113402**
					COC	CAT	
Lactobacillus fermentum		72.96	0.50	MAN 1	1	1	
Propionibacterium granulosum		24.67	0.38		1	90	
Gemella morbillorum		1.72	0.11		99	1	
07113420	**Low discrimination**						**07113420**
					COC	CAT	
Propionibacterium granulosum		35.16	0.38	ARA 1	1	90	
Lactobacillus fermentum		34.67	0.42		1	1	
Gemella morbillorum		26.97	0.29		99	1	
07113422	**Low discrimination**						**07113422**
					CAT		
Lactobacillus fermentum		77.10	0.42	MAN 1	1		
Propionibacterium granulosum		19.55	0.27		90		
Clostridium perfringens		3.35	0.07		1		
07113500	**Low discrimination**						**07113500**
Lactobacillus fermentum		69.00	0.54	MAN 1			
Propionibacterium propionicum		25.03	0.46				
Actinomyces meyeri		1.88	0.16				
07113502	**Low discrimination**						**07113502**
Lactobacillus fermentum		56.80	0.54	MAN 1			
Propionibacterium propionicum		30.90	0.49				
Bifidobacterium adolescentis		10.33	0.36				
07113512	**Low discrimination**						**07113512**
Bifidobacterium adolescentis		82.18	0.37	RAF 92			
Bifidobacterium longum subsp. longum		12.06	0.22	XYL 87			
Lactobacillus fermentum		3.74	0.20				
07113520	**Low discrimination**						**07113520**
Lactobacillus fermentum		52.43	0.46	MAN 1			
Clostridium perfringens		21.75	0.27				
Bifidobacterium longum subsp. longum		12.05	0.28				
07113522	**Low discrimination**						**07113522**
Lactobacillus fermentum		65.70	0.46	MAN 1			
Bifidobacterium longum subsp. longum		15.09	0.28				
Clostridium perfringens		12.83	0.22				
07113540	**Good identification**						**07113540**
Propionibacterium propionicum		97.29	0.46	NIT 86			
Lactobacillus fermentum		2.71	0.20	ARA 1			
07113542	**Acceptable identification**						**07113542**
Propionibacterium propionicum		94.35	0.49	NIT 86			
Bifidobacterium adolescentis		3.90	0.20	ARA 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		1.75	0.20					
07114000	**Very good identification**							**07114000**
					COC	GRA		
Gemella morbillorum		96.86	0.84	XYL 10	99	99		
Actinomyces meyeri		1.29	0.53		1	99		
Prevotella intermedia		0.72	0.46		1	1		
07114001	**Good identification**							**07114001**
					COC	GRA		
Gemella morbillorum		97.17	0.50	XYL 10	99	99		
Actinomyces meyeri		1.29	0.18	URE 1	1	99		
Prevotella intermedia		0.72	0.12		1	1		
07114002	**Low discrimination**							**07114002**
					COC	CAT		
Gemella morbillorum		73.93	0.50	XYL 10	99	1		
Lactobacillus fermentum		10.02	0.46	BGL 1	1	1		
Propionibacterium granulosum		7.64	0.40		1	90		
07114010	**Low discrimination**							**07114010**
					COC	SPO	CAT	
Gemella morbillorum		89.82	0.50	XYL 10	99	1	1	
Propionibacterium granulosum		5.06	0.35	MLZ 1	1	1	90	
Clostridium botulinum biovar B		2.88	0.18		1	99	1	
07114020	**Very good identification**							**07114020**
					COC	CAT		
Gemella morbillorum		96.60	0.67	XYL 10	99	1		
Actinomyces meyeri		1.58	0.38	NAG 10	1	1		
Propionibacterium granulosum		0.91	0.40		1	90		
07114040	**Low discrimination**							**07114040**
					COC	SPO		
Clostridium botulinum biovar B		83.91	0.58	XYL 1	1	99		
Gemella morbillorum		13.04	0.50		99	1		
Clostridium glycolicum		2.01	0.32		1	99		
07114050	**Good identification**							**07114050**
					COC	SPO		
Clostridium botulinum biovar B		99.63	0.52	XYL 1	1	99		
Gemella morbillorum		0.32	0.15		99	1		
Clostridium glycolicum		0.05	-.03		1	99		
07114060	**Good identification**							**07114060**
					COC	SPO		
Clostridium botulinum biovar B		96.51	0.52	XYL 1	1	99		
Gemella morbillorum		3.38	0.33		99	1		
Actinomyces meyeri		0.06	0.04		1	1		
07114070	**Good identification**							**07114070**
Clostridium botulinum biovar B		100.0	0.47	XYL 1				
07114100	**Low discrimination**							**07114100**
					COC	GRA		
Actinomyces meyeri		77.61	0.63	MNS 1	1	99		
Gemella morbillorum		14.76	0.50		99	99		
Fusobacterium mortiferum		3.63	0.49		1	1		
07114102	**Low discrimination**							**07114102**
					GRA			
Lactobacillus fermentum		49.61	0.50	GAL 99	99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		36.51	0.37		99	
Fusobacterium mortiferum		6.13	0.33		1	
07114120	**Low discrimination**					**07114120**
					COC	
Actinomyces meyeri		67.72	0.48	MNS 1	1	
Gemella morbillorum		10.49	0.33	NAG 12	99	
Lactobacillus fermentum		6.96	0.42		1	
07114122	**Low discrimination**					**07114122**
Lactobacillus fermentum		52.67	0.42	GAL 99		
Clostridium perfringens		17.51	0.22			
Actinomyces meyeri		15.86	0.22			
07114200	**Low discrimination**					**07114200**
					COC	
Gemella morbillorum		86.29	0.50	XYL 10	99	
Actinomyces odontolyticus		10.45	0.38	RHA 1	1	
Actinomyces meyeri		1.15	0.18		1	
07114400	**Very good identification**					**07114400**
					COC	
Gemella morbillorum		95.83	0.67	XYL 10	99	
Actinomyces meyeri		2.86	0.42	ARA 10	1	
Lactobacillus fermentum		1.18	0.46		1	
07114402	**Low discrimination**					**07114402**
					COC	
Lactobacillus fermentum		52.77	0.46	GAL 99	1	
Gemella morbillorum		43.28	0.33		99	
Actinomyces meyeri		3.96	0.16		1	
07114420	**Acceptable identification**					**07114420**
					COC	
Gemella morbillorum		93.14	0.51	XYL 10	99	
Lactobacillus fermentum		3.44	0.38	ARA 10	1	
Actinomyces meyeri		3.42	0.27	NAG 10	1	
07114422	**Low discrimination**					**07114422**
					COC	
Lactobacillus fermentum		76.68	0.38	GAL 99	1	
Gemella morbillorum		20.96	0.17		99	
Actinomyces meyeri		2.35	0.02		1	
07114500	**Low discrimination**					**07114500**
					COC	
Actinomyces meyeri		78.68	0.53	MNS 1	1	
Lactobacillus fermentum		13.23	0.50		1	
Gemella morbillorum		6.65	0.33		99	
07114502	**Low discrimination**					**07114502**
Lactobacillus fermentum		76.49	0.50	GAL 99		
Actinomyces meyeri		14.07	0.27			
Bifidobacterium longum subsp. longum		7.48	0.26			
07114520	**Low discrimination**					**07114520**
Actinomyces meyeri		44.89	0.38	MNS 1		
Bifidobacterium longum subsp. longum		21.65	0.36	NAG 12		
Lactobacillus fermentum		18.45	0.42			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07114522	**Low discrimination**						**07114522**
Bifidobacterium longum subsp. longum		52.19	0.36	RAF 97			
Lactobacillus fermentum		44.47	0.42	GAL 99			
Actinomyces meyeri		3.35	0.12				
07114530	**Good identification**						**07114530**
Bifidobacterium longum subsp. longum		98.79	0.40	RAF 97			
Actinomyces meyeri		1.22	0.04	GAL 99			
07114532	**Good identification**						**07114532**
Bifidobacterium longum subsp. longum		98.14	0.40	RAF 97			
Clostridium clostridioforme		1.86	0.09	GAL 99			
07115000	**Low discrimination**						**07115000**
					COC		
Gemella morbillorum		76.06	0.79	XYL 10	99		
Lactobacillus fermentum		19.14	0.81		1		
Actinomyces meyeri		2.81	0.56		1		
07115001	**Low discrimination**						**07115001**
					COC	SPO	
Gemella morbillorum		71.37	0.45	XYL 10	99	1	
Lactobacillus fermentum		17.96	0.46	URE 1	1	1	
Clostridium chauvoei		6.29	0.26		1	99	
07115002	**Good identification**						**07115002**
					COC	CAT	
Lactobacillus fermentum		94.52	0.81		1	1	
Gemella morbillorum		3.80	0.45		99	1	
Propionibacterium granulosum		0.73	0.40		1	90	
07115003	**Acceptable identification**						**07115003**
					COC	SPO	
Lactobacillus fermentum		90.75	0.46	URE 1	1	1	
Clostridium chauvoei		5.61	0.13		1	99	
Gemella morbillorum		3.64	0.11		99	1	
07115010	**Low discrimination**						**07115010**
					COC	CAT	
Gemella morbillorum		70.68	0.45	XYL 10	99	1	
Lactobacillus fermentum		17.78	0.46	MLZ 1	1	1	
Propionibacterium granulosum		7.40	0.35		1	90	
07115012	**Low discrimination**						**07115012**
					COC	CAT	
Lactobacillus fermentum		86.41	0.46	MLZ 1	1	1	
Propionibacterium granulosum		8.99	0.25		1	90	
Gemella morbillorum		3.47	0.11		99	1	
07115020	**Low discrimination**						**07115020**
					COC		
Gemella morbillorum		51.41	0.63	XYL 10	99		
Lactobacillus fermentum		38.81	0.72	NAG 10	1		
Clostridium perfringens		6.10	0.46		1		
07115022	**Good identification**						**07115022**
					COC		
Lactobacillus fermentum		90.48	0.72		1		
Clostridium perfringens		6.69	0.41		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		1.21	0.29		99		
07115023	**Low discrimination**						**07115023**
					SPO		
Lactobacillus fermentum		88.05	0.38	URE 1	1		
Clostridium perfringens		6.51	0.07		84		
Clostridium chauvoei		5.44	0.05		99		
07115032	**Low discrimination**						**07115032**
					CAT		
Lactobacillus fermentum		84.39	0.38	MLZ 1	1		
Propionibacterium granulosum		6.59	0.15		90		
Clostridium perfringens		6.24	0.07		1		
07115040	**Low discrimination**						**07115040**
					COC	SPO	
Gemella morbillorum		67.88	0.45	XYL 10	99	1	
Lactobacillus fermentum		17.08	0.46	SOR 1	1	1	
Clostridium botulinum biovar B		8.19	0.24		1	99	
07115042	**Acceptable identification**						**07115042**
					COC		
Lactobacillus fermentum		94.83	0.46	SOR 1	1		
Gemella morbillorum		3.81	0.11		99		
Clostridium perfringens		1.36	0.03		1		
07115100	**Low discrimination**						**07115100**
					COC		
Lactobacillus fermentum		67.75	0.84		1		
Actinomyces meyeri		24.42	0.66		1		
Atopobium parvulum		4.10	0.50		99		
07115101	**Good identification**						**07115101**
					SPO		
Clostridium chauvoei		93.49	0.60	XYL 1	99		
Lactobacillus fermentum		4.40	0.50		1		
Actinomyces meyeri		1.59	0.32		1		
07115102	**Very good identification**						**07115102**
					COC		
Lactobacillus fermentum		96.45	0.84		1		
Actinomyces meyeri		1.08	0.40		1		
Atopobium parvulum		1.03	0.38		99		
07115103	**Low discrimination**						**07115103**
					SPO		
Clostridium chauvoei		78.30	0.47	XYL 1	99		
Lactobacillus fermentum		20.87	0.50		1		
Actinomyces meyeri		0.23	0.06		1		
07115110	**Low discrimination**						**07115110**
Lactobacillus fermentum		60.59	0.50	MLZ 1			
Actinomyces meyeri		21.84	0.32				
Bifidobacterium longum subsp. longum		10.09	0.30				
07115112	**Low discrimination**						**07115112**
Lactobacillus fermentum		60.03	0.50	MLZ 1			
Bifidobacterium adolescentis		27.76	0.39				
Bifidobacterium longum subsp. longum		10.00	0.30				
07115120	**Low discrimination**						**07115120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Lactobacillus fermentum		42.75	0.76		99		
Clostridium perfringens		30.20	0.61		99		
Prevotella melaninogenica		15.75	0.50		1		
07115121	**Acceptable identification**						**07115121**
					SPO		
Clostridium chauvoei		90.11	0.52	XYL 1	99		
Lactobacillus fermentum		4.24	0.42		1		
Clostridium perfringens		2.99	0.27		84		
07115122	**Low discrimination**						**07115122**
					GRA		
Lactobacillus fermentum		62.26	0.76		99		
Clostridium perfringens		20.69	0.56		99		
Prevotella melaninogenica		4.37	0.37		1		
07115123	**Low discrimination**						**07115123**
					SPO		
Clostridium chauvoei		70.07	0.39	XYL 1	99		
Lactobacillus fermentum		18.68	0.42		1		
Clostridium perfringens		6.21	0.22		84		
07115130	**Low discrimination**						**07115130**
Bifidobacterium longum subsp. longum		45.98	0.40	RAF 97			
Lactobacillus fermentum		23.01	0.42	ARA 99			
Clostridium perfringens		16.25	0.27				
07115132	**Low discrimination**						**07115132**
					GRA		
Bifidobacterium longum subsp. longum		44.25	0.40	RAF 97	99		
Lactobacillus fermentum		22.15	0.42	ARA 99	99		
Parabacteroides distasonis		12.79	0.35		1		
07115140	**Low discrimination**						**07115140**
Lactobacillus fermentum		61.94	0.50	SOR 1			
Actinomyces meyeri		22.33	0.32				
Clostridium perfringens		8.51	0.23				
07115142	**Low discrimination**						**07115142**
Lactobacillus fermentum		85.46	0.50	SOR 1			
Clostridium perfringens		5.52	0.18				
Bifidobacterium adolescentis		4.00	0.22				
07115160	**Low discrimination**						**07115160**
					GRA		
Clostridium perfringens		89.03	0.49	XYL 1	99		
Lactobacillus fermentum		6.68	0.42		99		
Prevotella melaninogenica		2.46	0.15		1		
07115162	**Low discrimination**						**07115162**
					GRA	CAT	
Clostridium perfringens		79.59	0.43	XYL 1	99	1	
Lactobacillus fermentum		12.70	0.42		99	1	
Capnocytophaga ochracea		5.15	0.20		1	90	
07115200	**Low discrimination**						**07115200**
					COC		
Gemella morbillorum		67.62	0.45	XYL 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		17.01	0.46	RHA 1	1	
Actinomyces odontolyticus		10.14	0.35		1	
07115202	**Low discrimination**					**07115202**
					COC	
Lactobacillus fermentum		69.98	0.46	RHA 1	1	
Actinomyces odontolyticus		25.56	0.32		1	
Gemella morbillorum		2.81	0.11		99	
07115222	**Low discrimination**					**07115222**
					GRA	
Lactobacillus fermentum		63.38	0.38	RHA 1	99	
Prevotella oralis		17.26	0.13		1	
Actinomyces odontolyticus		11.31	0.18		99	
07115300	**Low discrimination**					**07115300**
Lactobacillus fermentum		47.46	0.50	RHA 1		
Eubacterium contortum		17.85	0.29			
Actinomyces meyeri		17.11	0.32			
07115302	**Low discrimination**					**07115302**
Lactobacillus fermentum		82.27	0.50	RHA 1		
Actinomyces odontolyticus		5.82	0.23			
Eubacterium contortum		5.46	0.16			
07115320	**Acceptable identification**					**07115320**
					GRA	
Bacteroides vulgatus		94.18	0.49	RAF 97	1	
Lactobacillus fermentum		2.17	0.42	ARA 93	99	
Clostridium perfringens		1.53	0.27		99	
07115322	**Low discrimination**					**07115322**
					GRA	
Parabacteroides distasonis		40.99	0.48	RAF 99	1	
Prevotella oralis		33.32	0.33		1	
Lactobacillus fermentum		12.74	0.42		99	
07115332	**Acceptable identification**					**07115332**
Parabacteroides distasonis		94.56	0.43	RAF 99		
Clostridium clostridioforme		2.90	0.12			
Prevotella oralis		1.44	-.01			
07115400	**Low discrimination**					**07115400**
					COC	
Lactobacillus fermentum		67.52	0.81		1	
Gemella morbillorum		29.82	0.63		99	
Actinomyces meyeri		2.48	0.45		1	
07115401	**Low discrimination**					**07115401**
					COC	
Lactobacillus fermentum		67.64	0.46	URE 1	1	
Gemella morbillorum		29.87	0.29		99	
Actinomyces meyeri		2.49	0.11		1	
07115402	**Very good identification**					**07115402**
					COC	
Lactobacillus fermentum		99.23	0.81		1	
Gemella morbillorum		0.44	0.29		99	
Bifidobacterium longum subsp. longum		0.16	0.26		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07115403	**Good identification**					**07115403**
Lactobacillus fermentum		100.0	0.46	URE 1		
07115410	**Low discrimination**					**07115410**
					COC	
Lactobacillus fermentum		57.14	0.46	MLZ 1	1	
Gemella morbillorum		25.23	0.29		99	
Bifidobacterium longum subsp. longum		15.52	0.30		1	
07115412	**Low discrimination**					**07115412**
Lactobacillus fermentum		74.57	0.46	MLZ 1		
Bifidobacterium longum subsp. longum		20.26	0.30			
Bifidobacterium adolescentis		3.80	0.19			
07115420	**Low discrimination**					**07115420**
					COC	
Lactobacillus fermentum		83.76	0.72		1	
Gemella morbillorum		12.33	0.46		99	
Bifidobacterium longum subsp. longum		1.62	0.36		1	
07115421	**Low discrimination**					**07115421**
					COC	
Lactobacillus fermentum		87.17	0.38	URE 1	1	
Gemella morbillorum		12.83	0.12		99	
07115422	**Very good identification**					**07115422**
					COC	
Lactobacillus fermentum		97.84	0.72		1	
Bifidobacterium longum subsp. longum		1.89	0.36		1	
Gemella morbillorum		0.15	0.12		99	
07115423	**Good identification**					**07115423**
Lactobacillus fermentum		100.0	0.38	URE 1		
07115430	**Low discrimination**					**07115430**
					COC	
Bifidobacterium longum subsp. longum		73.97	0.40	RAF 97	1	
Lactobacillus fermentum		22.69	0.38	LAC 99	1	
Gemella morbillorum		3.34	0.12		99	
07115432	**Low discrimination**					**07115432**
Bifidobacterium longum subsp. longum		75.55	0.40	RAF 97		
Lactobacillus fermentum		23.17	0.38	LAC 99		
Clostridium clostridioforme		1.28	0.08			
07115440	**Low discrimination**					**07115440**
					COC	
Lactobacillus fermentum		67.64	0.46	SOR 1	1	
Gemella morbillorum		29.87	0.29		99	
Actinomyces meyeri		2.49	0.11		1	
07115442	**Good identification**					**07115442**
Lactobacillus fermentum		100.0	0.46	SOR 1		
07115460	**Low discrimination**					**07115460**
					COC	
Lactobacillus fermentum		87.17	0.38	SOR 1	1	
Gemella morbillorum		12.83	0.12		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07115462	**Good identification**					**07115462**
Lactobacillus fermentum		100.0	0.38	SOR 1		
07115500	**Low discrimination**					**07115500**
Lactobacillus fermentum		83.74	0.84			
Bifidobacterium longum subsp. longum		8.19	0.60			
Actinomyces meyeri		7.55	0.56			
07115501	**Low discrimination**					**07115501**
					SPO	
Lactobacillus fermentum		71.29	0.50	URE 1	1	
Clostridium chauvoei		15.31	0.26		99	
Bifidobacterium longum subsp. longum		6.97	0.26		1	
07115502	**Low discrimination**					**07115502**
Lactobacillus fermentum		88.74	0.84			
Bifidobacterium longum subsp. longum		8.68	0.60			
Bifidobacterium adolescentis		2.27	0.52			
07115503	**Low discrimination**					**07115503**
					SPO	
Lactobacillus fermentum		86.11	0.50	URE 1	1	
Bifidobacterium longum subsp. longum		8.42	0.26		1	
Clostridium chauvoei		3.26	0.13		99	
07115510	**Good identification**					**07115510**
Bifidobacterium longum subsp. longum		93.63	0.64	RAF 97		
Lactobacillus fermentum		5.68	0.50			
Actinomyces meyeri		0.51	0.21			
07115512	**Low discrimination**					**07115512**
Bifidobacterium longum subsp. longum		79.96	0.64	RAF 97		
Bifidobacterium adolescentis		15.01	0.53			
Lactobacillus fermentum		4.85	0.50			
07115520	**Low discrimination**					**07115520**
					GRA	
Bifidobacterium longum subsp. longum		40.89	0.70	RAF 97	99	
Lactobacillus fermentum		34.85	0.76		99	
Bacteroides vulgatus		22.53	0.52		1	
07115521	**Low discrimination**					**07115521**
					GRA	
Bifidobacterium longum subsp. longum		38.21	0.36	RAF 97	99	
Lactobacillus fermentum		32.56	0.42	URE 1	99	
Bacteroides vulgatus		21.05	0.17		1	
07115522	**Low discrimination**					**07115522**
					GRA	
Bifidobacterium longum subsp. longum		53.32	0.70	RAF 97	99	
Lactobacillus fermentum		45.44	0.76		99	
Parabacteroides distasonis		0.30	0.36		1	
07115523	**Low discrimination**					**07115523**
					SPO	
Bifidobacterium longum subsp. longum		53.07	0.36	RAF 97	1	
Lactobacillus fermentum		45.22	0.42	URE 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		1.71	0.05		99		
07115530	**Very good identification**						**07115530**
					GRA		
Bifidobacterium longum subsp. longum		99.16	0.74	RAF 97	99		
Lactobacillus fermentum		0.50	0.42		99		
Bacteroides vulgatus		0.32	0.17		1		
07115531	**Good identification**						**07115531**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97			
07115532	**Very good identification**						**07115532**
					GRA		
Bifidobacterium longum subsp. longum		99.18	0.74	RAF 97	99		
Lactobacillus fermentum		0.50	0.42		99		
Parabacteroides distasonis		0.18	0.31		1		
07115533	**Good identification**						**07115533**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97			
07115540	**Low discrimination**						**07115540**
Lactobacillus fermentum		83.70	0.50	SOR 1			
Bifidobacterium longum subsp. longum		8.19	0.26				
Actinomyces meyeri		7.54	0.21				
07115542	**Low discrimination**						**07115542**
Lactobacillus fermentum		70.37	0.50	SOR 1			
Bifidobacterium adolescentis		22.02	0.36				
Bifidobacterium longum subsp. longum		6.88	0.26				
07115552	**Low discrimination**						**07115552**
Bifidobacterium adolescentis		68.41	0.38	RAF 92			
Bifidobacterium longum subsp. longum		29.78	0.30	SOR 11			
Lactobacillus fermentum		1.81	0.16				
07115560	**Low discrimination**						**07115560**
					GRA		
Bifidobacterium longum subsp. longum		39.23	0.36	RAF 97	99		
Lactobacillus fermentum		33.43	0.42	SOR 1	99		
Bacteroides vulgatus		21.61	0.17		1		
07115562	**Low discrimination**						**07115562**
					GRA	CAT	
Bifidobacterium longum subsp. longum		50.84	0.36	RAF 97	99	1	
Lactobacillus fermentum		43.32	0.42	SOR 1	99	1	
Capnocytophaga ochracea		3.10	0.07		1	90	
07115570	**Good identification**						**07115570**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97			
07115572	**Good identification**						**07115572**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97			
07115600	**Low discrimination**						**07115600**
					COC		
Lactobacillus fermentum		66.21	0.46	RHA 1	1		
Gemella morbillorum		29.24	0.29		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		2.43	0.11		1			
07115602	**Good identification**							**07115602**
Lactobacillus fermentum		97.12	0.46	RHA 1				
Clostridium clostridioforme		2.88	0.12					
07115622	**Low discrimination**							**07115622**
					GRA			
Lactobacillus fermentum		82.15	0.38	RHA 1	99			
Bacteroides vulgatus		7.88	-.01		1			
Clostridium clostridioforme		7.31	0.12		50			
07115702	**Low discrimination**							**07115702**
Lactobacillus fermentum		79.07	0.50	RHA 1				
Bifidobacterium longum subsp. longum		7.73	0.26					
Clostridium clostridioforme		4.81	0.21					
07115720	**Very good identification**							**07115720**
					GRA			
Bacteroides vulgatus		99.59	0.68	RAF 97	1			
Bifidobacterium longum subsp. longum		0.20	0.36		99			
Lactobacillus fermentum		0.17	0.42		99			
07115732	**Low discrimination**							**07115732**
					GRA			
Bifidobacterium longum subsp. longum		50.59	0.40	RAF 97	99			
Parabacteroides distasonis		26.88	0.39	RHA 1	1			
Clostridium clostridioforme		21.05	0.32		50			
07116000	**Low discrimination**							**07116000**
					COC	GRA	CAT	
Gemella morbillorum		70.67	0.50	MAN 1	99	99	1	
Propionibacterium granulosum		22.05	0.48	XYL 10	1	99	90	
Megasphaera elsdenii		5.30	0.32		99	1	1	
07116002	**Low discrimination**							**07116002**
					COC	GRA	CAT	
Propionibacterium granulosum		78.10	0.38	XYL 1	1	99	90	
Gemella morbillorum		10.12	0.15		99	99	1	
Megasphaera elsdenii		8.35	0.15		99	1	1	
07116100	**Low discrimination**							**07116100**
					COC			
Propionibacterium propionicum		65.76	0.46	NIT 86	1			
Actinomyces meyeri		25.59	0.29	XYL 1	1			
Gemella morbillorum		4.87	0.15		99			
07116102	**Good identification**							**07116102**
Propionibacterium propionicum		98.14	0.49	NIT 86				
Lactobacillus fermentum		1.07	0.16	XYL 1				
Actinomyces meyeri		0.79	0.03					
07116140	**Good identification**							**07116140**
Propionibacterium propionicum		100.0	0.46	NIT 86				
07116142	**Good identification**							**07116142**
Propionibacterium propionicum		100.0	0.49	NIT 86				
07117000	**Low discrimination**							**07117000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	CAT	
Gemella morbillorum		52.64	0.45	MAN 1	99	1	
Propionibacterium granulosum		30.49	0.48	XYL 10	1	90	
Lactobacillus fermentum		13.25	0.46		1	1	
07117002	**Low discrimination**						**07117002**
					CAT		
Lactobacillus fermentum		58.62	0.46	MAN 1	1		
Propionibacterium granulosum		33.74	0.38		90		
Actinomyces israelii		4.38	0.28		1		
07117020	**Low discrimination**						**07117020**
					COC	CAT	
Propionibacterium granulosum		40.45	0.38	XYL 1	1	90	
Gemella morbillorum		31.03	0.29		99	1	
Lactobacillus fermentum		23.43	0.38		1	1	
07117022	**Low discrimination**						**07117022**
					CAT		
Lactobacillus fermentum		66.42	0.38	MAN 1	1		
Propionibacterium granulosum		28.67	0.27		90		
Clostridium perfringens		4.91	0.07		1		
07117100	**Low discrimination**						**07117100**
Lactobacillus fermentum		44.69	0.50	MAN 1			
Propionibacterium propionicum		27.60	0.46				
Actinomyces meyeri		16.11	0.32				
07117102	**Low discrimination**						**07117102**
Lactobacillus fermentum		40.74	0.50	MAN 1			
Propionibacterium propionicum		37.74	0.49				
Bifidobacterium adolescentis		12.61	0.36				
07117112	**Low discrimination**						**07117112**
Bifidobacterium adolescentis		87.66	0.37	RAF 92			
Actinomyces israelii		6.35	0.24	ARA 87			
Lactobacillus fermentum		2.34	0.16				
07117120	**Low discrimination**						**07117120**
					GRA		
Lactobacillus fermentum		41.91	0.42	MAN 1	99		
Clostridium perfringens		29.60	0.27		99		
Prevotella melaninogenica		15.44	0.15		1		
07117122	**Low discrimination**						**07117122**
					GRA		
Lactobacillus fermentum		58.57	0.42	MAN 1	99		
Clostridium perfringens		19.47	0.22		99		
Prevotella melaninogenica		4.11	0.03		1		
07117140	**Acceptable identification**						**07117140**
Propionibacterium propionicum		91.47	0.46	NIT 86			
Actinomyces israelii		7.04	0.28	XYL 1			
Lactobacillus fermentum		1.50	0.16				
07117142	**Low discrimination**						**07117142**
Propionibacterium propionicum		88.56	0.49	NIT 86			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		6.82	0.31	XYL 1		
Bifidobacterium adolescentis		3.66	0.20			
07117400	**Low discrimination**					**07117400**
					COC	
Lactobacillus fermentum		64.83	0.46	MAN 1	1	
Gemella morbillorum		28.63	0.29		99	
Actinomyces israelii		2.64	0.24		1	
07117402	**Acceptable identification**					**07117402**
Lactobacillus fermentum		91.30	0.46	MAN 1		
Actinomyces israelii		5.59	0.27			
Bifidobacterium adolescentis		3.12	0.16			
07117420	**Low discrimination**					**07117420**
					COC	
Lactobacillus fermentum		87.17	0.38	MAN 1	1	
Gemella morbillorum		12.83	0.12		99	
07117422	**Good identification**					**07117422**
Lactobacillus fermentum		100.0	0.38	MAN 1		
07117500	**Low discrimination**					**07117500**
Lactobacillus fermentum		76.04	0.50	MAN 1		
Actinomyces israelii		7.60	0.34			
Bifidobacterium longum subsp. longum		7.44	0.26			
07117502	**Low discrimination**					**07117502**
Bifidobacterium adolescentis		62.24	0.50	RAF 92		
Lactobacillus fermentum		30.04	0.50			
Actinomyces israelii		4.51	0.37			
07117512	**Acceptable identification**					**07117512**
Bifidobacterium adolescentis		92.71	0.52	RAF 92		
Bifidobacterium longum subsp. longum		6.10	0.30			
Actinomyces israelii		0.82	0.23			
07117520	**Low discrimination**					**07117520**
					GRA	
Bifidobacterium longum subsp. longum		40.80	0.36	RAF 97	99	
Lactobacillus fermentum		34.77	0.42	MAN 1	99	
Bacteroides vulgatus		22.48	0.17		1	
07117522	**Low discrimination**					**07117522**
Bifidobacterium longum subsp. longum		51.81	0.36	RAF 97		
Lactobacillus fermentum		44.15	0.42	MAN 1		
Bifidobacterium adolescentis		2.77	0.16			
07117530	**Good identification**					**07117530**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97		
07117532	**Good identification**					**07117532**
Bifidobacterium longum subsp. longum		96.30	0.40	RAF 97		
Bifidobacterium adolescentis		3.70	0.17	MAN 1		
07117552	**Good identification**					**07117552**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium adolescentis		97.29	0.36	RAF 92				
Actinomyces israelii		2.71	0.16	SOR 11				
07120000	**Low discrimination**							**07120000**
					COC	SPO	CAT	
Gemella morbillorum		50.51	0.64	TRE 1	99	1	1	
Clostridium botulinum biovar B		33.11	0.56		1	99	1	
Propionibacterium granulosum		8.51	0.58		1	1	90	
07120002	**Low discrimination**							**07120002**
					COC	SPO	CAT	
Propionibacterium granulosum		68.95	0.48	FRU 99	1	1	90	
Gemella morbillorum		16.54	0.30		99	1	1	
Clostridium botulinum biovar B		10.84	0.21		1	99	1	
07120010	**Acceptable identification**							**07120010**
					COC	SPO	CAT	
Clostridium botulinum biovar B		90.32	0.50	SOR 99	1	99	1	
Propionibacterium granulosum		6.43	0.43		1	1	90	
Gemella morbillorum		2.83	0.30		99	1	1	
07120020	**Low discrimination**							**07120020**
					COC	SPO	CAT	
Clostridium botulinum biovar B		67.13	0.50	SOR 99	1	99	1	
Gemella morbillorum		23.11	0.48		99	1	1	
Propionibacterium granulosum		8.76	0.48		1	1	90	
07120022	**Low discrimination**							**07120022**
					COC	SPO	CAT	
Propionibacterium granulosum		66.83	0.38	FRU 99	1	1	90	
Clostridium botulinum biovar B		20.70	0.16		1	99	1	
Gemella morbillorum		7.12	0.14		99	1	1	
07120030	**Good identification**							**07120030**
					COC	SPO	CAT	
Clostridium botulinum biovar B		95.86	0.45	SOR 99	1	99	1	
Propionibacterium granulosum		3.46	0.33		1	1	90	
Gemella morbillorum		0.68	0.14		99	1	1	
07120040	**Very good identification**							**07120040**
					COC	SPO		
Clostridium botulinum biovar B		99.93	0.90		1	99		
Clostridium botulinum biovar A		0.05	0.36		1	99		
Gemella morbillorum		0.02	0.30		99	1		
07120041	**Good identification**							**07120041**
Clostridium botulinum biovar B		99.95	0.56	URE 1				
Clostridium botulinum biovar A		0.05	0.02					
07120042	**Good identification**							**07120042**
					SPO	CAT		
Clostridium botulinum biovar B		99.88	0.56	BGL 1	99	1		
Propionibacterium granulosum		0.07	0.14		1	90		
Clostridium botulinum biovar A		0.05	0.02		99	1		
07120050	**Very good identification**							**07120050**
					SPO	CAT		
Clostridium botulinum biovar B		100.0	0.84		99	1		
Propionibacterium granulosum		0.00	0.09		1	90		
Clostridium botulinum biovar A		0.00	0.02		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07120051	**Good identification**							**07120051**
Clostridium botulinum biovar B		100.0	0.50	URE 1				
07120052	**Good identification**							**07120052**
Clostridium botulinum biovar B		100.0	0.50	BGL 1				
07120060	**Very good identification**							**07120060**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.99	0.84		1	99	1	
Gemella morbillorum		0.00	0.14		99	1	1	
Propionibacterium granulosum		0.00	0.14		1	1	90	
07120061	**Good identification**							**07120061**
Clostridium botulinum biovar B		100.0	0.50	URE 1				
07120062	**Good identification**							**07120062**
Clostridium botulinum biovar B		100.0	0.50	BGL 1				
07120070	**Very good identification**							**07120070**
Clostridium botulinum biovar B		100.0	0.79					
07120071	**Good identification**							**07120071**
Clostridium botulinum biovar B		100.0	0.45	URE 1				
07120072	**Good identification**							**07120072**
Clostridium botulinum biovar B		100.0	0.45	BGL 1				
07120101	**Good identification**							**07120101**
Clostridium chauvoei		100.0	0.37	TRE 10				
07120140	**Good identification**							**07120140**
Clostridium botulinum biovar B		99.95	0.56	LAC 1				
Clostridium botulinum biovar A		0.05	0.02					
07120150	**Good identification**							**07120150**
Clostridium botulinum biovar B		100.0	0.50	LAC 1				
07120160	**Good identification**							**07120160**
					GRA	SPO		
Clostridium botulinum biovar B		98.98	0.50	LAC 1	99	99		
Clostridium perfringens		0.77	0.19		99	84		
Prevotella bivia		0.25	-.03		1	1		
07120170	**Good identification**							**07120170**
Clostridium botulinum biovar B		100.0	0.45	LAC 1				
07120240	**Good identification**							**07120240**
Clostridium botulinum biovar B		99.95	0.56	RHA 1				
Clostridium botulinum biovar A		0.05	0.02					
07120250	**Good identification**							**07120250**
Clostridium botulinum biovar B		100.0	0.50	RHA 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07120260	**Good identification**							**07120260**
Clostridium botulinum biovar B		100.0	0.50	RHA 1				
07120270	**Good identification**							**07120270**
Clostridium botulinum biovar B		100.0	0.45	RHA 1				
07120400	**Acceptable identification**							**07120400**
					COC	SPO	CAT	
Gemella morbillorum		91.86	0.48	TRE 1	99	1	1	
Clostridium botulinum biovar B		5.47	0.21	ARA 10	1	99	1	
Propionibacterium granulosum		1.41	0.24		1	1	90	
07120440	**Good identification**							**07120440**
					COC	SPO		
Clostridium botulinum biovar B		99.78	0.56	ARA 1	1	99		
Gemella morbillorum		0.17	0.14		99	1		
Clostridium botulinum biovar A		0.05	0.02		1	99		
07120450	**Good identification**							**07120450**
Clostridium botulinum biovar B		100.0	0.50	ARA 1				
07120460	**Good identification**							**07120460**
Clostridium botulinum biovar B		100.0	0.50	ARA 1				
07120470	**Good identification**							**07120470**
Clostridium botulinum biovar B		100.0	0.45	ARA 1				
07121000	**Low discrimination**							**07121000**
					COC	SPO	CAT	
Gemella morbillorum		73.96	0.60	TRE 1	99	1	1	
Propionibacterium granulosum		23.14	0.58		1	1	90	
Clostridium botulinum biovar B		0.91	0.21		1	99	1	
07121001	**Low discrimination**							**07121001**
					COC	SPO	CAT	
Clostridium chauvoei		78.61	0.37	TRE 10	1	99	1	
Gemella morbillorum		14.07	0.26	LAC 99	99	1	1	
Propionibacterium granulosum		4.40	0.24		1	1	90	
07121002	**Low discrimination**							**07121002**
					COC	CAT		
Propionibacterium granulosum		78.92	0.48	FRU 99	1	90		
Gemella morbillorum		10.19	0.26		99	1		
Clostridium perfringens		4.83	0.20		1	1		
07121010	**Low discrimination**							**07121010**
					COC	SPO	CAT	
Propionibacterium granulosum		70.13	0.43	FRU 99	1	1	90	
Gemella morbillorum		16.61	0.26	MLZ 12	99	1	1	
Clostridium botulinum biovar B		9.96	0.16		1	99	1	
07121020	**Low discrimination**							**07121020**
					COC	CAT		
Clostridium perfringens		62.31	0.51	FRU 99	1	1		
Gemella morbillorum		21.04	0.43	LAC 88	99	1		
Propionibacterium granulosum		14.81	0.48		1	90		
07121022	**Low discrimination**							**07121022**
					CAT			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		87.03	0.45	FRU 99	1			
Propionibacterium granulosum		10.99	0.38	LAC 88	90			
Actinomyces naeslundii		0.79	0.33		25			
07121040	**Good identification**							**07121040**
					COC	SPO	CAT	
Clostridium botulinum biovar B		98.72	0.56	GAL 1	1	99	1	
Gemella morbillorum		0.82	0.26		99	1	1	
Propionibacterium granulosum		0.26	0.24		1	1	90	
07121050	**Good identification**							**07121050**
					SPO	CAT		
Clostridium botulinum biovar B		99.93	0.50	GAL 1	99	1		
Propionibacterium granulosum		0.07	0.09		1	90		
07121060	**Acceptable identification**							**07121060**
					COC	SPO		
Clostridium botulinum biovar B		90.27	0.50	GAL 1	1	99		
Clostridium perfringens		9.44	0.39		1	84		
Gemella morbillorum		0.17	0.09		99	1		
07121070	**Good identification**							**07121070**
Clostridium botulinum biovar B		99.79	0.45	GAL 1				
Clostridium perfringens		0.21	0.05					
07121100	**Low discrimination**							**07121100**
					COC	SPO		
Atopobium parvulum		53.74	0.46	FRU 99	99	1		
Clostridium perfringens		22.32	0.40		1	84		
Clostridium chauvoei		16.88	0.37		1	99		
07121101	**Very good identification**							**07121101**
					COC	SPO		
Clostridium chauvoei		99.85	0.71	TRE 10	1	99		
Actinomyces naeslundii		0.11	0.44		1	1		
Atopobium parvulum		0.03	0.12		99	1		
07121103	**Very good identification**							**07121103**
					SPO			
Clostridium chauvoei		99.36	0.58	TRE 10	99			
Actinomyces naeslundii		0.60	0.44		1			
Clostridium perfringens		0.04	0.00		84			
07121111	**Good identification**							**07121111**
Clostridium chauvoei		100.0	0.37	TRE 10				
07121120	**Good identification**							**07121120**
					COC	GRA		
Clostridium perfringens		93.77	0.66	FRU 99	1	99		
Prevotella melaninogenica		2.93	0.33		1	1		
Atopobium parvulum		1.23	0.33		99	99		
07121121	**Very good identification**							**07121121**
					SPO			
Clostridium chauvoei		98.36	0.63	TRE 10	99			
Clostridium perfringens		1.29	0.32		84			
Actinomyces naeslundii		0.32	0.44		1			
07121122	**Very good identification**							**07121122**
					COC	GRA		
Clostridium perfringens		97.61	0.60	FRU 99	1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella melaninogenica		1.23	0.21		1	1		
Atopobium parvulum		0.48	0.20		99	99		
07121123	**Acceptable identification**							**07121123**
					SPO			
Clostridium chauvoei		94.96	0.50	TRE 10	99			
Clostridium perfringens		3.31	0.26		84			
Actinomyces naeslundii		1.72	0.44		1			
07121141	**Good identification**							**07121141**
					SPO			
Clostridium chauvoei		98.17	0.37	TRE 10	99			
Actinomyces naeslundii		1.83	0.31	SOR 1	1			
07121160	**Good identification**							**07121160**
					GRA	SPO		
Clostridium perfringens		98.39	0.54	FRU 99	99	84		
Clostridium botulinum biovar B		1.30	0.16		99	99		
Prevotella melaninogenica		0.16	-.01		1	1		
07121162	**Good identification**							**07121162**
Clostridium perfringens		99.68	0.48	FRU 99				
Actinomyces naeslundii		0.32	0.28					
07121301	**Good identification**							**07121301**
Clostridium chauvoei		100.0	0.37	TRE 10				
07121400	**Good identification**							**07121400**
					COC	CAT		
Gemella morbillorum		96.74	0.43	TRE 1	99	1		
Propionibacterium granulosum		2.75	0.24	ARA 10	1	90		
Lactobacillus fermentum		0.51	0.16		1	1		
07121501	**Good identification**							**07121501**
Clostridium chauvoei		100.0	0.37	TRE 10				
07121522	**Low discrimination**							**07121522**
Bifidobacterium longum subsp. longum		63.54	0.35	RAF 97				
Clostridium perfringens		36.46	0.26	FRU 90				
07121530	**Good identification**							**07121530**
Bifidobacterium longum subsp. longum		100.0	0.39	RAF 97				
07121532	**Good identification**							**07121532**
Bifidobacterium longum subsp. longum		100.0	0.39	RAF 97				
07122000	**Low discrimination**							**07122000**
					COC	SPO	CAT	
Propionibacterium granulosum		87.13	0.56	FRU 99	1	1	90	
Gemella morbillorum		6.93	0.30		99	1	1	
Clostridium botulinum biovar B		4.54	0.21		1	99	1	
07122002	**Good identification**							**07122002**
					CAT			
Propionibacterium granulosum		99.28	0.46	FRU 99	90			
Propionibacterium propionicum		0.72	0.14		1			
07122010	**Low discrimination**							**07122010**
					SPO	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		84.16	0.41	FRU 99	1	90		
Clostridium botulinum biovar B		15.84	0.16	MLZ 12	99	1		
07122020	**Low discrimination**							**07122020**
					COC	SPO	CAT	
Propionibacterium granulosum		87.88	0.46	FRU 99	1	1	90	
Clostridium botulinum biovar B		9.02	0.16		1	99	1	
Gemella morbillorum		3.10	0.14		99	1	1	
07122022	**Good identification**							**07122022**
Propionibacterium granulosum		100.0	0.35	FRU 99				
07122040	**Good identification**							**07122040**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.70	0.56	MAN 1	1	99	1	
Propionibacterium granulosum		0.20	0.22		1	1	90	
Megasphaera elsdenii		0.05	0.07		99	1	1	
07122050	**Good identification**							**07122050**
Clostridium botulinum biovar B		100.0	0.50	MAN 1				
07122060	**Good identification**							**07122060**
					SPO	CAT		
Clostridium botulinum biovar B		99.90	0.50	MAN 1	99	1		
Propionibacterium granulosum		0.10	0.12		1	90		
07122070	**Good identification**							**07122070**
Clostridium botulinum biovar B		100.0	0.45	MAN 1				
07122100	**Acceptable identification**							**07122100**
					CAT			
Propionibacterium propionicum		92.20	0.45	NIT 86	1			
Propionibacterium granulosum		7.80	0.22	FRU 99	90			
07122102	**Good identification**							**07122102**
					CAT			
Propionibacterium propionicum		98.61	0.48	NIT 86	1			
Propionibacterium granulosum		1.39	0.12	FRU 99	90			
07122140	**Low discrimination**							**07122140**
					SPO			
Propionibacterium propionicum		69.62	0.45	NIT 86	1			
Clostridium botulinum biovar B		30.39	0.21	FRU 99	99			
07122142	**Good identification**							**07122142**
Propionibacterium propionicum		100.0	0.48	NIT 86				
07123000	**Acceptable identification**							**07123000**
					COC	CAT		
Propionibacterium granulosum		94.85	0.56	FRU 99	1	90		
Gemella morbillorum		4.06	0.26		99	1		
Actinomyces naeslundii		1.09	0.38		1	25		
07123001	**Low discrimination**							**07123001**
					SPO	CAT		
Actinomyces naeslundii		57.94	0.41	RAF 92	1	25		
Propionibacterium granulosum		33.95	0.22	FRU 95	1	90		
Clostridium chauvoei		8.12	0.03		99	1		
07123002	**Acceptable identification**							**07123002**
					CAT			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		94.95	0.46	FRU 99	90	
Actinomyces naeslundii		4.37	0.38		25	
Propionibacterium propionicum		0.69	0.14		1	
07123003	**Low discrimination**					**07123003**
Actinomyces naeslundii		87.23	0.41	RAF 92		
Propionibacterium granulosum		12.78	0.12	FRU 95		
07123010	**Good identification**					**07123010**
Propionibacterium granulosum		100.0	0.41	FRU 99		
07123020	**Low discrimination**					**07123020**
					CAT	
Propionibacterium granulosum		89.18	0.46	FRU 99	90	
Clostridium perfringens		5.02	0.17		1	
Actinomyces naeslundii		4.10	0.38		25	
07123021	**Low discrimination**					**07123021**
Actinomyces naeslundii		87.23	0.41	RAF 92		
Propionibacterium granulosum		12.78	0.12	FRU 95		
07123022	**Low discrimination**					**07123022**
					CAT	
Propionibacterium granulosum		77.52	0.35	FRU 99	90	
Actinomyces naeslundii		14.26	0.38		25	
Clostridium perfringens		8.22	0.11		1	
07123023	**Good identification**					**07123023**
Actinomyces naeslundii		100.0	0.41	RAF 92		
07123100	**Low discrimination**					**07123100**
					CAT	
Propionibacterium propionicum		68.02	0.45	NIT 86	1	
Actinomyces naeslundii		18.64	0.45	FRU 99	25	
Propionibacterium granulosum		5.76	0.22		90	
07123101	**Low discrimination**					**07123101**
					SPO	
Clostridium chauvoei		79.32	0.37	TRE 10	99	
Actinomyces naeslundii		16.28	0.49	MAN 1	1	
Propionibacterium propionicum		4.40	0.29		1	
07123102	**Low discrimination**					**07123102**
					CAT	
Propionibacterium propionicum		82.97	0.48	NIT 86	1	
Actinomyces naeslundii		15.16	0.45	FRU 99	25	
Propionibacterium granulosum		1.17	0.12		90	
07123103	**Low discrimination**					**07123103**
					SPO	
Actinomyces naeslundii		44.14	0.49	RAF 92	1	
Clostridium chauvoei		37.96	0.24	FRU 95	99	
Propionibacterium propionicum		17.90	0.32		1	
07123123	**Low discrimination**					**07123123**
					SPO	
Actinomyces naeslundii		77.72	0.49	RAF 92	1	
Clostridium chauvoei		22.28	0.16	FRU 95	99	
07123140	**Good identification**					**07123140**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		95.39	0.45	NIT 86				
Actinomyces naeslundii		4.61	0.33	FRU 99				
07123142	**Good identification**							**07123142**
Propionibacterium propionicum		96.88	0.48	NIT 86				
Actinomyces naeslundii		3.12	0.33	FRU 99				
07123161	**Good identification**							**07123161**
Actinomyces naeslundii		100.0	0.36	RAF 92				
07123163	**Good identification**							**07123163**
Actinomyces naeslundii		100.0	0.36	RAF 92				
07124000	**Acceptable identification**							**07124000**
					COC	SPO	CAT	
Gemella morbillorum		91.86	0.48	TRE 1	99	1	1	
Clostridium botulinum biovar B		5.47	0.21	XYL 10	1	99	1	
Propionibacterium granulosum		1.41	0.24		1	1	90	
07124040	**Good identification**							**07124040**
					COC	SPO		
Clostridium botulinum biovar B		99.78	0.56	XYL 1	1	99		
Gemella morbillorum		0.17	0.14		99	1		
Clostridium botulinum biovar A		0.05	0.02		1	99		
07124050	**Good identification**							**07124050**
Clostridium botulinum biovar B		100.0	0.50	XYL 1				
07124060	**Good identification**							**07124060**
Clostridium botulinum biovar B		100.0	0.50	XYL 1				
07124070	**Good identification**							**07124070**
Clostridium botulinum biovar B		100.0	0.45	XYL 1				
07125000	**Acceptable identification**							**07125000**
					COC	CAT		
Gemella morbillorum		94.17	0.43	TRE 1	99	1		
Atopobium parvulum		2.86	0.16	XYL 10	99	1		
Propionibacterium granulosum		2.68	0.24		1	90		
07125100	**Good identification**							**07125100**
					COC			
Atopobium parvulum		98.80	0.50	FRU 99	99			
Gemella morbillorum		0.33	0.09		99			
Actinomyces meyeri		0.31	0.10		1			
07125101	**Acceptable identification**							**07125101**
					COC	SPO		
Clostridium chauvoei		94.36	0.37	TRE 10	1	99		
Atopobium parvulum		5.64	0.16	XYL 1	99	1		
07125102	**Good identification**							**07125102**
					COC			
Atopobium parvulum		98.47	0.38	FRU 99	99			
Lactobacillus fermentum		0.94	0.16		1			
Clostridium perfringens		0.59	0.00		1			
07125120	**Low discrimination**							**07125120**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Atopobium parvulum		69.58	0.38	FRU 99	99	99		
Clostridium perfringens		28.80	0.32		1	99		
Prevotella melaninogenica		0.90	-.01		1	1		
07125510	**Good identification**							**07125510**
Bifidobacterium longum subsp. longum		100.0	0.37	RAF 97				
07125512	**Good identification**							**07125512**
Bifidobacterium longum subsp. longum		95.65	0.37	RAF 97				
Bifidobacterium adolescentis		4.35	0.16	FRU 90				
07125520	**Good identification**							**07125520**
					COC	GRA		
Bifidobacterium longum subsp. longum		98.21	0.44	RAF 97	1	99		
Atopobium parvulum		0.97	0.03	FRU 90	99	99		
Bacteroides vulgatus		0.82	-.06		1	1		
07125522	**Good identification**							**07125522**
					GRA			
Bifidobacterium longum subsp. longum		99.07	0.44	RAF 97	99			
Parabacteroides distasonis		0.93	0.13	FRU 90	1			
07125530	**Good identification**							**07125530**
Bifidobacterium longum subsp. longum		100.0	0.48	RAF 97				
07125532	**Good identification**							**07125532**
Bifidobacterium longum subsp. longum		100.0	0.48	RAF 97				
07130000	**Low discrimination**							**07130000**
					COC	SPO	CAT	
Propionibacterium granulosum		80.39	0.92		1	1	90	
Clostridium botulinum biovar B		12.64	0.66		1	99	1	
Gemella morbillorum		5.89	0.66		99	1	1	
07130001	**Low discrimination**							**07130001**
					COC	SPO	CAT	
Propionibacterium granulosum		79.89	0.58	URE 1	1	1	90	
Clostridium botulinum biovar B		12.56	0.32		1	99	1	
Gemella morbillorum		5.86	0.32		99	1	1	
07130002	**Very good identification**							**07130002**
					COC	SPO	CAT	
Propionibacterium granulosum		98.63	0.82		1	1	90	
Clostridium botulinum biovar B		0.63	0.32		1	99	1	
Gemella morbillorum		0.29	0.32		99	1	1	
07130003	**Acceptable identification**							**07130003**
Propionibacterium granulosum		94.05	0.48	URE 1				
Actinomyces viscosus		3.35	0.14					
Actinomyces naeslundii		2.00	0.34					
07130010	**Low discrimination**							**07130010**
					COC	SPO	CAT	
Propionibacterium granulosum		63.51	0.77	MLZ 12	1	1	90	
Clostridium botulinum biovar B		36.06	0.61		1	99	1	
Gemella morbillorum		0.35	0.32		99	1	1	
07130011	**Low discrimination**							**07130011**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	CAT		
Propionibacterium granulosum		63.78	0.43	MLZ 12	1	90		
Clostridium botulinum biovar B		36.22	0.26	URE 1	99	1		
07130012	**Very good identification**							**07130012**
					SPO	CAT		
Propionibacterium granulosum		97.64	0.67	MLZ 12	1	90		
Clostridium botulinum biovar B		2.24	0.26		99	1		
Lactobacillus jensenii		0.12	0.11		1	1		
07130020	**Low discrimination**							**07130020**
					CAT	SPO		
Propionibacterium granulosum		72.16	0.82		90	1		
Clostridium botulinum biovar B		22.35	0.61		1	99		
Clostridium perfringens		3.07	0.51		1	84		
07130021	**Low discrimination**							**07130021**
					CAT	SPO		
Propionibacterium granulosum		70.84	0.48	URE 1	90	1		
Clostridium botulinum biovar B		21.94	0.26		1	99		
Clostridium perfringens		3.01	0.17		1	84		
07130022	**Good identification**							**07130022**
					CAT	SPO		
Propionibacterium granulosum		91.33	0.72		90	1		
Clostridium perfringens		7.31	0.45		1	84		
Clostridium botulinum biovar B		1.14	0.26		1	99		
07130023	**Low discrimination**							**07130023**
					CAT			
Propionibacterium granulosum		85.85	0.38	URE 1	90			
Actinomyces naeslundii		7.29	0.34		25			
Clostridium perfringens		6.87	0.11		1			
07130030	**Low discrimination**							**07130030**
					CAT	SPO		
Clostridium botulinum biovar B		52.65	0.55	SOR 99	1	99		
Propionibacterium granulosum		47.07	0.67		90	1		
Clostridium perfringens		0.15	0.17		1	84		
07130032	**Very good identification**							**07130032**
					CAT	SPO		
Propionibacterium granulosum		95.14	0.57	MLZ 12	90	1		
Clostridium botulinum biovar B		4.30	0.21		1	99		
Clostridium perfringens		0.56	0.11		1	84		
07130040	**Excellent identification**							**07130040**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.93	1.00		1	99	1	
Propionibacterium granulosum		0.07	0.58		1	1	90	
Gemella morbillorum		0.01	0.32		99	1	1	
07130041	**Very good identification**							**07130041**
					SPO	CAT		
Clostridium botulinum biovar B		99.94	0.66	URE 1	99	1		
Propionibacterium granulosum		0.07	0.24		1	90		
07130042	**Very good identification**							**07130042**
					COC	SPO	CAT	
Clostridium botulinum biovar B		98.36	0.66	BGL 1	1	99	1	
Propionibacterium granulosum		1.58	0.48		1	1	90	
Megasphaera elsdenii		0.03	0.12		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07130050	**Excellent identification**							**07130050**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.95		99	1		
Propionibacterium granulosum		0.02	0.43		1	90		
07130051	**Very good identification**							**07130051**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.61	URE 1	99	1		
Propionibacterium granulosum		0.02	0.09		1	90		
07130052	**Very good identification**							**07130052**
					SPO	CAT		
Clostridium botulinum biovar B		99.56	0.61	BGL 1	99	1		
Propionibacterium granulosum		0.44	0.33		1	90		
07130060	**Excellent identification**							**07130060**
					CAT	SPO		
Clostridium botulinum biovar B		99.94	0.95		1	99		
Propionibacterium granulosum		0.03	0.48		90	1		
Clostridium perfringens		0.03	0.39		1	84		
07130061	**Very good identification**							**07130061**
					CAT	SPO		
Clostridium botulinum biovar B		99.94	0.61	URE 1	1	99		
Propionibacterium granulosum		0.03	0.14		90	1		
Clostridium perfringens		0.03	0.05		1	84		
07130062	**Very good identification**							**07130062**
					CAT	SPO		
Clostridium botulinum biovar B		98.00	0.61	BGL 1	1	99		
Clostridium perfringens		1.21	0.33		1	84		
Propionibacterium granulosum		0.80	0.38		90	1		
07130070	**Very good identification**							**07130070**
					CAT	SPO		
Clostridium botulinum biovar B		99.99	0.90		1	99		
Propionibacterium granulosum		0.01	0.33		90	1		
Clostridium perfringens		0.00	0.05		1	84		
07130071	**Good identification**							**07130071**
Clostridium botulinum biovar B		100.0	0.55	URE 1				
07130072	**Good identification**							**07130072**
					SPO	CAT		
Clostridium botulinum biovar B		99.78	0.55	BGL 1	99	1		
Propionibacterium granulosum		0.23	0.23		1	90		
07130100	**Low discrimination**							**07130100**
					COC	GRA	CAT	
Propionibacterium granulosum		33.69	0.58	LAC 1	1	99	90	
Fusobacterium mortiferum		26.51	0.60		1	1	5	
Atopobium parvulum		19.35	0.46		99	99	1	
07130101	**Low discrimination**							**07130101**
					SPO	CAT		
Clostridium chauvoei		88.83	0.44	TRE 10	99	1		
Actinomyces naeslundii		3.19	0.42	GAL 99	1	25		
Propionibacterium granulosum		2.13	0.24		1	90		
07130102	**Low discrimination**							**07130102**
					CAT			
Propionibacterium granulosum		33.41	0.48	LAC 1	90			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		18.05	0.48		1			
Clostridium perfringens		15.00	0.34		1			
07130110	**Low discrimination**							**07130110**
					GRA	SPO	CAT	
Propionibacterium granulosum		59.30	0.43	LAC 1	99	1	90	
Clostridium botulinum biovar B		33.67	0.26	MLZ 12	99	99	1	
Fusobacterium mortiferum		3.46	0.25		1	1	5	
07130120	**Good identification**							**07130120**
					COC	CAT		
Clostridium perfringens		93.44	0.66	GAL 99	1	1		
Propionibacterium granulosum		3.03	0.48		1	90		
Atopobium parvulum		1.23	0.33		99	1		
07130121	**Low discrimination**							**07130121**
					SPO			
Clostridium chauvoei		59.49	0.35	TRE 10	99			
Clostridium perfringens		33.04	0.32	GAL 99	84			
Actinomyces naeslundii		6.40	0.42		1			
07130122	**Very good identification**							**07130122**
					COC	CAT		
Clostridium perfringens		97.38	0.60	GAL 99	1	1		
Propionibacterium granulosum		1.68	0.38		1	90		
Atopobium parvulum		0.48	0.20		99	1		
07130140	**Very good identification**							**07130140**
					SPO			
Clostridium botulinum biovar B		98.98	0.66	LAC 1	99			
Propionibacterium propionicum		0.57	0.45		1			
Clostridium perfringens		0.29	0.28		84			
07130150	**Very good identification**							**07130150**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.61	LAC 1	99	1		
Propionibacterium granulosum		0.02	0.09		1	90		
07130160	**Low discrimination**							**07130160**
					CAT	SPO		
Clostridium botulinum biovar B		83.88	0.61	LAC 1	1	99		
Clostridium perfringens		16.08	0.54		1	84		
Propionibacterium granulosum		0.03	0.14		90	1		
07130162	**Low discrimination**							**07130162**
					SPO			
Clostridium perfringens		89.56	0.48	GAL 99	84			
Clostridium botulinum biovar B		10.03	0.26		99			
Actinomyces naeslundii		0.23	0.26		1			
07130170	**Good identification**							**07130170**
Clostridium botulinum biovar B		99.61	0.55	LAC 1				
Clostridium perfringens		0.39	0.19					
07130200	**Good identification**							**07130200**
					COC	SPO	CAT	
Propionibacterium granulosum		92.61	0.66	RHA 3	1	1	90	
Clostridium botulinum biovar B		4.76	0.32		1	99	1	
Gemella morbillorum		2.22	0.32		99	1	1	
07130202	**Very good identification**							**07130202**
					SPO	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		99.63	0.56	RHA 3	1	90		
Clostridium botulinum biovar B		0.21	-.03		99	1		
Actinomyces odontolyticus		0.16	0.06		1	10		
07130210	**Low discrimination**							**07130210**
					SPO	CAT		
Propionibacterium granulosum		84.36	0.52	RHA 3	1	90		
Clostridium botulinum biovar B		15.64	0.26	MLZ 12	99	1		
07130212	**Good identification**							**07130212**
Propionibacterium granulosum		100.0	0.41	RHA 3				
07130220	**Low discrimination**							**07130220**
					CAT	SPO		
Propionibacterium granulosum		88.84	0.56	RHA 3	90	1		
Clostridium botulinum biovar B		8.99	0.26		1	99		
Clostridium perfringens		1.23	0.17		1	84		
07130222	**Good identification**							**07130222**
					CAT			
Propionibacterium granulosum		97.45	0.46	RHA 3	90			
Clostridium perfringens		2.55	0.11		1			
07130230	**Low discrimination**							**07130230**
					SPO	CAT		
Propionibacterium granulosum		73.24	0.41	RHA 3	1	90		
Clostridium botulinum biovar B		26.76	0.21	MLZ 12	99	1		
07130240	**Very good identification**							**07130240**
					SPO	CAT		
Clostridium botulinum biovar B		99.80	0.66	RHA 1	99	1		
Propionibacterium granulosum		0.20	0.32		1	90		
07130250	**Very good identification**							**07130250**
					SPO	CAT		
Clostridium botulinum biovar B		99.95	0.61	RHA 1	99	1		
Propionibacterium granulosum		0.06	0.17		1	90		
07130260	**Very good identification**							**07130260**
					CAT	SPO		
Clostridium botulinum biovar B		99.87	0.61	RHA 1	1	99		
Propionibacterium granulosum		0.10	0.22		90	1		
Clostridium perfringens		0.03	0.05		1	84		
07130270	**Good identification**							**07130270**
Clostridium botulinum biovar B		100.0	0.55	RHA 1				
07130400	**Low discrimination**							**07130400**
					COC	SPO	CAT	
Propionibacterium granulosum		50.55	0.58	ARA 1	1	1	90	
Gemella morbillorum		40.75	0.50		99	1	1	
Clostridium botulinum biovar B		7.95	0.32		1	99	1	
07130402	**Good identification**							**07130402**
					COC	CAT		
Propionibacterium granulosum		95.56	0.48	ARA 1	1	90		
Gemella morbillorum		3.11	0.15		99	1		
Lactobacillus fermentum		0.72	0.16		1	1		
07130410	**Low discrimination**							**07130410**
					COC	SPO	CAT	
Propionibacterium granulosum		61.44	0.43	ARA 1	1	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		34.89	0.26	MLZ 12	1	99	1	
Gemella morbillorum		3.67	0.15		99	1	1	
07130420	**Low discrimination**							**07130420**
					COC	SPO	CAT	
Propionibacterium granulosum		58.46	0.48	ARA 1	1	1	90	
Gemella morbillorum		20.95	0.33		99	1	1	
Clostridium botulinum biovar B		18.11	0.26		1	99	1	
07130422	**Acceptable identification**							**07130422**
					CAT			
Propionibacterium granulosum		92.59	0.38	ARA 1	90			
Clostridium perfringens		7.41	0.11		1			
07130440	**Very good identification**							**07130440**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.88	0.66	ARA 1	1	99	1	
Propionibacterium granulosum		0.07	0.24		1	1	90	
Gemella morbillorum		0.05	0.15		99	1	1	
07130450	**Very good identification**							**07130450**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.61	ARA 1	99	1		
Propionibacterium granulosum		0.02	0.09		1	90		
07130460	**Very good identification**							**07130460**
					CAT	SPO		
Clostridium botulinum biovar B		99.94	0.61	ARA 1	1	99		
Propionibacterium granulosum		0.03	0.14		90	1		
Clostridium perfringens		0.03	0.05		1	84		
07130470	**Good identification**							**07130470**
Clostridium botulinum biovar B		100.0	0.55	ARA 1				
07131000	**Low discrimination**							**07131000**
					COC	CAT		
Propionibacterium granulosum		89.88	0.92		1	90		
Gemella morbillorum		3.55	0.61		99	1		
Clostridium perfringens		2.92	0.59		1	1		
07131001	**Low discrimination**							**07131001**
					SPO	CAT		
Propionibacterium granulosum		60.25	0.58	URE 1	1	90		
Clostridium chauvoei		25.36	0.44		99	1		
Actinomyces naeslundii		7.67	0.58		1	25		
07131002	**Low discrimination**							**07131002**
					CAT			
Propionibacterium granulosum		88.27	0.82		90			
Clostridium perfringens		5.41	0.54		1			
Lactobacillus jensenii		4.54	0.53		1			
07131003	**Low discrimination**							**07131003**
					SPO	CAT		
Propionibacterium granulosum		51.33	0.48	URE 1	1	90		
Actinomyces naeslundii		26.13	0.58		1	25		
Clostridium chauvoei		15.25	0.31		99	1		
07131010	**Good identification**							**07131010**
					SPO	CAT		
Propionibacterium granulosum		94.34	0.77	MLZ 12	1	90		
Lactobacillus jensenii		4.45	0.48		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar B		0.54	0.26		99	1	
07131011	**Low discrimination**						**07131011**
					CAT		
Propionibacterium granulosum		88.70	0.43	MLZ 12	90		
Actinomyces naeslundii		4.36	0.36	URE 1	25		
Lactobacillus jensenii		4.18	0.14		1		
07131012	**Low discrimination**						**07131012**
					CAT		
Propionibacterium granulosum		88.33	0.67	MLZ 12	90		
Lactobacillus jensenii		11.10	0.45		1		
Clostridium perfringens		0.40	0.20		1		
07131020	**Low discrimination**						**07131020**
					CAT		
Clostridium perfringens		80.00	0.85	LAC 88	1		
Propionibacterium granulosum		19.01	0.82		90		
Lactobacillus jensenii		0.37	0.46		1		
07131021	**Low discrimination**						**07131021**
					CAT	SPO	
Clostridium perfringens		66.63	0.51	LAC 88	1	84	
Propionibacterium granulosum		15.84	0.48	URE 1	90	1	
Clostridium chauvoei		8.89	0.35		1	99	
07131022	**Low discrimination**						**07131022**
					CAT		
Clostridium perfringens		87.97	0.80	LAC 88	1		
Propionibacterium granulosum		11.11	0.72		90		
Lactobacillus jensenii		0.57	0.43		1		
07131023	**Low discrimination**						**07131023**
					CAT		
Clostridium perfringens		69.45	0.45	LAC 88	1		
Actinomyces naeslundii		17.86	0.58	URE 1	25		
Propionibacterium granulosum		8.77	0.38		90		
07131030	**Low discrimination**						**07131030**
					CAT		
Propionibacterium granulosum		72.85	0.67	MLZ 12	90		
Clostridium perfringens		22.70	0.51		1		
Lactobacillus jensenii		3.43	0.38		1		
07131032	**Low discrimination**						**07131032**
					CAT		
Propionibacterium granulosum		58.22	0.57	MLZ 12	90		
Clostridium perfringens		34.16	0.45		1		
Lactobacillus jensenii		7.32	0.35		1		
07131040	**Good identification**						**07131040**
					CAT	SPO	
Clostridium botulinum biovar B		90.12	0.66	GAL 1	1	99	
Propionibacterium granulosum		5.79	0.58		90	1	
Clostridium perfringens		3.55	0.47		1	84	
07131042	**Low discrimination**						**07131042**
					CAT	SPO	
Clostridium perfringens		39.65	0.41	LAC 88	1	84	
Propionibacterium granulosum		34.34	0.48	NAG 97	90	1	
Clostridium botulinum biovar B		21.59	0.32		1	99	
07131043	**Low discrimination**						**07131043**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Actinomyces naeslundii		76.17	0.45	RAF 92	25		
Clostridium perfringens		9.89	0.07		1		
Propionibacterium granulosum		8.56	0.14		90		
07131050	**Very good identification**						**07131050**
					SPO	CAT	
Clostridium botulinum biovar B		98.09	0.61	GAL 1	99	1	
Propionibacterium granulosum		1.75	0.43		1	90	
Lactobacillus jensenii		0.08	0.14		1	1	
07131060	**Low discrimination**						**07131060**
					CAT	SPO	
Clostridium perfringens		71.42	0.73	LAC 88	1	84	
Clostridium botulinum biovar B		27.60	0.61		1	99	
Propionibacterium granulosum		0.90	0.48		90	1	
07131061	**Low discrimination**						**07131061**
					SPO		
Clostridium perfringens		65.87	0.39	LAC 88	84		
Clostridium botulinum biovar B		25.45	0.26	URE 1	99		
Actinomyces naeslundii		7.39	0.45		1		
07131062	**Very good identification**						**07131062**
					CAT	SPO	
Clostridium perfringens		98.32	0.67	LAC 88	1	84	
Clostridium botulinum biovar B		0.82	0.26		1	99	
Propionibacterium granulosum		0.66	0.38		90	1	
07131070	**Acceptable identification**						**07131070**
					CAT	SPO	
Clostridium botulinum biovar B		94.12	0.55	GAL 1	1	99	
Clostridium perfringens		5.00	0.39		1	84	
Propionibacterium granulosum		0.85	0.33		90	1	
07131100	**Low discrimination**						**07131100**
					COC	GRA	
Atopobium parvulum		67.81	0.80		99	99	
Clostridium perfringens		28.17	0.74		1	99	
Fusobacterium mortiferum		1.41	0.63		1	1	
07131101	**Very good identification**						**07131101**
					COC	SPO	
Clostridium chauvoei		97.18	0.78	TRE 10	1	99	
Atopobium parvulum		1.34	0.46		99	1	
Actinomyces naeslundii		0.85	0.65		1	1	
07131102	**Low discrimination**						**07131102**
					COC		
Clostridium perfringens		49.50	0.69	NAG 97	1		
Atopobium parvulum		44.69	0.67		99		
Lactobacillus fermentum		1.35	0.54		1		
07131103	**Good identification**						**07131103**
					SPO		
Clostridium chauvoei		92.37	0.65	TRE 10	99		
Actinomyces naeslundii		4.55	0.65		1		
Clostridium perfringens		1.41	0.34		84		
07131110	**Low discrimination**						**07131110**
					COC	CAT	
Atopobium parvulum		57.82	0.46	MLZ 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		24.02	0.40		1	1	
Propionibacterium granulosum		13.73	0.43		1	90	
07131111	**Acceptable identification**						**07131111**
					COC	SPO	
Clostridium chauvoei		93.63	0.44	TRE 10	1	99	
Actinomyces naeslundii		4.24	0.44	MLZ 1	1	1	
Atopobium parvulum		1.29	0.12		99	1	
07131120	**Very good identification**						**07131120**
					COC	GRA	
Clostridium perfringens		98.25	1.00		1	99	
Atopobium parvulum		1.29	0.67		99	99	
Prevotella melaninogenica		0.28	0.50		1	1	
07131121	**Low discrimination**						**07131121**
					SPO		
Clostridium chauvoei		62.86	0.70	TRE 10	99		
Clostridium perfringens		34.91	0.66		84		
Actinomyces naeslundii		1.64	0.65		1		
07131122	**Excellent identification**						**07131122**
					COC	GRA	
Clostridium perfringens		99.18	0.94		1	99	
Atopobium parvulum		0.49	0.54		99	99	
Prevotella melaninogenica		0.11	0.37		1	1	
07131123	**Low discrimination**						**07131123**
					SPO		
Clostridium perfringens		56.13	0.60	URE 1	84		
Clostridium chauvoei		37.90	0.57		99		
Actinomyces naeslundii		5.60	0.65		1		
07131130	**Very good identification**						**07131130**
					COC	CAT	
Clostridium perfringens		97.60	0.66	MLZ 1	1	1	
Atopobium parvulum		1.28	0.33		99	1	
Propionibacterium granulosum		0.43	0.33		1	90	
07131131	**Low discrimination**						**07131131**
					SPO		
Clostridium chauvoei		59.13	0.35	TRE 10	99		
Clostridium perfringens		32.84	0.32	MLZ 1	84		
Actinomyces naeslundii		8.04	0.44		1		
07131132	**Very good identification**						**07131132**
					COC	GRA	
Clostridium perfringens		98.04	0.60	MLZ 1	1	99	
Atopobium parvulum		0.48	0.20		99	99	
Parabacteroides distasonis		0.47	0.30		1	1	
07131140	**Low discrimination**						**07131140**
					COC	SPO	
Clostridium perfringens		83.43	0.62	NAG 97	1	84	
Atopobium parvulum		10.65	0.46		99	1	
Clostridium botulinum biovar B		2.92	0.32		1	99	
07131141	**Low discrimination**						**07131141**
					SPO		
Clostridium chauvoei		76.85	0.44	TRE 10	99		
Actinomyces naeslundii		11.67	0.53	SOR 1	1		
Clostridium perfringens		8.30	0.28		84		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07131142	**Low discrimination**						**07131142**
					COC		
Clostridium perfringens		87.57	0.56	NAG 97	1		
Propionibacterium propionicum		5.59	0.48		1		
Atopobium parvulum		4.19	0.33		99		
07131160	**Very good identification**						**07131160**
					COC	SPO	
Clostridium perfringens		99.83	0.88		1	84	
Atopobium parvulum		0.07	0.33		99	1	
Clostridium botulinum biovar B		0.05	0.26		1	99	
07131161	**Low discrimination**						**07131161**
					SPO		
Clostridium perfringens		87.80	0.54	URE 1	84		
Clostridium chauvoei		8.38	0.35		99		
Actinomyces naeslundii		3.82	0.53		1		
07131162	**Very good identification**						**07131162**
					COC		
Clostridium perfringens		99.88	0.82		1		
Actinomyces naeslundii		0.06	0.50		1		
Atopobium parvulum		0.03	0.20		99		
07131163	**Low discrimination**						**07131163**
					SPO		
Clostridium perfringens		88.63	0.48	URE 1	84		
Actinomyces naeslundii		8.20	0.53		1		
Clostridium chauvoei		3.17	0.22		99		
07131170	**Good identification**						**07131170**
					SPO		
Clostridium perfringens		97.33	0.54	MLZ 1	84		
Clostridium botulinum biovar B		2.53	0.21		99		
Actinomyces naeslundii		0.15	0.28		1		
07131172	**Good identification**						**07131172**
Clostridium perfringens		99.68	0.48	MLZ 1			
Actinomyces naeslundii		0.32	0.28				
07131200	**Very good identification**						**07131200**
					COC	CAT	
Propionibacterium granulosum		96.44	0.66	RHA 3	1	90	
Gemella morbillorum		1.24	0.27		99	1	
Clostridium perfringens		1.03	0.25		1	1	
07131202	**Very good identification**						**07131202**
					CAT		
Propionibacterium granulosum		96.22	0.56	RHA 3	90		
Clostridium perfringens		1.92	0.20		1		
Lactobacillus jensenii		1.62	0.19		1		
07131210	**Good identification**						**07131210**
					CAT		
Propionibacterium granulosum		98.48	0.52	RHA 3	90		
Lactobacillus jensenii		1.52	0.14	MLZ 12	1		
07131212	**Good identification**						**07131212**
					CAT		
Propionibacterium granulosum		96.06	0.41	RHA 3	90		
Lactobacillus jensenii		3.94	0.11	MLZ 12	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07131220	**Low discrimination**						**07131220**
					CAT		
Clostridium perfringens		57.59	0.51	LAC 88	1		
Propionibacterium granulosum		41.91	0.56	RHA 1	90		
Lactobacillus jensenii		0.26	0.12		1		
07131222	**Low discrimination**						**07131222**
					GRA	CAT	
Clostridium perfringens		71.25	0.45	LAC 88	99	1	
Propionibacterium granulosum		27.55	0.46	RHA 1	99	90	
Prevotella oralis		0.74	0.08		1	1	
07131230	**Low discrimination**						**07131230**
					CAT		
Propionibacterium granulosum		89.51	0.41	RHA 3	90		
Clostridium perfringens		9.11	0.17	MLZ 12	1		
Lactobacillus jensenii		1.38	0.04		1		
07131260	**Low discrimination**						**07131260**
					CAT	SPO	
Clostridium perfringens		70.17	0.39	LAC 88	1	84	
Clostridium botulinum biovar B		27.12	0.26	RHA 1	1	99	
Propionibacterium granulosum		2.71	0.22		90	1	
07131300	**Low discrimination**						**07131300**
					COC	CAT	
Atopobium parvulum		66.36	0.46	RHA 1	99	1	
Clostridium perfringens		27.56	0.40		1	1	
Propionibacterium granulosum		3.57	0.32		1	90	
07131301	**Good identification**						**07131301**
					COC	SPO	
Clostridium chauvoei		97.26	0.44	TRE 10	1	99	
Atopobium parvulum		1.34	0.12	RHA 1	99	1	
Actinomyces naeslundii		0.85	0.31		1	1	
07131320	**Very good identification**						**07131320**
					COC	GRA	
Clostridium perfringens		98.12	0.66	RHA 1	1	99	
Atopobium parvulum		1.29	0.33		99	99	
Prevotella melaninogenica		0.28	0.15		1	1	
07131321	**Low discrimination**						**07131321**
					SPO		
Clostridium chauvoei		63.24	0.35	TRE 10	99		
Clostridium perfringens		35.12	0.32	RHA 1	84		
Actinomyces naeslundii		1.65	0.31		1		
07131322	**Good identification**						**07131322**
					GRA		
Clostridium perfringens		94.41	0.60	RHA 1	99		
Parabacteroides distasonis		2.52	0.43		1		
Prevotella oralis		2.09	0.29		1		
07131332	**Low discrimination**						**07131332**
					GRA		
Parabacteroides distasonis		57.22	0.38	RAF 99	1		
Clostridium perfringens		40.28	0.26	XYL 90	99		
Clostridium clostridioforme		2.51	0.09		50		
07131360	**Good identification**						**07131360**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		100.0	0.54	RHA 1			
07131362	**Good identification**						**07131362**
Clostridium perfringens		100.0	0.48	RHA 1			
07131400	**Low discrimination**						**07131400**
					COC	CAT	
Propionibacterium granulosum		59.04	0.58	ARA 1	1	90	
Gemella morbillorum		25.63	0.45		99	1	
Lactobacillus fermentum		10.98	0.50		1	1	
07131402	**Low discrimination**						**07131402**
					CAT		
Propionibacterium granulosum		53.20	0.48	ARA 1	90		
Lactobacillus fermentum		39.57	0.50		1		
Clostridium perfringens		3.26	0.20		1		
07131410	**Low discrimination**						**07131410**
					COC	CAT	
Propionibacterium granulosum		89.99	0.43	ARA 1	1	90	
Lactobacillus jensenii		4.24	0.14	MLZ 12	1	1	
Gemella morbillorum		2.89	0.11		99	1	
07131420	**Low discrimination**						**07131420**
					CAT		
Clostridium perfringens		73.93	0.51	LAC 88	1		
Propionibacterium granulosum		17.57	0.48	ARA 1	90		
Lactobacillus fermentum		4.36	0.42		1		
07131422	**Low discrimination**						**07131422**
					CAT		
Clostridium perfringens		78.76	0.45	LAC 88	1		
Propionibacterium granulosum		9.94	0.38	ARA 1	90		
Lactobacillus fermentum		9.86	0.42		1		
07131460	**Low discrimination**						**07131460**
					CAT	SPO	
Clostridium perfringens		71.48	0.39	LAC 88	1	84	
Clostridium botulinum biovar B		27.62	0.26	ARA 1	1	99	
Propionibacterium granulosum		0.90	0.14		90	1	
07131500	**Low discrimination**						**07131500**
					COC		
Atopobium parvulum		44.13	0.46	ARA 1	99		
Lactobacillus fermentum		23.31	0.54		1		
Clostridium perfringens		18.33	0.40		1		
07131501	**Good identification**						**07131501**
					COC	SPO	
Clostridium chauvoei		96.25	0.44	TRE 10	1	99	
Atopobium parvulum		1.33	0.12	ARA 1	99	1	
Actinomyces naeslundii		0.84	0.31		1	1	
07131502	**Low discrimination**						**07131502**
Lactobacillus fermentum		42.44	0.54	TRE 1			
Bifidobacterium longum subsp. longum		20.11	0.41				
Clostridium perfringens		15.70	0.34				
07131510	**Good identification**						**07131510**
					COC		
Bifidobacterium longum subsp. longum		95.09	0.45	RAF 97	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		2.25	0.12		99		
Lactobacillus fermentum		1.19	0.20		1		
07131512	**Low discrimination**						**07131512**
Bifidobacterium longum subsp. longum		79.27	0.45	RAF 97			
Bifidobacterium adolescentis		17.79	0.36				
Clostridium clostridioforme		1.58	0.15				
07131520	**Good identification**						**07131520**
Clostridium perfringens		90.43	0.66	ARA 1			
Bifidobacterium longum subsp. longum		6.74	0.52				
Lactobacillus fermentum		1.19	0.46				
07131521	**Low discrimination**						**07131521**
					SPO		
Clostridium chauvoei		61.62	0.35	TRE 10	99		
Clostridium perfringens		34.22	0.32	ARA 1	84		
Bifidobacterium longum subsp. longum		2.55	0.18		1		
07131522	**Low discrimination**						**07131522**
Clostridium perfringens		83.37	0.60	ARA 1			
Bifidobacterium longum subsp. longum		13.21	0.52				
Lactobacillus fermentum		2.32	0.46				
07131530	**Acceptable identification**						**07131530**
Bifidobacterium longum subsp. longum		92.63	0.56	RAF 97			
Clostridium perfringens		7.37	0.32				
07131532	**Acceptable identification**						**07131532**
					GRA		
Bifidobacterium longum subsp. longum		94.94	0.56	RAF 97	99		
Clostridium perfringens		3.56	0.26		99		
Parabacteroides distasonis		1.03	0.26		1		
07131560	**Good identification**						**07131560**
Clostridium perfringens		99.61	0.54	ARA 1			
Bifidobacterium longum subsp. longum		0.39	0.18				
07131562	**Good identification**						**07131562**
Clostridium perfringens		99.17	0.48	ARA 1			
Bifidobacterium longum subsp. longum		0.83	0.18				
07131722	**Low discrimination**						**07131722**
					GRA		
Parabacteroides distasonis		54.08	0.39	RAF 99	1		
Clostridium perfringens		33.45	0.26	XYL 90	99		
Clostridium clostridioforme		7.17	0.18		50		
07132000	**Very good identification**						**07132000**
					SPO	CAT	
Propionibacterium granulosum		99.47	0.90		1	90	
Clostridium botulinum biovar B		0.21	0.32		99	1	
Propionibacterium propionicum		0.12	0.45		1	1	
07132001	**Good identification**						**07132001**
					CAT		
Propionibacterium granulosum		97.18	0.56	URE 1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		1.33	0.39		25		
Propionibacterium propionicum		1.29	0.29		1		
07132002	**Very good identification**						**07132002**
					CAT		
Propionibacterium granulosum		99.20	0.80		90		
Propionibacterium propionicum		0.72	0.48		1		
Actinomyces naeslundii		0.04	0.36		25		
07132003	**Low discrimination**						**07132003**
					CAT		
Propionibacterium granulosum		88.17	0.46	URE 1	90		
Propionibacterium propionicum		7.02	0.32		1		
Actinomyces naeslundii		4.81	0.39		25		
07132010	**Very good identification**						**07132010**
					SPO	CAT	
Propionibacterium granulosum		99.21	0.75	MLZ 12	1	90	
Clostridium botulinum biovar B		0.75	0.26		99	1	
Clostridium difficile		0.04	0.04		99	1	
07132011	**Good identification**						**07132011**
Propionibacterium granulosum		100.0	0.41	MLZ 12			
07132012	**Very good identification**						**07132012**
					CAT		
Propionibacterium granulosum		99.95	0.65	MLZ 12	90		
Propionibacterium propionicum		0.05	0.14		1		
07132020	**Very good identification**						**07132020**
					CAT	SPO	
Propionibacterium granulosum		99.45	0.80		90	1	
Clostridium botulinum biovar B		0.41	0.26		1	99	
Clostridium perfringens		0.06	0.17		1	84	
07132021	**Acceptable identification**						**07132021**
Propionibacterium granulosum		94.82	0.46	URE 1			
Actinomyces naeslundii		5.18	0.39				
07132022	**Very good identification**						**07132022**
					CAT		
Propionibacterium granulosum		99.72	0.70		90		
Actinomyces naeslundii		0.15	0.36		25		
Clostridium perfringens		0.11	0.11		1		
07132023	**Low discrimination**						**07132023**
Propionibacterium granulosum		82.08	0.35	URE 1			
Actinomyces naeslundii		17.92	0.39				
07132030	**Very good identification**						**07132030**
					SPO	CAT	
Propionibacterium granulosum		98.52	0.65	MLZ 12	1	90	
Clostridium botulinum biovar B		1.48	0.21		99	1	
07132032	**Good identification**						**07132032**
Propionibacterium granulosum		100.0	0.55	MLZ 12			
07132040	**Good identification**						**07132040**
					SPO	CAT	
Clostridium botulinum biovar B		94.61	0.66	MAN 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		4.59	0.56		1	90	
Propionibacterium propionicum		0.55	0.45		1	1	
07132042	**Low discrimination**						**07132042**
					SPO	CAT	
Propionibacterium granulosum		38.76	0.46	SOR 1	1	90	
Clostridium botulinum biovar B		32.31	0.32		99	1	
Propionibacterium propionicum		27.76	0.48		1	1	
07132050	**Very good identification**						**07132050**
					SPO	CAT	
Clostridium botulinum biovar B		98.68	0.61	MAN 1	99	1	
Propionibacterium granulosum		1.32	0.41		1	90	
07132060	**Very good identification**						**07132060**
					CAT	SPO	
Clostridium botulinum biovar B		97.56	0.61	MAN 1	1	99	
Propionibacterium granulosum		2.40	0.46		90	1	
Clostridium perfringens		0.03	0.05		1	84	
07132070	**Good identification**						**07132070**
					SPO	CAT	
Clostridium botulinum biovar B		99.32	0.55	MAN 1	99	1	
Propionibacterium granulosum		0.68	0.31		1	90	
07132100	**Good identification**						**07132100**
					CAT		
Propionibacterium propionicum		91.84	0.79	NIT 86	1		
Propionibacterium granulosum		7.77	0.56		90		
Actinomyces naeslundii		0.20	0.44		25		
07132101	**Very good identification**						**07132101**
					CAT		
Propionibacterium propionicum		96.01	0.63	NIT 86	1		
Actinomyces naeslundii		2.84	0.47	URE 10	25		
Propionibacterium granulosum		0.74	0.22		90		
07132102	**Very good identification**						**07132102**
					CAT		
Propionibacterium propionicum		98.46	0.82	NIT 86	1		
Propionibacterium granulosum		1.39	0.46		90		
Actinomyces naeslundii		0.14	0.44		25		
07132103	**Very good identification**						**07132103**
					CAT		
Propionibacterium propionicum		97.94	0.66	NIT 86	1		
Actinomyces naeslundii		1.93	0.47	URE 10	25		
Propionibacterium granulosum		0.13	0.12		90		
07132110	**Low discrimination**						**07132110**
					CAT		
Propionibacterium granulosum		53.33	0.41	LAC 1	90		
Propionibacterium propionicum		46.67	0.45	MLZ 12	1		
07132112	**Low discrimination**						**07132112**
					CAT		
Propionibacterium propionicum		84.00	0.48	NIT 86	1		
Propionibacterium granulosum		16.00	0.31	MLZ 1	90		
07132120	**Low discrimination**						**07132120**
					CAT		
Propionibacterium granulosum		50.15	0.46	LAC 1	90		
Propionibacterium propionicum		23.94	0.45		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		20.72	0.32		1		
07132121	**Low discrimination**						**07132121**
					CAT		
Actinomyces naeslundii		68.74	0.47	RAF 92	25		
Propionibacterium propionicum		23.47	0.29	GAL 96	1		
Propionibacterium granulosum		4.47	0.12		90		
07132122	**Low discrimination**						**07132122**
					CAT		
Propionibacterium propionicum		56.65	0.48	NIT 86	1		
Propionibacterium granulosum		19.78	0.35	NAG 1	90		
Clostridium perfringens		15.38	0.26		1		
07132123	**Low discrimination**						**07132123**
Actinomyces naeslundii		66.13	0.47	RAF 92			
Propionibacterium propionicum		33.87	0.32	GAL 96			
07132140	**Very good identification**						**07132140**
					SPO	CAT	
Propionibacterium propionicum		98.15	0.79	NIT 86	1	1	
Clostridium botulinum biovar B		1.73	0.32		99	1	
Propionibacterium granulosum		0.08	0.22		1	90	
07132141	**Very good identification**						**07132141**
					SPO		
Propionibacterium propionicum		99.32	0.63	NIT 86	1		
Actinomyces naeslundii		0.52	0.34	URE 10	1		
Clostridium botulinum biovar B		0.16	-.03		99		
07132142	**Very good identification**						**07132142**
					CAT		
Propionibacterium propionicum		99.95	0.82	NIT 86	1		
Actinomyces naeslundii		0.03	0.31		25		
Propionibacterium granulosum		0.01	0.12		90		
07132143	**Very good identification**						**07132143**
Propionibacterium propionicum		99.65	0.66	NIT 86			
Actinomyces naeslundii		0.35	0.34	URE 10			
07132150	**Low discrimination**						**07132150**
					SPO		
Propionibacterium propionicum		53.77	0.45	NIT 86	1		
Clostridium botulinum biovar B		46.23	0.26	MLZ 1	99		
07132152	**Good identification**						**07132152**
Propionibacterium propionicum		100.0	0.48	NIT 86			
07132160	**Low discrimination**						**07132160**
					SPO		
Propionibacterium propionicum		47.98	0.45	NIT 86	1		
Clostridium botulinum biovar B		41.26	0.26	NAG 1	99		
Clostridium perfringens		7.91	0.19		84		
07132162	**Acceptable identification**						**07132162**
Propionibacterium propionicum		92.83	0.48	NIT 86			
Clostridium perfringens		4.80	0.14	NAG 1			
Actinomyces naeslundii		2.37	0.31				
07132200	**Very good identification**						**07132200**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	CAT		
Propionibacterium granulosum		99.93	0.64	RHA 3	1	90		
Clostridium botulinum biovar B		0.07	-.03		99	1		
07132202	**Good identification**							**07132202**
					CAT			
Propionibacterium granulosum		99.76	0.54	RHA 3	90			
Propionibacterium propionicum		0.24	0.14		1			
07132210	**Good identification**							**07132210**
Propionibacterium granulosum		100.0	0.50	RHA 3				
07132212	**Good identification**							**07132212**
Propionibacterium granulosum		100.0	0.39	RHA 3				
07132220	**Good identification**							**07132220**
Propionibacterium granulosum		100.0	0.54	RHA 3				
07132222	**Good identification**							**07132222**
Propionibacterium granulosum		100.0	0.44	RHA 3				
07132230	**Good identification**							**07132230**
Propionibacterium granulosum		100.0	0.39	RHA 3				
07132300	**Low discrimination**							**07132300**
					CAT			
Propionibacterium propionicum		79.42	0.45	NIT 86	1			
Propionibacterium granulosum		20.58	0.30	RHA 1	90			
07132302	**Good identification**							**07132302**
					CAT			
Propionibacterium propionicum		95.86	0.48	NIT 86	1			
Propionibacterium granulosum		4.14	0.20	RHA 1	90			
07132340	**Good identification**							**07132340**
					SPO			
Propionibacterium propionicum		98.27	0.45	NIT 86	1			
Clostridium botulinum biovar B		1.73	-.03	RHA 1	99			
07132342	**Good identification**							**07132342**
Propionibacterium propionicum		100.0	0.48	NIT 86				
07132400	**Good identification**							**07132400**
					COC	SPO	CAT	
Propionibacterium granulosum		98.73	0.56	ARA 1	1	1	90	
Gemella morbillorum		1.07	0.15		99	1	1	
Clostridium botulinum biovar B		0.21	-.03		1	99	1	
07132402	**Good identification**							**07132402**
					CAT			
Propionibacterium granulosum		99.28	0.46	ARA 1	90			
Propionibacterium propionicum		0.72	0.14		1			
07132410	**Good identification**							**07132410**
Propionibacterium granulosum		100.0	0.41	ARA 1				
07132420	**Good identification**							**07132420**
Propionibacterium granulosum		100.0	0.46	ARA 1				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07132422	**Good identification**					**07132422**
Propionibacterium granulosum		100.0	0.35	ARA 1		
07132500	**Acceptable identification**					**07132500**
					CAT	
Propionibacterium propionicum		92.20	0.45	NIT 86	1	
Propionibacterium granulosum		7.80	0.22	ARA 1	90	
07132502	**Good identification**					**07132502**
					CAT	
Propionibacterium propionicum		98.61	0.48	NIT 86	1	
Propionibacterium granulosum		1.39	0.12	ARA 1	90	
07132540	**Good identification**					**07132540**
					SPO	
Propionibacterium propionicum		98.27	0.45	NIT 86	1	
Clostridium botulinum biovar B		1.73	-.03	ARA 1	99	
07132542	**Good identification**					**07132542**
Propionibacterium propionicum		100.0	0.48	NIT 86		
07133000	**Very good identification**					**07133000**
					CAT	
Propionibacterium granulosum		99.51	0.90		90	
Actinomyces naeslundii		0.22	0.60		25	
Propionibacterium propionicum		0.12	0.45		1	
07133001	**Low discrimination**					**07133001**
					CAT	
Propionibacterium granulosum		74.27	0.56	URE 1	90	
Actinomyces naeslundii		24.33	0.63		25	
Propionibacterium propionicum		0.99	0.29		1	
07133002	**Very good identification**					**07133002**
					CAT	
Propionibacterium granulosum		98.25	0.80		90	
Actinomyces naeslundii		0.87	0.60		25	
Propionibacterium propionicum		0.71	0.48		1	
07133003	**Low discrimination**					**07133003**
					CAT	
Actinomyces naeslundii		54.83	0.63	RAF 92	25	
Propionibacterium granulosum		41.84	0.46		90	
Propionibacterium propionicum		3.33	0.32		1	
07133010	**Very good identification**					**07133010**
					CAT	
Propionibacterium granulosum		99.85	0.75	MLZ 12	90	
Actinomyces naeslundii		0.09	0.38		25	
Lactobacillus jensenii		0.06	0.14		1	
07133011	**Low discrimination**					**07133011**
Propionibacterium granulosum		88.78	0.41	MLZ 12		
Actinomyces naeslundii		11.22	0.41	URE 1		
07133012	**Very good identification**					**07133012**
					CAT	
Propionibacterium granulosum		99.44	0.65	MLZ 12	90	
Actinomyces naeslundii		0.34	0.38		25	
Lactobacillus jensenii		0.17	0.11		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07133020	**Good identification**						**07133020**
					CAT		
Propionibacterium granulosum		93.84	0.80		90		
Clostridium perfringens		5.29	0.51		1		
Actinomyces naeslundii		0.83	0.60		25		
07133021	**Low discrimination**						**07133021**
					CAT		
Actinomyces naeslundii		55.19	0.63	RAF 92	25		
Propionibacterium granulosum		42.12	0.46		90		
Clostridium perfringens		2.37	0.17		1		
07133022	**Low discrimination**						**07133022**
					CAT		
Propionibacterium granulosum		87.54	0.70		90		
Clostridium perfringens		9.28	0.45		1		
Actinomyces naeslundii		3.09	0.60		25		
07133023	**Low discrimination**						**07133023**
					CAT		
Actinomyces naeslundii		82.57	0.63	RAF 92	25		
Propionibacterium granulosum		15.76	0.35		90		
Clostridium perfringens		1.67	0.11		1		
07133030	**Very good identification**						**07133030**
					CAT		
Propionibacterium granulosum		99.19	0.65	MLZ 12	90		
Clostridium perfringens		0.41	0.17		1		
Actinomyces naeslundii		0.34	0.38		25		
07133032	**Good identification**						**07133032**
					CAT		
Propionibacterium granulosum		97.90	0.55	MLZ 12	90		
Actinomyces naeslundii		1.33	0.38		25		
Clostridium perfringens		0.77	0.11		1		
07133033	**Low discrimination**						**07133033**
Actinomyces naeslundii		66.92	0.41	RAF 92			
Propionibacterium granulosum		33.08	0.21	MLZ 5			
07133040	**Low discrimination**						**07133040**
					SPO	CAT	
Propionibacterium granulosum		72.76	0.56	SOR 1	1	90	
Clostridium botulinum biovar B		15.16	0.32		99	1	
Propionibacterium propionicum		8.68	0.45		1	1	
07133041	**Low discrimination**						**07133041**
					CAT		
Actinomyces naeslundii		69.42	0.50	RAF 92	25		
Propionibacterium propionicum		15.93	0.29		1		
Propionibacterium granulosum		12.13	0.22		90		
07133042	**Low discrimination**						**07133042**
					CAT		
Propionibacterium granulosum		52.79	0.46	SOR 1	90		
Propionibacterium propionicum		37.81	0.48		1		
Actinomyces naeslundii		8.14	0.47		25		
07133043	**Low discrimination**						**07133043**
					CAT		
Actinomyces naeslundii		72.06	0.50	RAF 92	25		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		24.80	0.32		1		
Propionibacterium granulosum		3.15	0.12		90		
07133050	**Low discrimination**						**07133050**
					SPO	CAT	
Propionibacterium granulosum		56.58	0.41	MLZ 12	1	90	
Clostridium botulinum biovar B		42.58	0.26	SOR 1	99	1	
Actinomyces naeslundii		0.84	0.25		1	25	
07133060	**Low discrimination**						**07133060**
					CAT	SPO	
Clostridium perfringens		40.44	0.39	MAN 1	1	84	
Propionibacterium granulosum		38.07	0.46	LAC 88	90	1	
Clostridium botulinum biovar B		15.63	0.26		1	99	
07133061	**Acceptable identification**						**07133061**
					CAT		
Actinomyces naeslundii		91.74	0.50	RAF 92	25		
Clostridium perfringens		4.26	0.05		1		
Propionibacterium granulosum		4.01	0.12		90		
07133063	**Good identification**						**07133063**
Actinomyces naeslundii		100.0	0.50	RAF 92			
07133100	**Low discrimination**						**07133100**
					COC	CAT	
Propionibacterium propionicum		81.20	0.79	NIT 86	1	1	
Propionibacterium granulosum		6.87	0.56		1	90	
Atopobium parvulum		5.23	0.46		99	1	
07133101	**Low discrimination**						**07133101**
					SPO		
Propionibacterium propionicum		46.49	0.63	NIT 86	1		
Actinomyces naeslundii		33.01	0.70	URE 10	1		
Clostridium chauvoei		19.76	0.44		99		
07133102	**Good identification**						**07133102**
					CAT		
Propionibacterium propionicum		93.69	0.82	NIT 86	1		
Actinomyces naeslundii		3.29	0.67		25		
Propionibacterium granulosum		1.32	0.46		90		
07133103	**Low discrimination**						**07133103**
					SPO		
Propionibacterium propionicum		65.55	0.66	NIT 86	1		
Actinomyces naeslundii		31.03	0.70	URE 10	1		
Clostridium chauvoei		3.28	0.31		99		
07133110	**Low discrimination**						**07133110**
					CAT		
Propionibacterium granulosum		45.07	0.41	LAC 1	90		
Propionibacterium propionicum		39.44	0.45	MLZ 12	1		
Actinomyces naeslundii		10.81	0.45		25		
07133111	**Low discrimination**						**07133111**
					SPO		
Actinomyces naeslundii		72.20	0.49	RAF 92	1		
Propionibacterium propionicum		19.51	0.29	MLZ 5	1		
Clostridium chauvoei		8.29	0.09		99		
07133112	**Low discrimination**						**07133112**
					CAT		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		66.75	0.48	NIT 86	1	
Propionibacterium granulosum		12.71	0.31	MLZ 1	90	
Actinomyces naeslundii		12.20	0.45		25	
07133113	**Low discrimination**					**07133113**
Actinomyces naeslundii		71.15	0.49	RAF 92		
Propionibacterium propionicum		28.85	0.32	MLZ 5		
07133120	**Low discrimination**					**07133120**
					CAT	
Clostridium perfringens		89.69	0.66	MAN 1	1	
Actinomyces naeslundii		5.45	0.67		25	
Propionibacterium granulosum		2.19	0.46		90	
07133121	**Low discrimination**					**07133121**
					SPO	
Actinomyces naeslundii		75.35	0.70	RAF 92	1	
Clostridium chauvoei		15.03	0.35		99	
Clostridium perfringens		8.35	0.32		84	
07133122	**Low discrimination**					**07133122**
Clostridium perfringens		83.66	0.60	MAN 1		
Actinomyces naeslundii		10.81	0.67			
Propionibacterium propionicum		3.11	0.48			
07133123	**Good identification**					**07133123**
					SPO	
Actinomyces naeslundii		90.20	0.70	RAF 92	1	
Clostridium perfringens		4.70	0.26		84	
Clostridium chauvoei		3.18	0.22		99	
07133131	**Good identification**					**07133131**
					SPO	
Actinomyces naeslundii		96.31	0.49	RAF 92	1	
Clostridium chauvoei		3.69	0.01	MLZ 5	99	
07133133	**Good identification**					**07133133**
Actinomyces naeslundii		100.0	0.49	RAF 92		
07133140	**Very good identification**					**07133140**
Propionibacterium propionicum		98.39	0.79	NIT 86		
Actinomyces naeslundii		0.91	0.54			
Clostridium perfringens		0.50	0.28			
07133141	**Low discrimination**					**07133141**
					SPO	
Propionibacterium propionicum		88.53	0.63	NIT 86	1	
Actinomyces naeslundii		11.09	0.57	URE 10	1	
Clostridium chauvoei		0.38	0.09		99	
07133142	**Very good identification**					**07133142**
Propionibacterium propionicum		99.18	0.82	NIT 86		
Actinomyces naeslundii		0.61	0.54			
Clostridium perfringens		0.16	0.22			
07133143	**Good identification**					**07133143**
Propionibacterium propionicum		92.29	0.66	NIT 86		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		7.71	0.57	URE 10		
07133150	**Good identification**					**07133150**
Propionibacterium propionicum		95.39	0.45	NIT 86		
Actinomyces naeslundii		4.61	0.33	MLZ 1		
07133152	**Good identification**					**07133152**
Propionibacterium propionicum		96.88	0.48	NIT 86		
Actinomyces naeslundii		3.12	0.33	MLZ 1		
07133160	**Low discrimination**					**07133160**
Clostridium perfringens		89.37	0.54	MAN 1		
Propionibacterium propionicum		5.48	0.45			
Actinomyces naeslundii		5.03	0.54			
07133161	**Low discrimination**					**07133161**
Actinomyces naeslundii		82.54	0.57	RAF 92		
Clostridium perfringens		9.87	0.19			
Propionibacterium propionicum		6.65	0.29			
07133162	**Low discrimination**					**07133162**
Clostridium perfringens		76.05	0.48	MAN 1		
Propionibacterium propionicum		14.85	0.48			
Actinomyces naeslundii		9.10	0.54			
07133163	**Low discrimination**					**07133163**
Actinomyces naeslundii		84.95	0.57	RAF 92		
Propionibacterium propionicum		10.27	0.32			
Clostridium perfringens		4.78	0.14			
07133171	**Good identification**					**07133171**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07133173	**Good identification**					**07133173**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07133200	**Very good identification**					**07133200**
Propionibacterium granulosum		99.93	0.64	RHA 3		
Actinomyces naeslundii		0.07	0.25			
07133202	**Good identification**					**07133202**
					CAT	
Propionibacterium granulosum		99.48	0.54	RHA 3	90	
Actinomyces naeslundii		0.29	0.25		25	
Propionibacterium propionicum		0.24	0.14		1	
07133210	**Good identification**					**07133210**
Propionibacterium granulosum		100.0	0.50	RHA 3		
07133212	**Good identification**					**07133212**
Propionibacterium granulosum		100.0	0.39	RHA 3		
07133220	**Good identification**					**07133220**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		97.92	0.54	RHA 3	90		
Clostridium perfringens		1.80	0.17		1		
Actinomyces naeslundii		0.28	0.25		25		
07133222	**Good identification**						**07133222**
					CAT		
Propionibacterium granulosum		95.59	0.44	RHA 3	90		
Clostridium perfringens		3.31	0.11		1		
Actinomyces naeslundii		1.10	0.25		25		
07133230	**Good identification**						**07133230**
Propionibacterium granulosum		100.0	0.39	RHA 3			
07133300	**Low discrimination**						**07133300**
					COC	CAT	
Propionibacterium propionicum		71.28	0.45	NIT 86	1	1	
Propionibacterium granulosum		18.47	0.30	RHA 1	1	90	
Atopobium parvulum		4.59	0.12		99	1	
07133302	**Acceptable identification**						**07133302**
					CAT		
Propionibacterium propionicum		92.03	0.48	NIT 86	1		
Propionibacterium granulosum		3.97	0.20	RHA 1	90		
Actinomyces naeslundii		3.23	0.33		25		
07133321	**Low discrimination**						**07133321**
					SPO		
Actinomyces naeslundii		83.37	0.36	RAF 92	1		
Clostridium chauvoei		16.63	0.01	RHA 1	99		
07133323	**Good identification**						**07133323**
Actinomyces naeslundii		100.0	0.36	RAF 92			
07133340	**Good identification**						**07133340**
Propionibacterium propionicum		100.0	0.45	NIT 86			
07133342	**Good identification**						**07133342**
Propionibacterium propionicum		100.0	0.48	NIT 86			
07133400	**Good identification**						**07133400**
					COC	CAT	
Propionibacterium granulosum		98.77	0.56	ARA 1	1	90	
Gemella morbillorum		0.57	0.11		99	1	
Lactobacillus fermentum		0.25	0.16		1	1	
07133402	**Good identification**						**07133402**
					CAT		
Propionibacterium granulosum		96.38	0.46	ARA 1	90		
Actinomyces israelii		1.12	0.18		1		
Lactobacillus fermentum		0.96	0.16		1		
07133410	**Good identification**						**07133410**
Propionibacterium granulosum		100.0	0.41	ARA 1			
07133420	**Acceptable identification**						**07133420**
					CAT		
Propionibacterium granulosum		93.88	0.46	ARA 1	90		
Clostridium perfringens		5.29	0.17		1		
Actinomyces naeslundii		0.83	0.25		25		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07133422	**Low discrimination**							**07133422**
					CAT			
Propionibacterium granulosum		87.62	0.35	ARA 1	90			
Clostridium perfringens		9.29	0.11		1			
Actinomyces naeslundii		3.09	0.25		25			
07133500	**Low discrimination**							**07133500**
					CAT			
Propionibacterium propionicum		74.44	0.45	NIT 86	1			
Propionibacterium granulosum		6.30	0.22	ARA 1	90			
Actinomyces israelii		4.82	0.26		1			
07133502	**Low discrimination**							**07133502**
Propionibacterium propionicum		70.03	0.48	NIT 86				
Bifidobacterium adolescentis		19.07	0.33	ARA 1				
Actinomyces israelii		4.53	0.29					
07133521	**Low discrimination**							**07133521**
					SPO			
Actinomyces naeslundii		83.37	0.36	RAF 92	1			
Clostridium chauvoei		16.63	0.01	ARA 1	99			
07133523	**Good identification**							**07133523**
Actinomyces naeslundii		100.0	0.36	RAF 92				
07133540	**Good identification**							**07133540**
Propionibacterium propionicum		97.54	0.45	NIT 86				
Actinomyces israelii		2.46	0.19	ARA 1				
07133542	**Acceptable identification**							**07133542**
Propionibacterium propionicum		94.44	0.48	NIT 86				
Bifidobacterium adolescentis		3.18	0.17	ARA 1				
Actinomyces israelii		2.38	0.22					
07134000	**Low discrimination**							**07134000**
					COC	SPO	CAT	
Propionibacterium granulosum		49.51	0.58	XYL 1	1	1	90	
Gemella morbillorum		39.92	0.50		99	1	1	
Clostridium botulinum biovar B		7.78	0.32		1	99	1	
07134002	**Acceptable identification**							**07134002**
					COC	GRA	CAT	
Propionibacterium granulosum		94.14	0.48	XYL 1	1	99	90	
Gemella morbillorum		3.07	0.15		99	99	1	
Fusobacterium mortiferum		1.06	0.18		1	1	5	
07134010	**Low discrimination**							**07134010**
					COC	SPO	CAT	
Propionibacterium granulosum		61.44	0.43	XYL 1	1	1	90	
Clostridium botulinum biovar B		34.89	0.26	MLZ 12	1	99	1	
Gemella morbillorum		3.67	0.15		99	1	1	
07134020	**Low discrimination**							**07134020**
					COC	SPO	CAT	
Propionibacterium granulosum		58.01	0.48	XYL 1	1	1	90	
Gemella morbillorum		20.78	0.33		99	1	1	
Clostridium botulinum biovar B		17.97	0.26		1	99	1	
07134022	**Acceptable identification**							**07134022**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					CAT			
Propionibacterium granulosum		92.59	0.38	XYL 1	90			
Clostridium perfringens		7.41	0.11		1			
07134040	**Very good identification**							**07134040**
					COC	SPO	CAT	
Clostridium botulinum biovar B		99.88	0.66	XYL 1	1	99	1	
Propionibacterium granulosum		0.07	0.24		1	1	90	
Gemella morbillorum		0.05	0.15		99	1	1	
07134050	**Very good identification**							**07134050**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.61	XYL 1	99	1		
Propionibacterium granulosum		0.02	0.09		1	90		
07134060	**Very good identification**							**07134060**
					CAT	SPO		
Clostridium botulinum biovar B		99.94	0.61	XYL 1	1	99		
Propionibacterium granulosum		0.03	0.14		90	1		
Clostridium perfringens		0.03	0.05		1	84		
07134070	**Good identification**							**07134070**
Clostridium botulinum biovar B		100.0	0.55	XYL 1				
07134100	**Low discrimination**							**07134100**
					COC	GRA		
Atopobium parvulum		89.43	0.50	GAL 99	99	99		
Fusobacterium mortiferum		4.96	0.40		1	1		
Actinomyces meyeri		3.59	0.29		1	99		
07134102	**Acceptable identification**							**07134102**
					COC	GRA	CAT	
Atopobium parvulum		92.74	0.38	GAL 99	99	99	1	
Fusobacterium mortiferum		3.24	0.24		1	1	5	
Propionibacterium granulosum		1.24	0.14		1	99	90	
07134120	**Low discrimination**							**07134120**
					COC			
Atopobium parvulum		68.57	0.38	GAL 99	99			
Clostridium perfringens		28.38	0.32		1			
Actinomyces meyeri		2.13	0.14		1			
07134712	**Good identification**							**07134712**
Clostridium clostridioforme		100.0	0.39	SAL 88				
07134732	**Good identification**							**07134732**
Clostridium clostridioforme		97.65	0.39	SAL 88				
Parabacteroides distasonis		2.35	0.17	RAF 94				
07135000	**Low discrimination**							**07135000**
					COC	CAT		
Atopobium parvulum		39.12	0.50	LAC 99	99	1		
Propionibacterium granulosum		36.67	0.58		1	90		
Gemella morbillorum		15.92	0.45		99	1		
07135002	**Low discrimination**							**07135002**
					COC	CAT		
Propionibacterium granulosum		42.40	0.48	XYL 1	1	90		
Atopobium parvulum		31.92	0.38		99	1		
Lactobacillus fermentum		18.52	0.46		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07135010	**Low discrimination**						**07135010**
					COC	CAT	
Propionibacterium granulosum		85.73	0.43	XYL 1	1	90	
Atopobium parvulum		6.77	0.16	MLZ 12	99	1	
Lactobacillus jensenii		4.04	0.14		1	1	
07135020	**Low discrimination**						**07135020**
					COC	CAT	
Clostridium perfringens		66.50	0.51	XYL 1	1	1	
Propionibacterium granulosum		15.81	0.48	LAC 88	1	90	
Atopobium parvulum		11.90	0.38		99	1	
07135022	**Low discrimination**						**07135022**
					CAT		
Clostridium perfringens		77.98	0.45	XYL 1	1		
Propionibacterium granulosum		9.85	0.38	LAC 88	90		
Lactobacillus fermentum		5.74	0.38		1		
07135060	**Low discrimination**						**07135060**
					CAT	SPO	
Clostridium perfringens		71.00	0.39	XYL 1	1	84	
Clostridium botulinum biovar B		27.44	0.26	LAC 88	1	99	
Propionibacterium granulosum		0.90	0.14		90	1	
07135100	**Very good identification**						**07135100**
					COC		
Atopobium parvulum		99.42	0.85		99		
Clostridium perfringens		0.23	0.40		1		
Lactobacillus fermentum		0.17	0.50		1		
07135101	**Low discrimination**						**07135101**
					COC	SPO	
Atopobium parvulum		71.33	0.50	URE 1	99	1	
Clostridium chauvoei		28.15	0.44		1	99	
Actinomyces naeslundii		0.25	0.31		1	1	
07135102	**Very good identification**						**07135102**
					COC		
Atopobium parvulum		97.98	0.72		99		
Lactobacillus fermentum		0.94	0.50		1		
Clostridium perfringens		0.59	0.34		1		
07135103	**Low discrimination**						**07135103**
					COC	SPO	
Atopobium parvulum		69.95	0.38	URE 1	99	1	
Clostridium chauvoei		27.60	0.31		1	99	
Actinomyces naeslundii		1.36	0.31		1	1	
07135110	**Good identification**						**07135110**
					COC		
Atopobium parvulum		98.44	0.50	MLZ 1	99		
Bifidobacterium longum subsp. longum		0.69	0.19		1		
Clostridium perfringens		0.22	0.06		1		
07135112	**Low discrimination**						**07135112**
					COC		
Atopobium parvulum		71.98	0.38	MLZ 1	99		
Bifidobacterium adolescentis		21.01	0.36		1		
Bifidobacterium longum subsp. longum		2.84	0.19		1		
07135120	**Low discrimination**						**07135120**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		70.36	0.72		99		
Clostridium perfringens		29.12	0.66		1		
Lactobacillus fermentum		0.22	0.42		1		
07135121	**Low discrimination**						**07135121**
					COC	SPO	
Atopobium parvulum		45.90	0.38	URE 1	99	1	
Clostridium chauvoei		34.21	0.35		1	99	
Clostridium perfringens		19.00	0.32		1	84	
07135122	**Low discrimination**						**07135122**
					COC	GRA	
Clostridium perfringens		49.57	0.60	XYL 1	1	99	
Atopobium parvulum		44.91	0.59		99	99	
Parabacteroides distasonis		3.96	0.51		1	1	
07135130	**Low discrimination**						**07135130**
					COC		
Atopobium parvulum		63.29	0.38	MLZ 1	99		
Clostridium perfringens		26.20	0.32		1		
Bifidobacterium longum subsp. longum		9.98	0.30		1		
07135132	**Low discrimination**						**07135132**
					COC	GRA	
Parabacteroides distasonis		60.34	0.46	RAF 99	1	1	
Clostridium perfringens		14.16	0.26		1	99	
Atopobium parvulum		12.83	0.25		99	99	
07135140	**Good identification**						**07135140**
					COC		
Atopobium parvulum		95.14	0.50	SOR 1	99		
Clostridium perfringens		4.05	0.28		1		
Actinomyces israelii		0.50	0.25		1		
07135142	**Low discrimination**						**07135142**
					COC		
Atopobium parvulum		83.01	0.38	SOR 1	99		
Clostridium perfringens		9.43	0.22		1		
Actinomyces israelii		3.72	0.28		1		
07135160	**Low discrimination**						**07135160**
					COC		
Clostridium perfringens		88.64	0.54	XYL 1	1		
Atopobium parvulum		11.36	0.38		99		
07135162	**Good identification**						**07135162**
					COC	GRA	
Clostridium perfringens		95.03	0.48	XYL 1	1	99	
Atopobium parvulum		4.57	0.25		99	99	
Parabacteroides distasonis		0.40	0.17		1	1	
07135300	**Good identification**						**07135300**
					COC		
Atopobium parvulum		99.61	0.50	RHA 1	99		
Clostridium perfringens		0.23	0.06		1		
Lactobacillus fermentum		0.17	0.16		1		
07135302	**Acceptable identification**						**07135302**
					COC	GRA	
Atopobium parvulum		91.08	0.38	RHA 1	99	99	
Parabacteroides distasonis		4.26	0.25		1	1	
Clostridium clostridioforme		2.46	0.15		1	50	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07135320	**Low discrimination**						**07135320**
					COC	GRA	
Atopobium parvulum		49.25	0.38	RHA 1	99	99	
Bacteroides vulgatus		28.07	0.25		1	1	
Clostridium perfringens		20.38	0.32		1	99	
07135322	**Good identification**						**07135322**
					COC	GRA	
Parabacteroides distasonis		91.12	0.59	RAF 99	1	1	
Clostridium perfringens		3.84	0.26		1	99	
Atopobium parvulum		3.48	0.25		99	99	
07135332	**Good identification**						**07135332**
Parabacteroides distasonis		95.30	0.54	RAF 99			
Clostridium clostridioforme		4.70	0.26				
07135400	**Low discrimination**						**07135400**
					COC		
Lactobacillus fermentum		57.59	0.46	TRE 1	1		
Gemella morbillorum		25.43	0.29		99		
Atopobium parvulum		5.68	0.16		99		
07135402	**Low discrimination**						**07135402**
Lactobacillus fermentum		84.85	0.46	TRE 1			
Actinomyces israelii		3.45	0.24				
Bifidobacterium longum subsp. longum		3.39	0.15				
07135420	**Low discrimination**						**07135420**
Lactobacillus fermentum		48.00	0.38	TRE 1			
Bifidobacterium longum subsp. longum		22.98	0.26				
Clostridium perfringens		14.01	0.17				
07135422	**Low discrimination**						**07135422**
Lactobacillus fermentum		57.00	0.38	TRE 1			
Bifidobacterium longum subsp. longum		27.29	0.26				
Clostridium perfringens		7.83	0.11				
07135500	**Low discrimination**						**07135500**
					COC		
Atopobium parvulum		62.28	0.50	ARA 1	99		
Bifidobacterium longum subsp. longum		25.18	0.50		1		
Lactobacillus fermentum		10.40	0.50		1		
07135502	**Low discrimination**						**07135502**
					COC		
Bifidobacterium longum subsp. longum		38.17	0.50	RAF 97	1		
Bifidobacterium adolescentis		26.62	0.49		1		
Atopobium parvulum		16.66	0.38		99		
07135510	**Good identification**						**07135510**
					COC		
Bifidobacterium longum subsp. longum		97.60	0.54	RAF 97	1		
Atopobium parvulum		1.43	0.16		99		
Bifidobacterium adolescentis		0.49	0.16		1		
07135512	**Low discrimination**						**07135512**
Bifidobacterium longum subsp. longum		63.43	0.54	RAF 97			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		31.76	0.50				
Clostridium clostridioforme		4.27	0.32				
07135520	**Low discrimination**						**07135520**
					COC	GRA	
Bifidobacterium longum subsp. longum		77.95	0.60	RAF 97	1	99	
Atopobium parvulum		8.51	0.38		99	99	
Bacteroides vulgatus		7.16	0.28		1	1	
07135522	**Low discrimination**						**07135522**
					GRA		
Bifidobacterium longum subsp. longum		84.20	0.60	RAF 97	99		
Parabacteroides distasonis		8.68	0.47		1		
Lactobacillus fermentum		2.90	0.42		99		
07135530	**Very good identification**						**07135530**
					COC	GRA	
Bifidobacterium longum subsp. longum		99.88	0.64	RAF 97	1	99	
Atopobium parvulum		0.07	0.03		99	99	
Bacteroides vulgatus		0.05	-.06		1	1	
07135532	**Very good identification**						**07135532**
					GRA		
Bifidobacterium longum subsp. longum		95.18	0.64	RAF 97	99		
Parabacteroides distasonis		3.10	0.43		1		
Clostridium clostridioforme		1.60	0.32		50		
07135612	**Good identification**						**07135612**
Clostridium clostridioforme		100.0	0.38	SAL 88			
07135632	**Good identification**						**07135632**
Clostridium clostridioforme		97.39	0.38	SAL 88			
Parabacteroides distasonis		2.61	0.17	RAF 94			
07135702	**Low discrimination**						**07135702**
					GRA		
Clostridium clostridioforme		81.69	0.36	SAL 88	50		
Parabacteroides distasonis		5.54	0.21	RAF 94	1		
Bifidobacterium longum subsp. longum		4.48	0.15		99		
07135712	**Good identification**						**07135712**
Clostridium clostridioforme		96.27	0.47	SAL 88			
Bifidobacterium longum subsp. longum		1.97	0.19	RAF 94			
Bifidobacterium adolescentis		0.99	0.16				
07135720	**Good identification**						**07135720**
					GRA		
Bacteroides vulgatus		98.30	0.44	RAF 97	1		
Bifidobacterium longum subsp. longum		1.20	0.26	TRE 4	99		
Parabacteroides distasonis		0.37	0.21		1		
07135722	**Low discrimination**						**07135722**
					GRA		
Parabacteroides distasonis		81.75	0.55	RAF 99	1		
Clostridium clostridioforme		12.18	0.36		50		
Bifidobacterium longum subsp. longum		2.67	0.26		99		
07135732	**Low discrimination**						**07135732**
					GRA		
Clostridium clostridioforme		53.08	0.47	SAL 88	50		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		42.12	0.51	RAF 94	1			
Bifidobacterium longum subsp. longum		4.35	0.30		99			
07136000	**Good identification**							**07136000**
					COC	SPO	CAT	
Propionibacterium granulosum		98.73	0.56	XYL 1	1	1	90	
Gemella morbillorum		1.07	0.15		99	1	1	
Clostridium botulinum biovar B		0.21	-.03		1	99	1	
07136002	**Good identification**							**07136002**
					CAT			
Propionibacterium granulosum		99.28	0.46	XYL 1	90			
Propionibacterium propionicum		0.72	0.14		1			
07136010	**Good identification**							**07136010**
Propionibacterium granulosum		100.0	0.41	XYL 1				
07136020	**Good identification**							**07136020**
Propionibacterium granulosum		100.0	0.46	XYL 1				
07136022	**Good identification**							**07136022**
Propionibacterium granulosum		100.0	0.35	XYL 1				
07136100	**Low discrimination**							**07136100**
					COC	CAT		
Propionibacterium propionicum		83.04	0.45	NIT 86	1	1		
Atopobium parvulum		9.94	0.16	XYL 1	99	1		
Propionibacterium granulosum		7.03	0.22		1	90		
07136102	**Good identification**							**07136102**
					COC	CAT		
Propionibacterium propionicum		96.40	0.48	NIT 86	1	1		
Propionibacterium granulosum		1.36	0.12	XYL 1	1	90		
Atopobium parvulum		1.36	0.03		99	1		
07136140	**Good identification**							**07136140**
					SPO			
Propionibacterium propionicum		98.27	0.45	NIT 86	1			
Clostridium botulinum biovar B		1.73	-.03	XYL 1	99			
07136142	**Good identification**							**07136142**
Propionibacterium propionicum		100.0	0.48	NIT 86				
07137000	**Good identification**							**07137000**
					COC	CAT		
Propionibacterium granulosum		95.16	0.56	XYL 1	1	90		
Actinomyces israelii		2.58	0.35		1	1		
Atopobium parvulum		1.36	0.16		99	1		
07137002	**Low discrimination**							**07137002**
					COC	CAT		
Propionibacterium granulosum		83.69	0.46	XYL 1	1	90		
Actinomyces israelii		13.63	0.38		1	1		
Atopobium parvulum		0.84	0.03		99	1		
07137010	**Good identification**							**07137010**
					CAT			
Propionibacterium granulosum		97.11	0.41	XYL 1	90			
Actinomyces israelii		2.89	0.21	MLZ 12	1			
07137020	**Acceptable identification**							**07137020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	CAT	
Propionibacterium granulosum		92.41	0.46	XYL 1	1	90	
Clostridium perfringens		5.21	0.17		1	1	
Atopobium parvulum		0.93	0.03		99	1	
07137022	**Low discrimination**						**07137022**
					CAT		
Propionibacterium granulosum		84.54	0.35	XYL 1	90		
Clostridium perfringens		8.97	0.11		1		
Actinomyces israelii		3.52	0.17		1		
07137100	**Low discrimination**						**07137100**
					COC		
Atopobium parvulum		84.85	0.50	MAN 1	99		
Propionibacterium propionicum		7.16	0.45		1		
Actinomyces israelii		6.52	0.45		1		
07137102	**Low discrimination**						**07137102**
					COC		
Atopobium parvulum		37.78	0.38	MAN 1	99		
Propionibacterium propionicum		27.10	0.48		1		
Actinomyces israelii		24.66	0.48		1		
07137120	**Low discrimination**						**07137120**
					COC	SPO	
Atopobium parvulum		66.33	0.38	MAN 1	99	1	
Clostridium perfringens		27.46	0.32		1	84	
Clostridium tertium		2.03	0.11		1	99	
07137121	**Low discrimination**						**07137121**
					COC	SPO	
Actinomyces naeslundii		68.16	0.36	RAF 92	1	1	
Atopobium parvulum		18.25	0.03	XYL 1	99	1	
Clostridium chauvoei		13.60	0.01		1	99	
07137123	**Good identification**						**07137123**
Actinomyces naeslundii		100.0	0.36	RAF 92			
07137140	**Low discrimination**						**07137140**
					COC		
Propionibacterium propionicum		67.87	0.45	NIT 86	1		
Actinomyces israelii		24.01	0.38	XYL 1	1		
Atopobium parvulum		8.12	0.16		99		
07137142	**Low discrimination**						**07137142**
Propionibacterium propionicum		71.35	0.48	NIT 86			
Actinomyces israelii		25.25	0.41	XYL 1			
Bifidobacterium adolescentis		2.40	0.17				
07137402	**Low discrimination**						**07137402**
					CAT		
Actinomyces israelii		82.06	0.36	SAL 92	1		
Bifidobacterium adolescentis		8.12	0.13	CEL 93	1		
Propionibacterium granulosum		6.22	0.12		90		
07137500	**Low discrimination**						**07137500**
					COC		
Actinomyces israelii		77.54	0.44	SAL 92	1		
Atopobium parvulum		12.47	0.16	CEL 93	99		
Bifidobacterium longum subsp. longum		5.04	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07137502	**Low discrimination**							**07137502**
Bifidobacterium adolescentis		69.12	0.47	RAF 92				
Actinomyces israelii		28.24	0.47					
Bifidobacterium longum subsp. longum		1.22	0.15					
07137512	**Acceptable identification**							**07137512**
Bifidobacterium adolescentis		93.06	0.49	RAF 92				
Actinomyces israelii		4.65	0.33					
Bifidobacterium longum subsp. longum		2.30	0.19					
07137540	**Good identification**							**07137540**
Actinomyces israelii		100.0	0.37	SAL 92				
07137542	**Low discrimination**							**07137542**
Actinomyces israelii		55.16	0.40	SAL 92				
Bifidobacterium adolescentis		42.92	0.32	CEL 93				
Propionibacterium propionicum		1.92	0.14					
07140000	**Very good identification**							**07140000**
					COC	GRA	SPO	
Gemella morbillorum		95.15	0.64	CEL 1	99	99	1	
Prevotella disiens		3.76	0.32		1	1	1	
Clostridium botulinum biovar A		0.43	0.21		1	99	99	
07140020	**Good identification**							**07140020**
					COC	GRA	SPO	
Gemella morbillorum		97.97	0.48	CEL 1	99	99	1	
Clostridium botulinum biovar B		0.96	0.08	NAG 10	1	99	99	
Prevotella bivia		0.72	-.03		1	1	1	
07140040	**Acceptable identification**							**07140040**
					COC	SPO		
Clostridium botulinum biovar B		94.71	0.47	CEL 1	1	99		
Gemella morbillorum		4.38	0.30		99	1		
Clostridium botulinum biovar A		0.58	0.12		1	99		
07140050	**Good identification**							**07140050**
Clostridium botulinum biovar B		100.0	0.42	CEL 1				
07140060	**Good identification**							**07140060**
					COC	SPO		
Clostridium botulinum biovar B		98.97	0.42	CEL 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
07140070	**Good identification**							**07140070**
Clostridium botulinum biovar B		100.0	0.37	CEL 1				
07140400	**Good identification**							**07140400**
					COC	GRA		
Gemella morbillorum		99.64	0.48	CEL 1	99	99		
Prevotella disiens		0.36	-.03	ARA 10	1	1		
07141000	**Good identification**							**07141000**
					COC			
Gemella morbillorum		94.36	0.60	CEL 1	99			
Lactobacillus jensenii		3.61	0.36		1			
Lactobacillus catenaformis		1.38	0.29		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07141020	**Low discrimination**						**07141020**
					COC	GRA	
Gemella morbillorum		78.38	0.43	CEL 1	99	99	
Lactobacillus jensenii		6.74	0.26	NAG 10	1	99	
Prevotella melaninogenica		5.90	0.17		1	1	
07141101	**Good identification**						**07141101**
					SPO		
Clostridium chauvoei		99.20	0.54	CEL 1	99		
Actinomyces naeslundii		0.80	0.41		1		
07141103	**Good identification**						**07141103**
					SPO		
Clostridium chauvoei		95.08	0.41	CEL 1	99		
Actinomyces naeslundii		4.32	0.41		1		
Actinomyces viscosus		0.60	0.02		1		
07141120	**Low discrimination**						**07141120**
					GRA		
Prevotella melaninogenica		89.28	0.51	RAF 90	1		
Clostridium perfringens		5.84	0.38	FRU 90	99		
Prevotella bivia		2.87	0.19	CEL 10	1		
07141121	**Acceptable identification**						**07141121**
					GRA	SPO	
Clostridium chauvoei		92.88	0.45	CEL 1	99	99	
Prevotella melaninogenica		4.59	0.17		1	1	
Actinomyces naeslundii		2.24	0.41		99	1	
07141122	**Low discrimination**						**07141122**
					GRA		
Prevotella melaninogenica		65.28	0.39	RAF 90	1		
Prevotella oralis		13.60	0.31	FRU 90	1		
Clostridium perfringens		10.54	0.32	CEL 10	99		
07141400	**Good identification**						**07141400**
					COC		
Gemella morbillorum		99.13	0.43	CEL 1	99		
Lactobacillus fermentum		0.52	0.16	ARA 10	1		
Lactobacillus jensenii		0.34	0.02		1		
07143001	**Good identification**						**07143001**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07143003	**Good identification**						**07143003**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07143021	**Good identification**						**07143021**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07143023	**Good identification**						**07143023**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07143100	**Acceptable identification**						**07143100**
Actinomyces naeslundii		93.81	0.43	RAF 92			
Propionibacterium propionicum		6.19	0.12	FRU 95			
07143101	**Low discrimination**						**07143101**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		60.65	0.46	RAF 92	1			
Clostridium chauvoei		39.35	0.19	FRU 95	99			
07143102	**Low discrimination**							**07143102**
Actinomyces naeslundii		83.62	0.43	RAF 92				
Propionibacterium propionicum		8.27	0.15	FRU 95				
Lactobacillus acidophilus		8.11	0.01					
07143103	**Low discrimination**							**07143103**
					SPO			
Actinomyces naeslundii		89.73	0.46	RAF 92	1			
Clostridium chauvoei		10.27	0.07	FRU 95	99			
07143121	**Low discrimination**							**07143121**
					SPO			
Actinomyces naeslundii		82.22	0.46	RAF 92	1			
Clostridium chauvoei		17.78	0.11	FRU 95	99			
07143122	**Low discrimination**							**07143122**
					GRA	SPO		
Actinomyces naeslundii		62.86	0.43	RAF 92	99	1		
Prevotella melaninogenica		19.00	0.05	FRU 95	1	1		
Clostridium barati		6.85	-.01		99	99		
07143123	**Good identification**							**07143123**
Actinomyces naeslundii		100.0	0.46	RAF 92				
07144000	**Good identification**							**07144000**
					COC	GRA		
Gemella morbillorum		99.64	0.48	CEL 1	99	99		
Prevotella disiens		0.36	-.03	XYL 10	1	1		
07145000	**Good identification**							**07145000**
					COC			
Gemella morbillorum		99.35	0.43	CEL 1	99			
Lactobacillus jensenii		0.35	0.02	XYL 10	1			
Lactobacillus fermentum		0.31	0.12		1			
07150000	**Low discrimination**							**07150000**
					COC	GRA		
Gemella morbillorum		79.85	0.66	CEL 1	99	99		
Prevotella intermedia		6.49	0.46		1	1		
Fusobacterium mortiferum		4.14	0.54		1	1		
07150020	**Low discrimination**							**07150020**
					COC	CAT		
Gemella morbillorum		79.79	0.50	CEL 1	99	1		
Propionibacterium granulosum		8.25	0.40	NAG 10	1	90		
Lactobacillus jensenii		3.02	0.26		1	1		
07150040	**Very good identification**							**07150040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		97.55	0.58	CEL 1	1	99	99	
Gemella morbillorum		1.38	0.32		99	99	1	
Megasphaera elsdenii		0.76	0.29		99	1	1	
07150050	**Good identification**							**07150050**
Clostridium botulinum biovar B		100.0	0.52	CEL 1				
07150060	**Good identification**							**07150060**
					COC	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar B		99.51	0.52	CEL 1	1	99	
Gemella morbillorum		0.32	0.15		99	1	
Clostridium perfringens		0.18	0.11		1	84	
07150070	**Good identification**						**07150070**
Clostridium botulinum biovar B		100.0	0.47	CEL 1			
07150100	**Low discrimination**						**07150100**
					COC	GRA	
Fusobacterium mortiferum		79.26	0.60	MLT 11	1	1	
Gemella morbillorum		6.62	0.32		99	99	
Lactobacillus catenaformis		6.32	0.32		1	99	
07150102	**Low discrimination**						**07150102**
					GRA		
Actinomyces viscosus		39.49	0.36	NIT 99	99		
Fusobacterium mortiferum		22.94	0.44	RAF 94	1		
Lactobacillus catenaformis		16.47	0.32		99		
07150123	**Low discrimination**						**07150123**
					SPO		
Actinomyces naeslundii		75.40	0.39	RAF 92	1		
Clostridium chauvoei		16.46	0.05	GAL 96	99		
Clostridium barati		8.13	-.04		99		
07150400	**Good identification**						**07150400**
					COC	GRA	
Gemella morbillorum		97.58	0.50	CEL 1	99	99	
Prevotella intermedia		0.72	0.12	ARA 10	1	1	
Fusobacterium mortiferum		0.46	0.20		1	1	
07151000	**Low discrimination**						**07151000**
					COC		
Lactobacillus jensenii		53.65	0.70	SAL 95	1		
Lactobacillus catenaformis		20.51	0.63		1		
Gemella morbillorum		17.33	0.61		99		
07151001	**Low discrimination**						**07151001**
Lactobacillus jensenii		34.86	0.36	SAL 95			
Actinomyces naeslundii		24.81	0.55	URE 1			
Lactobacillus catenaformis		13.33	0.29				
07151002	**Low discrimination**						**07151002**
Lactobacillus jensenii		58.63	0.67	SAL 95			
Lactobacillus catenaformis		33.62	0.63				
Lactobacillus acidophilus		1.94	0.39				
07151003	**Low discrimination**						**07151003**
Actinomyces naeslundii		35.05	0.55	RAF 92			
Lactobacillus jensenii		32.83	0.33				
Lactobacillus catenaformis		18.83	0.29				
07151010	**Very good identification**						**07151010**
					CAT		
Lactobacillus jensenii		96.45	0.62	SAL 95	1		
Lactobacillus catenaformis		1.12	0.29		1		
Propionibacterium granulosum		1.09	0.35		90		
07151012	**Very good identification**						**07151012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Lactobacillus jensenii		97.53	0.59	SAL 95	1	
Lactobacillus catenaformis		1.70	0.29		1	
Propionibacterium granulosum		0.41	0.25		90	
07151020	**Low discrimination**					**07151020**
Lactobacillus jensenii		39.32	0.60	SAL 95		
Clostridium perfringens		30.39	0.57			
Lactobacillus catenaformis		15.03	0.53			
07151021	**Low discrimination**					**07151021**
Actinomyces naeslundii		49.77	0.55	RAF 92		
Lactobacillus jensenii		17.48	0.26			
Clostridium perfringens		13.51	0.23			
07151022	**Low discrimination**					**07151022**
Lactobacillus jensenii		33.08	0.57	SAL 95		
Lactobacillus catenaformis		18.97	0.53			
Clostridium perfringens		18.05	0.51			
07151023	**Low discrimination**					**07151023**
Actinomyces naeslundii		58.44	0.55	RAF 92		
Lactobacillus jensenii		13.69	0.23			
Lactobacillus catenaformis		7.85	0.18			
07151030	**Acceptable identification**					**07151030**
Lactobacillus jensenii		94.48	0.52	SAL 95		
Clostridium perfringens		2.21	0.23			
Lactobacillus catenaformis		1.09	0.18			
07151032	**Acceptable identification**					**07151032**
Lactobacillus jensenii		93.71	0.49	SAL 95		
Lactobacillus catenaformis		1.63	0.18			
Clostridium perfringens		1.55	0.17			
07151040	**Low discrimination**					**07151040**
					SPO	
Lactobacillus jensenii		39.62	0.36	SAL 95	1	
Clostridium botulinum biovar B		17.00	0.24	SOR 1	99	
Lactobacillus catenaformis		15.14	0.29		1	
07151041	**Acceptable identification**					**07151041**
Actinomyces naeslundii		92.56	0.42	RAF 92		
Lactobacillus jensenii		7.44	0.02			
07151043	**Good identification**					**07151043**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07151060	**Low discrimination**					**07151060**
					SPO	
Clostridium perfringens		80.87	0.45	CEL 4	84	
Lactobacillus jensenii		5.55	0.26	LAC 88	1	
Clostridium botulinum biovar B		4.69	0.18		99	
07151061	**Low discrimination**					**07151061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		77.34	0.42	RAF 92			
Clostridium perfringens		22.66	0.11				
07151062	**Low discrimination**						**07151062**
					SPO		
Clostridium perfringens		59.49	0.39	CEL 4	84		
Clostridium barati		17.22	0.25	LAC 88	99		
Lactobacillus acidophilus		9.38	0.23		1		
07151063	**Low discrimination**						**07151063**
Actinomyces naeslundii		87.88	0.42	RAF 92			
Clostridium perfringens		12.12	0.05				
07151100	**Low discrimination**						**07151100**
					COC	GRA	
Lactobacillus catenaformis		58.31	0.66	SAL 99	1	99	
Atopobium parvulum		11.27	0.52		99	99	
Fusobacterium mortiferum		11.07	0.63		1	1	
07151101	**Good identification**						**07151101**
					SPO		
Clostridium chauvoei		91.24	0.60	CEL 1	99		
Actinomyces naeslundii		5.96	0.63		1		
Lactobacillus catenaformis		1.69	0.32		1		
07151102	**Low discrimination**						**07151102**
					SPO		
Lactobacillus catenaformis		54.59	0.66	SAL 99	1		
Lactobacillus acidophilus		24.20	0.57		1		
Clostridium barati		5.49	0.44		99		
07151103	**Low discrimination**						**07151103**
					SPO		
Clostridium chauvoei		62.29	0.47	CEL 1	99		
Actinomyces naeslundii		23.05	0.63		1		
Lactobacillus catenaformis		6.53	0.32		1		
07151113	**Low discrimination**						**07151113**
					SPO		
Actinomyces naeslundii		65.85	0.41	RAF 92	1		
Clostridium chauvoei		34.15	0.13	MLZ 5	99		
07151120	**Low discrimination**						**07151120**
					GRA		
Prevotella melaninogenica		52.25	0.67	RAF 90	1		
Clostridium perfringens		37.58	0.72	CEL 10	99		
Lactobacillus catenaformis		3.80	0.56		99		
07151121	**Low discrimination**						**07151121**
					GRA	SPO	
Clostridium chauvoei		64.23	0.52	CEL 1	99	99	
Actinomyces naeslundii		12.58	0.63		99	1	
Prevotella melaninogenica		12.24	0.33		1	1	
07151122	**Low discrimination**						**07151122**
					GRA		
Prevotella oralis		33.96	0.65	RAF 92	1		
Clostridium perfringens		26.34	0.66		99		
Prevotella melaninogenica		14.83	0.55		1		
07151123	**Low discrimination**						**07151123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Actinomyces naeslundii		31.90	0.63	RAF 92	99	1	
Clostridium chauvoei		28.73	0.39		99	99	
Prevotella oralis		13.54	0.31		1	1	
07151131	**Low discrimination**						**07151131**
					GRA	SPO	
Actinomyces naeslundii		43.47	0.41	RAF 92	99	1	
Clostridium chauvoei		42.58	0.18	MLZ 5	99	99	
Prevotella melaninogenica		8.12	-.01		1	1	
07151133	**Low discrimination**						**07151133**
					SPO		
Actinomyces naeslundii		85.26	0.41	RAF 92	1		
Clostridium chauvoei		14.74	0.05	MLZ 5	99		
07151141	**Low discrimination**						**07151141**
					SPO		
Actinomyces naeslundii		53.29	0.50	RAF 92	1		
Clostridium chauvoei		46.71	0.26		99		
07151142	**Low discrimination**						**07151142**
					SPO		
Lactobacillus acidophilus		62.36	0.41	SAL 99	1		
Lactobacillus catenaformis		11.50	0.32	SOR 11	1		
Clostridium barati		9.95	0.25		99		
07151143	**Low discrimination**						**07151143**
					SPO		
Actinomyces naeslundii		80.48	0.50	RAF 92	1		
Clostridium chauvoei		12.45	0.13		99		
Lactobacillus acidophilus		7.07	0.07		1		
07151160	**Low discrimination**						**07151160**
					GRA	SPO	
Clostridium perfringens		89.31	0.59	CEL 4	99	84	
Prevotella melaninogenica		6.59	0.33		1	1	
Clostridium barati		1.66	0.25		99	99	
07151161	**Low discrimination**						**07151161**
					SPO		
Actinomyces naeslundii		47.56	0.50	RAF 92	1		
Clostridium perfringens		35.90	0.25		84		
Clostridium chauvoei		13.89	0.18		99		
07151162	**Low discrimination**						**07151162**
					SPO		
Clostridium perfringens		65.92	0.54	CEL 4	84		
Lactobacillus acidophilus		16.30	0.41		1		
Clostridium barati		7.81	0.34		99		
07151163	**Low discrimination**						**07151163**
Actinomyces naeslundii		65.07	0.50	RAF 92			
Clostridium perfringens		23.12	0.20				
Lactobacillus acidophilus		5.72	0.07				
07151200	**Low discrimination**						**07151200**
					COC		
Lactobacillus jensenii		50.85	0.36	SAL 95	1		
Lactobacillus catenaformis		19.44	0.29	RHA 1	1		
Gemella morbillorum		16.43	0.27		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07151222	**Low discrimination**						**07151222**
					GRA	SPO	
Prevotella oralis		69.66	0.38	RAF 92	1	1	
Clostridium barati		14.85	0.25	LAC 94	99	99	
Lactobacillus jensenii		4.99	0.23		99	1	
07151302	**Low discrimination**						**07151302**
					GRA		
Prevotella oralis		38.60	0.37	RAF 92	1		
Lactobacillus acidophilus		38.07	0.38	NAG 95	99		
Lactobacillus catenaformis		11.53	0.32		99		
07151322	**Low discrimination**						**07151322**
					GRA	SPO	
Prevotella oralis		88.17	0.58	RAF 92	1	1	
Lactobacillus acidophilus		4.58	0.38		99	1	
Clostridium barati		3.60	0.34		99	99	
07151400	**Low discrimination**						**07151400**
					COC		
Gemella morbillorum		53.86	0.45	CEL 1	99		
Lactobacillus fermentum		23.07	0.50	ARA 10	1		
Lactobacillus jensenii		15.16	0.36		1		
07151402	**Low discrimination**						**07151402**
Lactobacillus fermentum		56.99	0.50	CEL 1			
Lactobacillus jensenii		24.95	0.33				
Lactobacillus catenaformis		14.31	0.29				
07151420	**Low discrimination**						**07151420**
					COC		
Lactobacillus fermentum		34.28	0.42	CEL 1	1		
Gemella morbillorum		26.68	0.29		99		
Lactobacillus jensenii		16.89	0.26		1		
07151422	**Low discrimination**						**07151422**
Lactobacillus fermentum		50.99	0.42	CEL 1			
Lactobacillus jensenii		16.75	0.23				
Lactobacillus catenaformis		9.60	0.18				
07151500	**Low discrimination**						**07151500**
					COC		
Lactobacillus fermentum		68.64	0.54	CEL 1	1		
Lactobacillus catenaformis		15.85	0.32		1		
Atopobium parvulum		3.06	0.18		99		
07151502	**Low discrimination**						**07151502**
Lactobacillus fermentum		51.96	0.54	CEL 1			
Bifidobacterium adolescentis		24.54	0.43				
Lactobacillus catenaformis		12.00	0.32				
07151512	**Acceptable identification**						**07151512**
Bifidobacterium adolescentis		92.02	0.45	RAF 92			
Bifidobacterium longum subsp. longum		5.20	0.22	XYL 87			
Lactobacillus fermentum		1.61	0.20				
07152000	**Low discrimination**						**07152000**
					COC	GRA CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		61.19	0.48	CEL 1	1	99	90	
Gemella morbillorum		17.83	0.32		99	99	1	
Megasphaera elsdenii		14.72	0.32		99	1	1	
07152001	**Low discrimination**							**07152001**
Actinomyces naeslundii		73.46	0.36	RAF 92				
Propionibacterium granulosum		26.54	0.14	GAL 96				
07152002	**Low discrimination**							**07152002**
					COC	SPO	CAT	
Propionibacterium granulosum		77.45	0.38	CEL 1	1	1	90	
Megasphaera elsdenii		8.28	0.15		99	1	1	
Clostridium innocuum		6.45	0.10		1	99	1	
07152003	**Good identification**							**07152003**
Actinomyces naeslundii		100.0	0.36	RAF 92				
07152020	**Low discrimination**							**07152020**
					COC	CAT		
Propionibacterium granulosum		83.05	0.38	CEL 1	1	90		
Gemella morbillorum		10.76	0.15		99	1		
Actinomyces naeslundii		6.19	0.33		1	25		
07152021	**Good identification**							**07152021**
Actinomyces naeslundii		100.0	0.36	RAF 92				
07152023	**Good identification**							**07152023**
Actinomyces naeslundii		100.0	0.36	RAF 92				
07152100	**Low discrimination**							**07152100**
					GRA			
Propionibacterium propionicum		81.64	0.46	NIT 86	99			
Actinomyces naeslundii		9.90	0.41	CEL 1	99			
Fusobacterium mortiferum		6.58	0.26		1			
07152101	**Low discrimination**							**07152101**
Actinomyces naeslundii		62.07	0.44	RAF 92				
Propionibacterium propionicum		37.93	0.30	GAL 96				
07152102	**Low discrimination**							**07152102**
					SPO			
Propionibacterium propionicum		89.26	0.49	NIT 86	1			
Actinomyces naeslundii		7.21	0.41	CEL 1	1			
Clostridium barati		1.73	0.04		99			
07152103	**Low discrimination**							**07152103**
Actinomyces naeslundii		52.18	0.44	RAF 92				
Propionibacterium propionicum		47.82	0.33	GAL 96				
07152120	**Low discrimination**							**07152120**
					GRA	SPO		
Actinomyces naeslundii		61.14	0.41	RAF 92	99	1		
Clostridium barati		14.62	0.04	GAL 96	99	99		
Prevotella melaninogenica		11.14	-.01		1	1		
07152121	**Good identification**							**07152121**
Actinomyces naeslundii		100.0	0.44	RAF 92				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07152122	**Low discrimination**					**07152122**
					SPO	
Actinomyces naeslundii		48.32	0.41	RAF 92	1	
Clostridium barati		34.67	0.12	GAL 96	99	
Propionibacterium propionicum		6.04	0.15		1	
07152123	**Good identification**					**07152123**
Actinomyces naeslundii		100.0	0.44	RAF 92		
07152140	**Good identification**					**07152140**
Propionibacterium propionicum		97.91	0.46	NIT 86		
Actinomyces naeslundii		2.09	0.28	CEL 1		
07152142	**Good identification**					**07152142**
Propionibacterium propionicum		98.59	0.49	NIT 86		
Actinomyces naeslundii		1.41	0.28	CEL 1		
07153000	**Low discrimination**					**07153000**
					CAT	
Propionibacterium granulosum		35.96	0.48	CEL 1	90	
Lactobacillus jensenii		17.47	0.36		1	
Actinomyces naeslundii		16.09	0.57		25	
07153001	**Very good identification**					**07153001**
					CAT	
Actinomyces naeslundii		97.81	0.60	RAF 92	25	
Propionibacterium granulosum		1.47	0.14		90	
Lactobacillus jensenii		0.72	0.02		1	
07153002	**Low discrimination**					**07153002**
					SPO	
Actinomyces naeslundii		24.99	0.57	RAF 92	1	
Clostridium innocuum		22.08	0.32		99	
Lactobacillus jensenii		18.09	0.33		1	
07153003	**Very good identification**					**07153003**
Actinomyces naeslundii		100.0	0.60	RAF 92		
07153011	**Good identification**					**07153011**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07153013	**Good identification**					**07153013**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07153020	**Low discrimination**					**07153020**
					CAT	
Actinomyces naeslundii		41.95	0.57	RAF 92	25	
Propionibacterium granulosum		23.44	0.38		90	
Lactobacillus jensenii		11.39	0.26		1	
07153021	**Very good identification**					**07153021**
Actinomyces naeslundii		100.0	0.60	RAF 92		
07153022	**Low discrimination**					**07153022**
					SPO	
Actinomyces naeslundii		44.27	0.57	RAF 92	1	
Clostridium barati		23.85	0.25		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		8.01	0.23		1	
07153023	**Very good identification**					**07153023**
Actinomyces naeslundii		100.0	0.60	RAF 92		
07153031	**Good identification**					**07153031**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07153033	**Good identification**					**07153033**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07153040	**Low discrimination**					**07153040**
					CAT	
Actinomyces naeslundii		73.03	0.44	RAF 92	25	
Propionibacterium granulosum		9.34	0.14		90	
Actinomyces israelii		9.00	0.19		1	
07153041	**Good identification**					**07153041**
Actinomyces naeslundii		100.0	0.47	RAF 92		
07153042	**Low discrimination**					**07153042**
					SPO	
Actinomyces naeslundii		69.21	0.44	RAF 92	1	
Actinomyces israelii		12.79	0.22		1	
Clostridium barati		6.13	-.01		99	
07153043	**Good identification**					**07153043**
Actinomyces naeslundii		100.0	0.47	RAF 92		
07153060	**Low discrimination**					**07153060**
					SPO	
Actinomyces naeslundii		76.05	0.44	RAF 92	1	
Clostridium perfringens		17.22	0.11		84	
Clostridium barati		6.73	-.01		99	
07153061	**Good identification**					**07153061**
Actinomyces naeslundii		100.0	0.47	RAF 92		
07153062	**Low discrimination**					**07153062**
					SPO	
Actinomyces naeslundii		68.51	0.44	RAF 92	1	
Clostridium barati		18.19	0.07		99	
Clostridium perfringens		7.30	0.05		84	
07153063	**Good identification**					**07153063**
Actinomyces naeslundii		100.0	0.47	RAF 92		
07153100	**Low discrimination**					**07153100**
Actinomyces naeslundii		52.56	0.65	RAF 92		
Propionibacterium propionicum		18.06	0.46			
Lactobacillus catenaformis		11.50	0.32			
07153101	**Good identification**					**07153101**
					SPO	
Actinomyces naeslundii		90.49	0.68	RAF 92	1	
Clostridium chauvoei		7.21	0.26		99	
Propionibacterium propionicum		2.30	0.30		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07153102	**Low discrimination**						**07153102**
Actinomyces naeslundii		35.82	0.65	RAF 92			
Propionibacterium propionicum		18.47	0.49				
Lactobacillus acidophilus		18.10	0.35				
07153103	**Good identification**						**07153103**
					SPO		
Actinomyces naeslundii		94.73	0.68	RAF 92	1		
Propionibacterium propionicum		3.62	0.33		1		
Clostridium chauvoei		1.33	0.13		99		
07153111	**Good identification**						**07153111**
Actinomyces naeslundii		100.0	0.46	RAF 92			
07153113	**Good identification**						**07153113**
Actinomyces naeslundii		100.0	0.46	RAF 92			
07153120	**Low discrimination**						**07153120**
					GRA		
Actinomyces naeslundii		35.38	0.65	RAF 92	99		
Prevotella melaninogenica		26.60	0.33		1		
Clostridium perfringens		19.13	0.38		99		
07153121	**Very good identification**						**07153121**
					GRA	SPO	
Actinomyces naeslundii		96.59	0.68	RAF 92	99	1	
Clostridium chauvoei		2.57	0.18		99	99	
Prevotella melaninogenica		0.49	-.01		1	1	
07153122	**Low discrimination**						**07153122**
					SPO		
Actinomyces naeslundii		27.60	0.65	RAF 92	1		
Clostridium barati		15.68	0.34		99		
Lactobacillus acidophilus		13.94	0.35		1		
07153123	**Very good identification**						**07153123**
					SPO		
Actinomyces naeslundii		98.54	0.68	RAF 92	1		
Clostridium chauvoei		0.46	0.05		99		
Clostridium barati		0.38	-.01		99		
07153131	**Good identification**						**07153131**
Actinomyces naeslundii		100.0	0.46	RAF 92			
07153132	**Acceptable identification**						**07153132**
					SPO		
Clostridium tertium		91.84	0.39	SAL 99	99		
Actinomyces naeslundii		6.31	0.43	TRE 90	1		
Clostridium barati		0.69	-.01		99		
07153133	**Acceptable identification**						**07153133**
					SPO		
Actinomyces naeslundii		91.08	0.46	RAF 92	1		
Clostridium tertium		8.92	0.05	MLZ 5	99		
07153140	**Low discrimination**						**07153140**
Propionibacterium propionicum		61.79	0.46	NIT 86			
Actinomyces naeslundii		31.73	0.52	CEL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		5.49	0.29				
07153141	**Low discrimination**						**07153141**
Actinomyces naeslundii		87.39	0.55	RAF 92			
Propionibacterium propionicum		12.61	0.30				
07153142	**Low discrimination**						**07153142**
Propionibacterium propionicum		62.60	0.49	NIT 86			
Actinomyces naeslundii		21.43	0.52	CEL 1			
Lactobacillus acidophilus		7.58	0.19				
07153143	**Low discrimination**						**07153143**
Actinomyces naeslundii		82.21	0.55	RAF 92			
Propionibacterium propionicum		17.79	0.33				
07153160	**Low discrimination**						**07153160**
					SPO		
Actinomyces naeslundii		57.49	0.52	RAF 92	1		
Clostridium perfringens		33.54	0.25		84		
Clostridium barati		5.36	0.07		99		
07153161	**Good identification**						**07153161**
Actinomyces naeslundii		100.0	0.55	RAF 92			
07153162	**Low discrimination**						**07153162**
					SPO		
Actinomyces naeslundii		50.30	0.52	RAF 92	1		
Lactobacillus acidophilus		17.80	0.19		1		
Clostridium barati		14.08	0.15		99		
07153163	**Good identification**						**07153163**
Actinomyces naeslundii		100.0	0.55	RAF 92			
07153502	**Low discrimination**						**07153502**
Bifidobacterium adolescentis		74.34	0.42	RAF 92			
Actinomyces israelii		19.58	0.38	XYL 87			
Actinomyces naeslundii		2.05	0.30				
07153512	**Good identification**						**07153512**
Bifidobacterium adolescentis		96.88	0.43	RAF 92			
Actinomyces israelii		3.12	0.24	XYL 87			
07154000	**Acceptable identification**						**07154000**
					COC	GRA	
Gemella morbillorum		93.83	0.50	CEL 1	99	99	
Fusobacterium mortiferum		3.30	0.35	XYL 10	1	1	
Actinomyces meyeri		1.25	0.18		1	99	
07154100	**Low discrimination**						**07154100**
					COC	GRA	
Fusobacterium mortiferum		45.89	0.41	MLT 11	1	1	
Actinomyces meyeri		29.75	0.29	XYL 7	1	99	
Atopobium parvulum		17.46	0.23		99	99	
07155000	**Low discrimination**						**07155000**
					COC		
Gemella morbillorum		53.07	0.45	CEL 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		14.93	0.36	XYL 10	1		
Lactobacillus fermentum		13.35	0.46		1		
07155002	**Low discrimination**						**07155002**
Lactobacillus fermentum		38.89	0.46	CEL 1			
Lactobacillus jensenii		29.00	0.33				
Lactobacillus catenaformis		16.63	0.29				
07155100	**Low discrimination**						**07155100**
					COC	GRA	
Atopobium parvulum		83.51	0.57	TRE 99	99	99	
Lactobacillus fermentum		5.98	0.50		1	99	
Fusobacterium mortiferum		3.33	0.44		1	1	
07155102	**Low discrimination**						**07155102**
					COC		
Atopobium parvulum		42.31	0.44	TRE 99	99		
Lactobacillus fermentum		17.17	0.50		1		
Bifidobacterium adolescentis		13.80	0.43		1		
07155112	**Low discrimination**						**07155112**
					COC		
Bifidobacterium adolescentis		89.63	0.45	RAF 92	1		
Actinomyces israelii		5.86	0.31	ARA 87	1		
Atopobium parvulum		2.27	0.10		99		
07155120	**Low discrimination**						**07155120**
					COC	GRA	
Atopobium parvulum		41.50	0.44	TRE 99	99	99	
Prevotella melaninogenica		22.74	0.33		1	1	
Clostridium perfringens		16.35	0.38		1	99	
07155122	**Low discrimination**						**07155122**
					CAT	GRA	
Prevotella oralis		67.84	0.49	RAF 92	1	1	
Bacteroides fragilis		9.28	0.25	XYL 11	99	1	
Clostridium perfringens		4.30	0.32		1	99	
07155142	**Low discrimination**						**07155142**
					COC		
Actinomyces israelii		51.74	0.38	SAL 92	1		
Bifidobacterium adolescentis		30.76	0.28		1		
Atopobium parvulum		7.71	0.10		99		
07155322	**Low discrimination**						**07155322**
Prevotella oralis		83.03	0.43	RAF 92			
Parabacteroides distasonis		15.52	0.43	XYL 11			
Clostridium clostridioforme		0.35	0.09				
07155332	**Low discrimination**						**07155332**
Parabacteroides distasonis		77.47	0.39	RAF 99			
Clostridium clostridioforme		14.75	0.21				
Prevotella oralis		7.78	0.09				
07155400	**Low discrimination**						**07155400**
					COC		
Lactobacillus fermentum		63.08	0.46	CEL 1	1		
Gemella morbillorum		27.86	0.29		99		
Actinomyces israelii		6.03	0.30		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07155402	**Low discrimination**						**07155402**
Lactobacillus fermentum		80.28	0.46	CEL 1			
Actinomyces israelii		11.52	0.33				
Bifidobacterium adolescentis		7.12	0.23				
07155420	**Low discrimination**						**07155420**
					COC	GRA	
Lactobacillus fermentum		83.15	0.38	CEL 1	1	99	
Gemella morbillorum		12.24	0.12		99	99	
Bacteroides vulgatus		4.62	-.05		1	1	
07155422	**Good identification**						**07155422**
Lactobacillus fermentum		96.10	0.38	CEL 1			
Clostridium clostridioforme		3.90	0.06				
07155500	**Low discrimination**						**07155500**
					COC		
Lactobacillus fermentum		59.28	0.50	CEL 1	1		
Actinomyces israelii		13.90	0.40		1		
Atopobium parvulum		8.36	0.23		99		
07155502	**Low discrimination**						**07155502**
Bifidobacterium adolescentis		77.92	0.57	RAF 92			
Lactobacillus fermentum		14.48	0.50				
Actinomyces israelii		5.09	0.43				
07155512	**Good identification**						**07155512**
Bifidobacterium adolescentis		94.87	0.59	RAF 92			
Bifidobacterium longum subsp. longum		2.40	0.30				
Clostridium clostridioforme		1.83	0.26				
07155522	**Low discrimination**						**07155522**
					GRA		
Bifidobacterium longum subsp. longum		35.73	0.36	RAF 97	99		
Lactobacillus fermentum		30.44	0.42	CEL 1	99		
Parabacteroides distasonis		10.78	0.31		1		
07155530	**Good identification**						**07155530**
					GRA		
Bifidobacterium longum subsp. longum		98.33	0.40	RAF 97	99		
Bacteroides vulgatus		1.67	-.05	CEL 1	1		
07155532	**Low discrimination**						**07155532**
Bifidobacterium longum subsp. longum		72.20	0.40	RAF 97			
Clostridium clostridioforme		13.71	0.26	CEL 1			
Bifidobacterium adolescentis		7.20	0.25				
07155542	**Low discrimination**						**07155542**
Bifidobacterium adolescentis		81.91	0.42	RAF 92			
Actinomyces israelii		16.84	0.36	SOR 11			
Lactobacillus fermentum		1.24	0.16				
07155552	**Good identification**						**07155552**
Bifidobacterium adolescentis		97.55	0.43	RAF 92			
Actinomyces israelii		2.45	0.22	SOR 11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07155712	**Acceptable identification**						**07155712**
Clostridium clostridioforme		93.32	0.41	SAL 88			
Bifidobacterium adolescentis		6.68	0.25	RAF 94			
07155720	**Good identification**						**07155720**
Bacteroides vulgatus		100.0	0.46	RAF 97			
07155722	**Low discrimination**						**07155722**
Parabacteroides distasonis		50.11	0.39	RAF 99			
Clostridium clostridioforme		28.83	0.30				
Bacteroides vulgatus		14.45	0.12				
07155732	**Low discrimination**						**07155732**
					GRA		
Clostridium clostridioforme		82.46	0.41	SAL 88	50		
Parabacteroides distasonis		16.94	0.35	RAF 94	1		
Bifidobacterium longum subsp. longum		0.60	0.06	TRE 88	99		
07157000	**Low discrimination**						**07157000**
					COC	SPO	
Actinomyces israelii		82.23	0.44	SAL 92	1	1	
Clostridium innocuum		5.95	0.10		1	99	
Gemella morbillorum		4.99	0.11		99	1	
07157002	**Acceptable identification**						**07157002**
					SPO		
Actinomyces israelii		92.61	0.47	SAL 92	1		
Clostridium innocuum		4.47	0.10		99		
Bifidobacterium adolescentis		1.01	0.07		1		
07157040	**Good identification**						**07157040**
Actinomyces israelii		100.0	0.37	SAL 92			
07157042	**Good identification**						**07157042**
Actinomyces israelii		100.0	0.40	SAL 92			
07157100	**Low discrimination**						**07157100**
					COC	SPO	
Actinomyces israelii		84.36	0.55	SAL 92	1	1	
Atopobium parvulum		7.33	0.23		99	1	
Clostridium tertium		6.17	0.21		1	99	
07157102	**Low discrimination**						**07157102**
					SPO		
Actinomyces israelii		69.17	0.58	SAL 92	1		
Bifidobacterium adolescentis		18.68	0.42		1		
Clostridium tertium		10.12	0.29		99		
07157110	**Low discrimination**						**07157110**
					SPO		
Actinomyces israelii		52.54	0.40	SAL 92	1		
Clostridium tertium		45.71	0.25	MLZ 13	99		
Bifidobacterium adolescentis		1.76	0.09		1		
07157112	**Low discrimination**						**07157112**
					SPO		
Bifidobacterium adolescentis		44.64	0.43	RAF 92	1		
Clostridium tertium		35.16	0.33	ARA 87	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		20.20	0.43		1		
07157122	**Low discrimination**						**07157122**
					GRA	SPO	
Clostridium tertium		79.09	0.37	SAL 99	99	99	
Actinomyces israelii		11.51	0.37	TRE 90	99	1	
Prevotella oralis		5.39	0.15		1	1	
07157132	**Good identification**						**07157132**
					SPO		
Clostridium tertium		98.38	0.41	SAL 99	99		
Actinomyces israelii		1.20	0.23	TRE 90	1		
Bifidobacterium adolescentis		0.42	0.09		1		
07157140	**Good identification**						**07157140**
Actinomyces israelii		99.02	0.48	SAL 92			
Propionibacterium propionicum		0.98	0.12				
07157142	**Acceptable identification**						**07157142**
Actinomyces israelii		91.26	0.51	SAL 92			
Bifidobacterium adolescentis		7.84	0.26				
Propionibacterium propionicum		0.90	0.15				
07157152	**Low discrimination**						**07157152**
					SPO		
Actinomyces israelii		57.23	0.36	SAL 92	1		
Bifidobacterium adolescentis		40.19	0.28	MLZ 13	1		
Clostridium tertium		2.59	-.01		99		
07157300	**Good identification**						**07157300**
Actinomyces israelii		100.0	0.35	SAL 92			
07157302	**Good identification**						**07157302**
Actinomyces israelii		96.50	0.38	SAL 92			
Bifidobacterium adolescentis		3.50	0.07	RHA 7			
07157400	**Good identification**						**07157400**
Actinomyces israelii		98.17	0.43	SAL 92			
Lactobacillus fermentum		1.83	0.12				
07157402	**Acceptable identification**						**07157402**
Actinomyces israelii		90.77	0.46	SAL 92			
Bifidobacterium adolescentis		8.10	0.22				
Lactobacillus fermentum		1.13	0.12				
07157440	**Good identification**						**07157440**
Actinomyces israelii		100.0	0.36	SAL 92			
07157442	**Good identification**						**07157442**
Actinomyces israelii		100.0	0.39	SAL 92			
07157500	**Good identification**						**07157500**
Actinomyces israelii		95.61	0.53	SAL 92			
Bifidobacterium adolescentis		3.20	0.22				
Lactobacillus fermentum		0.73	0.16				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07157502	**Low discrimination**					**07157502**
Bifidobacterium adolescentis		68.73	0.56	RAF 92		
Actinomyces israelii		31.11	0.56			
Lactobacillus fermentum		0.16	0.16			
07157510	**Low discrimination**					**07157510**
Actinomyces israelii		78.50	0.39	SAL 92		
Bifidobacterium adolescentis		21.50	0.23	MLZ 13		
07157512	**Good identification**					**07157512**
					SPO	
Bifidobacterium adolescentis		94.65	0.57	RAF 92	1	
Actinomyces israelii		5.24	0.42		1	
Clostridium tertium		0.11	-.01		99	
07157522	**Low discrimination**					**07157522**
					SPO	
Actinomyces israelii		69.71	0.36	SAL 92	1	
Bifidobacterium adolescentis		24.38	0.22	NAG 6	1	
Clostridium tertium		5.92	0.03		99	
07157540	**Good identification**					**07157540**
Actinomyces israelii		100.0	0.46	SAL 92		
07157542	**Low discrimination**					**07157542**
Actinomyces israelii		58.75	0.49	SAL 92		
Bifidobacterium adolescentis		41.25	0.40			
07157552	**Low discrimination**					**07157552**
Bifidobacterium adolescentis		85.17	0.42	RAF 92		
Actinomyces israelii		14.83	0.35	SOR 11		
07157702	**Low discrimination**					**07157702**
Actinomyces israelii		77.13	0.37	SAL 92		
Bifidobacterium adolescentis		22.87	0.22	RHA 7		
07160040	**Good identification**					**07160040**
Clostridium botulinum biovar B		99.95	0.56	CEL 1		
Clostridium botulinum biovar A		0.05	0.02			
07160050	**Good identification**					**07160050**
Clostridium botulinum biovar B		100.0	0.50	CEL 1		
07160060	**Good identification**					**07160060**
Clostridium botulinum biovar B		100.0	0.50	CEL 1		
07160070	**Good identification**					**07160070**
Clostridium botulinum biovar B		100.0	0.45	CEL 1		
07161000	**Low discrimination**					**07161000**
					COC	
Lactobacillus jensenii		84.95	0.44	SAL 95	1	
Gemella morbillorum		7.49	0.26	FRU 99	99	
Atopobium parvulum		3.14	0.18		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07161001	**Low discrimination**						**07161001**
					SPO		
Actinomyces naeslundii		61.66	0.38	RAF 92	1		
Lactobacillus jensenii		25.70	0.10	FRU 95	1		
Clostridium chauvoei		12.65	0.03		99		
07161002	**Acceptable identification**						**07161002**
Lactobacillus jensenii		92.48	0.41	SAL 95			
Lactobacillus acidophilus		3.42	0.14	FRU 99			
Actinomyces naeslundii		2.24	0.35				
07161003	**Low discrimination**						**07161003**
Actinomyces naeslundii		78.26	0.38	RAF 92			
Lactobacillus jensenii		21.74	0.07	FRU 95			
07161010	**Good identification**						**07161010**
					CAT		
Lactobacillus jensenii		98.89	0.36	SAL 95	1		
Propionibacterium granulosum		1.12	0.09	FRU 99	90		
07161021	**Good identification**						**07161021**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07161023	**Good identification**						**07161023**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07161100	**Good identification**						**07161100**
					COC		
Atopobium parvulum		97.36	0.52	FRU 99	99		
Actinomyces naeslundii		1.22	0.43		1		
Clostridium perfringens		0.72	0.16		1		
07161101	**Low discrimination**						**07161101**
					COC	SPO	
Clostridium chauvoei		82.29	0.37	TRE 10	1	99	
Actinomyces naeslundii		11.53	0.46	CEL 1	1	1	
Atopobium parvulum		6.18	0.18		99	1	
07161102	**Low discrimination**						**07161102**
					COC		
Atopobium parvulum		64.90	0.39	FRU 99	99		
Lactobacillus acidophilus		28.52	0.32		1		
Actinomyces naeslundii		4.62	0.43		1		
07161120	**Low discrimination**						**07161120**
					COC	GRA	
Clostridium perfringens		51.66	0.42	FRU 99	1	99	
Atopobium parvulum		38.02	0.39	CEL 4	99	99	
Prevotella melaninogenica		4.30	0.17		1	1	
07161122	**Low discrimination**						**07161122**
					COC		
Clostridium perfringens		45.81	0.37	FRU 99	1		
Lactobacillus acidophilus		31.48	0.32	CEL 4	1		
Atopobium parvulum		12.64	0.26		99		
07161123	**Low discrimination**						**07161123**
					SPO		
Actinomyces naeslundii		67.56	0.46	RAF 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		28.36	0.16	FRU 95	99		
Clostridium perfringens		4.08	0.03		84		
07163000	**Low discrimination**						**07163000**
					CAT		
Actinomyces naeslundii		46.62	0.40	RAF 92	25		
Propionibacterium granulosum		30.91	0.22	FRU 95	90		
Lactobacillus jensenii		15.01	0.10		1		
07163001	**Good identification**						**07163001**
Actinomyces naeslundii		100.0	0.43	RAF 92			
07163002	**Low discrimination**						**07163002**
					CAT		
Actinomyces naeslundii		64.92	0.40	RAF 92	25		
Lactobacillus jensenii		13.94	0.07	FRU 95	1		
Propionibacterium granulosum		10.76	0.12		90		
07163003	**Good identification**						**07163003**
Actinomyces naeslundii		100.0	0.43	RAF 92			
07163020	**Low discrimination**						**07163020**
Actinomyces naeslundii		85.78	0.40	RAF 92			
Propionibacterium granulosum		14.22	0.12	FRU 95			
07163021	**Good identification**						**07163021**
Actinomyces naeslundii		100.0	0.43	RAF 92			
07163022	**Good identification**						**07163022**
Actinomyces naeslundii		100.0	0.40	RAF 92			
07163023	**Good identification**						**07163023**
Actinomyces naeslundii		100.0	0.43	RAF 92			
07163100	**Low discrimination**						**07163100**
					COC	SPO	
Actinomyces naeslundii		68.66	0.48	RAF 92	1	1	
Atopobium parvulum		28.41	0.18	FRU 95	99	1	
Clostridium tertium		2.92	0.01		1	99	
07163101	**Good identification**						**07163101**
					SPO		
Actinomyces naeslundii		96.42	0.51	RAF 92	1		
Clostridium chauvoei		3.58	0.03	FRU 95	99		
07163102	**Low discrimination**						**07163102**
					SPO		
Actinomyces naeslundii		68.80	0.48	RAF 92	1		
Lactobacillus acidophilus		11.50	0.10	FRU 95	1		
Clostridium tertium		8.79	0.09		99		
07163103	**Good identification**						**07163103**
Actinomyces naeslundii		100.0	0.51	RAF 92			
07163120	**Low discrimination**						**07163120**
					SPO		
Actinomyces naeslundii		70.40	0.48	RAF 92	1		
Clostridium tertium		8.99	0.09	FRU 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		8.48	0.07		92		
07163121	**Good identification**						**07163121**
Actinomyces naeslundii		100.0	0.51	RAF 92			
07163122	**Low discrimination**						**07163122**
					SPO		
Actinomyces naeslundii		48.11	0.48	RAF 92	1		
Clostridium ramosum		23.17	0.17	FRU 95	92		
Clostridium tertium		18.43	0.17		99		
07163123	**Good identification**						**07163123**
Actinomyces naeslundii		100.0	0.51	RAF 92			
07163141	**Good identification**						**07163141**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07163143	**Good identification**						**07163143**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07163161	**Good identification**						**07163161**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07163163	**Good identification**						**07163163**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07165100	**Very good identification**						**07165100**
					COC		
Atopobium parvulum		99.92	0.57	FRU 99	99		
Actinomyces israelii		0.08	0.17		1		
07165102	**Good identification**						**07165102**
					COC		
Atopobium parvulum		99.34	0.44	FRU 99	99		
Actinomyces israelii		0.66	0.20		1		
07165120	**Good identification**						**07165120**
					COC		
Atopobium parvulum		99.27	0.44	FRU 99	99		
Clostridium perfringens		0.73	0.08		1		
07170000	**Low discrimination**						**07170000**
					SPO	CAT	
Propionibacterium granulosum		57.56	0.58	CEL 1	1	90	
Lactobacillus jensenii		21.09	0.44		1	1	
Clostridium botulinum biovar B		9.05	0.32		99	1	
07170002	**Low discrimination**						**07170002**
					GRA	CAT	
Propionibacterium granulosum		47.03	0.48	CEL 1	99	90	
Lactobacillus jensenii		45.96	0.41		99	1	
Fusobacterium mortiferum		2.35	0.29		1	5	
07170003	**Low discrimination**						**07170003**
					CAT		
Actinomyces naeslundii		48.85	0.36	RAF 92	25		
Propionibacterium granulosum		17.55	0.14	GAL 96	90		
Lactobacillus jensenii		17.14	0.07		1		
07170010	**Low discrimination**						**07170010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO	CAT	
Propionibacterium granulosum		40.46	0.43	CEL 1	1	90	
Lactobacillus jensenii		36.24	0.36	MLZ 12	1	1	
Clostridium botulinum biovar B		22.97	0.26		99	1	
07170020	**Low discrimination**						**07170020**
					SPO	CAT	
Propionibacterium granulosum		52.17	0.48	CEL 1	1	90	
Lactobacillus jensenii		19.12	0.34		1	1	
Clostridium botulinum biovar B		16.16	0.26		99	1	
07170021	**Low discrimination**						**07170021**
Actinomyces naeslundii		73.57	0.36	RAF 92			
Propionibacterium granulosum		26.43	0.14	GAL 96			
07170022	**Low discrimination**						**07170022**
					CAT		
Propionibacterium granulosum		38.79	0.38	CEL 1	90		
Lactobacillus jensenii		37.91	0.31		1		
Clostridium perfringens		12.80	0.22		1		
07170023	**Good identification**						**07170023**
Actinomyces naeslundii		100.0	0.36	RAF 92			
07170040	**Very good identification**						**07170040**
					SPO	CAT	
Clostridium botulinum biovar B		99.90	0.66	CEL 1	99	1	
Propionibacterium granulosum		0.07	0.24		1	90	
Lactobacillus jensenii		0.02	0.10		1	1	
07170050	**Very good identification**						**07170050**
					SPO	CAT	
Clostridium botulinum biovar B		99.97	0.61	CEL 1	99	1	
Propionibacterium granulosum		0.02	0.09		1	90	
Lactobacillus jensenii		0.02	0.02		1	1	
07170060	**Very good identification**						**07170060**
					CAT	SPO	
Clostridium botulinum biovar B		99.86	0.61	CEL 1	1	99	
Clostridium perfringens		0.11	0.15		1	84	
Propionibacterium granulosum		0.03	0.14		90	1	
07170070	**Good identification**						**07170070**
Clostridium botulinum biovar B		100.0	0.55	CEL 1			
07170100	**Low discrimination**						**07170100**
					COC	GRA	
Atopobium parvulum		81.67	0.52	GAL 99	99	99	
Fusobacterium mortiferum		15.98	0.51		1	1	
Actinomyces naeslundii		0.81	0.41		1	99	
07170101	**Low discrimination**						**07170101**
					COC	SPO	
Actinomyces naeslundii		49.47	0.44	RAF 92	1	1	
Atopobium parvulum		33.47	0.18	GAL 96	99	1	
Clostridium chauvoei		10.51	0.09		1	99	
07170102	**Low discrimination**						**07170102**
					COC	GRA	
Atopobium parvulum		57.72	0.39	GAL 99	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		25.36	0.32		1	99	
Fusobacterium mortiferum		7.11	0.35		1	1	
07170103	**Low discrimination**						**07170103**
					COC		
Actinomyces naeslundii		75.19	0.44	RAF 92	1		
Actinomyces viscosus		15.83	0.08	GAL 96	1		
Atopobium parvulum		8.98	0.05		99		
07170120	**Low discrimination**						**07170120**
					COC	SPO	
Clostridium perfringens		53.33	0.42	CEL 4	1	84	
Atopobium parvulum		39.24	0.39	GAL 99	99	1	
Clostridium septicum		3.42	0.15		1	99	
07170121	**Low discrimination**						**07170121**
					COC		
Actinomyces naeslundii		73.94	0.44	RAF 92	1		
Clostridium perfringens		12.00	0.08	GAL 96	1		
Atopobium parvulum		8.83	0.05		99		
07170122	**Low discrimination**						**07170122**
					COC		
Clostridium perfringens		45.93	0.37	CEL 4	1		
Lactobacillus acidophilus		31.56	0.32	GAL 99	1		
Atopobium parvulum		12.67	0.26		99		
07170123	**Acceptable identification**						**07170123**
Actinomyces naeslundii		92.91	0.44	RAF 92			
Clostridium perfringens		7.09	0.03	GAL 96			
07171000	**Good identification**						**07171000**
					COC	CAT	
Lactobacillus jensenii		92.67	0.78	SAL 95	1	1	
Atopobium parvulum		3.42	0.52		99	1	
Propionibacterium granulosum		2.55	0.58		1	90	
07171001	**Low discrimination**						**07171001**
					COC		
Lactobacillus jensenii		64.73	0.44	SAL 95	1		
Actinomyces naeslundii		29.81	0.60	URE 1	1		
Atopobium parvulum		2.39	0.18		99		
07171002	**Good identification**						**07171002**
					CAT		
Lactobacillus jensenii		93.59	0.75	SAL 95	1		
Lactobacillus acidophilus		3.46	0.48		1		
Propionibacterium granulosum		0.97	0.48		90		
07171003	**Low discrimination**						**07171003**
Lactobacillus jensenii		57.02	0.41	SAL 95			
Actinomyces naeslundii		39.39	0.60	URE 1			
Lactobacillus acidophilus		2.11	0.14				
07171010	**Very good identification**						**07171010**
					COC	CAT	
Lactobacillus jensenii		98.70	0.70	SAL 95	1	1	
Propionibacterium granulosum		1.11	0.43		1	90	
Atopobium parvulum		0.11	0.18		99	1	
07171011	**Acceptable identification**						**07171011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Lactobacillus jensenii		92.25	0.36	SAL 95	1	
Actinomyces naeslundii		6.71	0.38	URE 1	25	
Propionibacterium granulosum		1.04	0.09		90	
07171012	**Very good identification**					**07171012**
					CAT	
Lactobacillus jensenii		99.32	0.67	SAL 95	1	
Propionibacterium granulosum		0.42	0.33		90	
Lactobacillus acidophilus		0.11	0.14		1	
07171020	**Low discrimination**					**07171020**
					CAT	
Lactobacillus jensenii		64.41	0.68	SAL 95	1	
Clostridium perfringens		30.82	0.62		1	
Propionibacterium granulosum		1.78	0.48		90	
07171021	**Low discrimination**					**07171021**
Actinomyces naeslundii		54.34	0.60	RAF 92		
Lactobacillus jensenii		29.50	0.34			
Clostridium perfringens		14.12	0.27			
07171022	**Low discrimination**					**07171022**
Lactobacillus jensenii		64.90	0.65	SAL 95		
Clostridium perfringens		21.92	0.56			
Lactobacillus acidophilus		9.60	0.48			
07171023	**Low discrimination**					**07171023**
Actinomyces naeslundii		65.03	0.60	RAF 92		
Lactobacillus jensenii		23.54	0.31			
Clostridium perfringens		7.95	0.22			
07171030	**Very good identification**					**07171030**
					CAT	
Lactobacillus jensenii		97.23	0.60	SAL 95	1	
Clostridium perfringens		1.41	0.27		1	
Propionibacterium granulosum		1.10	0.33		90	
07171032	**Very good identification**					**07171032**
Lactobacillus jensenii		97.86	0.57	SAL 95		
Clostridium perfringens		1.00	0.22			
Lactobacillus acidophilus		0.44	0.14			
07171040	**Low discrimination**					**07171040**
					SPO	
Lactobacillus jensenii		61.37	0.44	SAL 95	1	
Clostridium botulinum biovar B		26.33	0.32	SOR 1	99	
Clostridium perfringens		4.28	0.23		84	
07171041	**Low discrimination**					**07171041**
					SPO	
Actinomyces naeslundii		84.92	0.47	RAF 92	1	
Lactobacillus jensenii		10.56	0.10		1	
Clostridium botulinum biovar B		4.53	-.03		99	
07171042	**Low discrimination**					**07171042**
Lactobacillus jensenii		61.47	0.41	SAL 95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		27.82	0.32	SOR 1			
Actinomyces naeslundii		5.00	0.44				
07171043	**Low discrimination**						**07171043**
Actinomyces naeslundii		89.26	0.47	RAF 92			
Lactobacillus jensenii		7.40	0.07				
Lactobacillus acidophilus		3.35	-.02				
07171050	**Low discrimination**						**07171050**
					SPO	CAT	
Lactobacillus jensenii		60.78	0.36	SAL 95	1	1	
Clostridium botulinum biovar B		38.53	0.26	SOR 1	99	1	
Propionibacterium granulosum		0.69	0.09		1	90	
07171060	**Low discrimination**						**07171060**
					SPO		
Clostridium perfringens		80.78	0.49	CEL 4	84		
Lactobacillus jensenii		8.95	0.34	LAC 88	1		
Clostridium botulinum biovar B		7.57	0.26		99		
07171061	**Low discrimination**						**07171061**
Actinomyces naeslundii		78.10	0.47	RAF 92			
Clostridium perfringens		21.90	0.15				
07171062	**Low discrimination**						**07171062**
Clostridium perfringens		66.40	0.44	CEL 4			
Lactobacillus acidophilus		18.88	0.32	LAC 88			
Lactobacillus jensenii		10.43	0.31				
07171063	**Low discrimination**						**07171063**
Actinomyces naeslundii		85.51	0.47	RAF 92			
Clostridium perfringens		11.28	0.10				
Lactobacillus acidophilus		3.21	-.02				
07171100	**Very good identification**						**07171100**
					COC	GRA	
Atopobium parvulum		98.28	0.86		99	99	
Clostridium perfringens		0.73	0.51		1	99	
Fusobacterium mortiferum		0.29	0.54		1	1	
07171101	**Low discrimination**						**07171101**
					COC	SPO	
Atopobium parvulum		59.33	0.52	URE 1	99	1	
Actinomyces naeslundii		21.26	0.68		1	1	
Clostridium chauvoei		18.63	0.44		1	99	
07171102	**Low discrimination**						**07171102**
					COC		
Atopobium parvulum		66.89	0.73		99		
Lactobacillus acidophilus		29.39	0.66		1		
Clostridium perfringens		1.32	0.45		1		
07171103	**Low discrimination**						**07171103**
					COC		
Actinomyces naeslundii		53.09	0.68	RAF 92	1		
Atopobium parvulum		26.14	0.39		99		
Lactobacillus acidophilus		11.49	0.32		1		
07171110	**Low discrimination**						**07171110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Atopobium parvulum		89.15	0.52	MLZ 1	99		
Lactobacillus jensenii		8.13	0.36		1		
Actinomyces naeslundii		1.12	0.43		1		
07171111	**Low discrimination**						**07171111**
					COC	SPO	
Actinomyces naeslundii		57.06	0.46	RAF 92	1	1	
Atopobium parvulum		30.56	0.18	MLZ 5	99	1	
Clostridium chauvoei		9.60	0.09		1	99	
07171113	**Low discrimination**						**07171113**
					COC	GRA	
Actinomyces naeslundii		80.08	0.46	RAF 92	1	99	
Leptotrichia buccalis		12.36	-.03	MLZ 5	1	1	
Atopobium parvulum		7.57	0.05		99	99	
07171120	**Low discrimination**						**07171120**
					COC	SPO	
Clostridium perfringens		54.58	0.76	CEL 4	1	84	
Atopobium parvulum		40.16	0.73		99	1	
Clostridium septicum		3.50	0.49		1	99	
07171121	**Low discrimination**						**07171121**
					COC		
Actinomyces naeslundii		39.97	0.68	RAF 92	1		
Clostridium perfringens		26.75	0.42		1		
Atopobium parvulum		19.68	0.39		99		
07171122	**Low discrimination**						**07171122**
					COC		
Clostridium perfringens		48.20	0.71	CEL 4	1		
Lactobacillus acidophilus		33.12	0.66		1		
Atopobium parvulum		13.30	0.61		99		
07171123	**Low discrimination**						**07171123**
Actinomyces naeslundii		59.06	0.68	RAF 92			
Clostridium perfringens		18.60	0.37				
Lactobacillus acidophilus		12.78	0.32				
07171130	**Low discrimination**						**07171130**
					COC		
Clostridium perfringens		50.55	0.42	CEL 4	1		
Atopobium parvulum		37.20	0.39	MLZ 1	99		
Lactobacillus jensenii		4.80	0.26		1		
07171131	**Low discrimination**						**07171131**
					COC		
Actinomyces naeslundii		78.18	0.46	RAF 92	1		
Clostridium perfringens		10.04	0.08	MLZ 5	1		
Atopobium parvulum		7.39	0.05		99		
07171132	**Low discrimination**						**07171132**
					GRA		
Clostridium perfringens		36.01	0.37	CEL 4	99		
Lactobacillus acidophilus		24.75	0.32	MLZ 1	99		
Leptotrichia buccalis		10.22	0.15		1		
07171133	**Acceptable identification**						**07171133**
Actinomyces naeslundii		94.30	0.46	RAF 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		5.70	0.03	MLZ 5			
07171140	**Low discrimination**						**07171140**
					COC		
Atopobium parvulum		83.01	0.52	SOR 1	99		
Clostridium perfringens		11.61	0.38		1		
Actinomyces naeslundii		3.50	0.52		1		
07171141	**Low discrimination**						**07171141**
					COC	SPO	
Actinomyces naeslundii		81.15	0.55	RAF 92	1	1	
Atopobium parvulum		12.96	0.18		99	1	
Clostridium chauvoei		4.07	0.09		1	99	
07171142	**Low discrimination**						**07171142**
					COC		
Lactobacillus acidophilus		75.99	0.50	SAL 99	1		
Atopobium parvulum		14.13	0.39	SOR 11	99		
Clostridium perfringens		5.27	0.33		1		
07171143	**Low discrimination**						**07171143**
					COC		
Actinomyces naeslundii		84.76	0.55	RAF 92	1		
Lactobacillus acidophilus		12.85	0.16		1		
Atopobium parvulum		2.39	0.05		99		
07171160	**Good identification**						**07171160**
					COC		
Clostridium perfringens		94.69	0.64	CEL 4	1		
Atopobium parvulum		3.70	0.39		99		
Actinomyces naeslundii		0.88	0.52		1		
07171161	**Low discrimination**						**07171161**
					COC		
Actinomyces naeslundii		56.59	0.55	RAF 92	1		
Clostridium perfringens		40.87	0.30		1		
Atopobium parvulum		1.60	0.05		99		
07171162	**Low discrimination**						**07171162**
Clostridium perfringens		67.11	0.58	CEL 4			
Lactobacillus acidophilus		29.92	0.50				
Actinomyces naeslundii		1.33	0.52				
07171163	**Low discrimination**						**07171163**
Actinomyces naeslundii		67.05	0.55	RAF 92			
Clostridium perfringens		22.79	0.24				
Lactobacillus acidophilus		10.16	0.16				
07171200	**Low discrimination**						**07171200**
					COC	CAT	
Lactobacillus jensenii		88.28	0.44	SAL 95	1	1	
Propionibacterium granulosum		7.45	0.32	RHA 1	1	90	
Atopobium parvulum		3.26	0.18		99	1	
07171202	**Low discrimination**						**07171202**
					CAT		
Lactobacillus jensenii		75.25	0.41	SAL 95	1		
Lactobacillus acidophilus		20.74	0.29	RHA 1	1		
Propionibacterium granulosum		2.38	0.22		90		
07171210	**Good identification**						**07171210**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Lactobacillus jensenii		96.66	0.36	SAL 95	1		
Propionibacterium granulosum		3.34	0.17	RHA 1	90		
07171300	**Good identification**						**07171300**
					COC		
Atopobium parvulum		97.69	0.52	RHA 1	99		
Clostridium perfringens		0.73	0.16		1		
Lactobacillus acidophilus		0.57	0.12		1		
07171302	**Low discrimination**						**07171302**
					COC	GRA	
Lactobacillus acidophilus		73.84	0.47	SAL 99	1	99	
Atopobium parvulum		22.55	0.39	RHA 7	99	99	
Prevotella oralis		1.43	0.16		1	1	
07171320	**Low discrimination**						**07171320**
					COC	SPO	
Clostridium perfringens		45.69	0.42	CEL 4	1	84	
Atopobium parvulum		33.62	0.39	RHA 1	99	1	
Clostridium ramosum		12.97	0.30		1	92	
07171322	**Low discrimination**						**07171322**
					SPO	GRA	
Lactobacillus acidophilus		45.84	0.47	SAL 99	1	99	
Clostridium ramosum		21.60	0.40	RHA 7	92	50	
Prevotella oralis		16.84	0.38		1	1	
07171400	**Acceptable identification**						**07171400**
					COC	CAT	
Lactobacillus jensenii		90.71	0.44	SAL 95	1	1	
Atopobium parvulum		3.35	0.18	ARA 1	99	1	
Propionibacterium granulosum		2.50	0.24		1	90	
07171402	**Acceptable identification**						**07171402**
Lactobacillus jensenii		91.71	0.41	SAL 95			
Lactobacillus acidophilus		3.39	0.14	ARA 1			
Actinomyces israelii		1.92	0.25				
07171410	**Good identification**						**07171410**
					CAT		
Lactobacillus jensenii		98.89	0.36	SAL 95	1		
Propionibacterium granulosum		1.12	0.09	ARA 1	90		
07171500	**Good identification**						**07171500**
					COC		
Atopobium parvulum		96.82	0.52	ARA 1	99		
Actinomyces israelii		0.99	0.32		1		
Clostridium perfringens		0.72	0.16		1		
07171502	**Low discrimination**						**07171502**
					COC		
Atopobium parvulum		49.60	0.39	ARA 1	99		
Lactobacillus acidophilus		21.79	0.32		1		
Bifidobacterium adolescentis		20.03	0.40		1		
07171512	**Low discrimination**						**07171512**
					GRA		
Bifidobacterium adolescentis		80.56	0.42	RAF 92	99		
Clostridium clostridioforme		11.28	0.24	XYL 87	50		
Leptotrichia buccalis		2.69	-.03		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07171520	**Low discrimination**						**07171520**
					COC	SPO	
Clostridium perfringens		54.47	0.42	CEL 4	1	84	
Atopobium parvulum		40.08	0.39	ARA 1	99	1	
Clostridium septicum		3.50	0.15		1	99	
07171522	**Low discrimination**						**07171522**
					COC		
Clostridium perfringens		45.45	0.37	CEL 4	1		
Lactobacillus acidophilus		31.23	0.32	ARA 1	1		
Atopobium parvulum		12.54	0.26		99		
07171712	**Good identification**						**07171712**
Clostridium clostridioforme		99.03	0.39	SAL 88			
Bifidobacterium adolescentis		0.97	0.07	RAF 94			
07171732	**Low discrimination**						**07171732**
Clostridium clostridioforme		87.46	0.39	SAL 88			
Parabacteroides distasonis		12.54	0.30	RAF 94			
07172000	**Good identification**						**07172000**
					CAT		
Propionibacterium granulosum		97.75	0.56	CEL 1	90		
Actinomyces naeslundii		1.18	0.38		25		
Lactobacillus jensenii		0.48	0.10		1		
07172001	**Low discrimination**						**07172001**
Actinomyces naeslundii		64.17	0.41	RAF 92			
Propionibacterium granulosum		35.83	0.22	GAL 96			
07172002	**Acceptable identification**						**07172002**
					CAT		
Propionibacterium granulosum		92.74	0.46	CEL 1	90		
Actinomyces naeslundii		4.47	0.38		25		
Lactobacillus jensenii		1.21	0.07		1		
07172003	**Low discrimination**						**07172003**
Actinomyces naeslundii		87.75	0.41	RAF 92			
Propionibacterium granulosum		12.25	0.12	GAL 96			
07172010	**Good identification**						**07172010**
					CAT		
Propionibacterium granulosum		98.82	0.41	CEL 1	90		
Lactobacillus jensenii		1.19	0.02	MLZ 12	1		
07172020	**Good identification**						**07172020**
Propionibacterium granulosum		95.40	0.46	CEL 1			
Actinomyces naeslundii		4.60	0.38				
07172021	**Low discrimination**						**07172021**
Actinomyces naeslundii		87.75	0.41	RAF 92			
Propionibacterium granulosum		12.25	0.12	GAL 96			
07172022	**Low discrimination**						**07172022**
Propionibacterium granulosum		83.82	0.35	CEL 1			
Actinomyces naeslundii		16.18	0.38				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07172023	**Good identification**					**07172023**
Actinomyces naeslundii		100.0	0.41	RAF 92		
07172100	**Low discrimination**					**07172100**
					COC	
Propionibacterium propionicum		63.80	0.45	NIT 86	1	
Actinomyces naeslundii		18.35	0.46	CEL 1	1	
Atopobium parvulum		9.59	0.18		99	
07172101	**Low discrimination**					**07172101**
Actinomyces naeslundii		79.52	0.49	RAF 92		
Propionibacterium propionicum		20.48	0.29	GAL 96		
07172102	**Low discrimination**					**07172102**
Propionibacterium propionicum		76.59	0.48	NIT 86		
Actinomyces naeslundii		14.69	0.46	CEL 1		
Lactobacillus acidophilus		3.10	0.10			
07172103	**Low discrimination**					**07172103**
Actinomyces naeslundii		72.13	0.49	RAF 92		
Propionibacterium propionicum		27.87	0.32	GAL 96		
07172120	**Low discrimination**					**07172120**
					SPO	
Actinomyces naeslundii		62.32	0.46	RAF 92	1	
Clostridium tertium		10.06	0.09	GAL 96	99	
Clostridium ramosum		9.48	0.07		92	
07172121	**Good identification**					**07172121**
Actinomyces naeslundii		100.0	0.49	RAF 92		
07172122	**Low discrimination**					**07172122**
					SPO	
Actinomyces naeslundii		41.40	0.46	RAF 92	1	
Clostridium ramosum		25.19	0.17	GAL 96	92	
Clostridium tertium		20.04	0.17		99	
07172123	**Good identification**					**07172123**
Actinomyces naeslundii		100.0	0.49	RAF 92		
07172140	**Acceptable identification**					**07172140**
					SPO	
Propionibacterium propionicum		93.60	0.45	NIT 86	1	
Actinomyces naeslundii		4.75	0.33	CEL 1	1	
Clostridium botulinum biovar B		1.65	-.03		99	
07172142	**Good identification**					**07172142**
Propionibacterium propionicum		96.73	0.48	NIT 86		
Actinomyces naeslundii		3.27	0.33	CEL 1		
07172161	**Good identification**					**07172161**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07172163	**Good identification**					**07172163**
Actinomyces naeslundii		100.0	0.36	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07173000	**Low discrimination**					**07173000**
					CAT	
Propionibacterium granulosum		53.29	0.56	CEL 1	90	
Lactobacillus jensenii		25.88	0.44		1	
Actinomyces naeslundii		15.42	0.62		25	
07173001	**Very good identification**					**07173001**
					CAT	
Actinomyces naeslundii		96.66	0.65	RAF 92	25	
Propionibacterium granulosum		2.25	0.22		90	
Lactobacillus jensenii		1.09	0.10		1	
07173002	**Low discrimination**					**07173002**
					CAT	
Lactobacillus jensenii		31.38	0.41	SAL 95	1	
Actinomyces naeslundii		28.05	0.62	MAN 1	25	
Propionibacterium granulosum		24.23	0.46		90	
07173003	**Very good identification**					**07173003**
					CAT	
Actinomyces naeslundii		98.68	0.65	RAF 92	25	
Lactobacillus jensenii		0.74	0.07		1	
Propionibacterium granulosum		0.57	0.12		90	
07173010	**Low discrimination**					**07173010**
					CAT	
Lactobacillus jensenii		50.06	0.36	SAL 95	1	
Propionibacterium granulosum		42.17	0.41	MAN 1	90	
Actinomyces naeslundii		4.71	0.40		25	
07173011	**Acceptable identification**					**07173011**
Actinomyces naeslundii		93.32	0.43	RAF 92		
Lactobacillus jensenii		6.68	0.02	MLZ 5		
07173013	**Good identification**					**07173013**
Actinomyces naeslundii		100.0	0.43	RAF 92		
07173020	**Low discrimination**					**07173020**
					CAT	
Actinomyces naeslundii		39.26	0.62	RAF 92	25	
Propionibacterium granulosum		33.91	0.46		90	
Lactobacillus jensenii		16.47	0.34		1	
07173021	**Very good identification**					**07173021**
Actinomyces naeslundii		99.42	0.65	RAF 92		
Propionibacterium granulosum		0.58	0.12			
07173022	**Low discrimination**					**07173022**
					CAT	
Actinomyces naeslundii		50.77	0.62	RAF 92	25	
Lactobacillus jensenii		14.20	0.31		1	
Propionibacterium granulosum		10.96	0.35		90	
07173023	**Very good identification**					**07173023**
Actinomyces naeslundii		100.0	0.65	RAF 92		
07173031	**Good identification**					**07173031**
Actinomyces naeslundii		100.0	0.43	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07173033	**Good identification**						**07173033**
Actinomyces naeslundii		100.0	0.43	RAF 92			
07173040	**Low discrimination**						**07173040**
					CAT		
Actinomyces naeslundii		63.54	0.49	RAF 92	25		
Actinomyces israelii		15.17	0.29		1		
Propionibacterium granulosum		12.57	0.22		90		
07173041	**Good identification**						**07173041**
Actinomyces naeslundii		100.0	0.52	RAF 92			
07173042	**Low discrimination**						**07173042**
Actinomyces naeslundii		60.31	0.49	RAF 92			
Actinomyces israelii		21.60	0.32				
Lactobacillus acidophilus		9.11	0.10				
07173043	**Good identification**						**07173043**
Actinomyces naeslundii		100.0	0.52	RAF 92			
07173060	**Low discrimination**						**07173060**
					CAT		
Actinomyces naeslundii		78.99	0.49	RAF 92	25		
Clostridium perfringens		17.11	0.15		1		
Propionibacterium granulosum		3.90	0.12		90		
07173061	**Good identification**						**07173061**
Actinomyces naeslundii		100.0	0.52	RAF 92			
07173062	**Low discrimination**						**07173062**
Actinomyces naeslundii		79.81	0.49	RAF 92			
Lactobacillus acidophilus		12.05	0.10				
Clostridium perfringens		8.14	0.10				
07173063	**Good identification**						**07173063**
Actinomyces naeslundii		100.0	0.52	RAF 92			
07173100	**Low discrimination**						**07173100**
					COC	SPO	
Atopobium parvulum		57.75	0.52	MAN 1	99	1	
Actinomyces naeslundii		26.78	0.69		1	1	
Clostridium tertium		5.94	0.35		1	99	
07173101	**Very good identification**						**07173101**
					COC		
Actinomyces naeslundii		96.97	0.72	RAF 92	1		
Atopobium parvulum		1.41	0.18		99		
Propionibacterium propionicum		1.04	0.29		1		
07173102	**Low discrimination**						**07173102**
					SPO		
Actinomyces naeslundii		29.15	0.69	RAF 92	1		
Lactobacillus acidophilus		25.40	0.44		1		
Clostridium tertium		19.40	0.43		99		
07173103	**Very good identification**						**07173103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.04	0.72	RAF 92		
Propionibacterium propionicum		1.56	0.32			
Lactobacillus acidophilus		0.57	0.10			
07173110	**Low discrimination**					**07173110**
					SPO	
Clostridium tertium		79.91	0.39	SAL 99	99	
Actinomyces naeslundii		10.66	0.48		1	
Actinomyces israelii		4.61	0.32		1	
07173111	**Good identification**					**07173111**
					SPO	
Actinomyces naeslundii		95.20	0.51	RAF 92	1	
Clostridium tertium		4.80	0.05	MLZ 5	99	
07173112	**Acceptable identification**					**07173112**
					SPO	
Clostridium tertium		90.04	0.47	SAL 99	99	
Actinomyces naeslundii		4.01	0.48		1	
Actinomyces israelii		2.60	0.35		1	
07173113	**Low discrimination**					**07173113**
					SPO	
Actinomyces naeslundii		86.85	0.51	RAF 92	1	
Clostridium tertium		13.15	0.13	MLZ 5	99	
07173120	**Low discrimination**					**07173120**
					SPO	
Actinomyces naeslundii		30.70	0.69	RAF 92	1	
Clostridium tertium		20.43	0.43		99	
Clostridium ramosum		19.26	0.41		92	
07173121	**Very good identification**					**07173121**
					SPO	
Actinomyces naeslundii		98.40	0.72	RAF 92	1	
Clostridium tertium		0.44	0.09		99	
Clostridium ramosum		0.42	0.07		92	
07173122	**Low discrimination**					**07173122**
					SPO	
Clostridium ramosum		37.23	0.51	SAL 99	92	
Clostridium tertium		29.62	0.51		99	
Actinomyces naeslundii		14.84	0.69		1	
07173123	**Very good identification**					**07173123**
					SPO	
Actinomyces naeslundii		96.35	0.72	RAF 92	1	
Clostridium ramosum		1.63	0.17		92	
Clostridium tertium		1.30	0.17		99	
07173130	**Acceptable identification**					**07173130**
					SPO	
Clostridium tertium		94.56	0.47	SAL 99	99	
Actinomyces naeslundii		4.21	0.48		1	
Clostridium ramosum		0.51	0.07		92	
07173131	**Low discrimination**					**07173131**
					SPO	
Actinomyces naeslundii		86.85	0.51	RAF 92	1	
Clostridium tertium		13.15	0.13	MLZ 5	99	
07173132	**Good identification**					**07173132**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		97.49	0.55	SAL 99	99	
Actinomyces naeslundii		1.45	0.48		1	
Clostridium ramosum		0.70	0.17		92	
07173133	**Low discrimination**					**07173133**
					SPO	
Actinomyces naeslundii		68.77	0.51	RAF 92	1	
Clostridium tertium		31.23	0.21	MLZ 5	99	
07173140	**Low discrimination**					**07173140**
Actinomyces naeslundii		43.29	0.57	RAF 92		
Propionibacterium propionicum		35.54	0.45			
Actinomyces israelii		14.53	0.39			
07173141	**Good identification**					**07173141**
Actinomyces naeslundii		94.27	0.60	RAF 92		
Propionibacterium propionicum		5.73	0.29			
07173142	**Low discrimination**					**07173142**
Propionibacterium propionicum		36.03	0.48	NIT 86		
Actinomyces naeslundii		29.26	0.57	CEL 1		
Lactobacillus acidophilus		17.85	0.28			
07173143	**Good identification**					**07173143**
Actinomyces naeslundii		91.64	0.60	RAF 92		
Propionibacterium propionicum		8.36	0.32			
07173151	**Good identification**					**07173151**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07173153	**Good identification**					**07173153**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07173160	**Low discrimination**					**07173160**
					SPO	
Actinomyces naeslundii		59.69	0.57	RAF 92	1	
Clostridium perfringens		33.32	0.30		84	
Clostridium tertium		2.27	0.09		99	
07173161	**Very good identification**					**07173161**
Actinomyces naeslundii		100.0	0.60	RAF 92		
07173162	**Low discrimination**					**07173162**
Actinomyces naeslundii		45.71	0.57	RAF 92		
Lactobacillus acidophilus		27.89	0.28			
Clostridium perfringens		12.01	0.24			
07173163	**Very good identification**					**07173163**
Actinomyces naeslundii		100.0	0.60	RAF 92		
07173171	**Good identification**					**07173171**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07173173	**Good identification**					**07173173**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	RAF 92		
07173301	**Good identification**					**07173301**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07173303	**Good identification**					**07173303**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07173320	**Good identification**					**07173320**
					SPO	
Clostridium ramosum		96.98	0.43	SAL 99	92	
Actinomyces naeslundii		1.18	0.35		1	
Clostridium tertium		0.78	0.09		99	
07173321	**Low discrimination**					**07173321**
					SPO	
Actinomyces naeslundii		64.33	0.38	RAF 92	1	
Clostridium ramosum		35.67	0.09	RHA 1	92	
07173322	**Good identification**					**07173322**
					SPO	
Clostridium ramosum		96.94	0.53	SAL 99	92	
Lactobacillus acidophilus		1.91	0.25		1	
Clostridium tertium		0.59	0.17		99	
07173402	**Low discrimination**					**07173402**
					SPO	
Actinomyces israelii		69.50	0.37	SAL 92	1	
Clostridium innocuum		14.75	0.11	XYL 92	99	
Lactobacillus jensenii		5.91	0.07		1	
07173500	**Low discrimination**					**07173500**
					COC	
Actinomyces israelii		78.50	0.45	SAL 92	1	
Atopobium parvulum		13.72	0.18	XYL 92	99	
Actinomyces naeslundii		6.36	0.35		1	
07173501	**Good identification**					**07173501**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07173502	**Low discrimination**					**07173502**
Actinomyces israelii		54.28	0.48	SAL 92		
Bifidobacterium adolescentis		36.52	0.39	XYL 92		
Actinomyces naeslundii		2.93	0.35			
07173503	**Low discrimination**					**07173503**
Actinomyces naeslundii		88.92	0.38	RAF 92		
Actinomyces israelii		11.08	0.14	ARA 1		
07173512	**Low discrimination**					**07173512**
					SPO	
Bifidobacterium adolescentis		79.40	0.40	RAF 92	1	
Actinomyces israelii		14.43	0.34	XYL 87	1	
Clostridium tertium		6.18	0.13		99	
07173520	**Low discrimination**					**07173520**
					SPO	
Actinomyces naeslundii		25.14	0.35	RAF 92	1	
Actinomyces israelii		19.79	0.24	ARA 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		16.73	0.09		99		
07173521	**Good identification**						**07173521**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07173523	**Good identification**						**07173523**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07173540	**Good identification**						**07173540**
Actinomyces israelii		96.45	0.38	SAL 92			
Actinomyces naeslundii		3.55	0.22	XYL 92			
07173542	**Low discrimination**						**07173542**
Actinomyces israelii		78.83	0.41	SAL 92			
Bifidobacterium adolescentis		16.86	0.23	XYL 92			
Propionibacterium propionicum		2.38	0.14				
07174100	**Very good identification**						**07174100**
					COC	GRA	
Atopobium parvulum		99.14	0.57	GAL 99	99	99	
Fusobacterium mortiferum		0.79	0.32		1	1	
Actinomyces israelii		0.08	0.17		1	99	
07174102	**Good identification**						**07174102**
					COC	GRA	
Atopobium parvulum		98.85	0.44	GAL 99	99	99	
Actinomyces israelii		0.66	0.20		1	99	
Fusobacterium mortiferum		0.49	0.16		1	1	
07174120	**Good identification**						**07174120**
					COC		
Atopobium parvulum		99.27	0.44	GAL 99	99		
Clostridium perfringens		0.73	0.08		1		
07174612	**Good identification**						**07174612**
Clostridium clostridioforme		100.0	0.39	SAL 88			
07174632	**Good identification**						**07174632**
Clostridium clostridioforme		100.0	0.39	SAL 88			
07174702	**Good identification**						**07174702**
Clostridium clostridioforme		100.0	0.36	SAL 88			
07174712	**Good identification**						**07174712**
Clostridium clostridioforme		100.0	0.48	SAL 88			
07174722	**Good identification**						**07174722**
Clostridium clostridioforme		96.83	0.36	SAL 88			
Parabacteroides distasonis		3.17	0.17	RAF 94			
07174732	**Good identification**						**07174732**
Clostridium clostridioforme		99.62	0.48	SAL 88			
Parabacteroides distasonis		0.39	0.12	RAF 94			
07175000	**Low discrimination**						**07175000**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		84.83	0.57	LAC 99	99		
Lactobacillus jensenii		12.49	0.44		1		
Actinomyces israelii		1.64	0.41		1		
07175002	**Low discrimination**						**07175002**
					COC		
Atopobium parvulum		56.99	0.44	LAC 99	99		
Lactobacillus jensenii		31.70	0.41		1		
Actinomyces israelii		9.35	0.44		1		
07175010	**Low discrimination**						**07175010**
					COC		
Lactobacillus jensenii		78.38	0.36	SAL 95	1		
Atopobium parvulum		16.13	0.23	XYL 1	99		
Actinomyces israelii		4.60	0.27		1		
07175020	**Low discrimination**						**07175020**
					COC		
Atopobium parvulum		75.55	0.44	LAC 99	99		
Lactobacillus jensenii		15.76	0.34		1		
Clostridium perfringens		7.54	0.27		1		
07175042	**Low discrimination**						**07175042**
					COC		
Actinomyces israelii		77.74	0.37	SAL 92	1		
Atopobium parvulum		12.31	0.10		99		
Lactobacillus jensenii		6.85	0.07		1		
07175100	**Very good identification**						**07175100**
					COC	GRA	
Atopobium parvulum		99.90	0.91		99	99	
Actinomyces israelii		0.08	0.52		1	99	
Fusobacterium mortiferum		0.01	0.35		1	1	
07175101	**Very good identification**						**07175101**
					COC	SPO	
Atopobium parvulum		99.56	0.57	URE 1	99	1	
Actinomyces naeslundii		0.19	0.33		1	1	
Clostridium chauvoei		0.17	0.09		1	99	
07175102	**Very good identification**						**07175102**
					COC		
Atopobium parvulum		98.81	0.78		99		
Actinomyces israelii		0.66	0.55		1		
Lactobacillus acidophilus		0.24	0.32		1		
07175103	**Good identification**						**07175103**
					COC		
Atopobium parvulum		98.26	0.44	URE 1	99		
Actinomyces naeslundii		1.09	0.33		1		
Actinomyces israelii		0.65	0.20		1		
07175110	**Very good identification**						**07175110**
					COC	SPO	
Atopobium parvulum		98.63	0.57	MLZ 1	99	1	
Actinomyces israelii		1.14	0.37		1	1	
Clostridium tertium		0.14	0.07		1	99	
07175112	**Low discrimination**						**07175112**
					COC		
Atopobium parvulum		70.42	0.44	MLZ 1	99		
Bifidobacterium adolescentis		18.72	0.42		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		6.90	0.40		1			
07175120	**Very good identification**							**07175120**
					COC	SPO		
Atopobium parvulum		99.15	0.78		99	1		
Clostridium perfringens		0.73	0.42		1	84		
Clostridium septicum		0.05	0.15		1	99		
07175121	**Good identification**							**07175121**
					COC			
Atopobium parvulum		97.88	0.44	URE 1	99			
Actinomyces naeslundii		1.08	0.33		1			
Clostridium perfringens		0.72	0.08		1			
07175122	**Good identification**							**07175122**
					COC	GRA		
Atopobium parvulum		93.41	0.65		99	99		
Parabacteroides distasonis		1.90	0.46		1	1		
Clostridium perfringens		1.84	0.37		1	99		
07175130	**Good identification**							**07175130**
					COC	SPO		
Atopobium parvulum		96.63	0.44	MLZ 1	99	1		
Clostridium tertium		2.26	0.15		1	99		
Clostridium perfringens		0.71	0.08		1	84		
07175132	**Low discrimination**							**07175132**
					COC	GRA	SPO	
Parabacteroides distasonis		40.37	0.42	RAF 99	1	1	1	
Atopobium parvulum		37.19	0.31		99	99	1	
Clostridium tertium		14.76	0.24		1	99	99	
07175140	**Very good identification**							**07175140**
					COC			
Atopobium parvulum		96.96	0.57	SOR 1	99			
Actinomyces israelii		2.91	0.45		1			
Clostridium perfringens		0.07	0.04		1			
07175142	**Low discrimination**							**07175142**
					COC			
Atopobium parvulum		76.27	0.44	SOR 1	99			
Actinomyces israelii		19.45	0.48		1			
Lactobacillus acidophilus		2.23	0.16		1			
07175160	**Low discrimination**							**07175160**
					COC			
Atopobium parvulum		86.94	0.44	SOR 1	99			
Clostridium perfringens		12.12	0.30		1			
Actinomyces israelii		0.94	0.24		1			
07175300	**Very good identification**							**07175300**
					COC			
Atopobium parvulum		99.42	0.57	RHA 1	99			
Actinomyces israelii		0.58	0.32		1			
07175302	**Low discrimination**							**07175302**
					COC			
Atopobium parvulum		89.22	0.44	RHA 1	99			
Actinomyces israelii		4.41	0.35		1			
Clostridium clostridioforme		3.45	0.24		1			
07175312	**Low discrimination**							**07175312**
					COC			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		87.15	0.35	SAL 88	1		
Atopobium parvulum		5.00	0.10	RAF 94	99		
Actinomyces israelii		3.65	0.21	ARA 94	1		
07175320	**Good identification**						**07175320**
					COC	GRA	
Atopobium parvulum		96.60	0.44	RHA 1	99	99	
Bacteroides vulgatus		1.24	0.03		1	1	
Parabacteroides distasonis		1.04	0.20		1	1	
07175322	**Low discrimination**						**07175322**
					COC	GRA	
Parabacteroides distasonis		76.49	0.54	RAF 99	1	1	
Atopobium parvulum		12.65	0.31		99	99	
Prevotella oralis		5.74	0.22		1	1	
07175332	**Low discrimination**						**07175332**
Parabacteroides distasonis		76.54	0.50	RAF 99			
Clostridium clostridioforme		23.46	0.35				
07175400	**Low discrimination**						**07175400**
					COC	SPO	
Actinomyces israelii		51.73	0.40	SAL 92	1	1	
Atopobium parvulum		33.10	0.23		99	1	
Clostridium butyricum		8.85	0.06		1	99	
07175402	**Low discrimination**						**07175402**
Actinomyces israelii		67.86	0.43	SAL 92			
Clostridium clostridioforme		12.48	0.21				
Bifidobacterium adolescentis		7.44	0.20				
07175442	**Good identification**						**07175442**
Actinomyces israelii		100.0	0.36	SAL 92			
07175500	**Low discrimination**						**07175500**
					COC	SPO	
Atopobium parvulum		88.26	0.57	ARA 1	99	1	
Clostridium butyricum		5.72	0.29		1	99	
Actinomyces israelii		5.57	0.50		1	1	
07175502	**Low discrimination**						**07175502**
					COC		
Bifidobacterium adolescentis		45.04	0.54	RAF 92	1		
Atopobium parvulum		30.95	0.44		99		
Actinomyces israelii		16.61	0.53		1		
07175512	**Low discrimination**						**07175512**
Bifidobacterium adolescentis		78.37	0.56	RAF 92			
Clostridium clostridioforme		16.58	0.41				
Actinomyces israelii		3.53	0.39				
07175520	**Low discrimination**						**07175520**
					COC	SPO	
Atopobium parvulum		89.22	0.44	ARA 1	99	1	
Bifidobacterium longum subsp. longum		3.54	0.26		1	1	
Clostridium butyricum		2.85	0.11		1	99	
07175522	**Low discrimination**						**07175522**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		37.31	0.43	RAF 99	1	1	
Atopobium parvulum		30.19	0.31		99	99	
Clostridium clostridioforme		14.13	0.30		1	50	
07175532	**Low discrimination**						**07175532**
					GRA		
Clostridium clostridioforme		64.40	0.41	SAL 88	50		
Parabacteroides distasonis		20.09	0.38	RAF 94	1		
Bifidobacterium longum subsp. longum		11.56	0.30	RHA 88	99		
07175540	**Low discrimination**						**07175540**
					COC	SPO	
Actinomyces israelii		60.95	0.43	SAL 92	1	1	
Atopobium parvulum		25.07	0.23		99	1	
Clostridium butyricum		13.98	0.11		1	99	
07175542	**Low discrimination**						**07175542**
					SPO		
Actinomyces israelii		50.77	0.46	SAL 92	1		
Bifidobacterium adolescentis		43.76	0.39		1		
Clostridium butyricum		3.02	0.04		99		
07175552	**Low discrimination**						**07175552**
Bifidobacterium adolescentis		86.27	0.40	RAF 92			
Actinomyces israelii		12.24	0.32	SOR 11			
Clostridium clostridioforme		1.49	0.07				
07175602	**Acceptable identification**						**07175602**
Clostridium clostridioforme		94.71	0.36	SAL 88			
Actinomyces israelii		5.29	0.24	RAF 94			
07175612	**Good identification**						**07175612**
Clostridium clostridioforme		100.0	0.47	SAL 88			
07175622	**Good identification**						**07175622**
Clostridium clostridioforme		96.48	0.36	SAL 88			
Parabacteroides distasonis		3.52	0.17	RAF 94			
07175632	**Good identification**						**07175632**
Clostridium clostridioforme		99.57	0.47	SAL 88			
Parabacteroides distasonis		0.43	0.12	RAF 94			
07175702	**Low discrimination**						**07175702**
Clostridium clostridioforme		88.83	0.45	SAL 88			
Actinomyces israelii		5.92	0.34	RAF 94			
Bifidobacterium adolescentis		2.16	0.20				
07175712	**Good identification**						**07175712**
Clostridium clostridioforme		99.01	0.56	SAL 88			
Bifidobacterium adolescentis		0.65	0.22	RAF 94			
Actinomyces israelii		0.22	0.20				
07175722	**Low discrimination**						**07175722**
Parabacteroides distasonis		49.62	0.51	RAF 99			
Clostridium clostridioforme		45.95	0.45				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		2.88	0.14				
07175732	**Low discrimination**						**07175732**
Clostridium clostridioforme		87.93	0.56	SAL 88			
Parabacteroides distasonis		11.22	0.46	RAF 94			
Bacteroides thetaiotaomicron		0.47	0.07				
07177000	**Low discrimination**						**07177000**
					COC	CAT	
Actinomyces israelii		86.51	0.54	SAL 92	1	1	
Atopobium parvulum		7.99	0.23		99	1	
Propionibacterium granulosum		2.42	0.22		1	90	
07177002	**Very good identification**						**07177002**
					COC	SPO	
Actinomyces israelii		95.64	0.57	SAL 92	1	1	
Clostridium tertium		1.35	0.11		1	99	
Atopobium parvulum		1.04	0.10		99	1	
07177010	**Low discrimination**						**07177010**
					SPO		
Actinomyces israelii		89.73	0.40	SAL 92	1		
Clostridium tertium		7.55	0.07	MLZ 13	99		
Lactobacillus jensenii		2.72	0.02		1		
07177012	**Low discrimination**						**07177012**
					SPO		
Actinomyces israelii		85.60	0.43	SAL 92	1		
Clostridium tertium		14.40	0.15	MLZ 13	99		
07177022	**Low discrimination**						**07177022**
					SPO		
Actinomyces israelii		57.17	0.37	SAL 92	1		
Clostridium tertium		37.99	0.19	NAG 6	99		
Actinomyces naeslundii		4.84	0.28		1		
07177040	**Good identification**						**07177040**
Actinomyces israelii		100.0	0.47	SAL 92			
07177042	**Good identification**						**07177042**
Actinomyces israelii		100.0	0.50	SAL 92			
07177052	**Good identification**						**07177052**
Actinomyces israelii		100.0	0.36	SAL 92			
07177100	**Low discrimination**						**07177100**
					COC	SPO	
Atopobium parvulum		65.92	0.57	MAN 1	99	1	
Actinomyces israelii		28.82	0.64		1	1	
Clostridium tertium		5.04	0.37		1	99	
07177102	**Low discrimination**						**07177102**
					COC	SPO	
Actinomyces israelii		59.71	0.67	SAL 92	1	1	
Clostridium tertium		20.89	0.45		1	99	
Atopobium parvulum		16.06	0.44		99	1	
07177110	**Low discrimination**						**07177110**
					COC	SPO	
Clostridium tertium		64.32	0.41	SAL 99	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		30.90	0.50		1	1	
Atopobium parvulum		4.78	0.23		99	1	
07177112	**Low discrimination**						**07177112**
					SPO		
Clostridium tertium		74.70	0.50	SAL 99	99		
Actinomyces israelii		17.95	0.53		1		
Bifidobacterium adolescentis		7.03	0.40		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07177120	**Low discrimination**						**07177120**
					COC	SPO	
Clostridium tertium		52.24	0.45	SAL 99	1	99	
Atopobium parvulum		40.17	0.44		99	1	
Actinomyces israelii		6.35	0.44		1	1	
07177121	**Low discrimination**						**07177121**
					COC	SPO	
Actinomyces naeslundii		47.99	0.38	RAF 92	1	1	
Clostridium tertium		29.40	0.11	XYL 1	1	99	
Atopobium parvulum		22.61	0.10		99	1	
07177122	**Low discrimination**						**07177122**
					COC	SPO	
Clostridium tertium		88.98	0.53	SAL 99	1	99	
Actinomyces israelii		5.41	0.47		1	1	
Atopobium parvulum		4.02	0.31		99	1	
07177130	**Good identification**						**07177130**
					COC	SPO	
Clostridium tertium		98.56	0.50	SAL 99	1	99	
Actinomyces israelii		1.01	0.30		1	1	
Atopobium parvulum		0.43	0.10		99	1	
07177132	**Very good identification**						**07177132**
					SPO		
Clostridium tertium		99.49	0.58	SAL 99	99		
Actinomyces israelii		0.51	0.33		1		
07177140	**Good identification**						**07177140**
					COC	SPO	
Actinomyces israelii		93.76	0.57	SAL 92	1	1	
Atopobium parvulum		5.57	0.23		99	1	
Clostridium tertium		0.43	0.03		1	99	
07177142	**Very good identification**						**07177142**
					SPO		
Actinomyces israelii		96.60	0.60	SAL 92	1		
Bifidobacterium adolescentis		1.47	0.23		1		
Clostridium tertium		0.88	0.11		99		
07177150	**Acceptable identification**						**07177150**
					SPO		
Actinomyces israelii		94.87	0.43	SAL 92	1		
Clostridium tertium		5.13	0.07	MLZ 13	99		
07177152	**Low discrimination**						**07177152**
					SPO		
Actinomyces israelii		81.13	0.46	SAL 92	1		
Bifidobacterium adolescentis		10.10	0.25	MLZ 13	1		
Clostridium tertium		8.77	0.15		99		
07177160	**Low discrimination**						**07177160**
					COC	SPO	
Actinomyces israelii		70.48	0.37	SAL 92	1	1	
Clostridium tertium		15.05	0.11	NAG 6	1	99	
Atopobium parvulum		11.57	0.10		99	1	
07177162	**Low discrimination**						**07177162**
					SPO		
Actinomyces israelii		68.75	0.40	SAL 92	1		
Clostridium tertium		29.37	0.19	NAG 6	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		1.88	0.22		1		
07177202	**Good identification**						**07177202**
Actinomyces israelii		100.0	0.38	SAL 92			
07177300	**Low discrimination**						**07177300**
					COC	SPO	
Actinomyces israelii		75.17	0.45	SAL 92	1	1	
Atopobium parvulum		23.07	0.23	RHA 7	99	1	
Clostridium tertium		1.77	0.03		1	99	
07177302	**Acceptable identification**						**07177302**
					COC	SPO	
Actinomyces israelii		92.33	0.48	SAL 92	1	1	
Clostridium tertium		4.34	0.11	RHA 7	1	99	
Atopobium parvulum		3.33	0.10		99	1	
07177340	**Good identification**						**07177340**
Actinomyces israelii		100.0	0.38	SAL 92			
07177342	**Good identification**						**07177342**
Actinomyces israelii		100.0	0.41	SAL 92			
07177400	**Good identification**						**07177400**
Actinomyces israelii		100.0	0.53	SAL 92			
07177402	**Good identification**						**07177402**
Actinomyces israelii		98.44	0.56	SAL 92			
Bifidobacterium adolescentis		1.56	0.19				
07177410	**Good identification**						**07177410**
Actinomyces israelii		100.0	0.38	SAL 92			
07177412	**Low discrimination**						**07177412**
Actinomyces israelii		88.54	0.42	SAL 92			
Bifidobacterium adolescentis		11.46	0.20	MLZ 13			
07177422	**Good identification**						**07177422**
Actinomyces israelii		100.0	0.35	SAL 92			
07177440	**Good identification**						**07177440**
Actinomyces israelii		100.0	0.46	SAL 92			
07177442	**Good identification**						**07177442**
Actinomyces israelii		100.0	0.49	SAL 92			
07177500	**Good identification**						**07177500**
					COC	SPO	
Actinomyces israelii		94.66	0.63	SAL 92	1	1	
Atopobium parvulum		2.67	0.23		99	1	
Clostridium butyricum		1.90	0.13		1	99	
07177502	**Low discrimination**						**07177502**
					SPO		
Actinomyces israelii		71.20	0.66	SAL 92	1		
Bifidobacterium adolescentis		27.88	0.53		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.37	0.06		99		
07177510	**Acceptable identification**						**07177510**
					SPO		
Actinomyces israelii		93.09	0.49	SAL 92	1		
Bifidobacterium adolescentis		4.52	0.20	MLZ 13	1		
Clostridium tertium		2.39	0.07		99		
07177512	**Low discrimination**						**07177512**
Bifidobacterium adolescentis		74.23	0.54	RAF 92			
Actinomyces israelii		23.18	0.52				
Bifidobacterium dentium		1.21	0.06				
07177520	**Low discrimination**						**07177520**
					COC	SPO	
Actinomyces israelii		82.85	0.42	SAL 92	1	1	
Clostridium tertium		8.41	0.11	NAG 6	1	99	
Atopobium parvulum		6.47	0.10		99	1	
07177522	**Low discrimination**						**07177522**
					SPO		
Actinomyces israelii		79.05	0.45	SAL 92	1		
Clostridium tertium		16.05	0.19	NAG 6	99		
Bifidobacterium adolescentis		4.90	0.19		1		
07177540	**Good identification**						**07177540**
					SPO		
Actinomyces israelii		99.55	0.56	SAL 92	1		
Clostridium butyricum		0.45	-.05		99		
07177542	**Low discrimination**						**07177542**
Actinomyces israelii		88.93	0.59	SAL 92			
Bifidobacterium adolescentis		11.07	0.37				
07177550	**Good identification**						**07177550**
					COC		
Actinomyces israelii		96.33	0.42	SAL 92	1		
Blautia producta		3.67	-.03	MLZ 13	99		
07177552	**Low discrimination**						**07177552**
Bifidobacterium adolescentis		50.44	0.39	RAF 92			
Actinomyces israelii		49.56	0.45	SOR 11			
07177560	**Good identification**						**07177560**
Actinomyces israelii		100.0	0.35	SAL 92			
07177562	**Good identification**						**07177562**
Actinomyces israelii		100.0	0.38	SAL 92			
07177602	**Good identification**						**07177602**
Actinomyces israelii		100.0	0.36	SAL 92			
07177700	**Good identification**						**07177700**
Actinomyces israelii		100.0	0.44	SAL 92			
07177702	**Acceptable identification**						**07177702**
Actinomyces israelii		92.66	0.47	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium adolescentis		4.87	0.19	RHA 7				
Clostridium clostridioforme		2.48	0.10					
07177740	**Good identification**							**07177740**
Actinomyces israelii		100.0	0.37	SAL 92				
07177742	**Good identification**							**07177742**
Actinomyces israelii		100.0	0.40	SAL 92				
07200000	**Low discrimination**							**07200000**
					COC	GRA		
Gemella morbillorum		76.44	0.52	SAL 1	99	99		
Fusobacterium mortiferum		13.07	0.49		1	1		
Atopobium minutum		4.11	0.35		1	99		
07200020	**Low discrimination**							**07200020**
					COC	SPO		
Gemella morbillorum		74.50	0.35	SAL 1	99	1		
Atopobium minutum		12.02	0.26	NAG 10	1	1		
Clostridium cadaveris		10.06	0.15		1	99		
07200100	**Good identification**							**07200100**
					COC	GRA		
Fusobacterium mortiferum		97.53	0.55	FRU 99	1	1		
Gemella morbillorum		2.47	0.17		99	99		
07200102	**Good identification**							**07200102**
Fusobacterium mortiferum		100.0	0.39	FRU 99				
07200400	**Good identification**							**07200400**
					COC	GRA		
Gemella morbillorum		98.47	0.35	SAL 1	99	99		
Fusobacterium mortiferum		1.53	0.14	ARA 10	1	1		
07201000	**Low discrimination**							**07201000**
					COC	GRA		
Gemella morbillorum		53.74	0.47	SAL 1	99	99		
Fusobacterium mortiferum		25.59	0.52		1	1		
Eubacterium tortuosum		16.57	0.32		1	99		
07201100	**Very good identification**							**07201100**
					COC	GRA		
Fusobacterium mortiferum		97.17	0.58	FRU 99	1	1		
Lactobacillus catenaformis		1.21	0.15		1	99		
Gemella morbillorum		0.88	0.13		99	99		
07201102	**Low discrimination**							**07201102**
					GRA			
Fusobacterium mortiferum		84.59	0.42	FRU 99	1			
Lactobacillus catenaformis		9.45	0.15	BGL 10	99			
Lactobacillus acidophilus		2.41	0.02		99			
07203000	**Good identification**							**07203000**
					COC	SPO	CAT	
Clostridium innocuum		97.04	0.44	FRU 95	1	99	1	
Gemella morbillorum		1.07	0.13	CEL 99	99	1	1	
Propionibacterium granulosum		1.06	0.20		1	1	90	
07203002	**Good identification**							**07203002**
					SPO	CAT		
Clostridium innocuum		99.73	0.44	FRU 95	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.27	0.09	CEL 99	1	90		
07204000	**Low discrimination**							**07204000**
					COC	GRA		
Gemella morbillorum		89.62	0.35	SAL 1	99	99		
Fusobacterium mortiferum		10.38	0.29	XYL 10	1	1		
07204100	**Acceptable identification**							**07204100**
					COC	GRA		
Fusobacterium mortiferum		90.83	0.36	FRU 99	1	1		
Actinomyces meyeri		5.78	0.06	XYL 7	1	99		
Gemella morbillorum		3.40	0.01		99	99		
07205100	**Acceptable identification**							**07205100**
					GRA			
Fusobacterium mortiferum		94.02	0.39	FRU 99	1			
Actinomyces meyeri		5.98	0.09	XYL 7	99			
07210000	**Low discrimination**							**07210000**
					COC	GRA	CAT	
Fusobacterium mortiferum		88.50	0.83		1	1	5	
Gemella morbillorum		6.39	0.53		99	99	1	
Propionibacterium granulosum		3.70	0.56		1	99	90	
07210001	**Low discrimination**							**07210001**
					COC	GRA	CAT	
Fusobacterium mortiferum		89.76	0.49	URE 1	1	1	5	
Gemella morbillorum		6.48	0.19		99	99	1	
Propionibacterium granulosum		3.76	0.22		1	99	90	
07210002	**Low discrimination**							**07210002**
					GRA	CAT		
Fusobacterium mortiferum		88.50	0.66	BGL 10	1	5		
Propionibacterium granulosum		8.33	0.46		99	90		
Atopobium minutum		1.56	0.31		99	1		
07210010	**Low discrimination**							**07210010**
					GRA	SPO	CAT	
Fusobacterium mortiferum		39.28	0.49	MLZ 1	1	1	5	
Clostridium difficile		35.22	0.32		99	99	1	
Propionibacterium granulosum		22.19	0.41		99	1	90	
07210020	**Low discrimination**							**07210020**
					COC	GRA	CAT	
Propionibacterium granulosum		33.85	0.46	SAL 1	1	99	90	
Fusobacterium mortiferum		32.67	0.49		1	1	5	
Gemella morbillorum		25.96	0.37		99	99	1	
07210022	**Low discrimination**							**07210022**
					GRA	CAT		
Propionibacterium granulosum		52.65	0.35	SAL 1	99	90		
Fusobacterium mortiferum		22.59	0.32		1	5		
Atopobium minutum		13.12	0.23		99	1		
07210040	**Low discrimination**							**07210040**
					COC	GRA	SPO	
Fusobacterium mortiferum		63.84	0.49	SOR 1	1	1	1	
Clostridium botulinum biovar B		18.68	0.24		1	99	99	
Megasphaera elsdenii		9.58	0.26		99	1	1	
07210100	**Very good identification**							**07210100**
					COC	GRA		
Fusobacterium mortiferum		99.88	0.89		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		0.05	0.23		1	99		
Gemella morbillorum		0.03	0.19		99	99		
07210101	**Good identification**							**07210101**
					GRA			
Fusobacterium mortiferum		99.58	0.55	URE 1	1			
Actinomyces naeslundii		0.42	0.26		99			
07210102	**Very good identification**							**07210102**
					GRA	SPO		
Fusobacterium mortiferum		99.52	0.73	BGL 10	1	1		
Lactobacillus catenaformis		0.17	0.15		99	1		
Clostridium barati		0.16	0.10		99	99		
07210103	**Good identification**							**07210103**
					GRA			
Fusobacterium mortiferum		96.35	0.39	URE 1	1			
Actinomyces naeslundii		3.65	0.26	BGL 10	99			
07210110	**Good identification**							**07210110**
					GRA	SPO		
Fusobacterium mortiferum		99.61	0.55	MLZ 1	1	1		
Clostridium difficile		0.39	-.02		99	99		
07210112	**Good identification**							**07210112**
Fusobacterium mortiferum		100.0	0.39	MLZ 1				
07210120	**Acceptable identification**							**07210120**
					GRA	SPO		
Fusobacterium mortiferum		93.91	0.55	NAG 1	1	1		
Clostridium perfringens		2.36	0.17		99	84		
Clostridium barati		1.69	0.10		99	99		
07210122	**Low discrimination**							**07210122**
					GRA	SPO		
Fusobacterium mortiferum		58.76	0.39	NAG 1	1	1		
Clostridium barati		28.54	0.18	BGL 10	99	99		
Clostridium perfringens		6.25	0.11		99	84		
07210140	**Good identification**							**07210140**
Fusobacterium mortiferum		100.0	0.55	SOR 1				
07210142	**Good identification**							**07210142**
Fusobacterium mortiferum		100.0	0.39	SOR 1				
07210200	**Low discrimination**							**07210200**
					COC	GRA	CAT	
Fusobacterium mortiferum		81.83	0.49	RHA 1	1	1	5	
Propionibacterium granulosum		10.49	0.30		1	99	90	
Gemella morbillorum		5.91	0.19		99	99	1	
07210300	**Good identification**							**07210300**
					GRA			
Fusobacterium mortiferum		99.71	0.55	RHA 1	1			
Actinomyces odontolyticus		0.29	0.06		99			
07210302	**Good identification**							**07210302**
Fusobacterium mortiferum		100.0	0.39	RHA 1				
07210400	**Low discrimination**							**07210400**
					COC	GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		54.25	0.49	ARA 1	1	1	5	
Gemella morbillorum		43.11	0.37		99	99	1	
Propionibacterium granulosum		2.27	0.22		1	99	90	
07210500	**Good identification**							**07210500**
					COC	GRA		
Fusobacterium mortiferum		98.51	0.55	ARA 1	1	1		
Actinomyces meyeri		1.15	0.13		1	99		
Gemella morbillorum		0.34	0.03		99	99		
07210502	**Good identification**							**07210502**
Fusobacterium mortiferum		100.0	0.39	ARA 1				
07211000	**Good identification**							**07211000**
					COC	GRA	CAT	
Fusobacterium mortiferum		90.45	0.86		1	1	5	
Propionibacterium granulosum		2.52	0.56		1	99	90	
Gemella morbillorum		2.35	0.48		99	99	1	
07211001	**Low discrimination**							**07211001**
					GRA	CAT		
Fusobacterium mortiferum		86.78	0.52	URE 1	1	5		
Actinomyces naeslundii		4.80	0.42		99	25		
Propionibacterium granulosum		2.42	0.22		99	90		
07211002	**Low discrimination**							**07211002**
					GRA			
Fusobacterium mortiferum		68.16	0.69	BGL 10	1			
Lactobacillus catenaformis		11.85	0.47		99			
Lactobacillus jensenii		9.79	0.45		99			
07211003	**Low discrimination**							**07211003**
					GRA			
Fusobacterium mortiferum		53.26	0.35	URE 1	1			
Actinomyces naeslundii		26.49	0.42	BGL 10	99			
Lactobacillus catenaformis		9.26	0.12		99			
07211010	**Low discrimination**							**07211010**
					GRA	CAT		
Fusobacterium mortiferum		44.78	0.52	MLZ 1	1	5		
Lactobacillus jensenii		35.37	0.40		99	1		
Propionibacterium granulosum		16.87	0.41		99	90		
07211012	**Low discrimination**							**07211012**
					GRA	CAT		
Lactobacillus jensenii		68.69	0.37	MLT 95	99	1		
Fusobacterium mortiferum		14.49	0.35	CEL 95	1	5		
Propionibacterium granulosum		12.29	0.31		99	90		
07211020	**Low discrimination**							**07211020**
					GRA	CAT		
Fusobacterium mortiferum		25.70	0.52	NAG 1	1	5		
Propionibacterium granulosum		17.75	0.46		99	90		
Lactobacillus jensenii		15.23	0.38		99	1		
07211021	**Low discrimination**							**07211021**
					GRA	CAT		
Actinomyces naeslundii		62.46	0.42	MLT 94	99	25		
Fusobacterium mortiferum		11.42	0.17	RAF 92	1	5		
Propionibacterium granulosum		7.89	0.11		99	90		
07211023	**Low discrimination**							**07211023**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Actinomyces naeslundii		84.42	0.42	MLT 94	1			
Clostridium barati		8.21	-.02	RAF 92	99			
Lactobacillus catenaformis		7.38	0.02		1			
07211040	**Acceptable identification**							**07211040**
					COC	GRA	CAT	
Fusobacterium mortiferum		90.19	0.52	SOR 1	1	1	5	
Propionibacterium granulosum		2.52	0.22		1	99	90	
Gemella morbillorum		2.34	0.14		99	99	1	
07211042	**Low discrimination**							**07211042**
					GRA			
Fusobacterium mortiferum		60.68	0.35	SOR 1	1			
Lactobacillus catenaformis		10.55	0.12	BGL 10	99			
Lactobacillus jensenii		8.71	0.11		99			
07211100	**Very good identification**							**07211100**
					GRA			
Fusobacterium mortiferum		98.34	0.92		1			
Lactobacillus catenaformis		1.22	0.50		99			
Lactobacillus fermentum		0.16	0.46		99			
07211101	**Low discrimination**							**07211101**
					GRA	SPO		
Fusobacterium mortiferum		87.65	0.58	URE 1	1	1		
Actinomyces naeslundii		5.91	0.49		99	1		
Clostridium chauvoei		5.35	0.26		99	99		
07211102	**Low discrimination**							**07211102**
					GRA			
Fusobacterium mortiferum		84.45	0.76	BGL 10	1			
Lactobacillus catenaformis		9.44	0.50		99			
Lactobacillus acidophilus		2.40	0.36		99			
07211103	**Low discrimination**							**07211103**
					GRA			
Fusobacterium mortiferum		53.63	0.42	URE 1	1			
Actinomyces naeslundii		32.53	0.49	BGL 10	99			
Lactobacillus catenaformis		5.99	0.15		99			
07211110	**Very good identification**							**07211110**
					GRA			
Fusobacterium mortiferum		98.21	0.58	MLZ 1	1			
Lactobacillus catenaformis		1.22	0.15		99			
Lactobacillus jensenii		0.34	0.06		99			
07211112	**Low discrimination**							**07211112**
					GRA			
Fusobacterium mortiferum		83.14	0.42	MLZ 1	1			
Lactobacillus catenaformis		9.29	0.15	BGL 10	99			
Lactobacillus acidophilus		2.37	0.02		99			
07211120	**Low discrimination**							**07211120**
					GRA			
Clostridium perfringens		48.17	0.51	MLT 99	99			
Fusobacterium mortiferum		29.07	0.58		1			
Lactobacillus catenaformis		8.94	0.39		99			
07211121	**Low discrimination**							**07211121**
					SPO			
Actinomyces naeslundii		56.17	0.49	MLT 94	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium chauvoei		16.96	0.18	RAF 92	99			
Clostridium perfringens		13.95	0.17		84			
07211122	**Low discrimination**							**07211122**
					SPO			
Clostridium perfringens		32.39	0.45	MLT 99	84			
Clostridium barati		28.39	0.40		99			
Lactobacillus acidophilus		13.00	0.36		1			
07211123	**Low discrimination**							**07211123**
					SPO			
Actinomyces naeslundii		73.23	0.49	MLT 94	1			
Clostridium perfringens		8.56	0.11	RAF 92	84			
Clostridium barati		7.50	0.06		99			
07211140	**Very good identification**							**07211140**
					GRA			
Fusobacterium mortiferum		96.46	0.58	SOR 1	1			
Lactobacillus catenaformis		1.20	0.15		99			
Clostridium perfringens		0.94	0.13		99			
07211141	**Low discrimination**							**07211141**
					GRA			
Actinomyces naeslundii		54.08	0.37	MLT 94	99			
Fusobacterium mortiferum		45.92	0.24	RAF 92	1			
07211142	**Low discrimination**							**07211142**
					GRA	SPO		
Fusobacterium mortiferum		57.11	0.42	SOR 1	1	1		
Lactobacillus acidophilus		19.89	0.21	BGL 10	99	1		
Clostridium barati		10.18	0.14		99	99		
07211143	**Good identification**							**07211143**
Actinomyces naeslundii		100.0	0.37	MLT 94				
07211160	**Low discrimination**							**07211160**
					GRA	SPO		
Clostridium perfringens		88.52	0.39	MLT 99	99	84		
Clostridium barati		5.56	0.14		99	99		
Fusobacterium mortiferum		2.83	0.24		1	1		
07211161	**Low discrimination**							**07211161**
Actinomyces naeslundii		78.86	0.37	MLT 94				
Clostridium perfringens		21.14	0.04	RAF 92				
07211163	**Good identification**							**07211163**
Actinomyces naeslundii		100.0	0.37	MLT 94				
07211200	**Low discrimination**							**07211200**
					COC	GRA	CAT	
Fusobacterium mortiferum		85.51	0.52	RHA 1	1	1	5	
Propionibacterium granulosum		7.31	0.30		1	99	90	
Gemella morbillorum		2.22	0.14		99	99	1	
07211202	**Low discrimination**							**07211202**
					GRA	CAT		
Fusobacterium mortiferum		57.32	0.35	RHA 1	1	5		
Propionibacterium granulosum		11.02	0.20	BGL 10	99	90		
Lactobacillus catenaformis		9.97	0.12		99	1		
07211300	**Very good identification**							**07211300**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Fusobacterium mortiferum		98.13	0.58	RHA 1	1	1		
Lactobacillus catenaformis		1.22	0.15		99	1		
Clostridium barati		0.65	0.05		99	99		
07211302	**Low discrimination**							**07211302**
					GRA	SPO		
Fusobacterium mortiferum		65.83	0.42	RHA 1	1	1		
Lactobacillus acidophilus		13.96	0.17	BGL 10	99	1		
Clostridium barati		11.74	0.14		99	99		
07211400	**Low discrimination**							**07211400**
					COC	GRA		
Fusobacterium mortiferum		60.27	0.52	ARA 1	1	1		
Gemella morbillorum		17.19	0.32		99	99		
Lactobacillus fermentum		13.91	0.42		1	99		
07211402	**Low discrimination**							**07211402**
					GRA	SPO		
Lactobacillus fermentum		51.35	0.42	SAL 1	99	1		
Fusobacterium mortiferum		24.72	0.35		1	1		
Clostridium innocuum		14.53	0.19		99	99		
07211500	**Low discrimination**							**07211500**
					GRA			
Fusobacterium mortiferum		84.35	0.58	ARA 1	1			
Lactobacillus fermentum		13.61	0.46		99			
Lactobacillus catenaformis		1.05	0.15		99			
07211502	**Low discrimination**							**07211502**
					GRA			
Lactobacillus fermentum		54.42	0.46	SAL 1	99			
Fusobacterium mortiferum		37.47	0.42		1			
Lactobacillus catenaformis		4.19	0.15		99			
07211520	**Low discrimination**							**07211520**
					GRA			
Lactobacillus fermentum		62.52	0.38	SAL 1	99			
Clostridium perfringens		19.45	0.17		99			
Fusobacterium mortiferum		11.74	0.24		1			
07211522	**Low discrimination**							**07211522**
					SPO			
Lactobacillus fermentum		69.90	0.38	SAL 1	1			
Clostridium perfringens		10.24	0.11		84			
Clostridium barati		8.97	0.06		99			
07212000	**Low discrimination**							**07212000**
					SPO	CAT		
Propionibacterium granulosum		32.20	0.54	SAL 1	1	90		
Clostridium innocuum		29.79	0.44		99	1		
Clostridium difficile		17.03	0.37		99	1		
07212002	**Low discrimination**							**07212002**
					GRA	SPO	CAT	
Clostridium innocuum		75.44	0.44	CEL 99	99	99	1	
Propionibacterium granulosum		20.38	0.44	GAL 95	99	1	90	
Fusobacterium mortiferum		2.90	0.32		1	1	5	
07212010	**Acceptable identification**							**07212010**
					SPO	CAT		
Clostridium difficile		93.47	0.48	SUC 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		4.94	0.39		1	90		
Clostridium innocuum		1.40	0.20		99	1		
07212020	**Acceptable identification**							**07212020**
					SPO	CAT		
Propionibacterium granulosum		91.62	0.44	SAL 1	1	90		
Clostridium innocuum		3.43	0.10		99	1		
Clostridium difficile		1.96	0.02		99	1		
07212100	**Good identification**							**07212100**
					GRA	SPO	CAT	
Fusobacterium mortiferum		95.74	0.55	MAN 1	1	1	5	
Propionibacterium granulosum		1.30	0.20		99	1	90	
Clostridium innocuum		1.20	0.10		99	99	1	
07212102	**Low discrimination**							**07212102**
					GRA	SPO		
Fusobacterium mortiferum		81.43	0.39	MAN 1	1	1		
Clostridium innocuum		9.18	0.10	BGL 10	99	99		
Actinomyces naeslundii		3.99	0.28		99	1		
07213000	**Good identification**							**07213000**
					GRA	SPO	CAT	
Clostridium innocuum		91.86	0.66	CEL 99	99	99	1	
Propionibacterium granulosum		5.23	0.54		99	1	90	
Fusobacterium mortiferum		2.51	0.52		1	1	5	
07213002	**Very good identification**							**07213002**
					GRA	SPO	CAT	
Clostridium innocuum		97.98	0.66	CEL 99	99	99	1	
Propionibacterium granulosum		1.39	0.44		99	1	90	
Fusobacterium mortiferum		0.30	0.35		1	1	5	
07213010	**Low discrimination**							**07213010**
					SPO	CAT		
Clostridium innocuum		81.06	0.42	CEL 99	99	1		
Propionibacterium granulosum		15.09	0.39	MLZ 4	1	90		
Clostridium difficile		2.89	0.14		99	1		
07213012	**Good identification**							**07213012**
					SPO	CAT		
Clostridium innocuum		95.24	0.42	CEL 99	99	1		
Propionibacterium granulosum		4.43	0.29	MLZ 4	1	90		
Lactobacillus jensenii		0.33	0.03		1	1		
07213020	**Low discrimination**							**07213020**
					SPO	CAT		
Propionibacterium granulosum		51.88	0.44	SAL 1	1	90		
Clostridium innocuum		36.85	0.32		99	1		
Actinomyces naeslundii		7.12	0.44		1	25		
07213021	**Good identification**							**07213021**
Actinomyces naeslundii		95.32	0.47	MLT 94				
Propionibacterium granulosum		4.68	0.09	RAF 92				
07213023	**Good identification**							**07213023**
Actinomyces naeslundii		100.0	0.47	MLT 94				
07213100	**Low discrimination**							**07213100**
					GRA	SPO		
Fusobacterium mortiferum		78.39	0.58	MAN 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		12.43	0.32		99	99	
Actinomyces naeslundii		6.84	0.51		99	1	
07213101	**Acceptable identification**						**07213101**
					GRA		
Actinomyces naeslundii		92.83	0.54	MLT 94	99		
Fusobacterium mortiferum		7.17	0.24	RAF 92	1		
07213103	**Good identification**						**07213103**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07213120	**Low discrimination**						**07213120**
					SPO		
Actinomyces naeslundii		62.65	0.51	MLT 94	1		
Clostridium barati		14.23	0.14	RAF 92	99		
Clostridium perfringens		12.02	0.17		84		
07213121	**Good identification**						**07213121**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07213122	**Low discrimination**						**07213122**
					SPO		
Actinomyces naeslundii		49.46	0.51	MLT 94	1		
Clostridium barati		33.71	0.22	RAF 92	99		
Lactobacillus acidophilus		9.34	0.15		1		
07213123	**Good identification**						**07213123**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07213140	**Low discrimination**						**07213140**
					GRA		
Actinomyces naeslundii		56.45	0.39	MLT 94	99		
Fusobacterium mortiferum		37.05	0.24	RAF 92	1		
Propionibacterium propionicum		6.50	0.12		99		
07213141	**Good identification**						**07213141**
Actinomyces naeslundii		100.0	0.42	MLT 94			
07213142	**Low discrimination**						**07213142**
Actinomyces naeslundii		76.62	0.39	MLT 94			
Propionibacterium propionicum		13.24	0.15	RAF 92			
Lactobacillus acidophilus		10.13	-.01				
07213143	**Good identification**						**07213143**
Actinomyces naeslundii		100.0	0.42	MLT 94			
07213160	**Low discrimination**						**07213160**
Actinomyces naeslundii		82.84	0.39	MLT 94			
Clostridium perfringens		17.16	0.04	RAF 92			
07213161	**Good identification**						**07213161**
Actinomyces naeslundii		100.0	0.42	MLT 94			
07213162	**Low discrimination**						**07213162**
					SPO		
Actinomyces naeslundii		68.12	0.39	MLT 94	1		
Clostridium barati		22.87	0.04	RAF 92	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus acidophilus		9.01	-.01		1			
07213163	**Good identification**							**07213163**
Actinomyces naeslundii		100.0	0.42	MLT 94				
07213400	**Good identification**							**07213400**
					GRA	SPO	CAT	
Clostridium innocuum		99.52	0.53	CEL 99	99	99	1	
Propionibacterium granulosum		0.32	0.20		99	1	90	
Fusobacterium mortiferum		0.16	0.17		1	1	5	
07213402	**Good identification**							**07213402**
					SPO	CAT		
Clostridium innocuum		99.92	0.53	CEL 99	99	1		
Propionibacterium granulosum		0.08	0.09		1	90		
07214000	**Low discrimination**							**07214000**
					COC	GRA		
Fusobacterium mortiferum		89.01	0.64	XYL 7	1	1		
Gemella morbillorum		9.49	0.37		99	99		
Actinomyces meyeri		0.63	0.18		1	99		
07214002	**Good identification**							**07214002**
					COC	GRA	CAT	
Fusobacterium mortiferum		97.82	0.47	XYL 7	1	1	5	
Propionibacterium granulosum		1.24	0.11	BGL 10	1	99	90	
Gemella morbillorum		0.95	0.03		99	99	1	
07214100	**Very good identification**							**07214100**
					COC	GRA		
Fusobacterium mortiferum		98.75	0.70	XYL 7	1	1		
Actinomyces meyeri		1.21	0.28		1	99		
Gemella morbillorum		0.05	0.03		99	99		
07214101	**Good identification**							**07214101**
Fusobacterium mortiferum		100.0	0.36	XYL 7				
07214102	**Good identification**							**07214102**
					GRA			
Fusobacterium mortiferum		99.66	0.54	XYL 7	1			
Actinomyces meyeri		0.34	0.02	BGL 10	99			
07214110	**Good identification**							**07214110**
Fusobacterium mortiferum		100.0	0.36	XYL 7				
07214120	**Low discrimination**							**07214120**
					GRA			
Fusobacterium mortiferum		85.86	0.36	XYL 7	1			
Actinomyces meyeri		14.14	0.13	NAG 1	99			
07214140	**Good identification**							**07214140**
Fusobacterium mortiferum		100.0	0.36	XYL 7				
07214300	**Good identification**							**07214300**
Fusobacterium mortiferum		100.0	0.36	XYL 7				
07214500	**Low discrimination**							**07214500**
					GRA			
Fusobacterium mortiferum		76.81	0.36	XYL 7	1			
Actinomyces meyeri		23.19	0.18	ARA 1	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07215000	**Good identification**						**07215000**
					COC	GRA	
Fusobacterium mortiferum		92.96	0.67	XYL 7	1	1	
Gemella morbillorum		3.56	0.32		99	99	
Lactobacillus fermentum		1.69	0.38		1	99	
07215002	**Low discrimination**						**07215002**
					GRA	SPO	
Fusobacterium mortiferum		80.76	0.50	XYL 7	1	1	
Lactobacillus fermentum		13.22	0.38	BGL 10	99	1	
Clostridium innocuum		1.90	0.10		99	99	
07215100	**Very good identification**						**07215100**
					GRA		
Fusobacterium mortiferum		97.12	0.73	XYL 7	1		
Lactobacillus fermentum		1.24	0.42		99		
Actinomyces meyeri		1.19	0.31		99		
07215101	**Good identification**						**07215101**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
07215102	**Low discrimination**						**07215102**
					GRA		
Fusobacterium mortiferum		86.14	0.57	XYL 7	1		
Lactobacillus fermentum		9.86	0.42	BGL 10	99		
Lactobacillus catenaformis		1.29	0.15		99		
07215110	**Good identification**						**07215110**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
07215120	**Low discrimination**						**07215120**
					GRA		
Fusobacterium mortiferum		51.76	0.39	XYL 7	1		
Lactobacillus fermentum		21.73	0.34	NAG 1	99		
Clostridium perfringens		11.51	0.17		99		
07215140	**Good identification**						**07215140**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
07215322	**Acceptable identification**						**07215322**
					GRA		
Parabacteroides distasonis		92.29	0.39	RAF 99	1		
Prevotella oralis		4.41	0.04		1		
Eubacterium contortum		3.30	0.03		99		
07215400	**Low discrimination**						**07215400**
					COC	GRA	
Lactobacillus fermentum		53.00	0.38	SAL 1	1	99	
Fusobacterium mortiferum		29.43	0.32		1	1	
Gemella morbillorum		12.39	0.16		99	99	
07215402	**Acceptable identification**						**07215402**
					GRA		
Lactobacillus fermentum		94.19	0.38	SAL 1	99		
Fusobacterium mortiferum		5.81	0.16		1		
07215500	**Low discrimination**						**07215500**
					GRA		
Lactobacillus fermentum		49.17	0.42	SAL 1	99		
Fusobacterium mortiferum		39.05	0.39		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		11.79	0.21		99			
07215502	**Low discrimination**							**07215502**
					GRA			
Lactobacillus fermentum		65.66	0.42	SAL 1	99			
Bifidobacterium adolescentis		28.55	0.30		99			
Fusobacterium mortiferum		5.79	0.22		1			
07215722	**Good identification**							**07215722**
					GRA			
Parabacteroides distasonis		95.45	0.35	RAF 99	1			
Eubacterium contortum		4.55	0.01		99			
07216100	**Good identification**							**07216100**
Fusobacterium mortiferum		100.0	0.36	MAN 1				
07217000	**Good identification**							**07217000**
					GRA	SPO	CAT	
Clostridium innocuum		95.24	0.44	CEL 99	99	99	1	
Fusobacterium mortiferum		3.72	0.32	XYL 5	1	1	5	
Propionibacterium granulosum		1.04	0.20		99	1	90	
07217002	**Good identification**							**07217002**
					GRA	SPO	CAT	
Clostridium innocuum		99.30	0.44	CEL 99	99	99	1	
Fusobacterium mortiferum		0.43	0.16	XYL 5	1	1	5	
Propionibacterium granulosum		0.27	0.09		99	1	90	
07217100	**Low discrimination**							**07217100**
					GRA	SPO		
Fusobacterium mortiferum		85.87	0.39	MAN 1	1	1		
Clostridium innocuum		9.52	0.10	XYL 7	99	99		
Actinomyces israelii		4.61	0.19		99	1		
07220000	**Low discrimination**							**07220000**
					COC	GRA	CAT	
Fusobacterium mortiferum		56.49	0.40	FRU 99	1	1	5	
Propionibacterium granulosum		23.75	0.30		1	99	90	
Gemella morbillorum		11.18	0.17		99	99	1	
07220100	**Good identification**							**07220100**
Fusobacterium mortiferum		100.0	0.46	FRU 99				
07221000	**Low discrimination**							**07221000**
					GRA	CAT		
Fusobacterium mortiferum		61.11	0.43	FRU 99	1	5		
Propionibacterium granulosum		17.13	0.30		99	90		
Lactobacillus jensenii		14.69	0.22		99	1		
07221100	**Acceptable identification**							**07221100**
					COC	GRA		
Fusobacterium mortiferum		94.66	0.49	FRU 99	1	1		
Atopobium parvulum		3.89	0.14		99	99		
Actinomyces naeslundii		1.46	0.30		1	99		
07223000	**Acceptable identification**							**07223000**
					SPO	CAT		
Clostridium innocuum		90.58	0.36	FRU 95	99	1		
Propionibacterium granulosum		8.44	0.28	CEL 99	1	90		
Actinomyces naeslundii		0.98	0.27		1	25		
07223002	**Good identification**							**07223002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	CAT		
Clostridium innocuum		96.70	0.36	FRU 95	99	1		
Propionibacterium granulosum		2.25	0.18	CEL 99	1	90		
Actinomyces naeslundii		1.04	0.27		1	25		
07223101	**Good identification**							**07223101**
Actinomyces naeslundii		100.0	0.37	MLT 94				
07223103	**Good identification**							**07223103**
Actinomyces naeslundii		100.0	0.37	MLT 94				
07223121	**Good identification**							**07223121**
Actinomyces naeslundii		100.0	0.37	MLT 94				
07223123	**Good identification**							**07223123**
Actinomyces naeslundii		100.0	0.37	MLT 94				
07230000	**Low discrimination**							**07230000**
					GRA	CAT		
Fusobacterium mortiferum		70.05	0.74		1	5		
Propionibacterium granulosum		29.45	0.64		99	90		
Lactobacillus jensenii		0.26	0.22		99	1		
07230001	**Low discrimination**							**07230001**
					GRA	CAT		
Fusobacterium mortiferum		69.31	0.40	URE 1	1	5		
Propionibacterium granulosum		29.14	0.30		99	90		
Actinomyces naeslundii		1.56	0.23		99	25		
07230002	**Low discrimination**							**07230002**
					GRA	CAT		
Fusobacterium mortiferum		50.74	0.57	BGL 10	1	5		
Propionibacterium granulosum		47.99	0.54		99	90		
Lactobacillus jensenii		1.11	0.19		99	1		
07230010	**Low discrimination**							**07230010**
					GRA	SPO	CAT	
Propionibacterium granulosum		80.83	0.49	SAL 1	99	1	90	
Fusobacterium mortiferum		14.24	0.40	MLZ 12	1	1	5	
Clostridium difficile		3.22	0.13		99	99	1	
07230012	**Low discrimination**							**07230012**
					GRA	CAT		
Propionibacterium granulosum		88.12	0.39	SAL 1	99	90		
Fusobacterium mortiferum		6.90	0.23	MLZ 12	1	5		
Lactobacillus jensenii		4.98	0.11		99	1		
07230020	**Low discrimination**							**07230020**
					GRA	CAT		
Propionibacterium granulosum		89.92	0.54	SAL 1	99	90		
Fusobacterium mortiferum		8.64	0.40		1	5		
Lactobacillus jensenii		0.78	0.12		99	1		
07230022	**Acceptable identification**							**07230022**
					GRA	CAT		
Propionibacterium granulosum		93.04	0.43	SAL 1	99	90		
Fusobacterium mortiferum		3.98	0.23		1	5		
Lactobacillus jensenii		2.15	0.09		99	1		
07230030	**Good identification**							**07230030**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					CAT			
Propionibacterium granulosum		97.93	0.39	SAL 1	90			
Lactobacillus jensenii		2.07	0.04	MLZ 12	1			
07230100	**Very good identification**							**07230100**
					COC	GRA	CAT	
Fusobacterium mortiferum		99.74	0.80		1	1	5	
Propionibacterium granulosum		0.18	0.30		1	99	90	
Atopobium parvulum		0.06	0.14		99	99	1	
07230101	**Good identification**							**07230101**
					GRA			
Fusobacterium mortiferum		97.34	0.46	URE 1	1			
Actinomyces naeslundii		2.67	0.31		99			
07230102	**Very good identification**							**07230102**
					GRA	CAT		
Fusobacterium mortiferum		98.86	0.64	BGL 10	1	5		
Lactobacillus acidophilus		0.48	0.11		99	1		
Propionibacterium granulosum		0.41	0.20		99	90		
07230110	**Good identification**							**07230110**
					GRA	CAT		
Fusobacterium mortiferum		97.60	0.46	MLZ 1	1	5		
Propionibacterium granulosum		2.40	0.15		99	90		
07230120	**Low discrimination**							**07230120**
					GRA	CAT		
Fusobacterium mortiferum		80.41	0.46	NAG 1	1	5		
Clostridium perfringens		12.55	0.21		99	1		
Propionibacterium granulosum		3.62	0.20		99	90		
07230140	**Good identification**							**07230140**
					GRA	SPO		
Fusobacterium mortiferum		98.74	0.46	SOR 1	1	1		
Clostridium botulinum biovar B		1.26	-.03		99	99		
07230200	**Low discrimination**							**07230200**
					GRA	CAT		
Propionibacterium granulosum		56.28	0.38	SAL 1	99	90		
Fusobacterium mortiferum		43.72	0.40	RHA 3	1	5		
07230300	**Good identification**							**07230300**
Fusobacterium mortiferum		100.0	0.46	RHA 1				
07230400	**Low discrimination**							**07230400**
					COC	GRA	CAT	
Fusobacterium mortiferum		69.10	0.40	ARA 1	1	1	5	
Propionibacterium granulosum		29.05	0.30		1	99	90	
Gemella morbillorum		1.86	0.03		99	99	1	
07230500	**Good identification**							**07230500**
Fusobacterium mortiferum		100.0	0.46	ARA 1				
07231000	**Low discrimination**							**07231000**
					GRA	CAT		
Fusobacterium mortiferum		65.31	0.77		1	5		
Propionibacterium granulosum		18.30	0.64		99	90		
Lactobacillus jensenii		15.70	0.56		99	1		
07231001	**Low discrimination**							**07231001**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		53.20	0.43	URE 1	1	5	
Actinomyces naeslundii		19.10	0.47		99	25	
Propionibacterium granulosum		14.91	0.30		99	90	
07231002	**Low discrimination**						**07231002**
					GRA	CAT	
Lactobacillus jensenii		44.86	0.53	MLT 95	99	1	
Fusobacterium mortiferum		31.10	0.60	CEL 95	1	5	
Propionibacterium granulosum		19.61	0.54		99	90	
07231003	**Low discrimination**						**07231003**
					GRA		
Actinomyces naeslundii		51.26	0.47	MLT 94	99		
Lactobacillus jensenii		22.88	0.19	RAF 92	99		
Fusobacterium mortiferum		15.86	0.26		1		
07231010	**Low discrimination**						**07231010**
					GRA	CAT	
Lactobacillus jensenii		62.30	0.48	MLT 95	99	1	
Propionibacterium granulosum		29.71	0.49	CEL 95	99	90	
Fusobacterium mortiferum		7.85	0.43		1	5	
07231012	**Low discrimination**						**07231012**
					GRA	CAT	
Lactobacillus jensenii		83.11	0.45	MLT 95	99	1	
Propionibacterium granulosum		14.86	0.39	CEL 95	99	90	
Fusobacterium mortiferum		1.75	0.26		1	5	
07231020	**Low discrimination**						**07231020**
					CAT		
Propionibacterium granulosum		39.62	0.54	SAL 1	90		
Lactobacillus jensenii		33.98	0.46		1		
Clostridium perfringens		18.72	0.41		1		
07231021	**Low discrimination**						**07231021**
					CAT		
Actinomyces naeslundii		68.74	0.47	MLT 94	25		
Propionibacterium granulosum		13.41	0.20	RAF 92	90		
Lactobacillus jensenii		11.50	0.12		1		
07231022	**Low discrimination**						**07231022**
					CAT		
Lactobacillus jensenii		46.80	0.43	MLT 95	1		
Propionibacterium granulosum		20.46	0.43	CEL 95	90		
Clostridium perfringens		18.20	0.35		1		
07231023	**Low discrimination**						**07231023**
					CAT		
Actinomyces naeslundii		83.08	0.47	MLT 94	25		
Lactobacillus jensenii		9.27	0.09	RAF 92	1		
Propionibacterium granulosum		4.05	0.09		90		
07231030	**Low discrimination**						**07231030**
					CAT		
Lactobacillus jensenii		66.95	0.38	MLT 95	1		
Propionibacterium granulosum		31.93	0.39	CEL 95	90		
Clostridium perfringens		1.12	0.07		1		
07231040	**Low discrimination**						**07231040**
					GRA	CAT	
Fusobacterium mortiferum		62.42	0.43	SOR 1	1	5	
Propionibacterium granulosum		17.49	0.30		99	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		15.01	0.22		99	1	
07231100	**Very good identification**						**07231100**
					COC	GRA	
Fusobacterium mortiferum		95.18	0.83		1	1	
Atopobium parvulum		3.91	0.48		99	99	
Clostridium perfringens		0.31	0.30		1	99	
07231101	**Low discrimination**						**07231101**
					COC	GRA	
Fusobacterium mortiferum		66.59	0.49	URE 1	1	1	
Actinomyces naeslundii		29.17	0.54		1	99	
Atopobium parvulum		2.74	0.14		99	99	
07231102	**Low discrimination**						**07231102**
					COC	GRA	
Fusobacterium mortiferum		69.29	0.67	BGL 10	1	1	
Lactobacillus acidophilus		22.10	0.45		1	99	
Atopobium parvulum		4.52	0.35		99	99	
07231103	**Low discrimination**						**07231103**
					GRA		
Actinomyces naeslundii		74.01	0.54	MLT 94	99		
Fusobacterium mortiferum		18.78	0.33	RAF 92	1		
Lactobacillus acidophilus		5.99	0.11		99		
07231110	**Acceptable identification**						**07231110**
					COC	GRA	
Fusobacterium mortiferum		90.32	0.49	MLZ 1	1	1	
Atopobium parvulum		3.71	0.14		99	99	
Lactobacillus jensenii		3.10	0.14		1	99	
07231120	**Low discrimination**						**07231120**
					GRA	SPO	
Clostridium perfringens		77.11	0.56	MLT 99	99	84	
Fusobacterium mortiferum		7.49	0.49		1	1	
Clostridium septicum		6.59	0.31		99	99	
07231121	**Low discrimination**						**07231121**
					GRA		
Actinomyces naeslundii		76.06	0.54	MLT 94	99		
Clostridium perfringens		18.07	0.21	RAF 92	99		
Fusobacterium mortiferum		1.75	0.15		1		
07231122	**Low discrimination**						**07231122**
Clostridium perfringens		52.56	0.50	MLT 99			
Lactobacillus acidophilus		38.04	0.45				
Actinomyces naeslundii		3.17	0.51				
07231123	**Low discrimination**						**07231123**
Actinomyces naeslundii		83.84	0.54	MLT 94			
Clostridium perfringens		9.37	0.16	RAF 92			
Lactobacillus acidophilus		6.78	0.11				
07231140	**Low discrimination**						**07231140**
					GRA		
Fusobacterium mortiferum		86.71	0.49	SOR 1	1		
Clostridium perfringens		5.26	0.17		99		
Actinomyces naeslundii		4.47	0.39		99		
07231141	**Low discrimination**						**07231141**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Actinomyces naeslundii		88.44	0.42	MLT 94	99			
Fusobacterium mortiferum		11.56	0.15	RAF 92	1			
07231143	**Good identification**							**07231143**
Actinomyces naeslundii		100.0	0.42	MLT 94				
07231160	**Good identification**							**07231160**
					GRA			
Clostridium perfringens		96.19	0.43	MLT 99	99			
Actinomyces naeslundii		2.53	0.39		99			
Fusobacterium mortiferum		0.50	0.15		1			
07231161	**Low discrimination**							**07231161**
Actinomyces naeslundii		79.59	0.42	MLT 94				
Clostridium perfringens		20.41	0.09	RAF 92				
07231162	**Low discrimination**							**07231162**
Clostridium perfringens		64.76	0.38	MLT 99				
Lactobacillus acidophilus		30.41	0.30					
Actinomyces naeslundii		3.61	0.39					
07231163	**Low discrimination**							**07231163**
Actinomyces naeslundii		89.23	0.42	MLT 94				
Clostridium perfringens		10.77	0.03	RAF 92				
07231200	**Low discrimination**							**07231200**
					GRA	CAT		
Fusobacterium mortiferum		47.65	0.43	RHA 1	1	5		
Propionibacterium granulosum		40.89	0.38		99	90		
Lactobacillus jensenii		11.46	0.22		99	1		
07231300	**Good identification**							**07231300**
					COC	GRA		
Fusobacterium mortiferum		96.05	0.49	RHA 1	1	1		
Atopobium parvulum		3.95	0.14		99	99		
07231400	**Low discrimination**							**07231400**
					GRA	CAT		
Fusobacterium mortiferum		62.60	0.43	ARA 1	1	5		
Propionibacterium granulosum		17.54	0.30		99	90		
Lactobacillus jensenii		15.05	0.22		99	1		
07231500	**Good identification**							**07231500**
					COC	GRA		
Fusobacterium mortiferum		96.05	0.49	ARA 1	1	1		
Atopobium parvulum		3.95	0.14		99	99		
07232000	**Low discrimination**							**07232000**
					GRA	SPO	CAT	
Propionibacterium granulosum		86.56	0.62	SAL 1	99	1	90	
Clostridium innocuum		9.38	0.36		99	99	1	
Fusobacterium mortiferum		2.76	0.40		1	1	5	
07232002	**Low discrimination**							**07232002**
					GRA	SPO	CAT	
Propionibacterium granulosum		68.90	0.52	SAL 1	99	1	90	
Clostridium innocuum		29.87	0.36		99	99	1	
Fusobacterium mortiferum		0.98	0.23		1	1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07232010	**Low discrimination**							**07232010**
					SPO	CAT		
Propionibacterium granulosum		66.31	0.47	SAL 1	1	90		
Clostridium difficile		31.50	0.29	MLZ 12	99	1		
Clostridium innocuum		2.20	0.13		99	1		
07232012	**Low discrimination**							**07232012**
					SPO	CAT		
Propionibacterium granulosum		86.83	0.37	SAL 1	1	90		
Clostridium innocuum		11.50	0.13	MLZ 12	99	1		
Clostridium difficile		1.67	-.05		99	1		
07232020	**Good identification**							**07232020**
					SPO	CAT		
Propionibacterium granulosum		99.20	0.52	SAL 1	1	90		
Clostridium innocuum		0.43	0.02		99	1		
Actinomyces naeslundii		0.37	0.25		1	25		
07232022	**Good identification**							**07232022**
					SPO	CAT		
Propionibacterium granulosum		96.87	0.41	SAL 1	1	90		
Clostridium innocuum		1.70	0.02		99	1		
Actinomyces naeslundii		1.43	0.25		1	25		
07232030	**Good identification**							**07232030**
					SPO	CAT		
Propionibacterium granulosum		98.12	0.37	SAL 1	1	90		
Clostridium difficile		1.88	-.05	MLZ 12	99	1		
07232100	**Low discrimination**							**07232100**
					GRA	CAT		
Fusobacterium mortiferum		84.31	0.46	MAN 1	1	5		
Propionibacterium granulosum		11.46	0.28		99	90		
Actinomyces naeslundii		2.99	0.33		99	25		
07232101	**Low discrimination**							**07232101**
					GRA			
Actinomyces naeslundii		84.03	0.36	MLT 94	99			
Fusobacterium mortiferum		15.97	0.12	RAF 92	1			
07232103	**Good identification**							**07232103**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07232121	**Good identification**							**07232121**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07232123	**Good identification**							**07232123**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07232200	**Good identification**							**07232200**
					SPO	CAT		
Propionibacterium granulosum		96.58	0.36	SAL 1	1	90		
Clostridium innocuum		3.42	0.02	RHA 3	99	1		
07233000	**Low discrimination**							**07233000**
					GRA	SPO	CAT	
Clostridium innocuum		65.57	0.58	CEL 99	99	99	1	
Propionibacterium granulosum		31.84	0.62		99	1	90	
Fusobacterium mortiferum		1.52	0.43		1	1	5	
07233001	**Low discrimination**							**07233001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO	CAT	
Actinomyces naeslundii		51.87	0.52	MLT 94	1	25	
Clostridium innocuum		32.40	0.24	RAF 92	99	1	
Propionibacterium granulosum		15.73	0.28		1	90	
07233002	**Low discrimination**						**07233002**
					SPO	CAT	
Clostridium innocuum		87.77	0.58	CEL 99	99	1	
Propionibacterium granulosum		10.66	0.52		1	90	
Actinomyces naeslundii		0.95	0.49		1	25	
07233003	**Low discrimination**						**07233003**
					SPO	CAT	
Actinomyces naeslundii		58.81	0.52	MLT 94	1	25	
Clostridium innocuum		36.73	0.24	RAF 92	99	1	
Propionibacterium granulosum		4.46	0.18		1	90	
07233010	**Low discrimination**						**07233010**
					SPO	CAT	
Propionibacterium granulosum		59.86	0.47	SAL 1	1	90	
Clostridium innocuum		37.66	0.34	MLZ 12	99	1	
Lactobacillus jensenii		1.68	0.14		1	1	
07233020	**Low discrimination**						**07233020**
					SPO	CAT	
Propionibacterium granulosum		84.05	0.52	SAL 1	1	90	
Actinomyces naeslundii		7.46	0.49		1	25	
Clostridium innocuum		6.99	0.24		99	1	
07233021	**Acceptable identification**						**07233021**
Actinomyces naeslundii		92.95	0.52	MLT 94			
Propionibacterium granulosum		7.05	0.18	RAF 92			
07233022	**Low discrimination**						**07233022**
					SPO	CAT	
Propionibacterium granulosum		56.84	0.41	SAL 1	1	90	
Actinomyces naeslundii		20.19	0.49		1	25	
Clostridium innocuum		18.92	0.24		99	1	
07233023	**Good identification**						**07233023**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07233030	**Acceptable identification**						**07233030**
					CAT		
Propionibacterium granulosum		91.93	0.37	SAL 1	90		
Actinomyces naeslundii		3.15	0.27	MLZ 12	25		
Lactobacillus jensenii		2.58	0.04		1		
07233041	**Good identification**						**07233041**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07233043	**Good identification**						**07233043**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07233060	**Low discrimination**						**07233060**
Actinomyces naeslundii		60.81	0.36	MLT 94			
Propionibacterium granulosum		39.19	0.18	RAF 92			
07233061	**Good identification**						**07233061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	MLT 94			
07233062	**Good identification**						**07233062**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07233063	**Good identification**						**07233063**
Actinomyces naeslundii		100.0	0.39	MLT 94			
07233100	**Low discrimination**						**07233100**
					GRA	SPO	
Fusobacterium mortiferum		52.84	0.49	MAN 1	1	1	
Actinomyces naeslundii		29.95	0.56		99	1	
Clostridium innocuum		9.86	0.24		99	99	
07233101	**Very good identification**						**07233101**
					GRA		
Actinomyces naeslundii		98.83	0.59	MLT 94	99		
Fusobacterium mortiferum		1.17	0.15	RAF 92	1		
07233102	**Low discrimination**						**07233102**
					SPO		
Actinomyces naeslundii		50.82	0.56	MLT 94	1		
Clostridium innocuum		16.73	0.24	RAF 92	99		
Lactobacillus acidophilus		16.56	0.24		1		
07233103	**Very good identification**						**07233103**
Actinomyces naeslundii		100.0	0.59	MLT 94			
07233111	**Good identification**						**07233111**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07233113	**Good identification**						**07233113**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07233120	**Low discrimination**						**07233120**
					CAT		
Actinomyces naeslundii		74.28	0.56	MLT 94	25		
Clostridium perfringens		13.64	0.21	RAF 92	1		
Propionibacterium granulosum		2.97	0.18		90		
07233121	**Very good identification**						**07233121**
Actinomyces naeslundii		100.0	0.59	MLT 94			
07233122	**Low discrimination**						**07233122**
					SPO		
Actinomyces naeslundii		58.51	0.56	MLT 94	1		
Lactobacillus acidophilus		19.06	0.24	RAF 92	1		
Clostridium ramosum		8.69	0.17		92		
07233123	**Very good identification**						**07233123**
Actinomyces naeslundii		100.0	0.59	MLT 94			
07233131	**Good identification**						**07233131**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07233133	**Good identification**						**07233133**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		100.0	0.37	MLT 94				
07233140	**Acceptable identification**							**07233140**
					GRA			
Actinomyces naeslundii		90.83	0.43	MLT 94	99			
Fusobacterium mortiferum		9.17	0.15	RAF 92	1			
07233141	**Good identification**							**07233141**
Actinomyces naeslundii		100.0	0.46	MLT 94				
07233142	**Low discrimination**							**07233142**
Actinomyces naeslundii		76.87	0.43	MLT 94				
Lactobacillus acidophilus		17.54	0.08	RAF 92				
Propionibacterium propionicum		5.60	0.14					
07233143	**Good identification**							**07233143**
Actinomyces naeslundii		100.0	0.46	MLT 94				
07233160	**Low discrimination**							**07233160**
Actinomyces naeslundii		83.46	0.43	MLT 94				
Clostridium perfringens		16.54	0.09	RAF 92				
07233161	**Good identification**							**07233161**
Actinomyces naeslundii		100.0	0.46	MLT 94				
07233162	**Low discrimination**							**07233162**
Actinomyces naeslundii		75.68	0.43	MLT 94				
Lactobacillus acidophilus		17.27	0.08	RAF 92				
Clostridium perfringens		7.06	0.03					
07233163	**Good identification**							**07233163**
Actinomyces naeslundii		100.0	0.46	MLT 94				
07233200	**Low discrimination**							**07233200**
					SPO	CAT		
Propionibacterium granulosum		59.79	0.36	SAL 1	1	90		
Clostridium innocuum		40.21	0.24	RHA 3	99	1		
07233400	**Good identification**							**07233400**
					SPO	CAT		
Clostridium innocuum		97.30	0.45	CEL 99	99	1		
Propionibacterium granulosum		2.71	0.28		1	90		
07233402	**Good identification**							**07233402**
					SPO	CAT		
Clostridium innocuum		99.31	0.45	CEL 99	99	1		
Propionibacterium granulosum		0.69	0.18		1	90		
07234000	**Acceptable identification**							**07234000**
					COC	GRA	CAT	
Fusobacterium mortiferum		94.34	0.55	XYL 7	1	1	5	
Propionibacterium granulosum		5.32	0.30		1	99	90	
Gemella morbillorum		0.34	0.03		99	99	1	
07234002	**Low discrimination**							**07234002**
					GRA	CAT		
Fusobacterium mortiferum		88.74	0.38	XYL 7	1	5		
Propionibacterium granulosum		11.26	0.20	BGL 10	99	90		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07234100	**Very good identification**						**07234100**
					COC	GRA	
Fusobacterium mortiferum		98.49	0.61	XYL 7	1	1	
Atopobium parvulum		1.51	0.18		99	99	
07234102	**Good identification**						**07234102**
					COC	GRA	
Fusobacterium mortiferum		97.62	0.45	XYL 7	1	1	
Atopobium parvulum		2.38	0.06	BGL 10	99	99	
07235000	**Good identification**						**07235000**
					GRA	CAT	
Fusobacterium mortiferum		91.17	0.58	XYL 7	1	5	
Propionibacterium granulosum		3.43	0.30		99	90	
Lactobacillus jensenii		2.94	0.22		99	1	
07235002	**Low discrimination**						**07235002**
					GRA	CAT	
Fusobacterium mortiferum		74.42	0.41	XYL 7	1	5	
Lactobacillus jensenii		14.40	0.19	BGL 10	99	1	
Propionibacterium granulosum		6.30	0.20		99	90	
07235100	**Low discrimination**						**07235100**
					COC	GRA	
Atopobium parvulum		50.31	0.53	MLT 99	99	99	
Fusobacterium mortiferum		49.63	0.64		1	1	
Actinomyces israelii		0.06	0.16		1	99	
07235102	**Low discrimination**						**07235102**
					COC	GRA	
Atopobium parvulum		59.95	0.40	MLT 99	99	99	
Fusobacterium mortiferum		37.24	0.48		1	1	
Lactobacillus acidophilus		1.59	0.11		1	99	
07235120	**Low discrimination**						**07235120**
					COC	GRA	
Atopobium parvulum		87.63	0.40	MLT 99	99	99	
Clostridium perfringens		6.84	0.21		1	99	
Fusobacterium mortiferum		4.95	0.30		1	1	
07235122	**Low discrimination**						**07235122**
					COC	GRA	
Parabacteroides distasonis		59.09	0.42	RAF 99	1	1	
Atopobium parvulum		29.34	0.27		99	99	
Clostridium perfringens		6.11	0.16		1	99	
07235132	**Good identification**						**07235132**
Parabacteroides distasonis		100.0	0.37	RAF 99			
07235322	**Good identification**						**07235322**
Parabacteroides distasonis		100.0	0.50	RAF 99			
07235332	**Good identification**						**07235332**
Parabacteroides distasonis		98.82	0.46	RAF 99			
Clostridium clostridioforme		1.18	0.07				
07235522	**Good identification**						**07235522**
Parabacteroides distasonis		100.0	0.38	RAF 99			
07235722	**Good identification**						**07235722**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		96.52	0.47	RAF 99				
Clostridium clostridioforme		3.49	0.16					
07235732	**Low discrimination**							**07235732**
Parabacteroides distasonis		76.60	0.42	RAF 99				
Clostridium clostridioforme		23.40	0.27					
07237000	**Low discrimination**							**07237000**
					GRA	SPO	CAT	
Clostridium innocuum		87.76	0.36	CEL 99	99	99	1	
Propionibacterium granulosum		8.18	0.28	XYL 5	99	1	90	
Fusobacterium mortiferum		2.91	0.23		1	1	5	
07237002	**Good identification**							**07237002**
					SPO	CAT		
Clostridium innocuum		95.89	0.36	CEL 99	99	1		
Propionibacterium granulosum		2.23	0.18	XYL 5	1	90		
Actinomyces israelii		1.88	0.22		1	1		
07240000	**Low discrimination**							**07240000**
					COC	GRA	SPO	
Fusobacterium mortiferum		57.42	0.40	FRU 99	1	1	1	
Clostridium innocuum		26.22	0.22		1	99	99	
Gemella morbillorum		10.18	0.17		99	99	1	
07240100	**Acceptable identification**							**07240100**
					GRA	SPO		
Fusobacterium mortiferum		93.12	0.47	FRU 99	1	1		
Lactobacillus catenaformis		5.20	0.15		99	1		
Clostridium barati		1.68	0.02		99	99		
07241000	**Low discrimination**							**07241000**
					SPO			
Clostridium innocuum		40.33	0.44	FRU 95	99			
Lactobacillus catenaformis		40.00	0.47	MAN 99	1			
Lactobacillus jensenii		9.51	0.36		1			
07241002	**Low discrimination**							**07241002**
					SPO			
Clostridium innocuum		43.49	0.44	FRU 95	99			
Lactobacillus catenaformis		43.13	0.47	MAN 99	1			
Lactobacillus jensenii		6.84	0.33		1			
07241020	**Low discrimination**							**07241020**
					SPO			
Lactobacillus catenaformis		59.67	0.36	MLT 90	1			
Clostridium barati		22.12	0.24	FRU 99	99			
Lactobacillus jensenii		14.19	0.26		1			
07241100	**Low discrimination**							**07241100**
					GRA	SPO		
Lactobacillus catenaformis		73.90	0.50	MLT 90	99	1		
Fusobacterium mortiferum		20.04	0.50	FRU 99	1	1		
Clostridium barati		4.57	0.24		99	99		
07241102	**Low discrimination**							**07241102**
					SPO			
Lactobacillus catenaformis		67.59	0.50	MLT 90	1			
Lactobacillus acidophilus		17.21	0.36	FRU 99	1			
Clostridium barati		12.53	0.32		99			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07241120	**Low discrimination**					**07241120**
					SPO	
Lactobacillus catenaformis		55.65	0.39	MLT 90	1	
Clostridium barati		41.26	0.32	FRU 99	99	
Clostridium paraputrificum		0.83	-.03		99	
07241122	**Low discrimination**					**07241122**
					SPO	
Clostridium barati		52.10	0.40	ESL 87	99	
Lactobacillus acidophilus		23.86	0.36	FRU 99	1	
Lactobacillus catenaformis		23.43	0.39		1	
07242000	**Very good identification**					**07242000**
Clostridium innocuum		100.0	0.56	FRU 95		
07242002	**Very good identification**					**07242002**
Clostridium innocuum		100.0	0.56	FRU 95		
07242400	**Good identification**					**07242400**
Clostridium innocuum		100.0	0.43	FRU 95		
07242402	**Good identification**					**07242402**
Clostridium innocuum		100.0	0.43	FRU 95		
07243000	**Very good identification**					**07243000**
					SPO	
Clostridium innocuum		99.98	0.78	FRU 95	99	
Lactobacillus catenaformis		0.01	0.12		1	
Actinomyces naeslundii		0.00	0.24		1	
07243001	**Good identification**					**07243001**
					SPO	
Clostridium innocuum		99.61	0.44	FRU 95	99	
Actinomyces naeslundii		0.39	0.27	URE 1	1	
07243002	**Very good identification**					**07243002**
					SPO	
Clostridium innocuum		99.98	0.78	FRU 95	99	
Lactobacillus catenaformis		0.01	0.12		1	
Clostridium barati		0.01	0.05		99	
07243003	**Good identification**					**07243003**
					SPO	
Clostridium innocuum		99.61	0.44	FRU 95	99	
Actinomyces naeslundii		0.39	0.27	URE 1	1	
07243010	**Good identification**					**07243010**
Clostridium innocuum		100.0	0.55	FRU 95		
07243012	**Good identification**					**07243012**
Clostridium innocuum		100.0	0.55	FRU 95		
07243020	**Good identification**					**07243020**
					SPO	
Clostridium innocuum		98.71	0.44	FRU 95	99	
Clostridium barati		0.79	0.05	NAG 1	99	
Actinomyces naeslundii		0.26	0.24		1	
07243022	**Good identification**					**07243022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium innocuum		97.18	0.44	FRU 95	99	
Clostridium barati		2.33	0.14	NAG 1	99	
Actinomyces naeslundii		0.25	0.24		1	
07243040	**Good identification**					**07243040**
Clostridium innocuum		100.0	0.44	FRU 95		
07243042	**Good identification**					**07243042**
Clostridium innocuum		100.0	0.44	FRU 95		
07243100	**Good identification**					**07243100**
					SPO	
Clostridium innocuum		96.67	0.44	FRU 95	99	
Lactobacillus catenaformis		1.45	0.15	LAC 1	1	
Clostridium barati		0.77	0.05		99	
07243102	**Acceptable identification**					**07243102**
					SPO	
Clostridium innocuum		93.78	0.44	FRU 95	99	
Clostridium barati		2.25	0.14	LAC 1	99	
Lactobacillus acidophilus		1.87	0.15		1	
07243200	**Good identification**					**07243200**
Clostridium innocuum		100.0	0.44	FRU 95		
07243202	**Good identification**					**07243202**
Clostridium innocuum		100.0	0.44	FRU 95		
07243400	**Very good identification**					**07243400**
Clostridium innocuum		100.0	0.65	FRU 95		
07243402	**Very good identification**					**07243402**
Clostridium innocuum		100.0	0.65	FRU 95		
07243410	**Good identification**					**07243410**
Clostridium innocuum		100.0	0.42	FRU 95		
07243412	**Good identification**					**07243412**
Clostridium innocuum		100.0	0.42	FRU 95		
07247000	**Very good identification**					**07247000**
Clostridium innocuum		100.0	0.56	FRU 95		
07247002	**Very good identification**					**07247002**
Clostridium innocuum		100.0	0.56	FRU 95		
07247400	**Good identification**					**07247400**
Clostridium innocuum		100.0	0.43	FRU 95		
07247402	**Good identification**					**07247402**
Clostridium innocuum		100.0	0.43	FRU 95		
07250000	**Low discrimination**					**07250000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		82.44	0.75		1	1	
Clostridium innocuum		7.23	0.44		99	99	
Lactobacillus catenaformis		7.17	0.47		99	1	
07250001	**Low discrimination**						**07250001**
					GRA	SPO	
Fusobacterium mortiferum		83.66	0.40	URE 1	1	1	
Clostridium innocuum		7.33	0.10		99	99	
Lactobacillus catenaformis		7.27	0.12		99	1	
07250002	**Low discrimination**						**07250002**
					GRA	SPO	
Fusobacterium mortiferum		32.24	0.58	BGL 10	1	1	
Clostridium innocuum		25.43	0.44		99	99	
Lactobacillus catenaformis		25.22	0.47		99	1	
07250010	**Low discrimination**						**07250010**
					GRA	SPO	
Fusobacterium mortiferum		44.01	0.40	MLZ 1	1	1	
Lactobacillus jensenii		30.02	0.28		99	1	
Clostridium innocuum		15.91	0.20		99	99	
07250020	**Low discrimination**						**07250020**
					GRA	SPO	
Clostridium barati		52.19	0.36	ESL 87	99	99	
Lactobacillus catenaformis		27.02	0.36	GAL 95	99	1	
Fusobacterium mortiferum		12.56	0.40		1	1	
07250022	**Low discrimination**						**07250022**
					SPO		
Clostridium barati		80.72	0.44	ESL 87	99		
Lactobacillus catenaformis		13.93	0.36	GAL 95	1		
Lactobacillus jensenii		2.21	0.23		1		
07250040	**Low discrimination**						**07250040**
					GRA	SPO	
Fusobacterium mortiferum		76.01	0.40	SOR 1	1	1	
Clostridium barati		9.15	0.10		99	99	
Clostridium innocuum		6.66	0.10		99	99	
07250100	**Good identification**						**07250100**
					GRA	SPO	
Fusobacterium mortiferum		93.03	0.81		1	1	
Lactobacillus catenaformis		5.20	0.50		99	1	
Clostridium barati		1.67	0.36		99	99	
07250101	**Acceptable identification**						**07250101**
					GRA	SPO	
Fusobacterium mortiferum		91.69	0.47	URE 1	1	1	
Lactobacillus catenaformis		5.12	0.15		99	1	
Clostridium barati		1.65	0.02		99	99	
07250102	**Low discrimination**						**07250102**
					GRA	SPO	
Fusobacterium mortiferum		47.15	0.65	BGL 10	1	1	
Lactobacillus catenaformis		23.71	0.50		99	1	
Clostridium barati		22.90	0.44		99	99	
07250110	**Acceptable identification**						**07250110**
					GRA	SPO	
Fusobacterium mortiferum		93.12	0.47	MLZ 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		5.20	0.15		99	1	
Clostridium barati		1.68	0.02		99	99	
07250120	**Low discrimination**						**07250120**
					GRA	SPO	
Clostridium barati		68.63	0.44	ESL 87	99	99	
Lactobacillus catenaformis		17.77	0.39	GAL 95	99	1	
Fusobacterium mortiferum		12.85	0.47		1	1	
07250122	**Low discrimination**						**07250122**
					SPO		
Clostridium barati		84.52	0.52	ESL 87	99		
Lactobacillus acidophilus		7.43	0.36	GAL 95	1		
Lactobacillus catenaformis		7.29	0.39		1		
07250140	**Low discrimination**						**07250140**
					GRA	SPO	
Fusobacterium mortiferum		82.60	0.47	SOR 1	1	1	
Clostridium barati		12.79	0.18		99	99	
Lactobacillus catenaformis		4.62	0.15		99	1	
07250200	**Low discrimination**						**07250200**
					GRA	SPO	
Fusobacterium mortiferum		76.01	0.40	RHA 1	1	1	
Clostridium barati		9.15	0.10		99	99	
Clostridium innocuum		6.66	0.10		99	99	
07250300	**Low discrimination**						**07250300**
					GRA	SPO	
Fusobacterium mortiferum		82.60	0.47	RHA 1	1	1	
Clostridium barati		12.79	0.18		99	99	
Lactobacillus catenaformis		4.62	0.15		99	1	
07250500	**Acceptable identification**						**07250500**
					GRA	SPO	
Fusobacterium mortiferum		93.12	0.47	ARA 1	1	1	
Lactobacillus catenaformis		5.20	0.15		99	1	
Clostridium barati		1.68	0.02		99	99	
07251000	**Low discrimination**						**07251000**
					SPO		
Lactobacillus catenaformis		61.07	0.81	MLT 90	1		
Lactobacillus jensenii		14.52	0.70		1		
Clostridium innocuum		11.82	0.66		99		
07251001	**Low discrimination**						**07251001**
					SPO		
Lactobacillus catenaformis		59.71	0.47	MLT 90	1		
Lactobacillus jensenii		14.20	0.36	URE 1	1		
Clostridium innocuum		11.56	0.32		99		
07251002	**Low discrimination**						**07251002**
					SPO		
Lactobacillus catenaformis		66.77	0.81	MLT 90	1		
Clostridium innocuum		12.92	0.66		99		
Lactobacillus jensenii		10.59	0.67		1		
07251003	**Low discrimination**						**07251003**
					SPO		
Lactobacillus catenaformis		65.12	0.47	MLT 90	1		
Clostridium innocuum		12.60	0.32	URE 1	99		
Lactobacillus jensenii		10.32	0.33		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07251010	**Low discrimination**					**07251010**
					SPO	
Lactobacillus jensenii		79.66	0.62	MLT 95	1	
Lactobacillus catenaformis		10.15	0.47		1	
Clostridium innocuum		8.10	0.42		99	
07251012	**Low discrimination**					**07251012**
					SPO	
Lactobacillus jensenii		72.91	0.59	MLT 95	1	
Lactobacillus catenaformis		13.94	0.47		1	
Clostridium innocuum		11.13	0.42		99	
07251020	**Low discrimination**					**07251020**
					SPO	
Lactobacillus catenaformis		61.47	0.70	MLT 90	1	
Clostridium barati		22.79	0.58		99	
Lactobacillus jensenii		14.62	0.60		1	
07251021	**Low discrimination**					**07251021**
					SPO	
Lactobacillus catenaformis		56.76	0.36	MLT 90	1	
Clostridium barati		21.04	0.24	URE 1	99	
Lactobacillus jensenii		13.50	0.26		1	
07251022	**Low discrimination**					**07251022**
					SPO	
Clostridium barati		46.06	0.66	ESL 87	99	
Lactobacillus catenaformis		41.42	0.70		1	
Lactobacillus jensenii		6.57	0.57		1	
07251030	**Low discrimination**					**07251030**
					SPO	
Lactobacillus jensenii		84.84	0.52	MLT 95	1	
Lactobacillus catenaformis		10.81	0.36		1	
Clostridium barati		4.01	0.24		99	
07251032	**Low discrimination**					**07251032**
					SPO	
Lactobacillus jensenii		69.68	0.49	MLT 95	1	
Clostridium barati		14.81	0.32		99	
Lactobacillus catenaformis		13.32	0.36		1	
07251040	**Low discrimination**					**07251040**
					SPO	
Lactobacillus catenaformis		53.19	0.47	MLT 90	1	
Clostridium barati		14.14	0.31	SOR 1	99	
Lactobacillus jensenii		12.65	0.36		1	
07251042	**Low discrimination**					**07251042**
					SPO	
Lactobacillus catenaformis		38.76	0.47	MLT 90	1	
Clostridium barati		30.92	0.40	SOR 1	99	
Lactobacillus acidophilus		15.75	0.37		1	
07251060	**Low discrimination**					**07251060**
					SPO	
Clostridium barati		71.36	0.40	ESL 87	99	
Lactobacillus catenaformis		22.36	0.36	SOR 8	1	
Lactobacillus jensenii		5.32	0.26		1	
07251062	**Low discrimination**					**07251062**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		77.36	0.48	ESL 87	99		
Lactobacillus acidophilus		13.14	0.37	SOR 8	1		
Lactobacillus catenaformis		8.08	0.36		1		
07251100	**Low discrimination**						**07251100**
					GRA	SPO	
Lactobacillus catenaformis		74.30	0.84	MLT 90	99	1	
Fusobacterium mortiferum		20.15	0.84		1	1	
Clostridium barati		4.59	0.58		99	99	
07251101	**Low discrimination**						**07251101**
					GRA		
Lactobacillus catenaformis		70.60	0.50	MLT 90	99		
Fusobacterium mortiferum		19.14	0.50	URE 1	1		
Actinomyces naeslundii		5.13	0.52		99		
07251102	**Low discrimination**						**07251102**
					SPO		
Lactobacillus catenaformis		67.85	0.84	MLT 90	1		
Lactobacillus acidophilus		17.28	0.71		1		
Clostridium barati		12.58	0.66		99		
07251103	**Low discrimination**						**07251103**
					SPO		
Lactobacillus catenaformis		64.82	0.50	MLT 90	1		
Lactobacillus acidophilus		16.50	0.36	URE 1	1		
Clostridium barati		12.01	0.32		99		
07251110	**Low discrimination**						**07251110**
					GRA	SPO	
Lactobacillus catenaformis		71.28	0.50	MLT 90	99	1	
Fusobacterium mortiferum		19.33	0.50	MLZ 1	1	1	
Clostridium barati		4.40	0.24		99	99	
07251112	**Low discrimination**						**07251112**
					SPO		
Lactobacillus catenaformis		65.78	0.50	MLT 90	1		
Lactobacillus acidophilus		16.75	0.36	MLZ 1	1		
Clostridium barati		12.19	0.32		99		
07251120	**Low discrimination**						**07251120**
					SPO		
Lactobacillus catenaformis		55.66	0.73	MLT 90	1		
Clostridium barati		41.27	0.66		99		
Clostridium paraputrificum		0.83	0.32		99		
07251121	**Low discrimination**						**07251121**
					SPO		
Lactobacillus catenaformis		48.14	0.39	MLT 90	1		
Clostridium barati		35.69	0.32	URE 1	99		
Actinomyces naeslundii		13.99	0.52		1		
07251122	**Low discrimination**						**07251122**
					SPO		
Clostridium barati		52.14	0.74	ESL 87	99		
Lactobacillus acidophilus		23.88	0.71		1		
Lactobacillus catenaformis		23.44	0.73		1		
07251123	**Low discrimination**						**07251123**
					SPO		
Clostridium barati		48.89	0.40	ESL 87	99		
Lactobacillus acidophilus		22.39	0.36	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		21.98	0.39		1		
07251130	**Low discrimination**						**07251130**
					SPO		
Lactobacillus catenaformis		54.06	0.39	MLT 90	1		
Clostridium barati		40.08	0.32	MLZ 1	99		
Lactobacillus jensenii		2.86	0.17		1		
07251132	**Low discrimination**						**07251132**
					SPO		
Clostridium barati		51.43	0.40	ESL 87	99		
Lactobacillus acidophilus		23.55	0.36	MLZ 1	1		
Lactobacillus catenaformis		23.12	0.39		1		
07251140	**Low discrimination**						**07251140**
					GRA	SPO	
Lactobacillus catenaformis		54.02	0.50	MLT 90	99	1	
Clostridium barati		28.73	0.40	SOR 1	99	99	
Fusobacterium mortiferum		14.65	0.50		1	1	
07251141	**Low discrimination**						**07251141**
					SPO		
Actinomyces naeslundii		41.32	0.39	MLT 94	1		
Lactobacillus catenaformis		32.54	0.15	RAF 92	1		
Clostridium barati		17.31	0.05		99		
07251142	**Low discrimination**						**07251142**
					SPO		
Lactobacillus acidophilus		54.19	0.55	MLT 94	1		
Clostridium barati		27.75	0.48	SOR 11	99		
Lactobacillus catenaformis		17.39	0.50		1		
07251160	**Low discrimination**						**07251160**
					SPO		
Clostridium barati		82.71	0.48	ESL 87	99		
Lactobacillus catenaformis		12.96	0.39	SOR 8	1		
Lactobacillus acidophilus		1.63	0.21		1		
07251162	**Low discrimination**						**07251162**
					SPO		
Clostridium barati		58.50	0.56	ESL 87	99		
Lactobacillus acidophilus		38.08	0.55	SOR 8	1		
Lactobacillus catenaformis		3.06	0.39		1		
07251200	**Low discrimination**						**07251200**
					SPO		
Lactobacillus catenaformis		53.36	0.47	MLT 90	1		
Clostridium barati		14.19	0.31	RHA 1	99		
Lactobacillus jensenii		12.69	0.36		1		
07251202	**Low discrimination**						**07251202**
					SPO		
Lactobacillus catenaformis		41.38	0.47	MLT 90	1		
Clostridium barati		33.01	0.40	RHA 1	99		
Lactobacillus acidophilus		10.24	0.33		1		
07251220	**Low discrimination**						**07251220**
					SPO		
Clostridium barati		71.89	0.40	ESL 87	99		
Lactobacillus catenaformis		22.53	0.36	RHA 8	1		
Lactobacillus jensenii		5.36	0.26		1		
07251222	**Low discrimination**						**07251222**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium barati		81.40	0.48	ESL 87	99		
Lactobacillus catenaformis		8.50	0.36	RHA 8	1		
Lactobacillus acidophilus		8.42	0.33		1		
07251300	**Low discrimination**						**07251300**
					GRA	SPO	
Lactobacillus catenaformis		54.63	0.50	MLT 90	99	1	
Clostridium barati		29.06	0.40	RHA 1	99	99	
Fusobacterium mortiferum		14.81	0.50		1	1	
07251302	**Low discrimination**						**07251302**
					SPO		
Lactobacillus acidophilus		41.90	0.51	MLT 94	1		
Clostridium barati		35.23	0.48	RHA 7	99		
Lactobacillus catenaformis		22.08	0.50		1		
07251320	**Low discrimination**						**07251320**
					SPO		
Clostridium barati		84.58	0.48	ESL 87	99		
Lactobacillus catenaformis		13.25	0.39	RHA 8	1		
Lactobacillus acidophilus		1.02	0.17		1		
07251322	**Low discrimination**						**07251322**
					SPO		
Clostridium barati		67.66	0.56	ESL 87	99		
Lactobacillus acidophilus		26.82	0.51	RHA 8	1		
Lactobacillus catenaformis		3.53	0.39		1		
07251342	**Low discrimination**						**07251342**
					SPO		
Lactobacillus acidophilus		61.17	0.36	MLT 94	1		
Clostridium barati		36.19	0.30	RHA 7	99		
Lactobacillus catenaformis		2.64	0.15	SOR 11	1		
07251362	**Low discrimination**						**07251362**
					SPO		
Clostridium barati		63.72	0.38	ESL 87	99		
Lactobacillus acidophilus		35.90	0.36	RHA 8	1		
Lactobacillus catenaformis		0.39	0.05	SOR 8	1		
07251400	**Low discrimination**						**07251400**
					SPO		
Clostridium innocuum		70.09	0.53	MAN 99	99		
Lactobacillus catenaformis		20.73	0.47		1		
Lactobacillus jensenii		4.93	0.36		1		
07251402	**Low discrimination**						**07251402**
					SPO		
Clostridium innocuum		72.17	0.53	MAN 99	99		
Lactobacillus catenaformis		21.35	0.47		1		
Lactobacillus jensenii		3.38	0.33		1		
07251420	**Low discrimination**						**07251420**
					SPO		
Lactobacillus catenaformis		57.30	0.36	MLT 90	1		
Clostridium barati		21.24	0.24	ARA 1	99		
Lactobacillus jensenii		13.63	0.26		1		
07251500	**Low discrimination**						**07251500**
					GRA	SPO	
Lactobacillus catenaformis		73.18	0.50	MLT 90	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		19.84	0.50	ARA 1	1	1	
Clostridium barati		4.52	0.24		99	99	
07251502	**Low discrimination**						**07251502**
					SPO		
Lactobacillus catenaformis		66.51	0.50	MLT 90	1		
Lactobacillus acidophilus		16.94	0.36	ARA 1	1		
Clostridium barati		12.33	0.32		99		
07251520	**Low discrimination**						**07251520**
					SPO		
Lactobacillus catenaformis		55.97	0.39	MLT 90	1		
Clostridium barati		41.49	0.32	ARA 1	99		
Clostridium paraputrificum		0.84	-.03		99		
07251522	**Low discrimination**						**07251522**
					SPO		
Clostridium barati		52.22	0.40	ESL 87	99		
Lactobacillus acidophilus		23.92	0.36	ARA 1	1		
Lactobacillus catenaformis		23.48	0.39		1		
07252000	**Very good identification**						**07252000**
					GRA	SPO	
Clostridium innocuum		99.81	0.78	GAL 95	99	99	
Fusobacterium mortiferum		0.12	0.40		1	1	
Clostridium difficile		0.03	0.15		99	99	
07252001	**Good identification**						**07252001**
					SPO		
Clostridium innocuum		99.69	0.44	GAL 95	99		
Actinomyces naeslundii		0.31	0.25	URE 1	1		
07252002	**Very good identification**						**07252002**
					GRA	SPO	
Clostridium innocuum		99.93	0.78	GAL 95	99	99	
Clostridium barati		0.04	0.18		99	99	
Fusobacterium mortiferum		0.01	0.24		1	1	
07252003	**Good identification**						**07252003**
					SPO		
Clostridium innocuum		99.69	0.44	GAL 95	99		
Actinomyces naeslundii		0.31	0.25	URE 1	1		
07252010	**Good identification**						**07252010**
Clostridium innocuum		96.56	0.55	GAL 95			
Clostridium difficile		3.44	0.26	MLZ 4			
07252012	**Good identification**						**07252012**
Clostridium innocuum		100.0	0.55	GAL 95			
07252020	**Good identification**						**07252020**
					SPO	CAT	
Clostridium innocuum		95.52	0.44	GAL 95	99	1	
Clostridium barati		3.98	0.18	NAG 1	99	1	
Propionibacterium granulosum		0.26	0.09		1	90	
07252022	**Low discrimination**						**07252022**
					SPO		
Clostridium innocuum		88.70	0.44	GAL 95	99		
Clostridium barati		11.08	0.26	NAG 1	99		
Lactobacillus catenaformis		0.22	0.02		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07252040	**Good identification**						**07252040**
Clostridium innocuum		100.0	0.44	GAL 95			
07252042	**Good identification**						**07252042**
Clostridium innocuum		99.64	0.44	GAL 95			
Clostridium barati		0.36	-.01	SOR 1			
07252100	**Low discrimination**						**07252100**
					GRA	SPO	
Clostridium innocuum		75.11	0.44	GAL 95	99	99	
Fusobacterium mortiferum		20.20	0.47	LAC 1	1	1	
Clostridium barati		3.13	0.18		99	99	
07252102	**Low discrimination**						**07252102**
					GRA	SPO	
Clostridium innocuum		83.64	0.44	GAL 95	99	99	
Clostridium barati		10.45	0.26	LAC 1	99	99	
Fusobacterium mortiferum		2.50	0.30		1	1	
07252200	**Good identification**						**07252200**
Clostridium innocuum		100.0	0.44	GAL 95			
07252202	**Good identification**						**07252202**
Clostridium innocuum		99.64	0.44	GAL 95			
Clostridium barati		0.36	-.01	RHA 1			
07252400	**Very good identification**						**07252400**
Clostridium innocuum		100.0	0.65	GAL 95			
07252402	**Very good identification**						**07252402**
Clostridium innocuum		100.0	0.65	GAL 95			
07252410	**Good identification**						**07252410**
Clostridium innocuum		100.0	0.42	GAL 95			
07252412	**Good identification**						**07252412**
Clostridium innocuum		100.0	0.42	GAL 95			
07253000	**Excellent identification**						**07253000**
					GRA	SPO	
Clostridium innocuum		99.91	1.00		99	99	
Lactobacillus catenaformis		0.05	0.47		99	1	
Fusobacterium mortiferum		0.01	0.43		1	1	
07253001	**Very good identification**						**07253001**
					SPO		
Clostridium innocuum		99.54	0.66	URE 1	99		
Actinomyces naeslundii		0.39	0.49		1		
Lactobacillus catenaformis		0.05	0.12		1		
07253002	**Excellent identification**						**07253002**
					SPO		
Clostridium innocuum		99.88	1.00		99		
Lactobacillus catenaformis		0.05	0.47		1		
Clostridium barati		0.04	0.40		99		
07253003	**Very good identification**						**07253003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium innocuum		99.52	0.66	URE 1	99	
Actinomyces naeslundii		0.39	0.49		1	
Lactobacillus catenaformis		0.05	0.12		1	
07253010	**Very good identification**					**07253010**
					SPO	
Clostridium innocuum		99.88	0.76	MLZ 4	99	
Lactobacillus jensenii		0.10	0.28		1	
Lactobacillus catenaformis		0.01	0.12		1	
07253011	**Good identification**					**07253011**
					SPO	
Clostridium innocuum		99.51	0.42	MLZ 4	99	
Actinomyces naeslundii		0.49	0.27	URE 1	1	
07253012	**Very good identification**					**07253012**
					SPO	
Clostridium innocuum		99.91	0.76	MLZ 4	99	
Lactobacillus jensenii		0.07	0.25		1	
Lactobacillus catenaformis		0.01	0.12		1	
07253013	**Good identification**					**07253013**
					SPO	
Clostridium innocuum		99.51	0.42	MLZ 4	99	
Actinomyces naeslundii		0.49	0.27	URE 1	1	
07253020	**Good identification**					**07253020**
					SPO	
Clostridium innocuum		94.29	0.66	NAG 1	99	
Clostridium barati		3.93	0.40		99	
Lactobacillus catenaformis		1.23	0.36		1	
07253022	**Low discrimination**					**07253022**
					SPO	
Clostridium innocuum		86.82	0.66	NAG 1	99	
Clostridium barati		10.85	0.48		99	
Lactobacillus catenaformis		1.13	0.36		1	
07253030	**Good identification**					**07253030**
					SPO	
Clostridium innocuum		96.03	0.42	MLZ 4	99	
Lactobacillus jensenii		2.38	0.17	NAG 1	1	
Clostridium barati		0.97	0.05		99	
07253032	**Acceptable identification**					**07253032**
					SPO	
Clostridium innocuum		94.40	0.42	MLZ 4	99	
Clostridium barati		2.86	0.14	NAG 1	99	
Lactobacillus jensenii		1.56	0.14		1	
07253040	**Very good identification**					**07253040**
					SPO	
Clostridium innocuum		99.77	0.66	SOR 1	99	
Clostridium barati		0.12	0.13		99	
Actinomyces naeslundii		0.05	0.33		1	
07253042	**Very good identification**					**07253042**
					SPO	
Clostridium innocuum		99.43	0.66	SOR 1	99	
Clostridium barati		0.36	0.21		99	
Lactobacillus acidophilus		0.11	0.15		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07253050	**Good identification**					**07253050**
Clostridium innocuum		100.0	0.42	MLZ 4		
07253052	**Good identification**					**07253052**
Clostridium innocuum		100.0	0.42	MLZ 4		
07253061	**Good identification**					**07253061**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07253063	**Good identification**					**07253063**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07253100	**Low discrimination**					**07253100**
					SPO	
Clostridium innocuum		86.80	0.66	LAC 1	99	
Lactobacillus catenaformis		6.80	0.50		1	
Clostridium barati		3.61	0.40		99	
07253101	**Low discrimination**					**07253101**
					SPO	
Actinomyces naeslundii		48.93	0.57	MLT 94	1	
Clostridium innocuum		44.75	0.32	RAF 92	99	
Lactobacillus catenaformis		3.50	0.15		1	
07253102	**Low discrimination**					**07253102**
					SPO	
Clostridium innocuum		75.65	0.66	LAC 1	99	
Clostridium barati		9.45	0.48		99	
Lactobacillus acidophilus		7.86	0.49		1	
07253103	**Low discrimination**					**07253103**
					SPO	
Actinomyces naeslundii		45.55	0.57	MLT 94	1	
Clostridium innocuum		41.66	0.32	RAF 92	99	
Clostridium barati		5.20	0.14		99	
07253110	**Acceptable identification**					**07253110**
					SPO	
Clostridium innocuum		90.10	0.42	LAC 1	99	
Clostridium tertium		5.99	0.23	MLZ 4	99	
Lactobacillus catenaformis		1.71	0.15		1	
07253112	**Low discrimination**					**07253112**
Clostridium innocuum		77.76	0.42	LAC 1		
Clostridium tertium		15.50	0.31	MLZ 4		
Clostridium barati		2.36	0.14			
07253120	**Low discrimination**					**07253120**
					SPO	
Clostridium barati		73.10	0.48	ESL 87	99	
Lactobacillus catenaformis		11.45	0.39	MAN 8	1	
Clostridium innocuum		5.91	0.32		99	
07253121	**Low discrimination**					**07253121**
					SPO	
Actinomyces naeslundii		88.33	0.57	MLT 94	1	
Clostridium barati		10.09	0.14	RAF 92	99	
Lactobacillus catenaformis		1.58	0.05		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07253122	**Low discrimination**					**07253122**
					SPO	
Clostridium barati		69.37	0.56	ESL 87	99	
Lactobacillus acidophilus		19.23	0.49	MAN 8	1	
Lactobacillus catenaformis		3.62	0.39		1	
07253123	**Low discrimination**					**07253123**
					SPO	
Actinomyces naeslundii		68.10	0.57	MLT 94	1	
Clostridium barati		23.34	0.22	RAF 92	99	
Lactobacillus acidophilus		6.47	0.15		1	
07253132	**Low discrimination**					**07253132**
					SPO	
Clostridium tertium		80.62	0.39	MLT 99	99	
Clostridium barati		12.25	0.22	TRE 90	99	
Lactobacillus acidophilus		3.40	0.15		1	
07253141	**Good identification**					**07253141**
Actinomyces naeslundii		100.0	0.44	MLT 94		
07253143	**Acceptable identification**					**07253143**
Actinomyces naeslundii		93.76	0.44	MLT 94		
Lactobacillus acidophilus		6.24	-.01	RAF 92		
07253161	**Good identification**					**07253161**
Actinomyces naeslundii		100.0	0.44	MLT 94		
07253162	**Low discrimination**					**07253162**
					SPO	
Clostridium barati		69.03	0.38	ESL 87	99	
Lactobacillus acidophilus		27.19	0.33	MAN 8	1	
Actinomyces naeslundii		2.75	0.41	SOR 8	1	
07253163	**Low discrimination**					**07253163**
					SPO	
Actinomyces naeslundii		80.95	0.44	MLT 94	1	
Clostridium barati		13.67	0.04	RAF 92	99	
Lactobacillus acidophilus		5.38	-.01		1	
07253200	**Very good identification**					**07253200**
					SPO	
Clostridium innocuum		99.81	0.66	RHA 1	99	
Clostridium barati		0.12	0.13		99	
Lactobacillus catenaformis		0.05	0.12		1	
07253202	**Very good identification**					**07253202**
					SPO	
Clostridium innocuum		99.52	0.66	RHA 1	99	
Clostridium barati		0.36	0.21		99	
Lactobacillus acidophilus		0.07	0.11		1	
07253210	**Good identification**					**07253210**
Clostridium innocuum		100.0	0.42	RHA 1		
07253212	**Good identification**					**07253212**
Clostridium innocuum		100.0	0.42	RHA 1		
07253322	**Low discrimination**					**07253322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium barati		60.17	0.38	ESL 87	99	
Clostridium ramosum		24.76	0.33	MAN 8	92	
Lactobacillus acidophilus		14.44	0.29	RHA 8	1	
07253400	**Very good identification**					**07253400**
					SPO	
Clostridium innocuum		100.0	0.87		99	
Lactobacillus catenaformis		0.00	0.12		1	
Lactobacillus jensenii		0.00	0.02		1	
07253401	**Good identification**					**07253401**
Clostridium innocuum		100.0	0.53	URE 1		
07253402	**Very good identification**					**07253402**
					SPO	
Clostridium innocuum		100.0	0.87		99	
Lactobacillus catenaformis		0.00	0.12		1	
Clostridium barati		0.00	0.05		99	
07253403	**Good identification**					**07253403**
Clostridium innocuum		100.0	0.53	URE 1		
07253410	**Very good identification**					**07253410**
Clostridium innocuum		100.0	0.64	MLZ 4		
07253412	**Very good identification**					**07253412**
Clostridium innocuum		100.0	0.64	MLZ 4		
07253420	**Good identification**					**07253420**
					SPO	
Clostridium innocuum		99.69	0.53	NAG 1	99	
Clostridium barati		0.24	0.05		99	
Lactobacillus catenaformis		0.07	0.02		1	
07253422	**Good identification**					**07253422**
					SPO	
Clostridium innocuum		99.22	0.53	NAG 1	99	
Clostridium barati		0.71	0.14		99	
Lactobacillus catenaformis		0.07	0.02		1	
07253440	**Good identification**					**07253440**
Clostridium innocuum		100.0	0.53	SOR 1		
07253442	**Good identification**					**07253442**
Clostridium innocuum		100.0	0.53	SOR 1		
07253500	**Good identification**					**07253500**
					SPO	
Clostridium innocuum		99.06	0.53	LAC 1	99	
Lactobacillus catenaformis		0.44	0.15		1	
Clostridium barati		0.24	0.05		99	
07253502	**Good identification**					**07253502**
					SPO	
Clostridium innocuum		97.70	0.53	LAC 1	99	
Clostridium barati		0.70	0.14		99	
Lactobacillus acidophilus		0.58	0.15		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07253600	**Good identification**						**07253600**
Clostridium innocuum		100.0	0.53	RHA 1			
07253602	**Good identification**						**07253602**
Clostridium innocuum		100.0	0.53	RHA 1			
07254000	**Acceptable identification**						**07254000**
					GRA	SPO	
Fusobacterium mortiferum		92.68	0.55	XYL 7	1	1	
Clostridium innocuum		5.68	0.22		99	99	
Lactobacillus catenaformis		1.08	0.12		99	1	
07254002	**Low discrimination**						**07254002**
					GRA	SPO	
Fusobacterium mortiferum		58.57	0.39	XYL 7	1	1	
Clostridium innocuum		32.31	0.22	BGL 10	99	99	
Lactobacillus catenaformis		6.15	0.12		99	1	
07254100	**Very good identification**						**07254100**
					GRA	SPO	
Fusobacterium mortiferum		99.02	0.62	XYL 7	1	1	
Lactobacillus catenaformis		0.74	0.15		99	1	
Clostridium barati		0.24	0.02		99	99	
07254102	**Low discrimination**						**07254102**
					GRA	SPO	
Fusobacterium mortiferum		86.97	0.45	XYL 7	1	1	
Lactobacillus catenaformis		5.87	0.15	BGL 10	99	1	
Clostridium barati		5.67	0.10		99	99	
07255000	**Low discrimination**						**07255000**
					GRA	SPO	
Fusobacterium mortiferum		36.29	0.58	XYL 7	1	1	
Clostridium innocuum		28.18	0.44		99	99	
Lactobacillus catenaformis		27.94	0.47		99	1	
07255002	**Low discrimination**						**07255002**
					SPO		
Clostridium innocuum		41.30	0.44	MAN 99	99		
Lactobacillus catenaformis		40.96	0.47	XYL 5	1		
Lactobacillus jensenii		6.49	0.33		1		
07255020	**Low discrimination**						**07255020**
					SPO		
Lactobacillus catenaformis		58.76	0.36	MLT 90	1		
Clostridium barati		21.78	0.24	XYL 1	99		
Lactobacillus jensenii		13.98	0.26		1		
07255100	**Low discrimination**						**07255100**
					GRA	SPO	
Fusobacterium mortiferum		64.10	0.65	XYL 7	1	1	
Lactobacillus catenaformis		31.72	0.50		99	1	
Clostridium barati		1.96	0.24		99	99	
07255102	**Low discrimination**						**07255102**
					GRA		
Lactobacillus catenaformis		58.73	0.50	MLT 90	99		
Lactobacillus acidophilus		14.95	0.36	XYL 1	99		
Fusobacterium mortiferum		13.19	0.48		1		
07255120	**Low discrimination**						**07255120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Lactobacillus catenaformis		52.79	0.39	MLT 90	99	1	
Clostridium barati		39.14	0.32	XYL 1	99	99	
Fusobacterium mortiferum		4.31	0.31		1	1	
07255122	**Low discrimination**						**07255122**
					SPO		
Clostridium barati		50.07	0.40	ESL 87	99		
Lactobacillus acidophilus		22.93	0.36	XYL 1	1		
Lactobacillus catenaformis		22.51	0.39		1		
07255502	**Low discrimination**						**07255502**
Bifidobacterium adolescentis		69.37	0.36	MLT 99			
Lactobacillus catenaformis		9.84	0.15	RAF 92			
Actinomyces israelii		9.20	0.27				
07255512	**Good identification**						**07255512**
Bifidobacterium adolescentis		97.57	0.37	MLT 99			
Clostridium clostridioforme		2.43	0.07	RAF 92			
07256000	**Very good identification**						**07256000**
					GRA	SPO	
Clostridium innocuum		99.83	0.56	GAL 95	99	99	
Fusobacterium mortiferum		0.17	0.21	XYL 5	1	1	
07256002	**Very good identification**						**07256002**
Clostridium innocuum		100.0	0.56	GAL 95			
07256400	**Good identification**						**07256400**
Clostridium innocuum		100.0	0.43	GAL 95			
07256402	**Good identification**						**07256402**
Clostridium innocuum		100.0	0.43	GAL 95			
07257000	**Very good identification**						**07257000**
					GRA	SPO	
Clostridium innocuum		99.96	0.78	XYL 5	99	99	
Actinomyces israelii		0.02	0.28		99	1	
Fusobacterium mortiferum		0.01	0.24		1	1	
07257001	**Good identification**						**07257001**
Clostridium innocuum		100.0	0.44	XYL 5			
07257002	**Very good identification**						**07257002**
					SPO		
Clostridium innocuum		99.96	0.78	XYL 5	99		
Actinomyces israelii		0.03	0.31		1		
Lactobacillus catenaformis		0.01	0.12		1		
07257003	**Good identification**						**07257003**
Clostridium innocuum		100.0	0.44	XYL 5			
07257010	**Good identification**						**07257010**
					SPO		
Clostridium innocuum		99.94	0.55	XYL 5	99		
Actinomyces israelii		0.06	0.14	MLZ 4	1		
07257012	**Good identification**						**07257012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium innocuum		99.86	0.55	XYL 5	99	
Actinomyces israelii		0.09	0.17	MLZ 4	1	
Clostridium tertium		0.05	-.01		99	
07257020	**Good identification**					**07257020**
					SPO	
Clostridium innocuum		98.96	0.44	XYL 5	99	
Clostridium barati		0.79	0.05	NAG 1	99	
Lactobacillus catenaformis		0.25	0.02		1	
07257022	**Good identification**					**07257022**
Clostridium innocuum		97.07	0.44	XYL 5		
Clostridium barati		2.33	0.14	NAG 1		
Clostridium tertium		0.36	0.03			
07257040	**Good identification**					**07257040**
					SPO	
Clostridium innocuum		99.37	0.44	XYL 5	99	
Actinomyces israelii		0.63	0.21	SOR 1	1	
07257042	**Good identification**					**07257042**
					SPO	
Clostridium innocuum		99.06	0.44	XYL 5	99	
Actinomyces israelii		0.94	0.24	SOR 1	1	
07257100	**Low discrimination**					**07257100**
					SPO	
Clostridium innocuum		86.27	0.44	XYL 5	99	
Actinomyces israelii		5.60	0.39	LAC 1	1	
Clostridium tertium		3.52	0.21		99	
07257102	**Low discrimination**					**07257102**
					SPO	
Clostridium innocuum		77.21	0.44	XYL 5	99	
Clostridium tertium		9.46	0.29	LAC 1	99	
Actinomyces israelii		7.52	0.42		1	
07257122	**Low discrimination**					**07257122**
					SPO	
Clostridium tertium		76.66	0.37	MLT 99	99	
Clostridium barati		15.00	0.22	TRE 90	99	
Lactobacillus acidophilus		4.16	0.15		1	
07257132	**Good identification**					**07257132**
Clostridium tertium		100.0	0.41	MLT 99		
07257200	**Good identification**					**07257200**
Clostridium innocuum		100.0	0.44	XYL 5		
07257202	**Good identification**					**07257202**
Clostridium innocuum		100.0	0.44	XYL 5		
07257400	**Very good identification**					**07257400**
					SPO	
Clostridium innocuum		99.92	0.65	XYL 5	99	
Actinomyces israelii		0.08	0.27		1	
07257402	**Very good identification**					**07257402**
					SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		99.89	0.65	XYL 5	99		
Actinomyces israelii		0.11	0.30		1		
07257410	**Good identification**						**07257410**
Clostridium innocuum		100.0	0.42	XYL 5			
07257412	**Good identification**						**07257412**
					SPO		
Clostridium innocuum		99.59	0.42	XYL 5	99		
Actinomyces israelii		0.41	0.16	MLZ 4	1		
07257512	**Low discrimination**						**07257512**
					SPO		
Bifidobacterium adolescentis		83.16	0.36	MLT 99	1		
Actinomyces israelii		9.34	0.26	RAF 92	1		
Clostridium innocuum		5.77	0.07		99		
07260100	**Good identification**						**07260100**
Fusobacterium mortiferum		100.0	0.38	FRU 99			
07261000	**Low discrimination**						**07261000**
					GRA	SPO	
Lactobacillus jensenii		62.85	0.44	MLT 95	99	1	
Clostridium innocuum		31.21	0.36	FRU 99	99	99	
Fusobacterium mortiferum		4.59	0.34		1	1	
07261002	**Low discrimination**						**07261002**
					SPO		
Lactobacillus jensenii		49.15	0.41	MLT 95	1		
Clostridium innocuum		36.61	0.36	FRU 99	99		
Lactobacillus acidophilus		11.49	0.27		1		
07261010	**Acceptable identification**						**07261010**
					SPO		
Lactobacillus jensenii		94.16	0.36	MLT 95	1		
Clostridium innocuum		5.85	0.13	FRU 99	99		
07261100	**Low discrimination**						**07261100**
					COC	GRA	
Fusobacterium mortiferum		57.64	0.41	FRU 99	1	1	
Atopobium parvulum		16.58	0.20		99	99	
Lactobacillus catenaformis		7.19	0.15		1	99	
07261102	**Acceptable identification**						**07261102**
					SPO		
Lactobacillus acidophilus		94.47	0.45	MLT 94	1		
Clostridium barati		1.79	0.14	FRU 99	99		
Lactobacillus catenaformis		1.12	0.15		1		
07261122	**Acceptable identification**						**07261122**
					SPO		
Lactobacillus acidophilus		93.48	0.45	MLT 94	1		
Clostridium barati		5.30	0.22	FRU 99	99		
Actinomyces naeslundii		0.54	0.32		1		
07262000	**Good identification**						**07262000**
Clostridium innocuum		100.0	0.48	FRU 95			
07262002	**Good identification**						**07262002**
Clostridium innocuum		100.0	0.48	FRU 95			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07262400	**Good identification**					**07262400**
Clostridium innocuum		100.0	0.36	FRU 95		
07262402	**Good identification**					**07262402**
Clostridium innocuum		100.0	0.36	FRU 95		
07263000	**Very good identification**					**07263000**
					SPO	
Clostridium innocuum		99.97	0.70	FRU 95	99	
Lactobacillus jensenii		0.02	0.10		1	
Actinomyces naeslundii		0.01	0.29		1	
07263001	**Good identification**					**07263001**
					SPO	
Clostridium innocuum		97.90	0.36	FRU 95	99	
Actinomyces naeslundii		2.10	0.32	URE 1	1	
07263002	**Very good identification**					**07263002**
					SPO	
Clostridium innocuum		99.96	0.70	FRU 95	99	
Lactobacillus acidophilus		0.02	0.05		1	
Actinomyces naeslundii		0.01	0.29		1	
07263003	**Good identification**					**07263003**
					SPO	
Clostridium innocuum		97.90	0.36	FRU 95	99	
Actinomyces naeslundii		2.10	0.32	URE 1	1	
07263010	**Good identification**					**07263010**
					SPO	
Clostridium innocuum		99.84	0.47	FRU 95	99	
Lactobacillus jensenii		0.16	0.02	MLZ 4	1	
07263012	**Good identification**					**07263012**
Clostridium innocuum		100.0	0.47	FRU 95		
07263020	**Good identification**					**07263020**
					SPO	
Clostridium innocuum		98.59	0.36	FRU 95	99	
Actinomyces naeslundii		1.41	0.29	NAG 1	1	
07263022	**Good identification**					**07263022**
					SPO	
Clostridium innocuum		97.01	0.36	FRU 95	99	
Lactobacillus acidophilus		1.60	0.05	NAG 1	1	
Actinomyces naeslundii		1.39	0.29		1	
07263040	**Good identification**					**07263040**
Clostridium innocuum		100.0	0.36	FRU 95		
07263042	**Good identification**					**07263042**
Clostridium innocuum		100.0	0.36	FRU 95		
07263100	**Good identification**					**07263100**
					SPO	
Clostridium innocuum		95.39	0.36	FRU 95	99	
Actinomyces naeslundii		3.88	0.37	LAC 1	1	
Clostridium tertium		0.73	0.01		99	
07263101	**Low discrimination**					**07263101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		85.80	0.40	MLT 94	1		
Clostridium innocuum		14.20	0.02	RAF 92	99		
07263102	**Low discrimination**						**07263102**
					SPO		
Clostridium innocuum		79.77	0.36	FRU 95	99		
Lactobacillus acidophilus		15.15	0.24	LAC 1	1		
Actinomyces naeslundii		3.25	0.37		1		
07263103	**Low discrimination**						**07263103**
					SPO		
Actinomyces naeslundii		85.80	0.40	MLT 94	1		
Clostridium innocuum		14.20	0.02	RAF 92	99		
07263120	**Low discrimination**						**07263120**
					SPO		
Actinomyces naeslundii		42.65	0.37	MLT 94	1		
Clostridium tertium		24.07	0.09	RAF 92	99		
Clostridium ramosum		22.69	0.07	FRU 95	92		
07263121	**Good identification**						**07263121**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07263123	**Good identification**						**07263123**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07263200	**Good identification**						**07263200**
Clostridium innocuum		100.0	0.36	FRU 95			
07263202	**Good identification**						**07263202**
Clostridium innocuum		100.0	0.36	FRU 95			
07263400	**Very good identification**						**07263400**
Clostridium innocuum		100.0	0.57	FRU 95			
07263402	**Very good identification**						**07263402**
Clostridium innocuum		100.0	0.57	FRU 95			
07267000	**Good identification**						**07267000**
Clostridium innocuum		100.0	0.48	FRU 95			
07267002	**Good identification**						**07267002**
Clostridium innocuum		100.0	0.48	FRU 95			
07267400	**Good identification**						**07267400**
Clostridium innocuum		100.0	0.36	FRU 95			
07267402	**Good identification**						**07267402**
Clostridium innocuum		100.0	0.36	FRU 95			
07270000	**Low discrimination**						**07270000**
					GRA	SPO	
Fusobacterium mortiferum		75.16	0.66		1	1	
Lactobacillus jensenii		15.60	0.44		99	1	
Clostridium innocuum		7.75	0.36		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07270002	**Low discrimination**							**07270002**
					GRA	SPO		
Lactobacillus jensenii		34.32	0.41	MLT 95	99	1		
Fusobacterium mortiferum		27.55	0.49	GAL 99	1	1		
Clostridium innocuum		25.56	0.36		99	99		
07270010	**Low discrimination**							**07270010**
					GRA	SPO		
Lactobacillus jensenii		81.09	0.36	MLT 95	99	1		
Fusobacterium mortiferum		11.84	0.31	GAL 99	1	1		
Clostridium innocuum		5.03	0.13		99	99		
07270100	**Very good identification**							**07270100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.51	0.72		1	1	1	
Clostridium barati		0.52	0.18		1	99	99	
Atopobium parvulum		0.43	0.20		99	99	1	
07270101	**Acceptable identification**							**07270101**
					GRA			
Fusobacterium mortiferum		90.18	0.38	URE 1	1			
Actinomyces naeslundii		9.82	0.33		99			
07270102	**Low discrimination**							**07270102**
					GRA	SPO		
Lactobacillus acidophilus		54.85	0.45	MLT 94	99	1		
Fusobacterium mortiferum		38.22	0.56	GAL 99	1	1		
Clostridium barati		5.40	0.26		99	99		
07270110	**Good identification**							**07270110**
					GRA			
Fusobacterium mortiferum		97.12	0.38	MLZ 1	1			
Lactobacillus jensenii		2.88	0.02		99			
07270122	**Low discrimination**							**07270122**
					GRA	SPO		
Lactobacillus acidophilus		76.08	0.45	MLT 94	99	1		
Clostridium barati		22.49	0.34	GAL 99	99	99		
Fusobacterium mortiferum		0.54	0.21		1	1		
07270140	**Good identification**							**07270140**
					GRA	SPO		
Fusobacterium mortiferum		95.69	0.38	SOR 1	1	1		
Clostridium barati		4.31	-.01		99	99		
07270300	**Good identification**							**07270300**
					GRA	SPO		
Fusobacterium mortiferum		95.69	0.38	RHA 1	1	1		
Clostridium barati		4.31	-.01		99	99		
07270500	**Good identification**							**07270500**
Fusobacterium mortiferum		100.0	0.38	ARA 1				
07271000	**Low discrimination**							**07271000**
					GRA	SPO		
Lactobacillus jensenii		84.06	0.78	MLT 95	99	1		
Clostridium innocuum		8.01	0.58		99	99		
Fusobacterium mortiferum		6.14	0.69		1	1		
07271001	**Low discrimination**							**07271001**
					SPO			
Lactobacillus jensenii		77.49	0.44	MLT 95	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		8.08	0.49	URE 1	1		
Clostridium innocuum		7.39	0.24		99		
07271002	**Low discrimination**						**07271002**
					SPO		
Lactobacillus jensenii		70.00	0.75	MLT 95	1		
Lactobacillus acidophilus		16.36	0.61		1		
Clostridium innocuum		10.01	0.58		99		
07271003	**Low discrimination**						**07271003**
Lactobacillus jensenii		63.16	0.41	MLT 95			
Lactobacillus acidophilus		14.76	0.27	URE 1			
Actinomyces naeslundii		9.87	0.49				
07271010	**Very good identification**						**07271010**
					GRA	SPO	
Lactobacillus jensenii		98.52	0.70	MLT 95	99	1	
Clostridium innocuum		1.17	0.34		99	99	
Fusobacterium mortiferum		0.22	0.34		1	1	
07271011	**Good identification**						**07271011**
					SPO		
Lactobacillus jensenii		97.24	0.36	MLT 95	1		
Actinomyces naeslundii		1.60	0.27	URE 1	1		
Clostridium innocuum		1.16	0.00		99		
07271012	**Very good identification**						**07271012**
					SPO		
Lactobacillus jensenii		97.36	0.67	MLT 95	1		
Clostridium innocuum		1.74	0.34		99		
Lactobacillus acidophilus		0.69	0.27		1		
07271020	**Good identification**						**07271020**
					SPO		
Lactobacillus jensenii		92.76	0.68	MLT 95	1		
Clostridium barati		4.19	0.40		99		
Lactobacillus catenaformis		1.31	0.36		1		
07271022	**Low discrimination**						**07271022**
					SPO		
Lactobacillus jensenii		46.05	0.65	MLT 95	1		
Lactobacillus acidophilus		43.06	0.61		1		
Clostridium barati		9.36	0.48		99		
07271030	**Very good identification**						**07271030**
					SPO		
Lactobacillus jensenii		99.73	0.60	MLT 95	1		
Clostridium barati		0.14	0.05		99		
Clostridium innocuum		0.05	0.00		99		
07271032	**Very good identification**						**07271032**
					SPO		
Lactobacillus jensenii		96.48	0.57	MLT 95	1		
Lactobacillus acidophilus		2.73	0.27		1		
Clostridium barati		0.59	0.14		99		
07271040	**Low discrimination**						**07271040**
					GRA	SPO	
Lactobacillus jensenii		80.23	0.44	MLT 95	99	1	
Clostridium innocuum		7.65	0.24	SOR 1	99	99	
Fusobacterium mortiferum		5.86	0.34		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07271041	**Low discrimination**						**07271041**
Actinomyces naeslundii		64.55	0.36	MLT 94			
Lactobacillus jensenii		35.45	0.10	RAF 92			
07271042	**Low discrimination**						**07271042**
					SPO		
Lactobacillus acidophilus		68.09	0.46	MLT 94	1		
Lactobacillus jensenii		23.80	0.41	LAC 92	1		
Clostridium barati		3.47	0.21	SOR 11	99		
07271050	**Good identification**						**07271050**
					SPO		
Lactobacillus jensenii		98.82	0.36	MLT 95	1		
Clostridium innocuum		1.18	0.00	SOR 1	99		
07271061	**Good identification**						**07271061**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07271062	**Low discrimination**						**07271062**
					SPO		
Lactobacillus acidophilus		80.09	0.46	MLT 94	1		
Clostridium barati		12.25	0.30	LAC 92	99		
Lactobacillus jensenii		7.00	0.31	SOR 11	1		
07271100	**Low discrimination**						**07271100**
					COC	GRA	
Fusobacterium mortiferum		60.09	0.75		1	1	
Atopobium parvulum		17.28	0.54		99	99	
Lactobacillus catenaformis		7.49	0.50		1	99	
07271101	**Low discrimination**						**07271101**
					COC	GRA	
Actinomyces naeslundii		51.44	0.57	MLT 94	1	99	
Fusobacterium mortiferum		29.53	0.41	RAF 92	1	1	
Atopobium parvulum		8.49	0.20		99	99	
07271102	**Very good identification**						**07271102**
					SPO		
Lactobacillus acidophilus		95.08	0.80	MLT 94	1		
Clostridium barati		1.80	0.48		99		
Lactobacillus catenaformis		1.13	0.50		1		
07271103	**Low discrimination**						**07271103**
					SPO		
Lactobacillus acidophilus		82.28	0.45	MLT 94	1		
Actinomyces naeslundii		13.62	0.57	URE 1	1		
Clostridium barati		1.56	0.14		99		
07271110	**Low discrimination**						**07271110**
					COC	GRA	
Lactobacillus jensenii		52.49	0.36	MLT 95	1	99	
Fusobacterium mortiferum		26.82	0.41	LAC 1	1	1	
Atopobium parvulum		7.71	0.20		99	99	
07271112	**Low discrimination**						**07271112**
					GRA		
Lactobacillus acidophilus		56.26	0.45	MLT 94	99		
Leptotrichia buccalis		33.08	0.32	MLZ 1	1		
Lactobacillus jensenii		6.98	0.33		99		
07271120	**Low discrimination**						**07271120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium barati		36.48	0.48	ESL 87	99		
Lactobacillus acidophilus		19.49	0.45	TRE 8	1		
Clostridium septicum		15.48	0.39		99		
07271121	**Low discrimination**						**07271121**
					SPO		
Actinomyces naeslundii		78.69	0.57	MLT 94	1		
Clostridium barati		8.99	0.14	RAF 92	99		
Lactobacillus acidophilus		4.80	0.11		1		
07271122	**Good identification**						**07271122**
					SPO		
Lactobacillus acidophilus		93.58	0.80	MLT 94	1		
Clostridium barati		5.31	0.56		99		
Clostridium ramosum		0.40	0.38		92		
07271123	**Low discrimination**						**07271123**
					SPO		
Lactobacillus acidophilus		81.34	0.45	MLT 94	1		
Actinomyces naeslundii		13.46	0.57	URE 1	1		
Clostridium barati		4.61	0.22		99		
07271132	**Low discrimination**						**07271132**
					GRA	SPO	
Lactobacillus acidophilus		83.03	0.45	MLT 94	99	1	
Leptotrichia buccalis		5.43	0.15	MLZ 1	1	1	
Clostridium barati		4.71	0.22		99	99	
07271141	**Good identification**						**07271141**
Actinomyces naeslundii		100.0	0.44	MLT 94			
07271142	**Very good identification**						**07271142**
					SPO		
Lactobacillus acidophilus		98.27	0.64	MLT 94	1		
Clostridium barati		1.31	0.30	SOR 11	99		
Actinomyces naeslundii		0.16	0.41		1		
07271161	**Good identification**						**07271161**
Actinomyces naeslundii		100.0	0.44	MLT 94			
07271162	**Very good identification**						**07271162**
					SPO		
Lactobacillus acidophilus		95.85	0.64	MLT 94	1		
Clostridium barati		3.83	0.38	SOR 11	99		
Actinomyces naeslundii		0.15	0.41		1		
07271200	**Low discrimination**						**07271200**
					GRA	SPO	
Lactobacillus jensenii		81.53	0.44	MLT 95	99	1	
Clostridium innocuum		7.77	0.24	RHA 1	99	99	
Fusobacterium mortiferum		5.95	0.34		1	1	
07271202	**Low discrimination**						**07271202**
					SPO		
Lactobacillus acidophilus		56.85	0.42	MLT 94	1		
Lactobacillus jensenii		32.64	0.41	LAC 92	1		
Clostridium barati		4.76	0.21	RHA 7	99		
07271210	**Good identification**						**07271210**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		98.82	0.36	MLT 95	1		
Clostridium innocuum		1.18	0.00	RHA 1	99		
07271222	**Low discrimination**						**07271222**
					SPO		
Lactobacillus acidophilus		71.10	0.42	MLT 94	1		
Clostridium barati		17.86	0.30	LAC 92	99		
Lactobacillus jensenii		10.21	0.31	RHA 7	1		
07271300	**Low discrimination**						**07271300**
					GRA	SPO	
Fusobacterium mortiferum		35.28	0.41	RHA 1	1	1	
Lactobacillus acidophilus		27.94	0.26		99	1	
Clostridium barati		20.14	0.21		99	99	
07271302	**Very good identification**						**07271302**
					SPO		
Lactobacillus acidophilus		97.39	0.60	MLT 94	1		
Clostridium barati		2.13	0.30	RHA 7	99		
Lactobacillus catenaformis		0.16	0.15		1		
07271322	**Low discrimination**						**07271322**
					SPO		
Lactobacillus acidophilus		87.35	0.60	MLT 94	1		
Clostridium ramosum		6.51	0.40	RHA 7	92		
Clostridium barati		5.72	0.38		99		
07271342	**Good identification**						**07271342**
					SPO		
Lactobacillus acidophilus		98.49	0.45	MLT 94	1		
Clostridium barati		1.51	0.11	RHA 7	99		
07271362	**Good identification**						**07271362**
					SPO		
Lactobacillus acidophilus		95.04	0.45	MLT 94	1		
Clostridium barati		4.38	0.20	RHA 7	99		
Clostridium ramosum		0.58	0.06	SOR 11	92		
07271400	**Low discrimination**						**07271400**
					GRA	SPO	
Clostridium innocuum		60.42	0.45	MAN 99	99	99	
Lactobacillus jensenii		36.28	0.44		99	1	
Fusobacterium mortiferum		2.65	0.34		1	1	
07271402	**Low discrimination**						**07271402**
					SPO		
Clostridium innocuum		66.05	0.45	MAN 99	99		
Lactobacillus jensenii		26.44	0.41		1		
Lactobacillus acidophilus		6.18	0.27		1		
07271410	**Low discrimination**						**07271410**
					SPO		
Lactobacillus jensenii		82.77	0.36	MLT 95	1		
Clostridium innocuum		17.23	0.22	ARA 1	99		
07271500	**Low discrimination**						**07271500**
					COC	GRA	
Fusobacterium mortiferum		54.10	0.41	ARA 1	1	1	
Atopobium parvulum		15.56	0.20		99	99	
Lactobacillus catenaformis		6.75	0.15		1	99	
07271502	**Acceptable identification**						**07271502**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus acidophilus		93.09	0.45	MLT 94	1			
Clostridium barati		1.76	0.14	ARA 1	99			
Lactobacillus catenaformis		1.10	0.15		1			
07271522	**Acceptable identification**							**07271522**
					GRA	SPO		
Lactobacillus acidophilus		93.56	0.45	MLT 94	99	1		
Clostridium barati		5.31	0.22	ARA 1	99	99		
Parabacteroides distasonis		0.46	0.17		1	1		
07272000	**Very good identification**							**07272000**
					GRA	SPO	CAT	
Clostridium innocuum		99.77	0.70	GAL 95	99	99	1	
Fusobacterium mortiferum		0.10	0.31		1	1	5	
Propionibacterium granulosum		0.09	0.28		99	1	90	
07272001	**Good identification**							**07272001**
					SPO			
Clostridium innocuum		98.33	0.36	GAL 95	99			
Actinomyces naeslundii		1.67	0.30	URE 1	1			
07272002	**Very good identification**							**07272002**
					SPO	CAT		
Clostridium innocuum		99.91	0.70	GAL 95	99	1		
Propionibacterium granulosum		0.02	0.18		1	90		
Lactobacillus acidophilus		0.02	0.05		1	1		
07272003	**Good identification**							**07272003**
					SPO			
Clostridium innocuum		98.33	0.36	GAL 95	99			
Actinomyces naeslundii		1.67	0.30	URE 1	1			
07272010	**Good identification**							**07272010**
					SPO	CAT		
Clostridium innocuum		98.78	0.47	GAL 95	99	1		
Clostridium difficile		0.75	0.07	MLZ 4	99	1		
Propionibacterium granulosum		0.30	0.13		1	90		
07272012	**Good identification**							**07272012**
Clostridium innocuum		100.0	0.47	GAL 95				
07272020	**Good identification**							**07272020**
					SPO	CAT		
Clostridium innocuum		95.70	0.36	GAL 95	99	1		
Propionibacterium granulosum		2.23	0.18	NAG 1	1	90		
Actinomyces naeslundii		1.08	0.27		1	25		
07272022	**Acceptable identification**							**07272022**
					SPO			
Clostridium innocuum		94.45	0.36	GAL 95	99			
Clostridium barati		2.92	0.08	NAG 1	99			
Lactobacillus acidophilus		1.56	0.05		1			
07272040	**Good identification**							**07272040**
Clostridium innocuum		100.0	0.36	GAL 95				
07272042	**Good identification**							**07272042**
Clostridium innocuum		100.0	0.36	GAL 95				
07272100	**Low discrimination**							**07272100**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		78.20	0.36	GAL 95	99	99	
Fusobacterium mortiferum		17.88	0.38	LAC 1	1	1	
Actinomyces naeslundii		2.52	0.35		99	1	
07272101	**Low discrimination**						**07272101**
					SPO		
Actinomyces naeslundii		82.71	0.38	MLT 94	1		
Clostridium innocuum		17.29	0.02	RAF 92	99		
07272102	**Low discrimination**						**07272102**
					SPO		
Clostridium innocuum		76.84	0.36	GAL 95	99		
Lactobacillus acidophilus		14.60	0.24	LAC 1	1		
Actinomyces naeslundii		2.48	0.35		1		
07272103	**Low discrimination**						**07272103**
					SPO		
Actinomyces naeslundii		82.71	0.38	MLT 94	1		
Clostridium innocuum		17.29	0.02	RAF 92	99		
07272121	**Good identification**						**07272121**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07272123	**Good identification**						**07272123**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07272200	**Good identification**						**07272200**
Clostridium innocuum		100.0	0.36	GAL 95			
07272202	**Good identification**						**07272202**
Clostridium innocuum		100.0	0.36	GAL 95			
07272400	**Very good identification**						**07272400**
Clostridium innocuum		100.0	0.57	GAL 95			
07272402	**Very good identification**						**07272402**
Clostridium innocuum		100.0	0.57	GAL 95			
07273000	**Excellent identification**						**07273000**
					SPO		
Clostridium innocuum		99.86	0.92		99		
Lactobacillus jensenii		0.11	0.44		1		
Actinomyces naeslundii		0.01	0.51		1		
07273001	**Very good identification**						**07273001**
					SPO		
Clostridium innocuum		97.80	0.58	URE 1	99		
Actinomyces naeslundii		2.10	0.54		1		
Lactobacillus jensenii		0.11	0.10		1		
07273002	**Excellent identification**						**07273002**
					SPO		
Clostridium innocuum		99.81	0.92		99		
Lactobacillus acidophilus		0.09	0.40		1		
Lactobacillus jensenii		0.07	0.41		1		
07273003	**Very good identification**						**07273003**
					SPO		
Clostridium innocuum		97.75	0.58	URE 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		2.10	0.54		1	
Lactobacillus acidophilus		0.09	0.05		1	
07273010	**Very good identification**					**07273010**
					SPO	
Clostridium innocuum		99.10	0.69	MLZ 4	99	
Lactobacillus jensenii		0.85	0.36		1	
Actinomyces naeslundii		0.02	0.29		1	
07273012	**Very good identification**					**07273012**
					SPO	
Clostridium innocuum		99.34	0.69	MLZ 4	99	
Lactobacillus jensenii		0.57	0.33		1	
Clostridium tertium		0.05	0.13		99	
07273020	**Good identification**					**07273020**
					SPO	
Clostridium innocuum		94.71	0.58	NAG 1	99	
Lactobacillus jensenii		2.51	0.34		1	
Actinomyces naeslundii		1.35	0.51		1	
07273021	**Low discrimination**					**07273021**
					SPO	
Actinomyces naeslundii		67.98	0.54	MLT 94	1	
Clostridium innocuum		32.03	0.24	RAF 92	99	
07273022	**Low discrimination**					**07273022**
					SPO	
Clostridium innocuum		86.39	0.58	NAG 1	99	
Lactobacillus acidophilus		7.44	0.40		1	
Clostridium barati		2.67	0.30		99	
07273023	**Low discrimination**					**07273023**
					SPO	
Actinomyces naeslundii		66.15	0.54	MLT 94	1	
Clostridium innocuum		31.17	0.24	RAF 92	99	
Lactobacillus acidophilus		2.68	0.05		1	
07273040	**Very good identification**					**07273040**
					SPO	
Clostridium innocuum		99.64	0.58	SOR 1	99	
Actinomyces naeslundii		0.25	0.38		1	
Lactobacillus jensenii		0.11	0.10		1	
07273042	**Very good identification**					**07273042**
					SPO	
Clostridium innocuum		98.50	0.58	SOR 1	99	
Lactobacillus acidophilus		1.05	0.24		1	
Actinomyces naeslundii		0.25	0.38		1	
07273061	**Good identification**					**07273061**
Actinomyces naeslundii		100.0	0.41	MLT 94		
07273063	**Good identification**					**07273063**
Actinomyces naeslundii		100.0	0.41	MLT 94		
07273100	**Low discrimination**					**07273100**
					SPO	
Clostridium innocuum		88.36	0.58	LAC 1	99	
Actinomyces naeslundii		3.60	0.58		1	
Clostridium tertium		3.52	0.35		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07273101	**Low discrimination**					**07273101**
					SPO	
Actinomyces naeslundii		85.31	0.61	MLT 94	1	
Clostridium innocuum		14.12	0.24	RAF 92	99	
Clostridium tertium		0.56	0.01		99	
07273102	**Low discrimination**					**07273102**
					SPO	
Clostridium innocuum		45.54	0.58	LAC 1	99	
Lactobacillus acidophilus		45.07	0.58		1	
Clostridium tertium		5.45	0.43		99	
07273103	**Low discrimination**					**07273103**
					SPO	
Actinomyces naeslundii		74.12	0.61	MLT 94	1	
Clostridium innocuum		12.27	0.24	RAF 92	99	
Lactobacillus acidophilus		12.14	0.24		1	
07273110	**Low discrimination**					**07273110**
					SPO	
Clostridium tertium		61.36	0.39	MLT 99	99	
Clostridium innocuum		36.06	0.34		99	
Actinomyces naeslundii		1.85	0.37		1	
07273111	**Low discrimination**					**07273111**
					SPO	
Actinomyces naeslundii		73.85	0.40	MLT 94	1	
Clostridium tertium		16.47	0.05	RAF 92	99	
Clostridium innocuum		9.68	0.00	MLZ 5	99	
07273112	**Low discrimination**					**07273112**
					SPO	
Clostridium tertium		79.26	0.47	MLT 99	99	
Clostridium innocuum		15.53	0.34		99	
Lactobacillus acidophilus		3.73	0.24		1	
07273113	**Low discrimination**					**07273113**
					SPO	
Actinomyces naeslundii		55.56	0.40	MLT 94	1	
Clostridium tertium		37.16	0.13	RAF 92	99	
Clostridium innocuum		7.28	0.00	MLZ 5	99	
07273120	**Low discrimination**					**07273120**
					SPO	
Clostridium tertium		36.58	0.43	MLT 99	99	
Clostridium ramosum		34.48	0.41		92	
Actinomyces naeslundii		12.44	0.58		1	
07273121	**Very good identification**					**07273121**
					SPO	
Actinomyces naeslundii		96.30	0.61	MLT 94	1	
Clostridium tertium		1.91	0.09	RAF 92	99	
Clostridium ramosum		1.80	0.07		92	
07273122	**Low discrimination**					**07273122**
					SPO	
Lactobacillus acidophilus		50.92	0.58	MLT 94	1	
Clostridium ramosum		23.21	0.51	MAN 5	92	
Clostridium tertium		18.47	0.51		99	
07273123	**Low discrimination**					**07273123**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		76.15	0.61	MLT 94	1	
Lactobacillus acidophilus		12.47	0.24	RAF 92	1	
Clostridium ramosum		5.69	0.17		92	
07273130	**Good identification**					**07273130**
					SPO	
Clostridium tertium		98.29	0.47	MLT 99	99	
Actinomyces naeslundii		0.99	0.37		1	
Clostridium ramosum		0.53	0.07		92	
07273131	**Low discrimination**					**07273131**
					SPO	
Actinomyces naeslundii		59.92	0.40	MLT 94	1	
Clostridium tertium		40.08	0.13	RAF 92	99	
07273132	**Good identification**					**07273132**
					SPO	
Clostridium tertium		97.25	0.55	MLT 99	99	
Lactobacillus acidophilus		1.52	0.24		1	
Clostridium ramosum		0.69	0.17		92	
07273141	**Good identification**					**07273141**
Actinomyces naeslundii		100.0	0.49	MLT 94		
07273142	**Low discrimination**					**07273142**
					SPO	
Lactobacillus acidophilus		84.15	0.42	MLT 94	1	
Clostridium innocuum		6.95	0.24	MAN 5	99	
Actinomyces naeslundii		4.94	0.46	SOR 11	1	
07273143	**Low discrimination**					**07273143**
Actinomyces naeslundii		89.71	0.49	MLT 94		
Lactobacillus acidophilus		10.29	0.08	RAF 92		
07273160	**Low discrimination**					**07273160**
					SPO	
Actinomyces naeslundii		53.37	0.46	MLT 94	1	
Clostridium barati		19.99	0.11	RAF 92	99	
Lactobacillus acidophilus		9.19	0.08		1	
07273161	**Good identification**					**07273161**
Actinomyces naeslundii		100.0	0.49	MLT 94		
07273162	**Low discrimination**					**07273162**
					SPO	
Lactobacillus acidophilus		83.93	0.42	MLT 94	1	
Clostridium barati		5.53	0.20	MAN 5	99	
Actinomyces naeslundii		4.93	0.46	SOR 11	1	
07273163	**Low discrimination**					**07273163**
Actinomyces naeslundii		89.71	0.49	MLT 94		
Lactobacillus acidophilus		10.29	0.08	RAF 92		
07273200	**Very good identification**					**07273200**
					SPO	
Clostridium innocuum		99.89	0.58	RHA 1	99	
Lactobacillus jensenii		0.11	0.10		1	
07273202	**Very good identification**					**07273202**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium innocuum		99.20	0.58	RHA 1	99	
Lactobacillus acidophilus		0.64	0.20		1	
Clostridium barati		0.09	0.03		99	
07273302	**Low discrimination**					**07273302**
					SPO	
Lactobacillus acidophilus		78.17	0.38	MLT 94	1	
Clostridium innocuum		10.60	0.24	MAN 5	99	
Clostridium ramosum		6.34	0.19	RHA 7	92	
07273320	**Good identification**					**07273320**
Clostridium ramosum		96.73	0.43	MLT 99		
Clostridium barati		1.74	0.11			
Clostridium tertium		0.78	0.09			
07273322	**Low discrimination**					**07273322**
					SPO	
Clostridium ramosum		87.35	0.53	MLT 99	92	
Lactobacillus acidophilus		10.88	0.38		1	
Clostridium barati		1.18	0.20		99	
07273400	**Very good identification**					**07273400**
					SPO	
Clostridium innocuum		99.99	0.79		99	
Lactobacillus jensenii		0.01	0.10		1	
Actinomyces israelii		0.00	0.18		1	
07273401	**Good identification**					**07273401**
Clostridium innocuum		100.0	0.45	URE 1		
07273402	**Very good identification**					**07273402**
					SPO	
Clostridium innocuum		99.99	0.79		99	
Actinomyces israelii		0.01	0.21		1	
Lactobacillus jensenii		0.00	0.07		1	
07273403	**Good identification**					**07273403**
Clostridium innocuum		100.0	0.45	URE 1		
07273410	**Good identification**					**07273410**
					SPO	
Clostridium innocuum		99.95	0.56	MLZ 4	99	
Lactobacillus jensenii		0.05	0.02		1	
07273412	**Good identification**					**07273412**
Clostridium innocuum		100.0	0.56	MLZ 4		
07273420	**Good identification**					**07273420**
Clostridium innocuum		100.0	0.45	NAG 1		
07273422	**Good identification**					**07273422**
					SPO	
Clostridium innocuum		99.51	0.45	NAG 1	99	
Lactobacillus acidophilus		0.49	0.05		1	
07273440	**Good identification**					**07273440**
Clostridium innocuum		100.0	0.45	SOR 1		
07273442	**Good identification**					**07273442**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium innocuum		99.79	0.45	SOR 1	99		
Actinomyces israelii		0.21	0.14		1		
07273500	**Good identification**						**07273500**
					SPO		
Clostridium innocuum		98.10	0.45	LAC 1	99		
Actinomyces israelii		1.45	0.29		1		
Actinomyces naeslundii		0.23	0.24		1		
07273502	**Acceptable identification**						**07273502**
					SPO		
Clostridium innocuum		91.30	0.45	LAC 1	99		
Lactobacillus acidophilus		5.17	0.24		1		
Actinomyces israelii		2.02	0.32		1		
07273600	**Good identification**						**07273600**
Clostridium innocuum		100.0	0.45	RHA 1			
07273602	**Good identification**						**07273602**
Clostridium innocuum		100.0	0.45	RHA 1			
07274000	**Acceptable identification**						**07274000**
					GRA	SPO	
Fusobacterium mortiferum		90.91	0.46	XYL 7	1	1	
Clostridium innocuum		6.55	0.14		99	99	
Lactobacillus jensenii		2.53	0.10		99	1	
07274100	**Acceptable identification**						**07274100**
					COC	GRA	
Fusobacterium mortiferum		90.29	0.53	XYL 7	1	1	
Atopobium parvulum		9.71	0.25		99	99	
07274102	**Low discrimination**						**07274102**
					COC	GRA	
Fusobacterium mortiferum		73.35	0.36	XYL 7	1	1	
Lactobacillus acidophilus		14.13	0.11	BGL 10	1	99	
Atopobium parvulum		12.53	0.12		99	99	
07275000	**Low discrimination**						**07275000**
					GRA	SPO	
Lactobacillus jensenii		46.20	0.44	MLT 95	99	1	
Fusobacterium mortiferum		25.12	0.49	XYL 1	1	1	
Clostridium innocuum		22.94	0.36		99	99	
07275002	**Low discrimination**						**07275002**
					SPO		
Lactobacillus jensenii		46.17	0.41	MLT 95	1		
Clostridium innocuum		34.39	0.36	XYL 1	99		
Lactobacillus acidophilus		10.79	0.27		1		
07275010	**Acceptable identification**						**07275010**
					GRA	SPO	
Lactobacillus jensenii		92.72	0.36	MLT 95	99	1	
Clostridium innocuum		5.76	0.13	XYL 1	99	99	
Fusobacterium mortiferum		1.53	0.15		1	1	
07275100	**Low discrimination**						**07275100**
					COC	GRA	
Atopobium parvulum		86.13	0.59	MLT 99	99	99	
Fusobacterium mortiferum		12.14	0.56		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.58	0.36		1	99	
07275102	**Low discrimination**						**07275102**
					COC	GRA	
Lactobacillus acidophilus		48.05	0.45	MLT 94	1	99	
Atopobium parvulum		42.60	0.46	XYL 1	99	99	
Fusobacterium mortiferum		3.78	0.39		1	1	
07275120	**Acceptable identification**						**07275120**
					COC	SPO	
Atopobium parvulum		93.62	0.46	MLT 99	99	1	
Clostridium barati		2.00	0.14		1	99	
Lactobacillus acidophilus		1.07	0.11		1	1	
07275122	**Low discrimination**						**07275122**
					COC	GRA	
Lactobacillus acidophilus		75.48	0.45	MLT 94	1	99	
Atopobium parvulum		11.81	0.33	XYL 1	99	99	
Parabacteroides distasonis		5.49	0.37		1	1	
07275322	**Low discrimination**						**07275322**
					SPO	GRA	
Parabacteroides distasonis		71.32	0.46	RAF 99	1	1	
Lactobacillus acidophilus		24.61	0.26		1	99	
Clostridium ramosum		1.84	0.06		92	50	
07275332	**Acceptable identification**						**07275332**
Parabacteroides distasonis		93.08	0.41	RAF 99			
Clostridium clostridioforme		6.92	0.16				
07275712	**Good identification**						**07275712**
Clostridium clostridioforme		100.0	0.36	MLT 99			
07275722	**Low discrimination**						**07275722**
Parabacteroides distasonis		81.67	0.42	RAF 99			
Clostridium clostridioforme		18.34	0.25				
07275732	**Low discrimination**						**07275732**
Clostridium clostridioforme		65.51	0.36	MLT 99			
Parabacteroides distasonis		34.49	0.37	RAF 94			
07276000	**Good identification**						**07276000**
					GRA	SPO	
Clostridium innocuum		99.86	0.48	GAL 95	99	99	
Fusobacterium mortiferum		0.14	0.12	XYL 5	1	1	
07276002	**Good identification**						**07276002**
Clostridium innocuum		100.0	0.48	GAL 95			
07276400	**Good identification**						**07276400**
Clostridium innocuum		100.0	0.36	GAL 95			
07276402	**Good identification**						**07276402**
Clostridium innocuum		100.0	0.36	GAL 95			
07277000	**Very good identification**						**07277000**
					SPO		
Clostridium innocuum		99.78	0.70	XYL 5	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.18	0.38		1	
Lactobacillus jensenii		0.02	0.10		1	
07277001	**Good identification**					**07277001**
Clostridium innocuum		100.0	0.36	XYL 5		
07277002	**Very good identification**					**07277002**
					SPO	
Clostridium innocuum		99.68	0.70	XYL 5	99	
Actinomyces israelii		0.26	0.41		1	
Clostridium tertium		0.03	0.11		99	
07277003	**Good identification**					**07277003**
Clostridium innocuum		100.0	0.36	XYL 5		
07277010	**Good identification**					**07277010**
					SPO	
Clostridium innocuum		98.77	0.47	XYL 5	99	
Actinomyces israelii		0.62	0.24	MLZ 4	1	
Clostridium tertium		0.45	0.07		99	
07277012	**Good identification**					**07277012**
					SPO	
Clostridium innocuum		97.74	0.47	XYL 5	99	
Clostridium tertium		1.34	0.15	MLZ 4	99	
Actinomyces israelii		0.92	0.27		1	
07277020	**Good identification**					**07277020**
					SPO	
Clostridium innocuum		95.90	0.36	XYL 5	99	
Clostridium tertium		3.04	0.11	NAG 1	99	
Actinomyces israelii		1.06	0.18		1	
07277022	**Low discrimination**					**07277022**
					SPO	
Clostridium innocuum		88.63	0.36	XYL 5	99	
Clostridium tertium		8.43	0.19	NAG 1	99	
Actinomyces israelii		1.47	0.21		1	
07277040	**Acceptable identification**					**07277040**
					SPO	
Clostridium innocuum		93.67	0.36	XYL 5	99	
Actinomyces israelii		6.33	0.31	SOR 1	1	
07277042	**Acceptable identification**					**07277042**
					SPO	
Clostridium innocuum		90.80	0.36	XYL 5	99	
Actinomyces israelii		9.20	0.34	SOR 1	1	
07277100	**Low discrimination**					**07277100**
					SPO	
Clostridium tertium		35.25	0.37	MLT 99	99	
Clostridium innocuum		33.68	0.36		99	
Actinomyces israelii		23.40	0.48		1	
07277102	**Low discrimination**					**07277102**
					SPO	
Clostridium tertium		57.75	0.45	MLT 99	99	
Actinomyces israelii		19.17	0.51		1	
Clostridium innocuum		18.40	0.36		99	
07277110	**Acceptable identification**					**07277110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium tertium		92.75	0.41	MLT 99	99		
Actinomyces israelii		5.18	0.34		1		
Clostridium innocuum		2.08	0.13		99		
07277112	**Good identification**						**07277112**
					SPO		
Clostridium tertium		96.08	0.50	MLT 99	99		
Actinomyces israelii		2.68	0.37		1		
Clostridium innocuum		0.72	0.13		99		
07277120	**Good identification**						**07277120**
					COC	SPO	
Clostridium tertium		96.43	0.45	MLT 99	1	99	
Actinomyces israelii		1.36	0.28		1	1	
Atopobium parvulum		0.99	0.12		99	1	
07277122	**Good identification**						**07277122**
					SPO		
Clostridium tertium		96.13	0.53	MLT 99	99		
Lactobacillus acidophilus		1.94	0.24		1		
Clostridium ramosum		0.88	0.17		92		
07277130	**Good identification**						**07277130**
					SPO		
Clostridium tertium		99.88	0.50	MLT 99	99		
Actinomyces israelii		0.12	0.14		1		
07277132	**Very good identification**						**07277132**
					SPO		
Clostridium tertium		99.94	0.58	MLT 99	99		
Actinomyces israelii		0.06	0.17		1		
07277140	**Acceptable identification**						**07277140**
					SPO		
Actinomyces israelii		92.89	0.41	MLT 99	1		
Clostridium tertium		3.63	0.03		99		
Clostridium innocuum		3.47	0.02		99		
07277142	**Low discrimination**						**07277142**
					SPO		
Actinomyces israelii		86.13	0.44	MLT 99	1		
Clostridium tertium		6.74	0.11		99		
Lactobacillus acidophilus		4.99	0.08		1		
07277200	**Good identification**						**07277200**
					SPO		
Clostridium innocuum		98.71	0.36	XYL 5	99		
Actinomyces israelii		1.29	0.19	RHA 1	1		
07277202	**Good identification**						**07277202**
					SPO		
Clostridium innocuum		98.08	0.36	XYL 5	99		
Actinomyces israelii		1.92	0.22	RHA 1	1		
07277400	**Very good identification**						**07277400**
					SPO		
Clostridium innocuum		99.19	0.57	XYL 5	99		
Actinomyces israelii		0.81	0.37		1		
07277402	**Very good identification**						**07277402**
					SPO		
Clostridium innocuum		98.80	0.57	XYL 5	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		1.21	0.40		1			
07277500	**Low discrimination**							**07277500**
					SPO			
Actinomyces israelii		66.57	0.47	MLT 99	1			
Clostridium innocuum		20.67	0.23		99			
Clostridium butyricum		11.53	0.13		99			
07277502	**Low discrimination**							**07277502**
					SPO			
Actinomyces israelii		67.48	0.50	MLT 99	1			
Clostridium innocuum		13.97	0.23		99			
Bifidobacterium adolescentis		13.02	0.31		1			
07277540	**Good identification**							**07277540**
					SPO			
Actinomyces israelii		96.27	0.40	MLT 99	1			
Clostridium butyricum		3.73	-.05		99			
07277542	**Acceptable identification**							**07277542**
Actinomyces israelii		94.22	0.43	MLT 99				
Bifidobacterium adolescentis		5.78	0.16					
07300000	**Good identification**							**07300000**
					COC	GRA	SPO	
Gemella morbillorum		93.75	0.64	SAL 1	99	99	1	
Prevotella disiens		3.70	0.32		1	1	1	
Clostridium bifermentans		0.96	0.27		1	99	99	
07300020	**Good identification**							**07300020**
					COC	GRA	SPO	
Gemella morbillorum		96.80	0.48	SAL 1	99	99	1	
Clostridium botulinum biovar B		0.95	0.08	NAG 10	1	99	99	
Prevotella bivia		0.71	-.03		1	1	1	
07300040	**Acceptable identification**							**07300040**
					COC	SPO		
Clostridium botulinum biovar B		94.45	0.47	SAL 1	1	99		
Gemella morbillorum		4.37	0.30		99	1		
Clostridium botulinum biovar A		0.58	0.12		1	99		
07300050	**Good identification**							**07300050**
Clostridium botulinum biovar B		99.80	0.42	SAL 1				
Clostridium novyi biovar A		0.20	0.11					
07300060	**Good identification**							**07300060**
					COC	SPO		
Clostridium botulinum biovar B		98.97	0.42	SAL 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
07300070	**Good identification**							**07300070**
Clostridium botulinum biovar B		100.0	0.37	SAL 1				
07300400	**Good identification**							**07300400**
					COC	GRA		
Gemella morbillorum		99.64	0.48	SAL 1	99	99		
Prevotella disiens		0.36	-.03	ARA 10	1	1		
07301000	**Good identification**							**07301000**
					COC			
Gemella morbillorum		92.98	0.60	SAL 1	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		3.55	0.36		1		
Lactobacillus catenaformis		1.36	0.29		1		
07301003	**Low discrimination**						**07301003**
					SPO		
Actinomyces naeslundii		80.87	0.40	RAF 92	1		
Clostridium chauvoei		19.13	0.07	FRU 95	99		
07301020	**Low discrimination**						**07301020**
					COC		
Gemella morbillorum		64.13	0.43	SAL 1	99		
Clostridium perfringens		25.57	0.36	NAG 10	1		
Lactobacillus jensenii		5.51	0.26		1		
07301021	**Low discrimination**						**07301021**
					COC	SPO	
Actinomyces naeslundii		53.86	0.40	RAF 92	1	1	
Clostridium chauvoei		24.06	0.11	FRU 95	1	99	
Gemella morbillorum		15.79	0.09		99	1	
07301023	**Good identification**						**07301023**
Actinomyces naeslundii		100.0	0.40	RAF 92			
07301100	**Low discrimination**						**07301100**
					COC	GRA	
Fusobacterium mortiferum		27.78	0.42	MLT 11	1	1	
Lactobacillus catenaformis		25.12	0.32	FRU 99	1	99	
Gemella morbillorum		11.58	0.26		99	99	
07301101	**Good identification**						**07301101**
					SPO		
Clostridium chauvoei		97.90	0.54	SAL 1	99		
Actinomyces naeslundii		2.10	0.48		1		
07301103	**Low discrimination**						**07301103**
					SPO		
Clostridium chauvoei		88.66	0.41	SAL 1	99		
Actinomyces naeslundii		10.78	0.48		1		
Actinomyces viscosus		0.56	0.02		1		
07301120	**Low discrimination**						**07301120**
					GRA		
Clostridium perfringens		72.29	0.51	FRU 99	99		
Prevotella melaninogenica		16.75	0.33		1		
Prevotella bivia		5.91	0.19		1		
07301121	**Acceptable identification**						**07301121**
					SPO		
Clostridium chauvoei		91.89	0.45	SAL 1	99		
Actinomyces naeslundii		5.91	0.48		1		
Clostridium perfringens		1.78	0.17		84		
07301122	**Low discrimination**						**07301122**
					GRA		
Clostridium perfringens		70.66	0.45	FRU 99	99		
Prevotella oralis		7.62	0.25		1		
Prevotella melaninogenica		6.63	0.21		1		
07301143	**Low discrimination**						**07301143**
					SPO		
Actinomyces naeslundii		67.99	0.35	RAF 92	1		
Clostridium chauvoei		32.01	0.07	FRU 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07301160	**Good identification**					**07301160**
					GRA	
Clostridium perfringens		96.81	0.39	FRU 99	99	
Actinomyces naeslundii		2.01	0.32		99	
Prevotella melaninogenica		1.19	-.01		1	
07301161	**Low discrimination**					**07301161**
					SPO	
Actinomyces naeslundii		45.16	0.35	RAF 92	1	
Clostridium chauvoei		40.17	0.11	FRU 95	99	
Clostridium perfringens		14.68	0.04		84	
07301163	**Good identification**					**07301163**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07301400	**Good identification**					**07301400**
					COC	
Gemella morbillorum		99.13	0.43	SAL 1	99	
Lactobacillus fermentum		0.52	0.16	ARA 10	1	
Lactobacillus jensenii		0.34	0.02		1	
07303001	**Good identification**					**07303001**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07303002	**Low discrimination**					**07303002**
					SPO	
Actinomyces naeslundii		75.17	0.42	RAF 92	1	
Clostridium innocuum		24.83	0.10	FRU 95	99	
07303003	**Good identification**					**07303003**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07303020	**Low discrimination**					**07303020**
					COC	
Actinomyces naeslundii		75.94	0.42	RAF 92	1	
Gemella morbillorum		17.20	0.09	FRU 95	99	
Clostridium perfringens		6.86	0.02		1	
07303021	**Good identification**					**07303021**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07303022	**Good identification**					**07303022**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07303023	**Good identification**					**07303023**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07303100	**Good identification**					**07303100**
Actinomyces naeslundii		97.59	0.50	RAF 92		
Propionibacterium propionicum		2.41	0.12	FRU 95		
07303101	**Low discrimination**					**07303101**
					SPO	
Actinomyces naeslundii		80.48	0.53	RAF 92	1	
Clostridium chauvoei		19.52	0.19	FRU 95	99	
07303102	**Acceptable identification**					**07303102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		93.18	0.50	RAF 92		
Propionibacterium propionicum		3.45	0.15	FRU 95		
Lactobacillus acidophilus		3.38	0.01			
07303103	**Good identification**					**07303103**
					SPO	
Actinomyces naeslundii		95.90	0.53	RAF 92	1	
Clostridium chauvoei		4.11	0.07	FRU 95	99	
07303120	**Low discrimination**					**07303120**
					GRA	
Actinomyces naeslundii		77.72	0.50	RAF 92	99	
Clostridium perfringens		18.09	0.17	FRU 95	99	
Prevotella melaninogenica		4.19	-.01		1	
07303121	**Acceptable identification**					**07303121**
					SPO	
Actinomyces naeslundii		92.52	0.53	RAF 92	1	
Clostridium chauvoei		7.48	0.11	FRU 95	99	
07303122	**Low discrimination**					**07303122**
					SPO	
Actinomyces naeslundii		82.14	0.50	RAF 92	1	
Clostridium perfringens		9.00	0.11	FRU 95	84	
Clostridium barati		3.35	-.01		99	
07303123	**Good identification**					**07303123**
Actinomyces naeslundii		100.0	0.53	RAF 92		
07303140	**Low discrimination**					**07303140**
Actinomyces naeslundii		87.74	0.37	RAF 92		
Propionibacterium propionicum		12.26	0.12	FRU 95		
07303141	**Good identification**					**07303141**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07303142	**Low discrimination**					**07303142**
Actinomyces naeslundii		82.67	0.37	RAF 92		
Propionibacterium propionicum		17.33	0.15	FRU 95		
07303143	**Good identification**					**07303143**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07303160	**Low discrimination**					**07303160**
Actinomyces naeslundii		79.93	0.37	RAF 92		
Clostridium perfringens		20.08	0.04	FRU 95		
07303161	**Good identification**					**07303161**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07303162	**Good identification**					**07303162**
Actinomyces naeslundii		100.0	0.37	RAF 92		
07303163	**Good identification**					**07303163**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07304000	**Good identification**					**07304000**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Gemella morbillorum		98.14	0.48	SAL 1	99	99		
Actinomyces odontolyticus		1.26	0.20	XYL 10	1	99		
Prevotella disiens		0.35	-.03		1	1		
07305000	**Good identification**							**07305000**
					COC	GRA		
Gemella morbillorum		97.13	0.43	SAL 1	99	99		
Actinomyces odontolyticus		1.55	0.17	XYL 10	1	99		
Fusobacterium mortiferum		0.68	0.17		1	1		
07310000	**Low discrimination**							**07310000**
					COC	GRA		
Gemella morbillorum		65.30	0.66	SAL 1	99	99		
Fusobacterium mortiferum		19.72	0.67		1	1		
Prevotella intermedia		5.31	0.46		1	1		
07310002	**Low discrimination**							**07310002**
					GRA			
Fusobacterium mortiferum		31.03	0.51	MLT 11	1			
Actinomyces odontolyticus		15.49	0.40	BGL 10	99			
Actinomyces viscosus		12.04	0.31		99			
07310020	**Low discrimination**							**07310020**
					COC	CAT		
Gemella morbillorum		68.29	0.50	SAL 1	99	1		
Clostridium perfringens		12.00	0.36	NAG 10	1	1		
Propionibacterium granulosum		7.06	0.40		1	90		
07310021	**Low discrimination**							**07310021**
					COC			
Actinomyces naeslundii		50.30	0.39	RAF 92	1			
Gemella morbillorum		42.28	0.15	GAL 96	99			
Clostridium perfringens		7.43	0.02		1			
07310023	**Good identification**							**07310023**
Actinomyces naeslundii		100.0	0.39	RAF 92				
07310040	**Very good identification**							**07310040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		97.16	0.58	SAL 1	1	99	99	
Gemella morbillorum		1.37	0.32		99	99	1	
Megasphaera elsdenii		0.76	0.29		99	1	1	
07310050	**Good identification**							**07310050**
Clostridium botulinum biovar B		100.0	0.52	SAL 1				
07310060	**Good identification**							**07310060**
					COC	SPO		
Clostridium botulinum biovar B		98.60	0.52	SAL 1	1	99		
Clostridium perfringens		1.04	0.24		1	84		
Gemella morbillorum		0.31	0.15		99	1		
07310070	**Good identification**							**07310070**
Clostridium botulinum biovar B		100.0	0.47	SAL 1				
07310100	**Good identification**							**07310100**
					COC	GRA		
Fusobacterium mortiferum		94.07	0.74	MLT 11	1	1		
Gemella morbillorum		1.35	0.32		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		1.29	0.32		1	99	
07310101	**Low discrimination**						**07310101**
					GRA	SPO	
Fusobacterium mortiferum		43.96	0.39	MLT 11	1	1	
Clostridium chauvoei		32.57	0.26	URE 1	99	99	
Actinomyces naeslundii		23.47	0.46		99	1	
07310102	**Low discrimination**						**07310102**
					GRA		
Fusobacterium mortiferum		59.16	0.57	MLT 11	1		
Actinomyces viscosus		17.49	0.36	BGL 10	99		
Lactobacillus catenaformis		7.29	0.32		99		
07310110	**Good identification**						**07310110**
Fusobacterium mortiferum		100.0	0.39	MLT 11			
07310120	**Low discrimination**						**07310120**
					GRA		
Clostridium perfringens		81.44	0.51	GAL 99	99		
Prevotella bivia		4.20	0.15		1		
Fusobacterium mortiferum		4.05	0.39		1		
07310121	**Low discrimination**						**07310121**
					SPO		
Actinomyces naeslundii		54.26	0.46	RAF 92	1		
Clostridium chauvoei		25.10	0.18	GAL 96	99		
Clostridium perfringens		20.64	0.17		84		
07310122	**Low discrimination**						**07310122**
					GRA	SPO	
Clostridium perfringens		65.36	0.45	GAL 99	99	84	
Clostridium barati		14.72	0.30		99	99	
Prevotella oralis		7.04	0.25		1	1	
07310123	**Low discrimination**						**07310123**
					SPO		
Actinomyces naeslundii		76.87	0.46	RAF 92	1		
Clostridium perfringens		13.76	0.11	GAL 96	84		
Clostridium chauvoei		6.27	0.05		99		
07310160	**Low discrimination**						**07310160**
					SPO		
Clostridium perfringens		85.98	0.39	GAL 99	84		
Clostridium botulinum biovar B		11.22	0.18		99		
Actinomyces naeslundii		1.41	0.30		1		
07310200	**Low discrimination**						**07310200**
					COC	GRA	
Actinomyces odontolyticus		76.95	0.49	NIT 99	1	99	
Gemella morbillorum		14.66	0.32	MNS 9	99	99	
Fusobacterium mortiferum		4.43	0.33		1	1	
07310202	**Good identification**						**07310202**
					GRA	CAT	
Actinomyces odontolyticus		97.91	0.45	NIT 99	99	10	
Propionibacterium granulosum		1.07	0.14	MNS 9	99	90	
Fusobacterium mortiferum		1.02	0.17		1	5	
07310220	**Low discrimination**						**07310220**
					COC	CAT	
Actinomyces odontolyticus		83.75	0.35	NIT 99	1	10	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		10.89	0.15	MNS 9	99	1	
Propionibacterium granulosum		3.45	0.14	NAG 14	1	90	
07310300	**Low discrimination**						**07310300**
					GRA		
Actinomyces odontolyticus		70.18	0.40	NIT 99	99		
Fusobacterium mortiferum		29.82	0.39	MNS 9	1		
07310302	**Low discrimination**						**07310302**
					GRA	SPO	
Actinomyces odontolyticus		84.03	0.36	NIT 99	99	1	
Fusobacterium mortiferum		6.47	0.23	MNS 9	1	1	
Clostridium barati		3.60	0.04		99	99	
07310400	**Good identification**						**07310400**
					COC	GRA	
Gemella morbillorum		95.24	0.50	SAL 1	99	99	
Fusobacterium mortiferum		2.61	0.33	ARA 10	1	1	
Prevotella intermedia		0.70	0.12		1	1	
07310500	**Low discrimination**						**07310500**
					COC	GRA	
Fusobacterium mortiferum		75.54	0.39	MLT 11	1	1	
Gemella morbillorum		11.91	0.15	ARA 1	99	99	
Actinomyces meyeri		8.07	0.13		1	99	
07311000	**Low discrimination**						**07311000**
					COC		
Lactobacillus jensenii		46.80	0.70	CEL 95	1		
Lactobacillus catenaformis		17.89	0.63		1		
Gemella morbillorum		15.12	0.61		99		
07311001	**Low discrimination**						**07311001**
Actinomyces naeslundii		43.94	0.62	RAF 92			
Lactobacillus jensenii		23.08	0.36				
Lactobacillus catenaformis		8.82	0.29				
07311002	**Low discrimination**						**07311002**
					GRA		
Lactobacillus jensenii		56.03	0.67	CEL 95	99		
Lactobacillus catenaformis		32.13	0.63		99		
Fusobacterium mortiferum		2.54	0.54		1		
07311003	**Low discrimination**						**07311003**
Actinomyces naeslundii		58.22	0.62	RAF 92			
Lactobacillus jensenii		20.39	0.33				
Lactobacillus catenaformis		11.69	0.29				
07311010	**Very good identification**						**07311010**
					CAT		
Lactobacillus jensenii		95.58	0.62	CEL 95	1		
Lactobacillus catenaformis		1.11	0.29		1		
Propionibacterium granulosum		1.08	0.35		90		
07311012	**Very good identification**						**07311012**
					CAT		
Lactobacillus jensenii		97.15	0.59	CEL 95	1		
Lactobacillus catenaformis		1.69	0.29		1		
Propionibacterium granulosum		0.41	0.25		90		
07311020	**Low discrimination**						**07311020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		72.71	0.70	LAC 88		
Lactobacillus jensenii		15.68	0.60			
Lactobacillus catenaformis		5.99	0.53			
07311021	**Low discrimination**					**07311021**
Actinomyces naeslundii		53.35	0.62	RAF 92		
Clostridium perfringens		32.49	0.36			
Lactobacillus jensenii		7.01	0.26			
07311022	**Low discrimination**					**07311022**
Clostridium perfringens		58.18	0.65	LAC 88		
Lactobacillus jensenii		17.77	0.57			
Lactobacillus catenaformis		10.19	0.53			
07311023	**Low discrimination**					**07311023**
Actinomyces naeslundii		67.10	0.62	RAF 92		
Clostridium perfringens		19.23	0.31			
Lactobacillus jensenii		5.88	0.23			
07311030	**Low discrimination**					**07311030**
Lactobacillus jensenii		84.88	0.52	CEL 95		
Clostridium perfringens		11.93	0.36			
Lactobacillus catenaformis		0.98	0.18			
07311031	**Low discrimination**					**07311031**
Actinomyces naeslundii		51.32	0.40	RAF 92		
Lactobacillus jensenii		42.68	0.17	MLZ 5		
Clostridium perfringens		6.00	0.02			
07311032	**Low discrimination**					**07311032**
Lactobacillus jensenii		86.77	0.49	CEL 95		
Clostridium perfringens		8.61	0.31			
Lactobacillus catenaformis		1.51	0.18			
07311033	**Low discrimination**					**07311033**
Actinomyces naeslundii		64.33	0.40	RAF 92		
Lactobacillus jensenii		35.67	0.14	MLZ 5		
07311040	**Low discrimination**					**07311040**
					SPO	
Lactobacillus jensenii		28.87	0.36	CEL 95	1	
Clostridium perfringens		19.52	0.32	SOR 1	84	
Clostridium botulinum biovar B		12.39	0.24		99	
07311041	**Good identification**					**07311041**
Actinomyces naeslundii		97.08	0.49	RAF 92		
Lactobacillus jensenii		2.92	0.02			
07311043	**Good identification**					**07311043**
Actinomyces naeslundii		100.0	0.49	RAF 92		
07311060	**Very good identification**					**07311060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		95.70	0.58	LAC 88			
Lactobacillus jensenii		1.09	0.26				
Actinomyces naeslundii		0.98	0.46				
07311061	**Low discrimination**						**07311061**
Actinomyces naeslundii		60.34	0.49	RAF 92			
Clostridium perfringens		39.66	0.24				
07311062	**Low discrimination**						**07311062**
					SPO		
Clostridium perfringens		88.93	0.52	LAC 88	84		
Clostridium barati		4.29	0.25		99		
Lactobacillus acidophilus		2.34	0.23		1		
07311063	**Low discrimination**						**07311063**
Actinomyces naeslundii		76.37	0.49	RAF 92			
Clostridium perfringens		23.63	0.18				
07311100	**Low discrimination**						**07311100**
					GRA		
Fusobacterium mortiferum		37.49	0.77	MLT 11	1		
Lactobacillus catenaformis		33.90	0.66		99		
Clostridium perfringens		15.54	0.59		99		
07311101	**Low discrimination**						**07311101**
					GRA	SPO	
Clostridium chauvoei		81.48	0.60	SAL 1	99	99	
Actinomyces naeslundii		14.23	0.70		99	1	
Fusobacterium mortiferum		1.67	0.42		1	1	
07311102	**Low discrimination**						**07311102**
Lactobacillus catenaformis		45.85	0.66	CEL 99			
Lactobacillus acidophilus		20.32	0.57				
Clostridium perfringens		9.90	0.54				
07311103	**Low discrimination**						**07311103**
					SPO		
Clostridium chauvoei		44.29	0.47	SAL 1	99		
Actinomyces naeslundii		43.84	0.70		1		
Lactobacillus catenaformis		4.64	0.32		1		
07311110	**Low discrimination**						**07311110**
					GRA		
Fusobacterium mortiferum		29.59	0.42	MLT 11	1		
Lactobacillus catenaformis		26.75	0.32	MLZ 1	99		
Lactobacillus jensenii		15.55	0.28		99		
07311113	**Low discrimination**						**07311113**
					SPO		
Actinomyces naeslundii		83.76	0.48	RAF 92	1		
Clostridium chauvoei		16.24	0.13	MLZ 5	99		
07311120	**Good identification**						**07311120**
					GRA		
Clostridium perfringens		93.60	0.85		99		
Prevotella melaninogenica		1.97	0.50		1		
Lactobacillus catenaformis		1.58	0.56		99		
07311121	**Low discrimination**						**07311121**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		41.74	0.52	SAL 1	99	
Clostridium perfringens		34.33	0.51		84	
Actinomyces naeslundii		21.88	0.70		1	
07311122	**Low discrimination**					**07311122**
					GRA	
Clostridium perfringens		77.15	0.80		99	
Prevotella oralis		8.31	0.60		1	
Lactobacillus acidophilus		4.90	0.57		99	
07311123	**Low discrimination**					**07311123**
					SPO	
Actinomyces naeslundii		43.79	0.70	RAF 92	1	
Clostridium perfringens		32.33	0.45		84	
Clostridium chauvoei		14.75	0.39		99	
07311130	**Acceptable identification**					**07311130**
					GRA	
Clostridium perfringens		91.58	0.51	MLZ 1	99	
Actinomyces naeslundii		2.05	0.45		99	
Prevotella melaninogenica		1.93	0.15		1	
07311131	**Low discrimination**					**07311131**
					SPO	
Actinomyces naeslundii		59.97	0.48	RAF 92	1	
Clostridium chauvoei		21.97	0.18	MLZ 5	99	
Clostridium perfringens		18.06	0.17		84	
07311132	**Low discrimination**					**07311132**
					GRA	
Clostridium perfringens		73.51	0.45	MLZ 1	99	
Prevotella oralis		7.92	0.25		1	
Lactobacillus acidophilus		4.67	0.23		99	
07311133	**Low discrimination**					**07311133**
					SPO	
Actinomyces naeslundii		82.90	0.48	RAF 92	1	
Clostridium perfringens		11.75	0.11	MLZ 5	84	
Clostridium chauvoei		5.36	0.05		99	
07311140	**Low discrimination**					**07311140**
					GRA	
Clostridium perfringens		69.23	0.47	NAG 97	99	
Actinomyces naeslundii		8.90	0.54		99	
Fusobacterium mortiferum		8.86	0.42		1	
07311141	**Low discrimination**					**07311141**
					SPO	
Actinomyces naeslundii		72.46	0.57	RAF 92	1	
Clostridium chauvoei		23.74	0.26		99	
Clostridium perfringens		3.80	0.13		84	
07311142	**Low discrimination**					**07311142**
Lactobacillus acidophilus		41.93	0.41	CEL 99		
Clostridium perfringens		31.46	0.41	SOR 11		
Actinomyces naeslundii		8.59	0.54			
07311143	**Low discrimination**					**07311143**
					SPO	
Actinomyces naeslundii		89.66	0.57	RAF 92	1	
Clostridium chauvoei		5.18	0.13		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		2.95	0.07		1	
07311151	**Good identification**					**07311151**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07311153	**Good identification**					**07311153**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07311160	**Very good identification**					**07311160**
					SPO	
Clostridium perfringens		98.91	0.73		84	
Actinomyces naeslundii		0.39	0.54		1	
Clostridium barati		0.31	0.25		99	
07311161	**Low discrimination**					**07311161**
					SPO	
Clostridium perfringens		60.43	0.39	URE 1	84	
Actinomyces naeslundii		35.68	0.57		1	
Clostridium chauvoei		3.90	0.18		99	
07311162	**Good identification**					**07311162**
					SPO	
Clostridium perfringens		92.62	0.67		84	
Lactobacillus acidophilus		3.82	0.41		1	
Clostridium barati		1.83	0.34		99	
07311163	**Low discrimination**					**07311163**
Actinomyces naeslundii		53.62	0.57	RAF 92		
Clostridium perfringens		42.74	0.33			
Lactobacillus acidophilus		1.76	0.07			
07311170	**Good identification**					**07311170**
Clostridium perfringens		97.97	0.39	MLZ 1		
Actinomyces naeslundii		2.03	0.32			
07311171	**Low discrimination**					**07311171**
Actinomyces naeslundii		75.47	0.35	RAF 92		
Clostridium perfringens		24.53	0.04	MLZ 5		
07311173	**Good identification**					**07311173**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07311200	**Low discrimination**					**07311200**
Actinomyces odontolyticus		48.51	0.46	NIT 99		
Lactobacillus jensenii		23.10	0.36	MNS 9		
Lactobacillus catenaformis		8.83	0.29			
07311202	**Low discrimination**					**07311202**
Actinomyces odontolyticus		45.58	0.42	NIT 99		
Lactobacillus jensenii		23.61	0.33	MNS 9		
Lactobacillus catenaformis		13.54	0.29			
07311220	**Low discrimination**					**07311220**
Clostridium perfringens		54.09	0.36	LAC 88		
Actinomyces odontolyticus		15.95	0.32	RHA 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		11.66	0.26				
07311300	**Low discrimination**						**07311300**
					GRA		
Actinomyces odontolyticus		21.43	0.37	NIT 99	99		
Fusobacterium mortiferum		20.50	0.42	MNS 9	1		
Lactobacillus catenaformis		18.53	0.32		99		
07311302	**Low discrimination**						**07311302**
					GRA		
Lactobacillus acidophilus		39.92	0.38	CEL 99	99		
Prevotella oralis		20.30	0.31	RHA 7	1		
Lactobacillus catenaformis		12.09	0.32		99		
07311320	**Low discrimination**						**07311320**
					GRA	SPO	
Clostridium perfringens		78.69	0.51	RHA 1	99	84	
Prevotella oralis		8.91	0.31		1	1	
Clostridium barati		4.59	0.25		99	99	
07311322	**Low discrimination**						**07311322**
					GRA		
Prevotella oralis		69.17	0.53	RAF 92	1		
Clostridium perfringens		15.13	0.45		99		
Lactobacillus acidophilus		7.16	0.38		99		
07311360	**Good identification**						**07311360**
					GRA	SPO	
Clostridium perfringens		96.84	0.39	RHA 1	99	84	
Clostridium barati		2.58	0.07		99	99	
Prevotella oralis		0.58	-.03		1	1	
07311400	**Low discrimination**						**07311400**
					COC		
Gemella morbillorum		51.63	0.45	SAL 1	99		
Lactobacillus fermentum		22.12	0.50	ARA 10	1		
Lactobacillus jensenii		14.53	0.36		1		
07311402	**Low discrimination**						**07311402**
Lactobacillus fermentum		55.37	0.50	SAL 1			
Lactobacillus jensenii		24.24	0.33				
Lactobacillus catenaformis		13.90	0.29				
07311420	**Low discrimination**						**07311420**
					COC		
Clostridium perfringens		47.53	0.36	LAC 88	1		
Lactobacillus fermentum		20.81	0.42	ARA 1	1		
Gemella morbillorum		16.19	0.29		99		
07311500	**Low discrimination**						**07311500**
					GRA		
Lactobacillus fermentum		57.32	0.54	SAL 1	99		
Fusobacterium mortiferum		14.63	0.42		1		
Lactobacillus catenaformis		13.23	0.32		99		
07311502	**Low discrimination**						**07311502**
Bifidobacterium adolescentis		45.37	0.50	RAF 92			
Lactobacillus fermentum		36.03	0.54	XYL 87			
Lactobacillus catenaformis		8.32	0.32				
07311512	**Good identification**						**07311512**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium adolescentis		96.32	0.52	RAF 92				
Bifidobacterium longum subsp. longum		2.04	0.22	XYL 87				
Clostridium clostridioforme		1.01	0.15					
07311520	**Low discrimination**							**07311520**
Clostridium perfringens		83.08	0.51	ARA 1				
Lactobacillus fermentum		8.09	0.46					
Bifidobacterium longum subsp. longum		1.86	0.28					
07311522	**Low discrimination**							**07311522**
					GRA			
Clostridium perfringens		63.83	0.45	ARA 1	99			
Lactobacillus fermentum		13.21	0.46		99			
Prevotella oralis		6.88	0.25		1			
07311523	**Low discrimination**							**07311523**
					SPO			
Actinomyces naeslundii		48.19	0.36	RAF 92	1			
Clostridium perfringens		35.59	0.11	ARA 1	84			
Clostridium chauvoei		16.23	0.05		99			
07311552	**Good identification**							**07311552**
Bifidobacterium adolescentis		100.0	0.36	RAF 92				
07311560	**Good identification**							**07311560**
Clostridium perfringens		100.0	0.39	ARA 1				
07312000	**Low discrimination**							**07312000**
					COC	GRA	CAT	
Propionibacterium granulosum		56.88	0.48	SAL 1	1	99	90	
Gemella morbillorum		16.58	0.32		99	99	1	
Megasphaera elsdenii		13.68	0.32		99	1	1	
07312001	**Low discrimination**							**07312001**
Actinomyces naeslundii		88.10	0.43	RAF 92				
Propionibacterium granulosum		11.90	0.14	GAL 96				
07312002	**Low discrimination**							**07312002**
					COC	GRA		
Propionibacterium granulosum		67.81	0.38	SAL 1	1	99		
Actinomyces naeslundii		13.52	0.40		1	99		
Megasphaera elsdenii		7.25	0.15		99	1		
07312003	**Good identification**							**07312003**
Actinomyces naeslundii		100.0	0.43	RAF 92				
07312020	**Low discrimination**							**07312020**
					COC	CAT		
Propionibacterium granulosum		73.98	0.38	SAL 1	1	90		
Actinomyces naeslundii		14.75	0.40		1	25		
Gemella morbillorum		9.58	0.15		99	1		
07312021	**Good identification**							**07312021**
Actinomyces naeslundii		100.0	0.43	RAF 92				
07312023	**Good identification**							**07312023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	RAF 92		
07312100	**Low discrimination**					**07312100**
					GRA	
Propionibacterium propionicum		55.05	0.46	NIT 86	99	
Fusobacterium mortiferum		25.83	0.39	SAL 1	1	
Actinomyces naeslundii		17.85	0.48		99	
07312101	**Low discrimination**					**07312101**
Actinomyces naeslundii		81.40	0.51	RAF 92		
Propionibacterium propionicum		18.60	0.30	GAL 96		
07312102	**Low discrimination**					**07312102**
					GRA	
Propionibacterium propionicum		77.49	0.49	NIT 86	99	
Actinomyces naeslundii		16.75	0.48	SAL 1	99	
Fusobacterium mortiferum		2.69	0.23		1	
07312103	**Low discrimination**					**07312103**
Actinomyces naeslundii		74.48	0.51	RAF 92		
Propionibacterium propionicum		25.52	0.33	GAL 96		
07312120	**Low discrimination**					**07312120**
					SPO	
Actinomyces naeslundii		70.70	0.48	RAF 92	1	
Clostridium perfringens		20.78	0.17	GAL 96	84	
Clostridium barati		6.32	0.04		99	
07312121	**Good identification**					**07312121**
Actinomyces naeslundii		100.0	0.51	RAF 92		
07312122	**Low discrimination**					**07312122**
					SPO	
Actinomyces naeslundii		65.01	0.48	RAF 92	1	
Clostridium barati		17.44	0.12	GAL 96	99	
Clostridium perfringens		8.99	0.11		84	
07312123	**Good identification**					**07312123**
Actinomyces naeslundii		100.0	0.51	RAF 92		
07312140	**Acceptable identification**					**07312140**
Propionibacterium propionicum		94.59	0.46	NIT 86		
Actinomyces naeslundii		5.41	0.35	SAL 1		
07312142	**Good identification**					**07312142**
Propionibacterium propionicum		96.33	0.49	NIT 86		
Actinomyces naeslundii		3.68	0.35	SAL 1		
07312160	**Low discrimination**					**07312160**
Actinomyces naeslundii		66.95	0.35	RAF 92		
Clostridium perfringens		21.24	0.04	GAL 96		
Propionibacterium propionicum		11.81	0.12			
07312161	**Good identification**					**07312161**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07312162	**Low discrimination**					**07312162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		79.07	0.35	RAF 92		
Propionibacterium propionicum		20.93	0.15	GAL 96		
07312163	**Good identification**					**07312163**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07313000	**Low discrimination**					**07313000**
					CAT	
Actinomyces naeslundii		32.70	0.64	RAF 92	25	
Propionibacterium granulosum		27.33	0.48		90	
Lactobacillus jensenii		13.28	0.36		1	
07313001	**Very good identification**					**07313001**
					CAT	
Actinomyces naeslundii		99.17	0.67	RAF 92	25	
Propionibacterium granulosum		0.56	0.14		90	
Lactobacillus jensenii		0.27	0.02		1	
07313002	**Low discrimination**					**07313002**
					SPO	
Actinomyces naeslundii		46.79	0.64	RAF 92	1	
Clostridium innocuum		15.46	0.32		99	
Lactobacillus jensenii		12.66	0.33		1	
07313003	**Very good identification**					**07313003**
Actinomyces naeslundii		100.0	0.67	RAF 92		
07313011	**Good identification**					**07313011**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07313013	**Good identification**					**07313013**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07313020	**Low discrimination**					**07313020**
					CAT	
Actinomyces naeslundii		52.59	0.64	RAF 92	25	
Clostridium perfringens		24.75	0.36		1	
Propionibacterium granulosum		10.99	0.38		90	
07313021	**Very good identification**					**07313021**
Actinomyces naeslundii		99.68	0.67	RAF 92		
Clostridium perfringens		0.32	0.02			
07313022	**Low discrimination**					**07313022**
					SPO	
Actinomyces naeslundii		60.82	0.64	RAF 92	1	
Clostridium perfringens		13.47	0.31		84	
Clostridium barati		12.25	0.25		99	
07313023	**Very good identification**					**07313023**
Actinomyces naeslundii		100.0	0.67	RAF 92		
07313030	**Low discrimination**					**07313030**
					CAT	
Actinomyces naeslundii		43.97	0.42	RAF 92	25	
Lactobacillus jensenii		28.26	0.17	MLZ 5	1	
Propionibacterium granulosum		23.80	0.23		90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07313031	**Good identification**					**07313031**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07313032	**Low discrimination**					**07313032**
					SPO	
Actinomyces naeslundii		58.13	0.42	RAF 92	1	
Lactobacillus jensenii		24.91	0.14	MLZ 5	1	
Clostridium tertium		9.09	0.05		99	
07313033	**Good identification**					**07313033**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07313040	**Low discrimination**					**07313040**
					CAT	
Actinomyces naeslundii		88.35	0.51	RAF 92	25	
Propionibacterium granulosum		4.23	0.14		90	
Actinomyces israelii		3.52	0.18		1	
07313041	**Good identification**					**07313041**
Actinomyces naeslundii		100.0	0.54	RAF 92		
07313042	**Low discrimination**					**07313042**
					SPO	
Actinomyces naeslundii		86.43	0.51	RAF 92	1	
Actinomyces israelii		5.17	0.21		1	
Clostridium barati		2.86	-.01		99	
07313043	**Good identification**					**07313043**
Actinomyces naeslundii		100.0	0.54	RAF 92		
07313060	**Low discrimination**					**07313060**
					SPO	
Actinomyces naeslundii		64.89	0.51	RAF 92	1	
Clostridium perfringens		32.96	0.24		84	
Clostridium barati		2.15	-.01		99	
07313061	**Good identification**					**07313061**
Actinomyces naeslundii		100.0	0.54	RAF 92		
07313062	**Low discrimination**					**07313062**
					SPO	
Actinomyces naeslundii		72.94	0.51	RAF 92	1	
Clostridium perfringens		17.44	0.18		84	
Clostridium barati		7.24	0.07		99	
07313063	**Good identification**					**07313063**
Actinomyces naeslundii		100.0	0.54	RAF 92		
07313100	**Low discrimination**					**07313100**
					GRA	
Actinomyces naeslundii		69.93	0.72	RAF 92	99	
Propionibacterium propionicum		8.99	0.46		99	
Fusobacterium mortiferum		6.33	0.42		1	
07313101	**Very good identification**					**07313101**
					SPO	
Actinomyces naeslundii		96.22	0.75	RAF 92	1	
Clostridium chauvoei		2.87	0.26		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		0.92	0.30		1	
07313102	**Low discrimination**					**07313102**
Actinomyces naeslundii		58.02	0.72	RAF 92		
Propionibacterium propionicum		11.19	0.49			
Lactobacillus acidophilus		10.96	0.35			
07313103	**Very good identification**					**07313103**
					SPO	
Actinomyces naeslundii		97.96	0.75	RAF 92	1	
Propionibacterium propionicum		1.40	0.33		1	
Clostridium chauvoei		0.52	0.13		99	
07313110	**Low discrimination**					**07313110**
					SPO	
Actinomyces naeslundii		58.79	0.50	RAF 92	1	
Clostridium tertium		35.52	0.23	MLZ 5	99	
Actinomyces israelii		4.24	0.21		1	
07313111	**Good identification**					**07313111**
Actinomyces naeslundii		100.0	0.53	RAF 92		
07313113	**Good identification**					**07313113**
Actinomyces naeslundii		100.0	0.53	RAF 92		
07313120	**Low discrimination**					**07313120**
					SPO	
Clostridium perfringens		49.79	0.51	MAN 1	84	
Actinomyces naeslundii		41.06	0.72		1	
Clostridium barati		2.91	0.25		99	
07313121	**Very good identification**					**07313121**
					SPO	
Actinomyces naeslundii		98.22	0.75	RAF 92	1	
Clostridium chauvoei		0.98	0.18		99	
Clostridium perfringens		0.80	0.17		84	
07313122	**Low discrimination**					**07313122**
					SPO	
Actinomyces naeslundii		42.46	0.72	RAF 92	1	
Clostridium perfringens		24.23	0.45		84	
Clostridium barati		9.02	0.34		99	
07313123	**Very good identification**					**07313123**
					SPO	
Actinomyces naeslundii		99.07	0.75	RAF 92	1	
Clostridium perfringens		0.38	0.11		84	
Clostridium chauvoei		0.17	0.05		99	
07313131	**Good identification**					**07313131**
Actinomyces naeslundii		100.0	0.53	RAF 92		
07313132	**Low discrimination**					**07313132**
					SPO	
Clostridium tertium		81.70	0.39	TRE 90	99	
Actinomyces naeslundii		15.02	0.50	CEL 99	1	
Clostridium perfringens		1.65	0.11		84	
07313133	**Good identification**					**07313133**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.47	0.53	RAF 92	1	
Clostridium tertium		3.53	0.05	MLZ 5	99	
07313140	**Low discrimination**					**07313140**
Actinomyces naeslundii		54.46	0.59	RAF 92		
Propionibacterium propionicum		39.66	0.46			
Actinomyces israelii		3.05	0.28			
07313141	**Good identification**					**07313141**
Actinomyces naeslundii		94.88	0.62	RAF 92		
Propionibacterium propionicum		5.12	0.30			
07313142	**Low discrimination**					**07313142**
Propionibacterium propionicum		45.50	0.49	NIT 86		
Actinomyces naeslundii		41.65	0.59	SAL 1		
Lactobacillus acidophilus		5.51	0.19			
07313143	**Good identification**					**07313143**
Actinomyces naeslundii		92.52	0.62	RAF 92		
Propionibacterium propionicum		7.49	0.33			
07313150	**Low discrimination**					**07313150**
Actinomyces naeslundii		77.00	0.37	RAF 92		
Actinomyces israelii		12.25	0.14	MLZ 5		
Propionibacterium propionicum		10.76	0.12			
07313151	**Good identification**					**07313151**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07313152	**Low discrimination**					**07313152**
Actinomyces naeslundii		42.82	0.37	RAF 92		
Bifidobacterium adolescentis		37.99	0.21	MLZ 5		
Actinomyces israelii		10.22	0.17			
07313153	**Good identification**					**07313153**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07313160	**Low discrimination**					**07313160**
					SPO	
Clostridium perfringens		55.67	0.39	MAN 1	84	
Actinomyces naeslundii		42.54	0.59		1	
Clostridium barati		1.48	0.07		99	
07313161	**Very good identification**					**07313161**
Actinomyces naeslundii		99.13	0.62	RAF 92		
Clostridium perfringens		0.87	0.04			
07313162	**Low discrimination**					**07313162**
Actinomyces naeslundii		53.12	0.59	RAF 92		
Clostridium perfringens		32.72	0.33			
Lactobacillus acidophilus		7.03	0.19			
07313163	**Very good identification**					**07313163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.62	RAF 92		
07313170	**Low discrimination**					**07313170**
Actinomyces naeslundii		79.93	0.37	RAF 92		
Clostridium perfringens		20.08	0.04	MLZ 5		
07313171	**Good identification**					**07313171**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07313172	**Low discrimination**					**07313172**
					SPO	
Actinomyces naeslundii		66.98	0.37	RAF 92	1	
Clostridium tertium		20.85	0.05	MLZ 5	99	
Bifidobacterium breve		12.17	0.10		1	
07313173	**Good identification**					**07313173**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07313300	**Low discrimination**					**07313300**
					SPO	
Actinomyces naeslundii		65.74	0.38	RAF 92	1	
Clostridium barati		13.35	-.01	RHA 1	99	
Actinomyces israelii		12.46	0.16		1	
07313301	**Good identification**					**07313301**
Actinomyces naeslundii		100.0	0.41	RAF 92		
07313303	**Good identification**					**07313303**
Actinomyces naeslundii		100.0	0.41	RAF 92		
07313321	**Good identification**					**07313321**
Actinomyces naeslundii		100.0	0.41	RAF 92		
07313323	**Acceptable identification**					**07313323**
					SPO	
Actinomyces naeslundii		90.69	0.41	RAF 92	1	
Clostridium ramosum		9.32	-.02	RHA 1	92	
07313501	**Good identification**					**07313501**
Actinomyces naeslundii		100.0	0.41	RAF 92		
07313502	**Low discrimination**					**07313502**
Bifidobacterium adolescentis		88.22	0.49	RAF 92		
Actinomyces israelii		7.54	0.37	XYL 87		
Actinomyces naeslundii		2.44	0.38			
07313503	**Low discrimination**					**07313503**
Actinomyces naeslundii		80.43	0.41	RAF 92		
Bifidobacterium adolescentis		19.57	0.15	ARA 1		
07313512	**Good identification**					**07313512**
Bifidobacterium adolescentis		98.97	0.50	RAF 92		
Actinomyces israelii		1.03	0.23	XYL 87		
07313521	**Good identification**					**07313521**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	RAF 92			
07313522	**Low discrimination**						**07313522**
Actinomyces naeslundii		37.10	0.38	RAF 92			
Clostridium perfringens		21.17	0.11	ARA 1			
Bifidobacterium adolescentis		13.54	0.15				
07313523	**Good identification**						**07313523**
Actinomyces naeslundii		100.0	0.41	RAF 92			
07314000	**Low discrimination**						**07314000**
					COC	GRA	
Gemella morbillorum		73.73	0.50	SAL 1	99	99	
Fusobacterium mortiferum		15.08	0.48	XYL 10	1	1	
Actinomyces odontolyticus		8.93	0.38		1	99	
07314100	**Low discrimination**						**07314100**
					GRA		
Fusobacterium mortiferum		79.72	0.54	MLT 11	1		
Actinomyces meyeri		8.87	0.29	XYL 7	99		
Actinomyces odontolyticus		6.39	0.30		99		
07314102	**Low discrimination**						**07314102**
					GRA		
Fusobacterium mortiferum		55.11	0.38	MLT 11	1		
Actinomyces odontolyticus		24.36	0.26	XYL 7	99		
Bifidobacterium longum subsp. infantis		6.10	0.04	BGL 10	99		
07314200	**Good identification**						**07314200**
					COC	GRA	
Actinomyces odontolyticus		95.08	0.43	NIT 99	1	99	
Gemella morbillorum		4.09	0.15	MNS 9	99	99	
Fusobacterium mortiferum		0.84	0.14		1	1	
07314202	**Good identification**						**07314202**
Actinomyces odontolyticus		100.0	0.40	NIT 99			
07314712	**Good identification**						**07314712**
Clostridium clostridioforme		100.0	0.39	RAF 94			
07314732	**Good identification**						**07314732**
Clostridium clostridioforme		100.0	0.39	RAF 94			
07315000	**Low discrimination**						**07315000**
					COC	GRA	
Gemella morbillorum		40.15	0.45	SAL 1	99	99	
Fusobacterium mortiferum		22.88	0.51	XYL 10	1	1	
Lactobacillus jensenii		11.30	0.36		1	99	
07315002	**Low discrimination**						**07315002**
Lactobacillus fermentum		31.42	0.46	SAL 1			
Lactobacillus jensenii		23.43	0.33				
Lactobacillus catenaformis		13.43	0.29				
07315020	**Low discrimination**						**07315020**
					COC		
Clostridium perfringens		48.44	0.36	XYL 1	1		
Gemella morbillorum		16.50	0.29	LAC 88	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		12.45	0.38		1		
07315100	**Low discrimination**						**07315100**
					COC	GRA	
Atopobium parvulum		55.71	0.53	TRE 99	99	99	
Fusobacterium mortiferum		22.74	0.57		1	1	
Lactobacillus fermentum		7.02	0.50		1	99	
07315102	**Low discrimination**						**07315102**
					COC		
Bifidobacterium longum subsp. infantis		29.34	0.38	ESL 99	1		
Bifidobacterium adolescentis		23.93	0.50	RAF 95	1		
Atopobium parvulum		15.63	0.40		99		
07315112	**Good identification**						**07315112**
Bifidobacterium adolescentis		95.26	0.52	RAF 92			
Actinomyces israelii		2.02	0.29	ARA 87			
Bifidobacterium longum subsp. infantis		0.97	0.04				
07315120	**Low discrimination**						**07315120**
					COC		
Clostridium perfringens		63.70	0.51	XYL 1	1		
Atopobium parvulum		15.31	0.40		99		
Bifidobacterium longum subsp. infantis		5.05	0.25		1		
07315122	**Low discrimination**						**07315122**
					GRA		
Bifidobacterium longum subsp. infantis		37.21	0.43	ESL 99	99		
Prevotella oralis		28.79	0.44	RAF 95	1		
Clostridium perfringens		21.83	0.45		99		
07315132	**Low discrimination**						**07315132**
					GRA		
Parabacteroides distasonis		53.47	0.36	RAF 99	1		
Bifidobacterium longum subsp. infantis		12.10	0.08		99		
Prevotella oralis		9.36	0.10		1		
07315152	**Acceptable identification**						**07315152**
Bifidobacterium adolescentis		93.74	0.36	RAF 92			
Actinomyces israelii		6.26	0.22	ARA 87			
07315160	**Good identification**						**07315160**
					COC		
Clostridium perfringens		98.20	0.39	XYL 1	1		
Atopobium parvulum		1.25	0.06		99		
Actinomyces israelii		0.55	0.13		1		
07315200	**Low discrimination**						**07315200**
					COC		
Actinomyces odontolyticus		62.37	0.40	NIT 99	1		
Eubacterium contortum		33.08	0.29	MNS 9	1		
Gemella morbillorum		2.17	0.11		99		
07315202	**Low discrimination**						**07315202**
Actinomyces odontolyticus		83.15	0.37	NIT 99			
Eubacterium contortum		12.70	0.16	MNS 9			
Clostridium clostridioforme		1.63	0.06				
07315300	**Very good identification**						**07315300**
					COC		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium contortum		98.93	0.63	ESL 99	1		
Actinomyces odontolyticus		0.60	0.32		1		
Atopobium parvulum		0.21	0.18		99		
07315302	**Acceptable identification**						**07315302**
					GRA		
Eubacterium contortum		94.63	0.50	ESL 99	99		
Actinomyces odontolyticus		1.98	0.28		99		
Prevotella oralis		1.18	0.16		1		
07315320	**Low discrimination**						**07315320**
					GRA		
Eubacterium contortum		63.26	0.50	ESL 99	99		
Bacteroides vulgatus		34.79	0.38		1		
Prevotella oralis		0.79	0.16		1		
07315322	**Low discrimination**						**07315322**
					GRA		
Prevotella oralis		38.20	0.38	RAF 92	1		
Parabacteroides distasonis		28.64	0.48	XYL 11	1		
Eubacterium contortum		28.58	0.37		99		
07315332	**Low discrimination**						**07315332**
Parabacteroides distasonis		79.93	0.44	RAF 99			
Clostridium clostridioforme		16.57	0.26				
Prevotella oralis		2.00	0.04				
07315400	**Low discrimination**						**07315400**
					COC		
Lactobacillus fermentum		62.68	0.46	SAL 1	1		
Gemella morbillorum		27.68	0.29		99		
Actinomyces israelii		5.19	0.29		1		
07315402	**Low discrimination**						**07315402**
Lactobacillus fermentum		71.93	0.46	SAL 1			
Bifidobacterium adolescentis		17.01	0.30				
Actinomyces israelii		8.93	0.32				
07315420	**Low discrimination**						**07315420**
					COC	GRA	
Lactobacillus fermentum		68.96	0.38	SAL 1	1	99	
Bacteroides vulgatus		18.18	0.07		1	1	
Gemella morbillorum		10.15	0.12		99	99	
07315422	**Acceptable identification**						**07315422**
Lactobacillus fermentum		91.83	0.38	SAL 1			
Clostridium clostridioforme		8.17	0.12				
07315500	**Low discrimination**						**07315500**
Lactobacillus fermentum		48.36	0.50	SAL 1			
Eubacterium contortum		14.88	0.27				
Actinomyces israelii		9.81	0.39				
07315502	**Good identification**						**07315502**
Bifidobacterium adolescentis		90.08	0.64	RAF 92			
Lactobacillus fermentum		6.28	0.50				
Actinomyces israelii		1.91	0.42				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07315512	**Very good identification**					**07315512**
Bifidobacterium adolescentis		97.23	0.66	RAF 92		
Clostridium clostridioforme		1.54	0.32			
Bifidobacterium longum subsp. longum		0.92	0.30			
07315520	**Low discrimination**					**07315520**
					GRA	
Bacteroides vulgatus		85.88	0.41	RAF 97	1	
Bifidobacterium longum subsp. longum		6.30	0.36	RHA 90	99	
Lactobacillus fermentum		5.37	0.42		99	
07315522	**Low discrimination**					**07315522**
					GRA	
Bifidobacterium longum subsp. longum		28.72	0.36	SAL 1	99	
Lactobacillus fermentum		24.47	0.42	RAF 97	99	
Parabacteroides distasonis		17.43	0.37		1	
07315530	**Acceptable identification**					**07315530**
					GRA	
Bifidobacterium longum subsp. longum		92.52	0.40	SAL 1	99	
Bacteroides vulgatus		7.48	0.07	RAF 97	1	
07315532	**Low discrimination**					**07315532**
Bifidobacterium longum subsp. longum		53.36	0.40	SAL 1		
Clostridium clostridioforme		22.20	0.32	RAF 97		
Bifidobacterium adolescentis		14.19	0.32			
07315542	**Acceptable identification**					**07315542**
Bifidobacterium adolescentis		93.25	0.49	RAF 92		
Actinomyces israelii		6.22	0.35	SOR 11		
Lactobacillus fermentum		0.53	0.16			
07315552	**Good identification**					**07315552**
Bifidobacterium adolescentis		99.19	0.50	RAF 92		
Actinomyces israelii		0.81	0.21	SOR 11		
07315612	**Good identification**					**07315612**
Clostridium clostridioforme		100.0	0.38	RAF 94		
07315632	**Good identification**					**07315632**
Clostridium clostridioforme		100.0	0.38	RAF 94		
07315700	**Very good identification**					**07315700**
					GRA	
Eubacterium contortum		98.35	0.61	ESL 99	99	
Bacteroides vulgatus		1.57	0.23		1	
Actinomyces israelii		0.05	0.20		99	
07315702	**Low discrimination**					**07315702**
Eubacterium contortum		89.49	0.48	ESL 99		
Clostridium clostridioforme		7.43	0.36			
Bifidobacterium adolescentis		2.41	0.30			
07315712	**Low discrimination**					**07315712**
Clostridium clostridioforme		89.78	0.47	RAF 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		7.83	0.32	TRE 88			
Eubacterium contortum		2.40	0.14				
07315720	**Low discrimination**						**07315720**
					GRA		
Bacteroides vulgatus		89.92	0.58	RAF 97	1		
Eubacterium contortum		10.07	0.48		99		
Parabacteroides distasonis		0.01	0.11		1		
07315722	**Low discrimination**						**07315722**
					GRA		
Eubacterium contortum		36.38	0.36	ESL 99	99		
Parabacteroides distasonis		27.32	0.45		1		
Bacteroides vulgatus		18.60	0.23		1		
07315732	**Low discrimination**						**07315732**
					GRA		
Clostridium clostridioforme		83.56	0.47	RAF 94	50		
Parabacteroides distasonis		15.77	0.40	TRE 88	1		
Eubacterium contortum		0.39	0.01		99		
07317000	**Low discrimination**						**07317000**
					COC	SPO	
Actinomyces israelii		75.76	0.43	CEL 93	1	1	
Clostridium innocuum		6.33	0.10		1	99	
Gemella morbillorum		5.31	0.11		99	1	
07317002	**Low discrimination**						**07317002**
					SPO		
Actinomyces israelii		88.23	0.46	CEL 93	1		
Clostridium innocuum		4.92	0.10		99		
Bifidobacterium adolescentis		2.97	0.15		1		
07317040	**Good identification**						**07317040**
Actinomyces israelii		100.0	0.36	CEL 93			
07317042	**Good identification**						**07317042**
Actinomyces israelii		100.0	0.39	CEL 93			
07317100	**Low discrimination**						**07317100**
					COC	SPO	
Actinomyces israelii		81.73	0.54	CEL 93	1	1	
Clostridium tertium		6.91	0.21		1	99	
Atopobium parvulum		4.66	0.18		99	1	
07317101	**Low discrimination**						**07317101**
Actinomyces naeslundii		83.69	0.41	RAF 92			
Actinomyces israelii		16.31	0.19	XYL 1			
07317102	**Low discrimination**						**07317102**
					SPO		
Actinomyces israelii		48.61	0.57	CEL 93	1		
Bifidobacterium adolescentis		40.46	0.49		1		
Clostridium tertium		8.21	0.29		99		
07317103	**Low discrimination**						**07317103**
Actinomyces naeslundii		65.11	0.41	RAF 92			
Actinomyces israelii		19.04	0.22	XYL 1			
Bifidobacterium adolescentis		15.85	0.15				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07317112	**Low discrimination**					**07317112**
					SPO	
Bifidobacterium adolescentis		69.34	0.50	RAF 92	1	
Clostridium tertium		20.48	0.33	ARA 87	99	
Actinomyces israelii		10.19	0.42		1	
07317121	**Good identification**					**07317121**
Actinomyces naeslundii		100.0	0.41	RAF 92		
07317122	**Low discrimination**					**07317122**
					SPO	
Clostridium tertium		76.31	0.37	TRE 90	99	
Actinomyces israelii		9.61	0.36	CEL 99	1	
Actinomyces naeslundii		3.47	0.38		1	
07317123	**Low discrimination**					**07317123**
					SPO	
Actinomyces naeslundii		87.09	0.41	RAF 92	1	
Clostridium tertium		12.91	0.03	XYL 1	99	
07317132	**Good identification**					**07317132**
					SPO	
Clostridium tertium		97.85	0.41	TRE 90	99	
Bifidobacterium adolescentis		1.12	0.16	CEL 99	1	
Actinomyces israelii		1.04	0.22		1	
07317140	**Good identification**					**07317140**
Actinomyces israelii		97.36	0.46	CEL 93		
Actinomyces naeslundii		1.53	0.25			
Propionibacterium propionicum		1.11	0.12			
07317142	**Low discrimination**					**07317142**
Actinomyces israelii		77.74	0.49	CEL 93		
Bifidobacterium adolescentis		20.56	0.33			
Propionibacterium propionicum		0.89	0.15			
07317302	**Low discrimination**					**07317302**
Actinomyces israelii		65.37	0.37	CEL 93		
Eubacterium contortum		27.33	0.16	RHA 7		
Bifidobacterium adolescentis		7.30	0.15			
07317400	**Good identification**					**07317400**
Actinomyces israelii		97.89	0.42	CEL 93		
Lactobacillus fermentum		2.11	0.12			
07317402	**Low discrimination**					**07317402**
Actinomyces israelii		77.56	0.45	CEL 93		
Bifidobacterium adolescentis		21.33	0.29			
Lactobacillus fermentum		1.11	0.12			
07317442	**Acceptable identification**					**07317442**
Actinomyces israelii		91.96	0.38	CEL 93		
Bifidobacterium adolescentis		8.04	0.13			
07317500	**Low discrimination**					**07317500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		89.49	0.52	CEL 93		
Bifidobacterium adolescentis		9.23	0.29			
Lactobacillus fermentum		0.79	0.16			
07317502	**Low discrimination**					**07317502**
Bifidobacterium adolescentis		87.13	0.63	RAF 92		
Actinomyces israelii		12.80	0.55			
Lactobacillus fermentum		0.08	0.16			
07317510	**Low discrimination**					**07317510**
Actinomyces israelii		54.24	0.38	CEL 93		
Bifidobacterium adolescentis		45.76	0.30	MLZ 13		
07317512	**Very good identification**					**07317512**
					SPO	
Bifidobacterium adolescentis		98.19	0.64	RAF 92	1	
Actinomyces israelii		1.76	0.41		1	
Clostridium tertium		0.04	-.01		99	
07317540	**Good identification**					**07317540**
Actinomyces israelii		96.83	0.45	CEL 93		
Bifidobacterium adolescentis		3.17	0.13			
07317542	**Low discrimination**					**07317542**
Bifidobacterium adolescentis		68.39	0.47	RAF 92		
Actinomyces israelii		31.61	0.48	SOR 11		
07317552	**Acceptable identification**					**07317552**
Bifidobacterium adolescentis		94.65	0.49	RAF 92		
Actinomyces israelii		5.35	0.34	SOR 11		
07317702	**Low discrimination**					**07317702**
Actinomyces israelii		42.89	0.36	CEL 93		
Bifidobacterium adolescentis		39.18	0.29	RHA 7		
Eubacterium contortum		17.94	0.14			
07320040	**Good identification**					**07320040**
Clostridium botulinum biovar B		99.95	0.56	SAL 1		
Clostridium botulinum biovar A		0.05	0.02			
07320050	**Good identification**					**07320050**
Clostridium botulinum biovar B		100.0	0.50	SAL 1		
07320060	**Good identification**					**07320060**
Clostridium botulinum biovar B		100.0	0.50	SAL 1		
07320070	**Good identification**					**07320070**
Clostridium botulinum biovar B		100.0	0.45	SAL 1		
07321000	**Low discrimination**					**07321000**
					COC	
Lactobacillus jensenii		81.36	0.44	FRU 99	1	
Gemella morbillorum		7.17	0.26	CEL 95	99	
Actinomyces naeslundii		3.52	0.42		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07321001	**Low discrimination**						**07321001**
					SPO		
Actinomyces naeslundii		81.13	0.45	RAF 92	1		
Lactobacillus jensenii		12.64	0.10	FRU 95	1		
Clostridium chauvoei		6.23	0.03		99		
07321002	**Low discrimination**						**07321002**
Lactobacillus jensenii		87.88	0.41	FRU 99			
Actinomyces naeslundii		5.70	0.42	CEL 95			
Lactobacillus acidophilus		3.25	0.14				
07321003	**Acceptable identification**						**07321003**
Actinomyces naeslundii		90.59	0.45	RAF 92			
Lactobacillus jensenii		9.41	0.07	FRU 95			
07321010	**Good identification**						**07321010**
					CAT		
Lactobacillus jensenii		98.89	0.36	FRU 99	1		
Propionibacterium granulosum		1.12	0.09	CEL 95	90		
07321020	**Low discrimination**						**07321020**
Clostridium perfringens		69.59	0.41	FRU 99			
Lactobacillus jensenii		24.24	0.34	LAC 88			
Actinomyces naeslundii		4.19	0.42				
07321021	**Low discrimination**						**07321021**
Actinomyces naeslundii		89.94	0.45	RAF 92			
Clostridium perfringens		10.06	0.07	FRU 95			
07321022	**Low discrimination**						**07321022**
Clostridium perfringens		59.02	0.35	FRU 99			
Lactobacillus jensenii		29.12	0.31	LAC 88			
Actinomyces naeslundii		7.55	0.42				
07321023	**Good identification**						**07321023**
Actinomyces naeslundii		95.00	0.45	RAF 92			
Clostridium perfringens		5.00	0.01	FRU 95			
07321100	**Low discrimination**						**07321100**
					COC		
Atopobium parvulum		85.09	0.48	FRU 99	99		
Clostridium perfringens		6.67	0.30		1		
Actinomyces naeslundii		5.04	0.50		1		
07321101	**Low discrimination**						**07321101**
					COC	SPO	
Clostridium chauvoei		70.55	0.37	SAL 1	1	99	
Actinomyces naeslundii		26.44	0.53	TRE 10	1	1	
Atopobium parvulum		3.01	0.14		99	1	
07321102	**Low discrimination**						**07321102**
					COC		
Atopobium parvulum		42.46	0.35	FRU 99	99		
Lactobacillus acidophilus		32.84	0.32		1		
Actinomyces naeslundii		14.24	0.50		1		
07321103	**Low discrimination**						**07321103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces naeslundii		67.07	0.53	RAF 92	1	1	
Clostridium chauvoei		31.58	0.24	FRU 95	1	99	
Atopobium parvulum		1.35	0.01		99	1	
07321120	**Acceptable identification**						**07321120**
					COC		
Clostridium perfringens		90.27	0.56	FRU 99	1		
Atopobium parvulum		6.29	0.35		99		
Actinomyces naeslundii		2.11	0.50		1		
07321121	**Low discrimination**						**07321121**
					SPO		
Actinomyces naeslundii		45.50	0.53	RAF 92	1		
Clostridium chauvoei		40.47	0.29	FRU 95	99		
Clostridium perfringens		13.11	0.21		84		
07321122	**Low discrimination**						**07321122**
Clostridium perfringens		83.40	0.50	FRU 99			
Lactobacillus acidophilus		9.55	0.32				
Actinomyces naeslundii		4.14	0.50				
07321123	**Low discrimination**						**07321123**
					SPO		
Actinomyces naeslundii		77.37	0.53	RAF 92	1		
Clostridium chauvoei		12.14	0.16	FRU 95	99		
Clostridium perfringens		10.49	0.16		84		
07321141	**Low discrimination**						**07321141**
					SPO		
Actinomyces naeslundii		86.75	0.40	RAF 92	1		
Clostridium chauvoei		13.25	0.03	FRU 95	99		
07321143	**Good identification**						**07321143**
Actinomyces naeslundii		100.0	0.40	RAF 92			
07321160	**Good identification**						**07321160**
					COC		
Clostridium perfringens		97.53	0.43	FRU 99	1		
Actinomyces naeslundii		2.11	0.37		1		
Atopobium parvulum		0.36	0.01		99		
07321161	**Low discrimination**						**07321161**
Actinomyces naeslundii		76.28	0.40	RAF 92			
Clostridium perfringens		23.72	0.09	FRU 95			
07321162	**Low discrimination**						**07321162**
Clostridium perfringens		89.26	0.38	FRU 99			
Lactobacillus acidophilus		6.63	0.16				
Actinomyces naeslundii		4.11	0.37				
07321163	**Low discrimination**						**07321163**
Actinomyces naeslundii		87.23	0.40	RAF 92			
Clostridium perfringens		12.77	0.03	FRU 95			
07323000	**Low discrimination**						**07323000**
					CAT		
Actinomyces naeslundii		70.02	0.47	RAF 92	25		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		17.36	0.22	FRU 95	90		
Lactobacillus jensenii		8.43	0.10		1		
07323001	**Good identification**						**07323001**
Actinomyces naeslundii		100.0	0.50	RAF 92			
07323002	**Low discrimination**						**07323002**
					CAT		
Actinomyces naeslundii		83.19	0.47	RAF 92	25		
Lactobacillus jensenii		6.68	0.07	FRU 95	1		
Propionibacterium granulosum		5.16	0.12		90		
07323003	**Good identification**						**07323003**
Actinomyces naeslundii		100.0	0.50	RAF 92			
07323020	**Low discrimination**						**07323020**
					CAT		
Actinomyces naeslundii		87.08	0.47	RAF 92	25		
Clostridium perfringens		7.53	0.07	FRU 95	1		
Propionibacterium granulosum		5.40	0.12		90		
07323021	**Good identification**						**07323021**
Actinomyces naeslundii		100.0	0.50	RAF 92			
07323022	**Good identification**						**07323022**
Actinomyces naeslundii		96.09	0.47	RAF 92			
Clostridium perfringens		3.91	0.01	FRU 95			
07323023	**Good identification**						**07323023**
Actinomyces naeslundii		100.0	0.50	RAF 92			
07323041	**Good identification**						**07323041**
Actinomyces naeslundii		100.0	0.37	RAF 92			
07323043	**Good identification**						**07323043**
Actinomyces naeslundii		100.0	0.37	RAF 92			
07323061	**Good identification**						**07323061**
Actinomyces naeslundii		100.0	0.37	RAF 92			
07323063	**Good identification**						**07323063**
Actinomyces naeslundii		100.0	0.37	RAF 92			
07323100	**Acceptable identification**						**07323100**
					COC	SPO	
Actinomyces naeslundii		90.59	0.55	RAF 92	1	1	
Atopobium parvulum		7.96	0.14	FRU 95	99	1	
Clostridium tertium		1.44	0.01		1	99	
07323101	**Very good identification**						**07323101**
					SPO		
Actinomyces naeslundii		98.63	0.58	RAF 92	1		
Clostridium chauvoei		1.37	0.03	FRU 95	99		
07323102	**Low discrimination**						**07323102**
					SPO		
Actinomyces naeslundii		86.54	0.55	RAF 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		5.41	0.10	FRU 95	1	
Clostridium tertium		4.13	0.09		99	
07323103	**Very good identification**					**07323103**
Actinomyces naeslundii		100.0	0.58	RAF 92		
07323111	**Good identification**					**07323111**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07323113	**Good identification**					**07323113**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07323120	**Low discrimination**					**07323120**
					SPO	
Actinomyces naeslundii		75.14	0.55	RAF 92	1	
Clostridium perfringens		16.73	0.21	FRU 95	84	
Clostridium tertium		3.59	0.09		99	
07323121	**Very good identification**					**07323121**
Actinomyces naeslundii		100.0	0.58	RAF 92		
07323122	**Low discrimination**					**07323122**
					SPO	
Actinomyces naeslundii		67.09	0.55	RAF 92	1	
Clostridium ramosum		12.08	0.17	FRU 95	92	
Clostridium tertium		9.61	0.17		99	
07323123	**Very good identification**					**07323123**
Actinomyces naeslundii		100.0	0.58	RAF 92		
07323131	**Good identification**					**07323131**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07323133	**Good identification**					**07323133**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07323140	**Good identification**					**07323140**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07323141	**Good identification**					**07323141**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07323142	**Acceptable identification**					**07323142**
Actinomyces naeslundii		91.88	0.42	RAF 92		
Propionibacterium propionicum		8.12	0.14	FRU 95		
07323143	**Good identification**					**07323143**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07323160	**Low discrimination**					**07323160**
Actinomyces naeslundii		80.62	0.42	RAF 92		
Clostridium perfringens		19.38	0.09	FRU 95		
07323161	**Good identification**					**07323161**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.45	RAF 92			
07323162	**Low discrimination**						**07323162**
Actinomyces naeslundii		89.84	0.42	RAF 92			
Clostridium perfringens		10.16	0.03	FRU 95			
07323163	**Good identification**						**07323163**
Actinomyces naeslundii		100.0	0.45	RAF 92			
07325100	**Good identification**						**07325100**
					COC	GRA	
Atopobium parvulum		99.76	0.53	FRU 99	99	99	
Fusobacterium mortiferum		0.12	0.14		1	1	
Actinomyces israelii		0.12	0.16		1	99	
07325102	**Good identification**						**07325102**
					COC		
Atopobium parvulum		98.00	0.40	FRU 99	99		
Bifidobacterium adolescentis		1.01	0.13		1		
Actinomyces israelii		0.99	0.19		1		
07325120	**Acceptable identification**						**07325120**
					COC		
Atopobium parvulum		92.76	0.40	FRU 99	99		
Clostridium perfringens		7.24	0.21		1		
07330000	**Low discrimination**						**07330000**
					GRA	CAT	
Propionibacterium granulosum		43.67	0.58	SAL 1	99	90	
Fusobacterium mortiferum		28.58	0.58		1	5	
Lactobacillus jensenii		16.00	0.44		99	1	
07330001	**Low discrimination**						**07330001**
					GRA	CAT	
Actinomyces naeslundii		46.09	0.43	RAF 92	99	25	
Propionibacterium granulosum		24.75	0.24	GAL 96	99	90	
Fusobacterium mortiferum		16.20	0.24		1	5	
07330002	**Low discrimination**						**07330002**
					GRA	CAT	
Propionibacterium granulosum		41.41	0.48	SAL 1	99	90	
Lactobacillus jensenii		40.47	0.41		99	1	
Fusobacterium mortiferum		12.05	0.42		1	5	
07330003	**Low discrimination**						**07330003**
					CAT		
Actinomyces naeslundii		71.86	0.43	RAF 92	25		
Propionibacterium granulosum		9.65	0.14	GAL 96	90		
Lactobacillus jensenii		9.43	0.07		1		
07330010	**Low discrimination**						**07330010**
					SPO	CAT	
Propionibacterium granulosum		39.81	0.43	SAL 1	1	90	
Lactobacillus jensenii		35.66	0.36	MLZ 12	1	1	
Clostridium botulinum biovar B		22.60	0.26		99	1	
07330020	**Low discrimination**						**07330020**
					CAT		
Clostridium perfringens		37.00	0.41	GAL 99	1		
Propionibacterium granulosum		35.18	0.48	LAC 88	90		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus jensenii		12.89	0.34		1			
07330021	**Low discrimination**							**07330021**
					CAT			
Actinomyces naeslundii		78.40	0.43	RAF 92	25			
Clostridium perfringens		11.08	0.07	GAL 96	1			
Propionibacterium granulosum		10.53	0.14		90			
07330022	**Low discrimination**							**07330022**
					CAT			
Clostridium perfringens		45.48	0.35	GAL 99	1			
Propionibacterium granulosum		22.97	0.38	LAC 88	90			
Lactobacillus jensenii		22.45	0.31		1			
07330023	**Acceptable identification**							**07330023**
Actinomyces naeslundii		93.77	0.43	RAF 92				
Clostridium perfringens		6.23	0.01	GAL 96				
07330040	**Very good identification**							**07330040**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		99.85	0.66	SAL 1	99	99	1	
Propionibacterium granulosum		0.07	0.24		99	1	90	
Fusobacterium mortiferum		0.04	0.24		1	1	5	
07330050	**Very good identification**							**07330050**
					SPO	CAT		
Clostridium botulinum biovar B		99.97	0.61	SAL 1	99	1		
Propionibacterium granulosum		0.02	0.09		1	90		
Lactobacillus jensenii		0.02	0.02		1	1		
07330060	**Very good identification**							**07330060**
					SPO			
Clostridium botulinum biovar B		99.29	0.61	SAL 1	99			
Clostridium perfringens		0.65	0.28		84			
Actinomyces naeslundii		0.03	0.28		1			
07330070	**Good identification**							**07330070**
Clostridium botulinum biovar B		100.0	0.55	SAL 1				
07330100	**Low discrimination**							**07330100**
					COC	GRA		
Fusobacterium mortiferum		63.65	0.65	MLT 11	1	1		
Atopobium parvulum		31.74	0.48		99	99		
Clostridium perfringens		2.49	0.30		1	99		
07330101	**Low discrimination**							**07330101**
					COC	GRA		
Actinomyces naeslundii		66.17	0.51	RAF 92	1	99		
Fusobacterium mortiferum		19.07	0.30	GAL 96	1	1		
Atopobium parvulum		9.51	0.14		99	99		
07330102	**Low discrimination**							**07330102**
					COC	GRA		
Fusobacterium mortiferum		34.35	0.48	MLT 11	1	1		
Atopobium parvulum		27.21	0.35	BGL 10	99	99		
Lactobacillus acidophilus		21.04	0.32		1	99		
07330103	**Low discrimination**							**07330103**
					GRA			
Actinomyces naeslundii		88.02	0.51	RAF 92	99			
Actinomyces viscosus		6.93	0.08	GAL 96	99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		2.82	0.14		1	
07330120	**Low discrimination**					**07330120**
					COC	
Clostridium perfringens		89.88	0.56	GAL 99	1	
Atopobium parvulum		6.26	0.35		99	
Actinomyces naeslundii		1.66	0.48		1	
07330121	**Low discrimination**					**07330121**
					SPO	
Actinomyces naeslundii		70.63	0.51	RAF 92	1	
Clostridium perfringens		25.71	0.21	GAL 96	84	
Clostridium chauvoei		1.87	0.01		99	
07330122	**Low discrimination**					**07330122**
Clostridium perfringens		83.39	0.50	GAL 99		
Lactobacillus acidophilus		9.55	0.32			
Actinomyces naeslundii		3.28	0.48			
07330123	**Low discrimination**					**07330123**
Actinomyces naeslundii		85.38	0.51	RAF 92		
Clostridium perfringens		14.62	0.16	GAL 96		
07330141	**Low discrimination**					**07330141**
					SPO	
Actinomyces naeslundii		85.61	0.38	RAF 92	1	
Clostridium botulinum biovar B		14.39	-.03	GAL 96	99	
07330143	**Good identification**					**07330143**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07330160	**Low discrimination**					**07330160**
					SPO	
Clostridium perfringens		81.19	0.43	GAL 99	84	
Clostridium botulinum biovar B		17.11	0.26		99	
Actinomyces naeslundii		1.39	0.35		1	
07330161	**Low discrimination**					**07330161**
Actinomyces naeslundii		71.79	0.38	RAF 92		
Clostridium perfringens		28.21	0.09	GAL 96		
07330162	**Acceptable identification**					**07330162**
Clostridium perfringens		90.03	0.38	GAL 99		
Lactobacillus acidophilus		6.69	0.16			
Actinomyces naeslundii		3.28	0.35			
07330163	**Low discrimination**					**07330163**
Actinomyces naeslundii		84.40	0.38	RAF 92		
Clostridium perfringens		15.61	0.03	GAL 96		
07330712	**Good identification**					**07330712**
Clostridium clostridioforme		100.0	0.36	RAF 94		
07330732	**Good identification**					**07330732**
Clostridium clostridioforme		100.0	0.36	RAF 94		
07331000	**Good identification**					**07331000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Lactobacillus jensenii		90.15	0.78	CEL 95	99	1	
Propionibacterium granulosum		2.49	0.58		99	90	
Fusobacterium mortiferum		2.44	0.61		1	5	
07331001	**Low discrimination**						**07331001**
					CAT		
Actinomyces naeslundii		52.61	0.67	RAF 92	25		
Lactobacillus jensenii		42.72	0.44		1		
Propionibacterium granulosum		1.18	0.24		90		
07331002	**Good identification**						**07331002**
Lactobacillus jensenii		91.84	0.75	CEL 95			
Lactobacillus acidophilus		3.40	0.48				
Clostridium perfringens		1.44	0.43				
07331003	**Low discrimination**						**07331003**
Actinomyces naeslundii		63.13	0.67	RAF 92			
Lactobacillus jensenii		34.17	0.41				
Lactobacillus acidophilus		1.26	0.14				
07331010	**Very good identification**						**07331010**
					CAT		
Lactobacillus jensenii		98.55	0.70	CEL 95	1		
Propionibacterium granulosum		1.11	0.43		90		
Actinomyces naeslundii		0.13	0.42		25		
07331011	**Low discrimination**						**07331011**
					CAT		
Lactobacillus jensenii		82.94	0.36	CEL 95	1		
Actinomyces naeslundii		16.13	0.45	URE 1	25		
Propionibacterium granulosum		0.94	0.09		90		
07331012	**Very good identification**						**07331012**
					CAT		
Lactobacillus jensenii		99.15	0.67	CEL 95	1		
Propionibacterium granulosum		0.42	0.33		90		
Actinomyces naeslundii		0.20	0.42		25		
07331020	**Low discrimination**						**07331020**
Clostridium perfringens		72.58	0.75	LAC 88			
Lactobacillus jensenii		25.28	0.68				
Actinomyces naeslundii		0.84	0.64				
07331021	**Low discrimination**						**07331021**
Actinomyces naeslundii		55.64	0.67	RAF 92			
Clostridium perfringens		32.42	0.41				
Lactobacillus jensenii		11.29	0.34				
07331022	**Low discrimination**						**07331022**
Clostridium perfringens		62.26	0.69	LAC 88			
Lactobacillus jensenii		30.73	0.65				
Lactobacillus acidophilus		4.55	0.48				
07331023	**Low discrimination**						**07331023**
Actinomyces naeslundii		69.95	0.67	RAF 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		19.18	0.35			
Lactobacillus jensenii		9.47	0.31			
07331030	**Good identification**					**07331030**
					CAT	
Lactobacillus jensenii		90.58	0.60	CEL 95	1	
Clostridium perfringens		7.88	0.41		1	
Propionibacterium granulosum		1.02	0.33		90	
07331032	**Good identification**					**07331032**
Lactobacillus jensenii		92.77	0.57	CEL 95		
Clostridium perfringens		5.70	0.35			
Actinomyces naeslundii		0.73	0.42			
07331033	**Low discrimination**					**07331033**
Actinomyces naeslundii		52.37	0.45	RAF 92		
Lactobacillus jensenii		44.88	0.23	MLZ 5		
Clostridium perfringens		2.76	0.01			
07331040	**Low discrimination**					**07331040**
					SPO	
Lactobacillus jensenii		48.19	0.44	CEL 95	1	
Clostridium botulinum biovar B		20.68	0.32	SOR 1	99	
Clostridium perfringens		20.17	0.37		84	
07331041	**Acceptable identification**					**07331041**
					SPO	
Actinomyces naeslundii		92.09	0.54	RAF 92	1	
Lactobacillus jensenii		4.28	0.10		1	
Clostridium botulinum biovar B		1.84	-.03		99	
07331042	**Low discrimination**					**07331042**
Lactobacillus jensenii		49.80	0.41	CEL 95		
Lactobacillus acidophilus		22.54	0.32	SOR 1		
Clostridium perfringens		14.71	0.31			
07331043	**Good identification**					**07331043**
Actinomyces naeslundii		95.69	0.54	RAF 92		
Lactobacillus jensenii		2.97	0.07			
Lactobacillus acidophilus		1.34	-.02			
07331050	**Low discrimination**					**07331050**
					SPO	
Lactobacillus jensenii		59.50	0.36	CEL 95	1	
Clostridium botulinum biovar B		37.72	0.26	SOR 1	99	
Actinomyces naeslundii		1.36	0.29		1	
07331060	**Very good identification**					**07331060**
					SPO	
Clostridium perfringens		95.59	0.63	LAC 88	84	
Lactobacillus jensenii		1.77	0.34		1	
Clostridium botulinum biovar B		1.49	0.26		99	
07331061	**Low discrimination**					**07331061**
Actinomyces naeslundii		61.39	0.54	RAF 92		
Clostridium perfringens		38.61	0.28			
07331062	**Good identification**					**07331062**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		91.03	0.57	LAC 88			
Lactobacillus acidophilus		4.31	0.32				
Lactobacillus jensenii		2.38	0.31				
07331063	**Low discrimination**						**07331063**
Actinomyces naeslundii		76.34	0.54	RAF 92			
Clostridium perfringens		22.59	0.23				
Lactobacillus acidophilus		1.07	-.02				
07331100	**Low discrimination**						**07331100**
					COC	GRA	
Atopobium parvulum		88.62	0.82		99	99	
Clostridium perfringens		6.94	0.64		1	99	
Fusobacterium mortiferum		2.69	0.68		1	1	
07331101	**Low discrimination**						**07331101**
					COC	SPO	
Actinomyces naeslundii		50.31	0.75	RAF 92	1	1	
Atopobium parvulum		29.82	0.48		99	1	
Clostridium chauvoei		16.49	0.44		1	99	
07331102	**Low discrimination**						**07331102**
					COC		
Atopobium parvulum		47.57	0.69		99		
Lactobacillus acidophilus		36.79	0.66		1		
Clostridium perfringens		9.94	0.58		1		
07331103	**Low discrimination**						**07331103**
					COC		
Actinomyces naeslundii		78.67	0.75	RAF 92	1		
Atopobium parvulum		8.23	0.35		99		
Lactobacillus acidophilus		6.37	0.32		1		
07331110	**Low discrimination**						**07331110**
					COC		
Atopobium parvulum		74.55	0.48	MLZ 1	99		
Lactobacillus jensenii		11.96	0.36		1		
Clostridium perfringens		5.84	0.30		1		
07331111	**Low discrimination**						**07331111**
					COC	SPO	
Actinomyces naeslundii		83.69	0.53	RAF 92	1	1	
Atopobium parvulum		9.52	0.14	MLZ 5	99	1	
Clostridium chauvoei		5.26	0.09		1	99	
07331113	**Acceptable identification**						**07331113**
					COC	GRA	
Actinomyces naeslundii		92.79	0.53	RAF 92	1	99	
Leptotrichia buccalis		5.35	-.03	MLZ 5	1	1	
Atopobium parvulum		1.86	0.01		99	99	
07331120	**Good identification**						**07331120**
					COC	SPO	
Clostridium perfringens		91.72	0.90		1	84	
Atopobium parvulum		6.39	0.69		99	1	
Clostridium septicum		1.24	0.51		1	99	
07331121	**Low discrimination**						**07331121**
					SPO		
Clostridium perfringens		54.88	0.56	URE 1	84		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		36.56	0.75		1	
Clostridium chauvoei		3.99	0.35		99	
07331122	**Low discrimination**					**07331122**
					COC	
Clostridium perfringens		86.09	0.84		1	
Lactobacillus acidophilus		9.86	0.66		1	
Atopobium parvulum		2.25	0.56		99	
07331123	**Low discrimination**					**07331123**
Actinomyces naeslundii		54.61	0.75	RAF 92		
Clostridium perfringens		38.58	0.50			
Lactobacillus acidophilus		4.42	0.32			
07331130	**Low discrimination**					**07331130**
					COC	
Clostridium perfringens		88.79	0.56	MLZ 1	1	
Atopobium parvulum		6.19	0.35		99	
Actinomyces naeslundii		2.08	0.50		1	
07331131	**Low discrimination**					**07331131**
					SPO	
Actinomyces naeslundii		75.24	0.53	RAF 92	1	
Clostridium perfringens		21.68	0.21	MLZ 5	84	
Clostridium chauvoei		1.58	0.01		99	
07331132	**Low discrimination**					**07331132**
Clostridium perfringens		75.95	0.50	MLZ 1		
Lactobacillus acidophilus		8.70	0.32			
Actinomyces naeslundii		3.77	0.50			
07331133	**Low discrimination**					**07331133**
Actinomyces naeslundii		88.06	0.53	RAF 92		
Clostridium perfringens		11.94	0.16	MLZ 5		
07331140	**Low discrimination**					**07331140**
					COC	
Clostridium perfringens		53.92	0.52	NAG 97	1	
Atopobium parvulum		36.50	0.48		99	
Actinomyces naeslundii		7.25	0.59		1	
07331141	**Good identification**					**07331141**
					COC	
Actinomyces naeslundii		90.68	0.62	RAF 92	1	
Clostridium perfringens		4.54	0.17		1	
Atopobium parvulum		3.08	0.14		99	
07331142	**Low discrimination**					**07331142**
Lactobacillus acidophilus		60.23	0.50	CEL 99		
Clostridium perfringens		25.07	0.46	SOR 11		
Actinomyces naeslundii		7.16	0.59			
07331143	**Good identification**					**07331143**
Actinomyces naeslundii		92.06	0.62	RAF 92		
Lactobacillus acidophilus		5.22	0.16			
Clostridium perfringens		2.17	0.12			
07331151	**Good identification**					**07331151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	RAF 92		
07331153	**Good identification**					**07331153**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07331160	**Very good identification**					**07331160**
					COC	
Clostridium perfringens		99.09	0.77		1	
Actinomyces naeslundii		0.41	0.59		1	
Atopobium parvulum		0.37	0.35		99	
07331161	**Low discrimination**					**07331161**
					SPO	
Clostridium perfringens		61.55	0.43	URE 1	84	
Actinomyces naeslundii		37.98	0.62		1	
Clostridium chauvoei		0.24	0.01		99	
07331162	**Good identification**					**07331162**
Clostridium perfringens		92.09	0.72			
Lactobacillus acidophilus		6.84	0.50			
Actinomyces naeslundii		0.81	0.59			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07331163	**Low discrimination**						**07331163**
Actinomyces naeslundii		54.97	0.62	RAF 92			
Clostridium perfringens		41.92	0.38				
Lactobacillus acidophilus		3.12	0.16				
07331170	**Good identification**						**07331170**
					COC		
Clostridium perfringens		97.53	0.43	MLZ 1	1		
Actinomyces naeslundii		2.11	0.37		1		
Atopobium parvulum		0.36	0.01		99		
07331171	**Low discrimination**						**07331171**
Actinomyces naeslundii		76.28	0.40	RAF 92			
Clostridium perfringens		23.72	0.09	MLZ 5			
07331172	**Low discrimination**						**07331172**
Clostridium perfringens		89.26	0.38	MLZ 1			
Lactobacillus acidophilus		6.63	0.16				
Actinomyces naeslundii		4.11	0.37				
07331173	**Low discrimination**						**07331173**
Actinomyces naeslundii		87.23	0.40	RAF 92			
Clostridium perfringens		12.77	0.03	MLZ 5			
07331200	**Low discrimination**						**07331200**
					GRA	CAT	
Lactobacillus jensenii		85.47	0.44	CEL 95	99	1	
Propionibacterium granulosum		7.21	0.32	RHA 1	99	90	
Fusobacterium mortiferum		2.31	0.27		1	5	
07331202	**Low discrimination**						**07331202**
					CAT		
Lactobacillus jensenii		73.33	0.41	CEL 95	1		
Lactobacillus acidophilus		20.21	0.29	RHA 1	1		
Propionibacterium granulosum		2.32	0.22		90		
07331210	**Good identification**						**07331210**
					CAT		
Lactobacillus jensenii		96.66	0.36	CEL 95	1		
Propionibacterium granulosum		3.34	0.17	RHA 1	90		
07331220	**Low discrimination**						**07331220**
					CAT		
Clostridium perfringens		71.37	0.41	LAC 88	1		
Lactobacillus jensenii		24.86	0.34	RHA 1	1		
Propionibacterium granulosum		2.10	0.22		90		
07331222	**Low discrimination**						**07331222**
Clostridium perfringens		44.79	0.35	LAC 88			
Lactobacillus acidophilus		24.37	0.29	RHA 1			
Lactobacillus jensenii		22.10	0.31				
07331300	**Low discrimination**						**07331300**
					COC	GRA	
Atopobium parvulum		87.91	0.48	RHA 1	99	99	
Clostridium perfringens		6.89	0.30		1	99	
Fusobacterium mortiferum		2.67	0.33		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07331301	**Low discrimination**						**07331301**
					COC	SPO	
Actinomyces naeslundii		52.07	0.41	RAF 92	1	1	
Atopobium parvulum		30.87	0.14	RHA 1	99	1	
Clostridium chauvoei		17.07	0.09		1	99	
07331302	**Low discrimination**						**07331302**
					COC		
Lactobacillus acidophilus		79.65	0.47	CEL 99	1		
Atopobium parvulum		13.82	0.35	RHA 7	99		
Clostridium perfringens		2.89	0.24		1		
07331303	**Low discrimination**						**07331303**
					COC		
Actinomyces naeslundii		58.57	0.41	RAF 92	1		
Lactobacillus acidophilus		35.31	0.13	RHA 1	1		
Atopobium parvulum		6.13	0.01		99		
07331320	**Low discrimination**						**07331320**
					COC	SPO	
Clostridium perfringens		87.29	0.56	RHA 1	1	84	
Atopobium parvulum		6.08	0.35		99	1	
Clostridium ramosum		4.13	0.30		1	92	
07331322	**Low discrimination**						**07331322**
					SPO		
Clostridium perfringens		38.68	0.50	RHA 1	84		
Lactobacillus acidophilus		33.01	0.47		1		
Clostridium ramosum		15.55	0.40		92		
07331323	**Low discrimination**						**07331323**
Actinomyces naeslundii		36.98	0.41	RAF 92			
Clostridium perfringens		26.12	0.16	RHA 1			
Lactobacillus acidophilus		22.29	0.13				
07331332	**Low discrimination**						**07331332**
					GRA		
Parabacteroides distasonis		59.21	0.39	RAF 99	1		
Clostridium clostridioforme		17.59	0.24	XYL 90	50		
Clostridium perfringens		9.62	0.16		99		
07331360	**Good identification**						**07331360**
					COC		
Clostridium perfringens		99.22	0.43	RHA 1	1		
Actinomyces naeslundii		0.41	0.25		1		
Atopobium parvulum		0.37	0.01		99		
07331362	**Low discrimination**						**07331362**
					SPO		
Clostridium perfringens		62.50	0.38	RHA 1	84		
Lactobacillus acidophilus		34.60	0.31		1		
Clostridium ramosum		1.33	0.06		92		
07331400	**Low discrimination**						**07331400**
					GRA	CAT	
Lactobacillus jensenii		88.40	0.44	CEL 95	99	1	
Propionibacterium granulosum		2.44	0.24	ARA 1	99	90	
Fusobacterium mortiferum		2.39	0.27		1	5	
07331402	**Low discrimination**						**07331402**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		88.38	0.41	CEL 95			
Lactobacillus acidophilus		3.27	0.14	ARA 1			
Actinomyces israelii		1.61	0.24				
07331410	**Good identification**						**07331410**
					CAT		
Lactobacillus jensenii		98.89	0.36	CEL 95	1		
Propionibacterium granulosum		1.12	0.09	ARA 1	90		
07331420	**Low discrimination**						**07331420**
Clostridium perfringens		72.74	0.41	LAC 88			
Lactobacillus jensenii		25.34	0.34	ARA 1			
Actinomyces naeslundii		0.84	0.30				
07331422	**Low discrimination**						**07331422**
Clostridium perfringens		62.11	0.35	LAC 88			
Lactobacillus jensenii		30.65	0.31	ARA 1			
Lactobacillus acidophilus		4.54	0.14				
07331500	**Low discrimination**						**07331500**
					COC	GRA	
Atopobium parvulum		86.48	0.48	ARA 1	99	99	
Clostridium perfringens		6.77	0.30		1	99	
Fusobacterium mortiferum		2.63	0.33		1	1	
07331501	**Low discrimination**						**07331501**
					COC	SPO	
Actinomyces naeslundii		52.07	0.41	RAF 92	1	1	
Atopobium parvulum		30.87	0.14	ARA 1	99	1	
Clostridium chauvoei		17.07	0.09		1	99	
07331502	**Low discrimination**						**07331502**
					COC		
Bifidobacterium adolescentis		44.94	0.47	RAF 92	1		
Atopobium parvulum		23.70	0.35	XYL 87	99		
Lactobacillus acidophilus		18.34	0.32		1		
07331503	**Low discrimination**						**07331503**
					COC		
Actinomyces naeslundii		76.75	0.41	RAF 92	1		
Bifidobacterium adolescentis		15.22	0.13	ARA 1	1		
Atopobium parvulum		8.03	0.01		99		
07331512	**Low discrimination**						**07331512**
					GRA		
Bifidobacterium adolescentis		87.08	0.49	RAF 92	99		
Clostridium clostridioforme		10.02	0.30	XYL 87	50		
Leptotrichia buccalis		1.09	-.03		1		
07331520	**Acceptable identification**						**07331520**
					COC	SPO	
Clostridium perfringens		91.69	0.56	ARA 1	1	84	
Atopobium parvulum		6.39	0.35		99	1	
Clostridium septicum		1.24	0.17		1	99	
07331522	**Low discrimination**						**07331522**
					COC		
Clostridium perfringens		83.22	0.50	ARA 1	1		
Lactobacillus acidophilus		9.53	0.32		1		
Atopobium parvulum		2.17	0.22		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07331523	**Low discrimination**						**07331523**
Actinomyces naeslundii		58.60	0.41	RAF 92			
Clostridium perfringens		41.40	0.16	ARA 1			
07331560	**Good identification**						**07331560**
					COC		
Clostridium perfringens		99.22	0.43	ARA 1	1		
Actinomyces naeslundii		0.41	0.25		1		
Atopobium parvulum		0.37	0.01		99		
07331562	**Acceptable identification**						**07331562**
Clostridium perfringens		92.32	0.38	ARA 1			
Lactobacillus acidophilus		6.86	0.16				
Actinomyces naeslundii		0.82	0.25				
07331612	**Good identification**						**07331612**
Clostridium clostridioforme		100.0	0.35	RAF 94			
07331632	**Good identification**						**07331632**
Clostridium clostridioforme		100.0	0.35	RAF 94			
07331712	**Good identification**						**07331712**
Clostridium clostridioforme		98.82	0.44	RAF 94			
Bifidobacterium adolescentis		1.18	0.15	XYL 91			
07331722	**Low discrimination**						**07331722**
					GRA		
Parabacteroides distasonis		44.61	0.40	RAF 99	1		
Clostridium clostridioforme		40.04	0.33	XYL 90	50		
Clostridium perfringens		6.37	0.16		99		
07331732	**Low discrimination**						**07331732**
Clostridium clostridioforme		88.36	0.44	RAF 94			
Parabacteroides distasonis		11.64	0.35	XYL 91			
07332000	**Good identification**						**07332000**
					GRA	CAT	
Propionibacterium granulosum		95.20	0.56	SAL 1	99	90	
Actinomyces naeslundii		3.07	0.45		99	25	
Fusobacterium mortiferum		0.83	0.24		1	5	
07332001	**Low discrimination**						**07332001**
Actinomyces naeslundii		82.73	0.48	RAF 92			
Propionibacterium granulosum		17.27	0.22	GAL 96			
07332002	**Low discrimination**						**07332002**
					CAT		
Propionibacterium granulosum		86.27	0.46	SAL 1	90		
Actinomyces naeslundii		11.13	0.45		25		
Lactobacillus jensenii		1.13	0.07		1		
07332003	**Good identification**						**07332003**
Actinomyces naeslundii		95.04	0.48	RAF 92			
Propionibacterium granulosum		4.96	0.12	GAL 96			
07332010	**Good identification**						**07332010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Propionibacterium granulosum		96.61	0.41	SAL 1	90	
Actinomyces naeslundii		1.20	0.23	MLZ 12	25	
Lactobacillus jensenii		1.16	0.02		1	
07332020	**Low discrimination**					**07332020**
					CAT	
Propionibacterium granulosum		87.48	0.46	SAL 1	90	
Actinomyces naeslundii		11.29	0.45		25	
Clostridium perfringens		1.23	0.07		1	
07332021	**Good identification**					**07332021**
Actinomyces naeslundii		95.04	0.48	RAF 92		
Propionibacterium granulosum		4.96	0.12	GAL 96		
07332022	**Low discrimination**					**07332022**
					CAT	
Propionibacterium granulosum		64.82	0.35	SAL 1	90	
Actinomyces naeslundii		33.46	0.45		25	
Clostridium perfringens		1.72	0.01		1	
07332023	**Good identification**					**07332023**
Actinomyces naeslundii		100.0	0.48	RAF 92		
07332041	**Low discrimination**					**07332041**
					SPO	
Actinomyces naeslundii		80.22	0.36	RAF 92	1	
Clostridium botulinum biovar B		19.78	-.03	GAL 96	99	
07332043	**Good identification**					**07332043**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07332061	**Good identification**					**07332061**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07332063	**Good identification**					**07332063**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07332100	**Low discrimination**					**07332100**
					GRA	
Propionibacterium propionicum		47.03	0.45	NIT 86	99	
Actinomyces naeslundii		36.18	0.53	SAL 1	99	
Fusobacterium mortiferum		8.06	0.30		1	
07332101	**Good identification**					**07332101**
Actinomyces naeslundii		91.22	0.56	RAF 92		
Propionibacterium propionicum		8.78	0.29	GAL 96		
07332102	**Low discrimination**					**07332102**
Propionibacterium propionicum		61.34	0.48	NIT 86		
Actinomyces naeslundii		31.46	0.53	SAL 1		
Lactobacillus acidophilus		2.48	0.10			
07332103	**Low discrimination**					**07332103**
Actinomyces naeslundii		87.38	0.56	RAF 92		
Propionibacterium propionicum		12.62	0.32	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07332120	**Low discrimination**					**07332120**
					SPO	
Actinomyces naeslundii		69.18	0.53	RAF 92	1	
Clostridium perfringens		19.46	0.21	GAL 96	84	
Clostridium tertium		4.17	0.09		99	
07332121	**Very good identification**					**07332121**
Actinomyces naeslundii		100.0	0.56	RAF 92		
07332122	**Low discrimination**					**07332122**
					SPO	
Actinomyces naeslundii		61.00	0.53	RAF 92	1	
Clostridium ramosum		13.87	0.17	GAL 96	92	
Clostridium tertium		11.04	0.17		99	
07332123	**Very good identification**					**07332123**
Actinomyces naeslundii		100.0	0.56	RAF 92		
07332140	**Low discrimination**					**07332140**
					SPO	
Propionibacterium propionicum		86.70	0.45	NIT 86	1	
Actinomyces naeslundii		11.77	0.40	SAL 1	1	
Clostridium botulinum biovar B		1.53	-.03		99	
07332141	**Low discrimination**					**07332141**
Actinomyces naeslundii		64.70	0.43	RAF 92		
Propionibacterium propionicum		35.30	0.29	GAL 96		
07332142	**Acceptable identification**					**07332142**
Propionibacterium propionicum		91.70	0.48	NIT 86		
Actinomyces naeslundii		8.30	0.40	SAL 1		
07332143	**Low discrimination**					**07332143**
Actinomyces naeslundii		54.99	0.43	RAF 92		
Propionibacterium propionicum		45.01	0.32	GAL 96		
07332160	**Low discrimination**					**07332160**
Actinomyces naeslundii		76.71	0.40	RAF 92		
Clostridium perfringens		23.29	0.09	GAL 96		
07332161	**Good identification**					**07332161**
Actinomyces naeslundii		100.0	0.43	RAF 92		
07332162	**Low discrimination**					**07332162**
Actinomyces naeslundii		79.72	0.40	RAF 92		
Clostridium perfringens		11.39	0.03	GAL 96		
Propionibacterium propionicum		8.90	0.14			
07332163	**Good identification**					**07332163**
Actinomyces naeslundii		100.0	0.43	RAF 92		
07333000	**Low discrimination**					**07333000**
					CAT	
Propionibacterium granulosum		42.21	0.56	SAL 1	90	
Actinomyces naeslundii		32.68	0.69		25	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		20.50	0.44		1	
07333001	**Very good identification**					**07333001**
					CAT	
Actinomyces naeslundii		98.72	0.72	RAF 92	25	
Propionibacterium granulosum		0.86	0.22		90	
Lactobacillus jensenii		0.42	0.10		1	
07333002	**Low discrimination**					**07333002**
					CAT	
Actinomyces naeslundii		51.09	0.69	RAF 92	25	
Lactobacillus jensenii		21.37	0.41		1	
Propionibacterium granulosum		16.50	0.46		90	
07333003	**Very good identification**					**07333003**
					CAT	
Actinomyces naeslundii		99.50	0.72	RAF 92	25	
Lactobacillus jensenii		0.28	0.07		1	
Propionibacterium granulosum		0.22	0.12		90	
07333010	**Low discrimination**					**07333010**
					CAT	
Lactobacillus jensenii		46.49	0.36	CEL 95	1	
Propionibacterium granulosum		39.15	0.41	MAN 1	90	
Actinomyces naeslundii		11.70	0.47		25	
07333011	**Good identification**					**07333011**
Actinomyces naeslundii		97.39	0.50	RAF 92		
Lactobacillus jensenii		2.61	0.02	MLZ 5		
07333013	**Good identification**					**07333013**
Actinomyces naeslundii		100.0	0.50	RAF 92		
07333020	**Low discrimination**					**07333020**
					CAT	
Actinomyces naeslundii		51.24	0.69	RAF 92	25	
Clostridium perfringens		23.07	0.41		1	
Propionibacterium granulosum		16.55	0.46		90	
07333021	**Very good identification**					**07333021**
					CAT	
Actinomyces naeslundii		99.48	0.72	RAF 92	25	
Clostridium perfringens		0.30	0.07		1	
Propionibacterium granulosum		0.22	0.12		90	
07333022	**Low discrimination**					**07333022**
Actinomyces naeslundii		64.99	0.69	RAF 92		
Clostridium perfringens		13.77	0.35			
Lactobacillus jensenii		6.80	0.31			
07333023	**Very good identification**					**07333023**
Actinomyces naeslundii		99.86	0.72	RAF 92		
Clostridium perfringens		0.14	0.01			
07333030	**Low discrimination**					**07333030**
					CAT	
Actinomyces naeslundii		31.67	0.47	RAF 92	25	
Lactobacillus jensenii		31.46	0.26	MLZ 5	1	
Propionibacterium granulosum		26.49	0.31		90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07333031	**Good identification**					**07333031**
Actinomyces naeslundii		100.0	0.50	RAF 92		
07333032	**Low discrimination**					**07333032**
					SPO	
Actinomyces naeslundii		37.92	0.47	RAF 92	1	
Clostridium tertium		27.49	0.21	MLZ 5	99	
Lactobacillus jensenii		25.11	0.23		1	
07333033	**Good identification**					**07333033**
Actinomyces naeslundii		100.0	0.50	RAF 92		
07333040	**Low discrimination**					**07333040**
					CAT	
Actinomyces naeslundii		82.13	0.56	RAF 92	25	
Actinomyces israelii		6.35	0.28		1	
Propionibacterium granulosum		6.07	0.22		90	
07333041	**Very good identification**					**07333041**
Actinomyces naeslundii		100.0	0.59	RAF 92		
07333042	**Low discrimination**					**07333042**
Actinomyces naeslundii		81.43	0.56	RAF 92		
Actinomyces israelii		9.44	0.31			
Lactobacillus acidophilus		4.60	0.10			
07333043	**Very good identification**					**07333043**
Actinomyces naeslundii		100.0	0.59	RAF 92		
07333051	**Good identification**					**07333051**
Actinomyces naeslundii		100.0	0.37	RAF 92		
07333053	**Good identification**					**07333053**
Actinomyces naeslundii		100.0	0.37	RAF 92		
07333060	**Low discrimination**					**07333060**
					CAT	
Actinomyces naeslundii		66.47	0.56	RAF 92	25	
Clostridium perfringens		32.30	0.28		1	
Propionibacterium granulosum		1.23	0.12		90	
07333061	**Very good identification**					**07333061**
Actinomyces naeslundii		100.0	0.59	RAF 92		
07333062	**Low discrimination**					**07333062**
Actinomyces naeslundii		77.81	0.56	RAF 92		
Clostridium perfringens		17.80	0.23			
Lactobacillus acidophilus		4.39	0.10			
07333063	**Very good identification**					**07333063**
Actinomyces naeslundii		100.0	0.59	RAF 92		
07333071	**Good identification**					**07333071**
Actinomyces naeslundii		100.0	0.37	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07333073	**Good identification**						**07333073**
Actinomyces naeslundii		100.0	0.37	RAF 92			
07333100	**Low discrimination**						**07333100**
					COC	SPO	
Actinomyces naeslundii		58.56	0.77	RAF 92	1	1	
Atopobium parvulum		26.82	0.48		99	1	
Clostridium tertium		4.86	0.35		1	99	
07333101	**Very good identification**						**07333101**
					COC		
Actinomyces naeslundii		99.07	0.80	RAF 92	1		
Propionibacterium propionicum		0.40	0.29		1		
Atopobium parvulum		0.31	0.14		99		
07333102	**Low discrimination**						**07333102**
					SPO		
Actinomyces naeslundii		53.68	0.77	RAF 92	1		
Lactobacillus acidophilus		17.49	0.44		1		
Clostridium tertium		13.36	0.43		99		
07333103	**Very good identification**						**07333103**
Actinomyces naeslundii		98.92	0.80	RAF 92			
Propionibacterium propionicum		0.60	0.32				
Lactobacillus acidophilus		0.22	0.10				
07333110	**Low discrimination**						**07333110**
					SPO		
Clostridium tertium		69.29	0.39	CEL 99	99		
Actinomyces naeslundii		24.73	0.55		1		
Actinomyces israelii		3.46	0.31		1		
07333111	**Very good identification**						**07333111**
					SPO		
Actinomyces naeslundii		98.15	0.58	RAF 92	1		
Clostridium tertium		1.85	0.05	MLZ 5	99		
07333112	**Low discrimination**						**07333112**
					SPO		
Clostridium tertium		82.50	0.47	CEL 99	99		
Actinomyces naeslundii		9.82	0.55		1		
Bifidobacterium adolescentis		4.27	0.33		1		
07333113	**Good identification**						**07333113**
					SPO		
Actinomyces naeslundii		94.64	0.58	RAF 92	1		
Clostridium tertium		5.36	0.13	MLZ 5	99		
07333120	**Low discrimination**						**07333120**
					SPO		
Clostridium perfringens		42.17	0.56	MAN 1	84		
Actinomyces naeslundii		36.35	0.77		1		
Clostridium tertium		9.05	0.43		99		
07333121	**Very good identification**						**07333121**
					SPO		
Actinomyces naeslundii		98.80	0.80	RAF 92	1		
Clostridium perfringens		0.77	0.21		84		
Clostridium tertium		0.17	0.09		99		
07333122	**Low discrimination**						**07333122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		27.87	0.77	RAF 92	1	
Clostridium ramosum		26.15	0.51		92	
Clostridium tertium		20.81	0.51		99	
07333123	**Very good identification**					**07333123**
					SPO	
Actinomyces naeslundii		98.31	0.80	RAF 92	1	
Clostridium ramosum		0.62	0.17		92	
Clostridium tertium		0.49	0.17		99	
07333130	**Low discrimination**					**07333130**
					SPO	
Clostridium tertium		86.76	0.47	CEL 99	99	
Actinomyces naeslundii		10.32	0.55		1	
Clostridium perfringens		2.30	0.21		84	
07333131	**Good identification**					**07333131**
					SPO	
Actinomyces naeslundii		94.64	0.58	RAF 92	1	
Clostridium tertium		5.36	0.13	MLZ 5	99	
07333132	**Acceptable identification**					**07333132**
					SPO	
Clostridium tertium		94.88	0.55	CEL 99	99	
Actinomyces naeslundii		3.76	0.55		1	
Clostridium ramosum		0.68	0.17		92	
07333133	**Low discrimination**					**07333133**
					SPO	
Actinomyces naeslundii		85.49	0.58	RAF 92	1	
Clostridium tertium		14.52	0.21	MLZ 5	99	
07333140	**Low discrimination**					**07333140**
Actinomyces naeslundii		67.33	0.64	RAF 92		
Propionibacterium propionicum		20.67	0.45			
Actinomyces israelii		7.31	0.38			
07333141	**Very good identification**					**07333141**
Actinomyces naeslundii		97.78	0.67	RAF 92		
Propionibacterium propionicum		2.22	0.29			
07333142	**Low discrimination**					**07333142**
Actinomyces naeslundii		52.59	0.64	RAF 92		
Propionibacterium propionicum		24.21	0.48			
Lactobacillus acidophilus		12.00	0.28			
07333143	**Very good identification**					**07333143**
Actinomyces naeslundii		96.70	0.67	RAF 92		
Propionibacterium propionicum		3.30	0.32			
07333150	**Low discrimination**					**07333150**
					SPO	
Actinomyces naeslundii		68.09	0.42	RAF 92	1	
Actinomyces israelii		20.99	0.24	MLZ 5	1	
Clostridium tertium		10.92	0.05		99	
07333151	**Good identification**					**07333151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	RAF 92		
07333152	**Low discrimination**					**07333152**
					SPO	
Actinomyces naeslundii		42.80	0.42	RAF 92	1	
Clostridium tertium		20.59	0.13	MLZ 5	99	
Actinomyces israelii		19.79	0.27		1	
07333153	**Good identification**					**07333153**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07333160	**Low discrimination**					**07333160**
					SPO	
Clostridium perfringens		54.65	0.43	MAN 1	84	
Actinomyces naeslundii		43.64	0.64		1	
Clostridium tertium		0.62	0.09		99	
07333161	**Very good identification**					**07333161**
Actinomyces naeslundii		99.16	0.67	RAF 92		
Clostridium perfringens		0.84	0.09			
07333162	**Low discrimination**					**07333162**
Actinomyces naeslundii		51.72	0.64	RAF 92		
Clostridium perfringens		30.48	0.38			
Lactobacillus acidophilus		11.80	0.28			
07333163	**Very good identification**					**07333163**
Actinomyces naeslundii		99.61	0.67	RAF 92		
Clostridium perfringens		0.40	0.03			
07333170	**Low discrimination**					**07333170**
					SPO	
Actinomyces naeslundii		58.09	0.42	RAF 92	1	
Clostridium tertium		27.95	0.13	MLZ 5	99	
Clostridium perfringens		13.96	0.09		84	
07333171	**Good identification**					**07333171**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07333173	**Good identification**					**07333173**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07333201	**Good identification**					**07333201**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333203	**Good identification**					**07333203**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333221	**Good identification**					**07333221**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333223	**Good identification**					**07333223**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333300	**Low discrimination**					**07333300**
					COC	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		45.06	0.43	RAF 92	1	
Atopobium parvulum		20.64	0.14	RHA 1	99	
Actinomyces israelii		16.55	0.26		1	
07333301	**Good identification**					**07333301**
Actinomyces naeslundii		100.0	0.46	RAF 92		
07333303	**Good identification**					**07333303**
Actinomyces naeslundii		100.0	0.46	RAF 92		
07333320	**Acceptable identification**					**07333320**
					SPO	
Clostridium ramosum		92.52	0.43	CEL 99	92	
Clostridium perfringens		3.49	0.21		84	
Actinomyces naeslundii		3.01	0.43		1	
07333321	**Low discrimination**					**07333321**
					SPO	
Actinomyces naeslundii		82.83	0.46	RAF 92	1	
Clostridium ramosum		17.17	0.09	RHA 1	92	
07333322	**Good identification**					**07333322**
					SPO	
Clostridium ramosum		96.18	0.53	CEL 99	92	
Lactobacillus acidophilus		1.90	0.25		1	
Actinomyces naeslundii		0.78	0.43		1	
07333323	**Low discrimination**					**07333323**
					SPO	
Actinomyces naeslundii		54.67	0.46	RAF 92	1	
Clostridium ramosum		45.33	0.19	RHA 1	92	
07333401	**Good identification**					**07333401**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333402	**Low discrimination**					**07333402**
					SPO	
Actinomyces israelii		57.55	0.36	CEL 93	1	
Clostridium innocuum		14.11	0.11	XYL 92	99	
Actinomyces naeslundii		13.51	0.35		1	
07333403	**Good identification**					**07333403**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333421	**Good identification**					**07333421**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333423	**Good identification**					**07333423**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07333500	**Low discrimination**					**07333500**
					COC	
Actinomyces israelii		70.55	0.44	CEL 93	1	
Actinomyces naeslundii		17.67	0.43	XYL 92	1	
Atopobium parvulum		8.10	0.14		99	
07333501	**Good identification**					**07333501**
Actinomyces naeslundii		100.0	0.46	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07333502	**Low discrimination**							**07333502**
Bifidobacterium adolescentis		61.64	0.46	RAF 92				
Actinomyces israelii		29.74	0.47	XYL 87				
Actinomyces naeslundii		4.97	0.43					
07333503	**Acceptable identification**							**07333503**
Actinomyces naeslundii		92.29	0.46	RAF 92				
Bifidobacterium adolescentis		7.71	0.12	ARA 1				
07333512	**Acceptable identification**							**07333512**
					SPO			
Bifidobacterium adolescentis		91.90	0.47	RAF 92	1			
Actinomyces israelii		5.42	0.33	XYL 87	1			
Clostridium tertium		2.68	0.13		99			
07333521	**Good identification**							**07333521**
Actinomyces naeslundii		100.0	0.46	RAF 92				
07333522	**Low discrimination**							**07333522**
					SPO			
Actinomyces naeslundii		24.61	0.43	RAF 92	1			
Clostridium ramosum		23.09	0.17	ARA 1	92			
Clostridium tertium		18.37	0.17		99			
07333523	**Good identification**							**07333523**
Actinomyces naeslundii		100.0	0.46	RAF 92				
07333540	**Low discrimination**							**07333540**
Actinomyces israelii		89.79	0.37	CEL 93				
Actinomyces naeslundii		10.21	0.30	XYL 92				
07333542	**Low discrimination**							**07333542**
Actinomyces israelii		56.51	0.40	CEL 93				
Bifidobacterium adolescentis		37.23	0.30	XYL 92				
Actinomyces naeslundii		4.28	0.30					
07334000	**Low discrimination**							**07334000**
					COC	GRA	CAT	
Fusobacterium mortiferum		56.57	0.39	MLT 11	1	1	5	
Atopobium parvulum		16.40	0.18	XYL 7	99	99	1	
Propionibacterium granulosum		11.60	0.24		1	99	90	
07334100	**Acceptable identification**							**07334100**
					COC	GRA		
Atopobium parvulum		92.38	0.53	GAL 99	99	99		
Fusobacterium mortiferum		7.51	0.45		1	1		
Actinomyces israelii		0.11	0.16		1	99		
07334102	**Acceptable identification**							**07334102**
					COC	GRA		
Atopobium parvulum		92.73	0.40	GAL 99	99	99		
Fusobacterium mortiferum		4.75	0.29		1	1		
Bifidobacterium adolescentis		0.96	0.13		1	99		
07334120	**Acceptable identification**							**07334120**
					COC	GRA		
Atopobium parvulum		92.36	0.40	GAL 99	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		7.21	0.21		1	99	
Fusobacterium mortiferum		0.43	0.11		1	1	
07334512	**Low discrimination**						**07334512**
Clostridium clostridioforme		85.15	0.39	RAF 94			
Bifidobacterium adolescentis		14.85	0.29	RHA 88			
07334532	**Good identification**						**07334532**
Clostridium clostridioforme		100.0	0.39	RAF 94			
07334612	**Good identification**						**07334612**
Clostridium clostridioforme		100.0	0.44	RAF 94			
07334632	**Good identification**						**07334632**
Clostridium clostridioforme		100.0	0.44	RAF 94			
07334702	**Good identification**						**07334702**
Clostridium clostridioforme		100.0	0.42	RAF 94			
07334712	**Good identification**						**07334712**
Clostridium clostridioforme		100.0	0.53	RAF 94			
07334722	**Good identification**						**07334722**
Clostridium clostridioforme		97.08	0.42	RAF 94			
Parabacteroides distasonis		2.92	0.22				
07334732	**Good identification**						**07334732**
Clostridium clostridioforme		99.65	0.53	RAF 94			
Parabacteroides distasonis		0.35	0.17				
07335000	**Low discrimination**						**07335000**
					COC	GRA	
Atopobium parvulum		73.55	0.53	LAC 99	99	99	
Lactobacillus jensenii		19.06	0.44		1	99	
Fusobacterium mortiferum		3.84	0.42		1	1	
07335002	**Low discrimination**						**07335002**
					COC		
Atopobium parvulum		42.44	0.40	LAC 99	99		
Lactobacillus jensenii		41.55	0.41		1		
Actinomyces israelii		10.60	0.43		1		
07335010	**Low discrimination**						**07335010**
					COC		
Lactobacillus jensenii		84.81	0.36	CEL 95	1		
Atopobium parvulum		9.92	0.18	XYL 1	99		
Actinomyces israelii		4.31	0.26		1		
07335020	**Low discrimination**						**07335020**
					COC		
Clostridium perfringens		42.95	0.41	XYL 1	1		
Atopobium parvulum		40.75	0.40	LAC 88	99		
Lactobacillus jensenii		14.96	0.34		1		
07335022	**Low discrimination**						**07335022**
					COC		
Clostridium perfringens		49.06	0.35	XYL 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		24.21	0.31	LAC 88	1		
Atopobium parvulum		17.46	0.27		99		
07335042	**Low discrimination**						**07335042**
					COC		
Actinomyces israelii		79.89	0.36	CEL 93	1		
Atopobium parvulum		8.30	0.06		99		
Lactobacillus jensenii		8.13	0.07		1		
07335100	**Very good identification**						**07335100**
					COC	GRA	
Atopobium parvulum		99.69	0.87		99	99	
Fusobacterium mortiferum		0.12	0.48		1	1	
Actinomyces israelii		0.12	0.50		1	99	
07335101	**Good identification**						**07335101**
					COC	SPO	
Atopobium parvulum		98.56	0.53	URE 1	99	1	
Actinomyces naeslundii		0.90	0.41		1	1	
Clostridium chauvoei		0.30	0.09		1	99	
07335102	**Very good identification**						**07335102**
					COC		
Atopobium parvulum		97.26	0.74		99		
Bifidobacterium adolescentis		1.00	0.47		1		
Actinomyces israelii		0.98	0.53		1		
07335103	**Acceptable identification**						**07335103**
					COC		
Atopobium parvulum		93.25	0.40	URE 1	99		
Actinomyces naeslundii		4.85	0.41		1		
Bifidobacterium adolescentis		0.96	0.13		1		
07335110	**Good identification**						**07335110**
					COC	SPO	
Atopobium parvulum		97.63	0.53	MLZ 1	99	1	
Actinomyces israelii		1.72	0.36		1	1	
Clostridium tertium		0.24	0.07		1	99	
07335112	**Low discrimination**						**07335112**
					COC		
Bifidobacterium adolescentis		48.87	0.49	RAF 92	1		
Atopobium parvulum		39.17	0.40	ARA 87	99		
Actinomyces israelii		5.85	0.39		1		
07335120	**Good identification**						**07335120**
					COC	SPO	
Atopobium parvulum		92.54	0.74		99	1	
Clostridium perfringens		7.22	0.56		1	84	
Clostridium septicum		0.10	0.17		1	99	
07335121	**Low discrimination**						**07335121**
					COC		
Atopobium parvulum		88.05	0.40	URE 1	99		
Clostridium perfringens		6.87	0.21		1		
Actinomyces naeslundii		4.58	0.41		1		
07335122	**Low discrimination**						**07335122**
					COC	GRA	
Atopobium parvulum		75.31	0.61		99	99	
Clostridium perfringens		15.68	0.50		1	99	
Parabacteroides distasonis		5.43	0.51		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07335130	**Low discrimination**							**07335130**
					COC	SPO		
Atopobium parvulum		88.31	0.40	MLZ 1	99	1		
Clostridium perfringens		6.89	0.21		1	84		
Clostridium tertium		3.63	0.15		1	99		
07335132	**Low discrimination**							**07335132**
					COC	GRA	SPO	
Parabacteroides distasonis		60.05	0.47	RAF 99	1	1	1	
Atopobium parvulum		15.62	0.27		99	99	1	
Clostridium tertium		10.92	0.24		1	99	99	
07335140	**Acceptable identification**							**07335140**
					COC			
Atopobium parvulum		94.59	0.53	SOR 1	99			
Actinomyces israelii		4.33	0.43		1			
Clostridium perfringens		0.76	0.17		1			
07335142	**Low discrimination**							**07335142**
					COC			
Atopobium parvulum		62.74	0.40	SOR 1	99			
Actinomyces israelii		24.39	0.46		1			
Bifidobacterium adolescentis		7.92	0.32		1			
07335160	**Low discrimination**							**07335160**
					COC			
Clostridium perfringens		59.01	0.43	XYL 1	1			
Atopobium parvulum		40.08	0.40		99			
Actinomyces israelii		0.66	0.23		1			
07335162	**Low discrimination**							**07335162**
					COC			
Clostridium perfringens		71.84	0.38	XYL 1	1			
Atopobium parvulum		18.30	0.27		99			
Lactobacillus acidophilus		5.34	0.16		1			
07335300	**Acceptable identification**							**07335300**
					COC			
Atopobium parvulum		94.57	0.53	RHA 1	99			
Eubacterium contortum		4.48	0.29		1			
Actinomyces israelii		0.84	0.31		1			
07335302	**Low discrimination**							**07335302**
					COC			
Atopobium parvulum		73.99	0.40	RHA 1	99			
Clostridium clostridioforme		11.03	0.30		1			
Actinomyces israelii		5.57	0.34		1			
07335312	**Acceptable identification**							**07335312**
					GRA			
Clostridium clostridioforme		92.57	0.41	RAF 94	50			
Parabacteroides distasonis		2.80	0.21	ARA 94	1			
Bifidobacterium adolescentis		1.72	0.15		99			
07335320	**Low discrimination**							**07335320**
					COC	GRA		
Atopobium parvulum		78.67	0.40	RHA 1	99	99		
Bacteroides vulgatus		8.46	0.15		1	1		
Clostridium perfringens		6.14	0.21		1	99		
07335322	**Low discrimination**							**07335322**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		88.48	0.60	RAF 99	1	1	
Atopobium parvulum		4.13	0.27		99	99	
Clostridium clostridioforme		3.49	0.30		1	50	
07335332	**Low discrimination**						**07335332**
Parabacteroides distasonis		74.98	0.55	RAF 99			
Clostridium clostridioforme		25.02	0.41				
07335400	**Low discrimination**						**07335400**
					COC	SPO	
Actinomyces israelii		56.86	0.39	CEL 93	1	1	
Atopobium parvulum		23.88	0.18		99	1	
Clostridium butyricum		11.23	0.06		1	99	
07335402	**Low discrimination**						**07335402**
Actinomyces israelii		50.67	0.42	CEL 93			
Clostridium clostridioforme		23.58	0.27				
Bifidobacterium adolescentis		17.10	0.27				
07335412	**Low discrimination**						**07335412**
Clostridium clostridioforme		79.05	0.38	RAF 94			
Bifidobacterium adolescentis		15.38	0.29	RHA 88			
Actinomyces israelii		5.57	0.28				
07335432	**Good identification**						**07335432**
Clostridium clostridioforme		100.0	0.38	RAF 94			
07335500	**Low discrimination**						**07335500**
					COC	SPO	
Atopobium parvulum		81.43	0.53	ARA 1	99	1	
Clostridium butyricum		9.29	0.29		1	99	
Actinomyces israelii		7.83	0.49		1	1	
07335502	**Low discrimination**						**07335502**
					COC		
Bifidobacterium adolescentis		73.84	0.61	RAF 92	1		
Atopobium parvulum		10.81	0.40		99		
Actinomyces israelii		8.84	0.52		1		
07335512	**Low discrimination**						**07335512**
Bifidobacterium adolescentis		83.70	0.63	RAF 92			
Clostridium clostridioforme		14.55	0.47				
Actinomyces israelii		1.23	0.38				
07335520	**Low discrimination**						**07335520**
					COC	GRA	
Atopobium parvulum		71.16	0.40	ARA 1	99	99	
Bacteroides vulgatus		11.29	0.18		1	1	
Clostridium perfringens		5.56	0.21		1	99	
07335522	**Low discrimination**						**07335522**
					COC	GRA	
Parabacteroides distasonis		50.91	0.48	RAF 99	1	1	
Clostridium clostridioforme		21.00	0.36		1	50	
Atopobium parvulum		11.63	0.27		99	99	
07335532	**Low discrimination**						**07335532**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		69.83	0.47	RAF 94	50		
Parabacteroides distasonis		20.01	0.43	RHA 88	1		
Bifidobacterium longum subsp. longum		5.72	0.30		99		
07335540	**Low discrimination**						**07335540**
					COC	SPO	
Actinomyces israelii		63.48	0.42	CEL 93	1	1	
Atopobium parvulum		17.14	0.18		99	1	
Clostridium butyricum		16.83	0.11		1	99	
07335542	**Low discrimination**						**07335542**
					SPO		
Bifidobacterium adolescentis		70.70	0.46	RAF 92	1		
Actinomyces israelii		26.63	0.45	SOR 11	1		
Clostridium butyricum		1.83	0.04		99		
07335552	**Acceptable identification**						**07335552**
Bifidobacterium adolescentis		94.32	0.47	RAF 92			
Actinomyces israelii		4.34	0.31	SOR 11			
Clostridium clostridioforme		1.34	0.13				
07335602	**Good identification**						**07335602**
Clostridium clostridioforme		97.84	0.41	RAF 94			
Actinomyces israelii		2.16	0.22				
07335612	**Good identification**						**07335612**
Clostridium clostridioforme		100.0	0.53	RAF 94			
07335622	**Good identification**						**07335622**
Clostridium clostridioforme		96.76	0.41	RAF 94			
Parabacteroides distasonis		3.25	0.22				
07335632	**Good identification**						**07335632**
Clostridium clostridioforme		99.61	0.53	RAF 94			
Parabacteroides distasonis		0.40	0.17				
07335702	**Acceptable identification**						**07335702**
Clostridium clostridioforme		91.73	0.50	RAF 94			
Bifidobacterium adolescentis		2.71	0.27				
Actinomyces israelii		2.42	0.33				
07335712	**Very good identification**						**07335712**
					GRA		
Clostridium clostridioforme		99.01	0.62	RAF 94	50		
Bifidobacterium adolescentis		0.79	0.29		99		
Parabacteroides distasonis		0.12	0.17		1		
07335722	**Low discrimination**						**07335722**
Clostridium clostridioforme		49.92	0.50	RAF 94			
Parabacteroides distasonis		49.51	0.56				
Bacteroides vulgatus		0.31	0.00				
07335732	**Low discrimination**						**07335732**
Clostridium clostridioforme		89.51	0.62	RAF 94			
Parabacteroides distasonis		10.49	0.51				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07337000	**Low discrimination**						**07337000**
					COC	CAT	
Actinomyces israelii		86.98	0.53	CEL 93	1	1	
Atopobium parvulum		5.28	0.18		99	1	
Propionibacterium granulosum		2.81	0.22		1	90	
07337001	**Low discrimination**						**07337001**
Actinomyces naeslundii		78.81	0.38	RAF 92			
Actinomyces israelii		21.19	0.19	XYL 1			
07337002	**Acceptable identification**						**07337002**
					SPO		
Actinomyces israelii		94.01	0.56	CEL 93	1		
Actinomyces naeslundii		1.57	0.35		1		
Clostridium tertium		1.54	0.11		99		
07337003	**Low discrimination**						**07337003**
Actinomyces naeslundii		71.27	0.38	RAF 92			
Actinomyces israelii		28.74	0.22	XYL 1			
07337010	**Low discrimination**						**07337010**
					SPO		
Actinomyces israelii		88.32	0.39	CEL 93	1		
Clostridium tertium		8.58	0.07	MLZ 13	99		
Lactobacillus jensenii		3.10	0.02		1		
07337012	**Low discrimination**						**07337012**
					SPO		
Actinomyces israelii		80.45	0.42	CEL 93	1		
Clostridium tertium		15.63	0.15	MLZ 13	99		
Bifidobacterium adolescentis		3.92	0.13		1		
07337021	**Good identification**						**07337021**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07337022	**Low discrimination**						**07337022**
					SPO		
Actinomyces israelii		47.97	0.36	CEL 93	1		
Clostridium tertium		36.82	0.19	NAG 6	99		
Actinomyces naeslundii		12.55	0.35		1		
07337023	**Good identification**						**07337023**
Actinomyces naeslundii		100.0	0.38	RAF 92			
07337040	**Good identification**						**07337040**
Actinomyces israelii		100.0	0.46	CEL 93			
07337042	**Good identification**						**07337042**
Actinomyces israelii		100.0	0.49	CEL 93			
07337100	**Low discrimination**						**07337100**
					COC	SPO	
Atopobium parvulum		55.04	0.53	MAN 1	99	1	
Actinomyces israelii		36.67	0.63		1	1	
Clostridium tertium		7.41	0.37		1	99	
07337101	**Low discrimination**						**07337101**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		49.48	0.46	RAF 92	1		
Atopobium parvulum		28.06	0.18	XYL 1	99		
Actinomyces israelii		18.69	0.29		1		
07337102	**Low discrimination**						**07337102**
					COC	SPO	
Actinomyces israelii		57.31	0.66	CEL 93	1	1	
Clostridium tertium		23.17	0.45		1	99	
Atopobium parvulum		10.12	0.40		99	1	
07337103	**Low discrimination**						**07337103**
					SPO		
Actinomyces naeslundii		50.52	0.46	RAF 92	1		
Actinomyces israelii		28.63	0.32	XYL 1	1		
Clostridium tertium		11.57	0.11		99		
07337110	**Low discrimination**						**07337110**
					COC	SPO	
Clostridium tertium		68.23	0.41	CEL 99	1	99	
Actinomyces israelii		28.37	0.49		1	1	
Atopobium parvulum		2.88	0.18		99	1	
07337112	**Low discrimination**						**07337112**
					SPO		
Clostridium tertium		68.44	0.50	CEL 99	99		
Bifidobacterium adolescentis		17.17	0.47		1		
Actinomyces israelii		14.23	0.52		1		
07337120	**Low discrimination**						**07337120**
					COC	SPO	
Clostridium tertium		62.02	0.45	CEL 99	1	99	
Atopobium parvulum		27.09	0.40		99	1	
Actinomyces israelii		6.53	0.43		1	1	
07337121	**Low discrimination**						**07337121**
					COC	SPO	
Actinomyces naeslundii		75.24	0.46	RAF 92	1	1	
Clostridium tertium		17.24	0.11	XYL 1	1	99	
Atopobium parvulum		7.53	0.06		99	1	
07337122	**Acceptable identification**						**07337122**
					COC	SPO	
Clostridium tertium		90.18	0.53	CEL 99	1	99	
Actinomyces israelii		4.75	0.46		1	1	
Atopobium parvulum		2.32	0.27		99	1	
07337123	**Low discrimination**						**07337123**
					SPO		
Actinomyces naeslundii		59.27	0.46	RAF 92	1		
Clostridium tertium		40.73	0.19	XYL 1	99		
07337130	**Good identification**						**07337130**
					COC	SPO	
Clostridium tertium		98.88	0.50	CEL 99	1	99	
Actinomyces israelii		0.88	0.29		1	1	
Atopobium parvulum		0.25	0.06		99	1	
07337132	**Very good identification**						**07337132**
					SPO		
Clostridium tertium		99.42	0.58	CEL 99	99		
Actinomyces israelii		0.44	0.32		1		
Bifidobacterium adolescentis		0.08	0.13		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07337140	**Very good identification**						**07337140**
					COC		
Actinomyces israelii		95.03	0.56	CEL 93	1		
Atopobium parvulum		3.71	0.18		99		
Actinomyces naeslundii		0.77	0.30		1		
07337142	**Good identification**						**07337142**
					SPO		
Actinomyces israelii		93.47	0.59	CEL 93	1		
Bifidobacterium adolescentis		4.38	0.30		1		
Clostridium tertium		0.98	0.11		99		
07337150	**Acceptable identification**						**07337150**
					SPO		
Actinomyces israelii		94.12	0.42	CEL 93	1		
Clostridium tertium		5.88	0.07	MLZ 13	99		
07337152	**Low discrimination**						**07337152**
					SPO		
Actinomyces israelii		66.30	0.45	CEL 93	1		
Bifidobacterium adolescentis		25.42	0.32	MLZ 13	1		
Clostridium tertium		8.28	0.15		99		
07337160	**Low discrimination**						**07337160**
					SPO		
Actinomyces israelii		60.98	0.36	CEL 93	1		
Clostridium tertium		15.05	0.11	NAG 6	99		
Clostridium perfringens		9.68	0.09		84		
07337162	**Low discrimination**						**07337162**
					SPO		
Actinomyces israelii		61.44	0.39	CEL 93	1		
Clostridium tertium		30.32	0.19	NAG 6	99		
Actinomyces naeslundii		5.19	0.30		1		
07337202	**Good identification**						**07337202**
Actinomyces israelii		100.0	0.37	CEL 93			
07337300	**Low discrimination**						**07337300**
					COC	SPO	
Actinomyces israelii		81.40	0.44	CEL 93	1	1	
Atopobium parvulum		16.40	0.18	RHA 7	99	1	
Clostridium tertium		2.21	0.03		1	99	
07337302	**Acceptable identification**						**07337302**
					COC	SPO	
Actinomyces israelii		91.10	0.47	CEL 93	1	1	
Clostridium tertium		4.94	0.11	RHA 7	1	99	
Atopobium parvulum		2.16	0.06		99	1	
07337340	**Good identification**						**07337340**
Actinomyces israelii		100.0	0.37	CEL 93			
07337342	**Good identification**						**07337342**
Actinomyces israelii		100.0	0.40	CEL 93			
07337400	**Good identification**						**07337400**
Actinomyces israelii		100.0	0.52	CEL 93			
07337402	**Good identification**						**07337402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		95.35	0.55	CEL 93			
Bifidobacterium adolescentis		4.65	0.26				
07337410	**Good identification**						**07337410**
Actinomyces israelii		100.0	0.37	CEL 93			
07337412	**Low discrimination**						**07337412**
Actinomyces israelii		71.50	0.40	CEL 93			
Bifidobacterium adolescentis		28.50	0.27	MLZ 13			
07337440	**Good identification**						**07337440**
Actinomyces israelii		100.0	0.45	CEL 93			
07337442	**Good identification**						**07337442**
Actinomyces israelii		98.48	0.48	CEL 93			
Bifidobacterium adolescentis		1.53	0.10				
07337500	**Good identification**						**07337500**
					COC	SPO	
Actinomyces israelii		94.11	0.62	CEL 93	1	1	
Clostridium butyricum		2.19	0.13		1	99	
Atopobium parvulum		1.74	0.18		99	1	
07337502	**Low discrimination**						**07337502**
					SPO		
Bifidobacterium adolescentis		54.35	0.60	RAF 92	1		
Actinomyces israelii		45.06	0.65		1		
Clostridium butyricum		0.27	0.06		99		
07337510	**Low discrimination**						**07337510**
					SPO		
Actinomyces israelii		84.80	0.48	CEL 93	1		
Bifidobacterium adolescentis		12.68	0.27	MLZ 13	1		
Clostridium tertium		2.52	0.07		99		
07337512	**Low discrimination**						**07337512**
Bifidobacterium adolescentis		89.64	0.61	RAF 92			
Actinomyces israelii		9.09	0.51				
Bifidobacterium dentium		0.55	0.06				
07337520	**Low discrimination**						**07337520**
					COC	SPO	
Actinomyces israelii		83.32	0.41	CEL 93	1	1	
Clostridium tertium		9.77	0.11	NAG 6	1	99	
Atopobium parvulum		4.27	0.06		99	1	
07337522	**Low discrimination**						**07337522**
					SPO		
Actinomyces israelii		68.88	0.44	CEL 93	1		
Clostridium tertium		16.16	0.19	NAG 6	99		
Bifidobacterium adolescentis		13.15	0.26		1		
07337540	**Good identification**						**07337540**
					SPO		
Actinomyces israelii		98.91	0.55	CEL 93	1		
Bifidobacterium adolescentis		0.58	0.10		1		
Clostridium butyricum		0.51	-.05		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07337542	**Low discrimination**					**07337542**
Actinomyces israelii		72.29	0.58	CEL 93		
Bifidobacterium adolescentis		27.71	0.44			
07337550	**Acceptable identification**					**07337550**
					COC	
Actinomyces israelii		91.62	0.41	CEL 93	1	
Bifidobacterium adolescentis		4.35	0.12	MLZ 13	1	
Blautia producta		4.03	-.03		99	
07337552	**Low discrimination**					**07337552**
Bifidobacterium adolescentis		75.82	0.46	RAF 92		
Actinomyces israelii		24.18	0.44	SOR 11		
07337562	**Acceptable identification**					**07337562**
Actinomyces israelii		94.28	0.37	CEL 93		
Bifidobacterium adolescentis		5.72	0.10	NAG 6		
07337602	**Acceptable identification**					**07337602**
Actinomyces israelii		92.52	0.35	CEL 93		
Clostridium clostridioforme		7.48	0.07	RHA 7		
07337700	**Good identification**					**07337700**
Actinomyces israelii		100.0	0.43	CEL 93		
07337702	**Low discrimination**					**07337702**
Actinomyces israelii		81.33	0.46	CEL 93		
Bifidobacterium adolescentis		13.17	0.26	RHA 7		
Clostridium clostridioforme		5.50	0.16			
07337740	**Good identification**					**07337740**
Actinomyces israelii		100.0	0.36	CEL 93		
07337742	**Good identification**					**07337742**
Actinomyces israelii		95.11	0.39	CEL 93		
Bifidobacterium adolescentis		4.89	0.10	RHA 7		
07341000	**Low discrimination**					**07341000**
					SPO	
Lactobacillus catenaformis		65.22	0.63	FRU 99	1	
Lactobacillus jensenii		32.74	0.58		1	
Clostridium barati		1.09	0.27		99	
07341002	**Low discrimination**					**07341002**
Lactobacillus catenaformis		69.13	0.63	FRU 99		
Lactobacillus jensenii		23.14	0.55			
Lactobacillus acidophilus		4.00	0.39			
07341010	**Acceptable identification**					**07341010**
Lactobacillus jensenii		94.31	0.50	FRU 99		
Lactobacillus catenaformis		5.69	0.29			
07341012	**Acceptable identification**					**07341012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		90.88	0.47	FRU 99		
Lactobacillus catenaformis		8.23	0.29			
Lactobacillus acidophilus		0.48	0.05			
07341020	**Low discrimination**					**07341020**
					SPO	
Lactobacillus catenaformis		57.95	0.53	FRU 99	1	
Lactobacillus jensenii		29.10	0.47		1	
Clostridium barati		11.65	0.35		99	
07341021	**Low discrimination**					**07341021**
Actinomyces naeslundii		44.93	0.42	RAF 92		
Lactobacillus catenaformis		32.33	0.18	FRU 95		
Lactobacillus jensenii		16.23	0.13			
07341022	**Low discrimination**					**07341022**
					SPO	
Lactobacillus catenaformis		45.61	0.53	FRU 99	1	
Clostridium barati		27.52	0.44		99	
Lactobacillus jensenii		15.27	0.44		1	
07341023	**Low discrimination**					**07341023**
					SPO	
Actinomyces naeslundii		39.05	0.42	RAF 92	1	
Lactobacillus catenaformis		28.10	0.18	FRU 95	1	
Clostridium barati		16.95	0.09		99	
07341030	**Acceptable identification**					**07341030**
					SPO	
Lactobacillus jensenii		93.24	0.39	FRU 99	1	
Lactobacillus catenaformis		5.63	0.18		1	
Clostridium barati		1.13	0.01		99	
07341032	**Low discrimination**					**07341032**
					SPO	
Lactobacillus jensenii		85.76	0.36	FRU 99	1	
Lactobacillus catenaformis		7.76	0.18		1	
Clostridium barati		4.68	0.09		99	
07341100	**Low discrimination**					**07341100**
					SPO	
Lactobacillus catenaformis		88.79	0.66	FRU 99	1	
Collinsella aerofaciens		6.53	0.32		1	
Clostridium barati		2.98	0.35		99	
07341102	**Low discrimination**					**07341102**
					SPO	
Lactobacillus catenaformis		64.04	0.66	FRU 99	1	
Lactobacillus acidophilus		28.39	0.57		1	
Clostridium barati		6.44	0.44		99	
07341120	**Low discrimination**					**07341120**
					SPO	
Lactobacillus catenaformis		58.66	0.56	FRU 99	1	
Clostridium barati		23.59	0.44		99	
Clostridium paraputrificum		9.66	0.32		99	
07341121	**Low discrimination**					**07341121**
					SPO	
Actinomyces naeslundii		58.39	0.50	RAF 92	1	
Lactobacillus catenaformis		22.14	0.21	FRU 95	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		8.91	0.09		99		
07341122	**Low discrimination**						**07341122**
					SPO		
Lactobacillus acidophilus		42.13	0.57	FRU 99	1		
Clostridium barati		28.67	0.52		99		
Lactobacillus catenaformis		23.77	0.56		1		
07341123	**Low discrimination**						**07341123**
					SPO		
Actinomyces naeslundii		38.81	0.50	RAF 92	1		
Lactobacillus acidophilus		26.09	0.23	FRU 95	1		
Clostridium barati		17.76	0.18		99		
07341141	**Good identification**						**07341141**
Actinomyces naeslundii		100.0	0.37	RAF 92			
07341142	**Low discrimination**						**07341142**
					SPO		
Lactobacillus acidophilus		73.62	0.41	FRU 99	1		
Lactobacillus catenaformis		13.58	0.32	SOR 11	1		
Clostridium barati		11.75	0.25		99		
07341143	**Low discrimination**						**07341143**
Actinomyces naeslundii		67.99	0.37	RAF 92			
Lactobacillus acidophilus		32.01	0.07	FRU 95			
07341161	**Good identification**						**07341161**
Actinomyces naeslundii		100.0	0.37	RAF 92			
07341162	**Low discrimination**						**07341162**
					SPO		
Lactobacillus acidophilus		63.20	0.41	FRU 99	1		
Clostridium barati		30.26	0.34	SOR 11	99		
Lactobacillus catenaformis		2.91	0.21		1		
07341163	**Low discrimination**						**07341163**
					SPO		
Actinomyces naeslundii		58.95	0.37	RAF 92	1		
Lactobacillus acidophilus		27.76	0.07	FRU 95	1		
Clostridium barati		13.29	-.01		99		
07341302	**Low discrimination**						**07341302**
					SPO		
Lactobacillus acidophilus		63.17	0.38	FRU 99	1		
Lactobacillus catenaformis		19.13	0.32	RHA 7	1		
Clostridium barati		16.56	0.25		99		
07341322	**Low discrimination**						**07341322**
					GRA	SPO	
Lactobacillus acidophilus		43.22	0.38	FRU 99	99	1	
Clostridium barati		33.98	0.34	RHA 7	99	99	
Prevotella oralis		14.95	0.29		1	1	
07343000	**Low discrimination**						**07343000**
					SPO		
Clostridium innocuum		84.11	0.44	MLT 1	99		
Lactobacillus catenaformis		7.58	0.29	FRU 95	1		
Lactobacillus jensenii		3.81	0.24		1		
07343001	**Low discrimination**						**07343001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		85.75	0.47	RAF 92	1	
Clostridium innocuum		14.25	0.10	FRU 95	99	
07343002	**Low discrimination**					**07343002**
					SPO	
Clostridium innocuum		81.36	0.44	MLT 1	99	
Lactobacillus catenaformis		7.33	0.29	FRU 95	1	
Actinomyces naeslundii		3.30	0.44		1	
07343003	**Low discrimination**					**07343003**
					SPO	
Actinomyces naeslundii		85.75	0.47	RAF 92	1	
Clostridium innocuum		14.25	0.10	FRU 95	99	
07343020	**Low discrimination**					**07343020**
					SPO	
Actinomyces naeslundii		32.82	0.44	RAF 92	1	
Clostridium barati		31.59	0.17	FRU 95	99	
Lactobacillus catenaformis		18.25	0.18		1	
07343021	**Good identification**					**07343021**
Actinomyces naeslundii		100.0	0.47	RAF 92		
07343023	**Good identification**					**07343023**
Actinomyces naeslundii		100.0	0.47	RAF 92		
07343101	**Good identification**					**07343101**
Actinomyces naeslundii		100.0	0.55	RAF 92		
07343102	**Low discrimination**					**07343102**
					SPO	
Lactobacillus acidophilus		39.67	0.35	FRU 99	1	
Lactobacillus catenaformis		17.18	0.32	MAN 5	1	
Clostridium barati		14.87	0.25		99	
07343103	**Good identification**					**07343103**
Actinomyces naeslundii		98.21	0.55	RAF 92		
Lactobacillus acidophilus		1.79	0.01	FRU 95		
07343121	**Good identification**					**07343121**
Actinomyces naeslundii		100.0	0.55	RAF 92		
07343123	**Acceptable identification**					**07343123**
					SPO	
Actinomyces naeslundii		94.85	0.55	RAF 92	1	
Clostridium barati		1.94	-.01	FRU 95	99	
Lactobacillus acidophilus		1.73	0.01		1	
07343132	**Good identification**					**07343132**
					SPO	
Clostridium tertium		96.82	0.39	FRU 99	99	
Actinomyces naeslundii		1.24	0.30	TRE 90	1	
Clostridium barati		0.73	-.01		99	
07343140	**Low discrimination**					**07343140**
					SPO	
Actinomyces naeslundii		78.47	0.39	RAF 92	1	
Clostridium barati		13.08	-.01	FRU 95	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		8.46	0.13		1	
07343141	**Good identification**					**07343141**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07343143	**Good identification**					**07343143**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07343160	**Low discrimination**					**07343160**
					SPO	
Actinomyces naeslundii		66.67	0.39	RAF 92	1	
Clostridium barati		33.33	0.07	FRU 95	99	
07343161	**Good identification**					**07343161**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07343163	**Good identification**					**07343163**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07345502	**Low discrimination**					**07345502**
					SPO	
Bifidobacterium adolescentis		82.95	0.36	RAF 92	1	
Actinomyces israelii		11.00	0.27	FRU 99	1	
Clostridium butyricum		6.05	0.02		99	
07345512	**Good identification**					**07345512**
Bifidobacterium adolescentis		97.57	0.37	RAF 92		
Clostridium clostridioforme		2.43	0.07	FRU 99		
07347100	**Low discrimination**					**07347100**
					SPO	
Actinomyces israelii		61.37	0.39	FRU 99	1	
Clostridium tertium		38.63	0.21		99	
07347122	**Good identification**					**07347122**
					SPO	
Clostridium tertium		96.61	0.37	FRU 99	99	
Actinomyces israelii		1.63	0.21	TRE 90	1	
Clostridium barati		0.93	-.01		99	
07347132	**Good identification**					**07347132**
Clostridium tertium		100.0	0.41	FRU 99		
07347500	**Good identification**					**07347500**
					SPO	
Actinomyces israelii		96.01	0.37	FRU 99	1	
Clostridium butyricum		3.99	-.08		99	
07347512	**Low discrimination**					**07347512**
					SPO	
Bifidobacterium adolescentis		88.26	0.36	RAF 92	1	
Actinomyces israelii		9.91	0.26	FRU 99	1	
Clostridium tertium		1.83	-.01		99	
07350000	**Low discrimination**					**07350000**
					GRA	
Lactobacillus catenaformis		56.47	0.63	GAL 99	99	
Lactobacillus jensenii		28.35	0.58		99	
Fusobacterium mortiferum		8.92	0.59		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07350002	**Low discrimination**						**07350002**
					SPO		
Lactobacillus catenaformis		59.50	0.63	GAL 99	1		
Lactobacillus jensenii		19.92	0.55		1		
Clostridium barati		15.59	0.48		99		
07350010	**Acceptable identification**						**07350010**
					GRA		
Lactobacillus jensenii		93.01	0.50	GAL 99	99		
Lactobacillus catenaformis		5.61	0.29		99		
Fusobacterium mortiferum		0.89	0.25		1		
07350012	**Low discrimination**						**07350012**
					SPO		
Lactobacillus jensenii		89.33	0.47	GAL 99	1		
Lactobacillus catenaformis		8.09	0.29		1		
Clostridium barati		2.12	0.14		99		
07350020	**Low discrimination**						**07350020**
					SPO		
Clostridium barati		40.60	0.48	ESL 87	99		
Lactobacillus catenaformis		38.75	0.53	GAL 95	1		
Lactobacillus jensenii		19.46	0.47		1		
07350021	**Low discrimination**						**07350021**
					SPO		
Actinomyces naeslundii		30.14	0.41	RAF 92	1		
Clostridium barati		28.71	0.14	GAL 96	99		
Lactobacillus catenaformis		27.40	0.18		1		
07350022	**Low discrimination**						**07350022**
					SPO		
Clostridium barati		66.45	0.56	ESL 87	99		
Lactobacillus catenaformis		21.14	0.53	GAL 95	1		
Lactobacillus jensenii		7.08	0.44		1		
07350030	**Low discrimination**						**07350030**
					SPO		
Lactobacillus jensenii		89.00	0.39	GAL 99	1		
Clostridium barati		5.63	0.14		99		
Lactobacillus catenaformis		5.37	0.18		1		
07350032	**Low discrimination**						**07350032**
					SPO		
Lactobacillus jensenii		71.63	0.36	GAL 99	1		
Clostridium barati		20.39	0.22		99		
Lactobacillus catenaformis		6.49	0.18		1		
07350062	**Low discrimination**						**07350062**
					SPO		
Clostridium barati		86.31	0.38	ESL 87	99		
Lactobacillus acidophilus		9.03	0.23	GAL 95	1		
Lactobacillus catenaformis		3.19	0.18	SOR 8	1		
07350100	**Low discrimination**						**07350100**
					GRA	SPO	
Lactobacillus catenaformis		66.29	0.66	GAL 99	99	1	
Fusobacterium mortiferum		16.29	0.65		1	1	
Clostridium barati		11.58	0.48		99	99	
07350102	**Low discrimination**						**07350102**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		49.58	0.66	GAL 99	1	
Clostridium barati		25.97	0.56		99	
Lactobacillus acidophilus		21.97	0.57		1	
07350120	**Low discrimination**					**07350120**
					SPO	
Clostridium barati		61.81	0.56	ESL 87	99	
Lactobacillus catenaformis		29.50	0.56	GAL 95	1	
Clostridium paraputrificum		4.86	0.32		99	
07350122	**Low discrimination**					**07350122**
					SPO	
Clostridium barati		67.73	0.64	ESL 87	99	
Lactobacillus acidophilus		19.10	0.57	GAL 95	1	
Lactobacillus catenaformis		10.77	0.56		1	
07350141	**Good identification**					**07350141**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07350142	**Low discrimination**					**07350142**
					SPO	
Lactobacillus acidophilus		49.20	0.41	GAL 99	1	
Clostridium barati		40.92	0.38	SOR 11	99	
Lactobacillus catenaformis		9.07	0.32		1	
07350143	**Low discrimination**					**07350143**
					SPO	
Actinomyces naeslundii		47.86	0.36	RAF 92	1	
Lactobacillus acidophilus		28.47	0.07	GAL 96	1	
Clostridium barati		23.67	0.04		99	
07350160	**Low discrimination**					**07350160**
					SPO	
Clostridium barati		85.09	0.38	ESL 87	99	
Clostridium paraputrificum		5.79	0.13	GAL 95	99	
Lactobacillus catenaformis		4.72	0.21	SOR 8	1	
07350161	**Low discrimination**					**07350161**
					SPO	
Actinomyces naeslundii		66.91	0.36	RAF 92	1	
Clostridium barati		33.09	0.04	GAL 96	99	
07350162	**Low discrimination**					**07350162**
					SPO	
Clostridium barati		69.39	0.46	ESL 87	99	
Lactobacillus acidophilus		27.81	0.41	GAL 95	1	
Lactobacillus catenaformis		1.28	0.21	SOR 8	1	
07350222	**Low discrimination**					**07350222**
					SPO	
Clostridium barati		88.56	0.38	ESL 87	99	
Lactobacillus acidophilus		5.64	0.20	GAL 95	1	
Lactobacillus catenaformis		3.27	0.18	RHA 8	1	
07350302	**Low discrimination**					**07350302**
					SPO	
Clostridium barati		50.67	0.38	ESL 87	99	
Lactobacillus acidophilus		37.11	0.38	GAL 95	1	
Lactobacillus catenaformis		11.24	0.32	RHA 8	1	
07350320	**Acceptable identification**					**07350320**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		90.60	0.38	ESL 87	99		
Lactobacillus catenaformis		5.02	0.21	GAL 95	1		
Clostridium ramosum		1.67	0.09	RHA 8	92		
07350322	**Low discrimination**						**07350322**
					GRA	SPO	
Clostridium barati		72.84	0.46	ESL 87	99	99	
Lactobacillus acidophilus		17.78	0.38	GAL 95	99	1	
Prevotella oralis		6.15	0.29	RHA 8	1	1	
07351000	**Low discrimination**						**07351000**
					SPO		
Lactobacillus catenaformis		65.63	0.97		1		
Lactobacillus jensenii		32.95	0.92		1		
Clostridium barati		1.10	0.61		99		
07351001	**Low discrimination**						**07351001**
Lactobacillus catenaformis		62.91	0.63	URE 1			
Lactobacillus jensenii		31.58	0.58				
Actinomyces naeslundii		4.20	0.64				
07351002	**Low discrimination**						**07351002**
Lactobacillus catenaformis		69.25	0.97				
Lactobacillus jensenii		23.18	0.89				
Lactobacillus acidophilus		4.00	0.73				
07351003	**Low discrimination**						**07351003**
Lactobacillus catenaformis		66.19	0.63	URE 1			
Lactobacillus jensenii		22.15	0.55				
Actinomyces naeslundii		4.41	0.64				
07351010	**Good identification**						**07351010**
					SPO		
Lactobacillus jensenii		94.18	0.84		1		
Lactobacillus catenaformis		5.68	0.63		1		
Clostridium barati		0.10	0.27		99		
07351011	**Acceptable identification**						**07351011**
Lactobacillus jensenii		92.48	0.50	URE 1			
Lactobacillus catenaformis		5.58	0.29				
Actinomyces naeslundii		1.94	0.42				
07351012	**Good identification**						**07351012**
Lactobacillus jensenii		90.84	0.81				
Lactobacillus catenaformis		8.23	0.63				
Lactobacillus acidophilus		0.48	0.39				
07351013	**Low discrimination**						**07351013**
Lactobacillus jensenii		88.36	0.47	URE 1			
Lactobacillus catenaformis		8.00	0.29				
Actinomyces naeslundii		2.78	0.42				
07351020	**Low discrimination**						**07351020**
					SPO		
Lactobacillus catenaformis		58.26	0.87		1		
Lactobacillus jensenii		29.25	0.82		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		11.72	0.70		99	
07351021	**Low discrimination**					**07351021**
Lactobacillus catenaformis		50.51	0.53	URE 1		
Lactobacillus jensenii		25.36	0.47			
Actinomyces naeslundii		13.47	0.64			
07351022	**Low discrimination**					**07351022**
					SPO	
Lactobacillus catenaformis		45.77	0.87		1	
Clostridium barati		27.61	0.78		99	
Lactobacillus jensenii		15.32	0.79		1	
07351023	**Low discrimination**					**07351023**
					SPO	
Lactobacillus catenaformis		40.82	0.53	URE 1	1	
Clostridium barati		24.63	0.44		99	
Lactobacillus jensenii		13.66	0.44		1	
07351030	**Good identification**					**07351030**
					SPO	
Lactobacillus jensenii		93.14	0.73		1	
Lactobacillus catenaformis		5.62	0.53		1	
Clostridium barati		1.13	0.35		99	
07351031	**Low discrimination**					**07351031**
Lactobacillus jensenii		86.48	0.39	URE 1		
Actinomyces naeslundii		7.25	0.42			
Lactobacillus catenaformis		5.22	0.18			
07351032	**Low discrimination**					**07351032**
					SPO	
Lactobacillus jensenii		85.58	0.70		1	
Lactobacillus catenaformis		7.75	0.53		1	
Clostridium barati		4.67	0.44		99	
07351033	**Low discrimination**					**07351033**
Lactobacillus jensenii		77.41	0.36	URE 1		
Actinomyces naeslundii		9.74	0.42			
Lactobacillus catenaformis		7.01	0.18			
07351040	**Low discrimination**					**07351040**
					SPO	
Lactobacillus catenaformis		60.00	0.63	SOR 1	1	
Lactobacillus jensenii		30.12	0.58		1	
Clostridium barati		8.66	0.43		99	
07351041	**Low discrimination**					**07351041**
Actinomyces naeslundii		41.44	0.51	RAF 92		
Lactobacillus catenaformis		35.57	0.29			
Lactobacillus jensenii		17.86	0.24			
07351042	**Low discrimination**					**07351042**
					SPO	
Lactobacillus catenaformis		40.24	0.63	SOR 1	1	
Lactobacillus acidophilus		28.47	0.57		1	
Clostridium barati		17.42	0.51		99	
07351043	**Low discrimination**					**07351043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		32.00	0.51	RAF 92		
Lactobacillus catenaformis		27.48	0.29			
Lactobacillus acidophilus		19.44	0.23			
07351050	**Acceptable identification**					**07351050**
					SPO	
Lactobacillus jensenii		93.32	0.50	SOR 1	1	
Lactobacillus catenaformis		5.63	0.29		1	
Clostridium barati		0.81	0.09		99	
07351052	**Low discrimination**					**07351052**
Lactobacillus jensenii		83.36	0.47	SOR 1		
Lactobacillus catenaformis		7.55	0.29			
Lactobacillus acidophilus		5.34	0.23			
07351060	**Low discrimination**					**07351060**
					SPO	
Clostridium barati		50.80	0.51	ESL 87	99	
Lactobacillus catenaformis		29.34	0.53	SOR 8	1	
Lactobacillus jensenii		14.73	0.47		1	
07351061	**Low discrimination**					**07351061**
					SPO	
Actinomyces naeslundii		58.32	0.51	RAF 92	1	
Clostridium barati		21.67	0.17		99	
Lactobacillus catenaformis		12.52	0.18		1	
07351062	**Low discrimination**					**07351062**
					SPO	
Clostridium barati		54.93	0.60	ESL 87	99	
Lactobacillus acidophilus		29.93	0.57	SOR 8	1	
Lactobacillus catenaformis		10.58	0.53		1	
07351070	**Low discrimination**					**07351070**
					SPO	
Lactobacillus jensenii		84.71	0.39	SOR 1	1	
Clostridium barati		8.85	0.17		99	
Lactobacillus catenaformis		5.11	0.18		1	
07351072	**Low discrimination**					**07351072**
					SPO	
Lactobacillus jensenii		54.26	0.36	SOR 1	1	
Clostridium barati		25.51	0.25		99	
Lactobacillus acidophilus		13.90	0.23		1	
07351100	**Low discrimination**					**07351100**
					SPO	
Lactobacillus catenaformis		89.06	1.00		1	
Collinsella aerofaciens		6.55	0.66		1	
Clostridium barati		2.99	0.70		99	
07351101	**Low discrimination**					**07351101**
Lactobacillus catenaformis		79.76	0.66	URE 1		
Actinomyces naeslundii		10.09	0.72			
Collinsella aerofaciens		5.86	0.32			
07351102	**Low discrimination**					**07351102**
					SPO	
Lactobacillus catenaformis		64.18	1.00		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		28.45	0.91		1	
Clostridium barati		6.45	0.78		99	
07351103	**Low discrimination**					**07351103**
Lactobacillus catenaformis		59.33	0.66	URE 1		
Lactobacillus acidophilus		26.29	0.57			
Actinomyces naeslundii		7.51	0.72			
07351110	**Low discrimination**					**07351110**
Lactobacillus catenaformis		80.56	0.66	MLZ 1		
Lactobacillus jensenii		8.99	0.50			
Collinsella aerofaciens		5.92	0.32			
07351112	**Low discrimination**					**07351112**
					SPO	
Lactobacillus catenaformis		57.63	0.66	MLZ 1	1	
Lactobacillus acidophilus		25.54	0.57		1	
Clostridium barati		5.79	0.44		99	
07351120	**Low discrimination**					**07351120**
					SPO	
Lactobacillus catenaformis		59.63	0.90		1	
Clostridium barati		23.98	0.78		99	
Clostridium paraputrificum		9.82	0.66		99	
07351121	**Low discrimination**					**07351121**
					SPO	
Lactobacillus catenaformis		45.74	0.56	URE 1	1	
Actinomyces naeslundii		23.15	0.72		1	
Clostridium barati		18.39	0.44		99	
07351122	**Low discrimination**					**07351122**
					SPO	
Lactobacillus acidophilus		42.30	0.91		1	
Clostridium barati		28.79	0.86		99	
Lactobacillus catenaformis		23.86	0.90		1	
07351123	**Low discrimination**					**07351123**
					SPO	
Lactobacillus acidophilus		37.77	0.57	URE 1	1	
Clostridium barati		25.70	0.52		99	
Lactobacillus catenaformis		21.30	0.56		1	
07351130	**Low discrimination**					**07351130**
					SPO	
Lactobacillus catenaformis		53.93	0.56	MLZ 1	1	
Clostridium barati		21.69	0.44		99	
Clostridium paraputrificum		8.88	0.32		99	
07351131	**Low discrimination**					**07351131**
					SPO	
Actinomyces naeslundii		61.11	0.50	RAF 92	1	
Lactobacillus catenaformis		23.17	0.21	MLZ 5	1	
Clostridium barati		9.32	0.09		99	
07351132	**Low discrimination**					**07351132**
					SPO	
Lactobacillus acidophilus		39.17	0.57	MLZ 1	1	
Clostridium barati		26.66	0.52		99	
Lactobacillus catenaformis		22.09	0.56		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07351133	**Low discrimination**					**07351133**
					SPO	
Actinomyces naeslundii		37.18	0.50	RAF 92	1	
Lactobacillus acidophilus		25.00	0.23	MLZ 5	1	
Clostridium barati		17.01	0.18		99	
07351140	**Low discrimination**					**07351140**
					SPO	
Lactobacillus catenaformis		68.86	0.66	SOR 1	1	
Clostridium barati		19.87	0.51		99	
Collinsella aerofaciens		5.06	0.32		1	
07351141	**Low discrimination**					**07351141**
					SPO	
Actinomyces naeslundii		60.94	0.59	RAF 92	1	
Lactobacillus catenaformis		27.57	0.32		1	
Clostridium barati		7.96	0.17		99	
07351142	**Low discrimination**					**07351142**
					SPO	
Lactobacillus acidophilus		73.91	0.76	SOR 11	1	
Lactobacillus catenaformis		13.63	0.66		1	
Clostridium barati		11.80	0.60		99	
07351143	**Low discrimination**					**07351143**
Lactobacillus acidophilus		57.09	0.41	SOR 11		
Actinomyces naeslundii		23.27	0.59	URE 1		
Lactobacillus catenaformis		10.53	0.32			
07351151	**Good identification**					**07351151**
Actinomyces naeslundii		100.0	0.37	RAF 92		
07351152	**Low discrimination**					**07351152**
Lactobacillus acidophilus		60.46	0.41	MLZ 1		
Bifidobacterium breve		12.81	0.40	SOR 11		
Lactobacillus catenaformis		11.15	0.32			
07351153	**Low discrimination**					**07351153**
Actinomyces naeslundii		67.99	0.37	RAF 92		
Lactobacillus acidophilus		32.01	0.07	MLZ 5		
07351160	**Low discrimination**					**07351160**
					SPO	
Clostridium barati		53.18	0.60	ESL 87	99	
Clostridium paraputrificum		18.84	0.47	SOR 8	99	
Lactobacillus catenaformis		15.36	0.56		1	
07351161	**Low discrimination**					**07351161**
					SPO	
Actinomyces naeslundii		58.09	0.59	RAF 92	1	
Clostridium barati		22.75	0.25		99	
Clostridium paraputrificum		8.06	0.13		99	
07351162	**Low discrimination**					**07351162**
					SPO	
Lactobacillus acidophilus		63.63	0.76	SOR 11	1	
Clostridium barati		30.46	0.68		99	
Lactobacillus catenaformis		2.93	0.56		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07351163	**Low discrimination**					**07351163**
					SPO	
Lactobacillus acidophilus		50.62	0.41	SOR 11	1	
Clostridium barati		24.24	0.34	URE 1	99	
Actinomyces naeslundii		20.63	0.59		1	
07351171	**Good identification**					**07351171**
Actinomyces naeslundii		100.0	0.37	RAF 92		
07351172	**Low discrimination**					**07351172**
					SPO	
Lactobacillus acidophilus		50.89	0.41	MLZ 1	1	
Clostridium barati		24.36	0.34	SOR 11	99	
Bifidobacterium breve		19.17	0.44		1	
07351173	**Low discrimination**					**07351173**
					SPO	
Actinomyces naeslundii		53.37	0.37	RAF 92	1	
Lactobacillus acidophilus		25.13	0.07	MLZ 5	1	
Clostridium barati		12.03	-.01		99	
07351200	**Low discrimination**					**07351200**
					SPO	
Lactobacillus catenaformis		60.41	0.63	RHA 1	1	
Lactobacillus jensenii		30.33	0.58		1	
Clostridium barati		8.71	0.43		99	
07351202	**Low discrimination**					**07351202**
					SPO	
Lactobacillus catenaformis		45.34	0.63	RHA 1	1	
Clostridium barati		19.62	0.51		99	
Lactobacillus acidophilus		19.53	0.54		1	
07351210	**Acceptable identification**					**07351210**
					SPO	
Lactobacillus jensenii		93.54	0.50	RHA 1	1	
Lactobacillus catenaformis		5.65	0.29		1	
Clostridium barati		0.81	0.09		99	
07351212	**Low discrimination**					**07351212**
					SPO	
Lactobacillus jensenii		85.24	0.47	RHA 1	1	
Lactobacillus catenaformis		7.72	0.29		1	
Clostridium barati		3.34	0.17		99	
07351220	**Low discrimination**					**07351220**
					SPO	
Clostridium barati		52.65	0.51	ESL 87	99	
Lactobacillus catenaformis		30.41	0.53	RHA 8	1	
Lactobacillus jensenii		15.27	0.47		1	
07351222	**Low discrimination**					**07351222**
					SPO	
Clostridium barati		59.60	0.60	ESL 87	99	
Lactobacillus acidophilus		19.77	0.54	RHA 8	1	
Lactobacillus catenaformis		11.48	0.53		1	
07351230	**Low discrimination**					**07351230**
					SPO	
Lactobacillus jensenii		85.85	0.39	RHA 1	1	
Clostridium barati		8.97	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		5.18	0.18		1		
07351232	**Low discrimination**						**07351232**
					SPO		
Lactobacillus jensenii		56.37	0.36	RHA 1	1		
Clostridium barati		26.50	0.25		99		
Lactobacillus acidophilus		8.79	0.20		1		
07351242	**Low discrimination**						**07351242**
					SPO		
Lactobacillus acidophilus		51.02	0.38	LAC 92	1		
Clostridium barati		36.06	0.33	RHA 7	99		
Lactobacillus catenaformis		9.68	0.29	SOR 11	1		
07351262	**Low discrimination**						**07351262**
					SPO		
Clostridium barati		66.17	0.41	ESL 87	99		
Lactobacillus acidophilus		31.21	0.38	RHA 8	1		
Lactobacillus catenaformis		1.48	0.18	SOR 8	1		
07351300	**Low discrimination**						**07351300**
					SPO		
Lactobacillus catenaformis		70.87	0.66	RHA 1	1		
Clostridium barati		20.45	0.51		99		
Collinsella aerofaciens		5.21	0.32		1		
07351302	**Low discrimination**						**07351302**
					SPO		
Lactobacillus acidophilus		62.97	0.72	RHA 7	1		
Lactobacillus catenaformis		19.06	0.66		1		
Clostridium barati		16.50	0.60		99		
07351303	**Low discrimination**						**07351303**
					SPO		
Lactobacillus acidophilus		61.68	0.38	RHA 7	1		
Lactobacillus catenaformis		18.67	0.32	URE 1	1		
Clostridium barati		16.16	0.25		99		
07351312	**Low discrimination**						**07351312**
					SPO		
Lactobacillus acidophilus		60.17	0.38	RHA 7	1		
Lactobacillus catenaformis		18.22	0.32	MLZ 1	1		
Clostridium barati		15.77	0.25		99		
07351320	**Low discrimination**						**07351320**
					SPO		
Clostridium barati		63.92	0.60	ESL 87	99		
Lactobacillus catenaformis		18.46	0.56	RHA 8	1		
Clostridium ramosum		6.14	0.44		92		
07351322	**Low discrimination**						**07351322**
					GRA	SPO	
Lactobacillus acidophilus		43.17	0.72	RHA 7	99	1	
Clostridium barati		33.95	0.68		99	99	
Prevotella oralis		14.94	0.63		1	1	
07351323	**Low discrimination**						**07351323**
					GRA	SPO	
Lactobacillus acidophilus		42.51	0.38	RHA 7	99	1	
Clostridium barati		33.43	0.34	URE 1	99	99	
Prevotella oralis		14.71	0.29		1	1	
07351332	**Low discrimination**						**07351332**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Lactobacillus acidophilus		41.70	0.38	RHA 7	99	1	
Clostridium barati		32.78	0.34	MLZ 1	99	99	
Prevotella oralis		14.42	0.29		1	1	
07351342	**Low discrimination**						**07351342**
					SPO		
Lactobacillus acidophilus		82.55	0.56	RHA 7	1		
Clostridium barati		15.22	0.41	SOR 11	99		
Lactobacillus catenaformis		2.04	0.32		1		
07351360	**Low discrimination**						**07351360**
					SPO		
Clostridium barati		85.87	0.41	ESL 87	99		
Lactobacillus acidophilus		4.70	0.22	RHA 8	1		
Clostridium paraputrificum		3.53	0.13	SOR 8	99		
07351362	**Low discrimination**						**07351362**
					GRA	SPO	
Lactobacillus acidophilus		62.46	0.56	RHA 7	99	1	
Clostridium barati		34.55	0.50	SOR 11	99	99	
Prevotella oralis		1.77	0.29		1	1	
07351400	**Low discrimination**						**07351400**
					SPO		
Lactobacillus catenaformis		65.32	0.63	ARA 1	1		
Lactobacillus jensenii		32.79	0.58		1		
Clostridium barati		1.10	0.27		99		
07351402	**Low discrimination**						**07351402**
Lactobacillus catenaformis		68.83	0.63	ARA 1			
Lactobacillus jensenii		23.04	0.55				
Lactobacillus acidophilus		3.98	0.39				
07351410	**Acceptable identification**						**07351410**
Lactobacillus jensenii		94.13	0.50	ARA 1			
Lactobacillus catenaformis		5.68	0.29				
Actinomyces israelii		0.19	0.16				
07351412	**Low discrimination**						**07351412**
Lactobacillus jensenii		88.20	0.47	ARA 1			
Lactobacillus catenaformis		7.99	0.29				
Bifidobacterium adolescentis		1.71	0.23				
07351420	**Low discrimination**						**07351420**
					SPO		
Lactobacillus catenaformis		58.31	0.53	ARA 1	1		
Lactobacillus jensenii		29.28	0.47		1		
Clostridium barati		11.73	0.35		99		
07351422	**Low discrimination**						**07351422**
					SPO		
Lactobacillus catenaformis		45.77	0.53	ARA 1	1		
Clostridium barati		27.61	0.44		99		
Lactobacillus jensenii		15.32	0.44		1		
07351430	**Acceptable identification**						**07351430**
					SPO		
Lactobacillus jensenii		93.24	0.39	ARA 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		5.63	0.18		1	
Clostridium barati		1.13	0.01		99	
07351432	**Low discrimination**					**07351432**
					SPO	
Lactobacillus jensenii		83.06	0.36	ARA 1	1	
Lactobacillus catenaformis		7.52	0.18		1	
Clostridium barati		4.54	0.09		99	
07351500	**Low discrimination**					**07351500**
					SPO	
Lactobacillus catenaformis		88.12	0.66	ARA 1	1	
Collinsella aerofaciens		6.48	0.32		1	
Clostridium barati		2.95	0.35		99	
07351502	**Low discrimination**					**07351502**
Lactobacillus catenaformis		59.30	0.66	ARA 1		
Lactobacillus acidophilus		26.28	0.57			
Bifidobacterium adolescentis		6.94	0.56			
07351512	**Low discrimination**					**07351512**
Bifidobacterium adolescentis		87.41	0.57	RAF 92		
Lactobacillus catenaformis		6.17	0.32	XYL 87		
Lactobacillus acidophilus		2.73	0.23			
07351520	**Low discrimination**					**07351520**
					SPO	
Lactobacillus catenaformis		59.54	0.56	ARA 1	1	
Clostridium barati		23.95	0.44		99	
Clostridium paraputrificum		9.80	0.32		99	
07351522	**Low discrimination**					**07351522**
					SPO	
Lactobacillus acidophilus		42.20	0.57	ARA 1	1	
Clostridium barati		28.72	0.52		99	
Lactobacillus catenaformis		23.80	0.56		1	
07351542	**Low discrimination**					**07351542**
Lactobacillus acidophilus		59.80	0.41	ARA 1		
Bifidobacterium adolescentis		15.80	0.40	SOR 11		
Lactobacillus catenaformis		11.03	0.32			
07351552	**Acceptable identification**					**07351552**
Bifidobacterium adolescentis		94.25	0.42	RAF 92		
Lactobacillus acidophilus		2.95	0.07	XYL 87		
Actinomyces israelii		2.80	0.22	SOR 11		
07351562	**Low discrimination**					**07351562**
					SPO	
Lactobacillus acidophilus		63.61	0.41	ARA 1	1	
Clostridium barati		30.45	0.34	SOR 11	99	
Lactobacillus catenaformis		2.93	0.21		1	
07351702	**Low discrimination**					**07351702**
					SPO	
Lactobacillus acidophilus		57.06	0.38	RHA 7	1	
Lactobacillus catenaformis		17.28	0.32	ARA 1	1	
Clostridium barati		14.95	0.25		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07351712	**Acceptable identification**						**07351712**
Clostridium clostridioforme		90.65	0.39	RAF 94			
Bifidobacterium adolescentis		7.58	0.23	TRE 88			
Lactobacillus acidophilus		1.77	0.04	XYL 91			
07351722	**Low discrimination**						**07351722**
					GRA	SPO	
Lactobacillus acidophilus		40.64	0.38	RHA 7	99	1	
Clostridium barati		31.95	0.34	ARA 1	99	99	
Prevotella oralis		14.06	0.29		1	1	
07351732	**Acceptable identification**						**07351732**
					GRA		
Clostridium clostridioforme		93.55	0.39	RAF 94	50		
Parabacteroides distasonis		3.19	0.19	TRE 88	1		
Lactobacillus acidophilus		1.82	0.04	XYL 91	99		
07352000	**Low discrimination**						**07352000**
					SPO		
Clostridium innocuum		79.72	0.44	MLT 1	99		
Lactobacillus catenaformis		7.19	0.29	GAL 95	1		
Clostridium barati		5.40	0.21		99		
07352001	**Low discrimination**						**07352001**
					SPO		
Actinomyces naeslundii		82.65	0.46	RAF 92	1		
Clostridium innocuum		17.35	0.10	GAL 96	99		
07352002	**Low discrimination**						**07352002**
					SPO		
Clostridium innocuum		72.20	0.44	MLT 1	99		
Clostridium barati		14.68	0.30	GAL 95	99		
Lactobacillus catenaformis		6.51	0.29		1		
07352003	**Low discrimination**						**07352003**
					SPO		
Actinomyces naeslundii		82.65	0.46	RAF 92	1		
Clostridium innocuum		17.35	0.10	GAL 96	99		
07352021	**Good identification**						**07352021**
Actinomyces naeslundii		100.0	0.46	RAF 92			
07352022	**Low discrimination**						**07352022**
					SPO		
Clostridium barati		85.66	0.38	ESL 87	99		
Actinomyces naeslundii		4.51	0.43	GAL 95	1		
Lactobacillus acidophilus		3.81	0.17	MAN 8	1		
07352023	**Low discrimination**						**07352023**
					SPO		
Actinomyces naeslundii		88.65	0.46	RAF 92	1		
Clostridium barati		11.35	0.04	GAL 96	99		
07352101	**Good identification**						**07352101**
Actinomyces naeslundii		100.0	0.53	RAF 92			
07352102	**Low discrimination**						**07352102**
					SPO		
Clostridium barati		48.29	0.38	ESL 87	99		
Lactobacillus acidophilus		24.73	0.35	GAL 95	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		10.71	0.32	MAN 8	1	
07352103	**Acceptable identification**					**07352103**
					SPO	
Actinomyces naeslundii		93.63	0.53	RAF 92	1	
Clostridium barati		4.21	0.04	GAL 96	99	
Lactobacillus acidophilus		2.16	0.01		1	
07352120	**Low discrimination**					**07352120**
					SPO	
Clostridium barati		69.26	0.38	ESL 87	99	
Actinomyces naeslundii		10.37	0.50	GAL 95	1	
Clostridium tertium		10.08	0.27	MAN 8	99	
07352121	**Good identification**					**07352121**
					SPO	
Actinomyces naeslundii		95.70	0.53	RAF 92	1	
Clostridium barati		4.30	0.04	GAL 96	99	
07352122	**Low discrimination**					**07352122**
					SPO	
Clostridium barati		66.91	0.46	ESL 87	99	
Lactobacillus acidophilus		11.42	0.35	GAL 95	1	
Clostridium tertium		9.74	0.35	MAN 8	99	
07352123	**Low discrimination**					**07352123**
					SPO	
Actinomyces naeslundii		84.92	0.53	RAF 92	1	
Clostridium barati		11.46	0.12	GAL 96	99	
Lactobacillus acidophilus		1.96	0.01		1	
07352132	**Acceptable identification**					**07352132**
					SPO	
Clostridium tertium		94.18	0.39	TRE 90	99	
Clostridium barati		3.68	0.12	GAL 99	99	
Actinomyces naeslundii		0.96	0.28		1	
07352141	**Good identification**					**07352141**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07352143	**Good identification**					**07352143**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07352161	**Good identification**					**07352161**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07352163	**Good identification**					**07352163**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07353000	**Low discrimination**					**07353000**
					SPO	
Clostridium innocuum		54.76	0.66	MLT 1	99	
Lactobacillus catenaformis		25.72	0.63		1	
Lactobacillus jensenii		12.92	0.58		1	
07353001	**Low discrimination**					**07353001**
					SPO	
Actinomyces naeslundii		77.24	0.69	RAF 92	1	
Clostridium innocuum		12.83	0.32		99	
Lactobacillus catenaformis		6.03	0.29		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07353002	**Low discrimination**					**07353002**
					SPO	
Clostridium innocuum		49.31	0.66	MLT 1	99	
Lactobacillus catenaformis		23.16	0.63		1	
Clostridium barati		10.02	0.51		99	
07353003	**Low discrimination**					**07353003**
					SPO	
Actinomyces naeslundii		75.22	0.69	RAF 92	1	
Clostridium innocuum		12.50	0.32		99	
Lactobacillus catenaformis		5.87	0.29		1	
07353010	**Low discrimination**					**07353010**
					SPO	
Lactobacillus jensenii		59.69	0.50	MAN 1	1	
Clostridium innocuum		31.64	0.42		99	
Lactobacillus catenaformis		3.60	0.29		1	
07353011	**Low discrimination**					**07353011**
					SPO	
Actinomyces naeslundii		72.48	0.47	RAF 92	1	
Lactobacillus jensenii		17.99	0.15	MLZ 5	1	
Clostridium innocuum		9.53	0.08		99	
07353012	**Low discrimination**					**07353012**
					SPO	
Lactobacillus jensenii		45.77	0.47	MAN 1	1	
Clostridium innocuum		36.39	0.42		99	
Clostridium tertium		7.25	0.31		99	
07353013	**Low discrimination**					**07353013**
					SPO	
Actinomyces naeslundii		77.10	0.47	RAF 92	1	
Lactobacillus jensenii		12.76	0.12	MLZ 5	1	
Clostridium innocuum		10.14	0.08		99	
07353020	**Low discrimination**					**07353020**
					SPO	
Clostridium barati		46.17	0.51	ESL 87	99	
Lactobacillus catenaformis		26.67	0.53	MAN 8	1	
Lactobacillus jensenii		13.39	0.47		1	
07353021	**Good identification**					**07353021**
					SPO	
Actinomyces naeslundii		94.07	0.69	RAF 92	1	
Clostridium barati		3.18	0.17		99	
Lactobacillus catenaformis		1.84	0.18		1	
07353022	**Low discrimination**					**07353022**
					SPO	
Clostridium barati		61.90	0.60	ESL 87	99	
Lactobacillus acidophilus		14.36	0.51	MAN 8	1	
Lactobacillus catenaformis		11.92	0.53		1	
07353023	**Low discrimination**					**07353023**
					SPO	
Actinomyces naeslundii		86.72	0.69	RAF 92	1	
Clostridium barati		8.79	0.25		99	
Lactobacillus acidophilus		2.04	0.17		1	
07353030	**Low discrimination**					**07353030**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		58.23	0.39	MAN 1	1	
Clostridium tertium		24.60	0.31		99	
Actinomyces naeslundii		6.32	0.44		1	
07353031	**Acceptable identification**					**07353031**
Actinomyces naeslundii		94.16	0.47	RAF 92		
Lactobacillus jensenii		5.84	0.05	MLZ 5		
07353032	**Low discrimination**					**07353032**
					SPO	
Clostridium tertium		50.34	0.39	TRE 90	99	
Lactobacillus jensenii		26.47	0.36	LAC 99	1	
Clostridium barati		12.45	0.25		99	
07353033	**Low discrimination**					**07353033**
					SPO	
Actinomyces naeslundii		89.29	0.47	RAF 92	1	
Clostridium tertium		7.02	0.05	MLZ 5	99	
Lactobacillus jensenii		3.69	0.02		1	
07353041	**Very good identification**					**07353041**
Actinomyces naeslundii		100.0	0.56	RAF 92		
07353043	**Very good identification**					**07353043**
					SPO	
Actinomyces naeslundii		96.80	0.56	RAF 92	1	
Clostridium barati		1.61	-.01		99	
Lactobacillus acidophilus		1.59	0.01		1	
07353061	**Very good identification**					**07353061**
					SPO	
Actinomyces naeslundii		98.36	0.56	RAF 92	1	
Clostridium barati		1.64	-.01		99	
07353062	**Low discrimination**					**07353062**
					SPO	
Clostridium barati		66.44	0.41	ESL 87	99	
Lactobacillus acidophilus		21.91	0.35	MAN 8	1	
Actinomyces naeslundii		8.96	0.53	SOR 8	1	
07353063	**Good identification**					**07353063**
					SPO	
Actinomyces naeslundii		93.77	0.56	RAF 92	1	
Clostridium barati		4.68	0.07		99	
Lactobacillus acidophilus		1.54	0.01		1	
07353100	**Low discrimination**					**07353100**
					SPO	
Lactobacillus catenaformis		54.38	0.66	MAN 1	1	
Clostridium barati		15.69	0.51		99	
Clostridium tertium		11.89	0.53		99	
07353101	**Good identification**					**07353101**
					SPO	
Actinomyces naeslundii		93.66	0.77	RAF 92	1	
Lactobacillus catenaformis		3.85	0.32		1	
Clostridium barati		1.11	0.17		99	
07353102	**Low discrimination**					**07353102**
					SPO	
Lactobacillus acidophilus		45.06	0.69	MAN 5	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		19.51	0.66		1	
Clostridium barati		16.89	0.60		99	
07353103	**Low discrimination**					**07353103**
Actinomyces naeslundii		83.24	0.77	RAF 92		
Lactobacillus acidophilus		7.91	0.35			
Lactobacillus catenaformis		3.42	0.32			
07353110	**Good identification**					**07353110**
					SPO	
Clostridium tertium		92.42	0.57	TRE 90	99	
Lactobacillus catenaformis		2.40	0.32		1	
Actinomyces naeslundii		2.05	0.52		1	
07353111	**Low discrimination**					**07353111**
					SPO	
Actinomyces naeslundii		76.69	0.55	RAF 92	1	
Clostridium tertium		23.31	0.23	MLZ 5	99	
07353112	**Good identification**					**07353112**
					SPO	
Clostridium tertium		92.75	0.65	TRE 90	99	
Lactobacillus acidophilus		1.86	0.35		1	
Bifidobacterium adolescentis		1.39	0.42		1	
07353113	**Low discrimination**					**07353113**
					SPO	
Actinomyces naeslundii		51.45	0.55	RAF 92	1	
Clostridium tertium		46.91	0.31	MLZ 5	99	
Lactobacillus acidophilus		0.94	0.01		1	
07353120	**Low discrimination**					**07353120**
Clostridium barati		36.35	0.60	ESL 87		
Clostridium tertium		27.56	0.61	MAN 8		
Clostridium ramosum		14.92	0.54			
07353121	**Good identification**					**07353121**
					SPO	
Actinomyces naeslundii		91.73	0.77	RAF 92	1	
Clostridium barati		3.27	0.25		99	
Clostridium tertium		2.48	0.27		99	
07353122	**Low discrimination**					**07353122**
					SPO	
Clostridium barati		29.56	0.68	ESL 87	99	
Lactobacillus acidophilus		26.29	0.69	MAN 8	1	
Clostridium tertium		22.41	0.69		99	
07353123	**Low discrimination**					**07353123**
					SPO	
Actinomyces naeslundii		73.84	0.77	RAF 92	1	
Clostridium barati		7.89	0.34		99	
Lactobacillus acidophilus		7.01	0.35		1	
07353130	**Very good identification**					**07353130**
					SPO	
Clostridium tertium		97.75	0.65	TRE 90	99	
Clostridium barati		0.73	0.25		99	
Actinomyces naeslundii		0.72	0.52		1	
07353131	**Low discrimination**					**07353131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		52.31	0.55	RAF 92	1	
Clostridium tertium		47.69	0.31	MLZ 5	99	
07353132	**Very good identification**					**07353132**
					SPO	
Clostridium tertium		97.30	0.73	TRE 90	99	
Clostridium barati		0.73	0.34		99	
Lactobacillus acidophilus		0.65	0.35		1	
07353133	**Low discrimination**					**07353133**
					SPO	
Clostridium tertium		72.17	0.39	TRE 90	99	
Actinomyces naeslundii		26.39	0.55	URE 1	1	
Clostridium barati		0.54	-.01		99	
07353140	**Low discrimination**					**07353140**
					SPO	
Actinomyces naeslundii		33.04	0.61	RAF 92	1	
Clostridium barati		28.69	0.33		99	
Actinomyces israelii		18.56	0.48		1	
07353141	**Very good identification**					**07353141**
					SPO	
Actinomyces naeslundii		99.05	0.64	RAF 92	1	
Clostridium barati		0.58	-.01		99	
Actinomyces israelii		0.38	0.13		1	
07353142	**Low discrimination**					**07353142**
					SPO	
Lactobacillus acidophilus		64.25	0.54	MAN 5	1	
Clostridium barati		16.94	0.41	SOR 11	99	
Actinomyces naeslundii		6.50	0.61		1	
07353143	**Good identification**					**07353143**
					SPO	
Actinomyces naeslundii		91.77	0.64	RAF 92	1	
Lactobacillus acidophilus		6.11	0.19		1	
Clostridium barati		1.61	0.07		99	
07353151	**Good identification**					**07353151**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07353153	**Good identification**					**07353153**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07353160	**Low discrimination**					**07353160**
					SPO	
Clostridium barati		59.20	0.41	ESL 87	99	
Actinomyces naeslundii		22.72	0.61	MAN 8	1	
Clostridium tertium		5.21	0.27	SOR 8	99	
07353161	**Very good identification**					**07353161**
					SPO	
Actinomyces naeslundii		98.28	0.64	RAF 92	1	
Clostridium barati		1.72	0.07		99	
07353162	**Low discrimination**					**07353162**
					SPO	
Lactobacillus acidophilus		47.47	0.54	MAN 5	1	
Clostridium barati		37.56	0.50	SOR 11	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		4.81	0.61		1	
07353163	**Low discrimination**					**07353163**
					SPO	
Actinomyces naeslundii		88.61	0.64	RAF 92	1	
Lactobacillus acidophilus		5.90	0.19		1	
Clostridium barati		4.66	0.15		99	
07353171	**Good identification**					**07353171**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07353172	**Low discrimination**					**07353172**
					SPO	
Clostridium tertium		56.20	0.39	TRE 90	99	
Bifidobacterium breve		32.80	0.44	SOR 1	1	
Lactobacillus acidophilus		4.58	0.19		1	
07353173	**Low discrimination**					**07353173**
					SPO	
Actinomyces naeslundii		80.13	0.42	RAF 92	1	
Clostridium tertium		12.55	0.05	MLZ 5	99	
Bifidobacterium breve		7.32	0.10		1	
07353201	**Good identification**					**07353201**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07353203	**Low discrimination**					**07353203**
					SPO	
Actinomyces naeslundii		77.48	0.35	RAF 92	1	
Clostridium barati		22.53	-.01	RHA 1	99	
07353221	**Low discrimination**					**07353221**
					SPO	
Actinomyces naeslundii		77.48	0.35	RAF 92	1	
Clostridium barati		22.53	-.01	RHA 1	99	
07353222	**Low discrimination**					**07353222**
					SPO	
Clostridium barati		66.50	0.41	ESL 87	99	
Clostridium ramosum		16.82	0.33	MAN 8	92	
Lactobacillus acidophilus		13.36	0.32	RHA 8	1	
07353223	**Low discrimination**					**07353223**
					SPO	
Actinomyces naeslundii		47.78	0.35	RAF 92	1	
Clostridium barati		41.68	0.07	RHA 1	99	
Clostridium ramosum		10.54	-.02		92	
07353301	**Acceptable identification**					**07353301**
					SPO	
Actinomyces naeslundii		90.73	0.43	RAF 92	1	
Clostridium barati		9.27	-.01	RHA 1	99	
07353302	**Low discrimination**					**07353302**
					SPO	
Lactobacillus acidophilus		59.57	0.50	MAN 5	1	
Clostridium barati		25.79	0.41	RHA 7	99	
Clostridium ramosum		6.52	0.33		92	
07353303	**Low discrimination**					**07353303**
					SPO	
Actinomyces naeslundii		47.81	0.43	RAF 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		33.84	0.16	RHA 1	1	
Clostridium barati		14.65	0.07		99	
07353320	**Low discrimination**					**07353320**
Clostridium ramosum		84.10	0.57	TRE 94		
Clostridium barati		13.44	0.41			
Clostridium tertium		1.18	0.27			
07353322	**Low discrimination**					**07353322**
					SPO	
Clostridium ramosum		81.21	0.67	TRE 94	92	
Clostridium barati		9.73	0.50		99	
Lactobacillus acidophilus		7.49	0.50		1	
07353332	**Low discrimination**					**07353332**
Clostridium tertium		60.30	0.39	TRE 90		
Clostridium ramosum		32.46	0.33	RHA 1		
Clostridium barati		3.89	0.15			
07353400	**Acceptable identification**					**07353400**
					SPO	
Clostridium innocuum		92.54	0.53	MLT 1	99	
Actinomyces israelii		3.15	0.43		1	
Lactobacillus catenaformis		2.49	0.29		1	
07353402	**Low discrimination**					**07353402**
					SPO	
Clostridium innocuum		89.85	0.53	MLT 1	99	
Actinomyces israelii		4.59	0.46		1	
Lactobacillus catenaformis		2.42	0.29		1	
07353421	**Good identification**					**07353421**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07353423	**Good identification**					**07353423**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07353440	**Low discrimination**					**07353440**
					SPO	
Actinomyces israelii		56.75	0.36	XYL 92	1	
Clostridium innocuum		43.25	0.19		99	
07353442	**Low discrimination**					**07353442**
					SPO	
Actinomyces israelii		62.14	0.39	XYL 92	1	
Clostridium innocuum		31.57	0.19		99	
Clostridium barati		3.16	-.01		99	
07353500	**Low discrimination**					**07353500**
					SPO	
Actinomyces israelii		60.97	0.53	XYL 92	1	
Lactobacillus catenaformis		18.04	0.32		1	
Clostridium barati		5.20	0.17		99	
07353501	**Low discrimination**					**07353501**
Actinomyces naeslundii		87.80	0.43	RAF 92		
Actinomyces israelii		12.20	0.19	ARA 1		
07353502	**Low discrimination**					**07353502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		47.86	0.54	RAF 92		
Actinomyces israelii		25.58	0.56	XYL 87		
Lactobacillus acidophilus		11.65	0.35			
07353503	**Low discrimination**					**07353503**
Actinomyces naeslundii		59.05	0.43	RAF 92		
Bifidobacterium adolescentis		23.03	0.20	ARA 1		
Actinomyces israelii		12.31	0.22			
07353510	**Low discrimination**					**07353510**
					SPO	
Actinomyces israelii		49.95	0.39	XYL 92	1	
Clostridium tertium		38.47	0.23	MLZ 13	99	
Bifidobacterium adolescentis		11.58	0.22		1	
07353512	**Low discrimination**					**07353512**
					SPO	
Bifidobacterium adolescentis		85.62	0.56	RAF 92	1	
Clostridium tertium		8.62	0.31	XYL 87	99	
Actinomyces israelii		5.60	0.42		1	
07353521	**Good identification**					**07353521**
Actinomyces naeslundii		100.0	0.43	RAF 92		
07353523	**Low discrimination**					**07353523**
					SPO	
Actinomyces naeslundii		77.96	0.43	RAF 92	1	
Clostridium barati		8.33	-.01	ARA 1	99	
Lactobacillus acidophilus		7.40	0.01		1	
07353532	**Acceptable identification**					**07353532**
					SPO	
Clostridium tertium		93.69	0.39	TRE 90	99	
Bifidobacterium adolescentis		3.14	0.22	ARA 1	1	
Actinomyces israelii		1.30	0.22		1	
07353540	**Good identification**					**07353540**
					SPO	
Actinomyces israelii		96.06	0.46	XYL 92	1	
Actinomyces naeslundii		2.11	0.27		1	
Clostridium barati		1.83	-.01		99	
07353542	**Low discrimination**					**07353542**
Actinomyces israelii		55.80	0.49	XYL 92		
Bifidobacterium adolescentis		33.18	0.39			
Lactobacillus acidophilus		8.08	0.19			
07353552	**Low discrimination**					**07353552**
Bifidobacterium adolescentis		82.95	0.40	RAF 92		
Actinomyces israelii		17.06	0.35	XYL 87		
07353702	**Low discrimination**					**07353702**
Actinomyces israelii		51.18	0.37	XYL 92		
Lactobacillus acidophilus		23.32	0.16	RHA 7		
Bifidobacterium adolescentis		12.85	0.20			
07354000	**Low discrimination**					**07354000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		39.60	0.40	MLT 11	1		
Lactobacillus catenaformis		33.64	0.29	XYL 7	99		
Lactobacillus jensenii		16.89	0.24		99		
07354100	**Low discrimination**						**07354100**
					COC	GRA	
Fusobacterium mortiferum		50.09	0.46	MLT 11	1	1	
Lactobacillus catenaformis		27.35	0.32	XYL 7	1	99	
Atopobium parvulum		13.01	0.25		99	99	
07354502	**Low discrimination**						**07354502**
Bifidobacterium adolescentis		70.13	0.36	RAF 92			
Clostridium clostridioforme		15.46	0.22	GAL 99			
Actinomyces israelii		9.30	0.27				
07354512	**Low discrimination**						**07354512**
Bifidobacterium adolescentis		54.90	0.37	RAF 92			
Clostridium clostridioforme		45.10	0.33	GAL 99			
07354612	**Good identification**						**07354612**
Clostridium clostridioforme		100.0	0.39	RAF 94			
07354632	**Good identification**						**07354632**
Clostridium clostridioforme		100.0	0.39	RAF 94			
07354702	**Good identification**						**07354702**
Clostridium clostridioforme		100.0	0.36	RAF 94			
07354712	**Good identification**						**07354712**
Clostridium clostridioforme		100.0	0.48	RAF 94			
07354722	**Good identification**						**07354722**
Clostridium clostridioforme		100.0	0.36	RAF 94			
07354732	**Good identification**						**07354732**
Clostridium clostridioforme		100.0	0.48	RAF 94			
07355000	**Low discrimination**						**07355000**
Lactobacillus catenaformis		63.25	0.63	XYL 1			
Lactobacillus jensenii		31.75	0.58				
Actinomyces israelii		2.01	0.50				
07355002	**Low discrimination**						**07355002**
Lactobacillus catenaformis		66.70	0.63	XYL 1			
Lactobacillus jensenii		22.32	0.55				
Lactobacillus acidophilus		3.86	0.39				
07355010	**Acceptable identification**						**07355010**
Lactobacillus jensenii		91.85	0.50	XYL 1			
Lactobacillus catenaformis		5.54	0.29				
Actinomyces israelii		2.60	0.35				
07355012	**Low discrimination**						**07355012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		83.90	0.47	XYL 1		
Lactobacillus catenaformis		7.60	0.29			
Actinomyces israelii		5.35	0.38			
07355020	**Low discrimination**					**07355020**
					SPO	
Lactobacillus catenaformis		57.65	0.53	XYL 1	1	
Lactobacillus jensenii		28.94	0.47		1	
Clostridium barati		11.59	0.35		99	
07355022	**Low discrimination**					**07355022**
					SPO	
Lactobacillus catenaformis		43.21	0.53	XYL 1	1	
Clostridium barati		26.07	0.44		99	
Lactobacillus jensenii		14.46	0.44		1	
07355030	**Acceptable identification**					**07355030**
					SPO	
Lactobacillus jensenii		92.11	0.39	XYL 1	1	
Lactobacillus catenaformis		5.56	0.18		1	
Clostridium barati		1.12	0.01		99	
07355032	**Low discrimination**					**07355032**
					SPO	
Lactobacillus jensenii		80.34	0.36	XYL 1	1	
Lactobacillus catenaformis		7.27	0.18		1	
Clostridium barati		4.39	0.09		99	
07355040	**Low discrimination**					**07355040**
Actinomyces israelii		42.61	0.43			
Lactobacillus catenaformis		34.86	0.29			
Lactobacillus jensenii		17.50	0.24			
07355042	**Low discrimination**					**07355042**
Actinomyces israelii		42.56	0.46			
Lactobacillus catenaformis		23.21	0.29			
Lactobacillus acidophilus		16.42	0.23			
07355100	**Low discrimination**					**07355100**
					COC	
Lactobacillus catenaformis		57.93	0.66	XYL 1	1	
Atopobium parvulum		27.56	0.59		99	
Actinomyces israelii		4.91	0.60		1	
07355102	**Low discrimination**					**07355102**
Lactobacillus catenaformis		52.07	0.66	XYL 1		
Lactobacillus acidophilus		23.08	0.57			
Actinomyces israelii		6.61	0.63			
07355110	**Low discrimination**					**07355110**
					SPO	
Actinomyces israelii		33.49	0.46	MLZ 13	1	
Lactobacillus catenaformis		26.74	0.32		1	
Clostridium tertium		14.36	0.25		99	
07355112	**Low discrimination**					**07355112**
					SPO	
Bifidobacterium adolescentis		72.22	0.57	RAF 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		9.58	0.49	ARA 87	1		
Clostridium tertium		8.21	0.33		99		
07355120	**Low discrimination**						**07355120**
					COC	SPO	
Lactobacillus catenaformis		46.70	0.56	XYL 1	1	1	
Clostridium barati		18.78	0.44		1	99	
Atopobium parvulum		15.69	0.46		99	1	
07355122	**Low discrimination**						**07355122**
					GRA	SPO	
Lactobacillus acidophilus		31.45	0.57	XYL 1	99	1	
Prevotella oralis		23.38	0.53		1	1	
Clostridium barati		21.40	0.52		99	99	
07355132	**Low discrimination**						**07355132**
					GRA	SPO	
Clostridium tertium		69.91	0.41	TRE 90	99	99	
Lactobacillus acidophilus		6.41	0.23	MAN 99	99	1	
Prevotella oralis		4.77	0.19		1	1	
07355140	**Low discrimination**						**07355140**
					COC		
Actinomyces israelii		61.27	0.53		1		
Lactobacillus catenaformis		18.79	0.32		1		
Atopobium parvulum		8.94	0.25		99		
07355142	**Low discrimination**						**07355142**
Lactobacillus acidophilus		39.39	0.41	XYL 1			
Actinomyces israelii		35.52	0.56	SOR 11			
Bifidobacterium adolescentis		10.41	0.40				
07355150	**Good identification**						**07355150**
Actinomyces israelii		96.50	0.39	MLZ 13			
Bifidobacterium adolescentis		3.51	0.08				
07355152	**Low discrimination**						**07355152**
Bifidobacterium adolescentis		68.59	0.42	RAF 92			
Actinomyces israelii		28.62	0.42	ARA 87			
Lactobacillus acidophilus		2.15	0.07	SOR 11			
07355162	**Low discrimination**						**07355162**
					GRA	SPO	
Lactobacillus acidophilus		59.29	0.41	XYL 1	99	1	
Clostridium barati		28.39	0.34	SOR 11	99	99	
Prevotella oralis		3.60	0.19		1	1	
07355302	**Low discrimination**						**07355302**
Lactobacillus acidophilus		44.05	0.38	XYL 1			
Lactobacillus catenaformis		13.34	0.32	RHA 7			
Actinomyces israelii		12.62	0.44				
07355312	**Low discrimination**						**07355312**
Clostridium clostridioforme		76.78	0.35	RAF 94			
Bifidobacterium adolescentis		9.95	0.23	TRE 88			
Actinomyces israelii		9.83	0.29	ARA 94			
07355322	**Low discrimination**						**07355322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Prevotella oralis		62.26	0.47	RAF 92	1	1	
Lactobacillus acidophilus		14.71	0.38	XYL 11	99	1	
Clostridium barati		11.57	0.34		99	99	
07355332	**Low discrimination**						**07355332**
Clostridium clostridioforme		49.96	0.35	RAF 94			
Parabacteroides distasonis		38.76	0.39	TRE 88			
Prevotella oralis		6.39	0.13	ARA 94			
07355400	**Low discrimination**						**07355400**
					SPO		
Actinomyces israelii		51.37	0.48		1		
Lactobacillus catenaformis		19.97	0.29		1		
Clostridium butyricum		18.16	0.19		99		
07355402	**Low discrimination**						**07355402**
Actinomyces israelii		54.45	0.51				
Bifidobacterium adolescentis		16.59	0.36				
Lactobacillus catenaformis		14.11	0.29				
07355412	**Low discrimination**						**07355412**
Bifidobacterium adolescentis		45.15	0.37	RAF 92			
Clostridium clostridioforme		33.23	0.32	LAC 99			
Actinomyces israelii		18.12	0.37				
07355440	**Acceptable identification**						**07355440**
					SPO		
Actinomyces israelii		92.68	0.41		1		
Clostridium butyricum		7.33	0.01		99		
07355442	**Low discrimination**						**07355442**
					SPO		
Actinomyces israelii		89.50	0.44		1		
Bifidobacterium adolescentis		8.67	0.20		1		
Clostridium butyricum		1.83	-.06		99		
07355500	**Low discrimination**						**07355500**
					SPO		
Clostridium butyricum		61.06	0.43	ESL 88	99		
Actinomyces israelii		28.79	0.58	RAF 95	1		
Lactobacillus catenaformis		4.20	0.32	TRE 94	1		
07355502	**Low discrimination**						**07355502**
					SPO		
Bifidobacterium adolescentis		81.05	0.70	RAF 92	1		
Actinomyces israelii		10.75	0.61		1		
Clostridium butyricum		5.91	0.36		99		
07355503	**Low discrimination**						**07355503**
					SPO		
Bifidobacterium adolescentis		82.95	0.36	RAF 92	1		
Actinomyces israelii		11.00	0.27	URE 1	1		
Clostridium butyricum		6.05	0.02		99		
07355510	**Low discrimination**						**07355510**
					SPO		
Actinomyces israelii		36.97	0.44	MLZ 13	1		
Bifidobacterium adolescentis		34.55	0.37		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		27.62	0.21		99	
07355512	**Very good identification**					**07355512**
Bifidobacterium adolescentis		95.75	0.72	RAF 92		
Clostridium clostridioforme		2.38	0.41			
Actinomyces israelii		1.55	0.47			
07355513	**Good identification**					**07355513**
Bifidobacterium adolescentis		97.57	0.37	RAF 92		
Clostridium clostridioforme		2.43	0.07	URE 1		
07355522	**Low discrimination**					**07355522**
Bifidobacterium adolescentis		18.12	0.36	RAF 92		
Actinomyces israelii		15.19	0.41	NAG 1		
Clostridium clostridioforme		11.98	0.30			
07355532	**Low discrimination**					**07355532**
					GRA	
Clostridium clostridioforme		64.08	0.41	RAF 94	50	
Bifidobacterium adolescentis		26.01	0.37	TRE 88	99	
Parabacteroides distasonis		4.75	0.27	RHA 88	1	
07355540	**Low discrimination**					**07355540**
					SPO	
Actinomyces israelii		66.20	0.51		1	
Clostridium butyricum		31.39	0.25		99	
Bifidobacterium adolescentis		2.40	0.20		1	
07355542	**Low discrimination**					**07355542**
					SPO	
Bifidobacterium adolescentis		67.83	0.55	RAF 92	1	
Actinomyces israelii		28.30	0.54	SOR 11	1	
Clostridium butyricum		3.48	0.18		99	
07355550	**Low discrimination**					**07355550**
					SPO	
Actinomyces israelii		68.30	0.37	MLZ 13	1	
Bifidobacterium adolescentis		20.29	0.22		1	
Clostridium butyricum		11.41	0.03		99	
07355552	**Acceptable identification**					**07355552**
					SPO	
Bifidobacterium adolescentis		94.76	0.56	RAF 92	1	
Actinomyces israelii		4.83	0.40	SOR 11	1	
Clostridium butyricum		0.21	-.04		99	
07355602	**Low discrimination**					**07355602**
Clostridium clostridioforme		85.43	0.36	RAF 94		
Actinomyces israelii		14.57	0.32	TRE 88		
07355612	**Good identification**					**07355612**
Clostridium clostridioforme		99.44	0.47	RAF 94		
Actinomyces israelii		0.56	0.18	TRE 88		
07355622	**Good identification**					**07355622**
Clostridium clostridioforme		100.0	0.36	RAF 94		
07355632	**Good identification**					**07355632**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.47	RAF 94			
07355702	**Low discrimination**						**07355702**
Clostridium clostridioforme		66.61	0.45	RAF 94			
Bifidobacterium adolescentis		13.74	0.36	TRE 88			
Actinomyces israelii		13.58	0.42				
07355712	**Acceptable identification**						**07355712**
Clostridium clostridioforme		94.16	0.56	RAF 94			
Bifidobacterium adolescentis		5.21	0.37	TRE 88			
Actinomyces israelii		0.63	0.28				
07355720	**Good identification**						**07355720**
					GRA	SPO	
Bacteroides vulgatus		96.28	0.36	RAF 97	1	1	
Eubacterium contortum		2.07	0.14	CEL 5	99	1	
Clostridium butyricum		0.59	-.01		99	99	
07355722	**Low discrimination**						**07355722**
Clostridium clostridioforme		69.79	0.45	RAF 94			
Parabacteroides distasonis		17.92	0.40	TRE 88			
Bacteroides uniformis		4.56	0.11				
07355732	**Good identification**						**07355732**
Clostridium clostridioforme		96.76	0.56	RAF 94			
Parabacteroides distasonis		2.94	0.35	TRE 88			
Bacteroides ovatus		0.30	0.03				
07355742	**Low discrimination**						**07355742**
Actinomyces israelii		69.01	0.35	RHA 7			
Bifidobacterium adolescentis		22.20	0.20				
Clostridium clostridioforme		8.79	0.10				
07356100	**Low discrimination**						**07356100**
					GRA	SPO	
Actinomyces israelii		59.95	0.39	GAL 99	99	1	
Clostridium tertium		37.74	0.21		99	99	
Fusobacterium mortiferum		2.31	0.12		1	1	
07356122	**Acceptable identification**						**07356122**
					SPO		
Clostridium tertium		92.96	0.37	TRE 90	99		
Clostridium barati		4.67	0.12	GAL 99	99		
Actinomyces israelii		1.57	0.21		1		
07356132	**Good identification**						**07356132**
Clostridium tertium		100.0	0.41	TRE 90			
07356500	**Good identification**						**07356500**
					SPO		
Actinomyces israelii		96.01	0.37	GAL 99	1		
Clostridium butyricum		3.99	-.08		99		
07356512	**Low discrimination**						**07356512**
					SPO		
Bifidobacterium adolescentis		88.26	0.36	RAF 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		9.91	0.26	GAL 99	1	
Clostridium tertium		1.83	-.01		99	
07357000	**Low discrimination**					**07357000**
					SPO	
Actinomyces israelii		57.30	0.62		1	
Clostridium innocuum		35.68	0.44		99	
Lactobacillus catenaformis		3.22	0.29		1	
07357002	**Low discrimination**					**07357002**
					SPO	
Actinomyces israelii		64.43	0.65		1	
Clostridium innocuum		26.75	0.44		99	
Clostridium tertium		3.28	0.29		99	
07357010	**Low discrimination**					**07357010**
					SPO	
Actinomyces israelii		64.97	0.48	MLZ 13	1	
Clostridium tertium		19.66	0.25		99	
Clostridium innocuum		11.28	0.20		99	
07357012	**Low discrimination**					**07357012**
					SPO	
Actinomyces israelii		55.78	0.51	MLZ 13	1	
Clostridium tertium		33.76	0.33		99	
Clostridium innocuum		6.46	0.20		99	
07357021	**Good identification**					**07357021**
Actinomyces naeslundii		100.0	0.35	RAF 92		
07357022	**Low discrimination**					**07357022**
					SPO	
Clostridium tertium		51.32	0.37	TRE 90	99	
Actinomyces israelii		21.46	0.45	LAC 99	1	
Clostridium barati		16.34	0.25		99	
07357023	**Low discrimination**					**07357023**
					SPO	
Actinomyces naeslundii		75.86	0.35	RAF 92	1	
Clostridium tertium		24.14	0.03	XYL 1	99	
07357032	**Good identification**					**07357032**
					SPO	
Clostridium tertium		96.25	0.41	TRE 90	99	
Actinomyces israelii		3.38	0.31	LAC 99	1	
Lactobacillus jensenii		0.37	0.02		1	
07357040	**Good identification**					**07357040**
					SPO	
Actinomyces israelii		98.41	0.55		1	
Clostridium innocuum		1.59	0.10		99	
07357042	**Very good identification**					**07357042**
					SPO	
Actinomyces israelii		98.23	0.58		1	
Clostridium innocuum		1.06	0.10		99	
Clostridium barati		0.36	-.01		99	
07357050	**Good identification**					**07357050**
Actinomyces israelii		100.0	0.41	MLZ 13		
07357052	**Good identification**					**07357052**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		98.45	0.44	MLZ 13	1	
Clostridium tertium		1.55	-.01		99	
07357062	**Low discrimination**					**07357062**
					SPO	
Actinomyces israelii		77.62	0.38	NAG 6	1	
Clostridium barati		13.21	0.07		99	
Clostridium tertium		4.82	0.03		99	
07357100	**Low discrimination**					**07357100**
					SPO	
Actinomyces israelii		59.52	0.73		1	
Clostridium tertium		37.47	0.55		99	
Lactobacillus catenaformis		1.25	0.32		1	
07357101	**Low discrimination**					**07357101**
					SPO	
Actinomyces israelii		46.70	0.39	URE 1	1	
Clostridium tertium		29.40	0.21		99	
Actinomyces naeslundii		23.90	0.43		1	
07357102	**Low discrimination**					**07357102**
					SPO	
Clostridium tertium		51.25	0.63	TRE 90	99	
Actinomyces israelii		40.71	0.76		1	
Bifidobacterium adolescentis		5.42	0.54		1	
07357110	**Low discrimination**					**07357110**
					SPO	
Clostridium tertium		88.05	0.59	TRE 90	99	
Actinomyces israelii		11.76	0.59		1	
Bifidobacterium adolescentis		0.19	0.22		1	
07357112	**Low discrimination**					**07357112**
					SPO	
Clostridium tertium		87.75	0.67	TRE 90	99	
Bifidobacterium adolescentis		6.38	0.56		1	
Actinomyces israelii		5.86	0.62		1	
07357120	**Good identification**					**07357120**
					SPO	
Clostridium tertium		94.89	0.63	TRE 90	99	
Actinomyces israelii		3.21	0.52		1	
Clostridium barati		0.92	0.25		99	
07357122	**Very good identification**					**07357122**
					SPO	
Clostridium tertium		95.82	0.71	TRE 90	99	
Actinomyces israelii		1.62	0.55		1	
Clostridium barati		0.92	0.34		99	
07357123	**Low discrimination**					**07357123**
					SPO	
Clostridium tertium		88.85	0.37	TRE 90	99	
Actinomyces naeslundii		8.03	0.43	URE 1	1	
Actinomyces israelii		1.50	0.21		1	
07357130	**Very good identification**					**07357130**
					SPO	
Clostridium tertium		99.72	0.67	TRE 90	99	
Actinomyces israelii		0.28	0.38		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07357132	**Very good identification**					**07357132**
					SPO	
Clostridium tertium		99.82	0.76	TRE 90	99	
Actinomyces israelii		0.14	0.41		1	
Bifidobacterium adolescentis		0.02	0.22		1	
07357133	**Good identification**					**07357133**
Clostridium tertium		100.0	0.41	TRE 90		
07357140	**Very good identification**					**07357140**
					SPO	
Actinomyces israelii		98.11	0.66		1	
Clostridium tertium		1.60	0.21		99	
Actinomyces naeslundii		0.15	0.27		1	
07357142	**Good identification**					**07357142**
					SPO	
Actinomyces israelii		91.82	0.69		1	
Bifidobacterium adolescentis		3.88	0.39		1	
Clostridium tertium		3.00	0.29		99	
07357150	**Low discrimination**					**07357150**
					SPO	
Actinomyces israelii		83.71	0.52	MLZ 13	1	
Clostridium tertium		16.29	0.25		99	
07357152	**Low discrimination**					**07357152**
					SPO	
Actinomyces israelii		57.63	0.55	MLZ 13	1	
Clostridium tertium		22.42	0.33		99	
Bifidobacterium adolescentis		19.94	0.40		1	
07357160	**Low discrimination**					**07357160**
					SPO	
Actinomyces israelii		53.85	0.45	NAG 6	1	
Clostridium tertium		41.39	0.29		99	
Clostridium barati		3.44	0.07		99	
07357162	**Low discrimination**					**07357162**
					SPO	
Clostridium tertium		53.85	0.37	TRE 90	99	
Actinomyces israelii		35.04	0.48	SOR 1	1	
Lactobacillus acidophilus		5.65	0.19		1	
07357172	**Acceptable identification**					**07357172**
					SPO	
Clostridium tertium		94.43	0.41	TRE 90	99	
Actinomyces israelii		5.16	0.34	SOR 1	1	
Bifidobacterium breve		0.40	0.10		1	
07357200	**Acceptable identification**					**07357200**
					SPO	
Actinomyces israelii		92.29	0.43	RHA 7	1	
Clostridium innocuum		7.71	0.10		99	
07357202	**Acceptable identification**					**07357202**
					SPO	
Actinomyces israelii		93.07	0.46	RHA 7	1	
Clostridium innocuum		5.19	0.10		99	
Clostridium barati		1.74	-.01		99	
07357240	**Good identification**					**07357240**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	RHA 7		
07357242	**Good identification**					**07357242**
Actinomyces israelii		100.0	0.39	RHA 7		
07357300	**Acceptable identification**					**07357300**
					SPO	
Actinomyces israelii		91.62	0.54	RHA 7	1	
Clostridium tertium		7.74	0.21		99	
Clostridium barati		0.64	-.01		99	
07357302	**Low discrimination**					**07357302**
					SPO	
Actinomyces israelii		80.85	0.57	RHA 7	1	
Clostridium tertium		13.66	0.29		99	
Lactobacillus acidophilus		2.62	0.16		1	
07357322	**Low discrimination**					**07357322**
					SPO	
Clostridium tertium		49.41	0.37	TRE 90	99	
Clostridium ramosum		34.24	0.33	RHA 1	92	
Actinomyces israelii		6.22	0.36		1	
07357332	**Good identification**					**07357332**
					SPO	
Clostridium tertium		98.57	0.41	TRE 90	99	
Actinomyces israelii		1.04	0.22	RHA 1	1	
Clostridium ramosum		0.39	-.02		92	
07357340	**Good identification**					**07357340**
Actinomyces israelii		100.0	0.46	RHA 7		
07357342	**Good identification**					**07357342**
Actinomyces israelii		98.98	0.49	RHA 7		
Lactobacillus acidophilus		1.02	0.00			
07357352	**Good identification**					**07357352**
					SPO	
Actinomyces israelii		95.04	0.35	RHA 7	1	
Clostridium tertium		4.96	-.01	MLZ 13	99	
07357400	**Low discrimination**					**07357400**
					SPO	
Actinomyces israelii		87.63	0.61		1	
Clostridium innocuum		11.77	0.31		99	
Clostridium butyricum		0.61	0.03		99	
07357402	**Low discrimination**					**07357402**
					SPO	
Actinomyces israelii		88.08	0.64		1	
Clostridium innocuum		7.89	0.31		99	
Bifidobacterium adolescentis		3.88	0.34		1	
07357410	**Good identification**					**07357410**
					SPO	
Actinomyces israelii		96.39	0.47	MLZ 13	1	
Clostridium innocuum		3.61	0.07		99	
07357412	**Low discrimination**					**07357412**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		71.83	0.50	MLZ 13	1	
Bifidobacterium adolescentis		25.84	0.36		1	
Clostridium innocuum		1.79	0.07		99	
07357420	**Good identification**					**07357420**
Actinomyces israelii		100.0	0.40	NAG 6		
07357422	**Good identification**					**07357422**
					SPO	
Actinomyces israelii		97.13	0.43	NAG 6	1	
Clostridium tertium		2.87	0.03		99	
07357440	**Good identification**					**07357440**
Actinomyces israelii		100.0	0.54			
07357442	**Very good identification**					**07357442**
Actinomyces israelii		98.62	0.57			
Bifidobacterium adolescentis		1.38	0.19			
07357450	**Good identification**					**07357450**
Actinomyces israelii		100.0	0.40	MLZ 13		
07357452	**Low discrimination**					**07357452**
Actinomyces israelii		89.74	0.43	MLZ 13		
Bifidobacterium adolescentis		10.26	0.20			
07357462	**Good identification**					**07357462**
Actinomyces israelii		100.0	0.36	NAG 6		
07357500	**Good identification**					**07357500**
					SPO	
Actinomyces israelii		93.82	0.71		1	
Clostridium butyricum		3.90	0.27		99	
Bifidobacterium adolescentis		1.55	0.34		1	
07357501	**Good identification**					**07357501**
					SPO	
Actinomyces israelii		96.01	0.37	URE 1	1	
Clostridium butyricum		3.99	-.08		99	
07357502	**Low discrimination**					**07357502**
					SPO	
Bifidobacterium adolescentis		51.47	0.69	RAF 92	1	
Actinomyces israelii		47.27	0.74		1	
Clostridium tertium		0.74	0.29		99	
07357510	**Low discrimination**					**07357510**
					SPO	
Actinomyces israelii		80.51	0.57	MLZ 13	1	
Bifidobacterium adolescentis		10.86	0.36		1	
Clostridium tertium		7.45	0.25		99	
07357512	**Low discrimination**					**07357512**
					SPO	
Bifidobacterium adolescentis		88.05	0.70	RAF 92	1	
Actinomyces israelii		9.89	0.60		1	
Clostridium tertium		1.83	0.33		99	
07357513	**Low discrimination**					**07357513**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Bifidobacterium adolescentis		88.26	0.36	RAF 92	1	
Actinomyces israelii		9.91	0.26	URE 1	1	
Clostridium tertium		1.83	-.01		99	
07357520	**Low discrimination**					**07357520**
					SPO	
Actinomyces israelii		70.33	0.51	NAG 6	1	
Clostridium tertium		25.69	0.29		99	
Clostridium butyricum		3.98	0.08		99	
07357522	**Low discrimination**					**07357522**
					SPO	
Actinomyces israelii		51.79	0.54	NAG 6	1	
Clostridium tertium		37.84	0.37		99	
Bifidobacterium adolescentis		8.92	0.34		1	
07357532	**Low discrimination**					**07357532**
					SPO	
Clostridium tertium		74.81	0.41	TRE 90	99	
Bifidobacterium adolescentis		12.13	0.36	ARA 1	1	
Actinomyces israelii		8.61	0.40		1	
07357540	**Very good identification**					**07357540**
					SPO	
Actinomyces israelii		98.57	0.64		1	
Clostridium butyricum		0.92	0.08		99	
Bifidobacterium adolescentis		0.52	0.19		1	
07357542	**Low discrimination**					**07357542**
					SPO	
Actinomyces israelii		74.16	0.67		1	
Bifidobacterium adolescentis		25.66	0.53		1	
Clostridium butyricum		0.18	0.01		99	
07357550	**Acceptable identification**					**07357550**
					COC	
Actinomyces israelii		94.76	0.50	MLZ 13	1	
Bifidobacterium adolescentis		4.06	0.20		1	
Blautia producta		1.18	-.03		99	
07357552	**Low discrimination**					**07357552**
					SPO	
Bifidobacterium adolescentis		73.80	0.55	RAF 92	1	
Actinomyces israelii		26.08	0.53	SOR 11	1	
Clostridium tertium		0.13	-.01		99	
07357560	**Good identification**					**07357560**
Actinomyces israelii		100.0	0.44	NAG 6		
07357562	**Acceptable identification**					**07357562**
					SPO	
Actinomyces israelii		93.13	0.47	NAG 6	1	
Bifidobacterium adolescentis		5.10	0.19		1	
Clostridium tertium		1.77	0.03		99	
07357600	**Good identification**					**07357600**
Actinomyces israelii		100.0	0.42	RHA 7		
07357602	**Good identification**					**07357602**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.45	RHA 7		
07357642	**Good identification**					**07357642**
Actinomyces israelii		100.0	0.38	RHA 7		
07357700	**Good identification**					**07357700**
					SPO	
Actinomyces israelii		98.32	0.52	RHA 7	1	
Clostridium butyricum		1.68	0.01		99	
07357702	**Low discrimination**					**07357702**
Actinomyces israelii		86.26	0.55	RHA 7		
Bifidobacterium adolescentis		12.60	0.34			
Clostridium clostridioforme		0.75	0.10			
07357710	**Good identification**					**07357710**
Actinomyces israelii		100.0	0.38	RHA 7		
07357712	**Low discrimination**					**07357712**
Bifidobacterium adolescentis		48.06	0.36	RAF 92		
Actinomyces israelii		40.22	0.41	RHA 1		
Clostridium clostridioforme		10.72	0.22			
07357740	**Good identification**					**07357740**
Actinomyces israelii		100.0	0.45	RHA 7		
07357742	**Good identification**					**07357742**
Actinomyces israelii		95.56	0.48	RHA 7		
Bifidobacterium adolescentis		4.44	0.19			
07361000	**Very good identification**					**07361000**
Lactobacillus jensenii		98.49	0.66	FRU 99		
Lactobacillus catenaformis		0.66	0.29			
Actinomyces naeslundii		0.30	0.44			
07361002	**Low discrimination**					**07361002**
Lactobacillus jensenii		82.46	0.63	FRU 99		
Lactobacillus acidophilus		15.89	0.48			
Lactobacillus catenaformis		0.83	0.29			
07361010	**Very good identification**					**07361010**
Lactobacillus jensenii		99.95	0.58	FRU 99		
Actinomyces naeslundii		0.05	0.22			
07361012	**Good identification**					**07361012**
					GRA	
Lactobacillus jensenii		99.10	0.55	FRU 99	99	
Lactobacillus acidophilus		0.58	0.14		99	
Leptotrichia buccalis		0.25	-.03		1	
07361020	**Good identification**					**07361020**
					SPO	
Lactobacillus jensenii		95.81	0.56	FRU 99	1	
Actinomyces naeslundii		1.16	0.44		1	
Clostridium barati		1.11	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07361021	**Low discrimination**						**07361021**
Actinomyces naeslundii		64.17	0.47	RAF 92			
Lactobacillus jensenii		35.83	0.21	FRU 95			
07361022	**Low discrimination**						**07361022**
					SPO		
Lactobacillus jensenii		53.81	0.53	FRU 99	1		
Lactobacillus acidophilus		41.49	0.48		1		
Clostridium barati		2.81	0.25		99		
07361023	**Low discrimination**						**07361023**
Actinomyces naeslundii		60.27	0.47	RAF 92			
Lactobacillus jensenii		22.43	0.18	FRU 95			
Lactobacillus acidophilus		17.30	0.14				
07361030	**Good identification**						**07361030**
Lactobacillus jensenii		99.81	0.47	FRU 99			
Actinomyces naeslundii		0.19	0.22				
07361032	**Good identification**						**07361032**
Lactobacillus jensenii		97.45	0.44	FRU 99			
Lactobacillus acidophilus		2.28	0.14				
Actinomyces naeslundii		0.28	0.22				
07361100	**Low discrimination**						**07361100**
					COC		
Atopobium parvulum		83.82	0.54	FRU 99	99		
Lactobacillus acidophilus		4.90	0.32		1		
Lactobacillus jensenii		3.32	0.32		1		
07361101	**Low discrimination**						**07361101**
					COC	SPO	
Actinomyces naeslundii		81.93	0.55	RAF 92	1	1	
Atopobium parvulum		16.41	0.20	FRU 95	99	1	
Clostridium chauvoei		1.67	0.03		1	99	
07361102	**Good identification**						**07361102**
					COC		
Lactobacillus acidophilus		94.79	0.66	FRU 99	1		
Atopobium parvulum		2.89	0.41		99		
Lactobacillus catenaformis		0.65	0.32		1		
07361120	**Low discrimination**						**07361120**
					COC	SPO	
Atopobium parvulum		35.44	0.41	FRU 99	99	1	
Lactobacillus acidophilus		11.75	0.32		1	1	
Clostridium septicum		9.33	0.25		1	99	
07361121	**Good identification**						**07361121**
					COC		
Actinomyces naeslundii		96.59	0.55	RAF 92	1		
Atopobium parvulum		3.41	0.07	FRU 95	99		
07361122	**Good identification**						**07361122**
					SPO		
Lactobacillus acidophilus		93.65	0.66	FRU 99	1		
Clostridium ramosum		2.51	0.38		92		
Clostridium barati		1.66	0.34		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07361141	**Good identification**					**07361141**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07361142	**Good identification**					**07361142**
					SPO	
Lactobacillus acidophilus		98.44	0.50	FRU 99	1	
Actinomyces naeslundii		0.82	0.39	SOR 11	1	
Clostridium barati		0.41	0.07		99	
07361143	**Low discrimination**					**07361143**
Actinomyces naeslundii		55.18	0.42	RAF 92		
Lactobacillus acidophilus		44.82	0.16	FRU 95		
07361161	**Good identification**					**07361161**
Actinomyces naeslundii		100.0	0.42	RAF 92		
07361162	**Good identification**					**07361162**
					SPO	
Lactobacillus acidophilus		97.04	0.50	FRU 99	1	
Clostridium barati		1.21	0.15	SOR 11	99	
Actinomyces naeslundii		0.81	0.39		1	
07361163	**Low discrimination**					**07361163**
Actinomyces naeslundii		55.18	0.42	RAF 92		
Lactobacillus acidophilus		44.82	0.16	FRU 95		
07361302	**Good identification**					**07361302**
					SPO	
Lactobacillus acidophilus		98.46	0.47	FRU 99	1	
Clostridium barati		0.67	0.07	RHA 7	99	
Clostridium ramosum		0.47	0.06		92	
07361322	**Low discrimination**					**07361322**
					SPO	
Lactobacillus acidophilus		66.75	0.47	FRU 99	1	
Clostridium ramosum		31.45	0.40	RHA 7	92	
Clostridium barati		1.36	0.15		99	
07362101	**Good identification**					**07362101**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07362103	**Good identification**					**07362103**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07362121	**Good identification**					**07362121**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07362123	**Good identification**					**07362123**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07363000	**Low discrimination**					**07363000**
					SPO	
Clostridium innocuum		61.23	0.36	MLT 1	99	
Lactobacillus jensenii		23.66	0.32	FRU 95	1	
Actinomyces naeslundii		13.71	0.49		1	
07363001	**Good identification**					**07363001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		97.08	0.52	RAF 92	1	
Clostridium innocuum		2.92	0.02	FRU 95	99	
07363002	**Low discrimination**					**07363002**
					SPO	
Clostridium innocuum		55.69	0.36	MLT 1	99	
Lactobacillus acidophilus		14.41	0.26	FRU 95	1	
Lactobacillus jensenii		14.35	0.29		1	
07363003	**Good identification**					**07363003**
					SPO	
Actinomyces naeslundii		97.08	0.52	RAF 92	1	
Clostridium innocuum		2.92	0.02	FRU 95	99	
07363020	**Low discrimination**					**07363020**
					SPO	
Actinomyces naeslundii		58.18	0.49	RAF 92	1	
Lactobacillus jensenii		25.10	0.21	FRU 95	1	
Clostridium tertium		5.97	0.09		99	
07363021	**Good identification**					**07363021**
Actinomyces naeslundii		100.0	0.52	RAF 92		
07363023	**Good identification**					**07363023**
Actinomyces naeslundii		100.0	0.52	RAF 92		
07363040	**Low discrimination**					**07363040**
					SPO	
Actinomyces naeslundii		79.65	0.36	RAF 92	1	
Clostridium innocuum		20.35	0.02	FRU 95	99	
07363041	**Good identification**					**07363041**
Actinomyces naeslundii		100.0	0.39	RAF 92		
07363042	**Low discrimination**					**07363042**
					SPO	
Actinomyces naeslundii		43.71	0.36	RAF 92	1	
Lactobacillus acidophilus		35.37	0.10	FRU 95	1	
Clostridium innocuum		11.17	0.02		99	
07363043	**Good identification**					**07363043**
Actinomyces naeslundii		100.0	0.39	RAF 92		
07363060	**Good identification**					**07363060**
Actinomyces naeslundii		100.0	0.36	RAF 92		
07363061	**Good identification**					**07363061**
Actinomyces naeslundii		100.0	0.39	RAF 92		
07363062	**Low discrimination**					**07363062**
Actinomyces naeslundii		55.27	0.36	RAF 92		
Lactobacillus acidophilus		44.73	0.10	FRU 95		
07363063	**Good identification**					**07363063**
Actinomyces naeslundii		100.0	0.39	RAF 92		
07363101	**Very good identification**					**07363101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		99.21	0.60	RAF 92	1	
Clostridium tertium		0.79	0.01	FRU 95	99	
07363102	**Low discrimination**					**07363102**
					SPO	
Lactobacillus acidophilus		48.62	0.44	FRU 99	1	
Clostridium tertium		37.14	0.43	MAN 5	99	
Actinomyces naeslundii		10.41	0.57		1	
07363103	**Good identification**					**07363103**
					SPO	
Actinomyces naeslundii		94.75	0.60	RAF 92	1	
Lactobacillus acidophilus		2.98	0.10	FRU 95	1	
Clostridium tertium		2.28	0.09		99	
07363110	**Good identification**					**07363110**
					SPO	
Clostridium tertium		96.94	0.39	FRU 99	99	
Actinomyces naeslundii		2.41	0.35		1	
Actinomyces israelii		0.65	0.16		1	
07363111	**Low discrimination**					**07363111**
					SPO	
Actinomyces naeslundii		78.71	0.38	RAF 92	1	
Clostridium tertium		21.29	0.05	FRU 95	99	
07363112	**Good identification**					**07363112**
					SPO	
Clostridium tertium		97.62	0.47	FRU 99	99	
Actinomyces naeslundii		0.81	0.35		1	
Lactobacillus acidophilus		0.73	0.10		1	
07363113	**Low discrimination**					**07363113**
					SPO	
Actinomyces naeslundii		55.21	0.38	RAF 92	1	
Clostridium tertium		44.79	0.13	FRU 95	99	
07363120	**Low discrimination**					**07363120**
					SPO	
Clostridium tertium		44.32	0.43	FRU 99	99	
Clostridium ramosum		41.77	0.41		92	
Actinomyces naeslundii		12.43	0.57		1	
07363121	**Very good identification**					**07363121**
					SPO	
Actinomyces naeslundii		95.54	0.60	RAF 92	1	
Clostridium tertium		2.30	0.09	FRU 95	99	
Clostridium ramosum		2.16	0.07		92	
07363122	**Low discrimination**					**07363122**
					SPO	
Clostridium ramosum		44.90	0.51	FRU 99	92	
Clostridium tertium		35.72	0.51		99	
Lactobacillus acidophilus		15.59	0.44		1	
07363123	**Low discrimination**					**07363123**
					SPO	
Actinomyces naeslundii		83.75	0.60	RAF 92	1	
Clostridium ramosum		7.58	0.17	FRU 95	92	
Clostridium tertium		6.03	0.17		99	
07363130	**Good identification**					**07363130**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium tertium		98.65	0.47	FRU 99	99		
Actinomyces naeslundii		0.82	0.35		1		
Clostridium ramosum		0.53	0.07		92		
07363131	**Low discrimination**						**07363131**
					SPO		
Actinomyces naeslundii		55.21	0.38	RAF 92	1		
Clostridium tertium		44.79	0.13	FRU 95	99		
07363132	**Good identification**						**07363132**
					SPO		
Clostridium tertium		98.78	0.55	FRU 99	99		
Clostridium ramosum		0.71	0.17		92		
Actinomyces naeslundii		0.27	0.35		1		
07363140	**Low discrimination**						**07363140**
					SPO		
Actinomyces naeslundii		78.31	0.44	RAF 92	1		
Actinomyces israelii		16.36	0.23	FRU 95	1		
Clostridium tertium		5.33	0.01		99		
07363141	**Good identification**						**07363141**
Actinomyces naeslundii		100.0	0.47	RAF 92			
07363143	**Good identification**						**07363143**
Actinomyces naeslundii		100.0	0.47	RAF 92			
07363160	**Low discrimination**						**07363160**
					SPO		
Actinomyces naeslundii		71.60	0.44	RAF 92	1		
Clostridium tertium		14.62	0.09	FRU 95	99		
Clostridium ramosum		13.78	0.07		92		
07363161	**Good identification**						**07363161**
Actinomyces naeslundii		100.0	0.47	RAF 92			
07363163	**Good identification**						**07363163**
Actinomyces naeslundii		100.0	0.47	RAF 92			
07363320	**Good identification**						**07363320**
					SPO		
Clostridium ramosum		98.98	0.43	FRU 99	92		
Clostridium tertium		0.80	0.09		99		
Actinomyces naeslundii		0.22	0.23		1		
07363322	**Good identification**						**07363322**
					SPO		
Clostridium ramosum		97.37	0.53	FRU 99	92		
Lactobacillus acidophilus		1.92	0.25		1		
Clostridium tertium		0.59	0.17		99		
07365100	**Very good identification**						**07365100**
					COC	SPO	
Atopobium parvulum		98.74	0.59	FRU 99	99	1	
Actinomyces israelii		0.67	0.36		1	1	
Clostridium butyricum		0.54	0.13		1	99	
07365102	**Low discrimination**						**07365102**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		78.91	0.46	FRU 99	99		
Lactobacillus acidophilus		14.09	0.32		1		
Actinomyces israelii		4.54	0.39		1		
07365120	**Good identification**						**07365120**
					COC	SPO	
Atopobium parvulum		98.52	0.46	FRU 99	99	1	
Clostridium tertium		0.98	0.11		1	99	
Clostridium butyricum		0.26	-.05		1	99	
07365712	**Good identification**						**07365712**
Clostridium clostridioforme		100.0	0.36	RAF 94			
07365732	**Good identification**						**07365732**
Clostridium clostridioforme		98.15	0.36	RAF 94			
Parabacteroides distasonis		1.85	0.13	FRU 99			
07367000	**Low discrimination**						**07367000**
					SPO		
Actinomyces israelii		73.32	0.38	FRU 99	1		
Clostridium innocuum		22.22	0.14		99		
Clostridium tertium		4.46	0.03		99		
07367002	**Low discrimination**						**07367002**
					SPO		
Actinomyces israelii		75.55	0.41	FRU 99	1		
Clostridium innocuum		15.26	0.14		99		
Clostridium tertium		9.19	0.11		99		
07367100	**Low discrimination**						**07367100**
					COC	SPO	
Clostridium tertium		53.89	0.37	FRU 99	1	99	
Actinomyces israelii		35.78	0.48		1	1	
Atopobium parvulum		9.43	0.25		99	1	
07367102	**Low discrimination**						**07367102**
					COC	SPO	
Clostridium tertium		73.44	0.45	FRU 99	1	99	
Actinomyces israelii		24.38	0.51		1	1	
Atopobium parvulum		0.76	0.12		99	1	
07367110	**Acceptable identification**						**07367110**
					SPO		
Clostridium tertium		94.71	0.41	FRU 99	99		
Actinomyces israelii		5.29	0.34		1		
07367112	**Good identification**						**07367112**
					SPO		
Clostridium tertium		96.78	0.50	FRU 99	99		
Actinomyces israelii		2.70	0.37		1		
Bifidobacterium adolescentis		0.52	0.19		1		
07367120	**Good identification**						**07367120**
					COC	SPO	
Clostridium tertium		96.77	0.45	FRU 99	1	99	
Actinomyces israelii		1.37	0.28		1	1	
Atopobium parvulum		1.00	0.12		99	1	
07367122	**Good identification**						**07367122**
					SPO		
Clostridium tertium		98.03	0.53	FRU 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		0.90	0.17		92	
Actinomyces israelii		0.69	0.31		1	
07367130	**Good identification**					**07367130**
					SPO	
Clostridium tertium		99.88	0.50	FRU 99	99	
Actinomyces israelii		0.12	0.14		1	
07367132	**Very good identification**					**07367132**
					SPO	
Clostridium tertium		99.94	0.58	FRU 99	99	
Actinomyces israelii		0.06	0.17		1	
07367140	**Good identification**					**07367140**
					SPO	
Actinomyces israelii		96.24	0.41	FRU 99	1	
Clostridium tertium		3.76	0.03		99	
07367142	**Acceptable identification**					**07367142**
					SPO	
Actinomyces israelii		92.74	0.44	FRU 99	1	
Clostridium tertium		7.26	0.11		99	
07367400	**Acceptable identification**					**07367400**
					SPO	
Actinomyces israelii		93.87	0.37	FRU 99	1	
Clostridium innocuum		6.13	0.01		99	
07367402	**Good identification**					**07367402**
					SPO	
Actinomyces israelii		95.83	0.40	FRU 99	1	
Clostridium innocuum		4.18	0.01		99	
07367500	**Low discrimination**					**07367500**
					SPO	
Actinomyces israelii		83.91	0.47	FRU 99	1	
Clostridium butyricum		14.53	0.13		99	
Clostridium tertium		1.56	0.03		99	
07367502	**Low discrimination**					**07367502**
					SPO	
Actinomyces israelii		78.43	0.50	FRU 99	1	
Bifidobacterium adolescentis		15.13	0.31		1	
Clostridium butyricum		3.52	0.06		99	
07367540	**Good identification**					**07367540**
					SPO	
Actinomyces israelii		96.27	0.40	FRU 99	1	
Clostridium butyricum		3.73	-.05		99	
07367542	**Acceptable identification**					**07367542**
Actinomyces israelii		94.22	0.43	FRU 99		
Bifidobacterium adolescentis		5.78	0.16			
07370000	**Very good identification**					**07370000**
					GRA	
Lactobacillus jensenii		95.12	0.66	GAL 99	99	
Fusobacterium mortiferum		2.98	0.50		1	
Lactobacillus catenaformis		0.64	0.29		99	
07370002	**Low discrimination**					**07370002**
					SPO	
Lactobacillus jensenii		80.89	0.63	GAL 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus acidophilus		15.59	0.48		1			
Clostridium barati		1.84	0.30		99			
07370010	**Very good identification**							**07370010**
					GRA	CAT		
Lactobacillus jensenii		99.85	0.58	GAL 99	99	1		
Fusobacterium mortiferum		0.10	0.16		1	5		
Propionibacterium granulosum		0.06	0.09		99	90		
07370012	**Good identification**							**07370012**
					GRA			
Lactobacillus jensenii		99.17	0.55	GAL 99	99			
Lactobacillus acidophilus		0.58	0.14		99			
Leptotrichia buccalis		0.25	-.03		1			
07370020	**Acceptable identification**							**07370020**
					SPO			
Lactobacillus jensenii		91.51	0.56	GAL 99	1			
Clostridium barati		5.54	0.30		99			
Actinomyces naeslundii		0.87	0.43		1			
07370021	**Low discrimination**							**07370021**
Actinomyces naeslundii		58.64	0.46	RAF 92				
Lactobacillus jensenii		41.36	0.21	GAL 96				
07370022	**Low discrimination**							**07370022**
					SPO			
Lactobacillus jensenii		48.21	0.53	GAL 99	1			
Lactobacillus acidophilus		37.17	0.48		1			
Clostridium barati		13.12	0.38		99			
07370023	**Low discrimination**							**07370023**
Actinomyces naeslundii		51.00	0.46	RAF 92				
Lactobacillus jensenii		23.98	0.18	GAL 96				
Lactobacillus acidophilus		18.49	0.14					
07370030	**Good identification**							**07370030**
Lactobacillus jensenii		100.0	0.47	GAL 99				
07370032	**Good identification**							**07370032**
					SPO			
Lactobacillus jensenii		96.94	0.44	GAL 99	1			
Lactobacillus acidophilus		2.27	0.14		1			
Clostridium barati		0.80	0.04		99			
07370100	**Low discrimination**							**07370100**
					COC	GRA	SPO	
Atopobium parvulum		65.96	0.54	GAL 99	99	99	1	
Fusobacterium mortiferum		18.90	0.56		1	1	1	
Clostridium barati		3.91	0.30		1	99	99	
07370101	**Low discrimination**							**07370101**
					COC	GRA		
Actinomyces naeslundii		75.45	0.53	RAF 92	1	99		
Atopobium parvulum		19.08	0.20	GAL 96	99	99		
Fusobacterium mortiferum		5.47	0.22		1	1		
07370102	**Good identification**							**07370102**
					COC	SPO		
Lactobacillus acidophilus		92.22	0.66	GAL 99	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		2.83	0.38		1	99	
Atopobium parvulum		2.81	0.41		99	1	
07370120	**Low discrimination**						**07370120**
					COC	SPO	
Clostridium barati		27.89	0.38	ESL 87	1	99	
Atopobium parvulum		27.68	0.41	TRE 8	99	1	
Lactobacillus acidophilus		9.18	0.32	GAL 95	1	1	
07370121	**Acceptable identification**						**07370121**
					COC	SPO	
Actinomyces naeslundii		91.78	0.53	RAF 92	1	1	
Clostridium barati		4.13	0.04	GAL 96	1	99	
Atopobium parvulum		4.10	0.07		99	1	
07370122	**Low discrimination**						**07370122**
					SPO		
Lactobacillus acidophilus		87.64	0.66	GAL 99	1		
Clostridium barati		8.07	0.46		99		
Clostridium ramosum		2.34	0.38		92		
07370141	**Good identification**						**07370141**
Actinomyces naeslundii		100.0	0.40	RAF 92			
07370142	**Good identification**						**07370142**
					SPO		
Lactobacillus acidophilus		96.94	0.50	GAL 99	1		
Clostridium barati		2.09	0.20	SOR 11	99		
Actinomyces naeslundii		0.64	0.37		1		
07370161	**Good identification**						**07370161**
Actinomyces naeslundii		100.0	0.40	RAF 92			
07370162	**Acceptable identification**						**07370162**
					SPO		
Lactobacillus acidophilus		92.50	0.50	GAL 99	1		
Clostridium barati		5.99	0.28	SOR 11	99		
Actinomyces naeslundii		0.61	0.37		1		
07370302	**Good identification**						**07370302**
					SPO		
Lactobacillus acidophilus		95.27	0.47	GAL 99	1		
Clostridium barati		3.38	0.20	RHA 7	99		
Clostridium clostridioforme		0.51	0.14		66		
07370322	**Low discrimination**						**07370322**
					SPO		
Lactobacillus acidophilus		62.92	0.47	GAL 99	1		
Clostridium ramosum		29.64	0.40	RHA 7	92		
Clostridium barati		6.70	0.28		99		
07370612	**Good identification**						**07370612**
Clostridium clostridioforme		100.0	0.36	RAF 94			
07370632	**Good identification**						**07370632**
Clostridium clostridioforme		100.0	0.36	RAF 94			
07370712	**Good identification**						**07370712**
Clostridium clostridioforme		100.0	0.45	RAF 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07370732	**Good identification**					**07370732**
Clostridium clostridioforme		100.0	0.45	RAF 94		
07371000	**Very good identification**					**07371000**
					COC	
Lactobacillus jensenii		98.74	1.00		1	
Lactobacillus catenaformis		0.66	0.63		1	
Atopobium parvulum		0.25	0.54		99	
07371001	**Good identification**					**07371001**
Lactobacillus jensenii		91.08	0.66	URE 1		
Actinomyces naeslundii		7.83	0.69			
Lactobacillus catenaformis		0.61	0.29			
07371002	**Low discrimination**					**07371002**
Lactobacillus jensenii		82.71	0.97			
Lactobacillus acidophilus		15.94	0.82			
Lactobacillus catenaformis		0.83	0.63			
07371003	**Low discrimination**					**07371003**
Lactobacillus jensenii		74.81	0.63	URE 1		
Lactobacillus acidophilus		14.42	0.48			
Actinomyces naeslundii		9.64	0.69			
07371010	**Excellent identification**					**07371010**
Lactobacillus jensenii		99.95	0.92			
Lactobacillus catenaformis		0.02	0.29			
Actinomyces naeslundii		0.01	0.44			
07371011	**Very good identification**					**07371011**
Lactobacillus jensenii		98.66	0.58	URE 1		
Actinomyces naeslundii		1.34	0.47			
07371012	**Very good identification**					**07371012**
					GRA	
Lactobacillus jensenii		99.10	0.89		99	
Lactobacillus acidophilus		0.58	0.48		99	
Leptotrichia buccalis		0.25	0.32		1	
07371013	**Good identification**					**07371013**
Lactobacillus jensenii		97.21	0.55	URE 1		
Actinomyces naeslundii		1.98	0.47			
Lactobacillus acidophilus		0.57	0.14			
07371020	**Very good identification**					**07371020**
					SPO	
Lactobacillus jensenii		96.59	0.90		1	
Clostridium barati		1.12	0.51		99	
Lactobacillus catenaformis		0.65	0.53		1	
07371021	**Low discrimination**					**07371021**
					SPO	
Lactobacillus jensenii		72.71	0.56	URE 1	1	
Actinomyces naeslundii		24.99	0.69		1	
Clostridium barati		0.84	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07371022	**Low discrimination**					**07371022**
					SPO	
Lactobacillus jensenii		54.20	0.87		1	
Lactobacillus acidophilus		41.78	0.82		1	
Clostridium barati		2.83	0.60		99	
07371023	**Low discrimination**					**07371023**
Lactobacillus jensenii		42.45	0.53	URE 1		
Lactobacillus acidophilus		32.73	0.48			
Actinomyces naeslundii		21.89	0.69			
07371030	**Very good identification**					**07371030**
					SPO	
Lactobacillus jensenii		99.86	0.82		1	
Actinomyces naeslundii		0.04	0.44		1	
Clostridium barati		0.04	0.17		99	
07371031	**Acceptable identification**					**07371031**
Lactobacillus jensenii		94.85	0.47	URE 1		
Actinomyces naeslundii		5.15	0.47			
07371032	**Very good identification**					**07371032**
					SPO	
Lactobacillus jensenii		97.30	0.79		1	
Lactobacillus acidophilus		2.27	0.48		1	
Clostridium barati		0.15	0.25		99	
07371033	**Acceptable identification**					**07371033**
Lactobacillus jensenii		90.52	0.44	URE 1		
Actinomyces naeslundii		7.37	0.47			
Lactobacillus acidophilus		2.12	0.14			
07371040	**Very good identification**					**07371040**
Lactobacillus jensenii		95.43	0.66	SOR 1		
Lactobacillus acidophilus		1.52	0.32			
Actinomyces naeslundii		0.97	0.53			
07371041	**Low discrimination**					**07371041**
Actinomyces naeslundii		59.64	0.56	RAF 92		
Lactobacillus jensenii		39.73	0.32			
Lactobacillus acidophilus		0.63	-.02			
07371042	**Low discrimination**					**07371042**
					SPO	
Lactobacillus acidophilus		68.79	0.66	LAC 92	1	
Lactobacillus jensenii		29.17	0.63	SOR 11	1	
Clostridium barati		1.09	0.33		99	
07371050	**Very good identification**					**07371050**
Lactobacillus jensenii		99.67	0.58	SOR 1		
Actinomyces naeslundii		0.16	0.31			
Actinomyces israelii		0.12	0.20			
07371052	**Acceptable identification**					**07371052**
Lactobacillus jensenii		92.57	0.55	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		6.62	0.32				
Actinomyces israelii		0.25	0.23				
07371060	**Low discrimination**						**07371060**
					SPO		
Lactobacillus jensenii		74.90	0.56	SOR 1	1		
Clostridium perfringens		8.89	0.39		84		
Clostridium barati		7.49	0.33		99		
07371061	**Low discrimination**						**07371061**
Actinomyces naeslundii		82.41	0.56	RAF 92			
Lactobacillus jensenii		13.72	0.21				
Clostridium perfringens		1.63	0.05				
07371062	**Low discrimination**						**07371062**
					SPO		
Lactobacillus acidophilus		85.42	0.66	LAC 92	1		
Lactobacillus jensenii		9.05	0.53	SOR 11	1		
Clostridium barati		4.07	0.41		99		
07371070	**Good identification**						**07371070**
Lactobacillus jensenii		98.53	0.47	SOR 1			
Actinomyces naeslundii		0.63	0.31				
Clostridium perfringens		0.35	0.05				
07371072	**Low discrimination**						**07371072**
					SPO		
Lactobacillus jensenii		76.39	0.44	SOR 1	1		
Lactobacillus acidophilus		21.84	0.32		1		
Clostridium barati		1.04	0.07		99		
07371100	**Low discrimination**						**07371100**
					COC		
Atopobium parvulum		85.65	0.88		99		
Lactobacillus acidophilus		5.01	0.66		1		
Lactobacillus jensenii		3.39	0.66		1		
07371101	**Low discrimination**						**07371101**
					COC		
Atopobium parvulum		47.16	0.54	URE 1	99		
Actinomyces naeslundii		45.21	0.77		1		
Lactobacillus acidophilus		2.76	0.32		1		
07371102	**Very good identification**						**07371102**
					COC		
Lactobacillus acidophilus		95.18	1.00		1		
Atopobium parvulum		2.90	0.76		99		
Lactobacillus catenaformis		0.65	0.66		1		
07371103	**Low discrimination**						**07371103**
					COC		
Lactobacillus acidophilus		82.33	0.66	URE 1	1		
Actinomyces naeslundii		13.63	0.77		1		
Atopobium parvulum		2.51	0.41		99		
07371110	**Low discrimination**						**07371110**
					COC	GRA	
Lactobacillus jensenii		46.10	0.58	LAC 1	1	99	
Atopobium parvulum		35.29	0.54		99	99	
Leptotrichia buccalis		7.67	0.32		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07371111	**Low discrimination**						**07371111**
					COC		
Actinomyces naeslundii		65.23	0.55	RAF 92	1		
Lactobacillus jensenii		17.06	0.24	MLZ 5	1		
Atopobium parvulum		13.06	0.20		99		
07371112	**Low discrimination**						**07371112**
					GRA		
Leptotrichia buccalis		74.21	0.66	ESL 99	1		
Lactobacillus acidophilus		19.97	0.66		99		
Lactobacillus jensenii		3.00	0.55		99		
07371120	**Low discrimination**						**07371120**
					COC	SPO	
Atopobium parvulum		37.42	0.76		99	1	
Lactobacillus acidophilus		12.40	0.66		1	1	
Clostridium septicum		9.85	0.60		1	99	
07371121	**Low discrimination**						**07371121**
					COC		
Actinomyces naeslundii		67.33	0.77	RAF 92	1		
Atopobium parvulum		12.40	0.41		99		
Lactobacillus acidophilus		4.11	0.32		1		
07371122	**Good identification**						**07371122**
					SPO		
Lactobacillus acidophilus		94.05	1.00		1		
Clostridium ramosum		2.52	0.72		92		
Clostridium barati		1.66	0.68		99		
07371123	**Low discrimination**						**07371123**
					SPO		
Lactobacillus acidophilus		81.47	0.66	URE 1	1		
Actinomyces naeslundii		13.48	0.77		1		
Clostridium ramosum		2.18	0.38		92		
07371130	**Low discrimination**						**07371130**
					COC	SPO	
Clostridium tertium		32.94	0.47	MAN 99	1	99	
Lactobacillus jensenii		25.99	0.47		1	1	
Atopobium parvulum		14.04	0.41		99	1	
07371131	**Low discrimination**						**07371131**
					SPO		
Actinomyces naeslundii		84.49	0.55	RAF 92	1		
Clostridium tertium		7.00	0.13	MLZ 5	99		
Lactobacillus jensenii		5.53	0.13		1		
07371132	**Low discrimination**						**07371132**
					GRA	SPO	
Lactobacillus acidophilus		57.42	0.66	MLZ 1	99	1	
Leptotrichia buccalis		23.70	0.50		1	1	
Clostridium tertium		12.31	0.55		99	99	
07371140	**Low discrimination**						**07371140**
					COC		
Atopobium parvulum		47.83	0.54	SOR 1	99		
Lactobacillus acidophilus		34.24	0.50		1		
Actinomyces naeslundii		5.39	0.61		1		
07371141	**Good identification**						**07371141**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		90.23	0.64	RAF 92	1		
Atopobium parvulum		5.39	0.20		99		
Lactobacillus acidophilus		3.86	0.16		1		
07371142	**Very good identification**						**07371142**
					COC	SPO	
Lactobacillus acidophilus		98.93	0.85	SOR 11	1	1	
Clostridium barati		0.41	0.41		1	99	
Atopobium parvulum		0.25	0.41		99	1	
07371143	**Low discrimination**						**07371143**
					SPO		
Lactobacillus acidophilus		80.40	0.50	SOR 11	1		
Actinomyces naeslundii		19.00	0.64	URE 1	1		
Clostridium barati		0.33	0.07		99		
07371151	**Good identification**						**07371151**
Actinomyces naeslundii		100.0	0.42	RAF 92			
07371152	**Low discrimination**						**07371152**
					GRA		
Lactobacillus acidophilus		71.34	0.50	MLZ 1	99		
Leptotrichia buccalis		21.66	0.32	SOR 11	1		
Bifidobacterium breve		3.01	0.37		99		
07371153	**Low discrimination**						**07371153**
					GRA		
Actinomyces naeslundii		48.57	0.42	RAF 92	99		
Lactobacillus acidophilus		39.45	0.16	MLZ 5	99		
Leptotrichia buccalis		11.98	-.03		1		
07371160	**Low discrimination**						**07371160**
					SPO		
Clostridium perfringens		32.99	0.54	CEL 4	84		
Lactobacillus acidophilus		27.69	0.50		1		
Clostridium paraputrificum		12.31	0.37		99		
07371161	**Low discrimination**						**07371161**
Actinomyces naeslundii		87.66	0.64	RAF 92			
Clostridium perfringens		4.47	0.20				
Lactobacillus acidophilus		3.75	0.16				
07371162	**Very good identification**						**07371162**
					SPO		
Lactobacillus acidophilus		97.51	0.85	SOR 11	1		
Clostridium barati		1.21	0.50		99		
Clostridium perfringens		0.55	0.48		84		
07371163	**Low discrimination**						**07371163**
					SPO		
Lactobacillus acidophilus		79.46	0.50	SOR 11	1		
Actinomyces naeslundii		18.78	0.64	URE 1	1		
Clostridium barati		0.99	0.15		99		
07371171	**Good identification**						**07371171**
Actinomyces naeslundii		100.0	0.42	RAF 92			
07371172	**Low discrimination**						**07371172**
					GRA		
Lactobacillus acidophilus		86.22	0.50	MLZ 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		6.47	0.41	SOR 11	99	
Leptotrichia buccalis		2.91	0.15		1	
07371173	**Low discrimination**					**07371173**
Actinomyces naeslundii		55.18	0.42	RAF 92		
Lactobacillus acidophilus		44.82	0.16	MLZ 5		
07371200	**Very good identification**					**07371200**
					SPO	
Lactobacillus jensenii		97.20	0.66	RHA 1	1	
Lactobacillus acidophilus		0.94	0.29		1	
Clostridium barati		0.81	0.25		99	
07371202	**Low discrimination**					**07371202**
					SPO	
Lactobacillus acidophilus		57.72	0.63	LAC 92	1	
Lactobacillus jensenii		40.19	0.63	RHA 7	1	
Clostridium barati		1.51	0.33		99	
07371210	**Very good identification**					**07371210**
Lactobacillus jensenii		100.0	0.58	RHA 1		
07371212	**Acceptable identification**					**07371212**
Lactobacillus jensenii		94.52	0.55	RHA 1		
Lactobacillus acidophilus		4.11	0.29			
Clostridium clostridioforme		1.02	0.24			
07371220	**Low discrimination**					**07371220**
					SPO	
Lactobacillus jensenii		82.70	0.56	RHA 1	1	
Clostridium barati		8.27	0.33		99	
Clostridium ramosum		4.33	0.30		92	
07371222	**Low discrimination**					**07371222**
					SPO	
Lactobacillus acidophilus		76.01	0.63	LAC 92	1	
Lactobacillus jensenii		13.23	0.53	RHA 7	1	
Clostridium barati		5.95	0.41		99	
07371230	**Good identification**					**07371230**
					SPO	
Lactobacillus jensenii		99.70	0.47	RHA 1	1	
Clostridium barati		0.30	-.01		99	
07371232	**Low discrimination**					**07371232**
Lactobacillus jensenii		80.61	0.44	RHA 1		
Lactobacillus acidophilus		14.03	0.29			
Clostridium clostridioforme		3.49	0.24			
07371242	**Acceptable identification**					**07371242**
					SPO	
Lactobacillus acidophilus		92.78	0.47	LAC 92	1	
Lactobacillus jensenii		5.28	0.29	RHA 7	1	
Clostridium barati		1.70	0.15	SOR 11	99	
07371262	**Acceptable identification**					**07371262**
					SPO	
Lactobacillus acidophilus		93.13	0.47	LAC 92	1	
Clostridium barati		5.13	0.23	RHA 7	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		1.33	0.18	SOR 11	1		
07371300	**Low discrimination**						**07371300**
					COC	SPO	
Atopobium parvulum		59.17	0.54	RHA 1	99	1	
Lactobacillus acidophilus		25.79	0.47		1	1	
Clostridium barati		5.79	0.33		1	99	
07371302	**Very good identification**						**07371302**
					SPO		
Lactobacillus acidophilus		98.21	0.81	RHA 7	1		
Clostridium barati		0.67	0.41		99		
Clostridium ramosum		0.47	0.40		92		
07371303	**Good identification**						**07371303**
					SPO		
Lactobacillus acidophilus		96.35	0.47	RHA 7	1		
Actinomyces naeslundii		2.14	0.43	URE 1	1		
Clostridium barati		0.66	0.07		99		
07371312	**Low discrimination**						**07371312**
					GRA		
Lactobacillus acidophilus		61.70	0.47	RHA 7	99		
Leptotrichia buccalis		30.77	0.32	MLZ 1	1		
Clostridium clostridioforme		4.47	0.33		50		
07371320	**Low discrimination**						**07371320**
					SPO		
Clostridium ramosum		82.42	0.64		92		
Lactobacillus acidophilus		7.07	0.47		1		
Clostridium barati		4.76	0.41		99		
07371322	**Low discrimination**						**07371322**
					SPO		
Lactobacillus acidophilus		66.60	0.81	RHA 7	1		
Clostridium ramosum		31.38	0.74		92		
Clostridium barati		1.36	0.50		99		
07371323	**Low discrimination**						**07371323**
					SPO		
Lactobacillus acidophilus		65.77	0.47	RHA 7	1		
Clostridium ramosum		30.99	0.40	URE 1	92		
Actinomyces naeslundii		1.46	0.43		1		
07371332	**Low discrimination**						**07371332**
					SPO		
Lactobacillus acidophilus		58.86	0.47	RHA 7	1		
Clostridium ramosum		27.73	0.40	MLZ 1	92		
Clostridium clostridioforme		4.27	0.33		66		
07371342	**Very good identification**						**07371342**
					SPO		
Lactobacillus acidophilus		99.34	0.65	RHA 7	1		
Clostridium barati		0.48	0.23	SOR 11	99		
Actinomyces israelii		0.09	0.28		1		
07371362	**Good identification**						**07371362**
					SPO		
Lactobacillus acidophilus		94.82	0.65	RHA 7	1		
Clostridium ramosum		3.65	0.40	SOR 11	92		
Clostridium barati		1.36	0.31		99		
07371400	**Very good identification**						**07371400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		97.84	0.66	ARA 1			
Lactobacillus catenaformis		0.66	0.29				
Actinomyces israelii		0.55	0.40				
07371402	**Low discrimination**						**07371402**
Lactobacillus jensenii		81.48	0.63	ARA 1			
Lactobacillus acidophilus		15.71	0.48				
Actinomyces israelii		1.03	0.43				
07371410	**Very good identification**						**07371410**
Lactobacillus jensenii		99.75	0.58	ARA 1			
Actinomyces israelii		0.25	0.26				
07371412	**Good identification**						**07371412**
Lactobacillus jensenii		96.01	0.55	ARA 1			
Clostridium clostridioforme		2.22	0.30				
Lactobacillus acidophilus		0.56	0.14				
07371420	**Good identification**						**07371420**
					SPO		
Lactobacillus jensenii		96.49	0.56	ARA 1	1		
Clostridium barati		1.12	0.17		99		
Lactobacillus catenaformis		0.65	0.18		1		
07371422	**Low discrimination**						**07371422**
					SPO		
Lactobacillus jensenii		53.95	0.53	ARA 1	1		
Lactobacillus acidophilus		41.59	0.48		1		
Clostridium barati		2.82	0.25		99		
07371430	**Good identification**						**07371430**
Lactobacillus jensenii		100.0	0.47	ARA 1			
07371432	**Low discrimination**						**07371432**
Lactobacillus jensenii		89.61	0.44	ARA 1			
Clostridium clostridioforme		8.30	0.30				
Lactobacillus acidophilus		2.09	0.14				
07371500	**Low discrimination**						**07371500**
					COC	SPO	
Atopobium parvulum		65.18	0.54	ARA 1	99	1	
Clostridium butyricum		18.11	0.38		1	99	
Actinomyces israelii		5.77	0.50		1	1	
07371502	**Low discrimination**						**07371502**
					COC		
Lactobacillus acidophilus		87.20	0.66	ARA 1	1		
Bifidobacterium adolescentis		4.59	0.53		1		
Atopobium parvulum		2.66	0.41		99		
07371512	**Low discrimination**						**07371512**
					GRA		
Bifidobacterium adolescentis		49.26	0.54	RAF 92	99		
Leptotrichia buccalis		28.75	0.32	XYL 87	1		
Clostridium clostridioforme		8.93	0.39		50		
07371520	**Low discrimination**						**07371520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Atopobium parvulum		35.25	0.41	ARA 1	99	1	
Lactobacillus acidophilus		11.69	0.32		1	1	
Clostridium septicum		9.28	0.25		1	99	
07371521	**Low discrimination**						**07371521**
					COC		
Actinomyces naeslundii		84.45	0.43	RAF 92	1		
Atopobium parvulum		15.55	0.07	ARA 1	99		
07371522	**Good identification**						**07371522**
					SPO		
Lactobacillus acidophilus		93.42	0.66	ARA 1	1		
Clostridium ramosum		2.50	0.38		92		
Clostridium barati		1.65	0.34		99		
07371532	**Low discrimination**						**07371532**
					GRA		
Clostridium clostridioforme		38.35	0.39	RAF 94	50		
Lactobacillus acidophilus		33.24	0.32	XYL 91	99		
Leptotrichia buccalis		13.72	0.15	RHA 88	1		
07371540	**Low discrimination**						**07371540**
					COC	SPO	
Actinomyces israelii		44.09	0.43	XYL 92	1	1	
Clostridium butyricum		30.97	0.19		1	99	
Atopobium parvulum		12.95	0.20		99	1	
07371542	**Low discrimination**						**07371542**
Lactobacillus acidophilus		87.26	0.50	ARA 1			
Actinomyces israelii		6.29	0.46	SOR 11			
Bifidobacterium adolescentis		4.59	0.37				
07371552	**Low discrimination**						**07371552**
Bifidobacterium adolescentis		72.09	0.39	RAF 92			
Actinomyces israelii		12.08	0.32	XYL 87			
Lactobacillus acidophilus		11.33	0.16	SOR 11			
07371562	**Good identification**						**07371562**
					SPO		
Lactobacillus acidophilus		97.13	0.50	ARA 1	1		
Clostridium barati		1.21	0.15	SOR 11	99		
Clostridium perfringens		0.55	0.14		84		
07371612	**Acceptable identification**						**07371612**
Clostridium clostridioforme		94.38	0.44	RAF 94			
Lactobacillus jensenii		5.62	0.21	XYL 91			
07371632	**Good identification**						**07371632**
Clostridium clostridioforme		98.53	0.44	RAF 94			
Lactobacillus jensenii		1.47	0.10	XYL 91			
07371702	**Low discrimination**						**07371702**
Lactobacillus acidophilus		76.77	0.47	RHA 7			
Clostridium clostridioforme		19.13	0.42	ARA 1			
Actinomyces israelii		1.76	0.34				
07371712	**Good identification**						**07371712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		97.66	0.53	RAF 94		
Lactobacillus acidophilus		0.87	0.13	XYL 91		
Bifidobacterium adolescentis		0.74	0.20			
07371722	**Low discrimination**					**07371722**
					SPO	
Lactobacillus acidophilus		55.76	0.47	RHA 7	1	
Clostridium ramosum		26.27	0.40	ARA 1	92	
Clostridium clostridioforme		13.90	0.42		66	
07371732	**Good identification**					**07371732**
					GRA	
Clostridium clostridioforme		96.69	0.53	RAF 94	50	
Parabacteroides distasonis		2.05	0.30	XYL 91	1	
Lactobacillus acidophilus		0.86	0.13		99	
07372000	**Low discrimination**					**07372000**
					SPO	
Clostridium innocuum		60.34	0.36	MLT 1	99	
Lactobacillus jensenii		23.32	0.32	GAL 95	1	
Actinomyces naeslundii		10.70	0.47		1	
07372001	**Good identification**					**07372001**
					SPO	
Actinomyces naeslundii		96.34	0.50	RAF 92	1	
Clostridium innocuum		3.66	0.02	GAL 96	99	
07372002	**Low discrimination**					**07372002**
					SPO	
Clostridium innocuum		55.55	0.36	MLT 1	99	
Lactobacillus acidophilus		14.38	0.26	GAL 95	1	
Lactobacillus jensenii		14.31	0.29		1	
07372003	**Good identification**					**07372003**
					SPO	
Actinomyces naeslundii		96.34	0.50	RAF 92	1	
Clostridium innocuum		3.66	0.02	GAL 96	99	
07372020	**Low discrimination**					**07372020**
					SPO	
Actinomyces naeslundii		45.53	0.47	RAF 92	1	
Lactobacillus jensenii		24.81	0.21	GAL 96	1	
Clostridium barati		12.92	0.11		99	
07372021	**Good identification**					**07372021**
Actinomyces naeslundii		100.0	0.50	RAF 92		
07372023	**Good identification**					**07372023**
Actinomyces naeslundii		100.0	0.50	RAF 92		
07372041	**Good identification**					**07372041**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07372043	**Good identification**					**07372043**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07372061	**Good identification**					**07372061**
Actinomyces naeslundii		100.0	0.38	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07372063	**Good identification**					**07372063**
Actinomyces naeslundii		100.0	0.38	RAF 92		
07372101	**Very good identification**					**07372101**
					SPO	
Actinomyces naeslundii		99.00	0.58	RAF 92	1	
Clostridium tertium		1.00	0.01	GAL 96	99	
07372102	**Low discrimination**					**07372102**
					SPO	
Lactobacillus acidophilus		48.65	0.44	GAL 99	1	
Clostridium tertium		37.16	0.43	MAN 5	99	
Actinomyces naeslundii		8.25	0.55		1	
07372103	**Good identification**					**07372103**
					SPO	
Actinomyces naeslundii		93.45	0.58	RAF 92	1	
Lactobacillus acidophilus		3.71	0.10	GAL 96	1	
Clostridium tertium		2.84	0.09		99	
07372110	**Good identification**					**07372110**
					SPO	
Clostridium tertium		97.43	0.39	GAL 99	99	
Actinomyces naeslundii		1.92	0.33		1	
Actinomyces israelii		0.65	0.16		1	
07372111	**Low discrimination**					**07372111**
					SPO	
Actinomyces naeslundii		74.54	0.36	RAF 92	1	
Clostridium tertium		25.46	0.05	GAL 96	99	
07372112	**Good identification**					**07372112**
					SPO	
Clostridium tertium		97.78	0.47	GAL 99	99	
Lactobacillus acidophilus		0.73	0.10		1	
Actinomyces naeslundii		0.64	0.33		1	
07372120	**Low discrimination**					**07372120**
					SPO	
Clostridium tertium		44.41	0.43	GAL 99	99	
Clostridium ramosum		41.86	0.41		92	
Actinomyces naeslundii		9.86	0.55		1	
07372121	**Good identification**					**07372121**
					SPO	
Actinomyces naeslundii		94.44	0.58	RAF 92	1	
Clostridium tertium		2.87	0.09	GAL 96	99	
Clostridium ramosum		2.70	0.07		92	
07372122	**Low discrimination**					**07372122**
					SPO	
Clostridium ramosum		44.35	0.51	GAL 99	92	
Clostridium tertium		35.29	0.51		99	
Lactobacillus acidophilus		15.40	0.44		1	
07372123	**Low discrimination**					**07372123**
					SPO	
Actinomyces naeslundii		80.31	0.58	RAF 92	1	
Clostridium ramosum		9.19	0.17	GAL 96	92	
Clostridium tertium		7.31	0.17		99	
07372130	**Good identification**					**07372130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium tertium		98.82	0.47	GAL 99	99	
Actinomyces naeslundii		0.65	0.33		1	
Clostridium ramosum		0.53	0.07		92	
07372132	**Good identification**					**07372132**
					SPO	
Clostridium tertium		98.83	0.55	GAL 99	99	
Clostridium ramosum		0.71	0.17		92	
Lactobacillus acidophilus		0.25	0.10		1	
07372140	**Low discrimination**					**07372140**
					SPO	
Actinomyces naeslundii		74.08	0.42	RAF 92	1	
Actinomyces israelii		19.55	0.23	GAL 96	1	
Clostridium tertium		6.37	0.01		99	
07372141	**Good identification**					**07372141**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07372143	**Good identification**					**07372143**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07372160	**Low discrimination**					**07372160**
					SPO	
Actinomyces naeslundii		60.67	0.42	RAF 92	1	
Clostridium tertium		15.64	0.09	GAL 96	99	
Clostridium ramosum		14.75	0.07		92	
07372161	**Good identification**					**07372161**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07372163	**Good identification**					**07372163**
Actinomyces naeslundii		100.0	0.45	RAF 92		
07372320	**Good identification**					**07372320**
Clostridium ramosum		98.75	0.43	GAL 99		
Clostridium tertium		0.80	0.09			
Clostridium barati		0.46	0.01			
07372322	**Good identification**					**07372322**
					SPO	
Clostridium ramosum		97.16	0.53	GAL 99	92	
Lactobacillus acidophilus		1.92	0.25		1	
Clostridium tertium		0.59	0.17		99	
07373000	**Low discrimination**					**07373000**
					SPO	
Lactobacillus jensenii		59.06	0.66	MAN 1	1	
Clostridium innocuum		29.33	0.58		99	
Actinomyces naeslundii		6.57	0.71		1	
07373001	**Good identification**					**07373001**
					SPO	
Actinomyces naeslundii		91.39	0.74	RAF 92	1	
Lactobacillus jensenii		5.53	0.32		1	
Clostridium innocuum		2.75	0.24		99	
07373002	**Low discrimination**					**07373002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Lactobacillus acidophilus		31.93	0.60	MAN 5	1	
Lactobacillus jensenii		31.80	0.63	LAC 92	1	
Clostridium innocuum		23.69	0.58		99	
07373003	**Low discrimination**					**07373003**
Actinomyces naeslundii		89.31	0.74	RAF 92		
Lactobacillus acidophilus		3.62	0.26			
Lactobacillus jensenii		3.61	0.29			
07373010	**Low discrimination**					**07373010**
					SPO	
Lactobacillus jensenii		83.14	0.58	MAN 1	1	
Clostridium tertium		8.78	0.39		99	
Clostridium innocuum		5.16	0.34		99	
07373011	**Low discrimination**					**07373011**
					SPO	
Actinomyces naeslundii		69.07	0.52	RAF 92	1	
Lactobacillus jensenii		26.49	0.24	MLZ 5	1	
Clostridium tertium		2.80	0.05		99	
07373012	**Low discrimination**					**07373012**
					SPO	
Lactobacillus jensenii		59.94	0.55	MAN 1	1	
Clostridium tertium		28.49	0.47		99	
Clostridium innocuum		5.58	0.34		99	
07373013	**Low discrimination**					**07373013**
					SPO	
Actinomyces naeslundii		71.38	0.52	RAF 92	1	
Lactobacillus jensenii		18.25	0.21	MLZ 5	1	
Clostridium tertium		8.67	0.13		99	
07373020	**Low discrimination**					**07373020**
					SPO	
Lactobacillus jensenii		48.09	0.56	MAN 1	1	
Actinomyces naeslundii		21.39	0.71		1	
Clostridium tertium		11.43	0.43		99	
07373021	**Very good identification**					**07373021**
					SPO	
Actinomyces naeslundii		97.69	0.74	RAF 92	1	
Lactobacillus jensenii		1.48	0.21		1	
Clostridium tertium		0.35	0.09		99	
07373022	**Low discrimination**					**07373022**
					SPO	
Lactobacillus acidophilus		46.63	0.60	MAN 5	1	
Clostridium ramosum		15.60	0.51	LAC 92	92	
Clostridium tertium		12.41	0.51		99	
07373023	**Good identification**					**07373023**
					SPO	
Actinomyces naeslundii		92.63	0.74	RAF 92	1	
Lactobacillus acidophilus		3.76	0.26		1	
Clostridium ramosum		1.26	0.17		92	
07373030	**Low discrimination**					**07373030**
					SPO	
Clostridium tertium		53.83	0.47	LAC 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		42.47	0.47		1	
Actinomyces naeslundii		2.98	0.49		1	
07373031	**Low discrimination**					**07373031**
					SPO	
Actinomyces naeslundii		82.14	0.52	RAF 92	1	
Clostridium tertium		9.98	0.13	MLZ 5	99	
Lactobacillus jensenii		7.88	0.13		1	
07373032	**Low discrimination**					**07373032**
					SPO	
Clostridium tertium		81.49	0.55	LAC 99	99	
Lactobacillus jensenii		14.29	0.44		1	
Lactobacillus acidophilus		1.74	0.26		1	
07373033	**Low discrimination**					**07373033**
					SPO	
Actinomyces naeslundii		70.00	0.52	RAF 92	1	
Clostridium tertium		25.52	0.21	MLZ 5	99	
Lactobacillus jensenii		4.48	0.10		1	
07373040	**Low discrimination**					**07373040**
Actinomyces naeslundii		37.96	0.58	RAF 92		
Actinomyces israelii		29.40	0.47			
Lactobacillus jensenii		19.54	0.32			
07373041	**Very good identification**					**07373041**
Actinomyces naeslundii		100.0	0.61	RAF 92		
07373042	**Low discrimination**					**07373042**
Lactobacillus acidophilus		59.24	0.44	MAN 5		
Actinomyces israelii		16.32	0.50	LAC 92		
Actinomyces naeslundii		14.05	0.58	SOR 11		
07373043	**Very good identification**					**07373043**
Actinomyces naeslundii		96.51	0.61	RAF 92		
Lactobacillus acidophilus		2.74	0.10			
Actinomyces israelii		0.76	0.16			
07373051	**Good identification**					**07373051**
Actinomyces naeslundii		100.0	0.39	RAF 92		
07373052	**Low discrimination**					**07373052**
					SPO	
Actinomyces israelii		38.73	0.36	XYL 92	1	
Lactobacillus jensenii		25.52	0.21	MLZ 13	1	
Clostridium tertium		12.13	0.13		99	
07373053	**Good identification**					**07373053**
Actinomyces naeslundii		100.0	0.39	RAF 92		
07373060	**Low discrimination**					**07373060**
					SPO	
Actinomyces naeslundii		71.12	0.58	RAF 92	1	
Lactobacillus jensenii		9.15	0.21		1	
Clostridium barati		7.87	0.15		99	
07373061	**Very good identification**					**07373061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.61	RAF 92			
07373062	**Low discrimination**						**07373062**
					SPO		
Lactobacillus acidophilus		71.28	0.44	MAN 5	1		
Actinomyces naeslundii		16.90	0.58	LAC 92	1		
Clostridium barati		5.61	0.23	SOR 11	99		
07373063	**Very good identification**						**07373063**
Actinomyces naeslundii		97.24	0.61	RAF 92			
Lactobacillus acidophilus		2.76	0.10				
07373071	**Good identification**						**07373071**
Actinomyces naeslundii		100.0	0.39	RAF 92			
07373073	**Good identification**						**07373073**
Actinomyces naeslundii		100.0	0.39	RAF 92			
07373100	**Low discrimination**						**07373100**
					COC	SPO	
Clostridium tertium		68.16	0.69		1	99	
Actinomyces naeslundii		11.00	0.79		1	1	
Atopobium parvulum		8.87	0.54		99	1	
07373101	**Good identification**						**07373101**
					COC	SPO	
Actinomyces naeslundii		94.89	0.82	RAF 92	1	1	
Clostridium tertium		3.96	0.35		1	99	
Atopobium parvulum		0.52	0.20		99	1	
07373102	**Low discrimination**						**07373102**
					SPO		
Lactobacillus acidophilus		53.04	0.78	MAN 5	1		
Clostridium tertium		40.52	0.77		99		
Actinomyces naeslundii		2.18	0.79		1		
07373103	**Low discrimination**						**07373103**
					SPO		
Actinomyces naeslundii		76.85	0.82	RAF 92	1		
Lactobacillus acidophilus		12.59	0.44		1		
Clostridium tertium		9.62	0.43		99		
07373110	**Very good identification**						**07373110**
					SPO		
Clostridium tertium		98.61	0.73		99		
Actinomyces israelii		0.66	0.50		1		
Actinomyces naeslundii		0.47	0.57		1		
07373111	**Low discrimination**						**07373111**
					SPO		
Clostridium tertium		58.26	0.39	URE 1	99		
Actinomyces naeslundii		41.35	0.60		1		
Actinomyces israelii		0.39	0.16		1		
07373112	**Very good identification**						**07373112**
					GRA	SPO	
Clostridium tertium		98.03	0.81		99	99	
Lactobacillus acidophilus		0.73	0.44		99	1	
Leptotrichia buccalis		0.52	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07373113	**Low discrimination**					**07373113**
					SPO	
Clostridium tertium		79.83	0.47	URE 1	99	
Actinomyces naeslundii		18.89	0.60		1	
Lactobacillus acidophilus		0.59	0.10		1	
07373120	**Low discrimination**					**07373120**
					SPO	
Clostridium tertium		49.05	0.77		99	
Clostridium ramosum		46.24	0.75		92	
Actinomyces naeslundii		2.64	0.79		1	
07373121	**Low discrimination**					**07373121**
					SPO	
Actinomyces naeslundii		80.17	0.82	RAF 92	1	
Clostridium tertium		10.03	0.43		99	
Clostridium ramosum		9.46	0.41		92	
07373122	**Low discrimination**					**07373122**
					SPO	
Clostridium ramosum		46.09	0.85		92	
Clostridium tertium		36.67	0.85		99	
Lactobacillus acidophilus		16.00	0.78		1	
07373123	**Low discrimination**					**07373123**
					SPO	
Actinomyces naeslundii		49.61	0.82	RAF 92	1	
Clostridium ramosum		23.41	0.51		92	
Clostridium tertium		18.62	0.51		99	
07373130	**Very good identification**					**07373130**
					SPO	
Clostridium tertium		99.26	0.81		99	
Clostridium ramosum		0.53	0.41		92	
Actinomyces naeslundii		0.16	0.57		1	
07373131	**Low discrimination**					**07373131**
					SPO	
Clostridium tertium		80.52	0.47	URE 1	99	
Actinomyces naeslundii		19.05	0.60		1	
Clostridium ramosum		0.43	0.07		92	
07373132	**Very good identification**					**07373132**
					SPO	
Clostridium tertium		98.92	0.89		99	
Clostridium ramosum		0.71	0.51		92	
Lactobacillus acidophilus		0.25	0.44		1	
07373133	**Acceptable identification**					**07373133**
					SPO	
Clostridium tertium		91.87	0.55	URE 1	99	
Actinomyces naeslundii		7.25	0.60		1	
Clostridium ramosum		0.66	0.17		92	
07373140	**Low discrimination**					**07373140**
					SPO	
Actinomyces israelii		39.96	0.57	XYL 92	1	
Actinomyces naeslundii		36.71	0.66		1	
Clostridium tertium		13.02	0.35		99	
07373141	**Very good identification**					**07373141**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.04	0.69	RAF 92	1	
Actinomyces israelii		0.73	0.23		1	
Clostridium tertium		0.24	0.01		99	
07373142	**Low discrimination**					**07373142**
					SPO	
Lactobacillus acidophilus		80.89	0.63	MAN 5	1	
Actinomyces israelii		7.75	0.60	SOR 11	1	
Clostridium tertium		5.05	0.43		99	
07373143	**Low discrimination**					**07373143**
Actinomyces naeslundii		88.27	0.69	RAF 92		
Lactobacillus acidophilus		10.13	0.28			
Actinomyces israelii		0.97	0.26			
07373150	**Low discrimination**					**07373150**
					SPO	
Clostridium tertium		73.36	0.39	SOR 1	99	
Actinomyces israelii		18.93	0.43		1	
Actinomyces naeslundii		6.13	0.44		1	
07373151	**Acceptable identification**					**07373151**
					SPO	
Actinomyces naeslundii		92.54	0.47	RAF 92	1	
Clostridium tertium		7.46	0.05	MLZ 5	99	
07373152	**Low discrimination**					**07373152**
					SPO	
Clostridium tertium		74.64	0.47	SOR 1	99	
Actinomyces israelii		9.63	0.46		1	
Lactobacillus acidophilus		6.79	0.28		1	
07373153	**Low discrimination**					**07373153**
					SPO	
Actinomyces naeslundii		80.52	0.47	RAF 92	1	
Clostridium tertium		19.48	0.13	MLZ 5	99	
07373160	**Low discrimination**					**07373160**
					SPO	
Clostridium tertium		30.30	0.43	SOR 1	99	
Clostridium ramosum		28.56	0.41		92	
Actinomyces naeslundii		28.48	0.66		1	
07373161	**Very good identification**					**07373161**
					SPO	
Actinomyces naeslundii		98.63	0.69	RAF 92	1	
Clostridium tertium		0.71	0.09		99	
Clostridium ramosum		0.67	0.07		92	
07373162	**Low discrimination**					**07373162**
					SPO	
Lactobacillus acidophilus		65.62	0.63	MAN 5	1	
Clostridium ramosum		15.45	0.51	SOR 11	92	
Clostridium tertium		12.29	0.51		99	
07373163	**Low discrimination**					**07373163**
					SPO	
Actinomyces naeslundii		85.79	0.69	RAF 92	1	
Lactobacillus acidophilus		9.84	0.28		1	
Clostridium ramosum		2.32	0.17		92	
07373170	**Good identification**					**07373170**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium tertium		95.10	0.47	SOR 1	99	
Actinomyces naeslundii		2.65	0.44		1	
Bifidobacterium breve		1.22	0.24		1	
07373171	**Low discrimination**					**07373171**
					SPO	
Actinomyces naeslundii		80.52	0.47	RAF 92	1	
Clostridium tertium		19.48	0.13	MLZ 5	99	
07373172	**Acceptable identification**					**07373172**
					SPO	
Clostridium tertium		91.46	0.55	SOR 1	99	
Bifidobacterium breve		3.95	0.41		1	
Lactobacillus acidophilus		2.78	0.28		1	
07373173	**Low discrimination**					**07373173**
					SPO	
Actinomyces naeslundii		57.94	0.47	RAF 92	1	
Clostridium tertium		42.06	0.21	MLZ 5	99	
07373201	**Good identification**					**07373201**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07373202	**Low discrimination**					**07373202**
					SPO	
Lactobacillus acidophilus		68.61	0.41	MAN 5	1	
Lactobacillus jensenii		9.17	0.29	LAC 92	1	
Clostridium innocuum		6.83	0.24	RHA 7	99	
07373203	**Low discrimination**					**07373203**
Actinomyces naeslundii		76.80	0.40	RAF 92		
Lactobacillus acidophilus		23.20	0.07	RHA 1		
07373220	**Acceptable identification**					**07373220**
					SPO	
Clostridium ramosum		91.25	0.43	LAC 99	92	
Lactobacillus jensenii		3.10	0.21		1	
Clostridium barati		2.67	0.15		99	
07373221	**Low discrimination**					**07373221**
					SPO	
Actinomyces naeslundii		69.20	0.40	RAF 92	1	
Clostridium ramosum		30.80	0.09	RHA 1	92	
07373222	**Low discrimination**					**07373222**
					SPO	
Clostridium ramosum		82.77	0.53	LAC 99	92	
Lactobacillus acidophilus		14.05	0.41		1	
Clostridium barati		1.82	0.23		99	
07373300	**Low discrimination**					**07373300**
					SPO	
Clostridium ramosum		62.07	0.43	NAG 99	92	
Clostridium tertium		16.56	0.35		99	
Actinomyces israelii		9.84	0.45		1	
07373301	**Low discrimination**					**07373301**
					SPO	
Actinomyces naeslundii		83.47	0.48	RAF 92	1	
Clostridium ramosum		13.05	0.09	RHA 1	92	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		3.48	0.01		99	
07373302	**Low discrimination**					**07373302**
					SPO	
Lactobacillus acidophilus		60.14	0.59	MAN 5	1	
Clostridium ramosum		30.81	0.53	RHA 7	92	
Clostridium tertium		6.17	0.43		99	
07373310	**Acceptable identification**					**07373310**
					SPO	
Clostridium tertium		92.94	0.39	RHA 1	99	
Actinomyces israelii		4.64	0.31		1	
Clostridium ramosum		1.98	0.09		92	
07373312	**Low discrimination**					**07373312**
					SPO	
Clostridium tertium		89.63	0.47	RHA 1	99	
Lactobacillus acidophilus		4.97	0.25		1	
Clostridium ramosum		2.55	0.19		92	
07373320	**Very good identification**					**07373320**
Clostridium ramosum		98.97	0.77			
Clostridium tertium		0.80	0.43			
Clostridium barati		0.09	0.23			
07373321	**Acceptable identification**					**07373321**
					SPO	
Clostridium ramosum		93.22	0.43	URE 1	92	
Actinomyces naeslundii		6.02	0.48		1	
Clostridium tertium		0.75	0.09		99	
07373322	**Very good identification**					**07373322**
					SPO	
Clostridium ramosum		97.41	0.87		92	
Lactobacillus acidophilus		1.92	0.59		1	
Clostridium tertium		0.59	0.51		99	
07373323	**Good identification**					**07373323**
					SPO	
Clostridium ramosum		95.92	0.53	URE 1	92	
Lactobacillus acidophilus		1.89	0.25		1	
Actinomyces naeslundii		1.55	0.48		1	
07373330	**Low discrimination**					**07373330**
					SPO	
Clostridium tertium		58.67	0.47	RHA 1	99	
Clostridium ramosum		41.24	0.43		92	
Actinomyces naeslundii		0.09	0.23		1	
07373332	**Low discrimination**					**07373332**
					SPO	
Clostridium tertium		51.09	0.55	RHA 1	99	
Clostridium ramosum		47.88	0.53		92	
Lactobacillus acidophilus		0.94	0.25		1	
07373340	**Low discrimination**					**07373340**
					SPO	
Actinomyces israelii		65.35	0.38	XYL 92	1	
Clostridium ramosum		10.71	0.09	RHA 7	92	
Lactobacillus acidophilus		10.33	0.09		1	
07373342	**Low discrimination**					**07373342**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Lactobacillus acidophilus		86.07	0.43	MAN 5	1	
Actinomyces israelii		8.25	0.41	RHA 7	1	
Clostridium ramosum		3.60	0.19	SOR 11	92	
07373360	**Good identification**					**07373360**
					SPO	
Clostridium ramosum		96.43	0.43	SOR 1	92	
Lactobacillus acidophilus		0.94	0.09		1	
Clostridium tertium		0.78	0.09		99	
07373362	**Low discrimination**					**07373362**
					SPO	
Clostridium ramosum		79.59	0.53	SOR 1	92	
Lactobacillus acidophilus		19.20	0.43		1	
Clostridium tertium		0.48	0.17		99	
07373400	**Low discrimination**					**07373400**
					SPO	
Clostridium innocuum		66.71	0.45	MLT 1	99	
Actinomyces israelii		24.34	0.53		1	
Lactobacillus jensenii		7.69	0.32		1	
07373401	**Low discrimination**					**07373401**
					SPO	
Actinomyces naeslundii		58.24	0.40	RAF 92	1	
Clostridium innocuum		30.60	0.11	ARA 1	99	
Actinomyces israelii		11.17	0.18		1	
07373402	**Low discrimination**					**07373402**
					SPO	
Clostridium innocuum		57.76	0.45	MLT 1	99	
Actinomyces israelii		31.61	0.56		1	
Lactobacillus acidophilus		4.46	0.26		1	
07373403	**Low discrimination**					**07373403**
					SPO	
Actinomyces naeslundii		55.16	0.40	RAF 92	1	
Clostridium innocuum		28.98	0.11	ARA 1	99	
Actinomyces israelii		15.86	0.21		1	
07373410	**Low discrimination**					**07373410**
					SPO	
Actinomyces israelii		39.32	0.38	XYL 92	1	
Clostridium innocuum		30.05	0.22	MLZ 13	99	
Lactobacillus jensenii		27.71	0.24		1	
07373412	**Low discrimination**					**07373412**
					SPO	
Actinomyces israelii		48.05	0.42	XYL 92	1	
Clostridium innocuum		24.48	0.22	MLZ 13	99	
Lactobacillus jensenii		15.05	0.21		1	
07373421	**Good identification**					**07373421**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07373423	**Good identification**					**07373423**
Actinomyces naeslundii		100.0	0.40	RAF 92		
07373440	**Acceptable identification**					**07373440**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		91.99	0.46	XYL 92	1	
Clostridium innocuum		6.55	0.11		99	
Actinomyces naeslundii		1.47	0.24		1	
07373442	**Acceptable identification**					**07373442**
					SPO	
Actinomyces israelii		90.67	0.49	XYL 92	1	
Clostridium innocuum		4.30	0.11		99	
Lactobacillus acidophilus		4.06	0.10		1	
07373500	**Low discrimination**					**07373500**
					SPO	
Actinomyces israelii		77.76	0.63	XYL 92	1	
Clostridium tertium		12.04	0.35		99	
Clostridium butyricum		4.79	0.21		99	
07373501	**Low discrimination**					**07373501**
					SPO	
Actinomyces naeslundii		76.27	0.48	RAF 92	1	
Actinomyces israelii		20.55	0.29	ARA 1	1	
Clostridium tertium		3.18	0.01		99	
07373502	**Low discrimination**					**07373502**
Actinomyces israelii		47.57	0.66	XYL 92		
Lactobacillus acidophilus		19.29	0.44			
Bifidobacterium adolescentis		15.77	0.51			
07373503	**Low discrimination**					**07373503**
Actinomyces naeslundii		54.73	0.48	RAF 92		
Actinomyces israelii		22.12	0.32	ARA 1		
Lactobacillus acidophilus		8.97	0.10			
07373510	**Low discrimination**					**07373510**
					SPO	
Clostridium tertium		63.22	0.39	ARA 1	99	
Actinomyces israelii		34.32	0.49		1	
Bifidobacterium adolescentis		1.41	0.19		1	
07373512	**Low discrimination**					**07373512**
					SPO	
Clostridium tertium		49.04	0.47	ARA 1	99	
Bifidobacterium adolescentis		36.10	0.53		1	
Actinomyces israelii		13.31	0.52		1	
07373520	**Low discrimination**					**07373520**
					SPO	
Clostridium tertium		45.89	0.43	ARA 1	99	
Clostridium ramosum		43.26	0.41		92	
Actinomyces israelii		6.31	0.42		1	
07373521	**Low discrimination**					**07373521**
					SPO	
Actinomyces naeslundii		80.44	0.48	RAF 92	1	
Clostridium tertium		10.07	0.09	ARA 1	99	
Clostridium ramosum		9.49	0.07		92	
07373522	**Low discrimination**					**07373522**
					SPO	
Clostridium ramosum		44.92	0.51	ARA 1	92	
Clostridium tertium		35.75	0.51		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		15.60	0.44		1		
07373523	**Low discrimination**						**07373523**
					SPO		
Actinomyces naeslundii		49.72	0.48	RAF 92	1		
Clostridium ramosum		23.46	0.17	ARA 1	92		
Clostridium tertium		18.67	0.17		99		
07373530	**Good identification**						**07373530**
					SPO		
Clostridium tertium		97.94	0.47	ARA 1	99		
Actinomyces israelii		1.13	0.28		1		
Clostridium ramosum		0.53	0.07		92		
07373532	**Good identification**						**07373532**
					SPO		
Clostridium tertium		97.34	0.55	ARA 1	99		
Bifidobacterium dentium		0.82	0.10		1		
Clostridium ramosum		0.70	0.17		92		
07373540	**Good identification**						**07373540**
					SPO		
Actinomyces israelii		97.17	0.56	XYL 92	1		
Clostridium butyricum		1.34	0.03		99		
Actinomyces naeslundii		1.10	0.32		1		
07373542	**Low discrimination**						**07373542**
Actinomyces israelii		79.78	0.59	XYL 92			
Lactobacillus acidophilus		10.28	0.28				
Bifidobacterium adolescentis		8.41	0.36				
07373550	**Acceptable identification**						**07373550**
					COC	SPO	
Actinomyces israelii		92.09	0.42	XYL 92	1	1	
Clostridium tertium		4.41	0.05	MLZ 13	1	99	
Blautia producta		3.51	-.03		99	1	
07373552	**Low discrimination**						**07373552**
					SPO		
Actinomyces israelii		51.08	0.45	XYL 92	1		
Bifidobacterium adolescentis		44.03	0.37	MLZ 13	1		
Clostridium tertium		4.89	0.13		99		
07373560	**Low discrimination**						**07373560**
					SPO		
Actinomyces israelii		64.73	0.35	XYL 92	1		
Clostridium tertium		12.24	0.09	NAG 6	99		
Clostridium ramosum		11.53	0.07		92		
07373602	**Low discrimination**						**07373602**
					SPO		
Actinomyces israelii		72.14	0.36	XYL 92	1		
Clostridium innocuum		17.69	0.11	RHA 7	99		
Lactobacillus acidophilus		10.17	0.07		1		
07373700	**Low discrimination**						**07373700**
					SPO		
Actinomyces israelii		88.97	0.44	XYL 92	1		
Clostridium ramosum		6.93	0.09	RHA 7	92		
Clostridium butyricum		2.25	-.05		99		
07373702	**Low discrimination**						**07373702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		58.19	0.47	XYL 92	1		
Lactobacillus acidophilus		23.59	0.25	RHA 7	1		
Clostridium ramosum		12.09	0.19		92		
07373720	**Good identification**						**07373720**
					SPO		
Clostridium ramosum		98.39	0.43	ARA 1	92		
Actinomyces israelii		0.81	0.23		1		
Clostridium tertium		0.80	0.09		99		
07373722	**Good identification**						**07373722**
					SPO		
Clostridium ramosum		97.04	0.53	ARA 1	92		
Lactobacillus acidophilus		1.91	0.25		1		
Clostridium tertium		0.59	0.17		99		
07373740	**Good identification**						**07373740**
Actinomyces israelii		100.0	0.37	XYL 92			
07373742	**Low discrimination**						**07373742**
Actinomyces israelii		88.59	0.40	XYL 92			
Lactobacillus acidophilus		11.41	0.09	RHA 7			
07374100	**Very good identification**						**07374100**
					COC	GRA	
Atopobium parvulum		97.62	0.59	GAL 99	99	99	
Fusobacterium mortiferum		1.13	0.37		1	1	
Actinomyces israelii		0.66	0.36		1	99	
07374102	**Low discrimination**						**07374102**
					COC		
Atopobium parvulum		77.53	0.46	GAL 99	99		
Lactobacillus acidophilus		13.84	0.32		1		
Actinomyces israelii		4.46	0.39		1		
07374120	**Good identification**						**07374120**
					COC	SPO	
Atopobium parvulum		97.99	0.46	GAL 99	99	1	
Clostridium tertium		0.97	0.11		1	99	
Clostridium barati		0.54	0.04		1	99	
07374312	**Good identification**						**07374312**
Clostridium clostridioforme		100.0	0.42	RAF 94			
07374332	**Good identification**						**07374332**
Clostridium clostridioforme		98.56	0.42	RAF 94			
Parabacteroides distasonis		1.44	0.16	ARA 94			
07374412	**Good identification**						**07374412**
Clostridium clostridioforme		100.0	0.39	RAF 94			
07374432	**Good identification**						**07374432**
Clostridium clostridioforme		100.0	0.39	RAF 94			
07374502	**Low discrimination**						**07374502**
					SPO		
Clostridium clostridioforme		40.09	0.36	RAF 94	66		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		28.34	0.22	RHA 88	99	
Bifidobacterium adolescentis		16.54	0.33		1	
07374512	**Low discrimination**					**07374512**
Clostridium clostridioforme		88.58	0.48	RAF 94		
Bifidobacterium adolescentis		9.80	0.34	RHA 88		
Actinomyces israelii		0.90	0.23			
07374522	**Acceptable identification**					**07374522**
					SPO	
Clostridium clostridioforme		92.49	0.36	RAF 94	66	
Clostridium butyricum		5.69	0.04	RHA 88	99	
Actinomyces israelii		1.82	0.17		1	
07374532	**Good identification**					**07374532**
Clostridium clostridioforme		100.0	0.48	RAF 94		
07374602	**Good identification**					**07374602**
Clostridium clostridioforme		100.0	0.42	RAF 94		
07374612	**Good identification**					**07374612**
Clostridium clostridioforme		100.0	0.53	RAF 94		
07374622	**Good identification**					**07374622**
Clostridium clostridioforme		100.0	0.42	RAF 94		
07374632	**Good identification**					**07374632**
Clostridium clostridioforme		100.0	0.53	RAF 94		
07374702	**Good identification**					**07374702**
					SPO	
Clostridium clostridioforme		99.39	0.51	RAF 94	66	
Actinomyces israelii		0.32	0.18		1	
Clostridium butyricum		0.30	-.04		99	
07374712	**Very good identification**					**07374712**
Clostridium clostridioforme		100.0	0.62	RAF 94		
07374722	**Good identification**					**07374722**
Clostridium clostridioforme		99.52	0.51	RAF 94		
Parabacteroides distasonis		0.48	0.17			
07374732	**Very good identification**					**07374732**
Clostridium clostridioforme		99.94	0.62	RAF 94		
Parabacteroides distasonis		0.06	0.13			
07375000	**Low discrimination**					**07375000**
					COC	
Lactobacillus jensenii		63.39	0.66	XYL 1	1	
Atopobium parvulum		30.04	0.59		99	
Actinomyces israelii		5.03	0.59		1	
07375002	**Low discrimination**					**07375002**
Lactobacillus jensenii		65.68	0.63	XYL 1		
Lactobacillus acidophilus		12.66	0.48			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		11.72	0.62				
07375010	**Good identification**						**07375010**
					COC		
Lactobacillus jensenii		94.96	0.58	XYL 1	1		
Actinomyces israelii		3.38	0.45		1		
Atopobium parvulum		1.36	0.25		99		
07375012	**Low discrimination**						**07375012**
Lactobacillus jensenii		89.36	0.55	XYL 1			
Actinomyces israelii		7.15	0.48				
Clostridium clostridioforme		1.34	0.26				
07375020	**Low discrimination**						**07375020**
					COC		
Lactobacillus jensenii		71.74	0.56	XYL 1	1		
Atopobium parvulum		24.00	0.46		99		
Actinomyces israelii		1.45	0.39		1		
07375022	**Low discrimination**						**07375022**
					COC		
Lactobacillus jensenii		49.78	0.53	XYL 1	1		
Lactobacillus acidophilus		38.38	0.48		1		
Atopobium parvulum		4.41	0.33		99		
07375030	**Good identification**						**07375030**
					COC	SPO	
Lactobacillus jensenii		96.44	0.47	XYL 1	1	1	
Clostridium tertium		1.71	0.15		1	99	
Atopobium parvulum		0.98	0.12		99	1	
07375032	**Low discrimination**						**07375032**
					SPO		
Lactobacillus jensenii		84.52	0.44	XYL 1	1		
Clostridium tertium		6.73	0.24		99		
Clostridium clostridioforme		5.05	0.26		66		
07375040	**Low discrimination**						**07375040**
					COC		
Actinomyces israelii		65.89	0.52		1		
Lactobacillus jensenii		21.58	0.32		1		
Atopobium parvulum		10.23	0.25		99		
07375042	**Low discrimination**						**07375042**
Actinomyces israelii		65.78	0.55				
Lactobacillus acidophilus		22.58	0.32				
Lactobacillus jensenii		9.57	0.29				
07375050	**Low discrimination**						**07375050**
Actinomyces israelii		57.79	0.38	MLZ 13			
Lactobacillus jensenii		42.22	0.24				
07375052	**Low discrimination**						**07375052**
Actinomyces israelii		74.19	0.41	MLZ 13			
Lactobacillus jensenii		24.09	0.21				
Lactobacillus acidophilus		1.72	-.02				
07375100	**Very good identification**						**07375100**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		98.64	0.93		99	1	
Actinomyces israelii		0.67	0.70		1	1	
Clostridium butyricum		0.54	0.47		1	99	
07375101	**Very good identification**						**07375101**
					COC	SPO	
Atopobium parvulum		98.23	0.59	URE 1	99	1	
Actinomyces israelii		0.66	0.36		1	1	
Clostridium butyricum		0.53	0.13		1	99	
07375102	**Low discrimination**						**07375102**
					COC		
Atopobium parvulum		78.67	0.80		99		
Lactobacillus acidophilus		14.04	0.66		1		
Actinomyces israelii		4.52	0.73		1		
07375103	**Low discrimination**						**07375103**
					COC		
Atopobium parvulum		77.11	0.46	URE 1	99		
Lactobacillus acidophilus		13.76	0.32		1		
Actinomyces israelii		4.43	0.39		1		
07375110	**Low discrimination**						**07375110**
					COC	SPO	
Atopobium parvulum		80.57	0.59	MLZ 1	99	1	
Clostridium tertium		8.27	0.41		1	99	
Actinomyces israelii		8.06	0.56		1	1	
07375112	**Low discrimination**						**07375112**
					COC	SPO	
Clostridium tertium		30.10	0.50	MAN 99	1	99	
Bifidobacterium adolescentis		19.61	0.54		1	1	
Atopobium parvulum		17.26	0.46		99	1	
07375120	**Very good identification**						**07375120**
					COC	SPO	
Atopobium parvulum		97.64	0.80		99	1	
Clostridium tertium		0.97	0.45		1	99	
Clostridium butyricum		0.26	0.29		1	99	
07375121	**Good identification**						**07375121**
					COC	SPO	
Atopobium parvulum		95.73	0.46	URE 1	99	1	
Actinomyces naeslundii		2.83	0.43		1	1	
Clostridium tertium		0.95	0.11		1	99	
07375122	**Low discrimination**						**07375122**
					COC	SPO	
Lactobacillus acidophilus		44.06	0.66	XYL 1	1	1	
Atopobium parvulum		43.56	0.67		99	1	
Clostridium tertium		7.34	0.53		1	99	
07375130	**Low discrimination**						**07375130**
					COC	SPO	
Clostridium tertium		62.20	0.50	MAN 99	1	99	
Atopobium parvulum		35.66	0.46		99	1	
Actinomyces israelii		1.29	0.35		1	1	
07375132	**Low discrimination**						**07375132**
					COC	SPO	
Clostridium tertium		86.65	0.58	MAN 99	1	99	
Lactobacillus acidophilus		2.96	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		2.92	0.33		99	1	
07375140	**Low discrimination**						**07375140**
					COC	SPO	
Atopobium parvulum		76.18	0.59	SOR 1	99	1	
Actinomyces israelii		19.83	0.63		1	1	
Clostridium butyricum		3.56	0.29		1	99	
07375142	**Low discrimination**						**07375142**
					COC		
Actinomyces israelii		39.26	0.66		1		
Lactobacillus acidophilus		38.74	0.50		1		
Atopobium parvulum		17.74	0.46		99		
07375150	**Low discrimination**						**07375150**
					COC	SPO	
Actinomyces israelii		74.10	0.49	MLZ 13	1	1	
Atopobium parvulum		19.24	0.25		99	1	
Clostridium butyricum		4.69	0.07		1	99	
07375152	**Low discrimination**						**07375152**
Actinomyces israelii		62.04	0.52	MLZ 13			
Bifidobacterium adolescentis		26.35	0.39				
Lactobacillus acidophilus		4.14	0.16				
07375160	**Low discrimination**						**07375160**
					COC		
Atopobium parvulum		83.49	0.46	SOR 1	99		
Actinomyces israelii		7.86	0.42		1		
Clostridium perfringens		2.20	0.20		1		
07375162	**Low discrimination**						**07375162**
					COC		
Lactobacillus acidophilus		84.07	0.50	XYL 1	1		
Atopobium parvulum		6.79	0.33	SOR 11	99		
Actinomyces israelii		5.44	0.45		1		
07375212	**Low discrimination**						**07375212**
Clostridium clostridioforme		87.17	0.41	RAF 94			
Lactobacillus jensenii		8.04	0.21	ARA 94			
Actinomyces israelii		4.79	0.29				
07375232	**Good identification**						**07375232**
					GRA		
Clostridium clostridioforme		96.21	0.41	RAF 94	50		
Lactobacillus jensenii		2.22	0.10	ARA 94	99		
Parabacteroides distasonis		1.57	0.16		1		
07375242	**Low discrimination**						**07375242**
Actinomyces israelii		74.44	0.36	RHA 7			
Lactobacillus acidophilus		25.56	0.13				
07375300	**Good identification**						**07375300**
					COC	SPO	
Atopobium parvulum		93.38	0.59	RHA 1	99	1	
Actinomyces israelii		4.71	0.50		1	1	
Clostridium butyricum		1.55	0.21		1	99	
07375302	**Low discrimination**						**07375302**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		43.49	0.47	XYL 1	1		
Atopobium parvulum		32.70	0.46	RHA 7	99		
Actinomyces israelii		14.01	0.53		1		
07375312	**Low discrimination**						**07375312**
					SPO		
Clostridium clostridioforme		88.23	0.50	RAF 94	66		
Actinomyces israelii		5.79	0.39	ARA 94	1		
Clostridium tertium		1.60	0.15		99		
07375320	**Low discrimination**						**07375320**
					COC	SPO	
Atopobium parvulum		81.00	0.46	RHA 1	99	1	
Clostridium ramosum		12.69	0.30		1	92	
Actinomyces israelii		1.48	0.30		1	1	
07375322	**Low discrimination**						**07375322**
					SPO	GRA	
Lactobacillus acidophilus		38.92	0.47	XYL 1	1	99	
Parabacteroides distasonis		25.53	0.55	RHA 7	1	1	
Clostridium ramosum		18.34	0.40		92	50	
07375332	**Low discrimination**						**07375332**
					GRA	SPO	
Clostridium clostridioforme		63.99	0.50	RAF 94	50	66	
Parabacteroides distasonis		30.84	0.50	ARA 94	1	1	
Clostridium tertium		3.47	0.24		99	99	
07375340	**Low discrimination**						**07375340**
					COC	SPO	
Actinomyces israelii		62.92	0.43	RHA 7	1	1	
Atopobium parvulum		32.44	0.25		99	1	
Clostridium butyricum		4.64	0.03		1	99	
07375342	**Low discrimination**						**07375342**
					COC		
Actinomyces israelii		48.09	0.46	RHA 7	1		
Lactobacillus acidophilus		47.46	0.31		1		
Atopobium parvulum		2.92	0.12		99		
07375400	**Low discrimination**						**07375400**
					SPO		
Clostridium butyricum		54.49	0.40	ESL 88	99		
Actinomyces israelii		37.02	0.58	RAF 95	1		
Lactobacillus jensenii		5.76	0.32	LAC 96	1		
07375402	**Low discrimination**						**07375402**
					SPO		
Actinomyces israelii		60.82	0.61		1		
Clostridium butyricum		23.21	0.33		99		
Clostridium clostridioforme		7.13	0.36		66		
07375410	**Low discrimination**						**07375410**
					SPO		
Actinomyces israelii		52.09	0.44	MLZ 13	1		
Clostridium butyricum		27.01	0.18		99		
Lactobacillus jensenii		18.09	0.24		1		
07375412	**Low discrimination**						**07375412**
Clostridium clostridioforme		67.39	0.47	RAF 94			
Actinomyces israelii		18.85	0.47	RHA 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		8.32	0.34				
07375422	**Low discrimination**						**07375422**
					SPO		
Clostridium clostridioforme		47.89	0.36	RAF 94	66		
Actinomyces israelii		26.06	0.40	RHA 88	1		
Clostridium butyricum		13.55	0.15		99		
07375432	**Good identification**						**07375432**
Clostridium clostridioforme		96.91	0.47	RAF 94			
Actinomyces israelii		1.73	0.26	RHA 88			
Lactobacillus jensenii		1.05	0.10				
07375440	**Low discrimination**						**07375440**
					SPO		
Actinomyces israelii		75.24	0.51		1		
Clostridium butyricum		24.76	0.22		99		
07375442	**Acceptable identification**						**07375442**
					SPO		
Actinomyces israelii		90.36	0.54		1		
Clostridium butyricum		7.71	0.15		99		
Bifidobacterium adolescentis		1.55	0.17		1		
07375450	**Low discrimination**						**07375450**
					SPO		
Actinomyces israelii		89.61	0.37	MLZ 13	1		
Clostridium butyricum		10.39	0.00		99		
07375452	**Low discrimination**						**07375452**
Actinomyces israelii		79.16	0.40	MLZ 13			
Bifidobacterium adolescentis		11.11	0.19				
Clostridium clostridioforme		7.35	0.13				
07375500	**Low discrimination**						**07375500**
					COC	SPO	
Clostridium butyricum		75.61	0.63	ESL 88	1	99	
Atopobium parvulum		15.63	0.59	RAF 95	99	1	
Actinomyces israelii		8.56	0.68		1	1	
07375502	**Low discrimination**						**07375502**
					SPO		
Clostridium butyricum		45.99	0.56	ESL 88	99		
Bifidobacterium adolescentis		26.84	0.67	RAF 95	1		
Actinomyces israelii		20.08	0.71		1		
07375510	**Low discrimination**						**07375510**
					SPO		
Clostridium butyricum		69.78	0.41	ESL 88	99		
Actinomyces israelii		22.43	0.54	RAF 95	1		
Bifidobacterium adolescentis		3.72	0.34	MLZ 5	1		
07375512	**Low discrimination**						**07375512**
Bifidobacterium adolescentis		69.16	0.69	RAF 92			
Clostridium clostridioforme		18.94	0.56				
Actinomyces israelii		6.33	0.57				
07375520	**Low discrimination**						**07375520**
					COC	SPO	
Clostridium butyricum		66.22	0.45	ESL 88	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		27.78	0.46	RAF 95	99	1	
Actinomyces israelii		5.50	0.48	NAG 8	1	1	
07375522	**Low discrimination**						**07375522**
					SPO		
Clostridium butyricum		39.91	0.38	ESL 88	99		
Clostridium clostridioforme		19.68	0.45	RAF 95	66		
Actinomyces israelii		12.80	0.51	NAG 8	1		
07375532	**Low discrimination**						**07375532**
					GRA		
Clostridium clostridioforme		85.47	0.56	RAF 94	50		
Parabacteroides distasonis		3.94	0.39	RHA 88	1		
Bifidobacterium adolescentis		3.15	0.34		99		
07375540	**Low discrimination**						**07375540**
					COC	SPO	
Clostridium butyricum		65.21	0.45	ESL 88	1	99	
Actinomyces israelii		33.02	0.61	RAF 95	1	1	
Atopobium parvulum		1.57	0.25	SOR 8	99	1	
07375542	**Low discrimination**						**07375542**
					SPO		
Actinomyces israelii		51.10	0.64		1		
Clostridium butyricum		26.16	0.38		99		
Bifidobacterium adolescentis		21.70	0.52		1		
07375550	**Low discrimination**						**07375550**
					SPO		
Actinomyces israelii		56.59	0.47	MLZ 13	1		
Clostridium butyricum		39.37	0.23		99		
Bifidobacterium adolescentis		2.98	0.19		1		
07375552	**Low discrimination**						**07375552**
					SPO		
Bifidobacterium adolescentis		73.41	0.53	RAF 92	1		
Actinomyces israelii		21.13	0.50	SOR 11	1		
Clostridium butyricum		3.81	0.16		99		
07375562	**Low discrimination**						**07375562**
					SPO		
Actinomyces israelii		50.11	0.44	NAG 6	1		
Clostridium butyricum		34.95	0.20		99		
Lactobacillus acidophilus		9.57	0.16		1		
07375600	**Low discrimination**						**07375600**
					SPO		
Actinomyces israelii		56.24	0.39	RHA 7	1		
Clostridium butyricum		34.02	0.14		99		
Clostridium clostridioforme		9.74	0.16		66		
07375602	**Acceptable identification**						**07375602**
					SPO		
Clostridium clostridioforme		90.81	0.50	RAF 94	66		
Actinomyces israelii		7.95	0.42		1		
Clostridium butyricum		1.25	0.07		99		
07375612	**Very good identification**						**07375612**
Clostridium clostridioforme		99.71	0.62	RAF 94			
Actinomyces israelii		0.29	0.28				
07375622	**Good identification**						**07375622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Clostridium clostridioforme		98.91	0.50	RAF 94	50		
Actinomyces israelii		0.55	0.21		99		
Parabacteroides distasonis		0.53	0.17		1		
07375632	**Very good identification**						**07375632**
Clostridium clostridioforme		99.94	0.62	RAF 94			
Parabacteroides distasonis		0.06	0.13				
07375700	**Low discrimination**						**07375700**
					COC	SPO	
Clostridium butyricum		72.31	0.38	ESL 88	1	99	
Actinomyces israelii		19.92	0.49	RAF 95	1	1	
Atopobium parvulum		4.88	0.25	RHA 3	99	1	
07375702	**Low discrimination**						**07375702**
					SPO		
Clostridium clostridioforme		81.28	0.59	RAF 94	66		
Actinomyces israelii		8.50	0.52		1		
Clostridium butyricum		8.00	0.30		99		
07375710	**Low discrimination**						**07375710**
					SPO		
Clostridium clostridioforme		65.98	0.36	RAF 94	66		
Clostridium butyricum		19.09	0.16	BGL 99	99		
Actinomyces israelii		14.93	0.35		1		
07375712	**Very good identification**						**07375712**
Clostridium clostridioforme		99.03	0.71	RAF 94			
Bifidobacterium adolescentis		0.50	0.34				
Actinomyces israelii		0.34	0.38				
07375713	**Good identification**						**07375713**
Clostridium clostridioforme		100.0	0.36	RAF 94			
07375722	**Low discrimination**						**07375722**
Clostridium clostridioforme		82.53	0.59	RAF 94			
Parabacteroides distasonis		13.17	0.51				
Bacteroides ovatus		2.43	0.23				
07375730	**Acceptable identification**						**07375730**
					GRA	SPO	
Clostridium clostridioforme		94.48	0.36	RAF 94	50	66	
Clostridium butyricum		2.38	-.03	BGL 99	99	99	
Parabacteroides distasonis		1.78	0.13		1	1	
07375732	**Very good identification**						**07375732**
Clostridium clostridioforme		97.70	0.71	RAF 94			
Parabacteroides distasonis		1.84	0.47				
Bacteroides ovatus		0.42	0.20				
07375733	**Good identification**						**07375733**
Clostridium clostridioforme		98.15	0.36	RAF 94			
Parabacteroides distasonis		1.85	0.13	URE 1			
07375740	**Low discrimination**						**07375740**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		55.20	0.42	RHA 7	1		
Clostridium butyricum		44.80	0.19		99		
07375742	**Low discrimination**						**07375742**
					SPO		
Actinomyces israelii		65.97	0.45	RHA 7	1		
Clostridium clostridioforme		16.39	0.25		66		
Clostridium butyricum		13.88	0.12		99		
07375752	**Low discrimination**						**07375752**
Clostridium clostridioforme		83.78	0.36	RAF 94			
Actinomyces israelii		11.06	0.31	SOR 1			
Bifidobacterium adolescentis		5.16	0.19				
07375772	**Good identification**						**07375772**
Clostridium clostridioforme		96.47	0.36	RAF 94			
Parabacteroides distasonis		1.82	0.13	SOR 1			
Bacteroides ovatus		1.72	-.04				
07376000	**Low discrimination**						**07376000**
					SPO		
Actinomyces israelii		73.32	0.38	GAL 99	1		
Clostridium innocuum		22.22	0.14		99		
Clostridium tertium		4.46	0.03		99		
07376002	**Low discrimination**						**07376002**
					SPO		
Actinomyces israelii		75.55	0.41	GAL 99	1		
Clostridium innocuum		15.26	0.14		99		
Clostridium tertium		9.19	0.11		99		
07376100	**Low discrimination**						**07376100**
					COC	SPO	
Clostridium tertium		54.07	0.37	GAL 99	1	99	
Actinomyces israelii		35.90	0.48		1	1	
Atopobium parvulum		9.46	0.25		99	1	
07376102	**Low discrimination**						**07376102**
					COC	SPO	
Clostridium tertium		73.55	0.45	GAL 99	1	99	
Actinomyces israelii		24.42	0.51		1	1	
Atopobium parvulum		0.76	0.12		99	1	
07376110	**Acceptable identification**						**07376110**
					SPO		
Clostridium tertium		94.71	0.41	GAL 99	99		
Actinomyces israelii		5.29	0.34		1		
07376112	**Good identification**						**07376112**
					SPO		
Clostridium tertium		96.78	0.50	GAL 99	99		
Actinomyces israelii		2.70	0.37		1		
Bifidobacterium adolescentis		0.52	0.19		1		
07376120	**Good identification**						**07376120**
					COC	SPO	
Clostridium tertium		96.96	0.45	GAL 99	1	99	
Actinomyces israelii		1.37	0.28		1	1	
Atopobium parvulum		1.00	0.12		99	1	
07376122	**Good identification**						**07376122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium tertium		98.09	0.53	GAL 99	99	
Clostridium ramosum		0.90	0.17		92	
Actinomyces israelii		0.69	0.31		1	
07376130	**Good identification**					**07376130**
					SPO	
Clostridium tertium		99.88	0.50	GAL 99	99	
Actinomyces israelii		0.12	0.14		1	
07376132	**Very good identification**					**07376132**
					SPO	
Clostridium tertium		99.94	0.58	GAL 99	99	
Actinomyces israelii		0.06	0.17		1	
07376140	**Good identification**					**07376140**
					SPO	
Actinomyces israelii		96.24	0.41	GAL 99	1	
Clostridium tertium		3.76	0.03		99	
07376142	**Acceptable identification**					**07376142**
					SPO	
Actinomyces israelii		92.74	0.44	GAL 99	1	
Clostridium tertium		7.26	0.11		99	
07376400	**Acceptable identification**					**07376400**
					SPO	
Actinomyces israelii		93.87	0.37	GAL 99	1	
Clostridium innocuum		6.13	0.01		99	
07376402	**Good identification**					**07376402**
					SPO	
Actinomyces israelii		95.83	0.40	GAL 99	1	
Clostridium innocuum		4.18	0.01		99	
07376500	**Low discrimination**					**07376500**
					SPO	
Actinomyces israelii		83.91	0.47	GAL 99	1	
Clostridium butyricum		14.53	0.13		99	
Clostridium tertium		1.56	0.03		99	
07376502	**Low discrimination**					**07376502**
					SPO	
Actinomyces israelii		78.43	0.50	GAL 99	1	
Bifidobacterium adolescentis		15.13	0.31		1	
Clostridium butyricum		3.52	0.06		99	
07376540	**Good identification**					**07376540**
					SPO	
Actinomyces israelii		96.27	0.40	GAL 99	1	
Clostridium butyricum		3.73	-.05		99	
07376542	**Acceptable identification**					**07376542**
Actinomyces israelii		94.22	0.43	GAL 99		
Bifidobacterium adolescentis		5.78	0.16			
07377000	**Low discrimination**					**07377000**
					SPO	
Actinomyces israelii		86.41	0.72		1	
Clostridium tertium		5.26	0.37		99	
Clostridium innocuum		5.03	0.36		99	
07377001	**Low discrimination**					**07377001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		67.11	0.38	URE 1	1	
Actinomyces naeslundii		24.91	0.40		1	
Clostridium tertium		4.08	0.03		99	
07377002	**Low discrimination**					**07377002**
					SPO	
Actinomyces israelii		84.36	0.75		1	
Clostridium tertium		10.27	0.45		99	
Clostridium innocuum		3.27	0.36		99	
07377003	**Low discrimination**					**07377003**
					SPO	
Actinomyces israelii		71.03	0.41	URE 1	1	
Actinomyces naeslundii		17.57	0.40		1	
Clostridium tertium		8.64	0.11		99	
07377010	**Low discrimination**					**07377010**
					SPO	
Actinomyces israelii		55.86	0.58	MLZ 13	1	
Clostridium tertium		40.44	0.41		99	
Lactobacillus jensenii		2.80	0.24		1	
07377012	**Low discrimination**					**07377012**
					SPO	
Clostridium tertium		58.18	0.50	LAC 99	99	
Actinomyces israelii		40.18	0.61		1	
Lactobacillus jensenii		0.90	0.21		1	
07377020	**Low discrimination**					**07377020**
					SPO	
Clostridium tertium		70.54	0.45	LAC 99	99	
Actinomyces israelii		24.66	0.52		1	
Lactobacillus jensenii		2.17	0.21		1	
07377021	**Low discrimination**					**07377021**
					SPO	
Actinomyces naeslundii		60.10	0.40	RAF 92	1	
Clostridium tertium		29.56	0.11	XYL 1	99	
Actinomyces israelii		10.34	0.18		1	
07377022	**Low discrimination**					**07377022**
					SPO	
Clostridium tertium		81.50	0.53	LAC 99	99	
Actinomyces israelii		14.25	0.55		1	
Lactobacillus acidophilus		2.24	0.26		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07377030	**Good identification**						**07377030**
					SPO		
Clostridium tertium		96.60	0.50	LAC 99	99		
Actinomyces israelii		2.84	0.38		1		
Lactobacillus jensenii		0.56	0.13		1		
07377032	**Very good identification**						**07377032**
					SPO		
Clostridium tertium		98.43	0.58	LAC 99	99		
Actinomyces israelii		1.45	0.41		1		
Lactobacillus jensenii		0.13	0.10		1		
07377040	**Very good identification**						**07377040**
					SPO		
Actinomyces israelii		99.58	0.65		1		
Clostridium tertium		0.16	0.03		99		
Clostridium innocuum		0.15	0.02		99		
07377042	**Very good identification**						**07377042**
					SPO		
Actinomyces israelii		99.20	0.68		1		
Lactobacillus acidophilus		0.32	0.10		1		
Clostridium tertium		0.31	0.11		99		
07377050	**Good identification**						**07377050**
					SPO		
Actinomyces israelii		98.15	0.51	MLZ 13	1		
Clostridium tertium		1.85	0.07		99		
07377052	**Good identification**						**07377052**
					SPO		
Actinomyces israelii		96.38	0.54	MLZ 13	1		
Clostridium tertium		3.63	0.15		99		
07377060	**Acceptable identification**						**07377060**
					SPO		
Actinomyces israelii		91.57	0.45	NAG 6	1		
Clostridium tertium		6.80	0.11		99		
Actinomyces naeslundii		1.63	0.24		1		
07377062	**Low discrimination**						**07377062**
					SPO		
Actinomyces israelii		82.62	0.48	NAG 6	1		
Clostridium tertium		12.28	0.19		99		
Lactobacillus acidophilus		4.13	0.10		1		
07377100	**Low discrimination**						**07377100**
					COC	SPO	
Clostridium tertium		54.01	0.71		1	99	
Actinomyces israelii		35.86	0.83		1	1	
Atopobium parvulum		9.45	0.59		99	1	
07377101	**Low discrimination**						**07377101**
					SPO		
Clostridium tertium		49.39	0.37	URE 1	99		
Actinomyces israelii		32.79	0.48		1		
Actinomyces naeslundii		8.66	0.48		1		
07377102	**Low discrimination**						**07377102**
					COC	SPO	
Clostridium tertium		73.43	0.79		1	99	
Actinomyces israelii		24.38	0.86		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		0.76	0.46		99	1	
07377103	**Low discrimination**						**07377103**
					SPO		
Clostridium tertium		70.52	0.45	URE 1	99		
Actinomyces israelii		23.41	0.51		1		
Actinomyces naeslundii		4.12	0.48		1		
07377110	**Good identification**						**07377110**
					COC	SPO	
Clostridium tertium		94.58	0.76		1	99	
Actinomyces israelii		5.28	0.68		1	1	
Atopobium parvulum		0.09	0.25		99	1	
07377111	**Acceptable identification**						**07377111**
					SPO		
Clostridium tertium		94.25	0.41	URE 1	99		
Actinomyces israelii		5.26	0.34		1		
Actinomyces naeslundii		0.49	0.26		1		
07377112	**Very good identification**						**07377112**
					SPO		
Clostridium tertium		96.75	0.84		99		
Actinomyces israelii		2.70	0.71		1		
Bifidobacterium adolescentis		0.52	0.53		1		
07377113	**Good identification**						**07377113**
					SPO		
Clostridium tertium		96.62	0.50	URE 1	99		
Actinomyces israelii		2.70	0.37		1		
Bifidobacterium adolescentis		0.52	0.19		1		
07377120	**Very good identification**						**07377120**
					COC	SPO	
Clostridium tertium		96.88	0.79		1	99	
Actinomyces israelii		1.37	0.62		1	1	
Atopobium parvulum		1.00	0.46		99	1	
07377121	**Acceptable identification**						**07377121**
					SPO		
Clostridium tertium		91.76	0.45	URE 1	99		
Actinomyces naeslundii		5.36	0.48		1		
Actinomyces israelii		1.30	0.28		1		
07377122	**Very good identification**						**07377122**
					SPO		
Clostridium tertium		98.00	0.88		99		
Clostridium ramosum		0.90	0.51		92		
Actinomyces israelii		0.69	0.65		1		
07377123	**Good identification**						**07377123**
					SPO		
Clostridium tertium		96.25	0.53	URE 1	99		
Actinomyces naeslundii		1.88	0.48		1		
Clostridium ramosum		0.89	0.17		92		
07377130	**Very good identification**						**07377130**
					COC	SPO	
Clostridium tertium		99.87	0.84		1	99	
Actinomyces israelii		0.12	0.48		1	1	
Atopobium parvulum		0.01	0.12		99	1	
07377131	**Good identification**						**07377131**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				SPO		
Clostridium tertium	99.71	0.50	URE 1	99		
Actinomyces naeslundii	0.17	0.26		1		
Actinomyces israelii	0.12	0.14		1		
07377132 Excellent identification						**07377132**
				SPO		
Clostridium tertium	99.93	0.92		99		
Actinomyces israelii	0.06	0.51		1		
Bifidobacterium dentium	0.01	0.10		1		
07377133 Very good identification						**07377133**
				SPO		
Clostridium tertium	99.88	0.58	URE 1	99		
Actinomyces israelii	0.06	0.17		1		
Actinomyces naeslundii	0.06	0.26		1		
07377140 Very good identification						**07377140**
				COC	SPO	
Actinomyces israelii	95.21	0.76		1	1	
Clostridium tertium	3.73	0.37		1	99	
Atopobium parvulum	0.65	0.25		99	1	
07377141 Low discrimination						**07377141**
				SPO		
Actinomyces israelii	86.29	0.41	URE 1	1		
Actinomyces naeslundii	10.34	0.35		1		
Clostridium tertium	3.38	0.03		99		
07377142 Good identification						**07377142**
				SPO		
Actinomyces israelii	91.15	0.79		1		
Clostridium tertium	7.13	0.45		99		
Lactobacillus acidophilus	0.84	0.28		1		
07377143 Low discrimination						**07377143**
				SPO		
Actinomyces israelii	86.35	0.44	URE 1	1		
Actinomyces naeslundii	6.90	0.35		1		
Clostridium tertium	6.76	0.11		99		
07377150 Low discrimination						**07377150**
				COC	SPO	
Actinomyces israelii	68.11	0.61	MLZ 13	1	1	
Clostridium tertium	31.70	0.41		1	99	
Blautia producta	0.18	-.03		99	1	
07377152 Low discrimination						**07377152**
				SPO		
Actinomyces israelii	50.20	0.64	MLZ 13	1		
Clostridium tertium	46.73	0.50		99		
Bifidobacterium adolescentis	3.08	0.37		1		
07377160 Low discrimination						**07377160**
				COC	SPO	
Clostridium tertium	63.67	0.45	SOR 1	1	99	
Actinomyces israelii	34.63	0.55		1	1	
Atopobium parvulum	0.66	0.12		99	1	
07377162 Low discrimination						**07377162**
				SPO		
Clostridium tertium	75.57	0.53	SOR 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		20.55	0.58		1		
Lactobacillus acidophilus		2.95	0.28		1		
07377170	**Good identification**						**07377170**
					SPO		
Clostridium tertium		95.63	0.50	SOR 1	99		
Actinomyces israelii		4.37	0.41		1		
07377172	**Very good identification**						**07377172**
					SPO		
Clostridium tertium		97.77	0.58	SOR 1	99		
Actinomyces israelii		2.24	0.44		1		
07377200	**Good identification**						**07377200**
					SPO		
Actinomyces israelii		98.43	0.53	RHA 7	1		
Clostridium tertium		0.80	0.03		99		
Clostridium innocuum		0.77	0.02		99		
07377202	**Good identification**						**07377202**
					SPO		
Actinomyces israelii		96.94	0.56	RHA 7	1		
Clostridium tertium		1.58	0.11		99		
Lactobacillus acidophilus		0.97	0.07		1		
07377210	**Acceptable identification**						**07377210**
					SPO		
Actinomyces israelii		91.14	0.39	RHA 7	1		
Clostridium tertium		8.86	0.07	MLZ 13	99		
07377212	**Low discrimination**						**07377212**
					SPO		
Actinomyces israelii		81.52	0.42	RHA 7	1		
Clostridium tertium		15.84	0.15	MLZ 13	99		
Clostridium clostridioforme		2.64	0.07		66		
07377222	**Low discrimination**						**07377222**
					SPO		
Actinomyces israelii		35.08	0.36	RHA 7	1		
Clostridium ramosum		32.48	0.19	NAG 6	92		
Clostridium tertium		26.93	0.19		99		
07377240	**Good identification**						**07377240**
Actinomyces israelii		100.0	0.46	RHA 7			
07377242	**Good identification**						**07377242**
Actinomyces israelii		100.0	0.49	RHA 7			
07377300	**Low discrimination**						**07377300**
					COC	SPO	
Actinomyces israelii		80.03	0.63	RHA 7	1	1	
Clostridium tertium		16.18	0.37		1	99	
Atopobium parvulum		2.83	0.25		99	1	
07377302	**Low discrimination**						**07377302**
					SPO		
Actinomyces israelii		68.60	0.66	RHA 7	1		
Clostridium tertium		27.73	0.45		99		
Lactobacillus acidophilus		1.98	0.25		1		
07377310	**Low discrimination**						**07377310**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		70.63	0.41	RHA 1	99		
Actinomyces israelii		29.37	0.49		1		
07377312	**Low discrimination**						**07377312**
					SPO		
Clostridium tertium		82.04	0.50	RHA 1	99		
Actinomyces israelii		17.06	0.52		1		
Clostridium clostridioforme		0.46	0.16		66		
07377320	**Low discrimination**						**07377320**
					SPO		
Clostridium tertium		49.50	0.45	RHA 1	99		
Clostridium ramosum		44.78	0.43		92		
Actinomyces israelii		5.21	0.43		1		
07377322	**Low discrimination**						**07377322**
					SPO		
Clostridium ramosum		52.74	0.53	XYL 1	92		
Clostridium tertium		43.72	0.53		99		
Actinomyces israelii		2.30	0.46		1		
07377330	**Good identification**						**07377330**
					SPO		
Clostridium tertium		98.62	0.50	RHA 1	99		
Actinomyces israelii		0.87	0.29		1		
Clostridium ramosum		0.51	0.09		92		
07377332	**Very good identification**						**07377332**
					SPO		
Clostridium tertium		98.61	0.58	RHA 1	99		
Clostridium ramosum		0.68	0.19		92		
Actinomyces israelii		0.44	0.32		1		
07377340	**Very good identification**						**07377340**
					SPO		
Actinomyces israelii		99.48	0.56	RHA 7	1		
Clostridium tertium		0.52	0.03		99		
07377342	**Very good identification**						**07377342**
					SPO		
Actinomyces israelii		98.07	0.59	RHA 7	1		
Clostridium tertium		1.03	0.11		99		
Lactobacillus acidophilus		0.90	0.09		1		
07377350	**Acceptable identification**						**07377350**
					COC	SPO	
Actinomyces israelii		91.04	0.42	RHA 7	1	1	
Clostridium tertium		5.69	0.07	MLZ 13	1	99	
Blautia producta		3.28	-.03		99	1	
07377352	**Low discrimination**						**07377352**
					SPO		
Actinomyces israelii		88.90	0.45	RHA 7	1		
Clostridium tertium		11.11	0.15	MLZ 13	99		
07377360	**Low discrimination**						**07377360**
					SPO		
Actinomyces israelii		68.03	0.36	RHA 7	1		
Clostridium tertium		16.79	0.11	NAG 6	99		
Clostridium ramosum		15.19	0.09		92		
07377362	**Low discrimination**						**07377362**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		44.80	0.39	RHA 7	1	
Clostridium ramosum		26.67	0.19	NAG 6	92	
Clostridium tertium		22.11	0.19		99	
07377400	**Very good identification**					**07377400**
					SPO	
Actinomyces israelii		95.96	0.71		1	
Clostridium butyricum		2.77	0.23		99	
Clostridium innocuum		1.20	0.23		99	
07377401	**Good identification**					**07377401**
Actinomyces israelii		100.0	0.37	URE 1		
07377402	**Very good identification**					**07377402**
					SPO	
Actinomyces israelii		97.55	0.74		1	
Clostridium innocuum		0.82	0.23		99	
Bifidobacterium adolescentis		0.76	0.31		1	
07377403	**Good identification**					**07377403**
Actinomyces israelii		100.0	0.40	URE 1		
07377410	**Very good identification**					**07377410**
					SPO	
Actinomyces israelii		97.79	0.57	MLZ 13	1	
Clostridium butyricum		0.99	0.02		99	
Clostridium tertium		0.87	0.07		99	
07377412	**Low discrimination**					**07377412**
Actinomyces israelii		90.00	0.60	MLZ 13		
Bifidobacterium adolescentis		5.74	0.33			
Bifidobacterium dentium		1.63	0.06			
07377420	**Acceptable identification**					**07377420**
					SPO	
Actinomyces israelii		93.06	0.50	NAG 6	1	
Clostridium butyricum		3.66	0.05		99	
Clostridium tertium		3.29	0.11		99	
07377422	**Acceptable identification**					**07377422**
					SPO	
Actinomyces israelii		92.52	0.53	NAG 6	1	
Clostridium tertium		6.53	0.19		99	
Clostridium butyricum		0.94	-.02		99	
07377430	**Low discrimination**					**07377430**
					SPO	
Actinomyces israelii		66.89	0.36	MLZ 13	1	
Clostridium tertium		28.10	0.15	NAG 6	99	
Bifidobacterium dentium		5.01	-.07		1	
07377432	**Low discrimination**					**07377432**
					SPO	
Actinomyces israelii		40.89	0.39	MLZ 13	1	
Clostridium tertium		34.36	0.24	NAG 6	99	
Bifidobacterium dentium		20.66	0.10		1	
07377440	**Very good identification**					**07377440**
					SPO	
Actinomyces israelii		99.36	0.64		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		0.64	0.05		99	
07377442	**Very good identification**					**07377442**
					SPO	
Actinomyces israelii		99.59	0.67		1	
Bifidobacterium adolescentis		0.25	0.16		1	
Clostridium butyricum		0.17	-.02		99	
07377450	**Good identification**					**07377450**
					COC	
Actinomyces israelii		98.69	0.50	MLZ 13	1	
Blautia producta		1.31	-.03		99	
07377452	**Good identification**					**07377452**
Actinomyces israelii		98.01	0.53	MLZ 13		
Bifidobacterium adolescentis		1.99	0.17			
07377460	**Good identification**					**07377460**
Actinomyces israelii		100.0	0.43	NAG 6		
07377462	**Good identification**					**07377462**
Actinomyces israelii		100.0	0.46	NAG 6		
07377500	**Low discrimination**					**07377500**
					SPO	
Actinomyces israelii		83.48	0.81		1	
Clostridium butyricum		14.46	0.47		99	
Clostridium tertium		1.55	0.37		99	
07377501	**Low discrimination**					**07377501**
					SPO	
Actinomyces israelii		83.91	0.47	URE 1	1	
Clostridium butyricum		14.53	0.13		99	
Clostridium tertium		1.56	0.03		99	
07377502	**Low discrimination**					**07377502**
					SPO	
Actinomyces israelii		78.33	0.84		1	
Bifidobacterium adolescentis		15.11	0.66		1	
Clostridium butyricum		3.52	0.40		99	
07377503	**Low discrimination**					**07377503**
					SPO	
Actinomyces israelii		78.43	0.50	URE 1	1	
Bifidobacterium adolescentis		15.13	0.31		1	
Clostridium butyricum		3.52	0.06		99	
07377510	**Low discrimination**					**07377510**
					SPO	
Actinomyces israelii		72.78	0.67	MLZ 13	1	
Clostridium tertium		16.10	0.41		99	
Bifidobacterium dentium		4.85	0.23		1	
07377512	**Low discrimination**					**07377512**
Bifidobacterium adolescentis		45.21	0.67	RAF 92		
Actinomyces israelii		28.65	0.70			
Bifidobacterium dentium		12.86	0.40			
07377520	**Low discrimination**					**07377520**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		47.00	0.61	NAG 6	1		
Clostridium tertium		41.07	0.45		99		
Clostridium butyricum		11.09	0.29		99		
07377522	**Low discrimination**						**07377522**
					SPO		
Clostridium tertium		60.63	0.53	ARA 1	99		
Actinomyces israelii		34.69	0.64		1		
Clostridium butyricum		2.12	0.22		99		
07377530	**Low discrimination**						**07377530**
					SPO		
Clostridium tertium		77.97	0.50	ARA 1	99		
Bifidobacterium dentium		13.91	0.27		1		
Actinomyces israelii		7.50	0.47		1		
07377532	**Low discrimination**						**07377532**
					SPO		
Clostridium tertium		60.00	0.58	ARA 1	99		
Bifidobacterium dentium		36.09	0.44		1		
Actinomyces israelii		2.89	0.50		1		
07377540	**Good identification**						**07377540**
					COC	SPO	
Actinomyces israelii		94.66	0.74		1	1	
Clostridium butyricum		3.67	0.29		1	99	
Blautia producta		1.55	0.24		99	1	
07377541	**Good identification**						**07377541**
					SPO		
Actinomyces israelii		96.27	0.40	URE 1	1		
Clostridium butyricum		3.73	-.05		99		
07377542	**Good identification**						**07377542**
					SPO		
Actinomyces israelii		93.15	0.77		1		
Bifidobacterium adolescentis		5.71	0.50		1		
Clostridium butyricum		0.94	0.22		99		
07377543	**Acceptable identification**						**07377543**
Actinomyces israelii		94.22	0.43	URE 1			
Bifidobacterium adolescentis		5.78	0.16				
07377550	**Low discrimination**						**07377550**
					COC	SPO	
Actinomyces israelii		73.82	0.60	MLZ 13	1	1	
Blautia producta		24.19	0.32		99	1	
Clostridium butyricum		1.01	0.07		1	99	
07377552	**Low discrimination**						**07377552**
					COC		
Actinomyces israelii		64.41	0.63	MLZ 13	1		
Bifidobacterium adolescentis		32.31	0.52		1		
Blautia producta		1.56	0.15		99		
07377560	**Acceptable identification**						**07377560**
					COC	SPO	
Actinomyces israelii		90.59	0.54	NAG 6	1	1	
Clostridium butyricum		4.78	0.11		1	99	
Blautia producta		2.57	0.07		99	1	
07377562	**Good identification**						**07377562**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		93.57	0.57	NAG 6	1		
Clostridium tertium		4.25	0.19		99		
Clostridium butyricum		1.28	0.04		99		
07377570	**Low discrimination**						**07377570**
					COC	SPO	
Actinomyces israelii		52.95	0.39	MLZ 13	1	1	
Blautia producta		30.20	0.15	NAG 6	99	1	
Clostridium tertium		14.30	0.15		1	99	
07377572	**Low discrimination**						**07377572**
					SPO		
Actinomyces israelii		50.34	0.42	MLZ 13	1		
Clostridium tertium		27.19	0.24	NAG 6	99		
Bifidobacterium dentium		16.35	0.10		1		
07377600	**Good identification**						**07377600**
					SPO		
Actinomyces israelii		98.83	0.52	RHA 7	1		
Clostridium butyricum		1.17	-.03		99		
07377602	**Good identification**						**07377602**
Actinomyces israelii		98.00	0.55	RHA 7			
Clostridium clostridioforme		2.00	0.16				
07377610	**Good identification**						**07377610**
Actinomyces israelii		100.0	0.37	RHA 7			
07377612	**Low discrimination**						**07377612**
Actinomyces israelii		61.69	0.40	RHA 7			
Clostridium clostridioforme		38.31	0.27	MLZ 13			
07377640	**Good identification**						**07377640**
Actinomyces israelii		100.0	0.45	RHA 7			
07377642	**Good identification**						**07377642**
Actinomyces israelii		100.0	0.48	RHA 7			
07377700	**Good identification**						**07377700**
					SPO		
Actinomyces israelii		93.14	0.62	RHA 7	1		
Clostridium butyricum		6.63	0.21		99		
Clostridium tertium		0.23	0.03		99		
07377702	**Good identification**						**07377702**
					SPO		
Actinomyces israelii		93.78	0.65	RHA 7	1		
Bifidobacterium adolescentis		2.43	0.31		1		
Clostridium butyricum		1.73	0.14		99		
07377710	**Acceptable identification**						**07377710**
					SPO		
Actinomyces israelii		93.30	0.48	RHA 7	1		
Clostridium tertium		2.77	0.07	MLZ 13	99		
Clostridium butyricum		2.34	-.01		99		
07377712	**Low discrimination**						**07377712**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		53.83	0.51	RHA 7				
Clostridium clostridioforme		27.98	0.36	MLZ 13				
Bifidobacterium adolescentis		11.40	0.33					
07377720	**Low discrimination**							**07377720**
					SPO			
Actinomyces israelii		75.74	0.41	RHA 7	1			
Clostridium tertium		8.88	0.11	NAG 6	99			
Clostridium ramosum		8.04	0.09		92			
07377722	**Low discrimination**							**07377722**
					SPO			
Actinomyces israelii		47.35	0.44	RHA 7	1			
Clostridium ramosum		13.40	0.19	NAG 6	92			
Clostridium clostridioforme		12.65	0.25		66			
07377732	**Low discrimination**							**07377732**
					SPO			
Clostridium clostridioforme		54.56	0.36	RAF 94	66			
Clostridium tertium		18.70	0.24	MAN 1	99			
Bifidobacterium dentium		11.25	0.10		1			
07377740	**Low discrimination**							**07377740**
					COC	SPO		
Actinomyces israelii		81.12	0.55	RHA 7	1	1		
Blautia producta		17.59	0.24		99	1		
Clostridium butyricum		1.29	0.03		1	99		
07377742	**Very good identification**							**07377742**
					COC			
Actinomyces israelii		97.24	0.58	RHA 7	1			
Blautia producta		1.56	0.07		99			
Bifidobacterium adolescentis		0.80	0.16		1			
07377752	**Low discrimination**							**07377752**
					COC			
Actinomyces israelii		71.95	0.44	RHA 7	1			
Blautia producta		23.20	0.15	MLZ 13	99			
Bifidobacterium adolescentis		4.84	0.17		1			
07377762	**Good identification**							**07377762**
Actinomyces israelii		100.0	0.37	RHA 7				
07400000	**Low discrimination**							**07400000**
					COC	GRA	CAT	
Gemella morbillorum		45.59	0.52	RAF 1	99	99	1	
Propionibacterium granulosum		41.10	0.58		1	99	90	
Fusobacterium mortiferum		7.79	0.49		1	1	5	
07400002	**Low discrimination**							**07400002**
					GRA	CAT		
Propionibacterium granulosum		76.33	0.47	FRU 99	99	90		
Atopobium minutum		12.14	0.32		99	1		
Fusobacterium mortiferum		6.43	0.32		1	5		
07400010	**Acceptable identification**							**07400010**
					COC	GRA	CAT	
Propionibacterium granulosum		91.22	0.43	FRU 99	1	99	90	
Gemella morbillorum		7.50	0.17	MLZ 12	99	99	1	
Fusobacterium mortiferum		1.28	0.14		1	1	5	
07400020	**Low discrimination**							**07400020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	CAT		
Propionibacterium granulosum		60.72	0.47	FRU 99	1	90		
Gemella morbillorum		29.94	0.35		99	1		
Atopobium minutum		4.83	0.26		1	1		
07400022	**Low discrimination**							**07400022**
					COC	CAT		
Propionibacterium granulosum		81.17	0.37	FRU 99	1	90		
Atopobium minutum		17.22	0.23		1	1		
Gemella morbillorum		1.62	0.01		99	1		
07400100	**Good identification**							**07400100**
					COC	GRA	CAT	
Fusobacterium mortiferum		95.04	0.55	FRU 99	1	1	5	
Gemella morbillorum		2.41	0.17		99	99	1	
Propionibacterium granulosum		2.17	0.24		1	99	90	
07400102	**Low discrimination**							**07400102**
					GRA	CAT		
Fusobacterium mortiferum		83.29	0.39	FRU 99	1	5		
Actinomyces viscosus		9.44	0.10	BGL 10	99	76		
Propionibacterium granulosum		4.28	0.13		99	90		
07400400	**Acceptable identification**							**07400400**
					COC	GRA	CAT	
Gemella morbillorum		91.12	0.35	RAF 1	99	99	1	
Propionibacterium granulosum		7.47	0.24	ARA 10	1	99	90	
Fusobacterium mortiferum		1.42	0.14		1	1	5	
07401000	**Low discrimination**							**07401000**
					COC	GRA	CAT	
Propionibacterium granulosum		49.50	0.58	FRU 99	1	99	90	
Gemella morbillorum		29.57	0.47		99	99	1	
Fusobacterium mortiferum		14.08	0.52		1	1	5	
07401002	**Low discrimination**							**07401002**
					GRA	CAT		
Propionibacterium granulosum		60.72	0.47	FRU 99	99	90		
Lactobacillus fermentum		26.02	0.46		99	1		
Fusobacterium mortiferum		7.68	0.35		1	5		
07401010	**Acceptable identification**							**07401010**
					COC	GRA	CAT	
Propionibacterium granulosum		93.87	0.43	FRU 99	1	99	90	
Gemella morbillorum		4.15	0.13	MLZ 12	99	99	1	
Fusobacterium mortiferum		1.98	0.17		1	1	5	
07401020	**Low discrimination**							**07401020**
					COC	CAT		
Propionibacterium granulosum		64.68	0.47	FRU 99	1	90		
Gemella morbillorum		17.17	0.31		99	1		
Lactobacillus fermentum		9.24	0.38		1	1		
07401022	**Low discrimination**							**07401022**
					CAT			
Propionibacterium granulosum		57.43	0.37	FRU 99	90			
Lactobacillus fermentum		32.81	0.38		1			
Atopobium minutum		4.06	0.15		1			
07401100	**Low discrimination**							**07401100**
					GRA			
Fusobacterium mortiferum		74.87	0.58	FRU 99	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		19.71	0.50		99		
Prevotella melaninogenica		2.61	0.15		1		
07401102	**Low discrimination**						**07401102**
					GRA		
Lactobacillus fermentum		66.21	0.50	FRU 99	99		
Fusobacterium mortiferum		27.94	0.42		1		
Actinomyces naeslundii		1.93	0.33		99		
07401103	**Low discrimination**						**07401103**
					SPO		
Actinomyces naeslundii		61.26	0.36	MLT 94	1		
Clostridium chauvoei		24.59	0.07	FRU 95	99		
Lactobacillus fermentum		14.16	0.15		1		
07401120	**Good identification**						**07401120**
					GRA		
Prevotella melaninogenica		96.59	0.50	MLT 99	1		
Lactobacillus fermentum		2.45	0.41	FRU 90	99		
Clostridium perfringens		0.36	0.15		99		
07401122	**Low discrimination**						**07401122**
					GRA		
Prevotella melaninogenica		85.67	0.37	MLT 99	1		
Lactobacillus fermentum		11.42	0.41	FRU 90	99		
Actinomyces naeslundii		1.00	0.33		99		
07401123	**Low discrimination**						**07401123**
					GRA		
Actinomyces naeslundii		63.37	0.36	MLT 94	99		
Prevotella melaninogenica		36.63	0.03	FRU 95	1		
07401400	**Low discrimination**						**07401400**
					COC	CAT	
Lactobacillus fermentum		57.44	0.46	FRU 99	1	1	
Gemella morbillorum		35.60	0.31		99	1	
Propionibacterium granulosum		5.42	0.24		1	90	
07401402	**Good identification**						**07401402**
					CAT		
Lactobacillus fermentum		97.70	0.46	FRU 99	1		
Propionibacterium granulosum		2.30	0.13		90		
07401420	**Low discrimination**						**07401420**
					COC	CAT	
Lactobacillus fermentum		78.29	0.38	FRU 99	1	1	
Gemella morbillorum		16.17	0.14		99	1	
Propionibacterium granulosum		5.54	0.13		1	90	
07401422	**Good identification**						**07401422**
Lactobacillus fermentum		100.0	0.38	FRU 99			
07401500	**Low discrimination**						**07401500**
					GRA		
Lactobacillus fermentum		85.85	0.50	FRU 99	99		
Bifidobacterium longum subsp. longum		10.86	0.27		99		
Fusobacterium mortiferum		3.29	0.24		1		
07401502	**Low discrimination**						**07401502**
Lactobacillus fermentum		88.77	0.50	FRU 99			
Bifidobacterium longum subsp. longum		11.23	0.27				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07401520	**Low discrimination**							**07401520**
					GRA			
Bifidobacterium longum subsp. longum		52.06	0.37	MLT 99	99			
Lactobacillus fermentum		34.30	0.41	FRU 90	99			
Prevotella melaninogenica		13.64	0.15		1			
07401522	**Low discrimination**							**07401522**
					GRA			
Bifidobacterium longum subsp. longum		58.02	0.37	MLT 99	99			
Lactobacillus fermentum		38.23	0.41	FRU 90	99			
Prevotella melaninogenica		2.90	0.03		1			
07401530	**Good identification**							**07401530**
Bifidobacterium longum subsp. longum		100.0	0.41	MLT 99				
07401532	**Good identification**							**07401532**
Bifidobacterium longum subsp. longum		100.0	0.41	MLT 99				
07402000	**Good identification**							**07402000**
					COC	GRA	CAT	
Propionibacterium granulosum		97.83	0.56	FRU 99	1	99	90	
Gemella morbillorum		1.45	0.17		99	99	1	
Megasphaera elsdenii		0.29	0.07		99	1	1	
07402002	**Good identification**							**07402002**
Propionibacterium granulosum		100.0	0.45	FRU 99				
07402010	**Good identification**							**07402010**
Propionibacterium granulosum		100.0	0.41	FRU 99				
07402020	**Good identification**							**07402020**
					COC	CAT		
Propionibacterium granulosum		99.34	0.45	FRU 99	1	90		
Gemella morbillorum		0.66	0.01		99	1		
07402022	**Good identification**							**07402022**
Propionibacterium granulosum		100.0	0.35	FRU 99				
07403000	**Good identification**							**07403000**
					COC	CAT		
Propionibacterium granulosum		98.39	0.56	FRU 99	1	90		
Gemella morbillorum		0.79	0.13		99	1		
Actinomyces naeslundii		0.45	0.30		1	25		
07403002	**Good identification**							**07403002**
Propionibacterium granulosum		98.19	0.45	FRU 99				
Actinomyces naeslundii		1.81	0.30					
07403010	**Good identification**							**07403010**
Propionibacterium granulosum		100.0	0.41	FRU 99				
07403020	**Good identification**							**07403020**
Propionibacterium granulosum		98.19	0.45	FRU 99				
Actinomyces naeslundii		1.81	0.30					
07403022	**Acceptable identification**							**07403022**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		93.14	0.35	FRU 99				
Actinomyces naeslundii		6.86	0.30					
07403100	**Low discrimination**							**07403100**
					GRA	CAT		
Actinomyces naeslundii		38.05	0.38	MLT 94	99	25		
Propionibacterium granulosum		29.34	0.21	FRU 95	99	90		
Fusobacterium mortiferum		25.81	0.24		1	5		
07403101	**Good identification**							**07403101**
Actinomyces naeslundii		100.0	0.41	MLT 94				
07403102	**Low discrimination**							**07403102**
					CAT			
Actinomyces naeslundii		67.09	0.38	MLT 94	25			
Propionibacterium granulosum		12.94	0.11	FRU 95	90			
Lactobacillus fermentum		11.98	0.15		1			
07403103	**Good identification**							**07403103**
Actinomyces naeslundii		100.0	0.41	MLT 94				
07403121	**Good identification**							**07403121**
Actinomyces naeslundii		100.0	0.41	MLT 94				
07403122	**Low discrimination**							**07403122**
					GRA			
Actinomyces naeslundii		69.13	0.38	MLT 94	99			
Prevotella melaninogenica		30.87	0.03	FRU 95	1			
07403123	**Good identification**							**07403123**
Actinomyces naeslundii		100.0	0.41	MLT 94				
07404000	**Low discrimination**							**07404000**
					COC	GRA	CAT	
Gemella morbillorum		83.49	0.35	RAF 1	99	99	1	
Fusobacterium mortiferum		9.67	0.29	XYL 10	1	1	5	
Propionibacterium granulosum		6.84	0.24		1	99	90	
07404100	**Acceptable identification**							**07404100**
					COC	GRA		
Fusobacterium mortiferum		90.83	0.36	FRU 99	1	1		
Actinomyces meyeri		5.78	0.06	XYL 7	1	99		
Gemella morbillorum		3.40	0.01		99	99		
07405002	**Acceptable identification**							**07405002**
					GRA	CAT		
Lactobacillus fermentum		92.77	0.42	FRU 99	99	1		
Propionibacterium granulosum		3.72	0.13		99	90		
Fusobacterium mortiferum		3.51	0.16		1	5		
07405100	**Low discrimination**							**07405100**
					GRA			
Lactobacillus fermentum		65.89	0.46	FRU 99	99			
Fusobacterium mortiferum		32.07	0.39		1			
Actinomyces meyeri		2.04	0.09		99			
07405102	**Acceptable identification**							**07405102**
					GRA			
Lactobacillus fermentum		94.87	0.46	FRU 99	99			
Fusobacterium mortiferum		5.13	0.22		1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07405120	**Low discrimination**					**07405120**
					GRA	
Lactobacillus fermentum		56.96	0.37	FRU 99	99	
Prevotella melaninogenica		38.58	0.15		1	
Bifidobacterium longum subsp. longum		4.46	0.11		99	
07405122	**Low discrimination**					**07405122**
					GRA	
Lactobacillus fermentum		52.86	0.37	FRU 99	99	
Parabacteroides distasonis		29.40	0.30		1	
Prevotella melaninogenica		6.82	0.03		1	
07405322	**Good identification**					**07405322**
Parabacteroides distasonis		100.0	0.38	FRU 99		
07405400	**Low discrimination**					**07405400**
					COC	
Lactobacillus fermentum		89.51	0.42	FRU 99	1	
Gemella morbillorum		10.49	0.14		99	
07405402	**Good identification**					**07405402**
Lactobacillus fermentum		100.0	0.42	FRU 99		
07405500	**Low discrimination**					**07405500**
Lactobacillus fermentum		60.75	0.46	FRU 99		
Bifidobacterium longum subsp. longum		39.25	0.35			
07405502	**Low discrimination**					**07405502**
Lactobacillus fermentum		60.75	0.46	FRU 99		
Bifidobacterium longum subsp. longum		39.25	0.35			
07405510	**Good identification**					**07405510**
Bifidobacterium longum subsp. longum		100.0	0.39	MLT 99		
07405512	**Good identification**					**07405512**
Bifidobacterium longum subsp. longum		100.0	0.39	MLT 99		
07405520	**Low discrimination**					**07405520**
					GRA	
Bifidobacterium longum subsp. longum		84.81	0.46	MLT 99	99	
Lactobacillus fermentum		10.94	0.37	FRU 90	99	
Bacteroides vulgatus		4.25	0.09		1	
07405522	**Low discrimination**					**07405522**
					GRA	
Bifidobacterium longum subsp. longum		85.25	0.46	MLT 99	99	
Lactobacillus fermentum		11.00	0.37	FRU 90	99	
Parabacteroides distasonis		3.75	0.27		1	
07405530	**Good identification**					**07405530**
Bifidobacterium longum subsp. longum		100.0	0.50	MLT 99		
07405532	**Good identification**					**07405532**
					GRA	
Bifidobacterium longum subsp. longum		98.63	0.50	MLT 99	99	
Parabacteroides distasonis		1.37	0.22	FRU 90	1	
07410000	**Low discrimination**					**07410000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	CAT	
Propionibacterium granulosum		82.22	0.92		1	99	90	
Fusobacterium mortiferum		15.59	0.83		1	1	5	
Gemella morbillorum		1.13	0.53		99	99	1	
07410001	**Low discrimination**							**07410001**
					COC	GRA	CAT	
Propionibacterium granulosum		82.31	0.58	URE 1	1	99	90	
Fusobacterium mortiferum		15.61	0.49		1	1	5	
Gemella morbillorum		1.13	0.19		99	99	1	
07410002	**Good identification**							**07410002**
					GRA	CAT		
Propionibacterium granulosum		91.44	0.82		99	90		
Fusobacterium mortiferum		7.71	0.66		1	5		
Lactobacillus fermentum		0.40	0.46		99	1		
07410003	**Low discrimination**							**07410003**
					GRA	CAT		
Propionibacterium granulosum		85.31	0.47	URE 1	99	90		
Fusobacterium mortiferum		7.19	0.32		1	5		
Actinomyces viscosus		6.77	0.19		99	76		
07410010	**Very good identification**							**07410010**
					COC	GRA	CAT	
Propionibacterium granulosum		98.41	0.77	MLZ 12	1	99	90	
Fusobacterium mortiferum		1.38	0.49		1	1	5	
Gemella morbillorum		0.10	0.19		99	99	1	
07410011	**Good identification**							**07410011**
					GRA	CAT		
Propionibacterium granulosum		98.62	0.43	MLZ 12	99	90		
Fusobacterium mortiferum		1.39	0.14	URE 1	1	5		
07410012	**Very good identification**							**07410012**
					GRA	CAT		
Propionibacterium granulosum		99.38	0.67	MLZ 12	99	90		
Fusobacterium mortiferum		0.62	0.32		1	5		
07410020	**Very good identification**							**07410020**
					COC	GRA	CAT	
Propionibacterium granulosum		98.31	0.82		1	99	90	
Fusobacterium mortiferum		0.75	0.49		1	1	5	
Gemella morbillorum		0.60	0.37		99	99	1	
07410021	**Good identification**							**07410021**
					GRA	CAT		
Propionibacterium granulosum		97.82	0.47	URE 1	99	90		
Actinomyces naeslundii		0.83	0.27		99	25		
Fusobacterium mortiferum		0.75	0.14		1	5		
07410022	**Very good identification**							**07410022**
					GRA	CAT		
Propionibacterium granulosum		98.85	0.71		99	90		
Lactobacillus fermentum		0.57	0.38		99	1		
Fusobacterium mortiferum		0.34	0.32		1	5		
07410023	**Good identification**							**07410023**
Propionibacterium granulosum		96.71	0.37	URE 1				
Actinomyces naeslundii		3.29	0.27					
07410030	**Very good identification**							**07410030**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	CAT	
Propionibacterium granulosum		99.90	0.67	MLZ 12	1	99	90	
Fusobacterium mortiferum		0.06	0.14		1	1	5	
Gemella morbillorum		0.05	0.03		99	99	1	
07410032	**Very good identification**							**07410032**
Propionibacterium granulosum		100.0	0.56	MLZ 12				
07410040	**Low discrimination**							**07410040**
					GRA	SPO	CAT	
Propionibacterium granulosum		77.12	0.58	SOR 1	99	1	90	
Fusobacterium mortiferum		14.62	0.49		1	1	5	
Clostridium botulinum biovar B		4.28	0.24		99	99	1	
07410042	**Acceptable identification**							**07410042**
					COC	GRA	CAT	
Propionibacterium granulosum		91.16	0.47	SOR 1	1	99	90	
Fusobacterium mortiferum		7.68	0.32		1	1	5	
Megasphaera elsdenii		1.15	0.09		99	1	1	
07410050	**Low discrimination**							**07410050**
					GRA	SPO	CAT	
Propionibacterium granulosum		82.34	0.43	MLZ 12	99	1	90	
Clostridium botulinum biovar B		16.50	0.18	SOR 1	99	99	1	
Fusobacterium mortiferum		1.16	0.14		1	1	5	
07410060	**Low discrimination**							**07410060**
					GRA	SPO	CAT	
Propionibacterium granulosum		89.04	0.47	SOR 1	99	1	90	
Clostridium botulinum biovar B		9.73	0.18		99	99	1	
Fusobacterium mortiferum		0.68	0.14		1	1	5	
07410062	**Good identification**							**07410062**
Propionibacterium granulosum		100.0	0.37	SOR 1				
07410100	**Very good identification**							**07410100**
					GRA	CAT		
Fusobacterium mortiferum		97.28	0.89		1	5		
Propionibacterium granulosum		2.22	0.58		99	90		
Lactobacillus fermentum		0.39	0.50		99	1		
07410101	**Good identification**							**07410101**
					GRA	CAT		
Fusobacterium mortiferum		96.11	0.55	URE 1	1	5		
Propionibacterium granulosum		2.19	0.24		99	90		
Actinomyces naeslundii		1.31	0.35		99	25		
07410102	**Good identification**							**07410102**
					GRA	CAT		
Fusobacterium mortiferum		90.08	0.73	BGL 10	1	5		
Propionibacterium granulosum		4.63	0.47		99	90		
Lactobacillus fermentum		3.23	0.50		99	1		
07410103	**Low discrimination**							**07410103**
					GRA			
Fusobacterium mortiferum		51.87	0.39	URE 1	1			
Actinomyces viscosus		37.23	0.24	BGL 10	99			
Actinomyces naeslundii		6.38	0.35		99			
07410110	**Low discrimination**							**07410110**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		76.21	0.55	MLZ 1	1	5		
Propionibacterium granulosum		23.49	0.43		99	90		
Lactobacillus fermentum		0.30	0.15		99	1		
07410112	**Low discrimination**							**07410112**
					GRA	CAT		
Fusobacterium mortiferum		57.83	0.39	MLZ 1	1	5		
Propionibacterium granulosum		40.10	0.33	BGL 10	99	90		
Lactobacillus fermentum		2.08	0.15		99	1		
07410120	**Low discrimination**							**07410120**
					GRA	CAT		
Fusobacterium mortiferum		45.33	0.55	NAG 1	1	5		
Propionibacterium granulosum		25.61	0.47		99	90		
Prevotella melaninogenica		21.36	0.32		1	1		
07410122	**Low discrimination**							**07410122**
					GRA	CAT		
Propionibacterium granulosum		27.98	0.37	LAC 1	99	90		
Lactobacillus fermentum		26.07	0.41		99	1		
Fusobacterium mortiferum		22.01	0.39		1	5		
07410140	**Good identification**							**07410140**
					GRA	CAT		
Fusobacterium mortiferum		97.39	0.55	SOR 1	1	5		
Propionibacterium granulosum		2.22	0.24		99	90		
Lactobacillus fermentum		0.39	0.15		99	1		
07410142	**Low discrimination**							**07410142**
					GRA	CAT		
Fusobacterium mortiferum		89.99	0.39	SOR 1	1	5		
Propionibacterium granulosum		4.62	0.13	BGL 10	99	90		
Lactobacillus fermentum		3.23	0.15		99	1		
07410200	**Good identification**							**07410200**
					COC	GRA	CAT	
Propionibacterium granulosum		93.50	0.66	RHA 3	1	99	90	
Fusobacterium mortiferum		5.79	0.49		1	1	5	
Gemella morbillorum		0.42	0.19		99	99	1	
07410202	**Good identification**							**07410202**
					GRA	CAT		
Propionibacterium granulosum		97.32	0.56	RHA 3	99	90		
Fusobacterium mortiferum		2.68	0.32		1	5		
07410210	**Good identification**							**07410210**
					GRA	CAT		
Propionibacterium granulosum		99.54	0.51	RHA 3	99	90		
Fusobacterium mortiferum		0.46	0.14	MLZ 12	1	5		
07410212	**Good identification**							**07410212**
Propionibacterium granulosum		100.0	0.41	RHA 3				
07410220	**Good identification**							**07410220**
					COC	GRA	CAT	
Propionibacterium granulosum		99.55	0.56	RHA 3	1	99	90	
Fusobacterium mortiferum		0.25	0.14		1	1	5	
Gemella morbillorum		0.20	0.03		99	99	1	
07410222	**Good identification**							**07410222**
Propionibacterium granulosum		100.0	0.45	RHA 3				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07410230	**Good identification**							**07410230**
Propionibacterium granulosum		100.0	0.41	RHA 3				
07410300	**Acceptable identification**							**07410300**
					GRA	CAT		
Fusobacterium mortiferum		93.12	0.55	RHA 1	1	5		
Propionibacterium granulosum		6.51	0.32		99	90		
Lactobacillus fermentum		0.37	0.15		99	1		
07410302	**Low discrimination**							**07410302**
					GRA	CAT		
Fusobacterium mortiferum		83.81	0.39	RHA 1	1	5		
Propionibacterium granulosum		13.18	0.22	BGL 10	99	90		
Lactobacillus fermentum		3.01	0.15		99	1		
07410400	**Low discrimination**							**07410400**
					COC	GRA	CAT	
Propionibacterium granulosum		68.58	0.58	ARA 1	1	99	90	
Fusobacterium mortiferum		13.00	0.49		1	1	5	
Gemella morbillorum		10.33	0.37		99	99	1	
07410402	**Low discrimination**							**07410402**
					GRA	CAT		
Propionibacterium granulosum		65.84	0.47	ARA 1	99	90		
Lactobacillus fermentum		28.21	0.46		99	1		
Fusobacterium mortiferum		5.55	0.32		1	5		
07410410	**Good identification**							**07410410**
					COC	GRA	CAT	
Propionibacterium granulosum		97.54	0.43	ARA 1	1	99	90	
Fusobacterium mortiferum		1.37	0.14	MLZ 12	1	1	5	
Gemella morbillorum		1.09	0.03		99	99	1	
07410420	**Low discrimination**							**07410420**
					COC	CAT		
Propionibacterium granulosum		82.14	0.47	ARA 1	1	90		
Lactobacillus fermentum		11.73	0.38		1	1		
Gemella morbillorum		5.50	0.20		99	1		
07410422	**Low discrimination**							**07410422**
					CAT			
Propionibacterium granulosum		63.64	0.37	ARA 1	90			
Lactobacillus fermentum		36.36	0.38		1			
07410500	**Low discrimination**							**07410500**
					GRA	CAT		
Fusobacterium mortiferum		69.57	0.55	ARA 1	1	5		
Lactobacillus fermentum		27.48	0.50		99	1		
Propionibacterium granulosum		1.59	0.24		99	90		
07410502	**Low discrimination**							**07410502**
					GRA	CAT		
Lactobacillus fermentum		76.50	0.50	GAL 99	99	1		
Fusobacterium mortiferum		21.52	0.39		1	5		
Propionibacterium granulosum		1.11	0.13		99	90		
07410520	**Low discrimination**							**07410520**
					GRA			
Lactobacillus fermentum		77.27	0.41	GAL 99	99			
Bifidobacterium longum subsp. longum		10.66	0.20		99			
Fusobacterium mortiferum		5.93	0.21		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07410522	**Low discrimination**						**07410522**
					GRA		
Lactobacillus fermentum		86.19	0.41	GAL 99	99		
Bifidobacterium longum subsp. longum		11.89	0.20		99		
Parabacteroides distasonis		1.92	0.10		1		
07411000	**Low discrimination**						**07411000**
					GRA	CAT	
Propionibacterium granulosum		70.97	0.92		99	90	
Fusobacterium mortiferum		20.19	0.86		1	5	
Lactobacillus fermentum		7.60	0.80		99	1	
07411001	**Low discrimination**						**07411001**
					GRA	CAT	
Propionibacterium granulosum		68.52	0.58	URE 1	99	90	
Fusobacterium mortiferum		19.49	0.52		1	5	
Lactobacillus fermentum		7.34	0.46		99	1	
07411002	**Low discrimination**						**07411002**
					GRA	CAT	
Propionibacterium granulosum		64.15	0.82		99	90	
Lactobacillus fermentum		27.48	0.80		99	1	
Fusobacterium mortiferum		8.11	0.69		1	5	
07411003	**Low discrimination**						**07411003**
					CAT		
Propionibacterium granulosum		55.33	0.47	URE 1	90		
Lactobacillus fermentum		23.70	0.46		1		
Actinomyces naeslundii		11.28	0.50		25		
07411010	**Very good identification**						**07411010**
					GRA	CAT	
Propionibacterium granulosum		97.05	0.77	MLZ 12	99	90	
Fusobacterium mortiferum		2.05	0.52		1	5	
Lactobacillus fermentum		0.77	0.46		99	1	
07411011	**Good identification**						**07411011**
					GRA	CAT	
Propionibacterium granulosum		96.09	0.43	MLZ 12	99	90	
Fusobacterium mortiferum		2.02	0.17	URE 1	1	5	
Actinomyces naeslundii		1.89	0.29		99	25	
07411012	**Very good identification**						**07411012**
					GRA	CAT	
Propionibacterium granulosum		96.00	0.67	MLZ 12	99	90	
Lactobacillus fermentum		3.05	0.46		99	1	
Fusobacterium mortiferum		0.90	0.35		1	5	
07411020	**Low discrimination**						**07411020**
					GRA	CAT	
Propionibacterium granulosum		85.27	0.82		99	90	
Lactobacillus fermentum		12.18	0.72		99	1	
Fusobacterium mortiferum		0.98	0.52		1	5	
07411021	**Low discrimination**						**07411021**
					CAT		
Propionibacterium granulosum		73.17	0.47	URE 1	90		
Actinomyces naeslundii		14.92	0.50		25		
Lactobacillus fermentum		10.45	0.38		1		
07411022	**Low discrimination**						**07411022**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		62.44	0.71		90		
Lactobacillus fermentum		35.67	0.72		1		
Clostridium perfringens		0.54	0.29		1		
07411023	**Low discrimination**						**07411023**
					CAT		
Propionibacterium granulosum		41.90	0.37	URE 1	90		
Actinomyces naeslundii		34.17	0.50		25		
Lactobacillus fermentum		23.93	0.38		1		
07411030	**Very good identification**						**07411030**
					GRA	CAT	
Propionibacterium granulosum		98.76	0.67	MLZ 12	99	90	
Lactobacillus fermentum		1.05	0.38		99	1	
Fusobacterium mortiferum		0.08	0.17		1	5	
07411032	**Very good identification**						**07411032**
					CAT		
Propionibacterium granulosum		95.75	0.56	MLZ 12	90		
Lactobacillus fermentum		4.05	0.38		1		
Actinomyces naeslundii		0.20	0.26		25		
07411040	**Low discrimination**						**07411040**
					GRA	CAT	
Propionibacterium granulosum		70.70	0.58	SOR 1	99	90	
Fusobacterium mortiferum		20.11	0.52		1	5	
Lactobacillus fermentum		7.57	0.46		99	1	
07411042	**Low discrimination**						**07411042**
					GRA	CAT	
Propionibacterium granulosum		63.34	0.47	SOR 1	99	90	
Lactobacillus fermentum		27.14	0.46		99	1	
Fusobacterium mortiferum		8.01	0.35		1	5	
07411043	**Low discrimination**						**07411043**
Actinomyces naeslundii		78.08	0.38	MLT 94			
Propionibacterium granulosum		21.92	0.13				
07411050	**Good identification**						**07411050**
					GRA	CAT	
Propionibacterium granulosum		97.94	0.43	MLZ 12	99	90	
Fusobacterium mortiferum		2.06	0.17	SOR 1	1	5	
07411060	**Low discrimination**						**07411060**
					CAT		
Propionibacterium granulosum		78.51	0.47	SOR 1	90		
Lactobacillus fermentum		11.21	0.38		1		
Clostridium perfringens		6.83	0.22		1		
07411061	**Low discrimination**						**07411061**
Actinomyces naeslundii		78.08	0.38	MLT 94			
Propionibacterium granulosum		21.92	0.13				
07411062	**Low discrimination**						**07411062**
					CAT		
Propionibacterium granulosum		54.61	0.37	SOR 1	90		
Lactobacillus fermentum		31.20	0.38		1		
Clostridium perfringens		8.95	0.16		1		
07411063	**Good identification**						**07411063**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	MLT 94			
07411100	**Low discrimination**						**07411100**
					GRA	CAT	
Fusobacterium mortiferum		77.86	0.92		1	5	
Lactobacillus fermentum		20.50	0.84		99	1	
Propionibacterium granulosum		1.19	0.58		99	90	
07411101	**Low discrimination**						**07411101**
					GRA		
Fusobacterium mortiferum		64.05	0.58	URE 1	1		
Lactobacillus fermentum		16.86	0.50		99		
Actinomyces naeslundii		14.00	0.58		99		
07411102	**Low discrimination**						**07411102**
					GRA	CAT	
Lactobacillus fermentum		68.93	0.84		99	1	
Fusobacterium mortiferum		29.09	0.76		1	5	
Propionibacterium granulosum		1.00	0.47		99	90	
07411103	**Low discrimination**						**07411103**
					GRA		
Lactobacillus fermentum		41.13	0.50	URE 1	99		
Actinomyces naeslundii		34.15	0.58		99		
Fusobacterium mortiferum		17.36	0.42		1		
07411110	**Low discrimination**						**07411110**
					GRA	CAT	
Fusobacterium mortiferum		67.35	0.58	MLZ 1	1	5	
Lactobacillus fermentum		17.73	0.50		99	1	
Propionibacterium granulosum		13.84	0.43		99	90	
07411111	**Low discrimination**						**07411111**
					GRA		
Actinomyces naeslundii		43.68	0.36	MLT 94	99		
Fusobacterium mortiferum		38.34	0.24	MLZ 5	1		
Lactobacillus fermentum		10.10	0.15		99		
07411112	**Low discrimination**						**07411112**
					GRA	CAT	
Lactobacillus fermentum		59.72	0.50	MLZ 1	99	1	
Fusobacterium mortiferum		25.20	0.42		1	5	
Propionibacterium granulosum		11.65	0.33		99	90	
07411113	**Low discrimination**						**07411113**
Actinomyces naeslundii		81.23	0.36	MLT 94			
Lactobacillus fermentum		18.77	0.15	MLZ 5			
07411120	**Low discrimination**						**07411120**
					GRA		
Prevotella melaninogenica		72.94	0.66	MLT 99	1		
Lactobacillus fermentum		20.37	0.76		99		
Clostridium perfringens		2.96	0.49		99		
07411122	**Low discrimination**						**07411122**
					GRA		
Lactobacillus fermentum		52.45	0.76		99		
Prevotella melaninogenica		35.77	0.54		1		
Clostridium perfringens		3.59	0.43		99		
07411123	**Low discrimination**						**07411123**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		56.86	0.58	MLT 94	99		
Lactobacillus fermentum		22.83	0.41		99		
Prevotella melaninogenica		15.57	0.19		1		
07411131	**Low discrimination**						**07411131**
					GRA		
Actinomyces naeslundii		78.38	0.36	MLT 94	99		
Prevotella melaninogenica		21.62	-.03	MLZ 5	1		
07411132	**Low discrimination**						**07411132**
					GRA		
Parabacteroides distasonis		44.22	0.43	XYL 90	1		
Lactobacillus fermentum		22.85	0.41		99		
Prevotella melaninogenica		15.58	0.19		1		
07411133	**Good identification**						**07411133**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07411140	**Low discrimination**						**07411140**
					GRA		
Fusobacterium mortiferum		76.05	0.58	SOR 1	1		
Lactobacillus fermentum		20.03	0.50		99		
Actinomyces naeslundii		1.96	0.42		99		
07411141	**Low discrimination**						**07411141**
					GRA		
Actinomyces naeslundii		75.15	0.45	MLT 94	99		
Fusobacterium mortiferum		19.67	0.24		1		
Lactobacillus fermentum		5.18	0.15		99		
07411142	**Low discrimination**						**07411142**
					GRA		
Lactobacillus fermentum		64.35	0.50	SOR 1	99		
Fusobacterium mortiferum		27.15	0.42		1		
Actinomyces naeslundii		6.29	0.42		99		
07411143	**Acceptable identification**						**07411143**
Actinomyces naeslundii		93.55	0.45	MLT 94			
Lactobacillus fermentum		6.45	0.15				
07411161	**Low discrimination**						**07411161**
					GRA		
Actinomyces naeslundii		87.32	0.45	MLT 94	99		
Prevotella melaninogenica		7.19	-.03		1		
Clostridium perfringens		5.50	0.02		99		
07411163	**Good identification**						**07411163**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07411200	**Low discrimination**						**07411200**
					GRA	CAT	
Propionibacterium granulosum		88.23	0.66	RHA 3	99	90	
Fusobacterium mortiferum		8.20	0.52		1	5	
Lactobacillus fermentum		3.09	0.46		99	1	
07411202	**Low discrimination**						**07411202**
					GRA	CAT	
Propionibacterium granulosum		84.66	0.56	RHA 3	99	90	
Lactobacillus fermentum		11.85	0.46		99	1	
Fusobacterium mortiferum		3.50	0.35		1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07411210	**Good identification**						**07411210**
					GRA	CAT	
Propionibacterium granulosum		99.32	0.51	RHA 3	99	90	
Fusobacterium mortiferum		0.68	0.17	MLZ 12	1	5	
07411212	**Good identification**						**07411212**
Propionibacterium granulosum		100.0	0.41	RHA 3			
07411220	**Acceptable identification**						**07411220**
					GRA	CAT	
Propionibacterium granulosum		94.95	0.56	RHA 3	99	90	
Lactobacillus fermentum		4.43	0.38		99	1	
Fusobacterium mortiferum		0.36	0.17		1	5	
07411222	**Low discrimination**						**07411222**
					GRA	CAT	
Propionibacterium granulosum		78.66	0.45	RHA 3	99	90	
Lactobacillus fermentum		14.68	0.38		99	1	
Prevotella oralis		4.06	0.13		1	1	
07411230	**Good identification**						**07411230**
Propionibacterium granulosum		100.0	0.41	RHA 3			
07411300	**Low discrimination**						**07411300**
					GRA	CAT	
Fusobacterium mortiferum		76.16	0.58	RHA 1	1	5	
Lactobacillus fermentum		20.05	0.50		99	1	
Propionibacterium granulosum		3.55	0.32		99	90	
07411302	**Low discrimination**						**07411302**
					GRA		
Lactobacillus fermentum		64.55	0.50	RHA 1	99		
Fusobacterium mortiferum		27.24	0.42		1		
Prevotella oralis		3.01	0.11		1		
07411322	**Low discrimination**						**07411322**
					GRA		
Parabacteroides distasonis		70.84	0.56	XYL 90	1		
Prevotella oralis		17.44	0.33		1		
Lactobacillus fermentum		6.57	0.41		99		
07411332	**Good identification**						**07411332**
Parabacteroides distasonis		99.54	0.52	XYL 90			
Prevotella oralis		0.46	-.01				
07411400	**Low discrimination**						**07411400**
					GRA	CAT	
Lactobacillus fermentum		88.49	0.80		99	1	
Propionibacterium granulosum		8.35	0.58		99	90	
Fusobacterium mortiferum		2.37	0.52		1	5	
07411401	**Low discrimination**						**07411401**
					GRA	CAT	
Lactobacillus fermentum		89.20	0.46	URE 1	99	1	
Propionibacterium granulosum		8.41	0.24		99	90	
Fusobacterium mortiferum		2.39	0.17		1	5	
07411402	**Very good identification**						**07411402**
					GRA	CAT	
Lactobacillus fermentum		97.40	0.80		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		2.30	0.47		99	90	
Fusobacterium mortiferum		0.29	0.35		1	5	
07411403	**Good identification**						**07411403**
					CAT		
Lactobacillus fermentum		97.70	0.46	URE 1	1		
Propionibacterium granulosum		2.30	0.13		90		
07411410	**Low discrimination**						**07411410**
					CAT		
Propionibacterium granulosum		54.60	0.43	ARA 1	90		
Lactobacillus fermentum		42.88	0.46	MLZ 12	1		
Bifidobacterium longum subsp. longum		1.37	0.13		1		
07411412	**Low discrimination**						**07411412**
					CAT		
Lactobacillus fermentum		74.06	0.46	MLZ 1	1		
Propionibacterium granulosum		23.57	0.33		90		
Bifidobacterium longum subsp. longum		2.37	0.13		1		
07411420	**Good identification**						**07411420**
					COC	CAT	
Lactobacillus fermentum		92.85	0.72		1	1	
Propionibacterium granulosum		6.57	0.47		1	90	
Gemella morbillorum		0.24	0.16		99	1	
07411421	**Acceptable identification**						**07411421**
					CAT		
Lactobacillus fermentum		93.39	0.38	URE 1	1		
Propionibacterium granulosum		6.61	0.13		90		
07411422	**Very good identification**						**07411422**
					CAT		
Lactobacillus fermentum		98.01	0.72		1		
Propionibacterium granulosum		1.73	0.37		90		
Bifidobacterium longum subsp. longum		0.22	0.20		1		
07411423	**Good identification**						**07411423**
Lactobacillus fermentum		100.0	0.38	URE 1			
07411430	**Low discrimination**						**07411430**
					CAT		
Lactobacillus fermentum		42.77	0.38	MLZ 1	1		
Propionibacterium granulosum		40.84	0.33		90		
Bifidobacterium longum subsp. longum		16.39	0.24		1		
07411432	**Low discrimination**						**07411432**
					CAT		
Lactobacillus fermentum		61.65	0.38	MLZ 1	1		
Bifidobacterium longum subsp. longum		23.63	0.24		1		
Propionibacterium granulosum		14.72	0.22		90		
07411440	**Low discrimination**						**07411440**
					GRA	CAT	
Lactobacillus fermentum		89.20	0.46	SOR 1	99	1	
Propionibacterium granulosum		8.41	0.24		99	90	
Fusobacterium mortiferum		2.39	0.17		1	5	
07411442	**Good identification**						**07411442**
					CAT		
Lactobacillus fermentum		97.70	0.46	SOR 1	1		
Propionibacterium granulosum		2.30	0.13		90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07411460	**Acceptable identification**					**07411460**
					CAT	
Lactobacillus fermentum		93.39	0.38	SOR 1	1	
Propionibacterium granulosum		6.61	0.13		90	
07411462	**Good identification**					**07411462**
Lactobacillus fermentum		100.0	0.38	SOR 1		
07411500	**Very good identification**					**07411500**
					GRA	
Lactobacillus fermentum		95.15	0.84		99	
Fusobacterium mortiferum		3.65	0.58		1	
Bifidobacterium longum subsp. longum		1.09	0.43		99	
07411501	**Acceptable identification**					**07411501**
					GRA	
Lactobacillus fermentum		94.50	0.50	URE 1	99	
Fusobacterium mortiferum		3.63	0.24		1	
Bifidobacterium longum subsp. longum		1.09	0.09		99	
07411502	**Very good identification**					**07411502**
					GRA	
Lactobacillus fermentum		98.37	0.84		99	
Bifidobacterium longum subsp. longum		1.13	0.43		99	
Fusobacterium mortiferum		0.42	0.42		1	
07411503	**Good identification**					**07411503**
Lactobacillus fermentum		98.05	0.50	URE 1		
Bifidobacterium longum subsp. longum		1.13	0.09			
Actinomyces naeslundii		0.82	0.24			
07411510	**Low discrimination**					**07411510**
					GRA	
Bifidobacterium longum subsp. longum		64.94	0.47	MLT 99	99	
Lactobacillus fermentum		33.51	0.50		99	
Fusobacterium mortiferum		1.29	0.24		1	
07411512	**Low discrimination**					**07411512**
Bifidobacterium longum subsp. longum		64.69	0.47	MLT 99		
Lactobacillus fermentum		33.38	0.50			
Bifidobacterium adolescentis		1.94	0.23			
07411520	**Low discrimination**					**07411520**
					GRA	
Lactobacillus fermentum		84.58	0.76		99	
Bifidobacterium longum subsp. longum		11.67	0.54		99	
Prevotella melaninogenica		3.06	0.32		1	
07411521	**Low discrimination**					**07411521**
					GRA	
Lactobacillus fermentum		83.38	0.41	URE 1	99	
Bifidobacterium longum subsp. longum		11.50	0.20		99	
Prevotella melaninogenica		3.02	-.03		1	
07411522	**Low discrimination**					**07411522**
					GRA	
Lactobacillus fermentum		85.40	0.76		99	
Bifidobacterium longum subsp. longum		11.78	0.54		99	
Parabacteroides distasonis		1.90	0.44		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07411523	**Low discrimination**						**07411523**
Lactobacillus fermentum		84.36	0.41	URE 1			
Bifidobacterium longum subsp. longum		11.64	0.20				
Actinomyces naeslundii		2.12	0.24				
07411530	**Very good identification**						**07411530**
					GRA		
Bifidobacterium longum subsp. longum		95.74	0.58	MLT 99	99		
Lactobacillus fermentum		4.12	0.41		99		
Prevotella melaninogenica		0.15	-.03		1		
07411532	**Good identification**						**07411532**
					GRA		
Bifidobacterium longum subsp. longum		91.31	0.58	MLT 99	99		
Parabacteroides distasonis		4.66	0.40		1		
Lactobacillus fermentum		3.93	0.41		99		
07411540	**Good identification**						**07411540**
					GRA		
Lactobacillus fermentum		95.25	0.50	SOR 1	99		
Fusobacterium mortiferum		3.65	0.24		1		
Bifidobacterium longum subsp. longum		1.10	0.09		99		
07411542	**Good identification**						**07411542**
Lactobacillus fermentum		98.86	0.50	SOR 1			
Bifidobacterium longum subsp. longum		1.14	0.09				
07411560	**Low discrimination**						**07411560**
					GRA		
Lactobacillus fermentum		83.21	0.41	SOR 1	99		
Bifidobacterium longum subsp. longum		11.48	0.20		99		
Prevotella melaninogenica		3.01	-.03		1		
07411562	**Low discrimination**						**07411562**
					GRA	CAT	
Lactobacillus fermentum		84.15	0.41	SOR 1	99	1	
Bifidobacterium longum subsp. longum		11.61	0.20		99	1	
Capnocytophaga ochracea		2.37	-.01		1	90	
07411600	**Low discrimination**						**07411600**
					GRA	CAT	
Lactobacillus fermentum		76.01	0.46	RHA 1	99	1	
Propionibacterium granulosum		21.95	0.32		99	90	
Fusobacterium mortiferum		2.04	0.17		1	5	
07411602	**Acceptable identification**						**07411602**
					CAT		
Lactobacillus fermentum		93.27	0.46	RHA 1	1		
Propionibacterium granulosum		6.73	0.22		90		
07411620	**Low discrimination**						**07411620**
					GRA	CAT	
Lactobacillus fermentum		77.83	0.38	RHA 1	99	1	
Propionibacterium granulosum		16.86	0.22		99	90	
Bacteroides vulgatus		5.32	-.04		1	3	
07411622	**Low discrimination**						**07411622**
					GRA	CAT	
Lactobacillus fermentum		85.98	0.38	RHA 1	99	1	
Parabacteroides distasonis		9.36	0.18		1	76	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		4.66	0.11		99	90		
07411700	**Acceptable identification**							**07411700**
					GRA			
Lactobacillus fermentum		94.00	0.50	RHA 1	99			
Fusobacterium mortiferum		3.61	0.24		1			
Bacteroides vulgatus		1.31	-.04		1			
07411702	**Good identification**							**07411702**
					GRA			
Lactobacillus fermentum		96.74	0.50	RHA 1	99			
Parabacteroides distasonis		2.15	0.18		1			
Bifidobacterium longum subsp. longum		1.11	0.09		99			
07411722	**Low discrimination**							**07411722**
					GRA			
Parabacteroides distasonis		84.56	0.53	XYL 90	1			
Lactobacillus fermentum		12.80	0.41		99			
Bifidobacterium longum subsp. longum		1.77	0.20		99			
07411732	**Acceptable identification**							**07411732**
					GRA			
Parabacteroides distasonis		93.81	0.48	XYL 90	1			
Bifidobacterium longum subsp. longum		6.19	0.24		99			
07412000	**Very good identification**							**07412000**
					GRA	CAT		
Propionibacterium granulosum		99.45	0.90		99	90		
Fusobacterium mortiferum		0.25	0.49		1	5		
Eubacterium limosum		0.19	0.29		99	1		
07412001	**Good identification**							**07412001**
					GRA	CAT		
Propionibacterium granulosum		99.02	0.56	URE 1	99	90		
Actinomyces naeslundii		0.54	0.32		99	25		
Fusobacterium mortiferum		0.25	0.14		1	5		
07412002	**Very good identification**							**07412002**
					COC	GRA	CAT	
Propionibacterium granulosum		99.83	0.79		1	99	90	
Fusobacterium mortiferum		0.11	0.32		1	1	5	
Megasphaera elsdenii		0.03	0.12		99	1	1	
07412003	**Good identification**							**07412003**
Propionibacterium granulosum		97.86	0.45	URE 1				
Actinomyces naeslundii		2.14	0.32					
07412010	**Very good identification**							**07412010**
					GRA	SPO	CAT	
Propionibacterium granulosum		99.43	0.75	MLZ 12	99	1	90	
Clostridium difficile		0.54	0.24		99	99	1	
Fusobacterium mortiferum		0.02	0.14		1	1	5	
07412011	**Good identification**							**07412011**
Propionibacterium granulosum		100.0	0.41	MLZ 12				
07412012	**Very good identification**							**07412012**
Propionibacterium granulosum		100.0	0.65	MLZ 12				
07412020	**Very good identification**							**07412020**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		99.96	0.79		99	90		
Actinomyces naeslundii		0.02	0.29		99	25		
Fusobacterium mortiferum		0.01	0.14		1	5		
07412021	**Good identification**							**07412021**
Propionibacterium granulosum		97.86	0.45	URE 1				
Actinomyces naeslundii		2.14	0.32					
07412022	**Very good identification**							**07412022**
Propionibacterium granulosum		99.94	0.69					
Actinomyces naeslundii		0.06	0.29					
07412023	**Acceptable identification**							**07412023**
Propionibacterium granulosum		91.96	0.35	URE 1				
Actinomyces naeslundii		8.04	0.32					
07412030	**Very good identification**							**07412030**
Propionibacterium granulosum		100.0	0.65	MLZ 12				
07412032	**Good identification**							**07412032**
Propionibacterium granulosum		100.0	0.54	MLZ 12				
07412040	**Good identification**							**07412040**
					COC	GRA	CAT	
Propionibacterium granulosum		95.96	0.56	SOR 1	1	99	90	
Megasphaera elsdenii		3.62	0.26		99	1	1	
Fusobacterium mortiferum		0.24	0.14		1	1	5	
07412042	**Good identification**							**07412042**
					COC	CAT	GRA	
Propionibacterium granulosum		97.75	0.45	SOR 1	1	90	99	
Megasphaera elsdenii		1.64	0.09		99	1	1	
Propionibacterium propionicum		0.61	0.12		1	1	99	
07412050	**Good identification**							**07412050**
					SPO	CAT		
Propionibacterium granulosum		97.27	0.41	MLZ 12	1	90		
Clostridium difficile		2.73	0.02	SOR 1	99	1		
07412060	**Good identification**							**07412060**
Propionibacterium granulosum		100.0	0.45	SOR 1				
07412062	**Good identification**							**07412062**
Propionibacterium granulosum		100.0	0.35	SOR 1				
07412100	**Low discrimination**							**07412100**
					GRA	CAT		
Propionibacterium granulosum		58.68	0.56	LAC 1	99	90		
Fusobacterium mortiferum		34.42	0.55		1	5		
Propionibacterium propionicum		6.04	0.43		99	1		
07412101	**Low discrimination**							**07412101**
					CAT			
Actinomyces naeslundii		36.16	0.40	MLT 94	25			
Propionibacterium propionicum		26.60	0.27	GAL 96	1			
Propionibacterium granulosum		23.48	0.21		90			
07412102	**Low discrimination**							**07412102**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				GRA	CAT	
Propionibacterium granulosum	51.83	0.45	LAC 1	99	90	
Propionibacterium propionicum	32.03	0.46		99	1	
Fusobacterium mortiferum	13.51	0.39		1	5	
07412110 Good identification						**07412110**
				GRA	CAT	
Propionibacterium granulosum	95.84	0.41	LAC 1	99	90	
Fusobacterium mortiferum	4.16	0.21	MLZ 12	1	5	
07412120 Acceptable identification						**07412120**
				GRA	CAT	
Propionibacterium granulosum	92.91	0.45	LAC 1	99	90	
Actinomyces naeslundii	3.85	0.37		99	25	
Fusobacterium mortiferum	2.20	0.21		1	5	
07412121 Low discrimination						**07412121**
Actinomyces naeslundii	86.03	0.40	MLT 94			
Propionibacterium granulosum	13.97	0.11	GAL 96			
07412122 Low discrimination						**07412122**
				CAT		
Propionibacterium granulosum	83.97	0.35	LAC 1	90		
Actinomyces naeslundii	13.93	0.37		25		
Propionibacterium propionicum	2.10	0.12		1		
07412123 Good identification						**07412123**
Actinomyces naeslundii	100.0	0.40	MLT 94			
07412140 Low discrimination						**07412140**
				GRA	CAT	
Propionibacterium propionicum	85.22	0.43	NIT 86	99	1	
Propionibacterium granulosum	8.36	0.21	MLT 99	99	90	
Fusobacterium mortiferum	4.90	0.21		1	5	
07412142 Good identification						**07412142**
				CAT		
Propionibacterium propionicum	97.26	0.46	NIT 86	1		
Propionibacterium granulosum	1.59	0.11	MLT 99	90		
Actinomyces naeslundii	1.15	0.24		25		
07412200 Very good identification						**07412200**
				GRA	CAT	
Propionibacterium granulosum	99.86	0.64	RHA 3	99	90	
Fusobacterium mortiferum	0.08	0.14		1	5	
Eubacterium limosum	0.06	-.06		99	1	
07412202 Good identification						**07412202**
Propionibacterium granulosum	100.0	0.54	RHA 3			
07412210 Good identification						**07412210**
Propionibacterium granulosum	100.0	0.49	RHA 3			
07412212 Good identification						**07412212**
Propionibacterium granulosum	100.0	0.39	RHA 3			
07412220 Good identification						**07412220**
Propionibacterium granulosum	100.0	0.54	RHA 3			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07412222	**Good identification**							**07412222**
Propionibacterium granulosum		100.0	0.43	RHA 3				
07412230	**Good identification**							**07412230**
Propionibacterium granulosum		100.0	0.39	RHA 3				
07412400	**Good identification**							**07412400**
					COC	GRA	CAT	
Propionibacterium granulosum		99.36	0.56	ARA 1	1	99	90	
Fusobacterium mortiferum		0.25	0.14		1	1	5	
Gemella morbillorum		0.20	0.03		99	99	1	
07412402	**Good identification**							**07412402**
Propionibacterium granulosum		100.0	0.45	ARA 1				
07412410	**Good identification**							**07412410**
Propionibacterium granulosum		100.0	0.41	ARA 1				
07412420	**Good identification**							**07412420**
Propionibacterium granulosum		100.0	0.45	ARA 1				
07412422	**Good identification**							**07412422**
Propionibacterium granulosum		100.0	0.35	ARA 1				
07413000	**Very good identification**							**07413000**
					GRA	CAT		
Propionibacterium granulosum		99.37	0.90		99	90		
Fusobacterium mortiferum		0.38	0.52		1	5		
Lactobacillus fermentum		0.14	0.46		99	1		
07413001	**Low discrimination**							**07413001**
					GRA	CAT		
Propionibacterium granulosum		88.11	0.56	URE 1	99	90		
Actinomyces naeslundii		11.56	0.55		99	25		
Fusobacterium mortiferum		0.34	0.17		1	5		
07413002	**Very good identification**							**07413002**
					CAT			
Propionibacterium granulosum		98.91	0.79		90			
Lactobacillus fermentum		0.57	0.46		1			
Actinomyces naeslundii		0.35	0.52		25			
07413003	**Low discrimination**							**07413003**
Propionibacterium granulosum		65.59	0.45	URE 1				
Actinomyces naeslundii		34.41	0.55					
07413010	**Very good identification**							**07413010**
					GRA	CAT		
Propionibacterium granulosum		99.94	0.75	MLZ 12	99	90		
Actinomyces naeslundii		0.03	0.30		99	25		
Fusobacterium mortiferum		0.03	0.17		1	5		
07413011	**Good identification**							**07413011**
Propionibacterium granulosum		95.18	0.41	MLZ 12				
Actinomyces naeslundii		4.82	0.33	URE 1				
07413012	**Very good identification**							**07413012**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		99.86	0.65	MLZ 12			
Actinomyces naeslundii		0.14	0.30				
07413020	**Very good identification**						**07413020**
					CAT		
Propionibacterium granulosum		99.43	0.79		90		
Actinomyces naeslundii		0.35	0.52		25		
Lactobacillus fermentum		0.19	0.38		1		
07413021	**Low discrimination**						**07413021**
Propionibacterium granulosum		65.59	0.45	URE 1			
Actinomyces naeslundii		34.41	0.55				
07413022	**Very good identification**						**07413022**
					CAT		
Propionibacterium granulosum		97.87	0.69		90		
Actinomyces naeslundii		1.38	0.52		25		
Lactobacillus fermentum		0.75	0.38		1		
07413023	**Low discrimination**						**07413023**
Actinomyces naeslundii		67.73	0.55	MLT 94			
Propionibacterium granulosum		32.27	0.35				
07413030	**Very good identification**						**07413030**
Propionibacterium granulosum		99.86	0.65	MLZ 12			
Actinomyces naeslundii		0.14	0.30				
07413032	**Good identification**						**07413032**
Propionibacterium granulosum		99.46	0.54	MLZ 12			
Actinomyces naeslundii		0.54	0.30				
07413040	**Good identification**						**07413040**
					GRA	CAT	
Propionibacterium granulosum		98.11	0.56	SOR 1	99	90	
Actinomyces naeslundii		1.51	0.39		99	25	
Fusobacterium mortiferum		0.37	0.17		1	5	
07413041	**Low discrimination**						**07413041**
Actinomyces naeslundii		69.62	0.42	MLT 94			
Propionibacterium granulosum		30.38	0.21				
07413042	**Acceptable identification**						**07413042**
					CAT		
Propionibacterium granulosum		93.64	0.45	SOR 1	90		
Actinomyces naeslundii		5.78	0.39		25		
Propionibacterium propionicum		0.58	0.12		1		
07413043	**Acceptable identification**						**07413043**
Actinomyces naeslundii		90.16	0.42	MLT 94			
Propionibacterium granulosum		9.84	0.11				
07413050	**Good identification**						**07413050**
Propionibacterium granulosum		100.0	0.41	MLZ 12			
07413060	**Acceptable identification**						**07413060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		94.19	0.45	SOR 1			
Actinomyces naeslundii		5.81	0.39				
07413061	**Acceptable identification**						**07413061**
Actinomyces naeslundii		90.16	0.42	MLT 94			
Propionibacterium granulosum		9.84	0.11				
07413062	**Low discrimination**						**07413062**
Propionibacterium granulosum		80.20	0.35	SOR 1			
Actinomyces naeslundii		19.80	0.39				
07413063	**Good identification**						**07413063**
Actinomyces naeslundii		100.0	0.42	MLT 94			
07413100	**Low discrimination**						**07413100**
					GRA	CAT	
Propionibacterium granulosum		40.55	0.56	LAC 1	99	90	
Fusobacterium mortiferum		35.67	0.58		1	5	
Actinomyces naeslundii		10.09	0.60		99	25	
07413101	**Good identification**						**07413101**
					CAT		
Actinomyces naeslundii		91.93	0.63	MLT 94	25		
Propionibacterium propionicum		2.82	0.27		1		
Propionibacterium granulosum		2.49	0.21		90		
07413102	**Low discrimination**						**07413102**
					CAT		
Propionibacterium granulosum		25.44	0.45	LAC 1	90		
Actinomyces naeslundii		25.33	0.60		25		
Lactobacillus fermentum		23.57	0.50		1		
07413103	**Good identification**						**07413103**
					CAT		
Actinomyces naeslundii		94.43	0.63	MLT 94	25		
Propionibacterium propionicum		4.34	0.30		1		
Propionibacterium granulosum		0.64	0.11		90		
07413110	**Low discrimination**						**07413110**
					GRA	CAT	
Propionibacterium granulosum		84.86	0.41	LAC 1	99	90	
Actinomyces naeslundii		8.15	0.38	MLZ 12	99	25	
Fusobacterium mortiferum		5.53	0.24		1	5	
07413111	**Good identification**						**07413111**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07413113	**Good identification**						**07413113**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07413121	**Very good identification**						**07413121**
					GRA	CAT	
Actinomyces naeslundii		98.60	0.63	MLT 94	99	25	
Prevotella melaninogenica		0.74	-.03		1	1	
Propionibacterium granulosum		0.67	0.11		99	90	
07413122	**Low discrimination**						**07413122**
					CAT		
Actinomyces naeslundii		54.61	0.60	MLT 94	25		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		16.94	0.41		1		
Propionibacterium granulosum		13.71	0.35		90		
07413123	**Very good identification**						**07413123**
Actinomyces naeslundii		100.0	0.63	MLT 94			
07413131	**Good identification**						**07413131**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07413132	**Low discrimination**						**07413132**
Actinomyces naeslundii		60.58	0.38	MLT 94			
Propionibacterium granulosum		39.42	0.20	MLZ 5			
07413133	**Good identification**						**07413133**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07413140	**Low discrimination**						**07413140**
					CAT		
Propionibacterium propionicum		61.22	0.43	NIT 86	1		
Actinomyces naeslundii		26.11	0.47	MLT 99	25		
Propionibacterium granulosum		6.00	0.21		90		
07413141	**Low discrimination**						**07413141**
Actinomyces naeslundii		85.20	0.50	MLT 94			
Propionibacterium propionicum		14.80	0.27				
07413142	**Low discrimination**						**07413142**
					CAT		
Propionibacterium propionicum		76.00	0.46	NIT 86	1		
Actinomyces naeslundii		21.61	0.47	MLT 99	25		
Propionibacterium granulosum		1.24	0.11		90		
07413143	**Low discrimination**						**07413143**
Actinomyces naeslundii		79.33	0.50	MLT 94			
Propionibacterium propionicum		20.67	0.30				
07413160	**Low discrimination**						**07413160**
					GRA	CAT	
Actinomyces naeslundii		85.49	0.47	MLT 94	99	25	
Prevotella melaninogenica		5.44	-.03		1	1	
Propionibacterium granulosum		4.92	0.11		99	90	
07413161	**Good identification**						**07413161**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07413162	**Good identification**						**07413162**
Actinomyces naeslundii		96.57	0.47	MLT 94			
Propionibacterium propionicum		3.43	0.12				
07413163	**Good identification**						**07413163**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07413200	**Very good identification**						**07413200**
					GRA	CAT	
Propionibacterium granulosum		99.88	0.64	RHA 3	99	90	
Fusobacterium mortiferum		0.12	0.17		1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07413202	**Good identification**						**07413202**
Propionibacterium granulosum		100.0	0.54	RHA 3			
07413210	**Good identification**						**07413210**
Propionibacterium granulosum		100.0	0.49	RHA 3			
07413212	**Good identification**						**07413212**
Propionibacterium granulosum		100.0	0.39	RHA 3			
07413220	**Good identification**						**07413220**
Propionibacterium granulosum		100.0	0.54	RHA 3			
07413222	**Good identification**						**07413222**
Propionibacterium granulosum		100.0	0.43	RHA 3			
07413230	**Good identification**						**07413230**
Propionibacterium granulosum		100.0	0.39	RHA 3			
07413400	**Low discrimination**						**07413400**
					GRA	CAT	
Propionibacterium granulosum		87.28	0.56	ARA 1	99	90	
Lactobacillus fermentum		12.39	0.46		99	1	
Fusobacterium mortiferum		0.33	0.17		1	5	
07413402	**Low discrimination**						**07413402**
					CAT		
Propionibacterium granulosum		63.78	0.45	ARA 1	90		
Lactobacillus fermentum		36.22	0.46		1		
07413410	**Good identification**						**07413410**
Propionibacterium granulosum		100.0	0.41	ARA 1			
07413420	**Low discrimination**						**07413420**
					CAT		
Propionibacterium granulosum		84.08	0.45	ARA 1	90		
Lactobacillus fermentum		15.92	0.38		1		
07413422	**Low discrimination**						**07413422**
					CAT		
Propionibacterium granulosum		56.91	0.35	ARA 1	90		
Lactobacillus fermentum		43.09	0.38		1		
07413500	**Acceptable identification**						**07413500**
					GRA	CAT	
Lactobacillus fermentum		90.55	0.50	MAN 1	99	1	
Propionibacterium granulosum		3.95	0.21		99	90	
Fusobacterium mortiferum		3.47	0.24		1	5	
07413502	**Acceptable identification**						**07413502**
Lactobacillus fermentum		92.69	0.50	MAN 1			
Bifidobacterium adolescentis		3.60	0.20				
Bifidobacterium longum subsp. longum		1.07	0.09				
07413520	**Low discrimination**						**07413520**
					GRA		
Lactobacillus fermentum		80.69	0.41	MAN 1	99		
Bifidobacterium longum subsp. longum		11.13	0.20		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella melaninogenica		2.92	-.03		1			
07413522	**Low discrimination**							**07413522**
Lactobacillus fermentum		83.84	0.41	MAN 1				
Bifidobacterium longum subsp. longum		11.57	0.20					
Actinomyces naeslundii		2.73	0.26					
07414000	**Low discrimination**							**07414000**
					COC	GRA	CAT	
Fusobacterium mortiferum		53.32	0.64	XYL 7	1	1	5	
Propionibacterium granulosum		37.73	0.58		1	99	90	
Gemella morbillorum		5.69	0.37		99	99	1	
07414002	**Low discrimination**							**07414002**
					GRA	CAT		
Propionibacterium granulosum		53.03	0.47	XYL 1	99	90		
Fusobacterium mortiferum		33.30	0.47		1	5		
Lactobacillus fermentum		13.34	0.42		99	1		
07414010	**Low discrimination**							**07414010**
					COC	GRA	CAT	
Propionibacterium granulosum		89.62	0.43	XYL 1	1	99	90	
Fusobacterium mortiferum		9.38	0.29	MLZ 12	1	1	5	
Gemella morbillorum		1.00	0.03		99	99	1	
07414020	**Low discrimination**							**07414020**
					COC	CAT		
Propionibacterium granulosum		82.41	0.47	XYL 1	1	90		
Lactobacillus fermentum		6.91	0.34		1	1		
Gemella morbillorum		5.52	0.20		99	1		
07414022	**Low discrimination**							**07414022**
					GRA	CAT		
Propionibacterium granulosum		73.48	0.37	XYL 1	99	90		
Lactobacillus fermentum		24.65	0.34		99	1		
Fusobacterium mortiferum		1.86	0.13		1	5		
07414100	**Very good identification**							**07414100**
					GRA			
Fusobacterium mortiferum		95.52	0.70	XYL 7	1			
Lactobacillus fermentum		2.97	0.46		99			
Actinomyces meyeri		1.17	0.28		99			
07414101	**Good identification**							**07414101**
Fusobacterium mortiferum		100.0	0.36	XYL 7				
07414102	**Low discrimination**							**07414102**
					GRA	CAT		
Fusobacterium mortiferum		77.49	0.54	XYL 7	1	5		
Lactobacillus fermentum		21.71	0.46	BGL 10	99	1		
Propionibacterium granulosum		0.53	0.13		99	90		
07414110	**Good identification**							**07414110**
					GRA	CAT		
Fusobacterium mortiferum		96.03	0.36	XYL 7	1	5		
Propionibacterium granulosum		3.97	0.09	MLZ 1	99	90		
07414120	**Low discrimination**							**07414120**
					GRA			
Lactobacillus fermentum		44.07	0.37	GAL 99	99			
Fusobacterium mortiferum		42.90	0.36		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		7.07	0.13		99		
07414122	**Low discrimination**						**07414122**
					GRA	CAT	
Lactobacillus fermentum		55.79	0.37	GAL 99	99	1	
Parabacteroides distasonis		31.03	0.30		1	76	
Bacteroides fragilis		7.15	-.03		1	99	
07414140	**Good identification**						**07414140**
Fusobacterium mortiferum		100.0	0.36	XYL 7			
07414300	**Good identification**						**07414300**
Fusobacterium mortiferum		100.0	0.36	XYL 7			
07414322	**Good identification**						**07414322**
Parabacteroides distasonis		100.0	0.38	GAL 99			
07414400	**Low discrimination**						**07414400**
					COC	GRA	
Lactobacillus fermentum		59.05	0.42	GAL 99	1	99	
Gemella morbillorum		15.71	0.20		99	99	
Fusobacterium mortiferum		13.40	0.29		1	1	
07414402	**Acceptable identification**						**07414402**
					GRA	CAT	
Lactobacillus fermentum		93.87	0.42	GAL 99	99	1	
Propionibacterium granulosum		3.77	0.13		99	90	
Fusobacterium mortiferum		2.37	0.13		1	5	
07414500	**Low discrimination**						**07414500**
					GRA		
Lactobacillus fermentum		67.51	0.46	GAL 99	99		
Fusobacterium mortiferum		21.91	0.36		1		
Actinomyces meyeri		6.62	0.18		99		
07414502	**Acceptable identification**						**07414502**
					GRA		
Lactobacillus fermentum		91.34	0.46	GAL 99	99		
Bifidobacterium longum subsp. longum		5.36	0.17		99		
Fusobacterium mortiferum		3.29	0.19		1		
07414520	**Low discrimination**						**07414520**
					GRA		
Lactobacillus fermentum		46.88	0.37	GAL 99	99		
Bifidobacterium longum subsp. longum		33.04	0.28		99		
Bacteroides vulgatus		18.20	0.09		1		
07414522	**Low discrimination**						**07414522**
					GRA		
Lactobacillus fermentum		48.88	0.37	GAL 99	99		
Bifidobacterium longum subsp. longum		34.45	0.28		99		
Parabacteroides distasonis		16.67	0.27		1		
07415000	**Low discrimination**						**07415000**
					GRA	CAT	
Lactobacillus fermentum		65.88	0.76		99	1	
Fusobacterium mortiferum		22.42	0.67		1	5	
Propionibacterium granulosum		10.58	0.58		99	90	
07415001	**Low discrimination**						**07415001**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		66.63	0.42	URE 1	99	1	
Fusobacterium mortiferum		22.68	0.32		1	5	
Propionibacterium granulosum		10.70	0.24		99	90	
07415002	**Good identification**						**07415002**
					GRA	CAT	
Lactobacillus fermentum		92.77	0.76		99	1	
Propionibacterium granulosum		3.72	0.47		99	90	
Fusobacterium mortiferum		3.51	0.50		1	5	
07415003	**Acceptable identification**						**07415003**
					GRA	CAT	
Lactobacillus fermentum		92.77	0.42	URE 1	99	1	
Propionibacterium granulosum		3.72	0.13		99	90	
Fusobacterium mortiferum		3.51	0.16		1	5	
07415010	**Low discrimination**						**07415010**
					GRA	CAT	
Propionibacterium granulosum		61.79	0.43	XYL 1	99	90	
Lactobacillus fermentum		28.51	0.42	MLZ 12	99	1	
Fusobacterium mortiferum		9.70	0.32		1	5	
07415012	**Low discrimination**						**07415012**
					GRA	CAT	
Lactobacillus fermentum		63.30	0.42	MLZ 1	99	1	
Propionibacterium granulosum		34.31	0.33		99	90	
Fusobacterium mortiferum		2.39	0.16		1	5	
07415020	**Low discrimination**						**07415020**
					GRA	CAT	
Lactobacillus fermentum		87.94	0.68		99	1	
Propionibacterium granulosum		10.59	0.47		99	90	
Fusobacterium mortiferum		0.91	0.32		1	5	
07415022	**Very good identification**						**07415022**
					GRA	CAT	
Lactobacillus fermentum		95.79	0.68		99	1	
Propionibacterium granulosum		2.88	0.37		99	90	
Parabacteroides distasonis		0.88	0.30		1	76	
07415040	**Low discrimination**						**07415040**
					GRA	CAT	
Lactobacillus fermentum		66.63	0.42	SOR 1	99	1	
Fusobacterium mortiferum		22.68	0.32		1	5	
Propionibacterium granulosum		10.70	0.24		99	90	
07415042	**Acceptable identification**						**07415042**
					GRA	CAT	
Lactobacillus fermentum		92.77	0.42	SOR 1	99	1	
Propionibacterium granulosum		3.72	0.13		99	90	
Fusobacterium mortiferum		3.51	0.16		1	5	
07415100	**Low discrimination**						**07415100**
					GRA		
Lactobacillus fermentum		66.91	0.80		99		
Fusobacterium mortiferum		32.57	0.73		1		
Actinomyces meyeri		0.40	0.31		99		
07415101	**Low discrimination**						**07415101**
					GRA		
Lactobacillus fermentum		66.62	0.46	URE 1	99		
Fusobacterium mortiferum		32.43	0.39		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.95	0.24		99		
07415102	**Good identification**						**07415102**
					GRA		
Lactobacillus fermentum		94.46	0.80		99		
Fusobacterium mortiferum		5.11	0.57		1		
Parabacteroides distasonis		0.18	0.30		1		
07415103	**Acceptable identification**						**07415103**
					GRA		
Lactobacillus fermentum		93.60	0.46	URE 1	99		
Fusobacterium mortiferum		5.06	0.22		1		
Actinomyces naeslundii		1.34	0.24		99		
07415110	**Low discrimination**						**07415110**
					GRA		
Lactobacillus fermentum		62.49	0.46	MLZ 1	99		
Fusobacterium mortiferum		30.42	0.39		1		
Bifidobacterium longum subsp. longum		6.25	0.21		99		
07415112	**Low discrimination**						**07415112**
					GRA		
Lactobacillus fermentum		73.93	0.46	MLZ 1	99		
Bifidobacterium longum subsp. longum		7.39	0.21		99		
Parabacteroides distasonis		7.38	0.26		1		
07415120	**Low discrimination**						**07415120**
					GRA		
Lactobacillus fermentum		88.33	0.72		99		
Prevotella melaninogenica		5.44	0.32		1		
Bacteroides vulgatus		2.58	0.24		1		
07415121	**Acceptable identification**						**07415121**
					GRA		
Lactobacillus fermentum		90.55	0.37	URE 1	99		
Prevotella melaninogenica		5.58	-.03		1		
Actinomyces naeslundii		3.88	0.24		99		
07415122	**Low discrimination**						**07415122**
					GRA	CAT	
Lactobacillus fermentum		57.94	0.72		99	1	
Parabacteroides distasonis		32.23	0.64		1	76	
Bacteroides fragilis		7.43	0.32		1	99	
07415123	**Low discrimination**						**07415123**
					GRA	CAT	
Lactobacillus fermentum		57.90	0.37	URE 1	99	1	
Parabacteroides distasonis		32.20	0.30		1	76	
Bacteroides fragilis		7.42	-.03		1	99	
07415132	**Good identification**						**07415132**
					GRA		
Parabacteroides distasonis		92.52	0.60		1		
Bifidobacterium longum subsp. longum		3.74	0.32		99		
Lactobacillus fermentum		3.12	0.37		99		
07415140	**Low discrimination**						**07415140**
					GRA		
Lactobacillus fermentum		67.26	0.46	SOR 1	99		
Fusobacterium mortiferum		32.74	0.39		1		
07415142	**Acceptable identification**						**07415142**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		94.87	0.46	SOR 1	99		
Fusobacterium mortiferum		5.13	0.22		1		
07415160	**Acceptable identification**						**07415160**
					GRA		
Lactobacillus fermentum		90.20	0.37	SOR 1	99		
Prevotella melaninogenica		5.55	-.03		1		
Clostridium perfringens		4.25	0.02		99		
07415162	**Low discrimination**						**07415162**
					GRA	CAT	
Lactobacillus fermentum		57.07	0.37	SOR 1	99	1	
Parabacteroides distasonis		31.74	0.30		1	76	
Bacteroides fragilis		7.32	-.03		1	99	
07415200	**Low discrimination**						**07415200**
					GRA	CAT	
Lactobacillus fermentum		54.59	0.42	RHA 1	99	1	
Propionibacterium granulosum		26.84	0.32		99	90	
Fusobacterium mortiferum		18.58	0.32		1	5	
07415202	**Low discrimination**						**07415202**
					GRA	CAT	
Lactobacillus fermentum		86.15	0.42	RHA 1	99	1	
Propionibacterium granulosum		10.59	0.22		99	90	
Fusobacterium mortiferum		3.26	0.16		1	5	
07415222	**Low discrimination**						**07415222**
					GRA	CAT	
Parabacteroides distasonis		70.30	0.38	LAC 99	1	76	
Lactobacillus fermentum		25.82	0.34		99	1	
Propionibacterium granulosum		2.38	0.11		99	90	
07415300	**Low discrimination**						**07415300**
					GRA		
Lactobacillus fermentum		63.52	0.46	RHA 1	99		
Fusobacterium mortiferum		30.92	0.39		1		
Bacteroides vulgatus		5.57	0.06		1		
07415302	**Low discrimination**						**07415302**
					GRA		
Lactobacillus fermentum		62.10	0.46	RHA 1	99		
Parabacteroides distasonis		34.54	0.38		1		
Fusobacterium mortiferum		3.36	0.22		1		
07415320	**Acceptable identification**						**07415320**
					GRA		
Bacteroides vulgatus		90.51	0.40	MLT 99	1		
Parabacteroides distasonis		5.80	0.38	ARA 93	1		
Lactobacillus fermentum		3.48	0.37		99		
07415322	**Very good identification**						**07415322**
					GRA		
Parabacteroides distasonis		98.83	0.72		1		
Lactobacillus fermentum		0.60	0.37		99		
Prevotella oralis		0.33	0.17		1		
07415323	**Good identification**						**07415323**
Parabacteroides distasonis		100.0	0.38	URE 1			
07415332	**Very good identification**						**07415332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.68				
07415362	**Good identification**						**07415362**
Parabacteroides distasonis		100.0	0.38	SOR 1			
07415400	**Very good identification**						**07415400**
					GRA	CAT	
Lactobacillus fermentum		99.20	0.76		99	1	
Fusobacterium mortiferum		0.34	0.32		1	5	
Propionibacterium granulosum		0.16	0.24		99	90	
07415401	**Good identification**						**07415401**
Lactobacillus fermentum		100.0	0.42	URE 1			
07415402	**Very good identification**						**07415402**
					GRA		
Lactobacillus fermentum		99.83	0.76		99		
Bifidobacterium longum subsp. longum		0.10	0.17		99		
Fusobacterium mortiferum		0.04	0.16		1		
07415403	**Good identification**						**07415403**
Lactobacillus fermentum		100.0	0.42	URE 1			
07415410	**Low discrimination**						**07415410**
					CAT		
Lactobacillus fermentum		84.39	0.42	MLZ 1	1		
Bifidobacterium longum subsp. longum		13.77	0.21		1		
Propionibacterium granulosum		1.85	0.09		90		
07415412	**Low discrimination**						**07415412**
Lactobacillus fermentum		85.97	0.42	MLZ 1			
Bifidobacterium longum subsp. longum		14.03	0.21				
07415420	**Very good identification**						**07415420**
					GRA		
Lactobacillus fermentum		98.11	0.68		99		
Bifidobacterium longum subsp. longum		1.14	0.28		99		
Bacteroides vulgatus		0.63	0.09		1		
07415422	**Very good identification**						**07415422**
					GRA		
Lactobacillus fermentum		98.31	0.68		99		
Bifidobacterium longum subsp. longum		1.14	0.28		99		
Parabacteroides distasonis		0.55	0.27		1		
07415440	**Good identification**						**07415440**
Lactobacillus fermentum		100.0	0.42	SOR 1			
07415442	**Good identification**						**07415442**
Lactobacillus fermentum		100.0	0.42	SOR 1			
07415500	**Good identification**						**07415500**
					GRA		
Lactobacillus fermentum		93.77	0.80		99		
Bifidobacterium longum subsp. longum		5.51	0.52		99		
Fusobacterium mortiferum		0.46	0.39		1		
07415501	**Acceptable identification**						**07415501**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		94.45	0.46	URE 1		
Bifidobacterium longum subsp. longum		5.55	0.17			
07415502	**Good identification**					**07415502**
Lactobacillus fermentum		93.82	0.80			
Bifidobacterium longum subsp. longum		5.51	0.52			
Bifidobacterium adolescentis		0.51	0.36			
07415503	**Acceptable identification**					**07415503**
Lactobacillus fermentum		94.45	0.46	URE 1		
Bifidobacterium longum subsp. longum		5.55	0.17			
07415510	**Acceptable identification**					**07415510**
Bifidobacterium longum subsp. longum		90.83	0.56	MLT 99		
Lactobacillus fermentum		9.18	0.46			
07415512	**Low discrimination**					**07415512**
Bifidobacterium longum subsp. longum		85.18	0.56	MLT 99		
Lactobacillus fermentum		8.61	0.46			
Bifidobacterium adolescentis		5.69	0.37			
07415520	**Low discrimination**					**07415520**
					GRA	
Lactobacillus fermentum		47.67	0.72		99	
Bifidobacterium longum subsp. longum		33.60	0.62		99	
Bacteroides vulgatus		18.51	0.43		1	
07415521	**Low discrimination**					**07415521**
					GRA	
Lactobacillus fermentum		47.78	0.37	URE 1	99	
Bifidobacterium longum subsp. longum		33.67	0.28		99	
Bacteroides vulgatus		18.55	0.09		1	
07415522	**Low discrimination**					**07415522**
					GRA	
Lactobacillus fermentum		48.74	0.72		99	
Bifidobacterium longum subsp. longum		34.34	0.62		99	
Parabacteroides distasonis		16.61	0.61		1	
07415523	**Low discrimination**					**07415523**
					GRA	
Lactobacillus fermentum		48.88	0.37	URE 1	99	
Bifidobacterium longum subsp. longum		34.45	0.28		99	
Parabacteroides distasonis		16.67	0.27		1	
07415530	**Very good identification**					**07415530**
					GRA	
Bifidobacterium longum subsp. longum		98.69	0.66	MLT 99	99	
Lactobacillus fermentum		0.83	0.37		99	
Bacteroides vulgatus		0.32	0.09		1	
07415532	**Low discrimination**					**07415532**
					GRA	
Bifidobacterium longum subsp. longum		86.10	0.66	MLT 99	99	
Parabacteroides distasonis		13.17	0.56		1	
Lactobacillus fermentum		0.73	0.37		99	
07415540	**Acceptable identification**					**07415540**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		94.45	0.46	SOR 1				
Bifidobacterium longum subsp. longum		5.55	0.17					
07415542	**Low discrimination**							**07415542**
Lactobacillus fermentum		88.84	0.46	SOR 1				
Bifidobacterium adolescentis		5.94	0.20					
Bifidobacterium longum subsp. longum		5.22	0.17					
07415560	**Low discrimination**							**07415560**
					GRA			
Lactobacillus fermentum		47.78	0.37	SOR 1	99			
Bifidobacterium longum subsp. longum		33.67	0.28		99			
Bacteroides vulgatus		18.55	0.09		1			
07415562	**Low discrimination**							**07415562**
					GRA			
Lactobacillus fermentum		48.88	0.37	SOR 1	99			
Bifidobacterium longum subsp. longum		34.45	0.28		99			
Parabacteroides distasonis		16.67	0.27		1			
07415600	**Good identification**							**07415600**
					GRA			
Lactobacillus fermentum		98.12	0.42	RHA 1	99			
Bacteroides vulgatus		1.88	-.09		1			
07415602	**Good identification**							**07415602**
Lactobacillus fermentum		100.0	0.42	RHA 1				
07415702	**Low discrimination**							**07415702**
					GRA			
Lactobacillus fermentum		69.97	0.46	RHA 1	99			
Parabacteroides distasonis		23.86	0.35		1			
Bifidobacterium longum subsp. longum		4.11	0.17		99			
07415720	**Very good identification**							**07415720**
					GRA			
Bacteroides vulgatus		99.22	0.60	MLT 99	1			
Parabacteroides distasonis		0.29	0.35		1			
Lactobacillus fermentum		0.29	0.37		99			
07415722	**Good identification**							**07415722**
					GRA			
Parabacteroides distasonis		94.97	0.69		1			
Bacteroides vulgatus		3.28	0.25		1			
Lactobacillus fermentum		0.94	0.37		99			
07415732	**Very good identification**							**07415732**
					GRA			
Parabacteroides distasonis		97.28	0.64		1			
Bifidobacterium longum subsp. longum		2.14	0.32		99			
Clostridium clostridioforme		0.43	0.18		50			
07416000	**Good identification**							**07416000**
					COC	GRA	CAT	
Propionibacterium granulosum		97.77	0.56	XYL 1	1	99	90	
Fusobacterium mortiferum		1.85	0.29		1	1	5	
Gemella morbillorum		0.20	0.03		99	99	1	
07416002	**Good identification**							**07416002**
					GRA	CAT		
Propionibacterium granulosum		99.17	0.45	XYL 1	99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.83	0.13		1	5	
07416010	**Good identification**						**07416010**
					SPO	CAT	
Propionibacterium granulosum		97.27	0.41	XYL 1	1	90	
Clostridium difficile		2.73	0.02	MLZ 12	99	1	
07416020	**Good identification**						**07416020**
Propionibacterium granulosum		100.0	0.45	XYL 1			
07416022	**Good identification**						**07416022**
Propionibacterium granulosum		100.0	0.35	XYL 1			
07416100	**Low discrimination**						**07416100**
					GRA	CAT	
Fusobacterium mortiferum		81.38	0.36	MAN 1	1	5	
Propionibacterium granulosum		18.62	0.21	XYL 7	99	90	
07417000	**Low discrimination**						**07417000**
					GRA	CAT	
Propionibacterium granulosum		89.95	0.56	XYL 1	99	90	
Lactobacillus fermentum		7.50	0.42		99	1	
Fusobacterium mortiferum		2.55	0.32		1	5	
07417002	**Low discrimination**						**07417002**
					GRA	CAT	
Propionibacterium granulosum		74.29	0.45	XYL 1	99	90	
Lactobacillus fermentum		24.78	0.42		99	1	
Fusobacterium mortiferum		0.94	0.16		1	5	
07417010	**Good identification**						**07417010**
Propionibacterium granulosum		100.0	0.41	XYL 1			
07417020	**Low discrimination**						**07417020**
					CAT		
Propionibacterium granulosum		90.00	0.45	XYL 1	90		
Lactobacillus fermentum		10.01	0.34		1		
07417022	**Low discrimination**						**07417022**
					CAT		
Propionibacterium granulosum		69.22	0.35	XYL 1	90		
Lactobacillus fermentum		30.78	0.34		1		
07417100	**Low discrimination**						**07417100**
					GRA	CAT	
Lactobacillus fermentum		63.31	0.46	MAN 1	99	1	
Fusobacterium mortiferum		30.82	0.39		1	5	
Propionibacterium granulosum		4.70	0.21		99	90	
07417102	**Low discrimination**						**07417102**
					GRA		
Lactobacillus fermentum		84.64	0.46	MAN 1	99		
Bifidobacterium adolescentis		5.60	0.20		99		
Fusobacterium mortiferum		4.58	0.22		1		
07417120	**Low discrimination**						**07417120**
					GRA	CAT	
Lactobacillus fermentum		85.27	0.37	MAN 1	99	1	
Prevotella melaninogenica		5.25	-.03		1	1	
Propionibacterium granulosum		4.75	0.11		99	90	
07417122	**Low discrimination**						**07417122**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Lactobacillus fermentum		57.48	0.37	MAN 1	99	1		
Parabacteroides distasonis		31.97	0.30		1	76		
Bacteroides fragilis		7.37	-.03		1	99		
07417322	**Good identification**							**07417322**
Parabacteroides distasonis		100.0	0.38	MAN 1				
07417400	**Low discrimination**							**07417400**
					CAT			
Lactobacillus fermentum		89.20	0.42	MAN 1	1			
Propionibacterium granulosum		10.81	0.21		90			
07417402	**Good identification**							**07417402**
					CAT			
Lactobacillus fermentum		97.06	0.42	MAN 1	1			
Propionibacterium granulosum		2.94	0.11		90			
07417500	**Acceptable identification**							**07417500**
Lactobacillus fermentum		94.45	0.46	MAN 1				
Bifidobacterium longum subsp. longum		5.55	0.17					
07417502	**Low discrimination**							**07417502**
Lactobacillus fermentum		66.61	0.46	MAN 1				
Bifidobacterium adolescentis		29.48	0.34					
Bifidobacterium longum subsp. longum		3.91	0.17					
07417512	**Low discrimination**							**07417512**
Bifidobacterium adolescentis		84.40	0.36	MLT 99				
Bifidobacterium longum subsp. longum		15.60	0.21					
07417520	**Low discrimination**							**07417520**
					GRA			
Lactobacillus fermentum		47.78	0.37	MAN 1	99			
Bifidobacterium longum subsp. longum		33.67	0.28		99			
Bacteroides vulgatus		18.55	0.09		1			
07417522	**Low discrimination**							**07417522**
					GRA			
Lactobacillus fermentum		48.88	0.37	MAN 1	99			
Bifidobacterium longum subsp. longum		34.45	0.28		99			
Parabacteroides distasonis		16.67	0.27		1			
07420000	**Very good identification**							**07420000**
					COC	GRA	CAT	
Propionibacterium granulosum		97.52	0.66	FRU 99	1	99	90	
Fusobacterium mortiferum		1.84	0.40		1	1	5	
Gemella morbillorum		0.36	0.17		99	99	1	
07420002	**Good identification**							**07420002**
					GRA	CAT		
Propionibacterium granulosum		99.17	0.56	FRU 99	99	90		
Fusobacterium mortiferum		0.83	0.23		1	5		
07420010	**Good identification**							**07420010**
Propionibacterium granulosum		100.0	0.51	FRU 99				
07420012	**Good identification**							**07420012**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.41	FRU 99				
07420020	**Good identification**							**07420020**
					COC	CAT		
Propionibacterium granulosum		99.83	0.56	FRU 99	1	90		
Gemella morbillorum		0.17	0.01		99	1		
07420022	**Good identification**							**07420022**
Propionibacterium granulosum		100.0	0.45	FRU 99				
07420030	**Good identification**							**07420030**
Propionibacterium granulosum		100.0	0.41	FRU 99				
07420100	**Low discrimination**							**07420100**
					GRA	CAT		
Fusobacterium mortiferum		81.35	0.46	FRU 99	1	5		
Propionibacterium granulosum		18.65	0.32		99	90		
07420200	**Good identification**							**07420200**
Propionibacterium granulosum		100.0	0.40	FRU 99				
07421000	**Very good identification**							**07421000**
					COC	GRA	CAT	
Propionibacterium granulosum		96.95	0.66	FRU 99	1	99	90	
Fusobacterium mortiferum		2.75	0.43		1	1	5	
Gemella morbillorum		0.20	0.13		99	99	1	
07421002	**Good identification**							**07421002**
					GRA	CAT		
Propionibacterium granulosum		98.31	0.56	FRU 99	99	90		
Fusobacterium mortiferum		1.24	0.26		1	5		
Actinomyces naeslundii		0.46	0.30		99	25		
07421010	**Good identification**							**07421010**
Propionibacterium granulosum		100.0	0.51	FRU 99				
07421012	**Good identification**							**07421012**
Propionibacterium granulosum		100.0	0.41	FRU 99				
07421020	**Good identification**							**07421020**
					CAT			
Propionibacterium granulosum		99.26	0.56	FRU 99	90			
Actinomyces naeslundii		0.46	0.30		25			
Clostridium perfringens		0.28	0.05		1			
07421022	**Good identification**							**07421022**
Propionibacterium granulosum		98.18	0.45	FRU 99				
Actinomyces naeslundii		1.82	0.30					
07421030	**Good identification**							**07421030**
Propionibacterium granulosum		100.0	0.41	FRU 99				
07421100	**Low discrimination**							**07421100**
					GRA	CAT		
Fusobacterium mortiferum		82.54	0.49	FRU 99	1	5		
Propionibacterium granulosum		12.62	0.32		99	90		
Actinomyces naeslundii		4.11	0.38		99	25		
07421101	**Low discrimination**							**07421101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Actinomyces naeslundii		79.03	0.41	MLT 94	99	1	
Fusobacterium mortiferum		10.68	0.15	FRU 95	1	1	
Clostridium chauvoei		10.29	0.03		99	99	
07421103	**Good identification**						**07421103**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07421121	**Good identification**						**07421121**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07421122	**Low discrimination**						**07421122**
					GRA		
Actinomyces naeslundii		29.75	0.38	MLT 94	99		
Parabacteroides distasonis		29.00	0.25	FRU 95	1		
Clostridium perfringens		22.27	0.14		99		
07421123	**Good identification**						**07421123**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07421200	**Good identification**						**07421200**
Propionibacterium granulosum		100.0	0.40	FRU 99			
07422000	**Very good identification**						**07422000**
Propionibacterium granulosum		100.0	0.64	FRU 99			
07422002	**Good identification**						**07422002**
Propionibacterium granulosum		100.0	0.53	FRU 99			
07422010	**Good identification**						**07422010**
Propionibacterium granulosum		100.0	0.49	FRU 99			
07422012	**Good identification**						**07422012**
Propionibacterium granulosum		100.0	0.39	FRU 99			
07422020	**Good identification**						**07422020**
Propionibacterium granulosum		100.0	0.53	FRU 99			
07422022	**Good identification**						**07422022**
Propionibacterium granulosum		100.0	0.43	FRU 99			
07422030	**Good identification**						**07422030**
Propionibacterium granulosum		100.0	0.39	FRU 99			
07422200	**Good identification**						**07422200**
Propionibacterium granulosum		100.0	0.38	FRU 99			
07423000	**Very good identification**						**07423000**
Propionibacterium granulosum		99.70	0.64	FRU 99			
Actinomyces naeslundii		0.30	0.35				
07423002	**Good identification**						**07423002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		98.82	0.53	FRU 99		
Actinomyces naeslundii		1.18	0.35			
07423003	**Low discrimination**					**07423003**
Actinomyces naeslundii		63.89	0.38	MLT 94		
Propionibacterium granulosum		36.11	0.19	FRU 95		
07423010	**Good identification**					**07423010**
Propionibacterium granulosum		100.0	0.49	FRU 99		
07423012	**Good identification**					**07423012**
Propionibacterium granulosum		100.0	0.39	FRU 99		
07423020	**Good identification**					**07423020**
Propionibacterium granulosum		98.82	0.53	FRU 99		
Actinomyces naeslundii		1.18	0.35			
07423021	**Low discrimination**					**07423021**
Actinomyces naeslundii		63.89	0.38	MLT 94		
Propionibacterium granulosum		36.11	0.19	FRU 95		
07423022	**Good identification**					**07423022**
Propionibacterium granulosum		95.45	0.43	FRU 99		
Actinomyces naeslundii		4.55	0.35			
07423023	**Low discrimination**					**07423023**
Actinomyces naeslundii		87.62	0.38	MLT 94		
Propionibacterium granulosum		12.38	0.09	FRU 95		
07423030	**Good identification**					**07423030**
Propionibacterium granulosum		100.0	0.39	FRU 99		
07423101	**Good identification**					**07423101**
Actinomyces naeslundii		100.0	0.46	MLT 94		
07423102	**Low discrimination**					**07423102**
Actinomyces naeslundii		77.05	0.43	MLT 94		
Propionibacterium granulosum		22.95	0.19	FRU 95		
07423103	**Good identification**					**07423103**
Actinomyces naeslundii		100.0	0.46	MLT 94		
07423120	**Low discrimination**					**07423120**
Actinomyces naeslundii		77.05	0.43	MLT 94		
Propionibacterium granulosum		22.95	0.19	FRU 95		
07423121	**Good identification**					**07423121**
Actinomyces naeslundii		100.0	0.46	MLT 94		
07423122	**Acceptable identification**					**07423122**
Actinomyces naeslundii		93.07	0.43	MLT 94		
Propionibacterium granulosum		6.93	0.09	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07423123	**Good identification**						**07423123**
Actinomyces naeslundii		100.0	0.46	MLT 94			
07423200	**Good identification**						**07423200**
Propionibacterium granulosum		100.0	0.38	FRU 99			
07425122	**Good identification**						**07425122**
Parabacteroides distasonis		100.0	0.41	FRU 99			
07425132	**Good identification**						**07425132**
Parabacteroides distasonis		100.0	0.37	FRU 99			
07425322	**Good identification**						**07425322**
Parabacteroides distasonis		100.0	0.50	FRU 99			
07425332	**Good identification**						**07425332**
Parabacteroides distasonis		100.0	0.45	FRU 99			
07425520	**Good identification**						**07425520**
Bifidobacterium longum subsp. longum		100.0	0.35	MLT 99			
07425522	**Low discrimination**						**07425522**
					GRA		
Bifidobacterium longum subsp. longum		55.51	0.35	MLT 99	99		
Parabacteroides distasonis		44.49	0.38	FRU 90	1		
07425530	**Good identification**						**07425530**
Bifidobacterium longum subsp. longum		100.0	0.39	MLT 99			
07425532	**Low discrimination**						**07425532**
					GRA		
Bifidobacterium longum subsp. longum		79.78	0.39	MLT 99	99		
Parabacteroides distasonis		20.22	0.33	FRU 90	1		
07425722	**Good identification**						**07425722**
Parabacteroides distasonis		100.0	0.46	FRU 99			
07425732	**Good identification**						**07425732**
Parabacteroides distasonis		100.0	0.41	FRU 99			
07430000	**Very good identification**						**07430000**
					GRA	CAT	
Propionibacterium granulosum		98.14	1.00		99	90	
Fusobacterium mortiferum		1.85	0.74		1	5	
Actinomyces naeslundii		0.00	0.29		99	25	
07430001	**Very good identification**						**07430001**
					GRA	CAT	
Propionibacterium granulosum		98.02	0.66	URE 1	99	90	
Fusobacterium mortiferum		1.85	0.40		1	5	
Actinomyces naeslundii		0.14	0.32		99	25	
07430002	**Very good identification**						**07430002**
					GRA	CAT	
Propionibacterium granulosum		99.16	0.90		99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		0.83	0.57		1	5		
Actinomyces viscosus		0.01	0.08		99	76		
07430003	**Good identification**							**07430003**
					GRA	CAT		
Propionibacterium granulosum		98.44	0.56	URE 1	99	90		
Fusobacterium mortiferum		0.83	0.23		1	5		
Actinomyces naeslundii		0.54	0.32		99	25		
07430010	**Very good identification**							**07430010**
					GRA	CAT		
Propionibacterium granulosum		99.86	0.85	MLZ 12	99	90		
Fusobacterium mortiferum		0.14	0.40		1	5		
07430011	**Good identification**							**07430011**
Propionibacterium granulosum		100.0	0.51	MLZ 12				
07430012	**Very good identification**							**07430012**
					GRA	CAT		
Propionibacterium granulosum		99.94	0.75	MLZ 12	99	90		
Fusobacterium mortiferum		0.06	0.23		1	5		
07430013	**Good identification**							**07430013**
Propionibacterium granulosum		100.0	0.41	MLZ 12				
07430020	**Very good identification**							**07430020**
					GRA	CAT		
Propionibacterium granulosum		99.92	0.90		99	90		
Fusobacterium mortiferum		0.08	0.40		1	5		
Actinomyces naeslundii		0.00	0.29		99	25		
07430021	**Good identification**							**07430021**
Propionibacterium granulosum		99.45	0.56	URE 1				
Actinomyces naeslundii		0.55	0.32					
07430022	**Very good identification**							**07430022**
					GRA	CAT		
Propionibacterium granulosum		99.95	0.79		99	90		
Fusobacterium mortiferum		0.03	0.23		1	5		
Actinomyces naeslundii		0.02	0.29		99	25		
07430023	**Good identification**							**07430023**
Propionibacterium granulosum		97.85	0.45	URE 1				
Actinomyces naeslundii		2.15	0.32					
07430030	**Very good identification**							**07430030**
Propionibacterium granulosum		100.0	0.75	MLZ 12				
07430031	**Good identification**							**07430031**
Propionibacterium granulosum		100.0	0.41	MLZ 12				
07430032	**Very good identification**							**07430032**
Propionibacterium granulosum		100.0	0.65	MLZ 12				
07430040	**Good identification**							**07430040**
					GRA	SPO	CAT	
Propionibacterium granulosum		93.08	0.66	SOR 1	99	1	90	
Clostridium botulinum biovar B		5.17	0.32		99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		1.76	0.40		1	1	5	
07430042	**Good identification**							**07430042**
					GRA	SPO	CAT	
Propionibacterium granulosum		98.95	0.56	SOR 1	99	1	90	
Fusobacterium mortiferum		0.83	0.23		1	1	5	
Clostridium botulinum biovar B		0.22	-.03		99	99	1	
07430050	**Low discrimination**							**07430050**
					SPO	CAT		
Propionibacterium granulosum		83.30	0.51	MLZ 12	1	90		
Clostridium botulinum biovar B		16.70	0.26	SOR 1	99	1		
07430052	**Good identification**							**07430052**
Propionibacterium granulosum		100.0	0.41	MLZ 12				
07430060	**Acceptable identification**							**07430060**
					SPO	CAT		
Propionibacterium granulosum		90.15	0.56	SOR 1	1	90		
Clostridium botulinum biovar B		9.86	0.26		99	1		
07430062	**Good identification**							**07430062**
Propionibacterium granulosum		100.0	0.45	SOR 1				
07430070	**Low discrimination**							**07430070**
					SPO	CAT		
Propionibacterium granulosum		71.69	0.41	MLZ 12	1	90		
Clostridium botulinum biovar B		28.31	0.21	SOR 1	99	1		
07430100	**Low discrimination**							**07430100**
					GRA	CAT		
Fusobacterium mortiferum		81.30	0.80		1	5		
Propionibacterium granulosum		18.64	0.66		99	90		
Actinomyces naeslundii		0.05	0.37		99	25		
07430101	**Low discrimination**							**07430101**
					GRA	CAT		
Fusobacterium mortiferum		75.87	0.46	URE 1	1	5		
Propionibacterium granulosum		17.40	0.32		99	90		
Actinomyces naeslundii		6.74	0.40		99	25		
07430102	**Low discrimination**							**07430102**
					GRA	CAT		
Fusobacterium mortiferum		65.45	0.64	BGL 10	1	5		
Propionibacterium granulosum		33.76	0.56		99	90		
Actinomyces viscosus		0.36	0.12		99	76		
07430110	**Low discrimination**							**07430110**
					GRA	CAT		
Propionibacterium granulosum		75.58	0.51	LAC 1	99	90		
Fusobacterium mortiferum		24.42	0.46	MLZ 12	1	5		
07430112	**Low discrimination**							**07430112**
					GRA	CAT		
Propionibacterium granulosum		87.45	0.41	LAC 1	99	90		
Fusobacterium mortiferum		12.56	0.30	MLZ 12	1	5		
07430120	**Low discrimination**							**07430120**
					GRA	CAT		
Propionibacterium granulosum		82.62	0.56	LAC 1	99	90		
Fusobacterium mortiferum		14.56	0.46		1	5		
Clostridium perfringens		1.73	0.19		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07430121	**Low discrimination**							**07430121**
					GRA	CAT		
Actinomyces naeslundii		56.84	0.40	MLT 94	99	25		
Propionibacterium granulosum		36.69	0.21	GAL 96	99	90		
Fusobacterium mortiferum		6.47	0.12		1	5		
07430122	**Low discrimination**							**07430122**
					GRA	CAT		
Propionibacterium granulosum		82.59	0.45	LAC 1	99	90		
Fusobacterium mortiferum		6.47	0.30		1	5		
Parabacteroides distasonis		4.24	0.25		1	76		
07430123	**Low discrimination**							**07430123**
Actinomyces naeslundii		86.11	0.40	MLT 94				
Propionibacterium granulosum		13.90	0.11	GAL 96				
07430130	**Good identification**							**07430130**
					GRA	CAT		
Propionibacterium granulosum		98.71	0.41	LAC 1	99	90		
Fusobacterium mortiferum		1.29	0.12	MLZ 12	1	5		
07430140	**Low discrimination**							**07430140**
					GRA	SPO	CAT	
Fusobacterium mortiferum		79.84	0.46	SOR 1	1	1	5	
Propionibacterium granulosum		18.31	0.32		99	1	90	
Clostridium botulinum biovar B		1.02	-.03		99	99	1	
07430200	**Very good identification**							**07430200**
					GRA	CAT		
Propionibacterium granulosum		99.39	0.74	RHA 3	99	90		
Fusobacterium mortiferum		0.61	0.40		1	5		
07430201	**Good identification**							**07430201**
Propionibacterium granulosum		100.0	0.40	RHA 3				
07430202	**Very good identification**							**07430202**
					GRA	CAT		
Propionibacterium granulosum		99.73	0.64	RHA 3	99	90		
Fusobacterium mortiferum		0.27	0.23		1	5		
07430210	**Very good identification**							**07430210**
Propionibacterium granulosum		100.0	0.59	RHA 3				
07430212	**Good identification**							**07430212**
Propionibacterium granulosum		100.0	0.49	RHA 3				
07430220	**Very good identification**							**07430220**
Propionibacterium granulosum		100.0	0.64	RHA 3				
07430222	**Good identification**							**07430222**
Propionibacterium granulosum		100.0	0.54	RHA 3				
07430230	**Good identification**							**07430230**
Propionibacterium granulosum		100.0	0.49	RHA 3				
07430232	**Good identification**							**07430232**
Propionibacterium granulosum		100.0	0.39	RHA 3				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07430240	**Good identification**							**07430240**
					SPO	CAT		
Propionibacterium granulosum		98.22	0.40	RHA 3	1	90		
Clostridium botulinum biovar B		1.78	-.03	SOR 1	99	1		
07430300	**Low discrimination**							**07430300**
					GRA	CAT		
Fusobacterium mortiferum		58.75	0.46	RHA 1	1	5		
Propionibacterium granulosum		41.25	0.40		99	90		
07430400	**Very good identification**							**07430400**
					COC	GRA	CAT	
Propionibacterium granulosum		98.10	0.66	ARA 1	1	99	90	
Fusobacterium mortiferum		1.85	0.40		1	1	5	
Gemella morbillorum		0.05	0.03		99	99	1	
07430402	**Good identification**							**07430402**
					GRA	CAT		
Propionibacterium granulosum		99.17	0.56	ARA 1	99	90		
Fusobacterium mortiferum		0.83	0.23		1	5		
07430410	**Good identification**							**07430410**
Propionibacterium granulosum		100.0	0.51	ARA 1				
07430412	**Good identification**							**07430412**
Propionibacterium granulosum		100.0	0.41	ARA 1				
07430420	**Good identification**							**07430420**
Propionibacterium granulosum		100.0	0.56	ARA 1				
07430422	**Good identification**							**07430422**
Propionibacterium granulosum		100.0	0.45	ARA 1				
07430430	**Good identification**							**07430430**
Propionibacterium granulosum		100.0	0.41	ARA 1				
07430500	**Low discrimination**							**07430500**
					GRA	CAT		
Fusobacterium mortiferum		80.47	0.46	ARA 1	1	5		
Propionibacterium granulosum		18.45	0.32		99	90		
Lactobacillus fermentum		1.08	0.15		99	1		
07430600	**Good identification**							**07430600**
Propionibacterium granulosum		100.0	0.40	RHA 3				
07431000	**Very good identification**							**07431000**
					GRA	CAT		
Propionibacterium granulosum		97.18	1.00		99	90		
Fusobacterium mortiferum		2.75	0.77		1	5		
Lactobacillus fermentum		0.04	0.46		99	1		
07431001	**Good identification**							**07431001**
					GRA	CAT		
Propionibacterium granulosum		94.22	0.66	URE 1	99	90		
Actinomyces naeslundii		3.11	0.55		99	25		
Fusobacterium mortiferum		2.67	0.43		1	5		
07431002	**Very good identification**							**07431002**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		98.52	0.90		99	90	
Fusobacterium mortiferum		1.24	0.60		1	5	
Lactobacillus fermentum		0.14	0.46		99	1	
07431003	**Low discrimination**						**07431003**
					GRA	CAT	
Propionibacterium granulosum		87.37	0.56	URE 1	99	90	
Actinomyces naeslundii		11.53	0.55		99	25	
Fusobacterium mortiferum		1.10	0.26		1	5	
07431010	**Very good identification**						**07431010**
					GRA	CAT	
Propionibacterium granulosum		99.78	0.85	MLZ 12	99	90	
Fusobacterium mortiferum		0.21	0.43		1	5	
Actinomyces naeslundii		0.01	0.30		99	25	
07431011	**Good identification**						**07431011**
Propionibacterium granulosum		98.74	0.51	MLZ 12			
Actinomyces naeslundii		1.26	0.33	URE 1			
07431012	**Very good identification**						**07431012**
					GRA	CAT	
Propionibacterium granulosum		99.87	0.75	MLZ 12	99	90	
Fusobacterium mortiferum		0.09	0.26		1	5	
Actinomyces naeslundii		0.03	0.30		99	25	
07431013	**Good identification**						**07431013**
Propionibacterium granulosum		95.16	0.41	MLZ 12			
Actinomyces naeslundii		4.85	0.33	URE 1			
07431020	**Very good identification**						**07431020**
					GRA	CAT	
Propionibacterium granulosum		99.46	0.90		99	90	
Clostridium perfringens		0.28	0.39		99	1	
Fusobacterium mortiferum		0.11	0.43		1	5	
07431021	**Low discrimination**						**07431021**
					CAT		
Propionibacterium granulosum		88.12	0.56	URE 1	90		
Actinomyces naeslundii		11.63	0.55		25		
Clostridium perfringens		0.25	0.05		1		
07431022	**Very good identification**						**07431022**
					CAT		
Propionibacterium granulosum		98.82	0.79		90		
Clostridium perfringens		0.53	0.33		1		
Actinomyces naeslundii		0.35	0.52		25		
07431023	**Low discrimination**						**07431023**
Propionibacterium granulosum		65.46	0.45	URE 1			
Actinomyces naeslundii		34.55	0.55				
07431030	**Very good identification**						**07431030**
					CAT		
Propionibacterium granulosum		99.95	0.75	MLZ 12	90		
Actinomyces naeslundii		0.03	0.30		25		
Clostridium perfringens		0.02	0.05		1		
07431031	**Good identification**						**07431031**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		95.16	0.41	MLZ 12			
Actinomyces naeslundii		4.85	0.33	URE 1			
07431032	**Very good identification**						**07431032**
					GRA		
Propionibacterium granulosum		99.66	0.65	MLZ 12	99		
Parabacteroides distasonis		0.20	0.20		1		
Actinomyces naeslundii		0.14	0.30		99		
07431040	**Very good identification**						**07431040**
					GRA	CAT	
Propionibacterium granulosum		96.79	0.66	SOR 1	99	90	
Fusobacterium mortiferum		2.74	0.43		1	5	
Actinomyces naeslundii		0.38	0.39		99	25	
07431042	**Good identification**						**07431042**
					GRA	CAT	
Propionibacterium granulosum		97.27	0.56	SOR 1	99	90	
Actinomyces naeslundii		1.51	0.39		99	25	
Fusobacterium mortiferum		1.22	0.26		1	5	
07431043	**Low discrimination**						**07431043**
Actinomyces naeslundii		69.74	0.42	MLT 94			
Propionibacterium granulosum		30.26	0.21				
07431050	**Good identification**						**07431050**
Propionibacterium granulosum		100.0	0.51	MLZ 12			
07431052	**Good identification**						**07431052**
Propionibacterium granulosum		100.0	0.41	MLZ 12			
07431060	**Acceptable identification**						**07431060**
					CAT		
Propionibacterium granulosum		93.51	0.56	SOR 1	90		
Clostridium perfringens		5.04	0.26		1		
Actinomyces naeslundii		1.45	0.39		25		
07431061	**Low discrimination**						**07431061**
Actinomyces naeslundii		69.74	0.42	MLT 94			
Propionibacterium granulosum		30.26	0.21				
07431062	**Low discrimination**						**07431062**
					CAT		
Propionibacterium granulosum		85.95	0.45	SOR 1	90		
Clostridium perfringens		8.72	0.21		1		
Actinomyces naeslundii		5.34	0.39		25		
07431063	**Acceptable identification**						**07431063**
Actinomyces naeslundii		90.22	0.42	MLT 94			
Propionibacterium granulosum		9.78	0.11				
07431070	**Good identification**						**07431070**
Propionibacterium granulosum		100.0	0.41	MLZ 12			
07431100	**Low discrimination**						**07431100**
					GRA	CAT	
Fusobacterium mortiferum		85.16	0.83		1	5	
Propionibacterium granulosum		13.02	0.66		99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.82	0.60		99	25	
07431101	**Low discrimination**						**07431101**
					GRA	CAT	
Actinomyces naeslundii		54.53	0.63	MLT 94	99	25	
Fusobacterium mortiferum		38.39	0.49		1	5	
Propionibacterium granulosum		5.87	0.32		99	90	
07431102	**Low discrimination**						**07431102**
					GRA	CAT	
Fusobacterium mortiferum		65.43	0.67	BGL 10	1	5	
Propionibacterium granulosum		22.50	0.56		99	90	
Actinomyces naeslundii		5.63	0.60		99	25	
07431103	**Low discrimination**						**07431103**
					GRA	CAT	
Actinomyces naeslundii		89.51	0.63	MLT 94	99	25	
Fusobacterium mortiferum		7.00	0.33		1	5	
Propionibacterium granulosum		2.41	0.21		99	90	
07431110	**Low discrimination**						**07431110**
					GRA	CAT	
Propionibacterium granulosum		66.09	0.51	LAC 1	99	90	
Fusobacterium mortiferum		32.03	0.49	MLZ 12	1	5	
Actinomyces naeslundii		1.60	0.38		99	25	
07431111	**Low discrimination**						**07431111**
					GRA	CAT	
Actinomyces naeslundii		70.73	0.41	MLT 94	99	25	
Propionibacterium granulosum		19.72	0.17	MLZ 5	99	90	
Fusobacterium mortiferum		9.56	0.15		1	5	
07431112	**Low discrimination**						**07431112**
					GRA	CAT	
Propionibacterium granulosum		72.47	0.41	LAC 1	99	90	
Fusobacterium mortiferum		15.61	0.33	MLZ 12	1	5	
Actinomyces naeslundii		7.00	0.38		99	25	
07431113	**Good identification**						**07431113**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07431120	**Low discrimination**						**07431120**
					GRA	CAT	
Clostridium perfringens		52.36	0.54	MLT 99	99	1	
Propionibacterium granulosum		25.24	0.56		99	90	
Prevotella melaninogenica		7.02	0.32		1	1	
07431121	**Good identification**						**07431121**
					CAT		
Actinomyces naeslundii		90.69	0.63	MLT 94	25		
Clostridium perfringens		5.06	0.19		1		
Propionibacterium granulosum		2.44	0.21		90		
07431122	**Low discrimination**						**07431122**
					GRA		
Parabacteroides distasonis		43.20	0.59	XYL 90	1		
Clostridium perfringens		33.17	0.48		99		
Actinomyces naeslundii		8.51	0.60		99		
07431123	**Good identification**						**07431123**
					GRA		
Actinomyces naeslundii		93.71	0.63	MLT 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		3.21	0.25		1		
Clostridium perfringens		2.46	0.14		99		
07431130	**Low discrimination**						**07431130**
					CAT		
Propionibacterium granulosum		72.75	0.41	LAC 1	90		
Clostridium perfringens		11.18	0.19	MLZ 12	1		
Actinomyces naeslundii		7.03	0.38		25		
07431131	**Good identification**						**07431131**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07431132	**Acceptable identification**						**07431132**
					GRA		
Parabacteroides distasonis		91.73	0.55	XYL 90	1		
Propionibacterium granulosum		4.57	0.30		99		
Actinomyces naeslundii		1.77	0.38		99		
07431133	**Low discrimination**						**07431133**
					GRA		
Actinomyces naeslundii		74.08	0.41	MLT 94	99		
Parabacteroides distasonis		25.92	0.20	MLZ 5	1		
07431140	**Low discrimination**						**07431140**
					GRA	CAT	
Fusobacterium mortiferum		72.72	0.49	SOR 1	1	5	
Actinomyces naeslundii		12.16	0.47		99	25	
Propionibacterium granulosum		11.12	0.32		99	90	
07431141	**Good identification**						**07431141**
					GRA		
Actinomyces naeslundii		96.13	0.50	MLT 94	99		
Fusobacterium mortiferum		3.87	0.15		1		
07431142	**Low discrimination**						**07431142**
					GRA	CAT	
Actinomyces naeslundii		48.15	0.47	MLT 94	99	25	
Fusobacterium mortiferum		32.01	0.33		1	5	
Propionibacterium granulosum		11.01	0.21		99	90	
07431143	**Good identification**						**07431143**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07431160	**Low discrimination**						**07431160**
					CAT		
Clostridium perfringens		86.86	0.41	MLT 99	1		
Actinomyces naeslundii		9.71	0.47		25		
Propionibacterium granulosum		2.22	0.21		90		
07431161	**Acceptable identification**						**07431161**
Actinomyces naeslundii		94.32	0.50	MLT 94			
Clostridium perfringens		5.68	0.07				
07431162	**Low discrimination**						**07431162**
					GRA		
Clostridium perfringens		75.75	0.36	MLT 99	99		
Actinomyces naeslundii		18.00	0.47		99		
Parabacteroides distasonis		5.23	0.25		1		
07431163	**Good identification**						**07431163**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		97.24	0.50	MLT 94			
Clostridium perfringens		2.76	0.01				
07431200	**Very good identification**						**07431200**
					GRA	CAT	
Propionibacterium granulosum		99.08	0.74	RHA 3	99	90	
Fusobacterium mortiferum		0.92	0.43		1	5	
07431201	**Good identification**						**07431201**
Propionibacterium granulosum		100.0	0.40	RHA 3			
07431202	**Very good identification**						**07431202**
					GRA	CAT	
Propionibacterium granulosum		99.59	0.64	RHA 3	99	90	
Fusobacterium mortiferum		0.41	0.26		1	5	
07431210	**Very good identification**						**07431210**
Propionibacterium granulosum		100.0	0.59	RHA 3			
07431212	**Good identification**						**07431212**
Propionibacterium granulosum		100.0	0.49	RHA 3			
07431220	**Very good identification**						**07431220**
					CAT		
Propionibacterium granulosum		99.91	0.64	RHA 3	90		
Clostridium perfringens		0.09	0.05		1		
07431222	**Good identification**						**07431222**
					GRA		
Propionibacterium granulosum		95.21	0.54	RHA 3	99		
Parabacteroides distasonis		4.79	0.33		1		
07431230	**Good identification**						**07431230**
Propionibacterium granulosum		100.0	0.49	RHA 3			
07431232	**Low discrimination**						**07431232**
					GRA		
Propionibacterium granulosum		83.42	0.39	RHA 3	99		
Parabacteroides distasonis		16.58	0.29	MLZ 12	1		
07431240	**Good identification**						**07431240**
Propionibacterium granulosum		100.0	0.40	RHA 3			
07431300	**Low discrimination**						**07431300**
					GRA	CAT	
Fusobacterium mortiferum		67.27	0.49	RHA 1	1	5	
Propionibacterium granulosum		31.49	0.40		99	90	
Actinomyces naeslundii		0.64	0.26		99	25	
07431322	**Very good identification**						**07431322**
					GRA		
Parabacteroides distasonis		99.14	0.67	XYL 90	1		
Prevotella oralis		0.34	0.13		1		
Clostridium perfringens		0.26	0.14		99		
07431332	**Very good identification**						**07431332**
Parabacteroides distasonis		100.0	0.63	XYL 90			
07431400	**Good identification**						**07431400**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		93.98	0.66	ARA 1	99	90	
Lactobacillus fermentum		3.36	0.46		99	1	
Fusobacterium mortiferum		2.66	0.43		1	5	
07431402	**Low discrimination**						**07431402**
					GRA	CAT	
Propionibacterium granulosum		86.55	0.56	ARA 1	99	90	
Lactobacillus fermentum		12.36	0.46		99	1	
Fusobacterium mortiferum		1.09	0.26		1	5	
07431410	**Good identification**						**07431410**
Propionibacterium granulosum		100.0	0.51	ARA 1			
07431412	**Good identification**						**07431412**
Propionibacterium granulosum		100.0	0.41	ARA 1			
07431420	**Acceptable identification**						**07431420**
					CAT		
Propionibacterium granulosum		94.95	0.56	ARA 1	90		
Lactobacillus fermentum		4.52	0.38		1		
Clostridium perfringens		0.27	0.05		1		
07431422	**Low discrimination**						**07431422**
					GRA	CAT	
Propionibacterium granulosum		81.13	0.45	ARA 1	99	90	
Lactobacillus fermentum		15.45	0.38		99	1	
Parabacteroides distasonis		2.55	0.21		1	76	
07431430	**Good identification**						**07431430**
					CAT		
Propionibacterium granulosum		96.76	0.41	ARA 1	90		
Bifidobacterium longum subsp. longum		3.24	0.13	MLZ 12	1		
07431500	**Low discrimination**						**07431500**
					GRA		
Fusobacterium mortiferum		43.58	0.49	ARA 1	1		
Lactobacillus fermentum		38.41	0.50		99		
Bifidobacterium longum subsp. longum		10.93	0.33		99		
07431502	**Low discrimination**						**07431502**
					GRA		
Lactobacillus fermentum		65.35	0.50	TRE 1	99		
Bifidobacterium longum subsp. longum		18.60	0.33		99		
Fusobacterium mortiferum		8.24	0.33		1		
07431510	**Acceptable identification**						**07431510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.47	0.37	MLT 99	99	1	
Propionibacterium granulosum		4.46	0.17		99	90	
Fusobacterium mortiferum		2.16	0.15		1	5	
07431512	**Low discrimination**						**07431512**
					GRA		
Bifidobacterium longum subsp. longum		87.86	0.37	MLT 99	99		
Bifidobacterium adolescentis		7.01	0.20		99		
Parabacteroides distasonis		3.30	0.17		1		
07431520	**Low discrimination**						**07431520**
Bifidobacterium longum subsp. longum		68.03	0.43	MLT 99			
Lactobacillus fermentum		19.93	0.41				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		5.38	0.19				
07431522	**Low discrimination**						**07431522**
					GRA		
Parabacteroides distasonis		68.53	0.56	XYL 90	1		
Bifidobacterium longum subsp. longum		23.32	0.43		99		
Lactobacillus fermentum		6.83	0.41		99		
07431530	**Good identification**						**07431530**
					GRA		
Bifidobacterium longum subsp. longum		99.07	0.47	MLT 99	99		
Parabacteroides distasonis		0.93	0.17		1		
07431532	**Low discrimination**						**07431532**
					GRA		
Bifidobacterium longum subsp. longum		51.84	0.47	MLT 99	99		
Parabacteroides distasonis		48.17	0.51		1		
07431600	**Good identification**						**07431600**
Propionibacterium granulosum		100.0	0.40	RHA 3			
07431722	**Very good identification**						**07431722**
					GRA	CAT	
Parabacteroides distasonis		99.69	0.64	XYL 90	1	76	
Bacteroides thetaiotaomicron		0.12	-.02		1	89	
Bifidobacterium longum subsp. longum		0.11	0.09		99	1	
07431732	**Very good identification**						**07431732**
					GRA		
Parabacteroides distasonis		99.01	0.59	XYL 90	1		
Clostridium clostridioforme		0.63	0.16		50		
Bifidobacterium longum subsp. longum		0.36	0.13		99		
07432000	**Excellent identification**						**07432000**
					GRA	CAT	
Propionibacterium granulosum		99.97	0.98		99	90	
Fusobacterium mortiferum		0.03	0.40		1	5	
Actinomyces naeslundii		0.00	0.34		99	25	
07432001	**Very good identification**						**07432001**
Propionibacterium granulosum		99.65	0.64	URE 1			
Actinomyces naeslundii		0.35	0.37				
07432002	**Very good identification**						**07432002**
					GRA	CAT	
Propionibacterium granulosum		99.98	0.88		99	90	
Fusobacterium mortiferum		0.01	0.23		1	5	
Actinomyces naeslundii		0.01	0.34		99	25	
07432003	**Good identification**						**07432003**
Propionibacterium granulosum		98.61	0.53	URE 1			
Actinomyces naeslundii		1.40	0.37				
07432010	**Very good identification**						**07432010**
					SPO	CAT	
Propionibacterium granulosum		99.99	0.83	MLZ 12	1	90	
Clostridium difficile		0.01	0.04		99	1	
07432011	**Good identification**						**07432011**
Propionibacterium granulosum		100.0	0.49	MLZ 12			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07432012	**Very good identification**						**07432012**
Propionibacterium granulosum		100.0	0.73	MLZ 12			
07432013	**Good identification**						**07432013**
Propionibacterium granulosum		100.0	0.39	MLZ 12			
07432020	**Very good identification**						**07432020**
Propionibacterium granulosum		99.99	0.88				
Actinomyces naeslundii		0.01	0.34				
07432021	**Good identification**						**07432021**
Propionibacterium granulosum		98.61	0.53	URE 1			
Actinomyces naeslundii		1.40	0.37				
07432022	**Very good identification**						**07432022**
Propionibacterium granulosum		99.96	0.77				
Actinomyces naeslundii		0.04	0.34				
07432023	**Acceptable identification**						**07432023**
Propionibacterium granulosum		94.65	0.43	URE 1			
Actinomyces naeslundii		5.36	0.37				
07432030	**Very good identification**						**07432030**
Propionibacterium granulosum		100.0	0.73	MLZ 12			
07432031	**Good identification**						**07432031**
Propionibacterium granulosum		100.0	0.39	MLZ 12			
07432032	**Very good identification**						**07432032**
Propionibacterium granulosum		100.0	0.63	MLZ 12			
07432040	**Very good identification**						**07432040**
					SPO	CAT	
Propionibacterium granulosum		99.93	0.64	SOR 1	1	90	
Clostridium botulinum biovar B		0.07	-.03		99	1	
07432042	**Good identification**						**07432042**
Propionibacterium granulosum		100.0	0.53	SOR 1			
07432050	**Good identification**						**07432050**
Propionibacterium granulosum		100.0	0.49	MLZ 12			
07432052	**Good identification**						**07432052**
Propionibacterium granulosum		100.0	0.39	MLZ 12			
07432060	**Good identification**						**07432060**
Propionibacterium granulosum		100.0	0.53	SOR 1			
07432062	**Good identification**						**07432062**
Propionibacterium granulosum		100.0	0.43	SOR 1			
07432070	**Good identification**						**07432070**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.39	MLZ 12			
07432100	**Good identification**						**07432100**
					GRA	CAT	
Propionibacterium granulosum		91.48	0.64	LAC 1	99	90	
Fusobacterium mortiferum		5.34	0.46		1	5	
Propionibacterium propionicum		2.57	0.42		99	1	
07432102	**Low discrimination**						**07432102**
					CAT		
Propionibacterium granulosum		81.88	0.53	LAC 1	90		
Propionibacterium propionicum		13.80	0.45		1		
Actinomyces naeslundii		2.20	0.41		25		
07432103	**Low discrimination**						**07432103**
					CAT		
Actinomyces naeslundii		58.28	0.44	MLT 94	25		
Propionibacterium propionicum		27.10	0.29	GAL 96	1		
Propionibacterium granulosum		14.62	0.19		90		
07432110	**Good identification**						**07432110**
					GRA	CAT	
Propionibacterium granulosum		99.57	0.49	LAC 1	99	90	
Fusobacterium mortiferum		0.43	0.12	MLZ 12	1	5	
07432112	**Good identification**						**07432112**
Propionibacterium granulosum		100.0	0.39	LAC 1			
07432120	**Good identification**						**07432120**
					GRA	CAT	
Propionibacterium granulosum		97.16	0.53	LAC 1	99	90	
Actinomyces naeslundii		2.61	0.41		99	25	
Fusobacterium mortiferum		0.23	0.12		1	5	
07432121	**Low discrimination**						**07432121**
Actinomyces naeslundii		79.94	0.44	MLT 94			
Propionibacterium granulosum		20.06	0.19	GAL 96			
07432122	**Acceptable identification**						**07432122**
Propionibacterium granulosum		90.30	0.43	LAC 1			
Actinomyces naeslundii		9.70	0.41				
07432123	**Acceptable identification**						**07432123**
Actinomyces naeslundii		94.10	0.44	MLT 94			
Propionibacterium granulosum		5.90	0.09	GAL 96			
07432130	**Good identification**						**07432130**
Propionibacterium granulosum		100.0	0.39	LAC 1			
07432140	**Low discrimination**						**07432140**
					CAT		
Propionibacterium propionicum		70.29	0.42	NIT 86	1		
Propionibacterium granulosum		25.28	0.30	MLT 99	90		
Actinomyces naeslundii		2.96	0.29		25		
07432142	**Acceptable identification**						**07432142**
					CAT		
Propionibacterium propionicum		91.91	0.45	NIT 86	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		5.51	0.19	MLT 99	90		
Actinomyces naeslundii		2.58	0.29		25		
07432200	**Very good identification**						**07432200**
Propionibacterium granulosum		100.0	0.72	RHA 3			
07432201	**Good identification**						**07432201**
Propionibacterium granulosum		100.0	0.38	RHA 3			
07432202	**Very good identification**						**07432202**
Propionibacterium granulosum		100.0	0.62	RHA 3			
07432210	**Very good identification**						**07432210**
Propionibacterium granulosum		100.0	0.57	RHA 3			
07432212	**Good identification**						**07432212**
Propionibacterium granulosum		100.0	0.47	RHA 3			
07432220	**Very good identification**						**07432220**
Propionibacterium granulosum		100.0	0.62	RHA 3			
07432222	**Good identification**						**07432222**
Propionibacterium granulosum		100.0	0.52	RHA 3			
07432230	**Good identification**						**07432230**
Propionibacterium granulosum		100.0	0.47	RHA 3			
07432232	**Good identification**						**07432232**
Propionibacterium granulosum		100.0	0.37	RHA 3			
07432240	**Good identification**						**07432240**
Propionibacterium granulosum		100.0	0.38	RHA 3			
07432300	**Good identification**						**07432300**
					GRA	CAT	
Propionibacterium granulosum		98.13	0.38	LAC 1	99	90	
Fusobacterium mortiferum		1.87	0.12	RHA 3	1	5	
07432400	**Very good identification**						**07432400**
Propionibacterium granulosum		100.0	0.64	ARA 1			
07432402	**Good identification**						**07432402**
Propionibacterium granulosum		100.0	0.53	ARA 1			
07432410	**Good identification**						**07432410**
Propionibacterium granulosum		100.0	0.49	ARA 1			
07432412	**Good identification**						**07432412**
Propionibacterium granulosum		100.0	0.39	ARA 1			
07432420	**Good identification**						**07432420**
Propionibacterium granulosum		100.0	0.53	ARA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07432422	**Good identification**						**07432422**
Propionibacterium granulosum		100.0	0.43	ARA 1			
07432430	**Good identification**						**07432430**
Propionibacterium granulosum		100.0	0.39	ARA 1			
07432600	**Good identification**						**07432600**
Propionibacterium granulosum		100.0	0.38	RHA 3			
07433000	**Excellent identification**						**07433000**
					GRA	CAT	
Propionibacterium granulosum		99.91	0.98		99	90	
Actinomyces naeslundii		0.06	0.57		99	25	
Fusobacterium mortiferum		0.04	0.43		1	5	
07433001	**Good identification**						**07433001**
Propionibacterium granulosum		92.18	0.64	URE 1			
Actinomyces naeslundii		7.82	0.60				
07433002	**Very good identification**						**07433002**
					GRA	CAT	
Propionibacterium granulosum		99.76	0.88		99	90	
Actinomyces naeslundii		0.23	0.57		99	25	
Fusobacterium mortiferum		0.02	0.26		1	5	
07433003	**Low discrimination**						**07433003**
Propionibacterium granulosum		74.65	0.53	URE 1			
Actinomyces naeslundii		25.35	0.60				
07433010	**Very good identification**						**07433010**
Propionibacterium granulosum		99.98	0.83	MLZ 12			
Actinomyces naeslundii		0.02	0.35				
07433011	**Good identification**						**07433011**
Propionibacterium granulosum		96.83	0.49	MLZ 12			
Actinomyces naeslundii		3.17	0.38	URE 1			
07433012	**Very good identification**						**07433012**
Propionibacterium granulosum		99.91	0.73	MLZ 12			
Actinomyces naeslundii		0.09	0.35				
07433013	**Low discrimination**						**07433013**
Propionibacterium granulosum		88.41	0.39	MLZ 12			
Actinomyces naeslundii		11.59	0.38	URE 1			
07433020	**Very good identification**						**07433020**
					CAT		
Propionibacterium granulosum		99.77	0.88		90		
Actinomyces naeslundii		0.23	0.57		25		
Clostridium perfringens		0.00	0.05		1		
07433021	**Low discrimination**						**07433021**
Propionibacterium granulosum		74.65	0.53	URE 1			
Actinomyces naeslundii		25.35	0.60				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07433022	**Very good identification**					**07433022**
Propionibacterium granulosum		99.09	0.77			
Actinomyces naeslundii		0.91	0.57			
07433023	**Low discrimination**					**07433023**
Actinomyces naeslundii		57.59	0.60	MLT 94		
Propionibacterium granulosum		42.41	0.43			
07433030	**Very good identification**					**07433030**
Propionibacterium granulosum		99.91	0.73	MLZ 12		
Actinomyces naeslundii		0.09	0.35			
07433031	**Low discrimination**					**07433031**
Propionibacterium granulosum		88.41	0.39	MLZ 12		
Actinomyces naeslundii		11.59	0.38	URE 1		
07433032	**Very good identification**					**07433032**
Propionibacterium granulosum		99.65	0.63	MLZ 12		
Actinomyces naeslundii		0.35	0.35			
07433040	**Very good identification**					**07433040**
Propionibacterium granulosum		99.01	0.64	SOR 1		
Actinomyces naeslundii		0.99	0.44			
07433041	**Low discrimination**					**07433041**
Actinomyces naeslundii		59.72	0.47	MLT 94		
Propionibacterium granulosum		40.28	0.30			
07433042	**Good identification**					**07433042**
Propionibacterium granulosum		96.16	0.53	SOR 1		
Actinomyces naeslundii		3.84	0.44			
07433043	**Low discrimination**					**07433043**
Actinomyces naeslundii		85.57	0.47	MLT 94		
Propionibacterium granulosum		14.43	0.19			
07433050	**Good identification**					**07433050**
Propionibacterium granulosum		99.62	0.49	MLZ 12		
Actinomyces naeslundii		0.38	0.23	SOR 1		
07433052	**Good identification**					**07433052**
Propionibacterium granulosum		98.48	0.39	MLZ 12		
Actinomyces naeslundii		1.52	0.23	SOR 1		
07433060	**Good identification**					**07433060**
Propionibacterium granulosum		96.16	0.53	SOR 1		
Actinomyces naeslundii		3.84	0.44			
07433061	**Low discrimination**					**07433061**
Actinomyces naeslundii		85.57	0.47	MLT 94		
Propionibacterium granulosum		14.43	0.19			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07433062	**Low discrimination**						**07433062**
Propionibacterium granulosum		86.22	0.43	SOR 1			
Actinomyces naeslundii		13.78	0.44				
07433063	**Good identification**						**07433063**
Actinomyces naeslundii		95.96	0.47	MLT 94			
Propionibacterium granulosum		4.04	0.09				
07433070	**Good identification**						**07433070**
Propionibacterium granulosum		98.48	0.39	MLZ 12			
Actinomyces naeslundii		1.52	0.23	SOR 1			
07433100	**Low discrimination**						**07433100**
					GRA	CAT	
Propionibacterium granulosum		78.28	0.64	LAC 1	99	90	
Actinomyces naeslundii		12.61	0.65		99	25	
Fusobacterium mortiferum		6.86	0.49		1	5	
07433101	**Good identification**						**07433101**
					CAT		
Actinomyces naeslundii		94.48	0.68	MLT 94	25		
Propionibacterium granulosum		3.95	0.30		90		
Propionibacterium propionicum		1.22	0.26		1		
07433102	**Low discrimination**						**07433102**
					CAT		
Propionibacterium granulosum		53.92	0.53	LAC 1	90		
Actinomyces naeslundii		34.73	0.65		25		
Propionibacterium propionicum		9.09	0.45		1		
07433103	**Very good identification**						**07433103**
					CAT		
Actinomyces naeslundii		97.10	0.68	MLT 94	25		
Propionibacterium propionicum		1.88	0.29		1		
Propionibacterium granulosum		1.02	0.19		90		
07433110	**Acceptable identification**						**07433110**
					GRA	CAT	
Propionibacterium granulosum		93.58	0.49	LAC 1	99	90	
Actinomyces naeslundii		5.82	0.43	MLZ 12	99	25	
Fusobacterium mortiferum		0.61	0.15		1	5	
07433111	**Acceptable identification**						**07433111**
Actinomyces naeslundii		90.23	0.46	MLT 94			
Propionibacterium granulosum		9.78	0.15	MLZ 5			
07433112	**Low discrimination**						**07433112**
Propionibacterium granulosum		80.09	0.39	LAC 1			
Actinomyces naeslundii		19.91	0.43	MLZ 12			
07433113	**Good identification**						**07433113**
Actinomyces naeslundii		100.0	0.46	MLT 94			
07433120	**Low discrimination**						**07433120**
					CAT		
Propionibacterium granulosum		59.55	0.53	LAC 1	90		
Actinomyces naeslundii		38.36	0.65		25		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		1.65	0.19		1	
07433121	**Very good identification**					**07433121**
Actinomyces naeslundii		98.97	0.68	MLT 94		
Propionibacterium granulosum		1.04	0.19			
07433122	**Low discrimination**					**07433122**
					GRA	
Actinomyces naeslundii		69.70	0.65	MLT 94	99	
Propionibacterium granulosum		27.05	0.43		99	
Parabacteroides distasonis		1.84	0.25		1	
07433123	**Very good identification**					**07433123**
Actinomyces naeslundii		99.74	0.68	MLT 94		
Propionibacterium granulosum		0.26	0.09			
07433130	**Low discrimination**					**07433130**
Propionibacterium granulosum		80.09	0.39	LAC 1		
Actinomyces naeslundii		19.91	0.43	MLZ 12		
07433131	**Good identification**					**07433131**
Actinomyces naeslundii		100.0	0.46	MLT 94		
07433133	**Good identification**					**07433133**
Actinomyces naeslundii		100.0	0.46	MLT 94		
07433140	**Low discrimination**					**07433140**
					CAT	
Actinomyces naeslundii		42.11	0.52	MLT 94	25	
Propionibacterium propionicum		41.62	0.42		1	
Propionibacterium granulosum		14.97	0.30		90	
07433141	**Acceptable identification**					**07433141**
Actinomyces naeslundii		93.18	0.55	MLT 94		
Propionibacterium propionicum		6.82	0.26			
07433142	**Low discrimination**					**07433142**
					CAT	
Propionibacterium propionicum		57.65	0.45	NIT 86	1	
Actinomyces naeslundii		38.89	0.52	MLT 99	25	
Propionibacterium granulosum		3.46	0.19		90	
07433143	**Acceptable identification**					**07433143**
Actinomyces naeslundii		90.11	0.55	MLT 94		
Propionibacterium propionicum		9.89	0.29			
07433160	**Low discrimination**					**07433160**
					CAT	
Actinomyces naeslundii		88.07	0.52	MLT 94	25	
Propionibacterium granulosum		7.83	0.19		90	
Clostridium perfringens		4.10	0.07		1	
07433161	**Good identification**					**07433161**
Actinomyces naeslundii		100.0	0.55	MLT 94		
07433162	**Good identification**					**07433162**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		95.78	0.52	MLT 94	25		
Propionibacterium granulosum		2.13	0.09		90		
Clostridium perfringens		2.10	0.01		1		
07433163	**Good identification**						**07433163**
Actinomyces naeslundii		100.0	0.55	MLT 94			
07433200	**Very good identification**						**07433200**
Propionibacterium granulosum		99.98	0.72	RHA 3			
Actinomyces naeslundii		0.02	0.23				
07433201	**Good identification**						**07433201**
Propionibacterium granulosum		97.30	0.38	RHA 3			
Actinomyces naeslundii		2.70	0.26	URE 1			
07433202	**Very good identification**						**07433202**
Propionibacterium granulosum		99.93	0.62	RHA 3			
Actinomyces naeslundii		0.08	0.23				
07433210	**Very good identification**						**07433210**
Propionibacterium granulosum		100.0	0.57	RHA 3			
07433212	**Good identification**						**07433212**
Propionibacterium granulosum		100.0	0.47	RHA 3			
07433220	**Very good identification**						**07433220**
Propionibacterium granulosum		99.93	0.62	RHA 3			
Actinomyces naeslundii		0.08	0.23				
07433222	**Good identification**						**07433222**
Propionibacterium granulosum		99.70	0.52	RHA 3			
Actinomyces naeslundii		0.30	0.23				
07433230	**Good identification**						**07433230**
Propionibacterium granulosum		100.0	0.47	RHA 3			
07433232	**Good identification**						**07433232**
Propionibacterium granulosum		100.0	0.37	RHA 3			
07433240	**Good identification**						**07433240**
Propionibacterium granulosum		100.0	0.38	RHA 3			
07433300	**Acceptable identification**						**07433300**
					GRA	CAT	
Propionibacterium granulosum		92.49	0.38	LAC 1	99	90	
Actinomyces naeslundii		4.87	0.31	RHA 3	99	25	
Fusobacterium mortiferum		2.65	0.15		1	5	
07433400	**Very good identification**						**07433400**
Propionibacterium granulosum		99.94	0.64	ARA 1			
Actinomyces naeslundii		0.06	0.23				
07433402	**Good identification**						**07433402**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		99.77	0.53	ARA 1				
Actinomyces naeslundii		0.23	0.23					
07433410	**Good identification**							**07433410**
Propionibacterium granulosum		100.0	0.49	ARA 1				
07433412	**Good identification**							**07433412**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07433420	**Good identification**							**07433420**
Propionibacterium granulosum		99.77	0.53	ARA 1				
Actinomyces naeslundii		0.23	0.23					
07433422	**Good identification**							**07433422**
Propionibacterium granulosum		99.09	0.43	ARA 1				
Actinomyces naeslundii		0.91	0.23					
07433430	**Good identification**							**07433430**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07433600	**Good identification**							**07433600**
Propionibacterium granulosum		100.0	0.38	RHA 3				
07434000	**Low discrimination**							**07434000**
					COC	GRA	CAT	
Propionibacterium granulosum		87.63	0.66	XYL 1	1	99	90	
Fusobacterium mortiferum		12.33	0.55		1	1	5	
Gemella morbillorum		0.04	0.03		99	99	1	
07434002	**Acceptable identification**							**07434002**
					GRA	CAT		
Propionibacterium granulosum		94.12	0.56	XYL 1	99	90		
Fusobacterium mortiferum		5.89	0.38		1	5		
07434010	**Good identification**							**07434010**
					GRA	CAT		
Propionibacterium granulosum		98.97	0.51	XYL 1	99	90		
Fusobacterium mortiferum		1.03	0.20	MLZ 12	1	5		
07434012	**Good identification**							**07434012**
Propionibacterium granulosum		100.0	0.41	XYL 1				
07434020	**Good identification**							**07434020**
					GRA	CAT		
Propionibacterium granulosum		99.44	0.56	XYL 1	99	90		
Fusobacterium mortiferum		0.57	0.20		1	5		
07434022	**Good identification**							**07434022**
Propionibacterium granulosum		100.0	0.45	XYL 1				
07434030	**Good identification**							**07434030**
Propionibacterium granulosum		100.0	0.41	XYL 1				
07434100	**Very good identification**							**07434100**
					GRA	CAT		
Fusobacterium mortiferum		97.02	0.61	XYL 7	1	5		
Propionibacterium granulosum		2.99	0.32		99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07434102	**Acceptable identification**						**07434102**
					GRA	CAT	
Fusobacterium mortiferum		93.53	0.45	XYL 7	1	5	
Propionibacterium granulosum		6.48	0.21	BGL 10	99	90	
07434122	**Good identification**						**07434122**
					GRA		
Parabacteroides distasonis		97.86	0.41	GAL 99	1		
Propionibacterium granulosum		2.14	0.11		99		
07434132	**Good identification**						**07434132**
Parabacteroides distasonis		100.0	0.37	GAL 99			
07434200	**Good identification**						**07434200**
					GRA	CAT	
Propionibacterium granulosum		95.61	0.40	XYL 1	99	90	
Fusobacterium mortiferum		4.39	0.20	RHA 3	1	5	
07434322	**Good identification**						**07434322**
Parabacteroides distasonis		100.0	0.50	GAL 99			
07434332	**Good identification**						**07434332**
Parabacteroides distasonis		99.09	0.45	GAL 99			
Clostridium clostridioforme		0.91	0.04				
07434522	**Low discrimination**						**07434522**
					GRA		
Parabacteroides distasonis		89.81	0.38	GAL 99	1		
Bifidobacterium longum subsp. longum		10.19	0.17		99		
07434722	**Good identification**						**07434722**
Parabacteroides distasonis		97.29	0.46	GAL 99			
Clostridium clostridioforme		2.71	0.14				
07434732	**Low discrimination**						**07434732**
Parabacteroides distasonis		80.93	0.41	GAL 99			
Clostridium clostridioforme		19.07	0.25				
07435000	**Low discrimination**						**07435000**
					GRA	CAT	
Propionibacterium granulosum		81.17	0.66	XYL 1	99	90	
Fusobacterium mortiferum		17.13	0.58		1	5	
Lactobacillus fermentum		1.70	0.42		99	1	
07435002	**Low discrimination**						**07435002**
					GRA	CAT	
Propionibacterium granulosum		84.91	0.56	XYL 1	99	90	
Fusobacterium mortiferum		7.96	0.41		1	5	
Lactobacillus fermentum		7.12	0.42		99	1	
07435010	**Good identification**						**07435010**
					GRA	CAT	
Propionibacterium granulosum		98.46	0.51	XYL 1	99	90	
Fusobacterium mortiferum		1.54	0.23	MLZ 12	1	5	
07435012	**Good identification**						**07435012**
Propionibacterium granulosum		100.0	0.41	XYL 1			
07435020	**Good identification**						**07435020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Propionibacterium granulosum		96.22	0.56	XYL 1	99	90	
Lactobacillus fermentum		2.69	0.34		99	1	
Fusobacterium mortiferum		0.82	0.23		1	5	
07435022	**Low discrimination**						**07435022**
					GRA	CAT	
Propionibacterium granulosum		63.53	0.45	XYL 1	99	90	
Parabacteroides distasonis		29.37	0.41		1	76	
Lactobacillus fermentum		7.10	0.34		99	1	
07435030	**Good identification**						**07435030**
Propionibacterium granulosum		100.0	0.41	XYL 1			
07435032	**Low discrimination**						**07435032**
					GRA		
Parabacteroides distasonis		64.61	0.37	LAC 99	1		
Propionibacterium granulosum		35.39	0.30		99		
07435100	**Good identification**						**07435100**
					GRA	CAT	
Fusobacterium mortiferum		90.91	0.64	XYL 7	1	5	
Lactobacillus fermentum		6.32	0.46		99	1	
Propionibacterium granulosum		1.87	0.32		99	90	
07435102	**Low discrimination**						**07435102**
					GRA		
Fusobacterium mortiferum		44.13	0.48	XYL 7	1		
Lactobacillus fermentum		27.59	0.46	BGL 10	99		
Parabacteroides distasonis		23.30	0.41		1		
07435112	**Low discrimination**						**07435112**
					GRA		
Parabacteroides distasonis		80.95	0.37	NAG 99	1		
Bifidobacterium adolescentis		11.72	0.20		99		
Bifidobacterium longum subsp. longum		4.45	0.11		99		
07435120	**Low discrimination**						**07435120**
					GRA		
Parabacteroides distasonis		49.46	0.41	BGL 99	1		
Lactobacillus fermentum		19.52	0.37		99		
Clostridium perfringens		8.97	0.19		99		
07435122	**Very good identification**						**07435122**
					GRA		
Parabacteroides distasonis		99.30	0.76		1		
Lactobacillus fermentum		0.40	0.37		99		
Clostridium perfringens		0.09	0.14		99		
07435123	**Good identification**						**07435123**
					GRA		
Parabacteroides distasonis		96.82	0.41	URE 1	1		
Actinomyces naeslundii		3.18	0.29		99		
07435130	**Low discrimination**						**07435130**
					GRA		
Parabacteroides distasonis		81.98	0.37	BGL 99	1		
Bifidobacterium longum subsp. longum		18.02	0.21		99		
07435132	**Very good identification**						**07435132**
					GRA		
Parabacteroides distasonis		99.78	0.71		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		0.22	0.21		99		
07435133	**Good identification**						**07435133**
Parabacteroides distasonis		100.0	0.37	URE 1			
07435162	**Good identification**						**07435162**
					GRA		
Parabacteroides distasonis		98.40	0.41	SOR 1	1		
Clostridium perfringens		1.60	0.01		99		
07435172	**Good identification**						**07435172**
Parabacteroides distasonis		100.0	0.37	SOR 1			
07435200	**Acceptable identification**						**07435200**
					GRA	CAT	
Propionibacterium granulosum		93.55	0.40	XYL 1	99	90	
Fusobacterium mortiferum		6.45	0.23	RHA 3	1	5	
07435222	**Good identification**						**07435222**
					GRA		
Parabacteroides distasonis		97.82	0.50	LAC 99	1		
Propionibacterium granulosum		2.18	0.19		99		
07435232	**Good identification**						**07435232**
Parabacteroides distasonis		100.0	0.45	LAC 99			
07435302	**Good identification**						**07435302**
Parabacteroides distasonis		99.37	0.50	NAG 99			
Fusobacterium mortiferum		0.63	0.13				
07435312	**Good identification**						**07435312**
Parabacteroides distasonis		97.31	0.45	NAG 99			
Clostridium clostridioforme		2.69	0.13				
07435320	**Low discrimination**						**07435320**
Parabacteroides distasonis		87.51	0.50	BGL 99			
Bacteroides vulgatus		12.49	0.17				
07435322	**Very good identification**						**07435322**
Parabacteroides distasonis		100.0	0.84				
Prevotella oralis		0.01	-.03				
07435323	**Good identification**						**07435323**
Parabacteroides distasonis		100.0	0.50	URE 1			
07435330	**Good identification**						**07435330**
Parabacteroides distasonis		100.0	0.45	BGL 99			
07435332	**Very good identification**						**07435332**
Parabacteroides distasonis		99.97	0.79				
Clostridium clostridioforme		0.03	0.13				
07435333	**Good identification**						**07435333**
Parabacteroides distasonis		100.0	0.45	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07435362	**Good identification**						**07435362**
Parabacteroides distasonis		100.0	0.50	SOR 1			
07435372	**Good identification**						**07435372**
Parabacteroides distasonis		100.0	0.45	SOR 1			
07435400	**Low discrimination**						**07435400**
					GRA	CAT	
Lactobacillus fermentum		62.22	0.42	TRE 1	99	1	
Propionibacterium granulosum		29.97	0.32		99	90	
Fusobacterium mortiferum		6.32	0.23		1	5	
07435402	**Low discrimination**						**07435402**
					CAT		
Lactobacillus fermentum		87.38	0.42	TRE 1	1		
Propionibacterium granulosum		10.52	0.21		90		
Bifidobacterium longum subsp. longum		2.09	0.07		1		
07435422	**Low discrimination**						**07435422**
					GRA		
Parabacteroides distasonis		64.78	0.38	LAC 99	1		
Lactobacillus fermentum		25.56	0.34		99		
Bifidobacterium longum subsp. longum		7.35	0.17		99		
07435500	**Low discrimination**						**07435500**
					GRA		
Bifidobacterium longum subsp. longum		55.93	0.41	MLT 99	99		
Lactobacillus fermentum		38.48	0.46		99		
Fusobacterium mortiferum		5.59	0.30		1		
07435502	**Low discrimination**						**07435502**
					GRA		
Bifidobacterium longum subsp. longum		43.19	0.41	MLT 99	99		
Lactobacillus fermentum		29.71	0.46		99		
Parabacteroides distasonis		15.38	0.38		1		
07435510	**Good identification**						**07435510**
Bifidobacterium longum subsp. longum		100.0	0.45	MLT 99			
07435512	**Low discrimination**						**07435512**
					GRA		
Bifidobacterium longum subsp. longum		75.57	0.45	MLT 99	99		
Bifidobacterium adolescentis		13.46	0.34		99		
Parabacteroides distasonis		8.51	0.33		1		
07435520	**Low discrimination**						**07435520**
					GRA		
Bifidobacterium longum subsp. longum		80.76	0.52	MLT 99	99		
Bacteroides vulgatus		7.42	0.20		1		
Parabacteroides distasonis		7.19	0.38		1		
07435522	**Low discrimination**						**07435522**
					GRA		
Parabacteroides distasonis		89.20	0.72		1		
Bifidobacterium longum subsp. longum		10.12	0.52		99		
Lactobacillus fermentum		0.58	0.37		99		
07435523	**Low discrimination**						**07435523**
					GRA		
Parabacteroides distasonis		89.81	0.38	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		10.19	0.17		99	
07435530	**Good identification**					**07435530**
					GRA	
Bifidobacterium longum subsp. longum		97.26	0.56	MLT 99	99	
Parabacteroides distasonis		2.74	0.33		1	
07435532	**Low discrimination**					**07435532**
					GRA	
Parabacteroides distasonis		73.41	0.67		1	
Bifidobacterium longum subsp. longum		26.34	0.56		99	
Clostridium clostridioforme		0.21	0.18		50	
07435562	**Low discrimination**					**07435562**
					GRA	
Parabacteroides distasonis		89.81	0.38	SOR 1	1	
Bifidobacterium longum subsp. longum		10.19	0.17		99	
07435622	**Good identification**					**07435622**
Parabacteroides distasonis		97.57	0.46	LAC 99		
Clostridium clostridioforme		2.44	0.13			
07435632	**Low discrimination**					**07435632**
Parabacteroides distasonis		82.56	0.41	LAC 99		
Clostridium clostridioforme		17.44	0.24			
07435702	**Acceptable identification**					**07435702**
					GRA	
Parabacteroides distasonis		91.49	0.46	NAG 99	1	
Clostridium clostridioforme		7.65	0.22		50	
Bifidobacterium longum subsp. longum		0.87	0.07		99	
07435712	**Low discrimination**					**07435712**
					GRA	
Parabacteroides distasonis		57.57	0.41	NAG 99	1	
Clostridium clostridioforme		40.71	0.33		50	
Bifidobacterium longum subsp. longum		1.72	0.11		99	
07435720	**Low discrimination**					**07435720**
					GRA	
Bacteroides vulgatus		74.88	0.36	MLT 99	1	
Parabacteroides distasonis		24.21	0.46	TRE 4	1	
Bifidobacterium longum subsp. longum		0.92	0.17		99	
07435722	**Very good identification**					**07435722**
Parabacteroides distasonis		99.52	0.80			
Bacteroides thetaiotaomicron		0.33	0.22			
Clostridium clostridioforme		0.08	0.22			
07435723	**Good identification**					**07435723**
Parabacteroides distasonis		100.0	0.46	URE 1		
07435730	**Low discrimination**					**07435730**
					GRA	
Parabacteroides distasonis		84.92	0.41	BGL 99	1	
Bifidobacterium longum subsp. longum		10.15	0.21		99	
Bacteroides vulgatus		4.93	0.02		1	
07435732	**Very good identification**					**07435732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		98.94	0.76				
Clostridium clostridioforme		0.71	0.33				
Bacteroides thetaiotaomicron		0.24	0.15				
07435733	**Good identification**						**07435733**
Parabacteroides distasonis		100.0	0.41	URE 1			
07435762	**Good identification**						**07435762**
Parabacteroides distasonis		100.0	0.46	SOR 1			
07435772	**Good identification**						**07435772**
Parabacteroides distasonis		100.0	0.41	SOR 1			
07436000	**Very good identification**						**07436000**
					GRA	CAT	
Propionibacterium granulosum		99.81	0.64	XYL 1	99	90	
Fusobacterium mortiferum		0.19	0.20		1	5	
07436002	**Good identification**						**07436002**
Propionibacterium granulosum		100.0	0.53	XYL 1			
07436010	**Good identification**						**07436010**
Propionibacterium granulosum		100.0	0.49	XYL 1			
07436012	**Good identification**						**07436012**
Propionibacterium granulosum		100.0	0.39	XYL 1			
07436020	**Good identification**						**07436020**
Propionibacterium granulosum		100.0	0.53	XYL 1			
07436022	**Good identification**						**07436022**
Propionibacterium granulosum		100.0	0.43	XYL 1			
07436030	**Good identification**						**07436030**
Propionibacterium granulosum		100.0	0.39	XYL 1			
07436200	**Good identification**						**07436200**
Propionibacterium granulosum		100.0	0.38	XYL 1			
07437000	**Very good identification**						**07437000**
					GRA	CAT	
Propionibacterium granulosum		99.66	0.64	XYL 1	99	90	
Fusobacterium mortiferum		0.28	0.23		1	5	
Actinomyces naeslundii		0.06	0.23		99	25	
07437002	**Good identification**						**07437002**
					CAT		
Propionibacterium granulosum		99.56	0.53	XYL 1	90		
Actinomyces naeslundii		0.23	0.23		25		
Actinomyces israelii		0.22	0.14		1		
07437010	**Good identification**						**07437010**
Propionibacterium granulosum		100.0	0.49	XYL 1			
07437012	**Good identification**						**07437012**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.39	XYL 1				
07437020	**Good identification**							**07437020**
Propionibacterium granulosum		99.77	0.53	XYL 1				
Actinomyces naeslundii		0.23	0.23					
07437022	**Good identification**							**07437022**
Propionibacterium granulosum		99.09	0.43	XYL 1				
Actinomyces naeslundii		0.91	0.23					
07437030	**Good identification**							**07437030**
Propionibacterium granulosum		100.0	0.39	XYL 1				
07437122	**Acceptable identification**							**07437122**
					GRA			
Parabacteroides distasonis		94.43	0.41	MAN 1	1			
Actinomyces naeslundii		4.01	0.31		99			
Propionibacterium granulosum		1.56	0.09		99			
07437132	**Good identification**							**07437132**
Parabacteroides distasonis		100.0	0.37	MAN 1				
07437200	**Good identification**							**07437200**
Propionibacterium granulosum		100.0	0.38	XYL 1				
07437322	**Good identification**							**07437322**
Parabacteroides distasonis		100.0	0.50	MAN 1				
07437332	**Good identification**							**07437332**
Parabacteroides distasonis		100.0	0.45	MAN 1				
07437522	**Low discrimination**							**07437522**
					GRA			
Parabacteroides distasonis		89.81	0.38	MAN 1	1			
Bifidobacterium longum subsp. longum		10.19	0.17		99			
07437722	**Good identification**							**07437722**
Parabacteroides distasonis		100.0	0.46	MAN 1				
07437732	**Good identification**							**07437732**
Parabacteroides distasonis		100.0	0.41	MAN 1				
07440000	**Low discrimination**							**07440000**
					COC	GRA	CAT	
Fusobacterium mortiferum		74.79	0.40	FRU 99	1	1	5	
Gemella morbillorum		13.26	0.17		99	99	1	
Propionibacterium granulosum		11.95	0.24		1	99	90	
07440100	**Good identification**							**07440100**
Fusobacterium mortiferum		100.0	0.47	FRU 99				
07441000	**Low discrimination**							**07441000**
					COC	GRA	CAT	
Fusobacterium mortiferum		83.67	0.43	FRU 99	1	1	5	
Propionibacterium granulosum		8.91	0.24		1	99	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		5.33	0.13		99	99	1	
07441100	**Good identification**							**07441100**
					GRA			
Fusobacterium mortiferum		95.23	0.50	FRU 99	1			
Actinomyces naeslundii		2.91	0.35		99			
Prevotella melaninogenica		1.11	-.01		1			
07441101	**Low discrimination**							**07441101**
					GRA			
Actinomyces naeslundii		81.92	0.38	MLT 94	99			
Fusobacterium mortiferum		18.08	0.16	FRU 95	1			
07441103	**Good identification**							**07441103**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07441121	**Low discrimination**							**07441121**
					GRA			
Actinomyces naeslundii		79.72	0.38	MLT 94	99			
Prevotella melaninogenica		20.28	-.01	FRU 95	1			
07441123	**Good identification**							**07441123**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07443001	**Good identification**							**07443001**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07443003	**Good identification**							**07443003**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07443021	**Good identification**							**07443021**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07443023	**Good identification**							**07443023**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07443100	**Low discrimination**							**07443100**
					GRA			
Actinomyces naeslundii		85.43	0.40	MLT 94	99			
Fusobacterium mortiferum		14.57	0.16	FRU 95	1			
07443101	**Good identification**							**07443101**
Actinomyces naeslundii		100.0	0.43	MLT 94				
07443102	**Good identification**							**07443102**
Actinomyces naeslundii		100.0	0.40	MLT 94				
07443103	**Good identification**							**07443103**
Actinomyces naeslundii		100.0	0.43	MLT 94				
07443120	**Low discrimination**							**07443120**
					GRA			
Actinomyces naeslundii		83.57	0.40	MLT 94	99			
Prevotella melaninogenica		16.43	-.01	FRU 95	1			
07443121	**Good identification**							**07443121**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		100.0	0.43	MLT 94				
07443122	**Good identification**							**07443122**
Actinomyces naeslundii		100.0	0.40	MLT 94				
07443123	**Good identification**							**07443123**
Actinomyces naeslundii		100.0	0.43	MLT 94				
07450000	**Low discrimination**							**07450000**
					COC	GRA	CAT	
Fusobacterium mortiferum		85.93	0.75		1	1	5	
Propionibacterium granulosum		13.73	0.58		1	99	90	
Gemella morbillorum		0.19	0.19		99	99	1	
07450001	**Low discrimination**							**07450001**
					GRA	CAT		
Fusobacterium mortiferum		83.03	0.40	URE 1	1	5		
Propionibacterium granulosum		13.27	0.24		99	90		
Actinomyces naeslundii		3.70	0.29		99	25		
07450002	**Low discrimination**							**07450002**
					GRA	CAT		
Fusobacterium mortiferum		72.20	0.58	BGL 10	1	5		
Propionibacterium granulosum		25.96	0.47		99	90		
Actinomyces viscosus		1.64	0.18		99	76		
07450010	**Low discrimination**							**07450010**
					GRA	CAT		
Propionibacterium granulosum		68.33	0.43	CEL 1	99	90		
Fusobacterium mortiferum		31.67	0.40	MLZ 12	1	5		
07450020	**Low discrimination**							**07450020**
					GRA	CAT		
Propionibacterium granulosum		78.96	0.47	CEL 1	99	90		
Fusobacterium mortiferum		19.96	0.40		1	5		
Actinomyces naeslundii		0.59	0.26		99	25		
07450022	**Low discrimination**							**07450022**
					GRA	CAT		
Propionibacterium granulosum		87.54	0.37	CEL 1	99	90		
Fusobacterium mortiferum		9.84	0.24		1	5		
Actinomyces naeslundii		2.63	0.26		99	25		
07450040	**Low discrimination**							**07450040**
					GRA	CAT		
Fusobacterium mortiferum		86.22	0.40	SOR 1	1	5		
Propionibacterium granulosum		13.78	0.24		99	90		
07450100	**Very good identification**							**07450100**
					GRA	CAT		
Fusobacterium mortiferum		99.88	0.81		1	5		
Propionibacterium granulosum		0.07	0.24		99	90		
Actinomyces naeslundii		0.04	0.34		99	25		
07450101	**Acceptable identification**							**07450101**
					GRA			
Fusobacterium mortiferum		94.85	0.47	URE 1	1			
Actinomyces naeslundii		5.15	0.37		99			
07450102	**Very good identification**							**07450102**
					GRA			
Fusobacterium mortiferum		97.73	0.65	BGL 10	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces viscosus		1.70	0.22		99			
Actinomyces naeslundii		0.32	0.34		99			
07450110	**Good identification**							**07450110**
					GRA	CAT		
Fusobacterium mortiferum		99.08	0.47	MLZ 1	1	5		
Propionibacterium granulosum		0.93	0.09		99	90		
07450120	**Low discrimination**							**07450120**
					GRA			
Fusobacterium mortiferum		82.61	0.47	NAG 1	1			
Prevotella melaninogenica		12.98	0.15		1			
Actinomyces naeslundii		2.99	0.34		99			
07450121	**Low discrimination**							**07450121**
					GRA			
Actinomyces naeslundii		84.33	0.37	MLT 94	99			
Fusobacterium mortiferum		15.68	0.13	GAL 96	1			
07450123	**Good identification**							**07450123**
Actinomyces naeslundii		100.0	0.37	MLT 94				
07450140	**Good identification**							**07450140**
Fusobacterium mortiferum		100.0	0.47	SOR 1				
07450200	**Low discrimination**							**07450200**
					GRA	CAT		
Fusobacterium mortiferum		67.15	0.40	RHA 1	1	5		
Propionibacterium granulosum		32.86	0.32		99	90		
07450300	**Good identification**							**07450300**
Fusobacterium mortiferum		100.0	0.47	RHA 1				
07450400	**Low discrimination**							**07450400**
					COC	GRA	CAT	
Fusobacterium mortiferum		84.47	0.40	ARA 1	1	1	5	
Propionibacterium granulosum		13.50	0.24		1	99	90	
Gemella morbillorum		2.03	0.03		99	99	1	
07450500	**Good identification**							**07450500**
					GRA			
Fusobacterium mortiferum		98.82	0.47	ARA 1	1			
Lactobacillus fermentum		1.18	0.15		99			
07451000	**Low discrimination**							**07451000**
					GRA	CAT		
Fusobacterium mortiferum		88.82	0.78		1	5		
Propionibacterium granulosum		9.46	0.58		99	90		
Lactobacillus fermentum		1.01	0.46		99	1		
07451001	**Low discrimination**							**07451001**
					GRA	CAT		
Fusobacterium mortiferum		54.97	0.43	URE 1	1	5		
Actinomyces naeslundii		39.18	0.53		99	25		
Propionibacterium granulosum		5.86	0.24		99	90		
07451002	**Low discrimination**							**07451002**
					GRA	CAT		
Fusobacterium mortiferum		70.82	0.61	BGL 10	1	5		
Propionibacterium granulosum		16.98	0.47		99	90		
Lactobacillus fermentum		7.27	0.46		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07451003	**Low discrimination**						**07451003**
					GRA		
Actinomyces naeslundii		80.57	0.53	MLT 94	99		
Fusobacterium mortiferum		12.56	0.27		1		
Actinomyces viscosus		3.86	0.06		99		
07451010	**Low discrimination**						**07451010**
					GRA	CAT	
Propionibacterium granulosum		57.83	0.43	CEL 1	99	90	
Fusobacterium mortiferum		40.21	0.43	MLZ 12	1	5	
Actinomyces naeslundii		1.01	0.28		99	25	
07451020	**Low discrimination**						**07451020**
					GRA	CAT	
Propionibacterium granulosum		54.25	0.47	CEL 1	99	90	
Fusobacterium mortiferum		20.57	0.43		1	5	
Actinomyces naeslundii		9.78	0.50		99	25	
07451021	**Good identification**						**07451021**
Actinomyces naeslundii		96.40	0.53	MLT 94			
Propionibacterium granulosum		3.60	0.13				
07451023	**Good identification**						**07451023**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07451040	**Low discrimination**						**07451040**
					GRA	CAT	
Fusobacterium mortiferum		84.01	0.43	SOR 1	1	5	
Propionibacterium granulosum		8.95	0.24		99	90	
Actinomyces naeslundii		7.04	0.37		99	25	
07451041	**Good identification**						**07451041**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07451043	**Good identification**						**07451043**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07451060	**Low discrimination**						**07451060**
Actinomyces naeslundii		75.89	0.37	MLT 94			
Propionibacterium granulosum		24.11	0.13				
07451061	**Good identification**						**07451061**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07451062	**Good identification**						**07451062**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07451063	**Good identification**						**07451063**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07451100	**Very good identification**						**07451100**
					GRA		
Fusobacterium mortiferum		98.44	0.84		1		
Lactobacillus fermentum		0.79	0.50		99		
Actinomyces naeslundii		0.58	0.57		99		
07451101	**Low discrimination**						**07451101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		53.27	0.50	URE 1	1	
Actinomyces naeslundii		46.31	0.60		99	
Lactobacillus fermentum		0.43	0.15		99	
07451102	**Low discrimination**					**07451102**
					GRA	
Fusobacterium mortiferum		84.43	0.68	BGL 10	1	
Lactobacillus fermentum		6.06	0.50		99	
Actinomyces naeslundii		4.45	0.57		99	
07451103	**Low discrimination**					**07451103**
					GRA	
Actinomyces naeslundii		85.70	0.60	MLT 94	99	
Fusobacterium mortiferum		10.95	0.33		1	
Actinomyces viscosus		2.56	0.10		99	
07451110	**Good identification**					**07451110**
					GRA	
Fusobacterium mortiferum		95.72	0.50	MLZ 1	1	
Actinomyces naeslundii		2.92	0.35		99	
Lactobacillus fermentum		0.76	0.15		99	
07451111	**Low discrimination**					**07451111**
					GRA	
Actinomyces naeslundii		81.92	0.38	MLT 94	99	
Fusobacterium mortiferum		18.08	0.16	MLZ 5	1	
07451113	**Good identification**					**07451113**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07451120	**Low discrimination**					**07451120**
					GRA	
Prevotella melaninogenica		81.57	0.50	MLT 99	1	
Fusobacterium mortiferum		7.87	0.50	CEL 10	1	
Actinomyces naeslundii		4.56	0.57		99	
07451121	**Low discrimination**					**07451121**
					GRA	
Actinomyces naeslundii		88.33	0.60	MLT 94	99	
Prevotella melaninogenica		10.64	0.15		1	
Fusobacterium mortiferum		1.03	0.16		1	
07451122	**Low discrimination**					**07451122**
					GRA	
Prevotella oralis		60.19	0.49	MLT 99	1	
Prevotella melaninogenica		20.56	0.37		1	
Actinomyces naeslundii		6.03	0.57		99	
07451123	**Good identification**					**07451123**
					GRA	
Actinomyces naeslundii		91.73	0.60	MLT 94	99	
Prevotella oralis		6.16	0.15		1	
Prevotella melaninogenica		2.11	0.03		1	
07451131	**Good identification**					**07451131**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07451132	**Low discrimination**					**07451132**
					GRA	
Parabacteroides distasonis		69.69	0.39	XYL 90	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		14.84	0.15		1		
Actinomyces naeslundii		7.75	0.35		99		
07451133	**Good identification**						**07451133**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07451140	**Acceptable identification**						**07451140**
					GRA		
Fusobacterium mortiferum		90.07	0.50	SOR 1	1		
Actinomyces naeslundii		9.21	0.44		99		
Lactobacillus fermentum		0.72	0.15		99		
07451141	**Acceptable identification**						**07451141**
					GRA		
Actinomyces naeslundii		93.82	0.47	MLT 94	99		
Fusobacterium mortiferum		6.18	0.16		1		
07451143	**Good identification**						**07451143**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07451161	**Good identification**						**07451161**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07451162	**Low discrimination**						**07451162**
					GRA		
Actinomyces naeslundii		44.98	0.44	MLT 94	99		
Prevotella oralis		25.69	0.15		1		
Lactobacillus acidophilus		9.72	0.09		99		
07451163	**Good identification**						**07451163**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07451200	**Low discrimination**						**07451200**
					GRA	CAT	
Fusobacterium mortiferum		75.40	0.43	RHA 1	1	5	
Propionibacterium granulosum		24.60	0.32		99	90	
07451300	**Good identification**						**07451300**
					GRA		
Fusobacterium mortiferum		97.97	0.50	RHA 1	1		
Lactobacillus fermentum		0.78	0.15		99		
Prevotella oralis		0.67	-.01		1		
07451322	**Low discrimination**						**07451322**
Prevotella oralis		61.25	0.42	MLT 99			
Parabacteroides distasonis		37.78	0.52				
Prevotella melaninogenica		0.49	0.03				
07451332	**Good identification**						**07451332**
Parabacteroides distasonis		97.05	0.47	XYL 90			
Prevotella oralis		2.95	0.08				
07451400	**Low discrimination**						**07451400**
					GRA	CAT	
Lactobacillus fermentum		50.52	0.46	CEL 1	99	1	
Fusobacterium mortiferum		44.72	0.43		1	5	
Propionibacterium granulosum		4.76	0.24		99	90	
07451402	**Low discrimination**						**07451402**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Lactobacillus fermentum		89.13	0.46	CEL 1	99	1		
Fusobacterium mortiferum		8.77	0.27		1	5		
Propionibacterium granulosum		2.10	0.13		99	90		
07451420	**Acceptable identification**							**07451420**
					CAT			
Lactobacillus fermentum		93.39	0.38	CEL 1	1			
Propionibacterium granulosum		6.61	0.13		90			
07451422	**Good identification**							**07451422**
Lactobacillus fermentum		100.0	0.38	CEL 1				
07451500	**Low discrimination**							**07451500**
					GRA			
Fusobacterium mortiferum		55.41	0.50	ARA 1	1			
Lactobacillus fermentum		43.77	0.50		99			
Bifidobacterium longum subsp. longum		0.50	0.09		99			
07451502	**Low discrimination**							**07451502**
					GRA			
Lactobacillus fermentum		79.34	0.50	CEL 1	99			
Fusobacterium mortiferum		11.16	0.33		1			
Bifidobacterium adolescentis		8.00	0.27		99			
07451520	**Low discrimination**							**07451520**
					GRA			
Lactobacillus fermentum		61.71	0.41	CEL 1	99			
Prevotella melaninogenica		24.55	0.15		1			
Bifidobacterium longum subsp. longum		8.51	0.20		99			
07451522	**Low discrimination**							**07451522**
					GRA			
Parabacteroides distasonis		43.81	0.40	XYL 90	1			
Lactobacillus fermentum		36.94	0.41		99			
Prevotella oralis		8.19	0.15		1			
07451532	**Low discrimination**							**07451532**
					GRA			
Parabacteroides distasonis		73.11	0.35	XYL 90	1			
Bifidobacterium longum subsp. longum		26.89	0.24		99			
07451722	**Good identification**							**07451722**
Parabacteroides distasonis		97.40	0.48	XYL 90				
Prevotella oralis		2.60	0.08					
07451732	**Good identification**							**07451732**
Parabacteroides distasonis		97.60	0.43	XYL 90				
Clostridium clostridioforme		2.40	0.10					
07452000	**Acceptable identification**							**07452000**
					GRA	SPO	CAT	
Propionibacterium granulosum		91.09	0.56	CEL 1	99	1	90	
Fusobacterium mortiferum		7.63	0.40		1	1	5	
Clostridium innocuum		0.67	0.10		99	99	1	
07452002	**Acceptable identification**							**07452002**
					GRA	SPO	CAT	
Propionibacterium granulosum		92.09	0.45	CEL 1	99	1	90	
Fusobacterium mortiferum		3.43	0.24		1	1	5	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium innocuum		2.71	0.10		99	99	1	
07452010	**Good identification**							**07452010**
					SPO	CAT		
Propionibacterium granulosum		97.27	0.41	CEL 1	1	90		
Clostridium difficile		2.73	0.02	MLZ 12	99	1		
07452020	**Good identification**							**07452020**
Propionibacterium granulosum		98.10	0.45	CEL 1				
Actinomyces naeslundii		1.90	0.31					
07452022	**Acceptable identification**							**07452022**
Propionibacterium granulosum		92.83	0.35	CEL 1				
Actinomyces naeslundii		7.17	0.31					
07452100	**Low discrimination**							**07452100**
					GRA	CAT		
Fusobacterium mortiferum		89.13	0.47	MAN 1	1	5		
Actinomyces naeslundii		6.27	0.39		99	25		
Propionibacterium granulosum		4.61	0.21		99	90		
07452101	**Acceptable identification**							**07452101**
					GRA			
Actinomyces naeslundii		91.26	0.42	MLT 94	99			
Fusobacterium mortiferum		8.74	0.13	GAL 96	1			
07452103	**Good identification**							**07452103**
Actinomyces naeslundii		100.0	0.42	MLT 94				
07452120	**Low discrimination**							**07452120**
					GRA	CAT		
Actinomyces naeslundii		75.34	0.39	MLT 94	99	25		
Propionibacterium granulosum		13.84	0.11	GAL 96	99	90		
Fusobacterium mortiferum		10.82	0.13		1	5		
07452121	**Good identification**							**07452121**
Actinomyces naeslundii		100.0	0.42	MLT 94				
07452122	**Good identification**							**07452122**
Actinomyces naeslundii		100.0	0.39	MLT 94				
07452123	**Good identification**							**07452123**
Actinomyces naeslundii		100.0	0.42	MLT 94				
07453000	**Low discrimination**							**07453000**
					GRA	SPO	CAT	
Propionibacterium granulosum		72.40	0.56	CEL 1	99	1	90	
Clostridium innocuum		10.10	0.32		99	99	1	
Fusobacterium mortiferum		9.10	0.43		1	1	5	
07453001	**Good identification**							**07453001**
Actinomyces naeslundii		94.51	0.57	MLT 94				
Propionibacterium granulosum		5.49	0.21					
07453002	**Low discrimination**							**07453002**
					SPO	CAT		
Propionibacterium granulosum		48.13	0.45	CEL 1	1	90		
Clostridium innocuum		26.86	0.32		99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		22.32	0.54		1	25	
07453003	**Very good identification**						**07453003**
Actinomyces naeslundii		98.57	0.57	MLT 94			
Propionibacterium granulosum		1.43	0.11				
07453010	**Acceptable identification**						**07453010**
					SPO	CAT	
Propionibacterium granulosum		91.97	0.41	CEL 1	1	90	
Actinomyces naeslundii		4.11	0.33	MLZ 12	1	25	
Clostridium innocuum		3.92	0.08		99	1	
07453011	**Good identification**						**07453011**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07453013	**Good identification**						**07453013**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07453020	**Low discrimination**						**07453020**
Propionibacterium granulosum		68.32	0.45	CEL 1			
Actinomyces naeslundii		31.68	0.54				
07453021	**Very good identification**						**07453021**
Actinomyces naeslundii		98.57	0.57	MLT 94			
Propionibacterium granulosum		1.43	0.11				
07453022	**Low discrimination**						**07453022**
Actinomyces naeslundii		64.97	0.54	MLT 94			
Propionibacterium granulosum		35.03	0.35				
07453023	**Very good identification**						**07453023**
Actinomyces naeslundii		100.0	0.57	MLT 94			
07453031	**Good identification**						**07453031**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07453033	**Good identification**						**07453033**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07453040	**Low discrimination**						**07453040**
Actinomyces naeslundii		66.94	0.42	MLT 94			
Propionibacterium granulosum		33.06	0.21				
07453041	**Good identification**						**07453041**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07453042	**Low discrimination**						**07453042**
Actinomyces naeslundii		89.01	0.42	MLT 94			
Propionibacterium granulosum		10.99	0.11				
07453043	**Good identification**						**07453043**
Actinomyces naeslundii		100.0	0.45	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07453060	**Low discrimination**						**07453060**
Actinomyces naeslundii		89.01	0.42	MLT 94			
Propionibacterium granulosum		10.99	0.11				
07453061	**Good identification**						**07453061**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07453062	**Good identification**						**07453062**
Actinomyces naeslundii		100.0	0.42	MLT 94			
07453063	**Good identification**						**07453063**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07453100	**Low discrimination**						**07453100**
					GRA	CAT	
Actinomyces naeslundii		51.91	0.62	MLT 94	99	25	
Fusobacterium mortiferum		46.14	0.50		1	5	
Propionibacterium granulosum		1.59	0.21		99	90	
07453101	**Very good identification**						**07453101**
					GRA		
Actinomyces naeslundii		99.41	0.65	MLT 94	99		
Fusobacterium mortiferum		0.60	0.16		1		
07453102	**Low discrimination**						**07453102**
					GRA		
Actinomyces naeslundii		87.62	0.62	MLT 94	99		
Fusobacterium mortiferum		8.65	0.33		1		
Bifidobacterium adolescentis		0.76	0.11		99		
07453103	**Very good identification**						**07453103**
Actinomyces naeslundii		100.0	0.65	MLT 94			
07453110	**Low discrimination**						**07453110**
					GRA		
Actinomyces naeslundii		85.43	0.40	MLT 94	99		
Fusobacterium mortiferum		14.57	0.16	MLZ 5	1		
07453111	**Good identification**						**07453111**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07453112	**Low discrimination**						**07453112**
Actinomyces naeslundii		83.26	0.40	MLT 94			
Bifidobacterium adolescentis		16.74	0.13	MLZ 5			
07453113	**Good identification**						**07453113**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07453120	**Good identification**						**07453120**
					GRA		
Actinomyces naeslundii		90.11	0.62	MLT 94	99		
Prevotella melaninogenica		8.39	0.15		1		
Fusobacterium mortiferum		0.81	0.16		1		
07453121	**Very good identification**						**07453121**
Actinomyces naeslundii		100.0	0.65	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07453122	**Good identification**							**07453122**
					GRA			
Actinomyces naeslundii		91.67	0.62	MLT 94	99			
Prevotella oralis		4.76	0.15		1			
Prevotella melaninogenica		1.63	0.03		1			
07453123	**Very good identification**							**07453123**
Actinomyces naeslundii		100.0	0.65	MLT 94				
07453130	**Good identification**							**07453130**
Actinomyces naeslundii		100.0	0.40	MLT 94				
07453131	**Good identification**							**07453131**
Actinomyces naeslundii		100.0	0.43	MLT 94				
07453132	**Good identification**							**07453132**
Actinomyces naeslundii		100.0	0.40	MLT 94				
07453133	**Good identification**							**07453133**
Actinomyces naeslundii		100.0	0.43	MLT 94				
07453140	**Good identification**							**07453140**
					GRA			
Actinomyces naeslundii		95.16	0.49	MLT 94	99			
Fusobacterium mortiferum		4.84	0.16		1			
07453141	**Good identification**							**07453141**
Actinomyces naeslundii		100.0	0.52	MLT 94				
07453142	**Good identification**							**07453142**
Actinomyces naeslundii		97.39	0.49	MLT 94				
Propionibacterium propionicum		2.61	0.12					
07453143	**Good identification**							**07453143**
Actinomyces naeslundii		100.0	0.52	MLT 94				
07453160	**Good identification**							**07453160**
Actinomyces naeslundii		100.0	0.49	MLT 94				
07453161	**Good identification**							**07453161**
Actinomyces naeslundii		100.0	0.52	MLT 94				
07453162	**Good identification**							**07453162**
Actinomyces naeslundii		100.0	0.49	MLT 94				
07453163	**Good identification**							**07453163**
Actinomyces naeslundii		100.0	0.52	MLT 94				
07454000	**Good identification**							**07454000**
					COC	GRA	CAT	
Fusobacterium mortiferum		97.59	0.55	XYL 7	1	1	5	
Propionibacterium granulosum		2.09	0.24		1	99	90	
Gemella morbillorum		0.32	0.03		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07454002	**Good identification**						**07454002**
					GRA	CAT	
Fusobacterium mortiferum		95.40	0.39	XYL 7	1	5	
Propionibacterium granulosum		4.60	0.13	BGL 10	99	90	
07454100	**Very good identification**						**07454100**
Fusobacterium mortiferum		100.0	0.62	XYL 7			
07454102	**Good identification**						**07454102**
Fusobacterium mortiferum		100.0	0.45	XYL 7			
07455000	**Good identification**						**07455000**
					GRA	CAT	
Fusobacterium mortiferum		90.63	0.58	XYL 7	1	5	
Lactobacillus fermentum		8.07	0.42		99	1	
Propionibacterium granulosum		1.30	0.24		99	90	
07455002	**Low discrimination**						**07455002**
					GRA	CAT	
Fusobacterium mortiferum		54.54	0.42	XYL 7	1	5	
Lactobacillus fermentum		43.71	0.42	BGL 10	99	1	
Propionibacterium granulosum		1.76	0.13		99	90	
07455100	**Good identification**						**07455100**
					GRA		
Fusobacterium mortiferum		93.96	0.65	XYL 7	1		
Lactobacillus fermentum		5.85	0.46		99		
Actinomyces israelii		0.12	0.17		99		
07455102	**Low discrimination**						**07455102**
					GRA		
Fusobacterium mortiferum		55.80	0.48	XYL 7	1		
Lactobacillus fermentum		31.27	0.46	BGL 10	99		
Bifidobacterium adolescentis		5.37	0.27		99		
07455120	**Low discrimination**						**07455120**
					GRA		
Lactobacillus fermentum		31.42	0.37	CEL 1	99		
Prevotella melaninogenica		21.28	0.15		1		
Fusobacterium mortiferum		15.29	0.31		1		
07455122	**Low discrimination**						**07455122**
					CAT		
Parabacteroides distasonis		72.17	0.60		76		
Bacteroides fragilis		13.24	0.25		99		
Prevotella oralis		11.25	0.33		1		
07455132	**Good identification**						**07455132**
					CAT		
Parabacteroides distasonis		99.37	0.55		76		
Bacteroides fragilis		0.34	-.09		99		
Prevotella oralis		0.29	-.01		1		
07455322	**Very good identification**						**07455322**
					CAT		
Parabacteroides distasonis		97.76	0.68		76		
Prevotella oralis		2.18	0.27		1		
Bacteroides fragilis		0.06	-.09		99		
07455332	**Very good identification**						**07455332**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		99.89	0.63			
Clostridium clostridioforme		0.11	0.07			
07455400	**Low discrimination**					**07455400**
					GRA	
Lactobacillus fermentum		89.81	0.42	CEL 1	99	
Fusobacterium mortiferum		10.19	0.24		1	
07455402	**Good identification**					**07455402**
Lactobacillus fermentum		98.14	0.42	CEL 1		
Bifidobacterium adolescentis		1.86	0.07			
07455500	**Low discrimination**					**07455500**
					GRA	
Lactobacillus fermentum		78.78	0.46	CEL 1	99	
Fusobacterium mortiferum		12.78	0.31		1	
Bifidobacterium longum subsp. longum		4.63	0.17		99	
07455502	**Low discrimination**					**07455502**
					GRA	
Bifidobacterium adolescentis		49.71	0.41	MLT 99	99	
Lactobacillus fermentum		43.26	0.46		99	
Parabacteroides distasonis		2.65	0.22		1	
07455512	**Low discrimination**					**07455512**
Bifidobacterium adolescentis		89.22	0.43	MLT 99		
Bifidobacterium longum subsp. longum		6.35	0.21			
Clostridium clostridioforme		2.34	0.12			
07455522	**Acceptable identification**					**07455522**
					GRA	
Parabacteroides distasonis		90.37	0.56		1	
Lactobacillus fermentum		4.97	0.37		99	
Bifidobacterium longum subsp. longum		3.50	0.28		99	
07455532	**Low discrimination**					**07455532**
					GRA	
Parabacteroides distasonis		87.85	0.51		1	
Bifidobacterium longum subsp. longum		10.77	0.32		99	
Clostridium clostridioforme		0.99	0.12		50	
07455720	**Good identification**					**07455720**
Bacteroides vulgatus		97.06	0.38	MLT 99		
Parabacteroides distasonis		2.94	0.30	CEL 5		
07455722	**Very good identification**					**07455722**
					CAT	
Parabacteroides distasonis		98.66	0.64		76	
Bacteroides thetaiotaomicron		0.44	0.08		89	
Bacteroides vulgatus		0.33	0.04		3	
07455732	**Very good identification**					**07455732**
Parabacteroides distasonis		97.01	0.60			
Clostridium clostridioforme		2.68	0.27			
Bacteroides thetaiotaomicron		0.32	0.01			
07457502	**Low discrimination**					**07457502**
Bifidobacterium adolescentis		87.10	0.40	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		12.90	0.32			
07457512	**Good identification**					**07457512**
Bifidobacterium adolescentis		98.22	0.41	MLT 99		
Actinomyces israelii		1.78	0.18			
07460100	**Good identification**					**07460100**
Fusobacterium mortiferum		100.0	0.38	FRU 99		
07461001	**Good identification**					**07461001**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07461003	**Good identification**					**07461003**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07461021	**Good identification**					**07461021**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07461023	**Good identification**					**07461023**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07461100	**Low discrimination**					**07461100**
					GRA	
Fusobacterium mortiferum		83.46	0.41	FRU 99	1	
Actinomyces naeslundii		16.54	0.40		99	
07461101	**Good identification**					**07461101**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07461102	**Low discrimination**					**07461102**
					GRA	
Actinomyces naeslundii		60.15	0.40	MLT 94	99	
Fusobacterium mortiferum		33.71	0.24	FRU 95	1	
Lactobacillus acidophilus		6.14	-.01		99	
07461103	**Good identification**					**07461103**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07461120	**Low discrimination**					**07461120**
					GRA	
Actinomyces naeslundii		83.57	0.40	MLT 94	99	
Prevotella melaninogenica		16.43	-.01	FRU 95	1	
07461121	**Good identification**					**07461121**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07461122	**Low discrimination**					**07461122**
					GRA	
Actinomyces naeslundii		66.75	0.40	MLT 94	99	
Parabacteroides distasonis		26.43	0.20	FRU 95	1	
Lactobacillus acidophilus		6.82	-.01		99	
07461123	**Good identification**					**07461123**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07463001	**Good identification**					**07463001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	MLT 94			
07463002	**Low discrimination**						**07463002**
					SPO	CAT	
Actinomyces naeslundii		53.84	0.38	MLT 94	1	25	
Propionibacterium granulosum		34.44	0.19	FRU 95	1	90	
Clostridium innocuum		11.73	0.02		99	1	
07463003	**Good identification**						**07463003**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07463020	**Low discrimination**						**07463020**
Actinomyces naeslundii		60.99	0.38	MLT 94			
Propionibacterium granulosum		39.01	0.19	FRU 95			
07463021	**Good identification**						**07463021**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07463022	**Low discrimination**						**07463022**
Actinomyces naeslundii		86.22	0.38	MLT 94			
Propionibacterium granulosum		13.78	0.09	FRU 95			
07463023	**Good identification**						**07463023**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07463100	**Good identification**						**07463100**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07463101	**Good identification**						**07463101**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07463102	**Good identification**						**07463102**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07463103	**Good identification**						**07463103**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07463120	**Good identification**						**07463120**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07463121	**Good identification**						**07463121**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07463122	**Good identification**						**07463122**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07463123	**Good identification**						**07463123**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07463141	**Good identification**						**07463141**
Actinomyces naeslundii		100.0	0.35	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07463143	**Good identification**						**07463143**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07463161	**Good identification**						**07463161**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07463163	**Good identification**						**07463163**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07465122	**Good identification**						**07465122**
Parabacteroides distasonis		100.0	0.37	FRU 99			
07465322	**Good identification**						**07465322**
Parabacteroides distasonis		100.0	0.45	FRU 99			
07465332	**Good identification**						**07465332**
Parabacteroides distasonis		100.0	0.40	FRU 99			
07465722	**Good identification**						**07465722**
Parabacteroides distasonis		100.0	0.41	FRU 99			
07465732	**Good identification**						**07465732**
Parabacteroides distasonis		95.75	0.37	FRU 99			
Clostridium clostridioforme		4.25	0.08				
07470000	**Low discrimination**						**07470000**
					GRA	CAT	
Propionibacterium granulosum		61.57	0.66	CEL 1	99	90	
Fusobacterium mortiferum		38.36	0.66		1	5	
Actinomyces naeslundii		0.08	0.31		99	25	
07470002	**Low discrimination**						**07470002**
					GRA	CAT	
Propionibacterium granulosum		77.92	0.56	CEL 1	99	90	
Fusobacterium mortiferum		21.58	0.49		1	5	
Actinomyces naeslundii		0.38	0.31		99	25	
07470010	**Good identification**						**07470010**
					GRA	CAT	
Propionibacterium granulosum		95.59	0.51	CEL 1	99	90	
Fusobacterium mortiferum		4.41	0.31	MLZ 12	1	5	
07470012	**Good identification**						**07470012**
					GRA	CAT	
Propionibacterium granulosum		97.99	0.41	CEL 1	99	90	
Fusobacterium mortiferum		2.01	0.15	MLZ 12	1	5	
07470020	**Good identification**						**07470020**
					GRA	CAT	
Propionibacterium granulosum		97.08	0.56	CEL 1	99	90	
Fusobacterium mortiferum		2.44	0.31		1	5	
Actinomyces naeslundii		0.47	0.31		99	25	
07470022	**Good identification**						**07470022**
					GRA	CAT	
Propionibacterium granulosum		97.03	0.45	CEL 1	99	90	
Actinomyces naeslundii		1.89	0.31		99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		1.09	0.15		1	5	
07470030	**Good identification**						**07470030**
Propionibacterium granulosum		100.0	0.41	CEL 1			
07470100	**Very good identification**						**07470100**
					GRA	CAT	
Fusobacterium mortiferum		99.08	0.72		1	5	
Propionibacterium granulosum		0.69	0.32		99	90	
Actinomyces naeslundii		0.24	0.39		99	25	
07470101	**Low discrimination**						**07470101**
					GRA		
Fusobacterium mortiferum		73.90	0.38	URE 1	1		
Actinomyces naeslundii		26.10	0.42		99		
07470102	**Good identification**						**07470102**
					GRA	CAT	
Fusobacterium mortiferum		95.77	0.56	BGL 10	1	5	
Actinomyces naeslundii		2.05	0.39		99	25	
Propionibacterium granulosum		1.50	0.21		99	90	
07470103	**Low discrimination**						**07470103**
					GRA		
Actinomyces naeslundii		76.07	0.42	MLT 94	99		
Fusobacterium mortiferum		23.94	0.21	GAL 96	1		
07470110	**Acceptable identification**						**07470110**
					GRA	CAT	
Fusobacterium mortiferum		91.43	0.38	MLZ 1	1	5	
Propionibacterium granulosum		8.58	0.17		99	90	
07470120	**Low discrimination**						**07470120**
					GRA	CAT	
Fusobacterium mortiferum		71.05	0.38	NAG 1	1	5	
Actinomyces naeslundii		16.73	0.39		99	25	
Propionibacterium granulosum		12.22	0.21		99	90	
07470121	**Good identification**						**07470121**
Actinomyces naeslundii		100.0	0.42	MLT 94			
07470122	**Low discrimination**						**07470122**
					GRA		
Actinomyces naeslundii		43.79	0.39	MLT 94	99		
Parabacteroides distasonis		21.90	0.20	GAL 96	1		
Fusobacterium mortiferum		20.67	0.21		1		
07470123	**Good identification**						**07470123**
Actinomyces naeslundii		100.0	0.42	MLT 94			
07470140	**Good identification**						**07470140**
					GRA		
Fusobacterium mortiferum		96.01	0.38	SOR 1	1		
Actinomyces naeslundii		3.99	0.26		99		
07470200	**Low discrimination**						**07470200**
					GRA	CAT	
Propionibacterium granulosum		83.09	0.40	CEL 1	99	90	
Fusobacterium mortiferum		16.91	0.31	RHA 3	1	5	
07470300	**Good identification**						**07470300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.38	RHA 1			
07470500	**Good identification**						**07470500**
Fusobacterium mortiferum		100.0	0.38	ARA 1			
07471000	**Low discrimination**						**07471000**
					GRA	CAT	
Propionibacterium granulosum		50.75	0.66	CEL 1	99	90	
Fusobacterium mortiferum		47.43	0.69		1	5	
Actinomyces naeslundii		1.48	0.54		99	25	
07471001	**Low discrimination**						**07471001**
					GRA	CAT	
Actinomyces naeslundii		69.11	0.57	MLT 94	99	25	
Propionibacterium granulosum		15.97	0.32		99	90	
Fusobacterium mortiferum		14.92	0.34		1	5	
07471002	**Low discrimination**						**07471002**
					GRA	CAT	
Propionibacterium granulosum		63.88	0.56	CEL 1	99	90	
Fusobacterium mortiferum		26.53	0.52		1	5	
Actinomyces naeslundii		7.45	0.54		99	25	
07471003	**Good identification**						**07471003**
					GRA	CAT	
Actinomyces naeslundii		92.44	0.57	MLT 94	99	25	
Propionibacterium granulosum		5.34	0.21		99	90	
Fusobacterium mortiferum		2.22	0.18		1	5	
07471010	**Acceptable identification**						**07471010**
					GRA	CAT	
Propionibacterium granulosum		91.15	0.51	CEL 1	99	90	
Fusobacterium mortiferum		6.31	0.34	MLZ 12	1	5	
Lactobacillus jensenii		1.52	0.14		99	1	
07471011	**Low discrimination**						**07471011**
Actinomyces naeslundii		62.56	0.36	MLT 94			
Propionibacterium granulosum		37.44	0.17	MLZ 5			
07471012	**Low discrimination**						**07471012**
					CAT		
Propionibacterium granulosum		89.27	0.41	CEL 1	90		
Actinomyces naeslundii		4.02	0.33	MLZ 12	25		
Lactobacillus jensenii		3.96	0.11		1		
07471013	**Good identification**						**07471013**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07471020	**Low discrimination**						**07471020**
					GRA	CAT	
Propionibacterium granulosum		85.26	0.56	CEL 1	99	90	
Actinomyces naeslundii		9.94	0.54		99	25	
Fusobacterium mortiferum		3.22	0.34		1	5	
07471021	**Good identification**						**07471021**
Actinomyces naeslundii		94.54	0.57	MLT 94			
Propionibacterium granulosum		5.46	0.21				
07471022	**Low discrimination**						**07471022**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		60.96	0.45	CEL 1	90		
Actinomyces naeslundii		28.43	0.54		25		
Lactobacillus acidophilus		3.75	0.15		1		
07471023	**Very good identification**						**07471023**
Actinomyces naeslundii		98.58	0.57	MLT 94			
Propionibacterium granulosum		1.42	0.11				
07471030	**Acceptable identification**						**07471030**
					CAT		
Propionibacterium granulosum		94.19	0.41	CEL 1	90		
Actinomyces naeslundii		4.24	0.33	MLZ 12	25		
Lactobacillus jensenii		1.57	0.04		1		
07471031	**Good identification**						**07471031**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07471033	**Good identification**						**07471033**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07471041	**Good identification**						**07471041**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07471042	**Low discrimination**						**07471042**
					GRA	CAT	
Actinomyces naeslundii		55.96	0.42	MLT 94	99	25	
Propionibacterium granulosum		27.47	0.21		99	90	
Fusobacterium mortiferum		11.41	0.18		1	5	
07471043	**Good identification**						**07471043**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07471060	**Low discrimination**						**07471060**
					CAT		
Actinomyces naeslundii		62.50	0.42	MLT 94	25		
Propionibacterium granulosum		30.68	0.21		90		
Clostridium perfringens		6.82	0.03		1		
07471061	**Good identification**						**07471061**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07471062	**Low discrimination**						**07471062**
					CAT		
Actinomyces naeslundii		82.31	0.42	MLT 94	25		
Propionibacterium granulosum		10.10	0.11		90		
Lactobacillus acidophilus		7.59	-.01		1		
07471063	**Good identification**						**07471063**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07471100	**Very good identification**						**07471100**
					GRA	CAT	
Fusobacterium mortiferum		95.63	0.75		1	5	
Actinomyces naeslundii		3.64	0.62		99	25	
Propionibacterium granulosum		0.44	0.32		99	90	
07471101	**Low discrimination**						**07471101**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		84.96	0.65	MLT 94	99		
Fusobacterium mortiferum		15.04	0.41		1		
07471102	**Low discrimination**						**07471102**
					GRA		
Fusobacterium mortiferum		64.18	0.59	BGL 10	1		
Actinomyces naeslundii		21.97	0.62		99		
Lactobacillus acidophilus		11.70	0.33		99		
07471103	**Very good identification**						**07471103**
					GRA		
Actinomyces naeslundii		97.73	0.65	MLT 94	99		
Fusobacterium mortiferum		1.92	0.24		1		
Lactobacillus acidophilus		0.35	-.01		99		
07471110	**Low discrimination**						**07471110**
					GRA	CAT	
Fusobacterium mortiferum		79.32	0.41	MLZ 1	1	5	
Actinomyces naeslundii		15.72	0.40		99	25	
Propionibacterium granulosum		4.96	0.17		99	90	
07471111	**Good identification**						**07471111**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07471112	**Low discrimination**						**07471112**
					GRA		
Actinomyces naeslundii		46.63	0.40	MLT 94	99		
Fusobacterium mortiferum		26.13	0.24	MLZ 5	1		
Parabacteroides distasonis		9.94	0.16		1		
07471113	**Good identification**						**07471113**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07471120	**Low discrimination**						**07471120**
					GRA		
Actinomyces naeslundii		58.22	0.62	MLT 94	99		
Fusobacterium mortiferum		15.46	0.41		1		
Clostridium perfringens		15.17	0.30		99		
07471121	**Very good identification**						**07471121**
Actinomyces naeslundii		100.0	0.65	MLT 94			
07471122	**Low discrimination**						**07471122**
					GRA		
Parabacteroides distasonis		49.88	0.55	XYL 90	1		
Actinomyces naeslundii		24.18	0.62		99		
Lactobacillus acidophilus		12.87	0.33		99		
07471123	**Very good identification**						**07471123**
					GRA		
Actinomyces naeslundii		98.28	0.65	MLT 94	99		
Parabacteroides distasonis		1.37	0.20		1		
Lactobacillus acidophilus		0.35	-.01		99		
07471130	**Low discrimination**						**07471130**
					GRA		
Actinomyces naeslundii		82.43	0.40	MLT 94	99		
Parabacteroides distasonis		17.57	0.16	MLZ 5	1		
07471131	**Good identification**						**07471131**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.43	MLT 94			
07471132	**Good identification**						**07471132**
					GRA		
Parabacteroides distasonis		95.04	0.50	XYL 90	1		
Actinomyces naeslundii		4.50	0.40		99		
Lactobacillus acidophilus		0.46	-.01		99		
07471133	**Low discrimination**						**07471133**
					GRA		
Actinomyces naeslundii		87.56	0.43	MLT 94	99		
Parabacteroides distasonis		12.44	0.16	MLZ 5	1		
07471140	**Low discrimination**						**07471140**
					GRA		
Fusobacterium mortiferum		60.07	0.41	SOR 1	1		
Actinomyces naeslundii		39.93	0.49		99		
07471141	**Good identification**						**07471141**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07471142	**Low discrimination**						**07471142**
					GRA		
Actinomyces naeslundii		64.94	0.49	MLT 94	99		
Lactobacillus acidophilus		24.21	0.18		99		
Fusobacterium mortiferum		10.86	0.24		1		
07471143	**Good identification**						**07471143**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07471160	**Low discrimination**						**07471160**
Actinomyces naeslundii		78.05	0.49	MLT 94			
Clostridium perfringens		21.95	0.18				
07471161	**Good identification**						**07471161**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07471162	**Low discrimination**						**07471162**
Actinomyces naeslundii		61.61	0.49	MLT 94			
Lactobacillus acidophilus		22.97	0.18				
Clostridium perfringens		8.15	0.12				
07471163	**Good identification**						**07471163**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07471200	**Low discrimination**						**07471200**
					GRA	CAT	
Propionibacterium granulosum		76.62	0.40	CEL 1	99	90	
Fusobacterium mortiferum		23.39	0.34	RHA 3	1	5	
07471300	**Good identification**						**07471300**
					GRA		
Fusobacterium mortiferum		96.34	0.41	RHA 1	1		
Actinomyces naeslundii		3.67	0.28		99		
07471322	**Very good identification**						**07471322**
					SPO		
Parabacteroides distasonis		96.30	0.63	XYL 90	1		
Prevotella oralis		2.19	0.22		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		0.73	0.14		92		
07471332	**Very good identification**						**07471332**
Parabacteroides distasonis		99.85	0.58	XYL 90			
Clostridium clostridioforme		0.15	0.04				
07471500	**Acceptable identification**						**07471500**
					GRA		
Fusobacterium mortiferum		93.92	0.41	ARA 1	1		
Actinomyces naeslundii		3.57	0.28		99		
Lactobacillus fermentum		2.51	0.15		99		
07471522	**Good identification**						**07471522**
					GRA		
Parabacteroides distasonis		98.17	0.51	XYL 90	1		
Actinomyces naeslundii		0.78	0.28		99		
Bifidobacterium longum subsp. longum		0.63	0.09		99		
07471532	**Good identification**						**07471532**
					GRA		
Parabacteroides distasonis		96.49	0.46	XYL 90	1		
Bifidobacterium longum subsp. longum		1.95	0.13		99		
Clostridium clostridioforme		1.56	0.10		50		
07471722	**Very good identification**						**07471722**
Parabacteroides distasonis		99.05	0.59	XYL 90			
Bacteroides thetaiotaomicron		0.49	0.04				
Clostridium clostridioforme		0.46	0.14				
07471732	**Good identification**						**07471732**
Parabacteroides distasonis		95.87	0.55	XYL 90			
Clostridium clostridioforme		3.79	0.25				
Bacteroides thetaiotaomicron		0.34	-.03				
07472000	**Very good identification**						**07472000**
					GRA	CAT	
Propionibacterium granulosum		98.78	0.64	CEL 1	99	90	
Fusobacterium mortiferum		0.82	0.31		1	5	
Actinomyces naeslundii		0.31	0.36		99	25	
07472002	**Good identification**						**07472002**
					GRA	CAT	
Propionibacterium granulosum		98.07	0.53	CEL 1	99	90	
Actinomyces naeslundii		1.23	0.36		99	25	
Fusobacterium mortiferum		0.36	0.15		1	5	
07472003	**Low discrimination**						**07472003**
Actinomyces naeslundii		64.99	0.39	MLT 94			
Propionibacterium granulosum		35.01	0.19	GAL 96			
07472010	**Good identification**						**07472010**
Propionibacterium granulosum		100.0	0.49	CEL 1			
07472012	**Good identification**						**07472012**
Propionibacterium granulosum		100.0	0.39	CEL 1			
07472020	**Good identification**						**07472020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		98.77	0.53	CEL 1			
Actinomyces naeslundii		1.24	0.36				
07472021	**Low discrimination**						**07472021**
Actinomyces naeslundii		64.99	0.39	MLT 94			
Propionibacterium granulosum		35.01	0.19	GAL 96			
07472022	**Good identification**						**07472022**
Propionibacterium granulosum		95.24	0.43	CEL 1			
Actinomyces naeslundii		4.76	0.36				
07472023	**Low discrimination**						**07472023**
Actinomyces naeslundii		88.13	0.39	MLT 94			
Propionibacterium granulosum		11.87	0.09	GAL 96			
07472030	**Good identification**						**07472030**
Propionibacterium granulosum		100.0	0.39	CEL 1			
07472100	**Low discrimination**						**07472100**
					GRA	CAT	
Fusobacterium mortiferum		50.61	0.38	MAN 1	1	5	
Propionibacterium granulosum		26.26	0.30		99	90	
Actinomyces naeslundii		23.13	0.44		99	25	
07472101	**Good identification**						**07472101**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07472102	**Low discrimination**						**07472102**
					GRA	CAT	
Actinomyces naeslundii		65.49	0.44	MLT 94	99	25	
Propionibacterium granulosum		18.59	0.19	GAL 96	99	90	
Fusobacterium mortiferum		15.92	0.21		1	5	
07472103	**Good identification**						**07472103**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07472120	**Low discrimination**						**07472120**
Actinomyces naeslundii		77.89	0.44	MLT 94			
Propionibacterium granulosum		22.11	0.19	GAL 96			
07472121	**Good identification**						**07472121**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07472122	**Acceptable identification**						**07472122**
Actinomyces naeslundii		93.37	0.44	MLT 94			
Propionibacterium granulosum		6.63	0.09	GAL 96			
07472123	**Good identification**						**07472123**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07472200	**Good identification**						**07472200**
Propionibacterium granulosum		100.0	0.38	CEL 1			
07473000	**Good identification**						**07473000**
					SPO	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		90.59	0.64	CEL 1	1	90	
Actinomyces naeslundii		6.80	0.59		1	25	
Clostridium innocuum		1.48	0.24		99	1	
07473001	**Good identification**						**07473001**
Actinomyces naeslundii		91.76	0.62	MLT 94			
Propionibacterium granulosum		8.24	0.30				
07473002	**Low discrimination**						**07473002**
					SPO	CAT	
Propionibacterium granulosum		72.81	0.53	CEL 1	1	90	
Actinomyces naeslundii		21.84	0.59		1	25	
Clostridium innocuum		4.76	0.24		99	1	
07473003	**Very good identification**						**07473003**
Actinomyces naeslundii		97.81	0.62	MLT 94			
Propionibacterium granulosum		2.20	0.19				
07473010	**Good identification**						**07473010**
					SPO	CAT	
Propionibacterium granulosum		96.72	0.49	CEL 1	1	90	
Actinomyces naeslundii		2.80	0.38	MLZ 12	1	25	
Clostridium innocuum		0.48	0.00		99	1	
07473011	**Low discrimination**						**07473011**
Actinomyces naeslundii		81.13	0.41	MLT 94			
Propionibacterium granulosum		18.87	0.15	MLZ 5			
07473012	**Low discrimination**						**07473012**
					SPO	CAT	
Propionibacterium granulosum		88.05	0.39	CEL 1	1	90	
Actinomyces naeslundii		10.20	0.38	MLZ 12	1	25	
Clostridium innocuum		1.76	0.00		99	1	
07473013	**Good identification**						**07473013**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07473020	**Low discrimination**						**07473020**
Propionibacterium granulosum		76.92	0.53	CEL 1			
Actinomyces naeslundii		23.08	0.59				
07473021	**Very good identification**						**07473021**
Actinomyces naeslundii		97.81	0.62	MLT 94			
Propionibacterium granulosum		2.20	0.19				
07473022	**Low discrimination**						**07473022**
Actinomyces naeslundii		54.55	0.59	MLT 94			
Propionibacterium granulosum		45.45	0.43				
07473023	**Very good identification**						**07473023**
Actinomyces naeslundii		99.44	0.62	MLT 94			
Propionibacterium granulosum		0.56	0.09				
07473030	**Low discrimination**						**07473030**
Propionibacterium granulosum		89.62	0.39	CEL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		10.38	0.38	MLZ 12		
07473031	**Good identification**					**07473031**
Actinomyces naeslundii		100.0	0.41	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07473033	**Good identification**						**07473033**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07473040	**Low discrimination**						**07473040**
Actinomyces naeslundii		56.72	0.47	MLT 94			
Propionibacterium granulosum		43.28	0.30				
07473041	**Good identification**						**07473041**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07473042	**Low discrimination**						**07473042**
Actinomyces naeslundii		83.98	0.47	MLT 94			
Propionibacterium granulosum		16.02	0.19				
07473043	**Good identification**						**07473043**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07473060	**Low discrimination**						**07473060**
Actinomyces naeslundii		83.98	0.47	MLT 94			
Propionibacterium granulosum		16.02	0.19				
07473061	**Good identification**						**07473061**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07473062	**Good identification**						**07473062**
Actinomyces naeslundii		95.45	0.47	MLT 94			
Propionibacterium granulosum		4.55	0.09				
07473063	**Good identification**						**07473063**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07473100	**Low discrimination**						**07473100**
					GRA	CAT	
Actinomyces naeslundii		83.90	0.67	MLT 94	99	25	
Fusobacterium mortiferum		11.47	0.41		1	5	
Propionibacterium granulosum		3.97	0.30		99	90	
07473101	**Very good identification**						**07473101**
Actinomyces naeslundii		100.0	0.70	MLT 94			
07473102	**Good identification**						**07473102**
					GRA		
Actinomyces naeslundii		94.67	0.67	MLT 94	99		
Fusobacterium mortiferum		1.44	0.24		1		
Lactobacillus acidophilus		1.37	0.11		99		
07473103	**Very good identification**						**07473103**
Actinomyces naeslundii		100.0	0.70	MLT 94			
07473110	**Low discrimination**						**07473110**
Actinomyces naeslundii		89.08	0.45	MLT 94			
Propionibacterium granulosum		10.92	0.15	MLZ 5			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07473111	**Good identification**						**07473111**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07473112	**Acceptable identification**						**07473112**
Actinomyces naeslundii		93.54	0.45	MLT 94			
Bifidobacterium adolescentis		6.46	0.10	MLZ 5			
07473113	**Good identification**						**07473113**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07473120	**Very good identification**						**07473120**
					SPO	CAT	
Actinomyces naeslundii		96.36	0.67	MLT 94	1	25	
Clostridium ramosum		2.50	0.15		92	1	
Propionibacterium granulosum		1.14	0.19		1	90	
07473121	**Very good identification**						**07473121**
Actinomyces naeslundii		100.0	0.70	MLT 94			
07473122	**Low discrimination**						**07473122**
					SPO		
Actinomyces naeslundii		88.36	0.67	MLT 94	1		
Clostridium ramosum		9.15	0.25		92		
Lactobacillus acidophilus		1.28	0.11		1		
07473123	**Very good identification**						**07473123**
Actinomyces naeslundii		100.0	0.70	MLT 94			
07473130	**Good identification**						**07473130**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07473131	**Good identification**						**07473131**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07473132	**Acceptable identification**						**07473132**
					GRA		
Actinomyces naeslundii		90.11	0.45	MLT 94	99		
Parabacteroides distasonis		9.90	0.16	MLZ 5	1		
07473133	**Good identification**						**07473133**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07473140	**Good identification**						**07473140**
Actinomyces naeslundii		98.29	0.54	MLT 94			
Actinomyces israelii		1.71	0.15				
07473141	**Very good identification**						**07473141**
Actinomyces naeslundii		100.0	0.57	MLT 94			
07473142	**Good identification**						**07473142**
Actinomyces naeslundii		97.46	0.54	MLT 94			
Actinomyces israelii		2.54	0.18				
07473143	**Very good identification**						**07473143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.57	MLT 94		
07473151	**Good identification**					**07473151**
Actinomyces naeslundii		100.0	0.35	MLT 94		
07473153	**Good identification**					**07473153**
Actinomyces naeslundii		100.0	0.35	MLT 94		
07473160	**Good identification**					**07473160**
Actinomyces naeslundii		100.0	0.54	MLT 94		
07473161	**Very good identification**					**07473161**
Actinomyces naeslundii		100.0	0.57	MLT 94		
07473162	**Good identification**					**07473162**
Actinomyces naeslundii		100.0	0.54	MLT 94		
07473163	**Very good identification**					**07473163**
Actinomyces naeslundii		100.0	0.57	MLT 94		
07473171	**Good identification**					**07473171**
Actinomyces naeslundii		100.0	0.35	MLT 94		
07473173	**Good identification**					**07473173**
Actinomyces naeslundii		100.0	0.35	MLT 94		
07473200	**Good identification**					**07473200**
Propionibacterium granulosum		97.61	0.38	CEL 1		
Actinomyces naeslundii		2.39	0.25	RHA 3		
07473301	**Good identification**					**07473301**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07473303	**Good identification**					**07473303**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07473321	**Good identification**					**07473321**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07473323	**Good identification**					**07473323**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07473501	**Good identification**					**07473501**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07473503	**Good identification**					**07473503**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07473521	**Good identification**					**07473521**
Actinomyces naeslundii		100.0	0.36	MLT 94		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07473523	**Good identification**						**07473523**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07474000	**Low discrimination**						**07474000**
					GRA	CAT	
Fusobacterium mortiferum		82.28	0.46	XYL 7	1	5	
Propionibacterium granulosum		17.72	0.32		99	90	
07474100	**Good identification**						**07474100**
Fusobacterium mortiferum		100.0	0.53	XYL 7			
07474102	**Good identification**						**07474102**
Fusobacterium mortiferum		100.0	0.36	XYL 7			
07474122	**Good identification**						**07474122**
Parabacteroides distasonis		100.0	0.37	GAL 99			
07474322	**Good identification**						**07474322**
Parabacteroides distasonis		100.0	0.45	GAL 99			
07474332	**Acceptable identification**						**07474332**
Parabacteroides distasonis		94.58	0.40	GAL 99			
Clostridium clostridioforme		5.42	0.13				
07474722	**Low discrimination**						**07474722**
Parabacteroides distasonis		85.24	0.41	GAL 99			
Clostridium clostridioforme		14.76	0.23				
07475000	**Low discrimination**						**07475000**
					GRA	CAT	
Fusobacterium mortiferum		86.75	0.49	XYL 7	1	5	
Propionibacterium granulosum		12.46	0.32		99	90	
Actinomyces israelii		0.79	0.17		99	1	
07475022	**Good identification**						**07475022**
					GRA		
Parabacteroides distasonis		96.10	0.37	LAC 99	1		
Propionibacterium granulosum		3.90	0.11		99		
07475100	**Acceptable identification**						**07475100**
					COC	GRA	
Fusobacterium mortiferum		92.40	0.56	XYL 7	1	1	
Atopobium parvulum		5.00	0.23		99	99	
Actinomyces israelii		1.45	0.27		1	99	
07475102	**Low discrimination**						**07475102**
					GRA		
Fusobacterium mortiferum		40.62	0.39	XYL 7	1		
Parabacteroides distasonis		34.65	0.37	BGL 10	1		
Bifidobacterium adolescentis		8.72	0.24		99		
07475120	**Low discrimination**						**07475120**
					COC	GRA	
Parabacteroides distasonis		79.30	0.37	BGL 99	1	1	
Fusobacterium mortiferum		8.45	0.22		1	1	
Atopobium parvulum		7.98	0.10		99	99	
07475122	**Very good identification**						**07475122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		99.66	0.71		1	
Prevotella oralis		0.22	0.13		1	
Actinomyces naeslundii		0.05	0.28		99	
07475123	**Acceptable identification**					**07475123**
					GRA	
Parabacteroides distasonis		92.52	0.37	URE	1	
Actinomyces naeslundii		7.48	0.31		99	
07475132	**Very good identification**					**07475132**
Parabacteroides distasonis		99.93	0.66			
Clostridium clostridioforme		0.07	0.07			
07475162	**Good identification**					**07475162**
Parabacteroides distasonis		100.0	0.37	SOR	1	
07475222	**Good identification**					**07475222**
Parabacteroides distasonis		100.0	0.45	LAC	99	
07475232	**Good identification**					**07475232**
Parabacteroides distasonis		95.11	0.40	LAC	99	
Clostridium clostridioforme		4.89	0.13			
07475302	**Good identification**					**07475302**
Parabacteroides distasonis		98.01	0.45	NAG	99	
Clostridium clostridioforme		1.99	0.10			
07475312	**Low discrimination**					**07475312**
Parabacteroides distasonis		85.33	0.40	NAG	99	
Clostridium clostridioforme		14.68	0.22			
07475320	**Good identification**					**07475320**
Parabacteroides distasonis		98.62	0.45	BGL	99	
Bacteroides vulgatus		1.38	-.05			
07475322	**Very good identification**					**07475322**
Parabacteroides distasonis		99.95	0.79			
Prevotella oralis		0.03	0.06			
Clostridium clostridioforme		0.02	0.10			
07475323	**Good identification**					**07475323**
Parabacteroides distasonis		100.0	0.45	URE	1	
07475330	**Good identification**					**07475330**
Parabacteroides distasonis		100.0	0.40	BGL	99	
07475332	**Very good identification**					**07475332**
Parabacteroides distasonis		99.83	0.75			
Clostridium clostridioforme		0.17	0.22			
07475333	**Good identification**					**07475333**
Parabacteroides distasonis		100.0	0.40	URE	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07475362	**Good identification**						**07475362**
Parabacteroides distasonis		100.0	0.45	SOR 1			
07475372	**Good identification**						**07475372**
Parabacteroides distasonis		100.0	0.40	SOR 1			
07475502	**Low discrimination**						**07475502**
					GRA	SPO	
Bifidobacterium adolescentis		57.37	0.38	MLT 99	99	1	
Parabacteroides distasonis		20.87	0.33		1	1	
Clostridium butyricum		9.30	0.10		99	99	
07475512	**Low discrimination**						**07475512**
					GRA		
Bifidobacterium adolescentis		67.12	0.40	MLT 99	99		
Clostridium clostridioforme		19.35	0.27		50		
Parabacteroides distasonis		10.76	0.29		1		
07475522	**Very good identification**						**07475522**
					GRA	CAT	
Parabacteroides distasonis		99.28	0.67		1	76	
Bacteroides thetaiotaomicron		0.29	0.08		1	89	
Bifidobacterium longum subsp. longum		0.21	0.17		99	1	
07475532	**Very good identification**						**07475532**
					GRA		
Parabacteroides distasonis		97.37	0.63		1		
Clostridium clostridioforme		1.77	0.27		50		
Bifidobacterium longum subsp. longum		0.66	0.21		99		
07475622	**Low discrimination**						**07475622**
Parabacteroides distasonis		84.62	0.41	LAC 99			
Clostridium clostridioforme		13.13	0.22				
Bacteroides thetaiotaomicron		2.25	-.02				
07475702	**Low discrimination**						**07475702**
Parabacteroides distasonis		65.81	0.41	NAG 99			
Clostridium clostridioforme		34.19	0.31				
07475712	**Low discrimination**						**07475712**
Clostridium clostridioforme		81.47	0.42	MLT 99			
Parabacteroides distasonis		18.53	0.37	SAL 88			
07475720	**Low discrimination**						**07475720**
Parabacteroides distasonis		76.78	0.41	BGL 99			
Bacteroides vulgatus		23.22	0.14				
07475722	**Very good identification**						**07475722**
Parabacteroides distasonis		98.10	0.76				
Bacteroides thetaiotaomicron		1.29	0.27				
Clostridium clostridioforme		0.52	0.31				
07475723	**Good identification**						**07475723**
Parabacteroides distasonis		100.0	0.41	URE 1			
07475730	**Good identification**						**07475730**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		95.75	0.37	BGL 99		
Clostridium clostridioforme		4.25	0.08			
07475732	**Good identification**					**07475732**
Parabacteroides distasonis		94.78	0.71			
Clostridium clostridioforme		4.21	0.42			
Bacteroides thetaiotaomicron		0.90	0.20			
07475733	**Good identification**					**07475733**
Parabacteroides distasonis		95.75	0.37	URE 1		
Clostridium clostridioforme		4.25	0.08			
07475762	**Good identification**					**07475762**
Parabacteroides distasonis		100.0	0.41	SOR 1		
07475772	**Good identification**					**07475772**
Parabacteroides distasonis		95.75	0.37	SOR 1		
Clostridium clostridioforme		4.25	0.08			
07477100	**Low discrimination**					**07477100**
					GRA	
Actinomyces israelii		81.63	0.40	MLT 99	99	
Fusobacterium mortiferum		9.27	0.22	SAL 92	1	
Actinomyces naeslundii		9.10	0.33		99	
07477101	**Good identification**					**07477101**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07477102	**Low discrimination**					**07477102**
Actinomyces israelii		81.93	0.43	MLT 99		
Bifidobacterium adolescentis		11.99	0.23	SAL 92		
Actinomyces naeslundii		6.09	0.33			
07477103	**Good identification**					**07477103**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07477121	**Good identification**					**07477121**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07477122	**Low discrimination**					**07477122**
					GRA	
Parabacteroides distasonis		83.72	0.37	MAN 1	1	
Actinomyces naeslundii		8.75	0.33		99	
Actinomyces israelii		7.52	0.23		99	
07477123	**Good identification**					**07477123**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07477142	**Good identification**					**07477142**
Actinomyces israelii		95.56	0.36	MLT 99		
Bifidobacterium adolescentis		4.44	0.07	SAL 92		
07477322	**Good identification**					**07477322**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		100.0	0.45	MAN 1				
07477332	**Good identification**							**07477332**
Parabacteroides distasonis		100.0	0.40	MAN 1				
07477500	**Acceptable identification**							**07477500**
					SPO			
Actinomyces israelii		90.78	0.39	MLT 99	1			
Clostridium butyricum		9.22	0.01	SAL 92	99			
07477502	**Low discrimination**							**07477502**
					SPO			
Bifidobacterium adolescentis		53.84	0.37	MLT 99	1			
Actinomyces israelii		44.98	0.42		1			
Clostridium butyricum		1.19	-.07		99			
07477512	**Low discrimination**							**07477512**
Bifidobacterium adolescentis		88.57	0.38	MLT 99				
Actinomyces israelii		9.05	0.28					
Bifidobacterium dentium		2.38	-.07					
07477722	**Good identification**							**07477722**
Parabacteroides distasonis		96.88	0.41	MAN 1				
Bacteroides ovatus		3.13	0.00					
07477732	**Acceptable identification**							**07477732**
Parabacteroides distasonis		92.22	0.37	MAN 1				
Clostridium clostridioforme		4.10	0.08					
Bacteroides ovatus		3.68	-.03					
07500000	**Low discrimination**							**07500000**
					COC	GRA	CAT	
Gemella morbillorum		78.50	0.64	RAF 1	99	99	1	
Prevotella intermedia		11.69	0.49		1	1	1	
Propionibacterium granulosum		5.61	0.52		1	99	90	
07500002	**Low discrimination**							**07500002**
					COC	CAT		
Actinomyces viscosus		61.43	0.40	NIT 99	1	76		
Propionibacterium granulosum		22.24	0.41	FRU 95	1	90		
Gemella morbillorum		12.57	0.30		99	1		
07500020	**Low discrimination**							**07500020**
					COC	GRA	CAT	
Gemella morbillorum		81.31	0.48	RAF 1	99	99	1	
Propionibacterium granulosum		13.07	0.41	NAG 10	1	99	90	
Prevotella melaninogenica		2.70	0.15		1	1	1	
07500040	**Acceptable identification**							**07500040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		93.81	0.47	RAF 1	1	99	99	
Gemella morbillorum		4.34	0.30		99	99	1	
Prevotella intermedia		0.65	0.15		1	1	1	
07500050	**Good identification**							**07500050**
Clostridium botulinum biovar B		100.0	0.42	RAF 1				
07500060	**Good identification**							**07500060**
					COC	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		98.97	0.42	RAF 1	1	99		
Gemella morbillorum		1.03	0.14		99	1		
07500070	**Good identification**							**07500070**
Clostridium botulinum biovar B		100.0	0.37	RAF 1				
07500102	**Good identification**							**07500102**
					GRA			
Actinomyces viscosus		96.45	0.44	NIT 99	99			
Fusobacterium mortiferum		1.06	0.23	FRU 95	1			
Lactobacillus fermentum		0.93	0.24		99			
07500103	**Good identification**							**07500103**
					SPO			
Actinomyces viscosus		95.81	0.36	NIT 99	1			
Actinomyces naeslundii		2.60	0.33	FRU 95	1			
Clostridium chauvoei		1.60	0.07		99			
07500120	**Low discrimination**							**07500120**
					GRA			
Prevotella melaninogenica		81.22	0.50	FRU 90	1			
Prevotella bivia		18.05	0.32	GAL 99	1			
Clostridium perfringens		0.30	0.15		99			
07500122	**Acceptable identification**							**07500122**
					GRA			
Prevotella melaninogenica		94.91	0.37	FRU 90	1			
Actinomyces viscosus		1.21	0.10	GAL 99	99			
Prevotella bivia		1.12	-.03		1			
07500400	**Good identification**							**07500400**
					COC	GRA	CAT	
Gemella morbillorum		97.25	0.48	RAF 1	99	99	1	
Prevotella intermedia		1.32	0.15	ARA 10	1	1	1	
Propionibacterium granulosum		0.63	0.18		1	99	90	
07500520	**Low discrimination**							**07500520**
					GRA			
Bifidobacterium longum subsp. longum		74.61	0.37	FRU 90	99			
Prevotella melaninogenica		19.56	0.15	GAL 99	1			
Prevotella bivia		4.35	-.03		1			
07500522	**Acceptable identification**							**07500522**
					GRA			
Bifidobacterium longum subsp. longum		93.47	0.37	FRU 90	99			
Prevotella melaninogenica		4.67	0.03	GAL 99	1			
Lactobacillus fermentum		1.87	0.15		99			
07500530	**Good identification**							**07500530**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
07500532	**Good identification**							**07500532**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
07501000	**Low discrimination**							**07501000**
					COC	GRA	CAT	
Gemella morbillorum		65.25	0.60	RAF 1	99	99	1	
Prevotella intermedia		18.05	0.49		1	1	1	
Propionibacterium granulosum		8.66	0.52		1	99	90	
07501001	**Low discrimination**							**07501001**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Actinomyces naeslundii		37.93	0.49	FRU 95	1	1		
Gemella morbillorum		30.85	0.26		99	1		
Clostridium chauvoei		15.67	0.19		1	99		
07501002	**Low discrimination**							**07501002**
					CAT			
Lactobacillus fermentum		45.47	0.54	FRU 99	1			
Actinomyces viscosus		26.91	0.36		76			
Propionibacterium granulosum		15.89	0.41		90			
07501010	**Low discrimination**							**07501010**
					COC	GRA	CAT	
Propionibacterium granulosum		55.88	0.37	FRU 99	1	99	90	
Gemella morbillorum		31.19	0.26	MLZ 12	99	99	1	
Prevotella intermedia		8.63	0.15		1	1	1	
07501020	**Low discrimination**							**07501020**
					COC	GRA		
Prevotella melaninogenica		75.40	0.50	FRU 90	1	1		
Gemella morbillorum		12.36	0.43	LAC 99	99	99		
Clostridium perfringens		3.76	0.34		1	99		
07501021	**Low discrimination**							**07501021**
					GRA	SPO		
Actinomyces naeslundii		56.20	0.49	FRU 95	99	1		
Prevotella melaninogenica		30.98	0.15		1	1		
Clostridium chauvoei		7.74	0.11		99	99		
07501022	**Low discrimination**							**07501022**
					GRA			
Prevotella melaninogenica		64.01	0.37	FRU 90	1			
Lactobacillus fermentum		15.69	0.46	LAC 99	99			
Clostridium perfringens		7.88	0.29		99			
07501023	**Acceptable identification**							**07501023**
					GRA			
Actinomyces naeslundii		90.50	0.49	FRU 95	99			
Prevotella melaninogenica		9.50	0.03		1			
07501041	**Good identification**							**07501041**
Actinomyces naeslundii		100.0	0.36	FRU 95				
07501043	**Good identification**							**07501043**
Actinomyces naeslundii		100.0	0.36	FRU 95				
07501061	**Good identification**							**07501061**
Actinomyces naeslundii		100.0	0.36	FRU 95				
07501063	**Good identification**							**07501063**
Actinomyces naeslundii		100.0	0.36	FRU 95				
07501100	**Low discrimination**							**07501100**
					COC	GRA		
Prevotella melaninogenica		69.52	0.50	FRU 90	1	1		
Lactobacillus fermentum		15.89	0.58	NAG 99	1	99		
Blautia hansenii		6.09	0.32		99	99		
07501101	**Acceptable identification**							**07501101**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		91.97	0.54	RAF 1	99	99	
Actinomyces naeslundii		6.40	0.57		99	1	
Prevotella melaninogenica		1.24	0.15		1	1	
07501102	**Low discrimination**						**07501102**
					GRA		
Lactobacillus fermentum		35.43	0.58	FRU 99	99		
Prevotella melaninogenica		29.53	0.37		1		
Actinomyces viscosus		22.85	0.40		99		
07501103	**Low discrimination**						**07501103**
					SPO		
Clostridium chauvoei		55.63	0.41	RAF 1	99		
Actinomyces naeslundii		21.93	0.57		1		
Actinomyces viscosus		20.66	0.32		1		
07501120	**Very good identification**						**07501120**
					GRA		
Prevotella melaninogenica		99.42	0.84	FRU 90	1		
Clostridium perfringens		0.37	0.49		99		
Lactobacillus fermentum		0.08	0.50		99		
07501121	**Low discrimination**						**07501121**
					GRA	SPO	
Prevotella melaninogenica		76.54	0.50	FRU 90	1	1	
Clostridium chauvoei		19.12	0.45	URE 1	99	99	
Actinomyces naeslundii		3.99	0.57		99	1	
07501122	**Very good identification**						**07501122**
					GRA		
Prevotella melaninogenica		97.30	0.71	FRU 90	1		
Clostridium perfringens		0.89	0.43		99		
Prevotella oralis		0.81	0.39		1		
07501123	**Low discrimination**						**07501123**
					GRA	SPO	
Prevotella melaninogenica		65.50	0.37	FRU 90	1	1	
Actinomyces naeslundii		17.93	0.57	URE 1	99	1	
Clostridium chauvoei		15.16	0.33		99	99	
07501130	**Acceptable identification**						**07501130**
					GRA		
Prevotella melaninogenica		93.29	0.50	FRU 90	1		
Bifidobacterium longum subsp. longum		6.12	0.41	MLZ 1	99		
Clostridium perfringens		0.34	0.15		99		
07501132	**Low discrimination**						**07501132**
					GRA	CAT	
Prevotella melaninogenica		71.12	0.37	FRU 90	1	1	
Bifidobacterium longum subsp. longum		24.50	0.41	MLZ 1	99	1	
Capnocytophaga ochracea		1.51	0.12		1	90	
07501141	**Low discrimination**						**07501141**
					SPO		
Actinomyces naeslundii		54.87	0.44	FRU 95	1		
Clostridium chauvoei		45.14	0.19		99		
07501142	**Low discrimination**						**07501142**
					GRA		
Actinomyces naeslundii		45.85	0.41	FRU 95	99		
Lactobacillus fermentum		17.25	0.24		99		
Prevotella melaninogenica		14.38	0.03		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07501143	**Low discrimination**					**07501143**
					SPO	
Actinomyces naeslundii		87.32	0.44	FRU 95	1	
Clostridium chauvoei		12.68	0.07		99	
07501160	**Acceptable identification**					**07501160**
					GRA	
Prevotella melaninogenica		92.90	0.50	FRU 90	1	
Clostridium perfringens		6.46	0.37	SOR 1	99	
Actinomyces naeslundii		0.57	0.41		99	
07501162	**Low discrimination**					**07501162**
					GRA	
Prevotella melaninogenica		80.79	0.37	FRU 90	1	
Clostridium perfringens		13.89	0.31	SOR 1	99	
Actinomyces naeslundii		2.60	0.41		99	
07501163	**Low discrimination**					**07501163**
					GRA	
Actinomyces naeslundii		82.71	0.44	FRU 95	99	
Prevotella melaninogenica		17.29	0.03		1	
07501320	**Good identification**					**07501320**
					GRA	
Prevotella melaninogenica		98.43	0.50	FRU 90	1	
Bacteroides vulgatus		0.78	0.11	RHA 1	1	
Clostridium perfringens		0.36	0.15		99	
07501322	**Low discrimination**					**07501322**
Prevotella melaninogenica		70.50	0.37	FRU 90		
Prevotella oralis		24.95	0.33	RHA 1		
Parabacteroides distasonis		3.63	0.31			
07501400	**Low discrimination**					**07501400**
					COC	
Gemella morbillorum		51.28	0.43	RAF 1	99	
Lactobacillus fermentum		43.80	0.54	ARA 10	1	
Bifidobacterium longum subsp. longum		3.01	0.27		1	
07501402	**Acceptable identification**					**07501402**
					COC	
Lactobacillus fermentum		92.03	0.54	FRU 99	1	
Bifidobacterium longum subsp. longum		6.33	0.27		1	
Gemella morbillorum		1.09	0.09		99	
07501420	**Low discrimination**					**07501420**
					COC	
Lactobacillus fermentum		41.12	0.46	FRU 99	1	
Bifidobacterium longum subsp. longum		33.94	0.37		1	
Gemella morbillorum		16.05	0.27		99	
07501422	**Low discrimination**					**07501422**
					GRA	
Lactobacillus fermentum		53.58	0.46	FRU 99	99	
Bifidobacterium longum subsp. longum		44.22	0.37		99	
Prevotella melaninogenica		2.21	0.03		1	
07501430	**Good identification**					**07501430**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90		
07501432	**Good identification**					**07501432**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90			
07501500	**Low discrimination**						**07501500**
					COC		
Bifidobacterium longum subsp. longum		78.86	0.61	FRU 90	1		
Lactobacillus fermentum		18.90	0.58		1		
Blautia hansenii		1.28	0.19		99		
07501502	**Low discrimination**						**07501502**
Bifidobacterium longum subsp. longum		80.03	0.61	FRU 90			
Lactobacillus fermentum		19.18	0.58				
Bifidobacterium adolescentis		0.30	0.21				
07501510	**Very good identification**						**07501510**
Bifidobacterium longum subsp. longum		99.86	0.65	FRU 90			
Lactobacillus fermentum		0.14	0.24				
07501512	**Very good identification**						**07501512**
Bifidobacterium longum subsp. longum		99.59	0.65	FRU 90			
Bifidobacterium adolescentis		0.27	0.23				
Lactobacillus fermentum		0.14	0.24				
07501520	**Low discrimination**						**07501520**
					GRA		
Bifidobacterium longum subsp. longum		77.72	0.72	FRU 90	99		
Prevotella melaninogenica		20.37	0.50		1		
Lactobacillus fermentum		1.55	0.50		99		
07501521	**Low discrimination**						**07501521**
					GRA	SPO	
Bifidobacterium longum subsp. longum		73.46	0.37	FRU 90	99	1	
Prevotella melaninogenica		19.26	0.15	URE 1	1	1	
Clostridium chauvoei		4.81	0.11		99	99	
07501522	**Good identification**						**07501522**
					GRA		
Bifidobacterium longum subsp. longum		93.20	0.72	FRU 90	99		
Prevotella melaninogenica		4.65	0.37		1		
Lactobacillus fermentum		1.86	0.50		99		
07501523	**Acceptable identification**						**07501523**
					GRA		
Bifidobacterium longum subsp. longum		92.29	0.37	FRU 90	99		
Prevotella melaninogenica		4.61	0.03	URE 1	1		
Lactobacillus fermentum		1.84	0.15		99		
07501530	**Very good identification**						**07501530**
					GRA		
Bifidobacterium longum subsp. longum		99.83	0.76	FRU 90	99		
Prevotella melaninogenica		0.16	0.15		1		
Lactobacillus fermentum		0.01	0.15		99		
07501531	**Good identification**						**07501531**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90			
07501532	**Very good identification**						**07501532**
					GRA		
Bifidobacterium longum subsp. longum		99.93	0.76	FRU 90	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella melaninogenica		0.03	0.03		1			
Parabacteroides distasonis		0.02	0.15		1			
07501533	**Good identification**							**07501533**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
07501560	**Low discrimination**							**07501560**
					GRA			
Bifidobacterium longum subsp. longum		76.91	0.37	FRU 90	99			
Prevotella melaninogenica		20.16	0.15	SOR 1	1			
Lactobacillus fermentum		1.54	0.15		99			
07501562	**Acceptable identification**							**07501562**
					GRA			
Bifidobacterium longum subsp. longum		91.88	0.37	FRU 90	99			
Prevotella melaninogenica		4.59	0.03	SOR 1	1			
Lactobacillus fermentum		1.83	0.15		99			
07501570	**Good identification**							**07501570**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
07501572	**Good identification**							**07501572**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
07501722	**Low discrimination**							**07501722**
					GRA			
Bifidobacterium longum subsp. longum		78.67	0.37	FRU 90	99			
Parabacteroides distasonis		12.27	0.28	RHA 1	1			
Prevotella melaninogenica		3.93	0.03		1			
07501730	**Good identification**							**07501730**
					GRA			
Bifidobacterium longum subsp. longum		98.41	0.41	FRU 90	99			
Bacteroides vulgatus		1.60	-.04	RHA 1	1			
07501732	**Good identification**							**07501732**
					GRA			
Bifidobacterium longum subsp. longum		95.30	0.41	FRU 90	99			
Parabacteroides distasonis		4.70	0.23	RHA 1	1			
07502000	**Low discrimination**							**07502000**
					COC	GRA	CAT	
Propionibacterium granulosum		80.69	0.50	FRU 99	1	99	90	
Gemella morbillorum		15.12	0.30		99	99	1	
Prevotella intermedia		2.25	0.15		1	1	1	
07502002	**Acceptable identification**							**07502002**
Propionibacterium granulosum		92.80	0.39	FRU 99				
Actinomyces viscosus		3.43	0.06					
Actinomyces naeslundii		2.48	0.27					
07502020	**Acceptable identification**							**07502020**
					COC	CAT		
Propionibacterium granulosum		90.09	0.39	FRU 99	1	90		
Gemella morbillorum		7.50	0.14		99	1		
Actinomyces naeslundii		2.41	0.27		1	25		
07502100	**Low discrimination**							**07502100**
					CAT			
Propionibacterium propionicum		88.72	0.43	NIT 86	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		7.37	0.35	FRU 99	25		
Propionibacterium granulosum		3.91	0.16		90		
07502101	**Low discrimination**						**07502101**
Actinomyces naeslundii		52.85	0.38	FRU 95			
Propionibacterium propionicum		47.15	0.27	GAL 96			
07502102	**Acceptable identification**						**07502102**
Propionibacterium propionicum		90.65	0.46	NIT 86			
Actinomyces naeslundii		5.02	0.35	FRU 99			
Actinomyces viscosus		4.33	0.10				
07502121	**Good identification**						**07502121**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07502122	**Low discrimination**						**07502122**
					GRA		
Actinomyces naeslundii		53.57	0.35	FRU 95	99		
Prevotella melaninogenica		36.65	0.03	GAL 96	1		
Propionibacterium propionicum		9.77	0.12		99		
07502123	**Good identification**						**07502123**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07502140	**Good identification**						**07502140**
Propionibacterium propionicum		100.0	0.43	NIT 86			
07502142	**Good identification**						**07502142**
Propionibacterium propionicum		100.0	0.46	NIT 86			
07503000	**Low discrimination**						**07503000**
					COC	CAT	
Propionibacterium granulosum		76.99	0.50	FRU 99	1	90	
Actinomyces naeslundii		12.36	0.51		1	25	
Gemella morbillorum		7.77	0.26		99	1	
07503001	**Good identification**						**07503001**
Actinomyces naeslundii		95.97	0.54	FRU 95			
Propionibacterium granulosum		4.03	0.16				
07503002	**Low discrimination**						**07503002**
					CAT		
Propionibacterium granulosum		58.25	0.39	FRU 99	90		
Actinomyces naeslundii		37.39	0.51		25		
Lactobacillus fermentum		2.23	0.20		1		
07503003	**Good identification**						**07503003**
Actinomyces naeslundii		100.0	0.54	FRU 95			
07503020	**Low discrimination**						**07503020**
					GRA	CAT	
Propionibacterium granulosum		51.01	0.39	FRU 99	99	90	
Actinomyces naeslundii		32.75	0.51		99	25	
Prevotella melaninogenica		13.95	0.15		1	1	
07503021	**Good identification**						**07503021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.54	FRU 95			
07503022	**Low discrimination**						**07503022**
					GRA	CAT	
Actinomyces naeslundii		68.00	0.51	FRU 95	99	25	
Propionibacterium granulosum		26.48	0.29		99	90	
Prevotella melaninogenica		5.52	0.03		1	1	
07503023	**Good identification**						**07503023**
Actinomyces naeslundii		100.0	0.54	FRU 95			
07503040	**Low discrimination**						**07503040**
Actinomyces naeslundii		73.71	0.38	FRU 95			
Propionibacterium granulosum		26.29	0.16				
07503041	**Good identification**						**07503041**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07503042	**Low discrimination**						**07503042**
Actinomyces naeslundii		89.08	0.38	FRU 95			
Propionibacterium propionicum		10.92	0.12				
07503043	**Good identification**						**07503043**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07503060	**Good identification**						**07503060**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07503061	**Good identification**						**07503061**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07503062	**Good identification**						**07503062**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07503063	**Good identification**						**07503063**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07503100	**Low discrimination**						**07503100**
					GRA		
Actinomyces naeslundii		58.11	0.59	FRU 95	99		
Propionibacterium propionicum		29.16	0.43		99		
Prevotella melaninogenica		8.70	0.15		1		
07503101	**Low discrimination**						**07503101**
					SPO		
Actinomyces naeslundii		89.93	0.62	FRU 95	1		
Clostridium chauvoei		6.73	0.19		99		
Propionibacterium propionicum		3.34	0.27		1		
07503102	**Low discrimination**						**07503102**
Actinomyces naeslundii		54.42	0.59	FRU 95			
Propionibacterium propionicum		40.97	0.46				
Lactobacillus fermentum		1.86	0.24				
07503103	**Good identification**						**07503103**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		93.55	0.62	FRU 95	1		
Propionibacterium propionicum		5.22	0.30		1		
Clostridium chauvoei		1.24	0.07		99		
07503110	**Good identification**						**07503110**
Actinomyces naeslundii		100.0	0.37	FRU 95			
07503111	**Good identification**						**07503111**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07503112	**Low discrimination**						**07503112**
Actinomyces naeslundii		77.13	0.37	FRU 95			
Bifidobacterium adolescentis		11.73	0.07	MLZ 5			
Propionibacterium propionicum		11.14	0.12				
07503113	**Good identification**						**07503113**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07503120	**Acceptable identification**						**07503120**
					GRA		
Prevotella melaninogenica		93.29	0.50	FRU 90	1		
Actinomyces naeslundii		6.30	0.59	MAN 1	99		
Clostridium perfringens		0.34	0.15		99		
07503121	**Low discrimination**						**07503121**
					GRA	SPO	
Actinomyces naeslundii		88.91	0.62	FRU 95	99	1	
Prevotella melaninogenica		8.87	0.15		1	1	
Clostridium chauvoei		2.22	0.11		99	99	
07503122	**Low discrimination**						**07503122**
					GRA		
Prevotella melaninogenica		72.54	0.37	FRU 90	1		
Actinomyces naeslundii		25.70	0.59	MAN 1	99		
Clostridium perfringens		0.66	0.09		99		
07503123	**Very good identification**						**07503123**
					GRA		
Actinomyces naeslundii		98.14	0.62	FRU 95	99		
Prevotella melaninogenica		1.87	0.03		1		
07503131	**Good identification**						**07503131**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07503132	**Low discrimination**						**07503132**
					GRA		
Actinomyces naeslundii		57.86	0.37	FRU 95	99		
Prevotella melaninogenica		31.34	0.03	MLZ 5	1		
Bifidobacterium longum subsp. longum		10.80	0.07		99		
07503133	**Good identification**						**07503133**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07503140	**Low discrimination**						**07503140**
Propionibacterium propionicum		73.98	0.43	NIT 86			
Actinomyces naeslundii		26.02	0.46	FRU 99			
07503141	**Low discrimination**						**07503141**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		82.60	0.49	FRU 95				
Propionibacterium propionicum		17.40	0.27					
07503142	**Low discrimination**							**07503142**
Propionibacterium propionicum		81.01	0.46	NIT 86				
Actinomyces naeslundii		18.99	0.46	FRU 99				
07503143	**Low discrimination**							**07503143**
Actinomyces naeslundii		75.99	0.49	FRU 95				
Propionibacterium propionicum		24.01	0.30					
07503160	**Low discrimination**							**07503160**
					GRA			
Actinomyces naeslundii		52.44	0.46	FRU 95	99			
Prevotella melaninogenica		44.47	0.15		1			
Clostridium perfringens		3.09	0.02		99			
07503161	**Good identification**							**07503161**
Actinomyces naeslundii		100.0	0.49	FRU 95				
07503162	**Low discrimination**							**07503162**
					GRA			
Actinomyces naeslundii		83.01	0.46	FRU 95	99			
Prevotella melaninogenica		13.41	0.03		1			
Propionibacterium propionicum		3.58	0.12		99			
07503163	**Good identification**							**07503163**
Actinomyces naeslundii		100.0	0.49	FRU 95				
07503520	**Low discrimination**							**07503520**
					GRA			
Bifidobacterium longum subsp. longum		76.94	0.37	FRU 90	99			
Prevotella melaninogenica		20.17	0.15	MAN 1	1			
Lactobacillus fermentum		1.54	0.15		99			
07503522	**Acceptable identification**							**07503522**
					GRA			
Bifidobacterium longum subsp. longum		91.95	0.37	FRU 90	99			
Prevotella melaninogenica		4.59	0.03	MAN 1	1			
Lactobacillus fermentum		1.84	0.15		99			
07503530	**Good identification**							**07503530**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
07503532	**Good identification**							**07503532**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
07504000	**Good identification**							**07504000**
					COC	GRA	CAT	
Gemella morbillorum		97.19	0.48	RAF 1	99	99	1	
Prevotella intermedia		1.32	0.15	XYL 10	1	1	1	
Propionibacterium granulosum		0.63	0.18		1	99	90	
07504500	**Good identification**							**07504500**
Bifidobacterium longum subsp. longum		95.52	0.35	FRU 90				
Lactobacillus fermentum		4.48	0.20	GAL 99				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07504502	**Good identification**						**07504502**
Bifidobacterium longum subsp. longum		95.52	0.35	FRU 90			
Lactobacillus fermentum		4.48	0.20	GAL 99			
07504510	**Good identification**						**07504510**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
07504512	**Good identification**						**07504512**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
07504520	**Good identification**						**07504520**
					GRA		
Bifidobacterium longum subsp. longum		95.23	0.46	FRU 90	99		
Bacteroides vulgatus		4.77	0.09	GAL 99	1		
07504522	**Good identification**						**07504522**
Bifidobacterium longum subsp. longum		100.0	0.46	FRU 90			
07504530	**Good identification**						**07504530**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
07504532	**Good identification**						**07504532**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
07505000	**Low discrimination**						**07505000**
					COC	GRA	
Gemella morbillorum		64.68	0.43	RAF 1	99	99	
Lactobacillus fermentum		32.45	0.50	XYL 10	1	99	
Prevotella intermedia		1.63	0.15		1	1	
07505002	**Good identification**						**07505002**
					COC		
Lactobacillus fermentum		97.06	0.50	FRU 99	1		
Gemella morbillorum		1.95	0.09		99		
Actinomyces viscosus		0.99	0.02		1		
07505020	**Low discrimination**						**07505020**
					COC	GRA	
Lactobacillus fermentum		48.19	0.42	FRU 99	1	99	
Gemella morbillorum		32.01	0.27		99	99	
Prevotella melaninogenica		17.75	0.15		1	1	
07505022	**Low discrimination**						**07505022**
					CAT	GRA	
Lactobacillus fermentum		81.45	0.42	FRU 99	1	99	
Prevotella melaninogenica		5.72	0.03		1	1	
Bacteroides fragilis		5.68	-.03		99	1	
07505100	**Low discrimination**						**07505100**
					COC		
Lactobacillus fermentum		68.98	0.54	FRU 99	1		
Bifidobacterium longum subsp. longum		14.85	0.35		1		
Blautia hansenii		7.94	0.19		99		
07505102	**Low discrimination**						**07505102**
Lactobacillus fermentum		76.34	0.54	FRU 99			
Bifidobacterium longum subsp. longum		16.44	0.35				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		2.06	0.21				
07505110	**Good identification**						**07505110**
Bifidobacterium longum subsp. longum		97.32	0.39	FRU 90			
Lactobacillus fermentum		2.68	0.20	ARA 99			
07505112	**Low discrimination**						**07505112**
Bifidobacterium longum subsp. longum		89.50	0.39	FRU 90			
Bifidobacterium adolescentis		8.04	0.23	ARA 99			
Lactobacillus fermentum		2.47	0.20				
07505120	**Low discrimination**						**07505120**
					GRA		
Prevotella melaninogenica		81.91	0.50	FRU 90	1		
Bifidobacterium longum subsp. longum		9.47	0.46	XYL 1	99		
Lactobacillus fermentum		3.67	0.46		99		
07505122	**Low discrimination**						**07505122**
					CAT	GRA	
Prevotella melaninogenica		31.72	0.37	FRU 90	1	1	
Bacteroides fragilis		31.52	0.32	XYL 1	99	1	
Bifidobacterium longum subsp. longum		19.25	0.46		1	99	
07505130	**Good identification**						**07505130**
					GRA		
Bifidobacterium longum subsp. longum		95.12	0.50	FRU 90	99		
Prevotella melaninogenica		4.88	0.15	ARA 99	1		
07505132	**Acceptable identification**						**07505132**
					GRA		
Bifidobacterium longum subsp. longum		90.49	0.50	FRU 90	99		
Parabacteroides distasonis		7.28	0.35	ARA 99	1		
Prevotella melaninogenica		0.88	0.03		1		
07505320	**Good identification**						**07505320**
					GRA		
Bacteroides vulgatus		96.96	0.40	FRU 99	1		
Prevotella melaninogenica		2.52	0.15	ARA 93	1		
Bifidobacterium longum subsp. longum		0.29	0.11		99		
07505322	**Low discrimination**						**07505322**
Parabacteroides distasonis		83.64	0.48	FRU 99			
Prevotella oralis		7.90	0.17				
Bacteroides vulgatus		3.72	0.06				
07505332	**Good identification**						**07505332**
					GRA		
Parabacteroides distasonis		95.98	0.43	FRU 99	1		
Bifidobacterium longum subsp. longum		4.02	0.15		99		
07505400	**Low discrimination**						**07505400**
					COC		
Lactobacillus fermentum		63.58	0.50	FRU 99	1		
Bifidobacterium longum subsp. longum		22.34	0.35		1		
Gemella morbillorum		14.08	0.27		99		
07505402	**Low discrimination**						**07505402**
Lactobacillus fermentum		74.00	0.50	FRU 99			
Bifidobacterium longum subsp. longum		26.00	0.35				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07505410	**Good identification**					**07505410**
Bifidobacterium longum subsp. longum		98.34	0.39	FRU 90		
Lactobacillus fermentum		1.66	0.16	LAC 99		
07505412	**Good identification**					**07505412**
Bifidobacterium longum subsp. longum		98.34	0.39	FRU 90		
Lactobacillus fermentum		1.66	0.16	LAC 99		
07505420	**Low discrimination**					**07505420**
					GRA	
Bifidobacterium longum subsp. longum		76.64	0.46	FRU 90	99	
Lactobacillus fermentum		18.18	0.42	LAC 99	99	
Bacteroides vulgatus		3.84	0.09		1	
07505422	**Low discrimination**					**07505422**
Bifidobacterium longum subsp. longum		80.83	0.46	FRU 90		
Lactobacillus fermentum		19.17	0.42	LAC 99		
07505430	**Good identification**					**07505430**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90		
07505432	**Good identification**					**07505432**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90		
07505500	**Very good identification**					**07505500**
					GRA	
Bifidobacterium longum subsp. longum		95.25	0.69	FRU 90	99	
Lactobacillus fermentum		4.47	0.54		99	
Bacteroides vulgatus		0.19	0.09		1	
07505501	**Good identification**					**07505501**
Bifidobacterium longum subsp. longum		95.52	0.35	FRU 90		
Lactobacillus fermentum		4.48	0.20	URE 1		
07505502	**Good identification**					**07505502**
Bifidobacterium longum subsp. longum		94.75	0.69	FRU 90		
Lactobacillus fermentum		4.45	0.54			
Bifidobacterium adolescentis		0.80	0.36			
07505503	**Good identification**					**07505503**
Bifidobacterium longum subsp. longum		95.52	0.35	FRU 90		
Lactobacillus fermentum		4.48	0.20	URE 1		
07505510	**Very good identification**					**07505510**
Bifidobacterium longum subsp. longum		99.97	0.73	FRU 90		
Lactobacillus fermentum		0.03	0.20			
07505511	**Good identification**					**07505511**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
07505512	**Very good identification**					**07505512**
Bifidobacterium longum subsp. longum		99.37	0.73	FRU 90		
Bifidobacterium adolescentis		0.60	0.37			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		0.03	0.20			
07505513	**Good identification**					**07505513**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
07505520	**Good identification**					**07505520**
					GRA	
Bifidobacterium longum subsp. longum		94.80	0.80	FRU 90	99	
Bacteroides vulgatus		4.75	0.43		1	
Lactobacillus fermentum		0.37	0.46		99	
07505521	**Good identification**					**07505521**
					GRA	
Bifidobacterium longum subsp. longum		95.23	0.46	FRU 90	99	
Bacteroides vulgatus		4.77	0.09	URE 1	1	
07505522	**Very good identification**					**07505522**
					GRA	
Bifidobacterium longum subsp. longum		99.36	0.80	FRU 90	99	
Lactobacillus fermentum		0.39	0.46		99	
Parabacteroides distasonis		0.16	0.36		1	
07505523	**Good identification**					**07505523**
Bifidobacterium longum subsp. longum		100.0	0.46	FRU 90		
07505530	**Very good identification**					**07505530**
					GRA	
Bifidobacterium longum subsp. longum		99.97	0.84	FRU 90	99	
Bacteroides vulgatus		0.03	0.09		1	
07505531	**Good identification**					**07505531**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90		
07505532	**Very good identification**					**07505532**
					GRA	
Bifidobacterium longum subsp. longum		99.95	0.84	FRU 90	99	
Parabacteroides distasonis		0.05	0.31		1	
07505533	**Good identification**					**07505533**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90		
07505540	**Good identification**					**07505540**
Bifidobacterium longum subsp. longum		95.52	0.35	FRU 90		
Lactobacillus fermentum		4.48	0.20	SOR 1		
07505542	**Low discrimination**					**07505542**
Bifidobacterium longum subsp. longum		86.93	0.35	FRU 90		
Bifidobacterium adolescentis		9.00	0.20	SOR 1		
Lactobacillus fermentum		4.08	0.20			
07505550	**Good identification**					**07505550**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
07505552	**Acceptable identification**					**07505552**
Bifidobacterium longum subsp. longum		93.09	0.39	FRU 90		
Bifidobacterium adolescentis		6.91	0.22	SOR 1		
07505560	**Good identification**					**07505560**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium longum subsp. longum		95.23	0.46	FRU 90	99	
Bacteroides vulgatus		4.77	0.09	SOR 1	1	
07505562	**Good identification**					**07505562**
Bifidobacterium longum subsp. longum		100.0	0.46	FRU 90		
07505570	**Good identification**					**07505570**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90		
07505572	**Good identification**					**07505572**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90		
07505702	**Low discrimination**					**07505702**
Bifidobacterium longum subsp. longum		89.98	0.35	FRU 90		
Lactobacillus fermentum		4.22	0.20	RHA 1		
Clostridium clostridioforme		2.47	0.07			
07505710	**Good identification**					**07505710**
					GRA	
Bifidobacterium longum subsp. longum		98.94	0.39	FRU 90	99	
Bacteroides vulgatus		1.06	-.09	RHA 1	1	
07505712	**Acceptable identification**					**07505712**
Bifidobacterium longum subsp. longum		93.17	0.39	FRU 90		
Clostridium clostridioforme		6.83	0.18	RHA 1		
07505720	**Very good identification**					**07505720**
					GRA	
Bacteroides vulgatus		97.80	0.60	FRU 99	1	
Bifidobacterium longum subsp. longum		2.19	0.46		99	
Parabacteroides distasonis		0.01	0.10		1	
07505722	**Low discrimination**					**07505722**
					GRA	
Bifidobacterium longum subsp. longum		51.94	0.46	FRU 90	99	
Parabacteroides distasonis		24.30	0.44	RHA 1	1	
Bacteroides vulgatus		23.41	0.25		1	
07505730	**Low discrimination**					**07505730**
					GRA	
Bifidobacterium longum subsp. longum		79.07	0.50	FRU 90	99	
Bacteroides vulgatus		20.93	0.25	RHA 1	1	
07505732	**Low discrimination**					**07505732**
					GRA	
Bifidobacterium longum subsp. longum		85.55	0.50	FRU 90	99	
Parabacteroides distasonis		12.66	0.39	RHA 1	1	
Clostridium clostridioforme		1.57	0.18		50	
07507500	**Good identification**					**07507500**
Bifidobacterium longum subsp. longum		95.52	0.35	FRU 90		
Lactobacillus fermentum		4.48	0.20	MAN 1		
07507502	**Low discrimination**					**07507502**
Bifidobacterium longum subsp. longum		57.74	0.35	FRU 90		
Bifidobacterium adolescentis		39.55	0.34	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		2.71	0.20					
07507510	**Good identification**							**07507510**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90				
07507512	**Low discrimination**							**07507512**
Bifidobacterium longum subsp. longum		67.04	0.39	FRU 90				
Bifidobacterium adolescentis		32.96	0.36	MAN 1				
07507520	**Good identification**							**07507520**
					GRA			
Bifidobacterium longum subsp. longum		95.23	0.46	FRU 90	99			
Bacteroides vulgatus		4.77	0.09	MAN 1	1			
07507522	**Good identification**							**07507522**
Bifidobacterium longum subsp. longum		100.0	0.46	FRU 90				
07507530	**Good identification**							**07507530**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90				
07507532	**Good identification**							**07507532**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90				
07510000	**Low discrimination**							**07510000**
					COC	GRA	CAT	
Prevotella intermedia		62.64	0.84		1	1	1	
Propionibacterium granulosum		30.06	0.86		1	99	90	
Gemella morbillorum		5.19	0.66		99	99	1	
07510001	**Low discrimination**							**07510001**
					COC	GRA	CAT	
Prevotella intermedia		60.58	0.49	URE 1	1	1	1	
Propionibacterium granulosum		29.07	0.52		1	99	90	
Gemella morbillorum		5.02	0.32		99	99	1	
07510002	**Low discrimination**							**07510002**
					GRA	CAT		
Propionibacterium granulosum		59.64	0.76		99	90		
Actinomyces viscosus		31.62	0.62		99	76		
Prevotella intermedia		5.02	0.49		1	1		
07510003	**Acceptable identification**							**07510003**
Actinomyces viscosus		92.43	0.53	NIT 99				
Propionibacterium granulosum		5.28	0.41					
Actinomyces naeslundii		1.57	0.47					
07510010	**Low discrimination**							**07510010**
					COC	GRA	CAT	
Propionibacterium granulosum		84.99	0.71	MLZ 12	1	99	90	
Prevotella intermedia		13.12	0.49		1	1	1	
Gemella morbillorum		1.09	0.32		99	99	1	
07510011	**Low discrimination**							**07510011**
					GRA	CAT		
Propionibacterium granulosum		84.54	0.37	MLZ 12	99	90		
Prevotella intermedia		13.05	0.15	URE 1	1	1		
Actinomyces naeslundii		2.42	0.25		99	25		
07510012	**Very good identification**							**07510012**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Propionibacterium granulosum		95.23	0.61	MLZ 12	99	90		
Actinomyces viscosus		3.74	0.27		99	76		
Prevotella intermedia		0.59	0.15		1	1		
07510020	**Low discrimination**							**07510020**
					COC	GRA	CAT	
Propionibacterium granulosum		82.88	0.76		1	99	90	
Prevotella intermedia		6.98	0.49		1	1	1	
Gemella morbillorum		6.36	0.50		99	99	1	
07510021	**Low discrimination**							**07510021**
					GRA	CAT		
Propionibacterium granulosum		67.75	0.41	URE 1	99	90		
Actinomyces naeslundii		20.07	0.47		99	25		
Prevotella intermedia		5.70	0.15		1	1		
07510022	**Low discrimination**							**07510022**
					CAT			
Propionibacterium granulosum		89.50	0.65		90			
Lactobacillus fermentum		3.45	0.46		1			
Actinomyces viscosus		1.92	0.27		76			
07510023	**Low discrimination**							**07510023**
Actinomyces naeslundii		40.98	0.47	GAL 96				
Propionibacterium granulosum		34.58	0.31					
Actinomyces viscosus		24.44	0.19					
07510030	**Very good identification**							**07510030**
					GRA	SPO	CAT	
Propionibacterium granulosum		97.76	0.61	MLZ 12	99	1	90	
Clostridium botulinum biovar B		0.87	0.13		99	99	1	
Prevotella intermedia		0.61	0.15		1	1	1	
07510032	**Good identification**							**07510032**
Propionibacterium granulosum		99.69	0.51	MLZ 12				
Actinomyces naeslundii		0.31	0.22					
07510040	**Low discrimination**							**07510040**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		78.12	0.58	RAF 1	99	99	1	
Prevotella intermedia		13.31	0.49		1	1	1	
Propionibacterium granulosum		6.39	0.52		99	1	90	
07510042	**Low discrimination**							**07510042**
					SPO	CAT		
Propionibacterium granulosum		44.48	0.41	SOR 1	1	90		
Actinomyces viscosus		23.58	0.27		1	76		
Clostridium botulinum biovar B		21.98	0.24		99	1		
07510050	**Good identification**							**07510050**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		97.45	0.52	RAF 1	99	99	1	
Propionibacterium granulosum		2.21	0.37		99	1	90	
Prevotella intermedia		0.34	0.15		1	1	1	
07510060	**Acceptable identification**							**07510060**
					CAT	SPO		
Clostridium botulinum biovar B		94.48	0.52	RAF 1	1	99		
Propionibacterium granulosum		3.92	0.41		90	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.76	0.22		1	84	
07510070	**Good identification**						**07510070**
					SPO	CAT	
Clostridium botulinum biovar B		98.86	0.47	RAF 1	99	1	
Propionibacterium granulosum		1.14	0.27		1	90	
07510100	**Low discrimination**						**07510100**
					GRA		
Fusobacterium mortiferum		68.44	0.74	MLT 11	1		
Prevotella intermedia		11.84	0.49		1		
Lactobacillus fermentum		6.63	0.58		99		
07510101	**Low discrimination**						**07510101**
					GRA	SPO	
Actinomyces naeslundii		37.11	0.55	GAL 96	99	1	
Fusobacterium mortiferum		21.43	0.39		1	1	
Clostridium chauvoei		15.88	0.26		99	99	
07510102	**Low discrimination**						**07510102**
					GRA		
Actinomyces viscosus		87.39	0.66	NIT 99	99		
Fusobacterium mortiferum		5.02	0.57		1		
Lactobacillus fermentum		4.37	0.58		99		
07510103	**Very good identification**						**07510103**
Actinomyces viscosus		96.51	0.58	NIT 99			
Actinomyces naeslundii		2.62	0.55				
Propionibacterium propionicum		0.36	0.30				
07510110	**Low discrimination**						**07510110**
					GRA	CAT	
Propionibacterium granulosum		43.90	0.37	LAC 1	99	90	
Fusobacterium mortiferum		39.19	0.39	MLZ 12	1	5	
Prevotella intermedia		6.78	0.15		1	1	
07510120	**Good identification**						**07510120**
					GRA		
Prevotella melaninogenica		93.82	0.66	GAL 99	1		
Clostridium perfringens		3.81	0.49		99		
Lactobacillus fermentum		0.79	0.50		99		
07510122	**Low discrimination**						**07510122**
					GRA		
Prevotella melaninogenica		76.03	0.54	GAL 99	1		
Clostridium perfringens		7.62	0.43		99		
Prevotella oralis		6.98	0.39		1		
07510123	**Low discrimination**						**07510123**
					GRA		
Actinomyces naeslundii		52.18	0.55	GAL 96	99		
Prevotella melaninogenica		21.89	0.19		1		
Actinomyces viscosus		19.44	0.24		99		
07510140	**Low discrimination**						**07510140**
					GRA	SPO	
Propionibacterium propionicum		34.48	0.43	NIT 86	99	1	
Clostridium botulinum biovar B		24.63	0.24	MAN 99	99	99	
Fusobacterium mortiferum		24.27	0.39		1	1	
07510141	**Low discrimination**						**07510141**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		65.94	0.42	GAL 96				
Propionibacterium propionicum		34.06	0.27					
07510142	**Low discrimination**							**07510142**
Propionibacterium propionicum		47.14	0.46	NIT 86				
Actinomyces viscosus		42.82	0.32	MAN 99				
Actinomyces naeslundii		4.51	0.39					
07510161	**Low discrimination**							**07510161**
					GRA			
Actinomyces naeslundii		81.80	0.42	GAL 96	99			
Prevotella melaninogenica		10.31	-.03		1			
Clostridium perfringens		7.89	0.02		99			
07510163	**Good identification**							**07510163**
Actinomyces naeslundii		100.0	0.42	GAL 96				
07510200	**Low discrimination**							**07510200**
					COC	GRA	CAT	
Propionibacterium granulosum		56.72	0.60	RHA 3	1	99	90	
Prevotella intermedia		38.61	0.49		1	1	1	
Gemella morbillorum		3.20	0.32		99	99	1	
07510202	**Low discrimination**							**07510202**
					GRA	CAT		
Propionibacterium granulosum		81.35	0.50	RHA 3	99	90		
Actinomyces viscosus		14.09	0.27		99	76		
Prevotella intermedia		2.24	0.15		1	1		
07510210	**Good identification**							**07510210**
					GRA	CAT		
Propionibacterium granulosum		95.20	0.45	RHA 3	99	90		
Prevotella intermedia		4.80	0.15	MLZ 12	1	1		
07510220	**Acceptable identification**							**07510220**
					COC	GRA	CAT	
Propionibacterium granulosum		94.46	0.50	RHA 3	1	99	90	
Prevotella intermedia		2.60	0.15		1	1	1	
Gemella morbillorum		2.37	0.15		99	99	1	
07510222	**Low discrimination**							**07510222**
					GRA	CAT		
Propionibacterium granulosum		89.70	0.39	RHA 3	99	90		
Prevotella oralis		10.30	0.13		1	1		
07510300	**Low discrimination**							**07510300**
					GRA	CAT		
Fusobacterium mortiferum		62.51	0.39	MLT 11	1	5		
Propionibacterium granulosum		15.88	0.26	RHA 1	99	90		
Prevotella intermedia		10.81	0.15		1	1		
07510400	**Low discrimination**							**07510400**
					COC	GRA	CAT	
Prevotella intermedia		36.15	0.49	ARA 1	1	1	1	
Gemella morbillorum		32.96	0.50		99	99	1	
Propionibacterium granulosum		17.35	0.52		1	99	90	
07510402	**Low discrimination**							**07510402**
					CAT			
Lactobacillus fermentum		62.29	0.54	GAL 99	1			
Propionibacterium granulosum		21.77	0.41		90			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces viscosus		11.54	0.27		76			
07510410	**Low discrimination**							**07510410**
					COC	GRA	CAT	
Propionibacterium granulosum		71.23	0.37	ARA 1	1	99	90	
Prevotella intermedia		11.00	0.15	MLZ 12	1	1	1	
Gemella morbillorum		10.03	0.15		99	99	1	
07510420	**Low discrimination**							**07510420**
					COC	CAT		
Propionibacterium granulosum		33.64	0.41	ARA 1	1	90		
Lactobacillus fermentum		32.08	0.46		1	1		
Gemella morbillorum		28.41	0.33		99	1		
07510422	**Low discrimination**							**07510422**
					CAT			
Lactobacillus fermentum		74.78	0.46	GAL 99	1			
Propionibacterium granulosum		19.61	0.31		90			
Bifidobacterium longum subsp. longum		5.61	0.20		1			
07510500	**Low discrimination**							**07510500**
					GRA			
Lactobacillus fermentum		64.85	0.58	GAL 99	99			
Bifidobacterium longum subsp. longum		24.60	0.43		99			
Fusobacterium mortiferum		6.76	0.39		1			
07510502	**Low discrimination**							**07510502**
Lactobacillus fermentum		61.95	0.58	GAL 99				
Bifidobacterium longum subsp. longum		23.51	0.43					
Actinomyces viscosus		12.51	0.32					
07510510	**Good identification**							**07510510**
Bifidobacterium longum subsp. longum		98.46	0.47	GAL 99				
Lactobacillus fermentum		1.54	0.24					
07510512	**Acceptable identification**							**07510512**
Bifidobacterium longum subsp. longum		94.66	0.47	GAL 99				
Bifidobacterium adolescentis		2.83	0.23					
Lactobacillus fermentum		1.48	0.24					
07510520	**Low discrimination**							**07510520**
					GRA			
Bifidobacterium longum subsp. longum		65.44	0.54	GAL 99	99			
Prevotella melaninogenica		17.15	0.32		1			
Lactobacillus fermentum		14.37	0.50		99			
07510522	**Low discrimination**							**07510522**
					GRA			
Bifidobacterium longum subsp. longum		76.72	0.54	GAL 99	99			
Lactobacillus fermentum		16.85	0.50		99			
Prevotella melaninogenica		3.83	0.19		1			
07510530	**Very good identification**							**07510530**
					GRA			
Bifidobacterium longum subsp. longum		99.72	0.58	GAL 99	99			
Prevotella melaninogenica		0.16	-.03		1			
Lactobacillus fermentum		0.13	0.15		99			
07510532	**Very good identification**							**07510532**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		99.30	0.58	GAL 99	99		
Clostridium clostridioforme		0.27	0.12		50		
Parabacteroides distasonis		0.18	0.15		1		
07511000	**Low discrimination**						**07511000**
					GRA	CAT	
Prevotella intermedia		52.14	0.84		1	1	
Propionibacterium granulosum		25.02	0.86		99	90	
Lactobacillus fermentum		17.90	0.88		99	1	
07511001	**Low discrimination**						**07511001**
					GRA	CAT	
Prevotella intermedia		35.45	0.49	URE 1	1	1	
Actinomyces naeslundii		30.24	0.71		99	25	
Propionibacterium granulosum		17.01	0.52		99	90	
07511002	**Low discrimination**						**07511002**
					CAT		
Lactobacillus fermentum		65.21	0.88		1		
Propionibacterium granulosum		22.79	0.76		90		
Actinomyces viscosus		7.41	0.58		76		
07511003	**Low discrimination**						**07511003**
Actinomyces viscosus		48.96	0.50	NIT 99			
Actinomyces naeslundii		32.46	0.71				
Lactobacillus fermentum		13.06	0.54				
07511010	**Low discrimination**						**07511010**
					GRA	CAT	
Propionibacterium granulosum		81.53	0.71	MLZ 12	99	90	
Prevotella intermedia		12.59	0.49		1	1	
Lactobacillus fermentum		4.32	0.54		99	1	
07511011	**Low discrimination**						**07511011**
					GRA	CAT	
Propionibacterium granulosum		52.81	0.37	MLZ 12	99	90	
Actinomyces naeslundii		36.24	0.49	URE 1	99	25	
Prevotella intermedia		8.15	0.15		1	1	
07511012	**Low discrimination**						**07511012**
					CAT		
Propionibacterium granulosum		78.78	0.61	MLZ 12	90		
Lactobacillus fermentum		16.69	0.54		1		
Actinomyces viscosus		1.90	0.24		76		
07511013	**Low discrimination**						**07511013**
Actinomyces naeslundii		57.77	0.49	MLZ 5			
Propionibacterium granulosum		21.05	0.27				
Actinomyces viscosus		16.72	0.16				
07511020	**Low discrimination**						**07511020**
					GRA	CAT	
Prevotella melaninogenica		37.01	0.66	LAC 99	1	1	
Clostridium perfringens		20.28	0.68		99	1	
Propionibacterium granulosum		19.93	0.76		99	90	
07511021	**Low discrimination**						**07511021**
					GRA		
Actinomyces naeslundii		58.31	0.71		99		
Prevotella melaninogenica		15.23	0.32		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		8.34	0.34		99		
07511022	**Low discrimination**						**07511022**
					GRA		
Lactobacillus fermentum		41.40	0.80		99		
Clostridium perfringens		20.78	0.63		99		
Prevotella melaninogenica		15.35	0.54		1		
07511023	**Low discrimination**						**07511023**
Actinomyces naeslundii		75.00	0.71				
Lactobacillus fermentum		10.06	0.46				
Clostridium perfringens		5.05	0.29				
07511030	**Low discrimination**						**07511030**
					GRA	CAT	
Propionibacterium granulosum		75.69	0.61	MLZ 12	99	90	
Prevotella melaninogenica		10.41	0.32		1	1	
Clostridium perfringens		5.70	0.34		99	1	
07511031	**Low discrimination**						**07511031**
					GRA	CAT	
Actinomyces naeslundii		70.70	0.49	MLZ 5	99	25	
Propionibacterium granulosum		25.76	0.27		99	90	
Prevotella melaninogenica		3.54	-.03		1	1	
07511032	**Low discrimination**						**07511032**
					CAT		
Propionibacterium granulosum		57.64	0.51	MLZ 12	90		
Lactobacillus fermentum		16.29	0.46		1		
Clostridium perfringens		8.18	0.29		1		
07511033	**Acceptable identification**						**07511033**
Actinomyces naeslundii		91.65	0.49	MLZ 5			
Propionibacterium granulosum		8.35	0.16				
07511040	**Low discrimination**						**07511040**
					GRA	CAT	
Prevotella intermedia		46.77	0.49	SOR 1	1	1	
Propionibacterium granulosum		22.44	0.52		99	90	
Lactobacillus fermentum		16.05	0.54		99	1	
07511041	**Low discrimination**						**07511041**
					GRA	CAT	
Actinomyces naeslundii		89.10	0.58		99	25	
Prevotella intermedia		5.98	0.15		1	1	
Propionibacterium granulosum		2.87	0.18		99	90	
07511042	**Low discrimination**						**07511042**
					CAT		
Lactobacillus fermentum		52.57	0.54	SOR 1	1		
Propionibacterium granulosum		18.38	0.41		90		
Actinomyces naeslundii		15.37	0.55		25		
07511043	**Good identification**						**07511043**
Actinomyces naeslundii		90.14	0.58				
Actinomyces viscosus		7.78	0.16				
Lactobacillus fermentum		2.08	0.20				
07511050	**Low discrimination**						**07511050**
					GRA	SPO CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		57.67	0.37	MLZ 12	99	1	90	
Clostridium botulinum biovar B		25.73	0.18	SOR 1	99	99	1	
Prevotella intermedia		8.90	0.15		1	1	1	
07511051	**Good identification**							**07511051**
Actinomyces naeslundii		100.0	0.36	MLZ 5				
07511053	**Good identification**							**07511053**
Actinomyces naeslundii		100.0	0.36	MLZ 5				
07511060	**Low discrimination**							**07511060**
					GRA	CAT		
Clostridium perfringens		79.24	0.56	LAC 88	99	1		
Prevotella melaninogenica		7.67	0.32		1	1		
Propionibacterium granulosum		4.13	0.41		99	90		
07511061	**Low discrimination**							**07511061**
					GRA			
Actinomyces naeslundii		85.52	0.58		99			
Clostridium perfringens		13.21	0.22		99			
Prevotella melaninogenica		1.28	-.03		1			
07511062	**Low discrimination**							**07511062**
Clostridium perfringens		77.10	0.50	LAC 88				
Lactobacillus fermentum		8.15	0.46					
Actinomyces naeslundii		7.15	0.55					
07511063	**Good identification**							**07511063**
Actinomyces naeslundii		93.23	0.58					
Clostridium perfringens		6.77	0.16					
07511071	**Good identification**							**07511071**
Actinomyces naeslundii		100.0	0.36	MLZ 5				
07511073	**Good identification**							**07511073**
Actinomyces naeslundii		100.0	0.36	MLZ 5				
07511100	**Low discrimination**							**07511100**
					GRA			
Lactobacillus fermentum		59.78	0.92		99			
Prevotella melaninogenica		23.78	0.66		1			
Fusobacterium mortiferum		9.35	0.77		1			
07511101	**Low discrimination**							**07511101**
					SPO			
Clostridium chauvoei		55.94	0.60	RAF 1	99			
Actinomyces naeslundii		31.69	0.79		1			
Lactobacillus fermentum		7.31	0.58		1			
07511102	**Low discrimination**							**07511102**
					GRA			
Lactobacillus fermentum		74.80	0.92		99			
Actinomyces viscosus		9.26	0.62		99			
Prevotella melaninogenica		5.67	0.54		1			
07511103	**Low discrimination**							**07511103**
					SPO			
Actinomyces naeslundii		39.51	0.79		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		37.22	0.54		1		
Clostridium chauvoei		12.31	0.47		99		
07511110	**Low discrimination**						**07511110**
					GRA		
Lactobacillus fermentum		38.97	0.58	MLZ 1	99		
Bifidobacterium longum subsp. longum		25.17	0.47		99		
Prevotella melaninogenica		15.50	0.32		1		
07511111	**Low discrimination**						**07511111**
					SPO		
Actinomyces naeslundii		69.97	0.57	MLZ 5	1		
Clostridium chauvoei		23.70	0.26		99		
Lactobacillus fermentum		3.10	0.24		1		
07511112	**Low discrimination**						**07511112**
Lactobacillus fermentum		38.51	0.58	MLZ 1			
Bifidobacterium longum subsp. longum		24.88	0.47				
Bifidobacterium adolescentis		11.02	0.43				
07511113	**Low discrimination**						**07511113**
					SPO		
Actinomyces naeslundii		75.41	0.57	MLZ 5	1		
Actinomyces viscosus		13.63	0.20		1		
Clostridium chauvoei		4.51	0.13		99		
07511120	**Very good identification**						**07511120**
					GRA		
Prevotella melaninogenica		95.08	1.00		1		
Clostridium perfringens		3.86	0.83		99		
Lactobacillus fermentum		0.81	0.84		99		
07511121	**Low discrimination**						**07511121**
					GRA	SPO	
Prevotella melaninogenica		81.58	0.66	URE 1	1	1	
Actinomyces naeslundii		8.98	0.79		99	1	
Clostridium chauvoei		5.29	0.52		99	99	
07511122	**Low discrimination**						**07511122**
					GRA		
Prevotella melaninogenica		78.90	0.88		1		
Clostridium perfringens		7.91	0.78		99		
Prevotella oralis		7.24	0.73		1		
07511123	**Low discrimination**						**07511123**
					GRA		
Prevotella melaninogenica		52.44	0.54	URE 1	1		
Actinomyces naeslundii		30.31	0.79		99		
Clostridium perfringens		5.26	0.43		99		
07511130	**Low discrimination**						**07511130**
					GRA		
Prevotella melaninogenica		88.86	0.66	MLZ 1	1		
Bifidobacterium longum subsp. longum		5.83	0.58		99		
Clostridium perfringens		3.61	0.49		99		
07511132	**Low discrimination**						**07511132**
					GRA	CAT	
Prevotella melaninogenica		51.56	0.54	MLZ 1	1	1	
Bifidobacterium longum subsp. longum		17.76	0.58		99	1	
Capnocytophaga ochracea		12.05	0.47		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07511133	**Low discrimination**						**07511133**
					GRA		
Actinomyces naeslundii		60.38	0.57	MLZ 5	99		
Prevotella melaninogenica		20.05	0.19		1		
Bifidobacterium longum subsp. longum		6.91	0.24		99		
07511140	**Low discrimination**						**07511140**
Lactobacillus fermentum		31.28	0.58	SOR 1			
Clostridium perfringens		29.16	0.45				
Actinomyces naeslundii		15.96	0.63				
07511141	**Low discrimination**						**07511141**
					SPO		
Actinomyces naeslundii		86.71	0.66		1		
Clostridium chauvoei		8.76	0.26		99		
Propionibacterium propionicum		1.87	0.27		1		
07511142	**Low discrimination**						**07511142**
Lactobacillus fermentum		37.68	0.58	SOR 1			
Actinomyces naeslundii		19.22	0.63				
Clostridium perfringens		16.53	0.39				
07511143	**Low discrimination**						**07511143**
Actinomyces naeslundii		89.05	0.66				
Actinomyces viscosus		4.80	0.20				
Propionibacterium propionicum		2.87	0.30				
07511150	**Low discrimination**						**07511150**
Actinomyces naeslundii		45.27	0.41	MLZ 5			
Lactobacillus fermentum		17.03	0.24				
Clostridium perfringens		15.88	0.11				
07511151	**Good identification**						**07511151**
Actinomyces naeslundii		100.0	0.44	MLZ 5			
07511153	**Good identification**						**07511153**
Actinomyces naeslundii		100.0	0.44	MLZ 5			
07511160	**Low discrimination**						**07511160**
					GRA		
Prevotella melaninogenica		55.77	0.66	SOR 1	1		
Clostridium perfringens		42.68	0.71		99		
Actinomyces naeslundii		0.72	0.63		99		
07511161	**Low discrimination**						**07511161**
					GRA		
Actinomyces naeslundii		51.13	0.66		99		
Prevotella melaninogenica		26.58	0.32		1		
Clostridium perfringens		20.34	0.37		99		
07511162	**Low discrimination**						**07511162**
					GRA	CAT	
Clostridium perfringens		56.70	0.65		99	1	
Prevotella melaninogenica		29.99	0.54		1	1	
Capnocytophaga ochracea		7.01	0.47		1	90	
07511163	**Low discrimination**						**07511163**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		75.26	0.66		99		
Clostridium perfringens		14.09	0.31		99		
Prevotella melaninogenica		7.45	0.19		1		
07511171	**Low discrimination**						**07511171**
					GRA		
Actinomyces naeslundii		85.03	0.44	MLZ 5	99		
Prevotella melaninogenica		8.48	-.03		1		
Clostridium perfringens		6.49	0.02		99		
07511173	**Good identification**						**07511173**
Actinomyces naeslundii		100.0	0.44	MLZ 5			
07511200	**Low discrimination**						**07511200**
					GRA	CAT	
Propionibacterium granulosum		50.42	0.60	RHA 3	99	90	
Prevotella intermedia		34.32	0.49		1	1	
Lactobacillus fermentum		11.78	0.54		99	1	
07511202	**Low discrimination**						**07511202**
					GRA	CAT	
Propionibacterium granulosum		44.31	0.50	RHA 3	99	90	
Lactobacillus fermentum		41.40	0.54		99	1	
Prevotella oralis		6.63	0.25		1	1	
07511210	**Acceptable identification**						**07511210**
					GRA	CAT	
Propionibacterium granulosum		93.66	0.45	RHA 3	99	90	
Prevotella intermedia		4.72	0.15	MLZ 12	1	1	
Lactobacillus fermentum		1.62	0.20		99	1	
07511220	**Low discrimination**						**07511220**
					GRA	CAT	
Propionibacterium granulosum		39.86	0.50	RHA 3	99	90	
Prevotella melaninogenica		24.17	0.32		1	1	
Clostridium perfringens		13.24	0.34		99	1	
07511221	**Low discrimination**						**07511221**
					GRA	CAT	
Actinomyces naeslundii		59.11	0.37	RHA 1	99	25	
Propionibacterium granulosum		25.46	0.16		99	90	
Prevotella melaninogenica		15.43	-.03		1	1	
07511222	**Low discrimination**						**07511222**
					GRA	CAT	
Prevotella oralis		75.65	0.47	LAC 94	1	1	
Lactobacillus fermentum		8.29	0.46		99	1	
Propionibacterium granulosum		6.66	0.39		99	90	
07511300	**Low discrimination**						**07511300**
					GRA		
Lactobacillus fermentum		55.57	0.58	RHA 1	99		
Prevotella melaninogenica		22.11	0.32		1		
Fusobacterium mortiferum		8.70	0.42		1		
07511302	**Low discrimination**						**07511302**
					GRA		
Prevotella oralis		53.28	0.45	NAG 95	1		
Lactobacillus fermentum		34.68	0.58		99		
Actinomyces viscosus		4.29	0.28		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07511303	**Low discrimination**						**07511303**
					GRA		
Actinomyces naeslundii		35.23	0.45	RHA 1	99		
Actinomyces viscosus		33.19	0.20		99		
Prevotella oralis		12.49	0.11		1		
07511320	**Low discrimination**						**07511320**
					GRA		
Prevotella melaninogenica		84.88	0.66	RHA 1	1		
Bacteroides vulgatus		7.40	0.46		1		
Clostridium perfringens		3.45	0.49		99		
07511322	**Low discrimination**						**07511322**
Prevotella oralis		69.06	0.67				
Prevotella melaninogenica		17.74	0.54				
Parabacteroides distasonis		10.04	0.66				
07511332	**Low discrimination**						**07511332**
Parabacteroides distasonis		84.01	0.61	XYL 90			
Prevotella oralis		10.84	0.33				
Prevotella melaninogenica		2.78	0.19				
07511400	**Good identification**						**07511400**
					COC	GRA	
Lactobacillus fermentum		93.83	0.88		1	99	
Prevotella intermedia		2.76	0.49		1	1	
Gemella morbillorum		1.36	0.45		99	99	
07511401	**Acceptable identification**						**07511401**
					GRA		
Lactobacillus fermentum		91.80	0.54	URE 1	99		
Prevotella intermedia		2.70	0.15		1		
Actinomyces naeslundii		2.31	0.37		99		
07511402	**Very good identification**						**07511402**
					CAT		
Lactobacillus fermentum		98.81	0.88		1		
Bifidobacterium longum subsp. longum		0.62	0.43		1		
Propionibacterium granulosum		0.35	0.41		90		
07511403	**Acceptable identification**						**07511403**
Lactobacillus fermentum		93.53	0.54	URE 1			
Actinomyces viscosus		3.54	0.16				
Actinomyces naeslundii		2.35	0.37				
07511410	**Low discrimination**						**07511410**
					CAT		
Bifidobacterium longum subsp. longum		46.06	0.47	LAC 99	1		
Lactobacillus fermentum		43.70	0.54		1		
Propionibacterium granulosum		8.33	0.37		90		
07511412	**Low discrimination**						**07511412**
					CAT		
Bifidobacterium longum subsp. longum		49.17	0.47	LAC 99	1		
Lactobacillus fermentum		46.65	0.54		1		
Propionibacterium granulosum		2.22	0.27		90		
07511420	**Low discrimination**						**07511420**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		88.88	0.80		99		
Bifidobacterium longum subsp. longum		6.67	0.54		99		
Prevotella melaninogenica		1.75	0.32		1		
07511421	**Low discrimination**						**07511421**
Lactobacillus fermentum		85.47	0.46	URE 1			
Actinomyces naeslundii		6.44	0.37				
Bifidobacterium longum subsp. longum		6.41	0.20				
07511422	**Good identification**						**07511422**
Lactobacillus fermentum		91.66	0.80				
Bifidobacterium longum subsp. longum		6.88	0.54				
Clostridium perfringens		0.47	0.29				
07511423	**Low discrimination**						**07511423**
Lactobacillus fermentum		86.93	0.46	URE 1			
Actinomyces naeslundii		6.55	0.37				
Bifidobacterium longum subsp. longum		6.52	0.20				
07511430	**Good identification**						**07511430**
					CAT		
Bifidobacterium longum subsp. longum		91.58	0.58	LAC 99	1		
Lactobacillus fermentum		7.24	0.46		1		
Propionibacterium granulosum		1.04	0.27		90		
07511432	**Good identification**						**07511432**
					CAT		
Bifidobacterium longum subsp. longum		91.97	0.58	LAC 99	1		
Lactobacillus fermentum		7.27	0.46		1		
Propionibacterium granulosum		0.26	0.16		90		
07511440	**Acceptable identification**						**07511440**
					COC	GRA	
Lactobacillus fermentum		93.96	0.54	SOR 1	1	99	
Prevotella intermedia		2.77	0.15		1	1	
Gemella morbillorum		1.36	0.11		99	99	
07511442	**Good identification**						**07511442**
Lactobacillus fermentum		99.38	0.54	SOR 1			
Bifidobacterium longum subsp. longum		0.62	0.09				
07511460	**Low discrimination**						**07511460**
Lactobacillus fermentum		77.05	0.46	SOR 1			
Clostridium perfringens		15.66	0.22				
Bifidobacterium longum subsp. longum		5.78	0.20				
07511462	**Low discrimination**						**07511462**
Lactobacillus fermentum		84.32	0.46	SOR 1			
Clostridium perfringens		8.06	0.16				
Bifidobacterium longum subsp. longum		6.33	0.20				
07511500	**Low discrimination**						**07511500**
					GRA		
Lactobacillus fermentum		72.08	0.92		99		
Bifidobacterium longum subsp. longum		27.35	0.78		99		
Prevotella melaninogenica		0.29	0.32		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07511501	**Low discrimination**					**07511501**
					SPO	
Lactobacillus fermentum		66.47	0.58	URE 1	1	
Bifidobacterium longum subsp. longum		25.22	0.43		1	
Clostridium chauvoei		5.14	0.26		99	
07511502	**Low discrimination**					**07511502**
Lactobacillus fermentum		71.35	0.92			
Bifidobacterium longum subsp. longum		27.07	0.78			
Bifidobacterium adolescentis		1.13	0.56			
07511503	**Low discrimination**					**07511503**
Lactobacillus fermentum		66.80	0.58	URE 1		
Bifidobacterium longum subsp. longum		25.34	0.43			
Actinomyces naeslundii		2.93	0.45			
07511510	**Very good identification**					**07511510**
Bifidobacterium longum subsp. longum		98.42	0.82			
Lactobacillus fermentum		1.54	0.58			
Bifidobacterium adolescentis		0.03	0.23			
07511511	**Good identification**					**07511511**
Bifidobacterium longum subsp. longum		98.12	0.47	URE 1		
Lactobacillus fermentum		1.53	0.24			
Actinomyces naeslundii		0.35	0.23			
07511512	**Very good identification**					**07511512**
Bifidobacterium longum subsp. longum		95.56	0.82			
Bifidobacterium adolescentis		2.86	0.57			
Lactobacillus fermentum		1.49	0.58			
07511513	**Good identification**					**07511513**
Bifidobacterium longum subsp. longum		95.32	0.47	URE 1		
Bifidobacterium adolescentis		2.85	0.23			
Lactobacillus fermentum		1.49	0.24			
07511520	**Low discrimination**					**07511520**
					GRA	
Bifidobacterium longum subsp. longum		65.42	0.88		99	
Prevotella melaninogenica		17.15	0.66		1	
Lactobacillus fermentum		14.37	0.84		99	
07511521	**Low discrimination**					**07511521**
					GRA	
Bifidobacterium longum subsp. longum		63.53	0.54	URE 1	99	
Prevotella melaninogenica		16.65	0.32		1	
Lactobacillus fermentum		13.96	0.50		99	
07511522	**Low discrimination**					**07511522**
					GRA	
Bifidobacterium longum subsp. longum		76.68	0.88		99	
Lactobacillus fermentum		16.84	0.84		99	
Prevotella melaninogenica		3.83	0.54		1	
07511523	**Low discrimination**					**07511523**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		74.89	0.54	URE 1	99		
Lactobacillus fermentum		16.45	0.50		99		
Prevotella melaninogenica		3.74	0.19		1		
07511530	**Excellent identification**						**07511530**
					GRA		
Bifidobacterium longum subsp. longum		99.67	0.92		99		
Prevotella melaninogenica		0.16	0.32		1		
Lactobacillus fermentum		0.13	0.50		99		
07511531	**Very good identification**						**07511531**
					GRA		
Bifidobacterium longum subsp. longum		99.63	0.58	URE 1	99		
Prevotella melaninogenica		0.16	-.03		1		
Lactobacillus fermentum		0.13	0.15		99		
07511532	**Excellent identification**						**07511532**
					GRA		
Bifidobacterium longum subsp. longum		99.52	0.92		99		
Parabacteroides distasonis		0.18	0.49		1		
Lactobacillus fermentum		0.13	0.50		99		
07511533	**Very good identification**						**07511533**
					GRA		
Bifidobacterium longum subsp. longum		99.48	0.58	URE 1	99		
Parabacteroides distasonis		0.18	0.15		1		
Lactobacillus fermentum		0.13	0.15		99		
07511540	**Low discrimination**						**07511540**
Lactobacillus fermentum		71.34	0.58	SOR 1			
Bifidobacterium longum subsp. longum		27.07	0.43				
Clostridium perfringens		0.67	0.11				
07511542	**Low discrimination**						**07511542**
Lactobacillus fermentum		61.76	0.58	SOR 1			
Bifidobacterium longum subsp. longum		23.43	0.43				
Bifidobacterium adolescentis		11.96	0.40				
07511550	**Good identification**						**07511550**
Bifidobacterium longum subsp. longum		98.10	0.47	SOR 1			
Lactobacillus fermentum		1.53	0.24				
Bifidobacterium adolescentis		0.36	0.07				
07511552	**Low discrimination**						**07511552**
Bifidobacterium longum subsp. longum		72.06	0.47	SOR 1			
Bifidobacterium adolescentis		26.39	0.42				
Lactobacillus fermentum		1.13	0.24				
07511560	**Low discrimination**						**07511560**
					GRA		
Bifidobacterium longum subsp. longum		57.40	0.54	SOR 1	99		
Prevotella melaninogenica		15.05	0.32		1		
Lactobacillus fermentum		12.61	0.50		99		
07511562	**Low discrimination**						**07511562**
					GRA	CAT	
Bifidobacterium longum subsp. longum		63.23	0.54	SOR 1	99	1	
Lactobacillus fermentum		13.89	0.50		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		12.88	0.34		1	90	
07511570	**Very good identification**						**07511570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.45	0.58	SOR 1	99	1	
Prevotella melaninogenica		0.16	-.03		1	1	
Capnocytophaga ochracea		0.15	0.01		1	90	
07511572	**Very good identification**						**07511572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.30	0.58	SOR 1	99	1	
Capnocytophaga ochracea		1.31	0.17		1	90	
Parabacteroides distasonis		0.18	0.15		1	76	
07511600	**Acceptable identification**						**07511600**
					GRA	CAT	
Lactobacillus fermentum		90.83	0.54	RHA 1	99	1	
Propionibacterium granulosum		3.93	0.26		99	90	
Prevotella intermedia		2.67	0.15		1	1	
07511602	**Good identification**						**07511602**
					CAT		
Lactobacillus fermentum		96.75	0.54	RHA 1	1		
Clostridium clostridioforme		1.60	0.15		33		
Propionibacterium granulosum		1.05	0.16		90		
07511622	**Low discrimination**						**07511622**
					GRA		
Lactobacillus fermentum		73.09	0.46	RHA 1	99		
Parabacteroides distasonis		9.41	0.28		1		
Prevotella oralis		6.73	0.13		1		
07511700	**Low discrimination**						**07511700**
					GRA		
Lactobacillus fermentum		54.23	0.58	RHA 1	99		
Bacteroides vulgatus		24.98	0.31		1		
Bifidobacterium longum subsp. longum		20.57	0.43		99		
07511702	**Low discrimination**						**07511702**
Lactobacillus fermentum		67.63	0.58	RHA 1			
Bifidobacterium longum subsp. longum		25.66	0.43				
Clostridium clostridioforme		2.29	0.24				
07511710	**Good identification**						**07511710**
					GRA		
Bifidobacterium longum subsp. longum		97.77	0.47	RHA 1	99		
Lactobacillus fermentum		1.53	0.24		99		
Bacteroides vulgatus		0.70	-.04		1		
07511712	**Low discrimination**						**07511712**
Bifidobacterium longum subsp. longum		76.53	0.47	RHA 1			
Clostridium clostridioforme		18.31	0.35				
Bifidobacterium adolescentis		2.29	0.23				
07511720	**Very good identification**						**07511720**
					GRA		
Bacteroides vulgatus		95.18	0.65	XYL 98	1		
Bifidobacterium longum subsp. longum		3.17	0.54		99		
Prevotella melaninogenica		0.83	0.32		1		
07511722	**Low discrimination**						**07511722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		48.56	0.62	XYL 90	1		
Bifidobacterium longum subsp. longum		28.32	0.54		99		
Bacteroides vulgatus		8.60	0.31		1		
07511730	**Low discrimination**						**07511730**
					GRA		
Bifidobacterium longum subsp. longum		84.27	0.58	RHA 1	99		
Bacteroides vulgatus		15.02	0.31		1		
Parabacteroides distasonis		0.46	0.23		1		
07511732	**Low discrimination**						**07511732**
					GRA		
Bifidobacterium longum subsp. longum		62.20	0.58	RHA 1	99		
Parabacteroides distasonis		33.74	0.57		1		
Clostridium clostridioforme		3.72	0.35		50		
07512000	**Very good identification**						**07512000**
					GRA	CAT	
Propionibacterium granulosum		96.47	0.84		99	90	
Prevotella intermedia		2.69	0.49		1	1	
Propionibacterium propionicum		0.22	0.43		99	1	
07512001	**Low discrimination**						**07512001**
					GRA	CAT	
Propionibacterium granulosum		80.39	0.50	URE 1	99	90	
Actinomyces naeslundii		15.32	0.52		99	25	
Prevotella intermedia		2.24	0.15		1	1	
07512002	**Very good identification**						**07512002**
					CAT		
Propionibacterium granulosum		97.20	0.74		90		
Propionibacterium propionicum		1.35	0.46		1		
Actinomyces viscosus		0.69	0.27		76		
07512003	**Low discrimination**						**07512003**
Propionibacterium granulosum		46.52	0.39	URE 1			
Actinomyces naeslundii		35.47	0.52				
Actinomyces viscosus		10.90	0.19				
07512010	**Very good identification**						**07512010**
					GRA	CAT	
Propionibacterium granulosum		99.74	0.69	MLZ 12	99	90	
Prevotella intermedia		0.21	0.15		1	1	
Actinomyces naeslundii		0.05	0.27		99	25	
07512012	**Very good identification**						**07512012**
					CAT		
Propionibacterium granulosum		99.70	0.59	MLZ 12	90		
Actinomyces naeslundii		0.20	0.27		25		
Propionibacterium propionicum		0.10	0.12		1		
07512020	**Very good identification**						**07512020**
					GRA	CAT	
Propionibacterium granulosum		99.25	0.74		99	90	
Actinomyces naeslundii		0.51	0.49		99	25	
Prevotella intermedia		0.11	0.15		1	1	
07512021	**Low discrimination**						**07512021**
Propionibacterium granulosum		56.74	0.39	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		43.26	0.52				
07512022	**Very good identification**						**07512022**
					CAT		
Propionibacterium granulosum		97.93	0.63		90		
Actinomyces naeslundii		2.01	0.49		25		
Propionibacterium propionicum		0.06	0.12		1		
07512023	**Low discrimination**						**07512023**
Actinomyces naeslundii		75.31	0.52	GAL 96			
Propionibacterium granulosum		24.70	0.29				
07512030	**Very good identification**						**07512030**
Propionibacterium granulosum		99.80	0.59	MLZ 12			
Actinomyces naeslundii		0.20	0.27				
07512032	**Good identification**						**07512032**
Propionibacterium granulosum		99.21	0.48	MLZ 12			
Actinomyces naeslundii		0.79	0.27				
07512040	**Low discrimination**						**07512040**
					SPO	CAT	
Propionibacterium granulosum		63.72	0.50	SOR 1	1	90	
Propionibacterium propionicum		14.61	0.43		1	1	
Clostridium botulinum biovar B		10.43	0.24		99	1	
07512041	**Low discrimination**						**07512041**
					CAT		
Actinomyces naeslundii		48.60	0.39	GAL 96	25		
Propionibacterium propionicum		36.80	0.27		1		
Propionibacterium granulosum		14.60	0.16		90		
07512042	**Low discrimination**						**07512042**
					CAT		
Propionibacterium propionicum		54.56	0.46	NIT 86	1		
Propionibacterium granulosum		39.67	0.39	LAC 99	90		
Actinomyces naeslundii		3.56	0.36		25		
07512060	**Low discrimination**						**07512060**
					SPO	CAT	
Propionibacterium granulosum		70.81	0.39	SOR 1	1	90	
Clostridium botulinum biovar B		22.84	0.18		99	1	
Actinomyces naeslundii		6.35	0.36		1	25	
07512061	**Good identification**						**07512061**
Actinomyces naeslundii		100.0	0.39	GAL 96			
07512063	**Good identification**						**07512063**
Actinomyces naeslundii		100.0	0.39	GAL 96			
07512100	**Good identification**						**07512100**
					CAT		
Propionibacterium propionicum		93.52	0.78	NIT 86	1		
Propionibacterium granulosum		4.12	0.50		90		
Actinomyces naeslundii		1.49	0.57		25		
07512101	**Low discrimination**						**07512101**
					CAT		
Propionibacterium propionicum		82.03	0.61	NIT 86	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		17.65	0.60	URE 10	25	
Propionibacterium granulosum		0.33	0.16		90	
07512102	**Very good identification**					**07512102**
Propionibacterium propionicum		97.26	0.81	NIT 86		
Actinomyces naeslundii		1.03	0.57			
Actinomyces viscosus		0.89	0.32			
07512103	**Low discrimination**					**07512103**
Propionibacterium propionicum		85.40	0.64	NIT 86		
Actinomyces naeslundii		12.25	0.60	URE 10		
Actinomyces viscosus		2.35	0.24			
07512110	**Low discrimination**					**07512110**
					CAT	
Propionibacterium propionicum		59.60	0.43	NIT 86	1	
Propionibacterium granulosum		35.45	0.35	MLZ 1	90	
Actinomyces naeslundii		4.95	0.35		25	
07512111	**Low discrimination**					**07512111**
Actinomyces naeslundii		52.85	0.38	GAL 96		
Propionibacterium propionicum		47.15	0.27	MLZ 5		
07512112	**Low discrimination**					**07512112**
					CAT	
Propionibacterium propionicum		85.83	0.46	NIT 86	1	
Propionibacterium granulosum		8.51	0.25	MLZ 1	90	
Actinomyces naeslundii		4.75	0.35		25	
07512121	**Good identification**					**07512121**
					GRA	
Actinomyces naeslundii		94.48	0.60	GAL 96	99	
Propionibacterium propionicum		4.44	0.27		99	
Prevotella melaninogenica		1.08	-.03		1	
07512122	**Low discrimination**					**07512122**
					GRA	
Actinomyces naeslundii		39.61	0.57	GAL 96	99	
Propionibacterium propionicum		37.67	0.46		99	
Prevotella melaninogenica		12.84	0.19		1	
07512123	**Good identification**					**07512123**
Actinomyces naeslundii		93.42	0.60	GAL 96		
Propionibacterium propionicum		6.58	0.30			
07512131	**Good identification**					**07512131**
Actinomyces naeslundii		100.0	0.38	GAL 96		
07512132	**Low discrimination**					**07512132**
					CAT	
Actinomyces naeslundii		61.34	0.35	GAL 96	25	
Propionibacterium granulosum		27.47	0.14	MLZ 5	90	
Propionibacterium propionicum		11.19	0.12		1	
07512133	**Good identification**					**07512133**
Actinomyces naeslundii		100.0	0.38	GAL 96		
07512140	**Very good identification**					**07512140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Propionibacterium propionicum		99.68	0.78	NIT 86	1		
Actinomyces naeslundii		0.28	0.44		25		
Propionibacterium granulosum		0.04	0.16		90		
07512141	**Very good identification**						**07512141**
Propionibacterium propionicum		96.34	0.61	NIT 86			
Actinomyces naeslundii		3.66	0.47	URE 10			
07512142	**Very good identification**						**07512142**
Propionibacterium propionicum		99.81	0.81	NIT 86			
Actinomyces naeslundii		0.19	0.44				
07512143	**Very good identification**						**07512143**
Propionibacterium propionicum		97.53	0.64	NIT 86			
Actinomyces naeslundii		2.47	0.47	URE 10			
07512150	**Good identification**						**07512150**
Propionibacterium propionicum		100.0	0.43	NIT 86			
07512152	**Good identification**						**07512152**
Propionibacterium propionicum		100.0	0.46	NIT 86			
07512160	**Low discrimination**						**07512160**
					GRA		
Propionibacterium propionicum		75.40	0.43	NIT 86	99		
Actinomyces naeslundii		20.99	0.44	NAG 1	99		
Prevotella melaninogenica		2.04	-.03		1		
07512161	**Low discrimination**						**07512161**
Actinomyces naeslundii		78.98	0.47	GAL 96			
Propionibacterium propionicum		21.02	0.27				
07512162	**Low discrimination**						**07512162**
Propionibacterium propionicum		84.35	0.46	NIT 86			
Actinomyces naeslundii		15.65	0.44	NAG 1			
07512163	**Low discrimination**						**07512163**
Actinomyces naeslundii		71.47	0.47	GAL 96			
Propionibacterium propionicum		28.53	0.30				
07512200	**Very good identification**						**07512200**
					GRA	CAT	
Propionibacterium granulosum		99.10	0.58	RHA 3	99	90	
Prevotella intermedia		0.90	0.15		1	1	
07512202	**Good identification**						**07512202**
					CAT		
Propionibacterium granulosum		99.55	0.48	RHA 3	90		
Propionibacterium propionicum		0.45	0.12		1		
07512210	**Good identification**						**07512210**
Propionibacterium granulosum		100.0	0.43	RHA 3			
07512220	**Good identification**						**07512220**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		100.0	0.48	RHA 3				
07512222	**Good identification**							**07512222**
Propionibacterium granulosum		100.0	0.37	RHA 3				
07512300	**Low discrimination**							**07512300**
					CAT			
Propionibacterium propionicum		86.89	0.43	NIT 86	1			
Propionibacterium granulosum		11.72	0.24	RHA 1	90			
Actinomyces naeslundii		1.39	0.23		25			
07512302	**Good identification**							**07512302**
					CAT			
Propionibacterium propionicum		96.80	0.46	NIT 86	1			
Propionibacterium granulosum		2.18	0.14	RHA 1	90			
Actinomyces naeslundii		1.03	0.23		25			
07512340	**Good identification**							**07512340**
Propionibacterium propionicum		100.0	0.43	NIT 86				
07512342	**Good identification**							**07512342**
Propionibacterium propionicum		100.0	0.46	NIT 86				
07512400	**Acceptable identification**							**07512400**
					COC	GRA	CAT	
Propionibacterium granulosum		94.08	0.50	ARA 1	1	99	90	
Prevotella intermedia		2.63	0.15		1	1	1	
Gemella morbillorum		2.39	0.15		99	99	1	
07512402	**Good identification**							**07512402**
					CAT			
Propionibacterium granulosum		95.04	0.39	ARA 1	90			
Lactobacillus fermentum		3.64	0.20		1			
Propionibacterium propionicum		1.32	0.12		1			
07512420	**Good identification**							**07512420**
Propionibacterium granulosum		100.0	0.39	ARA 1				
07512500	**Low discrimination**							**07512500**
					CAT			
Propionibacterium propionicum		86.65	0.43	NIT 86	1			
Lactobacillus fermentum		5.91	0.24	ARA 1	1			
Propionibacterium granulosum		3.82	0.16		90			
07512502	**Low discrimination**							**07512502**
Propionibacterium propionicum		88.38	0.46	NIT 86				
Bifidobacterium adolescentis		5.15	0.20	ARA 1				
Lactobacillus fermentum		4.02	0.24					
07512540	**Good identification**							**07512540**
Propionibacterium propionicum		100.0	0.43	NIT 86				
07512542	**Good identification**							**07512542**
Propionibacterium propionicum		100.0	0.46	NIT 86				
07513000	**Good identification**							**07513000**
					GRA	CAT		
Propionibacterium granulosum		93.19	0.84		99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		2.87	0.73		99	25	
Prevotella intermedia		2.60	0.49		1	1	
07513001	**Low discrimination**						**07513001**
					GRA	CAT	
Actinomyces naeslundii		81.14	0.76		99	25	
Propionibacterium granulosum		17.74	0.50		99	90	
Prevotella intermedia		0.50	0.15		1	1	
07513002	**Low discrimination**						**07513002**
					CAT		
Propionibacterium granulosum		84.60	0.74		90		
Actinomyces naeslundii		10.42	0.73		25		
Lactobacillus fermentum		3.24	0.54		1		
07513003	**Good identification**						**07513003**
					CAT		
Actinomyces naeslundii		93.20	0.76		25		
Propionibacterium granulosum		5.09	0.39		90		
Propionibacterium propionicum		0.78	0.30		1		
07513010	**Very good identification**						**07513010**
					GRA	CAT	
Propionibacterium granulosum		98.56	0.69	MLZ 12	99	90	
Actinomyces naeslundii		1.17	0.51		99	25	
Prevotella intermedia		0.20	0.15		1	1	
07513011	**Low discrimination**						**07513011**
Actinomyces naeslundii		63.84	0.54	MLZ 5			
Propionibacterium granulosum		36.16	0.35				
07513012	**Very good identification**						**07513012**
					CAT		
Propionibacterium granulosum		95.01	0.59	MLZ 12	90		
Actinomyces naeslundii		4.52	0.51		25		
Lactobacillus fermentum		0.27	0.20		1		
07513013	**Low discrimination**						**07513013**
Actinomyces naeslundii		87.60	0.54	MLZ 5			
Propionibacterium granulosum		12.40	0.25				
07513020	**Low discrimination**						**07513020**
					GRA	CAT	
Propionibacterium granulosum		85.01	0.74		99	90	
Actinomyces naeslundii		10.47	0.73		99	25	
Prevotella melaninogenica		2.11	0.32		1	1	
07513021	**Good identification**						**07513021**
					GRA	CAT	
Actinomyces naeslundii		94.70	0.76		99	25	
Propionibacterium granulosum		5.18	0.39		99	90	
Prevotella melaninogenica		0.13	-.03		1	1	
07513022	**Low discrimination**						**07513022**
					CAT		
Propionibacterium granulosum		62.50	0.63		90		
Actinomyces naeslundii		30.80	0.73		25		
Lactobacillus fermentum		3.19	0.46		1		
07513023	**Very good identification**						**07513023**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		98.65	0.76				
Propionibacterium granulosum		1.35	0.29				
07513030	**Very good identification**						**07513030**
					GRA	CAT	
Propionibacterium granulosum		95.29	0.59	MLZ 12	99	90	
Actinomyces naeslundii		4.53	0.51		99	25	
Prevotella melaninogenica		0.18	-.03		1	1	
07513031	**Low discrimination**						**07513031**
Actinomyces naeslundii		87.60	0.54	MLZ 5			
Propionibacterium granulosum		12.40	0.25				
07513032	**Low discrimination**						**07513032**
Propionibacterium granulosum		84.02	0.48	MLZ 12			
Actinomyces naeslundii		15.98	0.51				
07513033	**Good identification**						**07513033**
Actinomyces naeslundii		96.58	0.54	MLZ 5			
Propionibacterium granulosum		3.42	0.14				
07513040	**Low discrimination**						**07513040**
					CAT		
Propionibacterium granulosum		55.41	0.50	SOR 1	90		
Actinomyces naeslundii		29.82	0.60		25		
Propionibacterium propionicum		12.70	0.43		1		
07513041	**Very good identification**						**07513041**
					CAT		
Actinomyces naeslundii		95.78	0.63		25		
Propionibacterium propionicum		3.02	0.27		1		
Propionibacterium granulosum		1.20	0.16		90		
07513042	**Low discrimination**						**07513042**
					CAT		
Actinomyces naeslundii		47.14	0.60		25		
Propionibacterium propionicum		30.12	0.46		1		
Propionibacterium granulosum		21.90	0.39		90		
07513043	**Very good identification**						**07513043**
Actinomyces naeslundii		95.48	0.63				
Propionibacterium propionicum		4.52	0.30				
07513051	**Good identification**						**07513051**
Actinomyces naeslundii		100.0	0.41	MLZ 5			
07513053	**Good identification**						**07513053**
Actinomyces naeslundii		100.0	0.41	MLZ 5			
07513060	**Low discrimination**						**07513060**
					CAT		
Actinomyces naeslundii		62.68	0.60		25		
Propionibacterium granulosum		29.12	0.39		90		
Clostridium perfringens		7.48	0.22		1		
07513061	**Very good identification**						**07513061**
Actinomyces naeslundii		100.0	0.63				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07513062	**Low discrimination**					**07513062**
					CAT	
Actinomyces naeslundii		84.84	0.60		25	
Propionibacterium granulosum		9.85	0.29		90	
Clostridium perfringens		4.76	0.16		1	
07513063	**Very good identification**					**07513063**
Actinomyces naeslundii		100.0	0.63			
07513071	**Good identification**					**07513071**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
07513072	**Low discrimination**					**07513072**
Actinomyces naeslundii		76.87	0.38	MLZ 5		
Propionibacterium granulosum		23.13	0.14			
07513073	**Good identification**					**07513073**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
07513100	**Low discrimination**					**07513100**
Propionibacterium propionicum		65.01	0.78	NIT 86		
Actinomyces naeslundii		24.86	0.81			
Lactobacillus fermentum		4.43	0.58			
07513101	**Low discrimination**					**07513101**
					SPO	
Actinomyces naeslundii		82.97	0.84		1	
Propionibacterium propionicum		16.07	0.61		1	
Clostridium chauvoei		0.76	0.26		99	
07513102	**Low discrimination**					**07513102**
Propionibacterium propionicum		75.13	0.81	NIT 86		
Actinomyces naeslundii		19.16	0.81			
Lactobacillus fermentum		3.41	0.58			
07513103	**Low discrimination**					**07513103**
Actinomyces naeslundii		77.03	0.84			
Propionibacterium propionicum		22.38	0.64			
Actinomyces viscosus		0.38	0.20			
07513110	**Low discrimination**					**07513110**
					CAT	
Actinomyces naeslundii		53.24	0.59	MLZ 5	25	
Propionibacterium propionicum		26.72	0.43		1	
Propionibacterium granulosum		15.89	0.35		90	
07513111	**Very good identification**					**07513111**
Actinomyces naeslundii		96.42	0.62	MLZ 5		
Propionibacterium propionicum		3.58	0.27			
07513112	**Low discrimination**					**07513112**
Actinomyces naeslundii		37.22	0.59	MLZ 5		
Bifidobacterium adolescentis		29.49	0.41			
Propionibacterium propionicum		28.01	0.46			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07513113	**Good identification**						**07513113**
Actinomyces naeslundii		94.24	0.62	MLZ 5			
Propionibacterium propionicum		5.26	0.30				
Bifidobacterium adolescentis		0.50	0.07				
07513120	**Low discrimination**						**07513120**
					GRA		
Prevotella melaninogenica		83.25	0.66	MAN 1	1		
Actinomyces naeslundii		11.86	0.81		99		
Clostridium perfringens		3.38	0.49		99		
07513121	**Good identification**						**07513121**
					GRA	SPO	
Actinomyces naeslundii		94.82	0.84		99	1	
Prevotella melaninogenica		4.48	0.32		1	1	
Clostridium chauvoei		0.29	0.18		99	99	
07513122	**Low discrimination**						**07513122**
					GRA		
Prevotella melaninogenica		48.85	0.54	MAN 1	1		
Actinomyces naeslundii		36.54	0.81		99		
Clostridium perfringens		4.90	0.43		99		
07513123	**Very good identification**						**07513123**
					GRA		
Actinomyces naeslundii		98.56	0.84		99		
Prevotella melaninogenica		0.89	0.19		1		
Propionibacterium propionicum		0.29	0.30		99		
07513131	**Very good identification**						**07513131**
					GRA		
Actinomyces naeslundii		99.10	0.62	MLZ 5	99		
Prevotella melaninogenica		0.90	-.03		1		
07513132	**Low discrimination**						**07513132**
					GRA		
Actinomyces naeslundii		65.54	0.59	MLZ 5	99		
Prevotella melaninogenica		16.81	0.19		1		
Bifidobacterium longum subsp. longum		5.79	0.24		99		
07513133	**Very good identification**						**07513133**
Actinomyces naeslundii		100.0	0.62	MLZ 5			
07513140	**Good identification**						**07513140**
Propionibacterium propionicum		93.47	0.78	NIT 86			
Actinomyces naeslundii		6.31	0.68				
Lactobacillus fermentum		0.06	0.24				
07513141	**Low discrimination**						**07513141**
Propionibacterium propionicum		52.33	0.61	NIT 86			
Actinomyces naeslundii		47.67	0.71	URE 10			
07513142	**Very good identification**						**07513142**
Propionibacterium propionicum		95.50	0.81	NIT 86			
Actinomyces naeslundii		4.30	0.68				
Bifidobacterium adolescentis		0.10	0.24				
07513143	**Low discrimination**						**07513143**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		62.21	0.64	NIT 86		
Actinomyces naeslundii		37.79	0.71	URE 10		
07513150	**Low discrimination**					**07513150**
Propionibacterium propionicum		73.98	0.43	NIT 86		
Actinomyces naeslundii		26.02	0.46	MLZ 1		
07513151	**Low discrimination**					**07513151**
Actinomyces naeslundii		82.60	0.49	MLZ 5		
Propionibacterium propionicum		17.40	0.27			
07513152	**Low discrimination**					**07513152**
Propionibacterium propionicum		73.28	0.46	NIT 86		
Actinomyces naeslundii		17.18	0.46	MLZ 1		
Bifidobacterium adolescentis		9.54	0.26			
07513153	**Low discrimination**					**07513153**
Actinomyces naeslundii		75.99	0.49	MLZ 5		
Propionibacterium propionicum		24.01	0.30			
07513160	**Low discrimination**					**07513160**
					GRA	
Actinomyces naeslundii		53.70	0.68		99	
Prevotella melaninogenica		21.57	0.32		1	
Clostridium perfringens		16.51	0.37		99	
07513161	**Very good identification**					**07513161**
					GRA	
Actinomyces naeslundii		98.44	0.71		99	
Propionibacterium propionicum		1.09	0.27		99	
Prevotella melaninogenica		0.27	-.03		1	
07513162	**Low discrimination**					**07513162**
Actinomyces naeslundii		67.84	0.68			
Propionibacterium propionicum		15.23	0.46			
Clostridium perfringens		9.82	0.31			
07513163	**Very good identification**					**07513163**
Actinomyces naeslundii		98.36	0.71			
Propionibacterium propionicum		1.64	0.30			
07513170	**Low discrimination**					**07513170**
					GRA	
Actinomyces naeslundii		88.02	0.46	MLZ 5	99	
Prevotella melaninogenica		6.79	-.03		1	
Clostridium perfringens		5.19	0.02		99	
07513171	**Good identification**					**07513171**
Actinomyces naeslundii		100.0	0.49	MLZ 5		
07513172	**Good identification**					**07513172**
Actinomyces naeslundii		95.87	0.46	MLZ 5		
Propionibacterium propionicum		4.13	0.12			
07513173	**Good identification**					**07513173**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.49	MLZ 5			
07513200	**Very good identification**						**07513200**
					GRA	CAT	
Propionibacterium granulosum		97.82	0.58	RHA 3	99	90	
Actinomyces naeslundii		0.98	0.39		99	25	
Prevotella intermedia		0.89	0.15		1	1	
07513201	**Low discrimination**						**07513201**
Actinomyces naeslundii		59.90	0.42	RHA 1			
Propionibacterium granulosum		40.10	0.24				
07513202	**Acceptable identification**						**07513202**
					CAT		
Propionibacterium granulosum		94.58	0.48	RHA 3	90		
Actinomyces naeslundii		3.81	0.39		25		
Lactobacillus fermentum		1.18	0.20		1		
07513203	**Low discrimination**						**07513203**
Actinomyces naeslundii		85.66	0.42	RHA 1			
Propionibacterium granulosum		14.34	0.14				
07513210	**Good identification**						**07513210**
Propionibacterium granulosum		100.0	0.43	RHA 3			
07513220	**Good identification**						**07513220**
					GRA	CAT	
Propionibacterium granulosum		95.39	0.48	RHA 3	99	90	
Actinomyces naeslundii		3.84	0.39		99	25	
Prevotella melaninogenica		0.77	-.03		1	1	
07513221	**Low discrimination**						**07513221**
Actinomyces naeslundii		85.66	0.42	RHA 1			
Propionibacterium granulosum		14.34	0.14				
07513222	**Low discrimination**						**07513222**
					GRA	CAT	
Propionibacterium granulosum		76.16	0.37	RHA 3	99	90	
Actinomyces naeslundii		12.26	0.39		99	25	
Prevotella oralis		11.59	0.13		1	1	
07513223	**Good identification**						**07513223**
Actinomyces naeslundii		100.0	0.42	RHA 1			
07513300	**Low discrimination**						**07513300**
					CAT		
Propionibacterium propionicum		62.01	0.43	NIT 86	1		
Actinomyces naeslundii		23.72	0.46	RHA 1	25		
Propionibacterium granulosum		8.37	0.24		90		
07513301	**Low discrimination**						**07513301**
Actinomyces naeslundii		83.77	0.49	RHA 1			
Propionibacterium propionicum		16.23	0.27				
07513302	**Low discrimination**						**07513302**
					GRA		
Propionibacterium propionicum		71.80	0.46	NIT 86	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		18.31	0.46	RHA 1	99	
Prevotella oralis		5.01	0.11		1	
07513303	**Low discrimination**					**07513303**
Actinomyces naeslundii		77.49	0.49	RHA 1		
Propionibacterium propionicum		22.51	0.30			
07513321	**Good identification**					**07513321**
					GRA	
Actinomyces naeslundii		95.49	0.49	RHA 1	99	
Prevotella melaninogenica		4.51	-.03		1	
07513323	**Good identification**					**07513323**
					GRA	
Actinomyces naeslundii		96.62	0.49	RHA 1	99	
Prevotella oralis		3.39	-.01		1	
07513340	**Acceptable identification**					**07513340**
Propionibacterium propionicum		93.68	0.43	NIT 86		
Actinomyces naeslundii		6.32	0.34	RHA 1		
07513342	**Good identification**					**07513342**
Propionibacterium propionicum		95.69	0.46	NIT 86		
Actinomyces naeslundii		4.31	0.34	RHA 1		
07513361	**Good identification**					**07513361**
Actinomyces naeslundii		100.0	0.37	RHA 1		
07513363	**Good identification**					**07513363**
Actinomyces naeslundii		100.0	0.37	RHA 1		
07513400	**Low discrimination**					**07513400**
					CAT	
Propionibacterium granulosum		49.24	0.50	ARA 1	90	
Lactobacillus fermentum		46.68	0.54		1	
Actinomyces naeslundii		1.52	0.39		25	
07513401	**Low discrimination**					**07513401**
					CAT	
Actinomyces naeslundii		70.13	0.42	ARA 1	25	
Propionibacterium granulosum		15.33	0.16		90	
Lactobacillus fermentum		14.54	0.20		1	
07513402	**Low discrimination**					**07513402**
					CAT	
Lactobacillus fermentum		74.94	0.54	MAN 1	1	
Propionibacterium granulosum		19.76	0.39		90	
Actinomyces naeslundii		2.44	0.39		25	
07513403	**Low discrimination**					**07513403**
Actinomyces naeslundii		82.83	0.42	ARA 1		
Lactobacillus fermentum		17.17	0.20			
07513420	**Low discrimination**					**07513420**
					CAT	
Lactobacillus fermentum		50.42	0.46	MAN 1	1	
Propionibacterium granulosum		39.89	0.39		90	
Actinomyces naeslundii		4.92	0.39		25	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07513421	**Good identification**					**07513421**
Actinomyces naeslundii		100.0	0.42	ARA 1		
07513422	**Low discrimination**					**07513422**
					CAT	
Lactobacillus fermentum		72.98	0.46	MAN 1	1	
Propionibacterium granulosum		14.43	0.29		90	
Actinomyces naeslundii		7.11	0.39		25	
07513423	**Good identification**					**07513423**
Actinomyces naeslundii		100.0	0.42	ARA 1		
07513500	**Low discrimination**					**07513500**
Lactobacillus fermentum		61.75	0.58	MAN 1		
Bifidobacterium longum subsp. longum		23.43	0.43			
Propionibacterium propionicum		9.15	0.43			
07513501	**Low discrimination**					**07513501**
Actinomyces naeslundii		73.66	0.49	ARA 1		
Propionibacterium propionicum		14.27	0.27			
Lactobacillus fermentum		8.75	0.24			
07513502	**Low discrimination**					**07513502**
Bifidobacterium adolescentis		43.29	0.54	XYL 87		
Lactobacillus fermentum		33.78	0.58			
Bifidobacterium longum subsp. longum		12.81	0.43			
07513503	**Low discrimination**					**07513503**
Actinomyces naeslundii		62.24	0.49	ARA 1		
Propionibacterium propionicum		18.08	0.30			
Bifidobacterium adolescentis		9.48	0.20			
07513510	**Good identification**					**07513510**
Bifidobacterium longum subsp. longum		95.72	0.47	MAN 1		
Bifidobacterium adolescentis		2.34	0.21			
Lactobacillus fermentum		1.50	0.24			
07513512	**Low discrimination**					**07513512**
Bifidobacterium adolescentis		70.23	0.56	XYL 87		
Bifidobacterium longum subsp. longum		28.97	0.47			
Lactobacillus fermentum		0.45	0.24			
07513520	**Low discrimination**					**07513520**
					GRA	
Bifidobacterium longum subsp. longum		63.88	0.54	MAN 1	99	
Prevotella melaninogenica		16.74	0.32		1	
Lactobacillus fermentum		14.03	0.50		99	
07513521	**Low discrimination**					**07513521**
					GRA	
Actinomyces naeslundii		78.92	0.49	ARA 1	99	
Bifidobacterium longum subsp. longum		14.23	0.20		99	
Prevotella melaninogenica		3.73	-.03		1	
07513522	**Low discrimination**					**07513522**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Bifidobacterium longum subsp. longum		74.03	0.54	MAN 1	99			
Lactobacillus fermentum		16.26	0.50		99			
Prevotella melaninogenica		3.70	0.19		1			
07513523	**Low discrimination**							**07513523**
Actinomyces naeslundii		81.97	0.49	ARA 1				
Bifidobacterium longum subsp. longum		14.78	0.20					
Lactobacillus fermentum		3.25	0.15					
07513530	**Very good identification**							**07513530**
					GRA			
Bifidobacterium longum subsp. longum		99.60	0.58	MAN 1	99			
Prevotella melaninogenica		0.16	-.03		1			
Lactobacillus fermentum		0.13	0.15		99			
07513532	**Very good identification**							**07513532**
					GRA			
Bifidobacterium longum subsp. longum		98.85	0.58	MAN 1	99			
Bifidobacterium adolescentis		0.61	0.21		99			
Parabacteroides distasonis		0.18	0.15		1			
07513540	**Low discrimination**							**07513540**
Propionibacterium propionicum		83.87	0.43	NIT 86				
Lactobacillus fermentum		5.72	0.24	ARA 1				
Actinomyces naeslundii		5.66	0.34					
07513542	**Low discrimination**							**07513542**
Propionibacterium propionicum		54.02	0.46	NIT 86				
Bifidobacterium adolescentis		38.49	0.39	ARA 1				
Lactobacillus fermentum		2.46	0.24					
07513552	**Good identification**							**07513552**
Bifidobacterium adolescentis		95.67	0.40	XYL 87				
Bifidobacterium longum subsp. longum		3.22	0.13	SOR 11				
Propionibacterium propionicum		1.11	0.12					
07513561	**Good identification**							**07513561**
Actinomyces naeslundii		100.0	0.37	ARA 1				
07513563	**Good identification**							**07513563**
Actinomyces naeslundii		100.0	0.37	ARA 1				
07514000	**Low discrimination**							**07514000**
					COC	GRA	CAT	
Prevotella intermedia		35.76	0.49	XYL 1	1	1	1	
Gemella morbillorum		32.61	0.50		99	99	1	
Propionibacterium granulosum		17.16	0.52		1	99	90	
07514002	**Low discrimination**							**07514002**
					CAT			
Lactobacillus fermentum		47.41	0.50	GAL 99	1			
Propionibacterium granulosum		28.22	0.41		90			
Actinomyces viscosus		14.96	0.27		76			
07514010	**Low discrimination**							**07514010**
					COC	GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		73.80	0.37	XYL 1	1	99	90	
Prevotella intermedia		11.39	0.15	MLZ 12	1	1	1	
Gemella morbillorum		10.39	0.15		99	99	1	
07514020	**Low discrimination**							**07514020**
					COC	CAT		
Propionibacterium granulosum		39.42	0.41	XYL 1	1	90		
Gemella morbillorum		33.29	0.33		99	1		
Lactobacillus fermentum		22.08	0.42		1	1		
07514022	**Low discrimination**							**07514022**
					GRA	CAT		
Lactobacillus fermentum		64.05	0.42	GAL 99	99	1		
Propionibacterium granulosum		28.59	0.31		99	90		
Bacteroides fragilis		4.47	-.03		1	99		
07514100	**Low discrimination**							**07514100**
					GRA			
Fusobacterium mortiferum		51.48	0.54	MLT 11	1			
Lactobacillus fermentum		38.90	0.54	XYL 7	99			
Actinomyces meyeri		5.73	0.29		99			
07514102	**Low discrimination**							**07514102**
					GRA			
Lactobacillus fermentum		63.49	0.54	GAL 99	99			
Actinomyces viscosus		21.83	0.32		99			
Fusobacterium mortiferum		9.34	0.38		1			
07514320	**Good identification**							**07514320**
Bacteroides vulgatus		99.54	0.40	GAL 99				
Prevotella melaninogenica		0.24	-.03	ARA 93				
Parabacteroides distasonis		0.23	0.14					
07514322	**Low discrimination**							**07514322**
Parabacteroides distasonis		84.79	0.48	GAL 99				
Prevotella oralis		8.01	0.17					
Bacteroides vulgatus		3.77	0.06					
07514332	**Low discrimination**							**07514332**
Parabacteroides distasonis		86.21	0.43	GAL 99				
Clostridium clostridioforme		13.79	0.24					
07514400	**Low discrimination**							**07514400**
					COC	GRA		
Lactobacillus fermentum		61.23	0.50	GAL 99	1	99		
Gemella morbillorum		30.78	0.33		99	99		
Prevotella intermedia		3.07	0.15		1	1		
07514402	**Good identification**							**07514402**
Lactobacillus fermentum		95.71	0.50	GAL 99				
Bifidobacterium longum subsp. longum		3.06	0.17					
Clostridium clostridioforme		1.24	0.09					
07514420	**Low discrimination**							**07514420**
					GRA			
Lactobacillus fermentum		56.76	0.42	GAL 99	99			
Bifidobacterium longum subsp. longum		21.75	0.28		99			
Bacteroides vulgatus		11.98	0.09		1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07514422	**Low discrimination**					**07514422**
Lactobacillus fermentum		70.32	0.42	GAL 99		
Bifidobacterium longum subsp. longum		26.95	0.28			
Clostridium clostridioforme		2.73	0.09			
07514500	**Low discrimination**					**07514500**
					GRA	
Bifidobacterium longum subsp. longum		63.93	0.52	GAL 99	99	
Lactobacillus fermentum		32.99	0.54		99	
Bacteroides vulgatus		1.42	0.09		1	
07514502	**Low discrimination**					**07514502**
Bifidobacterium longum subsp. longum		61.63	0.52	GAL 99		
Lactobacillus fermentum		31.80	0.54			
Bifidobacterium adolescentis		5.73	0.36			
07514510	**Good identification**					**07514510**
Bifidobacterium longum subsp. longum		99.70	0.56	GAL 99		
Lactobacillus fermentum		0.31	0.20			
07514512	**Acceptable identification**					**07514512**
Bifidobacterium longum subsp. longum		90.38	0.56	GAL 99		
Bifidobacterium adolescentis		6.04	0.37			
Clostridium clostridioforme		3.31	0.29			
07514520	**Low discrimination**					**07514520**
					GRA	
Bifidobacterium longum subsp. longum		62.68	0.62	GAL 99	99	
Bacteroides vulgatus		34.53	0.43		1	
Lactobacillus fermentum		2.70	0.46		99	
07514522	**Good identification**					**07514522**
					GRA	
Bifidobacterium longum subsp. longum		93.22	0.62	GAL 99	99	
Lactobacillus fermentum		4.01	0.46		99	
Parabacteroides distasonis		1.62	0.36		1	
07514530	**Very good identification**					**07514530**
					GRA	
Bifidobacterium longum subsp. longum		99.67	0.66	GAL 99	99	
Bacteroides vulgatus		0.33	0.09		1	
07514532	**Very good identification**					**07514532**
					GRA	
Bifidobacterium longum subsp. longum		98.56	0.66	GAL 99	99	
Clostridium clostridioforme		0.90	0.29		50	
Parabacteroides distasonis		0.54	0.31		1	
07514612	**Good identification**					**07514612**
Clostridium clostridioforme		100.0	0.35	SAL 88		
07514632	**Good identification**					**07514632**
Clostridium clostridioforme		100.0	0.35	SAL 88		
07514712	**Good identification**					**07514712**
Clostridium clostridioforme		96.38	0.44	SAL 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		3.62	0.21	TRE 88			
07514720	**Very good identification**						**07514720**
					GRA		
Bacteroides vulgatus		99.79	0.60	GAL 99	1		
Bifidobacterium longum subsp. longum		0.20	0.28		99		
Parabacteroides distasonis		0.01	0.10		1		
07514722	**Low discrimination**						**07514722**
Parabacteroides distasonis		37.86	0.44	GAL 99			
Bacteroides vulgatus		36.48	0.25				
Clostridium clostridioforme		18.30	0.33				
07514732	**Low discrimination**						**07514732**
					GRA		
Clostridium clostridioforme		71.46	0.44	SAL 88	50		
Parabacteroides distasonis		17.48	0.39	TRE 88	1		
Bifidobacterium longum subsp. longum		10.74	0.32		99		
07515000	**Low discrimination**						**07515000**
					COC	GRA	
Lactobacillus fermentum		89.70	0.84		1	99	
Prevotella intermedia		4.50	0.49		1	1	
Gemella morbillorum		2.21	0.45		99	99	
07515001	**Low discrimination**						**07515001**
					GRA		
Lactobacillus fermentum		86.54	0.50	URE 1	99		
Prevotella intermedia		4.34	0.15		1		
Actinomyces naeslundii		3.70	0.37		99		
07515002	**Very good identification**						**07515002**
					CAT		
Lactobacillus fermentum		98.73	0.84		1		
Propionibacterium granulosum		0.59	0.41		90		
Actinomyces viscosus		0.19	0.24		76		
07515003	**Acceptable identification**						**07515003**
Lactobacillus fermentum		90.32	0.50	URE 1			
Actinomyces viscosus		5.82	0.16				
Actinomyces naeslundii		3.86	0.37				
07515010	**Low discrimination**						**07515010**
					CAT		
Lactobacillus fermentum		68.13	0.50	MLZ 1	1		
Propionibacterium granulosum		22.12	0.37		90		
Bifidobacterium longum subsp. longum		3.71	0.21		1		
07515012	**Low discrimination**						**07515012**
					CAT		
Lactobacillus fermentum		83.30	0.50	MLZ 1	1		
Propionibacterium granulosum		6.76	0.27		90		
Bifidobacterium longum subsp. longum		4.53	0.21		1		
07515020	**Good identification**						**07515020**
					GRA		
Lactobacillus fermentum		90.30	0.76		99		
Prevotella melaninogenica		3.02	0.32		1		
Clostridium perfringens		1.66	0.34		99		
07515021	**Low discrimination**						**07515021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Lactobacillus fermentum		86.08	0.42	URE 1	99		
Actinomyces naeslundii		11.04	0.37		99		
Prevotella melaninogenica		2.88	-.03		1		
07515022	**Low discrimination**						**07515022**
					CAT	GRA	
Lactobacillus fermentum		86.68	0.76		1	99	
Bacteroides fragilis		6.04	0.32		99	1	
Prevotella oralis		3.92	0.37		1	1	
07515023	**Low discrimination**						**07515023**
					GRA	CAT	
Lactobacillus fermentum		80.44	0.42	URE 1	99	1	
Actinomyces naeslundii		10.32	0.37		99	25	
Bacteroides fragilis		5.61	-.03		1	99	
07515030	**Low discrimination**						**07515030**
					CAT		
Lactobacillus fermentum		51.83	0.42	MLZ 1	1		
Bifidobacterium longum subsp. longum		33.82	0.32		1		
Propionibacterium granulosum		12.62	0.27		90		
07515032	**Low discrimination**						**07515032**
					GRA		
Lactobacillus fermentum		40.38	0.42	MLZ 1	99		
Bifidobacterium longum subsp. longum		26.35	0.32		99		
Parabacteroides distasonis		23.32	0.35		1		
07515040	**Low discrimination**						**07515040**
					COC	GRA	
Lactobacillus fermentum		89.23	0.50	SOR 1	1	99	
Prevotella intermedia		4.47	0.15		1	1	
Gemella morbillorum		2.20	0.11		99	99	
07515042	**Good identification**						**07515042**
Lactobacillus fermentum		98.83	0.50	SOR 1			
Actinomyces israelii		1.17	0.18				
07515060	**Low discrimination**						**07515060**
					GRA		
Lactobacillus fermentum		72.49	0.42	SOR 1	99		
Clostridium perfringens		25.08	0.22		99		
Prevotella melaninogenica		2.43	-.03		1		
07515062	**Low discrimination**						**07515062**
					CAT	GRA	
Lactobacillus fermentum		76.07	0.42	SOR 1	1	99	
Clostridium perfringens		12.38	0.16		1	99	
Bacteroides fragilis		5.30	-.03		99	1	
07515100	**Good identification**						**07515100**
					GRA		
Lactobacillus fermentum		94.57	0.88		99		
Fusobacterium mortiferum		1.90	0.57		1		
Bifidobacterium longum subsp. longum		1.85	0.52		99		
07515101	**Low discrimination**						**07515101**
					SPO		
Lactobacillus fermentum		79.83	0.54	URE 1	1		
Clostridium chauvoei		10.51	0.26		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		5.95	0.45		1		
07515102	**Good identification**						**07515102**
Lactobacillus fermentum		91.85	0.88				
Bifidobacterium adolescentis		2.47	0.56				
Bifidobacterium longum subsp. longum		1.80	0.52				
07515103	**Low discrimination**						**07515103**
Lactobacillus fermentum		80.39	0.54	URE 1			
Actinomyces naeslundii		6.00	0.45				
Actinomyces viscosus		5.65	0.20				
07515110	**Low discrimination**						**07515110**
Bifidobacterium longum subsp. longum		75.32	0.56	ARA 99			
Lactobacillus fermentum		22.83	0.54				
Bifidobacterium adolescentis		0.75	0.23				
07515112	**Low discrimination**						**07515112**
Bifidobacterium longum subsp. longum		42.22	0.56	ARA 99			
Bifidobacterium adolescentis		41.70	0.57				
Lactobacillus fermentum		12.79	0.54				
07515120	**Low discrimination**						**07515120**
					GRA		
Prevotella melaninogenica		44.24	0.66	XYL 1	1		
Lactobacillus fermentum		21.77	0.80		99		
Bacteroides vulgatus		20.99	0.58		1		
07515122	**Low discrimination**						**07515122**
					GRA	CAT	
Bacteroides fragilis		58.63	0.66	ESL 99	1	99	
Lactobacillus fermentum		13.86	0.80		99	1	
Parabacteroides distasonis		9.11	0.74		1	76	
07515130	**Low discrimination**						**07515130**
					GRA		
Bifidobacterium longum subsp. longum		88.94	0.66	ARA 99	99		
Prevotella melaninogenica		4.56	0.32		1		
Lactobacillus fermentum		2.25	0.46		99		
07515132	**Low discrimination**						**07515132**
					GRA	CAT	
Bifidobacterium longum subsp. longum		47.62	0.66	ARA 99	99	1	
Parabacteroides distasonis		42.14	0.69		1	76	
Bacteroides fragilis		5.09	0.32		1	99	
07515140	**Acceptable identification**						**07515140**
					GRA		
Lactobacillus fermentum		90.92	0.54	SOR 1	99		
Fusobacterium mortiferum		1.82	0.23		1		
Bifidobacterium longum subsp. longum		1.78	0.17		99		
07515142	**Low discrimination**						**07515142**
					GRA	CAT	
Lactobacillus fermentum		66.53	0.54	SOR 1	99	1	
Bifidobacterium adolescentis		21.93	0.40		99	1	
Capnocytophaga ochracea		5.51	0.20		1	90	
07515152	**Low discrimination**						**07515152**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		87.61	0.42	ARA 87			
Bifidobacterium longum subsp. longum		7.25	0.21	SOR 11			
Lactobacillus fermentum		2.20	0.20				
07515200	**Low discrimination**						**07515200**
					GRA	CAT	
Lactobacillus fermentum		81.88	0.50	RHA 1	99	1	
Propionibacterium granulosum		6.03	0.26		99	90	
Prevotella intermedia		4.10	0.15		1	1	
07515202	**Acceptable identification**						**07515202**
					GRA	CAT	
Lactobacillus fermentum		91.91	0.50	RHA 1	99	1	
Prevotella oralis		3.10	0.09		1	1	
Propionibacterium granulosum		1.69	0.16		99	90	
07515220	**Acceptable identification**						**07515220**
					GRA		
Bacteroides vulgatus		92.05	0.40	LAC 99	1		
Lactobacillus fermentum		6.50	0.42	ARA 93	99		
Prevotella oralis		0.66	0.09		1		
07515222	**Low discrimination**						**07515222**
					GRA		
Parabacteroides distasonis		50.24	0.48	LAC 99	1		
Prevotella oralis		29.97	0.31		1		
Lactobacillus fermentum		15.61	0.42		99		
07515232	**Low discrimination**						**07515232**
Parabacteroides distasonis		86.62	0.43	LAC 99			
Clostridium clostridioforme		12.42	0.23				
Prevotella oralis		0.97	-.03				
07515300	**Low discrimination**						**07515300**
					GRA		
Bacteroides vulgatus		70.78	0.40	ARA 93	1		
Lactobacillus fermentum		24.47	0.54	NAG 99	99		
Eubacterium contortum		3.04	0.24		99		
07515302	**Low discrimination**						**07515302**
					GRA		
Lactobacillus fermentum		46.66	0.54	RHA 1	99		
Parabacteroides distasonis		30.67	0.48		1		
Prevotella oralis		15.09	0.30		1		
07515312	**Low discrimination**						**07515312**
					GRA		
Parabacteroides distasonis		58.98	0.43	NAG 99	1		
Clostridium clostridioforme		28.30	0.32		50		
Bifidobacterium longum subsp. longum		5.55	0.21		99		
07515320	**Very good identification**						**07515320**
Bacteroides vulgatus		99.24	0.75	ARA 93			
Prevotella melaninogenica		0.24	0.32				
Parabacteroides distasonis		0.23	0.48				
07515321	**Good identification**						**07515321**
Bacteroides vulgatus		99.54	0.40	ARA 93			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		0.24	-.03	URE 1		
Parabacteroides distasonis		0.23	0.14			
07515322	**Low discrimination**					**07515322**
Parabacteroides distasonis		85.41	0.82			
Prevotella oralis		8.06	0.52			
Bacteroides vulgatus		3.80	0.40			
07515323	**Low discrimination**					**07515323**
Parabacteroides distasonis		86.17	0.48	URE 1		
Prevotella oralis		8.14	0.17			
Bacteroides vulgatus		3.83	0.06			
07515330	**Low discrimination**					**07515330**
					GRA	
Bacteroides vulgatus		85.41	0.40	ARA 93	1	
Parabacteroides distasonis		10.45	0.43	MLZ 1	1	
Bifidobacterium longum subsp. longum		3.94	0.32		99	
07515332	**Very good identification**					**07515332**
					GRA	
Parabacteroides distasonis		98.83	0.77		1	
Clostridium clostridioforme		0.48	0.32		50	
Bifidobacterium longum subsp. longum		0.38	0.32		99	
07515333	**Good identification**					**07515333**
Parabacteroides distasonis		100.0	0.43	URE 1		
07515360	**Good identification**					**07515360**
Bacteroides vulgatus		99.36	0.40	ARA 93		
Prevotella melaninogenica		0.24	-.03	SOR 1		
Parabacteroides distasonis		0.23	0.14			
07515362	**Low discrimination**					**07515362**
Parabacteroides distasonis		85.33	0.48	SOR 1		
Prevotella oralis		8.06	0.17			
Bacteroides vulgatus		3.79	0.06			
07515372	**Good identification**					**07515372**
Parabacteroides distasonis		100.0	0.43	SOR 1		
07515400	**Very good identification**					**07515400**
					COC	
Lactobacillus fermentum		96.46	0.84		1	
Bifidobacterium longum subsp. longum		3.08	0.52		1	
Gemella morbillorum		0.26	0.29		99	
07515401	**Good identification**					**07515401**
Lactobacillus fermentum		96.91	0.50	URE 1		
Bifidobacterium longum subsp. longum		3.10	0.17			
07515402	**Very good identification**					**07515402**
Lactobacillus fermentum		96.57	0.84			
Bifidobacterium longum subsp. longum		3.08	0.52			
Bifidobacterium adolescentis		0.29	0.36			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07515403	**Good identification**					**07515403**
Lactobacillus fermentum		96.91	0.50	URE 1		
Bifidobacterium longum subsp. longum		3.10	0.17			
07515410	**Low discrimination**					**07515410**
Bifidobacterium longum subsp. longum		84.34	0.56	LAC 99		
Lactobacillus fermentum		15.66	0.50			
07515412	**Low discrimination**					**07515412**
Bifidobacterium longum subsp. longum		77.80	0.56	LAC 99		
Lactobacillus fermentum		14.45	0.50			
Bifidobacterium adolescentis		5.20	0.37			
07515420	**Low discrimination**					**07515420**
					GRA	
Lactobacillus fermentum		62.67	0.76		99	
Bifidobacterium longum subsp. longum		24.02	0.62		99	
Bacteroides vulgatus		13.23	0.43		1	
07515421	**Low discrimination**					**07515421**
					GRA	
Lactobacillus fermentum		62.72	0.42	URE 1	99	
Bifidobacterium longum subsp. longum		24.04	0.28		99	
Bacteroides vulgatus		13.24	0.09		1	
07515422	**Low discrimination**					**07515422**
					GRA	
Lactobacillus fermentum		71.69	0.76		99	
Bifidobacterium longum subsp. longum		27.48	0.62		99	
Parabacteroides distasonis		0.48	0.36		1	
07515423	**Low discrimination**					**07515423**
Lactobacillus fermentum		72.29	0.42	URE 1		
Bifidobacterium longum subsp. longum		27.71	0.28			
07515430	**Very good identification**					**07515430**
					GRA	
Bifidobacterium longum subsp. longum		98.16	0.66	LAC 99	99	
Lactobacillus fermentum		1.52	0.42		99	
Bacteroides vulgatus		0.32	0.09		1	
07515432	**Very good identification**					**07515432**
Bifidobacterium longum subsp. longum		97.17	0.66	LAC 99		
Lactobacillus fermentum		1.50	0.42			
Clostridium clostridioforme		0.80	0.29			
07515440	**Good identification**					**07515440**
Lactobacillus fermentum		96.30	0.50	SOR 1		
Bifidobacterium longum subsp. longum		3.08	0.17			
Actinomyces israelii		0.62	0.13			
07515442	**Acceptable identification**					**07515442**
Lactobacillus fermentum		92.77	0.50	SOR 1		
Bifidobacterium adolescentis		3.37	0.20			
Bifidobacterium longum subsp. longum		2.96	0.17			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07515460	**Low discrimination**					**07515460**
					GRA	
Lactobacillus fermentum		62.72	0.42	SOR 1	99	
Bifidobacterium longum subsp. longum		24.04	0.28		99	
Bacteroides vulgatus		13.24	0.09		1	
07515462	**Low discrimination**					**07515462**
Lactobacillus fermentum		72.29	0.42	SOR 1		
Bifidobacterium longum subsp. longum		27.71	0.28			
07515500	**Low discrimination**					**07515500**
					GRA	
Bifidobacterium longum subsp. longum		64.94	0.86		99	
Lactobacillus fermentum		33.51	0.88		99	
Bacteroides vulgatus		1.45	0.43		1	
07515501	**Low discrimination**					**07515501**
					GRA	
Bifidobacterium longum subsp. longum		65.01	0.52	URE 1	99	
Lactobacillus fermentum		33.54	0.54		99	
Bacteroides vulgatus		1.45	0.09		1	
07515502	**Low discrimination**					**07515502**
Bifidobacterium longum subsp. longum		62.06	0.86			
Lactobacillus fermentum		32.02	0.88			
Bifidobacterium adolescentis		5.77	0.70			
07515503	**Low discrimination**					**07515503**
Bifidobacterium longum subsp. longum		62.15	0.52	URE 1		
Lactobacillus fermentum		32.07	0.54			
Bifidobacterium adolescentis		5.78	0.36			
07515510	**Very good identification**					**07515510**
Bifidobacterium longum subsp. longum		99.61	0.90			
Lactobacillus fermentum		0.31	0.54			
Bifidobacterium adolescentis		0.07	0.37			
07515511	**Good identification**					**07515511**
Bifidobacterium longum subsp. longum		99.70	0.56	URE 1		
Lactobacillus fermentum		0.31	0.20			
07515512	**Good identification**					**07515512**
Bifidobacterium longum subsp. longum		93.35	0.90			
Bifidobacterium adolescentis		6.23	0.71			
Lactobacillus fermentum		0.29	0.54			
07515513	**Acceptable identification**					**07515513**
Bifidobacterium longum subsp. longum		93.47	0.56	URE 1		
Bifidobacterium adolescentis		6.24	0.37			
Lactobacillus fermentum		0.29	0.20			
07515520	**Low discrimination**					**07515520**
					GRA	
Bifidobacterium longum subsp. longum		62.69	0.96		99	
Bacteroides vulgatus		34.53	0.77		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		2.70	0.80		99	
07515521	**Low discrimination**					**07515521**
					GRA	
Bifidobacterium longum subsp. longum		62.71	0.62	URE 1	99	
Bacteroides vulgatus		34.54	0.43		1	
Lactobacillus fermentum		2.70	0.46		99	
07515522	**Good identification**					**07515522**
					GRA	
Bifidobacterium longum subsp. longum		93.46	0.96		99	
Lactobacillus fermentum		4.02	0.80		99	
Parabacteroides distasonis		1.62	0.70		1	
07515523	**Good identification**					**07515523**
					GRA	
Bifidobacterium longum subsp. longum		93.52	0.62	URE 1	99	
Lactobacillus fermentum		4.02	0.46		99	
Parabacteroides distasonis		1.62	0.36		1	
07515530	**Excellent identification**					**07515530**
					GRA	
Bifidobacterium longum subsp. longum		99.64	1.00		99	
Bacteroides vulgatus		0.33	0.43		1	
Lactobacillus fermentum		0.03	0.46		99	
07515531	**Very good identification**					**07515531**
					GRA	
Bifidobacterium longum subsp. longum		99.67	0.66	URE 1	99	
Bacteroides vulgatus		0.33	0.09		1	
07515532	**Excellent identification**					**07515532**
					GRA	
Bifidobacterium longum subsp. longum		99.37	1.00		99	
Parabacteroides distasonis		0.54	0.65		1	
Clostridium clostridioforme		0.03	0.38		50	
07515533	**Very good identification**					**07515533**
					GRA	
Bifidobacterium longum subsp. longum		99.46	0.66	URE 1	99	
Parabacteroides distasonis		0.55	0.31		1	
07515540	**Low discrimination**					**07515540**
					GRA	
Bifidobacterium longum subsp. longum		64.19	0.52	SOR 1	99	
Lactobacillus fermentum		33.12	0.54		99	
Bacteroides vulgatus		1.43	0.09		1	
07515542	**Low discrimination**					**07515542**
Bifidobacterium adolescentis		42.57	0.54	SOR 11		
Bifidobacterium longum subsp. longum		37.40	0.52			
Lactobacillus fermentum		19.30	0.54			
07515550	**Good identification**					**07515550**
Bifidobacterium longum subsp. longum		98.88	0.56	SOR 1		
Bifidobacterium adolescentis		0.82	0.22			
Lactobacillus fermentum		0.30	0.20			
07515552	**Low discrimination**					**07515552**
Bifidobacterium longum subsp. longum		54.94	0.56	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		44.89	0.56			
Lactobacillus fermentum		0.17	0.20			
07515560	**Low discrimination**					**07515560**
					GRA	
Bifidobacterium longum subsp. longum		62.61	0.62	SOR 1	99	
Bacteroides vulgatus		34.49	0.43		1	
Lactobacillus fermentum		2.69	0.46		99	
07515562	**Good identification**					**07515562**
					GRA	
Bifidobacterium longum subsp. longum		91.95	0.62	SOR 1	99	
Lactobacillus fermentum		3.95	0.46		99	
Parabacteroides distasonis		1.59	0.36		1	
07515570	**Very good identification**					**07515570**
					GRA	
Bifidobacterium longum subsp. longum		99.67	0.66	SOR 1	99	
Bacteroides vulgatus		0.33	0.09		1	
07515572	**Very good identification**					**07515572**
					GRA	
Bifidobacterium longum subsp. longum		99.14	0.66	SOR 1	99	
Parabacteroides distasonis		0.54	0.31		1	
Bifidobacterium adolescentis		0.21	0.22		99	
07515600	**Low discrimination**					**07515600**
					GRA	
Lactobacillus fermentum		60.05	0.50	RHA 1	99	
Bacteroides vulgatus		38.03	0.25		1	
Bifidobacterium longum subsp. longum		1.92	0.17		99	
07515602	**Low discrimination**					**07515602**
Lactobacillus fermentum		75.21	0.50	RHA 1		
Clostridium clostridioforme		21.41	0.32			
Bifidobacterium longum subsp. longum		2.40	0.17			
07515612	**Good identification**					**07515612**
Clostridium clostridioforme		95.26	0.43	SAL 88		
Bifidobacterium longum subsp. longum		4.00	0.21	TRE 88		
Lactobacillus fermentum		0.74	0.16			
07515620	**Very good identification**					**07515620**
					GRA	
Bacteroides vulgatus		99.26	0.60	LAC 99	1	
Lactobacillus fermentum		0.53	0.42		99	
Bifidobacterium longum subsp. longum		0.20	0.28		99	
07515622	**Low discrimination**					**07515622**
					GRA	
Parabacteroides distasonis		32.07	0.44	LAC 99	1	
Bacteroides vulgatus		30.90	0.25		1	
Lactobacillus fermentum		16.26	0.42		99	
07515632	**Low discrimination**					**07515632**
					GRA	
Clostridium clostridioforme		69.17	0.43	SAL 88	50	
Parabacteroides distasonis		18.88	0.39	TRE 88	1	
Bifidobacterium longum subsp. longum		11.60	0.32		99	
07515700	**Good identification**					**07515700**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bacteroides vulgatus		92.65	0.60	NAG 99	1	
Bifidobacterium longum subsp. longum		4.67	0.52		99	
Lactobacillus fermentum		2.41	0.54		99	
07515702	**Low discrimination**					**07515702**
Bifidobacterium longum subsp. longum		42.91	0.52	RHA 1		
Lactobacillus fermentum		22.14	0.54			
Clostridium clostridioforme		12.93	0.41			
07515710	**Low discrimination**					**07515710**
					GRA	
Bifidobacterium longum subsp. longum		88.59	0.56	RHA 1	99	
Bacteroides vulgatus		10.42	0.25		1	
Clostridium clostridioforme		0.72	0.18		50	
07515712	**Low discrimination**					**07515712**
Bifidobacterium longum subsp. longum		51.46	0.56	RHA 1		
Clostridium clostridioforme		41.50	0.52			
Bifidobacterium adolescentis		3.44	0.37			
07515720	**Excellent identification**					**07515720**
					GRA	
Bacteroides vulgatus		99.78	0.94		1	
Bifidobacterium longum subsp. longum		0.20	0.62		99	
Lactobacillus fermentum		0.01	0.46		99	
07515721	**Very good identification**					**07515721**
					GRA	
Bacteroides vulgatus		99.79	0.60	URE 1	1	
Bifidobacterium longum subsp. longum		0.20	0.28		99	
Parabacteroides distasonis		0.01	0.10		1	
07515722	**Low discrimination**					**07515722**
					GRA	
Parabacteroides distasonis		45.70	0.78		1	
Bacteroides vulgatus		44.04	0.60		1	
Bifidobacterium longum subsp. longum		8.88	0.62		99	
07515723	**Low discrimination**					**07515723**
					GRA	
Parabacteroides distasonis		46.03	0.44	URE 1	1	
Bacteroides vulgatus		44.35	0.25		1	
Bifidobacterium longum subsp. longum		8.95	0.28		99	
07515730	**Low discrimination**					**07515730**
					GRA	
Bacteroides vulgatus		74.09	0.60	MLZ 1	1	
Bifidobacterium longum subsp. longum		25.44	0.66		99	
Parabacteroides distasonis		0.42	0.39		1	
07515732	**Low discrimination**					**07515732**
					GRA	
Parabacteroides distasonis		56.81	0.74		1	
Bifidobacterium longum subsp. longum		34.92	0.66		99	
Clostridium clostridioforme		7.04	0.52		50	
07515733	**Low discrimination**					**07515733**
					GRA	
Parabacteroides distasonis		56.93	0.39	URE 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		34.99	0.32		99			
Clostridium clostridioforme		7.05	0.18		50			
07515760	**Very good identification**							**07515760**
					GRA			
Bacteroides vulgatus		99.79	0.60	SOR 1	1			
Bifidobacterium longum subsp. longum		0.20	0.28		99			
Parabacteroides distasonis		0.01	0.10		1			
07515762	**Low discrimination**							**07515762**
					GRA			
Parabacteroides distasonis		46.03	0.44	SOR 1	1			
Bacteroides vulgatus		44.35	0.25		1			
Bifidobacterium longum subsp. longum		8.95	0.28		99			
07515772	**Low discrimination**							**07515772**
					GRA			
Parabacteroides distasonis		56.93	0.39	SOR 1	1			
Bifidobacterium longum subsp. longum		34.99	0.32		99			
Clostridium clostridioforme		7.05	0.18		50			
07516000	**Acceptable identification**							**07516000**
					COC	GRA	CAT	
Propionibacterium granulosum		93.97	0.50	XYL 1	1	99	90	
Prevotella intermedia		2.62	0.15		1	1	1	
Gemella morbillorum		2.39	0.15		99	99	1	
07516002	**Good identification**							**07516002**
					CAT			
Propionibacterium granulosum		96.49	0.39	XYL 1	90			
Lactobacillus fermentum		2.17	0.16		1			
Propionibacterium propionicum		1.34	0.12		1			
07516020	**Good identification**							**07516020**
Propionibacterium granulosum		100.0	0.39	XYL 1				
07516100	**Low discrimination**							**07516100**
					GRA	CAT		
Propionibacterium propionicum		86.73	0.43	NIT 86	99	1		
Fusobacterium mortiferum		4.60	0.20	XYL 1	1	5		
Propionibacterium granulosum		3.82	0.16		99	90		
07516102	**Acceptable identification**							**07516102**
Propionibacterium propionicum		90.35	0.46	NIT 86				
Bifidobacterium adolescentis		5.26	0.20	XYL 1				
Lactobacillus fermentum		2.41	0.20					
07516140	**Good identification**							**07516140**
Propionibacterium propionicum		100.0	0.43	NIT 86				
07516142	**Good identification**							**07516142**
Propionibacterium propionicum		100.0	0.46	NIT 86				
07516512	**Low discrimination**							**07516512**
Bifidobacterium adolescentis		84.40	0.36	GAL 99				
Bifidobacterium longum subsp. longum		15.60	0.21					
07517000	**Low discrimination**							**07517000**
					CAT			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		58.79	0.50	XYL 1	90	
Lactobacillus fermentum		32.73	0.50		1	
Actinomyces israelii		3.76	0.35		1	
07517001	**Low discrimination**					**07517001**
					CAT	
Actinomyces naeslundii		74.61	0.42	XYL 1	25	
Propionibacterium granulosum		16.31	0.16		90	
Lactobacillus fermentum		9.08	0.16		1	
07517002	**Low discrimination**					**07517002**
					CAT	
Lactobacillus fermentum		58.18	0.50	MAN 1	1	
Propionibacterium granulosum		26.12	0.39		90	
Actinomyces israelii		10.03	0.38		1	
07517003	**Low discrimination**					**07517003**
Actinomyces naeslundii		89.15	0.42	XYL 1		
Lactobacillus fermentum		10.85	0.16			
07517020	**Low discrimination**					**07517020**
					CAT	
Propionibacterium granulosum		52.44	0.39	XYL 1	90	
Lactobacillus fermentum		38.93	0.42		1	
Actinomyces naeslundii		6.46	0.39		25	
07517021	**Good identification**					**07517021**
Actinomyces naeslundii		100.0	0.42	XYL 1		
07517022	**Low discrimination**					**07517022**
					CAT	
Lactobacillus fermentum		60.58	0.42	MAN 1	1	
Propionibacterium granulosum		20.40	0.29		90	
Actinomyces naeslundii		10.06	0.39		25	
07517023	**Good identification**					**07517023**
Actinomyces naeslundii		100.0	0.42	XYL 1		
07517100	**Low discrimination**					**07517100**
Lactobacillus fermentum		58.47	0.54	MAN 1		
Actinomyces israelii		16.48	0.45			
Propionibacterium propionicum		14.76	0.43			
07517101	**Low discrimination**					**07517101**
Actinomyces naeslundii		79.15	0.49	XYL 1		
Propionibacterium propionicum		15.33	0.27			
Lactobacillus fermentum		5.52	0.20			
07517102	**Low discrimination**					**07517102**
Bifidobacterium adolescentis		52.85	0.54	ARA 87		
Lactobacillus fermentum		24.22	0.54			
Actinomyces israelii		10.24	0.48			
07517103	**Low discrimination**					**07517103**
Actinomyces naeslundii		64.85	0.49	XYL 1		
Propionibacterium propionicum		18.84	0.30			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		9.87	0.20			
07517112	**Good identification**					**07517112**
Bifidobacterium adolescentis		95.85	0.56	ARA 87		
Actinomyces israelii		2.27	0.34			
Bifidobacterium longum subsp. longum		1.20	0.21			
07517121	**Good identification**					**07517121**
					GRA	
Actinomyces naeslundii		95.49	0.49	XYL 1	99	
Prevotella melaninogenica		4.51	-.03		1	
07517123	**Acceptable identification**					**07517123**
					GRA	
Actinomyces naeslundii		91.04	0.49	XYL 1	99	
Bacteroides fragilis		8.96	-.03		1	
07517140	**Low discrimination**					**07517140**
Propionibacterium propionicum		64.86	0.43	NIT 86		
Actinomyces israelii		28.17	0.38	XYL 1		
Actinomyces naeslundii		4.38	0.34			
07517142	**Low discrimination**					**07517142**
Propionibacterium propionicum		45.08	0.46	NIT 86		
Bifidobacterium adolescentis		32.12	0.39	XYL 1		
Actinomyces israelii		19.57	0.41			
07517152	**Acceptable identification**					**07517152**
Bifidobacterium adolescentis		92.07	0.40	ARA 87		
Actinomyces israelii		6.86	0.27	SOR 11		
Propionibacterium propionicum		1.07	0.12			
07517161	**Good identification**					**07517161**
Actinomyces naeslundii		100.0	0.37	XYL 1		
07517163	**Good identification**					**07517163**
Actinomyces naeslundii		100.0	0.37	XYL 1		
07517320	**Good identification**					**07517320**
Bacteroides vulgatus		99.54	0.40	MAN 1		
Prevotella melaninogenica		0.24	-.03	ARA 93		
Parabacteroides distasonis		0.23	0.14			
07517322	**Low discrimination**					**07517322**
Parabacteroides distasonis		86.17	0.48	MAN 1		
Prevotella oralis		8.14	0.17			
Bacteroides vulgatus		3.83	0.06			
07517332	**Good identification**					**07517332**
Parabacteroides distasonis		100.0	0.43	MAN 1		
07517400	**Low discrimination**					**07517400**
Lactobacillus fermentum		87.40	0.50	MAN 1		
Actinomyces israelii		8.22	0.33			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		2.79	0.17			
07517402	**Low discrimination**					**07517402**
Lactobacillus fermentum		70.74	0.50	MAN 1		
Bifidobacterium adolescentis		17.02	0.34			
Actinomyces israelii		9.98	0.36			
07517412	**Low discrimination**					**07517412**
Bifidobacterium adolescentis		77.47	0.36	LAC 99		
Bifidobacterium longum subsp. longum		14.32	0.21			
Actinomyces israelii		5.55	0.22			
07517420	**Low discrimination**					**07517420**
					GRA	
Lactobacillus fermentum		62.72	0.42	MAN 1	99	
Bifidobacterium longum subsp. longum		24.04	0.28		99	
Bacteroides vulgatus		13.24	0.09		1	
07517422	**Low discrimination**					**07517422**
Lactobacillus fermentum		70.91	0.42	MAN 1		
Bifidobacterium longum subsp. longum		27.18	0.28			
Actinomyces israelii		1.92	0.16			
07517500	**Low discrimination**					**07517500**
Bifidobacterium longum subsp. longum		57.69	0.52	MAN 1		
Lactobacillus fermentum		29.77	0.54			
Actinomyces israelii		6.86	0.44			
07517502	**Low discrimination**					**07517502**
Bifidobacterium adolescentis		81.62	0.68			
Bifidobacterium longum subsp. longum		10.83	0.52			
Lactobacillus fermentum		5.59	0.54			
07517510	**Acceptable identification**					**07517510**
Bifidobacterium longum subsp. longum		93.62	0.56	MAN 1		
Bifidobacterium adolescentis		5.12	0.36			
Actinomyces israelii		0.98	0.30			
07517512	**Low discrimination**					**07517512**
Bifidobacterium adolescentis		84.15	0.70			
Bifidobacterium longum subsp. longum		15.56	0.56			
Actinomyces israelii		0.24	0.33			
07517513	**Low discrimination**					**07517513**
Bifidobacterium adolescentis		84.40	0.36	URE 1		
Bifidobacterium longum subsp. longum		15.60	0.21			
07517520	**Low discrimination**					**07517520**
					GRA	
Bifidobacterium longum subsp. longum		62.63	0.62	MAN 1	99	
Bacteroides vulgatus		34.50	0.43		1	
Lactobacillus fermentum		2.69	0.46		99	
07517522	**Good identification**					**07517522**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		91.65	0.62	MAN 1				
Lactobacillus fermentum		3.94	0.46					
Bifidobacterium adolescentis		1.74	0.34					
07517530	**Very good identification**							**07517530**
					GRA			
Bifidobacterium longum subsp. longum		99.67	0.66	MAN 1	99			
Bacteroides vulgatus		0.33	0.09		1			
07517532	**Very good identification**							**07517532**
					GRA			
Bifidobacterium longum subsp. longum		98.12	0.66	MAN 1	99			
Bifidobacterium adolescentis		1.34	0.36		99			
Parabacteroides distasonis		0.54	0.31		1			
07517540	**Low discrimination**							**07517540**
Actinomyces israelii		65.17	0.37	SAL 92				
Bifidobacterium longum subsp. longum		14.23	0.17	CEL 93				
Bifidobacterium adolescentis		13.25	0.19					
07517542	**Acceptable identification**							**07517542**
Bifidobacterium adolescentis		91.49	0.53	SOR 11				
Actinomyces israelii		6.82	0.40					
Bifidobacterium longum subsp. longum		0.99	0.17					
07517552	**Good identification**							**07517552**
Bifidobacterium adolescentis		97.64	0.54	SOR 11				
Bifidobacterium longum subsp. longum		1.48	0.21					
Actinomyces israelii		0.89	0.26					
07517712	**Low discrimination**							**07517712**
Bifidobacterium adolescentis		73.77	0.36	RHA 1				
Bifidobacterium longum subsp. longum		13.64	0.21					
Clostridium clostridioforme		11.00	0.18					
07517720	**Very good identification**							**07517720**
					GRA			
Bacteroides vulgatus		99.79	0.60	MAN 1	1			
Bifidobacterium longum subsp. longum		0.20	0.28		99			
Parabacteroides distasonis		0.01	0.10		1			
07517722	**Low discrimination**							**07517722**
					GRA			
Parabacteroides distasonis		46.03	0.44	MAN 1	1			
Bacteroides vulgatus		44.35	0.25		1			
Bifidobacterium longum subsp. longum		8.95	0.28		99			
07517732	**Low discrimination**							**07517732**
					GRA			
Parabacteroides distasonis		56.93	0.39	MAN 1	1			
Bifidobacterium longum subsp. longum		34.99	0.32		99			
Clostridium clostridioforme		7.05	0.18		50			
07520000	**Good identification**							**07520000**
					COC	SPO	CAT	
Propionibacterium granulosum		90.99	0.60	FRU 99	1	1	90	
Gemella morbillorum		4.29	0.30		99	1	1	
Clostridium botulinum biovar B		2.81	0.21		1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07520002	**Acceptable identification**							**07520002**
Propionibacterium granulosum		92.93	0.50	FRU 99				
Actinomyces viscosus		6.44	0.20					
Actinomyces naeslundii		0.63	0.27					
07520010	**Low discrimination**							**07520010**
					SPO	CAT		
Propionibacterium granulosum		89.97	0.45	FRU 99	1	90		
Clostridium botulinum biovar B		10.03	0.16	MLZ 12	99	1		
07520020	**Acceptable identification**							**07520020**
					COC	SPO	CAT	
Propionibacterium granulosum		91.34	0.50	FRU 99	1	1	90	
Clostridium botulinum biovar B		5.56	0.16		1	99	1	
Gemella morbillorum		1.91	0.14		99	1	1	
07520022	**Good identification**							**07520022**
Propionibacterium granulosum		97.38	0.39	FRU 99				
Actinomyces naeslundii		2.62	0.27					
07520040	**Good identification**							**07520040**
					SPO	CAT		
Clostridium botulinum biovar B		99.62	0.56	RAF 1	99	1		
Propionibacterium granulosum		0.33	0.26		1	90		
Clostridium botulinum biovar A		0.05	0.02		99	1		
07520050	**Good identification**							**07520050**
					SPO	CAT		
Clostridium botulinum biovar B		99.91	0.50	RAF 1	99	1		
Propionibacterium granulosum		0.09	0.11		1	90		
07520060	**Good identification**							**07520060**
					SPO	CAT		
Clostridium botulinum biovar B		99.83	0.50	RAF 1	99	1		
Propionibacterium granulosum		0.17	0.15		1	90		
07520070	**Good identification**							**07520070**
Clostridium botulinum biovar B		100.0	0.45	RAF 1				
07520101	**Low discrimination**							**07520101**
					SPO			
Actinomyces naeslundii		83.37	0.38	FRU 95	1			
Clostridium chauvoei		16.63	0.03	GAL 96	99			
07520121	**Good identification**							**07520121**
Actinomyces naeslundii		100.0	0.38	FRU 95				
07520123	**Good identification**							**07520123**
Actinomyces naeslundii		100.0	0.38	FRU 95				
07521000	**Good identification**							**07521000**
					COC	CAT		
Propionibacterium granulosum		91.93	0.60	FRU 99	1	90		
Actinomyces naeslundii		3.71	0.51		1	25		
Gemella morbillorum		2.33	0.26		99	1		
07521001	**Low discrimination**							**07521001**
					SPO	CAT		
Actinomyces naeslundii		84.00	0.54	FRU 95	1	25		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		14.02	0.26		1	90	
Clostridium chauvoei		1.99	0.03		99	1	
07521002	**Low discrimination**						**07521002**
Propionibacterium granulosum		81.55	0.50	FRU 99			
Actinomyces naeslundii		13.17	0.51				
Actinomyces viscosus		3.46	0.17				
07521003	**Acceptable identification**						**07521003**
Actinomyces naeslundii		90.89	0.54	FRU 95			
Actinomyces viscosus		5.32	0.09				
Propionibacterium granulosum		3.79	0.15				
07521010	**Good identification**						**07521010**
Propionibacterium granulosum		98.47	0.45	FRU 99			
Actinomyces naeslundii		1.53	0.29	MLZ 12			
07521020	**Low discrimination**						**07521020**
					CAT		
Propionibacterium granulosum		53.44	0.50	FRU 99	90		
Clostridium perfringens		33.65	0.39		1		
Actinomyces naeslundii		8.63	0.51		25		
07521021	**Acceptable identification**						**07521021**
					CAT		
Actinomyces naeslundii		93.64	0.54	FRU 95	25		
Propionibacterium granulosum		3.91	0.15		90		
Clostridium perfringens		2.46	0.05		1		
07521023	**Good identification**						**07521023**
Actinomyces naeslundii		100.0	0.54	FRU 95			
07521041	**Good identification**						**07521041**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07521042	**Low discrimination**						**07521042**
Actinomyces naeslundii		73.83	0.38	FRU 95			
Propionibacterium granulosum		26.18	0.15				
07521043	**Good identification**						**07521043**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07521061	**Good identification**						**07521061**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07521063	**Good identification**						**07521063**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07521100	**Low discrimination**						**07521100**
					GRA		
Actinomyces naeslundii		54.29	0.59	FRU 95	99		
Clostridium perfringens		16.88	0.28		99		
Fusobacterium mortiferum		8.59	0.33		1		
07521101	**Low discrimination**						**07521101**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		54.87	0.62	FRU 95	1		
Clostridium chauvoei		45.14	0.37		99		
07521102	**Low discrimination**						**07521102**
Actinomyces naeslundii		68.66	0.59	FRU 95			
Actinomyces viscosus		11.29	0.21				
Clostridium perfringens		10.04	0.22				
07521103	**Low discrimination**						**07521103**
					SPO		
Actinomyces naeslundii		84.62	0.62	FRU 95	1		
Clostridium chauvoei		12.29	0.24		99		
Actinomyces viscosus		3.09	0.13		1		
07521111	**Low discrimination**						**07521111**
					SPO		
Actinomyces naeslundii		86.36	0.40	FRU 95	1		
Clostridium chauvoei		13.64	0.03	MLZ 5	99		
07521113	**Good identification**						**07521113**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07521120	**Low discrimination**						**07521120**
					GRA		
Prevotella melaninogenica		56.88	0.50	FRU 90	1		
Clostridium perfringens		38.58	0.54	TRE 1	99		
Actinomyces naeslundii		3.84	0.59		99		
07521121	**Low discrimination**						**07521121**
					GRA	SPO	
Actinomyces naeslundii		69.36	0.62	FRU 95	99	1	
Clostridium chauvoei		19.02	0.29		99	99	
Prevotella melaninogenica		6.92	0.15		1	1	
07521122	**Low discrimination**						**07521122**
					GRA		
Clostridium perfringens		52.12	0.48	FRU 99	99		
Prevotella melaninogenica		31.10	0.37		1		
Actinomyces naeslundii		11.02	0.59		99		
07521123	**Good identification**						**07521123**
					SPO		
Actinomyces naeslundii		90.97	0.62	FRU 95	1		
Clostridium chauvoei		4.40	0.16		99		
Clostridium perfringens		2.90	0.14		84		
07521131	**Good identification**						**07521131**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07521133	**Good identification**						**07521133**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07521140	**Low discrimination**						**07521140**
Actinomyces naeslundii		74.88	0.46	FRU 95			
Clostridium perfringens		25.12	0.15				
07521141	**Good identification**						**07521141**
					SPO		
Actinomyces naeslundii		95.50	0.49	FRU 95	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		4.50	0.03		99	
07521142	**Low discrimination**					**07521142**
Actinomyces naeslundii		86.37	0.46	FRU 95		
Clostridium perfringens		13.64	0.10			
07521143	**Good identification**					**07521143**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07521160	**Low discrimination**					**07521160**
					GRA	
Clostridium perfringens		85.44	0.41	FRU 99	99	
Actinomyces naeslundii		7.88	0.46		99	
Prevotella melaninogenica		6.68	0.15		1	
07521161	**Acceptable identification**					**07521161**
Actinomyces naeslundii		93.19	0.49	FRU 95		
Clostridium perfringens		6.81	0.07			
07521162	**Low discrimination**					**07521162**
					GRA	
Clostridium perfringens		81.46	0.36	FRU 99	99	
Actinomyces naeslundii		15.96	0.46		99	
Prevotella melaninogenica		2.58	0.03		1	
07521163	**Good identification**					**07521163**
Actinomyces naeslundii		96.68	0.49	FRU 95		
Clostridium perfringens		3.32	0.01			
07521322	**Low discrimination**					**07521322**
					GRA	
Parabacteroides distasonis		81.51	0.43	FRU 99	1	
Prevotella oralis		7.85	0.13	XYL 90	1	
Clostridium perfringens		5.89	0.14		99	
07521332	**Good identification**					**07521332**
Parabacteroides distasonis		100.0	0.38	FRU 99		
07521500	**Good identification**					**07521500**
Bifidobacterium longum subsp. longum		98.76	0.51	FRU 90		
Lactobacillus fermentum		0.96	0.24			
Actinomyces naeslundii		0.28	0.25			
07521502	**Good identification**					**07521502**
Bifidobacterium longum subsp. longum		97.79	0.51	FRU 90		
Bifidobacterium adolescentis		0.99	0.18			
Lactobacillus fermentum		0.95	0.24			
07521510	**Good identification**					**07521510**
Bifidobacterium longum subsp. longum		100.0	0.55	FRU 90		
07521512	**Good identification**					**07521512**
Bifidobacterium longum subsp. longum		99.28	0.55	FRU 90		
Bifidobacterium adolescentis		0.72	0.20			
07521520	**Very good identification**					**07521520**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium longum subsp. longum		98.11	0.61	FRU 90	99	
Prevotella melaninogenica		1.04	0.15		1	
Clostridium perfringens		0.71	0.19		99	
07521522	**Very good identification**					**07521522**
					GRA	
Bifidobacterium longum subsp. longum		98.38	0.61	FRU 90	99	
Parabacteroides distasonis		0.94	0.31		1	
Clostridium perfringens		0.33	0.14		99	
07521530	**Very good identification**					**07521530**
Bifidobacterium longum subsp. longum		100.0	0.65	FRU 90		
07521532	**Very good identification**					**07521532**
					GRA	
Bifidobacterium longum subsp. longum		99.70	0.65	FRU 90	99	
Parabacteroides distasonis		0.30	0.26		1	
07521722	**Low discrimination**					**07521722**
					GRA	
Parabacteroides distasonis		72.84	0.39	FRU 99	1	
Bifidobacterium longum subsp. longum		25.64	0.27	XYL 90	99	
Clostridium clostridioforme		1.53	0.05		50	
07522000	**Very good identification**					**07522000**
Propionibacterium granulosum		99.57	0.58	FRU 99		
Actinomyces naeslundii		0.43	0.32			
07522002	**Good identification**					**07522002**
Propionibacterium granulosum		98.30	0.48	FRU 99		
Actinomyces naeslundii		1.70	0.32			
07522003	**Low discrimination**					**07522003**
Actinomyces naeslundii		71.99	0.35	FRU 95		
Propionibacterium granulosum		28.01	0.13	GAL 96		
07522010	**Good identification**					**07522010**
Propionibacterium granulosum		100.0	0.43	FRU 99		
07522020	**Good identification**					**07522020**
Propionibacterium granulosum		98.30	0.48	FRU 99		
Actinomyces naeslundii		1.70	0.32			
07522021	**Low discrimination**					**07522021**
Actinomyces naeslundii		71.99	0.35	FRU 95		
Propionibacterium granulosum		28.01	0.13	GAL 96		
07522022	**Acceptable identification**					**07522022**
Propionibacterium granulosum		93.53	0.37	FRU 99		
Actinomyces naeslundii		6.48	0.32			
07522023	**Good identification**					**07522023**
Actinomyces naeslundii		100.0	0.35	FRU 95		
07522100	**Low discrimination**					**07522100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Propionibacterium propionicum		73.61	0.42	NIT 86	1	
Actinomyces naeslundii		14.50	0.40	FRU 99	25	
Propionibacterium granulosum		11.89	0.24		90	
07522101	**Low discrimination**					**07522101**
Actinomyces naeslundii		72.67	0.43	FRU 95		
Propionibacterium propionicum		27.33	0.26	GAL 96		
07522102	**Low discrimination**					**07522102**
					CAT	
Propionibacterium propionicum		86.34	0.45	NIT 86	1	
Actinomyces naeslundii		11.34	0.40	FRU 99	25	
Propionibacterium granulosum		2.33	0.13		90	
07522103	**Low discrimination**					**07522103**
Actinomyces naeslundii		63.94	0.43	FRU 95		
Propionibacterium propionicum		36.06	0.29	GAL 96		
07522120	**Low discrimination**					**07522120**
Actinomyces naeslundii		82.99	0.40	FRU 95		
Propionibacterium granulosum		17.01	0.13	GAL 96		
07522121	**Good identification**					**07522121**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07522122	**Good identification**					**07522122**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07522123	**Good identification**					**07522123**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07522140	**Good identification**					**07522140**
Propionibacterium propionicum		96.64	0.42	NIT 86		
Actinomyces naeslundii		3.36	0.27	FRU 99		
07522142	**Good identification**					**07522142**
Propionibacterium propionicum		97.74	0.45	NIT 86		
Actinomyces naeslundii		2.27	0.27	FRU 99		
07523000	**Good identification**					**07523000**
Propionibacterium granulosum		90.59	0.58	FRU 99		
Actinomyces naeslundii		9.41	0.56			
07523001	**Good identification**					**07523001**
Actinomyces naeslundii		93.91	0.59	FRU 95		
Propionibacterium granulosum		6.09	0.24			
07523002	**Low discrimination**					**07523002**
Propionibacterium granulosum		70.65	0.48	FRU 99		
Actinomyces naeslundii		29.35	0.56			
07523003	**Very good identification**					**07523003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.41	0.59	FRU 95		
Propionibacterium granulosum		1.60	0.13			
07523010	**Good identification**					**07523010**
Propionibacterium granulosum		96.15	0.43	FRU 99		
Actinomyces naeslundii		3.85	0.34	MLZ 12		
07523011	**Low discrimination**					**07523011**
Actinomyces naeslundii		85.62	0.37	FRU 95		
Propionibacterium granulosum		14.38	0.09	MLZ 5		
07523013	**Good identification**					**07523013**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07523020	**Low discrimination**					**07523020**
					CAT	
Propionibacterium granulosum		70.23	0.48	FRU 99	90	
Actinomyces naeslundii		29.18	0.56		25	
Clostridium perfringens		0.59	0.05		1	
07523021	**Very good identification**					**07523021**
Actinomyces naeslundii		98.41	0.59	FRU 95		
Propionibacterium granulosum		1.60	0.13			
07523022	**Low discrimination**					**07523022**
Actinomyces naeslundii		62.43	0.56	FRU 95		
Propionibacterium granulosum		37.57	0.37			
07523023	**Very good identification**					**07523023**
Actinomyces naeslundii		100.0	0.59	FRU 95		
07523031	**Good identification**					**07523031**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07523033	**Good identification**					**07523033**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07523040	**Low discrimination**					**07523040**
Actinomyces naeslundii		64.47	0.43	FRU 95		
Propionibacterium granulosum		35.53	0.24			
07523041	**Good identification**					**07523041**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07523042	**Low discrimination**					**07523042**
Actinomyces naeslundii		87.89	0.43	FRU 95		
Propionibacterium granulosum		12.11	0.13			
07523043	**Good identification**					**07523043**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07523060	**Low discrimination**					**07523060**
Actinomyces naeslundii		87.89	0.43	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		12.11	0.13			
07523061	**Good identification**					**07523061**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07523062	**Good identification**					**07523062**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07523063	**Good identification**					**07523063**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07523100	**Low discrimination**					**07523100**
					CAT	
Actinomyces naeslundii		80.28	0.64	FRU 95	25	
Propionibacterium propionicum		16.98	0.42		1	
Propionibacterium granulosum		2.74	0.24		90	
07523101	**Very good identification**					**07523101**
					SPO	
Actinomyces naeslundii		98.04	0.67	FRU 95	1	
Propionibacterium propionicum		1.54	0.26		1	
Clostridium chauvoei		0.42	0.03		99	
07523102	**Low discrimination**					**07523102**
					CAT	
Actinomyces naeslundii		75.43	0.64	FRU 95	25	
Propionibacterium propionicum		23.93	0.45		1	
Propionibacterium granulosum		0.64	0.13		90	
07523103	**Very good identification**					**07523103**
Actinomyces naeslundii		97.70	0.67	FRU 95		
Propionibacterium propionicum		2.30	0.29			
07523110	**Acceptable identification**					**07523110**
Actinomyces naeslundii		91.87	0.42	FRU 95		
Propionibacterium granulosum		8.13	0.09	MLZ 5		
07523111	**Good identification**					**07523111**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07523112	**Good identification**					**07523112**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07523113	**Good identification**					**07523113**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07523120	**Low discrimination**					**07523120**
					GRA	
Actinomyces naeslundii		87.88	0.64	FRU 95	99	
Prevotella melaninogenica		6.78	0.15		1	
Clostridium perfringens		4.60	0.19		99	
07523121	**Very good identification**					**07523121**
Actinomyces naeslundii		100.0	0.67	FRU 95		
07523122	**Very good identification**					**07523122**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.22	0.64	FRU 95	99	
Clostridium perfringens		2.37	0.14		99	
Prevotella melaninogenica		1.41	0.03		1	
07523123	**Very good identification**					**07523123**
Actinomyces naeslundii		100.0	0.67	FRU 95		
07523130	**Good identification**					**07523130**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07523131	**Good identification**					**07523131**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07523132	**Good identification**					**07523132**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07523133	**Good identification**					**07523133**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07523140	**Low discrimination**					**07523140**
Propionibacterium propionicum		54.52	0.42	NIT 86		
Actinomyces naeslundii		45.48	0.51	FRU 99		
07523141	**Acceptable identification**					**07523141**
Actinomyces naeslundii		91.85	0.54	FRU 95		
Propionibacterium propionicum		8.15	0.26			
07523142	**Low discrimination**					**07523142**
Propionibacterium propionicum		64.26	0.45	NIT 86		
Actinomyces naeslundii		35.74	0.51	FRU 99		
07523143	**Low discrimination**					**07523143**
Actinomyces naeslundii		88.25	0.54	FRU 95		
Propionibacterium propionicum		11.75	0.29			
07523160	**Acceptable identification**					**07523160**
Actinomyces naeslundii		94.66	0.51	FRU 95		
Clostridium perfringens		5.34	0.07			
07523161	**Good identification**					**07523161**
Actinomyces naeslundii		100.0	0.54	FRU 95		
07523162	**Good identification**					**07523162**
Actinomyces naeslundii		97.41	0.51	FRU 95		
Clostridium perfringens		2.59	0.01			
07523163	**Good identification**					**07523163**
Actinomyces naeslundii		100.0	0.54	FRU 95		
07524520	**Good identification**					**07524520**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07524522	**Good identification**					**07524522**
					GRA	
Bifidobacterium longum subsp. longum		97.21	0.35	FRU 90	99	
Parabacteroides distasonis		2.79	0.13	GAL 99	1	
07524530	**Good identification**					**07524530**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
07524532	**Good identification**					**07524532**
Bifidobacterium longum subsp. longum		97.62	0.39	FRU 90		
Clostridium clostridioforme		2.39	0.10	GAL 99		
07525120	**Low discrimination**					**07525120**
					GRA	
Bifidobacterium longum subsp. longum		57.16	0.35	FRU 90	99	
Prevotella melaninogenica		19.98	0.15	ARA 99	1	
Clostridium perfringens		13.55	0.19		99	
07525122	**Low discrimination**					**07525122**
					GRA	
Parabacteroides distasonis		77.96	0.51	FRU 99	1	
Bifidobacterium longum subsp. longum		16.82	0.35		99	
Clostridium perfringens		1.88	0.14		99	
07525130	**Good identification**					**07525130**
					GRA	
Bifidobacterium longum subsp. longum		98.54	0.39	FRU 90	99	
Parabacteroides distasonis		1.46	0.12	ARA 99	1	
07525132	**Low discrimination**					**07525132**
					GRA	
Parabacteroides distasonis		59.45	0.46	FRU 99	1	
Bifidobacterium longum subsp. longum		40.55	0.39		99	
07525322	**Very good identification**					**07525322**
Parabacteroides distasonis		99.87	0.59	FRU 99		
Prevotella oralis		0.13	-.03			
07525332	**Good identification**					**07525332**
Parabacteroides distasonis		99.23	0.54	FRU 99		
Clostridium clostridioforme		0.77	0.13			
07525420	**Good identification**					**07525420**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90		
07525422	**Good identification**					**07525422**
					GRA	
Bifidobacterium longum subsp. longum		97.21	0.35	FRU 90	99	
Parabacteroides distasonis		2.79	0.13	LAC 99	1	
07525430	**Good identification**					**07525430**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
07525432	**Good identification**					**07525432**
Bifidobacterium longum subsp. longum		97.86	0.39	FRU 90		
Clostridium clostridioforme		2.14	0.09	LAC 99		
07525500	**Very good identification**					**07525500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		99.81	0.59	FRU 90		
Lactobacillus fermentum		0.19	0.20			
07525502	**Very good identification**					**07525502**
Bifidobacterium longum subsp. longum		97.40	0.59	FRU 90		
Bifidobacterium adolescentis		2.20	0.33			
Lactobacillus fermentum		0.19	0.20			
07525510	**Very good identification**					**07525510**
Bifidobacterium longum subsp. longum		100.0	0.63	FRU 90		
07525512	**Very good identification**					**07525512**
Bifidobacterium longum subsp. longum		98.12	0.63	FRU 90		
Bifidobacterium adolescentis		1.59	0.34			
Clostridium clostridioforme		0.29	0.18			
07525520	**Very good identification**					**07525520**
					GRA	
Bifidobacterium longum subsp. longum		99.14	0.69	FRU 90	99	
Bacteroides vulgatus		0.83	0.20		1	
Parabacteroides distasonis		0.03	0.13		1	
07525521	**Good identification**					**07525521**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90		
07525522	**Very good identification**					**07525522**
					GRA	
Bifidobacterium longum subsp. longum		97.18	0.69	FRU 90	99	
Parabacteroides distasonis		2.79	0.47		1	
Clostridium clostridioforme		0.03	0.07		50	
07525523	**Good identification**					**07525523**
					GRA	
Bifidobacterium longum subsp. longum		97.21	0.35	FRU 90	99	
Parabacteroides distasonis		2.79	0.13	URE 1	1	
07525530	**Very good identification**					**07525530**
Bifidobacterium longum subsp. longum		100.0	0.73	FRU 90		
07525531	**Good identification**					**07525531**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
07525532	**Very good identification**					**07525532**
					GRA	
Bifidobacterium longum subsp. longum		99.03	0.73	FRU 90	99	
Parabacteroides distasonis		0.90	0.43		1	
Clostridium clostridioforme		0.07	0.18		50	
07525533	**Good identification**					**07525533**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
07525560	**Good identification**					**07525560**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90		
07525562	**Good identification**					**07525562**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		97.21	0.35	FRU 90	99		
Parabacteroides distasonis		2.79	0.13	SOR 1	1		
07525570	**Good identification**						**07525570**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
07525572	**Good identification**						**07525572**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
07525720	**Low discrimination**						**07525720**
					GRA		
Bacteroides vulgatus		87.26	0.36	FRU 99	1		
Bifidobacterium longum subsp. longum		11.73	0.35	TRE 4	99		
Parabacteroides distasonis		1.01	0.21		1		
07525722	**Low discrimination**						**07525722**
					GRA		
Parabacteroides distasonis		85.88	0.55	FRU 99	1		
Bifidobacterium longum subsp. longum		10.08	0.35		99		
Clostridium clostridioforme		2.02	0.22		50		
07525730	**Acceptable identification**						**07525730**
					GRA		
Bifidobacterium longum subsp. longum		93.34	0.39	FRU 90	99		
Bacteroides vulgatus		4.12	0.02	RHA 1	1		
Parabacteroides distasonis		2.54	0.17		1		
07525732	**Low discrimination**						**07525732**
					GRA		
Parabacteroides distasonis		63.25	0.51	FRU 99	1		
Bifidobacterium longum subsp. longum		23.47	0.39		99		
Clostridium clostridioforme		12.61	0.33		50		
07527520	**Good identification**						**07527520**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90			
07527522	**Good identification**						**07527522**
					GRA		
Bifidobacterium longum subsp. longum		97.21	0.35	FRU 90	99		
Parabacteroides distasonis		2.79	0.13	MAN 1	1		
07527530	**Good identification**						**07527530**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
07527532	**Good identification**						**07527532**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
07530000	**Very good identification**						**07530000**
					GRA	CAT	
Propionibacterium granulosum		98.57	0.94		99	90	
Prevotella intermedia		0.69	0.49		1	1	
Fusobacterium mortiferum		0.51	0.58		1	5	
07530001	**Good identification**						**07530001**
					GRA	CAT	
Propionibacterium granulosum		94.12	0.60	URE 1	99	90	
Actinomyces naeslundii		4.51	0.52		99	25	
Prevotella intermedia		0.66	0.15		1	1	
07530002	**Very good identification**						**07530002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Propionibacterium granulosum		98.29	0.84		99	90		
Actinomyces viscosus		1.31	0.42		99	76		
Fusobacterium mortiferum		0.23	0.42		1	5		
07530003	**Low discrimination**							**07530003**
Propionibacterium granulosum		61.32	0.50	URE 1				
Actinomyces viscosus		26.92	0.34					
Actinomyces naeslundii		11.76	0.52					
07530010	**Very good identification**							**07530010**
					GRA	SPO	CAT	
Propionibacterium granulosum		99.45	0.79	MLZ 12	99	1	90	
Clostridium botulinum biovar B		0.45	0.26		99	99	1	
Prevotella intermedia		0.05	0.15		1	1	1	
07530011	**Good identification**							**07530011**
Propionibacterium granulosum		98.18	0.45	MLZ 12				
Actinomyces naeslundii		1.82	0.30	URE 1				
07530012	**Very good identification**							**07530012**
Propionibacterium granulosum		99.85	0.69	MLZ 12				
Actinomyces viscosus		0.10	0.08					
Actinomyces naeslundii		0.05	0.27					
07530020	**Very good identification**							**07530020**
					CAT	SPO		
Propionibacterium granulosum		98.92	0.84		90	1		
Clostridium perfringens		0.63	0.39		1	84		
Clostridium botulinum biovar B		0.24	0.26		1	99		
07530021	**Low discrimination**							**07530021**
					CAT			
Propionibacterium granulosum		83.47	0.50	URE 1	90			
Actinomyces naeslundii		16.00	0.52		25			
Clostridium perfringens		0.53	0.05		1			
07530022	**Very good identification**							**07530022**
					CAT			
Propionibacterium granulosum		98.25	0.73		90			
Clostridium perfringens		1.18	0.33		1			
Actinomyces naeslundii		0.51	0.49		25			
07530023	**Low discrimination**							**07530023**
Propionibacterium granulosum		56.04	0.39	URE 1				
Actinomyces naeslundii		42.97	0.52					
Actinomyces viscosus		0.99	0.00					
07530030	**Very good identification**							**07530030**
					SPO	CAT		
Propionibacterium granulosum		99.03	0.69	MLZ 12	1	90		
Clostridium botulinum biovar B		0.88	0.21		99	1		
Actinomyces naeslundii		0.05	0.27		1	25		
07530032	**Very good identification**							**07530032**
Propionibacterium granulosum		99.80	0.59	MLZ 12				
Actinomyces naeslundii		0.20	0.27					

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07530040	**Good identification**							**07530040**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		92.31	0.66	RAF 1	99	99	1	
Propionibacterium granulosum		7.55	0.60		99	1	90	
Prevotella intermedia		0.05	0.15		1	1	1	
07530042	**Low discrimination**							**07530042**
					SPO	CAT		
Propionibacterium granulosum		65.36	0.50	SOR 1	1	90		
Clostridium botulinum biovar B		32.29	0.32		99	1		
Actinomyces naeslundii		1.47	0.36		1	25		
07530043	**Low discrimination**							**07530043**
					SPO	CAT		
Actinomyces naeslundii		63.41	0.39	GAL 96	1	25		
Propionibacterium granulosum		18.93	0.15		1	90		
Clostridium botulinum biovar B		9.35	-.03		99	1		
07530050	**Very good identification**							**07530050**
					SPO	CAT		
Clostridium botulinum biovar B		97.79	0.61	RAF 1	99	1		
Propionibacterium granulosum		2.21	0.45		1	90		
07530060	**Very good identification**							**07530060**
					CAT	SPO		
Clostridium botulinum biovar B		95.47	0.61	RAF 1	1	99		
Propionibacterium granulosum		3.96	0.50		90	1		
Clostridium perfringens		0.48	0.26		1	84		
07530062	**Low discrimination**							**07530062**
					CAT	SPO		
Propionibacterium granulosum		43.68	0.39	SOR 1	90	1		
Clostridium botulinum biovar B		42.52	0.26		1	99		
Clostridium perfringens		9.86	0.21		1	84		
07530063	**Good identification**							**07530063**
Actinomyces naeslundii		100.0	0.39	GAL 96				
07530070	**Good identification**							**07530070**
					SPO	CAT		
Clostridium botulinum biovar B		98.86	0.55	RAF 1	99	1		
Propionibacterium granulosum		1.14	0.35		1	90		
07530100	**Low discrimination**							**07530100**
					GRA	CAT		
Fusobacterium mortiferum		50.04	0.65	MLT 11	1	5		
Propionibacterium granulosum		41.71	0.60		99	90		
Actinomyces naeslundii		3.79	0.57		99	25		
07530101	**Low discrimination**							**07530101**
					GRA	CAT		
Actinomyces naeslundii		80.63	0.60	GAL 96	99	25		
Fusobacterium mortiferum		7.16	0.30		1	5		
Propionibacterium granulosum		5.97	0.26		99	90		
07530102	**Low discrimination**							**07530102**
					GRA	CAT		
Actinomyces viscosus		50.81	0.47	NIT 99	99	76		
Propionibacterium granulosum		21.70	0.50	TRE 7	99	90		
Fusobacterium mortiferum		11.57	0.48		1	5		
07530103	**Low discrimination**							**07530103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		56.84	0.38	NIT 99			
Actinomyces naeslundii		39.74	0.60	TRE 7			
Propionibacterium propionicum		2.29	0.29				
07530110	**Low discrimination**						**07530110**
					GRA	CAT	
Propionibacterium granulosum		88.97	0.45	LAC 1	99	90	
Fusobacterium mortiferum		7.91	0.30	MLZ 12	1	5	
Actinomyces naeslundii		3.12	0.35		99	25	
07530111	**Low discrimination**						**07530111**
Actinomyces naeslundii		83.90	0.38	GAL 96			
Propionibacterium granulosum		16.10	0.11	MLZ 5			
07530113	**Low discrimination**						**07530113**
Actinomyces naeslundii		78.46	0.38	GAL 96			
Actinomyces viscosus		21.54	0.04	MLZ 5			
07530120	**Low discrimination**						**07530120**
					GRA	CAT	
Clostridium perfringens		69.30	0.54	GAL 99	99	1	
Propionibacterium granulosum		15.01	0.50		99	90	
Prevotella melaninogenica		9.29	0.32		1	1	
07530121	**Low discrimination**						**07530121**
					CAT		
Actinomyces naeslundii		89.00	0.60	GAL 96	25		
Clostridium perfringens		7.61	0.19		1		
Propionibacterium granulosum		1.65	0.15		90		
07530122	**Low discrimination**						**07530122**
					CAT		
Clostridium perfringens		69.77	0.48	GAL 99	1		
Actinomyces naeslundii		11.68	0.57		25		
Propionibacterium granulosum		8.03	0.39		90		
07530123	**Good identification**						**07530123**
Actinomyces naeslundii		94.82	0.60	GAL 96			
Clostridium perfringens		3.81	0.14				
Actinomyces viscosus		1.37	0.04				
07530131	**Good identification**						**07530131**
Actinomyces naeslundii		100.0	0.38	GAL 96			
07530133	**Good identification**						**07530133**
Actinomyces naeslundii		100.0	0.38	GAL 96			
07530141	**Low discrimination**						**07530141**
					SPO		
Actinomyces naeslundii		78.77	0.47	GAL 96	1		
Propionibacterium propionicum		17.15	0.26		1		
Clostridium botulinum biovar B		4.08	-.03		99		
07530142	**Low discrimination**						**07530142**
Propionibacterium propionicum		70.03	0.45	NIT 86			
Actinomyces naeslundii		15.89	0.44	MAN 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		5.85	0.12				
07530143	**Low discrimination**						**07530143**
Actinomyces naeslundii		71.00	0.47	GAL 96			
Propionibacterium propionicum		23.19	0.29				
Actinomyces viscosus		5.81	0.04				
07530160	**Low discrimination**						**07530160**
					SPO		
Clostridium perfringens		73.09	0.41	GAL 99	84		
Clostridium botulinum biovar B		20.22	0.26		99		
Actinomyces naeslundii		5.33	0.44		1		
07530161	**Acceptable identification**						**07530161**
Actinomyces naeslundii		91.55	0.47	GAL 96			
Clostridium perfringens		8.45	0.07				
07530162	**Low discrimination**						**07530162**
Clostridium perfringens		86.57	0.36	GAL 99			
Actinomyces naeslundii		13.43	0.44				
07530163	**Good identification**						**07530163**
Actinomyces naeslundii		95.84	0.47	GAL 96			
Clostridium perfringens		4.16	0.01				
07530200	**Very good identification**						**07530200**
					GRA	CAT	
Propionibacterium granulosum		99.56	0.68	RHA 3	99	90	
Prevotella intermedia		0.23	0.15		1	1	
Fusobacterium mortiferum		0.17	0.24		1	5	
07530202	**Very good identification**						**07530202**
Propionibacterium granulosum		99.57	0.58	RHA 3			
Actinomyces viscosus		0.43	0.08				
07530210	**Good identification**						**07530210**
Propionibacterium granulosum		100.0	0.53	RHA 3			
07530212	**Good identification**						**07530212**
Propionibacterium granulosum		97.39	0.43	RHA 3			
Clostridium clostridioforme		2.61	0.12	MLZ 12			
07530220	**Very good identification**						**07530220**
					CAT		
Propionibacterium granulosum		99.79	0.58	RHA 3	90		
Clostridium perfringens		0.21	0.05		1		
07530222	**Good identification**						**07530222**
Propionibacterium granulosum		100.0	0.48	RHA 3			
07530230	**Good identification**						**07530230**
Propionibacterium granulosum		100.0	0.43	RHA 3			
07530322	**Low discrimination**						**07530322**
					GRA		
Parabacteroides distasonis		81.09	0.43	GAL 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella oralis		7.81	0.13	XYL 90	1			
Clostridium perfringens		5.86	0.14		99			
07530332	**Low discrimination**							**07530332**
Parabacteroides distasonis		81.36	0.38	GAL 99				
Clostridium clostridioforme		18.65	0.21	XYL 90				
07530400	**Very good identification**							**07530400**
					COC	GRA	CAT	
Propionibacterium granulosum		97.82	0.60	ARA 1	1	99	90	
Prevotella intermedia		0.69	0.15		1	1	1	
Gemella morbillorum		0.63	0.15		99	99	1	
07530402	**Good identification**							**07530402**
					CAT			
Propionibacterium granulosum		97.26	0.50	ARA 1	90			
Actinomyces viscosus		1.29	0.08		76			
Lactobacillus fermentum		0.94	0.20		1			
07530410	**Good identification**							**07530410**
Propionibacterium granulosum		100.0	0.45	ARA 1				
07530420	**Good identification**							**07530420**
					CAT			
Propionibacterium granulosum		98.78	0.50	ARA 1	90			
Clostridium perfringens		0.63	0.05		1			
Bifidobacterium longum subsp. longum		0.59	0.09		1			
07530422	**Good identification**							**07530422**
					CAT			
Propionibacterium granulosum		95.71	0.39	ARA 1	90			
Bifidobacterium longum subsp. longum		2.28	0.09		1			
Clostridium clostridioforme		2.01	0.07		33			
07530510	**Good identification**							**07530510**
					CAT			
Bifidobacterium longum subsp. longum		97.85	0.37	GAL 99	1			
Propionibacterium granulosum		2.15	0.11		90			
07530512	**Low discrimination**							**07530512**
Bifidobacterium longum subsp. longum		71.49	0.37	GAL 99				
Clostridium clostridioforme		22.80	0.27					
Bifidobacterium adolescentis		5.71	0.20					
07530520	**Low discrimination**							**07530520**
					CAT			
Bifidobacterium longum subsp. longum		89.14	0.43	GAL 99	1			
Clostridium perfringens		7.04	0.19		1			
Propionibacterium granulosum		1.53	0.15		90			
07530522	**Low discrimination**							**07530522**
					GRA			
Bifidobacterium longum subsp. longum		84.22	0.43	GAL 99	99			
Parabacteroides distasonis		8.86	0.31		1			
Clostridium perfringens		3.13	0.14		99			
07530530	**Good identification**							**07530530**
Bifidobacterium longum subsp. longum		100.0	0.47	GAL 99				
07530532	**Low discrimination**							**07530532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium longum subsp. longum		89.85	0.47	GAL 99	99		
Clostridium clostridioforme		7.17	0.27		50		
Parabacteroides distasonis		2.99	0.26		1		
07530712	**Good identification**						**07530712**
Clostridium clostridioforme		100.0	0.42	SAL 88			
07530722	**Low discrimination**						**07530722**
					GRA		
Parabacteroides distasonis		57.99	0.39	GAL 99	1		
Clostridium clostridioforme		40.15	0.31	XYL 90	50		
Bifidobacterium longum subsp. longum		1.86	0.09		99		
07530732	**Low discrimination**						**07530732**
					GRA		
Clostridium clostridioforme		84.18	0.42	SAL 88	50		
Parabacteroides distasonis		14.37	0.34	XYL 91	1		
Bifidobacterium longum subsp. longum		1.45	0.13		99		
07531000	**Very good identification**						**07531000**
					GRA	CAT	
Propionibacterium granulosum		97.05	0.94		99	90	
Fusobacterium mortiferum		0.76	0.61		1	5	
Actinomyces naeslundii		0.75	0.73		99	25	
07531001	**Low discrimination**						**07531001**
					GRA	CAT	
Actinomyces naeslundii		52.87	0.76		99	25	
Propionibacterium granulosum		45.96	0.60		99	90	
Fusobacterium mortiferum		0.36	0.27		1	5	
07531002	**Good identification**						**07531002**
					CAT		
Propionibacterium granulosum		94.12	0.84		90		
Actinomyces naeslundii		2.92	0.73		25		
Lactobacillus fermentum		0.91	0.54		1		
07531003	**Low discrimination**						**07531003**
Actinomyces naeslundii		78.09	0.76				
Propionibacterium granulosum		16.97	0.50				
Actinomyces viscosus		4.57	0.31				
07531010	**Very good identification**						**07531010**
					GRA	CAT	
Propionibacterium granulosum		99.50	0.79	MLZ 12	99	90	
Actinomyces naeslundii		0.30	0.51		99	25	
Fusobacterium mortiferum		0.06	0.27		1	5	
07531011	**Low discrimination**						**07531011**
Propionibacterium granulosum		69.26	0.45	MLZ 12			
Actinomyces naeslundii		30.75	0.54	URE 1			
07531012	**Very good identification**						**07531012**
					CAT		
Propionibacterium granulosum		98.47	0.69	MLZ 12	90		
Actinomyces naeslundii		1.18	0.51		25		
Lactobacillus jensenii		0.10	0.11		1		
07531013	**Low discrimination**						**07531013**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		63.97	0.54	MLZ 5			
Propionibacterium granulosum		36.03	0.35				
07531020	**Low discrimination**						**07531020**
					CAT		
Propionibacterium granulosum		59.83	0.84		90		
Clostridium perfringens		37.68	0.73		1		
Actinomyces naeslundii		1.85	0.73		25		
07531021	**Low discrimination**						**07531021**
					CAT		
Actinomyces naeslundii		73.72	0.76		25		
Propionibacterium granulosum		16.02	0.50		90		
Clostridium perfringens		10.09	0.39		1		
07531022	**Low discrimination**						**07531022**
					CAT		
Clostridium perfringens		50.45	0.67	LAC 88	1		
Propionibacterium granulosum		42.56	0.73		90		
Actinomyces naeslundii		5.27	0.73		25		
07531023	**Low discrimination**						**07531023**
					CAT		
Actinomyces naeslundii		89.31	0.76		25		
Clostridium perfringens		5.75	0.33		1		
Propionibacterium granulosum		4.85	0.39		90		
07531030	**Good identification**						**07531030**
					CAT		
Propionibacterium granulosum		94.32	0.69	MLZ 12	90		
Clostridium perfringens		4.40	0.39		1		
Actinomyces naeslundii		1.13	0.51		25		
07531031	**Low discrimination**						**07531031**
					CAT		
Actinomyces naeslundii		62.92	0.54	MLZ 5	25		
Propionibacterium granulosum		35.43	0.35		90		
Clostridium perfringens		1.65	0.05		1		
07531032	**Low discrimination**						**07531032**
					CAT		
Propionibacterium granulosum		86.40	0.59	MLZ 12	90		
Clostridium perfringens		7.59	0.33		1		
Actinomyces naeslundii		4.13	0.51		25		
07531033	**Low discrimination**						**07531033**
Actinomyces naeslundii		87.66	0.54	MLZ 5			
Propionibacterium granulosum		12.34	0.25				
07531040	**Low discrimination**						**07531040**
					SPO	CAT	
Propionibacterium granulosum		73.11	0.60	SOR 1	1	90	
Actinomyces naeslundii		9.89	0.60		1	25	
Clostridium botulinum biovar B		9.03	0.32		99	1	
07531041	**Good identification**						**07531041**
					SPO	CAT	
Actinomyces naeslundii		94.30	0.63		1	25	
Propionibacterium granulosum		4.69	0.26		1	90	
Clostridium botulinum biovar B		0.58	-.03		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07531042	**Low discrimination**						**07531042**
					CAT		
Propionibacterium granulosum		57.46	0.50	SOR 1	90		
Actinomyces naeslundii		31.10	0.60		25		
Clostridium perfringens		9.93	0.29		1		
07531043	**Very good identification**						**07531043**
Actinomyces naeslundii		98.77	0.63				
Propionibacterium granulosum		1.23	0.15				
07531050	**Low discrimination**						**07531050**
					SPO	CAT	
Propionibacterium granulosum		66.44	0.45	MLZ 12	1	90	
Clostridium botulinum biovar B		29.64	0.26	SOR 1	99	1	
Actinomyces naeslundii		3.47	0.38		1	25	
07531051	**Low discrimination**						**07531051**
Actinomyces naeslundii		88.58	0.41	MLZ 5			
Propionibacterium granulosum		11.42	0.11				
07531053	**Good identification**						**07531053**
Actinomyces naeslundii		100.0	0.41	MLZ 5			
07531060	**Low discrimination**						**07531060**
					CAT		
Clostridium perfringens		86.90	0.61	LAC 88	1		
Propionibacterium granulosum		7.32	0.50		90		
Actinomyces naeslundii		3.96	0.60		25		
07531061	**Low discrimination**						**07531061**
					CAT		
Actinomyces naeslundii		86.19	0.63		25		
Clostridium perfringens		12.73	0.26		1		
Propionibacterium granulosum		1.07	0.15		90		
07531062	**Low discrimination**						**07531062**
					CAT		
Clostridium perfringens		87.54	0.55	LAC 88	1		
Actinomyces naeslundii		8.48	0.60		25		
Propionibacterium granulosum		3.92	0.39		90		
07531063	**Good identification**						**07531063**
Actinomyces naeslundii		93.50	0.63				
Clostridium perfringens		6.50	0.21				
07531071	**Good identification**						**07531071**
Actinomyces naeslundii		100.0	0.41	MLZ 5			
07531073	**Good identification**						**07531073**
Actinomyces naeslundii		100.0	0.41	MLZ 5			
07531100	**Low discrimination**						**07531100**
					GRA		
Clostridium perfringens		36.74	0.62	NAG 97	99		
Actinomyces naeslundii		22.69	0.81		99		
Fusobacterium mortiferum		18.71	0.68		1		
07531101	**Low discrimination**						**07531101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		88.84	0.84		1		
Clostridium chauvoei		8.98	0.44		99		
Clostridium perfringens		0.97	0.28		84		
07531102	**Low discrimination**						**07531102**
Actinomyces naeslundii		39.94	0.81				
Clostridium perfringens		30.44	0.56				
Lactobacillus fermentum		7.12	0.58				
07531103	**Good identification**						**07531103**
					SPO		
Actinomyces naeslundii		93.92	0.84		1		
Actinomyces viscosus		3.43	0.35		1		
Clostridium chauvoei		1.67	0.31		99		
07531110	**Low discrimination**						**07531110**
					CAT		
Propionibacterium granulosum		35.95	0.45	LAC 1	90		
Actinomyces naeslundii		30.28	0.59	MLZ 12	25		
Bifidobacterium longum subsp. longum		16.56	0.37		1		
07531111	**Very good identification**						**07531111**
					SPO	CAT	
Actinomyces naeslundii		97.33	0.62	MLZ 5	1	25	
Clostridium chauvoei		1.89	0.09		99	1	
Propionibacterium granulosum		0.78	0.11		1	90	
07531112	**Low discrimination**						**07531112**
Actinomyces naeslundii		34.63	0.59	MLZ 5			
Bifidobacterium adolescentis		22.36	0.40				
Bifidobacterium longum subsp. longum		18.94	0.37				
07531113	**Very good identification**						**07531113**
Actinomyces naeslundii		99.30	0.62	MLZ 5			
Actinomyces viscosus		0.70	0.01				
07531120	**Low discrimination**						**07531120**
					GRA		
Clostridium perfringens		86.22	0.88		99		
Prevotella melaninogenica		11.56	0.66		1		
Actinomyces naeslundii		1.65	0.81		99		
07531121	**Low discrimination**						**07531121**
					GRA		
Actinomyces naeslundii		69.66	0.84		99		
Clostridium perfringens		24.57	0.54		99		
Prevotella melaninogenica		3.29	0.32		1		
07531122	**Low discrimination**						**07531122**
					GRA		
Clostridium perfringens		84.66	0.82		99		
Prevotella melaninogenica		4.59	0.54		1		
Parabacteroides distasonis		3.95	0.69		1		
07531123	**Low discrimination**						**07531123**
					GRA		
Actinomyces naeslundii		83.68	0.84		99		
Clostridium perfringens		13.89	0.48		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		0.75	0.19		1	
07531130	**Low discrimination**					**07531130**
					GRA	
Clostridium perfringens		66.92	0.54	MLZ 1	99	
Bifidobacterium longum subsp. longum		14.56	0.47		99	
Prevotella melaninogenica		8.97	0.32		1	
07531131	**Good identification**					**07531131**
Actinomyces naeslundii		91.08	0.62	MLZ 5		
Clostridium perfringens		6.16	0.19			
Bifidobacterium longum subsp. longum		1.34	0.13			
07531132	**Low discrimination**					**07531132**
					GRA	
Parabacteroides distasonis		58.04	0.64	XYL 90	1	
Clostridium perfringens		23.35	0.48		99	
Bifidobacterium longum subsp. longum		10.80	0.47		99	
07531133	**Low discrimination**					**07531133**
					GRA	
Actinomyces naeslundii		88.83	0.62	MLZ 5	99	
Parabacteroides distasonis		7.03	0.30		1	
Clostridium perfringens		2.83	0.14		99	
07531140	**Low discrimination**					**07531140**
Clostridium perfringens		58.72	0.50	NAG 97		
Actinomyces naeslundii		33.59	0.68			
Propionibacterium propionicum		4.11	0.42			
07531141	**Very good identification**					**07531141**
Actinomyces naeslundii		97.41	0.71			
Clostridium perfringens		1.15	0.15			
Propionibacterium propionicum		0.88	0.26			
07531142	**Low discrimination**					**07531142**
Actinomyces naeslundii		48.31	0.68			
Clostridium perfringens		39.74	0.44			
Propionibacterium propionicum		8.87	0.45			
07531143	**Very good identification**					**07531143**
Actinomyces naeslundii		97.92	0.71			
Propionibacterium propionicum		1.33	0.29			
Clostridium perfringens		0.54	0.10			
07531150	**Low discrimination**					**07531150**
					CAT	
Actinomyces naeslundii		71.25	0.46	MLZ 5	25	
Clostridium perfringens		23.91	0.15		1	
Propionibacterium granulosum		4.84	0.11		90	
07531151	**Good identification**					**07531151**
Actinomyces naeslundii		100.0	0.49	MLZ 5		
07531152	**Low discrimination**					**07531152**
Actinomyces naeslundii		62.11	0.46	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		28.09	0.24				
Clostridium perfringens		9.81	0.10				
07531153	**Good identification**						**07531153**
Actinomyces naeslundii		100.0	0.49	MLZ 5			
07531160	**Very good identification**						**07531160**
					GRA		
Clostridium perfringens		97.55	0.75		99		
Actinomyces naeslundii		1.73	0.68		99		
Prevotella melaninogenica		0.69	0.32		1		
07531161	**Low discrimination**						**07531161**
					GRA		
Actinomyces naeslundii		72.19	0.71		99		
Clostridium perfringens		27.48	0.41		99		
Prevotella melaninogenica		0.20	-.03		1		
07531162	**Very good identification**						**07531162**
					GRA		
Clostridium perfringens		95.63	0.70		99		
Actinomyces naeslundii		3.60	0.68		99		
Prevotella melaninogenica		0.28	0.19		1		
07531163	**Low discrimination**						**07531163**
Actinomyces naeslundii		84.81	0.71				
Clostridium perfringens		15.19	0.36				
07531170	**Acceptable identification**						**07531170**
Clostridium perfringens		90.02	0.41	MLZ 1			
Actinomyces naeslundii		8.30	0.46				
Bifidobacterium longum subsp. longum		1.04	0.13				
07531171	**Acceptable identification**						**07531171**
Actinomyces naeslundii		93.19	0.49	MLZ 5			
Clostridium perfringens		6.81	0.07				
07531172	**Low discrimination**						**07531172**
					GRA		
Clostridium perfringens		73.95	0.36	MLZ 1	99		
Actinomyces naeslundii		14.49	0.46		99		
Parabacteroides distasonis		9.75	0.30		1		
07531173	**Good identification**						**07531173**
Actinomyces naeslundii		96.68	0.49	MLZ 5			
Clostridium perfringens		3.32	0.01				
07531200	**Very good identification**						**07531200**
					GRA	CAT	
Propionibacterium granulosum		99.04	0.68	RHA 3	99	90	
Fusobacterium mortiferum		0.25	0.27		1	5	
Actinomyces naeslundii		0.25	0.39		99	25	
07531202	**Very good identification**						**07531202**
					CAT		
Propionibacterium granulosum		97.50	0.58	RHA 3	90		
Actinomyces naeslundii		0.99	0.39		25		
Lactobacillus fermentum		0.31	0.20		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07531203	**Low discrimination**						**07531203**
Actinomyces naeslundii		60.04	0.42	RHA 1			
Propionibacterium granulosum		39.96	0.24				
07531210	**Good identification**						**07531210**
Propionibacterium granulosum		100.0	0.53	RHA 3			
07531212	**Acceptable identification**						**07531212**
Propionibacterium granulosum		92.55	0.43	RHA 3			
Clostridium clostridioforme		7.45	0.21	MLZ 12			
07531220	**Low discrimination**						**07531220**
					CAT		
Propionibacterium granulosum		81.90	0.58	RHA 3	90		
Clostridium perfringens		16.84	0.39		1		
Actinomyces naeslundii		0.83	0.39		25		
07531221	**Low discrimination**						**07531221**
					CAT		
Actinomyces naeslundii		55.48	0.42	RHA 1	25		
Propionibacterium granulosum		36.93	0.24		90		
Clostridium perfringens		7.59	0.05		1		
07531222	**Low discrimination**						**07531222**
					GRA	CAT	
Propionibacterium granulosum		48.17	0.48	RHA 3	99	90	
Parabacteroides distasonis		19.13	0.43		1	76	
Clostridium perfringens		18.65	0.33		99	1	
07531223	**Low discrimination**						**07531223**
Actinomyces naeslundii		85.74	0.42	RHA 1			
Propionibacterium granulosum		14.26	0.13				
07531230	**Good identification**						**07531230**
					CAT		
Propionibacterium granulosum		98.50	0.43	RHA 3	90		
Clostridium perfringens		1.50	0.05	MLZ 12	1		
07531232	**Low discrimination**						**07531232**
					GRA		
Parabacteroides distasonis		54.26	0.38	XYL 90	1		
Propionibacterium granulosum		34.60	0.33	LAC 99	99		
Clostridium clostridioforme		11.14	0.21		50		
07531301	**Acceptable identification**						**07531301**
					SPO		
Actinomyces naeslundii		90.82	0.49	RHA 1	1		
Clostridium chauvoei		9.18	0.09		99		
07531302	**Low discrimination**						**07531302**
					GRA		
Parabacteroides distasonis		42.78	0.43	XYL 90	1		
Prevotella oralis		21.47	0.25	NAG 99	1		
Actinomyces naeslundii		12.41	0.46		99		
07531303	**Good identification**						**07531303**
Actinomyces naeslundii		96.48	0.49	RHA 1			
Actinomyces viscosus		3.52	0.01				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07531312	**Low discrimination**					**07531312**
					GRA	
Parabacteroides distasonis		57.42	0.38	XYL 90	1	
Clostridium clostridioforme		39.48	0.30	NAG 99	50	
Actinomyces naeslundii		1.63	0.25		99	
07531320	**Low discrimination**					**07531320**
					GRA	
Clostridium perfringens		76.27	0.54	RHA 1	99	
Prevotella melaninogenica		10.22	0.32		1	
Parabacteroides distasonis		5.02	0.43		1	
07531321	**Low discrimination**					**07531321**
					GRA	
Actinomyces naeslundii		69.76	0.49	RHA 1	99	
Clostridium perfringens		24.60	0.19		99	
Prevotella melaninogenica		3.30	-.03		1	
07531322	**Low discrimination**					**07531322**
					GRA	
Parabacteroides distasonis		84.88	0.77	XYL 90	1	
Prevotella oralis		8.18	0.47		1	
Clostridium perfringens		6.13	0.48		99	
07531323	**Low discrimination**					**07531323**
					GRA	
Parabacteroides distasonis		62.36	0.43	XYL 90	1	
Actinomyces naeslundii		27.13	0.49	URE 1	99	
Prevotella oralis		6.01	0.13		1	
07531330	**Low discrimination**					**07531330**
					GRA	
Parabacteroides distasonis		69.48	0.38	XYL 90	1	
Clostridium perfringens		19.80	0.19	BGL 99	99	
Bifidobacterium longum subsp. longum		4.31	0.13		99	
07531332	**Very good identification**					**07531332**
Parabacteroides distasonis		98.91	0.72	XYL 90		
Clostridium clostridioforme		0.69	0.30			
Prevotella oralis		0.18	0.13			
07531333	**Good identification**					**07531333**
					GRA	
Parabacteroides distasonis		95.92	0.38	XYL 90	1	
Actinomyces naeslundii		4.08	0.28	URE 1	99	
07531341	**Good identification**					**07531341**
Actinomyces naeslundii		100.0	0.37	RHA 1		
07531343	**Good identification**					**07531343**
Actinomyces naeslundii		100.0	0.37	RHA 1		
07531360	**Good identification**					**07531360**
					GRA	
Clostridium perfringens		97.58	0.41	RHA 1	99	
Actinomyces naeslundii		1.73	0.34		99	
Prevotella melaninogenica		0.69	-.03		1	
07531361	**Low discrimination**					**07531361**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		72.43	0.37	RHA 1		
Clostridium perfringens		27.57	0.07			
07531362	**Low discrimination**					**07531362**
					GRA	
Clostridium perfringens		54.26	0.36	RHA 1	99	
Parabacteroides distasonis		39.86	0.43		1	
Prevotella oralis		3.84	0.13		1	
07531363	**Low discrimination**					**07531363**
Actinomyces naeslundii		84.81	0.37	RHA 1		
Clostridium perfringens		15.19	0.01			
07531372	**Good identification**					**07531372**
					GRA	
Parabacteroides distasonis		97.51	0.38	XYL 90	1	
Clostridium perfringens		2.49	0.01	SOR 1	99	
07531400	**Low discrimination**					**07531400**
					CAT	
Propionibacterium granulosum		76.56	0.60	ARA 1	90	
Lactobacillus fermentum		18.25	0.54		1	
Bifidobacterium longum subsp. longum		2.83	0.33		1	
07531401	**Low discrimination**					**07531401**
					CAT	
Actinomyces naeslundii		48.16	0.42	ARA 1	25	
Propionibacterium granulosum		41.87	0.26		90	
Lactobacillus fermentum		9.98	0.20		1	
07531402	**Low discrimination**					**07531402**
					CAT	
Propionibacterium granulosum		45.67	0.50	ARA 1	90	
Lactobacillus fermentum		43.55	0.54		1	
Bifidobacterium longum subsp. longum		6.74	0.33		1	
07531403	**Low discrimination**					**07531403**
					CAT	
Actinomyces naeslundii		70.19	0.42	ARA 1	25	
Propionibacterium granulosum		15.26	0.15		90	
Lactobacillus fermentum		14.55	0.20		1	
07531410	**Low discrimination**					**07531410**
					CAT	
Propionibacterium granulosum		67.64	0.45	ARA 1	90	
Bifidobacterium longum subsp. longum		31.16	0.37	MLZ 12	1	
Lactobacillus fermentum		1.20	0.20		1	
07531412	**Low discrimination**					**07531412**
					CAT	
Bifidobacterium longum subsp. longum		51.37	0.37	LAC 99	1	
Propionibacterium granulosum		27.88	0.35		90	
Clostridium clostridioforme		14.68	0.26		33	
07531420	**Low discrimination**					**07531420**
					CAT	
Propionibacterium granulosum		38.83	0.50	ARA 1	90	
Clostridium perfringens		24.45	0.39		1	
Bifidobacterium longum subsp. longum		22.92	0.43		1	
07531421	**Low discrimination**					**07531421**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		67.45	0.42	ARA 1	25	
Propionibacterium granulosum		14.66	0.15		90	
Clostridium perfringens		9.23	0.05		1	
07531422	**Low discrimination**					**07531422**
Bifidobacterium longum subsp. longum		37.65	0.43	LAC 99		
Lactobacillus fermentum		20.28	0.46			
Clostridium perfringens		18.90	0.33			
07531423	**Low discrimination**					**07531423**
Actinomyces naeslundii		88.63	0.42	ARA 1		
Bifidobacterium longum subsp. longum		11.37	0.09			
07531430	**Low discrimination**					**07531430**
					CAT	
Bifidobacterium longum subsp. longum		87.57	0.47	LAC 99	1	
Propionibacterium granulosum		11.88	0.35		90	
Clostridium perfringens		0.55	0.05		1	
07531432	**Low discrimination**					**07531432**
					CAT	
Bifidobacterium longum subsp. longum		87.82	0.47	LAC 99	1	
Clostridium clostridioforme		6.28	0.26		33	
Propionibacterium granulosum		2.98	0.25		90	
07531500	**Low discrimination**					**07531500**
Bifidobacterium longum subsp. longum		88.21	0.67			
Lactobacillus fermentum		9.39	0.58			
Clostridium perfringens		0.86	0.28			
07531502	**Low discrimination**					**07531502**
Bifidobacterium longum subsp. longum		80.63	0.67			
Bifidobacterium adolescentis		8.97	0.53			
Lactobacillus fermentum		8.59	0.58			
07531510	**Very good identification**					**07531510**
Bifidobacterium longum subsp. longum		99.82	0.71			
Bifidobacterium adolescentis		0.08	0.20			
Lactobacillus fermentum		0.06	0.24			
07531511	**Good identification**					**07531511**
Bifidobacterium longum subsp. longum		97.30	0.37	URE 1		
Actinomyces naeslundii		2.70	0.28			
07531512	**Good identification**					**07531512**
Bifidobacterium longum subsp. longum		91.60	0.71			
Bifidobacterium adolescentis		7.31	0.54			
Clostridium clostridioforme		0.89	0.35			
07531513	**Acceptable identification**					**07531513**
Bifidobacterium longum subsp. longum		90.29	0.37	URE 1		
Bifidobacterium adolescentis		7.21	0.20			
Actinomyces naeslundii		2.50	0.28			
07531520	**Good identification**					**07531520**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium longum subsp. longum		90.34	0.78		99	
Clostridium perfringens		7.14	0.54		99	
Prevotella melaninogenica		0.96	0.32		1	
07531521	**Low discrimination**					**07531521**
Bifidobacterium longum subsp. longum		75.18	0.43	URE	1	
Actinomyces naeslundii		16.85	0.49			
Clostridium perfringens		5.94	0.19			
07531522	**Low discrimination**					**07531522**
					GRA	
Bifidobacterium longum subsp. longum		86.35	0.78		99	
Parabacteroides distasonis		9.09	0.65		1	
Clostridium perfringens		3.21	0.48		99	
07531523	**Low discrimination**					**07531523**
					GRA	
Bifidobacterium longum subsp. longum		72.71	0.43	URE	1 99	
Actinomyces naeslundii		16.29	0.49		99	
Parabacteroides distasonis		7.65	0.31		1	
07531530	**Very good identification**					**07531530**
					GRA	
Bifidobacterium longum subsp. longum		99.90	0.82		99	
Clostridium perfringens		0.05	0.19		99	
Parabacteroides distasonis		0.03	0.26		1	
07531531	**Good identification**					**07531531**
Bifidobacterium longum subsp. longum		99.31	0.47	URE	1	
Actinomyces naeslundii		0.69	0.28			
07531532	**Very good identification**					**07531532**
					GRA	
Bifidobacterium longum subsp. longum		96.50	0.82		99	
Parabacteroides distasonis		3.21	0.60		1	
Clostridium clostridioforme		0.23	0.35		50	
07531533	**Good identification**					**07531533**
					GRA	
Bifidobacterium longum subsp. longum		96.14	0.47	URE	1 99	
Parabacteroides distasonis		3.20	0.26		1	
Actinomyces naeslundii		0.67	0.28		99	
07531541	**Good identification**					**07531541**
Actinomyces naeslundii		100.0	0.37	ARA	1	
07531542	**Low discrimination**					**07531542**
Bifidobacterium adolescentis		50.26	0.37	XYL	87	
Bifidobacterium longum subsp. longum		36.94	0.33	SOR	11	
Lactobacillus fermentum		3.93	0.24			
07531543	**Good identification**					**07531543**
Actinomyces naeslundii		100.0	0.37	ARA	1	
07531550	**Good identification**					**07531550**
Bifidobacterium longum subsp. longum		100.0	0.37	SOR	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07531552	**Low discrimination**					**07531552**
Bifidobacterium longum subsp. longum		50.59	0.37	SOR 1		
Bifidobacterium adolescentis		49.41	0.39			
07531560	**Low discrimination**					**07531560**
Clostridium perfringens		58.76	0.41	ARA 1		
Bifidobacterium longum subsp. longum		39.43	0.43			
Actinomyces naeslundii		1.04	0.34			
07531561	**Low discrimination**					**07531561**
Actinomyces naeslundii		61.12	0.37	ARA 1		
Clostridium perfringens		23.27	0.07			
Bifidobacterium longum subsp. longum		15.61	0.09			
07531562	**Low discrimination**					**07531562**
					GRA	
Bifidobacterium longum subsp. longum		54.06	0.43	SOR 1	99	
Clostridium perfringens		37.91	0.36		99	
Parabacteroides distasonis		5.69	0.31		1	
07531563	**Low discrimination**					**07531563**
Actinomyces naeslundii		69.71	0.37	ARA 1		
Bifidobacterium longum subsp. longum		17.81	0.09			
Clostridium perfringens		12.49	0.01			
07531570	**Good identification**					**07531570**
Bifidobacterium longum subsp. longum		99.12	0.47	SOR 1		
Clostridium perfringens		0.88	0.07			
07531572	**Good identification**					**07531572**
					GRA	
Bifidobacterium longum subsp. longum		96.39	0.47	SOR 1	99	
Parabacteroides distasonis		3.21	0.26		1	
Clostridium perfringens		0.40	0.01		99	
07531612	**Good identification**					**07531612**
Clostridium clostridioforme		99.21	0.41	SAL 88		
Propionibacterium granulosum		0.79	0.09	XYL 91		
07531622	**Low discrimination**					**07531622**
					GRA	
Parabacteroides distasonis		59.03	0.39	XYL 90	1	
Clostridium clostridioforme		36.63	0.30	LAC 99	50	
Propionibacterium granulosum		2.45	0.13		99	
07531632	**Low discrimination**					**07531632**
					GRA	
Clostridium clostridioforme		82.66	0.41	SAL 88	50	
Parabacteroides distasonis		15.75	0.34	XYL 91	1	
Bifidobacterium longum subsp. longum		1.59	0.13		99	
07531702	**Low discrimination**					**07531702**
					GRA	
Clostridium clostridioforme		51.40	0.39	SAL 88	50	
Parabacteroides distasonis		24.74	0.39	XYL 91	1	
Bifidobacterium longum subsp. longum		19.60	0.33		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07531710	**Acceptable identification**					**07531710**
Bifidobacterium longum subsp. longum		93.38	0.37	RHA 1		
Clostridium clostridioforme		6.62	0.16			
07531712	**Low discrimination**					**07531712**
					GRA	
Clostridium clostridioforme		82.59	0.50	SAL 88	50	
Bifidobacterium longum subsp. longum		11.77	0.37	XYL 91	99	
Parabacteroides distasonis		4.70	0.34		1	
07531720	**Low discrimination**					**07531720**
					GRA	
Bacteroides vulgatus		77.90	0.41	TRE 4	1	
Bifidobacterium longum subsp. longum		15.55	0.43	XYL 98	99	
Parabacteroides distasonis		4.91	0.39		1	
07531722	**Good identification**					**07531722**
					GRA	
Parabacteroides distasonis		94.06	0.73	XYL 90	1	
Bifidobacterium longum subsp. longum		3.01	0.43		99	
Clostridium clostridioforme		1.97	0.39		50	
07531723	**Acceptable identification**					**07531723**
					GRA	
Parabacteroides distasonis		94.97	0.39	XYL 90	1	
Bifidobacterium longum subsp. longum		3.04	0.09	URE 1	99	
Clostridium clostridioforme		1.99	0.05		50	
07531730	**Low discrimination**					**07531730**
					GRA	
Bifidobacterium longum subsp. longum		87.16	0.47	RHA 1	99	
Parabacteroides distasonis		8.70	0.34		1	
Bacteroides vulgatus		2.59	0.07		1	
07531732	**Low discrimination**					**07531732**
					GRA	
Parabacteroides distasonis		77.96	0.69	XYL 90	1	
Clostridium clostridioforme		13.84	0.50		50	
Bifidobacterium longum subsp. longum		7.89	0.47		99	
07531762	**Acceptable identification**					**07531762**
					GRA	
Parabacteroides distasonis		92.99	0.39	XYL 90	1	
Bifidobacterium longum subsp. longum		2.98	0.09	SOR 1	99	
Clostridium perfringens		2.09	0.01		99	
07532000	**Excellent identification**					**07532000**
					CAT	
Propionibacterium granulosum		99.84	0.92		90	
Actinomyces naeslundii		0.08	0.54		25	
Propionibacterium propionicum		0.06	0.42		1	
07532001	**Low discrimination**					**07532001**
					CAT	
Propionibacterium granulosum		88.47	0.58	URE 1	90	
Actinomyces naeslundii		10.91	0.57		25	
Propionibacterium propionicum		0.62	0.26		1	
07532002	**Very good identification**					**07532002**
					CAT	
Propionibacterium granulosum		99.28	0.82		90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		0.38	0.45		1		
Actinomyces naeslundii		0.33	0.54		25		
07532003	**Low discrimination**						**07532003**
					CAT		
Propionibacterium granulosum		64.90	0.48	URE 1	90		
Actinomyces naeslundii		32.01	0.57		25		
Propionibacterium propionicum		2.71	0.29		1		
07532010	**Very good identification**						**07532010**
Propionibacterium granulosum		99.97	0.77	MLZ 12			
Actinomyces naeslundii		0.03	0.32				
07532011	**Good identification**						**07532011**
Propionibacterium granulosum		95.46	0.43	MLZ 12			
Actinomyces naeslundii		4.54	0.35	URE 1			
07532012	**Very good identification**						**07532012**
Propionibacterium granulosum		99.87	0.67	MLZ 12			
Actinomyces naeslundii		0.13	0.32				
07532020	**Very good identification**						**07532020**
					CAT		
Propionibacterium granulosum		99.66	0.82		90		
Actinomyces naeslundii		0.33	0.54		25		
Clostridium perfringens		0.01	0.05		1		
07532021	**Low discrimination**						**07532021**
Propionibacterium granulosum		66.97	0.48	URE 1			
Actinomyces naeslundii		33.03	0.57				
07532022	**Very good identification**						**07532022**
Propionibacterium granulosum		98.69	0.71				
Actinomyces naeslundii		1.31	0.54				
07532023	**Low discrimination**						**07532023**
Actinomyces naeslundii		66.37	0.57	GAL 96			
Propionibacterium granulosum		33.63	0.37				
07532030	**Very good identification**						**07532030**
Propionibacterium granulosum		99.87	0.67	MLZ 12			
Actinomyces naeslundii		0.13	0.32				
07532032	**Very good identification**						**07532032**
Propionibacterium granulosum		99.49	0.57	MLZ 12			
Actinomyces naeslundii		0.51	0.32				
07532040	**Low discrimination**						**07532040**
					SPO	CAT	
Propionibacterium granulosum		80.60	0.58	SOR 1	1	90	
Clostridium botulinum biovar B		13.20	0.32		99	1	
Propionibacterium propionicum		5.04	0.42		1	1	
07532041	**Low discrimination**						**07532041**
					CAT		
Actinomyces naeslundii		53.78	0.44	GAL 96	25		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		24.96	0.24		90		
Propionibacterium propionicum		17.17	0.26		1		
07532042	**Low discrimination**						**07532042**
					CAT		
Propionibacterium granulosum		69.46	0.48	SOR 1	90		
Propionibacterium propionicum		26.05	0.45		1		
Actinomyces naeslundii		4.03	0.41		25		
07532043	**Low discrimination**						**07532043**
					CAT		
Actinomyces naeslundii		62.70	0.44	GAL 96	25		
Propionibacterium propionicum		30.02	0.29		1		
Propionibacterium granulosum		7.28	0.13		90		
07532050	**Low discrimination**						**07532050**
					SPO	CAT	
Propionibacterium granulosum		62.84	0.43	MLZ 12	1	90	
Clostridium botulinum biovar B		37.16	0.26	SOR 1	99	1	
07532060	**Low discrimination**						**07532060**
					SPO	CAT	
Propionibacterium granulosum		72.43	0.48	SOR 1	1	90	
Clostridium botulinum biovar B		23.36	0.26		99	1	
Actinomyces naeslundii		4.20	0.41		1	25	
07532061	**Low discrimination**						**07532061**
Actinomyces naeslundii		89.60	0.44	GAL 96			
Propionibacterium granulosum		10.40	0.13				
07532062	**Low discrimination**						**07532062**
Propionibacterium granulosum		81.16	0.37	SOR 1			
Actinomyces naeslundii		18.84	0.41				
07532063	**Good identification**						**07532063**
Actinomyces naeslundii		100.0	0.44	GAL 96			
07532100	**Low discrimination**						**07532100**
					CAT		
Propionibacterium propionicum		83.20	0.76	NIT 86	1		
Propionibacterium granulosum		13.44	0.58		90		
Actinomyces naeslundii		3.15	0.62		25		
07532101	**Low discrimination**						**07532101**
					CAT		
Propionibacterium propionicum		65.57	0.60	NIT 86	1		
Actinomyces naeslundii		33.47	0.65	URE 10	25		
Propionibacterium granulosum		0.96	0.24		90		
07532102	**Good identification**						**07532102**
					CAT		
Propionibacterium propionicum		94.95	0.79	NIT 86	1		
Propionibacterium granulosum		2.56	0.48		90		
Actinomyces naeslundii		2.39	0.62		25		
07532103	**Low discrimination**						**07532103**
Propionibacterium propionicum		74.34	0.63	NIT 86			
Actinomyces naeslundii		25.29	0.65	URE 10			
Actinomyces viscosus		0.19	0.04				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07532110	**Low discrimination**					**07532110**
					CAT	
Propionibacterium granulosum		64.57	0.43	LAC 1	90	
Propionibacterium propionicum		29.60	0.42	MLZ 12	1	
Actinomyces naeslundii		5.83	0.40		25	
07532111	**Low discrimination**					**07532111**
					CAT	
Actinomyces naeslundii		68.94	0.43	GAL 96	25	
Propionibacterium propionicum		25.92	0.26	MLZ 5	1	
Propionibacterium granulosum		5.14	0.09		90	
07532112	**Low discrimination**					**07532112**
					CAT	
Propionibacterium propionicum		66.90	0.45	NIT 86	1	
Propionibacterium granulosum		24.32	0.33	MLZ 1	90	
Actinomyces naeslundii		8.79	0.40		25	
07532113	**Low discrimination**					**07532113**
Actinomyces naeslundii		63.94	0.43	GAL 96		
Propionibacterium propionicum		36.06	0.29	MLZ 5		
07532120	**Low discrimination**					**07532120**
					CAT	
Propionibacterium granulosum		44.32	0.48	LAC 1	90	
Actinomyces naeslundii		41.49	0.62		25	
Propionibacterium propionicum		11.08	0.42		1	
07532121	**Very good identification**					**07532121**
					CAT	
Actinomyces naeslundii		97.37	0.65	GAL 96	25	
Propionibacterium propionicum		1.93	0.26		1	
Propionibacterium granulosum		0.70	0.13		90	
07532122	**Low discrimination**					**07532122**
					CAT	
Actinomyces naeslundii		58.86	0.62	GAL 96	25	
Propionibacterium propionicum		23.59	0.45		1	
Propionibacterium granulosum		15.72	0.37		90	
07532123	**Very good identification**					**07532123**
Actinomyces naeslundii		97.12	0.65	GAL 96		
Propionibacterium propionicum		2.88	0.29			
07532131	**Good identification**					**07532131**
Actinomyces naeslundii		100.0	0.43	GAL 96		
07532132	**Low discrimination**					**07532132**
Actinomyces naeslundii		59.10	0.40	GAL 96		
Propionibacterium granulosum		40.90	0.22	MLZ 5		
07532133	**Good identification**					**07532133**
Actinomyces naeslundii		100.0	0.43	GAL 96		
07532140	**Very good identification**					**07532140**
					CAT	
Propionibacterium propionicum		99.15	0.76	NIT 86	1	
Actinomyces naeslundii		0.66	0.49		25	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		0.16	0.24		90	
07532141	**Good identification**					**07532141**
Propionibacterium propionicum		91.74	0.60	NIT 86		
Actinomyces naeslundii		8.26	0.52	URE 10		
07532142	**Very good identification**					**07532142**
					CAT	
Propionibacterium propionicum		99.53	0.79	NIT 86	1	
Actinomyces naeslundii		0.44	0.49		25	
Propionibacterium granulosum		0.03	0.13		90	
07532143	**Good identification**					**07532143**
Propionibacterium propionicum		94.34	0.63	NIT 86		
Actinomyces naeslundii		5.66	0.52	URE 10		
07532150	**Acceptable identification**					**07532150**
					CAT	
Propionibacterium propionicum		94.63	0.42	NIT 86	1	
Actinomyces naeslundii		3.29	0.27	MLZ 1	25	
Propionibacterium granulosum		2.08	0.09		90	
07532152	**Good identification**					**07532152**
Propionibacterium propionicum		97.74	0.45	NIT 86		
Actinomyces naeslundii		2.27	0.27	MLZ 1		
07532160	**Low discrimination**					**07532160**
Propionibacterium propionicum		57.21	0.42	NIT 86		
Actinomyces naeslundii		37.79	0.49	NAG 1		
Clostridium perfringens		2.69	0.07			
07532161	**Low discrimination**					**07532161**
Actinomyces naeslundii		89.92	0.52	GAL 96		
Propionibacterium propionicum		10.08	0.26			
07532162	**Low discrimination**					**07532162**
Propionibacterium propionicum		68.72	0.45	NIT 86		
Actinomyces naeslundii		30.26	0.49	NAG 1		
Clostridium perfringens		1.02	0.01			
07532163	**Low discrimination**					**07532163**
Actinomyces naeslundii		85.60	0.52	GAL 96		
Propionibacterium propionicum		14.40	0.29			
07532200	**Very good identification**					**07532200**
Propionibacterium granulosum		100.0	0.66	RHA 3		
07532202	**Good identification**					**07532202**
Propionibacterium granulosum		100.0	0.56	RHA 3		
07532210	**Good identification**					**07532210**
Propionibacterium granulosum		100.0	0.51	RHA 3		
07532212	**Good identification**					**07532212**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		100.0	0.41	RHA 3		
07532220	**Good identification**					**07532220**
Propionibacterium granulosum		100.0	0.56	RHA 3		
07532222	**Good identification**					**07532222**
Propionibacterium granulosum		100.0	0.46	RHA 3		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07532230	**Good identification**					**07532230**
Propionibacterium granulosum		100.0	0.41	RHA 3		
07532300	**Low discrimination**					**07532300**
					CAT	
Propionibacterium propionicum		65.25	0.42	NIT 86	1	
Propionibacterium granulosum		32.28	0.32	RHA 1	90	
Actinomyces naeslundii		2.47	0.28		25	
07532302	**Acceptable identification**					**07532302**
					CAT	
Propionibacterium propionicum		90.28	0.45	NIT 86	1	
Propionibacterium granulosum		7.44	0.22	RHA 1	90	
Actinomyces naeslundii		2.28	0.28		25	
07532340	**Good identification**					**07532340**
Propionibacterium propionicum		100.0	0.42	NIT 86		
07532342	**Good identification**					**07532342**
Propionibacterium propionicum		100.0	0.45	NIT 86		
07532400	**Very good identification**					**07532400**
Propionibacterium granulosum		100.0	0.58	ARA 1		
07532402	**Good identification**					**07532402**
Propionibacterium granulosum		100.0	0.48	ARA 1		
07532410	**Good identification**					**07532410**
Propionibacterium granulosum		100.0	0.43	ARA 1		
07532420	**Good identification**					**07532420**
Propionibacterium granulosum		100.0	0.48	ARA 1		
07532422	**Good identification**					**07532422**
Propionibacterium granulosum		100.0	0.37	ARA 1		
07532500	**Low discrimination**					**07532500**
					CAT	
Propionibacterium propionicum		83.38	0.42	NIT 86	1	
Propionibacterium granulosum		13.47	0.24	ARA 1	90	
Actinomyces naeslundii		3.15	0.28		25	
07532502	**Acceptable identification**					**07532502**
					CAT	
Propionibacterium propionicum		90.94	0.45	NIT 86	1	
Bifidobacterium adolescentis		4.32	0.17	ARA 1	1	
Propionibacterium granulosum		2.45	0.13		90	
07532540	**Good identification**					**07532540**
Propionibacterium propionicum		100.0	0.42	NIT 86		
07532542	**Good identification**					**07532542**
Propionibacterium propionicum		100.0	0.45	NIT 86		
07533000	**Very good identification**					**07533000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Propionibacterium granulosum		97.95	0.92		90	
Actinomyces naeslundii		1.95	0.78		25	
Propionibacterium propionicum		0.06	0.42		1	
07533001	**Low discrimination**					**07533001**
					CAT	
Actinomyces naeslundii		74.62	0.81		25	
Propionibacterium granulosum		25.21	0.58		90	
Propionibacterium propionicum		0.18	0.26		1	
07533002	**Good identification**					**07533002**
					CAT	
Propionibacterium granulosum		92.22	0.82		90	
Actinomyces naeslundii		7.35	0.78		25	
Propionibacterium propionicum		0.35	0.45		1	
07533003	**Good identification**					**07533003**
					CAT	
Actinomyces naeslundii		91.91	0.81		25	
Propionibacterium granulosum		7.76	0.48		90	
Propionibacterium propionicum		0.32	0.29		1	
07533010	**Very good identification**					**07533010**
Propionibacterium granulosum		99.24	0.77	MLZ 12		
Actinomyces naeslundii		0.76	0.56			
07533011	**Low discrimination**					**07533011**
Actinomyces naeslundii		53.32	0.59	MLZ 5		
Propionibacterium granulosum		46.68	0.43			
07533012	**Very good identification**					**07533012**
					CAT	
Propionibacterium granulosum		96.97	0.67	MLZ 12	90	
Actinomyces naeslundii		2.98	0.56		25	
Actinomyces israelii		0.05	0.15		1	
07533013	**Low discrimination**					**07533013**
Actinomyces naeslundii		82.05	0.59	MLZ 5		
Propionibacterium granulosum		17.96	0.33			
07533020	**Good identification**					**07533020**
					CAT	
Propionibacterium granulosum		91.89	0.82		90	
Actinomyces naeslundii		7.33	0.78		25	
Clostridium perfringens		0.78	0.39		1	
07533021	**Good identification**					**07533021**
					CAT	
Actinomyces naeslundii		92.15	0.81		25	
Propionibacterium granulosum		7.78	0.48		90	
Clostridium perfringens		0.07	0.05		1	
07533022	**Low discrimination**					**07533022**
					CAT	
Propionibacterium granulosum		74.91	0.71		90	
Actinomyces naeslundii		23.89	0.78		25	
Clostridium perfringens		1.19	0.33		1	
07533023	**Very good identification**					**07533023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.93	0.81			
Propionibacterium granulosum		2.07	0.37			
07533030	**Very good identification**					**07533030**
					CAT	
Propionibacterium granulosum		96.96	0.67	MLZ 12	90	
Actinomyces naeslundii		2.98	0.56		25	
Clostridium perfringens		0.06	0.05		1	
07533031	**Low discrimination**					**07533031**
Actinomyces naeslundii		82.05	0.59	MLZ 5		
Propionibacterium granulosum		17.96	0.33			
07533032	**Low discrimination**					**07533032**
Propionibacterium granulosum		89.04	0.57	MLZ 12		
Actinomyces naeslundii		10.96	0.56			
07533033	**Good identification**					**07533033**
Actinomyces naeslundii		94.81	0.59	MLZ 5		
Propionibacterium granulosum		5.19	0.22			
07533040	**Low discrimination**					**07533040**
					CAT	
Propionibacterium granulosum		70.61	0.58	SOR 1	90	
Actinomyces naeslundii		24.59	0.65		25	
Propionibacterium propionicum		4.41	0.42		1	
07533041	**Very good identification**					**07533041**
					CAT	
Actinomyces naeslundii		96.84	0.68		25	
Propionibacterium granulosum		1.87	0.24		90	
Propionibacterium propionicum		1.29	0.26		1	
07533042	**Low discrimination**					**07533042**
					CAT	
Actinomyces naeslundii		50.03	0.65		25	
Propionibacterium granulosum		35.92	0.48		90	
Propionibacterium propionicum		13.47	0.45		1	
07533043	**Very good identification**					**07533043**
					CAT	
Actinomyces naeslundii		97.58	0.68		25	
Propionibacterium propionicum		1.95	0.29		1	
Propionibacterium granulosum		0.47	0.13		90	
07533050	**Low discrimination**					**07533050**
Propionibacterium granulosum		88.15	0.43	MLZ 12		
Actinomyces naeslundii		11.85	0.43	SOR 1		
07533051	**Good identification**					**07533051**
Actinomyces naeslundii		95.23	0.46	MLZ 5		
Propionibacterium granulosum		4.77	0.09			
07533053	**Good identification**					**07533053**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07533060	**Low discrimination**					**07533060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Actinomyces naeslundii		54.58	0.65		25	
Propionibacterium granulosum		39.19	0.48		90	
Clostridium perfringens		6.23	0.26		1	
07533061	**Very good identification**					**07533061**
Actinomyces naeslundii		99.52	0.68			
Propionibacterium granulosum		0.48	0.13			
07533062	**Low discrimination**					**07533062**
					CAT	
Actinomyces naeslundii		81.09	0.65		25	
Propionibacterium granulosum		14.56	0.37		90	
Clostridium perfringens		4.36	0.21		1	
07533063	**Very good identification**					**07533063**
Actinomyces naeslundii		100.0	0.68			
07533071	**Good identification**					**07533071**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07533072	**Low discrimination**					**07533072**
Actinomyces naeslundii		68.26	0.43	MLZ 5		
Propionibacterium granulosum		31.74	0.22			
07533073	**Good identification**					**07533073**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07533100	**Low discrimination**					**07533100**
					CAT	
Propionibacterium propionicum		47.91	0.76	NIT 86	1	
Actinomyces naeslundii		43.47	0.86		25	
Propionibacterium granulosum		7.74	0.58		90	
07533101	**Good identification**					**07533101**
					CAT	
Actinomyces naeslundii		92.31	0.89		25	
Propionibacterium propionicum		7.54	0.60		1	
Propionibacterium granulosum		0.11	0.24		90	
07533102	**Low discrimination**					**07533102**
					CAT	
Propionibacterium propionicum		60.70	0.79	NIT 86	1	
Actinomyces naeslundii		36.72	0.86		25	
Propionibacterium granulosum		1.64	0.48		90	
07533103	**Low discrimination**					**07533103**
					CAT	
Actinomyces naeslundii		89.05	0.89		25	
Propionibacterium propionicum		10.91	0.63		1	
Propionibacterium granulosum		0.03	0.13		90	
07533110	**Low discrimination**					**07533110**
					CAT	
Actinomyces naeslundii		59.28	0.64	MLZ 5	25	
Propionibacterium granulosum		27.35	0.43		90	
Propionibacterium propionicum		12.54	0.42		1	
07533111	**Very good identification**					**07533111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Actinomyces naeslundii		98.16	0.67	MLZ 5	25	
Propionibacterium propionicum		1.54	0.26		1	
Propionibacterium granulosum		0.31	0.09		90	
07533112	**Low discrimination**					**07533112**
Actinomyces naeslundii		57.94	0.64	MLZ 5		
Propionibacterium propionicum		18.38	0.45			
Bifidobacterium adolescentis		15.77	0.38			
07533113	**Very good identification**					**07533113**
Actinomyces naeslundii		97.70	0.67	MLZ 5		
Propionibacterium propionicum		2.30	0.29			
07533120	**Low discrimination**					**07533120**
					CAT	
Actinomyces naeslundii		73.10	0.86		25	
Clostridium perfringens		19.92	0.54		1	
Propionibacterium granulosum		3.25	0.48		90	
07533121	**Very good identification**					**07533121**
Actinomyces naeslundii		99.66	0.89			
Clostridium perfringens		0.18	0.19			
Propionibacterium propionicum		0.08	0.26			
07533122	**Low discrimination**					**07533122**
Actinomyces naeslundii		84.54	0.86			
Clostridium perfringens		10.84	0.48			
Propionibacterium propionicum		1.41	0.45			
07533123	**Very good identification**					**07533123**
Actinomyces naeslundii		99.79	0.89			
Propionibacterium propionicum		0.12	0.29			
Clostridium perfringens		0.09	0.14			
07533130	**Low discrimination**					**07533130**
					CAT	
Actinomyces naeslundii		84.32	0.64	MLZ 5	25	
Propionibacterium granulosum		9.73	0.33		90	
Clostridium perfringens		4.41	0.19		1	
07533131	**Very good identification**					**07533131**
Actinomyces naeslundii		100.0	0.67	MLZ 5		
07533132	**Low discrimination**					**07533132**
					GRA	
Actinomyces naeslundii		88.81	0.64	MLZ 5	99	
Parabacteroides distasonis		5.43	0.30		1	
Propionibacterium granulosum		2.56	0.22		99	
07533133	**Very good identification**					**07533133**
Actinomyces naeslundii		100.0	0.67	MLZ 5		
07533140	**Low discrimination**					**07533140**
Propionibacterium propionicum		85.84	0.76	NIT 86		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		13.75	0.73			
Actinomyces israelii		0.15	0.30			
07533141	**Low discrimination**					**07533141**
Actinomyces naeslundii		68.37	0.76			
Propionibacterium propionicum		31.63	0.60			
07533142	**Good identification**					**07533142**
Propionibacterium propionicum		90.08	0.79	NIT 86		
Actinomyces naeslundii		9.62	0.73			
Actinomyces israelii		0.16	0.33			
07533143	**Low discrimination**					**07533143**
Actinomyces naeslundii		59.04	0.76			
Propionibacterium propionicum		40.97	0.63			
07533150	**Low discrimination**					**07533150**
Propionibacterium propionicum		53.13	0.42	NIT 86		
Actinomyces naeslundii		44.33	0.51	MLZ 1		
Actinomyces israelii		1.38	0.16			
07533151	**Acceptable identification**					**07533151**
Actinomyces naeslundii		91.85	0.54	MLZ 5		
Propionibacterium propionicum		8.15	0.26			
07533152	**Low discrimination**					**07533152**
Propionibacterium propionicum		59.24	0.45	NIT 86		
Actinomyces naeslundii		32.95	0.51	MLZ 1		
Bifidobacterium adolescentis		6.28	0.23			
07533153	**Low discrimination**					**07533153**
Actinomyces naeslundii		88.25	0.54	MLZ 5		
Propionibacterium propionicum		11.75	0.29			
07533160	**Low discrimination**					**07533160**
Actinomyces naeslundii		73.43	0.73			
Clostridium perfringens		21.60	0.41			
Propionibacterium propionicum		4.63	0.42			
07533161	**Very good identification**					**07533161**
Actinomyces naeslundii		99.34	0.76			
Propionibacterium propionicum		0.46	0.26			
Clostridium perfringens		0.20	0.07			
07533162	**Low discrimination**					**07533162**
Actinomyces naeslundii		81.10	0.73			
Clostridium perfringens		11.23	0.36			
Propionibacterium propionicum		7.67	0.45			
07533163	**Very good identification**					**07533163**
Actinomyces naeslundii		99.21	0.76			
Propionibacterium propionicum		0.70	0.29			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		0.09	0.01			
07533170	**Acceptable identification**					**07533170**
Actinomyces naeslundii		94.66	0.51	MLZ 5		
Clostridium perfringens		5.34	0.07			
07533171	**Good identification**					**07533171**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
07533172	**Good identification**					**07533172**
Actinomyces naeslundii		97.41	0.51	MLZ 5		
Clostridium perfringens		2.59	0.01			
07533173	**Good identification**					**07533173**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
07533200	**Very good identification**					**07533200**
Propionibacterium granulosum		99.35	0.66	RHA 3		
Actinomyces naeslundii		0.65	0.44			
07533202	**Good identification**					**07533202**
Propionibacterium granulosum		97.46	0.56	RHA 3		
Actinomyces naeslundii		2.54	0.44			
07533203	**Low discrimination**					**07533203**
Actinomyces naeslundii		79.45	0.47	RHA 1		
Propionibacterium granulosum		20.55	0.22			
07533210	**Good identification**					**07533210**
Propionibacterium granulosum		100.0	0.51	RHA 3		
07533212	**Good identification**					**07533212**
Propionibacterium granulosum		100.0	0.41	RHA 3		
07533220	**Good identification**					**07533220**
					CAT	
Propionibacterium granulosum		97.20	0.56	RHA 3	90	
Actinomyces naeslundii		2.53	0.44		25	
Clostridium perfringens		0.27	0.05		1	
07533221	**Low discrimination**					**07533221**
Actinomyces naeslundii		79.45	0.47	RHA 1		
Propionibacterium granulosum		20.55	0.22			
07533222	**Acceptable identification**					**07533222**
Propionibacterium granulosum		90.57	0.46	RHA 3		
Actinomyces naeslundii		9.43	0.44			
07533223	**Acceptable identification**					**07533223**
Actinomyces naeslundii		93.93	0.47	RHA 1		
Propionibacterium granulosum		6.07	0.11			
07533230	**Good identification**					**07533230**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.41	RHA 3			
07533300	**Low discrimination**						**07533300**
					CAT		
Propionibacterium propionicum		41.06	0.42	NIT 86	1		
Actinomyces naeslundii		37.25	0.51	RHA 1	25		
Propionibacterium granulosum		20.31	0.32		90		
07533301	**Acceptable identification**						**07533301**
Actinomyces naeslundii		92.45	0.54	RHA 1			
Propionibacterium propionicum		7.55	0.26				
07533302	**Low discrimination**						**07533302**
					CAT		
Propionibacterium propionicum		58.11	0.45	NIT 86	1		
Actinomyces naeslundii		35.15	0.51	RHA 1	25		
Propionibacterium granulosum		4.79	0.22		90		
07533303	**Low discrimination**						**07533303**
Actinomyces naeslundii		89.09	0.54	RHA 1			
Propionibacterium propionicum		10.91	0.29				
07533320	**Low discrimination**						**07533320**
					SPO		
Actinomyces naeslundii		57.71	0.51	RHA 1	1		
Clostridium ramosum		16.60	0.17		92		
Clostridium perfringens		15.73	0.19		84		
07533321	**Good identification**						**07533321**
Actinomyces naeslundii		100.0	0.54	RHA 1			
07533322	**Low discrimination**						**07533322**
					SPO	GRA	
Parabacteroides distasonis		41.70	0.43	MAN 1	1	1	
Clostridium ramosum		27.00	0.27	XYL 90	92	50	
Actinomyces naeslundii		23.48	0.51		1	99	
07533323	**Good identification**						**07533323**
Actinomyces naeslundii		100.0	0.54	RHA 1			
07533332	**Acceptable identification**						**07533332**
					GRA		
Parabacteroides distasonis		94.78	0.38	MAN 1	1		
Actinomyces naeslundii		5.22	0.29	XYL 90	99		
07533340	**Low discrimination**						**07533340**
Propionibacterium propionicum		86.20	0.42	NIT 86			
Actinomyces naeslundii		13.80	0.38	RHA 1			
07533341	**Low discrimination**						**07533341**
Actinomyces naeslundii		68.37	0.41	RHA 1			
Propionibacterium propionicum		31.63	0.26				
07533342	**Low discrimination**						**07533342**
Propionibacterium propionicum		89.30	0.45	NIT 86			
Actinomyces naeslundii		9.53	0.38	RHA 1			
Actinomyces israelii		1.17	0.14				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07533343	**Low discrimination**					**07533343**
Actinomyces naeslundii		59.04	0.41	RHA 1		
Propionibacterium propionicum		40.97	0.29			
07533360	**Low discrimination**					**07533360**
Actinomyces naeslundii		77.27	0.38	RHA 1		
Clostridium perfringens		22.73	0.07			
07533361	**Good identification**					**07533361**
Actinomyces naeslundii		100.0	0.41	RHA 1		
07533362	**Low discrimination**					**07533362**
Actinomyces naeslundii		87.84	0.38	RHA 1		
Clostridium perfringens		12.16	0.01			
07533363	**Good identification**					**07533363**
Actinomyces naeslundii		100.0	0.41	RHA 1		
07533400	**Very good identification**					**07533400**
					CAT	
Propionibacterium granulosum		97.20	0.58	ARA 1	90	
Actinomyces naeslundii		1.94	0.44		25	
Actinomyces israelii		0.56	0.25		1	
07533401	**Low discrimination**					**07533401**
Actinomyces naeslundii		74.75	0.47	ARA 1		
Propionibacterium granulosum		25.26	0.24			
07533402	**Low discrimination**					**07533402**
					CAT	
Propionibacterium granulosum		87.37	0.48	ARA 1	90	
Actinomyces naeslundii		6.97	0.44		25	
Actinomyces israelii		2.99	0.28		1	
07533403	**Acceptable identification**					**07533403**
Actinomyces naeslundii		92.21	0.47	ARA 1		
Propionibacterium granulosum		7.79	0.13			
07533410	**Good identification**					**07533410**
Propionibacterium granulosum		100.0	0.43	ARA 1		
07533420	**Acceptable identification**					**07533420**
					CAT	
Propionibacterium granulosum		91.24	0.48	ARA 1	90	
Actinomyces naeslundii		7.27	0.44		25	
Clostridium perfringens		0.77	0.05		1	
07533421	**Acceptable identification**					**07533421**
Actinomyces naeslundii		92.21	0.47	ARA 1		
Propionibacterium granulosum		7.79	0.13			
07533422	**Low discrimination**					**07533422**
					CAT	
Propionibacterium granulosum		74.05	0.37	ARA 1	90	
Actinomyces naeslundii		23.61	0.44		25	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		2.34	0.09		1	
07533423	**Good identification**					**07533423**
Actinomyces naeslundii		100.0	0.47	ARA 1		
07533500	**Low discrimination**					**07533500**
Propionibacterium propionicum		29.67	0.42	NIT 86		
Actinomyces naeslundii		26.92	0.51	ARA 1		
Bifidobacterium longum subsp. longum		23.21	0.33			
07533501	**Acceptable identification**					**07533501**
Actinomyces naeslundii		92.45	0.54	ARA 1		
Propionibacterium propionicum		7.55	0.26			
07533502	**Low discrimination**					**07533502**
Bifidobacterium adolescentis		64.60	0.51	XYL 87		
Propionibacterium propionicum		13.75	0.45			
Actinomyces naeslundii		8.32	0.51			
07533503	**Low discrimination**					**07533503**
Actinomyces naeslundii		85.13	0.54	ARA 1		
Propionibacterium propionicum		10.42	0.29			
Bifidobacterium adolescentis		4.45	0.17			
07533510	**Low discrimination**					**07533510**
Bifidobacterium longum subsp. longum		86.30	0.37	MAN 1		
Bifidobacterium adolescentis		5.64	0.18			
Actinomyces israelii		3.54	0.21			
07533512	**Low discrimination**					**07533512**
Bifidobacterium adolescentis		85.49	0.53	XYL 87		
Bifidobacterium longum subsp. longum		13.22	0.37			
Actinomyces israelii		0.81	0.24			
07533520	**Low discrimination**					**07533520**
Bifidobacterium longum subsp. longum		70.98	0.43	MAN 1		
Actinomyces naeslundii		20.58	0.51			
Clostridium perfringens		5.61	0.19			
07533521	**Good identification**					**07533521**
Actinomyces naeslundii		97.73	0.54	ARA 1		
Bifidobacterium longum subsp. longum		2.27	0.09			
07533522	**Low discrimination**					**07533522**
					GRA	
Bifidobacterium longum subsp. longum		67.79	0.43	MAN 1	99	
Actinomyces naeslundii		19.66	0.51		99	
Parabacteroides distasonis		7.13	0.31		1	
07533523	**Good identification**					**07533523**
Actinomyces naeslundii		97.73	0.54	ARA 1		
Bifidobacterium longum subsp. longum		2.27	0.09			
07533530	**Good identification**					**07533530**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		99.11	0.47	MAN 1		
Actinomyces naeslundii		0.89	0.29			
07533532	**Acceptable identification**					**07533532**
					GRA	
Bifidobacterium longum subsp. longum		94.47	0.47	MAN 1	99	
Parabacteroides distasonis		3.14	0.26		1	
Bifidobacterium adolescentis		1.54	0.18		99	
07533540	**Low discrimination**					**07533540**
Propionibacterium propionicum		76.80	0.42	NIT 86		
Actinomyces naeslundii		12.30	0.38	ARA 1		
Actinomyces israelii		10.91	0.28			
07533541	**Low discrimination**					**07533541**
Actinomyces naeslundii		68.37	0.41	ARA 1		
Propionibacterium propionicum		31.63	0.26			
07533542	**Low discrimination**					**07533542**
Propionibacterium propionicum		54.66	0.45	NIT 86		
Bifidobacterium adolescentis		31.74	0.36	ARA 1		
Actinomyces israelii		7.77	0.31			
07533543	**Low discrimination**					**07533543**
Actinomyces naeslundii		59.04	0.41	ARA 1		
Propionibacterium propionicum		40.97	0.29			
07533552	**Good identification**					**07533552**
Bifidobacterium adolescentis		97.10	0.37	XYL 87		
Actinomyces israelii		2.90	0.17	SOR 11		
07533560	**Low discrimination**					**07533560**
Actinomyces naeslundii		67.05	0.38	ARA 1		
Clostridium perfringens		19.72	0.07			
Bifidobacterium longum subsp. longum		13.23	0.09			
07533561	**Good identification**					**07533561**
Actinomyces naeslundii		100.0	0.41	ARA 1		
07533562	**Low discrimination**					**07533562**
Actinomyces naeslundii		74.86	0.38	ARA 1		
Bifidobacterium longum subsp. longum		14.78	0.09			
Clostridium perfringens		10.36	0.01			
07533563	**Good identification**					**07533563**
Actinomyces naeslundii		100.0	0.41	ARA 1		
07533722	**Acceptable identification**					**07533722**
					GRA	
Parabacteroides distasonis		94.97	0.39	MAN 1	1	
Bifidobacterium longum subsp. longum		3.04	0.09	XYL 90	99	
Clostridium clostridioforme		1.99	0.05		50	
07534000	**Good identification**					**07534000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Propionibacterium granulosum		94.81	0.60	XYL 1	99	90		
Fusobacterium mortiferum		3.67	0.39		1	5		
Prevotella intermedia		0.67	0.15		1	1		
07534002	**Good identification**							**07534002**
					GRA	CAT		
Propionibacterium granulosum		96.51	0.50	XYL 1	99	90		
Fusobacterium mortiferum		1.66	0.23		1	5		
Actinomyces viscosus		1.28	0.08		99	76		
07534010	**Good identification**							**07534010**
Propionibacterium granulosum		100.0	0.45	XYL 1				
07534020	**Good identification**							**07534020**
					CAT			
Propionibacterium granulosum		99.37	0.50	XYL 1	90			
Clostridium perfringens		0.63	0.05		1			
07534022	**Good identification**							**07534022**
					GRA			
Propionibacterium granulosum		96.45	0.39	XYL 1	99			
Parabacteroides distasonis		3.55	0.17		1			
07534100	**Low discrimination**							**07534100**
					COC	GRA	CAT	
Fusobacterium mortiferum		79.76	0.45	MLT 11	1	1	5	
Propionibacterium granulosum		8.92	0.26	XYL 7	1	99	90	
Atopobium parvulum		7.48	0.16		99	99	1	
07534122	**Acceptable identification**							**07534122**
					GRA			
Parabacteroides distasonis		92.28	0.51	GAL 99	1			
Clostridium perfringens		2.22	0.14		99			
Bifidobacterium longum subsp. longum		1.81	0.17		99			
07534132	**Low discrimination**							**07534132**
					GRA			
Parabacteroides distasonis		85.67	0.46	GAL 99	1			
Clostridium clostridioforme		9.02	0.24		50			
Bifidobacterium longum subsp. longum		5.31	0.21		99			
07534312	**Good identification**							**07534312**
Clostridium clostridioforme		96.23	0.39	SAL 88				
Parabacteroides distasonis		3.78	0.20	ARA 94				
07534322	**Very good identification**							**07534322**
Parabacteroides distasonis		96.92	0.59	GAL 99				
Clostridium clostridioforme		2.95	0.27					
Prevotella oralis		0.13	-.03					
07534332	**Low discrimination**							**07534332**
Parabacteroides distasonis		79.53	0.54	GAL 99				
Clostridium clostridioforme		20.48	0.39					
07534412	**Good identification**							**07534412**
Clostridium clostridioforme		96.95	0.35	SAL 88				
Bifidobacterium longum subsp. longum		3.05	0.11	RHA 88				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07534432	**Low discrimination**					**07534432**
Clostridium clostridioforme		88.82	0.35	SAL 88		
Bifidobacterium longum subsp. longum		11.18	0.21	RHA 88		
07534500	**Good identification**					**07534500**
					GRA	
Bifidobacterium longum subsp. longum		97.17	0.41	GAL 99	99	
Lactobacillus fermentum		2.03	0.20		99	
Fusobacterium mortiferum		0.80	0.11		1	
07534502	**Low discrimination**					**07534502**
Bifidobacterium longum subsp. longum		59.40	0.41	GAL 99		
Clostridium clostridioforme		23.87	0.33			
Bifidobacterium adolescentis		14.74	0.33			
07534510	**Good identification**					**07534510**
Bifidobacterium longum subsp. longum		98.93	0.45	GAL 99		
Clostridium clostridioforme		1.07	0.10			
07534512	**Low discrimination**					**07534512**
Clostridium clostridioforme		47.72	0.44	SAL 88		
Bifidobacterium longum subsp. longum		44.38	0.45	RHA 88		
Bifidobacterium adolescentis		7.90	0.34			
07534520	**Acceptable identification**					**07534520**
					GRA	
Bifidobacterium longum subsp. longum		91.32	0.52	GAL 99	99	
Bacteroides vulgatus		8.38	0.20		1	
Parabacteroides distasonis		0.29	0.13		1	
07534522	**Low discrimination**					**07534522**
					GRA	
Bifidobacterium longum subsp. longum		70.61	0.52	GAL 99	99	
Parabacteroides distasonis		22.29	0.47		1	
Clostridium clostridioforme		7.09	0.33		50	
07534530	**Good identification**					**07534530**
Bifidobacterium longum subsp. longum		99.73	0.56	GAL 99		
Clostridium clostridioforme		0.27	0.10			
07534532	**Low discrimination**					**07534532**
					GRA	
Bifidobacterium longum subsp. longum		73.07	0.56	GAL 99	99	
Clostridium clostridioforme		19.64	0.44		50	
Parabacteroides distasonis		7.29	0.43		1	
07534602	**Good identification**					**07534602**
Clostridium clostridioforme		100.0	0.39	SAL 88		
07534612	**Good identification**					**07534612**
Clostridium clostridioforme		100.0	0.50	SAL 88		
07534622	**Good identification**					**07534622**
Clostridium clostridioforme		95.83	0.39	SAL 88		
Parabacteroides distasonis		4.17	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07534632	**Good identification**						**07534632**
Clostridium clostridioforme		99.49	0.50	SAL 88			
Parabacteroides distasonis		0.51	0.17				
07534702	**Good identification**						**07534702**
					GRA		
Clostridium clostridioforme		98.39	0.48	SAL 88	50		
Parabacteroides distasonis		1.28	0.21		1		
Bifidobacterium longum subsp. longum		0.34	0.07		99		
07534712	**Very good identification**						**07534712**
					GRA		
Clostridium clostridioforme		99.72	0.59	SAL 88	50		
Parabacteroides distasonis		0.15	0.17		1		
Bifidobacterium longum subsp. longum		0.13	0.11		99		
07534720	**Good identification**						**07534720**
					GRA		
Bacteroides vulgatus		96.82	0.36	TRE 4	1		
Bifidobacterium longum subsp. longum		1.18	0.17	GAL 99	99		
Parabacteroides distasonis		1.12	0.21		1		
07534722	**Low discrimination**						**07534722**
Parabacteroides distasonis		55.19	0.55	GAL 99			
Clostridium clostridioforme		42.93	0.48				
Bacteroides thetaiotaomicron		0.81	0.08				
07534732	**Low discrimination**						**07534732**
					GRA		
Clostridium clostridioforme		86.31	0.59	SAL 88	50		
Parabacteroides distasonis		13.11	0.51		1		
Bifidobacterium longum subsp. longum		0.44	0.21		99		
07535000	**Low discrimination**						**07535000**
					GRA	CAT	
Propionibacterium granulosum		80.55	0.60	XYL 1	99	90	
Lactobacillus fermentum		11.28	0.50		99	1	
Fusobacterium mortiferum		4.68	0.42		1	5	
07535001	**Low discrimination**						**07535001**
					CAT		
Actinomyces naeslundii		50.22	0.42	XYL 1	25		
Propionibacterium granulosum		43.66	0.26		90		
Lactobacillus fermentum		6.11	0.16		1		
07535002	**Low discrimination**						**07535002**
					CAT		
Propionibacterium granulosum		57.32	0.50	XYL 1	90		
Lactobacillus fermentum		32.10	0.50		1		
Actinomyces israelii		3.67	0.35		1		
07535003	**Low discrimination**						**07535003**
					CAT		
Actinomyces naeslundii		74.68	0.42	XYL 1	25		
Propionibacterium granulosum		16.23	0.15		90		
Lactobacillus fermentum		9.09	0.16		1		
07535010	**Good identification**						**07535010**
					CAT		
Propionibacterium granulosum		96.52	0.45	XYL 1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		1.35	0.11	MLZ 12	1		
Actinomyces israelii		1.13	0.18		1		
07535020	**Low discrimination**						**07535020**
					CAT		
Propionibacterium granulosum		52.66	0.50	XYL 1	90		
Clostridium perfringens		33.16	0.39		1		
Lactobacillus fermentum		9.83	0.42		1		
07535021	**Low discrimination**						**07535021**
					CAT		
Actinomyces naeslundii		73.84	0.42	XYL 1	25		
Propionibacterium granulosum		16.05	0.15		90		
Clostridium perfringens		10.11	0.05		1		
07535022	**Low discrimination**						**07535022**
					GRA	CAT	
Parabacteroides distasonis		51.40	0.51	LAC 99	1	76	
Clostridium perfringens		16.70	0.33		99	1	
Propionibacterium granulosum		14.09	0.39		99	90	
07535023	**Low discrimination**						**07535023**
					GRA		
Actinomyces naeslundii		83.46	0.42	XYL 1	99		
Parabacteroides distasonis		16.55	0.17		1		
07535032	**Low discrimination**						**07535032**
					GRA		
Parabacteroides distasonis		81.58	0.46	LAC 99	1		
Clostridium clostridioforme		7.70	0.23		50		
Propionibacterium granulosum		5.66	0.25		99		
07535100	**Low discrimination**						**07535100**
					COC	GRA	
Atopobium parvulum		58.29	0.50	RAF 1	99	99	
Lactobacillus fermentum		15.89	0.54		1	99	
Fusobacterium mortiferum		9.42	0.48		1	1	
07535101	**Low discrimination**						**07535101**
					COC	SPO	
Actinomyces naeslundii		66.58	0.49	XYL 1	1	1	
Atopobium parvulum		17.05	0.16		99	1	
Clostridium chauvoei		6.73	0.09		1	99	
07535102	**Low discrimination**						**07535102**
					GRA		
Bifidobacterium adolescentis		31.96	0.53	ARA 87	99		
Lactobacillus fermentum		17.98	0.54		99		
Parabacteroides distasonis		17.94	0.51		1		
07535103	**Low discrimination**						**07535103**
Actinomyces naeslundii		72.52	0.49	XYL 1			
Bifidobacterium adolescentis		9.00	0.18				
Lactobacillus fermentum		5.06	0.20				
07535110	**Low discrimination**						**07535110**
					COC		
Bifidobacterium longum subsp. longum		87.34	0.45	ARA 99	1		
Atopobium parvulum		3.92	0.16		99		
Actinomyces israelii		2.96	0.28		1		
07535112	**Low discrimination**						**07535112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium adolescentis		57.66	0.54	ARA 87	99	
Bifidobacterium longum subsp. longum		21.89	0.45		99	
Parabacteroides distasonis		14.26	0.46		1	
07535120	**Low discrimination**					**07535120**
					GRA	
Clostridium perfringens		45.01	0.54	XYL 1	99	
Bifidobacterium longum subsp. longum		17.26	0.52		99	
Bacteroides vulgatus		11.81	0.35		1	
07535121	**Low discrimination**					**07535121**
Actinomyces naeslundii		56.30	0.49	XYL 1		
Clostridium perfringens		19.86	0.19			
Bifidobacterium longum subsp. longum		7.61	0.17			
07535122	**Good identification**					**07535122**
					GRA	
Parabacteroides distasonis		92.97	0.85		1	
Clostridium perfringens		2.24	0.48		99	
Bifidobacterium longum subsp. longum		1.82	0.52		99	
07535123	**Low discrimination**					**07535123**
					GRA	
Parabacteroides distasonis		82.49	0.51	URE 1	1	
Actinomyces naeslundii		11.96	0.49		99	
Clostridium perfringens		1.99	0.14		99	
07535130	**Low discrimination**					**07535130**
					GRA	
Bifidobacterium longum subsp. longum		84.18	0.56	ARA 99	99	
Parabacteroides distasonis		13.71	0.46		1	
Clostridium perfringens		1.30	0.19		99	
07535132	**Good identification**					**07535132**
					GRA	
Parabacteroides distasonis		93.74	0.80		1	
Bifidobacterium longum subsp. longum		5.81	0.56		99	
Clostridium clostridioforme		0.30	0.32		50	
07535133	**Acceptable identification**					**07535133**
					GRA	
Parabacteroides distasonis		92.92	0.46	URE 1	1	
Bifidobacterium longum subsp. longum		5.76	0.21		99	
Actinomyces naeslundii		1.32	0.28		99	
07535140	**Low discrimination**					**07535140**
					COC	
Actinomyces israelii		40.96	0.35	SAL 92	1	
Atopobium parvulum		20.87	0.16	CEL 93	99	
Clostridium perfringens		16.77	0.15		1	
07535141	**Good identification**					**07535141**
Actinomyces naeslundii		100.0	0.37	XYL 1		
07535142	**Low discrimination**					**07535142**
Bifidobacterium adolescentis		56.13	0.37	ARA 87		
Actinomyces israelii		27.87	0.38	SOR 11		
Actinomyces naeslundii		4.35	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07535143	**Good identification**					**07535143**
Actinomyces naeslundii		100.0	0.37	XYL 1		
07535152	**Low discrimination**					**07535152**
Bifidobacterium adolescentis		89.93	0.39	ARA 87		
Actinomyces israelii		5.46	0.24	SOR 11		
Bifidobacterium longum subsp. longum		2.79	0.11			
07535160	**Acceptable identification**					**07535160**
Clostridium perfringens		92.50	0.41	XYL 1		
Bifidobacterium longum subsp. longum		1.88	0.17			
Actinomyces naeslundii		1.64	0.34			
07535161	**Low discrimination**					**07535161**
Actinomyces naeslundii		72.43	0.37	XYL 1		
Clostridium perfringens		27.57	0.07			
07535162	**Low discrimination**					**07535162**
					GRA	
Parabacteroides distasonis		65.42	0.51	SOR 1	1	
Clostridium perfringens		29.69	0.36		99	
Bifidobacterium longum subsp. longum		1.28	0.17		99	
07535163	**Low discrimination**					**07535163**
					GRA	
Actinomyces naeslundii		63.54	0.37	XYL 1	99	
Parabacteroides distasonis		25.08	0.17		1	
Clostridium perfringens		11.38	0.01		99	
07535172	**Acceptable identification**					**07535172**
					GRA	
Parabacteroides distasonis		93.41	0.46	SOR 1	1	
Bifidobacterium longum subsp. longum		5.79	0.21		99	
Clostridium perfringens		0.80	0.01		99	
07535212	**Acceptable identification**					**07535212**
					GRA	
Clostridium clostridioforme		94.68	0.38	SAL 88	50	
Parabacteroides distasonis		4.15	0.20	ARA 94	1	
Propionibacterium granulosum		1.18	0.09		99	
07535222	**Very good identification**					**07535222**
Parabacteroides distasonis		96.30	0.59	LAC 99		
Clostridium clostridioforme		2.63	0.27			
Prevotella oralis		0.81	0.11			
07535232	**Low discrimination**					**07535232**
Parabacteroides distasonis		81.25	0.54	LAC 99		
Clostridium clostridioforme		18.75	0.38			
07535302	**Low discrimination**					**07535302**
					GRA	
Parabacteroides distasonis		89.41	0.59	NAG 99	1	
Clostridium clostridioforme		8.16	0.36		50	
Actinomyces israelii		0.63	0.26		99	
07535312	**Low discrimination**					**07535312**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		55.83	0.54	NAG 99	1	
Clostridium clostridioforme		43.12	0.47		50	
Bifidobacterium adolescentis		0.76	0.20		99	
07535320	**Low discrimination**					**07535320**
					GRA	
Bacteroides vulgatus		79.38	0.51	TRE 4	1	
Parabacteroides distasonis		19.92	0.59	ARA 93	1	
Clostridium perfringens		0.34	0.19		99	
07535322	**Excellent identification**					**07535322**
Parabacteroides distasonis		99.71	0.93			
Prevotella oralis		0.13	0.31			
Clostridium clostridioforme		0.09	0.36			
07535323	**Very good identification**					**07535323**
Parabacteroides distasonis		99.87	0.59	URE 1		
Prevotella oralis		0.13	-.03			
07535330	**Acceptable identification**					**07535330**
					GRA	
Parabacteroides distasonis		90.64	0.54	BGL 99	1	
Bacteroides vulgatus		6.78	0.17		1	
Bifidobacterium longum subsp. longum		1.87	0.21		99	
07535332	**Very good identification**					**07535332**
					GRA	
Parabacteroides distasonis		99.20	0.89		1	
Clostridium clostridioforme		0.77	0.47		50	
Bifidobacterium longum subsp. longum		0.02	0.21		99	
07535333	**Good identification**					**07535333**
Parabacteroides distasonis		99.23	0.54	URE 1		
Clostridium clostridioforme		0.77	0.13			
07535362	**Very good identification**					**07535362**
					GRA	
Parabacteroides distasonis		99.72	0.59	SOR 1	1	
Clostridium perfringens		0.15	0.01		99	
Prevotella oralis		0.13	-.03		1	
07535372	**Good identification**					**07535372**
Parabacteroides distasonis		99.23	0.54	SOR 1		
Clostridium clostridioforme		0.77	0.13			
07535400	**Low discrimination**					**07535400**
					CAT	
Lactobacillus fermentum		51.75	0.50	TRE 1	1	
Bifidobacterium longum subsp. longum		40.91	0.41		1	
Propionibacterium granulosum		3.73	0.26		90	
07535402	**Low discrimination**					**07535402**
Lactobacillus fermentum		41.79	0.50	TRE 1		
Bifidobacterium longum subsp. longum		33.03	0.41			
Clostridium clostridioforme		11.90	0.32			
07535410	**Good identification**					**07535410**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		96.95	0.45	LAC 99		
Clostridium clostridioforme		0.94	0.09			
Lactobacillus fermentum		0.73	0.16			
07535412	**Low discrimination**					**07535412**
Bifidobacterium longum subsp. longum		46.31	0.45	LAC 99		
Clostridium clostridioforme		44.62	0.43			
Bifidobacterium adolescentis		8.25	0.34			
07535420	**Low discrimination**					**07535420**
					GRA	
Bifidobacterium longum subsp. longum		82.66	0.52	LAC 99	99	
Lactobacillus fermentum		8.71	0.42		99	
Bacteroides vulgatus		7.59	0.20		1	
07535422	**Low discrimination**					**07535422**
					GRA	
Bifidobacterium longum subsp. longum		66.08	0.52	LAC 99	99	
Parabacteroides distasonis		20.86	0.47		1	
Lactobacillus fermentum		6.97	0.42		99	
07535430	**Good identification**					**07535430**
Bifidobacterium longum subsp. longum		99.76	0.56	LAC 99		
Clostridium clostridioforme		0.24	0.09			
07535432	**Low discrimination**					**07535432**
					GRA	
Bifidobacterium longum subsp. longum		74.59	0.56	LAC 99	99	
Clostridium clostridioforme		17.97	0.43		50	
Parabacteroides distasonis		7.45	0.43		1	
07535500	**Very good identification**					**07535500**
					GRA	
Bifidobacterium longum subsp. longum		96.75	0.75		99	
Lactobacillus fermentum		2.02	0.54		99	
Bacteroides vulgatus		0.36	0.20		1	
07535501	**Good identification**					**07535501**
Bifidobacterium longum subsp. longum		97.96	0.41	URE 1		
Lactobacillus fermentum		2.04	0.20			
07535502	**Low discrimination**					**07535502**
Bifidobacterium longum subsp. longum		76.95	0.75			
Bifidobacterium adolescentis		19.09	0.67			
Lactobacillus fermentum		1.60	0.54			
07535503	**Low discrimination**					**07535503**
Bifidobacterium longum subsp. longum		77.29	0.41	URE 1		
Bifidobacterium adolescentis		19.17	0.33			
Lactobacillus fermentum		1.61	0.20			
07535510	**Very good identification**					**07535510**
Bifidobacterium longum subsp. longum		99.74	0.79			
Bifidobacterium adolescentis		0.18	0.34			
Actinomyces israelii		0.03	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07535511	**Good identification**					**07535511**
Bifidobacterium longum subsp. longum		100.0	0.45	URE 1		
07535512	**Low discrimination**					**07535512**
Bifidobacterium longum subsp. longum		82.29	0.79			
Bifidobacterium adolescentis		14.65	0.68			
Clostridium clostridioforme		2.68	0.52			
07535513	**Low discrimination**					**07535513**
Bifidobacterium longum subsp. longum		82.60	0.45	URE 1		
Bifidobacterium adolescentis		14.71	0.34			
Clostridium clostridioforme		2.69	0.18			
07535520	**Good identification**					**07535520**
					GRA	
Bifidobacterium longum subsp. longum		91.14	0.86		99	
Bacteroides vulgatus		8.37	0.54		1	
Parabacteroides distasonis		0.29	0.47		1	
07535521	**Acceptable identification**					**07535521**
					GRA	
Bifidobacterium longum subsp. longum		91.32	0.52	URE 1	99	
Bacteroides vulgatus		8.38	0.20		1	
Parabacteroides distasonis		0.29	0.13		1	
07535522	**Low discrimination**					**07535522**
					GRA	
Bifidobacterium longum subsp. longum		75.56	0.86		99	
Parabacteroides distasonis		23.85	0.81		1	
Clostridium clostridioforme		0.23	0.41		50	
07535523	**Low discrimination**					**07535523**
					GRA	
Bifidobacterium longum subsp. longum		75.83	0.52	URE 1	99	
Parabacteroides distasonis		23.94	0.47		1	
Clostridium clostridioforme		0.23	0.07		50	
07535530	**Very good identification**					**07535530**
					GRA	
Bifidobacterium longum subsp. longum		99.84	0.90		99	
Parabacteroides distasonis		0.10	0.43		1	
Bacteroides vulgatus		0.05	0.20		1	
07535531	**Good identification**					**07535531**
Bifidobacterium longum subsp. longum		100.0	0.56	URE 1		
07535532	**Good identification**					**07535532**
					GRA	
Bifidobacterium longum subsp. longum		90.20	0.90		99	
Parabacteroides distasonis		9.01	0.77		1	
Clostridium clostridioforme		0.74	0.52		50	
07535533	**Acceptable identification**					**07535533**
					GRA	
Bifidobacterium longum subsp. longum		90.26	0.56	URE 1	99	
Parabacteroides distasonis		9.01	0.43		1	
Clostridium clostridioforme		0.74	0.18		50	
07535540	**Low discrimination**					**07535540**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		83.79	0.41	SOR 1		
Actinomyces israelii		10.29	0.34			
Bifidobacterium adolescentis		2.57	0.17			
07535542	**Low discrimination**					**07535542**
Bifidobacterium adolescentis		71.16	0.51	SOR 11		
Bifidobacterium longum subsp. longum		23.44	0.41			
Actinomyces israelii		4.32	0.37			
07535550	**Good identification**					**07535550**
Bifidobacterium longum subsp. longum		96.83	0.45	SOR 1		
Bifidobacterium adolescentis		2.13	0.19			
Actinomyces israelii		1.04	0.20			
07535552	**Low discrimination**					**07535552**
Bifidobacterium adolescentis		67.51	0.53	SOR 11		
Bifidobacterium longum subsp. longum		30.98	0.45			
Clostridium clostridioforme		1.01	0.18			
07535560	**Acceptable identification**					**07535560**
					GRA	
Bifidobacterium longum subsp. longum		90.75	0.52	SOR 1	99	
Bacteroides vulgatus		8.33	0.20		1	
Clostridium perfringens		0.46	0.07		99	
07535562	**Low discrimination**					**07535562**
					GRA	
Bifidobacterium longum subsp. longum		75.09	0.52	SOR 1	99	
Parabacteroides distasonis		23.71	0.47		1	
Bifidobacterium adolescentis		0.58	0.17		99	
07535570	**Good identification**					**07535570**
Bifidobacterium longum subsp. longum		100.0	0.56	SOR 1		
07535572	**Low discrimination**					**07535572**
					GRA	
Bifidobacterium longum subsp. longum		89.81	0.56	SOR 1	99	
Parabacteroides distasonis		8.97	0.43		1	
Clostridium clostridioforme		0.73	0.18		50	
07535602	**Good identification**					**07535602**
					GRA	
Clostridium clostridioforme		97.42	0.47	SAL 88	50	
Parabacteroides distasonis		1.41	0.21		1	
Lactobacillus fermentum		0.47	0.16		99	
07535612	**Very good identification**					**07535612**
					GRA	
Clostridium clostridioforme		99.69	0.58	SAL 88	50	
Parabacteroides distasonis		0.17	0.17		1	
Bifidobacterium longum subsp. longum		0.14	0.11		99	
07535620	**Good identification**					**07535620**
					GRA	
Bacteroides vulgatus		96.91	0.36	TRE 4	1	
Bifidobacterium longum subsp. longum		1.19	0.17	LAC 99	99	
Parabacteroides distasonis		1.12	0.21		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07535622	**Low discrimination**					**07535622**
Parabacteroides distasonis		57.27	0.55	LAC 99		
Clostridium clostridioforme		39.92	0.47			
Bacteroides thetaiotaomicron		1.70	0.13			
07535632	**Low discrimination**					**07535632**
					GRA	
Clostridium clostridioforme		84.82	0.58	SAL 88	50	
Parabacteroides distasonis		14.38	0.51		1	
Bifidobacterium longum subsp. longum		0.49	0.21		99	
07535700	**Low discrimination**					**07535700**
					GRA	
Bacteroides vulgatus		73.82	0.36	TRE 4	1	
Bifidobacterium longum subsp. longum		22.34	0.41	NAG 99	99	
Clostridium clostridioforme		1.99	0.22		50	
07535702	**Low discrimination**					**07535702**
					GRA	
Clostridium clostridioforme		63.05	0.56	SAL 88	50	
Parabacteroides distasonis		27.02	0.55		1	
Bifidobacterium longum subsp. longum		7.13	0.41		99	
07535710	**Low discrimination**					**07535710**
					GRA	
Bifidobacterium longum subsp. longum		78.70	0.45	RHA 1	99	
Clostridium clostridioforme		18.80	0.33		50	
Bacteroides vulgatus		1.54	0.02		1	
07535712	**Good identification**					**07535712**
					GRA	
Clostridium clostridioforme		90.83	0.67	SAL 88	50	
Parabacteroides distasonis		4.60	0.51		1	
Bifidobacterium longum subsp. longum		3.84	0.45		99	
07535720	**Very good identification**					**07535720**
					GRA	
Bacteroides vulgatus		97.63	0.70	TRE 4	1	
Bifidobacterium longum subsp. longum		1.19	0.52		99	
Parabacteroides distasonis		1.13	0.55		1	
07535721	**Good identification**					**07535721**
					GRA	
Bacteroides vulgatus		97.68	0.36	TRE 4	1	
Bifidobacterium longum subsp. longum		1.19	0.17	URE 1	99	
Parabacteroides distasonis		1.13	0.21		1	
07535722	**Good identification**					**07535722**
Parabacteroides distasonis		94.53	0.90			
Clostridium clostridioforme		2.23	0.56			
Bacteroides thetaiotaomicron		1.39	0.42			
07535723	**Acceptable identification**					**07535723**
Parabacteroides distasonis		94.54	0.55	URE 1		
Clostridium clostridioforme		2.23	0.22			
Bacteroides thetaiotaomicron		1.39	0.08			
07535730	**Low discrimination**					**07535730**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		54.12	0.56	RHA 1	99	
Bacteroides vulgatus		26.26	0.36		1	
Parabacteroides distasonis		16.21	0.51		1	
07535732	**Low discrimination**					**07535732**
					GRA	
Parabacteroides distasonis		80.38	0.85		1	
Clostridium clostridioforme		16.03	0.67		50	
Bifidobacterium longum subsp. longum		2.71	0.56		99	
07535733	**Low discrimination**					**07535733**
					GRA	
Parabacteroides distasonis		80.40	0.51	URE 1	1	
Clostridium clostridioforme		16.03	0.33		50	
Bifidobacterium longum subsp. longum		2.71	0.21		99	
07535760	**Good identification**					**07535760**
					GRA	
Bacteroides vulgatus		97.68	0.36	TRE 4	1	
Bifidobacterium longum subsp. longum		1.19	0.17	SOR 1	99	
Parabacteroides distasonis		1.13	0.21		1	
07535762	**Acceptable identification**					**07535762**
Parabacteroides distasonis		94.54	0.55	SOR 1		
Clostridium clostridioforme		2.23	0.22			
Bacteroides thetaiotaomicron		1.39	0.08			
07535772	**Low discrimination**					**07535772**
					GRA	
Parabacteroides distasonis		80.40	0.51	SOR 1	1	
Clostridium clostridioforme		16.03	0.33		50	
Bifidobacterium longum subsp. longum		2.71	0.21		99	
07536000	**Very good identification**					**07536000**
Propionibacterium granulosum		100.0	0.58	XYL 1		
07536002	**Good identification**					**07536002**
					CAT	
Propionibacterium granulosum		99.52	0.48	XYL 1	90	
Actinomyces israelii		0.48	0.14		1	
07536010	**Good identification**					**07536010**
Propionibacterium granulosum		100.0	0.43	XYL 1		
07536020	**Good identification**					**07536020**
Propionibacterium granulosum		100.0	0.48	XYL 1		
07536022	**Good identification**					**07536022**
Propionibacterium granulosum		100.0	0.37	XYL 1		
07536100	**Low discrimination**					**07536100**
					CAT	
Propionibacterium propionicum		78.70	0.42	NIT 86	1	
Propionibacterium granulosum		12.72	0.24	XYL 1	90	
Actinomyces israelii		4.08	0.21		1	
07536102	**Low discrimination**					**07536102**
Propionibacterium propionicum		86.85	0.45	NIT 86		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		4.50	0.24	XYL 1		
Bifidobacterium adolescentis		4.12	0.17			
07536140	**Good identification**					**07536140**
Propionibacterium propionicum		98.02	0.42	NIT 86		
Actinomyces israelii		1.98	0.14	XYL 1		
07536142	**Good identification**					**07536142**
Propionibacterium propionicum		98.02	0.45	NIT 86		
Actinomyces israelii		1.98	0.17	XYL 1		
07537000	**Good identification**					**07537000**
					CAT	
Propionibacterium granulosum		90.74	0.58	XYL 1	90	
Actinomyces israelii		7.28	0.45		1	
Actinomyces naeslundii		1.81	0.44		25	
07537001	**Low discrimination**					**07537001**
Actinomyces naeslundii		74.75	0.47	XYL 1		
Propionibacterium granulosum		25.26	0.24			
07537002	**Low discrimination**					**07537002**
					CAT	
Propionibacterium granulosum		63.03	0.48	XYL 1	90	
Actinomyces israelii		30.35	0.48		1	
Actinomyces naeslundii		5.03	0.44		25	
07537003	**Low discrimination**					**07537003**
					CAT	
Actinomyces naeslundii		88.88	0.47	XYL 1	25	
Propionibacterium granulosum		7.51	0.13		90	
Actinomyces israelii		3.62	0.14		1	
07537010	**Acceptable identification**					**07537010**
					CAT	
Propionibacterium granulosum		91.92	0.43	XYL 1	90	
Actinomyces israelii		8.08	0.31	MLZ 12	1	
07537020	**Acceptable identification**					**07537020**
					CAT	
Propionibacterium granulosum		90.20	0.48	XYL 1	90	
Actinomyces naeslundii		7.19	0.44		25	
Actinomyces israelii		1.85	0.24		1	
07537021	**Acceptable identification**					**07537021**
Actinomyces naeslundii		92.21	0.47	XYL 1		
Propionibacterium granulosum		7.79	0.13			
07537022	**Low discrimination**					**07537022**
					CAT	
Propionibacterium granulosum		67.08	0.37	XYL 1	90	
Actinomyces naeslundii		21.39	0.44		25	
Actinomyces israelii		8.25	0.27		1	
07537023	**Good identification**					**07537023**
Actinomyces naeslundii		100.0	0.47	XYL 1		
07537040	**Low discrimination**					**07537040**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		69.62	0.38	SAL 92	1	
Propionibacterium granulosum		22.54	0.24	CEL 93	90	
Actinomyces naeslundii		7.85	0.31		25	
07537042	**Low discrimination**					**07537042**
					CAT	
Actinomyces israelii		88.57	0.41	SAL 92	1	
Actinomyces naeslundii		6.66	0.31	CEL 93	25	
Propionibacterium granulosum		4.78	0.13		90	
07537100	**Low discrimination**					**07537100**
Actinomyces israelii		67.95	0.55	SAL 92		
Propionibacterium propionicum		13.24	0.42	CEL 93		
Actinomyces naeslundii		12.01	0.51			
07537101	**Low discrimination**					**07537101**
Actinomyces naeslundii		89.31	0.54	XYL 1		
Propionibacterium propionicum		7.29	0.26			
Actinomyces israelii		3.40	0.21			
07537102	**Low discrimination**					**07537102**
Actinomyces israelii		44.39	0.58	SAL 92		
Bifidobacterium adolescentis		40.62	0.51	CEL 93		
Propionibacterium propionicum		8.65	0.45			
07537103	**Low discrimination**					**07537103**
Actinomyces naeslundii		81.18	0.54	XYL 1		
Propionibacterium propionicum		9.94	0.29			
Actinomyces israelii		4.64	0.24			
07537110	**Low discrimination**					**07537110**
Actinomyces israelii		79.56	0.41	SAL 92		
Bifidobacterium adolescentis		9.02	0.18	CEL 93		
Actinomyces naeslundii		4.95	0.29	MLZ 13		
07537112	**Low discrimination**					**07537112**
Bifidobacterium adolescentis		87.19	0.53	ARA 87		
Actinomyces israelii		11.65	0.44			
Actinomyces naeslundii		0.48	0.29			
07537120	**Low discrimination**					**07537120**
Actinomyces naeslundii		50.52	0.51	XYL 1		
Actinomyces israelii		18.25	0.35			
Clostridium perfringens		13.77	0.19			
07537121	**Good identification**					**07537121**
Actinomyces naeslundii		100.0	0.54	XYL 1		
07537122	**Low discrimination**					**07537122**
					GRA	
Parabacteroides distasonis		72.93	0.51	MAN 1	1	
Actinomyces naeslundii		13.69	0.51		99	
Actinomyces israelii		7.42	0.38		99	
07537123	**Good identification**					**07537123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		96.54	0.54	XYL 1	99	
Parabacteroides distasonis		3.46	0.17		1	
07537132	**Low discrimination**					**07537132**
					GRA	
Parabacteroides distasonis		87.57	0.46	MAN 1	1	
Bifidobacterium longum subsp. longum		5.43	0.21		99	
Bifidobacterium adolescentis		2.93	0.18		99	
07537140	**Low discrimination**					**07537140**
Actinomyces israelii		63.25	0.48	SAL 92		
Propionibacterium propionicum		31.68	0.42	CEL 93		
Actinomyces naeslundii		5.07	0.38			
07537141	**Low discrimination**					**07537141**
Actinomyces naeslundii		64.66	0.41	XYL 1		
Propionibacterium propionicum		29.91	0.26			
Actinomyces israelii		5.43	0.14			
07537142	**Low discrimination**					**07537142**
Actinomyces israelii		54.20	0.51	SAL 92		
Propionibacterium propionicum		27.15	0.45	CEL 93		
Bifidobacterium adolescentis		15.76	0.36			
07537143	**Low discrimination**					**07537143**
Actinomyces naeslundii		54.95	0.41	XYL 1		
Propionibacterium propionicum		38.13	0.29			
Actinomyces israelii		6.92	0.17			
07537152	**Low discrimination**					**07537152**
Bifidobacterium adolescentis		70.40	0.37	ARA 87		
Actinomyces israelii		29.60	0.37	SOR 11		
07537160	**Low discrimination**					**07537160**
Actinomyces naeslundii		47.85	0.38	XYL 1		
Actinomyces israelii		38.08	0.28			
Clostridium perfringens		14.07	0.07			
07537161	**Good identification**					**07537161**
Actinomyces naeslundii		100.0	0.41	XYL 1		
07537163	**Good identification**					**07537163**
Actinomyces naeslundii		100.0	0.41	XYL 1		
07537300	**Good identification**					**07537300**
Actinomyces israelii		100.0	0.36	SAL 92		
07537302	**Low discrimination**					**07537302**
					GRA	
Actinomyces israelii		72.70	0.39	SAL 92	99	
Parabacteroides distasonis		18.38	0.25	CEL 93	1	
Bifidobacterium adolescentis		8.93	0.17	RHA 7	99	
07537322	**Very good identification**					**07537322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		99.61	0.59	MAN 1	1	
Actinomyces israelii		0.25	0.18		99	
Prevotella oralis		0.13	-.03		1	
07537332	**Good identification**					**07537332**
Parabacteroides distasonis		99.23	0.54	MAN 1		
Clostridium clostridioforme		0.77	0.13			
07537400	**Low discrimination**					**07537400**
					CAT	
Actinomyces israelii		82.99	0.43	SAL 92	1	
Propionibacterium granulosum		12.77	0.24	CEL 93	90	
Lactobacillus fermentum		2.37	0.16		1	
07537402	**Low discrimination**					**07537402**
					CAT	
Actinomyces israelii		73.42	0.46	SAL 92	1	
Bifidobacterium adolescentis		22.20	0.31	CEL 93	1	
Propionibacterium granulosum		1.88	0.13		90	
07537440	**Good identification**					**07537440**
Actinomyces israelii		100.0	0.36	SAL 92		
07537442	**Acceptable identification**					**07537442**
Actinomyces israelii		91.23	0.39	SAL 92		
Bifidobacterium adolescentis		8.77	0.16	CEL 93		
07537500	**Low discrimination**					**07537500**
Actinomyces israelii		58.92	0.54	SAL 92		
Bifidobacterium longum subsp. longum		32.91	0.41	CEL 93		
Bifidobacterium adolescentis		6.68	0.31			
07537502	**Low discrimination**					**07537502**
Bifidobacterium adolescentis		84.31	0.65			
Actinomyces israelii		11.27	0.57			
Bifidobacterium longum subsp. longum		4.20	0.41			
07537510	**Low discrimination**					**07537510**
Bifidobacterium longum subsp. longum		76.76	0.45	MAN 1		
Actinomyces israelii		12.06	0.39			
Bifidobacterium adolescentis		11.18	0.33			
07537512	**Good identification**					**07537512**
Bifidobacterium adolescentis		91.87	0.67			
Bifidobacterium longum subsp. longum		6.37	0.45			
Actinomyces israelii		1.50	0.42			
07537520	**Low discrimination**					**07537520**
					GRA	
Bifidobacterium longum subsp. longum		89.00	0.52	MAN 1	99	
Bacteroides vulgatus		8.17	0.20		1	
Actinomyces israelii		2.54	0.33		99	
07537522	**Low discrimination**					**07537522**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		70.81	0.52	MAN 1	99	
Parabacteroides distasonis		22.35	0.47		1	
Bifidobacterium adolescentis		3.59	0.31		99	
07537530	**Good identification**					**07537530**
Bifidobacterium longum subsp. longum		99.75	0.56	MAN 1		
Actinomyces israelii		0.25	0.19			
07537532	**Low discrimination**					**07537532**
					GRA	
Bifidobacterium longum subsp. longum		86.82	0.56	MAN 1	99	
Parabacteroides distasonis		8.67	0.43		1	
Bifidobacterium adolescentis		3.16	0.33		99	
07537540	**Good identification**					**07537540**
Actinomyces israelii		95.19	0.47	SAL 92		
Bifidobacterium adolescentis		3.43	0.16	CEL 93		
Bifidobacterium longum subsp. longum		1.38	0.07			
07537542	**Low discrimination**					**07537542**
Bifidobacterium adolescentis		70.20	0.50	SOR 11		
Actinomyces israelii		29.52	0.50			
Bifidobacterium longum subsp. longum		0.29	0.07			
07537552	**Acceptable identification**					**07537552**
Bifidobacterium adolescentis		94.60	0.51	SOR 11		
Actinomyces israelii		4.86	0.35			
Bifidobacterium longum subsp. longum		0.54	0.11			
07537720	**Good identification**					**07537720**
					GRA	
Bacteroides vulgatus		97.43	0.36	TRE 4	1	
Bifidobacterium longum subsp. longum		1.19	0.17	MAN 1	99	
Parabacteroides distasonis		1.13	0.21		1	
07537722	**Acceptable identification**					**07537722**
Parabacteroides distasonis		93.80	0.55	MAN 1		
Clostridium clostridioforme		2.21	0.22			
Bacteroides thetaiotaomicron		1.38	0.08			
07537732	**Low discrimination**					**07537732**
					GRA	
Parabacteroides distasonis		80.01	0.51	MAN 1	1	
Clostridium clostridioforme		15.96	0.33		50	
Bifidobacterium longum subsp. longum		2.70	0.21		99	
07540101	**Good identification**					**07540101**
Actinomyces naeslundii		100.0	0.35	FRU 95		
07540121	**Low discrimination**					**07540121**
					GRA	
Actinomyces naeslundii		71.96	0.35	FRU 95	99	
Prevotella melaninogenica		28.04	-.01	GAL 96	1	
07540123	**Good identification**					**07540123**
Actinomyces naeslundii		100.0	0.35	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07541000	**Low discrimination**						**07541000**
					COC	GRA	
Actinomyces naeslundii		35.66	0.48	FRU 95	1	99	
Gemella morbillorum		32.82	0.26		99	99	
Fusobacterium mortiferum		11.25	0.28		1	1	
07541001	**Good identification**						**07541001**
Actinomyces naeslundii		100.0	0.51	FRU 95			
07541003	**Low discrimination**						**07541003**
Actinomyces naeslundii		76.78	0.51	FRU 95			
Actinomyces viscosus		23.22	0.18				
07541021	**Good identification**						**07541021**
					GRA		
Actinomyces naeslundii		95.58	0.51	FRU 95	99		
Prevotella melaninogenica		4.42	-.01		1		
07541023	**Good identification**						**07541023**
Actinomyces naeslundii		100.0	0.51	FRU 95			
07541040	**Good identification**						**07541040**
Actinomyces naeslundii		100.0	0.35	FRU 95			
07541041	**Good identification**						**07541041**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07541042	**Good identification**						**07541042**
Actinomyces naeslundii		100.0	0.35	FRU 95			
07541043	**Good identification**						**07541043**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07541060	**Low discrimination**						**07541060**
					GRA		
Actinomyces naeslundii		71.80	0.35	FRU 95	99		
Prevotella melaninogenica		28.20	-.01		1		
07541061	**Good identification**						**07541061**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07541062	**Good identification**						**07541062**
Actinomyces naeslundii		100.0	0.35	FRU 95			
07541063	**Good identification**						**07541063**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07541101	**Low discrimination**						**07541101**
					GRA	SPO	
Actinomyces naeslundii		88.83	0.59	FRU 95	99	1	
Clostridium chauvoei		9.73	0.19		99	99	
Prevotella melaninogenica		1.44	-.01		1	1	
07541102	**Low discrimination**						**07541102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		31.94	0.56	FRU 95	99	
Actinomyces viscosus		27.15	0.31		99	
Prevotella oralis		20.34	0.27		1	
07541103	**Low discrimination**					**07541103**
					SPO	
Actinomyces naeslundii		82.77	0.59	FRU 95	1	
Actinomyces viscosus		15.64	0.22		1	
Clostridium chauvoei		1.60	0.07		99	
07541111	**Good identification**					**07541111**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07541113	**Good identification**					**07541113**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07541120	**Very good identification**					**07541120**
					GRA	
Prevotella melaninogenica		99.17	0.67	FRU 90	1	
Actinomyces naeslundii		0.42	0.56	CEL 10	99	
Prevotella oralis		0.27	0.27		1	
07541122	**Low discrimination**					**07541122**
					GRA	
Prevotella melaninogenica		77.00	0.55	FRU 90	1	
Prevotella oralis		20.49	0.49	CEL 10	1	
Actinomyces naeslundii		1.69	0.56		99	
07541123	**Low discrimination**					**07541123**
					GRA	
Actinomyces naeslundii		72.07	0.59	FRU 95	99	
Prevotella melaninogenica		22.06	0.21		1	
Prevotella oralis		5.87	0.15		1	
07541131	**Low discrimination**					**07541131**
					GRA	
Actinomyces naeslundii		76.43	0.37	FRU 95	99	
Prevotella melaninogenica		23.58	-.01	MLZ 5	1	
07541133	**Good identification**					**07541133**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07541140	**Low discrimination**					**07541140**
					GRA	
Actinomyces naeslundii		87.87	0.43	FRU 95	99	
Prevotella melaninogenica		12.13	-.01		1	
07541141	**Good identification**					**07541141**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07541142	**Good identification**					**07541142**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07541143	**Good identification**					**07541143**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07541161	**Acceptable identification**					**07541161**
					GRA	
Actinomyces naeslundii		91.58	0.46	FRU 95	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		8.43	-.01		1	
07541163	**Good identification**					**07541163**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07541322	**Acceptable identification**					**07541322**
Prevotella oralis		90.04	0.42	FRU 99		
Prevotella melaninogenica		7.97	0.21			
Parabacteroides distasonis		1.99	0.27			
07541522	**Low discrimination**					**07541522**
					GRA	
Bifidobacterium longum subsp. longum		56.02	0.37	FRU 90	99	
Prevotella melaninogenica		30.77	0.21	CEL 1	1	
Prevotella oralis		8.19	0.15		1	
07541530	**Good identification**					**07541530**
					GRA	
Bifidobacterium longum subsp. longum		98.32	0.41	FRU 90	99	
Prevotella melaninogenica		1.68	-.01	CEL 1	1	
07541532	**Good identification**					**07541532**
					GRA	
Bifidobacterium longum subsp. longum		99.12	0.41	FRU 90	99	
Parabacteroides distasonis		0.88	0.10	CEL 1	1	
07542100	**Good identification**					**07542100**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07542101	**Good identification**					**07542101**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07542102	**Low discrimination**					**07542102**
Actinomyces naeslundii		76.29	0.37	FRU 95		
Actinomyces viscosus		13.21	0.00	GAL 96		
Propionibacterium propionicum		10.50	0.12			
07542103	**Good identification**					**07542103**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07542120	**Low discrimination**					**07542120**
					GRA	
Actinomyces naeslundii		76.86	0.37	FRU 95	99	
Prevotella melaninogenica		23.14	-.01	GAL 96	1	
07542121	**Good identification**					**07542121**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07542122	**Good identification**					**07542122**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07542123	**Good identification**					**07542123**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07543000	**Good identification**					**07543000**
Actinomyces naeslundii		95.47	0.53	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		4.53	0.16			
07543001	**Good identification**					**07543001**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07543002	**Good identification**					**07543002**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07543003	**Good identification**					**07543003**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07543020	**Good identification**					**07543020**
					GRA	
Actinomyces naeslundii		96.55	0.53	FRU 95	99	
Prevotella melaninogenica		3.45	-.01		1	
07543021	**Good identification**					**07543021**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07543022	**Good identification**					**07543022**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07543023	**Good identification**					**07543023**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07543040	**Good identification**					**07543040**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07543041	**Good identification**					**07543041**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07543042	**Good identification**					**07543042**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07543043	**Good identification**					**07543043**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07543060	**Good identification**					**07543060**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07543061	**Good identification**					**07543061**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07543062	**Good identification**					**07543062**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07543063	**Good identification**					**07543063**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07543100	**Very good identification**					**07543100**
					GRA	
Actinomyces naeslundii		98.76	0.61	FRU 95	99	
Prevotella melaninogenica		1.24	-.01		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07543101	**Very good identification**					**07543101**
Actinomyces naeslundii		100.0	0.64	FRU 95		
07543102	**Very good identification**					**07543102**
Actinomyces naeslundii		97.69	0.61	FRU 95		
Bifidobacterium adolescentis		1.03	0.11			
Actinomyces israelii		0.72	0.15			
07543103	**Very good identification**					**07543103**
Actinomyces naeslundii		100.0	0.64	FRU 95		
07543110	**Good identification**					**07543110**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07543111	**Good identification**					**07543111**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07543112	**Low discrimination**					**07543112**
Actinomyces naeslundii		80.40	0.39	FRU 95		
Bifidobacterium adolescentis		19.60	0.13	MLZ 5		
07543113	**Good identification**					**07543113**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07543121	**Very good identification**					**07543121**
					GRA	
Actinomyces naeslundii		99.17	0.64	FRU 95	99	
Prevotella melaninogenica		0.83	-.01		1	
07543122	**Low discrimination**					**07543122**
					GRA	
Actinomyces naeslundii		76.04	0.61	FRU 95	99	
Prevotella melaninogenica		17.99	0.21		1	
Prevotella oralis		4.79	0.15		1	
07543123	**Very good identification**					**07543123**
Actinomyces naeslundii		100.0	0.64	FRU 95		
07543130	**Low discrimination**					**07543130**
					GRA	
Actinomyces naeslundii		80.75	0.39	FRU 95	99	
Prevotella melaninogenica		19.25	-.01	MLZ 5	1	
07543131	**Good identification**					**07543131**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07543132	**Good identification**					**07543132**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07543133	**Good identification**					**07543133**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07543140	**Good identification**					**07543140**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.48	FRU 95		
07543141	**Good identification**					**07543141**
	Actinomyces naeslundii	100.0	0.51	FRU 95		
07543142	**Good identification**					**07543142**
	Actinomyces naeslundii	96.85	0.48	FRU 95		
	Propionibacterium propionicum	3.15	0.12			
07543143	**Good identification**					**07543143**
	Actinomyces naeslundii	100.0	0.51	FRU 95		
07543160	**Acceptable identification**					**07543160**
					GRA	
	Actinomyces naeslundii	93.36	0.48	FRU 95	99	
	Prevotella melaninogenica	6.64	-.01		1	
07543161	**Good identification**					**07543161**
	Actinomyces naeslundii	100.0	0.51	FRU 95		
07543162	**Good identification**					**07543162**
	Actinomyces naeslundii	100.0	0.48	FRU 95		
07543163	**Good identification**					**07543163**
	Actinomyces naeslundii	100.0	0.51	FRU 95		
07545322	**Low discrimination**					**07545322**
					CAT	
	Parabacteroides distasonis	61.02	0.43	FRU 99	76	
	Prevotella oralis	37.93	0.27		1	
	Bacteroides fragilis	1.05	-.09		99	
07545332	**Good identification**					**07545332**
	Parabacteroides distasonis	97.08	0.39	FRU 99		
	Clostridium clostridioforme	2.93	0.07			
07545500	**Low discrimination**					**07545500**
					SPO	
	Bifidobacterium longum subsp. longum	89.14	0.35	FRU 90	1	
	Lactobacillus fermentum	4.18	0.20	CEL 1	1	
	Clostridium butyricum	2.81	-.04		99	
07545502	**Low discrimination**					**07545502**
	Bifidobacterium adolescentis	62.12	0.41	FRU 99		
	Bifidobacterium longum subsp. longum	34.92	0.35			
	Lactobacillus fermentum	1.64	0.20			
07545510	**Good identification**					**07545510**
	Bifidobacterium longum subsp. longum	98.73	0.39	FRU 90		
	Bifidobacterium adolescentis	1.27	0.09	CEL 1		
07545512	**Low discrimination**					**07545512**
	Bifidobacterium adolescentis	55.27	0.43	FRU 99		
	Bifidobacterium longum subsp. longum	43.28	0.39			
	Clostridium clostridioforme	1.45	0.12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07545520	**Low discrimination**						**07545520**
					GRA		
Bifidobacterium longum subsp. longum		78.70	0.46	FRU 90	99		
Bacteroides vulgatus		20.54	0.21	CEL 1	1		
Prevotella melaninogenica		0.76	-.01		1		
07545522	**Acceptable identification**						**07545522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.78	0.46	FRU 90	99	1	
Parabacteroides distasonis		7.62	0.31	CEL 1	1	76	
Bacteroides fragilis		0.64	-.09		1	99	
07545530	**Good identification**						**07545530**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
07545532	**Good identification**						**07545532**
					GRA		
Bifidobacterium longum subsp. longum		96.33	0.50	FRU 90	99		
Parabacteroides distasonis		2.56	0.27	CEL 1	1		
Clostridium clostridioforme		0.81	0.12		50		
07545720	**Good identification**						**07545720**
					GRA		
Bacteroides vulgatus		99.57	0.38	FRU 99	1		
Bifidobacterium longum subsp. longum		0.43	0.11	CEL 5	99		
07545722	**Low discrimination**						**07545722**
Parabacteroides distasonis		80.31	0.39	FRU 99			
Bacteroides vulgatus		7.56	0.04				
Clostridium clostridioforme		7.31	0.16				
07547502	**Low discrimination**						**07547502**
Bifidobacterium adolescentis		87.10	0.40	FRU 99			
Actinomyces israelii		12.90	0.32				
07547512	**Good identification**						**07547512**
Bifidobacterium adolescentis		98.22	0.41	FRU 99			
Actinomyces israelii		1.78	0.18				
07550000	**Low discrimination**						**07550000**
					GRA	CAT	
Prevotella intermedia		40.90	0.49	CEL 1	1	1	
Fusobacterium mortiferum		33.79	0.59		1	5	
Propionibacterium granulosum		19.63	0.52		99	90	
07550001	**Low discrimination**						**07550001**
					GRA		
Actinomyces naeslundii		61.96	0.49	GAL 96	99		
Prevotella intermedia		13.28	0.15		1		
Fusobacterium mortiferum		10.97	0.25		1		
07550002	**Low discrimination**						**07550002**
					GRA	CAT	
Actinomyces viscosus		86.50	0.52	NIT 99	99	76	
Propionibacterium granulosum		6.20	0.41		99	90	
Fusobacterium mortiferum		4.74	0.43		1	5	
07550003	**Acceptable identification**						**07550003**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		92.21	0.44	NIT 99			
Actinomyces naeslundii		7.79	0.49				
07550010	**Low discrimination**						**07550010**
					GRA	CAT	
Propionibacterium granulosum		76.50	0.37	CEL 1	99	90	
Prevotella intermedia		11.81	0.15	MLZ 12	1	1	
Fusobacterium mortiferum		9.76	0.25		1	5	
07550020	**Low discrimination**						**07550020**
					GRA	CAT	
Propionibacterium granulosum		56.20	0.41	CEL 1	99	90	
Actinomyces naeslundii		14.72	0.46		99	25	
Prevotella melaninogenica		11.59	0.15		1	1	
07550021	**Good identification**						**07550021**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07550023	**Low discrimination**						**07550023**
Actinomyces naeslundii		89.32	0.49	GAL 96			
Actinomyces viscosus		10.68	0.09				
07550041	**Good identification**						**07550041**
Actinomyces naeslundii		100.0	0.36	GAL 96			
07550043	**Low discrimination**						**07550043**
Actinomyces naeslundii		59.60	0.36	GAL 96			
Actinomyces viscosus		40.40	0.09				
07550061	**Good identification**						**07550061**
Actinomyces naeslundii		100.0	0.36	GAL 96			
07550063	**Good identification**						**07550063**
Actinomyces naeslundii		100.0	0.36	GAL 96			
07550100	**Good identification**						**07550100**
					GRA		
Fusobacterium mortiferum		91.96	0.65	MLT 11	1		
Actinomyces naeslundii		4.27	0.54		99		
Actinomyces viscosus		1.43	0.22		99		
07550101	**Low discrimination**						**07550101**
					GRA		
Actinomyces naeslundii		81.97	0.57	GAL 96	99		
Fusobacterium mortiferum		11.90	0.31		1		
Actinomyces viscosus		6.13	0.14		99		
07550102	**Low discrimination**						**07550102**
					GRA		
Actinomyces viscosus		88.18	0.56	NIT 99	99		
Fusobacterium mortiferum		6.35	0.49		1		
Actinomyces naeslundii		2.65	0.54		99		
07550103	**Low discrimination**						**07550103**
					GRA		
Actinomyces viscosus		87.92	0.48	NIT 99	99		
Actinomyces naeslundii		11.89	0.57		99		
Fusobacterium mortiferum		0.19	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07550111	**Good identification**					**07550111**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07550120	**Acceptable identification**					**07550120**
					GRA	
Prevotella melaninogenica		91.68	0.50	CEL 10	1	
Actinomyces naeslundii		3.35	0.54	GAL 99	99	
Prevotella oralis		2.69	0.27		1	
07550121	**Low discrimination**					**07550121**
					GRA	
Actinomyces naeslundii		84.42	0.57	GAL 96	99	
Prevotella melaninogenica		15.58	0.15		1	
07550122	**Low discrimination**					**07550122**
					GRA	
Prevotella oralis		67.95	0.49	GAL 99	1	
Prevotella melaninogenica		23.21	0.37		1	
Actinomyces naeslundii		4.45	0.54		99	
07550123	**Low discrimination**					**07550123**
					GRA	
Actinomyces naeslundii		82.46	0.57	GAL 96	99	
Prevotella oralis		8.49	0.15		1	
Actinomyces viscosus		6.16	0.14		99	
07550131	**Good identification**					**07550131**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07550133	**Good identification**					**07550133**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07550141	**Good identification**					**07550141**
Actinomyces naeslundii		100.0	0.44	GAL 96		
07550143	**Low discrimination**					**07550143**
Actinomyces naeslundii		70.26	0.44	GAL 96		
Actinomyces viscosus		29.75	0.14			
07550161	**Good identification**					**07550161**
Actinomyces naeslundii		100.0	0.44	GAL 96		
07550162	**Low discrimination**					**07550162**
					GRA	
Actinomyces naeslundii		39.33	0.41	GAL 96	99	
Prevotella oralis		34.40	0.15		1	
Prevotella melaninogenica		11.75	0.03		1	
07550163	**Good identification**					**07550163**
Actinomyces naeslundii		100.0	0.44	GAL 96		
07550322	**Good identification**					**07550322**
Prevotella oralis		96.86	0.42	GAL 99		
Parabacteroides distasonis		2.14	0.27			
Prevotella melaninogenica		0.78	0.03			
07550712	**Good identification**					**07550712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.36	SAL 88		
07550732	**Good identification**					**07550732**
Clostridium clostridioforme		95.77	0.36	SAL 88		
Parabacteroides distasonis		4.23	0.19	TRE 88		
07551000	**Low discrimination**					**07551000**
					GRA	
Fusobacterium mortiferum		30.00	0.62	MLT 11	1	
Prevotella intermedia		24.21	0.49		1	
Actinomyces naeslundii		18.25	0.70		99	
07551001	**Very good identification**					**07551001**
					GRA	
Actinomyces naeslundii		96.93	0.73		99	
Fusobacterium mortiferum		1.07	0.28		1	
Prevotella intermedia		0.87	0.15		1	
07551002	**Low discrimination**					**07551002**
					GRA	
Actinomyces viscosus		33.37	0.48	NIT 99	99	
Actinomyces naeslundii		24.52	0.70		99	
Prevotella oralis		14.78	0.41		1	
07551003	**Low discrimination**					**07551003**
					GRA	
Actinomyces naeslundii		76.29	0.73		99	
Actinomyces viscosus		23.07	0.40		99	
Prevotella oralis		0.31	0.06		1	
07551010	**Low discrimination**					**07551010**
					CAT	
Propionibacterium granulosum		36.92	0.37	CEL 1	90	
Lactobacillus jensenii		25.98	0.28	MLZ 12	1	
Actinomyces naeslundii		22.38	0.48		25	
07551011	**Good identification**					**07551011**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
07551012	**Low discrimination**					**07551012**
					CAT	
Actinomyces naeslundii		36.07	0.48	MLZ 5	25	
Lactobacillus jensenii		27.92	0.25		1	
Propionibacterium granulosum		14.88	0.27		90	
07551013	**Acceptable identification**					**07551013**
Actinomyces naeslundii		94.51	0.51	MLZ 5		
Actinomyces viscosus		5.49	0.06			
07551020	**Low discrimination**					**07551020**
					GRA	
Prevotella melaninogenica		54.73	0.50	CEL 10	1	
Actinomyces naeslundii		16.84	0.70	LAC 99	99	
Clostridium perfringens		11.25	0.45		99	
07551021	**Very good identification**					**07551021**
					GRA	
Actinomyces naeslundii		97.05	0.73		99	
Prevotella melaninogenica		2.12	0.15		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.44	0.11		99		
07551022	**Low discrimination**						**07551022**
					GRA		
Prevotella oralis		83.19	0.62	LAC 94	1		
Actinomyces naeslundii		7.27	0.70		99		
Prevotella melaninogenica		4.50	0.37		1		
07551023	**Good identification**						**07551023**
					GRA		
Actinomyces naeslundii		92.04	0.73		99		
Prevotella oralis		7.10	0.28		1		
Prevotella melaninogenica		0.38	0.03		1		
07551030	**Low discrimination**						**07551030**
					GRA	CAT	
Actinomyces naeslundii		38.91	0.48	MLZ 5	99	25	
Prevotella melaninogenica		24.27	0.15		1	1	
Propionibacterium granulosum		16.04	0.27		99	90	
07551031	**Good identification**						**07551031**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
07551033	**Good identification**						**07551033**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
07551040	**Low discrimination**						**07551040**
					GRA		
Actinomyces naeslundii		78.18	0.57		99		
Fusobacterium mortiferum		7.36	0.28		1		
Prevotella intermedia		5.94	0.15		1		
07551041	**Very good identification**						**07551041**
Actinomyces naeslundii		100.0	0.60				
07551042	**Low discrimination**						**07551042**
Actinomyces naeslundii		81.06	0.57				
Actinomyces viscosus		6.32	0.14				
Lactobacillus acidophilus		4.34	0.11				
07551043	**Very good identification**						**07551043**
Actinomyces naeslundii		98.30	0.60				
Actinomyces viscosus		1.70	0.06				
07551050	**Good identification**						**07551050**
Actinomyces naeslundii		100.0	0.35	MLZ 5			
07551051	**Good identification**						**07551051**
Actinomyces naeslundii		100.0	0.38	MLZ 5			
07551052	**Good identification**						**07551052**
Actinomyces naeslundii		100.0	0.35	MLZ 5			
07551053	**Good identification**						**07551053**
Actinomyces naeslundii		100.0	0.38	MLZ 5			
07551060	**Low discrimination**						**07551060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		51.51	0.57		99		
Clostridium perfringens		37.13	0.32		99		
Prevotella melaninogenica		9.58	0.15		1		
07551061	**Very good identification**						**07551061**
Actinomyces naeslundii		100.0	0.60				
07551062	**Low discrimination**						**07551062**
					GRA		
Actinomyces naeslundii		47.57	0.57		99		
Prevotella oralis		31.18	0.28		1		
Clostridium perfringens		16.13	0.27		99		
07551063	**Very good identification**						**07551063**
Actinomyces naeslundii		100.0	0.60				
07551070	**Good identification**						**07551070**
Actinomyces naeslundii		100.0	0.35	MLZ 5			
07551071	**Good identification**						**07551071**
Actinomyces naeslundii		100.0	0.38	MLZ 5			
07551072	**Good identification**						**07551072**
Actinomyces naeslundii		100.0	0.35	MLZ 5			
07551073	**Good identification**						**07551073**
Actinomyces naeslundii		100.0	0.38	MLZ 5			
07551100	**Low discrimination**						**07551100**
					GRA		
Fusobacterium mortiferum		33.11	0.68	MLT 11	1		
Prevotella melaninogenica		28.06	0.50		1		
Actinomyces naeslundii		24.57	0.78		99		
07551101	**Very good identification**						**07551101**
					GRA	SPO	
Actinomyces naeslundii		96.61	0.81		99	1	
Clostridium chauvoei		1.30	0.26		99	99	
Fusobacterium mortiferum		0.88	0.34		1	1	
07551102	**Low discrimination**						**07551102**
					GRA		
Prevotella oralis		51.50	0.61	NAG 95	1		
Actinomyces naeslundii		15.52	0.78		99		
Actinomyces viscosus		13.20	0.52		99		
07551103	**Low discrimination**						**07551103**
					GRA		
Actinomyces naeslundii		81.86	0.81		99		
Actinomyces viscosus		15.46	0.44		99		
Prevotella oralis		1.83	0.27		1		
07551110	**Low discrimination**						**07551110**
					GRA		
Actinomyces naeslundii		59.13	0.56	MLZ 5	99		
Fusobacterium mortiferum		15.29	0.34		1		
Prevotella melaninogenica		12.96	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07551111	**Very good identification**					**07551111**
Actinomyces naeslundii		100.0	0.59	MLZ 5		
07551112	**Low discrimination**					**07551112**
					GRA	
Bifidobacterium adolescentis		54.65	0.49	XYL 87	99	
Actinomyces naeslundii		18.13	0.56	ARA 87	99	
Prevotella oralis		11.54	0.27		1	
07551113	**Good identification**					**07551113**
Actinomyces naeslundii		94.65	0.59	MLZ 5		
Actinomyces viscosus		3.43	0.10			
Bifidobacterium adolescentis		1.92	0.14			
07551120	**Good identification**					**07551120**
					GRA	
Prevotella melaninogenica		94.71	0.84	CEL 10	1	
Prevotella oralis		2.78	0.61		1	
Clostridium perfringens		1.44	0.59		99	
07551121	**Low discrimination**					**07551121**
					GRA	
Actinomyces naeslundii		55.52	0.81		99	
Prevotella melaninogenica		42.27	0.50		1	
Prevotella oralis		1.24	0.27		1	
07551122	**Low discrimination**					**07551122**
					CAT	
Prevotella oralis		71.49	0.83		1	
Prevotella melaninogenica		24.42	0.71		1	
Capnocytophaga ochracea		1.17	0.53		90	
07551123	**Low discrimination**					**07551123**
					GRA	
Actinomyces naeslundii		62.93	0.81		99	
Prevotella oralis		26.71	0.49		1	
Prevotella melaninogenica		9.13	0.37		1	
07551130	**Acceptable identification**					**07551130**
					GRA	
Prevotella melaninogenica		90.03	0.50	CEL 10	1	
Actinomyces naeslundii		4.15	0.56	MLZ 1	99	
Prevotella oralis		2.64	0.27		1	
07551131	**Low discrimination**					**07551131**
					GRA	
Actinomyces naeslundii		87.25	0.59	MLZ 5	99	
Prevotella melaninogenica		12.75	0.15		1	
07551132	**Low discrimination**					**07551132**
					CAT	
Prevotella oralis		54.63	0.49	MLZ 1	1	
Prevotella melaninogenica		18.66	0.37		1	
Capnocytophaga ochracea		9.81	0.36		90	
07551133	**Low discrimination**					**07551133**
					GRA	
Actinomyces naeslundii		87.90	0.59	MLZ 5	99	
Prevotella oralis		7.16	0.15		1	
Prevotella melaninogenica		2.45	0.03		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07551140	**Low discrimination**					**07551140**
					GRA	
Actinomyces naeslundii		79.57	0.65		99	
Fusobacterium mortiferum		6.14	0.34		1	
Prevotella melaninogenica		5.20	0.15		1	
07551141	**Very good identification**					**07551141**
Actinomyces naeslundii		100.0	0.68			
07551142	**Low discrimination**					**07551142**
					GRA	
Actinomyces naeslundii		56.85	0.65		99	
Lactobacillus acidophilus		12.29	0.29		99	
Prevotella oralis		10.80	0.27		1	
07551143	**Very good identification**					**07551143**
Actinomyces naeslundii		98.93	0.68			
Actinomyces viscosus		1.07	0.10			
07551150	**Acceptable identification**					**07551150**
Actinomyces naeslundii		92.02	0.43	MLZ 5		
Bifidobacterium breve		7.98	0.10			
07551151	**Good identification**					**07551151**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07551153	**Good identification**					**07551153**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07551160	**Low discrimination**					**07551160**
					GRA	
Prevotella melaninogenica		67.35	0.50	CEL 10	1	
Clostridium perfringens		19.33	0.47	SOR 1	99	
Actinomyces naeslundii		10.41	0.65		99	
07551161	**Good identification**					**07551161**
					GRA	
Actinomyces naeslundii		94.69	0.68		99	
Prevotella melaninogenica		4.13	0.15		1	
Clostridium perfringens		1.18	0.13		99	
07551162	**Low discrimination**					**07551162**
					GRA	
Prevotella oralis		46.93	0.49	SOR 1	1	
Prevotella melaninogenica		16.03	0.37		1	
Actinomyces naeslundii		13.01	0.65		99	
07551163	**Very good identification**					**07551163**
					GRA	
Actinomyces naeslundii		95.89	0.68		99	
Prevotella oralis		2.33	0.15		1	
Prevotella melaninogenica		0.80	0.03		1	
07551171	**Good identification**					**07551171**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07551173	**Good identification**					**07551173**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07551201	**Good identification**					**07551201**
Actinomyces naeslundii		100.0	0.39	RHA 1		
07551203	**Low discrimination**					**07551203**
					GRA	
Actinomyces naeslundii		67.81	0.39	RHA 1	99	
Actinomyces viscosus		20.51	0.06		99	
Prevotella oralis		11.68	0.00		1	
07551221	**Low discrimination**					**07551221**
					GRA	
Actinomyces naeslundii		85.31	0.39	RHA 1	99	
Prevotella oralis		14.69	0.00		1	
07551222	**Very good identification**					**07551222**
					GRA	
Prevotella oralis		99.14	0.56	LAC 94	1	
Parabacteroides distasonis		0.35	0.27		1	
Actinomyces naeslundii		0.20	0.36		99	
07551301	**Good identification**					**07551301**
					GRA	
Actinomyces naeslundii		95.25	0.47	RHA 1	99	
Prevotella oralis		4.75	-.01		1	
07551302	**Good identification**					**07551302**
					GRA	
Prevotella oralis		95.97	0.55	NAG 95	1	
Lactobacillus acidophilus		1.57	0.25		99	
Actinomyces naeslundii		0.68	0.44		99	
07551303	**Low discrimination**					**07551303**
					GRA	
Actinomyces naeslundii		46.80	0.47	RHA 1	99	
Prevotella oralis		44.36	0.21		1	
Actinomyces viscosus		8.84	0.10		99	
07551320	**Low discrimination**					**07551320**
Prevotella oralis		53.54	0.55	BGL 95		
Prevotella melaninogenica		43.00	0.50			
Bacteroides vulgatus		1.78	0.24			
07551321	**Low discrimination**					**07551321**
					GRA	
Actinomyces naeslundii		36.91	0.47	RHA 1	99	
Prevotella oralis		34.99	0.21		1	
Prevotella melaninogenica		28.10	0.15		1	
07551322	**Very good identification**					**07551322**
Prevotella oralis		96.75	0.77			
Parabacteroides distasonis		2.14	0.61			
Prevotella melaninogenica		0.78	0.37			
07551323	**Acceptable identification**					**07551323**
					GRA	
Prevotella oralis		92.11	0.42	URE 1	1	
Actinomyces naeslundii		5.11	0.47		99	
Parabacteroides distasonis		2.04	0.27		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07551332	**Low discrimination**					**07551332**
Parabacteroides distasonis		53.01	0.56	XYL 90		
Prevotella oralis		45.02	0.42			
Clostridium clostridioforme		1.42	0.24			
07551362	**Acceptable identification**					**07551362**
					GRA	
Prevotella oralis		94.60	0.42	SOR 1	1	
Parabacteroides distasonis		2.09	0.27		1	
Lactobacillus acidophilus		1.00	0.10		99	
07551400	**Low discrimination**					**07551400**
					GRA	
Lactobacillus fermentum		87.86	0.54	CEL 1	99	
Fusobacterium mortiferum		3.20	0.28		1	
Prevotella intermedia		2.59	0.15		1	
07551401	**Low discrimination**					**07551401**
Actinomyces naeslundii		76.72	0.39	ARA 1		
Lactobacillus fermentum		23.28	0.20			
07551402	**Low discrimination**					**07551402**
Lactobacillus fermentum		86.55	0.54	CEL 1		
Bifidobacterium adolescentis		4.75	0.27			
Actinomyces viscosus		2.61	0.14			
07551403	**Low discrimination**					**07551403**
Actinomyces naeslundii		62.27	0.39	ARA 1		
Lactobacillus fermentum		18.90	0.20			
Actinomyces viscosus		18.83	0.06			
07551420	**Low discrimination**					**07551420**
					GRA	
Lactobacillus fermentum		68.70	0.46	CEL 1	99	
Prevotella melaninogenica		14.86	0.15		1	
Bifidobacterium longum subsp. longum		5.16	0.20		99	
07551421	**Good identification**					**07551421**
Actinomyces naeslundii		100.0	0.39	ARA 1		
07551422	**Low discrimination**					**07551422**
					GRA	
Lactobacillus fermentum		48.87	0.46	CEL 1	99	
Prevotella oralis		37.25	0.28		1	
Bifidobacterium longum subsp. longum		3.67	0.20		99	
07551423	**Good identification**					**07551423**
Actinomyces naeslundii		100.0	0.39	ARA 1		
07551500	**Low discrimination**					**07551500**
					GRA	
Lactobacillus fermentum		62.27	0.58	CEL 1	99	
Bifidobacterium longum subsp. longum		23.63	0.43		99	
Fusobacterium mortiferum		3.25	0.34		1	
07551501	**Low discrimination**					**07551501**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		80.65	0.47	ARA 1			
Lactobacillus fermentum		14.03	0.24				
Bifidobacterium longum subsp. longum		5.32	0.09				
07551502	**Low discrimination**						**07551502**
Bifidobacterium adolescentis		65.55	0.61	XYL 87			
Lactobacillus fermentum		19.69	0.58				
Bifidobacterium longum subsp. longum		7.47	0.43				
07551503	**Low discrimination**						**07551503**
Actinomyces naeslundii		49.80	0.47	ARA 1			
Bifidobacterium adolescentis		28.84	0.27				
Actinomyces viscosus		9.41	0.10				
07551510	**Acceptable identification**						**07551510**
Bifidobacterium longum subsp. longum		92.11	0.47	CEL 1			
Bifidobacterium adolescentis		5.86	0.29				
Lactobacillus fermentum		1.44	0.24				
07551512	**Low discrimination**						**07551512**
Bifidobacterium adolescentis		84.54	0.63	XYL 87			
Bifidobacterium longum subsp. longum		13.43	0.47				
Clostridium clostridioforme		1.47	0.29				
07551520	**Low discrimination**						**07551520**
					GRA		
Prevotella melaninogenica		64.26	0.50	CEL 10	1		
Bifidobacterium longum subsp. longum		22.29	0.54	ARA 1	99		
Lactobacillus fermentum		4.90	0.50		99		
07551521	**Low discrimination**						**07551521**
					GRA		
Actinomyces naeslundii		46.96	0.47	ARA 1	99		
Prevotella melaninogenica		35.75	0.15		1		
Bifidobacterium longum subsp. longum		12.40	0.20		99		
07551522	**Low discrimination**						**07551522**
					GRA		
Prevotella oralis		37.96	0.49	ARA 1	1		
Bifidobacterium longum subsp. longum		23.61	0.54		99		
Prevotella melaninogenica		12.97	0.37		1		
07551523	**Low discrimination**						**07551523**
					GRA		
Actinomyces naeslundii		47.76	0.47	ARA 1	99		
Prevotella oralis		20.27	0.15		1		
Bifidobacterium longum subsp. longum		12.61	0.20		99		
07551530	**Very good identification**						**07551530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.68	0.58	CEL 1	99	1	
Prevotella melaninogenica		1.67	0.15		1	1	
Capnocytophaga ochracea		0.33	0.07		1	90	
07551532	**Low discrimination**						**07551532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		84.42	0.58	CEL 1	99	1	
Parabacteroides distasonis		8.22	0.45		1	76	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		2.53	0.23		1	90	
07551542	**Acceptable identification**						**07551542**
Bifidobacterium adolescentis		90.05	0.46	XYL 87			
Actinomyces israelii		3.53	0.28	SOR 11			
Lactobacillus fermentum		2.21	0.24				
07551552	**Good identification**						**07551552**
Bifidobacterium adolescentis		98.26	0.47	XYL 87			
Bifidobacterium longum subsp. longum		1.28	0.13	SOR 11			
Actinomyces israelii		0.47	0.14				
07551612	**Good identification**						**07551612**
Clostridium clostridioforme		100.0	0.35	SAL 88			
07551632	**Good identification**						**07551632**
Clostridium clostridioforme		95.30	0.35	SAL 88			
Parabacteroides distasonis		4.70	0.19	TRE 88			
07551712	**Acceptable identification**						**07551712**
					GRA		
Clostridium clostridioforme		90.35	0.44	SAL 88	50		
Bifidobacterium adolescentis		7.18	0.29	TRE 88	99		
Parabacteroides distasonis		1.33	0.19	XYL 91	1		
07551720	**Acceptable identification**						**07551720**
Bacteroides vulgatus		94.78	0.43	CEL 5			
Prevotella oralis		2.17	0.21	XYL 98			
Prevotella melaninogenica		1.75	0.15				
07551722	**Low discrimination**						**07551722**
Parabacteroides distasonis		53.30	0.57	XYL 90			
Prevotella oralis		39.76	0.42				
Clostridium clostridioforme		4.32	0.33				
07551732	**Low discrimination**						**07551732**
					GRA		
Parabacteroides distasonis		57.49	0.53	XYL 90	1		
Clostridium clostridioforme		39.41	0.44		50		
Bifidobacterium longum subsp. longum		1.99	0.24		99		
07552000	**Low discrimination**						**07552000**
					GRA	CAT	
Propionibacterium granulosum		82.01	0.50	CEL 1	99	90	
Actinomyces naeslundii		13.81	0.51		99	25	
Prevotella intermedia		2.29	0.15		1	1	
07552001	**Good identification**						**07552001**
Actinomyces naeslundii		96.16	0.54	GAL 96			
Propionibacterium granulosum		3.85	0.16				
07552002	**Low discrimination**						**07552002**
Propionibacterium granulosum		53.35	0.39	CEL 1			
Actinomyces naeslundii		35.94	0.51				
Actinomyces viscosus		9.97	0.18				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07552003	**Acceptable identification**					**07552003**
Actinomyces naeslundii		94.20	0.54	GAL 96		
Actinomyces viscosus		5.81	0.09			
07552020	**Low discrimination**					**07552020**
Propionibacterium granulosum		59.75	0.39	CEL 1		
Actinomyces naeslundii		40.25	0.51			
07552021	**Good identification**					**07552021**
Actinomyces naeslundii		100.0	0.54	GAL 96		
07552022	**Low discrimination**					**07552022**
Actinomyces naeslundii		72.94	0.51	GAL 96		
Propionibacterium granulosum		27.07	0.29			
07552023	**Good identification**					**07552023**
Actinomyces naeslundii		100.0	0.54	GAL 96		
07552040	**Low discrimination**					**07552040**
Actinomyces naeslundii		74.63	0.38	GAL 96		
Propionibacterium granulosum		25.37	0.16			
07552041	**Good identification**					**07552041**
Actinomyces naeslundii		100.0	0.41	GAL 96		
07552042	**Low discrimination**					**07552042**
Actinomyces naeslundii		89.54	0.38	GAL 96		
Propionibacterium propionicum		10.46	0.12			
07552043	**Good identification**					**07552043**
Actinomyces naeslundii		100.0	0.41	GAL 96		
07552060	**Good identification**					**07552060**
Actinomyces naeslundii		100.0	0.38	GAL 96		
07552061	**Good identification**					**07552061**
Actinomyces naeslundii		100.0	0.41	GAL 96		
07552062	**Good identification**					**07552062**
Actinomyces naeslundii		100.0	0.38	GAL 96		
07552063	**Good identification**					**07552063**
Actinomyces naeslundii		100.0	0.41	GAL 96		
07552100	**Low discrimination**					**07552100**
					GRA	
Actinomyces naeslundii		62.06	0.59	GAL 96	99	
Propionibacterium propionicum		29.68	0.43		99	
Fusobacterium mortiferum		6.96	0.31		1	
07552101	**Very good identification**					**07552101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.58	0.62	GAL 96		
Propionibacterium propionicum		3.42	0.27			
07552102	**Low discrimination**					**07552102**
Actinomyces naeslundii		51.93	0.59	GAL 96		
Propionibacterium propionicum		37.25	0.46			
Actinomyces viscosus		9.00	0.22			
07552103	**Good identification**					**07552103**
Actinomyces naeslundii		91.61	0.62	GAL 96		
Propionibacterium propionicum		4.87	0.30			
Actinomyces viscosus		3.53	0.14			
07552110	**Good identification**					**07552110**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07552111	**Good identification**					**07552111**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07552112	**Low discrimination**					**07552112**
Actinomyces naeslundii		69.18	0.37	GAL 96		
Bifidobacterium adolescentis		21.30	0.13	MLZ 5		
Propionibacterium propionicum		9.52	0.12			
07552113	**Good identification**					**07552113**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07552120	**Low discrimination**					**07552120**
					GRA	
Actinomyces naeslundii		87.52	0.59	GAL 96	99	
Prevotella melaninogenica		12.48	0.15		1	
07552121	**Very good identification**					**07552121**
Actinomyces naeslundii		100.0	0.62	GAL 96		
07552122	**Low discrimination**					**07552122**
					GRA	
Actinomyces naeslundii		88.22	0.59	GAL 96	99	
Prevotella oralis		7.02	0.15		1	
Prevotella melaninogenica		2.40	0.03		1	
07552123	**Very good identification**					**07552123**
Actinomyces naeslundii		100.0	0.62	GAL 96		
07552130	**Good identification**					**07552130**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07552131	**Good identification**					**07552131**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07552132	**Good identification**					**07552132**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07552133	**Good identification**					**07552133**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	GAL 96		
07552140	**Low discrimination**					**07552140**
Propionibacterium propionicum		73.04	0.43	NIT 86		
Actinomyces naeslundii		26.96	0.46	CEL 1		
07552141	**Low discrimination**					**07552141**
Actinomyces naeslundii		83.28	0.49	GAL 96		
Propionibacterium propionicum		16.72	0.27			
07552142	**Low discrimination**					**07552142**
Propionibacterium propionicum		80.26	0.46	NIT 86		
Actinomyces naeslundii		19.75	0.46	CEL 1		
07552143	**Low discrimination**					**07552143**
Actinomyces naeslundii		76.86	0.49	GAL 96		
Propionibacterium propionicum		23.14	0.30			
07552160	**Good identification**					**07552160**
Actinomyces naeslundii		100.0	0.46	GAL 96		
07552161	**Good identification**					**07552161**
Actinomyces naeslundii		100.0	0.49	GAL 96		
07552162	**Good identification**					**07552162**
Actinomyces naeslundii		96.06	0.46	GAL 96		
Propionibacterium propionicum		3.94	0.12			
07552163	**Good identification**					**07552163**
Actinomyces naeslundii		100.0	0.49	GAL 96		
07553000	**Low discrimination**					**07553000**
					CAT	
Actinomyces naeslundii		77.85	0.75		25	
Propionibacterium granulosum		19.26	0.50		90	
Actinomyces israelii		1.42	0.36		1	
07553001	**Very good identification**					**07553001**
Actinomyces naeslundii		99.83	0.78			
Propionibacterium granulosum		0.17	0.16			
07553002	**Good identification**					**07553002**
					CAT	
Actinomyces naeslundii		90.31	0.75		25	
Propionibacterium granulosum		5.59	0.39		90	
Actinomyces israelii		2.48	0.39		1	
07553003	**Very good identification**					**07553003**
Actinomyces naeslundii		99.84	0.78			
Actinomyces viscosus		0.16	0.06			
07553010	**Low discrimination**					**07553010**
					CAT	
Actinomyces naeslundii		59.07	0.53	MLZ 5	25	
Propionibacterium granulosum		37.86	0.35		90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		3.07	0.22		1		
07553011	**Good identification**						**07553011**
Actinomyces naeslundii		100.0	0.56	MLZ 5			
07553012	**Low discrimination**						**07553012**
					CAT		
Actinomyces naeslundii		78.45	0.53	MLZ 5	25		
Propionibacterium granulosum		12.57	0.25		90		
Actinomyces israelii		6.11	0.25		1		
07553013	**Good identification**						**07553013**
Actinomyces naeslundii		100.0	0.56	MLZ 5			
07553020	**Good identification**						**07553020**
					GRA	CAT	
Actinomyces naeslundii		92.04	0.75		99	25	
Propionibacterium granulosum		5.69	0.39		99	90	
Prevotella melaninogenica		1.56	0.15		1	1	
07553021	**Very good identification**						**07553021**
Actinomyces naeslundii		100.0	0.78				
07553022	**Good identification**						**07553022**
					GRA	CAT	
Actinomyces naeslundii		92.08	0.75		99	25	
Prevotella oralis		5.49	0.28		1	1	
Propionibacterium granulosum		1.42	0.29		99	90	
07553023	**Very good identification**						**07553023**
Actinomyces naeslundii		100.0	0.78				
07553030	**Low discrimination**						**07553030**
Actinomyces naeslundii		86.19	0.53	MLZ 5			
Propionibacterium granulosum		13.81	0.25				
07553031	**Good identification**						**07553031**
Actinomyces naeslundii		100.0	0.56	MLZ 5			
07553032	**Good identification**						**07553032**
Actinomyces naeslundii		96.15	0.53	MLZ 5			
Propionibacterium granulosum		3.85	0.14				
07553033	**Good identification**						**07553033**
Actinomyces naeslundii		100.0	0.56	MLZ 5			
07553040	**Good identification**						**07553040**
					CAT		
Actinomyces naeslundii		94.84	0.62		25		
Actinomyces israelii		3.82	0.29		1		
Propionibacterium granulosum		1.34	0.16		90		
07553041	**Very good identification**						**07553041**
Actinomyces naeslundii		100.0	0.65				
07553042	**Good identification**						**07553042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		93.87	0.62			
Actinomyces israelii		5.68	0.32			
Propionibacterium propionicum		0.46	0.12			
07553043	**Very good identification**					**07553043**
Actinomyces naeslundii		100.0	0.65			
07553050	**Low discrimination**					**07553050**
Actinomyces naeslundii		89.73	0.40	MLZ 5		
Actinomyces israelii		10.27	0.15			
07553051	**Good identification**					**07553051**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07553052	**Low discrimination**					**07553052**
Actinomyces naeslundii		85.35	0.40	MLZ 5		
Actinomyces israelii		14.65	0.18			
07553053	**Good identification**					**07553053**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07553060	**Very good identification**					**07553060**
Actinomyces naeslundii		100.0	0.62			
07553061	**Very good identification**					**07553061**
Actinomyces naeslundii		100.0	0.65			
07553062	**Very good identification**					**07553062**
Actinomyces naeslundii		100.0	0.62			
07553063	**Very good identification**					**07553063**
Actinomyces naeslundii		100.0	0.65			
07553070	**Good identification**					**07553070**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
07553071	**Good identification**					**07553071**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07553072	**Good identification**					**07553072**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
07553073	**Good identification**					**07553073**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07553100	**Good identification**					**07553100**
Actinomyces naeslundii		94.10	0.83			
Actinomyces israelii		2.42	0.46			
Propionibacterium propionicum		1.88	0.43			
07553101	**Very good identification**					**07553101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		99.85	0.86				
Propionibacterium propionicum		0.15	0.27				
07553102	**Low discrimination**						**07553102**
Actinomyces naeslundii		86.05	0.83				
Bifidobacterium adolescentis		4.71	0.46				
Actinomyces israelii		3.32	0.49				
07553103	**Very good identification**						**07553103**
Actinomyces naeslundii		99.62	0.86				
Propionibacterium propionicum		0.22	0.30				
Actinomyces viscosus		0.10	0.10				
07553110	**Good identification**						**07553110**
Actinomyces naeslundii		91.43	0.61	MLZ 5			
Actinomyces israelii		6.67	0.32				
Bifidobacterium adolescentis		1.17	0.13				
07553111	**Very good identification**						**07553111**
Actinomyces naeslundii		100.0	0.64	MLZ 5			
07553112	**Low discrimination**						**07553112**
Bifidobacterium adolescentis		51.53	0.47	XYL 87			
Actinomyces naeslundii		40.56	0.61	ARA 87			
Actinomyces israelii		4.44	0.35				
07553113	**Very good identification**						**07553113**
Actinomyces naeslundii		99.15	0.64	MLZ 5			
Bifidobacterium adolescentis		0.85	0.13				
07553120	**Low discrimination**						**07553120**
					SPO	GRA	
Actinomyces naeslundii		61.04	0.83		1	99	
Prevotella melaninogenica		35.91	0.50		1	1	
Clostridium ramosum		1.24	0.28		92	50	
07553121	**Very good identification**						**07553121**
					GRA		
Actinomyces naeslundii		99.61	0.86		99		
Prevotella melaninogenica		0.40	0.15		1		
07553122	**Low discrimination**						**07553122**
					GRA		
Actinomyces naeslundii		64.40	0.83		99		
Prevotella oralis		21.13	0.49		1		
Prevotella melaninogenica		7.22	0.37		1		
07553123	**Very good identification**						**07553123**
					GRA		
Actinomyces naeslundii		99.65	0.86		99		
Prevotella oralis		0.22	0.15		1		
Prevotella melaninogenica		0.08	0.03		1		
07553130	**Low discrimination**						**07553130**
					GRA		
Actinomyces naeslundii		88.74	0.61	MLZ 5	99		
Prevotella melaninogenica		10.02	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		1.24	0.15		99	
07553131	**Very good identification**					**07553131**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
07553132	**Low discrimination**					**07553132**
					GRA	
Actinomyces naeslundii		77.15	0.61	MLZ 5	99	
Bifidobacterium breve		10.93	0.32		99	
Prevotella oralis		4.86	0.15		1	
07553133	**Very good identification**					**07553133**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
07553140	**Low discrimination**					**07553140**
Actinomyces naeslundii		85.33	0.70			
Propionibacterium propionicum		9.64	0.43			
Actinomyces israelii		4.83	0.39			
07553141	**Very good identification**					**07553141**
Actinomyces naeslundii		99.17	0.73			
Propionibacterium propionicum		0.83	0.27			
07553142	**Low discrimination**					**07553142**
Actinomyces naeslundii		75.63	0.70			
Propionibacterium propionicum		12.81	0.46			
Actinomyces israelii		6.43	0.42			
07553143	**Very good identification**					**07553143**
Actinomyces naeslundii		98.76	0.73			
Propionibacterium propionicum		1.24	0.30			
07553150	**Low discrimination**					**07553150**
Actinomyces naeslundii		82.96	0.48	MLZ 5		
Actinomyces israelii		13.34	0.25			
Bifidobacterium breve		3.70	0.10			
07553151	**Good identification**					**07553151**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
07553152	**Low discrimination**					**07553152**
Actinomyces naeslundii		38.24	0.48	MLZ 5		
Bifidobacterium adolescentis		34.03	0.32			
Bifidobacterium breve		17.27	0.27			
07553153	**Good identification**					**07553153**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
07553160	**Very good identification**					**07553160**
					GRA	
Actinomyces naeslundii		95.14	0.70		99	
Prevotella melaninogenica		3.20	0.15		1	
Clostridium perfringens		0.92	0.13		99	
07553161	**Very good identification**					**07553161**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.73				
07553162	**Good identification**						**07553162**
					GRA		
Actinomyces naeslundii		91.49	0.70		99		
Bifidobacterium breve		3.87	0.32		99		
Prevotella oralis		1.72	0.15		1		
07553163	**Very good identification**						**07553163**
Actinomyces naeslundii		100.0	0.73				
07553170	**Acceptable identification**						**07553170**
Actinomyces naeslundii		92.64	0.48	MLZ 5			
Bifidobacterium breve		7.36	0.15				
07553171	**Good identification**						**07553171**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
07553172	**Low discrimination**						**07553172**
Actinomyces naeslundii		55.47	0.48	MLZ 5			
Bifidobacterium breve		44.53	0.32				
07553173	**Good identification**						**07553173**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
07553200	**Low discrimination**						**07553200**
					CAT		
Actinomyces naeslundii		52.81	0.41	RHA 1	25		
Propionibacterium granulosum		40.00	0.24		90		
Actinomyces israelii		7.20	0.17		1		
07553201	**Good identification**						**07553201**
Actinomyces naeslundii		100.0	0.44	RHA 1			
07553202	**Low discrimination**						**07553202**
					CAT		
Actinomyces naeslundii		65.50	0.41	RHA 1	25		
Actinomyces israelii		13.39	0.20		1		
Propionibacterium granulosum		12.40	0.14		90		
07553203	**Good identification**						**07553203**
Actinomyces naeslundii		100.0	0.44	RHA 1			
07553220	**Low discrimination**						**07553220**
					GRA	CAT	
Actinomyces naeslundii		75.62	0.41	RHA 1	99	25	
Propionibacterium granulosum		14.32	0.14		99	90	
Prevotella oralis		10.06	0.00		1	1	
07553221	**Good identification**						**07553221**
Actinomyces naeslundii		100.0	0.44	RHA 1			
07553223	**Good identification**						**07553223**
Actinomyces naeslundii		100.0	0.44	RHA 1			
07553300	**Low discrimination**						**07553300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		81.30	0.49	RHA 1	99		
Actinomyces israelii		15.57	0.27		99		
Prevotella oralis		3.14	-.01		1		
07553301	**Good identification**						**07553301**
Actinomyces naeslundii		100.0	0.52	RHA 1			
07553302	**Low discrimination**						**07553302**
					GRA		
Actinomyces naeslundii		43.97	0.49	RHA 1	99		
Prevotella oralis		32.21	0.21		1		
Actinomyces israelii		12.63	0.30		99		
07553303	**Good identification**						**07553303**
Actinomyces naeslundii		100.0	0.52	RHA 1			
07553321	**Good identification**						**07553321**
Actinomyces naeslundii		100.0	0.52	RHA 1			
07553322	**Low discrimination**						**07553322**
					SPO	GRA	
Prevotella oralis		53.44	0.42	MAN 1	1	1	
Clostridium ramosum		40.87	0.41		92	50	
Actinomyces naeslundii		3.84	0.49		1	99	
07553323	**Low discrimination**						**07553323**
					SPO	GRA	
Actinomyces naeslundii		85.81	0.52	RHA 1	1	99	
Prevotella oralis		8.04	0.08		1	1	
Clostridium ramosum		6.15	0.07		92	50	
07553340	**Low discrimination**						**07553340**
Actinomyces naeslundii		70.32	0.36	RHA 1			
Actinomyces israelii		29.68	0.20				
07553341	**Good identification**						**07553341**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07553342	**Low discrimination**						**07553342**
Actinomyces naeslundii		55.48	0.36	RHA 1			
Actinomyces israelii		35.13	0.23				
Propionibacterium propionicum		9.39	0.12				
07553343	**Good identification**						**07553343**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07553360	**Good identification**						**07553360**
Actinomyces naeslundii		100.0	0.36	RHA 1			
07553361	**Good identification**						**07553361**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07553362	**Low discrimination**						**07553362**
					SPO	GRA	
Actinomyces naeslundii		41.57	0.36	RHA 1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		33.11	0.08		1	1	
Clostridium ramosum		25.32	0.07		92	50	
07553363	**Good identification**						**07553363**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07553401	**Good identification**						**07553401**
Actinomyces naeslundii		100.0	0.44	ARA 1			
07553402	**Low discrimination**						**07553402**
Actinomyces israelii		49.39	0.37	SAL 92			
Bifidobacterium adolescentis		23.16	0.26	XYL 92			
Actinomyces naeslundii		22.23	0.41				
07553403	**Good identification**						**07553403**
Actinomyces naeslundii		100.0	0.44	ARA 1			
07553420	**Acceptable identification**						**07553420**
Actinomyces naeslundii		91.36	0.41	ARA 1			
Actinomyces israelii		8.64	0.14				
07553421	**Good identification**						**07553421**
Actinomyces naeslundii		100.0	0.44	ARA 1			
07553422	**Low discrimination**						**07553422**
Actinomyces naeslundii		87.58	0.41	ARA 1			
Actinomyces israelii		12.42	0.17				
07553423	**Good identification**						**07553423**
Actinomyces naeslundii		100.0	0.44	ARA 1			
07553500	**Low discrimination**						**07553500**
Actinomyces israelii		57.30	0.45	SAL 92			
Actinomyces naeslundii		27.52	0.49	XYL 92			
Bifidobacterium adolescentis		10.08	0.26				
07553501	**Good identification**						**07553501**
Actinomyces naeslundii		100.0	0.52	ARA 1			
07553502	**Low discrimination**						**07553502**
Bifidobacterium adolescentis		89.31	0.60	XYL 87			
Actinomyces israelii		7.70	0.48				
Actinomyces naeslundii		2.46	0.49				
07553503	**Low discrimination**						**07553503**
Actinomyces naeslundii		79.05	0.52	ARA 1			
Bifidobacterium adolescentis		19.29	0.26				
Actinomyces israelii		1.66	0.14				
07553512	**Very good identification**						**07553512**
Bifidobacterium adolescentis		98.65	0.61	XYL 87			
Actinomyces israelii		1.04	0.34				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		0.19	0.13			
07553520	**Low discrimination**					**07553520**
					GRA	
Actinomyces naeslundii		49.85	0.49	ARA 1	99	
Prevotella melaninogenica		29.33	0.15		1	
Bifidobacterium longum subsp. longum		10.17	0.20		99	
07553521	**Good identification**					**07553521**
Actinomyces naeslundii		100.0	0.52	ARA 1		
07553522	**Low discrimination**					**07553522**
					GRA	
Actinomyces naeslundii		40.13	0.49	ARA 1	99	
Bifidobacterium adolescentis		14.69	0.26		99	
Prevotella oralis		13.16	0.15		1	
07553523	**Good identification**					**07553523**
Actinomyces naeslundii		100.0	0.52	ARA 1		
07553540	**Low discrimination**					**07553540**
Actinomyces israelii		78.50	0.38	SAL 92		
Actinomyces naeslundii		17.11	0.36	XYL 92		
Bifidobacterium adolescentis		4.39	0.10			
07553541	**Good identification**					**07553541**
Actinomyces naeslundii		100.0	0.39	ARA 1		
07553542	**Low discrimination**					**07553542**
Bifidobacterium adolescentis		75.92	0.44	XYL 87		
Actinomyces israelii		20.58	0.41	SOR 11		
Actinomyces naeslundii		2.99	0.36			
07553543	**Low discrimination**					**07553543**
Actinomyces naeslundii		85.40	0.39	ARA 1		
Bifidobacterium adolescentis		14.60	0.10			
07553552	**Good identification**					**07553552**
Bifidobacterium adolescentis		96.79	0.46	XYL 87		
Actinomyces israelii		3.21	0.27	SOR 11		
07553560	**Low discrimination**					**07553560**
Actinomyces naeslundii		77.35	0.36	ARA 1		
Actinomyces israelii		22.65	0.17			
07553561	**Good identification**					**07553561**
Actinomyces naeslundii		100.0	0.39	ARA 1		
07553562	**Low discrimination**					**07553562**
Actinomyces naeslundii		58.97	0.36	ARA 1		
Actinomyces israelii		25.91	0.20			
Bifidobacterium adolescentis		15.12	0.10			
07553563	**Good identification**					**07553563**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	ARA 1			
07554000	**Low discrimination**						**07554000**
					COC	GRA	
Fusobacterium mortiferum		70.36	0.40	MLT 11	1	1	
Prevotella intermedia		11.43	0.15	XYL 7	1	1	
Gemella morbillorum		10.42	0.15		99	99	
07554100	**Good identification**						**07554100**
					GRA		
Fusobacterium mortiferum		96.34	0.46	MLT 11	1		
Lactobacillus fermentum		2.21	0.20	XYL 7	99		
Actinomyces israelii		1.46	0.17		99		
07554322	**Low discrimination**						**07554322**
Parabacteroides distasonis		56.93	0.43	GAL 99			
Prevotella oralis		35.39	0.27				
Clostridium clostridioforme		6.69	0.22				
07554332	**Low discrimination**						**07554332**
Parabacteroides distasonis		50.14	0.39	GAL 99			
Clostridium clostridioforme		49.86	0.33				
07554502	**Low discrimination**						**07554502**
Bifidobacterium adolescentis		75.14	0.41	GAL 99			
Clostridium clostridioforme		17.43	0.27				
Bifidobacterium longum subsp. longum		3.84	0.17				
07554512	**Low discrimination**						**07554512**
Bifidobacterium adolescentis		51.66	0.43	GAL 99			
Clostridium clostridioforme		44.67	0.38				
Bifidobacterium longum subsp. longum		3.68	0.21				
07554532	**Low discrimination**						**07554532**
					GRA		
Clostridium clostridioforme		69.58	0.38	SAL 88	50		
Bifidobacterium longum subsp. longum		22.91	0.32	TRE 88	99		
Parabacteroides distasonis		6.69	0.27	RHA 88	1		
07554612	**Good identification**						**07554612**
Clostridium clostridioforme		100.0	0.44	SAL 88			
07554632	**Good identification**						**07554632**
Clostridium clostridioforme		100.0	0.44	SAL 88			
07554702	**Good identification**						**07554702**
Clostridium clostridioforme		99.41	0.42	SAL 88			
Bifidobacterium adolescentis		0.59	0.07	TRE 88			
07554712	**Good identification**						**07554712**
Clostridium clostridioforme		99.84	0.53	SAL 88			
Bifidobacterium adolescentis		0.16	0.09	TRE 88			
07554720	**Good identification**						**07554720**
Bacteroides vulgatus		99.68	0.38	CEL 5			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.32	0.08	GAL 99			
07554722	**Low discrimination**						**07554722**
Clostridium clostridioforme		72.95	0.42	SAL 88			
Parabacteroides distasonis		24.28	0.39	TRE 88			
Bacteroides vulgatus		2.29	0.04				
07554732	**Good identification**						**07554732**
Clostridium clostridioforme		96.22	0.53	SAL 88			
Parabacteroides distasonis		3.79	0.35	TRE 88			
07555000	**Low discrimination**						**07555000**
					GRA		
Lactobacillus fermentum		52.90	0.50	CEL 1	99		
Fusobacterium mortiferum		24.48	0.43		1		
Actinomyces israelii		14.26	0.41		99		
07555001	**Low discrimination**						**07555001**
Actinomyces naeslundii		84.87	0.39	XYL 1			
Lactobacillus fermentum		15.13	0.16				
07555002	**Low discrimination**						**07555002**
					GRA		
Lactobacillus fermentum		51.68	0.50	CEL 1	99		
Actinomyces israelii		20.89	0.44		99		
Prevotella oralis		14.40	0.25		1		
07555003	**Low discrimination**						**07555003**
Actinomyces naeslundii		67.54	0.39	XYL 1			
Actinomyces viscosus		20.43	0.06				
Lactobacillus fermentum		12.04	0.16				
07555020	**Low discrimination**						**07555020**
					GRA		
Lactobacillus fermentum		37.27	0.42	CEL 1	99		
Prevotella oralis		31.15	0.25		1		
Prevotella melaninogenica		13.73	0.15		1		
07555021	**Good identification**						**07555021**
Actinomyces naeslundii		100.0	0.39	XYL 1			
07555022	**Low discrimination**						**07555022**
					CAT	GRA	
Prevotella oralis		76.27	0.47	XYL 11	1	1	
Bacteroides fragilis		14.21	0.25	LAC 94	99	1	
Lactobacillus fermentum		4.80	0.42		1	99	
07555023	**Low discrimination**						**07555023**
					GRA	CAT	
Actinomyces naeslundii		47.19	0.39	XYL 1	99	25	
Prevotella oralis		44.52	0.13		1	1	
Bacteroides fragilis		8.29	-.09		1	99	
07555042	**Low discrimination**						**07555042**
Actinomyces israelii		84.74	0.37	SAL 92			
Bifidobacterium adolescentis		6.22	0.12				
Lactobacillus fermentum		5.45	0.16				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07555100	**Low discrimination**						**07555100**
					GRA		
Lactobacillus fermentum		34.34	0.54	CEL 1	99		
Fusobacterium mortiferum		22.73	0.49		1		
Actinomyces israelii		22.69	0.52		99		
07555101	**Low discrimination**						**07555101**
					GRA		
Actinomyces naeslundii		80.82	0.47	XYL 1	99		
Lactobacillus fermentum		8.26	0.20		99		
Fusobacterium mortiferum		5.47	0.15		1		
07555102	**Low discrimination**						**07555102**
					GRA		
Bifidobacterium adolescentis		49.05	0.61	ARA 87	99		
Prevotella oralis		23.16	0.45		1		
Lactobacillus fermentum		8.66	0.54		99		
07555103	**Low discrimination**						**07555103**
					GRA		
Actinomyces naeslundii		43.35	0.47	XYL 1	99		
Bifidobacterium adolescentis		25.11	0.27		99		
Prevotella oralis		11.85	0.11		1		
07555110	**Low discrimination**						**07555110**
Actinomyces israelii		44.32	0.37	SAL 92			
Bifidobacterium adolescentis		31.41	0.29	MLZ 13			
Bifidobacterium longum subsp. longum		14.97	0.21				
07555112	**Good identification**						**07555112**
Bifidobacterium adolescentis		94.72	0.63	ARA 87			
Actinomyces israelii		2.03	0.40				
Clostridium clostridioforme		1.06	0.26				
07555120	**Low discrimination**						**07555120**
					CAT		
Prevotella melaninogenica		46.81	0.50	CEL 10	1		
Bacteroides fragilis		19.79	0.38	XYL 1	99		
Prevotella oralis		16.81	0.45		1		
07555121	**Low discrimination**						**07555121**
					GRA	CAT	
Actinomyces naeslundii		39.56	0.47	XYL 1	99	25	
Prevotella melaninogenica		30.12	0.15		1	1	
Bacteroides fragilis		12.73	0.04		1	99	
07555122	**Low discrimination**						**07555122**
					CAT		
Bacteroides fragilis		47.42	0.60	ESL 99	99		
Prevotella oralis		40.29	0.67		1		
Parabacteroides distasonis		9.26	0.69		76		
07555132	**Low discrimination**						**07555132**
					CAT		
Parabacteroides distasonis		79.63	0.65		76		
Bacteroides fragilis		7.65	0.25		99		
Prevotella oralis		6.50	0.33		1		
07555140	**Low discrimination**						**07555140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		85.95	0.45	SAL 92		
Actinomyces naeslundii		3.89	0.31			
Lactobacillus fermentum		3.38	0.20			
07555142	**Low discrimination**					**07555142**
					GRA	
Bifidobacterium adolescentis		60.31	0.46	ARA 87	99	
Actinomyces israelii		33.18	0.48	SOR 11	99	
Prevotella oralis		2.33	0.11		1	
07555152	**Acceptable identification**					**07555152**
Bifidobacterium adolescentis		93.70	0.47	ARA 87		
Actinomyces israelii		6.30	0.33	SOR 11		
07555222	**Low discrimination**					**07555222**
Prevotella oralis		77.78	0.41	XYL 11		
Parabacteroides distasonis		19.80	0.43	LAC 94		
Clostridium clostridioforme		2.09	0.21			
07555232	**Low discrimination**					**07555232**
Parabacteroides distasonis		50.90	0.39	LAC 99		
Clostridium clostridioforme		45.35	0.32			
Prevotella oralis		3.75	0.06			
07555302	**Low discrimination**					**07555302**
Prevotella oralis		64.65	0.39	XYL 11		
Parabacteroides distasonis		19.96	0.43	NAG 95		
Clostridium clostridioforme		7.04	0.30			
07555312	**Low discrimination**					**07555312**
					GRA	
Clostridium clostridioforme		66.75	0.41	SAL 88	50	
Parabacteroides distasonis		22.37	0.39	TRE 88	1	
Bifidobacterium adolescentis		8.21	0.29	ARA 94	99	
07555320	**Acceptable identification**					**07555320**
Bacteroides vulgatus		90.36	0.53	CEL 5		
Prevotella oralis		6.87	0.39	ARA 93		
Parabacteroides distasonis		2.12	0.43			
07555322	**Low discrimination**					**07555322**
					CAT	
Parabacteroides distasonis		60.66	0.77		76	
Prevotella oralis		37.71	0.61		1	
Bacteroides fragilis		1.05	0.25		99	
07555323	**Low discrimination**					**07555323**
					CAT	
Parabacteroides distasonis		61.02	0.43	URE 1	76	
Prevotella oralis		37.93	0.27		1	
Bacteroides fragilis		1.05	-.09		99	
07555330	**Low discrimination**					**07555330**
Parabacteroides distasonis		52.92	0.39	BGL 99		
Bacteroides vulgatus		42.27	0.18			
Prevotella oralis		3.22	0.05			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07555332	**Very good identification**					**07555332**
Parabacteroides distasonis		95.93	0.73			
Clostridium clostridioforme		2.89	0.41			
Prevotella oralis		1.12	0.27			
07555333	**Good identification**					**07555333**
Parabacteroides distasonis		97.08	0.39	URE 1		
Clostridium clostridioforme		2.93	0.07			
07555362	**Low discrimination**					**07555362**
					CAT	
Parabacteroides distasonis		61.02	0.43	SOR 1	76	
Prevotella oralis		37.93	0.27		1	
Bacteroides fragilis		1.05	-.09		99	
07555372	**Good identification**					**07555372**
Parabacteroides distasonis		97.08	0.39	SOR 1		
Clostridium clostridioforme		2.93	0.07			
07555400	**Low discrimination**					**07555400**
					SPO	
Lactobacillus fermentum		76.66	0.50	CEL 1	1	
Actinomyces israelii		16.90	0.40		1	
Clostridium butyricum		3.51	0.07		99	
07555402	**Low discrimination**					**07555402**
Lactobacillus fermentum		46.83	0.50	CEL 1		
Bifidobacterium adolescentis		29.27	0.41			
Actinomyces israelii		15.49	0.43			
07555412	**Low discrimination**					**07555412**
Bifidobacterium adolescentis		51.98	0.43	LAC 99		
Clostridium clostridioforme		40.28	0.38			
Bifidobacterium longum subsp. longum		3.70	0.21			
07555422	**Low discrimination**					**07555422**
Lactobacillus fermentum		41.67	0.42	CEL 1		
Clostridium clostridioforme		16.25	0.26			
Bifidobacterium longum subsp. longum		15.97	0.28			
07555432	**Low discrimination**					**07555432**
					GRA	
Clostridium clostridioforme		67.21	0.38	SAL 88	50	
Bifidobacterium longum subsp. longum		24.70	0.32	TRE 88	99	
Parabacteroides distasonis		7.21	0.27	RHA 88	1	
07555442	**Low discrimination**					**07555442**
Actinomyces israelii		59.55	0.36	SAL 92		
Bifidobacterium adolescentis		35.77	0.26			
Lactobacillus fermentum		4.68	0.16			
07555500	**Low discrimination**					**07555500**
					SPO	
Bifidobacterium longum subsp. longum		40.67	0.52	CEL 1	1	
Lactobacillus fermentum		20.98	0.54		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		14.11	0.31		99		
07555502	**Low discrimination**						**07555502**
Bifidobacterium adolescentis		89.57	0.75				
Bifidobacterium longum subsp. longum		4.58	0.52				
Lactobacillus fermentum		2.36	0.54				
07555503	**Acceptable identification**						**07555503**
Bifidobacterium adolescentis		91.01	0.41	URE 1			
Bifidobacterium longum subsp. longum		4.65	0.17				
Lactobacillus fermentum		2.40	0.20				
07555510	**Low discrimination**						**07555510**
Bifidobacterium longum subsp. longum		84.47	0.56	CEL 1			
Bifidobacterium adolescentis		11.99	0.43				
Actinomyces israelii		2.07	0.36				
07555512	**Good identification**						**07555512**
Bifidobacterium adolescentis		90.80	0.77				
Bifidobacterium longum subsp. longum		6.46	0.56				
Clostridium clostridioforme		2.38	0.47				
07555513	**Acceptable identification**						**07555513**
Bifidobacterium adolescentis		91.13	0.43	URE 1			
Bifidobacterium longum subsp. longum		6.49	0.21				
Clostridium clostridioforme		2.39	0.12				
07555520	**Low discrimination**						**07555520**
					GRA		
Bacteroides vulgatus		72.54	0.56	CEL 5	1		
Bifidobacterium longum subsp. longum		25.27	0.62	RHA 90	99		
Lactobacillus fermentum		1.09	0.46		99		
07555522	**Low discrimination**						**07555522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		43.58	0.62	CEL 1	99	1	
Parabacteroides distasonis		40.24	0.65		1	76	
Bacteroides fragilis		3.40	0.25		1	99	
07555530	**Very good identification**						**07555530**
					GRA		
Bifidobacterium longum subsp. longum		97.88	0.66	CEL 1	99		
Bacteroides vulgatus		1.67	0.21		1		
Parabacteroides distasonis		0.29	0.27		1		
07555532	**Low discrimination**						**07555532**
					GRA		
Bifidobacterium longum subsp. longum		70.17	0.66	CEL 1	99		
Parabacteroides distasonis		20.49	0.61		1		
Clostridium clostridioforme		6.46	0.47		50		
07555540	**Low discrimination**						**07555540**
					SPO		
Actinomyces israelii		60.80	0.43	SAL 92	1		
Clostridium butyricum		16.92	0.12		99		
Bifidobacterium adolescentis		13.70	0.26		1		
07555542	**Good identification**						**07555542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium adolescentis		92.73	0.60	SOR 11	1		
Actinomyces israelii		6.24	0.46		1		
Clostridium butyricum		0.45	0.05		99		
07555552	**Very good identification**						**07555552**
Bifidobacterium adolescentis		98.41	0.61	SOR 11			
Actinomyces israelii		0.81	0.32				
Bifidobacterium longum subsp. longum		0.57	0.21				
07555602	**Good identification**						**07555602**
Clostridium clostridioforme		95.85	0.41	SAL 88			
Actinomyces israelii		2.50	0.24	TRE 88			
Lactobacillus fermentum		1.02	0.16				
07555612	**Good identification**						**07555612**
Clostridium clostridioforme		99.82	0.52	SAL 88			
Bifidobacterium adolescentis		0.18	0.09	TRE 88			
07555620	**Good identification**						**07555620**
Bacteroides vulgatus		99.71	0.38	CEL 5			
Clostridium clostridioforme		0.29	0.07	LAC 99			
07555622	**Low discrimination**						**07555622**
Clostridium clostridioforme		69.19	0.41	SAL 88			
Parabacteroides distasonis		25.70	0.39	TRE 88			
Bacteroides vulgatus		2.42	0.04				
07555632	**Good identification**						**07555632**
Clostridium clostridioforme		95.79	0.52	SAL 88			
Parabacteroides distasonis		4.21	0.35	TRE 88			
07555700	**Acceptable identification**						**07555700**
					GRA	SPO	
Bacteroides vulgatus		94.63	0.38	CEL 5	1	1	
Actinomyces israelii		1.90	0.31	NAG 99	99	1	
Clostridium butyricum		0.97	0.05		99	99	
07555702	**Low discrimination**						**07555702**
					GRA		
Clostridium clostridioforme		72.68	0.50	SAL 88	50		
Bifidobacterium adolescentis		14.24	0.41	TRE 88	99		
Parabacteroides distasonis		8.06	0.39		1		
07555712	**Good identification**						**07555712**
					GRA		
Clostridium clostridioforme		93.42	0.61	SAL 88	50		
Bifidobacterium adolescentis		4.91	0.43	TRE 88	99		
Parabacteroides distasonis		1.23	0.35		1		
07555720	**Very good identification**						**07555720**
					GRA		
Bacteroides vulgatus		99.84	0.72	CEL 5	1		
Parabacteroides distasonis		0.11	0.39		1		
Bifidobacterium longum subsp. longum		0.04	0.28		99		
07555721	**Good identification**						**07555721**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		100.0	0.38	CEL 5		
07555722	**Low discrimination**					**07555722**
Parabacteroides distasonis		80.66	0.74			
Bacteroides vulgatus		7.60	0.38			
Clostridium clostridioforme		7.34	0.50			
07555723	**Low discrimination**					**07555723**
Parabacteroides distasonis		82.98	0.39	URE 1		
Bacteroides vulgatus		7.82	0.04			
Clostridium clostridioforme		7.56	0.16			
07555730	**Low discrimination**					**07555730**
					GRA	
Bacteroides vulgatus		85.61	0.38	CEL 5	1	
Bifidobacterium longum subsp. longum		5.64	0.32	MLZ 1	99	
Parabacteroides distasonis		4.94	0.35		1	
07555732	**Low discrimination**					**07555732**
Parabacteroides distasonis		55.14	0.69			
Clostridium clostridioforme		42.47	0.61			
Bacteroides ovatus		0.84	0.22			
07555760	**Good identification**					**07555760**
Bacteroides vulgatus		100.0	0.38	CEL 5		
07555762	**Low discrimination**					**07555762**
Parabacteroides distasonis		79.63	0.39	SOR 1		
Bacteroides vulgatus		7.50	0.04			
Clostridium clostridioforme		7.25	0.16			
07556502	**Low discrimination**					**07556502**
Bifidobacterium adolescentis		87.10	0.40	GAL 99		
Actinomyces israelii		12.90	0.32			
07556512	**Good identification**					**07556512**
Bifidobacterium adolescentis		97.20	0.41	GAL 99		
Actinomyces israelii		1.76	0.18			
Clostridium clostridioforme		1.04	0.04			
07557000	**Acceptable identification**					**07557000**
					CAT	
Actinomyces israelii		93.76	0.54	SAL 92	1	
Actinomyces naeslundii		4.50	0.41		25	
Propionibacterium granulosum		1.11	0.16		90	
07557001	**Low discrimination**					**07557001**
Actinomyces naeslundii		87.70	0.44	XYL 1		
Actinomyces israelii		12.30	0.20			
07557002	**Good identification**					**07557002**
Actinomyces israelii		93.48	0.57	SAL 92		
Bifidobacterium adolescentis		3.12	0.26			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		2.99	0.41			
07557003	**Low discrimination**					**07557003**
Actinomyces naeslundii		82.62	0.44	XYL 1		
Actinomyces israelii		17.38	0.23			
07557010	**Good identification**					**07557010**
Actinomyces israelii		100.0	0.40	SAL 92		
07557012	**Low discrimination**					**07557012**
Actinomyces israelii		78.56	0.43	SAL 92		
Bifidobacterium adolescentis		21.44	0.27	MLZ 13		
07557021	**Good identification**					**07557021**
Actinomyces naeslundii		100.0	0.44	XYL 1		
07557022	**Low discrimination**					**07557022**
					GRA	
Actinomyces israelii		51.67	0.37	SAL 92	99	
Actinomyces naeslundii		25.92	0.41	NAG 6	99	
Prevotella oralis		18.90	0.13		1	
07557023	**Good identification**					**07557023**
Actinomyces naeslundii		100.0	0.44	XYL 1		
07557040	**Good identification**					**07557040**
Actinomyces israelii		97.87	0.47	SAL 92		
Actinomyces naeslundii		2.13	0.28			
07557042	**Good identification**					**07557042**
Actinomyces israelii		97.55	0.50	SAL 92		
Actinomyces naeslundii		1.42	0.28			
Bifidobacterium adolescentis		1.03	0.10			
07557052	**Acceptable identification**					**07557052**
Actinomyces israelii		92.02	0.36	SAL 92		
Bifidobacterium adolescentis		7.98	0.12	MLZ 13		
07557100	**Good identification**					**07557100**
Actinomyces israelii		94.93	0.64	SAL 92		
Actinomyces naeslundii		3.24	0.49			
Bifidobacterium adolescentis		1.19	0.26			
07557101	**Low discrimination**					**07557101**
Actinomyces naeslundii		83.54	0.52	XYL 1		
Actinomyces israelii		16.46	0.30			
07557102	**Low discrimination**					**07557102**
Actinomyces israelii		53.85	0.67	SAL 92		
Bifidobacterium adolescentis		44.47	0.60			
Actinomyces naeslundii		1.23	0.49			
07557103	**Low discrimination**					**07557103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		64.95	0.52	XYL 1			
Actinomyces israelii		19.20	0.33				
Bifidobacterium adolescentis		15.85	0.26				
07557110	**Low discrimination**						**07557110**
Actinomyces israelii		89.74	0.50	SAL 92			
Bifidobacterium adolescentis		9.18	0.27	MLZ 13			
Actinomyces naeslundii		1.08	0.27				
07557112	**Low discrimination**						**07557112**
Bifidobacterium adolescentis		86.96	0.61	ARA 87			
Actinomyces israelii		12.88	0.53				
Actinomyces naeslundii		0.10	0.27				
07557120	**Low discrimination**						**07557120**
					GRA		
Actinomyces israelii		46.14	0.44	SAL 92	99		
Actinomyces naeslundii		24.70	0.49	NAG 6	99		
Prevotella melaninogenica		14.53	0.15		1		
07557121	**Good identification**						**07557121**
Actinomyces naeslundii		100.0	0.52	XYL 1			
07557123	**Acceptable identification**						**07557123**
					GRA	CAT	
Actinomyces naeslundii		94.44	0.52	XYL 1	99	25	
Bacteroides fragilis		3.00	-.09		1	99	
Prevotella oralis		2.55	-.01		1	1	
07557140	**Very good identification**						**07557140**
Actinomyces israelii		98.09	0.57	SAL 92			
Actinomyces naeslundii		1.52	0.36				
Bifidobacterium adolescentis		0.39	0.10				
07557141	**Low discrimination**						**07557141**
Actinomyces naeslundii		69.73	0.39	XYL 1			
Actinomyces israelii		30.28	0.23				
07557142	**Low discrimination**						**07557142**
Actinomyces israelii		78.46	0.60	SAL 92			
Bifidobacterium adolescentis		20.59	0.44				
Actinomyces naeslundii		0.81	0.36				
07557143	**Low discrimination**						**07557143**
Actinomyces naeslundii		54.88	0.39	XYL 1			
Actinomyces israelii		35.74	0.26				
Bifidobacterium adolescentis		9.38	0.10				
07557150	**Good identification**						**07557150**
Actinomyces israelii		96.85	0.43	SAL 92			
Bifidobacterium adolescentis		3.15	0.12	MLZ 13			
07557152	**Low discrimination**						**07557152**
Bifidobacterium adolescentis		68.22	0.46	ARA 87			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		31.78	0.46	SOR 11			
07557160	**Low discrimination**						**07557160**
Actinomyces israelii		80.45	0.37	SAL 92			
Actinomyces naeslundii		19.55	0.36	NAG 6			
07557161	**Good identification**						**07557161**
Actinomyces naeslundii		100.0	0.39	XYL 1			
07557162	**Low discrimination**						**07557162**
					GRA	CAT	
Actinomyces israelii		77.85	0.40	SAL 92	99	1	
Actinomyces naeslundii		12.61	0.36	NAG 6	99	25	
Bacteroides fragilis		3.41	-.09		1	99	
07557163	**Good identification**						**07557163**
Actinomyces naeslundii		100.0	0.39	XYL 1			
07557202	**Good identification**						**07557202**
Actinomyces israelii		100.0	0.38	SAL 92			
07557300	**Good identification**						**07557300**
Actinomyces israelii		100.0	0.45	SAL 92			
07557302	**Low discrimination**						**07557302**
					GRA		
Actinomyces israelii		87.86	0.48	SAL 92	99		
Bifidobacterium adolescentis		9.74	0.26	RHA 7	99		
Prevotella oralis		2.41	0.05		1		
07557322	**Low discrimination**						**07557322**
					GRA		
Parabacteroides distasonis		57.01	0.43	MAN 1	1		
Prevotella oralis		35.44	0.27		1		
Actinomyces israelii		4.35	0.28		99		
07557332	**Good identification**						**07557332**
					GRA		
Parabacteroides distasonis		95.12	0.39	MAN 1	1		
Clostridium clostridioforme		2.87	0.07		50		
Actinomyces israelii		2.01	0.14		99		
07557340	**Good identification**						**07557340**
Actinomyces israelii		100.0	0.38	SAL 92			
07557342	**Good identification**						**07557342**
Actinomyces israelii		96.60	0.41	SAL 92			
Bifidobacterium adolescentis		3.40	0.10	RHA 7			
07557400	**Good identification**						**07557400**
Actinomyces israelii		99.20	0.53	SAL 92			
Lactobacillus fermentum		0.80	0.16				
07557402	**Low discrimination**						**07557402**
Actinomyces israelii		78.23	0.56	SAL 92			
Bifidobacterium adolescentis		21.35	0.40				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		0.42	0.16			
07557410	**Good identification**					**07557410**
Actinomyces israelii		96.73	0.38	SAL 92		
Bifidobacterium adolescentis		3.27	0.07	MLZ 13		
07557412	**Low discrimination**					**07557412**
Bifidobacterium adolescentis		69.06	0.41	LAC 99		
Actinomyces israelii		30.94	0.42			
07557422	**Good identification**					**07557422**
Actinomyces israelii		100.0	0.35	SAL 92		
07557440	**Good identification**					**07557440**
Actinomyces israelii		100.0	0.46	SAL 92		
07557442	**Acceptable identification**					**07557442**
Actinomyces israelii		92.02	0.49	SAL 92		
Bifidobacterium adolescentis		7.98	0.24			
07557500	**Low discrimination**					**07557500**
					SPO	
Actinomyces israelii		88.00	0.63	SAL 92	1	
Bifidobacterium adolescentis		9.01	0.40		1	
Clostridium butyricum		2.15	0.14		99	
07557502	**Low discrimination**					**07557502**
					SPO	
Bifidobacterium adolescentis		86.96	0.74		1	
Actinomyces israelii		12.88	0.66		1	
Clostridium butyricum		0.08	0.07		99	
07557503	**Low discrimination**					**07557503**
Bifidobacterium adolescentis		87.10	0.40	URE 1		
Actinomyces israelii		12.90	0.32			
07557510	**Low discrimination**					**07557510**
Actinomyces israelii		52.12	0.49	SAL 92		
Bifidobacterium adolescentis		43.63	0.41	MLZ 13		
Bifidobacterium longum subsp. longum		3.80	0.21			
07557512	**Very good identification**					**07557512**
Bifidobacterium adolescentis		98.09	0.75			
Actinomyces israelii		1.78	0.52			
Bifidobacterium longum subsp. longum		0.09	0.21			
07557513	**Good identification**					**07557513**
Bifidobacterium adolescentis		98.22	0.41	URE 1		
Actinomyces israelii		1.78	0.18			
07557522	**Low discrimination**					**07557522**
Bifidobacterium adolescentis		40.54	0.40	NAG 1		
Actinomyces israelii		37.93	0.45			
Bifidobacterium longum subsp. longum		10.13	0.28			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07557532	**Low discrimination**					**07557532**
Bifidobacterium adolescentis		61.43	0.41	NAG 1		
Bifidobacterium longum subsp. longum		21.38	0.32			
Actinomyces israelii		7.03	0.31			
07557540	**Good identification**					**07557540**
					SPO	
Actinomyces israelii		96.34	0.56	SAL 92	1	
Bifidobacterium adolescentis		3.13	0.24		1	
Clostridium butyricum		0.53	-.04		99	
07557542	**Low discrimination**					**07557542**
Bifidobacterium adolescentis		68.22	0.58	SOR 11		
Actinomyces israelii		31.78	0.59			
07557550	**Low discrimination**					**07557550**
					COC	
Actinomyces israelii		76.68	0.42	SAL 92	1	
Bifidobacterium adolescentis		20.40	0.26	MLZ 13	1	
Blautia producta		2.92	-.03		99	
07557552	**Good identification**					**07557552**
Bifidobacterium adolescentis		94.61	0.60	SOR 11		
Actinomyces israelii		5.39	0.45			
07557560	**Good identification**					**07557560**
Actinomyces israelii		100.0	0.35	SAL 92		
07557562	**Low discrimination**					**07557562**
Actinomyces israelii		74.64	0.38	SAL 92		
Bifidobacterium adolescentis		25.36	0.24	NAG 6		
07557602	**Acceptable identification**					**07557602**
Actinomyces israelii		93.61	0.36	SAL 92		
Clostridium clostridioforme		6.39	0.07	RHA 7		
07557700	**Acceptable identification**					**07557700**
					GRA	
Actinomyces israelii		91.86	0.44	SAL 92	99	
Bacteroides vulgatus		8.14	0.04	RHA 7	1	
07557702	**Low discrimination**					**07557702**
Actinomyces israelii		50.93	0.47	SAL 92		
Bifidobacterium adolescentis		46.16	0.40	RHA 7		
Clostridium clostridioforme		2.91	0.16			
07557712	**Low discrimination**					**07557712**
Bifidobacterium adolescentis		73.01	0.41	RHA 1		
Clostridium clostridioforme		17.15	0.27			
Actinomyces israelii		9.85	0.33			
07557720	**Good identification**					**07557720**
					GRA	
Bacteroides vulgatus		99.28	0.38	CEL 5	1	
Actinomyces israelii		0.72	0.23	MAN 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07557722	**Low discrimination**					**07557722**
					GRA	
Parabacteroides distasonis		58.38	0.39	MAN 1	1	
Bacteroides ovatus		23.70	0.17		1	
Actinomyces israelii		5.94	0.26		99	
07557740	**Good identification**					**07557740**
Actinomyces israelii		100.0	0.37	SAL 92		
07557742	**Low discrimination**					**07557742**
Actinomyces israelii		77.64	0.40	SAL 92		
Bifidobacterium adolescentis		22.37	0.24	RHA 7		
07560100	**Low discrimination**					**07560100**
					GRA	
Actinomyces naeslundii		61.10	0.37	FRU 95	99	
Fusobacterium mortiferum		38.90	0.22	GAL 96	1	
07560101	**Good identification**					**07560101**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07560103	**Low discrimination**					**07560103**
Actinomyces naeslundii		77.71	0.40	FRU 95		
Actinomyces viscosus		22.29	0.07	GAL 96		
07560120	**Low discrimination**					**07560120**
					GRA	
Actinomyces naeslundii		76.86	0.37	FRU 95	99	
Prevotella melaninogenica		23.14	-.01	GAL 96	1	
07560121	**Good identification**					**07560121**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07560122	**Good identification**					**07560122**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07560123	**Good identification**					**07560123**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07561000	**Low discrimination**					**07561000**
					CAT	
Actinomyces naeslundii		77.42	0.53	FRU 95	25	
Propionibacterium granulosum		14.62	0.26		90	
Lactobacillus jensenii		4.21	0.10		1	
07561001	**Good identification**					**07561001**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07561002	**Low discrimination**					**07561002**
Actinomyces naeslundii		86.10	0.53	FRU 95		
Actinomyces viscosus		4.54	0.07			
Propionibacterium granulosum		4.06	0.15			
07561003	**Good identification**					**07561003**
Actinomyces naeslundii		98.84	0.56	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		1.16	-.01				
07561020	**Low discrimination**						**07561020**
					CAT		
Actinomyces naeslundii		82.95	0.53	FRU 95	25		
Clostridium perfringens		10.17	0.15		1		
Propionibacterium granulosum		3.92	0.15		90		
07561021	**Good identification**						**07561021**
Actinomyces naeslundii		100.0	0.56	FRU 95			
07561022	**Low discrimination**						**07561022**
					GRA		
Actinomyces naeslundii		86.56	0.53	FRU 95	99		
Prevotella oralis		6.26	0.08		1		
Clostridium perfringens		4.99	0.10		99		
07561023	**Good identification**						**07561023**
Actinomyces naeslundii		100.0	0.56	FRU 95			
07561040	**Good identification**						**07561040**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07561041	**Good identification**						**07561041**
Actinomyces naeslundii		100.0	0.43	FRU 95			
07561042	**Good identification**						**07561042**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07561043	**Good identification**						**07561043**
Actinomyces naeslundii		100.0	0.43	FRU 95			
07561060	**Low discrimination**						**07561060**
Actinomyces naeslundii		88.31	0.40	FRU 95			
Clostridium perfringens		11.69	0.03				
07561061	**Good identification**						**07561061**
Actinomyces naeslundii		100.0	0.43	FRU 95			
07561062	**Good identification**						**07561062**
Actinomyces naeslundii		100.0	0.40	FRU 95			
07561063	**Good identification**						**07561063**
Actinomyces naeslundii		100.0	0.43	FRU 95			
07561100	**Low discrimination**						**07561100**
					COC	GRA	
Actinomyces naeslundii		88.34	0.61	FRU 95	1	99	
Atopobium parvulum		6.17	0.18		99	99	
Fusobacterium mortiferum		3.52	0.25		1	1	
07561101	**Very good identification**						**07561101**
					SPO		
Actinomyces naeslundii		99.38	0.64	FRU 95	1		
Clostridium chauvoei		0.62	0.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07561102	**Low discrimination**					**07561102**
Actinomyces naeslundii		84.97	0.61	FRU 95		
Lactobacillus acidophilus		8.68	0.19			
Actinomyces viscosus		2.80	0.11			
07561103	**Very good identification**					**07561103**
Actinomyces naeslundii		99.27	0.64	FRU 95		
Actinomyces viscosus		0.73	0.03			
07561110	**Good identification**					**07561110**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07561111	**Good identification**					**07561111**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07561112	**Low discrimination**					**07561112**
					GRA	
Actinomyces naeslundii		75.41	0.39	FRU 95	99	
Bifidobacterium adolescentis		14.98	0.11	MLZ 5	99	
Leptotrichia buccalis		9.61	-.11		1	
07561113	**Good identification**					**07561113**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07561121	**Very good identification**					**07561121**
					GRA	
Actinomyces naeslundii		99.17	0.64	FRU 95	99	
Prevotella melaninogenica		0.83	-.01		1	
07561122	**Low discrimination**					**07561122**
					GRA	
Actinomyces naeslundii		49.56	0.61	FRU 95	99	
Prevotella oralis		19.72	0.28		1	
Prevotella melaninogenica		11.72	0.21		1	
07561123	**Very good identification**					**07561123**
Actinomyces naeslundii		100.0	0.64	FRU 95		
07561130	**Low discrimination**					**07561130**
					GRA	
Actinomyces naeslundii		80.75	0.39	FRU 95	99	
Prevotella melaninogenica		19.25	-.01	MLZ 5	1	
07561131	**Good identification**					**07561131**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07561132	**Low discrimination**					**07561132**
					GRA	
Actinomyces naeslundii		52.17	0.39	FRU 95	99	
Parabacteroides distasonis		47.83	0.25	MLZ 5	1	
07561133	**Good identification**					**07561133**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07561140	**Good identification**					**07561140**
Actinomyces naeslundii		100.0	0.48	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07561141	**Good identification**					**07561141**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07561142	**Acceptable identification**					**07561142**
Actinomyces naeslundii		93.32	0.48	FRU 95		
Lactobacillus acidophilus		6.68	0.04			
07561143	**Good identification**					**07561143**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07561160	**Low discrimination**					**07561160**
					GRA	
Actinomyces naeslundii		70.82	0.48	FRU 95	99	
Clostridium perfringens		24.15	0.18		99	
Prevotella melaninogenica		5.03	-.01		1	
07561161	**Good identification**					**07561161**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07561162	**Low discrimination**					**07561162**
Actinomyces naeslundii		81.17	0.48	FRU 95		
Clostridium perfringens		13.02	0.12			
Lactobacillus acidophilus		5.81	0.04			
07561163	**Good identification**					**07561163**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07561322	**Low discrimination**					**07561322**
					SPO	
Parabacteroides distasonis		52.28	0.38	FRU 99	1	
Prevotella oralis		33.16	0.22	XYL 90	1	
Clostridium ramosum		11.10	0.14		92	
07562001	**Good identification**					**07562001**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07562003	**Good identification**					**07562003**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07562021	**Good identification**					**07562021**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07562023	**Good identification**					**07562023**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07562100	**Good identification**					**07562100**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07562101	**Good identification**					**07562101**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07562102	**Good identification**					**07562102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	FRU 95		
07562103	**Good identification**					**07562103**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07562120	**Good identification**					**07562120**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07562121	**Good identification**					**07562121**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07562122	**Good identification**					**07562122**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07562123	**Good identification**					**07562123**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07563000	**Good identification**					**07563000**
Actinomyces naeslundii		93.16	0.58	FRU 95		
Propionibacterium granulosum		6.84	0.24			
07563001	**Very good identification**					**07563001**
Actinomyces naeslundii		100.0	0.61	FRU 95		
07563002	**Very good identification**					**07563002**
					CAT	
Actinomyces naeslundii		97.23	0.58	FRU 95	25	
Propionibacterium granulosum		1.78	0.13		90	
Actinomyces israelii		0.99	0.15		1	
07563003	**Very good identification**					**07563003**
Actinomyces naeslundii		100.0	0.61	FRU 95		
07563010	**Low discrimination**					**07563010**
Actinomyces naeslundii		84.03	0.36	FRU 95		
Propionibacterium granulosum		15.97	0.09	MLZ 5		
07563011	**Good identification**					**07563011**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07563012	**Good identification**					**07563012**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07563013	**Good identification**					**07563013**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07563020	**Very good identification**					**07563020**
Actinomyces naeslundii		98.20	0.58	FRU 95		
Propionibacterium granulosum		1.80	0.13			
07563021	**Very good identification**					**07563021**
Actinomyces naeslundii		100.0	0.61	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07563022	**Very good identification**					**07563022**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07563023	**Very good identification**					**07563023**
Actinomyces naeslundii		100.0	0.61	FRU 95		
07563030	**Good identification**					**07563030**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07563031	**Good identification**					**07563031**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07563032	**Good identification**					**07563032**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07563033	**Good identification**					**07563033**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07563040	**Good identification**					**07563040**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07563041	**Good identification**					**07563041**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07563042	**Good identification**					**07563042**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07563043	**Good identification**					**07563043**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07563060	**Good identification**					**07563060**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07563061	**Good identification**					**07563061**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07563062	**Good identification**					**07563062**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07563063	**Good identification**					**07563063**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07563100	**Very good identification**					**07563100**
Actinomyces naeslundii		99.05	0.66	FRU 95		
Actinomyces israelii		0.95	0.22			
07563101	**Very good identification**					**07563101**
Actinomyces naeslundii		100.0	0.69	FRU 95		
07563102	**Very good identification**					**07563102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		98.24	0.66	FRU 95			
Actinomyces israelii		1.41	0.25				
Bifidobacterium adolescentis		0.35	0.08				
07563103	**Very good identification**						**07563103**
Actinomyces naeslundii		100.0	0.69	FRU 95			
07563110	**Good identification**						**07563110**
Actinomyces naeslundii		100.0	0.44	FRU 95			
07563111	**Good identification**						**07563111**
Actinomyces naeslundii		100.0	0.47	FRU 95			
07563112	**Acceptable identification**						**07563112**
Actinomyces naeslundii		92.27	0.44	FRU 95			
Bifidobacterium adolescentis		7.73	0.10	MLZ 5			
07563113	**Good identification**						**07563113**
Actinomyces naeslundii		100.0	0.47	FRU 95			
07563120	**Very good identification**						**07563120**
					SPO	GRA	
Actinomyces naeslundii		96.35	0.66	FRU 95	1	99	
Clostridium ramosum		3.03	0.15		92	50	
Prevotella melaninogenica		0.62	-.01		1	1	
07563121	**Very good identification**						**07563121**
Actinomyces naeslundii		100.0	0.69	FRU 95			
07563122	**Low discrimination**						**07563122**
					SPO		
Actinomyces naeslundii		88.84	0.66	FRU 95	1		
Clostridium ramosum		11.16	0.25		92		
07563123	**Very good identification**						**07563123**
Actinomyces naeslundii		100.0	0.69	FRU 95			
07563130	**Good identification**						**07563130**
Actinomyces naeslundii		100.0	0.44	FRU 95			
07563131	**Good identification**						**07563131**
Actinomyces naeslundii		100.0	0.47	FRU 95			
07563132	**Good identification**						**07563132**
Actinomyces naeslundii		100.0	0.44	FRU 95			
07563133	**Good identification**						**07563133**
Actinomyces naeslundii		100.0	0.47	FRU 95			
07563140	**Good identification**						**07563140**
Actinomyces naeslundii		97.94	0.53	FRU 95			
Actinomyces israelii		2.06	0.15				
07563141	**Good identification**						**07563141**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.56	FRU 95			
07563142	**Good identification**						**07563142**
Actinomyces naeslundii		96.94	0.53	FRU 95			
Actinomyces israelii		3.06	0.18				
07563143	**Good identification**						**07563143**
Actinomyces naeslundii		100.0	0.56	FRU 95			
07563160	**Good identification**						**07563160**
Actinomyces naeslundii		100.0	0.53	FRU 95			
07563161	**Good identification**						**07563161**
Actinomyces naeslundii		100.0	0.56	FRU 95			
07563162	**Good identification**						**07563162**
Actinomyces naeslundii		100.0	0.53	FRU 95			
07563163	**Good identification**						**07563163**
Actinomyces naeslundii		100.0	0.56	FRU 95			
07565122	**Acceptable identification**						**07565122**
					GRA		
Parabacteroides distasonis		92.18	0.46	FRU 99	1		
Prevotella oralis		5.62	0.13		1		
Actinomyces naeslundii		1.16	0.27		99		
07565132	**Good identification**						**07565132**
Parabacteroides distasonis		98.06	0.42	FRU 99			
Clostridium clostridioforme		1.95	0.07				
07565322	**Good identification**						**07565322**
Parabacteroides distasonis		98.58	0.54	FRU 99			
Prevotella oralis		0.86	0.06				
Clostridium clostridioforme		0.57	0.10				
07565332	**Good identification**						**07565332**
Parabacteroides distasonis		95.37	0.50	FRU 99			
Clostridium clostridioforme		4.63	0.22				
07565502	**Low discrimination**						**07565502**
					SPO		
Bifidobacterium adolescentis		63.66	0.38	FRU 99	1		
Bifidobacterium longum subsp. longum		13.42	0.25		1		
Clostridium butyricum		10.32	0.10		99		
07565512	**Low discrimination**						**07565512**
Bifidobacterium adolescentis		62.63	0.40	FRU 99			
Bifidobacterium longum subsp. longum		18.39	0.29				
Clostridium clostridioforme		18.05	0.27				
07565520	**Acceptable identification**						**07565520**
					GRA	SPO	
Bifidobacterium longum subsp. longum		92.04	0.35	FRU 90	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		4.00	-.02	CEL 1	1	1	
Clostridium butyricum		3.96	-.01		99	99	
07565522	**Low discrimination**						**07565522**
					GRA		
Parabacteroides distasonis		57.91	0.43	FRU 99	1		
Bifidobacterium longum subsp. longum		37.86	0.35		99		
Clostridium clostridioforme		3.47	0.16		50		
07565530	**Good identification**						**07565530**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
07565532	**Low discrimination**						**07565532**
					GRA		
Bifidobacterium longum subsp. longum		57.83	0.39	FRU 90	99		
Parabacteroides distasonis		27.98	0.38	CEL 1	1		
Clostridium clostridioforme		14.19	0.27		50		
07565712	**Good identification**						**07565712**
Clostridium clostridioforme		99.19	0.42	SAL 88			
Parabacteroides distasonis		0.81	0.12	FRU 99			
07565722	**Low discrimination**						**07565722**
Parabacteroides distasonis		81.18	0.51	FRU 99			
Clostridium clostridioforme		11.90	0.31				
Bacteroides thetaiotaomicron		4.71	0.14				
07565732	**Low discrimination**						**07565732**
Clostridium clostridioforme		53.55	0.42	SAL 88			
Parabacteroides distasonis		43.18	0.46	FRU 99			
Bacteroides thetaiotaomicron		1.81	0.07				
07567100	**Acceptable identification**						**07567100**
Actinomyces israelii		91.59	0.40	SAL 92			
Actinomyces naeslundii		8.41	0.32	FRU 99			
07567102	**Low discrimination**						**07567102**
Actinomyces israelii		82.81	0.43	SAL 92			
Bifidobacterium adolescentis		12.12	0.23	FRU 99			
Actinomyces naeslundii		5.07	0.32				
07567142	**Good identification**						**07567142**
Actinomyces israelii		95.56	0.36	SAL 92			
Bifidobacterium adolescentis		4.44	0.07	FRU 99			
07567500	**Acceptable identification**						**07567500**
					SPO		
Actinomyces israelii		90.78	0.39	SAL 92	1		
Clostridium butyricum		9.22	0.01	FRU 99	99		
07567502	**Low discrimination**						**07567502**
					SPO		
Bifidobacterium adolescentis		53.84	0.37	FRU 99	1		
Actinomyces israelii		44.98	0.42		1		
Clostridium butyricum		1.18	-.07		99		
07567512	**Low discrimination**						**07567512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		88.57	0.38	FRU 99			
Actinomyces israelii		9.05	0.28				
Bifidobacterium dentium		2.38	-.07				
07570000	**Low discrimination**						**07570000**
					GRA	CAT	
Propionibacterium granulosum		81.58	0.60	CEL 1	99	90	
Fusobacterium mortiferum		13.98	0.50		1	5	
Actinomyces naeslundii		3.46	0.51		99	25	
07570001	**Low discrimination**						**07570001**
					GRA	CAT	
Actinomyces naeslundii		84.30	0.54	GAL 96	99	25	
Propionibacterium granulosum		13.40	0.26		99	90	
Fusobacterium mortiferum		2.30	0.16		1	5	
07570002	**Low discrimination**						**07570002**
Propionibacterium granulosum		62.16	0.50	CEL 1			
Actinomyces viscosus		21.76	0.33				
Actinomyces naeslundii		10.53	0.51				
07570003	**Low discrimination**						**07570003**
Actinomyces naeslundii		66.71	0.54	GAL 96			
Actinomyces viscosus		30.63	0.24				
Propionibacterium granulosum		2.65	0.15				
07570010	**Good identification**						**07570010**
					GRA	CAT	
Propionibacterium granulosum		96.51	0.45	CEL 1	99	90	
Actinomyces naeslundii		1.58	0.29	MLZ 12	99	25	
Fusobacterium mortiferum		1.23	0.16		1	5	
07570020	**Low discrimination**						**07570020**
					CAT		
Propionibacterium granulosum		83.16	0.50	CEL 1	90		
Actinomyces naeslundii		14.09	0.51		25		
Clostridium perfringens		2.18	0.15		1		
07570021	**Good identification**						**07570021**
Actinomyces naeslundii		96.18	0.54	GAL 96			
Propionibacterium granulosum		3.82	0.15				
07570022	**Low discrimination**						**07570022**
					GRA	CAT	
Propionibacterium granulosum		54.80	0.39	CEL 1	99	90	
Actinomyces naeslundii		37.14	0.51		99	25	
Prevotella oralis		3.39	0.08		1	1	
07570023	**Good identification**						**07570023**
Actinomyces naeslundii		100.0	0.54	GAL 96			
07570041	**Low discrimination**						**07570041**
					SPO		
Actinomyces naeslundii		89.99	0.41	GAL 96	1		
Clostridium botulinum biovar B		10.01	-.03		99		
07570042	**Low discrimination**						**07570042**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		61.61	0.38	GAL 96	1	25	
Propionibacterium granulosum		20.82	0.15		1	90	
Clostridium botulinum biovar B		10.29	-.03		99	1	
07570043	**Good identification**						**07570043**
Actinomyces naeslundii		100.0	0.41	GAL 96			
07570061	**Good identification**						**07570061**
Actinomyces naeslundii		100.0	0.41	GAL 96			
07570062	**Good identification**						**07570062**
Actinomyces naeslundii		100.0	0.38	GAL 96			
07570063	**Good identification**						**07570063**
Actinomyces naeslundii		100.0	0.41	GAL 96			
07570100	**Low discrimination**						**07570100**
					COC	GRA	
Fusobacterium mortiferum		73.48	0.56	MLT 11	1	1	
Actinomyces naeslundii		22.15	0.59		1	99	
Atopobium parvulum		1.95	0.18		99	99	
07570101	**Very good identification**						**07570101**
					GRA		
Actinomyces naeslundii		97.82	0.62	GAL 96	99		
Fusobacterium mortiferum		2.19	0.22		1		
07570102	**Low discrimination**						**07570102**
					GRA		
Actinomyces viscosus		44.97	0.37	NIT 99	99		
Actinomyces naeslundii		34.84	0.59	TRE 7	99		
Fusobacterium mortiferum		12.85	0.40		1		
07570103	**Low discrimination**						**07570103**
Actinomyces naeslundii		77.71	0.62	GAL 96			
Actinomyces viscosus		22.29	0.29				
07570110	**Low discrimination**						**07570110**
					GRA	CAT	
Actinomyces naeslundii		53.94	0.37	GAL 96	99	25	
Fusobacterium mortiferum		34.35	0.22	MLZ 5	1	5	
Propionibacterium granulosum		11.71	0.11		99	90	
07570111	**Good identification**						**07570111**
Actinomyces naeslundii		100.0	0.40	GAL 96			
07570112	**Low discrimination**						**07570112**
Actinomyces naeslundii		42.84	0.37	GAL 96			
Clostridium clostridioforme		28.91	0.16	MLZ 5			
Bifidobacterium adolescentis		10.75	0.11				
07570113	**Good identification**						**07570113**
Actinomyces naeslundii		100.0	0.40	GAL 96			
07570120	**Low discrimination**						**07570120**
					GRA		
Actinomyces naeslundii		60.54	0.59	GAL 96	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		24.16	0.30		99		
Prevotella melaninogenica		8.63	0.15		1		
07570121	**Very good identification**						**07570121**
Actinomyces naeslundii		100.0	0.62	GAL 96			
07570122	**Low discrimination**						**07570122**
					GRA		
Actinomyces naeslundii		49.14	0.59	GAL 96	99		
Prevotella oralis		24.69	0.28		1		
Clostridium perfringens		9.23	0.24		99		
07570123	**Very good identification**						**07570123**
Actinomyces naeslundii		100.0	0.62	GAL 96			
07570130	**Good identification**						**07570130**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07570131	**Good identification**						**07570131**
Actinomyces naeslundii		100.0	0.40	GAL 96			
07570133	**Good identification**						**07570133**
Actinomyces naeslundii		100.0	0.40	GAL 96			
07570140	**Low discrimination**						**07570140**
					GRA	SPO	
Actinomyces naeslundii		80.09	0.46	GAL 96	99	1	
Fusobacterium mortiferum		15.21	0.22		1	1	
Clostridium botulinum biovar B		4.70	-.03		99	99	
07570141	**Good identification**						**07570141**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07570142	**Low discrimination**						**07570142**
Actinomyces naeslundii		85.89	0.46	GAL 96			
Lactobacillus acidophilus		7.76	0.04				
Actinomyces viscosus		6.35	0.03				
07570143	**Good identification**						**07570143**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07570160	**Low discrimination**						**07570160**
Actinomyces naeslundii		69.90	0.46	GAL 96			
Clostridium perfringens		30.11	0.18				
07570161	**Good identification**						**07570161**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07570162	**Low discrimination**						**07570162**
Actinomyces naeslundii		77.34	0.46	GAL 96			
Clostridium perfringens		15.68	0.12				
Lactobacillus acidophilus		6.99	0.04				
07570163	**Good identification**						**07570163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	GAL 96		
07570322	**Low discrimination**					**07570322**
					SPO	
Parabacteroides distasonis		48.23	0.38	GAL 99	1	
Prevotella oralis		30.59	0.22	XYL 90	1	
Clostridium ramosum		10.24	0.14		92	
07570512	**Low discrimination**					**07570512**
Clostridium clostridioforme		86.29	0.36	SAL 88		
Bifidobacterium adolescentis		13.71	0.26	XYL 91		
07570532	**Acceptable identification**					**07570532**
					GRA	
Clostridium clostridioforme		90.50	0.36	SAL 88	50	
Parabacteroides distasonis		6.07	0.22	XYL 91	1	
Bifidobacterium longum subsp. longum		3.42	0.13	RHA 88	99	
07570612	**Good identification**					**07570612**
Clostridium clostridioforme		100.0	0.42	SAL 88		
07570632	**Good identification**					**07570632**
Clostridium clostridioforme		100.0	0.42	SAL 88		
07570702	**Good identification**					**07570702**
Clostridium clostridioforme		100.0	0.40	SAL 88		
07570712	**Good identification**					**07570712**
Clostridium clostridioforme		100.0	0.51	SAL 88		
07570722	**Low discrimination**					**07570722**
Clostridium clostridioforme		81.15	0.40	SAL 88		
Parabacteroides distasonis		18.85	0.34	XYL 91		
07570732	**Good identification**					**07570732**
Clostridium clostridioforme		97.33	0.51	SAL 88		
Parabacteroides distasonis		2.67	0.30	XYL 91		
07571000	**Low discrimination**					**07571000**
					CAT	
Actinomyces naeslundii		38.85	0.75		25	
Propionibacterium granulosum		38.22	0.60		90	
Lactobacillus jensenii		11.00	0.44		1	
07571001	**Very good identification**					**07571001**
					CAT	
Actinomyces naeslundii		98.99	0.78		25	
Propionibacterium granulosum		0.66	0.26		90	
Lactobacillus jensenii		0.19	0.10		1	
07571002	**Low discrimination**					**07571002**
					CAT	
Actinomyces naeslundii		58.82	0.75		25	
Propionibacterium granulosum		14.47	0.50		90	
Lactobacillus jensenii		11.11	0.41		1	
07571003	**Very good identification**					**07571003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.47	0.78			
Actinomyces viscosus		1.16	0.21			
Propionibacterium granulosum		0.16	0.15			
07571010	**Low discrimination**					**07571010**
					CAT	
Propionibacterium granulosum		46.98	0.45	CEL 1	90	
Lactobacillus jensenii		33.06	0.36	MLZ 12	1	
Actinomyces naeslundii		18.43	0.53		25	
07571011	**Good identification**					**07571011**
					CAT	
Actinomyces naeslundii		97.16	0.56	MLZ 5	25	
Propionibacterium granulosum		1.67	0.11		90	
Lactobacillus jensenii		1.17	0.02		1	
07571013	**Good identification**					**07571013**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
07571020	**Low discrimination**					**07571020**
					CAT	
Actinomyces naeslundii		50.01	0.75		25	
Clostridium perfringens		31.95	0.49		1	
Propionibacterium granulosum		12.30	0.50		90	
07571021	**Very good identification**					**07571021**
					CAT	
Actinomyces naeslundii		99.41	0.78		25	
Clostridium perfringens		0.43	0.15		1	
Propionibacterium granulosum		0.17	0.15		90	
07571022	**Low discrimination**					**07571022**
					GRA	
Actinomyces naeslundii		51.50	0.75		99	
Prevotella oralis		19.39	0.42		1	
Clostridium perfringens		15.48	0.44		99	
07571023	**Very good identification**					**07571023**
					GRA	
Actinomyces naeslundii		99.46	0.78		99	
Prevotella oralis		0.25	0.08		1	
Clostridium perfringens		0.20	0.10		99	
07571030	**Low discrimination**					**07571030**
					CAT	
Actinomyces naeslundii		45.28	0.53	MLZ 5	25	
Propionibacterium granulosum		28.86	0.35		90	
Lactobacillus jensenii		20.31	0.26		1	
07571031	**Good identification**					**07571031**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
07571032	**Low discrimination**					**07571032**
					CAT	
Actinomyces naeslundii		54.87	0.53	MLZ 5	25	
Lactobacillus jensenii		16.40	0.23		1	
Propionibacterium granulosum		8.74	0.25		90	
07571033	**Good identification**					**07571033**
Actinomyces naeslundii		100.0	0.56	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07571040	**Low discrimination**					**07571040**
					CAT	
Actinomyces naeslundii		87.56	0.62		25	
Propionibacterium granulosum		4.93	0.26		90	
Actinomyces israelii		2.34	0.26		1	
07571041	**Very good identification**					**07571041**
Actinomyces naeslundii		100.0	0.65			
07571042	**Low discrimination**					**07571042**
Actinomyces naeslundii		85.67	0.62			
Lactobacillus acidophilus		7.90	0.20			
Actinomyces israelii		3.44	0.29			
07571043	**Very good identification**					**07571043**
Actinomyces naeslundii		100.0	0.65			
07571050	**Low discrimination**					**07571050**
					CAT	
Actinomyces naeslundii		80.09	0.40	MLZ 5	25	
Propionibacterium granulosum		11.69	0.11		90	
Lactobacillus jensenii		8.22	0.02		1	
07571051	**Good identification**					**07571051**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07571052	**Low discrimination**					**07571052**
Actinomyces naeslundii		89.77	0.40	MLZ 5		
Actinomyces israelii		10.23	0.15			
07571053	**Good identification**					**07571053**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07571060	**Low discrimination**					**07571060**
					CAT	
Actinomyces naeslundii		58.70	0.62		25	
Clostridium perfringens		40.48	0.37		1	
Propionibacterium granulosum		0.83	0.15		90	
07571061	**Very good identification**					**07571061**
Actinomyces naeslundii		99.54	0.65			
Clostridium perfringens		0.46	0.03			
07571062	**Low discrimination**					**07571062**
Actinomyces naeslundii		69.53	0.62			
Clostridium perfringens		22.56	0.31			
Lactobacillus acidophilus		6.41	0.20			
07571063	**Very good identification**					**07571063**
Actinomyces naeslundii		100.0	0.65			
07571070	**Low discrimination**					**07571070**
Actinomyces naeslundii		88.31	0.40	MLZ 5		
Clostridium perfringens		11.69	0.03			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07571071	**Good identification**						**07571071**
Actinomyces naeslundii		100.0	0.43	MLZ 5			
07571072	**Good identification**						**07571072**
Actinomyces naeslundii		100.0	0.40	MLZ 5			
07571073	**Good identification**						**07571073**
Actinomyces naeslundii		100.0	0.43	MLZ 5			
07571100	**Low discrimination**						**07571100**
					COC	GRA	
Actinomyces naeslundii		60.01	0.83		1	99	
Atopobium parvulum		21.84	0.52		99	99	
Fusobacterium mortiferum		12.44	0.59		1	1	
07571101	**Very good identification**						**07571101**
					COC	GRA	
Actinomyces naeslundii		99.51	0.86		1	99	
Atopobium parvulum		0.24	0.18		99	99	
Fusobacterium mortiferum		0.14	0.25		1	1	
07571102	**Low discrimination**						**07571102**
					GRA		
Actinomyces naeslundii		53.40	0.83		99		
Lactobacillus acidophilus		28.42	0.54		99		
Prevotella oralis		5.83	0.41		1		
07571103	**Very good identification**						**07571103**
Actinomyces naeslundii		98.78	0.86				
Actinomyces viscosus		0.72	0.25				
Lactobacillus acidophilus		0.35	0.19				
07571110	**Low discrimination**						**07571110**
					COC		
Actinomyces naeslundii		83.30	0.61	MLZ 5	1		
Atopobium parvulum		5.82	0.18		99		
Actinomyces israelii		4.04	0.29		1		
07571111	**Very good identification**						**07571111**
Actinomyces naeslundii		100.0	0.64	MLZ 5			
07571112	**Low discrimination**						**07571112**
					GRA		
Bifidobacterium adolescentis		33.40	0.46	XYL 87	99		
Actinomyces naeslundii		32.26	0.61	ARA 87	99		
Leptotrichia buccalis		21.43	0.24		1		
07571113	**Very good identification**						**07571113**
					GRA		
Actinomyces naeslundii		98.87	0.64	MLZ 5	99		
Bifidobacterium adolescentis		0.69	0.11		99		
Leptotrichia buccalis		0.44	-.11		1		
07571120	**Low discrimination**						**07571120**
					GRA		
Clostridium perfringens		47.20	0.64	CEL 4	99		
Actinomyces naeslundii		28.68	0.83		99		
Prevotella melaninogenica		16.87	0.50		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07571121	**Very good identification**					**07571121**
					GRA	
Actinomyces naeslundii		98.34	0.86		99	
Clostridium perfringens		1.09	0.30		99	
Prevotella melaninogenica		0.39	0.15		1	
07571122	**Low discrimination**					**07571122**
					GRA	
Prevotella oralis		39.78	0.62	TRE 6	1	
Actinomyces naeslundii		19.19	0.83		99	
Clostridium perfringens		14.87	0.58		99	
07571123	**Very good identification**					**07571123**
					GRA	
Actinomyces naeslundii		97.26	0.86		99	
Prevotella oralis		1.36	0.28		1	
Clostridium perfringens		0.51	0.24		99	
07571130	**Low discrimination**					**07571130**
					GRA	
Actinomyces naeslundii		64.64	0.61	MLZ 5	99	
Clostridium perfringens		20.42	0.30		99	
Prevotella melaninogenica		7.30	0.15		1	
07571131	**Very good identification**					**07571131**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
07571132	**Low discrimination**					**07571132**
					GRA	
Parabacteroides distasonis		70.44	0.59	XYL 90	1	
Actinomyces naeslundii		14.75	0.61		99	
Prevotella oralis		5.87	0.28		1	
07571133	**Very good identification**					**07571133**
					GRA	
Actinomyces naeslundii		96.88	0.64	MLZ 5	99	
Parabacteroides distasonis		3.12	0.25		1	
07571140	**Low discrimination**					**07571140**
Actinomyces naeslundii		88.57	0.70			
Clostridium perfringens		4.87	0.26			
Actinomyces israelii		3.33	0.36			
07571141	**Very good identification**					**07571141**
Actinomyces naeslundii		100.0	0.73			
07571142	**Low discrimination**					**07571142**
Actinomyces naeslundii		66.06	0.70			
Lactobacillus acidophilus		24.62	0.38			
Actinomyces israelii		3.73	0.39			
07571143	**Very good identification**					**07571143**
Actinomyces naeslundii		99.75	0.73			
Lactobacillus acidophilus		0.25	0.04			
07571150	**Acceptable identification**					**07571150**
Actinomyces naeslundii		90.35	0.48	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		9.65	0.22			
07571151	**Good identification**					**07571151**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
07571152	**Low discrimination**					**07571152**
Actinomyces naeslundii		43.56	0.48	MLZ 5		
Bifidobacterium adolescentis		31.58	0.30			
Bifidobacterium breve		13.11	0.24			
07571153	**Good identification**					**07571153**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
07571160	**Low discrimination**					**07571160**
					GRA	
Clostridium perfringens		62.73	0.52	CEL 4	99	
Actinomyces naeslundii		35.31	0.70		99	
Prevotella melaninogenica		1.19	0.15		1	
07571161	**Very good identification**					**07571161**
Actinomyces naeslundii		98.82	0.73			
Clostridium perfringens		1.18	0.18			
07571162	**Low discrimination**					**07571162**
Actinomyces naeslundii		41.52	0.70			
Clostridium perfringens		34.71	0.46			
Lactobacillus acidophilus		15.48	0.38			
07571163	**Very good identification**					**07571163**
Actinomyces naeslundii		99.19	0.73			
Clostridium perfringens		0.56	0.12			
Lactobacillus acidophilus		0.25	0.04			
07571170	**Low discrimination**					**07571170**
Actinomyces naeslundii		71.74	0.48	MLZ 5		
Clostridium perfringens		24.46	0.18			
Bifidobacterium breve		3.80	0.12			
07571171	**Good identification**					**07571171**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
07571172	**Low discrimination**					**07571172**
					GRA	
Actinomyces naeslundii		49.01	0.48	MLZ 5	99	
Bifidobacterium breve		26.23	0.29		99	
Parabacteroides distasonis		13.40	0.25		1	
07571173	**Good identification**					**07571173**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
07571201	**Good identification**					**07571201**
Actinomyces naeslundii		100.0	0.44	RHA 1		
07571202	**Low discrimination**					**07571202**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		22.21	0.41	RHA 1	99		
Lactobacillus acidophilus		21.80	0.16		99		
Prevotella oralis		18.68	0.14		1		
07571203	**Good identification**						**07571203**
Actinomyces naeslundii		100.0	0.44	RHA 1			
07571220	**Low discrimination**						**07571220**
					GRA	CAT	
Actinomyces naeslundii		29.03	0.41	RHA 1	99	25	
Prevotella oralis		24.41	0.14		1	1	
Propionibacterium granulosum		21.86	0.24		99	90	
07571221	**Good identification**						**07571221**
Actinomyces naeslundii		100.0	0.44	RHA 1			
07571222	**Low discrimination**						**07571222**
					GRA		
Prevotella oralis		66.90	0.36	TRE 6	1		
Parabacteroides distasonis		16.69	0.38	LAC 94	1		
Actinomyces naeslundii		4.19	0.41		99		
07571223	**Acceptable identification**						**07571223**
					GRA		
Actinomyces naeslundii		90.29	0.44	RHA 1	99		
Prevotella oralis		9.71	0.02		1		
07571300	**Low discrimination**						**07571300**
					COC	GRA	
Actinomyces naeslundii		47.45	0.49	RHA 1	1	99	
Atopobium parvulum		17.27	0.18		99	99	
Prevotella oralis		11.56	0.12		1	1	
07571301	**Good identification**						**07571301**
Actinomyces naeslundii		100.0	0.52	RHA 1			
07571303	**Acceptable identification**						**07571303**
					GRA		
Actinomyces naeslundii		94.53	0.52	RHA 1	99		
Prevotella oralis		2.95	0.00		1		
Lactobacillus acidophilus		2.52	0.00		99		
07571312	**Low discrimination**						**07571312**
					GRA		
Clostridium clostridioforme		76.63	0.39	SAL 88	50		
Parabacteroides distasonis		17.93	0.33	XYL 91	1		
Bifidobacterium adolescentis		1.30	0.11	ARA 94	99		
07571320	**Low discrimination**						**07571320**
					GRA	SPO	
Clostridium ramosum		43.73	0.38	SAL 99	50	92	
Prevotella oralis		27.51	0.34		1	1	
Clostridium perfringens		9.78	0.30		99	84	
07571321	**Acceptable identification**						**07571321**
					SPO	GRA	
Actinomyces naeslundii		92.53	0.52	RHA 1	1	99	
Clostridium ramosum		4.58	0.04		92	50	
Prevotella oralis		2.88	0.00		1	1	
07571322	**Low discrimination**						**07571322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Parabacteroides distasonis		52.80	0.72	XYL 90	1	
Prevotella oralis		33.49	0.56		1	
Clostridium ramosum		11.21	0.48		92	
07571323	**Low discrimination**					**07571323**
					GRA	
Actinomyces naeslundii		36.35	0.52	RHA 1	99	
Parabacteroides distasonis		33.94	0.38		1	
Prevotella oralis		21.53	0.22		1	
07571332	**Good identification**					**07571332**
Parabacteroides distasonis		94.31	0.68	XYL 90		
Clostridium clostridioforme		4.07	0.39			
Prevotella oralis		1.12	0.22			
07571340	**Low discrimination**					**07571340**
Actinomyces naeslundii		78.11	0.36	RHA 1		
Actinomyces israelii		21.89	0.17			
07571341	**Good identification**					**07571341**
Actinomyces naeslundii		100.0	0.39	RHA 1		
07571343	**Good identification**					**07571343**
Actinomyces naeslundii		100.0	0.39	RHA 1		
07571361	**Good identification**					**07571361**
Actinomyces naeslundii		100.0	0.39	RHA 1		
07571362	**Low discrimination**					**07571362**
					GRA	
Parabacteroides distasonis		41.19	0.38	XYL 90	1	
Prevotella oralis		26.13	0.22	SOR 1	1	
Lactobacillus acidophilus		14.42	0.19		99	
07571363	**Good identification**					**07571363**
Actinomyces naeslundii		100.0	0.39	RHA 1		
07571400	**Low discrimination**					**07571400**
					CAT	
Actinomyces naeslundii		24.75	0.41	ARA 1	25	
Propionibacterium granulosum		24.35	0.26		90	
Actinomyces israelii		24.34	0.31		1	
07571401	**Good identification**					**07571401**
Actinomyces naeslundii		100.0	0.44	ARA 1		
07571403	**Good identification**					**07571403**
Actinomyces naeslundii		100.0	0.44	ARA 1		
07571412	**Low discrimination**					**07571412**
Clostridium clostridioforme		82.31	0.35	SAL 88		
Bifidobacterium adolescentis		14.59	0.26	XYL 91		
Actinomyces israelii		3.10	0.20	RHA 88		
07571420	**Low discrimination**					**07571420**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Actinomyces naeslundii		49.26	0.41	ARA 1	25	
Clostridium perfringens		31.47	0.15		1	
Propionibacterium granulosum		12.12	0.15		90	
07571421	**Good identification**					**07571421**
Actinomyces naeslundii		100.0	0.44	ARA 1		
07571423	**Good identification**					**07571423**
Actinomyces naeslundii		100.0	0.44	ARA 1		
07571432	**Low discrimination**					**07571432**
					GRA	
Clostridium clostridioforme		89.52	0.35	SAL 88	50	
Parabacteroides distasonis		6.71	0.22	XYL 91	1	
Bifidobacterium longum subsp. longum		3.78	0.13	RHA 88	99	
07571501	**Good identification**					**07571501**
Actinomyces naeslundii		100.0	0.52	ARA 1		
07571502	**Low discrimination**					**07571502**
Bifidobacterium adolescentis		78.69	0.58	XYL 87		
Actinomyces israelii		5.52	0.45			
Clostridium clostridioforme		4.03	0.33			
07571503	**Low discrimination**					**07571503**
Actinomyces naeslundii		83.41	0.52	ARA 1		
Bifidobacterium adolescentis		16.59	0.24			
07571510	**Low discrimination**					**07571510**
Bifidobacterium longum subsp. longum		72.21	0.37	CEL 1		
Bifidobacterium adolescentis		12.25	0.26			
Actinomyces israelii		6.94	0.28			
07571512	**Low discrimination**					**07571512**
Bifidobacterium adolescentis		78.92	0.60	XYL 87		
Clostridium clostridioforme		15.06	0.44			
Bifidobacterium longum subsp. longum		4.70	0.37			
07571520	**Low discrimination**					**07571520**
Bifidobacterium longum subsp. longum		54.29	0.43	CEL 1		
Clostridium perfringens		17.70	0.30			
Actinomyces naeslundii		10.75	0.49			
07571521	**Good identification**					**07571521**
Actinomyces naeslundii		96.71	0.52	ARA 1		
Bifidobacterium longum subsp. longum		3.29	0.09			
07571522	**Low discrimination**					**07571522**
					GRA	
Parabacteroides distasonis		70.12	0.60	XYL 90	1	
Bifidobacterium longum subsp. longum		12.50	0.43		99	
Prevotella oralis		5.13	0.28		1	
07571523	**Low discrimination**					**07571523**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
				GRA	
Actinomyces naeslundii	81.65	0.52	ARA 1	99	
Parabacteroides distasonis	15.58	0.26		1	
Bifidobacterium longum subsp. longum	2.78	0.09		99	
07571530 Good identification					**07571530**
				GRA	
Bifidobacterium longum subsp. longum	96.89	0.47	CEL 1	99	
Parabacteroides distasonis	1.74	0.22		1	
Clostridium clostridioforme	0.78	0.10		50	
07571532 Low discrimination					**07571532**
				GRA	
Parabacteroides distasonis	48.87	0.56	XYL 90	1	
Bifidobacterium longum subsp. longum	27.55	0.47		99	
Clostridium clostridioforme	22.07	0.44		50	
07571541 Good identification					**07571541**
Actinomyces naeslundii	100.0	0.39	ARA 1		
07571542 Low discrimination					**07571542**
Bifidobacterium adolescentis	76.29	0.43	XYL 87		
Actinomyces israelii	16.85	0.38	SOR 11		
Actinomyces naeslundii	3.69	0.36			
07571543 Low discrimination					**07571543**
Actinomyces naeslundii	87.77	0.39	ARA 1		
Bifidobacterium adolescentis	12.23	0.09			
07571552 Good identification					**07571552**
Bifidobacterium adolescentis	95.92	0.44	XYL 87		
Actinomyces israelii	2.59	0.24	SOR 11		
Clostridium clostridioforme	1.50	0.10			
07571561 Good identification					**07571561**
Actinomyces naeslundii	100.0	0.39	ARA 1		
07571563 Good identification					**07571563**
Actinomyces naeslundii	100.0	0.39	ARA 1		
07571602 Good identification					**07571602**
Clostridium clostridioforme	98.84	0.39	SAL 88		
Actinomyces israelii	1.16	0.15	XYL 91		
07571612 Good identification					**07571612**
Clostridium clostridioforme	100.0	0.50	SAL 88		
07571622 Low discrimination					**07571622**
Clostridium clostridioforme	77.66	0.39	SAL 88		
Parabacteroides distasonis	20.13	0.34	XYL 91		
Prevotella oralis	1.33	0.02			
07571632 Good identification					**07571632**
Clostridium clostridioforme	97.03	0.50	SAL 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		2.97	0.30	XYL 91			
07571702	**Low discrimination**						**07571702**
					GRA		
Clostridium clostridioforme		88.57	0.48	SAL 88	50		
Parabacteroides distasonis		6.86	0.34	XYL 91	1		
Bifidobacterium adolescentis		2.38	0.24		99		
07571712	**Very good identification**						**07571712**
					GRA		
Clostridium clostridioforme		98.39	0.59	SAL 88	50		
Parabacteroides distasonis		0.90	0.30	XYL 91	1		
Bifidobacterium adolescentis		0.71	0.26		99		
07571722	**Low discrimination**						**07571722**
Parabacteroides distasonis		85.49	0.69	XYL 90			
Clostridium clostridioforme		11.15	0.48				
Bacteroides thetaiotaomicron		1.86	0.24				
07571732	**Low discrimination**						**07571732**
Clostridium clostridioforme		51.95	0.59	SAL 88			
Parabacteroides distasonis		47.07	0.64	XYL 91			
Bacteroides thetaiotaomicron		0.74	0.17				
07572000	**Low discrimination**						**07572000**
					GRA	CAT	
Propionibacterium granulosum		89.99	0.58	CEL 1	99	90	
Actinomyces naeslundii		9.81	0.56		99	25	
Fusobacterium mortiferum		0.21	0.16		1	5	
07572001	**Good identification**						**07572001**
Actinomyces naeslundii		94.18	0.59	GAL 96			
Propionibacterium granulosum		5.82	0.24				
07572002	**Low discrimination**						**07572002**
					CAT		
Propionibacterium granulosum		69.15	0.48	CEL 1	90		
Actinomyces naeslundii		30.14	0.56		25		
Actinomyces israelii		0.39	0.15		1		
07572003	**Very good identification**						**07572003**
Actinomyces naeslundii		98.48	0.59	GAL 96			
Propionibacterium granulosum		1.52	0.13				
07572010	**Good identification**						**07572010**
Propionibacterium granulosum		95.96	0.43	CEL 1			
Actinomyces naeslundii		4.04	0.34	MLZ 12			
07572011	**Low discrimination**						**07572011**
Actinomyces naeslundii		86.20	0.37	GAL 96			
Propionibacterium granulosum		13.80	0.09	MLZ 5			
07572013	**Good identification**						**07572013**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07572020	**Low discrimination**						**07572020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		69.64	0.48	CEL 1			
Actinomyces naeslundii		30.36	0.56				
07572021	**Very good identification**						**07572021**
Actinomyces naeslundii		98.48	0.59	GAL 96			
Propionibacterium granulosum		1.52	0.13				
07572022	**Low discrimination**						**07572022**
Actinomyces naeslundii		63.55	0.56	GAL 96			
Propionibacterium granulosum		36.45	0.37				
07572023	**Very good identification**						**07572023**
Actinomyces naeslundii		100.0	0.59	GAL 96			
07572031	**Good identification**						**07572031**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07572033	**Good identification**						**07572033**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07572040	**Low discrimination**						**07572040**
					SPO	CAT	
Actinomyces naeslundii		62.07	0.43	GAL 96	1	25	
Propionibacterium granulosum		32.60	0.24		1	90	
Clostridium botulinum biovar B		5.34	-.03		99	1	
07572041	**Good identification**						**07572041**
Actinomyces naeslundii		100.0	0.46	GAL 96			
07572042	**Low discrimination**						**07572042**
Actinomyces naeslundii		88.40	0.43	GAL 96			
Propionibacterium granulosum		11.61	0.13				
07572043	**Good identification**						**07572043**
Actinomyces naeslundii		100.0	0.46	GAL 96			
07572060	**Low discrimination**						**07572060**
Actinomyces naeslundii		88.40	0.43	GAL 96			
Propionibacterium granulosum		11.61	0.13				
07572061	**Good identification**						**07572061**
Actinomyces naeslundii		100.0	0.46	GAL 96			
07572062	**Good identification**						**07572062**
Actinomyces naeslundii		100.0	0.43	GAL 96			
07572063	**Good identification**						**07572063**
Actinomyces naeslundii		100.0	0.46	GAL 96			
07572100	**Low discrimination**						**07572100**
					CAT		
Actinomyces naeslundii		79.15	0.64	GAL 96	25		
Propionibacterium propionicum		15.95	0.42		1		
Propionibacterium granulosum		2.58	0.24		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07572101	**Very good identification**						**07572101**
Actinomyces naeslundii		98.53	0.67	GAL 96			
Propionibacterium propionicum		1.47	0.26				
07572102	**Low discrimination**						**07572102**
Actinomyces naeslundii		74.64	0.64	GAL 96			
Propionibacterium propionicum		22.56	0.45				
Actinomyces israelii		1.35	0.25				
07572103	**Very good identification**						**07572103**
Actinomyces naeslundii		97.81	0.67	GAL 96			
Propionibacterium propionicum		2.19	0.29				
07572110	**Acceptable identification**						**07572110**
Actinomyces naeslundii		92.22	0.42	GAL 96			
Propionibacterium granulosum		7.78	0.09	MLZ 5			
07572111	**Good identification**						**07572111**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07572112	**Acceptable identification**						**07572112**
Actinomyces naeslundii		90.43	0.42	GAL 96			
Bifidobacterium adolescentis		9.57	0.10	MLZ 5			
07572113	**Good identification**						**07572113**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07572120	**Very good identification**						**07572120**
					SPO	CAT	
Actinomyces naeslundii		95.44	0.64	GAL 96	1	25	
Clostridium ramosum		3.79	0.15		92	1	
Propionibacterium granulosum		0.78	0.13		1	90	
07572121	**Very good identification**						**07572121**
Actinomyces naeslundii		100.0	0.67	GAL 96			
07572122	**Low discrimination**						**07572122**
					SPO		
Actinomyces naeslundii		86.30	0.64	GAL 96	1		
Clostridium ramosum		13.70	0.25		92		
07572123	**Very good identification**						**07572123**
Actinomyces naeslundii		100.0	0.67	GAL 96			
07572130	**Good identification**						**07572130**
Actinomyces naeslundii		100.0	0.42	GAL 96			
07572131	**Good identification**						**07572131**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07572132	**Good identification**						**07572132**
Actinomyces naeslundii		100.0	0.42	GAL 96			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07572133	**Good identification**					**07572133**
Actinomyces naeslundii		100.0	0.45	GAL 96		
07572140	**Low discrimination**					**07572140**
Propionibacterium propionicum		52.66	0.42	NIT 86		
Actinomyces naeslundii		46.11	0.51	CEL 1		
Actinomyces israelii		1.23	0.15			
07572141	**Acceptable identification**					**07572141**
Actinomyces naeslundii		92.20	0.54	GAL 96		
Propionibacterium propionicum		7.80	0.26			
07572142	**Low discrimination**					**07572142**
Propionibacterium propionicum		62.23	0.45	NIT 86		
Actinomyces naeslundii		36.32	0.51	CEL 1		
Actinomyces israelii		1.45	0.18			
07572143	**Low discrimination**					**07572143**
Actinomyces naeslundii		88.74	0.54	GAL 96		
Propionibacterium propionicum		11.26	0.29			
07572160	**Good identification**					**07572160**
Actinomyces naeslundii		100.0	0.51	GAL 96		
07572161	**Good identification**					**07572161**
Actinomyces naeslundii		100.0	0.54	GAL 96		
07572162	**Good identification**					**07572162**
Actinomyces naeslundii		100.0	0.51	GAL 96		
07572163	**Good identification**					**07572163**
Actinomyces naeslundii		100.0	0.54	GAL 96		
07573000	**Low discrimination**					**07573000**
					CAT	
Actinomyces naeslundii		70.40	0.80		25	
Propionibacterium granulosum		26.91	0.58		90	
Actinomyces israelii		2.50	0.46		1	
07573001	**Very good identification**					**07573001**
Actinomyces naeslundii		99.74	0.83			
Propionibacterium granulosum		0.26	0.24			
07573002	**Low discrimination**					**07573002**
					CAT	
Actinomyces naeslundii		86.67	0.80		25	
Propionibacterium granulosum		8.28	0.48		90	
Actinomyces israelii		4.61	0.49		1	
07573003	**Very good identification**					**07573003**
					CAT	
Actinomyces naeslundii		99.90	0.83		25	
Propionibacterium granulosum		0.06	0.13		90	
Actinomyces israelii		0.04	0.15		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07573010	**Low discrimination**						**07573010**
					CAT		
Actinomyces naeslundii		47.61	0.58	MLZ 5	25		
Propionibacterium granulosum		47.16	0.43		90		
Actinomyces israelii		4.79	0.32		1		
07573011	**Very good identification**						**07573011**
Actinomyces naeslundii		99.34	0.61	MLZ 5			
Propionibacterium granulosum		0.66	0.09				
07573012	**Low discrimination**						**07573012**
					CAT		
Actinomyces naeslundii		70.87	0.58	MLZ 5	25		
Propionibacterium granulosum		17.55	0.33		90		
Actinomyces israelii		10.70	0.35		1		
07573013	**Very good identification**						**07573013**
Actinomyces naeslundii		100.0	0.61	MLZ 5			
07573020	**Good identification**						**07573020**
					SPO	CAT	
Actinomyces naeslundii		90.43	0.80		1	25	
Propionibacterium granulosum		8.64	0.48		1	90	
Clostridium ramosum		0.43	0.15		92	1	
07573021	**Very good identification**						**07573021**
Actinomyces naeslundii		99.94	0.83				
Propionibacterium granulosum		0.06	0.13				
07573022	**Good identification**						**07573022**
					SPO	CAT	
Actinomyces naeslundii		94.95	0.80		1	25	
Propionibacterium granulosum		2.27	0.37		1	90	
Clostridium ramosum		1.79	0.25		92	1	
07573023	**Very good identification**						**07573023**
Actinomyces naeslundii		100.0	0.83				
07573030	**Low discrimination**						**07573030**
Actinomyces naeslundii		80.15	0.58	MLZ 5			
Propionibacterium granulosum		19.85	0.33				
07573031	**Very good identification**						**07573031**
Actinomyces naeslundii		100.0	0.61	MLZ 5			
07573032	**Good identification**						**07573032**
					CAT		
Actinomyces naeslundii		93.32	0.58	MLZ 5	25		
Propionibacterium granulosum		5.78	0.22		90		
Actinomyces israelii		0.90	0.14		1		
07573033	**Very good identification**						**07573033**
Actinomyces naeslundii		100.0	0.61	MLZ 5			
07573040	**Good identification**						**07573040**
					CAT		
Actinomyces naeslundii		90.91	0.67		25		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		7.10	0.39		1	
Propionibacterium granulosum		1.99	0.24		90	
07573041	**Very good identification**					**07573041**
Actinomyces naeslundii		100.0	0.70			
07573042	**Low discrimination**					**07573042**
					CAT	
Actinomyces naeslundii		88.88	0.67		25	
Actinomyces israelii		10.41	0.42		1	
Propionibacterium granulosum		0.49	0.13		90	
07573043	**Very good identification**					**07573043**
Actinomyces naeslundii		100.0	0.70			
07573050	**Low discrimination**					**07573050**
					CAT	
Actinomyces naeslundii		78.22	0.45	MLZ 5	25	
Actinomyces israelii		17.35	0.25		1	
Propionibacterium granulosum		4.43	0.09		90	
07573051	**Good identification**					**07573051**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07573052	**Low discrimination**					**07573052**
Actinomyces naeslundii		75.04	0.45	MLZ 5		
Actinomyces israelii		24.96	0.28			
07573053	**Good identification**					**07573053**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07573060	**Very good identification**					**07573060**
					CAT	
Actinomyces naeslundii		98.62	0.67		25	
Propionibacterium granulosum		0.54	0.13		90	
Actinomyces israelii		0.49	0.18		1	
07573061	**Very good identification**					**07573061**
Actinomyces naeslundii		100.0	0.70			
07573062	**Very good identification**					**07573062**
Actinomyces naeslundii		99.01	0.67			
Actinomyces israelii		0.74	0.21			
Lactobacillus acidophilus		0.25	-.02			
07573063	**Very good identification**					**07573063**
Actinomyces naeslundii		100.0	0.70			
07573070	**Good identification**					**07573070**
Actinomyces naeslundii		100.0	0.45	MLZ 5		
07573071	**Good identification**					**07573071**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07573072	**Good identification**					**07573072**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	MLZ 5		
07573073	**Good identification**					**07573073**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07573100	**Good identification**					**07573100**
Actinomyces naeslundii		93.90	0.88			
Actinomyces israelii		4.68	0.56			
Propionibacterium propionicum		0.79	0.42			
07573101	**Very good identification**					**07573101**
Actinomyces naeslundii		99.90	0.91			
Propionibacterium propionicum		0.06	0.26			
Actinomyces israelii		0.03	0.22			
07573102	**Low discrimination**					**07573102**
Actinomyces naeslundii		88.45	0.88			
Actinomyces israelii		6.61	0.59			
Bifidobacterium adolescentis		1.66	0.43			
07573103	**Very good identification**					**07573103**
Actinomyces naeslundii		99.84	0.91			
Propionibacterium propionicum		0.09	0.29			
Actinomyces israelii		0.05	0.25			
07573110	**Low discrimination**					**07573110**
					SPO	
Actinomyces naeslundii		86.51	0.66	MLZ 5	1	
Actinomyces israelii		12.23	0.42		1	
Clostridium tertium		0.57	0.05		99	
07573111	**Very good identification**					**07573111**
Actinomyces naeslundii		100.0	0.69	MLZ 5		
07573112	**Low discrimination**					**07573112**
Actinomyces naeslundii		58.86	0.66	MLZ 5		
Bifidobacterium adolescentis		25.68	0.44			
Actinomyces israelii		12.49	0.45			
07573113	**Very good identification**					**07573113**
Actinomyces naeslundii		99.71	0.69	MLZ 5		
Bifidobacterium adolescentis		0.29	0.10			
07573120	**Low discrimination**					**07573120**
					SPO	
Actinomyces naeslundii		84.71	0.88		1	
Clostridium ramosum		13.86	0.49		92	
Clostridium perfringens		0.73	0.30		84	
07573121	**Very good identification**					**07573121**
					SPO	
Actinomyces naeslundii		99.89	0.91		1	
Clostridium ramosum		0.11	0.15		92	
07573122	**Low discrimination**					**07573122**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		58.96	0.88		1	
Clostridium ramosum		38.59	0.59		92	
Lactobacillus acidophilus		0.85	0.32		1	
07573123	**Very good identification**					**07573123**
					SPO	
Actinomyces naeslundii		99.56	0.91		1	
Clostridium ramosum		0.44	0.25		92	
07573130	**Good identification**					**07573130**
					SPO	
Actinomyces naeslundii		93.90	0.66	MLZ 5	1	
Clostridium ramosum		2.95	0.15		92	
Clostridium tertium		1.85	0.13		99	
07573131	**Very good identification**					**07573131**
Actinomyces naeslundii		100.0	0.69	MLZ 5		
07573132	**Low discrimination**					**07573132**
					SPO	
Actinomyces naeslundii		78.35	0.66	MLZ 5	1	
Clostridium ramosum		9.84	0.25		92	
Clostridium tertium		4.64	0.21		99	
07573133	**Very good identification**					**07573133**
Actinomyces naeslundii		100.0	0.69	MLZ 5		
07573140	**Low discrimination**					**07573140**
Actinomyces naeslundii		86.40	0.75			
Actinomyces israelii		9.49	0.49			
Propionibacterium propionicum		4.11	0.42			
07573141	**Very good identification**					**07573141**
Actinomyces naeslundii		99.58	0.78			
Propionibacterium propionicum		0.35	0.26			
Actinomyces israelii		0.07	0.15			
07573142	**Low discrimination**					**07573142**
Actinomyces naeslundii		78.90	0.75			
Actinomyces israelii		12.99	0.52			
Propionibacterium propionicum		5.63	0.45			
07573143	**Very good identification**					**07573143**
Actinomyces naeslundii		99.36	0.78			
Propionibacterium propionicum		0.53	0.29			
Actinomyces israelii		0.11	0.18			
07573150	**Low discrimination**					**07573150**
Actinomyces naeslundii		76.24	0.53	MLZ 5		
Actinomyces israelii		23.76	0.35			
07573151	**Good identification**					**07573151**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
07573152	**Low discrimination**					**07573152**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		51.86	0.53	MLZ 5			
Actinomyces israelii		24.25	0.38				
Bifidobacterium adolescentis		15.85	0.29				
07573153	**Good identification**						**07573153**
Actinomyces naeslundii		100.0	0.56	MLZ 5			
07573160	**Very good identification**						**07573160**
					SPO		
Actinomyces naeslundii		97.37	0.75		1		
Clostridium ramosum		0.91	0.15		92		
Clostridium perfringens		0.90	0.18		84		
07573161	**Very good identification**						**07573161**
Actinomyces naeslundii		100.0	0.78				
07573162	**Good identification**						**07573162**
					SPO		
Actinomyces naeslundii		92.86	0.75		1		
Clostridium ramosum		3.48	0.25		92		
Bifidobacterium breve		1.35	0.29		1		
07573163	**Very good identification**						**07573163**
Actinomyces naeslundii		100.0	0.78				
07573170	**Good identification**						**07573170**
Actinomyces naeslundii		95.50	0.53	MLZ 5			
Bifidobacterium breve		2.60	0.12				
Actinomyces israelii		1.90	0.15				
07573171	**Good identification**						**07573171**
Actinomyces naeslundii		100.0	0.56	MLZ 5			
07573172	**Low discrimination**						**07573172**
Actinomyces naeslundii		76.60	0.53	MLZ 5			
Bifidobacterium breve		21.12	0.29				
Actinomyces israelii		2.29	0.18				
07573173	**Good identification**						**07573173**
Actinomyces naeslundii		100.0	0.56	MLZ 5			
07573201	**Good identification**						**07573201**
Actinomyces naeslundii		100.0	0.49	RHA 1			
07573202	**Low discrimination**						**07573202**
					CAT		
Actinomyces naeslundii		59.21	0.46	RHA 1	25		
Actinomyces israelii		23.46	0.30		1		
Propionibacterium granulosum		17.33	0.22		90		
07573203	**Good identification**						**07573203**
Actinomyces naeslundii		100.0	0.49	RHA 1			
07573220	**Low discrimination**						**07573220**
					SPO	CAT	
Actinomyces naeslundii		52.35	0.46	RHA 1	1	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		32.33	0.17		92	1	
Propionibacterium granulosum		15.32	0.22		1	90	
07573221	**Good identification**						**07573221**
Actinomyces naeslundii		100.0	0.49	RHA 1			
07573223	**Good identification**						**07573223**
Actinomyces naeslundii		100.0	0.49	RHA 1			
07573241	**Good identification**						**07573241**
Actinomyces naeslundii		100.0	0.36	RHA 1			
07573243	**Good identification**						**07573243**
Actinomyces naeslundii		100.0	0.36	RHA 1			
07573261	**Good identification**						**07573261**
Actinomyces naeslundii		100.0	0.36	RHA 1			
07573263	**Good identification**						**07573263**
Actinomyces naeslundii		100.0	0.36	RHA 1			
07573300	**Low discrimination**						**07573300**
					SPO		
Actinomyces naeslundii		62.97	0.53	RHA 1	1		
Actinomyces israelii		23.37	0.37		1		
Clostridium ramosum		13.66	0.17		92		
07573301	**Very good identification**						**07573301**
Actinomyces naeslundii		100.0	0.56	RHA 1			
07573302	**Low discrimination**						**07573302**
					SPO		
Actinomyces naeslundii		38.84	0.53	RHA 1	1		
Clostridium ramosum		33.70	0.27		92		
Actinomyces israelii		21.62	0.40		1		
07573303	**Very good identification**						**07573303**
Actinomyces naeslundii		100.0	0.56	RHA 1			
07573320	**Good identification**						**07573320**
					SPO	GRA	
Clostridium ramosum		95.35	0.51	SAL 99	92	50	
Actinomyces naeslundii		4.44	0.53		1	99	
Prevotella oralis		0.11	0.00		1	1	
07573321	**Low discrimination**						**07573321**
					SPO		
Actinomyces naeslundii		87.37	0.56	RHA 1	1		
Clostridium ramosum		12.64	0.17		92		
07573322	**Very good identification**						**07573322**
					SPO	GRA	
Clostridium ramosum		97.37	0.61	SAL 99	92	50	
Actinomyces naeslundii		1.13	0.53		1	99	
Parabacteroides distasonis		0.82	0.38		1	1	
07573323	**Low discrimination**						**07573323**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		63.35	0.56	RHA 1	1	
Clostridium ramosum		36.65	0.27		92	
07573340	**Low discrimination**					**07573340**
Actinomyces naeslundii		55.01	0.41	RHA 1		
Actinomyces israelii		44.99	0.30			
07573341	**Good identification**					**07573341**
Actinomyces naeslundii		100.0	0.44	RHA 1		
07573343	**Good identification**					**07573343**
Actinomyces naeslundii		100.0	0.44	RHA 1		
07573361	**Good identification**					**07573361**
Actinomyces naeslundii		100.0	0.44	RHA 1		
07573363	**Good identification**					**07573363**
Actinomyces naeslundii		100.0	0.44	RHA 1		
07573400	**Low discrimination**					**07573400**
					CAT	
Actinomyces israelii		67.50	0.44	SAL 92	1	
Actinomyces naeslundii		23.51	0.46	XYL 92	25	
Propionibacterium granulosum		8.99	0.24		90	
07573401	**Good identification**					**07573401**
Actinomyces naeslundii		100.0	0.49	ARA 1		
07573402	**Low discrimination**					**07573402**
Actinomyces israelii		74.77	0.47	SAL 92		
Actinomyces naeslundii		17.36	0.46	XYL 92		
Bifidobacterium adolescentis		6.21	0.23			
07573403	**Good identification**					**07573403**
Actinomyces naeslundii		97.18	0.49	ARA 1		
Actinomyces israelii		2.82	0.13			
07573420	**Low discrimination**					**07573420**
					CAT	
Actinomyces naeslundii		78.20	0.46	ARA 1	25	
Actinomyces israelii		14.33	0.24		1	
Propionibacterium granulosum		7.47	0.13		90	
07573421	**Good identification**					**07573421**
Actinomyces naeslundii		100.0	0.49	ARA 1		
07573422	**Low discrimination**					**07573422**
Actinomyces naeslundii		78.44	0.46	ARA 1		
Actinomyces israelii		21.56	0.27			
07573423	**Good identification**					**07573423**
Actinomyces naeslundii		100.0	0.49	ARA 1		
07573440	**Low discrimination**					**07573440**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		86.35	0.37	SAL 92		
Actinomyces naeslundii		13.65	0.33	XYL 92		
07573441	**Good identification**					**07573441**
Actinomyces naeslundii		100.0	0.36	ARA 1		
07573442	**Low discrimination**					**07573442**
Actinomyces israelii		88.36	0.40	SAL 92		
Actinomyces naeslundii		9.31	0.33	XYL 92		
Bifidobacterium adolescentis		2.33	0.07			
07573443	**Good identification**					**07573443**
Actinomyces naeslundii		100.0	0.36	ARA 1		
07573461	**Good identification**					**07573461**
Actinomyces naeslundii		100.0	0.36	ARA 1		
07573463	**Good identification**					**07573463**
Actinomyces naeslundii		100.0	0.36	ARA 1		
07573500	**Low discrimination**					**07573500**
					SPO	
Actinomyces israelii		76.04	0.55	SAL 92	1	
Actinomyces naeslundii		18.85	0.53	XYL 92	1	
Clostridium butyricum		2.75	0.09		99	
07573501	**Very good identification**					**07573501**
Actinomyces naeslundii		97.36	0.56	ARA 1		
Actinomyces israelii		2.65	0.20			
07573502	**Low discrimination**					**07573502**
Bifidobacterium adolescentis		63.65	0.57	XYL 87		
Actinomyces israelii		30.95	0.58			
Actinomyces naeslundii		5.11	0.53			
07573503	**Low discrimination**					**07573503**
Actinomyces naeslundii		88.92	0.56	ARA 1		
Bifidobacterium adolescentis		7.45	0.23			
Actinomyces israelii		3.62	0.23			
07573510	**Low discrimination**					**07573510**
Actinomyces israelii		74.51	0.40	SAL 92		
Bifidobacterium adolescentis		18.99	0.24	XYL 92		
Actinomyces naeslundii		6.50	0.32	MLZ 13		
07573512	**Good identification**					**07573512**
Bifidobacterium adolescentis		93.71	0.58	XYL 87		
Actinomyces israelii		5.57	0.44			
Actinomyces naeslundii		0.32	0.32			
07573520	**Low discrimination**					**07573520**
					SPO	
Actinomyces naeslundii		69.09	0.53	ARA 1	1	
Actinomyces israelii		17.79	0.34		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		11.31	0.15		92	
07573521	**Very good identification**					**07573521**
Actinomyces naeslundii		100.0	0.56	ARA 1		
07573522	**Low discrimination**					**07573522**
					SPO	
Actinomyces naeslundii		42.73	0.53	ARA 1	1	
Clostridium ramosum		27.97	0.25		92	
Actinomyces israelii		16.51	0.37		1	
07573523	**Very good identification**					**07573523**
Actinomyces naeslundii		100.0	0.56	ARA 1		
07573540	**Low discrimination**					**07573540**
Actinomyces israelii		89.10	0.48	SAL 92		
Actinomyces naeslundii		10.02	0.41	XYL 92		
Bifidobacterium adolescentis		0.88	0.07			
07573541	**Acceptable identification**					**07573541**
Actinomyces naeslundii		94.35	0.44	ARA 1		
Actinomyces israelii		5.65	0.13			
07573542	**Low discrimination**					**07573542**
Actinomyces israelii		57.85	0.51	SAL 92		
Bifidobacterium adolescentis		37.82	0.41	XYL 92		
Actinomyces naeslundii		4.34	0.41			
07573543	**Low discrimination**					**07573543**
Actinomyces naeslundii		87.07	0.44	ARA 1		
Actinomyces israelii		7.82	0.16			
Bifidobacterium adolescentis		5.11	0.07			
07573552	**Low discrimination**					**07573552**
Bifidobacterium adolescentis		84.25	0.43	XYL 87		
Actinomyces israelii		15.76	0.36	SOR 11		
07573560	**Low discrimination**					**07573560**
Actinomyces naeslundii		63.79	0.41	ARA 1		
Actinomyces israelii		36.21	0.27			
07573561	**Good identification**					**07573561**
Actinomyces naeslundii		100.0	0.44	ARA 1		
07573562	**Low discrimination**					**07573562**
Actinomyces naeslundii		51.56	0.41	ARA 1		
Actinomyces israelii		43.90	0.30			
Bifidobacterium adolescentis		4.54	0.07			
07573563	**Good identification**					**07573563**
Actinomyces naeslundii		100.0	0.44	ARA 1		
07573700	**Good identification**					**07573700**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.35	SAL 92			
07573702	**Low discrimination**						**07573702**
Actinomyces israelii		71.29	0.38	SAL 92			
Bifidobacterium adolescentis		19.68	0.23	XYL 92			
Clostridium clostridioforme		9.03	0.14	RHA 7			
07574100	**Low discrimination**						**07574100**
					COC	GRA	
Fusobacterium mortiferum		50.58	0.37	MLT 11	1	1	
Atopobium parvulum		33.19	0.23	XYL 7	99	99	
Actinomyces israelii		9.64	0.27		1	99	
07574122	**Low discrimination**						**07574122**
Parabacteroides distasonis		86.23	0.46	GAL 99			
Clostridium clostridioforme		6.68	0.22				
Prevotella oralis		5.26	0.13				
07574132	**Low discrimination**						**07574132**
Parabacteroides distasonis		60.43	0.42	GAL 99			
Clostridium clostridioforme		39.57	0.33				
07574212	**Good identification**						**07574212**
Clostridium clostridioforme		100.0	0.39	SAL 88			
07574232	**Good identification**						**07574232**
Clostridium clostridioforme		97.93	0.39	SAL 88			
Parabacteroides distasonis		2.07	0.16	ARA 94			
07574302	**Acceptable identification**						**07574302**
Clostridium clostridioforme		94.93	0.36	SAL 88			
Parabacteroides distasonis		5.07	0.20	ARA 94			
07574312	**Good identification**						**07574312**
Clostridium clostridioforme		99.37	0.48	SAL 88			
Parabacteroides distasonis		0.63	0.16	ARA 94			
07574322	**Low discrimination**						**07574322**
Parabacteroides distasonis		83.48	0.54	GAL 99			
Clostridium clostridioforme		15.80	0.36				
Prevotella oralis		0.73	0.06				
07574332	**Low discrimination**						**07574332**
Clostridium clostridioforme		61.55	0.48	SAL 88			
Parabacteroides distasonis		38.45	0.50	ARA 94			
07574412	**Good identification**						**07574412**
Clostridium clostridioforme		100.0	0.44	SAL 88			
07574432	**Good identification**						**07574432**
Clostridium clostridioforme		100.0	0.44	SAL 88			
07574502	**Low discrimination**						**07574502**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		66.22	0.42	SAL 88	66	
Bifidobacterium adolescentis		25.95	0.38	RHA 88	1	
Clostridium butyricum		4.21	0.10		99	
07574512	**Acceptable identification**					**07574512**
Clostridium clostridioforme		90.13	0.53	SAL 88		
Bifidobacterium adolescentis		9.48	0.40	RHA 88		
Bifidobacterium longum subsp. longum		0.25	0.11			
07574522	**Low discrimination**					**07574522**
					GRA	
Clostridium clostridioforme		64.85	0.42	SAL 88	50	
Parabacteroides distasonis		32.78	0.43	RHA 88	1	
Bifidobacterium longum subsp. longum		1.95	0.17		99	
07574532	**Acceptable identification**					**07574532**
					GRA	
Clostridium clostridioforme		93.37	0.53	SAL 88	50	
Parabacteroides distasonis		5.58	0.38	RHA 88	1	
Bifidobacterium longum subsp. longum		1.05	0.21		99	
07574602	**Good identification**					**07574602**
Clostridium clostridioforme		100.0	0.48	SAL 88		
07574612	**Very good identification**					**07574612**
Clostridium clostridioforme		100.0	0.59	SAL 88		
07574622	**Good identification**					**07574622**
Clostridium clostridioforme		99.31	0.48	SAL 88		
Parabacteroides distasonis		0.69	0.17			
07574632	**Very good identification**					**07574632**
Clostridium clostridioforme		99.92	0.59	SAL 88		
Parabacteroides distasonis		0.08	0.12			
07574702	**Very good identification**					**07574702**
Clostridium clostridioforme		99.79	0.57	SAL 88		
Parabacteroides distasonis		0.21	0.17			
07574712	**Very good identification**					**07574712**
Clostridium clostridioforme		99.98	0.68	SAL 88		
Parabacteroides distasonis		0.03	0.12			
07574722	**Low discrimination**					**07574722**
Clostridium clostridioforme		81.67	0.57	SAL 88		
Parabacteroides distasonis		16.89	0.51			
Bacteroides thetaiotaomicron		0.98	0.14			
07574732	**Very good identification**					**07574732**
Clostridium clostridioforme		97.44	0.68	SAL 88		
Parabacteroides distasonis		2.38	0.46			
Bacteroides thetaiotaomicron		0.10	0.07			
07575000	**Low discrimination**					**07575000**
					COC GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		65.44	0.51	SAL 92	1	99	
Atopobium parvulum		9.11	0.23		99	99	
Fusobacterium mortiferum		8.92	0.34		1	1	
07575001	**Acceptable identification**						**07575001**
Actinomyces naeslundii		91.48	0.44	XYL 1			
Actinomyces israelii		8.52	0.17				
07575002	**Low discrimination**						**07575002**
Actinomyces israelii		81.47	0.54	SAL 92			
Actinomyces naeslundii		3.93	0.41				
Bifidobacterium adolescentis		3.34	0.24				
07575003	**Low discrimination**						**07575003**
Actinomyces naeslundii		87.75	0.44	XYL 1			
Actinomyces israelii		12.25	0.20				
07575010	**Acceptable identification**						**07575010**
					CAT		
Actinomyces israelii		90.04	0.37	SAL 92	1		
Propionibacterium granulosum		5.85	0.11	MLZ 13	90		
Lactobacillus jensenii		4.12	0.02		1		
07575020	**Low discrimination**						**07575020**
Actinomyces naeslundii		28.06	0.41	XYL 1			
Actinomyces israelii		24.76	0.31				
Clostridium perfringens		17.92	0.15				
07575021	**Good identification**						**07575021**
Actinomyces naeslundii		100.0	0.44	XYL 1			
07575022	**Low discrimination**						**07575022**
					GRA		
Parabacteroides distasonis		58.95	0.46	LAC 99	1		
Prevotella oralis		22.71	0.26		1		
Actinomyces israelii		6.52	0.34		99		
07575023	**Acceptable identification**						**07575023**
					GRA		
Actinomyces naeslundii		92.55	0.44	XYL 1	99		
Parabacteroides distasonis		7.45	0.12		1		
07575032	**Low discrimination**						**07575032**
					GRA		
Parabacteroides distasonis		61.82	0.42	LAC 99	1		
Clostridium clostridioforme		36.28	0.32		50		
Actinomyces israelii		1.90	0.20		99		
07575040	**Good identification**						**07575040**
					SPO		
Actinomyces israelii		95.20	0.44	SAL 92	1		
Actinomyces naeslundii		3.13	0.28		1		
Clostridium butyricum		1.67	-.07		99		
07575042	**Good identification**						**07575042**
Actinomyces israelii		96.63	0.47	SAL 92			
Actinomyces naeslundii		2.12	0.28				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		1.26	0.09				
07575100	**Low discrimination**						**07575100**
					COC	SPO	
Atopobium parvulum		67.86	0.57	RAF 1	99	1	
Actinomyces israelii		19.70	0.61		1	1	
Clostridium butyricum		9.28	0.35		1	99	
07575101	**Low discrimination**						**07575101**
					COC		
Actinomyces naeslundii		60.86	0.52	XYL 1	1		
Atopobium parvulum		27.43	0.23		99		
Actinomyces israelii		7.96	0.27		1		
07575102	**Low discrimination**						**07575102**
					COC		
Bifidobacterium adolescentis		35.82	0.58	ARA 87	1		
Actinomyces israelii		35.34	0.64		1		
Atopobium parvulum		14.32	0.44		99		
07575103	**Low discrimination**						**07575103**
Actinomyces naeslundii		65.49	0.52	XYL 1			
Bifidobacterium adolescentis		13.03	0.24				
Actinomyces israelii		12.85	0.30				
07575110	**Low discrimination**						**07575110**
					COC	SPO	
Actinomyces israelii		63.27	0.47	SAL 92	1	1	
Atopobium parvulum		14.73	0.23	MLZ 13	99	1	
Clostridium butyricum		10.50	0.13		1	99	
07575112	**Low discrimination**						**07575112**
Bifidobacterium adolescentis		75.81	0.60	ARA 87			
Clostridium clostridioforme		9.33	0.41				
Actinomyces israelii		9.14	0.50				
07575120	**Low discrimination**						**07575120**
					COC	GRA	
Atopobium parvulum		50.00	0.44	RAF 1	99	99	
Parabacteroides distasonis		17.79	0.46		1	1	
Clostridium perfringens		6.97	0.30		1	99	
07575121	**Acceptable identification**						**07575121**
					COC	GRA	
Actinomyces naeslundii		90.27	0.52	XYL 1	1	99	
Atopobium parvulum		7.18	0.10		99	99	
Parabacteroides distasonis		2.56	0.12		1	1	
07575122	**Good identification**						**07575122**
					CAT		
Parabacteroides distasonis		91.51	0.80		76		
Prevotella oralis		5.58	0.47		1		
Bacteroides fragilis		1.04	0.25		99		
07575123	**Low discrimination**						**07575123**
					GRA		
Parabacteroides distasonis		69.96	0.46	URE 1	1		
Actinomyces naeslundii		24.98	0.52		99		
Prevotella oralis		4.27	0.13		1		
07575130	**Low discrimination**						**07575130**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		76.25	0.42	BGL 99	1		
Bifidobacterium longum subsp. longum		8.78	0.21		99		
Actinomyces israelii		6.25	0.27		99		
07575132	**Very good identification**						**07575132**
					GRA		
Parabacteroides distasonis		97.52	0.76		1		
Clostridium clostridioforme		1.94	0.41		50		
Bifidobacterium adolescentis		0.16	0.26		99		
07575133	**Acceptable identification**						**07575133**
					GRA		
Parabacteroides distasonis		94.81	0.42	URE 1	1		
Actinomyces naeslundii		3.31	0.30		99		
Clostridium clostridioforme		1.88	0.07		50		
07575140	**Low discrimination**						**07575140**
					COC	SPO	
Actinomyces israelii		81.76	0.54	SAL 92	1	1	
Clostridium butyricum		8.61	0.17		1	99	
Atopobium parvulum		7.32	0.23		99	1	
07575141	**Low discrimination**						**07575141**
Actinomyces naeslundii		77.62	0.39	XYL 1			
Actinomyces israelii		22.38	0.20				
07575142	**Low discrimination**						**07575142**
					SPO		
Actinomyces israelii		72.20	0.57	SAL 92	1		
Bifidobacterium adolescentis		23.26	0.43		1		
Clostridium butyricum		1.97	0.10		99		
07575143	**Low discrimination**						**07575143**
Actinomyces naeslundii		63.62	0.39	XYL 1			
Actinomyces israelii		27.51	0.23				
Bifidobacterium adolescentis		8.86	0.09				
07575150	**Acceptable identification**						**07575150**
					SPO		
Actinomyces israelii		92.85	0.40	SAL 92	1		
Bifidobacterium adolescentis		3.71	0.10	MLZ 13	1		
Clostridium butyricum		3.44	-.05		99		
07575152	**Low discrimination**						**07575152**
Bifidobacterium adolescentis		71.96	0.44	ARA 87			
Actinomyces israelii		27.31	0.43	SOR 11			
Clostridium clostridioforme		0.72	0.07				
07575161	**Good identification**						**07575161**
Actinomyces naeslundii		100.0	0.39	XYL 1			
07575162	**Low discrimination**						**07575162**
					GRA		
Parabacteroides distasonis		74.29	0.46	SOR 1	1		
Actinomyces israelii		12.79	0.37		99		
Prevotella oralis		4.53	0.13		1		
07575163	**Low discrimination**						**07575163**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		86.18	0.39	XYL 1	99		
Parabacteroides distasonis		13.82	0.12		1		
07575172	**Acceptable identification**						**07575172**
					GRA		
Parabacteroides distasonis		91.95	0.42	SOR 1	1		
Actinomyces israelii		4.39	0.23		99		
Bifidobacterium adolescentis		1.83	0.10		99		
07575202	**Low discrimination**						**07575202**
					GRA		
Clostridium clostridioforme		75.01	0.36	SAL 88	50		
Actinomyces israelii		19.25	0.35	ARA 94	99		
Parabacteroides distasonis		4.47	0.20		1		
07575212	**Good identification**						**07575212**
					GRA		
Clostridium clostridioforme		98.48	0.47	SAL 88	50		
Actinomyces israelii		0.83	0.21	ARA 94	99		
Parabacteroides distasonis		0.69	0.16		1		
07575222	**Low discrimination**						**07575222**
Parabacteroides distasonis		81.48	0.54	LAC 99			
Clostridium clostridioforme		13.81	0.36				
Prevotella oralis		4.49	0.20				
07575232	**Low discrimination**						**07575232**
Clostridium clostridioforme		58.92	0.47	SAL 88			
Parabacteroides distasonis		41.08	0.50	ARA 94			
07575300	**Low discrimination**						**07575300**
					COC	SPO	
Actinomyces israelii		54.15	0.42	SAL 92	1	1	
Atopobium parvulum		25.03	0.23	RHA 7	99	1	
Clostridium butyricum		10.48	0.09		1	99	
07575302	**Low discrimination**						**07575302**
					GRA		
Parabacteroides distasonis		54.58	0.54	NAG 99	1		
Clostridium clostridioforme		30.98	0.45		50		
Actinomyces israelii		9.50	0.45		99		
07575312	**Low discrimination**						**07575312**
					GRA		
Clostridium clostridioforme		81.34	0.56	SAL 88	50		
Parabacteroides distasonis		16.94	0.50	ARA 94	1		
Bifidobacterium adolescentis		0.91	0.26		99		
07575320	**Low discrimination**						**07575320**
Parabacteroides distasonis		68.34	0.54	BGL 99			
Bacteroides vulgatus		26.62	0.29				
Prevotella oralis		3.10	0.19				
07575322	**Very good identification**						**07575322**
Parabacteroides distasonis		98.45	0.89				
Prevotella oralis		0.86	0.41				
Clostridium clostridioforme		0.56	0.45				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07575323	**Good identification**					**07575323**
Parabacteroides distasonis		98.58	0.54	URE 1		
Prevotella oralis		0.86	0.06			
Clostridium clostridioforme		0.57	0.10			
07575330	**Acceptable identification**					**07575330**
Parabacteroides distasonis		94.71	0.50	BGL 99		
Clostridium clostridioforme		4.59	0.22			
Bacteroides vulgatus		0.69	-.05			
07575332	**Very good identification**					**07575332**
Parabacteroides distasonis		95.28	0.84			
Clostridium clostridioforme		4.62	0.56			
Bacteroides ovatus		0.06	0.13			
07575333	**Good identification**					**07575333**
Parabacteroides distasonis		95.37	0.50	URE 1		
Clostridium clostridioforme		4.63	0.22			
07575340	**Good identification**					**07575340**
Actinomyces israelii		100.0	0.35	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07575342	**Low discrimination**					**07575342**
					GRA	
Actinomyces israelii		78.30	0.38	SAL 92	99	
Parabacteroides distasonis		11.69	0.20	RHA 7	1	
Clostridium clostridioforme		6.63	0.10		50	
07575362	**Good identification**					**07575362**
Parabacteroides distasonis		98.16	0.54	SOR 1		
Prevotella oralis		0.86	0.06			
Clostridium clostridioforme		0.56	0.10			
07575372	**Good identification**					**07575372**
Parabacteroides distasonis		95.37	0.50	SOR 1		
Clostridium clostridioforme		4.63	0.22			
07575400	**Low discrimination**					**07575400**
					SPO	
Actinomyces israelii		52.71	0.50	SAL 92	1	
Clostridium butyricum		45.53	0.27		99	
Lactobacillus fermentum		0.64	0.16		1	
07575402	**Low discrimination**					**07575402**
Actinomyces israelii		42.75	0.53	SAL 92		
Clostridium clostridioforme		32.74	0.41			
Bifidobacterium adolescentis		14.32	0.38			
07575410	**Low discrimination**					**07575410**
					SPO	
Actinomyces israelii		56.57	0.35	SAL 92	1	
Clostridium clostridioforme		20.02	0.18	MLZ 13	66	
Clostridium butyricum		17.21	0.06		99	
07575412	**Low discrimination**					**07575412**
Clostridium clostridioforme		85.72	0.52	SAL 88		
Bifidobacterium adolescentis		10.06	0.40	RHA 88		
Actinomyces israelii		3.67	0.38			
07575422	**Low discrimination**					**07575422**
					GRA	
Clostridium clostridioforme		58.10	0.41	SAL 88	50	
Parabacteroides distasonis		32.77	0.43	RHA 88	1	
Actinomyces israelii		4.84	0.32		99	
07575432	**Acceptable identification**					**07575432**
					GRA	
Clostridium clostridioforme		92.43	0.52	SAL 88	50	
Parabacteroides distasonis		6.16	0.38	RHA 88	1	
Bifidobacterium longum subsp. longum		1.16	0.21		99	
07575440	**Low discrimination**					**07575440**
					SPO	
Actinomyces israelii		83.81	0.43	SAL 92	1	
Clostridium butyricum		16.19	0.09		99	
07575442	**Low discrimination**					**07575442**
					SPO	
Actinomyces israelii		85.00	0.46	SAL 92	1	
Bifidobacterium adolescentis		9.05	0.23		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		4.26	0.02		99		
07575500	**Low discrimination**						**07575500**
					SPO		
Clostridium butyricum		78.42	0.51	ESL 88	99		
Actinomyces israelii		15.13	0.60	SAL 99	1		
Bifidobacterium longum subsp. longum		3.61	0.41		1		
07575502	**Low discrimination**						**07575502**
					SPO		
Bifidobacterium adolescentis		71.77	0.72		1		
Clostridium butyricum		11.63	0.44		99		
Actinomyces israelii		8.66	0.63		1		
07575503	**Low discrimination**						**07575503**
					SPO		
Bifidobacterium adolescentis		72.51	0.38	URE 1	1		
Clostridium butyricum		11.75	0.10		99		
Actinomyces israelii		8.75	0.29		1		
07575510	**Low discrimination**						**07575510**
					SPO		
Bifidobacterium longum subsp. longum		39.56	0.45	CEL 1	1		
Clostridium butyricum		26.59	0.29		99		
Bifidobacterium adolescentis		14.97	0.40		1		
07575512	**Low discrimination**						**07575512**
Bifidobacterium adolescentis		74.48	0.74				
Clostridium clostridioforme		21.47	0.61				
Bifidobacterium longum subsp. longum		1.99	0.45				
07575513	**Low discrimination**						**07575513**
Bifidobacterium adolescentis		75.21	0.40	URE 1			
Clostridium clostridioforme		21.68	0.27				
Bifidobacterium longum subsp. longum		2.01	0.11				
07575520	**Low discrimination**						**07575520**
					GRA	SPO	
Bifidobacterium longum subsp. longum		45.28	0.52	CEL 1	99	1	
Bacteroides vulgatus		21.66	0.32		1	1	
Clostridium butyricum		21.41	0.33		99	99	
07575522	**Low discrimination**						**07575522**
					GRA		
Parabacteroides distasonis		85.99	0.77		1		
Clostridium clostridioforme		5.16	0.50		50		
Bifidobacterium longum subsp. longum		5.11	0.52		99		
07575523	**Low discrimination**						**07575523**
					GRA		
Parabacteroides distasonis		88.30	0.43	URE 1	1		
Clostridium clostridioforme		5.29	0.16		50		
Bifidobacterium longum subsp. longum		5.25	0.17		99		
07575530	**Acceptable identification**						**07575530**
					GRA		
Bifidobacterium longum subsp. longum		90.55	0.56	CEL 1	99		
Parabacteroides distasonis		4.87	0.38		1		
Clostridium clostridioforme		2.47	0.27		50		
07575532	**Low discrimination**						**07575532**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		57.77	0.72		1	
Clostridium clostridioforme		29.30	0.61		50	
Bifidobacterium longum subsp. longum		10.85	0.56		99	
07575533	**Low discrimination**					**07575533**
					GRA	
Parabacteroides distasonis		58.99	0.38	URE 1	1	
Clostridium clostridioforme		29.92	0.27		50	
Bifidobacterium longum subsp. longum		11.09	0.21		99	
07575542	**Low discrimination**					**07575542**
					SPO	
Bifidobacterium adolescentis		66.60	0.57	SOR 11	1	
Actinomyces israelii		25.28	0.56		1	
Clostridium butyricum		7.59	0.26		99	
07575550	**Low discrimination**					**07575550**
					SPO	
Actinomyces israelii		53.69	0.39	SAL 92	1	
Clostridium butyricum		21.91	0.11	MLZ 13	99	
Bifidobacterium adolescentis		17.53	0.24		1	
07575552	**Good identification**					**07575552**
Bifidobacterium adolescentis		92.85	0.58	SOR 11		
Actinomyces israelii		4.31	0.42			
Clostridium clostridioforme		2.19	0.27			
07575562	**Low discrimination**					**07575562**
					GRA	
Parabacteroides distasonis		63.70	0.43	SOR 1	1	
Actinomyces israelii		14.64	0.35		99	
Bifidobacterium adolescentis		6.10	0.23		99	
07575572	**Low discrimination**					**07575572**
					GRA	
Parabacteroides distasonis		50.04	0.38	SOR 1	1	
Clostridium clostridioforme		25.38	0.27		50	
Bifidobacterium adolescentis		10.88	0.24		99	
07575602	**Good identification**					**07575602**
					GRA	
Clostridium clostridioforme		98.33	0.56	SAL 88	50	
Actinomyces israelii		1.32	0.33		99	
Parabacteroides distasonis		0.23	0.17		1	
07575612	**Very good identification**					**07575612**
					GRA	
Clostridium clostridioforme		99.93	0.67	SAL 88	50	
Actinomyces israelii		0.04	0.19		99	
Parabacteroides distasonis		0.03	0.12		1	
07575622	**Low discrimination**					**07575622**
Clostridium clostridioforme		79.11	0.56	SAL 88		
Parabacteroides distasonis		18.25	0.51			
Bacteroides thetaiotaomicron		2.14	0.19			
07575632	**Very good identification**					**07575632**
Clostridium clostridioforme		97.04	0.67	SAL 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		2.65	0.46			
Bacteroides thetaiotaomicron		0.22	0.12			
07575702	**Low discrimination**					**07575702**
					GRA	
Clostridium clostridioforme		89.46	0.65	SAL 88	50	
Parabacteroides distasonis		6.17	0.51		1	
Bifidobacterium adolescentis		1.59	0.38		99	
07575710	**Acceptable identification**					**07575710**
					SPO	
Clostridium clostridioforme		92.47	0.42	SAL 88	66	
Actinomyces israelii		3.21	0.27	BGL 99	1	
Clostridium butyricum		2.40	0.03		99	
07575712	**Very good identification**					**07575712**
					GRA	
Clostridium clostridioforme		98.59	0.76	SAL 88	50	
Parabacteroides distasonis		0.80	0.46		1	
Bifidobacterium adolescentis		0.47	0.40		99	
07575713	**Good identification**					**07575713**
Clostridium clostridioforme		99.19	0.42	SAL 88		
Parabacteroides distasonis		0.81	0.12	URE 1		
07575720	**Low discrimination**					**07575720**
Bacteroides vulgatus		86.74	0.48	TRE 4		
Parabacteroides distasonis		10.27	0.51	CEL 5		
Clostridium clostridioforme		1.51	0.31			
07575722	**Low discrimination**					**07575722**
Parabacteroides distasonis		81.08	0.85			
Clostridium clostridioforme		11.88	0.65			
Bacteroides thetaiotaomicron		4.71	0.48			
07575723	**Low discrimination**					**07575723**
Parabacteroides distasonis		81.18	0.51	URE 1		
Clostridium clostridioforme		11.90	0.31			
Bacteroides thetaiotaomicron		4.71	0.14			
07575730	**Low discrimination**					**07575730**
Clostridium clostridioforme		48.88	0.42	SAL 88		
Parabacteroides distasonis		39.42	0.46	BGL 99		
Bacteroides vulgatus		6.24	0.14			
07575732	**Low discrimination**					**07575732**
Clostridium clostridioforme		53.53	0.76	SAL 88		
Parabacteroides distasonis		43.17	0.80			
Bacteroides thetaiotaomicron		1.81	0.41			
07575733	**Low discrimination**					**07575733**
Clostridium clostridioforme		53.55	0.42	SAL 88		
Parabacteroides distasonis		43.18	0.46	URE 1		
Bacteroides thetaiotaomicron		1.81	0.07			
07575752	**Acceptable identification**					**07575752**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		92.01	0.42	SAL 88		
Bifidobacterium adolescentis		5.38	0.24	SOR 1		
Actinomyces israelii		1.86	0.23			
07575762	**Low discrimination**					**07575762**
Parabacteroides distasonis		75.59	0.51	SOR 1		
Clostridium clostridioforme		11.08	0.31			
Bacteroides ovatus		8.51	0.19			
07575772	**Low discrimination**					**07575772**
Clostridium clostridioforme		51.21	0.42	SAL 88		
Parabacteroides distasonis		41.30	0.46	SOR 1		
Bacteroides ovatus		5.76	0.16			
07576100	**Acceptable identification**					**07576100**
Actinomyces israelii		93.22	0.40	SAL 92		
Actinomyces naeslundii		6.78	0.30	GAL 99		
07576102	**Low discrimination**					**07576102**
Actinomyces israelii		83.69	0.43	SAL 92		
Bifidobacterium adolescentis		12.25	0.23	GAL 99		
Actinomyces naeslundii		4.06	0.30			
07576142	**Good identification**					**07576142**
Actinomyces israelii		95.56	0.36	SAL 92		
Bifidobacterium adolescentis		4.44	0.07	GAL 99		
07576500	**Acceptable identification**					**07576500**
					SPO	
Actinomyces israelii		90.78	0.39	SAL 92	1	
Clostridium butyricum		9.22	0.01	GAL 99	99	
07576502	**Low discrimination**					**07576502**
Bifidobacterium adolescentis		52.94	0.37	GAL 99		
Actinomyces israelii		44.23	0.42			
Clostridium clostridioforme		1.67	0.08			
07576512	**Low discrimination**					**07576512**
Bifidobacterium adolescentis		80.23	0.38	GAL 99		
Clostridium clostridioforme		9.42	0.19			
Actinomyces israelii		8.19	0.28			
07577000	**Very good identification**					**07577000**
					CAT	
Actinomyces israelii		96.54	0.64	SAL 92	1	
Actinomyces naeslundii		2.39	0.46		25	
Propionibacterium granulosum		0.92	0.24		90	
07577001	**Low discrimination**					**07577001**
Actinomyces naeslundii		78.63	0.49	XYL 1		
Actinomyces israelii		21.37	0.30			
07577002	**Very good identification**					**07577002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		97.66	0.67	SAL 92		
Actinomyces naeslundii		1.61	0.46			
Bifidobacterium adolescentis		0.58	0.23			
07577003	**Low discrimination**					**07577003**
Actinomyces naeslundii		71.04	0.49	XYL 1		
Actinomyces israelii		28.96	0.33			
07577010	**Good identification**					**07577010**
					CAT	
Actinomyces israelii		98.29	0.50	SAL 92	1	
Actinomyces naeslundii		0.86	0.24	MLZ 13	25	
Propionibacterium granulosum		0.85	0.09		90	
07577012	**Acceptable identification**					**07577012**
Actinomyces israelii		94.86	0.53	SAL 92		
Bifidobacterium adolescentis		4.59	0.24	MLZ 13		
Actinomyces naeslundii		0.55	0.24			
07577020	**Low discrimination**					**07577020**
					CAT	
Actinomyces israelii		70.16	0.44	SAL 92	1	
Actinomyces naeslundii		27.24	0.46	NAG 6	25	
Propionibacterium granulosum		2.60	0.13		90	
07577021	**Good identification**					**07577021**
Actinomyces naeslundii		100.0	0.49	XYL 1		
07577022	**Low discrimination**					**07577022**
					GRA	
Actinomyces israelii		78.44	0.47	SAL 92	99	
Actinomyces naeslundii		20.30	0.46	NAG 6	99	
Parabacteroides distasonis		1.26	0.12		1	
07577023	**Good identification**					**07577023**
Actinomyces naeslundii		100.0	0.49	XYL 1		
07577040	**Very good identification**					**07577040**
Actinomyces israelii		98.89	0.57	SAL 92		
Actinomyces naeslundii		1.11	0.33			
07577041	**Low discrimination**					**07577041**
Actinomyces naeslundii		62.55	0.36	XYL 1		
Actinomyces israelii		37.45	0.23			
07577042	**Very good identification**					**07577042**
Actinomyces israelii		99.07	0.60	SAL 92		
Actinomyces naeslundii		0.74	0.33			
Bifidobacterium adolescentis		0.19	0.07			
07577043	**Low discrimination**					**07577043**
Actinomyces naeslundii		52.68	0.36	XYL 1		
Actinomyces israelii		47.32	0.26			
07577050	**Good identification**					**07577050**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.43	SAL 92		
07577052	**Good identification**					**07577052**
Actinomyces israelii		98.49	0.46	SAL 92		
Bifidobacterium adolescentis		1.51	0.09	MLZ 13		
07577060	**Low discrimination**					**07577060**
Actinomyces israelii		85.02	0.36	SAL 92		
Actinomyces naeslundii		14.98	0.33	NAG 6		
07577061	**Good identification**					**07577061**
Actinomyces naeslundii		100.0	0.36	XYL 1		
07577062	**Low discrimination**					**07577062**
Actinomyces israelii		89.49	0.40	SAL 92		
Actinomyces naeslundii		10.51	0.33	NAG 6		
07577063	**Good identification**					**07577063**
Actinomyces naeslundii		100.0	0.36	XYL 1		
07577100	**Very good identification**					**07577100**
					SPO	
Actinomyces israelii		96.55	0.74	SAL 92	1	
Actinomyces naeslundii		1.70	0.53		1	
Clostridium butyricum		0.89	0.18		99	
07577101	**Low discrimination**					**07577101**
Actinomyces naeslundii		72.36	0.56	XYL 1		
Actinomyces israelii		27.64	0.40			
07577102	**Low discrimination**					**07577102**
Actinomyces israelii		86.03	0.77	SAL 92		
Bifidobacterium adolescentis		12.59	0.57			
Actinomyces naeslundii		1.01	0.53			
07577103	**Low discrimination**					**07577103**
Actinomyces naeslundii		60.36	0.56	XYL 1		
Actinomyces israelii		34.58	0.43			
Bifidobacterium adolescentis		5.06	0.23			
07577110	**Very good identification**					**07577110**
Actinomyces israelii		96.79	0.60	SAL 92		
Bifidobacterium adolescentis		1.76	0.24	MLZ 13		
Actinomyces naeslundii		0.60	0.32			
07577112	**Low discrimination**					**07577112**
					SPO	
Bifidobacterium adolescentis		54.01	0.58	ARA 87	1	
Actinomyces israelii		45.12	0.63		1	
Clostridium tertium		0.50	0.15		99	
07577120	**Low discrimination**					**07577120**
					SPO	
Actinomyces israelii		72.51	0.54	SAL 92	1	
Actinomyces naeslundii		20.03	0.53	NAG 6	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		3.28	0.15		92		
07577121	**Very good identification**						**07577121**
Actinomyces naeslundii		97.62	0.56	XYL 1			
Actinomyces israelii		2.38	0.20				
07577122	**Low discrimination**						**07577122**
					GRA		
Actinomyces israelii		55.47	0.57	SAL 92	99		
Parabacteroides distasonis		22.11	0.46	NAG 6	1		
Actinomyces naeslundii		10.22	0.53		99		
07577123	**Very good identification**						**07577123**
					GRA		
Actinomyces naeslundii		95.14	0.56	XYL 1	99		
Actinomyces israelii		3.48	0.23		99		
Parabacteroides distasonis		1.39	0.12		1		
07577130	**Low discrimination**						**07577130**
					SPO		
Actinomyces israelii		73.53	0.40	SAL 92	1		
Clostridium tertium		19.32	0.15	MLZ 13	99		
Actinomyces naeslundii		7.16	0.32	NAG 6	1		
07577132	**Low discrimination**						**07577132**
					GRA	SPO	
Parabacteroides distasonis		43.46	0.42	MAN 1	1	1	
Actinomyces israelii		30.26	0.43		99	1	
Clostridium tertium		15.90	0.24		99	99	
07577140	**Very good identification**						**07577140**
					SPO		
Actinomyces israelii		98.93	0.67	SAL 92	1		
Actinomyces naeslundii		0.79	0.41		1		
Clostridium butyricum		0.20	0.00		99		
07577141	**Low discrimination**						**07577141**
Actinomyces naeslundii		54.30	0.44	XYL 1			
Actinomyces israelii		45.70	0.33				
07577142	**Very good identification**						**07577142**
Actinomyces israelii		95.02	0.70	SAL 92			
Bifidobacterium adolescentis		4.42	0.41				
Actinomyces naeslundii		0.51	0.41				
07577143	**Low discrimination**						**07577143**
Actinomyces israelii		54.39	0.36	SAL 92			
Actinomyces naeslundii		43.08	0.44	URE 1			
Bifidobacterium adolescentis		2.53	0.07				
07577150	**Good identification**						**07577150**
					COC		
Actinomyces israelii		98.62	0.53	SAL 92	1		
Blautia producta		0.82	-.03	MLZ 13	99		
Bifidobacterium adolescentis		0.57	0.09		1		
07577152	**Low discrimination**						**07577152**
Actinomyces israelii		72.44	0.56	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		27.56	0.43	MLZ 13		
07577160	**Low discrimination**					**07577160**
Actinomyces israelii		88.86	0.47	SAL 92		
Actinomyces naeslundii		11.14	0.41	NAG 6		
07577161	**Good identification**					**07577161**
Actinomyces naeslundii		100.0	0.44	XYL 1		
07577162	**Acceptable identification**					**07577162**
					GRA	
Actinomyces israelii		90.80	0.50	SAL 92	99	
Actinomyces naeslundii		7.59	0.41	NAG 6	99	
Parabacteroides distasonis		0.94	0.12		1	
07577163	**Acceptable identification**					**07577163**
Actinomyces naeslundii		92.54	0.44	XYL 1		
Actinomyces israelii		7.46	0.16			
07577172	**Acceptable identification**					**07577172**
Actinomyces israelii		94.32	0.36	SAL 92		
Bifidobacterium adolescentis		5.68	0.09	MLZ 13		
07577200	**Good identification**					**07577200**
Actinomyces israelii		100.0	0.45	SAL 92		
07577202	**Good identification**					**07577202**
Actinomyces israelii		100.0	0.48	SAL 92		
07577240	**Good identification**					**07577240**
Actinomyces israelii		100.0	0.38	SAL 92		
07577242	**Good identification**					**07577242**
Actinomyces israelii		100.0	0.41	SAL 92		
07577300	**Good identification**					**07577300**
					SPO	
Actinomyces israelii		99.62	0.55	SAL 92	1	
Clostridium butyricum		0.38	-.08	RHA 7	99	
07577302	**Very good identification**					**07577302**
					GRA	
Actinomyces israelii		96.55	0.58	SAL 92	99	
Bifidobacterium adolescentis		1.90	0.23	RHA 7	99	
Parabacteroides distasonis		0.99	0.20		1	
07577310	**Good identification**					**07577310**
Actinomyces israelii		100.0	0.41	SAL 92		
07577312	**Low discrimination**					**07577312**
Actinomyces israelii		72.74	0.44	SAL 92		
Clostridium clostridioforme		12.90	0.22	RHA 7		
Bifidobacterium adolescentis		11.68	0.24	MLZ 13		
07577322	**Low discrimination**					**07577322**
					SPO GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		81.43	0.54	MAN 1	1	1	
Clostridium ramosum		10.88	0.27		92	50	
Actinomyces israelii		5.13	0.38		1	99	
07577332	**Acceptable identification**						**07577332**
Parabacteroides distasonis		91.99	0.50	MAN 1			
Clostridium clostridioforme		4.46	0.22				
Bacteroides ovatus		1.95	0.05				
07577340	**Good identification**						**07577340**
Actinomyces israelii		100.0	0.48	SAL 92			
07577342	**Good identification**						**07577342**
Actinomyces israelii		99.38	0.51	SAL 92			
Bifidobacterium adolescentis		0.62	0.07	RHA 7			
07577352	**Good identification**						**07577352**
Actinomyces israelii		95.14	0.37	SAL 92			
Bifidobacterium adolescentis		4.86	0.09	RHA 7			
07577400	**Very good identification**						**07577400**
					SPO		
Actinomyces israelii		98.34	0.62	SAL 92	1		
Clostridium butyricum		1.67	0.11		99		
07577402	**Good identification**						**07577402**
					SPO		
Actinomyces israelii		94.87	0.65	SAL 92	1		
Bifidobacterium adolescentis		4.59	0.37		1		
Clostridium butyricum		0.42	0.04		99		
07577410	**Good identification**						**07577410**
Actinomyces israelii		100.0	0.48	SAL 92			
07577412	**Low discrimination**						**07577412**
Actinomyces israelii		69.07	0.51	SAL 92			
Bifidobacterium adolescentis		27.32	0.38	MLZ 13			
Clostridium clostridioforme		2.88	0.18				
07577420	**Good identification**						**07577420**
					SPO		
Actinomyces israelii		97.75	0.42	SAL 92	1		
Clostridium butyricum		2.26	-.07	NAG 6	99		
07577422	**Good identification**						**07577422**
Actinomyces israelii		97.91	0.45	SAL 92			
Clostridium clostridioforme		2.09	0.07	NAG 6			
07577440	**Good identification**						**07577440**
					SPO		
Actinomyces israelii		99.62	0.55	SAL 92	1		
Clostridium butyricum		0.38	-.07		99		
07577442	**Very good identification**						**07577442**
Actinomyces israelii		98.49	0.58	SAL 92			
Bifidobacterium adolescentis		1.51	0.21				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07577450	**Good identification**					**07577450**
					COC	
Actinomyces israelii		96.11	0.41	SAL 92	1	
Blautia producta		3.89	-.03	MLZ 13	99	
07577452	**Low discrimination**					**07577452**
Actinomyces israelii		88.83	0.44	SAL 92		
Bifidobacterium adolescentis		11.17	0.23	MLZ 13		
07577460	**Good identification**					**07577460**
Actinomyces israelii		100.0	0.35	SAL 92		
07577462	**Good identification**					**07577462**
Actinomyces israelii		100.0	0.38	SAL 92		
07577500	**Low discrimination**					**07577500**
					SPO	
Actinomyces israelii		89.27	0.73	SAL 92	1	
Clostridium butyricum		9.07	0.35		99	
Bifidobacterium adolescentis		1.62	0.37		1	
07577501	**Acceptable identification**					**07577501**
					SPO	
Actinomyces israelii		90.78	0.39	SAL 92	1	
Clostridium butyricum		9.22	0.01	URE 1	99	
07577502	**Low discrimination**					**07577502**
					SPO	
Bifidobacterium adolescentis		53.79	0.71		1	
Actinomyces israelii		44.94	0.76		1	
Clostridium butyricum		1.18	0.28		99	
07577503	**Low discrimination**					**07577503**
					SPO	
Bifidobacterium adolescentis		53.84	0.37	URE 1	1	
Actinomyces israelii		44.98	0.42		1	
Clostridium butyricum		1.18	-.07		99	
07577510	**Low discrimination**					**07577510**
Actinomyces israelii		81.17	0.59	SAL 92		
Bifidobacterium adolescentis		12.04	0.38	MLZ 13		
Bifidobacterium dentium		3.17	0.10			
07577512	**Low discrimination**					**07577512**
Bifidobacterium adolescentis		88.19	0.72			
Actinomyces israelii		9.01	0.62			
Bifidobacterium dentium		2.37	0.27			
07577513	**Low discrimination**					**07577513**
Bifidobacterium adolescentis		88.57	0.38	URE 1		
Actinomyces israelii		9.05	0.28			
Bifidobacterium dentium		2.38	-.07			
07577520	**Low discrimination**					**07577520**
					SPO	
Actinomyces israelii		84.91	0.52	SAL 92	1	
Clostridium butyricum		11.75	0.17	NAG 6	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		2.26	0.17		1		
07577522	**Low discrimination**						**07577522**
					GRA		
Actinomyces israelii		61.84	0.55	SAL 92	99		
Parabacteroides distasonis		18.47	0.43	NAG 6	1		
Bifidobacterium adolescentis		11.71	0.37		99		
07577540	**Good identification**						**07577540**
					COC	SPO	
Actinomyces israelii		92.73	0.66	SAL 92	1	1	
Blautia producta		4.63	0.24		99	1	
Clostridium butyricum		2.11	0.17		1	99	
07577542	**Low discrimination**						**07577542**
					SPO		
Actinomyces israelii		71.94	0.69	SAL 92	1		
Bifidobacterium adolescentis		27.37	0.55		1		
Clostridium butyricum		0.42	0.10		99		
07577552	**Low discrimination**						**07577552**
					COC		
Bifidobacterium adolescentis		74.22	0.57	SOR 11	1		
Actinomyces israelii		23.85	0.55		1		
Blautia producta		1.77	0.15		99		
07577560	**Low discrimination**						**07577560**
					COC	SPO	
Actinomyces israelii		89.46	0.45	SAL 92	1	1	
Blautia producta		7.77	0.07	NAG 6	99	1	
Clostridium butyricum		2.77	-.02		1	99	
07577562	**Acceptable identification**						**07577562**
Actinomyces israelii		94.32	0.48	SAL 92			
Bifidobacterium adolescentis		5.68	0.21	NAG 6			
07577600	**Good identification**						**07577600**
Actinomyces israelii		100.0	0.43	SAL 92			
07577602	**Low discrimination**						**07577602**
Actinomyces israelii		88.26	0.46	SAL 92			
Clostridium clostridioforme		11.74	0.22	RHA 7			
07577640	**Good identification**						**07577640**
Actinomyces israelii		100.0	0.36	SAL 92			
07577642	**Good identification**						**07577642**
Actinomyces israelii		100.0	0.39	SAL 92			
07577700	**Good identification**						**07577700**
					SPO		
Actinomyces israelii		95.99	0.54	SAL 92	1		
Clostridium butyricum		4.01	0.09	RHA 7	99		
07577702	**Low discrimination**						**07577702**
Actinomyces israelii		77.08	0.57	SAL 92			
Bifidobacterium adolescentis		12.38	0.37	RHA 7			
Clostridium clostridioforme		8.58	0.31				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
07577710	**Acceptable identification**							**07577710**
					COC			
Actinomyces israelii		90.60	0.39	SAL 92	1			
Blautia producta		4.74	-.03	RHA 7	99			
Clostridium clostridioforme		4.66	0.08	MLZ 13	1			
07577712	**Low discrimination**							**07577712**
Clostridium clostridioforme		58.50	0.42	SAL 88				
Bifidobacterium adolescentis		22.65	0.38	MAN 1				
Actinomyces israelii		17.24	0.42					
07577722	**Low discrimination**							**07577722**
Parabacteroides distasonis		45.67	0.51	MAN 1				
Bacteroides ovatus		41.15	0.34					
Clostridium clostridioforme		6.69	0.31					
07577732	**Low discrimination**							**07577732**
Clostridium clostridioforme		36.01	0.42	SAL 88				
Bacteroides ovatus		32.39	0.31	MAN 1				
Parabacteroides distasonis		29.04	0.46					
07577740	**Low discrimination**							**07577740**
					COC			
Actinomyces israelii		60.14	0.47	SAL 92	1			
Blautia producta		39.87	0.24	RHA 7	99			
07577742	**Acceptable identification**							**07577742**
					COC			
Actinomyces israelii		90.90	0.50	SAL 92	1			
Bifidobacterium adolescentis		4.64	0.21	RHA 7	1			
Blautia producta		4.46	0.07		99			
07577752	**Low discrimination**							**07577752**
					COC			
Actinomyces israelii		40.14	0.35	SAL 92	1			
Blautia producta		39.57	0.15	RHA 7	99			
Bifidobacterium adolescentis		16.76	0.23	MLZ 13	1			
07600000	**Acceptable identification**							**07600000**
					COC	GRA	CAT	
Fusobacterium mortiferum		94.53	0.54	FRU 99	1	1	5	
Gemella morbillorum		2.88	0.17		99	99	1	
Propionibacterium granulosum		2.59	0.24		1	99	90	
07600002	**Acceptable identification**							**07600002**
					GRA	CAT		
Fusobacterium mortiferum		94.18	0.37	FRU 99	1	5		
Propionibacterium granulosum		5.82	0.13	BGL 10	99	90		
07600100	**Very good identification**							**07600100**
Fusobacterium mortiferum		100.0	0.60	FRU 99				
07600102	**Good identification**							**07600102**
					GRA			
Fusobacterium mortiferum		98.47	0.44	FRU 99	1			
Actinomyces viscosus		1.53	0.00	BGL 10	99			
07601000	**Low discrimination**							**07601000**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		86.81	0.57	FRU 99	1	5	
Eubacterium tortuosum		9.65	0.24		99	1	
Propionibacterium granulosum		1.59	0.24		99	90	
07601001	**Low discrimination**						**07601001**
					GRA		
Actinomyces naeslundii		63.04	0.38	MLT 94	99		
Fusobacterium mortiferum		36.96	0.22	FRU 95	1		
07601002	**Low discrimination**						**07601002**
					GRA		
Fusobacterium mortiferum		75.69	0.40	FRU 99	1		
Eubacterium tortuosum		13.37	0.11	BGL 10	99		
Actinomyces naeslundii		7.83	0.35		99		
07601003	**Good identification**						**07601003**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07601021	**Good identification**						**07601021**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07601023	**Good identification**						**07601023**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07601100	**Very good identification**						**07601100**
					GRA		
Fusobacterium mortiferum		98.49	0.63	FRU 99	1		
Actinomyces naeslundii		1.38	0.43		99		
Lactobacillus fermentum		0.14	0.15		99		
07601101	**Low discrimination**						**07601101**
					GRA		
Actinomyces naeslundii		67.54	0.46	MLT 94	99		
Fusobacterium mortiferum		32.46	0.29	FRU 95	1		
07601102	**Low discrimination**						**07601102**
					GRA		
Fusobacterium mortiferum		87.21	0.47	FRU 99	1		
Actinomyces naeslundii		11.00	0.43	BGL 10	99		
Lactobacillus fermentum		1.08	0.15		99		
07601103	**Acceptable identification**						**07601103**
					GRA		
Actinomyces naeslundii		94.93	0.46	MLT 94	99		
Fusobacterium mortiferum		5.07	0.12	FRU 95	1		
07601121	**Good identification**						**07601121**
Actinomyces naeslundii		100.0	0.46	MLT 94			
07601122	**Low discrimination**						**07601122**
					GRA		
Actinomyces naeslundii		51.91	0.43	MLT 94	99		
Prevotella oralis		18.63	0.09	FRU 95	1		
Prevotella melaninogenica		12.69	0.03		1		
07601123	**Good identification**						**07601123**
Actinomyces naeslundii		100.0	0.46	MLT 94			
07603000	**Low discrimination**						**07603000**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		39.65	0.40	MLT 94	99	25	
Propionibacterium granulosum		24.54	0.21	FRU 95	99	90	
Fusobacterium mortiferum		17.96	0.22		1	5	
07603001	**Good identification**						**07603001**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07603002	**Low discrimination**						**07603002**
					SPO	CAT	
Actinomyces naeslundii		62.31	0.40	MLT 94	1	25	
Clostridium innocuum		28.04	0.10	FRU 95	99	1	
Propionibacterium granulosum		9.64	0.11		1	90	
07603003	**Good identification**						**07603003**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07603020	**Low discrimination**						**07603020**
Actinomyces naeslundii		86.60	0.40	MLT 94			
Propionibacterium granulosum		13.40	0.11	FRU 95			
07603021	**Good identification**						**07603021**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07603022	**Good identification**						**07603022**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07603023	**Good identification**						**07603023**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07603100	**Low discrimination**						**07603100**
					GRA		
Actinomyces naeslundii		72.92	0.48	MLT 94	99		
Fusobacterium mortiferum		27.08	0.29	FRU 95	1		
07603101	**Good identification**						**07603101**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07603102	**Good identification**						**07603102**
					GRA		
Actinomyces naeslundii		96.04	0.48	MLT 94	99		
Fusobacterium mortiferum		3.96	0.12	FRU 95	1		
07603103	**Good identification**						**07603103**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07603120	**Good identification**						**07603120**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07603121	**Good identification**						**07603121**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07603122	**Good identification**						**07603122**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07603123	**Good identification**						**07603123**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		100.0	0.51	MLT 94				
07603141	**Good identification**							**07603141**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07603143	**Good identification**							**07603143**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07603161	**Good identification**							**07603161**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07603163	**Good identification**							**07603163**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07604100	**Good identification**							**07604100**
Fusobacterium mortiferum		100.0	0.41	FRU 99				
07605000	**Good identification**							**07605000**
Fusobacterium mortiferum		100.0	0.37	FRU 99				
07605100	**Good identification**							**07605100**
Fusobacterium mortiferum		100.0	0.44	FRU 99				
07605322	**Good identification**							**07605322**
Parabacteroides distasonis		100.0	0.39	FRU 99				
07605722	**Good identification**							**07605722**
Parabacteroides distasonis		100.0	0.35	FRU 99				
07610000	**Very good identification**							**07610000**
					COC	GRA	CAT	
Fusobacterium mortiferum		97.26	0.88		1	1	5	
Propionibacterium granulosum		2.67	0.58		1	99	90	
Gemella morbillorum		0.04	0.19		99	99	1	
07610001	**Good identification**							**07610001**
					GRA	CAT		
Fusobacterium mortiferum		95.43	0.54	URE 1	1	5		
Propionibacterium granulosum		2.62	0.24		99	90		
Actinomyces naeslundii		1.95	0.36		99	25		
07610002	**Good identification**							**07610002**
					GRA	CAT		
Fusobacterium mortiferum		93.73	0.71	BGL 10	1	5		
Propionibacterium granulosum		5.79	0.47		99	90		
Actinomyces viscosus		0.37	0.18		99	76		
07610003	**Low discrimination**							**07610003**
					GRA			
Fusobacterium mortiferum		72.73	0.37	URE 1	1			
Actinomyces naeslundii		13.39	0.36	BGL 10	99			
Actinomyces viscosus		9.38	0.09		99			
07610010	**Low discrimination**							**07610010**
					GRA	SPO	CAT	
Fusobacterium mortiferum		72.72	0.54	MLZ 1	1	1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		26.94	0.43		99	1	90	
Clostridium difficile		0.34	-.02		99	99	1	
07610012	**Low discrimination**							**07610012**
					GRA	CAT		
Fusobacterium mortiferum		54.54	0.37	MLZ 1	1	5		
Propionibacterium granulosum		45.46	0.33	BGL 10	99	90		
07610020	**Low discrimination**							**07610020**
					GRA	CAT		
Fusobacterium mortiferum		58.93	0.54	NAG 1	1	5		
Propionibacterium granulosum		40.02	0.47		99	90		
Actinomyces naeslundii		0.80	0.33		99	25		
07610021	**Low discrimination**							**07610021**
					GRA	CAT		
Actinomyces naeslundii		54.67	0.36	MLT 94	99	25		
Fusobacterium mortiferum		26.99	0.19	GAL 96	1	5		
Propionibacterium granulosum		18.33	0.13		99	90		
07610022	**Low discrimination**							**07610022**
					GRA	CAT		
Propionibacterium granulosum		57.65	0.37	SAL 1	99	90		
Fusobacterium mortiferum		37.72	0.37		1	5		
Actinomyces naeslundii		4.63	0.33		99	25		
07610023	**Good identification**							**07610023**
Actinomyces naeslundii		100.0	0.36	MLT 94				
07610040	**Good identification**							**07610040**
					GRA	CAT		
Fusobacterium mortiferum		97.33	0.54	SOR 1	1	5		
Propionibacterium granulosum		2.67	0.24		99	90		
07610042	**Acceptable identification**							**07610042**
					GRA	CAT		
Fusobacterium mortiferum		94.18	0.37	SOR 1	1	5		
Propionibacterium granulosum		5.82	0.13	BGL 10	99	90		
07610100	**Excellent identification**							**07610100**
					GRA	CAT		
Fusobacterium mortiferum		99.97	0.94		1	5		
Actinomyces naeslundii		0.02	0.41		99	25		
Propionibacterium granulosum		0.01	0.24		99	90		
07610101	**Very good identification**							**07610101**
					GRA			
Fusobacterium mortiferum		97.57	0.60	URE 1	1			
Actinomyces naeslundii		2.44	0.44		99			
07610102	**Very good identification**							**07610102**
					GRA			
Fusobacterium mortiferum		99.50	0.78	BGL 10	1			
Actinomyces viscosus		0.30	0.22		99			
Actinomyces naeslundii		0.15	0.41		99			
07610103	**Low discrimination**							**07610103**
					GRA			
Fusobacterium mortiferum		75.59	0.44	URE 1	1			
Actinomyces naeslundii		16.98	0.44	BGL 10	99			
Actinomyces viscosus		7.43	0.14		99			
07610110	**Very good identification**							**07610110**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Fusobacterium mortiferum		99.84	0.60	MLZ 1	1	5		
Propionibacterium granulosum		0.16	0.09		99	90		
07610112	**Good identification**							**07610112**
Fusobacterium mortiferum		100.0	0.44	MLZ 1				
07610120	**Very good identification**							**07610120**
					GRA	CAT		
Fusobacterium mortiferum		97.61	0.60	NAG 1	1	5		
Actinomyces naeslundii		1.62	0.41		99	25		
Propionibacterium granulosum		0.29	0.13		99	90		
07610121	**Low discrimination**							**07610121**
					GRA			
Actinomyces naeslundii		71.19	0.44	MLT 94	99			
Fusobacterium mortiferum		28.81	0.26	GAL 96	1			
07610122	**Low discrimination**							**07610122**
					GRA			
Fusobacterium mortiferum		79.14	0.44	NAG 1	1			
Actinomyces naeslundii		11.85	0.41	BGL 10	99			
Prevotella oralis		5.37	0.09		1			
07610123	**Good identification**							**07610123**
Actinomyces naeslundii		100.0	0.44	MLT 94				
07610140	**Very good identification**							**07610140**
					GRA			
Fusobacterium mortiferum		99.71	0.60	SOR 1	1			
Actinomyces naeslundii		0.29	0.28		99			
07610142	**Good identification**							**07610142**
					GRA			
Fusobacterium mortiferum		97.43	0.44	SOR 1	1			
Actinomyces naeslundii		2.57	0.28	BGL 10	99			
07610200	**Acceptable identification**							**07610200**
					GRA	CAT		
Fusobacterium mortiferum		92.25	0.54	RHA 1	1	5		
Propionibacterium granulosum		7.75	0.32		99	90		
07610202	**Low discrimination**							**07610202**
					GRA	CAT		
Fusobacterium mortiferum		84.10	0.37	RHA 1	1	5		
Propionibacterium granulosum		15.90	0.22	BGL 10	99	90		
07610300	**Very good identification**							**07610300**
Fusobacterium mortiferum		100.0	0.60	RHA 1				
07610302	**Good identification**							**07610302**
Fusobacterium mortiferum		100.0	0.44	RHA 1				
07610400	**Good identification**							**07610400**
					COC	GRA	CAT	
Fusobacterium mortiferum		96.94	0.54	ARA 1	1	1	5	
Propionibacterium granulosum		2.66	0.24		1	99	90	
Gemella morbillorum		0.40	0.03		99	99	1	
07610402	**Acceptable identification**							**07610402**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		94.18	0.37	ARA 1	1	5	
Propionibacterium granulosum		5.82	0.13	BGL 10	99	90	
07610500	**Very good identification**						**07610500**
					GRA		
Fusobacterium mortiferum		99.80	0.60	ARA 1	1		
Lactobacillus fermentum		0.21	0.15		99		
07610502	**Good identification**						**07610502**
					GRA		
Fusobacterium mortiferum		98.18	0.44	ARA 1	1		
Lactobacillus fermentum		1.82	0.15	BGL 10	99		
07611000	**Very good identification**						**07611000**
					GRA	CAT	
Fusobacterium mortiferum		97.75	0.91		1	5	
Propionibacterium granulosum		1.79	0.58		99	90	
Actinomyces naeslundii		0.22	0.57		99	25	
07611001	**Low discrimination**						**07611001**
					GRA	CAT	
Fusobacterium mortiferum		74.31	0.57	URE 1	1	5	
Actinomyces naeslundii		24.33	0.60		99	25	
Propionibacterium granulosum		1.36	0.24		99	90	
07611002	**Good identification**						**07611002**
					GRA	CAT	
Fusobacterium mortiferum		92.34	0.74	BGL 10	1	5	
Propionibacterium granulosum		3.80	0.47		99	90	
Actinomyces naeslundii		1.83	0.57		99	25	
07611003	**Low discrimination**						**07611003**
					GRA		
Actinomyces naeslundii		72.92	0.60	MLT 94	99		
Fusobacterium mortiferum		24.75	0.40		1		
Actinomyces viscosus		1.31	0.06		99		
07611010	**Low discrimination**						**07611010**
					GRA	CAT	
Fusobacterium mortiferum		79.20	0.57	MLZ 1	1	5	
Propionibacterium granulosum		19.56	0.43		99	90	
Actinomyces naeslundii		0.91	0.35		99	25	
07611011	**Low discrimination**						**07611011**
					GRA	CAT	
Actinomyces naeslundii		57.77	0.38	MLT 94	99	25	
Fusobacterium mortiferum		33.87	0.22	MLZ 5	1	5	
Propionibacterium granulosum		8.37	0.09		99	90	
07611012	**Low discrimination**						**07611012**
					GRA	CAT	
Fusobacterium mortiferum		59.39	0.40	MLZ 1	1	5	
Propionibacterium granulosum		33.00	0.33	BGL 10	99	90	
Actinomyces naeslundii		6.14	0.35		99	25	
07611013	**Good identification**						**07611013**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07611020	**Low discrimination**						**07611020**
					GRA	CAT	
Fusobacterium mortiferum		55.53	0.57	NAG 1	1	5	
Propionibacterium granulosum		25.15	0.47		99	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		12.12	0.57		99	25	
07611021	**Very good identification**						**07611021**
					GRA	CAT	
Actinomyces naeslundii		95.71	0.60	MLT 94	99	25	
Fusobacterium mortiferum		2.95	0.22		1	5	
Propionibacterium granulosum		1.34	0.13		99	90	
07611022	**Low discrimination**						**07611022**
					GRA	CAT	
Actinomyces naeslundii		34.91	0.57	MLT 94	99	25	
Propionibacterium granulosum		18.11	0.37		99	90	
Fusobacterium mortiferum		17.78	0.40		1	5	
07611023	**Very good identification**						**07611023**
Actinomyces naeslundii		100.0	0.60	MLT 94			
07611031	**Good identification**						**07611031**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07611033	**Good identification**						**07611033**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07611040	**Good identification**						**07611040**
					GRA	CAT	
Fusobacterium mortiferum		94.63	0.57	SOR 1	1	5	
Actinomyces naeslundii		3.64	0.44		99	25	
Propionibacterium granulosum		1.73	0.24		99	90	
07611041	**Low discrimination**						**07611041**
					GRA		
Actinomyces naeslundii		85.12	0.47	MLT 94	99		
Fusobacterium mortiferum		14.88	0.22		1		
07611042	**Low discrimination**						**07611042**
					GRA	CAT	
Fusobacterium mortiferum		72.06	0.40	SOR 1	1	5	
Actinomyces naeslundii		24.98	0.44	BGL 10	99	25	
Propionibacterium granulosum		2.97	0.13		99	90	
07611043	**Good identification**						**07611043**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07611060	**Low discrimination**						**07611060**
					GRA		
Actinomyces naeslundii		61.09	0.44	MLT 94	99		
Fusobacterium mortiferum		16.02	0.22		1		
Clostridium perfringens		15.63	0.12		99		
07611061	**Good identification**						**07611061**
Actinomyces naeslundii		100.0	0.47	MLT 94			
07611062	**Low discrimination**						**07611062**
Actinomyces naeslundii		89.25	0.44	MLT 94			
Clostridium perfringens		10.75	0.06				
07611063	**Good identification**						**07611063**
Actinomyces naeslundii		100.0	0.47	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07611100	**Excellent identification**					**07611100**
					GRA	
Fusobacterium mortiferum		99.56	0.97		1	
Actinomyces naeslundii		0.27	0.65		99	
Lactobacillus fermentum		0.14	0.50		99	
07611101	**Low discrimination**					**07611101**
					GRA	
Fusobacterium mortiferum		71.40	0.63	URE 1	1	
Actinomyces naeslundii		28.51	0.68		99	
Lactobacillus fermentum		0.10	0.15		99	
07611102	**Good identification**					**07611102**
					GRA	
Fusobacterium mortiferum		94.98	0.81	BGL 10	1	
Actinomyces naeslundii		2.30	0.65		99	
Lactobacillus fermentum		1.17	0.50		99	
07611103	**Low discrimination**					**07611103**
					GRA	
Actinomyces naeslundii		77.21	0.68	MLT 94	99	
Fusobacterium mortiferum		21.49	0.47		1	
Actinomyces viscosus		0.86	0.10		99	
07611110	**Very good identification**					**07611110**
					GRA	
Fusobacterium mortiferum		98.38	0.63	MLZ 1	1	
Actinomyces naeslundii		1.38	0.43		99	
Lactobacillus fermentum		0.14	0.15		99	
07611111	**Low discrimination**					**07611111**
					GRA	
Actinomyces naeslundii		67.54	0.46	MLT 94	99	
Fusobacterium mortiferum		32.46	0.29	MLZ 5	1	
07611112	**Low discrimination**					**07611112**
					GRA	
Fusobacterium mortiferum		82.29	0.47	MLZ 1	1	
Actinomyces naeslundii		10.38	0.43	BGL 10	99	
Bifidobacterium adolescentis		4.94	0.22		99	
07611113	**Acceptable identification**					**07611113**
					GRA	
Actinomyces naeslundii		94.93	0.46	MLT 94	99	
Fusobacterium mortiferum		5.07	0.12	MLZ 5	1	
07611120	**Low discrimination**					**07611120**
					GRA	
Fusobacterium mortiferum		59.37	0.63	NAG 1	1	
Actinomyces naeslundii		15.80	0.65		99	
Clostridium perfringens		9.65	0.39		99	
07611121	**Very good identification**					**07611121**
					GRA	
Actinomyces naeslundii		96.76	0.68	MLT 94	99	
Fusobacterium mortiferum		2.45	0.29		1	
Clostridium perfringens		0.40	0.04		99	
07611122	**Low discrimination**					**07611122**
					GRA	
Prevotella oralis		35.55	0.44	MLT 99	1	
Actinomyces naeslundii		19.01	0.65		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		12.57	0.49		1	
07611123	**Very good identification**					**07611123**
					GRA	
Actinomyces naeslundii		97.66	0.68	MLT 94	99	
Prevotella oralis		1.23	0.09		1	
Parabacteroides distasonis		0.44	0.14		1	
07611130	**Low discrimination**					**07611130**
					GRA	
Actinomyces naeslundii		49.43	0.43	MLT 94	99	
Fusobacterium mortiferum		35.64	0.29	MLZ 5	1	
Clostridium perfringens		5.79	0.04		99	
07611131	**Good identification**					**07611131**
Actinomyces naeslundii		100.0	0.46	MLT 94		
07611132	**Low discrimination**					**07611132**
					GRA	
Parabacteroides distasonis		81.31	0.44	XYL 90	1	
Actinomyces naeslundii		12.02	0.43		99	
Prevotella oralis		4.31	0.09		1	
07611133	**Good identification**					**07611133**
					GRA	
Actinomyces naeslundii		95.64	0.46	MLT 94	99	
Parabacteroides distasonis		4.36	0.10	MLZ 5	1	
07611140	**Very good identification**					**07611140**
					GRA	
Fusobacterium mortiferum		95.30	0.63	SOR 1	1	
Actinomyces naeslundii		4.48	0.52		99	
Lactobacillus fermentum		0.13	0.15		99	
07611141	**Low discrimination**					**07611141**
					GRA	
Actinomyces naeslundii		87.46	0.55	MLT 94	99	
Fusobacterium mortiferum		12.54	0.29		1	
07611142	**Low discrimination**					**07611142**
					GRA	
Fusobacterium mortiferum		67.68	0.47	SOR 1	1	
Actinomyces naeslundii		28.62	0.52	BGL 10	99	
Lactobacillus acidophilus		2.31	0.09		99	
07611143	**Good identification**					**07611143**
					GRA	
Actinomyces naeslundii		98.43	0.55	MLT 94	99	
Fusobacterium mortiferum		1.57	0.12		1	
07611160	**Low discrimination**					**07611160**
					GRA	
Actinomyces naeslundii		52.38	0.52	MLT 94	99	
Clostridium perfringens		34.53	0.26		99	
Fusobacterium mortiferum		11.26	0.29		1	
07611161	**Good identification**					**07611161**
Actinomyces naeslundii		100.0	0.55	MLT 94		
07611162	**Low discrimination**					**07611162**
					GRA	
Actinomyces naeslundii		62.72	0.52	MLT 94	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		19.46	0.21		99		
Prevotella oralis		6.71	0.09		1		
07611163	**Good identification**						**07611163**
Actinomyces naeslundii		100.0	0.55	MLT 94			
07611200	**Good identification**						**07611200**
					GRA	CAT	
Fusobacterium mortiferum		94.50	0.57	RHA 1	1	5	
Propionibacterium granulosum		5.29	0.32		99	90	
Actinomyces naeslundii		0.21	0.23		99	25	
07611202	**Low discrimination**						**07611202**
					GRA	CAT	
Fusobacterium mortiferum		87.27	0.40	RHA 1	1	5	
Propionibacterium granulosum		11.00	0.22	BGL 10	99	90	
Actinomyces naeslundii		1.73	0.23		99	25	
07611300	**Very good identification**						**07611300**
					GRA		
Fusobacterium mortiferum		99.60	0.63	RHA 1	1		
Actinomyces naeslundii		0.27	0.30		99		
Lactobacillus fermentum		0.14	0.15		99		
07611302	**Low discrimination**						**07611302**
Fusobacterium mortiferum		82.33	0.47	RHA 1			
Prevotella oralis		8.32	0.15	BGL 10			
Parabacteroides distasonis		3.95	0.23				
07611322	**Low discrimination**						**07611322**
					GRA		
Parabacteroides distasonis		70.49	0.57	XYL 90	1		
Prevotella oralis		28.48	0.37		1		
Actinomyces naeslundii		0.36	0.30		99		
07611332	**Good identification**						**07611332**
Parabacteroides distasonis		99.25	0.52	XYL 90			
Prevotella oralis		0.75	0.03				
07611400	**Low discrimination**						**07611400**
					GRA	CAT	
Fusobacterium mortiferum		82.34	0.57	ARA 1	1	5	
Lactobacillus fermentum		15.97	0.46		99	1	
Propionibacterium granulosum		1.51	0.24		99	90	
07611402	**Low discrimination**						**07611402**
					GRA	CAT	
Lactobacillus fermentum		62.20	0.46	SAL 1	99	1	
Fusobacterium mortiferum		35.63	0.40		1	5	
Propionibacterium granulosum		1.47	0.13		99	90	
07611420	**Low discrimination**						**07611420**
					GRA	CAT	
Lactobacillus fermentum		79.30	0.38	SAL 1	99	1	
Fusobacterium mortiferum		12.39	0.22		1	5	
Propionibacterium granulosum		5.61	0.13		99	90	
07611422	**Acceptable identification**						**07611422**
					GRA		
Lactobacillus fermentum		93.16	0.38	SAL 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		3.66	0.11		1		
Actinomyces naeslundii		3.18	0.23		99		
07611500	**Low discrimination**						**07611500**
					GRA		
Fusobacterium mortiferum		87.73	0.63	ARA 1	1		
Lactobacillus fermentum		11.90	0.50		99		
Actinomyces naeslundii		0.24	0.30		99		
07611502	**Low discrimination**						**07611502**
					GRA		
Lactobacillus fermentum		46.86	0.50	SAL 1	99		
Fusobacterium mortiferum		38.38	0.47		1		
Bifidobacterium adolescentis		12.60	0.34		99		
07611512	**Low discrimination**						**07611512**
Bifidobacterium adolescentis		89.65	0.36	MLT 99			
Bifidobacterium longum subsp. longum		5.34	0.13	XYL 87			
Lactobacillus fermentum		2.76	0.15				
07611520	**Low discrimination**						**07611520**
					GRA		
Lactobacillus fermentum		61.25	0.41	SAL 1	99		
Fusobacterium mortiferum		13.68	0.29		1		
Bifidobacterium longum subsp. longum		8.45	0.20		99		
07611522	**Low discrimination**						**07611522**
					GRA		
Parabacteroides distasonis		63.54	0.45	XYL 90	1		
Lactobacillus fermentum		26.63	0.41		99		
Bifidobacterium longum subsp. longum		3.67	0.20		99		
07611532	**Low discrimination**						**07611532**
					GRA		
Parabacteroides distasonis		84.54	0.40	XYL 90	1		
Bifidobacterium longum subsp. longum		15.46	0.24		99		
07611722	**Good identification**						**07611722**
Parabacteroides distasonis		99.03	0.53	XYL 90			
Prevotella oralis		0.66	0.03				
Clostridium clostridioforme		0.31	0.05				
07611732	**Good identification**						**07611732**
Parabacteroides distasonis		97.40	0.48	XYL 90			
Clostridium clostridioforme		2.60	0.16				
07612000	**Low discrimination**						**07612000**
					GRA	CAT	
Propionibacterium granulosum		66.04	0.56	SAL 1	99	90	
Fusobacterium mortiferum		32.22	0.54		1	5	
Actinomyces naeslundii		0.85	0.38		99	25	
07612001	**Low discrimination**						**07612001**
					GRA	CAT	
Actinomyces naeslundii		56.32	0.41	MLT 94	99	25	
Propionibacterium granulosum		29.36	0.21	GAL 96	99	90	
Fusobacterium mortiferum		14.33	0.19		1	5	
07612002	**Low discrimination**						**07612002**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		77.05	0.45	SAL 1	99	90		
Fusobacterium mortiferum		16.71	0.37		1	5		
Actinomyces naeslundii		3.98	0.38		99	25		
07612003	**Low discrimination**							**07612003**
Actinomyces naeslundii		88.47	0.41	MLT 94				
Propionibacterium granulosum		11.53	0.11	GAL 96				
07612010	**Low discrimination**							**07612010**
					GRA	SPO	CAT	
Propionibacterium granulosum		84.29	0.41	SAL 1	99	1	90	
Clostridium difficile		12.67	0.14	MLZ 12	99	99	1	
Fusobacterium mortiferum		3.05	0.19		1	1	5	
07612020	**Acceptable identification**							**07612020**
					GRA	CAT		
Propionibacterium granulosum		93.34	0.45	SAL 1	99	90		
Actinomyces naeslundii		4.82	0.38		99	25		
Fusobacterium mortiferum		1.84	0.19		1	5		
07612021	**Low discrimination**							**07612021**
Actinomyces naeslundii		88.47	0.41	MLT 94				
Propionibacterium granulosum		11.53	0.11	GAL 96				
07612022	**Low discrimination**							**07612022**
Propionibacterium granulosum		82.87	0.35	SAL 1				
Actinomyces naeslundii		17.13	0.38					
07612023	**Good identification**							**07612023**
Actinomyces naeslundii		100.0	0.41	MLT 94				
07612100	**Very good identification**							**07612100**
					GRA	CAT		
Fusobacterium mortiferum		96.05	0.60	MAN 1	1	5		
Actinomyces naeslundii		3.10	0.46		99	25		
Propionibacterium granulosum		0.85	0.21		99	90		
07612101	**Low discrimination**							**07612101**
					GRA			
Actinomyces naeslundii		82.75	0.49	MLT 94	99			
Fusobacterium mortiferum		17.25	0.26	GAL 96	1			
07612102	**Low discrimination**							**07612102**
					GRA	CAT		
Fusobacterium mortiferum		75.59	0.44	MAN 1	1	5		
Actinomyces naeslundii		21.97	0.46	BGL 10	99	25		
Propionibacterium granulosum		1.51	0.11		99	90		
07612103	**Good identification**							**07612103**
Actinomyces naeslundii		100.0	0.49	MLT 94				
07612120	**Low discrimination**							**07612120**
					GRA	CAT		
Actinomyces naeslundii		72.39	0.46	MLT 94	99	25		
Fusobacterium mortiferum		22.64	0.26	GAL 96	1	5		
Propionibacterium granulosum		4.97	0.11		99	90		
07612121	**Good identification**							**07612121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.49	MLT 94			
07612122	**Good identification**						**07612122**
Actinomyces naeslundii		100.0	0.46	MLT 94			
07612123	**Good identification**						**07612123**
Actinomyces naeslundii		100.0	0.49	MLT 94			
07612141	**Good identification**						**07612141**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07612143	**Good identification**						**07612143**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07612161	**Good identification**						**07612161**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07612163	**Good identification**						**07612163**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07613000	**Low discrimination**						**07613000**
					GRA	CAT	
Propionibacterium granulosum		45.84	0.56	SAL 1	99	90	
Fusobacterium mortiferum		33.55	0.57		1	5	
Actinomyces naeslundii		14.21	0.62		99	25	
07613001	**Very good identification**						**07613001**
					GRA	CAT	
Actinomyces naeslundii		96.38	0.65	MLT 94	99	25	
Propionibacterium granulosum		2.09	0.21		99	90	
Fusobacterium mortiferum		1.53	0.22		1	5	
07613002	**Low discrimination**						**07613002**
					SPO	CAT	
Actinomyces naeslundii		39.70	0.62	MLT 94	1	25	
Propionibacterium granulosum		32.01	0.45		1	90	
Clostridium innocuum		17.87	0.32		99	1	
07613003	**Very good identification**						**07613003**
Actinomyces naeslundii		99.46	0.65	MLT 94			
Propionibacterium granulosum		0.54	0.11				
07613010	**Low discrimination**						**07613010**
					GRA	CAT	
Propionibacterium granulosum		82.20	0.41	SAL 1	99	90	
Actinomyces naeslundii		9.84	0.40	MLZ 12	99	25	
Fusobacterium mortiferum		4.46	0.22		1	5	
07613011	**Good identification**						**07613011**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07613013	**Good identification**						**07613013**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07613020	**Low discrimination**						**07613020**
					GRA	CAT	
Actinomyces naeslundii		54.64	0.62	MLT 94	99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		44.06	0.45		99	90	
Fusobacterium mortiferum		1.30	0.22		1	5	
07613021	**Very good identification**						**07613021**
Actinomyces naeslundii		99.46	0.65	MLT 94			
Propionibacterium granulosum		0.54	0.11				
07613022	**Low discrimination**						**07613022**
Actinomyces naeslundii		83.22	0.62	MLT 94			
Propionibacterium granulosum		16.78	0.35				
07613023	**Very good identification**						**07613023**
Actinomyces naeslundii		100.0	0.65	MLT 94			
07613031	**Good identification**						**07613031**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07613032	**Low discrimination**						**07613032**
Actinomyces naeslundii		65.69	0.40	MLT 94			
Propionibacterium granulosum		34.31	0.20	MLZ 5			
07613033	**Good identification**						**07613033**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07613040	**Low discrimination**						**07613040**
					GRA	CAT	
Actinomyces naeslundii		75.77	0.49	MLT 94	99	25	
Propionibacterium granulosum		13.99	0.21		99	90	
Fusobacterium mortiferum		10.24	0.22		1	5	
07613041	**Good identification**						**07613041**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07613042	**Good identification**						**07613042**
Actinomyces naeslundii		95.59	0.49	MLT 94			
Propionibacterium granulosum		4.41	0.11				
07613043	**Good identification**						**07613043**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07613060	**Good identification**						**07613060**
Actinomyces naeslundii		95.59	0.49	MLT 94			
Propionibacterium granulosum		4.41	0.11				
07613061	**Good identification**						**07613061**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07613062	**Good identification**						**07613062**
Actinomyces naeslundii		100.0	0.49	MLT 94			
07613063	**Good identification**						**07613063**
Actinomyces naeslundii		100.0	0.52	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07613100	**Low discrimination**						**07613100**
					GRA	CAT	
Fusobacterium mortiferum		65.62	0.63	MAN 1	1	5	
Actinomyces naeslundii		33.91	0.70		99	25	
Propionibacterium granulosum		0.39	0.21		99	90	
07613101	**Very good identification**						**07613101**
					GRA		
Actinomyces naeslundii		98.71	0.73	MLT 94	99		
Fusobacterium mortiferum		1.29	0.29		1		
07613102	**Low discrimination**						**07613102**
					GRA		
Actinomyces naeslundii		80.79	0.70	MLT 94	99		
Fusobacterium mortiferum		17.37	0.47		1		
Bifidobacterium adolescentis		0.70	0.19		99		
07613103	**Very good identification**						**07613103**
					GRA		
Actinomyces naeslundii		99.86	0.73	MLT 94	99		
Fusobacterium mortiferum		0.15	0.12		1		
07613110	**Low discrimination**						**07613110**
					GRA		
Actinomyces naeslundii		72.92	0.48	MLT 94	99		
Fusobacterium mortiferum		27.08	0.29	MLZ 5	1		
07613111	**Good identification**						**07613111**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07613112	**Low discrimination**						**07613112**
					GRA		
Actinomyces naeslundii		80.53	0.48	MLT 94	99		
Bifidobacterium adolescentis		16.14	0.20	MLZ 5	99		
Fusobacterium mortiferum		3.32	0.12		1		
07613113	**Good identification**						**07613113**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07613120	**Very good identification**						**07613120**
					GRA		
Actinomyces naeslundii		97.21	0.70	MLT 94	99		
Fusobacterium mortiferum		1.90	0.29		1		
Clostridium perfringens		0.31	0.04		99		
07613121	**Very good identification**						**07613121**
Actinomyces naeslundii		100.0	0.73	MLT 94			
07613122	**Very good identification**						**07613122**
					SPO	GRA	
Actinomyces naeslundii		97.43	0.70	MLT 94	1	99	
Prevotella oralis		0.95	0.09		1	1	
Clostridium ramosum		0.47	0.05		92	50	
07613123	**Very good identification**						**07613123**
Actinomyces naeslundii		100.0	0.73	MLT 94			
07613130	**Good identification**						**07613130**
Actinomyces naeslundii		100.0	0.48	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07613131	**Good identification**					**07613131**
Actinomyces naeslundii		100.0	0.51	MLT 94		
07613132	**Good identification**					**07613132**
					GRA	
Actinomyces naeslundii		96.60	0.48	MLT 94	99	
Parabacteroides distasonis		3.40	0.10	MLZ 5	1	
07613133	**Good identification**					**07613133**
Actinomyces naeslundii		100.0	0.51	MLT 94		
07613140	**Good identification**					**07613140**
					GRA	
Actinomyces naeslundii		90.03	0.57	MLT 94	99	
Fusobacterium mortiferum		9.97	0.29		1	
07613141	**Very good identification**					**07613141**
Actinomyces naeslundii		100.0	0.60	MLT 94		
07613142	**Very good identification**					**07613142**
					GRA	
Actinomyces naeslundii		97.82	0.57	MLT 94	99	
Fusobacterium mortiferum		1.20	0.12		1	
Propionibacterium propionicum		0.98	0.12		99	
07613143	**Very good identification**					**07613143**
Actinomyces naeslundii		100.0	0.60	MLT 94		
07613151	**Good identification**					**07613151**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07613153	**Good identification**					**07613153**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07613160	**Very good identification**					**07613160**
Actinomyces naeslundii		100.0	0.57	MLT 94		
07613161	**Very good identification**					**07613161**
Actinomyces naeslundii		100.0	0.60	MLT 94		
07613162	**Very good identification**					**07613162**
Actinomyces naeslundii		100.0	0.57	MLT 94		
07613163	**Very good identification**					**07613163**
Actinomyces naeslundii		100.0	0.60	MLT 94		
07613171	**Good identification**					**07613171**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07613173	**Good identification**					**07613173**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07613301	**Good identification**					**07613301**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		100.0	0.38	MLT 94				
07613302	**Low discrimination**							**07613302**
					GRA			
Actinomyces naeslundii		82.30	0.35	MLT 94	99			
Fusobacterium mortiferum		17.70	0.12	RHA 1	1			
07613303	**Good identification**							**07613303**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07613320	**Good identification**							**07613320**
Actinomyces naeslundii		100.0	0.35	MLT 94				
07613321	**Good identification**							**07613321**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07613323	**Good identification**							**07613323**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07613501	**Good identification**							**07613501**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07613503	**Good identification**							**07613503**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07613520	**Good identification**							**07613520**
Actinomyces naeslundii		100.0	0.35	MLT 94				
07613521	**Good identification**							**07613521**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07613522	**Low discrimination**							**07613522**
					GRA			
Actinomyces naeslundii		82.73	0.35	MLT 94	99			
Parabacteroides distasonis		17.27	0.11	ARA 1	1			
07613523	**Good identification**							**07613523**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07614000	**Very good identification**							**07614000**
					COC	GRA	CAT	
Fusobacterium mortiferum		99.58	0.68	XYL 7	1	1	5	
Propionibacterium granulosum		0.37	0.24		1	99	90	
Gemella morbillorum		0.06	0.03		99	99	1	
07614002	**Good identification**							**07614002**
					GRA	CAT		
Fusobacterium mortiferum		99.18	0.52	XYL 7	1	5		
Propionibacterium granulosum		0.82	0.13	BGL 10	99	90		
07614100	**Very good identification**							**07614100**
Fusobacterium mortiferum		100.0	0.75	XYL 7				
07614101	**Good identification**							**07614101**
Fusobacterium mortiferum		100.0	0.41	XYL 7				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07614102	**Very good identification**						**07614102**
Fusobacterium mortiferum		100.0	0.58	XYL 7			
07614110	**Good identification**						**07614110**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
07614120	**Good identification**						**07614120**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
07614140	**Good identification**						**07614140**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
07614300	**Good identification**						**07614300**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
07614322	**Good identification**						**07614322**
Parabacteroides distasonis		100.0	0.39	GAL 99			
07614500	**Good identification**						**07614500**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
07614722	**Low discrimination**						**07614722**
Parabacteroides distasonis		89.51	0.35	GAL 99			
Clostridium clostridioforme		10.49	0.14				
07615000	**Very good identification**						**07615000**
					GRA	CAT	
Fusobacterium mortiferum		98.23	0.72	XYL 7	1	5	
Lactobacillus fermentum		1.50	0.42		99	1	
Propionibacterium granulosum		0.24	0.24		99	90	
07615001	**Good identification**						**07615001**
					GRA		
Fusobacterium mortiferum		95.79	0.37	XYL 7	1		
Actinomyces naeslundii		4.21	0.26	URE 1	99		
07615002	**Low discrimination**						**07615002**
					GRA	CAT	
Fusobacterium mortiferum		87.28	0.55	XYL 7	1	5	
Lactobacillus fermentum		12.01	0.42	BGL 10	99	1	
Propionibacterium granulosum		0.48	0.13		99	90	
07615010	**Good identification**						**07615010**
					GRA	CAT	
Fusobacterium mortiferum		96.79	0.37	XYL 7	1	5	
Propionibacterium granulosum		3.21	0.09	MLZ 1	99	90	
07615020	**Low discrimination**						**07615020**
					GRA	CAT	
Fusobacterium mortiferum		62.71	0.37	XYL 7	1	5	
Lactobacillus fermentum		31.64	0.34	NAG 1	99	1	
Propionibacterium granulosum		3.81	0.13		99	90	
07615040	**Good identification**						**07615040**
Fusobacterium mortiferum		100.0	0.37	XYL 7			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07615100	**Very good identification**					**07615100**
					GRA	
Fusobacterium mortiferum		98.79	0.78	XYL 7	1	
Lactobacillus fermentum		1.06	0.46		99	
Bifidobacterium longum subsp. infantis		0.10	0.09		99	
07615101	**Acceptable identification**					**07615101**
					GRA	
Fusobacterium mortiferum		94.91	0.44	XYL 7	1	
Actinomyces naeslundii		5.09	0.33	URE 1	99	
07615102	**Low discrimination**					**07615102**
					GRA	
Fusobacterium mortiferum		79.46	0.61	XYL 7	1	
Lactobacillus fermentum		7.65	0.46	BGL 10	99	
Bifidobacterium longum subsp. infantis		7.09	0.26		99	
07615110	**Good identification**					**07615110**
Fusobacterium mortiferum		100.0	0.44	XYL 7		
07615112	**Low discrimination**					**07615112**
					GRA	
Bifidobacterium adolescentis		72.42	0.36	MLT 99	99	
Parabacteroides distasonis		13.99	0.26	ARA 87	1	
Fusobacterium mortiferum		13.58	0.27		1	
07615120	**Low discrimination**					**07615120**
					GRA	
Fusobacterium mortiferum		47.27	0.44	XYL 7	1	
Lactobacillus fermentum		16.68	0.37	NAG 1	99	
Bacteroides vulgatus		12.06	0.14		1	
07615122	**Low discrimination**					**07615122**
					GRA	
Parabacteroides distasonis		87.13	0.65		1	
Bifidobacterium longum subsp. infantis		7.23	0.30		99	
Prevotella oralis		3.38	0.28		1	
07615132	**Very good identification**					**07615132**
					GRA	
Parabacteroides distasonis		99.85	0.60		1	
Bifidobacterium longum subsp. infantis		0.16	-.04		99	
07615140	**Good identification**					**07615140**
Fusobacterium mortiferum		100.0	0.44	XYL 7		
07615200	**Good identification**					**07615200**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
07615222	**Good identification**					**07615222**
Parabacteroides distasonis		96.61	0.39	LAC 99		
Prevotella oralis		3.39	0.01			
07615300	**Low discrimination**					**07615300**
					GRA	
Fusobacterium mortiferum		70.56	0.44	XYL 7	1	
Eubacterium contortum		27.80	0.24	RHA 1	99	
Bacteroides vulgatus		1.64	-.04		1	
07615302	**Low discrimination**					**07615302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		76.21	0.39	NAG 99	1	
Fusobacterium mortiferum		13.28	0.27		1	
Eubacterium contortum		8.31	0.11		99	
07615322	**Very good identification**					**07615322**
Parabacteroides distasonis		99.37	0.73			
Prevotella oralis		0.55	0.22			
Bacteroides vulgatus		0.04	-.04			
07615323	**Good identification**					**07615323**
Parabacteroides distasonis		100.0	0.39	URE 1		
07615332	**Very good identification**					**07615332**
Parabacteroides distasonis		99.88	0.68			
Clostridium clostridioforme		0.12	0.13			
07615362	**Good identification**					**07615362**
Parabacteroides distasonis		100.0	0.39	SOR 1		
07615400	**Low discrimination**					**07615400**
					GRA	
Lactobacillus fermentum		60.22	0.42	SAL 1	99	
Fusobacterium mortiferum		39.78	0.37		1	
07615402	**Low discrimination**					**07615402**
					GRA	
Lactobacillus fermentum		88.98	0.42	SAL 1	99	
Fusobacterium mortiferum		6.53	0.21		1	
Bifidobacterium adolescentis		4.49	0.14		99	
07615500	**Low discrimination**					**07615500**
					GRA	
Lactobacillus fermentum		47.59	0.46	SAL 1	99	
Fusobacterium mortiferum		44.96	0.44		1	
Bifidobacterium longum subsp. longum		2.80	0.17		99	
07615502	**Low discrimination**					**07615502**
					GRA	
Bifidobacterium adolescentis		69.80	0.48	MLT 99	99	
Lactobacillus fermentum		22.78	0.46		99	
Parabacteroides distasonis		2.80	0.27		1	
07615512	**Acceptable identification**					**07615512**
Bifidobacterium adolescentis		93.82	0.50	MLT 99		
Bifidobacterium longum subsp. longum		2.50	0.21			
Clostridium clostridioforme		2.02	0.18			
07615522	**Good identification**					**07615522**
					GRA	
Parabacteroides distasonis		94.82	0.61		1	
Lactobacillus fermentum		2.59	0.37		99	
Bifidobacterium longum subsp. longum		1.83	0.28		99	
07615532	**Good identification**					**07615532**
					GRA	
Parabacteroides distasonis		92.68	0.57		1	
Bifidobacterium longum subsp. longum		5.65	0.32		99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium clostridioforme		1.14	0.18		50			
07615622	**Acceptable identification**							**07615622**
Parabacteroides distasonis		90.50	0.35	LAC 99				
Clostridium clostridioforme		9.50	0.13					
07615702	**Low discrimination**							**07615702**
					GRA			
Parabacteroides distasonis		63.26	0.35	NAG 99	1			
Clostridium clostridioforme		22.23	0.22		50			
Eubacterium contortum		9.21	0.10		99			
07615720	**Good identification**							**07615720**
					GRA			
Bacteroides vulgatus		98.55	0.49	MLT 99	1			
Parabacteroides distasonis		1.26	0.35		1			
Eubacterium contortum		0.18	0.10		99			
07615722	**Very good identification**							**07615722**
Parabacteroides distasonis		98.86	0.69					
Bacteroides vulgatus		0.79	0.15					
Clostridium clostridioforme		0.35	0.22					
07615723	**Good identification**							**07615723**
Parabacteroides distasonis		100.0	0.35	URE 1				
07615732	**Very good identification**							**07615732**
Parabacteroides distasonis		97.08	0.65					
Clostridium clostridioforme		2.92	0.33					
07615762	**Good identification**							**07615762**
Parabacteroides distasonis		100.0	0.35	SOR 1				
07616100	**Good identification**							**07616100**
Fusobacterium mortiferum		100.0	0.41	MAN 1				
07617000	**Low discrimination**							**07617000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		69.73	0.37	MAN 1	1	1	5	
Propionibacterium granulosum		12.79	0.21	XYL 7	99	1	90	
Clostridium innocuum		9.30	0.10		99	99	1	
07617100	**Low discrimination**							**07617100**
					GRA			
Fusobacterium mortiferum		85.23	0.44	MAN 1	1			
Actinomyces israelii		8.86	0.29	XYL 7	99			
Actinomyces naeslundii		5.91	0.35		99			
07617101	**Good identification**							**07617101**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07617103	**Good identification**							**07617103**
Actinomyces naeslundii		100.0	0.38	MLT 94				
07617120	**Good identification**							**07617120**
Actinomyces naeslundii		100.0	0.35	MLT 94				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07617121	**Good identification**						**07617121**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07617123	**Good identification**						**07617123**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07617322	**Good identification**						**07617322**
Parabacteroides distasonis		100.0	0.39	MAN 1			
07617502	**Good identification**						**07617502**
Bifidobacterium adolescentis		95.42	0.47	MLT 99			
Actinomyces israelii		4.59	0.31				
07617512	**Good identification**						**07617512**
Bifidobacterium adolescentis		99.42	0.48	MLT 99			
Actinomyces israelii		0.58	0.17				
07617722	**Good identification**						**07617722**
Parabacteroides distasonis		100.0	0.35	MAN 1			
07620000	**Low discrimination**						**07620000**
					GRA	CAT	
Fusobacterium mortiferum		78.39	0.45	FRU 99	1	5	
Propionibacterium granulosum		21.61	0.32		99	90	
07620100	**Good identification**						**07620100**
					GRA		
Fusobacterium mortiferum		99.43	0.51	FRU 99	1		
Actinomyces naeslundii		0.57	0.24		99		
07621000	**Low discrimination**						**07621000**
					GRA	CAT	
Fusobacterium mortiferum		79.47	0.48	FRU 99	1	5	
Propionibacterium granulosum		14.60	0.32		99	90	
Actinomyces naeslundii		5.93	0.40		99	25	
07621001	**Acceptable identification**						**07621001**
					GRA		
Actinomyces naeslundii		91.73	0.43	MLT 94	99		
Fusobacterium mortiferum		8.27	0.13	FRU 95	1		
07621003	**Good identification**						**07621003**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07621020	**Low discrimination**						**07621020**
					GRA	CAT	
Actinomyces naeslundii		57.12	0.40	MLT 94	99	25	
Propionibacterium granulosum		35.15	0.21	FRU 95	99	90	
Fusobacterium mortiferum		7.73	0.13		1	5	
07621021	**Good identification**						**07621021**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07621022	**Low discrimination**						**07621022**
Actinomyces naeslundii		86.67	0.40	MLT 94			
Propionibacterium granulosum		13.33	0.11	FRU 95			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07621023	**Good identification**						**07621023**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07621100	**Acceptable identification**						**07621100**
					GRA		
Fusobacterium mortiferum		91.66	0.54	FRU 99	1		
Actinomyces naeslundii		8.35	0.48		99		
07621101	**Acceptable identification**						**07621101**
					GRA		
Actinomyces naeslundii		93.11	0.51	MLT 94	99		
Fusobacterium mortiferum		6.89	0.20	FRU 95	1		
07621102	**Low discrimination**						**07621102**
					GRA		
Fusobacterium mortiferum		54.03	0.38	FRU 99	1		
Actinomyces naeslundii		44.28	0.48	BGL 10	99		
Lactobacillus acidophilus		1.69	-.01		99		
07621103	**Good identification**						**07621103**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07621120	**Low discrimination**						**07621120**
					GRA		
Actinomyces naeslundii		81.76	0.48	MLT 94	99		
Clostridium perfringens		9.17	0.09	FRU 95	99		
Fusobacterium mortiferum		9.07	0.20		1		
07621121	**Good identification**						**07621121**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07621122	**Low discrimination**						**07621122**
					GRA		
Actinomyces naeslundii		72.01	0.48	MLT 94	99		
Parabacteroides distasonis		21.44	0.26	FRU 95	1		
Clostridium perfringens		3.80	0.03		99		
07621123	**Good identification**						**07621123**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07621141	**Good identification**						**07621141**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07621143	**Good identification**						**07621143**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07621161	**Good identification**						**07621161**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07621163	**Good identification**						**07621163**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07623000	**Low discrimination**						**07623000**
					SPO	CAT	
Actinomyces naeslundii		47.44	0.45	MLT 94	1	25	
Propionibacterium granulosum		45.39	0.30	FRU 95	1	90	
Clostridium innocuum		3.86	0.02		99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07623001	**Good identification**						**07623001**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07623002	**Low discrimination**						**07623002**
					SPO	CAT	
Actinomyces naeslundii		75.72	0.45	MLT 94	1	25	
Propionibacterium granulosum		18.11	0.19	FRU 95	1	90	
Clostridium innocuum		6.17	0.02		99	1	
07623003	**Good identification**						**07623003**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07623020	**Low discrimination**						**07623020**
Actinomyces naeslundii		80.70	0.45	MLT 94			
Propionibacterium granulosum		19.30	0.19	FRU 95			
07623021	**Good identification**						**07623021**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07623022	**Acceptable identification**						**07623022**
Actinomyces naeslundii		94.36	0.45	MLT 94			
Propionibacterium granulosum		5.64	0.09	FRU 95			
07623023	**Good identification**						**07623023**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07623100	**Acceptable identification**						**07623100**
					GRA		
Actinomyces naeslundii		94.59	0.53	MLT 94	99		
Fusobacterium mortiferum		5.41	0.20	FRU 95	1		
07623101	**Good identification**						**07623101**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07623102	**Good identification**						**07623102**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07623103	**Good identification**						**07623103**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07623120	**Good identification**						**07623120**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07623121	**Good identification**						**07623121**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07623122	**Good identification**						**07623122**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07623123	**Good identification**						**07623123**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07623140	**Good identification**						**07623140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.40	MLT 94			
07623141	**Good identification**						**07623141**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07623142	**Good identification**						**07623142**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07623143	**Good identification**						**07623143**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07623160	**Good identification**						**07623160**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07623161	**Good identification**						**07623161**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07623162	**Good identification**						**07623162**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07623163	**Good identification**						**07623163**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07625122	**Good identification**						**07625122**
Parabacteroides distasonis		100.0	0.42	FRU 99			
07625132	**Good identification**						**07625132**
Parabacteroides distasonis		100.0	0.37	FRU 99			
07625322	**Good identification**						**07625322**
Parabacteroides distasonis		100.0	0.50	FRU 99			
07625332	**Good identification**						**07625332**
Parabacteroides distasonis		100.0	0.46	FRU 99			
07625522	**Good identification**						**07625522**
Parabacteroides distasonis		100.0	0.38	FRU 99			
07625722	**Good identification**						**07625722**
Parabacteroides distasonis		100.0	0.47	FRU 99			
07625732	**Good identification**						**07625732**
Parabacteroides distasonis		95.39	0.42	FRU 99			
Clostridium clostridioforme		4.61	0.14				
07630000	**Low discrimination**						**07630000**
					GRA	CAT	
Fusobacterium mortiferum		78.34	0.79		1	5	
Propionibacterium granulosum		21.59	0.66		99	90	
Actinomyces naeslundii		0.07	0.38		99	25	
07630001	**Low discrimination**						**07630001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Fusobacterium mortiferum		70.99	0.45	URE 1	1	5	
Propionibacterium granulosum		19.57	0.32		99	90	
Actinomyces naeslundii		9.44	0.41		99	25	
07630002	**Low discrimination**						**07630002**
					GRA	CAT	
Fusobacterium mortiferum		61.38	0.62	BGL 10	1	5	
Propionibacterium granulosum		38.06	0.56		99	90	
Actinomyces naeslundii		0.50	0.38		99	25	
07630003	**Low discrimination**						**07630003**
					GRA	CAT	
Actinomyces naeslundii		42.48	0.41	MLT 94	99	25	
Fusobacterium mortiferum		35.50	0.28	GAL 96	1	5	
Propionibacterium granulosum		22.02	0.21		99	90	
07630010	**Low discrimination**						**07630010**
					GRA	CAT	
Propionibacterium granulosum		78.82	0.51	SAL 1	99	90	
Fusobacterium mortiferum		21.18	0.45	MLZ 12	1	5	
07630012	**Low discrimination**						**07630012**
					GRA	CAT	
Propionibacterium granulosum		89.33	0.41	SAL 1	99	90	
Fusobacterium mortiferum		10.67	0.28	MLZ 12	1	5	
07630020	**Low discrimination**						**07630020**
					GRA	CAT	
Propionibacterium granulosum		86.24	0.56	SAL 1	99	90	
Fusobacterium mortiferum		12.64	0.45		1	5	
Actinomyces naeslundii		1.12	0.38		99	25	
07630021	**Low discrimination**						**07630021**
Actinomyces naeslundii		65.87	0.41	MLT 94			
Propionibacterium granulosum		34.14	0.21	GAL 96			
07630022	**Low discrimination**						**07630022**
					GRA	CAT	
Propionibacterium granulosum		89.52	0.45	SAL 1	99	90	
Fusobacterium mortiferum		5.83	0.28		1	5	
Actinomyces naeslundii		4.65	0.38		99	25	
07630023	**Low discrimination**						**07630023**
Actinomyces naeslundii		88.53	0.41	MLT 94			
Propionibacterium granulosum		11.47	0.11	GAL 96			
07630030	**Good identification**						**07630030**
Propionibacterium granulosum		100.0	0.41	SAL 1			
07630040	**Low discrimination**						**07630040**
					GRA	CAT	
Fusobacterium mortiferum		76.54	0.45	SOR 1	1	5	
Propionibacterium granulosum		21.10	0.32		99	90	
Actinomyces naeslundii		1.20	0.25		99	25	
07630100	**Very good identification**						**07630100**
					GRA	CAT	
Fusobacterium mortiferum		99.77	0.85		1	5	
Propionibacterium granulosum		0.12	0.32		99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.11	0.46		99	25	
07630101	**Low discrimination**						**07630101**
					GRA		
Fusobacterium mortiferum		86.05	0.51	URE 1	1		
Actinomyces naeslundii		13.96	0.49		99		
07630102	**Very good identification**						**07630102**
					GRA	CAT	
Fusobacterium mortiferum		98.65	0.69	BGL 10	1	5	
Actinomyces naeslundii		0.97	0.46		99	25	
Propionibacterium granulosum		0.27	0.21		99	90	
07630103	**Low discrimination**						**07630103**
					GRA		
Actinomyces naeslundii		59.34	0.49	MLT 94	99		
Fusobacterium mortiferum		40.66	0.35	GAL 96	1		
07630110	**Good identification**						**07630110**
					GRA	CAT	
Fusobacterium mortiferum		97.87	0.51	MLZ 1	1	5	
Propionibacterium granulosum		1.58	0.17		99	90	
Actinomyces naeslundii		0.56	0.24		99	25	
07630120	**Low discrimination**						**07630120**
					GRA	CAT	
Fusobacterium mortiferum		86.73	0.51	NAG 1	1	5	
Actinomyces naeslundii		9.38	0.46		99	25	
Propionibacterium granulosum		2.56	0.21		99	90	
07630121	**Acceptable identification**						**07630121**
					GRA		
Actinomyces naeslundii		94.14	0.49	MLT 94	99		
Fusobacterium mortiferum		5.86	0.17	GAL 96	1		
07630123	**Good identification**						**07630123**
Actinomyces naeslundii		100.0	0.49	MLT 94			
07630140	**Good identification**						**07630140**
					GRA		
Fusobacterium mortiferum		98.13	0.51	SOR 1	1		
Actinomyces naeslundii		1.87	0.33		99		
07630141	**Low discrimination**						**07630141**
					GRA		
Actinomyces naeslundii		73.91	0.36	MLT 94	99		
Fusobacterium mortiferum		26.09	0.17	GAL 96	1		
07630143	**Good identification**						**07630143**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07630161	**Good identification**						**07630161**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07630163	**Good identification**						**07630163**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07630200	**Low discrimination**						**07630200**
					GRA	CAT	
Fusobacterium mortiferum		54.23	0.45	RHA 1	1	5	
Propionibacterium granulosum		45.77	0.40		99	90	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07630300	**Good identification**						**07630300**
Fusobacterium mortiferum		100.0	0.51	RHA 1			
07630400	**Low discrimination**						**07630400**
					GRA	CAT	
Fusobacterium mortiferum		78.39	0.45	ARA 1	1	5	
Propionibacterium granulosum		21.61	0.32		99	90	
07630500	**Good identification**						**07630500**
Fusobacterium mortiferum		100.0	0.51	ARA 1			
07631000	**Low discrimination**						**07631000**
					GRA	CAT	
Fusobacterium mortiferum		83.37	0.82		1	5	
Propionibacterium granulosum		15.32	0.66		99	90	
Actinomyces naeslundii		1.19	0.62		99	25	
07631001	**Low discrimination**						**07631001**
					GRA	CAT	
Actinomyces naeslundii		64.25	0.65	MLT 94	99	25	
Fusobacterium mortiferum		30.20	0.48		1	5	
Propionibacterium granulosum		5.55	0.32		99	90	
07631002	**Low discrimination**						**07631002**
					GRA	CAT	
Fusobacterium mortiferum		64.26	0.65	BGL 10	1	5	
Propionibacterium granulosum		26.57	0.56		99	90	
Actinomyces naeslundii		8.29	0.62		99	25	
07631003	**Good identification**						**07631003**
					GRA	CAT	
Actinomyces naeslundii		93.13	0.65	MLT 94	99	25	
Fusobacterium mortiferum		4.86	0.31		1	5	
Propionibacterium granulosum		2.01	0.21		99	90	
07631010	**Low discrimination**						**07631010**
					GRA	CAT	
Propionibacterium granulosum		68.97	0.51	SAL 1	99	90	
Fusobacterium mortiferum		27.81	0.48	MLZ 12	1	5	
Actinomyces naeslundii		2.08	0.40		99	25	
07631011	**Low discrimination**						**07631011**
					GRA	CAT	
Actinomyces naeslundii		76.11	0.43	MLT 94	99	25	
Propionibacterium granulosum		17.03	0.17	MLZ 5	99	90	
Fusobacterium mortiferum		6.87	0.13		1	5	
07631012	**Low discrimination**						**07631012**
					GRA	CAT	
Propionibacterium granulosum		74.41	0.41	SAL 1	99	90	
Fusobacterium mortiferum		13.33	0.31	MLZ 12	1	5	
Actinomyces naeslundii		8.96	0.40		99	25	
07631013	**Good identification**						**07631013**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07631020	**Low discrimination**						**07631020**
					GRA	CAT	
Propionibacterium granulosum		62.14	0.56	SAL 1	99	90	
Actinomyces naeslundii		19.38	0.62		99	25	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		13.66	0.48		1	5	
07631021	**Very good identification**						**07631021**
					GRA	CAT	
Actinomyces naeslundii		97.43	0.65	MLT 94	99	25	
Propionibacterium granulosum		2.10	0.21		99	90	
Fusobacterium mortiferum		0.46	0.13		1	5	
07631022	**Low discrimination**						**07631022**
					CAT		
Actinomyces naeslundii		47.48	0.62	MLT 94	25		
Propionibacterium granulosum		38.06	0.45		90		
Clostridium perfringens		5.07	0.23		1		
07631023	**Very good identification**						**07631023**
Actinomyces naeslundii		99.46	0.65	MLT 94			
Propionibacterium granulosum		0.54	0.11				
07631030	**Low discrimination**						**07631030**
					CAT		
Propionibacterium granulosum		86.71	0.41	SAL 1	90		
Actinomyces naeslundii		10.44	0.40	MLZ 12	25		
Lactobacillus jensenii		1.44	0.04		1		
07631031	**Good identification**						**07631031**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07631033	**Good identification**						**07631033**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07631040	**Low discrimination**						**07631040**
					GRA	CAT	
Fusobacterium mortiferum		69.73	0.48	SOR 1	1	5	
Actinomyces naeslundii		17.45	0.49		99	25	
Propionibacterium granulosum		12.81	0.32		99	90	
07631041	**Good identification**						**07631041**
					GRA		
Actinomyces naeslundii		97.38	0.52	MLT 94	99		
Fusobacterium mortiferum		2.62	0.13		1		
07631042	**Low discrimination**						**07631042**
					GRA	CAT	
Actinomyces naeslundii		60.17	0.49	MLT 94	99	25	
Fusobacterium mortiferum		26.71	0.31		1	5	
Propionibacterium granulosum		11.04	0.21		99	90	
07631043	**Good identification**						**07631043**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07631060	**Low discrimination**						**07631060**
					CAT		
Actinomyces naeslundii		68.09	0.49	MLT 94	25		
Clostridium perfringens		16.67	0.16		1		
Propionibacterium granulosum		12.50	0.21		90		
07631061	**Good identification**						**07631061**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07631062	**Low discrimination**						**07631062**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Actinomyces naeslundii		83.64	0.49	MLT 94	25		
Clostridium perfringens		9.64	0.11		1		
Propionibacterium granulosum		3.84	0.11		90		
07631063	**Good identification**						**07631063**
Actinomyces naeslundii		100.0	0.52	MLT 94			
07631100	**Very good identification**						**07631100**
					GRA	CAT	
Fusobacterium mortiferum		98.15	0.88		1	5	
Actinomyces naeslundii		1.72	0.70		99	25	
Propionibacterium granulosum		0.08	0.32		99	90	
07631101	**Low discrimination**						**07631101**
					GRA		
Actinomyces naeslundii		72.18	0.73	MLT 94	99		
Fusobacterium mortiferum		27.82	0.54		1		
07631102	**Low discrimination**						**07631102**
					GRA		
Fusobacterium mortiferum		83.54	0.72	BGL 10	1		
Actinomyces naeslundii		13.14	0.70		99		
Lactobacillus acidophilus		2.61	0.33		99		
07631103	**Very good identification**						**07631103**
					GRA		
Actinomyces naeslundii		95.77	0.73	MLT 94	99		
Fusobacterium mortiferum		4.10	0.38		1		
Lactobacillus acidophilus		0.13	-.01		99		
07631110	**Acceptable identification**						**07631110**
					GRA	CAT	
Fusobacterium mortiferum		90.76	0.54	MLZ 1	1	5	
Actinomyces naeslundii		8.26	0.48		99	25	
Propionibacterium granulosum		0.98	0.17		99	90	
07631111	**Acceptable identification**						**07631111**
					GRA		
Actinomyces naeslundii		93.11	0.51	MLT 94	99		
Fusobacterium mortiferum		6.89	0.20	MLZ 5	1		
07631112	**Low discrimination**						**07631112**
					GRA		
Fusobacterium mortiferum		46.55	0.38	MLZ 1	1		
Actinomyces naeslundii		38.15	0.48	BGL 10	99		
Bifidobacterium adolescentis		6.23	0.19		99		
07631113	**Good identification**						**07631113**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07631120	**Low discrimination**						**07631120**
					GRA		
Actinomyces naeslundii		45.33	0.70	MLT 94	99		
Clostridium perfringens		26.49	0.43		99		
Fusobacterium mortiferum		26.20	0.54		1		
07631121	**Very good identification**						**07631121**
					GRA		
Actinomyces naeslundii		99.22	0.73	MLT 94	99		
Clostridium perfringens		0.39	0.09		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		0.39	0.20		1	
07631122	**Low discrimination**					**07631122**
					GRA	
Parabacteroides distasonis		48.94	0.60	XYL 90	1	
Actinomyces naeslundii		31.54	0.70		99	
Clostridium perfringens		8.67	0.38		99	
07631123	**Very good identification**					**07631123**
					GRA	
Actinomyces naeslundii		98.65	0.73	MLT 94	99	
Parabacteroides distasonis		1.03	0.26		1	
Clostridium perfringens		0.18	0.03		99	
07631130	**Low discrimination**					**07631130**
					GRA	
Actinomyces naeslundii		72.28	0.48	MLT 94	99	
Parabacteroides distasonis		11.59	0.21	MLZ 5	1	
Clostridium perfringens		8.11	0.09		99	
07631131	**Good identification**					**07631131**
Actinomyces naeslundii		100.0	0.51	MLT 94		
07631132	**Acceptable identification**					**07631132**
					GRA	
Parabacteroides distasonis		93.57	0.55	XYL 90	1	
Actinomyces naeslundii		5.89	0.48		99	
Clostridium perfringens		0.31	0.03		99	
07631133	**Acceptable identification**					**07631133**
					GRA	
Actinomyces naeslundii		90.34	0.51	MLT 94	99	
Parabacteroides distasonis		9.66	0.21	MLZ 5	1	
07631140	**Low discrimination**					**07631140**
					GRA	
Fusobacterium mortiferum		76.26	0.54	SOR 1	1	
Actinomyces naeslundii		23.28	0.57		99	
Clostridium perfringens		0.45	0.05		99	
07631141	**Very good identification**					**07631141**
					GRA	
Actinomyces naeslundii		97.84	0.60	MLT 94	99	
Fusobacterium mortiferum		2.16	0.20		1	
07631142	**Low discrimination**					**07631142**
					GRA	
Actinomyces naeslundii		66.52	0.57	MLT 94	99	
Fusobacterium mortiferum		24.21	0.38		1	
Lactobacillus acidophilus		9.27	0.18		99	
07631143	**Very good identification**					**07631143**
Actinomyces naeslundii		100.0	0.60	MLT 94		
07631151	**Good identification**					**07631151**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07631153	**Good identification**					**07631153**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07631160	**Low discrimination**					**07631160**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		60.10	0.57	MLT 94	99		
Clostridium perfringens		37.91	0.31		99		
Fusobacterium mortiferum		1.99	0.20		1		
07631161	**Very good identification**						**07631161**
Actinomyces naeslundii		100.0	0.60	MLT 94			
07631162	**Low discrimination**						**07631162**
Actinomyces naeslundii		65.57	0.57	MLT 94			
Clostridium perfringens		19.46	0.25				
Lactobacillus acidophilus		9.14	0.18				
07631163	**Very good identification**						**07631163**
Actinomyces naeslundii		100.0	0.60	MLT 94			
07631171	**Good identification**						**07631171**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07631173	**Good identification**						**07631173**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07631200	**Low discrimination**						**07631200**
					GRA	CAT	
Fusobacterium mortiferum		63.42	0.48	RHA 1	1	5	
Propionibacterium granulosum		35.68	0.40		99	90	
Actinomyces naeslundii		0.91	0.28		99	25	
07631300	**Good identification**						**07631300**
					GRA		
Fusobacterium mortiferum		98.28	0.54	RHA 1	1		
Actinomyces naeslundii		1.72	0.35		99		
07631301	**Low discrimination**						**07631301**
					GRA		
Actinomyces naeslundii		72.18	0.38	MLT 94	99		
Fusobacterium mortiferum		27.82	0.20	RHA 1	1		
07631302	**Low discrimination**						**07631302**
					GRA		
Fusobacterium mortiferum		47.11	0.38	RHA 1	1		
Parabacteroides distasonis		34.49	0.34	BGL 10	1		
Lactobacillus acidophilus		10.99	0.14		99		
07631303	**Good identification**						**07631303**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07631321	**Good identification**						**07631321**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07631322	**Very good identification**						**07631322**
					SPO		
Parabacteroides distasonis		98.48	0.68	XYL 90	1		
Prevotella oralis		0.56	0.17		1		
Clostridium ramosum		0.37	0.14		92		
07631332	**Very good identification**						**07631332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		99.83	0.63	XYL 90			
Clostridium clostridioforme		0.17	0.10				
07631400	**Low discrimination**						**07631400**
					GRA	CAT	
Fusobacterium mortiferum		83.47	0.48	ARA 1	1	5	
Propionibacterium granulosum		15.34	0.32		99	90	
Actinomyces naeslundii		1.20	0.28		99	25	
07631500	**Good identification**						**07631500**
					GRA		
Fusobacterium mortiferum		97.84	0.54	ARA 1	1		
Actinomyces naeslundii		1.71	0.35		99		
Lactobacillus fermentum		0.45	0.15		99		
07631501	**Low discrimination**						**07631501**
					GRA		
Actinomyces naeslundii		72.18	0.38	MLT 94	99		
Fusobacterium mortiferum		27.82	0.20	ARA 1	1		
07631502	**Low discrimination**						**07631502**
					GRA		
Fusobacterium mortiferum		46.66	0.38	ARA 1	1		
Bifidobacterium adolescentis		34.20	0.31	BGL 10	99		
Actinomyces naeslundii		7.34	0.35		99		
07631503	**Good identification**						**07631503**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07631520	**Low discrimination**						**07631520**
					GRA		
Actinomyces naeslundii		29.31	0.35	MLT 94	99		
Parabacteroides distasonis		27.87	0.22	ARA 1	1		
Clostridium perfringens		17.13	0.09		99		
07631521	**Good identification**						**07631521**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07631522	**Very good identification**						**07631522**
					GRA		
Parabacteroides distasonis		97.95	0.56	XYL 90	1		
Actinomyces naeslundii		1.04	0.35		99		
Bifidobacterium longum subsp. longum		0.31	0.09		99		
07631523	**Low discrimination**						**07631523**
					GRA		
Actinomyces naeslundii		61.20	0.38	MLT 94	99		
Parabacteroides distasonis		38.80	0.22	ARA 1	1		
07631532	**Good identification**						**07631532**
					GRA		
Parabacteroides distasonis		97.31	0.52	XYL 90	1		
Clostridium clostridioforme		1.71	0.16		50		
Bifidobacterium longum subsp. longum		0.98	0.13		99		
07631722	**Very good identification**						**07631722**
Parabacteroides distasonis		99.49	0.64	XYL 90			
Clostridium clostridioforme		0.51	0.19				
07631732	**Very good identification**						**07631732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		95.87	0.60	XYL 90			
Clostridium clostridioforme		4.13	0.31				
07632000	**Good identification**						**07632000**
					GRA	CAT	
Propionibacterium granulosum		94.54	0.64	SAL 1	99	90	
Fusobacterium mortiferum		4.59	0.45		1	5	
Actinomyces naeslundii		0.79	0.43		99	25	
07632001	**Low discrimination**						**07632001**
Actinomyces naeslundii		55.38	0.46	MLT 94			
Propionibacterium granulosum		44.62	0.30	GAL 96			
07632002	**Acceptable identification**						**07632002**
					GRA	CAT	
Propionibacterium granulosum		94.48	0.53	SAL 1	99	90	
Actinomyces naeslundii		3.16	0.43		99	25	
Fusobacterium mortiferum		2.04	0.28		1	5	
07632003	**Low discrimination**						**07632003**
Actinomyces naeslundii		83.24	0.46	MLT 94			
Propionibacterium granulosum		16.76	0.19	GAL 96			
07632010	**Good identification**						**07632010**
					SPO	CAT	
Propionibacterium granulosum		99.62	0.49	SAL 1	1	90	
Clostridium difficile		0.38	-.05	MLZ 12	99	1	
07632012	**Good identification**						**07632012**
Propionibacterium granulosum		100.0	0.39	SAL 1			
07632020	**Good identification**						**07632020**
Propionibacterium granulosum		96.77	0.53	SAL 1			
Actinomyces naeslundii		3.24	0.43				
07632021	**Low discrimination**						**07632021**
Actinomyces naeslundii		83.24	0.46	MLT 94			
Propionibacterium granulosum		16.76	0.19	GAL 96			
07632022	**Low discrimination**						**07632022**
Propionibacterium granulosum		88.20	0.43	SAL 1			
Actinomyces naeslundii		11.80	0.43				
07632023	**Good identification**						**07632023**
Actinomyces naeslundii		95.21	0.46	MLT 94			
Propionibacterium granulosum		4.79	0.09	GAL 96			
07632030	**Good identification**						**07632030**
Propionibacterium granulosum		100.0	0.39	SAL 1			
07632100	**Low discrimination**						**07632100**
					GRA	CAT	
Fusobacterium mortiferum		76.98	0.51	MAN 1	1	5	
Actinomyces naeslundii		16.16	0.51		99	25	
Propionibacterium granulosum		6.86	0.30		99	90	
07632101	**Good identification**						**07632101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		96.89	0.54	MLT 94	99		
Fusobacterium mortiferum		3.11	0.17	GAL 96	1		
07632102	**Low discrimination**						**07632102**
					GRA	CAT	
Actinomyces naeslundii		61.14	0.51	MLT 94	99	25	
Fusobacterium mortiferum		32.37	0.35	GAL 96	1	5	
Propionibacterium granulosum		6.49	0.19		99	90	
07632103	**Good identification**						**07632103**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07632120	**Low discrimination**						**07632120**
					GRA	CAT	
Actinomyces naeslundii		86.63	0.51	MLT 94	99	25	
Propionibacterium granulosum		9.20	0.19	GAL 96	99	90	
Fusobacterium mortiferum		4.17	0.17		1	5	
07632121	**Good identification**						**07632121**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07632122	**Good identification**						**07632122**
Actinomyces naeslundii		97.42	0.51	MLT 94			
Propionibacterium granulosum		2.58	0.09	GAL 96			
07632123	**Good identification**						**07632123**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07632140	**Low discrimination**						**07632140**
					GRA		
Actinomyces naeslundii		78.57	0.38	MLT 94	99		
Fusobacterium mortiferum		21.43	0.17	GAL 96	1		
07632141	**Good identification**						**07632141**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07632142	**Good identification**						**07632142**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07632143	**Good identification**						**07632143**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07632160	**Good identification**						**07632160**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07632161	**Good identification**						**07632161**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07632162	**Good identification**						**07632162**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07632163	**Good identification**						**07632163**
Actinomyces naeslundii		100.0	0.41	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07632200	**Good identification**						**07632200**
Propionibacterium granulosum		100.0	0.38	SAL 1			
07633000	**Low discrimination**						**07633000**
					GRA	CAT	
Propionibacterium granulosum		77.53	0.64	SAL 1	99	90	
Actinomyces naeslundii		15.55	0.67		99	25	
Fusobacterium mortiferum		5.65	0.48		1	5	
07633001	**Very good identification**						**07633001**
					GRA	CAT	
Actinomyces naeslundii		96.52	0.70	MLT 94	99	25	
Propionibacterium granulosum		3.24	0.30		99	90	
Fusobacterium mortiferum		0.24	0.13		1	5	
07633002	**Low discrimination**						**07633002**
					SPO	CAT	
Propionibacterium granulosum		52.57	0.53	SAL 1	1	90	
Actinomyces naeslundii		42.18	0.67		1	25	
Clostridium innocuum		3.44	0.24		99	1	
07633003	**Very good identification**						**07633003**
Actinomyces naeslundii		99.17	0.70	MLT 94			
Propionibacterium granulosum		0.83	0.19				
07633010	**Acceptable identification**						**07633010**
					GRA	CAT	
Propionibacterium granulosum		91.93	0.49	SAL 1	99	90	
Actinomyces naeslundii		7.12	0.45	MLZ 12	99	25	
Fusobacterium mortiferum		0.50	0.13		1	5	
07633011	**Acceptable identification**						**07633011**
Actinomyces naeslundii		92.00	0.48	MLT 94			
Propionibacterium granulosum		8.00	0.15	MLZ 5			
07633012	**Low discrimination**						**07633012**
					SPO	CAT	
Propionibacterium granulosum		75.21	0.39	SAL 1	1	90	
Actinomyces naeslundii		23.29	0.45	MLZ 12	1	25	
Clostridium innocuum		1.50	0.00		99	1	
07633013	**Good identification**						**07633013**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07633020	**Low discrimination**						**07633020**
					GRA	CAT	
Propionibacterium granulosum		55.39	0.53	SAL 1	99	90	
Actinomyces naeslundii		44.45	0.67		99	25	
Fusobacterium mortiferum		0.16	0.13		1	5	
07633021	**Very good identification**						**07633021**
Actinomyces naeslundii		99.17	0.70	MLT 94			
Propionibacterium granulosum		0.83	0.19				
07633022	**Low discrimination**						**07633022**
Actinomyces naeslundii		76.25	0.67	MLT 94			
Propionibacterium granulosum		23.75	0.43				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07633023	**Very good identification**						**07633023**
Actinomyces naeslundii		99.79	0.70	MLT 94			
Propionibacterium granulosum		0.21	0.09				
07633030	**Low discrimination**						**07633030**
Propionibacterium granulosum		76.35	0.39	SAL 1			
Actinomyces naeslundii		23.65	0.45	MLZ 12			
07633031	**Good identification**						**07633031**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07633032	**Low discrimination**						**07633032**
Actinomyces naeslundii		55.34	0.45	MLT 94			
Propionibacterium granulosum		44.67	0.28	MLZ 5			
07633033	**Good identification**						**07633033**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07633040	**Low discrimination**						**07633040**
					GRA	CAT	
Actinomyces naeslundii		76.56	0.54	MLT 94	99	25	
Propionibacterium granulosum		21.85	0.30		99	90	
Fusobacterium mortiferum		1.59	0.13		1	5	
07633041	**Very good identification**						**07633041**
Actinomyces naeslundii		100.0	0.57	MLT 94			
07633042	**Acceptable identification**						**07633042**
Actinomyces naeslundii		93.34	0.54	MLT 94			
Propionibacterium granulosum		6.66	0.19				
07633043	**Very good identification**						**07633043**
Actinomyces naeslundii		100.0	0.57	MLT 94			
07633060	**Acceptable identification**						**07633060**
Actinomyces naeslundii		93.34	0.54	MLT 94			
Propionibacterium granulosum		6.66	0.19				
07633061	**Very good identification**						**07633061**
Actinomyces naeslundii		100.0	0.57	MLT 94			
07633062	**Good identification**						**07633062**
Actinomyces naeslundii		98.25	0.54	MLT 94			
Propionibacterium granulosum		1.75	0.09				
07633063	**Very good identification**						**07633063**
Actinomyces naeslundii		100.0	0.57	MLT 94			
07633100	**Low discrimination**						**07633100**
					GRA	CAT	
Actinomyces naeslundii		75.87	0.74	MLT 94	99	25	
Fusobacterium mortiferum		22.59	0.54		1	5	
Propionibacterium granulosum		1.34	0.30		99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07633101	**Very good identification**						**07633101**
					GRA		
Actinomyces naeslundii		99.80	0.78	MLT 94	99		
Fusobacterium mortiferum		0.20	0.20		1		
07633102	**Very good identification**						**07633102**
					GRA		
Actinomyces naeslundii		95.27	0.74	MLT 94	99		
Fusobacterium mortiferum		3.15	0.38		1		
Lactobacillus acidophilus		0.51	0.11		99		
07633103	**Very good identification**						**07633103**
Actinomyces naeslundii		100.0	0.78	MLT 94			
07633110	**Acceptable identification**						**07633110**
					GRA	CAT	
Actinomyces naeslundii		90.66	0.53	MLT 94	99	25	
Fusobacterium mortiferum		5.18	0.20	MLZ 5	1	5	
Propionibacterium granulosum		4.16	0.15		99	90	
07633111	**Good identification**						**07633111**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07633112	**Acceptable identification**						**07633112**
Actinomyces naeslundii		93.56	0.53	MLT 94			
Bifidobacterium adolescentis		6.44	0.17	MLZ 5			
07633113	**Good identification**						**07633113**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07633120	**Very good identification**						**07633120**
					SPO	CAT	
Actinomyces naeslundii		98.02	0.74	MLT 94	1	25	
Clostridium ramosum		0.95	0.15		92	1	
Propionibacterium granulosum		0.43	0.19		1	90	
07633121	**Very good identification**						**07633121**
Actinomyces naeslundii		100.0	0.78	MLT 94			
07633122	**Good identification**						**07633122**
					SPO	GRA	
Actinomyces naeslundii		94.81	0.74	MLT 94	1	99	
Clostridium ramosum		3.67	0.25		92	50	
Parabacteroides distasonis		0.77	0.26		1	1	
07633123	**Very good identification**						**07633123**
Actinomyces naeslundii		100.0	0.78	MLT 94			
07633130	**Good identification**						**07633130**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07633131	**Good identification**						**07633131**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07633132	**Acceptable identification**						**07633132**
					GRA		
Actinomyces naeslundii		92.37	0.53	MLT 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		7.63	0.21	MLZ 5	1	
07633133	**Good identification**					**07633133**
Actinomyces naeslundii		100.0	0.56	MLT 94		
07633140	**Very good identification**					**07633140**
					GRA	
Actinomyces naeslundii		97.78	0.62	MLT 94	99	
Fusobacterium mortiferum		1.67	0.20		1	
Actinomyces israelii		0.55	0.14		99	
07633141	**Very good identification**					**07633141**
Actinomyces naeslundii		100.0	0.65	MLT 94		
07633142	**Very good identification**					**07633142**
Actinomyces naeslundii		99.16	0.62	MLT 94		
Actinomyces israelii		0.84	0.17			
07633143	**Very good identification**					**07633143**
Actinomyces naeslundii		100.0	0.65	MLT 94		
07633150	**Good identification**					**07633150**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07633151	**Good identification**					**07633151**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07633152	**Good identification**					**07633152**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07633153	**Good identification**					**07633153**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07633160	**Very good identification**					**07633160**
Actinomyces naeslundii		100.0	0.62	MLT 94		
07633161	**Very good identification**					**07633161**
Actinomyces naeslundii		100.0	0.65	MLT 94		
07633162	**Very good identification**					**07633162**
Actinomyces naeslundii		100.0	0.62	MLT 94		
07633163	**Very good identification**					**07633163**
Actinomyces naeslundii		100.0	0.65	MLT 94		
07633170	**Good identification**					**07633170**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07633171	**Good identification**					**07633171**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07633172	**Good identification**					**07633172**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.40	MLT 94			
07633173	**Good identification**						**07633173**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633200	**Acceptable identification**						**07633200**
					GRA	CAT	
Propionibacterium granulosum		91.80	0.38	SAL 1	99	90	
Actinomyces naeslundii		6.02	0.33	RHA 3	99	25	
Fusobacterium mortiferum		2.18	0.13		1	5	
07633201	**Good identification**						**07633201**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633203	**Good identification**						**07633203**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633221	**Good identification**						**07633221**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633223	**Good identification**						**07633223**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633300	**Low discrimination**						**07633300**
					GRA		
Actinomyces naeslundii		77.06	0.40	MLT 94	99		
Fusobacterium mortiferum		22.95	0.20	RHA 1	1		
07633301	**Good identification**						**07633301**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633302	**Good identification**						**07633302**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07633303	**Good identification**						**07633303**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633321	**Good identification**						**07633321**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633323	**Good identification**						**07633323**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633401	**Good identification**						**07633401**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633403	**Good identification**						**07633403**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633421	**Good identification**						**07633421**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633423	**Good identification**						**07633423**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.36	MLT 94			
07633500	**Low discrimination**						**07633500**
					GRA		
Actinomyces naeslundii		66.47	0.40	MLT 94	99		
Fusobacterium mortiferum		19.79	0.20	ARA 1	1		
Actinomyces israelii		13.74	0.19		99		
07633501	**Good identification**						**07633501**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633503	**Good identification**						**07633503**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633520	**Good identification**						**07633520**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07633521	**Good identification**						**07633521**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07633522	**Low discrimination**						**07633522**
					GRA		
Actinomyces naeslundii		67.12	0.40	MLT 94	99		
Parabacteroides distasonis		32.88	0.22	ARA 1	1		
07633523	**Good identification**						**07633523**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07634000	**Very good identification**						**07634000**
					GRA	CAT	
Fusobacterium mortiferum		96.43	0.60	XYL 7	1	5	
Propionibacterium granulosum		3.57	0.32		99	90	
07634002	**Acceptable identification**						**07634002**
					GRA	CAT	
Fusobacterium mortiferum		92.32	0.43	XYL 7	1	5	
Propionibacterium granulosum		7.68	0.21	BGL 10	99	90	
07634100	**Very good identification**						**07634100**
Fusobacterium mortiferum		100.0	0.66	XYL 7			
07634102	**Good identification**						**07634102**
Fusobacterium mortiferum		100.0	0.49	XYL 7			
07634122	**Good identification**						**07634122**
Parabacteroides distasonis		97.77	0.42	GAL 99			
Fusobacterium mortiferum		2.23	0.15				
07634132	**Good identification**						**07634132**
Parabacteroides distasonis		97.51	0.37	GAL 99			
Clostridium clostridioforme		2.49	0.04				
07634322	**Good identification**						**07634322**
Parabacteroides distasonis		99.27	0.50	GAL 99			
Clostridium clostridioforme		0.73	0.08				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07634332	**Acceptable identification**						**07634332**
Parabacteroides distasonis		94.13	0.46	GAL 99			
Clostridium clostridioforme		5.88	0.19				
07634522	**Acceptable identification**						**07634522**
Parabacteroides distasonis		92.84	0.38	GAL 99			
Clostridium clostridioforme		7.16	0.14				
07634712	**Good identification**						**07634712**
Clostridium clostridioforme		100.0	0.40	MLT 99			
07634722	**Low discrimination**						**07634722**
Parabacteroides distasonis		84.13	0.47	GAL 99			
Clostridium clostridioforme		15.87	0.28				
07634732	**Low discrimination**						**07634732**
Clostridium clostridioforme		61.47	0.40	MLT 99			
Parabacteroides distasonis		38.53	0.42				
07635000	**Very good identification**						**07635000**
					GRA	CAT	
Fusobacterium mortiferum		97.28	0.63	XYL 7	1	5	
Propionibacterium granulosum		2.40	0.32		99	90	
Actinomyces naeslundii		0.19	0.28		99	25	
07635002	**Acceptable identification**						**07635002**
					GRA	CAT	
Fusobacterium mortiferum		91.65	0.46	XYL 7	1	5	
Propionibacterium granulosum		5.09	0.21	BGL 10	99	90	
Actinomyces israelii		1.68	0.19		99	1	
07635022	**Acceptable identification**						**07635022**
					GRA		
Parabacteroides distasonis		94.34	0.42	LAC 99	1		
Actinomyces naeslundii		2.37	0.28		99		
Propionibacterium granulosum		1.90	0.11		99		
07635032	**Good identification**						**07635032**
Parabacteroides distasonis		100.0	0.37	LAC 99			
07635100	**Very good identification**						**07635100**
					COC	GRA	
Fusobacterium mortiferum		98.89	0.69	XYL 7	1	1	
Atopobium parvulum		0.52	0.18		99	99	
Actinomyces naeslundii		0.23	0.35		1	99	
07635102	**Low discrimination**						**07635102**
					GRA		
Fusobacterium mortiferum		68.37	0.52	XYL 7	1		
Parabacteroides distasonis		20.15	0.42	BGL 10	1		
Bifidobacterium adolescentis		6.72	0.31		99		
07635103	**Low discrimination**						**07635103**
					GRA		
Actinomyces naeslundii		75.81	0.38	MLT 94	99		
Fusobacterium mortiferum		24.19	0.18	XYL 1	1		
07635112	**Low discrimination**						**07635112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		51.89	0.37	NAG 99	1	
Bifidobacterium adolescentis		39.30	0.33		99	
Clostridium clostridioforme		3.97	0.13		50	
07635120	**Low discrimination**					**07635120**
					GRA	
Parabacteroides distasonis		68.95	0.42	BGL 99	1	
Fusobacterium mortiferum		21.27	0.35		1	
Actinomyces naeslundii		4.94	0.35		99	
07635121	**Good identification**					**07635121**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07635122	**Very good identification**					**07635122**
					GRA	
Parabacteroides distasonis		99.79	0.76		1	
Actinomyces naeslundii		0.07	0.35		99	
Prevotella oralis		0.05	0.08		1	
07635123	**Acceptable identification**					**07635123**
					GRA	
Parabacteroides distasonis		90.30	0.42	URE 1	1	
Actinomyces naeslundii		9.70	0.38		99	
07635130	**Good identification**					**07635130**
Parabacteroides distasonis		100.0	0.37	BGL 99		
07635132	**Very good identification**					**07635132**
Parabacteroides distasonis		99.92	0.72			
Clostridium clostridioforme		0.08	0.13			
07635133	**Good identification**					**07635133**
Parabacteroides distasonis		100.0	0.37	URE 1		
07635162	**Good identification**					**07635162**
					GRA	
Parabacteroides distasonis		98.75	0.42	SOR 1	1	
Actinomyces naeslundii		1.25	0.22		99	
07635172	**Good identification**					**07635172**
Parabacteroides distasonis		100.0	0.37	SOR 1		
07635222	**Good identification**					**07635222**
Parabacteroides distasonis		99.34	0.50	LAC 99		
Clostridium clostridioforme		0.66	0.07			
07635232	**Acceptable identification**					**07635232**
Parabacteroides distasonis		94.70	0.46	LAC 99		
Clostridium clostridioforme		5.30	0.18			
07635302	**Good identification**					**07635302**
Parabacteroides distasonis		96.75	0.50	NAG 99		
Clostridium clostridioforme		2.14	0.16			
Fusobacterium mortiferum		1.11	0.18			
07635312	**Low discrimination**					**07635312**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		84.23	0.46	NAG 99		
Clostridium clostridioforme		15.77	0.27			
07635320	**Good identification**					**07635320**
Parabacteroides distasonis		96.81	0.50	BGL 99		
Bacteroides vulgatus		3.19	0.06			
07635322	**Very good identification**					**07635322**
Parabacteroides distasonis		99.97	0.84			
Clostridium clostridioforme		0.02	0.16			
Prevotella oralis		0.01	0.01			
07635323	**Good identification**					**07635323**
Parabacteroides distasonis		100.0	0.50	URE 1		
07635330	**Good identification**					**07635330**
Parabacteroides distasonis		100.0	0.46	BGL 99		
07635332	**Very good identification**					**07635332**
Parabacteroides distasonis		99.81	0.80			
Clostridium clostridioforme		0.19	0.27			
07635333	**Good identification**					**07635333**
Parabacteroides distasonis		100.0	0.46	URE 1		
07635362	**Good identification**					**07635362**
Parabacteroides distasonis		100.0	0.50	SOR 1		
07635372	**Good identification**					**07635372**
Parabacteroides distasonis		100.0	0.46	SOR 1		
07635422	**Acceptable identification**					**07635422**
Parabacteroides distasonis		93.53	0.38	LAC 99		
Clostridium clostridioforme		6.47	0.13			
07635502	**Low discrimination**					**07635502**
					GRA	
Bifidobacterium adolescentis		68.47	0.45	MLT 99	99	
Parabacteroides distasonis		18.79	0.38		1	
Clostridium clostridioforme		4.35	0.22		50	
07635512	**Low discrimination**					**07635512**
					GRA	
Bifidobacterium adolescentis		72.86	0.47	MLT 99	99	
Clostridium clostridioforme		17.25	0.33		50	
Parabacteroides distasonis		8.81	0.34		1	
07635520	**Low discrimination**					**07635520**
					GRA	
Parabacteroides distasonis		71.83	0.38	BGL 99	1	
Bacteroides vulgatus		17.09	0.09		1	
Bifidobacterium longum subsp. longum		7.52	0.17		99	
07635522	**Very good identification**					**07635522**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		99.63	0.73		1	
Clostridium clostridioforme		0.23	0.22		50	
Bifidobacterium longum subsp. longum		0.11	0.17		99	
07635523	**Good identification**					**07635523**
Parabacteroides distasonis		100.0	0.38	URE 1		
07635532	**Very good identification**					**07635532**
					GRA	
Parabacteroides distasonis		97.66	0.68		1	
Clostridium clostridioforme		1.93	0.33		50	
Bifidobacterium longum subsp. longum		0.33	0.21		99	
07635562	**Good identification**					**07635562**
Parabacteroides distasonis		100.0	0.38	SOR 1		
07635612	**Good identification**					**07635612**
Clostridium clostridioforme		100.0	0.39	MLT 99		
07635622	**Low discrimination**					**07635622**
Parabacteroides distasonis		85.54	0.47	LAC 99		
Clostridium clostridioforme		14.46	0.28			
07635632	**Low discrimination**					**07635632**
Clostridium clostridioforme		58.84	0.39	MLT 99		
Parabacteroides distasonis		41.16	0.42			
07635702	**Low discrimination**					**07635702**
					GRA	
Parabacteroides distasonis		63.37	0.47	NAG 99	1	
Clostridium clostridioforme		35.85	0.37		50	
Bifidobacterium adolescentis		0.78	0.11		99	
07635712	**Low discrimination**					**07635712**
					GRA	
Clostridium clostridioforme		82.32	0.48	MLT 99	50	
Parabacteroides distasonis		17.20	0.42		1	
Bifidobacterium adolescentis		0.48	0.13		99	
07635720	**Low discrimination**					**07635720**
Parabacteroides distasonis		58.35	0.47	BGL 99		
Bacteroides vulgatus		41.65	0.26			
07635722	**Very good identification**					**07635722**
Parabacteroides distasonis		99.37	0.81			
Clostridium clostridioforme		0.57	0.37			
Bacteroides thetaiotaomicron		0.05	0.09			
07635723	**Good identification**					**07635723**
Parabacteroides distasonis		100.0	0.47	URE 1		
07635730	**Acceptable identification**					**07635730**
Parabacteroides distasonis		94.19	0.42	BGL 99		
Clostridium clostridioforme		4.55	0.14			
Bacteroides vulgatus		1.26	-.09			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07635732	**Very good identification**					**07635732**
Parabacteroides distasonis		95.35	0.76			
Clostridium clostridioforme		4.61	0.48			
Bacteroides thetaiotaomicron		0.04	0.02			
07635733	**Good identification**					**07635733**
Parabacteroides distasonis		95.39	0.42	URE 1		
Clostridium clostridioforme		4.61	0.14			
07635762	**Good identification**					**07635762**
Parabacteroides distasonis		100.0	0.47	SOR 1		
07635772	**Good identification**					**07635772**
Parabacteroides distasonis		95.39	0.42	SOR 1		
Clostridium clostridioforme		4.61	0.14			
07637001	**Good identification**					**07637001**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07637003	**Good identification**					**07637003**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07637021	**Good identification**					**07637021**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07637023	**Good identification**					**07637023**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07637100	**Low discrimination**					**07637100**
					GRA	
Actinomyces israelii		47.43	0.39	MLT 99	99	
Fusobacterium mortiferum		36.24	0.35	CEL 93	1	
Actinomyces naeslundii		16.33	0.40		99	
07637101	**Good identification**					**07637101**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07637102	**Low discrimination**					**07637102**
Actinomyces israelii		57.57	0.42	MLT 99		
Bifidobacterium adolescentis		25.96	0.30	CEL 93		
Actinomyces naeslundii		13.22	0.40			
07637103	**Good identification**					**07637103**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07637120	**Low discrimination**					**07637120**
Actinomyces naeslundii		84.36	0.40	MLT 94		
Actinomyces israelii		15.64	0.19	XYL 1		
07637121	**Good identification**					**07637121**
Actinomyces naeslundii		100.0	0.43	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07637122	**Low discrimination**					**07637122**
					GRA	
Parabacteroides distasonis		84.92	0.42	MAN 1	1	
Actinomyces naeslundii		11.80	0.40		99	
Actinomyces israelii		3.28	0.22		99	
07637123	**Good identification**					**07637123**
Actinomyces naeslundii		100.0	0.43	MLT 94		
07637132	**Good identification**					**07637132**
Parabacteroides distasonis		100.0	0.37	MAN 1		
07637142	**Low discrimination**					**07637142**
Actinomyces israelii		80.16	0.35	MLT 99		
Bifidobacterium adolescentis		11.49	0.14	CEL 93		
Actinomyces naeslundii		8.35	0.27			
07637322	**Good identification**					**07637322**
Parabacteroides distasonis		100.0	0.50	MAN 1		
07637332	**Good identification**					**07637332**
Parabacteroides distasonis		100.0	0.46	MAN 1		
07637500	**Low discrimination**					**07637500**
					SPO	
Actinomyces israelii		85.23	0.38	MLT 99	1	
Clostridium butyricum		10.00	0.01	CEL 93	99	
Bifidobacterium adolescentis		4.76	0.10		1	
07637502	**Low discrimination**					**07637502**
					SPO	
Bifidobacterium adolescentis		78.16	0.44	MLT 99	1	
Actinomyces israelii		21.20	0.41		1	
Clostridium butyricum		0.65	-.07		99	
07637512	**Good identification**					**07637512**
Bifidobacterium adolescentis		95.86	0.45	MLT 99		
Actinomyces israelii		3.18	0.26			
Bifidobacterium dentium		0.97	-.07			
07637522	**Acceptable identification**					**07637522**
					GRA	
Parabacteroides distasonis		92.45	0.38	MAN 1	1	
Actinomyces israelii		4.77	0.20		99	
Bifidobacterium adolescentis		2.78	0.10		99	
07637722	**Good identification**					**07637722**
Parabacteroides distasonis		100.0	0.47	MAN 1		
07637732	**Good identification**					**07637732**
Parabacteroides distasonis		95.39	0.42	MAN 1		
Clostridium clostridioforme		4.61	0.14			
07640000	**Good identification**					**07640000**
Fusobacterium mortiferum		100.0	0.45	FRU 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07640100	**Good identification**					**07640100**
Fusobacterium mortiferum		100.0	0.52	FRU 99		
07640102	**Good identification**					**07640102**
Fusobacterium mortiferum		100.0	0.35	FRU 99		
07641000	**Low discrimination**					**07641000**
					GRA	
Fusobacterium mortiferum		86.97	0.48	FRU 99	1	
Actinomyces naeslundii		3.97	0.37		99	
Eubacterium tortuosum		3.22	0.07		99	
07641001	**Low discrimination**					**07641001**
					GRA	
Actinomyces naeslundii		87.15	0.40	MLT 94	99	
Fusobacterium mortiferum		12.85	0.14	FRU 95	1	
07641003	**Good identification**					**07641003**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07641020	**Low discrimination**					**07641020**
					GRA	
Actinomyces naeslundii		65.45	0.37	MLT 94	99	
Fusobacterium mortiferum		14.48	0.14	FRU 95	1	
Lactobacillus catenaformis		10.70	0.02		99	
07641021	**Good identification**					**07641021**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07641022	**Low discrimination**					**07641022**
					SPO	
Actinomyces naeslundii		51.84	0.37	MLT 94	1	
Lactobacillus acidophilus		21.22	0.06	FRU 95	1	
Clostridium barati		9.42	-.02		99	
07641023	**Good identification**					**07641023**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07641100	**Acceptable identification**					**07641100**
					GRA	
Fusobacterium mortiferum		93.03	0.55	FRU 99	1	
Actinomyces naeslundii		5.18	0.45		99	
Lactobacillus catenaformis		1.79	0.15		99	
07641101	**Low discrimination**					**07641101**
					GRA	
Actinomyces naeslundii		89.22	0.48	MLT 94	99	
Fusobacterium mortiferum		10.78	0.21	FRU 95	1	
07641102	**Low discrimination**					**07641102**
					GRA	
Fusobacterium mortiferum		39.05	0.38	FRU 99	1	
Lactobacillus acidophilus		32.38	0.24	BGL 10	99	
Actinomyces naeslundii		19.58	0.45		99	
07641103	**Good identification**					**07641103**
Actinomyces naeslundii		100.0	0.48	MLT 94		
07641120	**Low discrimination**					**07641120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		67.14	0.45	MLT 94	99		
Fusobacterium mortiferum		12.17	0.21	FRU 95	1		
Prevotella melaninogenica		7.23	-.01		1		
07641121	**Good identification**						**07641121**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07641123	**Good identification**						**07641123**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07643000	**Good identification**						**07643000**
					GRA	SPO	
Clostridium innocuum		96.79	0.44	RAF 1	99	99	
Actinomyces naeslundii		2.88	0.42	FRU 95	99	1	
Fusobacterium mortiferum		0.33	0.14		1	1	
07643001	**Low discrimination**						**07643001**
					SPO		
Actinomyces naeslundii		81.54	0.45	MLT 94	1		
Clostridium innocuum		18.46	0.10	FRU 95	99		
07643002	**Good identification**						**07643002**
					SPO		
Clostridium innocuum		97.11	0.44	RAF 1	99		
Actinomyces naeslundii		2.89	0.42	FRU 95	1		
07643003	**Low discrimination**						**07643003**
					SPO		
Actinomyces naeslundii		81.54	0.45	MLT 94	1		
Clostridium innocuum		18.46	0.10	FRU 95	99		
07643020	**Low discrimination**						**07643020**
					SPO		
Actinomyces naeslundii		74.65	0.42	MLT 94	1		
Clostridium innocuum		25.35	0.10	FRU 95	99		
07643021	**Good identification**						**07643021**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07643022	**Low discrimination**						**07643022**
					SPO		
Actinomyces naeslundii		74.65	0.42	MLT 94	1		
Clostridium innocuum		25.35	0.10	FRU 95	99		
07643023	**Good identification**						**07643023**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07643100	**Low discrimination**						**07643100**
					GRA	SPO	
Actinomyces naeslundii		82.46	0.50	MLT 94	99	1	
Clostridium innocuum		9.84	0.10	FRU 95	99	99	
Fusobacterium mortiferum		7.70	0.21		1	1	
07643101	**Good identification**						**07643101**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07643102	**Low discrimination**						**07643102**
					SPO		
Actinomyces naeslundii		85.90	0.50	MLT 94	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		10.25	0.10	FRU 95	99		
Lactobacillus acidophilus		3.85	0.02		1		
07643103	**Good identification**						**07643103**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07643120	**Good identification**						**07643120**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07643121	**Good identification**						**07643121**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07643122	**Low discrimination**						**07643122**
					SPO		
Actinomyces naeslundii		89.80	0.50	MLT 94	1		
Clostridium ramosum		6.18	0.05	FRU 95	92		
Lactobacillus acidophilus		4.03	0.02		1		
07643123	**Good identification**						**07643123**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07643140	**Good identification**						**07643140**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07643141	**Good identification**						**07643141**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07643142	**Good identification**						**07643142**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07643143	**Good identification**						**07643143**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07643160	**Good identification**						**07643160**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07643161	**Good identification**						**07643161**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07643162	**Good identification**						**07643162**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07643163	**Good identification**						**07643163**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07645100	**Good identification**						**07645100**
Fusobacterium mortiferum		100.0	0.35	FRU 99			
07650000	**Very good identification**						**07650000**
					GRA	CAT	
Fusobacterium mortiferum		99.76	0.80		1	5	
Propionibacterium granulosum		0.08	0.24		99	90	
Actinomyces naeslundii		0.06	0.35		99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07650001	**Acceptable identification**						**07650001**
					GRA		
Fusobacterium mortiferum		92.48	0.45	URE 1	1		
Actinomyces naeslundii		7.52	0.38		99		
07650002	**Very good identification**						**07650002**
					GRA		
Fusobacterium mortiferum		97.79	0.63	BGL 10	1		
Actinomyces naeslundii		0.48	0.35		99		
Lactobacillus catenaformis		0.40	0.12		99		
07650010	**Good identification**						**07650010**
					GRA	CAT	
Fusobacterium mortiferum		98.89	0.45	MLZ 1	1	5	
Propionibacterium granulosum		1.11	0.09		99	90	
07650020	**Acceptable identification**						**07650020**
					GRA	SPO	
Fusobacterium mortiferum		90.28	0.45	NAG 1	1	1	
Actinomyces naeslundii		4.90	0.35		99	1	
Clostridium barati		1.95	0.02		99	99	
07650021	**Low discrimination**						**07650021**
					GRA		
Actinomyces naeslundii		88.95	0.38	MLT 94	99		
Fusobacterium mortiferum		11.05	0.11	GAL 96	1		
07650023	**Good identification**						**07650023**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07650040	**Good identification**						**07650040**
					GRA		
Fusobacterium mortiferum		99.05	0.45	SOR 1	1		
Actinomyces naeslundii		0.95	0.22		99		
07650100	**Very good identification**						**07650100**
					GRA		
Fusobacterium mortiferum		99.90	0.86		1		
Actinomyces naeslundii		0.07	0.43		99		
Lactobacillus catenaformis		0.03	0.15		99		
07650101	**Acceptable identification**						**07650101**
					GRA		
Fusobacterium mortiferum		90.97	0.52	URE 1	1		
Actinomyces naeslundii		9.03	0.46		99		
07650102	**Very good identification**						**07650102**
					GRA		
Fusobacterium mortiferum		97.42	0.70	BGL 10	1		
Lactobacillus acidophilus		1.22	0.24		99		
Actinomyces naeslundii		0.59	0.43		99		
07650103	**Low discrimination**						**07650103**
					GRA		
Fusobacterium mortiferum		50.72	0.35	URE 1	1		
Actinomyces naeslundii		45.30	0.46	BGL 10	99		
Actinomyces viscosus		3.98	0.04		99		
07650110	**Good identification**						**07650110**
Fusobacterium mortiferum		100.0	0.52	MLZ 1			
07650112	**Good identification**						**07650112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.35	MLZ 1			
07650120	**Acceptable identification**						**07650120**
					GRA	SPO	
Fusobacterium mortiferum		90.47	0.52	NAG 1	1	1	
Actinomyces naeslundii		5.99	0.43		99	1	
Clostridium barati		2.52	0.10		99	99	
07650121	**Acceptable identification**						**07650121**
					GRA		
Actinomyces naeslundii		90.76	0.46	MLT 94	99		
Fusobacterium mortiferum		9.24	0.18	GAL 96	1		
07650123	**Good identification**						**07650123**
Actinomyces naeslundii		100.0	0.46	MLT 94			
07650140	**Good identification**						**07650140**
					GRA		
Fusobacterium mortiferum		98.85	0.52	SOR 1	1		
Actinomyces naeslundii		1.15	0.30		99		
07650142	**Low discrimination**						**07650142**
					GRA		
Fusobacterium mortiferum		79.44	0.35	SOR 1	1		
Lactobacillus acidophilus		12.21	0.09	BGL 10	99		
Actinomyces naeslundii		8.35	0.30		99		
07650200	**Good identification**						**07650200**
Fusobacterium mortiferum		100.0	0.45	RHA 1			
07650300	**Good identification**						**07650300**
Fusobacterium mortiferum		100.0	0.52	RHA 1			
07650302	**Acceptable identification**						**07650302**
					GRA		
Fusobacterium mortiferum		91.44	0.35	RHA 1	1		
Lactobacillus acidophilus		8.56	0.05	BGL 10	99		
07650400	**Good identification**						**07650400**
Fusobacterium mortiferum		100.0	0.45	ARA 1			
07650500	**Good identification**						**07650500**
Fusobacterium mortiferum		100.0	0.52	ARA 1			
07650502	**Good identification**						**07650502**
Fusobacterium mortiferum		100.0	0.35	ARA 1			
07651000	**Good identification**						**07651000**
					GRA		
Fusobacterium mortiferum		94.95	0.83		1		
Lactobacillus catenaformis		2.84	0.47		99		
Actinomyces naeslundii		0.83	0.59		99		
07651001	**Low discrimination**						**07651001**
					GRA		
Actinomyces naeslundii		55.66	0.62	MLT 94	99		
Fusobacterium mortiferum		42.76	0.48		1		
Lactobacillus catenaformis		1.28	0.12		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07651002	**Low discrimination**					**07651002**
					GRA	
Fusobacterium mortiferum		60.92	0.66	BGL 10	1	
Lactobacillus catenaformis		16.37	0.47		99	
Lactobacillus acidophilus		10.25	0.40		99	
07651003	**Low discrimination**					**07651003**
					GRA	
Actinomyces naeslundii		89.08	0.62	MLT 94	99	
Fusobacterium mortiferum		7.60	0.32		1	
Lactobacillus catenaformis		2.04	0.12		99	
07651010	**Low discrimination**					**07651010**
					GRA	
Fusobacterium mortiferum		74.57	0.48	MLZ 1	1	
Lactobacillus jensenii		17.47	0.28		99	
Actinomyces naeslundii		3.41	0.37		99	
07651011	**Low discrimination**					**07651011**
					GRA	
Actinomyces naeslundii		87.15	0.40	MLT 94	99	
Fusobacterium mortiferum		12.85	0.14	MLZ 5	1	
07651013	**Good identification**					**07651013**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07651020	**Low discrimination**					**07651020**
					GRA	
Fusobacterium mortiferum		31.88	0.48	NAG 1	1	
Actinomyces naeslundii		27.67	0.59		99	
Lactobacillus catenaformis		23.56	0.36		99	
07651021	**Very good identification**					**07651021**
					GRA	
Actinomyces naeslundii		98.67	0.62	MLT 94	99	
Fusobacterium mortiferum		0.77	0.14		1	
Lactobacillus catenaformis		0.57	0.02		99	
07651022	**Low discrimination**					**07651022**
					SPO	
Lactobacillus acidophilus		34.90	0.40	MLT 94	1	
Actinomyces naeslundii		16.37	0.59	LAC 92	1	
Clostridium barati		15.49	0.32		99	
07651023	**Very good identification**					**07651023**
					SPO	
Actinomyces naeslundii		96.84	0.62	MLT 94	1	
Lactobacillus acidophilus		1.39	0.06		1	
Clostridium barati		0.62	-.02		99	
07651031	**Good identification**					**07651031**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07651032	**Low discrimination**					**07651032**
Actinomyces naeslundii		35.92	0.37	MLT 94		
Lactobacillus jensenii		30.71	0.14	MLZ 5		
Lactobacillus acidophilus		14.70	0.06			
07651033	**Good identification**					**07651033**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.40	MLT 94			
07651040	**Low discrimination**						**07651040**
					GRA		
Fusobacterium mortiferum		83.47	0.48	SOR 1	1		
Actinomyces naeslundii		12.78	0.46		99		
Lactobacillus catenaformis		2.49	0.12		99		
07651041	**Good identification**						**07651041**
					GRA		
Actinomyces naeslundii		95.79	0.49	MLT 94	99		
Fusobacterium mortiferum		4.21	0.14		1		
07651043	**Good identification**						**07651043**
Actinomyces naeslundii		100.0	0.49	MLT 94			
07651060	**Low discrimination**						**07651060**
					GRA	SPO	
Actinomyces naeslundii		78.73	0.46	MLT 94	99	1	
Clostridium barati		12.24	0.05		99	99	
Fusobacterium mortiferum		5.19	0.14		1	1	
07651061	**Good identification**						**07651061**
Actinomyces naeslundii		100.0	0.49	MLT 94			
07651063	**Good identification**						**07651063**
Actinomyces naeslundii		100.0	0.49	MLT 94			
07651100	**Very good identification**						**07651100**
					GRA		
Fusobacterium mortiferum		96.86	0.89		1		
Lactobacillus catenaformis		1.86	0.50		99		
Actinomyces naeslundii		1.04	0.67		99		
07651101	**Low discrimination**						**07651101**
					GRA		
Actinomyces naeslundii		60.91	0.70	MLT 94	99		
Fusobacterium mortiferum		38.36	0.55		1		
Lactobacillus catenaformis		0.74	0.15		99		
07651102	**Low discrimination**						**07651102**
					GRA		
Fusobacterium mortiferum		46.20	0.73	BGL 10	1		
Lactobacillus acidophilus		38.31	0.58		99		
Lactobacillus catenaformis		7.98	0.50		99		
07651103	**Low discrimination**						**07651103**
					GRA		
Actinomyces naeslundii		87.23	0.70	MLT 94	99		
Fusobacterium mortiferum		6.10	0.38		1		
Lactobacillus acidophilus		5.06	0.24		99		
07651110	**Acceptable identification**						**07651110**
					GRA		
Fusobacterium mortiferum		92.71	0.55	MLZ 1	1		
Actinomyces naeslundii		5.17	0.45		99		
Lactobacillus catenaformis		1.78	0.15		99		
07651111	**Low discrimination**						**07651111**
					GRA		
Actinomyces naeslundii		89.22	0.48	MLT 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		10.78	0.21	MLZ 5	1	
07651112	**Low discrimination**					**07651112**
					GRA	
Fusobacterium mortiferum		30.17	0.38	MLZ 1	1	
Lactobacillus acidophilus		25.02	0.24	BGL 10	99	
Actinomyces naeslundii		15.13	0.45		99	
07651113	**Good identification**					**07651113**
Actinomyces naeslundii		100.0	0.48	MLT 94		
07651120	**Low discrimination**					**07651120**
					GRA	
Actinomyces naeslundii		30.86	0.67	MLT 94	99	
Fusobacterium mortiferum		29.15	0.55		1	
Lactobacillus catenaformis		13.85	0.39		99	
07651121	**Very good identification**					**07651121**
					GRA	
Actinomyces naeslundii		98.68	0.70	MLT 94	99	
Fusobacterium mortiferum		0.63	0.21		1	
Lactobacillus catenaformis		0.30	0.05		99	
07651122	**Low discrimination**					**07651122**
					GRA	
Lactobacillus acidophilus		51.69	0.58	MLT 94	99	
Prevotella oralis		30.00	0.53		1	
Actinomyces naeslundii		6.00	0.67		99	
07651123	**Good identification**					**07651123**
					GRA	
Actinomyces naeslundii		90.64	0.70	MLT 94	99	
Lactobacillus acidophilus		5.26	0.24		99	
Prevotella oralis		3.05	0.19		1	
07651130	**Low discrimination**					**07651130**
					GRA	
Actinomyces naeslundii		66.58	0.45	MLT 94	99	
Fusobacterium mortiferum		12.07	0.21	MLZ 5	1	
Bifidobacterium breve		8.00	0.15		99	
07651131	**Good identification**					**07651131**
Actinomyces naeslundii		100.0	0.48	MLT 94		
07651132	**Low discrimination**					**07651132**
					GRA	
Parabacteroides distasonis		34.99	0.39	XYL 90	1	
Lactobacillus acidophilus		21.06	0.24		99	
Bifidobacterium breve		15.47	0.32		99	
07651133	**Good identification**					**07651133**
Actinomyces naeslundii		100.0	0.48	MLT 94		
07651140	**Low discrimination**					**07651140**
					GRA	
Fusobacterium mortiferum		81.21	0.55	SOR 1	1	
Actinomyces naeslundii		15.17	0.54		99	
Lactobacillus catenaformis		1.56	0.15		99	
07651141	**Very good identification**					**07651141**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.52	0.57	MLT 94	99	
Fusobacterium mortiferum		3.48	0.21		1	
07651142	**Low discrimination**					**07651142**
					GRA	
Lactobacillus acidophilus		74.02	0.43	MLT 94	99	
Actinomyces naeslundii		12.27	0.54	SOR 11	99	
Fusobacterium mortiferum		7.30	0.38		1	
07651143	**Very good identification**					**07651143**
Actinomyces naeslundii		96.10	0.57	MLT 94		
Lactobacillus acidophilus		3.91	0.09			
07651160	**Low discrimination**					**07651160**
					SPO	
Actinomyces naeslundii		71.55	0.54	MLT 94	1	
Clostridium barati		11.73	0.14		99	
Lactobacillus acidophilus		4.36	0.09		1	
07651161	**Very good identification**					**07651161**
Actinomyces naeslundii		100.0	0.57	MLT 94		
07651162	**Low discrimination**					**07651162**
					SPO	
Lactobacillus acidophilus		73.45	0.43	MLT 94	1	
Actinomyces naeslundii		12.17	0.54	SOR 11	1	
Clostridium barati		5.98	0.22		99	
07651163	**Very good identification**					**07651163**
Actinomyces naeslundii		96.10	0.57	MLT 94		
Lactobacillus acidophilus		3.91	0.09			
07651200	**Acceptable identification**					**07651200**
					GRA	
Fusobacterium mortiferum		94.91	0.48	RHA 1	1	
Lactobacillus catenaformis		2.83	0.12		99	
Actinomyces naeslundii		0.83	0.25		99	
07651300	**Acceptable identification**					**07651300**
					GRA	
Fusobacterium mortiferum		94.94	0.55	RHA 1	1	
Lactobacillus catenaformis		1.82	0.15		99	
Actinomyces naeslundii		1.02	0.33		99	
07651301	**Low discrimination**					**07651301**
					GRA	
Actinomyces naeslundii		61.36	0.36	MLT 94	99	
Fusobacterium mortiferum		38.64	0.21	RHA 1	1	
07651302	**Low discrimination**					**07651302**
					GRA	
Lactobacillus acidophilus		69.62	0.39	MLT 94	99	
Prevotella oralis		12.11	0.25	RHA 7	1	
Fusobacterium mortiferum		11.27	0.38		1	
07651303	**Low discrimination**					**07651303**
Actinomyces naeslundii		69.81	0.36	MLT 94		
Lactobacillus acidophilus		30.19	0.05	RHA 1		
07651321	**Good identification**					**07651321**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.36	MLT 94			
07651322	**Low discrimination**						**07651322**
					GRA		
Prevotella oralis		55.50	0.47	MLT 99	1		
Parabacteroides distasonis		20.86	0.52		1		
Lactobacillus acidophilus		16.79	0.39		99		
07651332	**Acceptable identification**						**07651332**
					GRA		
Parabacteroides distasonis		93.33	0.48	XYL 90	1		
Prevotella oralis		4.66	0.13		1		
Lactobacillus acidophilus		1.41	0.05		99		
07651400	**Low discrimination**						**07651400**
					GRA	SPO	
Fusobacterium mortiferum		87.21	0.48	ARA 1	1	1	
Clostridium innocuum		8.80	0.19		99	99	
Lactobacillus catenaformis		2.60	0.12		99	1	
07651500	**Acceptable identification**						**07651500**
					GRA	SPO	
Fusobacterium mortiferum		94.98	0.55	ARA 1	1	1	
Lactobacillus catenaformis		1.82	0.15		99	1	
Clostridium butyricum		1.42	0.05		99	99	
07651501	**Low discrimination**						**07651501**
					GRA		
Actinomyces naeslundii		61.36	0.36	MLT 94	99		
Fusobacterium mortiferum		38.64	0.21	ARA 1	1		
07651502	**Low discrimination**						**07651502**
					GRA		
Bifidobacterium adolescentis		47.67	0.40	MLT 99	99		
Fusobacterium mortiferum		22.78	0.38	XYL 87	1		
Lactobacillus acidophilus		18.89	0.24		99		
07651503	**Good identification**						**07651503**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07651512	**Good identification**						**07651512**
Bifidobacterium adolescentis		97.81	0.41	MLT 99			
Clostridium clostridioforme		2.20	0.10	XYL 87			
07651521	**Good identification**						**07651521**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07651522	**Low discrimination**						**07651522**
					GRA		
Parabacteroides distasonis		50.36	0.40	XYL 90	1		
Lactobacillus acidophilus		26.63	0.24		99		
Prevotella oralis		15.46	0.19		1		
07651523	**Good identification**						**07651523**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07651532	**Acceptable identification**						**07651532**
					GRA		
Parabacteroides distasonis		91.03	0.36	XYL 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		6.19	0.10		50		
Bifidobacterium adolescentis		2.79	0.07		99		
07651722	**Acceptable identification**						**07651722**
Parabacteroides distasonis		92.87	0.48	XYL 90			
Prevotella oralis		4.07	0.13				
Clostridium clostridioforme		1.82	0.14				
07651732	**Low discrimination**						**07651732**
Parabacteroides distasonis		85.75	0.44	XYL 90			
Clostridium clostridioforme		14.25	0.25				
07652000	**Low discrimination**						**07652000**
					GRA	SPO	
Clostridium innocuum		79.29	0.44	RAF 1	99	99	
Fusobacterium mortiferum		17.74	0.45	GAL 95	1	1	
Actinomyces naeslundii		1.87	0.40		99	1	
07652001	**Low discrimination**						**07652001**
					GRA	SPO	
Actinomyces naeslundii		74.08	0.43	MLT 94	99	1	
Clostridium innocuum		21.18	0.10	GAL 96	99	99	
Fusobacterium mortiferum		4.74	0.11		1	1	
07652002	**Good identification**						**07652002**
					GRA	SPO	
Clostridium innocuum		95.07	0.44	RAF 1	99	99	
Fusobacterium mortiferum		2.36	0.29	GAL 95	1	1	
Actinomyces naeslundii		2.24	0.40		99	1	
07652003	**Low discrimination**						**07652003**
					SPO		
Actinomyces naeslundii		77.77	0.43	MLT 94	1		
Clostridium innocuum		22.23	0.10	GAL 96	99		
07652020	**Low discrimination**						**07652020**
					SPO	CAT	
Actinomyces naeslundii		59.80	0.40	MLT 94	1	25	
Clostridium innocuum		25.65	0.10	GAL 96	99	1	
Propionibacterium granulosum		8.82	0.11		1	90	
07652021	**Good identification**						**07652021**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07652022	**Low discrimination**						**07652022**
					SPO		
Actinomyces naeslundii		69.99	0.40	MLT 94	1		
Clostridium innocuum		30.01	0.10	GAL 96	99		
07652023	**Good identification**						**07652023**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07652100	**Low discrimination**						**07652100**
					GRA	SPO	
Fusobacterium mortiferum		87.13	0.52	MAN 1	1	1	
Actinomyces naeslundii		11.19	0.48		99	1	
Clostridium innocuum		1.69	0.10		99	99	
07652101	**Good identification**						**07652101**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		95.02	0.51	MLT 94	99		
Fusobacterium mortiferum		4.98	0.18	GAL 96	1		
07652102	**Low discrimination**						**07652102**
					GRA	SPO	
Actinomyces naeslundii		48.25	0.48	MLT 94	99	1	
Fusobacterium mortiferum		41.75	0.35	GAL 96	1	1	
Clostridium innocuum		7.27	0.10		99	99	
07652103	**Good identification**						**07652103**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07652120	**Acceptable identification**						**07652120**
					GRA		
Actinomyces naeslundii		92.71	0.48	MLT 94	99		
Fusobacterium mortiferum		7.29	0.18	GAL 96	1		
07652121	**Good identification**						**07652121**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07652122	**Low discrimination**						**07652122**
					SPO		
Actinomyces naeslundii		83.33	0.48	MLT 94	1		
Clostridium ramosum		7.24	0.05	GAL 96	92		
Lactobacillus acidophilus		4.72	0.02		1		
07652123	**Good identification**						**07652123**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07652140	**Low discrimination**						**07652140**
					GRA		
Actinomyces naeslundii		69.18	0.35	MLT 94	99		
Fusobacterium mortiferum		30.83	0.18	GAL 96	1		
07652141	**Good identification**						**07652141**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07652142	**Good identification**						**07652142**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07652143	**Good identification**						**07652143**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07652160	**Good identification**						**07652160**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07652161	**Good identification**						**07652161**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07652162	**Good identification**						**07652162**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07652163	**Good identification**						**07652163**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07653000	**Very good identification**						**07653000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium innocuum		95.31	0.66	RAF 1	99	99	
Actinomyces naeslundii		2.84	0.64		99	1	
Fusobacterium mortiferum		1.68	0.48		1	1	
07653001	**Low discrimination**						**07653001**
					GRA	SPO	
Actinomyces naeslundii		81.28	0.67	MLT 94	99	1	
Clostridium innocuum		18.40	0.32		99	99	
Fusobacterium mortiferum		0.33	0.14		1	1	
07653002	**Very good identification**						**07653002**
					GRA	SPO	
Clostridium innocuum		96.62	0.66	RAF 1	99	99	
Actinomyces naeslundii		2.88	0.64		99	1	
Fusobacterium mortiferum		0.19	0.32		1	1	
07653003	**Low discrimination**						**07653003**
					SPO		
Actinomyces naeslundii		81.54	0.67	MLT 94	1		
Clostridium innocuum		18.46	0.32		99		
07653010	**Good identification**						**07653010**
					GRA	SPO	
Clostridium innocuum		95.98	0.42	RAF 1	99	99	
Actinomyces naeslundii		3.61	0.42	MLZ 4	99	1	
Fusobacterium mortiferum		0.41	0.14		1	1	
07653011	**Low discrimination**						**07653011**
					SPO		
Actinomyces naeslundii		84.80	0.45	MLT 94	1		
Clostridium innocuum		15.20	0.08	MLZ 5	99		
07653012	**Good identification**						**07653012**
					SPO		
Clostridium innocuum		96.38	0.42	RAF 1	99		
Actinomyces naeslundii		3.62	0.42	MLZ 4	1		
07653013	**Low discrimination**						**07653013**
					SPO		
Actinomyces naeslundii		84.80	0.45	MLT 94	1		
Clostridium innocuum		15.20	0.08	MLZ 5	99		
07653020	**Low discrimination**						**07653020**
					SPO		
Actinomyces naeslundii		72.98	0.64	MLT 94	1		
Clostridium innocuum		24.78	0.32		99		
Clostridium barati		1.03	0.05		99		
07653021	**Very good identification**						**07653021**
Actinomyces naeslundii		100.0	0.67	MLT 94			
07653022	**Low discrimination**						**07653022**
					SPO		
Actinomyces naeslundii		68.53	0.64	MLT 94	1		
Clostridium innocuum		23.27	0.32		99		
Lactobacillus acidophilus		3.96	0.18		1		
07653023	**Very good identification**						**07653023**
Actinomyces naeslundii		100.0	0.67	MLT 94			
07653030	**Low discrimination**						**07653030**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		78.82	0.42	MLT 94	1		
Clostridium innocuum		21.18	0.08	MLZ 5	99		
07653031	**Good identification**						**07653031**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07653032	**Low discrimination**						**07653032**
					SPO		
Actinomyces naeslundii		78.82	0.42	MLT 94	1		
Clostridium innocuum		21.18	0.08	MLZ 5	99		
07653033	**Good identification**						**07653033**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07653041	**Good identification**						**07653041**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07653043	**Good identification**						**07653043**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07653060	**Good identification**						**07653060**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07653061	**Good identification**						**07653061**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07653062	**Good identification**						**07653062**
Actinomyces naeslundii		96.11	0.51	MLT 94			
Lactobacillus acidophilus		3.89	0.03				
07653063	**Good identification**						**07653063**
Actinomyces naeslundii		100.0	0.54	MLT 94			
07653100	**Low discrimination**						**07653100**
					GRA	SPO	
Actinomyces naeslundii		61.06	0.72	MLT 94	99	1	
Fusobacterium mortiferum		29.72	0.55		1	1	
Clostridium innocuum		7.28	0.32		99	99	
07653101	**Very good identification**						**07653101**
					GRA		
Actinomyces naeslundii		99.67	0.75	MLT 94	99		
Fusobacterium mortiferum		0.33	0.21		1		
07653102	**Low discrimination**						**07653102**
					SPO		
Actinomyces naeslundii		67.19	0.72	MLT 94	1		
Lactobacillus acidophilus		15.70	0.36		1		
Clostridium innocuum		8.02	0.32		99		
07653103	**Very good identification**						**07653103**
Actinomyces naeslundii		99.84	0.75	MLT 94			
Lactobacillus acidophilus		0.16	0.02				
07653110	**Low discrimination**						**07653110**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		79.36	0.50	MLT 94	99	1	
Clostridium innocuum		7.49	0.08	MLZ 5	99	99	
Fusobacterium mortiferum		7.41	0.21		1	1	
07653111	**Good identification**						**07653111**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07653112	**Low discrimination**						**07653112**
Actinomyces naeslundii		53.51	0.50	MLT 94			
Bifidobacterium breve		18.84	0.27	MLZ 5			
Bifidobacterium adolescentis		17.19	0.26				
07653113	**Good identification**						**07653113**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07653120	**Low discrimination**						**07653120**
					SPO		
Actinomyces naeslundii		89.39	0.72	MLT 94	1		
Clostridium ramosum		8.01	0.28		92		
Clostridium barati		1.33	0.14		99		
07653121	**Very good identification**						**07653121**
Actinomyces naeslundii		100.0	0.75	MLT 94			
07653122	**Low discrimination**						**07653122**
					SPO		
Actinomyces naeslundii		58.79	0.72	MLT 94	1		
Clostridium ramosum		21.07	0.39		92		
Lactobacillus acidophilus		13.73	0.36		1		
07653123	**Very good identification**						**07653123**
					SPO		
Actinomyces naeslundii		99.60	0.75	MLT 94	1		
Clostridium ramosum		0.24	0.05		92		
Lactobacillus acidophilus		0.16	0.02		1		
07653130	**Acceptable identification**						**07653130**
Actinomyces naeslundii		94.17	0.50	MLT 94			
Bifidobacterium breve		5.83	0.15	MLZ 5			
07653131	**Good identification**						**07653131**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07653132	**Low discrimination**						**07653132**
					SPO		
Actinomyces naeslundii		55.68	0.50	MLT 94	1		
Bifidobacterium breve		34.86	0.32	MLZ 5	1		
Clostridium ramosum		3.83	0.05		92		
07653133	**Good identification**						**07653133**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07653140	**Good identification**						**07653140**
					GRA		
Actinomyces naeslundii		93.69	0.59	MLT 94	99		
Actinomyces israelii		2.72	0.23		99		
Fusobacterium mortiferum		2.61	0.21		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07653141	**Very good identification**					**07653141**
Actinomyces naeslundii		100.0	0.62	MLT 94		
07653142	**Low discrimination**					**07653142**
Actinomyces naeslundii		75.65	0.59	MLT 94		
Lactobacillus acidophilus		12.38	0.21			
Bifidobacterium breve		7.95	0.27			
07653143	**Very good identification**					**07653143**
Actinomyces naeslundii		100.0	0.62	MLT 94		
07653150	**Low discrimination**					**07653150**
Actinomyces naeslundii		83.52	0.37	MLT 94		
Bifidobacterium breve		16.48	0.10	MLZ 5		
07653151	**Good identification**					**07653151**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07653153	**Good identification**					**07653153**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07653160	**Very good identification**					**07653160**
Actinomyces naeslundii		98.19	0.59	MLT 94		
Bifidobacterium breve		1.81	0.15			
07653161	**Very good identification**					**07653161**
Actinomyces naeslundii		100.0	0.62	MLT 94		
07653162	**Low discrimination**					**07653162**
Actinomyces naeslundii		71.79	0.59	MLT 94		
Bifidobacterium breve		13.41	0.32			
Lactobacillus acidophilus		11.75	0.21			
07653163	**Very good identification**					**07653163**
Actinomyces naeslundii		100.0	0.62	MLT 94		
07653170	**Low discrimination**					**07653170**
Actinomyces naeslundii		74.03	0.37	MLT 94		
Bifidobacterium breve		25.97	0.15	MLZ 5		
07653171	**Good identification**					**07653171**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07653173	**Good identification**					**07653173**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07653300	**Low discrimination**					**07653300**
					GRA	
Actinomyces naeslundii		67.27	0.37	MLT 94	99	
Fusobacterium mortiferum		32.74	0.21	RHA 1	1	
07653301	**Good identification**					**07653301**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	MLT 94			
07653303	**Good identification**						**07653303**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07653321	**Good identification**						**07653321**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07653322	**Acceptable identification**						**07653322**
					SPO	GRA	
Clostridium ramosum		91.01	0.41	MLT 99	92	50	
Lactobacillus acidophilus		3.37	0.17	TRE 94	1	99	
Prevotella oralis		2.14	0.13		1	1	
07653323	**Low discrimination**						**07653323**
					SPO		
Actinomyces naeslundii		75.95	0.41	MLT 94	1		
Clostridium ramosum		24.05	0.07	RHA 1	92		
07653400	**Good identification**						**07653400**
					SPO		
Clostridium innocuum		99.60	0.53	RAF 1	99		
Actinomyces naeslundii		0.17	0.30		1		
Actinomyces israelii		0.13	0.18		1		
07653402	**Good identification**						**07653402**
					SPO		
Clostridium innocuum		99.64	0.53	RAF 1	99		
Actinomyces israelii		0.19	0.21		1		
Actinomyces naeslundii		0.17	0.30		1		
07653501	**Good identification**						**07653501**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07653502	**Low discrimination**						**07653502**
					SPO		
Bifidobacterium adolescentis		65.07	0.38	MLT 99	1		
Clostridium innocuum		14.79	0.19	XYL 87	99		
Actinomyces israelii		11.38	0.32		1		
07653503	**Good identification**						**07653503**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07653512	**Good identification**						**07653512**
Bifidobacterium adolescentis		97.91	0.40	MLT 99			
Actinomyces israelii		2.09	0.18	XYL 87			
07653520	**Good identification**						**07653520**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07653521	**Good identification**						**07653521**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07653522	**Low discrimination**						**07653522**
					SPO		
Actinomyces naeslundii		62.81	0.37	MLT 94	1		
Clostridium ramosum		22.51	0.05	ARA 1	92		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		14.68	0.02		1		
07653523	**Good identification**						**07653523**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07654000	**Very good identification**						**07654000**
Fusobacterium mortiferum		100.0	0.60	XYL 7			
07654002	**Good identification**						**07654002**
Fusobacterium mortiferum		100.0	0.44	XYL 7			
07654100	**Very good identification**						**07654100**
Fusobacterium mortiferum		100.0	0.67	XYL 7			
07654102	**Good identification**						**07654102**
Fusobacterium mortiferum		100.0	0.50	XYL 7			
07655000	**Very good identification**						**07655000**
					GRA	SPO	
Fusobacterium mortiferum		98.57	0.63	XYL 7	1	1	
Actinomyces israelii		0.43	0.25		99	1	
Clostridium innocuum		0.40	0.10		99	99	
07655002	**Low discrimination**						**07655002**
					GRA	SPO	
Fusobacterium mortiferum		85.93	0.47	XYL 7	1	1	
Actinomyces israelii		5.00	0.28	BGL 10	99	1	
Clostridium innocuum		3.12	0.10		99	99	
07655100	**Very good identification**						**07655100**
					GRA	SPO	
Fusobacterium mortiferum		98.18	0.70	XYL 7	1	1	
Actinomyces israelii		0.73	0.36		99	1	
Clostridium butyricum		0.71	0.14		99	99	
07655101	**Low discrimination**						**07655101**
					GRA		
Fusobacterium mortiferum		82.43	0.35	XYL 7	1		
Actinomyces naeslundii		17.57	0.36	URE 1	99		
07655102	**Low discrimination**						**07655102**
					GRA		
Fusobacterium mortiferum		61.34	0.53	XYL 7	1		
Bifidobacterium adolescentis		17.23	0.40	BGL 10	99		
Lactobacillus acidophilus		6.83	0.24		99		
07655103	**Low discrimination**						**07655103**
					GRA		
Actinomyces naeslundii		65.73	0.36	MLT 94	99		
Fusobacterium mortiferum		34.27	0.19	XYL 1	1		
07655110	**Low discrimination**						**07655110**
					GRA		
Fusobacterium mortiferum		84.40	0.35	XYL 7	1		
Actinomyces israelii		9.22	0.21	MLZ 1	99		
Bifidobacterium adolescentis		3.22	0.07		99		
07655112	**Low discrimination**						**07655112**
					GRA		
Bifidobacterium adolescentis		88.01	0.41	MLT 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		4.31	0.22	ARA 87	1		
Actinomyces israelii		3.82	0.24		99		
07655120	**Low discrimination**						**07655120**
Fusobacterium mortiferum		45.40	0.35	XYL 7			
Prevotella oralis		20.77	0.16	NAG 1			
Parabacteroides distasonis		15.59	0.26				
07655121	**Good identification**						**07655121**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07655122	**Low discrimination**						**07655122**
					GRA		
Parabacteroides distasonis		76.13	0.60		1		
Prevotella oralis		19.47	0.37		1		
Lactobacillus acidophilus		2.74	0.24		99		
07655132	**Good identification**						**07655132**
Parabacteroides distasonis		99.02	0.56				
Prevotella oralis		0.48	0.03				
Clostridium clostridioforme		0.30	0.07				
07655140	**Low discrimination**						**07655140**
					GRA	SPO	
Fusobacterium mortiferum		74.29	0.35	XYL 7	1	1	
Actinomyces israelii		21.12	0.29	SOR 1	99	1	
Clostridium butyricum		4.60	-.04		99	99	
07655300	**Acceptable identification**						**07655300**
					GRA		
Fusobacterium mortiferum		94.79	0.35	XYL 7	1		
Actinomyces israelii		5.22	0.16	RHA 1	99		
07655322	**Very good identification**						**07655322**
Parabacteroides distasonis		96.19	0.68				
Prevotella oralis		3.51	0.31				
Bacteroides uniformis		0.13	0.01				
07655332	**Very good identification**						**07655332**
Parabacteroides distasonis		99.21	0.64				
Clostridium clostridioforme		0.73	0.22				
Prevotella oralis		0.07	-.03				
07655502	**Low discrimination**						**07655502**
					SPO		
Bifidobacterium adolescentis		84.14	0.54	MLT 99	1		
Clostridium butyricum		10.14	0.24		99		
Actinomyces israelii		3.65	0.37		1		
07655512	**Good identification**						**07655512**
Bifidobacterium adolescentis		95.34	0.56	MLT 99			
Clostridium clostridioforme		3.23	0.27				
Actinomyces israelii		0.51	0.23				
07655522	**Good identification**						**07655522**
					GRA	SPO	
Parabacteroides distasonis		91.57	0.57		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides uniformis		5.10	0.16		1	1	
Clostridium butyricum		0.95	0.05		99	99	
07655532	**Acceptable identification**						**07655532**
					GRA		
Parabacteroides distasonis		90.98	0.52		1		
Clostridium clostridioforme		6.95	0.27		50		
Bifidobacterium adolescentis		2.07	0.21		99		
07655542	**Low discrimination**						**07655542**
					SPO		
Bifidobacterium adolescentis		81.88	0.39	MLT 99	1		
Actinomyces israelii		11.18	0.30	SOR 11	1		
Clostridium butyricum		6.94	0.05		99		
07655552	**Good identification**						**07655552**
Bifidobacterium adolescentis		98.36	0.40	MLT 99			
Actinomyces israelii		1.64	0.16	SOR 11			
07655712	**Acceptable identification**						**07655712**
					GRA		
Clostridium clostridioforme		91.35	0.42	MLT 99	50		
Parabacteroides distasonis		4.94	0.26	TRE 88	1		
Bifidobacterium adolescentis		3.71	0.21		99		
07655722	**Good identification**						**07655722**
Parabacteroides distasonis		94.99	0.65				
Bacteroides uniformis		2.44	0.19				
Clostridium clostridioforme		2.10	0.31				
07655732	**Low discrimination**						**07655732**
Parabacteroides distasonis		84.00	0.60				
Clostridium clostridioforme		15.69	0.42				
Bacteroides ovatus		0.31	0.03				
07657000	**Acceptable identification**						**07657000**
					GRA	SPO	
Clostridium innocuum		91.58	0.44	RAF 1	99	99	
Actinomyces israelii		5.59	0.38	XYL 5	99	1	
Fusobacterium mortiferum		2.31	0.29		1	1	
07657002	**Acceptable identification**						**07657002**
					SPO		
Clostridium innocuum		90.91	0.44	RAF 1	99		
Actinomyces israelii		8.32	0.41	XYL 5	1		
Actinomyces naeslundii		0.52	0.30		1		
07657100	**Low discrimination**						**07657100**
					GRA		
Actinomyces israelii		73.07	0.48	MLT 99	99		
Fusobacterium mortiferum		17.64	0.35		1		
Actinomyces naeslundii		4.87	0.37		99		
07657101	**Acceptable identification**						**07657101**
Actinomyces naeslundii		90.81	0.41	MLT 94			
Actinomyces israelii		9.19	0.14	XYL 1			
07657102	**Low discrimination**						**07657102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		66.35	0.51	MLT 99		
Bifidobacterium adolescentis		27.00	0.38			
Actinomyces naeslundii		2.95	0.37			
07657103	**Low discrimination**					**07657103**
Actinomyces naeslundii		86.83	0.41	MLT 94		
Actinomyces israelii		13.17	0.17	XYL 1		
07657112	**Low discrimination**					**07657112**
					SPO	
Bifidobacterium adolescentis		76.19	0.40	MLT 99	1	
Actinomyces israelii		22.89	0.37	ARA 87	1	
Clostridium tertium		0.92	-.01		99	
07657120	**Low discrimination**					**07657120**
Actinomyces naeslundii		51.05	0.37	MLT 94		
Actinomyces israelii		48.95	0.28	XYL 1		
07657121	**Good identification**					**07657121**
Actinomyces naeslundii		100.0	0.41	MLT 94		
07657123	**Good identification**					**07657123**
Actinomyces naeslundii		100.0	0.41	MLT 94		
07657140	**Good identification**					**07657140**
Actinomyces israelii		97.07	0.41	MLT 99		
Actinomyces naeslundii		2.93	0.25			
07657142	**Low discrimination**					**07657142**
Actinomyces israelii		87.00	0.44	MLT 99		
Bifidobacterium adolescentis		11.25	0.23			
Actinomyces naeslundii		1.75	0.25			
07657500	**Low discrimination**					**07657500**
					SPO	
Actinomyces israelii		79.34	0.47	MLT 99	1	
Clostridium butyricum		16.66	0.14		99	
Bifidobacterium adolescentis		4.00	0.18		1	
07657502	**Low discrimination**					**07657502**
					SPO	
Bifidobacterium adolescentis		75.94	0.53	MLT 99	1	
Actinomyces israelii		22.82	0.50		1	
Clostridium butyricum		1.24	0.07		99	
07657512	**Good identification**					**07657512**
Bifidobacterium adolescentis		96.16	0.54	MLT 99		
Actinomyces israelii		3.53	0.36			
Bifidobacterium dentium		0.30	-.07			
07657540	**Good identification**					**07657540**
					SPO	
Actinomyces israelii		95.52	0.40	MLT 99	1	
Clostridium butyricum		4.49	-.04		99	
07657542	**Low discrimination**					**07657542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		51.40	0.37	MLT 99		
Actinomyces israelii		48.60	0.43	SOR 11		
07657552	**Low discrimination**					**07657552**
Bifidobacterium adolescentis		89.64	0.39	MLT 99		
Actinomyces israelii		10.36	0.29	SOR 11		
07660000	**Good identification**					**07660000**
Fusobacterium mortiferum		100.0	0.36	FRU 99		
07660100	**Good identification**					**07660100**
					GRA	
Fusobacterium mortiferum		97.79	0.43	FRU 99	1	
Actinomyces naeslundii		2.21	0.26		99	
07661000	**Low discrimination**					**07661000**
					GRA	
Fusobacterium mortiferum		71.25	0.39	FRU 99	1	
Actinomyces naeslundii		21.15	0.42		99	
Lactobacillus jensenii		5.08	0.10		99	
07661001	**Good identification**					**07661001**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07661002	**Low discrimination**					**07661002**
					GRA	
Actinomyces naeslundii		42.38	0.42	MLT 94	99	
Lactobacillus acidophilus		29.92	0.15	FRU 95	99	
Fusobacterium mortiferum		15.86	0.23		1	
07661003	**Good identification**					**07661003**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07661020	**Good identification**					**07661020**
Actinomyces naeslundii		100.0	0.42	MLT 94		
07661021	**Good identification**					**07661021**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07661022	**Low discrimination**					**07661022**
Actinomyces naeslundii		58.62	0.42	MLT 94		
Lactobacillus acidophilus		41.38	0.15	FRU 95		
07661023	**Good identification**					**07661023**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07661100	**Low discrimination**					**07661100**
					GRA	
Fusobacterium mortiferum		72.86	0.46	FRU 99	1	
Actinomyces naeslundii		26.38	0.50		99	
Lactobacillus acidophilus		0.76	-.01		99	
07661101	**Good identification**					**07661101**
					GRA	
Actinomyces naeslundii		98.18	0.53	MLT 94	99	
Fusobacterium mortiferum		1.83	0.12	FRU 95	1	
07661103	**Good identification**					**07661103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.12	0.53	MLT 94		
Lactobacillus acidophilus		1.89	-.01	FRU 95		
07661120	**Acceptable identification**					**07661120**
					SPO	
Actinomyces naeslundii		90.56	0.50	MLT 94	1	
Clostridium ramosum		4.31	0.02	FRU 95	92	
Lactobacillus acidophilus		2.61	-.01		1	
07661121	**Good identification**					**07661121**
Actinomyces naeslundii		100.0	0.53	MLT 94		
07661123	**Good identification**					**07661123**
Actinomyces naeslundii		98.12	0.53	MLT 94		
Lactobacillus acidophilus		1.89	-.01	FRU 95		
07661140	**Low discrimination**					**07661140**
					GRA	
Actinomyces naeslundii		86.35	0.37	MLT 94	99	
Fusobacterium mortiferum		13.65	0.12	FRU 95	1	
07661141	**Good identification**					**07661141**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07661143	**Good identification**					**07661143**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07661160	**Good identification**					**07661160**
Actinomyces naeslundii		100.0	0.37	MLT 94		
07661161	**Good identification**					**07661161**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07661163	**Good identification**					**07661163**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07663000	**Low discrimination**					**07663000**
					SPO	
Clostridium innocuum		85.88	0.36	RAF 1	99	
Actinomyces naeslundii		14.12	0.47	FRU 95	1	
07663001	**Good identification**					**07663001**
					SPO	
Actinomyces naeslundii		96.07	0.50	MLT 94	1	
Clostridium innocuum		3.94	0.02	FRU 95	99	
07663002	**Low discrimination**					**07663002**
					SPO	
Clostridium innocuum		85.88	0.36	RAF 1	99	
Actinomyces naeslundii		14.12	0.47	FRU 95	1	
07663003	**Good identification**					**07663003**
					SPO	
Actinomyces naeslundii		96.07	0.50	MLT 94	1	
Clostridium innocuum		3.94	0.02	FRU 95	99	
07663020	**Acceptable identification**					**07663020**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		94.21	0.47	MLT 94	1		
Clostridium innocuum		5.79	0.02	FRU 95	99		
07663021	**Good identification**						**07663021**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07663022	**Acceptable identification**						**07663022**
					SPO		
Actinomyces naeslundii		94.21	0.47	MLT 94	1		
Clostridium innocuum		5.79	0.02	FRU 95	99		
07663023	**Good identification**						**07663023**
Actinomyces naeslundii		100.0	0.50	MLT 94			
07663041	**Good identification**						**07663041**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07663043	**Good identification**						**07663043**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07663061	**Good identification**						**07663061**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07663063	**Good identification**						**07663063**
Actinomyces naeslundii		100.0	0.37	MLT 94			
07663100	**Good identification**						**07663100**
					GRA	SPO	
Actinomyces naeslundii		96.53	0.55	MLT 94	99	1	
Clostridium innocuum		2.08	0.02	FRU 95	99	99	
Fusobacterium mortiferum		1.39	0.12		1	1	
07663101	**Very good identification**						**07663101**
Actinomyces naeslundii		100.0	0.58	MLT 94			
07663102	**Acceptable identification**						**07663102**
					SPO		
Actinomyces naeslundii		91.00	0.55	MLT 94	1		
Lactobacillus acidophilus		7.04	0.11	FRU 95	1		
Clostridium innocuum		1.96	0.02		99		
07663103	**Very good identification**						**07663103**
Actinomyces naeslundii		100.0	0.58	MLT 94			
07663111	**Good identification**						**07663111**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07663113	**Good identification**						**07663113**
Actinomyces naeslundii		100.0	0.36	MLT 94			
07663120	**Low discrimination**						**07663120**
					SPO		
Actinomyces naeslundii		87.81	0.55	MLT 94	1		
Clostridium ramosum		12.19	0.15	FRU 95	92		
07663121	**Very good identification**						**07663121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.58	MLT 94		
07663122	**Low discrimination**					**07663122**
					SPO	
Actinomyces naeslundii		61.26	0.55	MLT 94	1	
Clostridium ramosum		34.01	0.25	FRU 95	92	
Lactobacillus acidophilus		4.74	0.11		1	
07663123	**Very good identification**					**07663123**
Actinomyces naeslundii		100.0	0.58	MLT 94		
07663131	**Good identification**					**07663131**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07663133	**Good identification**					**07663133**
Actinomyces naeslundii		100.0	0.36	MLT 94		
07663140	**Good identification**					**07663140**
Actinomyces naeslundii		100.0	0.42	MLT 94		
07663141	**Good identification**					**07663141**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07663142	**Good identification**					**07663142**
Actinomyces naeslundii		100.0	0.42	MLT 94		
07663143	**Good identification**					**07663143**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07663160	**Good identification**					**07663160**
Actinomyces naeslundii		100.0	0.42	MLT 94		
07663161	**Good identification**					**07663161**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07663162	**Good identification**					**07663162**
Actinomyces naeslundii		100.0	0.42	MLT 94		
07663163	**Good identification**					**07663163**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07665122	**Good identification**					**07665122**
					GRA	
Parabacteroides distasonis		97.42	0.37	FRU 99	1	
Lactobacillus acidophilus		2.58	-.01		99	
07665322	**Good identification**					**07665322**
Parabacteroides distasonis		100.0	0.46	FRU 99		
07665332	**Good identification**					**07665332**
Parabacteroides distasonis		100.0	0.41	FRU 99		
07665722	**Good identification**					**07665722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		96.57	0.42	FRU 99			
Clostridium clostridioforme		3.43	0.11				
07665732	**Low discrimination**						**07665732**
Parabacteroides distasonis		76.89	0.37	FRU 99			
Clostridium clostridioforme		23.11	0.23				
07670000	**Very good identification**						**07670000**
					GRA	CAT	
Fusobacterium mortiferum		98.67	0.71		1	5	
Propionibacterium granulosum		0.82	0.32		99	90	
Actinomyces naeslundii		0.35	0.40		99	25	
07670001	**Low discrimination**						**07670001**
					GRA		
Fusobacterium mortiferum		65.41	0.36	URE 1	1		
Actinomyces naeslundii		34.59	0.43		99		
07670002	**Acceptable identification**						**07670002**
					GRA		
Fusobacterium mortiferum		91.68	0.54	BGL 10	1		
Actinomyces naeslundii		2.94	0.40		99		
Lactobacillus acidophilus		2.62	0.15		99		
07670003	**Low discrimination**						**07670003**
					GRA		
Actinomyces naeslundii		82.63	0.43	MLT 94	99		
Fusobacterium mortiferum		17.37	0.20	GAL 96	1		
07670010	**Low discrimination**						**07670010**
					GRA	CAT	
Fusobacterium mortiferum		87.08	0.36	MLZ 1	1	5	
Propionibacterium granulosum		9.82	0.17		99	90	
Lactobacillus jensenii		3.11	0.02		99	1	
07670020	**Low discrimination**						**07670020**
					GRA	CAT	
Fusobacterium mortiferum		64.14	0.36	NAG 1	1	5	
Actinomyces naeslundii		22.61	0.40		99	25	
Propionibacterium granulosum		13.26	0.21		99	90	
07670021	**Good identification**						**07670021**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07670022	**Low discrimination**						**07670022**
					GRA		
Actinomyces naeslundii		42.49	0.40	MLT 94	99		
Lactobacillus acidophilus		37.89	0.15	GAL 96	99		
Fusobacterium mortiferum		13.39	0.20		1		
07670023	**Good identification**						**07670023**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07670040	**Acceptable identification**						**07670040**
					GRA		
Fusobacterium mortiferum		94.14	0.36	SOR 1	1		
Actinomyces naeslundii		5.86	0.27		99		
07670100	**Very good identification**						**07670100**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		99.55	0.77		1		
Actinomyces naeslundii		0.43	0.48		99		
Lactobacillus acidophilus		0.02	-.01		99		
07670101	**Low discrimination**						**07670101**
					GRA		
Fusobacterium mortiferum		60.79	0.43	URE 1	1		
Actinomyces naeslundii		39.21	0.51		99		
07670102	**Low discrimination**						**07670102**
					GRA		
Fusobacterium mortiferum		84.75	0.61	BGL 10	1		
Lactobacillus acidophilus		11.94	0.33		99		
Actinomyces naeslundii		3.31	0.48		99		
07670103	**Low discrimination**						**07670103**
					GRA		
Actinomyces naeslundii		83.58	0.51	MLT 94	99		
Fusobacterium mortiferum		14.40	0.26	GAL 96	1		
Lactobacillus acidophilus		2.03	-.01		99		
07670110	**Good identification**						**07670110**
					GRA		
Fusobacterium mortiferum		97.79	0.43	MLZ 1	1		
Actinomyces naeslundii		2.21	0.26		99		
07670120	**Low discrimination**						**07670120**
					SPO	GRA	
Fusobacterium mortiferum		67.96	0.43	NAG 1	1	1	
Actinomyces naeslundii		29.22	0.48		1	99	
Clostridium ramosum		1.76	0.02		92	50	
07670121	**Good identification**						**07670121**
Actinomyces naeslundii		100.0	0.51	MLT 94			
07670123	**Good identification**						**07670123**
Actinomyces naeslundii		97.63	0.51	MLT 94			
Lactobacillus acidophilus		2.37	-.01	GAL 96			
07670140	**Acceptable identification**						**07670140**
					GRA		
Fusobacterium mortiferum		92.95	0.43	SOR 1	1		
Actinomyces naeslundii		7.05	0.35		99		
07670141	**Good identification**						**07670141**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07670143	**Good identification**						**07670143**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07670160	**Good identification**						**07670160**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07670161	**Good identification**						**07670161**
Actinomyces naeslundii		100.0	0.38	MLT 94			
07670163	**Good identification**						**07670163**
Actinomyces naeslundii		100.0	0.38	MLT 94			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07670200	**Good identification**					**07670200**
Fusobacterium mortiferum		100.0	0.36	RHA 1		
07670300	**Good identification**					**07670300**
Fusobacterium mortiferum		100.0	0.43	RHA 1		
07670400	**Good identification**					**07670400**
Fusobacterium mortiferum		100.0	0.36	ARA 1		
07670500	**Good identification**					**07670500**
Fusobacterium mortiferum		100.0	0.43	ARA 1		
07671000	**Low discrimination**					**07671000**
					GRA	
Fusobacterium mortiferum		87.38	0.74		1	
Lactobacillus jensenii		6.23	0.44		99	
Actinomyces naeslundii		4.98	0.64		99	
07671001	**Low discrimination**					**07671001**
					GRA	
Actinomyces naeslundii		88.76	0.67	MLT 94	99	
Fusobacterium mortiferum		10.49	0.39		1	
Lactobacillus jensenii		0.75	0.10		99	
07671002	**Low discrimination**					**07671002**
					GRA	
Lactobacillus acidophilus		48.09	0.49	MLT 94	99	
Fusobacterium mortiferum		25.50	0.57	LAC 92	1	
Actinomyces naeslundii		13.08	0.64		99	
07671003	**Very good identification**					**07671003**
					GRA	
Actinomyces naeslundii		95.83	0.67	MLT 94	99	
Lactobacillus acidophilus		2.37	0.15		99	
Fusobacterium mortiferum		1.26	0.23		1	
07671010	**Low discrimination**					**07671010**
					GRA	
Lactobacillus jensenii		62.70	0.36	MLT 95	99	
Fusobacterium mortiferum		26.65	0.39	RAF 1	1	
Actinomyces naeslundii		7.91	0.42		99	
07671011	**Acceptable identification**					**07671011**
Actinomyces naeslundii		94.93	0.45	MLT 94		
Lactobacillus jensenii		5.07	0.02	MLZ 5		
07671013	**Good identification**					**07671013**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07671020	**Low discrimination**					**07671020**
					GRA	
Actinomyces naeslundii		63.11	0.64	MLT 94	99	
Lactobacillus jensenii		19.75	0.34		99	
Fusobacterium mortiferum		11.19	0.39		1	
07671021	**Very good identification**					**07671021**
Actinomyces naeslundii		100.0	0.67	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07671022	**Low discrimination**					**07671022**
Lactobacillus acidophilus		73.04	0.49	MLT 94		
Actinomyces naeslundii		19.86	0.64	LAC 92		
Lactobacillus jensenii		4.14	0.31			
07671023	**Very good identification**					**07671023**
Actinomyces naeslundii		97.58	0.67	MLT 94		
Lactobacillus acidophilus		2.42	0.15			
07671031	**Good identification**					**07671031**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07671033	**Good identification**					**07671033**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07671040	**Low discrimination**					**07671040**
					GRA	
Fusobacterium mortiferum		47.79	0.39	SOR 1	1	
Actinomyces naeslundii		47.56	0.51		99	
Lactobacillus jensenii		3.41	0.10		99	
07671041	**Good identification**					**07671041**
Actinomyces naeslundii		100.0	0.54	MLT 94		
07671043	**Good identification**					**07671043**
Actinomyces naeslundii		98.30	0.54	MLT 94		
Lactobacillus acidophilus		1.71	-.01			
07671060	**Good identification**					**07671060**
Actinomyces naeslundii		97.46	0.51	MLT 94		
Lactobacillus acidophilus		2.54	-.01			
07671061	**Good identification**					**07671061**
Actinomyces naeslundii		100.0	0.54	MLT 94		
07671063	**Good identification**					**07671063**
Actinomyces naeslundii		98.30	0.54	MLT 94		
Lactobacillus acidophilus		1.71	-.01			
07671100	**Good identification**					**07671100**
					GRA	
Fusobacterium mortiferum		92.22	0.80		1	
Actinomyces naeslundii		6.41	0.72		99	
Lactobacillus acidophilus		0.96	0.33		99	
07671101	**Good identification**					**07671101**
					GRA	
Actinomyces naeslundii		91.08	0.75	MLT 94	99	
Fusobacterium mortiferum		8.83	0.46		1	
Lactobacillus acidophilus		0.09	-.01		99	
07671102	**Low discrimination**					**07671102**
					GRA	
Lactobacillus acidophilus		84.63	0.67	MLT 94	99	
Fusobacterium mortiferum		9.11	0.64		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		5.70	0.72		99		
07671103	**Good identification**						**07671103**
					GRA		
Actinomyces naeslundii		90.02	0.75	MLT 94	99		
Lactobacillus acidophilus		9.01	0.33		99		
Fusobacterium mortiferum		0.97	0.29		1		
07671110	**Low discrimination**						**07671110**
					GRA		
Fusobacterium mortiferum		71.77	0.46	MLZ 1	1		
Actinomyces naeslundii		25.99	0.50		99		
Leptotrichia buccalis		0.77	-.11		1		
07671111	**Good identification**						**07671111**
					GRA		
Actinomyces naeslundii		98.18	0.53	MLT 94	99		
Fusobacterium mortiferum		1.83	0.12	MLZ 5	1		
07671113	**Good identification**						**07671113**
					GRA		
Actinomyces naeslundii		96.25	0.53	MLT 94	99		
Leptotrichia buccalis		1.90	-.11	MLZ 5	1		
Lactobacillus acidophilus		1.85	-.01		99		
07671120	**Low discrimination**						**07671120**
					SPO		
Actinomyces naeslundii		62.00	0.72	MLT 94	1		
Clostridium ramosum		15.36	0.36		92		
Lactobacillus acidophilus		9.31	0.33		1		
07671121	**Very good identification**						**07671121**
					SPO	GRA	
Actinomyces naeslundii		99.64	0.75	MLT 94	1	99	
Clostridium ramosum		0.17	0.02		92	50	
Fusobacterium mortiferum		0.10	0.12		1	1	
07671122	**Low discrimination**						**07671122**
					SPO		
Lactobacillus acidophilus		83.76	0.67	MLT 94	1		
Actinomyces naeslundii		5.64	0.72		1		
Clostridium ramosum		5.58	0.46		92		
07671123	**Low discrimination**						**07671123**
					SPO		
Actinomyces naeslundii		89.92	0.75	MLT 94	1		
Lactobacillus acidophilus		9.00	0.33		1		
Clostridium ramosum		0.60	0.12		92		
07671130	**Low discrimination**						**07671130**
					SPO	GRA	
Actinomyces naeslundii		82.60	0.50	MLT 94	1	99	
Parabacteroides distasonis		5.38	0.16	MLZ 5	1	1	
Clostridium ramosum		3.93	0.02		92	50	
07671131	**Good identification**						**07671131**
Actinomyces naeslundii		100.0	0.53	MLT 94			
07671132	**Low discrimination**						**07671132**
					GRA		
Parabacteroides distasonis		57.10	0.51	XYL 90	1		
Lactobacillus acidophilus		25.26	0.33		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		8.85	0.50		99	
07671133	**Acceptable identification**					**07671133**
					GRA	
Actinomyces naeslundii		94.11	0.53	MLT 94	99	
Parabacteroides distasonis		4.09	0.16	MLZ 5	1	
Lactobacillus acidophilus		1.81	-.01		99	
07671140	**Low discrimination**					**07671140**
					GRA	
Actinomyces naeslundii		51.33	0.59	MLT 94	99	
Fusobacterium mortiferum		42.28	0.46		1	
Lactobacillus acidophilus		5.40	0.18		99	
07671141	**Very good identification**					**07671141**
					GRA	
Actinomyces naeslundii		99.45	0.62	MLT 94	99	
Fusobacterium mortiferum		0.55	0.12		1	
07671142	**Low discrimination**					**07671142**
					GRA	
Lactobacillus acidophilus		89.68	0.52	MLT 94	99	
Actinomyces naeslundii		8.62	0.59	SOR 11	99	
Fusobacterium mortiferum		0.79	0.29		1	
07671143	**Good identification**					**07671143**
Actinomyces naeslundii		93.45	0.62	MLT 94		
Lactobacillus acidophilus		6.55	0.18			
07671150	**Low discrimination**					**07671150**
					GRA	
Actinomyces naeslundii		86.35	0.37	MLT 94	99	
Fusobacterium mortiferum		13.65	0.12	MLZ 5	1	
07671151	**Good identification**					**07671151**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07671153	**Good identification**					**07671153**
Actinomyces naeslundii		100.0	0.40	MLT 94		
07671160	**Low discrimination**					**07671160**
Actinomyces naeslundii		86.22	0.59	MLT 94		
Lactobacillus acidophilus		9.07	0.18			
Clostridium perfringens		1.71	0.07			
07671161	**Very good identification**					**07671161**
Actinomyces naeslundii		100.0	0.62	MLT 94		
07671162	**Low discrimination**					**07671162**
Lactobacillus acidophilus		89.22	0.52	MLT 94		
Actinomyces naeslundii		8.57	0.59	SOR 11		
Bifidobacterium breve		1.07	0.29			
07671163	**Good identification**					**07671163**
Actinomyces naeslundii		93.45	0.62	MLT 94		
Lactobacillus acidophilus		6.55	0.18			
07671170	**Low discrimination**					**07671170**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		81.04	0.37	MLT 94			
Bifidobacterium breve		18.96	0.12	MLZ 5			
07671171	**Good identification**						**07671171**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07671173	**Good identification**						**07671173**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07671200	**Low discrimination**						**07671200**
					GRA		
Fusobacterium mortiferum		88.63	0.39	RHA 1	1		
Lactobacillus jensenii		6.32	0.10		99		
Actinomyces naeslundii		5.05	0.30		99		
07671300	**Low discrimination**						**07671300**
					GRA		
Fusobacterium mortiferum		85.47	0.46	RHA 1	1		
Lactobacillus acidophilus		6.64	0.14		99		
Actinomyces naeslundii		5.94	0.37		99		
07671301	**Acceptable identification**						**07671301**
					GRA		
Actinomyces naeslundii		91.17	0.41	MLT 94	99		
Fusobacterium mortiferum		8.83	0.12	RHA 1	1		
07671302	**Acceptable identification**						**07671302**
					GRA		
Lactobacillus acidophilus		94.54	0.48	MLT 94	99		
Parabacteroides distasonis		1.61	0.29	RHA 7	1		
Fusobacterium mortiferum		1.37	0.29		1		
07671303	**Low discrimination**						**07671303**
Actinomyces naeslundii		57.28	0.41	MLT 94			
Lactobacillus acidophilus		42.72	0.14	RHA 1			
07671320	**Low discrimination**						**07671320**
					SPO	GRA	
Clostridium ramosum		87.79	0.38	MLT 99	92	50	
Parabacteroides distasonis		5.11	0.29		1	1	
Lactobacillus acidophilus		3.02	0.14		1	99	
07671321	**Low discrimination**						**07671321**
					SPO		
Actinomyces naeslundii		82.04	0.41	MLT 94	1		
Clostridium ramosum		17.96	0.04	RHA 1	92		
07671322	**Low discrimination**						**07671322**
					SPO	GRA	
Parabacteroides distasonis		42.88	0.63	XYL 90	1	1	
Clostridium ramosum		29.79	0.48		92	50	
Lactobacillus acidophilus		25.37	0.48		1	99	
07671332	**Very good identification**						**07671332**
					SPO	GRA	
Parabacteroides distasonis		96.66	0.59	XYL 90	1	1	
Clostridium ramosum		1.26	0.14		92	50	
Lactobacillus acidophilus		1.07	0.14		1	99	
07671400	**Low discrimination**						**07671400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		73.98	0.39	ARA 1	1	1	
Clostridium innocuum		8.78	0.11		99	99	
Clostridium butyricum		5.63	0.02		99	99	
07671500	**Low discrimination**						**07671500**
					GRA	SPO	
Fusobacterium mortiferum		52.32	0.46	ARA 1	1	1	
Clostridium butyricum		40.92	0.25		99	99	
Actinomyces naeslundii		3.64	0.37		99	1	
07671501	**Acceptable identification**						**07671501**
					GRA		
Actinomyces naeslundii		91.17	0.41	MLT 94	99		
Fusobacterium mortiferum		8.83	0.12	ARA 1	1		
07671503	**Acceptable identification**						**07671503**
Actinomyces naeslundii		90.90	0.41	MLT 94			
Lactobacillus acidophilus		9.10	-.01	ARA 1			
07671512	**Low discrimination**						**07671512**
					SPO		
Bifidobacterium adolescentis		74.31	0.38	MLT 99	1		
Clostridium clostridioforme		18.35	0.25	XYL 87	66		
Clostridium butyricum		1.88	-.04		99		
07671520	**Low discrimination**						**07671520**
					GRA	SPO	
Actinomyces naeslundii		34.38	0.37	MLT 94	99	1	
Clostridium butyricum		33.66	0.07	ARA 1	99	99	
Parabacteroides distasonis		13.28	0.17		1	1	
07671521	**Good identification**						**07671521**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07671522	**Low discrimination**						**07671522**
					GRA		
Parabacteroides distasonis		68.13	0.52	XYL 90	1		
Lactobacillus acidophilus		26.47	0.33		99		
Actinomyces naeslundii		1.78	0.37		99		
07671523	**Low discrimination**						**07671523**
					GRA		
Actinomyces naeslundii		73.66	0.41	MLT 94	99		
Parabacteroides distasonis		18.97	0.17	ARA 1	1		
Lactobacillus acidophilus		7.37	-.01		99		
07671532	**Low discrimination**						**07671532**
					GRA		
Parabacteroides distasonis		89.55	0.47	XYL 90	1		
Clostridium clostridioforme		9.80	0.25		50		
Lactobacillus acidophilus		0.65	-.01		99		
07671712	**Good identification**						**07671712**
Clostridium clostridioforme		96.36	0.40	MLT 99			
Parabacteroides distasonis		3.64	0.21	XYL 91			
07671722	**Good identification**						**07671722**
					SPO		
Parabacteroides distasonis		94.98	0.60	XYL 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		3.00	0.28		66		
Clostridium ramosum		1.09	0.14		92		
07671732	**Low discrimination**						**07671732**
Parabacteroides distasonis		78.89	0.55	XYL 90			
Clostridium clostridioforme		21.11	0.40				
07672000	**Low discrimination**						**07672000**
					GRA	SPO	
Clostridium innocuum		69.50	0.36	RAF 1	99	99	
Fusobacterium mortiferum		13.22	0.36	GAL 95	1	1	
Actinomyces naeslundii		9.04	0.45		99	1	
07672001	**Good identification**						**07672001**
					SPO		
Actinomyces naeslundii		95.08	0.48	MLT 94	1		
Clostridium innocuum		4.92	0.02	GAL 96	99		
07672002	**Low discrimination**						**07672002**
					SPO	CAT	
Clostridium innocuum		84.68	0.36	RAF 1	99	1	
Actinomyces naeslundii		11.02	0.45	GAL 95	1	25	
Propionibacterium granulosum		2.51	0.19		1	90	
07672003	**Good identification**						**07672003**
					SPO		
Actinomyces naeslundii		95.08	0.48	MLT 94	1		
Clostridium innocuum		4.92	0.02	GAL 96	99		
07672020	**Low discrimination**						**07672020**
					SPO	CAT	
Actinomyces naeslundii		76.60	0.45	MLT 94	1	25	
Propionibacterium granulosum		17.46	0.19	GAL 96	1	90	
Clostridium innocuum		5.95	0.02		99	1	
07672021	**Good identification**						**07672021**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07672022	**Low discrimination**						**07672022**
					SPO	CAT	
Actinomyces naeslundii		88.14	0.45	MLT 94	1	25	
Clostridium innocuum		6.84	0.02	GAL 96	99	1	
Propionibacterium granulosum		5.02	0.09		1	90	
07672023	**Good identification**						**07672023**
Actinomyces naeslundii		100.0	0.48	MLT 94			
07672041	**Good identification**						**07672041**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07672043	**Good identification**						**07672043**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07672061	**Good identification**						**07672061**
Actinomyces naeslundii		100.0	0.35	MLT 94			
07672063	**Good identification**						**07672063**
Actinomyces naeslundii		100.0	0.35	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07672100	**Low discrimination**						**07672100**
					GRA	SPO	
Fusobacterium mortiferum		53.84	0.43	MAN 1	1	1	
Actinomyces naeslundii		44.94	0.53		99	1	
Clostridium innocuum		1.23	0.02		99	99	
07672101	**Good identification**						**07672101**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07672102	**Low discrimination**						**07672102**
					GRA		
Actinomyces naeslundii		79.49	0.53	MLT 94	99		
Fusobacterium mortiferum		10.58	0.26	GAL 96	1		
Lactobacillus acidophilus		7.76	0.11		99		
07672103	**Good identification**						**07672103**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07672120	**Low discrimination**						**07672120**
					SPO		
Actinomyces naeslundii		85.08	0.53	MLT 94	1		
Clostridium ramosum		14.92	0.15	GAL 96	92		
07672121	**Good identification**						**07672121**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07672122	**Low discrimination**						**07672122**
					SPO		
Actinomyces naeslundii		55.59	0.53	MLT 94	1		
Clostridium ramosum		38.98	0.25	GAL 96	92		
Lactobacillus acidophilus		5.43	0.11		1		
07672123	**Good identification**						**07672123**
Actinomyces naeslundii		100.0	0.56	MLT 94			
07672140	**Good identification**						**07672140**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07672141	**Good identification**						**07672141**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07672142	**Good identification**						**07672142**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07672143	**Good identification**						**07672143**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07672160	**Good identification**						**07672160**
Actinomyces naeslundii		100.0	0.40	MLT 94			
07672161	**Good identification**						**07672161**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07672162	**Good identification**						**07672162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.40	MLT 94			
07672163	**Good identification**						**07672163**
Actinomyces naeslundii		100.0	0.43	MLT 94			
07673000	**Low discrimination**						**07673000**
					GRA	SPO	
Clostridium innocuum		84.06	0.58	RAF 1	99	99	
Actinomyces naeslundii		13.82	0.69		99	1	
Fusobacterium mortiferum		1.26	0.39		1	1	
07673001	**Very good identification**						**07673001**
					SPO		
Actinomyces naeslundii		96.06	0.72	MLT 94	1		
Clostridium innocuum		3.94	0.24		99		
07673002	**Low discrimination**						**07673002**
					SPO		
Clostridium innocuum		84.09	0.58	RAF 1	99		
Actinomyces naeslundii		13.82	0.69		1		
Lactobacillus acidophilus		1.38	0.27		1		
07673003	**Very good identification**						**07673003**
					SPO		
Actinomyces naeslundii		96.06	0.72	MLT 94	1		
Clostridium innocuum		3.94	0.24		99		
07673011	**Good identification**						**07673011**
					SPO		
Actinomyces naeslundii		96.86	0.50	MLT 94	1		
Clostridium innocuum		3.14	0.00	MLZ 5	99		
07673013	**Good identification**						**07673013**
					SPO		
Actinomyces naeslundii		96.86	0.50	MLT 94	1		
Clostridium innocuum		3.14	0.00	MLZ 5	99		
07673020	**Good identification**						**07673020**
					SPO		
Actinomyces naeslundii		91.60	0.69	MLT 94	1		
Clostridium innocuum		5.63	0.24		99		
Clostridium ramosum		1.90	0.15		92		
07673021	**Very good identification**						**07673021**
Actinomyces naeslundii		100.0	0.72	MLT 94			
07673022	**Low discrimination**						**07673022**
					SPO		
Actinomyces naeslundii		80.21	0.69	MLT 94	1		
Lactobacillus acidophilus		8.00	0.27		1		
Clostridium ramosum		6.67	0.25		92		
07673023	**Very good identification**						**07673023**
Actinomyces naeslundii		100.0	0.72	MLT 94			
07673030	**Good identification**						**07673030**
					SPO		
Actinomyces naeslundii		95.36	0.47	MLT 94	1		
Clostridium innocuum		4.64	0.00	MLZ 5	99		
07673031	**Good identification**						**07673031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MLT 94		
07673032	**Good identification**					**07673032**
					SPO	
Actinomyces naeslundii		95.36	0.47	MLT 94	1	
Clostridium innocuum		4.64	0.00	MLZ 5	99	
07673033	**Good identification**					**07673033**
Actinomyces naeslundii		100.0	0.50	MLT 94		
07673040	**Low discrimination**					**07673040**
					SPO	
Actinomyces naeslundii		72.03	0.56	MLT 94	1	
Clostridium innocuum		25.08	0.24		99	
Actinomyces israelii		2.89	0.23		1	
07673041	**Very good identification**					**07673041**
Actinomyces naeslundii		100.0	0.59	MLT 94		
07673042	**Low discrimination**					**07673042**
					SPO	
Actinomyces naeslundii		67.65	0.56	MLT 94	1	
Clostridium innocuum		23.56	0.24		99	
Lactobacillus acidophilus		4.73	0.12		1	
07673043	**Very good identification**					**07673043**
Actinomyces naeslundii		100.0	0.59	MLT 94		
07673051	**Good identification**					**07673051**
Actinomyces naeslundii		100.0	0.37	MLT 94		
07673053	**Good identification**					**07673053**
Actinomyces naeslundii		100.0	0.37	MLT 94		
07673060	**Good identification**					**07673060**
Actinomyces naeslundii		100.0	0.56	MLT 94		
07673061	**Very good identification**					**07673061**
Actinomyces naeslundii		100.0	0.59	MLT 94		
07673062	**Acceptable identification**					**07673062**
Actinomyces naeslundii		93.47	0.56	MLT 94		
Lactobacillus acidophilus		6.53	0.12			
07673063	**Very good identification**					**07673063**
Actinomyces naeslundii		100.0	0.59	MLT 94		
07673071	**Good identification**					**07673071**
Actinomyces naeslundii		100.0	0.37	MLT 94		
07673073	**Good identification**					**07673073**
Actinomyces naeslundii		100.0	0.37	MLT 94		
07673100	**Low discrimination**					**07673100**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		88.11	0.77	MLT 94	99	
Fusobacterium mortiferum		6.60	0.46		1	
Actinomyces israelii		2.25	0.40		99	
07673101	**Very good identification**					**07673101**
					GRA	
Actinomyces naeslundii		99.95	0.80	MLT 94	99	
Fusobacterium mortiferum		0.05	0.12		1	
07673102	**Low discrimination**					**07673102**
Actinomyces naeslundii		66.05	0.77	MLT 94		
Lactobacillus acidophilus		26.61	0.45			
Actinomyces israelii		2.53	0.43			
07673103	**Very good identification**					**07673103**
Actinomyces naeslundii		99.73	0.80	MLT 94		
Lactobacillus acidophilus		0.27	0.11			
07673110	**Low discrimination**					**07673110**
					SPO	
Actinomyces naeslundii		88.25	0.55	MLT 94	1	
Actinomyces israelii		6.41	0.26	MLZ 5	1	
Clostridium tertium		2.57	0.05		99	
07673111	**Very good identification**					**07673111**
Actinomyces naeslundii		100.0	0.58	MLT 94		
07673112	**Low discrimination**					**07673112**
Actinomyces naeslundii		65.06	0.55	MLT 94		
Bifidobacterium breve		7.87	0.24	MLZ 5		
Bifidobacterium adolescentis		7.18	0.23			
07673113	**Very good identification**					**07673113**
Actinomyces naeslundii		100.0	0.58	MLT 94		
07673120	**Low discrimination**					**07673120**
					SPO	
Actinomyces naeslundii		57.69	0.77	MLT 94	1	
Clostridium ramosum		41.72	0.49		92	
Lactobacillus acidophilus		0.24	0.11		1	
07673121	**Very good identification**					**07673121**
					SPO	
Actinomyces naeslundii		99.52	0.80	MLT 94	1	
Clostridium ramosum		0.49	0.15		92	
07673122	**Low discrimination**					**07673122**
					SPO	
Clostridium ramosum		66.91	0.59	MLT 99	92	
Actinomyces naeslundii		23.13	0.77		1	
Lactobacillus acidophilus		9.32	0.45		1	
07673123	**Very good identification**					**07673123**
					SPO	
Actinomyces naeslundii		97.83	0.80	MLT 94	1	
Clostridium ramosum		1.91	0.25		92	
Lactobacillus acidophilus		0.27	0.11		1	
07673130	**Low discrimination**					**07673130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		80.17	0.55	MLT 94	1	
Clostridium ramosum		11.13	0.15	MLZ 5	92	
Clostridium tertium		7.00	0.13		99	
07673131	**Very good identification**					**07673131**
Actinomyces naeslundii		100.0	0.58	MLT 94		
07673132	**Low discrimination**					**07673132**
					SPO	
Actinomyces naeslundii		46.67	0.55	MLT 94	1	
Clostridium ramosum		25.91	0.25	MLZ 5	92	
Clostridium tertium		12.22	0.21		99	
07673133	**Very good identification**					**07673133**
Actinomyces naeslundii		100.0	0.58	MLT 94		
07673140	**Good identification**					**07673140**
					GRA	
Actinomyces naeslundii		94.28	0.64	MLT 94	99	
Actinomyces israelii		5.31	0.33		99	
Fusobacterium mortiferum		0.40	0.12		1	
07673141	**Very good identification**					**07673141**
Actinomyces naeslundii		100.0	0.67	MLT 94		
07673142	**Low discrimination**					**07673142**
Actinomyces naeslundii		71.29	0.64	MLT 94		
Lactobacillus acidophilus		20.12	0.30			
Actinomyces israelii		6.03	0.36			
07673143	**Very good identification**					**07673143**
Actinomyces naeslundii		100.0	0.67	MLT 94		
07673150	**Low discrimination**					**07673150**
Actinomyces naeslundii		86.21	0.42	MLT 94		
Actinomyces israelii		13.79	0.19	MLZ 5		
07673151	**Good identification**					**07673151**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673152	**Low discrimination**					**07673152**
Actinomyces naeslundii		49.94	0.42	MLT 94		
Bifidobacterium breve		34.22	0.24	MLZ 5		
Actinomyces israelii		11.98	0.22			
07673153	**Good identification**					**07673153**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673160	**Very good identification**					**07673160**
					SPO	
Actinomyces naeslundii		95.44	0.64	MLT 94	1	
Clostridium ramosum		3.95	0.15		92	
Bifidobacterium breve		0.61	0.12		1	
07673161	**Very good identification**					**07673161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.67	MLT 94		
07673162	**Low discrimination**					**07673162**
					SPO	
Actinomyces naeslundii		65.91	0.64	MLT 94	1	
Lactobacillus acidophilus		18.60	0.30		1	
Clostridium ramosum		10.91	0.25		92	
07673163	**Very good identification**					**07673163**
Actinomyces naeslundii		100.0	0.67	MLT 94		
07673170	**Low discrimination**					**07673170**
Actinomyces naeslundii		89.25	0.42	MLT 94		
Bifidobacterium breve		10.75	0.12	MLZ 5		
07673171	**Good identification**					**07673171**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673173	**Good identification**					**07673173**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673201	**Good identification**					**07673201**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07673203	**Good identification**					**07673203**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07673221	**Good identification**					**07673221**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07673223	**Good identification**					**07673223**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07673300	**Low discrimination**					**07673300**
					SPO	
Actinomyces naeslundii		44.96	0.42	MLT 94	1	
Clostridium ramosum		43.10	0.17	RHA 1	92	
Actinomyces israelii		8.57	0.21		1	
07673301	**Good identification**					**07673301**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673303	**Good identification**					**07673303**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673320	**Good identification**					**07673320**
					SPO	
Clostridium ramosum		98.96	0.51	MLT 99	92	
Actinomyces naeslundii		1.04	0.42		1	
07673321	**Low discrimination**					**07673321**
					SPO	
Actinomyces naeslundii		61.01	0.45	MLT 94	1	
Clostridium ramosum		38.99	0.17	RHA 1	92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07673322	**Very good identification**					**07673322**
					SPO	
Clostridium ramosum		98.71	0.61	MLT 99	92	
Lactobacillus acidophilus		0.78	0.26		1	
Actinomyces naeslundii		0.26	0.42		1	
07673400	**Good identification**					**07673400**
					SPO	
Clostridium innocuum		97.73	0.45	RAF 1	99	
Actinomyces israelii		1.36	0.28		1	
Actinomyces naeslundii		0.92	0.35		1	
07673401	**Low discrimination**					**07673401**
					SPO	
Actinomyces naeslundii		58.28	0.38	MLT 94	1	
Clostridium innocuum		41.72	0.11	ARA 1	99	
07673402	**Good identification**					**07673402**
					SPO	
Clostridium innocuum		97.07	0.45	RAF 1	99	
Actinomyces israelii		2.02	0.31		1	
Actinomyces naeslundii		0.91	0.35		1	
07673403	**Low discrimination**					**07673403**
					SPO	
Actinomyces naeslundii		58.28	0.38	MLT 94	1	
Clostridium innocuum		41.72	0.11	ARA 1	99	
07673421	**Good identification**					**07673421**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07673423	**Good identification**					**07673423**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07673500	**Low discrimination**					**07673500**
					SPO	
Actinomyces israelii		49.70	0.39	MLT 99	1	
Actinomyces naeslundii		24.00	0.42	XYL 92	1	
Clostridium butyricum		15.46	0.09		99	
07673501	**Good identification**					**07673501**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673502	**Low discrimination**					**07673502**
Bifidobacterium adolescentis		38.00	0.35	MLT 99		
Actinomyces israelii		37.49	0.42	XYL 87		
Actinomyces naeslundii		12.07	0.42			
07673503	**Good identification**					**07673503**
Actinomyces naeslundii		100.0	0.45	MLT 94		
07673512	**Low discrimination**					**07673512**
Bifidobacterium adolescentis		86.78	0.37	MLT 99		
Actinomyces israelii		10.47	0.28	XYL 87		
Bifidobacterium dentium		2.76	-.07			
07673520	**Low discrimination**					**07673520**
					SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		53.90	0.42	MLT 94	1		
Clostridium ramosum		38.98	0.15	ARA 1	92		
Actinomyces israelii		7.13	0.18		1		
07673521	**Good identification**						**07673521**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07673523	**Good identification**						**07673523**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07674000	**Good identification**						**07674000**
Fusobacterium mortiferum		100.0	0.51	XYL 7			
07674002	**Good identification**						**07674002**
Fusobacterium mortiferum		100.0	0.35	XYL 7			
07674100	**Very good identification**						**07674100**
					GRA	SPO	
Fusobacterium mortiferum		99.43	0.58	XYL 7	1	1	
Clostridium butyricum		0.57	0.01		99	99	
07674102	**Acceptable identification**						**07674102**
					GRA	SPO	
Fusobacterium mortiferum		94.53	0.41	XYL 7	1	1	
Clostridium butyricum		1.89	-.07	BGL 10	99	99	
Actinomyces israelii		1.80	0.14		99	1	
07674122	**Good identification**						**07674122**
					GRA		
Parabacteroides distasonis		97.42	0.37	GAL 99	1		
Lactobacillus acidophilus		2.58	-.01		99		
07674322	**Good identification**						**07674322**
Parabacteroides distasonis		95.61	0.46	GAL 99			
Clostridium clostridioforme		4.39	0.17				
07674332	**Low discrimination**						**07674332**
Parabacteroides distasonis		72.04	0.41	GAL 99			
Clostridium clostridioforme		27.96	0.28				
07674612	**Good identification**						**07674612**
Clostridium clostridioforme		100.0	0.40	MLT 99			
07674632	**Good identification**						**07674632**
Clostridium clostridioforme		100.0	0.40	MLT 99			
07674702	**Good identification**						**07674702**
Clostridium clostridioforme		100.0	0.37	MLT 99			
07674712	**Good identification**						**07674712**
Clostridium clostridioforme		100.0	0.49	MLT 99			
07674722	**Low discrimination**						**07674722**
Clostridium clostridioforme		53.97	0.37	MLT 99			
Parabacteroides distasonis		46.03	0.42				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07674732	**Acceptable identification**						**07674732**
Clostridium clostridioforme		90.84	0.49	MLT 99			
Parabacteroides distasonis		9.16	0.37				
07675000	**Low discrimination**						**07675000**
					GRA	SPO	
Fusobacterium mortiferum		89.86	0.54	XYL 7	1	1	
Actinomyces israelii		4.87	0.35		99	1	
Clostridium butyricum		3.29	0.11		99	99	
07675002	**Low discrimination**						**07675002**
					GRA		
Fusobacterium mortiferum		43.81	0.38	XYL 7	1		
Actinomyces israelii		32.08	0.38	BGL 10	99		
Lactobacillus acidophilus		11.09	0.15		99		
07675022	**Low discrimination**						**07675022**
					GRA		
Parabacteroides distasonis		75.09	0.37	LAC 99	1		
Lactobacillus acidophilus		17.11	0.15		99		
Actinomyces naeslundii		4.65	0.30		99		
07675100	**Low discrimination**						**07675100**
					GRA	SPO	
Fusobacterium mortiferum		65.89	0.61	XYL 7	1	1	
Clostridium butyricum		24.85	0.35		99	99	
Actinomyces israelii		6.13	0.45		99	1	
07675101	**Low discrimination**						**07675101**
					GRA	SPO	
Actinomyces naeslundii		50.14	0.41	MLT 94	99	1	
Fusobacterium mortiferum		36.21	0.26	XYL 1	1	1	
Clostridium butyricum		13.65	0.01		99	99	
07675103	**Low discrimination**						**07675103**
					SPO		
Actinomyces naeslundii		76.53	0.41	MLT 94	1		
Clostridium butyricum		8.10	-.07	XYL 1	99		
Actinomyces israelii		7.71	0.14		1		
07675112	**Low discrimination**						**07675112**
					GRA		
Bifidobacterium adolescentis		53.69	0.38	MLT 99	99		
Parabacteroides distasonis		17.96	0.33	ARA 87	1		
Actinomyces israelii		13.14	0.34		99		
07675120	**Low discrimination**						**07675120**
					GRA	SPO	
Parabacteroides distasonis		43.62	0.37	BGL 99	1	1	
Clostridium butyricum		27.04	0.17		99	99	
Fusobacterium mortiferum		8.33	0.26		1	1	
07675121	**Good identification**						**07675121**
Actinomyces naeslundii		100.0	0.41	MLT 94			
07675122	**Very good identification**						**07675122**
					GRA		
Parabacteroides distasonis		96.31	0.72		1		
Lactobacillus acidophilus		2.55	0.33		99		
Prevotella oralis		0.35	0.17		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07675123	**Low discrimination**						**07675123**
					GRA		
Parabacteroides distasonis		77.47	0.37	URE 1	1		
Actinomyces naeslundii		20.48	0.41		99		
Lactobacillus acidophilus		2.05	-.01		99		
07675132	**Very good identification**						**07675132**
					GRA		
Parabacteroides distasonis		99.43	0.67		1		
Clostridium clostridioforme		0.48	0.22		50		
Actinomyces israelii		0.05	0.14		99		
07675140	**Low discrimination**						**07675140**
					GRA	SPO	
Actinomyces israelii		44.81	0.38	MLT 99	99	1	
Clostridium butyricum		40.63	0.17		99	99	
Fusobacterium mortiferum		12.52	0.26		1	1	
07675142	**Low discrimination**						**07675142**
					SPO		
Actinomyces israelii		57.47	0.41	MLT 99	1		
Lactobacillus acidophilus		18.15	0.18		1		
Clostridium butyricum		13.51	0.10		99		
07675162	**Low discrimination**						**07675162**
					GRA		
Parabacteroides distasonis		70.41	0.37	SOR 1	1		
Lactobacillus acidophilus		22.80	0.18		99		
Actinomyces israelii		4.61	0.21		99		
07675222	**Good identification**						**07675222**
Parabacteroides distasonis		96.05	0.46	LAC 99			
Clostridium clostridioforme		3.95	0.16				
07675232	**Low discrimination**						**07675232**
Parabacteroides distasonis		74.20	0.41	LAC 99			
Clostridium clostridioforme		25.80	0.27				
07675302	**Low discrimination**						**07675302**
					GRA		
Parabacteroides distasonis		77.04	0.46	NAG 99	1		
Clostridium clostridioforme		10.60	0.25		50		
Actinomyces israelii		5.10	0.29		99		
07675312	**Low discrimination**						**07675312**
					GRA		
Clostridium clostridioforme		53.34	0.36	MLT 99	50		
Parabacteroides distasonis		45.82	0.41	ARA 94	1		
Actinomyces israelii		0.84	0.15		99		
07675320	**Good identification**						**07675320**
					SPO		
Parabacteroides distasonis		98.11	0.46	BGL 99	1		
Clostridium ramosum		1.89	0.04		92		
07675322	**Very good identification**						**07675322**
					SPO		
Parabacteroides distasonis		99.65	0.80		1		
Clostridium clostridioforme		0.14	0.25		66		
Clostridium ramosum		0.08	0.14		92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07675323	**Good identification**						**07675323**
Parabacteroides distasonis		100.0	0.46	URE 1			
07675330	**Good identification**						**07675330**
Parabacteroides distasonis		100.0	0.41	BGL 99			
07675332	**Very good identification**						**07675332**
Parabacteroides distasonis		98.82	0.75				
Clostridium clostridioforme		1.16	0.36				
Bacteroides ovatus		0.02	-.06				
07675333	**Good identification**						**07675333**
Parabacteroides distasonis		100.0	0.41	URE 1			
07675362	**Good identification**						**07675362**
					GRA		
Parabacteroides distasonis		99.19	0.46	SOR 1	1		
Lactobacillus acidophilus		0.81	-.02		99		
07675372	**Good identification**						**07675372**
Parabacteroides distasonis		100.0	0.41	SOR 1			
07675500	**Good identification**						**07675500**
					GRA	SPO	
Clostridium butyricum		97.39	0.51	ESL 88	99	99	
Actinomyces israelii		2.18	0.44	MLT 99	99	1	
Fusobacterium mortiferum		0.29	0.26		1	1	
07675502	**Low discrimination**						**07675502**
					SPO		
Clostridium butyricum		66.64	0.44	ESL 88	99		
Bifidobacterium adolescentis		23.54	0.51	MLT 99	1		
Actinomyces israelii		5.76	0.47		1		
07675512	**Low discrimination**						**07675512**
					SPO		
Bifidobacterium adolescentis		64.13	0.53	MLT 99	1		
Clostridium clostridioforme		23.92	0.42		66		
Clostridium butyricum		7.82	0.22		99		
07675522	**Good identification**						**07675522**
					GRA	SPO	
Parabacteroides distasonis		94.70	0.68		1	1	
Clostridium butyricum		3.39	0.26		99	99	
Clostridium clostridioforme		1.38	0.31		50	66	
07675532	**Low discrimination**						**07675532**
					GRA	SPO	
Parabacteroides distasonis		88.23	0.63		1	1	
Clostridium clostridioforme		10.85	0.42		50	66	
Clostridium butyricum		0.31	0.04		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07675552	**Low discrimination**						**07675552**
					SPO		
Bifidobacterium adolescentis		82.62	0.37	MLT 99	1		
Actinomyces israelii		7.78	0.26	SOR 11	1		
Clostridium butyricum		7.09	0.04		99		
07675602	**Good identification**						**07675602**
					SPO		
Clostridium clostridioforme		96.37	0.37	MLT 99	66		
Actinomyces israelii		2.03	0.17		1		
Clostridium butyricum		1.60	-.06		99		
07675612	**Good identification**						**07675612**
Clostridium clostridioforme		100.0	0.48	MLT 99			
07675622	**Low discrimination**						**07675622**
Clostridium clostridioforme		51.24	0.37	MLT 99			
Parabacteroides distasonis		48.77	0.42				
07675632	**Low discrimination**						**07675632**
Clostridium clostridioforme		89.89	0.48	MLT 99			
Parabacteroides distasonis		10.11	0.37				
07675702	**Low discrimination**						**07675702**
					GRA	SPO	
Clostridium clostridioforme		69.33	0.46	MLT 99	50	66	
Parabacteroides distasonis		19.71	0.42		1	1	
Clostridium butyricum		8.27	0.18		99	99	
07675712	**Very good identification**						**07675712**
					GRA		
Clostridium clostridioforme		96.20	0.57	MLT 99	50		
Parabacteroides distasonis		3.23	0.37		1		
Bifidobacterium adolescentis		0.36	0.18		99		
07675720	**Low discrimination**						**07675720**
					GRA	SPO	
Parabacteroides distasonis		83.39	0.42	BGL 99	1	1	
Clostridium butyricum		7.83	0.07		99	99	
Bacteroides vulgatus		5.82	0.04		1	1	
07675722	**Very good identification**						**07675722**
Parabacteroides distasonis		95.73	0.76				
Clostridium clostridioforme		3.40	0.46				
Bacteroides ovatus		0.63	0.23				
07675723	**Good identification**						**07675723**
Parabacteroides distasonis		96.57	0.42	URE 1			
Clostridium clostridioforme		3.43	0.11				
07675730	**Low discrimination**						**07675730**
Parabacteroides distasonis		76.89	0.37	BGL 99			
Clostridium clostridioforme		23.11	0.23				
07675732	**Low discrimination**						**07675732**
Parabacteroides distasonis		76.32	0.71				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		22.94	0.57			
Bacteroides ovatus		0.63	0.20			
07675733	**Low discrimination**					**07675733**
Parabacteroides distasonis		76.89	0.37	URE 1		
Clostridium clostridioforme		23.11	0.23			
07675762	**Acceptable identification**					**07675762**
Parabacteroides distasonis		94.09	0.42	SOR 1		
Clostridium clostridioforme		3.34	0.11			
Bacteroides ovatus		2.57	-.01			
07675772	**Low discrimination**					**07675772**
Parabacteroides distasonis		74.94	0.37	SOR 1		
Clostridium clostridioforme		22.53	0.23			
Bacteroides ovatus		2.53	-.04			
07677000	**Low discrimination**					**07677000**
					SPO	
Clostridium innocuum		58.31	0.36	RAF 1	99	
Actinomyces israelii		38.10	0.48	XYL 5	1	
Actinomyces naeslundii		1.84	0.35		1	
07677001	**Low discrimination**					**07677001**
					SPO	
Actinomyces naeslundii		73.92	0.38	MLT 94	1	
Clostridium innocuum		15.78	0.02	XYL 1	99	
Actinomyces israelii		10.31	0.14		1	
07677002	**Low discrimination**					**07677002**
					SPO	
Clostridium innocuum		49.71	0.36	RAF 1	99	
Actinomyces israelii		48.72	0.51	XYL 5	1	
Actinomyces naeslundii		1.57	0.35		1	
07677003	**Low discrimination**					**07677003**
					SPO	
Actinomyces naeslundii		70.29	0.38	MLT 94	1	
Clostridium innocuum		15.00	0.02	XYL 1	99	
Actinomyces israelii		14.70	0.17		1	
07677012	**Low discrimination**					**07677012**
					SPO	
Actinomyces israelii		77.85	0.37	MLT 99	1	
Clostridium innocuum		22.15	0.13	MLZ 13	99	
07677021	**Good identification**					**07677021**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07677023	**Good identification**					**07677023**
Actinomyces naeslundii		100.0	0.38	MLT 94		
07677040	**Good identification**					**07677040**
					SPO	
Actinomyces israelii		96.18	0.41	MLT 99	1	
Clostridium innocuum		3.82	0.02		99	
07677042	**Good identification**					**07677042**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		97.42	0.44	MLT 99	1		
Clostridium innocuum		2.58	0.02		99		
07677100	**Low discrimination**						**07677100**
					SPO		
Actinomyces israelii		87.65	0.58	MLT 99	1		
Clostridium butyricum		6.97	0.18		99		
Actinomyces naeslundii		3.01	0.42		1		
07677101	**Low discrimination**						**07677101**
Actinomyces naeslundii		83.61	0.45	MLT 94			
Actinomyces israelii		16.39	0.24	XYL 1			
07677102	**Low discrimination**						**07677102**
Actinomyces israelii		88.07	0.61	MLT 99			
Bifidobacterium adolescentis		6.35	0.35				
Actinomyces naeslundii		2.02	0.42				
07677103	**Low discrimination**						**07677103**
Actinomyces naeslundii		77.28	0.45	MLT 94			
Actinomyces israelii		22.72	0.27	XYL 1			
07677110	**Acceptable identification**						**07677110**
					SPO		
Actinomyces israelii		92.93	0.44	MLT 99	1		
Clostridium tertium		4.47	0.07	MLZ 13	99		
Clostridium butyricum		2.60	-.04		99		
07677112	**Low discrimination**						**07677112**
					SPO		
Actinomyces israelii		58.66	0.47	MLT 99	1		
Bifidobacterium adolescentis		34.60	0.37	MLZ 13	1		
Clostridium tertium		5.65	0.15		99		
07677120	**Low discrimination**						**07677120**
					SPO		
Actinomyces israelii		44.93	0.38	MLT 99	1		
Actinomyces naeslundii		24.18	0.42	NAG 6	1		
Clostridium ramosum		17.49	0.15		92		
07677121	**Good identification**						**07677121**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07677122	**Low discrimination**						**07677122**
					SPO	GRA	
Parabacteroides distasonis		26.23	0.37	MAN 1	1	1	
Clostridium ramosum		25.96	0.25		92	50	
Actinomyces israelii		25.01	0.41		1	99	
07677123	**Good identification**						**07677123**
Actinomyces naeslundii		100.0	0.45	MLT 94			
07677140	**Good identification**						**07677140**
					SPO		
Actinomyces israelii		96.77	0.51	MLT 99	1		
Clostridium butyricum		1.72	0.00		99		
Actinomyces naeslundii		1.51	0.30		1		
07677142	**Good identification**						**07677142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		96.35	0.54	MLT 99			
Bifidobacterium adolescentis		2.21	0.20				
Actinomyces naeslundii		1.00	0.30				
07677150	**Acceptable identification**						**07677150**
					COC		
Actinomyces israelii		93.35	0.37	MLT 99	1		
Blautia producta		6.65	-.03	MLZ 13	99		
07677152	**Low discrimination**						**07677152**
Actinomyces israelii		84.21	0.40	MLT 99			
Bifidobacterium adolescentis		15.79	0.21	MLZ 13			
07677300	**Good identification**						**07677300**
					SPO		
Actinomyces israelii		96.84	0.39	MLT 99	1		
Clostridium butyricum		3.16	-.08	RHA 7	99		
07677302	**Good identification**						**07677302**
					GRA		
Actinomyces israelii		97.38	0.42	MLT 99	99		
Parabacteroides distasonis		2.62	0.11	RHA 7	1		
07677322	**Low discrimination**						**07677322**
					SPO	GRA	
Parabacteroides distasonis		68.43	0.46	MAN 1	1	1	
Clostridium ramosum		29.93	0.27		92	50	
Actinomyces israelii		1.64	0.22		1	99	
07677332	**Good identification**						**07677332**
Parabacteroides distasonis		100.0	0.41	MAN 1			
07677342	**Good identification**						**07677342**
Actinomyces israelii		100.0	0.35	MLT 99			
07677400	**Low discrimination**						**07677400**
					SPO		
Actinomyces israelii		67.76	0.46	MLT 99	1		
Clostridium innocuum		22.37	0.23		99		
Clostridium butyricum		9.88	0.11		99		
07677402	**Low discrimination**						**07677402**
					SPO		
Actinomyces israelii		78.02	0.49	MLT 99	1		
Clostridium innocuum		17.17	0.23		99		
Clostridium butyricum		2.95	0.04		99		
07677412	**Low discrimination**						**07677412**
Actinomyces israelii		74.19	0.35	MLT 99			
Bifidobacterium adolescentis		14.46	0.17	MLZ 13			
Bifidobacterium dentium		6.80	-.07				
07677440	**Good identification**						**07677440**
					SPO		
Actinomyces israelii		96.84	0.39	MLT 99	1		
Clostridium butyricum		3.16	-.07		99		
07677442	**Good identification**						**07677442**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.42	MLT 99			
07677500	**Low discrimination**						**07677500**
					SPO		
Actinomyces israelii		53.09	0.57	MLT 99	1		
Clostridium butyricum		46.44	0.35		99		
Bifidobacterium adolescentis		0.47	0.15		1		
07677502	**Low discrimination**						**07677502**
					SPO		
Actinomyces israelii		54.95	0.60	MLT 99	1		
Bifidobacterium adolescentis		32.41	0.50		1		
Clostridium butyricum		12.46	0.28		99		
07677510	**Low discrimination**						**07677510**
					SPO		
Actinomyces israelii		57.09	0.43	MLT 99	1		
Bifidobacterium dentium		19.21	0.10	MLZ 13	1		
Clostridium butyricum		17.59	0.13		99		
07677512	**Low discrimination**						**07677512**
Bifidobacterium adolescentis		58.87	0.51	MLT 99			
Bifidobacterium dentium		27.68	0.27				
Actinomyces israelii		12.20	0.46				
07677522	**Low discrimination**						**07677522**
					GRA	SPO	
Actinomyces israelii		43.80	0.39	MLT 99	99	1	
Parabacteroides distasonis		34.40	0.34	NAG 6	1	1	
Clostridium butyricum		13.53	0.10		99	99	
07677540	**Low discrimination**						**07677540**
					COC	SPO	
Actinomyces israelii		61.53	0.50	MLT 99	1	1	
Blautia producta		26.43	0.24		99	1	
Clostridium butyricum		12.04	0.17		1	99	
07677542	**Low discrimination**						**07677542**
					SPO		
Actinomyces israelii		78.74	0.53	MLT 99	1		
Bifidobacterium adolescentis		14.76	0.34		1		
Clostridium butyricum		3.99	0.10		99		
07677552	**Low discrimination**						**07677552**
					COC		
Bifidobacterium adolescentis		47.46	0.36	MLT 99	1		
Actinomyces israelii		30.95	0.39	SOR 11	1		
Blautia producta		19.77	0.15		99		
07677700	**Low discrimination**						**07677700**
					SPO		
Actinomyces israelii		73.56	0.38	MLT 99	1		
Clostridium butyricum		26.44	0.09	RHA 7	99		
07677702	**Low discrimination**						**07677702**
					SPO		
Actinomyces israelii		80.43	0.41	MLT 99	1		
Clostridium butyricum		7.49	0.02	RHA 7	99		
Bifidobacterium adolescentis		6.37	0.15		1		
07677722	**Low discrimination**						**07677722**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		77.77	0.42	MAN 1			
Bacteroides ovatus		16.99	0.15				
Clostridium clostridioforme		2.76	0.11				
07677732	**Low discrimination**						**07677732**
Parabacteroides distasonis		60.75	0.37	MAN 1			
Clostridium clostridioforme		18.26	0.23				
Bacteroides ovatus		16.43	0.12				
07700100	**Low discrimination**						**07700100**
					GRA		
Fusobacterium mortiferum		88.89	0.44	MLT 11	1		
Actinomyces naeslundii		9.87	0.40	FRU 99	99		
Actinomyces viscosus		1.24	0.00		99		
07700101	**Good identification**						**07700101**
Actinomyces naeslundii		100.0	0.43	FRU 95			
07700121	**Good identification**						**07700121**
Actinomyces naeslundii		100.0	0.43	FRU 95			
07700122	**Low discrimination**						**07700122**
					GRA		
Actinomyces naeslundii		44.89	0.40	FRU 95	99		
Prevotella oralis		24.68	0.09	GAL 96	1		
Prevotella melaninogenica		16.82	0.03		1		
07700123	**Good identification**						**07700123**
Actinomyces naeslundii		100.0	0.43	FRU 95			
07701000	**Low discrimination**						**07701000**
					COC	GRA	
Actinomyces naeslundii		45.09	0.55	FRU 95	1	99	
Fusobacterium mortiferum		30.97	0.41		1	1	
Gemella morbillorum		15.52	0.26		99	99	
07701001	**Very good identification**						**07701001**
Actinomyces naeslundii		100.0	0.58	FRU 95			
07701002	**Low discrimination**						**07701002**
					GRA		
Actinomyces naeslundii		60.56	0.55	FRU 95	99		
Actinomyces viscosus		30.81	0.26		99		
Fusobacterium mortiferum		4.62	0.25		1		
07701003	**Low discrimination**						**07701003**
Actinomyces naeslundii		89.84	0.58	FRU 95			
Actinomyces viscosus		10.16	0.18				
07701011	**Good identification**						**07701011**
Actinomyces naeslundii		100.0	0.37	FRU 95			
07701013	**Good identification**						**07701013**
Actinomyces naeslundii		100.0	0.37	FRU 95			
07701020	**Low discrimination**						**07701020**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		63.27	0.55	FRU 95	99	
Clostridium perfringens		18.19	0.24		99	
Prevotella melaninogenica		14.75	0.15		1	
07701021	**Very good identification**					**07701021**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07701022	**Low discrimination**					**07701022**
					GRA	
Actinomyces naeslundii		62.01	0.55	FRU 95	99	
Prevotella oralis		25.55	0.23		1	
Clostridium perfringens		8.39	0.18		99	
07701023	**Very good identification**					**07701023**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07701031	**Good identification**					**07701031**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07701033	**Good identification**					**07701033**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07701040	**Good identification**					**07701040**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07701041	**Good identification**					**07701041**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07701042	**Good identification**					**07701042**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07701043	**Good identification**					**07701043**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07701060	**Low discrimination**					**07701060**
Actinomyces naeslundii		76.32	0.43	FRU 95		
Clostridium perfringens		23.68	0.12			
07701061	**Good identification**					**07701061**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07701062	**Low discrimination**					**07701062**
Actinomyces naeslundii		87.26	0.43	FRU 95		
Clostridium perfringens		12.74	0.06			
07701063	**Good identification**					**07701063**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07701100	**Low discrimination**					**07701100**
					GRA	
Actinomyces naeslundii		57.44	0.63	FRU 95	99	
Fusobacterium mortiferum		32.35	0.47		1	
Prevotella melaninogenica		4.71	0.15		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07701101	**Very good identification**						**07701101**
					GRA	SPO	
Actinomyces naeslundii		95.72	0.66	FRU 95	99	1	
Clostridium chauvoei		3.92	0.19		99	99	
Fusobacterium mortiferum		0.36	0.13		1	1	
07701102	**Low discrimination**						**07701102**
Actinomyces naeslundii		49.92	0.63	FRU 95			
Bifidobacterium longum subsp. infantis		20.54	0.27				
Actinomyces viscosus		15.86	0.31				
07701103	**Good identification**						**07701103**
					SPO		
Actinomyces naeslundii		92.54	0.66	FRU 95	1		
Actinomyces viscosus		6.54	0.22		1		
Clostridium chauvoei		0.67	0.07		99		
07701110	**Acceptable identification**						**07701110**
					GRA		
Actinomyces naeslundii		90.25	0.41	FRU 95	99		
Fusobacterium mortiferum		9.75	0.13	MLZ 5	1		
07701111	**Good identification**						**07701111**
Actinomyces naeslundii		100.0	0.44	FRU 95			
07701112	**Low discrimination**						**07701112**
Actinomyces naeslundii		60.40	0.41	FRU 95			
Bifidobacterium adolescentis		34.84	0.22	MLZ 5			
Bifidobacterium longum subsp. infantis		4.77	-.07				
07701113	**Good identification**						**07701113**
Actinomyces naeslundii		100.0	0.44	FRU 95			
07701120	**Low discrimination**						**07701120**
					GRA		
Prevotella melaninogenica		80.67	0.50	SAL 1	1		
Actinomyces naeslundii		9.94	0.63	FRU 90	99		
Clostridium perfringens		7.37	0.39		99		
07701121	**Good identification**						**07701121**
					GRA	SPO	
Actinomyces naeslundii		93.17	0.66	FRU 95	99	1	
Prevotella melaninogenica		5.09	0.15		1	1	
Clostridium chauvoei		1.27	0.11		99	99	
07701122	**Low discrimination**						**07701122**
					GRA		
Prevotella oralis		38.12	0.44	FRU 99	1		
Prevotella melaninogenica		25.97	0.37		1		
Actinomyces naeslundii		16.81	0.63		99		
07701123	**Very good identification**						**07701123**
					GRA		
Actinomyces naeslundii		97.03	0.66	FRU 95	99		
Prevotella oralis		1.48	0.09		1		
Prevotella melaninogenica		1.01	0.03		1		
07701131	**Good identification**						**07701131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	FRU 95		
07701132	**Low discrimination**					**07701132**
					GRA	
Actinomyces naeslundii		44.09	0.41	FRU 95	99	
Prevotella oralis		19.19	0.09	MLZ 5	1	
Prevotella melaninogenica		13.07	0.03		1	
07701133	**Good identification**					**07701133**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07701140	**Acceptable identification**					**07701140**
					GRA	
Actinomyces naeslundii		94.61	0.50	FRU 95	99	
Fusobacterium mortiferum		3.05	0.13		1	
Clostridium perfringens		2.34	0.00		99	
07701141	**Good identification**					**07701141**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07701142	**Good identification**					**07701142**
Actinomyces naeslundii		97.70	0.50	FRU 95		
Bifidobacterium longum subsp. infantis		2.30	-.07			
07701143	**Good identification**					**07701143**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07701160	**Low discrimination**					**07701160**
					GRA	
Actinomyces naeslundii		44.17	0.50	FRU 95	99	
Clostridium perfringens		35.32	0.26		99	
Prevotella melaninogenica		20.51	0.15		1	
07701161	**Good identification**					**07701161**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07701162	**Low discrimination**					**07701162**
					GRA	
Actinomyces naeslundii		61.11	0.50	FRU 95	99	
Clostridium perfringens		22.99	0.21		99	
Prevotella oralis		7.93	0.09		1	
07701163	**Good identification**					**07701163**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07701322	**Low discrimination**					**07701322**
Prevotella oralis		89.07	0.37	FRU 99		
Parabacteroides distasonis		7.90	0.32			
Prevotella melaninogenica		1.43	0.03			
07701512	**Low discrimination**					**07701512**
Bifidobacterium adolescentis		60.42	0.36	FRU 99		
Bifidobacterium longum subsp. longum		39.58	0.31	XYL 87		
07701520	**Low discrimination**					**07701520**
					GRA	
Bifidobacterium longum subsp. longum		70.71	0.37	SAL 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		18.53	0.15	FRU 90	1	
Bacteroides vulgatus		5.37	0.04		1	
07701522	**Low discrimination**					**07701522**
					GRA	
Bifidobacterium longum subsp. longum		79.65	0.37	SAL 1	99	
Prevotella oralis		5.84	0.09	FRU 90	1	
Parabacteroides distasonis		4.49	0.20		1	
07701530	**Good identification**					**07701530**
Bifidobacterium longum subsp. longum		100.0	0.41	SAL 1		
07701532	**Good identification**					**07701532**
					GRA	
Bifidobacterium longum subsp. longum		98.25	0.41	SAL 1	99	
Parabacteroides distasonis		1.75	0.16	FRU 90	1	
07702000	**Low discrimination**					**07702000**
Actinomyces naeslundii		70.12	0.37	FRU 95		
Propionibacterium granulosum		29.88	0.16	GAL 96		
07702001	**Good identification**					**07702001**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07702002	**Good identification**					**07702002**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07702003	**Good identification**					**07702003**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07702020	**Good identification**					**07702020**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07702021	**Good identification**					**07702021**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07702022	**Good identification**					**07702022**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07702023	**Good identification**					**07702023**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07702100	**Good identification**					**07702100**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07702101	**Good identification**					**07702101**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07702102	**Low discrimination**					**07702102**
Actinomyces naeslundii		89.59	0.44	FRU 95		
Actinomyces viscosus		5.80	0.00	GAL 96		
Propionibacterium propionicum		4.61	0.12			
07702103	**Good identification**					**07702103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	FRU 95		
07702120	**Good identification**					**07702120**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07702121	**Good identification**					**07702121**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07702122	**Good identification**					**07702122**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07702123	**Good identification**					**07702123**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07703000	**Very good identification**					**07703000**
Actinomyces naeslundii		98.26	0.60	FRU 95		
Propionibacterium granulosum		1.74	0.16			
07703001	**Very good identification**					**07703001**
Actinomyces naeslundii		100.0	0.63	FRU 95		
07703002	**Very good identification**					**07703002**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07703003	**Very good identification**					**07703003**
Actinomyces naeslundii		100.0	0.63	FRU 95		
07703010	**Good identification**					**07703010**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07703011	**Good identification**					**07703011**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07703012	**Good identification**					**07703012**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07703013	**Good identification**					**07703013**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07703020	**Very good identification**					**07703020**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07703021	**Very good identification**					**07703021**
Actinomyces naeslundii		100.0	0.63	FRU 95		
07703022	**Very good identification**					**07703022**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07703023	**Very good identification**					**07703023**
Actinomyces naeslundii		100.0	0.63	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07703030	**Good identification**					**07703030**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07703031	**Good identification**					**07703031**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07703032	**Good identification**					**07703032**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07703033	**Good identification**					**07703033**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07703040	**Good identification**					**07703040**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07703041	**Good identification**					**07703041**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07703042	**Good identification**					**07703042**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07703043	**Good identification**					**07703043**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07703060	**Good identification**					**07703060**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07703061	**Good identification**					**07703061**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07703062	**Good identification**					**07703062**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07703063	**Good identification**					**07703063**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07703100	**Very good identification**					**07703100**
					GRA	
Actinomyces naeslundii		99.71	0.68	FRU 95	99	
Fusobacterium mortiferum		0.29	0.13		1	
07703101	**Very good identification**					**07703101**
Actinomyces naeslundii		100.0	0.71	FRU 95		
07703102	**Very good identification**					**07703102**
Actinomyces naeslundii		98.31	0.68	FRU 95		
Bifidobacterium adolescentis		1.03	0.19			
Actinomyces israelii		0.24	0.14			
07703103	**Very good identification**					**07703103**
Actinomyces naeslundii		100.0	0.71	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07703110	**Good identification**					**07703110**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07703111	**Good identification**					**07703111**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07703112	**Low discrimination**					**07703112**
Actinomyces naeslundii		80.44	0.46	FRU 95		
Bifidobacterium adolescentis		19.56	0.20	MLZ 5		
07703113	**Good identification**					**07703113**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07703120	**Very good identification**					**07703120**
					GRA	
Actinomyces naeslundii		95.60	0.68	FRU 95	99	
Prevotella melaninogenica		4.04	0.15		1	
Clostridium perfringens		0.37	0.04		99	
07703121	**Very good identification**					**07703121**
Actinomyces naeslundii		100.0	0.71	FRU 95		
07703122	**Very good identification**					**07703122**
					GRA	
Actinomyces naeslundii		97.14	0.68	FRU 95	99	
Prevotella oralis		1.15	0.09		1	
Prevotella melaninogenica		0.78	0.03		1	
07703123	**Very good identification**					**07703123**
Actinomyces naeslundii		100.0	0.71	FRU 95		
07703130	**Good identification**					**07703130**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07703131	**Good identification**					**07703131**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07703132	**Good identification**					**07703132**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07703133	**Good identification**					**07703133**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07703140	**Good identification**					**07703140**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07703141	**Very good identification**					**07703141**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07703142	**Good identification**					**07703142**
Actinomyces naeslundii		98.80	0.55	FRU 95		
Propionibacterium propionicum		1.20	0.12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07703143	**Very good identification**					**07703143**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07703151	**Good identification**					**07703151**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07703153	**Good identification**					**07703153**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07703160	**Good identification**					**07703160**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07703161	**Very good identification**					**07703161**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07703162	**Good identification**					**07703162**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07703163	**Very good identification**					**07703163**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07703171	**Good identification**					**07703171**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07703173	**Good identification**					**07703173**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07703301	**Good identification**					**07703301**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07703303	**Good identification**					**07703303**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07703321	**Good identification**					**07703321**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07703323	**Good identification**					**07703323**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07703501	**Good identification**					**07703501**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07703503	**Good identification**					**07703503**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07703521	**Good identification**					**07703521**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07703523	**Good identification**					**07703523**
Actinomyces naeslundii		100.0	0.37	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07705112	**Good identification**					**07705112**
Bifidobacterium adolescentis		100.0	0.36	FRU 99		
07705132	**Low discrimination**					**07705132**
					GRA	
Parabacteroides distasonis		86.26	0.36	FRU 99	1	
Bifidobacterium longum subsp. longum		10.00	0.15		99	
Bifidobacterium longum subsp. infantis		3.75	-.04		99	
07705322	**Low discrimination**					**07705322**
Parabacteroides distasonis		84.86	0.48	FRU 99		
Prevotella oralis		13.15	0.22			
Bacteroides vulgatus		0.87	-.04			
07705332	**Good identification**					**07705332**
Parabacteroides distasonis		96.82	0.44	FRU 99		
Clostridium clostridioforme		3.18	0.13			
07705500	**Low discrimination**					**07705500**
					GRA	
Bifidobacterium longum subsp. longum		83.45	0.35	SAL 1	99	
Bacteroides vulgatus		4.18	-.01	FRU 90	1	
Bifidobacterium adolescentis		4.00	0.14		99	
07705502	**Low discrimination**					**07705502**
Bifidobacterium adolescentis		81.08	0.48	FRU 99		
Bifidobacterium longum subsp. longum		17.09	0.35			
Lactobacillus fermentum		0.80	0.20			
07705510	**Good identification**					**07705510**
Bifidobacterium longum subsp. longum		96.67	0.39	SAL 1		
Bifidobacterium adolescentis		3.33	0.16	FRU 90		
07705512	**Low discrimination**					**07705512**
Bifidobacterium adolescentis		76.04	0.50	FRU 99		
Bifidobacterium longum subsp. longum		22.33	0.39			
Clostridium clostridioforme		1.64	0.18			
07705522	**Low discrimination**					**07705522**
					GRA	
Bifidobacterium longum subsp. longum		82.87	0.46	SAL 1	99	
Parabacteroides distasonis		14.00	0.37	FRU 90	1	
Bacteroides vulgatus		1.04	-.01		1	
07705530	**Good identification**					**07705530**
					GRA	
Bifidobacterium longum subsp. longum		99.27	0.50	SAL 1	99	
Bacteroides vulgatus		0.73	-.01	FRU 90	1	
07705532	**Acceptable identification**					**07705532**
					GRA	
Bifidobacterium longum subsp. longum		92.56	0.50	SAL 1	99	
Parabacteroides distasonis		4.94	0.32	FRU 90	1	
Clostridium clostridioforme		1.70	0.18		50	
07705720	**Good identification**					**07705720**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		99.68	0.49	FRU 99	1		
Eubacterium contortum		0.19	0.10		99		
Bifidobacterium longum subsp. longum		0.09	0.11		99		
07705722	**Low discrimination**						**07705722**
Parabacteroides distasonis		74.55	0.45	FRU 99			
Bacteroides vulgatus		16.58	0.15				
Clostridium clostridioforme		7.39	0.22				
07705732	**Low discrimination**						**07705732**
					GRA		
Parabacteroides distasonis		52.59	0.40	FRU 99	1		
Clostridium clostridioforme		44.10	0.33		50		
Bifidobacterium longum subsp. longum		3.32	0.15		99		
07707101	**Good identification**						**07707101**
Actinomyces naeslundii		100.0	0.37	FRU 95			
07707103	**Good identification**						**07707103**
Actinomyces naeslundii		100.0	0.37	FRU 95			
07707121	**Good identification**						**07707121**
Actinomyces naeslundii		100.0	0.37	FRU 95			
07707123	**Good identification**						**07707123**
Actinomyces naeslundii		100.0	0.37	FRU 95			
07707502	**Good identification**						**07707502**
Bifidobacterium adolescentis		95.42	0.47	FRU 99			
Actinomyces israelii		4.59	0.31				
07707512	**Good identification**						**07707512**
Bifidobacterium adolescentis		99.42	0.48	FRU 99			
Actinomyces israelii		0.58	0.17				
07710000	**Low discrimination**						**07710000**
					GRA	CAT	
Fusobacterium mortiferum		74.14	0.72	MLT 11	1	5	
Prevotella intermedia		15.41	0.49		1	1	
Propionibacterium granulosum		7.40	0.52		99	90	
07710001	**Low discrimination**						**07710001**
					GRA		
Actinomyces naeslundii		64.56	0.57	GAL 96	99		
Fusobacterium mortiferum		24.90	0.38		1		
Prevotella intermedia		5.18	0.15		1		
07710002	**Low discrimination**						**07710002**
					GRA	CAT	
Actinomyces viscosus		68.78	0.52	NIT 99	99	76	
Fusobacterium mortiferum		21.96	0.56		1	5	
Propionibacterium granulosum		4.93	0.41		99	90	
07710003	**Low discrimination**						**07710003**
					GRA		
Actinomyces viscosus		80.94	0.44	NIT 99	99		
Actinomyces naeslundii		18.28	0.57		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.78	0.22		1		
07710010	**Low discrimination**						**07710010**
					GRA	CAT	
Propionibacterium granulosum		50.90	0.37	SAL 1	99	90	
Fusobacterium mortiferum		37.80	0.38	MLZ 12	1	5	
Prevotella intermedia		7.86	0.15		1	1	
07710020	**Low discrimination**						**07710020**
					GRA	CAT	
Propionibacterium granulosum		39.09	0.41	SAL 1	99	90	
Actinomyces naeslundii		27.37	0.54		99	25	
Fusobacterium mortiferum		15.83	0.38		1	5	
07710021	**Very good identification**						**07710021**
Actinomyces naeslundii		100.0	0.57	GAL 96			
07710022	**Low discrimination**						**07710022**
					GRA	CAT	
Actinomyces naeslundii		42.73	0.54	GAL 96	99	25	
Prevotella oralis		22.24	0.23		1	1	
Propionibacterium granulosum		15.25	0.31		99	90	
07710023	**Very good identification**						**07710023**
Actinomyces naeslundii		95.72	0.57	GAL 96			
Actinomyces viscosus		4.28	0.09				
07710041	**Good identification**						**07710041**
Actinomyces naeslundii		100.0	0.44	GAL 96			
07710043	**Low discrimination**						**07710043**
Actinomyces naeslundii		79.78	0.44	GAL 96			
Actinomyces viscosus		20.22	0.09				
07710061	**Good identification**						**07710061**
Actinomyces naeslundii		100.0	0.44	GAL 96			
07710062	**Low discrimination**						**07710062**
Actinomyces naeslundii		84.43	0.41	GAL 96			
Clostridium perfringens		15.57	0.06				
07710063	**Good identification**						**07710063**
Actinomyces naeslundii		100.0	0.44	GAL 96			
07710100	**Very good identification**						**07710100**
					GRA		
Fusobacterium mortiferum		97.27	0.78	MLT 11	1		
Actinomyces naeslundii		2.07	0.61		99		
Actinomyces viscosus		0.26	0.22		99		
07710101	**Low discrimination**						**07710101**
					GRA		
Actinomyces naeslundii		74.40	0.64	GAL 96	99		
Fusobacterium mortiferum		23.52	0.44		1		
Actinomyces viscosus		2.08	0.14		99		
07710102	**Low discrimination**						**07710102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		64.02	0.56	NIT 99	99	
Fusobacterium mortiferum		26.83	0.62		1	
Actinomyces naeslundii		5.14	0.61		99	
07710103	**Low discrimination**					**07710103**
					GRA	
Actinomyces viscosus		72.70	0.48	NIT 99	99	
Actinomyces naeslundii		26.29	0.64		99	
Fusobacterium mortiferum		0.92	0.28		1	
07710110	**Acceptable identification**					**07710110**
					GRA	
Fusobacterium mortiferum		90.01	0.44	MLT 11	1	
Actinomyces naeslundii		9.99	0.40	MLZ 1	99	
07710111	**Good identification**					**07710111**
Actinomyces naeslundii		100.0	0.43	GAL 96		
07710113	**Low discrimination**					**07710113**
Actinomyces naeslundii		65.33	0.43	GAL 96		
Actinomyces viscosus		34.67	0.14	MLZ 5		
07710120	**Low discrimination**					**07710120**
					GRA	
Actinomyces naeslundii		27.84	0.61	GAL 96	99	
Clostridium perfringens		26.05	0.39		99	
Prevotella melaninogenica		25.94	0.32		1	
07710121	**Very good identification**					**07710121**
					GRA	
Actinomyces naeslundii		98.76	0.64	GAL 96	99	
Clostridium perfringens		0.62	0.04		99	
Prevotella melaninogenica		0.62	-.03		1	
07710122	**Low discrimination**					**07710122**
					GRA	
Prevotella oralis		50.59	0.44	GAL 99	1	
Actinomyces naeslundii		17.66	0.61		99	
Bifidobacterium longum subsp. infantis		16.32	0.32		99	
07710123	**Good identification**					**07710123**
					GRA	
Actinomyces naeslundii		94.93	0.64	GAL 96	99	
Actinomyces viscosus		2.65	0.14		99	
Prevotella oralis		1.83	0.09		1	
07710130	**Low discrimination**					**07710130**
					GRA	
Actinomyces naeslundii		73.62	0.40	GAL 96	99	
Clostridium perfringens		13.22	0.04	MLZ 5	99	
Prevotella melaninogenica		13.16	-.03		1	
07710131	**Good identification**					**07710131**
Actinomyces naeslundii		100.0	0.43	GAL 96		
07710132	**Low discrimination**					**07710132**
					GRA	
Actinomyces naeslundii		44.55	0.40	GAL 96	99	
Prevotella oralis		24.50	0.09	MLZ 5	1	
Parabacteroides distasonis		16.53	0.19		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07710133	**Good identification**						**07710133**
Actinomyces naeslundii		100.0	0.43	GAL 96			
07710140	**Low discrimination**						**07710140**
					GRA		
Fusobacterium mortiferum		72.27	0.44	MLT 11	1		
Actinomyces naeslundii		26.89	0.49	SOR 1	99		
Clostridium perfringens		0.84	0.00		99		
07710141	**Good identification**						**07710141**
Actinomyces naeslundii		100.0	0.52	GAL 96			
07710142	**Low discrimination**						**07710142**
					GRA		
Actinomyces naeslundii		48.30	0.49	GAL 96	99		
Actinomyces viscosus		34.41	0.22		99		
Fusobacterium mortiferum		14.42	0.28		1		
07710143	**Low discrimination**						**07710143**
Actinomyces naeslundii		86.33	0.52	GAL 96			
Actinomyces viscosus		13.67	0.14				
07710161	**Good identification**						**07710161**
Actinomyces naeslundii		100.0	0.52	GAL 96			
07710162	**Low discrimination**						**07710162**
					GRA		
Actinomyces naeslundii		59.10	0.49	GAL 96	99		
Clostridium perfringens		28.09	0.21		99		
Prevotella oralis		9.69	0.09		1		
07710163	**Good identification**						**07710163**
Actinomyces naeslundii		100.0	0.52	GAL 96			
07710200	**Low discrimination**						**07710200**
					GRA	CAT	
Fusobacterium mortiferum		55.86	0.38	MLT 11	1	5	
Propionibacterium granulosum		17.06	0.26	RHA 1	99	90	
Actinomyces odontolyticus		15.47	0.23		99	10	
07710300	**Acceptable identification**						**07710300**
					GRA		
Fusobacterium mortiferum		94.45	0.44	MLT 11	1		
Actinomyces odontolyticus		3.54	0.14	RHA 1	99		
Actinomyces naeslundii		2.01	0.27		99		
07710322	**Low discrimination**						**07710322**
Prevotella oralis		89.72	0.37	GAL 99			
Parabacteroides distasonis		7.95	0.32				
Clostridium clostridioforme		0.91	0.10				
07710400	**Low discrimination**						**07710400**
					COC	GRA	
Fusobacterium mortiferum		63.75	0.38	MLT 11	1	1	
Prevotella intermedia		13.25	0.15	ARA 1	1	1	
Gemella morbillorum		12.09	0.15		99	99	
07710500	**Acceptable identification**						**07710500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		91.73	0.44	MLT 11	1		
Lactobacillus fermentum		4.58	0.24	ARA 1	99		
Actinomyces naeslundii		1.95	0.27		99		
07710512	**Low discrimination**						**07710512**
Bifidobacterium adolescentis		65.37	0.36	GAL 99			
Clostridium clostridioforme		30.74	0.27	XYL 87			
Bifidobacterium longum subsp. longum		3.89	0.13				
07710712	**Good identification**						**07710712**
Clostridium clostridioforme		100.0	0.42	TRE 88			
07710732	**Good identification**						**07710732**
Clostridium clostridioforme		96.10	0.42	TRE 88			
Parabacteroides distasonis		3.90	0.24	XYL 91			
07711000	**Low discrimination**						**07711000**
					GRA		
Fusobacterium mortiferum		63.13	0.75	MLT 11	1		
Actinomyces naeslundii		17.64	0.77		99		
Prevotella intermedia		8.75	0.49		1		
07711001	**Very good identification**						**07711001**
					GRA		
Actinomyces naeslundii		96.92	0.80		99		
Fusobacterium mortiferum		2.34	0.41		1		
Prevotella intermedia		0.32	0.15		1		
07711002	**Low discrimination**						**07711002**
					GRA		
Actinomyces naeslundii		40.86	0.77		99		
Actinomyces viscosus		20.79	0.48		99		
Fusobacterium mortiferum		16.25	0.59		1		
07711003	**Low discrimination**						**07711003**
					GRA		
Actinomyces naeslundii		89.44	0.80		99		
Actinomyces viscosus		10.11	0.40		99		
Fusobacterium mortiferum		0.24	0.25		1		
07711010	**Low discrimination**						**07711010**
					GRA	CAT	
Actinomyces naeslundii		34.90	0.55	MLZ 5	99	25	
Fusobacterium mortiferum		23.97	0.41		1	5	
Propionibacterium granulosum		21.52	0.37		99	90	
07711011	**Very good identification**						**07711011**
Actinomyces naeslundii		100.0	0.58	MLZ 5			
07711012	**Low discrimination**						**07711012**
					CAT		
Actinomyces naeslundii		56.00	0.55	MLZ 5	25		
Lactobacillus jensenii		16.20	0.25		1		
Propionibacterium granulosum		8.63	0.27		90		
07711013	**Very good identification**						**07711013**
Actinomyces naeslundii		97.88	0.58	MLZ 5			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		2.12	0.06			
07711020	**Low discrimination**					**07711020**
					GRA	
Clostridium perfringens		51.35	0.58	LAC 88	99	
Actinomyces naeslundii		34.28	0.77		99	
Prevotella oralis		3.87	0.35		1	
07711021	**Very good identification**					**07711021**
					GRA	
Actinomyces naeslundii		98.85	0.80		99	
Clostridium perfringens		1.00	0.24		99	
Prevotella oralis		0.08	0.01		1	
07711022	**Low discrimination**					**07711022**
					GRA	
Prevotella oralis		52.13	0.57	LAC 94	1	
Actinomyces naeslundii		24.28	0.77		99	
Clostridium perfringens		17.12	0.52		99	
07711023	**Very good identification**					**07711023**
					GRA	
Actinomyces naeslundii		97.94	0.80		99	
Prevotella oralis		1.42	0.23		1	
Clostridium perfringens		0.47	0.18		99	
07711030	**Low discrimination**					**07711030**
					CAT	
Actinomyces naeslundii		62.77	0.55	MLZ 5	25	
Clostridium perfringens		18.05	0.24		1	
Propionibacterium granulosum		9.68	0.27		90	
07711031	**Very good identification**					**07711031**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07711032	**Low discrimination**					**07711032**
					GRA	
Actinomyces naeslundii		57.51	0.55	MLZ 5	99	
Prevotella oralis		23.70	0.23		1	
Clostridium perfringens		7.78	0.18		99	
07711033	**Very good identification**					**07711033**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07711040	**Low discrimination**					**07711040**
					GRA	
Actinomyces naeslundii		76.05	0.64		99	
Fusobacterium mortiferum		15.58	0.41		1	
Clostridium perfringens		3.80	0.20		99	
07711041	**Very good identification**					**07711041**
Actinomyces naeslundii		100.0	0.67			
07711042	**Low discrimination**					**07711042**
Actinomyces naeslundii		89.14	0.64			
Actinomyces viscosus		2.60	0.14			
Clostridium perfringens		2.10	0.14			
07711043	**Very good identification**					**07711043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.36	0.67			
Actinomyces viscosus		0.64	0.06			
07711050	**Good identification**					**07711050**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07711051	**Good identification**					**07711051**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07711052	**Good identification**					**07711052**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
07711053	**Good identification**					**07711053**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07711060	**Low discrimination**					**07711060**
					GRA	
Clostridium perfringens		61.49	0.46	LAC 88	99	
Actinomyces naeslundii		38.03	0.64		99	
Prevotella oralis		0.25	0.01		1	
07711061	**Very good identification**					**07711061**
Actinomyces naeslundii		98.92	0.67			
Clostridium perfringens		1.08	0.12			
07711062	**Low discrimination**					**07711062**
					GRA	
Actinomyces naeslundii		52.36	0.64		99	
Clostridium perfringens		39.84	0.40		99	
Prevotella oralis		6.43	0.23		1	
07711063	**Very good identification**					**07711063**
Actinomyces naeslundii		99.49	0.67			
Clostridium perfringens		0.51	0.06			
07711070	**Low discrimination**					**07711070**
Actinomyces naeslundii		76.32	0.43	MLZ 5		
Clostridium perfringens		23.68	0.12			
07711071	**Good identification**					**07711071**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07711072	**Low discrimination**					**07711072**
Actinomyces naeslundii		87.26	0.43	MLZ 5		
Clostridium perfringens		12.74	0.06			
07711073	**Good identification**					**07711073**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
07711100	**Low discrimination**					**07711100**
					GRA	
Fusobacterium mortiferum		65.73	0.81	MLT 11	1	
Actinomyces naeslundii		22.40	0.85		99	
Bifidobacterium longum subsp. infantis		4.75	0.44		99	
07711101	**Very good identification**					**07711101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Actinomyces naeslundii		97.19	0.88		99	1	
Fusobacterium mortiferum		1.92	0.47		1	1	
Clostridium chauvoei		0.49	0.26		99	99	
07711102	**Low discrimination**						**07711102**
					GRA		
Bifidobacterium longum subsp. infantis		44.28	0.62	ESL 99	99		
Actinomyces naeslundii		20.65	0.85		99		
Prevotella oralis		12.84	0.56		1		
07711103	**Good identification**						**07711103**
Actinomyces naeslundii		91.32	0.88				
Actinomyces viscosus		6.45	0.44				
Bifidobacterium longum subsp. infantis		1.32	0.27				
07711110	**Low discrimination**						**07711110**
					GRA		
Actinomyces naeslundii		58.29	0.63	MLZ 5	99		
Fusobacterium mortiferum		32.83	0.47		1		
Bifidobacterium longum subsp. infantis		2.37	0.10		99		
07711111	**Very good identification**						**07711111**
					GRA		
Actinomyces naeslundii		99.62	0.66	MLZ 5	99		
Fusobacterium mortiferum		0.38	0.13		1		
07711112	**Low discrimination**						**07711112**
Bifidobacterium adolescentis		62.19	0.56	XYL 87			
Actinomyces naeslundii		20.70	0.63	ARA 87			
Bifidobacterium longum subsp. infantis		8.52	0.27				
07711113	**Very good identification**						**07711113**
Actinomyces naeslundii		96.47	0.66	MLZ 5			
Bifidobacterium adolescentis		1.95	0.22				
Actinomyces viscosus		1.31	0.10				
07711120	**Low discrimination**						**07711120**
					GRA		
Clostridium perfringens		39.41	0.73		99		
Prevotella melaninogenica		39.24	0.66		1		
Actinomyces naeslundii		10.21	0.85		99		
07711121	**Good identification**						**07711121**
					GRA		
Actinomyces naeslundii		94.28	0.88		99		
Clostridium perfringens		2.45	0.39		99		
Prevotella melaninogenica		2.44	0.32		1		
07711122	**Low discrimination**						**07711122**
					GRA		
Prevotella oralis		60.52	0.78		1		
Bifidobacterium longum subsp. infantis		19.52	0.66		99		
Clostridium perfringens		9.30	0.67		99		
07711123	**Low discrimination**						**07711123**
					GRA		
Actinomyces naeslundii		88.86	0.88		99		
Prevotella oralis		7.07	0.44		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. infantis		2.28	0.32		99	
07711130	**Low discrimination**					**07711130**
					GRA	
Actinomyces naeslundii		36.29	0.63	MLZ 5	99	
Clostridium perfringens		26.88	0.39		99	
Prevotella melaninogenica		26.76	0.32		1	
07711131	**Very good identification**					**07711131**
					GRA	
Actinomyces naeslundii		99.01	0.66	MLZ 5	99	
Clostridium perfringens		0.49	0.04		99	
Prevotella melaninogenica		0.49	-.03		1	
07711132	**Low discrimination**					**07711132**
					GRA	
Prevotella oralis		36.11	0.44	MLZ 1	1	
Parabacteroides distasonis		24.38	0.53		1	
Actinomyces naeslundii		15.92	0.63		99	
07711133	**Very good identification**					**07711133**
					GRA	
Actinomyces naeslundii		97.04	0.66	MLZ 5	99	
Prevotella oralis		1.48	0.09		1	
Parabacteroides distasonis		1.00	0.19		1	
07711140	**Low discrimination**					**07711140**
					GRA	
Actinomyces naeslundii		75.38	0.72		99	
Fusobacterium mortiferum		12.66	0.47		1	
Clostridium perfringens		9.71	0.35		99	
07711141	**Very good identification**					**07711141**
					GRA	
Actinomyces naeslundii		99.80	0.75		99	
Fusobacterium mortiferum		0.11	0.13		1	
Clostridium perfringens		0.09	0.00		99	
07711142	**Low discrimination**					**07711142**
Actinomyces naeslundii		68.46	0.72			
Bifidobacterium longum subsp. infantis		8.40	0.27			
Bifidobacterium adolescentis		6.21	0.39			
07711143	**Very good identification**					**07711143**
Actinomyces naeslundii		99.52	0.75			
Actinomyces viscosus		0.40	0.10			
Bifidobacterium longum subsp. infantis		0.08	-.07			
07711150	**Acceptable identification**					**07711150**
					GRA	
Actinomyces naeslundii		91.80	0.50	MLZ 5	99	
Bifidobacterium breve		2.98	0.10		99	
Fusobacterium mortiferum		2.96	0.13		1	
07711151	**Good identification**					**07711151**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07711152	**Low discrimination**					**07711152**
Bifidobacterium adolescentis		60.43	0.40	XYL 87		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		28.71	0.50	ARA 87			
Bifidobacterium breve		9.41	0.27	SOR 11			
07711153	**Good identification**						**07711153**
Actinomyces naeslundii		100.0	0.53	MLZ 5			
07711160	**Low discrimination**						**07711160**
					GRA		
Clostridium perfringens		76.42	0.60		99		
Actinomyces naeslundii		18.34	0.72		99		
Prevotella melaninogenica		4.04	0.32		1		
07711161	**Very good identification**						**07711161**
					GRA		
Actinomyces naeslundii		97.13	0.75		99		
Clostridium perfringens		2.73	0.26		99		
Prevotella melaninogenica		0.14	-.03		1		
07711162	**Low discrimination**						**07711162**
					GRA		
Clostridium perfringens		48.64	0.55		99		
Actinomyces naeslundii		24.81	0.72		99		
Prevotella oralis		16.78	0.44		1		
07711163	**Very good identification**						**07711163**
					GRA		
Actinomyces naeslundii		98.11	0.75		99		
Clostridium perfringens		1.30	0.21		99		
Prevotella oralis		0.45	0.09		1		
07711170	**Low discrimination**						**07711170**
Actinomyces naeslundii		52.65	0.50	MLZ 5			
Clostridium perfringens		42.09	0.26				
Bifidobacterium breve		3.03	0.15				
07711171	**Good identification**						**07711171**
Actinomyces naeslundii		100.0	0.53	MLZ 5			
07711172	**Low discrimination**						**07711172**
Actinomyces naeslundii		45.08	0.50	MLZ 5			
Bifidobacterium breve		26.27	0.32				
Clostridium perfringens		16.96	0.21				
07711173	**Good identification**						**07711173**
Actinomyces naeslundii		100.0	0.53	MLZ 5			
07711200	**Low discrimination**						**07711200**
					GRA	CAT	
Fusobacterium mortiferum		53.14	0.41	MLT 11	1	5	
Actinomyces naeslundii		14.85	0.43	RHA 1	99	25	
Propionibacterium granulosum		10.82	0.26		99	90	
07711201	**Good identification**						**07711201**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07711203	**Low discrimination**						**07711203**
Actinomyces naeslundii		89.84	0.46	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		10.16	0.06			
07711221	**Good identification**					**07711221**
Actinomyces naeslundii		100.0	0.46	RHA 1		
07711222	**Good identification**					**07711222**
					GRA	
Prevotella oralis		96.37	0.51	LAC 94	1	
Parabacteroides distasonis		1.35	0.32		1	
Actinomyces naeslundii		1.06	0.43		99	
07711223	**Low discrimination**					**07711223**
					GRA	
Actinomyces naeslundii		61.98	0.46	RHA 1	99	
Prevotella oralis		38.02	0.17		1	
07711300	**Low discrimination**					**07711300**
					GRA	
Fusobacterium mortiferum		44.27	0.47	MLT 11	1	
Prevotella oralis		20.96	0.28	RHA 1	1	
Actinomyces naeslundii		15.09	0.51		99	
07711301	**Good identification**					**07711301**
					GRA	
Actinomyces naeslundii		98.06	0.54	RHA 1	99	
Fusobacterium mortiferum		1.94	0.13		1	
07711302	**Low discrimination**					**07711302**
					GRA	
Prevotella oralis		82.19	0.50	NAG 95	1	
Bifidobacterium longum subsp. infantis		6.68	0.27		99	
Actinomyces naeslundii		3.11	0.51		99	
07711303	**Low discrimination**					**07711303**
					GRA	
Actinomyces naeslundii		79.19	0.54	RHA 1	99	
Prevotella oralis		14.07	0.15		1	
Actinomyces viscosus		5.59	0.10		99	
07711320	**Low discrimination**					**07711320**
					GRA	
Prevotella oralis		58.99	0.50	BGL 95	1	
Bacteroides vulgatus		18.53	0.35		1	
Clostridium perfringens		8.63	0.39		99	
07711321	**Low discrimination**					**07711321**
					GRA	
Actinomyces naeslundii		77.80	0.54	RHA 1	99	
Prevotella oralis		13.83	0.15		1	
Bacteroides vulgatus		4.34	0.01		1	
07711322	**Good identification**					**07711322**
					GRA	
Prevotella oralis		90.22	0.71		1	
Parabacteroides distasonis		8.00	0.66		1	
Bifidobacterium longum subsp. infantis		0.69	0.32		99	
07711323	**Low discrimination**					**07711323**
					GRA	
Prevotella oralis		71.82	0.37	URE 1	1	
Actinomyces naeslundii		21.27	0.54		99	
Parabacteroides distasonis		6.37	0.32		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07711332	**Low discrimination**					**07711332**
Parabacteroides distasonis		80.34	0.62	XYL 90		
Prevotella oralis		17.00	0.37			
Clostridium clostridioforme		2.35	0.30			
07711340	**Low discrimination**					**07711340**
					GRA	
Actinomyces naeslundii		77.11	0.38	RHA 1	99	
Fusobacterium mortiferum		12.95	0.13		1	
Clostridium perfringens		9.94	0.00		99	
07711341	**Good identification**					**07711341**
Actinomyces naeslundii		100.0	0.41	RHA 1		
07711343	**Good identification**					**07711343**
Actinomyces naeslundii		100.0	0.41	RHA 1		
07711361	**Good identification**					**07711361**
Actinomyces naeslundii		100.0	0.41	RHA 1		
07711362	**Low discrimination**					**07711362**
					GRA	
Prevotella oralis		81.94	0.37	SOR 1	1	
Parabacteroides distasonis		7.26	0.32		1	
Clostridium perfringens		5.60	0.21		99	
07711363	**Low discrimination**					**07711363**
					GRA	
Actinomyces naeslundii		83.81	0.41	RHA 1	99	
Prevotella oralis		16.20	0.03		1	
07711400	**Low discrimination**					**07711400**
					GRA	
Lactobacillus fermentum		74.09	0.54	SAL 1	99	
Fusobacterium mortiferum		15.73	0.41		1	
Actinomyces naeslundii		4.40	0.43		99	
07711401	**Low discrimination**					**07711401**
Actinomyces naeslundii		89.81	0.46	ARA 1		
Lactobacillus fermentum		10.19	0.20			
07711402	**Low discrimination**					**07711402**
Lactobacillus fermentum		76.50	0.54	SAL 1		
Bifidobacterium adolescentis		11.19	0.34			
Actinomyces naeslundii		4.54	0.43			
07711403	**Low discrimination**					**07711403**
Actinomyces naeslundii		81.53	0.46	ARA 1		
Lactobacillus fermentum		9.25	0.20			
Actinomyces viscosus		9.22	0.06			
07711412	**Low discrimination**					**07711412**
Bifidobacterium adolescentis		65.05	0.36	XYL 87		
Clostridium clostridioforme		27.40	0.26	LAC 99		
Bifidobacterium longum subsp. longum		3.87	0.13			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07711420	**Low discrimination**					**07711420**
Lactobacillus fermentum		61.65	0.46	SAL 1		
Clostridium perfringens		16.44	0.24			
Actinomyces naeslundii		10.98	0.43			
07711421	**Good identification**					**07711421**
Actinomyces naeslundii		100.0	0.46	ARA 1		
07711422	**Low discrimination**					**07711422**
					GRA	
Lactobacillus fermentum		53.30	0.46	SAL 1	99	
Prevotella oralis		20.37	0.23		1	
Actinomyces naeslundii		9.49	0.43		99	
07711423	**Good identification**					**07711423**
Actinomyces naeslundii		100.0	0.46	ARA 1		
07711500	**Low discrimination**					**07711500**
					GRA	
Lactobacillus fermentum		50.84	0.58	SAL 1	99	
Bifidobacterium longum subsp. longum		19.29	0.43		99	
Fusobacterium mortiferum		15.44	0.47		1	
07711501	**Acceptable identification**					**07711501**
Actinomyces naeslundii		90.13	0.54	ARA 1		
Lactobacillus fermentum		5.86	0.24			
Bifidobacterium longum subsp. longum		2.23	0.09			
07711502	**Low discrimination**					**07711502**
Bifidobacterium adolescentis		82.06	0.69	XYL 87		
Lactobacillus fermentum		9.25	0.58			
Bifidobacterium longum subsp. longum		3.51	0.43			
07711503	**Low discrimination**					**07711503**
Actinomyces naeslundii		57.07	0.54	ARA 1		
Bifidobacterium adolescentis		32.95	0.34			
Actinomyces viscosus		4.03	0.10			
07711510	**Low discrimination**					**07711510**
Bifidobacterium longum subsp. longum		83.06	0.47	SAL 1		
Bifidobacterium adolescentis		14.09	0.36			
Lactobacillus fermentum		1.30	0.24			
07711512	**Good identification**					**07711512**
Bifidobacterium adolescentis		92.90	0.70	XYL 87		
Bifidobacterium longum subsp. longum		5.53	0.47			
Clostridium clostridioforme		1.32	0.35			
07711513	**Low discrimination**					**07711513**
Bifidobacterium adolescentis		88.17	0.36	XYL 87		
Actinomyces naeslundii		6.58	0.32	URE 1		
Bifidobacterium longum subsp. longum		5.25	0.13			
07711520	**Low discrimination**					**07711520**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium longum subsp. longum		36.68	0.54	SAL 1	99	
Bacteroides vulgatus		30.62	0.38		1	
Clostridium perfringens		9.66	0.39		99	
07711521	**Low discrimination**					**07711521**
					GRA	
Actinomyces naeslundii		79.70	0.54	ARA 1	99	
Bifidobacterium longum subsp. longum		7.87	0.20		99	
Bacteroides vulgatus		6.57	0.04		1	
07711522	**Low discrimination**					**07711522**
					GRA	
Bifidobacterium longum subsp. longum		30.63	0.54	SAL 1	99	
Prevotella oralis		24.69	0.44		1	
Parabacteroides distasonis		18.97	0.54		1	
07711523	**Low discrimination**					**07711523**
					GRA	
Actinomyces naeslundii		77.71	0.54	ARA 1	99	
Bifidobacterium longum subsp. longum		7.67	0.20		99	
Prevotella oralis		6.18	0.09		1	
07711530	**Very good identification**					**07711530**
					GRA	
Bifidobacterium longum subsp. longum		98.67	0.58	SAL 1	99	
Bacteroides vulgatus		0.49	0.04		1	
Actinomyces naeslundii		0.21	0.29		99	
07711532	**Low discrimination**					**07711532**
					GRA	
Bifidobacterium longum subsp. longum		76.39	0.58	SAL 1	99	
Parabacteroides distasonis		14.96	0.50		1	
Clostridium clostridioforme		4.57	0.35		50	
07711540	**Low discrimination**					**07711540**
Actinomyces naeslundii		31.50	0.38	ARA 1		
Bifidobacterium adolescentis		19.11	0.19			
Lactobacillus fermentum		17.42	0.24			
07711541	**Good identification**					**07711541**
Actinomyces naeslundii		100.0	0.41	ARA 1		
07711542	**Good identification**					**07711542**
Bifidobacterium adolescentis		95.78	0.53	XYL 87		
Actinomyces naeslundii		1.60	0.38	SOR 11		
Actinomyces israelii		1.22	0.27			
07711543	**Low discrimination**					**07711543**
Actinomyces naeslundii		71.21	0.41	ARA 1		
Bifidobacterium adolescentis		28.79	0.19			
07711552	**Good identification**					**07711552**
Bifidobacterium adolescentis		99.52	0.54	XYL 87		
Bifidobacterium longum subsp. longum		0.48	0.13	SOR 11		
07711561	**Good identification**					**07711561**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	ARA 1			
07711563	**Good identification**						**07711563**
Actinomyces naeslundii		100.0	0.41	ARA 1			
07711612	**Good identification**						**07711612**
Clostridium clostridioforme		100.0	0.41	TRE 88			
07711632	**Good identification**						**07711632**
Clostridium clostridioforme		95.67	0.41	TRE 88			
Parabacteroides distasonis		4.33	0.24	XYL 91			
07711702	**Low discrimination**						**07711702**
					GRA		
Clostridium clostridioforme		58.69	0.39	TRE 88	50		
Bifidobacterium adolescentis		21.15	0.34	XYL 91	99		
Parabacteroides distasonis		6.72	0.28		1		
07711712	**Low discrimination**						**07711712**
					GRA		
Clostridium clostridioforme		89.61	0.50	TRE 88	50		
Bifidobacterium adolescentis		8.66	0.36	XYL 91	99		
Parabacteroides distasonis		1.21	0.24		1		
07711720	**Good identification**						**07711720**
Bacteroides vulgatus		99.17	0.55	XYL 98			
Parabacteroides distasonis		0.25	0.28				
Prevotella oralis		0.24	0.15				
07711722	**Low discrimination**						**07711722**
Parabacteroides distasonis		75.58	0.63	XYL 90			
Prevotella oralis		14.05	0.37				
Clostridium clostridioforme		6.67	0.39				
07711732	**Low discrimination**						**07711732**
					GRA		
Parabacteroides distasonis		56.59	0.58	XYL 90	1		
Clostridium clostridioforme		42.24	0.50		50		
Bifidobacterium longum subsp. longum		0.97	0.24		99		
07712000	**Low discrimination**						**07712000**
					GRA	CAT	
Propionibacterium granulosum		62.01	0.50	SAL 1	99	90	
Actinomyces naeslundii		27.94	0.59		99	25	
Fusobacterium mortiferum		8.32	0.38		1	5	
07712001	**Very good identification**						**07712001**
Actinomyces naeslundii		98.53	0.62	GAL 96			
Propionibacterium granulosum		1.47	0.16				
07712002	**Low discrimination**						**07712002**
Actinomyces naeslundii		58.84	0.59	GAL 96			
Propionibacterium granulosum		32.66	0.39				
Actinomyces viscosus		6.10	0.18				
07712003	**Very good identification**						**07712003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.75	0.62	GAL 96		
Actinomyces viscosus		2.25	0.09			
07712011	**Good identification**					**07712011**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07712013	**Good identification**					**07712013**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07712020	**Low discrimination**					**07712020**
Actinomyces naeslundii		64.31	0.59	GAL 96		
Propionibacterium granulosum		35.69	0.39			
07712021	**Very good identification**					**07712021**
Actinomyces naeslundii		100.0	0.62	GAL 96		
07712022	**Low discrimination**					**07712022**
Actinomyces naeslundii		87.82	0.59	GAL 96		
Propionibacterium granulosum		12.18	0.29			
07712023	**Very good identification**					**07712023**
Actinomyces naeslundii		100.0	0.62	GAL 96		
07712031	**Good identification**					**07712031**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07712032	**Low discrimination**					**07712032**
Actinomyces naeslundii		73.56	0.37	GAL 96		
Propionibacterium granulosum		26.44	0.14	MLZ 5		
07712033	**Good identification**					**07712033**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07712040	**Low discrimination**					**07712040**
Actinomyces naeslundii		88.73	0.46	GAL 96		
Propionibacterium granulosum		11.28	0.16			
07712041	**Good identification**					**07712041**
Actinomyces naeslundii		100.0	0.49	GAL 96		
07712042	**Good identification**					**07712042**
Actinomyces naeslundii		95.82	0.46	GAL 96		
Propionibacterium propionicum		4.19	0.12			
07712043	**Good identification**					**07712043**
Actinomyces naeslundii		100.0	0.49	GAL 96		
07712060	**Good identification**					**07712060**
Actinomyces naeslundii		100.0	0.46	GAL 96		
07712061	**Good identification**					**07712061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.49	GAL 96			
07712062	**Good identification**						**07712062**
Actinomyces naeslundii		100.0	0.46	GAL 96			
07712063	**Good identification**						**07712063**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07712100	**Low discrimination**						**07712100**
					GRA		
Actinomyces naeslundii		69.88	0.66	GAL 96	99		
Fusobacterium mortiferum		17.07	0.44		1		
Propionibacterium propionicum		12.49	0.43		99		
07712101	**Very good identification**						**07712101**
Actinomyces naeslundii		98.69	0.69	GAL 96			
Propionibacterium propionicum		1.31	0.27				
07712102	**Low discrimination**						**07712102**
Actinomyces naeslundii		72.51	0.66	GAL 96			
Propionibacterium propionicum		19.45	0.46				
Actinomyces viscosus		4.70	0.22				
07712103	**Very good identification**						**07712103**
Actinomyces naeslundii		96.69	0.69	GAL 96			
Propionibacterium propionicum		1.92	0.30				
Actinomyces viscosus		1.39	0.14				
07712110	**Good identification**						**07712110**
Actinomyces naeslundii		100.0	0.44	GAL 96			
07712111	**Good identification**						**07712111**
Actinomyces naeslundii		100.0	0.48	GAL 96			
07712112	**Low discrimination**						**07712112**
Actinomyces naeslundii		73.61	0.44	GAL 96			
Bifidobacterium adolescentis		22.60	0.20	MLZ 5			
Propionibacterium propionicum		3.79	0.12				
07712113	**Good identification**						**07712113**
Actinomyces naeslundii		100.0	0.48	GAL 96			
07712120	**Very good identification**						**07712120**
					GRA		
Actinomyces naeslundii		99.04	0.66	GAL 96	99		
Clostridium perfringens		0.48	0.04		99		
Prevotella melaninogenica		0.48	-.03		1		
07712121	**Very good identification**						**07712121**
Actinomyces naeslundii		100.0	0.69	GAL 96			
07712122	**Very good identification**						**07712122**
					SPO	GRA	
Actinomyces naeslundii		97.11	0.66	GAL 96	1	99	
Prevotella oralis		1.45	0.09		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		0.71	0.05		92	50	
07712123	**Very good identification**						**07712123**
Actinomyces naeslundii		100.0	0.69	GAL	96		
07712130	**Good identification**						**07712130**
Actinomyces naeslundii		100.0	0.44	GAL	96		
07712131	**Good identification**						**07712131**
Actinomyces naeslundii		100.0	0.48	GAL	96		
07712132	**Good identification**						**07712132**
Actinomyces naeslundii		100.0	0.44	GAL	96		
07712133	**Good identification**						**07712133**
Actinomyces naeslundii		100.0	0.48	GAL	96		
07712140	**Low discrimination**						**07712140**
Propionibacterium propionicum		50.33	0.43	NIT	86		
Actinomyces naeslundii		49.67	0.53	SAL	1		
07712141	**Good identification**						**07712141**
Actinomyces naeslundii		93.02	0.57	GAL	96		
Propionibacterium propionicum		6.98	0.27				
07712142	**Low discrimination**						**07712142**
Propionibacterium propionicum		60.31	0.46	NIT	86		
Actinomyces naeslundii		39.69	0.53	SAL	1		
07712143	**Low discrimination**						**07712143**
Actinomyces naeslundii		89.88	0.57	GAL	96		
Propionibacterium propionicum		10.12	0.30				
07712160	**Good identification**						**07712160**
Actinomyces naeslundii		100.0	0.53	GAL	96		
07712161	**Very good identification**						**07712161**
Actinomyces naeslundii		100.0	0.57	GAL	96		
07712162	**Good identification**						**07712162**
Actinomyces naeslundii		98.49	0.53	GAL	96		
Propionibacterium propionicum		1.51	0.12				
07712163	**Very good identification**						**07712163**
Actinomyces naeslundii		100.0	0.57	GAL	96		
07712301	**Good identification**						**07712301**
Actinomyces naeslundii		100.0	0.35	GAL	96		
07712303	**Good identification**						**07712303**
Actinomyces naeslundii		100.0	0.35	GAL	96		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07712321	**Good identification**						**07712321**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07712323	**Good identification**						**07712323**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07712501	**Good identification**						**07712501**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07712503	**Good identification**						**07712503**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07712521	**Good identification**						**07712521**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07712523	**Good identification**						**07712523**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07713000	**Low discrimination**						**07713000**
					GRA	CAT	
Actinomyces naeslundii		89.22	0.82		99	25	
Propionibacterium granulosum		8.25	0.50		99	90	
Fusobacterium mortiferum		1.66	0.41		1	5	
07713001	**Very good identification**						**07713001**
Actinomyces naeslundii		99.94	0.85				
Propionibacterium granulosum		0.06	0.16				
07713002	**Very good identification**						**07713002**
					CAT		
Actinomyces naeslundii		96.05	0.82		25		
Propionibacterium granulosum		2.22	0.39		90		
Actinomyces israelii		0.85	0.38		1		
07713003	**Very good identification**						**07713003**
Actinomyces naeslundii		99.94	0.85				
Actinomyces viscosus		0.06	0.06				
07713010	**Low discrimination**						**07713010**
					CAT		
Actinomyces naeslundii		79.59	0.60	MLZ 5	25		
Propionibacterium granulosum		19.07	0.35		90		
Actinomyces israelii		1.34	0.21		1		
07713011	**Very good identification**						**07713011**
Actinomyces naeslundii		100.0	0.63	MLZ 5			
07713012	**Low discrimination**						**07713012**
					CAT		
Actinomyces naeslundii		89.16	0.60	MLZ 5	25		
Propionibacterium granulosum		5.34	0.25		90		
Bifidobacterium adolescentis		3.25	0.20		1		
07713013	**Very good identification**						**07713013**
Actinomyces naeslundii		100.0	0.63	MLZ 5			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07713020	**Very good identification**						**07713020**
					CAT		
Actinomyces naeslundii		96.86	0.82		25		
Propionibacterium granulosum		2.24	0.39		90		
Clostridium perfringens		0.76	0.24		1		
07713021	**Very good identification**						**07713021**
Actinomyces naeslundii		100.0	0.85				
07713022	**Very good identification**						**07713022**
					GRA	CAT	
Actinomyces naeslundii		97.71	0.82		99	25	
Prevotella oralis		1.09	0.23		1	1	
Propionibacterium granulosum		0.57	0.29		99	90	
07713023	**Very good identification**						**07713023**
Actinomyces naeslundii		100.0	0.85				
07713030	**Good identification**						**07713030**
Actinomyces naeslundii		94.35	0.60	MLZ 5			
Propionibacterium granulosum		5.65	0.25				
07713031	**Very good identification**						**07713031**
Actinomyces naeslundii		100.0	0.63	MLZ 5			
07713032	**Very good identification**						**07713032**
Actinomyces naeslundii		98.52	0.60	MLZ 5			
Propionibacterium granulosum		1.48	0.14				
07713033	**Very good identification**						**07713033**
Actinomyces naeslundii		100.0	0.63	MLZ 5			
07713040	**Very good identification**						**07713040**
					CAT		
Actinomyces naeslundii		98.20	0.69		25		
Actinomyces israelii		1.28	0.28		1		
Propionibacterium granulosum		0.52	0.16		90		
07713041	**Very good identification**						**07713041**
Actinomyces naeslundii		100.0	0.72				
07713042	**Very good identification**						**07713042**
Actinomyces naeslundii		97.91	0.69				
Actinomyces israelii		1.92	0.31				
Propionibacterium propionicum		0.18	0.12				
07713043	**Very good identification**						**07713043**
Actinomyces naeslundii		100.0	0.72				
07713050	**Good identification**						**07713050**
Actinomyces naeslundii		96.43	0.47	MLZ 5			
Actinomyces israelii		3.57	0.14				
07713051	**Good identification**						**07713051**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07713052	**Acceptable identification**					**07713052**
Actinomyces naeslundii		94.74	0.47	MLZ 5		
Actinomyces israelii		5.26	0.17			
07713053	**Good identification**					**07713053**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07713060	**Very good identification**					**07713060**
Actinomyces naeslundii		99.17	0.69			
Clostridium perfringens		0.83	0.12			
07713061	**Very good identification**					**07713061**
Actinomyces naeslundii		100.0	0.72			
07713062	**Very good identification**					**07713062**
Actinomyces naeslundii		99.61	0.69			
Clostridium perfringens		0.39	0.06			
07713063	**Very good identification**					**07713063**
Actinomyces naeslundii		100.0	0.72			
07713070	**Good identification**					**07713070**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
07713071	**Good identification**					**07713071**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07713072	**Good identification**					**07713072**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
07713073	**Good identification**					**07713073**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07713100	**Very good identification**					**07713100**
					GRA	
Actinomyces naeslundii		96.65	0.90		99	
Fusobacterium mortiferum		1.48	0.47		1	
Actinomyces israelii		0.80	0.45		99	
07713101	**Excellent identification**					**07713101**
					GRA	
Actinomyces naeslundii		99.94	0.93		99	
Propionibacterium propionicum		0.06	0.27		99	
Fusobacterium mortiferum		0.01	0.13		1	
07713102	**Good identification**					**07713102**
Actinomyces naeslundii		90.77	0.90			
Bifidobacterium adolescentis		4.95	0.53			
Actinomyces israelii		1.13	0.48			
07713103	**Excellent identification**					**07713103**
Actinomyces naeslundii		99.83	0.93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		0.08	0.30				
Bifidobacterium adolescentis		0.04	0.19				
07713110	**Very good identification**						**07713110**
Actinomyces naeslundii		95.94	0.68	MLZ 5			
Actinomyces israelii		2.27	0.31				
Bifidobacterium adolescentis		1.23	0.20				
07713111	**Very good identification**						**07713111**
Actinomyces naeslundii		100.0	0.71	MLZ 5			
07713112	**Low discrimination**						**07713112**
Bifidobacterium adolescentis		54.22	0.54	XYL 87			
Actinomyces naeslundii		42.81	0.68	ARA 87			
Actinomyces israelii		1.52	0.34				
07713113	**Very good identification**						**07713113**
Actinomyces naeslundii		99.15	0.71	MLZ 5			
Bifidobacterium adolescentis		0.85	0.20				
07713120	**Good identification**						**07713120**
					GRA		
Actinomyces naeslundii		94.88	0.90		99		
Clostridium perfringens		1.91	0.39		99		
Prevotella melaninogenica		1.90	0.32		1		
07713121	**Excellent identification**						**07713121**
					GRA		
Actinomyces naeslundii		99.97	0.93		99		
Clostridium perfringens		0.01	0.04		99		
Prevotella melaninogenica		0.01	-.03		1		
07713122	**Low discrimination**						**07713122**
					SPO	GRA	
Actinomyces naeslundii		88.22	0.90		1	99	
Prevotella oralis		5.43	0.44		1	1	
Clostridium ramosum		2.68	0.39		92	50	
07713123	**Excellent identification**						**07713123**
					SPO	GRA	
Actinomyces naeslundii		99.93	0.93		1	99	
Prevotella oralis		0.04	0.09		1	1	
Clostridium ramosum		0.02	0.05		92	50	
07713130	**Very good identification**						**07713130**
Actinomyces naeslundii		98.72	0.68	MLZ 5			
Bifidobacterium breve		0.52	0.15				
Clostridium perfringens		0.38	0.04				
07713131	**Very good identification**						**07713131**
Actinomyces naeslundii		100.0	0.71	MLZ 5			
07713132	**Good identification**						**07713132**
Actinomyces naeslundii		90.73	0.68	MLZ 5			
Bifidobacterium breve		4.81	0.32				
Bifidobacterium adolescentis		1.16	0.20				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07713133	**Very good identification**					**07713133**
Actinomyces naeslundii		100.0	0.71	MLZ 5		
07713140	**Good identification**					**07713140**
Actinomyces naeslundii		94.08	0.77			
Propionibacterium propionicum		3.97	0.43			
Actinomyces israelii		1.73	0.38			
07713141	**Very good identification**					**07713141**
Actinomyces naeslundii		99.69	0.80			
Propionibacterium propionicum		0.31	0.27			
07713142	**Low discrimination**					**07713142**
Actinomyces naeslundii		87.66	0.77			
Propionibacterium propionicum		5.55	0.46			
Bifidobacterium adolescentis		3.35	0.37			
07713143	**Very good identification**					**07713143**
Actinomyces naeslundii		99.53	0.80			
Propionibacterium propionicum		0.47	0.30			
07713150	**Acceptable identification**					**07713150**
Actinomyces naeslundii		93.57	0.55	MLZ 5		
Actinomyces israelii		4.87	0.24			
Bifidobacterium breve		1.56	0.10			
07713151	**Very good identification**					**07713151**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07713152	**Low discrimination**					**07713152**
Actinomyces naeslundii		46.59	0.55	MLZ 5		
Bifidobacterium adolescentis		41.34	0.39			
Bifidobacterium breve		7.87	0.27			
07713153	**Very good identification**					**07713153**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07713160	**Very good identification**					**07713160**
Actinomyces naeslundii		97.51	0.77			
Clostridium perfringens		2.11	0.26			
Bifidobacterium breve		0.15	0.15			
07713161	**Very good identification**					**07713161**
Actinomyces naeslundii		100.0	0.80			
07713162	**Very good identification**					**07713162**
Actinomyces naeslundii		96.43	0.77			
Bifidobacterium breve		1.52	0.32			
Clostridium perfringens		0.98	0.21			
07713163	**Very good identification**					**07713163**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.80				
07713170	**Good identification**						**07713170**
Actinomyces naeslundii		97.12	0.55	MLZ 5			
Bifidobacterium breve		2.88	0.15				
07713171	**Very good identification**						**07713171**
Actinomyces naeslundii		100.0	0.58	MLZ 5			
07713172	**Low discrimination**						**07713172**
Actinomyces naeslundii		76.91	0.55	MLZ 5			
Bifidobacterium breve		23.09	0.32				
07713173	**Very good identification**						**07713173**
Actinomyces naeslundii		100.0	0.58	MLZ 5			
07713200	**Low discrimination**						**07713200**
					CAT		
Actinomyces naeslundii		75.34	0.48	RHA 1	25		
Propionibacterium granulosum		21.34	0.24		90		
Actinomyces israelii		3.32	0.16		1		
07713201	**Good identification**						**07713201**
Actinomyces naeslundii		100.0	0.51	RHA 1			
07713202	**Low discrimination**						**07713202**
					CAT		
Actinomyces naeslundii		87.95	0.48	RHA 1	25		
Propionibacterium granulosum		6.23	0.14		90		
Actinomyces israelii		5.82	0.19		1		
07713203	**Good identification**						**07713203**
Actinomyces naeslundii		100.0	0.51	RHA 1			
07713220	**Acceptable identification**						**07713220**
Actinomyces naeslundii		93.39	0.48	RHA 1			
Propionibacterium granulosum		6.61	0.14				
07713221	**Good identification**						**07713221**
Actinomyces naeslundii		100.0	0.51	RHA 1			
07713222	**Low discrimination**						**07713222**
					SPO	GRA	
Actinomyces naeslundii		62.96	0.48	RHA 1	1	99	
Prevotella oralis		29.84	0.17		1	1	
Clostridium ramosum		7.20	0.07		92	50	
07713223	**Good identification**						**07713223**
Actinomyces naeslundii		100.0	0.51	RHA 1			
07713240	**Good identification**						**07713240**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07713241	**Good identification**						**07713241**
Actinomyces naeslundii		100.0	0.38	RHA 1			

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
07713242	**Good identification**						**07713242**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07713243	**Good identification**						**07713243**
Actinomyces naeslundii		100.0	0.38	RHA 1			
07713260	**Good identification**						**07713260**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07713261	**Good identification**						**07713261**
Actinomyces naeslundii		100.0	0.38	RHA 1			
07713262	**Good identification**						**07713262**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07713263	**Good identification**						**07713263**
Actinomyces naeslundii		100.0	0.38	RHA 1			
07713300	**Acceptable identification**						**07713300**
					GRA		
Actinomyces naeslundii		92.83	0.56	RHA 1	99		
Actinomyces israelii		5.75	0.26		99		
Fusobacterium mortiferum		1.42	0.13		1		
07713301	**Very good identification**						**07713301**
Actinomyces naeslundii		100.0	0.59	RHA 1			
07713302	**Low discrimination**						**07713302**
					GRA		
Actinomyces naeslundii		72.96	0.56	RHA 1	99		
Prevotella oralis		10.02	0.15		1		
Actinomyces israelii		6.78	0.29		99		
07713303	**Very good identification**						**07713303**
Actinomyces naeslundii		100.0	0.59	RHA 1			
07713311	**Good identification**						**07713311**
Actinomyces naeslundii		100.0	0.37	RHA 1			
07713313	**Good identification**						**07713313**
Actinomyces naeslundii		100.0	0.37	RHA 1			
07713320	**Low discrimination**						**07713320**
					SPO	GRA	
Actinomyces naeslundii		45.14	0.56	RHA 1	1	99	
Clostridium ramosum		44.91	0.31		92	50	
Prevotella oralis		6.20	0.15		1	1	
07713321	**Very good identification**						**07713321**
Actinomyces naeslundii		100.0	0.59	RHA 1			
07713322	**Low discrimination**						**07713322**
					SPO	GRA	
Clostridium ramosum		50.61	0.41	TRE 94	92	50	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		33.18	0.37	CEL 99	1	1	
Actinomyces naeslundii		12.72	0.56		1	99	
07713323	**Very good identification**						**07713323**
					SPO	GRA	
Actinomyces naeslundii		95.75	0.59	RHA 1	1	99	
Clostridium ramosum		2.57	0.07		92	50	
Prevotella oralis		1.68	0.03		1	1	
07713331	**Good identification**						**07713331**
Actinomyces naeslundii		100.0	0.37	RHA 1			
07713333	**Good identification**						**07713333**
Actinomyces naeslundii		100.0	0.37	RHA 1			
07713340	**Low discrimination**						**07713340**
Actinomyces naeslundii		87.98	0.43	RHA 1			
Actinomyces israelii		12.02	0.19				
07713341	**Good identification**						**07713341**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07713342	**Low discrimination**						**07713342**
Actinomyces naeslundii		78.85	0.43	RHA 1			
Actinomyces israelii		16.16	0.22				
Propionibacterium propionicum		4.99	0.12				
07713343	**Good identification**						**07713343**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07713360	**Good identification**						**07713360**
Actinomyces naeslundii		100.0	0.43	RHA 1			
07713361	**Good identification**						**07713361**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07713362	**Low discrimination**						**07713362**
					SPO	GRA	
Actinomyces naeslundii		72.62	0.43	RHA 1	1	99	
Clostridium ramosum		16.54	0.07		92	50	
Prevotella oralis		10.84	0.03		1	1	
07713363	**Good identification**						**07713363**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07713400	**Low discrimination**						**07713400**
					CAT		
Actinomyces naeslundii		60.26	0.48	ARA 1	25		
Actinomyces israelii		28.89	0.33		1		
Propionibacterium granulosum		5.57	0.16		90		
07713401	**Good identification**						**07713401**
Actinomyces naeslundii		100.0	0.51	ARA 1			
07713403	**Good identification**						**07713403**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	ARA 1		
07713420	**Good identification**					**07713420**
Actinomyces naeslundii		100.0	0.48	ARA 1		
07713421	**Good identification**					**07713421**
Actinomyces naeslundii		100.0	0.51	ARA 1		
07713422	**Good identification**					**07713422**
Actinomyces naeslundii		95.61	0.48	ARA 1		
Actinomyces israelii		4.39	0.16			
07713423	**Good identification**					**07713423**
Actinomyces naeslundii		100.0	0.51	ARA 1		
07713441	**Good identification**					**07713441**
Actinomyces naeslundii		100.0	0.38	ARA 1		
07713443	**Good identification**					**07713443**
Actinomyces naeslundii		100.0	0.38	ARA 1		
07713460	**Good identification**					**07713460**
Actinomyces naeslundii		100.0	0.35	ARA 1		
07713461	**Good identification**					**07713461**
Actinomyces naeslundii		100.0	0.38	ARA 1		
07713462	**Good identification**					**07713462**
Actinomyces naeslundii		100.0	0.35	ARA 1		
07713463	**Good identification**					**07713463**
Actinomyces naeslundii		100.0	0.38	ARA 1		
07713500	**Low discrimination**					**07713500**
Actinomyces naeslundii		47.09	0.56	ARA 1		
Actinomyces israelii		31.73	0.44			
Bifidobacterium adolescentis		17.19	0.33			
07713501	**Very good identification**					**07713501**
Actinomyces naeslundii		100.0	0.59	ARA 1		
07713502	**Good identification**					**07713502**
Bifidobacterium adolescentis		94.50	0.67	XYL 87		
Actinomyces israelii		2.64	0.47			
Actinomyces naeslundii		2.62	0.56			
07713503	**Low discrimination**					**07713503**
Actinomyces naeslundii		80.43	0.59	ARA 1		
Bifidobacterium adolescentis		19.57	0.33			
07713511	**Good identification**					**07713511**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ARA 1		
07713512	**Very good identification**					**07713512**
Bifidobacterium adolescentis		99.47	0.69	XYL 87		
Actinomyces israelii		0.34	0.33			
Actinomyces naeslundii		0.12	0.34			
07713520	**Low discrimination**					**07713520**
					GRA	
Actinomyces naeslundii		80.66	0.56	ARA 1	99	
Bifidobacterium longum subsp. longum		6.15	0.20		99	
Bacteroides vulgatus		5.14	0.04		1	
07713521	**Very good identification**					**07713521**
Actinomyces naeslundii		100.0	0.59	ARA 1		
07713522	**Low discrimination**					**07713522**
Actinomyces naeslundii		59.47	0.56	ARA 1		
Bifidobacterium adolescentis		21.71	0.33			
Bifidobacterium longum subsp. longum		4.54	0.20			
07713523	**Very good identification**					**07713523**
Actinomyces naeslundii		100.0	0.59	ARA 1		
07713531	**Good identification**					**07713531**
Actinomyces naeslundii		100.0	0.37	ARA 1		
07713533	**Good identification**					**07713533**
Actinomyces naeslundii		100.0	0.37	ARA 1		
07713540	**Low discrimination**					**07713540**
Actinomyces israelii		54.18	0.37	CEL 93		
Actinomyces naeslundii		36.49	0.43	XYL 92		
Bifidobacterium adolescentis		9.33	0.17			
07713541	**Good identification**					**07713541**
Actinomyces naeslundii		100.0	0.46	ARA 1		
07713542	**Low discrimination**					**07713542**
Bifidobacterium adolescentis		88.50	0.51	XYL 87		
Actinomyces israelii		7.79	0.40	SOR 11		
Actinomyces naeslundii		3.50	0.43			
07713543	**Low discrimination**					**07713543**
Actinomyces naeslundii		85.44	0.46	ARA 1		
Bifidobacterium adolescentis		14.56	0.17			
07713552	**Good identification**					**07713552**
Bifidobacterium adolescentis		98.94	0.53	XYL 87		
Actinomyces israelii		1.06	0.26	SOR 11		
07713560	**Acceptable identification**					**07713560**
Actinomyces naeslundii		91.34	0.43	ARA 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		8.66	0.16				
07713561	**Good identification**						**07713561**
Actinomyces naeslundii		100.0	0.46	ARA 1			
07713562	**Low discrimination**						**07713562**
Actinomyces naeslundii		71.54	0.43	ARA 1			
Bifidobacterium adolescentis		18.29	0.17				
Actinomyces israelii		10.17	0.19				
07713563	**Good identification**						**07713563**
Actinomyces naeslundii		100.0	0.46	ARA 1			
07714000	**Acceptable identification**						**07714000**
					COC	GRA	
Fusobacterium mortiferum		92.44	0.53	MLT 11	1	1	
Prevotella intermedia		2.58	0.15	XYL 7	1	1	
Gemella morbillorum		2.35	0.15		99	99	
07714002	**Low discrimination**						**07714002**
					GRA		
Fusobacterium mortiferum		65.67	0.37	MLT 11	1		
Actinomyces viscosus		27.60	0.18	XYL 7	99		
Actinomyces odontolyticus		3.42	0.09	BGL 10	99		
07714100	**Very good identification**						**07714100**
					GRA		
Fusobacterium mortiferum		97.95	0.59	MLT 11	1		
Bifidobacterium longum subsp. infantis		1.17	0.09	XYL 7	99		
Lactobacillus fermentum		0.39	0.20		99		
07714112	**Low discrimination**						**07714112**
Bifidobacterium adolescentis		76.72	0.36	GAL 99			
Clostridium clostridioforme		23.28	0.24	ARA 87			
07714132	**Low discrimination**						**07714132**
					GRA		
Parabacteroides distasonis		67.28	0.36	GAL 99	1		
Clostridium clostridioforme		29.80	0.24		50		
Bifidobacterium longum subsp. infantis		2.92	-.04		99		
07714312	**Good identification**						**07714312**
Clostridium clostridioforme		99.08	0.39	TRE 88			
Parabacteroides distasonis		0.93	0.10	ARA 94			
07714322	**Low discrimination**						**07714322**
Parabacteroides distasonis		76.55	0.48	GAL 99			
Prevotella oralis		11.86	0.22				
Clostridium clostridioforme		9.80	0.27				
07714332	**Low discrimination**						**07714332**
Clostridium clostridioforme		51.99	0.39	TRE 88			
Parabacteroides distasonis		48.02	0.44	ARA 94			
07714412	**Good identification**						**07714412**
Clostridium clostridioforme		95.46	0.35	TRE 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		4.55	0.16	RHA 88		
07714432	**Good identification**					**07714432**
Clostridium clostridioforme		100.0	0.35	TRE 88		
07714502	**Low discrimination**					**07714502**
Bifidobacterium adolescentis		81.53	0.48	GAL 99		
Clostridium clostridioforme		15.54	0.33			
Bifidobacterium longum subsp. longum		1.56	0.17			
07714512	**Low discrimination**					**07714512**
Bifidobacterium adolescentis		57.57	0.50	GAL 99		
Clostridium clostridioforme		40.89	0.44			
Bifidobacterium longum subsp. longum		1.54	0.21			
07714532	**Low discrimination**					**07714532**
					GRA	
Clostridium clostridioforme		79.82	0.44	TRE 88	50	
Bifidobacterium longum subsp. longum		12.00	0.32	RHA 88	99	
Parabacteroides distasonis		7.05	0.32		1	
07714602	**Good identification**					**07714602**
Clostridium clostridioforme		100.0	0.39	TRE 88		
07714612	**Good identification**					**07714612**
Clostridium clostridioforme		100.0	0.50	TRE 88		
07714622	**Good identification**					**07714622**
Clostridium clostridioforme		98.98	0.39	TRE 88		
Parabacteroides distasonis		1.02	0.11			
07714632	**Good identification**					**07714632**
Clostridium clostridioforme		100.0	0.50	TRE 88		
07714702	**Good identification**					**07714702**
Clostridium clostridioforme		97.77	0.48	TRE 88		
Eubacterium contortum		1.23	0.10			
Bifidobacterium adolescentis		0.71	0.14			
07714712	**Very good identification**					**07714712**
Clostridium clostridioforme		99.81	0.59	TRE 88		
Bifidobacterium adolescentis		0.19	0.16			
07714720	**Good identification**					**07714720**
					GRA	
Bacteroides vulgatus		99.62	0.49	GAL 99	1	
Eubacterium contortum		0.19	0.10		99	
Clostridium clostridioforme		0.15	0.14		50	
07714722	**Low discrimination**					**07714722**
Clostridium clostridioforme		72.80	0.48	TRE 88		
Parabacteroides distasonis		22.25	0.45			
Bacteroides vulgatus		4.95	0.15			
07714732	**Very good identification**					**07714732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		96.51	0.59	TRE 88		
Parabacteroides distasonis		3.49	0.40			
07715000	**Low discrimination**					**07715000**
					GRA	
Fusobacterium mortiferum		64.11	0.56	MLT 11	1	
Lactobacillus fermentum		23.79	0.50	XYL 7	99	
Actinomyces israelii		5.55	0.40		99	
07715001	**Low discrimination**					**07715001**
					GRA	
Actinomyces naeslundii		80.24	0.46	XYL 1	99	
Fusobacterium mortiferum		14.41	0.22		1	
Lactobacillus fermentum		5.35	0.16		99	
07715002	**Low discrimination**					**07715002**
Lactobacillus fermentum		36.76	0.50	SAL 1		
Bifidobacterium longum subsp. infantis		18.55	0.26			
Actinomyces israelii		12.86	0.43			
07715003	**Low discrimination**					**07715003**
Actinomyces naeslundii		84.77	0.46	XYL 1		
Actinomyces viscosus		9.59	0.06			
Lactobacillus fermentum		5.65	0.16			
07715012	**Low discrimination**					**07715012**
Bifidobacterium adolescentis		67.71	0.36	LAC 99		
Clostridium clostridioforme		18.41	0.23	ARA 87		
Actinomyces israelii		11.63	0.29			
07715020	**Low discrimination**					**07715020**
					GRA	
Lactobacillus fermentum		33.38	0.42	SAL 1	99	
Clostridium perfringens		15.16	0.24		99	
Prevotella oralis		13.99	0.20		1	
07715021	**Good identification**					**07715021**
Actinomyces naeslundii		100.0	0.46	XYL 1		
07715022	**Low discrimination**					**07715022**
					GRA	
Prevotella oralis		58.88	0.42	XYL 11	1	
Bifidobacterium longum subsp. infantis		19.90	0.30	LAC 94	99	
Parabacteroides distasonis		8.59	0.40		1	
07715023	**Low discrimination**					**07715023**
					GRA	
Actinomyces naeslundii		80.86	0.46	XYL 1	99	
Prevotella oralis		14.30	0.08		1	
Bifidobacterium longum subsp. infantis		4.83	-.04		99	
07715032	**Low discrimination**					**07715032**
Parabacteroides distasonis		63.74	0.36	LAC 99		
Clostridium clostridioforme		25.30	0.23			
Prevotella oralis		8.20	0.08			
07715042	**Low discrimination**					**07715042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		69.86	0.36	CEL 93		
Bifidobacterium adolescentis		15.80	0.19			
Actinomyces naeslundii		9.16	0.30			
07715100	**Low discrimination**					**07715100**
					GRA	
Fusobacterium mortiferum		41.76	0.62	MLT 11	1	
Bifidobacterium longum subsp. infantis		32.81	0.43	XYL 7	99	
Lactobacillus fermentum		10.84	0.54		99	
07715101	**Low discrimination**					**07715101**
					GRA	
Actinomyces naeslundii		75.59	0.54	XYL 1	99	
Fusobacterium mortiferum		11.13	0.28		1	
Bifidobacterium longum subsp. infantis		8.75	0.09		99	
07715102	**Low discrimination**					**07715102**
					GRA	
Bifidobacterium longum subsp. infantis		61.24	0.60	ESL 99	99	
Bifidobacterium adolescentis		30.23	0.69		99	
Prevotella oralis		2.68	0.40		1	
07715110	**Low discrimination**					**07715110**
					GRA	
Bifidobacterium adolescentis		46.86	0.36	ARA 87	99	
Actinomyces israelii		21.46	0.36	BGL 99	99	
Fusobacterium mortiferum		9.78	0.28		1	
07715112	**Very good identification**					**07715112**
Bifidobacterium adolescentis		95.82	0.70	ARA 87		
Bifidobacterium longum subsp. infantis		1.60	0.26			
Clostridium clostridioforme		0.88	0.32			
07715113	**Acceptable identification**					**07715113**
Bifidobacterium adolescentis		93.06	0.36	ARA 87		
Actinomyces naeslundii		6.94	0.32	URE 1		
07715120	**Low discrimination**					**07715120**
					GRA	
Bacteroides vulgatus		46.64	0.48	RHA 90	1	
Bifidobacterium longum subsp. infantis		31.56	0.47	ARA 93	99	
Prevotella oralis		7.87	0.40		1	
07715121	**Low discrimination**					**07715121**
					GRA	
Actinomyces naeslundii		62.01	0.54	XYL 1	99	
Bacteroides vulgatus		18.85	0.14		1	
Bifidobacterium longum subsp. infantis		12.75	0.13		99	
07715122	**Low discrimination**					**07715122**
					GRA	
Bifidobacterium longum subsp. infantis		51.21	0.64	ESL 99	99	
Prevotella oralis		23.98	0.62		1	
Parabacteroides distasonis		22.11	0.74		1	
07715132	**Good identification**					**07715132**
					GRA	
Parabacteroides distasonis		90.45	0.70		1	
Bifidobacterium longum subsp. infantis		3.93	0.30		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		1.84	0.28		1	
07715133	**Low discrimination**					**07715133**
					GRA	
Parabacteroides distasonis		86.78	0.36	URE 1	1	
Actinomyces naeslundii		9.45	0.32		99	
Bifidobacterium longum subsp. infantis		3.77	-.04		99	
07715140	**Low discrimination**					**07715140**
					GRA	
Actinomyces israelii		61.35	0.43	CEL 93	99	
Fusobacterium mortiferum		10.74	0.28		1	
Actinomyces naeslundii		8.58	0.38		99	
07715141	**Good identification**					**07715141**
Actinomyces naeslundii		100.0	0.41	XYL 1		
07715142	**Low discrimination**					**07715142**
Bifidobacterium adolescentis		72.69	0.53	ARA 87		
Actinomyces israelii		12.98	0.46	SOR 11		
Bifidobacterium longum subsp. infantis		12.03	0.26			
07715143	**Low discrimination**					**07715143**
Actinomyces naeslundii		71.21	0.41	XYL 1		
Bifidobacterium adolescentis		28.79	0.19			
07715152	**Good identification**					**07715152**
Bifidobacterium adolescentis		97.86	0.54	ARA 87		
Actinomyces israelii		2.14	0.32	SOR 11		
07715161	**Good identification**					**07715161**
Actinomyces naeslundii		100.0	0.41	XYL 1		
07715163	**Low discrimination**					**07715163**
Actinomyces naeslundii		89.36	0.41	XYL 1		
Bifidobacterium longum subsp. infantis		10.64	-.04			
07715172	**Low discrimination**					**07715172**
					GRA	
Parabacteroides distasonis		81.71	0.36	SOR 1	1	
Bifidobacterium adolescentis		14.74	0.20		99	
Bifidobacterium longum subsp. infantis		3.55	-.04		99	
07715212	**Good identification**					**07715212**
Clostridium clostridioforme		98.97	0.38	TRE 88		
Parabacteroides distasonis		1.03	0.10	ARA 94		
07715222	**Low discrimination**					**07715222**
Parabacteroides distasonis		47.23	0.48	LAC 99		
Prevotella oralis		46.24	0.35			
Clostridium clostridioforme		5.42	0.27			
07715232	**Low discrimination**					**07715232**
Parabacteroides distasonis		50.29	0.44	LAC 99		
Clostridium clostridioforme		48.79	0.38			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		0.92	0.01			
07715300	**Good identification**					**07715300**
					GRA	
Eubacterium contortum		93.26	0.58	ESL 99	99	
Bacteroides vulgatus		5.49	0.30		1	
Actinomyces israelii		0.33	0.31		99	
07715302	**Low discrimination**					**07715302**
					GRA	
Eubacterium contortum		50.69	0.45	ESL 99	99	
Parabacteroides distasonis		16.65	0.48		1	
Prevotella oralis		13.44	0.34		1	
07715312	**Low discrimination**					**07715312**
					GRA	
Clostridium clostridioforme		67.22	0.47	TRE 88	50	
Parabacteroides distasonis		20.70	0.44	ARA 94	1	
Bifidobacterium adolescentis		10.07	0.36		99	
07715320	**Very good identification**					**07715320**
					GRA	
Bacteroides vulgatus		95.31	0.64	ARA 93	1	
Eubacterium contortum		2.89	0.45		99	
Parabacteroides distasonis		0.95	0.48		1	
07715322	**Low discrimination**					**07715322**
Parabacteroides distasonis		84.56	0.82			
Prevotella oralis		13.10	0.56			
Bacteroides vulgatus		0.87	0.30			
07715323	**Low discrimination**					**07715323**
Parabacteroides distasonis		84.86	0.48	URE 1		
Prevotella oralis		13.15	0.22			
Bacteroides vulgatus		0.87	-.04			
07715332	**Very good identification**					**07715332**
Parabacteroides distasonis		96.51	0.78			
Clostridium clostridioforme		3.17	0.47			
Prevotella oralis		0.28	0.22			
07715333	**Good identification**					**07715333**
Parabacteroides distasonis		96.82	0.44	URE 1		
Clostridium clostridioforme		3.18	0.13			
07715362	**Low discrimination**					**07715362**
Parabacteroides distasonis		84.86	0.48	SOR 1		
Prevotella oralis		13.15	0.22			
Bacteroides vulgatus		0.87	-.04			
07715372	**Good identification**					**07715372**
Parabacteroides distasonis		96.82	0.44	SOR 1		
Clostridium clostridioforme		3.18	0.13			
07715400	**Low discrimination**					**07715400**
					SPO	
Lactobacillus fermentum		75.18	0.50	SAL 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		14.35	0.39		1	
Clostridium butyricum		3.44	0.07		99	
07715402	**Low discrimination**					**07715402**
Bifidobacterium adolescentis		50.70	0.48	LAC 99		
Lactobacillus fermentum		30.42	0.50			
Actinomyces israelii		8.71	0.42			
07715412	**Low discrimination**					**07715412**
Bifidobacterium adolescentis		59.20	0.50	LAC 99		
Clostridium clostridioforme		37.68	0.43			
Bifidobacterium longum subsp. longum		1.58	0.21			
07715422	**Low discrimination**					**07715422**
					GRA	
Lactobacillus fermentum		31.10	0.42	SAL 1	99	
Clostridium clostridioforme		26.56	0.32		50	
Parabacteroides distasonis		22.15	0.37		1	
07715432	**Low discrimination**					**07715432**
					GRA	
Clostridium clostridioforme		77.99	0.43	TRE 88	50	
Bifidobacterium longum subsp. longum		13.08	0.32	RHA 88	99	
Parabacteroides distasonis		7.69	0.32		1	
07715500	**Low discrimination**					**07715500**
					GRA	
Bifidobacterium longum subsp. longum		30.55	0.52	SAL 1	99	
Bacteroides vulgatus		16.83	0.33		1	
Bifidobacterium adolescentis		16.10	0.48		99	
07715502	**Very good identification**					**07715502**
Bifidobacterium adolescentis		95.30	0.83			
Bifidobacterium longum subsp. longum		1.83	0.52			
Lactobacillus fermentum		0.94	0.54			
07715503	**Good identification**					**07715503**
Bifidobacterium adolescentis		95.99	0.48	URE 1		
Bifidobacterium longum subsp. longum		1.84	0.17			
Lactobacillus fermentum		0.95	0.20			
07715510	**Low discrimination**					**07715510**
Bifidobacterium longum subsp. longum		70.19	0.56	SAL 1		
Bifidobacterium adolescentis		26.56	0.50			
Actinomyces israelii		1.49	0.35			
07715512	**Very good identification**					**07715512**
Bifidobacterium adolescentis		95.25	0.84			
Bifidobacterium longum subsp. longum		2.54	0.56			
Clostridium clostridioforme		2.05	0.52			
07715513	**Good identification**					**07715513**
Bifidobacterium adolescentis		95.40	0.50	URE 1		
Bifidobacterium longum subsp. longum		2.55	0.21			
Clostridium clostridioforme		2.05	0.18			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07715520	**Good identification**					**07715520**
					GRA	
Bacteroides vulgatus		92.61	0.67	RHA 90	1	
Bifidobacterium longum subsp. longum		6.79	0.62		99	
Lactobacillus fermentum		0.29	0.46		99	
07715522	**Low discrimination**					**07715522**
					GRA	
Parabacteroides distasonis		55.23	0.71		1	
Bifidobacterium longum subsp. longum		29.73	0.62		99	
Bacteroides vulgatus		4.09	0.33		1	
07715523	**Low discrimination**					**07715523**
					GRA	
Parabacteroides distasonis		56.76	0.37	URE 1	1	
Bifidobacterium longum subsp. longum		30.55	0.28		99	
Bacteroides vulgatus		4.21	-.01		1	
07715530	**Good identification**					**07715530**
					GRA	
Bifidobacterium longum subsp. longum		91.73	0.66	SAL 1	99	
Bacteroides vulgatus		7.42	0.33		1	
Parabacteroides distasonis		0.54	0.32		1	
07715532	**Low discrimination**					**07715532**
					GRA	
Bifidobacterium longum subsp. longum		53.04	0.66	SAL 1	99	
Parabacteroides distasonis		31.16	0.66		1	
Clostridium clostridioforme		10.69	0.52		50	
07715540	**Low discrimination**					**07715540**
					SPO	
Actinomyces israelii		44.69	0.42	CEL 93	1	
Bifidobacterium adolescentis		31.01	0.33		1	
Clostridium butyricum		14.36	0.12		99	
07715542	**Very good identification**					**07715542**
					SPO	
Bifidobacterium adolescentis		97.42	0.67	SOR 11	1	
Actinomyces israelii		2.13	0.45		1	
Clostridium butyricum		0.18	0.05		99	
07715552	**Very good identification**					**07715552**
Bifidobacterium adolescentis		99.34	0.69	SOR 11		
Actinomyces israelii		0.27	0.31			
Bifidobacterium longum subsp. longum		0.22	0.21			
07715562	**Low discrimination**					**07715562**
					GRA	
Parabacteroides distasonis		37.33	0.37	SOR 1	1	
Bifidobacterium adolescentis		32.40	0.33		99	
Bifidobacterium longum subsp. longum		20.09	0.28		99	
07715602	**Good identification**					**07715602**
Clostridium clostridioforme		96.09	0.47	TRE 88		
Eubacterium contortum		1.35	0.10			
Actinomyces israelii		0.99	0.22			
07715612	**Very good identification**					**07715612**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		99.78	0.58	TRE 88		
Bifidobacterium adolescentis		0.22	0.16			
07715620	**Good identification**					**07715620**
					GRA	
Bacteroides vulgatus		99.63	0.49	LAC 99	1	
Eubacterium contortum		0.19	0.10		99	
Clostridium clostridioforme		0.13	0.13		50	
07715622	**Low discrimination**					**07715622**
Clostridium clostridioforme		70.30	0.47	TRE 88		
Parabacteroides distasonis		23.98	0.45			
Bacteroides vulgatus		5.33	0.15			
07715632	**Very good identification**					**07715632**
Clostridium clostridioforme		96.12	0.58	TRE 88		
Parabacteroides distasonis		3.88	0.40			
07715700	**Low discrimination**					**07715700**
					GRA	
Eubacterium contortum		50.76	0.57	ESL 99	99	
Bacteroides vulgatus		48.52	0.49		1	
Clostridium clostridioforme		0.22	0.22		50	
07715702	**Low discrimination**					**07715702**
Clostridium clostridioforme		55.57	0.56	TRE 88		
Eubacterium contortum		23.01	0.44			
Bifidobacterium adolescentis		13.25	0.48			
07715712	**Good identification**					**07715712**
					GRA	
Clostridium clostridioforme		92.67	0.67	TRE 88	50	
Bifidobacterium adolescentis		5.93	0.50		99	
Parabacteroides distasonis		1.12	0.40		1	
07715720	**Very good identification**					**07715720**
					GRA	
Bacteroides vulgatus		99.76	0.83		1	
Eubacterium contortum		0.19	0.44		99	
Parabacteroides distasonis		0.05	0.45		1	
07715721	**Good identification**					**07715721**
					GRA	
Bacteroides vulgatus		99.77	0.49	URE 1	1	
Eubacterium contortum		0.19	0.10		99	
Parabacteroides distasonis		0.05	0.11		1	
07715722	**Low discrimination**					**07715722**
Parabacteroides distasonis		74.90	0.79			
Bacteroides vulgatus		16.65	0.49			
Clostridium clostridioforme		7.42	0.56			
07715723	**Low discrimination**					**07715723**
Parabacteroides distasonis		75.67	0.45	URE 1		
Bacteroides vulgatus		16.83	0.15			
Clostridium clostridioforme		7.50	0.22			
07715730	**Acceptable identification**					**07715730**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		94.28	0.49	MLZ 1		
Parabacteroides distasonis		2.31	0.40			
Clostridium clostridioforme		1.93	0.33			
07715732	**Low discrimination**					**07715732**
					GRA	
Parabacteroides distasonis		54.05	0.74		1	
Clostridium clostridioforme		45.33	0.67		50	
Bifidobacterium longum subsp. longum		0.31	0.32		99	
07715733	**Low discrimination**					**07715733**
Parabacteroides distasonis		54.39	0.40	URE 1		
Clostridium clostridioforme		45.61	0.33			
07715760	**Good identification**					**07715760**
					GRA	
Bacteroides vulgatus		99.77	0.49	SOR 1	1	
Eubacterium contortum		0.19	0.10		99	
Parabacteroides distasonis		0.05	0.11		1	
07715762	**Low discrimination**					**07715762**
Parabacteroides distasonis		75.67	0.45	SOR 1		
Bacteroides vulgatus		16.83	0.15			
Clostridium clostridioforme		7.50	0.22			
07715772	**Low discrimination**					**07715772**
Parabacteroides distasonis		54.39	0.40	SOR 1		
Clostridium clostridioforme		45.61	0.33			
07716101	**Good identification**					**07716101**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07716103	**Good identification**					**07716103**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07716121	**Good identification**					**07716121**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07716123	**Good identification**					**07716123**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07716502	**Good identification**					**07716502**
Bifidobacterium adolescentis		95.42	0.47	GAL 99		
Actinomyces israelii		4.59	0.31			
07716512	**Good identification**					**07716512**
Bifidobacterium adolescentis		98.56	0.48	GAL 99		
Clostridium clostridioforme		0.86	0.10			
Actinomyces israelii		0.58	0.17			
07717000	**Low discrimination**					**07717000**
					GRA	
Actinomyces israelii		84.01	0.53	CEL 93	99	
Actinomyces naeslundii		12.47	0.48		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		1.73	0.22		1	
07717001	**Good identification**					**07717001**
Actinomyces naeslundii		95.66	0.51	XYL 1		
Actinomyces israelii		4.34	0.19			
07717002	**Low discrimination**					**07717002**
Actinomyces israelii		82.86	0.56	CEL 93		
Bifidobacterium adolescentis		8.52	0.33			
Actinomyces naeslundii		8.20	0.48			
07717003	**Acceptable identification**					**07717003**
Actinomyces naeslundii		93.63	0.51	XYL 1		
Actinomyces israelii		6.37	0.22			
07717010	**Good identification**					**07717010**
Actinomyces israelii		95.03	0.39	CEL 93		
Actinomyces naeslundii		4.97	0.26	MLZ 13		
07717012	**Low discrimination**					**07717012**
Actinomyces israelii		53.32	0.42	CEL 93		
Bifidobacterium adolescentis		44.83	0.34	MLZ 13		
Actinomyces naeslundii		1.86	0.26			
07717020	**Low discrimination**					**07717020**
Actinomyces naeslundii		69.92	0.48	XYL 1		
Actinomyces israelii		30.08	0.33			
07717021	**Good identification**					**07717021**
Actinomyces naeslundii		100.0	0.51	XYL 1		
07717022	**Low discrimination**					**07717022**
					GRA	
Actinomyces naeslundii		54.70	0.48	XYL 1	99	
Actinomyces israelii		35.29	0.36		99	
Prevotella oralis		7.48	0.08		1	
07717023	**Good identification**					**07717023**
Actinomyces naeslundii		100.0	0.51	XYL 1		
07717040	**Acceptable identification**					**07717040**
Actinomyces israelii		93.69	0.46	CEL 93		
Actinomyces naeslundii		6.31	0.35			
07717041	**Good identification**					**07717041**
Actinomyces naeslundii		100.0	0.38	XYL 1		
07717042	**Acceptable identification**					**07717042**
Actinomyces israelii		92.80	0.49	CEL 93		
Actinomyces naeslundii		4.17	0.35			
Bifidobacterium adolescentis		3.03	0.17			
07717043	**Low discrimination**					**07717043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		86.96	0.38	XYL 1		
Actinomyces israelii		13.04	0.15			
07717060	**Low discrimination**					**07717060**
Actinomyces naeslundii		51.34	0.35	XYL 1		
Actinomyces israelii		48.66	0.26			
07717061	**Good identification**					**07717061**
Actinomyces naeslundii		100.0	0.38	XYL 1		
07717063	**Good identification**					**07717063**
Actinomyces naeslundii		100.0	0.38	XYL 1		
07717100	**Low discrimination**					**07717100**
Actinomyces israelii		85.38	0.63	CEL 93		
Actinomyces naeslundii		9.02	0.56			
Bifidobacterium adolescentis		3.29	0.33			
07717101	**Good identification**					**07717101**
Actinomyces naeslundii		94.01	0.59	XYL 1		
Actinomyces israelii		6.00	0.29			
07717102	**Low discrimination**					**07717102**
Bifidobacterium adolescentis		69.05	0.67	ARA 87		
Actinomyces israelii		27.14	0.66			
Actinomyces naeslundii		1.91	0.56			
07717103	**Low discrimination**					**07717103**
Actinomyces naeslundii		74.68	0.59	XYL 1		
Bifidobacterium adolescentis		18.17	0.33			
Actinomyces israelii		7.14	0.32			
07717110	**Low discrimination**					**07717110**
Actinomyces israelii		73.94	0.49	CEL 93		
Bifidobacterium adolescentis		23.31	0.34	MLZ 13		
Actinomyces naeslundii		2.75	0.34			
07717111	**Low discrimination**					**07717111**
Actinomyces naeslundii		84.67	0.37	XYL 1		
Actinomyces israelii		15.33	0.15	MLZ 5		
07717112	**Very good identification**					**07717112**
Bifidobacterium adolescentis		95.28	0.69	ARA 87		
Actinomyces israelii		4.58	0.52			
Actinomyces naeslundii		0.11	0.34			
07717120	**Low discrimination**					**07717120**
					GRA	
Actinomyces naeslundii		48.13	0.56	XYL 1	99	
Actinomyces israelii		29.09	0.43		99	
Bacteroides vulgatus		11.30	0.14		1	
07717121	**Very good identification**					**07717121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.59	XYL 1		
07717123	**Very good identification**					**07717123**
Actinomyces naeslundii		98.93	0.59	XYL 1		
Bifidobacterium longum subsp. infantis		1.07	-.04			
07717131	**Good identification**					**07717131**
Actinomyces naeslundii		100.0	0.37	XYL 1		
07717133	**Good identification**					**07717133**
Actinomyces naeslundii		100.0	0.37	XYL 1		
07717140	**Good identification**					**07717140**
Actinomyces israelii		94.33	0.56	CEL 93		
Actinomyces naeslundii		4.52	0.43			
Bifidobacterium adolescentis		1.16	0.17			
07717141	**Low discrimination**					**07717141**
Actinomyces naeslundii		87.68	0.46	XYL 1		
Actinomyces israelii		12.32	0.22			
07717142	**Low discrimination**					**07717142**
Actinomyces israelii		54.28	0.59	CEL 93		
Bifidobacterium adolescentis		43.88	0.51			
Actinomyces naeslundii		1.73	0.43			
07717143	**Low discrimination**					**07717143**
Actinomyces naeslundii		72.40	0.46	XYL 1		
Actinomyces israelii		15.26	0.25			
Bifidobacterium adolescentis		12.34	0.17			
07717150	**Acceptable identification**					**07717150**
Actinomyces israelii		90.89	0.42	CEL 93		
Bifidobacterium adolescentis		9.11	0.19	MLZ 13		
07717152	**Low discrimination**					**07717152**
Bifidobacterium adolescentis		86.87	0.53	ARA 87		
Actinomyces israelii		13.14	0.45	SOR 11		
07717160	**Low discrimination**					**07717160**
Actinomyces israelii		57.12	0.36	CEL 93		
Actinomyces naeslundii		42.88	0.43	NAG 6		
07717161	**Good identification**					**07717161**
Actinomyces naeslundii		100.0	0.46	XYL 1		
07717162	**Low discrimination**					**07717162**
Actinomyces israelii		59.72	0.39	CEL 93		
Actinomyces naeslundii		29.89	0.43	NAG 6		
Bifidobacterium adolescentis		7.64	0.17			
07717163	**Good identification**					**07717163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	XYL 1		
07717202	**Good identification**					**07717202**
Actinomyces israelii		100.0	0.37	CEL 93		
07717300	**Low discrimination**					**07717300**
					GRA	
Actinomyces israelii		64.99	0.44	CEL 93	99	
Eubacterium contortum		33.07	0.24	RHA 7	99	
Bacteroides vulgatus		1.95	-.04		1	
07717302	**Low discrimination**					**07717302**
Actinomyces israelii		69.60	0.47	CEL 93		
Bifidobacterium adolescentis		23.76	0.33	RHA 7		
Eubacterium contortum		4.17	0.11			
07717322	**Low discrimination**					**07717322**
					GRA	
Parabacteroides distasonis		81.27	0.48	MAN 1	1	
Prevotella oralis		12.59	0.22		1	
Actinomyces israelii		2.67	0.27		99	
07717332	**Good identification**					**07717332**
Parabacteroides distasonis		96.82	0.44	MAN 1		
Clostridium clostridioforme		3.18	0.13			
07717340	**Good identification**					**07717340**
Actinomyces israelii		100.0	0.37	CEL 93		
07717342	**Acceptable identification**					**07717342**
Actinomyces israelii		90.21	0.40	CEL 93		
Bifidobacterium adolescentis		9.79	0.17	RHA 7		
07717400	**Good identification**					**07717400**
Actinomyces israelii		97.84	0.52	CEL 93		
Bifidobacterium adolescentis		1.25	0.13			
Lactobacillus fermentum		0.91	0.16			
07717402	**Low discrimination**					**07717402**
Actinomyces israelii		54.14	0.55	CEL 93		
Bifidobacterium adolescentis		45.52	0.47			
Lactobacillus fermentum		0.34	0.16			
07717410	**Acceptable identification**					**07717410**
Actinomyces israelii		90.56	0.37	CEL 93		
Bifidobacterium adolescentis		9.44	0.14	MLZ 13		
07717412	**Low discrimination**					**07717412**
Bifidobacterium adolescentis		86.71	0.48	LAC 99		
Actinomyces israelii		12.61	0.40			
Clostridium clostridioforme		0.68	0.09			
07717440	**Good identification**					**07717440**
Actinomyces israelii		100.0	0.45	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07717442	**Low discrimination**					**07717442**
Actinomyces israelii		78.91	0.48	CEL 93		
Bifidobacterium adolescentis		21.09	0.31			
07717500	**Low discrimination**					**07717500**
					SPO	
Actinomyces israelii		73.60	0.62	CEL 93	1	
Bifidobacterium adolescentis		23.20	0.47		1	
Clostridium butyricum		2.07	0.14		99	
07717502	**Very good identification**					**07717502**
					SPO	
Bifidobacterium adolescentis		95.34	0.81		1	
Actinomyces israelii		4.58	0.65		1	
Clostridium butyricum		0.03	0.07		99	
07717503	**Acceptable identification**					**07717503**
Bifidobacterium adolescentis		94.85	0.47	URE 1		
Actinomyces israelii		4.56	0.31			
Actinomyces naeslundii		0.59	0.25			
07717510	**Low discrimination**					**07717510**
Bifidobacterium adolescentis		70.21	0.48	BGL 99		
Actinomyces israelii		27.23	0.48			
Bifidobacterium longum subsp. longum		2.29	0.21			
07717512	**Very good identification**					**07717512**
Bifidobacterium adolescentis		99.35	0.83			
Actinomyces israelii		0.58	0.51			
Bifidobacterium longum subsp. longum		0.03	0.21			
07717513	**Good identification**					**07717513**
Bifidobacterium adolescentis		99.42	0.48	URE 1		
Actinomyces israelii		0.58	0.17			
07717522	**Low discrimination**					**07717522**
					GRA	
Bifidobacterium adolescentis		62.58	0.47	NAG 1	99	
Actinomyces israelii		19.00	0.44		99	
Parabacteroides distasonis		10.89	0.37		1	
07717532	**Low discrimination**					**07717532**
					GRA	
Bifidobacterium adolescentis		77.97	0.48	NAG 1	99	
Bifidobacterium longum subsp. longum		10.18	0.32		99	
Parabacteroides distasonis		5.98	0.32		1	
07717540	**Acceptable identification**					**07717540**
					SPO	
Actinomyces israelii		90.38	0.55	CEL 93	1	
Bifidobacterium adolescentis		9.06	0.31		1	
Clostridium butyricum		0.57	-.04		99	
07717542	**Low discrimination**					**07717542**
Bifidobacterium adolescentis		86.87	0.66	SOR 11		
Actinomyces israelii		13.14	0.58			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07717550	**Low discrimination**					**07717550**
					COC	
Actinomyces israelii		53.66	0.41	CEL 93	1	
Bifidobacterium adolescentis		43.98	0.33	MLZ 13	1	
Blautia producta		2.36	-.03		99	
07717552	**Very good identification**					**07717552**
Bifidobacterium adolescentis		98.19	0.67	SOR 11		
Actinomyces israelii		1.82	0.44			
07717602	**Low discrimination**					**07717602**
Actinomyces israelii		77.79	0.35	CEL 93		
Clostridium clostridioforme		13.43	0.13	RHA 7		
Bifidobacterium adolescentis		8.78	0.13			
07717700	**Low discrimination**					**07717700**
					GRA	
Actinomyces israelii		49.08	0.43	CEL 93	99	
Eubacterium contortum		24.97	0.23	RHA 7	99	
Bacteroides vulgatus		23.87	0.15		1	
07717702	**Low discrimination**					**07717702**
Bifidobacterium adolescentis		69.61	0.47	RHA 1		
Actinomyces israelii		24.93	0.46			
Clostridium clostridioforme		3.60	0.22			
07717712	**Low discrimination**					**07717712**
Bifidobacterium adolescentis		80.86	0.48	RHA 1		
Clostridium clostridioforme		15.60	0.33			
Actinomyces israelii		3.54	0.32			
07717720	**Good identification**					**07717720**
					GRA	
Bacteroides vulgatus		99.64	0.49	MAN 1	1	
Eubacterium contortum		0.19	0.10		99	
Actinomyces israelii		0.13	0.22		99	
07717722	**Low discrimination**					**07717722**
Parabacteroides distasonis		71.45	0.45	MAN 1		
Bacteroides vulgatus		15.89	0.15			
Clostridium clostridioforme		7.08	0.22			
07717732	**Low discrimination**					**07717732**
					GRA	
Parabacteroides distasonis		52.60	0.40	MAN 1	1	
Clostridium clostridioforme		44.11	0.33		50	
Bifidobacterium adolescentis		2.31	0.14		99	
07717740	**Good identification**					**07717740**
Actinomyces israelii		100.0	0.36	CEL 93		
07717742	**Low discrimination**					**07717742**
Actinomyces israelii		52.98	0.39	CEL 93		
Bifidobacterium adolescentis		47.02	0.31	RHA 7		
07720001	**Good identification**					**07720001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	FRU 95		
07720002	**Low discrimination**					**07720002**
Actinomyces naeslundii		41.95	0.37	FRU 95		
Actinomyces viscosus		32.41	0.11	GAL 96		
Propionibacterium granulosum		17.77	0.15			
07720003	**Low discrimination**					**07720003**
Actinomyces naeslundii		85.35	0.40	FRU 95		
Actinomyces viscosus		14.65	0.02	GAL 96		
07720020	**Low discrimination**					**07720020**
Actinomyces naeslundii		70.25	0.37	FRU 95		
Propionibacterium granulosum		29.75	0.15	GAL 96		
07720021	**Good identification**					**07720021**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07720022	**Good identification**					**07720022**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07720023	**Good identification**					**07720023**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07720100	**Low discrimination**					**07720100**
					GRA	
Fusobacterium mortiferum		58.10	0.35	MLT 11	1	
Actinomyces naeslundii		41.91	0.44	FRU 99	99	
07720101	**Good identification**					**07720101**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07720102	**Low discrimination**					**07720102**
					GRA	
Actinomyces naeslundii		61.10	0.44	FRU 95	99	
Actinomyces viscosus		29.49	0.15	GAL 96	99	
Fusobacterium mortiferum		9.41	0.19		1	
07720103	**Acceptable identification**					**07720103**
Actinomyces naeslundii		90.32	0.48	FRU 95		
Actinomyces viscosus		9.68	0.07	GAL 96		
07720120	**Low discrimination**					**07720120**
Actinomyces naeslundii		85.34	0.44	FRU 95		
Clostridium perfringens		14.66	0.09	GAL 96		
07720121	**Good identification**					**07720121**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07720122	**Acceptable identification**					**07720122**
Actinomyces naeslundii		92.52	0.44	FRU 95		
Clostridium perfringens		7.48	0.03	GAL 96		
07720123	**Good identification**					**07720123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.48	FRU 95			
07721000	**Low discrimination**						**07721000**
					GRA	CAT	
Actinomyces naeslundii		82.98	0.60	FRU 95	99	25	
Fusobacterium mortiferum		8.77	0.32		1	5	
Propionibacterium granulosum		5.86	0.26		99	90	
07721001	**Very good identification**						**07721001**
Actinomyces naeslundii		100.0	0.63	FRU 95			
07721002	**Good identification**						**07721002**
Actinomyces naeslundii		93.27	0.60	FRU 95			
Actinomyces viscosus		1.84	0.07				
Propionibacterium granulosum		1.65	0.15				
07721003	**Very good identification**						**07721003**
Actinomyces naeslundii		99.56	0.63	FRU 95			
Actinomyces viscosus		0.44	-.01				
07721010	**Low discrimination**						**07721010**
					CAT		
Actinomyces naeslundii		76.24	0.38	FRU 95	25		
Propionibacterium granulosum		13.95	0.11	MLZ 5	90		
Lactobacillus jensenii		9.81	0.02		1		
07721011	**Good identification**						**07721011**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07721012	**Good identification**						**07721012**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07721013	**Good identification**						**07721013**
Actinomyces naeslundii		100.0	0.41	FRU 95			
07721020	**Low discrimination**						**07721020**
					CAT		
Actinomyces naeslundii		77.36	0.60	FRU 95	25		
Clostridium perfringens		21.28	0.28		1		
Propionibacterium granulosum		1.37	0.15		90		
07721021	**Very good identification**						**07721021**
Actinomyces naeslundii		100.0	0.63	FRU 95			
07721022	**Low discrimination**						**07721022**
					GRA		
Actinomyces naeslundii		86.77	0.60	FRU 95	99		
Clostridium perfringens		11.23	0.23		99		
Prevotella oralis		1.18	0.03		1		
07721023	**Very good identification**						**07721023**
Actinomyces naeslundii		100.0	0.63	FRU 95			
07721030	**Good identification**						**07721030**
Actinomyces naeslundii		100.0	0.38	FRU 95			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07721031	**Good identification**					**07721031**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07721032	**Good identification**					**07721032**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07721033	**Good identification**					**07721033**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07721040	**Good identification**					**07721040**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07721041	**Good identification**					**07721041**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07721042	**Good identification**					**07721042**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07721043	**Good identification**					**07721043**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07721060	**Low discrimination**					**07721060**
Actinomyces naeslundii		77.11	0.47	FRU 95		
Clostridium perfringens		22.89	0.16			
07721061	**Good identification**					**07721061**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07721062	**Low discrimination**					**07721062**
Actinomyces naeslundii		87.74	0.47	FRU 95		
Clostridium perfringens		12.26	0.11			
07721063	**Good identification**					**07721063**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07721100	**Low discrimination**					**07721100**
					GRA	
Actinomyces naeslundii		89.02	0.68	FRU 95	99	
Fusobacterium mortiferum		7.71	0.38		1	
Clostridium perfringens		1.95	0.17		99	
07721101	**Very good identification**					**07721101**
					SPO	
Actinomyces naeslundii		99.77	0.71	FRU 95	1	
Clostridium chauvoei		0.23	0.03		99	
07721102	**Good identification**					**07721102**
Actinomyces naeslundii		91.94	0.68	FRU 95		
Lactobacillus acidophilus		3.51	0.19			
Actinomyces viscosus		1.13	0.11			
07721103	**Very good identification**					**07721103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.73	0.71	FRU 95		
Actinomyces viscosus		0.27	0.03			
07721110	**Good identification**					**07721110**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07721111	**Good identification**					**07721111**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07721112	**Low discrimination**					**07721112**
					GRA	
Actinomyces naeslundii		80.27	0.46	FRU 95	99	
Bifidobacterium adolescentis		15.90	0.19	MLZ 5	99	
Leptotrichia buccalis		3.83	-.11		1	
07721113	**Good identification**					**07721113**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07721120	**Low discrimination**					**07721120**
					GRA	
Actinomyces naeslundii		56.77	0.68	FRU 95	99	
Clostridium perfringens		40.23	0.43		99	
Prevotella melaninogenica		2.40	0.15		1	
07721121	**Very good identification**					**07721121**
Actinomyces naeslundii		99.53	0.71	FRU 95		
Clostridium perfringens		0.48	0.09			
07721122	**Low discrimination**					**07721122**
					GRA	
Actinomyces naeslundii		64.87	0.68	FRU 95	99	
Clostridium perfringens		21.63	0.38		99	
Prevotella oralis		4.84	0.23		1	
07721123	**Very good identification**					**07721123**
Actinomyces naeslundii		99.78	0.71	FRU 95		
Clostridium perfringens		0.22	0.03			
07721130	**Low discrimination**					**07721130**
Actinomyces naeslundii		88.03	0.46	FRU 95		
Clostridium perfringens		11.97	0.09	MLZ 5		
07721131	**Good identification**					**07721131**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07721132	**Low discrimination**					**07721132**
					GRA	
Actinomyces naeslundii		57.03	0.46	FRU 95	99	
Parabacteroides distasonis		39.32	0.30	MLZ 5	1	
Clostridium perfringens		3.65	0.03		99	
07721133	**Good identification**					**07721133**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07721140	**Good identification**					**07721140**
Actinomyces naeslundii		97.69	0.55	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		2.31	0.05			
07721141	**Very good identification**					**07721141**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07721142	**Good identification**					**07721142**
Actinomyces naeslundii		97.40	0.55	FRU 95		
Lactobacillus acidophilus		2.61	0.04			
07721143	**Very good identification**					**07721143**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07721151	**Good identification**					**07721151**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07721153	**Good identification**					**07721153**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07721160	**Low discrimination**					**07721160**
Actinomyces naeslundii		56.66	0.55	FRU 95		
Clostridium perfringens		43.34	0.31			
07721161	**Very good identification**					**07721161**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07721162	**Low discrimination**					**07721162**
Actinomyces naeslundii		72.11	0.55	FRU 95		
Clostridium perfringens		25.96	0.25			
Lactobacillus acidophilus		1.93	0.04			
07721163	**Very good identification**					**07721163**
Actinomyces naeslundii		100.0	0.58	FRU 95		
07721171	**Good identification**					**07721171**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07721173	**Good identification**					**07721173**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07721301	**Good identification**					**07721301**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07721303	**Good identification**					**07721303**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07721321	**Good identification**					**07721321**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07721322	**Low discrimination**					**07721322**
					SPO	
Parabacteroides distasonis		74.38	0.43	FRU 99	1	
Prevotella oralis		11.76	0.17	XYL 90	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Clostridium ramosum	7.85	0.14		92	
07721323	**Good identification**					**07721323**
	Actinomyces naeslundii	100.0	0.37	FRU 95		
07721332	**Good identification**					**07721332**
	Parabacteroides distasonis	95.51	0.39	FRU 99		
	Clostridium clostridioforme	4.49	0.10	XYL 90		
07721501	**Good identification**					**07721501**
	Actinomyces naeslundii	100.0	0.37	FRU 95		
07721503	**Good identification**					**07721503**
	Actinomyces naeslundii	100.0	0.37	FRU 95		
07721521	**Good identification**					**07721521**
	Actinomyces naeslundii	100.0	0.37	FRU 95		
07721523	**Good identification**					**07721523**
	Actinomyces naeslundii	100.0	0.37	FRU 95		
07721722	**Low discrimination**					**07721722**
	Parabacteroides distasonis	87.56	0.40	FRU 99		
	Clostridium clostridioforme	12.44	0.19	XYL 90		
07722000	**Low discrimination**					**07722000**
	Actinomyces naeslundii	60.30	0.42	FRU 95		
	Propionibacterium granulosum	39.70	0.24	GAL 96		
07722001	**Good identification**					**07722001**
	Actinomyces naeslundii	100.0	0.45	FRU 95		
07722002	**Low discrimination**					**07722002**
	Actinomyces naeslundii	85.87	0.42	FRU 95		
	Propionibacterium granulosum	14.13	0.13	GAL 96		
07722003	**Good identification**					**07722003**
	Actinomyces naeslundii	100.0	0.45	FRU 95		
07722020	**Low discrimination**					**07722020**
	Actinomyces naeslundii	85.87	0.42	FRU 95		
	Propionibacterium granulosum	14.13	0.13	GAL 96		
07722021	**Good identification**					**07722021**
	Actinomyces naeslundii	100.0	0.45	FRU 95		
07722022	**Good identification**					**07722022**
	Actinomyces naeslundii	100.0	0.42	FRU 95		
07722023	**Good identification**					**07722023**
	Actinomyces naeslundii	100.0	0.45	FRU 95		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07722100	**Good identification**					**07722100**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07722101	**Good identification**					**07722101**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07722102	**Good identification**					**07722102**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07722103	**Good identification**					**07722103**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07722120	**Good identification**					**07722120**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07722121	**Good identification**					**07722121**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07722122	**Good identification**					**07722122**
Actinomyces naeslundii		100.0	0.49	FRU 95		
07722123	**Good identification**					**07722123**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07722140	**Good identification**					**07722140**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07722141	**Good identification**					**07722141**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07722142	**Good identification**					**07722142**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07722143	**Good identification**					**07722143**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07722160	**Good identification**					**07722160**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07722161	**Good identification**					**07722161**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07722162	**Good identification**					**07722162**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07722163	**Good identification**					**07722163**
Actinomyces naeslundii		100.0	0.40	FRU 95		
07723000	**Very good identification**					**07723000**
Actinomyces naeslundii		97.33	0.65	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		2.67	0.24			
07723001	**Very good identification**					**07723001**
Actinomyces naeslundii		100.0	0.68	FRU 95		
07723002	**Very good identification**					**07723002**
					CAT	
Actinomyces naeslundii		98.99	0.65	FRU 95	25	
Propionibacterium granulosum		0.68	0.13		90	
Actinomyces israelii		0.33	0.14		1	
07723003	**Very good identification**					**07723003**
Actinomyces naeslundii		100.0	0.68	FRU 95		
07723010	**Acceptable identification**					**07723010**
Actinomyces naeslundii		93.37	0.43	FRU 95		
Propionibacterium granulosum		6.64	0.09	MLZ 5		
07723011	**Good identification**					**07723011**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07723012	**Good identification**					**07723012**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07723013	**Good identification**					**07723013**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07723020	**Very good identification**					**07723020**
Actinomyces naeslundii		99.32	0.65	FRU 95		
Propionibacterium granulosum		0.68	0.13			
07723021	**Very good identification**					**07723021**
Actinomyces naeslundii		100.0	0.68	FRU 95		
07723022	**Very good identification**					**07723022**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07723023	**Very good identification**					**07723023**
Actinomyces naeslundii		100.0	0.68	FRU 95		
07723030	**Good identification**					**07723030**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07723031	**Good identification**					**07723031**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07723032	**Good identification**					**07723032**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07723033	**Good identification**					**07723033**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07723040	**Good identification**					**07723040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	FRU 95		
07723041	**Good identification**					**07723041**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07723042	**Good identification**					**07723042**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07723043	**Good identification**					**07723043**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07723060	**Good identification**					**07723060**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07723061	**Good identification**					**07723061**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07723062	**Good identification**					**07723062**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07723063	**Good identification**					**07723063**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07723100	**Very good identification**					**07723100**
Actinomyces naeslundii		99.69	0.73	FRU 95		
Actinomyces israelii		0.31	0.21			
07723101	**Very good identification**					**07723101**
Actinomyces naeslundii		100.0	0.76	FRU 95		
07723102	**Very good identification**					**07723102**
Actinomyces naeslundii		99.18	0.73	FRU 95		
Actinomyces israelii		0.46	0.24			
Bifidobacterium adolescentis		0.36	0.16			
07723103	**Very good identification**					**07723103**
Actinomyces naeslundii		100.0	0.76	FRU 95		
07723110	**Good identification**					**07723110**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07723111	**Good identification**					**07723111**
Actinomyces naeslundii		100.0	0.54	FRU 95		
07723112	**Acceptable identification**					**07723112**
Actinomyces naeslundii		92.29	0.51	FRU 95		
Bifidobacterium adolescentis		7.71	0.17	MLZ 5		
07723113	**Good identification**					**07723113**
Actinomyces naeslundii		100.0	0.54	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07723120	**Very good identification**					**07723120**
					SPO	
Actinomyces naeslundii		98.48	0.73	FRU 95	1	
Clostridium ramosum		1.16	0.15		92	
Clostridium perfringens		0.36	0.09		84	
07723121	**Very good identification**					**07723121**
Actinomyces naeslundii		100.0	0.76	FRU 95		
07723122	**Very good identification**					**07723122**
					SPO	
Actinomyces naeslundii		95.36	0.73	FRU 95	1	
Clostridium ramosum		4.48	0.25		92	
Clostridium perfringens		0.17	0.03		84	
07723123	**Very good identification**					**07723123**
Actinomyces naeslundii		100.0	0.76	FRU 95		
07723130	**Good identification**					**07723130**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07723131	**Good identification**					**07723131**
Actinomyces naeslundii		100.0	0.54	FRU 95		
07723132	**Good identification**					**07723132**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07723133	**Good identification**					**07723133**
Actinomyces naeslundii		100.0	0.54	FRU 95		
07723140	**Very good identification**					**07723140**
Actinomyces naeslundii		99.32	0.60	FRU 95		
Actinomyces israelii		0.68	0.14			
07723141	**Very good identification**					**07723141**
Actinomyces naeslundii		100.0	0.63	FRU 95		
07723142	**Very good identification**					**07723142**
Actinomyces naeslundii		98.99	0.60	FRU 95		
Actinomyces israelii		1.01	0.17			
07723143	**Very good identification**					**07723143**
Actinomyces naeslundii		100.0	0.63	FRU 95		
07723150	**Good identification**					**07723150**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07723151	**Good identification**					**07723151**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07723152	**Good identification**					**07723152**
Actinomyces naeslundii		100.0	0.38	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07723153	**Good identification**					**07723153**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07723160	**Very good identification**					**07723160**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07723161	**Very good identification**					**07723161**
Actinomyces naeslundii		100.0	0.63	FRU 95		
07723162	**Very good identification**					**07723162**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07723163	**Very good identification**					**07723163**
Actinomyces naeslundii		100.0	0.63	FRU 95		
07723170	**Good identification**					**07723170**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07723171	**Good identification**					**07723171**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07723172	**Good identification**					**07723172**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07723173	**Good identification**					**07723173**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07723300	**Good identification**					**07723300**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07723301	**Good identification**					**07723301**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07723302	**Good identification**					**07723302**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07723303	**Good identification**					**07723303**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07723321	**Good identification**					**07723321**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07723323	**Good identification**					**07723323**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07723500	**Low discrimination**					**07723500**
Actinomyces naeslundii		79.96	0.39	FRU 95		
Actinomyces israelii		20.04	0.19	ARA 1		
07723501	**Good identification**					**07723501**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	FRU 95		
07723503	**Good identification**					**07723503**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07723520	**Good identification**					**07723520**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07723521	**Good identification**					**07723521**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07723522	**Good identification**					**07723522**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07723523	**Good identification**					**07723523**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07724712	**Good identification**					**07724712**
Clostridium clostridioforme		100.0	0.40	FRU 99		
07724732	**Good identification**					**07724732**
Clostridium clostridioforme		97.81	0.40	FRU 99		
Parabacteroides distasonis		2.20	0.17			
07725101	**Good identification**					**07725101**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07725103	**Good identification**					**07725103**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07725121	**Good identification**					**07725121**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07725122	**Good identification**					**07725122**
					GRA	
Parabacteroides distasonis		95.92	0.51	FRU 99	1	
Actinomyces naeslundii		1.60	0.34		99	
Prevotella oralis		1.46	0.08		1	
07725123	**Low discrimination**					**07725123**
					GRA	
Actinomyces naeslundii		71.20	0.37	FRU 95	99	
Parabacteroides distasonis		28.80	0.17	XYL 1	1	
07725132	**Good identification**					**07725132**
Parabacteroides distasonis		97.89	0.47	FRU 99		
Clostridium clostridioforme		2.12	0.13			
07725322	**Very good identification**					**07725322**
Parabacteroides distasonis		99.17	0.60	FRU 99		
Clostridium clostridioforme		0.62	0.16			
Prevotella oralis		0.22	0.01			
07725332	**Acceptable identification**					**07725332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		94.98	0.55	FRU 99			
Clostridium clostridioforme		5.02	0.27				
07725502	**Low discrimination**						**07725502**
Bifidobacterium adolescentis		79.85	0.45	FRU 99			
Bifidobacterium longum subsp. longum		6.31	0.25				
Clostridium clostridioforme		5.07	0.22				
07725512	**Low discrimination**						**07725512**
Bifidobacterium adolescentis		73.98	0.47	FRU 99			
Clostridium clostridioforme		17.52	0.33				
Bifidobacterium longum subsp. longum		8.15	0.29				
07725520	**Low discrimination**						**07725520**
					GRA	SPO	
Bifidobacterium longum subsp. longum		78.09	0.35	SAL 1	99	1	
Bacteroides vulgatus		16.13	0.09	FRU 90	1	1	
Clostridium butyricum		3.36	-.01		99	99	
07725522	**Low discrimination**						**07725522**
					GRA		
Parabacteroides distasonis		71.40	0.48	FRU 99	1		
Bifidobacterium longum subsp. longum		23.20	0.35		99		
Clostridium clostridioforme		4.66	0.22		50		
07725530	**Good identification**						**07725530**
Bifidobacterium longum subsp. longum		100.0	0.39	SAL 1			
07725532	**Low discrimination**						**07725532**
					GRA		
Bifidobacterium longum subsp. longum		39.47	0.39	SAL 1	99		
Parabacteroides distasonis		38.41	0.43	FRU 90	1		
Clostridium clostridioforme		21.22	0.33		50		
07725612	**Good identification**						**07725612**
Clostridium clostridioforme		100.0	0.39	FRU 99			
07725632	**Good identification**						**07725632**
Clostridium clostridioforme		97.56	0.39	FRU 99			
Parabacteroides distasonis		2.44	0.17				
07725702	**Acceptable identification**						**07725702**
					GRA		
Clostridium clostridioforme		92.17	0.37	FRU 99	50		
Parabacteroides distasonis		5.84	0.22		1		
Bifidobacterium adolescentis		2.00	0.11		99		
07725712	**Good identification**						**07725712**
					GRA		
Clostridium clostridioforme		98.69	0.48	FRU 99	50		
Parabacteroides distasonis		0.74	0.17		1		
Bifidobacterium adolescentis		0.57	0.13		99		
07725722	**Low discrimination**						**07725722**
Parabacteroides distasonis		85.91	0.56	FRU 99			
Clostridium clostridioforme		13.71	0.37				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		0.21	-.05			
07725732	**Low discrimination**					**07725732**
Clostridium clostridioforme		57.45	0.48	FRU 99		
Parabacteroides distasonis		42.55	0.51			
07727100	**Low discrimination**					**07727100**
Actinomyces israelii		77.89	0.39	FRU 99		
Actinomyces naeslundii		22.11	0.39	CEL 93		
07727101	**Good identification**					**07727101**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07727102	**Low discrimination**					**07727102**
Actinomyces israelii		60.97	0.42	FRU 99		
Bifidobacterium adolescentis		27.49	0.30	CEL 93		
Actinomyces naeslundii		11.54	0.39			
07727103	**Good identification**					**07727103**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07727120	**Low discrimination**					**07727120**
Actinomyces naeslundii		81.65	0.39	FRU 95		
Actinomyces israelii		18.36	0.19	XYL 1		
07727121	**Good identification**					**07727121**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07727122	**Low discrimination**					**07727122**
					GRA	
Actinomyces naeslundii		60.62	0.39	FRU 95	99	
Actinomyces israelii		20.44	0.22	XYL 1	99	
Parabacteroides distasonis		18.94	0.17		1	
07727123	**Good identification**					**07727123**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07727142	**Low discrimination**					**07727142**
Actinomyces israelii		81.35	0.35	FRU 99		
Bifidobacterium adolescentis		11.66	0.14	CEL 93		
Actinomyces naeslundii		6.99	0.26			
07727500	**Low discrimination**					**07727500**
					SPO	
Actinomyces israelii		85.23	0.38	FRU 99	1	
Clostridium butyricum		10.00	0.01	CEL 93	99	
Bifidobacterium adolescentis		4.76	0.10		1	
07727502	**Low discrimination**					**07727502**
					SPO	
Bifidobacterium adolescentis		78.16	0.44	FRU 99	1	
Actinomyces israelii		21.20	0.41		1	
Clostridium butyricum		0.65	-.07		99	
07727512	**Good identification**					**07727512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		95.86	0.45	FRU 99			
Actinomyces israelii		3.18	0.26				
Bifidobacterium dentium		0.97	-.07				
07730000	**Low discrimination**						**07730000**
					GRA	CAT	
Propionibacterium granulosum		47.07	0.60	SAL 1	99	90	
Fusobacterium mortiferum		46.98	0.63		1	5	
Actinomyces naeslundii		5.33	0.59		99	25	
07730001	**Low discrimination**						**07730001**
					GRA	CAT	
Actinomyces naeslundii		89.38	0.62	GAL 96	99	25	
Propionibacterium granulosum		5.31	0.26		99	90	
Fusobacterium mortiferum		5.30	0.29		1	5	
07730002	**Low discrimination**						**07730002**
					GRA	CAT	
Propionibacterium granulosum		44.25	0.50	SAL 1	99	90	
Actinomyces naeslundii		20.05	0.59		99	25	
Fusobacterium mortiferum		19.63	0.47		1	5	
07730003	**Low discrimination**						**07730003**
Actinomyces naeslundii		83.81	0.62	GAL 96			
Actinomyces viscosus		14.39	0.24				
Propionibacterium granulosum		1.25	0.15				
07730010	**Low discrimination**						**07730010**
					GRA	CAT	
Propionibacterium granulosum		88.91	0.45	SAL 1	99	90	
Fusobacterium mortiferum		6.57	0.29	MLZ 12	1	5	
Actinomyces naeslundii		3.89	0.37		99	25	
07730011	**Low discrimination**						**07730011**
Actinomyces naeslundii		86.65	0.40	GAL 96			
Propionibacterium granulosum		13.35	0.11	MLZ 5			
07730013	**Good identification**						**07730013**
Actinomyces naeslundii		100.0	0.40	GAL 96			
07730020	**Low discrimination**						**07730020**
					CAT		
Propionibacterium granulosum		60.57	0.50	SAL 1	90		
Actinomyces naeslundii		27.45	0.59		25		
Clostridium perfringens		9.54	0.28		1		
07730021	**Very good identification**						**07730021**
Actinomyces naeslundii		98.54	0.62	GAL 96			
Propionibacterium granulosum		1.46	0.15				
07730022	**Low discrimination**						**07730022**
					CAT		
Actinomyces naeslundii		56.76	0.59	GAL 96	25		
Propionibacterium granulosum		31.31	0.39		90		
Clostridium perfringens		9.28	0.23		1		
07730023	**Very good identification**						**07730023**
Actinomyces naeslundii		100.0	0.62	GAL 96			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07730031	**Good identification**						**07730031**
Actinomyces naeslundii		100.0	0.40	GAL 96			
07730033	**Good identification**						**07730033**
Actinomyces naeslundii		100.0	0.40	GAL 96			
07730041	**Good identification**						**07730041**
					SPO		
Actinomyces naeslundii		96.01	0.49	GAL 96	1		
Clostridium botulinum biovar B		4.00	-.03		99		
07730042	**Low discrimination**						**07730042**
					SPO	CAT	
Actinomyces naeslundii		77.58	0.46	GAL 96	1	25	
Propionibacterium granulosum		9.80	0.15		1	90	
Clostridium botulinum biovar B		4.84	-.03		99	1	
07730043	**Good identification**						**07730043**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07730061	**Good identification**						**07730061**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07730062	**Low discrimination**						**07730062**
Actinomyces naeslundii		85.00	0.46	GAL 96			
Clostridium perfringens		15.00	0.11				
07730063	**Good identification**						**07730063**
Actinomyces naeslundii		100.0	0.49	GAL 96			
07730100	**Low discrimination**						**07730100**
					GRA	CAT	
Fusobacterium mortiferum		86.97	0.69	MLT 11	1	5	
Actinomyces naeslundii		12.04	0.66		99	25	
Propionibacterium granulosum		0.38	0.26		99	90	
07730101	**Very good identification**						**07730101**
					GRA		
Actinomyces naeslundii		95.36	0.69	GAL 96	99		
Fusobacterium mortiferum		4.64	0.35		1		
07730102	**Low discrimination**						**07730102**
					GRA		
Actinomyces naeslundii		41.97	0.66	GAL 96	99		
Fusobacterium mortiferum		33.69	0.53		1		
Actinomyces viscosus		20.26	0.37		99		
07730103	**Low discrimination**						**07730103**
					GRA		
Actinomyces naeslundii		89.88	0.69	GAL 96	99		
Actinomyces viscosus		9.64	0.29		99		
Fusobacterium mortiferum		0.49	0.19		1		
07730110	**Low discrimination**						**07730110**
					GRA	CAT	
Fusobacterium mortiferum		56.18	0.35	MLT 11	1	5	
Actinomyces naeslundii		40.53	0.44	MLZ 1	99	25	
Propionibacterium granulosum		3.29	0.11		99	90	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07730111	**Good identification**					**07730111**
Actinomyces naeslundii		100.0	0.48	GAL 96		
07730112	**Low discrimination**					**07730112**
Actinomyces naeslundii		47.40	0.44	GAL 96		
Clostridium clostridioforme		26.20	0.21	MLZ 5		
Bifidobacterium adolescentis		11.86	0.19			
07730113	**Good identification**					**07730113**
Actinomyces naeslundii		100.0	0.48	GAL 96		
07730120	**Low discrimination**					**07730120**
					GRA	
Actinomyces naeslundii		50.15	0.66	GAL 96	99	
Clostridium perfringens		44.89	0.43		99	
Fusobacterium mortiferum		3.66	0.35		1	
07730121	**Very good identification**					**07730121**
Actinomyces naeslundii		99.40	0.69	GAL 96		
Clostridium perfringens		0.60	0.09			
07730122	**Low discrimination**					**07730122**
					GRA	
Actinomyces naeslundii		59.06	0.66	GAL 96	99	
Clostridium perfringens		24.88	0.38		99	
Prevotella oralis		5.56	0.23		1	
07730123	**Very good identification**					**07730123**
Actinomyces naeslundii		99.72	0.69	GAL 96		
Clostridium perfringens		0.28	0.03			
07730130	**Low discrimination**					**07730130**
Actinomyces naeslundii		85.34	0.44	GAL 96		
Clostridium perfringens		14.66	0.09	MLZ 5		
07730131	**Good identification**					**07730131**
Actinomyces naeslundii		100.0	0.48	GAL 96		
07730132	**Low discrimination**					**07730132**
					GRA	
Actinomyces naeslundii		39.93	0.44	GAL 96	99	
Parabacteroides distasonis		34.77	0.30	MLZ 5	1	
Clostridium clostridioforme		22.07	0.21		50	
07730133	**Good identification**					**07730133**
Actinomyces naeslundii		100.0	0.48	GAL 96		
07730140	**Low discrimination**					**07730140**
					GRA	
Actinomyces naeslundii		68.25	0.53	GAL 96	99	
Fusobacterium mortiferum		28.22	0.35		1	
Clostridium perfringens		2.04	0.05		99	
07730141	**Very good identification**					**07730141**
Actinomyces naeslundii		100.0	0.57	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07730142	**Acceptable identification**					**07730142**
					GRA	
Actinomyces naeslundii		90.31	0.53	GAL 96	99	
Fusobacterium mortiferum		4.15	0.19		1	
Lactobacillus acidophilus		3.05	0.04		99	
07730143	**Very good identification**					**07730143**
Actinomyces naeslundii		100.0	0.57	GAL 96		
07730160	**Low discrimination**					**07730160**
Actinomyces naeslundii		50.86	0.53	GAL 96		
Clostridium perfringens		49.14	0.31			
07730161	**Very good identification**					**07730161**
Actinomyces naeslundii		100.0	0.57	GAL 96		
07730162	**Low discrimination**					**07730162**
Actinomyces naeslundii		67.18	0.53	GAL 96		
Clostridium perfringens		30.55	0.25			
Lactobacillus acidophilus		2.27	0.04			
07730163	**Very good identification**					**07730163**
Actinomyces naeslundii		100.0	0.57	GAL 96		
07730300	**Low discrimination**					**07730300**
					GRA	
Fusobacterium mortiferum		87.84	0.35	MLT 11	1	
Actinomyces naeslundii		12.16	0.32	RHA 1	99	
07730301	**Good identification**					**07730301**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730303	**Good identification**					**07730303**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730312	**Good identification**					**07730312**
Clostridium clostridioforme		100.0	0.36	XYL 91		
07730321	**Good identification**					**07730321**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730322	**Low discrimination**					**07730322**
Parabacteroides distasonis		65.90	0.43	GAL 99		
Clostridium clostridioforme		12.08	0.25	XYL 90		
Prevotella oralis		10.42	0.17			
07730323	**Good identification**					**07730323**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730332	**Low discrimination**					**07730332**
Clostridium clostridioforme		60.81	0.36	XYL 91		
Parabacteroides distasonis		39.20	0.39	ARA 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07730500	**Low discrimination**					**07730500**
					GRA	
Fusobacterium mortiferum		87.84	0.35	MLT 11	1	
Actinomyces naeslundii		12.16	0.32	ARA 1	99	
07730501	**Good identification**					**07730501**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730503	**Good identification**					**07730503**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730512	**Low discrimination**					**07730512**
Clostridium clostridioforme		83.80	0.42	XYL 91		
Bifidobacterium adolescentis		16.20	0.33	RHA 88		
07730521	**Good identification**					**07730521**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730523	**Good identification**					**07730523**
Actinomyces naeslundii		100.0	0.35	GAL 96		
07730532	**Acceptable identification**					**07730532**
					GRA	
Clostridium clostridioforme		92.69	0.42	XYL 91	50	
Parabacteroides distasonis		5.71	0.27	RHA 88	1	
Bifidobacterium longum subsp. longum		1.60	0.13		99	
07730602	**Good identification**					**07730602**
Clostridium clostridioforme		100.0	0.36	XYL 91		
07730612	**Good identification**					**07730612**
Clostridium clostridioforme		100.0	0.48	XYL 91		
07730622	**Good identification**					**07730622**
Clostridium clostridioforme		100.0	0.36	XYL 91		
07730632	**Good identification**					**07730632**
Clostridium clostridioforme		100.0	0.48	XYL 91		
07730702	**Good identification**					**07730702**
Clostridium clostridioforme		100.0	0.45	XYL 91		
07730712	**Very good identification**					**07730712**
Clostridium clostridioforme		100.0	0.57	XYL 91		
07730722	**Low discrimination**					**07730722**
Clostridium clostridioforme		82.42	0.45	XYL 91		
Parabacteroides distasonis		17.58	0.40			
07730732	**Very good identification**					**07730732**
Clostridium clostridioforme		97.54	0.57	XYL 91		
Parabacteroides distasonis		2.46	0.35			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07731000	**Low discrimination**						**07731000**
					GRA	CAT	
Actinomyces naeslundii		48.10	0.82		99	25	
Fusobacterium mortiferum		26.49	0.66		1	5	
Propionibacterium granulosum		17.69	0.60		99	90	
07731001	**Very good identification**						**07731001**
					GRA	CAT	
Actinomyces naeslundii		99.29	0.85		99	25	
Fusobacterium mortiferum		0.37	0.32		1	5	
Propionibacterium granulosum		0.25	0.26		99	90	
07731002	**Low discrimination**						**07731002**
					CAT		
Actinomyces naeslundii		75.37	0.82		25		
Propionibacterium granulosum		6.93	0.50		90		
Lactobacillus jensenii		5.32	0.41		1		
07731003	**Very good identification**						**07731003**
Actinomyces naeslundii		99.38	0.85				
Actinomyces viscosus		0.44	0.21				
Propionibacterium granulosum		0.06	0.15				
07731010	**Low discrimination**						**07731010**
					CAT		
Actinomyces naeslundii		36.38	0.60	MLZ 5	25		
Propionibacterium granulosum		34.67	0.45		90		
Lactobacillus jensenii		24.40	0.36		1		
07731011	**Very good identification**						**07731011**
					CAT		
Actinomyces naeslundii		98.92	0.63	MLZ 5	25		
Propionibacterium granulosum		0.64	0.11		90		
Lactobacillus jensenii		0.45	0.02		1		
07731012	**Low discrimination**						**07731012**
					CAT		
Actinomyces naeslundii		54.98	0.60	MLZ 5	25		
Lactobacillus jensenii		24.58	0.33		1		
Propionibacterium granulosum		13.10	0.35		90		
07731013	**Very good identification**						**07731013**
Actinomyces naeslundii		100.0	0.63	MLZ 5			
07731020	**Low discrimination**						**07731020**
					CAT		
Clostridium perfringens		55.92	0.63	LAC 88	1		
Actinomyces naeslundii		39.02	0.82		25		
Propionibacterium granulosum		3.59	0.50		90		
07731021	**Very good identification**						**07731021**
					CAT		
Actinomyces naeslundii		98.98	0.85		25		
Clostridium perfringens		0.96	0.28		1		
Propionibacterium granulosum		0.06	0.15		90		
07731022	**Low discrimination**						**07731022**
					GRA		
Actinomyces naeslundii		54.06	0.82		99		
Clostridium perfringens		36.46	0.57		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		3.82	0.37		1	
07731023	**Very good identification**					**07731023**
					GRA	
Actinomyces naeslundii		99.47	0.85		99	
Clostridium perfringens		0.45	0.23		99	
Prevotella oralis		0.05	0.03		1	
07731030	**Low discrimination**					**07731030**
					CAT	
Actinomyces naeslundii		59.49	0.60	MLZ 5	25	
Clostridium perfringens		16.36	0.28		1	
Propionibacterium granulosum		14.17	0.35		90	
07731031	**Very good identification**					**07731031**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
07731032	**Low discrimination**					**07731032**
Actinomyces naeslundii		67.30	0.60	MLZ 5		
Clostridium perfringens		8.71	0.23			
Lactobacillus jensenii		7.52	0.23			
07731033	**Very good identification**					**07731033**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
07731040	**Low discrimination**					**07731040**
					GRA	
Actinomyces naeslundii		89.47	0.69		99	
Clostridium perfringens		4.28	0.24		99	
Fusobacterium mortiferum		2.82	0.32		1	
07731041	**Very good identification**					**07731041**
Actinomyces naeslundii		100.0	0.72			
07731042	**Good identification**					**07731042**
Actinomyces naeslundii		92.35	0.69			
Lactobacillus acidophilus		3.19	0.20			
Clostridium perfringens		2.08	0.19			
07731043	**Very good identification**					**07731043**
Actinomyces naeslundii		100.0	0.72			
07731050	**Acceptable identification**					**07731050**
					CAT	
Actinomyces naeslundii		91.50	0.47	MLZ 5	25	
Propionibacterium granulosum		4.99	0.11		90	
Lactobacillus jensenii		3.51	0.02		1	
07731051	**Good identification**					**07731051**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07731052	**Good identification**					**07731052**
Actinomyces naeslundii		96.44	0.47	MLZ 5		
Actinomyces israelii		3.56	0.14			
07731053	**Good identification**					**07731053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07731060	**Low discrimination**					**07731060**
					CAT	
Clostridium perfringens		60.61	0.50	LAC 88	1	
Actinomyces naeslundii		39.18	0.69		25	
Propionibacterium granulosum		0.21	0.15		90	
07731061	**Very good identification**					**07731061**
Actinomyces naeslundii		98.97	0.72			
Clostridium perfringens		1.03	0.16			
07731062	**Low discrimination**					**07731062**
Actinomyces naeslundii		56.61	0.69			
Clostridium perfringens		41.21	0.45			
Lactobacillus acidophilus		1.95	0.20			
07731063	**Very good identification**					**07731063**
Actinomyces naeslundii		99.51	0.72			
Clostridium perfringens		0.49	0.11			
07731070	**Low discrimination**					**07731070**
Actinomyces naeslundii		77.11	0.47	MLZ 5		
Clostridium perfringens		22.89	0.16			
07731071	**Good identification**					**07731071**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07731072	**Low discrimination**					**07731072**
Actinomyces naeslundii		87.74	0.47	MLZ 5		
Clostridium perfringens		12.26	0.11			
07731073	**Good identification**					**07731073**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
07731100	**Low discrimination**					**07731100**
					GRA	
Actinomyces naeslundii		60.35	0.90		99	
Fusobacterium mortiferum		27.25	0.73		1	
Clostridium perfringens		6.89	0.52		99	
07731101	**Excellent identification**					**07731101**
					GRA	
Actinomyces naeslundii		99.54	0.93		99	
Fusobacterium mortiferum		0.30	0.38		1	
Clostridium perfringens		0.08	0.17		99	
07731102	**Low discrimination**					**07731102**
Actinomyces naeslundii		70.37	0.90			
Lactobacillus acidophilus		14.00	0.54			
Clostridium perfringens		3.78	0.46			
07731103	**Excellent identification**					**07731103**
Actinomyces naeslundii		99.47	0.93			
Actinomyces viscosus		0.27	0.25			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		0.13	0.19			
07731110	**Low discrimination**					**07731110**
					GRA	
Actinomyces naeslundii		86.37	0.68	MLZ 5	99	
Fusobacterium mortiferum		7.48	0.38		1	
Clostridium perfringens		1.89	0.17		99	
07731111	**Very good identification**					**07731111**
Actinomyces naeslundii		100.0	0.71	MLZ 5		
07731112	**Low discrimination**					**07731112**
					GRA	
Bifidobacterium adolescentis		41.31	0.53	XYL 87	99	
Actinomyces naeslundii		40.03	0.68	ARA 87	99	
Leptotrichia buccalis		9.94	0.24		1	
07731113	**Very good identification**					**07731113**
					GRA	
Actinomyces naeslundii		99.15	0.71	MLZ 5	99	
Bifidobacterium adolescentis		0.69	0.19		99	
Leptotrichia buccalis		0.17	-.11		1	
07731120	**Low discrimination**					**07731120**
					SPO	
Clostridium perfringens		77.18	0.77		84	
Actinomyces naeslundii		20.90	0.90		1	
Clostridium ramosum		0.44	0.36		92	
07731121	**Very good identification**					**07731121**
					SPO	
Actinomyces naeslundii		97.52	0.93		1	
Clostridium perfringens		2.43	0.43		84	
Clostridium ramosum		0.01	0.02		92	
07731122	**Low discrimination**					**07731122**
					GRA	
Clostridium perfringens		45.83	0.72		99	
Actinomyces naeslundii		26.38	0.90		99	
Prevotella oralis		10.25	0.57		1	
07731123	**Very good identification**					**07731123**
					GRA	
Actinomyces naeslundii		98.16	0.93		99	
Clostridium perfringens		1.15	0.38		99	
Prevotella oralis		0.26	0.23		1	
07731130	**Low discrimination**					**07731130**
					GRA	
Actinomyces naeslundii		56.53	0.68	MLZ 5	99	
Clostridium perfringens		40.06	0.43		99	
Parabacteroides distasonis		2.05	0.30		1	
07731131	**Very good identification**					**07731131**
Actinomyces naeslundii		99.53	0.71	MLZ 5		
Clostridium perfringens		0.48	0.09			
07731132	**Low discrimination**					**07731132**
					GRA	
Parabacteroides distasonis		68.99	0.65	XYL 90	1	
Actinomyces naeslundii		19.20	0.68		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		6.40	0.38		99	
07731133	**Very good identification**					**07731133**
					GRA	
Actinomyces naeslundii		97.42	0.71	MLZ 5	99	
Parabacteroides distasonis		2.36	0.30		1	
Clostridium perfringens		0.22	0.03		99	
07731140	**Low discrimination**					**07731140**
					GRA	
Actinomyces naeslundii		85.68	0.77		99	
Clostridium perfringens		10.56	0.39		99	
Fusobacterium mortiferum		2.21	0.38		1	
07731141	**Very good identification**					**07731141**
Actinomyces naeslundii		99.92	0.80			
Clostridium perfringens		0.08	0.05			
07731142	**Low discrimination**					**07731142**
Actinomyces naeslundii		79.44	0.77			
Lactobacillus acidophilus		11.07	0.38			
Clostridium perfringens		4.61	0.34			
07731143	**Very good identification**					**07731143**
Actinomyces naeslundii		99.91	0.80			
Lactobacillus acidophilus		0.09	0.04			
07731150	**Acceptable identification**					**07731150**
Actinomyces naeslundii		94.50	0.55	MLZ 5		
Actinomyces israelii		3.27	0.21			
Clostridium perfringens		2.24	0.05			
07731151	**Very good identification**					**07731151**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07731152	**Low discrimination**					**07731152**
Actinomyces naeslundii		51.86	0.55	MLZ 5		
Bifidobacterium adolescentis		37.49	0.37			
Bifidobacterium breve		5.84	0.24			
07731153	**Very good identification**					**07731153**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07731160	**Low discrimination**					**07731160**
Clostridium perfringens		79.82	0.65			
Actinomyces naeslundii		20.03	0.77			
Lactobacillus acidophilus		0.03	0.04			
07731161	**Very good identification**					**07731161**
Actinomyces naeslundii		97.39	0.80			
Clostridium perfringens		2.61	0.31			
07731162	**Low discrimination**					**07731162**
Clostridium perfringens		61.00	0.59			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		32.52	0.77			
Lactobacillus acidophilus		4.53	0.38			
07731163	**Very good identification**					**07731163**
Actinomyces naeslundii		98.66	0.80			
Clostridium perfringens		1.25	0.25			
Lactobacillus acidophilus		0.09	0.04			
07731170	**Low discrimination**					**07731170**
Actinomyces naeslundii		56.03	0.55	MLZ 5		
Clostridium perfringens		42.86	0.31			
Bifidobacterium breve		1.11	0.12			
07731171	**Very good identification**					**07731171**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07731172	**Low discrimination**					**07731172**
					GRA	
Actinomyces naeslundii		55.79	0.55	MLZ 5	99	
Clostridium perfringens		20.08	0.25		99	
Parabacteroides distasonis		11.47	0.30		1	
07731173	**Very good identification**					**07731173**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
07731201	**Good identification**					**07731201**
Actinomyces naeslundii		100.0	0.51	RHA 1		
07731202	**Low discrimination**					**07731202**
Actinomyces naeslundii		40.41	0.48	RHA 1		
Clostridium clostridioforme		19.91	0.24			
Lactobacillus acidophilus		14.83	0.16			
07731203	**Good identification**					**07731203**
Actinomyces naeslundii		100.0	0.51	RHA 1		
07731212	**Good identification**					**07731212**
Clostridium clostridioforme		96.10	0.35	XYL 91		
Actinomyces naeslundii		2.25	0.26	ARA 94		
Propionibacterium granulosum		1.64	0.09			
07731221	**Good identification**					**07731221**
Actinomyces naeslundii		100.0	0.51	RHA 1		
07731222	**Low discrimination**					**07731222**
					GRA	
Parabacteroides distasonis		33.64	0.43	XYL 90	1	
Prevotella oralis		33.60	0.31	LAC 99	1	
Actinomyces naeslundii		11.22	0.48		99	
07731223	**Good identification**					**07731223**
Actinomyces naeslundii		100.0	0.51	RHA 1		
07731232	**Low discrimination**					**07731232**
					GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		57.38	0.35	XYL 91	50	
Parabacteroides distasonis		41.28	0.39	ARA 94	1	
Actinomyces naeslundii		1.35	0.26		99	
07731240	**Good identification**					**07731240**
Actinomyces naeslundii		100.0	0.35	RHA 1		
07731241	**Good identification**					**07731241**
Actinomyces naeslundii		100.0	0.38	RHA 1		
07731242	**Low discrimination**					**07731242**
Actinomyces naeslundii		79.55	0.35	RHA 1		
Lactobacillus acidophilus		20.45	0.01			
07731243	**Good identification**					**07731243**
Actinomyces naeslundii		100.0	0.38	RHA 1		
07731261	**Good identification**					**07731261**
Actinomyces naeslundii		100.0	0.38	RHA 1		
07731262	**Low discrimination**					**07731262**
Actinomyces naeslundii		50.38	0.35	RHA 1		
Clostridium perfringens		36.67	0.11			
Lactobacillus acidophilus		12.95	0.01			
07731263	**Good identification**					**07731263**
Actinomyces naeslundii		100.0	0.38	RHA 1		
07731300	**Low discrimination**					**07731300**
					GRA	
Actinomyces naeslundii		56.42	0.56	RHA 1	99	
Fusobacterium mortiferum		25.47	0.38		1	
Clostridium perfringens		6.44	0.17		99	
07731301	**Very good identification**					**07731301**
Actinomyces naeslundii		100.0	0.59	RHA 1		
07731303	**Very good identification**					**07731303**
Actinomyces naeslundii		99.01	0.59	RHA 1		
Lactobacillus acidophilus		0.99	0.00			
07731311	**Good identification**					**07731311**
Actinomyces naeslundii		100.0	0.37	RHA 1		
07731312	**Low discrimination**					**07731312**
					GRA	
Clostridium clostridioforme		78.99	0.44	XYL 91	50	
Parabacteroides distasonis		16.97	0.39	ARA 94	1	
Bifidobacterium adolescentis		1.63	0.19		99	
07731313	**Low discrimination**					**07731313**
Actinomyces naeslundii		74.75	0.37	RHA 1		
Clostridium clostridioforme		25.25	0.10	MLZ 5		
07731320	**Low discrimination**					**07731320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium perfringens		38.50	0.43	RHA 1	99	84	
Clostridium ramosum		28.68	0.38		50	92	
Parabacteroides distasonis		10.98	0.43		1	1	
07731321	**Very good identification**						**07731321**
					SPO		
Actinomyces naeslundii		95.84	0.59	RHA 1	1		
Clostridium perfringens		2.38	0.09		84		
Clostridium ramosum		1.78	0.04		92		
07731322	**Low discrimination**						**07731322**
					SPO		
Parabacteroides distasonis		76.35	0.77	XYL 90	1		
Prevotella oralis		12.07	0.51		1		
Clostridium ramosum		8.05	0.48		92		
07731323	**Low discrimination**						**07731323**
					GRA		
Actinomyces naeslundii		52.38	0.59	RHA 1	99		
Parabacteroides distasonis		36.79	0.43		1		
Prevotella oralis		5.82	0.17		1		
07731330	**Low discrimination**						**07731330**
					GRA		
Parabacteroides distasonis		79.71	0.39	XYL 90	1		
Actinomyces naeslundii		7.40	0.34	BGL 99	99		
Clostridium perfringens		5.24	0.09		99		
07731331	**Good identification**						**07731331**
Actinomyces naeslundii		100.0	0.37	RHA 1			
07731332	**Good identification**						**07731332**
Parabacteroides distasonis		94.92	0.73	XYL 90			
Clostridium clostridioforme		4.46	0.44				
Prevotella oralis		0.28	0.17				
07731333	**Low discrimination**						**07731333**
					GRA		
Parabacteroides distasonis		84.31	0.39	XYL 90	1		
Actinomyces naeslundii		11.73	0.37	URE 1	99		
Clostridium clostridioforme		3.96	0.10		50		
07731340	**Low discrimination**						**07731340**
Actinomyces naeslundii		82.38	0.43	RHA 1			
Clostridium perfringens		10.15	0.05				
Actinomyces israelii		7.47	0.16				
07731341	**Good identification**						**07731341**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07731343	**Good identification**						**07731343**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07731361	**Good identification**						**07731361**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07731362	**Low discrimination**						**07731362**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		52.10	0.43	XYL 90	1		
Clostridium perfringens		16.37	0.25	SOR 1	99		
Lactobacillus acidophilus		9.06	0.19		99		
07731363	**Good identification**						**07731363**
Actinomyces naeslundii		100.0	0.46	RHA 1			
07731372	**Good identification**						**07731372**
Parabacteroides distasonis		95.51	0.39	XYL 90			
Clostridium clostridioforme		4.49	0.10	SOR 1			
07731400	**Low discrimination**						**07731400**
					GRA	CAT	
Actinomyces naeslundii		38.65	0.48	ARA 1	99	25	
Fusobacterium mortiferum		21.28	0.32		1	5	
Propionibacterium granulosum		14.22	0.26		99	90	
07731401	**Good identification**						**07731401**
Actinomyces naeslundii		100.0	0.51	ARA 1			
07731403	**Good identification**						**07731403**
Actinomyces naeslundii		100.0	0.51	ARA 1			
07731412	**Low discrimination**						**07731412**
Clostridium clostridioforme		80.54	0.41	XYL 91			
Bifidobacterium adolescentis		17.38	0.33	RHA 88			
Actinomyces israelii		1.20	0.19				
07731421	**Good identification**						**07731421**
Actinomyces naeslundii		100.0	0.51	ARA 1			
07731423	**Good identification**						**07731423**
Actinomyces naeslundii		100.0	0.51	ARA 1			
07731432	**Acceptable identification**						**07731432**
					GRA		
Clostridium clostridioforme		90.99	0.41	XYL 91	50		
Parabacteroides distasonis		6.26	0.27	RHA 88	1		
Bifidobacterium longum subsp. longum		1.75	0.13		99		
07731440	**Low discrimination**						**07731440**
Actinomyces naeslundii		58.78	0.35	ARA 1			
Actinomyces israelii		41.23	0.23				
07731441	**Good identification**						**07731441**
Actinomyces naeslundii		100.0	0.38	ARA 1			
07731443	**Good identification**						**07731443**
Actinomyces naeslundii		100.0	0.38	ARA 1			
07731461	**Good identification**						**07731461**
Actinomyces naeslundii		100.0	0.38	ARA 1			
07731462	**Low discrimination**						**07731462**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		57.87	0.35	ARA 1		
Clostridium perfringens		42.13	0.11			
07731463	**Good identification**					**07731463**
Actinomyces naeslundii		100.0	0.38	ARA 1		
07731500	**Low discrimination**					**07731500**
					SPO	
Actinomyces naeslundii		25.81	0.56	ARA 1	1	
Clostridium butyricum		24.58	0.25		99	
Bifidobacterium longum subsp. longum		12.18	0.33		1	
07731501	**Very good identification**					**07731501**
Actinomyces naeslundii		100.0	0.59	ARA 1		
07731502	**Low discrimination**					**07731502**
Bifidobacterium adolescentis		87.06	0.66	XYL 87		
Clostridium clostridioforme		3.66	0.39			
Actinomyces naeslundii		2.96	0.56			
07731503	**Low discrimination**					**07731503**
Actinomyces naeslundii		82.88	0.59	ARA 1		
Bifidobacterium adolescentis		16.43	0.31			
Clostridium clostridioforme		0.69	0.05			
07731510	**Low discrimination**					**07731510**
Bifidobacterium longum subsp. longum		57.66	0.37	SAL 1		
Bifidobacterium adolescentis		26.08	0.33			
Actinomyces israelii		4.80	0.27			
07731511	**Good identification**					**07731511**
Actinomyces naeslundii		100.0	0.37	ARA 1		
07731512	**Low discrimination**					**07731512**
Bifidobacterium adolescentis		84.22	0.67	XYL 87		
Clostridium clostridioforme		13.20	0.50			
Bifidobacterium longum subsp. longum		1.88	0.37			
07731520	**Low discrimination**					**07731520**
Clostridium perfringens		50.76	0.43	ARA 1		
Bifidobacterium longum subsp. longum		25.96	0.43			
Actinomyces naeslundii		13.75	0.56			
07731521	**Very good identification**					**07731521**
Actinomyces naeslundii		96.38	0.59	ARA 1		
Clostridium perfringens		2.40	0.09			
Bifidobacterium longum subsp. longum		1.23	0.09			
07731522	**Low discrimination**					**07731522**
					GRA	
Parabacteroides distasonis		75.42	0.66	XYL 90	1	
Bifidobacterium longum subsp. longum		6.68	0.43		99	
Clostridium perfringens		6.15	0.38		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07731523	**Low discrimination**					**07731523**
					GRA	
Actinomyces naeslundii		85.02	0.59	ARA 1	99	
Parabacteroides distasonis		12.20	0.31		1	
Bifidobacterium longum subsp. longum		1.08	0.09		99	
07731530	**Acceptable identification**					**07731530**
					GRA	
Bifidobacterium longum subsp. longum		92.44	0.47	SAL 1	99	
Parabacteroides distasonis		3.33	0.27		1	
Clostridium clostridioforme		1.64	0.16		50	
07731531	**Low discrimination**					**07731531**
Actinomyces naeslundii		70.86	0.37	ARA 1		
Bifidobacterium longum subsp. longum		29.14	0.13	MLZ 5		
07731532	**Low discrimination**					**07731532**
					GRA	
Parabacteroides distasonis		55.26	0.61	XYL 90	1	
Clostridium clostridioforme		27.17	0.50		50	
Bifidobacterium longum subsp. longum		15.48	0.47		99	
07731540	**Low discrimination**					**07731540**
					SPO	
Actinomyces naeslundii		35.89	0.43	ARA 1	1	
Actinomyces israelii		35.37	0.34		1	
Clostridium butyricum		16.84	0.07		99	
07731541	**Good identification**					**07731541**
Actinomyces naeslundii		100.0	0.46	ARA 1		
07731542	**Low discrimination**					**07731542**
Bifidobacterium adolescentis		87.78	0.50	XYL 87		
Actinomyces israelii		6.29	0.37	SOR 11		
Actinomyces naeslundii		4.26	0.43			
07731543	**Low discrimination**					**07731543**
Actinomyces naeslundii		87.81	0.46	ARA 1		
Bifidobacterium adolescentis		12.20	0.16			
07731552	**Good identification**					**07731552**
Bifidobacterium adolescentis		97.89	0.51	XYL 87		
Clostridium clostridioforme		1.25	0.16	SOR 11		
Actinomyces israelii		0.86	0.23			
07731561	**Good identification**					**07731561**
Actinomyces naeslundii		100.0	0.46	ARA 1		
07731563	**Good identification**					**07731563**
Actinomyces naeslundii		100.0	0.46	ARA 1		
07731602	**Good identification**					**07731602**
Clostridium clostridioforme		99.54	0.45	XYL 91		
Actinomyces israelii		0.46	0.14			
07731612	**Good identification**					**07731612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.56	XYL 91			
07731622	**Low discrimination**						**07731622**
Clostridium clostridioforme		80.77	0.45	XYL 91			
Parabacteroides distasonis		19.23	0.40				
07731632	**Good identification**						**07731632**
Clostridium clostridioforme		97.26	0.56	XYL 91			
Parabacteroides distasonis		2.74	0.35				
07731702	**Low discrimination**						**07731702**
					GRA		
Clostridium clostridioforme		89.90	0.54	XYL 91	50		
Parabacteroides distasonis		6.39	0.40		1		
Bifidobacterium adolescentis		2.95	0.31		99		
07731712	**Very good identification**						**07731712**
					GRA		
Clostridium clostridioforme		98.31	0.65	XYL 91	50		
Bifidobacterium adolescentis		0.86	0.33		99		
Parabacteroides distasonis		0.83	0.35		1		
07731722	**Low discrimination**						**07731722**
Parabacteroides distasonis		87.07	0.74	XYL 90			
Clostridium clostridioforme		12.37	0.54				
Prevotella oralis		0.23	0.17				
07731723	**Low discrimination**						**07731723**
					GRA		
Parabacteroides distasonis		85.80	0.40	XYL 90	1		
Clostridium clostridioforme		12.19	0.19	URE 1	50		
Actinomyces naeslundii		2.01	0.25		99		
07731732	**Low discrimination**						**07731732**
					GRA		
Clostridium clostridioforme		54.54	0.65	XYL 91	50		
Parabacteroides distasonis		45.39	0.69		1		
Bifidobacterium longum subsp. longum		0.04	0.13		99		
07731762	**Low discrimination**						**07731762**
Parabacteroides distasonis		87.56	0.40	XYL 90			
Clostridium clostridioforme		12.44	0.19	SOR 1			
07732000	**Low discrimination**						**07732000**
					GRA	CAT	
Propionibacterium granulosum		76.64	0.58	SAL 1	99	90	
Actinomyces naeslundii		22.34	0.64		99	25	
Fusobacterium mortiferum		1.02	0.29		1	5	
07732001	**Very good identification**						**07732001**
Actinomyces naeslundii		97.74	0.67	GAL 96			
Propionibacterium granulosum		2.26	0.24				
07732002	**Low discrimination**						**07732002**
					GRA	CAT	
Actinomyces naeslundii		53.45	0.64	GAL 96	99	25	
Propionibacterium granulosum		45.84	0.48		99	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.27	0.13		1	5	
07732003	**Very good identification**						**07732003**
Actinomyces naeslundii		99.43	0.67	GAL 96			
Propionibacterium granulosum		0.57	0.13				
07732010	**Low discrimination**						**07732010**
Propionibacterium granulosum		89.89	0.43	SAL 1			
Actinomyces naeslundii		10.11	0.42	MLZ 12			
07732011	**Acceptable identification**						**07732011**
Actinomyces naeslundii		94.35	0.45	GAL 96			
Propionibacterium granulosum		5.65	0.09	MLZ 5			
07732013	**Good identification**						**07732013**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07732020	**Low discrimination**						**07732020**
Actinomyces naeslundii		53.83	0.64	GAL 96			
Propionibacterium granulosum		46.17	0.48				
07732021	**Very good identification**						**07732021**
Actinomyces naeslundii		99.43	0.67	GAL 96			
Propionibacterium granulosum		0.57	0.13				
07732022	**Low discrimination**						**07732022**
Actinomyces naeslundii		82.34	0.64	GAL 96			
Propionibacterium granulosum		17.66	0.37				
07732023	**Very good identification**						**07732023**
Actinomyces naeslundii		100.0	0.67	GAL 96			
07732031	**Good identification**						**07732031**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07732032	**Low discrimination**						**07732032**
Actinomyces naeslundii		64.29	0.42	GAL 96			
Propionibacterium granulosum		35.71	0.22	MLZ 5			
07732033	**Good identification**						**07732033**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07732040	**Low discrimination**						**07732040**
					SPO	CAT	
Actinomyces naeslundii		81.40	0.51	GAL 96	1	25	
Propionibacterium granulosum		15.98	0.24		1	90	
Clostridium botulinum biovar B		2.62	-.03		99	1	
07732041	**Good identification**						**07732041**
Actinomyces naeslundii		100.0	0.54	GAL 96			
07732042	**Good identification**						**07732042**
Actinomyces naeslundii		95.32	0.51	GAL 96			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		4.68	0.13			
07732043	**Good identification**					**07732043**
Actinomyces naeslundii		100.0	0.54	GAL 96		
07732060	**Good identification**					**07732060**
Actinomyces naeslundii		95.32	0.51	GAL 96		
Propionibacterium granulosum		4.68	0.13			
07732061	**Good identification**					**07732061**
Actinomyces naeslundii		100.0	0.54	GAL 96		
07732062	**Good identification**					**07732062**
Actinomyces naeslundii		100.0	0.51	GAL 96		
07732063	**Good identification**					**07732063**
Actinomyces naeslundii		100.0	0.54	GAL 96		
07732100	**Low discrimination**					**07732100**
					GRA	
Actinomyces naeslundii		88.57	0.71	GAL 96	99	
Propionibacterium propionicum		6.68	0.42		99	
Fusobacterium mortiferum		3.33	0.35		1	
07732101	**Very good identification**					**07732101**
Actinomyces naeslundii		99.45	0.74	GAL 96		
Propionibacterium propionicum		0.56	0.26			
07732102	**Low discrimination**					**07732102**
Actinomyces naeslundii		88.25	0.71	GAL 96		
Propionibacterium propionicum		9.98	0.45			
Actinomyces israelii		0.52	0.24			
07732103	**Very good identification**					**07732103**
Actinomyces naeslundii		99.17	0.74	GAL 96		
Propionibacterium propionicum		0.83	0.29			
07732110	**Good identification**					**07732110**
Actinomyces naeslundii		96.94	0.49	GAL 96		
Propionibacterium granulosum		3.06	0.09	MLZ 5		
07732111	**Good identification**					**07732111**
Actinomyces naeslundii		100.0	0.52	GAL 96		
07732112	**Acceptable identification**					**07732112**
Actinomyces naeslundii		90.46	0.49	GAL 96		
Bifidobacterium adolescentis		9.54	0.17	MLZ 5		
07732113	**Good identification**					**07732113**
Actinomyces naeslundii		100.0	0.52	GAL 96		
07732120	**Very good identification**					**07732120**
					SPO	
Actinomyces naeslundii		97.80	0.71	GAL 96	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		1.45	0.15		92	
Clostridium perfringens		0.46	0.09		84	
07732121	**Very good identification**					**07732121**
Actinomyces naeslundii		100.0	0.74	GAL 96		
07732122	**Good identification**					**07732122**
					SPO	
Actinomyces naeslundii		94.20	0.71	GAL 96	1	
Clostridium ramosum		5.59	0.25		92	
Clostridium perfringens		0.21	0.03		84	
07732123	**Very good identification**					**07732123**
Actinomyces naeslundii		100.0	0.74	GAL 96		
07732130	**Good identification**					**07732130**
Actinomyces naeslundii		100.0	0.49	GAL 96		
07732131	**Good identification**					**07732131**
Actinomyces naeslundii		100.0	0.52	GAL 96		
07732132	**Good identification**					**07732132**
Actinomyces naeslundii		100.0	0.49	GAL 96		
07732133	**Good identification**					**07732133**
Actinomyces naeslundii		100.0	0.52	GAL 96		
07732140	**Low discrimination**					**07732140**
Actinomyces naeslundii		69.66	0.58	GAL 96		
Propionibacterium propionicum		29.75	0.42			
Actinomyces israelii		0.60	0.14			
07732141	**Very good identification**					**07732141**
Actinomyces naeslundii		96.93	0.61	GAL 96		
Propionibacterium propionicum		3.07	0.26			
07732142	**Low discrimination**					**07732142**
Actinomyces naeslundii		60.48	0.58	GAL 96		
Propionibacterium propionicum		38.74	0.45			
Actinomyces israelii		0.78	0.17			
07732143	**Very good identification**					**07732143**
Actinomyces naeslundii		95.47	0.61	GAL 96		
Propionibacterium propionicum		4.53	0.29			
07732150	**Good identification**					**07732150**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07732151	**Good identification**					**07732151**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732152	**Good identification**					**07732152**
Actinomyces naeslundii		100.0	0.37	GAL 96		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07732153	**Good identification**					**07732153**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732160	**Very good identification**					**07732160**
Actinomyces naeslundii		100.0	0.58	GAL 96		
07732161	**Very good identification**					**07732161**
Actinomyces naeslundii		100.0	0.61	GAL 96		
07732162	**Very good identification**					**07732162**
Actinomyces naeslundii		100.0	0.58	GAL 96		
07732163	**Very good identification**					**07732163**
Actinomyces naeslundii		100.0	0.61	GAL 96		
07732170	**Good identification**					**07732170**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07732171	**Good identification**					**07732171**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732172	**Good identification**					**07732172**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07732173	**Good identification**					**07732173**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732300	**Good identification**					**07732300**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07732301	**Good identification**					**07732301**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732302	**Good identification**					**07732302**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07732303	**Good identification**					**07732303**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732321	**Good identification**					**07732321**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732323	**Good identification**					**07732323**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732500	**Low discrimination**					**07732500**
Actinomyces naeslundii		75.96	0.37	GAL 96		
Actinomyces israelii		24.04	0.19	ARA 1		
07732501	**Good identification**					**07732501**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732503	**Good identification**					**07732503**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732520	**Good identification**					**07732520**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07732521	**Good identification**					**07732521**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07732522	**Good identification**					**07732522**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07732523	**Good identification**					**07732523**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07733000	**Low discrimination**					**07733000**
					CAT	
Actinomyces naeslundii		86.35	0.87		25	
Propionibacterium granulosum		12.34	0.58		90	
Actinomyces israelii		0.99	0.45		1	
07733001	**Very good identification**					**07733001**
Actinomyces naeslundii		99.90	0.90			
Propionibacterium granulosum		0.10	0.24			
07733002	**Good identification**					**07733002**
					CAT	
Actinomyces naeslundii		94.74	0.87		25	
Propionibacterium granulosum		3.39	0.48		90	
Actinomyces israelii		1.63	0.48		1	
07733003	**Very good identification**					**07733003**
					CAT	
Actinomyces naeslundii		99.96	0.90		25	
Propionibacterium granulosum		0.02	0.13		90	
Actinomyces israelii		0.01	0.14		1	
07733010	**Low discrimination**					**07733010**
					CAT	
Actinomyces naeslundii		71.10	0.65	MLZ 5	25	
Propionibacterium granulosum		26.33	0.43		90	
Actinomyces israelii		2.32	0.31		1	
07733011	**Very good identification**					**07733011**
Actinomyces naeslundii		99.75	0.68	MLZ 5		
Propionibacterium granulosum		0.25	0.09			
07733012	**Low discrimination**					**07733012**
					CAT	
Actinomyces naeslundii		86.66	0.65	MLZ 5	25	
Propionibacterium granulosum		8.02	0.33		90	
Actinomyces israelii		4.23	0.34		1	
07733013	**Very good identification**					**07733013**
Actinomyces naeslundii		100.0	0.68	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07733020	**Very good identification**						**07733020**
					CAT		
Actinomyces naeslundii		95.63	0.87		25		
Propionibacterium granulosum		3.42	0.48		90		
Clostridium perfringens		0.71	0.28		1		
07733021	**Very good identification**						**07733021**
Actinomyces naeslundii		99.98	0.90				
Propionibacterium granulosum		0.02	0.13				
07733022	**Very good identification**						**07733022**
					SPO	CAT	
Actinomyces naeslundii		97.82	0.87		1	25	
Propionibacterium granulosum		0.87	0.37		1	90	
Clostridium ramosum		0.69	0.25		92	1	
07733023	**Very good identification**						**07733023**
Actinomyces naeslundii		100.0	0.90				
07733030	**Good identification**						**07733030**
Actinomyces naeslundii		91.53	0.65	MLZ 5			
Propionibacterium granulosum		8.47	0.33				
07733031	**Very good identification**						**07733031**
Actinomyces naeslundii		100.0	0.68	MLZ 5			
07733032	**Very good identification**						**07733032**
					CAT		
Actinomyces naeslundii		97.44	0.65	MLZ 5	25		
Propionibacterium granulosum		2.26	0.22		90		
Actinomyces israelii		0.30	0.13		1		
07733033	**Very good identification**						**07733033**
Actinomyces naeslundii		100.0	0.68	MLZ 5			
07733040	**Very good identification**						**07733040**
					CAT		
Actinomyces naeslundii		96.76	0.74		25		
Actinomyces israelii		2.45	0.38		1		
Propionibacterium granulosum		0.79	0.24		90		
07733041	**Very good identification**						**07733041**
Actinomyces naeslundii		100.0	0.77				
07733042	**Very good identification**						**07733042**
					CAT		
Actinomyces naeslundii		96.07	0.74		25		
Actinomyces israelii		3.64	0.41		1		
Propionibacterium granulosum		0.20	0.13		90		
07733043	**Very good identification**						**07733043**
Actinomyces naeslundii		100.0	0.77				
07733050	**Acceptable identification**						**07733050**
					CAT		
Actinomyces naeslundii		91.50	0.52	MLZ 5	25		
Actinomyces israelii		6.57	0.24		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		1.94	0.09		90	
07733051	**Good identification**					**07733051**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
07733052	**Acceptable identification**					**07733052**
Actinomyces naeslundii		90.28	0.52	MLZ 5		
Actinomyces israelii		9.72	0.27			
07733053	**Good identification**					**07733053**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
07733060	**Very good identification**					**07733060**
					CAT	
Actinomyces naeslundii		98.84	0.74		25	
Clostridium perfringens		0.80	0.16		1	
Propionibacterium granulosum		0.20	0.13		90	
07733061	**Very good identification**					**07733061**
Actinomyces naeslundii		100.0	0.77			
07733062	**Very good identification**					**07733062**
Actinomyces naeslundii		99.29	0.74			
Clostridium perfringens		0.38	0.11			
Actinomyces israelii		0.24	0.20			
07733063	**Very good identification**					**07733063**
Actinomyces naeslundii		100.0	0.77			
07733070	**Good identification**					**07733070**
Actinomyces naeslundii		100.0	0.52	MLZ 5		
07733071	**Good identification**					**07733071**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
07733072	**Good identification**					**07733072**
Actinomyces naeslundii		100.0	0.52	MLZ 5		
07733073	**Good identification**					**07733073**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
07733100	**Very good identification**					**07733100**
Actinomyces naeslundii		97.65	0.95			
Actinomyces israelii		1.57	0.55			
Propionibacterium propionicum		0.31	0.42			
07733101	**Excellent identification**					**07733101**
Actinomyces naeslundii		99.97	0.98			
Propionibacterium propionicum		0.02	0.26			
Actinomyces israelii		0.01	0.21			
07733102	**Good identification**					**07733102**
Actinomyces naeslundii		94.59	0.95			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		2.29	0.58			
Bifidobacterium adolescentis		1.77	0.50			
07733103	**Excellent identification**					**07733103**
Actinomyces naeslundii		99.94	0.98			
Propionibacterium propionicum		0.04	0.29			
Actinomyces israelii		0.02	0.24			
07733110	**Good identification**					**07733110**
Actinomyces naeslundii		94.88	0.73	MLZ 5		
Actinomyces israelii		4.34	0.41			
Bifidobacterium adolescentis		0.42	0.17			
07733111	**Very good identification**					**07733111**
Actinomyces naeslundii		100.0	0.76	MLZ 5		
07733112	**Low discrimination**					**07733112**
Actinomyces naeslundii		65.68	0.73	MLZ 5		
Bifidobacterium adolescentis		28.57	0.51			
Actinomyces israelii		4.51	0.44			
07733113	**Very good identification**					**07733113**
Actinomyces naeslundii		99.71	0.76	MLZ 5		
Bifidobacterium adolescentis		0.29	0.17			
07733120	**Good identification**					**07733120**
					SPO	
Actinomyces naeslundii		92.41	0.95		1	
Clostridium ramosum		5.65	0.49		92	
Clostridium perfringens		1.78	0.43		84	
07733121	**Excellent identification**					**07733121**
					SPO	
Actinomyces naeslundii		99.95	0.98		1	
Clostridium ramosum		0.04	0.15		92	
Clostridium perfringens		0.01	0.09		84	
07733122	**Low discrimination**					**07733122**
					SPO	
Actinomyces naeslundii		78.96	0.95		1	
Clostridium ramosum		19.33	0.59		92	
Clostridium perfringens		0.71	0.38		84	
07733123	**Excellent identification**					**07733123**
					SPO	
Actinomyces naeslundii		99.83	0.98		1	
Clostridium ramosum		0.17	0.25		92	
Clostridium perfringens		0.01	0.03		84	
07733130	**Very good identification**					**07733130**
					SPO	
Actinomyces naeslundii		97.32	0.73	MLZ 5	1	
Clostridium ramosum		1.14	0.15		92	
Clostridium tertium		0.72	0.13		99	
07733131	**Very good identification**					**07733131**
Actinomyces naeslundii		100.0	0.76	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07733132	**Low discrimination**					**07733132**
					SPO	
Actinomyces naeslundii		89.56	0.73	MLZ 5	1	
Clostridium ramosum		4.21	0.25		92	
Clostridium tertium		1.98	0.21		99	
07733133	**Very good identification**					**07733133**
Actinomyces naeslundii		100.0	0.76	MLZ 5		
07733140	**Good identification**					**07733140**
Actinomyces naeslundii		94.88	0.82			
Actinomyces israelii		3.37	0.48			
Propionibacterium propionicum		1.69	0.42			
07733141	**Very good identification**					**07733141**
Actinomyces naeslundii		99.85	0.85			
Propionibacterium propionicum		0.13	0.26			
Actinomyces israelii		0.02	0.14			
07733142	**Good identification**					**07733142**
Actinomyces naeslundii		90.92	0.82			
Actinomyces israelii		4.85	0.51			
Propionibacterium propionicum		2.43	0.45			
07733143	**Very good identification**					**07733143**
Actinomyces naeslundii		99.77	0.85			
Propionibacterium propionicum		0.20	0.29			
Actinomyces israelii		0.04	0.17			
07733150	**Good identification**					**07733150**
Actinomyces naeslundii		90.84	0.60	MLZ 5		
Actinomyces israelii		9.16	0.34			
07733151	**Very good identification**					**07733151**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
07733152	**Low discrimination**					**07733152**
Actinomyces naeslundii		66.05	0.60	MLZ 5		
Bifidobacterium adolescentis		20.13	0.36			
Actinomyces israelii		9.99	0.37			
07733153	**Very good identification**					**07733153**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
07733160	**Very good identification**					**07733160**
					SPO	
Actinomyces naeslundii		97.37	0.82		1	
Clostridium perfringens		2.02	0.31		84	
Clostridium ramosum		0.34	0.15		92	
07733161	**Very good identification**					**07733161**
Actinomyces naeslundii		100.0	0.85			
07733162	**Very good identification**					**07733162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		96.49	0.82		1		
Clostridium ramosum		1.35	0.25		92		
Clostridium perfringens		0.94	0.25		84		
07733163	**Very good identification**						**07733163**
Actinomyces naeslundii		100.0	0.85				
07733170	**Very good identification**						**07733170**
Actinomyces naeslundii		98.36	0.60	MLZ 5			
Bifidobacterium breve		1.00	0.12				
Actinomyces israelii		0.63	0.13				
07733171	**Very good identification**						**07733171**
Actinomyces naeslundii		100.0	0.63	MLZ 5			
07733172	**Low discrimination**						**07733172**
Actinomyces naeslundii		89.87	0.60	MLZ 5			
Bifidobacterium breve		9.26	0.29				
Actinomyces israelii		0.87	0.16				
07733173	**Very good identification**						**07733173**
Actinomyces naeslundii		100.0	0.63	MLZ 5			
07733200	**Low discrimination**						**07733200**
					CAT		
Actinomyces naeslundii		65.65	0.53	RHA 1	25		
Propionibacterium granulosum		28.74	0.32		90		
Actinomyces israelii		5.61	0.26		1		
07733201	**Good identification**						**07733201**
Actinomyces naeslundii		100.0	0.56	RHA 1			
07733202	**Low discrimination**						**07733202**
					CAT		
Actinomyces naeslundii		80.80	0.53	RHA 1	25		
Actinomyces israelii		10.36	0.29		1		
Propionibacterium granulosum		8.84	0.22		90		
07733203	**Good identification**						**07733203**
Actinomyces naeslundii		100.0	0.56	RHA 1			
07733220	**Low discrimination**						**07733220**
					SPO	CAT	
Actinomyces naeslundii		74.61	0.53	RHA 1	1	25	
Clostridium ramosum		17.22	0.17		92	1	
Propionibacterium granulosum		8.16	0.22		1	90	
07733221	**Good identification**						**07733221**
Actinomyces naeslundii		100.0	0.56	RHA 1			
07733222	**Low discrimination**						**07733222**
					SPO	CAT	
Actinomyces naeslundii		51.26	0.53	RHA 1	1	25	
Clostridium ramosum		47.34	0.27		92	1	
Propionibacterium granulosum		1.40	0.11		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07733223	**Good identification**					**07733223**
Actinomyces naeslundii		100.0	0.56	RHA 1		
07733240	**Low discrimination**					**07733240**
Actinomyces naeslundii		84.15	0.40	RHA 1		
Actinomyces israelii		15.85	0.18			
07733241	**Good identification**					**07733241**
Actinomyces naeslundii		100.0	0.43	RHA 1		
07733242	**Low discrimination**					**07733242**
Actinomyces naeslundii		77.97	0.40	RHA 1		
Actinomyces israelii		22.03	0.22			
07733243	**Good identification**					**07733243**
Actinomyces naeslundii		100.0	0.43	RHA 1		
07733260	**Good identification**					**07733260**
Actinomyces naeslundii		100.0	0.40	RHA 1		
07733261	**Good identification**					**07733261**
Actinomyces naeslundii		100.0	0.43	RHA 1		
07733262	**Good identification**					**07733262**
Actinomyces naeslundii		100.0	0.40	RHA 1		
07733263	**Good identification**					**07733263**
Actinomyces naeslundii		100.0	0.43	RHA 1		
07733300	**Low discrimination**					**07733300**
					SPO	
Actinomyces naeslundii		83.25	0.61	RHA 1	1	
Actinomyces israelii		10.00	0.36		1	
Clostridium ramosum		6.75	0.17		92	
07733301	**Very good identification**					**07733301**
Actinomyces naeslundii		100.0	0.64	RHA 1		
07733302	**Low discrimination**					**07733302**
					SPO	
Actinomyces naeslundii		63.96	0.61	RHA 1	1	
Clostridium ramosum		20.75	0.27		92	
Actinomyces israelii		11.52	0.39		1	
07733303	**Very good identification**					**07733303**
Actinomyces naeslundii		100.0	0.64	RHA 1		
07733310	**Low discrimination**					**07733310**
Actinomyces naeslundii		74.57	0.39	RHA 1		
Actinomyces israelii		25.43	0.22	MLZ 5		
07733311	**Good identification**					**07733311**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.42	RHA 1			
07733312	**Low discrimination**						**07733312**
Actinomyces naeslundii		45.30	0.39	RHA 1			
Actinomyces israelii		23.17	0.25	MLZ 5			
Bifidobacterium adolescentis		19.71	0.17				
07733313	**Good identification**						**07733313**
Actinomyces naeslundii		100.0	0.42	RHA 1			
07733320	**Low discrimination**						**07733320**
					SPO		
Clostridium ramosum		88.66	0.51	CEL 99	92		
Actinomyces naeslundii		11.04	0.61		1		
Clostridium perfringens		0.21	0.09		84		
07733321	**Good identification**						**07733321**
					SPO		
Actinomyces naeslundii		94.87	0.64	RHA 1	1		
Clostridium ramosum		5.13	0.17		92		
07733322	**Very good identification**						**07733322**
					SPO	GRA	
Clostridium ramosum		95.00	0.61	CEL 99	92	50	
Actinomyces naeslundii		2.96	0.61		1	99	
Parabacteroides distasonis		1.61	0.43		1	1	
07733323	**Low discrimination**						**07733323**
					SPO		
Actinomyces naeslundii		82.22	0.64	RHA 1	1		
Clostridium ramosum		17.78	0.27		92		
07733331	**Good identification**						**07733331**
Actinomyces naeslundii		100.0	0.42	RHA 1			
07733333	**Good identification**						**07733333**
Actinomyces naeslundii		100.0	0.42	RHA 1			
07733340	**Low discrimination**						**07733340**
Actinomyces naeslundii		79.07	0.48	RHA 1			
Actinomyces israelii		20.93	0.29				
07733341	**Good identification**						**07733341**
Actinomyces naeslundii		100.0	0.51	RHA 1			
07733342	**Low discrimination**						**07733342**
Actinomyces naeslundii		71.58	0.48	RHA 1			
Actinomyces israelii		28.42	0.32				
07733343	**Good identification**						**07733343**
Actinomyces naeslundii		100.0	0.51	RHA 1			
07733360	**Low discrimination**						**07733360**
					SPO		
Actinomyces naeslundii		68.51	0.48	RHA 1	1		
Clostridium ramosum		31.49	0.17		92		
07733361	**Good identification**						**07733361**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	RHA 1		
07733363	**Good identification**					**07733363**
Actinomyces naeslundii		100.0	0.51	RHA 1		
07733400	**Low discrimination**					**07733400**
					CAT	
Actinomyces naeslundii		48.26	0.53	ARA 1	25	
Actinomyces israelii		44.84	0.43		1	
Propionibacterium granulosum		6.90	0.24		90	
07733401	**Good identification**					**07733401**
Actinomyces naeslundii		100.0	0.56	ARA 1		
07733402	**Low discrimination**					**07733402**
Actinomyces israelii		50.02	0.46	CEL 93		
Actinomyces naeslundii		35.89	0.53	XYL 92		
Bifidobacterium adolescentis		12.81	0.30			
07733403	**Good identification**					**07733403**
Actinomyces naeslundii		100.0	0.56	ARA 1		
07733420	**Acceptable identification**					**07733420**
					CAT	
Actinomyces naeslundii		91.32	0.53	ARA 1	25	
Actinomyces israelii		5.42	0.23		1	
Propionibacterium granulosum		3.26	0.13		90	
07733421	**Good identification**					**07733421**
Actinomyces naeslundii		100.0	0.56	ARA 1		
07733422	**Acceptable identification**					**07733422**
Actinomyces naeslundii		91.83	0.53	ARA 1		
Actinomyces israelii		8.17	0.26			
07733423	**Good identification**					**07733423**
Actinomyces naeslundii		100.0	0.56	ARA 1		
07733440	**Low discrimination**					**07733440**
Actinomyces israelii		67.19	0.36	CEL 93		
Actinomyces naeslundii		32.81	0.40	XYL 92		
07733441	**Good identification**					**07733441**
Actinomyces naeslundii		100.0	0.43	ARA 1		
07733442	**Low discrimination**					**07733442**
Actinomyces israelii		71.07	0.39	CEL 93		
Actinomyces naeslundii		23.14	0.40	XYL 92		
Bifidobacterium adolescentis		5.78	0.14			
07733443	**Good identification**					**07733443**
Actinomyces naeslundii		100.0	0.43	ARA 1		
07733460	**Low discrimination**					**07733460**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		88.44	0.40	ARA 1		
Actinomyces israelii		11.56	0.16			
07733461	**Good identification**					**07733461**
Actinomyces naeslundii		100.0	0.43	ARA 1		
07733462	**Low discrimination**					**07733462**
Actinomyces naeslundii		83.61	0.40	ARA 1		
Actinomyces israelii		16.39	0.19			
07733463	**Good identification**					**07733463**
Actinomyces naeslundii		100.0	0.43	ARA 1		
07733500	**Low discrimination**					**07733500**
Actinomyces israelii		52.53	0.54	CEL 93		
Actinomyces naeslundii		40.23	0.61	XYL 92		
Bifidobacterium adolescentis		5.04	0.30			
07733501	**Very good identification**					**07733501**
Actinomyces naeslundii		99.13	0.64	ARA 1		
Actinomyces israelii		0.87	0.19			
07733502	**Low discrimination**					**07733502**
Bifidobacterium adolescentis		80.61	0.64	XYL 87		
Actinomyces israelii		12.72	0.57			
Actinomyces naeslundii		6.49	0.61			
07733503	**Good identification**					**07733503**
Actinomyces naeslundii		91.18	0.64	ARA 1		
Bifidobacterium adolescentis		7.62	0.30			
Actinomyces israelii		1.20	0.22			
07733510	**Low discrimination**					**07733510**
Actinomyces israelii		48.66	0.39	CEL 93		
Bifidobacterium adolescentis		38.21	0.31	XYL 92		
Actinomyces naeslundii		13.13	0.39	MLZ 13		
07733511	**Good identification**					**07733511**
Actinomyces naeslundii		100.0	0.42	ARA 1		
07733512	**Very good identification**					**07733512**
Bifidobacterium adolescentis		97.53	0.66	XYL 87		
Actinomyces israelii		1.88	0.42			
Actinomyces naeslundii		0.34	0.39			
07733520	**Low discrimination**					**07733520**
					SPO	
Actinomyces naeslundii		85.21	0.61	ARA 1	1	
Actinomyces israelii		7.10	0.33		1	
Clostridium ramosum		5.21	0.15		92	
07733521	**Very good identification**					**07733521**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.64	ARA 1		
07733522	**Low discrimination**					**07733522**
					SPO	
Actinomyces naeslundii		61.30	0.61	ARA 1	1	
Clostridium ramosum		15.00	0.25		92	
Bifidobacterium adolescentis		7.69	0.30		1	
07733523	**Very good identification**					**07733523**
Actinomyces naeslundii		100.0	0.64	ARA 1		
07733530	**Low discrimination**					**07733530**
Actinomyces naeslundii		64.33	0.39	ARA 1		
Bifidobacterium longum subsp. longum		20.45	0.13	MLZ 5		
Actinomyces israelii		15.22	0.19			
07733531	**Good identification**					**07733531**
Actinomyces naeslundii		100.0	0.42	ARA 1		
07733533	**Good identification**					**07733533**
Actinomyces naeslundii		100.0	0.42	ARA 1		
07733540	**Low discrimination**					**07733540**
Actinomyces israelii		72.57	0.47	CEL 93		
Actinomyces naeslundii		25.22	0.48	XYL 92		
Bifidobacterium adolescentis		2.22	0.14			
07733541	**Good identification**					**07733541**
Actinomyces naeslundii		100.0	0.51	ARA 1		
07733542	**Low discrimination**					**07733542**
Bifidobacterium adolescentis		62.05	0.48	XYL 87		
Actinomyces israelii		30.81	0.50	SOR 11		
Actinomyces naeslundii		7.14	0.48			
07733543	**Acceptable identification**					**07733543**
Actinomyces naeslundii		91.95	0.51	ARA 1		
Bifidobacterium adolescentis		5.38	0.14			
Actinomyces israelii		2.67	0.15			
07733552	**Acceptable identification**					**07733552**
Bifidobacterium adolescentis		93.84	0.50	XYL 87		
Actinomyces israelii		5.70	0.35	SOR 11		
Actinomyces naeslundii		0.47	0.26			
07733560	**Low discrimination**					**07733560**
Actinomyces naeslundii		84.48	0.48	ARA 1		
Actinomyces israelii		15.52	0.26			
07733561	**Good identification**					**07733561**
Actinomyces naeslundii		100.0	0.51	ARA 1		
07733562	**Low discrimination**					**07733562**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		73.35	0.48	ARA 1			
Actinomyces israelii		20.21	0.29				
Bifidobacterium adolescentis		6.44	0.14				
07733563	**Good identification**						**07733563**
Actinomyces naeslundii		100.0	0.51	ARA 1			
07733702	**Low discrimination**						**07733702**
Actinomyces israelii		44.66	0.37	CEL 93			
Bifidobacterium adolescentis		37.98	0.30	XYL 92			
Clostridium clostridioforme		14.31	0.19	RHA 7			
07733722	**Low discrimination**						**07733722**
					SPO		
Parabacteroides distasonis		45.96	0.40	MAN 1	1		
Clostridium ramosum		44.82	0.27	XYL 90	92		
Clostridium clostridioforme		6.53	0.19		66		
07734000	**Low discrimination**						**07734000**
					GRA	CAT	
Fusobacterium mortiferum		85.75	0.44	MLT 11	1	5	
Propionibacterium granulosum		11.53	0.26	XYL 7	99	90	
Actinomyces israelii		1.42	0.16		99	1	
07734100	**Low discrimination**						**07734100**
					COC	GRA	
Fusobacterium mortiferum		88.79	0.50	MLT 11	1	1	
Atopobium parvulum		5.68	0.18	XYL 7	99	99	
Actinomyces israelii		2.51	0.26		1	99	
07734101	**Low discrimination**						**07734101**
					GRA		
Actinomyces naeslundii		73.40	0.35	GAL 96	99		
Fusobacterium mortiferum		26.61	0.16	XYL 1	1		
07734103	**Good identification**						**07734103**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07734112	**Low discrimination**						**07734112**
					GRA		
Clostridium clostridioforme		75.44	0.39	RHA 88	50		
Bifidobacterium adolescentis		22.60	0.33	ARA 94	99		
Parabacteroides distasonis		1.07	0.13		1		
07734121	**Good identification**						**07734121**
Actinomyces naeslundii		100.0	0.35	GAL 96			
07734122	**Low discrimination**						**07734122**
Parabacteroides distasonis		89.02	0.51	GAL 99			
Clostridium clostridioforme		7.50	0.27				
Prevotella oralis		1.35	0.08				
07734123	**Low discrimination**						**07734123**
					GRA		
Actinomyces naeslundii		66.19	0.35	GAL 96	99		
Parabacteroides distasonis		33.81	0.17	XYL 1	1		
07734132	**Low discrimination**						**07734132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		58.38	0.47	GAL 99		
Clostridium clostridioforme		41.62	0.39			
07734212	**Good identification**					**07734212**
Clostridium clostridioforme		100.0	0.44	ARA 94		
07734232	**Good identification**					**07734232**
Clostridium clostridioforme		98.10	0.44	ARA 94		
Parabacteroides distasonis		1.90	0.21			
07734302	**Good identification**					**07734302**
Clostridium clostridioforme		95.33	0.42	ARA 94		
Parabacteroides distasonis		4.67	0.25			
07734312	**Good identification**					**07734312**
Clostridium clostridioforme		99.42	0.53	ARA 94		
Parabacteroides distasonis		0.58	0.21			
07734322	**Low discrimination**					**07734322**
Parabacteroides distasonis		82.77	0.60	GAL 99		
Clostridium clostridioforme		17.05	0.42			
Prevotella oralis		0.18	0.01			
07734332	**Low discrimination**					**07734332**
Clostridium clostridioforme		63.54	0.53	ARA 94		
Parabacteroides distasonis		36.46	0.55			
07734402	**Good identification**					**07734402**
Clostridium clostridioforme		96.92	0.39	RHA 88		
Bifidobacterium adolescentis		1.56	0.11			
Actinomyces israelii		1.52	0.17			
07734412	**Good identification**					**07734412**
Clostridium clostridioforme		99.57	0.50	RHA 88		
Bifidobacterium adolescentis		0.43	0.13			
07734422	**Good identification**					**07734422**
Clostridium clostridioforme		98.45	0.39	RHA 88		
Parabacteroides distasonis		1.55	0.14			
07734432	**Good identification**					**07734432**
Clostridium clostridioforme		100.0	0.50	RHA 88		
07734502	**Low discrimination**					**07734502**
					SPO	
Clostridium clostridioforme		65.24	0.48	RHA 88	66	
Bifidobacterium adolescentis		31.12	0.45		1	
Clostridium butyricum		1.89	0.10		99	
07734512	**Low discrimination**					**07734512**
Clostridium clostridioforme		88.51	0.59	RHA 88		
Bifidobacterium adolescentis		11.33	0.47			
Bifidobacterium longum subsp. longum		0.11	0.11			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07734522	**Low discrimination**					**07734522**
					GRA	
Clostridium clostridioforme		67.44	0.48	RHA 88	50	
Parabacteroides distasonis		31.31	0.48		1	
Bifidobacterium longum subsp. longum		0.93	0.17		99	
07734532	**Good identification**					**07734532**
					GRA	
Clostridium clostridioforme		94.23	0.59	RHA 88	50	
Parabacteroides distasonis		5.17	0.43		1	
Bifidobacterium longum subsp. longum		0.48	0.21		99	
07734602	**Good identification**					**07734602**
Clostridium clostridioforme		100.0	0.54			
07734612	**Very good identification**					**07734612**
Clostridium clostridioforme		100.0	0.65			
07734622	**Good identification**					**07734622**
Clostridium clostridioforme		99.36	0.54			
Parabacteroides distasonis		0.64	0.22			
07734632	**Very good identification**					**07734632**
Clostridium clostridioforme		99.92	0.65			
Parabacteroides distasonis		0.08	0.17			
07734702	**Very good identification**					**07734702**
					GRA	
Clostridium clostridioforme		99.74	0.63		50	
Parabacteroides distasonis		0.19	0.22		1	
Bifidobacterium adolescentis		0.07	0.11		99	
07734710	**Good identification**					**07734710**
Clostridium clostridioforme		100.0	0.40	BGL 99		
07734712	**Very good identification**					**07734712**
					GRA	
Clostridium clostridioforme		99.96	0.74		50	
Parabacteroides distasonis		0.02	0.17		1	
Bifidobacterium adolescentis		0.02	0.13		99	
07734713	**Good identification**					**07734713**
Clostridium clostridioforme		100.0	0.40	URE 1		
07734722	**Low discrimination**					**07734722**
Clostridium clostridioforme		83.98	0.63			
Parabacteroides distasonis		15.95	0.56			
Bacteroides thetaiotaomicron		0.04	-.05			
07734730	**Good identification**					**07734730**
Clostridium clostridioforme		97.01	0.40	BGL 99		
Parabacteroides distasonis		2.18	0.17			
Bacteroides vulgatus		0.81	-.09			
07734732	**Very good identification**					**07734732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		97.81	0.74			
Parabacteroides distasonis		2.20	0.51			
07734733	**Good identification**					**07734733**
Clostridium clostridioforme		97.81	0.40	URE 1		
Parabacteroides distasonis		2.20	0.17			
07734752	**Good identification**					**07734752**
Clostridium clostridioforme		100.0	0.40	SOR 1		
07734772	**Good identification**					**07734772**
Clostridium clostridioforme		97.81	0.40	SOR 1		
Parabacteroides distasonis		2.20	0.17			
07735000	**Low discrimination**					**07735000**
					GRA	
Actinomyces israelii		40.81	0.50	CEL 93	99	
Fusobacterium mortiferum		37.44	0.47		1	
Actinomyces naeslundii		9.12	0.48		99	
07735001	**Acceptable identification**					**07735001**
					GRA	
Actinomyces naeslundii		94.54	0.51	XYL 1	99	
Actinomyces israelii		2.85	0.16		99	
Fusobacterium mortiferum		2.61	0.13		1	
07735002	**Low discrimination**					**07735002**
Actinomyces israelii		65.01	0.53	CEL 93		
Actinomyces naeslundii		9.69	0.48			
Bifidobacterium adolescentis		8.20	0.31			
07735003	**Good identification**					**07735003**
Actinomyces naeslundii		95.68	0.51	XYL 1		
Actinomyces israelii		4.32	0.19			
07735010	**Low discrimination**					**07735010**
					CAT	
Actinomyces israelii		78.83	0.36	CEL 93	1	
Actinomyces naeslundii		6.21	0.26	MLZ 13	25	
Propionibacterium granulosum		5.92	0.11		90	
07735012	**Low discrimination**					**07735012**
Clostridium clostridioforme		59.13	0.38	RHA 88		
Bifidobacterium adolescentis		19.77	0.33	ARA 94		
Actinomyces israelii		19.16	0.39			
07735021	**Good identification**					**07735021**
Actinomyces naeslundii		100.0	0.51	XYL 1		
07735022	**Low discrimination**					**07735022**
					GRA	
Parabacteroides distasonis		70.63	0.51	LAC 99	1	
Actinomyces naeslundii		7.85	0.48		99	
Prevotella oralis		6.78	0.21		1	
07735023	**Acceptable identification**					**07735023**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		94.29	0.51	XYL 1	99		
Parabacteroides distasonis		5.71	0.17		1		
07735032	**Low discrimination**						**07735032**
					GRA		
Parabacteroides distasonis		60.13	0.47	LAC 99	1		
Clostridium clostridioforme		38.42	0.38		50		
Actinomyces israelii		0.80	0.18		99		
07735040	**Low discrimination**						**07735040**
					GRA		
Actinomyces israelii		87.30	0.43	CEL 93	99		
Actinomyces naeslundii		8.86	0.35		99		
Fusobacterium mortiferum		2.08	0.13		1		
07735041	**Good identification**						**07735041**
Actinomyces naeslundii		100.0	0.38	XYL 1			
07735042	**Acceptable identification**						**07735042**
Actinomyces israelii		90.28	0.46	CEL 93			
Actinomyces naeslundii		6.11	0.35				
Bifidobacterium adolescentis		3.62	0.16				
07735043	**Good identification**						**07735043**
Actinomyces naeslundii		100.0	0.38	XYL 1			
07735061	**Good identification**						**07735061**
Actinomyces naeslundii		100.0	0.38	XYL 1			
07735062	**Low discrimination**						**07735062**
Actinomyces naeslundii		31.38	0.35	XYL 1			
Actinomyces israelii		29.62	0.26				
Clostridium perfringens		22.84	0.11				
07735063	**Good identification**						**07735063**
Actinomyces naeslundii		100.0	0.38	XYL 1			
07735100	**Low discrimination**						**07735100**
					COC	SPO	
Atopobium parvulum		49.22	0.53	RAF 1	99	1	
Actinomyces israelii		21.77	0.60		1	1	
Clostridium butyricum		11.85	0.35		1	99	
07735101	**Low discrimination**						**07735101**
					COC		
Actinomyces naeslundii		84.48	0.59	XYL 1	1		
Atopobium parvulum		8.09	0.18		99		
Actinomyces israelii		3.58	0.26		1		
07735102	**Low discrimination**						**07735102**
					GRA		
Bifidobacterium adolescentis		59.47	0.66	ARA 87	99		
Actinomyces israelii		19.05	0.63		99		
Parabacteroides distasonis		6.38	0.51		1		
07735103	**Low discrimination**						**07735103**
Actinomyces naeslundii		76.40	0.59	XYL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		15.15	0.31				
Actinomyces israelii		4.85	0.29				
07735110	**Low discrimination**						**07735110**
					SPO		
Actinomyces israelii		52.44	0.46	CEL 93	1		
Bifidobacterium adolescentis		20.29	0.33	MLZ 13	1		
Clostridium butyricum		10.05	0.13		99		
07735111	**Good identification**						**07735111**
Actinomyces naeslundii		100.0	0.37	XYL 1			
07735112	**Low discrimination**						**07735112**
					GRA		
Bifidobacterium adolescentis		83.84	0.67	ARA 87	99		
Clostridium clostridioforme		8.48	0.47		50		
Parabacteroides distasonis		3.96	0.47		1		
07735120	**Low discrimination**						**07735120**
					COC	GRA	
Clostridium perfringens		29.38	0.43	XYL 1	1	99	
Parabacteroides distasonis		25.14	0.51		1	1	
Atopobium parvulum		19.96	0.40		99	99	
07735121	**Good identification**						**07735121**
					GRA		
Actinomyces naeslundii		94.07	0.59	XYL 1	99		
Clostridium perfringens		2.34	0.09		99		
Parabacteroides distasonis		2.00	0.17		1		
07735122	**Very good identification**						**07735122**
					GRA		
Parabacteroides distasonis		96.38	0.86		1		
Prevotella oralis		1.47	0.42		1		
Clostridium perfringens		0.54	0.38		99		
07735123	**Low discrimination**						**07735123**
					GRA		
Parabacteroides distasonis		66.65	0.51	URE 1	1		
Actinomyces naeslundii		31.63	0.59		99		
Prevotella oralis		1.01	0.08		1		
07735130	**Low discrimination**						**07735130**
					GRA		
Parabacteroides distasonis		83.96	0.47	BGL 99	1		
Bifidobacterium longum subsp. longum		4.81	0.21		99		
Actinomyces israelii		2.96	0.26		99		
07735131	**Low discrimination**						**07735131**
					GRA		
Actinomyces naeslundii		82.12	0.37	XYL 1	99		
Parabacteroides distasonis		17.88	0.13	MLZ 5	1		
07735132	**Very good identification**						**07735132**
					GRA		
Parabacteroides distasonis		97.50	0.81		1		
Clostridium clostridioforme		2.11	0.47		50		
Bifidobacterium adolescentis		0.21	0.33		99		
07735133	**Acceptable identification**						**07735133**
					GRA		
Parabacteroides distasonis		93.63	0.47	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		4.34	0.37		99	
Clostridium clostridioforme		2.02	0.13		50	
07735140	**Low discrimination**					**07735140**
					SPO	
Actinomyces israelii		77.49	0.53	CEL 93	1	
Clostridium butyricum		9.43	0.17		99	
Actinomyces naeslundii		5.59	0.43		1	
07735141	**Acceptable identification**					**07735141**
Actinomyces naeslundii		91.47	0.46	XYL 1		
Actinomyces israelii		8.53	0.19			
07735142	**Low discrimination**					**07735142**
Actinomyces israelii		47.75	0.56	CEL 93		
Bifidobacterium adolescentis		47.38	0.50			
Actinomyces naeslundii		2.30	0.43			
07735143	**Low discrimination**					**07735143**
Actinomyces naeslundii		78.20	0.46	XYL 1		
Actinomyces israelii		10.94	0.22			
Bifidobacterium adolescentis		10.86	0.16			
07735150	**Low discrimination**					**07735150**
					SPO	
Actinomyces israelii		85.77	0.39	CEL 93	1	
Bifidobacterium adolescentis		10.55	0.17	MLZ 13	1	
Clostridium butyricum		3.68	-.05		99	
07735152	**Low discrimination**					**07735152**
Bifidobacterium adolescentis		88.08	0.51	ARA 87		
Actinomyces israelii		10.85	0.42	SOR 11		
Clostridium clostridioforme		0.73	0.13			
07735161	**Good identification**					**07735161**
Actinomyces naeslundii		100.0	0.46	XYL 1		
07735162	**Low discrimination**					**07735162**
					GRA	
Parabacteroides distasonis		78.47	0.51	SOR 1	1	
Clostridium perfringens		8.22	0.25		99	
Actinomyces israelii		5.81	0.36		99	
07735163	**Low discrimination**					**07735163**
					GRA	
Actinomyces naeslundii		89.24	0.46	XYL 1	99	
Parabacteroides distasonis		10.76	0.17		1	
07735172	**Acceptable identification**					**07735172**
					GRA	
Parabacteroides distasonis		93.58	0.47	SOR 1	1	
Bifidobacterium adolescentis		2.47	0.17		99	
Clostridium clostridioforme		2.02	0.13		50	
07735202	**Low discrimination**					**07735202**
					GRA	
Clostridium clostridioforme		86.50	0.41	ARA 94	50	
Actinomyces israelii		8.77	0.34		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		4.73	0.25		1	
07735212	**Good identification**					**07735212**
					GRA	
Clostridium clostridioforme		99.03	0.53	ARA 94	50	
Parabacteroides distasonis		0.64	0.21		1	
Actinomyces israelii		0.33	0.20		99	
07735222	**Low discrimination**					**07735222**
Parabacteroides distasonis		83.37	0.60	LAC 99		
Clostridium clostridioforme		15.39	0.41			
Prevotella oralis		1.14	0.15			
07735232	**Low discrimination**					**07735232**
Clostridium clostridioforme		60.97	0.53	ARA 94		
Parabacteroides distasonis		39.04	0.55			
07735300	**Low discrimination**					**07735300**
					COC	
Actinomyces israelii		34.74	0.41	CEL 93	1	
Eubacterium contortum		26.62	0.24	RHA 7	1	
Atopobium parvulum		10.54	0.18		99	
07735302	**Low discrimination**					**07735302**
					GRA	
Parabacteroides distasonis		57.11	0.60	NAG 99	1	
Clostridium clostridioforme		35.30	0.50		50	
Actinomyces israelii		4.28	0.44		99	
07735312	**Low discrimination**					**07735312**
					GRA	
Clostridium clostridioforme		82.72	0.62	ARA 94	50	
Parabacteroides distasonis		15.82	0.55		1	
Bifidobacterium adolescentis		1.13	0.33		99	
07735320	**Low discrimination**					**07735320**
Parabacteroides distasonis		51.11	0.60	BGL 99		
Bacteroides vulgatus		47.00	0.41			
Prevotella oralis		0.58	0.14			
07735322	**Excellent identification**					**07735322**
Parabacteroides distasonis		99.13	0.94			
Clostridium clostridioforme		0.62	0.50			
Prevotella oralis		0.22	0.35			
07735323	**Very good identification**					**07735323**
Parabacteroides distasonis		99.01	0.60	URE 1		
Clostridium clostridioforme		0.62	0.16			
Prevotella oralis		0.22	0.01			
07735330	**Acceptable identification**					**07735330**
Parabacteroides distasonis		93.45	0.55	BGL 99		
Clostridium clostridioforme		4.94	0.27			
Bacteroides vulgatus		1.61	0.06			
07735332	**Good identification**					**07735332**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		94.98	0.89			
Clostridium clostridioforme		5.02	0.62			
Prevotella oralis		0.00	0.01			
07735333	**Acceptable identification**					**07735333**
Parabacteroides distasonis		94.98	0.55	URE 1		
Clostridium clostridioforme		5.02	0.27			
07735342	**Low discrimination**					**07735342**
					GRA	
Actinomyces israelii		59.01	0.37	CEL 93	99	
Parabacteroides distasonis		20.47	0.25	RHA 7	1	
Clostridium clostridioforme		12.65	0.16		50	
07735362	**Very good identification**					**07735362**
Parabacteroides distasonis		98.98	0.60	SOR 1		
Clostridium clostridioforme		0.62	0.16			
Prevotella oralis		0.22	0.01			
07735372	**Acceptable identification**					**07735372**
Parabacteroides distasonis		94.98	0.55	SOR 1		
Clostridium clostridioforme		5.02	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07735400	**Low discrimination**					**07735400**
					SPO	
Actinomyces israelii		48.09	0.49	CEL 93	1	
Clostridium butyricum		47.99	0.27		99	
Clostridium clostridioforme		1.41	0.13		66	
07735402	**Low discrimination**					**07735402**
Clostridium clostridioforme		45.55	0.47	RHA 88		
Bifidobacterium adolescentis		24.25	0.45			
Actinomyces israelii		23.50	0.52			
07735412	**Low discrimination**					**07735412**
Clostridium clostridioforme		86.01	0.58	RHA 88		
Bifidobacterium adolescentis		12.28	0.47			
Actinomyces israelii		1.46	0.37			
07735422	**Low discrimination**					**07735422**
					GRA	
Clostridium clostridioforme		63.16	0.47	RHA 88	50	
Parabacteroides distasonis		32.72	0.48		1	
Actinomyces israelii		2.08	0.31		99	
07735432	**Good identification**					**07735432**
					GRA	
Clostridium clostridioforme		93.50	0.58	RHA 88	50	
Parabacteroides distasonis		5.73	0.43		1	
Bifidobacterium longum subsp. longum		0.54	0.21		99	
07735440	**Low discrimination**					**07735440**
					SPO	
Actinomyces israelii		81.76	0.41	CEL 93	1	
Clostridium butyricum		18.24	0.09		99	
07735442	**Low discrimination**					**07735442**
					SPO	
Actinomyces israelii		69.63	0.45	CEL 93	1	
Bifidobacterium adolescentis		22.84	0.30		1	
Clostridium butyricum		4.03	0.02		99	
07735500	**Low discrimination**					**07735500**
					SPO	
Clostridium butyricum		77.32	0.51	ESL 88	99	
Actinomyces israelii		12.92	0.59	CEL 99	1	
Bifidobacterium adolescentis		5.00	0.45		1	
07735502	**Low discrimination**					**07735502**
					SPO	
Bifidobacterium adolescentis		84.69	0.80		1	
Clostridium clostridioforme		5.38	0.56		66	
Clostridium butyricum		5.15	0.44		99	
07735503	**Low discrimination**					**07735503**
					SPO	
Bifidobacterium adolescentis		84.17	0.45	URE 1	1	
Clostridium clostridioforme		5.35	0.22		66	
Clostridium butyricum		5.12	0.10		99	
07735510	**Low discrimination**					**07735510**
					SPO	
Bifidobacterium adolescentis		31.15	0.47	BGL 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		30.87	0.45		1		
Clostridium butyricum		20.75	0.29		99		
07735512	**Low discrimination**						**07735512**
Bifidobacterium adolescentis		79.45	0.81				
Clostridium clostridioforme		18.81	0.67				
Bifidobacterium longum subsp. longum		0.80	0.45				
07735513	**Low discrimination**						**07735513**
Bifidobacterium adolescentis		79.90	0.47	URE 1			
Clostridium clostridioforme		18.92	0.33				
Bifidobacterium longum subsp. longum		0.80	0.11				
07735520	**Low discrimination**						**07735520**
					GRA	SPO	
Bacteroides vulgatus		54.34	0.43	TRE 4	1	1	
Bifidobacterium longum subsp. longum		23.91	0.52	RHA 90	99	1	
Clostridium butyricum		11.31	0.33		99	99	
07735522	**Low discrimination**						**07735522**
					GRA		
Parabacteroides distasonis		89.69	0.82		1		
Clostridium clostridioforme		5.86	0.56		50		
Bifidobacterium longum subsp. longum		2.65	0.52		99		
07735523	**Low discrimination**						**07735523**
					GRA		
Parabacteroides distasonis		89.85	0.48	URE 1	1		
Clostridium clostridioforme		5.87	0.22		50		
Bifidobacterium longum subsp. longum		2.65	0.17		99		
07735530	**Low discrimination**						**07735530**
					GRA		
Bifidobacterium longum subsp. longum		83.08	0.56	SAL 1	99		
Parabacteroides distasonis		8.99	0.43		1		
Clostridium clostridioforme		4.96	0.33		50		
07735532	**Low discrimination**						**07735532**
					GRA		
Parabacteroides distasonis		59.83	0.77		1		
Clostridium clostridioforme		33.04	0.67		50		
Bifidobacterium longum subsp. longum		5.59	0.56		99		
07735533	**Low discrimination**						**07735533**
					GRA		
Parabacteroides distasonis		59.91	0.43	URE 1	1		
Clostridium clostridioforme		33.09	0.33		50		
Bifidobacterium longum subsp. longum		5.60	0.21		99		
07735542	**Low discrimination**						**07735542**
					SPO		
Bifidobacterium adolescentis		85.29	0.64	SOR 11	1		
Actinomyces israelii		10.51	0.55		1		
Clostridium butyricum		3.65	0.26		99		
07735552	**Very good identification**						**07735552**
Bifidobacterium adolescentis		96.43	0.66	SOR 11			
Clostridium clostridioforme		1.87	0.33				
Actinomyces israelii		1.45	0.41				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07735562	**Low discrimination**					**07735562**
					GRA	
Parabacteroides distasonis		73.13	0.48	SOR 1	1	
Bifidobacterium adolescentis		9.29	0.30		99	
Actinomyces israelii		7.23	0.34		99	
07735572	**Low discrimination**					**07735572**
					GRA	
Parabacteroides distasonis		50.98	0.43	SOR 1	1	
Clostridium clostridioforme		28.16	0.33		50	
Bifidobacterium adolescentis		14.70	0.31		99	
07735602	**Very good identification**					**07735602**
					GRA	
Clostridium clostridioforme		99.13	0.62		50	
Actinomyces israelii		0.53	0.32		99	
Parabacteroides distasonis		0.21	0.22		1	
07735610	**Good identification**					**07735610**
Clostridium clostridioforme		98.87	0.39	BGL 99		
Actinomyces israelii		1.13	0.15			
07735612	**Very good identification**					**07735612**
					GRA	
Clostridium clostridioforme		99.94	0.73		50	
Parabacteroides distasonis		0.03	0.17		1	
Actinomyces israelii		0.02	0.18		99	
07735613	**Good identification**					**07735613**
Clostridium clostridioforme		100.0	0.39	URE 1		
07735622	**Low discrimination**					**07735622**
Clostridium clostridioforme		82.41	0.62			
Parabacteroides distasonis		17.47	0.56			
Bacteroides thetaiotaomicron		0.09	0.00			
07735630	**Good identification**					**07735630**
Clostridium clostridioforme		96.67	0.39	BGL 99		
Parabacteroides distasonis		2.42	0.17			
Bacteroides vulgatus		0.91	-.09			
07735632	**Very good identification**					**07735632**
Clostridium clostridioforme		97.56	0.73			
Parabacteroides distasonis		2.44	0.51			
07735633	**Good identification**					**07735633**
Clostridium clostridioforme		97.56	0.39	URE 1		
Parabacteroides distasonis		2.44	0.17			
07735652	**Good identification**					**07735652**
Clostridium clostridioforme		100.0	0.39	SOR 1		
07735672	**Good identification**					**07735672**
Clostridium clostridioforme		97.56	0.39	SOR 1		
Parabacteroides distasonis		2.44	0.17			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07735702	**Good identification**					**07735702**
					GRA	
Clostridium clostridioforme		91.21	0.71		50	
Parabacteroides distasonis		5.77	0.56		1	
Bifidobacterium adolescentis		1.98	0.45		99	
07735703	**Acceptable identification**					**07735703**
					GRA	
Clostridium clostridioforme		92.17	0.37	URE 1	50	
Parabacteroides distasonis		5.84	0.22		1	
Bifidobacterium adolescentis		2.00	0.11		99	
07735710	**Good identification**					**07735710**
					SPO	
Clostridium clostridioforme		95.47	0.48	BGL 99	66	
Actinomyces israelii		1.31	0.25		1	
Clostridium butyricum		1.13	0.03		99	
07735712	**Very good identification**					**07735712**
					GRA	
Clostridium clostridioforme		98.66	0.82		50	
Parabacteroides distasonis		0.74	0.51		1	
Bifidobacterium adolescentis		0.57	0.47		99	
07735713	**Good identification**					**07735713**
					GRA	
Clostridium clostridioforme		98.69	0.48	URE 1	50	
Parabacteroides distasonis		0.74	0.17		1	
Bifidobacterium adolescentis		0.57	0.13		99	
07735720	**Good identification**					**07735720**
Bacteroides vulgatus		94.33	0.60	TRE 4		
Parabacteroides distasonis		4.73	0.56			
Clostridium clostridioforme		0.76	0.37			
07735722	**Low discrimination**					**07735722**
Parabacteroides distasonis		85.82	0.90			
Clostridium clostridioforme		13.69	0.71			
Bacteroides thetaiotaomicron		0.21	0.29			
07735723	**Low discrimination**					**07735723**
Parabacteroides distasonis		85.91	0.56	URE 1		
Clostridium clostridioforme		13.71	0.37			
Bacteroides thetaiotaomicron		0.21	-.05			
07735730	**Low discrimination**					**07735730**
Clostridium clostridioforme		49.00	0.48	BGL 99		
Parabacteroides distasonis		36.30	0.51			
Bacteroides vulgatus		13.57	0.26			
07735732	**Low discrimination**					**07735732**
Clostridium clostridioforme		57.36	0.82			
Parabacteroides distasonis		42.49	0.86			
Bacteroides thetaiotaomicron		0.07	0.22			
07735733	**Low discrimination**					**07735733**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		57.45	0.48	URE 1		
Parabacteroides distasonis		42.55	0.51			
07735742	**Low discrimination**					**07735742**
Clostridium clostridioforme		62.23	0.37	SOR 1		
Bifidobacterium adolescentis		16.51	0.30			
Actinomyces israelii		15.16	0.36			
07735752	**Acceptable identification**					**07735752**
Clostridium clostridioforme		92.03	0.48	SOR 1		
Bifidobacterium adolescentis		6.55	0.31			
Actinomyces israelii		0.74	0.21			
07735762	**Low discrimination**					**07735762**
Parabacteroides distasonis		85.43	0.56	SOR 1		
Clostridium clostridioforme		13.63	0.37			
Bacteroides ovatus		0.35	-.01			
07735772	**Low discrimination**					**07735772**
Clostridium clostridioforme		57.32	0.48	SOR 1		
Parabacteroides distasonis		42.46	0.51			
Bacteroides ovatus		0.22	-.04			
07736100	**Low discrimination**					**07736100**
					GRA	
Actinomyces israelii		77.66	0.39	CEL 93	99	
Actinomyces naeslundii		17.45	0.37	GAL 99	99	
Fusobacterium mortiferum		4.89	0.16		1	
07736101	**Good identification**					**07736101**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07736102	**Low discrimination**					**07736102**
Actinomyces israelii		62.47	0.42	CEL 93		
Bifidobacterium adolescentis		28.17	0.30	GAL 99		
Actinomyces naeslundii		9.36	0.37			
07736103	**Good identification**					**07736103**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07736120	**Low discrimination**					**07736120**
Actinomyces naeslundii		77.88	0.37	GAL 96		
Actinomyces israelii		22.12	0.19	XYL 1		
07736121	**Good identification**					**07736121**
Actinomyces naeslundii		100.0	0.40	GAL 96		
07736122	**Low discrimination**					**07736122**
					GRA	
Actinomyces naeslundii		54.92	0.37	GAL 96	99	
Actinomyces israelii		23.40	0.22	XYL 1	99	
Parabacteroides distasonis		21.68	0.17		1	
07736123	**Good identification**					**07736123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	GAL 96		
07736142	**Low discrimination**					**07736142**
Actinomyces israelii		82.56	0.35	CEL 93		
Bifidobacterium adolescentis		11.83	0.14	GAL 99		
Actinomyces naeslundii		5.61	0.24			
07736500	**Low discrimination**					**07736500**
					SPO	
Actinomyces israelii		85.23	0.38	CEL 93	1	
Clostridium butyricum		10.00	0.01	GAL 99	99	
Bifidobacterium adolescentis		4.76	0.10		1	
07736502	**Low discrimination**					**07736502**
Bifidobacterium adolescentis		76.61	0.44	GAL 99		
Actinomyces israelii		20.78	0.41			
Clostridium clostridioforme		1.98	0.14			
07736512	**Low discrimination**					**07736512**
Bifidobacterium adolescentis		87.74	0.45	GAL 99		
Clostridium clostridioforme		8.46	0.25			
Actinomyces israelii		2.91	0.26			
07736712	**Good identification**					**07736712**
Clostridium clostridioforme		98.59	0.40	MAN 1		
Bifidobacterium adolescentis		1.41	0.11			
07736732	**Good identification**					**07736732**
Clostridium clostridioforme		97.81	0.40	MAN 1		
Parabacteroides distasonis		2.20	0.17			
07737000	**Good identification**					**07737000**
					CAT	
Actinomyces israelii		91.66	0.63	CEL 93	1	
Actinomyces naeslundii		7.02	0.53		25	
Propionibacterium granulosum		1.00	0.24		90	
07737001	**Acceptable identification**					**07737001**
Actinomyces naeslundii		91.92	0.56	XYL 1		
Actinomyces israelii		8.08	0.29			
07737002	**Good identification**					**07737002**
Actinomyces israelii		93.36	0.66	CEL 93		
Actinomyces naeslundii		4.77	0.53			
Bifidobacterium adolescentis		1.70	0.30			
07737003	**Low discrimination**					**07737003**
Actinomyces naeslundii		88.35	0.56	XYL 1		
Actinomyces israelii		11.65	0.32			
07737010	**Good identification**					**07737010**
					CAT	
Actinomyces israelii		96.44	0.49	CEL 93	1	
Actinomyces naeslundii		2.60	0.31	MLZ 13	25	
Propionibacterium granulosum		0.96	0.09		90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07737012	**Low discrimination**					**07737012**
Actinomyces israelii		85.69	0.52	CEL 93		
Bifidobacterium adolescentis		12.77	0.31	MLZ 13		
Actinomyces naeslundii		1.54	0.31			
07737020	**Low discrimination**					**07737020**
					CAT	
Actinomyces naeslundii		53.50	0.53	XYL 1	25	
Actinomyces israelii		44.59	0.42		1	
Propionibacterium granulosum		1.91	0.13		90	
07737021	**Good identification**					**07737021**
Actinomyces naeslundii		100.0	0.56	XYL 1		
07737022	**Low discrimination**					**07737022**
					GRA	
Actinomyces israelii		54.43	0.45	CEL 93	99	
Actinomyces naeslundii		43.53	0.53	NAG 6	99	
Parabacteroides distasonis		2.04	0.17		1	
07737023	**Good identification**					**07737023**
Actinomyces naeslundii		100.0	0.56	XYL 1		
07737040	**Good identification**					**07737040**
Actinomyces israelii		96.64	0.56	CEL 93		
Actinomyces naeslundii		3.36	0.40			
07737041	**Low discrimination**					**07737041**
Actinomyces naeslundii		83.77	0.43	XYL 1		
Actinomyces israelii		16.23	0.22			
07737042	**Very good identification**					**07737042**
Actinomyces israelii		97.19	0.59	CEL 93		
Actinomyces naeslundii		2.25	0.40			
Bifidobacterium adolescentis		0.56	0.14			
07737043	**Low discrimination**					**07737043**
Actinomyces naeslundii		77.48	0.43	XYL 1		
Actinomyces israelii		22.52	0.25			
07737050	**Good identification**					**07737050**
Actinomyces israelii		100.0	0.42	CEL 93		
07737052	**Good identification**					**07737052**
Actinomyces israelii		95.48	0.45	CEL 93		
Bifidobacterium adolescentis		4.52	0.16	MLZ 13		
07737060	**Low discrimination**					**07737060**
Actinomyces israelii		64.75	0.35	CEL 93		
Actinomyces naeslundii		35.25	0.40	NAG 6		
07737061	**Good identification**					**07737061**
Actinomyces naeslundii		100.0	0.43	XYL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07737062	**Low discrimination**					**07737062**
Actinomyces israelii		73.37	0.38	CEL 93		
Actinomyces naeslundii		26.63	0.40	NAG 6		
07737063	**Good identification**					**07737063**
Actinomyces naeslundii		100.0	0.43	XYL 1		
07737100	**Good identification**					**07737100**
					SPO	
Actinomyces israelii		92.81	0.73	CEL 93	1	
Actinomyces naeslundii		5.06	0.61		1	
Clostridium butyricum		0.99	0.18		99	
07737101	**Low discrimination**					**07737101**
Actinomyces naeslundii		89.00	0.64	XYL 1		
Actinomyces israelii		11.00	0.39			
07737102	**Low discrimination**					**07737102**
Actinomyces israelii		67.02	0.76	CEL 93		
Bifidobacterium adolescentis		30.22	0.64			
Actinomyces naeslundii		2.43	0.61			
07737103	**Low discrimination**					**07737103**
Actinomyces naeslundii		78.80	0.64	XYL 1		
Actinomyces israelii		14.61	0.42			
Bifidobacterium adolescentis		6.59	0.30			
07737110	**Good identification**					**07737110**
Actinomyces israelii		92.14	0.59	CEL 93		
Bifidobacterium adolescentis		5.15	0.31	MLZ 13		
Actinomyces naeslundii		1.77	0.39			
07737111	**Low discrimination**					**07737111**
Actinomyces naeslundii		74.03	0.42	XYL 1		
Actinomyces israelii		25.97	0.25	MLZ 5		
07737112	**Low discrimination**					**07737112**
Bifidobacterium adolescentis		78.08	0.66	ARA 87		
Actinomyces israelii		21.18	0.62			
Actinomyces naeslundii		0.27	0.39			
07737120	**Low discrimination**					**07737120**
					SPO	
Actinomyces israelii		50.32	0.53	CEL 93	1	
Actinomyces naeslundii		42.95	0.61	NAG 6	1	
Clostridium ramosum		2.63	0.15		92	
07737121	**Very good identification**					**07737121**
Actinomyces naeslundii		99.22	0.64	XYL 1		
Actinomyces israelii		0.78	0.19			
07737122	**Low discrimination**					**07737122**
					GRA	
Actinomyces israelii		35.91	0.56	CEL 93	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		33.27	0.51	NAG 6	1		
Actinomyces naeslundii		20.43	0.61		99		
07737123	**Very good identification**						**07737123**
					GRA		
Actinomyces naeslundii		97.77	0.64	XYL 1	99		
Actinomyces israelii		1.16	0.22		99		
Parabacteroides distasonis		1.07	0.17		1		
07737130	**Low discrimination**						**07737130**
					SPO		
Actinomyces israelii		60.43	0.39	CEL 93	1		
Clostridium tertium		18.34	0.15	MLZ 13	99		
Actinomyces naeslundii		18.17	0.39	NAG 6	1		
07737131	**Good identification**						**07737131**
Actinomyces naeslundii		100.0	0.42	XYL 1			
07737132	**Low discrimination**						**07737132**
					GRA	SPO	
Parabacteroides distasonis		56.86	0.47	MAN 1	1	1	
Actinomyces israelii		17.03	0.42		99	1	
Clostridium tertium		10.34	0.24		99	99	
07737133	**Low discrimination**						**07737133**
					GRA		
Actinomyces naeslundii		89.92	0.42	XYL 1	99		
Parabacteroides distasonis		10.09	0.13	MLZ 5	1		
07737140	**Very good identification**						**07737140**
					SPO		
Actinomyces israelii		97.16	0.66	CEL 93	1		
Actinomyces naeslundii		2.40	0.48		1		
Clostridium butyricum		0.23	0.00		99		
07737141	**Low discrimination**						**07737141**
Actinomyces naeslundii		78.59	0.51	XYL 1			
Actinomyces israelii		21.41	0.32				
07737142	**Low discrimination**						**07737142**
Actinomyces israelii		86.18	0.69	CEL 93			
Bifidobacterium adolescentis		12.35	0.48				
Actinomyces naeslundii		1.42	0.48				
07737143	**Low discrimination**						**07737143**
Actinomyces naeslundii		68.16	0.51	XYL 1			
Actinomyces israelii		27.85	0.35				
Bifidobacterium adolescentis		3.99	0.14				
07737150	**Good identification**						**07737150**
					COC		
Actinomyces israelii		96.52	0.52	CEL 93	1		
Bifidobacterium adolescentis		1.71	0.16	MLZ 13	1		
Blautia producta		0.92	-.03		99		
07737152	**Low discrimination**						**07737152**
Bifidobacterium adolescentis		53.82	0.50	ARA 87			
Actinomyces israelii		45.92	0.55	SOR 11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.27	0.26				
07737160	**Low discrimination**						**07737160**
Actinomyces israelii		72.08	0.46	CEL 93			
Actinomyces naeslundii		27.92	0.48	NAG 6			
07737161	**Good identification**						**07737161**
Actinomyces naeslundii		100.0	0.51	XYL 1			
07737162	**Low discrimination**						**07737162**
					GRA		
Actinomyces israelii		76.63	0.49	CEL 93	99		
Actinomyces naeslundii		19.79	0.48	NAG 6	99		
Parabacteroides distasonis		1.84	0.17		1		
07737163	**Good identification**						**07737163**
Actinomyces naeslundii		97.46	0.51	XYL 1			
Actinomyces israelii		2.54	0.15				
07737200	**Good identification**						**07737200**
Actinomyces israelii		100.0	0.44	CEL 93			
07737202	**Good identification**						**07737202**
Actinomyces israelii		98.27	0.47	CEL 93			
Clostridium clostridioforme		1.73	0.07	RHA 7			
07737240	**Good identification**						**07737240**
Actinomyces israelii		100.0	0.37	CEL 93			
07737242	**Good identification**						**07737242**
Actinomyces israelii		100.0	0.40	CEL 93			
07737300	**Good identification**						**07737300**
					SPO		
Actinomyces israelii		98.84	0.54	CEL 93	1		
Actinomyces naeslundii		0.72	0.27	RHA 7	1		
Clostridium butyricum		0.43	-.08		99		
07737302	**Good identification**						**07737302**
					GRA		
Actinomyces israelii		90.59	0.57	CEL 93	99		
Bifidobacterium adolescentis		5.48	0.30	RHA 7	99		
Parabacteroides distasonis		2.16	0.25		1		
07737310	**Good identification**						**07737310**
Actinomyces israelii		100.0	0.40	CEL 93			
07737312	**Low discrimination**						**07737312**
Actinomyces israelii		49.28	0.43	CEL 93			
Bifidobacterium adolescentis		24.39	0.31	RHA 7			
Clostridium clostridioforme		22.11	0.27	MLZ 13			
07737322	**Good identification**						**07737322**
					SPO	GRA	
Parabacteroides distasonis		90.58	0.60	MAN 1	1	1	
Clostridium ramosum		6.02	0.27		92	50	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		2.45	0.37		1	99	
07737332	**Acceptable identification**						**07737332**
					GRA		
Parabacteroides distasonis		94.31	0.55	MAN 1	1		
Clostridium clostridioforme		4.98	0.27		50		
Actinomyces israelii		0.71	0.22		99		
07737340	**Good identification**						**07737340**
Actinomyces israelii		100.0	0.47	CEL 93			
07737342	**Good identification**						**07737342**
Actinomyces israelii		98.11	0.50	CEL 93			
Bifidobacterium adolescentis		1.89	0.14	RHA 7			
07737352	**Low discrimination**						**07737352**
Actinomyces israelii		86.41	0.36	CEL 93			
Bifidobacterium adolescentis		13.59	0.16	RHA 7			
07737400	**Very good identification**						**07737400**
					SPO		
Actinomyces israelii		97.86	0.61	CEL 93	1		
Clostridium butyricum		1.92	0.11		99		
Bifidobacterium adolescentis		0.22	0.10		1		
07737402	**Low discrimination**						**07737402**
					SPO		
Actinomyces israelii		86.39	0.64	CEL 93	1		
Bifidobacterium adolescentis		12.87	0.44		1		
Clostridium butyricum		0.44	0.04		99		
07737410	**Good identification**						**07737410**
Actinomyces israelii		98.19	0.47	CEL 93			
Bifidobacterium adolescentis		1.81	0.11	MLZ 13			
07737412	**Low discrimination**						**07737412**
Bifidobacterium adolescentis		52.16	0.45	LAC 99			
Actinomyces israelii		42.80	0.50				
Clostridium clostridioforme		4.51	0.24				
07737420	**Good identification**						**07737420**
					SPO		
Actinomyces israelii		97.40	0.41	CEL 93	1		
Clostridium butyricum		2.60	-.07	NAG 6	99		
07737422	**Acceptable identification**						**07737422**
					GRA		
Actinomyces israelii		90.44	0.44	CEL 93	99		
Clostridium clostridioforme		4.90	0.13	NAG 6	50		
Parabacteroides distasonis		2.54	0.14		1		
07737440	**Good identification**						**07737440**
					SPO		
Actinomyces israelii		99.57	0.54	CEL 93	1		
Clostridium butyricum		0.44	-.07		99		
07737442	**Very good identification**						**07737442**
Actinomyces israelii		95.48	0.57	CEL 93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		4.52	0.28				
07737450	**Good identification**						**07737450**
					COC		
Actinomyces israelii		95.53	0.40	CEL 93	1		
Blautia producta		4.47	-.03	MLZ 13	99		
07737452	**Low discrimination**						**07737452**
Actinomyces israelii		72.08	0.43	CEL 93			
Bifidobacterium adolescentis		27.92	0.30	MLZ 13			
07737462	**Good identification**						**07737462**
Actinomyces israelii		100.0	0.37	CEL 93			
07737500	**Low discrimination**						**07737500**
					SPO		
Actinomyces israelii		85.15	0.72	CEL 93	1		
Clostridium butyricum		10.00	0.35		99		
Bifidobacterium adolescentis		4.76	0.44		1		
07737501	**Low discrimination**						**07737501**
					SPO		
Actinomyces israelii		78.55	0.38	CEL 93	1		
Clostridium butyricum		9.22	0.01	URE 1	99		
Actinomyces naeslundii		7.85	0.30		1		
07737502	**Low discrimination**						**07737502**
					SPO		
Bifidobacterium adolescentis		78.08	0.78		1		
Actinomyces israelii		21.18	0.75		1		
Clostridium butyricum		0.64	0.28		99		
07737503	**Low discrimination**						**07737503**
Bifidobacterium adolescentis		77.07	0.44	URE 1			
Actinomyces israelii		20.90	0.41				
Actinomyces naeslundii		1.39	0.30				
07737510	**Low discrimination**						**07737510**
Actinomyces israelii		64.37	0.58	CEL 93			
Bifidobacterium adolescentis		29.42	0.45	MLZ 13			
Bifidobacterium dentium		2.91	0.10				
07737512	**Very good identification**						**07737512**
Bifidobacterium adolescentis		95.54	0.80				
Actinomyces israelii		3.17	0.61				
Bifidobacterium dentium		0.96	0.27				
07737513	**Good identification**						**07737513**
Bifidobacterium adolescentis		95.86	0.45	URE 1			
Actinomyces israelii		3.18	0.26				
Bifidobacterium dentium		0.97	-.07				
07737520	**Low discrimination**						**07737520**
					GRA	SPO	
Actinomyces israelii		77.49	0.51	CEL 93	99	1	
Clostridium butyricum		12.39	0.17	NAG 6	99	99	
Bacteroides vulgatus		5.42	0.09		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07737522	**Low discrimination**						**07737522**
					GRA		
Actinomyces israelii		41.59	0.54	CEL 93	99		
Parabacteroides distasonis		28.86	0.48	NAG 6	1		
Bifidobacterium adolescentis		24.27	0.44		99		
07737540	**Good identification**						**07737540**
					COC	SPO	
Actinomyces israelii		90.77	0.65	CEL 93	1	1	
Blautia producta		5.23	0.24		99	1	
Clostridium butyricum		2.38	0.17		1	99	
07737542	**Low discrimination**						**07737542**
					SPO		
Bifidobacterium adolescentis		53.69	0.63	SOR 11	1		
Actinomyces israelii		45.81	0.68		1		
Clostridium butyricum		0.31	0.10		99		
07737552	**Low discrimination**						**07737552**
					COC		
Bifidobacterium adolescentis		89.76	0.64	SOR 11	1		
Actinomyces israelii		9.36	0.54		1		
Blautia producta		0.80	0.15		99		
07737560	**Low discrimination**						**07737560**
					COC	SPO	
Actinomyces israelii		88.02	0.44	CEL 93	1	1	
Blautia producta		8.83	0.07	NAG 6	99	1	
Clostridium butyricum		3.15	-.02		1	99	
07737562	**Low discrimination**						**07737562**
					GRA		
Actinomyces israelii		83.09	0.47	CEL 93	99		
Bifidobacterium adolescentis		15.41	0.28	NAG 6	99		
Parabacteroides distasonis		1.50	0.14		1		
07737600	**Good identification**						**07737600**
Actinomyces israelii		100.0	0.42	CEL 93			
07737602	**Low discrimination**						**07737602**
Actinomyces israelii		73.71	0.45	CEL 93			
Clostridium clostridioforme		24.82	0.28	RHA 7			
Bifidobacterium adolescentis		1.47	0.10				
07737612	**Low discrimination**						**07737612**
Clostridium clostridioforme		89.82	0.39	MAN 1			
Actinomyces israelii		8.75	0.31				
Bifidobacterium adolescentis		1.43	0.11				
07737632	**Good identification**						**07737632**
Clostridium clostridioforme		97.56	0.39	MAN 1			
Parabacteroides distasonis		2.44	0.17				
07737640	**Good identification**						**07737640**
Actinomyces israelii		100.0	0.35	CEL 93			
07737642	**Good identification**						**07737642**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.38	CEL 93		
07737700	**Acceptable identification**					**07737700**
					SPO	
Actinomyces israelii		94.24	0.52	CEL 93	1	
Clostridium butyricum		4.55	0.09	RHA 7	99	
Bifidobacterium adolescentis		0.71	0.10		1	
07737702	**Low discrimination**					**07737702**
Actinomyces israelii		55.34	0.55	CEL 93		
Bifidobacterium adolescentis		27.39	0.44	RHA 7		
Clostridium clostridioforme		15.59	0.37			
07737710	**Low discrimination**					**07737710**
Actinomyces israelii		79.88	0.38	CEL 93		
Clostridium clostridioforme		10.40	0.14	RHA 7		
Bifidobacterium adolescentis		4.90	0.11	MLZ 13		
07737712	**Low discrimination**					**07737712**
Clostridium clostridioforme		62.51	0.48	MAN 1		
Bifidobacterium adolescentis		29.45	0.45			
Actinomyces israelii		7.28	0.41			
07737722	**Low discrimination**					**07737722**
					GRA	
Parabacteroides distasonis		80.90	0.56	MAN 1	1	
Clostridium clostridioforme		12.91	0.37		50	
Actinomyces israelii		2.92	0.35		99	
07737732	**Low discrimination**					**07737732**
Clostridium clostridioforme		55.81	0.48	MAN 1		
Parabacteroides distasonis		41.34	0.51			
Bacteroides ovatus		1.70	0.12			
07737740	**Low discrimination**					**07737740**
					COC	
Actinomyces israelii		56.63	0.45	CEL 93	1	
Blautia producta		43.37	0.24	RHA 7	99	
07737742	**Low discrimination**					**07737742**
					COC	
Actinomyces israelii		82.37	0.48	CEL 93	1	
Bifidobacterium adolescentis		12.96	0.28	RHA 7	1	
Blautia producta		4.67	0.07		99	
07740001	**Good identification**					**07740001**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07740003	**Low discrimination**					**07740003**
Actinomyces naeslundii		52.97	0.37	FRU 95		
Actinomyces viscosus		47.03	0.12	GAL 96		
07740021	**Good identification**					**07740021**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07740023	**Good identification**					**07740023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	FRU 95		
07740100	**Low discrimination**					**07740100**
					GRA	
Fusobacterium mortiferum		69.38	0.36	MLT 11	1	
Actinomyces naeslundii		30.62	0.42	FRU 99	99	
07740101	**Good identification**					**07740101**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07740103	**Low discrimination**					**07740103**
Actinomyces naeslundii		64.33	0.45	FRU 95		
Actinomyces viscosus		35.68	0.16	GAL 96		
07740120	**Low discrimination**					**07740120**
					GRA	
Actinomyces naeslundii		80.50	0.42	FRU 95	99	
Prevotella melaninogenica		13.27	-.01	GAL 96	1	
Prevotella oralis		6.23	-.03		1	
07740121	**Good identification**					**07740121**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07740123	**Good identification**					**07740123**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07741000	**Low discrimination**					**07741000**
Actinomyces naeslundii		57.58	0.58	FRU 95		
Lactobacillus catenaformis		21.62	0.29			
Lactobacillus jensenii		10.85	0.24			
07741001	**Very good identification**					**07741001**
Actinomyces naeslundii		100.0	0.61	FRU 95		
07741002	**Low discrimination**					**07741002**
Actinomyces naeslundii		47.43	0.58	FRU 95		
Lactobacillus acidophilus		19.41	0.26			
Lactobacillus catenaformis		17.81	0.29			
07741003	**Very good identification**					**07741003**
Actinomyces naeslundii		97.78	0.61	FRU 95		
Actinomyces viscosus		2.22	0.08			
07741011	**Good identification**					**07741011**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07741012	**Low discrimination**					**07741012**
Actinomyces naeslundii		55.68	0.36	FRU 95		
Lactobacillus jensenii		44.32	0.12	MLZ 5		
07741013	**Good identification**					**07741013**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07741020	**Low discrimination**					**07741020**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		80.23	0.58	FRU 95	99	
Lactobacillus catenaformis		7.53	0.18		99	
Prevotella oralis		4.65	0.11		1	
07741021	**Very good identification**					**07741021**
Actinomyces naeslundii		100.0	0.61	FRU 95		
07741023	**Very good identification**					**07741023**
					GRA	
Actinomyces naeslundii		99.26	0.61	FRU 95	99	
Prevotella oralis		0.74	-.02		1	
07741030	**Low discrimination**					**07741030**
Actinomyces naeslundii		77.01	0.36	FRU 95		
Lactobacillus jensenii		22.99	0.05	MLZ 5		
07741031	**Good identification**					**07741031**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07741032	**Low discrimination**					**07741032**
					GRA	
Actinomyces naeslundii		70.90	0.36	FRU 95	99	
Prevotella oralis		15.00	-.02	MLZ 5	1	
Lactobacillus jensenii		14.11	0.02		99	
07741033	**Good identification**					**07741033**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07741040	**Good identification**					**07741040**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07741041	**Good identification**					**07741041**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07741042	**Low discrimination**					**07741042**
Actinomyces naeslundii		77.72	0.45	FRU 95		
Lactobacillus acidophilus		22.28	0.11			
07741043	**Good identification**					**07741043**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07741060	**Good identification**					**07741060**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07741061	**Good identification**					**07741061**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07741062	**Low discrimination**					**07741062**
					GRA	
Actinomyces naeslundii		74.09	0.45	FRU 95	99	
Lactobacillus acidophilus		21.24	0.11		99	
Prevotella oralis		4.67	-.02		1	
07741063	**Good identification**					**07741063**
Actinomyces naeslundii		100.0	0.48	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07741100	**Low discrimination**					**07741100**
					GRA	
Actinomyces naeslundii		70.88	0.65	FRU 95	99	
Lactobacillus catenaformis		14.03	0.32		99	
Fusobacterium mortiferum		10.04	0.39		1	
07741101	**Very good identification**					**07741101**
Actinomyces naeslundii		100.0	0.68	FRU 95		
07741102	**Low discrimination**					**07741102**
					GRA	
Lactobacillus acidophilus		49.15	0.45	FRU 99	99	
Actinomyces naeslundii		29.72	0.65		99	
Prevotella oralis		9.49	0.31		1	
07741103	**Very good identification**					**07741103**
Actinomyces naeslundii		97.33	0.68	FRU 95		
Actinomyces viscosus		1.38	0.12			
Lactobacillus acidophilus		1.08	0.10			
07741110	**Acceptable identification**					**07741110**
Actinomyces naeslundii		92.42	0.43	FRU 95		
Bifidobacterium breve		7.58	0.10	MLZ 5		
07741111	**Good identification**					**07741111**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07741112	**Low discrimination**					**07741112**
Actinomyces naeslundii		31.23	0.43	FRU 95		
Bifidobacterium adolescentis		28.88	0.27	MLZ 5		
Bifidobacterium breve		25.89	0.27			
07741113	**Good identification**					**07741113**
Actinomyces naeslundii		100.0	0.46	FRU 95		
07741120	**Low discrimination**					**07741120**
					GRA	
Actinomyces naeslundii		45.16	0.65	FRU 95	99	
Prevotella melaninogenica		30.71	0.33		1	
Prevotella oralis		14.42	0.31		1	
07741121	**Very good identification**					**07741121**
					GRA	
Actinomyces naeslundii		99.33	0.68	FRU 95	99	
Prevotella melaninogenica		0.46	-.01		1	
Prevotella oralis		0.21	-.03		1	
07741122	**Low discrimination**					**07741122**
					GRA	
Prevotella oralis		65.66	0.53	FRU 99	1	
Lactobacillus acidophilus		17.90	0.45		99	
Actinomyces naeslundii		10.82	0.65		99	
07741123	**Very good identification**					**07741123**
					GRA	
Actinomyces naeslundii		95.06	0.68	FRU 95	99	
Prevotella oralis		3.88	0.19		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		1.06	0.10		99		
07741130	**Low discrimination**						**07741130**
					GRA		
Actinomyces naeslundii		74.77	0.43	FRU 95	99		
Bifidobacterium breve		10.90	0.15	MLZ 5	99		
Prevotella melaninogenica		9.76	-.01		1		
07741131	**Good identification**						**07741131**
Actinomyces naeslundii		100.0	0.46	FRU 95			
07741133	**Good identification**						**07741133**
Actinomyces naeslundii		100.0	0.46	FRU 95			
07741140	**Good identification**						**07741140**
Actinomyces naeslundii		97.61	0.52	FRU 95			
Bifidobacterium breve		2.39	0.10				
07741141	**Good identification**						**07741141**
Actinomyces naeslundii		100.0	0.55	FRU 95			
07741143	**Good identification**						**07741143**
Actinomyces naeslundii		100.0	0.55	FRU 95			
07741160	**Low discrimination**						**07741160**
					GRA		
Actinomyces naeslundii		86.83	0.52	FRU 95	99		
Bifidobacterium breve		3.77	0.15		99		
Prevotella melaninogenica		3.38	-.01		1		
07741161	**Good identification**						**07741161**
Actinomyces naeslundii		100.0	0.55	FRU 95			
07741163	**Good identification**						**07741163**
Actinomyces naeslundii		100.0	0.55	FRU 95			
07741322	**Low discrimination**						**07741322**
					SPO	GRA	
Prevotella oralis		87.48	0.47	FRU 99	1	1	
Clostridium ramosum		6.64	0.28		92	50	
Lactobacillus acidophilus		4.19	0.25		1	99	
07741502	**Low discrimination**						**07741502**
Bifidobacterium adolescentis		85.99	0.40	FRU 99			
Lactobacillus acidophilus		5.39	0.10	XYL 87			
Actinomyces naeslundii		3.26	0.31				
07741512	**Good identification**						**07741512**
Bifidobacterium adolescentis		97.81	0.41	FRU 99			
Clostridium clostridioforme		2.20	0.10	XYL 87			
07742000	**Good identification**						**07742000**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07742001	**Good identification**						**07742001**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	FRU 95		
07742002	**Good identification**					**07742002**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07742003	**Good identification**					**07742003**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07742020	**Good identification**					**07742020**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07742021	**Good identification**					**07742021**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07742022	**Good identification**					**07742022**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07742023	**Good identification**					**07742023**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07742100	**Good identification**					**07742100**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07742101	**Good identification**					**07742101**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07742102	**Good identification**					**07742102**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07742103	**Good identification**					**07742103**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07742120	**Good identification**					**07742120**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07742121	**Good identification**					**07742121**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07742122	**Acceptable identification**					**07742122**
					SPO	
Actinomyces naeslundii		90.47	0.47	FRU 95	1	
Clostridium ramosum		9.53	0.05	GAL 96	92	
07742123	**Good identification**					**07742123**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07742141	**Good identification**					**07742141**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07742143	**Good identification**					**07742143**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07742161	**Good identification**					**07742161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	FRU 95		
07742163	**Good identification**					**07742163**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07743000	**Very good identification**					**07743000**
					SPO	
Actinomyces naeslundii		96.80	0.62	FRU 95	1	
Clostridium innocuum		2.10	0.10		99	
Actinomyces israelii		1.10	0.20		1	
07743001	**Very good identification**					**07743001**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07743002	**Very good identification**					**07743002**
					SPO	
Actinomyces naeslundii		95.25	0.62	FRU 95	1	
Clostridium innocuum		2.06	0.10		99	
Actinomyces israelii		1.63	0.23		1	
07743003	**Very good identification**					**07743003**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07743010	**Good identification**					**07743010**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07743011	**Good identification**					**07743011**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07743012	**Good identification**					**07743012**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07743013	**Good identification**					**07743013**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07743020	**Very good identification**					**07743020**
Actinomyces naeslundii		100.0	0.62	FRU 95		
07743021	**Very good identification**					**07743021**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07743022	**Very good identification**					**07743022**
					SPO	
Actinomyces naeslundii		97.15	0.62	FRU 95	1	
Clostridium ramosum		1.21	0.05		92	
Lactobacillus acidophilus		1.08	0.05		1	
07743023	**Very good identification**					**07743023**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07743030	**Good identification**					**07743030**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07743031	**Good identification**					**07743031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	FRU 95		
07743032	**Good identification**					**07743032**
Actinomyces naeslundii		100.0	0.41	FRU 95		
07743033	**Good identification**					**07743033**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07743040	**Good identification**					**07743040**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07743041	**Good identification**					**07743041**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07743042	**Good identification**					**07743042**
Actinomyces naeslundii		96.37	0.50	FRU 95		
Actinomyces israelii		3.63	0.16			
07743043	**Good identification**					**07743043**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07743060	**Good identification**					**07743060**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07743061	**Good identification**					**07743061**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07743062	**Good identification**					**07743062**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07743063	**Good identification**					**07743063**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07743100	**Very good identification**					**07743100**
Actinomyces naeslundii		98.21	0.70	FRU 95		
Actinomyces israelii		1.57	0.30			
Bifidobacterium breve		0.22	0.10			
07743101	**Very good identification**					**07743101**
Actinomyces naeslundii		100.0	0.73	FRU 95		
07743102	**Low discrimination**					**07743102**
Actinomyces naeslundii		89.75	0.70	FRU 95		
Lactobacillus acidophilus		4.03	0.23			
Actinomyces israelii		2.15	0.33			
07743103	**Very good identification**					**07743103**
Actinomyces naeslundii		100.0	0.73	FRU 95		
07743110	**Acceptable identification**					**07743110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.94	0.48	FRU 95		
Actinomyces israelii		4.18	0.16	MLZ 5		
Bifidobacterium breve		3.88	0.10			
07743111	**Good identification**					**07743111**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07743112	**Low discrimination**					**07743112**
Actinomyces naeslundii		53.05	0.48	FRU 95		
Bifidobacterium breve		22.66	0.27	MLZ 5		
Bifidobacterium adolescentis		20.68	0.26			
07743113	**Good identification**					**07743113**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07743120	**Low discrimination**					**07743120**
					SPO	
Actinomyces naeslundii		89.46	0.70	FRU 95	1	
Clostridium ramosum		9.72	0.28		92	
Bifidobacterium breve		0.35	0.15		1	
07743121	**Very good identification**					**07743121**
Actinomyces naeslundii		100.0	0.73	FRU 95		
07743122	**Low discrimination**					**07743122**
					SPO	
Actinomyces naeslundii		64.28	0.70	FRU 95	1	
Clostridium ramosum		27.94	0.39		92	
Lactobacillus acidophilus		2.88	0.23		1	
07743123	**Very good identification**					**07743123**
					SPO	
Actinomyces naeslundii		99.71	0.73	FRU 95	1	
Clostridium ramosum		0.29	0.05		92	
07743130	**Acceptable identification**					**07743130**
Actinomyces naeslundii		93.02	0.48	FRU 95		
Bifidobacterium breve		6.98	0.15	MLZ 5		
07743131	**Good identification**					**07743131**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07743132	**Low discrimination**					**07743132**
					SPO	
Actinomyces naeslundii		52.33	0.48	FRU 95	1	
Bifidobacterium breve		39.73	0.32	MLZ 5	1	
Clostridium ramosum		4.37	0.05		92	
07743133	**Good identification**					**07743133**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07743140	**Very good identification**					**07743140**
Actinomyces naeslundii		95.43	0.57	FRU 95		
Actinomyces israelii		3.37	0.23			
Bifidobacterium breve		1.20	0.10			
07743141	**Very good identification**					**07743141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.60	FRU 95		
07743142	**Low discrimination**					**07743142**
Actinomyces naeslundii		81.74	0.57	FRU 95		
Bifidobacterium breve		10.41	0.27			
Actinomyces israelii		4.32	0.26			
07743143	**Very good identification**					**07743143**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07743150	**Low discrimination**					**07743150**
Actinomyces naeslundii		80.68	0.35	FRU 95		
Bifidobacterium breve		19.32	0.10	MLZ 5		
07743151	**Good identification**					**07743151**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07743153	**Good identification**					**07743153**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07743160	**Very good identification**					**07743160**
Actinomyces naeslundii		97.81	0.57	FRU 95		
Bifidobacterium breve		2.19	0.15			
07743161	**Very good identification**					**07743161**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07743162	**Low discrimination**					**07743162**
Actinomyces naeslundii		77.96	0.57	FRU 95		
Bifidobacterium breve		17.65	0.32			
Lactobacillus acidophilus		2.45	0.07			
07743163	**Very good identification**					**07743163**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07743170	**Low discrimination**					**07743170**
Actinomyces naeslundii		70.15	0.35	FRU 95		
Bifidobacterium breve		29.85	0.15	MLZ 5		
07743171	**Good identification**					**07743171**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07743173	**Good identification**					**07743173**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07743300	**Good identification**					**07743300**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07743301	**Good identification**					**07743301**
Actinomyces naeslundii		100.0	0.39	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07743302	**Low discrimination**						**07743302**
					SPO		
Actinomyces naeslundii		47.87	0.36	FRU 95	1		
Clostridium ramosum		27.58	0.07	RHA 1	92		
Lactobacillus acidophilus		16.00	0.03		1		
07743303	**Good identification**						**07743303**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07743321	**Good identification**						**07743321**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07743322	**Good identification**						**07743322**
					SPO	GRA	
Clostridium ramosum		95.52	0.41	FRU 99	92	50	
Prevotella oralis		2.24	0.13	TRE 94	1	1	
Actinomyces naeslundii		1.68	0.36		1	99	
07743323	**Low discrimination**						**07743323**
					SPO		
Actinomyces naeslundii		72.25	0.39	FRU 95	1		
Clostridium ramosum		27.75	0.07	RHA 1	92		
07743501	**Good identification**						**07743501**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07743502	**Low discrimination**						**07743502**
Bifidobacterium adolescentis		79.07	0.38	FRU 99			
Actinomyces israelii		13.82	0.32	XYL 87			
Actinomyces naeslundii		7.11	0.36				
07743503	**Good identification**						**07743503**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07743512	**Good identification**						**07743512**
Bifidobacterium adolescentis		97.91	0.40	FRU 99			
Actinomyces israelii		2.09	0.18	XYL 87			
07743520	**Good identification**						**07743520**
Actinomyces naeslundii		100.0	0.36	FRU 95			
07743521	**Good identification**						**07743521**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07743522	**Low discrimination**						**07743522**
					SPO		
Actinomyces naeslundii		69.70	0.36	FRU 95	1		
Clostridium ramosum		30.30	0.05	ARA 1	92		
07743523	**Good identification**						**07743523**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07745100	**Low discrimination**						**07745100**
					GRA	SPO	
Actinomyces israelii		43.42	0.36	FRU 99	99	1	
Clostridium butyricum		42.28	0.14		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		7.34	0.20		1 1	
07745102	**Low discrimination**					**07745102**
					GRA	
Bifidobacterium adolescentis		59.06	0.40	FRU 99	99	
Actinomyces israelii		20.96	0.39	ARA 87	99	
Prevotella oralis		8.75	0.16		1	
07745112	**Acceptable identification**					**07745112**
Bifidobacterium adolescentis		94.53	0.41	FRU 99		
Actinomyces israelii		4.10	0.24	ARA 87		
Clostridium clostridioforme		1.37	0.07			
07745122	**Low discrimination**					**07745122**
					GRA	
Prevotella oralis		83.57	0.37	FRU 99	1	
Parabacteroides distasonis		11.70	0.36	XYL 11	1	
Lactobacillus acidophilus		1.86	0.10		99	
07745332	**Low discrimination**					**07745332**
Parabacteroides distasonis		81.76	0.39	FRU 99		
Clostridium clostridioforme		16.68	0.22			
Prevotella oralis		1.56	-.03			
07745502	**Low discrimination**					**07745502**
					SPO	
Bifidobacterium adolescentis		85.26	0.54	FRU 99	1	
Clostridium butyricum		10.27	0.24		99	
Actinomyces israelii		3.70	0.37		1	
07745512	**Good identification**					**07745512**
Bifidobacterium adolescentis		95.75	0.56	FRU 99		
Clostridium clostridioforme		3.25	0.27			
Actinomyces israelii		0.51	0.23			
07745542	**Low discrimination**					**07745542**
					SPO	
Bifidobacterium adolescentis		81.88	0.39	FRU 99	1	
Actinomyces israelii		11.18	0.30	SOR 11	1	
Clostridium butyricum		6.94	0.05		99	
07745552	**Good identification**					**07745552**
Bifidobacterium adolescentis		98.36	0.40	FRU 99		
Actinomyces israelii		1.64	0.16	SOR 11		
07745712	**Good identification**					**07745712**
Clostridium clostridioforme		96.10	0.42	FRU 99		
Bifidobacterium adolescentis		3.90	0.21	TRE 88		
07745722	**Low discrimination**					**07745722**
Parabacteroides distasonis		54.91	0.40	FRU 99		
Clostridium clostridioforme		33.84	0.31			
Bacteroides ovatus		4.57	0.06			
07745732	**Low discrimination**					**07745732**
Clostridium clostridioforme		82.54	0.42	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		15.83	0.35	TRE 88		
Bacteroides ovatus		1.63	0.03			
07747000	**Acceptable identification**					**07747000**
Actinomyces israelii		92.84	0.38	FRU 99		
Actinomyces naeslundii		7.16	0.28			
07747002	**Good identification**					**07747002**
Actinomyces israelii		95.11	0.41	FRU 99		
Actinomyces naeslundii		4.89	0.28			
07747100	**Acceptable identification**					**07747100**
					SPO	
Actinomyces israelii		93.11	0.48	FRU 99	1	
Actinomyces naeslundii		5.11	0.36		1	
Clostridium butyricum		1.78	-.02		99	
07747101	**Low discrimination**					**07747101**
Actinomyces naeslundii		89.07	0.39	FRU 95		
Actinomyces israelii		10.93	0.14	XYL 1		
07747102	**Low discrimination**					**07747102**
Actinomyces israelii		69.28	0.51	FRU 99		
Bifidobacterium adolescentis		28.19	0.38			
Actinomyces naeslundii		2.54	0.36			
07747103	**Low discrimination**					**07747103**
Actinomyces naeslundii		84.46	0.39	FRU 95		
Actinomyces israelii		15.54	0.17	XYL 1		
07747112	**Low discrimination**					**07747112**
					SPO	
Bifidobacterium adolescentis		76.19	0.40	FRU 99	1	
Actinomyces israelii		22.89	0.37	ARA 87	1	
Clostridium tertium		0.92	-.01		99	
07747121	**Good identification**					**07747121**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07747123	**Good identification**					**07747123**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07747140	**Good identification**					**07747140**
Actinomyces israelii		97.57	0.41	FRU 99		
Actinomyces naeslundii		2.43	0.23			
07747142	**Low discrimination**					**07747142**
Actinomyces israelii		87.27	0.44	FRU 99		
Bifidobacterium adolescentis		11.29	0.23			
Actinomyces naeslundii		1.45	0.23			
07747400	**Good identification**					**07747400**
Actinomyces israelii		100.0	0.37	FRU 99		
07747402	**Low discrimination**					**07747402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		88.15	0.40	FRU 99		
Bifidobacterium adolescentis		11.85	0.18			
07747500	**Low discrimination**					**07747500**
					SPO	
Actinomyces israelii		79.34	0.47	FRU 99	1	
Clostridium butyricum		16.66	0.14		99	
Bifidobacterium adolescentis		4.00	0.18		1	
07747502	**Low discrimination**					**07747502**
					SPO	
Bifidobacterium adolescentis		75.94	0.53	FRU 99	1	
Actinomyces israelii		22.82	0.50		1	
Clostridium butyricum		1.24	0.07		99	
07747512	**Good identification**					**07747512**
Bifidobacterium adolescentis		96.16	0.54	FRU 99		
Actinomyces israelii		3.53	0.36			
Bifidobacterium dentium		0.30	-.07			
07747540	**Good identification**					**07747540**
					SPO	
Actinomyces israelii		95.52	0.40	FRU 99	1	
Clostridium butyricum		4.49	-.04		99	
07747542	**Low discrimination**					**07747542**
Bifidobacterium adolescentis		51.40	0.37	FRU 99		
Actinomyces israelii		48.60	0.43	SOR 11		
07747552	**Low discrimination**					**07747552**
Bifidobacterium adolescentis		89.64	0.39	FRU 99		
Actinomyces israelii		10.36	0.29	SOR 11		
07750000	**Low discrimination**					**07750000**
					GRA	
Fusobacterium mortiferum		88.19	0.64	MLT 11	1	
Actinomyces naeslundii		6.13	0.56		99	
Lactobacillus catenaformis		2.91	0.29		99	
07750001	**Good identification**					**07750001**
					GRA	
Actinomyces naeslundii		90.42	0.59	GAL 96	99	
Fusobacterium mortiferum		8.77	0.30		1	
Actinomyces viscosus		0.81	-.01		99	
07750002	**Low discrimination**					**07750002**
					GRA	
Actinomyces viscosus		50.30	0.42	NIT 99	99	
Fusobacterium mortiferum		20.14	0.48		1	
Actinomyces naeslundii		12.59	0.56		99	
07750003	**Low discrimination**					**07750003**
					GRA	
Actinomyces naeslundii		52.67	0.59	GAL 96	99	
Actinomyces viscosus		46.76	0.34		99	
Fusobacterium mortiferum		0.57	0.14		1	
07750011	**Good identification**					**07750011**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	GAL 96			
07750013	**Low discrimination**						**07750013**
Actinomyces naeslundii		85.44	0.37	GAL 96			
Actinomyces viscosus		14.56	-.01	MLZ 5			
07750020	**Low discrimination**						**07750020**
					GRA	SPO	
Actinomyces naeslundii		65.26	0.56	GAL 96	99	1	
Fusobacterium mortiferum		9.49	0.30		1	1	
Clostridium barati		8.11	0.14		99	99	
07750021	**Very good identification**						**07750021**
Actinomyces naeslundii		100.0	0.59	GAL 96			
07750023	**Very good identification**						**07750023**
					GRA		
Actinomyces naeslundii		98.20	0.59	GAL 96	99		
Prevotella oralis		0.92	-.02		1		
Actinomyces viscosus		0.88	-.01		99		
07750031	**Good identification**						**07750031**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07750033	**Good identification**						**07750033**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07750040	**Low discrimination**						**07750040**
					GRA		
Actinomyces naeslundii		54.82	0.43	GAL 96	99		
Fusobacterium mortiferum		45.18	0.30		1		
07750041	**Good identification**						**07750041**
Actinomyces naeslundii		100.0	0.46	GAL 96			
07750042	**Low discrimination**						**07750042**
Actinomyces naeslundii		59.44	0.43	GAL 96			
Lactobacillus acidophilus		21.52	0.11				
Actinomyces viscosus		13.59	0.08				
07750043	**Good identification**						**07750043**
Actinomyces naeslundii		95.16	0.46	GAL 96			
Actinomyces viscosus		4.84	-.01				
07750060	**Good identification**						**07750060**
Actinomyces naeslundii		100.0	0.43	GAL 96			
07750061	**Good identification**						**07750061**
Actinomyces naeslundii		100.0	0.46	GAL 96			
07750062	**Low discrimination**						**07750062**
					SPO		
Actinomyces naeslundii		61.52	0.43	GAL 96	1		
Lactobacillus acidophilus		22.28	0.11		1		
Clostridium barati		11.30	0.04		99		
07750063	**Good identification**						**07750063**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.46	GAL 96			
07750100	**Low discrimination**						**07750100**
					GRA		
Fusobacterium mortiferum		89.45	0.70	MLT 11	1		
Actinomyces naeslundii		7.58	0.64		99		
Lactobacillus catenaformis		1.89	0.32		99		
07750101	**Good identification**						**07750101**
					GRA		
Actinomyces naeslundii		92.16	0.67	GAL 96	99		
Fusobacterium mortiferum		7.33	0.36		1		
Actinomyces viscosus		0.52	0.04		99		
07750102	**Low discrimination**						**07750102**
					GRA		
Actinomyces viscosus		31.98	0.46	NIT 99	99		
Lactobacillus acidophilus		26.77	0.45		99		
Fusobacterium mortiferum		16.81	0.54		1		
07750103	**Low discrimination**						**07750103**
Actinomyces naeslundii		63.28	0.67	GAL 96			
Actinomyces viscosus		35.10	0.38				
Lactobacillus acidophilus		0.89	0.10				
07750110	**Low discrimination**						**07750110**
					GRA		
Fusobacterium mortiferum		67.24	0.36	MLT 11	1		
Actinomyces naeslundii		29.68	0.42	MLZ 1	99		
Bifidobacterium breve		3.08	0.10		99		
07750111	**Good identification**						**07750111**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07750113	**Acceptable identification**						**07750113**
Actinomyces naeslundii		90.38	0.45	GAL 96			
Actinomyces viscosus		9.62	0.04	MLZ 5			
07750120	**Low discrimination**						**07750120**
					GRA	SPO	
Actinomyces naeslundii		50.56	0.64	GAL 96	99	1	
Prevotella oralis		20.39	0.31		1	1	
Clostridium barati		6.62	0.22		99	99	
07750121	**Very good identification**						**07750121**
					GRA		
Actinomyces naeslundii		99.73	0.67	GAL 96	99		
Prevotella oralis		0.27	-.03		1		
07750122	**Low discrimination**						**07750122**
					GRA		
Prevotella oralis		65.95	0.53	GAL 99	1		
Lactobacillus acidophilus		17.98	0.45		99		
Actinomyces naeslundii		8.61	0.64		99		
07750123	**Good identification**						**07750123**
					GRA		
Actinomyces naeslundii		93.12	0.67	GAL 96	99		
Prevotella oralis		4.81	0.19		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		1.31	0.10		99		
07750130	**Low discrimination**						**07750130**
					GRA		
Actinomyces naeslundii		79.27	0.42	GAL 96	99		
Bifidobacterium breve		14.59	0.15	MLZ 5	99		
Prevotella oralis		6.14	-.03		1		
07750131	**Good identification**						**07750131**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07750133	**Good identification**						**07750133**
Actinomyces naeslundii		100.0	0.45	GAL 96			
07750140	**Low discrimination**						**07750140**
					GRA		
Actinomyces naeslundii		58.60	0.51	GAL 96	99		
Fusobacterium mortiferum		39.59	0.36		1		
Bifidobacterium breve		1.81	0.10		99		
07750141	**Good identification**						**07750141**
Actinomyces naeslundii		100.0	0.54	GAL 96			
07750143	**Good identification**						**07750143**
Actinomyces naeslundii		96.92	0.54	GAL 96			
Actinomyces viscosus		3.08	0.04				
07750160	**Low discrimination**						**07750160**
					SPO		
Actinomyces naeslundii		82.85	0.51	GAL 96	1		
Clostridium barati		5.35	0.04		99		
Bifidobacterium breve		4.55	0.15		1		
07750161	**Good identification**						**07750161**
Actinomyces naeslundii		100.0	0.54	GAL 96			
07750163	**Good identification**						**07750163**
Actinomyces naeslundii		100.0	0.54	GAL 96			
07750300	**Low discrimination**						**07750300**
					GRA		
Fusobacterium mortiferum		86.13	0.36	MLT 11	1		
Actinomyces naeslundii		7.30	0.29	RHA 1	99		
Prevotella oralis		6.57	0.03		1		
07750322	**Low discrimination**						**07750322**
					SPO	GRA	
Prevotella oralis		86.19	0.47	GAL 99	1	1	
Clostridium ramosum		6.54	0.28		92	50	
Lactobacillus acidophilus		4.13	0.25		1	99	
07750500	**Low discrimination**						**07750500**
					GRA	SPO	
Fusobacterium mortiferum		76.15	0.36	MLT 11	1	1	
Clostridium butyricum		13.78	0.05	ARA 1	99	99	
Actinomyces naeslundii		6.45	0.29		99	1	
07750502	**Low discrimination**						**07750502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		68.12	0.40	GAL 99		
Clostridium clostridioforme		13.54	0.25	XYL 87		
Actinomyces viscosus		5.10	0.12			
07750512	**Low discrimination**					**07750512**
Bifidobacterium adolescentis		57.45	0.41	GAL 99		
Clostridium clostridioforme		42.55	0.36	XYL 87		
07750532	**Good identification**					**07750532**
					GRA	
Clostridium clostridioforme		97.13	0.36	TRE 88	50	
Parabacteroides distasonis		1.55	0.11	XYL 91	1	
Bifidobacterium adolescentis		1.33	0.07	RHA 88	99	
07750612	**Good identification**					**07750612**
Clostridium clostridioforme		100.0	0.42	TRE 88		
07750632	**Good identification**					**07750632**
Clostridium clostridioforme		100.0	0.42	TRE 88		
07750702	**Good identification**					**07750702**
Clostridium clostridioforme		100.0	0.40	TRE 88		
07750712	**Good identification**					**07750712**
Clostridium clostridioforme		99.81	0.51	TRE 88		
Bifidobacterium adolescentis		0.19	0.07	XYL 91		
07750722	**Low discrimination**					**07750722**
Clostridium clostridioforme		89.06	0.40	TRE 88		
Prevotella oralis		6.02	0.13	XYL 91		
Parabacteroides distasonis		4.92	0.24			
07750732	**Good identification**					**07750732**
Clostridium clostridioforme		99.35	0.51	TRE 88		
Parabacteroides distasonis		0.65	0.19	XYL 91		
07751000	**Low discrimination**					**07751000**
Lactobacillus catenaformis		39.58	0.63	RAF 1		
Actinomyces naeslundii		20.23	0.79			
Lactobacillus jensenii		19.87	0.58			
07751001	**Very good identification**					**07751001**
Actinomyces naeslundii		97.48	0.82			
Lactobacillus catenaformis		1.28	0.29			
Lactobacillus jensenii		0.65	0.24			
07751002	**Low discrimination**					**07751002**
Lactobacillus acidophilus		33.37	0.61	LAC 92		
Lactobacillus catenaformis		30.61	0.63			
Actinomyces naeslundii		15.65	0.79			
07751003	**Good identification**					**07751003**
Actinomyces naeslundii		94.52	0.82			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		2.14	0.30			
Lactobacillus acidophilus		1.36	0.26			
07751010	**Low discrimination**					**07751010**
Lactobacillus jensenii		79.36	0.50	RAF 1		
Actinomyces naeslundii		12.76	0.58			
Lactobacillus catenaformis		4.79	0.29			
07751011	**Very good identification**					**07751011**
Actinomyces naeslundii		95.98	0.61	MLZ 5		
Lactobacillus jensenii		4.02	0.15			
07751012	**Low discrimination**					**07751012**
Lactobacillus jensenii		64.33	0.47	RAF 1		
Actinomyces naeslundii		15.51	0.58			
Lactobacillus acidophilus		6.35	0.26			
07751013	**Very good identification**					**07751013**
Actinomyces naeslundii		97.28	0.61	MLZ 5		
Lactobacillus jensenii		2.72	0.12			
07751020	**Low discrimination**					**07751020**
					GRA	
Actinomyces naeslundii		44.66	0.79		99	
Lactobacillus catenaformis		21.84	0.53		99	
Prevotella oralis		13.50	0.45		1	
07751021	**Very good identification**					**07751021**
					GRA	
Actinomyces naeslundii		99.21	0.82		99	
Lactobacillus catenaformis		0.33	0.18		99	
Prevotella oralis		0.20	0.11		1	
07751022	**Low discrimination**					**07751022**
					GRA	
Prevotella oralis		58.06	0.67	LAC 94	1	
Lactobacillus acidophilus		21.56	0.61		99	
Actinomyces naeslundii		10.11	0.79		99	
07751023	**Good identification**					**07751023**
					GRA	
Actinomyces naeslundii		94.39	0.82		99	
Prevotella oralis		3.65	0.33		1	
Lactobacillus acidophilus		1.36	0.26		99	
07751030	**Low discrimination**					**07751030**
Lactobacillus jensenii		56.41	0.39	RAF 1		
Actinomyces naeslundii		36.28	0.58			
Lactobacillus catenaformis		3.41	0.18			
07751031	**Very good identification**					**07751031**
Actinomyces naeslundii		98.96	0.61	MLZ 5		
Lactobacillus jensenii		1.04	0.05			
07751033	**Very good identification**					**07751033**
					GRA	
Actinomyces naeslundii		98.58	0.61	MLZ 5	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		0.73	-.02		1	
Lactobacillus jensenii		0.69	0.02		99	
07751040	**Low discrimination**					**07751040**
Actinomyces naeslundii		76.96	0.67			
Lactobacillus catenaformis		8.62	0.29			
Lactobacillus jensenii		4.33	0.24			
07751041	**Very good identification**					**07751041**
Actinomyces naeslundii		100.0	0.70			
07751042	**Low discrimination**					**07751042**
Lactobacillus acidophilus		52.80	0.45	LAC 92		
Actinomyces naeslundii		35.35	0.67	SOR 11		
Lactobacillus catenaformis		3.96	0.29			
07751043	**Very good identification**					**07751043**
Actinomyces naeslundii		99.00	0.70			
Lactobacillus acidophilus		1.00	0.11			
07751050	**Low discrimination**					**07751050**
Actinomyces naeslundii		64.40	0.45	MLZ 5		
Lactobacillus jensenii		22.93	0.15			
Actinomyces israelii		8.19	0.20			
07751051	**Good identification**					**07751051**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07751052	**Low discrimination**					**07751052**
Actinomyces naeslundii		39.75	0.45	MLZ 5		
Bifidobacterium breve		27.98	0.27			
Lactobacillus acidophilus		11.40	0.11			
07751053	**Good identification**					**07751053**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07751060	**Low discrimination**					**07751060**
					SPO	
Actinomyces naeslundii		84.25	0.67		1	
Clostridium perfringens		4.28	0.22		84	
Clostridium barati		4.08	0.17		99	
07751061	**Very good identification**					**07751061**
Actinomyces naeslundii		100.0	0.70			
07751062	**Low discrimination**					**07751062**
					GRA	
Lactobacillus acidophilus		48.20	0.45	LAC 92	99	
Actinomyces naeslundii		32.27	0.67	SOR 11	99	
Prevotella oralis		10.61	0.33		1	
07751063	**Very good identification**					**07751063**
					GRA	
Actinomyces naeslundii		98.79	0.70		99	
Lactobacillus acidophilus		0.99	0.11		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		0.22	-.02		1	
07751070	**Low discrimination**					**07751070**
Actinomyces naeslundii		82.46	0.45	MLZ 5		
Bifidobacterium breve		10.20	0.15			
Lactobacillus jensenii		7.34	0.05			
07751071	**Good identification**					**07751071**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07751073	**Good identification**					**07751073**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
07751100	**Low discrimination**					**07751100**
					GRA	
Lactobacillus catenaformis		32.86	0.66	RAF 1	99	
Actinomyces naeslundii		31.87	0.87		99	
Fusobacterium mortiferum		23.51	0.73		1	
07751101	**Very good identification**					**07751101**
					GRA	
Actinomyces naeslundii		98.59	0.90		99	
Lactobacillus catenaformis		0.68	0.32		99	
Fusobacterium mortiferum		0.49	0.39		1	
07751102	**Low discrimination**					**07751102**
					GRA	
Lactobacillus acidophilus		65.86	0.79		99	
Prevotella oralis		12.71	0.65		1	
Lactobacillus catenaformis		7.88	0.66		99	
07751103	**Good identification**					**07751103**
Actinomyces naeslundii		91.33	0.90			
Lactobacillus acidophilus		5.30	0.45			
Actinomyces viscosus		1.29	0.34			
07751110	**Low discrimination**					**07751110**
Actinomyces naeslundii		50.33	0.65	MLZ 5		
Bifidobacterium breve		21.50	0.44			
Lactobacillus catenaformis		9.96	0.32			
07751111	**Very good identification**					**07751111**
Actinomyces naeslundii		99.71	0.68	MLZ 5		
Bifidobacterium breve		0.29	0.10			
07751112	**Low discrimination**					**07751112**
Bifidobacterium adolescentis		37.30	0.61	XYL 87		
Bifidobacterium breve		33.45	0.62	ARA 87		
Lactobacillus acidophilus		12.81	0.45			
07751113	**Good identification**					**07751113**
Actinomyces naeslundii		92.59	0.68	MLZ 5		
Bifidobacterium adolescentis		3.00	0.27			
Bifidobacterium breve		2.69	0.27			
07751120	**Low discrimination**					**07751120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Prevotella oralis		40.24	0.65	BGL 95	1	
Actinomyces naeslundii		24.19	0.87		99	
Prevotella melaninogenica		7.79	0.50		1	
07751121	**Very good identification**					**07751121**
					GRA	
Actinomyces naeslundii		97.95	0.90		99	
Prevotella oralis		1.10	0.31		1	
Prevotella melaninogenica		0.21	0.15		1	
07751122	**Low discrimination**					**07751122**
					GRA	
Prevotella oralis		72.95	0.87		1	
Lactobacillus acidophilus		19.89	0.79		99	
Actinomyces naeslundii		2.31	0.87		99	
07751123	**Low discrimination**					**07751123**
					GRA	
Actinomyces naeslundii		77.81	0.90		99	
Prevotella oralis		16.56	0.53		1	
Lactobacillus acidophilus		4.52	0.45		99	
07751130	**Low discrimination**					**07751130**
					GRA	
Actinomyces naeslundii		42.29	0.65	MLZ 5	99	
Bifidobacterium breve		32.12	0.49		99	
Prevotella oralis		13.50	0.31		1	
07751131	**Very good identification**					**07751131**
					GRA	
Actinomyces naeslundii		99.28	0.68	MLZ 5	99	
Bifidobacterium breve		0.51	0.15		99	
Prevotella oralis		0.21	-.03		1	
07751132	**Low discrimination**					**07751132**
					GRA	
Bifidobacterium breve		43.90	0.66	ESL 99	99	
Prevotella oralis		34.68	0.53		1	
Lactobacillus acidophilus		9.45	0.45		99	
07751133	**Good identification**					**07751133**
					GRA	
Actinomyces naeslundii		90.26	0.68	MLZ 5	99	
Bifidobacterium breve		4.67	0.32		99	
Prevotella oralis		3.69	0.19		1	
07751140	**Low discrimination**					**07751140**
Actinomyces naeslundii		71.91	0.74			
Bifidobacterium breve		9.16	0.44			
Actinomyces israelii		4.53	0.45			
07751141	**Very good identification**					**07751141**
Actinomyces naeslundii		99.91	0.77			
Bifidobacterium breve		0.09	0.10			
07751142	**Low discrimination**					**07751142**
Lactobacillus acidophilus		68.73	0.63	SOR 11		
Bifidobacterium breve		14.67	0.62			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		11.39	0.74			
07751143	**Very good identification**					**07751143**
Actinomyces naeslundii		95.02	0.77			
Lactobacillus acidophilus		3.86	0.29			
Bifidobacterium breve		0.82	0.27			
07751150	**Low discrimination**					**07751150**
Bifidobacterium breve		66.62	0.44	ESL 99		
Actinomyces naeslundii		27.52	0.52	BGL 91		
Actinomyces israelii		4.92	0.30			
07751151	**Good identification**					**07751151**
Actinomyces naeslundii		98.40	0.55	MLZ 5		
Bifidobacterium breve		1.60	0.10			
07751152	**Low discrimination**					**07751152**
Bifidobacterium breve		80.74	0.62	ESL 99		
Bifidobacterium adolescentis		11.13	0.46			
Lactobacillus acidophilus		3.82	0.29			
07751153	**Low discrimination**					**07751153**
Actinomyces naeslundii		84.21	0.55	MLZ 5		
Bifidobacterium breve		13.88	0.27			
Bifidobacterium adolescentis		1.91	0.12			
07751160	**Low discrimination**					**07751160**
					SPO	
Actinomyces naeslundii		57.21	0.74		1	
Bifidobacterium breve		12.96	0.49		1	
Clostridium paraputrificum		7.72	0.27		99	
07751161	**Very good identification**					**07751161**
Actinomyces naeslundii		99.61	0.77			
Bifidobacterium breve		0.15	0.15			
Clostridium perfringens		0.09	0.03			
07751162	**Low discrimination**					**07751162**
					GRA	
Lactobacillus acidophilus		52.55	0.63	SOR 11	99	
Bifidobacterium breve		19.94	0.66		99	
Prevotella oralis		15.75	0.53		1	
07751163	**Good identification**					**07751163**
Actinomyces naeslundii		93.52	0.77			
Lactobacillus acidophilus		3.80	0.29			
Bifidobacterium breve		1.44	0.32			
07751170	**Low discrimination**					**07751170**
					SPO	
Bifidobacterium breve		80.10	0.49	ESL 99	1	
Actinomyces naeslundii		18.61	0.52	BGL 91	1	
Clostridium paraputrificum		0.48	-.07		99	
07751171	**Good identification**					**07751171**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		97.18	0.55	MLZ 5			
Bifidobacterium breve		2.82	0.15				
07751172	**Good identification**						**07751172**
Bifidobacterium breve		94.35	0.66	ESL 99			
Lactobacillus acidophilus		2.51	0.29				
Actinomyces naeslundii		2.17	0.52				
07751173	**Low discrimination**						**07751173**
Actinomyces naeslundii		77.34	0.55	MLZ 5			
Bifidobacterium breve		22.66	0.32				
07751201	**Good identification**						**07751201**
Actinomyces naeslundii		100.0	0.48	RHA 1			
07751202	**Low discrimination**						**07751202**
					GRA		
Lactobacillus acidophilus		46.84	0.41	LAC 92	99		
Prevotella oralis		37.79	0.39	RHA 7	1		
Lactobacillus catenaformis		5.77	0.29		99		
07751203	**Low discrimination**						**07751203**
					GRA		
Actinomyces naeslundii		83.80	0.48	RHA 1	99		
Lactobacillus acidophilus		8.97	0.07		99		
Prevotella oralis		7.24	0.04		1		
07751220	**Low discrimination**						**07751220**
					GRA	SPO	
Prevotella oralis		78.46	0.39	LAC 94	1	1	
Actinomyces naeslundii		6.12	0.45	BGL 95	99	1	
Clostridium barati		5.18	0.17		99	99	
07751221	**Acceptable identification**						**07751221**
					GRA		
Actinomyces naeslundii		92.05	0.48	RHA 1	99		
Prevotella oralis		7.95	0.04		1		
07751222	**Good identification**						**07751222**
					SPO	GRA	
Prevotella oralis		91.08	0.60	LAC 94	1	1	
Lactobacillus acidophilus		5.94	0.41		1	99	
Clostridium ramosum		1.09	0.28		92	50	
07751241	**Good identification**						**07751241**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07751243	**Good identification**						**07751243**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07751261	**Good identification**						**07751261**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07751263	**Good identification**						**07751263**
Actinomyces naeslundii		100.0	0.35	RHA 1			
07751300	**Low discrimination**						**07751300**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		46.27	0.37	NAG 95	1		
Lactobacillus catenaformis		12.84	0.32	BGL 95	99		
Actinomyces naeslundii		12.46	0.53		99		
07751301	**Good identification**						**07751301**
					GRA		
Actinomyces naeslundii		97.56	0.56	RHA 1	99		
Prevotella oralis		2.44	0.03		1		
07751302	**Low discrimination**						**07751302**
					GRA		
Prevotella oralis		50.33	0.59	NAG 95	1		
Lactobacillus acidophilus		45.79	0.60		99		
Lactobacillus catenaformis		0.74	0.32		99		
07751303	**Low discrimination**						**07751303**
					GRA		
Actinomyces naeslundii		52.03	0.56	RHA 1	99		
Prevotella oralis		24.73	0.25		1		
Lactobacillus acidophilus		22.50	0.25		99		
07751312	**Low discrimination**						**07751312**
					GRA		
Clostridium clostridioforme		46.14	0.39	TRE 88	50		
Prevotella oralis		18.64	0.25	XYL 91	1		
Lactobacillus acidophilus		16.96	0.25	ARA 94	99		
07751320	**Low discrimination**						**07751320**
					SPO	GRA	
Prevotella oralis		70.48	0.59	BGL 95	1	1	
Clostridium ramosum		25.42	0.52		92	50	
Actinomyces naeslundii		1.00	0.53		1	99	
07751321	**Low discrimination**						**07751321**
					SPO	GRA	
Actinomyces naeslundii		60.73	0.56	RHA 1	1	99	
Prevotella oralis		28.86	0.25		1	1	
Clostridium ramosum		10.41	0.18		92	50	
07751322	**Low discrimination**						**07751322**
					SPO	GRA	
Prevotella oralis		87.64	0.81		1	1	
Clostridium ramosum		6.65	0.62		92	50	
Lactobacillus acidophilus		4.20	0.60		1	99	
07751323	**Low discrimination**						**07751323**
					SPO	GRA	
Prevotella oralis		80.00	0.47	URE 1	1	1	
Actinomyces naeslundii		8.86	0.56		1	99	
Clostridium ramosum		6.07	0.28		92	50	
07751332	**Low discrimination**						**07751332**
Prevotella oralis		49.82	0.47	MLZ 1			
Parabacteroides distasonis		35.75	0.57				
Clostridium clostridioforme		6.49	0.39				
07751340	**Low discrimination**						**07751340**
Actinomyces naeslundii		41.50	0.40	RHA 1			
Actinomyces israelii		19.46	0.25				
Lactobacillus acidophilus		18.85	0.10				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07751341	**Good identification**						**07751341**
Actinomyces naeslundii		100.0	0.43	RHA 1			
07751342	**Low discrimination**						**07751342**
					GRA		
Lactobacillus acidophilus		85.52	0.44	RHA 7	99		
Prevotella oralis		7.68	0.25	SOR 11	1		
Bifidobacterium breve		2.45	0.27		99		
07751343	**Low discrimination**						**07751343**
Actinomyces naeslundii		76.76	0.43	RHA 1			
Lactobacillus acidophilus		23.25	0.10				
07751361	**Good identification**						**07751361**
Actinomyces naeslundii		100.0	0.43	RHA 1			
07751362	**Low discrimination**						**07751362**
					SPO	GRA	
Prevotella oralis		57.63	0.47	SOR 1	1	1	
Lactobacillus acidophilus		33.76	0.44		1	99	
Clostridium ramosum		4.38	0.28		92	50	
07751363	**Low discrimination**						**07751363**
					GRA		
Actinomyces naeslundii		54.95	0.43	RHA 1	99		
Prevotella oralis		28.41	0.13		1		
Lactobacillus acidophilus		16.64	0.10		99		
07751401	**Good identification**						**07751401**
Actinomyces naeslundii		100.0	0.48	ARA 1			
07751402	**Low discrimination**						**07751402**
Bifidobacterium adolescentis		28.21	0.40	XYL 87			
Actinomyces israelii		17.63	0.43	LAC 99			
Lactobacillus acidophilus		15.23	0.26				
07751403	**Good identification**						**07751403**
Actinomyces naeslundii		100.0	0.48	ARA 1			
07751412	**Low discrimination**						**07751412**
Bifidobacterium adolescentis		55.67	0.41	XYL 87			
Clostridium clostridioforme		36.95	0.35	LAC 99			
Actinomyces israelii		4.25	0.29				
07751420	**Low discrimination**						**07751420**
					GRA		
Actinomyces naeslundii		42.05	0.45	ARA 1	99		
Lactobacillus catenaformis		20.56	0.18		99		
Prevotella oralis		12.71	0.11		1		
07751421	**Good identification**						**07751421**
Actinomyces naeslundii		100.0	0.48	ARA 1			
07751423	**Good identification**						**07751423**
					GRA		
Actinomyces naeslundii		96.28	0.48	ARA 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		3.72	-.02		1	
07751432	**Acceptable identification**					**07751432**
Clostridium clostridioforme		92.20	0.35	TRE 88		
Prevotella oralis		1.67	-.02	XYL 91		
Parabacteroides distasonis		1.64	0.11	RHA 88		
07751441	**Good identification**					**07751441**
Actinomyces naeslundii		100.0	0.35	ARA 1		
07751442	**Low discrimination**					**07751442**
Actinomyces israelii		50.84	0.36	XYL 92		
Bifidobacterium adolescentis		25.86	0.24			
Lactobacillus acidophilus		13.96	0.11			
07751443	**Good identification**					**07751443**
Actinomyces naeslundii		100.0	0.35	ARA 1		
07751461	**Good identification**					**07751461**
Actinomyces naeslundii		100.0	0.35	ARA 1		
07751463	**Good identification**					**07751463**
Actinomyces naeslundii		100.0	0.35	ARA 1		
07751500	**Low discrimination**					**07751500**
					SPO	
Clostridium butyricum		56.16	0.39	ESL 88	99	
Actinomyces israelii		14.74	0.50	TRE 94	1	
Bifidobacterium adolescentis		8.85	0.40	XYL 97	1	
07751501	**Acceptable identification**					**07751501**
					SPO	
Actinomyces naeslundii		93.03	0.56	ARA 1	1	
Clostridium butyricum		5.52	0.05		99	
Actinomyces israelii		1.45	0.16		1	
07751502	**Low discrimination**					**07751502**
Bifidobacterium adolescentis		86.93	0.74	XYL 87		
Lactobacillus acidophilus		5.45	0.45			
Actinomyces israelii		2.20	0.53			
07751503	**Low discrimination**					**07751503**
Actinomyces naeslundii		48.65	0.56	ARA 1		
Bifidobacterium adolescentis		45.03	0.40			
Lactobacillus acidophilus		2.82	0.10			
07751510	**Low discrimination**					**07751510**
					SPO	
Bifidobacterium adolescentis		63.62	0.41	XYL 87	1	
Clostridium butyricum		17.39	0.17	BGL 99	99	
Actinomyces israelii		12.96	0.36		1	
07751512	**Very good identification**					**07751512**
Bifidobacterium adolescentis		97.14	0.76	XYL 87		
Clostridium clostridioforme		2.18	0.44			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	0.30	0.39			
07751513	**Acceptable identification**					**07751513**
	Bifidobacterium adolescentis	93.55	0.41	XYL 87		
	Actinomyces naeslundii	4.35	0.34	URE 1		
	Clostridium clostridioforme	2.10	0.10			
07751521	**Good identification**					**07751521**
					GRA	
	Actinomyces naeslundii	98.89	0.56	ARA 1	99	
	Prevotella oralis	1.11	-.03		1	
07751522	**Low discrimination**					**07751522**
					GRA	
	Prevotella oralis	63.31	0.53	ARA 1	1	
	Lactobacillus acidophilus	17.26	0.45		99	
	Parabacteroides distasonis	7.39	0.50		1	
07751523	**Low discrimination**					**07751523**
					GRA	
	Actinomyces naeslundii	77.18	0.56	ARA 1	99	
	Prevotella oralis	16.43	0.19		1	
	Lactobacillus acidophilus	4.48	0.10		99	
07751532	**Low discrimination**					**07751532**
					GRA	
	Clostridium clostridioforme	44.15	0.44	TRE 88	50	
	Parabacteroides distasonis	23.26	0.45	XYL 91	1	
	Bifidobacterium adolescentis	19.87	0.41	RHA 88	99	
07751540	**Low discrimination**					**07751540**
					SPO	
	Actinomyces israelii	44.17	0.43	XYL 92	1	
	Clostridium butyricum	37.63	0.21		99	
	Actinomyces naeslundii	8.67	0.40		1	
07751541	**Good identification**					**07751541**
	Actinomyces naeslundii	100.0	0.43	ARA 1		
07751542	**Low discrimination**					**07751542**
	Bifidobacterium adolescentis	84.43	0.59	XYL 87		
	Actinomyces israelii	6.71	0.46	SOR 11		
	Lactobacillus acidophilus	5.29	0.29			
07751543	**Low discrimination**					**07751543**
	Actinomyces naeslundii	60.67	0.43	ARA 1		
	Bifidobacterium adolescentis	39.33	0.24			
07751552	**Very good identification**					**07751552**
	Bifidobacterium adolescentis	97.80	0.60	XYL 87		
	Bifidobacterium breve	1.07	0.27	SOR 11		
	Actinomyces israelii	0.95	0.32			
07751560	**Low discrimination**					**07751560**
					SPO	
	Actinomyces naeslundii	43.66	0.40	ARA 1	1	
	Clostridium butyricum	16.49	0.03		99	
	Actinomyces israelii	14.21	0.23		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07751561	**Good identification**						**07751561**
Actinomyces naeslundii		100.0	0.43	ARA 1			
07751563	**Good identification**						**07751563**
Actinomyces naeslundii		100.0	0.43	ARA 1			
07751602	**Acceptable identification**						**07751602**
Clostridium clostridioforme		91.56	0.39	TRE 88			
Actinomyces israelii		3.30	0.24	XYL 91			
Lactobacillus acidophilus		2.85	0.07				
07751612	**Good identification**						**07751612**
Clostridium clostridioforme		99.79	0.50	TRE 88			
Bifidobacterium adolescentis		0.21	0.07	XYL 91			
07751622	**Low discrimination**						**07751622**
Clostridium clostridioforme		63.71	0.39	TRE 88			
Prevotella oralis		30.38	0.26	XYL 91			
Parabacteroides distasonis		3.93	0.24				
07751632	**Good identification**						**07751632**
Clostridium clostridioforme		99.28	0.50	TRE 88			
Parabacteroides distasonis		0.72	0.19	XYL 91			
07751702	**Low discrimination**						**07751702**
					GRA		
Clostridium clostridioforme		65.28	0.48	TRE 88	50		
Bifidobacterium adolescentis		14.93	0.40	XYL 91	99		
Prevotella oralis		7.67	0.25		1		
07751712	**Good identification**						**07751712**
					GRA		
Clostridium clostridioforme		93.90	0.59	TRE 88	50		
Bifidobacterium adolescentis		5.76	0.41	XYL 91	99		
Parabacteroides distasonis		0.20	0.19		1		
07751722	**Low discrimination**						**07751722**
Prevotella oralis		41.54	0.47	ARA 1			
Parabacteroides distasonis		33.93	0.58				
Clostridium clostridioforme		18.61	0.48				
07751732	**Low discrimination**						**07751732**
Clostridium clostridioforme		81.76	0.59	TRE 88			
Parabacteroides distasonis		17.62	0.53	XYL 91			
Prevotella oralis		0.41	0.13				
07752000	**Low discrimination**						**07752000**
					GRA	SPO	
Actinomyces naeslundii		88.21	0.61	GAL 96	99	1	
Fusobacterium mortiferum		6.61	0.30		1	1	
Clostridium innocuum		2.42	0.10		99	99	
07752001	**Very good identification**						**07752001**
Actinomyces naeslundii		100.0	0.64	GAL 96			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07752002	**Good identification**					**07752002**
					SPO	
Actinomyces naeslundii		91.57	0.61	GAL 96	1	
Clostridium innocuum		2.51	0.10		99	
Actinomyces israelii		1.98	0.23		1	
07752003	**Very good identification**					**07752003**
Actinomyces naeslundii		99.54	0.64	GAL 96		
Actinomyces viscosus		0.46	-.01			
07752010	**Good identification**					**07752010**
Actinomyces naeslundii		100.0	0.39	GAL 96		
07752011	**Good identification**					**07752011**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07752012	**Good identification**					**07752012**
Actinomyces naeslundii		100.0	0.39	GAL 96		
07752013	**Good identification**					**07752013**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07752020	**Very good identification**					**07752020**
Actinomyces naeslundii		100.0	0.61	GAL 96		
07752021	**Very good identification**					**07752021**
Actinomyces naeslundii		100.0	0.64	GAL 96		
07752022	**Good identification**					**07752022**
					SPO	
Actinomyces naeslundii		94.90	0.61	GAL 96	1	
Clostridium barati		1.58	0.04		99	
Clostridium ramosum		1.50	0.05		92	
07752023	**Very good identification**					**07752023**
Actinomyces naeslundii		100.0	0.64	GAL 96		
07752030	**Good identification**					**07752030**
Actinomyces naeslundii		100.0	0.39	GAL 96		
07752031	**Good identification**					**07752031**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07752032	**Good identification**					**07752032**
Actinomyces naeslundii		100.0	0.39	GAL 96		
07752033	**Good identification**					**07752033**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07752040	**Good identification**					**07752040**
Actinomyces naeslundii		100.0	0.48	GAL 96		
07752041	**Good identification**					**07752041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	GAL 96		
07752042	**Good identification**					**07752042**
Actinomyces naeslundii		95.46	0.48	GAL 96		
Actinomyces israelii		4.54	0.16			
07752043	**Good identification**					**07752043**
Actinomyces naeslundii		100.0	0.51	GAL 96		
07752060	**Good identification**					**07752060**
Actinomyces naeslundii		100.0	0.48	GAL 96		
07752061	**Good identification**					**07752061**
Actinomyces naeslundii		100.0	0.51	GAL 96		
07752062	**Good identification**					**07752062**
Actinomyces naeslundii		100.0	0.48	GAL 96		
07752063	**Good identification**					**07752063**
Actinomyces naeslundii		100.0	0.51	GAL 96		
07752100	**Good identification**					**07752100**
					GRA	
Actinomyces naeslundii		92.21	0.68	GAL 96	99	
Fusobacterium mortiferum		5.66	0.36		1	
Actinomyces israelii		1.86	0.30		99	
07752101	**Very good identification**					**07752101**
Actinomyces naeslundii		100.0	0.71	GAL 96		
07752102	**Low discrimination**					**07752102**
Actinomyces naeslundii		85.33	0.68	GAL 96		
Lactobacillus acidophilus		4.83	0.23			
Actinomyces israelii		2.59	0.33			
07752103	**Very good identification**					**07752103**
Actinomyces naeslundii		99.71	0.71	GAL 96		
Actinomyces viscosus		0.29	0.04			
07752110	**Acceptable identification**					**07752110**
Actinomyces naeslundii		90.03	0.47	GAL 96		
Actinomyces israelii		5.17	0.16	MLZ 5		
Bifidobacterium breve		4.81	0.10			
07752111	**Good identification**					**07752111**
Actinomyces naeslundii		100.0	0.50	GAL 96		
07752112	**Low discrimination**					**07752112**
Actinomyces naeslundii		47.22	0.47	GAL 96		
Bifidobacterium breve		25.47	0.27	MLZ 5		
Bifidobacterium adolescentis		23.24	0.26			
07752113	**Good identification**					**07752113**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	GAL 96		
07752120	**Low discrimination**					**07752120**
					SPO	
Actinomyces naeslundii		86.94	0.68	GAL 96	1	
Clostridium ramosum		11.93	0.28		92	
Clostridium barati		0.51	0.04		99	
07752121	**Very good identification**					**07752121**
Actinomyces naeslundii		100.0	0.71	GAL 96		
07752122	**Low discrimination**					**07752122**
					SPO	
Actinomyces naeslundii		58.27	0.68	GAL 96	1	
Clostridium ramosum		31.99	0.39		92	
Lactobacillus acidophilus		3.30	0.23		1	
07752123	**Very good identification**					**07752123**
					SPO	
Actinomyces naeslundii		99.63	0.71	GAL 96	1	
Clostridium ramosum		0.37	0.05		92	
07752130	**Acceptable identification**					**07752130**
Actinomyces naeslundii		91.34	0.47	GAL 96		
Bifidobacterium breve		8.66	0.15	MLZ 5		
07752131	**Good identification**					**07752131**
Actinomyces naeslundii		100.0	0.50	GAL 96		
07752132	**Low discrimination**					**07752132**
					SPO	
Actinomyces naeslundii		46.49	0.47	GAL 96	1	
Bifidobacterium breve		44.59	0.32	MLZ 5	1	
Clostridium ramosum		4.90	0.05		92	
07752133	**Good identification**					**07752133**
Actinomyces naeslundii		100.0	0.50	GAL 96		
07752140	**Acceptable identification**					**07752140**
Actinomyces naeslundii		94.30	0.56	GAL 96		
Actinomyces israelii		4.20	0.23			
Bifidobacterium breve		1.50	0.10			
07752141	**Very good identification**					**07752141**
Actinomyces naeslundii		100.0	0.59	GAL 96		
07752142	**Low discrimination**					**07752142**
Actinomyces naeslundii		77.29	0.56	GAL 96		
Bifidobacterium breve		12.44	0.27			
Actinomyces israelii		5.16	0.26			
07752143	**Very good identification**					**07752143**
Actinomyces naeslundii		100.0	0.59	GAL 96		
07752151	**Good identification**					**07752151**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752153	**Good identification**						**07752153**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752160	**Good identification**						**07752160**
Actinomyces naeslundii		97.25	0.56	GAL 96			
Bifidobacterium breve		2.75	0.15				
07752161	**Very good identification**						**07752161**
Actinomyces naeslundii		100.0	0.59	GAL 96			
07752162	**Low discrimination**						**07752162**
Actinomyces naeslundii		73.69	0.56	GAL 96			
Bifidobacterium breve		21.08	0.32				
Lactobacillus acidophilus		2.92	0.07				
07752163	**Very good identification**						**07752163**
Actinomyces naeslundii		100.0	0.59	GAL 96			
07752171	**Good identification**						**07752171**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752173	**Good identification**						**07752173**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752301	**Good identification**						**07752301**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752303	**Good identification**						**07752303**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752321	**Good identification**						**07752321**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752322	**Good identification**						**07752322**
					SPO	GRA	
Clostridium ramosum		95.86	0.41	TRE 94	92	50	
Prevotella oralis		2.25	0.13	GAL 99	1	1	
Actinomyces naeslundii		1.33	0.34		1	99	
07752323	**Low discrimination**						**07752323**
					SPO		
Actinomyces naeslundii		67.33	0.37	GAL 96	1		
Clostridium ramosum		32.67	0.07	RHA 1	92		
07752501	**Good identification**						**07752501**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07752502	**Low discrimination**						**07752502**
Bifidobacterium adolescentis		80.25	0.38	GAL 99			
Actinomyces israelii		14.03	0.32	XYL 87			
Actinomyces naeslundii		5.71	0.34				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07752503	**Good identification**					**07752503**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07752512	**Good identification**					**07752512**
Bifidobacterium adolescentis		97.91	0.40	GAL 99		
Actinomyces israelii		2.09	0.18	XYL 87		
07752521	**Good identification**					**07752521**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07752523	**Good identification**					**07752523**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07753000	**Good identification**					**07753000**
					SPO	
Actinomyces naeslundii		90.58	0.84		1	
Actinomyces israelii		5.37	0.54		1	
Clostridium innocuum		1.96	0.32		99	
07753001	**Very good identification**					**07753001**
Actinomyces naeslundii		99.96	0.87			
Actinomyces israelii		0.04	0.20			
07753002	**Low discrimination**					**07753002**
Actinomyces naeslundii		83.64	0.84			
Actinomyces israelii		7.44	0.57			
Lactobacillus acidophilus		4.84	0.39			
07753003	**Very good identification**					**07753003**
Actinomyces naeslundii		99.90	0.87			
Actinomyces israelii		0.06	0.23			
Lactobacillus acidophilus		0.04	0.05			
07753010	**Low discrimination**					**07753010**
Actinomyces naeslundii		81.68	0.62	MLZ 5		
Actinomyces israelii		13.76	0.40			
Lactobacillus jensenii		2.64	0.15			
07753011	**Very good identification**					**07753011**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
07753012	**Low discrimination**					**07753012**
Actinomyces naeslundii		70.18	0.62	MLZ 5		
Actinomyces israelii		17.73	0.43			
Bifidobacterium breve		4.49	0.27			
07753013	**Very good identification**					**07753013**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
07753020	**Very good identification**					**07753020**
					SPO	
Actinomyces naeslundii		96.97	0.84		1	
Clostridium ramosum		1.58	0.28		92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		0.43	0.17		99	
07753021	**Very good identification**					**07753021**
Actinomyces naeslundii		100.0	0.87			
07753022	**Low discrimination**					**07753022**
					SPO	
Actinomyces naeslundii		84.65	0.84		1	
Clostridium ramosum		5.51	0.39		92	
Lactobacillus acidophilus		4.90	0.39		1	
07753023	**Very good identification**					**07753023**
					SPO	
Actinomyces naeslundii		99.90	0.87		1	
Clostridium ramosum		0.04	0.05		92	
Lactobacillus acidophilus		0.04	0.05		1	
07753030	**Very good identification**					**07753030**
Actinomyces naeslundii		97.08	0.62	MLZ 5		
Bifidobacterium breve		1.09	0.15			
Actinomyces israelii		1.04	0.20			
07753031	**Very good identification**					**07753031**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
07753032	**Low discrimination**					**07753032**
Actinomyces naeslundii		85.12	0.62	MLZ 5		
Bifidobacterium breve		9.68	0.32			
Actinomyces israelii		1.37	0.23			
07753033	**Very good identification**					**07753033**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
07753040	**Low discrimination**					**07753040**
Actinomyces naeslundii		88.29	0.71			
Actinomyces israelii		11.54	0.47			
Bifidobacterium breve		0.17	0.10			
07753041	**Very good identification**					**07753041**
Actinomyces naeslundii		100.0	0.74			
07753042	**Low discrimination**					**07753042**
Actinomyces naeslundii		79.39	0.71			
Actinomyces israelii		15.57	0.50			
Lactobacillus acidophilus		3.22	0.23			
07753043	**Very good identification**					**07753043**
Actinomyces naeslundii		99.87	0.74			
Actinomyces israelii		0.13	0.16			
07753050	**Low discrimination**					**07753050**
Actinomyces naeslundii		71.07	0.50	MLZ 5		
Actinomyces israelii		26.38	0.33			
Bifidobacterium breve		2.55	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07753051	**Good identification**					**07753051**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07753052	**Low discrimination**					**07753052**
Actinomyces naeslundii		51.02	0.50	MLZ 5		
Actinomyces israelii		28.40	0.36			
Bifidobacterium breve		18.50	0.27			
07753053	**Good identification**					**07753053**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07753060	**Very good identification**					**07753060**
Actinomyces naeslundii		98.63	0.71			
Actinomyces israelii		0.82	0.27			
Bifidobacterium breve		0.33	0.15			
07753061	**Very good identification**					**07753061**
Actinomyces naeslundii		100.0	0.74			
07753062	**Good identification**					**07753062**
Actinomyces naeslundii		91.00	0.71			
Lactobacillus acidophilus		3.69	0.23			
Bifidobacterium breve		3.09	0.32			
07753063	**Very good identification**					**07753063**
Actinomyces naeslundii		100.0	0.74			
07753070	**Acceptable identification**					**07753070**
Actinomyces naeslundii		94.01	0.50	MLZ 5		
Bifidobacterium breve		5.99	0.15			
07753071	**Good identification**					**07753071**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07753072	**Low discrimination**					**07753072**
Actinomyces naeslundii		59.52	0.50	MLZ 5		
Bifidobacterium breve		38.36	0.32			
Actinomyces israelii		2.12	0.16			
07753073	**Good identification**					**07753073**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07753100	**Low discrimination**					**07753100**
Actinomyces naeslundii		89.51	0.92			
Actinomyces israelii		7.46	0.64			
Bifidobacterium breve		1.04	0.44			
07753101	**Excellent identification**					**07753101**
Actinomyces naeslundii		99.94	0.95			
Actinomyces israelii		0.06	0.30			
Bifidobacterium breve		0.01	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07753102	**Low discrimination**					**07753102**
Actinomyces naeslundii		62.13	0.92			
Lactobacillus acidophilus		14.52	0.57			
Actinomyces israelii		7.77	0.67			
07753103	**Excellent identification**					**07753103**
Actinomyces naeslundii		99.59	0.95			
Lactobacillus acidophilus		0.16	0.23			
Actinomyces israelii		0.08	0.33			
07753110	**Low discrimination**					**07753110**
Actinomyces naeslundii		65.92	0.70	MLZ 5		
Actinomyces israelii		15.60	0.50			
Bifidobacterium breve		14.51	0.44			
07753111	**Very good identification**					**07753111**
Actinomyces naeslundii		99.69	0.73	MLZ 5		
Actinomyces israelii		0.16	0.16			
Bifidobacterium breve		0.15	0.10			
07753112	**Low discrimination**					**07753112**
Bifidobacterium breve		38.48	0.62	ESL 99		
Bifidobacterium adolescentis		35.12	0.60			
Actinomyces naeslundii		17.29	0.70			
07753113	**Very good identification**					**07753113**
Actinomyces naeslundii		96.99	0.73	MLZ 5		
Bifidobacterium breve		1.45	0.27			
Bifidobacterium adolescentis		1.33	0.26			
07753120	**Low discrimination**					**07753120**
					SPO	
Actinomyces naeslundii		61.86	0.92		1	
Clostridium ramosum		35.03	0.63		92	
Bifidobacterium breve		1.27	0.49		1	
07753121	**Excellent identification**					**07753121**
					SPO	
Actinomyces naeslundii		99.60	0.95		1	
Clostridium ramosum		0.38	0.28		92	
Bifidobacterium breve		0.01	0.15		1	
07753122	**Low discrimination**					**07753122**
					SPO	
Clostridium ramosum		57.96	0.73	TRE 94	92	
Actinomyces naeslundii		25.59	0.92		1	
Lactobacillus acidophilus		5.98	0.57		1	
07753123	**Very good identification**					**07753123**
					SPO	
Actinomyces naeslundii		98.09	0.95		1	
Clostridium ramosum		1.50	0.39		92	
Lactobacillus acidophilus		0.15	0.23		1	
07753130	**Low discrimination**					**07753130**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		61.14	0.70	MLZ 5	1	
Bifidobacterium breve		23.92	0.49		1	
Clostridium tertium		7.27	0.31		99	
07753131	**Very good identification**					**07753131**
Actinomyces naeslundii		99.74	0.73	MLZ 5		
Bifidobacterium breve		0.26	0.15			
07753132	**Low discrimination**					**07753132**
					SPO	
Bifidobacterium breve		67.36	0.66	ESL 99	1	
Actinomyces naeslundii		17.03	0.70		1	
Clostridium ramosum		7.40	0.39		92	
07753133	**Very good identification**					**07753133**
					SPO	
Actinomyces naeslundii		96.90	0.73	MLZ 5	1	
Bifidobacterium breve		2.58	0.32		1	
Clostridium ramosum		0.28	0.05		92	
07753140	**Low discrimination**					**07753140**
Actinomyces naeslundii		79.85	0.79			
Actinomyces israelii		14.67	0.57			
Bifidobacterium breve		5.24	0.44			
07753141	**Very good identification**					**07753141**
Actinomyces naeslundii		99.83	0.82			
Actinomyces israelii		0.12	0.23			
Bifidobacterium breve		0.04	0.10			
07753142	**Low discrimination**					**07753142**
Actinomyces naeslundii		46.11	0.79			
Bifidobacterium breve		30.60	0.62			
Actinomyces israelii		12.71	0.60			
07753143	**Very good identification**					**07753143**
Actinomyces naeslundii		99.22	0.82			
Bifidobacterium breve		0.44	0.27			
Actinomyces israelii		0.18	0.26			
07753150	**Low discrimination**					**07753150**
Bifidobacterium breve		44.81	0.44	ESL 99		
Actinomyces naeslundii		35.93	0.57	BGL 91		
Actinomyces israelii		18.74	0.43			
07753151	**Very good identification**					**07753151**
Actinomyces naeslundii		99.17	0.60	MLZ 5		
Bifidobacterium breve		0.83	0.10			
07753152	**Low discrimination**					**07753152**
Bifidobacterium breve		79.58	0.62	ESL 99		
Bifidobacterium adolescentis		8.98	0.44			
Actinomyces naeslundii		6.31	0.57			
07753153	**Good identification**					**07753153**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.37	0.60	MLZ 5		
Bifidobacterium breve		7.76	0.27			
Bifidobacterium adolescentis		0.88	0.10			
07753160	**Low discrimination**					**07753160**
					SPO	
Actinomyces naeslundii		85.72	0.79		1	
Bifidobacterium breve		10.00	0.49		1	
Clostridium ramosum		2.78	0.28		92	
07753161	**Very good identification**					**07753161**
Actinomyces naeslundii		99.92	0.82			
Bifidobacterium breve		0.08	0.15			
07753162	**Low discrimination**					**07753162**
Bifidobacterium breve		47.04	0.66	ESL 99		
Actinomyces naeslundii		39.87	0.79			
Lactobacillus acidophilus		6.52	0.41			
07753163	**Very good identification**					**07753163**
Actinomyces naeslundii		99.02	0.82			
Bifidobacterium breve		0.79	0.32			
Lactobacillus acidophilus		0.11	0.07			
07753170	**Low discrimination**					**07753170**
Bifidobacterium breve		68.21	0.49	ESL 99		
Actinomyces naeslundii		30.77	0.57	BGL 91		
Actinomyces israelii		1.02	0.23			
07753171	**Very good identification**					**07753171**
Actinomyces naeslundii		98.53	0.60	MLZ 5		
Bifidobacterium breve		1.47	0.15			
07753172	**Very good identification**					**07753172**
Bifidobacterium breve		95.16	0.66	ESL 99		
Actinomyces naeslundii		4.25	0.57			
Actinomyces israelii		0.21	0.26			
07753173	**Low discrimination**					**07753173**
Actinomyces naeslundii		86.88	0.60	MLZ 5		
Bifidobacterium breve		13.12	0.32			
07753200	**Low discrimination**					**07753200**
Actinomyces naeslundii		69.35	0.50	RHA 1		
Actinomyces israelii		30.65	0.35			
07753201	**Good identification**					**07753201**
Actinomyces naeslundii		100.0	0.53	RHA 1		
07753202	**Low discrimination**					**07753202**
Actinomyces naeslundii		43.77	0.50	RHA 1		
Actinomyces israelii		29.02	0.38			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		18.86	0.19				
07753203	**Good identification**						**07753203**
Actinomyces naeslundii		100.0	0.53	RHA 1			
07753221	**Good identification**						**07753221**
Actinomyces naeslundii		100.0	0.53	RHA 1			
07753222	**Low discrimination**						**07753222**
					SPO	GRA	
Clostridium ramosum		74.96	0.41	TRE 94	92	50	
Prevotella oralis		11.12	0.26	LAC 99	1	1	
Actinomyces naeslundii		8.77	0.50		1	99	
07753223	**Acceptable identification**						**07753223**
					SPO		
Actinomyces naeslundii		94.56	0.53	RHA 1	1		
Clostridium ramosum		5.44	0.07		92		
07753240	**Low discrimination**						**07753240**
Actinomyces naeslundii		50.66	0.37	RHA 1			
Actinomyces israelii		49.34	0.28				
07753241	**Good identification**						**07753241**
Actinomyces naeslundii		100.0	0.40	RHA 1			
07753243	**Good identification**						**07753243**
Actinomyces naeslundii		100.0	0.40	RHA 1			
07753260	**Good identification**						**07753260**
Actinomyces naeslundii		100.0	0.37	RHA 1			
07753261	**Good identification**						**07753261**
Actinomyces naeslundii		100.0	0.40	RHA 1			
07753262	**Low discrimination**						**07753262**
					SPO		
Actinomyces naeslundii		55.84	0.37	RHA 1	1		
Clostridium ramosum		27.31	0.07		92		
Lactobacillus acidophilus		16.85	0.04		1		
07753263	**Good identification**						**07753263**
Actinomyces naeslundii		100.0	0.40	RHA 1			
07753300	**Low discrimination**						**07753300**
					SPO		
Actinomyces naeslundii		41.09	0.58	RHA 1	1		
Clostridium ramosum		30.84	0.31		92		
Actinomyces israelii		25.53	0.45		1		
07753301	**Very good identification**						**07753301**
Actinomyces naeslundii		100.0	0.61	RHA 1			
07753302	**Low discrimination**						**07753302**
					SPO		
Clostridium ramosum		41.20	0.41	TRE 94	92		
Lactobacillus acidophilus		23.89	0.38	NAG 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		13.72	0.58		1		
07753303	**Very good identification**						**07753303**
					SPO		
Actinomyces naeslundii		96.32	0.61	RHA 1	1		
Clostridium ramosum		1.95	0.07		92		
Lactobacillus acidophilus		1.13	0.03		1		
07753311	**Good identification**						**07753311**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07753313	**Good identification**						**07753313**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07753320	**Very good identification**						**07753320**
					SPO	GRA	
Clostridium ramosum		98.04	0.65	TRE 94	92	50	
Actinomyces naeslundii		1.32	0.58		1	99	
Prevotella oralis		0.49	0.25		1	1	
07753321	**Low discrimination**						**07753321**
					SPO		
Actinomyces naeslundii		66.65	0.61	RHA 1	1		
Clostridium ramosum		33.35	0.31		92		
07753322	**Very good identification**						**07753322**
					SPO	GRA	
Clostridium ramosum		96.68	0.75	TRE 94	92	50	
Prevotella oralis		2.27	0.47		1	1	
Lactobacillus acidophilus		0.57	0.38		1	99	
07753323	**Low discrimination**						**07753323**
					SPO	GRA	
Clostridium ramosum		65.40	0.41	TRE 94	92	50	
Actinomyces naeslundii		32.68	0.61	URE 1	1	99	
Prevotella oralis		1.54	0.13		1	1	
07753331	**Good identification**						**07753331**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07753332	**Low discrimination**						**07753332**
					SPO	GRA	
Clostridium ramosum		87.31	0.41	TRE 94	92	50	
Bifidobacterium breve		6.06	0.32	MLZ 1	1	99	
Prevotella oralis		2.05	0.13		1	1	
07753333	**Low discrimination**						**07753333**
					SPO		
Actinomyces naeslundii		72.25	0.39	RHA 1	1		
Clostridium ramosum		27.75	0.07	MLZ 5	92		
07753340	**Low discrimination**						**07753340**
Actinomyces israelii		56.23	0.38	XYL 92			
Actinomyces naeslundii		41.07	0.45	RHA 7			
Bifidobacterium breve		2.70	0.10				
07753341	**Good identification**						**07753341**
Actinomyces naeslundii		100.0	0.48	RHA 1			
07753342	**Low discrimination**						**07753342**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii	39.72	0.41	XYL 92		
Lactobacillus acidophilus	23.59	0.22	RHA 7		
Actinomyces naeslundii	19.34	0.45			
07753343 Good identification					**07753343**
Actinomyces naeslundii	100.0	0.48	RHA 1		
07753361 Good identification					**07753361**
Actinomyces naeslundii	100.0	0.48	RHA 1		
07753362 Low discrimination					**07753362**
				SPO	
Clostridium ramosum	81.01	0.41	TRE 94	92	
Lactobacillus acidophilus	5.81	0.22	SOR 1	1	
Bifidobacterium breve	5.62	0.32		1	
07753363 Low discrimination					**07753363**
				SPO	
Actinomyces naeslundii	89.72	0.48	RHA 1	1	
Clostridium ramosum	10.28	0.07		92	
07753400 Low discrimination					**07753400**
				SPO	
Actinomyces israelii	76.96	0.53	XYL 92	1	
Actinomyces naeslundii	16.02	0.50		1	
Clostridium innocuum	6.06	0.19		99	
07753401 Good identification					**07753401**
Actinomyces naeslundii	96.87	0.53	ARA 1		
Actinomyces israelii	3.13	0.18			
07753402 Low discrimination					**07753402**
Actinomyces israelii	69.91	0.56	XYL 92		
Bifidobacterium adolescentis	16.16	0.38			
Actinomyces naeslundii	9.70	0.50			
07753403 Good identification					**07753403**
Actinomyces naeslundii	95.37	0.53	ARA 1		
Actinomyces israelii	4.63	0.21			
07753410 Acceptable identification					**07753410**
Actinomyces israelii	93.17	0.38	XYL 92		
Actinomyces naeslundii	6.83	0.28	MLZ 13		
07753412 Low discrimination					**07753412**
Bifidobacterium adolescentis	64.31	0.40	XYL 87		
Actinomyces israelii	34.03	0.42	LAC 99		
Actinomyces naeslundii	1.66	0.28			
07753420 Low discrimination					**07753420**
Actinomyces naeslundii	76.53	0.50	ARA 1		
Actinomyces israelii	23.47	0.32			
07753421 Good identification					**07753421**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	ARA 1		
07753422	**Low discrimination**					**07753422**
					SPO	
Actinomyces naeslundii		62.00	0.50	ARA 1	1	
Actinomyces israelii		28.52	0.35		1	
Clostridium ramosum		4.04	0.05		92	
07753423	**Good identification**					**07753423**
Actinomyces naeslundii		100.0	0.53	ARA 1		
07753440	**Acceptable identification**					**07753440**
Actinomyces israelii		91.37	0.46	XYL 92		
Actinomyces naeslundii		8.63	0.37			
07753441	**Good identification**					**07753441**
Actinomyces naeslundii		100.0	0.40	ARA 1		
07753442	**Low discrimination**					**07753442**
Actinomyces israelii		88.00	0.49	XYL 92		
Bifidobacterium adolescentis		6.46	0.23			
Actinomyces naeslundii		5.54	0.37			
07753443	**Acceptable identification**					**07753443**
Actinomyces naeslundii		90.34	0.40	ARA 1		
Actinomyces israelii		9.66	0.14			
07753460	**Low discrimination**					**07753460**
Actinomyces naeslundii		59.67	0.37	ARA 1		
Actinomyces israelii		40.33	0.25			
07753461	**Good identification**					**07753461**
Actinomyces naeslundii		100.0	0.40	ARA 1		
07753463	**Good identification**					**07753463**
Actinomyces naeslundii		100.0	0.40	ARA 1		
07753500	**Low discrimination**					**07753500**
Actinomyces israelii		76.27	0.63	XYL 92		
Actinomyces naeslundii		11.29	0.58			
Bifidobacterium adolescentis		6.61	0.38			
07753501	**Very good identification**					**07753501**
Actinomyces naeslundii		95.65	0.61	ARA 1		
Actinomyces israelii		4.35	0.29			
07753502	**Low discrimination**					**07753502**
Bifidobacterium adolescentis		83.19	0.73	XYL 87		
Actinomyces israelii		14.55	0.66			
Actinomyces naeslundii		1.44	0.58			
07753503	**Low discrimination**					**07753503**
Actinomyces naeslundii		68.57	0.61	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		26.75	0.38			
Actinomyces israelii		4.68	0.32			
07753510	**Low discrimination**					**07753510**
Actinomyces israelii		55.59	0.49	XYL 92		
Bifidobacterium adolescentis		39.41	0.40	MLZ 13		
Actinomyces naeslundii		2.90	0.36			
07753511	**Low discrimination**					**07753511**
Actinomyces naeslundii		88.56	0.39	ARA 1		
Actinomyces israelii		11.44	0.15	MLZ 5		
07753512	**Very good identification**					**07753512**
Bifidobacterium adolescentis		97.62	0.74	XYL 87		
Actinomyces israelii		2.09	0.52			
Bifidobacterium breve		0.16	0.27			
07753513	**Low discrimination**					**07753513**
Bifidobacterium adolescentis		88.36	0.40	XYL 87		
Actinomyces naeslundii		9.75	0.39	URE 1		
Actinomyces israelii		1.89	0.18			
07753520	**Low discrimination**					**07753520**
					SPO	
Actinomyces naeslundii		48.30	0.58	ARA 1	1	
Clostridium ramosum		27.35	0.28		92	
Actinomyces israelii		20.82	0.42		1	
07753521	**Very good identification**					**07753521**
Actinomyces naeslundii		100.0	0.61	ARA 1		
07753522	**Low discrimination**					**07753522**
					SPO	
Clostridium ramosum		43.86	0.39	TRE 94	92	
Actinomyces naeslundii		19.37	0.58	ARA 1	1	
Actinomyces israelii		12.52	0.45		1	
07753523	**Very good identification**					**07753523**
					SPO	
Actinomyces naeslundii		98.50	0.61	ARA 1	1	
Clostridium ramosum		1.50	0.05		92	
07753531	**Good identification**					**07753531**
Actinomyces naeslundii		100.0	0.39	ARA 1		
07753532	**Low discrimination**					**07753532**
Bifidobacterium adolescentis		61.07	0.40	XYL 87		
Bifidobacterium breve		17.78	0.32	NAG 1		
Actinomyces israelii		8.25	0.31			
07753533	**Good identification**					**07753533**
Actinomyces naeslundii		100.0	0.39	ARA 1		
07753540	**Low discrimination**					**07753540**
Actinomyces israelii		89.62	0.56	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		6.02	0.45			
Bifidobacterium adolescentis		2.47	0.23			
07753541	**Acceptable identification**					**07753541**
Actinomyces naeslundii		90.89	0.48	ARA 1		
Actinomyces israelii		9.11	0.22			
07753542	**Low discrimination**					**07753542**
Bifidobacterium adolescentis		62.60	0.57	XYL 87		
Actinomyces israelii		34.44	0.59	SOR 11		
Actinomyces naeslundii		1.54	0.45			
07753543	**Low discrimination**					**07753543**
Actinomyces naeslundii		70.25	0.48	ARA 1		
Bifidobacterium adolescentis		19.19	0.23			
Actinomyces israelii		10.56	0.25			
07753550	**Low discrimination**					**07753550**
					COC	
Actinomyces israelii		75.96	0.42	XYL 92	1	
Bifidobacterium adolescentis		17.11	0.24	MLZ 13	1	
Blautia producta		2.89	-.03		99	
07753552	**Good identification**					**07753552**
Bifidobacterium adolescentis		92.45	0.59	XYL 87		
Actinomyces israelii		6.22	0.45	SOR 11		
Bifidobacterium breve		1.24	0.27			
07753560	**Low discrimination**					**07753560**
Actinomyces naeslundii		48.39	0.45	ARA 1		
Actinomyces israelii		45.97	0.35			
Bifidobacterium breve		5.65	0.15			
07753561	**Good identification**					**07753561**
Actinomyces naeslundii		100.0	0.48	ARA 1		
07753562	**Low discrimination**					**07753562**
Actinomyces israelii		33.08	0.38	XYL 92		
Bifidobacterium breve		27.39	0.32	NAG 6		
Actinomyces naeslundii		23.21	0.45			
07753563	**Good identification**					**07753563**
Actinomyces naeslundii		100.0	0.48	ARA 1		
07753602	**Good identification**					**07753602**
Actinomyces israelii		95.29	0.36	XYL 92		
Clostridium clostridioforme		4.72	0.05	RHA 7		
07753700	**Good identification**					**07753700**
					SPO	
Actinomyces israelii		95.19	0.44	XYL 92	1	
Clostridium butyricum		2.92	-.03	RHA 7	99	
Actinomyces naeslundii		1.89	0.24		1	
07753702	**Low discrimination**					**07753702**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		53.05	0.47	XYL 92			
Bifidobacterium adolescentis		40.72	0.38	RHA 7			
Clostridium clostridioforme		2.20	0.14				
07753712	**Low discrimination**						**07753712**
Bifidobacterium adolescentis		73.51	0.40	XYL 87			
Clostridium clostridioforme		14.79	0.25	RHA 1			
Actinomyces israelii		11.71	0.33				
07753722	**Acceptable identification**						**07753722**
					SPO		
Clostridium ramosum		92.30	0.41	TRE 94	92		
Prevotella oralis		2.17	0.13	ARA 1	1		
Parabacteroides distasonis		1.77	0.24		1		
07753740	**Good identification**						**07753740**
Actinomyces israelii		100.0	0.37	XYL 92			
07753742	**Low discrimination**						**07753742**
Actinomyces israelii		80.39	0.40	XYL 92			
Bifidobacterium adolescentis		19.61	0.23	RHA 7			
07754000	**Good identification**						**07754000**
					GRA		
Fusobacterium mortiferum		95.03	0.45	MLT 11	1		
Actinomyces israelii		4.97	0.25	XYL 7	99		
07754100	**Low discrimination**						**07754100**
					GRA	SPO	
Fusobacterium mortiferum		84.16	0.51	MLT 11	1	1	
Actinomyces israelii		7.54	0.36	XYL 7	99	1	
Clostridium butyricum		7.34	0.14		99	99	
07754102	**Low discrimination**						**07754102**
					GRA		
Bifidobacterium adolescentis		45.84	0.40	GAL 99	99		
Actinomyces israelii		16.27	0.39	ARA 87	99		
Fusobacterium mortiferum		13.45	0.35		1		
07754112	**Low discrimination**						**07754112**
Bifidobacterium adolescentis		65.73	0.41	GAL 99			
Clostridium clostridioforme		31.42	0.33	ARA 87			
Actinomyces israelii		2.85	0.24				
07754122	**Low discrimination**						**07754122**
Prevotella oralis		80.41	0.37	GAL 99			
Parabacteroides distasonis		11.26	0.36	XYL 11			
Clostridium clostridioforme		3.67	0.22				
07754212	**Good identification**						**07754212**
Clostridium clostridioforme		100.0	0.39	TRE 88			
07754232	**Good identification**						**07754232**
Clostridium clostridioforme		100.0	0.39	TRE 88			
07754302	**Acceptable identification**						**07754302**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Clostridium clostridioforme		90.21	0.36	TRE 88	50		
Prevotella oralis		6.09	0.09	ARA 94	1		
Actinomyces israelii		2.56	0.19		99		
07754312	**Good identification**						**07754312**
Clostridium clostridioforme		99.71	0.48	TRE 88			
Bifidobacterium adolescentis		0.29	0.07	ARA 94			
07754332	**Low discrimination**						**07754332**
Clostridium clostridioforme		86.85	0.48	TRE 88			
Parabacteroides distasonis		12.90	0.39	ARA 94			
Prevotella oralis		0.25	-.03				
07754412	**Good identification**						**07754412**
Clostridium clostridioforme		97.07	0.44	TRE 88			
Bifidobacterium adolescentis		2.93	0.21	RHA 88			
07754432	**Good identification**						**07754432**
Clostridium clostridioforme		100.0	0.44	TRE 88			
07754502	**Low discrimination**						**07754502**
					SPO		
Bifidobacterium adolescentis		68.30	0.54	GAL 99	1		
Clostridium clostridioforme		20.51	0.42		66		
Clostridium butyricum		8.23	0.24		99		
07754512	**Low discrimination**						**07754512**
Clostridium clostridioforme		52.55	0.53	TRE 88			
Bifidobacterium adolescentis		46.96	0.56	RHA 88			
Actinomyces israelii		0.25	0.23				
07754522	**Low discrimination**						**07754522**
					GRA	SPO	
Clostridium clostridioforme		81.05	0.42	TRE 88	50	66	
Parabacteroides distasonis		9.74	0.32	RHA 88	1	1	
Clostridium butyricum		2.83	0.05		99	99	
07754532	**Good identification**						**07754532**
					GRA		
Clostridium clostridioforme		97.73	0.53	TRE 88	50		
Parabacteroides distasonis		1.39	0.27	RHA 88	1		
Bifidobacterium adolescentis		0.88	0.21		99		
07754542	**Low discrimination**						**07754542**
					SPO		
Bifidobacterium adolescentis		80.27	0.39	GAL 99	1		
Actinomyces israelii		10.96	0.30	SOR 11	1		
Clostridium butyricum		6.80	0.05		99		
07754552	**Acceptable identification**						**07754552**
Bifidobacterium adolescentis		90.24	0.40	GAL 99			
Clostridium clostridioforme		8.25	0.19	SOR 11			
Actinomyces israelii		1.51	0.16				
07754602	**Good identification**						**07754602**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.48	TRE 88			
07754612	**Very good identification**						**07754612**
Clostridium clostridioforme		100.0	0.59	TRE 88			
07754622	**Good identification**						**07754622**
Clostridium clostridioforme		100.0	0.48	TRE 88			
07754632	**Very good identification**						**07754632**
Clostridium clostridioforme		100.0	0.59	TRE 88			
07754702	**Very good identification**						**07754702**
					SPO		
Clostridium clostridioforme		99.23	0.57	TRE 88	66		
Bifidobacterium adolescentis		0.46	0.20		1		
Clostridium butyricum		0.17	-.02		99		
07754712	**Very good identification**						**07754712**
Clostridium clostridioforme		99.88	0.68	TRE 88			
Bifidobacterium adolescentis		0.12	0.21				
07754722	**Good identification**						**07754722**
Clostridium clostridioforme		94.41	0.57	TRE 88			
Parabacteroides distasonis		4.64	0.40				
Bacteroides ovatus		0.39	0.06				
07754732	**Very good identification**						**07754732**
Clostridium clostridioforme		99.36	0.68	TRE 88			
Parabacteroides distasonis		0.58	0.35				
Bacteroides ovatus		0.06	0.03				
07755000	**Low discrimination**						**07755000**
					GRA	SPO	
Actinomyces israelii		61.20	0.59		99	1	
Fusobacterium mortiferum		17.74	0.48		1	1	
Clostridium butyricum		9.93	0.25		99	99	
07755001	**Low discrimination**						**07755001**
					GRA		
Actinomyces naeslundii		83.27	0.48	XYL 1	99		
Actinomyces israelii		12.97	0.25		99		
Fusobacterium mortiferum		3.76	0.14		1		
07755002	**Low discrimination**						**07755002**
					GRA		
Actinomyces israelii		68.00	0.62		99		
Bifidobacterium adolescentis		7.74	0.40		99		
Prevotella oralis		7.25	0.29		1		
07755003	**Low discrimination**						**07755003**
Actinomyces naeslundii		81.06	0.48	XYL 1			
Actinomyces israelii		18.94	0.28				
07755010	**Low discrimination**						**07755010**
					SPO		
Actinomyces israelii		83.28	0.45	MLZ 13	1		
Lactobacillus jensenii		7.89	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		4.76	0.03		99	
07755012	**Low discrimination**					**07755012**
Actinomyces israelii		41.71	0.48	MLZ 13		
Bifidobacterium adolescentis		38.85	0.41			
Clostridium clostridioforme		16.64	0.32			
07755021	**Good identification**					**07755021**
Actinomyces naeslundii		100.0	0.48	XYL 1		
07755022	**Low discrimination**					**07755022**
					GRA	
Prevotella oralis		89.12	0.51	XYL 11	1	
Actinomyces israelii		2.81	0.42	LAC 94	99	
Lactobacillus acidophilus		2.71	0.26		99	
07755023	**Low discrimination**					**07755023**
					GRA	
Actinomyces naeslundii		67.88	0.48	XYL 1	99	
Prevotella oralis		32.12	0.17		1	
07755040	**Acceptable identification**					**07755040**
					SPO	
Actinomyces israelii		94.04	0.52		1	
Clostridium butyricum		3.41	0.07		99	
Actinomyces naeslundii		1.84	0.32		1	
07755041	**Low discrimination**					**07755041**
Actinomyces naeslundii		74.44	0.35	XYL 1		
Actinomyces israelii		25.56	0.18			
07755042	**Acceptable identification**					**07755042**
Actinomyces israelii		92.75	0.55			
Bifidobacterium adolescentis		3.36	0.24			
Lactobacillus acidophilus		1.81	0.11			
07755043	**Low discrimination**					**07755043**
Actinomyces naeslundii		66.01	0.35	XYL 1		
Actinomyces israelii		33.99	0.21			
07755050	**Good identification**					**07755050**
Actinomyces israelii		100.0	0.38	MLZ 13		
07755052	**Low discrimination**					**07755052**
Actinomyces israelii		77.16	0.41	MLZ 13		
Bifidobacterium adolescentis		22.84	0.26			
07755061	**Good identification**					**07755061**
Actinomyces naeslundii		100.0	0.35	XYL 1		
07755063	**Good identification**					**07755063**
Actinomyces naeslundii		100.0	0.35	XYL 1		
07755100	**Low discrimination**					**07755100**
					GRA SPO	
Actinomyces israelii		43.31	0.70		99 1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		42.18	0.48		99	99	
Fusobacterium mortiferum		7.32	0.54		1	1	
07755101	**Low discrimination**						**07755101**
					SPO		
Actinomyces naeslundii		68.06	0.56	XYL 1	1		
Actinomyces israelii		14.90	0.36		1		
Clostridium butyricum		14.51	0.14		99		
07755102	**Low discrimination**						**07755102**
					GRA		
Bifidobacterium adolescentis		58.97	0.74	ARA 87	99		
Actinomyces israelii		20.92	0.73		99		
Prevotella oralis		8.73	0.50		1		
07755103	**Low discrimination**						**07755103**
Actinomyces naeslundii		39.27	0.56	XYL 1			
Bifidobacterium adolescentis		36.35	0.40				
Actinomyces israelii		12.90	0.39				
07755110	**Low discrimination**						**07755110**
					SPO		
Actinomyces israelii		58.00	0.56	MLZ 13	1		
Bifidobacterium adolescentis		20.26	0.41		1		
Clostridium butyricum		19.89	0.27		99		
07755112	**Good identification**						**07755112**
Bifidobacterium adolescentis		93.71	0.76	ARA 87			
Actinomyces israelii		4.07	0.59				
Clostridium clostridioforme		1.36	0.41				
07755113	**Acceptable identification**						**07755113**
Bifidobacterium adolescentis		90.55	0.41	ARA 87			
Actinomyces naeslundii		4.21	0.34	URE 1			
Actinomyces israelii		3.93	0.24				
07755120	**Low discrimination**						**07755120**
					GRA	SPO	
Prevotella oralis		66.79	0.50	XYL 11	1	1	
Clostridium butyricum		9.03	0.30	BGL 95	99	99	
Actinomyces israelii		6.81	0.49		99	1	
07755121	**Low discrimination**						**07755121**
					GRA	SPO	
Actinomyces naeslundii		84.63	0.56	XYL 1	99	1	
Prevotella oralis		11.60	0.16		1	1	
Clostridium butyricum		1.57	-.04		99	99	
07755122	**Low discrimination**						**07755122**
					GRA		
Prevotella oralis		83.31	0.72	XYL 11	1		
Parabacteroides distasonis		11.67	0.70		1		
Lactobacillus acidophilus		1.86	0.45		99		
07755123	**Low discrimination**						**07755123**
					GRA		
Prevotella oralis		63.81	0.37	XYL 11	1		
Actinomyces naeslundii		24.50	0.56	URE 1	99		
Parabacteroides distasonis		8.94	0.36		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07755130	**Low discrimination**						**07755130**
					GRA		
Actinomyces israelii		27.58	0.35	MLZ 13	99		
Parabacteroides distasonis		26.20	0.31	NAG 6	1		
Prevotella oralis		18.29	0.16		1		
07755132	**Low discrimination**						**07755132**
Parabacteroides distasonis		76.01	0.65				
Prevotella oralis		10.18	0.37				
Clostridium clostridioforme		6.34	0.41				
07755140	**Low discrimination**						**07755140**
					SPO		
Actinomyces israelii		79.91	0.63		1		
Clostridium butyricum		17.40	0.30		99		
Actinomyces naeslundii		1.12	0.40		1		
07755141	**Low discrimination**						**07755141**
					SPO		
Actinomyces naeslundii		62.99	0.43	XYL 1	1		
Actinomyces israelii		30.39	0.29		1		
Clostridium butyricum		6.62	-.04		99		
07755142	**Low discrimination**						**07755142**
					SPO		
Actinomyces israelii		49.01	0.66		1		
Bifidobacterium adolescentis		43.90	0.59		1		
Clostridium butyricum		2.77	0.23		99		
07755143	**Low discrimination**						**07755143**
Actinomyces naeslundii		42.16	0.43	XYL 1			
Actinomyces israelii		30.51	0.32				
Bifidobacterium adolescentis		27.33	0.24				
07755150	**Low discrimination**						**07755150**
					SPO		
Actinomyces israelii		83.36	0.49	MLZ 13	1		
Bifidobacterium adolescentis		9.25	0.26		1		
Clostridium butyricum		6.39	0.08		99		
07755152	**Low discrimination**						**07755152**
Bifidobacterium adolescentis		86.86	0.60	ARA 87			
Actinomyces israelii		11.86	0.52	SOR 11			
Bifidobacterium breve		0.95	0.27				
07755160	**Low discrimination**						**07755160**
					GRA	SPO	
Actinomyces israelii		52.63	0.42	NAG 6	99	1	
Clostridium butyricum		15.61	0.12		99	99	
Prevotella oralis		13.41	0.16		1	1	
07755161	**Good identification**						**07755161**
Actinomyces naeslundii		100.0	0.43	XYL 1			
07755162	**Low discrimination**						**07755162**
					GRA		
Prevotella oralis		51.06	0.37	XYL 11	1		
Actinomyces israelii		15.82	0.45	SOR 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		13.92	0.29		99		
07755163	**Low discrimination**						**07755163**
					GRA		
Actinomyces naeslundii		87.03	0.43	XYL 1	99		
Prevotella oralis		12.98	0.03		1		
07755200	**Low discrimination**						**07755200**
					GRA	SPO	
Actinomyces israelii		86.69	0.40	RHA 7	99	1	
Clostridium butyricum		5.78	-.01		99	99	
Prevotella oralis		4.15	0.01		1	1	
07755202	**Low discrimination**						**07755202**
					GRA		
Clostridium clostridioforme		43.06	0.36	TRE 88	50		
Actinomyces israelii		33.77	0.43	ARA 94	99		
Prevotella oralis		20.49	0.23		1		
07755212	**Good identification**						**07755212**
Clostridium clostridioforme		97.19	0.47	TRE 88			
Actinomyces israelii		2.50	0.29	ARA 94			
Bifidobacterium adolescentis		0.31	0.07				
07755222	**Low discrimination**						**07755222**
Prevotella oralis		78.33	0.45	XYL 11			
Parabacteroides distasonis		12.15	0.44	LAC 94			
Clostridium clostridioforme		8.66	0.36				
07755232	**Low discrimination**						**07755232**
Clostridium clostridioforme		84.33	0.47	TRE 88			
Parabacteroides distasonis		13.98	0.39	ARA 94			
Prevotella oralis		1.69	0.11				
07755242	**Good identification**						**07755242**
Actinomyces israelii		100.0	0.36	RHA 7			
07755300	**Low discrimination**						**07755300**
					SPO		
Actinomyces israelii		52.17	0.50	RHA 7	1		
Clostridium butyricum		20.87	0.23		99		
Eubacterium contortum		11.32	0.24		1		
07755302	**Low discrimination**						**07755302**
					GRA		
Prevotella oralis		42.63	0.43	XYL 11	1		
Clostridium clostridioforme		19.14	0.45	NAG 95	50		
Actinomyces israelii		17.94	0.53		99		
07755310	**Low discrimination**						**07755310**
					SPO		
Actinomyces israelii		59.50	0.36	RHA 7	1		
Clostridium clostridioforme		29.33	0.22	MLZ 13	66		
Clostridium butyricum		8.39	0.01		99		
07755312	**Low discrimination**						**07755312**
					GRA		
Clostridium clostridioforme		84.65	0.56	TRE 88	50		
Bifidobacterium adolescentis		8.05	0.41	ARA 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		4.19	0.39		1	
07755320	**Low discrimination**					**07755320**
Bacteroides vulgatus		59.46	0.42	CEL 5		
Prevotella oralis		32.16	0.43	ARA 93		
Parabacteroides distasonis		6.05	0.44			
07755322	**Low discrimination**					**07755322**
Prevotella oralis		49.43	0.65	XYL 11		
Parabacteroides distasonis		48.45	0.78			
Clostridium clostridioforme		1.17	0.45			
07755330	**Low discrimination**					**07755330**
Parabacteroides distasonis		66.03	0.39	BGL 99		
Clostridium clostridioforme		13.47	0.22			
Bacteroides vulgatus		12.17	0.08			
07755332	**Low discrimination**					**07755332**
Parabacteroides distasonis		81.59	0.73			
Clostridium clostridioforme		16.64	0.56			
Prevotella oralis		1.56	0.31			
07755333	**Low discrimination**					**07755333**
Parabacteroides distasonis		81.76	0.39	URE 1		
Clostridium clostridioforme		16.68	0.22			
Prevotella oralis		1.56	-.03			
07755340	**Acceptable identification**					**07755340**
					SPO	
Actinomyces israelii		91.79	0.43	RHA 7	1	
Clostridium butyricum		8.21	0.04		99	
07755342	**Low discrimination**					**07755342**
					GRA	
Actinomyces israelii		76.89	0.46	RHA 7	99	
Bifidobacterium adolescentis		9.24	0.24		99	
Prevotella oralis		4.75	0.09		1	
07755372	**Low discrimination**					**07755372**
Parabacteroides distasonis		81.76	0.39	SOR 1		
Clostridium clostridioforme		16.68	0.22			
Prevotella oralis		1.56	-.03			
07755400	**Low discrimination**					**07755400**
					SPO	
Clostridium butyricum		63.52	0.41	ESL 88	99	
Actinomyces israelii		35.58	0.58	TRE 94	1	
Bifidobacterium adolescentis		0.50	0.20	LAC 96	1	
07755402	**Low discrimination**					**07755402**
					SPO	
Actinomyces israelii		37.77	0.61		1	
Bifidobacterium adolescentis		35.18	0.54		1	
Clostridium butyricum		17.48	0.34		99	
07755410	**Low discrimination**					**07755410**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		53.76	0.44	MLZ 13	1	
Clostridium butyricum		33.81	0.19		99	
Clostridium clostridioforme		6.22	0.18		66	
07755412	**Low discrimination**					**07755412**
Clostridium clostridioforme		46.52	0.52	TRE 88		
Bifidobacterium adolescentis		46.39	0.56	RHA 88		
Actinomyces israelii		6.09	0.47			
07755422	**Low discrimination**					**07755422**
					SPO	
Clostridium clostridioforme		57.61	0.41	TRE 88	66	
Actinomyces israelii		14.68	0.40	RHA 88	1	
Clostridium butyricum		9.25	0.16		99	
07755432	**Good identification**					**07755432**
					GRA	
Clostridium clostridioforme		96.51	0.52	TRE 88	50	
Parabacteroides distasonis		1.53	0.27	RHA 88	1	
Bifidobacterium adolescentis		0.97	0.21		99	
07755440	**Low discrimination**					**07755440**
					SPO	
Actinomyces israelii		71.47	0.51		1	
Clostridium butyricum		28.53	0.23		99	
07755442	**Low discrimination**					**07755442**
					SPO	
Actinomyces israelii		71.12	0.54		1	
Bifidobacterium adolescentis		21.05	0.39		1	
Clostridium butyricum		7.36	0.16		99	
07755450	**Low discrimination**					**07755450**
					SPO	
Actinomyces israelii		87.67	0.37	MLZ 13	1	
Clostridium butyricum		12.33	0.01		99	
07755452	**Low discrimination**					**07755452**
Bifidobacterium adolescentis		66.22	0.40	LAC 99		
Actinomyces israelii		27.35	0.40	SOR 11		
Clostridium clostridioforme		5.43	0.18			
07755500	**Low discrimination**					**07755500**
					SPO	
Clostridium butyricum		88.67	0.65	ESL 88	99	
Actinomyces israelii		8.28	0.68	TRE 94	1	
Bifidobacterium adolescentis		2.89	0.54		1	
07755502	**Low discrimination**					**07755502**
					SPO	
Bifidobacterium adolescentis		85.19	0.88		1	
Clostridium butyricum		10.26	0.58		99	
Actinomyces israelii		3.70	0.71		1	
07755503	**Low discrimination**					**07755503**
					SPO	
Bifidobacterium adolescentis		85.26	0.54	URE 1	1	
Clostridium butyricum		10.27	0.24		99	
Actinomyces israelii		3.70	0.37		1	
07755510	**Low discrimination**					**07755510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium butyricum		48.61	0.43	ESL 88	99		
Bifidobacterium adolescentis		36.80	0.56	TRE 94	1		
Actinomyces israelii		12.88	0.54	MLZ 5	1		
07755512	**Very good identification**						**07755512**
Bifidobacterium adolescentis		95.72	0.90				
Clostridium clostridioforme		3.25	0.61				
Actinomyces israelii		0.51	0.57				
07755513	**Good identification**						**07755513**
Bifidobacterium adolescentis		95.75	0.56	URE 1			
Clostridium clostridioforme		3.25	0.27				
Actinomyces israelii		0.51	0.23				
07755520	**Low discrimination**						**07755520**
					GRA	SPO	
Clostridium butyricum		48.21	0.47	ESL 88	99	99	
Bacteroides vulgatus		46.32	0.45	TRE 94	1	1	
Actinomyces israelii		3.30	0.48	NAG 8	99	1	
07755522	**Low discrimination**						**07755522**
					GRA	SPO	
Parabacteroides distasonis		44.79	0.66		1	1	
Clostridium butyricum		13.00	0.40		99	99	
Bifidobacterium adolescentis		12.54	0.54		99	1	
07755532	**Low discrimination**						**07755532**
					GRA		
Clostridium clostridioforme		55.05	0.61	TRE 88	50		
Parabacteroides distasonis		25.81	0.61	RHA 88	1		
Bifidobacterium adolescentis		16.40	0.56		99		
07755540	**Low discrimination**						**07755540**
					SPO		
Clostridium butyricum		68.31	0.47	ESL 88	99		
Actinomyces israelii		28.52	0.61	TRE 94	1		
Bifidobacterium adolescentis		3.17	0.39	SOR 8	1		
07755542	**Low discrimination**						**07755542**
					SPO		
Bifidobacterium adolescentis		81.83	0.73	SOR 11	1		
Actinomyces israelii		11.17	0.64		1		
Clostridium butyricum		6.94	0.40		99		
07755543	**Low discrimination**						**07755543**
					SPO		
Bifidobacterium adolescentis		81.88	0.39	SOR 11	1		
Actinomyces israelii		11.18	0.30	URE 1	1		
Clostridium butyricum		6.94	0.05		99		
07755550	**Low discrimination**						**07755550**
					SPO		
Actinomyces israelii		36.10	0.47	MLZ 13	1		
Bifidobacterium adolescentis		32.77	0.40		1		
Clostridium butyricum		30.45	0.25		99		
07755552	**Very good identification**						**07755552**
					SPO		
Bifidobacterium adolescentis		97.74	0.74	SOR 11	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.63	0.50		1		
Clostridium butyricum		0.36	0.18		99		
07755553	**Good identification**						**07755553**
Bifidobacterium adolescentis		98.36	0.40	SOR 11			
Actinomyces israelii		1.64	0.16	URE 1			
07755562	**Low discrimination**						**07755562**
					SPO		
Bifidobacterium adolescentis		32.67	0.39	NAG 1	1		
Actinomyces israelii		28.18	0.44	SOR 11	1		
Clostridium butyricum		23.84	0.21		99		
07755572	**Low discrimination**						**07755572**
					GRA		
Bifidobacterium adolescentis		64.94	0.40	NAG 1	99		
Clostridium clostridioforme		17.82	0.27	SOR 11	50		
Parabacteroides distasonis		8.35	0.27		1		
07755600	**Low discrimination**						**07755600**
					SPO		
Actinomyces israelii		46.57	0.39	RHA 7	1		
Clostridium butyricum		34.17	0.15		99		
Clostridium clostridioforme		17.23	0.22		66		
07755602	**Acceptable identification**						**07755602**
					SPO		
Clostridium clostridioforme		94.89	0.56	TRE 88	66		
Actinomyces israelii		3.89	0.42		1		
Clostridium butyricum		0.74	0.08		99		
07755612	**Very good identification**						**07755612**
Clostridium clostridioforme		99.73	0.67	TRE 88			
Bifidobacterium adolescentis		0.14	0.21				
Actinomyces israelii		0.13	0.28				
07755622	**Acceptable identification**						**07755622**
Clostridium clostridioforme		92.74	0.56	TRE 88			
Parabacteroides distasonis		5.09	0.40				
Bacteroides uniformis		0.86	0.08				
07755632	**Very good identification**						**07755632**
Clostridium clostridioforme		99.29	0.67	TRE 88			
Parabacteroides distasonis		0.64	0.35				
Bacteroides ovatus		0.07	0.03				
07755700	**Low discrimination**						**07755700**
					GRA	SPO	
Clostridium butyricum		62.38	0.39	ESL 88	99	99	
Bacteroides vulgatus		15.32	0.27	TRE 94	1	1	
Actinomyces israelii		14.17	0.49	RHA 3	99	1	
07755702	**Low discrimination**						**07755702**
					SPO		
Clostridium clostridioforme		78.23	0.65	TRE 88	66		
Bifidobacterium adolescentis		11.84	0.54		1		
Clostridium butyricum		4.37	0.32		99		
07755710	**Low discrimination**						**07755710**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium clostridioforme		75.84	0.42	TRE 88	66		
Clostridium butyricum		12.46	0.17	BGL 99	99		
Actinomyces israelii		8.04	0.35		1		
07755712	**Very good identification**						**07755712**
					GRA		
Clostridium clostridioforme		95.69	0.76	TRE 88	50		
Bifidobacterium adolescentis		3.89	0.56		99		
Parabacteroides distasonis		0.19	0.35		1		
07755713	**Good identification**						**07755713**
Clostridium clostridioforme		96.10	0.42	TRE 88			
Bifidobacterium adolescentis		3.90	0.21	URE 1			
07755720	**Very good identification**						**07755720**
					GRA	SPO	
Bacteroides vulgatus		98.69	0.62	CEL 5	1	1	
Parabacteroides distasonis		0.46	0.40		1	1	
Clostridium butyricum		0.35	0.21		99	99	
07755722	**Low discrimination**						**07755722**
Parabacteroides distasonis		54.63	0.74				
Clostridium clostridioforme		33.67	0.65				
Bacteroides ovatus		4.55	0.40				
07755723	**Low discrimination**						**07755723**
Parabacteroides distasonis		54.91	0.40	URE 1			
Clostridium clostridioforme		33.84	0.31				
Bacteroides ovatus		4.57	0.06				
07755730	**Low discrimination**						**07755730**
Clostridium clostridioforme		50.03	0.42	TRE 88			
Bacteroides vulgatus		38.34	0.27	BGL 99			
Parabacteroides distasonis		9.59	0.35				
07755732	**Low discrimination**						**07755732**
Clostridium clostridioforme		82.44	0.76	TRE 88			
Parabacteroides distasonis		15.81	0.70				
Bacteroides ovatus		1.63	0.37				
07755733	**Low discrimination**						**07755733**
Clostridium clostridioforme		82.54	0.42	TRE 88			
Parabacteroides distasonis		15.83	0.35	URE 1			
Bacteroides ovatus		1.63	0.03				
07755740	**Low discrimination**						**07755740**
					GRA	SPO	
Actinomyces israelii		49.70	0.42	RHA 7	99	1	
Clostridium butyricum		48.91	0.21		99	99	
Bacteroides vulgatus		1.40	-.07		1	1	
07755742	**Low discrimination**						**07755742**
Actinomyces israelii		36.11	0.45	RHA 7			
Bifidobacterium adolescentis		35.51	0.39				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		19.17	0.31			
07755752	**Low discrimination**					**07755752**
Clostridium clostridioforme		64.15	0.42	TRE 88		
Bifidobacterium adolescentis		31.88	0.40	SOR 1		
Actinomyces israelii		3.97	0.31			
07755762	**Low discrimination**					**07755762**
Parabacteroides distasonis		45.84	0.40	SOR 1		
Clostridium clostridioforme		28.25	0.31			
Bacteroides ovatus		15.75	0.16			
07755772	**Low discrimination**					**07755772**
Clostridium clostridioforme		78.53	0.42	TRE 88		
Parabacteroides distasonis		15.06	0.35	SOR 1		
Bacteroides ovatus		6.41	0.13			
07756000	**Acceptable identification**					**07756000**
Actinomyces israelii		94.24	0.38	GAL 99		
Actinomyces naeslundii		5.76	0.27			
07756002	**Good identification**					**07756002**
Actinomyces israelii		96.09	0.41	GAL 99		
Actinomyces naeslundii		3.91	0.27			
07756100	**Acceptable identification**					**07756100**
					GRA	
Actinomyces israelii		92.38	0.48	GAL 99	99	
Actinomyces naeslundii		4.02	0.34		99	
Fusobacterium mortiferum		1.84	0.17		1	
07756101	**Low discrimination**					**07756101**
Actinomyces naeslundii		86.58	0.37	GAL 96		
Actinomyces israelii		13.42	0.14	XYL 1		
07756102	**Low discrimination**					**07756102**
Actinomyces israelii		69.64	0.51	GAL 99		
Bifidobacterium adolescentis		28.34	0.38			
Actinomyces naeslundii		2.02	0.34			
07756103	**Low discrimination**					**07756103**
Actinomyces naeslundii		81.14	0.37	GAL 96		
Actinomyces israelii		18.86	0.17	XYL 1		
07756112	**Low discrimination**					**07756112**
					SPO	
Bifidobacterium adolescentis		76.19	0.40	GAL 99	1	
Actinomyces israelii		22.89	0.37	ARA 87	1	
Clostridium tertium		0.92	-.01		99	
07756121	**Good identification**					**07756121**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07756123	**Good identification**					**07756123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GAL 96		
07756140	**Good identification**					**07756140**
Actinomyces israelii		100.0	0.41	GAL 99		
07756142	**Low discrimination**					**07756142**
Actinomyces israelii		88.55	0.44	GAL 99		
Bifidobacterium adolescentis		11.45	0.23			
07756400	**Good identification**					**07756400**
Actinomyces israelii		100.0	0.37	GAL 99		
07756402	**Low discrimination**					**07756402**
Actinomyces israelii		88.15	0.40	GAL 99		
Bifidobacterium adolescentis		11.85	0.18			
07756500	**Low discrimination**					**07756500**
					SPO	
Actinomyces israelii		79.34	0.47	GAL 99	1	
Clostridium butyricum		16.66	0.14		99	
Bifidobacterium adolescentis		4.00	0.18		1	
07756502	**Low discrimination**					**07756502**
					SPO	
Bifidobacterium adolescentis		75.73	0.53	GAL 99	1	
Actinomyces israelii		22.75	0.50		1	
Clostridium butyricum		1.24	0.07		99	
07756512	**Acceptable identification**					**07756512**
Bifidobacterium adolescentis		94.90	0.54	GAL 99		
Actinomyces israelii		3.49	0.36			
Clostridium clostridioforme		1.31	0.19			
07756540	**Good identification**					**07756540**
					SPO	
Actinomyces israelii		95.52	0.40	GAL 99	1	
Clostridium butyricum		4.49	-.04		99	
07756542	**Low discrimination**					**07756542**
Bifidobacterium adolescentis		51.40	0.37	GAL 99		
Actinomyces israelii		48.60	0.43	SOR 11		
07756552	**Low discrimination**					**07756552**
Bifidobacterium adolescentis		89.64	0.39	GAL 99		
Actinomyces israelii		10.36	0.29	SOR 11		
07757000	**Very good identification**					**07757000**
					SPO	
Actinomyces israelii		98.02	0.72		1	
Actinomyces naeslundii		1.45	0.50		1	
Clostridium butyricum		0.31	0.08		99	
07757001	**Low discrimination**					**07757001**
Actinomyces naeslundii		68.74	0.53	XYL 1		
Actinomyces israelii		31.26	0.38			
07757002	**Very good identification**					**07757002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		97.20	0.75			
Bifidobacterium adolescentis		1.60	0.38			
Actinomyces naeslundii		0.96	0.50			
07757003	**Low discrimination**					**07757003**
Actinomyces naeslundii		59.45	0.53	XYL 1		
Actinomyces israelii		40.55	0.41			
07757010	**Very good identification**					**07757010**
Actinomyces israelii		99.48	0.58	MLZ 13		
Actinomyces naeslundii		0.52	0.28			
07757012	**Low discrimination**					**07757012**
Actinomyces israelii		87.75	0.61	MLZ 13		
Bifidobacterium adolescentis		11.80	0.40			
Actinomyces naeslundii		0.31	0.28			
07757020	**Low discrimination**					**07757020**
Actinomyces israelii		81.17	0.52	NAG 6		
Actinomyces naeslundii		18.83	0.50			
07757021	**Good identification**					**07757021**
Actinomyces naeslundii		97.18	0.53	XYL 1		
Actinomyces israelii		2.82	0.18			
07757022	**Low discrimination**					**07757022**
					GRA	
Actinomyces israelii		80.87	0.55	NAG 6	99	
Actinomyces naeslundii		12.51	0.50		99	
Prevotella oralis		4.57	0.17		1	
07757023	**Good identification**					**07757023**
Actinomyces naeslundii		95.83	0.53	XYL 1		
Actinomyces israelii		4.17	0.21			
07757030	**Low discrimination**					**07757030**
					SPO	
Actinomyces israelii		89.29	0.38	MLZ 13	1	
Actinomyces naeslundii		7.30	0.28	NAG 6	1	
Clostridium tertium		3.41	-.01		99	
07757032	**Low discrimination**					**07757032**
					SPO	
Actinomyces israelii		88.43	0.41	MLZ 13	1	
Clostridium tertium		6.76	0.07	NAG 6	99	
Actinomyces naeslundii		4.82	0.28		1	
07757040	**Very good identification**					**07757040**
Actinomyces israelii		99.33	0.65			
Actinomyces naeslundii		0.67	0.37			
07757042	**Very good identification**					**07757042**
Actinomyces israelii		99.04	0.68			
Bifidobacterium adolescentis		0.52	0.23			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.44	0.37			
07757050	**Good identification**					**07757050**
Actinomyces israelii		100.0	0.51	MLZ 13		
07757052	**Good identification**					**07757052**
Actinomyces israelii		95.90	0.54	MLZ 13		
Bifidobacterium adolescentis		4.10	0.24			
07757060	**Acceptable identification**					**07757060**
Actinomyces israelii		90.48	0.45	NAG 6		
Actinomyces naeslundii		9.52	0.37			
07757061	**Good identification**					**07757061**
Actinomyces naeslundii		100.0	0.40	XYL 1		
07757062	**Acceptable identification**					**07757062**
Actinomyces israelii		93.44	0.48	NAG 6		
Actinomyces naeslundii		6.56	0.37			
07757063	**Acceptable identification**					**07757063**
Actinomyces naeslundii		91.25	0.40	XYL 1		
Actinomyces israelii		8.76	0.14			
07757100	**Very good identification**					**07757100**
					SPO	
Actinomyces israelii		96.35	0.83		1	
Clostridium butyricum		1.84	0.32		99	
Actinomyces naeslundii		1.02	0.58		1	
07757101	**Low discrimination**					**07757101**
					SPO	
Actinomyces naeslundii		60.56	0.61	XYL 1	1	
Actinomyces israelii		38.71	0.48		1	
Clostridium butyricum		0.74	-.02		99	
07757102	**Low discrimination**					**07757102**
Actinomyces israelii		70.15	0.86			
Bifidobacterium adolescentis		28.54	0.73			
Actinomyces naeslundii		0.49	0.58			
07757103	**Low discrimination**					**07757103**
Actinomyces naeslundii		42.57	0.61	XYL 1		
Actinomyces israelii		40.82	0.51			
Bifidobacterium adolescentis		16.61	0.38			
07757110	**Good identification**					**07757110**
					SPO	
Actinomyces israelii		92.44	0.68	MLZ 13	1	
Bifidobacterium adolescentis		4.66	0.40		1	
Clostridium tertium		1.86	0.25		99	
07757112	**Low discrimination**					**07757112**
					SPO	
Bifidobacterium adolescentis		76.00	0.74	ARA 87	1	
Actinomyces israelii		22.83	0.71		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		0.92	0.33		99		
07757113	**Low discrimination**						**07757113**
Bifidobacterium adolescentis		70.28	0.40	ARA 87			
Actinomyces israelii		21.12	0.37	URE 1			
Actinomyces naeslundii		7.76	0.39				
07757120	**Low discrimination**						**07757120**
					SPO		
Actinomyces israelii		72.13	0.62	NAG 6	1		
Actinomyces naeslundii		11.91	0.58		1		
Clostridium ramosum		6.74	0.28		92		
07757121	**Very good identification**						**07757121**
Actinomyces naeslundii		96.08	0.61	XYL 1			
Actinomyces israelii		3.92	0.28				
07757122	**Low discrimination**						**07757122**
					SPO	GRA	
Actinomyces israelii		52.78	0.65	NAG 6	1	99	
Clostridium ramosum		13.15	0.39		92	50	
Prevotella oralis		11.69	0.37		1	1	
07757123	**Good identification**						**07757123**
					SPO		
Actinomyces naeslundii		90.93	0.61	XYL 1	1		
Actinomyces israelii		5.57	0.31		1		
Clostridium ramosum		1.39	0.05		92		
07757130	**Low discrimination**						**07757130**
					SPO		
Actinomyces israelii		49.36	0.48	MLZ 13	1		
Clostridium tertium		46.65	0.33	NAG 6	99		
Actinomyces naeslundii		2.87	0.36		1		
07757131	**Low discrimination**						**07757131**
					SPO		
Actinomyces naeslundii		81.61	0.39	XYL 1	1		
Actinomyces israelii		9.45	0.14	MLZ 5	1		
Clostridium tertium		8.94	-.01		99		
07757132	**Low discrimination**						**07757132**
					SPO		
Clostridium tertium		49.99	0.41	RAF 1	99		
Actinomyces israelii		26.44	0.51	TRE 90	1		
Bifidobacterium adolescentis		13.93	0.40		1		
07757133	**Low discrimination**						**07757133**
					SPO		
Actinomyces naeslundii		66.57	0.39	XYL 1	1		
Clostridium tertium		21.87	0.07	MLZ 5	99		
Actinomyces israelii		11.57	0.17		1		
07757140	**Very good identification**						**07757140**
					SPO		
Actinomyces israelii		98.88	0.76		1		
Actinomyces naeslundii		0.47	0.45		1		
Clostridium butyricum		0.42	0.14		99		
07757141	**Low discrimination**						**07757141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		58.48	0.41	URE 1		
Actinomyces naeslundii		41.52	0.48			
07757142	**Low discrimination**					**07757142**
Actinomyces israelii		88.01	0.79			
Bifidobacterium adolescentis		11.38	0.57			
Actinomyces naeslundii		0.28	0.45			
07757143	**Low discrimination**					**07757143**
Actinomyces israelii		62.40	0.44	URE 1		
Actinomyces naeslundii		29.53	0.48			
Bifidobacterium adolescentis		8.07	0.23			
07757150	**Very good identification**					**07757150**
					COC	
Actinomyces israelii		97.80	0.61	MLZ 13	1	
Bifidobacterium adolescentis		1.57	0.24		1	
Blautia producta		0.27	-.03		99	
07757152	**Low discrimination**					**07757152**
Bifidobacterium adolescentis		51.00	0.59	ARA 87		
Actinomyces israelii		48.21	0.64	SOR 11		
Bifidobacterium breve		0.68	0.27			
07757160	**Acceptable identification**					**07757160**
Actinomyces israelii		91.79	0.55	NAG 6		
Actinomyces naeslundii		6.88	0.45			
Bifidobacterium breve		0.80	0.15			
07757161	**Acceptable identification**					**07757161**
Actinomyces naeslundii		91.75	0.48	XYL 1		
Actinomyces israelii		8.25	0.21			
07757162	**Low discrimination**					**07757162**
Actinomyces israelii		86.67	0.58	NAG 6		
Bifidobacterium breve		5.11	0.32			
Actinomyces naeslundii		4.33	0.45			
07757163	**Low discrimination**					**07757163**
Actinomyces naeslundii		88.12	0.48	XYL 1		
Actinomyces israelii		11.88	0.24			
07757170	**Acceptable identification**					**07757170**
Actinomyces israelii		90.14	0.41	MLZ 13		
Bifidobacterium breve		5.27	0.15	NAG 6		
Actinomyces naeslundii		2.38	0.23			
07757172	**Low discrimination**					**07757172**
Actinomyces israelii		61.41	0.44	MLZ 13		
Bifidobacterium breve		24.21	0.32	NAG 6		
Bifidobacterium adolescentis		10.28	0.24			
07757200	**Good identification**					**07757200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.53	RHA 7			
07757202	**Good identification**						**07757202**
Actinomyces israelii		100.0	0.56	RHA 7			
07757210	**Good identification**						**07757210**
Actinomyces israelii		100.0	0.39	RHA 7			
07757212	**Acceptable identification**						**07757212**
Actinomyces israelii		93.52	0.42	RHA 7			
Clostridium clostridioforme		6.48	0.13	MLZ 13			
07757222	**Low discrimination**						**07757222**
					SPO	GRA	
Actinomyces israelii		64.55	0.36	RHA 7	1	99	
Prevotella oralis		20.78	0.11	NAG 6	1	1	
Clostridium ramosum		11.45	0.07		92	50	
07757240	**Good identification**						**07757240**
Actinomyces israelii		100.0	0.46	RHA 7			
07757242	**Good identification**						**07757242**
Actinomyces israelii		100.0	0.49	RHA 7			
07757300	**Very good identification**						**07757300**
					SPO		
Actinomyces israelii		99.08	0.63	RHA 7	1		
Clostridium butyricum		0.78	0.06		99		
Actinomyces naeslundii		0.14	0.24		1		
07757302	**Good identification**						**07757302**
					GRA		
Actinomyces israelii		93.55	0.66	RHA 7	99		
Bifidobacterium adolescentis		5.11	0.38		99		
Prevotella oralis		0.40	0.09		1		
07757310	**Good identification**						**07757310**
Actinomyces israelii		100.0	0.49	RHA 7			
07757312	**Low discrimination**						**07757312**
Actinomyces israelii		66.22	0.52	RHA 7			
Bifidobacterium adolescentis		29.58	0.40	MLZ 13			
Clostridium clostridioforme		3.84	0.22				
07757322	**Low discrimination**						**07757322**
					SPO		
Clostridium ramosum		54.58	0.41	TRE 94	92		
Prevotella oralis		15.68	0.31	XYL 1	1		
Parabacteroides distasonis		15.37	0.44		1		
07757332	**Low discrimination**						**07757332**
					GRA		
Parabacteroides distasonis		61.14	0.39	MAN 1	1		
Actinomyces israelii		13.73	0.32		99		
Clostridium clostridioforme		12.47	0.22		50		
07757340	**Very good identification**						**07757340**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.56	RHA 7			
07757342	**Very good identification**						**07757342**
Actinomyces israelii		98.29	0.59	RHA 7			
Bifidobacterium adolescentis		1.71	0.23				
07757350	**Good identification**						**07757350**
					COC		
Actinomyces israelii		96.53	0.42	RHA 7	1		
Blautia producta		3.47	-.03	MLZ 13	99		
07757352	**Low discrimination**						**07757352**
Actinomyces israelii		87.57	0.45	RHA 7			
Bifidobacterium adolescentis		12.43	0.24	MLZ 13			
07757360	**Good identification**						**07757360**
Actinomyces israelii		100.0	0.36	RHA 7			
07757362	**Low discrimination**						**07757362**
					SPO	GRA	
Actinomyces israelii		84.83	0.39	RHA 7	1	99	
Clostridium ramosum		9.67	0.07	NAG 6	92	50	
Prevotella oralis		2.78	-.03		1	1	
07757400	**Very good identification**						**07757400**
					SPO		
Actinomyces israelii		96.43	0.71		1		
Clostridium butyricum		3.38	0.25		99		
Bifidobacterium adolescentis		0.20	0.18		1		
07757401	**Good identification**						**07757401**
Actinomyces israelii		100.0	0.37	URE 1			
07757402	**Low discrimination**						**07757402**
					SPO		
Actinomyces israelii		87.41	0.74		1		
Bifidobacterium adolescentis		11.76	0.53		1		
Clostridium butyricum		0.79	0.18		99		
07757403	**Low discrimination**						**07757403**
Actinomyces israelii		88.15	0.40	URE 1			
Bifidobacterium adolescentis		11.85	0.18				
07757410	**Very good identification**						**07757410**
					SPO		
Actinomyces israelii		97.18	0.57	MLZ 13	1		
Bifidobacterium adolescentis		1.62	0.20		1		
Clostridium butyricum		1.20	0.03		99		
07757412	**Low discrimination**						**07757412**
Bifidobacterium adolescentis		51.88	0.54	LAC 99			
Actinomyces israelii		47.17	0.60				
Clostridium clostridioforme		0.64	0.18				
07757420	**Good identification**						**07757420**
					SPO		
Actinomyces israelii		95.45	0.50	NAG 6	1		
Clostridium butyricum		4.55	0.07		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07757422	**Good identification**					**07757422**
					SPO	
Actinomyces israelii		96.10	0.53	NAG 6	1	
Bifidobacterium adolescentis		2.05	0.18		1	
Clostridium butyricum		1.19	0.00		99	
07757430	**Good identification**					**07757430**
Actinomyces israelii		100.0	0.36	MLZ 13		
07757432	**Low discrimination**					**07757432**
Actinomyces israelii		67.37	0.39	MLZ 13		
Clostridium clostridioforme		14.37	0.18	NAG 6		
Bifidobacterium adolescentis		11.73	0.20			
07757440	**Very good identification**					**07757440**
					SPO	
Actinomyces israelii		99.22	0.64		1	
Clostridium butyricum		0.78	0.07		99	
07757442	**Very good identification**					**07757442**
					SPO	
Actinomyces israelii		95.72	0.67		1	
Bifidobacterium adolescentis		4.09	0.37		1	
Clostridium butyricum		0.19	0.00		99	
07757450	**Good identification**					**07757450**
					COC	
Actinomyces israelii		98.69	0.50	MLZ 13	1	
Blautia producta		1.31	-.03		99	
07757452	**Low discrimination**					**07757452**
Actinomyces israelii		74.10	0.53	MLZ 13		
Bifidobacterium adolescentis		25.90	0.39			
07757460	**Good identification**					**07757460**
Actinomyces israelii		100.0	0.43	NAG 6		
07757462	**Good identification**					**07757462**
Actinomyces israelii		100.0	0.46	NAG 6		
07757500	**Low discrimination**					**07757500**
					SPO	
Actinomyces israelii		79.33	0.81		1	
Clostridium butyricum		16.66	0.48		99	
Bifidobacterium adolescentis		4.00	0.53		1	
07757501	**Low discrimination**					**07757501**
					SPO	
Actinomyces israelii		78.14	0.47	URE 1	1	
Clostridium butyricum		16.41	0.14		99	
Bifidobacterium adolescentis		3.94	0.18		1	
07757502	**Low discrimination**					**07757502**
					SPO	
Bifidobacterium adolescentis		75.93	0.87		1	
Actinomyces israelii		22.81	0.84		1	
Clostridium butyricum		1.24	0.41		99	
07757503	**Low discrimination**					**07757503**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium adolescentis		75.72	0.53	URE 1	1		
Actinomyces israelii		22.75	0.50		1		
Clostridium butyricum		1.24	0.07		99		
07757510	**Low discrimination**						**07757510**
					SPO		
Actinomyces israelii		66.64	0.67	MLZ 13	1		
Bifidobacterium adolescentis		27.49	0.54		1		
Clostridium butyricum		4.93	0.27		99		
07757512	**Very good identification**						**07757512**
Bifidobacterium adolescentis		96.06	0.88				
Actinomyces israelii		3.53	0.70				
Bifidobacterium dentium		0.30	0.27				
07757513	**Good identification**						**07757513**
Bifidobacterium adolescentis		96.16	0.54	URE 1			
Actinomyces israelii		3.53	0.36				
Bifidobacterium dentium		0.30	-.07				
07757520	**Low discrimination**						**07757520**
					GRA	SPO	
Actinomyces israelii		75.69	0.61	NAG 6	99	1	
Clostridium butyricum		21.66	0.30		99	99	
Bacteroides vulgatus		1.89	0.11		1	1	
07757522	**Low discrimination**						**07757522**
					SPO		
Actinomyces israelii		60.35	0.64	NAG 6	1		
Bifidobacterium adolescentis		31.79	0.53		1		
Clostridium butyricum		4.48	0.23		99		
07757530	**Low discrimination**						**07757530**
					SPO		
Actinomyces israelii		65.20	0.47	MLZ 13	1		
Bifidobacterium dentium		23.22	0.15	NAG 6	1		
Clostridium butyricum		6.57	0.08		99		
07757532	**Low discrimination**						**07757532**
Bifidobacterium adolescentis		53.42	0.54	NAG 1			
Bifidobacterium dentium		29.76	0.32				
Actinomyces israelii		12.40	0.50				
07757540	**Good identification**						**07757540**
					COC	SPO	
Actinomyces israelii		92.65	0.74		1	1	
Clostridium butyricum		4.35	0.30		1	99	
Blautia producta		1.51	0.24		99	1	
07757541	**Good identification**						**07757541**
					SPO		
Actinomyces israelii		95.52	0.40	URE 1	1		
Clostridium butyricum		4.49	-.04		99		
07757542	**Low discrimination**						**07757542**
					SPO		
Bifidobacterium adolescentis		51.07	0.71	SOR 11	1		
Actinomyces israelii		48.28	0.77		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.59	0.23		99		
07757543	**Low discrimination**						**07757543**
Bifidobacterium adolescentis		51.40	0.37	SOR 11			
Actinomyces israelii		48.60	0.43	URE 1			
07757550	**Low discrimination**						**07757550**
					COC		
Actinomyces israelii		67.78	0.60	MLZ 13	1		
Blautia producta		22.21	0.32		99		
Bifidobacterium adolescentis		8.89	0.39		1		
07757552	**Low discrimination**						**07757552**
					COC		
Bifidobacterium adolescentis		89.35	0.73	SOR 11	1		
Actinomyces israelii		10.33	0.63		1		
Blautia producta		0.25	0.15		99		
07757553	**Low discrimination**						**07757553**
Bifidobacterium adolescentis		89.64	0.39	SOR 11			
Actinomyces israelii		10.36	0.29	URE 1			
07757560	**Acceptable identification**						**07757560**
					COC	SPO	
Actinomyces israelii		91.54	0.54	NAG 6	1	1	
Clostridium butyricum		5.86	0.12		1	99	
Blautia producta		2.60	0.07		99	1	
07757562	**Low discrimination**						**07757562**
					SPO		
Actinomyces israelii		84.46	0.57	NAG 6	1		
Bifidobacterium adolescentis		14.14	0.37		1		
Clostridium butyricum		1.40	0.05		99		
07757570	**Low discrimination**						**07757570**
					COC		
Actinomyces israelii		63.68	0.39	MLZ 13	1		
Blautia producta		36.32	0.15	NAG 6	99		
07757572	**Low discrimination**						**07757572**
Bifidobacterium adolescentis		55.35	0.39	NAG 1			
Actinomyces israelii		40.42	0.42	SOR 11			
Bifidobacterium dentium		2.52	-.03				
07757600	**Good identification**						**07757600**
					SPO		
Actinomyces israelii		98.58	0.52	RHA 7	1		
Clostridium butyricum		1.42	-.01		99		
07757602	**Acceptable identification**						**07757602**
Actinomyces israelii		94.20	0.55	RHA 7			
Clostridium clostridioforme		4.10	0.22				
Bifidobacterium adolescentis		1.70	0.18				
07757610	**Good identification**						**07757610**
Actinomyces israelii		100.0	0.37	RHA 7			
07757640	**Good identification**						**07757640**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.45	RHA 7			
07757642	**Good identification**						**07757642**
Actinomyces israelii		100.0	0.48	RHA 7			
07757700	**Good identification**						**07757700**
					SPO		
Actinomyces israelii		91.33	0.62	RHA 7	1		
Clostridium butyricum		7.88	0.23		99		
Bifidobacterium adolescentis		0.62	0.18		1		
07757702	**Low discrimination**						**07757702**
Actinomyces israelii		66.35	0.65	RHA 7			
Bifidobacterium adolescentis		29.64	0.53				
Clostridium clostridioforme		2.42	0.31				
07757710	**Low discrimination**						**07757710**
					SPO		
Actinomyces israelii		89.31	0.48	RHA 7	1		
Bifidobacterium adolescentis		4.94	0.20	MLZ 13	1		
Clostridium butyricum		2.72	0.01		99		
07757712	**Low discrimination**						**07757712**
Bifidobacterium adolescentis		63.09	0.54	RHA 1			
Clostridium clostridioforme		19.18	0.42				
Actinomyces israelii		17.27	0.51				
07757732	**Low discrimination**						**07757732**
Clostridium clostridioforme		51.19	0.42	TRE 88			
Bacteroides ovatus		33.40	0.29	MAN 1			
Parabacteroides distasonis		9.82	0.35				
07757740	**Low discrimination**						**07757740**
					COC	SPO	
Actinomyces israelii		80.90	0.55	RHA 7	1	1	
Blautia producta		17.54	0.24		99	1	
Clostridium butyricum		1.56	0.04		1	99	
07757742	**Low discrimination**						**07757742**
					COC		
Actinomyces israelii		85.98	0.58	RHA 7	1		
Bifidobacterium adolescentis		12.21	0.37		1		
Blautia producta		1.38	0.07		99		
07757752	**Low discrimination**						**07757752**
					COC		
Bifidobacterium adolescentis		46.21	0.39	RHA 1	1		
Actinomyces israelii		39.80	0.44	SOR 11	1		
Blautia producta		12.84	0.15		99		
07757762	**Low discrimination**						**07757762**
					GRA		
Actinomyces israelii		78.58	0.37	RHA 7	99		
Bacteroides ovatus		21.42	0.08	NAG 6	1		
07760000	**Low discrimination**						**07760000**
					GRA		
Actinomyces naeslundii		70.16	0.39	FRU 95	99		
Fusobacterium mortiferum		29.84	0.21	GAL 96	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07760001	**Good identification**					**07760001**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07760002	**Low discrimination**					**07760002**
Actinomyces naeslundii		75.41	0.39	FRU 95		
Lactobacillus acidophilus		12.91	0.01	GAL 96		
Actinomyces viscosus		11.68	0.01			
07760003	**Good identification**					**07760003**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07760020	**Good identification**					**07760020**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07760021	**Good identification**					**07760021**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07760022	**Low discrimination**					**07760022**
Actinomyces naeslundii		85.39	0.39	FRU 95		
Lactobacillus acidophilus		14.61	0.01	GAL 96		
07760023	**Good identification**					**07760023**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07760100	**Low discrimination**					**07760100**
					GRA	
Actinomyces naeslundii		74.15	0.47	FRU 95	99	
Fusobacterium mortiferum		25.85	0.27	GAL 96	1	
07760101	**Good identification**					**07760101**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07760102	**Low discrimination**					**07760102**
Actinomyces naeslundii		54.74	0.47	FRU 95		
Lactobacillus acidophilus		37.85	0.19	GAL 96		
Actinomyces viscosus		5.30	0.05			
07760103	**Good identification**					**07760103**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07760120	**Acceptable identification**					**07760120**
					SPO	
Actinomyces naeslundii		93.21	0.47	FRU 95	1	
Clostridium ramosum		6.79	0.02	GAL 96	92	
07760121	**Good identification**					**07760121**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07760122	**Low discrimination**					**07760122**
					SPO	
Actinomyces naeslundii		49.23	0.47	FRU 95	1	
Lactobacillus acidophilus		34.04	0.19	GAL 96	1	
Clostridium ramosum		14.34	0.12		92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07760123	**Good identification**					**07760123**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07760141	**Good identification**					**07760141**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07760143	**Good identification**					**07760143**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07760161	**Good identification**					**07760161**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07760163	**Good identification**					**07760163**
Actinomyces naeslundii		100.0	0.37	FRU 95		
07761000	**Low discrimination**					**07761000**
					GRA	
Actinomyces naeslundii		75.04	0.62	FRU 95	99	
Lactobacillus jensenii		21.86	0.32		99	
Fusobacterium mortiferum		1.99	0.24		1	
07761001	**Very good identification**					**07761001**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07761002	**Low discrimination**					**07761002**
Actinomyces naeslundii		52.32	0.62	FRU 95		
Lactobacillus acidophilus		36.93	0.35			
Lactobacillus jensenii		10.16	0.29			
07761003	**Very good identification**					**07761003**
Actinomyces naeslundii		99.53	0.65	FRU 95		
Lactobacillus acidophilus		0.47	0.01			
07761011	**Good identification**					**07761011**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07761013	**Good identification**					**07761013**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07761020	**Good identification**					**07761020**
Actinomyces naeslundii		91.14	0.62	FRU 95		
Lactobacillus jensenii		6.64	0.21			
Clostridium perfringens		0.79	0.05			
07761021	**Very good identification**					**07761021**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07761022	**Low discrimination**					**07761022**
Actinomyces naeslundii		54.79	0.62	FRU 95		
Lactobacillus acidophilus		38.67	0.35			
Lactobacillus jensenii		2.66	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07761023	**Very good identification**					**07761023**
Actinomyces naeslundii		99.53	0.65	FRU 95		
Lactobacillus acidophilus		0.47	0.01			
07761030	**Low discrimination**					**07761030**
Actinomyces naeslundii		68.43	0.41	FRU 95		
Lactobacillus jensenii		31.57	0.13	MLZ 5		
07761031	**Good identification**					**07761031**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07761032	**Low discrimination**					**07761032**
Actinomyces naeslundii		69.30	0.41	FRU 95		
Lactobacillus jensenii		21.31	0.10	MLZ 5		
Lactobacillus acidophilus		9.39	0.01			
07761033	**Good identification**					**07761033**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07761040	**Good identification**					**07761040**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07761041	**Good identification**					**07761041**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07761042	**Low discrimination**					**07761042**
Actinomyces naeslundii		66.92	0.50	FRU 95		
Lactobacillus acidophilus		33.08	0.20			
07761043	**Good identification**					**07761043**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07761060	**Good identification**					**07761060**
Actinomyces naeslundii		100.0	0.50	FRU 95		
07761061	**Good identification**					**07761061**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07761062	**Low discrimination**					**07761062**
Actinomyces naeslundii		66.92	0.50	FRU 95		
Lactobacillus acidophilus		33.08	0.20			
07761063	**Good identification**					**07761063**
Actinomyces naeslundii		100.0	0.53	FRU 95		
07761100	**Good identification**					**07761100**
					COC	
Actinomyces naeslundii		90.44	0.70	FRU 95	1	
Lactobacillus acidophilus		2.61	0.19		1	
Atopobium parvulum		2.36	0.20		99	
07761101	**Very good identification**					**07761101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.73	FRU 95		
07761102	**Low discrimination**					**07761102**
Lactobacillus acidophilus		72.45	0.54	FRU 99		
Actinomyces naeslundii		25.40	0.70			
Actinomyces israelii		0.41	0.30			
07761103	**Very good identification**					**07761103**
Actinomyces naeslundii		98.12	0.73	FRU 95		
Lactobacillus acidophilus		1.88	0.19			
07761110	**Good identification**					**07761110**
					GRA	
Actinomyces naeslundii		96.53	0.48	FRU 95	99	
Leptotrichia buccalis		3.47	-.11	MLZ 5	1	
07761111	**Good identification**					**07761111**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07761113	**Good identification**					**07761113**
					GRA	
Actinomyces naeslundii		97.66	0.51	FRU 95	99	
Leptotrichia buccalis		2.34	-.11	MLZ 5	1	
07761120	**Low discrimination**					**07761120**
					SPO	
Actinomyces naeslundii		72.41	0.70	FRU 95	1	
Clostridium ramosum		21.75	0.36		92	
Lactobacillus acidophilus		2.09	0.19		1	
07761121	**Very good identification**					**07761121**
					SPO	
Actinomyces naeslundii		99.80	0.73	FRU 95	1	
Clostridium ramosum		0.20	0.02		92	
07761122	**Low discrimination**					**07761122**
					SPO	
Lactobacillus acidophilus		53.57	0.54	FRU 99	1	
Clostridium ramosum		22.57	0.46		92	
Actinomyces naeslundii		18.78	0.70		1	
07761123	**Very good identification**					**07761123**
					SPO	
Actinomyces naeslundii		97.22	0.73	FRU 95	1	
Lactobacillus acidophilus		1.87	0.19		1	
Clostridium ramosum		0.79	0.12		92	
07761130	**Acceptable identification**					**07761130**
					SPO	
Actinomyces naeslundii		90.28	0.48	FRU 95	1	
Clostridium ramosum		5.21	0.02	MLZ 5	92	
Bifidobacterium breve		4.52	0.12		1	
07761131	**Good identification**					**07761131**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07761132	**Low discrimination**					**07761132**
Actinomyces naeslundii		32.99	0.48	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		18.06	0.19	MLZ 5		
Bifidobacterium breve		16.70	0.29			
07761133	**Good identification**					**07761133**
Actinomyces naeslundii		100.0	0.51	FRU 95		
07761140	**Very good identification**					**07761140**
Actinomyces naeslundii		95.82	0.57	FRU 95		
Actinomyces israelii		2.24	0.20			
Lactobacillus acidophilus		1.93	0.04			
07761141	**Very good identification**					**07761141**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07761142	**Low discrimination**					**07761142**
Lactobacillus acidophilus		63.88	0.38	FRU 99		
Actinomyces naeslundii		31.98	0.57	SOR 11		
Bifidobacterium breve		2.72	0.24			
07761143	**Very good identification**					**07761143**
Actinomyces naeslundii		98.67	0.60	FRU 95		
Lactobacillus acidophilus		1.33	0.04			
07761150	**Good identification**					**07761150**
Actinomyces naeslundii		100.0	0.35	FRU 95		
07761151	**Good identification**					**07761151**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07761153	**Good identification**					**07761153**
Actinomyces naeslundii		100.0	0.38	FRU 95		
07761160	**Good identification**					**07761160**
Actinomyces naeslundii		92.91	0.57	FRU 95		
Clostridium perfringens		2.23	0.07			
Lactobacillus acidophilus		1.87	0.04			
07761161	**Very good identification**					**07761161**
Actinomyces naeslundii		100.0	0.60	FRU 95		
07761162	**Low discrimination**					**07761162**
Lactobacillus acidophilus		61.65	0.38	FRU 99		
Actinomyces naeslundii		30.86	0.57	SOR 11		
Bifidobacterium breve		4.66	0.29			
07761163	**Very good identification**					**07761163**
Actinomyces naeslundii		98.67	0.60	FRU 95		
Lactobacillus acidophilus		1.33	0.04			
07761170	**Low discrimination**					**07761170**
Actinomyces naeslundii		77.90	0.35	FRU 95		
Bifidobacterium breve		22.10	0.12	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07761171	**Good identification**						**07761171**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07761173	**Good identification**						**07761173**
Actinomyces naeslundii		100.0	0.38	FRU 95			
07761300	**Low discrimination**						**07761300**
					SPO		
Actinomyces naeslundii		62.00	0.36	FRU 95	1		
Clostridium ramosum		24.69	0.04	RHA 1	92		
Lactobacillus acidophilus		13.31	0.00		1		
07761301	**Good identification**						**07761301**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07761303	**Low discrimination**						**07761303**
Actinomyces naeslundii		87.48	0.39	FRU 95			
Lactobacillus acidophilus		12.52	0.00	RHA 1			
07761320	**Good identification**						**07761320**
					SPO	GRA	
Clostridium ramosum		95.96	0.38	FRU 99	92	50	
Actinomyces naeslundii		2.43	0.36		1	99	
Prevotella oralis		1.08	0.04		1	1	
07761321	**Low discrimination**						**07761321**
					SPO		
Actinomyces naeslundii		79.02	0.39	FRU 95	1		
Clostridium ramosum		20.98	0.04	RHA 1	92		
07761322	**Low discrimination**						**07761322**
					SPO	GRA	
Clostridium ramosum		80.12	0.48	FRU 99	92	50	
Lactobacillus acidophilus		10.80	0.34		1	99	
Prevotella oralis		4.30	0.26		1	1	
07761501	**Good identification**						**07761501**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07761502	**Low discrimination**						**07761502**
					SPO		
Bifidobacterium adolescentis		45.05	0.37	FRU 99	1		
Clostridium butyricum		26.40	0.18	XYL 87	99		
Lactobacillus acidophilus		14.18	0.19		1		
07761503	**Good identification**						**07761503**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07761512	**Low discrimination**						**07761512**
					SPO		
Bifidobacterium adolescentis		76.54	0.38	FRU 99	1		
Clostridium clostridioforme		18.90	0.25	XYL 87	66		
Clostridium butyricum		1.93	-.04		99		
07761521	**Good identification**						**07761521**
Actinomyces naeslundii		100.0	0.39	FRU 95			
07761523	**Good identification**						**07761523**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	FRU 95		
07761712	**Good identification**					**07761712**
Clostridium clostridioforme		100.0	0.40	FRU 99		
07761722	**Low discrimination**					**07761722**
					SPO	
Parabacteroides distasonis		44.53	0.35	FRU 99	1	
Clostridium clostridioforme		39.32	0.28	XYL 90	66	
Clostridium ramosum		14.23	0.14		92	
07761732	**Low discrimination**					**07761732**
Clostridium clostridioforme		88.20	0.40	FRU 99		
Parabacteroides distasonis		11.81	0.30	XYL 91		
07762000	**Good identification**					**07762000**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07762001	**Good identification**					**07762001**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07762002	**Good identification**					**07762002**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07762003	**Good identification**					**07762003**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07762020	**Good identification**					**07762020**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07762021	**Good identification**					**07762021**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07762022	**Good identification**					**07762022**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07762023	**Good identification**					**07762023**
Actinomyces naeslundii		100.0	0.47	FRU 95		
07762100	**Good identification**					**07762100**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07762101	**Good identification**					**07762101**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07762102	**Good identification**					**07762102**
Actinomyces naeslundii		100.0	0.52	FRU 95		
07762103	**Good identification**					**07762103**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07762120	**Low discrimination**					**07762120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		82.47	0.52	FRU 95	1	
Clostridium ramosum		17.53	0.15	GAL 96	92	
07762121	**Good identification**					**07762121**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07762122	**Low discrimination**					**07762122**
					SPO	
Actinomyces naeslundii		54.04	0.52	FRU 95	1	
Clostridium ramosum		45.96	0.25	GAL 96	92	
07762123	**Good identification**					**07762123**
Actinomyces naeslundii		100.0	0.55	FRU 95		
07762140	**Good identification**					**07762140**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07762141	**Good identification**					**07762141**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07762142	**Good identification**					**07762142**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07762143	**Good identification**					**07762143**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07762160	**Good identification**					**07762160**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07762161	**Good identification**					**07762161**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07762162	**Good identification**					**07762162**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07762163	**Good identification**					**07762163**
Actinomyces naeslundii		100.0	0.42	FRU 95		
07763000	**Very good identification**					**07763000**
					SPO	
Actinomyces naeslundii		97.47	0.67	FRU 95	1	
Actinomyces israelii		2.15	0.30		1	
Clostridium innocuum		0.38	0.02		99	
07763001	**Very good identification**					**07763001**
Actinomyces naeslundii		100.0	0.70	FRU 95		
07763002	**Good identification**					**07763002**
Actinomyces naeslundii		94.68	0.67	FRU 95		
Actinomyces israelii		3.13	0.33			
Lactobacillus acidophilus		1.81	0.14			
07763003	**Very good identification**					**07763003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.70	FRU 95		
07763010	**Acceptable identification**					**07763010**
Actinomyces naeslundii		94.10	0.45	FRU 95		
Actinomyces israelii		5.90	0.16	MLZ 5		
07763011	**Good identification**					**07763011**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07763012	**Acceptable identification**					**07763012**
Actinomyces naeslundii		91.41	0.45	FRU 95		
Actinomyces israelii		8.59	0.19	MLZ 5		
07763013	**Good identification**					**07763013**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07763020	**Very good identification**					**07763020**
					SPO	
Actinomyces naeslundii		97.54	0.67	FRU 95	1	
Clostridium ramosum		2.46	0.15		92	
07763021	**Very good identification**					**07763021**
Actinomyces naeslundii		100.0	0.70	FRU 95		
07763022	**Low discrimination**					**07763022**
					SPO	
Actinomyces naeslundii		89.29	0.67	FRU 95	1	
Clostridium ramosum		9.01	0.25		92	
Lactobacillus acidophilus		1.71	0.14		1	
07763023	**Very good identification**					**07763023**
Actinomyces naeslundii		100.0	0.70	FRU 95		
07763030	**Good identification**					**07763030**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07763031	**Good identification**					**07763031**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07763032	**Good identification**					**07763032**
Actinomyces naeslundii		100.0	0.45	FRU 95		
07763033	**Good identification**					**07763033**
Actinomyces naeslundii		100.0	0.48	FRU 95		
07763040	**Good identification**					**07763040**
Actinomyces naeslundii		95.36	0.54	FRU 95		
Actinomyces israelii		4.64	0.23			
07763041	**Very good identification**					**07763041**
Actinomyces naeslundii		100.0	0.57	FRU 95		
07763042	**Acceptable identification**					**07763042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.05	0.54	FRU 95		
Actinomyces israelii		6.71	0.26			
Lactobacillus acidophilus		1.23	-.02			
07763043	**Very good identification**					**07763043**
Actinomyces naeslundii		100.0	0.57	FRU 95		
07763051	**Good identification**					**07763051**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763053	**Good identification**					**07763053**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763060	**Good identification**					**07763060**
Actinomyces naeslundii		100.0	0.54	FRU 95		
07763061	**Very good identification**					**07763061**
Actinomyces naeslundii		100.0	0.57	FRU 95		
07763062	**Good identification**					**07763062**
Actinomyces naeslundii		98.68	0.54	FRU 95		
Lactobacillus acidophilus		1.32	-.02			
07763063	**Very good identification**					**07763063**
Actinomyces naeslundii		100.0	0.57	FRU 95		
07763071	**Good identification**					**07763071**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763073	**Good identification**					**07763073**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763100	**Very good identification**					**07763100**
					SPO	
Actinomyces naeslundii		95.99	0.75	FRU 95	1	
Actinomyces israelii		2.98	0.40		1	
Clostridium ramosum		0.85	0.15		92	
07763101	**Very good identification**					**07763101**
Actinomyces naeslundii		100.0	0.78	FRU 95		
07763102	**Low discrimination**					**07763102**
Actinomyces naeslundii		85.04	0.75	FRU 95		
Lactobacillus acidophilus		6.58	0.32			
Actinomyces israelii		3.96	0.43			
07763103	**Very good identification**					**07763103**
Actinomyces naeslundii		100.0	0.78	FRU 95		
07763110	**Low discrimination**					**07763110**
					SPO	
Actinomyces naeslundii		89.02	0.53	FRU 95	1	
Actinomyces israelii		7.84	0.26	MLZ 5	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		3.14	0.05		99	
07763111	**Very good identification**					**07763111**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07763112	**Low discrimination**					**07763112**
Actinomyces naeslundii		65.07	0.53	FRU 95		
Bifidobacterium breve		9.54	0.24	MLZ 5		
Bifidobacterium adolescentis		8.71	0.23			
07763113	**Very good identification**					**07763113**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07763120	**Low discrimination**					**07763120**
					SPO	
Actinomyces naeslundii		53.09	0.75	FRU 95	1	
Clostridium ramosum		46.56	0.49		92	
Clostridium tertium		0.17	0.09		99	
07763121	**Very good identification**					**07763121**
					SPO	
Actinomyces naeslundii		99.41	0.78	FRU 95	1	
Clostridium ramosum		0.59	0.15		92	
07763122	**Low discrimination**					**07763122**
					SPO	
Clostridium ramosum		76.06	0.59	FRU 99	92	
Actinomyces naeslundii		21.68	0.75		1	
Lactobacillus acidophilus		1.68	0.32		1	
07763123	**Very good identification**					**07763123**
					SPO	
Actinomyces naeslundii		97.69	0.78	FRU 95	1	
Clostridium ramosum		2.31	0.25		92	
07763130	**Low discrimination**					**07763130**
					SPO	
Actinomyces naeslundii		76.93	0.53	FRU 95	1	
Clostridium ramosum		12.95	0.15	MLZ 5	92	
Clostridium tertium		8.14	0.13		99	
07763131	**Very good identification**					**07763131**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07763132	**Low discrimination**					**07763132**
					SPO	
Actinomyces naeslundii		44.42	0.53	FRU 95	1	
Clostridium ramosum		29.90	0.25	MLZ 5	92	
Clostridium tertium		14.10	0.21		99	
07763133	**Very good identification**					**07763133**
Actinomyces naeslundii		100.0	0.56	FRU 95		
07763140	**Good identification**					**07763140**
Actinomyces naeslundii		93.60	0.62	FRU 95		
Actinomyces israelii		6.40	0.33			
07763141	**Very good identification**					**07763141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.65	FRU 95		
07763142	**Low discrimination**					**07763142**
Actinomyces naeslundii		83.31	0.62	FRU 95		
Actinomyces israelii		8.54	0.36			
Lactobacillus acidophilus		4.51	0.16			
07763143	**Very good identification**					**07763143**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07763150	**Low discrimination**					**07763150**
Actinomyces naeslundii		83.75	0.40	FRU 95		
Actinomyces israelii		16.25	0.19	MLZ 5		
07763151	**Good identification**					**07763151**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07763152	**Low discrimination**					**07763152**
Actinomyces naeslundii		45.13	0.40	FRU 95		
Bifidobacterium breve		37.51	0.24	MLZ 5		
Actinomyces israelii		13.13	0.22			
07763153	**Good identification**					**07763153**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07763160	**Good identification**					**07763160**
					SPO	
Actinomyces naeslundii		94.53	0.62	FRU 95	1	
Clostridium ramosum		4.75	0.15		92	
Bifidobacterium breve		0.73	0.12		1	
07763161	**Very good identification**					**07763161**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07763162	**Low discrimination**					**07763162**
					SPO	
Actinomyces naeslundii		74.67	0.62	FRU 95	1	
Clostridium ramosum		14.99	0.25		92	
Bifidobacterium breve		5.81	0.29		1	
07763163	**Very good identification**					**07763163**
Actinomyces naeslundii		100.0	0.65	FRU 95		
07763170	**Low discrimination**					**07763170**
Actinomyces naeslundii		87.25	0.40	FRU 95		
Bifidobacterium breve		12.75	0.12	MLZ 5		
07763171	**Good identification**					**07763171**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07763173	**Good identification**					**07763173**
Actinomyces naeslundii		100.0	0.43	FRU 95		
07763201	**Good identification**					**07763201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763203	**Good identification**					**07763203**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763221	**Good identification**					**07763221**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763223	**Good identification**					**07763223**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763301	**Good identification**					**07763301**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07763303	**Good identification**					**07763303**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07763320	**Good identification**					**07763320**
					SPO	
Clostridium ramosum		99.14	0.51	FRU 99	92	
Actinomyces naeslundii		0.86	0.41		1	
07763321	**Low discrimination**					**07763321**
					SPO	
Actinomyces naeslundii		56.34	0.44	FRU 95	1	
Clostridium ramosum		43.66	0.17	RHA 1	92	
07763322	**Very good identification**					**07763322**
					SPO	
Clostridium ramosum		99.66	0.61	FRU 99	92	
Actinomyces naeslundii		0.22	0.41		1	
Lactobacillus acidophilus		0.13	0.12		1	
07763401	**Good identification**					**07763401**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763403	**Good identification**					**07763403**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763421	**Good identification**					**07763421**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763423	**Good identification**					**07763423**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07763500	**Low discrimination**					**07763500**
					SPO	
Actinomyces israelii		58.51	0.39	FRU 99	1	
Actinomyces naeslundii		23.29	0.41	XYL 92	1	
Clostridium butyricum		18.20	0.09		99	
07763501	**Good identification**					**07763501**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07763502	**Low discrimination**					**07763502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		42.95	0.35	FRU 99		
Actinomyces israelii		42.38	0.42	XYL 87		
Actinomyces naeslundii		11.25	0.41			
07763503	**Good identification**					**07763503**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07763512	**Low discrimination**					**07763512**
Bifidobacterium adolescentis		86.78	0.37	FRU 99		
Actinomyces israelii		10.47	0.28	XYL 87		
Bifidobacterium dentium		2.76	-.07			
07763520	**Low discrimination**					**07763520**
					SPO	
Actinomyces naeslundii		49.08	0.41	FRU 95	1	
Clostridium ramosum		43.05	0.15	ARA 1	92	
Actinomyces israelii		7.87	0.18		1	
07763521	**Good identification**					**07763521**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07763523	**Good identification**					**07763523**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07764612	**Good identification**					**07764612**
Clostridium clostridioforme		100.0	0.40	FRU 99		
07764632	**Good identification**					**07764632**
Clostridium clostridioforme		100.0	0.40	FRU 99		
07764702	**Good identification**					**07764702**
Clostridium clostridioforme		100.0	0.37	FRU 99		
07764712	**Good identification**					**07764712**
Clostridium clostridioforme		100.0	0.49	FRU 99		
07764722	**Good identification**					**07764722**
Clostridium clostridioforme		97.04	0.37	FRU 99		
Parabacteroides distasonis		2.96	0.17			
07764732	**Good identification**					**07764732**
Clostridium clostridioforme		99.64	0.49	FRU 99		
Parabacteroides distasonis		0.36	0.13			
07765000	**Low discrimination**					**07765000**
					SPO	
Actinomyces israelii		55.80	0.35	FRU 99	1	
Clostridium butyricum		37.72	0.11		99	
Actinomyces naeslundii		6.49	0.28		1	
07765002	**Low discrimination**					**07765002**
					SPO	
Actinomyces israelii		76.49	0.38	FRU 99	1	
Clostridium butyricum		13.40	0.04		99	
Actinomyces naeslundii		5.93	0.28		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07765101	**Low discrimination**					**07765101**
					SPO	
Actinomyces naeslundii		75.18	0.39	FRU 95	1	
Clostridium butyricum		24.83	0.01	XYL 1	99	
07765103	**Low discrimination**					**07765103**
					SPO	
Actinomyces naeslundii		79.96	0.39	FRU 95	1	
Clostridium butyricum		10.27	-.07	XYL 1	99	
Actinomyces israelii		9.77	0.14		1	
07765112	**Low discrimination**					**07765112**
Bifidobacterium adolescentis		66.15	0.38	FRU 99		
Actinomyces israelii		16.19	0.34	ARA 87		
Clostridium clostridioforme		10.54	0.22			
07765121	**Good identification**					**07765121**
Actinomyces naeslundii		100.0	0.39	FRU 95		
07765122	**Low discrimination**					**07765122**
					GRA	
Parabacteroides distasonis		69.51	0.47	FRU 99	1	
Lactobacillus acidophilus		8.13	0.19		99	
Prevotella oralis		6.96	0.17		1	
07765123	**Low discrimination**					**07765123**
					GRA	
Actinomyces naeslundii		85.89	0.39	FRU 95	99	
Parabacteroides distasonis		14.11	0.13	XYL 1	1	
07765132	**Low discrimination**					**07765132**
					GRA	
Parabacteroides distasonis		87.15	0.42	FRU 99	1	
Clostridium clostridioforme		11.71	0.22		50	
Actinomyces israelii		1.15	0.14		99	
07765140	**Low discrimination**					**07765140**
					SPO	
Actinomyces israelii		51.43	0.38	FRU 99	1	
Clostridium butyricum		46.64	0.17		99	
Actinomyces naeslundii		1.93	0.23		1	
07765142	**Low discrimination**					**07765142**
					SPO	
Actinomyces israelii		68.08	0.41	FRU 99	1	
Clostridium butyricum		16.01	0.10		99	
Bifidobacterium adolescentis		10.81	0.21		1	
07765312	**Good identification**					**07765312**
					GRA	
Clostridium clostridioforme		95.55	0.36	FRU 99	50	
Parabacteroides distasonis		2.94	0.16	ARA 94	1	
Actinomyces israelii		1.51	0.15		99	
07765322	**Acceptable identification**					**07765322**
					SPO	
Parabacteroides distasonis		92.48	0.55	FRU 99	1	
Clostridium clostridioforme		3.59	0.25		66	
Clostridium ramosum		2.01	0.14		92	
07765332	**Low discrimination**					**07765332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		75.04	0.50	FRU 99			
Clostridium clostridioforme		24.64	0.36				
Bacteroides ovatus		0.33	-.06				
07765500	**Good identification**						**07765500**
					SPO		
Clostridium butyricum		97.68	0.51	ESL 88	99		
Actinomyces israelii		2.19	0.44	FRU 99	1		
Bifidobacterium adolescentis		0.14	0.17		1		
07765502	**Low discrimination**						**07765502**
					SPO		
Clostridium butyricum		67.80	0.44	ESL 88	99		
Bifidobacterium adolescentis		23.95	0.51	FRU 99	1		
Actinomyces israelii		5.86	0.47		1		
07765512	**Low discrimination**						**07765512**
					SPO		
Bifidobacterium adolescentis		65.41	0.53	FRU 99	1		
Clostridium clostridioforme		24.40	0.42		66		
Clostridium butyricum		7.98	0.22		99		
07765522	**Low discrimination**						**07765522**
					GRA	SPO	
Parabacteroides distasonis		39.41	0.43	FRU 99	1	1	
Clostridium butyricum		39.34	0.26		99	99	
Clostridium clostridioforme		15.99	0.31		50	66	
07765532	**Low discrimination**						**07765532**
					GRA	SPO	
Clostridium clostridioforme		72.69	0.42	FRU 99	50	66	
Parabacteroides distasonis		21.17	0.39	RHA 88	1	1	
Clostridium butyricum		2.07	0.04		99	99	
07765552	**Low discrimination**						**07765552**
					SPO		
Bifidobacterium adolescentis		82.62	0.37	FRU 99	1		
Actinomyces israelii		7.78	0.26	SOR 11	1		
Clostridium butyricum		7.09	0.04		99		
07765602	**Good identification**						**07765602**
					SPO		
Clostridium clostridioforme		96.37	0.37	FRU 99	66		
Actinomyces israelii		2.03	0.17		1		
Clostridium butyricum		1.60	-.06		99		
07765612	**Good identification**						**07765612**
Clostridium clostridioforme		100.0	0.48	FRU 99			
07765622	**Good identification**						**07765622**
Clostridium clostridioforme		96.70	0.37	FRU 99			
Parabacteroides distasonis		3.30	0.17				
07765632	**Good identification**						**07765632**
Clostridium clostridioforme		99.60	0.48	FRU 99			
Parabacteroides distasonis		0.40	0.13				
07765702	**Low discrimination**						**07765702**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		85.59	0.46	FRU 99	66	
Clostridium butyricum		10.21	0.18		99	
Actinomyces israelii		2.15	0.28		1	
07765712	**Very good identification**					**07765712**
					SPO	
Clostridium clostridioforme		99.30	0.57	FRU 99	66	
Bifidobacterium adolescentis		0.37	0.18		1	
Clostridium butyricum		0.14	-.04		99	
07765722	**Low discrimination**					**07765722**
Parabacteroides distasonis		45.52	0.51	FRU 99		
Clostridium clostridioforme		45.16	0.46			
Bacteroides ovatus		8.41	0.23			
07765732	**Low discrimination**					**07765732**
Clostridium clostridioforme		87.17	0.57	FRU 99		
Parabacteroides distasonis		10.39	0.47			
Bacteroides ovatus		2.38	0.20			
07767000	**Acceptable identification**					**07767000**
					SPO	
Actinomyces israelii		94.96	0.48	FRU 99	1	
Actinomyces naeslundii		3.78	0.33		1	
Clostridium butyricum		1.26	-.05		99	
07767001	**Low discrimination**					**07767001**
Actinomyces naeslundii		85.53	0.36	FRU 95		
Actinomyces israelii		14.47	0.14	XYL 1		
07767002	**Good identification**					**07767002**
Actinomyces israelii		97.42	0.51	FRU 99		
Actinomyces naeslundii		2.59	0.33			
07767003	**Low discrimination**					**07767003**
Actinomyces naeslundii		79.77	0.36	FRU 95		
Actinomyces israelii		20.23	0.17	XYL 1		
07767012	**Good identification**					**07767012**
Actinomyces israelii		100.0	0.37	FRU 99		
07767021	**Good identification**					**07767021**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07767023	**Good identification**					**07767023**
Actinomyces naeslundii		100.0	0.36	FRU 95		
07767040	**Good identification**					**07767040**
Actinomyces israelii		100.0	0.41	FRU 99		
07767042	**Good identification**					**07767042**
Actinomyces israelii		100.0	0.44	FRU 99		
07767100	**Low discrimination**					**07767100**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		89.94	0.58	FRU 99	1	
Clostridium butyricum		7.15	0.18		99	
Actinomyces naeslundii		2.55	0.41		1	
07767101	**Low discrimination**					**07767101**
Actinomyces naeslundii		80.79	0.44	FRU 95		
Actinomyces israelii		19.21	0.24	XYL 1		
07767102	**Low discrimination**					**07767102**
					SPO	
Actinomyces israelii		89.31	0.61	FRU 99	1	
Bifidobacterium adolescentis		6.44	0.35		1	
Clostridium butyricum		1.84	0.11		99	
07767103	**Low discrimination**					**07767103**
Actinomyces naeslundii		73.71	0.44	FRU 95		
Actinomyces israelii		26.29	0.27	XYL 1		
07767110	**Acceptable identification**					**07767110**
					SPO	
Actinomyces israelii		92.93	0.44	FRU 99	1	
Clostridium tertium		4.47	0.07	MLZ 13	99	
Clostridium butyricum		2.60	-.04		99	
07767112	**Low discrimination**					**07767112**
					SPO	
Actinomyces israelii		58.66	0.47	FRU 99	1	
Bifidobacterium adolescentis		34.60	0.37	MLZ 13	1	
Clostridium tertium		5.65	0.15		99	
07767120	**Low discrimination**					**07767120**
					SPO	
Actinomyces israelii		46.92	0.38	FRU 99	1	
Actinomyces naeslundii		20.82	0.41	NAG 6	1	
Clostridium ramosum		18.26	0.15		92	
07767121	**Good identification**					**07767121**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07767123	**Good identification**					**07767123**
Actinomyces naeslundii		100.0	0.44	FRU 95		
07767140	**Good identification**					**07767140**
					SPO	
Actinomyces israelii		97.03	0.51	FRU 99	1	
Clostridium butyricum		1.73	0.00		99	
Actinomyces naeslundii		1.25	0.28		1	
07767142	**Good identification**					**07767142**
Actinomyces israelii		96.52	0.54	FRU 99		
Bifidobacterium adolescentis		2.21	0.20			
Actinomyces naeslundii		0.83	0.28			
07767150	**Acceptable identification**					**07767150**
					COC	
Actinomyces israelii		93.35	0.37	FRU 99	1	
Blautia producta		6.65	-.03	MLZ 13	99	
07767152	**Low discrimination**					**07767152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		84.21	0.40	FRU 99		
Bifidobacterium adolescentis		15.79	0.21	MLZ 13		
07767300	**Good identification**					**07767300**
					SPO	
Actinomyces israelii		96.84	0.39	FRU 99	1	
Clostridium butyricum		3.16	-.08	RHA 7	99	
07767302	**Good identification**					**07767302**
Actinomyces israelii		100.0	0.42	FRU 99		
07767342	**Good identification**					**07767342**
Actinomyces israelii		100.0	0.35	FRU 99		
07767400	**Low discrimination**					**07767400**
					SPO	
Actinomyces israelii		87.28	0.46	FRU 99	1	
Clostridium butyricum		12.72	0.11		99	
07767402	**Acceptable identification**					**07767402**
					SPO	
Actinomyces israelii		94.20	0.49	FRU 99	1	
Clostridium butyricum		3.56	0.04		99	
Bifidobacterium adolescentis		2.25	0.15		1	
07767412	**Low discrimination**					**07767412**
Actinomyces israelii		77.73	0.35	FRU 99		
Bifidobacterium adolescentis		15.15	0.17	MLZ 13		
Bifidobacterium dentium		7.12	-.07			
07767440	**Good identification**					**07767440**
					SPO	
Actinomyces israelii		96.84	0.39	FRU 99	1	
Clostridium butyricum		3.16	-.07		99	
07767442	**Good identification**					**07767442**
Actinomyces israelii		100.0	0.42	FRU 99		
07767500	**Low discrimination**					**07767500**
					SPO	
Actinomyces israelii		53.09	0.57	FRU 99	1	
Clostridium butyricum		46.44	0.35		99	
Bifidobacterium adolescentis		0.47	0.15		1	
07767502	**Low discrimination**					**07767502**
					SPO	
Actinomyces israelii		54.95	0.60	FRU 99	1	
Bifidobacterium adolescentis		32.41	0.50		1	
Clostridium butyricum		12.46	0.28		99	
07767510	**Low discrimination**					**07767510**
					SPO	
Actinomyces israelii		57.09	0.43	FRU 99	1	
Bifidobacterium dentium		19.21	0.10	MLZ 13	1	
Clostridium butyricum		17.59	0.13		99	
07767512	**Low discrimination**					**07767512**
Bifidobacterium adolescentis		58.87	0.51	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		27.68	0.27				
Actinomyces israelii		12.20	0.46				
07767522	**Low discrimination**						**07767522**
					SPO		
Actinomyces israelii		66.77	0.39	FRU 99	1		
Clostridium butyricum		20.63	0.10	NAG 6	99		
Bifidobacterium dentium		6.37	-.03		1		
07767540	**Low discrimination**						**07767540**
					COC	SPO	
Actinomyces israelii		61.53	0.50	FRU 99	1	1	
Blautia producta		26.43	0.24		99	1	
Clostridium butyricum		12.04	0.17		1	99	
07767542	**Low discrimination**						**07767542**
					SPO		
Actinomyces israelii		78.74	0.53	FRU 99	1		
Bifidobacterium adolescentis		14.76	0.34		1		
Clostridium butyricum		3.99	0.10		99		
07767552	**Low discrimination**						**07767552**
					COC		
Bifidobacterium adolescentis		47.46	0.36	FRU 99	1		
Actinomyces israelii		30.95	0.39	SOR 11	1		
Blautia producta		19.77	0.15		99		
07767700	**Low discrimination**						**07767700**
					SPO		
Actinomyces israelii		73.56	0.38	FRU 99	1		
Clostridium butyricum		26.44	0.09	RHA 7	99		
07767702	**Low discrimination**						**07767702**
					SPO		
Actinomyces israelii		80.43	0.41	FRU 99	1		
Clostridium butyricum		7.49	0.02	RHA 7	99		
Bifidobacterium adolescentis		6.37	0.15		1		
07770000	**Low discrimination**						**07770000**
					GRA		
Fusobacterium mortiferum		60.38	0.55	MLT 11	1		
Actinomyces naeslundii		27.25	0.61		99		
Lactobacillus jensenii		10.03	0.32		99		
07770001	**Very good identification**						**07770001**
					GRA		
Actinomyces naeslundii		98.53	0.64	GAL 96	99		
Fusobacterium mortiferum		1.47	0.21		1		
07770002	**Low discrimination**						**07770002**
					GRA		
Actinomyces naeslundii		38.72	0.61	GAL 96	99		
Lactobacillus acidophilus		34.52	0.35		99		
Fusobacterium mortiferum		9.53	0.39		1		
07770003	**Very good identification**						**07770003**
Actinomyces naeslundii		96.11	0.64	GAL 96			
Actinomyces viscosus		3.31	0.15				
Lactobacillus acidophilus		0.58	0.01				
07770011	**Good identification**						**07770011**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.42	GAL 96			
07770013	**Good identification**						**07770013**
Actinomyces naeslundii		100.0	0.42	GAL 96			
07770020	**Low discrimination**						**07770020**
					GRA		
Actinomyces naeslundii		86.06	0.61	GAL 96	99		
Lactobacillus jensenii		7.92	0.21		99		
Fusobacterium mortiferum		1.93	0.21		1		
07770021	**Very good identification**						**07770021**
Actinomyces naeslundii		100.0	0.64	GAL 96			
07770022	**Low discrimination**						**07770022**
Actinomyces naeslundii		48.24	0.61	GAL 96			
Lactobacillus acidophilus		43.02	0.35				
Lactobacillus jensenii		2.96	0.18				
07770023	**Very good identification**						**07770023**
Actinomyces naeslundii		99.40	0.64	GAL 96			
Lactobacillus acidophilus		0.60	0.01				
07770030	**Low discrimination**						**07770030**
Actinomyces naeslundii		63.19	0.39	GAL 96			
Lactobacillus jensenii		36.82	0.13	MLZ 5			
07770031	**Good identification**						**07770031**
Actinomyces naeslundii		100.0	0.42	GAL 96			
07770033	**Good identification**						**07770033**
Actinomyces naeslundii		100.0	0.42	GAL 96			
07770040	**Low discrimination**						**07770040**
					GRA	SPO	
Actinomyces naeslundii		85.19	0.48	GAL 96	99	1	
Fusobacterium mortiferum		10.80	0.21		1	1	
Clostridium botulinum biovar B		4.01	-.03		99	99	
07770041	**Good identification**						**07770041**
Actinomyces naeslundii		100.0	0.51	GAL 96			
07770042	**Low discrimination**						**07770042**
Actinomyces naeslundii		61.56	0.48	GAL 96			
Lactobacillus acidophilus		38.44	0.20				
07770043	**Good identification**						**07770043**
Actinomyces naeslundii		100.0	0.51	GAL 96			
07770060	**Good identification**						**07770060**
Actinomyces naeslundii		100.0	0.48	GAL 96			
07770061	**Good identification**						**07770061**
Actinomyces naeslundii		100.0	0.51	GAL 96			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07770062	**Low discrimination**					**07770062**
Actinomyces naeslundii		61.56	0.48	GAL 96		
Lactobacillus acidophilus		38.44	0.20			
07770063	**Good identification**					**07770063**
Actinomyces naeslundii		100.0	0.51	GAL 96		
07770100	**Low discrimination**					**07770100**
					GRA	
Fusobacterium mortiferum		62.16	0.61	MLT 11	1	
Actinomyces naeslundii		34.22	0.68		99	
Lactobacillus acidophilus		1.25	0.19		99	
07770101	**Very good identification**					**07770101**
					GRA	
Actinomyces naeslundii		98.79	0.71	GAL 96	99	
Fusobacterium mortiferum		1.21	0.27		1	
07770102	**Low discrimination**					**07770102**
					GRA	
Lactobacillus acidophilus		71.71	0.54	GAL 99	99	
Actinomyces naeslundii		19.90	0.68		99	
Fusobacterium mortiferum		4.02	0.45		1	
07770103	**Very good identification**					**07770103**
Actinomyces naeslundii		95.50	0.71	GAL 96		
Lactobacillus acidophilus		2.32	0.19			
Actinomyces viscosus		2.05	0.19			
07770110	**Low discrimination**					**07770110**
					GRA	
Actinomyces naeslundii		71.74	0.47	GAL 96	99	
Fusobacterium mortiferum		25.01	0.27	MLZ 5	1	
Leptotrichia buccalis		3.26	-.11		1	
07770111	**Good identification**					**07770111**
Actinomyces naeslundii		100.0	0.50	GAL 96		
07770113	**Good identification**					**07770113**
					GRA	
Actinomyces naeslundii		97.06	0.50	GAL 96	99	
Leptotrichia buccalis		2.94	-.11	MLZ 5	1	
07770120	**Low discrimination**					**07770120**
					SPO	
Actinomyces naeslundii		66.62	0.68	GAL 96	1	
Clostridium ramosum		25.28	0.36		92	
Lactobacillus acidophilus		2.42	0.19		1	
07770121	**Very good identification**					**07770121**
					SPO	
Actinomyces naeslundii		99.75	0.71	GAL 96	1	
Clostridium ramosum		0.26	0.02		92	
07770122	**Low discrimination**					**07770122**
					SPO	
Lactobacillus acidophilus		55.48	0.54	GAL 99	1	
Clostridium ramosum		23.37	0.46		92	
Actinomyces naeslundii		15.40	0.68		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07770123	**Very good identification**					**07770123**
					SPO	
Actinomyces naeslundii		96.51	0.71	GAL 96	1	
Lactobacillus acidophilus		2.34	0.19		1	
Clostridium ramosum		0.99	0.12		92	
07770130	**Low discrimination**					**07770130**
					SPO	
Actinomyces naeslundii		88.02	0.47	GAL 96	1	
Clostridium ramosum		6.41	0.02	MLZ 5	92	
Bifidobacterium breve		5.57	0.12		1	
07770131	**Good identification**					**07770131**
Actinomyces naeslundii		100.0	0.50	GAL 96		
07770133	**Good identification**					**07770133**
Actinomyces naeslundii		100.0	0.50	GAL 96		
07770140	**Low discrimination**					**07770140**
					GRA	
Actinomyces naeslundii		86.28	0.56	GAL 96	99	
Fusobacterium mortiferum		8.97	0.27		1	
Actinomyces israelii		2.55	0.20		99	
07770141	**Very good identification**					**07770141**
Actinomyces naeslundii		100.0	0.59	GAL 96		
07770142	**Low discrimination**					**07770142**
Lactobacillus acidophilus		68.22	0.38	GAL 99		
Actinomyces naeslundii		27.04	0.56	SOR 11		
Bifidobacterium breve		2.90	0.24			
07770143	**Very good identification**					**07770143**
Actinomyces naeslundii		98.33	0.59	GAL 96		
Lactobacillus acidophilus		1.67	0.04			
07770151	**Good identification**					**07770151**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07770153	**Good identification**					**07770153**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07770160	**Acceptable identification**					**07770160**
Actinomyces naeslundii		91.20	0.56	GAL 96		
Clostridium perfringens		2.77	0.07			
Lactobacillus acidophilus		2.32	0.04			
07770161	**Very good identification**					**07770161**
Actinomyces naeslundii		100.0	0.59	GAL 96		
07770162	**Low discrimination**					**07770162**
Lactobacillus acidophilus		65.89	0.38	GAL 99		
Actinomyces naeslundii		26.11	0.56	SOR 11		
Bifidobacterium breve		4.98	0.29			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07770163	**Very good identification**						**07770163**
Actinomyces naeslundii		98.33	0.59	GAL 96			
Lactobacillus acidophilus		1.67	0.04				
07770171	**Good identification**						**07770171**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07770173	**Good identification**						**07770173**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07770212	**Good identification**						**07770212**
Clostridium clostridioforme		100.0	0.36	XYL 91			
07770232	**Good identification**						**07770232**
Clostridium clostridioforme		100.0	0.36	XYL 91			
07770301	**Good identification**						**07770301**
Actinomyces naeslundii		100.0	0.37	GAL 96			
07770303	**Low discrimination**						**07770303**
Actinomyces naeslundii		84.69	0.37	GAL 96			
Lactobacillus acidophilus		15.31	0.00	RHA 1			
07770312	**Good identification**						**07770312**
					GRA		
Clostridium clostridioforme		99.06	0.45	XYL 91	50		
Lactobacillus acidophilus		0.50	0.00	ARA 94	99		
Leptotrichia buccalis		0.44	-.11		1		
07770320	**Good identification**						**07770320**
					SPO	GRA	
Clostridium ramosum		96.45	0.38	GAL 99	92	50	
Actinomyces naeslundii		1.94	0.34		1	99	
Prevotella oralis		1.09	0.04		1	1	
07770321	**Low discrimination**						**07770321**
					SPO		
Actinomyces naeslundii		74.89	0.37	GAL 96	1		
Clostridium ramosum		25.11	0.04	RHA 1	92		
07770322	**Low discrimination**						**07770322**
					SPO		
Clostridium ramosum		76.70	0.48	GAL 99	92		
Lactobacillus acidophilus		10.34	0.34		1		
Clostridium clostridioforme		4.51	0.34		66		
07770332	**Low discrimination**						**07770332**
					SPO		
Clostridium clostridioforme		87.22	0.45	XYL 91	66		
Parabacteroides distasonis		9.04	0.34	ARA 94	1		
Clostridium ramosum		3.29	0.14		92		
07770412	**Good identification**						**07770412**
Clostridium clostridioforme		100.0	0.42	XYL 91			
07770432	**Good identification**						**07770432**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.42	XYL 91		
07770501	**Good identification**					**07770501**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07770502	**Low discrimination**					**07770502**
					SPO	
Clostridium clostridioforme		50.43	0.40	XYL 91	66	
Bifidobacterium adolescentis		23.07	0.37	RHA 88	1	
Clostridium butyricum		13.52	0.18		99	
07770503	**Low discrimination**					**07770503**
Actinomyces naeslundii		85.58	0.37	GAL 96		
Clostridium clostridioforme		14.42	0.05	ARA 1		
07770512	**Low discrimination**					**07770512**
					SPO	
Clostridium clostridioforme		88.49	0.51	XYL 91	66	
Bifidobacterium adolescentis		10.86	0.38	RHA 88	1	
Clostridium butyricum		0.27	-.04		99	
07770521	**Good identification**					**07770521**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07770522	**Low discrimination**					**07770522**
					GRA	
Clostridium clostridioforme		73.42	0.40	XYL 91	50	
Lactobacillus acidophilus		10.58	0.19	RHA 88	99	
Parabacteroides distasonis		6.16	0.27		1	
07770523	**Low discrimination**					**07770523**
Actinomyces naeslundii		85.58	0.37	GAL 96		
Clostridium clostridioforme		14.42	0.05	ARA 1		
07770532	**Good identification**					**07770532**
Clostridium clostridioforme		99.02	0.51	XYL 91		
Parabacteroides distasonis		0.98	0.22	RHA 88		
07770602	**Good identification**					**07770602**
Clostridium clostridioforme		100.0	0.45	XYL 91		
07770612	**Very good identification**					**07770612**
Clostridium clostridioforme		100.0	0.57	XYL 91		
07770622	**Good identification**					**07770622**
Clostridium clostridioforme		100.0	0.45	XYL 91		
07770632	**Very good identification**					**07770632**
Clostridium clostridioforme		100.0	0.57	XYL 91		
07770702	**Good identification**					**07770702**
					SPO	
Clostridium clostridioforme		99.74	0.54	XYL 91	66	
Lactobacillus acidophilus		0.15	0.00		1	
Clostridium butyricum		0.11	-.08		99	
07770712	**Very good identification**					**07770712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.66	XYL 91		
07770722	**Good identification**					**07770722**
					SPO	
Clostridium clostridioforme		95.53	0.54	XYL 91	66	
Parabacteroides distasonis		3.28	0.35		1	
Clostridium ramosum		1.05	0.14		92	
07770732	**Very good identification**					**07770732**
Clostridium clostridioforme		99.60	0.66	XYL 91		
Parabacteroides distasonis		0.40	0.30			
07771000	**Low discrimination**					**07771000**
					GRA	
Lactobacillus jensenii		54.93	0.66	RAF 1	99	
Actinomyces naeslundii		36.19	0.84		99	
Fusobacterium mortiferum		5.01	0.58		1	
07771001	**Very good identification**					**07771001**
					GRA	
Actinomyces naeslundii		98.85	0.87		99	
Lactobacillus jensenii		1.01	0.32		99	
Fusobacterium mortiferum		0.09	0.24		1	
07771002	**Low discrimination**					**07771002**
Lactobacillus acidophilus		63.34	0.70	LAC 92		
Lactobacillus jensenii		17.43	0.63			
Actinomyces naeslundii		17.22	0.84			
07771003	**Very good identification**					**07771003**
Actinomyces naeslundii		96.82	0.87			
Lactobacillus acidophilus		2.40	0.35			
Lactobacillus jensenii		0.66	0.29			
07771010	**Low discrimination**					**07771010**
Lactobacillus jensenii		89.17	0.58	RAF 1		
Actinomyces naeslundii		9.28	0.62			
Actinomyces israelii		1.04	0.37			
07771011	**Good identification**					**07771011**
Actinomyces naeslundii		93.92	0.65	MLZ 5		
Lactobacillus jensenii		6.08	0.24			
07771012	**Low discrimination**					**07771012**
Lactobacillus jensenii		70.78	0.55	RAF 1		
Actinomyces naeslundii		11.05	0.62			
Lactobacillus acidophilus		7.80	0.35			
07771013	**Very good identification**					**07771013**
Actinomyces naeslundii		95.10	0.65	MLZ 5		
Lactobacillus jensenii		4.10	0.21			
Lactobacillus acidophilus		0.45	0.01			
07771020	**Low discrimination**					**07771020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		65.30	0.84			
Lactobacillus jensenii		24.78	0.56			
Clostridium perfringens		2.94	0.39			
07771021	**Very good identification**					**07771021**
Actinomyces naeslundii		99.66	0.87			
Lactobacillus jensenii		0.26	0.21			
Clostridium perfringens		0.03	0.05			
07771022	**Low discrimination**					**07771022**
Lactobacillus acidophilus		68.73	0.70	LAC 92		
Actinomyces naeslundii		18.69	0.84			
Lactobacillus jensenii		4.73	0.53			
07771023	**Very good identification**					**07771023**
Actinomyces naeslundii		97.19	0.87			
Lactobacillus acidophilus		2.41	0.35			
Lactobacillus jensenii		0.17	0.18			
07771030	**Low discrimination**					**07771030**
Lactobacillus jensenii		69.82	0.47	RAF 1		
Actinomyces naeslundii		29.05	0.62			
Clostridium perfringens		0.25	0.05			
07771031	**Very good identification**					**07771031**
Actinomyces naeslundii		98.41	0.65	MLZ 5		
Lactobacillus jensenii		1.59	0.13			
07771032	**Low discrimination**					**07771032**
Lactobacillus jensenii		43.13	0.44	RAF 1		
Actinomyces naeslundii		26.92	0.62			
Lactobacillus acidophilus		19.00	0.35			
07771033	**Very good identification**					**07771033**
Actinomyces naeslundii		98.47	0.65	MLZ 5		
Lactobacillus jensenii		1.06	0.10			
Lactobacillus acidophilus		0.47	0.01			
07771040	**Low discrimination**					**07771040**
Actinomyces naeslundii		82.39	0.71			
Lactobacillus jensenii		7.16	0.32			
Actinomyces israelii		7.15	0.44			
07771041	**Very good identification**					**07771041**
Actinomyces naeslundii		100.0	0.74			
07771042	**Low discrimination**					**07771042**
Lactobacillus acidophilus		68.11	0.54	LAC 92		
Actinomyces naeslundii		26.44	0.71	SOR 11		
Actinomyces israelii		3.44	0.47			
07771043	**Very good identification**					**07771043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.30	0.74			
Lactobacillus acidophilus		1.71	0.20			
07771050	**Low discrimination**					**07771050**
Actinomyces naeslundii		55.66	0.50	MLZ 5		
Lactobacillus jensenii		30.63	0.24			
Actinomyces israelii		13.71	0.30			
07771051	**Good identification**					**07771051**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07771052	**Low discrimination**					**07771052**
Actinomyces naeslundii		39.43	0.50	MLZ 5		
Lactobacillus acidophilus		19.49	0.20			
Actinomyces israelii		14.57	0.33			
07771053	**Good identification**					**07771053**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07771060	**Good identification**					**07771060**
Actinomyces naeslundii		90.18	0.71			
Clostridium perfringens		4.39	0.27			
Lactobacillus acidophilus		2.35	0.20			
07771061	**Very good identification**					**07771061**
Actinomyces naeslundii		100.0	0.74			
07771062	**Low discrimination**					**07771062**
Lactobacillus acidophilus		70.16	0.54	LAC 92		
Actinomyces naeslundii		27.23	0.71	SOR 11		
Bifidobacterium breve		0.62	0.29			
07771063	**Very good identification**					**07771063**
Actinomyces naeslundii		98.30	0.74			
Lactobacillus acidophilus		1.71	0.20			
07771070	**Low discrimination**					**07771070**
Actinomyces naeslundii		84.74	0.50	MLZ 5		
Lactobacillus jensenii		11.66	0.13			
Bifidobacterium breve		3.60	0.12			
07771071	**Good identification**					**07771071**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07771072	**Low discrimination**					**07771072**
Actinomyces naeslundii		49.61	0.50	MLZ 5		
Lactobacillus acidophilus		24.53	0.20			
Bifidobacterium breve		21.32	0.29			
07771073	**Good identification**					**07771073**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
07771100	**Low discrimination**					**07771100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces naeslundii		63.33	0.92		1		
Lactobacillus acidophilus		9.51	0.54		1		
Atopobium parvulum		8.62	0.54		99		
07771101	**Excellent identification**						**07771101**
					COC		
Actinomyces naeslundii		99.63	0.95		1		
Lactobacillus acidophilus		0.10	0.19		1		
Atopobium parvulum		0.09	0.20		99		
07771102	**Good identification**						**07771102**
Lactobacillus acidophilus		91.11	0.88				
Actinomyces naeslundii		6.13	0.92				
Actinomyces israelii		0.51	0.64				
07771103	**Good identification**						**07771103**
Actinomyces naeslundii		90.62	0.95				
Lactobacillus acidophilus		9.07	0.54				
Actinomyces israelii		0.05	0.30				
07771110	**Low discrimination**						**07771110**
					GRA		
Actinomyces naeslundii		58.58	0.70	MLZ 5	99		
Leptotrichia buccalis		10.97	0.24		1		
Actinomyces israelii		9.21	0.47		99		
07771111	**Very good identification**						**07771111**
					GRA		
Actinomyces naeslundii		99.87	0.73	MLZ 5	99		
Leptotrichia buccalis		0.13	-.11		1		
07771112	**Low discrimination**						**07771112**
					GRA		
Leptotrichia buccalis		71.18	0.58	ESL 99	1		
Lactobacillus acidophilus		10.95	0.54		99		
Bifidobacterium adolescentis		6.35	0.58		99		
07771113	**Low discrimination**						**07771113**
					GRA		
Actinomyces naeslundii		85.62	0.73	MLZ 5	99		
Leptotrichia buccalis		10.69	0.24		1		
Lactobacillus acidophilus		1.64	0.19		99		
07771120	**Low discrimination**						**07771120**
					SPO		
Clostridium ramosum		52.25	0.70		92		
Actinomyces naeslundii		33.38	0.92		1		
Lactobacillus acidophilus		5.01	0.54		1		
07771121	**Very good identification**						**07771121**
					SPO		
Actinomyces naeslundii		98.69	0.95		1		
Clostridium ramosum		1.04	0.36		92		
Lactobacillus acidophilus		0.10	0.19		1		
07771122	**Low discrimination**						**07771122**
					SPO	GRA	
Lactobacillus acidophilus		63.10	0.88		1	99	
Clostridium ramosum		26.59	0.80		92	50	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		4.41	0.67		1 1	
07771123	**Low discrimination**					**07771123**
					SPO	
Actinomyces naeslundii		86.83	0.95		1	
Lactobacillus acidophilus		8.69	0.54		1	
Clostridium ramosum		3.66	0.46		92	
07771130	**Low discrimination**					**07771130**
					SPO	
Actinomyces naeslundii		58.50	0.70	MLZ 5	1	
Clostridium ramosum		17.58	0.36		92	
Bifidobacterium breve		15.26	0.46		1	
07771131	**Very good identification**					**07771131**
					SPO	
Actinomyces naeslundii		99.62	0.73	MLZ 5	1	
Clostridium ramosum		0.20	0.02		92	
Bifidobacterium breve		0.18	0.12		1	
07771132	**Low discrimination**					**07771132**
					GRA	
Lactobacillus acidophilus		24.43	0.54	MLZ 1	99	
Bifidobacterium breve		22.59	0.63		99	
Leptotrichia buccalis		17.65	0.41		1	
07771133	**Good identification**					**07771133**
Actinomyces naeslundii		93.35	0.73	MLZ 5		
Lactobacillus acidophilus		1.79	0.19			
Bifidobacterium breve		1.66	0.29			
07771140	**Low discrimination**					**07771140**
Actinomyces naeslundii		75.16	0.79			
Actinomyces israelii		9.17	0.54			
Lactobacillus acidophilus		7.90	0.38			
07771141	**Very good identification**					**07771141**
Actinomyces naeslundii		99.85	0.82			
Actinomyces israelii		0.08	0.20			
Lactobacillus acidophilus		0.07	0.04			
07771142	**Low discrimination**					**07771142**
Lactobacillus acidophilus		85.93	0.72	SOR 11		
Actinomyces naeslundii		8.26	0.79			
Bifidobacterium breve		3.65	0.59			
07771143	**Good identification**					**07771143**
Actinomyces naeslundii		93.05	0.82			
Lactobacillus acidophilus		6.52	0.38			
Bifidobacterium breve		0.28	0.24			
07771150	**Low discrimination**					**07771150**
Actinomyces naeslundii		44.26	0.57	MLZ 5		
Bifidobacterium breve		36.80	0.41			
Actinomyces israelii		15.33	0.40			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07771151	**Very good identification**					**07771151**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
07771152	**Low discrimination**					**07771152**
Bifidobacterium breve		59.33	0.59	ESL 99		
Lactobacillus acidophilus		14.10	0.38			
Bifidobacterium adolescentis		8.18	0.43			
07771153	**Good identification**					**07771153**
Actinomyces naeslundii		92.17	0.60	MLZ 5		
Bifidobacterium breve		5.22	0.24			
Lactobacillus acidophilus		1.24	0.04			
07771160	**Low discrimination**					**07771160**
Actinomyces naeslundii		69.72	0.79			
Clostridium perfringens		8.74	0.41			
Lactobacillus acidophilus		7.33	0.38			
07771161	**Very good identification**					**07771161**
Actinomyces naeslundii		99.73	0.82			
Clostridium perfringens		0.08	0.07			
Lactobacillus acidophilus		0.07	0.04			
07771162	**Low discrimination**					**07771162**
Lactobacillus acidophilus		81.90	0.72	SOR 11		
Actinomyces naeslundii		7.87	0.79			
Bifidobacterium breve		6.19	0.63			
07771163	**Good identification**					**07771163**
Actinomyces naeslundii		92.71	0.82			
Lactobacillus acidophilus		6.50	0.38			
Bifidobacterium breve		0.49	0.29			
07771170	**Low discrimination**					**07771170**
Bifidobacterium breve		57.70	0.46	ESL 99		
Actinomyces naeslundii		39.04	0.57	BGL 91		
Clostridium perfringens		0.94	0.07			
07771171	**Very good identification**					**07771171**
Actinomyces naeslundii		99.01	0.60	MLZ 5		
Bifidobacterium breve		0.99	0.12			
07771172	**Low discrimination**					**07771172**
Bifidobacterium breve		81.71	0.63	ESL 99		
Lactobacillus acidophilus		10.92	0.38			
Actinomyces naeslundii		5.47	0.57			
07771173	**Low discrimination**					**07771173**
Actinomyces naeslundii		89.76	0.60	MLZ 5		
Bifidobacterium breve		9.03	0.29			
Lactobacillus acidophilus		1.21	0.04			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07771201	**Good identification**						**07771201**
Actinomyces naeslundii		100.0	0.53	RHA 1			
07771202	**Low discrimination**						**07771202**
Lactobacillus acidophilus		86.18	0.50	LAC 92			
Clostridium clostridioforme		3.91	0.33	RHA 7			
Lactobacillus jensenii		3.18	0.29				
07771203	**Low discrimination**						**07771203**
Actinomyces naeslundii		84.42	0.53	RHA 1			
Lactobacillus acidophilus		15.58	0.16				
07771212	**Low discrimination**						**07771212**
Clostridium clostridioforme		88.17	0.44	XYL 91			
Lactobacillus jensenii		5.25	0.21	ARA 94			
Lactobacillus acidophilus		4.31	0.16				
07771220	**Low discrimination**						**07771220**
					SPO	GRA	
Clostridium ramosum		73.21	0.38	LAC 99	92	50	
Actinomyces naeslundii		12.40	0.50		1	99	
Prevotella oralis		5.23	0.18		1	1	
07771221	**Good identification**						**07771221**
					SPO		
Actinomyces naeslundii		96.18	0.53	RHA 1	1		
Clostridium ramosum		3.83	0.04		92		
07771222	**Low discrimination**						**07771222**
					SPO	GRA	
Lactobacillus acidophilus		43.57	0.50	LAC 92	1	99	
Clostridium ramosum		37.56	0.48	RHA 7	92	50	
Prevotella oralis		12.74	0.40		1	1	
07771223	**Low discrimination**						**07771223**
					SPO		
Actinomyces naeslundii		71.55	0.53	RHA 1	1		
Lactobacillus acidophilus		13.21	0.16		1		
Clostridium ramosum		11.38	0.14		92		
07771232	**Low discrimination**						**07771232**
					GRA		
Clostridium clostridioforme		80.32	0.44	XYL 91	50		
Parabacteroides distasonis		9.29	0.34	ARA 94	1		
Lactobacillus acidophilus		3.92	0.16		99		
07771240	**Low discrimination**						**07771240**
Actinomyces naeslundii		54.33	0.37	RHA 1			
Actinomyces israelii		35.13	0.25				
Lactobacillus acidophilus		10.54	0.01				
07771241	**Good identification**						**07771241**
Actinomyces naeslundii		100.0	0.40	RHA 1			
07771243	**Low discrimination**						**07771243**
Actinomyces naeslundii		88.55	0.40	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		11.45	0.01				
07771260	**Low discrimination**						**07771260**
					SPO		
Actinomyces naeslundii		65.27	0.37	RHA 1	1		
Clostridium ramosum		22.07	0.04		92		
Lactobacillus acidophilus		12.66	0.01		1		
07771261	**Good identification**						**07771261**
Actinomyces naeslundii		100.0	0.40	RHA 1			
07771263	**Low discrimination**						**07771263**
Actinomyces naeslundii		88.55	0.40	RHA 1			
Lactobacillus acidophilus		11.45	0.01				
07771300	**Low discrimination**						**07771300**
					SPO		
Clostridium ramosum		39.52	0.38	NAG 99	92		
Lactobacillus acidophilus		21.31	0.34		1		
Actinomyces naeslundii		19.05	0.58		1		
07771301	**Very good identification**						**07771301**
					SPO		
Actinomyces naeslundii		97.90	0.61	RHA 1	1		
Clostridium ramosum		1.37	0.04		92		
Lactobacillus acidophilus		0.74	0.00		1		
07771302	**Low discrimination**						**07771302**
					SPO	GRA	
Lactobacillus acidophilus		88.45	0.68	RHA 7	1	99	
Clostridium ramosum		6.63	0.48		92	50	
Prevotella oralis		1.85	0.39		1	1	
07771303	**Low discrimination**						**07771303**
					SPO		
Actinomyces naeslundii		54.73	0.61	RHA 1	1		
Lactobacillus acidophilus		40.82	0.34		1		
Clostridium ramosum		3.06	0.14		92		
07771311	**Good identification**						**07771311**
Actinomyces naeslundii		100.0	0.39	RHA 1			
07771312	**Low discrimination**						**07771312**
					GRA		
Clostridium clostridioforme		70.86	0.53	XYL 91	50		
Lactobacillus acidophilus		11.89	0.34	ARA 94	99		
Leptotrichia buccalis		10.37	0.24		1		
07771313	**Low discrimination**						**07771313**
Actinomyces naeslundii		47.15	0.39	RHA 1			
Clostridium clostridioforme		40.22	0.19	MLZ 5			
Lactobacillus acidophilus		6.75	0.00				
07771320	**Very good identification**						**07771320**
					SPO	GRA	
Clostridium ramosum		97.55	0.72		92	50	
Prevotella oralis		1.10	0.39		1	1	
Lactobacillus acidophilus		0.53	0.34		1	99	
07771321	**Low discrimination**						**07771321**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO	GRA	
Clostridium ramosum		57.48	0.38	URE 1	92	50	
Actinomyces naeslundii		41.55	0.61		1	99	
Prevotella oralis		0.65	0.04		1	1	
07771322	**Low discrimination**						**07771322**
					SPO	GRA	
Clostridium ramosum		80.42	0.83		92	50	
Lactobacillus acidophilus		10.84	0.68		1	99	
Prevotella oralis		4.32	0.60		1	1	
07771323	**Low discrimination**						**07771323**
					SPO		
Clostridium ramosum		70.30	0.48	URE 1	92		
Actinomyces naeslundii		12.70	0.61		1		
Lactobacillus acidophilus		9.47	0.34		1		
07771330	**Low discrimination**						**07771330**
					SPO		
Clostridium ramosum		83.88	0.38	MLZ 1	92		
Parabacteroides distasonis		9.31	0.34		1		
Clostridium clostridioforme		2.72	0.19		66		
07771331	**Low discrimination**						**07771331**
					SPO		
Actinomyces naeslundii		79.02	0.39	RHA 1	1		
Clostridium ramosum		20.98	0.04	MLZ 5	92		
07771332	**Low discrimination**						**07771332**
					SPO		
Parabacteroides distasonis		57.53	0.68	XYL 90	1		
Clostridium ramosum		20.93	0.48		92		
Clostridium clostridioforme		16.82	0.53		66		
07771340	**Low discrimination**						**07771340**
Actinomyces naeslundii		34.11	0.45	RHA 1			
Actinomyces israelii		31.00	0.35				
Lactobacillus acidophilus		26.73	0.19				
07771341	**Good identification**						**07771341**
Actinomyces naeslundii		100.0	0.48	RHA 1			
07771342	**Good identification**						**07771342**
Lactobacillus acidophilus		95.63	0.53	RHA 7			
Actinomyces israelii		1.68	0.38	SOR 11			
Actinomyces naeslundii		1.23	0.45				
07771343	**Low discrimination**						**07771343**
Actinomyces naeslundii		65.69	0.48	RHA 1			
Lactobacillus acidophilus		34.31	0.19				
07771360	**Low discrimination**						**07771360**
					SPO		
Clostridium ramosum		84.38	0.38	SOR 1	92		
Actinomyces naeslundii		7.18	0.45		1		
Lactobacillus acidophilus		5.62	0.19		1		
07771361	**Acceptable identification**						**07771361**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		92.66	0.48	RHA 1	1		
Clostridium ramosum		7.34	0.04		92		
07771362	**Low discrimination**						**07771362**
					SPO	GRA	
Lactobacillus acidophilus		58.93	0.53	RHA 7	1	99	
Clostridium ramosum		35.74	0.48	SOR 11	92	50	
Prevotella oralis		1.92	0.26		1	1	
07771363	**Low discrimination**						**07771363**
					SPO		
Actinomyces naeslundii		54.38	0.48	RHA 1	1		
Lactobacillus acidophilus		28.40	0.19		1		
Clostridium ramosum		17.22	0.14		92		
07771400	**Low discrimination**						**07771400**
					SPO		
Clostridium butyricum		58.80	0.36	ESL 88	99		
Actinomyces israelii		22.24	0.50	XYL 97	1		
Lactobacillus jensenii		10.58	0.32	LAC 96	1		
07771401	**Acceptable identification**						**07771401**
					SPO		
Actinomyces naeslundii		92.74	0.53	ARA 1	1		
Clostridium butyricum		5.27	0.02		99		
Actinomyces israelii		1.99	0.15		1		
07771402	**Low discrimination**						**07771402**
					SPO		
Actinomyces israelii		26.93	0.53	XYL 92	1		
Lactobacillus acidophilus		20.70	0.35		1		
Clostridium butyricum		18.46	0.29		99		
07771403	**Acceptable identification**						**07771403**
Actinomyces naeslundii		91.16	0.53	ARA 1			
Actinomyces israelii		2.94	0.18				
Lactobacillus acidophilus		2.26	0.01				
07771412	**Low discrimination**						**07771412**
Clostridium clostridioforme		80.14	0.50	XYL 91			
Bifidobacterium adolescentis		10.98	0.38	RHA 88			
Actinomyces israelii		4.73	0.38				
07771420	**Low discrimination**						**07771420**
					SPO		
Actinomyces naeslundii		40.48	0.50	ARA 1	1		
Clostridium butyricum		29.69	0.18		99		
Lactobacillus jensenii		15.36	0.21		1		
07771421	**Good identification**						**07771421**
Actinomyces naeslundii		100.0	0.53	ARA 1			
07771422	**Low discrimination**						**07771422**
Lactobacillus acidophilus		41.81	0.35	LAC 92			
Clostridium clostridioforme		30.22	0.39	ARA 1			
Actinomyces naeslundii		11.37	0.50				
07771423	**Good identification**						**07771423**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.91	0.53	ARA 1		
Lactobacillus acidophilus		2.38	0.01			
Clostridium clostridioforme		1.72	0.05			
07771432	**Good identification**					**07771432**
					GRA	
Clostridium clostridioforme		97.16	0.50	XYL 91	50	
Parabacteroides distasonis		1.08	0.22	RHA 88	1	
Lactobacillus jensenii		0.68	0.10		99	
07771440	**Low discrimination**					**07771440**
					SPO	
Actinomyces israelii		57.69	0.43	XYL 92	1	
Clostridium butyricum		34.11	0.18		99	
Actinomyces naeslundii		8.21	0.37		1	
07771441	**Good identification**					**07771441**
Actinomyces naeslundii		100.0	0.40	ARA 1		
07771442	**Low discrimination**					**07771442**
					SPO	
Actinomyces israelii		62.62	0.46	XYL 92	1	
Lactobacillus acidophilus		15.30	0.20		1	
Clostridium butyricum		9.60	0.11		99	
07771443	**Good identification**					**07771443**
Actinomyces naeslundii		100.0	0.40	ARA 1		
07771460	**Low discrimination**					**07771460**
					SPO	
Actinomyces naeslundii		55.24	0.37	ARA 1	1	
Actinomyces israelii		24.79	0.22		1	
Clostridium butyricum		19.97	-.01		99	
07771461	**Good identification**					**07771461**
Actinomyces naeslundii		100.0	0.40	ARA 1		
07771463	**Good identification**					**07771463**
Actinomyces naeslundii		100.0	0.40	ARA 1		
07771500	**Good identification**					**07771500**
					SPO	
Clostridium butyricum		91.74	0.59	ESL 88	99	
Actinomyces israelii		5.78	0.60	XYL 97	1	
Actinomyces naeslundii		1.29	0.58		1	
07771501	**Low discrimination**					**07771501**
					SPO	
Actinomyces naeslundii		66.26	0.61	ARA 1	1	
Clostridium butyricum		31.74	0.25		99	
Actinomyces israelii		2.00	0.26		1	
07771502	**Low discrimination**					**07771502**
					SPO	
Bifidobacterium adolescentis		46.78	0.71	XYL 87	1	
Clostridium butyricum		27.41	0.52		99	
Lactobacillus acidophilus		14.73	0.54		1	
07771503	**Low discrimination**					**07771503**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		59.86	0.61	ARA 1	1		
Bifidobacterium adolescentis		19.03	0.37		1		
Clostridium butyricum		11.15	0.18		99		
07771510	**Low discrimination**						**07771510**
					SPO		
Clostridium butyricum		71.96	0.38	ESL 88	99		
Actinomyces israelii		12.88	0.46	XYL 97	1		
Bifidobacterium adolescentis		11.20	0.38	MLZ 5	1		
07771511	**Low discrimination**						**07771511**
					SPO		
Actinomyces naeslundii		67.62	0.39	ARA 1	1		
Clostridium butyricum		32.39	0.03	MLZ 5	99		
07771512	**Low discrimination**						**07771512**
					SPO		
Bifidobacterium adolescentis		76.20	0.73	XYL 87	1		
Clostridium clostridioforme		18.81	0.59		66		
Clostridium butyricum		1.92	0.30		99		
07771513	**Low discrimination**						**07771513**
Bifidobacterium adolescentis		69.35	0.38	XYL 87			
Clostridium clostridioforme		17.12	0.25	URE 1			
Actinomyces naeslundii		9.40	0.39				
07771520	**Low discrimination**						**07771520**
					SPO		
Clostridium butyricum		64.55	0.41	ESL 88	99		
Clostridium ramosum		16.34	0.36	XYL 97	92		
Actinomyces naeslundii		10.44	0.58	NAG 8	1		
07771521	**Very good identification**						**07771521**
					SPO		
Actinomyces naeslundii		95.04	0.61	ARA 1	1		
Clostridium butyricum		3.96	0.07		99		
Clostridium ramosum		1.00	0.02		92		
07771522	**Low discrimination**						**07771522**
					SPO	GRA	
Lactobacillus acidophilus		38.58	0.54	ARA 1	1	99	
Parabacteroides distasonis		22.47	0.61		1	1	
Clostridium ramosum		16.26	0.46		92	50	
07771523	**Low discrimination**						**07771523**
					GRA		
Actinomyces naeslundii		80.34	0.61	ARA 1	99		
Lactobacillus acidophilus		8.04	0.19		99		
Parabacteroides distasonis		4.68	0.27		1		
07771531	**Good identification**						**07771531**
Actinomyces naeslundii		100.0	0.39	ARA 1			
07771532	**Low discrimination**						**07771532**
					GRA		
Clostridium clostridioforme		70.28	0.59	XYL 91	50		
Parabacteroides distasonis		22.99	0.56	RHA 88	1		
Bifidobacterium adolescentis		2.88	0.38		99		
07771540	**Low discrimination**						**07771540**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		75.59	0.41	ESL 88	99	
Actinomyces israelii		21.31	0.53	XYL 97	1	
Actinomyces naeslundii		2.16	0.45	SOR 8	1	
07771541	**Low discrimination**					**07771541**
					SPO	
Actinomyces naeslundii		76.77	0.48	ARA 1	1	
Clostridium butyricum		18.12	0.07		99	
Actinomyces israelii		5.11	0.19		1	
07771542	**Low discrimination**					**07771542**
					SPO	
Bifidobacterium adolescentis		44.96	0.56	XYL 87	1	
Actinomyces israelii		20.15	0.56	SOR 11	1	
Clostridium butyricum		18.53	0.34		99	
07771543	**Low discrimination**					**07771543**
Actinomyces naeslundii		67.37	0.48	ARA 1		
Bifidobacterium adolescentis		15.00	0.21			
Actinomyces israelii		6.72	0.22			
07771552	**Good identification**					**07771552**
Bifidobacterium adolescentis		90.17	0.57	XYL 87		
Actinomyces israelii		4.94	0.42	SOR 11		
Clostridium clostridioforme		1.82	0.25			
07771561	**Good identification**					**07771561**
Actinomyces naeslundii		100.0	0.48	ARA 1		
07771562	**Low discrimination**					**07771562**
					SPO	
Lactobacillus acidophilus		65.83	0.38	ARA 1	1	
Clostridium butyricum		7.50	0.16	SOR 11	99	
Actinomyces naeslundii		6.33	0.45		1	
07771563	**Acceptable identification**					**07771563**
Actinomyces naeslundii		93.45	0.48	ARA 1		
Lactobacillus acidophilus		6.55	0.04			
07771602	**Good identification**					**07771602**
Clostridium clostridioforme		96.35	0.54	XYL 91		
Actinomyces israelii		1.78	0.33			
Lactobacillus acidophilus		1.37	0.16			
07771612	**Very good identification**					**07771612**
Clostridium clostridioforme		99.94	0.65	XYL 91		
Actinomyces israelii		0.06	0.19			
07771622	**Acceptable identification**					**07771622**
					GRA	
Clostridium clostridioforme		93.56	0.54	XYL 91	50	
Parabacteroides distasonis		3.58	0.35		1	
Lactobacillus acidophilus		1.33	0.16		99	
07771632	**Very good identification**					**07771632**
Clostridium clostridioforme		99.55	0.65	XYL 91		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		0.45	0.30			
07771702	**Low discrimination**					**07771702**
					SPO	
Clostridium clostridioforme		87.30	0.63	XYL 91	66	
Lactobacillus acidophilus		4.26	0.34		1	
Clostridium butyricum		3.26	0.27		99	
07771710	**Low discrimination**					**07771710**
					SPO	
Clostridium clostridioforme		86.39	0.40	XYL 91	66	
Clostridium butyricum		9.48	0.12	BGL 99	99	
Actinomyces israelii		4.13	0.27		1	
07771712	**Very good identification**					**07771712**
					GRA	
Clostridium clostridioforme		99.18	0.74	XYL 91	50	
Bifidobacterium adolescentis		0.55	0.38		99	
Parabacteroides distasonis		0.13	0.30		1	
07771713	**Good identification**					**07771713**
Clostridium clostridioforme		100.0	0.40	XYL 91		
07771720	**Low discrimination**					**07771720**
					SPO	
Clostridium ramosum		71.05	0.38	ARA 1	92	
Parabacteroides distasonis		8.98	0.35		1	
Clostridium clostridioforme		7.93	0.28		66	
07771722	**Low discrimination**					**07771722**
					SPO	
Parabacteroides distasonis		43.71	0.69	XYL 90	1	
Clostridium clostridioforme		38.60	0.63		66	
Clostridium ramosum		13.97	0.48		92	
07771723	**Low discrimination**					**07771723**
					SPO	
Parabacteroides distasonis		43.41	0.35	XYL 90	1	
Clostridium clostridioforme		38.34	0.28	URE 1	66	
Clostridium ramosum		13.88	0.14		92	
07771730	**Low discrimination**					**07771730**
					SPO	
Clostridium clostridioforme		85.97	0.40	XYL 91	66	
Parabacteroides distasonis		11.51	0.30	BGL 99	1	
Clostridium ramosum		1.71	0.04		92	
07771732	**Low discrimination**					**07771732**
Clostridium clostridioforme		87.88	0.74	XYL 91		
Parabacteroides distasonis		11.76	0.65			
Bacteroides ovatus		0.25	0.20			
07771733	**Low discrimination**					**07771733**
Clostridium clostridioforme		88.20	0.40	XYL 91		
Parabacteroides distasonis		11.81	0.30	URE 1		
07771752	**Acceptable identification**					**07771752**
Clostridium clostridioforme		91.23	0.40	XYL 91		
Bifidobacterium adolescentis		6.23	0.23	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		2.54	0.23				
07771762	**Low discrimination**						**07771762**
					GRA		
Parabacteroides distasonis		35.12	0.35	XYL 90	1		
Clostridium clostridioforme		31.02	0.28	SOR 1	50		
Lactobacillus acidophilus		18.52	0.19		99		
07771772	**Low discrimination**						**07771772**
Clostridium clostridioforme		87.31	0.40	XYL 91			
Parabacteroides distasonis		11.69	0.30	SOR 1			
Bacteroides ovatus		1.00	-.04				
07772000	**Good identification**						**07772000**
					CAT		
Actinomyces naeslundii		93.42	0.66	GAL 96	25		
Actinomyces israelii		2.60	0.30		1		
Propionibacterium granulosum		2.44	0.24		90		
07772001	**Very good identification**						**07772001**
Actinomyces naeslundii		100.0	0.69	GAL 96			
07772002	**Good identification**						**07772002**
Actinomyces naeslundii		92.81	0.66	GAL 96			
Actinomyces israelii		3.88	0.33				
Lactobacillus acidophilus		2.24	0.14				
07772003	**Very good identification**						**07772003**
Actinomyces naeslundii		100.0	0.69	GAL 96			
07772010	**Low discrimination**						**07772010**
					CAT		
Actinomyces naeslundii		87.20	0.44	GAL 96	25		
Actinomyces israelii		6.90	0.16	MLZ 5	1		
Propionibacterium granulosum		5.90	0.09		90		
07772011	**Good identification**						**07772011**
Actinomyces naeslundii		100.0	0.47	GAL 96			
07772012	**Low discrimination**						**07772012**
Actinomyces naeslundii		89.39	0.44	GAL 96			
Actinomyces israelii		10.61	0.19	MLZ 5			
07772013	**Good identification**						**07772013**
Actinomyces naeslundii		100.0	0.47	GAL 96			
07772020	**Very good identification**						**07772020**
					SPO	CAT	
Actinomyces naeslundii		96.30	0.66	GAL 96	1	25	
Clostridium ramosum		3.07	0.15		92	1	
Propionibacterium granulosum		0.63	0.13		1	90	
07772021	**Very good identification**						**07772021**
Actinomyces naeslundii		100.0	0.69	GAL 96			
07772022	**Low discrimination**						**07772022**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		86.84	0.66	GAL 96	1	
Clostridium ramosum		11.06	0.25		92	
Lactobacillus acidophilus		2.10	0.14		1	
07772023	**Very good identification**					**07772023**
Actinomyces naeslundii		100.0	0.69	GAL 96		
07772030	**Good identification**					**07772030**
Actinomyces naeslundii		100.0	0.44	GAL 96		
07772031	**Good identification**					**07772031**
Actinomyces naeslundii		100.0	0.47	GAL 96		
07772032	**Good identification**					**07772032**
Actinomyces naeslundii		100.0	0.44	GAL 96		
07772033	**Good identification**					**07772033**
Actinomyces naeslundii		100.0	0.47	GAL 96		
07772040	**Acceptable identification**					**07772040**
Actinomyces naeslundii		94.21	0.53	GAL 96		
Actinomyces israelii		5.79	0.23			
07772041	**Good identification**					**07772041**
Actinomyces naeslundii		100.0	0.56	GAL 96		
07772042	**Acceptable identification**					**07772042**
Actinomyces naeslundii		90.17	0.53	GAL 96		
Actinomyces israelii		8.31	0.26			
Lactobacillus acidophilus		1.53	-.02			
07772043	**Good identification**					**07772043**
Actinomyces naeslundii		100.0	0.56	GAL 96		
07772060	**Good identification**					**07772060**
Actinomyces naeslundii		100.0	0.53	GAL 96		
07772061	**Good identification**					**07772061**
Actinomyces naeslundii		100.0	0.56	GAL 96		
07772062	**Good identification**					**07772062**
Actinomyces naeslundii		98.34	0.53	GAL 96		
Lactobacillus acidophilus		1.66	-.02			
07772063	**Good identification**					**07772063**
Actinomyces naeslundii		100.0	0.56	GAL 96		
07772100	**Good identification**					**07772100**
					SPO	
Actinomyces naeslundii		94.14	0.73	GAL 96	1	
Actinomyces israelii		3.69	0.40		1	
Clostridium ramosum		1.05	0.15		92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07772101	**Very good identification**					**07772101**
Actinomyces naeslundii		100.0	0.76	GAL 96		
07772102	**Low discrimination**					**07772102**
Actinomyces naeslundii		81.75	0.73	GAL 96		
Lactobacillus acidophilus		7.99	0.32			
Actinomyces israelii		4.80	0.43			
07772103	**Very good identification**					**07772103**
Actinomyces naeslundii		100.0	0.76	GAL 96		
07772110	**Low discrimination**					**07772110**
					SPO	
Actinomyces naeslundii		86.52	0.52	GAL 96	1	
Actinomyces israelii		9.62	0.26	MLZ 5	1	
Clostridium tertium		3.86	0.05		99	
07772111	**Good identification**					**07772111**
Actinomyces naeslundii		100.0	0.55	GAL 96		
07772112	**Low discrimination**					**07772112**
Actinomyces naeslundii		59.59	0.52	GAL 96		
Bifidobacterium breve		11.04	0.24	MLZ 5		
Bifidobacterium adolescentis		10.07	0.23			
07772113	**Good identification**					**07772113**
Actinomyces naeslundii		100.0	0.55	GAL 96		
07772120	**Low discrimination**					**07772120**
					SPO	
Clostridium ramosum		52.35	0.49	GAL 99	92	
Actinomyces naeslundii		47.26	0.73		1	
Clostridium tertium		0.19	0.09		99	
07772121	**Very good identification**					**07772121**
					SPO	
Actinomyces naeslundii		99.26	0.76	GAL 96	1	
Clostridium ramosum		0.74	0.15		92	
07772122	**Low discrimination**					**07772122**
					SPO	
Clostridium ramosum		79.65	0.59	GAL 99	92	
Actinomyces naeslundii		17.98	0.73		1	
Lactobacillus acidophilus		1.76	0.32		1	
07772123	**Very good identification**					**07772123**
					SPO	
Actinomyces naeslundii		97.10	0.76	GAL 96	1	
Clostridium ramosum		2.90	0.25		92	
07772130	**Low discrimination**					**07772130**
					SPO	
Actinomyces naeslundii		72.52	0.52	GAL 96	1	
Clostridium ramosum		15.42	0.15	MLZ 5	92	
Clostridium tertium		9.69	0.13		99	
07772131	**Good identification**					**07772131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	GAL 96		
07772132	**Low discrimination**					**07772132**
					SPO	
Actinomyces naeslundii		38.75	0.52	GAL 96	1	
Clostridium ramosum		32.95	0.25	MLZ 5	92	
Clostridium tertium		15.54	0.21		99	
07772133	**Good identification**					**07772133**
Actinomyces naeslundii		100.0	0.55	GAL 96		
07772140	**Good identification**					**07772140**
Actinomyces naeslundii		92.05	0.61	GAL 96		
Actinomyces israelii		7.95	0.33			
07772141	**Very good identification**					**07772141**
Actinomyces naeslundii		100.0	0.64	GAL 96		
07772142	**Low discrimination**					**07772142**
Actinomyces naeslundii		79.80	0.61	GAL 96		
Actinomyces israelii		10.33	0.36			
Lactobacillus acidophilus		5.46	0.16			
07772143	**Very good identification**					**07772143**
Actinomyces naeslundii		100.0	0.64	GAL 96		
07772150	**Low discrimination**					**07772150**
Actinomyces naeslundii		80.32	0.39	GAL 96		
Actinomyces israelii		19.68	0.19	MLZ 5		
07772151	**Good identification**					**07772151**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772153	**Good identification**					**07772153**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772160	**Good identification**					**07772160**
					SPO	
Actinomyces naeslundii		93.19	0.61	GAL 96	1	
Clostridium ramosum		5.91	0.15		92	
Bifidobacterium breve		0.91	0.12		1	
07772161	**Very good identification**					**07772161**
Actinomyces naeslundii		100.0	0.64	GAL 96		
07772162	**Low discrimination**					**07772162**
					SPO	
Actinomyces naeslundii		70.00	0.61	GAL 96	1	
Clostridium ramosum		17.76	0.25		92	
Bifidobacterium breve		6.88	0.29		1	
07772163	**Very good identification**					**07772163**
Actinomyces naeslundii		100.0	0.64	GAL 96		
07772170	**Low discrimination**					**07772170**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.42	0.39	GAL 96		
Bifidobacterium breve		15.58	0.12	MLZ 5		
07772171	**Good identification**					**07772171**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772173	**Good identification**					**07772173**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772301	**Good identification**					**07772301**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772303	**Good identification**					**07772303**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772320	**Good identification**					**07772320**
					SPO	
Clostridium ramosum		99.32	0.51	GAL 99	92	
Actinomyces naeslundii		0.68	0.39		1	
07772321	**Low discrimination**					**07772321**
					SPO	
Actinomyces naeslundii		50.53	0.42	GAL 96	1	
Clostridium ramosum		49.47	0.17	RHA 1	92	
07772322	**Very good identification**					**07772322**
					SPO	
Clostridium ramosum		99.70	0.61	GAL 99	92	
Actinomyces naeslundii		0.17	0.39		1	
Lactobacillus acidophilus		0.13	0.12		1	
07772500	**Low discrimination**					**07772500**
					SPO	
Actinomyces israelii		61.49	0.39	GAL 99	1	
Actinomyces naeslundii		19.38	0.39	XYL 92	1	
Clostridium butyricum		19.13	0.09		99	
07772501	**Good identification**					**07772501**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772502	**Low discrimination**					**07772502**
Bifidobacterium adolescentis		43.47	0.35	GAL 99		
Actinomyces israelii		42.89	0.42	XYL 87		
Actinomyces naeslundii		9.01	0.39			
07772503	**Good identification**					**07772503**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772512	**Low discrimination**					**07772512**
Bifidobacterium adolescentis		79.81	0.37	GAL 99		
Actinomyces israelii		9.63	0.28	XYL 87		
Clostridium clostridioforme		8.03	0.17			
07772521	**Good identification**					**07772521**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07772523	**Good identification**					**07772523**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	GAL 96		
07773000	**Low discrimination**					**07773000**
Actinomyces naeslundii		88.37	0.89			
Actinomyces israelii		10.16	0.64			
Lactobacillus jensenii		0.70	0.32			
07773001	**Excellent identification**					**07773001**
Actinomyces naeslundii		99.92	0.92			
Actinomyces israelii		0.08	0.30			
07773002	**Low discrimination**					**07773002**
Actinomyces naeslundii		77.53	0.89			
Actinomyces israelii		13.37	0.67			
Lactobacillus acidophilus		7.73	0.48			
07773003	**Excellent identification**					**07773003**
Actinomyces naeslundii		99.82	0.92			
Actinomyces israelii		0.12	0.33			
Lactobacillus acidophilus		0.07	0.14			
07773010	**Low discrimination**					**07773010**
Actinomyces naeslundii		72.07	0.67	MLZ 5		
Actinomyces israelii		23.52	0.50			
Lactobacillus jensenii		3.61	0.24			
07773011	**Very good identification**					**07773011**
Actinomyces naeslundii		99.78	0.70	MLZ 5		
Actinomyces israelii		0.22	0.16			
07773012	**Low discrimination**					**07773012**
Actinomyces naeslundii		62.27	0.67	MLZ 5		
Actinomyces israelii		30.49	0.53			
Lactobacillus jensenii		2.08	0.21			
07773013	**Very good identification**					**07773013**
Actinomyces naeslundii		99.67	0.70	MLZ 5		
Actinomyces israelii		0.33	0.19			
07773020	**Low discrimination**					**07773020**
					SPO	
Actinomyces naeslundii		87.48	0.89		1	
Clostridium ramosum		11.49	0.49		92	
Actinomyces israelii		0.64	0.44		1	
07773021	**Excellent identification**					**07773021**
					SPO	
Actinomyces naeslundii		99.91	0.92		1	
Clostridium ramosum		0.09	0.15		92	
07773022	**Low discrimination**					**07773022**
					SPO	
Actinomyces naeslundii		60.88	0.89		1	
Clostridium ramosum		31.99	0.59		92	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		6.07	0.48		1	
07773023	**Excellent identification**					**07773023**
					SPO	
Actinomyces naeslundii		99.58	0.92		1	
Clostridium ramosum		0.35	0.25		92	
Lactobacillus acidophilus		0.07	0.14		1	
07773030	**Good identification**					**07773030**
					SPO	
Actinomyces naeslundii		92.74	0.67	MLZ 5	1	
Clostridium ramosum		2.34	0.15		92	
Actinomyces israelii		1.93	0.29		1	
07773031	**Very good identification**					**07773031**
Actinomyces naeslundii		100.0	0.70	MLZ 5		
07773032	**Low discrimination**					**07773032**
					SPO	
Actinomyces naeslundii		80.24	0.67	MLZ 5	1	
Clostridium ramosum		8.09	0.25		92	
Clostridium tertium		3.82	0.21		99	
07773033	**Very good identification**					**07773033**
Actinomyces naeslundii		100.0	0.70	MLZ 5		
07773040	**Low discrimination**					**07773040**
Actinomyces naeslundii		79.74	0.76			
Actinomyces israelii		20.20	0.57			
Lactobacillus acidophilus		0.06	-.02			
07773041	**Very good identification**					**07773041**
Actinomyces naeslundii		99.83	0.79			
Actinomyces israelii		0.17	0.23			
07773042	**Low discrimination**					**07773042**
Actinomyces naeslundii		68.66	0.76			
Actinomyces israelii		26.09	0.60			
Lactobacillus acidophilus		4.80	0.32			
07773043	**Very good identification**					**07773043**
Actinomyces naeslundii		99.70	0.79			
Actinomyces israelii		0.26	0.26			
Lactobacillus acidophilus		0.05	-.02			
07773050	**Low discrimination**					**07773050**
Actinomyces naeslundii		58.16	0.54	MLZ 5		
Actinomyces israelii		41.84	0.43			
07773051	**Very good identification**					**07773051**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
07773052	**Low discrimination**					**07773052**
Actinomyces israelii		48.36	0.46	XYL 92		
Actinomyces naeslundii		44.83	0.54	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		5.58	0.24			
07773053	**Very good identification**					**07773053**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
07773060	**Very good identification**					**07773060**
					SPO	
Actinomyces naeslundii		97.51	0.76		1	
Actinomyces israelii		1.58	0.36		1	
Clostridium ramosum		0.73	0.15		92	
07773061	**Very good identification**					**07773061**
Actinomyces naeslundii		100.0	0.79			
07773062	**Low discrimination**					**07773062**
					SPO	
Actinomyces naeslundii		88.04	0.76		1	
Lactobacillus acidophilus		6.15	0.32		1	
Clostridium ramosum		2.65	0.25		92	
07773063	**Very good identification**					**07773063**
Actinomyces naeslundii		99.95	0.79			
Lactobacillus acidophilus		0.05	-.02			
07773070	**Acceptable identification**					**07773070**
Actinomyces naeslundii		93.65	0.54	MLZ 5		
Actinomyces israelii		4.30	0.22			
Bifidobacterium breve		2.05	0.12			
07773071	**Very good identification**					**07773071**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
07773072	**Low discrimination**					**07773072**
Actinomyces naeslundii		76.71	0.54	MLZ 5		
Bifidobacterium breve		16.98	0.29			
Actinomyces israelii		5.28	0.25			
07773073	**Very good identification**					**07773073**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
07773100	**Low discrimination**					**07773100**
					SPO	
Actinomyces naeslundii		81.47	0.97		1	
Actinomyces israelii		13.16	0.74		1	
Clostridium ramosum		3.76	0.49		92	
07773101	**Excellent identification**					**07773101**
					SPO	
Actinomyces naeslundii		99.85	1.00		1	
Actinomyces israelii		0.11	0.40		1	
Clostridium ramosum		0.03	0.15		92	
07773102	**Low discrimination**					**07773102**
Actinomyces naeslundii		52.11	0.97			
Lactobacillus acidophilus		21.00	0.66			
Actinomyces israelii		12.63	0.77			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
07773103	**Excellent identification**					**07773103**
Actinomyces naeslundii		99.39	1.00			
Lactobacillus acidophilus		0.27	0.32			
Actinomyces israelii		0.16	0.43			
07773110	**Low discrimination**					**07773110**
					SPO	
Actinomyces naeslundii		57.49	0.75	MLZ 5	1	
Actinomyces israelii		26.37	0.60		1	
Clostridium tertium		10.57	0.39		99	
07773111	**Very good identification**					**07773111**
					SPO	
Actinomyces naeslundii		99.57	0.78	MLZ 5	1	
Actinomyces israelii		0.31	0.26		1	
Clostridium tertium		0.12	0.05		99	
07773112	**Low discrimination**					**07773112**
Actinomyces naeslundii		25.55	0.75	MLZ 5		
Bifidobacterium breve		19.52	0.59			
Bifidobacterium adolescentis		17.82	0.57			
07773113	**Very good identification**					**07773113**
Actinomyces naeslundii		98.15	0.78	MLZ 5		
Bifidobacterium breve		0.51	0.24			
Bifidobacterium adolescentis		0.46	0.23			
07773120	**Low discrimination**					**07773120**
					SPO	
Clostridium ramosum		81.47	0.83		92	
Actinomyces naeslundii		17.83	0.97		1	
Clostridium tertium		0.29	0.43		99	
07773121	**Very good identification**					**07773121**
					SPO	
Actinomyces naeslundii		96.99	1.00		1	
Clostridium ramosum		2.99	0.49		92	
Actinomyces israelii		0.01	0.20		1	
07773122	**Good identification**					**07773122**
					SPO	
Clostridium ramosum		92.20	0.93		92	
Actinomyces naeslundii		5.04	0.97		1	
Lactobacillus acidophilus		2.03	0.66		1	
07773123	**Low discrimination**					**07773123**
					SPO	
Actinomyces naeslundii		88.75	1.00		1	
Clostridium ramosum		10.92	0.59		92	
Lactobacillus acidophilus		0.24	0.32		1	
07773130	**Low discrimination**					**07773130**
					SPO	
Actinomyces naeslundii		38.58	0.75	MLZ 5	1	
Clostridium ramosum		33.84	0.49		92	
Clostridium tertium		21.27	0.47		99	
07773131	**Very good identification**					**07773131**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.96	0.78	MLZ 5	1	
Clostridium ramosum		0.58	0.15		92	
Clostridium tertium		0.37	0.13		99	
07773132	**Low discrimination**					**07773132**
					SPO	
Clostridium ramosum		45.73	0.59	MLZ 1	92	
Clostridium tertium		21.56	0.55		99	
Bifidobacterium breve		17.71	0.63		1	
07773133	**Very good identification**					**07773133**
					SPO	
Actinomyces naeslundii		95.79	0.78	MLZ 5	1	
Clostridium ramosum		2.26	0.25		92	
Clostridium tertium		1.07	0.21		99	
07773140	**Low discrimination**					**07773140**
Actinomyces naeslundii		72.12	0.84			
Actinomyces israelii		25.68	0.67			
Bifidobacterium breve		1.63	0.41			
07773141	**Very good identification**					**07773141**
Actinomyces naeslundii		99.76	0.87			
Actinomyces israelii		0.24	0.33			
07773142	**Low discrimination**					**07773142**
Actinomyces naeslundii		48.13	0.84			
Actinomyces israelii		25.71	0.70			
Lactobacillus acidophilus		13.58	0.50			
07773143	**Very good identification**					**07773143**
Actinomyces naeslundii		99.30	0.87			
Actinomyces israelii		0.36	0.36			
Lactobacillus acidophilus		0.19	0.16			
07773150	**Low discrimination**					**07773150**
Actinomyces israelii		41.04	0.53	XYL 92		
Actinomyces naeslundii		40.60	0.62	MLZ 13		
Bifidobacterium breve		17.39	0.41			
07773151	**Very good identification**					**07773151**
Actinomyces naeslundii		99.32	0.65	MLZ 5		
Actinomyces israelii		0.68	0.19			
07773152	**Low discrimination**					**07773152**
Bifidobacterium breve		58.31	0.59	ESL 99		
Actinomyces israelii		20.42	0.56			
Actinomyces naeslundii		13.47	0.62			
07773153	**Very good identification**					**07773153**
Actinomyces naeslundii		95.91	0.65	MLZ 5		
Bifidobacterium breve		2.80	0.24			
Actinomyces israelii		0.98	0.22			
07773160	**Low discrimination**					**07773160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		75.26	0.84		1	
Clostridium ramosum		19.68	0.49		92	
Bifidobacterium breve		3.02	0.46		1	
07773161	**Very good identification**					**07773161**
					SPO	
Actinomyces naeslundii		99.80	0.87		1	
Clostridium ramosum		0.18	0.15		92	
Bifidobacterium breve		0.03	0.12		1	
07773162	**Low discrimination**					**07773162**
					SPO	
Clostridium ramosum		37.75	0.59	SOR 1	92	
Actinomyces naeslundii		36.08	0.84		1	
Bifidobacterium breve		14.62	0.63		1	
07773163	**Very good identification**					**07773163**
					SPO	
Actinomyces naeslundii		98.82	0.87		1	
Clostridium ramosum		0.70	0.25		92	
Bifidobacterium breve		0.27	0.29		1	
07773170	**Low discrimination**					**07773170**
Actinomyces naeslundii		52.42	0.62	MLZ 5		
Bifidobacterium breve		39.91	0.46			
Actinomyces israelii		3.38	0.33			
07773171	**Very good identification**					**07773171**
Actinomyces naeslundii		99.49	0.65	MLZ 5		
Bifidobacterium breve		0.51	0.12			
07773172	**Low discrimination**					**07773172**
					SPO	
Bifidobacterium breve		84.14	0.63	ESL 99	1	
Actinomyces naeslundii		10.93	0.62		1	
Clostridium ramosum		2.19	0.25		92	
07773173	**Very good identification**					**07773173**
Actinomyces naeslundii		95.07	0.65	MLZ 5		
Bifidobacterium breve		4.93	0.29			
07773200	**Low discrimination**					**07773200**
					SPO	
Actinomyces naeslundii		49.24	0.55	RHA 1	1	
Actinomyces israelii		42.18	0.45		1	
Clostridium ramosum		8.58	0.17		92	
07773201	**Very good identification**					**07773201**
Actinomyces naeslundii		100.0	0.58	RHA 1		
07773202	**Low discrimination**					**07773202**
Actinomyces israelii		34.50	0.48	XYL 92		
Actinomyces naeslundii		26.85	0.55	RHA 7		
Lactobacillus acidophilus		19.95	0.28			
07773203	**Very good identification**					**07773203**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.14	0.58	RHA 1		
Actinomyces israelii		0.86	0.14			
07773211	**Good identification**					**07773211**
Actinomyces naeslundii		100.0	0.36	RHA 1		
07773213	**Good identification**					**07773213**
Actinomyces naeslundii		100.0	0.36	RHA 1		
07773220	**Acceptable identification**					**07773220**
					SPO	
Clostridium ramosum		94.24	0.51	LAC 99	92	
Actinomyces naeslundii		5.47	0.55		1	
Actinomyces israelii		0.30	0.24		1	
07773221	**Low discrimination**					**07773221**
					SPO	
Actinomyces naeslundii		89.60	0.58	RHA 1	1	
Clostridium ramosum		10.40	0.17		92	
07773222	**Very good identification**					**07773222**
					SPO	
Clostridium ramosum		97.36	0.61	LAC 99	92	
Actinomyces naeslundii		1.41	0.55		1	
Lactobacillus acidophilus		1.05	0.28		1	
07773223	**Low discrimination**					**07773223**
					SPO	
Actinomyces naeslundii		68.29	0.58	RHA 1	1	
Clostridium ramosum		31.71	0.27		92	
07773231	**Good identification**					**07773231**
Actinomyces naeslundii		100.0	0.36	RHA 1		
07773233	**Good identification**					**07773233**
Actinomyces naeslundii		100.0	0.36	RHA 1		
07773240	**Low discrimination**					**07773240**
Actinomyces israelii		65.37	0.38	XYL 92		
Actinomyces naeslundii		34.63	0.42	RHA 7		
07773241	**Good identification**					**07773241**
Actinomyces naeslundii		100.0	0.45	RHA 1		
07773242	**Low discrimination**					**07773242**
Actinomyces israelii		65.07	0.41	XYL 92		
Actinomyces naeslundii		22.98	0.42	RHA 7		
Lactobacillus acidophilus		11.96	0.13			
07773243	**Good identification**					**07773243**
Actinomyces naeslundii		100.0	0.45	RHA 1		
07773260	**Low discrimination**					**07773260**
					SPO	
Actinomyces naeslundii		47.45	0.42	RHA 1	1	
Clostridium ramosum		46.83	0.17		92	
Actinomyces israelii		5.72	0.17		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07773261	**Good identification**					**07773261**
Actinomyces naeslundii		100.0	0.45	RHA 1		
07773263	**Good identification**					**07773263**
Actinomyces naeslundii		100.0	0.45	RHA 1		
07773300	**Low discrimination**					**07773300**
					SPO	
Clostridium ramosum		72.88	0.51	NAG 99	92	
Actinomyces israelii		14.49	0.55		1	
Actinomyces naeslundii		12.03	0.63		1	
07773301	**Very good identification**					**07773301**
					SPO	
Actinomyces naeslundii		95.34	0.66	RHA 1	1	
Clostridium ramosum		3.89	0.17		92	
Actinomyces israelii		0.77	0.21		1	
07773302	**Low discrimination**					**07773302**
					SPO	
Clostridium ramosum		80.36	0.61	NAG 99	92	
Lactobacillus acidophilus		9.96	0.47		1	
Actinomyces israelii		5.99	0.58		1	
07773303	**Low discrimination**					**07773303**
					SPO	
Actinomyces naeslundii		83.64	0.66	RHA 1	1	
Clostridium ramosum		13.65	0.27		92	
Lactobacillus acidophilus		1.69	0.12		1	
07773310	**Low discrimination**					**07773310**
					SPO	
Actinomyces israelii		59.30	0.41	XYL 92	1	
Clostridium ramosum		20.17	0.17	RHA 7	92	
Actinomyces naeslundii		17.35	0.41	MLZ 13	1	
07773311	**Good identification**					**07773311**
Actinomyces naeslundii		100.0	0.44	RHA 1		
07773312	**Low discrimination**					**07773312**
					SPO	
Actinomyces israelii		36.31	0.44	XYL 92	1	
Clostridium ramosum		32.93	0.27	RHA 7	92	
Actinomyces naeslundii		7.08	0.41	MLZ 13	1	
07773313	**Good identification**					**07773313**
Actinomyces naeslundii		100.0	0.44	RHA 1		
07773320	**Very good identification**					**07773320**
					SPO	
Clostridium ramosum		99.81	0.85		92	
Actinomyces naeslundii		0.17	0.63		1	
Actinomyces israelii		0.01	0.35		1	
07773321	**Low discrimination**					**07773321**
					SPO	
Clostridium ramosum		80.15	0.51	URE 1	92	
Actinomyces naeslundii		19.85	0.66		1	
07773322	**Excellent identification**					**07773322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium ramosum		99.80	0.95		92		
Lactobacillus acidophilus		0.13	0.47		1		
Actinomyces naeslundii		0.04	0.63		1		
07773323	**Good identification**						**07773323**
					SPO		
Clostridium ramosum		94.06	0.61	URE 1	92		
Actinomyces naeslundii		5.82	0.66		1		
Lactobacillus acidophilus		0.12	0.12		1		
07773330	**Good identification**						**07773330**
					SPO		
Clostridium ramosum		98.37	0.51	MLZ 1	92		
Actinomyces naeslundii		0.86	0.41		1		
Clostridium tertium		0.47	0.13		99		
07773331	**Low discrimination**						**07773331**
					SPO		
Actinomyces naeslundii		56.34	0.44	RHA 1	1		
Clostridium ramosum		43.66	0.17	MLZ 5	92		
07773332	**Very good identification**						**07773332**
					SPO	GRA	
Clostridium ramosum		98.33	0.61	MLZ 1	92	50	
Parabacteroides distasonis		0.48	0.34		1	1	
Clostridium tertium		0.35	0.21		99	99	
07773340	**Low discrimination**						**07773340**
					SPO		
Actinomyces israelii		66.33	0.48	XYL 92	1		
Actinomyces naeslundii		25.00	0.50	RHA 7	1		
Clostridium ramosum		8.67	0.17		92		
07773341	**Good identification**						**07773341**
Actinomyces naeslundii		98.24	0.53	RHA 1			
Actinomyces israelii		1.76	0.14				
07773342	**Low discrimination**						**07773342**
					SPO		
Actinomyces israelii		45.76	0.51	XYL 92	1		
Lactobacillus acidophilus		24.18	0.31	RHA 7	1		
Clostridium ramosum		15.95	0.27		92		
07773343	**Good identification**						**07773343**
Actinomyces naeslundii		97.39	0.53	RHA 1			
Actinomyces israelii		2.61	0.17				
07773352	**Low discrimination**						**07773352**
Actinomyces israelii		66.01	0.37	XYL 92			
Bifidobacterium breve		25.30	0.24	RHA 7			
Actinomyces naeslundii		5.84	0.28	MLZ 13			
07773360	**Good identification**						**07773360**
					SPO		
Clostridium ramosum		96.60	0.51	SOR 1	92		
Actinomyces naeslundii		2.81	0.50		1		
Actinomyces israelii		0.48	0.28		1		
07773361	**Low discrimination**						**07773361**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		81.23	0.53	RHA 1	1	
Clostridium ramosum		18.78	0.17		92	
07773362	**Very good identification**					**07773362**
					SPO	
Clostridium ramosum		97.33	0.61	SOR 1	92	
Lactobacillus acidophilus		1.49	0.31		1	
Actinomyces naeslundii		0.71	0.50		1	
07773363	**Low discrimination**					**07773363**
					SPO	
Actinomyces naeslundii		51.96	0.53	RHA 1	1	
Clostridium ramosum		48.04	0.27		92	
07773400	**Low discrimination**					**07773400**
					SPO	
Actinomyces israelii		85.72	0.62	XYL 92	1	
Actinomyces naeslundii		9.21	0.55		1	
Clostridium butyricum		4.45	0.19		99	
07773401	**Good identification**					**07773401**
Actinomyces naeslundii		94.10	0.58	ARA 1		
Actinomyces israelii		5.90	0.28			
07773402	**Low discrimination**					**07773402**
Actinomyces israelii		87.80	0.65	XYL 92		
Actinomyces naeslundii		6.29	0.55			
Bifidobacterium adolescentis		3.60	0.35			
07773403	**Good identification**					**07773403**
Actinomyces naeslundii		91.40	0.58	ARA 1		
Actinomyces israelii		8.60	0.31			
07773410	**Acceptable identification**					**07773410**
					SPO	
Actinomyces israelii		94.69	0.48	XYL 92	1	
Actinomyces naeslundii		3.58	0.33	MLZ 13	1	
Clostridium butyricum		1.73	-.03		99	
07773411	**Low discrimination**					**07773411**
Actinomyces naeslundii		84.89	0.36	ARA 1		
Actinomyces israelii		15.11	0.14	MLZ 5		
07773412	**Low discrimination**					**07773412**
Actinomyces israelii		71.38	0.51	XYL 92		
Bifidobacterium adolescentis		23.91	0.37	MLZ 13		
Clostridium clostridioforme		2.16	0.16			
07773413	**Low discrimination**					**07773413**
Actinomyces naeslundii		78.92	0.36	ARA 1		
Actinomyces israelii		21.08	0.17	MLZ 5		
07773420	**Low discrimination**					**07773420**
					SPO	
Actinomyces naeslundii		56.57	0.55	ARA 1	1	
Actinomyces israelii		33.62	0.42		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		7.43	0.15		92	
07773421	**Very good identification**					**07773421**
Actinomyces naeslundii		100.0	0.58	ARA 1		
07773422	**Low discrimination**					**07773422**
					SPO	
Actinomyces naeslundii		39.26	0.55	ARA 1	1	
Actinomyces israelii		35.01	0.45		1	
Clostridium ramosum		20.63	0.25		92	
07773423	**Very good identification**					**07773423**
Actinomyces naeslundii		100.0	0.58	ARA 1		
07773431	**Good identification**					**07773431**
Actinomyces naeslundii		100.0	0.36	ARA 1		
07773433	**Good identification**					**07773433**
Actinomyces naeslundii		100.0	0.36	ARA 1		
07773440	**Acceptable identification**					**07773440**
					SPO	
Actinomyces israelii		94.31	0.55	XYL 92	1	
Actinomyces naeslundii		4.60	0.42		1	
Clostridium butyricum		1.09	0.01		99	
07773441	**Low discrimination**					**07773441**
Actinomyces naeslundii		87.86	0.45	ARA 1		
Actinomyces israelii		12.14	0.21			
07773442	**Very good identification**					**07773442**
Actinomyces israelii		95.17	0.58	XYL 92		
Actinomyces naeslundii		3.09	0.42			
Bifidobacterium adolescentis		1.24	0.20			
07773443	**Low discrimination**					**07773443**
Actinomyces naeslundii		82.83	0.45	ARA 1		
Actinomyces israelii		17.17	0.24			
07773450	**Good identification**					**07773450**
					COC	
Actinomyces israelii		96.11	0.41	XYL 92	1	
Blautia producta		3.89	-.03	MLZ 13	99	
07773452	**Acceptable identification**					**07773452**
Actinomyces israelii		90.38	0.44	XYL 92		
Bifidobacterium adolescentis		9.62	0.21	MLZ 13		
07773460	**Low discrimination**					**07773460**
Actinomyces israelii		56.71	0.35	XYL 92		
Actinomyces naeslundii		43.29	0.42	NAG 6		
07773461	**Good identification**					**07773461**
Actinomyces naeslundii		100.0	0.45	ARA 1		
07773462	**Low discrimination**					**07773462**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		64.75	0.38	XYL 92		
Actinomyces naeslundii		32.95	0.42	NAG 6		
Lactobacillus acidophilus		2.30	-.02			
07773463	**Good identification**					**07773463**
Actinomyces naeslundii		100.0	0.45	ARA 1		
07773500	**Low discrimination**					**07773500**
					SPO	
Actinomyces israelii		71.08	0.73	XYL 92	1	
Clostridium butyricum		22.11	0.43		99	
Actinomyces naeslundii		5.43	0.63		1	
07773501	**Low discrimination**					**07773501**
					SPO	
Actinomyces naeslundii		89.64	0.66	ARA 1	1	
Actinomyces israelii		7.90	0.39		1	
Clostridium butyricum		2.46	0.09		99	
07773502	**Low discrimination**					**07773502**
					SPO	
Bifidobacterium adolescentis		46.51	0.70	XYL 87	1	
Actinomyces israelii		45.89	0.76		1	
Clostridium butyricum		3.70	0.36		99	
07773503	**Low discrimination**					**07773503**
Actinomyces naeslundii		78.32	0.66	ARA 1		
Bifidobacterium adolescentis		10.49	0.35			
Actinomyces israelii		10.35	0.42			
07773510	**Low discrimination**					**07773510**
					SPO	
Actinomyces israelii		76.33	0.59	XYL 92	1	
Bifidobacterium adolescentis		9.59	0.37	MLZ 13	1	
Clostridium butyricum		8.36	0.21		99	
07773511	**Low discrimination**					**07773511**
Actinomyces naeslundii		79.99	0.44	ARA 1		
Actinomyces israelii		20.01	0.24	MLZ 5		
07773512	**Low discrimination**					**07773512**
Bifidobacterium adolescentis		85.90	0.71	XYL 87		
Actinomyces israelii		10.36	0.62			
Bifidobacterium dentium		2.73	0.27			
07773513	**Low discrimination**					**07773513**
Bifidobacterium adolescentis		67.85	0.37	XYL 87		
Actinomyces naeslundii		21.81	0.44	URE 1		
Actinomyces israelii		8.19	0.28			
07773520	**Low discrimination**					**07773520**
					SPO	
Clostridium ramosum		67.38	0.49	ARA 1	92	
Actinomyces naeslundii		14.74	0.63		1	
Actinomyces israelii		12.32	0.52		1	
07773521	**Very good identification**					**07773521**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		96.49	0.66	ARA 1	1		
Clostridium ramosum		2.97	0.15		92		
Actinomyces israelii		0.54	0.18		1		
07773522	**Low discrimination**						**07773522**
					SPO		
Clostridium ramosum		85.12	0.59	ARA 1	92		
Actinomyces israelii		5.84	0.55		1		
Actinomyces naeslundii		4.66	0.63		1		
07773523	**Low discrimination**						**07773523**
					SPO		
Actinomyces naeslundii		88.38	0.66	ARA 1	1		
Clostridium ramosum		10.88	0.25		92		
Actinomyces israelii		0.75	0.21		1		
07773531	**Good identification**						**07773531**
Actinomyces naeslundii		100.0	0.44	ARA 1			
07773533	**Low discrimination**						**07773533**
Actinomyces naeslundii		85.05	0.44	ARA 1			
Bifidobacterium dentium		14.95	-.03	MLZ 5			
07773540	**Low discrimination**						**07773540**
					COC	SPO	
Actinomyces israelii		86.28	0.66	XYL 92	1	1	
Clostridium butyricum		6.00	0.25		1	99	
Blautia producta		4.31	0.24		99	1	
07773541	**Low discrimination**						**07773541**
Actinomyces naeslundii		83.74	0.53	ARA 1			
Actinomyces israelii		16.26	0.32				
07773542	**Low discrimination**						**07773542**
Actinomyces israelii		72.53	0.69	XYL 92			
Bifidobacterium adolescentis		23.36	0.54				
Actinomyces naeslundii		1.68	0.50				
07773543	**Low discrimination**						**07773543**
Actinomyces naeslundii		72.19	0.53	ARA 1			
Actinomyces israelii		21.03	0.35				
Bifidobacterium adolescentis		6.77	0.20				
07773552	**Low discrimination**						**07773552**
					COC		
Bifidobacterium adolescentis		69.97	0.56	XYL 87	1		
Actinomyces israelii		26.56	0.55	SOR 11	1		
Blautia producta		1.97	0.15		99		
07773560	**Low discrimination**						**07773560**
					SPO		
Actinomyces israelii		52.95	0.45	XYL 92	1		
Actinomyces naeslundii		28.76	0.50	NAG 6	1		
Clostridium ramosum		7.52	0.15		92		
07773561	**Good identification**						**07773561**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.53	ARA 1			
07773562	**Low discrimination**						**07773562**
					SPO		
Actinomyces israelii		48.41	0.48	XYL 92	1		
Clostridium ramosum		18.34	0.25	NAG 6	92		
Actinomyces naeslundii		17.53	0.50		1		
07773563	**Good identification**						**07773563**
Actinomyces naeslundii		98.18	0.53	ARA 1			
Actinomyces israelii		1.83	0.14				
07773600	**Good identification**						**07773600**
					SPO		
Actinomyces israelii		97.91	0.43	XYL 92	1		
Clostridium butyricum		2.09	-.07	RHA 7	99		
07773602	**Acceptable identification**						**07773602**
Actinomyces israelii		91.20	0.46	XYL 92			
Clostridium clostridioforme		8.80	0.19	RHA 7			
07773640	**Good identification**						**07773640**
Actinomyces israelii		100.0	0.36	XYL 92			
07773642	**Good identification**						**07773642**
Actinomyces israelii		100.0	0.39	XYL 92			
07773700	**Low discrimination**						**07773700**
					SPO		
Actinomyces israelii		83.32	0.54	XYL 92	1		
Clostridium butyricum		10.65	0.17	RHA 7	99		
Clostridium ramosum		5.18	0.17		92		
07773702	**Low discrimination**						**07773702**
					SPO		
Actinomyces israelii		69.30	0.57	XYL 92	1		
Clostridium ramosum		11.48	0.27	RHA 7	92		
Bifidobacterium adolescentis		9.43	0.35		1		
07773710	**Low discrimination**						**07773710**
					COC	SPO	
Actinomyces israelii		88.14	0.39	XYL 92	1	1	
Blautia producta		4.61	-.03	RHA 7	99	1	
Clostridium butyricum		3.97	-.05	MLZ 13	1	99	
07773712	**Low discrimination**						**07773712**
Clostridium clostridioforme		53.41	0.40	MAN 1			
Bifidobacterium adolescentis		24.13	0.37	XYL 91			
Actinomyces israelii		21.69	0.42				
07773720	**Good identification**						**07773720**
					SPO		
Clostridium ramosum		98.63	0.51	ARA 1	92		
Actinomyces israelii		1.02	0.33		1		
Clostridium butyricum		0.18	-.01		99		
07773722	**Very good identification**						**07773722**
					SPO		
Clostridium ramosum		98.12	0.61	ARA 1	92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		0.55	0.35		1		
Clostridium clostridioforme		0.48	0.28		66		
07773732	**Low discrimination**						**07773732**
					SPO		
Clostridium clostridioforme		57.89	0.40	MAN 1	66		
Clostridium ramosum		26.07	0.27	XYL 91	92		
Parabacteroides distasonis		7.75	0.30		1		
07773740	**Low discrimination**						**07773740**
					COC	SPO	
Actinomyces israelii		59.12	0.47	XYL 92	1	1	
Blautia producta		39.19	0.24	RHA 7	99	1	
Clostridium butyricum		1.69	-.01		1	99	
07773742	**Acceptable identification**						**07773742**
					COC		
Actinomyces israelii		91.55	0.50	XYL 92	1		
Blautia producta		4.50	0.07	RHA 7	99		
Bifidobacterium adolescentis		3.96	0.20		1		
07773752	**Low discrimination**						**07773752**
					COC		
Actinomyces israelii		41.61	0.35	XYL 92	1		
Blautia producta		41.02	0.15	RHA 7	99		
Bifidobacterium adolescentis		14.71	0.21	MLZ 13	1		
07774000	**Low discrimination**						**07774000**
					GRA	SPO	
Fusobacterium mortiferum		46.21	0.36	MLT 11	1	1	
Actinomyces israelii		30.42	0.35	XYL 7	99	1	
Clostridium butyricum		20.56	0.11		99	99	
07774002	**Low discrimination**						**07774002**
					SPO		
Actinomyces israelii		44.53	0.38	GAL 99	1		
Clostridium clostridioforme		37.49	0.27		66		
Clostridium butyricum		7.80	0.04		99		
07774012	**Good identification**						**07774012**
Clostridium clostridioforme		96.25	0.39	RHA 88			
Actinomyces israelii		3.75	0.24	ARA 94			
07774032	**Good identification**						**07774032**
Clostridium clostridioforme		100.0	0.39	RHA 88			
07774101	**Low discrimination**						**07774101**
					SPO		
Actinomyces naeslundii		70.56	0.37	GAL 96	1		
Clostridium butyricum		29.44	0.01	XYL 1	99		
07774103	**Low discrimination**						**07774103**
					SPO		
Actinomyces naeslundii		75.96	0.37	GAL 96	1		
Clostridium butyricum		12.32	-.07	XYL 1	99		
Actinomyces israelii		11.72	0.14		1		
07774112	**Low discrimination**						**07774112**
Clostridium clostridioforme		79.54	0.48	RHA 88			
Bifidobacterium adolescentis		15.13	0.38	ARA 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		3.70	0.34			
07774121	**Good identification**					**07774121**
Actinomyces naeslundii		100.0	0.37	GAL 96		
07774122	**Low discrimination**					**07774122**
					GRA	
Parabacteroides distasonis		51.60	0.47	GAL 99	1	
Clostridium clostridioforme		27.03	0.36		50	
Lactobacillus acidophilus		6.03	0.19		99	
07774123	**Low discrimination**					**07774123**
					GRA	
Actinomyces naeslundii		82.82	0.37	GAL 96	99	
Parabacteroides distasonis		17.18	0.13	XYL 1	1	
07774132	**Low discrimination**					**07774132**
					GRA	
Clostridium clostridioforme		81.40	0.48	RHA 88	50	
Parabacteroides distasonis		18.36	0.42	ARA 94	1	
Actinomyces israelii		0.24	0.14		99	
07774140	**Low discrimination**					**07774140**
					SPO	
Actinomyces israelii		52.44	0.38	GAL 99	1	
Clostridium butyricum		47.56	0.17		99	
07774142	**Low discrimination**					**07774142**
					SPO	
Actinomyces israelii		69.26	0.41	GAL 99	1	
Clostridium butyricum		16.28	0.10		99	
Bifidobacterium adolescentis		10.99	0.21		1	
07774202	**Good identification**					**07774202**
Clostridium clostridioforme		98.80	0.42	ARA 94		
Actinomyces israelii		1.20	0.19			
07774212	**Good identification**					**07774212**
Clostridium clostridioforme		100.0	0.53	ARA 94		
07774222	**Good identification**					**07774222**
Clostridium clostridioforme		97.43	0.42	ARA 94		
Parabacteroides distasonis		2.57	0.21			
07774232	**Good identification**					**07774232**
Clostridium clostridioforme		99.69	0.53	ARA 94		
Parabacteroides distasonis		0.31	0.16			
07774302	**Good identification**					**07774302**
					GRA	
Clostridium clostridioforme		96.99	0.51	ARA 94	50	
Actinomyces israelii		1.41	0.29		99	
Parabacteroides distasonis		0.77	0.21		1	
07774312	**Very good identification**					**07774312**
					GRA	
Clostridium clostridioforme		99.86	0.62	ARA 94	50	
Parabacteroides distasonis		0.09	0.16		1	
Actinomyces israelii		0.05	0.15		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07774322	**Low discrimination**						**07774322**
					SPO		
Clostridium clostridioforme		55.13	0.51	ARA 94	66		
Parabacteroides distasonis		43.04	0.55		1		
Clostridium ramosum		0.94	0.14		92		
07774332	**Good identification**						**07774332**
Clostridium clostridioforme		91.52	0.62	ARA 94			
Parabacteroides distasonis		8.45	0.50				
Bacteroides ovatus		0.04	-.06				
07774402	**Low discrimination**						**07774402**
					SPO		
Clostridium clostridioforme		83.90	0.48	RHA 88	66		
Clostridium butyricum		10.03	0.20		99		
Actinomyces israelii		5.21	0.37		1		
07774412	**Very good identification**						**07774412**
Clostridium clostridioforme		99.39	0.59	RHA 88			
Bifidobacterium adolescentis		0.27	0.18				
Actinomyces israelii		0.20	0.22				
07774422	**Good identification**						**07774422**
					SPO		
Clostridium clostridioforme		98.59	0.48	RHA 88	66		
Clostridium butyricum		1.03	0.02		99		
Actinomyces israelii		0.39	0.16		1		
07774432	**Very good identification**						**07774432**
Clostridium clostridioforme		100.0	0.59	RHA 88			
07774500	**Good identification**						**07774500**
					SPO		
Clostridium butyricum		97.24	0.51	ESL 88	99		
Actinomyces israelii		2.18	0.44	GAL 99	1		
Clostridium clostridioforme		0.45	0.23		66		
07774502	**Low discrimination**						**07774502**
					SPO		
Clostridium clostridioforme		44.76	0.57	RHA 88	66		
Clostridium butyricum		38.37	0.44		99		
Bifidobacterium adolescentis		13.55	0.51		1		
07774512	**Good identification**						**07774512**
					SPO		
Clostridium clostridioforme		91.42	0.68	RHA 88	66		
Bifidobacterium adolescentis		7.43	0.53		1		
Clostridium butyricum		0.91	0.22		99		
07774522	**Low discrimination**						**07774522**
					GRA	SPO	
Clostridium clostridioforme		86.27	0.57	RHA 88	50	66	
Parabacteroides distasonis		6.44	0.43		1	1	
Clostridium butyricum		6.43	0.26		99	99	
07774532	**Very good identification**						**07774532**
					GRA		
Clostridium clostridioforme		98.87	0.68	RHA 88	50		
Parabacteroides distasonis		0.87	0.39		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		0.08	0.18		99	
07774602	**Very good identification**					**07774602**
					SPO	
Clostridium clostridioforme		99.89	0.63		66	
Actinomyces israelii		0.06	0.17		1	
Clostridium butyricum		0.05	-.06		99	
07774610	**Good identification**					**07774610**
Clostridium clostridioforme		100.0	0.40	BGL 99		
07774612	**Very good identification**					**07774612**
Clostridium clostridioforme		100.0	0.74			
07774613	**Good identification**					**07774613**
Clostridium clostridioforme		100.0	0.40	URE 1		
07774622	**Very good identification**					**07774622**
Clostridium clostridioforme		99.90	0.63			
Parabacteroides distasonis		0.10	0.17			
07774630	**Good identification**					**07774630**
Clostridium clostridioforme		100.0	0.40	BGL 99		
07774632	**Very good identification**					**07774632**
Clostridium clostridioforme		99.99	0.74			
Parabacteroides distasonis		0.01	0.13			
07774633	**Good identification**					**07774633**
Clostridium clostridioforme		100.0	0.40	URE 1		
07774652	**Good identification**					**07774652**
Clostridium clostridioforme		100.0	0.40	SOR 1		
07774672	**Good identification**					**07774672**
Clostridium clostridioforme		100.0	0.40	SOR 1		
07774700	**Low discrimination**					**07774700**
					SPO	
Clostridium clostridioforme		50.75	0.37	BGL 99	66	
Clostridium butyricum		46.71	0.25		99	
Actinomyces israelii		2.55	0.25		1	
07774702	**Very good identification**					**07774702**
					SPO	
Clostridium clostridioforme		99.49	0.72		66	
Clostridium butyricum		0.36	0.18		99	
Actinomyces israelii		0.08	0.28		1	
07774703	**Good identification**					**07774703**
Clostridium clostridioforme		100.0	0.37	URE 1		
07774710	**Good identification**					**07774710**
Clostridium clostridioforme		98.95	0.49	BGL 99		
Clostridium butyricum		1.05	0.03			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
07774712	**Very good identification**						**07774712**
Clostridium clostridioforme		99.98	0.83				
Bifidobacterium adolescentis		0.01	0.18				
Actinomyces israelii		0.00	0.14				
07774713	**Good identification**						**07774713**
Clostridium clostridioforme		100.0	0.49	URE 1			
07774720	**Low discrimination**						**07774720**
					GRA	SPO	
Clostridium clostridioforme		85.46	0.37	BGL 99	50	66	
Clostridium butyricum		6.84	0.07		99	99	
Bacteroides vulgatus		5.09	0.04		1	1	
07774722	**Very good identification**						**07774722**
Clostridium clostridioforme		96.45	0.72				
Parabacteroides distasonis		2.95	0.51				
Bacteroides ovatus		0.54	0.23				
07774723	**Good identification**						**07774723**
Clostridium clostridioforme		97.04	0.37	URE 1			
Parabacteroides distasonis		2.96	0.17				
07774730	**Good identification**						**07774730**
Clostridium clostridioforme		99.64	0.49	BGL 99			
Parabacteroides distasonis		0.36	0.13				
07774732	**Very good identification**						**07774732**
Clostridium clostridioforme		99.56	0.83				
Parabacteroides distasonis		0.36	0.47				
Bacteroides ovatus		0.08	0.20				
07774733	**Good identification**						**07774733**
Clostridium clostridioforme		99.64	0.49	URE 1			
Parabacteroides distasonis		0.36	0.13				
07774742	**Acceptable identification**						**07774742**
					SPO		
Clostridium clostridioforme		94.30	0.37	SOR 1	66		
Clostridium butyricum		2.94	0.00		99		
Actinomyces israelii		2.76	0.21		1		
07774752	**Good identification**						**07774752**
Clostridium clostridioforme		100.0	0.49	SOR 1			
07774762	**Acceptable identification**						**07774762**
Clostridium clostridioforme		94.89	0.37	SOR 1			
Parabacteroides distasonis		2.90	0.17				
Bacteroides ovatus		2.21	-.01				
07774772	**Good identification**						**07774772**
Clostridium clostridioforme		99.30	0.49	SOR 1			
Parabacteroides distasonis		0.36	0.13				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		0.34	-.04			
07775000	**Low discrimination**					**07775000**
					SPO	
Actinomyces israelii		56.44	0.69		1	
Clostridium butyricum		38.14	0.45		99	
Lactobacillus jensenii		1.91	0.32		1	
07775001	**Low discrimination**					**07775001**
					SPO	
Actinomyces naeslundii		66.40	0.53	XYL 1	1	
Actinomyces israelii		20.05	0.35		1	
Clostridium butyricum		13.55	0.11		99	
07775002	**Low discrimination**					**07775002**
					SPO	
Actinomyces israelii		76.25	0.72		1	
Clostridium butyricum		13.36	0.38		99	
Lactobacillus acidophilus		4.17	0.35		1	
07775003	**Low discrimination**					**07775003**
					SPO	
Actinomyces naeslundii		64.23	0.53	XYL 1	1	
Actinomyces israelii		29.09	0.38		1	
Clostridium butyricum		5.10	0.04		99	
07775010	**Low discrimination**					**07775010**
					SPO	
Actinomyces israelii		75.01	0.55	MLZ 13	1	
Clostridium butyricum		17.86	0.23		99	
Lactobacillus jensenii		5.67	0.24		1	
07775012	**Low discrimination**					**07775012**
Actinomyces israelii		48.79	0.58	MLZ 13		
Clostridium clostridioforme		37.94	0.47			
Bifidobacterium adolescentis		8.05	0.38			
07775020	**Low discrimination**					**07775020**
					SPO	
Actinomyces israelii		39.26	0.49	NAG 6	1	
Clostridium butyricum		36.15	0.27		99	
Actinomyces naeslundii		13.72	0.50		1	
07775021	**Good identification**					**07775021**
					SPO	
Actinomyces naeslundii		96.43	0.53	XYL 1	1	
Actinomyces israelii		1.86	0.15		1	
Clostridium butyricum		1.71	-.07		99	
07775022	**Low discrimination**					**07775022**
					GRA	
Actinomyces israelii		23.71	0.52	NAG 6	99	
Lactobacillus acidophilus		20.31	0.35		99	
Parabacteroides distasonis		20.18	0.47		1	
07775023	**Acceptable identification**					**07775023**
Actinomyces naeslundii		92.74	0.53	XYL 1		
Actinomyces israelii		2.68	0.18			
Lactobacillus acidophilus		2.30	0.01			
07775032	**Low discrimination**					**07775032**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Clostridium clostridioforme		73.37	0.47	RHA 88	50		
Parabacteroides distasonis		18.47	0.42	ARA 94	1		
Actinomyces israelii		6.02	0.38		99		
07775040	**Low discrimination**						**07775040**
					SPO		
Actinomyces israelii		86.11	0.62		1		
Clostridium butyricum		13.01	0.27		99		
Actinomyces naeslundii		0.87	0.37		1		
07775041	**Low discrimination**						**07775041**
					SPO		
Actinomyces naeslundii		56.63	0.40	XYL 1	1		
Actinomyces israelii		37.68	0.28		1		
Clostridium butyricum		5.69	-.07		99		
07775042	**Good identification**						**07775042**
					SPO		
Actinomyces israelii		93.48	0.65		1		
Clostridium butyricum		3.66	0.20		99		
Lactobacillus acidophilus		1.63	0.20		1		
07775043	**Low discrimination**						**07775043**
Actinomyces naeslundii		50.05	0.40	XYL 1			
Actinomyces israelii		49.95	0.31				
07775050	**Acceptable identification**						**07775050**
					SPO		
Actinomyces israelii		94.95	0.48	MLZ 13	1		
Clostridium butyricum		5.05	0.05		99		
07775052	**Acceptable identification**						**07775052**
Actinomyces israelii		92.04	0.51	MLZ 13			
Bifidobacterium adolescentis		4.83	0.23				
Clostridium clostridioforme		1.86	0.13				
07775060	**Low discrimination**						**07775060**
					SPO		
Actinomyces israelii		73.29	0.42	NAG 6	1		
Clostridium butyricum		15.09	0.09		99		
Actinomyces naeslundii		11.62	0.37		1		
07775061	**Good identification**						**07775061**
Actinomyces naeslundii		100.0	0.40	XYL 1			
07775062	**Low discrimination**						**07775062**
Actinomyces israelii		68.80	0.45	NAG 6			
Lactobacillus acidophilus		18.74	0.20				
Actinomyces naeslundii		7.27	0.37				
07775063	**Good identification**						**07775063**
Actinomyces naeslundii		100.0	0.40	XYL 1			
07775100	**Low discrimination**						**07775100**
					COC	SPO	
Clostridium butyricum		73.82	0.69	ESL 88	1	99	
Actinomyces israelii		18.20	0.80	ARA 90	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		7.23	0.59		99	1	
07775102	**Low discrimination**						**07775102**
					SPO		
Clostridium butyricum		37.41	0.62	ESL 88	99		
Actinomyces israelii		35.58	0.83	ARA 90	1		
Bifidobacterium adolescentis		17.77	0.71		1		
07775103	**Low discrimination**						**07775103**
					SPO		
Actinomyces naeslundii		36.03	0.61	XYL 1	1		
Clostridium butyricum		24.11	0.28		99		
Actinomyces israelii		22.93	0.48		1		
07775110	**Low discrimination**						**07775110**
					SPO		
Clostridium butyricum		56.30	0.47	ESL 88	99		
Actinomyces israelii		39.41	0.65	ARA 90	1		
Bifidobacterium adolescentis		2.44	0.38	MLZ 5	1		
07775112	**Low discrimination**						**07775112**
Bifidobacterium adolescentis		65.32	0.73	ARA 87			
Actinomyces israelii		15.99	0.68				
Clostridium clostridioforme		10.41	0.56				
07775113	**Low discrimination**						**07775113**
Bifidobacterium adolescentis		60.71	0.38	ARA 87			
Actinomyces israelii		14.86	0.34	URE 1			
Clostridium clostridioforme		9.67	0.22				
07775120	**Low discrimination**						**07775120**
					COC	SPO	
Clostridium butyricum		62.40	0.51	ESL 88	1	99	
Atopobium parvulum		12.40	0.46	ARA 90	99	1	
Actinomyces israelii		11.30	0.59	NAG 8	1	1	
07775121	**Low discrimination**						**07775121**
					COC	SPO	
Actinomyces naeslundii		81.59	0.61	XYL 1	1	1	
Clostridium butyricum		12.21	0.17		1	99	
Atopobium parvulum		2.43	0.12		99	1	
07775122	**Low discrimination**						**07775122**
					GRA		
Parabacteroides distasonis		70.48	0.81		1		
Lactobacillus acidophilus		8.24	0.54		99		
Prevotella oralis		7.05	0.51		1		
07775123	**Low discrimination**						**07775123**
					GRA		
Actinomyces naeslundii		45.53	0.61	XYL 1	99		
Parabacteroides distasonis		38.97	0.47		1		
Lactobacillus acidophilus		4.56	0.19		99		
07775131	**Low discrimination**						**07775131**
					SPO		
Actinomyces naeslundii		86.99	0.39	XYL 1	1		
Clostridium butyricum		13.01	-.05	MLZ 5	99		
07775132	**Low discrimination**						**07775132**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		85.00	0.76		1		
Clostridium clostridioforme		11.42	0.56		50		
Actinomyces israelii		1.12	0.48		99		
07775133	**Low discrimination**						**07775133**
					GRA		
Parabacteroides distasonis		79.26	0.42	URE 1	1		
Clostridium clostridioforme		10.65	0.22		50		
Actinomyces naeslundii		9.05	0.39		99		
07775140	**Low discrimination**						**07775140**
					COC	SPO	
Actinomyces israelii		51.87	0.73		1	1	
Clostridium butyricum		47.04	0.51		1	99	
Atopobium parvulum		0.54	0.25		99	1	
07775141	**Low discrimination**						**07775141**
					SPO		
Actinomyces naeslundii		35.93	0.48	XYL 1	1		
Actinomyces israelii		33.60	0.38		1		
Clostridium butyricum		30.47	0.17		99		
07775142	**Low discrimination**						**07775142**
					SPO		
Actinomyces israelii		68.83	0.76		1		
Clostridium butyricum		16.18	0.44		99		
Bifidobacterium adolescentis		10.93	0.56		1		
07775143	**Low discrimination**						**07775143**
					SPO		
Actinomyces israelii		46.37	0.41	URE 1	1		
Actinomyces naeslundii		33.06	0.48		1		
Clostridium butyricum		10.90	0.10		99		
07775150	**Low discrimination**						**07775150**
					SPO		
Actinomyces israelii		74.31	0.58	MLZ 13	1		
Clostridium butyricum		23.74	0.29		99		
Bifidobacterium adolescentis		1.46	0.23		1		
07775152	**Low discrimination**						**07775152**
					SPO		
Bifidobacterium adolescentis		53.67	0.57	ARA 87	1		
Actinomyces israelii		41.34	0.61	SOR 11	1		
Clostridium butyricum		3.42	0.22		99		
07775161	**Low discrimination**						**07775161**
					SPO		
Actinomyces naeslundii		88.23	0.48	XYL 1	1		
Clostridium butyricum		6.51	-.02		99		
Actinomyces israelii		5.27	0.18		1		
07775162	**Low discrimination**						**07775162**
					GRA		
Actinomyces israelii		34.43	0.55	NAG 6	99		
Lactobacillus acidophilus		26.95	0.38		99		
Parabacteroides distasonis		18.84	0.47		1		
07775163	**Low discrimination**						**07775163**
Actinomyces naeslundii		82.74	0.48	XYL 1			
Actinomyces israelii		7.41	0.21				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		5.80	0.04				
07775170	**Low discrimination**						**07775170**
					SPO		
Actinomyces israelii		65.21	0.38	MLZ 13	1		
Clostridium butyricum		28.37	0.11	NAG 6	99		
Bifidobacterium breve		3.83	0.12		1		
07775172	**Low discrimination**						**07775172**
					GRA		
Parabacteroides distasonis		48.93	0.42	SOR 1	1		
Actinomyces israelii		24.81	0.41		99		
Bifidobacterium breve		9.82	0.29		99		
07775200	**Low discrimination**						**07775200**
					SPO		
Actinomyces israelii		76.02	0.50	RHA 7	1		
Clostridium butyricum		21.11	0.19		99		
Clostridium clostridioforme		2.86	0.16		66		
07775202	**Low discrimination**						**07775202**
					SPO		
Clostridium clostridioforme		68.20	0.50	ARA 94	66		
Actinomyces israelii		27.44	0.53		1		
Clostridium butyricum		1.98	0.12		99		
07775212	**Very good identification**						**07775212**
					GRA		
Clostridium clostridioforme		98.60	0.62	ARA 94	50		
Actinomyces israelii		1.30	0.39		99		
Parabacteroides distasonis		0.10	0.16		1		
07775222	**Low discrimination**						**07775222**
Clostridium clostridioforme		49.45	0.50	ARA 94			
Parabacteroides distasonis		43.09	0.55				
Prevotella oralis		3.89	0.24				
07775232	**Good identification**						**07775232**
					GRA		
Clostridium clostridioforme		90.56	0.62	ARA 94	50		
Parabacteroides distasonis		9.33	0.50		1		
Actinomyces israelii		0.08	0.18		99		
07775240	**Acceptable identification**						**07775240**
					SPO		
Actinomyces israelii		94.15	0.43	RHA 7	1		
Clostridium butyricum		5.85	0.01		99		
07775242	**Acceptable identification**						**07775242**
Actinomyces israelii		91.07	0.46	RHA 7			
Clostridium clostridioforme		5.88	0.16				
Lactobacillus acidophilus		1.58	0.01				
07775300	**Low discrimination**						**07775300**
					COC	SPO	
Clostridium butyricum		60.09	0.43	ESL 88	1	99	
Actinomyces israelii		36.06	0.60	RHA 3	1	1	
Atopobium parvulum		1.92	0.25	ARA 90	99	1	
07775302	**Low discrimination**						**07775302**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		47.94	0.59	ARA 94	50		
Actinomyces israelii		23.04	0.63		99		
Parabacteroides distasonis		12.48	0.55		1		
07775310	**Low discrimination**						**07775310**
					SPO		
Actinomyces israelii		38.80	0.46	RHA 7	1		
Clostridium clostridioforme		37.28	0.36	MLZ 13	66		
Clostridium butyricum		22.77	0.21		99		
07775312	**Good identification**						**07775312**
					GRA		
Clostridium clostridioforme		94.52	0.71	ARA 94	50		
Parabacteroides distasonis		2.91	0.50		1		
Actinomyces israelii		1.49	0.49		99		
07775313	**Good identification**						**07775313**
					GRA		
Clostridium clostridioforme		95.55	0.36	ARA 94	50		
Parabacteroides distasonis		2.94	0.16	URE 1	1		
Actinomyces israelii		1.51	0.15		99		
07775320	**Low discrimination**						**07775320**
					SPO	GRA	
Parabacteroides distasonis		49.51	0.55	BGL 99	1	1	
Clostridium ramosum		26.68	0.38		92	50	
Clostridium butyricum		8.84	0.25		99	99	
07775322	**Good identification**						**07775322**
					SPO		
Parabacteroides distasonis		92.31	0.89		1		
Clostridium clostridioforme		3.58	0.59		66		
Clostridium ramosum		2.01	0.48		92		
07775323	**Acceptable identification**						**07775323**
					SPO		
Parabacteroides distasonis		92.15	0.55	URE 1	1		
Clostridium clostridioforme		3.58	0.25		66		
Clostridium ramosum		2.01	0.14		92		
07775330	**Low discrimination**						**07775330**
					GRA		
Parabacteroides distasonis		72.36	0.50	BGL 99	1		
Clostridium clostridioforme		23.76	0.36		50		
Actinomyces israelii		1.58	0.26		99		
07775332	**Low discrimination**						**07775332**
Parabacteroides distasonis		74.97	0.85				
Clostridium clostridioforme		24.62	0.71				
Bacteroides ovatus		0.33	0.28				
07775333	**Low discrimination**						**07775333**
Parabacteroides distasonis		75.04	0.50	URE 1			
Clostridium clostridioforme		24.64	0.36				
Bacteroides ovatus		0.33	-.06				
07775340	**Low discrimination**						**07775340**
					SPO		
Actinomyces israelii		72.86	0.53	RHA 7	1		
Clostridium butyricum		27.15	0.25		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
07775342	**Low discrimination**						**07775342**
					SPO		
Actinomyces israelii		80.91	0.56	RHA 7	1		
Clostridium butyricum		7.82	0.18		99		
Clostridium clostridioforme		4.37	0.25		66		
07775350	**Low discrimination**						**07775350**
					COC	SPO	
Actinomyces israelii		84.36	0.39	RHA 7	1	1	
Clostridium butyricum		11.07	0.03	MLZ 13	1	99	
Blautia producta		4.57	-.03		99	1	
07775352	**Low discrimination**						**07775352**
Clostridium clostridioforme		56.68	0.36	ARA 94			
Actinomyces israelii		34.41	0.42	SOR 1			
Bifidobacterium adolescentis		6.00	0.23				
07775362	**Low discrimination**						**07775362**
					GRA		
Parabacteroides distasonis		84.95	0.55	SOR 1	1		
Actinomyces israelii		3.90	0.36		99		
Clostridium clostridioforme		3.30	0.25		50		
07775372	**Low discrimination**						**07775372**
Parabacteroides distasonis		73.59	0.50	SOR 1			
Clostridium clostridioforme		24.16	0.36				
Bacteroides ovatus		1.32	0.05				
07775400	**Low discrimination**						**07775400**
					SPO		
Clostridium butyricum		88.04	0.62	ESL 88	99		
Actinomyces israelii		11.84	0.68	LAC 96	1		
Clostridium clostridioforme		0.09	0.22		66		
07775402	**Low discrimination**						**07775402**
					SPO		
Clostridium butyricum		53.82	0.55	ESL 88	99		
Actinomyces israelii		27.92	0.71	LAC 96	1		
Clostridium clostridioforme		13.64	0.56		66		
07775410	**Low discrimination**						**07775410**
					SPO		
Clostridium butyricum		67.76	0.40	ESL 88	99		
Actinomyces israelii		25.87	0.54	LAC 96	1		
Clostridium clostridioforme		5.84	0.33	MLZ 5	66		
07775412	**Low discrimination**						**07775412**
Clostridium clostridioforme		83.07	0.67	RHA 88			
Bifidobacterium adolescentis		7.53	0.53				
Actinomyces israelii		5.58	0.57				
07775420	**Acceptable identification**						**07775420**
					SPO		
Clostridium butyricum		90.08	0.43	ESL 88	99		
Actinomyces israelii		8.89	0.47	LAC 96	1		
Clostridium clostridioforme		1.03	0.22	NAG 8	66		
07775422	**Low discrimination**						**07775422**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		63.92	0.56	RHA 88	66		
Clostridium butyricum		21.94	0.36		99		
Actinomyces israelii		8.35	0.50		1		
07775432	**Very good identification**						**07775432**
					GRA		
Clostridium clostridioforme		98.04	0.67	RHA 88	50		
Parabacteroides distasonis		0.97	0.39		1		
Actinomyces israelii		0.42	0.36		99		
07775440	**Low discrimination**						**07775440**
					SPO		
Clostridium butyricum		62.44	0.43	ESL 88	99		
Actinomyces israelii		37.56	0.61	LAC 96	1		
07775442	**Low discrimination**						**07775442**
					SPO		
Actinomyces israelii		66.84	0.64		1		
Clostridium butyricum		28.81	0.36		99		
Bifidobacterium adolescentis		3.51	0.36		1		
07775450	**Low discrimination**						**07775450**
					COC	SPO	
Actinomyces israelii		62.29	0.47	MLZ 13	1	1	
Clostridium butyricum		36.47	0.22		1	99	
Blautia producta		1.24	-.03		99	1	
07775452	**Low discrimination**						**07775452**
Actinomyces israelii		50.82	0.50	MLZ 13			
Bifidobacterium adolescentis		21.81	0.37				
Clostridium clostridioforme		19.66	0.33				
07775462	**Low discrimination**						**07775462**
					SPO		
Actinomyces israelii		55.99	0.43	NAG 6	1		
Clostridium butyricum		32.88	0.18		99		
Clostridium clostridioforme		11.13	0.22		66		
07775500	**Very good identification**						**07775500**
					SPO		
Clostridium butyricum		97.65	0.85	ESL 88	99		
Actinomyces israelii		2.19	0.78		1		
Bifidobacterium adolescentis		0.14	0.51		1		
07775501	**Good identification**						**07775501**
					SPO		
Clostridium butyricum		97.55	0.51	ESL 88	99		
Actinomyces israelii		2.19	0.44	URE 1	1		
Bifidobacterium adolescentis		0.14	0.17		1		
07775502	**Low discrimination**						**07775502**
					SPO		
Clostridium butyricum		67.74	0.78	ESL 88	99		
Bifidobacterium adolescentis		23.93	0.85		1		
Actinomyces israelii		5.86	0.81		1		
07775503	**Low discrimination**						**07775503**
					SPO		
Clostridium butyricum		67.72	0.44	ESL 88	99		
Bifidobacterium adolescentis		23.92	0.51	URE 1	1		
Actinomyces israelii		5.86	0.47		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07775510	**Good identification**						**07775510**
					SPO		
Clostridium butyricum		90.11	0.63	ESL 88	99		
Actinomyces israelii		5.74	0.64	MLZ 5	1		
Bifidobacterium adolescentis		2.90	0.53		1		
07775512	**Low discrimination**						**07775512**
					SPO		
Bifidobacterium adolescentis		65.36	0.87		1		
Clostridium clostridioforme		24.38	0.76		66		
Clostridium butyricum		7.97	0.56		99		
07775513	**Low discrimination**						**07775513**
					SPO		
Bifidobacterium adolescentis		65.41	0.53	URE 1	1		
Clostridium clostridioforme		24.40	0.42		66		
Clostridium butyricum		7.98	0.22		99		
07775520	**Very good identification**						**07775520**
					GRA	SPO	
Clostridium butyricum		97.51	0.67	ESL 88	99	99	
Actinomyces israelii		1.61	0.58	NAG 8	99	1	
Parabacteroides distasonis		0.38	0.43		1	1	
07775522	**Low discrimination**						**07775522**
					GRA	SPO	
Parabacteroides distasonis		39.28	0.77		1	1	
Clostridium butyricum		39.22	0.60		99	99	
Clostridium clostridioforme		15.94	0.65		50	66	
07775523	**Low discrimination**						**07775523**
					GRA	SPO	
Parabacteroides distasonis		39.11	0.43	URE 1	1	1	
Clostridium butyricum		39.05	0.26		99	99	
Clostridium clostridioforme		15.87	0.31		50	66	
07775530	**Low discrimination**						**07775530**
					SPO		
Clostridium butyricum		79.15	0.45	ESL 88	99		
Clostridium clostridioforme		10.94	0.42	MLZ 5	66		
Actinomyces israelii		3.70	0.43	NAG 8	1		
07775532	**Low discrimination**						**07775532**
					GRA	SPO	
Clostridium clostridioforme		72.37	0.76	RHA 88	50	66	
Parabacteroides distasonis		21.08	0.73		1	1	
Clostridium butyricum		2.06	0.38		99	99	
07775533	**Low discrimination**						**07775533**
					GRA	SPO	
Clostridium clostridioforme		72.69	0.42	RHA 88	50	66	
Parabacteroides distasonis		21.17	0.39	URE 1	1	1	
Clostridium butyricum		2.07	0.04		99	99	
07775540	**Good identification**						**07775540**
					COC	SPO	
Clostridium butyricum		90.47	0.67	ESL 88	1	99	
Actinomyces israelii		9.07	0.71	SOR 8	1	1	
Blautia producta		0.22	0.24		99	1	
07775542	**Low discrimination**						**07775542**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		52.80	0.60	ESL 88	99		
Bifidobacterium adolescentis		26.51	0.70	SOR 8	1		
Actinomyces israelii		20.42	0.74		1		
07775550	**Low discrimination**						**07775550**
					COC	SPO	
Clostridium butyricum		67.73	0.45	ESL 88	1	99	
Actinomyces israelii		19.28	0.57	MLZ 5	1	1	
Blautia producta		9.52	0.32	SOR 8	99	1	
07775552	**Low discrimination**						**07775552**
					SPO		
Bifidobacterium adolescentis		82.36	0.71	SOR 11	1		
Actinomyces israelii		7.76	0.60		1		
Clostridium butyricum		7.06	0.38		99		
07775553	**Low discrimination**						**07775553**
					SPO		
Bifidobacterium adolescentis		82.62	0.37	SOR 11	1		
Actinomyces israelii		7.78	0.26	URE 1	1		
Clostridium butyricum		7.09	0.04		99		
07775560	**Acceptable identification**						**07775560**
					COC	SPO	
Clostridium butyricum		92.71	0.49	ESL 88	1	99	
Actinomyces israelii		6.82	0.51	NAG 8	1	1	
Blautia producta		0.29	0.07	SOR 8	99	1	
07775562	**Low discrimination**						**07775562**
					GRA	SPO	
Clostridium butyricum		65.68	0.42	ESL 88	99	99	
Actinomyces israelii		18.64	0.54	NAG 8	99	1	
Parabacteroides distasonis		7.64	0.43	SOR 8	1	1	
07775572	**Low discrimination**						**07775572**
					GRA		
Clostridium clostridioforme		46.75	0.42	RHA 88	50		
Bifidobacterium adolescentis		15.49	0.37	SOR 1	99		
Parabacteroides distasonis		13.62	0.39		1		
07775600	**Low discrimination**						**07775600**
					SPO		
Clostridium butyricum		63.96	0.36	ESL 88	99		
Actinomyces israelii		20.94	0.49	LAC 96	1		
Clostridium clostridioforme		15.10	0.37	RHA 3	66		
07775602	**Very good identification**						**07775602**
					SPO		
Clostridium clostridioforme		96.30	0.71		66		
Actinomyces israelii		2.02	0.52		1		
Clostridium butyricum		1.60	0.29		99		
07775603	**Good identification**						**07775603**
					SPO		
Clostridium clostridioforme		96.37	0.37	URE 1	66		
Actinomyces israelii		2.03	0.17		1		
Clostridium butyricum		1.60	-.06		99		
07775610	**Acceptable identification**						**07775610**
					SPO		
Clostridium clostridioforme		91.37	0.48	BGL 99	66		
Clostridium butyricum		4.47	0.14		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		4.16	0.34		1		
07775612	**Very good identification**						**07775612**
					SPO		
Clostridium clostridioforme		99.90	0.82		66		
Actinomyces israelii		0.07	0.37		1		
Clostridium butyricum		0.02	0.07		99		
07775613	**Good identification**						**07775613**
Clostridium clostridioforme		100.0	0.48	URE 1			
07775620	**Low discrimination**						**07775620**
					SPO		
Clostridium clostridioforme		64.21	0.37	BGL 99	66		
Clostridium butyricum		23.66	0.18		99		
Actinomyces israelii		5.68	0.28		1		
07775622	**Very good identification**						**07775622**
Clostridium clostridioforme		95.80	0.71				
Parabacteroides distasonis		3.27	0.51				
Bacteroides ovatus		0.60	0.23				
07775623	**Good identification**						**07775623**
Clostridium clostridioforme		96.70	0.37	URE 1			
Parabacteroides distasonis		3.30	0.17				
07775630	**Good identification**						**07775630**
					GRA	SPO	
Clostridium clostridioforme		98.89	0.48	BGL 99	50	66	
Clostridium butyricum		0.42	-.04		99	99	
Parabacteroides distasonis		0.40	0.13		1	1	
07775632	**Very good identification**						**07775632**
Clostridium clostridioforme		99.50	0.82				
Parabacteroides distasonis		0.40	0.47				
Bacteroides ovatus		0.09	0.20				
07775633	**Good identification**						**07775633**
Clostridium clostridioforme		99.60	0.48	URE 1			
Parabacteroides distasonis		0.40	0.13				
07775640	**Low discrimination**						**07775640**
					SPO		
Actinomyces israelii		59.41	0.41	RHA 7	1		
Clostridium butyricum		40.59	0.18		99		
07775642	**Low discrimination**						**07775642**
					SPO		
Clostridium clostridioforme		51.22	0.37	SOR 1	66		
Actinomyces israelii		41.44	0.45		1		
Clostridium butyricum		7.34	0.11		99		
07775652	**Good identification**						**07775652**
Clostridium clostridioforme		97.42	0.48	SOR 1			
Actinomyces israelii		2.59	0.30				
07775662	**Low discrimination**						**07775662**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		88.96	0.37	SOR 1	50		
Actinomyces israelii		4.59	0.24		99		
Parabacteroides distasonis		3.03	0.17		1		
07775672	**Good identification**						**07775672**
Clostridium clostridioforme		99.22	0.48	SOR 1			
Parabacteroides distasonis		0.40	0.13				
Bacteroides ovatus		0.38	-.04				
07775700	**Good identification**						**07775700**
					SPO		
Clostridium butyricum		91.55	0.59	ESL 88	99		
Actinomyces israelii		5.00	0.59	RHA 3	1		
Clostridium clostridioforme		3.01	0.46		66		
07775702	**Low discrimination**						**07775702**
					SPO		
Clostridium clostridioforme		85.43	0.80		66		
Clostridium butyricum		10.19	0.52		99		
Actinomyces israelii		2.15	0.62		1		
07775703	**Low discrimination**						**07775703**
					SPO		
Clostridium clostridioforme		85.59	0.46	URE 1	66		
Clostridium butyricum		10.21	0.18		99		
Actinomyces israelii		2.15	0.28		1		
07775710	**Low discrimination**						**07775710**
					SPO		
Clostridium clostridioforme		70.47	0.57	BGL 99	66		
Clostridium butyricum		24.73	0.38		99		
Actinomyces israelii		3.83	0.45		1		
07775712	**Very good identification**						**07775712**
					SPO		
Clostridium clostridioforme		99.27	0.91		66		
Bifidobacterium adolescentis		0.37	0.53		1		
Clostridium butyricum		0.14	0.30		99		
07775713	**Very good identification**						**07775713**
					SPO		
Clostridium clostridioforme		99.30	0.57	URE 1	66		
Bifidobacterium adolescentis		0.37	0.18		1		
Clostridium butyricum		0.14	-.04		99		
07775720	**Low discrimination**						**07775720**
					GRA	SPO	
Clostridium butyricum		38.01	0.41	ESL 88	99	99	
Bacteroides vulgatus		28.26	0.38	RHA 3	1	1	
Parabacteroides distasonis		14.51	0.51	NAG 8	1	1	
07775722	**Low discrimination**						**07775722**
Parabacteroides distasonis		45.47	0.86				
Clostridium clostridioforme		45.11	0.80				
Bacteroides ovatus		8.40	0.57				
07775723	**Low discrimination**						**07775723**
Parabacteroides distasonis		45.52	0.51	URE 1			
Clostridium clostridioforme		45.16	0.46				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		8.41	0.23				
07775730	**Low discrimination**						**07775730**
					GRA	SPO	
Clostridium clostridioforme		84.22	0.57	BGL 99	50	66	
Parabacteroides distasonis		10.03	0.47		1	1	
Clostridium butyricum		2.57	0.19		99	99	
07775732	**Low discrimination**						**07775732**
Clostridium clostridioforme		87.15	0.91				
Parabacteroides distasonis		10.38	0.81				
Bacteroides ovatus		2.38	0.54				
07775733	**Low discrimination**						**07775733**
Clostridium clostridioforme		87.17	0.57	URE 1			
Parabacteroides distasonis		10.39	0.47				
Bacteroides ovatus		2.38	0.20				
07775740	**Low discrimination**						**07775740**
					COC	SPO	
Clostridium butyricum		73.21	0.41	ESL 88	1	99	
Actinomyces israelii		17.86	0.52	RHA 3	1	1	
Blautia producta		5.84	0.24	SOR 8	99	1	
07775752	**Good identification**						**07775752**
Clostridium clostridioforme		90.31	0.57	SOR 1			
Bifidobacterium adolescentis		4.08	0.37				
Actinomyces israelii		2.86	0.41				
07775762	**Low discrimination**						**07775762**
Parabacteroides distasonis		34.32	0.51	SOR 1			
Clostridium clostridioforme		34.05	0.46				
Bacteroides ovatus		26.16	0.33				
07775772	**Low discrimination**						**07775772**
Clostridium clostridioforme		80.83	0.57	SOR 1			
Parabacteroides distasonis		9.63	0.47				
Bacteroides ovatus		9.09	0.30				
07776000	**Good identification**						**07776000**
					SPO		
Actinomyces israelii		95.72	0.48	GAL 99	1		
Actinomyces naeslundii		3.02	0.31		1		
Clostridium butyricum		1.27	-.05		99		
07776002	**Good identification**						**07776002**
Actinomyces israelii		97.94	0.51	GAL 99			
Actinomyces naeslundii		2.06	0.31				
07776012	**Good identification**						**07776012**
Actinomyces israelii		95.63	0.37	GAL 99			
Clostridium clostridioforme		4.38	0.04	MLZ 13			
07776040	**Good identification**						**07776040**
Actinomyces israelii		100.0	0.41	GAL 99			

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
07776042	**Good identification**					**07776042**
Actinomyces israelii		100.0	0.44	GAL 99		
07776100	**Good identification**					**07776100**
					SPO	
Actinomyces israelii		90.42	0.58	GAL 99	1	
Clostridium butyricum		7.19	0.18		99	
Actinomyces naeslundii		2.03	0.39		1	
07776101	**Low discrimination**					**07776101**
Actinomyces naeslundii		76.91	0.42	GAL 96		
Actinomyces israelii		23.10	0.24	XYL 1		
07776102	**Low discrimination**					**07776102**
					SPO	
Actinomyces israelii		89.63	0.61	GAL 99	1	
Bifidobacterium adolescentis		6.46	0.35		1	
Clostridium butyricum		1.85	0.11		99	
07776103	**Low discrimination**					**07776103**
Actinomyces naeslundii		68.94	0.42	GAL 96		
Actinomyces israelii		31.06	0.27	XYL 1		
07776110	**Acceptable identification**					**07776110**
					SPO	
Actinomyces israelii		92.93	0.44	GAL 99	1	
Clostridium tertium		4.47	0.07	MLZ 13	99	
Clostridium butyricum		2.60	-.04		99	
07776112	**Low discrimination**					**07776112**
					SPO	
Actinomyces israelii		57.37	0.47	GAL 99	1	
Bifidobacterium adolescentis		33.84	0.37	MLZ 13	1	
Clostridium tertium		5.52	0.15		99	
07776120	**Low discrimination**					**07776120**
					SPO	
Actinomyces israelii		49.05	0.38	GAL 99	1	
Clostridium ramosum		19.09	0.15	NAG 6	92	
Actinomyces naeslundii		17.23	0.39		1	
07776121	**Good identification**					**07776121**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07776123	**Good identification**					**07776123**
Actinomyces naeslundii		100.0	0.42	GAL 96		
07776140	**Good identification**					**07776140**
					SPO	
Actinomyces israelii		97.28	0.51	GAL 99	1	
Clostridium butyricum		1.73	0.00		99	
Actinomyces naeslundii		0.99	0.26		1	
07776142	**Good identification**					**07776142**
Actinomyces israelii		96.68	0.54	GAL 99		
Bifidobacterium adolescentis		2.22	0.20			
Actinomyces naeslundii		0.66	0.26			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
07776150	**Acceptable identification**					**07776150**
					COC	
Actinomyces israelii		93.35	0.37	GAL 99	1	
Blautia producta		6.65	-.03	MLZ 13	99	
07776152	**Low discrimination**					**07776152**
Actinomyces israelii		84.21	0.40	GAL 99		
Bifidobacterium adolescentis		15.79	0.21	MLZ 13		
07776300	**Good identification**					**07776300**
					SPO	
Actinomyces israelii		96.84	0.39	GAL 99	1	
Clostridium butyricum		3.16	-.08	RHA 7	99	
07776302	**Low discrimination**					**07776302**
Actinomyces israelii		89.10	0.42	GAL 99		
Clostridium clostridioforme		10.90	0.17	RHA 7		
07776342	**Good identification**					**07776342**
Actinomyces israelii		100.0	0.35	GAL 99		
07776400	**Low discrimination**					**07776400**
					SPO	
Actinomyces israelii		87.28	0.46	GAL 99	1	
Clostridium butyricum		12.72	0.11		99	
07776402	**Acceptable identification**					**07776402**
					SPO	
Actinomyces israelii		91.71	0.49	GAL 99	1	
Clostridium butyricum		3.47	0.04		99	
Clostridium clostridioforme		2.64	0.14		66	
07776412	**Low discrimination**					**07776412**
Actinomyces israelii		46.24	0.35	GAL 99		
Clostridium clostridioforme		40.51	0.25	MLZ 13		
Bifidobacterium adolescentis		9.01	0.17			
07776440	**Good identification**					**07776440**
					SPO	
Actinomyces israelii		96.84	0.39	GAL 99	1	
Clostridium butyricum		3.16	-.07		99	
07776442	**Good identification**					**07776442**
Actinomyces israelii		100.0	0.42	GAL 99		
07776500	**Low discrimination**					**07776500**
					SPO	
Actinomyces israelii		53.09	0.57	GAL 99	1	
Clostridium butyricum		46.44	0.35		99	
Bifidobacterium adolescentis		0.47	0.15		1	
07776502	**Low discrimination**					**07776502**
					SPO	
Actinomyces israelii		54.23	0.60	GAL 99	1	
Bifidobacterium adolescentis		31.99	0.50		1	
Clostridium butyricum		12.30	0.28		99	
07776510	**Low discrimination**					**07776510**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		57.09	0.43	GAL 99	1		
Bifidobacterium dentium		19.21	0.10	MLZ 13	1		
Clostridium butyricum		17.59	0.13		99		
07776512	**Low discrimination**						**07776512**
Bifidobacterium adolescentis		54.17	0.51	GAL 99			
Bifidobacterium dentium		25.47	0.27				
Actinomyces israelii		11.23	0.46				
07776522	**Low discrimination**						**07776522**
					SPO		
Actinomyces israelii		53.35	0.39	GAL 99	1		
Clostridium clostridioforme		20.10	0.23	NAG 6	66		
Clostridium butyricum		16.48	0.10		99		
07776540	**Low discrimination**						**07776540**
					COC	SPO	
Actinomyces israelii		61.53	0.50	GAL 99	1	1	
Blautia producta		26.43	0.24		99	1	
Clostridium butyricum		12.04	0.17		1	99	
07776542	**Low discrimination**						**07776542**
					SPO		
Actinomyces israelii		78.74	0.53	GAL 99	1		
Bifidobacterium adolescentis		14.76	0.34		1		
Clostridium butyricum		3.99	0.10		99		
07776552	**Low discrimination**						**07776552**
					COC		
Bifidobacterium adolescentis		47.46	0.36	GAL 99	1		
Actinomyces israelii		30.95	0.39	SOR 11	1		
Blautia producta		19.77	0.15		99		
07776612	**Good identification**						**07776612**
Clostridium clostridioforme		98.84	0.40	MAN 1			
Actinomyces israelii		1.16	0.16				
07776632	**Good identification**						**07776632**
Clostridium clostridioforme		100.0	0.40	MAN 1			
07776700	**Low discrimination**						**07776700**
					SPO		
Actinomyces israelii		73.56	0.38	GAL 99	1		
Clostridium butyricum		26.44	0.09	RHA 7	99		
07776702	**Low discrimination**						**07776702**
					SPO		
Clostridium clostridioforme		66.65	0.37	MAN 1	66		
Actinomyces israelii		28.45	0.41		1		
Clostridium butyricum		2.65	0.02		99		
07776712	**Good identification**						**07776712**
Clostridium clostridioforme		97.34	0.49	MAN 1			
Actinomyces israelii		1.36	0.26				
Bifidobacterium adolescentis		0.88	0.17				
07776722	**Low discrimination**						**07776722**
Clostridium clostridioforme		80.38	0.37	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		14.97	0.15			
Parabacteroides distasonis		2.46	0.17			
07776732	**Good identification**					**07776732**
					GRA	
Clostridium clostridioforme		96.30	0.49	MAN 1	50	
Bacteroides ovatus		2.63	0.12		1	
Bifidobacterium dentium		0.73	-.03		99	
07777000	**Very good identification**					**07777000**
					SPO	
Actinomyces israelii		97.92	0.82		1	
Clostridium butyricum		1.30	0.29		99	
Actinomyces naeslundii		0.75	0.55		1	
07777001	**Low discrimination**					**07777001**
					SPO	
Actinomyces naeslundii		52.83	0.58	XYL 1	1	
Actinomyces israelii		46.56	0.48		1	
Clostridium butyricum		0.62	-.05		99	
07777002	**Very good identification**					**07777002**
					SPO	
Actinomyces israelii		98.78	0.85		1	
Actinomyces naeslundii		0.50	0.55		1	
Clostridium butyricum		0.34	0.22		99	
07777003	**Low discrimination**					**07777003**
Actinomyces israelii		56.94	0.51	URE 1		
Actinomyces naeslundii		43.07	0.58			
07777010	**Very good identification**					**07777010**
					SPO	
Actinomyces israelii		99.08	0.68	MLZ 13	1	
Clostridium butyricum		0.46	0.07		99	
Actinomyces naeslundii		0.27	0.33		1	
07777012	**Very good identification**					**07777012**
					SPO	
Actinomyces israelii		96.82	0.71	MLZ 13	1	
Bifidobacterium adolescentis		2.31	0.37		1	
Clostridium tertium		0.38	0.15		99	
07777013	**Low discrimination**					**07777013**
Actinomyces israelii		78.96	0.37	MLZ 13		
Actinomyces naeslundii		21.04	0.36	URE 1		
07777020	**Low discrimination**					**07777020**
					SPO	
Actinomyces israelii		86.12	0.62	NAG 6	1	
Actinomyces naeslundii		10.31	0.55		1	
Clostridium butyricum		1.56	0.11		99	
07777021	**Good identification**					**07777021**
Actinomyces naeslundii		94.67	0.58	XYL 1		
Actinomyces israelii		5.33	0.28			
07777022	**Low discrimination**					**07777022**
					SPO	
Actinomyces israelii		86.86	0.65	NAG 6	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		6.93	0.55		1	
Clostridium ramosum		3.64	0.25		92	
07777023	**Good identification**					**07777023**
Actinomyces naeslundii		92.22	0.58	XYL 1		
Actinomyces israelii		7.78	0.31			
07777030	**Low discrimination**					**07777030**
					SPO	
Actinomyces israelii		88.22	0.48	MLZ 13	1	
Clostridium tertium		8.06	0.15	NAG 6	99	
Actinomyces naeslundii		3.72	0.33		1	
07777031	**Low discrimination**					**07777031**
Actinomyces naeslundii		86.23	0.36	XYL 1		
Actinomyces israelii		13.77	0.13	MLZ 5		
07777032	**Low discrimination**					**07777032**
					SPO	
Actinomyces israelii		79.77	0.51	MLZ 13	1	
Clostridium tertium		14.58	0.24	NAG 6	99	
Actinomyces naeslundii		2.24	0.33		1	
07777033	**Low discrimination**					**07777033**
Actinomyces naeslundii		80.67	0.36	XYL 1		
Actinomyces israelii		19.33	0.16	MLZ 5		
07777040	**Very good identification**					**07777040**
					SPO	
Actinomyces israelii		99.36	0.75		1	
Actinomyces naeslundii		0.34	0.42		1	
Clostridium butyricum		0.29	0.11		99	
07777041	**Low discrimination**					**07777041**
Actinomyces israelii		66.01	0.41	URE 1		
Actinomyces naeslundii		33.99	0.45			
07777042	**Very good identification**					**07777042**
Actinomyces israelii		99.59	0.78			
Actinomyces naeslundii		0.23	0.42			
Bifidobacterium adolescentis		0.09	0.20			
07777043	**Low discrimination**					**07777043**
Actinomyces israelii		74.45	0.44	URE 1		
Actinomyces naeslundii		25.55	0.45			
07777050	**Very good identification**					**07777050**
					COC	
Actinomyces israelii		99.71	0.61	MLZ 13	1	
Blautia producta		0.29	-.03		99	
07777052	**Very good identification**					**07777052**
Actinomyces israelii		99.25	0.64	MLZ 13		
Bifidobacterium adolescentis		0.75	0.21			
07777060	**Acceptable identification**					**07777060**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		94.49	0.55	NAG 6	1	
Actinomyces naeslundii		5.13	0.42		1	
Clostridium butyricum		0.38	-.07		99	
07777061	**Low discrimination**					**07777061**
Actinomyces naeslundii		88.97	0.45	XYL 1		
Actinomyces israelii		11.03	0.20			
07777062	**Very good identification**					**07777062**
Actinomyces israelii		96.27	0.58	NAG 6		
Actinomyces naeslundii		3.49	0.42			
Lactobacillus acidophilus		0.24	-.02			
07777063	**Low discrimination**					**07777063**
Actinomyces naeslundii		84.32	0.45	XYL 1		
Actinomyces israelii		15.68	0.24			
07777070	**Good identification**					**07777070**
Actinomyces israelii		100.0	0.41	MLZ 13		
07777072	**Good identification**					**07777072**
Actinomyces israelii		100.0	0.44	MLZ 13		
07777100	**Good identification**					**07777100**
					SPO	
Actinomyces israelii		91.65	0.92		1	
Clostridium butyricum		7.29	0.53		99	
Actinomyces naeslundii		0.50	0.63		1	
07777101	**Low discrimination**					**07777101**
					SPO	
Actinomyces israelii		52.89	0.58	URE 1	1	
Actinomyces naeslundii		42.70	0.66		1	
Clostridium butyricum		4.21	0.18		99	
07777102	**Good identification**					**07777102**
					SPO	
Actinomyces israelii		90.34	0.95		1	
Bifidobacterium adolescentis		6.52	0.70		1	
Clostridium butyricum		1.86	0.46		99	
07777103	**Low discrimination**					**07777103**
Actinomyces israelii		61.01	0.61	URE 1		
Actinomyces naeslundii		32.84	0.66			
Bifidobacterium adolescentis		4.40	0.35			
07777110	**Good identification**					**07777110**
					SPO	
Actinomyces israelii		91.74	0.78	MLZ 13	1	
Clostridium tertium		4.41	0.41		99	
Clostridium butyricum		2.57	0.31		99	
07777111	**Low discrimination**					**07777111**
					SPO	
Actinomyces israelii		73.50	0.44	MLZ 13	1	
Actinomyces naeslundii		20.90	0.44	URE 1	1	
Clostridium tertium		3.54	0.07		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
07777112	**Low discrimination**						**07777112**
					SPO		
Actinomyces israelii		58.29	0.81	MLZ 13	1		
Bifidobacterium adolescentis		34.38	0.71		1		
Clostridium tertium		5.61	0.50		99		
07777113	**Low discrimination**						**07777113**
Actinomyces israelii		52.79	0.47	MLZ 13			
Bifidobacterium adolescentis		31.14	0.37	URE 1			
Actinomyces naeslundii		10.01	0.44				
07777120	**Low discrimination**						**07777120**
					SPO		
Actinomyces israelii		56.31	0.72	NAG 6	1		
Clostridium ramosum		21.92	0.49		92		
Clostridium tertium		10.70	0.45		99		
07777121	**Low discrimination**						**07777121**
					SPO		
Actinomyces naeslundii		88.23	0.66	XYL 1	1		
Actinomyces israelii		6.98	0.38		1		
Clostridium ramosum		2.72	0.15		92		
07777122	**Low discrimination**						**07777122**
					SPO		
Clostridium ramosum		40.16	0.59	XYL 1	92		
Actinomyces israelii		38.69	0.75		1		
Clostridium tertium		14.71	0.53		99		
07777123	**Low discrimination**						**07777123**
					SPO		
Actinomyces naeslundii		77.25	0.66	XYL 1	1		
Clostridium ramosum		9.51	0.25		92		
Actinomyces israelii		9.16	0.41		1		
07777130	**Low discrimination**						**07777130**
					SPO		
Clostridium tertium		65.79	0.50	RAF 1	99		
Actinomyces israelii		29.09	0.58		1		
Bifidobacterium dentium		2.25	0.15		1		
07777131	**Low discrimination**						**07777131**
					SPO		
Actinomyces naeslundii		57.73	0.44	XYL 1	1		
Clostridium tertium		29.31	0.15	MLZ 5	99		
Actinomyces israelii		12.96	0.24		1		
07777132	**Low discrimination**						**07777132**
					SPO		
Clostridium tertium		70.43	0.58	RAF 1	99		
Actinomyces israelii		15.57	0.61		1		
Bifidobacterium dentium		8.13	0.32		1		
07777140	**Very good identification**						**07777140**
					COC	SPO	
Actinomyces israelii		97.63	0.85		1	1	
Clostridium butyricum		1.74	0.34		1	99	
Blautia producta		0.35	0.24		99	1	
07777141	**Low discrimination**						**07777141**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		72.25	0.51	URE 1	1		
Actinomyces naeslundii		26.47	0.53		1		
Clostridium butyricum		1.29	0.00		99		
07777142	**Very good identification**						**07777142**
					SPO		
Actinomyces israelii		97.04	0.88		1		
Bifidobacterium adolescentis		2.22	0.54		1		
Clostridium butyricum		0.45	0.27		99		
07777143	**Low discrimination**						**07777143**
Actinomyces israelii		78.63	0.54	URE 1			
Actinomyces naeslundii		19.20	0.53				
Bifidobacterium adolescentis		1.80	0.20				
07777150	**Good identification**						**07777150**
					COC	SPO	
Actinomyces israelii		92.38	0.71	MLZ 13	1	1	
Blautia producta		6.58	0.32		99	1	
Clostridium butyricum		0.58	0.13		1	99	
07777151	**Low discrimination**						**07777151**
					COC		
Actinomyces israelii		83.31	0.37	MLZ 13	1		
Actinomyces naeslundii		10.75	0.31	URE 1	1		
Blautia producta		5.94	-.03		99		
07777152	**Low discrimination**						**07777152**
					COC		
Actinomyces israelii		83.30	0.74	MLZ 13	1		
Bifidobacterium adolescentis		15.62	0.56		1		
Blautia producta		0.44	0.15		99		
07777153	**Low discrimination**						**07777153**
Actinomyces israelii		78.53	0.40	MLZ 13			
Bifidobacterium adolescentis		14.72	0.21	URE 1			
Actinomyces naeslundii		6.75	0.31				
07777160	**Good identification**						**07777160**
					SPO		
Actinomyces israelii		92.11	0.65	NAG 6	1		
Actinomyces naeslundii		3.56	0.50		1		
Clostridium butyricum		2.23	0.16		99		
07777161	**Low discrimination**						**07777161**
Actinomyces naeslundii		85.16	0.53	XYL 1			
Actinomyces israelii		14.84	0.31				
07777162	**Good identification**						**07777162**
					SPO		
Actinomyces israelii		91.64	0.68	NAG 6	1		
Clostridium ramosum		2.47	0.25		92		
Actinomyces naeslundii		2.36	0.50		1		
07777163	**Low discrimination**						**07777163**
Actinomyces naeslundii		79.28	0.53	XYL 1			
Actinomyces israelii		20.72	0.34				
07777170	**Low discrimination**						**07777170**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		82.28	0.51	MLZ 13	1	1	
Blautia producta		10.21	0.15	NAG 6	99	1	
Clostridium tertium		4.83	0.15		1	99	
07777172	**Low discrimination**						**07777172**
					SPO		
Actinomyces israelii		80.08	0.54	MLZ 13	1		
Clostridium tertium		9.41	0.24	NAG 6	99		
Bifidobacterium breve		5.60	0.29		1		
07777200	**Very good identification**						**07777200**
					SPO		
Actinomyces israelii		99.46	0.63	RHA 7	1		
Clostridium butyricum		0.54	0.03		99		
07777202	**Very good identification**						**07777202**
					SPO		
Actinomyces israelii		99.42	0.66	RHA 7	1		
Clostridium clostridioforme		0.44	0.16		66		
Clostridium butyricum		0.14	-.04		99		
07777210	**Good identification**						**07777210**
Actinomyces israelii		100.0	0.49	RHA 7			
07777212	**Low discrimination**						**07777212**
Actinomyces israelii		88.10	0.52	RHA 7			
Clostridium clostridioforme		11.90	0.27	MLZ 13			
07777220	**Low discrimination**						**07777220**
					SPO		
Actinomyces israelii		78.31	0.42	RHA 7	1		
Clostridium ramosum		21.69	0.17	NAG 6	92		
07777222	**Low discrimination**						**07777222**
					SPO		
Actinomyces israelii		53.52	0.45	RHA 7	1		
Clostridium ramosum		39.53	0.27	NAG 6	92		
Clostridium clostridioforme		3.72	0.16		66		
07777240	**Good identification**						**07777240**
Actinomyces israelii		100.0	0.56	RHA 7			
07777242	**Very good identification**						**07777242**
Actinomyces israelii		100.0	0.59	RHA 7			
07777250	**Good identification**						**07777250**
					COC		
Actinomyces israelii		96.32	0.42	RHA 7	1		
Blautia producta		3.68	-.03	MLZ 13	99		
07777252	**Good identification**						**07777252**
Actinomyces israelii		100.0	0.45	RHA 7			
07777260	**Good identification**						**07777260**
Actinomyces israelii		100.0	0.35	RHA 7			
07777262	**Good identification**						**07777262**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.38	RHA 7			
07777300	**Very good identification**						**07777300**
					SPO		
Actinomyces israelii		96.31	0.73	RHA 7	1		
Clostridium butyricum		3.15	0.27		99		
Clostridium ramosum		0.43	0.17		92		
07777301	**Low discrimination**						**07777301**
					SPO		
Actinomyces israelii		87.64	0.39	RHA 7	1		
Actinomyces naeslundii		9.50	0.32	URE 1	1		
Clostridium butyricum		2.86	-.08		99		
07777302	**Very good identification**						**07777302**
					SPO		
Actinomyces israelii		96.37	0.76	RHA 7	1		
Clostridium ramosum		1.14	0.27		92		
Bifidobacterium adolescentis		0.93	0.35		1		
07777303	**Acceptable identification**						**07777303**
Actinomyces israelii		93.26	0.42	RHA 7			
Actinomyces naeslundii		6.74	0.32	URE 1			
07777310	**Very good identification**						**07777310**
					SPO		
Actinomyces israelii		97.88	0.59	RHA 7	1		
Clostridium butyricum		1.13	0.05	MLZ 13	99		
Clostridium tertium		0.63	0.07		99		
07777312	**Low discrimination**						**07777312**
Actinomyces israelii		82.37	0.62	RHA 7			
Clostridium clostridioforme		9.31	0.36	MLZ 13			
Bifidobacterium adolescentis		6.52	0.37				
07777320	**Low discrimination**						**07777320**
					SPO		
Clostridium ramosum		86.13	0.51	XYL 1	92		
Actinomyces israelii		12.57	0.53		1		
Clostridium butyricum		0.56	0.09		99		
07777322	**Low discrimination**						**07777322**
					SPO	GRA	
Clostridium ramosum		86.54	0.61	XYL 1	92	50	
Parabacteroides distasonis		7.09	0.55		1	1	
Actinomyces israelii		4.74	0.56		1	99	
07777330	**Low discrimination**						**07777330**
					SPO		
Actinomyces israelii		54.10	0.39	RHA 7	1		
Clostridium ramosum		25.07	0.17	MLZ 13	92		
Clostridium tertium		16.42	0.15	NAG 6	99		
07777332	**Low discrimination**						**07777332**
					SPO		
Parabacteroides distasonis		49.71	0.50	MAN 1	1		
Clostridium clostridioforme		16.32	0.36		66		
Clostridium ramosum		11.39	0.27		92		
07777340	**Good identification**						**07777340**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		94.83	0.66	RHA 7	1	1	
Blautia producta		4.47	0.24		99	1	
Clostridium butyricum		0.69	0.09		1	99	
07777342	**Very good identification**						**07777342**
					COC		
Actinomyces israelii		99.16	0.69	RHA 7	1		
Blautia producta		0.35	0.07		99		
Bifidobacterium adolescentis		0.31	0.20		1		
07777343	**Good identification**						**07777343**
Actinomyces israelii		100.0	0.35	RHA 7			
07777350	**Low discrimination**						**07777350**
					COC		
Actinomyces israelii		51.36	0.52	RHA 7	1		
Blautia producta		48.64	0.32	MLZ 13	99		
07777352	**Acceptable identification**						**07777352**
					COC		
Actinomyces israelii		91.30	0.55	RHA 7	1		
Blautia producta		6.40	0.15	MLZ 13	99		
Bifidobacterium adolescentis		2.30	0.21		1		
07777360	**Low discrimination**						**07777360**
					COC	SPO	
Actinomyces israelii		79.36	0.46	RHA 7	1	1	
Clostridium ramosum		14.13	0.17	NAG 6	1	92	
Blautia producta		6.52	0.07		99	1	
07777362	**Low discrimination**						**07777362**
					SPO	GRA	
Actinomyces israelii		65.27	0.49	RHA 7	1	99	
Clostridium ramosum		30.99	0.27	NAG 6	92	50	
Parabacteroides distasonis		2.54	0.21		1	1	
07777400	**Low discrimination**						**07777400**
					SPO		
Actinomyces israelii		87.25	0.81		1		
Clostridium butyricum		12.72	0.45		99		
Bifidobacterium adolescentis		0.03	0.15		1		
07777401	**Low discrimination**						**07777401**
					SPO		
Actinomyces israelii		86.22	0.46	URE 1	1		
Clostridium butyricum		12.57	0.11		99		
Actinomyces naeslundii		1.21	0.24		1		
07777402	**Good identification**						**07777402**
					SPO		
Actinomyces israelii		94.11	0.84		1		
Clostridium butyricum		3.56	0.38		99		
Bifidobacterium adolescentis		2.24	0.50		1		
07777403	**Acceptable identification**						**07777403**
					SPO		
Actinomyces israelii		93.37	0.49	URE 1	1		
Clostridium butyricum		3.53	0.04		99		
Bifidobacterium adolescentis		2.23	0.15		1		
07777410	**Good identification**						**07777410**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		93.52	0.67	MLZ 13	1		
Clostridium butyricum		4.80	0.23		99		
Bifidobacterium dentium		1.27	0.10		1		
07777412	**Low discrimination**						**07777412**
Actinomyces israelii		75.39	0.70	MLZ 13			
Bifidobacterium adolescentis		14.70	0.51				
Bifidobacterium dentium		6.91	0.27				
07777413	**Low discrimination**						**07777413**
Actinomyces israelii		77.73	0.35	MLZ 13			
Bifidobacterium adolescentis		15.15	0.17	URE 1			
Bifidobacterium dentium		7.12	-.07				
07777420	**Low discrimination**						**07777420**
					SPO		
Actinomyces israelii		83.43	0.60	NAG 6	1		
Clostridium butyricum		16.57	0.27		99		
07777422	**Good identification**						**07777422**
					SPO		
Actinomyces israelii		93.22	0.63	NAG 6	1		
Clostridium butyricum		4.80	0.20		99		
Clostridium clostridioforme		1.27	0.22		66		
07777430	**Low discrimination**						**07777430**
					SPO		
Actinomyces israelii		69.03	0.46	MLZ 13	1		
Bifidobacterium dentium		26.14	0.15	NAG 6	1		
Clostridium butyricum		4.83	0.05		99		
07777440	**Very good identification**						**07777440**
					COC	SPO	
Actinomyces israelii		95.24	0.74		1	1	
Clostridium butyricum		3.11	0.27		1	99	
Blautia producta		1.65	0.24		99	1	
07777441	**Good identification**						**07777441**
					SPO		
Actinomyces israelii		96.84	0.39	URE 1	1		
Clostridium butyricum		3.16	-.07		99		
07777442	**Very good identification**						**07777442**
					SPO		
Actinomyces israelii		98.30	0.77		1		
Clostridium butyricum		0.83	0.20		99		
Bifidobacterium adolescentis		0.75	0.34		1		
07777443	**Good identification**						**07777443**
Actinomyces israelii		100.0	0.42	URE 1			
07777450	**Low discrimination**						**07777450**
					COC	SPO	
Actinomyces israelii		73.53	0.59	MLZ 13	1	1	
Blautia producta		25.62	0.32		99	1	
Clostridium butyricum		0.84	0.05		1	99	
07777452	**Good identification**						**07777452**
					COC		
Actinomyces israelii		91.48	0.62	MLZ 13	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		5.67	0.36		1		
Blautia producta		2.36	0.15		99		
07777460	**Acceptable identification**						**07777460**
					COC	SPO	
Actinomyces israelii		93.06	0.53	NAG 6	1	1	
Clostridium butyricum		4.13	0.09		1	99	
Blautia producta		2.81	0.07		99	1	
07777462	**Very good identification**						**07777462**
					SPO		
Actinomyces israelii		98.86	0.56	NAG 6	1		
Clostridium butyricum		1.14	0.02		99		
07777470	**Low discrimination**						**07777470**
					COC		
Actinomyces israelii		62.25	0.39	MLZ 13	1		
Blautia producta		37.75	0.15	NAG 6	99		
07777472	**Low discrimination**						**07777472**
					COC		
Actinomyces israelii		89.99	0.42	MLZ 13	1		
Bifidobacterium dentium		5.97	-.03	NAG 6	1		
Blautia producta		4.04	-.01		99		
07777500	**Low discrimination**						**07777500**
					SPO		
Actinomyces israelii		53.00	0.91		1		
Clostridium butyricum		46.35	0.69		99		
Bifidobacterium adolescentis		0.47	0.50		1		
07777501	**Low discrimination**						**07777501**
					GRA	SPO	
Actinomyces israelii		52.50	0.57	URE 1	99	1	
Clostridium butyricum		45.92	0.35		99	99	
Mitsuokella multacida		0.59	0.11		1	1	
07777502	**Low discrimination**						**07777502**
					SPO		
Actinomyces israelii		54.91	0.94		1		
Bifidobacterium adolescentis		32.39	0.84		1		
Clostridium butyricum		12.45	0.62		99		
07777503	**Low discrimination**						**07777503**
					SPO		
Actinomyces israelii		54.75	0.60	URE 1	1		
Bifidobacterium adolescentis		32.29	0.50		1		
Clostridium butyricum		12.41	0.28		99		
07777510	**Low discrimination**						**07777510**
					SPO		
Actinomyces israelii		56.99	0.77	MLZ 13	1		
Bifidobacterium dentium		19.17	0.44		1		
Clostridium butyricum		17.56	0.47		99		
07777511	**Low discrimination**						**07777511**
					SPO		
Actinomyces israelii		56.29	0.43	MLZ 13	1		
Bifidobacterium dentium		18.94	0.10	URE 1	1		
Clostridium butyricum		17.34	0.13		99		
07777512	**Low discrimination**						**07777512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		58.85	0.85				
Bifidobacterium dentium		27.66	0.62				
Actinomyces israelii		12.20	0.80				
07777513	**Low discrimination**						**07777513**
Bifidobacterium adolescentis		58.87	0.51	URE 1			
Bifidobacterium dentium		27.68	0.27				
Actinomyces israelii		12.20	0.46				
07777520	**Low discrimination**						**07777520**
					SPO		
Clostridium butyricum		53.62	0.51	ESL 88	99		
Actinomyces israelii		45.00	0.70	MAN 10	1		
Bifidobacterium dentium		0.64	0.15	NAG 8	1		
07777522	**Low discrimination**						**07777522**
					SPO		
Actinomyces israelii		63.19	0.74	NAG 6	1		
Clostridium butyricum		19.52	0.44		99		
Bifidobacterium dentium		6.03	0.32		1		
07777523	**Low discrimination**						**07777523**
					SPO		
Actinomyces israelii		62.43	0.39	NAG 6	1		
Clostridium butyricum		19.29	0.10	URE 1	99		
Actinomyces naeslundii		6.50	0.32		1		
07777530	**Low discrimination**						**07777530**
					SPO		
Bifidobacterium dentium		85.95	0.49	ESL 99	1		
Actinomyces israelii		9.17	0.56	BGL 91	1		
Clostridium butyricum		3.85	0.29		99		
07777532	**Very good identification**						**07777532**
Bifidobacterium dentium		96.19	0.66	ESL 99			
Actinomyces israelii		1.52	0.59				
Bifidobacterium adolescentis		1.16	0.51				
07777540	**Low discrimination**						**07777540**
					COC	SPO	
Actinomyces israelii		61.33	0.84		1	1	
Blautia producta		26.34	0.58		99	1	
Clostridium butyricum		11.99	0.51		1	99	
07777541	**Low discrimination**						**07777541**
					COC	SPO	
Actinomyces israelii		60.46	0.50	URE 1	1	1	
Blautia producta		25.97	0.24		99	1	
Clostridium butyricum		11.82	0.17		1	99	
07777542	**Low discrimination**						**07777542**
					SPO		
Actinomyces israelii		78.72	0.87		1		
Bifidobacterium adolescentis		14.76	0.68		1		
Clostridium butyricum		3.99	0.44		99		
07777543	**Low discrimination**						**07777543**
					SPO		
Actinomyces israelii		78.74	0.53	URE 1	1		
Bifidobacterium adolescentis		14.76	0.34		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		3.99	0.10		99		
07777550	**Low discrimination**						**07777550**
					COC	SPO	
Blautia producta		88.63	0.66	ESL 99	99	1	
Actinomyces israelii		10.28	0.70		1	1	
Clostridium butyricum		0.71	0.29		1	99	
07777552	**Low discrimination**						**07777552**
					COC		
Bifidobacterium adolescentis		47.18	0.70	SOR 11	1		
Actinomyces israelii		30.77	0.73		1		
Blautia producta		19.66	0.50		99		
07777553	**Low discrimination**						**07777553**
					COC		
Bifidobacterium adolescentis		47.46	0.36	SOR 11	1		
Actinomyces israelii		30.95	0.39	URE 1	1		
Blautia producta		19.77	0.15		99		
07777560	**Low discrimination**						**07777560**
					COC	SPO	
Actinomyces israelii		49.40	0.63	NAG 6	1	1	
Blautia producta		36.93	0.41		99	1	
Clostridium butyricum		13.16	0.33		1	99	
07777562	**Low discrimination**						**07777562**
					COC	SPO	
Actinomyces israelii		86.24	0.66	NAG 6	1	1	
Clostridium butyricum		5.96	0.26		1	99	
Blautia producta		4.78	0.25		99	1	
07777570	**Acceptable identification**						**07777570**
					COC		
Blautia producta		91.69	0.50	ESL 99	99		
Actinomyces israelii		6.11	0.49	NAG 10	1		
Bifidobacterium dentium		1.49	0.15		1		
07777600	**Good identification**						**07777600**
					SPO		
Actinomyces israelii		94.35	0.61	RHA 7	1		
Clostridium butyricum		5.65	0.19		99		
07777602	**Good identification**						**07777602**
					SPO		
Actinomyces israelii		90.62	0.64	RHA 7	1		
Clostridium clostridioforme		7.69	0.37		66		
Clostridium butyricum		1.41	0.12		99		
07777610	**Acceptable identification**						**07777610**
					SPO		
Actinomyces israelii		92.72	0.47	RHA 7	1		
Clostridium clostridioforme		3.63	0.14	MLZ 13	66		
Clostridium butyricum		1.96	-.03		99		
07777612	**Low discrimination**						**07777612**
Clostridium clostridioforme		71.35	0.48	MAN 1			
Actinomyces israelii		27.59	0.50				
Bifidobacterium adolescentis		0.72	0.17				
07777620	**Acceptable identification**						**07777620**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		92.46	0.41	RHA 7	1		
Clostridium butyricum		7.55	0.01	NAG 6	99		
07777622	**Low discrimination**						**07777622**
					GRA		
Clostridium clostridioforme		49.74	0.37	MAN 1	50		
Actinomyces israelii		37.43	0.44		99		
Bacteroides ovatus		10.34	0.15		1		
07777632	**Acceptable identification**						**07777632**
					GRA		
Clostridium clostridioforme		93.67	0.48	MAN 1	50		
Bacteroides ovatus		2.85	0.12		1		
Actinomyces israelii		2.31	0.30		99		
07777640	**Low discrimination**						**07777640**
					COC	SPO	
Actinomyces israelii		80.39	0.54	RHA 7	1	1	
Blautia producta		18.54	0.24		99	1	
Clostridium butyricum		1.08	0.01		1	99	
07777642	**Very good identification**						**07777642**
					COC	SPO	
Actinomyces israelii		97.99	0.57	RHA 7	1	1	
Blautia producta		1.67	0.07		99	1	
Clostridium butyricum		0.34	-.06		1	99	
07777652	**Low discrimination**						**07777652**
					COC		
Actinomyces israelii		70.92	0.43	RHA 7	1		
Blautia producta		24.32	0.15	MLZ 13	99		
Clostridium clostridioforme		4.76	0.14		1		
07777662	**Good identification**						**07777662**
Actinomyces israelii		100.0	0.37	RHA 7			
07777700	**Low discrimination**						**07777700**
					GRA	SPO	
Actinomyces israelii		70.96	0.72	RHA 7	99	1	
Clostridium butyricum		25.50	0.43		99	99	
Mitsuokella multacida		1.79	0.32		1	1	
07777701	**Low discrimination**						**07777701**
					GRA	SPO	
Actinomyces israelii		61.09	0.38	RHA 7	99	1	
Clostridium butyricum		21.96	0.09	URE 1	99	99	
Mitsuokella multacida		16.95	0.15		1	1	
07777702	**Low discrimination**						**07777702**
					SPO		
Actinomyces israelii		79.84	0.75	RHA 7	1		
Clostridium butyricum		7.44	0.36		99		
Bifidobacterium adolescentis		6.32	0.50		1		
07777703	**Low discrimination**						**07777703**
					SPO		
Actinomyces israelii		80.43	0.41	RHA 7	1		
Clostridium butyricum		7.49	0.02	URE 1	99		
Bifidobacterium adolescentis		6.37	0.15		1		
07777710	**Low discrimination**						**07777710**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		58.12	0.58	RHA 7	1	1	
Blautia producta		26.16	0.32	MLZ 13	99	1	
Clostridium butyricum		7.36	0.21		1	99	
07777712	**Low discrimination**						**07777712**
Clostridium clostridioforme		51.43	0.57	MAN 1			
Actinomyces israelii		23.76	0.61				
Bifidobacterium adolescentis		15.38	0.51				
07777720	**Low discrimination**						**07777720**
					GRA	SPO	
Actinomyces israelii		53.12	0.51	RHA 7	99	1	
Clostridium butyricum		26.01	0.25	NAG 6	99	99	
Bacteroides ovatus		5.50	0.15		1	1	
07777722	**Low discrimination**						**07777722**
Bacteroides ovatus		65.33	0.49	IND 97			
Parabacteroides distasonis		10.71	0.51	ESL 91			
Clostridium clostridioforme		10.63	0.46				
07777732	**Low discrimination**						**07777732**
					GRA		
Clostridium clostridioforme		43.38	0.57	MAN 1	50		
Bacteroides ovatus		39.03	0.46		1		
Bifidobacterium dentium		10.84	0.32		99		
07777740	**Low discrimination**						**07777740**
					COC	SPO	
Blautia producta		83.29	0.58	ESL 99	99	1	
Actinomyces israelii		14.59	0.65		1	1	
Clostridium butyricum		1.17	0.25		1	99	
07777742	**Low discrimination**						**07777742**
					COC		
Actinomyces israelii		67.78	0.68	RHA 7	1		
Blautia producta		28.65	0.41		99		
Bifidobacterium adolescentis		1.71	0.34		1		
07777750	**Very good identification**						**07777750**
					COC	GRA	
Blautia producta		98.99	0.66	ESL 99	99	99	
Actinomyces israelii		0.86	0.51		1	99	
Mitsuokella multacida		0.13	0.24		1	1	
07777752	**Low discrimination**						**07777752**
					COC		
Blautia producta		86.82	0.50	ESL 99	99		
Actinomyces israelii		10.23	0.54	BGL 10	1		
Bifidobacterium adolescentis		2.11	0.36		1		
07777760	**Low discrimination**						**07777760**
					COC	GRA	
Blautia producta		87.93	0.41	ESL 99	99	99	
Actinomyces israelii		8.85	0.44	NAG 10	1	99	
Mitsuokella multacida		2.25	0.21		1	1	
07777762	**Low discrimination**						**07777762**
					COC	GRA	
Actinomyces israelii		38.86	0.47	RHA 7	1	99	
Blautia producta		28.59	0.25	NAG 6	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides ovatus		28.47	0.25		1	1		
07777770	**Good identification**							**07777770**
					COC	GRA		
Blautia producta		99.22	0.50	ESL 99	99	99		
Actinomyces israelii		0.50	0.30	NAG 10	1	99		
Mitsuokella multacida		0.28	0.13		1	1		
10000000	**Low discrimination**							**10000000**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.46	1.00		1	1		
Peptoniphilus asaccharolyticus		24.46	1.00		99	99		
Porphyromonas asaccharolytica		24.21	1.00		1	1		
10000001	**Low discrimination**							**10000001**
					COC	GRA		
Anaerococcus prevotii		20.51	0.75	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.89	0.66		1	1		
Peptoniphilus asaccharolyticus		15.89	0.66		99	99		
10000002	**Low discrimination**							**10000002**
					COC	GRA		
Porphyromonas asaccharolytica		54.11	0.76	BGL 4	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.25	0.66		1	1		
Peptoniphilus asaccharolyticus		13.25	0.66		99	99		
10000003	**Low discrimination**							**10000003**
					COC	GRA		
Anaerococcus prevotii		77.12	0.64	IND 9	99	99		
Porphyromonas asaccharolytica		9.85	0.42		1	1		
Peptococcus niger		2.99	0.36		99	99		
10000010	**Low discrimination**							**10000010**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.46	0.66	MLZ 1	1	1		
Peptoniphilus asaccharolyticus		24.46	0.66		99	99		
Porphyromonas asaccharolytica		24.20	0.66		1	1		
10000011	**Low discrimination**							**10000011**
					COC	GRA		
Anaerococcus prevotii		20.40	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.80	0.32	MLZ 1	1	1		
Peptoniphilus asaccharolyticus		15.80	0.32		99	99		
10000012	**Low discrimination**							**10000012**
					COC	GRA		
Porphyromonas asaccharolytica		54.09	0.42	MLZ 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.25	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.25	0.32		99	99		
10000020	**Low discrimination**							**10000020**
					COC	GRA	SPO	
Clostridium cadaveris		24.19	0.78	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.08	0.66		1	1	1	
Peptoniphilus asaccharolyticus		18.08	0.66		99	99	1	
10000021	**Low discrimination**							**10000021**
					COC	GRA	SPO	
Clostridium cadaveris		15.96	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		15.40	0.41	URE 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		11.93	0.32		1	1	1	
10000022	**Low discrimination**							**10000022**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Pseudoflavonifractor capillosus		79.36	0.66	IND 1	1	1		
Porphyromonas asaccharolytica		7.68	0.42		1	1		
Peptococcus niger		2.62	0.37		99	99		
10000030	**Low discrimination**							**10000030**
					COC	GRA	SPO	
Clostridium cadaveris		24.16	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.06	0.32	MLZ 1	1	1	1	
Peptoniphilus asaccharolyticus		18.06	0.32		99	99	1	
10000040	**Low discrimination**							**10000040**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.39	0.66	SOR 1	1	1		
Peptoniphilus asaccharolyticus		24.39	0.66		99	99		
Porphyromonas asaccharolytica		24.14	0.66		1	1		
10000041	**Low discrimination**							**10000041**
					COC	GRA		
Anaerococcus prevotii		18.55	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		14.37	0.32	SOR 1	1	1		
Peptoniphilus asaccharolyticus		14.37	0.32		99	99		
10000042	**Low discrimination**							**10000042**
					COC	GRA		
Porphyromonas asaccharolytica		54.02	0.42	SOR 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.23	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.23	0.32		99	99		
10000060	**Low discrimination**							**10000060**
					COC	GRA	SPO	
Clostridium cadaveris		23.94	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		17.90	0.32	SOR 1	1	1	1	
Peptoniphilus asaccharolyticus		17.90	0.32		99	99	1	
10000100	**Low discrimination**							**10000100**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.46	0.66	LAC 1	1	1		
Peptoniphilus asaccharolyticus		24.46	0.66		99	99		
Porphyromonas asaccharolytica		24.21	0.66		1	1		
10000101	**Low discrimination**							**10000101**
					COC	GRA		
Anaerococcus prevotii		20.51	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.89	0.32	LAC 1	1	1		
Peptoniphilus asaccharolyticus		15.89	0.32		99	99		
10000102	**Low discrimination**							**10000102**
					COC	GRA		
Porphyromonas asaccharolytica		54.13	0.42	LAC 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.26	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.26	0.32		99	99		
10000120	**Low discrimination**							**10000120**
					COC	GRA	SPO	
Clostridium cadaveris		24.20	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.09	0.32	LAC 1	1	1	1	
Peptoniphilus asaccharolyticus		18.09	0.32		99	99	1	
10000200	**Low discrimination**							**10000200**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.46	0.66	RHA 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptoniphilus asaccharolyticus		24.46	0.66		99	99		
Porphyromonas asaccharolytica		24.21	0.66		1	1		
10000201	**Low discrimination**							**10000201**
					COC	GRA		
Anaerococcus prevotii		20.51	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.89	0.32	RHA 1	1	1		
Peptoniphilus asaccharolyticus		15.89	0.32		99	99		
10000202	**Low discrimination**							**10000202**
					COC	GRA		
Porphyromonas asaccharolytica		54.13	0.42	RHA 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.26	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.26	0.32		99	99		
10000220	**Low discrimination**							**10000220**
					COC	GRA	SPO	
Clostridium cadaveris		24.20	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.09	0.32	RHA 1	1	1	1	
Peptoniphilus asaccharolyticus		18.09	0.32		99	99	1	
10000400	**Low discrimination**							**10000400**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.46	0.66	ARA 1	1	1		
Peptoniphilus asaccharolyticus		24.46	0.66		99	99		
Porphyromonas asaccharolytica		24.21	0.66		1	1		
10000401	**Low discrimination**							**10000401**
					COC	GRA		
Anaerococcus prevotii		20.51	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.89	0.32	ARA 1	1	1		
Peptoniphilus asaccharolyticus		15.89	0.32		99	99		
10000402	**Low discrimination**							**10000402**
					COC	GRA		
Porphyromonas asaccharolytica		54.13	0.42	ARA 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.26	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.26	0.32		99	99		
10000420	**Low discrimination**							**10000420**
					COC	GRA	SPO	
Clostridium cadaveris		24.20	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.09	0.32	ARA 1	1	1	1	
Peptoniphilus asaccharolyticus		18.09	0.32		99	99	1	
10001000	**Good identification**							**10001000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.57	1.00		1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.66		1	1		
Peptoniphilus asaccharolyticus		2.04	0.66		99	99		
10001001	**Low discrimination**							**10001001**
					COC	GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		83.18	0.66	URE 1	1	1	1	
Clostridium sordellii		6.46	0.49		1	99	99	
Anaerococcus prevotii		2.37	0.41		99	99	1	
10001002	**Low discrimination**							**10001002**
Fusobacterium necrophorum subsp. necrophorum		85.13	0.66	BGL 1				
Porphyromonas asaccharolytica		7.66	0.42					
Fusobacterium nucleatum subsp. nucleatum		1.88	0.32					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
10001010	**Good identification**						**10001010**
					COC	GRA	
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	MLZ 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1	
Peptoniphilus asaccharolyticus		2.04	0.32		99	99	
10001020	**Low discrimination**						**10001020**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		89.16	0.66	NAG 1	1	1	
Clostridium cadaveris		2.63	0.44		99	99	
Fusobacterium nucleatum subsp. nucleatum		1.97	0.32		1	1	
10001022	**Low discrimination**						**10001022**
					GRA		
Pseudoflavonifractor capillosus		85.01	0.50	IND 1	1		
Fusobacterium necrophorum subsp. necrophorum		8.31	0.32	GAL 10	1		
Atopobium minutum		4.96	0.43		99		
10001040	**Good identification**						**10001040**
					COC	GRA	
Fusobacterium necrophorum subsp. necrophorum		92.55	0.66	SOR 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1	
Peptoniphilus asaccharolyticus		2.04	0.32		99	99	
10001100	**Good identification**						**10001100**
					COC	GRA	
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	LAC 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1	
Peptoniphilus asaccharolyticus		2.04	0.32		99	99	
10001200	**Good identification**						**10001200**
					COC	GRA	
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	RHA 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1	
Peptoniphilus asaccharolyticus		2.04	0.32		99	99	
10001400	**Good identification**						**10001400**
					COC	GRA	
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	ARA 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1	
Peptoniphilus asaccharolyticus		2.04	0.32		99	99	
10002000	**Low discrimination**						**10002000**
					COC	GRA	
Fusobacterium nucleatum subsp. nucleatum		24.46	0.66	MAN 1	1	1	
Peptoniphilus asaccharolyticus		24.46	0.66		99	99	
Porphyromonas asaccharolytica		24.21	0.66		1	1	
10002001	**Low discrimination**						**10002001**
					COC	GRA	
Anaerococcus prevotii		20.51	0.41	IND 9	99	99	
Fusobacterium nucleatum subsp. nucleatum		15.89	0.32	MAN 1	1	1	
Peptoniphilus asaccharolyticus		15.89	0.32		99	99	
10002002	**Low discrimination**						**10002002**
					COC	GRA	
Porphyromonas asaccharolytica		54.13	0.42	MAN 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		13.26	0.32	BGL 4	1	1	
Peptoniphilus asaccharolyticus		13.26	0.32		99	99	
10002020	**Low discrimination**						**10002020**
					COC	GRA	SPO

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium cadaveris		24.19	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.08	0.32	MAN 1	1	1	1	
Peptoniphilus asaccharolyticus		18.08	0.32		99	99	1	
10003000	**Good identification**							**10003000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	MAN 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1		
Peptoniphilus asaccharolyticus		2.04	0.32		99	99		
10004000	**Low discrimination**							**10004000**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.45	0.66	XYL 1	1	1		
Peptoniphilus asaccharolyticus		24.45	0.66		99	99		
Porphyromonas asaccharolytica		24.20	0.66		1	1		
10004001	**Low discrimination**							**10004001**
					COC	GRA		
Anaerococcus prevotii		20.51	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.89	0.32	XYL 1	1	1		
Peptoniphilus asaccharolyticus		15.89	0.32		99	99		
10004002	**Low discrimination**							**10004002**
					COC	GRA		
Porphyromonas asaccharolytica		54.12	0.42	XYL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.26	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.26	0.32		99	99		
10004020	**Low discrimination**							**10004020**
					COC	GRA	SPO	
Clostridium cadaveris		24.19	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.09	0.32	XYL 1	1	1	1	
Peptoniphilus asaccharolyticus		18.09	0.32		99	99	1	
10005000	**Good identification**							**10005000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	XYL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1		
Peptoniphilus asaccharolyticus		2.04	0.32		99	99		
10010000	**Low discrimination**							**10010000**
					COC	GRA		
Fusobacterium varium		53.32	0.84		1	1		
Fusobacterium nucleatum subsp. nucleatum		11.23	0.66		1	1		
Peptoniphilus asaccharolyticus		11.23	0.66		99	99		
10010001	**Low discrimination**							**10010001**
					COC	GRA	SPO	
Clostridium sordellii		64.68	0.62	GLU 95	1	99	99	
Fusobacterium varium		15.38	0.50	MLT 90	1	1	1	
Anaerococcus prevotii		4.18	0.41		99	99	1	
10010002	**Low discrimination**							**10010002**
					GRA			
Fusobacterium varium		32.34	0.50	BGL 1	1			
Porphyromonas asaccharolytica		27.81	0.42		1			
Atopobium minutum		15.46	0.58		99			
10010010	**Low discrimination**							**10010010**
					COC	GRA		
Fusobacterium varium		53.30	0.50	MLZ 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.23	0.32		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptoniphilus asaccharolyticus		11.23	0.32		99	99		
10010020	**Good identification**							**10010020**
					GRA	SPO		
Clostridium cadaveris		92.48	0.78	GLU 95	99	99		
Fusobacterium varium		3.32	0.50		1	1		
Atopobium minutum		0.79	0.53		99	1		
10010021	**Low discrimination**							**10010021**
					GRA	SPO		
Clostridium cadaveris		79.99	0.44	GLU 95	99	99		
Clostridium sordellii		12.06	0.28	URE 1	99	99		
Fusobacterium varium		2.87	0.16		1	1		
10010022	**Low discrimination**							**10010022**
					GRA	SPO		
Pseudoflavonifractor capillosus		67.64	0.50	IND 1	1	1		
Clostridium cadaveris		19.31	0.44	FRU 10	99	99		
Atopobium minutum		10.92	0.50		99	1		
10010030	**Acceptable identification**							**10010030**
					GRA	SPO		
Clostridium cadaveris		92.70	0.44	GLU 95	99	99		
Fusobacterium varium		3.32	0.16	MLZ 1	1	1		
Atopobium minutum		0.79	0.19		99	1		
10010040	**Low discrimination**							**10010040**
					COC	GRA		
Fusobacterium varium		52.12	0.50	SOR 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		10.98	0.32		1	1		
Peptoniphilus asaccharolyticus		10.98	0.32		99	99		
10010041	**Low discrimination**							**10010041**
					COC	GRA	SPO	
Clostridium sordellii		88.25	0.38	GLU 95	1	99	99	
Fusobacterium varium		5.09	0.16	MLT 90	1	1	1	
Anaerococcus prevotii		1.38	0.06	SOR 4	99	99	1	
10010060	**Low discrimination**							**10010060**
					GRA	SPO	CAT	
Clostridium cadaveris		84.88	0.44	GLU 95	99	99	1	
Propionibacterium acnes		7.37	0.29	SOR 1	99	1	95	
Fusobacterium varium		3.04	0.16		1	1	1	
10010100	**Low discrimination**							**10010100**
					COC	GRA		
Fusobacterium varium		53.25	0.50	LAC 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.22	0.32		1	1		
Peptoniphilus asaccharolyticus		11.22	0.32		99	99		
10010120	**Acceptable identification**							**10010120**
					GRA	SPO		
Clostridium cadaveris		92.54	0.44	GLU 95	99	99		
Fusobacterium varium		3.32	0.16	LAC 1	1	1		
Atopobium minutum		0.79	0.19		99	1		
10010200	**Low discrimination**							**10010200**
					COC	GRA		
Fusobacterium varium		53.36	0.50	RHA 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.24	0.32		1	1		
Peptoniphilus asaccharolyticus		11.24	0.32		99	99		
10010220	**Acceptable identification**							**10010220**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Clostridium cadaveris		92.70	0.44	GLU 95	99	99		
Fusobacterium varium		3.32	0.16	RHA 1	1	1		
Atopobium minutum		0.79	0.19		99	1		
10010400	**Low discrimination**							**10010400**
					COC	GRA		
Fusobacterium varium		53.36	0.50	ARA 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.24	0.32		1	1		
Peptoniphilus asaccharolyticus		11.24	0.32		99	99		
10010420	**Acceptable identification**							**10010420**
					GRA	SPO		
Clostridium cadaveris		92.70	0.44	GLU 95	99	99		
Fusobacterium varium		3.32	0.16	ARA 1	1	1		
Atopobium minutum		0.79	0.19		99	1		
10011000	**Low discrimination**							**10011000**
					GRA			
Fusobacterium necrophorum subsp. necrophorum		82.29	0.66	FRU 1	1			
Fusobacterium varium		8.61	0.50		1			
Atopobium minutum		2.06	0.53		99			
10011001	**Low discrimination**							**10011001**
					GRA	SPO		
Clostridium sordellii		86.01	0.49	GLU 95	99	99		
Fusobacterium necrophorum subsp. necrophorum		11.18	0.32	MLT 90	1	1		
Fusobacterium varium		1.17	0.16		1	1		
10011002	**Low discrimination**							**10011002**
					GRA			
Atopobium minutum		56.67	0.50	IND 1	99			
Fusobacterium necrophorum subsp. necrophorum		34.32	0.32		1			
Fusobacterium varium		3.59	0.16		1			
10011020	**Low discrimination**							**10011020**
					GRA	SPO	CAT	
Clostridium cadaveris		47.90	0.44	GLU 95	99	99	1	
Propionibacterium acnes		18.95	0.40	GAL 1	99	1	95	
Fusobacterium necrophorum subsp. necrophorum		16.41	0.32		1	1	1	
10012000	**Low discrimination**							**10012000**
					COC	GRA		
Fusobacterium varium		50.93	0.50	MAN 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		10.73	0.32		1	1		
Peptoniphilus asaccharolyticus		10.73	0.32		99	99		
10012020	**Low discrimination**							**10012020**
					GRA	SPO	CAT	
Clostridium cadaveris		75.34	0.44	GLU 95	99	99	1	
Propionibacterium acnes		16.77	0.36	MAN 1	99	1	95	
Fusobacterium varium		2.70	0.16		1	1	1	
10013020	**Acceptable identification**							**10013020**
					GRA	SPO	CAT	
Propionibacterium acnes		93.13	0.36	NIT 92	99	1	95	
Clostridium cadaveris		4.23	0.10	GLU 99	99	99	1	
Fusobacterium necrophorum subsp. necrophorum		1.45	-.03		1	1	1	
10014000	**Low discrimination**							**10014000**
					COC	GRA		
Fusobacterium varium		82.45	0.60	XYL 4	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium nucleatum subsp. nucleatum		4.21	0.32		1	1		
Peptoniphilus asaccharolyticus		4.21	0.32		99	99		
10014020	**Low discrimination**							**10014020**
					GRA	SPO		
Clostridium cadaveris		83.98	0.44	GLU 95	99	99		
Fusobacterium varium		12.42	0.26	XYL 1	1	1		
Atopobium minutum		0.72	0.19		99	1		
10020000	**Low discrimination**							**10020000**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.43	0.66	TRE 1	1	1		
Peptoniphilus asaccharolyticus		24.43	0.66		99	99		
Porphyromonas asaccharolytica		24.18	0.66		1	1		
10020001	**Low discrimination**							**10020001**
					COC	GRA		
Anaerococcus prevotii		20.50	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.88	0.32	TRE 1	1	1		
Peptoniphilus asaccharolyticus		15.88	0.32		99	99		
10020002	**Low discrimination**							**10020002**
					COC	GRA		
Porphyromonas asaccharolytica		53.79	0.42	TRE 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.18	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.18	0.32		99	99		
10020020	**Low discrimination**							**10020020**
					COC	GRA	SPO	
Clostridium cadaveris		62.44	0.56	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		8.96	0.32	TRE 5	1	1	1	
Peptoniphilus asaccharolyticus		8.96	0.32		99	99	1	
10021000	**Good identification**							**10021000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	TRE 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1		
Peptoniphilus asaccharolyticus		2.04	0.32		99	99		
10030000	**Low discrimination**							**10030000**
					COC	GRA		
Fusobacterium varium		52.78	0.50	TRE 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.12	0.32		1	1		
Peptoniphilus asaccharolyticus		11.12	0.32		99	99		
10030020	**Very good identification**							**10030020**
					GRA	SPO		
Clostridium cadaveris		98.51	0.56	GLU 95	99	99		
Fusobacterium varium		0.68	0.16	TRE 5	1	1		
Atopobium minutum		0.16	0.19		99	1		
10040000	**Low discrimination**							**10040000**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.46	0.66	CEL 1	1	1		
Peptoniphilus asaccharolyticus		24.46	0.66		99	99		
Porphyromonas asaccharolytica		24.21	0.66		1	1		
10040001	**Low discrimination**							**10040001**
					COC	GRA		
Anaerococcus prevotii		20.51	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.89	0.32	CEL 1	1	1		
Peptoniphilus asaccharolyticus		15.89	0.32		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
10040002	**Low discrimination**							**10040002**
					COC	GRA		
Porphyromonas asaccharolytica		54.13	0.42	CEL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.26	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.26	0.32		99	99		
10040020	**Low discrimination**							**10040020**
					COC	GRA	SPO	
Clostridium cadaveris		24.20	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		18.09	0.32	CEL 1	1	1	1	
Peptoniphilus asaccharolyticus		18.09	0.32		99	99	1	
10041000	**Good identification**							**10041000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.57	0.66	CEL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1		
Peptoniphilus asaccharolyticus		2.04	0.32		99	99		
10050000	**Low discrimination**							**10050000**
					COC	GRA		
Fusobacterium varium		53.36	0.50	CEL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.24	0.32		1	1		
Peptoniphilus asaccharolyticus		11.24	0.32		99	99		
10050020	**Acceptable identification**							**10050020**
					GRA	SPO		
Clostridium cadaveris		92.70	0.44	GLU 95	99	99		
Fusobacterium varium		3.32	0.16	CEL 1	1	1		
Atopobium minutum		0.79	0.19		99	1		
10100000	**Low discrimination**							**10100000**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		20.47	0.66	MLT 1	1	1		
Peptoniphilus asaccharolyticus		20.47	0.66		99	99		
Porphyromonas asaccharolytica		20.26	0.66		1	1		
10100001	**Very good identification**							**10100001**
					COC	GRA	SPO	
Clostridium sordellii		96.30	0.78	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.41		99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.54	0.32		1	1	1	
10100002	**Low discrimination**							**10100002**
					COC	GRA		
Porphyromonas asaccharolytica		47.57	0.42	MLT 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.65	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		11.65	0.32		99	99		
10100003	**Low discrimination**							**10100003**
					COC	GRA	SPO	
Clostridium sordellii		80.84	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		14.37	0.30	BGL 1	99	99	1	
Porphyromonas asaccharolytica		1.84	0.08		1	1	1	
10100011	**Acceptable identification**							**10100011**
					COC	SPO		
Clostridium sordellii		94.57	0.44	GLU 95	1	99		
Clostridium novyi biovar A		1.90	0.32	MLZ 1	1	99		
Anaerococcus prevotii		0.68	0.06		99	1		
10100020	**Low discrimination**							**10100020**
					GRA	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium cadaveris		46.35	0.56	GLU 95	99	99		
Clostridium bifermentans		24.40	0.54	MLT 5	99	99		
Fusobacterium nucleatum subsp. nucleatum		6.65	0.32		1	1		
10100021	**Acceptable identification**							**10100021**
Clostridium sordellii		90.68	0.44	GLU 95				
Clostridium cadaveris		3.52	0.22	NAG 1				
Clostridium bifermentans		1.85	0.19					
10100040	**Low discrimination**							**10100040**
					COC	GRA	SPO	
Clostridium bifermentans		34.10	0.50	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		14.24	0.32	SOR 4	1	1	1	
Peptoniphilus asaccharolyticus		14.24	0.32		99	99	1	
10100041	**Good identification**							**10100041**
Clostridium sordellii		98.51	0.55	GLU 95				
Clostridium novyi biovar A		0.32	0.29	SOR 4				
Clostridium bifermentans		0.32	0.16					
10100101	**Good identification**							**10100101**
					COC	GRA	SPO	
Clostridium sordellii		96.40	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	LAC 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.54	-.03		1	1	1	
10100201	**Good identification**							**10100201**
					COC	GRA	SPO	
Clostridium sordellii		96.40	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	RHA 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.54	-.03		1	1	1	
10100401	**Good identification**							**10100401**
					COC	GRA	SPO	
Clostridium sordellii		96.40	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	ARA 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.54	-.03		1	1	1	
10101000	**Good identification**							**10101000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		90.23	0.66	MLT 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		1.99	0.32		1	1		
Peptoniphilus asaccharolyticus		1.99	0.32		99	99		
10101001	**Very good identification**							**10101001**
					COC	GRA	SPO	
Clostridium sordellii		98.37	0.65	GLU 95	1	99	99	
Fusobacterium necrophorum subsp. necrophorum		1.42	0.32		1	1	1	
Anaerococcus prevotii		0.04	0.06		99	99	1	
10101041	**Good identification**							**10101041**
					GRA	SPO		
Clostridium sordellii		99.45	0.42	GLU 95	99	99		
Fusobacterium necrophorum subsp. necrophorum		0.35	-.03	SOR 4	1	1		
Clostridium novyi biovar A		0.20	0.12		99	99		
10102001	**Good identification**							**10102001**
					COC	GRA	SPO	
Clostridium sordellii		96.40	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	MAN 1	99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium nucleatum subsp. nucleatum		0.54	-.03		1	1	1	
10104001	**Good identification**							**10104001**
					COC	GRA	SPO	
Clostridium sordellii		96.40	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	XYL 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.54	-.03		1	1	1	
10110000	**Low discrimination**							**10110000**
					GRA	SPO		
Prevotella intermedia		54.97	0.68	GLU 90	1	1		
Fusobacterium varium		14.90	0.50		1	1		
Clostridium bifermentans		9.50	0.52		99	99		
10110001	**Very good identification**							**10110001**
					GRA	SPO		
Clostridium sordellii		99.82	0.78	GLU 95	99	99		
Prevotella intermedia		0.10	0.34		1	1		
Fusobacterium varium		0.03	0.16		1	1		
10110003	**Good identification**							**10110003**
					COC	SPO		
Clostridium sordellii		99.82	0.44	GLU 95	1	99		
Anaerococcus prevotii		0.18	-.04	BGL 1	99	1		
10110011	**Good identification**							**10110011**
Clostridium sordellii		99.78	0.44	GLU 95				
Clostridium novyi biovar A		0.22	0.15	MLZ 1				
10110020	**Good identification**							**10110020**
					GRA	SPO		
Clostridium cadaveris		93.36	0.56	GLU 95	99	99		
Clostridium bifermentans		2.59	0.32	MLT 5	99	99		
Prevotella intermedia		2.37	0.34		1	1		
10110021	**Good identification**							**10110021**
Clostridium sordellii		96.17	0.44	GLU 95				
Clostridium cadaveris		3.73	0.22	NAG 1				
Clostridium bifermentans		0.10	-.02					
10110041	**Good identification**							**10110041**
Clostridium sordellii		99.96	0.55	GLU 95				
Clostridium novyi biovar A		0.04	0.12	SOR 4				
10110101	**Good identification**							**10110101**
Clostridium sordellii		100.0	0.44	GLU 95				
10110201	**Good identification**							**10110201**
Clostridium sordellii		100.0	0.44	GLU 95				
10110401	**Good identification**							**10110401**
Clostridium sordellii		100.0	0.44	GLU 95				
10111000	**Very good identification**							**10111000**
					GRA	SPO		
Prevotella intermedia		95.04	0.68	GLU 90	1	1		
Fusobacterium necrophorum subsp. necrophorum		2.49	0.32		1	1		
Clostridium sordellii		1.74	0.31		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
10111001	**Very good identification**							**10111001**
					GRA	SPO		
Clostridium sordellii		99.43	0.65	GLU 95	99	99		
Prevotella intermedia		0.56	0.34		1	1		
Fusobacterium necrophorum subsp. necrophorum		0.02	-.03		1	1		
10111041	**Good identification**							**10111041**
Clostridium sordellii		100.0	0.42	GLU 95				
10112001	**Good identification**							**10112001**
Clostridium sordellii		100.0	0.44	GLU 95				
10114001	**Good identification**							**10114001**
					GRA	SPO		
Clostridium sordellii		99.89	0.44	GLU 95	99	99		
Fusobacterium varium		0.11	-.08	XYL 1	1	1		
10120001	**Good identification**							**10120001**
					COC	GRA	SPO	
Clostridium sordellii		96.16	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	TRE 1	99	99	1	
Porphyromonas asaccharolytica		0.53	-.03		1	1	1	
10130001	**Good identification**							**10130001**
Clostridium sordellii		100.0	0.44	GLU 95				
10140001	**Good identification**							**10140001**
					COC	GRA	SPO	
Clostridium sordellii		96.40	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	CEL 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.54	-.03		1	1	1	
10150001	**Good identification**							**10150001**
Clostridium sordellii		100.0	0.44	GLU 95				
10200000	**Low discrimination**							**10200000**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.31	0.66	SAL 1	1	1		
Peptoniphilus asaccharolyticus		24.31	0.66		99	99		
Porphyromonas asaccharolytica		24.06	0.66		1	1		
10200001	**Low discrimination**							**10200001**
					COC	GRA		
Anaerococcus prevotii		20.42	0.41	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		15.82	0.32	SAL 1	1	1		
Peptoniphilus asaccharolyticus		15.82	0.32		99	99		
10200002	**Low discrimination**							**10200002**
					COC	GRA		
Porphyromonas asaccharolytica		53.95	0.42	SAL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		13.22	0.32	BGL 4	1	1		
Peptoniphilus asaccharolyticus		13.22	0.32		99	99		
10200020	**Low discrimination**							**10200020**
					COC	GRA	SPO	
Clostridium cadaveris		23.52	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		17.59	0.32	SAL 1	1	1	1	
Peptoniphilus asaccharolyticus		17.59	0.32		99	99	1	
10201000	**Good identification**							**10201000**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.53	0.66	SAL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1		
Peptoniphilus asaccharolyticus		2.04	0.32		99	99		
10210000	**Low discrimination**							**10210000**
					COC	GRA		
Fusobacterium varium		52.58	0.50	SAL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		11.08	0.32		1	1		
Peptoniphilus asaccharolyticus		11.08	0.32		99	99		
10210020	**Acceptable identification**							**10210020**
					GRA	SPO		
Clostridium cadaveris		92.11	0.44	GLU 95	99	99		
Fusobacterium varium		3.30	0.16	SAL 1	1	1		
Atopobium minutum		0.79	0.19		99	1		
10300000	**Low discrimination**							**10300000**
					COC	GRA	SPO	
Clostridium bifermentans		59.77	0.58	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		9.36	0.32	SAL 10	1	1	1	
Peptoniphilus asaccharolyticus		9.36	0.32		99	99	1	
10300001	**Acceptable identification**							**10300001**
					COC	SPO		
Clostridium sordellii		93.33	0.44	GLU 95	1	99		
Clostridium bifermentans		3.32	0.24	SAL 1	1	99		
Anaerococcus prevotii		0.67	0.06		99	1		
10300020	**Low discrimination**							**10300020**
					GRA	SPO		
Clostridium bifermentans		78.33	0.37	GLU 92	99	99		
Clostridium cadaveris		13.53	0.22	SAL 10	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.94	-.03	NAG 6	1	1		
10310000	**Low discrimination**							**10310000**
					GRA	SPO		
Clostridium bifermentans		54.07	0.36	GLU 92	99	99		
Prevotella intermedia		28.45	0.34	SAL 10	1	1		
Fusobacterium varium		7.71	0.16	FRU 5	1	1		
10310001	**Good identification**							**10310001**
Clostridium sordellii		99.81	0.44	GLU 95				
Clostridium bifermentans		0.19	0.02	SAL 1				
10400000	**Low discrimination**							**10400000**
					COC	GRA		
Fusobacterium nucleatum subsp. nucleatum		24.18	0.66	RAF 1	1	1		
Peptoniphilus asaccharolyticus		24.18	0.66		99	99		
Porphyromonas asaccharolytica		23.93	0.66		1	1		
10400001	**Low discrimination**							**10400001**
					COC	GRA		
Anaerococcus prevotii		89.49	0.67	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		2.10	0.32		1	1		
Peptoniphilus asaccharolyticus		2.10	0.32		99	99		
10400002	**Low discrimination**							**10400002**
					COC	GRA		
Porphyromonas asaccharolytica		46.99	0.42	RAF 1	1	1		
Anaerococcus prevotii		13.62	0.40	BGL 4	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium nucleatum subsp. nucleatum		11.51	0.32		1	1		
10400003	**Very good identification**							**10400003**
					COC	GRA		
Anaerococcus prevotii		99.14	0.56	IND 9	99	99		
Porphyromonas asaccharolytica		0.38	0.08		1	1		
Peptococcus niger		0.12	0.02		99	99		
10400020	**Low discrimination**							**10400020**
					COC	GRA	SPO	
Clostridium cadaveris		23.99	0.44	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		17.94	0.32	RAF 1	1	1	1	
Peptoniphilus asaccharolyticus		17.94	0.32		99	99	1	
10401000	**Good identification**							**10401000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.48	0.66	RAF 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1		
Peptoniphilus asaccharolyticus		2.04	0.32		99	99		
10410000	**Low discrimination**							**10410000**
					COC	GRA		
Fusobacterium varium		51.54	0.50	RAF 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		10.86	0.32		1	1		
Peptoniphilus asaccharolyticus		10.86	0.32		99	99		
10410020	**Acceptable identification**							**10410020**
					GRA	SPO		
Clostridium cadaveris		92.53	0.44	GLU 95	99	99		
Fusobacterium varium		3.32	0.16	RAF 1	1	1		
Atopobium minutum		0.79	0.19		99	1		
10500001	**Low discrimination**							**10500001**
					COC	GRA	SPO	
Clostridium sordellii		78.82	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		18.69	0.32	RAF 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.44	-.03		1	1	1	
10510000	**Very good identification**							**10510000**
					GRA	SPO		
Prevotella intermedia		99.47	0.71	GLU 90	1	1		
Fusobacterium varium		0.18	0.16		1	1		
Clostridium bifermentans		0.12	0.18		99	99		
10510001	**Low discrimination**							**10510001**
					COC	GRA	SPO	
Clostridium sordellii		87.07	0.44	GLU 95	1	99	99	
Prevotella intermedia		12.73	0.37	RAF 1	1	1	1	
Anaerococcus prevotii		0.21	-.02		99	99	1	
10510002	**Good identification**							**10510002**
Prevotella intermedia		100.0	0.37	GLU 90				
10510010	**Good identification**							**10510010**
Prevotella intermedia		100.0	0.37	GLU 90				
10510020	**Low discrimination**							**10510020**
					GRA	SPO		
Prevotella intermedia		78.58	0.37	GLU 90	1	1		
Clostridium cadaveris		20.85	0.22	NAG 1	99	99		
Clostridium bifermentans		0.58	-.02		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
10510040	**Good identification**						**10510040**
Prevotella intermedia		100.0	0.37	GLU 90			
10510100	**Good identification**						**10510100**
Prevotella intermedia		100.0	0.37	GLU 90			
10510200	**Good identification**						**10510200**
Prevotella intermedia		100.0	0.37	GLU 90			
10510400	**Good identification**						**10510400**
Prevotella intermedia		100.0	0.37	GLU 90			
10511000	**Very good identification**						**10511000**
					GRA		
Prevotella intermedia		99.96	0.71	GLU 90	1		
Fusobacterium necrophorum subsp. necrophorum		0.02	-.03		1		
Lactobacillus fermentum		0.01	0.14		99		
10511002	**Good identification**						**10511002**
					GRA		
Prevotella intermedia		99.15	0.37	GLU 90	1		
Lactobacillus fermentum		0.85	0.14	BGL 1	99		
10511010	**Good identification**						**10511010**
Prevotella intermedia		100.0	0.37	GLU 90			
10511020	**Good identification**						**10511020**
Prevotella intermedia		100.0	0.37	GLU 90			
10511040	**Good identification**						**10511040**
Prevotella intermedia		100.0	0.37	GLU 90			
10511100	**Good identification**						**10511100**
					COC	GRA	
Prevotella intermedia		97.47	0.37	GLU 90	1	1	
Lactobacillus fermentum		1.36	0.18	LAC 1	1	99	
Blautia hansenii		1.17	-.03		99	99	
10511200	**Good identification**						**10511200**
Prevotella intermedia		100.0	0.37	GLU 90			
10511400	**Good identification**						**10511400**
					GRA		
Prevotella intermedia		99.15	0.37	GLU 90	1		
Lactobacillus fermentum		0.85	0.14	ARA 1	99		
10512000	**Good identification**						**10512000**
Prevotella intermedia		100.0	0.37	GLU 90			
10513000	**Good identification**						**10513000**
Prevotella intermedia		100.0	0.37	GLU 90			
10514000	**Good identification**						**10514000**
Prevotella intermedia		99.25	0.37	GLU 90			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium varium		0.75	-.08	XYL 1			
10515000	**Good identification**						**10515000**
Prevotella intermedia		100.0	0.37	GLU 90			
10530000	**Good identification**						**10530000**
Prevotella intermedia		100.0	0.37	GLU 90			
10531000	**Good identification**						**10531000**
Prevotella intermedia		100.0	0.37	GLU 90			
10550000	**Good identification**						**10550000**
Prevotella intermedia		100.0	0.37	GLU 90			
10551000	**Good identification**						**10551000**
Prevotella intermedia		100.0	0.37	GLU 90			
10710000	**Good identification**						**10710000**
					GRA	SPO	
Prevotella intermedia		98.74	0.37	GLU 90	1	1	
Clostridium bifermentans		1.26	0.02	SAL 1	99	99	
10711000	**Good identification**						**10711000**
Prevotella intermedia		100.0	0.37	GLU 90			
11000000	**Low discrimination**						**11000000**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		45.18	0.85	GLU 12	1	1	
Clostridium bifermentans		20.90	0.86		99	99	
Fusobacterium nucleatum subsp. nucleatum		7.30	0.66		1	1	
11000001	**Very good identification**						**11000001**
					GRA	SPO	
Clostridium sordellii		95.18	0.84	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		1.55	0.51		1	1	
Clostridium novyi biovar A		0.91	0.66		99	99	
11000002	**Low discrimination**						**11000002**
					GRA		
Fusobacterium necrophorum subsp. necrophorum		27.31	0.51	GLU 12	1		
Atopobium minutum		18.49	0.63	BGL 1	99		
Porphyromonas asaccharolytica		18.03	0.42		1		
11000003	**Low discrimination**						**11000003**
					COC	SPO	
Clostridium sordellii		76.26	0.50	MLT 90	1	99	
Clostridium novyi biovar A		12.67	0.53	BGL 1	1	99	
Anaerococcus prevotii		6.42	0.30		99	1	
11000010	**Low discrimination**						**11000010**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		35.83	0.51	GLU 12	1	1	
Clostridium novyi biovar A		20.90	0.66	MLZ 1	99	99	
Clostridium bifermentans		16.58	0.52		99	99	
11000011	**Low discrimination**						**11000011**
					GRA	SPO	
Clostridium sordellii		50.46	0.50	MLT 90	99	99	
Clostridium novyi biovar A		47.53	0.66	MLZ 1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium necrophorum subsp. necrophorum		0.82	0.17		1	1	
11000012	**Low discrimination**						**11000012**
					GRA	SPO	
Clostridium novyi biovar A		74.81	0.53	IND 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		7.34	0.17		1	1	
Atopobium minutum		4.97	0.29		99	1	
11000013	**Acceptable identification**						**11000013**
					COC	SPO	
Clostridium novyi biovar A		93.82	0.53	IND 1	1	99	
Clostridium sordellii		5.70	0.15		1	99	
Anaerococcus prevotii		0.48	-.04		99	1	
11000020	**Very good identification**						**11000020**
					GRA	SPO	
Clostridium cadaveris		98.74	1.00		99	99	
Clostridium bifermentans		0.71	0.66		99	99	
Fusobacterium necrophorum subsp. necrophorum		0.24	0.51		1	1	
11000021	**Low discrimination**						**11000021**
Clostridium cadaveris		83.36	0.66	URE 1			
Clostridium sordellii		12.57	0.50				
Clostridium novyi biovar A		2.96	0.56				
11000022	**Good identification**						**11000022**
					GRA	SPO	
Clostridium cadaveris		91.60	0.66	BGL 1	99	99	
Atopobium minutum		5.03	0.55		99	1	
Pseudoflavonifractor capillosus		1.54	0.32		1	1	
11000030	**Very good identification**						**11000030**
Clostridium cadaveris		95.44	0.66	MLZ 1			
Clostridium novyi biovar A		3.39	0.56				
Clostridium bifermentans		0.69	0.31				
11000031	**Low discrimination**						**11000031**
Clostridium novyi biovar A		75.22	0.56	IND 1			
Clostridium cadaveris		21.40	0.32				
Clostridium sordellii		3.23	0.15				
11000033	**Good identification**						**11000033**
Clostridium novyi biovar A		98.40	0.43	IND 1			
Clostridium cadaveris		1.60	-.03				
11000040	**Low discrimination**						**11000040**
					GRA	SPO	
Clostridium bifermentans		41.82	0.62	SOR 4	99	99	
Fusobacterium necrophorum subsp. necrophorum		21.91	0.51		1	1	
Clostridium botulinum biovar A		10.05	0.52		99	99	
11000041	**Low discrimination**						**11000041**
Clostridium sordellii		85.51	0.60	MLT 90			
Clostridium novyi biovar A		13.02	0.63	SOR 4			
Clostridium bifermentans		0.65	0.28				
11000042	**Low discrimination**						**11000042**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		53.95	0.50	IND 1	99	99	
Clostridium bifermentans		15.15	0.28		99	99	
Fusobacterium necrophorum subsp. necrophorum		7.94	0.17		1	1	
11000043	**Low discrimination**						**11000043**
					COC	SPO	
Clostridium novyi biovar A		72.27	0.50	IND 1	1	99	
Clostridium sordellii		27.18	0.26		1	99	
Anaerococcus prevotii		0.56	-.04		99	1	
11000050	**Good identification**						**11000050**
					GRA	SPO	
Clostridium novyi biovar A		90.51	0.63	IND 1	99	99	
Clostridium bifermentans		4.49	0.28		99	99	
Fusobacterium necrophorum subsp. necrophorum		2.35	0.17		1	1	
11000051	**Good identification**						**11000051**
Clostridium novyi biovar A		93.78	0.63	IND 1			
Clostridium sordellii		6.22	0.26				
11000052	**Good identification**						**11000052**
Clostridium novyi biovar A		100.0	0.50	IND 1			
11000053	**Good identification**						**11000053**
Clostridium novyi biovar A		100.0	0.50	IND 1			
11000060	**Good identification**						**11000060**
Clostridium cadaveris		93.93	0.66	SOR 1			
Clostridium bifermentans		2.79	0.42				
Clostridium novyi biovar A		2.22	0.53				
11000061	**Low discrimination**						**11000061**
Clostridium novyi biovar A		58.66	0.53	IND 1			
Clostridium cadaveris		25.03	0.32				
Clostridium sordellii		15.57	0.26				
11000063	**Good identification**						**11000063**
Clostridium novyi biovar A		97.62	0.40	IND 1			
Clostridium cadaveris		2.38	-.03				
11000070	**Low discrimination**						**11000070**
Clostridium novyi biovar A		69.47	0.53	IND 1			
Clostridium cadaveris		29.65	0.32				
Clostridium bifermentans		0.88	0.08				
11000071	**Good identification**						**11000071**
Clostridium novyi biovar A		99.57	0.53	IND 1			
Clostridium cadaveris		0.43	-.03				
11000072	**Good identification**						**11000072**
Clostridium novyi biovar A		97.62	0.40	IND 1			
Clostridium cadaveris		2.38	-.03				
11000073	**Good identification**						**11000073**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.40	IND 1			
11000100	**Low discrimination**						**11000100**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		45.15	0.51	GLU 12	1	1	
Clostridium bifermentans		20.89	0.52	LAC 1	99	99	
Fusobacterium nucleatum subsp. nucleatum		7.30	0.32		1	1	
11000101	**Good identification**						**11000101**
					GRA	SPO	
Clostridium sordellii		95.29	0.50	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		1.55	0.17	LAC 1	1	1	
Clostridium novyi biovar A		0.91	0.32		99	99	
11000120	**Very good identification**						**11000120**
					GRA	SPO	
Clostridium cadaveris		98.73	0.66	LAC 1	99	99	
Clostridium bifermentans		0.71	0.31		99	99	
Fusobacterium necrophorum subsp. necrophorum		0.24	0.17		1	1	
11000200	**Low discrimination**						**11000200**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		45.19	0.51	GLU 12	1	1	
Clostridium bifermentans		20.91	0.52	RHA 1	99	99	
Fusobacterium nucleatum subsp. nucleatum		7.31	0.32		1	1	
11000201	**Good identification**						**11000201**
					GRA	SPO	
Clostridium sordellii		95.29	0.50	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		1.55	0.17	RHA 1	1	1	
Clostridium novyi biovar A		0.91	0.32		99	99	
11000220	**Very good identification**						**11000220**
					GRA	SPO	
Clostridium cadaveris		98.76	0.66	RHA 1	99	99	
Clostridium bifermentans		0.71	0.31		99	99	
Fusobacterium necrophorum subsp. necrophorum		0.24	0.17		1	1	
11000400	**Low discrimination**						**11000400**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		45.18	0.51	GLU 12	1	1	
Clostridium bifermentans		20.90	0.52	ARA 1	99	99	
Fusobacterium nucleatum subsp. nucleatum		7.31	0.32		1	1	
11000401	**Good identification**						**11000401**
					GRA	SPO	
Clostridium sordellii		95.29	0.50	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		1.55	0.17	ARA 1	1	1	
Clostridium novyi biovar A		0.91	0.32		99	99	
11000420	**Very good identification**						**11000420**
					GRA	SPO	
Clostridium cadaveris		98.76	0.66	ARA 1	99	99	
Clostridium bifermentans		0.71	0.31		99	99	
Fusobacterium necrophorum subsp. necrophorum		0.24	0.17		1	1	
11001000	**Very good identification**						**11001000**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		98.30	0.85	GLU 12	1	1	
Clostridium bifermentans		0.46	0.52		99	99	
Atopobium minutum		0.34	0.58		99	1	
11001001	**Good identification**						**11001001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium sordellii		90.91	0.71	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.40	0.51		1	1	
Clostridium novyi biovar A		0.55	0.50		99	99	
11001002	**Low discrimination**						**11001002**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		79.62	0.51	GLU 12	1	1	
Atopobium minutum		17.96	0.55	BGL 1	99	1	
Clostridium novyi biovar A		0.91	0.37		99	99	
11001003	**Low discrimination**						**11001003**
					GRA	SPO	
Clostridium sordellii		81.79	0.37	MLT 90	99	99	
Clostridium novyi biovar A		8.56	0.37	BGL 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		7.56	0.17		1	1	
11001010	**Acceptable identification**						**11001010**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		92.52	0.51	GLU 12	1	1	
Clostridium novyi biovar A		6.00	0.50	MLZ 1	99	99	
Clostridium bifermentans		0.43	0.18		99	99	
11001011	**Low discrimination**						**11001011**
					GRA	SPO	
Clostridium sordellii		59.33	0.37	MLT 90	99	99	
Clostridium novyi biovar A		35.19	0.50	MLZ 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		5.48	0.17		1	1	
11001012	**Low discrimination**						**11001012**
					GRA	SPO	
Clostridium novyi biovar A		48.03	0.37	IND 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		42.40	0.17	GAL 10	1	1	
Atopobium minutum		9.57	0.21		99	1	
11001013	**Acceptable identification**						**11001013**
Clostridium novyi biovar A		91.20	0.37	IND 1			
Clostridium sordellii		8.80	0.03	GAL 10			
11001020	**Low discrimination**						**11001020**
					GRA	SPO	
Clostridium cadaveris		76.65	0.66	GAL 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		18.66	0.51		1	1	
Atopobium minutum		2.11	0.50		99	1	
11001021	**Low discrimination**						**11001021**
Clostridium sordellii		60.93	0.37	MLT 90			
Clostridium cadaveris		23.13	0.32	NAG 1			
Clostridium novyi biovar A		9.03	0.39				
11001022	**Low discrimination**						**11001022**
					GRA	SPO	
Atopobium minutum		54.31	0.47	IND 1	99	1	
Clostridium cadaveris		29.97	0.32		99	99	
Fusobacterium necrophorum subsp. necrophorum		7.29	0.17		1	1	
11001031	**Acceptable identification**						**11001031**
Clostridium novyi biovar A		91.41	0.39	IND 1			
Clostridium sordellii		6.23	0.03	GAL 10			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium cadaveris		2.36	-.03				
11001040	**Acceptable identification**						**11001040**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		92.26	0.51	GLU 12	1	1	
Clostridium novyi biovar A		3.99	0.47	SOR 1	99	99	
Clostridium bifermentans		1.78	0.28		99	99	
11001041	**Low discrimination**						**11001041**
					GRA	SPO	
Clostridium sordellii		89.43	0.47	MLT 90	99	99	
Clostridium novyi biovar A		8.57	0.47	SOR 4	99	99	
Fusobacterium necrophorum subsp. necrophorum		2.00	0.17		1	1	
11001050	**Low discrimination**						**11001050**
					GRA	SPO	
Clostridium novyi biovar A		81.06	0.47	IND 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		18.94	0.17	GAL 10	1	1	
11001051	**Acceptable identification**						**11001051**
Clostridium novyi biovar A		90.47	0.47	IND 1			
Clostridium sordellii		9.54	0.13	GAL 10			
11001061	**Low discrimination**						**11001061**
Clostridium novyi biovar A		68.48	0.36	IND 1			
Clostridium sordellii		28.87	0.13	GAL 10			
Clostridium cadaveris		2.66	-.03				
11001070	**Good identification**						**11001070**
Clostridium novyi biovar A		96.27	0.36	IND 1			
Clostridium cadaveris		3.73	-.03	GAL 10			
11001071	**Good identification**						**11001071**
Clostridium novyi biovar A		100.0	0.36	IND 1			
11001100	**Good identification**						**11001100**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		97.97	0.51	GLU 12	1	1	
Clostridium bifermentans		0.46	0.18	LAC 1	99	99	
Atopobium minutum		0.34	0.24		99	1	
11001101	**Acceptable identification**						**11001101**
					GRA	SPO	
Clostridium sordellii		91.04	0.37	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	LAC 1	1	1	
Clostridium novyi biovar A		0.55	0.15		99	99	
11001200	**Good identification**						**11001200**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		98.37	0.51	GLU 12	1	1	
Clostridium bifermentans		0.46	0.18	RHA 1	99	99	
Atopobium minutum		0.34	0.24		99	1	
11001201	**Acceptable identification**						**11001201**
					GRA	SPO	
Clostridium sordellii		91.04	0.37	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	RHA 1	1	1	
Clostridium novyi biovar A		0.55	0.15		99	99	
11001400	**Good identification**						**11001400**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		98.37	0.51	GLU 12	1	1		
Clostridium bifermentans		0.46	0.18	ARA 1	99	99		
Atopobium minutum		0.34	0.24		99	1		
11001401	**Acceptable identification**							**11001401**
					GRA	SPO		
Clostridium sordellii		91.04	0.37	MLT 90	99	99		
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	ARA 1	1	1		
Clostridium novyi biovar A		0.55	0.15		99	99		
11002000	**Low discrimination**							**11002000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		43.26	0.51	GLU 12	1	1		
Clostridium bifermentans		20.02	0.52	MAN 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		6.99	0.32		1	1		
11002001	**Good identification**							**11002001**
					GRA	SPO		
Clostridium sordellii		95.18	0.50	MLT 90	99	99		
Fusobacterium necrophorum subsp. necrophorum		1.55	0.17	MAN 1	1	1		
Clostridium novyi biovar A		0.91	0.32		99	99		
11002020	**Very good identification**							**11002020**
					SPO	CAT		
Clostridium cadaveris		97.49	0.66	MAN 1	99	1		
Propionibacterium acnes		1.14	0.36		1	95		
Clostridium bifermentans		0.70	0.31		99	1		
11003000	**Good identification**							**11003000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		98.25	0.51	GLU 12	1	1		
Clostridium bifermentans		0.46	0.18	MAN 1	99	99		
Atopobium minutum		0.34	0.24		99	1		
11003001	**Acceptable identification**							**11003001**
					GRA	SPO		
Clostridium sordellii		91.04	0.37	MLT 90	99	99		
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	MAN 1	1	1		
Clostridium novyi biovar A		0.55	0.15		99	99		
11003020	**Low discrimination**							**11003020**
					GRA	SPO	CAT	
Propionibacterium acnes		47.57	0.36	NIT 92	99	1	95	
Clostridium cadaveris		41.02	0.32	FRU 99	99	99	1	
Fusobacterium necrophorum subsp. necrophorum		9.99	0.17		1	1	1	
11004000	**Low discrimination**							**11004000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		43.23	0.51	GLU 12	1	1		
Clostridium bifermentans		20.00	0.52	XYL 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		6.99	0.32		1	1		
11004001	**Good identification**							**11004001**
					GRA	SPO		
Clostridium sordellii		95.13	0.50	MLT 90	99	99		
Fusobacterium necrophorum subsp. necrophorum		1.55	0.17	XYL 1	1	1		
Clostridium novyi biovar A		0.91	0.32		99	99		
11004020	**Very good identification**							**11004020**
					GRA	SPO		
Clostridium cadaveris		98.73	0.66	XYL 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		0.71	0.31		99	99	
Fusobacterium necrophorum subsp. necrophorum		0.24	0.17		1	1	
11004040	**Low discrimination**						**11004040**
					GRA	SPO	
Clostridium glycolicum		88.08	0.49	IND 1	99	99	
Clostridium bifermentans		5.02	0.28	FRU 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		2.63	0.17		1	1	
11005000	**Good identification**						**11005000**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		97.93	0.51	GLU 12	1	1	
Clostridium bifermentans		0.46	0.18	XYL 1	99	99	
Atopobium minutum		0.34	0.24		99	1	
11005001	**Acceptable identification**						**11005001**
					GRA	SPO	
Clostridium sordellii		91.04	0.37	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	XYL 1	1	1	
Clostridium novyi biovar A		0.55	0.15		99	99	
11010000	**Good identification**						**11010000**
					GRA	SPO	
Fusobacterium varium		93.31	0.93		1	1	
Pseudoramibacter alactolyticus		2.48	0.66		99	1	
Clostridium cadaveris		1.51	0.66		99	99	
11010001	**Very good identification**						**11010001**
					GRA	SPO	
Clostridium sordellii		95.61	0.84	MLT 90	99	99	
Fusobacterium varium		4.01	0.59		1	1	
Pseudoramibacter alactolyticus		0.11	0.32		99	1	
11010002	**Low discrimination**						**11010002**
					GRA		
Fusobacterium varium		61.75	0.59	BGL 1	1		
Pseudoramibacter alactolyticus		18.06	0.50		99		
Atopobium minutum		15.02	0.62		99		
11010003	**Acceptable identification**						**11010003**
					GRA	SPO	
Clostridium sordellii		92.11	0.50	MLT 90	99	99	
Fusobacterium varium		3.86	0.25	BGL 1	1	1	
Clostridium novyi biovar A		1.70	0.37		99	99	
11010010	**Low discrimination**						**11010010**
					GRA	SPO	
Fusobacterium varium		88.73	0.59	MLZ 1	1	1	
Clostridium difficile		2.71	0.36		99	99	
Pseudoramibacter alactolyticus		2.36	0.32		99	1	
11010011	**Low discrimination**						**11010011**
					GRA	SPO	
Clostridium sordellii		86.99	0.50	MLT 90	99	99	
Clostridium novyi biovar A		9.10	0.50	MLZ 1	99	99	
Fusobacterium varium		3.64	0.25		1	1	
11010013	**Low discrimination**						**11010013**
Clostridium novyi biovar A		64.64	0.37	IND 1			
Clostridium sordellii		35.36	0.15	FRU 10			
11010020	**Very good identification**						**11010020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Clostridium cadaveris		96.96	1.00		99	99	1	
Propionibacterium acnes		2.02	0.74		99	1	95	
Fusobacterium varium		0.61	0.59		1	1	1	
11010021	**Low discrimination**							**11010021**
					SPO	CAT		
Clostridium cadaveris		84.32	0.66	URE 1	99	1		
Clostridium sordellii		12.72	0.50		99	1		
Propionibacterium acnes		1.76	0.40		1	95		
11010022	**Low discrimination**							**11010022**
					SPO			
Clostridium cadaveris		89.83	0.66	BGL 1	99			
Atopobium minutum		4.55	0.54		1			
Pseudoramibacter alactolyticus		2.90	0.37		1			
11010030	**Very good identification**							**11010030**
					GRA	SPO	CAT	
Clostridium cadaveris		96.59	0.66	MLZ 1	99	99	1	
Propionibacterium acnes		2.01	0.40		99	1	95	
Fusobacterium varium		0.61	0.25		1	1	1	
11010040	**Low discrimination**							**11010040**
					COC	GRA		
Fusobacterium varium		49.52	0.59	SOR 1	1	1		
Pseudoramibacter alactolyticus		32.59	0.56		1	99		
Megasphaera elsdenii		9.15	0.57		99	1		
11010041	**Very good identification**							**11010041**
					GRA	SPO		
Clostridium sordellii		96.44	0.60	MLT 90	99	99		
Clostridium novyi biovar A		1.63	0.47	SOR 4	99	99		
Fusobacterium varium		0.98	0.25		1	1		
11010042	**Low discrimination**							**11010042**
					COC	GRA		
Pseudoramibacter alactolyticus		66.19	0.39	IND 1	1	99		
Megasphaera elsdenii		18.58	0.41	BGL 10	99	1		
Fusobacterium varium		9.14	0.25		1	1		
11010050	**Low discrimination**							**11010050**
					GRA	SPO		
Clostridium novyi biovar A		43.87	0.47	IND 1	99	99		
Fusobacterium varium		26.34	0.25	FRU 10	1	1		
Pseudoramibacter alactolyticus		17.34	0.21		99	1		
11010051	**Low discrimination**							**11010051**
Clostridium novyi biovar A		62.61	0.47	IND 1				
Clostridium sordellii		37.39	0.26	FRU 10				
11010060	**Low discrimination**							**11010060**
					SPO	CAT		
Clostridium cadaveris		65.02	0.66	SOR 1	99	1		
Propionibacterium acnes		29.42	0.63		1	95		
Pseudoramibacter alactolyticus		4.72	0.43		1	1		
11010071	**Good identification**							**11010071**
Clostridium novyi biovar A		96.27	0.36	IND 1				
Clostridium cadaveris		3.73	-.03	FRU 10				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
11010100	**Good identification**							**11010100**
					GRA	SPO		
Fusobacterium varium		91.85	0.59	LAC 1	1	1		
Pseudoramibacter alactolyticus		2.44	0.32		99	1		
Clostridium cadaveris		1.48	0.32		99	99		
11010101	**Good identification**							**11010101**
					GRA	SPO		
Clostridium sordellii		95.66	0.50	MLT 90	99	99		
Fusobacterium varium		4.01	0.25	LAC 1	1	1		
Pseudoramibacter alactolyticus		0.11	-.03		99	1		
11010120	**Very good identification**							**11010120**
					GRA	SPO	CAT	
Clostridium cadaveris		96.85	0.66	LAC 1	99	99	1	
Propionibacterium acnes		2.02	0.40		99	1	95	
Fusobacterium varium		0.61	0.25		1	1	1	
11010200	**Good identification**							**11010200**
					GRA	SPO		
Fusobacterium varium		93.33	0.59	RHA 1	1	1		
Pseudoramibacter alactolyticus		2.48	0.32		99	1		
Clostridium cadaveris		1.51	0.32		99	99		
11010201	**Good identification**							**11010201**
					GRA	SPO		
Clostridium sordellii		95.72	0.50	MLT 90	99	99		
Fusobacterium varium		4.01	0.25	RHA 1	1	1		
Pseudoramibacter alactolyticus		0.11	-.03		99	1		
11010220	**Very good identification**							**11010220**
					GRA	SPO	CAT	
Clostridium cadaveris		96.97	0.66	RHA 1	99	99	1	
Propionibacterium acnes		2.02	0.40		99	1	95	
Fusobacterium varium		0.61	0.25		1	1	1	
11010400	**Good identification**							**11010400**
					GRA	SPO		
Fusobacterium varium		93.23	0.59	ARA 1	1	1		
Pseudoramibacter alactolyticus		2.48	0.32		99	1		
Clostridium cadaveris		1.51	0.32		99	99		
11010401	**Good identification**							**11010401**
					GRA	SPO		
Clostridium sordellii		95.72	0.50	MLT 90	99	99		
Fusobacterium varium		4.01	0.25	ARA 1	1	1		
Pseudoramibacter alactolyticus		0.11	-.03		99	1		
11010420	**Very good identification**							**11010420**
					GRA	SPO	CAT	
Clostridium cadaveris		96.97	0.66	ARA 1	99	99	1	
Propionibacterium acnes		2.02	0.40		99	1	95	
Fusobacterium varium		0.61	0.25		1	1	1	
11011000	**Low discrimination**							**11011000**
					GRA			
Fusobacterium varium		56.80	0.59	GAL 1	1			
Fusobacterium necrophorum subsp. necrophorum		22.11	0.51		1			
Atopobium minutum		6.91	0.57		99			
11011001	**Very good identification**							**11011001**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sordellii		99.53	0.71	MLT 90	99	99	
Fusobacterium varium		0.24	0.25		1	1	
Fusobacterium necrophorum subsp. necrophorum		0.09	0.17		1	1	
11011002	**Low discrimination**						**11011002**
					GRA		
Atopobium minutum		77.93	0.54	IND 1	99		
Fusobacterium varium		9.71	0.25		1		
Fusobacterium necrophorum subsp. necrophorum		3.78	0.17		1		
11011003	**Good identification**						**11011003**
					SPO		
Clostridium sordellii		97.01	0.37	MLT 90	99		
Atopobium minutum		1.87	0.20	BGL 1	1		
Clostridium novyi biovar A		1.13	0.20		99		
11011011	**Acceptable identification**						**11011011**
Clostridium sordellii		93.82	0.37	MLT 90			
Clostridium novyi biovar A		6.18	0.33	MLZ 1			
11011020	**Low discrimination**						**11011020**
					SPO	CAT	
Propionibacterium acnes		66.47	0.74	NIT 92	1	95	
Clostridium cadaveris		32.24	0.66		99	1	
Atopobium minutum		0.82	0.49		1	1	
11011021	**Low discrimination**						**11011021**
					SPO	CAT	
Clostridium sordellii		45.70	0.37	MLT 90	99	1	
Propionibacterium acnes		35.76	0.40	NAG 1	1	95	
Clostridium cadaveris		17.35	0.32		99	1	
11011022	**Low discrimination**						**11011022**
					SPO	CAT	
Propionibacterium acnes		42.47	0.40	NIT 92	1	95	
Atopobium minutum		34.46	0.46	BGL 1	1	1	
Clostridium cadaveris		20.60	0.32		99	1	
11011030	**Low discrimination**						**11011030**
					SPO	CAT	
Propionibacterium acnes		65.86	0.40	NIT 92	1	95	
Clostridium cadaveris		31.95	0.32	MLZ 1	99	1	
Clostridium novyi biovar A		1.39	0.23		99	1	
11011041	**Good identification**						**11011041**
Clostridium sordellii		98.95	0.47	MLT 90			
Clostridium novyi biovar A		1.05	0.30	SOR 4			
11011060	**Very good identification**						**11011060**
					SPO	CAT	
Propionibacterium acnes		97.55	0.63	NIT 92	1	95	
Clostridium cadaveris		2.18	0.32		99	1	
Pseudoramibacter alactolyticus		0.16	0.09		1	1	
11011100	**Low discrimination**						**11011100**
					GRA		
Actinomyces meyeri		46.84	0.41	IND 1	99		
Fusobacterium varium		20.78	0.25	SUC 93	1		
Lactobacillus fermentum		9.71	0.40		99		
11011101	**Good identification**						**11011101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium sordellii		99.46	0.37	MLT 90	99		
Actinomyces meyeri		0.54	0.07	LAC 1	1		
11011102	**Low discrimination**						**11011102**
Lactobacillus fermentum		71.71	0.40	IND 1			
Atopobium minutum		12.44	0.20	SUC 87			
Actinomyces meyeri		10.70	0.15				
11011120	**Low discrimination**						**11011120**
					SPO	CAT	
Propionibacterium acnes		60.71	0.40	NIT 92	1	95	
Clostridium cadaveris		29.45	0.32	LAC 1	99	1	
Actinomyces meyeri		5.66	0.26		1	1	
11011201	**Good identification**						**11011201**
Clostridium sordellii		100.0	0.37	MLT 90			
11011220	**Low discrimination**						**11011220**
					SPO	CAT	
Propionibacterium acnes		66.79	0.40	NIT 92	1	95	
Clostridium cadaveris		32.39	0.32	RHA 1	99	1	
Atopobium minutum		0.82	0.15		1	1	
11011401	**Good identification**						**11011401**
Clostridium sordellii		100.0	0.37	MLT 90			
11011402	**Low discrimination**						**11011402**
Lactobacillus fermentum		77.95	0.36	IND 1			
Atopobium minutum		22.05	0.20	SUC 87			
11011420	**Low discrimination**						**11011420**
					SPO	CAT	
Propionibacterium acnes		64.87	0.40	NIT 92	1	95	
Clostridium cadaveris		31.47	0.32	ARA 1	99	1	
Lactobacillus fermentum		1.88	0.28		1	1	
11011502	**Good identification**						**11011502**
Lactobacillus fermentum		96.41	0.40	IND 1			
Actinomyces meyeri		3.60	0.05	SUC 87			
11012000	**Low discrimination**						**11012000**
					GRA		
Pseudoramibacter alactolyticus		38.19	0.61	IND 1	99		
Fusobacterium varium		26.94	0.59		1		
Eubacterium limosum		24.29	0.60		99		
11012001	**Low discrimination**						**11012001**
					GRA	SPO	
Clostridium sordellii		86.54	0.50	MLT 90	99	99	
Pseudoramibacter alactolyticus		5.14	0.27	MAN 1	99	1	
Fusobacterium varium		3.63	0.25		1	1	
11012002	**Low discrimination**						**11012002**
					COC	GRA	
Pseudoramibacter alactolyticus		74.69	0.45	IND 1	1	99	
Megasphaera elsdenii		14.60	0.44	BGL 10	99	1	
Fusobacterium varium		4.79	0.25		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
11012010	**Low discrimination**						**11012010**
					GRA	SPO	
Clostridium difficile		88.15	0.53	IND 1	99	99	
Pseudoramibacter alactolyticus		4.59	0.27		99	1	
Fusobacterium varium		3.24	0.25		1	1	
11012020	**Low discrimination**						**11012020**
					SPO	CAT	
Propionibacterium acnes		49.74	0.70	NIT 92	1	95	
Clostridium cadaveris		42.89	0.66		99	1	
Pseudoramibacter alactolyticus		6.71	0.48		1	1	
11012021	**Low discrimination**						**11012021**
					SPO	CAT	
Propionibacterium acnes		47.01	0.36	NIT 92	1	95	
Clostridium cadaveris		40.54	0.32	URE 1	99	1	
Pseudoramibacter alactolyticus		6.34	0.14		1	1	
11012030	**Low discrimination**						**11012030**
					SPO	CAT	
Propionibacterium acnes		46.61	0.36	NIT 92	1	95	
Clostridium cadaveris		40.19	0.32	MLZ 1	99	1	
Clostridium difficile		6.91	0.19		99	1	
11012040	**Low discrimination**						**11012040**
					COC	GRA	
Pseudoramibacter alactolyticus		62.52	0.51	IND 1	1	99	
Megasphaera elsdenii		32.59	0.57		99	1	
Fusobacterium varium		1.78	0.25		1	1	
11012060	**Low discrimination**						**11012060**
					SPO	CAT	
Propionibacterium acnes		83.48	0.59	NIT 92	1	95	
Pseudoramibacter alactolyticus		12.83	0.38		1	1	
Clostridium cadaveris		3.31	0.32		99	1	
11012120	**Low discrimination**						**11012120**
					SPO	CAT	
Propionibacterium acnes		50.07	0.36	NIT 92	1	95	
Clostridium cadaveris		43.18	0.32	LAC 1	99	1	
Pseudoramibacter alactolyticus		6.76	0.14		1	1	
11012220	**Low discrimination**						**11012220**
					SPO	CAT	
Propionibacterium acnes		50.07	0.36	NIT 92	1	95	
Clostridium cadaveris		43.18	0.32	RHA 1	99	1	
Pseudoramibacter alactolyticus		6.76	0.14		1	1	
11012420	**Low discrimination**						**11012420**
					SPO	CAT	
Propionibacterium acnes		50.07	0.36	NIT 92	1	95	
Clostridium cadaveris		43.18	0.32	ARA 1	99	1	
Pseudoramibacter alactolyticus		6.76	0.14		1	1	
11013000	**Low discrimination**						**11013000**
					GRA	CAT	
Propionibacterium acnes		30.02	0.36	NIT 92	99	95	
Pseudoramibacter alactolyticus		22.95	0.27	NAG 99	99	1	
Fusobacterium varium		16.19	0.25		1	1	
11013001	**Good identification**						**11013001**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sordellii		99.22	0.37	MLT 90	99	1	
Propionibacterium acnes		0.44	0.02	MAN 1	1	95	
Pseudoramibacter alactolyticus		0.34	-.07		1	1	
11013020	**Very good identification**						**11013020**
					SPO	CAT	
Propionibacterium acnes		98.98	0.70	NIT 92	1	95	
Clostridium cadaveris		0.86	0.32		99	1	
Pseudoramibacter alactolyticus		0.14	0.14		1	1	
11013021	**Good identification**						**11013021**
					SPO	CAT	
Propionibacterium acnes		96.93	0.36	NIT 92	1	95	
Clostridium sordellii		2.22	0.03	URE 1	99	1	
Clostridium cadaveris		0.84	-.03		99	1	
11013022	**Good identification**						**11013022**
					CAT		
Propionibacterium acnes		96.31	0.36	NIT 92	95		
Pseudoramibacter alactolyticus		1.44	-.02	BGL 1	1		
Atopobium minutum		1.40	0.12		1		
11013030	**Good identification**						**11013030**
					SPO	CAT	
Propionibacterium acnes		99.14	0.36	NIT 92	1	95	
Clostridium cadaveris		0.86	-.03	MLZ 1	99	1	
11013060	**Very good identification**						**11013060**
					SPO	CAT	
Propionibacterium acnes		99.81	0.59	NIT 92	1	95	
Pseudoramibacter alactolyticus		0.16	0.04		1	1	
Clostridium cadaveris		0.04	-.03		99	1	
11013120	**Good identification**						**11013120**
					SPO	CAT	
Propionibacterium acnes		99.14	0.36	NIT 92	1	95	
Clostridium cadaveris		0.86	-.03	LAC 1	99	1	
11013220	**Good identification**						**11013220**
					SPO	CAT	
Propionibacterium acnes		99.14	0.36	NIT 92	1	95	
Clostridium cadaveris		0.86	-.03	RHA 1	99	1	
11013420	**Good identification**						**11013420**
					SPO	CAT	
Propionibacterium acnes		99.14	0.36	NIT 92	1	95	
Clostridium cadaveris		0.86	-.03	ARA 1	99	1	
11014000	**Very good identification**						**11014000**
					GRA	SPO	
Fusobacterium varium		96.68	0.69	XYL 4	1	1	
Clostridium glycolicum		1.46	0.43		99	99	
Pseudoramibacter alactolyticus		0.62	0.32		99	1	
11014001	**Low discrimination**						**11014001**
					GRA	SPO	
Clostridium sordellii		84.87	0.50	MLT 90	99	99	
Fusobacterium varium		14.67	0.35	XYL 1	1	1	
Clostridium glycolicum		0.22	0.09		99	99	
11014002	**Low discrimination**						**11014002**
					GRA		
Fusobacterium varium		86.07	0.35	XYL 4	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		6.10	0.15	BGL 1	99			
Atopobium minutum		5.07	0.28		99			
11014010	**Acceptable identification**							**11014010**
					GRA	SPO		
Fusobacterium varium		93.44	0.35	XYL 4	1	1		
Clostridium difficile		3.60	0.15	MLZ 1	99	99		
Clostridium glycolicum		1.41	0.09		99	99		
11014020	**Very good identification**							**11014020**
					GRA	SPO	CAT	
Clostridium cadaveris		95.06	0.66	XYL 1	99	99	1	
Fusobacterium varium		2.48	0.35		1	1	1	
Propionibacterium acnes		1.98	0.40		99	1	95	
11014040	**Very good identification**							**11014040**
					GRA	SPO		
Clostridium glycolicum		95.87	0.65	IND 1	99	99		
Fusobacterium varium		3.37	0.35		1	1		
Pseudoramibacter alactolyticus		0.54	0.21		99	1		
11014100	**Low discrimination**							**11014100**
					GRA	SPO		
Fusobacterium varium		57.59	0.35	XYL 4	1	1		
Actinomyces meyeri		40.73	0.43	LAC 1	99	1		
Clostridium glycolicum		0.87	0.09		99	99		
11014200	**Good identification**							**11014200**
					GRA	SPO		
Fusobacterium varium		97.52	0.35	XYL 4	1	1		
Clostridium glycolicum		1.47	0.09	RHA 1	99	99		
Pseudoramibacter alactolyticus		0.63	-.03		99	1		
11014400	**Acceptable identification**							**11014400**
					GRA	SPO		
Fusobacterium varium		93.49	0.35	XYL 4	1	1		
Actinomyces meyeri		4.13	0.22	ARA 1	99	1		
Clostridium glycolicum		1.41	0.09		99	99		
11015000	**Low discrimination**							**11015000**
					GRA	SPO		
Fusobacterium varium		60.52	0.35	GAL 1	1	1		
Actinomyces meyeri		16.05	0.35	XYL 4	99	1		
Clostridium glycolicum		10.06	0.27		99	99		
11015001	**Good identification**							**11015001**
					GRA	SPO		
Clostridium sordellii		99.02	0.37	MLT 90	99	99		
Fusobacterium varium		0.98	0.01	XYL 1	1	1		
11015020	**Low discrimination**							**11015020**
					SPO	CAT		
Propionibacterium acnes		63.60	0.40	NIT 92	1	95		
Clostridium cadaveris		30.85	0.32	XYL 1	99	1		
Actinomyces meyeri		2.88	0.21		1	1		
11015040	**Good identification**							**11015040**
					GRA	SPO		
Clostridium glycolicum		99.68	0.49	IND 1	99	99		
Fusobacterium varium		0.32	0.01	GAL 10	1	1		
11015100	**Acceptable identification**							**11015100**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		92.19	0.46	IND 1	99		
Lactobacillus fermentum		5.78	0.36	SUC 93	99		
Fusobacterium varium		0.88	0.01		1		
11015102	**Low discrimination**						**11015102**
Lactobacillus fermentum		66.98	0.36	IND 1			
Actinomyces meyeri		33.03	0.20	SUC 87			
11015500	**Low discrimination**						**11015500**
Actinomyces meyeri		79.94	0.35	IND 1			
Lactobacillus fermentum		20.06	0.36	SUC 93			
11015502	**Low discrimination**						**11015502**
Lactobacillus fermentum		89.03	0.36	IND 1			
Actinomyces meyeri		10.98	0.10	SUC 87			
11016000	**Low discrimination**						**11016000**
					GRA		
Fusobacterium varium		58.00	0.35	MAN 1	1		
Pseudoramibacter alactolyticus		19.93	0.27	XYL 4	99		
Eubacterium limosum		12.68	0.26		99		
11016020	**Low discrimination**						**11016020**
					SPO	CAT	
Propionibacterium acnes		49.51	0.36	NIT 92	1	95	
Clostridium cadaveris		42.70	0.32	XYL 1	99	1	
Pseudoramibacter alactolyticus		6.68	0.14		1	1	
11017020	**Good identification**						**11017020**
					SPO	CAT	
Propionibacterium acnes		99.14	0.36	NIT 92	1	95	
Clostridium cadaveris		0.86	-.03	XYL 1	99	1	
11020000	**Low discrimination**						**11020000**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		34.77	0.51	GLU 12	1	1	
Clostridium bifermentans		16.09	0.52	TRE 1	99	99	
Clostridium botulinum biovar A		13.35	0.51		99	99	
11020001	**Acceptable identification**						**11020001**
					GRA	SPO	
Clostridium sordellii		94.29	0.50	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		1.54	0.17	TRE 1	1	1	
Clostridium novyi biovar A		0.90	0.32		99	99	
11020020	**Very good identification**						**11020020**
					GRA	SPO	
Clostridium cadaveris		99.74	0.78	TRE 5	99	99	
Clostridium bifermentans		0.14	0.31		99	99	
Fusobacterium necrophorum subsp. necrophorum		0.05	0.17		1	1	
11020021	**Good identification**						**11020021**
Clostridium cadaveris		96.42	0.44	TRE 5			
Clostridium sordellii		2.79	0.15	URE 1			
Clostridium novyi biovar A		0.66	0.21				
11020022	**Good identification**						**11020022**
					GRA	SPO	
Clostridium cadaveris		98.32	0.44	TRE 5	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		1.04	0.21	BGL 1	99	1	
Pseudoflavonifractor capillosus		0.32	-.03		1	1	
11020030	**Good identification**						**11020030**
Clostridium cadaveris		99.19	0.44	TRE 5			
Clostridium novyi biovar A		0.68	0.21	MLZ 1			
Clostridium bifermentans		0.14	-.03				
11020040	**Low discrimination**						**11020040**
					GRA	SPO	
Clostridium botulinum biovar A		72.69	0.42	IND 1	99	99	
Clostridium bifermentans		12.22	0.28		99	99	
Fusobacterium necrophorum subsp. necrophorum		6.40	0.17		1	1	
11020060	**Good identification**						**11020060**
Clostridium cadaveris		98.47	0.44	TRE 5			
Clostridium bifermentans		0.56	0.08	SOR 1			
Clostridium botulinum biovar A		0.53	0.08				
11020120	**Good identification**						**11020120**
Clostridium cadaveris		99.86	0.44	TRE 5			
Clostridium bifermentans		0.14	-.03	LAC 1			
11020220	**Good identification**						**11020220**
Clostridium cadaveris		99.86	0.44	TRE 5			
Clostridium bifermentans		0.14	-.03	RHA 1			
11020420	**Good identification**						**11020420**
Clostridium cadaveris		99.86	0.44	TRE 5			
Clostridium bifermentans		0.14	-.03	ARA 1			
11021000	**Good identification**						**11021000**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		97.70	0.51	GLU 12	1	1	
Clostridium bifermentans		0.46	0.18	TRE 1	99	99	
Clostridium botulinum biovar A		0.38	0.17		99	99	
11021001	**Acceptable identification**						**11021001**
					GRA	SPO	
Clostridium sordellii		91.04	0.37	MLT 90	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	TRE 1	1	1	
Clostridium novyi biovar A		0.55	0.15		99	99	
11021020	**Acceptable identification**						**11021020**
					GRA	SPO	
Clostridium cadaveris		94.58	0.44	TRE 5	99	99	
Fusobacterium necrophorum subsp. necrophorum		4.42	0.17	GAL 1	1	1	
Atopobium minutum		0.50	0.15		99	1	
11022020	**Good identification**						**11022020**
					SPO	CAT	
Clostridium cadaveris		99.64	0.44	TRE 5	99	1	
Propionibacterium acnes		0.22	0.02	MAN 1	1	95	
Clostridium bifermentans		0.14	-.03		99	1	
11024020	**Good identification**						**11024020**
Clostridium cadaveris		99.86	0.44	TRE 5			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		0.14	-.03	XYL 1				
11030000	**Low discrimination**							**11030000**
					GRA	SPO		
Fusobacterium varium		86.37	0.59	TRE 1	1	1		
Clostridium cadaveris		7.27	0.44		99	99		
Pseudoramibacter alactolyticus		2.30	0.32		99	1		
11030001	**Good identification**							**11030001**
					GRA	SPO		
Clostridium sordellii		95.46	0.50	MLT 90	99	99		
Fusobacterium varium		4.00	0.25	TRE 1	1	1		
Clostridium cadaveris		0.34	0.10		99	99		
11030020	**Very good identification**							**11030020**
					GRA	SPO	CAT	
Clostridium cadaveris		99.40	0.78	TRE 5	99	99	1	
Propionibacterium acnes		0.40	0.40		99	1	95	
Fusobacterium varium		0.12	0.25		1	1	1	
11030021	**Good identification**							**11030021**
					SPO	CAT		
Clostridium cadaveris		96.81	0.44	TRE 5	99	1		
Clostridium sordellii		2.80	0.15	URE 1	99	1		
Propionibacterium acnes		0.39	0.06		1	95		
11030022	**Good identification**							**11030022**
					SPO			
Clostridium cadaveris		98.05	0.44	TRE 5	99			
Atopobium minutum		0.95	0.20	BGL 1	1			
Pseudoramibacter alactolyticus		0.61	0.03		1			
11030030	**Good identification**							**11030030**
					SPO	CAT		
Clostridium cadaveris		99.60	0.44	TRE 5	99	1		
Propionibacterium acnes		0.40	0.06	MLZ 1	1	95		
11030060	**Acceptable identification**							**11030060**
					SPO	CAT		
Clostridium cadaveris		90.84	0.44	TRE 5	99	1		
Propionibacterium acnes		7.89	0.29	SOR 1	1	95		
Pseudoramibacter alactolyticus		1.27	0.09		1	1		
11030120	**Good identification**							**11030120**
					SPO	CAT		
Clostridium cadaveris		99.60	0.44	TRE 5	99	1		
Propionibacterium acnes		0.40	0.06	LAC 1	1	95		
11030220	**Good identification**							**11030220**
					SPO	CAT		
Clostridium cadaveris		99.60	0.44	TRE 5	99	1		
Propionibacterium acnes		0.40	0.06	RHA 1	1	95		
11030420	**Good identification**							**11030420**
					SPO	CAT		
Clostridium cadaveris		99.60	0.44	TRE 5	99	1		
Propionibacterium acnes		0.40	0.06	ARA 1	1	95		
11031001	**Good identification**							**11031001**
Clostridium sordellii		100.0	0.37	MLT 90				
11031020	**Low discrimination**							**11031020**
					SPO	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium cadaveris		71.40	0.44	TRE 5	99	1		
Propionibacterium acnes		28.25	0.40	GAL 1	1	95		
Atopobium minutum		0.35	0.15		1	1		
11032020	**Low discrimination**							**11032020**
					SPO	CAT		
Clostridium cadaveris		79.84	0.44	TRE 5	99	1		
Propionibacterium acnes		17.77	0.36	MAN 1	1	95		
Pseudoramibacter alactolyticus		2.40	0.14		1	1		
11033020	**Good identification**							**11033020**
					SPO	CAT		
Propionibacterium acnes		95.66	0.36	NIT 92	1	95		
Clostridium cadaveris		4.34	0.10	TRE 1	99	1		
11034000	**Good identification**							**11034000**
					GRA	SPO		
Fusobacterium varium		95.97	0.35	TRE 1	1	1		
Clostridium cadaveris		1.96	0.10	XYL 4	99	99		
Clostridium glycolicum		1.45	0.09		99	99		
11034020	**Good identification**							**11034020**
					GRA	SPO	CAT	
Clostridium cadaveris		99.11	0.44	TRE 5	99	99	1	
Fusobacterium varium		0.50	0.01	XYL 1	1	1	1	
Propionibacterium acnes		0.40	0.06		99	1	95	
11040000	**Low discrimination**							**11040000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		45.19	0.51	GLU 12	1	1		
Clostridium bifermentans		20.91	0.52	CEL 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		7.31	0.32		1	1		
11040001	**Good identification**							**11040001**
					GRA	SPO		
Clostridium sordellii		95.29	0.50	MLT 90	99	99		
Fusobacterium necrophorum subsp. necrophorum		1.55	0.17	CEL 1	1	1		
Clostridium novyi biovar A		0.91	0.32		99	99		
11040020	**Very good identification**							**11040020**
					GRA	SPO		
Clostridium cadaveris		98.76	0.66	CEL 1	99	99		
Clostridium bifermentans		0.71	0.31		99	99		
Fusobacterium necrophorum subsp. necrophorum		0.24	0.17		1	1		
11041000	**Good identification**							**11041000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		98.37	0.51	GLU 12	1	1		
Clostridium bifermentans		0.46	0.18	CEL 1	99	99		
Atopobium minutum		0.34	0.24		99	1		
11041001	**Acceptable identification**							**11041001**
					GRA	SPO		
Clostridium sordellii		91.04	0.37	MLT 90	99	99		
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	CEL 1	1	1		
Clostridium novyi biovar A		0.55	0.15		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
11050000	**Good identification**							**11050000**
					GRA	SPO		
Fusobacterium varium		93.27	0.59	CEL 1	1	1		
Pseudoramibacter alactolyticus		2.48	0.32		99	1		
Clostridium cadaveris		1.51	0.32		99	99		
11050001	**Good identification**							**11050001**
					GRA	SPO		
Clostridium sordellii		95.72	0.50	MLT 90	99	99		
Fusobacterium varium		4.01	0.25	CEL 1	1	1		
Pseudoramibacter alactolyticus		0.11	-.03		99	1		
11050020	**Very good identification**							**11050020**
					GRA	SPO	CAT	
Clostridium cadaveris		96.97	0.66	CEL 1	99	99	1	
Propionibacterium acnes		2.02	0.40		99	1	95	
Fusobacterium varium		0.61	0.25		1	1	1	
11051001	**Good identification**							**11051001**
Clostridium sordellii		100.0	0.37	MLT 90				
11051020	**Low discrimination**							**11051020**
					SPO	CAT		
Propionibacterium acnes		66.79	0.40	NIT 92	1	95		
Clostridium cadaveris		32.39	0.32	CEL 1	99	1		
Atopobium minutum		0.82	0.15		1	1		
11052020	**Low discrimination**							**11052020**
					SPO	CAT		
Propionibacterium acnes		50.07	0.36	NIT 92	1	95		
Clostridium cadaveris		43.18	0.32	CEL 1	99	1		
Pseudoramibacter alactolyticus		6.76	0.14		1	1		
11053020	**Good identification**							**11053020**
					SPO	CAT		
Propionibacterium acnes		99.14	0.36	NIT 92	1	95		
Clostridium cadaveris		0.86	-.03	CEL 1	99	1		
11054000	**Good identification**							**11054000**
					GRA	SPO		
Fusobacterium varium		97.52	0.35	CEL 1	1	1		
Clostridium glycolicum		1.47	0.09	XYL 4	99	99		
Pseudoramibacter alactolyticus		0.63	-.03		99	1		
11060020	**Good identification**							**11060020**
Clostridium cadaveris		99.86	0.44	TRE 5				
Clostridium bifermentans		0.14	-.03	CEL 1				
11070020	**Good identification**							**11070020**
					SPO	CAT		
Clostridium cadaveris		99.60	0.44	TRE 5	99	1		
Propionibacterium acnes		0.40	0.06	CEL 1	1	95		
11100000	**Low discrimination**							**11100000**
					COC	SPO		
Clostridium bifermentans		87.13	0.92		1	99		
Clostridium sordellii		4.55	0.66		1	99		
Peptostreptococcus anaerobius		2.90	0.63		99	1		
11100001	**Excellent identification**							**11100001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sordellii		99.68	1.00			
Clostridium bifermentans		0.20	0.58			
Clostridium novyi biovar A		0.11	0.66			
11100002	**Low discrimination**					**11100002**
Clostridium bifermentans		68.53	0.58	BGL 1		
Clostridium sporogenes		14.55	0.45			
Clostridium novyi biovar A		6.54	0.53			
11100003	**Very good identification**					**11100003**
Clostridium sordellii		97.94	0.66	BGL 1		
Clostridium novyi biovar A		1.81	0.53			
Clostridium bifermentans		0.19	0.24			
11100010	**Low discrimination**					**11100010**
Clostridium bifermentans		59.45	0.58	MLZ 1		
Clostridium novyi biovar A		32.13	0.66			
Clostridium sordellii		3.10	0.32			
11100011	**Good identification**					**11100011**
Clostridium sordellii		90.36	0.66	MLZ 1		
Clostridium novyi biovar A		9.46	0.66			
Clostridium bifermentans		0.18	0.24			
11100012	**Low discrimination**					**11100012**
Clostridium novyi biovar A		87.92	0.53	IND 1		
Clostridium bifermentans		9.31	0.24			
Clostridium sporogenes		1.98	0.11			
11100013	**Low discrimination**					**11100013**
Clostridium novyi biovar A		64.64	0.53	IND 1		
Clostridium sordellii		35.36	0.32			
11100020	**Low discrimination**					**11100020**
Clostridium cadaveris		75.03	0.78	MLT 5		
Clostridium bifermentans		23.91	0.72			
Clostridium novyi biovar A		0.51	0.56			
11100021	**Good identification**					**11100021**
Clostridium sordellii		92.82	0.66	NAG 1		
Clostridium cadaveris		3.60	0.44			
Clostridium novyi biovar A		2.43	0.56			
11100022	**Low discrimination**					**11100022**
Clostridium cadaveris		68.19	0.44	MLT 5		
Clostridium bifermentans		21.73	0.38	BGL 1		
Clostridium novyi biovar A		8.12	0.43			
11100030	**Low discrimination**					**11100030**
Clostridium cadaveris		50.06	0.44	MLT 5		
Clostridium novyi biovar A		33.77	0.56	MLZ 1		
Clostridium bifermentans		15.95	0.38			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
11100031	**Low discrimination**						**11100031**
Clostridium novyi biovar A		71.13	0.56	IND 1			
Clostridium sordellii		27.46	0.32				
Clostridium cadaveris		1.07	0.10				
11100032	**Low discrimination**						**11100032**
Clostridium novyi biovar A		89.94	0.43	IND 1			
Clostridium cadaveris		7.63	0.10				
Clostridium bifermentans		2.43	0.03				
11100033	**Good identification**						**11100033**
Clostridium novyi biovar A		97.84	0.43	IND 1			
Clostridium sordellii		2.16	-.03				
11100040	**Low discrimination**						**11100040**
Clostridium bifermentans		79.66	0.69	SOR 4			
Clostridium novyi biovar A		6.96	0.63				
Clostridium botulinum biovar A		6.72	0.51				
11100041	**Very good identification**						**11100041**
Clostridium sordellii		98.13	0.76	SOR 4			
Clostridium novyi biovar A		1.66	0.63				
Clostridium bifermentans		0.19	0.34				
11100042	**Low discrimination**						**11100042**
					COC	GRA	
Clostridium novyi biovar A		54.22	0.50	IND 1	1	99	
Clostridium bifermentans		35.53	0.34		1	99	
Megasphaera elsdenii		3.29	0.22		99	1	
11100043	**Low discrimination**						**11100043**
Clostridium sordellii		77.08	0.42	SOR 4			
Clostridium novyi biovar A		22.77	0.50	BGL 1			
Clostridium bifermentans		0.15	0.00				
11100050	**Low discrimination**						**11100050**
Clostridium novyi biovar A		88.24	0.63	IND 1			
Clostridium bifermentans		10.20	0.34				
Clostridium botulinum biovar A		0.86	0.17				
11100051	**Low discrimination**						**11100051**
Clostridium novyi biovar A		62.56	0.63	IND 1			
Clostridium sordellii		37.36	0.42				
Clostridium bifermentans		0.07	0.00				
11100052	**Good identification**						**11100052**
Clostridium novyi biovar A		99.34	0.50	IND 1			
Clostridium bifermentans		0.66	0.00				
11100053	**Good identification**						**11100053**
Clostridium novyi biovar A		96.69	0.50	IND 1			
Clostridium sordellii		3.31	0.08				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
11100060	**Low discrimination**						**11100060**
Clostridium bifermentans		47.01	0.48	NAG 6			
Clostridium cadaveris		35.77	0.44	SOR 4			
Clostridium novyi biovar A		16.09	0.53				
11100061	**Low discrimination**						**11100061**
Clostridium sordellii		69.43	0.42	NAG 1			
Clostridium novyi biovar A		29.06	0.53	SOR 4			
Clostridium bifermentans		0.86	0.14				
11100062	**Low discrimination**						**11100062**
Clostridium novyi biovar A		77.25	0.40	IND 1			
Clostridium bifermentans		12.92	0.14				
Clostridium cadaveris		9.83	0.10				
11100063	**Low discrimination**						**11100063**
Clostridium novyi biovar A		87.97	0.40	IND 1			
Clostridium sordellii		12.03	0.08				
11100070	**Good identification**						**11100070**
Clostridium novyi biovar A		95.06	0.53	IND 1			
Clostridium bifermentans		2.81	0.14				
Clostridium cadaveris		2.14	0.10				
11100071	**Good identification**						**11100071**
Clostridium novyi biovar A		97.64	0.53	IND 1			
Clostridium sordellii		2.36	0.08				
11100072	**Good identification**						**11100072**
Clostridium novyi biovar A		100.0	0.40	IND 1			
11100073	**Good identification**						**11100073**
Clostridium novyi biovar A		100.0	0.40	IND 1			
11100100	**Low discrimination**						**11100100**
					COC	SPO	
Clostridium bifermentans		86.89	0.58	LAC 1	1	99	
Clostridium sordellii		4.53	0.32		1	99	
Peptostreptococcus anaerobius		2.89	0.29		99	1	
11100101	**Very good identification**						**11100101**
Clostridium sordellii		99.69	0.66	LAC 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
11100120	**Low discrimination**						**11100120**
					GRA	SPO	
Clostridium cadaveris		51.90	0.44	MLT 5	99	99	
Prevotella bivia		30.91	0.32	LAC 1	1	1	
Clostridium bifermentans		16.54	0.38		99	99	
11100141	**Good identification**						**11100141**
Clostridium sordellii		98.15	0.42	LAC 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		1.66	0.29	SOR 4			
Clostridium bifermentans		0.19	0.00				
11100200	**Low discrimination**						**11100200**
					COC	SPO	
Clostridium bifermentans		87.18	0.58	RHA 1	1	99	
Clostridium sordellii		4.55	0.32		1	99	
Peptostreptococcus anaerobius		2.90	0.29		99	1	
11100201	**Very good identification**						**11100201**
Clostridium sordellii		99.69	0.66	RHA 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
11100220	**Low discrimination**						**11100220**
Clostridium cadaveris		75.20	0.44	MLT 5			
Clostridium bifermentans		23.96	0.38	RHA 1			
Clostridium novyi biovar A		0.51	0.21				
11100241	**Good identification**						**11100241**
Clostridium sordellii		98.15	0.42	RHA 1			
Clostridium novyi biovar A		1.66	0.29	SOR 4			
Clostridium bifermentans		0.19	0.00				
11100400	**Low discrimination**						**11100400**
					COC	SPO	
Clostridium bifermentans		87.05	0.58	ARA 1	1	99	
Clostridium sordellii		4.54	0.32		1	99	
Peptostreptococcus anaerobius		2.90	0.29		99	1	
11100401	**Very good identification**						**11100401**
Clostridium sordellii		99.69	0.66	ARA 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
11100420	**Low discrimination**						**11100420**
Clostridium cadaveris		75.20	0.44	MLT 5			
Clostridium bifermentans		23.96	0.38	ARA 1			
Clostridium novyi biovar A		0.51	0.21				
11100441	**Good identification**						**11100441**
Clostridium sordellii		98.15	0.42	ARA 1			
Clostridium novyi biovar A		1.66	0.29	SOR 4			
Clostridium bifermentans		0.19	0.00				
11101000	**Low discrimination**						**11101000**
					GRA	SPO	
Clostridium bifermentans		30.99	0.58	GAL 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		28.70	0.51		1	1	
Clostridium sordellii		28.24	0.53		99	99	
11101001	**Very good identification**						**11101001**
Clostridium sordellii		99.91	0.87				
Clostridium novyi biovar A		0.07	0.50				
Clostridium bifermentans		0.01	0.24				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
11101002	**Low discrimination**						**11101002**
					GRA	SPO	
Clostridium novyi biovar A		22.91	0.37	IND 1	99	99	
Clostridium bifermentans		21.83	0.24	GAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		20.23	0.17		1	1	
11101003	**Good identification**						**11101003**
Clostridium sordellii		98.85	0.53	BGL 1			
Clostridium novyi biovar A		1.15	0.37				
11101010	**Low discrimination**						**11101010**
					GRA	SPO	
Clostridium novyi biovar A		65.18	0.50	IND 1	99	99	
Clostridium bifermentans		10.96	0.24	GAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		10.16	0.17		1	1	
11101011	**Acceptable identification**						**11101011**
Clostridium sordellii		93.82	0.53	MLZ 1			
Clostridium novyi biovar A		6.18	0.50				
11101012	**Good identification**						**11101012**
Clostridium novyi biovar A		100.0	0.37	IND 1			
11101013	**Low discrimination**						**11101013**
Clostridium novyi biovar A		53.51	0.37	IND 1			
Clostridium sordellii		46.49	0.19	GAL 10			
11101020	**Low discrimination**						**11101020**
Clostridium cadaveris		66.06	0.44	MLT 5			
Clostridium bifermentans		21.05	0.38	GAL 1			
Clostridium novyi biovar A		4.95	0.39				
11101021	**Good identification**						**11101021**
Clostridium sordellii		98.10	0.53	NAG 1			
Clostridium novyi biovar A		1.62	0.39				
Clostridium cadaveris		0.22	0.10				
11101030	**Low discrimination**						**11101030**
Clostridium novyi biovar A		84.91	0.39	IND 1			
Clostridium cadaveris		11.44	0.10	GAL 10			
Clostridium bifermentans		3.65	0.03				
11101031	**Low discrimination**						**11101031**
Clostridium novyi biovar A		61.99	0.39	IND 1			
Clostridium sordellii		38.01	0.19	GAL 10			
11101041	**Very good identification**						**11101041**
Clostridium sordellii		98.94	0.64	SOR 4			
Clostridium novyi biovar A		1.05	0.47				
Clostridium bifermentans		0.01	0.00				
11101050	**Good identification**						**11101050**
Clostridium novyi biovar A		98.03	0.47	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		1.03	0.00	GAL 10			
Clostridium sordellii		0.94	-.05				
11101051	**Low discrimination**						**11101051**
Clostridium novyi biovar A		51.32	0.47	IND 1			
Clostridium sordellii		48.68	0.29	GAL 10			
11101060	**Low discrimination**						**11101060**
Clostridium novyi biovar A		66.39	0.36	IND 1			
Clostridium bifermentans		17.64	0.14	GAL 10			
Clostridium cadaveris		13.42	0.10				
11101070	**Good identification**						**11101070**
Clostridium novyi biovar A		100.0	0.36	IND 1			
11101071	**Good identification**						**11101071**
Clostridium novyi biovar A		96.31	0.36	IND 1			
Clostridium sordellii		3.69	-.05	GAL 10			
11101101	**Good identification**						**11101101**
Clostridium sordellii		99.83	0.53	LAC 1			
Clostridium novyi biovar A		0.07	0.15				
Clostridium chauvoei		0.05	0.03				
11101201	**Good identification**						**11101201**
Clostridium sordellii		99.93	0.53	RHA 1			
Clostridium novyi biovar A		0.07	0.15				
11101401	**Good identification**						**11101401**
Clostridium sordellii		99.93	0.53	ARA 1			
Clostridium novyi biovar A		0.07	0.15				
11102000	**Low discrimination**						**11102000**
					COC	GRA	
Clostridium bifermentans		83.40	0.58	MAN 1	1	99	
Clostridium sordellii		4.35	0.32		1	99	
Megasphaera elsdenii		4.34	0.41		99	1	
11102001	**Very good identification**						**11102001**
Clostridium sordellii		99.68	0.66	MAN 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
11102020	**Low discrimination**						**11102020**
Clostridium cadaveris		74.93	0.44	MLT 5			
Clostridium bifermentans		23.88	0.38	MAN 1			
Clostridium novyi biovar A		0.51	0.21				
11102041	**Good identification**						**11102041**
Clostridium sordellii		97.99	0.42	MAN 1			
Clostridium novyi biovar A		1.66	0.29	SOR 4			
Clostridium bifermentans		0.19	0.00				
11103001	**Good identification**						**11103001**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sordellii		99.93	0.53	MAN 1				
Clostridium novyi biovar A		0.07	0.15					
11104000	**Low discrimination**							**11104000**
					COC	SPO		
Clostridium bifermentans		85.57	0.58	XYL 1	1	99		
Clostridium sordellii		4.46	0.32		1	99		
Peptostreptococcus anaerobius		2.85	0.29		99	1		
11104001	**Very good identification**							**11104001**
Clostridium sordellii		99.69	0.66	XYL 1				
Clostridium bifermentans		0.20	0.24					
Clostridium novyi biovar A		0.11	0.32					
11104020	**Low discrimination**							**11104020**
Clostridium cadaveris		75.20	0.44	MLT 5				
Clostridium bifermentans		23.96	0.38	XYL 1				
Clostridium novyi biovar A		0.51	0.21					
11104040	**Low discrimination**							**11104040**
Clostridium glycolicum		86.79	0.50	IND 1				
Clostridium bifermentans		10.65	0.34	FRU 90				
Clostridium novyi biovar A		0.93	0.29					
11104041	**Good identification**							**11104041**
Clostridium sordellii		96.64	0.42	XYL 1				
Clostridium novyi biovar A		1.63	0.29	SOR 4				
Clostridium glycolicum		1.54	0.15					
11105001	**Good identification**							**11105001**
Clostridium sordellii		99.93	0.53	XYL 1				
Clostridium novyi biovar A		0.07	0.15					
11105722	**Good identification**							**11105722**
Bacteroides eggerthii		100.0	0.39	ESL 90				
11110000	**Low discrimination**							**11110000**
					GRA	SPO		
Prevotella intermedia		59.35	0.84		1	1		
Clostridium bifermentans		13.10	0.70		99	99		
Clostridium sordellii		12.99	0.66		99	99		
11110001	**Excellent identification**							**11110001**
					GRA	SPO		
Clostridium sordellii		99.92	1.00		99	99		
Prevotella intermedia		0.05	0.50		1	1		
Clostridium novyi biovar A		0.01	0.50		99	99		
11110002	**Low discrimination**							**11110002**
					COC	GRA	SPO	
Prevotella intermedia		42.58	0.50	BGL 1	1	1	1	
Megasphaera elsdenii		19.64	0.47		99	1	1	
Clostridium bifermentans		9.40	0.36		1	99	99	
11110003	**Very good identification**							**11110003**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sordellii		99.71	0.66	BGL 1	99	99		
Clostridium novyi biovar A		0.21	0.37		99	99		
Prevotella intermedia		0.05	0.16		1	1		
11110010	**Low discrimination**							**11110010**
					GRA	SPO		
Prevotella intermedia		51.69	0.50	MLZ 1	1	1		
Clostridium novyi biovar A		13.02	0.50		99	99		
Clostridium bifermentans		11.41	0.36		99	99		
11110011	**Very good identification**							**11110011**
					GRA	SPO		
Clostridium sordellii		98.80	0.66	MLZ 1	99	99		
Clostridium novyi biovar A		1.15	0.50		99	99		
Prevotella intermedia		0.05	0.16		1	1		
11110012	**Low discrimination**							**11110012**
					COC	GRA	SPO	
Clostridium novyi biovar A		68.14	0.37	IND 1	1	99	99	
Prevotella intermedia		15.48	0.16	FRU 10	1	1	1	
Megasphaera elsdenii		7.14	0.12		99	1	1	
11110020	**Very good identification**							**11110020**
					GRA	SPO		
Clostridium cadaveris		96.17	0.78	MLT 5	99	99		
Clostridium bifermentans		1.61	0.50		99	99		
Prevotella intermedia		1.16	0.50		1	1		
11110021	**Very good identification**							**11110021**
Clostridium sordellii		95.88	0.66	NAG 1				
Clostridium cadaveris		3.72	0.44					
Clostridium novyi biovar A		0.28	0.39					
11110022	**Acceptable identification**							**11110022**
Clostridium cadaveris		93.28	0.44	MLT 5				
Clostridium bifermentans		1.56	0.16	BGL 1				
Clostridium novyi biovar A		1.23	0.26					
11110030	**Acceptable identification**							**11110030**
Clostridium cadaveris		90.06	0.44	MLT 5				
Clostridium novyi biovar A		6.75	0.39	MLZ 1				
Clostridium bifermentans		1.51	0.16					
11110040	**Low discrimination**							**11110040**
					COC	GRA	SPO	
Megasphaera elsdenii		40.81	0.60	IND 1	99	1	1	
Prevotella intermedia		14.75	0.50		1	1	1	
Clostridium bifermentans		13.43	0.47		1	99	99	
11110041	**Very good identification**							**11110041**
					COC	GRA		
Clostridium sordellii		99.75	0.76	SOR 4	1	99		
Clostridium novyi biovar A		0.19	0.47		1	99		
Megasphaera elsdenii		0.03	0.26		99	1		
11110042	**Low discrimination**							**11110042**
					COC	GRA	SPO	
Megasphaera elsdenii		80.64	0.44	IND 1	99	1	1	
Clostridium novyi biovar A		7.77	0.34	BGL 10	1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		2.65	0.16		1	1	1	
11110043	**Good identification**							**11110043**
					COC	GRA		
Clostridium sordellii		96.50	0.42	SOR 4	1	99		
Clostridium novyi biovar A		3.17	0.34	BGL 1	1	99		
Megasphaera elsdenii		0.33	0.09		99	1		
11110050	**Low discrimination**							**11110050**
					COC	GRA	SPO	
Clostridium novyi biovar A		71.74	0.47	IND 1	1	99	99	
Megasphaera elsdenii		11.94	0.26	FRU 10	99	1	1	
Prevotella intermedia		4.32	0.16		1	1	1	
11110051	**Low discrimination**							**11110051**
Clostridium sordellii		84.31	0.42	MLZ 1				
Clostridium novyi biovar A		15.69	0.47	SOR 4				
11110060	**Low discrimination**							**11110060**
					SPO	CAT		
Clostridium cadaveris		77.66	0.44	MLT 5	99	1		
Propionibacterium acnes		6.75	0.29	SOR 1	1	95		
Clostridium bifermentans		5.37	0.26		99	1		
11110061	**Acceptable identification**							**11110061**
Clostridium sordellii		94.71	0.42	NAG 1				
Clostridium novyi biovar A		4.41	0.36	SOR 4				
Clostridium cadaveris		0.89	0.10					
11110070	**Low discrimination**							**11110070**
Clostridium novyi biovar A		80.83	0.36	IND 1				
Clostridium cadaveris		16.34	0.10	FRU 10				
Clostridium botulinum biovar B		2.83	-.03					
11110071	**Low discrimination**							**11110071**
Clostridium novyi biovar A		82.16	0.36	IND 1				
Clostridium sordellii		17.84	0.08	FRU 10				
11110100	**Low discrimination**							**11110100**
					GRA	SPO		
Prevotella intermedia		53.85	0.50	LAC 1	1	1		
Clostridium bifermentans		11.89	0.36		99	99		
Clostridium sordellii		11.78	0.32		99	99		
11110101	**Very good identification**							**11110101**
					GRA	SPO		
Clostridium sordellii		99.92	0.66	LAC 1	99	99		
Prevotella intermedia		0.05	0.16		1	1		
Clostridium novyi biovar A		0.01	0.15		99	99		
11110120	**Low discrimination**							**11110120**
					GRA	SPO		
Clostridium cadaveris		88.55	0.44	MLT 5	99	99		
Prevotella bivia		5.86	0.15	LAC 1	1	1		
Actinomyces meyeri		2.46	0.24		99	1		
11110141	**Good identification**							**11110141**
Clostridium sordellii		99.81	0.42	LAC 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		0.19	0.12	SOR 4			
11110200	**Low discrimination**						**11110200**
					GRA	SPO	
Prevotella intermedia		59.44	0.50	RHA 1	1	1	
Clostridium bifermentans		13.12	0.36		99	99	
Clostridium sordellii		13.01	0.32		99	99	
11110201	**Very good identification**						**11110201**
					GRA	SPO	
Clostridium sordellii		99.93	0.66	RHA 1	99	99	
Prevotella intermedia		0.05	0.16		1	1	
Clostridium novyi biovar A		0.01	0.15		99	99	
11110220	**Good identification**						**11110220**
					GRA	SPO	
Clostridium cadaveris		96.58	0.44	MLT 5	99	99	
Clostridium bifermentans		1.62	0.16	RHA 1	99	99	
Prevotella intermedia		1.16	0.16		1	1	
11110241	**Good identification**						**11110241**
Clostridium sordellii		99.81	0.42	RHA 1			
Clostridium novyi biovar A		0.19	0.12	SOR 4			
11110400	**Low discrimination**						**11110400**
					GRA	SPO	
Prevotella intermedia		58.77	0.50	ARA 1	1	1	
Clostridium bifermentans		12.98	0.36		99	99	
Clostridium sordellii		12.86	0.32		99	99	
11110401	**Very good identification**						**11110401**
					GRA	SPO	
Clostridium sordellii		99.93	0.66	ARA 1	99	99	
Prevotella intermedia		0.05	0.16		1	1	
Clostridium novyi biovar A		0.01	0.15		99	99	
11110420	**Good identification**						**11110420**
					GRA	SPO	
Clostridium cadaveris		96.42	0.44	MLT 5	99	99	
Clostridium bifermentans		1.62	0.16	ARA 1	99	99	
Prevotella intermedia		1.16	0.16		1	1	
11110441	**Good identification**						**11110441**
Clostridium sordellii		99.81	0.42	ARA 1			
Clostridium novyi biovar A		0.19	0.12	SOR 4			
11111000	**Very good identification**						**11111000**
					GRA	SPO	
Prevotella intermedia		95.58	0.84		1	1	
Clostridium sordellii		3.69	0.53		99	99	
Clostridium bifermentans		0.21	0.36		99	99	
11111001	**Very good identification**						**11111001**
					GRA	SPO	
Clostridium sordellii		99.73	0.87		99	99	
Prevotella intermedia		0.26	0.50		1	1	
Clostridium novyi biovar A		0.01	0.33		99	99	
11111002	**Low discrimination**						**11111002**
					GRA	SPO	
Prevotella intermedia		86.91	0.50	BGL 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		7.53	0.44		99	1		
Clostridium sordellii		3.36	0.19		99	99		
11111003	**Good identification**							**11111003**
					GRA	SPO		
Clostridium sordellii		99.61	0.53	BGL 1	99	99		
Prevotella intermedia		0.26	0.16		1	1		
Clostridium novyi biovar A		0.13	0.20		99	99		
11111010	**Acceptable identification**							**11111010**
					GRA	SPO		
Prevotella intermedia		92.83	0.50	MLZ 1	1	1		
Clostridium sordellii		3.59	0.19		99	99		
Clostridium novyi biovar A		2.60	0.33		99	99		
11111011	**Good identification**							**11111011**
					GRA	SPO		
Clostridium sordellii		99.01	0.53	MLZ 1	99	99		
Clostridium novyi biovar A		0.73	0.33		99	99		
Prevotella intermedia		0.26	0.16		1	1		
11111020	**Low discrimination**							**11111020**
					GRA	SPO	CAT	
Prevotella intermedia		43.85	0.50	NAG 1	1	1	1	
Clostridium cadaveris		36.84	0.44		99	99	1	
Propionibacterium acnes		14.58	0.40		99	1	95	
11111021	**Good identification**							**11111021**
					GRA	SPO		
Clostridium sordellii		99.25	0.53	NAG 1	99	99		
Prevotella intermedia		0.26	0.16		1	1		
Clostridium cadaveris		0.22	0.10		99	99		
11111022	**Low discrimination**							**11111022**
					GRA	SPO		
Lactobacillus fermentum		52.72	0.36	IND 1	99	1		
Prevotella intermedia		18.44	0.16	SUC 87	1	1		
Clostridium cadaveris		15.49	0.10		99	99		
11111040	**Low discrimination**							**11111040**
					GRA	SPO		
Prevotella intermedia		77.52	0.50	SOR 1	1	1		
Clostridium sordellii		12.35	0.29		99	99		
Clostridium glycolicum		5.81	0.28		99	99		
11111041	**Very good identification**							**11111041**
					GRA	SPO		
Clostridium sordellii		99.82	0.64	SOR 4	99	99		
Clostridium novyi biovar A		0.12	0.30		99	99		
Prevotella intermedia		0.06	0.16		1	1		
11111100	**Low discrimination**							**11111100**
					GRA			
Prevotella intermedia		68.10	0.50	LAC 1	1			
Actinomyces meyeri		17.46	0.42		99			
Lactobacillus fermentum		9.63	0.48		99			
11111101	**Good identification**							**11111101**
					GRA	SPO		
Clostridium sordellii		99.52	0.53	LAC 1	99	99		
Prevotella intermedia		0.26	0.16		1	1		
Clostridium chauvoei		0.12	0.09		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
11111102	**Low discrimination**						**11111102**
					GRA		
Lactobacillus fermentum		86.40	0.48	IND 1	99		
Prevotella intermedia		6.17	0.16	SUC 87	1		
Actinomyces meyeri		4.85	0.16		99		
11111120	**Low discrimination**						**11111120**
					GRA		
Lactobacillus fermentum		37.06	0.40	IND 1	99		
Actinomyces meyeri		27.49	0.27	SUC 87	99		
Prevotella intermedia		7.94	0.16		1		
11111122	**Good identification**						**11111122**
Lactobacillus fermentum		95.87	0.40	IND 1			
Clostridium perfringens		4.13	0.04	SUC 87			
11111200	**Good identification**						**11111200**
					GRA	SPO	
Prevotella intermedia		96.08	0.50	RHA 1	1	1	
Clostridium sordellii		3.71	0.19		99	99	
Clostridium bifermentans		0.21	0.02		99	99	
11111201	**Good identification**						**11111201**
					GRA	SPO	
Clostridium sordellii		99.74	0.53	RHA 1	99	99	
Prevotella intermedia		0.26	0.16		1	1	
11111400	**Low discrimination**						**11111400**
					GRA	SPO	
Prevotella intermedia		87.10	0.50	ARA 1	1	1	
Lactobacillus fermentum		7.55	0.44		99	1	
Clostridium sordellii		3.36	0.19		99	99	
11111401	**Good identification**						**11111401**
					GRA	SPO	
Clostridium sordellii		99.74	0.53	ARA 1	99	99	
Prevotella intermedia		0.26	0.16		1	1	
11111402	**Low discrimination**						**11111402**
					GRA		
Lactobacillus fermentum		89.56	0.44	IND 1	99		
Prevotella intermedia		10.44	0.16	SUC 87	1		
11111420	**Low discrimination**						**11111420**
					GRA	SPO	
Lactobacillus fermentum		54.49	0.36	IND 1	99	1	
Prevotella intermedia		19.05	0.16	SUC 87	1	1	
Clostridium cadaveris		16.01	0.10		99	99	
11111422	**Good identification**						**11111422**
Lactobacillus fermentum		100.0	0.36	IND 1			
11111500	**Low discrimination**						**11111500**
					GRA		
Lactobacillus fermentum		65.59	0.48	IND 1	99		
Actinomyces meyeri		29.73	0.31	SUC 87	99		
Prevotella intermedia		4.69	0.16		1		
11111502	**Good identification**						**11111502**
Lactobacillus fermentum		98.62	0.48	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		1.38	0.06	SUC 87				
11111520	**Low discrimination**							**11111520**
					GRA			
Lactobacillus fermentum		79.89	0.40	IND 1	99			
Actinomyces meyeri		14.81	0.17	SUC 87	99			
Odoribacter splanchnicus		5.30	-.03		1			
11111522	**Good identification**							**11111522**
Lactobacillus fermentum		100.0	0.40	IND 1				
11112000	**Low discrimination**							**11112000**
					COC	GRA	SPO	
Megasphaera elsdenii		68.83	0.63	IND 1	99	1	1	
Prevotella intermedia		16.59	0.50		1	1	1	
Clostridium bifermentans		3.66	0.36		1	99	99	
11112001	**Very good identification**							**11112001**
					COC	GRA	SPO	
Clostridium sordellii		99.73	0.66	MAN 1	1	99	99	
Megasphaera elsdenii		0.19	0.29		99	1	1	
Prevotella intermedia		0.05	0.16		1	1	1	
11112002	**Acceptable identification**							**11112002**
					COC	GRA		
Megasphaera elsdenii		93.51	0.47	IND 1	99	1		
Pseudoramibacter alactolyticus		3.22	0.11	BGL 10	1	99		
Prevotella intermedia		2.05	0.16		1	1		
11112020	**Low discrimination**							**11112020**
					COC	SPO	CAT	
Clostridium cadaveris		74.97	0.44	MLT 5	1	99	1	
Propionibacterium acnes		16.69	0.36	MAN 1	1	1	95	
Megasphaera elsdenii		3.74	0.29		99	1	1	
11112040	**Very good identification**							**11112040**
					COC	GRA		
Megasphaera elsdenii		97.44	0.60	IND 1	99	1		
Pseudoramibacter alactolyticus		1.26	0.17		1	99		
Prevotella intermedia		0.36	0.16		1	1		
11112041	**Good identification**							**11112041**
					COC	GRA		
Clostridium sordellii		96.82	0.42	MAN 1	1	99		
Megasphaera elsdenii		3.00	0.26	SOR 4	99	1		
Clostridium novyi biovar A		0.18	0.12		1	99		
11112042	**Good identification**							**11112042**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	IND 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
11113000	**Acceptable identification**							**11113000**
					COC	GRA	SPO	
Prevotella intermedia		92.07	0.50	MAN 1	1	1	1	
Megasphaera elsdenii		3.86	0.29		99	1	1	
Clostridium sordellii		3.56	0.19		1	99	99	
11113001	**Good identification**							**11113001**
					GRA	SPO		
Clostridium sordellii		99.74	0.53	MAN 1	99	99		
Prevotella intermedia		0.26	0.16		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
11113020	**Acceptable identification**							**11113020**
					GRA	SPO	CAT	
Propionibacterium acnes		90.96	0.36	NIT 92	99	1	95	
Prevotella intermedia		4.91	0.16	MLT 1	1	1	1	
Clostridium cadaveris		4.13	0.10		99	99	1	
11114000	**Low discrimination**							**11114000**
					GRA	SPO		
Prevotella intermedia		36.19	0.50	XYL 1	1	1		
Clostridium glycolicum		24.41	0.44		99	99		
Fusobacterium varium		15.05	0.35		1	1		
11114001	**Very good identification**							**11114001**
					GRA	SPO		
Clostridium sordellii		99.88	0.66	XYL 1	99	99		
Prevotella intermedia		0.05	0.16		1	1		
Clostridium glycolicum		0.03	0.10		99	99		
11114020	**Acceptable identification**							**11114020**
					SPO			
Clostridium cadaveris		94.16	0.44	MLT 5	99			
Clostridium bifermentans		1.58	0.16	XYL 1	99			
Actinomyces meyeri		1.27	0.19		1			
11114040	**Very good identification**							**11114040**
					COC	GRA	SPO	
Clostridium glycolicum		99.52	0.66	IND 1	1	99	99	
Megasphaera elsdenii		0.22	0.26		99	1	1	
Prevotella intermedia		0.08	0.16		1	1	1	
11114041	**Low discrimination**							**11114041**
Clostridium sordellii		87.32	0.42	XYL 1				
Clostridium glycolicum		12.52	0.32	SOR 4				
Clostridium novyi biovar A		0.16	0.12					
11114100	**Acceptable identification**							**11114100**
					GRA	SPO		
Actinomyces meyeri		92.85	0.44	IND 1	99	1		
Prevotella intermedia		2.83	0.16	SUC 93	1	1		
Clostridium glycolicum		1.91	0.10		99	99		
11115000	**Low discrimination**							**11115000**
					GRA	SPO		
Prevotella intermedia		76.61	0.50	XYL 1	1	1		
Actinomyces meyeri		9.53	0.36		99	1		
Clostridium glycolicum		5.74	0.28		99	99		
11115001	**Good identification**							**11115001**
					GRA	SPO		
Clostridium sordellii		99.70	0.53	XYL 1	99	99		
Prevotella intermedia		0.26	0.16		1	1		
Actinomyces meyeri		0.03	0.02		99	1		
11115002	**Low discrimination**							**11115002**
					GRA			
Lactobacillus fermentum		78.49	0.41	IND 1	99			
Prevotella intermedia		15.58	0.16	SUC 87	1			
Actinomyces meyeri		5.93	0.11		99			
11115040	**Good identification**							**11115040**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium glycolicum		99.10	0.50	IND 1	99	99	
Prevotella intermedia		0.70	0.16	GAL 10	1	1	
Clostridium sordellii		0.11	-.05		99	99	
11115100	**Low discrimination**						**11115100**
Actinomyces meyeri		82.43	0.47	IND 1			
Lactobacillus fermentum		13.76	0.44	SUC 93			
Eubacterium saburreum		2.14	0.27				
11115102	**Low discrimination**						**11115102**
Lactobacillus fermentum		82.45	0.44	IND 1			
Actinomyces meyeri		15.28	0.21	SUC 87			
Eubacterium saburreum		2.26	0.14				
11115122	**Acceptable identification**						**11115122**
Lactobacillus fermentum		92.95	0.36	IND 1			
Actinomyces meyeri		7.05	0.06	SUC 87			
11115400	**Low discrimination**						**11115400**
					GRA		
Lactobacillus fermentum		55.26	0.41	IND 1	99		
Actinomyces meyeri		33.77	0.26	SUC 87	99		
Prevotella intermedia		10.97	0.16		1		
11115402	**Good identification**						**11115402**
Lactobacillus fermentum		100.0	0.41	IND 1			
11115500	**Low discrimination**						**11115500**
Actinomyces meyeri		59.97	0.36	IND 1			
Lactobacillus fermentum		40.03	0.44	SUC 93			
11115502	**Good identification**						**11115502**
Lactobacillus fermentum		95.57	0.44	IND 1			
Actinomyces meyeri		4.43	0.11	SUC 87			
11115520	**Low discrimination**						**11115520**
Lactobacillus fermentum		62.00	0.36	IND 1			
Actinomyces meyeri		38.00	0.22	SUC 87			
11115522	**Good identification**						**11115522**
Lactobacillus fermentum		100.0	0.36	IND 1			
11120000	**Low discrimination**						**11120000**
Clostridium bifermentans		60.95	0.58	TRE 1			
Clostridium botulinum biovar A		17.74	0.49				
Clostridium sporogenes		12.94	0.45				
11120001	**Very good identification**						**11120001**
Clostridium sordellii		99.60	0.66	TRE 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
11120002	**Low discrimination**						**11120002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		82.64	0.37	IND 1			
Clostridium bifermentans		11.80	0.24				
Clostridium botulinum biovar A		3.43	0.15				
11120020	**Good identification**						**11120020**
Clostridium cadaveris		93.62	0.56	MLT 5			
Clostridium bifermentans		5.73	0.38	TRE 5			
Clostridium botulinum biovar A		0.26	0.15				
11120040	**Low discrimination**						**11120040**
Clostridium botulinum biovar A		62.85	0.41	IND 1			
Clostridium bifermentans		30.12	0.34				
Clostridium novyi biovar A		2.63	0.29				
11120041	**Good identification**						**11120041**
Clostridium sordellii		97.76	0.42	TRE 1			
Clostridium novyi biovar A		1.65	0.29	SOR 4			
Clostridium botulinum biovar A		0.40	0.06				
11121001	**Good identification**						**11121001**
Clostridium sordellii		99.93	0.53	TRE 1			
Clostridium novyi biovar A		0.07	0.15				
11130000	**Low discrimination**						**11130000**
					GRA	SPO	
Prevotella intermedia		51.41	0.50	TRE 1	1	1	
Clostridium bifermentans		11.35	0.36		99	99	
Clostridium sordellii		11.25	0.32		99	99	
11130001	**Very good identification**						**11130001**
					GRA	SPO	
Clostridium sordellii		99.92	0.66	TRE 1	99	99	
Prevotella intermedia		0.05	0.16		1	1	
Clostridium novyi biovar A		0.01	0.15		99	99	
11130020	**Very good identification**						**11130020**
					GRA	SPO	
Clostridium cadaveris		99.29	0.56	MLT 5	99	99	
Clostridium bifermentans		0.32	0.16	TRE 5	99	99	
Prevotella intermedia		0.23	0.16		1	1	
11130041	**Good identification**						**11130041**
Clostridium sordellii		99.81	0.42	TRE 1			
Clostridium novyi biovar A		0.19	0.12	SOR 4			
11131000	**Good identification**						**11131000**
					SPO	GRA	
Prevotella intermedia		95.30	0.50	TRE 1	1	1	
Clostridium sordellii		3.68	0.19		99	99	
Eubacterium saburreum		0.65	0.22		1	85	
11131001	**Good identification**						**11131001**
					GRA	SPO	
Clostridium sordellii		99.74	0.53	TRE 1	99	99	
Prevotella intermedia		0.26	0.16		1	1	
11140000	**Low discrimination**						**11140000**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		87.18	0.58	CEL 1	1	99	
Clostridium sordellii		4.55	0.32		1	99	
Peptostreptococcus anaerobius		2.90	0.29		99	1	
11140001	**Very good identification**						**11140001**
Clostridium sordellii		99.69	0.66	CEL 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
11140020	**Low discrimination**						**11140020**
Clostridium cadaveris		75.20	0.44	MLT 5			
Clostridium bifermentans		23.96	0.38	CEL 1			
Clostridium novyi biovar A		0.51	0.21				
11140041	**Good identification**						**11140041**
Clostridium sordellii		98.15	0.42	CEL 1			
Clostridium novyi biovar A		1.66	0.29	SOR 4			
Clostridium bifermentans		0.19	0.00				
11141001	**Good identification**						**11141001**
Clostridium sordellii		99.93	0.53	CEL 1			
Clostridium novyi biovar A		0.07	0.15				
11145722	**Good identification**						**11145722**
Bacteroides eggerthii		100.0	0.50	ESL 90			
11150000	**Low discrimination**						**11150000**
					GRA	SPO	
Prevotella intermedia		59.44	0.50	CEL 1	1	1	
Clostridium bifermentans		13.12	0.36		99	99	
Clostridium sordellii		13.01	0.32		99	99	
11150001	**Very good identification**						**11150001**
					GRA	SPO	
Clostridium sordellii		99.93	0.66	CEL 1	99	99	
Prevotella intermedia		0.05	0.16		1	1	
Clostridium novyi biovar A		0.01	0.15		99	99	
11150020	**Good identification**						**11150020**
					GRA	SPO	
Clostridium cadaveris		96.58	0.44	MLT 5	99	99	
Clostridium bifermentans		1.62	0.16	CEL 1	99	99	
Prevotella intermedia		1.16	0.16		1	1	
11150041	**Good identification**						**11150041**
Clostridium sordellii		99.81	0.42	CEL 1			
Clostridium novyi biovar A		0.19	0.12	SOR 4			
11151000	**Good identification**						**11151000**
					GRA	SPO	
Prevotella intermedia		96.08	0.50	CEL 1	1	1	
Clostridium sordellii		3.71	0.19		99	99	
Clostridium bifermentans		0.21	0.02		99	99	
11151001	**Good identification**						**11151001**
					GRA	SPO	
Clostridium sordellii		99.74	0.53	CEL 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		0.26	0.16		1	1	
11200000	**Low discrimination**						**11200000**
					GRA	SPO	
Clostridium bifermentans		73.93	0.70	SAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		14.53	0.51		1	1	
Fusobacterium nucleatum subsp. nucleatum		2.35	0.32		1	1	
11200001	**Low discrimination**						**11200001**
Clostridium sordellii		83.53	0.50	MLT 90			
Clostridium bifermentans		6.93	0.35	SAL 1			
Clostridium novyi biovar A		6.84	0.48				
11200002	**Low discrimination**						**11200002**
					GRA	SPO	
Clostridium bifermentans		56.89	0.35	SAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		11.18	0.17	BGL 1	1	1	
Clostridium novyi biovar A		9.91	0.35		99	99	
11200010	**Low discrimination**						**11200010**
					GRA	SPO	
Clostridium bifermentans		43.19	0.35	SAL 10	99	99	
Clostridium novyi biovar A		42.63	0.48	MLZ 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.49	0.17		1	1	
11200011	**Low discrimination**						**11200011**
Clostridium novyi biovar A		88.22	0.48	IND 1			
Clostridium sordellii		10.88	0.15	SAL 8			
Clostridium bifermentans		0.90	0.01				
11200020	**Good identification**						**11200020**
Clostridium cadaveris		91.98	0.66	SAL 1			
Clostridium bifermentans		7.27	0.49				
Clostridium novyi biovar A		0.28	0.37				
11200031	**Good identification**						**11200031**
Clostridium novyi biovar A		96.80	0.37	IND 1			
Clostridium cadaveris		3.20	-.03	SAL 8			
11200040	**Low discrimination**						**11200040**
					GRA	SPO	
Clostridium bifermentans		79.31	0.46	SAL 10	99	99	
Clostridium novyi biovar A		12.65	0.45	SOR 4	99	99	
Fusobacterium necrophorum subsp. necrophorum		3.78	0.17		1	1	
11200041	**Low discrimination**						**11200041**
Clostridium novyi biovar A		54.75	0.45	IND 1			
Clostridium sordellii		41.78	0.26	SAL 8			
Clostridium bifermentans		3.47	0.12				
11200050	**Acceptable identification**						**11200050**
Clostridium novyi biovar A		94.05	0.45	IND 1			
Clostridium bifermentans		5.96	0.12	SAL 8			
11200051	**Good identification**						**11200051**
Clostridium novyi biovar A		100.0	0.45	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
11200100	**Low discrimination**							**11200100**
					GRA	SPO		
Clostridium bifermentans		74.73	0.35	SAL 10	99	99		
Fusobacterium necrophorum subsp. necrophorum		14.68	0.17	LAC 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.37	-.03		1	1		
11200200	**Low discrimination**							**11200200**
					GRA	SPO		
Clostridium bifermentans		74.73	0.35	SAL 10	99	99		
Fusobacterium necrophorum subsp. necrophorum		14.68	0.17	RHA 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.37	-.03		1	1		
11200400	**Low discrimination**							**11200400**
					GRA	SPO		
Clostridium bifermentans		74.73	0.35	SAL 10	99	99		
Fusobacterium necrophorum subsp. necrophorum		14.68	0.17	ARA 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.37	-.03		1	1		
11201000	**Acceptable identification**							**11201000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		93.56	0.51	GLU 12	1	1		
Clostridium bifermentans		4.81	0.35	SAL 1	99	99		
Clostridium novyi biovar A		0.53	0.31		99	99		
11201001	**Low discrimination**							**11201001**
					GRA	SPO		
Clostridium sordellii		87.05	0.37	MLT 90	99	99		
Fusobacterium necrophorum subsp. necrophorum		8.04	0.17	SAL 1	1	1		
Clostridium novyi biovar A		4.49	0.31		99	99		
11202000	**Low discrimination**							**11202000**
					GRA	SPO		
Clostridium bifermentans		73.49	0.35	SAL 10	99	99		
Fusobacterium necrophorum subsp. necrophorum		14.44	0.17	MAN 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.33	-.03		1	1		
11204000	**Low discrimination**							**11204000**
					GRA	SPO		
Clostridium bifermentans		73.57	0.35	SAL 10	99	99		
Fusobacterium necrophorum subsp. necrophorum		14.46	0.17	XYL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.34	-.03		1	1		
11210000	**Low discrimination**							**11210000**
					GRA	SPO		
Fusobacterium varium		85.35	0.59	SAL 1	1	1		
Clostridium bifermentans		8.90	0.48		99	99		
Pseudoramibacter alactolyticus		2.27	0.32		99	1		
11210001	**Acceptable identification**							**11210001**
					GRA	SPO		
Clostridium sordellii		94.59	0.50	MLT 90	99	99		
Fusobacterium varium		3.96	0.25	SAL 1	1	1		
Clostridium novyi biovar A		0.86	0.31		99	99		
11210020	**Very good identification**							**11210020**
					GRA	SPO	CAT	
Clostridium cadaveris		96.59	0.66	SAL 1	99	99	1	
Propionibacterium acnes		2.01	0.40		99	1	95	
Fusobacterium varium		0.61	0.25		1	1	1	
11211001	**Good identification**							**11211001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sordellii		99.43	0.37	MLT 90			
Clostridium novyi biovar A		0.57	0.15	SAL 1			
11211020	**Low discrimination**						**11211020**
					SPO	CAT	
Propionibacterium acnes		66.79	0.40	NIT 92	1	95	
Clostridium cadaveris		32.39	0.32	SAL 1	99	1	
Atopobium minutum		0.82	0.15		1	1	
11211100	**Good identification**						**11211100**
					GRA		
Fusobacterium mortiferum		96.27	0.40	IND 1	1		
Actinomyces meyeri		3.73	0.07		99		
11212010	**Good identification**						**11212010**
					SPO		
Clostridium difficile		99.81	0.43	IND 1	99		
Pseudoramibacter alactolyticus		0.19	-.07		1		
11212020	**Low discrimination**						**11212020**
					SPO	CAT	
Propionibacterium acnes		49.86	0.36	NIT 92	1	95	
Clostridium cadaveris		42.99	0.32	SAL 1	99	1	
Pseudoramibacter alactolyticus		6.73	0.14		1	1	
11213020	**Good identification**						**11213020**
					SPO	CAT	
Propionibacterium acnes		99.14	0.36	NIT 92	1	95	
Clostridium cadaveris		0.86	-.03	SAL 1	99	1	
11214000	**Acceptable identification**						**11214000**
					GRA	SPO	
Fusobacterium varium		94.89	0.35	SAL 1	1	1	
Clostridium bifermentans		2.40	0.14	XYL 4	99	99	
Clostridium glycolicum		1.44	0.09		99	99	
11220000	**Low discrimination**						**11220000**
					GRA	SPO	
Clostridium bifermentans		67.92	0.35	SAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		13.35	0.17	TRE 1	1	1	
Clostridium botulinum biovar A		5.13	0.17		99	99	
11220020	**Good identification**						**11220020**
Clostridium cadaveris		98.51	0.44	SAL 1			
Clostridium bifermentans		1.50	0.15	TRE 5			
11230020	**Good identification**						**11230020**
					SPO	CAT	
Clostridium cadaveris		99.60	0.44	SAL 1	99	1	
Propionibacterium acnes		0.40	0.06	TRE 5	1	95	
11240000	**Low discrimination**						**11240000**
					GRA	SPO	
Clostridium bifermentans		74.73	0.35	SAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		14.68	0.17	CEL 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		2.37	-.03		1	1	
11300000	**Very good identification**						**11300000**
Clostridium bifermentans		98.31	0.76	SAL 10			
Clostridium sordellii		0.47	0.32				
Clostridium novyi biovar A		0.42	0.48				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
11300001	**Very good identification**					**11300001**
Clostridium sordellii		97.02	0.66	SAL 1		
Clostridium bifermentans		2.09	0.42			
Clostridium novyi biovar A		0.88	0.48			
11300002	**Acceptable identification**					**11300002**
Clostridium bifermentans		90.77	0.42	SAL 10		
Clostridium novyi biovar A		6.78	0.35	BGL 1		
Clostridium sporogenes		1.75	0.11			
11300010	**Low discrimination**					**11300010**
Clostridium bifermentans		69.89	0.42	SAL 10		
Clostridium novyi biovar A		29.57	0.48	MLZ 1		
Clostridium sordellii		0.33	-.03			
11300020	**Low discrimination**					**11300020**
Clostridium bifermentans		76.73	0.55	SAL 10		
Clostridium cadaveris		21.89	0.44	NAG 6		
Clostridium novyi biovar A		1.28	0.37			
11300030	**Low discrimination**					**11300030**
Clostridium novyi biovar A		56.31	0.37	IND 1		
Clostridium bifermentans		33.99	0.21	SAL 8		
Clostridium cadaveris		9.70	0.10			
11300031	**Good identification**					**11300031**
Clostridium novyi biovar A		95.71	0.37	IND 1		
Clostridium sordellii		4.29	-.03	SAL 8		
11300040	**Acceptable identification**					**11300040**
Clostridium bifermentans		92.40	0.52	SAL 10		
Clostridium novyi biovar A		6.32	0.45	SOR 4		
Clostridium botulinum biovar A		0.71	0.17			
11300041	**Low discrimination**					**11300041**
Clostridium sordellii		85.68	0.42	SAL 1		
Clostridium novyi biovar A		12.48	0.45	SOR 4		
Clostridium bifermentans		1.84	0.18			
11300050	**Low discrimination**					**11300050**
Clostridium novyi biovar A		87.13	0.45	IND 1		
Clostridium bifermentans		12.87	0.18	SAL 8		
11300051	**Acceptable identification**					**11300051**
Clostridium novyi biovar A		93.51	0.45	IND 1		
Clostridium sordellii		6.49	0.08	SAL 8		
11300100	**Good identification**					**11300100**
Clostridium bifermentans		98.81	0.42	SAL 10		
Clostridium sordellii		0.47	-.03	LAC 1		
Clostridium novyi biovar A		0.42	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
11300200	**Good identification**						**11300200**
Clostridium bifermentans		98.81	0.42	SAL 10			
Clostridium sordellii		0.47	-.03	RHA 1			
Clostridium novyi biovar A		0.42	0.14				
11300400	**Good identification**						**11300400**
Clostridium bifermentans		98.81	0.42	SAL 10			
Clostridium sordellii		0.47	-.03	ARA 1			
Clostridium novyi biovar A		0.42	0.14				
11301000	**Low discrimination**						**11301000**
					GRA	SPO	
Clostridium bifermentans		80.73	0.42	SAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		6.80	0.17	GAL 1	1	1	
Clostridium sordellii		6.69	0.19		99	99	
11301001	**Good identification**						**11301001**
Clostridium sordellii		99.31	0.53	SAL 1			
Clostridium novyi biovar A		0.57	0.31				
Clostridium bifermentans		0.12	0.08				
11302000	**Good identification**						**11302000**
					COC	GRA	
Clostridium bifermentans		98.35	0.42	SAL 10	1	99	
Megasphaera elsdenii		0.47	0.07	MAN 1	99	1	
Clostridium sordellii		0.47	-.03		1	99	
11304000	**Good identification**						**11304000**
Clostridium bifermentans		98.81	0.42	SAL 10			
Clostridium sordellii		0.47	-.03	XYL 1			
Clostridium novyi biovar A		0.42	0.14				
11310000	**Low discrimination**						**11310000**
					GRA	SPO	
Clostridium bifermentans		62.09	0.54	SAL 10	99	99	
Prevotella intermedia		25.57	0.50	FRU 5	1	1	
Clostridium sordellii		5.60	0.32		99	99	
11310001	**Very good identification**						**11310001**
Clostridium sordellii		99.74	0.66	SAL 1			
Clostridium bifermentans		0.11	0.20				
Clostridium novyi biovar A		0.10	0.31				
11310020	**Low discrimination**						**11310020**
					GRA	SPO	
Clostridium cadaveris		82.67	0.44	MLT 5	99	99	
Clostridium bifermentans		15.25	0.34	SAL 1	99	99	
Prevotella intermedia		0.99	0.16		1	1	
11310041	**Good identification**						**11310041**
Clostridium sordellii		98.41	0.42	SAL 1			
Clostridium novyi biovar A		1.59	0.28	SOR 4			
11311000	**Acceptable identification**						**11311000**
					GRA	SPO	
Prevotella intermedia		93.57	0.50	SAL 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sordellii		3.61	0.19		99	99		
Clostridium bifermentans		2.30	0.20		99	99		
11311001	**Good identification**							**11311001**
					GRA	SPO		
Clostridium sordellii		99.67	0.53	SAL 1	99	99		
Prevotella intermedia		0.26	0.16		1	1		
Clostridium novyi biovar A		0.06	0.15		99	99		
11320000	**Acceptable identification**							**11320000**
Clostridium bifermentans		94.54	0.42	SAL 10				
Clostridium botulinum biovar A		2.50	0.15	TRE 1				
Clostridium sporogenes		1.82	0.11					
11340000	**Good identification**							**11340000**
Clostridium bifermentans		98.81	0.42	SAL 10				
Clostridium sordellii		0.47	-.03	CEL 1				
Clostridium novyi biovar A		0.42	0.14					
11400000	**Low discrimination**							**11400000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		44.95	0.51	GLU 12	1	1		
Clostridium bifermentans		20.80	0.52	RAF 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		7.27	0.32		1	1		
11400001	**Low discrimination**							**11400001**
					COC	GRA	SPO	
Clostridium sordellii		86.33	0.50	MLT 90	1	99	99	
Anaerococcus prevotii		9.70	0.32	RAF 1	99	99	1	
Fusobacterium necrophorum subsp. necrophorum		1.41	0.17		1	1	1	
11400020	**Very good identification**							**11400020**
					GRA	SPO		
Clostridium cadaveris		98.76	0.66	RAF 1	99	99		
Clostridium bifermentans		0.71	0.31		99	99		
Fusobacterium necrophorum subsp. necrophorum		0.24	0.17		1	1		
11401000	**Good identification**							**11401000**
					GRA	SPO		
Fusobacterium necrophorum subsp. necrophorum		97.94	0.51	GLU 12	1	1		
Clostridium bifermentans		0.46	0.18	RAF 1	99	99		
Prevotella intermedia		0.38	0.19		1	1		
11401001	**Acceptable identification**							**11401001**
					COC	GRA	SPO	
Clostridium sordellii		90.51	0.37	MLT 90	1	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.36	0.17	RAF 1	1	1	1	
Anaerococcus prevotii		0.58	-.02		99	99	1	
11410000	**Low discrimination**							**11410000**
					GRA			
Fusobacterium varium		81.80	0.59	RAF 1	1			
Prevotella intermedia		12.29	0.53		1			
Pseudoramibacter alactolyticus		2.18	0.32		99			
11410001	**Good identification**							**11410001**
					GRA	SPO		
Clostridium sordellii		95.04	0.50	MLT 90	99	99		
Fusobacterium varium		3.98	0.25	RAF 1	1	1		
Prevotella intermedia		0.60	0.19		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
11410020	**Very good identification**							**11410020**
					GRA	SPO	CAT	
Clostridium cadaveris		96.88	0.66	RAF 1	99	99	1	
Propionibacterium acnes		2.02	0.40		99	1	95	
Fusobacterium varium		0.61	0.25		1	1	1	
11411000	**Low discrimination**							**11411000**
					GRA			
Prevotella intermedia		86.30	0.53	MLT 99	1			
Fusobacterium varium		5.80	0.25		1			
Lactobacillus fermentum		2.71	0.40		99			
11411001	**Good identification**							**11411001**
					GRA	SPO		
Clostridium sordellii		96.56	0.37	MLT 90	99	99		
Prevotella intermedia		3.44	0.19	RAF 1	1	1		
11411002	**Low discrimination**							**11411002**
					GRA			
Lactobacillus fermentum		64.32	0.40	IND 1	99			
Prevotella intermedia		20.68	0.19	SUC 87	1			
Atopobium minutum		11.15	0.20		99			
11411020	**Low discrimination**							**11411020**
					SPO	CAT		
Propionibacterium acnes		62.89	0.40	NIT 92	1	95		
Clostridium cadaveris		30.50	0.32	RAF 1	99	1		
Lactobacillus fermentum		2.97	0.32		1	1		
11411100	**Low discrimination**							**11411100**
					GRA			
Lactobacillus fermentum		48.43	0.44	IND 1	99			
Fusobacterium mortiferum		37.30	0.40	SUC 87	1			
Prevotella intermedia		9.54	0.19		1			
11411102	**Acceptable identification**							**11411102**
					GRA			
Lactobacillus fermentum		92.12	0.44	IND 1	99			
Fusobacterium mortiferum		7.88	0.24	SUC 87	1			
11411120	**Low discrimination**							**11411120**
					SPO	CAT		
Lactobacillus fermentum		83.72	0.35	IND 1	1	1		
Propionibacterium acnes		10.97	0.06	SUC 87	1	95		
Clostridium cadaveris		5.32	-.03		99	1		
11411122	**Good identification**							**11411122**
Lactobacillus fermentum		100.0	0.35	IND 1				
11411400	**Low discrimination**							**11411400**
					GRA			
Lactobacillus fermentum		75.67	0.40	IND 1	99			
Prevotella intermedia		24.33	0.19	SUC 87	1			
11411402	**Good identification**							**11411402**
Lactobacillus fermentum		100.0	0.40	IND 1				
11411500	**Good identification**							**11411500**
Lactobacillus fermentum		100.0	0.44	IND 1				
11411502	**Good identification**							**11411502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		100.0	0.44	IND 1			
11411520	**Low discrimination**						**11411520**
					GRA		
Lactobacillus fermentum		89.14	0.35	IND 1	99		
Odoribacter splanchnicus		10.86	-.03	SUC 87	1		
11411522	**Good identification**						**11411522**
Lactobacillus fermentum		100.0	0.35	IND 1			
11412020	**Low discrimination**						**11412020**
					SPO	CAT	
Propionibacterium acnes		50.07	0.36	NIT 92	1	95	
Clostridium cadaveris		43.18	0.32	RAF 1	99	1	
Pseudoramibacter alactolyticus		6.76	0.14		1	1	
11413020	**Good identification**						**11413020**
					SPO	CAT	
Propionibacterium acnes		99.14	0.36	NIT 92	1	95	
Clostridium cadaveris		0.86	-.03	RAF 1	99	1	
11414000	**Acceptable identification**						**11414000**
					GRA	SPO	
Fusobacterium varium		93.90	0.35	RAF 1	1	1	
Prevotella intermedia		3.42	0.19	XYL 4	1	1	
Clostridium glycolicum		1.42	0.09		99	99	
11415000	**Low discrimination**						**11415000**
					GRA		
Lactobacillus fermentum		56.56	0.36	IND 1	99		
Prevotella intermedia		30.96	0.19	SUC 87	1		
Fusobacterium varium		8.59	0.01		1		
11415002	**Good identification**						**11415002**
Lactobacillus fermentum		100.0	0.36	IND 1			
11415100	**Low discrimination**						**11415100**
					GRA		
Lactobacillus fermentum		80.91	0.40	IND 1	99		
Fusobacterium mortiferum		7.99	0.21	SUC 87	1		
Actinomyces meyeri		7.99	0.12		99		
11415102	**Good identification**						**11415102**
Lactobacillus fermentum		100.0	0.40	IND 1			
11415400	**Good identification**						**11415400**
Lactobacillus fermentum		100.0	0.36	IND 1			
11415402	**Good identification**						**11415402**
Lactobacillus fermentum		100.0	0.36	IND 1			
11415500	**Good identification**						**11415500**
Lactobacillus fermentum		100.0	0.40	IND 1			
11415502	**Good identification**						**11415502**
Lactobacillus fermentum		100.0	0.40	IND 1			
11420020	**Good identification**						**11420020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium cadaveris		99.86	0.44	RAF 1				
Clostridium bifermentans		0.14	-.03	TRE 5				
11430020	**Good identification**							**11430020**
					SPO	CAT		
Clostridium cadaveris		99.60	0.44	RAF 1	99	1		
Propionibacterium acnes		0.40	0.06	TRE 5	1	95		
11500000	**Low discrimination**							**11500000**
					GRA	SPO		
Clostridium bifermentans		66.54	0.58	RAF 1	99	99		
Prevotella intermedia		23.80	0.53		1	1		
Clostridium sordellii		3.47	0.32		99	99		
11500001	**Very good identification**							**11500001**
Clostridium sordellii		99.61	0.66	RAF 1				
Clostridium bifermentans		0.20	0.24					
Clostridium novyi biovar A		0.11	0.32					
11500020	**Low discrimination**							**11500020**
					GRA	SPO		
Clostridium cadaveris		74.19	0.44	MLT 5	99	99		
Clostridium bifermentans		23.64	0.38	RAF 1	99	99		
Prevotella intermedia		1.34	0.19		1	1		
11500041	**Good identification**							**11500041**
Clostridium sordellii		98.15	0.42	RAF 1				
Clostridium novyi biovar A		1.66	0.29	SOR 4				
Clostridium bifermentans		0.19	0.00					
11501000	**Low discrimination**							**11501000**
					COC	GRA	SPO	
Prevotella intermedia		82.20	0.53	FRU 99	1	1	1	
Blautia hansenii		10.81	0.32		99	99	1	
Clostridium bifermentans		2.32	0.24		1	99	99	
11501001	**Good identification**							**11501001**
					GRA	SPO		
Clostridium sordellii		99.49	0.53	RAF 1	99	99		
Prevotella intermedia		0.39	0.19		1	1		
Clostridium novyi biovar A		0.07	0.15		99	99		
11501100	**Very good identification**							**11501100**
					COC	GRA		
Blautia hansenii		99.89	0.66	IND 1	99	99		
Prevotella intermedia		0.08	0.19		1	1		
Eubacterium saburreum		0.02	0.19		1	85		
11501102	**Good identification**							**11501102**
					COC			
Blautia hansenii		99.89	0.50	IND 1	99			
Lactobacillus fermentum		0.11	0.18	BGL 10	1			
11501120	**Good identification**							**11501120**
					COC	GRA		
Blautia hansenii		99.88	0.50	IND 1	99	99		
Prevotella melaninogenica		0.12	-.01	NAG 10	1	1		
11501500	**Good identification**							**11501500**
					COC			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		99.93	0.53	IND 1	99			
Lactobacillus fermentum		0.07	0.18		1			
11501502	**Good identification**							**11501502**
					COC			
Blautia hansenii		99.40	0.37	IND 1	99			
Lactobacillus fermentum		0.61	0.18	BGL 10	1			
11501520	**Good identification**							**11501520**
Blautia hansenii		100.0	0.37	IND 1				
11505100	**Good identification**							**11505100**
					COC			
Blautia hansenii		99.89	0.53	IND 1	99			
Eubacterium saburreum		0.07	0.14		1			
Lactobacillus fermentum		0.04	0.14		1			
11505102	**Good identification**							**11505102**
					COC			
Blautia hansenii		99.64	0.37	IND 1	99			
Lactobacillus fermentum		0.36	0.14	BGL 10	1			
11505120	**Good identification**							**11505120**
Blautia hansenii		100.0	0.37	IND 1				
11505500	**Good identification**							**11505500**
					COC			
Blautia hansenii		99.78	0.40	IND 1	99			
Lactobacillus fermentum		0.23	0.14		1			
11510000	**Very good identification**							**11510000**
					GRA	SPO		
Prevotella intermedia		99.54	0.87		1	1		
Clostridium bifermentans		0.15	0.36		99	99		
Clostridium sordellii		0.15	0.32		99	99		
11510001	**Good identification**							**11510001**
					GRA	SPO		
Clostridium sordellii		93.51	0.66	RAF 1	99	99		
Prevotella intermedia		6.47	0.53		1	1		
Clostridium novyi biovar A		0.01	0.15		99	99		
11510002	**Good identification**							**11510002**
					COC	GRA	SPO	
Prevotella intermedia		99.20	0.53	BGL 1	1	1	1	
Megasphaera elsdenii		0.31	0.12		99	1	1	
Clostridium bifermentans		0.15	0.02		1	99	99	
11510010	**Good identification**							**11510010**
					GRA	SPO		
Prevotella intermedia		99.54	0.53	MLZ 1	1	1		
Clostridium novyi biovar A		0.17	0.15		99	99		
Clostridium bifermentans		0.15	0.02		99	99		
11510020	**Low discrimination**							**11510020**
					GRA	SPO		
Prevotella intermedia		63.56	0.53	NAG 1	1	1		
Clostridium cadaveris		35.60	0.44		99	99		
Clostridium bifermentans		0.60	0.16		99	99		
11510040	**Good identification**							**11510040**
					COC	GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		96.48	0.53	SOR 1	1	1	1	
Megasphaera elsdenii		1.80	0.26		99	1	1	
Clostridium bifermentans		0.59	0.13		1	99	99	
11510041	**Good identification**							**11510041**
					GRA	SPO		
Clostridium sordellii		98.17	0.42	RAF 1	99	99		
Prevotella intermedia		1.65	0.19	SOR 4	1	1		
Clostridium novyi biovar A		0.18	0.12		99	99		
11510100	**Good identification**							**11510100**
					GRA			
Prevotella intermedia		98.99	0.53	LAC 1	1			
Fusobacterium mortiferum		0.32	0.22		1			
Lactobacillus fermentum		0.15	0.18		99			
11510200	**Good identification**							**11510200**
					GRA	SPO		
Prevotella intermedia		99.71	0.53	RHA 1	1	1		
Clostridium bifermentans		0.15	0.02		99	99		
Clostridium sordellii		0.15	-.03		99	99		
11510400	**Good identification**							**11510400**
					GRA	SPO		
Prevotella intermedia		99.61	0.53	ARA 1	1	1		
Clostridium bifermentans		0.15	0.02		99	99		
Clostridium sordellii		0.15	-.03		99	99		
11511000	**Very good identification**							**11511000**
					GRA	SPO		
Prevotella intermedia		99.87	0.87		1	1		
Lactobacillus fermentum		0.10	0.48		99	1		
Clostridium sordellii		0.03	0.19		99	99		
11511001	**Low discrimination**							**11511001**
					GRA	SPO		
Clostridium sordellii		71.80	0.53	RAF 1	99	99		
Prevotella intermedia		28.17	0.53		1	1		
Lactobacillus fermentum		0.03	0.14		99	1		
11511002	**Acceptable identification**							**11511002**
					GRA			
Prevotella intermedia		91.39	0.53	BGL 1	1			
Lactobacillus fermentum		8.61	0.48		99			
11511010	**Good identification**							**11511010**
					GRA			
Prevotella intermedia		99.60	0.53	MLZ 1	1			
Eubacterium saburreum		0.30	0.19		85			
Lactobacillus fermentum		0.10	0.14		99			
11511020	**Good identification**							**11511020**
					GRA	SPO		
Prevotella intermedia		96.22	0.53	NAG 1	1	1		
Lactobacillus fermentum		3.02	0.40		99	1		
Clostridium cadaveris		0.54	0.10		99	99		
11511022	**Low discrimination**							**11511022**
					GRA			
Lactobacillus fermentum		75.67	0.40	IND 1	99			
Prevotella intermedia		24.33	0.19	SUC 87	1			
11511040	**Good identification**							**11511040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Prevotella intermedia		99.80	0.53	SOR 1	1	1	
Clostridium sordellii		0.11	-.05		99	99	
Lactobacillus fermentum		0.10	0.14		99	1	
11511100	**Low discrimination**						**11511100**
					COC	GRA	
Prevotella intermedia		76.89	0.53	LAC 1	1	1	
Lactobacillus fermentum		11.83	0.52		1	99	
Blautia hansenii		10.11	0.32		99	99	
11511102	**Low discrimination**						**11511102**
					COC	GRA	
Lactobacillus fermentum		85.43	0.52	IND 1	1	99	
Blautia hansenii		8.12	0.15	SUC 87	99	99	
Prevotella intermedia		5.61	0.19		1	1	
11511120	**Low discrimination**						**11511120**
					COC	GRA	
Lactobacillus fermentum		55.33	0.44	IND 1	1	99	
Prevotella melaninogenica		16.36	0.15	SUC 87	1	1	
Blautia hansenii		15.77	0.15		99	99	
11511122	**Acceptable identification**						**11511122**
					COC	GRA	
Lactobacillus fermentum		91.91	0.44	IND 1	1	99	
Prevotella melaninogenica		5.18	0.03	SUC 87	1	1	
Blautia hansenii		2.91	-.01		99	99	
11511200	**Good identification**						**11511200**
					GRA		
Prevotella intermedia		99.91	0.53	RHA 1	1		
Lactobacillus fermentum		0.10	0.14		99		
11511400	**Acceptable identification**						**11511400**
					GRA		
Prevotella intermedia		91.39	0.53	ARA 1	1		
Lactobacillus fermentum		8.61	0.48		99		
11511402	**Acceptable identification**						**11511402**
					GRA		
Lactobacillus fermentum		90.32	0.48	IND 1	99		
Prevotella intermedia		9.68	0.19	SUC 87	1		
11511420	**Low discrimination**						**11511420**
					GRA		
Lactobacillus fermentum		75.67	0.40	IND 1	99		
Prevotella intermedia		24.33	0.19	SUC 87	1		
11511422	**Good identification**						**11511422**
Lactobacillus fermentum		100.0	0.40	IND 1			
11511500	**Low discrimination**						**11511500**
					COC	GRA	
Lactobacillus fermentum		82.20	0.52	IND 1	1	99	
Blautia hansenii		12.40	0.19	SUC 87	99	99	
Prevotella intermedia		5.40	0.19		1	1	
11511502	**Good identification**						**11511502**
					COC		
Lactobacillus fermentum		98.35	0.52	IND 1	1		
Blautia hansenii		1.65	0.03	SUC 87	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
11511520	**Acceptable identification**							**11511520**
					COC			
Lactobacillus fermentum		93.84	0.44	IND 1	1			
Blautia hansenii		4.72	0.03	SUC 87	99			
Bifidobacterium longum subsp. longum		1.44	0.05		1			
11511522	**Good identification**							**11511522**
Lactobacillus fermentum		98.49	0.44	IND 1				
Bifidobacterium longum subsp. longum		1.51	0.05	SUC 87				
11512000	**Good identification**							**11512000**
					COC	GRA	SPO	
Prevotella intermedia		96.91	0.53	MAN 1	1	1	1	
Megasphaera elsdenii		2.71	0.29		99	1	1	
Clostridium bifermentans		0.14	0.02		1	99	99	
11513000	**Good identification**							**11513000**
					GRA			
Prevotella intermedia		99.91	0.53	MAN 1	1			
Lactobacillus fermentum		0.10	0.14		99			
11514000	**Good identification**							**11514000**
					GRA	SPO		
Prevotella intermedia		98.98	0.53	XYL 1	1	1		
Clostridium glycolicum		0.45	0.10		99	99		
Fusobacterium varium		0.28	0.01		1	1		
11515000	**Acceptable identification**							**11515000**
					GRA			
Prevotella intermedia		94.50	0.53	XYL 1	1			
Lactobacillus fermentum		5.23	0.44		99			
Eubacterium saburreum		0.19	0.16		85			
11515002	**Low discrimination**							**11515002**
					GRA			
Lactobacillus fermentum		84.57	0.44	IND 1	99			
Prevotella intermedia		15.43	0.19	SUC 87	1			
11515020	**Low discrimination**							**11515020**
					GRA			
Lactobacillus fermentum		64.63	0.36	IND 1	99			
Prevotella intermedia		35.38	0.19	SUC 87	1			
11515022	**Good identification**							**11515022**
Lactobacillus fermentum		100.0	0.36	IND 1				
11515100	**Low discrimination**							**11515100**
					COC	GRA		
Lactobacillus fermentum		68.62	0.48	IND 1	1	99		
Blautia hansenii		17.63	0.19	SUC 87	99	99		
Prevotella intermedia		7.67	0.19		1	1		
11515102	**Good identification**							**11515102**
					COC			
Lactobacillus fermentum		96.38	0.48	IND 1	1			
Blautia hansenii		2.75	0.03	SUC 87	99			
Eubacterium saburreum		0.87	0.10		1			
11515120	**Low discrimination**							**11515120**
					COC			
Lactobacillus fermentum		89.86	0.40	IND 1	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		7.70	0.03	SUC 87	99		
Eubacterium saburreum		2.44	0.10		1		
11515122	**Good identification**						**11515122**
Lactobacillus fermentum		100.0	0.40	IND 1			
11515400	**Low discrimination**						**11515400**
					GRA		
Lactobacillus fermentum		84.57	0.44	IND 1	99		
Prevotella intermedia		15.43	0.19	SUC 87	1		
11515402	**Good identification**						**11515402**
Lactobacillus fermentum		100.0	0.44	IND 1			
11515420	**Good identification**						**11515420**
Lactobacillus fermentum		100.0	0.36	IND 1			
11515422	**Good identification**						**11515422**
Lactobacillus fermentum		100.0	0.36	IND 1			
11515500	**Acceptable identification**						**11515500**
					COC		
Lactobacillus fermentum		94.82	0.48	IND 1	1		
Blautia hansenii		4.30	0.06	SUC 87	99		
Actinomyces meyeri		0.88	0.02		1		
11515502	**Good identification**						**11515502**
Lactobacillus fermentum		100.0	0.48	IND 1			
11515520	**Acceptable identification**						**11515520**
Lactobacillus fermentum		92.74	0.40	IND 1			
Bifidobacterium longum subsp. longum		7.26	0.14	SUC 87			
11515522	**Acceptable identification**						**11515522**
Lactobacillus fermentum		92.74	0.40	IND 1			
Bifidobacterium longum subsp. longum		7.26	0.14	SUC 87			
11530000	**Good identification**						**11530000**
					GRA	SPO	
Prevotella intermedia		99.60	0.53	TRE 1	1	1	
Clostridium bifermentans		0.15	0.02		99	99	
Clostridium sordellii		0.15	-.03		99	99	
11531000	**Good identification**						**11531000**
					GRA		
Prevotella intermedia		99.66	0.53	TRE 1	1		
Eubacterium saburreum		0.25	0.18		85		
Lactobacillus fermentum		0.10	0.14		99		
11550000	**Good identification**						**11550000**
					GRA	SPO	
Prevotella intermedia		99.71	0.53	CEL 1	1	1	
Clostridium bifermentans		0.15	0.02		99	99	
Clostridium sordellii		0.15	-.03		99	99	
11551000	**Good identification**						**11551000**
					GRA		
Prevotella intermedia		99.91	0.53	CEL 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		0.10	0.14		99		
11600000	**Low discrimination**						**11600000**
					GRA	SPO	
Clostridium bifermentans		74.73	0.35	SAL 10	99	99	
Fusobacterium necrophorum subsp. necrophorum		14.68	0.17	RAF 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		2.37	-.03		1	1	
11610100	**Good identification**						**11610100**
Fusobacterium mortiferum		100.0	0.42	IND 1			
11611000	**Low discrimination**						**11611000**
Fusobacterium mortiferum		76.48	0.39	IND 1			
Prevotella intermedia		23.52	0.19				
11611100	**Good identification**						**11611100**
Fusobacterium mortiferum		100.0	0.45	IND 1			
11631100	**Good identification**						**11631100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
11651100	**Good identification**						**11651100**
Fusobacterium mortiferum		100.0	0.37	IND 1			
11700000	**Good identification**						**11700000**
					GRA	SPO	
Clostridium bifermentans		95.74	0.42	SAL 10	99	99	
Prevotella intermedia		3.11	0.19	RAF 1	1	1	
Clostridium sordellii		0.45	-.03		99	99	
11710000	**Good identification**						**11710000**
					GRA	SPO	
Prevotella intermedia		97.99	0.53	SAL 1	1	1	
Clostridium bifermentans		1.60	0.20		99	99	
Fusobacterium mortiferum		0.27	0.20		1	1	
11711000	**Good identification**						**11711000**
					GRA		
Prevotella intermedia		99.50	0.53	SAL 1	1		
Fusobacterium mortiferum		0.40	0.23		1		
Lactobacillus fermentum		0.10	0.14		99		
12000000	**Low discrimination**						**12000000**
					COC	GRA	
Fusobacterium nucleatum subsp. nucleatum		24.44	0.66	SUC 1	1	1	
Peptoniphilus asaccharolyticus		24.44	0.66		99	99	
Porphyromonas asaccharolytica		24.19	0.66		1	1	
12000001	**Low discrimination**						**12000001**
					COC	GRA	
Anaerococcus prevotii		20.51	0.41	IND 9	99	99	
Fusobacterium nucleatum subsp. nucleatum		15.89	0.32	SUC 1	1	1	
Peptoniphilus asaccharolyticus		15.89	0.32		99	99	
12000002	**Low discrimination**						**12000002**
					COC	GRA	
Porphyromonas asaccharolytica		53.39	0.42	SUC 1	1	1	
Fusobacterium nucleatum subsp. nucleatum		13.08	0.32	BGL 4	1	1	
Peptoniphilus asaccharolyticus		13.08	0.32		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
12000020	**Low discrimination**							**12000020**
					COC	GRA	SPO	
Clostridium cadaveris		70.14	0.59	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		7.04	0.32	SUC 7	1	1	1	
Peptoniphilus asaccharolyticus		7.04	0.32		99	99	1	
12001000	**Good identification**							**12001000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		92.47	0.66	SUC 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.04	0.32		1	1		
Peptoniphilus asaccharolyticus		2.04	0.32		99	99		
12010000	**Low discrimination**							**12010000**
					COC	GRA		
Fusobacterium varium		50.78	0.50	SUC 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		10.70	0.32		1	1		
Peptoniphilus asaccharolyticus		10.70	0.32		99	99		
12010002	**Low discrimination**							**12010002**
					GRA			
Atopobium minutum		48.07	0.36	IND 1	99			
Fusobacterium varium		19.30	0.16	SUC 5	1			
Porphyromonas asaccharolytica		16.60	0.08		1			
12010020	**Very good identification**							**12010020**
					GRA	SPO		
Clostridium cadaveris		98.40	0.59	GLU 95	99	99		
Atopobium minutum		0.59	0.31	SUC 7	99	1		
Fusobacterium varium		0.47	0.16		1	1		
12010100	**Low discrimination**							**12010100**
					GRA			
Actinomyces meyeri		72.75	0.38	IND 1	99			
Fusobacterium varium		14.46	0.16	GLU 93	1			
Fusobacterium nucleatum subsp. nucleatum		3.05	-.03		1			
12011100	**Acceptable identification**							**12011100**
Actinomyces meyeri		94.26	0.41	IND 1				
Eubacterium saburreum		2.49	0.21	GLU 93				
Lactobacillus fermentum		1.32	0.20					
12014100	**Good identification**							**12014100**
					GRA			
Actinomyces meyeri		99.58	0.43	IND 1	99			
Fusobacterium varium		0.43	-.08	GLU 93	1			
12015000	**Acceptable identification**							**12015000**
					GRA			
Actinomyces meyeri		94.28	0.35	IND 1	99			
Fusobacterium necrophorum subsp. necrophorum		2.48	-.03	GLU 93	1			
Eubacterium saburreum		1.18	0.11		85			
12015100	**Good identification**							**12015100**
Actinomyces meyeri		98.86	0.46	IND 1				
Eubacterium saburreum		0.72	0.17	GLU 93				
Lactobacillus fermentum		0.42	0.16					
12015500	**Good identification**							**12015500**
Actinomyces meyeri		98.33	0.35	IND 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		1.67	0.16	GLU 93				
12020020	**Acceptable identification**							**12020020**
					COC	GRA	SPO	
Clostridium cadaveris		92.82	0.37	GLU 95	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.79	-.03	SUC 7	1	1	1	
Peptoniphilus asaccharolyticus		1.79	-.03	TRE 5	99	99	1	
12030020	**Good identification**							**12030020**
Clostridium cadaveris		100.0	0.37	GLU 95				
12100001	**Good identification**							**12100001**
					COC	GRA	SPO	
Clostridium sordellii		96.40	0.44	GLU 95	1	99	99	
Anaerococcus prevotii		0.69	0.06	SUC 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		0.54	-.03		1	1	1	
12100020	**Low discrimination**							**12100020**
					GRA	SPO		
Clostridium cadaveris		86.85	0.37	GLU 95	99	99		
Clostridium bifermentans		6.14	0.19	SUC 7	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.67	-.03	MLT 5	1	1		
12110000	**Very good identification**							**12110000**
					COC	GRA		
Prevotella intermedia		96.46	0.63	GLU 90	1	1		
Megasphaera elsdenii		1.12	0.32		99	1		
Actinomyces meyeri		0.76	0.28		1	99		
12110001	**Good identification**							**12110001**
					GRA	SPO		
Clostridium sordellii		95.42	0.44	GLU 95	99	99		
Prevotella intermedia		4.58	0.28	SUC 1	1	1		
12110020	**Low discrimination**							**12110020**
					GRA	SPO		
Clostridium cadaveris		84.21	0.37	GLU 95	99	99		
Prevotella intermedia		13.99	0.28	SUC 7	1	1		
Actinomyces meyeri		1.49	0.14	MLT 5	99	1		
12110100	**Low discrimination**							**12110100**
					GRA			
Actinomyces meyeri		75.75	0.39	IND 1	99			
Prevotella intermedia		24.25	0.28	GLU 93	1			
12111000	**Very good identification**							**12111000**
					GRA			
Prevotella intermedia		98.45	0.63	GLU 90	1			
Actinomyces meyeri		1.17	0.31		99			
Eubacterium saburreum		0.19	0.25		85			
12111100	**Low discrimination**							**12111100**
					GRA			
Actinomyces meyeri		74.39	0.42	IND 1	99			
Prevotella intermedia		15.88	0.28	GLU 93	1			
Eubacterium saburreum		6.96	0.32		85			
12114100	**Good identification**							**12114100**
Actinomyces meyeri		100.0	0.44	IND 1				
12115000	**Low discrimination**							**12115000**
					GRA			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		66.14	0.36	IND 1	99		
Prevotella intermedia		29.09	0.28	GLU 93	1		
Eubacterium saburreum		2.94	0.21		85		
12115100	**Good identification**						**12115100**
Actinomyces meyeri		96.41	0.47	IND 1			
Eubacterium saburreum		2.50	0.27	GLU 93			
Lactobacillus fermentum		1.09	0.24				
12115500	**Good identification**						**12115500**
Actinomyces meyeri		95.68	0.36	IND 1			
Lactobacillus fermentum		4.32	0.24	GLU 93			
12510000	**Very good identification**						**12510000**
Prevotella intermedia		100.0	0.66	GLU 90			
12511000	**Very good identification**						**12511000**
					GRA		
Prevotella intermedia		99.82	0.66	GLU 90	1		
Lactobacillus fermentum		0.12	0.28		99		
Eubacterium saburreum		0.07	0.21		85		
13000000	**Low discrimination**						**13000000**
					GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		37.17	0.51	GLU 12	1	1	
Clostridium bifermentans		17.20	0.52	SUC 1	99	99	
Clostridium cadaveris		11.49	0.47		99	99	
13000001	**Acceptable identification**						**13000001**
					GRA	SPO	
Clostridium sordellii		94.57	0.50	SUC 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		1.54	0.17	MLT 90	1	1	
Clostridium novyi biovar A		0.90	0.32		99	99	
13000002	**Low discrimination**						**13000002**
					GRA		
Atopobium minutum		47.74	0.41	IND 1	99		
Fusobacterium necrophorum subsp. necrophorum		13.54	0.17	SUC 5	1		
Porphyromonas asaccharolytica		8.93	0.08		1		
13000020	**Very good identification**						**13000020**
					SPO		
Clostridium cadaveris		99.78	0.81	SUC 7	99		
Clostridium bifermentans		0.10	0.31		99		
Atopobium minutum		0.06	0.36		1		
13000021	**Good identification**						**13000021**
Clostridium cadaveris		97.47	0.47	SUC 7			
Clostridium sordellii		1.97	0.15	URE 1			
Clostridium novyi biovar A		0.46	0.21				
13000022	**Good identification**						**13000022**
					GRA	SPO	
Clostridium cadaveris		96.01	0.47	SUC 7	99	99	
Atopobium minutum		3.69	0.33	BGL 1	99	1	
Pseudoflavonifractor capillosus		0.22	-.03		1	1	
13000030	**Good identification**						**13000030**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium cadaveris		99.43	0.47	SUC 7			
Clostridium novyi biovar A		0.47	0.21	MLZ 1			
Clostridium bifermentans		0.10	-.03				
13000040	**Low discrimination**						**13000040**
					COC	GRA	
Megasphaera elsdenii		32.49	0.38	IND 1	99	1	
Clostridium botulinum biovar A		29.94	0.33	FRU 95	1	99	
Clostridium bifermentans		16.71	0.28		1	99	
13000060	**Good identification**						**13000060**
Clostridium cadaveris		99.06	0.47	SUC 7			
Clostridium bifermentans		0.39	0.08	SOR 1			
Clostridium novyi biovar A		0.32	0.18				
13000100	**Low discrimination**						**13000100**
					GRA	SPO	
Actinomyces meyeri		49.94	0.35	IND 1	99	1	
Fusobacterium necrophorum subsp. necrophorum		18.50	0.17	FRU 95	1	1	
Clostridium bifermentans		8.56	0.18		99	99	
13000120	**Good identification**						**13000120**
					SPO		
Clostridium cadaveris		98.72	0.47	SUC 7	99		
Actinomyces meyeri		1.19	0.20	LAC 1	1		
Clostridium bifermentans		0.10	-.03		99		
13000220	**Good identification**						**13000220**
Clostridium cadaveris		99.90	0.47	SUC 7			
Clostridium bifermentans		0.10	-.03	RHA 1			
13000420	**Good identification**						**13000420**
Clostridium cadaveris		99.90	0.47	SUC 7			
Clostridium bifermentans		0.10	-.03	ARA 1			
13001000	**Acceptable identification**						**13001000**
					GRA		
Fusobacterium necrophorum subsp. necrophorum		93.35	0.51	GLU 12	1		
Eubacterium saburreum		2.04	0.41	SUC 1	85		
Atopobium minutum		1.66	0.36		99		
13001001	**Acceptable identification**						**13001001**
					GRA	SPO	
Clostridium sordellii		91.04	0.37	SUC 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		8.41	0.17	MLT 90	1	1	
Clostridium novyi biovar A		0.55	0.15		99	99	
13001010	**Low discrimination**						**13001010**
					SPO	GRA	
Eubacterium saburreum		62.42	0.40	ESL 99	1	85	
Fusobacterium necrophorum subsp. necrophorum		35.29	0.17		1	1	
Clostridium novyi biovar A		2.29	0.15		99	99	
13001020	**Acceptable identification**						**13001020**
					GRA	SPO	
Clostridium cadaveris		93.23	0.47	SUC 7	99	99	
Fusobacterium necrophorum subsp. necrophorum		3.05	0.17	GAL 1	1	1	
Atopobium minutum		1.79	0.28		99	1	
13001100	**Low discrimination**						**13001100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Eubacterium saburreum		49.35	0.47	ESL 99	85			
Actinomyces meyeri		39.63	0.38		99			
Fusobacterium necrophorum subsp. necrophorum		9.79	0.17		1			
13001110	**Good identification**							**13001110**
Eubacterium saburreum		99.02	0.46	ESL 99				
Actinomyces meyeri		0.98	0.04					
13002000	**Low discrimination**							**13002000**
					COC	GRA	SPO	
Megasphaera elsdenii		68.10	0.41	IND 1	99	1	1	
Fusobacterium necrophorum subsp. necrophorum		12.24	0.17	FRU 95	1	1	1	
Clostridium bifermentans		5.66	0.18		1	99	99	
13002020	**Good identification**							**13002020**
					COC	SPO	CAT	
Clostridium cadaveris		99.57	0.47	SUC 7	1	99	1	
Megasphaera elsdenii		0.18	0.07	MAN 1	99	1	1	
Propionibacterium acnes		0.16	0.02		1	1	95	
13002040	**Good identification**							**13002040**
					COC	GRA		
Megasphaera elsdenii		99.08	0.38	IND 1	99	1		
Clostridium botulinum biovar A		0.92	-.01	FRU 95	1	99		
13004020	**Good identification**							**13004020**
					SPO			
Clostridium cadaveris		99.32	0.47	SUC 7	99			
Actinomyces meyeri		0.58	0.15	XYL 1	1			
Clostridium bifermentans		0.10	-.03		99			
13004100	**Good identification**							**13004100**
Actinomyces meyeri		100.0	0.40	IND 1				
13005010	**Good identification**							**13005010**
Eubacterium saburreum		100.0	0.35	ESL 99				
13005100	**Low discrimination**							**13005100**
Actinomyces meyeri		73.99	0.43	IND 1				
Eubacterium saburreum		25.56	0.43	FRU 95				
Lactobacillus fermentum		0.45	0.16					
13005110	**Good identification**							**13005110**
Eubacterium saburreum		96.55	0.41	ESL 99				
Actinomyces meyeri		3.45	0.09					
13010000	**Low discrimination**							**13010000**
					COC	GRA	SPO	
Fusobacterium varium		58.38	0.59	SUC 1	1	1	1	
Megasphaera elsdenii		24.27	0.63		99	1	1	
Clostridium cadaveris		7.03	0.47		1	99	99	
13010001	**Acceptable identification**							**13010001**
					COC	GRA	SPO	
Clostridium sordellii		93.33	0.50	SUC 1	1	99	99	
Fusobacterium varium		3.91	0.25	MLT 90	1	1	1	
Megasphaera elsdenii		1.63	0.29		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
13010002	**Low discrimination**							**13010002**
					COC	GRA		
Megasphaera elsdenii		60.57	0.47	IND 1	99	1		
Atopobium minutum		16.78	0.40	BGL 10	1	99		
Fusobacterium varium		13.25	0.25		1	1		
13010020	**Very good identification**							**13010020**
					GRA	SPO	CAT	
Clostridium cadaveris		99.44	0.81	SUC 7	99	99	1	
Propionibacterium acnes		0.28	0.40		99	1	95	
Fusobacterium varium		0.08	0.25		1	1	1	
13010021	**Good identification**							**13010021**
					SPO	CAT		
Clostridium cadaveris		97.75	0.47	SUC 7	99	1		
Clostridium sordellii		1.98	0.15	URE 1	99	1		
Propionibacterium acnes		0.27	0.06		1	95		
13010022	**Good identification**							**13010022**
					SPO			
Clostridium cadaveris		95.40	0.47	SUC 7	99			
Atopobium minutum		3.38	0.32	BGL 1	1			
Pseudoramibacter alactolyticus		0.41	0.03		1			
13010030	**Good identification**							**13010030**
					SPO	CAT		
Clostridium cadaveris		99.72	0.47	SUC 7	99	1		
Propionibacterium acnes		0.28	0.06	MLZ 1	1	95		
13010040	**Good identification**							**13010040**
					COC	GRA		
Megasphaera elsdenii		92.89	0.60	IND 1	99	1		
Fusobacterium varium		3.39	0.25		1	1		
Pseudoramibacter alactolyticus		2.23	0.21		1	99		
13010042	**Good identification**							**13010042**
					COC	GRA		
Megasphaera elsdenii		97.26	0.44	IND 1	99	1		
Pseudoramibacter alactolyticus		2.33	0.05	BGL 10	1	99		
Atopobium minutum		0.41	0.06		1	99		
13010060	**Acceptable identification**							**13010060**
					COC	SPO	CAT	
Clostridium cadaveris		91.43	0.47	SUC 7	1	99	1	
Propionibacterium acnes		5.55	0.29	SOR 1	1	1	95	
Megasphaera elsdenii		2.13	0.26		99	1	1	
13010100	**Low discrimination**							**13010100**
					GRA			
Actinomyces meyeri		87.45	0.57	IND 1	99			
Fusobacterium mortiferum		5.01	0.45		1			
Fusobacterium varium		4.38	0.25		1			
13010120	**Low discrimination**							**13010120**
					SPO	CAT		
Clostridium cadaveris		81.15	0.47	SUC 7	99	1		
Actinomyces meyeri		18.55	0.42	LAC 1	1	1		
Propionibacterium acnes		0.23	0.06		1	95		
13010220	**Good identification**							**13010220**
					SPO	CAT		
Clostridium cadaveris		99.72	0.47	SUC 7	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		0.28	0.06	RHA 1	1	95	
13010400	**Low discrimination**						**13010400**
					COC	GRA	
Actinomyces meyeri		42.84	0.36	IND 1	1	99	
Fusobacterium varium		34.32	0.25		1	1	
Megasphaera elsdenii		14.27	0.29		99	1	
13010420	**Good identification**						**13010420**
					SPO	CAT	
Clostridium cadaveris		98.32	0.47	SUC 7	99	1	
Actinomyces meyeri		1.40	0.22	ARA 1	1	1	
Propionibacterium acnes		0.28	0.06		1	95	
13010500	**Good identification**						**13010500**
					GRA		
Actinomyces meyeri		99.14	0.47	IND 1	99		
Lactobacillus fermentum		0.63	0.20		99		
Fusobacterium mortiferum		0.23	0.11		1		
13011000	**Low discrimination**						**13011000**
					GRA		
Actinomyces meyeri		36.78	0.50	IND 1	99		
Prevotella intermedia		23.99	0.45		1		
Eubacterium saburreum		12.40	0.49		85		
13011001	**Good identification**						**13011001**
					GRA	SPO	
Clostridium sordellii		96.13	0.37	SUC 1	99	99	
Actinomyces meyeri		1.73	0.16	MLT 90	99	1	
Prevotella intermedia		1.13	0.11		1	1	
13011002	**Low discrimination**						**13011002**
Lactobacillus fermentum		59.67	0.50	IND 1			
Eubacterium saburreum		13.86	0.36				
Atopobium minutum		13.15	0.32				
13011010	**Acceptable identification**						**13011010**
					GRA		
Eubacterium saburreum		93.47	0.48	ESL 99	85		
Actinomyces meyeri		3.42	0.16		99		
Prevotella intermedia		2.23	0.11		1		
13011020	**Low discrimination**						**13011020**
					SPO	CAT	
Clostridium cadaveris		67.71	0.47	SUC 7	99	1	
Propionibacterium acnes		18.73	0.40	GAL 1	1	95	
Actinomyces meyeri		5.80	0.35		1	1	
13011022	**Low discrimination**						**13011022**
					SPO		
Lactobacillus fermentum		61.23	0.42	IND 1	1		
Atopobium minutum		13.49	0.24		1		
Clostridium cadaveris		11.53	0.13		99		
13011100	**Low discrimination**						**13011100**
Actinomyces meyeri		73.53	0.60	IND 1			
Eubacterium saburreum		14.46	0.56				
Lactobacillus fermentum		7.68	0.54				
13011102	**Low discrimination**						**13011102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		59.20	0.54	IND 1		
Eubacterium saburreum		19.67	0.43			
Actinomyces meyeri		17.53	0.34			
13011110	**Acceptable identification**					**13011110**
Eubacterium saburreum		93.20	0.54	ESL 99		
Actinomyces meyeri		5.85	0.26			
Lactobacillus fermentum		0.61	0.20			
13011112	**Good identification**					**13011112**
Eubacterium saburreum		96.42	0.41	ESL 99		
Lactobacillus fermentum		3.58	0.20			
13011120	**Low discrimination**					**13011120**
Actinomyces meyeri		64.20	0.45	IND 1		
Lactobacillus fermentum		16.39	0.46	NAG 12		
Eubacterium saburreum		16.33	0.43			
13011122	**Low discrimination**					**13011122**
Lactobacillus fermentum		77.10	0.46	IND 1		
Eubacterium saburreum		13.56	0.30			
Actinomyces meyeri		9.34	0.19			
13011130	**Good identification**					**13011130**
Eubacterium saburreum		95.37	0.41	ESL 99		
Actinomyces meyeri		4.63	0.11			
13011400	**Low discrimination**					**13011400**
					GRA	
Lactobacillus fermentum		49.62	0.50	IND 1	99	
Actinomyces meyeri		48.45	0.39		99	
Prevotella intermedia		1.28	0.11		1	
13011402	**Good identification**					**13011402**
Lactobacillus fermentum		97.07	0.50	IND 1		
Actinomyces meyeri		2.93	0.14			
13011420	**Low discrimination**					**13011420**
					SPO	
Lactobacillus fermentum		60.98	0.42	IND 1	1	
Actinomyces meyeri		24.36	0.25		1	
Clostridium cadaveris		11.49	0.13		99	
13011422	**Good identification**					**13011422**
Lactobacillus fermentum		100.0	0.42	IND 1		
13011500	**Low discrimination**					**13011500**
Actinomyces meyeri		70.03	0.50	IND 1		
Lactobacillus fermentum		29.25	0.54			
Eubacterium saburreum		0.56	0.21			
13011502	**Acceptable identification**					**13011502**
Lactobacillus fermentum		93.11	0.54	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		6.89	0.24					
13011520	**Low discrimination**							**13011520**
					GRA			
Lactobacillus fermentum		49.78	0.46	IND 1	99			
Actinomyces meyeri		48.75	0.35		99			
Odoribacter splanchnicus		1.48	-.03		1			
13011522	**Good identification**							**13011522**
Lactobacillus fermentum		97.06	0.46	IND 1				
Actinomyces meyeri		2.94	0.09					
13012000	**Good identification**							**13012000**
					COC	GRA		
Megasphaera elsdenii		91.77	0.63	IND 1	99	1		
Pseudoramibacter alactolyticus		3.16	0.27		1	99		
Fusobacterium varium		2.23	0.25		1	1		
13012002	**Good identification**							**13012002**
					COC	CAT	GRA	
Megasphaera elsdenii		96.11	0.47	IND 1	99	1	1	
Pseudoramibacter alactolyticus		3.31	0.11	BGL 10	1	1	99	
Propionibacterium granulosum		0.31	0.09		1	90	99	
13012020	**Low discrimination**							**13012020**
					COC	SPO	CAT	
Clostridium cadaveris		82.46	0.47	SUC 7	1	99	1	
Propionibacterium acnes		12.83	0.36	MAN 1	1	1	95	
Megasphaera elsdenii		2.88	0.29		99	1	1	
13012040	**Very good identification**							**13012040**
					COC	GRA		
Megasphaera elsdenii		98.73	0.60	IND 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.17		1	99		
13012042	**Good identification**							**13012042**
					COC	GRA		
Megasphaera elsdenii		98.73	0.44	IND 1	99	1		
Pseudoramibacter alactolyticus		1.28	0.00	BGL 10	1	99		
13013020	**Acceptable identification**							**13013020**
					SPO	CAT		
Propionibacterium acnes		93.22	0.36	NIT 92	1	95		
Clostridium cadaveris		6.05	0.13	SUC 1	99	1		
Propionibacterium granulosum		0.73	0.09		1	90		
13014000	**Low discrimination**							**13014000**
					COC	GRA		
Actinomyces meyeri		66.44	0.52	IND 1	1	99		
Fusobacterium varium		28.28	0.35		1	1		
Megasphaera elsdenii		2.85	0.29		99	1		
13014020	**Low discrimination**							**13014020**
					GRA	SPO		
Clostridium cadaveris		89.51	0.47	SUC 7	99	99		
Actinomyces meyeri		9.93	0.37	XYL 1	99	1		
Fusobacterium varium		0.31	0.01		1	1		
13014100	**Very good identification**							**13014100**
					GRA			
Actinomyces meyeri		99.54	0.62	IND 1	99			
Fusobacterium mortiferum		0.22	0.26		1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium varium		0.11	0.01		1	
13014102	**Good identification**					**13014102**
Actinomyces meyeri		98.48	0.36	IND 1		
Lactobacillus fermentum		1.52	0.16	BGL 3		
13014120	**Good identification**					**13014120**
					SPO	
Actinomyces meyeri		97.77	0.47	IND 1	1	
Clostridium cadaveris		2.23	0.13	NAG 12	99	
13014400	**Good identification**					**13014400**
					GRA	
Actinomyces meyeri		97.86	0.41	IND 1	99	
Fusobacterium varium		1.68	0.01		1	
Lactobacillus fermentum		0.46	0.12		99	
13014500	**Good identification**					**13014500**
Actinomyces meyeri		99.81	0.52	IND 1		
Lactobacillus fermentum		0.19	0.16			
13014520	**Good identification**					**13014520**
Actinomyces meyeri		100.0	0.37	IND 1		
13015000	**Low discrimination**					**13015000**
Actinomyces meyeri		84.67	0.55	IND 1		
Eubacterium saburreum		7.92	0.45			
Lactobacillus fermentum		6.56	0.46			
13015002	**Low discrimination**					**13015002**
Lactobacillus fermentum		62.02	0.46	IND 1		
Actinomyces meyeri		24.77	0.29			
Eubacterium saburreum		13.21	0.32			
13015010	**Low discrimination**					**13015010**
Eubacterium saburreum		87.55	0.43	ESL 99		
Actinomyces meyeri		11.56	0.21			
Lactobacillus fermentum		0.90	0.12			
13015020	**Low discrimination**					**13015020**
Actinomyces meyeri		72.05	0.40	IND 1		
Lactobacillus fermentum		13.64	0.38	NAG 12		
Eubacterium saburreum		8.72	0.32			
13015022	**Low discrimination**					**13015022**
Lactobacillus fermentum		78.36	0.38	IND 1		
Actinomyces meyeri		12.80	0.14			
Eubacterium saburreum		8.84	0.19			
13015100	**Good identification**					**13015100**
Actinomyces meyeri		91.88	0.65	IND 1		
Eubacterium saburreum		5.01	0.51			
Lactobacillus fermentum		2.90	0.50			
13015102	**Low discrimination**					**13015102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		43.64	0.50	IND 1		
Actinomyces meyeri		42.72	0.39			
Eubacterium saburreum		13.29	0.38			
13015110	**Low discrimination**					**13015110**
Eubacterium saburreum		81.07	0.49	ESL 99		
Actinomyces meyeri		18.35	0.31			
Lactobacillus fermentum		0.58	0.16			
13015112	**Acceptable identification**					**13015112**
Eubacterium saburreum		92.58	0.37	ESL 99		
Lactobacillus fermentum		3.75	0.16			
Actinomyces meyeri		3.67	0.05			
13015120	**Low discrimination**					**13015120**
Actinomyces meyeri		87.12	0.50	IND 1		
Lactobacillus fermentum		6.73	0.42	NAG 12		
Eubacterium saburreum		6.15	0.38			
13015122	**Low discrimination**					**13015122**
Lactobacillus fermentum		64.03	0.42	IND 1		
Actinomyces meyeri		25.64	0.24			
Eubacterium saburreum		10.33	0.25			
13015130	**Low discrimination**					**13015130**
Eubacterium saburreum		85.11	0.37	ESL 99		
Actinomyces meyeri		14.89	0.16			
13015400	**Low discrimination**					**13015400**
Actinomyces meyeri		76.13	0.44	IND 1		
Lactobacillus fermentum		23.59	0.46			
Eubacterium saburreum		0.29	0.11			
13015402	**Acceptable identification**					**13015402**
Lactobacillus fermentum		90.92	0.46	IND 1		
Actinomyces meyeri		9.08	0.19			
13015422	**Good identification**					**13015422**
Lactobacillus fermentum		96.08	0.38	IND 1		
Actinomyces meyeri		3.92	0.04			
13015500	**Low discrimination**					**13015500**
Actinomyces meyeri		88.61	0.55	IND 1		
Lactobacillus fermentum		11.20	0.50			
Eubacterium saburreum		0.20	0.17			
13015502	**Low discrimination**					**13015502**
Lactobacillus fermentum		80.34	0.50	IND 1		
Actinomyces meyeri		19.66	0.29			
13015520	**Low discrimination**					**13015520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		76.40	0.40	IND 1			
Lactobacillus fermentum		23.60	0.42	NAG 12			
13015522	**Acceptable identification**						**13015522**
Lactobacillus fermentum		90.90	0.42	IND 1			
Actinomyces meyeri		9.10	0.14				
13020020	**Very good identification**						**13020020**
Clostridium cadaveris		99.98	0.59	SUC 7			
Clostridium bifermentans		0.02	-.03	TRE 5			
13021000	**Low discrimination**						**13021000**
Eubacterium saburreum		59.03	0.38	ESL 99			
Fusobacterium necrophorum subsp. necrophorum		40.97	0.17				
13021010	**Good identification**						**13021010**
Eubacterium saburreum		100.0	0.37	ESL 99			
13021100	**Good identification**						**13021100**
Eubacterium saburreum		98.80	0.44	ESL 99			
Actinomyces meyeri		1.20	0.04				
13021110	**Good identification**						**13021110**
Eubacterium saburreum		100.0	0.43	ESL 99			
13025100	**Good identification**						**13025100**
Eubacterium saburreum		95.80	0.40	ESL 99			
Actinomyces meyeri		4.20	0.09				
13025110	**Good identification**						**13025110**
Eubacterium saburreum		100.0	0.38	ESL 99			
13030020	**Very good identification**						**13030020**
					SPO	CAT	
Clostridium cadaveris		99.85	0.59	SUC 7	99	1	
Propionibacterium granulosum		0.09	0.20	TRE 5	1	90	
Propionibacterium acnes		0.05	0.06		1	95	
13030100	**Low discrimination**						**13030100**
					GRA		
Fusobacterium mortiferum		58.59	0.36	IND 1	1		
Actinomyces meyeri		34.61	0.23		99		
Eubacterium saburreum		6.81	0.18		85		
13031000	**Low discrimination**						**13031000**
					CAT	GRA	
Eubacterium saburreum		73.86	0.46	ESL 99	1	85	
Propionibacterium granulosum		10.17	0.30		90	99	
Fusobacterium mortiferum		9.63	0.33		5	1	
13031010	**Good identification**						**13031010**
					CAT		
Eubacterium saburreum		97.76	0.45	ESL 99	1		
Propionibacterium granulosum		2.24	0.15		90		
13031100	**Low discrimination**						**13031100**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		82.27	0.53	ESL 99	85	
Fusobacterium mortiferum		10.73	0.39		1	
Actinomyces meyeri		6.34	0.26		99	
13031102	**Low discrimination**					**13031102**
					GRA	
Eubacterium saburreum		88.67	0.40	ESL 99	85	
Fusobacterium mortiferum		7.28	0.23		1	
Lactobacillus fermentum		4.04	0.20		99	
13031110	**Good identification**					**13031110**
Eubacterium saburreum		100.0	0.51	ESL 99		
13031112	**Good identification**					**13031112**
Eubacterium saburreum		100.0	0.38	ESL 99		
13031120	**Acceptable identification**					**13031120**
Eubacterium saburreum		94.38	0.40	ESL 99		
Actinomyces meyeri		5.62	0.11			
13031130	**Good identification**					**13031130**
Eubacterium saburreum		100.0	0.38	ESL 99		
13035000	**Low discrimination**					**13035000**
					GRA	
Eubacterium saburreum		83.83	0.42	ESL 99	85	
Actinomyces meyeri		13.59	0.21		99	
Fusobacterium mortiferum		1.53	0.14		1	
13035010	**Good identification**					**13035010**
Eubacterium saburreum		100.0	0.40	ESL 99		
13035100	**Low discrimination**					**13035100**
					GRA	
Eubacterium saburreum		76.64	0.48	ESL 99	85	
Actinomyces meyeri		21.29	0.31		99	
Fusobacterium mortiferum		1.40	0.20		1	
13035102	**Acceptable identification**					**13035102**
Eubacterium saburreum		91.04	0.35	ESL 99		
Lactobacillus fermentum		4.53	0.16			
Actinomyces meyeri		4.43	0.05			
13035110	**Good identification**					**13035110**
Eubacterium saburreum		100.0	0.46	ESL 99		
13035120	**Low discrimination**					**13035120**
Eubacterium saburreum		82.33	0.35	ESL 99		
Actinomyces meyeri		17.67	0.16			
13040020	**Good identification**					**13040020**
Clostridium cadaveris		99.90	0.47	SUC 7		
Clostridium bifermentans		0.10	-.03	CEL 1		
13050020	**Good identification**					**13050020**
					SPO CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium cadaveris		99.72	0.47	SUC 7	99	1	
Propionibacterium acnes		0.28	0.06	CEL 1	1	95	
13050100	**Low discrimination**						**13050100**
					GRA		
Fusobacterium mortiferum		65.39	0.37	IND 1	1		
Actinomyces meyeri		34.61	0.23		99		
13051100	**Low discrimination**						**13051100**
					GRA		
Fusobacterium mortiferum		59.22	0.40	IND 1	1		
Actinomyces meyeri		31.35	0.26		99		
Eubacterium saburreum		6.16	0.21		85		
13100000	**Low discrimination**						**13100000**
					GRA	SPO	
Clostridium bifermentans		68.87	0.58	SUC 1	99	99	
Prevotella intermedia		8.09	0.45		1	1	
Clostridium botulinum biovar A		6.04	0.41		99	99	
13100001	**Very good identification**						**13100001**
Clostridium sordellii		99.64	0.66	SUC 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
13100020	**Very good identification**						**13100020**
					COC	SPO	
Clostridium cadaveris		95.43	0.59	SUC 7	1	99	
Clostridium bifermentans		4.08	0.38	MLT 5	1	99	
Gemella morbillorum		0.11	0.14		99	1	
13100040	**Low discrimination**						**13100040**
					COC	GRA	
Megasphaera elsdenii		39.91	0.41	IND 1	99	1	
Clostridium bifermentans		31.93	0.34	FRU 95	1	99	
Clostridium botulinum biovar A		20.06	0.32		1	99	
13100041	**Good identification**						**13100041**
					COC	GRA	
Clostridium sordellii		97.91	0.42	SUC 1	1	99	
Clostridium novyi biovar A		1.66	0.29	SOR 4	1	99	
Megasphaera elsdenii		0.24	0.07		99	1	
13100100	**Low discrimination**						**13100100**
					SPO		
Actinomyces meyeri		64.43	0.36	IND 1	1		
Clostridium bifermentans		22.84	0.24	FRU 95	99		
Eubacterium saburreum		4.31	0.24		1		
13101000	**Low discrimination**						**13101000**
					GRA		
Prevotella intermedia		47.03	0.45	FRU 99	1		
Eubacterium saburreum		32.40	0.51		85		
Actinomyces meyeri		4.28	0.29		99		
13101001	**Good identification**						**13101001**
					SPO	GRA	
Clostridium sordellii		99.67	0.53	SUC 1	99	99	
Prevotella intermedia		0.13	0.11		1	1	
Eubacterium saburreum		0.09	0.17		1	85	
13101002	**Low discrimination**						**13101002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Eubacterium saburreum		83.05	0.39	ESL 99	85			
Lactobacillus fermentum		8.13	0.24		99			
Prevotella intermedia		6.90	0.11		1			
13101010	**Good identification**							**13101010**
					SPO	GRA		
Eubacterium saburreum		97.36	0.50	ESL 99	1	85		
Prevotella intermedia		1.75	0.11		1	1		
Clostridium novyi biovar A		0.89	0.15		99	99		
13101012	**Good identification**							**13101012**
Eubacterium saburreum		100.0	0.37	ESL 99				
13101030	**Good identification**							**13101030**
Eubacterium saburreum		100.0	0.37	ESL 99				
13101100	**Low discrimination**							**13101100**
Eubacterium saburreum		80.02	0.58	ESL 99				
Actinomyces meyeri		18.12	0.39					
Lactobacillus fermentum		0.97	0.28					
13101101	**Low discrimination**							**13101101**
					SPO			
Clostridium chauvoei		79.48	0.37	IND 1	99			
Clostridium sordellii		16.37	0.19	MNS 90	99			
Eubacterium saburreum		3.39	0.24		1			
13101102	**Acceptable identification**							**13101102**
Eubacterium saburreum		90.25	0.45	ESL 99				
Lactobacillus fermentum		6.17	0.28					
Actinomyces meyeri		3.58	0.13					
13101110	**Very good identification**							**13101110**
Eubacterium saburreum		99.72	0.56	ESL 99				
Actinomyces meyeri		0.28	0.05					
13101112	**Good identification**							**13101112**
Eubacterium saburreum		100.0	0.43	ESL 99				
13101120	**Low discrimination**							**13101120**
					GRA			
Eubacterium saburreum		71.60	0.45	ESL 99	85			
Prevotella melaninogenica		14.25	0.17		1			
Actinomyces meyeri		12.53	0.24		99			
13101130	**Good identification**							**13101130**
Eubacterium saburreum		100.0	0.43	ESL 99				
13102000	**Low discrimination**							**13102000**
					COC	GRA	SPO	
Megasphaera elsdenii		85.70	0.44	IND 1	99	1	1	
Clostridium bifermentans		11.08	0.24	FRU 95	1	99	99	
Prevotella intermedia		1.30	0.11		1	1	1	
13102040	**Good identification**							**13102040**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		99.20	0.41	IND 1	99	1		
Clostridium bifermentans		0.80	0.00	FRU 95	1	99		
13104100	**Good identification**							**13104100**
Actinomyces meyeri		98.18	0.41	IND 1				
Eubacterium saburreum		1.82	0.19	FRU 95				
13105000	**Low discrimination**							**13105000**
					GRA			
Eubacterium saburreum		64.83	0.47	ESL 99	85			
Actinomyces meyeri		30.87	0.34		99			
Prevotella intermedia		1.77	0.11		1			
13105010	**Good identification**							**13105010**
Eubacterium saburreum		100.0	0.45	ESL 99				
13105100	**Low discrimination**							**13105100**
Eubacterium saburreum		54.66	0.53	ESL 99				
Actinomyces meyeri		44.62	0.44					
Lactobacillus fermentum		0.72	0.24					
13105102	**Low discrimination**							**13105102**
Eubacterium saburreum		82.13	0.40	ESL 99				
Actinomyces meyeri		11.75	0.18					
Lactobacillus fermentum		6.13	0.24					
13105110	**Good identification**							**13105110**
Eubacterium saburreum		99.00	0.52	ESL 99				
Actinomyces meyeri		1.00	0.10					
13105112	**Good identification**							**13105112**
Eubacterium saburreum		100.0	0.39	ESL 99				
13105120	**Low discrimination**							**13105120**
Eubacterium saburreum		60.40	0.40	ESL 99				
Actinomyces meyeri		38.10	0.29					
Lactobacillus fermentum		1.50	0.16					
13105130	**Good identification**							**13105130**
Eubacterium saburreum		100.0	0.39	ESL 99				
13110000	**Low discrimination**							**13110000**
					COC	GRA		
Prevotella intermedia		86.48	0.79		1	1		
Megasphaera elsdenii		11.04	0.66		99	1		
Actinomyces meyeri		1.01	0.48		1	99		
13110001	**Very good identification**							**13110001**
					COC	GRA	SPO	
Clostridium sordellii		97.45	0.66	SUC 1	1	99	99	
Prevotella intermedia		2.22	0.45		1	1	1	
Megasphaera elsdenii		0.28	0.32		99	1	1	
13110002	**Low discrimination**							**13110002**
					COC	GRA		
Megasphaera elsdenii		56.54	0.50	IND 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		40.26	0.45	BGL 10	1	1		
Actinomyces meyeri		1.44	0.22		1	99		
13110010	**Low discrimination**							**13110010**
					COC			
Prevotella intermedia		85.43	0.45	MLZ 1	1			
Megasphaera elsdenii		10.91	0.32		99			
Eubacterium saburreum		1.46	0.24		1			
13110020	**Good identification**							**13110020**
					GRA	SPO		
Clostridium cadaveris		90.38	0.59	SUC 7	99	99		
Prevotella intermedia		7.11	0.45	MLT 5	1	1		
Actinomyces meyeri		1.12	0.33		99	1		
13110040	**Low discrimination**							**13110040**
					COC	GRA	SPO	
Megasphaera elsdenii		87.62	0.63	IND 1	99	1	1	
Prevotella intermedia		10.40	0.45		1	1	1	
Clostridium botulinum biovar B		1.15	0.24		1	99	99	
13110041	**Acceptable identification**							**13110041**
					COC	GRA	SPO	
Clostridium sordellii		94.89	0.42	SUC 1	1	99	99	
Megasphaera elsdenii		4.41	0.29	SOR 4	99	1	1	
Prevotella intermedia		0.52	0.11		1	1	1	
13110042	**Good identification**							**13110042**
					COC			
Megasphaera elsdenii		98.93	0.47	IND 1	99			
Prevotella intermedia		1.07	0.11	BGL 10	1			
13110100	**Low discrimination**							**13110100**
					COC	GRA		
Actinomyces meyeri		78.75	0.58	IND 1	1	99		
Prevotella intermedia		17.08	0.45		1	1		
Megasphaera elsdenii		2.18	0.32		99	1		
13110120	**Low discrimination**							**13110120**
					GRA	SPO		
Actinomyces meyeri		79.85	0.43	IND 1	99	1		
Clostridium cadaveris		16.31	0.25	NAG 12	99	99		
Prevotella intermedia		1.28	0.11		1	1		
13110200	**Low discrimination**							**13110200**
					COC	GRA		
Prevotella intermedia		83.96	0.45	RHA 1	1	1		
Megasphaera elsdenii		10.72	0.32		99	1		
Actinomyces odontolyticus		3.59	0.27		1	99		
13110400	**Low discrimination**							**13110400**
					COC	GRA		
Prevotella intermedia		68.78	0.45	ARA 1	1	1		
Actinomyces meyeri		19.82	0.37		1	99		
Megasphaera elsdenii		8.78	0.32		99	1		
13110500	**Good identification**							**13110500**
					GRA			
Actinomyces meyeri		97.50	0.48	IND 1	99			
Lactobacillus fermentum		1.64	0.28		99			
Prevotella intermedia		0.86	0.11		1			
13111000	**Good identification**							**13111000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Prevotella intermedia		94.91	0.79		1		
Eubacterium saburreum		1.98	0.60		85		
Actinomyces meyeri		1.66	0.51		99		
13111001	**Low discrimination**						**13111001**
					SPO	GRA	
Clostridium sordellii		87.99	0.53	SUC 1	99	99	
Prevotella intermedia		11.34	0.45		1	1	
Eubacterium saburreum		0.24	0.25		1	85	
13111002	**Low discrimination**						**13111002**
					GRA		
Lactobacillus fermentum		44.96	0.59	IND 1	99		
Prevotella intermedia		38.19	0.45		1		
Eubacterium saburreum		13.93	0.47		85		
13111010	**Low discrimination**						**13111010**
					GRA		
Eubacterium saburreum		62.16	0.58	ESL 99	85		
Prevotella intermedia		36.76	0.45		1		
Actinomyces meyeri		0.64	0.16		99		
13111012	**Acceptable identification**						**13111012**
					GRA		
Eubacterium saburreum		93.14	0.45	ESL 99	85		
Lactobacillus fermentum		3.71	0.24		99		
Prevotella intermedia		3.15	0.11		1		
13111020	**Low discrimination**						**13111020**
					GRA		
Prevotella intermedia		46.36	0.45	NAG 1	1		
Lactobacillus fermentum		18.20	0.50		99		
Eubacterium saburreum		16.91	0.47		85		
13111022	**Low discrimination**						**13111022**
					GRA		
Lactobacillus fermentum		81.78	0.50	IND 1	99		
Eubacterium saburreum		13.41	0.34		85		
Prevotella intermedia		2.11	0.11		1		
13111030	**Acceptable identification**						**13111030**
					GRA		
Eubacterium saburreum		94.78	0.45	ESL 99	85		
Prevotella intermedia		3.21	0.11		1		
Lactobacillus fermentum		1.26	0.16		99		
13111040	**Low discrimination**						**13111040**
					COC		
Prevotella intermedia		87.83	0.45	SOR 1	1		
Megasphaera elsdenii		7.48	0.29		99		
Eubacterium saburreum		1.83	0.25		1		
13111100	**Low discrimination**						**13111100**
Actinomyces meyeri		46.95	0.61	IND 1			
Eubacterium saburreum		32.75	0.66				
Lactobacillus fermentum		13.05	0.62				
13111101	**Low discrimination**						**13111101**
					SPO		
Clostridium chauvoei		79.69	0.44	IND 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sordellii		7.03	0.19	MNS 90	99	
Actinomyces meyeri		6.27	0.27		1	
13111102	**Low discrimination**					**13111102**
Lactobacillus fermentum		63.84	0.62	IND 1		
Eubacterium saburreum		28.28	0.53			
Actinomyces meyeri		7.11	0.35			
13111110	**Very good identification**					**13111110**
Eubacterium saburreum		97.54	0.64	ESL 99		
Actinomyces meyeri		1.73	0.27			
Lactobacillus fermentum		0.48	0.28			
13111112	**Good identification**					**13111112**
Eubacterium saburreum		97.29	0.52	ESL 99		
Lactobacillus fermentum		2.71	0.28			
13111120	**Low discrimination**					**13111120**
Actinomyces meyeri		30.98	0.46	IND 1		
Eubacterium saburreum		27.96	0.53	NAG 12		
Lactobacillus fermentum		21.04	0.54			
13111121	**Acceptable identification**					**13111121**
					SPO	
Clostridium chauvoei		90.88	0.35	IND 1	99	
Actinomyces meyeri		2.92	0.12	MNS 90	1	
Eubacterium saburreum		2.64	0.19		1	
13111122	**Low discrimination**					**13111122**
					GRA	
Lactobacillus fermentum		66.84	0.54	IND 1	99	
Eubacterium saburreum		15.67	0.40		85	
Prevotella melaninogenica		10.10	0.21		1	
13111130	**Good identification**					**13111130**
Eubacterium saburreum		97.06	0.52	ESL 99		
Actinomyces meyeri		1.33	0.12			
Lactobacillus fermentum		0.90	0.20			
13111132	**Acceptable identification**					**13111132**
Eubacterium saburreum		95.00	0.39	ESL 99		
Lactobacillus fermentum		5.00	0.20			
13111200	**Acceptable identification**					**13111200**
					GRA	
Prevotella intermedia		92.70	0.45	RHA 1	1	
Actinomyces odontolyticus		2.64	0.24		99	
Eubacterium saburreum		1.94	0.25		85	
13111400	**Low discrimination**					**13111400**
					GRA	
Lactobacillus fermentum		44.61	0.59	IND 1	99	
Prevotella intermedia		37.88	0.45		1	
Actinomyces meyeri		16.37	0.40		99	
13111402	**Very good identification**					**13111402**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		97.75	0.59	IND 1	99		
Actinomyces meyeri		1.11	0.15		99		
Prevotella intermedia		0.84	0.11		1		
13111420	**Low discrimination**						**13111420**
					GRA		
Lactobacillus fermentum		84.37	0.50	IND 1	99		
Actinomyces meyeri		12.67	0.26		99		
Prevotella intermedia		2.17	0.11		1		
13111422	**Good identification**						**13111422**
Lactobacillus fermentum		100.0	0.50	IND 1			
13111500	**Low discrimination**						**13111500**
Lactobacillus fermentum		51.80	0.62	IND 1			
Actinomyces meyeri		46.61	0.51				
Eubacterium saburreum		1.31	0.32				
13111502	**Very good identification**						**13111502**
Lactobacillus fermentum		96.87	0.62	IND 1			
Actinomyces meyeri		2.70	0.25				
Eubacterium saburreum		0.43	0.19				
13111520	**Low discrimination**						**13111520**
Lactobacillus fermentum		71.38	0.54	IND 1			
Actinomyces meyeri		26.28	0.36				
Eubacterium saburreum		0.96	0.19				
13111522	**Good identification**						**13111522**
Lactobacillus fermentum		97.77	0.54	IND 1			
Bifidobacterium longum subsp. longum		1.12	0.14				
Actinomyces meyeri		1.11	0.10				
13112000	**Good identification**						**13112000**
					COC	GRA	
Megasphaera elsdenii		92.45	0.66	IND 1	99	1	
Prevotella intermedia		7.32	0.45		1	1	
Actinomyces meyeri		0.09	0.13		1	99	
13112002	**Good identification**						**13112002**
					COC	GRA	
Megasphaera elsdenii		98.92	0.50	IND 1	99	1	
Prevotella intermedia		0.71	0.11	BGL 10	1	1	
Propionibacterium propionicum		0.37	0.18		1	99	
13112040	**Very good identification**						**13112040**
					COC	GRA	
Megasphaera elsdenii		99.84	0.63	IND 1	99	1	
Prevotella intermedia		0.12	0.11		1	1	
Propionibacterium propionicum		0.04	0.15		1	99	
13112042	**Good identification**						**13112042**
					COC	GRA	
Megasphaera elsdenii		99.44	0.47	IND 1	99	1	
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99	
13112100	**Low discrimination**						**13112100**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		65.40	0.49	IND 1	1	99	
Megasphaera elsdenii		23.96	0.32	NIT 86	99	1	
Actinomyces meyeri		8.74	0.24		1	99	
13112102	**Good identification**						**13112102**
					COC	GRA	
Propionibacterium propionicum		97.36	0.52	IND 1	1	99	
Megasphaera elsdenii		2.64	0.15	NIT 86	99	1	
13112103	**Good identification**						**13112103**
Propionibacterium propionicum		100.0	0.36	IND 1			
13112140	**Low discrimination**						**13112140**
					COC	GRA	
Propionibacterium propionicum		80.37	0.49	IND 1	1	99	
Megasphaera elsdenii		19.63	0.29	NIT 86	99	1	
13112142	**Good identification**						**13112142**
					COC	GRA	
Propionibacterium propionicum		98.22	0.52	IND 1	1	99	
Megasphaera elsdenii		1.78	0.12	NIT 86	99	1	
13112143	**Good identification**						**13112143**
Propionibacterium propionicum		100.0	0.36	IND 1			
13113000	**Low discrimination**						**13113000**
					COC		
Prevotella intermedia		84.26	0.45	MAN 1	1		
Megasphaera elsdenii		10.76	0.32		99		
Eubacterium saburreum		1.76	0.25		1		
13113100	**Low discrimination**						**13113100**
Propionibacterium propionicum		70.18	0.49	IND 1			
Actinomyces meyeri		14.07	0.27	NIT 86			
Eubacterium saburreum		9.81	0.32				
13113102	**Acceptable identification**						**13113102**
Propionibacterium propionicum		94.91	0.52	IND 1			
Lactobacillus fermentum		3.52	0.28	NIT 86			
Eubacterium saburreum		1.56	0.19				
13113103	**Good identification**						**13113103**
Propionibacterium propionicum		100.0	0.36	IND 1			
13113140	**Good identification**						**13113140**
Propionibacterium propionicum		100.0	0.49	IND 1			
13113142	**Good identification**						**13113142**
Propionibacterium propionicum		100.0	0.52	IND 1			
13113143	**Good identification**						**13113143**
Propionibacterium propionicum		100.0	0.36	IND 1			
13114000	**Low discrimination**						**13114000**
					COC	GRA	
Actinomyces meyeri		64.99	0.53	IND 1	1	99	
Prevotella intermedia		29.05	0.45		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		3.71	0.32		99	1	
13114020	**Low discrimination**						**13114020**
					GRA	SPO	
Actinomyces meyeri		68.42	0.38	IND 1	99	1	
Clostridium cadaveris		28.79	0.25	NAG 12	99	99	
Prevotella intermedia		2.27	0.11		1	1	
13114100	**Very good identification**						**13114100**
Actinomyces meyeri		99.43	0.63	IND 1			
Eubacterium saburreum		0.29	0.27				
Lactobacillus fermentum		0.13	0.24				
13114102	**Acceptable identification**						**13114102**
Actinomyces meyeri		94.53	0.37	IND 1			
Lactobacillus fermentum		3.89	0.24	BGL 3			
Eubacterium saburreum		1.58	0.14				
13114120	**Good identification**						**13114120**
Actinomyces meyeri		99.31	0.48	IND 1			
Eubacterium saburreum		0.38	0.14	NAG 12			
Lactobacillus fermentum		0.31	0.16				
13114400	**Good identification**						**13114400**
					GRA		
Actinomyces meyeri		97.04	0.42	IND 1	99		
Prevotella intermedia		1.75	0.11		1		
Lactobacillus fermentum		1.21	0.20		99		
13114500	**Good identification**						**13114500**
Actinomyces meyeri		99.49	0.53	IND 1			
Lactobacillus fermentum		0.51	0.24				
13114520	**Good identification**						**13114520**
Actinomyces meyeri		98.77	0.38	IND 1			
Lactobacillus fermentum		1.23	0.16	NAG 12			
13115000	**Low discrimination**						**13115000**
					GRA		
Actinomyces meyeri		54.33	0.56	IND 1	99		
Eubacterium saburreum		18.02	0.55		85		
Prevotella intermedia		16.19	0.45		1		
13115002	**Low discrimination**						**13115002**
Lactobacillus fermentum		68.72	0.55	IND 1			
Eubacterium saburreum		19.52	0.42				
Actinomyces meyeri		10.31	0.30				
13115010	**Acceptable identification**						**13115010**
					GRA		
Eubacterium saburreum		94.70	0.53	ESL 99	85		
Actinomyces meyeri		3.53	0.21		99		
Prevotella intermedia		1.05	0.11		1		
13115012	**Good identification**						**13115012**
Eubacterium saburreum		95.83	0.41	ESL 99			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		4.17	0.20			
13115020	**Low discrimination**					**13115020**
Actinomyces meyeri		51.15	0.41	IND 1		
Lactobacillus fermentum		25.77	0.46	NAG 12		
Eubacterium saburreum		21.95	0.42			
13115022	**Low discrimination**					**13115022**
Lactobacillus fermentum		82.53	0.46	IND 1		
Eubacterium saburreum		12.41	0.29			
Actinomyces meyeri		5.07	0.15			
13115030	**Good identification**					**13115030**
Eubacterium saburreum		95.85	0.41	ESL 99		
Actinomyces meyeri		2.76	0.07			
Lactobacillus fermentum		1.39	0.12			
13115100	**Low discrimination**					**13115100**
Actinomyces meyeri		78.22	0.66	IND 1		
Eubacterium saburreum		15.13	0.61			
Lactobacillus fermentum		6.57	0.58			
13115102	**Low discrimination**					**13115102**
Lactobacillus fermentum		56.37	0.58	IND 1		
Eubacterium saburreum		22.89	0.48			
Actinomyces meyeri		20.74	0.40			
13115110	**Good identification**					**13115110**
Eubacterium saburreum		93.53	0.60	ESL 99		
Actinomyces meyeri		5.97	0.32			
Lactobacillus fermentum		0.50	0.24			
13115112	**Good identification**					**13115112**
Eubacterium saburreum		96.01	0.47	ESL 99		
Lactobacillus fermentum		2.92	0.24			
Actinomyces meyeri		1.07	0.06			
13115120	**Low discrimination**					**13115120**
Actinomyces meyeri		68.56	0.51	IND 1		
Eubacterium saburreum		17.16	0.48	NAG 12		
Lactobacillus fermentum		14.09	0.50			
13115122	**Low discrimination**					**13115122**
Lactobacillus fermentum		73.23	0.50	IND 1		
Eubacterium saburreum		15.75	0.35			
Actinomyces meyeri		11.02	0.25			
13115130	**Acceptable identification**					**13115130**
Eubacterium saburreum		94.39	0.47	ESL 99		
Actinomyces meyeri		4.66	0.17			
Lactobacillus fermentum		0.96	0.16			
13115400	**Low discrimination**					**13115400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		54.06	0.45	IND 1		
Lactobacillus fermentum		44.56	0.55			
Eubacterium saburreum		0.72	0.21			
13115402	**Good identification**					**13115402**
Lactobacillus fermentum		96.38	0.55	IND 1		
Actinomyces meyeri		3.62	0.19			
13115420	**Low discrimination**					**13115420**
Lactobacillus fermentum		66.83	0.46	IND 1		
Actinomyces meyeri		33.17	0.30			
13115422	**Good identification**					**13115422**
Lactobacillus fermentum		98.49	0.46	IND 1		
Actinomyces meyeri		1.51	0.05			
13115500	**Low discrimination**					**13115500**
Actinomyces meyeri		74.31	0.56	IND 1		
Lactobacillus fermentum		24.99	0.58			
Eubacterium saburreum		0.58	0.27			
13115502	**Good identification**					**13115502**
Lactobacillus fermentum		90.83	0.58	IND 1		
Actinomyces meyeri		8.36	0.30			
Bifidobacterium longum subsp. longum		0.44	0.12			
13115520	**Low discrimination**					**13115520**
Actinomyces meyeri		53.19	0.41	IND 1		
Lactobacillus fermentum		43.72	0.50	NAG 12		
Bifidobacterium longum subsp. longum		2.56	0.22			
13115522	**Acceptable identification**					**13115522**
Lactobacillus fermentum		91.23	0.50	IND 1		
Bifidobacterium longum subsp. longum		5.33	0.22			
Actinomyces meyeri		3.43	0.15			
13120020	**Good identification**					**13120020**
Clostridium cadaveris		99.19	0.37	SUC 7		
Clostridium bifermentans		0.81	0.03	MLT 5		
13121000	**Good identification**					**13121000**
Eubacterium saburreum		97.85	0.48	ESL 99		
Prevotella intermedia		2.15	0.11			
13121002	**Good identification**					**13121002**
Eubacterium saburreum		100.0	0.36	ESL 99		
13121010	**Good identification**					**13121010**
Eubacterium saburreum		100.0	0.47	ESL 99		
13121020	**Acceptable identification**					**13121020**

SPO

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium saburreum		92.30	0.36	ESL 99	1			
Clostridium cadaveris		7.70	0.03		99			
13121100	**Good identification**							**13121100**
Eubacterium saburreum		99.66	0.55	ESL 99				
Actinomyces meyeri		0.34	0.05					
13121102	**Good identification**							**13121102**
Eubacterium saburreum		100.0	0.42	ESL 99				
13121110	**Good identification**							**13121110**
Eubacterium saburreum		100.0	0.53	ESL 99				
13121112	**Good identification**							**13121112**
Eubacterium saburreum		100.0	0.40	ESL 99				
13121120	**Good identification**							**13121120**
Eubacterium saburreum		100.0	0.42	ESL 99				
13121130	**Good identification**							**13121130**
Eubacterium saburreum		100.0	0.40	ESL 99				
13125000	**Good identification**							**13125000**
Eubacterium saburreum		100.0	0.44	ESL 99				
13125010	**Good identification**							**13125010**
Eubacterium saburreum		100.0	0.42	ESL 99				
13125100	**Good identification**							**13125100**
Eubacterium saburreum		98.78	0.50	ESL 99				
Actinomyces meyeri		1.22	0.10					
13125102	**Good identification**							**13125102**
Eubacterium saburreum		100.0	0.37	ESL 99				
13125110	**Good identification**							**13125110**
Eubacterium saburreum		100.0	0.49	ESL 99				
13125112	**Good identification**							**13125112**
Eubacterium saburreum		100.0	0.36	ESL 99				
13125120	**Good identification**							**13125120**
Eubacterium saburreum		100.0	0.37	ESL 99				
13125130	**Good identification**							**13125130**
Eubacterium saburreum		100.0	0.36	ESL 99				
13130000	**Low discrimination**							**13130000**
					COC	GRA	CAT	
Prevotella intermedia		82.47	0.45	TRE 1	1	1	1	
Megasphaera elsdenii		10.53	0.32		99	1	1	
Propionibacterium granulosum		1.76	0.24		1	99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
13130020	**Good identification**							**13130020**
					GRA	SPO	CAT	
Clostridium cadaveris		97.40	0.37	SUC 7	99	99	1	
Prevotella intermedia		1.47	0.11	MLT 5	1	1	1	
Propionibacterium granulosum		0.78	0.14	TRE 5	99	1	90	
13131000	**Low discrimination**							**13131000**
					CAT	GRA		
Eubacterium saburreum		56.61	0.57	ESL 99	1	85		
Prevotella intermedia		41.08	0.45		1	1		
Propionibacterium granulosum		0.88	0.24		90	99		
13131002	**Low discrimination**							**13131002**
					GRA			
Eubacterium saburreum		89.90	0.44	ESL 99	85			
Lactobacillus fermentum		4.40	0.24		99			
Prevotella intermedia		3.74	0.11		1			
13131010	**Good identification**							**13131010**
					CAT	GRA		
Eubacterium saburreum		98.86	0.55	ESL 99	1	85		
Prevotella intermedia		0.89	0.11		1	1		
Propionibacterium granulosum		0.26	0.09		90	99		
13131012	**Good identification**							**13131012**
Eubacterium saburreum		100.0	0.42	ESL 99				
13131020	**Low discrimination**							**13131020**
					GRA			
Eubacterium saburreum		82.67	0.44	ESL 99	85			
Clostridium perfringens		7.63	0.17		99			
Prevotella intermedia		3.43	0.11		1			
13131030	**Good identification**							**13131030**
Eubacterium saburreum		100.0	0.42	ESL 99				
13131100	**Very good identification**							**13131100**
Eubacterium saburreum		96.04	0.63	ESL 99				
Actinomyces meyeri		2.09	0.27					
Lactobacillus fermentum		0.58	0.28					
13131102	**Good identification**							**13131102**
Eubacterium saburreum		95.60	0.50	ESL 99				
Lactobacillus fermentum		3.27	0.28					
Propionibacterium propionicum		0.73	0.17					
13131110	**Very good identification**							**13131110**
Eubacterium saburreum		100.0	0.61	ESL 99				
13131112	**Good identification**							**13131112**
Eubacterium saburreum		100.0	0.48	ESL 99				
13131120	**Low discrimination**							**13131120**
Eubacterium saburreum		75.35	0.50	ESL 99				
Clostridium perfringens		21.84	0.32					
Actinomyces meyeri		1.27	0.12					

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
13131122	**Low discrimination**					**13131122**
Eubacterium saburreum		54.42	0.37	ESL 99		
Clostridium perfringens		42.06	0.26			
Lactobacillus fermentum		3.52	0.20			
13131130	**Good identification**					**13131130**
Eubacterium saburreum		100.0	0.48	ESL 99		
13131132	**Good identification**					**13131132**
Eubacterium saburreum		100.0	0.36	ESL 99		
13132100	**Good identification**					**13132100**
Propionibacterium propionicum		100.0	0.48	IND 1		
13132102	**Good identification**					**13132102**
Propionibacterium propionicum		100.0	0.51	IND 1		
13132140	**Good identification**					**13132140**
Propionibacterium propionicum		100.0	0.48	IND 1		
13132142	**Good identification**					**13132142**
Propionibacterium propionicum		100.0	0.51	IND 1		
13133100	**Low discrimination**					**13133100**
Propionibacterium propionicum		89.77	0.48	IND 1		
Eubacterium saburreum		10.23	0.29	NIT 86		
13133102	**Good identification**					**13133102**
Propionibacterium propionicum		98.68	0.51	IND 1		
Eubacterium saburreum		1.32	0.16	NIT 86		
13133140	**Good identification**					**13133140**
Propionibacterium propionicum		100.0	0.48	IND 1		
13133142	**Good identification**					**13133142**
Propionibacterium propionicum		100.0	0.51	IND 1		
13135000	**Acceptable identification**					**13135000**
					GRA	
Eubacterium saburreum		93.57	0.52	ESL 99	85	
Actinomyces meyeri		4.28	0.21		99	
Prevotella intermedia		1.27	0.11		1	
13135002	**Acceptable identification**					**13135002**
Eubacterium saburreum		94.94	0.39	ESL 99		
Lactobacillus fermentum		5.06	0.20			
13135010	**Good identification**					**13135010**
Eubacterium saburreum		100.0	0.50	ESL 99		
13135012	**Good identification**					**13135012**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.38	ESL 99			
13135020	**Acceptable identification**						**13135020**
Eubacterium saburreum		94.96	0.39	ESL 99			
Actinomyces meyeri		3.35	0.07				
Lactobacillus fermentum		1.69	0.12				
13135030	**Good identification**						**13135030**
Eubacterium saburreum		100.0	0.38	ESL 99			
13135100	**Good identification**						**13135100**
					COC		
Eubacterium saburreum		91.07	0.58	ESL 99	1		
Actinomyces meyeri		7.13	0.32		1		
Atopobium parvulum		1.20	0.16		99		
13135102	**Acceptable identification**						**13135102**
Eubacterium saburreum		93.97	0.45	ESL 99			
Lactobacillus fermentum		3.51	0.24				
Actinomyces meyeri		1.29	0.06				
13135110	**Very good identification**						**13135110**
Eubacterium saburreum		100.0	0.57	ESL 99			
13135112	**Good identification**						**13135112**
Eubacterium saburreum		100.0	0.44	ESL 99			
13135120	**Acceptable identification**						**13135120**
					COC		
Eubacterium saburreum		92.07	0.45	ESL 99	1		
Actinomyces meyeri		5.57	0.17		1		
Atopobium parvulum		1.21	0.03		99		
13135130	**Good identification**						**13135130**
Eubacterium saburreum		100.0	0.44	ESL 99			
13150000	**Low discrimination**						**13150000**
					COC	GRA	
Prevotella intermedia		86.77	0.45	CEL 1	1	1	
Megasphaera elsdenii		11.08	0.32		99	1	
Actinomyces meyeri		1.01	0.13		1	99	
13151000	**Acceptable identification**						**13151000**
					GRA		
Prevotella intermedia		94.43	0.45	CEL 1	1		
Eubacterium saburreum		1.97	0.25		85		
Actinomyces meyeri		1.65	0.16		99		
13200000	**Low discrimination**						**13200000**
					GRA	SPO	
Clostridium bifermentans		69.24	0.35	SUC 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		13.61	0.17	SAL 10	1	1	
Clostridium cadaveris		4.21	0.13		99	99	
13200020	**Good identification**						**13200020**
Clostridium cadaveris		98.95	0.47	SUC 7			
Clostridium bifermentans		1.05	0.15	SAL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
13210020	**Good identification**						**13210020**
					SPO	CAT	
Clostridium cadaveris		99.72	0.47	SUC 7	99	1	
Propionibacterium acnes		0.28	0.06	SAL 1	1	95	
13210100	**Acceptable identification**						**13210100**
					GRA		
Fusobacterium mortiferum		91.67	0.50	IND 1	1		
Actinomyces meyeri		8.33	0.23		99		
13211000	**Low discrimination**						**13211000**
					GRA		
Fusobacterium mortiferum		88.36	0.47	IND 1	1		
Actinomyces meyeri		4.69	0.16		99		
Prevotella intermedia		3.06	0.11		1		
13211100	**Low discrimination**						**13211100**
					GRA		
Fusobacterium mortiferum		89.42	0.53	IND 1	1		
Actinomyces meyeri		8.13	0.26		99		
Eubacterium saburreum		1.60	0.21		85		
13211102	**Acceptable identification**						**13211102**
					GRA		
Fusobacterium mortiferum		92.13	0.37	IND 1	1		
Lactobacillus fermentum		7.87	0.20	BGL 10	99		
13230100	**Good identification**						**13230100**
Fusobacterium mortiferum		100.0	0.41	IND 1			
13231000	**Good identification**						**13231000**
Fusobacterium mortiferum		96.16	0.38	IND 1			
Eubacterium saburreum		3.84	0.12				
13231100	**Good identification**						**13231100**
Fusobacterium mortiferum		96.16	0.44	IND 1			
Eubacterium saburreum		3.84	0.18				
13250000	**Good identification**						**13250000**
Fusobacterium mortiferum		100.0	0.36	IND 1			
13250100	**Good identification**						**13250100**
Fusobacterium mortiferum		100.0	0.42	IND 1			
13251000	**Low discrimination**						**13251000**
					GRA		
Fusobacterium mortiferum		87.78	0.39	IND 1	1		
Lactobacillus catenaformis		9.87	0.12		99		
Lactobacillus jensenii		2.35	0.02		99		
13251100	**Acceptable identification**						**13251100**
					GRA		
Fusobacterium mortiferum		93.26	0.45	IND 1	1		
Lactobacillus catenaformis		6.74	0.15		99		
13271100	**Good identification**						**13271100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
13300000	**Good identification**						**13300000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Clostridium bifermentans		96.00	0.42	SUC 1	99	99		
Prevotella intermedia		1.03	0.11	SAL 10	1	1		
Clostridium botulinum biovar A		0.77	0.06		99	99		
13310000	**Low discrimination**							**13310000**
					COC	GRA	SPO	
Prevotella intermedia		79.34	0.45	SAL 1	1	1	1	
Megasphaera elsdenii		10.13	0.32		99	1	1	
Clostridium bifermentans		3.95	0.20		1	99	99	
13311000	**Low discrimination**							**13311000**
					GRA			
Prevotella intermedia		89.49	0.45	SAL 1	1			
Fusobacterium mortiferum		3.23	0.31		1			
Actinomyces odontolyticus		2.55	0.24		99			
13311100	**Low discrimination**							**13311100**
					GRA			
Fusobacterium mortiferum		34.90	0.37	IND 1	1			
Actinomyces meyeri		28.95	0.27	MLT 11	99			
Eubacterium saburreum		20.19	0.32		85			
13377100	**Good identification**							**13377100**
					SPO			
Actinomyces israelii		96.41	0.42	IND 1	1			
Clostridium tertium		3.59	0.03		99			
13377102	**Acceptable identification**							**13377102**
					SPO			
Actinomyces israelii		93.07	0.45	IND 1	1			
Clostridium tertium		6.93	0.11		99			
13377142	**Good identification**							**13377142**
Actinomyces israelii		100.0	0.38	IND 1				
13377500	**Good identification**							**13377500**
Actinomyces israelii		100.0	0.40	IND 1				
13377502	**Acceptable identification**							**13377502**
Actinomyces israelii		93.41	0.43	IND 1				
Bifidobacterium adolescentis		6.59	0.17					
13377542	**Good identification**							**13377542**
Actinomyces israelii		100.0	0.36	IND 1				
13400020	**Good identification**							**13400020**
Clostridium cadaveris		99.90	0.47	SUC 7				
Clostridium bifermentans		0.10	-.03	RAF 1				
13401000	**Low discrimination**							**13401000**
Eubacterium saburreum		47.08	0.37	ESL 99				
Fusobacterium necrophorum subsp. necrophorum		40.45	0.17					
Prevotella intermedia		7.69	0.14					
13401010	**Good identification**							**13401010**
Eubacterium saburreum		100.0	0.35	ESL 99				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
13401100	**Low discrimination**							**13401100**
					GRA			
Eubacterium saburreum		89.74	0.43	ESL 99	85			
Lactobacillus fermentum		4.38	0.24		99			
Fusobacterium mortiferum		2.85	0.19		1			
13401110	**Good identification**							**13401110**
Eubacterium saburreum		100.0	0.41	ESL 99				
13405100	**Acceptable identification**							**13405100**
Eubacterium saburreum		90.29	0.38	ESL 99				
Actinomyces meyeri		4.90	0.09					
Lactobacillus fermentum		4.80	0.20					
13405110	**Good identification**							**13405110**
Eubacterium saburreum		100.0	0.37	ESL 99				
13410000	**Low discrimination**							**13410000**
Prevotella intermedia		72.88	0.48	MLT 99				
Fusobacterium varium		9.95	0.25					
Fusobacterium mortiferum		9.46	0.44					
13410020	**Good identification**							**13410020**
					GRA	SPO	CAT	
Clostridium cadaveris		98.91	0.47	SUC 7	99	99	1	
Prevotella intermedia		0.61	0.14	RAF 1	1	1	1	
Propionibacterium acnes		0.28	0.06		99	1	95	
13410100	**Low discrimination**							**13410100**
					GRA			
Fusobacterium mortiferum		86.06	0.50	IND 1	1			
Actinomyces meyeri		7.82	0.23		99			
Prevotella intermedia		2.87	0.14		1			
13411000	**Low discrimination**							**13411000**
					GRA			
Prevotella intermedia		54.62	0.48	MLT 99	1			
Lactobacillus fermentum		23.32	0.54		99			
Fusobacterium mortiferum		10.63	0.47		1			
13411002	**Low discrimination**							**13411002**
					GRA			
Lactobacillus fermentum		86.29	0.54	IND 1	99			
Eubacterium saburreum		6.62	0.32		85			
Fusobacterium mortiferum		4.37	0.30		1			
13411010	**Low discrimination**							**13411010**
					GRA			
Eubacterium saburreum		89.29	0.43	ESL 99	85			
Prevotella intermedia		5.94	0.14		1			
Lactobacillus fermentum		2.54	0.20		99			
13411020	**Low discrimination**							**13411020**
					SPO			
Lactobacillus fermentum		67.31	0.46	IND 1	1			
Eubacterium saburreum		15.48	0.32		1			
Clostridium cadaveris		7.77	0.13		99			
13411022	**Good identification**							**13411022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		96.10	0.46	IND 1		
Eubacterium saburreum		3.90	0.19			
13411100	**Low discrimination**					**13411100**
					GRA	
Lactobacillus fermentum		42.60	0.58	IND 1	99	
Fusobacterium mortiferum		27.78	0.53		1	
Eubacterium saburreum		26.47	0.51		85	
13411102	**Low discrimination**					**13411102**
					GRA	
Lactobacillus fermentum		84.60	0.58	IND 1	99	
Eubacterium saburreum		9.28	0.38		85	
Fusobacterium mortiferum		6.13	0.37		1	
13411110	**Good identification**					**13411110**
					GRA	
Eubacterium saburreum		96.82	0.49	ESL 99	85	
Lactobacillus fermentum		1.92	0.24		99	
Fusobacterium mortiferum		1.26	0.19		1	
13411112	**Low discrimination**					**13411112**
Eubacterium saburreum		89.88	0.37	ESL 99		
Lactobacillus fermentum		10.12	0.24			
13411120	**Low discrimination**					**13411120**
					GRA	
Lactobacillus fermentum		66.42	0.50	IND 1	99	
Eubacterium saburreum		21.85	0.38		85	
Prevotella melaninogenica		8.81	0.15		1	
13411122	**Acceptable identification**					**13411122**
					GRA	
Lactobacillus fermentum		92.31	0.50	IND 1	99	
Eubacterium saburreum		5.36	0.25		85	
Prevotella melaninogenica		2.33	0.03		1	
13411130	**Good identification**					**13411130**
Eubacterium saburreum		96.38	0.37	ESL 99		
Lactobacillus fermentum		3.62	0.15			
13411400	**Good identification**					**13411400**
					GRA	
Lactobacillus fermentum		96.27	0.54	IND 1	99	
Prevotella intermedia		2.28	0.14		1	
Actinomyces meyeri		0.58	0.05		99	
13411402	**Good identification**					**13411402**
Lactobacillus fermentum		100.0	0.54	IND 1		
13411420	**Good identification**					**13411420**
Lactobacillus fermentum		100.0	0.46	IND 1		
13411422	**Good identification**					**13411422**
Lactobacillus fermentum		100.0	0.46	IND 1		
13411500	**Very good identification**					**13411500**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		97.31	0.58	IND 1	99	
Actinomyces meyeri		1.44	0.16		99	
Fusobacterium mortiferum		0.64	0.19		1	
13411502	**Very good identification**					**13411502**
Lactobacillus fermentum		100.0	0.58	IND 1		
13411520	**Good identification**					**13411520**
Lactobacillus fermentum		99.32	0.50	IND 1		
Bifidobacterium longum subsp. longum		0.68	0.05			
13411522	**Good identification**					**13411522**
Lactobacillus fermentum		99.32	0.50	IND 1		
Bifidobacterium longum subsp. longum		0.68	0.05			
13415000	**Low discrimination**					**13415000**
Lactobacillus fermentum		63.41	0.50	IND 1		
Eubacterium saburreum		25.26	0.40			
Actinomyces meyeri		5.07	0.21			
13415002	**Acceptable identification**					**13415002**
					GRA	
Lactobacillus fermentum		92.87	0.50	IND 1	99	
Eubacterium saburreum		6.53	0.27		85	
Fusobacterium mortiferum		0.60	0.11		1	
13415010	**Good identification**					**13415010**
Eubacterium saburreum		96.99	0.39	ESL 99		
Lactobacillus fermentum		3.01	0.16			
13415020	**Low discrimination**					**13415020**
Lactobacillus fermentum		80.41	0.42	IND 1		
Eubacterium saburreum		16.96	0.27			
Actinomyces meyeri		2.63	0.06			
13415022	**Good identification**					**13415022**
Lactobacillus fermentum		96.41	0.42	IND 1		
Eubacterium saburreum		3.59	0.14			
13415100	**Low discrimination**					**13415100**
Lactobacillus fermentum		54.08	0.54	IND 1		
Eubacterium saburreum		30.81	0.46			
Actinomyces meyeri		10.60	0.31			
13415102	**Low discrimination**					**13415102**
					GRA	
Lactobacillus fermentum		89.62	0.54	IND 1	99	
Eubacterium saburreum		9.01	0.33		85	
Fusobacterium mortiferum		0.83	0.17		1	
13415110	**Good identification**					**13415110**
Eubacterium saburreum		97.88	0.45	ESL 99		
Lactobacillus fermentum		2.12	0.20			
13415120	**Low discrimination**					**13415120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		72.37	0.46	IND 1		
Eubacterium saburreum		21.83	0.33			
Actinomyces meyeri		5.80	0.16			
13415122	**Acceptable identification**					**13415122**
Lactobacillus fermentum		94.95	0.46	IND 1		
Eubacterium saburreum		5.05	0.21			
13415400	**Good identification**					**13415400**
Lactobacillus fermentum		98.04	0.50	IND 1		
Actinomyces meyeri		1.96	0.10			
13415402	**Good identification**					**13415402**
Lactobacillus fermentum		100.0	0.50	IND 1		
13415420	**Good identification**					**13415420**
Lactobacillus fermentum		100.0	0.42	IND 1		
13415422	**Good identification**					**13415422**
Lactobacillus fermentum		100.0	0.42	IND 1		
13415500	**Acceptable identification**					**13415500**
Lactobacillus fermentum		94.81	0.54	IND 1		
Actinomyces meyeri		4.64	0.21			
Eubacterium saburreum		0.55	0.12			
13415502	**Good identification**					**13415502**
Lactobacillus fermentum		100.0	0.54	IND 1		
13415520	**Acceptable identification**					**13415520**
Lactobacillus fermentum		94.77	0.46	IND 1		
Bifidobacterium longum subsp. longum		3.33	0.14			
Actinomyces meyeri		1.90	0.06			
13415522	**Good identification**					**13415522**
Lactobacillus fermentum		96.61	0.46	IND 1		
Bifidobacterium longum subsp. longum		3.39	0.14			
13421100	**Good identification**					**13421100**
Eubacterium saburreum		100.0	0.40	ESL 99		
13421110	**Good identification**					**13421110**
Eubacterium saburreum		100.0	0.38	ESL 99		
13425100	**Good identification**					**13425100**
Eubacterium saburreum		100.0	0.35	ESL 99		
13430100	**Good identification**					**13430100**
Fusobacterium mortiferum		96.88	0.41	IND 1		
Eubacterium saburreum		3.12	0.14			
13431000	**Low discrimination**					**13431000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT	GRA	
Eubacterium saburreum		52.29	0.42	ESL 99	1	85	
Fusobacterium mortiferum		24.59	0.38		5	1	
Propionibacterium granulosum		17.03	0.32		90	99	
13431010	**Acceptable identification**						**13431010**
					CAT		
Eubacterium saburreum		94.85	0.40	ESL 99	1		
Propionibacterium granulosum		5.15	0.17		90		
13431100	**Low discrimination**						**13431100**
					GRA		
Eubacterium saburreum		66.91	0.48	ESL 99	85		
Fusobacterium mortiferum		31.46	0.44		1		
Lactobacillus fermentum		1.63	0.24		99		
13431102	**Low discrimination**						**13431102**
					GRA		
Eubacterium saburreum		69.72	0.35	ESL 99	85		
Fusobacterium mortiferum		20.64	0.28		1		
Lactobacillus fermentum		9.64	0.24		99		
13431110	**Good identification**						**13431110**
Eubacterium saburreum		100.0	0.46	ESL 99			
13431120	**Good identification**						**13431120**
Eubacterium saburreum		95.60	0.35	ESL 99			
Lactobacillus fermentum		4.40	0.15				
13435000	**Acceptable identification**						**13435000**
					GRA		
Eubacterium saburreum		90.60	0.37	ESL 99	85		
Fusobacterium mortiferum		5.96	0.19		1		
Lactobacillus fermentum		3.45	0.16		99		
13435010	**Good identification**						**13435010**
Eubacterium saburreum		100.0	0.36	ESL 99			
13435100	**Acceptable identification**						**13435100**
					GRA		
Eubacterium saburreum		91.55	0.43	ESL 99	85		
Fusobacterium mortiferum		6.02	0.25		1		
Lactobacillus fermentum		2.44	0.20		99		
13435110	**Good identification**						**13435110**
Eubacterium saburreum		100.0	0.42	ESL 99			
13450000	**Low discrimination**						**13450000**
Fusobacterium mortiferum		81.07	0.36	IND 1			
Prevotella intermedia		18.93	0.14				
13450100	**Good identification**						**13450100**
Fusobacterium mortiferum		100.0	0.42	IND 1			
13451000	**Low discrimination**						**13451000**
					GRA		
Fusobacterium mortiferum		79.94	0.39	IND 1	1		
Prevotella intermedia		12.44	0.14		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		5.31	0.20		99		
13451100	**Acceptable identification**						**13451100**
					GRA		
Fusobacterium mortiferum		92.99	0.45	IND 1	1		
Lactobacillus fermentum		4.32	0.24		99		
Eubacterium saburreum		2.69	0.17		85		
13471100	**Acceptable identification**						**13471100**
Fusobacterium mortiferum		93.95	0.36	IND 1			
Eubacterium saburreum		6.06	0.14				
13500000	**Acceptable identification**						**13500000**
					GRA	SPO	
Prevotella intermedia		93.09	0.48	FRU 99	1	1	
Clostridium bifermentans		5.34	0.24		99	99	
Clostridium botulinum biovar A		0.47	0.06		99	99	
13501000	**Low discrimination**						**13501000**
					GRA		
Prevotella intermedia		79.18	0.48	FRU 99	1		
Eubacterium saburreum		19.58	0.47		85		
Lactobacillus fermentum		1.02	0.28		99		
13501010	**Good identification**						**13501010**
Eubacterium saburreum		95.25	0.45	ESL 99			
Prevotella intermedia		4.76	0.14				
13501100	**Low discrimination**						**13501100**
					COC		
Eubacterium saburreum		65.49	0.53	ESL 99	1		
Blautia hansenii		30.30	0.32		99		
Lactobacillus fermentum		2.40	0.32		1		
13501102	**Low discrimination**						**13501102**
					COC		
Eubacterium saburreum		66.73	0.40	ESL 99	1		
Blautia hansenii		19.44	0.15		99		
Lactobacillus fermentum		13.83	0.32		1		
13501110	**Good identification**						**13501110**
					COC		
Eubacterium saburreum		99.43	0.52	ESL 99	1		
Blautia hansenii		0.57	-.03		99		
13501112	**Good identification**						**13501112**
Eubacterium saburreum		100.0	0.39	ESL 99			
13501130	**Good identification**						**13501130**
Eubacterium saburreum		96.06	0.39	ESL 99			
Prevotella melaninogenica		3.94	-.01				
13505000	**Low discrimination**						**13505000**
					GRA		
Eubacterium saburreum		88.27	0.42	ESL 99	85		
Prevotella intermedia		6.70	0.14		1		
Lactobacillus fermentum		5.04	0.24		99		
13505010	**Good identification**						**13505010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.41	ESL 99			
13505100	**Low discrimination**						**13505100**
					COC		
Eubacterium saburreum		82.86	0.49	ESL 99	1		
Blautia hansenii		12.57	0.19		99		
Lactobacillus fermentum		3.31	0.28		1		
13505102	**Low discrimination**						**13505102**
					COC		
Eubacterium saburreum		75.67	0.36	ESL 99	1		
Lactobacillus fermentum		17.11	0.28		1		
Blautia hansenii		7.22	0.03		99		
13505110	**Good identification**						**13505110**
Eubacterium saburreum		100.0	0.47	ESL 99			
13505120	**Low discrimination**						**13505120**
					COC		
Eubacterium saburreum		81.09	0.36	ESL 99	1		
Blautia hansenii		7.74	0.03		99		
Lactobacillus fermentum		6.11	0.20		1		
13505530	**Good identification**						**13505530**
Bifidobacterium longum subsp. longum		100.0	0.35	IND 1			
13505532	**Good identification**						**13505532**
Bifidobacterium longum subsp. longum		100.0	0.35	IND 1			
13510000	**Very good identification**						**13510000**
					COC		
Prevotella intermedia		99.86	0.82		1		
Megasphaera elsdenii		0.09	0.32		99		
Fusobacterium mortiferum		0.02	0.28		1		
13510001	**Low discrimination**						**13510001**
					GRA	SPO	
Prevotella intermedia		77.15	0.48	URE 1	1	1	
Clostridium sordellii		22.85	0.32		99	99	
13510002	**Good identification**						**13510002**
					COC	GRA	
Prevotella intermedia		97.33	0.48	BGL 1	1	1	
Lactobacillus fermentum		1.26	0.28		1	99	
Megasphaera elsdenii		0.92	0.15		99	1	
13510010	**Good identification**						**13510010**
Prevotella intermedia		99.39	0.48	MLZ 1			
Eubacterium saburreum		0.61	0.19				
13510020	**Acceptable identification**						**13510020**
					GRA	SPO	
Prevotella intermedia		91.75	0.48	NAG 1	1	1	
Clostridium cadaveris		7.85	0.25		99	99	
Lactobacillus fermentum		0.40	0.20		99	1	
13510040	**Acceptable identification**						**13510040**
					COC		
Prevotella intermedia		94.63	0.48	SOR 1	1		
Megasphaera elsdenii		5.37	0.29		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
13510100	**Acceptable identification**						**13510100**
					GRA		
Prevotella intermedia		90.20	0.48	LAC 1	1		
Fusobacterium mortiferum		3.38	0.34		1		
Actinomyces meyeri		2.80	0.24		99		
13510200	**Good identification**						**13510200**
Prevotella intermedia		100.0	0.48	RHA 1			
13510400	**Good identification**						**13510400**
					GRA		
Prevotella intermedia		98.53	0.48	ARA 1	1		
Lactobacillus fermentum		1.28	0.28		99		
Actinomyces meyeri		0.19	0.03		99		
13511000	**Very good identification**						**13511000**
					GRA		
Prevotella intermedia		97.96	0.82		1		
Lactobacillus fermentum		1.27	0.62		99		
Eubacterium saburreum		0.73	0.55		85		
13511001	**Acceptable identification**						**13511001**
					GRA	SPO	
Prevotella intermedia		93.22	0.48	URE 1	1	1	
Clostridium sordellii		4.87	0.19		99	99	
Lactobacillus fermentum		1.21	0.28		99	1	
13511002	**Low discrimination**						**13511002**
					GRA		
Lactobacillus fermentum		53.00	0.62	IND 1	99		
Prevotella intermedia		41.38	0.48		1		
Eubacterium saburreum		5.42	0.42		85		
13511010	**Low discrimination**						**13511010**
					GRA		
Prevotella intermedia		61.73	0.48	MLZ 1	1		
Eubacterium saburreum		37.47	0.53		85		
Lactobacillus fermentum		0.80	0.28		99		
13511012	**Low discrimination**						**13511012**
					GRA		
Eubacterium saburreum		82.30	0.41	ESL 99	85		
Lactobacillus fermentum		9.94	0.28		99		
Prevotella intermedia		7.76	0.14		1		
13511020	**Low discrimination**						**13511020**
					GRA		
Prevotella intermedia		62.88	0.48	NAG 1	1		
Lactobacillus fermentum		26.85	0.54		99		
Eubacterium saburreum		8.23	0.42		85		
13511022	**Acceptable identification**						**13511022**
					GRA		
Lactobacillus fermentum		91.62	0.54	IND 1	99		
Eubacterium saburreum		4.96	0.29		85		
Prevotella intermedia		2.17	0.14		1		
13511030	**Low discrimination**						**13511030**
					GRA		
Eubacterium saburreum		88.14	0.41	ESL 99	85		
Prevotella intermedia		8.31	0.14		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		3.55	0.20		99	
13511040	**Good identification**					**13511040**
					GRA	
Prevotella intermedia		98.00	0.48	SOR 1	1	
Lactobacillus fermentum		1.27	0.28		99	
Eubacterium saburreum		0.73	0.21		85	
13511100	**Low discrimination**					**13511100**
					GRA	
Lactobacillus fermentum		42.12	0.66	IND 1	99	
Eubacterium saburreum		34.89	0.61		85	
Prevotella intermedia		20.16	0.48		1	
13511102	**Low discrimination**					**13511102**
					GRA	
Lactobacillus fermentum		86.13	0.66	IND 1	99	
Eubacterium saburreum		12.59	0.48		85	
Prevotella intermedia		0.42	0.14		1	
13511110	**Very good identification**					**13511110**
					GRA	
Eubacterium saburreum		97.84	0.60	ESL 99	85	
Lactobacillus fermentum		1.46	0.32		99	
Prevotella intermedia		0.70	0.14		1	
13511112	**Acceptable identification**					**13511112**
Eubacterium saburreum		92.21	0.47	ESL 99		
Lactobacillus fermentum		7.79	0.32			
13511120	**Low discrimination**					**13511120**
					GRA	
Prevotella melaninogenica		74.67	0.50	IND 1	1	
Lactobacillus fermentum		17.07	0.58	MNS 90	99	
Eubacterium saburreum		7.48	0.48		85	
13511122	**Low discrimination**					**13511122**
					GRA	
Lactobacillus fermentum		51.30	0.58	IND 1	99	
Prevotella melaninogenica		42.75	0.37		1	
Eubacterium saburreum		3.97	0.35		85	
13511130	**Low discrimination**					**13511130**
					GRA	
Eubacterium saburreum		86.05	0.47	ESL 99	85	
Prevotella melaninogenica		10.60	0.15		1	
Lactobacillus fermentum		2.42	0.24		99	
13511200	**Good identification**					**13511200**
					GRA	
Prevotella intermedia		98.00	0.48	RHA 1	1	
Lactobacillus fermentum		1.27	0.28		99	
Eubacterium saburreum		0.73	0.21		85	
13511400	**Low discrimination**					**13511400**
					GRA	
Lactobacillus fermentum		55.90	0.62	IND 1	99	
Prevotella intermedia		43.65	0.48		1	
Eubacterium saburreum		0.33	0.21		85	
13511402	**Very good identification**					**13511402**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		99.22	0.62	IND 1	99			
Prevotella intermedia		0.78	0.14		1			
13511420	**Good identification**							**13511420**
					GRA			
Lactobacillus fermentum		97.33	0.54	IND 1	99			
Prevotella intermedia		2.30	0.14		1			
Bifidobacterium longum subsp. longum		0.36	0.05		99			
13511422	**Good identification**							**13511422**
Lactobacillus fermentum		99.63	0.54	IND 1				
Bifidobacterium longum subsp. longum		0.37	0.05					
13511500	**Very good identification**							**13511500**
Lactobacillus fermentum		96.37	0.66	IND 1				
Bifidobacterium longum subsp. longum		1.82	0.29					
Eubacterium saburreum		0.81	0.27					
13511502	**Very good identification**							**13511502**
Lactobacillus fermentum		97.93	0.66	IND 1				
Bifidobacterium longum subsp. longum		1.85	0.29					
Eubacterium saburreum		0.15	0.14					
13511520	**Low discrimination**							**13511520**
					GRA			
Lactobacillus fermentum		78.27	0.58	IND 1	99			
Bifidobacterium longum subsp. longum		17.75	0.40		99			
Prevotella melaninogenica		3.46	0.15		1			
13511522	**Low discrimination**							**13511522**
					GRA			
Lactobacillus fermentum		80.76	0.58	IND 1	99			
Bifidobacterium longum subsp. longum		18.31	0.40		99			
Prevotella melaninogenica		0.68	0.03		1			
13511530	**Good identification**							**13511530**
Bifidobacterium longum subsp. longum		96.57	0.44	IND 1				
Lactobacillus fermentum		2.53	0.24	MNS 87				
Eubacterium saburreum		0.91	0.13					
13511532	**Good identification**							**13511532**
Bifidobacterium longum subsp. longum		97.45	0.44	IND 1				
Lactobacillus fermentum		2.55	0.24	MNS 87				
13512000	**Acceptable identification**							**13512000**
					COC	GRA	CAT	
Prevotella intermedia		91.64	0.48	MAN 1	1	1	1	
Megasphaera elsdenii		7.80	0.32		99	1	1	
Propionibacterium granulosum		0.42	0.16		1	99	90	
13512100	**Acceptable identification**							**13512100**
					GRA			
Propionibacterium propionicum		93.88	0.46	IND 1	99			
Prevotella intermedia		6.12	0.14	NIT 86	1			
13512102	**Good identification**							**13512102**
Propionibacterium propionicum		100.0	0.49	IND 1				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
13512140	**Good identification**					**13512140**
Propionibacterium propionicum		100.0	0.46	IND 1		
13512142	**Good identification**					**13512142**
Propionibacterium propionicum		100.0	0.49	IND 1		
13513000	**Good identification**					**13513000**
					GRA	
Prevotella intermedia		97.24	0.48	MAN 1	1	
Lactobacillus fermentum		1.26	0.28		99	
Eubacterium saburreum		0.73	0.21		85	
13513100	**Low discrimination**					**13513100**
Propionibacterium propionicum		74.12	0.46	IND 1		
Lactobacillus fermentum		10.10	0.32	NIT 86		
Eubacterium saburreum		8.37	0.27			
13513102	**Low discrimination**					**13513102**
Propionibacterium propionicum		88.70	0.49	IND 1		
Lactobacillus fermentum		8.06	0.32	NIT 86		
Actinomyces naeslundii		2.06	0.32			
13513140	**Good identification**					**13513140**
Propionibacterium propionicum		100.0	0.46	IND 1		
13513142	**Good identification**					**13513142**
Propionibacterium propionicum		100.0	0.49	IND 1		
13514000	**Good identification**					**13514000**
					GRA	
Prevotella intermedia		97.40	0.48	XYL 1	1	
Actinomyces meyeri		1.47	0.18		99	
Lactobacillus fermentum		0.74	0.24		99	
13515000	**Low discrimination**					**13515000**
					GRA	
Prevotella intermedia		45.95	0.48	XYL 1	1	
Lactobacillus fermentum		34.57	0.58		99	
Eubacterium saburreum		18.36	0.50		85	
13515002	**Good identification**					**13515002**
					GRA	
Lactobacillus fermentum		90.32	0.58	IND 1	99	
Eubacterium saburreum		8.47	0.37		85	
Prevotella intermedia		1.21	0.14		1	
13515010	**Acceptable identification**					**13515010**
					GRA	
Eubacterium saburreum		94.86	0.49	ESL 99	85	
Prevotella intermedia		2.93	0.14		1	
Lactobacillus fermentum		2.21	0.24		99	
13515012	**Low discrimination**					**13515012**
Eubacterium saburreum		88.36	0.36	ESL 99		
Lactobacillus fermentum		11.64	0.24			
13515020	**Low discrimination**					**13515020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Lactobacillus fermentum		74.98	0.50	IND 1	99	
Eubacterium saburreum		21.08	0.37		85	
Prevotella intermedia		3.02	0.14		1	
13515022	**Good identification**					**13515022**
Lactobacillus fermentum		95.27	0.50	IND 1		
Eubacterium saburreum		4.73	0.25			
13515030	**Good identification**					**13515030**
Eubacterium saburreum		95.80	0.36	ESL 99		
Lactobacillus fermentum		4.21	0.16			
13515100	**Low discrimination**					**13515100**
Lactobacillus fermentum		54.20	0.62	IND 1		
Eubacterium saburreum		41.17	0.57			
Actinomyces meyeri		3.99	0.32			
13515102	**Low discrimination**					**13515102**
Lactobacillus fermentum		87.90	0.62	IND 1		
Eubacterium saburreum		11.78	0.44			
Actinomyces meyeri		0.20	0.06			
13515110	**Good identification**					**13515110**
Eubacterium saburreum		98.14	0.55	ESL 99		
Lactobacillus fermentum		1.60	0.28			
Bifidobacterium longum subsp. longum		0.26	0.07			
13515112	**Low discrimination**					**13515112**
Eubacterium saburreum		89.11	0.42	ESL 99		
Lactobacillus fermentum		8.21	0.28			
Bifidobacterium longum subsp. longum		1.35	0.07			
13515120	**Low discrimination**					**13515120**
					GRA	
Lactobacillus fermentum		65.82	0.54	IND 1	99	
Eubacterium saburreum		26.47	0.44		85	
Prevotella melaninogenica		4.95	0.15		1	
13515122	**Low discrimination**					**13515122**
					GRA	
Lactobacillus fermentum		89.03	0.54	IND 1	99	
Eubacterium saburreum		6.32	0.31		85	
Prevotella melaninogenica		1.28	0.03		1	
13515130	**Acceptable identification**					**13515130**
Eubacterium saburreum		91.64	0.42	ESL 99		
Bifidobacterium longum subsp. longum		5.55	0.18			
Lactobacillus fermentum		2.81	0.20			
13515400	**Very good identification**					**13515400**
					GRA	
Lactobacillus fermentum		97.44	0.58	IND 1	99	
Prevotella intermedia		1.31	0.14		1	
Actinomyces meyeri		0.73	0.11		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
13515402	**Very good identification**						**13515402**
Lactobacillus fermentum		100.0	0.58	IND 1			
13515420	**Good identification**						**13515420**
Lactobacillus fermentum		98.13	0.50	IND 1			
Bifidobacterium longum subsp. longum		1.87	0.14				
13515422	**Good identification**						**13515422**
Lactobacillus fermentum		98.13	0.50	IND 1			
Bifidobacterium longum subsp. longum		1.87	0.14				
13515500	**Low discrimination**						**13515500**
Lactobacillus fermentum		89.08	0.62	IND 1			
Bifidobacterium longum subsp. longum		8.60	0.37				
Actinomyces meyeri		1.64	0.21				
13515502	**Good identification**						**13515502**
Lactobacillus fermentum		90.35	0.62	IND 1			
Bifidobacterium longum subsp. longum		8.72	0.37				
Bifidobacterium adolescentis		0.81	0.21				
13515510	**Acceptable identification**						**13515510**
Bifidobacterium longum subsp. longum		90.94	0.41	IND 1			
Lactobacillus fermentum		5.59	0.28	MNS 87			
Eubacterium saburreum		3.47	0.21				
13515512	**Low discrimination**						**13515512**
Bifidobacterium longum subsp. longum		88.63	0.41	IND 1			
Bifidobacterium adolescentis		5.92	0.23	MNS 87			
Lactobacillus fermentum		5.45	0.28				
13515520	**Low discrimination**						**13515520**
					GRA		
Bifidobacterium longum subsp. longum		52.34	0.48	IND 1	99		
Lactobacillus fermentum		45.19	0.54	MNS 87	99		
Bacteroides vulgatus		1.95	0.09		1		
13515522	**Low discrimination**						**13515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		53.29	0.48	IND 1	99	1	
Lactobacillus fermentum		46.01	0.54	MNS 87	99	1	
Bacteroides thetaiotaomicron		0.70	0.04		1	89	
13515530	**Good identification**						**13515530**
Bifidobacterium longum subsp. longum		99.49	0.52	IND 1			
Lactobacillus fermentum		0.51	0.20	MNS 87			
13515532	**Good identification**						**13515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.20	0.52	IND 1	99	1	
Lactobacillus fermentum		0.51	0.20	MNS 87	99	1	
Bacteroides thetaiotaomicron		0.30	-.03		1	89	
13521000	**Acceptable identification**						**13521000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		94.23	0.44	ESL 99			
Prevotella intermedia		5.77	0.14				
13521010	**Good identification**						**13521010**
Eubacterium saburreum		100.0	0.42	ESL 99			
13521100	**Good identification**						**13521100**
					COC		
Eubacterium saburreum		99.30	0.50	ESL 99	1		
Blautia hansenii		0.70	-.03		99		
13521102	**Good identification**						**13521102**
Eubacterium saburreum		100.0	0.37	ESL 99			
13521110	**Good identification**						**13521110**
Eubacterium saburreum		100.0	0.49	ESL 99			
13521112	**Good identification**						**13521112**
Eubacterium saburreum		100.0	0.36	ESL 99			
13521120	**Good identification**						**13521120**
Eubacterium saburreum		95.20	0.37	ESL 99			
Prevotella melaninogenica		4.80	-.01				
13521130	**Good identification**						**13521130**
Eubacterium saburreum		100.0	0.36	ESL 99			
13525000	**Good identification**						**13525000**
Eubacterium saburreum		100.0	0.39	ESL 99			
13525010	**Good identification**						**13525010**
Eubacterium saburreum		100.0	0.38	ESL 99			
13525100	**Good identification**						**13525100**
Eubacterium saburreum		100.0	0.46	ESL 99			
13525110	**Good identification**						**13525110**
Eubacterium saburreum		100.0	0.44	ESL 99			
13530000	**Good identification**						**13530000**
					CAT	GRA	
Prevotella intermedia		97.29	0.48	TRE 1	1	1	
Propionibacterium granulosum		1.76	0.26		90	99	
Eubacterium saburreum		0.49	0.18		1	85	
13531000	**Low discrimination**						**13531000**
					CAT	GRA	
Prevotella intermedia		65.16	0.48	TRE 1	1	1	
Eubacterium saburreum		32.23	0.52		1	85	
Propionibacterium granulosum		1.18	0.26		90	99	
13531002	**Low discrimination**						**13531002**
					GRA		
Eubacterium saburreum		74.77	0.39	ESL 99	85		
Lactobacillus fermentum		11.08	0.28		99		
Prevotella intermedia		8.65	0.14		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
13531010	**Good identification**						**13531010**
					CAT	GRA	
Eubacterium saburreum		96.99	0.50	ESL 99	1	85	
Prevotella intermedia		2.42	0.14		1	1	
Propionibacterium granulosum		0.59	0.11		90	99	
13531012	**Good identification**						**13531012**
Eubacterium saburreum		100.0	0.38	ESL 99			
13531020	**Low discrimination**						**13531020**
					CAT	GRA	
Eubacterium saburreum		78.66	0.39	ESL 99	1	85	
Prevotella intermedia		9.10	0.14		1	1	
Propionibacterium granulosum		4.08	0.15		90	99	
13531030	**Good identification**						**13531030**
Eubacterium saburreum		100.0	0.38	ESL 99			
13531100	**Very good identification**						**13531100**
					GRA		
Eubacterium saburreum		95.58	0.58	ESL 99	85		
Lactobacillus fermentum		1.75	0.32		99		
Fusobacterium mortiferum		1.39	0.28		1		
13531102	**Low discrimination**						**13531102**
Eubacterium saburreum		87.06	0.45	ESL 99			
Lactobacillus fermentum		9.02	0.32				
Actinomyces naeslundii		2.30	0.32				
13531110	**Very good identification**						**13531110**
Eubacterium saburreum		100.0	0.57	ESL 99			
13531112	**Good identification**						**13531112**
Eubacterium saburreum		100.0	0.44	ESL 99			
13531120	**Low discrimination**						**13531120**
					GRA		
Eubacterium saburreum		76.07	0.45	ESL 99	85		
Prevotella melaninogenica		11.50	0.15		1		
Clostridium perfringens		7.80	0.19		99		
13531130	**Good identification**						**13531130**
Eubacterium saburreum		100.0	0.44	ESL 99			
13532100	**Good identification**						**13532100**
Propionibacterium propionicum		100.0	0.45	IND 1			
13532102	**Good identification**						**13532102**
Propionibacterium propionicum		100.0	0.48	IND 1			
13532140	**Good identification**						**13532140**
Propionibacterium propionicum		100.0	0.45	IND 1			
13532142	**Good identification**						**13532142**
Propionibacterium propionicum		100.0	0.48	IND 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
13533100	**Low discrimination**					**13533100**
Propionibacterium propionicum		85.14	0.45	IND 1		
Eubacterium saburreum		7.83	0.24	NIT 86		
Actinomyces naeslundii		7.02	0.37			
13533101	**Low discrimination**					**13533101**
Actinomyces naeslundii		52.69	0.40	IND 1		
Propionibacterium propionicum		47.31	0.29	MNS 88		
13533102	**Acceptable identification**					**13533102**
Propionibacterium propionicum		93.82	0.48	IND 1		
Actinomyces naeslundii		5.16	0.37	NIT 86		
Eubacterium saburreum		1.02	0.11			
13533120	**Low discrimination**					**13533120**
Actinomyces naeslundii		83.55	0.37	IND 1		
Eubacterium saburreum		16.45	0.11	MNS 88		
13533121	**Good identification**					**13533121**
Actinomyces naeslundii		100.0	0.40	IND 1		
13533122	**Low discrimination**					**13533122**
Actinomyces naeslundii		84.48	0.37	IND 1		
Propionibacterium propionicum		15.52	0.14	MNS 88		
13533123	**Good identification**					**13533123**
Actinomyces naeslundii		100.0	0.40	IND 1		
13533140	**Good identification**					**13533140**
Propionibacterium propionicum		98.57	0.45	IND 1		
Actinomyces naeslundii		1.44	0.24	NIT 86		
13533142	**Good identification**					**13533142**
Propionibacterium propionicum		99.04	0.48	IND 1		
Actinomyces naeslundii		0.96	0.24	NIT 86		
13535000	**Acceptable identification**					**13535000**
					GRA	
Eubacterium saburreum		93.77	0.47	ESL 99	85	
Prevotella intermedia		3.56	0.14		1	
Lactobacillus fermentum		2.68	0.24		99	
13535010	**Good identification**					**13535010**
Eubacterium saburreum		100.0	0.46	ESL 99		
13535100	**Good identification**					**13535100**
Eubacterium saburreum		98.04	0.54	ESL 99		
Lactobacillus fermentum		1.96	0.28			
13535102	**Low discrimination**					**13535102**
Eubacterium saburreum		89.84	0.41	ESL 99		
Lactobacillus fermentum		10.16	0.28			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
13535110	**Good identification**						**13535110**
Eubacterium saburreum		100.0	0.52	ESL 99			
13535112	**Good identification**						**13535112**
Eubacterium saburreum		100.0	0.39	ESL 99			
13535120	**Good identification**						**13535120**
Eubacterium saburreum		96.37	0.41	ESL 99			
Lactobacillus fermentum		3.63	0.20				
13535130	**Good identification**						**13535130**
Eubacterium saburreum		97.78	0.39	ESL 99			
Bifidobacterium longum subsp. longum		2.22	0.07				
13535520	**Good identification**						**13535520**
Bifidobacterium longum subsp. longum		96.63	0.37	IND 1			
Lactobacillus fermentum		3.37	0.20	MNS 87			
13535522	**Low discrimination**						**13535522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		56.42	0.37	IND 1	99	1	
Bacteroides thetaiotaomicron		39.18	0.23	MNS 87	1	89	
Parabacteroides distasonis		2.43	0.18		1	76	
13535530	**Good identification**						**13535530**
Bifidobacterium longum subsp. longum		100.0	0.41	IND 1			
13535532	**Low discrimination**						**13535532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		85.31	0.41	IND 1	99	1	
Bacteroides thetaiotaomicron		13.53	0.16	MNS 87	1	89	
Parabacteroides distasonis		1.16	0.14		1	76	
13535722	**Good identification**						**13535722**
Bacteroides thetaiotaomicron		98.30	0.42	ESL 92			
Parabacteroides distasonis		1.38	0.26	MNS 99			
Bacteroides ovatus		0.32	0.00				
13535732	**Good identification**						**13535732**
Bacteroides thetaiotaomicron		97.10	0.35	ESL 92			
Parabacteroides distasonis		1.89	0.22	MNS 99			
Bacteroides ovatus		0.55	-.03				
13550000	**Good identification**						**13550000**
Prevotella intermedia		99.47	0.48	CEL 1			
Fusobacterium mortiferum		0.53	0.20				
13551000	**Good identification**						**13551000**
					GRA		
Prevotella intermedia		97.23	0.48	CEL 1	1		
Lactobacillus fermentum		1.26	0.28		99		
Fusobacterium mortiferum		0.78	0.23		1		
13553101	**Good identification**						**13553101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	IND 1		
13553103	**Good identification**					**13553103**
Actinomyces naeslundii		100.0	0.37	IND 1		
13553121	**Good identification**					**13553121**
Actinomyces naeslundii		100.0	0.37	IND 1		
13553123	**Good identification**					**13553123**
Actinomyces naeslundii		100.0	0.37	IND 1		
13571101	**Good identification**					**13571101**
Actinomyces naeslundii		100.0	0.37	IND 1		
13571103	**Good identification**					**13571103**
Actinomyces naeslundii		100.0	0.37	IND 1		
13571121	**Good identification**					**13571121**
Actinomyces naeslundii		100.0	0.37	IND 1		
13571123	**Good identification**					**13571123**
Actinomyces naeslundii		100.0	0.37	IND 1		
13573100	**Low discrimination**					**13573100**
Actinomyces naeslundii		87.01	0.39	IND 1		
Actinomyces israelii		12.99	0.15	MNS 88		
13573101	**Good identification**					**13573101**
Actinomyces naeslundii		100.0	0.42	IND 1		
13573102	**Low discrimination**					**13573102**
Actinomyces naeslundii		73.40	0.39	IND 1		
Actinomyces israelii		16.43	0.18	MNS 88		
Propionibacterium propionicum		10.17	0.14			
13573103	**Good identification**					**13573103**
Actinomyces naeslundii		100.0	0.42	IND 1		
13573120	**Good identification**					**13573120**
Actinomyces naeslundii		100.0	0.39	IND 1		
13573121	**Good identification**					**13573121**
Actinomyces naeslundii		100.0	0.42	IND 1		
13573122	**Good identification**					**13573122**
Actinomyces naeslundii		100.0	0.39	IND 1		
13573123	**Good identification**					**13573123**
Actinomyces naeslundii		100.0	0.42	IND 1		
13575722	**Low discrimination**					**13575722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		86.42	0.48	ESL 92		
Bacteroides ovatus		13.28	0.34	MNS 99		
Parabacteroides distasonis		0.31	0.22			
13575732	**Low discrimination**					**13575732**
Bacteroides thetaiotaomicron		78.66	0.41	ESL 92		
Bacteroides ovatus		20.72	0.31	MNS 99		
Parabacteroides distasonis		0.39	0.17			
13577102	**Acceptable identification**					**13577102**
Actinomyces israelii		94.92	0.36	IND 1		
Bifidobacterium adolescentis		5.08	0.08	SAL 92		
13610000	**Good identification**					**13610000**
Fusobacterium mortiferum		96.15	0.49	IND 1		
Prevotella intermedia		3.85	0.14			
13610100	**Good identification**					**13610100**
Fusobacterium mortiferum		100.0	0.55	IND 1		
13610102	**Good identification**					**13610102**
Fusobacterium mortiferum		100.0	0.39	IND 1		
13611000	**Good identification**					**13611000**
					GRA	
Fusobacterium mortiferum		95.87	0.52	IND 1	1	
Prevotella intermedia		2.56	0.14		1	
Lactobacillus fermentum		1.09	0.20		99	
13611002	**Acceptable identification**					**13611002**
					GRA	
Fusobacterium mortiferum		90.69	0.35	IND 1	1	
Lactobacillus fermentum		9.31	0.20	BGL 10	99	
13611100	**Very good identification**					**13611100**
					GRA	
Fusobacterium mortiferum		98.72	0.58	IND 1	1	
Lactobacillus fermentum		0.79	0.24		99	
Eubacterium saburreum		0.49	0.17		85	
13611102	**Acceptable identification**					**13611102**
					GRA	
Fusobacterium mortiferum		93.30	0.42	IND 1	1	
Lactobacillus fermentum		6.70	0.24	BGL 10	99	
13614100	**Good identification**					**13614100**
Fusobacterium mortiferum		100.0	0.36	IND 1		
13615100	**Acceptable identification**					**13615100**
					GRA	
Fusobacterium mortiferum		91.10	0.39	IND 1	1	
Lactobacillus fermentum		5.67	0.20	XYL 7	99	
Eubacterium saburreum		3.23	0.12		85	
13630000	**Good identification**					**13630000**
Fusobacterium mortiferum		100.0	0.40	IND 1		
13630100	**Good identification**					**13630100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.46	IND 1		
13631000	**Good identification**					**13631000**
Fusobacterium mortiferum		100.0	0.43	IND 1		
13631100	**Good identification**					**13631100**
Fusobacterium mortiferum		98.91	0.49	IND 1		
Eubacterium saburreum		1.10	0.14			
13650000	**Good identification**					**13650000**
Fusobacterium mortiferum		100.0	0.40	IND 1		
13650100	**Good identification**					**13650100**
Fusobacterium mortiferum		100.0	0.47	IND 1		
13651000	**Good identification**					**13651000**
Fusobacterium mortiferum		100.0	0.43	IND 1		
13651100	**Good identification**					**13651100**
Fusobacterium mortiferum		100.0	0.50	IND 1		
13670100	**Good identification**					**13670100**
Fusobacterium mortiferum		100.0	0.38	IND 1		
13671100	**Good identification**					**13671100**
					GRA	
Fusobacterium mortiferum		98.19	0.41	IND 1	1	
Actinomyces naeslundii		1.81	0.23		99	
13710000	**Good identification**					**13710000**
Prevotella intermedia		96.98	0.48	SAL 1		
Fusobacterium mortiferum		3.02	0.33			
13710100	**Low discrimination**					**13710100**
Fusobacterium mortiferum		87.79	0.39	IND 1		
Prevotella intermedia		12.21	0.14	MLT 11		
13711000	**Acceptable identification**					**13711000**
					GRA	
Prevotella intermedia		93.41	0.48	SAL 1	1	
Fusobacterium mortiferum		4.36	0.36		1	
Lactobacillus fermentum		1.21	0.28		99	
13711100	**Low discrimination**					**13711100**
					GRA	
Fusobacterium mortiferum		65.06	0.42	IND 1	1	
Lactobacillus fermentum		12.60	0.32	MLT 11	99	
Eubacterium saburreum		10.44	0.27		85	
13711101	**Good identification**					**13711101**
Actinomyces naeslundii		100.0	0.39	IND 1		
13711103	**Good identification**					**13711103**
Actinomyces naeslundii		100.0	0.39	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
13711121	**Good identification**					**13711121**
Actinomyces naeslundii		100.0	0.39	IND 1		
13711122	**Low discrimination**					**13711122**
					GRA	
Actinomyces naeslundii		28.82	0.36	IND 1	99	
Prevotella oralis		25.23	0.09	MNS 88	1	
Lactobacillus fermentum		20.62	0.24		99	
13711123	**Good identification**					**13711123**
Actinomyces naeslundii		100.0	0.39	IND 1		
13713001	**Good identification**					**13713001**
Actinomyces naeslundii		100.0	0.36	IND 1		
13713003	**Good identification**					**13713003**
Actinomyces naeslundii		100.0	0.36	IND 1		
13713021	**Good identification**					**13713021**
Actinomyces naeslundii		100.0	0.36	IND 1		
13713023	**Good identification**					**13713023**
Actinomyces naeslundii		100.0	0.36	IND 1		
13713100	**Acceptable identification**					**13713100**
Actinomyces naeslundii		92.43	0.41	IND 1		
Propionibacterium propionicum		7.57	0.12	MNS 88		
13713101	**Good identification**					**13713101**
Actinomyces naeslundii		100.0	0.44	IND 1		
13713102	**Low discrimination**					**13713102**
Actinomyces naeslundii		89.06	0.41	IND 1		
Propionibacterium propionicum		10.95	0.15	MNS 88		
13713103	**Good identification**					**13713103**
Actinomyces naeslundii		100.0	0.44	IND 1		
13713120	**Good identification**					**13713120**
Actinomyces naeslundii		100.0	0.41	IND 1		
13713121	**Good identification**					**13713121**
Actinomyces naeslundii		100.0	0.44	IND 1		
13713122	**Good identification**					**13713122**
Actinomyces naeslundii		100.0	0.41	IND 1		
13713123	**Good identification**					**13713123**
Actinomyces naeslundii		100.0	0.44	IND 1		
13715512	**Good identification**					**13715512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		97.40	0.36	IND 1		
Bifidobacterium longum subsp. longum		2.60	0.07	MNS 87		
13731001	**Good identification**					**13731001**
Actinomyces naeslundii		100.0	0.36	IND 1		
13731003	**Good identification**					**13731003**
Actinomyces naeslundii		100.0	0.36	IND 1		
13731021	**Good identification**					**13731021**
Actinomyces naeslundii		100.0	0.36	IND 1		
13731023	**Good identification**					**13731023**
Actinomyces naeslundii		100.0	0.36	IND 1		
13731101	**Good identification**					**13731101**
Actinomyces naeslundii		100.0	0.44	IND 1		
13731102	**Low discrimination**					**13731102**
					GRA	
Actinomyces naeslundii		77.08	0.41	IND 1	99	
Fusobacterium mortiferum		14.61	0.17	MNS 88	1	
Eubacterium saburreum		8.31	0.11		85	
13731103	**Good identification**					**13731103**
Actinomyces naeslundii		100.0	0.44	IND 1		
13731120	**Low discrimination**					**13731120**
Actinomyces naeslundii		72.39	0.41	IND 1		
Clostridium perfringens		19.80	0.09	MNS 88		
Eubacterium saburreum		7.80	0.11			
13731121	**Good identification**					**13731121**
Actinomyces naeslundii		100.0	0.44	IND 1		
13731122	**Low discrimination**					**13731122**
Actinomyces naeslundii		88.59	0.41	IND 1		
Clostridium perfringens		11.41	0.03	MNS 88		
13731123	**Good identification**					**13731123**
Actinomyces naeslundii		100.0	0.44	IND 1		
13733000	**Good identification**					**13733000**
Actinomyces naeslundii		100.0	0.38	IND 1		
13733001	**Good identification**					**13733001**
Actinomyces naeslundii		100.0	0.41	IND 1		
13733002	**Good identification**					**13733002**
Actinomyces naeslundii		100.0	0.38	IND 1		
13733003	**Good identification**					**13733003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	IND 1		
13733020	**Good identification**					**13733020**
Actinomyces naeslundii		100.0	0.38	IND 1		
13733021	**Good identification**					**13733021**
Actinomyces naeslundii		100.0	0.41	IND 1		
13733022	**Good identification**					**13733022**
Actinomyces naeslundii		100.0	0.38	IND 1		
13733023	**Good identification**					**13733023**
Actinomyces naeslundii		100.0	0.41	IND 1		
13733100	**Good identification**					**13733100**
Actinomyces naeslundii		95.40	0.46	IND 1		
Actinomyces israelii		4.61	0.14	MNS 88		
13733101	**Good identification**					**13733101**
Actinomyces naeslundii		100.0	0.49	IND 1		
13733102	**Low discrimination**					**13733102**
Actinomyces naeslundii		88.95	0.46	IND 1		
Actinomyces israelii		6.44	0.17	MNS 88		
Propionibacterium propionicum		4.61	0.14			
13733103	**Good identification**					**13733103**
Actinomyces naeslundii		100.0	0.49	IND 1		
13733120	**Good identification**					**13733120**
Actinomyces naeslundii		100.0	0.46	IND 1		
13733121	**Good identification**					**13733121**
Actinomyces naeslundii		100.0	0.49	IND 1		
13733122	**Good identification**					**13733122**
Actinomyces naeslundii		100.0	0.46	IND 1		
13733123	**Good identification**					**13733123**
Actinomyces naeslundii		100.0	0.49	IND 1		
13733141	**Good identification**					**13733141**
Actinomyces naeslundii		100.0	0.36	IND 1		
13733143	**Good identification**					**13733143**
Actinomyces naeslundii		100.0	0.36	IND 1		
13733161	**Good identification**					**13733161**
Actinomyces naeslundii		100.0	0.36	IND 1		
13733163	**Good identification**					**13733163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	IND 1		
13737102	**Low discrimination**					**13737102**
Actinomyces israelii		85.84	0.35	IND 1		
Bifidobacterium adolescentis		14.16	0.16	CEL 93		
13751101	**Good identification**					**13751101**
Actinomyces naeslundii		100.0	0.41	IND 1		
13751102	**Low discrimination**					**13751102**
					GRA	
Actinomyces naeslundii		41.20	0.38	IND 1	99	
Lactobacillus acidophilus		26.30	0.10	MNS 88	99	
Fusobacterium mortiferum		12.76	0.18		1	
13751103	**Good identification**					**13751103**
Actinomyces naeslundii		100.0	0.41	IND 1		
13751120	**Low discrimination**					**13751120**
					GRA	
Actinomyces naeslundii		72.17	0.38	IND 1	99	
Prevotella melaninogenica		18.94	-.01	MNS 88	1	
Prevotella oralis		8.90	-.03		1	
13751121	**Good identification**					**13751121**
Actinomyces naeslundii		100.0	0.41	IND 1		
13751123	**Good identification**					**13751123**
Actinomyces naeslundii		100.0	0.41	IND 1		
13753000	**Low discrimination**					**13753000**
Actinomyces naeslundii		84.91	0.35	IND 1		
Actinomyces israelii		15.09	0.13	MNS 88		
13753001	**Good identification**					**13753001**
Actinomyces naeslundii		100.0	0.38	IND 1		
13753002	**Low discrimination**					**13753002**
Actinomyces naeslundii		78.96	0.35	IND 1		
Actinomyces israelii		21.05	0.16	MNS 88		
13753003	**Good identification**					**13753003**
Actinomyces naeslundii		100.0	0.38	IND 1		
13753020	**Good identification**					**13753020**
Actinomyces naeslundii		100.0	0.35	IND 1		
13753021	**Good identification**					**13753021**
Actinomyces naeslundii		100.0	0.38	IND 1		
13753022	**Good identification**					**13753022**
Actinomyces naeslundii		100.0	0.35	IND 1		
13753023	**Good identification**					**13753023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	IND 1		
13753100	**Low discrimination**					**13753100**
Actinomyces naeslundii		80.02	0.43	IND 1		
Actinomyces israelii		19.98	0.24	MNS 88		
13753101	**Good identification**					**13753101**
Actinomyces naeslundii		100.0	0.46	IND 1		
13753102	**Low discrimination**					**13753102**
Actinomyces naeslundii		68.01	0.43	IND 1		
Actinomyces israelii		25.48	0.27	MNS 88		
Bifidobacterium adolescentis		6.52	0.10			
13753103	**Good identification**					**13753103**
Actinomyces naeslundii		100.0	0.46	IND 1		
13753120	**Good identification**					**13753120**
Actinomyces naeslundii		100.0	0.43	IND 1		
13753121	**Good identification**					**13753121**
Actinomyces naeslundii		100.0	0.46	IND 1		
13753122	**Low discrimination**					**13753122**
					SPO	
Actinomyces naeslundii		85.63	0.43	IND 1	1	
Clostridium ramosum		14.37	0.05	MNS 88	92	
13753123	**Good identification**					**13753123**
Actinomyces naeslundii		100.0	0.46	IND 1		
13755502	**Low discrimination**					**13755502**
					SPO	
Bifidobacterium adolescentis		85.83	0.40	IND 1	1	
Actinomyces israelii		10.18	0.30	MNS 87	1	
Clostridium butyricum		2.14	-.02		99	
13755512	**Good identification**					**13755512**
Bifidobacterium adolescentis		98.57	0.41	IND 1		
Actinomyces israelii		1.43	0.16	MNS 87		
13757100	**Good identification**					**13757100**
Actinomyces israelii		100.0	0.42	IND 1		
13757102	**Low discrimination**					**13757102**
Actinomyces israelii		87.04	0.45	IND 1		
Bifidobacterium adolescentis		12.96	0.24			
13757142	**Good identification**					**13757142**
Actinomyces israelii		95.48	0.38	IND 1		
Bifidobacterium adolescentis		4.52	0.09			
13757500	**Good identification**					**13757500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.40	IND 1		
13757502	**Low discrimination**					**13757502**
Bifidobacterium adolescentis		54.91	0.38	IND 1		
Actinomyces israelii		45.09	0.43	MNS 87		
13757512	**Acceptable identification**					**13757512**
Bifidobacterium adolescentis		90.88	0.40	IND 1		
Actinomyces israelii		9.13	0.29	MNS 87		
13757542	**Low discrimination**					**13757542**
Actinomyces israelii		72.10	0.36	IND 1		
Bifidobacterium adolescentis		27.90	0.23			
13771000	**Low discrimination**					**13771000**
					GRA	
Actinomyces naeslundii		65.65	0.35	IND 1	99	
Fusobacterium mortiferum		34.35	0.19	MNS 88	1	
13771001	**Good identification**					**13771001**
Actinomyces naeslundii		100.0	0.38	IND 1		
13771002	**Low discrimination**					**13771002**
Actinomyces naeslundii		68.99	0.35	IND 1		
Lactobacillus acidophilus		18.80	0.01	MNS 88		
Actinomyces israelii		12.21	0.13			
13771003	**Good identification**					**13771003**
Actinomyces naeslundii		100.0	0.38	IND 1		
13771020	**Good identification**					**13771020**
Actinomyces naeslundii		100.0	0.35	IND 1		
13771021	**Good identification**					**13771021**
Actinomyces naeslundii		100.0	0.38	IND 1		
13771022	**Low discrimination**					**13771022**
Actinomyces naeslundii		78.59	0.35	IND 1		
Lactobacillus acidophilus		21.41	0.01	MNS 88		
13771023	**Good identification**					**13771023**
Actinomyces naeslundii		100.0	0.38	IND 1		
13771100	**Low discrimination**					**13771100**
					GRA	
Actinomyces naeslundii		62.71	0.43	IND 1	99	
Fusobacterium mortiferum		26.90	0.25	MNS 88	1	
Actinomyces israelii		10.40	0.21		99	
13771101	**Good identification**					**13771101**
Actinomyces naeslundii		100.0	0.46	IND 1		
13771103	**Good identification**					**13771103**
Actinomyces naeslundii		100.0	0.46	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
13771120	**Low discrimination**					**13771120**
					SPO	
Actinomyces naeslundii		89.61	0.43	IND 1	1	
Clostridium ramosum		10.39	0.02	MNS 88	92	
13771121	**Good identification**					**13771121**
Actinomyces naeslundii		100.0	0.46	IND 1		
13771123	**Good identification**					**13771123**
Actinomyces naeslundii		100.0	0.46	IND 1		
13773000	**Low discrimination**					**13773000**
Actinomyces naeslundii		74.39	0.40	IND 1		
Actinomyces israelii		25.61	0.23	MNS 88		
13773001	**Good identification**					**13773001**
Actinomyces naeslundii		100.0	0.43	IND 1		
13773002	**Low discrimination**					**13773002**
Actinomyces naeslundii		65.94	0.40	IND 1		
Actinomyces israelii		34.06	0.26	MNS 88		
13773003	**Good identification**					**13773003**
Actinomyces naeslundii		100.0	0.43	IND 1		
13773020	**Good identification**					**13773020**
Actinomyces naeslundii		100.0	0.40	IND 1		
13773021	**Good identification**					**13773021**
Actinomyces naeslundii		100.0	0.43	IND 1		
13773022	**Good identification**					**13773022**
Actinomyces naeslundii		100.0	0.40	IND 1		
13773023	**Good identification**					**13773023**
Actinomyces naeslundii		100.0	0.43	IND 1		
13773100	**Low discrimination**					**13773100**
Actinomyces naeslundii		67.39	0.48	IND 1		
Actinomyces israelii		32.61	0.33	MNS 88		
13773101	**Good identification**					**13773101**
Actinomyces naeslundii		100.0	0.51	IND 1		
13773102	**Low discrimination**					**13773102**
Actinomyces naeslundii		56.86	0.48	IND 1		
Actinomyces israelii		41.27	0.36	MNS 88		
Bifidobacterium adolescentis		1.87	0.07			
13773103	**Good identification**					**13773103**
Actinomyces naeslundii		100.0	0.51	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
13773120	**Low discrimination**					**13773120**
					SPO	
Actinomyces naeslundii		74.71	0.48	IND 1	1	
Clostridium ramosum		25.29	0.15	MNS 88	92	
13773121	**Good identification**					**13773121**
Actinomyces naeslundii		100.0	0.51	IND 1		
13773123	**Good identification**					**13773123**
Actinomyces naeslundii		100.0	0.51	IND 1		
13773141	**Good identification**					**13773141**
Actinomyces naeslundii		100.0	0.38	IND 1		
13773143	**Good identification**					**13773143**
Actinomyces naeslundii		100.0	0.38	IND 1		
13773160	**Good identification**					**13773160**
Actinomyces naeslundii		100.0	0.35	IND 1		
13773161	**Good identification**					**13773161**
Actinomyces naeslundii		100.0	0.38	IND 1		
13773162	**Good identification**					**13773162**
Actinomyces naeslundii		100.0	0.35	IND 1		
13773163	**Good identification**					**13773163**
Actinomyces naeslundii		100.0	0.38	IND 1		
13775100	**Low discrimination**					**13775100**
					SPO	
Actinomyces israelii		76.51	0.39	IND 1	1	
Clostridium butyricum		23.49	0.09		99	
13775102	**Low discrimination**					**13775102**
					SPO	
Actinomyces israelii		79.22	0.42	IND 1	1	
Bifidobacterium adolescentis		14.47	0.23		1	
Clostridium butyricum		6.31	0.02		99	
13775502	**Low discrimination**					**13775502**
					SPO	
Bifidobacterium adolescentis		44.34	0.37	IND 1	1	
Actinomyces israelii		29.67	0.40	MNS 87	1	
Clostridium butyricum		25.99	0.18		99	
13775512	**Low discrimination**					**13775512**
Bifidobacterium adolescentis		88.32	0.38	IND 1		
Actinomyces israelii		7.23	0.26	MNS 87		
Clostridium clostridioforme		2.23	0.08			
13775722	**Low discrimination**					**13775722**
Bacteroides ovatus		85.97	0.49	ESL 91		
Bacteroides thetaiotaomicron		13.46	0.35	MNS 99		
Parabacteroides distasonis		0.29	0.22			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
13775732	**Acceptable identification**					**13775732**
Bacteroides ovatus		90.45	0.46	ESL 91		
Bacteroides thetaiotaomicron		8.26	0.28	MNS 99		
Clostridium clostridioforme		1.04	0.23			
13777000	**Good identification**					**13777000**
Actinomyces israelii		100.0	0.41	IND 1		
13777002	**Good identification**					**13777002**
Actinomyces israelii		100.0	0.44	IND 1		
13777042	**Good identification**					**13777042**
Actinomyces israelii		100.0	0.37	IND 1		
13777100	**Good identification**					**13777100**
					SPO	
Actinomyces israelii		99.40	0.52	IND 1	1	
Clostridium butyricum		0.60	-.08		99	
13777102	**Good identification**					**13777102**
Actinomyces israelii		97.43	0.55	IND 1		
Bifidobacterium adolescentis		2.57	0.21			
13777110	**Good identification**					**13777110**
Actinomyces israelii		100.0	0.37	IND 1		
13777112	**Low discrimination**					**13777112**
Actinomyces israelii		82.25	0.41	IND 1		
Bifidobacterium adolescentis		17.75	0.23	MLZ 13		
13777140	**Good identification**					**13777140**
Actinomyces israelii		100.0	0.45	IND 1		
13777142	**Good identification**					**13777142**
Actinomyces israelii		100.0	0.48	IND 1		
13777302	**Good identification**					**13777302**
Actinomyces israelii		100.0	0.35	IND 1		
13777400	**Good identification**					**13777400**
Actinomyces israelii		100.0	0.40	IND 1		
13777402	**Good identification**					**13777402**
Actinomyces israelii		100.0	0.43	IND 1		
13777442	**Good identification**					**13777442**
Actinomyces israelii		100.0	0.36	IND 1		
13777500	**Acceptable identification**					**13777500**
					SPO	
Actinomyces israelii		93.79	0.50	IND 1	1	
Clostridium butyricum		6.21	0.09		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
13777502	**Low discrimination**						**13777502**
					SPO		
Actinomyces israelii		81.11	0.53	IND 1	1		
Bifidobacterium adolescentis		17.50	0.35		1		
Clostridium butyricum		1.39	0.02		99		
13777510	**Good identification**						**13777510**
Actinomyces israelii		100.0	0.36	IND 1			
13777512	**Low discrimination**						**13777512**
Bifidobacterium adolescentis		62.56	0.37	IND 1			
Actinomyces israelii		35.45	0.39	MNS 87			
Bifidobacterium dentium		1.99	-.07				
13777540	**Good identification**						**13777540**
Actinomyces israelii		100.0	0.43	IND 1			
13777542	**Acceptable identification**						**13777542**
Actinomyces israelii		93.58	0.46	IND 1			
Bifidobacterium adolescentis		6.42	0.20				
13777722	**Good identification**						**13777722**
					GRA	CAT	
Bacteroides ovatus		99.35	0.41	ESL 91	1	68	
Bacteroides thetaiotaomicron		0.47	0.01	MNS 99	1	89	
Actinomyces israelii		0.18	0.13		99	1	
13777732	**Good identification**						**13777732**
Bacteroides ovatus		99.72	0.38	ESL 91			
Bacteroides thetaiotaomicron		0.28	-.06	MNS 99			
14000000	**Low discrimination**						**14000000**
					COC	GRA	
Fusobacterium nucleatum subsp. nucleatum		19.72	0.66	MNS 1	1	1	
Peptoniphilus asaccharolyticus		19.72	0.66		99	99	
Porphyromonas asaccharolytica		19.52	0.66		1	1	
14000001	**Very good identification**						**14000001**
					COC	GRA	
Anaerococcus prevotii		98.38	0.83	IND 9	99	99	
Fusobacterium nucleatum subsp. nucleatum		0.26	0.32		1	1	
Peptoniphilus asaccharolyticus		0.26	0.32		99	99	
14000002	**Low discrimination**						**14000002**
					COC	GRA	
Anaerococcus prevotii		52.04	0.56	IND 9	99	99	
Porphyromonas asaccharolytica		19.95	0.42	URE 90	1	1	
Parvimonas micra		5.04	0.33		99	99	
14000003	**Very good identification**						**14000003**
					COC	GRA	
Anaerococcus prevotii		99.85	0.73	IND 9	99	99	
Porphyromonas asaccharolytica		0.04	0.08		1	1	
Peptococcus niger		0.04	0.10		99	99	
14000011	**Good identification**						**14000011**
					COC	GRA	
Anaerococcus prevotii		98.45	0.49	IND 9	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	MLZ 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14000013	**Good identification**							**14000013**
Anaerococcus prevotii		100.0	0.38	IND 9				
14000020	**Good identification**							**14000020**
					GRA	SPO		
Clostridium cadaveris		92.40	0.75	GLU 95	99	99		
Clostridium bifermentans		2.01	0.49		99	99		
Fusobacterium nucleatum subsp. nucleatum		1.05	0.32		1	1		
14000021	**Low discrimination**							**14000021**
					COC	SPO		
Anaerococcus prevotii		79.57	0.49	IND 9	99	1		
Clostridium cadaveris		18.32	0.41	NAG 1	1	99		
Peptococcus niger		0.50	0.07		99	1		
14000022	**Low discrimination**							**14000022**
					GRA	SPO		
Clostridium cadaveris		49.51	0.41	GLU 95	99	99		
Pseudoflavonifractor capillosus		23.65	0.32	BGL 1	1	1		
Atopobium minutum		12.84	0.41		99	1		
14000023	**Good identification**							**14000023**
					COC	GRA	SPO	
Anaerococcus prevotii		97.99	0.38	IND 9	99	99	1	
Clostridium cadaveris		0.91	0.07	NAG 1	1	99	99	
Pseudoflavonifractor capillosus		0.44	-.03		1	1	1	
14000030	**Acceptable identification**							**14000030**
					GRA	SPO		
Clostridium cadaveris		92.69	0.41	GLU 95	99	99		
Clostridium bifermentans		2.02	0.15	MLZ 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.05	-.03		1	1		
14000040	**Low discrimination**							**14000040**
					COC	GRA	SPO	
Clostridium bifermentans		20.76	0.46	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		16.55	0.32	SOR 4	1	1	1	
Peptoniphilus asaccharolyticus		16.55	0.32		99	99	1	
14000041	**Good identification**							**14000041**
					COC	SPO		
Anaerococcus prevotii		97.57	0.49	IND 9	99	1		
Clostridium sordellii		0.87	0.15	SOR 1	1	99		
Clostridium bifermentans		0.32	0.12		1	99		
14000043	**Good identification**							**14000043**
Anaerococcus prevotii		100.0	0.38	IND 9				
14000060	**Low discrimination**							**14000060**
					GRA	SPO		
Clostridium cadaveris		86.93	0.41	GLU 95	99	99		
Clostridium bifermentans		7.81	0.25	SOR 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.99	-.03		1	1		
14000101	**Good identification**							**14000101**
					COC	GRA		
Anaerococcus prevotii		98.45	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	LAC 1	1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14000103	**Good identification**							**14000103**
Anaerococcus prevotii		100.0	0.38	IND 9				
14000120	**Acceptable identification**							**14000120**
					GRA	SPO		
Clostridium cadaveris		91.93	0.41	GLU 95	99	99		
Clostridium bifermentans		2.00	0.15	LAC 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.04	-.03		1	1		
14000201	**Good identification**							**14000201**
					COC	GRA		
Anaerococcus prevotii		98.45	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	RHA 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14000203	**Good identification**							**14000203**
Anaerococcus prevotii		100.0	0.38	IND 9				
14000220	**Acceptable identification**							**14000220**
					GRA	SPO		
Clostridium cadaveris		92.69	0.41	GLU 95	99	99		
Clostridium bifermentans		2.02	0.15	RHA 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.05	-.03		1	1		
14000401	**Good identification**							**14000401**
					COC	GRA		
Anaerococcus prevotii		98.45	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	ARA 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14000403	**Good identification**							**14000403**
Anaerococcus prevotii		100.0	0.38	IND 9				
14000420	**Acceptable identification**							**14000420**
					GRA	SPO		
Clostridium cadaveris		92.69	0.41	GLU 95	99	99		
Clostridium bifermentans		2.02	0.15	ARA 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.05	-.03		1	1		
14001000	**Good identification**							**14001000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		90.13	0.66	MNS 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		1.99	0.32		1	1		
Peptoniphilus asaccharolyticus		1.99	0.32		99	99		
14001001	**Low discrimination**							**14001001**
					COC	GRA	SPO	
Anaerococcus prevotii		85.51	0.49	IND 9	99	99	1	
Fusobacterium necrophorum subsp. necrophorum		10.12	0.32	GAL 1	1	1	1	
Clostridium sordellii		3.24	0.25		1	99	99	
14001003	**Good identification**							**14001003**
					COC	GRA		
Anaerococcus prevotii		99.29	0.38	IND 9	99	99		
Fusobacterium necrophorum subsp. necrophorum		0.48	-.03	GAL 1	1	1		
Atopobium minutum		0.24	0.07		1	99		
14001020	**Low discrimination**							**14001020**
					GRA	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium cadaveris		57.82	0.41	GLU 95	99	99		
Fusobacterium necrophorum subsp. necrophorum		29.71	0.32	GAL 1	1	1		
Atopobium minutum		7.50	0.36		99	1		
14002001	**Good identification**							**14002001**
					COC	GRA		
Anaerococcus prevotii		98.45	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	MAN 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14002003	**Good identification**							**14002003**
Anaerococcus prevotii		100.0	0.38	IND 9				
14002020	**Acceptable identification**							**14002020**
					GRA	SPO		
Clostridium cadaveris		91.92	0.41	GLU 95	99	99		
Clostridium bifermentans		2.00	0.15	MAN 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.04	-.03		1	1		
14004001	**Good identification**							**14004001**
					COC	GRA		
Anaerococcus prevotii		98.33	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	XYL 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14004003	**Good identification**							**14004003**
Anaerococcus prevotii		100.0	0.38	IND 9				
14004020	**Acceptable identification**							**14004020**
					GRA	SPO		
Clostridium cadaveris		92.20	0.41	GLU 95	99	99		
Clostridium bifermentans		2.01	0.15	XYL 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.04	-.03		1	1		
14010000	**Very good identification**							**14010000**
					GRA	SPO		
Fusobacterium varium		99.25	0.91		1	1		
Clostridium bifermentans		0.13	0.47		99	99		
Atopobium minutum		0.08	0.52		99	1		
14010001	**Low discrimination**							**14010001**
					COC	GRA	SPO	
Fusobacterium varium		71.78	0.57	URE 1	1	1	1	
Anaerococcus prevotii		22.77	0.49		99	99	1	
Clostridium sordellii		4.89	0.38		1	99	99	
14010002	**Good identification**							**14010002**
					COC	GRA		
Fusobacterium varium		93.35	0.57	BGL 1	1	1		
Atopobium minutum		4.90	0.49		1	99		
Anaerococcus prevotii		0.82	0.22		99	99		
14010003	**Low discrimination**							**14010003**
					COC	GRA	SPO	
Anaerococcus prevotii		87.51	0.38	IND 9	99	99	1	
Fusobacterium varium		11.15	0.23	FRU 1	1	1	1	
Clostridium sordellii		0.76	0.04		1	99	99	
14010010	**Very good identification**							**14010010**
					GRA	SPO		
Fusobacterium varium		99.19	0.57	MLZ 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		0.13	0.13		99	99		
Atopobium minutum		0.08	0.18		99	1		
14010020	**Low discrimination**							**14010020**
					GRA	SPO	CAT	
Clostridium cadaveris		86.21	0.75	GLU 95	99	99	1	
Fusobacterium varium		11.92	0.57		1	1	1	
Propionibacterium acnes		1.40	0.48		99	1	95	
14010021	**Low discrimination**							**14010021**
					COC	GRA	SPO	
Clostridium cadaveris		82.55	0.41	GLU 95	1	99	99	
Fusobacterium varium		11.42	0.23	URE 1	1	1	1	
Anaerococcus prevotii		3.62	0.15		99	99	1	
14010022	**Low discrimination**							**14010022**
					GRA	SPO		
Clostridium cadaveris		69.11	0.41	GLU 95	99	99		
Atopobium minutum		16.54	0.41	BGL 1	99	1		
Fusobacterium varium		9.56	0.23		1	1		
14010030	**Low discrimination**							**14010030**
					GRA	SPO	CAT	
Clostridium cadaveris		86.35	0.41	GLU 95	99	99	1	
Fusobacterium varium		11.94	0.23	MLZ 1	1	1	1	
Propionibacterium acnes		1.40	0.14		99	1	95	
14010040	**Very good identification**							**14010040**
					GRA	SPO		
Fusobacterium varium		98.84	0.57	SOR 1	1	1		
Clostridium bifermentans		0.54	0.24		99	99		
Atopobium minutum		0.08	0.18		99	1		
14010060	**Low discrimination**							**14010060**
					GRA	SPO	CAT	
Clostridium cadaveris		66.72	0.41	GLU 95	99	99	1	
Propionibacterium acnes		23.50	0.36	SOR 1	99	1	95	
Fusobacterium varium		9.23	0.23		1	1	1	
14010100	**Very good identification**							**14010100**
					GRA	SPO		
Fusobacterium varium		99.27	0.57	LAC 1	1	1		
Clostridium bifermentans		0.13	0.13		99	99		
Atopobium minutum		0.08	0.18		99	1		
14010120	**Low discrimination**							**14010120**
					GRA	SPO	CAT	
Clostridium cadaveris		86.35	0.41	GLU 95	99	99	1	
Fusobacterium varium		11.94	0.23	LAC 1	1	1	1	
Propionibacterium acnes		1.40	0.14		99	1	95	
14010200	**Very good identification**							**14010200**
					GRA	SPO		
Fusobacterium varium		99.27	0.57	RHA 1	1	1		
Clostridium bifermentans		0.13	0.13		99	99		
Atopobium minutum		0.08	0.18		99	1		
14010220	**Low discrimination**							**14010220**
					GRA	SPO	CAT	
Clostridium cadaveris		86.35	0.41	GLU 95	99	99	1	
Fusobacterium varium		11.94	0.23	RHA 1	1	1	1	
Propionibacterium acnes		1.40	0.14		99	1	95	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
14010400	**Very good identification**							**14010400**
					GRA	SPO		
Fusobacterium varium		99.25	0.57	ARA 1	1	1		
Clostridium bifermentans		0.13	0.13		99	99		
Atopobium minutum		0.08	0.18		99	1		
14010420	**Low discrimination**							**14010420**
					GRA	SPO	CAT	
Clostridium cadaveris		86.35	0.41	GLU 95	99	99	1	
Fusobacterium varium		11.94	0.23	ARA 1	1	1	1	
Propionibacterium acnes		1.40	0.14		99	1	95	
14011000	**Low discrimination**							**14011000**
Fusobacterium varium		86.70	0.57	GAL 1				
Prevotella intermedia		7.12	0.47					
Fusobacterium necrophorum subsp. necrophorum		3.25	0.32					
14011002	**Low discrimination**							**14011002**
					GRA			
Atopobium minutum		60.72	0.41	IND 1	99			
Fusobacterium varium		35.09	0.23		1			
Prevotella intermedia		2.88	0.12		1			
14011020	**Low discrimination**							**14011020**
					GRA	SPO	CAT	
Propionibacterium acnes		55.73	0.48	NIT 92	99	1	95	
Clostridium cadaveris		34.73	0.41	GLU 99	99	99	1	
Fusobacterium varium		4.80	0.23		1	1	1	
14011060	**Good identification**							**14011060**
					SPO	CAT		
Propionibacterium acnes		97.21	0.36	NIT 92	1	95		
Clostridium cadaveris		2.79	0.07	GLU 99	99	1		
14012000	**Very good identification**							**14012000**
					GRA	SPO		
Fusobacterium varium		97.33	0.57	MAN 1	1	1		
Eubacterium limosum		1.73	0.29		99	1		
Clostridium difficile		0.14	0.12		99	99		
14012020	**Low discrimination**							**14012020**
					GRA	SPO	CAT	
Clostridium cadaveris		48.85	0.41	GLU 95	99	99	1	
Propionibacterium acnes		44.10	0.43	MAN 1	99	1	95	
Fusobacterium varium		6.76	0.23		1	1	1	
14013020	**Good identification**							**14013020**
					SPO	CAT		
Propionibacterium acnes		98.89	0.43	NIT 92	1	95		
Clostridium cadaveris		1.11	0.07	GLU 99	99	1		
14014000	**Very good identification**							**14014000**
					GRA	SPO		
Fusobacterium varium		99.82	0.67	XYL 4	1	1		
Clostridium bifermentans		0.03	0.13		99	99		
Atopobium minutum		0.02	0.18		99	1		
14014020	**Low discrimination**							**14014020**
					GRA	SPO	CAT	
Clostridium cadaveris		62.88	0.41	GLU 95	99	99	1	
Fusobacterium varium		35.87	0.33	XYL 1	1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		1.02	0.14		99	1	95	
14020001	**Good identification**							**14020001**
					COC	GRA		
Anaerococcus prevotii		98.45	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	TRE 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14020003	**Good identification**							**14020003**
Anaerococcus prevotii		100.0	0.38	IND 9				
14020020	**Good identification**							**14020020**
					GRA	SPO		
Clostridium cadaveris		98.51	0.53	GLU 95	99	99		
Clostridium bifermentans		0.41	0.15	TRE 5	99	99		
Porphyromonas asaccharolytica		0.21	-.03		1	1		
14030000	**Very good identification**							**14030000**
					GRA	SPO		
Fusobacterium varium		98.94	0.57	TRE 1	1	1		
Clostridium cadaveris		0.38	0.19		99	99		
Clostridium bifermentans		0.13	0.13		99	99		
14030020	**Good identification**							**14030020**
					GRA	SPO	CAT	
Clostridium cadaveris		97.06	0.53	GLU 95	99	99	1	
Fusobacterium varium		2.58	0.23	TRE 5	1	1	1	
Propionibacterium acnes		0.30	0.14		99	1	95	
14040001	**Good identification**							**14040001**
					COC	GRA		
Anaerococcus prevotii		98.45	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	CEL 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
14040003	**Good identification**							**14040003**
Anaerococcus prevotii		100.0	0.38	IND 9				
14040020	**Acceptable identification**							**14040020**
					GRA	SPO		
Clostridium cadaveris		92.69	0.41	GLU 95	99	99		
Clostridium bifermentans		2.02	0.15	CEL 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.05	-.03		1	1		
14050000	**Very good identification**							**14050000**
					GRA	SPO		
Fusobacterium varium		99.27	0.57	CEL 1	1	1		
Clostridium bifermentans		0.13	0.13		99	99		
Atopobium minutum		0.08	0.18		99	1		
14050020	**Low discrimination**							**14050020**
					GRA	SPO	CAT	
Clostridium cadaveris		86.35	0.41	GLU 95	99	99	1	
Fusobacterium varium		11.94	0.23	CEL 1	1	1	1	
Propionibacterium acnes		1.40	0.14		99	1	95	
14100000	**Good identification**							**14100000**
					GRA	SPO		
Clostridium bifermentans		90.73	0.76	GLU 92	99	99		
Prevotella intermedia		1.29	0.47		1	1		
Fusobacterium nucleatum subsp. nucleatum		1.29	0.32		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
14100001	**Low discrimination**							**14100001**
					COC	SPO		
Clostridium sordellii		61.67	0.55	GLU 95	1	99		
Anaerococcus prevotii		31.90	0.49	MNS 4	99	1		
Clostridium bifermentans		5.85	0.41		1	99		
14100002	**Low discrimination**							**14100002**
					COC	GRA	SPO	
Clostridium bifermentans		75.94	0.41	GLU 92	1	99	99	
Anaerococcus prevotii		11.51	0.22	BGL 1	99	99	1	
Porphyromonas asaccharolytica		4.41	0.08		1	1	1	
14100003	**Acceptable identification**							**14100003**
					COC	SPO		
Anaerococcus prevotii		92.12	0.38	IND 9	99	1		
Clostridium sordellii		7.20	0.20	MLT 1	1	99		
Clostridium bifermentans		0.68	0.07		1	99		
14100010	**Acceptable identification**							**14100010**
					GRA	SPO		
Clostridium bifermentans		91.43	0.41	GLU 92	99	99		
Prevotella intermedia		1.30	0.12	MLZ 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		1.30	-.03		1	1		
14100020	**Low discrimination**							**14100020**
					COC	SPO		
Clostridium cadaveris		50.38	0.53	GLU 95	1	99		
Clostridium bifermentans		48.63	0.55	MLT 5	1	99		
Gemella morbillorum		0.12	0.14		99	1		
14100040	**Good identification**							**14100040**
					GRA	SPO		
Clostridium bifermentans		96.97	0.52	GLU 92	99	99		
Clostridium botulinum biovar A		0.64	0.15	SOR 4	99	99		
Prevotella intermedia		0.34	0.12		1	1		
14100100	**Acceptable identification**							**14100100**
					GRA	SPO		
Clostridium bifermentans		90.49	0.41	GLU 92	99	99		
Prevotella intermedia		1.29	0.12	LAC 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		1.29	-.03		1	1		
14100200	**Acceptable identification**							**14100200**
					GRA	SPO		
Clostridium bifermentans		91.43	0.41	GLU 92	99	99		
Prevotella intermedia		1.30	0.12	RHA 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		1.30	-.03		1	1		
14100400	**Acceptable identification**							**14100400**
					COC	GRA	SPO	
Clostridium bifermentans		90.15	0.41	GLU 92	1	99	99	
Gemella morbillorum		1.40	0.14	ARA 1	99	99	1	
Prevotella intermedia		1.29	0.12		1	1	1	
14101000	**Low discrimination**							**14101000**
					GRA	SPO		
Prevotella intermedia		43.73	0.47	GLU 90	1	1		
Clostridium bifermentans		30.96	0.41	FRU 99	99	99		
Fusobacterium necrophorum subsp. necrophorum		19.99	0.32		1	1		
14101001	**Good identification**							**14101001**
					COC	GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sordellii		95.58	0.42	GLU 95	1	99	99	
Anaerococcus prevotii		2.83	0.15	MNS 4	99	99	1	
Prevotella intermedia		0.73	0.12		1	1	1	
14102000	**Acceptable identification**							**14102000**
					GRA	SPO		
Clostridium bifermentans		91.43	0.41	GLU 92	99	99		
Prevotella intermedia		1.30	0.12	MAN 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		1.30	-.03		1	1		
14104000	**Low discrimination**							**14104000**
					COC	GRA	SPO	
Clostridium bifermentans		89.57	0.41	GLU 92	1	99	99	
Gemella morbillorum		1.39	0.14	XYL 1	99	99	1	
Prevotella intermedia		1.28	0.12		1	1	1	
14105722	**Good identification**							**14105722**
Bacteroides eggerthii		100.0	0.39	ESL 90				
14110000	**Low discrimination**							**14110000**
					GRA	SPO		
Prevotella intermedia		85.67	0.81	GLU 90	1	1		
Fusobacterium varium		10.54	0.57		1	1		
Clostridium bifermentans		3.19	0.54		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
14110001	**Low discrimination**						**14110001**
					GRA	SPO	
Clostridium sordellii		86.13	0.55	GLU 95	99	99	
Prevotella intermedia		11.53	0.47	MNS 4	1	1	
Fusobacterium varium		1.42	0.23		1	1	
14110002	**Low discrimination**						**14110002**
					GRA	SPO	
Prevotella intermedia		85.42	0.47	GLU 90	1	1	
Fusobacterium varium		10.51	0.23	BGL 1	1	1	
Clostridium bifermentans		3.18	0.20		99	99	
14110010	**Low discrimination**						**14110010**
					GRA	SPO	
Prevotella intermedia		85.90	0.47	GLU 90	1	1	
Fusobacterium varium		10.57	0.23	MLZ 1	1	1	
Clostridium bifermentans		3.20	0.20		99	99	
14110020	**Low discrimination**						**14110020**
					GRA	SPO	
Clostridium cadaveris		76.89	0.53	GLU 95	99	99	
Prevotella intermedia		16.59	0.47	MLT 5	1	1	
Clostridium bifermentans		3.91	0.33		99	99	
14110040	**Low discrimination**						**14110040**
					GRA	SPO	
Prevotella intermedia		77.32	0.47	GLU 90	1	1	
Clostridium bifermentans		11.88	0.30	SOR 1	99	99	
Fusobacterium varium		9.51	0.23		1	1	
14110100	**Low discrimination**						**14110100**
					GRA	SPO	
Prevotella intermedia		85.90	0.47	GLU 90	1	1	
Fusobacterium varium		10.57	0.23	LAC 1	1	1	
Clostridium bifermentans		3.20	0.20		99	99	
14110200	**Low discrimination**						**14110200**
					GRA	SPO	
Prevotella intermedia		85.90	0.47	GLU 90	1	1	
Fusobacterium varium		10.57	0.23	RHA 1	1	1	
Clostridium bifermentans		3.20	0.20		99	99	
14110400	**Low discrimination**						**14110400**
					GRA	SPO	
Prevotella intermedia		84.92	0.47	GLU 90	1	1	
Fusobacterium varium		10.45	0.23	ARA 1	1	1	
Clostridium bifermentans		3.16	0.20		99	99	
14111000	**Very good identification**						**14111000**
					COC	GRA	
Prevotella intermedia		99.75	0.81	GLU 90	1	1	
Fusobacterium varium		0.12	0.23		1	1	
Gemella morbillorum		0.07	0.27		99	99	
14111001	**Low discrimination**						**14111001**
					GRA	SPO	
Clostridium sordellii		56.86	0.42	GLU 95	99	99	
Prevotella intermedia		43.14	0.47	MNS 4	1	1	
14111002	**Good identification**						**14111002**
					GRA		
Prevotella intermedia		99.79	0.47	GLU 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium minutum		0.22	0.07	BGL 1	99			
14111010	**Good identification**							**14111010**
Prevotella intermedia		100.0	0.47	GLU 90				
14111020	**Acceptable identification**							**14111020**
					GRA	SPO	CAT	
Prevotella intermedia		93.38	0.47	GLU 90	1	1	1	
Clostridium cadaveris		4.37	0.19	NAG 1	99	99	1	
Propionibacterium acnes		1.35	0.14		99	1	95	
14111040	**Good identification**							**14111040**
Prevotella intermedia		100.0	0.47	GLU 90				
14111100	**Good identification**							**14111100**
Prevotella intermedia		100.0	0.47	GLU 90				
14111200	**Good identification**							**14111200**
Prevotella intermedia		100.0	0.47	GLU 90				
14111400	**Good identification**							**14111400**
					COC	GRA		
Prevotella intermedia		99.28	0.47	GLU 90	1	1		
Gemella morbillorum		0.72	0.11	ARA 1	99	99		
14112000	**Low discrimination**							**14112000**
					GRA	SPO		
Prevotella intermedia		85.74	0.47	GLU 90	1	1		
Fusobacterium varium		10.55	0.23	MAN 1	1	1		
Clostridium bifermentans		3.20	0.20		99	99		
14113000	**Good identification**							**14113000**
Prevotella intermedia		100.0	0.47	GLU 90				
14114000	**Low discrimination**							**14114000**
					GRA	SPO		
Prevotella intermedia		64.02	0.47	GLU 90	1	1		
Fusobacterium varium		32.49	0.33	XYL 1	1	1		
Clostridium bifermentans		2.39	0.20		99	99		
14115000	**Good identification**							**14115000**
					COC	GRA		
Prevotella intermedia		98.78	0.47	GLU 90	1	1		
Gemella morbillorum		0.72	0.11	XYL 1	99	99		
Fusobacterium varium		0.51	-.01		1	1		
14120000	**Low discrimination**							**14120000**
					GRA	SPO		
Clostridium bifermentans		88.99	0.41	GLU 92	99	99		
Clostridium botulinum biovar A		2.04	0.14	TRE 1	99	99		
Prevotella intermedia		1.27	0.12		1	1		
14130000	**Low discrimination**							**14130000**
					GRA	SPO		
Prevotella intermedia		85.72	0.47	GLU 90	1	1		
Fusobacterium varium		10.54	0.23	TRE 1	1	1		
Clostridium bifermentans		3.19	0.20		99	99		
14131000	**Good identification**							**14131000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		100.0	0.47	GLU 90				
14140000	**Acceptable identification**							**14140000**
					GRA	SPO		
Clostridium bifermentans		91.43	0.41	GLU 92	99	99		
Prevotella intermedia		1.30	0.12	CEL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		1.30	-.03		1	1		
14145722	**Good identification**							**14145722**
Bacteroides eggerthii		100.0	0.50	ESL 90				
14150000	**Low discrimination**							**14150000**
					GRA	SPO		
Prevotella intermedia		85.90	0.47	GLU 90	1	1		
Fusobacterium varium		10.57	0.23	CEL 1	1	1		
Clostridium bifermentans		3.20	0.20		99	99		
14151000	**Good identification**							**14151000**
Prevotella intermedia		100.0	0.47	GLU 90				
14200000	**Low discrimination**							**14200000**
					COC	GRA	SPO	
Clostridium bifermentans		41.25	0.53	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		12.33	0.32	SAL 10	1	1	1	
Peptoniphilus asaccharolyticus		12.33	0.32		99	99	1	
14200001	**Good identification**							**14200001**
					COC	GRA	SPO	
Anaerococcus prevotii		97.69	0.49	IND 9	99	99	1	
Clostridium bifermentans		0.85	0.19	SAL 1	1	99	99	
Porphyromonas asaccharolytica		0.25	-.03		1	1	1	
14200003	**Good identification**							**14200003**
Anaerococcus prevotii		100.0	0.38	IND 9				
14200020	**Low discrimination**							**14200020**
					GRA	SPO		
Clostridium cadaveris		77.13	0.41	GLU 95	99	99		
Clostridium bifermentans		18.47	0.33	SAL 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.87	-.03		1	1		
14210000	**Very good identification**							**14210000**
					GRA	SPO		
Fusobacterium varium		98.00	0.57	SAL 1	1	1		
Clostridium bifermentans		1.41	0.31		99	99		
Atopobium minutum		0.08	0.18		99	1		
14210020	**Low discrimination**							**14210020**
					GRA	SPO	CAT	
Clostridium cadaveris		85.42	0.41	GLU 95	99	99	1	
Fusobacterium varium		11.81	0.23	SAL 1	1	1	1	
Propionibacterium acnes		1.38	0.14		99	1	95	
14300000	**Very good identification**							**14300000**
					GRA	SPO		
Clostridium bifermentans		99.16	0.59	GLU 92	99	99		
Prevotella intermedia		0.13	0.12	SAL 10	1	1		
Porphyromonas asaccharolytica		0.13	-.03		1	1		
14300020	**Acceptable identification**							**14300020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		91.39	0.39	GLU 92				
Clostridium cadaveris		8.61	0.19	SAL 10				
14300040	**Good identification**							**14300040**
Clostridium bifermentans		100.0	0.36	GLU 92				
14310000	**Low discrimination**							**14310000**
					GRA	SPO		
Prevotella intermedia		65.07	0.47	GLU 90	1	1		
Clostridium bifermentans		26.67	0.37	SAL 1	99	99		
Fusobacterium varium		8.00	0.23		1	1		
14311000	**Good identification**							**14311000**
					GRA	SPO		
Prevotella intermedia		99.59	0.47	GLU 90	1	1		
Clostridium bifermentans		0.41	0.03	SAL 1	99	99		
14400000	**Low discrimination**							**14400000**
					COC	GRA		
Anaerococcus prevotii		75.32	0.58	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		5.30	0.32	URE 90	1	1		
Peptoniphilus asaccharolyticus		5.30	0.32		99	99		
14400001	**Very good identification**							**14400001**
					COC	GRA	SPO	
Anaerococcus prevotii		99.95	0.75	IND 9	99	99	1	
Porphyromonas asaccharolytica		0.01	-.03		1	1	1	
Clostridium sordellii		0.01	0.04		1	99	99	
14400002	**Good identification**							**14400002**
					COC	GRA		
Anaerococcus prevotii		97.36	0.48	IND 9	99	99		
Porphyromonas asaccharolytica		1.13	0.08	URE 90	1	1		
Parvimonas micra		0.29	-.01		99	99		
14400003	**Very good identification**							**14400003**
Anaerococcus prevotii		100.0	0.64	IND 9				
14400011	**Good identification**							**14400011**
Anaerococcus prevotii		100.0	0.41	IND 9				
14400020	**Low discrimination**							**14400020**
					COC	SPO		
Clostridium cadaveris		80.98	0.41	GLU 95	1	99		
Anaerococcus prevotii		13.03	0.24	RAF 1	99	1		
Clostridium bifermentans		1.76	0.15		1	99		
14400021	**Good identification**							**14400021**
					COC	SPO		
Anaerococcus prevotii		99.31	0.41	IND 9	99	1		
Clostridium cadaveris		0.69	0.07	NAG 1	1	99		
14400041	**Good identification**							**14400041**
Anaerococcus prevotii		100.0	0.41	IND 9				
14400101	**Good identification**							**14400101**
Anaerococcus prevotii		100.0	0.41	IND 9				
14400201	**Good identification**							**14400201**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Anaerococcus prevotii		100.0	0.41	IND 9				
14400401	**Good identification**							**14400401**
Anaerococcus prevotii		100.0	0.41	IND 9				
14401001	**Good identification**							**14401001**
					COC	GRA		
Anaerococcus prevotii		99.64	0.41	IND 9	99	99		
Fusobacterium necrophorum subsp. necrophorum		0.36	-.03	GAL 1	1	1		
14402001	**Good identification**							**14402001**
Anaerococcus prevotii		100.0	0.41	IND 9				
14404001	**Good identification**							**14404001**
Anaerococcus prevotii		100.0	0.41	IND 9				
14410000	**Low discrimination**							**14410000**
					COC	GRA		
Fusobacterium varium		87.64	0.57	RAF 1	1	1		
Prevotella intermedia		10.79	0.50		1	1		
Anaerococcus prevotii		1.03	0.24		99	99		
14410001	**Low discrimination**							**14410001**
					COC	GRA		
Anaerococcus prevotii		89.78	0.41	IND 9	99	99		
Fusobacterium varium		8.58	0.23	FRU 1	1	1		
Prevotella intermedia		1.06	0.15		1	1		
14410020	**Low discrimination**							**14410020**
					GRA	SPO		
Clostridium cadaveris		85.10	0.41	GLU 95	99	99		
Fusobacterium varium		11.77	0.23	RAF 1	1	1		
Prevotella intermedia		1.45	0.15		1	1		
14411000	**Acceptable identification**							**14411000**
Prevotella intermedia		91.98	0.50	GLU 90				
Fusobacterium varium		7.54	0.23	MLT 99				
Fusobacterium necrophorum subsp. necrophorum		0.28	-.03					
14420001	**Good identification**							**14420001**
Anaerococcus prevotii		100.0	0.41	IND 9				
14440001	**Good identification**							**14440001**
Anaerococcus prevotii		100.0	0.41	IND 9				
14500000	**Low discrimination**							**14500000**
					COC	GRA	SPO	
Prevotella intermedia		62.42	0.50	GLU 90	1	1	1	
Clostridium bifermentans		29.46	0.41	FRU 99	1	99	99	
Anaerococcus prevotii		5.96	0.24		99	99	1	
14500001	**Acceptable identification**							**14500001**
					COC	GRA	SPO	
Anaerococcus prevotii		92.95	0.41	IND 9	99	99	1	
Clostridium sordellii		5.45	0.20	MLT 1	1	99	99	
Prevotella intermedia		1.09	0.15		1	1	1	
14501000	**Good identification**							**14501000**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		99.22	0.50	GLU 90	1	1		
Clostridium bifermentans		0.47	0.07	FRU 99	99	99		
Fusobacterium necrophorum subsp. necrophorum		0.31	-.03		1	1		
14510000	**Very good identification**							**14510000**
					GRA	SPO		
Prevotella intermedia		99.89	0.84	GLU 90	1	1		
Fusobacterium varium		0.08	0.23		1	1		
Clostridium bifermentans		0.03	0.20		99	99		
14510001	**Acceptable identification**							**14510001**
					COC	GRA	SPO	
Prevotella intermedia		94.43	0.50	GLU 90	1	1	1	
Clostridium sordellii		4.75	0.20	URE 1	1	99	99	
Anaerococcus prevotii		0.82	0.06		99	99	1	
14510002	**Good identification**							**14510002**
Prevotella intermedia		100.0	0.50	GLU 90				
14510010	**Good identification**							**14510010**
Prevotella intermedia		100.0	0.50	GLU 90				
14510020	**Good identification**							**14510020**
					GRA	SPO		
Prevotella intermedia		96.82	0.50	GLU 90	1	1		
Clostridium cadaveris		3.02	0.19	NAG 1	99	99		
Clostridium bifermentans		0.15	-.01		99	99		
14510040	**Good identification**							**14510040**
Prevotella intermedia		100.0	0.50	GLU 90				
14510100	**Good identification**							**14510100**
Prevotella intermedia		100.0	0.50	GLU 90				
14510200	**Good identification**							**14510200**
Prevotella intermedia		100.0	0.50	GLU 90				
14510400	**Good identification**							**14510400**
Prevotella intermedia		100.0	0.50	GLU 90				
14511000	**Very good identification**							**14511000**
Prevotella intermedia		100.0	0.84	GLU 90				
14511001	**Good identification**							**14511001**
					GRA	SPO		
Prevotella intermedia		99.12	0.50	GLU 90	1	1		
Clostridium sordellii		0.88	0.07	URE 1	99	99		
14511002	**Good identification**							**14511002**
Prevotella intermedia		100.0	0.50	GLU 90				
14511010	**Good identification**							**14511010**
Prevotella intermedia		100.0	0.50	GLU 90				
14511020	**Good identification**							**14511020**
Prevotella intermedia		100.0	0.50	GLU 90				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
14511040	**Good identification**						**14511040**
Prevotella intermedia		100.0	0.50	GLU 90			
14511100	**Good identification**						**14511100**
Prevotella intermedia		100.0	0.50	GLU 90			
14511200	**Good identification**						**14511200**
Prevotella intermedia		100.0	0.50	GLU 90			
14511400	**Good identification**						**14511400**
Prevotella intermedia		100.0	0.50	GLU 90			
14512000	**Good identification**						**14512000**
Prevotella intermedia		100.0	0.50	GLU 90			
14513000	**Good identification**						**14513000**
Prevotella intermedia		100.0	0.50	GLU 90			
14514000	**Good identification**						**14514000**
Prevotella intermedia		99.66	0.50	GLU 90			
Fusobacterium varium		0.34	-.01	XYL 1			
14515000	**Good identification**						**14515000**
Prevotella intermedia		100.0	0.50	GLU 90			
14530000	**Good identification**						**14530000**
Prevotella intermedia		100.0	0.50	GLU 90			
14531000	**Good identification**						**14531000**
Prevotella intermedia		100.0	0.50	GLU 90			
14550000	**Good identification**						**14550000**
Prevotella intermedia		100.0	0.50	GLU 90			
14551000	**Good identification**						**14551000**
Prevotella intermedia		100.0	0.50	GLU 90			
14600001	**Good identification**						**14600001**
Anaerococcus prevotii		100.0	0.41	IND 9			
14710000	**Good identification**						**14710000**
					GRA	SPO	
Prevotella intermedia		99.73	0.50	GLU 90	1	1	
Clostridium bifermentans		0.28	0.03	SAL 1	99	99	
14711000	**Good identification**						**14711000**
Prevotella intermedia		100.0	0.50	GLU 90			
15000000	**Low discrimination**						**15000000**
					GRA	SPO	
Clostridium bifermentans		82.14	0.87		99	99	
Fusobacterium varium		9.69	0.66		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium cadaveris		4.02	0.63		99	99	
15000001	**Low discrimination**						**15000001**
					COC	SPO	
Clostridium sordellii		65.94	0.60	MNS 4	1	99	
Anaerococcus prevotii		16.15	0.49	MLT 90	99	1	
Clostridium bifermentans		14.59	0.53		1	99	
15000002	**Low discrimination**						**15000002**
					GRA	SPO	
Clostridium bifermentans		62.70	0.53	BGL 1	99	99	
Atopobium minutum		21.17	0.53		99	1	
Fusobacterium varium		7.40	0.32		1	1	
15000003	**Low discrimination**						**15000003**
					COC	SPO	
Anaerococcus prevotii		81.65	0.38	IND 9	99	1	
Clostridium sordellii		13.47	0.26	GLU 1	1	99	
Clostridium bifermentans		2.98	0.19		1	99	
15000010	**Low discrimination**						**15000010**
					GRA	SPO	
Clostridium bifermentans		81.36	0.53	MLZ 1	99	99	
Fusobacterium varium		9.60	0.32		1	1	
Clostridium cadaveris		3.98	0.29		99	99	
15000020	**Very good identification**						**15000020**
					SPO	CAT	
Clostridium cadaveris		98.54	0.97		99	1	
Clostridium bifermentans		1.30	0.67		99	1	
Propionibacterium acnes		0.08	0.48		1	95	
15000021	**Very good identification**						**15000021**
Clostridium cadaveris		97.37	0.63	URE 1			
Clostridium bifermentans		1.28	0.33				
Clostridium sordellii		0.92	0.26				
15000022	**Very good identification**						**15000022**
					SPO		
Clostridium cadaveris		96.35	0.63	BGL 1	99		
Atopobium minutum		2.24	0.45		1		
Clostridium bifermentans		1.27	0.33		99		
15000030	**Very good identification**						**15000030**
					SPO	CAT	
Clostridium cadaveris		98.50	0.63	MLZ 1	99	1	
Clostridium bifermentans		1.30	0.33		99	1	
Propionibacterium acnes		0.08	0.14		1	95	
15000040	**Good identification**						**15000040**
					GRA	SPO	
Clostridium bifermentans		93.53	0.64	SOR 4	99	99	
Fusobacterium varium		2.68	0.32		1	1	
Clostridium botulinum biovar A		1.39	0.33		99	99	
15000041	**Low discrimination**						**15000041**
					COC	SPO	
Clostridium sordellii		75.35	0.36	MNS 4	1	99	
Clostridium bifermentans		16.67	0.30	MLT 90	1	99	
Anaerococcus prevotii		4.48	0.15	SOR 4	99	1	
15000060	**Good identification**						**15000060**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	CAT		
Clostridium cadaveris		93.10	0.63	SOR 1	99	1		
Clostridium bifermentans		5.06	0.43		99	1		
Propionibacterium acnes		1.73	0.36		1	95		
15000100	**Low discrimination**							**15000100**
					GRA	SPO		
Clostridium bifermentans		82.09	0.53	LAC 1	99	99		
Fusobacterium varium		9.69	0.32		1	1		
Clostridium cadaveris		4.02	0.29		99	99		
15000120	**Good identification**							**15000120**
					GRA	SPO		
Clostridium cadaveris		94.17	0.63	LAC 1	99	99		
Prevotella bivia		4.43	0.32		1	1		
Clostridium bifermentans		1.24	0.33		99	99		
15000200	**Low discrimination**							**15000200**
					GRA	SPO		
Clostridium bifermentans		82.24	0.53	RHA 1	99	99		
Fusobacterium varium		9.71	0.32		1	1		
Clostridium cadaveris		4.02	0.29		99	99		
15000220	**Very good identification**							**15000220**
					SPO	CAT		
Clostridium cadaveris		98.55	0.63	RHA 1	99	1		
Clostridium bifermentans		1.30	0.33		99	1		
Propionibacterium acnes		0.08	0.14		1	95		
15000400	**Low discrimination**							**15000400**
					COC	GRA	SPO	
Clostridium bifermentans		78.99	0.53	ARA 1	1	99	99	
Fusobacterium varium		9.32	0.32		1	1	1	
Gemella morbillorum		4.35	0.35		99	99	1	
15000420	**Very good identification**							**15000420**
					COC	SPO		
Clostridium cadaveris		98.42	0.63	ARA 1	1	99		
Clostridium bifermentans		1.30	0.33		1	99		
Gemella morbillorum		0.12	0.19		99	1		
15001000	**Low discrimination**							**15001000**
					COC	GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		51.54	0.51	GLU 12	1	1	1	
Clostridium bifermentans		29.14	0.53	MNS 1	1	99	99	
Gemella morbillorum		7.78	0.47		99	99	1	
15001001	**Acceptable identification**							**15001001**
					COC	GRA	SPO	
Clostridium sordellii		94.56	0.47	MNS 4	1	99	99	
Fusobacterium necrophorum subsp. necrophorum		2.12	0.17	MLT 90	1	1	1	
Anaerococcus prevotii		1.33	0.15		99	99	1	
15001002	**Low discrimination**							**15001002**
					GRA	SPO		
Atopobium minutum		77.94	0.45	IND 1	99	1		
Fusobacterium necrophorum subsp. necrophorum		12.37	0.17		1	1		
Clostridium bifermentans		7.00	0.19		99	99		
15001020	**Low discrimination**							**15001020**
					SPO	CAT		
Clostridium cadaveris		89.33	0.63	GAL 1	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		7.54	0.48		1	95		
Clostridium bifermentans		1.18	0.33		99	1		
15001060	**Low discrimination**							**15001060**
					SPO	CAT		
Propionibacterium acnes		63.26	0.36	NIT 92	1	95		
Clostridium cadaveris		34.47	0.29	FRU 99	99	1		
Clostridium bifermentans		1.87	0.09		99	1		
15001520	**Very good identification**							**15001520**
Odoribacter splanchnicus		100.0	0.61	ESL 99				
15001560	**Good identification**							**15001560**
Odoribacter splanchnicus		100.0	0.45	ESL 99				
15002000	**Low discrimination**							**15002000**
					GRA	SPO		
Clostridium bifermentans		77.72	0.53	MAN 1	99	99		
Fusobacterium varium		9.17	0.32		1	1		
Eubacterium limosum		4.82	0.29		99	1		
15002020	**Good identification**							**15002020**
					SPO	CAT		
Clostridium cadaveris		94.20	0.63	MAN 1	99	1		
Propionibacterium acnes		4.48	0.43		1	95		
Clostridium bifermentans		1.24	0.33		99	1		
15003020	**Low discrimination**							**15003020**
					SPO	CAT		
Propionibacterium acnes		82.11	0.43	NIT 92	1	95		
Clostridium cadaveris		17.46	0.29	FRU 99	99	1		
Clostridium bifermentans		0.23	-.01		99	1		
15004000	**Low discrimination**							**15004000**
					COC	GRA	SPO	
Clostridium bifermentans		61.17	0.53	XYL 1	1	99	99	
Fusobacterium varium		29.78	0.42		1	1	1	
Gemella morbillorum		3.37	0.35		99	99	1	
15004020	**Very good identification**							**15004020**
					COC	SPO		
Clostridium cadaveris		98.36	0.63	XYL 1	1	99		
Clostridium bifermentans		1.30	0.33		1	99		
Gemella morbillorum		0.12	0.19		99	1		
15005722	**Good identification**							**15005722**
Bacteroides eggerthii		100.0	0.39	ESL 90				
15010000	**Very good identification**							**15010000**
					GRA	SPO		
Fusobacterium varium		98.66	1.00		1	1		
Clostridium bifermentans		0.44	0.66		99	99		
Clostridium cadaveris		0.41	0.63		99	99		
15010001	**Low discrimination**							**15010001**
					GRA	SPO		
Fusobacterium varium		71.36	0.66	URE 1	1	1		
Clostridium sordellii		27.60	0.60		99	99		
Clostridium bifermentans		0.32	0.31		99	99		
15010002	**Very good identification**							**15010002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Fusobacterium varium		96.01	0.66	BGL 1	1	1		
Atopobium minutum		2.56	0.53		99	1		
Clostridium bifermentans		0.43	0.31		99	99		
15010010	**Very good identification**							**15010010**
					GRA	SPO		
Fusobacterium varium		96.44	0.66	MLZ 1	1	1		
Clostridium difficile		2.22	0.41		99	99		
Clostridium bifermentans		0.43	0.31		99	99		
15010020	**Good identification**							**15010020**
					GRA	SPO	CAT	
Clostridium cadaveris		90.08	0.97		99	99	1	
Propionibacterium acnes		7.61	0.82		99	1	95	
Fusobacterium varium		2.20	0.66		1	1	1	
15010021	**Low discrimination**							**15010021**
					GRA	SPO	CAT	
Clostridium cadaveris		89.33	0.63	URE 1	99	99	1	
Propionibacterium acnes		7.55	0.48		99	1	95	
Fusobacterium varium		2.18	0.32		1	1	1	
15010022	**Low discrimination**							**15010022**
					GRA	SPO	CAT	
Clostridium cadaveris		88.35	0.63	BGL 1	99	99	1	
Propionibacterium acnes		7.46	0.48		99	1	95	
Fusobacterium varium		2.15	0.32		1	1	1	
15010030	**Good identification**							**15010030**
					GRA	SPO	CAT	
Clostridium cadaveris		90.03	0.63	MLZ 1	99	99	1	
Propionibacterium acnes		7.60	0.48		99	1	95	
Fusobacterium varium		2.19	0.32		1	1	1	
15010040	**Very good identification**							**15010040**
					GRA	SPO	CAT	
Fusobacterium varium		96.21	0.66	SOR 1	1	1	1	
Clostridium bifermentans		1.79	0.42		99	99	1	
Propionibacterium acnes		0.74	0.36		99	1	95	
15010041	**Low discrimination**							**15010041**
					GRA	SPO		
Clostridium sordellii		60.71	0.36	MNS 4	99	99		
Fusobacterium varium		38.05	0.32	MLT 90	1	1		
Clostridium bifermentans		0.71	0.08	SOR 4	99	99		
15010060	**Low discrimination**							**15010060**
					GRA	SPO	CAT	
Propionibacterium acnes		64.07	0.71	NIT 92	99	1	95	
Clostridium cadaveris		34.91	0.63		99	99	1	
Fusobacterium varium		0.85	0.32		1	1	1	
15010061	**Low discrimination**							**15010061**
					SPO	CAT		
Propionibacterium acnes		63.32	0.36	NIT 92	1	95		
Clostridium cadaveris		34.50	0.29	URE 1	99	1		
Clostridium sordellii		1.34	0.02		99	1		
15010062	**Low discrimination**							**15010062**
					GRA	SPO	CAT	
Propionibacterium acnes		63.44	0.36	NIT 92	99	1	95	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium cadaveris		34.56	0.29	BGL 1	99	99	1	
Fusobacterium varium		0.84	-.03		1	1	1	
15010070	**Low discrimination**							**15010070**
					GRA	SPO	CAT	
Propionibacterium acnes		63.87	0.36	NIT 92	99	1	95	
Clostridium cadaveris		34.80	0.29	MLZ 1	99	99	1	
Fusobacterium varium		0.85	-.03		1	1	1	
15010100	**Very good identification**							**15010100**
					GRA	SPO		
Fusobacterium varium		98.51	0.66	LAC 1	1	1		
Clostridium bifermentans		0.44	0.31		99	99		
Clostridium cadaveris		0.41	0.29		99	99		
15010120	**Low discrimination**							**15010120**
					GRA	SPO	CAT	
Clostridium cadaveris		89.65	0.63	LAC 1	99	99	1	
Propionibacterium acnes		7.57	0.48		99	1	95	
Fusobacterium varium		2.18	0.32		1	1	1	
15010160	**Low discrimination**							**15010160**
					GRA	SPO	CAT	
Propionibacterium acnes		64.18	0.36	NIT 92	99	1	95	
Clostridium cadaveris		34.97	0.29	LAC 1	99	99	1	
Fusobacterium varium		0.85	-.03		1	1	1	
15010200	**Very good identification**							**15010200**
					GRA	SPO		
Fusobacterium varium		98.66	0.66	RHA 1	1	1		
Clostridium bifermentans		0.44	0.31		99	99		
Clostridium cadaveris		0.41	0.29		99	99		
15010220	**Good identification**							**15010220**
					GRA	SPO	CAT	
Clostridium cadaveris		90.09	0.63	RHA 1	99	99	1	
Propionibacterium acnes		7.61	0.48		99	1	95	
Fusobacterium varium		2.20	0.32		1	1	1	
15010260	**Low discrimination**							**15010260**
					GRA	SPO	CAT	
Propionibacterium acnes		64.18	0.36	NIT 92	99	1	95	
Clostridium cadaveris		34.97	0.29	RHA 1	99	99	1	
Fusobacterium varium		0.85	-.03		1	1	1	
15010400	**Very good identification**							**15010400**
					COC	GRA	SPO	
Fusobacterium varium		98.16	0.66	ARA 1	1	1	1	
Gemella morbillorum		0.57	0.37		99	99	1	
Clostridium bifermentans		0.44	0.31		1	99	99	
15010420	**Low discrimination**							**15010420**
					GRA	SPO	CAT	
Clostridium cadaveris		89.95	0.63	ARA 1	99	99	1	
Propionibacterium acnes		7.60	0.48		99	1	95	
Fusobacterium varium		2.19	0.32		1	1	1	
15010460	**Low discrimination**							**15010460**
					GRA	SPO	CAT	
Propionibacterium acnes		64.18	0.36	NIT 92	99	1	95	
Clostridium cadaveris		34.97	0.29	ARA 1	99	99	1	
Fusobacterium varium		0.85	-.03		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
15011000	**Low discrimination**							**15011000**
					GRA	CAT		
Fusobacterium varium		76.17	0.66	GAL 1	1	1		
Prevotella intermedia		16.81	0.63		1	1		
Propionibacterium acnes		2.67	0.48		99	95		
15011001	**Low discrimination**							**15011001**
					GRA	SPO		
Clostridium sordellii		83.85	0.47	MNS 4	99	99		
Fusobacterium varium		12.41	0.32	MLT 90	1	1		
Prevotella intermedia		2.74	0.29		1	1		
15011020	**Low discrimination**							**15011020**
					GRA	SPO	CAT	
Propionibacterium acnes		88.84	0.82	NIT 92	99	1	95	
Clostridium cadaveris		10.63	0.63		99	99	1	
Fusobacterium varium		0.26	0.32		1	1	1	
15011021	**Low discrimination**							**15011021**
					SPO	CAT		
Propionibacterium acnes		87.45	0.48	NIT 92	1	95		
Clostridium cadaveris		10.46	0.29	URE 1	99	1		
Clostridium sordellii		1.72	0.13		99	1		
15011022	**Low discrimination**							**15011022**
					SPO	CAT		
Propionibacterium acnes		82.32	0.48	NIT 92	1	95		
Clostridium cadaveris		9.85	0.29	BGL 1	99	1		
Atopobium minutum		6.97	0.37		1	1		
15011030	**Low discrimination**							**15011030**
					GRA	SPO	CAT	
Propionibacterium acnes		88.81	0.48	NIT 92	99	1	95	
Clostridium cadaveris		10.62	0.29	MLZ 1	99	99	1	
Fusobacterium varium		0.26	-.03		1	1	1	
15011060	**Very good identification**							**15011060**
					SPO	CAT		
Propionibacterium acnes		99.43	0.71	NIT 92	1	95		
Clostridium cadaveris		0.55	0.29		99	1		
Atopobium minutum		0.01	0.05		1	1		
15011061	**Good identification**							**15011061**
Propionibacterium acnes		100.0	0.36	NIT 92				
15011062	**Good identification**							**15011062**
Propionibacterium acnes		100.0	0.36	NIT 92				
15011070	**Good identification**							**15011070**
Propionibacterium acnes		100.0	0.36	NIT 92				
15011120	**Low discrimination**							**15011120**
					GRA	SPO	CAT	
Propionibacterium acnes		66.43	0.48	NIT 92	99	1	95	
Odoribacter splanchnicus		22.86	0.32	LAC 1	1	1	1	
Clostridium cadaveris		7.95	0.29		99	99	1	
15011160	**Low discrimination**							**15011160**
					GRA	CAT		
Propionibacterium acnes		85.17	0.36	NIT 92	99	95		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Odoribacter splanchnicus		14.83	0.15	LAC 1	1	1		
15011220	**Low discrimination**							**15011220**
					GRA	SPO	CAT	
Propionibacterium acnes		88.98	0.48	NIT 92	99	1	95	
Clostridium cadaveris		10.64	0.29	RHA 1	99	99	1	
Fusobacterium varium		0.26	-.03		1	1	1	
15011260	**Good identification**							**15011260**
Propionibacterium acnes		100.0	0.36	NIT 92				
15011420	**Low discrimination**							**15011420**
					GRA	SPO	CAT	
Propionibacterium acnes		67.51	0.48	NIT 92	99	1	95	
Odoribacter splanchnicus		23.23	0.32	ARA 1	1	1	1	
Clostridium cadaveris		8.07	0.29		99	99	1	
15011460	**Low discrimination**							**15011460**
					GRA	CAT		
Propionibacterium acnes		85.17	0.36	NIT 92	99	95		
Odoribacter splanchnicus		14.83	0.15	ARA 1	1	1		
15011520	**Very good identification**							**15011520**
					GRA	CAT		
Odoribacter splanchnicus		99.95	0.66	ESL 99	1	1		
Propionibacterium acnes		0.03	0.14		99	95		
Lactobacillus fermentum		0.02	0.24		99	1		
15011560	**Good identification**							**15011560**
					GRA	CAT		
Odoribacter splanchnicus		99.94	0.50	ESL 99	1	1		
Propionibacterium acnes		0.06	0.02	SOR 10	99	95		
15012000	**Low discrimination**							**15012000**
					GRA	SPO		
Fusobacterium varium		62.17	0.66	MAN 1	1	1		
Eubacterium limosum		32.69	0.63		99	1		
Clostridium difficile		2.64	0.46		99	99		
15012010	**Good identification**							**15012010**
					GRA	SPO		
Clostridium difficile		92.86	0.58	IND 1	99	99		
Fusobacterium varium		4.52	0.32		1	1		
Eubacterium limosum		2.38	0.29		99	1		
15012020	**Low discrimination**							**15012020**
					GRA	SPO	CAT	
Propionibacterium acnes		81.85	0.78	NIT 92	99	1	95	
Clostridium cadaveris		17.40	0.63		99	99	1	
Fusobacterium varium		0.42	0.32		1	1	1	
15012021	**Low discrimination**							**15012021**
					GRA	SPO	CAT	
Propionibacterium acnes		81.93	0.43	NIT 92	99	1	95	
Clostridium cadaveris		17.42	0.29	URE 1	99	99	1	
Fusobacterium varium		0.42	-.03		1	1	1	
15012022	**Low discrimination**							**15012022**
					SPO	CAT		
Propionibacterium acnes		80.90	0.43	NIT 92	1	95		
Clostridium cadaveris		17.20	0.29	BGL 1	99	1		
Pseudoramibacter alactolyticus		0.45	-.02		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
15012030	**Low discrimination**							**15012030**
					SPO	CAT		
Propionibacterium acnes		75.19	0.43	NIT 92	1	95		
Clostridium cadaveris		15.99	0.29	MLZ 1	99	1		
Clostridium difficile		8.00	0.24		99	1		
15012050	**Good identification**							**15012050**
					GRA	SPO		
Clostridium difficile		98.59	0.36	IND 1	99	99		
Fusobacterium varium		0.92	-.03	SOR 5	1	1		
Eubacterium limosum		0.49	-.06		99	1		
15012060	**Very good identification**							**15012060**
					SPO	CAT		
Propionibacterium acnes		98.94	0.66	NIT 92	1	95		
Clostridium cadaveris		0.97	0.29		99	1		
Pseudoramibacter alactolyticus		0.06	0.04		1	1		
15012120	**Low discrimination**							**15012120**
					GRA	SPO	CAT	
Propionibacterium acnes		81.93	0.43	NIT 92	99	1	95	
Clostridium cadaveris		17.42	0.29	LAC 1	99	99	1	
Fusobacterium varium		0.42	-.03		1	1	1	
15012220	**Low discrimination**							**15012220**
					GRA	SPO	CAT	
Propionibacterium acnes		81.93	0.43	NIT 92	99	1	95	
Clostridium cadaveris		17.42	0.29	RHA 1	99	99	1	
Fusobacterium varium		0.42	-.03		1	1	1	
15012420	**Low discrimination**							**15012420**
					GRA	SPO	CAT	
Propionibacterium acnes		81.93	0.43	NIT 92	99	1	95	
Clostridium cadaveris		17.42	0.29	ARA 1	99	99	1	
Fusobacterium varium		0.42	-.03		1	1	1	
15013000	**Low discrimination**							**15013000**
					GRA	CAT		
Propionibacterium acnes		49.25	0.43	NIT 92	99	95		
Fusobacterium varium		25.26	0.32	NAG 99	1	1		
Eubacterium limosum		13.28	0.29		99	1		
15013020	**Very good identification**							**15013020**
					SPO	CAT		
Propionibacterium acnes		99.75	0.78	NIT 92	1	95		
Clostridium cadaveris		0.21	0.29		99	1		
Propionibacterium granulosum		0.02	0.24		1	90		
15013021	**Good identification**							**15013021**
Propionibacterium acnes		100.0	0.43	NIT 92				
15013022	**Good identification**							**15013022**
Propionibacterium acnes		99.46	0.43	NIT 92				
Propionibacterium granulosum		0.54	0.14	BGL 1				
15013030	**Good identification**							**15013030**
Propionibacterium acnes		99.71	0.43	NIT 92				
Propionibacterium granulosum		0.29	0.09	MLZ 1				
15013060	**Very good identification**							**15013060**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		100.0	0.66	NIT 92				
15013120	**Good identification**							**15013120**
					GRA	CAT		
Propionibacterium acnes		99.39	0.43	NIT 92	99	95		
Odoribacter splanchnicus		0.61	-.03	LAC 1	1	1		
15013220	**Good identification**							**15013220**
Propionibacterium acnes		100.0	0.43	NIT 92				
15013420	**Good identification**							**15013420**
					GRA	CAT		
Propionibacterium acnes		99.39	0.43	NIT 92	99	95		
Odoribacter splanchnicus		0.61	-.03	ARA 1	1	1		
15014000	**Very good identification**							**15014000**
					COC	GRA	SPO	
Fusobacterium varium		99.54	0.76	XYL 4	1	1	1	
Gemella morbillorum		0.14	0.37		99	99	1	
Clostridium bifermentans		0.11	0.31		1	99	99	
15014001	**Acceptable identification**							**15014001**
					COC	GRA	SPO	
Fusobacterium varium		91.22	0.42	XYL 4	1	1	1	
Clostridium sordellii		8.55	0.26	URE 1	1	99	99	
Gemella morbillorum		0.13	0.03		99	99	1	
15014002	**Good identification**							**15014002**
					COC	GRA		
Fusobacterium varium		99.11	0.42	XYL 4	1	1		
Atopobium minutum		0.64	0.19	BGL 1	1	99		
Gemella morbillorum		0.14	0.03		99	99		
15014010	**Good identification**							**15014010**
					COC	GRA	SPO	
Fusobacterium varium		96.94	0.42	XYL 4	1	1	1	
Clostridium difficile		2.82	0.19	MLZ 1	1	99	99	
Gemella morbillorum		0.14	0.03		99	99	1	
15014020	**Low discrimination**							**15014020**
					GRA	SPO	CAT	
Clostridium cadaveris		84.21	0.63	XYL 1	99	99	1	
Fusobacterium varium		8.46	0.42		1	1	1	
Propionibacterium acnes		7.11	0.48		99	1	95	
15014040	**Low discrimination**							**15014040**
					GRA	SPO		
Fusobacterium varium		89.33	0.42	XYL 4	1	1		
Clostridium glycolicum		9.98	0.31	SOR 1	99	99		
Clostridium bifermentans		0.40	0.08		99	99		
15014060	**Low discrimination**							**15014060**
					GRA	SPO	CAT	
Propionibacterium acnes		62.28	0.36	NIT 92	99	1	95	
Clostridium cadaveris		33.93	0.29	XYL 1	99	99	1	
Fusobacterium varium		3.41	0.08		1	1	1	
15014100	**Good identification**							**15014100**
					GRA			
Fusobacterium varium		99.20	0.42	XYL 4	1			
Fusobacterium mortiferum		0.28	0.18	LAC 1	1			

Code / Taxon		ID score	T index	tests against		Differentiation tests			Code
Actinomyces meyeri		0.28	0.09			99			
15014200	**Good identification**								**15014200**
						COC	GRA	SPO	
Fusobacterium varium		99.75	0.42	XYL	4	1	1	1	
Gemella morbillorum		0.14	0.03	RHA	1	99	99	1	
Clostridium bifermentans		0.11	-.03			1	99	99	
15014400	**Good identification**								**15014400**
						COC	GRA	SPO	
Fusobacterium varium		98.38	0.42	XYL	4	1	1	1	
Gemella morbillorum		1.51	0.20	ARA	1	99	99	1	
Clostridium bifermentans		0.11	-.03			1	99	99	
15015000	**Low discrimination**								**15015000**
						COC	GRA		
Fusobacterium varium		87.12	0.42	GAL	1	1	1		
Gemella morbillorum		6.49	0.32	XYL	4	99	99		
Prevotella intermedia		4.66	0.29			1	1		
15015020	**Low discrimination**								**15015020**
						GRA	SPO	CAT	
Propionibacterium acnes		87.35	0.48	NIT	92	99	1	95	
Clostridium cadaveris		10.45	0.29	XYL	1	99	99	1	
Fusobacterium varium		1.05	0.08			1	1	1	
15015060	**Good identification**								**15015060**
Propionibacterium acnes		100.0	0.36	NIT	92				
15016000	**Low discrimination**								**15016000**
						GRA	SPO		
Fusobacterium varium		84.05	0.42	MAN	1	1	1		
Eubacterium limosum		10.72	0.29	XYL	4	99	1		
Clostridium difficile		4.51	0.24			99	99		
15016010	**Good identification**								**15016010**
						GRA	SPO		
Clostridium difficile		95.84	0.36	IND	1	99	99		
Fusobacterium varium		3.69	0.08	XYL	5	1	1		
Eubacterium limosum		0.47	-.06			99	1		
15016020	**Low discrimination**								**15016020**
						GRA	SPO	CAT	
Propionibacterium acnes		80.86	0.43	NIT	92	99	1	95	
Clostridium cadaveris		17.19	0.29	XYL	1	99	99	1	
Fusobacterium varium		1.73	0.08			1	1	1	
15017020	**Good identification**								**15017020**
Propionibacterium acnes		100.0	0.43	NIT	92				
15020000	**Low discrimination**								**15020000**
						GRA	SPO		
Clostridium bifermentans		67.84	0.53	TRE	1	99	99		
Clostridium cadaveris		17.30	0.41			99	99		
Fusobacterium varium		8.01	0.32			1	1		
15020020	**Very good identification**								**15020020**
						SPO	CAT		
Clostridium cadaveris		99.72	0.75	TRE	5	99	1		
Clostridium bifermentans		0.25	0.33			99	1		
Propionibacterium acnes		0.02	0.14			1	95		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
15020021	**Good identification**							**15020021**
Clostridium cadaveris		99.75	0.41	TRE 5				
Clostridium bifermentans		0.25	-.01	URE 1				
15020022	**Good identification**							**15020022**
					SPO			
Clostridium cadaveris		99.31	0.41	TRE 5	99			
Atopobium minutum		0.44	0.11	BGL 1	1			
Clostridium bifermentans		0.25	-.01		99			
15020030	**Good identification**							**15020030**
Clostridium cadaveris		99.75	0.41	TRE 5				
Clostridium bifermentans		0.25	-.01	MLZ 1				
15020060	**Good identification**							**15020060**
Clostridium cadaveris		98.23	0.41	TRE 5				
Clostridium bifermentans		1.03	0.09	SOR 1				
Clostridium botulinum biovar B		0.39	0.00					
15020120	**Good identification**							**15020120**
					GRA	SPO		
Clostridium cadaveris		98.86	0.41	TRE 5	99	99		
Prevotella bivia		0.89	-.03	LAC 1	1	1		
Clostridium bifermentans		0.25	-.01		99	99		
15020220	**Good identification**							**15020220**
Clostridium cadaveris		99.75	0.41	TRE 5				
Clostridium bifermentans		0.25	-.01	RHA 1				
15020420	**Good identification**							**15020420**
Clostridium cadaveris		99.75	0.41	TRE 5				
Clostridium bifermentans		0.25	-.01	ARA 1				
15021020	**Good identification**							**15021020**
					SPO	CAT		
Clostridium cadaveris		97.95	0.41	TRE 5	99	1		
Propionibacterium acnes		1.59	0.14	GAL 1	1	95		
Clostridium bifermentans		0.25	-.01		99	1		
15022020	**Good identification**							**15022020**
					SPO	CAT		
Clostridium cadaveris		98.85	0.41	TRE 5	99	1		
Propionibacterium acnes		0.90	0.09	MAN 1	1	95		
Clostridium bifermentans		0.25	-.01		99	1		
15024020	**Good identification**							**15024020**
Clostridium cadaveris		99.75	0.41	TRE 5				
Clostridium bifermentans		0.25	-.01	XYL 1				
15030000	**Very good identification**							**15030000**
					GRA	SPO	CAT	
Fusobacterium varium		96.27	0.66	TRE 1	1	1	1	
Clostridium cadaveris		2.10	0.41		99	99	1	
Propionibacterium granulosum		0.65	0.45		99	1	90	
15030020	**Very good identification**							**15030020**
					GRA	SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium cadaveris		97.86	0.75	TRE 5	99	99	1	
Propionibacterium acnes		1.59	0.48		99	1	95	
Fusobacterium varium		0.46	0.32		1	1	1	
15030021	**Good identification**							**15030021**
					GRA	SPO	CAT	
Clostridium cadaveris		97.95	0.41	TRE 5	99	99	1	
Propionibacterium acnes		1.59	0.14	URE 1	99	1	95	
Fusobacterium varium		0.46	-.03		1	1	1	
15030022	**Good identification**							**15030022**
					SPO	CAT		
Clostridium cadaveris		95.75	0.41	TRE 5	99	1		
Propionibacterium granulosum		1.86	0.24	BGL 1	1	90		
Propionibacterium acnes		1.55	0.14		1	95		
15030030	**Good identification**							**15030030**
					SPO	CAT		
Clostridium cadaveris		96.95	0.41	TRE 5	99	1		
Propionibacterium acnes		1.57	0.14	MLZ 1	1	95		
Propionibacterium granulosum		1.03	0.20		1	90		
15030060	**Low discrimination**							**15030060**
					SPO	CAT		
Clostridium cadaveris		72.84	0.41	TRE 5	99	1		
Propionibacterium acnes		25.66	0.36	SOR 1	1	95		
Clostridium botulinum biovar B		1.16	0.10		99	1		
15030120	**Good identification**							**15030120**
					GRA	SPO	CAT	
Clostridium cadaveris		97.95	0.41	TRE 5	99	99	1	
Propionibacterium acnes		1.59	0.14	LAC 1	99	1	95	
Fusobacterium varium		0.46	-.03		1	1	1	
15030220	**Good identification**							**15030220**
					GRA	SPO	CAT	
Clostridium cadaveris		97.72	0.41	TRE 5	99	99	1	
Propionibacterium acnes		1.58	0.14	RHA 1	99	1	95	
Fusobacterium varium		0.46	-.03		1	1	1	
15030420	**Good identification**							**15030420**
					GRA	SPO	CAT	
Clostridium cadaveris		97.95	0.41	TRE 5	99	99	1	
Propionibacterium acnes		1.59	0.14	ARA 1	99	1	95	
Fusobacterium varium		0.46	-.03		1	1	1	
15031020	**Low discrimination**							**15031020**
					SPO	CAT		
Propionibacterium acnes		59.67	0.48	NIT 92	1	95		
Clostridium cadaveris		37.19	0.41	TRE 1	99	1		
Propionibacterium granulosum		2.89	0.35		1	90		
15031060	**Good identification**							**15031060**
					SPO	CAT		
Propionibacterium acnes		97.21	0.36	NIT 92	1	95		
Clostridium cadaveris		2.79	0.07	TRE 1	99	1		
15032010	**Acceptable identification**							**15032010**
					GRA	SPO	CAT	
Clostridium difficile		94.82	0.38	IND 1	99	99	1	
Propionibacterium granulosum		4.24	0.28	TRE 7	99	1	90	
Fusobacterium varium		0.62	-.03		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
15032020	**Low discrimination**							**15032020**
					SPO	CAT		
Clostridium cadaveris		50.80	0.41	TRE 5	99	1		
Propionibacterium acnes		45.86	0.43	MAN 1	1	95		
Propionibacterium granulosum		2.98	0.32		1	90		
15033000	**Acceptable identification**							**15033000**
					GRA	CAT		
Propionibacterium granulosum		93.52	0.43	IND 1	99	90		
Propionibacterium acnes		3.64	0.09	SUC 93	99	95		
Fusobacterium varium		1.86	-.03		1	1		
15033020	**Acceptable identification**							**15033020**
					SPO	CAT		
Propionibacterium acnes		92.92	0.43	NIT 92	1	95		
Propionibacterium granulosum		6.04	0.32	TRE 1	1	90		
Clostridium cadaveris		1.04	0.07		99	1		
15034000	**Good identification**							**15034000**
					GRA	SPO	CAT	
Fusobacterium varium		99.07	0.42	TRE 1	1	1	1	
Clostridium cadaveris		0.52	0.07	XYL 4	99	99	1	
Propionibacterium granulosum		0.16	0.11		99	1	90	
15034020	**Good identification**							**15034020**
					GRA	SPO	CAT	
Clostridium cadaveris		96.57	0.41	TRE 5	99	99	1	
Fusobacterium varium		1.86	0.08	XYL 1	1	1	1	
Propionibacterium acnes		1.57	0.14		99	1	95	
15040000	**Low discrimination**							**15040000**
					GRA	SPO		
Clostridium bifermentans		82.24	0.53	CEL 1	99	99		
Fusobacterium varium		9.71	0.32		1	1		
Clostridium cadaveris		4.02	0.29		99	99		
15040020	**Very good identification**							**15040020**
					SPO	CAT		
Clostridium cadaveris		98.55	0.63	CEL 1	99	1		
Clostridium bifermentans		1.30	0.33		99	1		
Propionibacterium acnes		0.08	0.14		1	95		
15045722	**Good identification**							**15045722**
Bacteroides eggerthii		100.0	0.50	ESL 90				
15050000	**Very good identification**							**15050000**
					GRA	SPO		
Fusobacterium varium		98.63	0.66	CEL 1	1	1		
Clostridium bifermentans		0.44	0.31		99	99		
Clostridium cadaveris		0.41	0.29		99	99		
15050020	**Good identification**							**15050020**
					GRA	SPO	CAT	
Clostridium cadaveris		90.09	0.63	CEL 1	99	99	1	
Propionibacterium acnes		7.61	0.48		99	1	95	
Fusobacterium varium		2.20	0.32		1	1	1	
15050060	**Low discrimination**							**15050060**
					GRA	SPO	CAT	
Propionibacterium acnes		64.18	0.36	NIT 92	99	1	95	
Clostridium cadaveris		34.97	0.29	CEL 1	99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium varium		0.85	-.03		1	1	1	
15051020	**Low discrimination**							**15051020**
					GRA	SPO	CAT	
Propionibacterium acnes		88.98	0.48	NIT 92	99	1	95	
Clostridium cadaveris		10.64	0.29	CEL 1	99	99	1	
Fusobacterium varium		0.26	-.03		1	1	1	
15051060	**Good identification**							**15051060**
Propionibacterium acnes		100.0	0.36	NIT 92				
15052010	**Good identification**							**15052010**
					GRA	SPO		
Clostridium difficile		98.59	0.36	IND 1	99	99		
Fusobacterium varium		0.92	-.03	CEL 5	1	1		
Eubacterium limosum		0.49	-.06		99	1		
15052020	**Low discrimination**							**15052020**
					GRA	SPO	CAT	
Propionibacterium acnes		81.93	0.43	NIT 92	99	1	95	
Clostridium cadaveris		17.42	0.29	CEL 1	99	99	1	
Fusobacterium varium		0.42	-.03		1	1	1	
15053020	**Good identification**							**15053020**
Propionibacterium acnes		100.0	0.43	NIT 92				
15054000	**Good identification**							**15054000**
					COC	GRA	SPO	
Fusobacterium varium		99.75	0.42	CEL 1	1	1	1	
Gemella morbillorum		0.14	0.03	XYL 4	99	99	1	
Clostridium bifermentans		0.11	-.03		1	99	99	
15060020	**Good identification**							**15060020**
Clostridium cadaveris		99.75	0.41	TRE 5				
Clostridium bifermentans		0.25	-.01	CEL 1				
15070020	**Good identification**							**15070020**
					GRA	SPO	CAT	
Clostridium cadaveris		97.95	0.41	TRE 5	99	99	1	
Propionibacterium acnes		1.59	0.14	CEL 1	99	1	95	
Fusobacterium varium		0.46	-.03		1	1	1	
15100000	**Good identification**							**15100000**
					COC	GRA	SPO	
Clostridium bifermentans		92.91	0.94		1	99	99	
Prevotella disiens		4.41	0.66		1	1	1	
Gemella morbillorum		1.13	0.64		99	99	1	
15100001	**Good identification**							**15100001**
					GRA	SPO		
Clostridium sordellii		94.15	0.76	MNS 4	99	99		
Clostridium bifermentans		5.40	0.60		99	99		
Prevotella disiens		0.26	0.32		1	1		
15100002	**Good identification**							**15100002**
					COC	GRA	SPO	
Clostridium bifermentans		92.59	0.60	BGL 1	1	99	99	
Prevotella disiens		4.40	0.32		1	1	1	
Gemella morbillorum		1.13	0.30		99	99	1	
15100003	**Acceptable identification**							**15100003**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Clostridium sordellii		93.34	0.42	MNS 4	1	99		
Clostridium bifermentans		5.35	0.25	BGL 1	1	99		
Anaerococcus prevotii		0.64	0.04		99	1		
15100010	**Good identification**							**15100010**
					COC	GRA	SPO	
Clostridium bifermentans		92.55	0.60	MLZ 1	1	99	99	
Prevotella disiens		4.40	0.32		1	1	1	
Gemella morbillorum		1.13	0.30		99	99	1	
15100011	**Acceptable identification**							**15100011**
Clostridium sordellii		92.13	0.42	MNS 4				
Clostridium bifermentans		5.29	0.25	MLZ 1				
Clostridium novyi biovar A		2.34	0.32					
15100020	**Low discrimination**							**15100020**
					COC	SPO		
Clostridium cadaveris		62.04	0.75	MLT 5	1	99		
Clostridium bifermentans		36.25	0.73		1	99		
Gemella morbillorum		0.77	0.48		99	1		
15100021	**Low discrimination**							**15100021**
Clostridium sordellii		49.87	0.42	MNS 4				
Clostridium cadaveris		30.94	0.41	NAG 1				
Clostridium bifermentans		18.08	0.39					
15100022	**Low discrimination**							**15100022**
					COC	SPO		
Clostridium cadaveris		61.94	0.41	MLT 5	1	99		
Clostridium bifermentans		36.19	0.39	BGL 1	1	99		
Gemella morbillorum		0.77	0.14		99	1		
15100030	**Low discrimination**							**15100030**
					COC	SPO		
Clostridium cadaveris		61.72	0.41	MLT 5	1	99		
Clostridium bifermentans		36.06	0.39	MLZ 1	1	99		
Gemella morbillorum		0.76	0.14		99	1		
15100040	**Very good identification**							**15100040**
					GRA	SPO		
Clostridium bifermentans		96.84	0.70	SOR 4	99	99		
Prevotella disiens		1.12	0.32		1	1		
Clostridium botulinum biovar B		0.69	0.31		99	99		
15100041	**Acceptable identification**							**15100041**
Clostridium sordellii		94.12	0.53	MNS 4				
Clostridium bifermentans		5.40	0.36	SOR 4				
Clostridium novyi biovar A		0.39	0.29					
15100042	**Good identification**							**15100042**
					GRA	SPO		
Clostridium bifermentans		96.97	0.36	SOR 4	99	99		
Clostridium novyi biovar A		1.22	0.16	BGL 1	99	99		
Prevotella disiens		1.12	-.03		1	1		
15100050	**Low discrimination**							**15100050**
Clostridium bifermentans		70.01	0.36	MLZ 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		24.18	0.26	SOR 4				
Clostridium novyi biovar A		5.00	0.29					
15100060	**Low discrimination**							**15100060**
Clostridium bifermentans		67.36	0.50	NAG 6				
Clostridium cadaveris		27.95	0.41	SOR 4				
Clostridium botulinum biovar B		3.68	0.26					
15100100	**Low discrimination**							**15100100**
					GRA	SPO		
Clostridium bifermentans		85.20	0.60	LAC 1	99	99		
Prevotella bivia		8.32	0.32		1	1		
Prevotella disiens		4.05	0.32		1	1		
15100101	**Acceptable identification**							**15100101**
					GRA	SPO		
Clostridium sordellii		93.84	0.42	MNS 4	99	99		
Clostridium bifermentans		5.38	0.25	LAC 1	99	99		
Prevotella bivia		0.53	-.03		1	1		
15100120	**Very good identification**							**15100120**
					GRA	SPO		
Prevotella bivia		98.22	0.66	IND 1	1	1		
Clostridium cadaveris		1.11	0.41		99	99		
Clostridium bifermentans		0.65	0.39		99	99		
15100140	**Good identification**							**15100140**
					GRA	SPO		
Clostridium bifermentans		95.95	0.36	LAC 1	99	99		
Prevotella bivia		2.27	-.03	SOR 4	1	1		
Prevotella disiens		1.11	-.03		1	1		
15100200	**Good identification**							**15100200**
					COC	GRA	SPO	
Clostridium bifermentans		92.93	0.60	RHA 1	1	99	99	
Prevotella disiens		4.42	0.32		1	1	1	
Gemella morbillorum		1.13	0.30		99	99	1	
15100201	**Acceptable identification**							**15100201**
					GRA	SPO		
Clostridium sordellii		94.33	0.42	MNS 4	99	99		
Clostridium bifermentans		5.41	0.25	RHA 1	99	99		
Prevotella disiens		0.26	-.03		1	1		
15100220	**Low discrimination**							**15100220**
					COC	SPO		
Clostridium cadaveris		62.11	0.41	MLT 5	1	99		
Clostridium bifermentans		36.29	0.39	RHA 1	1	99		
Gemella morbillorum		0.77	0.14		99	1		
15100240	**Good identification**							**15100240**
					GRA	SPO		
Clostridium bifermentans		98.17	0.36	RHA 1	99	99		
Prevotella disiens		1.13	-.03	SOR 4	1	1		
Clostridium botulinum biovar B		0.70	-.03		99	99		
15100400	**Low discrimination**							**15100400**
					COC	GRA	SPO	
Clostridium bifermentans		83.50	0.60	ARA 1	1	99	99	
Gemella morbillorum		11.17	0.48		99	99	1	
Prevotella disiens		3.97	0.32		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
15100401	**Acceptable identification**							**15100401**
					COC	SPO		
Clostridium sordellii		93.65	0.42	MNS 4	1	99		
Clostridium bifermentans		5.37	0.25	ARA 1	1	99		
Gemella morbillorum		0.72	0.14		99	1		
15100420	**Low discrimination**							**15100420**
					COC	SPO		
Clostridium cadaveris		57.68	0.41	MLT 5	1	99		
Clostridium bifermentans		33.70	0.39	ARA 1	1	99		
Gemella morbillorum		7.85	0.32		99	1		
15100440	**Good identification**							**15100440**
					COC	GRA	SPO	
Clostridium bifermentans		95.15	0.36	ARA 1	1	99	99	
Gemella morbillorum		3.09	0.14	SOR 4	99	99	1	
Prevotella disiens		1.10	-.03		1	1	1	
15101000	**Low discrimination**							**15101000**
					COC	GRA	SPO	
Prevotella intermedia		38.86	0.63	FRU 99	1	1	1	
Clostridium bifermentans		35.16	0.60		1	99	99	
Gemella morbillorum		22.79	0.60		99	99	1	
15101001	**Very good identification**							**15101001**
					GRA	SPO		
Clostridium sordellii		99.07	0.64	MNS 4	99	99		
Prevotella intermedia		0.36	0.29		1	1		
Clostridium bifermentans		0.33	0.25		99	99		
15101020	**Low discrimination**							**15101020**
					COC	SPO		
Clostridium cadaveris		39.54	0.41	MLT 5	1	99		
Gemella morbillorum		26.07	0.43	GAL 1	99	1		
Clostridium bifermentans		23.10	0.39		1	99		
15101040	**Low discrimination**							**15101040**
					COC	GRA	SPO	
Clostridium bifermentans		67.44	0.36	GAL 1	1	99	99	
Prevotella intermedia		18.07	0.29	SOR 4	1	1	1	
Gemella morbillorum		10.60	0.26		99	99	1	
15101041	**Good identification**							**15101041**
Clostridium sordellii		99.42	0.40	MNS 4				
Clostridium bifermentans		0.33	0.02	SOR 4				
Clostridium novyi biovar A		0.26	0.12					
15101120	**Good identification**							**15101120**
					GRA	SPO		
Prevotella bivia		99.73	0.53	IND 1	1	1		
Clostridium cadaveris		0.07	0.07		99	99		
Clostridium perfringens		0.05	0.10		99	84		
15101400	**Low discrimination**							**15101400**
					COC	GRA	SPO	
Gemella morbillorum		76.55	0.43	IND 1	99	99	1	
Prevotella intermedia		11.87	0.29	ARA 10	1	1	1	
Clostridium bifermentans		10.74	0.25		1	99	99	
15101722	**Good identification**							**15101722**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides eggerthii		100.0	0.54	ESL 90				
15102000	**Good identification**							**15102000**
					COC	GRA	SPO	
Clostridium bifermentans		92.87	0.60	MAN 1	1	99	99	
Prevotella disiens		4.41	0.32		1	1	1	
Gemella morbillorum		1.13	0.30		99	99	1	
15102001	**Acceptable identification**							**15102001**
					GRA	SPO		
Clostridium sordellii		94.33	0.42	MNS 4	99	99		
Clostridium bifermentans		5.41	0.25	MAN 1	99	99		
Prevotella disiens		0.26	-.03		1	1		
15102020	**Low discrimination**							**15102020**
					COC	SPO		
Clostridium cadaveris		61.76	0.41	MLT 5	1	99		
Clostridium bifermentans		36.08	0.39	MAN 1	1	99		
Gemella morbillorum		0.76	0.14		99	1		
15102040	**Good identification**							**15102040**
					GRA	SPO		
Clostridium bifermentans		97.52	0.36	MAN 1	99	99		
Prevotella disiens		1.12	-.03	SOR 4	1	1		
Clostridium botulinum biovar B		0.69	-.03		99	99		
15104000	**Low discrimination**							**15104000**
					COC	GRA	SPO	
Clostridium bifermentans		83.38	0.60	XYL 1	1	99	99	
Gemella morbillorum		11.15	0.48		99	99	1	
Prevotella disiens		3.96	0.32		1	1	1	
15104001	**Acceptable identification**							**15104001**
					COC	SPO		
Clostridium sordellii		93.65	0.42	MNS 4	1	99		
Clostridium bifermentans		5.37	0.25	XYL 1	1	99		
Gemella morbillorum		0.72	0.14		99	1		
15104020	**Low discrimination**							**15104020**
					COC	SPO		
Clostridium cadaveris		57.68	0.41	MLT 5	1	99		
Clostridium bifermentans		33.70	0.39	XYL 1	1	99		
Gemella morbillorum		7.85	0.32		99	1		
15104040	**Low discrimination**							**15104040**
					COC	SPO		
Clostridium bifermentans		89.42	0.36	XYL 1	1	99		
Clostridium glycolicum		6.02	0.15	SOR 4	1	99		
Gemella morbillorum		2.90	0.14		99	1		
15104722	**Good identification**							**15104722**
Bacteroides eggerthii		100.0	0.39	ESL 90				
15105000	**Low discrimination**							**15105000**
					COC	GRA	SPO	
Gemella morbillorum		76.55	0.43	IND 1	99	99	1	
Prevotella intermedia		11.87	0.29	XYL 10	1	1	1	
Clostridium bifermentans		10.74	0.25		1	99	99	
15105322	**Good identification**							**15105322**
Bacteroides eggerthii		100.0	0.39	ESL 90				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
15105522	**Good identification**						**15105522**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15105622	**Good identification**						**15105622**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15105702	**Good identification**						**15105702**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15105720	**Good identification**						**15105720**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15105722	**Very good identification**						**15105722**
Bacteroides eggerthii		100.0	0.73	ESL 90			
15105723	**Good identification**						**15105723**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15105732	**Good identification**						**15105732**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15105762	**Good identification**						**15105762**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15107722	**Good identification**						**15107722**
Bacteroides eggerthii		100.0	0.39	ESL 90			
15110000	**Low discrimination**						**15110000**
					GRA	SPO	
Prevotella intermedia		86.75	0.97		1	1	
Clostridium bifermentans		4.13	0.72		99	99	
Fusobacterium varium		3.97	0.66		1	1	
15110001	**Good identification**						**15110001**
					GRA	SPO	
Clostridium sordellii		93.20	0.76	MNS 4	99	99	
Prevotella intermedia		5.91	0.63		1	1	
Clostridium bifermentans		0.28	0.38		99	99	
15110002	**Low discrimination**						**15110002**
					GRA	SPO	
Prevotella intermedia		86.51	0.63	BGL 1	1	1	
Clostridium bifermentans		4.12	0.38		99	99	
Fusobacterium varium		3.96	0.32		1	1	
15110003	**Acceptable identification**						**15110003**
					GRA	SPO	
Clostridium sordellii		93.28	0.42	MNS 4	99	99	
Prevotella intermedia		5.92	0.29	BGL 1	1	1	
Clostridium bifermentans		0.28	0.04		99	99	
15110010	**Low discrimination**						**15110010**
					GRA	SPO	
Prevotella intermedia		86.58	0.63	MLZ 1	1	1	
Clostridium bifermentans		4.12	0.38		99	99	
Fusobacterium varium		3.96	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
15110011	**Acceptable identification**						**15110011**
					GRA	SPO	
Clostridium sordellii		93.03	0.42	MNS 4	99	99	
Prevotella intermedia		5.90	0.29	MLZ 1	1	1	
Clostridium bifermentans		0.28	0.04		99	99	
15110020	**Low discrimination**						**15110020**
					GRA	SPO	
Clostridium cadaveris		85.13	0.75	MLT 5	99	99	
Prevotella intermedia		8.70	0.63		1	1	
Clostridium bifermentans		2.62	0.51		99	99	
15110021	**Low discrimination**						**15110021**
					GRA	SPO	
Clostridium sordellii		57.87	0.42	MNS 4	99	99	
Clostridium cadaveris		35.91	0.41	NAG 1	99	99	
Prevotella intermedia		3.67	0.29		1	1	
15110022	**Low discrimination**						**15110022**
					GRA	SPO	
Clostridium cadaveris		84.72	0.41	MLT 5	99	99	
Prevotella intermedia		8.66	0.29	BGL 1	1	1	
Clostridium bifermentans		2.61	0.17		99	99	
15110030	**Low discrimination**						**15110030**
					GRA	SPO	
Clostridium cadaveris		85.04	0.41	MLT 5	99	99	
Prevotella intermedia		8.69	0.29	MLZ 1	1	1	
Clostridium bifermentans		2.62	0.17		99	99	
15110040	**Low discrimination**						**15110040**
					GRA	SPO	
Prevotella intermedia		70.42	0.63	SOR 1	1	1	
Clostridium bifermentans		13.83	0.48		99	99	
Clostridium botulinum biovar B		7.45	0.41		99	99	
15110041	**Good identification**						**15110041**
					GRA	SPO	
Clostridium sordellii		97.86	0.53	MNS 4	99	99	
Prevotella intermedia		1.50	0.29	SOR 4	1	1	
Clostridium bifermentans		0.30	0.14		99	99	
15110050	**Low discrimination**						**15110050**
					GRA	SPO	
Clostridium botulinum biovar B		79.68	0.36	IND 1	99	99	
Prevotella intermedia		15.45	0.29	SUC 90	1	1	
Clostridium bifermentans		3.04	0.14		99	99	
15110060	**Low discrimination**						**15110060**
					SPO	CAT	
Clostridium cadaveris		46.90	0.41	MLT 5	99	1	
Clostridium botulinum biovar B		24.71	0.36	SOR 1	99	1	
Propionibacterium acnes		16.52	0.36		1	95	
15110100	**Low discrimination**						**15110100**
					GRA	SPO	
Prevotella intermedia		85.96	0.63	LAC 1	1	1	
Clostridium bifermentans		4.09	0.38		99	99	
Fusobacterium varium		3.93	0.32		1	1	
15110101	**Acceptable identification**						**15110101**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sordellii		93.28	0.42	MNS 4	99	99		
Prevotella intermedia		5.92	0.29	LAC 1	1	1		
Clostridium bifermentans		0.28	0.04		99	99		
15110120	**Low discrimination**							**15110120**
					GRA	SPO		
Prevotella bivia		89.24	0.50	IND 1	1	1		
Clostridium cadaveris		9.08	0.41	FRU 10	99	99		
Prevotella intermedia		0.93	0.29		1	1		
15110200	**Low discrimination**							**15110200**
					GRA	SPO		
Prevotella intermedia		86.78	0.63	RHA 1	1	1		
Clostridium bifermentans		4.13	0.38		99	99		
Fusobacterium varium		3.97	0.32		1	1		
15110201	**Acceptable identification**							**15110201**
					GRA	SPO		
Clostridium sordellii		93.28	0.42	MNS 4	99	99		
Prevotella intermedia		5.92	0.29	RHA 1	1	1		
Clostridium bifermentans		0.28	0.04		99	99		
15110220	**Low discrimination**							**15110220**
					GRA	SPO		
Clostridium cadaveris		85.23	0.41	MLT 5	99	99		
Prevotella intermedia		8.71	0.29	RHA 1	1	1		
Clostridium bifermentans		2.62	0.17		99	99		
15110400	**Low discrimination**							**15110400**
					COC	GRA	SPO	
Prevotella intermedia		77.71	0.63	ARA 1	1	1	1	
Gemella morbillorum		11.49	0.50		99	99	1	
Clostridium bifermentans		3.70	0.38		1	99	99	
15110401	**Acceptable identification**							**15110401**
					COC	GRA	SPO	
Clostridium sordellii		92.47	0.42	MNS 4	1	99	99	
Prevotella intermedia		5.86	0.29	ARA 1	1	1	1	
Gemella morbillorum		0.87	0.15		99	99	1	
15110420	**Low discrimination**							**15110420**
					COC	GRA	SPO	
Clostridium cadaveris		75.50	0.41	MLT 5	1	99	99	
Gemella morbillorum		12.55	0.33	ARA 1	99	99	1	
Prevotella intermedia		7.72	0.29		1	1	1	
15111000	**Excellent identification**							**15111000**
					COC	GRA	SPO	
Prevotella intermedia		99.10	0.97		1	1	1	
Gemella morbillorum		0.72	0.61		99	99	1	
Clostridium bifermentans		0.05	0.38		1	99	99	
15111001	**Low discrimination**							**15111001**
					COC	GRA	SPO	
Clostridium sordellii		73.40	0.64	MNS 4	1	99	99	
Prevotella intermedia		26.37	0.63		1	1	1	
Gemella morbillorum		0.19	0.27		99	99	1	
15111002	**Very good identification**							**15111002**
					COC	GRA		
Prevotella intermedia		98.47	0.63	BGL 1	1	1		
Gemella morbillorum		0.71	0.27		99	99		

Code / Taxon		ID score	T index	tests against		Differentiation tests			Code
Lactobacillus fermentum		0.50	0.36		1	99			
15111010	**Very good identification**								**15111010**
					COC	GRA	CAT		
Prevotella intermedia		99.02	0.63	MLZ 1	1	1	1		
Gemella morbillorum		0.72	0.27		99	99	1		
Propionibacterium granulosum		0.06	0.16		1	99	90		
15111020	**Low discrimination**								**15111020**
					GRA	SPO	CAT		
Prevotella intermedia		73.71	0.63	NAG 1	1	1	1		
Propionibacterium acnes		11.69	0.48		99	1	95		
Clostridium cadaveris		7.29	0.41		99	99	1		
15111040	**Very good identification**								**15111040**
					COC	GRA	SPO		
Prevotella intermedia		98.75	0.63	SOR 1	1	1	1		
Gemella morbillorum		0.72	0.27		99	99	1		
Clostridium bifermentans		0.20	0.14		1	99	99		
15111041	**Acceptable identification**								**15111041**
					GRA	SPO			
Clostridium sordellii		91.99	0.40	MNS 4	99	99			
Prevotella intermedia		8.01	0.29	SOR 4	1	1			
15111060	**Low discrimination**								**15111060**
					GRA	CAT			
Propionibacterium acnes		70.76	0.36	NIT 92	99	95			
Prevotella intermedia		20.53	0.29	MLT 1	1	1			
Clostridium perfringens		3.98	0.17		99	1			
15111100	**Very good identification**								**15111100**
					COC	GRA			
Prevotella intermedia		97.61	0.63	LAC 1	1	1			
Lactobacillus fermentum		0.81	0.40		1	99			
Gemella morbillorum		0.71	0.27		99	99			
15111120	**Low discrimination**								**15111120**
					GRA	SPO			
Prevotella bivia		55.82	0.37	IND 1	1	1			
Clostridium perfringens		24.54	0.44	FRU 10	99	84			
Clostridium septicum		12.85	0.33		99	99			
15111122	**Low discrimination**								**15111122**
					GRA				
Clostridium perfringens		84.42	0.38	IND 1	99				
Lactobacillus fermentum		6.60	0.32	SUC 91	99				
Prevotella bivia		4.12	0.03		1				
15111200	**Very good identification**								**15111200**
					COC	GRA	SPO		
Prevotella intermedia		99.12	0.63	RHA 1	1	1	1		
Gemella morbillorum		0.72	0.27		99	99	1		
Clostridium bifermentans		0.05	0.04		1	99	99		
15111400	**Good identification**								**15111400**
					COC	GRA			
Prevotella intermedia		92.05	0.63	ARA 1	1	1			
Gemella morbillorum		7.33	0.45		99	99			
Lactobacillus fermentum		0.47	0.36		1	99			
15111502	**Good identification**								**15111502**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		100.0	0.40	IND 1				
15112000	**Low discrimination**							**15112000**
					GRA	SPO		
Prevotella intermedia		84.00	0.63	MAN 1	1	1		
Clostridium bifermentans		4.00	0.38		99	99		
Fusobacterium varium		3.84	0.32		1	1		
15112001	**Acceptable identification**							**15112001**
					GRA	SPO		
Clostridium sordellii		93.15	0.42	MNS 4	99	99		
Prevotella intermedia		5.91	0.29	MAN 1	1	1		
Clostridium bifermentans		0.28	0.04		99	99		
15112020	**Low discrimination**							**15112020**
					GRA	SPO	CAT	
Clostridium cadaveris		48.29	0.41	MLT 5	99	99	1	
Propionibacterium acnes		43.58	0.43	MAN 1	99	1	95	
Prevotella intermedia		4.94	0.29		1	1	1	
15113000	**Very good identification**							**15113000**
					COC	GRA	CAT	
Prevotella intermedia		98.67	0.63	MAN 1	1	1	1	
Gemella morbillorum		0.71	0.27		99	99	1	
Propionibacterium granulosum		0.34	0.29		1	99	90	
15113020	**Low discrimination**							**15113020**
					GRA	SPO	CAT	
Propionibacterium acnes		87.48	0.43	NIT 92	99	1	95	
Prevotella intermedia		9.90	0.29	MLT 1	1	1	1	
Clostridium cadaveris		0.98	0.07		99	99	1	
15114000	**Low discrimination**							**15114000**
					COC	GRA		
Prevotella intermedia		69.88	0.63	XYL 1	1	1		
Fusobacterium varium		13.19	0.42		1	1		
Gemella morbillorum		10.33	0.50		99	99		
15114001	**Acceptable identification**							**15114001**
					GRA	SPO		
Clostridium sordellii		91.70	0.42	MNS 4	99	99		
Prevotella intermedia		5.82	0.29	XYL 1	1	1		
Fusobacterium varium		1.10	0.08		1	1		
15114020	**Low discrimination**							**15114020**
					COC	GRA	SPO	
Clostridium cadaveris		74.68	0.41	MLT 5	1	99	99	
Gemella morbillorum		12.42	0.33	XYL 1	99	99	1	
Prevotella intermedia		7.63	0.29		1	1	1	
15115000	**Good identification**							**15115000**
					COC	GRA		
Prevotella intermedia		92.06	0.63	XYL 1	1	1		
Gemella morbillorum		7.33	0.45		99	99		
Lactobacillus fermentum		0.28	0.32		1	99		
15115102	**Good identification**							**15115102**
Lactobacillus fermentum		100.0	0.36	IND 1				
15115500	**Good identification**							**15115500**
Lactobacillus fermentum		95.66	0.36	IND 1				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		4.34	0.02	SUC 87				
15115502	**Good identification**							**15115502**
Lactobacillus fermentum		100.0	0.36	IND 1				
15115722	**Good identification**							**15115722**
Bacteroides eggerthii		99.90	0.39	ESL 90				
Bacteroides thetaiotaomicron		0.10	-.06	FRU 1				
15120000	**Good identification**							**15120000**
					GRA	SPO		
Clostridium bifermentans		90.86	0.60	TRE 1	99	99		
Prevotella disiens		4.32	0.32		1	1		
Clostridium botulinum biovar A		1.63	0.30		99	99		
15120001	**Acceptable identification**							**15120001**
					GRA	SPO		
Clostridium sordellii		94.33	0.42	MNS 4	99	99		
Clostridium bifermentans		5.41	0.25	TRE 1	99	99		
Prevotella disiens		0.26	-.03		1	1		
15120020	**Low discrimination**							**15120020**
					COC	SPO		
Clostridium cadaveris		89.46	0.53	MLT 5	1	99		
Clostridium bifermentans		10.03	0.39	TRE 5	1	99		
Gemella morbillorum		0.21	0.14		99	1		
15120040	**Low discrimination**							**15120040**
Clostridium botulinum biovar B		64.86	0.39	IND 1				
Clostridium bifermentans		30.83	0.36	SUC 90				
Clostridium botulinum biovar A		3.96	0.21					
15125722	**Good identification**							**15125722**
Bacteroides eggerthii		100.0	0.39	ESL 90				
15130000	**Low discrimination**							**15130000**
					GRA	SPO		
Prevotella intermedia		85.11	0.63	TRE 1	1	1		
Clostridium bifermentans		4.05	0.38		99	99		
Fusobacterium varium		3.90	0.32		1	1		
15130001	**Acceptable identification**							**15130001**
					GRA	SPO		
Clostridium sordellii		93.28	0.42	MNS 4	99	99		
Prevotella intermedia		5.92	0.29	TRE 1	1	1		
Clostridium bifermentans		0.28	0.04		99	99		
15130020	**Good identification**							**15130020**
					GRA	SPO	CAT	
Clostridium cadaveris		95.87	0.53	MLT 5	99	99	1	
Prevotella intermedia		1.88	0.29	TRE 5	1	1	1	
Propionibacterium granulosum		0.64	0.29		99	1	90	
15130040	**Good identification**							**15130040**
					GRA	SPO		
Clostridium botulinum biovar B		95.98	0.50	IND 1	99	99		
Prevotella intermedia		3.06	0.29	SUC 90	1	1		
Clostridium bifermentans		0.60	0.14		99	99		
15130050	**Good identification**							**15130050**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		100.0	0.44	IND 1				
15130060	**Good identification**							**15130060**
					SPO	CAT		
Clostridium botulinum biovar B		96.57	0.44	IND 1	99	1		
Clostridium cadaveris		3.22	0.19	SUC 90	99	1		
Propionibacterium acnes		0.22	0.02		1	95		
15130070	**Good identification**							**15130070**
Clostridium botulinum biovar B		100.0	0.39	IND 1				
15131000	**Very good identification**							**15131000**
					COC	GRA	CAT	
Prevotella intermedia		97.66	0.63	TRE 1	1	1	1	
Propionibacterium granulosum		1.35	0.39		1	99	90	
Gemella morbillorum		0.71	0.27		99	99	1	
15131120	**Low discrimination**							**15131120**
					GRA	SPO		
Clostridium perfringens		53.66	0.49	IND 1	99	84		
Clostridium septicum		45.38	0.41	SUC 91	99	99		
Prevotella bivia		0.66	0.03		1	1		
15131122	**Good identification**							**15131122**
Clostridium perfringens		96.46	0.43	IND 1				
Clostridium septicum		3.54	0.12	SUC 91				
15131160	**Good identification**							**15131160**
Clostridium perfringens		95.71	0.36	IND 1				
Clostridium septicum		4.29	0.07	SUC 91				
15132000	**Low discrimination**							**15132000**
					GRA	SPO	CAT	
Propionibacterium granulosum		47.05	0.37	IND 1	99	1	90	
Prevotella intermedia		45.63	0.29	SUC 93	1	1	1	
Clostridium bifermentans		2.17	0.04		99	99	1	
15133000	**Low discrimination**							**15133000**
					GRA	CAT		
Propionibacterium granulosum		50.77	0.37	IND 1	99	90		
Prevotella intermedia		49.23	0.29	SUC 93	1	1		
15140000	**Good identification**							**15140000**
					COC	GRA	SPO	
Clostridium bifermentans		92.93	0.60	CEL 1	1	99	99	
Prevotella disiens		4.42	0.32		1	1	1	
Gemella morbillorum		1.13	0.30		99	99	1	
15140001	**Acceptable identification**							**15140001**
					GRA	SPO		
Clostridium sordellii		94.33	0.42	MNS 4	99	99		
Clostridium bifermentans		5.41	0.25	CEL 1	99	99		
Prevotella disiens		0.26	-.03		1	1		
15140020	**Low discrimination**							**15140020**
					COC	SPO		
Clostridium cadaveris		62.11	0.41	MLT 5	1	99		
Clostridium bifermentans		36.29	0.39	CEL 1	1	99		
Gemella morbillorum		0.77	0.14		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
15140040	**Good identification**						**15140040**
					GRA	SPO	
Clostridium bifermentans		98.17	0.36	CEL 1	99	99	
Prevotella disiens		1.13	-.03	SOR 4	1	1	
Clostridium botulinum biovar B		0.70	-.03		99	99	
15141722	**Very good identification**						**15141722**
Bacteroides eggerthii		100.0	0.64	ESL 90			
15144722	**Good identification**						**15144722**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145322	**Good identification**						**15145322**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145522	**Good identification**						**15145522**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145622	**Good identification**						**15145622**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145702	**Good identification**						**15145702**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145720	**Good identification**						**15145720**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145722	**Very good identification**						**15145722**
Bacteroides eggerthii		100.0	0.84	ESL 90			
15145723	**Good identification**						**15145723**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145732	**Good identification**						**15145732**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15145762	**Good identification**						**15145762**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15147722	**Good identification**						**15147722**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15150000	**Low discrimination**						**15150000**
					GRA	SPO	
Prevotella intermedia		86.78	0.63	CEL 1	1	1	
Clostridium bifermentans		4.13	0.38		99	99	
Fusobacterium varium		3.97	0.32		1	1	
15150001	**Acceptable identification**						**15150001**
					GRA	SPO	
Clostridium sordellii		93.28	0.42	MNS 4	99	99	
Prevotella intermedia		5.92	0.29	CEL 1	1	1	
Clostridium bifermentans		0.28	0.04		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
15150020	**Low discrimination**						**15150020**
					GRA	SPO	
Clostridium cadaveris		85.23	0.41	MLT 5	99	99	
Prevotella intermedia		8.71	0.29	CEL 1	1	1	
Clostridium bifermentans		2.62	0.17		99	99	
15151000	**Very good identification**						**15151000**
					COC	GRA	
Prevotella intermedia		98.87	0.63	CEL 1	1	1	
Gemella morbillorum		0.72	0.27		99	99	
Lactobacillus jensenii		0.25	0.19		1	99	
15151120	**Good identification**						**15151120**
					GRA	SPO	
Clostridium septicum		95.92	0.41	IND 1	99	99	
Clostridium perfringens		2.41	0.20		99	84	
Prevotella bivia		1.33	0.03		1	1	
15155722	**Good identification**						**15155722**
Bacteroides eggerthii		99.95	0.50	ESL 90			
Bacteroides thetaiotaomicron		0.05	-.01	FRU 1			
15165722	**Good identification**						**15165722**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15171120	**Good identification**						**15171120**
					COC	SPO	
Clostridium septicum		98.00	0.49	IND 1	1	99	
Clostridium perfringens		1.53	0.25		1	84	
Atopobium parvulum		0.47	0.16		99	1	
15200000	**Very good identification**						**15200000**
					GRA	SPO	
Clostridium bifermentans		98.07	0.71	SAL 10	99	99	
Fusobacterium varium		1.05	0.32		1	1	
Clostridium cadaveris		0.44	0.29		99	99	
15200001	**Low discrimination**						**15200001**
					COC	SPO	
Clostridium bifermentans		65.34	0.37	SAL 10	1	99	
Clostridium sordellii		26.85	0.26	URE 1	1	99	
Anaerococcus prevotii		6.58	0.15		99	1	
15200002	**Good identification**						**15200002**
					GRA	SPO	
Clostridium bifermentans		96.02	0.37	SAL 10	99	99	
Atopobium minutum		2.95	0.19	BGL 1	99	1	
Fusobacterium varium		1.03	-.03		1	1	
15200010	**Good identification**						**15200010**
					GRA	SPO	
Clostridium bifermentans		97.49	0.37	SAL 10	99	99	
Fusobacterium varium		1.05	-.03	MLZ 1	1	1	
Clostridium novyi biovar A		0.80	0.14		99	99	
15200020	**Low discrimination**						**15200020**
					SPO	CAT	
Clostridium cadaveris		87.22	0.63	SAL 1	99	1	
Clostridium bifermentans		12.64	0.51		99	1	
Propionibacterium acnes		0.07	0.14		1	95	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
15200040	**Good identification**							**15200040**
					GRA	SPO		
Clostridium bifermentans		99.48	0.47	SAL 10	99	99		
Fusobacterium varium		0.26	-.03	SOR 4	1	1		
Clostridium novyi biovar A		0.13	0.11		99	99		
15200100	**Good identification**							**15200100**
					GRA	SPO		
Clostridium bifermentans		98.94	0.37	SAL 10	99	99		
Fusobacterium varium		1.06	-.03	LAC 1	1	1		
15200200	**Good identification**							**15200200**
					GRA	SPO		
Clostridium bifermentans		98.94	0.37	SAL 10	99	99		
Fusobacterium varium		1.06	-.03	RHA 1	1	1		
15200400	**Good identification**							**15200400**
					COC	GRA	SPO	
Clostridium bifermentans		98.45	0.37	SAL 10	1	99	99	
Fusobacterium varium		1.06	-.03	ARA 1	1	1	1	
Gemella morbillorum		0.49	0.01		99	99	1	
15201000	**Low discrimination**							**15201000**
					COC	GRA	SPO	
Clostridium bifermentans		82.57	0.37	SAL 10	1	99	99	
Fusobacterium necrophorum subsp. necrophorum		13.28	0.17	GAL 1	1	1	1	
Gemella morbillorum		2.00	0.13		99	99	1	
15202000	**Good identification**							**15202000**
					GRA	SPO		
Clostridium bifermentans		97.17	0.37	SAL 10	99	99		
Clostridium difficile		1.24	0.02	MAN 1	99	99		
Fusobacterium varium		1.04	-.03		1	1		
15204000	**Good identification**							**15204000**
					COC	GRA	SPO	
Clostridium bifermentans		95.31	0.37	SAL 10	1	99	99	
Fusobacterium varium		4.22	0.08	XYL 1	1	1	1	
Gemella morbillorum		0.48	0.01		99	99	1	
15210000	**Good identification**							**15210000**
					GRA	SPO		
Fusobacterium varium		94.24	0.66	SAL 1	1	1		
Clostridium bifermentans		4.67	0.49		99	99		
Clostridium cadaveris		0.40	0.29		99	99		
15210020	**Low discrimination**							**15210020**
					GRA	SPO	CAT	
Clostridium cadaveris		89.53	0.63	SAL 1	99	99	1	
Propionibacterium acnes		7.56	0.48		99	1	95	
Fusobacterium varium		2.18	0.32		1	1	1	
15210060	**Low discrimination**							**15210060**
					SPO	CAT		
Propionibacterium acnes		63.48	0.36	NIT 92	1	95		
Clostridium cadaveris		34.59	0.29	SAL 1	99	1		
Clostridium bifermentans		1.09	0.05		99	1		
15211020	**Low discrimination**							**15211020**
					GRA	SPO	CAT	
Propionibacterium acnes		88.98	0.48	NIT 92	99	1	95	
Clostridium cadaveris		10.64	0.29	SAL 1	99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium varium		0.26	-.03		1	1	1	
15211060	**Good identification**							**15211060**
Propionibacterium acnes		100.0	0.36	NIT 92				
15211100	**Good identification**							**15211100**
Fusobacterium mortiferum		97.77	0.45	IND 1				
Fusobacterium varium		2.23	-.03					
15212000	**Low discrimination**							**15212000**
					GRA	SPO		
Clostridium difficile		42.45	0.37	IND 1	99	99		
Fusobacterium varium		35.77	0.32		1	1		
Eubacterium limosum		18.81	0.29		99	1		
15212010	**Good identification**							**15212010**
					GRA	SPO		
Clostridium difficile		99.74	0.48	IND 1	99	99		
Fusobacterium varium		0.17	-.03		1	1		
Eubacterium limosum		0.09	-.06		99	1		
15212020	**Low discrimination**							**15212020**
					SPO	CAT		
Propionibacterium acnes		81.52	0.43	NIT 92	1	95		
Clostridium cadaveris		17.33	0.29	SAL 1	99	1		
Clostridium difficile		0.50	0.02		99	1		
15213020	**Good identification**							**15213020**
Propionibacterium acnes		100.0	0.43	NIT 92				
15214000	**Good identification**							**15214000**
					GRA	SPO		
Fusobacterium varium		98.29	0.42	SAL 1	1	1		
Clostridium bifermentans		1.18	0.15	XYL 4	99	99		
Fusobacterium mortiferum		0.23	0.17		1	1		
15220000	**Good identification**							**15220000**
					GRA	SPO		
Clostridium bifermentans		96.72	0.37	SAL 10	99	99		
Clostridium cadaveris		2.24	0.07	TRE 1	99	99		
Fusobacterium varium		1.04	-.03		1	1		
15220020	**Good identification**							**15220020**
Clostridium cadaveris		97.29	0.41	SAL 1				
Clostridium bifermentans		2.71	0.17	TRE 5				
15230020	**Good identification**							**15230020**
					GRA	SPO	CAT	
Clostridium cadaveris		97.95	0.41	SAL 1	99	99	1	
Propionibacterium acnes		1.59	0.14	TRE 5	99	1	95	
Fusobacterium varium		0.46	-.03		1	1	1	
15231100	**Good identification**							**15231100**
					GRA	SPO		
Fusobacterium mortiferum		95.20	0.36	IND 1	1	1		
Clostridium septicum		4.80	-.04		99	99		
15240000	**Good identification**							**15240000**
					GRA	SPO		
Clostridium bifermentans		98.94	0.37	SAL 10	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium varium		1.06	-.03	CEL 1	1	1	
15243000	**Good identification**						**15243000**
Clostridium innocuum		100.0	0.37	IND 1			
15243002	**Good identification**						**15243002**
Clostridium innocuum		100.0	0.37	IND 1			
15251100	**Low discrimination**						**15251100**
					GRA	SPO	
Fusobacterium mortiferum		79.44	0.37	IND 1	1	1	
Lactobacillus catenaformis		16.77	0.15		99	1	
Clostridium septicum		3.79	-.03		99	99	
15252000	**Acceptable identification**						**15252000**
					GRA	SPO	
Clostridium innocuum		91.21	0.37	IND 1	99	99	
Clostridium difficile		7.05	0.15	GAL 95	99	99	
Fusobacterium varium		1.14	-.03		1	1	
15252002	**Good identification**						**15252002**
Clostridium innocuum		100.0	0.37	IND 1			
15253000	**Very good identification**						**15253000**
Clostridium innocuum		100.0	0.59	IND 1			
15253002	**Very good identification**						**15253002**
Clostridium innocuum		100.0	0.59	IND 1			
15253010	**Good identification**						**15253010**
Clostridium innocuum		100.0	0.35	IND 1			
15253012	**Good identification**						**15253012**
Clostridium innocuum		100.0	0.35	IND 1			
15253400	**Good identification**						**15253400**
Clostridium innocuum		100.0	0.46	IND 1			
15253402	**Good identification**						**15253402**
Clostridium innocuum		100.0	0.46	IND 1			
15257000	**Good identification**						**15257000**
Clostridium innocuum		100.0	0.37	IND 1			
15257002	**Good identification**						**15257002**
Clostridium innocuum		100.0	0.37	IND 1			
15271120	**Good identification**						**15271120**
Clostridium septicum		100.0	0.39	IND 1			
15273000	**Good identification**						**15273000**
Clostridium innocuum		100.0	0.51	IND 1			
15273002	**Good identification**						**15273002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium innocuum		100.0	0.51	IND 1				
15273400	**Good identification**							**15273400**
Clostridium innocuum		100.0	0.38	IND 1				
15273402	**Good identification**							**15273402**
Clostridium innocuum		100.0	0.38	IND 1				
15300000	**Very good identification**							**15300000**
					COC	GRA	SPO	
Clostridium bifermentans		99.31	0.77	SAL 10	1	99	99	
Prevotella disiens		0.43	0.32		1	1	1	
Gemella morbillorum		0.11	0.30		99	99	1	
15300001	**Low discrimination**							**15300001**
					GRA	SPO		
Clostridium sordellii		61.13	0.42	MNS 4	99	99		
Clostridium bifermentans		38.57	0.43	SAL 1	99	99		
Prevotella disiens		0.17	-.03		1	1		
15300002	**Good identification**							**15300002**
					GRA	SPO		
Clostridium bifermentans		99.57	0.43	SAL 10	99	99		
Prevotella disiens		0.43	-.03	BGL 1	1	1		
15300010	**Good identification**							**15300010**
					GRA	SPO		
Clostridium bifermentans		99.23	0.43	SAL 10	99	99		
Prevotella disiens		0.43	-.03	MLZ 1	1	1		
Clostridium novyi biovar A		0.35	0.14		99	99		
15300020	**Low discrimination**							**15300020**
					COC	SPO		
Clostridium bifermentans		86.24	0.57	SAL 10	1	99		
Clostridium cadaveris		13.42	0.41	NAG 6	1	99		
Gemella morbillorum		0.17	0.14		99	1		
15300040	**Good identification**							**15300040**
					GRA	SPO		
Clostridium bifermentans		99.78	0.54	SAL 10	99	99		
Prevotella disiens		0.10	-.03	SOR 4	1	1		
Clostridium novyi biovar A		0.06	0.11		99	99		
15300100	**Good identification**							**15300100**
					GRA	SPO		
Clostridium bifermentans		98.70	0.43	SAL 10	99	99		
Prevotella bivia		0.88	-.03	LAC 1	1	1		
Prevotella disiens		0.43	-.03		1	1		
15300200	**Good identification**							**15300200**
					GRA	SPO		
Clostridium bifermentans		99.57	0.43	SAL 10	99	99		
Prevotella disiens		0.43	-.03	RHA 1	1	1		
15300400	**Good identification**							**15300400**
					COC	GRA	SPO	
Clostridium bifermentans		98.38	0.43	SAL 10	1	99	99	
Gemella morbillorum		1.20	0.14	ARA 1	99	99	1	
Prevotella disiens		0.43	-.03		1	1	1	
15301000	**Low discrimination**							**15301000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Clostridium bifermentans		85.72	0.43	SAL 10	1	99	99	
Prevotella intermedia		8.61	0.29	GAL 1	1	1	1	
Gemella morbillorum		5.05	0.26		99	99	1	
15302000	**Good identification**							**15302000**
					GRA	SPO		
Clostridium bifermentans		99.57	0.43	SAL 10	99	99		
Prevotella disiens		0.43	-.03	MAN 1	1	1		
15304000	**Good identification**							**15304000**
					COC	GRA	SPO	
Clostridium bifermentans		98.38	0.43	SAL 10	1	99	99	
Gemella morbillorum		1.20	0.14	XYL 1	99	99	1	
Prevotella disiens		0.43	-.03		1	1	1	
15305722	**Good identification**							**15305722**
Bacteroides eggerthii		100.0	0.39	ESL 90				
15310000	**Low discrimination**							**15310000**
					GRA	SPO		
Prevotella intermedia		61.38	0.63	SAL 1	1	1		
Clostridium bifermentans		32.15	0.56		99	99		
Fusobacterium varium		2.81	0.32		1	1		
15310001	**Acceptable identification**							**15310001**
					GRA	SPO		
Clostridium sordellii		90.72	0.42	MNS 4	99	99		
Prevotella intermedia		5.75	0.29	SAL 1	1	1		
Clostridium bifermentans		3.01	0.21		99	99		
15310020	**Low discrimination**							**15310020**
					GRA	SPO		
Clostridium cadaveris		67.53	0.41	MLT 5	99	99		
Clostridium bifermentans		22.84	0.35	SAL 1	99	99		
Prevotella intermedia		6.90	0.29		1	1		
15311000	**Very good identification**							**15311000**
					COC	GRA	SPO	
Prevotella intermedia		98.30	0.63	SAL 1	1	1	1	
Gemella morbillorum		0.71	0.27		99	99	1	
Clostridium bifermentans		0.52	0.21		1	99	99	
15311120	**Low discrimination**							**15311120**
					GRA	SPO		
Clostridium septicum		88.47	0.43	IND 1	99	99		
Clostridium perfringens		10.56	0.34		99	84		
Prevotella bivia		0.97	0.03		1	1		
15320000	**Good identification**							**15320000**
					GRA	SPO		
Clostridium bifermentans		99.57	0.43	SAL 10	99	99		
Prevotella disiens		0.43	-.03	TRE 1	1	1		
15331120	**Acceptable identification**							**15331120**
					COC	SPO		
Clostridium septicum		92.93	0.51	IND 1	1	99		
Clostridium perfringens		6.87	0.38		1	84		
Atopobium parvulum		0.20	0.11		99	1		
15340000	**Good identification**							**15340000**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		99.57	0.43	SAL 10	99	99	
Prevotella disiens		0.43	-.03	CEL 1	1	1	
15345722	**Good identification**						**15345722**
Bacteroides eggerthii		100.0	0.50	ESL 90			
15351000	**Low discrimination**						**15351000**
					GRA		
Lactobacillus jensenii		71.32	0.41	IND 1	99		
Prevotella intermedia		15.23	0.29	SUC 90	1		
Lactobacillus catenaformis		12.91	0.29		99		
15351002	**Low discrimination**						**15351002**
Lactobacillus jensenii		76.87	0.38	IND 1			
Lactobacillus catenaformis		20.88	0.29	SUC 90			
Lactobacillus acidophilus		1.21	0.05				
15351120	**Good identification**						**15351120**
					SPO		
Clostridium septicum		98.07	0.51	IND 1	99		
Clostridium paraputrificum		0.89	0.14		99		
Lactobacillus catenaformis		0.63	0.21		1		
15371000	**Good identification**						**15371000**
Lactobacillus jensenii		100.0	0.50	IND 1			
15371002	**Good identification**						**15371002**
Lactobacillus jensenii		98.28	0.47	IND 1			
Lactobacillus acidophilus		1.72	0.14	SUC 90			
15371010	**Good identification**						**15371010**
Lactobacillus jensenii		100.0	0.41	IND 1			
15371012	**Good identification**						**15371012**
Lactobacillus jensenii		100.0	0.38	IND 1			
15371020	**Low discrimination**						**15371020**
					SPO		
Lactobacillus jensenii		70.09	0.39	IND 1	1		
Clostridium septicum		29.92	0.25	SUC 90	99		
15371022	**Acceptable identification**						**15371022**
					SPO		
Lactobacillus jensenii		92.32	0.36	IND 1	1		
Lactobacillus acidophilus		6.47	0.14	SUC 90	1		
Clostridium septicum		1.21	-.04		99		
15371120	**Very good identification**						**15371120**
					COC	SPO	
Clostridium septicum		99.64	0.60	IND 1	1	99	
Atopobium parvulum		0.16	0.18		99	1	
Clostridium perfringens		0.10	0.15		1	84	
15375100	**Good identification**						**15375100**
Atopobium parvulum		100.0	0.35	IND 1			
15400000	**Low discrimination**						**15400000**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		77.24	0.53	RAF 1	99	99		
Fusobacterium varium		9.12	0.32		1	1		
Clostridium cadaveris		3.78	0.29		99	99		
15400001	**Low discrimination**							**15400001**
					COC	SPO		
Anaerococcus prevotii		86.63	0.41	IND 9	99	1		
Clostridium sordellii		10.72	0.26	GLU 1	1	99		
Clostridium bifermentans		2.37	0.19		1	99		
15400020	**Very good identification**							**15400020**
					SPO	CAT		
Clostridium cadaveris		98.55	0.63	RAF 1	99	1		
Clostridium bifermentans		1.30	0.33		99	1		
Propionibacterium acnes		0.08	0.14		1	95		
15410000	**Low discrimination**							**15410000**
					GRA	SPO		
Fusobacterium varium		74.26	0.66	RAF 1	1	1		
Prevotella intermedia		24.59	0.66		1	1		
Clostridium bifermentans		0.34	0.31		99	99		
15410020	**Low discrimination**							**15410020**
					GRA	SPO	CAT	
Clostridium cadaveris		89.30	0.63	RAF 1	99	99	1	
Propionibacterium acnes		7.54	0.48		99	1	95	
Fusobacterium varium		2.18	0.32		1	1	1	
15410060	**Low discrimination**							**15410060**
					GRA	SPO	CAT	
Propionibacterium acnes		64.18	0.36	NIT 92	99	1	95	
Clostridium cadaveris		34.97	0.29	RAF 1	99	99	1	
Fusobacterium varium		0.85	-.03		1	1	1	
15411000	**Very good identification**							**15411000**
					GRA	CAT		
Prevotella intermedia		95.30	0.66	MLT 99	1	1		
Fusobacterium varium		2.91	0.32		1	1		
Propionibacterium granulosum		0.83	0.38		99	90		
15411020	**Low discrimination**							**15411020**
					GRA	SPO	CAT	
Propionibacterium acnes		80.28	0.48	NIT 92	99	1	95	
Clostridium cadaveris		9.60	0.29	RAF 1	99	99	1	
Prevotella intermedia		7.67	0.32		1	1	1	
15411060	**Good identification**							**15411060**
Propionibacterium acnes		100.0	0.36	NIT 92				
15411100	**Low discrimination**							**15411100**
					GRA			
Fusobacterium mortiferum		50.19	0.45	IND 1	1			
Prevotella intermedia		37.48	0.32		1			
Lactobacillus fermentum		11.19	0.35		99			
15411102	**Low discrimination**							**15411102**
					GRA			
Lactobacillus fermentum		66.74	0.35	IND 1	99			
Fusobacterium mortiferum		33.26	0.29	SUC 87	1			
15411500	**Low discrimination**							**15411500**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		85.68	0.35	IND 1	99		
Odoribacter splanchnicus		10.44	-.03	SUC 87	1		
Fusobacterium mortiferum		3.88	0.11		1		
15411502	**Good identification**						**15411502**
Lactobacillus fermentum		100.0	0.35	IND 1			
15412020	**Low discrimination**						**15412020**
					SPO	CAT	
Propionibacterium acnes		80.12	0.43	NIT 92	1	95	
Clostridium cadaveris		17.04	0.29	RAF 1	99	1	
Propionibacterium granulosum		2.21	0.26		1	90	
15413020	**Good identification**						**15413020**
Propionibacterium acnes		97.32	0.43	NIT 92			
Propionibacterium granulosum		2.68	0.26	RAF 1			
15414000	**Acceptable identification**						**15414000**
Fusobacterium varium		92.16	0.42	RAF 1			
Prevotella intermedia		7.40	0.32	XYL 4			
Fusobacterium mortiferum		0.21	0.17				
15420020	**Good identification**						**15420020**
Clostridium cadaveris		99.75	0.41	RAF 1			
Clostridium bifermentans		0.25	-.01	TRE 5			
15430002	**Good identification**						**15430002**
					GRA	CAT	
Propionibacterium granulosum		95.49	0.36	IND 1	99	90	
Fusobacterium varium		4.51	-.03	SUC 93	1	1	
15430020	**Low discrimination**						**15430020**
					SPO	CAT	
Clostridium cadaveris		89.30	0.41	RAF 1	99	1	
Propionibacterium granulosum		8.84	0.36	TRE 5	1	90	
Propionibacterium acnes		1.45	0.14		1	95	
15431000	**Low discrimination**						**15431000**
					GRA	CAT	
Propionibacterium granulosum		68.23	0.47	IND 1	99	90	
Prevotella intermedia		26.43	0.32	SUC 93	1	1	
Fusobacterium mortiferum		4.53	0.30		1	5	
15431002	**Good identification**						**15431002**
					GRA	CAT	
Propionibacterium granulosum		97.13	0.36	IND 1	99	90	
Fusobacterium mortiferum		2.87	0.13	SUC 93	1	5	
15431020	**Low discrimination**						**15431020**
					SPO	CAT	
Propionibacterium granulosum		78.99	0.36	IND 1	1	90	
Propionibacterium acnes		12.94	0.14	SUC 93	1	95	
Clostridium cadaveris		8.07	0.07		99	1	
15431100	**Acceptable identification**						**15431100**
					GRA	CAT	
Fusobacterium mortiferum		93.88	0.36	IND 1	1	5	
Propionibacterium granulosum		6.12	0.13		99	90	
15432000	**Good identification**						**15432000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Propionibacterium granulosum		97.67	0.45	IND 1	99	90		
Fusobacterium varium		1.53	-.03	SUC 93	1	1		
Eubacterium limosum		0.81	-.06		99	1		
15433000	**Good identification**							**15433000**
Propionibacterium granulosum		100.0	0.45	IND 1				
15435322	**Good identification**							**15435322**
Parabacteroides distasonis		100.0	0.39	IND 1				
15451100	**Good identification**							**15451100**
Fusobacterium mortiferum		100.0	0.37	IND 1				
15500000	**Low discrimination**							**15500000**
					GRA	SPO		
Prevotella intermedia		60.89	0.66	FRU 99	1	1		
Clostridium bifermentans		36.73	0.60		99	99		
Prevotella disiens		1.75	0.32		1	1		
15500001	**Low discrimination**							**15500001**
					GRA	SPO		
Clostridium sordellii		85.89	0.42	MNS 4	99	99		
Prevotella intermedia		8.17	0.32	RAF 1	1	1		
Clostridium bifermentans		4.93	0.25		99	99		
15500020	**Low discrimination**							**15500020**
					GRA	SPO		
Clostridium cadaveris		56.71	0.41	MLT 5	99	99		
Clostridium bifermentans		33.13	0.39	RAF 1	99	99		
Prevotella intermedia		8.69	0.32		1	1		
15500040	**Low discrimination**							**15500040**
					GRA	SPO		
Clostridium bifermentans		70.40	0.36	RAF 1	99	99		
Prevotella intermedia		28.30	0.32	SOR 4	1	1		
Prevotella disiens		0.81	-.03		1	1		
15501000	**Very good identification**							**15501000**
					COC	GRA	SPO	
Prevotella intermedia		98.94	0.66	FRU 99	1	1	1	
Clostridium bifermentans		0.60	0.25		1	99	99	
Gemella morbillorum		0.39	0.26		99	99	1	
15505722	**Good identification**							**15505722**
Bacteroides eggerthii		99.90	0.39	ESL 90				
Bacteroides thetaiotaomicron		0.10	-.06	RAF 1				
15510000	**Excellent identification**							**15510000**
					GRA	SPO		
Prevotella intermedia		99.89	1.00		1	1		
Clostridium bifermentans		0.03	0.38		99	99		
Prevotella disiens		0.03	0.32		1	1		
15510001	**Good identification**							**15510001**
					GRA	SPO		
Prevotella intermedia		90.33	0.66	URE 1	1	1		
Clostridium sordellii		9.59	0.42		99	99		
Clostridium bifermentans		0.03	0.04		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
15510002	**Very good identification**							**15510002**
					GRA	SPO	CAT	
Prevotella intermedia		99.81	0.66	BGL 1	1	1	1	
Propionibacterium granulosum		0.10	0.22		99	1	90	
Clostridium bifermentans		0.03	0.04		99	99	1	
15510010	**Very good identification**							**15510010**
					GRA	SPO	CAT	
Prevotella intermedia		99.86	0.66	MLZ 1	1	1	1	
Propionibacterium granulosum		0.05	0.18		99	1	90	
Clostridium bifermentans		0.03	0.04		99	99	1	
15510020	**Good identification**							**15510020**
					GRA	SPO		
Prevotella intermedia		93.32	0.66	NAG 1	1	1		
Clostridium cadaveris		6.15	0.41		99	99		
Clostridium bifermentans		0.19	0.17		99	99		
15510040	**Very good identification**							**15510040**
					GRA	SPO		
Prevotella intermedia		99.74	0.66	SOR 1	1	1		
Clostridium bifermentans		0.13	0.14		99	99		
Clostridium botulinum biovar B		0.07	0.07		99	99		
15510100	**Very good identification**							**15510100**
					GRA	SPO		
Prevotella intermedia		99.80	0.66	LAC 1	1	1		
Fusobacterium mortiferum		0.11	0.26		1	1		
Clostridium bifermentans		0.03	0.04		99	99		
15510200	**Very good identification**							**15510200**
					GRA	SPO		
Prevotella intermedia		99.91	0.66	RHA 1	1	1		
Clostridium bifermentans		0.03	0.04		99	99		
Prevotella disiens		0.03	-.03		1	1		
15510400	**Very good identification**							**15510400**
					COC	GRA	SPO	
Prevotella intermedia		99.81	0.66	ARA 1	1	1	1	
Gemella morbillorum		0.10	0.15		99	99	1	
Clostridium bifermentans		0.03	0.04		1	99	99	
15511000	**Excellent identification**							**15511000**
					GRA	CAT		
Prevotella intermedia		99.98	1.00		1	1		
Lactobacillus fermentum		0.01	0.40		99	1		
Propionibacterium granulosum		0.00	0.33		99	90		
15511001	**Very good identification**							**15511001**
					GRA	SPO		
Prevotella intermedia		98.16	0.66	URE 1	1	1		
Clostridium sordellii		1.84	0.29		99	99		
15511002	**Very good identification**							**15511002**
					GRA	CAT		
Prevotella intermedia		99.35	0.66	BGL 1	1	1		
Lactobacillus fermentum		0.55	0.40		99	1		
Propionibacterium granulosum		0.10	0.22		99	90		
15511010	**Very good identification**							**15511010**
					GRA	CAT		
Prevotella intermedia		99.95	0.66	MLZ 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		0.05	0.18		99	90		
15511020	**Very good identification**							**15511020**
					GRA			
Prevotella intermedia		99.34	0.66	NAG 1	1			
Lactobacillus fermentum		0.18	0.32		99			
Clostridium perfringens		0.13	0.17		99			
15511040	**Very good identification**							**15511040**
Prevotella intermedia		100.0	0.66	SOR 1				
15511100	**Very good identification**							**15511100**
					GRA			
Prevotella intermedia		98.83	0.66	LAC 1	1			
Lactobacillus fermentum		0.89	0.44		99			
Fusobacterium mortiferum		0.17	0.30		1			
15511200	**Very good identification**							**15511200**
Prevotella intermedia		100.0	0.66	RHA 1				
15511400	**Very good identification**							**15511400**
					COC	GRA		
Prevotella intermedia		99.40	0.66	ARA 1	1	1		
Lactobacillus fermentum		0.55	0.40		1	99		
Gemella morbillorum		0.05	0.11		99	99		
15511502	**Good identification**							**15511502**
Lactobacillus fermentum		97.50	0.44	IND 1				
Bifidobacterium longum subsp. longum		2.50	0.09	SUC 87				
15511520	**Low discrimination**							**15511520**
					GRA			
Lactobacillus fermentum		66.21	0.35	IND 1	99			
Bifidobacterium longum subsp. longum		20.38	0.20	SUC 87	99			
Odoribacter splanchnicus		8.07	-.03		1			
15511522	**Low discrimination**							**15511522**
Lactobacillus fermentum		76.47	0.35	IND 1				
Bifidobacterium longum subsp. longum		23.53	0.20	SUC 87				
15512000	**Very good identification**							**15512000**
					GRA	SPO	CAT	
Prevotella intermedia		99.60	0.66	MAN 1	1	1	1	
Propionibacterium granulosum		0.29	0.30		99	1	90	
Clostridium bifermentans		0.03	0.04		99	99	1	
15513000	**Very good identification**							**15513000**
					GRA	CAT		
Prevotella intermedia		99.71	0.66	MAN 1	1	1		
Propionibacterium granulosum		0.29	0.30		99	90		
15514000	**Very good identification**							**15514000**
					COC	GRA		
Prevotella intermedia		99.71	0.66	XYL 1	1	1		
Fusobacterium varium		0.13	0.08		1	1		
Gemella morbillorum		0.10	0.15		99	99		
15515000	**Very good identification**							**15515000**
					COC	GRA		
Prevotella intermedia		99.62	0.66	XYL 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		0.32	0.36		1	99		
Gemella morbillorum		0.05	0.11		99	99		
15515102	**Good identification**							**15515102**
Lactobacillus fermentum		100.0	0.40	IND 1				
15515322	**Acceptable identification**							**15515322**
					CAT			
Parabacteroides distasonis		94.53	0.37	IND 1	76			
Prevotella oralis		3.10	-.01		1			
Bacteroides thetaiotaomicron		2.36	-.06		89			
15515402	**Good identification**							**15515402**
Lactobacillus fermentum		100.0	0.36	IND 1				
15515500	**Low discrimination**							**15515500**
Lactobacillus fermentum		88.42	0.40	IND 1				
Bifidobacterium longum subsp. longum		11.58	0.17	SUC 87				
15515502	**Low discrimination**							**15515502**
Lactobacillus fermentum		88.42	0.40	IND 1				
Bifidobacterium longum subsp. longum		11.58	0.17	SUC 87				
15515720	**Good identification**							**15515720**
					CAT			
Bacteroides vulgatus		99.74	0.36	IND 1	3			
Bacteroides thetaiotaomicron		0.26	-.06	SUC 96	89			
15530000	**Very good identification**							**15530000**
					GRA	SPO	CAT	
Prevotella intermedia		98.74	0.66	TRE 1	1	1	1	
Propionibacterium granulosum		1.16	0.41		99	1	90	
Clostridium bifermentans		0.03	0.04		99	99	1	
15531000	**Very good identification**							**15531000**
					GRA	CAT		
Prevotella intermedia		98.82	0.66	TRE 1	1	1		
Propionibacterium granulosum		1.16	0.41		99	90		
Fusobacterium mortiferum		0.02	0.14		1	5		
15531120	**Acceptable identification**							**15531120**
					GRA	SPO		
Clostridium perfringens		94.47	0.36	IND 1	99	84		
Clostridium septicum		4.24	0.07	SUC 91	99	99		
Prevotella melaninogenica		1.29	-.03		1	1		
15535122	**Good identification**							**15535122**
Parabacteroides distasonis		98.38	0.41	IND 1				
Bacteroides thetaiotaomicron		1.62	-.06					
15535132	**Good identification**							**15535132**
Parabacteroides distasonis		100.0	0.36	IND 1				
15535322	**Acceptable identification**							**15535322**
Parabacteroides distasonis		93.21	0.49	IND 1				
Bacteroides thetaiotaomicron		6.79	0.13					
15535332	**Good identification**							**15535332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		95.00	0.44	IND 1			
Bacteroides thetaiotaomicron		5.00	0.06				
15535722	**Acceptable identification**						**15535722**
Bacteroides thetaiotaomicron		92.03	0.47	ESL 92			
Parabacteroides distasonis		7.82	0.45	SUC 98			
Bacteroides ovatus		0.15	0.00				
15535732	**Low discrimination**						**15535732**
Bacteroides thetaiotaomicron		89.25	0.40	ESL 92			
Parabacteroides distasonis		10.50	0.40	SUC 98			
Bacteroides ovatus		0.25	-.03				
15545722	**Good identification**						**15545722**
Bacteroides eggerthii		99.95	0.50	ESL 90			
Bacteroides thetaiotaomicron		0.05	-.01	RAF 1			
15550000	**Very good identification**						**15550000**
					GRA	SPO	
Prevotella intermedia		99.89	0.66	CEL 1	1	1	
Clostridium bifermentans		0.03	0.04		99	99	
Prevotella disiens		0.03	-.03		1	1	
15551000	**Very good identification**						**15551000**
Prevotella intermedia		99.98	0.66	CEL 1			
Fusobacterium mortiferum		0.02	0.15				
15575122	**Acceptable identification**						**15575122**
Parabacteroides distasonis		93.91	0.36	IND 1			
Bacteroides thetaiotaomicron		6.10	-.01				
15575322	**Low discrimination**						**15575322**
Parabacteroides distasonis		73.84	0.44	IND 1			
Bacteroides thetaiotaomicron		21.22	0.19				
Bacteroides ovatus		4.94	0.08				
15575332	**Low discrimination**						**15575332**
Parabacteroides distasonis		77.49	0.39	IND 1			
Bacteroides thetaiotaomicron		16.09	0.12				
Bacteroides ovatus		6.42	0.05				
15575722	**Acceptable identification**						**15575722**
Bacteroides thetaiotaomicron		91.11	0.53	ESL 92			
Bacteroides ovatus		6.93	0.34	SUC 98			
Parabacteroides distasonis		1.96	0.40				
15575732	**Low discrimination**						**15575732**
Bacteroides thetaiotaomicron		86.08	0.46	ESL 92			
Bacteroides ovatus		11.22	0.31	SUC 98			
Parabacteroides distasonis		2.57	0.36				
15600000	**Good identification**						**15600000**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		98.94	0.37	SAL 10	99	99	
Fusobacterium varium		1.06	-.03	RAF 1	1	1	
15610100	**Good identification**						**15610100**
Fusobacterium mortiferum		98.27	0.47	IND 1			
Fusobacterium varium		1.73	-.03				
15611000	**Low discrimination**						**15611000**
Fusobacterium mortiferum		51.95	0.44	IND 1			
Prevotella intermedia		46.63	0.32				
Fusobacterium varium		1.42	-.03				
15611100	**Good identification**						**15611100**
Fusobacterium mortiferum		100.0	0.50	IND 1			
15630100	**Good identification**						**15630100**
Fusobacterium mortiferum		100.0	0.38	IND 1			
15631100	**Good identification**						**15631100**
Fusobacterium mortiferum		100.0	0.41	IND 1			
15635322	**Good identification**						**15635322**
Parabacteroides distasonis		100.0	0.40	IND 1			
15635332	**Good identification**						**15635332**
Parabacteroides distasonis		100.0	0.35	IND 1			
15635722	**Low discrimination**						**15635722**
Parabacteroides distasonis		69.75	0.36	IND 1			
Bacteroides thetaiotaomicron		30.25	0.14				
15650100	**Good identification**						**15650100**
Fusobacterium mortiferum		100.0	0.39	IND 1			
15651000	**Good identification**						**15651000**
Fusobacterium mortiferum		100.0	0.36	IND 1			
15651100	**Good identification**						**15651100**
Fusobacterium mortiferum		100.0	0.42	IND 1			
15675322	**Good identification**						**15675322**
Parabacteroides distasonis		100.0	0.35	IND 1			
15700000	**Low discrimination**						**15700000**
					GRA	SPO	
Clostridium bifermentans		86.58	0.43	SAL 10	99	99	
Prevotella intermedia		13.05	0.32	RAF 1	1	1	
Prevotella disiens		0.37	-.03		1	1	
15710000	**Very good identification**						**15710000**
					GRA	SPO	
Prevotella intermedia		99.50	0.66	SAL 1	1	1	
Clostridium bifermentans		0.35	0.21		99	99	
Fusobacterium mortiferum		0.09	0.25		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
15711000	**Very good identification**					**15711000**
Prevotella intermedia		99.86	0.66	SAL 1		
Fusobacterium mortiferum		0.14	0.28			
15715322	**Acceptable identification**					**15715322**
Parabacteroides distasonis		94.89	0.38	IND 1		
Prevotella oralis		5.11	0.04			
15735122	**Good identification**					**15735122**
Parabacteroides distasonis		100.0	0.41	IND 1		
15735132	**Good identification**					**15735132**
Parabacteroides distasonis		100.0	0.37	IND 1		
15735322	**Good identification**					**15735322**
Parabacteroides distasonis		98.83	0.49	IND 1		
Bacteroides thetaiotaomicron		1.17	0.00			
15735332	**Good identification**					**15735332**
Parabacteroides distasonis		100.0	0.45	IND 1		
15735522	**Low discrimination**					**15735522**
Parabacteroides distasonis		69.80	0.37	IND 1		
Bacteroides thetaiotaomicron		30.20	0.15			
15775122	**Acceptable identification**					**15775122**
Parabacteroides distasonis		90.39	0.37	IND 1		
Bacteroides ovatus		6.47	0.01			
Prevotella oralis		3.15	-.01			
15775322	**Low discrimination**					**15775322**
Parabacteroides distasonis		66.67	0.45	IND 1		
Bacteroides ovatus		30.21	0.23			
Bacteroides thetaiotaomicron		3.12	0.06			
15775332	**Low discrimination**					**15775332**
Parabacteroides distasonis		62.71	0.40	IND 1		
Bacteroides ovatus		35.18	0.20			
Bacteroides thetaiotaomicron		2.12	-.01			
15775722	**Low discrimination**					**15775722**
Bacteroides ovatus		73.45	0.49	ESL 91		
Bacteroides thetaiotaomicron		23.24	0.40	SUC 99		
Parabacteroides distasonis		3.07	0.41			
15775732	**Low discrimination**					**15775732**
Bacteroides ovatus		81.31	0.46	ESL 91		
Bacteroides thetaiotaomicron		15.01	0.33	SUC 99		
Parabacteroides distasonis		2.75	0.36			
15777722	**Good identification**					**15777722**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides ovatus		99.05	0.41	ESL 91				
Bacteroides thetaiotaomicron		0.95	0.06	SUC 99				
15777732	**Good identification**							**15777732**
Bacteroides ovatus		99.44	0.38	ESL 91				
Bacteroides thetaiotaomicron		0.56	-.01	SUC 99				
16000001	**Good identification**							**16000001**
					COC	GRA		
Anaerococcus prevotii		98.45	0.49	IND 9	99	99		
Fusobacterium nucleatum subsp. nucleatum		0.26	-.03	SUC 1	1	1		
Peptoniphilus asaccharolyticus		0.26	-.03		99	99		
16000003	**Good identification**							**16000003**
Anaerococcus prevotii		100.0	0.38	IND 9				
16000020	**Good identification**							**16000020**
					SPO			
Clostridium cadaveris		98.71	0.56	GLU 95	99			
Clostridium bifermentans		0.29	0.15	SUC 7	99			
Atopobium minutum		0.27	0.22		1			
16010000	**Good identification**							**16010000**
					GRA	SPO		
Fusobacterium varium		94.53	0.57	SUC 1	1	1		
Prevotella intermedia		3.82	0.41		1	1		
Clostridium cadaveris		0.52	0.22		99	99		
16010020	**Good identification**							**16010020**
					GRA	SPO		
Clostridium cadaveris		97.58	0.56	GLU 95	99	99		
Fusobacterium varium		1.81	0.23	SUC 7	1	1		
Atopobium minutum		0.25	0.22		99	1		
16011000	**Low discrimination**							**16011000**
					GRA			
Prevotella intermedia		74.27	0.41	GLU 90	1			
Fusobacterium varium		18.55	0.23	MLT 99	1			
Atopobium minutum		2.53	0.22		99			
16100000	**Low discrimination**							**16100000**
					COC	GRA	SPO	
Clostridium bifermentans		51.90	0.41	GLU 92	1	99	99	
Prevotella intermedia		36.10	0.41	SUC 1	1	1	1	
Gemella morbillorum		7.26	0.30		99	99	1	
16101000	**Low discrimination**							**16101000**
					COC	GRA	SPO	
Prevotella intermedia		88.33	0.41	GLU 90	1	1	1	
Gemella morbillorum		9.56	0.26	FRU 99	99	99	1	
Clostridium bifermentans		1.28	0.07		1	99	99	
16110000	**Very good identification**							**16110000**
					COC	GRA		
Prevotella intermedia		99.40	0.75	GLU 90	1	1		
Fusobacterium varium		0.25	0.23		1	1		
Gemella morbillorum		0.25	0.32		99	99		
16110001	**Low discrimination**							**16110001**
					GRA	SPO		
Prevotella intermedia		86.72	0.41	GLU 90	1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sordellii		13.28	0.20	URE 1	99	99		
16110002	**Good identification**							**16110002**
Prevotella intermedia		100.0	0.41	GLU 90				
16110010	**Good identification**							**16110010**
Prevotella intermedia		100.0	0.41	GLU 90				
16110020	**Low discrimination**							**16110020**
					COC	GRA	SPO	
Prevotella intermedia		57.46	0.41	GLU 90	1	1	1	
Clostridium cadaveris		40.70	0.34	NAG 1	1	99	99	
Gemella morbillorum		1.57	0.15		99	99	1	
16110040	**Low discrimination**							**16110040**
					GRA	SPO		
Prevotella intermedia		85.06	0.41	GLU 90	1	1		
Clostridium botulinum biovar B		14.94	0.24	SOR 1	99	99		
16110100	**Good identification**							**16110100**
					GRA			
Prevotella intermedia		99.45	0.41	GLU 90	1			
Actinomyces meyeri		0.55	0.05	LAC 1	99			
16110200	**Good identification**							**16110200**
Prevotella intermedia		100.0	0.41	GLU 90				
16110400	**Good identification**							**16110400**
					COC	GRA		
Prevotella intermedia		97.34	0.41	GLU 90	1	1		
Gemella morbillorum		2.66	0.15	ARA 1	99	99		
16111000	**Very good identification**							**16111000**
					COC	GRA		
Prevotella intermedia		99.84	0.75	GLU 90	1	1		
Gemella morbillorum		0.13	0.27		99	99		
Lactobacillus fermentum		0.01	0.16		1	99		
16111001	**Good identification**							**16111001**
					GRA	SPO		
Prevotella intermedia		97.37	0.41	GLU 90	1	1		
Clostridium sordellii		2.63	0.07	URE 1	99	99		
16111002	**Good identification**							**16111002**
					GRA			
Prevotella intermedia		99.37	0.41	GLU 90	1			
Lactobacillus fermentum		0.63	0.16	BGL 1	99			
16111010	**Good identification**							**16111010**
Prevotella intermedia		99.33	0.41	GLU 90				
Eubacterium saburreum		0.67	0.14	MLZ 1				
16111020	**Good identification**							**16111020**
					COC	GRA		
Prevotella intermedia		95.31	0.41	GLU 90	1	1		
Clostridium perfringens		1.83	0.12	NAG 1	1	99		
Gemella morbillorum		1.40	0.11		99	99		
16111040	**Good identification**							**16111040**
Prevotella intermedia		100.0	0.41	GLU 90				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
16111100	**Good identification**							**16111100**
					GRA			
Prevotella intermedia		95.20	0.41	GLU 90	1			
Eubacterium saburreum		1.84	0.21	LAC 1	85			
Lactobacillus fermentum		0.98	0.20		99			
16111200	**Good identification**							**16111200**
Prevotella intermedia		100.0	0.41	GLU 90				
16111400	**Good identification**							**16111400**
					COC	GRA		
Prevotella intermedia		97.94	0.41	GLU 90	1	1		
Gemella morbillorum		1.44	0.11	ARA 1	99	99		
Lactobacillus fermentum		0.62	0.16		1	99		
16112000	**Good identification**							**16112000**
					GRA	CAT		
Prevotella intermedia		99.16	0.41	GLU 90	1	1		
Propionibacterium granulosum		0.84	0.14	MAN 1	99	90		
16113000	**Good identification**							**16113000**
					GRA	CAT		
Prevotella intermedia		99.16	0.41	GLU 90	1	1		
Propionibacterium granulosum		0.84	0.14	MAN 1	99	90		
16114000	**Good identification**							**16114000**
					COC	GRA		
Prevotella intermedia		96.37	0.41	GLU 90	1	1		
Gemella morbillorum		2.63	0.15	XYL 1	99	99		
Fusobacterium varium		1.00	-.01		1	1		
16115000	**Good identification**							**16115000**
					COC	GRA		
Prevotella intermedia		97.38	0.41	GLU 90	1	1		
Gemella morbillorum		1.43	0.11	XYL 1	99	99		
Eubacterium saburreum		0.44	0.11		1	85		
16130000	**Good identification**							**16130000**
					GRA	SPO	CAT	
Prevotella intermedia		96.23	0.41	GLU 90	1	1	1	
Propionibacterium granulosum		3.26	0.24	TRE 1	99	1	90	
Clostridium botulinum biovar B		0.51	-.03		99	99	1	
16131000	**Good identification**							**16131000**
					CAT	GRA		
Prevotella intermedia		96.21	0.41	GLU 90	1	1		
Propionibacterium granulosum		3.26	0.24	TRE 1	90	99		
Eubacterium saburreum		0.53	0.12		1	85		
16150000	**Good identification**							**16150000**
Prevotella intermedia		100.0	0.41	GLU 90				
16151000	**Good identification**							**16151000**
					GRA			
Prevotella intermedia		99.59	0.41	GLU 90	1			
Lactobacillus jensenii		0.41	0.02	CEL 1	99			
16310000	**Good identification**							**16310000**
					GRA	SPO		
Prevotella intermedia		99.17	0.41	GLU 90	1	1		
Clostridium bifermentans		0.83	0.03	SAL 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
16311000	**Good identification**						**16311000**
					GRA		
Prevotella intermedia		99.59	0.41	GLU 90	1		
Lactobacillus jensenii		0.41	0.02	SAL 1	99		
16400001	**Good identification**						**16400001**
Anaerococcus prevotii		100.0	0.41	IND 9			
16410000	**Low discrimination**						**16410000**
					GRA	CAT	
Prevotella intermedia		83.90	0.44	GLU 90	1	1	
Fusobacterium varium		13.97	0.23	MLT 99	1	1	
Propionibacterium granulosum		1.79	0.24		99	90	
16411000	**Good identification**						**16411000**
					GRA	CAT	
Prevotella intermedia		97.34	0.44	GLU 90	1	1	
Propionibacterium granulosum		2.08	0.24	MLT 99	99	90	
Fusobacterium mortiferum		0.59	0.17		1	5	
16475722	**Good identification**						**16475722**
Bacteroides thetaiotaomicron		99.21	0.36	ESL 92			
Bacteroides ovatus		0.79	0.00	GLU 97			
16500000	**Good identification**						**16500000**
					GRA	SPO	
Prevotella intermedia		99.04	0.44	GLU 90	1	1	
Clostridium bifermentans		0.96	0.07	FRU 99	99	99	
16501000	**Good identification**						**16501000**
Prevotella intermedia		100.0	0.44	GLU 90			
16510000	**Very good identification**						**16510000**
					GRA	CAT	
Prevotella intermedia		99.99	0.78	GLU 90	1	1	
Propionibacterium granulosum		0.01	0.18		99	90	
16510001	**Good identification**						**16510001**
Prevotella intermedia		100.0	0.44	GLU 90			
16510002	**Good identification**						**16510002**
Prevotella intermedia		100.0	0.44	GLU 90			
16510010	**Good identification**						**16510010**
Prevotella intermedia		100.0	0.44	GLU 90			
16510020	**Good identification**						**16510020**
					GRA	SPO	
Prevotella intermedia		99.53	0.44	GLU 90	1	1	
Clostridium cadaveris		0.48	0.00	NAG 1	99	99	
16510040	**Good identification**						**16510040**
Prevotella intermedia		100.0	0.44	GLU 90			
16510100	**Good identification**						**16510100**
Prevotella intermedia		100.0	0.44	GLU 90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
16510200	**Good identification**						**16510200**
Prevotella intermedia		100.0	0.44	GLU 90			
16510400	**Good identification**						**16510400**
Prevotella intermedia		100.0	0.44	GLU 90			
16511000	**Very good identification**						**16511000**
					GRA	CAT	
Prevotella intermedia		99.98	0.78	GLU 90	1	1	
Propionibacterium granulosum		0.01	0.18		99	90	
Lactobacillus fermentum		0.01	0.20		99	1	
16511001	**Good identification**						**16511001**
Prevotella intermedia		100.0	0.44	GLU 90			
16511002	**Good identification**						**16511002**
					GRA		
Prevotella intermedia		99.32	0.44	GLU 90	1		
Lactobacillus fermentum		0.68	0.20	BGL 1	99		
16511010	**Good identification**						**16511010**
Prevotella intermedia		100.0	0.44	GLU 90			
16511020	**Good identification**						**16511020**
Prevotella intermedia		95.34	0.44	GLU 90			
Prevotella melaninogenica		4.66	0.15	NAG 1			
16511040	**Good identification**						**16511040**
Prevotella intermedia		100.0	0.44	GLU 90			
16511100	**Acceptable identification**						**16511100**
					GRA		
Prevotella intermedia		93.72	0.44	GLU 90	1		
Prevotella melaninogenica		4.58	0.15	LAC 1	1		
Lactobacillus fermentum		1.05	0.24		99		
16511120	**Good identification**						**16511120**
					GRA		
Prevotella melaninogenica		99.35	0.50	IND 1	1		
Clostridium perfringens		0.37	0.15	GLU 90	99		
Prevotella intermedia		0.21	0.10		1		
16511122	**Good identification**						**16511122**
					GRA		
Prevotella melaninogenica		97.90	0.37	IND 1	1		
Clostridium perfringens		0.89	0.09	GLU 90	99		
Prevotella oralis		0.82	0.05		1		
16511200	**Good identification**						**16511200**
Prevotella intermedia		100.0	0.44	GLU 90			
16511400	**Good identification**						**16511400**
					GRA		
Prevotella intermedia		99.32	0.44	GLU 90	1		
Lactobacillus fermentum		0.68	0.20	ARA 1	99		
16512000	**Good identification**						**16512000**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		99.28	0.44	GLU 90	1	1	
Propionibacterium granulosum		0.72	0.16	MAN 1	99	90	
16513000	**Good identification**						**16513000**
					GRA	CAT	
Prevotella intermedia		99.28	0.44	GLU 90	1	1	
Propionibacterium granulosum		0.72	0.16	MAN 1	99	90	
16514000	**Good identification**						**16514000**
Prevotella intermedia		100.0	0.44	GLU 90			
16515000	**Good identification**						**16515000**
					GRA		
Prevotella intermedia		99.60	0.44	GLU 90	1		
Lactobacillus fermentum		0.40	0.16	XYL 1	99		
16530000	**Good identification**						**16530000**
					GRA	CAT	
Prevotella intermedia		97.21	0.44	GLU 90	1	1	
Propionibacterium granulosum		2.79	0.26	TRE 1	99	90	
16531000	**Good identification**						**16531000**
					GRA	CAT	
Prevotella intermedia		97.21	0.44	GLU 90	1	1	
Propionibacterium granulosum		2.79	0.26	TRE 1	99	90	
16535722	**Good identification**						**16535722**
Bacteroides thetaiotaomicron		99.60	0.50	ESL 92			
Parabacteroides distasonis		0.23	0.21	GLU 97			
Bacteroides ovatus		0.11	0.00				
16535732	**Good identification**						**16535732**
Bacteroides thetaiotaomicron		98.67	0.43	ESL 92			
Clostridium clostridioforme		0.83	0.18	GLU 97			
Parabacteroides distasonis		0.31	0.17				
16550000	**Good identification**						**16550000**
Prevotella intermedia		100.0	0.44	GLU 90			
16551000	**Good identification**						**16551000**
Prevotella intermedia		100.0	0.44	GLU 90			
16555722	**Low discrimination**						**16555722**
					CAT		
Bacteroides thetaiotaomicron		88.53	0.37	ESL 92	89		
Bacteroides ovatus		5.84	0.17	GLU 97	68		
Bacteroides uniformis		5.63	0.14	TRE 93	5		
16575522	**Good identification**						**16575522**
Bacteroides thetaiotaomicron		96.61	0.37	ESL 92			
Bacteroides ovatus		3.39	0.12	GLU 97			
16575722	**Good identification**						**16575722**
Bacteroides thetaiotaomicron		95.07	0.56	ESL 92			
Bacteroides ovatus		4.77	0.34	GLU 97			
Clostridium clostridioforme		0.11	0.16				
16575732	**Acceptable identification**						**16575732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		90.90	0.49	ESL 92			
Bacteroides ovatus		7.82	0.31	GLU 97			
Clostridium clostridioforme		1.21	0.27				
16710000	**Good identification**						**16710000**
Prevotella intermedia		100.0	0.44	GLU 90			
16711000	**Good identification**						**16711000**
Prevotella intermedia		100.0	0.44	GLU 90			
16735722	**Acceptable identification**						**16735722**
Bacteroides thetaiotaomicron		91.86	0.38	ESL 92			
Bacteroides ovatus		4.11	0.15	GLU 97			
Clostridium clostridioforme		2.75	0.22				
16775712	**Acceptable identification**						**16775712**
Clostridium clostridioforme		92.29	0.42	IND 1			
Bacteroides ovatus		6.03	0.12	GLU 88			
Bacteroides thetaiotaomicron		1.69	0.02				
16775722	**Low discrimination**						**16775722**
Bacteroides ovatus		66.43	0.49	ESL 91			
Bacteroides thetaiotaomicron		31.85	0.43	GLU 99			
Clostridium clostridioforme		1.50	0.31				
16775732	**Low discrimination**						**16775732**
Bacteroides ovatus		69.65	0.46	ESL 91			
Bacteroides thetaiotaomicron		19.48	0.36	GLU 99			
Clostridium clostridioforme		10.78	0.42				
16777722	**Good identification**						**16777722**
Bacteroides ovatus		98.57	0.41	ESL 91			
Bacteroides thetaiotaomicron		1.43	0.09	GLU 99			
16777732	**Good identification**						**16777732**
Bacteroides ovatus		98.70	0.38	ESL 91			
Bacteroides thetaiotaomicron		0.84	0.02	GLU 99			
Clostridium clostridioforme		0.46	0.08				
17000000	**Low discrimination**						**17000000**
					COC	SPO	
Clostridium bifermentans		48.21	0.53	SUC 1	1	99	
Gemella morbillorum		23.89	0.52		99	1	
Clostridium cadaveris		17.58	0.44		1	99	
17000020	**Very good identification**						**17000020**
					COC	SPO	
Clostridium cadaveris		99.63	0.78	SUC 7	1	99	
Clostridium bifermentans		0.18	0.33		1	99	
Gemella morbillorum		0.15	0.35		99	1	
17000021	**Good identification**						**17000021**
					COC	SPO	
Clostridium cadaveris		99.67	0.44	SUC 7	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		0.18	-.01	URE 1	1	99		
Gemella morbillorum		0.15	0.01		99	1		
17000022	**Good identification**							**17000022**
					SPO			
Clostridium cadaveris		98.08	0.44	SUC 7	99			
Atopobium minutum		1.59	0.23	BGL 1	1			
Clostridium bifermentans		0.17	-.01		99			
17000030	**Good identification**							**17000030**
					COC	SPO		
Clostridium cadaveris		99.67	0.44	SUC 7	1	99		
Clostridium bifermentans		0.18	-.01	MLZ 1	1	99		
Gemella morbillorum		0.15	0.01		99	1		
17000060	**Good identification**							**17000060**
Clostridium cadaveris		98.07	0.44	SUC 7				
Clostridium botulinum biovar B		0.82	0.08	SOR 1				
Clostridium bifermentans		0.72	0.09					
17000120	**Good identification**							**17000120**
					GRA	SPO		
Clostridium cadaveris		99.05	0.44	SUC 7	99	99		
Prevotella bivia		0.63	-.03	LAC 1	1	1		
Clostridium bifermentans		0.18	-.01		99	99		
17000220	**Good identification**							**17000220**
					COC	SPO		
Clostridium cadaveris		99.67	0.44	SUC 7	1	99		
Clostridium bifermentans		0.18	-.01	RHA 1	1	99		
Gemella morbillorum		0.15	0.01		99	1		
17000400	**Low discrimination**							**17000400**
					COC	SPO		
Gemella morbillorum		78.62	0.35	IND 1	99	1		
Clostridium bifermentans		14.42	0.19	ARA 10	1	99		
Clostridium cadaveris		5.26	0.09		1	99		
17000420	**Good identification**							**17000420**
					COC	SPO		
Clostridium cadaveris		98.18	0.44	SUC 7	1	99		
Gemella morbillorum		1.65	0.19	ARA 1	99	1		
Clostridium bifermentans		0.17	-.01		1	99		
17001000	**Low discrimination**							**17001000**
					COC	GRA		
Gemella morbillorum		79.60	0.47	IND 1	99	99		
Fusobacterium necrophorum subsp. necrophorum		5.33	0.17		1	1		
Prevotella intermedia		3.83	0.23		1	1		
17001020	**Acceptable identification**							**17001020**
					COC	SPO	CAT	
Clostridium cadaveris		90.12	0.44	SUC 7	1	99	1	
Gemella morbillorum		7.33	0.31	GAL 1	99	1	1	
Propionibacterium acnes		1.02	0.14		1	1	95	
17001100	**Low discrimination**							**17001100**
					COC	GRA		
Eubacterium saburreum		81.92	0.37	ESL 99	1	85		
Gemella morbillorum		9.81	0.13		99	99		
Fusobacterium mortiferum		5.61	0.19		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
17001110	**Good identification**							**17001110**
Eubacterium saburreum		100.0	0.36	ESL 99				
17002020	**Good identification**							**17002020**
					SPO	CAT		
Clostridium cadaveris		98.86	0.44	SUC 7	99	1		
Propionibacterium acnes		0.63	0.09	MAN 1	1	95		
Propionibacterium granulosum		0.18	0.09		1	90		
17004000	**Low discrimination**							**17004000**
					COC	GRA	SPO	
Gemella morbillorum		74.65	0.35	IND 1	99	99	1	
Clostridium bifermentans		13.69	0.19	XYL 10	1	99	99	
Fusobacterium varium		6.67	0.08		1	1	1	
17004020	**Good identification**							**17004020**
					COC	SPO		
Clostridium cadaveris		98.18	0.44	SUC 7	1	99		
Gemella morbillorum		1.65	0.19	XYL 1	99	1		
Clostridium bifermentans		0.17	-.01		1	99		
17010000	**Low discrimination**							**17010000**
					COC	GRA		
Fusobacterium varium		80.92	0.66	SUC 1	1	1		
Prevotella intermedia		8.80	0.58		1	1		
Gemella morbillorum		4.20	0.53		99	99		
17010020	**Very good identification**							**17010020**
					GRA	SPO	CAT	
Clostridium cadaveris		98.08	0.78	SUC 7	99	99	1	
Propionibacterium acnes		1.11	0.48		99	1	95	
Fusobacterium varium		0.32	0.32		1	1	1	
17010021	**Good identification**							**17010021**
					GRA	SPO	CAT	
Clostridium cadaveris		98.15	0.44	SUC 7	99	99	1	
Propionibacterium acnes		1.11	0.14	URE 1	99	1	95	
Fusobacterium varium		0.32	-.03		1	1	1	
17010022	**Acceptable identification**							**17010022**
					SPO	CAT		
Clostridium cadaveris		91.61	0.44	SUC 7	99	1		
Propionibacterium granulosum		5.51	0.35	BGL 1	1	90		
Atopobium minutum		1.37	0.23		1	1		
17010030	**Good identification**							**17010030**
					SPO	CAT		
Clostridium cadaveris		95.30	0.44	SUC 7	99	1		
Propionibacterium granulosum		3.13	0.31	MLZ 1	1	90		
Propionibacterium acnes		1.08	0.14		1	95		
17010060	**Low discrimination**							**17010060**
					SPO	CAT		
Clostridium cadaveris		77.68	0.44	SUC 7	99	1		
Propionibacterium acnes		19.13	0.36	SOR 1	1	95		
Clostridium botulinum biovar B		2.60	0.18		99	1		
17010120	**Good identification**							**17010120**
					SPO	CAT		
Clostridium cadaveris		97.54	0.44	SUC 7	99	1		
Propionibacterium acnes		1.11	0.14	LAC 1	1	95		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		0.34	0.08		1	1		
17010220	**Good identification**							**17010220**
					SPO	CAT		
Clostridium cadaveris		97.67	0.44	SUC 7	99	1		
Propionibacterium acnes		1.11	0.14	RHA 1	1	95		
Propionibacterium granulosum		0.73	0.20		1	90		
17010420	**Good identification**							**17010420**
					COC	SPO	CAT	
Clostridium cadaveris		96.38	0.44	SUC 7	1	99	1	
Gemella morbillorum		1.98	0.20	ARA 1	99	1	1	
Propionibacterium acnes		1.09	0.14		1	1	95	
17011000	**Low discrimination**							**17011000**
					COC	GRA	CAT	
Prevotella intermedia		57.50	0.58	MLT 99	1	1	1	
Propionibacterium granulosum		15.89	0.56		1	99	90	
Gemella morbillorum		14.76	0.48		99	99	1	
17011002	**Low discrimination**							**17011002**
					GRA	CAT		
Propionibacterium granulosum		57.64	0.46	IND 1	99	90		
Lactobacillus fermentum		19.26	0.42		99	1		
Prevotella intermedia		8.43	0.23		1	1		
17011010	**Low discrimination**							**17011010**
					CAT	GRA		
Propionibacterium granulosum		53.46	0.41	IND 1	90	99		
Eubacterium saburreum		26.46	0.37	MLZ 12	1	85		
Prevotella intermedia		14.33	0.23		1	1		
17011020	**Low discrimination**							**17011020**
					SPO	CAT		
Propionibacterium acnes		43.86	0.48	NIT 92	1	95		
Clostridium cadaveris		39.09	0.44	SUC 1	99	1		
Propionibacterium granulosum		9.41	0.46		1	90		
17011022	**Low discrimination**							**17011022**
					CAT			
Propionibacterium granulosum		50.40	0.35	IND 1	90			
Lactobacillus fermentum		22.46	0.34		1			
Propionibacterium acnes		9.49	0.14		95			
17011060	**Good identification**							**17011060**
					SPO	CAT		
Propionibacterium acnes		95.16	0.36	NIT 92	1	95		
Clostridium cadaveris		3.90	0.09	SUC 1	99	1		
Propionibacterium granulosum		0.94	0.11		1	90		
17011100	**Low discrimination**							**17011100**
					GRA			
Fusobacterium mortiferum		50.32	0.53	IND 1	1			
Eubacterium saburreum		22.26	0.45		85			
Lactobacillus fermentum		15.77	0.46		99			
17011102	**Low discrimination**							**17011102**
					GRA			
Lactobacillus fermentum		61.64	0.46	IND 1	99			
Fusobacterium mortiferum		21.86	0.37		1			
Eubacterium saburreum		15.36	0.32		85			
17011110	**Good identification**							**17011110**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium saburreum		97.29	0.44	ESL 99				
Fusobacterium mortiferum		2.72	0.19					
17011122	**Low discrimination**							**17011122**
Lactobacillus fermentum		54.31	0.38	IND 1				
Clostridium perfringens		31.81	0.22					
Eubacterium saburreum		7.17	0.19					
17011402	**Good identification**							**17011402**
					CAT			
Lactobacillus fermentum		97.07	0.42	IND 1	1			
Propionibacterium granulosum		2.93	0.11		90			
17011500	**Low discrimination**							**17011500**
					GRA			
Lactobacillus fermentum		87.05	0.46	IND 1	99			
Actinomyces meyeri		6.32	0.16		99			
Fusobacterium mortiferum		2.81	0.19		1			
17011502	**Good identification**							**17011502**
Lactobacillus fermentum		100.0	0.46	IND 1				
17011522	**Acceptable identification**							**17011522**
					GRA			
Lactobacillus fermentum		91.82	0.38	IND 1	99			
Odoribacter splanchnicus		8.18	-.03		1			
17012000	**Low discrimination**							**17012000**
					GRA	CAT		
Propionibacterium granulosum		53.48	0.54	IND 1	99	90		
Fusobacterium varium		23.84	0.32		1	1		
Eubacterium limosum		12.54	0.29		99	1		
17012002	**Acceptable identification**							**17012002**
					COC	GRA	CAT	
Propionibacterium granulosum		94.38	0.44	IND 1	1	99	90	
Megasphaera elsdenii		3.02	0.12		99	1	1	
Fusobacterium varium		1.70	-.03		1	1	1	
17012010	**Low discrimination**							**17012010**
					GRA	SPO	CAT	
Propionibacterium granulosum		57.86	0.39	IND 1	99	1	90	
Clostridium difficile		39.23	0.24	MLZ 12	99	99	1	
Fusobacterium varium		1.91	-.03		1	1	1	
17012020	**Low discrimination**							**17012020**
					SPO	CAT		
Clostridium cadaveris		54.96	0.44	SUC 7	99	1		
Propionibacterium acnes		34.69	0.43	MAN 1	1	95		
Propionibacterium granulosum		9.98	0.44		1	90		
17013000	**Acceptable identification**							**17013000**
					COC	GRA	CAT	
Propionibacterium granulosum		92.68	0.54	IND 1	1	99	90	
Prevotella intermedia		4.49	0.23		1	1	1	
Gemella morbillorum		1.15	0.14		99	99	1	
17013002	**Good identification**							**17013002**
Propionibacterium granulosum		100.0	0.44	IND 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
17013010	**Good identification**							**17013010**
Propionibacterium granulosum		100.0	0.39	IND 1				
17013020	**Low discrimination**							**17013020**
					SPO	CAT		
Propionibacterium acnes		76.70	0.43	NIT 92	1	95		
Propionibacterium granulosum		22.07	0.44	SUC 1	1	90		
Clostridium cadaveris		1.23	0.09		99	1		
17014000	**Low discrimination**							**17014000**
					COC	GRA		
Fusobacterium varium		83.57	0.42	SUC 1	1	1		
Gemella morbillorum		11.56	0.37	XYL 4	99	99		
Prevotella intermedia		2.20	0.23		1	1		
17014020	**Good identification**							**17014020**
					COC	GRA	SPO	
Clostridium cadaveris		95.29	0.44	SUC 7	1	99	99	
Gemella morbillorum		1.96	0.20	XYL 1	99	99	1	
Fusobacterium varium		1.29	0.08		1	1	1	
17015002	**Low discrimination**							**17015002**
					CAT			
Lactobacillus fermentum		80.45	0.38	IND 1	1			
Eubacterium saburreum		12.85	0.21		1			
Propionibacterium granulosum		4.14	0.11		90			
17015100	**Low discrimination**							**17015100**
Eubacterium saburreum		35.35	0.41	ESL 99				
Lactobacillus fermentum		27.30	0.42					
Actinomyces meyeri		26.19	0.31					
17015102	**Low discrimination**							**17015102**
					GRA			
Lactobacillus fermentum		76.71	0.42	IND 1	99			
Eubacterium saburreum		17.53	0.28		85			
Fusobacterium mortiferum		3.49	0.17		1			
17015110	**Good identification**							**17015110**
Eubacterium saburreum		100.0	0.39	ESL 99				
17015400	**Low discrimination**							**17015400**
					COC			
Lactobacillus fermentum		75.10	0.38	IND 1	1			
Gemella morbillorum		17.56	0.16		99			
Actinomyces meyeri		7.35	0.10		1			
17015402	**Good identification**							**17015402**
Lactobacillus fermentum		100.0	0.38	IND 1				
17015500	**Low discrimination**							**17015500**
Lactobacillus fermentum		80.66	0.42	IND 1				
Actinomyces meyeri		19.34	0.21					
17015502	**Good identification**							**17015502**
Lactobacillus fermentum		100.0	0.42	IND 1				
17020020	**Good identification**							**17020020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO	CAT	
Clostridium cadaveris		99.80	0.56	SUC 7	1	99	1	
Propionibacterium granulosum		0.14	0.20	TRE 5	1	1	90	
Gemella morbillorum		0.03	0.01		99	1	1	
17030000	**Low discrimination**							**17030000**
					GRA	SPO	CAT	
Propionibacterium granulosum		86.07	0.64	IND 1	99	1	90	
Fusobacterium varium		9.65	0.32		1	1	1	
Clostridium cadaveris		1.57	0.22		99	99	1	
17030002	**Good identification**							**17030002**
					GRA	CAT		
Propionibacterium granulosum		99.01	0.54	IND 1	99	90		
Fusobacterium mortiferum		0.55	0.18		1	5		
Fusobacterium varium		0.45	-.03		1	1		
17030010	**Good identification**							**17030010**
					GRA	CAT		
Propionibacterium granulosum		99.18	0.49	IND 1	99	90		
Fusobacterium varium		0.82	-.03	MLZ 12	1	1		
17030012	**Good identification**							**17030012**
Propionibacterium granulosum		100.0	0.39	IND 1				
17030020	**Low discrimination**							**17030020**
					SPO	CAT		
Clostridium cadaveris		87.58	0.56	SUC 7	99	1		
Propionibacterium granulosum		12.14	0.54	TRE 5	1	90		
Propionibacterium acnes		0.19	0.14		1	95		
17030022	**Low discrimination**							**17030022**
					SPO	CAT		
Propionibacterium granulosum		77.43	0.43	IND 1	1	90		
Clostridium cadaveris		22.57	0.22		99	1		
17030030	**Low discrimination**							**17030030**
					SPO	CAT		
Propionibacterium granulosum		65.17	0.39	IND 1	1	90		
Clostridium cadaveris		34.83	0.22	MLZ 12	99	1		
17030100	**Low discrimination**							**17030100**
					GRA	CAT		
Fusobacterium mortiferum		72.00	0.41	IND 1	1	5		
Propionibacterium granulosum		25.18	0.30		99	90		
Fusobacterium varium		2.82	-.03		1	1		
17030200	**Good identification**							**17030200**
					GRA	CAT		
Propionibacterium granulosum		96.47	0.38	IND 1	99	90		
Fusobacterium varium		3.53	-.03	RHA 3	1	1		
17031000	**Good identification**							**17031000**
					CAT	GRA		
Propionibacterium granulosum		94.94	0.64	IND 1	90	99		
Fusobacterium mortiferum		1.76	0.38		5	1		
Eubacterium saburreum		1.74	0.36		1	85		
17031002	**Good identification**							**17031002**
					CAT	GRA		
Propionibacterium granulosum		97.93	0.54	IND 1	90	99		
Eubacterium saburreum		1.27	0.23		1	85		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.81	0.21		5	1	
17031010	**Acceptable identification**						**17031010**
					CAT		
Propionibacterium granulosum		90.09	0.49	IND 1	90		
Eubacterium saburreum		9.91	0.34	MLZ 12	1		
17031012	**Acceptable identification**						**17031012**
					CAT		
Propionibacterium granulosum		92.80	0.39	IND 1	90		
Eubacterium saburreum		7.20	0.22	MLZ 12	1		
17031020	**Low discrimination**						**17031020**
					CAT	SPO	
Propionibacterium granulosum		89.24	0.54	IND 1	90	1	
Clostridium cadaveris		6.50	0.22		1	99	
Clostridium perfringens		1.70	0.17		1	84	
17031022	**Good identification**						**17031022**
					CAT		
Propionibacterium granulosum		95.69	0.43	IND 1	90		
Clostridium perfringens		3.44	0.11		1		
Eubacterium saburreum		0.87	0.10		1		
17031030	**Acceptable identification**						**17031030**
					CAT		
Propionibacterium granulosum		92.80	0.39	IND 1	90		
Eubacterium saburreum		7.20	0.22	MLZ 12	1		
17031100	**Low discrimination**						**17031100**
					CAT	GRA	
Fusobacterium mortiferum		43.37	0.44	IND 1	5	1	
Eubacterium saburreum		42.82	0.42		1	85	
Propionibacterium granulosum		10.11	0.30		90	99	
17031110	**Good identification**						**17031110**
					CAT		
Eubacterium saburreum		96.21	0.41	ESL 99	1		
Propionibacterium granulosum		3.79	0.15		90		
17031200	**Good identification**						**17031200**
Propionibacterium granulosum		100.0	0.38	IND 1			
17032000	**Very good identification**						**17032000**
					GRA	CAT	
Propionibacterium granulosum		99.77	0.62	IND 1	99	90	
Fusobacterium varium		0.15	-.03		1	1	
Eubacterium limosum		0.08	-.06		99	1	
17032002	**Good identification**						**17032002**
Propionibacterium granulosum		100.0	0.52	IND 1			
17032010	**Good identification**						**17032010**
					SPO	CAT	
Propionibacterium granulosum		98.33	0.47	IND 1	1	90	
Clostridium difficile		1.67	0.04	MLZ 12	99	1	
17032012	**Good identification**						**17032012**
Propionibacterium granulosum		100.0	0.37	IND 1			
17032020	**Acceptable identification**						**17032020**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		90.23	0.52	IND 1	1	90	
Clostridium cadaveris		8.72	0.22		99	1	
Propionibacterium acnes		1.06	0.09		1	95	
17032022	**Good identification**						**17032022**
Propionibacterium granulosum		100.0	0.41	IND 1			
17032030	**Good identification**						**17032030**
Propionibacterium granulosum		100.0	0.37	IND 1			
17032200	**Good identification**						**17032200**
Propionibacterium granulosum		100.0	0.36	IND 1			
17033000	**Very good identification**						**17033000**
Propionibacterium granulosum		100.0	0.62	IND 1			
17033002	**Good identification**						**17033002**
Propionibacterium granulosum		100.0	0.52	IND 1			
17033010	**Good identification**						**17033010**
Propionibacterium granulosum		100.0	0.47	IND 1			
17033012	**Good identification**						**17033012**
Propionibacterium granulosum		100.0	0.37	IND 1			
17033020	**Good identification**						**17033020**
Propionibacterium granulosum		98.84	0.52	IND 1			
Propionibacterium acnes		1.16	0.09				
17033022	**Good identification**						**17033022**
Propionibacterium granulosum		100.0	0.41	IND 1			
17033030	**Good identification**						**17033030**
Propionibacterium granulosum		100.0	0.37	IND 1			
17033200	**Good identification**						**17033200**
Propionibacterium granulosum		100.0	0.36	IND 1			
17035100	**Low discrimination**						**17035100**
					COC	GRA	
Eubacterium saburreum		73.98	0.38	ESL 99	1	85	
Atopobium parvulum		15.55	0.16		99	99	
Fusobacterium mortiferum		10.47	0.25		1	1	
17035110	**Good identification**						**17035110**
Eubacterium saburreum		100.0	0.36	ESL 99			
17040020	**Good identification**						**17040020**
					COC	SPO	
Clostridium cadaveris		99.67	0.44	SUC 7	1	99	
Clostridium bifermentans		0.18	-.01	CEL 1	1	99	
Gemella morbillorum		0.15	0.01		99	1	
17050020	**Good identification**						**17050020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Clostridium cadaveris		98.15	0.44	SUC 7	99	99	1	
Propionibacterium acnes		1.11	0.14	CEL 1	99	1	95	
Fusobacterium varium		0.32	-.03		1	1	1	
17050100	**Good identification**							**17050100**
Fusobacterium mortiferum		96.61	0.42	IND 1				
Fusobacterium varium		3.39	-.03					
17051000	**Low discrimination**							**17051000**
					GRA	CAT		
Fusobacterium mortiferum		45.29	0.39	IND 1	1	5		
Prevotella intermedia		26.63	0.23		1	1		
Propionibacterium granulosum		7.36	0.22		99	90		
17051100	**Acceptable identification**							**17051100**
					GRA			
Fusobacterium mortiferum		92.11	0.45	IND 1	1			
Lactobacillus catenaformis		6.66	0.15		99			
Eubacterium saburreum		1.24	0.11		85			
17071100	**Low discrimination**							**17071100**
					COC	GRA		
Fusobacterium mortiferum		70.37	0.36	IND 1	1	1		
Atopobium parvulum		29.63	0.18		99	99		
17100000	**Low discrimination**							**17100000**
					COC	GRA	SPO	
Gemella morbillorum		42.74	0.64	IND 1	99	99	1	
Clostridium bifermentans		35.50	0.60		1	99	99	
Prevotella intermedia		19.33	0.58		1	1	1	
17100001	**Low discrimination**							**17100001**
					COC	SPO		
Clostridium sordellii		86.18	0.42	SUC 1	1	99		
Gemella morbillorum		5.95	0.30	MNS 4	99	1		
Clostridium bifermentans		4.94	0.25		1	99		
17100020	**Low discrimination**							**17100020**
					COC	SPO		
Clostridium cadaveris		79.82	0.56	SUC 7	1	99		
Gemella morbillorum		13.11	0.48	MLT 5	99	1		
Clostridium bifermentans		6.26	0.39		1	99		
17100040	**Low discrimination**							**17100040**
					COC	SPO		
Clostridium botulinum biovar B		80.96	0.47	IND 1	1	99		
Clostridium bifermentans		12.83	0.36		1	99		
Gemella morbillorum		3.74	0.30		99	1		
17100050	**Good identification**							**17100050**
Clostridium botulinum biovar B		99.68	0.42	IND 1				
Clostridium bifermentans		0.32	0.02					
17100060	**Acceptable identification**							**17100060**
Clostridium botulinum biovar B		91.37	0.42	IND 1				
Clostridium cadaveris		5.80	0.22					
Clostridium bifermentans		1.88	0.16					
17100070	**Good identification**							**17100070**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		100.0	0.37	IND 1				
17100400	**Low discrimination**							**17100400**
					COC	GRA	SPO	
Gemella morbillorum		89.27	0.48	IND 1	99	99	1	
Clostridium bifermentans		6.74	0.25	ARA 10	1	99	99	
Prevotella intermedia		3.67	0.23		1	1	1	
17101000	**Low discrimination**							**17101000**
					COC	GRA		
Gemella morbillorum		53.08	0.60	IND 1	99	99		
Prevotella intermedia		44.59	0.58		1	1		
Eubacterium saburreum		1.36	0.41		1	85		
17101010	**Low discrimination**							**17101010**
					COC	GRA		
Eubacterium saburreum		52.92	0.40	ESL 99	1	85		
Gemella morbillorum		25.59	0.26		99	99		
Prevotella intermedia		21.49	0.23		1	1		
17101020	**Low discrimination**							**17101020**
					COC	GRA	SPO	
Gemella morbillorum		79.18	0.43	IND 1	99	99	1	
Clostridium cadaveris		9.04	0.22	NAG 10	1	99	99	
Prevotella intermedia		6.05	0.23		1	1	1	
17101100	**Low discrimination**							**17101100**
					COC	GRA		
Eubacterium saburreum		69.30	0.47	ESL 99	1	85		
Gemella morbillorum		11.75	0.26		99	99		
Prevotella intermedia		9.87	0.23		1	1		
17101101	**Good identification**							**17101101**
					SPO			
Clostridium chauvoei		99.79	0.54	IND 1	99			
Eubacterium saburreum		0.12	0.13		1			
Clostridium sordellii		0.10	-.05		99			
17101103	**Good identification**							**17101103**
Clostridium chauvoei		100.0	0.41	IND 1				
17101110	**Good identification**							**17101110**
Eubacterium saburreum		100.0	0.46	ESL 99				
17101121	**Good identification**							**17101121**
					GRA	SPO		
Clostridium chauvoei		99.55	0.45	IND 1	99	99		
Prevotella melaninogenica		0.45	-.01		1	1		
17101400	**Acceptable identification**							**17101400**
					COC	GRA		
Gemella morbillorum		92.45	0.43	IND 1	99	99		
Prevotella intermedia		7.06	0.23	ARA 10	1	1		
Lactobacillus fermentum		0.49	0.16		1	99		
17104000	**Low discrimination**							**17104000**
					COC	GRA	SPO	
Gemella morbillorum		89.27	0.48	IND 1	99	99	1	
Clostridium bifermentans		6.74	0.25	XYL 10	1	99	99	
Prevotella intermedia		3.67	0.23		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
17105000	**Low discrimination**							**17105000**
					COC	GRA		
Gemella morbillorum		83.11	0.43	IND 1	99	99		
Eubacterium saburreum		10.28	0.37	XYL 10	1	85		
Prevotella intermedia		6.35	0.23		1	1		
17105010	**Acceptable identification**							**17105010**
					COC			
Eubacterium saburreum		90.93	0.35	ESL 99	1			
Gemella morbillorum		9.07	0.09		99			
17105100	**Acceptable identification**							**17105100**
					COC			
Eubacterium saburreum		92.12	0.43	ESL 99	1			
Gemella morbillorum		3.22	0.09		99			
Actinomyces meyeri		3.04	0.10		1			
17105110	**Good identification**							**17105110**
Eubacterium saburreum		100.0	0.41	ESL 99				
17105722	**Good identification**							**17105722**
Bacteroides eggerthii		100.0	0.39	ESL 90				
17110000	**Very good identification**							**17110000**
					COC	GRA	CAT	
Prevotella intermedia		96.90	0.92		1	1	1	
Gemella morbillorum		2.65	0.66		99	99	1	
Propionibacterium granulosum		0.12	0.50		1	99	90	
17110001	**Low discrimination**							**17110001**
					COC	GRA	SPO	
Prevotella intermedia		73.81	0.58	URE 1	1	1	1	
Clostridium sordellii		23.87	0.42		1	99	99	
Gemella morbillorum		2.02	0.32		99	99	1	
17110002	**Good identification**							**17110002**
					COC	GRA	CAT	
Prevotella intermedia		93.80	0.58	BGL 1	1	1	1	
Propionibacterium granulosum		2.91	0.40		1	99	90	
Gemella morbillorum		2.56	0.32		99	99	1	
17110010	**Good identification**							**17110010**
					COC	GRA	CAT	
Prevotella intermedia		94.57	0.58	MLZ 1	1	1	1	
Gemella morbillorum		2.58	0.32		99	99	1	
Propionibacterium granulosum		1.60	0.35		1	99	90	
17110020	**Low discrimination**							**17110020**
					COC	GRA	SPO	
Clostridium cadaveris		52.48	0.56	SUC 7	1	99	99	
Prevotella intermedia		35.09	0.58	MLT 5	1	1	1	
Gemella morbillorum		10.54	0.50		99	99	1	
17110040	**Low discrimination**							**17110040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		64.80	0.58	IND 1	1	99	99	
Prevotella intermedia		33.54	0.58		1	1	1	
Gemella morbillorum		0.92	0.32		99	99	1	
17110050	**Good identification**							**17110050**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		98.95	0.52	IND 1	99	99		
Prevotella intermedia		1.05	0.23		1	1		
17110060	**Good identification**							**17110060**
					GRA	SPO		
Clostridium botulinum biovar B		97.05	0.52	IND 1	99	99		
Clostridium cadaveris		1.54	0.22		99	99		
Prevotella intermedia		1.03	0.23		1	1		
17110070	**Good identification**							**17110070**
Clostridium botulinum biovar B		100.0	0.47	IND 1				
17110100	**Good identification**							**17110100**
					COC	GRA		
Prevotella intermedia		94.91	0.58	LAC 1	1	1		
Gemella morbillorum		2.59	0.32		99	99		
Fusobacterium mortiferum		0.94	0.34		1	1		
17110200	**Very good identification**							**17110200**
					COC	GRA	CAT	
Prevotella intermedia		96.65	0.58	RHA 1	1	1	1	
Gemella morbillorum		2.64	0.32		99	99	1	
Propionibacterium granulosum		0.37	0.24		1	99	90	
17110400	**Low discrimination**							**17110400**
					COC	GRA	CAT	
Prevotella intermedia		76.60	0.58	ARA 1	1	1	1	
Gemella morbillorum		23.00	0.50		99	99	1	
Propionibacterium granulosum		0.10	0.16		1	99	90	
17111000	**Very good identification**							**17111000**
					COC	GRA	CAT	
Prevotella intermedia		98.25	0.92		1	1	1	
Gemella morbillorum		1.44	0.61		99	99	1	
Propionibacterium granulosum		0.12	0.50		1	99	90	
17111001	**Good identification**							**17111001**
					COC	GRA	SPO	
Prevotella intermedia		91.84	0.58	URE 1	1	1	1	
Clostridium sordellii		5.24	0.29		1	99	99	
Gemella morbillorum		1.35	0.27		99	99	1	
17111002	**Low discrimination**							**17111002**
					GRA	CAT		
Prevotella intermedia		87.93	0.58	BGL 1	1	1		
Lactobacillus fermentum		6.09	0.50		99	1		
Propionibacterium granulosum		2.73	0.40		99	90		
17111010	**Good identification**							**17111010**
					CAT	GRA		
Prevotella intermedia		90.28	0.58	MLZ 1	1	1		
Eubacterium saburreum		6.74	0.48		1	85		
Propionibacterium granulosum		1.53	0.35		90	99		
17111020	**Low discrimination**							**17111020**
					COC	GRA		
Prevotella intermedia		68.01	0.58	NAG 1	1	1		
Clostridium perfringens		14.35	0.46		1	99		
Gemella morbillorum		11.00	0.45		99	99		
17111022	**Low discrimination**							**17111022**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		67.56	0.41	IND 1	99			
Lactobacillus fermentum		15.71	0.42	LAC 88	99			
Prevotella intermedia		6.87	0.23		1			
17111040	**Very good identification**							**17111040**
					COC	GRA	SPO	
Prevotella intermedia		96.31	0.58	SOR 1	1	1	1	
Clostridium botulinum biovar B		1.88	0.24		1	99	99	
Gemella morbillorum		1.42	0.27		99	99	1	
17111100	**Low discrimination**							**17111100**
					GRA			
Prevotella intermedia		69.34	0.58	LAC 1	1			
Eubacterium saburreum		14.75	0.56		85			
Lactobacillus fermentum		7.84	0.54		99			
17111101	**Very good identification**							**17111101**
					SPO	GRA		
Clostridium chauvoei		98.94	0.60	IND 1	99	99		
Prevotella intermedia		0.71	0.23		1	1		
Eubacterium saburreum		0.15	0.21		1	85		
17111102	**Low discrimination**							**17111102**
Lactobacillus fermentum		58.85	0.54	IND 1				
Eubacterium saburreum		19.55	0.43					
Clostridium perfringens		11.73	0.30					
17111103	**Good identification**							**17111103**
					SPO			
Clostridium chauvoei		99.07	0.47	IND 1	99			
Lactobacillus fermentum		0.45	0.20		1			
Actinomyces naeslundii		0.28	0.26		1			
17111110	**Acceptable identification**							**17111110**
					GRA			
Eubacterium saburreum		93.69	0.54	ESL 99	85			
Prevotella intermedia		5.44	0.23		1			
Lactobacillus fermentum		0.61	0.20		99			
17111112	**Good identification**							**17111112**
Eubacterium saburreum		96.42	0.41	ESL 99				
Lactobacillus fermentum		3.58	0.20					
17111120	**Low discrimination**							**17111120**
					GRA			
Clostridium perfringens		63.08	0.61	IND 1	99			
Prevotella melaninogenica		32.90	0.50		1			
Lactobacillus fermentum		1.54	0.46		99			
17111121	**Good identification**							**17111121**
					GRA	SPO		
Clostridium chauvoei		95.05	0.52	IND 1	99	99		
Clostridium perfringens		3.16	0.27		99	84		
Prevotella melaninogenica		1.65	0.15		1	1		
17111122	**Low discrimination**							**17111122**
					GRA			
Clostridium perfringens		77.32	0.56	IND 1	99			
Prevotella melaninogenica		16.32	0.37		1			
Lactobacillus fermentum		4.00	0.46		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
17111123	**Low discrimination**							**17111123**
					GRA	SPO		
Clostridium chauvoei		89.63	0.39	IND 1	99	99		
Clostridium perfringens		7.94	0.22		99	84		
Prevotella melaninogenica		1.68	0.03		1	1		
17111130	**Low discrimination**							**17111130**
					GRA			
Eubacterium saburreum		56.37	0.41	ESL 99	85			
Clostridium perfringens		28.68	0.27		99			
Prevotella melaninogenica		14.95	0.15		1			
17111160	**Good identification**							**17111160**
					GRA			
Clostridium perfringens		97.31	0.49	IND 1	99			
Prevotella melaninogenica		2.69	0.15		1			
17111162	**Good identification**							**17111162**
					GRA			
Clostridium perfringens		98.89	0.43	IND 1	99			
Prevotella melaninogenica		1.11	0.03		1			
17111200	**Very good identification**							**17111200**
					COC	GRA	CAT	
Prevotella intermedia		97.98	0.58	RHA 1	1	1	1	
Gemella morbillorum		1.44	0.27		99	99	1	
Propionibacterium granulosum		0.38	0.24		1	99	90	
17111400	**Low discrimination**							**17111400**
					COC	GRA		
Prevotella intermedia		81.05	0.58	ARA 1	1	1		
Gemella morbillorum		13.10	0.45		99	99		
Lactobacillus fermentum		5.61	0.50		1	99		
17111402	**Low discrimination**							**17111402**
					COC	GRA		
Lactobacillus fermentum		85.51	0.50	IND 1	1	99		
Prevotella intermedia		12.47	0.23		1	1		
Gemella morbillorum		2.02	0.11		99	99		
17111420	**Low discrimination**							**17111420**
					COC	GRA		
Lactobacillus fermentum		43.33	0.42	IND 1	1	99		
Gemella morbillorum		33.71	0.29		99	99		
Prevotella intermedia		18.96	0.23		1	1		
17111422	**Good identification**							**17111422**
Lactobacillus fermentum		95.84	0.42	IND 1				
Clostridium perfringens		4.16	0.07					
17111500	**Low discrimination**							**17111500**
					GRA			
Lactobacillus fermentum		85.21	0.54	IND 1	99			
Prevotella intermedia		7.62	0.23		1			
Actinomyces meyeri		2.32	0.16		99			
17111502	**Good identification**							**17111502**
Lactobacillus fermentum		98.12	0.54	IND 1				
Bifidobacterium longum subsp. longum		1.88	0.18					
17111520	**Low discrimination**							**17111520**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		52.58	0.46	IND 1				
Clostridium perfringens		21.81	0.27					
Bifidobacterium longum subsp. longum		12.08	0.28					
17111522	**Low discrimination**							**17111522**
Lactobacillus fermentum		67.50	0.46	IND 1				
Bifidobacterium longum subsp. longum		15.51	0.28					
Clostridium perfringens		13.18	0.22					
17112000	**Low discrimination**							**17112000**
					COC	GRA	CAT	
Prevotella intermedia		87.17	0.58	MAN 1	1	1	1	
Propionibacterium granulosum		8.16	0.48		1	99	90	
Gemella morbillorum		2.38	0.32		99	99	1	
17112002	**Low discrimination**							**17112002**
					COC	GRA	CAT	
Propionibacterium granulosum		63.91	0.38	IND 1	1	99	90	
Prevotella intermedia		27.57	0.23		1	1	1	
Megasphaera elsdenii		6.83	0.15		99	1	1	
17112020	**Low discrimination**							**17112020**
					GRA	SPO	CAT	
Propionibacterium granulosum		43.78	0.38	IND 1	99	1	90	
Clostridium cadaveris		28.24	0.22		99	99	1	
Prevotella intermedia		18.89	0.23		1	1	1	
17112100	**Low discrimination**							**17112100**
					GRA	CAT		
Propionibacterium propionicum		78.77	0.46	IND 1	99	1		
Prevotella intermedia		19.41	0.23	NIT 86	1	1		
Propionibacterium granulosum		1.82	0.14		99	90		
17112102	**Good identification**							**17112102**
Propionibacterium propionicum		100.0	0.49	IND 1				
17112140	**Good identification**							**17112140**
Propionibacterium propionicum		100.0	0.46	IND 1				
17112142	**Good identification**							**17112142**
Propionibacterium propionicum		100.0	0.49	IND 1				
17113000	**Good identification**							**17113000**
					COC	GRA	CAT	
Prevotella intermedia		90.06	0.58	MAN 1	1	1	1	
Propionibacterium granulosum		8.44	0.48		1	99	90	
Gemella morbillorum		1.32	0.27		99	99	1	
17113002	**Low discrimination**							**17113002**
					GRA	CAT		
Propionibacterium granulosum		67.21	0.38	IND 1	99	90		
Prevotella intermedia		29.00	0.23		1	1		
Lactobacillus fermentum		2.01	0.16		99	1		
17113020	**Low discrimination**							**17113020**
					GRA	CAT		
Propionibacterium granulosum		59.87	0.38	IND 1	99	90		
Prevotella intermedia		25.83	0.23		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		5.45	0.12		99	1	
17113100	**Low discrimination**						**17113100**
					GRA		
Propionibacterium propionicum		72.27	0.46	IND 1	99		
Prevotella intermedia		17.81	0.23	NIT 86	1		
Eubacterium saburreum		3.79	0.21		85		
17113102	**Good identification**						**17113102**
Propionibacterium propionicum		96.77	0.49	IND 1			
Lactobacillus fermentum		1.80	0.20	NIT 86			
Actinomyces naeslundii		1.43	0.28				
17113140	**Good identification**						**17113140**
Propionibacterium propionicum		100.0	0.46	IND 1			
17113142	**Good identification**						**17113142**
Propionibacterium propionicum		100.0	0.49	IND 1			
17114000	**Low discrimination**						**17114000**
					COC	GRA	
Prevotella intermedia		76.21	0.58	XYL 1	1	1	
Gemella morbillorum		22.88	0.50		99	99	
Actinomyces meyeri		0.30	0.18		1	99	
17115000	**Low discrimination**						**17115000**
					COC	GRA	
Prevotella intermedia		79.42	0.58	XYL 1	1	1	
Gemella morbillorum		12.84	0.45		99	99	
Eubacterium saburreum		3.90	0.45		1	85	
17115002	**Low discrimination**						**17115002**
					GRA		
Lactobacillus fermentum		66.60	0.46	IND 1	99		
Prevotella intermedia		16.54	0.23		1		
Eubacterium saburreum		14.19	0.32		85		
17115010	**Low discrimination**						**17115010**
					COC	GRA	
Eubacterium saburreum		76.79	0.43	ESL 99	1	85	
Prevotella intermedia		19.31	0.23		1	1	
Gemella morbillorum		3.12	0.11		99	99	
17115022	**Low discrimination**						**17115022**
Lactobacillus fermentum		84.27	0.38	IND 1			
Eubacterium saburreum		9.50	0.19				
Clostridium perfringens		6.23	0.07				
17115100	**Low discrimination**						**17115100**
Eubacterium saburreum		51.56	0.51	ESL 99			
Lactobacillus fermentum		29.87	0.50				
Actinomyces meyeri		10.77	0.32				
17115102	**Low discrimination**						**17115102**
Lactobacillus fermentum		75.39	0.50	IND 1			
Eubacterium saburreum		22.97	0.38				
Actinomyces meyeri		0.84	0.06				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17115110	**Good identification**						**17115110**
Eubacterium saburreum		99.29	0.49	ESL 99			
Lactobacillus fermentum		0.71	0.16				
17115112	**Good identification**						**17115112**
Eubacterium saburreum		96.11	0.37	ESL 99			
Lactobacillus fermentum		3.90	0.16				
17115120	**Low discrimination**						**17115120**
Lactobacillus fermentum		31.09	0.42	IND 1			
Eubacterium saburreum		28.41	0.38				
Clostridium perfringens		21.96	0.27				
17115122	**Low discrimination**						**17115122**
Lactobacillus fermentum		57.74	0.42	IND 1			
Clostridium perfringens		19.19	0.22				
Eubacterium saburreum		9.31	0.25				
17115130	**Good identification**						**17115130**
Eubacterium saburreum		97.37	0.37	ESL 99			
Bifidobacterium longum subsp. longum		2.63	0.06				
17115400	**Low discrimination**						**17115400**
					COC	GRA	
Lactobacillus fermentum		57.51	0.46	IND 1	1	99	
Gemella morbillorum		25.40	0.29		99	99	
Prevotella intermedia		14.28	0.23		1	1	
17115402	**Good identification**						**17115402**
Lactobacillus fermentum		100.0	0.46	IND 1			
17115420	**Low discrimination**						**17115420**
					COC		
Lactobacillus fermentum		87.17	0.38	IND 1	1		
Gemella morbillorum		12.83	0.12		99		
17115422	**Good identification**						**17115422**
Lactobacillus fermentum		100.0	0.38	IND 1			
17115500	**Low discrimination**						**17115500**
Lactobacillus fermentum		82.96	0.50	IND 1			
Bifidobacterium longum subsp. longum		8.11	0.26				
Actinomyces meyeri		7.48	0.21				
17115502	**Low discrimination**						**17115502**
Lactobacillus fermentum		89.02	0.50	IND 1			
Bifidobacterium longum subsp. longum		8.71	0.26				
Bifidobacterium adolescentis		2.28	0.17				
17115520	**Low discrimination**						**17115520**
					GRA		
Bifidobacterium longum subsp. longum		41.08	0.36	IND 1	99		
Lactobacillus fermentum		35.00	0.42	RAF 97	99		
Bacteroides vulgatus		22.63	0.17		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17115522	**Low discrimination**						**17115522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		52.21	0.36	IND 1	99	1	
Lactobacillus fermentum		44.49	0.42	RAF 97	99	1	
Bacteroides thetaiotaomicron		3.30	0.04		1	89	
17115530	**Good identification**						**17115530**
Bifidobacterium longum subsp. longum		100.0	0.40	IND 1			
17115532	**Good identification**						**17115532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.58	0.40	IND 1	99	1	
Bacteroides thetaiotaomicron		1.42	-.03	RAF 97	1	89	
17120040	**Good identification**						**17120040**
Clostridium botulinum biovar B		99.90	0.56	IND 1			
Clostridium bifermentans		0.05	0.02				
Clostridium botulinum biovar A		0.05	0.02				
17120050	**Good identification**						**17120050**
Clostridium botulinum biovar B		100.0	0.50	IND 1			
17120060	**Good identification**						**17120060**
Clostridium botulinum biovar B		99.89	0.50	IND 1			
Clostridium cadaveris		0.11	0.00				
17120070	**Good identification**						**17120070**
Clostridium botulinum biovar B		100.0	0.45	IND 1			
17121000	**Low discrimination**						**17121000**
					COC	GRA	
Eubacterium saburreum		43.91	0.38	ESL 99	1	85	
Gemella morbillorum		26.06	0.26		99	99	
Prevotella intermedia		21.89	0.23		1	1	
17121010	**Good identification**						**17121010**
					CAT		
Eubacterium saburreum		97.00	0.37	ESL 99	1		
Propionibacterium granulosum		3.00	0.09		90		
17121100	**Acceptable identification**						**17121100**
					COC		
Eubacterium saburreum		92.67	0.44	ESL 99	1		
Atopobium parvulum		4.24	0.12		99		
Clostridium perfringens		1.76	0.06		1		
17121101	**Good identification**						**17121101**
					SPO		
Clostridium chauvoei		99.29	0.37	IND 1	99		
Eubacterium saburreum		0.71	0.10	TRE 10	1		
17121110	**Good identification**						**17121110**
Eubacterium saburreum		100.0	0.43	ESL 99			
17125100	**Low discrimination**						**17125100**
					COC		
Eubacterium saburreum		86.38	0.40	ESL 99	1		
Atopobium parvulum		13.62	0.16		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
17125110	**Good identification**							**17125110**
Eubacterium saburreum		100.0	0.38	ESL 99				
17130000	**Low discrimination**							**17130000**
					GRA	SPO	CAT	
Prevotella intermedia		68.32	0.58	TRE 1	1	1	1	
Propionibacterium granulosum		25.45	0.58		99	1	90	
Clostridium botulinum biovar B		4.00	0.32		99	99	1	
17130002	**Low discrimination**							**17130002**
					GRA	SPO	CAT	
Propionibacterium granulosum		89.70	0.48	IND 1	99	1	90	
Prevotella intermedia		9.73	0.23		1	1	1	
Clostridium botulinum biovar B		0.57	-.03		99	99	1	
17130010	**Low discrimination**							**17130010**
					GRA	SPO	CAT	
Propionibacterium granulosum		56.29	0.43	IND 1	99	1	90	
Clostridium botulinum biovar B		31.96	0.26	MLZ 12	99	99	1	
Prevotella intermedia		11.20	0.23		1	1	1	
17130020	**Low discrimination**							**17130020**
					SPO	CAT		
Propionibacterium granulosum		42.76	0.48	IND 1	1	90		
Clostridium cadaveris		36.15	0.34		99	1		
Clostridium botulinum biovar B		13.24	0.26		99	1		
17130022	**Low discrimination**							**17130022**
					CAT	SPO		
Propionibacterium granulosum		89.75	0.38	IND 1	90	1		
Clostridium perfringens		7.18	0.11		1	84		
Clostridium cadaveris		3.07	0.00		1	99		
17130040	**Very good identification**							**17130040**
					GRA	SPO	CAT	
Clostridium botulinum biovar B		99.76	0.66	IND 1	99	99	1	
Prevotella intermedia		0.17	0.23		1	1	1	
Propionibacterium granulosum		0.07	0.24		99	1	90	
17130050	**Very good identification**							**17130050**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.61	IND 1	99	1		
Propionibacterium granulosum		0.02	0.09		1	90		
17130060	**Very good identification**							**17130060**
					CAT	SPO		
Clostridium botulinum biovar B		99.91	0.61	IND 1	1	99		
Propionibacterium granulosum		0.03	0.14		90	1		
Clostridium perfringens		0.03	0.05		1	84		
17130070	**Good identification**							**17130070**
Clostridium botulinum biovar B		100.0	0.55	IND 1				
17131000	**Low discrimination**							**17131000**
					CAT	GRA		
Prevotella intermedia		67.91	0.58	TRE 1	1	1		
Propionibacterium granulosum		25.29	0.58		90	99		
Eubacterium saburreum		4.13	0.46		1	85		
17131002	**Low discrimination**							**17131002**
					CAT	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		74.41	0.48	IND 1	90	99	
Eubacterium saburreum		8.57	0.33		1	85	
Prevotella intermedia		8.07	0.23		1	1	
17131010	**Low discrimination**						**17131010**
					CAT	GRA	
Propionibacterium granulosum		44.94	0.43	IND 1	90	99	
Eubacterium saburreum		44.01	0.45	MLZ 12	1	85	
Prevotella intermedia		8.94	0.23		1	1	
17131020	**Low discrimination**						**17131020**
					CAT		
Clostridium perfringens		76.83	0.51	IND 1	1		
Propionibacterium granulosum		18.26	0.48	LAC 88	90		
Eubacterium saburreum		2.10	0.33		1		
17131022	**Low discrimination**						**17131022**
					CAT		
Clostridium perfringens		87.49	0.45	IND 1	1		
Propionibacterium granulosum		11.05	0.38	LAC 88	90		
Eubacterium saburreum		0.90	0.20		1		
17131060	**Low discrimination**						**17131060**
					CAT	SPO	
Clostridium perfringens		71.48	0.39	IND 1	1	84	
Clostridium botulinum biovar B		27.62	0.26	LAC 88	1	99	
Propionibacterium granulosum		0.90	0.14		90	1	
17131100	**Low discrimination**						**17131100**
					COC		
Atopobium parvulum		45.85	0.46	IND 1	99		
Eubacterium saburreum		30.40	0.53		1		
Clostridium perfringens		19.05	0.40		1		
17131101	**Good identification**						**17131101**
					COC	SPO	
Clostridium chauvoei		96.40	0.44	IND 1	1	99	
Atopobium parvulum		1.33	0.12	TRE 10	99	1	
Eubacterium saburreum		0.88	0.18		1	1	
17131110	**Good identification**						**17131110**
					COC		
Eubacterium saburreum		97.01	0.51	ESL 99	1		
Atopobium parvulum		1.81	0.12		99		
Clostridium perfringens		0.75	0.06		1		
17131112	**Good identification**						**17131112**
Eubacterium saburreum		97.98	0.38	ESL 99			
Clostridium perfringens		2.02	0.00				
17131120	**Very good identification**						**17131120**
					COC		
Clostridium perfringens		97.43	0.66	IND 1	1		
Atopobium parvulum		1.28	0.33		99		
Eubacterium saburreum		0.85	0.40		1		
17131121	**Low discrimination**						**17131121**
					SPO		
Clostridium chauvoei		63.24	0.35	IND 1	99		
Clostridium perfringens		35.12	0.32	TRE 10	84		
Actinomyces naeslundii		1.65	0.31		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
17131122	**Very good identification**							**17131122**
					COC			
Clostridium perfringens		98.96	0.60	IND 1	1			
Atopobium parvulum		0.49	0.20		99			
Eubacterium saburreum		0.32	0.27		1			
17131160	**Good identification**							**17131160**
Clostridium perfringens		100.0	0.54	IND 1				
17131162	**Good identification**							**17131162**
Clostridium perfringens		100.0	0.48	IND 1				
17132000	**Good identification**							**17132000**
					GRA	SPO	CAT	
Propionibacterium granulosum		96.33	0.56	IND 1	99	1	90	
Prevotella intermedia		3.46	0.23		1	1	1	
Clostridium botulinum biovar B		0.20	-.03		99	99	1	
17132002	**Good identification**							**17132002**
					CAT			
Propionibacterium granulosum		99.28	0.46	IND 1	90			
Propionibacterium propionicum		0.72	0.14		1			
17132010	**Good identification**							**17132010**
Propionibacterium granulosum		100.0	0.41	IND 1				
17132020	**Good identification**							**17132020**
					SPO	CAT		
Propionibacterium granulosum		98.88	0.46	IND 1	1	90		
Clostridium cadaveris		1.12	0.00		99	1		
17132022	**Good identification**							**17132022**
Propionibacterium granulosum		100.0	0.35	IND 1				
17132100	**Acceptable identification**							**17132100**
					CAT			
Propionibacterium propionicum		92.20	0.45	IND 1	1			
Propionibacterium granulosum		7.80	0.22	NIT 86	90			
17132102	**Good identification**							**17132102**
					CAT			
Propionibacterium propionicum		98.61	0.48	IND 1	1			
Propionibacterium granulosum		1.39	0.12	NIT 86	90			
17132140	**Good identification**							**17132140**
					SPO			
Propionibacterium propionicum		98.27	0.45	IND 1	1			
Clostridium botulinum biovar B		1.73	-.03	NIT 86	99			
17132142	**Good identification**							**17132142**
Propionibacterium propionicum		100.0	0.48	IND 1				
17133000	**Good identification**							**17133000**
					GRA	CAT		
Propionibacterium granulosum		96.12	0.56	IND 1	99	90		
Prevotella intermedia		3.46	0.23		1	1		
Actinomyces naeslundii		0.21	0.25		99	25		
17133002	**Good identification**							**17133002**
					CAT			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		98.42	0.46	IND 1	90		
Actinomyces naeslundii		0.87	0.25		25		
Propionibacterium propionicum		0.71	0.14		1		
17133010	**Good identification**						**17133010**
					CAT		
Propionibacterium granulosum		98.71	0.41	IND 1	90		
Eubacterium saburreum		1.29	0.11	MLZ 12	1		
17133020	**Acceptable identification**						**17133020**
					CAT		
Propionibacterium granulosum		93.88	0.46	IND 1	90		
Clostridium perfringens		5.29	0.17		1		
Actinomyces naeslundii		0.83	0.25		25		
17133022	**Low discrimination**						**17133022**
					CAT		
Propionibacterium granulosum		87.62	0.35	IND 1	90		
Clostridium perfringens		9.29	0.11		1		
Actinomyces naeslundii		3.09	0.25		25		
17133100	**Low discrimination**						**17133100**
					COC	CAT	
Propionibacterium propionicum		78.67	0.45	IND 1	1	1	
Propionibacterium granulosum		6.66	0.22	NIT 86	1	90	
Atopobium parvulum		5.07	0.12		99	1	
17133102	**Acceptable identification**						**17133102**
					CAT		
Propionibacterium propionicum		94.56	0.48	IND 1	1		
Actinomyces naeslundii		3.32	0.33	NIT 86	25		
Propionibacterium granulosum		1.33	0.12		90		
17133121	**Low discrimination**						**17133121**
					SPO		
Actinomyces naeslundii		83.37	0.36	IND 1	1		
Clostridium chauvoei		16.63	0.01	RAF 92	99		
17133123	**Good identification**						**17133123**
Actinomyces naeslundii		100.0	0.36	IND 1			
17133140	**Good identification**						**17133140**
Propionibacterium propionicum		100.0	0.45	IND 1			
17133142	**Good identification**						**17133142**
Propionibacterium propionicum		100.0	0.48	IND 1			
17135000	**Low discrimination**						**17135000**
					COC	GRA	
Eubacterium saburreum		62.17	0.42	ESL 99	1	85	
Prevotella intermedia		19.18	0.23		1	1	
Atopobium parvulum		7.62	0.16		99	99	
17135010	**Good identification**						**17135010**
					CAT		
Eubacterium saburreum		98.12	0.40	ESL 99	1		
Propionibacterium granulosum		1.88	0.09		90		
17135100	**Low discrimination**						**17135100**
					COC		
Atopobium parvulum		83.60	0.50	IND 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		16.07	0.48		1		
Clostridium perfringens		0.19	0.06		1		
17135102	**Low discrimination**						**17135102**
					COC		
Atopobium parvulum		82.79	0.38	IND 1	99		
Eubacterium saburreum		15.92	0.35		1		
Lactobacillus fermentum		0.79	0.16		1		
17135110	**Acceptable identification**						**17135110**
					COC		
Eubacterium saburreum		93.97	0.46	ESL 99	1		
Atopobium parvulum		6.03	0.16		99		
17135120	**Low discrimination**						**17135120**
					COC		
Atopobium parvulum		62.26	0.38	IND 1	99		
Clostridium perfringens		25.77	0.32		1		
Eubacterium saburreum		11.97	0.35		1		
17135722	**Good identification**						**17135722**
Bacteroides thetaiotaomicron		98.99	0.42	ESL 92			
Parabacteroides distasonis		0.69	0.21	RAF 99			
Bacteroides ovatus		0.33	0.00				
17135732	**Good identification**						**17135732**
Bacteroides thetaiotaomicron		97.34	0.35	ESL 92			
Clostridium clostridioforme		1.18	0.13	RAF 99			
Parabacteroides distasonis		0.94	0.17				
17145722	**Good identification**						**17145722**
Bacteroides eggerthii		99.97	0.50	ESL 90			
Bacteroides thetaiotaomicron		0.03	-.06	SUC 1			
17150000	**Very good identification**						**17150000**
					COC	GRA	
Prevotella intermedia		96.79	0.58	CEL 1	1	1	
Gemella morbillorum		2.64	0.32		99	99	
Fusobacterium mortiferum		0.14	0.20		1	1	
17151000	**Good identification**						**17151000**
					GRA		
Prevotella intermedia		92.37	0.58	CEL 1	1		
Lactobacillus jensenii		4.20	0.36		99		
Lactobacillus catenaformis		1.61	0.29		99		
17171000	**Low discrimination**						**17171000**
					COC	GRA	
Lactobacillus jensenii		86.62	0.44	IND 1	1	99	
Prevotella intermedia		6.41	0.23	SAL 95	1	1	
Atopobium parvulum		3.20	0.18		99	99	
17171002	**Acceptable identification**						**17171002**
					CAT		
Lactobacillus jensenii		94.18	0.41	IND 1	1		
Lactobacillus acidophilus		3.48	0.14	SAL 95	1		
Propionibacterium granulosum		0.97	0.14		90		
17171010	**Good identification**						**17171010**
					CAT		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		97.82	0.36	IND 1	1		
Propionibacterium granulosum		1.10	0.09	SAL 95	90		
Eubacterium saburreum		1.08	0.11		1		
17171100	**Good identification**						**17171100**
					COC	GRA	
Atopobium parvulum		98.19	0.52	IND 1	99	99	
Clostridium perfringens		0.73	0.16		1	99	
Fusobacterium mortiferum		0.29	0.20		1	1	
17171102	**Low discrimination**						**17171102**
					COC		
Atopobium parvulum		67.42	0.39	IND 1	99		
Lactobacillus acidophilus		29.62	0.32		1		
Clostridium perfringens		1.33	0.11		1		
17171120	**Low discrimination**						**17171120**
					COC	SPO	
Clostridium perfringens		55.02	0.42	IND 1	1	84	
Atopobium parvulum		40.48	0.39	CEL 4	99	1	
Clostridium septicum		3.53	0.15		1	99	
17171122	**Low discrimination**						**17171122**
					COC		
Clostridium perfringens		48.56	0.37	IND 1	1		
Lactobacillus acidophilus		33.37	0.32	CEL 4	1		
Atopobium parvulum		13.40	0.26		99		
17173101	**Good identification**						**17173101**
Actinomyces naeslundii		100.0	0.38	IND 1			
17173102	**Low discrimination**						**17173102**
					SPO		
Actinomyces naeslundii		29.74	0.35	IND 1	1		
Lactobacillus acidophilus		25.91	0.10	RAF 92	1		
Clostridium tertium		19.79	0.09		99		
17173103	**Good identification**						**17173103**
Actinomyces naeslundii		100.0	0.38	IND 1			
17173120	**Low discrimination**						**17173120**
					SPO		
Actinomyces naeslundii		31.34	0.35	IND 1	1		
Clostridium tertium		20.86	0.09	RAF 92	99		
Clostridium ramosum		19.66	0.07		92		
17173121	**Good identification**						**17173121**
Actinomyces naeslundii		100.0	0.38	IND 1			
17173123	**Good identification**						**17173123**
Actinomyces naeslundii		100.0	0.38	IND 1			
17175100	**Very good identification**						**17175100**
					COC		
Atopobium parvulum		99.84	0.57	IND 1	99		
Eubacterium saburreum		0.08	0.14		1		
Actinomyces israelii		0.08	0.17		1		
17175102	**Good identification**						**17175102**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		99.34	0.44	IND 1	99			
Actinomyces israelii		0.66	0.20		1			
17175120	**Good identification**							**17175120**
					COC			
Atopobium parvulum		99.27	0.44	IND 1	99			
Clostridium perfringens		0.73	0.08		1			
17175722	**Low discrimination**							**17175722**
Bacteroides thetaiotaomicron		86.43	0.48	ESL 92				
Bacteroides ovatus		13.28	0.34	RAF 99				
Parabacteroides distasonis		0.15	0.17					
17175732	**Low discrimination**							**17175732**
Bacteroides thetaiotaomicron		77.83	0.41	ESL 92				
Bacteroides ovatus		20.50	0.31	RAF 99				
Clostridium clostridioforme		1.49	0.22					
17200000	**Acceptable identification**							**17200000**
					COC	SPO		
Clostridium bifermentans		91.19	0.37	SUC 1	1	99		
Gemella morbillorum		4.11	0.17	SAL 10	99	1		
Clostridium cadaveris		3.02	0.09		1	99		
17200020	**Good identification**							**17200020**
					COC	SPO		
Clostridium cadaveris		97.95	0.44	SUC 7	1	99		
Clostridium bifermentans		1.91	0.17	SAL 1	1	99		
Gemella morbillorum		0.15	0.01		99	1		
17210020	**Good identification**							**17210020**
					GRA	SPO	CAT	
Clostridium cadaveris		97.92	0.44	SUC 7	99	99	1	
Propionibacterium acnes		1.11	0.14	SAL 1	99	1	95	
Fusobacterium varium		0.32	-.03		1	1	1	
17210100	**Good identification**							**17210100**
Fusobacterium mortiferum		99.40	0.55	IND 1				
Fusobacterium varium		0.60	-.03					
17210102	**Good identification**							**17210102**
Fusobacterium mortiferum		100.0	0.39	IND 1				
17211000	**Low discrimination**							**17211000**
					GRA	CAT		
Fusobacterium mortiferum		82.82	0.52	IND 1	1	5		
Prevotella intermedia		8.36	0.23		1	1		
Propionibacterium granulosum		2.31	0.22		99	90		
17211002	**Low discrimination**							**17211002**
					GRA			
Fusobacterium mortiferum		72.45	0.35	IND 1	1			
Lactobacillus catenaformis		12.60	0.12	BGL 10	99			
Lactobacillus jensenii		10.41	0.11		99			
17211100	**Very good identification**							**17211100**
					GRA			
Fusobacterium mortiferum		98.55	0.58	IND 1	1			
Lactobacillus catenaformis		1.22	0.15		99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		0.23	0.11		85		
17211102	**Low discrimination**						**17211102**
					GRA		
Fusobacterium mortiferum		86.14	0.42	IND 1	1		
Lactobacillus catenaformis		9.63	0.15	BGL 10	99		
Lactobacillus acidophilus		2.45	0.02		99		
17214100	**Good identification**						**17214100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
17215100	**Good identification**						**17215100**
Fusobacterium mortiferum		100.0	0.39	IND 1			
17230000	**Low discrimination**						**17230000**
					GRA	CAT	
Fusobacterium mortiferum		68.14	0.40	IND 1	1	5	
Propionibacterium granulosum		28.65	0.30		99	90	
Fusobacterium varium		3.21	-.03		1	1	
17230100	**Good identification**						**17230100**
Fusobacterium mortiferum		100.0	0.46	IND 1			
17231000	**Low discrimination**						**17231000**
					GRA	CAT	
Fusobacterium mortiferum		65.76	0.43	IND 1	1	5	
Propionibacterium granulosum		18.43	0.30		99	90	
Lactobacillus jensenii		15.81	0.22		99	1	
17231100	**Good identification**						**17231100**
					COC	GRA	
Fusobacterium mortiferum		96.05	0.49	IND 1	1	1	
Atopobium parvulum		3.95	0.14		99	99	
17243000	**Good identification**						**17243000**
Clostridium innocuum		100.0	0.44	IND 1			
17243002	**Good identification**						**17243002**
Clostridium innocuum		100.0	0.44	IND 1			
17250000	**Low discrimination**						**17250000**
					GRA	SPO	
Fusobacterium mortiferum		80.81	0.40	IND 1	1	1	
Clostridium innocuum		7.08	0.10		99	99	
Lactobacillus catenaformis		7.02	0.12		99	1	
17250100	**Acceptable identification**						**17250100**
					GRA	SPO	
Fusobacterium mortiferum		93.12	0.47	IND 1	1	1	
Lactobacillus catenaformis		5.20	0.15		99	1	
Clostridium barati		1.68	0.02		99	99	
17251000	**Low discrimination**						**17251000**
					SPO		
Lactobacillus catenaformis		61.10	0.47	IND 1	1		
Lactobacillus jensenii		14.53	0.36	MLT 90	1		
Clostridium innocuum		11.83	0.32		99		
17251002	**Low discrimination**						**17251002**
					SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		66.79	0.47	IND 1	1		
Clostridium innocuum		12.93	0.32	MLT 90	99		
Lactobacillus jensenii		10.59	0.33		1		
17251020	**Low discrimination**						**17251020**
					SPO		
Lactobacillus catenaformis		62.17	0.36	IND 1	1		
Clostridium barati		23.05	0.24	MLT 90	99		
Lactobacillus jensenii		14.79	0.26		1		
17251100	**Low discrimination**						**17251100**
					GRA	SPO	
Lactobacillus catenaformis		74.42	0.50	IND 1	99	1	
Fusobacterium mortiferum		20.18	0.50	MLT 90	1	1	
Clostridium barati		4.60	0.24		99	99	
17251102	**Low discrimination**						**17251102**
					SPO		
Lactobacillus catenaformis		68.02	0.50	IND 1	1		
Lactobacillus acidophilus		17.32	0.36	MLT 90	1		
Clostridium barati		12.61	0.32		99		
17251120	**Low discrimination**						**17251120**
					SPO		
Lactobacillus catenaformis		55.97	0.39	IND 1	1		
Clostridium barati		41.49	0.32	MLT 90	99		
Clostridium paraputrificum		0.84	-.03		99		
17251122	**Low discrimination**						**17251122**
					SPO		
Clostridium barati		52.22	0.40	IND 1	99		
Lactobacillus acidophilus		23.92	0.36	ESL 87	1		
Lactobacillus catenaformis		23.48	0.39		1		
17252000	**Good identification**						**17252000**
Clostridium innocuum		100.0	0.44	IND 1			
17252002	**Good identification**						**17252002**
Clostridium innocuum		100.0	0.44	IND 1			
17253000	**Very good identification**						**17253000**
					SPO		
Clostridium innocuum		99.92	0.66	IND 1	99		
Lactobacillus catenaformis		0.05	0.12		1		
Lactobacillus jensenii		0.01	0.02		1		
17253002	**Very good identification**						**17253002**
					SPO		
Clostridium innocuum		99.91	0.66	IND 1	99		
Lactobacillus catenaformis		0.05	0.12		1		
Clostridium barati		0.04	0.05		99		
17253010	**Good identification**						**17253010**
Clostridium innocuum		100.0	0.42	IND 1			
17253012	**Good identification**						**17253012**
Clostridium innocuum		100.0	0.42	IND 1			
17253400	**Good identification**						**17253400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.53	IND 1			
17253402	**Good identification**						**17253402**
Clostridium innocuum		100.0	0.53	IND 1			
17257000	**Good identification**						**17257000**
Clostridium innocuum		100.0	0.44	IND 1			
17257002	**Good identification**						**17257002**
Clostridium innocuum		100.0	0.44	IND 1			
17263000	**Good identification**						**17263000**
Clostridium innocuum		100.0	0.36	IND 1			
17263002	**Good identification**						**17263002**
Clostridium innocuum		100.0	0.36	IND 1			
17270100	**Good identification**						**17270100**
Fusobacterium mortiferum		100.0	0.38	IND 1			
17271000	**Low discrimination**						**17271000**
					GRA	SPO	
Lactobacillus jensenii		84.30	0.44	IND 1	99	1	
Clostridium innocuum		8.04	0.24	MLT 95	99	99	
Fusobacterium mortiferum		6.15	0.34		1	1	
17271002	**Low discrimination**						**17271002**
					SPO		
Lactobacillus jensenii		70.07	0.41	IND 1	1		
Lactobacillus acidophilus		16.38	0.27	MLT 95	1		
Clostridium innocuum		10.02	0.24		99		
17271010	**Good identification**						**17271010**
					SPO		
Lactobacillus jensenii		98.82	0.36	IND 1	1		
Clostridium innocuum		1.18	0.00	MLT 95	99		
17271100	**Low discrimination**						**17271100**
					COC	GRA	
Fusobacterium mortiferum		60.82	0.41	IND 1	1	1	
Atopobium parvulum		17.49	0.20		99	99	
Lactobacillus catenaformis		7.58	0.15		1	99	
17271102	**Good identification**						**17271102**
					SPO		
Lactobacillus acidophilus		95.25	0.45	IND 1	1		
Clostridium barati		1.80	0.14	MLT 94	99		
Lactobacillus catenaformis		1.13	0.15		1		
17271122	**Acceptable identification**						**17271122**
					SPO		
Lactobacillus acidophilus		93.99	0.45	IND 1	1		
Clostridium barati		5.33	0.22	MLT 94	99		
Clostridium ramosum		0.40	0.04		92		
17272000	**Good identification**						**17272000**
Clostridium innocuum		100.0	0.36	IND 1			
17272002	**Good identification**						**17272002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium innocuum		100.0	0.36	IND 1				
17273000	**Very good identification**							**17273000**
					SPO			
Clostridium innocuum		99.89	0.58	IND 1	99			
Lactobacillus jensenii		0.11	0.10		1			
17273002	**Very good identification**							**17273002**
					SPO			
Clostridium innocuum		99.84	0.58	IND 1	99			
Lactobacillus acidophilus		0.09	0.05		1			
Lactobacillus jensenii		0.07	0.07		1			
17273400	**Good identification**							**17273400**
Clostridium innocuum		100.0	0.45	IND 1				
17273402	**Good identification**							**17273402**
Clostridium innocuum		100.0	0.45	IND 1				
17277000	**Good identification**							**17277000**
Clostridium innocuum		100.0	0.36	IND 1				
17277002	**Good identification**							**17277002**
Clostridium innocuum		100.0	0.36	IND 1				
17300000	**Low discrimination**							**17300000**
					COC	GRA	SPO	
Clostridium bifermentans		85.97	0.43	SUC 1	1	99	99	
Gemella morbillorum		9.41	0.30	SAL 10	99	99	1	
Prevotella intermedia		4.26	0.23		1	1	1	
17310000	**Very good identification**							**17310000**
					COC	GRA	SPO	
Prevotella intermedia		95.19	0.58	SAL 1	1	1	1	
Gemella morbillorum		2.60	0.32		99	99	1	
Clostridium bifermentans		1.02	0.21		1	99	99	
17310100	**Low discrimination**							**17310100**
Fusobacterium mortiferum		65.57	0.39	IND 1				
Prevotella intermedia		34.44	0.23	MLT 11				
17311000	**Good identification**							**17311000**
					GRA			
Prevotella intermedia		91.28	0.58	SAL 1	1			
Lactobacillus jensenii		4.15	0.36		99			
Lactobacillus catenaformis		1.59	0.29		99			
17311020	**Low discrimination**							**17311020**
					GRA			
Clostridium perfringens		64.59	0.36	IND 1	99			
Lactobacillus jensenii		13.93	0.26	LAC 88	99			
Prevotella intermedia		12.37	0.23		1			
17311100	**Low discrimination**							**17311100**
					GRA			
Fusobacterium mortiferum		32.76	0.42	IND 1	1			
Lactobacillus catenaformis		29.62	0.32	MLT 11	99			
Clostridium perfringens		13.58	0.25		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17311120	**Acceptable identification**						**17311120**
					GRA		
Clostridium perfringens		93.97	0.51	IND 1	99		
Prevotella melaninogenica		1.98	0.15		1		
Lactobacillus catenaformis		1.59	0.21		99		
17311122	**Low discrimination**						**17311122**
					GRA		
Clostridium perfringens		77.43	0.45	IND 1	99		
Prevotella oralis		8.34	0.25		1		
Lactobacillus acidophilus		4.92	0.23		99		
17311123	**Low discrimination**						**17311123**
					SPO		
Actinomyces naeslundii		48.19	0.36	IND 1	1		
Clostridium perfringens		35.59	0.11	RAF 92	84		
Clostridium chauvoei		16.23	0.05		99		
17311160	**Good identification**						**17311160**
Clostridium perfringens		100.0	0.39	IND 1			
17313100	**Low discrimination**						**17313100**
Actinomyces naeslundii		88.62	0.38	IND 1			
Propionibacterium propionicum		11.38	0.12	RAF 92			
17313101	**Good identification**						**17313101**
Actinomyces naeslundii		100.0	0.41	IND 1			
17313102	**Low discrimination**						**17313102**
Actinomyces naeslundii		72.38	0.38	IND 1			
Propionibacterium propionicum		13.95	0.15	RAF 92			
Lactobacillus acidophilus		13.67	0.01				
17313103	**Good identification**						**17313103**
Actinomyces naeslundii		100.0	0.41	IND 1			
17313121	**Good identification**						**17313121**
Actinomyces naeslundii		100.0	0.41	IND 1			
17313122	**Low discrimination**						**17313122**
					SPO		
Actinomyces naeslundii		46.89	0.38	IND 1	1		
Clostridium perfringens		26.75	0.11	RAF 92	84		
Clostridium barati		9.96	-.01		99		
17313123	**Good identification**						**17313123**
Actinomyces naeslundii		100.0	0.41	IND 1			
17331000	**Low discrimination**						**17331000**
					GRA	CAT	
Lactobacillus jensenii		84.42	0.44	IND 1	99	1	
Prevotella intermedia		6.25	0.23	CEL 95	1	1	
Propionibacterium granulosum		2.33	0.24		99	90	
17331002	**Acceptable identification**						**17331002**
Lactobacillus jensenii		92.12	0.41	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		3.41	0.14	CEL 95			
Clostridium perfringens		1.44	0.09				
17331010	**Good identification**						**17331010**
					CAT		
Lactobacillus jensenii		97.82	0.36	IND 1	1		
Propionibacterium granulosum		1.10	0.09	CEL 95	90		
Eubacterium saburreum		1.08	0.11		1		
17331020	**Low discrimination**						**17331020**
Clostridium perfringens		72.74	0.41	IND 1			
Lactobacillus jensenii		25.34	0.34	LAC 88			
Actinomyces naeslundii		0.84	0.30				
17331022	**Low discrimination**						**17331022**
Clostridium perfringens		62.85	0.35	IND 1			
Lactobacillus jensenii		31.02	0.31	LAC 88			
Lactobacillus acidophilus		4.59	0.14				
17331100	**Low discrimination**						**17331100**
					COC	GRA	
Atopobium parvulum		88.50	0.48	IND 1	99	99	
Clostridium perfringens		6.93	0.30		1	99	
Fusobacterium mortiferum		2.69	0.33		1	1	
17331101	**Low discrimination**						**17331101**
					COC	SPO	
Actinomyces naeslundii		52.07	0.41	IND 1	1	1	
Atopobium parvulum		30.87	0.14	RAF 92	99	1	
Clostridium chauvoei		17.07	0.09		1	99	
17331102	**Low discrimination**						**17331102**
					COC		
Atopobium parvulum		47.98	0.35	IND 1	99		
Lactobacillus acidophilus		37.11	0.32		1		
Clostridium perfringens		10.02	0.24		1		
17331103	**Acceptable identification**						**17331103**
					COC		
Actinomyces naeslundii		90.53	0.41	IND 1	1		
Atopobium parvulum		9.47	0.01	RAF 92	99		
17331120	**Acceptable identification**						**17331120**
					COC	SPO	
Clostridium perfringens		91.94	0.56	IND 1	1	84	
Atopobium parvulum		6.41	0.35		99	1	
Clostridium septicum		1.24	0.17		1	99	
17331122	**Low discrimination**						**17331122**
					COC		
Clostridium perfringens		86.28	0.50	IND 1	1		
Lactobacillus acidophilus		9.88	0.32		1		
Atopobium parvulum		2.25	0.22		99		
17331123	**Low discrimination**						**17331123**
Actinomyces naeslundii		58.60	0.41	IND 1			
Clostridium perfringens		41.40	0.16	RAF 92			
17331160	**Good identification**						**17331160**
					COC		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		99.22	0.43	IND 1	1		
Actinomyces naeslundii		0.41	0.25		1		
Atopobium parvulum		0.37	0.01		99		
17331162	**Acceptable identification**						**17331162**
Clostridium perfringens		92.32	0.38	IND 1			
Lactobacillus acidophilus		6.86	0.16				
Actinomyces naeslundii		0.82	0.25				
17333001	**Good identification**						**17333001**
Actinomyces naeslundii		100.0	0.38	IND 1			
17333003	**Good identification**						**17333003**
Actinomyces naeslundii		100.0	0.38	IND 1			
17333021	**Good identification**						**17333021**
Actinomyces naeslundii		100.0	0.38	IND 1			
17333023	**Good identification**						**17333023**
Actinomyces naeslundii		100.0	0.38	IND 1			
17333100	**Low discrimination**						**17333100**
					COC	SPO	
Actinomyces naeslundii		64.89	0.43	IND 1	1	1	
Atopobium parvulum		29.72	0.14	RAF 92	99	1	
Clostridium tertium		5.38	0.01		1	99	
17333101	**Good identification**						**17333101**
Actinomyces naeslundii		100.0	0.46	IND 1			
17333102	**Low discrimination**						**17333102**
					SPO		
Actinomyces naeslundii		55.23	0.43	IND 1	1		
Lactobacillus acidophilus		17.99	0.10	RAF 92	1		
Clostridium tertium		13.74	0.09		99		
17333103	**Good identification**						**17333103**
Actinomyces naeslundii		100.0	0.46	IND 1			
17333121	**Good identification**						**17333121**
Actinomyces naeslundii		100.0	0.46	IND 1			
17333122	**Low discrimination**						**17333122**
					SPO		
Actinomyces naeslundii		28.12	0.43	IND 1	1		
Clostridium ramosum		26.38	0.17	RAF 92	92		
Clostridium tertium		20.99	0.17		99		
17333123	**Good identification**						**17333123**
Actinomyces naeslundii		100.0	0.46	IND 1			
17335100	**Good identification**						**17335100**
					COC		
Atopobium parvulum		99.61	0.53	IND 1	99		
Eubacterium saburreum		0.15	0.14		1		
Actinomyces israelii		0.12	0.16		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17335102	**Good identification**					**17335102**
					COC	
Atopobium parvulum		98.00	0.40	IND 1	99	
Bifidobacterium adolescentis		1.01	0.13		1	
Actinomyces israelii		0.99	0.19		1	
17335120	**Acceptable identification**					**17335120**
					COC	
Atopobium parvulum		92.76	0.40	IND 1	99	
Clostridium perfringens		7.24	0.21		1	
17351000	**Low discrimination**					**17351000**
					SPO	
Lactobacillus catenaformis		65.39	0.63	IND 1	1	
Lactobacillus jensenii		32.83	0.58		1	
Clostridium barati		1.10	0.27		99	
17351002	**Low discrimination**					**17351002**
Lactobacillus catenaformis		69.28	0.63	IND 1		
Lactobacillus jensenii		23.19	0.55			
Lactobacillus acidophilus		4.01	0.39			
17351010	**Acceptable identification**					**17351010**
Lactobacillus jensenii		94.31	0.50	IND 1		
Lactobacillus catenaformis		5.69	0.29			
17351012	**Acceptable identification**					**17351012**
Lactobacillus jensenii		90.88	0.47	IND 1		
Lactobacillus catenaformis		8.23	0.29			
Lactobacillus acidophilus		0.48	0.05			
17351020	**Low discrimination**					**17351020**
					SPO	
Lactobacillus catenaformis		58.31	0.53	IND 1	1	
Lactobacillus jensenii		29.28	0.47		1	
Clostridium barati		11.73	0.35		99	
17351022	**Low discrimination**					**17351022**
					SPO	
Lactobacillus catenaformis		45.77	0.53	IND 1	1	
Clostridium barati		27.61	0.44		99	
Lactobacillus jensenii		15.32	0.44		1	
17351030	**Acceptable identification**					**17351030**
					SPO	
Lactobacillus jensenii		93.24	0.39	IND 1	1	
Lactobacillus catenaformis		5.63	0.18		1	
Clostridium barati		1.13	0.01		99	
17351032	**Low discrimination**					**17351032**
					SPO	
Lactobacillus jensenii		85.76	0.36	IND 1	1	
Lactobacillus catenaformis		7.76	0.18		1	
Clostridium barati		4.68	0.09		99	
17351100	**Low discrimination**					**17351100**
					SPO	
Lactobacillus catenaformis		89.08	0.66	IND 1	1	
Collinsella aerofaciens		6.55	0.32		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		2.99	0.35		99		
17351102	**Low discrimination**						**17351102**
					SPO		
Lactobacillus catenaformis		64.20	0.66	IND 1	1		
Lactobacillus acidophilus		28.45	0.57		1		
Clostridium barati		6.46	0.44		99		
17351120	**Low discrimination**						**17351120**
					SPO		
Lactobacillus catenaformis		59.65	0.56	IND 1	1		
Clostridium barati		23.99	0.44		99		
Clostridium paraputrificum		9.82	0.32		99		
17351122	**Low discrimination**						**17351122**
					SPO		
Lactobacillus acidophilus		42.31	0.57	IND 1	1		
Clostridium barati		28.79	0.52		99		
Lactobacillus catenaformis		23.86	0.56		1		
17351142	**Low discrimination**						**17351142**
					SPO		
Lactobacillus acidophilus		74.41	0.41	IND 1	1		
Lactobacillus catenaformis		13.72	0.32	SOR 11	1		
Clostridium barati		11.88	0.25		99		
17351162	**Low discrimination**						**17351162**
					SPO		
Lactobacillus acidophilus		63.77	0.41	IND 1	1		
Clostridium barati		30.53	0.34	SOR 11	99		
Lactobacillus catenaformis		2.94	0.21		1		
17351302	**Low discrimination**						**17351302**
					SPO		
Lactobacillus acidophilus		63.17	0.38	IND 1	1		
Lactobacillus catenaformis		19.13	0.32	RHA 7	1		
Clostridium barati		16.56	0.25		99		
17351322	**Low discrimination**						**17351322**
					GRA	SPO	
Lactobacillus acidophilus		43.22	0.38	IND 1	99	1	
Clostridium barati		33.98	0.34	RHA 7	99	99	
Prevotella oralis		14.95	0.29		1	1	
17353001	**Good identification**						**17353001**
Actinomyces naeslundii		100.0	0.35	IND 1			
17353003	**Good identification**						**17353003**
Actinomyces naeslundii		100.0	0.35	IND 1			
17353021	**Good identification**						**17353021**
Actinomyces naeslundii		100.0	0.35	IND 1			
17353023	**Good identification**						**17353023**
Actinomyces naeslundii		100.0	0.35	IND 1			
17353101	**Good identification**						**17353101**
Actinomyces naeslundii		100.0	0.43	IND 1			
17353102	**Low discrimination**						**17353102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Lactobacillus acidophilus		45.66	0.35	IND 1	1	
Lactobacillus catenaformis		19.77	0.32	MAN 5	1	
Clostridium barati		17.11	0.25		99	
17353103	**Acceptable identification**					**17353103**
Actinomyces naeslundii		91.33	0.43	IND 1		
Lactobacillus acidophilus		8.67	0.01	RAF 92		
17353121	**Good identification**					**17353121**
Actinomyces naeslundii		100.0	0.43	IND 1		
17353123	**Low discrimination**					**17353123**
					SPO	
Actinomyces naeslundii		77.96	0.43	IND 1	1	
Clostridium barati		8.33	-.01	RAF 92	99	
Lactobacillus acidophilus		7.40	0.01		1	
17353132	**Good identification**					**17353132**
					SPO	
Clostridium tertium		98.03	0.39	IND 1	99	
Clostridium barati		0.74	-.01	TRE 90	99	
Lactobacillus acidophilus		0.65	0.01		1	
17355502	**Low discrimination**					**17355502**
					SPO	
Bifidobacterium adolescentis		68.05	0.36	IND 1	1	
Clostridium butyricum		15.19	0.10	RAF 92	99	
Actinomyces israelii		9.03	0.27		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17355512	**Good identification**					**17355512**
Bifidobacterium adolescentis		97.57	0.37	IND 1		
Clostridium clostridioforme		2.43	0.07	RAF 92		
17355522	**Good identification**					**17355522**
					CAT	
Bacteroides uniformis		98.20	0.38	ESL 99	5	
Bacteroides ovatus		1.54	0.10	RAF 97	68	
Bacteroides thetaiotaomicron		0.26	-.04		89	
17355722	**Low discrimination**					**17355722**
					CAT	
Bacteroides uniformis		80.34	0.40	ESL 99	5	
Bacteroides ovatus		17.34	0.31	RAF 97	68	
Bacteroides thetaiotaomicron		2.07	0.16		89	
17357100	**Low discrimination**					**17357100**
					SPO	
Actinomyces israelii		61.37	0.39	IND 1	1	
Clostridium tertium		38.63	0.21		99	
17357122	**Good identification**					**17357122**
					SPO	
Clostridium tertium		96.61	0.37	IND 1	99	
Actinomyces israelii		1.63	0.21	TRE 90	1	
Clostridium barati		0.93	-.01		99	
17357132	**Good identification**					**17357132**
Clostridium tertium		100.0	0.41	IND 1		
17357500	**Low discrimination**					**17357500**
					SPO	
Actinomyces israelii		88.71	0.37	IND 1	1	
Clostridium butyricum		11.30	0.01		99	
17357512	**Low discrimination**					**17357512**
					SPO	
Bifidobacterium adolescentis		88.26	0.36	IND 1	1	
Actinomyces israelii		9.91	0.26	RAF 92	1	
Clostridium tertium		1.83	-.01		99	
17371000	**Very good identification**					**17371000**
					COC	
Lactobacillus jensenii		98.76	0.66	IND 1	1	
Lactobacillus catenaformis		0.66	0.29		1	
Atopobium parvulum		0.26	0.20		99	
17371002	**Low discrimination**					**17371002**
Lactobacillus jensenii		82.73	0.63	IND 1		
Lactobacillus acidophilus		15.95	0.48			
Lactobacillus catenaformis		0.83	0.29			
17371010	**Very good identification**					**17371010**
Lactobacillus jensenii		100.0	0.58	IND 1		
17371012	**Good identification**					**17371012**
					GRA	
Lactobacillus jensenii		99.17	0.55	IND 1	99	
Lactobacillus acidophilus		0.58	0.14		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		0.25	-.03		1		
17371020	**Good identification**						**17371020**
					SPO		
Lactobacillus jensenii		96.72	0.56	IND 1	1		
Clostridium barati		1.12	0.17		99		
Lactobacillus catenaformis		0.65	0.18		1		
17371022	**Low discrimination**						**17371022**
					SPO		
Lactobacillus jensenii		54.24	0.53	IND 1	1		
Lactobacillus acidophilus		41.82	0.48		1		
Clostridium barati		2.83	0.25		99		
17371030	**Good identification**						**17371030**
Lactobacillus jensenii		100.0	0.47	IND 1			
17371032	**Good identification**						**17371032**
Lactobacillus jensenii		97.72	0.44	IND 1			
Lactobacillus acidophilus		2.28	0.14				
17371100	**Low discrimination**						**17371100**
					COC		
Atopobium parvulum		85.70	0.54	IND 1	99		
Lactobacillus acidophilus		5.01	0.32		1		
Lactobacillus jensenii		3.39	0.32		1		
17371102	**Very good identification**						**17371102**
					COC		
Lactobacillus acidophilus		95.21	0.66	IND 1	1		
Atopobium parvulum		2.90	0.41		99		
Lactobacillus catenaformis		0.65	0.32		1		
17371120	**Low discrimination**						**17371120**
					COC	SPO	
Atopobium parvulum		37.49	0.41	IND 1	99	1	
Lactobacillus acidophilus		12.43	0.32		1	1	
Clostridium septicum		9.87	0.25		1	99	
17371121	**Low discrimination**						**17371121**
					COC		
Actinomyces naeslundii		84.45	0.43	IND 1	1		
Atopobium parvulum		15.55	0.07	RAF 92	99		
17371122	**Good identification**						**17371122**
					SPO		
Lactobacillus acidophilus		94.06	0.66	IND 1	1		
Clostridium ramosum		2.52	0.38		92		
Clostridium barati		1.66	0.34		99		
17371142	**Good identification**						**17371142**
					COC	SPO	
Lactobacillus acidophilus		99.10	0.50	IND 1	1	1	
Clostridium barati		0.41	0.07	SOR 11	1	99	
Atopobium parvulum		0.25	0.07		99	1	
17371162	**Good identification**						**17371162**
					SPO		
Lactobacillus acidophilus		97.68	0.50	IND 1	1		
Clostridium barati		1.22	0.15	SOR 11	99		
Clostridium perfringens		0.55	0.14		84		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17371302	**Good identification**						**17371302**
					SPO		
Lactobacillus acidophilus		98.46	0.47	IND 1	1		
Clostridium barati		0.67	0.07	RHA 7	99		
Clostridium ramosum		0.47	0.06		92		
17371322	**Low discrimination**						**17371322**
					SPO		
Lactobacillus acidophilus		66.75	0.47	IND 1	1		
Clostridium ramosum		31.45	0.40	RHA 7	92		
Clostridium barati		1.36	0.15		99		
17373001	**Good identification**						**17373001**
Actinomyces naeslundii		100.0	0.40	IND 1			
17373003	**Good identification**						**17373003**
Actinomyces naeslundii		100.0	0.40	IND 1			
17373021	**Good identification**						**17373021**
Actinomyces naeslundii		100.0	0.40	IND 1			
17373023	**Good identification**						**17373023**
Actinomyces naeslundii		100.0	0.40	IND 1			
17373101	**Good identification**						**17373101**
					SPO		
Actinomyces naeslundii		96.00	0.48	IND 1	1		
Clostridium tertium		4.01	0.01	RAF 92	99		
17373102	**Low discrimination**						**17373102**
					SPO		
Lactobacillus acidophilus		53.19	0.44	IND 1	1		
Clostridium tertium		40.63	0.43	MAN 5	99		
Actinomyces naeslundii		2.19	0.45		1		
17373103	**Low discrimination**						**17373103**
					SPO		
Actinomyces naeslundii		77.58	0.48	IND 1	1		
Lactobacillus acidophilus		12.71	0.10	RAF 92	1		
Clostridium tertium		9.71	0.09		99		
17373110	**Good identification**						**17373110**
					SPO		
Clostridium tertium		98.87	0.39	IND 1	99		
Actinomyces israelii		0.66	0.16		1		
Actinomyces naeslundii		0.47	0.23		1		
17373112	**Good identification**						**17373112**
					GRA	SPO	
Clostridium tertium		98.26	0.47	IND 1	99	99	
Lactobacillus acidophilus		0.73	0.10		99	1	
Leptotrichia buccalis		0.52	-.03		1	1	
17373120	**Low discrimination**						**17373120**
					SPO		
Clostridium tertium		49.26	0.43	IND 1	99		
Clostridium ramosum		46.43	0.41		92		
Actinomyces naeslundii		2.65	0.45		1		
17373121	**Low discrimination**						**17373121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		80.44	0.48	IND 1	1		
Clostridium tertium		10.07	0.09	RAF 92	99		
Clostridium ramosum		9.49	0.07		92		
17373122	**Low discrimination**						**17373122**
					SPO		
Clostridium ramosum		46.14	0.51	IND 1	92		
Clostridium tertium		36.71	0.51		99		
Lactobacillus acidophilus		16.02	0.44		1		
17373123	**Low discrimination**						**17373123**
					SPO		
Actinomyces naeslundii		49.72	0.48	IND 1	1		
Clostridium ramosum		23.46	0.17	RAF 92	92		
Clostridium tertium		18.67	0.17		99		
17373130	**Good identification**						**17373130**
					SPO		
Clostridium tertium		99.31	0.47	IND 1	99		
Clostridium ramosum		0.53	0.07		92		
Actinomyces naeslundii		0.16	0.23		1		
17373132	**Good identification**						**17373132**
					SPO		
Clostridium tertium		99.00	0.55	IND 1	99		
Clostridium ramosum		0.71	0.17		92		
Lactobacillus acidophilus		0.25	0.10		1		
17373320	**Good identification**						**17373320**
Clostridium ramosum		99.20	0.43	IND 1			
Clostridium tertium		0.80	0.09				
17373322	**Good identification**						**17373322**
					SPO		
Clostridium ramosum		97.42	0.53	IND 1	92		
Lactobacillus acidophilus		1.92	0.25		1		
Clostridium tertium		0.59	0.17		99		
17375100	**Very good identification**						**17375100**
					COC	SPO	
Atopobium parvulum		97.66	0.59	IND 1	99	1	
Clostridium butyricum		1.62	0.21		1	99	
Actinomyces israelii		0.66	0.36		1	1	
17375102	**Low discrimination**						**17375102**
					COC		
Atopobium parvulum		77.41	0.46	IND 1	99		
Lactobacillus acidophilus		13.82	0.32		1		
Actinomyces israelii		4.45	0.39		1		
17375120	**Good identification**						**17375120**
					COC	SPO	
Atopobium parvulum		97.99	0.46	IND 1	99	1	
Clostridium tertium		0.97	0.11		1	99	
Clostridium butyricum		0.80	0.03		1	99	
17375500	**Acceptable identification**						**17375500**
					COC	SPO	
Clostridium butyricum		90.54	0.38	IND 3	1	99	
Atopobium parvulum		6.11	0.25	ESL 88	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		3.35	0.34	RAF 95	1	1	
17375712	**Low discrimination**						**17375712**
Clostridium clostridioforme		86.79	0.36	IND 1			
Bacteroides ovatus		12.11	0.12	RAF 94			
Bacteroides thetaiotaomicron		1.11	-.06				
17375722	**Low discrimination**						**17375722**
Bacteroides ovatus		85.21	0.49	ESL 91			
Bacteroides thetaiotaomicron		13.35	0.35	RAF 99			
Clostridium clostridioforme		0.90	0.25				
17375732	**Low discrimination**						**17375732**
Bacteroides ovatus		85.83	0.46	ESL 91			
Bacteroides thetaiotaomicron		7.84	0.28	RAF 99			
Clostridium clostridioforme		6.22	0.36				
17377000	**Low discrimination**						**17377000**
					SPO		
Actinomyces israelii		89.37	0.38	IND 1	1		
Clostridium tertium		5.44	0.03		99		
Clostridium innocuum		5.20	0.02		99		
17377002	**Low discrimination**						**17377002**
					SPO		
Actinomyces israelii		86.17	0.41	IND 1	1		
Clostridium tertium		10.49	0.11		99		
Clostridium innocuum		3.34	0.02		99		
17377100	**Low discrimination**						**17377100**
					COC	SPO	
Clostridium tertium		53.45	0.37	IND 1	1	99	
Actinomyces israelii		35.49	0.48		1	1	
Atopobium parvulum		9.35	0.25		99	1	
17377102	**Low discrimination**						**17377102**
					COC	SPO	
Clostridium tertium		73.32	0.45	IND 1	1	99	
Actinomyces israelii		24.34	0.51		1	1	
Atopobium parvulum		0.76	0.12		99	1	
17377110	**Acceptable identification**						**17377110**
					SPO		
Clostridium tertium		94.71	0.41	IND 1	99		
Actinomyces israelii		5.29	0.34		1		
17377112	**Good identification**						**17377112**
					SPO		
Clostridium tertium		96.78	0.50	IND 1	99		
Actinomyces israelii		2.70	0.37		1		
Bifidobacterium adolescentis		0.52	0.19		1		
17377120	**Good identification**						**17377120**
					COC	SPO	
Clostridium tertium		96.96	0.45	IND 1	1	99	
Actinomyces israelii		1.37	0.28		1	1	
Atopobium parvulum		1.00	0.12		99	1	
17377122	**Good identification**						**17377122**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		98.09	0.53	IND 1	99	
Clostridium ramosum		0.90	0.17		92	
Actinomyces israelii		0.69	0.31		1	
17377130	**Good identification**					**17377130**
					SPO	
Clostridium tertium		99.88	0.50	IND 1	99	
Actinomyces israelii		0.12	0.14		1	
17377132	**Very good identification**					**17377132**
					SPO	
Clostridium tertium		99.94	0.58	IND 1	99	
Actinomyces israelii		0.06	0.17		1	
17377140	**Good identification**					**17377140**
					SPO	
Actinomyces israelii		96.24	0.41	IND 1	1	
Clostridium tertium		3.76	0.03		99	
17377142	**Acceptable identification**					**17377142**
					SPO	
Actinomyces israelii		92.74	0.44	IND 1	1	
Clostridium tertium		7.26	0.11		99	
17377400	**Acceptable identification**					**17377400**
					SPO	
Actinomyces israelii		91.88	0.37	IND 1	1	
Clostridium butyricum		8.12	-.03		99	
17377402	**Good identification**					**17377402**
Actinomyces israelii		100.0	0.40	IND 1		
17377500	**Low discrimination**					**17377500**
					SPO	
Actinomyces israelii		64.56	0.47	IND 1	1	
Clostridium butyricum		34.24	0.21		99	
Clostridium tertium		1.20	0.03		99	
17377502	**Low discrimination**					**17377502**
					SPO	
Actinomyces israelii		73.12	0.50	IND 1	1	
Bifidobacterium adolescentis		14.11	0.31		1	
Clostridium butyricum		10.05	0.14		99	
17377540	**Low discrimination**					**17377540**
					SPO	
Actinomyces israelii		89.40	0.40	IND 1	1	
Clostridium butyricum		10.60	0.03		99	
17377542	**Acceptable identification**					**17377542**
					SPO	
Actinomyces israelii		91.57	0.43	IND 1	1	
Bifidobacterium adolescentis		5.62	0.16		1	
Clostridium butyricum		2.81	-.04		99	
17377722	**Good identification**					**17377722**
Bacteroides ovatus		99.53	0.41	ESL 91		
Bacteroides thetaiotaomicron		0.47	0.01	RAF 99		
17377732	**Good identification**					**17377732**
Bacteroides ovatus		99.72	0.38	ESL 91		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides thetaiotaomicron		0.28	-.06	RAF 99				
17400020	**Good identification**							**17400020**
					SPO	CAT		
Clostridium cadaveris		99.37	0.44	SUC 7	99	1		
Propionibacterium granulosum		0.31	0.13	RAF 1	1	90		
Clostridium bifermentans		0.18	-.01		99	1		
17410000	**Low discrimination**							**17410000**
					GRA	CAT		
Prevotella intermedia		74.28	0.61	MLT 99	1	1		
Propionibacterium granulosum		17.42	0.58		99	90		
Fusobacterium varium		4.60	0.32		1	1		
17410002	**Low discrimination**							**17410002**
					GRA	CAT		
Propionibacterium granulosum		78.91	0.47	IND 1	99	90		
Prevotella intermedia		13.60	0.26		1	1		
Fusobacterium mortiferum		6.65	0.32		1	5		
17410010	**Low discrimination**							**17410010**
					GRA	CAT		
Propionibacterium granulosum		74.10	0.43	IND 1	99	90		
Prevotella intermedia		23.41	0.26	MLZ 12	1	1		
Fusobacterium varium		1.45	-.03		1	1		
17410020	**Low discrimination**							**17410020**
					GRA	SPO	CAT	
Clostridium cadaveris		72.57	0.44	SUC 7	99	99	1	
Propionibacterium granulosum		22.23	0.47	RAF 1	99	1	90	
Prevotella intermedia		3.83	0.26		1	1	1	
17410022	**Low discrimination**							**17410022**
					SPO	CAT		
Propionibacterium granulosum		88.35	0.37	IND 1	1	90		
Clostridium cadaveris		11.65	0.09		99	1		
17410100	**Low discrimination**							**17410100**
					GRA	CAT		
Fusobacterium mortiferum		88.48	0.55	IND 1	1	5		
Prevotella intermedia		8.62	0.26		1	1		
Propionibacterium granulosum		2.02	0.24		99	90		
17410102	**Acceptable identification**							**17410102**
					GRA	CAT		
Fusobacterium mortiferum		91.97	0.39	IND 1	1	5		
Propionibacterium granulosum		4.73	0.13	BGL 10	99	90		
Lactobacillus fermentum		3.30	0.15		99	1		
17411000	**Low discrimination**							**17411000**
					GRA	CAT		
Prevotella intermedia		74.81	0.61	MLT 99	1	1		
Propionibacterium granulosum		17.54	0.58		99	90		
Fusobacterium mortiferum		4.99	0.52		1	5		
17411002	**Low discrimination**							**17411002**
					GRA	CAT		
Propionibacterium granulosum		57.08	0.47	IND 1	99	90		
Lactobacillus fermentum		24.46	0.46		99	1		
Prevotella intermedia		9.84	0.26		1	1		
17411010	**Low discrimination**							**17411010**
					CAT	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		64.67	0.43	IND 1	90	99	
Prevotella intermedia		20.43	0.26	MLZ 12	1	1	
Eubacterium saburreum		13.54	0.33		1	85	
17411020	**Low discrimination**						**17411020**
					GRA	CAT	
Propionibacterium granulosum		69.95	0.47	IND 1	99	90	
Prevotella intermedia		12.05	0.26		1	1	
Lactobacillus fermentum		9.99	0.38		99	1	
17411022	**Low discrimination**						**17411022**
					CAT		
Propionibacterium granulosum		63.64	0.37	IND 1	90		
Lactobacillus fermentum		36.36	0.38		1		
17411100	**Low discrimination**						**17411100**
					GRA		
Fusobacterium mortiferum		68.06	0.58	IND 1	1		
Lactobacillus fermentum		17.92	0.50		99		
Eubacterium saburreum		8.35	0.41		85		
17411102	**Low discrimination**						**17411102**
					GRA		
Lactobacillus fermentum		65.85	0.50	IND 1	99		
Fusobacterium mortiferum		27.79	0.42		1		
Eubacterium saburreum		5.42	0.28		85		
17411110	**Low discrimination**						**17411110**
					GRA		
Eubacterium saburreum		87.13	0.39	ESL 99	85		
Fusobacterium mortiferum		8.77	0.24		1		
Lactobacillus fermentum		2.31	0.15		99		
17411122	**Low discrimination**						**17411122**
					GRA		
Lactobacillus fermentum		52.84	0.41	IND 1	99		
Prevotella melaninogenica		36.03	0.19		1		
Clostridium perfringens		3.61	0.09		99		
17411400	**Low discrimination**						**17411400**
					GRA	CAT	
Lactobacillus fermentum		65.65	0.46	IND 1	99	1	
Prevotella intermedia		26.40	0.26		1	1	
Propionibacterium granulosum		6.19	0.24		99	90	
17411402	**Good identification**						**17411402**
					CAT		
Lactobacillus fermentum		97.70	0.46	IND 1	1		
Propionibacterium granulosum		2.30	0.13		90		
17411420	**Acceptable identification**						**17411420**
					CAT		
Lactobacillus fermentum		93.39	0.38	IND 1	1		
Propionibacterium granulosum		6.61	0.13		90		
17411422	**Good identification**						**17411422**
Lactobacillus fermentum		100.0	0.38	IND 1			
17411500	**Good identification**						**17411500**
					GRA		
Lactobacillus fermentum		95.25	0.50	IND 1	99		
Fusobacterium mortiferum		3.65	0.24		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		1.10	0.09		99		
17411502	**Good identification**						**17411502**
Lactobacillus fermentum		98.86	0.50	IND 1			
Bifidobacterium longum subsp. longum		1.14	0.09				
17411520	**Low discrimination**						**17411520**
					GRA		
Lactobacillus fermentum		81.38	0.41	IND 1	99		
Bifidobacterium longum subsp. longum		11.23	0.20		99		
Odoribacter splanchnicus		4.45	-.03		1		
17411522	**Low discrimination**						**17411522**
					GRA		
Lactobacillus fermentum		84.65	0.41	IND 1	99		
Bifidobacterium longum subsp. longum		11.68	0.20		99		
Parabacteroides distasonis		1.88	0.10		1		
17412000	**Acceptable identification**						**17412000**
					GRA	CAT	
Propionibacterium granulosum		93.89	0.56	IND 1	99	90	
Prevotella intermedia		5.36	0.26		1	1	
Fusobacterium varium		0.33	-.03		1	1	
17412002	**Good identification**						**17412002**
Propionibacterium granulosum		100.0	0.45	IND 1			
17412010	**Good identification**						**17412010**
Propionibacterium granulosum		100.0	0.41	IND 1			
17412020	**Acceptable identification**						**17412020**
					SPO	CAT	
Propionibacterium granulosum		93.35	0.45	IND 1	1	90	
Clostridium cadaveris		4.08	0.09		99	1	
Propionibacterium acnes		2.58	0.09		1	95	
17412022	**Good identification**						**17412022**
Propionibacterium granulosum		100.0	0.35	IND 1			
17413000	**Acceptable identification**						**17413000**
					GRA	CAT	
Propionibacterium granulosum		94.26	0.56	IND 1	99	90	
Prevotella intermedia		5.38	0.26		1	1	
Fusobacterium mortiferum		0.36	0.17		1	5	
17413002	**Good identification**						**17413002**
Propionibacterium granulosum		100.0	0.45	IND 1			
17413010	**Good identification**						**17413010**
Propionibacterium granulosum		100.0	0.41	IND 1			
17413020	**Good identification**						**17413020**
Propionibacterium granulosum		97.32	0.45	IND 1			
Propionibacterium acnes		2.68	0.09				
17413022	**Good identification**						**17413022**
Propionibacterium granulosum		100.0	0.35	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17414100	**Good identification**						**17414100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
17415000	**Low discrimination**						**17415000**
					GRA		
Lactobacillus fermentum		40.25	0.42	IND 1	99		
Prevotella intermedia		27.57	0.26		1		
Fusobacterium mortiferum		13.70	0.32		1		
17415002	**Low discrimination**						**17415002**
					CAT		
Lactobacillus fermentum		88.44	0.42	IND 1	1		
Eubacterium saburreum		4.66	0.17		1		
Propionibacterium granulosum		3.55	0.13		90		
17415100	**Low discrimination**						**17415100**
					GRA		
Lactobacillus fermentum		52.25	0.46	IND 1	99		
Fusobacterium mortiferum		25.43	0.39		1		
Eubacterium saburreum		22.32	0.36		85		
17415102	**Low discrimination**						**17415102**
					GRA		
Lactobacillus fermentum		88.54	0.46	IND 1	99		
Eubacterium saburreum		6.68	0.23		85		
Fusobacterium mortiferum		4.79	0.22		1		
17415120	**Low discrimination**						**17415120**
					GRA		
Lactobacillus fermentum		77.65	0.37	IND 1	99		
Eubacterium saburreum		17.57	0.23		85		
Prevotella melaninogenica		4.78	-.03		1		
17415122	**Low discrimination**						**17415122**
					GRA	CAT	
Lactobacillus fermentum		57.99	0.37	IND 1	99	1	
Parabacteroides distasonis		32.26	0.30		1	76	
Bacteroides fragilis		7.44	-.03		1	99	
17415322	**Acceptable identification**						**17415322**
Parabacteroides distasonis		93.53	0.38	IND 1			
Bacteroides thetaiotaomicron		6.48	0.02				
17415400	**Good identification**						**17415400**
Lactobacillus fermentum		100.0	0.42	IND 1			
17415402	**Good identification**						**17415402**
Lactobacillus fermentum		100.0	0.42	IND 1			
17415500	**Acceptable identification**						**17415500**
Lactobacillus fermentum		94.45	0.46	IND 1			
Bifidobacterium longum subsp. longum		5.55	0.17				
17415502	**Acceptable identification**						**17415502**
Lactobacillus fermentum		94.45	0.46	IND 1			
Bifidobacterium longum subsp. longum		5.55	0.17				
17415520	**Low discrimination**						**17415520**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				GRA		
Lactobacillus fermentum	47.78	0.37	IND 1	99		
Bifidobacterium longum subsp. longum	33.67	0.28		99		
Bacteroides vulgatus	18.55	0.09		1		
17415522 Low discrimination						**17415522**
				GRA	CAT	
Lactobacillus fermentum	33.86	0.37	IND 1	99	1	
Bacteroides thetaiotaomicron	29.15	0.17		1	89	
Bifidobacterium longum subsp. longum	23.86	0.28		99	1	
17415722 Acceptable identification						**17415722**
				CAT		
Bacteroides thetaiotaomicron	91.06	0.37	ESL 92	89		
Parabacteroides distasonis	8.14	0.35	MLT 94	76		
Bacteroides uniformis	0.52	-.05	TRE 93	5		
17430000 Very good identification						**17430000**
				GRA	CAT	
Propionibacterium granulosum	96.70	0.66	IND 1	99	90	
Fusobacterium mortiferum	1.83	0.40		1	5	
Prevotella intermedia	1.39	0.26		1	1	
17430002 Good identification						**17430002**
				GRA	CAT	
Propionibacterium granulosum	99.17	0.56	IND 1	99	90	
Fusobacterium mortiferum	0.83	0.23		1	5	
17430010 Good identification						**17430010**
Propionibacterium granulosum	100.0	0.51	IND 1			
17430012 Good identification						**17430012**
Propionibacterium granulosum	100.0	0.41	IND 1			
17430020 Acceptable identification						**17430020**
				SPO	CAT	
Propionibacterium granulosum	94.58	0.56	IND 1	1	90	
Clostridium cadaveris	5.42	0.22		99	1	
17430022 Good identification						**17430022**
Propionibacterium granulosum	100.0	0.45	IND 1			
17430030 Good identification						**17430030**
Propionibacterium granulosum	100.0	0.41	IND 1			
17430100 Low discrimination						**17430100**
				GRA	CAT	
Fusobacterium mortiferum	81.35	0.46	IND 1	1	5	
Propionibacterium granulosum	18.65	0.32		99	90	
17430200 Good identification						**17430200**
Propionibacterium granulosum	100.0	0.40	IND 1			
17431000 Very good identification						**17431000**
				GRA	CAT	
Propionibacterium granulosum	95.20	0.66	IND 1	99	90	
Fusobacterium mortiferum	2.70	0.43		1	5	
Prevotella intermedia	1.37	0.26		1	1	
17431002 Good identification						**17431002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT	GRA	
Propionibacterium granulosum		98.23	0.56	IND 1	90	99	
Fusobacterium mortiferum		1.24	0.26		5	1	
Eubacterium saburreum		0.54	0.18		1	85	
17431010	**Good identification**						**17431010**
					CAT		
Propionibacterium granulosum		95.55	0.51	IND 1	90		
Eubacterium saburreum		4.45	0.30	MLZ 12	1		
17431012	**Good identification**						**17431012**
					CAT		
Propionibacterium granulosum		96.82	0.41	IND 1	90		
Eubacterium saburreum		3.18	0.17	MLZ 12	1		
17431020	**Good identification**						**17431020**
					CAT		
Propionibacterium granulosum		99.17	0.56	IND 1	90		
Eubacterium saburreum		0.54	0.18		1		
Clostridium perfringens		0.28	0.05		1		
17431022	**Good identification**						**17431022**
Propionibacterium granulosum		100.0	0.45	IND 1			
17431030	**Good identification**						**17431030**
					CAT		
Propionibacterium granulosum		96.82	0.41	IND 1	90		
Eubacterium saburreum		3.18	0.17	MLZ 12	1		
17431100	**Low discrimination**						**17431100**
					CAT	GRA	
Fusobacterium mortiferum		69.20	0.49	IND 1	5	1	
Eubacterium saburreum		18.95	0.38		1	85	
Propionibacterium granulosum		10.58	0.32		90	99	
17431110	**Low discrimination**						**17431110**
					CAT	GRA	
Eubacterium saburreum		87.87	0.36	ESL 99	1	85	
Propionibacterium granulosum		8.17	0.17		90	99	
Fusobacterium mortiferum		3.96	0.15		5	1	
17431200	**Good identification**						**17431200**
Propionibacterium granulosum		100.0	0.40	IND 1			
17432000	**Very good identification**						**17432000**
Propionibacterium granulosum		100.0	0.64	IND 1			
17432002	**Good identification**						**17432002**
Propionibacterium granulosum		100.0	0.53	IND 1			
17432010	**Good identification**						**17432010**
Propionibacterium granulosum		100.0	0.49	IND 1			
17432012	**Good identification**						**17432012**
Propionibacterium granulosum		100.0	0.39	IND 1			
17432020	**Good identification**						**17432020**
Propionibacterium granulosum		100.0	0.53	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17432022	**Good identification**					**17432022**
Propionibacterium granulosum		100.0	0.43	IND 1		
17432030	**Good identification**					**17432030**
Propionibacterium granulosum		100.0	0.39	IND 1		
17432200	**Good identification**					**17432200**
Propionibacterium granulosum		100.0	0.38	IND 1		
17433000	**Very good identification**					**17433000**
Propionibacterium granulosum		99.94	0.64	IND 1		
Actinomyces naeslundii		0.06	0.23			
17433002	**Good identification**					**17433002**
Propionibacterium granulosum		99.77	0.53	IND 1		
Actinomyces naeslundii		0.23	0.23			
17433010	**Good identification**					**17433010**
Propionibacterium granulosum		100.0	0.49	IND 1		
17433012	**Good identification**					**17433012**
Propionibacterium granulosum		100.0	0.39	IND 1		
17433020	**Good identification**					**17433020**
Propionibacterium granulosum		99.77	0.53	IND 1		
Actinomyces naeslundii		0.23	0.23			
17433022	**Good identification**					**17433022**
Propionibacterium granulosum		99.09	0.43	IND 1		
Actinomyces naeslundii		0.91	0.23			
17433030	**Good identification**					**17433030**
Propionibacterium granulosum		100.0	0.39	IND 1		
17433200	**Good identification**					**17433200**
Propionibacterium granulosum		100.0	0.38	IND 1		
17435122	**Good identification**					**17435122**
Parabacteroides distasonis		95.64	0.41	IND 1		
Bacteroides thetaiotaomicron		4.36	0.02			
17435132	**Good identification**					**17435132**
Parabacteroides distasonis		96.81	0.37	IND 1		
Bacteroides thetaiotaomicron		3.19	-.05			
17435322	**Low discrimination**					**17435322**
Parabacteroides distasonis		83.20	0.50	IND 1		
Bacteroides thetaiotaomicron		16.80	0.22			
17435332	**Low discrimination**					**17435332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		87.27	0.45	IND 1			
Bacteroides thetaiotaomicron		12.73	0.15				
17435522	**Low discrimination**						**17435522**
					GRA	CAT	
Bacteroides thetaiotaomicron		86.87	0.37	ESL 92	1	89	
Parabacteroides distasonis		11.80	0.38	MLT 94	1	76	
Bifidobacterium longum subsp. longum		1.34	0.17	RHA 93	99	1	
17435722	**Good identification**						**17435722**
Bacteroides thetaiotaomicron		96.98	0.56	ESL 92			
Parabacteroides distasonis		2.97	0.46	MLT 94			
Bacteroides ovatus		0.05	0.00				
17435732	**Good identification**						**17435732**
Bacteroides thetaiotaomicron		95.85	0.49	ESL 92			
Parabacteroides distasonis		4.07	0.41	MLT 94			
Bacteroides ovatus		0.09	-.03				
17450000	**Low discrimination**						**17450000**
					GRA	CAT	
Fusobacterium mortiferum		53.09	0.40	IND 1	1	5	
Prevotella intermedia		36.19	0.26		1	1	
Propionibacterium granulosum		8.49	0.24		99	90	
17450100	**Good identification**						**17450100**
Fusobacterium mortiferum		100.0	0.47	IND 1			
17451000	**Low discrimination**						**17451000**
					GRA	CAT	
Fusobacterium mortiferum		64.06	0.43	IND 1	1	5	
Prevotella intermedia		29.11	0.26		1	1	
Propionibacterium granulosum		6.83	0.24		99	90	
17451100	**Good identification**						**17451100**
					GRA		
Fusobacterium mortiferum		98.64	0.50	IND 1	1		
Lactobacillus fermentum		0.79	0.15		99		
Actinomyces naeslundii		0.58	0.23		99		
17455722	**Low discrimination**						**17455722**
					CAT		
Bacteroides thetaiotaomicron		75.76	0.42	ESL 92	89		
Bacteroides uniformis		20.11	0.30	MLT 94	5		
Bacteroides ovatus		2.42	0.17	TRE 93	68		
17455732	**Acceptable identification**						**17455732**
Bacteroides thetaiotaomicron		91.49	0.35	ESL 92			
Bacteroides ovatus		5.01	0.14	MLT 94			
Parabacteroides distasonis		2.87	0.25	TRE 93			
17470100	**Good identification**						**17470100**
Fusobacterium mortiferum		100.0	0.38	IND 1			
17471100	**Good identification**						**17471100**
					GRA		
Fusobacterium mortiferum		96.34	0.41	IND 1	1		
Actinomyces naeslundii		3.67	0.28		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17471722	**Good identification**					**17471722**
Bacteroides thetaiotaomicron		97.41	0.38	ESL 92		
Parabacteroides distasonis		2.02	0.25	MLT 94		
Bacteroides ovatus		0.57	0.00	XYL 96		
17473101	**Good identification**					**17473101**
Actinomyces naeslundii		100.0	0.36	IND 1		
17473103	**Good identification**					**17473103**
Actinomyces naeslundii		100.0	0.36	IND 1		
17473121	**Good identification**					**17473121**
Actinomyces naeslundii		100.0	0.36	IND 1		
17473123	**Good identification**					**17473123**
Actinomyces naeslundii		100.0	0.36	IND 1		
17475122	**Low discrimination**					**17475122**
Parabacteroides distasonis		84.75	0.37	IND 1		
Bacteroides thetaiotaomicron		15.25	0.08			
17475322	**Low discrimination**					**17475322**
Parabacteroides distasonis		53.87	0.45	IND 1		
Bacteroides thetaiotaomicron		42.93	0.27			
Bacteroides ovatus		3.20	0.08			
17475332	**Low discrimination**					**17475332**
Parabacteroides distasonis		60.65	0.40	IND 1		
Bacteroides thetaiotaomicron		34.90	0.20			
Bacteroides ovatus		4.45	0.05			
17475522	**Acceptable identification**					**17475522**
Bacteroides thetaiotaomicron		94.94	0.42	ESL 92		
Parabacteroides distasonis		3.27	0.33	MLT 94		
Bacteroides ovatus		1.61	0.12	RHA 93		
17475532	**Acceptable identification**					**17475532**
Bacteroides thetaiotaomicron		92.87	0.35	ESL 92		
Parabacteroides distasonis		4.42	0.29	MLT 94		
Bacteroides ovatus		2.71	0.09	RHA 93		
17475722	**Very good identification**					**17475722**
Bacteroides thetaiotaomicron		96.87	0.61	ESL 92		
Bacteroides ovatus		2.36	0.34	MLT 94		
Parabacteroides distasonis		0.75	0.41			
17475732	**Acceptable identification**					**17475732**
Bacteroides thetaiotaomicron		94.97	0.54	ESL 92		
Bacteroides ovatus		3.96	0.31	MLT 94		
Parabacteroides distasonis		1.02	0.37			
17500000	**Very good identification**					**17500000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Prevotella intermedia		97.19	0.61	FRU 99	1	1	1	
Gemella morbillorum		1.45	0.30		99	99	1	
Clostridium bifermentans		1.20	0.25		1	99	99	
17501000	**Very good identification**							**17501000**
					COC	GRA		
Prevotella intermedia		97.97	0.61	FRU 99	1	1		
Eubacterium saburreum		1.07	0.37		1	85		
Gemella morbillorum		0.79	0.26		99	99		
17501100	**Low discrimination**							**17501100**
Eubacterium saburreum		59.61	0.43	ESL 99				
Prevotella intermedia		23.65	0.26					
Prevotella melaninogenica		12.72	0.15					
17501110	**Good identification**							**17501110**
Eubacterium saburreum		100.0	0.41	ESL 99				
17501120	**Good identification**							**17501120**
					GRA			
Prevotella melaninogenica		98.74	0.50	IND 1	1			
Eubacterium saburreum		0.83	0.30	FRU 90	85			
Clostridium perfringens		0.36	0.15		99			
17501122	**Good identification**							**17501122**
					GRA			
Prevotella melaninogenica		97.16	0.37	IND 1	1			
Clostridium perfringens		0.89	0.09	FRU 90	99			
Prevotella oralis		0.81	0.05		1			
17501520	**Low discrimination**							**17501520**
					GRA			
Bifidobacterium longum subsp. longum		78.00	0.37	IND 1	99			
Prevotella melaninogenica		20.44	0.15	FRU 90	1			
Lactobacillus fermentum		1.56	0.15		99			
17501522	**Acceptable identification**							**17501522**
					GRA			
Bifidobacterium longum subsp. longum		93.47	0.37	IND 1	99			
Prevotella melaninogenica		4.67	0.03	FRU 90	1			
Lactobacillus fermentum		1.87	0.15		99			
17501530	**Good identification**							**17501530**
Bifidobacterium longum subsp. longum		100.0	0.41	IND 1				
17501532	**Good identification**							**17501532**
Bifidobacterium longum subsp. longum		100.0	0.41	IND 1				
17505100	**Acceptable identification**							**17505100**
Eubacterium saburreum		94.95	0.38	ESL 99				
Lactobacillus fermentum		5.05	0.20					
17505110	**Good identification**							**17505110**
Eubacterium saburreum		100.0	0.37	ESL 99				
17505500	**Good identification**							**17505500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		95.52	0.35	IND 1				
Lactobacillus fermentum		4.48	0.20	FRU 90				
17505502	**Good identification**							**17505502**
Bifidobacterium longum subsp. longum		95.52	0.35	IND 1				
Lactobacillus fermentum		4.48	0.20	FRU 90				
17505510	**Good identification**							**17505510**
Bifidobacterium longum subsp. longum		100.0	0.39	IND 1				
17505512	**Good identification**							**17505512**
Bifidobacterium longum subsp. longum		100.0	0.39	IND 1				
17505520	**Good identification**							**17505520**
					GRA			
Bifidobacterium longum subsp. longum		95.23	0.46	IND 1	99			
Bacteroides vulgatus		4.77	0.09	FRU 90	1			
17505522	**Good identification**							**17505522**
					GRA	CAT		
Bifidobacterium longum subsp. longum		98.27	0.46	IND 1	99	1		
Bacteroides thetaiotaomicron		1.73	0.04	FRU 90	1	89		
17505530	**Good identification**							**17505530**
Bifidobacterium longum subsp. longum		100.0	0.50	IND 1				
17505532	**Good identification**							**17505532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		99.60	0.50	IND 1	99	1		
Bacteroides thetaiotaomicron		0.40	-.03	FRU 90	1	89		
17510000	**Excellent identification**							**17510000**
					COC	GRA	CAT	
Prevotella intermedia		99.87	0.95		1	1	1	
Propionibacterium granulosum		0.11	0.52		1	99	90	
Gemella morbillorum		0.02	0.32		99	99	1	
17510001	**Very good identification**							**17510001**
					GRA	SPO	CAT	
Prevotella intermedia		99.68	0.61	URE 1	1	1	1	
Clostridium sordellii		0.22	0.08		99	99	1	
Propionibacterium granulosum		0.11	0.18		99	1	90	
17510002	**Very good identification**							**17510002**
					GRA	CAT		
Prevotella intermedia		96.00	0.61	BGL 1	1	1		
Propionibacterium granulosum		2.53	0.41		99	90		
Actinomyces viscosus		1.34	0.27		99	76		
17510010	**Very good identification**							**17510010**
					GRA	CAT		
Prevotella intermedia		98.58	0.61	MLZ 1	1	1		
Propionibacterium granulosum		1.42	0.37		99	90		
17510020	**Very good identification**							**17510020**
					GRA	SPO	CAT	
Prevotella intermedia		96.25	0.61	NAG 1	1	1	1	
Propionibacterium granulosum		2.54	0.41		99	1	90	
Clostridium cadaveris		0.97	0.22		99	99	1	
17510040	**Very good identification**							**17510040**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Prevotella intermedia		98.61	0.61	SOR 1	1	1	1	
Clostridium botulinum biovar B		1.28	0.24		99	99	1	
Propionibacterium granulosum		0.11	0.18		99	1	90	
17510100	**Very good identification**							**17510100**
					GRA			
Prevotella intermedia		98.39	0.61	LAC 1	1			
Fusobacterium mortiferum		1.26	0.39		1			
Lactobacillus fermentum		0.12	0.24		99			
17510200	**Very good identification**							**17510200**
					GRA	CAT		
Prevotella intermedia		99.68	0.61	RHA 1	1	1		
Propionibacterium granulosum		0.33	0.26		99	90		
17510400	**Very good identification**							**17510400**
					COC	GRA	CAT	
Prevotella intermedia		99.62	0.61	ARA 1	1	1	1	
Gemella morbillorum		0.20	0.15		99	99	1	
Propionibacterium granulosum		0.11	0.18		1	99	90	
17511000	**Excellent identification**							**17511000**
					GRA	CAT		
Prevotella intermedia		99.76	0.95		1	1		
Propionibacterium granulosum		0.11	0.52		99	90		
Lactobacillus fermentum		0.08	0.54		99	1		
17511001	**Very good identification**							**17511001**
					GRA	CAT		
Prevotella intermedia		99.56	0.61	URE 1	1	1		
Actinomyces naeslundii		0.19	0.37		99	25		
Propionibacterium granulosum		0.11	0.18		99	90		
17511002	**Low discrimination**							**17511002**
					GRA	CAT		
Prevotella intermedia		89.38	0.61	BGL 1	1	1		
Lactobacillus fermentum		6.73	0.54		99	1		
Propionibacterium granulosum		2.35	0.41		99	90		
17511010	**Very good identification**							**17511010**
					CAT	GRA		
Prevotella intermedia		95.98	0.61	MLZ 1	1	1		
Eubacterium saburreum		2.57	0.43		1	85		
Propionibacterium granulosum		1.38	0.37		90	99		
17511020	**Low discrimination**							**17511020**
					GRA			
Prevotella intermedia		88.06	0.61	NAG 1	1			
Prevotella melaninogenica		4.31	0.32		1			
Clostridium perfringens		2.36	0.34		99			
17511022	**Low discrimination**							**17511022**
					GRA			
Lactobacillus fermentum		35.17	0.46	IND 1	99			
Clostridium perfringens		17.66	0.29		99			
Prevotella intermedia		14.15	0.26		1			
17511023	**Good identification**							**17511023**
Actinomyces naeslundii		100.0	0.37	IND 1				
17511040	**Very good identification**							**17511040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Prevotella intermedia		99.79	0.61	SOR 1	1	1	
Propionibacterium granulosum		0.11	0.18		99	90	
Lactobacillus fermentum		0.08	0.20		99	1	
17511100	**Low discrimination**						**17511100**
					GRA		
Prevotella intermedia		78.21	0.61	LAC 1	1		
Lactobacillus fermentum		9.61	0.58		99		
Eubacterium saburreum		5.97	0.51		85		
17511102	**Low discrimination**						**17511102**
Lactobacillus fermentum		65.54	0.58	IND 1			
Actinomyces viscosus		8.11	0.28				
Eubacterium saburreum		7.19	0.38				
17511103	**Low discrimination**						**17511103**
					SPO		
Actinomyces naeslundii		40.26	0.45	IND 1	1		
Actinomyces viscosus		37.92	0.20		1		
Clostridium chauvoei		12.54	0.13		99		
17511110	**Low discrimination**						**17511110**
					GRA		
Eubacterium saburreum		83.18	0.49	ESL 99	85		
Prevotella intermedia		13.45	0.26		1		
Lactobacillus fermentum		1.65	0.24		99		
17511112	**Low discrimination**						**17511112**
Eubacterium saburreum		82.13	0.37	ESL 99			
Lactobacillus fermentum		9.25	0.24				
Bifidobacterium longum subsp. longum		5.97	0.13				
17511120	**Good identification**						**17511120**
					GRA		
Prevotella melaninogenica		94.65	0.66	IND 1	1		
Clostridium perfringens		3.84	0.49		99		
Lactobacillus fermentum		0.80	0.50		99		
17511122	**Low discrimination**						**17511122**
					GRA		
Prevotella melaninogenica		78.76	0.54	IND 1	1		
Clostridium perfringens		7.90	0.43		99		
Prevotella oralis		7.23	0.39		1		
17511200	**Very good identification**						**17511200**
					GRA	CAT	
Prevotella intermedia		99.57	0.61	RHA 1	1	1	
Propionibacterium granulosum		0.32	0.26		99	90	
Lactobacillus fermentum		0.08	0.20		99	1	
17511400	**Good identification**						**17511400**
					GRA	CAT	
Prevotella intermedia		92.74	0.61	ARA 1	1	1	
Lactobacillus fermentum		6.99	0.54		99	1	
Propionibacterium granulosum		0.10	0.18		99	90	
17511402	**Low discrimination**						**17511402**
					GRA		
Lactobacillus fermentum		87.69	0.54	IND 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		11.76	0.26		1		
Bifidobacterium longum subsp. longum		0.55	0.09		99		
17511420	**Low discrimination**						**17511420**
					GRA		
Lactobacillus fermentum		66.80	0.46	IND 1	99		
Prevotella intermedia		26.87	0.26		1		
Bifidobacterium longum subsp. longum		5.01	0.20		99		
17511422	**Acceptable identification**						**17511422**
Lactobacillus fermentum		93.02	0.46	IND 1			
Bifidobacterium longum subsp. longum		6.98	0.20				
17511500	**Low discrimination**						**17511500**
					GRA		
Lactobacillus fermentum		67.94	0.58	IND 1	99		
Bifidobacterium longum subsp. longum		25.78	0.43		99		
Prevotella intermedia		5.58	0.26		1		
17511502	**Low discrimination**						**17511502**
Lactobacillus fermentum		71.51	0.58	IND 1			
Bifidobacterium longum subsp. longum		27.13	0.43				
Bifidobacterium adolescentis		1.13	0.21				
17511510	**Good identification**						**17511510**
Bifidobacterium longum subsp. longum		97.70	0.47	IND 1			
Lactobacillus fermentum		1.53	0.24				
Eubacterium saburreum		0.78	0.15				
17511512	**Good identification**						**17511512**
Bifidobacterium longum subsp. longum		95.64	0.47	IND 1			
Bifidobacterium adolescentis		2.86	0.23				
Lactobacillus fermentum		1.50	0.24				
17511520	**Low discrimination**						**17511520**
					GRA		
Bifidobacterium longum subsp. longum		65.44	0.54	IND 1	99		
Prevotella melaninogenica		17.15	0.32		1		
Lactobacillus fermentum		14.37	0.50		99		
17511522	**Low discrimination**						**17511522**
					GRA		
Bifidobacterium longum subsp. longum		75.32	0.54	IND 1	99		
Lactobacillus fermentum		16.55	0.50		99		
Prevotella melaninogenica		3.76	0.19		1		
17511530	**Very good identification**						**17511530**
					GRA		
Bifidobacterium longum subsp. longum		99.72	0.58	IND 1	99		
Prevotella melaninogenica		0.16	-.03		1		
Lactobacillus fermentum		0.13	0.15		99		
17511532	**Very good identification**						**17511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.02	0.58	IND 1	99	1	
Bacteroides thetaiotaomicron		0.55	0.07		1	89	
Parabacteroides distasonis		0.18	0.15		1	76	
17512000	**Good identification**						**17512000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Prevotella intermedia		92.64	0.61	MAN 1	1	1	
Propionibacterium granulosum		7.36	0.50		99	90	
17512002	**Low discrimination**						**17512002**
					GRA	CAT	
Propionibacterium granulosum		65.69	0.39	IND 1	99	90	
Prevotella intermedia		33.40	0.26		1	1	
Propionibacterium propionicum		0.91	0.12		99	1	
17512020	**Low discrimination**						**17512020**
					GRA	CAT	
Propionibacterium granulosum		66.29	0.39	IND 1	99	90	
Prevotella intermedia		33.71	0.26		1	1	
17512100	**Low discrimination**						**17512100**
					GRA	CAT	
Propionibacterium propionicum		61.94	0.43	IND 1	99	1	
Prevotella intermedia		34.34	0.26	NIT 86	1	1	
Propionibacterium granulosum		2.73	0.16		99	90	
17512102	**Good identification**						**17512102**
Propionibacterium propionicum		98.95	0.46	IND 1			
Actinomyces naeslundii		1.05	0.23	NIT 86			
17512140	**Good identification**						**17512140**
Propionibacterium propionicum		100.0	0.43	IND 1			
17512142	**Good identification**						**17512142**
Propionibacterium propionicum		100.0	0.46	IND 1			
17513000	**Good identification**						**17513000**
					GRA	CAT	
Prevotella intermedia		92.34	0.61	MAN 1	1	1	
Propionibacterium granulosum		7.34	0.50		99	90	
Actinomyces naeslundii		0.23	0.39		99	25	
17513002	**Low discrimination**						**17513002**
					GRA	CAT	
Propionibacterium granulosum		59.39	0.39	IND 1	99	90	
Prevotella intermedia		30.20	0.26		1	1	
Actinomyces naeslundii		7.32	0.39		99	25	
17513003	**Good identification**						**17513003**
Actinomyces naeslundii		100.0	0.42	IND 1			
17513020	**Low discrimination**						**17513020**
					GRA	CAT	
Propionibacterium granulosum		60.37	0.39	IND 1	99	90	
Prevotella intermedia		30.69	0.26		1	1	
Actinomyces naeslundii		7.44	0.39		99	25	
17513021	**Good identification**						**17513021**
Actinomyces naeslundii		100.0	0.42	IND 1			
17513023	**Good identification**						**17513023**
Actinomyces naeslundii		100.0	0.42	IND 1			
17513100	**Low discrimination**						**17513100**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Propionibacterium propionicum		47.20	0.43	IND 1	99			
Prevotella intermedia		26.17	0.26	NIT 86	1			
Actinomyces naeslundii		18.05	0.46		99			
17513101	**Low discrimination**							**17513101**
Actinomyces naeslundii		83.77	0.49	IND 1				
Propionibacterium propionicum		16.23	0.27					
17513102	**Low discrimination**							**17513102**
Propionibacterium propionicum		76.90	0.46	IND 1				
Actinomyces naeslundii		19.61	0.46	NIT 86				
Lactobacillus fermentum		3.50	0.24					
17513103	**Low discrimination**							**17513103**
Actinomyces naeslundii		77.49	0.49	IND 1				
Propionibacterium propionicum		22.51	0.30					
17513121	**Good identification**							**17513121**
					GRA			
Actinomyces naeslundii		95.49	0.49	IND 1	99			
Prevotella melaninogenica		4.51	-.03		1			
17513123	**Good identification**							**17513123**
Actinomyces naeslundii		100.0	0.49	IND 1				
17513140	**Acceptable identification**							**17513140**
Propionibacterium propionicum		93.68	0.43	IND 1				
Actinomyces naeslundii		6.32	0.34	NIT 86				
17513142	**Good identification**							**17513142**
Propionibacterium propionicum		95.69	0.46	IND 1				
Actinomyces naeslundii		4.31	0.34	NIT 86				
17513161	**Good identification**							**17513161**
Actinomyces naeslundii		100.0	0.37	IND 1				
17513163	**Good identification**							**17513163**
Actinomyces naeslundii		100.0	0.37	IND 1				
17514000	**Very good identification**							**17514000**
					COC	GRA	CAT	
Prevotella intermedia		99.61	0.61	XYL 1	1	1	1	
Gemella morbillorum		0.20	0.15		99	99	1	
Propionibacterium granulosum		0.11	0.18		1	99	90	
17515000	**Good identification**							**17515000**
					GRA			
Prevotella intermedia		93.93	0.61	XYL 1	1			
Lactobacillus fermentum		4.16	0.50		99			
Eubacterium saburreum		1.66	0.40		85			
17515002	**Low discrimination**							**17515002**
					GRA			
Lactobacillus fermentum		77.01	0.50	IND 1	99			
Prevotella intermedia		17.58	0.26		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		5.41	0.27		85		
17515010	**Low discrimination**						**17515010**
					GRA		
Eubacterium saburreum		57.76	0.39	ESL 99	85		
Prevotella intermedia		40.46	0.26		1		
Lactobacillus fermentum		1.79	0.16		99		
17515020	**Low discrimination**						**17515020**
					GRA		
Lactobacillus fermentum		51.83	0.42	IND 1	99		
Prevotella intermedia		35.50	0.26		1		
Eubacterium saburreum		10.93	0.27		85		
17515022	**Low discrimination**						**17515022**
					CAT	GRA	
Lactobacillus fermentum		86.79	0.42	IND 1	1	99	
Bacteroides fragilis		6.05	-.03		99	1	
Prevotella oralis		3.93	0.03		1	1	
17515100	**Low discrimination**						**17515100**
					GRA		
Lactobacillus fermentum		56.95	0.54	IND 1	99		
Eubacterium saburreum		32.44	0.46		85		
Prevotella intermedia		7.97	0.26		1		
17515102	**Low discrimination**						**17515102**
Lactobacillus fermentum		85.00	0.54	IND 1			
Eubacterium saburreum		8.55	0.33				
Bifidobacterium adolescentis		2.29	0.21				
17515110	**Acceptable identification**						**17515110**
Eubacterium saburreum		91.48	0.45	ESL 99			
Bifidobacterium longum subsp. longum		6.54	0.21				
Lactobacillus fermentum		1.98	0.20				
17515320	**Good identification**						**17515320**
Bacteroides vulgatus		99.54	0.40	IND 1			
Prevotella melaninogenica		0.24	-.03	ARA 93			
Parabacteroides distasonis		0.23	0.14				
17515322	**Low discrimination**						**17515322**
					CAT		
Parabacteroides distasonis		67.68	0.48	IND 1	76		
Bacteroides thetaiotaomicron		20.71	0.23		89		
Prevotella oralis		6.39	0.17		1		
17515332	**Low discrimination**						**17515332**
Parabacteroides distasonis		81.90	0.43	IND 1			
Bacteroides thetaiotaomicron		18.10	0.16				
17515400	**Low discrimination**						**17515400**
					GRA		
Lactobacillus fermentum		79.35	0.50	IND 1	99		
Prevotella intermedia		18.12	0.26		1		
Bifidobacterium longum subsp. longum		2.53	0.17		99		
17515402	**Good identification**						**17515402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		96.91	0.50	IND 1			
Bifidobacterium longum subsp. longum		3.10	0.17				
17515420	**Low discrimination**						**17515420**
					GRA		
Lactobacillus fermentum		62.72	0.42	IND 1	99		
Bifidobacterium longum subsp. longum		24.04	0.28		99		
Bacteroides vulgatus		13.24	0.09		1		
17515422	**Low discrimination**						**17515422**
					GRA	CAT	
Lactobacillus fermentum		65.23	0.42	IND 1	99	1	
Bifidobacterium longum subsp. longum		25.00	0.28		99	1	
Bacteroides thetaiotaomicron		9.77	0.09		1	89	
17515500	**Low discrimination**						**17515500**
					GRA		
Bifidobacterium longum subsp. longum		64.88	0.52	IND 1	99		
Lactobacillus fermentum		33.48	0.54		99		
Bacteroides vulgatus		1.44	0.09		1		
17515502	**Low discrimination**						**17515502**
Bifidobacterium longum subsp. longum		61.85	0.52	IND 1			
Lactobacillus fermentum		31.91	0.54				
Bifidobacterium adolescentis		5.75	0.36				
17515510	**Good identification**						**17515510**
Bifidobacterium longum subsp. longum		99.55	0.56	IND 1			
Lactobacillus fermentum		0.31	0.20				
Eubacterium saburreum		0.14	0.11				
17515512	**Acceptable identification**						**17515512**
Bifidobacterium longum subsp. longum		93.32	0.56	IND 1			
Bifidobacterium adolescentis		6.23	0.37				
Lactobacillus fermentum		0.29	0.20				
17515520	**Low discrimination**						**17515520**
					GRA		
Bifidobacterium longum subsp. longum		62.63	0.62	IND 1	99		
Bacteroides vulgatus		34.50	0.43		1		
Lactobacillus fermentum		2.69	0.46		99		
17515522	**Low discrimination**						**17515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		78.68	0.62	IND 1	99	1	
Bacteroides thetaiotaomicron		15.21	0.38		1	89	
Lactobacillus fermentum		3.38	0.46		99	1	
17515530	**Very good identification**						**17515530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.63	0.66	IND 1	99	1	
Bacteroides vulgatus		0.33	0.09		1	3	
Bacteroides thetaiotaomicron		0.04	-.03		1	89	
17515532	**Very good identification**						**17515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		95.27	0.66	IND 1	99	1	
Bacteroides thetaiotaomicron		4.21	0.31		1	89	
Parabacteroides distasonis		0.52	0.31		1	76	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17515720	**Very good identification**						**17515720**
					GRA	CAT	
Bacteroides vulgatus		99.27	0.60	IND 1	1	3	
Bacteroides thetaiotaomicron		0.52	0.23		1	89	
Bifidobacterium longum subsp. longum		0.20	0.28		99	1	
17515722	**Very good identification**						**17515722**
					CAT		
Bacteroides thetaiotaomicron		95.01	0.57	ESL 92	89		
Parabacteroides distasonis		1.92	0.44	TRE 93	76		
Bacteroides vulgatus		1.85	0.25		3		
17515732	**Acceptable identification**						**17515732**
					GRA	CAT	
Bacteroides thetaiotaomicron		94.64	0.50	ESL 92	1	89	
Parabacteroides distasonis		2.65	0.39	TRE 93	1	76	
Bifidobacterium longum subsp. longum		1.63	0.32		99	1	
17517512	**Low discrimination**						**17517512**
Bifidobacterium adolescentis		84.40	0.36	IND 1			
Bifidobacterium longum subsp. longum		15.60	0.21				
17521100	**Good identification**						**17521100**
Eubacterium saburreum		97.89	0.40	ESL 99			
Actinomyces naeslundii		2.12	0.25				
17521110	**Good identification**						**17521110**
Eubacterium saburreum		100.0	0.38	ESL 99			
17525100	**Good identification**						**17525100**
Eubacterium saburreum		100.0	0.35	ESL 99			
17525520	**Good identification**						**17525520**
Bifidobacterium longum subsp. longum		100.0	0.35	IND 1			
17525522	**Low discrimination**						**17525522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		50.95	0.35	IND 1	99	1	
Bacteroides thetaiotaomicron		47.59	0.23	FRU 90	1	89	
Parabacteroides distasonis		1.46	0.13		1	76	
17525530	**Good identification**						**17525530**
Bifidobacterium longum subsp. longum		100.0	0.39	IND 1			
17525532	**Low discrimination**						**17525532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		82.42	0.39	IND 1	99	1	
Bacteroides thetaiotaomicron		17.58	0.16	FRU 90	1	89	
17525722	**Good identification**						**17525722**
Bacteroides thetaiotaomicron		98.99	0.42	ESL 92			
Parabacteroides distasonis		0.69	0.21	FRU 99			
Bacteroides ovatus		0.33	0.00				
17525732	**Good identification**						**17525732**
Bacteroides thetaiotaomicron		98.50	0.35	ESL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		0.95	0.17	FRU 99				
Bacteroides ovatus		0.56	-.03					
17530000	**Low discrimination**							**17530000**
					GRA	CAT		
Prevotella intermedia		75.87	0.61	TRE 1	1	1		
Propionibacterium granulosum		23.98	0.60		99	90		
Fusobacterium mortiferum		0.13	0.24		1	5		
17530002	**Low discrimination**							**17530002**
					GRA	CAT		
Propionibacterium granulosum		87.63	0.50	IND 1	99	90		
Prevotella intermedia		11.20	0.26		1	1		
Actinomyces viscosus		1.17	0.08		99	76		
17530010	**Low discrimination**							**17530010**
					GRA	CAT		
Propionibacterium granulosum		81.01	0.45	IND 1	99	90		
Prevotella intermedia		18.99	0.26	MLZ 12	1	1		
17530020	**Low discrimination**							**17530020**
					GRA	SPO	CAT	
Propionibacterium granulosum		87.65	0.50	IND 1	99	1	90	
Prevotella intermedia		11.21	0.26		1	1	1	
Clostridium cadaveris		0.59	0.00		99	99	1	
17530022	**Good identification**							**17530022**
Propionibacterium granulosum		100.0	0.39	IND 1				
17531000	**Low discrimination**							**17531000**
					CAT	GRA		
Prevotella intermedia		74.35	0.61	TRE 1	1	1		
Propionibacterium granulosum		23.50	0.60		90	99		
Eubacterium saburreum		1.62	0.42		1	85		
17531002	**Low discrimination**							**17531002**
					CAT	GRA		
Propionibacterium granulosum		80.78	0.50	IND 1	90	99		
Prevotella intermedia		10.33	0.26		1	1		
Eubacterium saburreum		3.94	0.29		1	85		
17531003	**Low discrimination**							**17531003**
Actinomyces naeslundii		82.15	0.42	IND 1				
Propionibacterium granulosum		17.86	0.15					
17531010	**Low discrimination**							**17531010**
					CAT	GRA		
Propionibacterium granulosum		60.65	0.45	IND 1	90	99		
Eubacterium saburreum		25.13	0.40	MLZ 12	1	85		
Prevotella intermedia		14.22	0.26		1	1		
17531020	**Low discrimination**							**17531020**
					GRA	CAT		
Propionibacterium granulosum		54.24	0.50	IND 1	99	90		
Clostridium perfringens		34.16	0.39		99	1		
Prevotella intermedia		6.94	0.26		1	1		
17531021	**Low discrimination**							**17531021**
					CAT			
Actinomyces naeslundii		73.84	0.42	IND 1	25			
Propionibacterium granulosum		16.05	0.15		90			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		10.11	0.05		1	
17531023	**Good identification**					**17531023**
Actinomyces naeslundii		100.0	0.42	IND 1		
17531100	**Low discrimination**					**17531100**
					GRA	
Eubacterium saburreum		55.72	0.48	ESL 99	85	
Clostridium perfringens		12.35	0.28		99	
Prevotella intermedia		11.05	0.26		1	
17531101	**Low discrimination**					**17531101**
					SPO	
Actinomyces naeslundii		86.94	0.49	IND 1	1	
Clostridium chauvoei		8.78	0.09		99	
Eubacterium saburreum		4.28	0.14		1	
17531102	**Low discrimination**					**17531102**
Eubacterium saburreum		35.42	0.35	ESL 99		
Actinomyces naeslundii		27.46	0.46			
Clostridium perfringens		20.93	0.22			
17531103	**Good identification**					**17531103**
Actinomyces naeslundii		96.48	0.49	IND 1		
Actinomyces viscosus		3.52	0.01			
17531110	**Good identification**					**17531110**
					CAT	
Eubacterium saburreum		98.11	0.46	ESL 99	1	
Propionibacterium granulosum		1.03	0.11		90	
Actinomyces naeslundii		0.86	0.25		25	
17531120	**Low discrimination**					**17531120**
					GRA	
Clostridium perfringens		84.52	0.54	IND 1	99	
Prevotella melaninogenica		11.33	0.32		1	
Eubacterium saburreum		2.08	0.35		85	
17531121	**Low discrimination**					**17531121**
					GRA	
Actinomyces naeslundii		69.76	0.49	IND 1	99	
Clostridium perfringens		24.60	0.19		99	
Prevotella melaninogenica		3.30	-.03		1	
17531122	**Low discrimination**					**17531122**
					GRA	
Clostridium perfringens		83.98	0.48	IND 1	99	
Prevotella melaninogenica		4.56	0.19		1	
Parabacteroides distasonis		3.92	0.34		1	
17531123	**Low discrimination**					**17531123**
Actinomyces naeslundii		85.77	0.49	IND 1		
Clostridium perfringens		14.23	0.14			
17531141	**Good identification**					**17531141**
Actinomyces naeslundii		100.0	0.37	IND 1		
17531143	**Good identification**					**17531143**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	IND 1			
17531160	**Good identification**						**17531160**
					GRA		
Clostridium perfringens		97.58	0.41	IND 1	99		
Actinomyces naeslundii		1.73	0.34		99		
Prevotella melaninogenica		0.69	-.03		1		
17531161	**Low discrimination**						**17531161**
Actinomyces naeslundii		72.43	0.37	IND 1			
Clostridium perfringens		27.57	0.07				
17531162	**Good identification**						**17531162**
Clostridium perfringens		96.38	0.36	IND 1			
Actinomyces naeslundii		3.62	0.34				
17531163	**Low discrimination**						**17531163**
Actinomyces naeslundii		84.81	0.37	IND 1			
Clostridium perfringens		15.19	0.01				
17531322	**Low discrimination**						**17531322**
					CAT		
Parabacteroides distasonis		66.53	0.43	IND 1	76		
Bacteroides thetaiotaomicron		22.26	0.19	XYL 90	89		
Prevotella oralis		6.41	0.13		1		
17531332	**Low discrimination**						**17531332**
Parabacteroides distasonis		80.54	0.38	IND 1			
Bacteroides thetaiotaomicron		19.46	0.11	XYL 90			
17531510	**Good identification**						**17531510**
Bifidobacterium longum subsp. longum		97.92	0.37	IND 1			
Eubacterium saburreum		2.08	0.12				
17531512	**Acceptable identification**						**17531512**
Bifidobacterium longum subsp. longum		92.61	0.37	IND 1			
Bifidobacterium adolescentis		7.39	0.20				
17531520	**Low discrimination**						**17531520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.97	0.43	IND 1	99	1	
Clostridium perfringens		7.11	0.19		99	1	
Bacteroides thetaiotaomicron		1.17	-.01		1	89	
17531530	**Good identification**						**17531530**
Bifidobacterium longum subsp. longum		100.0	0.47	IND 1			
17531532	**Low discrimination**						**17531532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		75.38	0.47	IND 1	99	1	
Bacteroides thetaiotaomicron		22.11	0.26		1	89	
Parabacteroides distasonis		2.51	0.26		1	76	
17531722	**Good identification**						**17531722**
Bacteroides thetaiotaomicron		98.01	0.53	ESL 92			
Parabacteroides distasonis		1.81	0.39	XYL 96			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Bacteroides ovatus	0.08	0.00				
17531732	**Good identification**						**17531732**
	Bacteroides thetaiotaomicron	96.70	0.46	ESL 92			
	Parabacteroides distasonis	2.48	0.34	XYL 96			
	Clostridium clostridioforme	0.44	0.16				
17532000	**Very good identification**						**17532000**
					GRA	CAT	
	Propionibacterium granulosum	95.94	0.58	IND 1	99	90	
	Prevotella intermedia	4.07	0.26		1	1	
17532002	**Good identification**						**17532002**
	Propionibacterium granulosum	100.0	0.48	IND 1			
17532010	**Good identification**						**17532010**
	Propionibacterium granulosum	100.0	0.43	IND 1			
17532020	**Good identification**						**17532020**
	Propionibacterium granulosum	100.0	0.48	IND 1			
17532022	**Good identification**						**17532022**
	Propionibacterium granulosum	100.0	0.37	IND 1			
17532100	**Low discrimination**						**17532100**
					CAT		
	Propionibacterium propionicum	83.38	0.42	IND 1	1		
	Propionibacterium granulosum	13.47	0.24	NIT 86	90		
	Actinomyces naeslundii	3.15	0.28		25		
17532102	**Good identification**						**17532102**
					CAT		
	Propionibacterium propionicum	95.05	0.45	IND 1	1		
	Propionibacterium granulosum	2.56	0.13	NIT 86	90		
	Actinomyces naeslundii	2.40	0.28		25		
17532140	**Good identification**						**17532140**
	Propionibacterium propionicum	100.0	0.42	IND 1			
17532142	**Good identification**						**17532142**
	Propionibacterium propionicum	100.0	0.45	IND 1			
17533000	**Good identification**						**17533000**
					GRA	CAT	
	Propionibacterium granulosum	94.14	0.58	IND 1	99	90	
	Prevotella intermedia	3.99	0.26		1	1	
	Actinomyces naeslundii	1.88	0.44		99	25	
17533001	**Low discrimination**						**17533001**
	Actinomyces naeslundii	74.75	0.47	IND 1			
	Propionibacterium granulosum	25.26	0.24				
17533002	**Acceptable identification**						**17533002**
	Propionibacterium granulosum	92.62	0.48	IND 1			
	Actinomyces naeslundii	7.38	0.44				
17533003	**Acceptable identification**						**17533003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.21	0.47	IND 1		
Propionibacterium granulosum		7.79	0.13			
17533010	**Good identification**					**17533010**
Propionibacterium granulosum		100.0	0.43	IND 1		
17533020	**Acceptable identification**					**17533020**
					CAT	
Propionibacterium granulosum		91.90	0.48	IND 1	90	
Actinomyces naeslundii		7.33	0.44		25	
Clostridium perfringens		0.78	0.05		1	
17533021	**Acceptable identification**					**17533021**
Actinomyces naeslundii		92.21	0.47	IND 1		
Propionibacterium granulosum		7.79	0.13			
17533022	**Low discrimination**					**17533022**
Propionibacterium granulosum		75.82	0.37	IND 1		
Actinomyces naeslundii		24.18	0.44			
17533023	**Good identification**					**17533023**
Actinomyces naeslundii		100.0	0.47	IND 1		
17533100	**Low discrimination**					**17533100**
					CAT	
Propionibacterium propionicum		47.54	0.42	IND 1	1	
Actinomyces naeslundii		43.14	0.51	NIT 86	25	
Propionibacterium granulosum		7.68	0.24		90	
17533101	**Acceptable identification**					**17533101**
Actinomyces naeslundii		92.45	0.54	IND 1		
Propionibacterium propionicum		7.55	0.26			
17533102	**Low discrimination**					**17533102**
					CAT	
Propionibacterium propionicum		61.28	0.45	IND 1	1	
Actinomyces naeslundii		37.07	0.51	NIT 86	25	
Propionibacterium granulosum		1.65	0.13		90	
17533103	**Low discrimination**					**17533103**
Actinomyces naeslundii		89.09	0.54	IND 1		
Propionibacterium propionicum		10.91	0.29			
17533120	**Low discrimination**					**17533120**
					CAT	
Actinomyces naeslundii		73.88	0.51	IND 1	25	
Clostridium perfringens		20.13	0.19		1	
Propionibacterium granulosum		3.29	0.13		90	
17533121	**Good identification**					**17533121**
Actinomyces naeslundii		100.0	0.54	IND 1		
17533122	**Low discrimination**					**17533122**
Actinomyces naeslundii		88.63	0.51	IND 1		
Clostridium perfringens		11.37	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17533123	**Good identification**						**17533123**
Actinomyces naeslundii		100.0	0.54	IND 1			
17533140	**Low discrimination**						**17533140**
Propionibacterium propionicum		86.20	0.42	IND 1			
Actinomyces naeslundii		13.80	0.38	NIT 86			
17533141	**Low discrimination**						**17533141**
Actinomyces naeslundii		68.37	0.41	IND 1			
Propionibacterium propionicum		31.63	0.26				
17533142	**Acceptable identification**						**17533142**
Propionibacterium propionicum		90.36	0.45	IND 1			
Actinomyces naeslundii		9.65	0.38	NIT 86			
17533143	**Low discrimination**						**17533143**
Actinomyces naeslundii		59.04	0.41	IND 1			
Propionibacterium propionicum		40.97	0.29				
17533160	**Low discrimination**						**17533160**
Actinomyces naeslundii		77.27	0.38	IND 1			
Clostridium perfringens		22.73	0.07				
17533161	**Good identification**						**17533161**
Actinomyces naeslundii		100.0	0.41	IND 1			
17533162	**Low discrimination**						**17533162**
Actinomyces naeslundii		87.84	0.38	IND 1			
Clostridium perfringens		12.16	0.01				
17533163	**Good identification**						**17533163**
Actinomyces naeslundii		100.0	0.41	IND 1			
17534722	**Good identification**						**17534722**
Bacteroides thetaiotaomicron		98.46	0.42	ESL 92			
Parabacteroides distasonis		0.68	0.21	GAL 99			
Clostridium clostridioforme		0.53	0.14				
17534732	**Acceptable identification**						**17534732**
Bacteroides thetaiotaomicron		92.72	0.35	ESL 92			
Clostridium clostridioforme		5.86	0.25	GAL 99			
Parabacteroides distasonis		0.89	0.17				
17535000	**Low discrimination**						**17535000**
					CAT	GRA	
Eubacterium saburreum		46.10	0.37	ESL 99	1	85	
Prevotella intermedia		39.63	0.26		1	1	
Propionibacterium granulosum		12.52	0.26		90	99	
17535010	**Good identification**						**17535010**
					CAT		
Eubacterium saburreum		95.67	0.36	ESL 99	1		
Propionibacterium granulosum		4.33	0.11		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17535100	**Low discrimination**						**17535100**
					COC		
Eubacterium saburreum		86.74	0.43	ESL 99	1		
Atopobium parvulum		8.46	0.16		99		
Lactobacillus fermentum		2.31	0.20		1		
17535110	**Good identification**						**17535110**
Eubacterium saburreum		97.39	0.42	ESL 99			
Bifidobacterium longum subsp. longum		2.61	0.11				
17535122	**Low discrimination**						**17535122**
					GRA	CAT	
Parabacteroides distasonis		78.09	0.51	IND 1	1	76	
Bacteroides thetaiotaomicron		15.73	0.23		1	89	
Clostridium perfringens		1.88	0.14		99	1	
17535132	**Low discrimination**						**17535132**
					GRA	CAT	
Parabacteroides distasonis		81.60	0.46	IND 1	1	76	
Bacteroides thetaiotaomicron		11.87	0.16		1	89	
Bifidobacterium longum subsp. longum		5.06	0.21		99	1	
17535322	**Low discrimination**						**17535322**
Parabacteroides distasonis		52.56	0.59	IND 1			
Bacteroides thetaiotaomicron		46.90	0.42				
Bacteroides ovatus		0.47	0.08				
17535332	**Low discrimination**						**17535332**
Parabacteroides distasonis		60.12	0.54	IND 1			
Bacteroides thetaiotaomicron		38.74	0.35				
Bacteroides ovatus		0.67	0.05				
17535500	**Good identification**						**17535500**
Bifidobacterium longum subsp. longum		97.20	0.41	IND 1			
Lactobacillus fermentum		2.03	0.20				
Eubacterium saburreum		0.77	0.09				
17535502	**Low discrimination**						**17535502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		58.52	0.41	IND 1	99	1	
Bacteroides thetaiotaomicron		24.29	0.23		1	89	
Bifidobacterium adolescentis		14.52	0.33		99	1	
17535510	**Good identification**						**17535510**
Bifidobacterium longum subsp. longum		100.0	0.45	IND 1			
17535512	**Low discrimination**						**17535512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		76.60	0.45	IND 1	99	1	
Bifidobacterium adolescentis		13.64	0.34		99	1	
Bacteroides thetaiotaomicron		7.26	0.16		1	89	
17535520	**Low discrimination**						**17535520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.42	0.52	IND 1	99	1	
Bacteroides thetaiotaomicron		8.66	0.23		1	89	
Bacteroides vulgatus		7.66	0.20		1	3	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17535522	**Low discrimination**						**17535522**
					GRA	CAT	
Bacteroides thetaiotaomicron		88.44	0.57	ESL 92	1	89	
Bifidobacterium longum subsp. longum		8.61	0.52	RHA 93	99	1	
Parabacteroides distasonis		2.72	0.47		1	76	
17535530	**Good identification**						**17535530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.69	0.56	IND 1	99	1	
Bacteroides thetaiotaomicron		2.32	0.16		1	89	
17535532	**Low discrimination**						**17535532**
					GRA	CAT	
Bacteroides thetaiotaomicron		67.74	0.50	ESL 92	1	89	
Bifidobacterium longum subsp. longum		28.87	0.56	RHA 93	99	1	
Parabacteroides distasonis		2.88	0.43		1	76	
17535622	**Good identification**						**17535622**
Bacteroides thetaiotaomicron		99.26	0.47	ESL 92			
Parabacteroides distasonis		0.34	0.21	LAC 98			
Clostridium clostridioforme		0.24	0.13				
17535632	**Good identification**						**17535632**
Bacteroides thetaiotaomicron		96.56	0.40	ESL 92			
Clostridium clostridioforme		2.71	0.24	LAC 98			
Parabacteroides distasonis		0.46	0.17				
17535702	**Good identification**						**17535702**
Bacteroides thetaiotaomicron		97.25	0.42	ESL 92			
Clostridium clostridioforme		1.58	0.22	NAG 99			
Parabacteroides distasonis		0.68	0.21				
17535712	**Low discrimination**						**17535712**
Bacteroides thetaiotaomicron		82.44	0.35	ESL 92			
Clostridium clostridioforme		15.64	0.33	NAG 99			
Parabacteroides distasonis		0.79	0.17				
17535720	**Low discrimination**						**17535720**
					GRA	CAT	
Bacteroides thetaiotaomicron		61.85	0.42	ESL 92	1	89	
Bacteroides vulgatus		37.07	0.36	BGL 99	1	3	
Bifidobacterium longum subsp. longum		0.45	0.17		99	1	
17535722	**Very good identification**						**17535722**
Bacteroides thetaiotaomicron		98.96	0.76	ESL 92			
Parabacteroides distasonis		0.69	0.55				
Bacteroides ovatus		0.33	0.34				
17535723	**Good identification**						**17535723**
Bacteroides thetaiotaomicron		98.99	0.42	ESL 92			
Parabacteroides distasonis		0.69	0.21	URE 1			
Bacteroides ovatus		0.33	0.00				
17535730	**Acceptable identification**						**17535730**
					GRA	CAT	
Bacteroides thetaiotaomicron		94.08	0.35	ESL 92	1	89	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		3.02	0.21	BGL 99	99	1	
Bacteroides vulgatus		1.47	0.02		1	3	
17535732	**Very good identification**						**17535732**
Bacteroides thetaiotaomicron		98.28	0.69	ESL 92			
Parabacteroides distasonis		0.94	0.51				
Bacteroides ovatus		0.56	0.31				
17535733	**Good identification**						**17535733**
Bacteroides thetaiotaomicron		98.50	0.35	ESL 92			
Parabacteroides distasonis		0.95	0.17	URE 1			
Bacteroides ovatus		0.56	-.03				
17535762	**Good identification**						**17535762**
Bacteroides thetaiotaomicron		97.99	0.42	ESL 92			
Bacteroides ovatus		1.33	0.10	SOR 1			
Parabacteroides distasonis		0.68	0.21				
17535772	**Good identification**						**17535772**
Bacteroides thetaiotaomicron		96.81	0.35	ESL 92			
Bacteroides ovatus		2.26	0.07	SOR 1			
Parabacteroides distasonis		0.93	0.17				
17537722	**Low discrimination**						**17537722**
Bacteroides thetaiotaomicron		89.62	0.42	ESL 92			
Bacteroides ovatus		9.76	0.26	MAN 1			
Parabacteroides distasonis		0.62	0.21				
17537732	**Low discrimination**						**17537732**
Bacteroides thetaiotaomicron		83.60	0.35	ESL 92			
Bacteroides ovatus		15.60	0.23	MAN 1			
Parabacteroides distasonis		0.80	0.17				
17550000	**Very good identification**						**17550000**
					GRA	CAT	
Prevotella intermedia		99.71	0.61	CEL 1	1	1	
Fusobacterium mortiferum		0.18	0.25		1	5	
Propionibacterium granulosum		0.11	0.18		99	90	
17551000	**Very good identification**						**17551000**
					GRA		
Prevotella intermedia		99.32	0.61	CEL 1	1		
Fusobacterium mortiferum		0.27	0.28		1		
Actinomyces naeslundii		0.17	0.36		99		
17551003	**Low discrimination**						**17551003**
Actinomyces naeslundii		76.78	0.39	IND 1			
Actinomyces viscosus		23.22	0.06				
17551021	**Good identification**						**17551021**
Actinomyces naeslundii		100.0	0.39	IND 1			
17551023	**Good identification**						**17551023**
Actinomyces naeslundii		100.0	0.39	IND 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
17551101	**Good identification**					**17551101**
Actinomyces naeslundii		100.0	0.47	IND 1		
17551103	**Low discrimination**					**17551103**
Actinomyces naeslundii		84.11	0.47	IND 1		
Actinomyces viscosus		15.89	0.10			
17551120	**Acceptable identification**					**17551120**
					GRA	
Prevotella melaninogenica		94.77	0.50	IND 1	1	
Prevotella oralis		2.78	0.27	CEL 10	1	
Clostridium perfringens		1.44	0.25		99	
17551121	**Low discrimination**					**17551121**
					GRA	
Actinomyces naeslundii		56.78	0.47	IND 1	99	
Prevotella melaninogenica		43.22	0.15		1	
17551122	**Low discrimination**					**17551122**
					CAT	
Prevotella oralis		71.49	0.49	IND 1	1	
Prevotella melaninogenica		24.42	0.37		1	
Capnocytophaga ochracea		1.17	0.18		90	
17551123	**Low discrimination**					**17551123**
					GRA	
Actinomyces naeslundii		63.72	0.47	IND 1	99	
Prevotella oralis		27.05	0.15		1	
Prevotella melaninogenica		9.24	0.03		1	
17551322	**Good identification**					**17551322**
					CAT	
Prevotella oralis		95.77	0.42	IND 1	1	
Parabacteroides distasonis		2.12	0.27		76	
Bacteroides thetaiotaomicron		0.96	0.05		89	
17551722	**Low discrimination**					**17551722**
					CAT	
Bacteroides thetaiotaomicron		89.28	0.39	ESL 92	89	
Bacteroides ovatus		4.37	0.17	TRE 93	68	
Bacteroides uniformis		4.22	0.14	XYL 96	5	
17553001	**Good identification**					**17553001**
Actinomyces naeslundii		100.0	0.44	IND 1		
17553002	**Good identification**					**17553002**
Actinomyces naeslundii		100.0	0.41	IND 1		
17553003	**Good identification**					**17553003**
Actinomyces naeslundii		100.0	0.44	IND 1		
17553020	**Good identification**					**17553020**
Actinomyces naeslundii		100.0	0.41	IND 1		
17553021	**Good identification**					**17553021**
Actinomyces naeslundii		100.0	0.44	IND 1		
17553022	**Good identification**					**17553022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	IND 1		
17553023	**Good identification**					**17553023**
Actinomyces naeslundii		100.0	0.44	IND 1		
17553100	**Good identification**					**17553100**
Actinomyces naeslundii		100.0	0.49	IND 1		
17553101	**Good identification**					**17553101**
Actinomyces naeslundii		100.0	0.52	IND 1		
17553102	**Low discrimination**					**17553102**
Actinomyces naeslundii		89.04	0.49	IND 1		
Bifidobacterium adolescentis		4.87	0.11			
Actinomyces israelii		3.43	0.15			
17553103	**Good identification**					**17553103**
Actinomyces naeslundii		100.0	0.52	IND 1		
17553120	**Low discrimination**					**17553120**
					GRA	
Actinomyces naeslundii		62.96	0.49	IND 1	99	
Prevotella melaninogenica		37.04	0.15		1	
17553121	**Good identification**					**17553121**
Actinomyces naeslundii		100.0	0.52	IND 1		
17553122	**Low discrimination**					**17553122**
					GRA	
Actinomyces naeslundii		65.74	0.49	IND 1	99	
Prevotella oralis		21.56	0.15		1	
Prevotella melaninogenica		7.37	0.03		1	
17553123	**Good identification**					**17553123**
Actinomyces naeslundii		100.0	0.52	IND 1		
17553140	**Good identification**					**17553140**
Actinomyces naeslundii		100.0	0.36	IND 1		
17553141	**Good identification**					**17553141**
Actinomyces naeslundii		100.0	0.39	IND 1		
17553142	**Low discrimination**					**17553142**
Actinomyces naeslundii		85.52	0.36	IND 1		
Propionibacterium propionicum		14.48	0.12			
17553143	**Good identification**					**17553143**
Actinomyces naeslundii		100.0	0.39	IND 1		
17553160	**Good identification**					**17553160**
Actinomyces naeslundii		100.0	0.36	IND 1		
17553161	**Good identification**					**17553161**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	IND 1			
17553162	**Good identification**						**17553162**
Actinomyces naeslundii		100.0	0.36	IND 1			
17553163	**Good identification**						**17553163**
Actinomyces naeslundii		100.0	0.39	IND 1			
17555332	**Low discrimination**						**17555332**
Parabacteroides distasonis		34.91	0.39	IND 1			
Bacteroides ovatus		32.26	0.22				
Bacteroides thetaiotaomicron		30.44	0.21				
17555502	**Low discrimination**						**17555502**
					GRA	CAT	
Bifidobacterium adolescentis		72.83	0.41	IND 1	99	1	
Bacteroides uniformis		11.46	0.11		1	5	
Bacteroides thetaiotaomicron		6.12	0.09		1	89	
17555512	**Low discrimination**						**17555512**
					GRA	CAT	
Bifidobacterium adolescentis		88.96	0.43	IND 1	99	1	
Bifidobacterium longum subsp. longum		6.33	0.21		99	1	
Bacteroides thetaiotaomicron		2.38	0.02		1	89	
17555522	**Low discrimination**						**17555522**
					CAT		
Bacteroides uniformis		61.10	0.45	ESL 99	5		
Bacteroides thetaiotaomicron		32.60	0.43	SAL 92	89		
Bacteroides ovatus		4.60	0.29		68		
17555532	**Low discrimination**						**17555532**
					GRA	CAT	
Bacteroides thetaiotaomicron		69.27	0.36	ESL 92	1	89	
Bacteroides ovatus		16.77	0.26	TRE 93	1	68	
Bifidobacterium longum subsp. longum		7.46	0.32	RHA 93	99	1	
17555720	**Low discrimination**						**17555720**
					CAT		
Bacteroides vulgatus		77.22	0.38	IND 1	3		
Bacteroides thetaiotaomicron		16.31	0.29	CEL 5	89		
Bacteroides ovatus		3.29	0.17		68		
17555722	**Low discrimination**						**17555722**
					CAT		
Bacteroides thetaiotaomicron		71.29	0.63	ESL 92	89		
Bacteroides ovatus		14.39	0.51	TRE 93	68		
Bacteroides uniformis		13.89	0.48		5		
17555732	**Low discrimination**						**17555732**
Bacteroides thetaiotaomicron		73.33	0.56	ESL 92			
Bacteroides ovatus		25.38	0.48	TRE 93			
Parabacteroides distasonis		0.52	0.35				
17557502	**Low discrimination**						**17557502**
Bifidobacterium adolescentis		87.10	0.40	IND 1			
Actinomyces israelii		12.90	0.32				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17557512	**Good identification**					**17557512**
Bifidobacterium adolescentis		98.22	0.41	IND 1		
Actinomyces israelii		1.78	0.18			
17557722	**Low discrimination**					**17557722**
					CAT	
Bacteroides ovatus		84.79	0.43	ESL 91	68	
Bacteroides thetaiotaomicron		12.73	0.29	SAL 88	89	
Bacteroides uniformis		2.48	0.14	TRE 91	5	
17557732	**Acceptable identification**					**17557732**
Bacteroides ovatus		91.95	0.40	ESL 91		
Bacteroides thetaiotaomicron		8.05	0.21	SAL 88		
17565722	**Low discrimination**					**17565722**
Bacteroides thetaiotaomicron		86.55	0.48	ESL 92		
Bacteroides ovatus		13.30	0.34	FRU 99		
Parabacteroides distasonis		0.15	0.17			
17565732	**Low discrimination**					**17565732**
Bacteroides thetaiotaomicron		78.81	0.41	ESL 92		
Bacteroides ovatus		20.76	0.31	FRU 99		
Clostridium clostridioforme		0.24	0.08			
17571001	**Good identification**					**17571001**
Actinomyces naeslundii		100.0	0.44	IND 1		
17571002	**Low discrimination**					**17571002**
					CAT	
Actinomyces naeslundii		63.84	0.41	IND 1	25	
Propionibacterium granulosum		15.70	0.15		90	
Lactobacillus jensenii		12.05	0.07		1	
17571003	**Good identification**					**17571003**
Actinomyces naeslundii		100.0	0.44	IND 1		
17571020	**Low discrimination**					**17571020**
					CAT	
Actinomyces naeslundii		53.06	0.41	IND 1	25	
Clostridium perfringens		33.89	0.15		1	
Propionibacterium granulosum		13.05	0.15		90	
17571021	**Good identification**					**17571021**
Actinomyces naeslundii		100.0	0.44	IND 1		
17571022	**Low discrimination**					**17571022**
					GRA	
Actinomyces naeslundii		55.28	0.41	IND 1	99	
Prevotella oralis		20.82	0.08		1	
Clostridium perfringens		16.62	0.10		99	
17571023	**Good identification**					**17571023**
Actinomyces naeslundii		100.0	0.44	IND 1		
17571100	**Low discrimination**					**17571100**
					COC GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		59.60	0.49	IND 1	1	99	
Atopobium parvulum		21.69	0.18		99	99	
Fusobacterium mortiferum		12.36	0.25		1	1	
17571101	**Good identification**						**17571101**
Actinomyces naeslundii		100.0	0.52	IND 1			
17571102	**Low discrimination**						**17571102**
					GRA		
Actinomyces naeslundii		56.08	0.49	IND 1	99		
Lactobacillus acidophilus		29.85	0.19		99		
Prevotella oralis		6.12	0.06		1		
17571103	**Good identification**						**17571103**
Actinomyces naeslundii		100.0	0.52	IND 1			
17571121	**Good identification**						**17571121**
Actinomyces naeslundii		100.0	0.52	IND 1			
17571123	**Good identification**						**17571123**
Actinomyces naeslundii		100.0	0.52	IND 1			
17571140	**Good identification**						**17571140**
Actinomyces naeslundii		100.0	0.36	IND 1			
17571141	**Good identification**						**17571141**
Actinomyces naeslundii		100.0	0.39	IND 1			
17571142	**Low discrimination**						**17571142**
Actinomyces naeslundii		72.85	0.36	IND 1			
Lactobacillus acidophilus		27.15	0.04				
17571143	**Good identification**						**17571143**
Actinomyces naeslundii		100.0	0.39	IND 1			
17571161	**Good identification**						**17571161**
Actinomyces naeslundii		100.0	0.39	IND 1			
17571162	**Low discrimination**						**17571162**
Actinomyces naeslundii		45.27	0.36	IND 1			
Clostridium perfringens		37.85	0.12				
Lactobacillus acidophilus		16.87	0.04				
17571163	**Good identification**						**17571163**
Actinomyces naeslundii		100.0	0.39	IND 1			
17571522	**Good identification**						**17571522**
Bacteroides thetaiotaomicron		95.19	0.39	ESL 92			
Bacteroides ovatus		2.48	0.12	XYL 96			
Parabacteroides distasonis		1.98	0.26	RHA 93			
17571722	**Very good identification**						**17571722**
Bacteroides thetaiotaomicron		95.92	0.58	ESL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		3.57	0.34	XYL 96		
Parabacteroides distasonis		0.45	0.34			
17571732	**Acceptable identification**					**17571732**
Bacteroides thetaiotaomicron		92.81	0.51	ESL 92		
Bacteroides ovatus		5.93	0.31	XYL 96		
Clostridium clostridioforme		0.67	0.25			
17573000	**Low discrimination**					**17573000**
Actinomyces naeslundii		72.34	0.46	IND 1		
Propionibacterium granulosum		27.66	0.24			
17573001	**Good identification**					**17573001**
Actinomyces naeslundii		100.0	0.49	IND 1		
17573002	**Low discrimination**					**17573002**
					CAT	
Actinomyces naeslundii		87.05	0.46	IND 1	25	
Propionibacterium granulosum		8.32	0.13		90	
Actinomyces israelii		4.63	0.15		1	
17573003	**Good identification**					**17573003**
Actinomyces naeslundii		100.0	0.49	IND 1		
17573020	**Acceptable identification**					**17573020**
Actinomyces naeslundii		91.28	0.46	IND 1		
Propionibacterium granulosum		8.72	0.13			
17573021	**Good identification**					**17573021**
Actinomyces naeslundii		100.0	0.49	IND 1		
17573022	**Good identification**					**17573022**
Actinomyces naeslundii		100.0	0.46	IND 1		
17573023	**Good identification**					**17573023**
Actinomyces naeslundii		100.0	0.49	IND 1		
17573041	**Good identification**					**17573041**
Actinomyces naeslundii		100.0	0.36	IND 1		
17573043	**Good identification**					**17573043**
Actinomyces naeslundii		100.0	0.36	IND 1		
17573061	**Good identification**					**17573061**
Actinomyces naeslundii		100.0	0.36	IND 1		
17573063	**Good identification**					**17573063**
Actinomyces naeslundii		100.0	0.36	IND 1		
17573100	**Good identification**					**17573100**
Actinomyces naeslundii		95.26	0.53	IND 1		
Actinomyces israelii		4.75	0.22			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17573101	**Very good identification**					**17573101**
Actinomyces naeslundii		100.0	0.56	IND 1		
17573102	**Acceptable identification**					**17573102**
Actinomyces naeslundii		91.45	0.53	IND 1		
Actinomyces israelii		6.83	0.25			
Bifidobacterium adolescentis		1.72	0.08			
17573103	**Very good identification**					**17573103**
Actinomyces naeslundii		100.0	0.56	IND 1		
17573120	**Low discrimination**					**17573120**
					SPO	
Actinomyces naeslundii		85.94	0.53	IND 1	1	
Clostridium ramosum		14.06	0.15		92	
17573121	**Very good identification**					**17573121**
Actinomyces naeslundii		100.0	0.56	IND 1		
17573122	**Low discrimination**					**17573122**
					SPO	
Actinomyces naeslundii		60.44	0.53	IND 1	1	
Clostridium ramosum		39.56	0.25		92	
17573123	**Very good identification**					**17573123**
Actinomyces naeslundii		100.0	0.56	IND 1		
17573140	**Acceptable identification**					**17573140**
Actinomyces naeslundii		90.11	0.41	IND 1		
Actinomyces israelii		9.89	0.15			
17573141	**Good identification**					**17573141**
Actinomyces naeslundii		100.0	0.44	IND 1		
17573142	**Low discrimination**					**17573142**
Actinomyces naeslundii		85.86	0.41	IND 1		
Actinomyces israelii		14.14	0.18			
17573143	**Good identification**					**17573143**
Actinomyces naeslundii		100.0	0.44	IND 1		
17573160	**Good identification**					**17573160**
Actinomyces naeslundii		100.0	0.41	IND 1		
17573161	**Good identification**					**17573161**
Actinomyces naeslundii		100.0	0.44	IND 1		
17573162	**Good identification**					**17573162**
Actinomyces naeslundii		100.0	0.41	IND 1		
17573163	**Good identification**					**17573163**
Actinomyces naeslundii		100.0	0.44	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17574722	**Low discrimination**						**17574722**
Bacteroides thetaiotaomicron		85.92	0.48	ESL 92			
Bacteroides ovatus		13.20	0.34	GAL 99			
Clostridium clostridioforme		0.73	0.23				
17574732	**Low discrimination**						**17574732**
Bacteroides thetaiotaomicron		73.24	0.41	ESL 92			
Bacteroides ovatus		19.29	0.31	GAL 99			
Clostridium clostridioforme		7.30	0.34				
17575122	**Low discrimination**						**17575122**
Parabacteroides distasonis		46.76	0.46	IND 1			
Bacteroides thetaiotaomicron		37.18	0.29				
Bacteroides ovatus		12.23	0.20				
17575132	**Low discrimination**						**17575132**
Parabacteroides distasonis		52.14	0.42	IND 1			
Bacteroides thetaiotaomicron		29.94	0.21				
Bacteroides ovatus		16.89	0.17				
17575322	**Low discrimination**						**17575322**
Bacteroides thetaiotaomicron		56.82	0.48	ESL 92			
Bacteroides ovatus		26.73	0.42	ARA 99			
Parabacteroides distasonis		16.14	0.54				
17575332	**Low discrimination**						**17575332**
Bacteroides thetaiotaomicron		45.07	0.41	ESL 92			
Bacteroides ovatus		36.35	0.39	ARA 99			
Parabacteroides distasonis		17.72	0.50				
17575512	**Low discrimination**						**17575512**
					GRA	CAT	
Bifidobacterium adolescentis		50.47	0.40	IND 1	99	1	
Bacteroides thetaiotaomicron		26.87	0.21		1	89	
Clostridium clostridioforme		14.55	0.27		50	33	
17575522	**Low discrimination**						**17575522**
Bacteroides thetaiotaomicron		89.46	0.63	ESL 92			
Bacteroides ovatus		9.61	0.46	RHA 93			
Parabacteroides distasonis		0.70	0.43				
17575532	**Low discrimination**						**17575532**
Bacteroides thetaiotaomicron		83.16	0.56	ESL 92			
Bacteroides ovatus		15.32	0.43	RHA 93			
Parabacteroides distasonis		0.90	0.38				
17575622	**Acceptable identification**						**17575622**
Bacteroides thetaiotaomicron		92.53	0.53	ESL 92			
Bacteroides ovatus		7.04	0.34	LAC 98			
Clostridium clostridioforme		0.35	0.22				
17575632	**Low discrimination**						**17575632**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		85.05	0.46	ESL 92		
Bacteroides ovatus		11.09	0.31	LAC 98		
Clostridium clostridioforme		3.76	0.33			
17575702	**Low discrimination**					**17575702**
Bacteroides thetaiotaomicron		84.63	0.48	ESL 92		
Bacteroides ovatus		13.00	0.34	NAG 99		
Clostridium clostridioforme		2.16	0.31			
17575712	**Low discrimination**					**17575712**
Bacteroides thetaiotaomicron		63.91	0.41	ESL 92		
Clostridium clostridioforme		19.10	0.42	NAG 99		
Bacteroides ovatus		16.84	0.31			
17575720	**Low discrimination**					**17575720**
					CAT	
Bacteroides thetaiotaomicron		85.45	0.48	ESL 92	89	
Bacteroides ovatus		13.13	0.34	BGL 99	68	
Bacteroides vulgatus		1.27	0.14		3	
17575722	**Low discrimination**					**17575722**
Bacteroides thetaiotaomicron		86.52	0.82	ESL 92		
Bacteroides ovatus		13.29	0.68			
Parabacteroides distasonis		0.15	0.51			
17575723	**Low discrimination**					**17575723**
Bacteroides thetaiotaomicron		86.55	0.48	ESL 92		
Bacteroides ovatus		13.30	0.34	URE 1		
Parabacteroides distasonis		0.15	0.17			
17575730	**Low discrimination**					**17575730**
Bacteroides thetaiotaomicron		78.81	0.41	ESL 92		
Bacteroides ovatus		20.76	0.31	BGL 99		
Clostridium clostridioforme		0.24	0.08			
17575732	**Low discrimination**					**17575732**
Bacteroides thetaiotaomicron		78.81	0.75	ESL 92		
Bacteroides ovatus		20.76	0.65			
Clostridium clostridioforme		0.24	0.42			
17575733	**Low discrimination**					**17575733**
Bacteroides thetaiotaomicron		78.81	0.41	ESL 92		
Bacteroides ovatus		20.76	0.31	URE 1		
Clostridium clostridioforme		0.24	0.08			
17575762	**Low discrimination**					**17575762**
Bacteroides thetaiotaomicron		61.14	0.48	ESL 92		
Bacteroides ovatus		38.75	0.44	SOR 1		
Parabacteroides distasonis		0.11	0.17			
17575772	**Low discrimination**					**17575772**
Bacteroides ovatus		51.94	0.41	ESL 91		
Bacteroides thetaiotaomicron		47.80	0.41	SAL 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.14	0.08	SOR 4			
17577100	**Good identification**						**17577100**
					SPO		
Actinomyces israelii		97.25	0.40	IND 1	1		
Clostridium butyricum		2.75	-.08	SAL 92	99		
17577102	**Low discrimination**						**17577102**
Actinomyces israelii		87.23	0.43	IND 1			
Bifidobacterium adolescentis		12.77	0.23	SAL 92			
17577142	**Good identification**						**17577142**
Actinomyces israelii		95.56	0.36	IND 1			
Bifidobacterium adolescentis		4.44	0.07	SAL 92			
17577500	**Low discrimination**						**17577500**
					SPO		
Actinomyces israelii		76.27	0.39	IND 1	1		
Clostridium butyricum		23.73	0.09	SAL 92	99		
17577502	**Low discrimination**						**17577502**
					GRA		
Bifidobacterium adolescentis		50.08	0.37	IND 1	99		
Actinomyces israelii		41.84	0.42		99		
Bacteroides ovatus		3.67	0.04		1		
17577512	**Low discrimination**						**17577512**
					GRA		
Bifidobacterium adolescentis		85.54	0.38	IND 1	99		
Actinomyces israelii		8.74	0.28		99		
Bacteroides ovatus		3.42	0.01		1		
17577522	**Low discrimination**						**17577522**
					GRA	CAT	
Bacteroides ovatus		77.56	0.38	ESL 91	1	68	
Bacteroides thetaiotaomicron		21.87	0.29	SAL 88	1	89	
Actinomyces israelii		0.57	0.21	RHA 95	99	1	
17577722	**Low discrimination**						**17577722**
Bacteroides ovatus		83.50	0.60	ESL 91			
Bacteroides thetaiotaomicron		16.47	0.48	SAL 88			
Parabacteroides distasonis		0.03	0.17				
17577732	**Low discrimination**						**17577732**
Bacteroides ovatus		89.63	0.57	ESL 91			
Bacteroides thetaiotaomicron		10.31	0.41	SAL 88			
Clostridium clostridioforme		0.03	0.08				
17577762	**Good identification**						**17577762**
Bacteroides ovatus		95.44	0.36	ESL 91			
Bacteroides thetaiotaomicron		4.56	0.14	SAL 88			
17610000	**Low discrimination**						**17610000**
					GRA	CAT	
Fusobacterium mortiferum		86.83	0.54	IND 1	1	5	
Prevotella intermedia		10.16	0.26		1	1	
Propionibacterium granulosum		2.38	0.24		99	90	
17610002	**Acceptable identification**						**17610002**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Fusobacterium mortiferum		94.18	0.37	IND 1	1	5	
Propionibacterium granulosum		5.82	0.13	BGL 10	99	90	
17610100	**Very good identification**						**17610100**
Fusobacterium mortiferum		100.0	0.60	IND 1			
17610102	**Good identification**						**17610102**
Fusobacterium mortiferum		100.0	0.44	IND 1			
17611000	**Good identification**						**17611000**
					GRA	CAT	
Fusobacterium mortiferum		91.03	0.57	IND 1	1	5	
Prevotella intermedia		7.10	0.26		1	1	
Propionibacterium granulosum		1.67	0.24		99	90	
17611002	**Acceptable identification**						**17611002**
					GRA	CAT	
Fusobacterium mortiferum		94.25	0.40	IND 1	1	5	
Propionibacterium granulosum		3.88	0.13	BGL 10	99	90	
Actinomyces naeslundii		1.87	0.23		99	25	
17611100	**Very good identification**						**17611100**
					GRA		
Fusobacterium mortiferum		99.60	0.63	IND 1	1		
Actinomyces naeslundii		0.27	0.30		99		
Lactobacillus fermentum		0.14	0.15		99		
17611102	**Good identification**						**17611102**
					GRA		
Fusobacterium mortiferum		95.72	0.47	IND 1	1		
Actinomyces naeslundii		2.32	0.30	BGL 10	99		
Lactobacillus fermentum		1.18	0.15		99		
17613101	**Good identification**						**17613101**
Actinomyces naeslundii		100.0	0.38	IND 1			
17613102	**Low discrimination**						**17613102**
					GRA		
Actinomyces naeslundii		82.30	0.35	IND 1	99		
Fusobacterium mortiferum		17.70	0.12	MLT 94	1		
17613103	**Good identification**						**17613103**
Actinomyces naeslundii		100.0	0.38	IND 1			
17613120	**Good identification**						**17613120**
Actinomyces naeslundii		100.0	0.35	IND 1			
17613121	**Good identification**						**17613121**
Actinomyces naeslundii		100.0	0.38	IND 1			
17613122	**Good identification**						**17613122**
Actinomyces naeslundii		100.0	0.35	IND 1			
17613123	**Good identification**						**17613123**
Actinomyces naeslundii		100.0	0.38	IND 1			
17614100	**Good identification**						**17614100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.41	IND 1			
17615000	**Good identification**						**17615000**
Fusobacterium mortiferum		100.0	0.37	IND 1			
17615100	**Good identification**						**17615100**
Fusobacterium mortiferum		100.0	0.44	IND 1			
17615322	**Good identification**						**17615322**
Parabacteroides distasonis		96.52	0.39	IND 1			
Bacteroides uniformis		3.48	-.05				
17630000	**Low discrimination**						**17630000**
					GRA	CAT	
Fusobacterium mortiferum		78.39	0.45	IND 1	1	5	
Propionibacterium granulosum		21.61	0.32		99	90	
17630100	**Good identification**						**17630100**
Fusobacterium mortiferum		100.0	0.51	IND 1			
17631000	**Low discrimination**						**17631000**
					GRA	CAT	
Fusobacterium mortiferum		83.47	0.48	IND 1	1	5	
Propionibacterium granulosum		15.34	0.32		99	90	
Actinomyces naeslundii		1.20	0.28		99	25	
17631100	**Good identification**						**17631100**
					GRA		
Fusobacterium mortiferum		98.28	0.54	IND 1	1		
Actinomyces naeslundii		1.72	0.35		99		
17631101	**Low discrimination**						**17631101**
					GRA		
Actinomyces naeslundii		72.18	0.38	IND 1	99		
Fusobacterium mortiferum		27.82	0.20	MLT 94	1		
17631102	**Low discrimination**						**17631102**
					GRA		
Fusobacterium mortiferum		84.14	0.38	IND 1	1		
Actinomyces naeslundii		13.23	0.35	BGL 10	99		
Lactobacillus acidophilus		2.63	-.01		99		
17631103	**Good identification**						**17631103**
Actinomyces naeslundii		100.0	0.38	IND 1			
17631120	**Low discrimination**						**17631120**
					GRA		
Actinomyces naeslundii		46.25	0.35	IND 1	99		
Clostridium perfringens		27.03	0.09	MLT 94	99		
Fusobacterium mortiferum		26.73	0.20		1		
17631121	**Good identification**						**17631121**
Actinomyces naeslundii		100.0	0.38	IND 1			
17631123	**Good identification**						**17631123**
Actinomyces naeslundii		100.0	0.38	IND 1			
17633001	**Good identification**						**17633001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	IND 1		
17633003	**Good identification**					**17633003**
Actinomyces naeslundii		100.0	0.36	IND 1		
17633021	**Good identification**					**17633021**
Actinomyces naeslundii		100.0	0.36	IND 1		
17633023	**Good identification**					**17633023**
Actinomyces naeslundii		100.0	0.36	IND 1		
17633100	**Low discrimination**					**17633100**
					GRA	
Actinomyces naeslundii		77.06	0.40	IND 1	99	
Fusobacterium mortiferum		22.95	0.20	MLT 94	1	
17633101	**Good identification**					**17633101**
Actinomyces naeslundii		100.0	0.43	IND 1		
17633102	**Good identification**					**17633102**
Actinomyces naeslundii		100.0	0.40	IND 1		
17633103	**Good identification**					**17633103**
Actinomyces naeslundii		100.0	0.43	IND 1		
17633120	**Good identification**					**17633120**
Actinomyces naeslundii		100.0	0.40	IND 1		
17633121	**Good identification**					**17633121**
Actinomyces naeslundii		100.0	0.43	IND 1		
17633122	**Good identification**					**17633122**
Actinomyces naeslundii		100.0	0.40	IND 1		
17633123	**Good identification**					**17633123**
Actinomyces naeslundii		100.0	0.43	IND 1		
17635122	**Good identification**					**17635122**
Parabacteroides distasonis		100.0	0.42	IND 1		
17635132	**Good identification**					**17635132**
Parabacteroides distasonis		100.0	0.37	IND 1		
17635322	**Good identification**					**17635322**
Parabacteroides distasonis		96.82	0.50	IND 1		
Bacteroides thetaiotaomicron		3.18	0.09			
17635332	**Good identification**					**17635332**
Parabacteroides distasonis		97.68	0.46	IND 1		
Bacteroides thetaiotaomicron		2.32	0.02			
17635722	**Low discrimination**					**17635722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		82.65	0.43	ESL 92		
Parabacteroides distasonis		15.56	0.47	MLT 94		
Bacteroides ovatus		1.79	0.15			
17635732	**Low discrimination**					**17635732**
Bacteroides thetaiotaomicron		76.31	0.36	ESL 92		
Parabacteroides distasonis		19.89	0.42	MLT 94		
Bacteroides ovatus		2.84	0.12			
17650000	**Good identification**					**17650000**
Fusobacterium mortiferum		100.0	0.45	IND 1		
17650100	**Good identification**					**17650100**
Fusobacterium mortiferum		100.0	0.52	IND 1		
17650102	**Good identification**					**17650102**
Fusobacterium mortiferum		100.0	0.35	IND 1		
17651000	**Good identification**					**17651000**
					GRA	
Fusobacterium mortiferum		95.63	0.48	IND 1	1	
Lactobacillus catenaformis		2.86	0.12		99	
Actinomyces naeslundii		0.84	0.25		99	
17651100	**Good identification**					**17651100**
					GRA	
Fusobacterium mortiferum		97.10	0.55	IND 1	1	
Lactobacillus catenaformis		1.86	0.15		99	
Actinomyces naeslundii		1.04	0.33		99	
17651101	**Low discrimination**					**17651101**
					GRA	
Actinomyces naeslundii		61.36	0.36	IND 1	99	
Fusobacterium mortiferum		38.64	0.21	MLT 94	1	
17651102	**Low discrimination**					**17651102**
					GRA	
Fusobacterium mortiferum		46.39	0.38	IND 1	1	
Lactobacillus acidophilus		38.47	0.24	BGL 10	99	
Lactobacillus catenaformis		8.01	0.15		99	
17651103	**Good identification**					**17651103**
Actinomyces naeslundii		100.0	0.36	IND 1		
17651121	**Good identification**					**17651121**
Actinomyces naeslundii		100.0	0.36	IND 1		
17651123	**Good identification**					**17651123**
Actinomyces naeslundii		100.0	0.36	IND 1		
17653100	**Low discrimination**					**17653100**
					GRA	
Actinomyces naeslundii		67.27	0.37	IND 1	99	
Fusobacterium mortiferum		32.74	0.21	MLT 94	1	
17653101	**Good identification**					**17653101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	IND 1		
17653102	**Low discrimination**					**17653102**
Actinomyces naeslundii		81.06	0.37	IND 1		
Lactobacillus acidophilus		18.94	0.02	MLT 94		
17653103	**Good identification**					**17653103**
Actinomyces naeslundii		100.0	0.41	IND 1		
17653120	**Good identification**					**17653120**
Actinomyces naeslundii		100.0	0.37	IND 1		
17653121	**Good identification**					**17653121**
Actinomyces naeslundii		100.0	0.41	IND 1		
17653122	**Low discrimination**					**17653122**
					SPO	
Actinomyces naeslundii		62.81	0.37	IND 1	1	
Clostridium ramosum		22.51	0.05	MLT 94	92	
Lactobacillus acidophilus		14.68	0.02		1	
17653123	**Good identification**					**17653123**
Actinomyces naeslundii		100.0	0.41	IND 1		
17655100	**Good identification**					**17655100**
Fusobacterium mortiferum		100.0	0.35	IND 1		
17655522	**Good identification**					**17655522**
					CAT	
Bacteroides uniformis		98.49	0.45	ESL 99	5	
Bacteroides thetaiotaomicron		0.59	0.10	MLT 92	89	
Bacteroides ovatus		0.55	0.10		68	
17655722	**Low discrimination**					**17655722**
					CAT	
Bacteroides uniformis		87.52	0.48	ESL 99	5	
Bacteroides ovatus		6.72	0.31	MLT 92	68	
Bacteroides thetaiotaomicron		5.06	0.29		89	
17670000	**Good identification**					**17670000**
Fusobacterium mortiferum		100.0	0.36	IND 1		
17670100	**Good identification**					**17670100**
Fusobacterium mortiferum		100.0	0.43	IND 1		
17671000	**Low discrimination**					**17671000**
					GRA	
Fusobacterium mortiferum		88.63	0.39	IND 1	1	
Lactobacillus jensenii		6.32	0.10		99	
Actinomyces naeslundii		5.05	0.30		99	
17671100	**Acceptable identification**					**17671100**
					GRA	
Fusobacterium mortiferum		92.60	0.46	IND 1	1	
Actinomyces naeslundii		6.44	0.37		99	
Lactobacillus acidophilus		0.97	-.01		99	
17671101	**Acceptable identification**					**17671101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		91.17	0.41	IND 1	99	
Fusobacterium mortiferum		8.83	0.12	MLT 94	1	
17671103	**Acceptable identification**					**17671103**
Actinomyces naeslundii		90.90	0.41	IND 1		
Lactobacillus acidophilus		9.10	-.01	MLT 94		
17671120	**Low discrimination**					**17671120**
					SPO	
Actinomyces naeslundii		64.80	0.37	IND 1	1	
Clostridium ramosum		16.06	0.02	MLT 94	92	
Lactobacillus acidophilus		9.73	-.01		1	
17671121	**Good identification**					**17671121**
Actinomyces naeslundii		100.0	0.41	IND 1		
17671123	**Acceptable identification**					**17671123**
Actinomyces naeslundii		90.90	0.41	IND 1		
Lactobacillus acidophilus		9.10	-.01	MLT 94		
17673001	**Good identification**					**17673001**
Actinomyces naeslundii		100.0	0.38	IND 1		
17673003	**Good identification**					**17673003**
Actinomyces naeslundii		100.0	0.38	IND 1		
17673021	**Good identification**					**17673021**
Actinomyces naeslundii		100.0	0.38	IND 1		
17673023	**Good identification**					**17673023**
Actinomyces naeslundii		100.0	0.38	IND 1		
17673100	**Acceptable identification**					**17673100**
					GRA	
Actinomyces naeslundii		93.04	0.42	IND 1	99	
Fusobacterium mortiferum		6.97	0.12	MLT 94	1	
17673101	**Good identification**					**17673101**
Actinomyces naeslundii		100.0	0.45	IND 1		
17673102	**Low discrimination**					**17673102**
Actinomyces naeslundii		71.28	0.42	IND 1		
Lactobacillus acidophilus		28.72	0.11	MLT 94		
17673103	**Good identification**					**17673103**
Actinomyces naeslundii		100.0	0.45	IND 1		
17673120	**Low discrimination**					**17673120**
					SPO	
Actinomyces naeslundii		58.03	0.42	IND 1	1	
Clostridium ramosum		41.97	0.15	MLT 94	92	
17673121	**Good identification**					**17673121**
Actinomyces naeslundii		100.0	0.45	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17673123	**Good identification**						**17673123**
Actinomyces naeslundii		100.0	0.45	IND 1			
17675122	**Low discrimination**						**17675122**
Parabacteroides distasonis		86.87	0.37	IND 1			
Bacteroides ovatus		5.51	0.01				
Bacteroides uniformis		2.77	-.07				
17675322	**Low discrimination**						**17675322**
Parabacteroides distasonis		64.68	0.46	IND 1			
Bacteroides ovatus		25.98	0.23				
Bacteroides thetaiotaomicron		8.40	0.14				
17675332	**Low discrimination**						**17675332**
Parabacteroides distasonis		62.86	0.41	IND 1			
Bacteroides ovatus		31.25	0.20				
Bacteroides thetaiotaomicron		5.89	0.07				
17675722	**Low discrimination**						**17675722**
Bacteroides ovatus		48.74	0.49	ESL 91			
Bacteroides thetaiotaomicron		48.24	0.49	MLT 99			
Parabacteroides distasonis		2.30	0.42				
17675732	**Low discrimination**						**17675732**
Bacteroides ovatus		61.47	0.46	ESL 91			
Bacteroides thetaiotaomicron		35.49	0.42	MLT 99			
Parabacteroides distasonis		2.34	0.37				
17677722	**Good identification**						**17677722**
Bacteroides ovatus		97.09	0.41	ESL 91			
Bacteroides thetaiotaomicron		2.91	0.14	MLT 99			
17677732	**Good identification**						**17677732**
Bacteroides ovatus		98.28	0.38	ESL 91			
Bacteroides thetaiotaomicron		1.72	0.07	MLT 99			
17703101	**Good identification**						**17703101**
Actinomyces naeslundii		100.0	0.37	IND 1			
17703103	**Good identification**						**17703103**
Actinomyces naeslundii		100.0	0.37	IND 1			
17703121	**Good identification**						**17703121**
Actinomyces naeslundii		100.0	0.37	IND 1			
17703123	**Good identification**						**17703123**
Actinomyces naeslundii		100.0	0.37	IND 1			
17710000	**Very good identification**						**17710000**
					GRA	CAT	
Prevotella intermedia		98.84	0.61	SAL 1	1	1	
Fusobacterium mortiferum		1.05	0.38		1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		0.11	0.18		99	90	
17710100	**Low discrimination**						**17710100**
					GRA		
Fusobacterium mortiferum		70.07	0.44	IND 1	1		
Prevotella intermedia		28.44	0.26	MLT 11	1		
Actinomyces naeslundii		1.49	0.27		99		
17711000	**Very good identification**						**17711000**
					GRA		
Prevotella intermedia		97.76	0.61	SAL 1	1		
Fusobacterium mortiferum		1.56	0.41		1		
Actinomyces naeslundii		0.44	0.43		99		
17711003	**Low discrimination**						**17711003**
Actinomyces naeslundii		89.84	0.46	IND 1			
Actinomyces viscosus		10.16	0.06				
17711021	**Good identification**						**17711021**
Actinomyces naeslundii		100.0	0.46	IND 1			
17711023	**Good identification**						**17711023**
Actinomyces naeslundii		100.0	0.46	IND 1			
17711100	**Low discrimination**						**17711100**
					GRA		
Fusobacterium mortiferum		55.47	0.47	IND 1	1		
Actinomyces naeslundii		18.91	0.51	MLT 11	99		
Prevotella intermedia		15.01	0.26		1		
17711101	**Good identification**						**17711101**
					GRA		
Actinomyces naeslundii		98.06	0.54	IND 1	99		
Fusobacterium mortiferum		1.94	0.13		1		
17711103	**Acceptable identification**						**17711103**
Actinomyces naeslundii		92.16	0.54	IND 1			
Actinomyces viscosus		6.51	0.10				
Bifidobacterium longum subsp. infantis		1.33	-.07				
17711120	**Low discrimination**						**17711120**
					GRA		
Clostridium perfringens		39.53	0.39	IND 1	99		
Prevotella melaninogenica		39.36	0.32		1		
Actinomyces naeslundii		10.24	0.51		99		
17711121	**Good identification**						**17711121**
					GRA		
Actinomyces naeslundii		95.07	0.54	IND 1	99		
Clostridium perfringens		2.47	0.04		99		
Prevotella melaninogenica		2.46	-.03		1		
17711122	**Low discrimination**						**17711122**
					GRA		
Prevotella oralis		60.68	0.44	IND 1	1		
Bifidobacterium longum subsp. infantis		19.58	0.32		99		
Clostridium perfringens		9.33	0.33		99		
17711123	**Acceptable identification**						**17711123**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		90.48	0.54	IND 1	99	
Prevotella oralis		7.20	0.09		1	
Bifidobacterium longum subsp. infantis		2.32	-.03		99	
17711140	**Low discrimination**					**17711140**
					GRA	
Actinomyces naeslundii		77.11	0.38	IND 1	99	
Fusobacterium mortiferum		12.95	0.13		1	
Clostridium perfringens		9.94	0.00		99	
17711141	**Good identification**					**17711141**
Actinomyces naeslundii		100.0	0.41	IND 1		
17711142	**Low discrimination**					**17711142**
Actinomyces naeslundii		89.07	0.38	IND 1		
Bifidobacterium longum subsp. infantis		10.93	-.07			
17711143	**Good identification**					**17711143**
Actinomyces naeslundii		100.0	0.41	IND 1		
17711161	**Good identification**					**17711161**
Actinomyces naeslundii		100.0	0.41	IND 1		
17711163	**Good identification**					**17711163**
Actinomyces naeslundii		100.0	0.41	IND 1		
17711322	**Acceptable identification**					**17711322**
					GRA	
Prevotella oralis		91.22	0.37	IND 1	1	
Parabacteroides distasonis		8.09	0.32		1	
Bifidobacterium longum subsp. infantis		0.69	-.03		99	
17711512	**Acceptable identification**					**17711512**
Bifidobacterium adolescentis		94.38	0.36	IND 1		
Bifidobacterium longum subsp. longum		5.62	0.13	XYL 87		
17712101	**Good identification**					**17712101**
Actinomyces naeslundii		100.0	0.35	IND 1		
17712103	**Good identification**					**17712103**
Actinomyces naeslundii		100.0	0.35	IND 1		
17712121	**Good identification**					**17712121**
Actinomyces naeslundii		100.0	0.35	IND 1		
17712123	**Good identification**					**17712123**
Actinomyces naeslundii		100.0	0.35	IND 1		
17713001	**Good identification**					**17713001**
Actinomyces naeslundii		100.0	0.51	IND 1		
17713002	**Good identification**					**17713002**
Actinomyces naeslundii		100.0	0.48	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17713003	**Good identification**					**17713003**
Actinomyces naeslundii		100.0	0.51	IND 1		
17713020	**Good identification**					**17713020**
Actinomyces naeslundii		100.0	0.48	IND 1		
17713021	**Good identification**					**17713021**
Actinomyces naeslundii		100.0	0.51	IND 1		
17713022	**Good identification**					**17713022**
Actinomyces naeslundii		100.0	0.48	IND 1		
17713023	**Good identification**					**17713023**
Actinomyces naeslundii		100.0	0.51	IND 1		
17713040	**Good identification**					**17713040**
Actinomyces naeslundii		100.0	0.35	IND 1		
17713041	**Good identification**					**17713041**
Actinomyces naeslundii		100.0	0.38	IND 1		
17713042	**Good identification**					**17713042**
Actinomyces naeslundii		100.0	0.35	IND 1		
17713043	**Good identification**					**17713043**
Actinomyces naeslundii		100.0	0.38	IND 1		
17713060	**Good identification**					**17713060**
Actinomyces naeslundii		100.0	0.35	IND 1		
17713061	**Good identification**					**17713061**
Actinomyces naeslundii		100.0	0.38	IND 1		
17713062	**Good identification**					**17713062**
Actinomyces naeslundii		100.0	0.35	IND 1		
17713063	**Good identification**					**17713063**
Actinomyces naeslundii		100.0	0.38	IND 1		
17713100	**Good identification**					**17713100**
					GRA	
Actinomyces naeslundii		98.50	0.56	IND 1	99	
Fusobacterium mortiferum		1.50	0.13		1	
17713101	**Very good identification**					**17713101**
Actinomyces naeslundii		100.0	0.59	IND 1		
17713102	**Acceptable identification**					**17713102**
Actinomyces naeslundii		91.80	0.56	IND 1		
Bifidobacterium adolescentis		5.01	0.19			
Actinomyces israelii		1.15	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17713103	**Very good identification**						**17713103**
Actinomyces naeslundii		100.0	0.59	IND 1			
17713111	**Good identification**						**17713111**
Actinomyces naeslundii		100.0	0.37	IND 1			
17713113	**Good identification**						**17713113**
Actinomyces naeslundii		100.0	0.37	IND 1			
17713120	**Good identification**						**17713120**
					GRA		
Actinomyces naeslundii		96.15	0.56	IND 1	99		
Clostridium perfringens		1.93	0.04		99		
Prevotella melaninogenica		1.92	-.03		1		
17713121	**Very good identification**						**17713121**
Actinomyces naeslundii		100.0	0.59	IND 1			
17713122	**Low discrimination**						**17713122**
					SPO	GRA	
Actinomyces naeslundii		89.96	0.56	IND 1	1	99	
Prevotella oralis		5.53	0.09		1	1	
Clostridium ramosum		2.73	0.05		92	50	
17713123	**Very good identification**						**17713123**
Actinomyces naeslundii		100.0	0.59	IND 1			
17713131	**Good identification**						**17713131**
Actinomyces naeslundii		100.0	0.37	IND 1			
17713133	**Good identification**						**17713133**
Actinomyces naeslundii		100.0	0.37	IND 1			
17713140	**Good identification**						**17713140**
Actinomyces naeslundii		100.0	0.43	IND 1			
17713141	**Good identification**						**17713141**
Actinomyces naeslundii		100.0	0.46	IND 1			
17713142	**Acceptable identification**						**17713142**
Actinomyces naeslundii		94.05	0.43	IND 1			
Propionibacterium propionicum		5.95	0.12				
17713143	**Good identification**						**17713143**
Actinomyces naeslundii		100.0	0.46	IND 1			
17713160	**Good identification**						**17713160**
Actinomyces naeslundii		100.0	0.43	IND 1			
17713161	**Good identification**						**17713161**
Actinomyces naeslundii		100.0	0.46	IND 1			
17713162	**Good identification**						**17713162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.43	IND 1			
17713163	**Good identification**						**17713163**
Actinomyces naeslundii		100.0	0.46	IND 1			
17715112	**Good identification**						**17715112**
Bifidobacterium adolescentis		100.0	0.36	IND 1			
17715132	**Good identification**						**17715132**
					GRA		
Parabacteroides distasonis		95.84	0.36	IND 1	1		
Bifidobacterium longum subsp. infantis		4.16	-.04		99		
17715322	**Low discrimination**						**17715322**
Parabacteroides distasonis		72.85	0.48	IND 1			
Prevotella oralis		11.29	0.22				
Bacteroides uniformis		8.52	0.14				
17715332	**Acceptable identification**						**17715332**
Parabacteroides distasonis		90.68	0.44	IND 1			
Bacteroides thetaiotaomicron		3.26	0.03				
Bacteroides ovatus		3.09	0.03				
17715502	**Acceptable identification**						**17715502**
Bifidobacterium adolescentis		94.65	0.48	IND 1			
Bifidobacterium longum subsp. longum		1.81	0.17				
Lactobacillus fermentum		0.94	0.20				
17715512	**Good identification**						**17715512**
Bifidobacterium adolescentis		95.40	0.50	IND 1			
Bifidobacterium longum subsp. longum		2.55	0.21				
Clostridium clostridioforme		2.05	0.18				
17715720	**Good identification**						**17715720**
					GRA	CAT	
Bacteroides vulgatus		99.24	0.49	IND 1	1	3	
Bacteroides thetaiotaomicron		0.37	0.10		1	89	
Eubacterium contortum		0.19	0.10		99	1	
17715722	**Low discrimination**						**17715722**
					CAT		
Bacteroides thetaiotaomicron		61.85	0.44	ESL 92	89		
Bacteroides uniformis		16.86	0.32	TRE 93	5		
Bacteroides ovatus		11.14	0.31		68		
17715732	**Low discrimination**						**17715732**
Bacteroides thetaiotaomicron		61.27	0.37	ESL 92			
Bacteroides ovatus		18.92	0.28	TRE 93			
Parabacteroides distasonis		10.54	0.40				
17717502	**Good identification**						**17717502**
Bifidobacterium adolescentis		95.42	0.47	IND 1			
Actinomyces israelii		4.59	0.31				
17717512	**Good identification**						**17717512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		99.42	0.48	IND 1		
Actinomyces israelii		0.58	0.17			
17721101	**Good identification**					**17721101**
Actinomyces naeslundii		100.0	0.37	IND 1		
17721103	**Good identification**					**17721103**
Actinomyces naeslundii		100.0	0.37	IND 1		
17721121	**Good identification**					**17721121**
Actinomyces naeslundii		100.0	0.37	IND 1		
17721123	**Good identification**					**17721123**
Actinomyces naeslundii		100.0	0.37	IND 1		
17723100	**Good identification**					**17723100**
Actinomyces naeslundii		100.0	0.39	IND 1		
17723101	**Good identification**					**17723101**
Actinomyces naeslundii		100.0	0.42	IND 1		
17723102	**Good identification**					**17723102**
Actinomyces naeslundii		100.0	0.39	IND 1		
17723103	**Good identification**					**17723103**
Actinomyces naeslundii		100.0	0.42	IND 1		
17723120	**Good identification**					**17723120**
Actinomyces naeslundii		100.0	0.39	IND 1		
17723121	**Good identification**					**17723121**
Actinomyces naeslundii		100.0	0.42	IND 1		
17723122	**Good identification**					**17723122**
Actinomyces naeslundii		100.0	0.39	IND 1		
17723123	**Good identification**					**17723123**
Actinomyces naeslundii		100.0	0.42	IND 1		
17730100	**Low discrimination**					**17730100**
					GRA	
Fusobacterium mortiferum		87.84	0.35	IND 1	1	
Actinomyces naeslundii		12.16	0.32	MLT 11	99	
17730101	**Good identification**					**17730101**
Actinomyces naeslundii		100.0	0.35	IND 1		
17730103	**Good identification**					**17730103**
Actinomyces naeslundii		100.0	0.35	IND 1		
17730121	**Good identification**					**17730121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	IND 1		
17730123	**Good identification**					**17730123**
Actinomyces naeslundii		100.0	0.35	IND 1		
17731001	**Good identification**					**17731001**
Actinomyces naeslundii		100.0	0.51	IND 1		
17731002	**Low discrimination**					**17731002**
					CAT	
Actinomyces naeslundii		78.56	0.48	IND 1	25	
Propionibacterium granulosum		7.22	0.15		90	
Lactobacillus jensenii		5.55	0.07		1	
17731003	**Good identification**					**17731003**
Actinomyces naeslundii		100.0	0.51	IND 1		
17731021	**Good identification**					**17731021**
Actinomyces naeslundii		100.0	0.51	IND 1		
17731022	**Low discrimination**					**17731022**
					GRA	
Actinomyces naeslundii		55.73	0.48	IND 1	99	
Clostridium perfringens		37.59	0.23		99	
Prevotella oralis		3.94	0.03		1	
17731023	**Good identification**					**17731023**
Actinomyces naeslundii		100.0	0.51	IND 1		
17731040	**Good identification**					**17731040**
Actinomyces naeslundii		100.0	0.35	IND 1		
17731041	**Good identification**					**17731041**
Actinomyces naeslundii		100.0	0.38	IND 1		
17731042	**Good identification**					**17731042**
Actinomyces naeslundii		100.0	0.35	IND 1		
17731043	**Good identification**					**17731043**
Actinomyces naeslundii		100.0	0.38	IND 1		
17731061	**Good identification**					**17731061**
Actinomyces naeslundii		100.0	0.38	IND 1		
17731062	**Low discrimination**					**17731062**
Actinomyces naeslundii		57.87	0.35	IND 1		
Clostridium perfringens		42.13	0.11			
17731063	**Good identification**					**17731063**
Actinomyces naeslundii		100.0	0.38	IND 1		
17731100	**Low discrimination**					**17731100**
					GRA	
Actinomyces naeslundii		60.10	0.56	IND 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		27.14	0.38		1	
Clostridium perfringens		6.86	0.17		99	
17731101	**Very good identification**					**17731101**
Actinomyces naeslundii		100.0	0.59	IND 1		
17731102	**Low discrimination**					**17731102**
Actinomyces naeslundii		71.18	0.56	IND 1		
Lactobacillus acidophilus		14.16	0.19			
Clostridium perfringens		3.83	0.12			
17731103	**Very good identification**					**17731103**
Actinomyces naeslundii		100.0	0.59	IND 1		
17731111	**Good identification**					**17731111**
Actinomyces naeslundii		100.0	0.37	IND 1		
17731113	**Good identification**					**17731113**
Actinomyces naeslundii		100.0	0.37	IND 1		
17731120	**Low discrimination**					**17731120**
					SPO	
Clostridium perfringens		77.45	0.43	IND 1	84	
Actinomyces naeslundii		20.98	0.56		1	
Clostridium ramosum		0.44	0.02		92	
17731121	**Very good identification**					**17731121**
Actinomyces naeslundii		97.57	0.59	IND 1		
Clostridium perfringens		2.43	0.09			
17731122	**Low discrimination**					**17731122**
					GRA	
Clostridium perfringens		45.97	0.38	IND 1	99	
Actinomyces naeslundii		26.45	0.56		99	
Prevotella oralis		10.28	0.23		1	
17731123	**Very good identification**					**17731123**
Actinomyces naeslundii		98.84	0.59	IND 1		
Clostridium perfringens		1.16	0.03			
17731131	**Good identification**					**17731131**
Actinomyces naeslundii		100.0	0.37	IND 1		
17731133	**Good identification**					**17731133**
Actinomyces naeslundii		100.0	0.37	IND 1		
17731140	**Low discrimination**					**17731140**
Actinomyces naeslundii		89.03	0.43	IND 1		
Clostridium perfringens		10.97	0.05			
17731141	**Good identification**					**17731141**
Actinomyces naeslundii		100.0	0.46	IND 1		
17731142	**Low discrimination**					**17731142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		87.77	0.43	IND 1		
Lactobacillus acidophilus		12.23	0.04			
17731143	**Good identification**					**17731143**
Actinomyces naeslundii		100.0	0.46	IND 1		
17731161	**Good identification**					**17731161**
Actinomyces naeslundii		100.0	0.46	IND 1		
17731163	**Good identification**					**17731163**
Actinomyces naeslundii		100.0	0.46	IND 1		
17731322	**Low discrimination**					**17731322**
					SPO	
Parabacteroides distasonis		74.13	0.43	IND 1	1	
Prevotella oralis		11.72	0.17	XYL 90	1	
Clostridium ramosum		7.82	0.14		92	
17731332	**Acceptable identification**					**17731332**
Parabacteroides distasonis		92.05	0.39	IND 1		
Clostridium clostridioforme		4.33	0.10	XYL 90		
Bacteroides thetaiotaomicron		3.62	-.02			
17731722	**Low discrimination**					**17731722**
Bacteroides thetaiotaomicron		85.99	0.40	ESL 92		
Parabacteroides distasonis		9.77	0.40	XYL 96		
Bacteroides ovatus		2.86	0.15			
17732100	**Good identification**					**17732100**
Actinomyces naeslundii		100.0	0.37	IND 1		
17732101	**Good identification**					**17732101**
Actinomyces naeslundii		100.0	0.40	IND 1		
17732102	**Good identification**					**17732102**
Actinomyces naeslundii		100.0	0.37	IND 1		
17732103	**Good identification**					**17732103**
Actinomyces naeslundii		100.0	0.40	IND 1		
17732120	**Good identification**					**17732120**
Actinomyces naeslundii		100.0	0.37	IND 1		
17732121	**Good identification**					**17732121**
Actinomyces naeslundii		100.0	0.40	IND 1		
17732122	**Good identification**					**17732122**
Actinomyces naeslundii		100.0	0.37	IND 1		
17732123	**Good identification**					**17732123**
Actinomyces naeslundii		100.0	0.40	IND 1		
17733000	**Low discrimination**					**17733000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		87.49	0.53	IND 1		
Propionibacterium granulosum		12.51	0.24			
17733001	**Good identification**					**17733001**
Actinomyces naeslundii		100.0	0.56	IND 1		
17733002	**Acceptable identification**					**17733002**
					CAT	
Actinomyces naeslundii		94.97	0.53	IND 1	25	
Propionibacterium granulosum		3.39	0.13		90	
Actinomyces israelii		1.63	0.14		1	
17733003	**Good identification**					**17733003**
Actinomyces naeslundii		100.0	0.56	IND 1		
17733020	**Good identification**					**17733020**
Actinomyces naeslundii		96.55	0.53	IND 1		
Propionibacterium granulosum		3.45	0.13			
17733021	**Good identification**					**17733021**
Actinomyces naeslundii		100.0	0.56	IND 1		
17733022	**Good identification**					**17733022**
Actinomyces naeslundii		100.0	0.53	IND 1		
17733023	**Good identification**					**17733023**
Actinomyces naeslundii		100.0	0.56	IND 1		
17733040	**Good identification**					**17733040**
Actinomyces naeslundii		100.0	0.40	IND 1		
17733041	**Good identification**					**17733041**
Actinomyces naeslundii		100.0	0.43	IND 1		
17733042	**Good identification**					**17733042**
Actinomyces naeslundii		100.0	0.40	IND 1		
17733043	**Good identification**					**17733043**
Actinomyces naeslundii		100.0	0.43	IND 1		
17733060	**Good identification**					**17733060**
Actinomyces naeslundii		100.0	0.40	IND 1		
17733061	**Good identification**					**17733061**
Actinomyces naeslundii		100.0	0.43	IND 1		
17733062	**Good identification**					**17733062**
Actinomyces naeslundii		100.0	0.40	IND 1		
17733063	**Good identification**					**17733063**
Actinomyces naeslundii		100.0	0.43	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17733100	**Very good identification**					**17733100**
Actinomyces naeslundii		98.41	0.61	IND 1		
Actinomyces israelii		1.59	0.21			
17733101	**Very good identification**					**17733101**
Actinomyces naeslundii		100.0	0.64	IND 1		
17733102	**Very good identification**					**17733102**
Actinomyces naeslundii		95.89	0.61	IND 1		
Actinomyces israelii		2.32	0.24			
Bifidobacterium adolescentis		1.80	0.16			
17733103	**Very good identification**					**17733103**
Actinomyces naeslundii		100.0	0.64	IND 1		
17733110	**Good identification**					**17733110**
Actinomyces naeslundii		100.0	0.39	IND 1		
17733111	**Good identification**					**17733111**
Actinomyces naeslundii		100.0	0.42	IND 1		
17733112	**Low discrimination**					**17733112**
Actinomyces naeslundii		69.69	0.39	IND 1		
Bifidobacterium adolescentis		30.32	0.17	MLZ 5		
17733113	**Good identification**					**17733113**
Actinomyces naeslundii		100.0	0.42	IND 1		
17733120	**Good identification**					**17733120**
					SPO	
Actinomyces naeslundii		92.56	0.61	IND 1	1	
Clostridium ramosum		5.66	0.15		92	
Clostridium perfringens		1.78	0.09		84	
17733121	**Very good identification**					**17733121**
Actinomyces naeslundii		100.0	0.64	IND 1		
17733122	**Low discrimination**					**17733122**
					SPO	
Actinomyces naeslundii		79.76	0.61	IND 1	1	
Clostridium ramosum		19.52	0.25		92	
Clostridium perfringens		0.72	0.03		84	
17733123	**Very good identification**					**17733123**
Actinomyces naeslundii		100.0	0.64	IND 1		
17733130	**Good identification**					**17733130**
Actinomyces naeslundii		100.0	0.39	IND 1		
17733131	**Good identification**					**17733131**
Actinomyces naeslundii		100.0	0.42	IND 1		
17733132	**Good identification**					**17733132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	IND 1		
17733133	**Good identification**					**17733133**
Actinomyces naeslundii		100.0	0.42	IND 1		
17733140	**Good identification**					**17733140**
Actinomyces naeslundii		96.57	0.48	IND 1		
Actinomyces israelii		3.43	0.14			
17733141	**Good identification**					**17733141**
Actinomyces naeslundii		100.0	0.51	IND 1		
17733142	**Acceptable identification**					**17733142**
Actinomyces naeslundii		94.94	0.48	IND 1		
Actinomyces israelii		5.06	0.17			
17733143	**Good identification**					**17733143**
Actinomyces naeslundii		100.0	0.51	IND 1		
17733160	**Good identification**					**17733160**
Actinomyces naeslundii		100.0	0.48	IND 1		
17733161	**Good identification**					**17733161**
Actinomyces naeslundii		100.0	0.51	IND 1		
17733162	**Good identification**					**17733162**
Actinomyces naeslundii		100.0	0.48	IND 1		
17733163	**Good identification**					**17733163**
Actinomyces naeslundii		100.0	0.51	IND 1		
17734712	**Good identification**					**17734712**
Clostridium clostridioforme		100.0	0.40	IND 1		
17734732	**Low discrimination**					**17734732**
Clostridium clostridioforme		67.00	0.40	IND 1		
Bacteroides thetaiotaomicron		25.50	0.22			
Bacteroides ovatus		5.99	0.12			
17735122	**Acceptable identification**					**17735122**
					CAT	
Parabacteroides distasonis		93.60	0.51	IND 1	76	
Bacteroides thetaiotaomicron		3.07	0.10		89	
Prevotella oralis		1.42	0.08		1	
17735132	**Acceptable identification**					**17735132**
Parabacteroides distasonis		94.59	0.47	IND 1		
Bacteroides thetaiotaomicron		2.24	0.03			
Clostridium clostridioforme		2.04	0.13			
17735322	**Low discrimination**					**17735322**
Parabacteroides distasonis		82.32	0.60	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		11.97	0.29				
Bacteroides ovatus		5.02	0.23				
17735332	**Low discrimination**						**17735332**
Parabacteroides distasonis		81.08	0.55	IND 1			
Bacteroides thetaiotaomicron		8.51	0.22				
Bacteroides ovatus		6.12	0.20				
17735502	**Low discrimination**						**17735502**
					SPO		
Bifidobacterium adolescentis		73.65	0.45	IND 1	1		
Clostridium butyricum		13.70	0.18		99		
Clostridium clostridioforme		4.68	0.22		66		
17735512	**Low discrimination**						**17735512**
					GRA	CAT	
Bifidobacterium adolescentis		78.36	0.47	IND 1	99	1	
Clostridium clostridioforme		18.55	0.33		50	33	
Bacteroides thetaiotaomicron		1.30	0.03		1	89	
17735522	**Low discrimination**						**17735522**
Bacteroides thetaiotaomicron		76.37	0.44	ESL 92			
Parabacteroides distasonis		14.41	0.48	RHA 93			
Bacteroides ovatus		7.32	0.27				
17735532	**Low discrimination**						**17735532**
Bacteroides thetaiotaomicron		62.49	0.37	ESL 92			
Parabacteroides distasonis		16.32	0.43	RHA 93			
Bacteroides ovatus		10.27	0.24				
17735612	**Good identification**						**17735612**
Clostridium clostridioforme		100.0	0.39	IND 1			
17735632	**Low discrimination**						**17735632**
Clostridium clostridioforme		50.43	0.39	IND 1			
Bacteroides thetaiotaomicron		43.27	0.27				
Bacteroides ovatus		5.03	0.12				
17735712	**Low discrimination**						**17735712**
Clostridium clostridioforme		85.47	0.48	IND 1			
Bacteroides thetaiotaomicron		10.84	0.22				
Bacteroides ovatus		2.55	0.12				
17735722	**Low discrimination**						**17735722**
Bacteroides thetaiotaomicron		84.26	0.63	ESL 92			
Bacteroides ovatus		11.55	0.49				
Parabacteroides distasonis		3.59	0.56				
17735732	**Low discrimination**						**17735732**
Bacteroides thetaiotaomicron		72.80	0.56	ESL 92			
Bacteroides ovatus		17.11	0.46				
Clostridium clostridioforme		5.80	0.48				
17737100	**Acceptable identification**						**17737100**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		91.98	0.39	IND 1	1	
Actinomyces naeslundii		5.01	0.27	CEL 93	1	
Clostridium butyricum		3.01	-.08		99	
17737102	**Low discrimination**					**17737102**
Actinomyces israelii		67.24	0.42	IND 1		
Bifidobacterium adolescentis		30.32	0.30	CEL 93		
Actinomyces naeslundii		2.44	0.27			
17737142	**Low discrimination**					**17737142**
Actinomyces israelii		87.47	0.35	IND 1		
Bifidobacterium adolescentis		12.53	0.14	CEL 93		
17737500	**Low discrimination**					**17737500**
					SPO	
Actinomyces israelii		70.66	0.38	IND 1	1	
Clostridium butyricum		25.40	0.09	CEL 93	99	
Bifidobacterium adolescentis		3.95	0.10		1	
17737502	**Low discrimination**					**17737502**
					SPO	
Bifidobacterium adolescentis		77.13	0.44	IND 1	1	
Actinomyces israelii		20.92	0.41		1	
Clostridium butyricum		1.95	0.02		99	
17737512	**Good identification**					**17737512**
Bifidobacterium adolescentis		95.86	0.45	IND 1		
Actinomyces israelii		3.18	0.26			
Bifidobacterium dentium		0.97	-.07			
17737722	**Low discrimination**					**17737722**
Bacteroides ovatus		81.27	0.41	ESL 91		
Bacteroides thetaiotaomicron		17.97	0.29	CEL 99		
Parabacteroides distasonis		0.77	0.22			
17737732	**Low discrimination**					**17737732**
Bacteroides ovatus		87.20	0.38	ESL 91		
Bacteroides thetaiotaomicron		11.25	0.22	CEL 99		
Clostridium clostridioforme		0.90	0.14			
17743100	**Good identification**					**17743100**
Actinomyces naeslundii		100.0	0.36	IND 1		
17743101	**Good identification**					**17743101**
Actinomyces naeslundii		100.0	0.39	IND 1		
17743102	**Good identification**					**17743102**
Actinomyces naeslundii		100.0	0.36	IND 1		
17743103	**Good identification**					**17743103**
Actinomyces naeslundii		100.0	0.39	IND 1		
17743120	**Good identification**					**17743120**
Actinomyces naeslundii		100.0	0.36	IND 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
17743121	**Good identification**					**17743121**
Actinomyces naeslundii		100.0	0.39	IND 1		
17743122	**Low discrimination**					**17743122**
					SPO	
Actinomyces naeslundii		69.70	0.36	IND 1	1	
Clostridium ramosum		30.30	0.05	FRU 95	92	
17743123	**Good identification**					**17743123**
Actinomyces naeslundii		100.0	0.39	IND 1		
17750100	**Acceptable identification**					**17750100**
					GRA	
Fusobacterium mortiferum		92.19	0.36	IND 1	1	
Actinomyces naeslundii		7.81	0.29	MLT 11	99	
17751001	**Good identification**					**17751001**
Actinomyces naeslundii		100.0	0.48	IND 1		
17751003	**Good identification**					**17751003**
Actinomyces naeslundii		100.0	0.48	IND 1		
17751020	**Low discrimination**					**17751020**
					GRA	
Actinomyces naeslundii		45.82	0.45	IND 1	99	
Lactobacillus catenaformis		22.41	0.18		99	
Prevotella oralis		13.85	0.11		1	
17751021	**Good identification**					**17751021**
Actinomyces naeslundii		100.0	0.48	IND 1		
17751023	**Good identification**					**17751023**
					GRA	
Actinomyces naeslundii		96.28	0.48	IND 1	99	
Prevotella oralis		3.72	-.02		1	
17751041	**Good identification**					**17751041**
Actinomyces naeslundii		100.0	0.35	IND 1		
17751043	**Good identification**					**17751043**
Actinomyces naeslundii		100.0	0.35	IND 1		
17751061	**Good identification**					**17751061**
Actinomyces naeslundii		100.0	0.35	IND 1		
17751063	**Good identification**					**17751063**
Actinomyces naeslundii		100.0	0.35	IND 1		
17751101	**Good identification**					**17751101**
Actinomyces naeslundii		100.0	0.56	IND 1		
17751102	**Low discrimination**					**17751102**
					GRA	
Lactobacillus acidophilus		66.01	0.45	IND 1	99	
Prevotella oralis		12.74	0.31		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Lactobacillus catenaformis	7.90	0.32		99		
17751103	**Acceptable identification**						**17751103**
	Actinomyces naeslundii	92.30	0.56	IND 1			
	Lactobacillus acidophilus	5.36	0.10				
	Actinomyces viscosus	1.31	0.00				
17751121	**Good identification**						**17751121**
					GRA		
	Actinomyces naeslundii	98.89	0.56	IND 1	99		
	Prevotella oralis	1.11	-.03		1		
17751122	**Low discrimination**						**17751122**
					GRA		
	Prevotella oralis	72.54	0.53	IND 1	1		
	Lactobacillus acidophilus	19.78	0.45		99		
	Actinomyces naeslundii	2.30	0.53		99		
17751123	**Low discrimination**						**17751123**
					GRA		
	Actinomyces naeslundii	78.69	0.56	IND 1	99		
	Prevotella oralis	16.75	0.19		1		
	Lactobacillus acidophilus	4.57	0.10		99		
17751140	**Low discrimination**						**17751140**
	Actinomyces naeslundii	88.70	0.40	IND 1			
	Bifidobacterium breve	11.30	0.10				
17751141	**Good identification**						**17751141**
	Actinomyces naeslundii	100.0	0.43	IND 1			
17751143	**Good identification**						**17751143**
	Actinomyces naeslundii	100.0	0.43	IND 1			
17751160	**Low discrimination**						**17751160**
					SPO		
	Actinomyces naeslundii	62.99	0.40	IND 1	1		
	Bifidobacterium breve	14.27	0.15		1		
	Clostridium paraputrificum	8.50	-.07		99		
17751161	**Good identification**						**17751161**
	Actinomyces naeslundii	100.0	0.43	IND 1			
17751163	**Good identification**						**17751163**
	Actinomyces naeslundii	100.0	0.43	IND 1			
17751322	**Low discrimination**						**17751322**
					SPO	GRA	
	Prevotella oralis	85.34	0.47	IND 1	1	1	
	Clostridium ramosum	6.48	0.28		92	50	
	Lactobacillus acidophilus	4.09	0.25		1	99	
17751502	**Low discrimination**						**17751502**
					SPO		
	Bifidobacterium adolescentis	83.47	0.40	IND 1	1		
	Clostridium butyricum	6.37	0.06	XYL 87	99		
	Lactobacillus acidophilus	5.23	0.10		1		
17751512	**Good identification**						**17751512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		97.81	0.41	IND 1		
Clostridium clostridioforme		2.20	0.10	XYL 87		
17752101	**Good identification**					**17752101**
Actinomyces naeslundii		100.0	0.37	IND 1		
17752103	**Good identification**					**17752103**
Actinomyces naeslundii		100.0	0.37	IND 1		
17752121	**Good identification**					**17752121**
Actinomyces naeslundii		100.0	0.37	IND 1		
17752123	**Good identification**					**17752123**
Actinomyces naeslundii		100.0	0.37	IND 1		
17753000	**Acceptable identification**					**17753000**
Actinomyces naeslundii		94.40	0.50	IND 1		
Actinomyces israelii		5.60	0.20			
17753001	**Good identification**					**17753001**
Actinomyces naeslundii		100.0	0.53	IND 1		
17753002	**Low discrimination**					**17753002**
Actinomyces naeslundii		87.20	0.50	IND 1		
Actinomyces israelii		7.76	0.23			
Lactobacillus acidophilus		5.04	0.05			
17753003	**Good identification**					**17753003**
Actinomyces naeslundii		100.0	0.53	IND 1		
17753020	**Good identification**					**17753020**
Actinomyces naeslundii		100.0	0.50	IND 1		
17753021	**Good identification**					**17753021**
Actinomyces naeslundii		100.0	0.53	IND 1		
17753022	**Low discrimination**					**17753022**
					SPO	
Actinomyces naeslundii		86.74	0.50	IND 1	1	
Clostridium ramosum		5.65	0.05		92	
Lactobacillus acidophilus		5.02	0.05		1	
17753023	**Good identification**					**17753023**
Actinomyces naeslundii		100.0	0.53	IND 1		
17753040	**Good identification**					**17753040**
Actinomyces naeslundii		100.0	0.37	IND 1		
17753041	**Good identification**					**17753041**
Actinomyces naeslundii		100.0	0.40	IND 1		
17753042	**Low discrimination**					**17753042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		83.61	0.37	IND 1		
Actinomyces israelii		16.39	0.16			
17753043	**Good identification**					**17753043**
Actinomyces naeslundii		100.0	0.40	IND 1		
17753060	**Good identification**					**17753060**
Actinomyces naeslundii		100.0	0.37	IND 1		
17753061	**Good identification**					**17753061**
Actinomyces naeslundii		100.0	0.40	IND 1		
17753062	**Good identification**					**17753062**
Actinomyces naeslundii		100.0	0.37	IND 1		
17753063	**Good identification**					**17753063**
Actinomyces naeslundii		100.0	0.40	IND 1		
17753100	**Good identification**					**17753100**
Actinomyces naeslundii		91.33	0.58	IND 1		
Actinomyces israelii		7.61	0.30			
Bifidobacterium breve		1.06	0.10			
17753101	**Very good identification**					**17753101**
Actinomyces naeslundii		100.0	0.61	IND 1		
17753102	**Low discrimination**					**17753102**
Actinomyces naeslundii		62.70	0.58	IND 1		
Lactobacillus acidophilus		14.65	0.23			
Actinomyces israelii		7.84	0.33			
17753103	**Very good identification**					**17753103**
Actinomyces naeslundii		100.0	0.61	IND 1		
17753110	**Low discrimination**					**17753110**
Actinomyces naeslundii		68.65	0.36	IND 1		
Actinomyces israelii		16.25	0.16	MLZ 5		
Bifidobacterium breve		15.11	0.10			
17753111	**Good identification**					**17753111**
Actinomyces naeslundii		100.0	0.39	IND 1		
17753113	**Good identification**					**17753113**
Actinomyces naeslundii		100.0	0.39	IND 1		
17753120	**Low discrimination**					**17753120**
					SPO	
Actinomyces naeslundii		62.68	0.58	IND 1	1	
Clostridium ramosum		35.49	0.28		92	
Bifidobacterium breve		1.29	0.15		1	
17753121	**Very good identification**					**17753121**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.61	IND 1			
17753122	**Low discrimination**						**17753122**
					SPO		
Clostridium ramosum		58.14	0.39	IND 1	92		
Actinomyces naeslundii		25.67	0.58	TRE 94	1		
Lactobacillus acidophilus		6.00	0.23		1		
17753123	**Very good identification**						**17753123**
					SPO		
Actinomyces naeslundii		98.50	0.61	IND 1	1		
Clostridium ramosum		1.50	0.05		92		
17753130	**Low discrimination**						**17753130**
Actinomyces naeslundii		71.88	0.36	IND 1			
Bifidobacterium breve		28.12	0.15	MLZ 5			
17753131	**Good identification**						**17753131**
Actinomyces naeslundii		100.0	0.39	IND 1			
17753133	**Good identification**						**17753133**
Actinomyces naeslundii		100.0	0.39	IND 1			
17753140	**Low discrimination**						**17753140**
Actinomyces naeslundii		80.04	0.45	IND 1			
Actinomyces israelii		14.71	0.23				
Bifidobacterium breve		5.25	0.10				
17753141	**Good identification**						**17753141**
Actinomyces naeslundii		100.0	0.48	IND 1			
17753142	**Low discrimination**						**17753142**
Actinomyces naeslundii		46.21	0.45	IND 1			
Bifidobacterium breve		30.67	0.27				
Actinomyces israelii		12.73	0.26				
17753143	**Good identification**						**17753143**
Actinomyces naeslundii		100.0	0.48	IND 1			
17753160	**Low discrimination**						**17753160**
Actinomyces naeslundii		89.55	0.45	IND 1			
Bifidobacterium breve		10.45	0.15				
17753161	**Good identification**						**17753161**
Actinomyces naeslundii		100.0	0.48	IND 1			
17753163	**Good identification**						**17753163**
Actinomyces naeslundii		100.0	0.48	IND 1			
17753322	**Good identification**						**17753322**
					SPO	GRA	
Clostridium ramosum		96.39	0.41	IND 1	92	50	
Prevotella oralis		2.26	0.13	TRE 94	1	1	
Lactobacillus acidophilus		0.57	0.03		1	99	
17753502	**Low discrimination**						**17753502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		83.89	0.38	IND 1			
Actinomyces israelii		14.67	0.32	XYL 87			
Actinomyces naeslundii		1.45	0.24				
17753512	**Good identification**						**17753512**
Bifidobacterium adolescentis		97.91	0.40	IND 1			
Actinomyces israelii		2.09	0.18	XYL 87			
17755102	**Low discrimination**						**17755102**
					SPO		
Bifidobacterium adolescentis		49.36	0.40	IND 1	1		
Actinomyces israelii		17.52	0.39	ARA 87	1		
Clostridium butyricum		13.54	0.16		99		
17755112	**Acceptable identification**						**17755112**
Bifidobacterium adolescentis		93.49	0.41	IND 1			
Actinomyces israelii		4.06	0.24	ARA 87			
Clostridium clostridioforme		1.35	0.07				
17755122	**Low discrimination**						**17755122**
Bacteroides uniformis		83.32	0.45	ESL 99			
Prevotella oralis		12.74	0.37	ARA 92			
Parabacteroides distasonis		1.78	0.36				
17755322	**Low discrimination**						**17755322**
Bacteroides uniformis		74.67	0.48	ESL 99			
Bacteroides ovatus		17.55	0.40	ARA 92			
Prevotella oralis		3.54	0.31				
17755332	**Low discrimination**						**17755332**
Bacteroides ovatus		77.96	0.37	ESL 91			
Parabacteroides distasonis		12.46	0.39	TRE 91			
Bacteroides uniformis		5.03	0.14	ARA 97			
17755500	**Good identification**						**17755500**
					SPO		
Clostridium butyricum		95.74	0.39	IND 3	99		
Actinomyces israelii		2.92	0.34	ESL 88	1		
Bifidobacterium adolescentis		1.02	0.20	TRE 94	1		
17755502	**Low discrimination**						**17755502**
					GRA	SPO	
Bifidobacterium adolescentis		56.86	0.54	IND 1	99	1	
Clostridium butyricum		20.97	0.32		99	99	
Bacteroides uniformis		18.16	0.29		1	1	
17755512	**Acceptable identification**						**17755512**
					SPO		
Bifidobacterium adolescentis		93.80	0.56	IND 1	1		
Clostridium clostridioforme		3.18	0.27		66		
Clostridium butyricum		1.49	0.10		99		
17755522	**Good identification**						**17755522**
					CAT		
Bacteroides uniformis		94.41	0.63	ESL 99	5		
Bacteroides ovatus		4.53	0.44		68		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		0.77	0.31		89	
17755532	**Low discrimination**					**17755532**
					CAT	
Bacteroides ovatus		67.24	0.41	ESL 91	68	
Bacteroides uniformis		21.21	0.29	TRE 91	5	
Bacteroides thetaiotaomicron		6.69	0.24	RHA 95	89	
17755542	**Low discrimination**					**17755542**
					SPO	
Bifidobacterium adolescentis		70.32	0.39	IND 1	1	
Clostridium butyricum		18.25	0.14	SOR 11	99	
Actinomyces israelii		9.60	0.30		1	
17755552	**Good identification**					**17755552**
Bifidobacterium adolescentis		98.36	0.40	IND 1		
Actinomyces israelii		1.64	0.16	SOR 11		
17755622	**Low discrimination**					**17755622**
					CAT	
Bacteroides uniformis		70.02	0.37	ESL 99	5	
Bacteroides ovatus		22.90	0.31	LAC 98	68	
Bacteroides thetaiotaomicron		5.52	0.21		89	
17755712	**Low discrimination**					**17755712**
Clostridium clostridioforme		57.50	0.42	IND 1		
Bacteroides ovatus		36.76	0.28	TRE 88		
Bacteroides thetaiotaomicron		2.56	0.09			
17755722	**Low discrimination**					**17755722**
					CAT	
Bacteroides uniformis		57.35	0.66	ESL 99	5	
Bacteroides ovatus		37.90	0.66		68	
Bacteroides thetaiotaomicron		4.52	0.50		89	
17755732	**Low discrimination**					**17755732**
					CAT	
Bacteroides ovatus		89.99	0.63	ESL 91	68	
Bacteroides thetaiotaomicron		6.26	0.43	TRE 91	89	
Bacteroides uniformis		2.06	0.32		5	
17755762	**Low discrimination**					**17755762**
					CAT	
Bacteroides ovatus		71.65	0.42	ESL 91	68	
Bacteroides uniformis		26.28	0.32	TRE 91	5	
Bacteroides thetaiotaomicron		2.07	0.16	SOR 4	89	
17755772	**Good identification**					**17755772**
					CAT	
Bacteroides ovatus		97.44	0.39	ESL 91	68	
Bacteroides thetaiotaomicron		1.64	0.09	TRE 91	89	
Bacteroides uniformis		0.54	-.03	SOR 4	5	
17757000	**Good identification**					**17757000**
Actinomyces israelii		100.0	0.38	IND 1		
17757002	**Good identification**					**17757002**
Actinomyces israelii		100.0	0.41	IND 1		
17757100	**Acceptable identification**					**17757100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		93.55	0.48	IND 1	1	
Clostridium butyricum		5.47	0.06		99	
Actinomyces naeslundii		0.99	0.24		1	
17757102	**Low discrimination**					**17757102**
					SPO	
Actinomyces israelii		69.98	0.51	IND 1	1	
Bifidobacterium adolescentis		28.47	0.38		1	
Clostridium butyricum		1.06	-.01		99	
17757112	**Low discrimination**					**17757112**
					SPO	
Bifidobacterium adolescentis		76.19	0.40	IND 1	1	
Actinomyces israelii		22.89	0.37	ARA 87	1	
Clostridium tertium		0.92	-.01		99	
17757140	**Good identification**					**17757140**
Actinomyces israelii		100.0	0.41	IND 1		
17757142	**Low discrimination**					**17757142**
Actinomyces israelii		88.55	0.44	IND 1		
Bifidobacterium adolescentis		11.45	0.23			
17757400	**Acceptable identification**					**17757400**
					SPO	
Actinomyces israelii		90.32	0.37	IND 1	1	
Clostridium butyricum		9.68	-.01		99	
17757402	**Low discrimination**					**17757402**
Actinomyces israelii		88.15	0.40	IND 1		
Bifidobacterium adolescentis		11.85	0.18			
17757500	**Low discrimination**					**17757500**
					SPO	
Actinomyces israelii		59.05	0.47	IND 1	1	
Clostridium butyricum		37.97	0.23		99	
Bifidobacterium adolescentis		2.98	0.18		1	
17757502	**Low discrimination**					**17757502**
					SPO	
Bifidobacterium adolescentis		73.49	0.53	IND 1	1	
Actinomyces israelii		22.08	0.50		1	
Clostridium butyricum		3.68	0.16		99	
17757512	**Good identification**					**17757512**
					GRA	
Bifidobacterium adolescentis		95.65	0.54	IND 1	99	
Actinomyces israelii		3.51	0.36		99	
Bacteroides ovatus		0.33	-.02		1	
17757522	**Low discrimination**					**17757522**
					GRA	
Bacteroides ovatus		59.16	0.36	ESL 91	1	
Bacteroides uniformis		37.32	0.29	TRE 91	1	
Actinomyces israelii		1.83	0.29	RHA 95	99	
17757540	**Low discrimination**					**17757540**
					SPO	
Actinomyces israelii		87.43	0.40	IND 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		12.57	0.04		99	
17757542	**Low discrimination**					**17757542**
					SPO	
Bifidobacterium adolescentis		50.49	0.37	IND 1	1	
Actinomyces israelii		47.73	0.43	SOR 11	1	
Clostridium butyricum		1.78	-.03		99	
17757552	**Low discrimination**					**17757552**
Bifidobacterium adolescentis		89.64	0.39	IND 1		
Actinomyces israelii		10.36	0.29	SOR 11		
17757722	**Very good identification**					**17757722**
					CAT	
Bacteroides ovatus		95.29	0.57	ESL 91	68	
Bacteroides uniformis		4.37	0.32	TRE 91	5	
Bacteroides thetaiotaomicron		0.34	0.16		89	
17757732	**Good identification**					**17757732**
					CAT	
Bacteroides ovatus		99.67	0.54	ESL 91	68	
Bacteroides thetaiotaomicron		0.21	0.09	TRE 91	89	
Bacteroides uniformis		0.07	-.03		5	
17761100	**Good identification**					**17761100**
Actinomyces naeslundii		100.0	0.36	IND 1		
17761101	**Good identification**					**17761101**
Actinomyces naeslundii		100.0	0.39	IND 1		
17761103	**Good identification**					**17761103**
Actinomyces naeslundii		100.0	0.39	IND 1		
17761120	**Low discrimination**					**17761120**
					SPO	
Actinomyces naeslundii		76.90	0.36	IND 1	1	
Clostridium ramosum		23.11	0.02	FRU 95	92	
17761121	**Good identification**					**17761121**
Actinomyces naeslundii		100.0	0.39	IND 1		
17761123	**Good identification**					**17761123**
Actinomyces naeslundii		100.0	0.39	IND 1		
17763001	**Good identification**					**17763001**
Actinomyces naeslundii		100.0	0.36	IND 1		
17763003	**Good identification**					**17763003**
Actinomyces naeslundii		100.0	0.36	IND 1		
17763021	**Good identification**					**17763021**
Actinomyces naeslundii		100.0	0.36	IND 1		
17763023	**Good identification**					**17763023**
Actinomyces naeslundii		100.0	0.36	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17763100	**Good identification**					**17763100**
Actinomyces naeslundii		100.0	0.41	IND 1		
17763101	**Good identification**					**17763101**
Actinomyces naeslundii		100.0	0.44	IND 1		
17763102	**Good identification**					**17763102**
Actinomyces naeslundii		100.0	0.41	IND 1		
17763103	**Good identification**					**17763103**
Actinomyces naeslundii		100.0	0.44	IND 1		
17763120	**Low discrimination**					**17763120**
					SPO	
Actinomyces naeslundii		53.28	0.41	IND 1	1	
Clostridium ramosum		46.72	0.15	FRU 95	92	
17763121	**Good identification**					**17763121**
Actinomyces naeslundii		100.0	0.44	IND 1		
17763123	**Good identification**					**17763123**
Actinomyces naeslundii		100.0	0.44	IND 1		
17765722	**Low discrimination**					**17765722**
Bacteroides ovatus		86.10	0.49	ESL 91		
Bacteroides thetaiotaomicron		13.48	0.35	FRU 99		
Parabacteroides distasonis		0.15	0.17			
17765732	**Acceptable identification**					**17765732**
Bacteroides ovatus		90.57	0.46	ESL 91		
Bacteroides thetaiotaomicron		8.27	0.28	FRU 99		
Clostridium clostridioforme		1.04	0.23			
17767722	**Good identification**					**17767722**
Bacteroides ovatus		99.53	0.41	ESL 91		
Bacteroides thetaiotaomicron		0.47	0.01	FRU 99		
17767732	**Good identification**					**17767732**
Bacteroides ovatus		99.72	0.38	ESL 91		
Bacteroides thetaiotaomicron		0.28	-.06	FRU 99		
17770101	**Good identification**					**17770101**
Actinomyces naeslundii		100.0	0.37	IND 1		
17770103	**Good identification**					**17770103**
Actinomyces naeslundii		100.0	0.37	IND 1		
17770121	**Good identification**					**17770121**
Actinomyces naeslundii		100.0	0.37	IND 1		
17770123	**Good identification**					**17770123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	IND 1		
17771001	**Good identification**					**17771001**
Actinomyces naeslundii		100.0	0.53	IND 1		
17771002	**Low discrimination**					**17771002**
Lactobacillus acidophilus		63.98	0.35	IND 1		
Lactobacillus jensenii		17.60	0.29	LAC 92		
Actinomyces naeslundii		17.39	0.50			
17771003	**Good identification**					**17771003**
Actinomyces naeslundii		97.58	0.53	IND 1		
Lactobacillus acidophilus		2.42	0.01			
17771020	**Low discrimination**					**17771020**
Actinomyces naeslundii		66.37	0.50	IND 1		
Lactobacillus jensenii		25.19	0.21			
Clostridium perfringens		2.99	0.05			
17771021	**Good identification**					**17771021**
Actinomyces naeslundii		100.0	0.53	IND 1		
17771022	**Low discrimination**					**17771022**
Lactobacillus acidophilus		69.40	0.35	IND 1		
Actinomyces naeslundii		18.87	0.50	LAC 92		
Lactobacillus jensenii		4.77	0.18			
17771023	**Good identification**					**17771023**
Actinomyces naeslundii		97.58	0.53	IND 1		
Lactobacillus acidophilus		2.42	0.01			
17771040	**Good identification**					**17771040**
Actinomyces naeslundii		100.0	0.37	IND 1		
17771041	**Good identification**					**17771041**
Actinomyces naeslundii		100.0	0.40	IND 1		
17771043	**Good identification**					**17771043**
Actinomyces naeslundii		100.0	0.40	IND 1		
17771060	**Good identification**					**17771060**
Actinomyces naeslundii		100.0	0.37	IND 1		
17771061	**Good identification**					**17771061**
Actinomyces naeslundii		100.0	0.40	IND 1		
17771063	**Good identification**					**17771063**
Actinomyces naeslundii		100.0	0.40	IND 1		
17771100	**Low discrimination**					**17771100**
					SPO	
Actinomyces naeslundii		58.30	0.58	IND 1	1	
Clostridium butyricum		14.25	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		8.75	0.19		1		
17771101	**Very good identification**						**17771101**
Actinomyces naeslundii		100.0	0.61	IND 1			
17771102	**Acceptable identification**						**17771102**
					SPO		
Lactobacillus acidophilus		90.88	0.54	IND 1	1		
Actinomyces naeslundii		6.12	0.58		1		
Clostridium butyricum		0.58	0.10		99		
17771103	**Good identification**						**17771103**
Actinomyces naeslundii		90.90	0.61	IND 1			
Lactobacillus acidophilus		9.10	0.19				
17771110	**Low discrimination**						**17771110**
					GRA	SPO	
Actinomyces naeslundii		69.85	0.36	IND 1	99	1	
Clostridium butyricum		17.07	-.05	MLZ 5	99	99	
Leptotrichia buccalis		13.08	-.11		1	1	
17771111	**Good identification**						**17771111**
Actinomyces naeslundii		100.0	0.39	IND 1			
17771113	**Low discrimination**						**17771113**
					GRA		
Actinomyces naeslundii		88.90	0.39	IND 1	99		
Leptotrichia buccalis		11.10	-.11	MLZ 5	1		
17771120	**Low discrimination**						**17771120**
					SPO		
Clostridium ramosum		52.38	0.36	IND 1	92		
Actinomyces naeslundii		33.46	0.58		1		
Lactobacillus acidophilus		5.02	0.19		1		
17771121	**Very good identification**						**17771121**
					SPO		
Actinomyces naeslundii		98.96	0.61	IND 1	1		
Clostridium ramosum		1.04	0.02		92		
17771122	**Low discrimination**						**17771122**
					SPO	GRA	
Lactobacillus acidophilus		63.10	0.54	IND 1	1	99	
Clostridium ramosum		26.58	0.46		92	50	
Prevotella oralis		4.41	0.33		1	1	
17771123	**Low discrimination**						**17771123**
					SPO		
Actinomyces naeslundii		87.01	0.61	IND 1	1		
Lactobacillus acidophilus		8.71	0.19		1		
Clostridium ramosum		3.67	0.12		92		
17771130	**Low discrimination**						**17771130**
					SPO		
Actinomyces naeslundii		64.05	0.36	IND 1	1		
Clostridium ramosum		19.25	0.02	MLZ 5	92		
Bifidobacterium breve		16.71	0.12		1		
17771131	**Good identification**						**17771131**
Actinomyces naeslundii		100.0	0.39	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17771133	**Good identification**						**17771133**
Actinomyces naeslundii		100.0	0.39	IND 1			
17771140	**Low discrimination**						**17771140**
					SPO		
Actinomyces naeslundii		74.21	0.45	IND 1	1		
Actinomyces israelii		9.05	0.20		1		
Clostridium butyricum		8.94	-.01		99		
17771141	**Good identification**						**17771141**
Actinomyces naeslundii		100.0	0.48	IND 1			
17771142	**Low discrimination**						**17771142**
Lactobacillus acidophilus		86.14	0.38	IND 1			
Actinomyces naeslundii		8.28	0.45	SOR 11			
Bifidobacterium breve		3.66	0.24				
17771143	**Acceptable identification**						**17771143**
Actinomyces naeslundii		93.45	0.48	IND 1			
Lactobacillus acidophilus		6.55	0.04				
17771160	**Low discrimination**						**17771160**
Actinomyces naeslundii		71.54	0.45	IND 1			
Clostridium perfringens		8.96	0.07				
Lactobacillus acidophilus		7.52	0.04				
17771161	**Good identification**						**17771161**
Actinomyces naeslundii		100.0	0.48	IND 1			
17771162	**Low discrimination**						**17771162**
Lactobacillus acidophilus		82.14	0.38	IND 1			
Actinomyces naeslundii		7.89	0.45	SOR 11			
Bifidobacterium breve		6.21	0.29				
17771163	**Acceptable identification**						**17771163**
Actinomyces naeslundii		93.45	0.48	IND 1			
Lactobacillus acidophilus		6.55	0.04				
17771320	**Good identification**						**17771320**
					SPO	GRA	
Clostridium ramosum		97.89	0.38	IND 1	92	50	
Prevotella oralis		1.11	0.04		1	1	
Lactobacillus acidophilus		0.53	0.00		1	99	
17771322	**Low discrimination**						**17771322**
					SPO	GRA	
Clostridium ramosum		76.53	0.48	IND 1	92	50	
Lactobacillus acidophilus		10.31	0.34		1	99	
Prevotella oralis		4.11	0.26		1	1	
17771512	**Low discrimination**						**17771512**
					SPO		
Bifidobacterium adolescentis		73.61	0.38	IND 1	1		
Clostridium clostridioforme		18.17	0.25	XYL 87	66		
Clostridium butyricum		5.69	0.05		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17771712	**Low discrimination**					**17771712**
Clostridium clostridioforme		88.97	0.40	IND 1		
Bacteroides ovatus		8.01	0.12	XYL 91		
Bacteroides thetaiotaomicron		3.02	0.04			
17771722	**Low discrimination**					**17771722**
Bacteroides ovatus		59.20	0.49	ESL 91		
Bacteroides thetaiotaomicron		38.24	0.45	XYL 99		
Parabacteroides distasonis		1.10	0.35			
17771732	**Low discrimination**					**17771732**
Bacteroides ovatus		66.49	0.46	ESL 91		
Bacteroides thetaiotaomicron		25.06	0.38	XYL 99		
Clostridium clostridioforme		7.46	0.40			
17772100	**Good identification**					**17772100**
Actinomyces naeslundii		100.0	0.39	IND 1		
17772101	**Good identification**					**17772101**
Actinomyces naeslundii		100.0	0.42	IND 1		
17772102	**Good identification**					**17772102**
Actinomyces naeslundii		100.0	0.39	IND 1		
17772103	**Good identification**					**17772103**
Actinomyces naeslundii		100.0	0.42	IND 1		
17772121	**Good identification**					**17772121**
Actinomyces naeslundii		100.0	0.42	IND 1		
17772123	**Good identification**					**17772123**
Actinomyces naeslundii		100.0	0.42	IND 1		
17773000	**Low discrimination**					**17773000**
Actinomyces naeslundii		89.69	0.55	IND 1		
Actinomyces israelii		10.31	0.30			
17773001	**Very good identification**					**17773001**
Actinomyces naeslundii		100.0	0.58	IND 1		
17773002	**Low discrimination**					**17773002**
Actinomyces naeslundii		78.61	0.55	IND 1		
Actinomyces israelii		13.56	0.33			
Lactobacillus acidophilus		7.84	0.14			
17773003	**Very good identification**					**17773003**
Actinomyces naeslundii		100.0	0.58	IND 1		
17773011	**Good identification**					**17773011**
Actinomyces naeslundii		100.0	0.36	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
17773013	**Good identification**					**17773013**
Actinomyces naeslundii		100.0	0.36	IND 1		
17773020	**Low discrimination**					**17773020**
					SPO	
Actinomyces naeslundii		88.39	0.55	IND 1	1	
Clostridium ramosum		11.61	0.15		92	
17773021	**Very good identification**					**17773021**
Actinomyces naeslundii		100.0	0.58	IND 1		
17773022	**Low discrimination**					**17773022**
					SPO	
Actinomyces naeslundii		61.53	0.55	IND 1	1	
Clostridium ramosum		32.33	0.25		92	
Lactobacillus acidophilus		6.14	0.14		1	
17773023	**Very good identification**					**17773023**
Actinomyces naeslundii		100.0	0.58	IND 1		
17773031	**Good identification**					**17773031**
Actinomyces naeslundii		100.0	0.36	IND 1		
17773033	**Good identification**					**17773033**
Actinomyces naeslundii		100.0	0.36	IND 1		
17773040	**Low discrimination**					**17773040**
Actinomyces naeslundii		79.79	0.42	IND 1		
Actinomyces israelii		20.21	0.23			
17773041	**Good identification**					**17773041**
Actinomyces naeslundii		100.0	0.45	IND 1		
17773042	**Low discrimination**					**17773042**
Actinomyces naeslundii		68.97	0.42	IND 1		
Actinomyces israelii		26.21	0.26			
Lactobacillus acidophilus		4.82	-.02			
17773043	**Good identification**					**17773043**
Actinomyces naeslundii		100.0	0.45	IND 1		
17773060	**Good identification**					**17773060**
Actinomyces naeslundii		100.0	0.42	IND 1		
17773061	**Good identification**					**17773061**
Actinomyces naeslundii		100.0	0.45	IND 1		
17773062	**Acceptable identification**					**17773062**
Actinomyces naeslundii		93.47	0.42	IND 1		
Lactobacillus acidophilus		6.53	-.02			
17773063	**Good identification**					**17773063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	IND 1		
17773100	**Low discrimination**					**17773100**
					SPO	
Actinomyces naeslundii		81.49	0.63	IND 1	1	
Actinomyces israelii		13.17	0.40		1	
Clostridium ramosum		3.76	0.15		92	
17773101	**Very good identification**					**17773101**
Actinomyces naeslundii		100.0	0.66	IND 1		
17773102	**Low discrimination**					**17773102**
Actinomyces naeslundii		52.03	0.63	IND 1		
Lactobacillus acidophilus		20.97	0.32			
Actinomyces israelii		12.61	0.43			
17773103	**Very good identification**					**17773103**
Actinomyces naeslundii		100.0	0.66	IND 1		
17773110	**Low discrimination**					**17773110**
					SPO	
Actinomyces naeslundii		60.88	0.41	IND 1	1	
Actinomyces israelii		27.93	0.26	MLZ 5	1	
Clostridium tertium		11.19	0.05		99	
17773111	**Good identification**					**17773111**
Actinomyces naeslundii		100.0	0.44	IND 1		
17773112	**Low discrimination**					**17773112**
Actinomyces naeslundii		26.33	0.41	IND 1		
Bifidobacterium breve		20.13	0.24	MLZ 5		
Bifidobacterium adolescentis		18.36	0.23			
17773113	**Good identification**					**17773113**
Actinomyces naeslundii		100.0	0.44	IND 1		
17773120	**Low discrimination**					**17773120**
					SPO	
Clostridium ramosum		81.55	0.49	IND 1	92	
Actinomyces naeslundii		17.85	0.63		1	
Clostridium tertium		0.29	0.09		99	
17773121	**Very good identification**					**17773121**
					SPO	
Actinomyces naeslundii		97.02	0.66	IND 1	1	
Clostridium ramosum		2.99	0.15		92	
17773122	**Good identification**					**17773122**
					SPO	
Clostridium ramosum		92.21	0.59	IND 1	92	
Actinomyces naeslundii		5.05	0.63		1	
Lactobacillus acidophilus		2.03	0.32		1	
17773123	**Low discrimination**					**17773123**
					SPO	
Actinomyces naeslundii		89.04	0.66	IND 1	1	
Clostridium ramosum		10.96	0.25		92	
17773130	**Low discrimination**					**17773130**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		39.02	0.41	IND 1	1		
Clostridium ramosum		34.22	0.15	MLZ 5	92		
Clostridium tertium		21.52	0.13		99		
17773131	**Good identification**						**17773131**
Actinomyces naeslundii		100.0	0.44	IND 1			
17773133	**Good identification**						**17773133**
Actinomyces naeslundii		100.0	0.44	IND 1			
17773140	**Low discrimination**						**17773140**
Actinomyces naeslundii		73.74	0.50	IND 1			
Actinomyces israelii		26.26	0.33				
17773141	**Good identification**						**17773141**
Actinomyces naeslundii		100.0	0.53	IND 1			
17773142	**Low discrimination**						**17773142**
Actinomyces naeslundii		48.92	0.50	IND 1			
Actinomyces israelii		26.13	0.36				
Lactobacillus acidophilus		13.81	0.16				
17773143	**Good identification**						**17773143**
Actinomyces naeslundii		100.0	0.53	IND 1			
17773160	**Low discrimination**						**17773160**
					SPO		
Actinomyces naeslundii		76.83	0.50	IND 1	1		
Clostridium ramosum		20.10	0.15		92		
Bifidobacterium breve		3.08	0.12		1		
17773161	**Good identification**						**17773161**
Actinomyces naeslundii		100.0	0.53	IND 1			
17773163	**Good identification**						**17773163**
Actinomyces naeslundii		100.0	0.53	IND 1			
17773320	**Good identification**						**17773320**
					SPO		
Clostridium ramosum		99.83	0.51	IND 1	92		
Actinomyces naeslundii		0.17	0.29		1		
17773322	**Very good identification**						**17773322**
					SPO	GRA	
Clostridium ramosum		99.53	0.61	IND 1	92	50	
Bacteroides ovatus		0.31	0.15		1	1	
Lactobacillus acidophilus		0.13	0.12		1	99	
17773500	**Low discrimination**						**17773500**
					SPO		
Actinomyces israelii		49.29	0.39	IND 1	1		
Clostridium butyricum		46.95	0.17	XYL 92	99		
Actinomyces naeslundii		3.77	0.29		1		
17773502	**Low discrimination**						**17773502**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		43.85	0.35	IND 1	1		
Actinomyces israelii		43.26	0.42	XYL 87	1		
Clostridium butyricum		10.68	0.10		99		
17773512	**Low discrimination**						**17773512**
Bifidobacterium adolescentis		86.78	0.37	IND 1			
Actinomyces israelii		10.47	0.28	XYL 87			
Bifidobacterium dentium		2.76	-.07				
17773722	**Low discrimination**						**17773722**
					SPO	CAT	
Bacteroides ovatus		89.24	0.41	ESL 91	1	68	
Clostridium ramosum		9.02	0.27	XYL 99	92	1	
Bacteroides thetaiotaomicron		1.75	0.11		1	89	
17773732	**Good identification**						**17773732**
Bacteroides ovatus		98.54	0.38	ESL 91			
Bacteroides thetaiotaomicron		1.13	0.04	XYL 99			
Clostridium clostridioforme		0.34	0.06				
17774612	**Good identification**						**17774612**
Clostridium clostridioforme		100.0	0.40	IND 1			
17774632	**Acceptable identification**						**17774632**
Clostridium clostridioforme		90.42	0.40	IND 1			
Bacteroides ovatus		8.09	0.12				
Bacteroides thetaiotaomicron		1.49	-.01				
17774702	**Low discrimination**						**17774702**
Clostridium clostridioforme		81.82	0.37	IND 1			
Bacteroides ovatus		14.93	0.15				
Bacteroides thetaiotaomicron		2.34	0.01				
17774712	**Good identification**						**17774712**
Clostridium clostridioforme		97.17	0.49	IND 1			
Bacteroides ovatus		2.60	0.12				
Bacteroides thetaiotaomicron		0.24	-.06				
17774722	**Low discrimination**						**17774722**
Bacteroides ovatus		82.29	0.49	ESL 91			
Bacteroides thetaiotaomicron		12.89	0.35	GAL 99			
Clostridium clostridioforme		4.55	0.37				
17774732	**Low discrimination**						**17774732**
Bacteroides ovatus		67.99	0.46	ESL 91			
Clostridium clostridioforme		25.71	0.49	GAL 99			
Bacteroides thetaiotaomicron		6.21	0.28				
17775002	**Low discrimination**						**17775002**
					SPO		
Actinomyces israelii		62.85	0.38	IND 1	1		
Clostridium butyricum		33.72	0.12		99		
Lactobacillus acidophilus		3.44	0.01		1		
17775100	**Low discrimination**						**17775100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium butyricum		89.70	0.43	IND 3	1	99	
Actinomyces israelii		7.22	0.45	ESL 88	1	1	
Atopobium parvulum		2.87	0.25	ARA 90	99	1	
17775102	**Low discrimination**						**17775102**
					SPO		
Clostridium butyricum		64.49	0.36	IND 3	99		
Actinomyces israelii		20.03	0.48	ESL 88	1		
Bifidobacterium adolescentis		10.00	0.37	ARA 90	1		
17775112	**Low discrimination**						**17775112**
					SPO		
Bifidobacterium adolescentis		57.98	0.38	IND 1	1		
Clostridium butyricum		16.09	0.14	ARA 87	99		
Actinomyces israelii		14.19	0.34		1		
17775122	**Low discrimination**						**17775122**
					GRA	SPO	
Bacteroides ovatus		49.89	0.35	ESL 91	1	1	
Parabacteroides distasonis		28.18	0.47	RHA 95	1	1	
Clostridium butyricum		5.87	0.18	ARA 97	99	99	
17775142	**Low discrimination**						**17775142**
					SPO		
Actinomyces israelii		51.85	0.41	IND 1	1		
Clostridium butyricum		37.33	0.18		99		
Bifidobacterium adolescentis		8.23	0.21		1		
17775312	**Low discrimination**						**17775312**
Clostridium clostridioforme		67.28	0.36	IND 1			
Bacteroides ovatus		28.73	0.20	ARA 94			
Parabacteroides distasonis		2.07	0.16				
17775322	**Low discrimination**						**17775322**
Bacteroides ovatus		86.84	0.57	ESL 91			
Parabacteroides distasonis		7.74	0.55	ARA 97			
Bacteroides thetaiotaomicron		4.44	0.35				
17775332	**Low discrimination**						**17775332**
Bacteroides ovatus		88.85	0.54	ESL 91			
Parabacteroides distasonis		6.40	0.50	ARA 97			
Bacteroides thetaiotaomicron		2.65	0.28				
17775400	**Good identification**						**17775400**
					SPO		
Clostridium butyricum		95.79	0.36	IND 3	99		
Actinomyces israelii		4.21	0.34	ESL 88	1		
17775500	**Very good identification**						**17775500**
					SPO		
Clostridium butyricum		99.23	0.59	IND 3	99		
Actinomyces israelii		0.73	0.44	ESL 88	1		
Bifidobacterium adolescentis		0.05	0.17		1		
17775502	**Low discrimination**						**17775502**
					SPO		
Clostridium butyricum		84.14	0.52	IND 3	99		
Bifidobacterium adolescentis		9.71	0.51	ESL 88	1		

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii	2.38	0.47		1		
17775510 Good identification						**17775510**
				SPO		
Clostridium butyricum	96.60	0.38	IND 3	99		
Actinomyces israelii	2.01	0.30	ESL 88	1		
Bifidobacterium adolescentis	1.02	0.18	MLZ 5	1		
17775512 Low discrimination						**17775512**
				SPO		
Bifidobacterium adolescentis	51.99	0.53	IND 1	1		
Clostridium butyricum	19.41	0.30		99		
Clostridium clostridioforme	19.39	0.42		66		
17775520 Low discrimination						**17775520**
				GRA	SPO	
Clostridium butyricum	86.60	0.41	IND 3	99	99	
Bacteroides ovatus	10.39	0.27	ESL 88	1	1	
Bacteroides thetaiotaomicron	2.33	0.16	NAG 8	1	1	
17775522 Low discrimination						**17775522**
				GRA	SPO	
Bacteroides ovatus	77.30	0.61	ESL 91	1	1	
Bacteroides thetaiotaomicron	17.31	0.50	RHA 95	1	1	
Clostridium butyricum	2.53	0.34		99	99	
17775532 Low discrimination						**17775532**
Bacteroides ovatus	85.23	0.58	ESL 91			
Bacteroides thetaiotaomicron	11.14	0.43	RHA 95			
Clostridium clostridioforme	2.53	0.42				
17775540 Good identification						**17775540**
				SPO		
Clostridium butyricum	96.83	0.41	IND 3	99		
Actinomyces israelii	3.17	0.37	ESL 88	1		
17775552 Low discrimination						**17775552**
				SPO		
Bifidobacterium adolescentis	69.93	0.37	IND 1	1		
Clostridium butyricum	18.37	0.12	SOR 11	99		
Actinomyces israelii	6.59	0.26		1		
17775562 Low discrimination						**17775562**
				GRA	SPO	
Bacteroides ovatus	88.19	0.37	ESL 91	1	1	
Clostridium butyricum	6.02	0.16	RHA 95	99	99	
Bacteroides thetaiotaomicron	4.79	0.16	SOR 4	1	1	
17775602 Low discrimination						**17775602**
Clostridium clostridioforme	74.62	0.37	IND 1			
Bacteroides ovatus	15.20	0.15				
Bacteroides thetaiotaomicron	4.81	0.06				
17775612 Good identification						**17775612**
Clostridium clostridioforme	96.59	0.48	IND 1			
Bacteroides ovatus	2.88	0.12				
Bacteroides thetaiotaomicron	0.53	-.01				
17775622 Low discrimination						**17775622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		72.95	0.49	ESL 91			
Bacteroides thetaiotaomicron		23.08	0.40	LAC 99			
Clostridium clostridioforme		3.62	0.37				
17775632	**Low discrimination**						**17775632**
Bacteroides ovatus		65.58	0.46	ESL 91			
Clostridium clostridioforme		22.22	0.48	LAC 99			
Bacteroides thetaiotaomicron		12.11	0.33				
17775702	**Low discrimination**						**17775702**
Bacteroides ovatus		71.79	0.49	ESL 91			
Clostridium clostridioforme		11.92	0.46	NAG 99			
Bacteroides thetaiotaomicron		11.24	0.35				
17775712	**Low discrimination**						**17775712**
Clostridium clostridioforme		50.71	0.57	IND 1			
Bacteroides ovatus		44.70	0.46				
Bacteroides thetaiotaomicron		4.08	0.28				
17775720	**Low discrimination**						**17775720**
					GRA	SPO	
Bacteroides ovatus		84.87	0.49	ESL 91	1	1	
Bacteroides thetaiotaomicron		13.29	0.35	BGL 99	1	1	
Clostridium butyricum		1.15	0.15		99	99	
17775722	**Low discrimination**						**17775722**
Bacteroides ovatus		86.09	0.83	ESL 91			
Bacteroides thetaiotaomicron		13.48	0.69				
Parabacteroides distasonis		0.15	0.51				
17775723	**Low discrimination**						**17775723**
Bacteroides ovatus		86.10	0.49	ESL 91			
Bacteroides thetaiotaomicron		13.48	0.35	URE 1			
Parabacteroides distasonis		0.15	0.17				
17775730	**Acceptable identification**						**17775730**
Bacteroides ovatus		90.48	0.46	ESL 91			
Bacteroides thetaiotaomicron		8.27	0.28	BGL 99			
Clostridium clostridioforme		1.04	0.23				
17775732	**Good identification**						**17775732**
Bacteroides ovatus		90.56	0.80	ESL 91			
Bacteroides thetaiotaomicron		8.27	0.62				
Clostridium clostridioforme		1.04	0.57				
17775733	**Acceptable identification**						**17775733**
Bacteroides ovatus		90.57	0.46	ESL 91			
Bacteroides thetaiotaomicron		8.27	0.28	URE 1			
Clostridium clostridioforme		1.04	0.23				
17775762	**Very good identification**						**17775762**
Bacteroides ovatus		96.23	0.59	ESL 91			
Bacteroides thetaiotaomicron		3.65	0.35	SOR 4			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		0.04	0.17			
17775772	**Very good identification**					**17775772**
Bacteroides ovatus		97.54	0.56	ESL 91		
Bacteroides thetaiotaomicron		2.16	0.28	SOR 4		
Clostridium clostridioforme		0.27	0.23			
17776722	**Good identification**					**17776722**
Bacteroides ovatus		99.53	0.41	ESL 91		
Bacteroides thetaiotaomicron		0.47	0.01	GAL 99		
17776732	**Good identification**					**17776732**
Bacteroides ovatus		98.60	0.38	ESL 91		
Clostridium clostridioforme		1.13	0.15	GAL 99		
Bacteroides thetaiotaomicron		0.27	-.06			
17777000	**Good identification**					**17777000**
					SPO	
Actinomyces israelii		96.10	0.48	IND 1	1	
Clostridium butyricum		3.90	0.03		99	
17777002	**Good identification**					**17777002**
					SPO	
Actinomyces israelii		98.96	0.51	IND 1	1	
Clostridium butyricum		1.04	-.04		99	
17777012	**Good identification**					**17777012**
Actinomyces israelii		100.0	0.37	IND 1		
17777040	**Good identification**					**17777040**
Actinomyces israelii		100.0	0.41	IND 1		
17777042	**Good identification**					**17777042**
Actinomyces israelii		100.0	0.44	IND 1		
17777100	**Low discrimination**					**17777100**
					SPO	
Actinomyces israelii		79.81	0.58	IND 1	1	
Clostridium butyricum		19.43	0.27		99	
Actinomyces naeslundii		0.43	0.29		1	
17777102	**Low discrimination**					**17777102**
					SPO	
Actinomyces israelii		87.19	0.61	IND 1	1	
Bifidobacterium adolescentis		6.29	0.35		1	
Clostridium butyricum		5.50	0.20		99	
17777110	**Low discrimination**					**17777110**
					SPO	
Actinomyces israelii		88.19	0.44	IND 1	1	
Clostridium butyricum		7.56	0.05	MLZ 13	99	
Clostridium tertium		4.24	0.07		99	
17777112	**Low discrimination**					**17777112**
					SPO	
Actinomyces israelii		57.90	0.47	IND 1	1	
Bifidobacterium adolescentis		34.15	0.37	MLZ 13	1	
Clostridium tertium		5.57	0.15		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
17777120	**Low discrimination**						**17777120**
					SPO		
Actinomyces israelii		50.10	0.38	IND 1	1		
Clostridium ramosum		19.50	0.15	NAG 6	92		
Clostridium butyricum		16.62	0.09		99		
17777140	**Acceptable identification**						**17777140**
					SPO		
Actinomyces israelii		94.84	0.51	IND 1	1		
Clostridium butyricum		5.16	0.09		99		
17777142	**Good identification**						**17777142**
					SPO		
Actinomyces israelii		96.43	0.54	IND 1	1		
Bifidobacterium adolescentis		2.21	0.20		1		
Clostridium butyricum		1.36	0.02		99		
17777150	**Acceptable identification**						**17777150**
					COC		
Actinomyces israelii		93.35	0.37	IND 1	1		
Blautia producta		6.65	-.03	MLZ 13	99		
17777152	**Low discrimination**						**17777152**
Actinomyces israelii		84.21	0.40	IND 1			
Bifidobacterium adolescentis		15.79	0.21	MLZ 13			
17777300	**Acceptable identification**						**17777300**
					SPO		
Actinomyces israelii		90.91	0.39	IND 1	1		
Clostridium butyricum		9.09	0.01	RHA 7	99		
17777302	**Low discrimination**						**17777302**
					GRA	SPO	
Actinomyces israelii		72.30	0.42	IND 1	99	1	
Bacteroides ovatus		25.82	0.15	RHA 7	1	1	
Clostridium butyricum		1.88	-.06		99	99	
17777322	**Good identification**						**17777322**
					SPO		
Bacteroides ovatus		96.24	0.49	ESL 91	1		
Clostridium ramosum		3.18	0.27	ARA 97	92		
Parabacteroides distasonis		0.26	0.21		1		
17777332	**Good identification**						**17777332**
Bacteroides ovatus		99.69	0.46	ESL 91			
Parabacteroides distasonis		0.22	0.16	ARA 97			
Bacteroides thetaiotaomicron		0.09	-.06				
17777342	**Good identification**						**17777342**
Actinomyces israelii		100.0	0.35	IND 1			
17777400	**Low discrimination**						**17777400**
					SPO		
Actinomyces israelii		69.14	0.46	IND 1	1		
Clostridium butyricum		30.86	0.19		99		
17777402	**Low discrimination**						**17777402**
					SPO		
Actinomyces israelii		87.75	0.49	IND 1	1		
Clostridium butyricum		10.16	0.12		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		2.09	0.15		1		
17777412	**Low discrimination**						**17777412**
Actinomyces israelii		77.73	0.35	IND 1			
Bifidobacterium adolescentis		15.15	0.17	MLZ 13			
Bifidobacterium dentium		7.12	-.07				
17777440	**Acceptable identification**						**17777440**
					SPO		
Actinomyces israelii		90.93	0.39	IND 1	1		
Clostridium butyricum		9.08	0.01		99		
17777442	**Good identification**						**17777442**
					SPO		
Actinomyces israelii		97.48	0.42	IND 1	1		
Clostridium butyricum		2.52	-.06		99		
17777500	**Low discrimination**						**17777500**
					SPO		
Clostridium butyricum		72.64	0.43	IND 3	99		
Actinomyces israelii		27.12	0.57	ESL 88	1		
Bifidobacterium adolescentis		0.24	0.15	MAN 10	1		
17777502	**Low discrimination**						**17777502**
					SPO		
Actinomyces israelii		42.67	0.60	IND 1	1		
Clostridium butyricum		29.63	0.36		99		
Bifidobacterium adolescentis		25.17	0.50		1		
17777510	**Low discrimination**						**17777510**
					SPO		
Actinomyces israelii		41.90	0.43	IND 1	1		
Clostridium butyricum		39.52	0.21	MLZ 13	99		
Bifidobacterium dentium		14.10	0.10		1		
17777512	**Low discrimination**						**17777512**
Bifidobacterium adolescentis		56.04	0.51	IND 1			
Bifidobacterium dentium		26.35	0.27				
Actinomyces israelii		11.62	0.46				
17777522	**Good identification**						**17777522**
					GRA	SPO	
Bacteroides ovatus		96.90	0.53	ESL 91	1	1	
Actinomyces israelii		1.12	0.39	RHA 95	99	1	
Clostridium butyricum		1.06	0.18		99	99	
17777532	**Low discrimination**						**17777532**
					GRA	CAT	
Bacteroides ovatus		85.21	0.50	ESL 91	1	68	
Bifidobacterium dentium		13.91	0.32	RHA 95	99	1	
Bacteroides thetaiotaomicron		0.34	0.09		1	89	
17777540	**Low discrimination**						**17777540**
					COC	SPO	
Actinomyces israelii		49.30	0.50	IND 1	1	1	
Clostridium butyricum		29.52	0.25		1	99	
Blautia producta		21.18	0.24		99	1	
17777542	**Low discrimination**						**17777542**
					SPO		
Actinomyces israelii		72.44	0.53	IND 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		13.58	0.34		1		
Clostridium butyricum		11.25	0.18		99		
17777552	**Low discrimination**						**17777552**
					COC		
Bifidobacterium adolescentis		46.67	0.36	IND 1	1		
Actinomyces israelii		30.44	0.39	SOR 11	1		
Blautia producta		19.44	0.15		99		
17777622	**Good identification**						**17777622**
Bacteroides ovatus		99.05	0.41	ESL 91			
Bacteroides thetaiotaomicron		0.95	0.06	LAC 99			
17777632	**Good identification**						**17777632**
Bacteroides ovatus		98.44	0.38	ESL 91			
Clostridium clostridioforme		1.01	0.14	LAC 99			
Bacteroides thetaiotaomicron		0.55	-.01				
17777702	**Acceptable identification**						**17777702**
					GRA	SPO	
Bacteroides ovatus		90.38	0.41	ESL 91	1	1	
Actinomyces israelii		6.40	0.41	NAG 99	99	1	
Clostridium butyricum		1.83	0.10		99	99	
17777712	**Acceptable identification**						**17777712**
					GRA		
Bacteroides ovatus		93.62	0.38	ESL 91	1		
Clostridium clostridioforme		3.22	0.23	NAG 99	50		
Actinomyces israelii		1.49	0.26		99		
17777720	**Good identification**						**17777720**
					GRA	SPO	
Bacteroides ovatus		98.79	0.41	ESL 91	1	1	
Bacteroides thetaiotaomicron		0.47	0.01	BGL 99	1	1	
Clostridium butyricum		0.45	-.01		99	99	
17777722	**Very good identification**						**17777722**
					GRA	CAT	
Bacteroides ovatus		99.51	0.75	ESL 91	1	68	
Bacteroides thetaiotaomicron		0.47	0.35		1	89	
Actinomyces israelii		0.01	0.20		99	1	
17777723	**Good identification**						**17777723**
Bacteroides ovatus		99.53	0.41	ESL 91			
Bacteroides thetaiotaomicron		0.47	0.01	URE 1			
17777730	**Good identification**						**17777730**
Bacteroides ovatus		99.72	0.38	ESL 91			
Bacteroides thetaiotaomicron		0.28	-.06	BGL 99			
17777732	**Very good identification**						**17777732**
Bacteroides ovatus		99.68	0.72	ESL 91			
Bacteroides thetaiotaomicron		0.28	0.28				
Clostridium clostridioforme		0.04	0.23				
17777733	**Good identification**						**17777733**
Bacteroides ovatus		99.72	0.38	ESL 91			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		0.28	-.06	URE 1			
17777762	**Good identification**						**17777762**
Bacteroides ovatus		99.89	0.51	ESL 91			
Bacteroides thetaiotaomicron		0.12	0.01	SOR 4			
17777772	**Good identification**						**17777772**
					COC	GRA	
Bacteroides ovatus		99.79	0.48	ESL 91	1	1	
Blautia producta		0.14	-.01	SOR 4	99	99	
Bacteroides thetaiotaomicron		0.07	-.06		1	1	
20000000	**Low discrimination**						**20000000**
					COC	GRA	
Veillonella parvula		51.40	1.00		99	1	
Campylobacter gracilis		37.76	1.00		1	1	
Eggerthella lenta		6.23	0.85		1	99	
20000001	**Good identification**						**20000001**
					COC		
Campylobacter ureolyticus		93.37	0.94		1		
Veillonella parvula		3.14	0.66		99		
Campylobacter gracilis		2.31	0.66		1		
20000002	**Low discrimination**						**20000002**
					COC	GRA	
Campylobacter gracilis		84.02	0.84	BGL 10	1	1	
Veillonella parvula		10.40	0.66		99	1	
Peptococcus niger		1.29	0.53		99	99	
20000003	**Low discrimination**						**20000003**
					COC	GRA	
Campylobacter ureolyticus		68.18	0.60	BGL 1	1	1	
Campylobacter gracilis		18.53	0.50		1	1	
Anaerococcus prevotii		7.33	0.47		99	99	
20000010	**Low discrimination**						**20000010**
					COC	GRA	
Veillonella parvula		51.39	0.66	MLZ 1	99	1	
Campylobacter gracilis		37.75	0.66		1	1	
Eggerthella lenta		6.23	0.51		1	99	
20000011	**Good identification**						**20000011**
					COC		
Campylobacter ureolyticus		93.27	0.60	MLZ 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.30	0.32		1		
20000012	**Low discrimination**						**20000012**
					COC	GRA	
Campylobacter gracilis		83.99	0.50	MLZ 1	1	1	
Veillonella parvula		10.39	0.32	BGL 10	99	1	
Peptococcus niger		1.29	0.19		99	99	
20000020	**Good identification**						**20000020**
					COC	GRA	
Campylobacter gracilis		93.34	0.90		1	1	
Veillonella parvula		5.13	0.66		99	1	
Eggerthella lenta		0.62	0.51		1	99	
20000021	**Low discrimination**						**20000021**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Campylobacter ureolyticus		59.46	0.60	NAG 1	1		
Campylobacter gracilis		36.35	0.56		1		
Veillonella parvula		2.00	0.32		99		
20000022	**Low discrimination**						**20000022**
					COC	GRA	
Campylobacter gracilis		81.68	0.73	BGL 10	1	1	
Pseudoflavonifractor capillosus		16.88	0.66		1	1	
Peptococcus niger		0.56	0.37		99	99	
20000023	**Low discrimination**						**20000023**
Campylobacter gracilis		69.10	0.39	URE 1			
Pseudoflavonifractor capillosus		14.28	0.32	BGL 10			
Campylobacter ureolyticus		10.28	0.26				
20000030	**Acceptable identification**						**20000030**
					COC	GRA	
Campylobacter gracilis		93.30	0.56	MLZ 1	1	1	
Veillonella parvula		5.13	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
20000032	**Low discrimination**						**20000032**
					COC	GRA	
Campylobacter gracilis		81.80	0.39	MLZ 1	1	1	
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1	
Peptococcus niger		0.56	0.03		99	99	
20000040	**Low discrimination**						**20000040**
					COC	GRA	
Veillonella parvula		51.31	0.66	SOR 1	99	1	
Campylobacter gracilis		37.70	0.66		1	1	
Eggerthella lenta		6.22	0.51		1	99	
20000041	**Good identification**						**20000041**
					COC		
Campylobacter ureolyticus		93.30	0.60	SOR 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20000042	**Low discrimination**						**20000042**
					COC	GRA	
Campylobacter gracilis		83.98	0.50	SOR 1	1	1	
Veillonella parvula		10.39	0.32	BGL 10	99	1	
Peptococcus niger		1.29	0.19		99	99	
20000060	**Acceptable identification**						**20000060**
					COC	GRA	
Campylobacter gracilis		93.29	0.56	SOR 1	1	1	
Veillonella parvula		5.13	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
20000062	**Low discrimination**						**20000062**
					COC	GRA	
Campylobacter gracilis		81.80	0.39	SOR 1	1	1	
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1	
Peptococcus niger		0.56	0.03		99	99	
20000100	**Low discrimination**						**20000100**
					COC	GRA	
Veillonella parvula		51.40	0.66	LAC 1	99	1	
Campylobacter gracilis		37.76	0.66		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eggerthella lenta		6.23	0.51		1	99	
20000101	**Good identification**						**20000101**
					COC		
Campylobacter ureolyticus		93.38	0.60	LAC 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20000102	**Low discrimination**						**20000102**
					COC	GRA	
Campylobacter gracilis		84.05	0.50	LAC 1	1	1	
Veillonella parvula		10.40	0.32	BGL 10	99	1	
Peptococcus niger		1.29	0.19		99	99	
20000120	**Acceptable identification**						**20000120**
					COC	GRA	
Campylobacter gracilis		93.34	0.56	LAC 1	1	1	
Veillonella parvula		5.13	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
20000122	**Low discrimination**						**20000122**
					COC	GRA	
Campylobacter gracilis		81.80	0.39	LAC 1	1	1	
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1	
Peptococcus niger		0.56	0.03		99	99	
20000200	**Low discrimination**						**20000200**
					COC	GRA	
Veillonella parvula		51.40	0.66	RHA 1	99	1	
Campylobacter gracilis		37.76	0.66		1	1	
Eggerthella lenta		6.23	0.51		1	99	
20000201	**Good identification**						**20000201**
					COC		
Campylobacter ureolyticus		93.38	0.60	RHA 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20000202	**Low discrimination**						**20000202**
					COC	GRA	
Campylobacter gracilis		84.05	0.50	RHA 1	1	1	
Veillonella parvula		10.40	0.32	BGL 10	99	1	
Peptococcus niger		1.29	0.19		99	99	
20000220	**Acceptable identification**						**20000220**
					COC	GRA	
Campylobacter gracilis		93.34	0.56	RHA 1	1	1	
Veillonella parvula		5.13	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
20000222	**Low discrimination**						**20000222**
					COC	GRA	
Campylobacter gracilis		81.80	0.39	RHA 1	1	1	
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1	
Peptococcus niger		0.56	0.03		99	99	
20000400	**Low discrimination**						**20000400**
					COC	GRA	
Veillonella parvula		51.40	0.66	ARA 1	99	1	
Campylobacter gracilis		37.76	0.66		1	1	
Eggerthella lenta		6.23	0.51		1	99	
20000401	**Good identification**						**20000401**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Campylobacter ureolyticus		93.38	0.60	ARA 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20000402	**Low discrimination**						**20000402**
					COC	GRA	
Campylobacter gracilis		84.05	0.50	ARA 1	1	1	
Veillonella parvula		10.40	0.32	BGL 10	99	1	
Peptococcus niger		1.29	0.19		99	99	
20000420	**Acceptable identification**						**20000420**
					COC	GRA	
Campylobacter gracilis		93.34	0.56	ARA 1	1	1	
Veillonella parvula		5.13	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
20000422	**Low discrimination**						**20000422**
					COC	GRA	
Campylobacter gracilis		81.80	0.39	ARA 1	1	1	
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1	
Peptococcus niger		0.56	0.03		99	99	
20001000	**Low discrimination**						**20001000**
					COC	GRA	
Veillonella parvula		50.97	0.66	GAL 1	99	1	
Campylobacter gracilis		37.44	0.66		1	1	
Eggerthella lenta		6.18	0.51		1	99	
20001001	**Good identification**						**20001001**
					COC		
Campylobacter ureolyticus		93.32	0.60	GAL 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20001002	**Low discrimination**						**20001002**
					COC	GRA	
Campylobacter gracilis		77.40	0.50	GAL 1	1	1	
Veillonella parvula		9.58	0.32	BGL 10	99	1	
Atopobium minutum		7.69	0.51		1	99	
20001020	**Acceptable identification**						**20001020**
					COC	GRA	
Campylobacter gracilis		91.26	0.56	GAL 1	1	1	
Veillonella parvula		5.02	0.32		99	1	
Atopobium minutum		2.02	0.46		1	99	
20001022	**Low discrimination**						**20001022**
					GRA		
Pseudoflavonifractor capillosus		66.51	0.50	NIT 1	1		
Campylobacter gracilis		29.27	0.39	GAL 10	1		
Atopobium minutum		3.88	0.43		99		
20002000	**Low discrimination**						**20002000**
					COC	GRA	
Veillonella parvula		51.40	0.66	MAN 1	99	1	
Campylobacter gracilis		37.76	0.66		1	1	
Eggerthella lenta		6.23	0.51		1	99	
20002001	**Good identification**						**20002001**
					COC		
Campylobacter ureolyticus		93.38	0.60	MAN 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20002002	**Low discrimination**						**20002002**
					COC	GRA	
Campylobacter gracilis		84.05	0.50	MAN 1	1	1	
Veillonella parvula		10.40	0.32	BGL 10	99	1	
Peptococcus niger		1.29	0.19		99	99	
20002020	**Acceptable identification**						**20002020**
					COC	GRA	
Campylobacter gracilis		93.32	0.56	MAN 1	1	1	
Veillonella parvula		5.13	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
20002022	**Low discrimination**						**20002022**
					COC	GRA	
Campylobacter gracilis		81.80	0.39	MAN 1	1	1	
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1	
Peptococcus niger		0.56	0.03		99	99	
20004000	**Low discrimination**						**20004000**
					COC	GRA	
Veillonella parvula		51.40	0.66	XYL 1	99	1	
Campylobacter gracilis		37.76	0.66		1	1	
Eggerthella lenta		6.23	0.51		1	99	
20004001	**Good identification**						**20004001**
					COC		
Campylobacter ureolyticus		93.38	0.60	XYL 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20004002	**Low discrimination**						**20004002**
					COC	GRA	
Campylobacter gracilis		84.05	0.50	XYL 1	1	1	
Veillonella parvula		10.40	0.32	BGL 10	99	1	
Peptococcus niger		1.29	0.19		99	99	
20004020	**Acceptable identification**						**20004020**
					COC	GRA	
Campylobacter gracilis		93.34	0.56	XYL 1	1	1	
Veillonella parvula		5.13	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
20004022	**Low discrimination**						**20004022**
					COC	GRA	
Campylobacter gracilis		81.80	0.39	XYL 1	1	1	
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1	
Peptococcus niger		0.56	0.03		99	99	
20010000	**Low discrimination**						**20010000**
					COC	GRA	
Veillonella parvula		49.94	0.66	FRU 1	99	1	
Campylobacter gracilis		36.68	0.66		1	1	
Eggerthella lenta		6.05	0.51		1	99	
20010001	**Good identification**						**20010001**
					COC		
Campylobacter ureolyticus		93.20	0.60	FRU 1	1		
Veillonella parvula		3.13	0.32		99		
Campylobacter gracilis		2.30	0.32		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
20010002	**Low discrimination**						**20010002**
					COC	GRA	
Campylobacter gracilis		67.63	0.50	FRU 1	1	1	
Atopobium minutum		18.61	0.58	BGL 10	1	99	
Veillonella parvula		8.37	0.32		99	1	
20010020	**Low discrimination**						**20010020**
					COC	GRA	
Campylobacter gracilis		84.06	0.56	FRU 1	1	1	
Atopobium minutum		5.14	0.53		1	99	
Veillonella parvula		4.62	0.32		99	1	
20010022	**Low discrimination**						**20010022**
					GRA		
Pseudoflavonifractor capillosus		62.10	0.50	NIT 1	1		
Campylobacter gracilis		27.32	0.39	FRU 10	1		
Atopobium minutum		10.03	0.50		99		
20011000	**Low discrimination**						**20011000**
					COC	GRA	
Atopobium minutum		35.89	0.53	NIT 1	1	99	
Veillonella parvula		32.28	0.32		99	1	
Campylobacter gracilis		23.71	0.32		1	1	
20011002	**Low discrimination**						**20011002**
					COC	GRA	
Atopobium minutum		88.44	0.50	NIT 1	1	99	
Campylobacter gracilis		9.74	0.15		1	1	
Veillonella parvula		1.21	-.03		99	1	
20011020	**Low discrimination**						**20011020**
					GRA	CAT	
Propionibacterium acnes		61.01	0.44	IND 87	99	95	
Atopobium minutum		25.38	0.45	GLU 99	99	1	
Campylobacter gracilis		12.57	0.21		1	1	
20012020	**Low discrimination**						**20012020**
					GRA	CAT	
Propionibacterium acnes		67.23	0.40	IND 87	99	95	
Campylobacter gracilis		24.63	0.21	GLU 99	1	1	
Pseudoramibacter alactolyticus		5.28	0.14		99	1	
20013020	**Good identification**						**20013020**
					GRA	CAT	
Propionibacterium acnes		98.90	0.40	IND 87	99	95	
Atopobium minutum		0.74	0.11	GLU 99	99	1	
Campylobacter gracilis		0.37	-.13		1	1	
20020000	**Low discrimination**						**20020000**
					COC	GRA	
Veillonella parvula		51.30	0.66	TRE 1	99	1	
Campylobacter gracilis		37.69	0.66		1	1	
Eggerthella lenta		6.22	0.51		1	99	
20020001	**Good identification**						**20020001**
					COC		
Campylobacter ureolyticus		93.37	0.60	TRE 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
20020002	**Low discrimination**						**20020002**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Campylobacter gracilis		83.64	0.50	TRE 1	1	1		
Veillonella parvula		10.35	0.32	BGL 10	99	1		
Peptococcus niger		1.28	0.19		99	99		
20020020	**Acceptable identification**							**20020020**
					COC	GRA		
Campylobacter gracilis		93.33	0.56	TRE 1	1	1		
Veillonella parvula		5.13	0.32		99	1		
Eggerthella lenta		0.62	0.17		1	99		
20020022	**Low discrimination**							**20020022**
					COC	GRA		
Campylobacter gracilis		81.80	0.39	TRE 1	1	1		
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1		
Peptococcus niger		0.56	0.03		99	99		
20040000	**Low discrimination**							**20040000**
					COC	GRA		
Veillonella parvula		51.40	0.66	CEL 1	99	1		
Campylobacter gracilis		37.76	0.66		1	1		
Eggerthella lenta		6.23	0.51		1	99		
20040001	**Good identification**							**20040001**
					COC			
Campylobacter ureolyticus		93.38	0.60	CEL 1	1			
Veillonella parvula		3.14	0.32		99			
Campylobacter gracilis		2.31	0.32		1			
20040002	**Low discrimination**							**20040002**
					COC	GRA		
Campylobacter gracilis		84.05	0.50	CEL 1	1	1		
Veillonella parvula		10.40	0.32	BGL 10	99	1		
Peptococcus niger		1.29	0.19		99	99		
20040020	**Acceptable identification**							**20040020**
					COC	GRA		
Campylobacter gracilis		93.34	0.56	CEL 1	1	1		
Veillonella parvula		5.13	0.32		99	1		
Eggerthella lenta		0.62	0.17		1	99		
20040022	**Low discrimination**							**20040022**
					COC	GRA		
Campylobacter gracilis		81.80	0.39	CEL 1	1	1		
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1		
Peptococcus niger		0.56	0.03		99	99		
20100000	**Low discrimination**							**20100000**
					COC	GRA		
Veillonella parvula		46.75	0.66	MLT 1	99	1		
Campylobacter gracilis		34.34	0.66		1	1		
Peptostreptococcus anaerobius		8.33	0.61		99	99		
20100001	**Good identification**							**20100001**
					COC			
Campylobacter ureolyticus		92.66	0.60	MLT 1	1			
Veillonella parvula		3.12	0.32		99			
Campylobacter gracilis		2.29	0.32		1			
20100002	**Low discrimination**							**20100002**
					COC	GRA	SPO	
Campylobacter gracilis		80.41	0.50	MLT 1	1	1	1	
Veillonella parvula		9.95	0.32	BGL 10	99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sporogenes		2.44	0.32		1	99	99	
20100020	**Acceptable identification**							**20100020**
					COC	GRA		
Campylobacter gracilis		92.37	0.56	MLT 1	1	1		
Veillonella parvula		5.08	0.32		99	1		
Peptostreptococcus anaerobius		0.91	0.27		99	99		
20100022	**Low discrimination**							**20100022**
					COC	GRA		
Campylobacter gracilis		81.66	0.39	MLT 1	1	1		
Pseudoflavonifractor capillosus		16.87	0.32	BGL 10	1	1		
Peptococcus niger		0.56	0.03		99	99		
20200000	**Low discrimination**							**20200000**
					COC	GRA		
Veillonella parvula		51.40	0.66	SAL 1	99	1		
Campylobacter gracilis		37.76	0.66		1	1		
Eggerthella lenta		6.23	0.51		1	99		
20200001	**Good identification**							**20200001**
					COC			
Campylobacter ureolyticus		93.37	0.60	SAL 1	1			
Veillonella parvula		3.14	0.32		99			
Campylobacter gracilis		2.31	0.32		1			
20200002	**Low discrimination**							**20200002**
					COC	GRA		
Campylobacter gracilis		84.05	0.50	SAL 1	1	1		
Veillonella parvula		10.40	0.32	BGL 10	99	1		
Peptococcus niger		1.29	0.19		99	99		
20200020	**Acceptable identification**							**20200020**
					COC	GRA		
Campylobacter gracilis		93.34	0.56	SAL 1	1	1		
Veillonella parvula		5.13	0.32		99	1		
Eggerthella lenta		0.62	0.17		1	99		
20200022	**Low discrimination**							**20200022**
					COC	GRA		
Campylobacter gracilis		81.80	0.39	SAL 1	1	1		
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1		
Peptococcus niger		0.56	0.03		99	99		
20400000	**Low discrimination**							**20400000**
					COC	GRA		
Veillonella parvula		51.28	0.66	RAF 1	99	1		
Campylobacter gracilis		37.67	0.66		1	1		
Eggerthella lenta		6.22	0.51		1	99		
20400001	**Low discrimination**							**20400001**
					COC	GRA		
Campylobacter ureolyticus		82.65	0.60	RAF 1	1	1		
Anaerococcus prevotii		11.85	0.50		99	99		
Veillonella parvula		2.78	0.32		99	1		
20400002	**Low discrimination**							**20400002**
					COC	GRA		
Campylobacter gracilis		83.05	0.50	RAF 1	1	1		
Veillonella parvula		10.28	0.32	BGL 10	99	1		
Peptococcus niger		1.28	0.19		99	99		
20400003	**Low discrimination**							**20400003**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Anaerococcus prevotii		72.49	0.39	NIT 1	99	99		
Campylobacter ureolyticus		20.43	0.26		1	1		
Campylobacter gracilis		5.55	0.15		1	1		
20400020	**Acceptable identification**							**20400020**
					COC	GRA		
Campylobacter gracilis		93.32	0.56	RAF 1	1	1		
Veillonella parvula		5.13	0.32		99	1		
Eggerthella lenta		0.62	0.17		1	99		
20400022	**Low discrimination**							**20400022**
					COC	GRA		
Campylobacter gracilis		81.80	0.39	RAF 1	1	1		
Pseudoflavonifractor capillosus		16.90	0.32	BGL 10	1	1		
Peptococcus niger		0.56	0.03		99	99		
21000000	**Low discrimination**							**21000000**
					COC	GRA		
Veillonella parvula		40.02	0.66	GLU 1	99	1		
Campylobacter gracilis		29.40	0.66		1	1		
Peptostreptococcus anaerobius		13.07	0.66		99	99		
21000001	**Low discrimination**							**21000001**
					COC	GRA	SPO	
Campylobacter ureolyticus		82.75	0.60	GLU 1	1	1	1	
Clostridium novyi biovar A		9.84	0.66		1	99	99	
Veillonella parvula		2.78	0.32		99	1	1	
21000002	**Low discrimination**							**21000002**
					COC	GRA		
Campylobacter gracilis		51.93	0.50	GLU 1	1	1		
Atopobium minutum		26.36	0.63	BGL 10	1	99		
Veillonella parvula		6.43	0.32		99	1		
21000003	**Low discrimination**							**21000003**
					GRA	SPO		
Clostridium novyi biovar A		56.07	0.53	NIT 1	99	99		
Campylobacter ureolyticus		26.99	0.26		1	1		
Campylobacter gracilis		7.33	0.15		1	1		
21000010	**Low discrimination**							**21000010**
					COC	GRA	SPO	
Clostridium novyi biovar A		58.93	0.66	NIT 1	1	99	99	
Veillonella parvula		16.66	0.32		99	1	1	
Campylobacter gracilis		12.24	0.32		1	1	1	
21000011	**Good identification**							**21000011**
					COC	GRA	SPO	
Clostridium novyi biovar A		91.67	0.66	NIT 1	1	99	99	
Campylobacter ureolyticus		7.79	0.26		1	1	1	
Veillonella parvula		0.26	-.03		99	1	1	
21000012	**Low discrimination**							**21000012**
					GRA	SPO		
Clostridium novyi biovar A		81.25	0.53	NIT 1	99	99		
Campylobacter gracilis		10.63	0.15		1	1		
Atopobium minutum		5.40	0.29		99	1		
21000013	**Good identification**							**21000013**
					GRA	SPO		
Clostridium novyi biovar A		99.52	0.53	NIT 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Campylobacter ureolyticus		0.48	-.08		1	1	
21000020	**Low discrimination**						**21000020**
					COC	GRA	
Campylobacter gracilis		74.69	0.56	GLU 1	1	1	
Atopobium minutum		8.43	0.58		1	99	
Veillonella parvula		4.11	0.32		99	1	
21000021	**Low discrimination**						**21000021**
					GRA	SPO	
Clostridium novyi biovar A		62.08	0.56	NIT 1	99	99	
Campylobacter ureolyticus		21.09	0.26		1	1	
Campylobacter gracilis		12.90	0.21		1	1	
21000022	**Low discrimination**						**21000022**
					GRA		
Campylobacter gracilis		50.28	0.39	GLU 1	1		
Atopobium minutum		34.03	0.55	BGL 10	99		
Pseudoflavonifractor capillosus		10.39	0.32		1		
21000023	**Low discrimination**						**21000023**
					GRA	SPO	
Clostridium novyi biovar A		79.00	0.43	NIT 1	99	99	
Campylobacter gracilis		10.33	0.05		1	1	
Atopobium minutum		6.99	0.21		99	1	
21000030	**Low discrimination**						**21000030**
					GRA	SPO	
Clostridium novyi biovar A		79.22	0.56	NIT 1	99	99	
Campylobacter gracilis		16.46	0.21		1	1	
Atopobium minutum		1.86	0.24		99	1	
21000031	**Good identification**						**21000031**
					GRA	SPO	
Clostridium novyi biovar A		99.45	0.56	NIT 1	99	99	
Campylobacter ureolyticus		0.34	-.08		1	1	
Campylobacter gracilis		0.21	-.13		1	1	
21000032	**Low discrimination**						**21000032**
					GRA	SPO	
Clostridium novyi biovar A		80.23	0.43	NIT 1	99	99	
Campylobacter gracilis		10.49	0.05		1	1	
Atopobium minutum		7.10	0.21		99	1	
21000033	**Good identification**						**21000033**
Clostridium novyi biovar A		100.0	0.43	NIT 1			
21000040	**Low discrimination**						**21000040**
					COC	GRA	
Clostridium botulinum biovar A		35.32	0.52	NIT 1	1	99	
Clostridium novyi biovar A		29.95	0.63		1	99	
Veillonella parvula		12.70	0.32		99	1	
21000041	**Low discrimination**						**21000041**
					GRA	SPO	
Clostridium novyi biovar A		87.10	0.63	NIT 1	99	99	
Campylobacter ureolyticus		11.10	0.26		1	1	
Clostridium botulinum biovar A		1.04	0.18		99	99	
21000042	**Low discrimination**						**21000042**
					GRA	SPO	
Clostridium novyi biovar A		67.49	0.50	NIT 1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Campylobacter gracilis		13.24	0.15		1	1		
Atopobium minutum		6.72	0.29		99	1		
21000043	**Good identification**							**21000043**
					GRA	SPO		
Clostridium novyi biovar A		99.28	0.50	NIT 1	99	99		
Campylobacter ureolyticus		0.72	-.08		1	1		
21000050	**Very good identification**							**21000050**
					COC	GRA		
Clostridium novyi biovar A		97.97	0.63	NIT 1	1	99		
Clostridium botulinum biovar A		1.17	0.18		1	99		
Veillonella parvula		0.42	-.03		99	1		
21000051	**Very good identification**							**21000051**
					GRA	SPO		
Clostridium novyi biovar A		99.87	0.63	NIT 1	99	99		
Campylobacter ureolyticus		0.13	-.08		1	1		
21000052	**Good identification**							**21000052**
Clostridium novyi biovar A		100.0	0.50	NIT 1				
21000053	**Good identification**							**21000053**
Clostridium novyi biovar A		100.0	0.50	NIT 1				
21000060	**Low discrimination**							**21000060**
					GRA	SPO	CAT	
Clostridium novyi biovar A		55.91	0.53	NIT 1	99	99	1	
Propionibacterium acnes		18.55	0.33		99	1	95	
Campylobacter gracilis		17.42	0.21		1	1	1	
21000061	**Good identification**							**21000061**
					GRA	SPO	CAT	
Clostridium novyi biovar A		98.85	0.53	NIT 1	99	99	1	
Campylobacter ureolyticus		0.51	-.08		1	1	10	
Propionibacterium acnes		0.33	-.01		99	1	95	
21000062	**Low discrimination**							**21000062**
					GRA	SPO		
Clostridium novyi biovar A		71.02	0.40	NIT 1	99	99		
Campylobacter gracilis		13.93	0.05		1	1		
Atopobium minutum		9.43	0.21		99	1		
21000063	**Good identification**							**21000063**
Clostridium novyi biovar A		100.0	0.40	NIT 1				
21000070	**Good identification**							**21000070**
					GRA	SPO	CAT	
Clostridium novyi biovar A		99.35	0.53	NIT 1	99	99	1	
Propionibacterium acnes		0.33	-.01		99	1	95	
Campylobacter gracilis		0.31	-.13		1	1	1	
21000071	**Good identification**							**21000071**
Clostridium novyi biovar A		100.0	0.53	NIT 1				
21000072	**Good identification**							**21000072**
Clostridium novyi biovar A		100.0	0.40	NIT 1				
21000073	**Good identification**							**21000073**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.40	NIT 1				
21001000	**Low discrimination**							**21001000**
					COC	GRA		
Atopobium minutum		41.62	0.58	NIT 1	1	99		
Veillonella parvula		20.29	0.32		99	1		
Campylobacter gracilis		14.91	0.32		1	1		
21001001	**Low discrimination**							**21001001**
					GRA	SPO		
Clostridium novyi biovar A		52.99	0.50	NIT 1	99	99		
Campylobacter ureolyticus		40.51	0.26	GAL 10	1	1		
Atopobium minutum		2.79	0.24		99	1		
21001002	**Low discrimination**							**21001002**
					GRA	SPO		
Atopobium minutum		88.83	0.55	NIT 1	99	1		
Campylobacter gracilis		5.30	0.15		1	1		
Clostridium novyi biovar A		4.51	0.37		99	99		
21001003	**Low discrimination**							**21001003**
					GRA	SPO		
Clostridium novyi biovar A		80.45	0.37	NIT 1	99	99		
Atopobium minutum		16.03	0.21	GAL 10	99	1		
Campylobacter ureolyticus		3.52	-.08		1	1		
21001010	**Acceptable identification**							**21001010**
					COC	SPO	GRA	
Clostridium novyi biovar A		90.44	0.50	NIT 1	1	99	99	
Atopobium minutum		4.77	0.24	GAL 10	1	1	99	
Veillonella parvula		2.33	-.03		99	1	1	
21001011	**Good identification**							**21001011**
					GRA	SPO		
Clostridium novyi biovar A		99.23	0.50	NIT 1	99	99		
Campylobacter ureolyticus		0.77	-.08	GAL 10	1	1		
21001012	**Low discrimination**							**21001012**
					SPO			
Clostridium novyi biovar A		83.39	0.37	NIT 1	99			
Atopobium minutum		16.61	0.21	GAL 10	1			
21001013	**Good identification**							**21001013**
Clostridium novyi biovar A		100.0	0.37	NIT 1				
21001020	**Low discrimination**							**21001020**
					GRA	CAT		
Propionibacterium acnes		47.52	0.44	IND 87	99	95		
Atopobium minutum		36.46	0.50	FRU 99	99	1		
Campylobacter gracilis		9.79	0.21		1	1		
21001021	**Low discrimination**							**21001021**
					SPO	CAT		
Clostridium novyi biovar A		82.53	0.39	NIT 1	99	1		
Propionibacterium acnes		7.56	0.10	GAL 10	1	95		
Atopobium minutum		5.80	0.15		1	1		
21001022	**Low discrimination**							**21001022**
					GRA			
Atopobium minutum		83.04	0.47	NIT 1	99			
Pseudoflavonifractor capillosus		8.45	0.15		1			
Campylobacter gracilis		3.72	0.05		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
21001030	**Low discrimination**						**21001030**
					SPO	CAT	
Clostridium novyi biovar A		84.69	0.39	NIT 1	99	1	
Propionibacterium acnes		7.76	0.10	GAL 10	1	95	
Atopobium minutum		5.95	0.15		1	1	
21001031	**Good identification**						**21001031**
Clostridium novyi biovar A		100.0	0.39	NIT 1			
21001040	**Low discrimination**						**21001040**
					SPO		
Clostridium novyi biovar A		75.81	0.47	NIT 1	99		
Clostridium botulinum biovar A		8.13	0.18	GAL 10	99		
Atopobium minutum		6.00	0.24		1		
21001041	**Good identification**						**21001041**
					GRA	SPO	
Clostridium novyi biovar A		98.86	0.47	NIT 1	99	99	
Campylobacter ureolyticus		1.15	-.08	GAL 10	1	1	
21001050	**Good identification**						**21001050**
Clostridium novyi biovar A		100.0	0.47	NIT 1			
21001051	**Good identification**						**21001051**
Clostridium novyi biovar A		100.0	0.47	NIT 1			
21001061	**Good identification**						**21001061**
					SPO	CAT	
Clostridium novyi biovar A		97.07	0.36	NIT 1	99	1	
Propionibacterium acnes		2.93	-.01	GAL 10	1	95	
21001070	**Good identification**						**21001070**
					SPO	CAT	
Clostridium novyi biovar A		97.07	0.36	NIT 1	99	1	
Propionibacterium acnes		2.93	-.01	GAL 10	1	95	
21001071	**Good identification**						**21001071**
Clostridium novyi biovar A		100.0	0.36	NIT 1			
21002020	**Low discrimination**						**21002020**
					GRA	CAT	
Propionibacterium acnes		64.70	0.40	IND 87	99	95	
Campylobacter gracilis		23.71	0.21	FRU 99	1	1	
Pseudoramibacter alactolyticus		5.08	0.14		99	1	
21003020	**Good identification**						**21003020**
					GRA	CAT	
Propionibacterium acnes		98.28	0.40	IND 87	99	95	
Atopobium minutum		1.35	0.15	FRU 99	99	1	
Campylobacter gracilis		0.36	-.13		1	1	
21004040	**Good identification**						**21004040**
Clostridium glycolicum		96.57	0.49	NIT 1			
Clostridium botulinum biovar A		1.32	0.18	FRU 90			
Clostridium novyi biovar A		1.12	0.29				
21010000	**Low discrimination**						**21010000**
					GRA		
Pseudoramibacter alactolyticus		63.83	0.66	NIT 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium varium		8.97	0.51		1		
Atopobium minutum		8.84	0.65		99		
21010002	**Low discrimination**						**21010002**
					COC	GRA	
Pseudoramibacter alactolyticus		49.35	0.50	NIT 1	1	99	
Atopobium minutum		41.03	0.62	BGL 10	1	99	
Megasphaera elsdenii		5.20	0.44		99	1	
21010003	**Low discrimination**						**21010003**
					SPO		
Clostridium novyi biovar A		42.94	0.37	NIT 1	99		
Pseudoramibacter alactolyticus		28.50	0.15	FRU 10	1		
Atopobium minutum		23.69	0.28		1		
21010010	**Low discrimination**						**21010010**
					SPO		
Clostridium difficile		31.53	0.36	NIT 1	99		
Pseudoramibacter alactolyticus		27.46	0.32	MAN 90	1		
Clostridium novyi biovar A		26.05	0.50		99		
21010011	**Good identification**						**21010011**
					SPO		
Clostridium novyi biovar A		97.03	0.50	NIT 1	99		
Clostridium difficile		1.19	0.02	FRU 10	99		
Pseudoramibacter alactolyticus		1.03	-.03		1		
21010012	**Low discrimination**						**21010012**
					SPO		
Clostridium novyi biovar A		42.48	0.37	NIT 1	99		
Pseudoramibacter alactolyticus		28.19	0.15	FRU 10	1		
Atopobium minutum		23.43	0.28		1		
21010013	**Good identification**						**21010013**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
21010020	**Low discrimination**						**21010020**
					CAT		
Propionibacterium acnes		89.74	0.79	IND 87	95		
Pseudoramibacter alactolyticus		7.36	0.53		1		
Atopobium minutum		1.93	0.57		1		
21010021	**Low discrimination**						**21010021**
					SPO	CAT	
Propionibacterium acnes		81.63	0.44	IND 87	1	95	
Clostridium novyi biovar A		9.00	0.39	URE 1	99	1	
Pseudoramibacter alactolyticus		6.70	0.19		1	1	
21010022	**Low discrimination**						**21010022**
					CAT		
Atopobium minutum		41.37	0.54	NIT 1	1		
Propionibacterium acnes		29.20	0.44		95		
Pseudoramibacter alactolyticus		26.35	0.37		1		
21010030	**Low discrimination**						**21010030**
					SPO	CAT	
Propionibacterium acnes		81.50	0.44	IND 87	1	95	
Clostridium novyi biovar A		8.99	0.39	MLZ 1	99	1	
Pseudoramibacter alactolyticus		6.69	0.19		1	1	
21010031	**Acceptable identification**						**21010031**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		91.61	0.39	NIT 1	99	1		
Propionibacterium acnes		8.39	0.10	FRU 10	1	95		
21010040	**Low discrimination**							**21010040**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		68.77	0.56	NIT 1	1	1	99	
Megasphaera elsdenii		19.30	0.57		99	1	1	
Clostridium glycolicum		6.52	0.43		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
21010041	**Low discrimination**							**21010041**
					COC	SPO	GRA	
Clostridium novyi biovar A		63.71	0.47	NIT 1	1	99	99	
Pseudoramibacter alactolyticus		25.18	0.21	FRU 10	1	1	99	
Megasphaera elsdenii		7.07	0.23		99	1	1	
21010042	**Low discrimination**							**21010042**
					COC	SPO	GRA	
Pseudoramibacter alactolyticus		73.16	0.39	NIT 1	1	1	99	
Megasphaera elsdenii		20.54	0.41	BGL 10	99	1	1	
Clostridium novyi biovar A		2.97	0.34		1	99	99	
21010050	**Low discrimination**							**21010050**
					COC	SPO	GRA	
Clostridium novyi biovar A		60.48	0.47	NIT 1	1	99	99	
Pseudoramibacter alactolyticus		23.90	0.21	FRU 10	1	1	99	
Megasphaera elsdenii		6.71	0.23		99	1	1	
21010051	**Good identification**							**21010051**
Clostridium novyi biovar A		100.0	0.47	NIT 1				
21010060	**Good identification**							**21010060**
					SPO	CAT		
Propionibacterium acnes		90.89	0.67	IND 87	1	95		
Pseudoramibacter alactolyticus		8.49	0.43		1	1		
Clostridium novyi biovar A		0.31	0.36		99	1		
21010071	**Good identification**							**21010071**
					SPO	CAT		
Clostridium novyi biovar A		97.07	0.36	NIT 1	99	1		
Propionibacterium acnes		2.93	-.01	FRU 10	1	95		
21010120	**Low discrimination**							**21010120**
					CAT			
Propionibacterium acnes		87.15	0.44	IND 87	95			
Pseudoramibacter alactolyticus		7.15	0.19	LAC 1	1			
Actinomyces meyeri		3.15	0.23		1			
21010220	**Low discrimination**							**21010220**
					CAT			
Propionibacterium acnes		89.50	0.44	IND 87	95			
Pseudoramibacter alactolyticus		7.34	0.19	RHA 1	1			
Atopobium minutum		1.92	0.23		1			
21010420	**Low discrimination**							**21010420**
					CAT			
Propionibacterium acnes		89.80	0.44	IND 87	95			
Pseudoramibacter alactolyticus		7.37	0.19	ARA 1	1			
Atopobium minutum		1.93	0.23		1			
21011000	**Low discrimination**							**21011000**
					CAT			
Atopobium minutum		52.78	0.57	NIT 1	1			
Propionibacterium acnes		24.84	0.44		95			
Pseudoramibacter alactolyticus		11.54	0.32		1			
21011002	**Acceptable identification**							**21011002**
Atopobium minutum		91.02	0.54	NIT 1				
Pseudoramibacter alactolyticus		3.32	0.15					
Lactobacillus fermentum		3.25	0.36					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
21011020	**Very good identification**						**21011020**
					CAT		
Propionibacterium acnes		99.16	0.79	IND 87	95		
Atopobium minutum		0.71	0.49		1		
Pseudoramibacter alactolyticus		0.08	0.19		1		
21011021	**Good identification**						**21011021**
					SPO	CAT	
Propionibacterium acnes		98.10	0.44	IND 87	1	95	
Clostridium novyi biovar A		1.20	0.23	URE 1	99	1	
Atopobium minutum		0.70	0.15		1	1	
21011022	**Low discrimination**						**21011022**
					CAT		
Propionibacterium acnes		66.40	0.44	IND 87	95		
Atopobium minutum		31.35	0.46	BGL 1	1		
Lactobacillus fermentum		1.12	0.28		1		
21011030	**Good identification**						**21011030**
					SPO	CAT	
Propionibacterium acnes		98.10	0.44	IND 87	1	95	
Clostridium novyi biovar A		1.20	0.23	MLZ 1	99	1	
Atopobium minutum		0.70	0.15		1	1	
21011060	**Very good identification**						**21011060**
					SPO	CAT	
Propionibacterium acnes		99.84	0.67	IND 87	1	95	
Pseudoramibacter alactolyticus		0.09	0.09		1	1	
Clostridium novyi biovar A		0.04	0.20		99	1	
21011100	**Low discrimination**						**21011100**
					GRA		
Actinomyces meyeri		69.32	0.41	NIT 1	99		
Lactobacillus fermentum		14.37	0.40	SUC 93	99		
Fusobacterium mortiferum		9.30	0.35		1		
21011102	**Low discrimination**						**21011102**
Lactobacillus fermentum		70.21	0.40	NIT 1			
Atopobium minutum		12.18	0.20	SUC 87			
Actinomyces meyeri		10.48	0.15				
21011120	**Acceptable identification**						**21011120**
					CAT		
Propionibacterium acnes		91.63	0.44	IND 87	95		
Actinomyces meyeri		4.97	0.26	LAC 1	1		
Lactobacillus fermentum		2.52	0.32		1		
21011220	**Good identification**						**21011220**
					CAT		
Propionibacterium acnes		98.90	0.44	IND 87	95		
Atopobium minutum		0.71	0.15	RHA 1	1		
Actinomyces odontolyticus		0.40	0.10		10		
21011400	**Low discrimination**						**21011400**
Lactobacillus fermentum		45.27	0.36	NIT 1			
Actinomyces meyeri		22.27	0.20	SUC 87			
Atopobium minutum		19.21	0.23				
21011402	**Low discrimination**						**21011402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		77.95	0.36	NIT 1			
Atopobium minutum		22.05	0.20	SUC 87			
21011420	**Good identification**						**21011420**
					CAT		
Propionibacterium acnes		97.33	0.44	IND 87	95		
Lactobacillus fermentum		1.64	0.28	ARA 1	1		
Atopobium minutum		0.70	0.15		1		
21011502	**Good identification**						**21011502**
Lactobacillus fermentum		96.41	0.40	NIT 1			
Actinomyces meyeri		3.60	0.05	SUC 87			
21012000	**Low discrimination**						**21012000**
					COC	GRA	
Pseudoramibacter alactolyticus		52.61	0.61	NIT 1	1	99	
Eubacterium limosum		33.46	0.60		1	99	
Megasphaera elsdenii		10.28	0.60		99	1	
21012002	**Low discrimination**						**21012002**
					COC	GRA	
Pseudoramibacter alactolyticus		78.46	0.45	NIT 1	1	99	
Megasphaera elsdenii		15.34	0.44	BGL 10	99	1	
Eubacterium limosum		4.54	0.26		1	99	
21012010	**Acceptable identification**						**21012010**
					SPO		
Clostridium difficile		91.11	0.53	NIT 1	99		
Pseudoramibacter alactolyticus		4.75	0.27		1		
Eubacterium limosum		3.02	0.26		1		
21012020	**Good identification**						**21012020**
					CAT		
Propionibacterium acnes		92.34	0.74	IND 87	95		
Pseudoramibacter alactolyticus		7.25	0.48		1		
Eubacterium limosum		0.26	0.26		1		
21012021	**Acceptable identification**						**21012021**
					CAT		
Propionibacterium acnes		92.72	0.40	IND 87	95		
Pseudoramibacter alactolyticus		7.28	0.14	URE 1	1		
21012022	**Low discrimination**						**21012022**
					CAT		
Propionibacterium acnes		52.67	0.40	IND 87	95		
Pseudoramibacter alactolyticus		45.49	0.32	BGL 1	1		
Atopobium minutum		1.34	0.20		1		
21012030	**Low discrimination**						**21012030**
					SPO	CAT	
Propionibacterium acnes		85.85	0.40	IND 87	1	95	
Clostridium difficile		7.41	0.19	MLZ 1	99	1	
Pseudoramibacter alactolyticus		6.74	0.14		1	1	
21012040	**Low discrimination**						**21012040**
					COC	GRA	
Pseudoramibacter alactolyticus		63.38	0.51	NIT 1	1	99	
Megasphaera elsdenii		33.04	0.57		99	1	
Eubacterium limosum		1.63	0.26		1	99	
21012060	**Good identification**						**21012060**
					COC	CAT	GRA

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		91.57	0.63	IND 87	1	95	99	
Pseudoramibacter alactolyticus		8.19	0.38		1	1	99	
Megasphaera elsdenii		0.24	0.23		99	1	1	
21012120	**Acceptable identification**							**21012120**
					CAT			
Propionibacterium acnes		92.72	0.40	IND 87	95			
Pseudoramibacter alactolyticus		7.28	0.14	LAC 1	1			
21012220	**Acceptable identification**							**21012220**
					CAT			
Propionibacterium acnes		92.72	0.40	IND 87	95			
Pseudoramibacter alactolyticus		7.28	0.14	RHA 1	1			
21012420	**Acceptable identification**							**21012420**
					CAT			
Propionibacterium acnes		92.72	0.40	IND 87	95			
Pseudoramibacter alactolyticus		7.28	0.14	ARA 1	1			
21013000	**Low discrimination**							**21013000**
					CAT			
Propionibacterium acnes		53.46	0.40	IND 87	95			
Pseudoramibacter alactolyticus		23.78	0.27	NAG 99	1			
Eubacterium limosum		15.13	0.26		1			
21013020	**Very good identification**							**21013020**
					CAT			
Propionibacterium acnes		99.91	0.74	IND 87	95			
Pseudoramibacter alactolyticus		0.08	0.14		1			
Atopobium minutum		0.01	0.15		1			
21013021	**Good identification**							**21013021**
Propionibacterium acnes		100.0	0.40	IND 87				
21013022	**Good identification**							**21013022**
					CAT			
Propionibacterium acnes		98.31	0.40	IND 87	95			
Pseudoramibacter alactolyticus		0.86	-.02	BGL 1	1			
Atopobium minutum		0.83	0.12		1			
21013030	**Good identification**							**21013030**
Propionibacterium acnes		100.0	0.40	IND 87				
21013060	**Very good identification**							**21013060**
					CAT			
Propionibacterium acnes		99.91	0.63	IND 87	95			
Pseudoramibacter alactolyticus		0.09	0.04		1			
21013120	**Good identification**							**21013120**
Propionibacterium acnes		100.0	0.40	IND 87				
21013220	**Good identification**							**21013220**
Propionibacterium acnes		100.0	0.40	IND 87				
21013420	**Good identification**							**21013420**
Propionibacterium acnes		100.0	0.40	IND 87				
21014000	**Low discrimination**							**21014000**
					GRA	SPO		
Clostridium glycolicum		50.39	0.43	NIT 1	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Pseudoramibacter alactolyticus		21.49	0.32	SOR 95	99	1		
Fusobacterium varium		12.45	0.28		1	1		
21014020	**Low discrimination**							**21014020**
					CAT			
Propionibacterium acnes		87.52	0.44	IND 87	95			
Pseudoramibacter alactolyticus		7.18	0.19	XYL 1	1			
Atopobium minutum		1.88	0.23		1			
21014040	**Very good identification**							**21014040**
					COC	SPO	GRA	
Clostridium glycolicum		99.24	0.65	NIT 1	1	99	99	
Pseudoramibacter alactolyticus		0.56	0.21		1	1	99	
Megasphaera elsdenii		0.16	0.23		99	1	1	
21014100	**Good identification**							**21014100**
					SPO			
Actinomyces meyeri		95.54	0.43	NIT 1	1			
Clostridium glycolicum		2.04	0.09	SUC 93	99			
Pseudoramibacter alactolyticus		0.87	-.03		1			
21015000	**Low discrimination**							**21015000**
					SPO			
Actinomyces meyeri		49.63	0.35	NIT 1	1			
Clostridium glycolicum		31.12	0.27	SUC 93	99			
Lactobacillus fermentum		7.63	0.32		1			
21015020	**Good identification**							**21015020**
					CAT			
Propionibacterium acnes		95.84	0.44	IND 87	95			
Actinomyces meyeri		2.52	0.21	XYL 1	1			
Lactobacillus fermentum		0.95	0.24		1			
21015040	**Good identification**							**21015040**
					SPO	CAT		
Clostridium glycolicum		99.90	0.49	NIT 1	99	1		
Propionibacterium acnes		0.10	-.01	GAL 10	1	95		
21015100	**Acceptable identification**							**21015100**
					GRA			
Actinomyces meyeri		93.64	0.46	NIT 1	99			
Lactobacillus fermentum		5.87	0.36	SUC 93	99			
Fusobacterium mortiferum		0.49	0.16		1			
21015102	**Low discrimination**							**21015102**
Lactobacillus fermentum		66.98	0.36	NIT 1				
Actinomyces meyeri		33.03	0.20	SUC 87				
21015500	**Low discrimination**							**21015500**
Actinomyces meyeri		79.94	0.35	NIT 1				
Lactobacillus fermentum		20.06	0.36	SUC 93				
21015502	**Low discrimination**							**21015502**
Lactobacillus fermentum		89.03	0.36	NIT 1				
Actinomyces meyeri		10.98	0.10	SUC 87				
21016020	**Acceptable identification**							**21016020**
					CAT			
Propionibacterium acnes		92.72	0.40	IND 87	95			
Pseudoramibacter alactolyticus		7.28	0.14	XYL 1	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
21017020	**Good identification**							**21017020**
Propionibacterium acnes		100.0	0.40	IND 87				
21020000	**Low discrimination**							**21020000**
					COC	GRA		
Clostridium botulinum biovar A		42.67	0.51	NIT 1	1	99		
Veillonella parvula		18.33	0.32		99	1		
Clostridium sporogenes		13.71	0.40		1	99		
21020040	**Acceptable identification**							**21020040**
					COC	GRA		
Clostridium botulinum biovar A		93.02	0.42	NIT 1	1	99		
Clostridium novyi biovar A		3.19	0.29		1	99		
Veillonella parvula		1.35	-.03		99	1		
21030020	**Low discrimination**							**21030020**
					SPO	CAT		
Propionibacterium acnes		88.06	0.44	IND 87	1	95		
Pseudoramibacter alactolyticus		7.22	0.19	TRE 1	1	1		
Clostridium cadaveris		2.64	0.15		99	1		
21031020	**Good identification**							**21031020**
					CAT			
Propionibacterium acnes		99.29	0.44	IND 87	95			
Atopobium minutum		0.71	0.15	TRE 1	1			
21032020	**Acceptable identification**							**21032020**
					CAT			
Propionibacterium acnes		92.72	0.40	IND 87	95			
Pseudoramibacter alactolyticus		7.28	0.14	TRE 1	1			
21033020	**Good identification**							**21033020**
Propionibacterium acnes		100.0	0.40	IND 87				
21050020	**Low discrimination**							**21050020**
					CAT			
Propionibacterium acnes		89.98	0.44	IND 87	95			
Pseudoramibacter alactolyticus		7.38	0.19	CEL 1	1			
Atopobium minutum		1.93	0.23		1			
21051020	**Good identification**							**21051020**
					CAT			
Propionibacterium acnes		99.29	0.44	IND 87	95			
Atopobium minutum		0.71	0.15	CEL 1	1			
21052020	**Acceptable identification**							**21052020**
					CAT			
Propionibacterium acnes		92.72	0.40	IND 87	95			
Pseudoramibacter alactolyticus		7.28	0.14	CEL 1	1			
21053020	**Good identification**							**21053020**
Propionibacterium acnes		100.0	0.40	IND 87				
21100000	**Low discrimination**							**21100000**
					COC	GRA	SPO	
Peptostreptococcus anaerobius		41.38	0.63	NIT 1	99	99	1	
Prevotella disiens		22.80	0.57		1	1	1	
Clostridium botulinum biovar A		14.63	0.60		1	99	99	
21100001	**Low discrimination**							**21100001**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		75.84	0.66	NIT 1	99	99		
Clostridium sordellii		7.32	0.32		99	99		
Campylobacter ureolyticus		6.44	0.26		1	1		
21100002	**Low discrimination**							**21100002**
					COC	SPO		
Clostridium sporogenes		51.53	0.45	NIT 1	1	99		
Clostridium novyi biovar A		23.16	0.53		1	99		
Peptostreptococcus anaerobius		8.08	0.29		99	1		
21100003	**Good identification**							**21100003**
Clostridium novyi biovar A		96.49	0.53	NIT 1				
Clostridium sporogenes		2.17	0.11					
Clostridium sordellii		0.53	-.03					
21100010	**Low discrimination**							**21100010**
					COC	GRA	SPO	
Clostridium novyi biovar A		87.94	0.66	NIT 1	1	99	99	
Peptostreptococcus anaerobius		5.41	0.29		99	99	1	
Prevotella disiens		2.98	0.23		1	1	1	
21100011	**Very good identification**							**21100011**
					GRA	SPO		
Clostridium novyi biovar A		99.76	0.66	NIT 1	99	99		
Clostridium sordellii		0.10	-.03		99	99		
Campylobacter ureolyticus		0.09	-.08		1	1		
21100012	**Good identification**							**21100012**
					COC	SPO		
Clostridium novyi biovar A		97.47	0.53	NIT 1	1	99		
Clostridium sporogenes		2.19	0.11		1	99		
Peptostreptococcus anaerobius		0.34	-.06		99	1		
21100013	**Good identification**							**21100013**
Clostridium novyi biovar A		100.0	0.53	NIT 1				
21100020	**Low discrimination**							**21100020**
					COC	GRA	SPO	
Clostridium novyi biovar A		52.52	0.56	NIT 1	1	99	99	
Peptostreptococcus anaerobius		12.93	0.29		99	99	1	
Campylobacter gracilis		10.91	0.21		1	1	1	
21100021	**Good identification**							**21100021**
					GRA	SPO		
Clostridium novyi biovar A		98.82	0.56	NIT 1	99	99		
Clostridium sordellii		0.39	-.03		99	99		
Campylobacter ureolyticus		0.34	-.08		1	1		
21100022	**Low discrimination**							**21100022**
					GRA	SPO		
Clostridium novyi biovar A		69.28	0.43	NIT 1	99	99		
Campylobacter gracilis		9.06	0.05		1	1		
Actinomyces odontolyticus		6.45	0.21		99	1		
21100023	**Good identification**							**21100023**
Clostridium novyi biovar A		100.0	0.43	NIT 1				
21100030	**Good identification**							**21100030**
					COC	GRA	SPO	
Clostridium novyi biovar A		99.54	0.56	NIT 1	1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptostreptococcus anaerobius		0.25	-.06		99	99	1	
Campylobacter gracilis		0.21	-.13		1	1	1	
21100031	**Good identification**							**21100031**
Clostridium novyi biovar A		100.0	0.56	NIT 1				
21100032	**Good identification**							**21100032**
Clostridium novyi biovar A		100.0	0.43	NIT 1				
21100033	**Good identification**							**21100033**
Clostridium novyi biovar A		100.0	0.43	NIT 1				
21100040	**Low discrimination**							**21100040**
					COC	GRA		
Clostridium novyi biovar A		43.61	0.63	NIT 1	1	99		
Clostridium botulinum biovar A		42.08	0.51		1	99		
Megasphaera elsdenii		4.20	0.38		99	1		
21100041	**Very good identification**							**21100041**
Clostridium novyi biovar A		98.14	0.63	NIT 1				
Clostridium botulinum biovar A		0.96	0.17					
Clostridium sordellii		0.59	0.08					
21100042	**Low discrimination**							**21100042**
					COC	GRA		
Clostridium novyi biovar A		86.59	0.50	NIT 1	1	99		
Megasphaera elsdenii		5.25	0.22		99	1		
Clostridium botulinum biovar A		4.78	0.17		1	99		
21100043	**Good identification**							**21100043**
Clostridium novyi biovar A		100.0	0.50	NIT 1				
21100050	**Very good identification**							**21100050**
					COC	GRA		
Clostridium novyi biovar A		98.85	0.63	NIT 1	1	99		
Clostridium botulinum biovar A		0.96	0.17		1	99		
Megasphaera elsdenii		0.10	0.04		99	1		
21100051	**Very good identification**							**21100051**
Clostridium novyi biovar A		100.0	0.63	NIT 1				
21100052	**Good identification**							**21100052**
Clostridium novyi biovar A		100.0	0.50	NIT 1				
21100053	**Good identification**							**21100053**
Clostridium novyi biovar A		100.0	0.50	NIT 1				
21100060	**Acceptable identification**							**21100060**
					COC	GRA		
Clostridium novyi biovar A		94.66	0.53	NIT 1	1	99		
Clostridium botulinum biovar A		3.69	0.17		1	99		
Megasphaera elsdenii		0.37	0.04		99	1		
21100061	**Good identification**							**21100061**
Clostridium novyi biovar A		100.0	0.53	NIT 1				
21100062	**Good identification**							**21100062**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.40	NIT 1				
21100063	**Good identification**							**21100063**
Clostridium novyi biovar A		100.0	0.40	NIT 1				
21100070	**Good identification**							**21100070**
Clostridium novyi biovar A		100.0	0.53	NIT 1				
21100071	**Good identification**							**21100071**
Clostridium novyi biovar A		100.0	0.53	NIT 1				
21100072	**Good identification**							**21100072**
Clostridium novyi biovar A		100.0	0.40	NIT 1				
21100073	**Good identification**							**21100073**
Clostridium novyi biovar A		100.0	0.40	NIT 1				
21100200	**Low discrimination**							**21100200**
					COC	GRA		
Actinomyces odontolyticus		35.18	0.43	SUC 99	1	99		
Peptostreptococcus anaerobius		27.10	0.29	FRU 92	99	99		
Prevotella disiens		14.94	0.23		1	1		
21100202	**Low discrimination**							**21100202**
					SPO			
Actinomyces odontolyticus		88.49	0.39	SUC 99	1			
Clostridium sporogenes		7.17	0.11	FRU 92	99			
Clostridium novyi biovar A		3.22	0.19		99			
21101000	**Low discrimination**							**21101000**
					COC	GRA	SPO	
Clostridium novyi biovar A		37.07	0.50	NIT 1	1	99	99	
Peptostreptococcus anaerobius		20.54	0.29	GAL 10	99	99	1	
Prevotella disiens		11.32	0.23		1	1	1	
21101001	**Low discrimination**							**21101001**
Clostridium novyi biovar A		84.90	0.50	NIT 1				
Clostridium sordellii		13.02	0.19	GAL 10				
Clostridium chauvoei		0.96	0.06					
21101002	**Low discrimination**							**21101002**
					SPO			
Clostridium novyi biovar A		43.65	0.37	NIT 1	99			
Actinomyces odontolyticus		37.44	0.31	GAL 10	1			
Clostridium sporogenes		8.83	0.11		99			
21101003	**Good identification**							**21101003**
Clostridium novyi biovar A		100.0	0.37	NIT 1				
21101010	**Good identification**							**21101010**
					COC	SPO		
Clostridium novyi biovar A		99.44	0.50	NIT 1	1	99		
Peptostreptococcus anaerobius		0.56	-.06	GAL 10	99	1		
21101011	**Good identification**							**21101011**
Clostridium novyi biovar A		100.0	0.50	NIT 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
21101012	**Good identification**						**21101012**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
21101013	**Good identification**						**21101013**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
21101020	**Low discrimination**						**21101020**
					SPO	CAT	
Clostridium novyi biovar A		70.26	0.39	NIT 1	99	1	
Actinomyces odontolyticus		11.30	0.21	GAL 10	1	10	
Propionibacterium acnes		6.44	0.10		1	95	
21101021	**Good identification**						**21101021**
Clostridium novyi biovar A		100.0	0.39	NIT 1			
21101030	**Good identification**						**21101030**
Clostridium novyi biovar A		100.0	0.39	NIT 1			
21101031	**Good identification**						**21101031**
Clostridium novyi biovar A		100.0	0.39	NIT 1			
21101040	**Low discrimination**						**21101040**
Clostridium novyi biovar A		86.62	0.47	NIT 1			
Clostridium botulinum biovar A		7.60	0.17	GAL 10			
Clostridium glycolicum		4.30	0.11				
21101041	**Good identification**						**21101041**
Clostridium novyi biovar A		99.05	0.47	NIT 1			
Clostridium sordellii		0.95	-.05	GAL 10			
21101050	**Good identification**						**21101050**
Clostridium novyi biovar A		100.0	0.47	NIT 1			
21101051	**Good identification**						**21101051**
Clostridium novyi biovar A		100.0	0.47	NIT 1			
21101060	**Good identification**						**21101060**
					SPO	CAT	
Clostridium novyi biovar A		97.07	0.36	NIT 1	99	1	
Propionibacterium acnes		2.93	-.01	GAL 10	1	95	
21101061	**Good identification**						**21101061**
Clostridium novyi biovar A		100.0	0.36	NIT 1			
21101070	**Good identification**						**21101070**
Clostridium novyi biovar A		100.0	0.36	NIT 1			
21101071	**Good identification**						**21101071**
Clostridium novyi biovar A		100.0	0.36	NIT 1			
21101101	**Good identification**						**21101101**
					COC	SPO	
Clostridium chauvoei		98.41	0.40	SUC 99	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		0.89	0.15	MNS 90	1	99		
Blautia hansenii		0.70	-.03		99	1		
21101200	**Good identification**							**21101200**
					COC	SPO		
Actinomyces odontolyticus		96.83	0.40	SUC 99	1	1		
Clostridium novyi biovar A		2.04	0.15	FRU 92	1	99		
Peptostreptococcus anaerobius		1.13	-.06		99	1		
21101202	**Good identification**							**21101202**
Actinomyces odontolyticus		100.0	0.36	SUC 99				
21102000	**Low discrimination**							**21102000**
					COC	GRA		
Megasphaera elsdenii		39.61	0.41	NIT 1	99	1		
Peptostreptococcus anaerobius		25.31	0.29	FRU 95	99	99		
Prevotella disiens		13.95	0.23		1	1		
21102040	**Low discrimination**							**21102040**
					COC	GRA		
Megasphaera elsdenii		82.26	0.38	NIT 1	99	1		
Clostridium novyi biovar A		8.62	0.29	FRU 95	1	99		
Clostridium botulinum biovar A		8.32	0.17		1	99		
21104040	**Good identification**							**21104040**
Clostridium glycolicum		97.74	0.50	NIT 1				
Clostridium novyi biovar A		1.05	0.29	FRU 90				
Clostridium botulinum biovar A		1.01	0.17					
21104200	**Good identification**							**21104200**
					COC			
Actinomyces odontolyticus		98.45	0.38	SUC 99	1			
Peptostreptococcus anaerobius		1.56	-.06	FRU 92	99			
21110000	**Low discrimination**							**21110000**
					COC	GRA		
Prevotella disiens		51.50	0.57	NIT 1	1	1		
Megasphaera elsdenii		28.07	0.63		99	1		
Actinomyces odontolyticus		7.26	0.56		1	99		
21110001	**Low discrimination**							**21110001**
					GRA	SPO		
Clostridium novyi biovar A		40.86	0.50	NIT 1	99	99		
Clostridium sordellii		35.49	0.32	FRU 10	99	99		
Prevotella disiens		12.47	0.23		1	1		
21110002	**Low discrimination**							**21110002**
					COC	SPO	GRA	
Actinomyces odontolyticus		44.60	0.52	SUC 99	1	1	99	
Megasphaera elsdenii		31.26	0.47		99	1	1	
Clostridium sporogenes		10.65	0.32		1	99	99	
21110003	**Low discrimination**							**21110003**
					SPO			
Clostridium novyi biovar A		68.72	0.37	NIT 1	99			
Actinomyces odontolyticus		10.27	0.18	FRU 10	1			
Actinomyces viscosus		7.93	0.09		1			
21110010	**Low discrimination**							**21110010**
					COC	GRA	SPO	
Clostridium novyi biovar A		63.15	0.50	NIT 1	1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella disiens		19.28	0.23	FRU 10	1	1	1	
Megasphaera elsdenii		10.51	0.29		99	1	1	
21110011	**Good identification**							**21110011**
Clostridium novyi biovar A		99.13	0.50	NIT 1				
Clostridium sordellii		0.87	-.03	FRU 10				
21110012	**Low discrimination**							**21110012**
					COC	SPO	GRA	
Clostridium novyi biovar A		77.52	0.37	NIT 1	1	99	99	
Actinomyces odontolyticus		11.59	0.18	FRU 10	1	1	99	
Megasphaera elsdenii		8.12	0.12		99	1	1	
21110013	**Good identification**							**21110013**
Clostridium novyi biovar A		100.0	0.37	NIT 1				
21110020	**Low discrimination**							**21110020**
					GRA	CAT		
Propionibacterium acnes		55.32	0.44	IND 87	99	95		
Actinomyces odontolyticus		16.92	0.43	MLT 1	99	10		
Prevotella disiens		7.45	0.23		1	1		
21110021	**Low discrimination**							**21110021**
					SPO	CAT		
Clostridium novyi biovar A		86.60	0.39	NIT 1	99	1		
Propionibacterium acnes		7.94	0.10	FRU 10	1	95		
Clostridium sordellii		3.04	-.03		99	1		
21110022	**Low discrimination**							**21110022**
					SPO			
Actinomyces odontolyticus		74.58	0.39	SUC 99	1			
Clostridium novyi biovar A		7.74	0.26	NAG 14	99			
Atopobium minutum		5.69	0.20		1			
21110030	**Low discrimination**							**21110030**
					SPO	CAT		
Clostridium novyi biovar A		89.31	0.39	NIT 1	99	1		
Propionibacterium acnes		8.18	0.10	FRU 10	1	95		
Actinomyces odontolyticus		2.50	0.08		1	10		
21110031	**Good identification**							**21110031**
Clostridium novyi biovar A		100.0	0.39	NIT 1				
21110040	**Low discrimination**							**21110040**
					COC	GRA		
Megasphaera elsdenii		71.46	0.60	NIT 1	99	1		
Clostridium glycolicum		17.42	0.44		1	99		
Clostridium novyi biovar A		4.34	0.47		1	99		
21110041	**Low discrimination**							**21110041**
					COC	GRA		
Clostridium novyi biovar A		79.29	0.47	NIT 1	1	99		
Megasphaera elsdenii		13.19	0.26	FRU 10	99	1		
Clostridium sordellii		4.31	0.08		1	99		
21110042	**Low discrimination**							**21110042**
					COC	SPO	GRA	
Megasphaera elsdenii		85.52	0.44	NIT 1	99	1	1	
Clostridium novyi biovar A		8.24	0.34	BGL 10	1	99	99	
Pseudoramibacter alactolyticus		2.05	0.05		1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
21110050	**Low discrimination**							**21110050**
					COC	GRA		
Clostridium novyi biovar A		81.65	0.47	NIT 1	1	99		
Megasphaera elsdenii		13.59	0.26	FRU 10	99	1		
Clostridium glycolicum		3.31	0.10		1	99		
21110051	**Good identification**							**21110051**
Clostridium novyi biovar A		100.0	0.47	NIT 1				
21110061	**Good identification**							**21110061**
					SPO	CAT		
Clostridium novyi biovar A		97.07	0.36	NIT 1	99	1		
Propionibacterium acnes		2.93	-.01	FRU 10	1	95		
21110070	**Acceptable identification**							**21110070**
					SPO	CAT		
Clostridium novyi biovar A		93.89	0.36	NIT 1	99	1		
Clostridium botulinum biovar B		3.28	-.03	FRU 10	99	1		
Propionibacterium acnes		2.83	-.01		1	95		
21110071	**Good identification**							**21110071**
Clostridium novyi biovar A		100.0	0.36	NIT 1				
21110100	**Low discrimination**							**21110100**
					GRA			
Actinomyces odontolyticus		51.35	0.47	SUC 99	99			
Actinomyces meyeri		25.90	0.39		99			
Prevotella disiens		11.65	0.23		1			
21110102	**Low discrimination**							**21110102**
Actinomyces odontolyticus		78.68	0.44	SUC 99				
Actinomyces viscosus		10.36	0.22					
Propionibacterium propionicum		6.05	0.32					
21110200	**Good identification**							**21110200**
					COC	GRA		
Actinomyces odontolyticus		93.89	0.61	SUC 99	1	99		
Prevotella disiens		3.47	0.23		1	1		
Megasphaera elsdenii		1.89	0.29		99	1		
21110202	**Very good identification**							**21110202**
					COC	SPO	GRA	
Actinomyces odontolyticus		99.51	0.57	SUC 99	1	1	99	
Megasphaera elsdenii		0.36	0.12		99	1	1	
Clostridium sporogenes		0.12	-.02		1	99	99	
21110220	**Good identification**							**21110220**
					CAT			
Actinomyces odontolyticus		98.33	0.47	SUC 99	10			
Propionibacterium acnes		1.67	0.10	NAG 14	95			
21110222	**Good identification**							**21110222**
Actinomyces odontolyticus		100.0	0.44	SUC 99				
21110300	**Good identification**							**21110300**
Actinomyces odontolyticus		99.74	0.52	SUC 99				
Actinomyces meyeri		0.26	0.05					
21110302	**Good identification**							**21110302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.49	SUC 99		
21110320	**Good identification**					**21110320**
Actinomyces odontolyticus		100.0	0.39	SUC 99		
21110322	**Good identification**					**21110322**
Actinomyces odontolyticus		100.0	0.35	SUC 99		
21111000	**Low discrimination**					**21111000**
					GRA	
Actinomyces odontolyticus		54.65	0.53	SUC 99	99	
Prevotella intermedia		16.77	0.39		1	
Lactobacillus fermentum		6.62	0.44		99	
21111002	**Low discrimination**					**21111002**
Actinomyces odontolyticus		78.56	0.49	SUC 99		
Lactobacillus fermentum		15.53	0.44			
Actinomyces viscosus		1.69	0.14			
21111020	**Low discrimination**					**21111020**
					CAT	
Propionibacterium acnes		75.01	0.44	IND 87	95	
Actinomyces odontolyticus		15.29	0.40	MLT 1	10	
Lactobacillus fermentum		3.79	0.36		1	
21111022	**Low discrimination**					**21111022**
Actinomyces odontolyticus		60.16	0.36	SUC 99		
Lactobacillus fermentum		24.34	0.36	NAG 14		
Clostridium perfringens		7.29	0.15			
21111100	**Low discrimination**					**21111100**
Actinomyces meyeri		39.25	0.42	NIT 1		
Actinomyces odontolyticus		34.59	0.44	SUC 93		
Lactobacillus fermentum		21.65	0.48			
21111101	**Good identification**					**21111101**
					SPO	
Clostridium chauvoei		98.87	0.47	SUC 99	99	
Actinomyces meyeri		0.39	0.08	MNS 90	1	
Actinomyces odontolyticus		0.34	0.10		1	
21111102	**Low discrimination**					**21111102**
Lactobacillus fermentum		43.68	0.48	NIT 1		
Actinomyces odontolyticus		42.78	0.41	SUC 87		
Actinomyces viscosus		5.18	0.18			
21111120	**Low discrimination**					**21111120**
Clostridium perfringens		48.81	0.36	SUC 91		
Lactobacillus fermentum		17.06	0.40	MNS 99		
Actinomyces odontolyticus		13.31	0.31			
21111121	**Good identification**					**21111121**
Clostridium chauvoei		99.38	0.38	SUC 99		
Clostridium perfringens		0.62	0.01	MNS 90		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
21111200	**Very good identification**					**21111200**
					GRA	
Actinomyces odontolyticus		99.84	0.58	SUC 99	99	
Prevotella intermedia		0.16	0.05		1	
21111202	**Good identification**					**21111202**
Actinomyces odontolyticus		100.0	0.54	SUC 99		
21111220	**Good identification**					**21111220**
					CAT	
Actinomyces odontolyticus		97.51	0.44	SUC 99	10	
Propionibacterium acnes		2.49	0.10	NAG 14	95	
21111222	**Good identification**					**21111222**
Actinomyces odontolyticus		100.0	0.41	SUC 99		
21111300	**Good identification**					**21111300**
Actinomyces odontolyticus		99.09	0.49	SUC 99		
Actinomyces meyeri		0.59	0.08			
Lactobacillus fermentum		0.32	0.14			
21111302	**Good identification**					**21111302**
Actinomyces odontolyticus		99.47	0.46	SUC 99		
Lactobacillus fermentum		0.53	0.14			
21111320	**Good identification**					**21111320**
Actinomyces odontolyticus		98.13	0.36	SUC 99		
Clostridium perfringens		1.87	0.01	NAG 14		
21111400	**Low discrimination**					**21111400**
Lactobacillus fermentum		74.27	0.44	NIT 1		
Actinomyces meyeri		13.73	0.21	SUC 87		
Actinomyces odontolyticus		6.19	0.19			
21111402	**Good identification**					**21111402**
Lactobacillus fermentum		95.14	0.44	NIT 1		
Actinomyces odontolyticus		4.86	0.15	SUC 87		
21111420	**Low discrimination**					**21111420**
					CAT	
Lactobacillus fermentum		75.98	0.36	NIT 1	1	
Propionibacterium acnes		15.18	0.10	SUC 87	95	
Actinomyces meyeri		5.75	0.06		1	
21111422	**Good identification**					**21111422**
Lactobacillus fermentum		100.0	0.36	NIT 1		
21111500	**Low discrimination**					**21111500**
Lactobacillus fermentum		68.05	0.48	NIT 1		
Actinomyces meyeri		30.85	0.31	SUC 87		
Actinomyces odontolyticus		1.10	0.10			
21111502	**Good identification**					**21111502**
Lactobacillus fermentum		97.66	0.48	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces meyeri		1.37	0.06	SUC 87				
Actinomyces odontolyticus		0.97	0.07					
21111520	**Low discrimination**							**21111520**
Lactobacillus fermentum		82.35	0.40	NIT 1				
Actinomyces meyeri		15.27	0.17	SUC 87				
Clostridium perfringens		2.38	0.01					
21111522	**Good identification**							**21111522**
Lactobacillus fermentum		100.0	0.40	NIT 1				
21112000	**Good identification**							**21112000**
					COC	GRA		
Megasphaera elsdenii		92.11	0.63	NIT 1	99	1		
Pseudoramibacter alactolyticus		3.17	0.27		1	99		
Eubacterium limosum		2.02	0.26		1	99		
21112002	**Acceptable identification**							**21112002**
					COC	GRA		
Megasphaera elsdenii		92.00	0.47	NIT 1	99	1		
Propionibacterium propionicum		3.19	0.32	BGL 10	1	99		
Pseudoramibacter alactolyticus		3.17	0.11		1	99		
21112020	**Low discrimination**							**21112020**
					COC	CAT	GRA	
Propionibacterium acnes		82.34	0.40	IND 87	1	95	99	
Megasphaera elsdenii		10.74	0.29	MLT 1	99	1	1	
Pseudoramibacter alactolyticus		6.47	0.14		1	1	99	
21112040	**Very good identification**							**21112040**
					COC	GRA		
Megasphaera elsdenii		98.05	0.60	NIT 1	99	1		
Pseudoramibacter alactolyticus		1.27	0.17		1	99		
Propionibacterium propionicum		0.38	0.29		1	99		
21112042	**Acceptable identification**							**21112042**
					COC	GRA		
Megasphaera elsdenii		93.91	0.44	NIT 1	99	1		
Propionibacterium propionicum		4.88	0.32	BGL 10	1	99		
Pseudoramibacter alactolyticus		1.21	0.00		1	99		
21112100	**Very good identification**							**21112100**
					COC	GRA		
Propionibacterium propionicum		95.60	0.63	SUC 99	1	99		
Megasphaera elsdenii		3.80	0.29		99	1		
Actinomyces odontolyticus		0.31	0.13		1	99		
21112101	**Good identification**							**21112101**
Propionibacterium propionicum		100.0	0.47	SUC 99				
21112102	**Very good identification**							**21112102**
					COC	GRA		
Propionibacterium propionicum		99.58	0.66	SUC 99	1	99		
Megasphaera elsdenii		0.29	0.12		99	1		
Actinomyces odontolyticus		0.13	0.10		1	99		
21112103	**Good identification**							**21112103**
Propionibacterium propionicum		100.0	0.50	SUC 99				
21112140	**Very good identification**							**21112140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Propionibacterium propionicum		97.42	0.63	SUC 99	1	99	
Megasphaera elsdenii		2.58	0.26		99	1	
21112141	**Good identification**						**21112141**
Propionibacterium propionicum		100.0	0.47	SUC 99			
21112142	**Very good identification**						**21112142**
					COC	GRA	
Propionibacterium propionicum		99.80	0.66	SUC 99	1	99	
Megasphaera elsdenii		0.20	0.09		99	1	
21112143	**Good identification**						**21112143**
Propionibacterium propionicum		100.0	0.50	SUC 99			
21113020	**Good identification**						**21113020**
					CAT		
Propionibacterium acnes		99.64	0.40	IND 87	95		
Actinomyces odontolyticus		0.37	0.05	MLT 1	10		
21113100	**Very good identification**						**21113100**
Propionibacterium propionicum		99.41	0.63	SUC 99			
Actinomyces meyeri		0.24	0.08				
Actinomyces odontolyticus		0.22	0.10				
21113101	**Acceptable identification**						**21113101**
					SPO		
Propionibacterium propionicum		94.64	0.47	SUC 99	1		
Clostridium chauvoei		5.36	0.12	URE 10	99		
21113102	**Very good identification**						**21113102**
Propionibacterium propionicum		99.82	0.66	SUC 99			
Lactobacillus fermentum		0.09	0.14				
Actinomyces odontolyticus		0.09	0.07				
21113103	**Good identification**						**21113103**
					SPO		
Propionibacterium propionicum		99.34	0.50	SUC 99	1		
Clostridium chauvoei		0.66	-.01	URE 10	99		
21113140	**Very good identification**						**21113140**
Propionibacterium propionicum		100.0	0.63	SUC 99			
21113141	**Good identification**						**21113141**
Propionibacterium propionicum		100.0	0.47	SUC 99			
21113142	**Very good identification**						**21113142**
Propionibacterium propionicum		100.0	0.66	SUC 99			
21113143	**Good identification**						**21113143**
Propionibacterium propionicum		100.0	0.50	SUC 99			
21114000	**Low discrimination**						**21114000**
					SPO		
Clostridium glycolicum		47.11	0.44	NIT 1	99		
Actinomyces odontolyticus		36.93	0.51	SOR 95	1		
Actinomyces meyeri		5.79	0.33		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
21114002	**Good identification**							**21114002**
					COC	SPO	GRA	
Actinomyces odontolyticus		95.40	0.47	SUC 99	1	1	99	
Clostridium glycolicum		2.01	0.10		1	99	99	
Megasphaera elsdenii		1.37	0.12		99	1	1	
21114020	**Low discrimination**							**21114020**
					SPO			
Actinomyces odontolyticus		78.27	0.37	SUC 99	1			
Actinomyces meyeri		10.29	0.19	NAG 14	1			
Clostridium glycolicum		6.20	0.10		99			
21114040	**Very good identification**							**21114040**
					COC	SPO	GRA	
Clostridium glycolicum		99.73	0.66	NIT 1	1	99	99	
Megasphaera elsdenii		0.22	0.26		99	1	1	
Actinomyces odontolyticus		0.04	0.17		1	1	99	
21114100	**Low discrimination**							**21114100**
					SPO			
Actinomyces meyeri		65.63	0.44	NIT 1	1			
Actinomyces odontolyticus		33.02	0.42	SUC 93	1			
Clostridium glycolicum		1.35	0.10		99			
21114102	**Acceptable identification**							**21114102**
Actinomyces odontolyticus		90.89	0.39	SUC 99				
Actinomyces meyeri		9.12	0.18					
21114200	**Good identification**							**21114200**
					SPO			
Actinomyces odontolyticus		99.34	0.56	SUC 99	1			
Clostridium glycolicum		0.66	0.10		99			
21114202	**Good identification**							**21114202**
Actinomyces odontolyticus		100.0	0.52	SUC 99				
21114220	**Good identification**							**21114220**
Actinomyces odontolyticus		100.0	0.42	SUC 99				
21114222	**Good identification**							**21114222**
Actinomyces odontolyticus		100.0	0.39	SUC 99				
21114300	**Good identification**							**21114300**
Actinomyces odontolyticus		98.98	0.47	SUC 99				
Actinomyces meyeri		1.02	0.09					
21114302	**Good identification**							**21114302**
Actinomyces odontolyticus		100.0	0.43	SUC 99				
21115000	**Low discrimination**							**21115000**
					SPO			
Actinomyces odontolyticus		57.83	0.48	SUC 99	1			
Actinomyces meyeri		20.42	0.36		1			
Clostridium glycolicum		12.30	0.28		99			
21115002	**Low discrimination**							**21115002**
Actinomyces odontolyticus		79.78	0.44	SUC 99				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		18.80	0.41			
Actinomyces meyeri		1.42	0.11			
21115040	**Good identification**					**21115040**
					SPO	
Clostridium glycolicum		99.66	0.50	NIT 1	99	
Actinomyces odontolyticus		0.25	0.14	GAL 10	1	
Actinomyces meyeri		0.09	0.02		1	
21115100	**Low discrimination**					**21115100**
Actinomyces meyeri		71.92	0.47	NIT 1		
Actinomyces odontolyticus		16.08	0.39	SUC 93		
Lactobacillus fermentum		12.00	0.44			
21115102	**Low discrimination**					**21115102**
Lactobacillus fermentum		49.83	0.44	NIT 1		
Actinomyces odontolyticus		40.93	0.36	SUC 87		
Actinomyces meyeri		9.24	0.21			
21115122	**Low discrimination**					**21115122**
Lactobacillus fermentum		67.71	0.36	NIT 1		
Actinomyces odontolyticus		27.16	0.22	SUC 87		
Actinomyces meyeri		5.13	0.06			
21115200	**Good identification**					**21115200**
Actinomyces odontolyticus		99.82	0.53	SUC 99		
Actinomyces meyeri		0.18	0.02			
21115202	**Good identification**					**21115202**
Actinomyces odontolyticus		100.0	0.49	SUC 99		
21115220	**Good identification**					**21115220**
Actinomyces odontolyticus		100.0	0.39	SUC 99		
21115222	**Good identification**					**21115222**
Actinomyces odontolyticus		100.0	0.36	SUC 99		
21115300	**Good identification**					**21115300**
Actinomyces odontolyticus		97.73	0.44	SUC 99		
Actinomyces meyeri		2.27	0.13			
21115302	**Good identification**					**21115302**
Actinomyces odontolyticus		100.0	0.40	SUC 99		
21115400	**Low discrimination**					**21115400**
Lactobacillus fermentum		59.49	0.41	NIT 1		
Actinomyces meyeri		36.35	0.26	SUC 87		
Actinomyces odontolyticus		4.16	0.14			
21115402	**Good identification**					**21115402**
Lactobacillus fermentum		95.89	0.41	NIT 1		
Actinomyces odontolyticus		4.11	0.10	SUC 87		
21115500	**Low discrimination**					**21115500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		59.97	0.36	NIT 1			
Lactobacillus fermentum		40.03	0.44	SUC 93			
21115502	**Good identification**						**21115502**
Lactobacillus fermentum		95.57	0.44	NIT 1			
Actinomyces meyeri		4.43	0.11	SUC 87			
21115520	**Low discrimination**						**21115520**
Lactobacillus fermentum		62.00	0.36	NIT 1			
Actinomyces meyeri		38.00	0.22	SUC 87			
21115522	**Good identification**						**21115522**
Lactobacillus fermentum		100.0	0.36	NIT 1			
21120000	**Low discrimination**						**21120000**
					COC	SPO	
Clostridium botulinum biovar A		51.54	0.49	NIT 1	1	99	
Clostridium sporogenes		37.59	0.45		1	99	
Peptostreptococcus anaerobius		5.89	0.29		99	1	
21120002	**Acceptable identification**						**21120002**
Clostridium sporogenes		94.35	0.37	NIT 1			
Clostridium botulinum biovar A		3.92	0.15				
Clostridium novyi biovar A		1.29	0.19				
21120040	**Acceptable identification**						**21120040**
Clostridium botulinum biovar A		92.01	0.41	NIT 1			
Clostridium novyi biovar A		3.85	0.29				
Clostridium sporogenes		2.63	0.11				
21131120	**Good identification**						**21131120**
Clostridium perfringens		97.37	0.40	SUC 91			
Clostridium septicum		2.63	0.07	MNS 99			
21132100	**Very good identification**						**21132100**
Propionibacterium propionicum		100.0	0.61	SUC 99			
21132101	**Good identification**						**21132101**
Propionibacterium propionicum		100.0	0.45	SUC 99			
21132102	**Very good identification**						**21132102**
Propionibacterium propionicum		100.0	0.64	SUC 99			
21132103	**Good identification**						**21132103**
Propionibacterium propionicum		100.0	0.48	SUC 99			
21132140	**Very good identification**						**21132140**
Propionibacterium propionicum		100.0	0.61	SUC 99			
21132141	**Good identification**						**21132141**
Propionibacterium propionicum		100.0	0.45	SUC 99			
21132142	**Very good identification**						**21132142**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		100.0	0.64	SUC 99				
21132143	**Good identification**							**21132143**
Propionibacterium propionicum		100.0	0.48	SUC 99				
21133100	**Very good identification**							**21133100**
Propionibacterium propionicum		100.0	0.61	SUC 99				
21133101	**Good identification**							**21133101**
Propionibacterium propionicum		100.0	0.45	SUC 99				
21133102	**Very good identification**							**21133102**
Propionibacterium propionicum		100.0	0.64	SUC 99				
21133103	**Good identification**							**21133103**
Propionibacterium propionicum		100.0	0.48	SUC 99				
21133140	**Very good identification**							**21133140**
Propionibacterium propionicum		100.0	0.61	SUC 99				
21133141	**Good identification**							**21133141**
Propionibacterium propionicum		100.0	0.45	SUC 99				
21133142	**Very good identification**							**21133142**
Propionibacterium propionicum		100.0	0.64	SUC 99				
21133143	**Good identification**							**21133143**
Propionibacterium propionicum		100.0	0.48	SUC 99				
21200001	**Low discrimination**							**21200001**
					COC	GRA	SPO	
Clostridium novyi biovar A		48.91	0.48	NIT 1	1	99	99	
Campylobacter ureolyticus		47.78	0.26	SAL 8	1	1	1	
Veillonella parvula		1.61	-.03		99	1	1	
21200010	**Acceptable identification**							**21200010**
					COC	GRA	SPO	
Clostridium novyi biovar A		93.66	0.48	NIT 1	1	99	99	
Veillonella parvula		3.08	-.03	SAL 8	99	1	1	
Campylobacter gracilis		2.26	-.03		1	1	1	
21200011	**Good identification**							**21200011**
					GRA	SPO		
Clostridium novyi biovar A		99.02	0.48	NIT 1	99	99		
Campylobacter ureolyticus		0.98	-.08	SAL 8	1	1		
21200021	**Acceptable identification**							**21200021**
					GRA	SPO		
Clostridium novyi biovar A		94.02	0.37	NIT 1	99	99		
Campylobacter ureolyticus		3.71	-.08	SAL 8	1	1		
Campylobacter gracilis		2.27	-.13		1	1		
21200030	**Good identification**							**21200030**
					GRA	SPO		
Clostridium novyi biovar A		97.64	0.37	NIT 1	99	99		
Campylobacter gracilis		2.36	-.13	SAL 8	1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
21200031	**Good identification**						**21200031**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
21200040	**Low discrimination**						**21200040**
					COC	GRA	
Clostridium novyi biovar A		80.74	0.45	NIT 1	1	99	
Clostridium botulinum biovar A		11.06	0.18	SAL 8	1	99	
Veillonella parvula		3.98	-.03		99	1	
21200041	**Good identification**						**21200041**
					GRA	SPO	
Clostridium novyi biovar A		98.54	0.45	NIT 1	99	99	
Campylobacter ureolyticus		1.46	-.08	SAL 8	1	1	
21200050	**Good identification**						**21200050**
Clostridium novyi biovar A		100.0	0.45	NIT 1			
21200051	**Good identification**						**21200051**
Clostridium novyi biovar A		100.0	0.45	NIT 1			
21210020	**Low discrimination**						**21210020**
					CAT		
Propionibacterium acnes		88.74	0.44	IND 87	95		
Pseudoramibacter alactolyticus		7.28	0.19	SAL 1	1		
Atopobium minutum		1.91	0.23		1		
21210100	**Low discrimination**						**21210100**
					GRA		
Fusobacterium mortiferum		78.43	0.37	NIT 1	1		
Actinomyces odontolyticus		13.20	0.18		99		
Pseudoramibacter alactolyticus		5.33	-.03		99		
21211020	**Good identification**						**21211020**
					CAT		
Propionibacterium acnes		98.90	0.44	IND 87	95		
Atopobium minutum		0.71	0.15	SAL 1	1		
Actinomyces odontolyticus		0.40	0.10		10		
21211100	**Low discrimination**						**21211100**
					GRA		
Fusobacterium mortiferum		89.80	0.40	NIT 1	1		
Actinomyces odontolyticus		6.72	0.15		99		
Actinomyces meyeri		3.48	0.07		99		
21212010	**Good identification**						**21212010**
					SPO		
Clostridium difficile		99.81	0.43	NIT 1	99		
Pseudoramibacter alactolyticus		0.19	-.07		1		
21212020	**Acceptable identification**						**21212020**
					SPO	CAT	
Propionibacterium acnes		92.29	0.40	IND 87	1	95	
Pseudoramibacter alactolyticus		7.25	0.14	SAL 1	1	1	
Clostridium difficile		0.46	-.03		99	1	
21213020	**Good identification**						**21213020**
Propionibacterium acnes		100.0	0.40	IND 87			
21300000	**Low discrimination**						**21300000**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		26.78	0.48	NIT 1	1	99	
Actinomyces odontolyticus		24.61	0.43	SAL 8	1	1	
Peptostreptococcus anaerobius		18.95	0.29		99	1	
21300001	**Good identification**						**21300001**
					GRA	SPO	
Clostridium novyi biovar A		96.39	0.48	NIT 1	99	99	
Clostridium sordellii		1.08	-.03	SAL 8	99	99	
Campylobacter ureolyticus		0.95	-.08		1	1	
21300002	**Low discrimination**						**21300002**
					SPO		
Actinomyces odontolyticus		71.07	0.39	SUC 99	1		
Clostridium novyi biovar A		22.27	0.35	FRU 92	99		
Clostridium sporogenes		5.76	0.11		99		
21300010	**Good identification**						**21300010**
					COC	SPO	
Clostridium novyi biovar A		98.38	0.48	NIT 1	1	99	
Actinomyces odontolyticus		0.91	0.09	SAL 8	1	1	
Peptostreptococcus anaerobius		0.70	-.06		99	1	
21300011	**Good identification**						**21300011**
Clostridium novyi biovar A		100.0	0.48	NIT 1			
21300020	**Low discrimination**						**21300020**
					SPO		
Clostridium novyi biovar A		58.20	0.37	NIT 1	99		
Actinomyces odontolyticus		34.83	0.29	SAL 8	1		
Clostridium bifermentans		3.90	0.05		99		
21300021	**Good identification**						**21300021**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
21300030	**Good identification**						**21300030**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
21300031	**Good identification**						**21300031**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
21300040	**Low discrimination**						**21300040**
Clostridium novyi biovar A		85.89	0.45	NIT 1			
Clostridium botulinum biovar A		9.63	0.17	SAL 8			
Clostridium bifermentans		1.41	0.02				
21300041	**Good identification**						**21300041**
Clostridium novyi biovar A		100.0	0.45	NIT 1			
21300050	**Good identification**						**21300050**
Clostridium novyi biovar A		100.0	0.45	NIT 1			
21300051	**Good identification**						**21300051**
Clostridium novyi biovar A		100.0	0.45	NIT 1			
21300200	**Good identification**						**21300200**
					COC	SPO	
Actinomyces odontolyticus		99.04	0.48	SUC 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		0.56	0.14	FRU 92	1	99	
Peptostreptococcus anaerobius		0.40	-.06		99	1	
21300202	**Good identification**						**21300202**
Actinomyces odontolyticus		100.0	0.44	SUC 99			
21300300	**Good identification**						**21300300**
Actinomyces odontolyticus		100.0	0.39	SUC 99			
21300302	**Good identification**						**21300302**
Actinomyces odontolyticus		100.0	0.36	SUC 99			
21301000	**Low discrimination**						**21301000**
					SPO		
Actinomyces odontolyticus		82.96	0.40	SUC 99	1		
Clostridium novyi biovar A		15.05	0.31	FRU 92	99		
Eubacterium tortuosum		1.03	-.04		1		
21301002	**Good identification**						**21301002**
					SPO		
Actinomyces odontolyticus		95.04	0.36	SUC 99	1		
Clostridium novyi biovar A		4.96	0.19	FRU 92	99		
21301200	**Good identification**						**21301200**
Actinomyces odontolyticus		100.0	0.45	SUC 99			
21301202	**Good identification**						**21301202**
Actinomyces odontolyticus		100.0	0.41	SUC 99			
21301300	**Good identification**						**21301300**
Actinomyces odontolyticus		100.0	0.36	SUC 99			
21304000	**Good identification**						**21304000**
					COC	SPO	
Actinomyces odontolyticus		96.33	0.38	SUC 99	1	1	
Clostridium novyi biovar A		2.15	0.14	FRU 92	1	99	
Peptostreptococcus anaerobius		1.52	-.06		99	1	
21304200	**Good identification**						**21304200**
Actinomyces odontolyticus		100.0	0.42	SUC 99			
21304202	**Good identification**						**21304202**
Actinomyces odontolyticus		100.0	0.39	SUC 99			
21305200	**Good identification**						**21305200**
Actinomyces odontolyticus		100.0	0.39	SUC 99			
21305202	**Good identification**						**21305202**
Actinomyces odontolyticus		100.0	0.36	SUC 99			
21310000	**Good identification**						**21310000**
					COC	GRA	
Actinomyces odontolyticus		92.88	0.61	SUC 99	1	99	
Prevotella disiens		3.43	0.23		1	1	
Megasphaera elsdenii		1.87	0.29		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
21310002	**Very good identification**						**21310002**
					COC	GRA	
Actinomyces odontolyticus		98.90	0.57	SUC 99	1	99	
Megasphaera elsdenii		0.36	0.12		99	1	
Actinomyces viscosus		0.32	0.08		1	99	
21310020	**Good identification**						**21310020**
					SPO	CAT	
Actinomyces odontolyticus		96.79	0.47	SUC 99	1	10	
Propionibacterium acnes		1.65	0.10	NAG 14	1	95	
Clostridium novyi biovar A		1.56	0.21		99	1	
21310022	**Good identification**						**21310022**
Actinomyces odontolyticus		100.0	0.44	SUC 99			
21310100	**Good identification**						**21310100**
					GRA		
Actinomyces odontolyticus		99.01	0.52	SUC 99	99		
Fusobacterium mortiferum		0.73	0.22		1		
Actinomyces meyeri		0.26	0.05		99		
21310102	**Good identification**						**21310102**
Actinomyces odontolyticus		98.23	0.49	SUC 99			
Actinomyces viscosus		1.77	0.12				
21310120	**Good identification**						**21310120**
Actinomyces odontolyticus		100.0	0.39	SUC 99			
21310122	**Good identification**						**21310122**
Actinomyces odontolyticus		100.0	0.35	SUC 99			
21310200	**Very good identification**						**21310200**
Actinomyces odontolyticus		100.0	0.66	SUC 99			
21310202	**Very good identification**						**21310202**
Actinomyces odontolyticus		100.0	0.62	SUC 99			
21310220	**Good identification**						**21310220**
Actinomyces odontolyticus		100.0	0.52	SUC 99			
21310222	**Good identification**						**21310222**
Actinomyces odontolyticus		100.0	0.49	SUC 99			
21310300	**Very good identification**						**21310300**
Actinomyces odontolyticus		100.0	0.57	SUC 99			
21310302	**Good identification**						**21310302**
Actinomyces odontolyticus		100.0	0.54	SUC 99			
21310320	**Good identification**						**21310320**
Actinomyces odontolyticus		100.0	0.44	SUC 99			
21310322	**Good identification**						**21310322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.40	SUC 99			
21311000	**Very good identification**						**21311000**
					GRA	SPO	
Actinomyces odontolyticus		99.44	0.58	SUC 99	99	1	
Fusobacterium mortiferum		0.23	0.18		1	1	
Clostridium novyi biovar A		0.17	0.15		99	99	
21311002	**Good identification**						**21311002**
Actinomyces odontolyticus		99.71	0.54	SUC 99			
Actinomyces viscosus		0.29	0.04				
21311020	**Good identification**						**21311020**
					CAT		
Actinomyces odontolyticus		95.62	0.44	SUC 99	10		
Propionibacterium acnes		2.44	0.10	NAG 14	95		
Clostridium perfringens		1.94	0.10		1		
21311022	**Good identification**						**21311022**
Actinomyces odontolyticus		98.46	0.41	SUC 99			
Clostridium perfringens		1.54	0.05	NAG 14			
21311100	**Good identification**						**21311100**
					GRA		
Actinomyces odontolyticus		97.48	0.49	SUC 99	99		
Fusobacterium mortiferum		1.63	0.25		1		
Actinomyces meyeri		0.58	0.08		99		
21311102	**Good identification**						**21311102**
Actinomyces odontolyticus		97.86	0.46	SUC 99			
Actinomyces viscosus		1.62	0.08				
Lactobacillus fermentum		0.52	0.14				
21311120	**Low discrimination**						**21311120**
					SPO		
Actinomyces odontolyticus		62.55	0.36	SUC 99	1		
Clostridium perfringens		29.54	0.25	NAG 14	84		
Clostridium septicum		7.91	0.09		99		
21311200	**Very good identification**						**21311200**
Actinomyces odontolyticus		100.0	0.63	SUC 99			
21311202	**Very good identification**						**21311202**
Actinomyces odontolyticus		100.0	0.59	SUC 99			
21311220	**Good identification**						**21311220**
Actinomyces odontolyticus		100.0	0.49	SUC 99			
21311222	**Good identification**						**21311222**
Actinomyces odontolyticus		100.0	0.46	SUC 99			
21311300	**Good identification**						**21311300**
Actinomyces odontolyticus		100.0	0.54	SUC 99			
21311302	**Good identification**						**21311302**
Actinomyces odontolyticus		100.0	0.51	SUC 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
21311320	**Good identification**					**21311320**
Actinomyces odontolyticus		100.0	0.41	SUC 99		
21311322	**Good identification**					**21311322**
Actinomyces odontolyticus		100.0	0.37	SUC 99		
21314000	**Good identification**					**21314000**
					SPO	
Actinomyces odontolyticus		99.34	0.56	SUC 99	1	
Clostridium glycolicum		0.66	0.10		99	
21314002	**Good identification**					**21314002**
Actinomyces odontolyticus		100.0	0.52	SUC 99		
21314020	**Good identification**					**21314020**
Actinomyces odontolyticus		100.0	0.42	SUC 99		
21314022	**Good identification**					**21314022**
Actinomyces odontolyticus		100.0	0.39	SUC 99		
21314100	**Good identification**					**21314100**
Actinomyces odontolyticus		98.98	0.47	SUC 99		
Actinomyces meyeri		1.02	0.09			
21314102	**Good identification**					**21314102**
Actinomyces odontolyticus		100.0	0.43	SUC 99		
21314200	**Very good identification**					**21314200**
Actinomyces odontolyticus		100.0	0.61	SUC 99		
21314202	**Very good identification**					**21314202**
Actinomyces odontolyticus		100.0	0.57	SUC 99		
21314220	**Good identification**					**21314220**
Actinomyces odontolyticus		100.0	0.47	SUC 99		
21314222	**Good identification**					**21314222**
Actinomyces odontolyticus		100.0	0.43	SUC 99		
21314300	**Good identification**					**21314300**
Actinomyces odontolyticus		100.0	0.52	SUC 99		
21314302	**Good identification**					**21314302**
Actinomyces odontolyticus		100.0	0.48	SUC 99		
21314320	**Good identification**					**21314320**
Actinomyces odontolyticus		100.0	0.39	SUC 99		
21315000	**Good identification**					**21315000**
Actinomyces odontolyticus		99.82	0.53	SUC 99		
Actinomyces meyeri		0.18	0.02			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
21315002	**Good identification**					**21315002**
Actinomyces odontolyticus		100.0	0.49	SUC 99		
21315020	**Good identification**					**21315020**
Actinomyces odontolyticus		100.0	0.39	SUC 99		
21315022	**Good identification**					**21315022**
Actinomyces odontolyticus		100.0	0.36	SUC 99		
21315100	**Good identification**					**21315100**
Actinomyces odontolyticus		97.73	0.44	SUC 99		
Actinomyces meyeri		2.27	0.13			
21315102	**Good identification**					**21315102**
Actinomyces odontolyticus		100.0	0.40	SUC 99		
21315200	**Very good identification**					**21315200**
Actinomyces odontolyticus		100.0	0.58	SUC 99		
21315202	**Good identification**					**21315202**
Actinomyces odontolyticus		100.0	0.54	SUC 99		
21315220	**Good identification**					**21315220**
Actinomyces odontolyticus		100.0	0.44	SUC 99		
21315222	**Good identification**					**21315222**
Actinomyces odontolyticus		100.0	0.40	SUC 99		
21315300	**Good identification**					**21315300**
Actinomyces odontolyticus		100.0	0.49	SUC 99		
21315302	**Good identification**					**21315302**
Actinomyces odontolyticus		100.0	0.45	SUC 99		
21315320	**Good identification**					**21315320**
Actinomyces odontolyticus		100.0	0.36	SUC 99		
21377100	**Good identification**					**21377100**
Actinomyces israelii		100.0	0.42	SUC 98		
21377102	**Good identification**					**21377102**
Actinomyces israelii		100.0	0.45	SUC 98		
21377142	**Good identification**					**21377142**
Actinomyces israelii		100.0	0.38	SUC 98		
21377500	**Good identification**					**21377500**
Actinomyces israelii		100.0	0.40	SUC 98		
21377502	**Good identification**					**21377502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.43	SUC 98		
21377542	**Good identification**					**21377542**
Actinomyces israelii		100.0	0.36	SUC 98		
21410020	**Low discrimination**					**21410020**
					CAT	
Propionibacterium acnes		89.98	0.44	IND 87	95	
Pseudoramibacter alactolyticus		7.38	0.19	RAF 1	1	
Atopobium minutum		1.93	0.23		1	
21410100	**Acceptable identification**					**21410100**
					GRA	
Fusobacterium mortiferum		90.36	0.37	NIT 1	1	
Pseudoramibacter alactolyticus		6.14	-.03		99	
Actinomyces meyeri		3.50	0.04		99	
21411000	**Low discrimination**					**21411000**
					GRA	
Lactobacillus fermentum		48.81	0.40	NIT 1	99	
Fusobacterium mortiferum		26.29	0.34	SUC 87	1	
Atopobium minutum		12.70	0.23		99	
21411002	**Low discrimination**					**21411002**
					GRA	
Lactobacillus fermentum		79.06	0.40	NIT 1	99	
Atopobium minutum		13.71	0.20	SUC 87	99	
Fusobacterium mortiferum		4.73	0.17		1	
21411020	**Good identification**					**21411020**
					CAT	
Propionibacterium acnes		96.65	0.44	IND 87	95	
Lactobacillus fermentum		2.66	0.32	RAF 1	1	
Atopobium minutum		0.69	0.15		1	
21411100	**Low discrimination**					**21411100**
					GRA	
Lactobacillus fermentum		54.77	0.44	NIT 1	99	
Fusobacterium mortiferum		42.18	0.40	SUC 87	1	
Actinomyces meyeri		1.64	0.07		99	
21411102	**Low discrimination**					**21411102**
					GRA	
Lactobacillus fermentum		89.28	0.44	NIT 1	99	
Fusobacterium mortiferum		7.64	0.24	SUC 87	1	
Actinomyces viscosus		3.08	0.04		99	
21411120	**Low discrimination**					**21411120**
					CAT	
Lactobacillus fermentum		81.63	0.35	NIT 1	1	
Propionibacterium acnes		18.38	0.10	SUC 87	95	
21411122	**Good identification**					**21411122**
Lactobacillus fermentum		100.0	0.35	NIT 1		
21411400	**Good identification**					**21411400**
Lactobacillus fermentum		100.0	0.40	NIT 1		
21411402	**Good identification**					**21411402**
Lactobacillus fermentum		100.0	0.40	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
21411500	**Good identification**					**21411500**
Lactobacillus fermentum		100.0	0.44	NIT 1		
21411502	**Good identification**					**21411502**
Lactobacillus fermentum		100.0	0.44	NIT 1		
21411520	**Good identification**					**21411520**
Lactobacillus fermentum		100.0	0.35	NIT 1		
21411522	**Good identification**					**21411522**
Lactobacillus fermentum		100.0	0.35	NIT 1		
21412020	**Acceptable identification**					**21412020**
					CAT	
Propionibacterium acnes		92.72	0.40	IND 87	95	
Pseudoramibacter alactolyticus		7.28	0.14	RAF 1	1	
21413020	**Good identification**					**21413020**
Propionibacterium acnes		100.0	0.40	IND 87		
21415000	**Acceptable identification**					**21415000**
					GRA	
Lactobacillus fermentum		93.54	0.36	NIT 1	99	
Fusobacterium mortiferum		6.46	0.15	SUC 87	1	
21415002	**Good identification**					**21415002**
Lactobacillus fermentum		100.0	0.36	NIT 1		
21415100	**Low discrimination**					**21415100**
					GRA	
Lactobacillus fermentum		83.51	0.40	NIT 1	99	
Fusobacterium mortiferum		8.24	0.21	SUC 87	1	
Actinomyces meyeri		8.24	0.12		99	
21415102	**Good identification**					**21415102**
Lactobacillus fermentum		100.0	0.40	NIT 1		
21415400	**Good identification**					**21415400**
Lactobacillus fermentum		100.0	0.36	NIT 1		
21415402	**Good identification**					**21415402**
Lactobacillus fermentum		100.0	0.36	NIT 1		
21415500	**Good identification**					**21415500**
Lactobacillus fermentum		100.0	0.40	NIT 1		
21415502	**Good identification**					**21415502**
Lactobacillus fermentum		100.0	0.40	NIT 1		
21501100	**Very good identification**					**21501100**
					COC	
Blautia hansenii		99.99	0.66	NIT 1	99	
Lactobacillus fermentum		0.01	0.18		1	
21501102	**Good identification**					**21501102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Blautia hansenii		99.26	0.50	NIT 1	99		
Actinomyces viscosus		0.63	0.17	BGL 10	1		
Lactobacillus fermentum		0.11	0.18		1		
21501120	**Good identification**						**21501120**
					COC	GRA	
Blautia hansenii		99.88	0.50	NIT 1	99	99	
Prevotella melaninogenica		0.12	-.01	NAG 10	1	1	
21501500	**Good identification**						**21501500**
					COC		
Blautia hansenii		99.93	0.53	NIT 1	99		
Lactobacillus fermentum		0.07	0.18		1		
21501502	**Good identification**						**21501502**
					COC		
Blautia hansenii		99.40	0.37	NIT 1	99		
Lactobacillus fermentum		0.61	0.18	BGL 10	1		
21501520	**Good identification**						**21501520**
Blautia hansenii		100.0	0.37	NIT 1			
21505100	**Good identification**						**21505100**
					COC		
Blautia hansenii		99.96	0.53	NIT 1	99		
Lactobacillus fermentum		0.04	0.14		1		
21505102	**Good identification**						**21505102**
					COC		
Blautia hansenii		99.64	0.37	NIT 1	99		
Lactobacillus fermentum		0.36	0.14	BGL 10	1		
21505120	**Good identification**						**21505120**
Blautia hansenii		100.0	0.37	NIT 1			
21505500	**Good identification**						**21505500**
					COC		
Blautia hansenii		99.78	0.40	NIT 1	99		
Lactobacillus fermentum		0.23	0.14		1		
21510000	**Low discrimination**						**21510000**
					COC		
Prevotella intermedia		65.62	0.42	NIT 1	1		
Prevotella disiens		15.33	0.23		1		
Megasphaera elsdenii		8.35	0.29		99		
21510002	**Low discrimination**						**21510002**
					COC	GRA	
Actinomyces viscosus		88.50	0.38	SUC 99	1	99	
Actinomyces odontolyticus		7.47	0.27	MNS 96	1	99	
Megasphaera elsdenii		1.71	0.12		99	1	
21510102	**Good identification**						**21510102**
Actinomyces viscosus		95.67	0.42	SUC 99			
Propionibacterium propionicum		2.38	0.29	MNS 96			
Actinomyces odontolyticus		1.43	0.18				
21510200	**Good identification**						**21510200**
					GRA		
Actinomyces odontolyticus		95.09	0.35	SUC 99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		4.91	0.08	RAF 3	1		
21511000	**Low discrimination**						**21511000**
					GRA		
Prevotella intermedia		65.93	0.42	NIT 1	1		
Lactobacillus fermentum		28.31	0.48		99		
Actinomyces odontolyticus		4.43	0.27		99		
21511020	**Low discrimination**						**21511020**
					CAT		
Lactobacillus fermentum		74.42	0.40	NIT 1	1		
Propionibacterium acnes		9.11	0.10	SUC 87	95		
Actinomyces odontolyticus		5.69	0.14		10		
21511022	**Low discrimination**						**21511022**
Lactobacillus fermentum		89.81	0.40	NIT 1			
Actinomyces odontolyticus		4.21	0.10	SUC 87			
Clostridium perfringens		3.14	0.03				
21511100	**Low discrimination**						**21511100**
					COC	GRA	
Lactobacillus fermentum		50.15	0.52	NIT 1	1	99	
Blautia hansenii		42.89	0.32	SUC 87	99	99	
Fusobacterium mortiferum		1.59	0.25		1	1	
21511102	**Low discrimination**						**21511102**
					COC		
Actinomyces viscosus		49.22	0.39	SUC 99	1		
Lactobacillus fermentum		43.21	0.52	MNS 96	1		
Blautia hansenii		4.11	0.15		99		
21511120	**Low discrimination**						**21511120**
					GRA		
Lactobacillus fermentum		50.99	0.44	NIT 1	99		
Clostridium perfringens		17.03	0.23	SUC 87	99		
Prevotella melaninogenica		15.08	0.15		1		
21511122	**Low discrimination**						**21511122**
					GRA	CAT	
Lactobacillus fermentum		67.38	0.44	NIT 1	99	1	
Capnocytophaga ochracea		11.97	0.15	SUC 87	1	90	
Clostridium perfringens		10.59	0.18		99	1	
21511400	**Good identification**						**21511400**
					GRA		
Lactobacillus fermentum		97.70	0.48	NIT 1	99		
Prevotella intermedia		2.30	0.08	SUC 87	1		
21511402	**Good identification**						**21511402**
Lactobacillus fermentum		98.96	0.48	NIT 1			
Actinomyces viscosus		1.05	0.00	SUC 87			
21511420	**Good identification**						**21511420**
Lactobacillus fermentum		100.0	0.40	NIT 1			
21511422	**Good identification**						**21511422**
Lactobacillus fermentum		100.0	0.40	NIT 1			
21511500	**Low discrimination**						**21511500**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		86.89	0.52	NIT 1	1		
Blautia hansenii		13.11	0.19	SUC 87	99		
21511502	**Good identification**						**21511502**
					COC		
Lactobacillus fermentum		97.25	0.52	NIT 1	1		
Blautia hansenii		1.63	0.03	SUC 87	99		
Actinomyces viscosus		1.12	0.04		1		
21511520	**Acceptable identification**						**21511520**
					COC		
Lactobacillus fermentum		93.84	0.44	NIT 1	1		
Blautia hansenii		4.72	0.03	SUC 87	99		
Bifidobacterium longum subsp. longum		1.44	0.05		1		
21511522	**Good identification**						**21511522**
					GRA	CAT	
Lactobacillus fermentum		95.54	0.44	NIT 1	99	1	
Capnocytophaga ochracea		3.00	0.02	SUC 87	1	90	
Bifidobacterium longum subsp. longum		1.47	0.05		99	1	
21512100	**Very good identification**						**21512100**
Propionibacterium propionicum		100.0	0.60	SUC 99			
21512101	**Good identification**						**21512101**
Propionibacterium propionicum		100.0	0.44	SUC 99			
21512102	**Very good identification**						**21512102**
Propionibacterium propionicum		99.59	0.63	SUC 99			
Actinomyces viscosus		0.41	0.08				
21512103	**Good identification**						**21512103**
Propionibacterium propionicum		98.78	0.47	SUC 99			
Actinomyces viscosus		1.22	0.00	URE 10			
21512140	**Very good identification**						**21512140**
Propionibacterium propionicum		100.0	0.60	SUC 99			
21512141	**Good identification**						**21512141**
Propionibacterium propionicum		100.0	0.44	SUC 99			
21512142	**Very good identification**						**21512142**
Propionibacterium propionicum		100.0	0.63	SUC 99			
21512143	**Good identification**						**21512143**
Propionibacterium propionicum		100.0	0.47	SUC 99			
21513100	**Very good identification**						**21513100**
Propionibacterium propionicum		98.83	0.60	SUC 99			
Actinomyces naeslundii		0.56	0.32				
Lactobacillus fermentum		0.33	0.18				
21513101	**Acceptable identification**						**21513101**
Propionibacterium propionicum		92.90	0.44	SUC 99			
Actinomyces naeslundii		7.10	0.35	URE 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
21513102	**Very good identification**					**21513102**
Propionibacterium propionicum		99.16	0.63	SUC 99		
Actinomyces naeslundii		0.37	0.32			
Actinomyces viscosus		0.25	0.04			
21513103	**Good identification**					**21513103**
Propionibacterium propionicum		95.15	0.47	SUC 99		
Actinomyces naeslundii		4.85	0.35	URE 10		
21513140	**Very good identification**					**21513140**
Propionibacterium propionicum		100.0	0.60	SUC 99		
21513141	**Good identification**					**21513141**
Propionibacterium propionicum		100.0	0.44	SUC 99		
21513142	**Very good identification**					**21513142**
Propionibacterium propionicum		100.0	0.63	SUC 99		
21513143	**Good identification**					**21513143**
Propionibacterium propionicum		100.0	0.47	SUC 99		
21515000	**Low discrimination**					**21515000**
					GRA	
Lactobacillus fermentum		84.29	0.44	NIT 1	99	
Actinomyces odontolyticus		11.06	0.22	SUC 87	99	
Prevotella intermedia		3.38	0.08		1	
21515002	**Acceptable identification**					**21515002**
Lactobacillus fermentum		91.04	0.44	NIT 1		
Actinomyces odontolyticus		7.32	0.18	SUC 87		
Actinomyces viscosus		1.64	0.00			
21515020	**Acceptable identification**					**21515020**
Lactobacillus fermentum		93.98	0.36	NIT 1		
Actinomyces odontolyticus		6.02	0.08	SUC 87		
21515022	**Good identification**					**21515022**
Lactobacillus fermentum		100.0	0.36	NIT 1		
21515100	**Low discrimination**					**21515100**
					COC	
Lactobacillus fermentum		75.79	0.48	NIT 1	1	
Blautia hansenii		19.47	0.19	SUC 87	99	
Actinomyces meyeri		2.81	0.13		1	
21515102	**Acceptable identification**					**21515102**
					COC	
Lactobacillus fermentum		94.01	0.48	NIT 1	1	
Blautia hansenii		2.68	0.03	SUC 87	99	
Actinomyces viscosus		1.84	0.04		1	
21515120	**Acceptable identification**					**21515120**
					COC	
Lactobacillus fermentum		92.11	0.40	NIT 1	1	
Blautia hansenii		7.89	0.03	SUC 87	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
21515122	**Acceptable identification**						**21515122**
					GRA	CAT	
Lactobacillus fermentum		92.97	0.40	NIT 1	99	1	
Capnocytophaga ochracea		7.03	0.05	SUC 87	1	90	
21515400	**Good identification**						**21515400**
Lactobacillus fermentum		100.0	0.44	NIT 1			
21515402	**Good identification**						**21515402**
Lactobacillus fermentum		100.0	0.44	NIT 1			
21515420	**Good identification**						**21515420**
Lactobacillus fermentum		100.0	0.36	NIT 1			
21515422	**Good identification**						**21515422**
Lactobacillus fermentum		100.0	0.36	NIT 1			
21515500	**Acceptable identification**						**21515500**
					COC		
Lactobacillus fermentum		94.82	0.48	NIT 1	1		
Blautia hansenii		4.30	0.06	SUC 87	99		
Actinomyces meyeri		0.88	0.02		1		
21515502	**Good identification**						**21515502**
Lactobacillus fermentum		100.0	0.48	NIT 1			
21515520	**Acceptable identification**						**21515520**
Lactobacillus fermentum		92.74	0.40	NIT 1			
Bifidobacterium longum subsp. longum		7.26	0.14	SUC 87			
21515522	**Acceptable identification**						**21515522**
Lactobacillus fermentum		92.74	0.40	NIT 1			
Bifidobacterium longum subsp. longum		7.26	0.14	SUC 87			
21532100	**Very good identification**						**21532100**
Propionibacterium propionicum		100.0	0.58	SUC 99			
21532101	**Good identification**						**21532101**
Propionibacterium propionicum		100.0	0.42	SUC 99			
21532102	**Very good identification**						**21532102**
Propionibacterium propionicum		100.0	0.61	SUC 99			
21532103	**Good identification**						**21532103**
Propionibacterium propionicum		100.0	0.45	SUC 99			
21532140	**Very good identification**						**21532140**
Propionibacterium propionicum		100.0	0.58	SUC 99			
21532141	**Good identification**						**21532141**
Propionibacterium propionicum		100.0	0.42	SUC 99			
21532142	**Very good identification**						**21532142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.61	SUC 99		
21532143	**Good identification**					**21532143**
Propionibacterium propionicum		100.0	0.45	SUC 99		
21533100	**Very good identification**					**21533100**
Propionibacterium propionicum		98.68	0.58	SUC 99		
Actinomyces naeslundii		1.33	0.37			
21533101	**Low discrimination**					**21533101**
Propionibacterium propionicum		84.65	0.42	SUC 99		
Actinomyces naeslundii		15.35	0.40	URE 10		
21533102	**Very good identification**					**21533102**
Propionibacterium propionicum		99.11	0.61	SUC 99		
Actinomyces naeslundii		0.89	0.37			
21533103	**Low discrimination**					**21533103**
Propionibacterium propionicum		89.22	0.45	SUC 99		
Actinomyces naeslundii		10.78	0.40	URE 10		
21533120	**Low discrimination**					**21533120**
Actinomyces naeslundii		57.07	0.37	SUC 99		
Propionibacterium propionicum		42.93	0.24	MNS 88		
21533121	**Good identification**					**21533121**
Actinomyces naeslundii		100.0	0.40	SUC 99		
21533123	**Good identification**					**21533123**
Actinomyces naeslundii		100.0	0.40	SUC 99		
21533140	**Very good identification**					**21533140**
Propionibacterium propionicum		99.76	0.58	SUC 99		
Actinomyces naeslundii		0.24	0.24			
21533141	**Good identification**					**21533141**
Propionibacterium propionicum		96.90	0.42	SUC 99		
Actinomyces naeslundii		3.10	0.27	URE 10		
21533142	**Very good identification**					**21533142**
Propionibacterium propionicum		99.84	0.61	SUC 99		
Actinomyces naeslundii		0.16	0.24			
21533143	**Good identification**					**21533143**
Propionibacterium propionicum		97.91	0.45	SUC 99		
Actinomyces naeslundii		2.09	0.27	URE 10		
21553101	**Good identification**					**21553101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
21553103	**Low discrimination**					**21553103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		86.99	0.37	SUC 99		
Propionibacterium propionicum		13.01	0.12	MNS 88		
21553121	**Good identification**					**21553121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
21553123	**Good identification**					**21553123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
21571101	**Good identification**					**21571101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
21571103	**Low discrimination**					**21571103**
Actinomyces naeslundii		81.86	0.37	SUC 99		
Actinomyces viscosus		18.14	0.01	MNS 88		
21571121	**Good identification**					**21571121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
21571123	**Good identification**					**21571123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
21573100	**Low discrimination**					**21573100**
Actinomyces naeslundii		58.28	0.39	SUC 99		
Propionibacterium propionicum		33.07	0.24	MNS 88		
Actinomyces israelii		8.65	0.15			
21573101	**Good identification**					**21573101**
Actinomyces naeslundii		100.0	0.42	SUC 99		
21573102	**Low discrimination**					**21573102**
Actinomyces naeslundii		48.22	0.39	SUC 99		
Propionibacterium propionicum		41.04	0.27	MNS 88		
Actinomyces israelii		10.74	0.18			
21573103	**Good identification**					**21573103**
Actinomyces naeslundii		100.0	0.42	SUC 99		
21573120	**Good identification**					**21573120**
Actinomyces naeslundii		100.0	0.39	SUC 99		
21573121	**Good identification**					**21573121**
Actinomyces naeslundii		100.0	0.42	SUC 99		
21573122	**Good identification**					**21573122**
Actinomyces naeslundii		100.0	0.39	SUC 99		
21573123	**Good identification**					**21573123**
Actinomyces naeslundii		100.0	0.42	SUC 99		
21577102	**Good identification**					**21577102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	SUC 98		
21610000	**Acceptable identification**					**21610000**
					GRA	
Fusobacterium mortiferum		92.45	0.36	NIT 1	1	
Pseudoramibacter alactolyticus		7.55	-.03		99	
21610100	**Good identification**					**21610100**
Fusobacterium mortiferum		100.0	0.42	NIT 1		
21611000	**Good identification**					**21611000**
Fusobacterium mortiferum		100.0	0.39	NIT 1		
21611100	**Good identification**					**21611100**
Fusobacterium mortiferum		100.0	0.45	NIT 1		
21631100	**Good identification**					**21631100**
Fusobacterium mortiferum		100.0	0.36	NIT 1		
21651100	**Good identification**					**21651100**
Fusobacterium mortiferum		100.0	0.37	NIT 1		
21710000	**Low discrimination**					**21710000**
					GRA	
Actinomyces odontolyticus		89.72	0.35	SUC 99	99	
Fusobacterium mortiferum		5.65	0.20	RAF 3	1	
Prevotella intermedia		4.63	0.08		1	
21710200	**Good identification**					**21710200**
Actinomyces odontolyticus		100.0	0.40	SUC 99		
21710202	**Good identification**					**21710202**
Actinomyces odontolyticus		100.0	0.36	SUC 99		
21711101	**Good identification**					**21711101**
Actinomyces naeslundii		100.0	0.39	SUC 99		
21711120	**Low discrimination**					**21711120**
Actinomyces naeslundii		45.70	0.36	SUC 99		
Clostridium perfringens		40.40	0.13	MNS 88		
Actinomyces odontolyticus		13.89	0.10			
21711121	**Good identification**					**21711121**
Actinomyces naeslundii		100.0	0.39	SUC 99		
21711122	**Low discrimination**					**21711122**
Actinomyces naeslundii		62.41	0.36	SUC 99		
Clostridium perfringens		25.96	0.07	MNS 88		
Actinomyces odontolyticus		11.63	0.06			
21711123	**Good identification**					**21711123**
Actinomyces naeslundii		100.0	0.39	SUC 99		
21711200	**Good identification**					**21711200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.37	SUC 99		
21713001	**Good identification**					**21713001**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21713003	**Good identification**					**21713003**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21713021	**Good identification**					**21713021**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21713023	**Good identification**					**21713023**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21713100	**Low discrimination**					**21713100**
Actinomyces naeslundii		66.52	0.41	SUC 99		
Propionibacterium propionicum		33.48	0.26	MNS 88		
21713101	**Good identification**					**21713101**
Actinomyces naeslundii		100.0	0.44	SUC 99		
21713102	**Low discrimination**					**21713102**
Actinomyces naeslundii		56.98	0.41	SUC 99		
Propionibacterium propionicum		43.02	0.29	MNS 88		
21713103	**Acceptable identification**					**21713103**
Actinomyces naeslundii		94.70	0.44	SUC 99		
Propionibacterium propionicum		5.30	0.12	MNS 88		
21713120	**Good identification**					**21713120**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21713121	**Good identification**					**21713121**
Actinomyces naeslundii		100.0	0.44	SUC 99		
21713122	**Good identification**					**21713122**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21713123	**Good identification**					**21713123**
Actinomyces naeslundii		100.0	0.44	SUC 99		
21731001	**Good identification**					**21731001**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21731003	**Good identification**					**21731003**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21731021	**Good identification**					**21731021**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21731023	**Good identification**					**21731023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	SUC 99		
21731100	**Low discrimination**					**21731100**
					GRA	
Actinomyces naeslundii		76.86	0.41	SUC 99	99	
Fusobacterium mortiferum		23.14	0.21	MNS 88	1	
21731101	**Good identification**					**21731101**
Actinomyces naeslundii		100.0	0.44	SUC 99		
21731102	**Low discrimination**					**21731102**
Actinomyces naeslundii		72.84	0.41	SUC 99		
Actinomyces viscosus		27.16	0.10	MNS 88		
21731103	**Acceptable identification**					**21731103**
Actinomyces naeslundii		92.35	0.44	SUC 99		
Actinomyces viscosus		7.65	0.01	MNS 88		
21731120	**Low discrimination**					**21731120**
Actinomyces naeslundii		54.18	0.41	SUC 99		
Clostridium perfringens		45.82	0.17	MNS 88		
21731121	**Good identification**					**21731121**
Actinomyces naeslundii		100.0	0.44	SUC 99		
21731122	**Low discrimination**					**21731122**
Actinomyces naeslundii		71.53	0.41	SUC 99		
Clostridium perfringens		28.47	0.12	MNS 88		
21731123	**Good identification**					**21731123**
Actinomyces naeslundii		100.0	0.44	SUC 99		
21733000	**Good identification**					**21733000**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21733001	**Good identification**					**21733001**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21733002	**Good identification**					**21733002**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21733003	**Good identification**					**21733003**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21733020	**Good identification**					**21733020**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21733021	**Good identification**					**21733021**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21733022	**Good identification**					**21733022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	SUC 99		
21733023	**Good identification**					**21733023**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21733100	**Low discrimination**					**21733100**
Actinomyces naeslundii		79.35	0.46	SUC 99		
Propionibacterium propionicum		16.84	0.24	MNS 88		
Actinomyces israelii		3.81	0.14			
21733101	**Good identification**					**21733101**
Actinomyces naeslundii		100.0	0.49	SUC 99		
21733102	**Low discrimination**					**21733102**
Actinomyces naeslundii		71.93	0.46	SUC 99		
Propionibacterium propionicum		22.89	0.27	MNS 88		
Actinomyces israelii		5.18	0.17			
21733103	**Good identification**					**21733103**
Actinomyces naeslundii		100.0	0.49	SUC 99		
21733120	**Good identification**					**21733120**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21733121	**Good identification**					**21733121**
Actinomyces naeslundii		100.0	0.49	SUC 99		
21733122	**Good identification**					**21733122**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21733123	**Good identification**					**21733123**
Actinomyces naeslundii		100.0	0.49	SUC 99		
21733141	**Good identification**					**21733141**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21733143	**Good identification**					**21733143**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21733161	**Good identification**					**21733161**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21733163	**Good identification**					**21733163**
Actinomyces naeslundii		100.0	0.36	SUC 99		
21737102	**Good identification**					**21737102**
Actinomyces israelii		100.0	0.35	SUC 98		
21751100	**Low discrimination**					**21751100**
					GRA	
Actinomyces naeslundii		67.03	0.38	SUC 99	99	
Fusobacterium mortiferum		32.97	0.21	MNS 88	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
21751101	**Good identification**					**21751101**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21751103	**Low discrimination**					**21751103**
Actinomyces naeslundii		70.01	0.41	SUC 99		
Actinomyces viscosus		29.99	0.11	MNS 88		
21751120	**Good identification**					**21751120**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21751121	**Good identification**					**21751121**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21751122	**Low discrimination**					**21751122**
					GRA	
Actinomyces naeslundii		83.09	0.38	SUC 99	99	
Prevotella oralis		16.92	0.01	MNS 88	1	
21751123	**Good identification**					**21751123**
Actinomyces naeslundii		100.0	0.41	SUC 99		
21753000	**Low discrimination**					**21753000**
Actinomyces naeslundii		84.97	0.35	SUC 99		
Actinomyces israelii		15.03	0.13	MNS 88		
21753001	**Good identification**					**21753001**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21753002	**Low discrimination**					**21753002**
Actinomyces naeslundii		79.04	0.35	SUC 99		
Actinomyces israelii		20.96	0.16	MNS 88		
21753003	**Good identification**					**21753003**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21753020	**Good identification**					**21753020**
Actinomyces naeslundii		100.0	0.35	SUC 99		
21753021	**Good identification**					**21753021**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21753022	**Good identification**					**21753022**
Actinomyces naeslundii		100.0	0.35	SUC 99		
21753023	**Good identification**					**21753023**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21753100	**Low discrimination**					**21753100**
Actinomyces naeslundii		80.10	0.43	SUC 99		
Actinomyces israelii		19.90	0.24	MNS 88		
21753101	**Good identification**					**21753101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	SUC 99		
21753102	**Low discrimination**					**21753102**
Actinomyces naeslundii		72.85	0.43	SUC 99		
Actinomyces israelii		27.15	0.27	MNS 88		
21753103	**Good identification**					**21753103**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21753120	**Good identification**					**21753120**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21753121	**Good identification**					**21753121**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21753122	**Good identification**					**21753122**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21753123	**Good identification**					**21753123**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21757100	**Good identification**					**21757100**
Actinomyces israelii		100.0	0.42	SUC 98		
21757102	**Good identification**					**21757102**
Actinomyces israelii		100.0	0.45	SUC 98		
21757142	**Good identification**					**21757142**
Actinomyces israelii		100.0	0.38	SUC 98		
21757500	**Good identification**					**21757500**
Actinomyces israelii		100.0	0.40	SUC 98		
21757502	**Good identification**					**21757502**
Actinomyces israelii		100.0	0.43	SUC 98		
21757542	**Good identification**					**21757542**
Actinomyces israelii		100.0	0.36	SUC 98		
21771000	**Good identification**					**21771000**
Actinomyces naeslundii		100.0	0.35	SUC 99		
21771001	**Good identification**					**21771001**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21771002	**Low discrimination**					**21771002**
Actinomyces naeslundii		85.03	0.35	SUC 99		
Actinomyces israelii		14.98	0.13	MNS 88		
21771003	**Good identification**					**21771003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	SUC 99		
21771020	**Good identification**					**21771020**
Actinomyces naeslundii		100.0	0.35	SUC 99		
21771021	**Good identification**					**21771021**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21771022	**Good identification**					**21771022**
Actinomyces naeslundii		100.0	0.35	SUC 99		
21771023	**Good identification**					**21771023**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21771100	**Low discrimination**					**21771100**
					GRA	
Actinomyces naeslundii		80.60	0.43	SUC 99	99	
Actinomyces israelii		13.30	0.21	MNS 88	99	
Fusobacterium mortiferum		6.10	0.12		1	
21771101	**Good identification**					**21771101**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21771102	**Low discrimination**					**21771102**
Actinomyces naeslundii		75.63	0.43	SUC 99		
Actinomyces israelii		18.71	0.24	MNS 88		
Actinomyces viscosus		5.65	0.00			
21771103	**Good identification**					**21771103**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21771120	**Good identification**					**21771120**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21771121	**Good identification**					**21771121**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21771122	**Good identification**					**21771122**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21771123	**Good identification**					**21771123**
Actinomyces naeslundii		100.0	0.46	SUC 99		
21773000	**Low discrimination**					**21773000**
Actinomyces naeslundii		74.48	0.40	SUC 99		
Actinomyces israelii		25.52	0.23	MNS 88		
21773001	**Good identification**					**21773001**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21773002	**Low discrimination**					**21773002**
Actinomyces naeslundii		66.05	0.40	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		33.95	0.26	MNS 88		
21773003	**Good identification**					**21773003**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21773020	**Good identification**					**21773020**
Actinomyces naeslundii		100.0	0.40	SUC 99		
21773021	**Good identification**					**21773021**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21773022	**Good identification**					**21773022**
Actinomyces naeslundii		100.0	0.40	SUC 99		
21773023	**Good identification**					**21773023**
Actinomyces naeslundii		100.0	0.43	SUC 99		
21773100	**Low discrimination**					**21773100**
Actinomyces naeslundii		67.50	0.48	SUC 99		
Actinomyces israelii		32.51	0.33	MNS 88		
21773101	**Good identification**					**21773101**
Actinomyces naeslundii		100.0	0.51	SUC 99		
21773102	**Low discrimination**					**21773102**
Actinomyces naeslundii		58.06	0.48	SUC 99		
Actinomyces israelii		41.94	0.36	MNS 88		
21773103	**Good identification**					**21773103**
Actinomyces naeslundii		100.0	0.51	SUC 99		
21773120	**Good identification**					**21773120**
Actinomyces naeslundii		100.0	0.48	SUC 99		
21773121	**Good identification**					**21773121**
Actinomyces naeslundii		100.0	0.51	SUC 99		
21773122	**Good identification**					**21773122**
Actinomyces naeslundii		95.59	0.48	SUC 99		
Actinomyces israelii		4.41	0.16	MNS 88		
21773123	**Good identification**					**21773123**
Actinomyces naeslundii		100.0	0.51	SUC 99		
21773141	**Good identification**					**21773141**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21773143	**Good identification**					**21773143**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21773160	**Good identification**					**21773160**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	SUC 99		
21773161	**Good identification**					**21773161**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21773162	**Good identification**					**21773162**
Actinomyces naeslundii		100.0	0.35	SUC 99		
21773163	**Good identification**					**21773163**
Actinomyces naeslundii		100.0	0.38	SUC 99		
21775100	**Good identification**					**21775100**
Actinomyces israelii		100.0	0.39	SUC 98		
21775102	**Good identification**					**21775102**
Actinomyces israelii		100.0	0.42	SUC 98		
21775500	**Good identification**					**21775500**
Actinomyces israelii		100.0	0.37	SUC 98		
21775502	**Good identification**					**21775502**
Actinomyces israelii		100.0	0.40	SUC 98		
21777000	**Good identification**					**21777000**
Actinomyces israelii		100.0	0.41	SUC 98		
21777002	**Good identification**					**21777002**
Actinomyces israelii		100.0	0.44	SUC 98		
21777042	**Good identification**					**21777042**
Actinomyces israelii		100.0	0.37	SUC 98		
21777100	**Good identification**					**21777100**
Actinomyces israelii		100.0	0.52	SUC 98		
21777102	**Good identification**					**21777102**
Actinomyces israelii		100.0	0.55	SUC 98		
21777110	**Good identification**					**21777110**
Actinomyces israelii		100.0	0.37	SUC 98		
21777112	**Good identification**					**21777112**
Actinomyces israelii		100.0	0.40	SUC 98		
21777140	**Good identification**					**21777140**
Actinomyces israelii		100.0	0.45	SUC 98		
21777142	**Good identification**					**21777142**
Actinomyces israelii		100.0	0.48	SUC 98		
21777302	**Good identification**					**21777302**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.35	SUC 98			
21777400	**Good identification**						**21777400**
Actinomyces israelii		100.0	0.40	SUC 98			
21777402	**Good identification**						**21777402**
Actinomyces israelii		100.0	0.43	SUC 98			
21777442	**Good identification**						**21777442**
Actinomyces israelii		100.0	0.36	SUC 98			
21777500	**Good identification**						**21777500**
Actinomyces israelii		100.0	0.50	SUC 98			
21777502	**Good identification**						**21777502**
Actinomyces israelii		100.0	0.53	SUC 98			
21777510	**Good identification**						**21777510**
Actinomyces israelii		100.0	0.36	SUC 98			
21777512	**Good identification**						**21777512**
Actinomyces israelii		100.0	0.39	SUC 98			
21777540	**Good identification**						**21777540**
Actinomyces israelii		100.0	0.43	SUC 98			
21777542	**Good identification**						**21777542**
Actinomyces israelii		100.0	0.46	SUC 98			
22000000	**Low discrimination**						**22000000**
					COC	GRA	
Veillonella parvula		51.34	0.66	SUC 1	99	1	
Campylobacter gracilis		37.71	0.66		1	1	
Eggerthella lenta		6.22	0.51		1	99	
22000001	**Good identification**						**22000001**
					COC		
Campylobacter ureolyticus		93.37	0.60	SUC 1	1		
Veillonella parvula		3.14	0.32		99		
Campylobacter gracilis		2.31	0.32		1		
22000002	**Low discrimination**						**22000002**
					COC	GRA	
Campylobacter gracilis		83.16	0.50	SUC 1	1	1	
Veillonella parvula		10.29	0.32	BGL 10	99	1	
Atopobium minutum		1.31	0.37		1	99	
22000020	**Acceptable identification**						**22000020**
					COC	GRA	
Campylobacter gracilis		93.09	0.56	SUC 1	1	1	
Veillonella parvula		5.12	0.32		99	1	
Eggerthella lenta		0.62	0.17		1	99	
22000022	**Low discrimination**						**22000022**
					GRA		
Campylobacter gracilis		80.69	0.39	SUC 1	1		
Pseudoflavonifractor capillosus		16.67	0.32	BGL 10	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		1.69	0.29		99		
22010002	**Low discrimination**						**22010002**
					GRA		
Atopobium minutum		51.28	0.36	NIT 1	99		
Campylobacter gracilis		35.77	0.15	SUC 5	1		
Actinomyces odontolyticus		4.54	0.18		99		
22010100	**Acceptable identification**						**22010100**
					COC	GRA	
Actinomyces meyeri		90.55	0.38	NIT 1	1	99	
Veillonella parvula		3.87	-.03	GLU 93	99	1	
Campylobacter gracilis		2.84	-.03		1	1	
22011100	**Good identification**						**22011100**
Actinomyces meyeri		96.92	0.41	NIT 1			
Lactobacillus fermentum		1.36	0.20	GLU 93			
Actinomyces odontolyticus		0.97	0.10				
22014100	**Good identification**						**22014100**
Actinomyces meyeri		99.43	0.43	NIT 1			
Actinomyces odontolyticus		0.57	0.08	GLU 93			
22015000	**Good identification**						**22015000**
Actinomyces meyeri		95.91	0.35	NIT 1			
Actinomyces odontolyticus		3.10	0.14	GLU 93			
Lactobacillus fermentum		1.00	0.12				
22015100	**Good identification**						**22015100**
Actinomyces meyeri		99.58	0.46	NIT 1			
Lactobacillus fermentum		0.42	0.16	GLU 93			
22015500	**Good identification**						**22015500**
Actinomyces meyeri		98.33	0.35	NIT 1			
Lactobacillus fermentum		1.67	0.16	GLU 93			
22100200	**Acceptable identification**						**22100200**
					COC	GRA	
Actinomyces odontolyticus		93.60	0.43	GLU 99	1	99	
Veillonella parvula		3.35	-.03	FRU 92	99	1	
Campylobacter gracilis		2.46	-.03		1	1	
22100202	**Good identification**						**22100202**
Actinomyces odontolyticus		100.0	0.39	GLU 99			
22101101	**Good identification**						**22101101**
Clostridium chauvoei		100.0	0.40	GLU 99			
22101200	**Good identification**						**22101200**
Actinomyces odontolyticus		100.0	0.40	GLU 99			
22101202	**Good identification**						**22101202**
Actinomyces odontolyticus		100.0	0.36	GLU 99			
22104200	**Good identification**						**22104200**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.38	GLU 99			
22110000	**Low discrimination**						**22110000**
					COC	GRA	
Actinomyces odontolyticus		89.00	0.56	GLU 99	1	99	
Megasphaera elsdenii		5.21	0.32		99	1	
Actinomyces meyeri		3.54	0.28		1	99	
22110002	**Good identification**						**22110002**
					COC	GRA	
Actinomyces odontolyticus		96.52	0.52	GLU 99	1	99	
Actinomyces viscosus		2.26	0.18		1	99	
Megasphaera elsdenii		1.03	0.15		99	1	
22110020	**Good identification**						**22110020**
					GRA		
Actinomyces odontolyticus		96.14	0.43	GLU 99	99		
Actinomyces meyeri		3.21	0.14	NAG 14	99		
Campylobacter gracilis		0.65	-.13		1		
22110022	**Good identification**						**22110022**
Actinomyces odontolyticus		100.0	0.39	GLU 99			
22110100	**Low discrimination**						**22110100**
Actinomyces odontolyticus		65.12	0.47	GLU 99			
Actinomyces meyeri		32.84	0.39				
Propionibacterium propionicum		2.05	0.29				
22110102	**Low discrimination**						**22110102**
Actinomyces odontolyticus		81.03	0.44	GLU 99			
Actinomyces viscosus		10.67	0.22				
Propionibacterium propionicum		6.23	0.32				
22110200	**Very good identification**						**22110200**
Actinomyces odontolyticus		100.0	0.61	GLU 99			
22110202	**Very good identification**						**22110202**
Actinomyces odontolyticus		100.0	0.57	GLU 99			
22110220	**Good identification**						**22110220**
Actinomyces odontolyticus		100.0	0.47	GLU 99			
22110222	**Good identification**						**22110222**
Actinomyces odontolyticus		100.0	0.44	GLU 99			
22110300	**Good identification**						**22110300**
Actinomyces odontolyticus		99.74	0.52	GLU 99			
Actinomyces meyeri		0.26	0.05				
22110302	**Good identification**						**22110302**
Actinomyces odontolyticus		100.0	0.49	GLU 99			
22110320	**Good identification**						**22110320**
Actinomyces odontolyticus		100.0	0.39	GLU 99			
22110322	**Good identification**						**22110322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.35	GLU 99		
22111000	**Low discrimination**					**22111000**
					GRA	
Actinomyces odontolyticus		89.72	0.53	GLU 99	99	
Actinomyces meyeri		8.04	0.31		99	
Prevotella intermedia		1.51	0.17		1	
22111002	**Good identification**					**22111002**
Actinomyces odontolyticus		96.21	0.49	GLU 99		
Actinomyces viscosus		2.07	0.14			
Lactobacillus fermentum		1.29	0.24			
22111020	**Acceptable identification**					**22111020**
Actinomyces odontolyticus		90.26	0.40	GLU 99		
Actinomyces meyeri		6.77	0.17	NAG 14		
Lactobacillus fermentum		1.51	0.16			
22111022	**Good identification**					**22111022**
Actinomyces odontolyticus		97.34	0.36	GLU 99		
Lactobacillus fermentum		2.66	0.16	NAG 14		
22111100	**Low discrimination**					**22111100**
Actinomyces meyeri		50.35	0.42	NIT 1		
Actinomyces odontolyticus		44.38	0.44	GLU 93		
Propionibacterium propionicum		2.09	0.29			
22111101	**Good identification**					**22111101**
					SPO	
Clostridium chauvoei		99.09	0.47	GLU 99	99	
Actinomyces meyeri		0.39	0.08	MNS 90	1	
Actinomyces odontolyticus		0.35	0.10		1	
22111102	**Low discrimination**					**22111102**
Actinomyces odontolyticus		72.93	0.41	GLU 99		
Actinomyces viscosus		8.83	0.18			
Propionibacterium propionicum		8.41	0.32			
22111121	**Good identification**					**22111121**
Clostridium chauvoei		100.0	0.38	GLU 99		
22111200	**Very good identification**					**22111200**
Actinomyces odontolyticus		100.0	0.58	GLU 99		
22111202	**Good identification**					**22111202**
Actinomyces odontolyticus		100.0	0.54	GLU 99		
22111220	**Good identification**					**22111220**
Actinomyces odontolyticus		100.0	0.44	GLU 99		
22111222	**Good identification**					**22111222**
Actinomyces odontolyticus		100.0	0.41	GLU 99		
22111300	**Good identification**					**22111300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.41	0.49	GLU 99		
Actinomyces meyeri		0.59	0.08			
22111302	**Good identification**					**22111302**
Actinomyces odontolyticus		100.0	0.46	GLU 99		
22111320	**Good identification**					**22111320**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22112100	**Very good identification**					**22112100**
Propionibacterium propionicum		99.51	0.63	GLU 99		
Actinomyces odontolyticus		0.32	0.13			
Actinomyces meyeri		0.16	0.05			
22112101	**Good identification**					**22112101**
Propionibacterium propionicum		100.0	0.47	GLU 99		
22112102	**Very good identification**					**22112102**
Propionibacterium propionicum		99.87	0.66	GLU 99		
Actinomyces odontolyticus		0.13	0.10			
22112103	**Good identification**					**22112103**
Propionibacterium propionicum		100.0	0.50	GLU 99		
22112140	**Very good identification**					**22112140**
Propionibacterium propionicum		100.0	0.63	GLU 99		
22112141	**Good identification**					**22112141**
Propionibacterium propionicum		100.0	0.47	GLU 99		
22112142	**Very good identification**					**22112142**
Propionibacterium propionicum		100.0	0.66	GLU 99		
22112143	**Good identification**					**22112143**
Propionibacterium propionicum		100.0	0.50	GLU 99		
22113100	**Very good identification**					**22113100**
Propionibacterium propionicum		99.54	0.63	GLU 99		
Actinomyces meyeri		0.25	0.08			
Actinomyces odontolyticus		0.22	0.10			
22113101	**Acceptable identification**					**22113101**
					SPO	
Propionibacterium propionicum		94.64	0.47	GLU 99	1	
Clostridium chauvoei		5.36	0.12	URE 10	99	
22113102	**Very good identification**					**22113102**
Propionibacterium propionicum		99.91	0.66	GLU 99		
Actinomyces odontolyticus		0.09	0.07			
22113103	**Good identification**					**22113103**
					SPO	
Propionibacterium propionicum		99.34	0.50	GLU 99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Clostridium chauvoei	0.66	-.01	URE 10	99	
22113140	**Very good identification**					**22113140**
	Propionibacterium propionicum	100.0	0.63	GLU 99		
22113141	**Good identification**					**22113141**
	Propionibacterium propionicum	100.0	0.47	GLU 99		
22113142	**Very good identification**					**22113142**
	Propionibacterium propionicum	100.0	0.66	GLU 99		
22113143	**Good identification**					**22113143**
	Propionibacterium propionicum	100.0	0.50	GLU 99		
22114000	**Low discrimination**					**22114000**
	Actinomyces odontolyticus	86.44	0.51	GLU 99		
	Actinomyces meyeri	13.56	0.33			
22114002	**Good identification**					**22114002**
	Actinomyces odontolyticus	99.21	0.47	GLU 99		
	Actinomyces meyeri	0.79	0.07			
22114020	**Low discrimination**					**22114020**
	Actinomyces odontolyticus	88.38	0.37	GLU 99		
	Actinomyces meyeri	11.62	0.19	NAG 14		
22114100	**Low discrimination**					**22114100**
	Actinomyces meyeri	66.53	0.44	NIT 1		
	Actinomyces odontolyticus	33.47	0.42	GLU 93		
22114102	**Acceptable identification**					**22114102**
	Actinomyces odontolyticus	90.89	0.39	GLU 99		
	Actinomyces meyeri	9.12	0.18			
22114200	**Good identification**					**22114200**
	Actinomyces odontolyticus	100.0	0.56	GLU 99		
22114202	**Good identification**					**22114202**
	Actinomyces odontolyticus	100.0	0.52	GLU 99		
22114220	**Good identification**					**22114220**
	Actinomyces odontolyticus	100.0	0.42	GLU 99		
22114222	**Good identification**					**22114222**
	Actinomyces odontolyticus	100.0	0.39	GLU 99		
22114300	**Good identification**					**22114300**
	Actinomyces odontolyticus	98.98	0.47	GLU 99		
	Actinomyces meyeri	1.02	0.09			
22114302	**Good identification**					**22114302**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.43	GLU 99		
22115000	**Low discrimination**					**22115000**
Actinomyces odontolyticus		73.38	0.48	GLU 99		
Actinomyces meyeri		25.91	0.36			
Lactobacillus fermentum		0.72	0.20			
22115002	**Good identification**					**22115002**
Actinomyces odontolyticus		96.74	0.44	GLU 99		
Actinomyces meyeri		1.72	0.11			
Lactobacillus fermentum		1.54	0.20			
22115100	**Low discrimination**					**22115100**
Actinomyces meyeri		80.98	0.47	NIT 1		
Actinomyces odontolyticus		18.11	0.39	GLU 93		
Lactobacillus fermentum		0.91	0.24			
22115102	**Low discrimination**					**22115102**
Actinomyces odontolyticus		76.46	0.36	GLU 99		
Actinomyces meyeri		17.25	0.21			
Lactobacillus fermentum		6.29	0.24			
22115200	**Good identification**					**22115200**
Actinomyces odontolyticus		99.82	0.53	GLU 99		
Actinomyces meyeri		0.18	0.02			
22115202	**Good identification**					**22115202**
Actinomyces odontolyticus		100.0	0.49	GLU 99		
22115220	**Good identification**					**22115220**
Actinomyces odontolyticus		100.0	0.39	GLU 99		
22115222	**Good identification**					**22115222**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22115300	**Good identification**					**22115300**
Actinomyces odontolyticus		97.73	0.44	GLU 99		
Actinomyces meyeri		2.27	0.13			
22115302	**Good identification**					**22115302**
Actinomyces odontolyticus		100.0	0.40	GLU 99		
22115500	**Good identification**					**22115500**
Actinomyces meyeri		95.68	0.36	NIT 1		
Lactobacillus fermentum		4.32	0.24	GLU 93		
22132100	**Very good identification**					**22132100**
Propionibacterium propionicum		100.0	0.61	GLU 99		
22132101	**Good identification**					**22132101**
Propionibacterium propionicum		100.0	0.45	GLU 99		
22132102	**Very good identification**					**22132102**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		100.0	0.64	GLU 99			
22132103	**Good identification**						**22132103**
Propionibacterium propionicum		100.0	0.48	GLU 99			
22132140	**Very good identification**						**22132140**
Propionibacterium propionicum		100.0	0.61	GLU 99			
22132141	**Good identification**						**22132141**
Propionibacterium propionicum		100.0	0.45	GLU 99			
22132142	**Very good identification**						**22132142**
Propionibacterium propionicum		100.0	0.64	GLU 99			
22132143	**Good identification**						**22132143**
Propionibacterium propionicum		100.0	0.48	GLU 99			
22133100	**Very good identification**						**22133100**
Propionibacterium propionicum		100.0	0.61	GLU 99			
22133101	**Good identification**						**22133101**
Propionibacterium propionicum		100.0	0.45	GLU 99			
22133102	**Very good identification**						**22133102**
Propionibacterium propionicum		100.0	0.64	GLU 99			
22133103	**Good identification**						**22133103**
Propionibacterium propionicum		100.0	0.48	GLU 99			
22133140	**Very good identification**						**22133140**
Propionibacterium propionicum		100.0	0.61	GLU 99			
22133141	**Good identification**						**22133141**
Propionibacterium propionicum		100.0	0.45	GLU 99			
22133142	**Very good identification**						**22133142**
Propionibacterium propionicum		100.0	0.64	GLU 99			
22133143	**Good identification**						**22133143**
Propionibacterium propionicum		100.0	0.48	GLU 99			
22300000	**Acceptable identification**						**22300000**
					COC	GRA	
Actinomyces odontolyticus		93.60	0.43	GLU 99	1	99	
Veillonella parvula		3.35	-.03	FRU 92	99	1	
Campylobacter gracilis		2.46	-.03		1	1	
22300002	**Good identification**						**22300002**
Actinomyces odontolyticus		100.0	0.39	GLU 99			
22300200	**Good identification**						**22300200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.48	GLU 99		
22300202	**Good identification**					**22300202**
Actinomyces odontolyticus		100.0	0.44	GLU 99		
22300300	**Good identification**					**22300300**
Actinomyces odontolyticus		100.0	0.39	GLU 99		
22300302	**Good identification**					**22300302**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22301000	**Good identification**					**22301000**
Actinomyces odontolyticus		98.78	0.40	GLU 99		
Eubacterium tortuosum		1.22	-.04	FRU 92		
22301002	**Good identification**					**22301002**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22301200	**Good identification**					**22301200**
Actinomyces odontolyticus		100.0	0.45	GLU 99		
22301202	**Good identification**					**22301202**
Actinomyces odontolyticus		100.0	0.41	GLU 99		
22301300	**Good identification**					**22301300**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22304000	**Good identification**					**22304000**
Actinomyces odontolyticus		100.0	0.38	GLU 99		
22304200	**Good identification**					**22304200**
Actinomyces odontolyticus		100.0	0.42	GLU 99		
22304202	**Good identification**					**22304202**
Actinomyces odontolyticus		100.0	0.39	GLU 99		
22305200	**Good identification**					**22305200**
Actinomyces odontolyticus		100.0	0.39	GLU 99		
22305202	**Good identification**					**22305202**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22310000	**Very good identification**					**22310000**
Actinomyces odontolyticus		100.0	0.61	GLU 99		
22310002	**Very good identification**					**22310002**
Actinomyces odontolyticus		99.68	0.57	GLU 99		
Actinomyces viscosus		0.32	0.08			
22310020	**Good identification**					**22310020**
Actinomyces odontolyticus		100.0	0.47	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22310022	**Good identification**					**22310022**
Actinomyces odontolyticus		100.0	0.44	GLU 99		
22310100	**Good identification**					**22310100**
Actinomyces odontolyticus		99.74	0.52	GLU 99		
Actinomyces meyeri		0.26	0.05			
22310102	**Good identification**					**22310102**
Actinomyces odontolyticus		98.23	0.49	GLU 99		
Actinomyces viscosus		1.77	0.12			
22310120	**Good identification**					**22310120**
Actinomyces odontolyticus		100.0	0.39	GLU 99		
22310122	**Good identification**					**22310122**
Actinomyces odontolyticus		100.0	0.35	GLU 99		
22310200	**Very good identification**					**22310200**
Actinomyces odontolyticus		100.0	0.66	GLU 99		
22310202	**Very good identification**					**22310202**
Actinomyces odontolyticus		100.0	0.62	GLU 99		
22310220	**Good identification**					**22310220**
Actinomyces odontolyticus		100.0	0.52	GLU 99		
22310222	**Good identification**					**22310222**
Actinomyces odontolyticus		100.0	0.49	GLU 99		
22310300	**Very good identification**					**22310300**
Actinomyces odontolyticus		100.0	0.57	GLU 99		
22310302	**Good identification**					**22310302**
Actinomyces odontolyticus		100.0	0.54	GLU 99		
22310320	**Good identification**					**22310320**
Actinomyces odontolyticus		100.0	0.44	GLU 99		
22310322	**Good identification**					**22310322**
Actinomyces odontolyticus		100.0	0.40	GLU 99		
22311000	**Very good identification**					**22311000**
Actinomyces odontolyticus		100.0	0.58	GLU 99		
22311002	**Good identification**					**22311002**
Actinomyces odontolyticus		99.71	0.54	GLU 99		
Actinomyces viscosus		0.29	0.04			
22311020	**Good identification**					**22311020**
Actinomyces odontolyticus		100.0	0.44	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22311022	**Good identification**					**22311022**
Actinomyces odontolyticus		100.0	0.41	GLU 99		
22311100	**Good identification**					**22311100**
Actinomyces odontolyticus		99.41	0.49	GLU 99		
Actinomyces meyeri		0.59	0.08			
22311102	**Good identification**					**22311102**
Actinomyces odontolyticus		98.37	0.46	GLU 99		
Actinomyces viscosus		1.63	0.08			
22311120	**Good identification**					**22311120**
Actinomyces odontolyticus		95.40	0.36	GLU 99		
Clostridium perfringens		4.60	0.08	NAG 14		
22311200	**Very good identification**					**22311200**
Actinomyces odontolyticus		100.0	0.63	GLU 99		
22311202	**Very good identification**					**22311202**
Actinomyces odontolyticus		100.0	0.59	GLU 99		
22311220	**Good identification**					**22311220**
Actinomyces odontolyticus		100.0	0.49	GLU 99		
22311222	**Good identification**					**22311222**
Actinomyces odontolyticus		100.0	0.46	GLU 99		
22311300	**Good identification**					**22311300**
Actinomyces odontolyticus		100.0	0.54	GLU 99		
22311302	**Good identification**					**22311302**
Actinomyces odontolyticus		100.0	0.51	GLU 99		
22311320	**Good identification**					**22311320**
Actinomyces odontolyticus		100.0	0.41	GLU 99		
22311322	**Good identification**					**22311322**
Actinomyces odontolyticus		100.0	0.37	GLU 99		
22314000	**Good identification**					**22314000**
Actinomyces odontolyticus		100.0	0.56	GLU 99		
22314002	**Good identification**					**22314002**
Actinomyces odontolyticus		100.0	0.52	GLU 99		
22314020	**Good identification**					**22314020**
Actinomyces odontolyticus		100.0	0.42	GLU 99		
22314022	**Good identification**					**22314022**
Actinomyces odontolyticus		100.0	0.39	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22314100	**Good identification**					**22314100**
Actinomyces odontolyticus		98.98	0.47	GLU 99		
Actinomyces meyeri		1.02	0.09			
22314102	**Good identification**					**22314102**
Actinomyces odontolyticus		100.0	0.43	GLU 99		
22314200	**Very good identification**					**22314200**
Actinomyces odontolyticus		100.0	0.61	GLU 99		
22314202	**Very good identification**					**22314202**
Actinomyces odontolyticus		100.0	0.57	GLU 99		
22314220	**Good identification**					**22314220**
Actinomyces odontolyticus		100.0	0.47	GLU 99		
22314222	**Good identification**					**22314222**
Actinomyces odontolyticus		100.0	0.43	GLU 99		
22314300	**Good identification**					**22314300**
Actinomyces odontolyticus		100.0	0.52	GLU 99		
22314302	**Good identification**					**22314302**
Actinomyces odontolyticus		100.0	0.48	GLU 99		
22314320	**Good identification**					**22314320**
Actinomyces odontolyticus		100.0	0.39	GLU 99		
22315000	**Good identification**					**22315000**
Actinomyces odontolyticus		99.82	0.53	GLU 99		
Actinomyces meyeri		0.18	0.02			
22315002	**Good identification**					**22315002**
Actinomyces odontolyticus		100.0	0.49	GLU 99		
22315020	**Good identification**					**22315020**
Actinomyces odontolyticus		100.0	0.39	GLU 99		
22315022	**Good identification**					**22315022**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22315100	**Good identification**					**22315100**
Actinomyces odontolyticus		97.73	0.44	GLU 99		
Actinomyces meyeri		2.27	0.13			
22315102	**Good identification**					**22315102**
Actinomyces odontolyticus		100.0	0.40	GLU 99		
22315200	**Very good identification**					**22315200**
Actinomyces odontolyticus		100.0	0.58	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22315202	**Good identification**					**22315202**
Actinomyces odontolyticus		100.0	0.54	GLU 99		
22315220	**Good identification**					**22315220**
Actinomyces odontolyticus		100.0	0.44	GLU 99		
22315222	**Good identification**					**22315222**
Actinomyces odontolyticus		100.0	0.40	GLU 99		
22315300	**Good identification**					**22315300**
Actinomyces odontolyticus		100.0	0.49	GLU 99		
22315302	**Good identification**					**22315302**
Actinomyces odontolyticus		100.0	0.45	GLU 99		
22315320	**Good identification**					**22315320**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22377100	**Good identification**					**22377100**
Actinomyces israelii		100.0	0.37	GLU 99		
22377102	**Good identification**					**22377102**
Actinomyces israelii		100.0	0.40	GLU 99		
22377500	**Good identification**					**22377500**
Actinomyces israelii		100.0	0.35	GLU 99		
22377502	**Good identification**					**22377502**
Actinomyces israelii		100.0	0.38	GLU 99		
22510002	**Acceptable identification**					**22510002**
Actinomyces viscosus		92.22	0.38	GLU 99		
Actinomyces odontolyticus		7.78	0.27	MNS 96		
22510102	**Good identification**					**22510102**
Actinomyces viscosus		96.17	0.42	GLU 99		
Propionibacterium propionicum		2.39	0.29	MNS 96		
Actinomyces odontolyticus		1.44	0.18			
22510200	**Good identification**					**22510200**
Actinomyces odontolyticus		100.0	0.35	GLU 99		
22511102	**Low discrimination**					**22511102**
Actinomyces viscosus		88.51	0.39	GLU 99		
Lactobacillus fermentum		5.25	0.32	MNS 96		
Propionibacterium propionicum		3.59	0.29			
22512100	**Very good identification**					**22512100**
Propionibacterium propionicum		100.0	0.60	GLU 99		
22512101	**Good identification**					**22512101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.44	GLU 99		
22512102	**Very good identification**					**22512102**
Propionibacterium propionicum		99.59	0.63	GLU 99		
Actinomyces viscosus		0.41	0.08			
22512103	**Good identification**					**22512103**
Propionibacterium propionicum		98.78	0.47	GLU 99		
Actinomyces viscosus		1.22	0.00	URE 10		
22512140	**Very good identification**					**22512140**
Propionibacterium propionicum		100.0	0.60	GLU 99		
22512141	**Good identification**					**22512141**
Propionibacterium propionicum		100.0	0.44	GLU 99		
22512142	**Very good identification**					**22512142**
Propionibacterium propionicum		100.0	0.63	GLU 99		
22512143	**Good identification**					**22512143**
Propionibacterium propionicum		100.0	0.47	GLU 99		
22513100	**Very good identification**					**22513100**
Propionibacterium propionicum		99.44	0.60	GLU 99		
Actinomyces naeslundii		0.56	0.32			
22513101	**Acceptable identification**					**22513101**
Propionibacterium propionicum		92.90	0.44	GLU 99		
Actinomyces naeslundii		7.10	0.35	URE 10		
22513102	**Very good identification**					**22513102**
Propionibacterium propionicum		99.38	0.63	GLU 99		
Actinomyces naeslundii		0.38	0.32			
Actinomyces viscosus		0.25	0.04			
22513103	**Good identification**					**22513103**
Propionibacterium propionicum		95.15	0.47	GLU 99		
Actinomyces naeslundii		4.85	0.35	URE 10		
22513140	**Very good identification**					**22513140**
Propionibacterium propionicum		100.0	0.60	GLU 99		
22513141	**Good identification**					**22513141**
Propionibacterium propionicum		100.0	0.44	GLU 99		
22513142	**Very good identification**					**22513142**
Propionibacterium propionicum		100.0	0.63	GLU 99		
22513143	**Good identification**					**22513143**
Propionibacterium propionicum		100.0	0.47	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22532100	**Very good identification**					**22532100**
Propionibacterium propionicum		100.0	0.58	GLU 99		
22532101	**Good identification**					**22532101**
Propionibacterium propionicum		100.0	0.42	GLU 99		
22532102	**Very good identification**					**22532102**
Propionibacterium propionicum		100.0	0.61	GLU 99		
22532103	**Good identification**					**22532103**
Propionibacterium propionicum		100.0	0.45	GLU 99		
22532140	**Very good identification**					**22532140**
Propionibacterium propionicum		100.0	0.58	GLU 99		
22532141	**Good identification**					**22532141**
Propionibacterium propionicum		100.0	0.42	GLU 99		
22532142	**Very good identification**					**22532142**
Propionibacterium propionicum		100.0	0.61	GLU 99		
22532143	**Good identification**					**22532143**
Propionibacterium propionicum		100.0	0.45	GLU 99		
22533100	**Very good identification**					**22533100**
Propionibacterium propionicum		98.68	0.58	GLU 99		
Actinomyces naeslundii		1.33	0.37			
22533101	**Low discrimination**					**22533101**
Propionibacterium propionicum		84.65	0.42	GLU 99		
Actinomyces naeslundii		15.35	0.40	URE 10		
22533102	**Very good identification**					**22533102**
Propionibacterium propionicum		99.11	0.61	GLU 99		
Actinomyces naeslundii		0.89	0.37			
22533103	**Low discrimination**					**22533103**
Propionibacterium propionicum		89.22	0.45	GLU 99		
Actinomyces naeslundii		10.78	0.40	URE 10		
22533120	**Low discrimination**					**22533120**
Actinomyces naeslundii		57.07	0.37	GLU 99		
Propionibacterium propionicum		42.93	0.24	MNS 88		
22533121	**Good identification**					**22533121**
Actinomyces naeslundii		100.0	0.40	GLU 99		
22533123	**Good identification**					**22533123**
Actinomyces naeslundii		100.0	0.40	GLU 99		
22533140	**Very good identification**					**22533140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.76	0.58	GLU 99		
Actinomyces naeslundii		0.24	0.24			
22533141	**Good identification**					**22533141**
Propionibacterium propionicum		96.90	0.42	GLU 99		
Actinomyces naeslundii		3.10	0.27	URE 10		
22533142	**Very good identification**					**22533142**
Propionibacterium propionicum		99.84	0.61	GLU 99		
Actinomyces naeslundii		0.16	0.24			
22533143	**Good identification**					**22533143**
Propionibacterium propionicum		97.91	0.45	GLU 99		
Actinomyces naeslundii		2.09	0.27	URE 10		
22553101	**Good identification**					**22553101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
22553103	**Low discrimination**					**22553103**
Actinomyces naeslundii		86.99	0.37	GLU 99		
Propionibacterium propionicum		13.01	0.12	MNS 88		
22553121	**Good identification**					**22553121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
22553123	**Good identification**					**22553123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
22571101	**Good identification**					**22571101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
22571103	**Low discrimination**					**22571103**
Actinomyces naeslundii		81.86	0.37	GLU 99		
Actinomyces viscosus		18.14	0.01	MNS 88		
22571121	**Good identification**					**22571121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
22571123	**Good identification**					**22571123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
22573100	**Low discrimination**					**22573100**
Actinomyces naeslundii		63.80	0.39	GLU 99		
Propionibacterium propionicum		36.20	0.24	MNS 88		
22573101	**Good identification**					**22573101**
Actinomyces naeslundii		100.0	0.42	GLU 99		
22573102	**Low discrimination**					**22573102**
Actinomyces naeslundii		54.02	0.39	GLU 99		
Propionibacterium propionicum		45.98	0.27	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22573103	**Good identification**					**22573103**
Actinomyces naeslundii		100.0	0.42	GLU 99		
22573120	**Good identification**					**22573120**
Actinomyces naeslundii		100.0	0.39	GLU 99		
22573121	**Good identification**					**22573121**
Actinomyces naeslundii		100.0	0.42	GLU 99		
22573122	**Good identification**					**22573122**
Actinomyces naeslundii		100.0	0.39	GLU 99		
22573123	**Good identification**					**22573123**
Actinomyces naeslundii		100.0	0.42	GLU 99		
22710000	**Good identification**					**22710000**
Actinomyces odontolyticus		100.0	0.35	GLU 99		
22710200	**Good identification**					**22710200**
Actinomyces odontolyticus		100.0	0.40	GLU 99		
22710202	**Good identification**					**22710202**
Actinomyces odontolyticus		100.0	0.36	GLU 99		
22711101	**Good identification**					**22711101**
Actinomyces naeslundii		100.0	0.39	GLU 99		
22711120	**Low discrimination**					**22711120**
					GRA	
Actinomyces naeslundii		60.54	0.36	GLU 99	99	
Prevotella melaninogenica		21.06	-.01	MNS 88	1	
Actinomyces odontolyticus		18.40	0.10		99	
22711121	**Good identification**					**22711121**
Actinomyces naeslundii		100.0	0.39	GLU 99		
22711122	**Low discrimination**					**22711122**
Actinomyces naeslundii		84.30	0.36	GLU 99		
Actinomyces odontolyticus		15.70	0.06	MNS 88		
22711123	**Good identification**					**22711123**
Actinomyces naeslundii		100.0	0.39	GLU 99		
22711200	**Good identification**					**22711200**
Actinomyces odontolyticus		100.0	0.37	GLU 99		
22713001	**Good identification**					**22713001**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22713003	**Good identification**					**22713003**
Actinomyces naeslundii		100.0	0.36	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22713021	**Good identification**					**22713021**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22713023	**Good identification**					**22713023**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22713100	**Low discrimination**					**22713100**
Actinomyces naeslundii		66.52	0.41	GLU 99		
Propionibacterium propionicum		33.48	0.26	MNS 88		
22713101	**Good identification**					**22713101**
Actinomyces naeslundii		100.0	0.44	GLU 99		
22713102	**Low discrimination**					**22713102**
Actinomyces naeslundii		56.98	0.41	GLU 99		
Propionibacterium propionicum		43.02	0.29	MNS 88		
22713103	**Acceptable identification**					**22713103**
Actinomyces naeslundii		94.70	0.44	GLU 99		
Propionibacterium propionicum		5.30	0.12	MNS 88		
22713120	**Good identification**					**22713120**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22713121	**Good identification**					**22713121**
Actinomyces naeslundii		100.0	0.44	GLU 99		
22713122	**Good identification**					**22713122**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22713123	**Good identification**					**22713123**
Actinomyces naeslundii		100.0	0.44	GLU 99		
22731001	**Good identification**					**22731001**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22731003	**Good identification**					**22731003**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22731021	**Good identification**					**22731021**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22731023	**Good identification**					**22731023**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22731100	**Good identification**					**22731100**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22731101	**Good identification**					**22731101**
Actinomyces naeslundii		100.0	0.44	GLU 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
22731102	**Low discrimination**					**22731102**
Actinomyces naeslundii		72.84	0.41	GLU 99		
Actinomyces viscosus		27.16	0.10	MNS 88		
22731103	**Acceptable identification**					**22731103**
Actinomyces naeslundii		92.35	0.44	GLU 99		
Actinomyces viscosus		7.65	0.01	MNS 88		
22731120	**Acceptable identification**					**22731120**
Actinomyces naeslundii		92.05	0.41	GLU 99		
Clostridium perfringens		7.95	0.01	MNS 88		
22731121	**Good identification**					**22731121**
Actinomyces naeslundii		100.0	0.44	GLU 99		
22731122	**Good identification**					**22731122**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22731123	**Good identification**					**22731123**
Actinomyces naeslundii		100.0	0.44	GLU 99		
22733000	**Good identification**					**22733000**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22733001	**Good identification**					**22733001**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22733002	**Good identification**					**22733002**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22733003	**Good identification**					**22733003**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22733020	**Good identification**					**22733020**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22733021	**Good identification**					**22733021**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22733022	**Good identification**					**22733022**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22733023	**Good identification**					**22733023**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22733100	**Low discrimination**					**22733100**
Actinomyces naeslundii		82.50	0.46	GLU 99		
Propionibacterium propionicum		17.50	0.24	MNS 88		
22733101	**Good identification**					**22733101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	GLU 99		
22733102	**Low discrimination**					**22733102**
Actinomyces naeslundii		75.86	0.46	GLU 99		
Propionibacterium propionicum		24.14	0.27	MNS 88		
22733103	**Good identification**					**22733103**
Actinomyces naeslundii		100.0	0.49	GLU 99		
22733120	**Good identification**					**22733120**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22733121	**Good identification**					**22733121**
Actinomyces naeslundii		100.0	0.49	GLU 99		
22733122	**Good identification**					**22733122**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22733123	**Good identification**					**22733123**
Actinomyces naeslundii		100.0	0.49	GLU 99		
22733141	**Good identification**					**22733141**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22733143	**Good identification**					**22733143**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22733161	**Good identification**					**22733161**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22733163	**Good identification**					**22733163**
Actinomyces naeslundii		100.0	0.36	GLU 99		
22751100	**Good identification**					**22751100**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22751101	**Good identification**					**22751101**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22751103	**Low discrimination**					**22751103**
Actinomyces naeslundii		70.01	0.41	GLU 99		
Actinomyces viscosus		29.99	0.11	MNS 88		
22751120	**Good identification**					**22751120**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22751121	**Good identification**					**22751121**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22751122	**Good identification**					**22751122**
Actinomyces naeslundii		100.0	0.38	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22751123	**Good identification**					**22751123**
Actinomyces naeslundii		100.0	0.41	GLU 99		
22753000	**Good identification**					**22753000**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22753001	**Good identification**					**22753001**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22753002	**Good identification**					**22753002**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22753003	**Good identification**					**22753003**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22753020	**Good identification**					**22753020**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22753021	**Good identification**					**22753021**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22753022	**Good identification**					**22753022**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22753023	**Good identification**					**22753023**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22753100	**Low discrimination**					**22753100**
Actinomyces naeslundii		89.05	0.43	GLU 99		
Actinomyces israelii		10.95	0.18	MNS 88		
22753101	**Good identification**					**22753101**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22753102	**Low discrimination**					**22753102**
Actinomyces naeslundii		84.42	0.43	GLU 99		
Actinomyces israelii		15.58	0.21	MNS 88		
22753103	**Good identification**					**22753103**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22753120	**Good identification**					**22753120**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22753121	**Good identification**					**22753121**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22753122	**Good identification**					**22753122**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22753123	**Good identification**					**22753123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	GLU 99		
22757100	**Good identification**					**22757100**
Actinomyces israelii		100.0	0.37	GLU 99		
22757102	**Good identification**					**22757102**
Actinomyces israelii		100.0	0.40	GLU 99		
22757500	**Good identification**					**22757500**
Actinomyces israelii		100.0	0.35	GLU 99		
22757502	**Good identification**					**22757502**
Actinomyces israelii		100.0	0.38	GLU 99		
22771000	**Good identification**					**22771000**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22771001	**Good identification**					**22771001**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22771002	**Good identification**					**22771002**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22771003	**Good identification**					**22771003**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22771020	**Good identification**					**22771020**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22771021	**Good identification**					**22771021**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22771022	**Good identification**					**22771022**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22771023	**Good identification**					**22771023**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22771100	**Acceptable identification**					**22771100**
Actinomyces naeslundii		92.45	0.43	GLU 99		
Actinomyces israelii		7.55	0.15	MNS 88		
22771101	**Good identification**					**22771101**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22771102	**Low discrimination**					**22771102**
Actinomyces naeslundii		83.53	0.43	GLU 99		
Actinomyces israelii		10.23	0.18	MNS 88		
Actinomyces viscosus		6.24	0.00			
22771103	**Good identification**					**22771103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	GLU 99		
22771120	**Good identification**					**22771120**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22771121	**Good identification**					**22771121**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22771122	**Good identification**					**22771122**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22771123	**Good identification**					**22771123**
Actinomyces naeslundii		100.0	0.46	GLU 99		
22773000	**Low discrimination**					**22773000**
Actinomyces naeslundii		85.50	0.40	GLU 99		
Actinomyces israelii		14.50	0.18	MNS 88		
22773001	**Good identification**					**22773001**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22773002	**Low discrimination**					**22773002**
Actinomyces naeslundii		79.72	0.40	GLU 99		
Actinomyces israelii		20.28	0.21	MNS 88		
22773003	**Good identification**					**22773003**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22773020	**Good identification**					**22773020**
Actinomyces naeslundii		100.0	0.40	GLU 99		
22773021	**Good identification**					**22773021**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22773022	**Good identification**					**22773022**
Actinomyces naeslundii		100.0	0.40	GLU 99		
22773023	**Good identification**					**22773023**
Actinomyces naeslundii		100.0	0.43	GLU 99		
22773100	**Low discrimination**					**22773100**
Actinomyces naeslundii		80.75	0.48	GLU 99		
Actinomyces israelii		19.25	0.28	MNS 88		
22773101	**Good identification**					**22773101**
Actinomyces naeslundii		100.0	0.51	GLU 99		
22773102	**Low discrimination**					**22773102**
Actinomyces naeslundii		73.66	0.48	GLU 99		
Actinomyces israelii		26.34	0.31	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
22773103	**Good identification**					**22773103**
Actinomyces naeslundii		100.0	0.51	GLU 99		
22773120	**Good identification**					**22773120**
Actinomyces naeslundii		100.0	0.48	GLU 99		
22773121	**Good identification**					**22773121**
Actinomyces naeslundii		100.0	0.51	GLU 99		
22773122	**Good identification**					**22773122**
Actinomyces naeslundii		100.0	0.48	GLU 99		
22773123	**Good identification**					**22773123**
Actinomyces naeslundii		100.0	0.51	GLU 99		
22773140	**Low discrimination**					**22773140**
Actinomyces naeslundii		65.56	0.35	GLU 99		
Actinomyces israelii		34.44	0.21	MNS 88		
22773141	**Good identification**					**22773141**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22773142	**Low discrimination**					**22773142**
Actinomyces naeslundii		55.93	0.35	GLU 99		
Actinomyces israelii		44.07	0.24	MNS 88		
22773143	**Good identification**					**22773143**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22773160	**Good identification**					**22773160**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22773161	**Good identification**					**22773161**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22773162	**Good identification**					**22773162**
Actinomyces naeslundii		100.0	0.35	GLU 99		
22773163	**Good identification**					**22773163**
Actinomyces naeslundii		100.0	0.38	GLU 99		
22775102	**Good identification**					**22775102**
Actinomyces israelii		100.0	0.36	GLU 99		
22777000	**Good identification**					**22777000**
Actinomyces israelii		100.0	0.36	GLU 99		
22777002	**Good identification**					**22777002**
Actinomyces israelii		100.0	0.39	GLU 99		
22777100	**Good identification**					**22777100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.46	GLU 99			
22777102	**Good identification**						**22777102**
Actinomyces israelii		100.0	0.49	GLU 99			
22777112	**Good identification**						**22777112**
Actinomyces israelii		100.0	0.35	GLU 99			
22777140	**Good identification**						**22777140**
Actinomyces israelii		100.0	0.39	GLU 99			
22777142	**Good identification**						**22777142**
Actinomyces israelii		100.0	0.42	GLU 99			
22777402	**Good identification**						**22777402**
Actinomyces israelii		100.0	0.38	GLU 99			
22777500	**Good identification**						**22777500**
Actinomyces israelii		100.0	0.45	GLU 99			
22777502	**Good identification**						**22777502**
Actinomyces israelii		100.0	0.48	GLU 99			
22777540	**Good identification**						**22777540**
Actinomyces israelii		100.0	0.38	GLU 99			
22777542	**Good identification**						**22777542**
Actinomyces israelii		100.0	0.41	GLU 99			
23000002	**Low discrimination**						**23000002**
					GRA		
Atopobium minutum		47.04	0.41	NIT 1	99		
Actinomyces odontolyticus		19.44	0.34	SUC 5	99		
Campylobacter gracilis		17.79	0.15		1		
23000040	**Low discrimination**						**23000040**
					COC	GRA	
Megasphaera elsdenii		46.90	0.38	NIT 1	99	1	
Clostridium botulinum biovar A		43.21	0.33	FRU 95	1	99	
Clostridium novyi biovar A		4.92	0.29		1	99	
23000100	**Low discrimination**						**23000100**
					COC	GRA	
Actinomyces meyeri		66.74	0.35	NIT 1	1	99	
Actinomyces odontolyticus		18.56	0.29	FRU 95	1	99	
Veillonella parvula		4.08	-.03		99	1	
23000200	**Low discrimination**						**23000200**
					COC	GRA	
Actinomyces odontolyticus		89.09	0.43	MLT 99	1	99	
Veillonella parvula		3.19	-.03	FRU 92	99	1	
Campylobacter gracilis		2.34	-.03		1	1	
23000202	**Good identification**						**23000202**
Actinomyces odontolyticus		98.76	0.39	MLT 99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		1.24	0.07	FRU 92			
23001100	**Low discrimination**						**23001100**
Actinomyces meyeri		84.61	0.38	NIT 1			
Actinomyces odontolyticus		10.46	0.26	FRU 95			
Lactobacillus fermentum		1.70	0.20				
23001101	**Good identification**						**23001101**
					SPO		
Clostridium chauvoei		99.43	0.40	MNS 90	99		
Actinomyces meyeri		0.57	0.04	MLT 99	1		
23001200	**Good identification**						**23001200**
Actinomyces odontolyticus		98.78	0.40	MLT 99			
Eubacterium tortuosum		1.22	-.04	FRU 92			
23001202	**Good identification**						**23001202**
Actinomyces odontolyticus		100.0	0.36	MLT 99			
23002000	**Acceptable identification**						**23002000**
					COC		
Megasphaera elsdenii		91.07	0.41	NIT 1	99		
Veillonella parvula		2.70	-.03	FRU 95	99		
Campylobacter gracilis		1.98	-.03		1		
23002040	**Good identification**						**23002040**
					COC	GRA	
Megasphaera elsdenii		99.08	0.38	NIT 1	99	1	
Clostridium botulinum biovar A		0.92	-.01	FRU 95	1	99	
23004100	**Acceptable identification**						**23004100**
Actinomyces meyeri		93.41	0.40	NIT 1			
Actinomyces odontolyticus		6.59	0.24	FRU 95			
23004200	**Good identification**						**23004200**
Actinomyces odontolyticus		100.0	0.38	MLT 99			
23005100	**Good identification**						**23005100**
Actinomyces meyeri		96.39	0.43	NIT 1			
Actinomyces odontolyticus		3.02	0.21	FRU 95			
Lactobacillus fermentum		0.58	0.16				
23010000	**Low discrimination**						**23010000**
					COC	GRA	
Megasphaera elsdenii		65.21	0.63	NIT 1	99	1	
Actinomyces odontolyticus		16.87	0.56		1	99	
Actinomyces meyeri		7.91	0.47		1	99	
23010002	**Low discrimination**						**23010002**
					COC	GRA	
Actinomyces odontolyticus		49.97	0.52	MLT 99	1	99	
Megasphaera elsdenii		35.03	0.47		99	1	
Atopobium minutum		9.71	0.40		1	99	
23010020	**Low discrimination**						**23010020**
					CAT		
Propionibacterium acnes		57.33	0.44	IND 87	95		
Actinomyces odontolyticus		17.54	0.43	SUC 1	10		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		6.89	0.32		1		
23010022	**Low discrimination**						**23010022**
					CAT		
Actinomyces odontolyticus		63.96	0.39	MLT 99	10		
Atopobium minutum		25.44	0.32	NAG 14	1		
Propionibacterium acnes		3.45	0.10		95		
23010040	**Very good identification**						**23010040**
					COC	GRA	
Megasphaera elsdenii		96.44	0.60	NIT 1	99	1	
Pseudoramibacter alactolyticus		2.31	0.21		1	99	
Actinomyces odontolyticus		0.38	0.22		1	99	
23010042	**Good identification**						**23010042**
					COC	GRA	
Megasphaera elsdenii		95.26	0.44	NIT 1	99	1	
Pseudoramibacter alactolyticus		2.29	0.05	BGL 10	1	99	
Actinomyces odontolyticus		2.06	0.18		1	99	
23010100	**Low discrimination**						**23010100**
					GRA		
Actinomyces meyeri		79.65	0.57	NIT 1	99		
Actinomyces odontolyticus		13.41	0.47		99		
Fusobacterium mortiferum		4.56	0.45		1		
23010102	**Low discrimination**						**23010102**
Actinomyces odontolyticus		61.96	0.44	MLT 99			
Actinomyces meyeri		18.58	0.31				
Actinomyces viscosus		8.16	0.22				
23010120	**Low discrimination**						**23010120**
					CAT		
Actinomyces meyeri		81.55	0.42	NIT 1	1		
Actinomyces odontolyticus		16.39	0.34	NAG 12	10		
Propionibacterium acnes		1.71	0.10		95		
23010200	**Very good identification**						**23010200**
					COC	GRA	
Actinomyces odontolyticus		97.58	0.61	MLT 99	1	99	
Megasphaera elsdenii		1.96	0.29		99	1	
Actinomyces meyeri		0.24	0.13		1	99	
23010202	**Very good identification**						**23010202**
					COC	GRA	
Actinomyces odontolyticus		99.54	0.57	MLT 99	1	99	
Megasphaera elsdenii		0.36	0.12		99	1	
Atopobium minutum		0.10	0.06		1	99	
23010220	**Good identification**						**23010220**
					CAT		
Actinomyces odontolyticus		98.33	0.47	MLT 99	10		
Propionibacterium acnes		1.67	0.10	NAG 14	95		
23010222	**Good identification**						**23010222**
Actinomyces odontolyticus		100.0	0.44	MLT 99			
23010300	**Good identification**						**23010300**
					GRA		
Actinomyces odontolyticus		96.84	0.52	MLT 99	99		
Actinomyces meyeri		2.99	0.23		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.17	0.11		1		
23010302	**Good identification**						**23010302**
Actinomyces odontolyticus		100.0	0.49	MLT 99			
23010320	**Good identification**						**23010320**
Actinomyces odontolyticus		97.48	0.39	MLT 99			
Actinomyces meyeri		2.52	0.08	NAG 14			
23010322	**Good identification**						**23010322**
Actinomyces odontolyticus		100.0	0.35	MLT 99			
23010400	**Low discrimination**						**23010400**
					COC	GRA	
Actinomyces meyeri		67.42	0.36	NIT 1	1	99	
Megasphaera elsdenii		22.46	0.29		99	1	
Actinomyces odontolyticus		5.81	0.22		1	99	
23010500	**Good identification**						**23010500**
Actinomyces meyeri		98.48	0.47	NIT 1			
Actinomyces odontolyticus		0.67	0.13				
Lactobacillus fermentum		0.62	0.20				
23011000	**Low discrimination**						**23011000**
Actinomyces meyeri		40.41	0.50	NIT 1			
Actinomyces odontolyticus		38.29	0.53				
Lactobacillus fermentum		10.35	0.50				
23011002	**Low discrimination**						**23011002**
Actinomyces odontolyticus		60.73	0.49	MLT 99			
Lactobacillus fermentum		26.77	0.50				
Atopobium minutum		5.90	0.32				
23011020	**Low discrimination**						**23011020**
					CAT		
Propionibacterium acnes		64.90	0.44	IND 87	95		
Actinomyces odontolyticus		13.23	0.40	SUC 1	10		
Actinomyces meyeri		11.70	0.35		1		
23011022	**Low discrimination**						**23011022**
Actinomyces odontolyticus		44.91	0.36	MLT 99			
Lactobacillus fermentum		40.53	0.42	NAG 14			
Atopobium minutum		8.93	0.24				
23011100	**Low discrimination**						**23011100**
Actinomyces meyeri		80.50	0.60	NIT 1			
Lactobacillus fermentum		8.41	0.54				
Actinomyces odontolyticus		6.02	0.44				
23011101	**Acceptable identification**						**23011101**
					SPO		
Clostridium chauvoei		94.18	0.47	MNS 90	99		
Actinomyces meyeri		4.38	0.26	MLT 99	1		
Lactobacillus fermentum		0.46	0.20		1		
23011102	**Low discrimination**						**23011102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		52.53	0.54	NIT 1		
Actinomyces odontolyticus		23.06	0.41			
Actinomyces meyeri		15.56	0.34			
23011120	**Low discrimination**					**23011120**
Actinomyces meyeri		71.03	0.45	NIT 1		
Lactobacillus fermentum		18.13	0.46	NAG 12		
Actinomyces odontolyticus		6.34	0.31			
23011121	**Good identification**					**23011121**
					SPO	
Clostridium chauvoei		97.40	0.38	MNS 90	99	
Actinomyces meyeri		1.85	0.11	MLT 99	1	
Actinomyces naeslundii		0.75	0.25		1	
23011122	**Low discrimination**					**23011122**
Lactobacillus fermentum		70.84	0.46	NIT 1		
Actinomyces odontolyticus		15.19	0.27			
Actinomyces meyeri		8.58	0.19			
23011200	**Very good identification**					**23011200**
Actinomyces odontolyticus		99.32	0.58	MLT 99		
Actinomyces meyeri		0.55	0.16			
Lactobacillus fermentum		0.14	0.16			
23011202	**Good identification**					**23011202**
Actinomyces odontolyticus		99.77	0.54	MLT 99		
Lactobacillus fermentum		0.23	0.16			
23011220	**Good identification**					**23011220**
					CAT	
Actinomyces odontolyticus		97.51	0.44	MLT 99	10	
Propionibacterium acnes		2.49	0.10	NAG 14	95	
23011222	**Good identification**					**23011222**
Actinomyces odontolyticus		100.0	0.41	MLT 99		
23011300	**Acceptable identification**					**23011300**
Actinomyces odontolyticus		92.52	0.49	MLT 99		
Actinomyces meyeri		6.44	0.26			
Lactobacillus fermentum		0.67	0.20			
23011302	**Good identification**					**23011302**
Actinomyces odontolyticus		98.83	0.46	MLT 99		
Lactobacillus fermentum		1.17	0.20			
23011320	**Acceptable identification**					**23011320**
Actinomyces odontolyticus		94.49	0.36	MLT 99		
Actinomyces meyeri		5.51	0.11	NAG 14		
23011400	**Low discrimination**					**23011400**
Lactobacillus fermentum		49.66	0.50	NIT 1		
Actinomyces meyeri		48.49	0.39			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces odontolyticus		1.86	0.19					
23011402	**Acceptable identification**							**23011402**
Lactobacillus fermentum		94.96	0.50	NIT 1				
Actinomyces meyeri		2.87	0.14					
Actinomyces odontolyticus		2.18	0.15					
23011420	**Low discrimination**							**23011420**
					CAT			
Lactobacillus fermentum		66.35	0.42	NIT 1	1			
Actinomyces meyeri		26.50	0.25		1			
Propionibacterium acnes		5.94	0.10		95			
23011422	**Good identification**							**23011422**
Lactobacillus fermentum		100.0	0.42	NIT 1				
23011500	**Low discrimination**							**23011500**
Actinomyces meyeri		70.27	0.50	NIT 1				
Lactobacillus fermentum		29.35	0.54					
Actinomyces odontolyticus		0.21	0.10					
23011502	**Acceptable identification**							**23011502**
Lactobacillus fermentum		92.72	0.54	NIT 1				
Actinomyces meyeri		6.87	0.24					
Actinomyces odontolyticus		0.41	0.07					
23011520	**Low discrimination**							**23011520**
Lactobacillus fermentum		50.52	0.46	NIT 1				
Actinomyces meyeri		49.48	0.35					
23011522	**Good identification**							**23011522**
Lactobacillus fermentum		97.06	0.46	NIT 1				
Actinomyces meyeri		2.94	0.09					
23012000	**Good identification**							**23012000**
					COC	GRA		
Megasphaera elsdenii		93.74	0.63	NIT 1	99	1		
Pseudoramibacter alactolyticus		3.23	0.27		1	99		
Eubacterium limosum		2.05	0.26		1	99		
23012002	**Acceptable identification**							**23012002**
					COC	GRA		
Megasphaera elsdenii		91.79	0.47	NIT 1	99	1		
Propionibacterium propionicum		3.18	0.32	BGL 10	1	99		
Pseudoramibacter alactolyticus		3.16	0.11		1	99		
23012020	**Low discrimination**							**23012020**
					COC	CAT	GRA	
Propionibacterium acnes		82.03	0.40	IND 87	1	95	99	
Megasphaera elsdenii		10.70	0.29	SUC 1	99	1	1	
Pseudoramibacter alactolyticus		6.44	0.14		1	1	99	
23012040	**Very good identification**							**23012040**
					COC	GRA		
Megasphaera elsdenii		98.35	0.60	NIT 1	99	1		
Pseudoramibacter alactolyticus		1.27	0.17		1	99		
Propionibacterium propionicum		0.38	0.29		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23012042	**Acceptable identification**						**23012042**
					COC	GRA	
Megasphaera elsdenii		93.91	0.44	NIT 1	99	1	
Propionibacterium propionicum		4.88	0.32	BGL 10	1	99	
Pseudoramibacter alactolyticus		1.21	0.00		1	99	
23012100	**Good identification**						**23012100**
					COC	GRA	
Propionibacterium propionicum		93.92	0.63	MLT 99	1	99	
Megasphaera elsdenii		3.74	0.29		99	1	
Actinomyces meyeri		1.81	0.23		1	99	
23012101	**Good identification**						**23012101**
Propionibacterium propionicum		100.0	0.47	MLT 99			
23012102	**Very good identification**						**23012102**
					COC	GRA	
Propionibacterium propionicum		99.58	0.66	MLT 99	1	99	
Megasphaera elsdenii		0.29	0.12		99	1	
Actinomyces odontolyticus		0.13	0.10		1	99	
23012103	**Good identification**						**23012103**
Propionibacterium propionicum		100.0	0.50	MLT 99			
23012140	**Very good identification**						**23012140**
					COC	GRA	
Propionibacterium propionicum		97.42	0.63	MLT 99	1	99	
Megasphaera elsdenii		2.58	0.26		99	1	
23012141	**Good identification**						**23012141**
Propionibacterium propionicum		100.0	0.47	MLT 99			
23012142	**Very good identification**						**23012142**
					COC	GRA	
Propionibacterium propionicum		99.80	0.66	MLT 99	1	99	
Megasphaera elsdenii		0.20	0.09		99	1	
23012143	**Good identification**						**23012143**
Propionibacterium propionicum		100.0	0.50	MLT 99			
23013020	**Good identification**						**23013020**
					CAT		
Propionibacterium acnes		99.18	0.40	IND 87	95		
Propionibacterium granulosum		0.45	0.09	SUC 1	90		
Actinomyces odontolyticus		0.36	0.05		10		
23013100	**Very good identification**						**23013100**
Propionibacterium propionicum		96.35	0.63	MLT 99			
Actinomyces meyeri		2.79	0.26				
Lactobacillus fermentum		0.29	0.20				
23013101	**Acceptable identification**						**23013101**
					SPO		
Propionibacterium propionicum		92.20	0.47	MLT 99	1		
Clostridium chauvoei		5.22	0.12	URE 10	99		
Actinomyces naeslundii		2.58	0.30		1		
23013102	**Very good identification**						**23013102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		99.57	0.66	MLT 99			
Lactobacillus fermentum		0.20	0.20				
Actinomyces naeslundii		0.14	0.27				
23013103	**Good identification**						**23013103**
					SPO		
Propionibacterium propionicum		97.53	0.50	MLT 99	1		
Actinomyces naeslundii		1.82	0.30	URE 10	1		
Clostridium chauvoei		0.65	-.01		99		
23013140	**Very good identification**						**23013140**
Propionibacterium propionicum		100.0	0.63	MLT 99			
23013141	**Good identification**						**23013141**
Propionibacterium propionicum		100.0	0.47	MLT 99			
23013142	**Very good identification**						**23013142**
Propionibacterium propionicum		100.0	0.66	MLT 99			
23013143	**Good identification**						**23013143**
Propionibacterium propionicum		100.0	0.50	MLT 99			
23014000	**Low discrimination**						**23014000**
					COC	GRA	
Actinomyces meyeri		62.38	0.52	NIT 1	1	99	
Actinomyces odontolyticus		33.74	0.51		1	99	
Megasphaera elsdenii		2.68	0.29		99	1	
23014002	**Low discrimination**						**23014002**
					COC	GRA	
Actinomyces odontolyticus		89.67	0.47	MLT 99	1	99	
Actinomyces meyeri		8.37	0.26		1	99	
Megasphaera elsdenii		1.29	0.12		99	1	
23014020	**Low discrimination**						**23014020**
					CAT		
Actinomyces meyeri		59.20	0.37	NIT 1	1		
Actinomyces odontolyticus		38.23	0.37	NAG 12	10		
Propionibacterium acnes		2.56	0.10		95		
23014100	**Very good identification**						**23014100**
					GRA		
Actinomyces meyeri		95.66	0.62	NIT 1	99		
Actinomyces odontolyticus		4.09	0.42		99		
Fusobacterium mortiferum		0.21	0.26		1		
23014102	**Low discrimination**						**23014102**
Actinomyces meyeri		53.71	0.36	NIT 1			
Actinomyces odontolyticus		45.46	0.39	BGL 3			
Lactobacillus fermentum		0.83	0.16				
23014120	**Good identification**						**23014120**
Actinomyces meyeri		95.15	0.47	NIT 1			
Actinomyces odontolyticus		4.85	0.29	NAG 12			
23014200	**Good identification**						**23014200**
Actinomyces odontolyticus		99.05	0.56	MLT 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		0.95	0.18			
23014202	**Good identification**					**23014202**
Actinomyces odontolyticus		100.0	0.52	MLT 99		
23014220	**Good identification**					**23014220**
Actinomyces odontolyticus		99.20	0.42	MLT 99		
Actinomyces meyeri		0.80	0.03	NAG 14		
23014222	**Good identification**					**23014222**
Actinomyces odontolyticus		100.0	0.39	MLT 99		
23014300	**Low discrimination**					**23014300**
Actinomyces odontolyticus		89.14	0.47	MLT 99		
Actinomyces meyeri		10.86	0.28			
23014302	**Good identification**					**23014302**
Actinomyces odontolyticus		99.39	0.43	MLT 99		
Actinomyces meyeri		0.61	0.02			
23014400	**Good identification**					**23014400**
Actinomyces meyeri		97.41	0.41	NIT 1		
Actinomyces odontolyticus		2.13	0.17			
Lactobacillus fermentum		0.46	0.12			
23014500	**Good identification**					**23014500**
Actinomyces meyeri		99.64	0.52	NIT 1		
Lactobacillus fermentum		0.19	0.16			
Actinomyces odontolyticus		0.17	0.08			
23014520	**Good identification**					**23014520**
Actinomyces meyeri		100.0	0.37	NIT 1		
23015000	**Low discrimination**					**23015000**
Actinomyces meyeri		75.66	0.55	NIT 1		
Actinomyces odontolyticus		18.19	0.48			
Lactobacillus fermentum		5.86	0.46			
23015002	**Low discrimination**					**23015002**
Actinomyces odontolyticus		57.62	0.44	MLT 99		
Lactobacillus fermentum		30.29	0.46			
Actinomyces meyeri		12.09	0.29			
23015020	**Low discrimination**					**23015020**
Actinomyces meyeri		66.44	0.40	NIT 1		
Actinomyces odontolyticus		19.07	0.34	NAG 12		
Lactobacillus fermentum		12.58	0.38			
23015022	**Low discrimination**					**23015022**
Lactobacillus fermentum		47.79	0.38	NIT 1		
Actinomyces odontolyticus		44.40	0.31			
Actinomyces meyeri		7.81	0.14			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23015100	**Good identification**					**23015100**
Actinomyces meyeri		94.99	0.65	NIT 1		
Lactobacillus fermentum		3.00	0.50			
Actinomyces odontolyticus		1.80	0.39			
23015102	**Low discrimination**					**23015102**
Lactobacillus fermentum		42.47	0.50	NIT 1		
Actinomyces meyeri		41.57	0.39			
Actinomyces odontolyticus		15.64	0.36			
23015120	**Acceptable identification**					**23015120**
Actinomyces meyeri		90.92	0.50	NIT 1		
Lactobacillus fermentum		7.02	0.42	NAG 12		
Actinomyces odontolyticus		2.06	0.26			
23015122	**Low discrimination**					**23015122**
Lactobacillus fermentum		63.28	0.42	NIT 1		
Actinomyces meyeri		25.34	0.24			
Actinomyces odontolyticus		11.38	0.22			
23015200	**Good identification**					**23015200**
Actinomyces odontolyticus		97.72	0.53	MLT 99		
Actinomyces meyeri		2.12	0.21			
Lactobacillus fermentum		0.16	0.12			
23015202	**Good identification**					**23015202**
Actinomyces odontolyticus		99.73	0.49	MLT 99		
Lactobacillus fermentum		0.27	0.12			
23015220	**Good identification**					**23015220**
Actinomyces odontolyticus		98.22	0.39	MLT 99		
Actinomyces meyeri		1.78	0.06	NAG 14		
23015222	**Good identification**					**23015222**
Actinomyces odontolyticus		100.0	0.36	MLT 99		
23015300	**Low discrimination**					**23015300**
Actinomyces odontolyticus		77.95	0.44	MLT 99		
Actinomyces meyeri		21.37	0.31			
Lactobacillus fermentum		0.68	0.16			
23015302	**Good identification**					**23015302**
Actinomyces odontolyticus		97.28	0.40	MLT 99		
Lactobacillus fermentum		1.38	0.16			
Actinomyces meyeri		1.35	0.05			
23015400	**Low discrimination**					**23015400**
Actinomyces meyeri		75.78	0.44	NIT 1		
Lactobacillus fermentum		23.48	0.46			
Actinomyces odontolyticus		0.74	0.14			
23015402	**Low discrimination**					**23015402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		89.36	0.46	NIT 1		
Actinomyces meyeri		8.92	0.19			
Actinomyces odontolyticus		1.72	0.10			
23015422	**Good identification**					**23015422**
Lactobacillus fermentum		96.08	0.38	NIT 1		
Actinomyces meyeri		3.92	0.04			
23015500	**Low discrimination**					**23015500**
Actinomyces meyeri		88.78	0.55	NIT 1		
Lactobacillus fermentum		11.22	0.50			
23015502	**Low discrimination**					**23015502**
Lactobacillus fermentum		80.34	0.50	NIT 1		
Actinomyces meyeri		19.66	0.29			
23015520	**Low discrimination**					**23015520**
Actinomyces meyeri		76.40	0.40	NIT 1		
Lactobacillus fermentum		23.60	0.42	NAG 12		
23015522	**Acceptable identification**					**23015522**
Lactobacillus fermentum		90.90	0.42	NIT 1		
Actinomyces meyeri		9.10	0.14			
23030100	**Low discrimination**					**23030100**
					GRA	
Fusobacterium mortiferum		51.46	0.36	NIT 1	1	
Actinomyces meyeri		30.40	0.23		99	
Propionibacterium propionicum		13.02	0.27		99	
23031100	**Low discrimination**					**23031100**
					GRA	
Fusobacterium mortiferum		52.42	0.39	NIT 1	1	
Actinomyces meyeri		30.97	0.26		99	
Propionibacterium propionicum		8.84	0.27		99	
23032100	**Very good identification**					**23032100**
Propionibacterium propionicum		100.0	0.61	MLT 99		
23032101	**Good identification**					**23032101**
Propionibacterium propionicum		100.0	0.45	MLT 99		
23032102	**Very good identification**					**23032102**
Propionibacterium propionicum		100.0	0.64	MLT 99		
23032103	**Good identification**					**23032103**
Propionibacterium propionicum		100.0	0.48	MLT 99		
23032140	**Very good identification**					**23032140**
Propionibacterium propionicum		100.0	0.61	MLT 99		
23032141	**Good identification**					**23032141**
Propionibacterium propionicum		100.0	0.45	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23032142	**Very good identification**					**23032142**
Propionibacterium propionicum		100.0	0.64	MLT 99		
23032143	**Good identification**					**23032143**
Propionibacterium propionicum		100.0	0.48	MLT 99		
23033100	**Very good identification**					**23033100**
Propionibacterium propionicum		99.51	0.61	MLT 99		
Actinomyces naeslundii		0.49	0.32			
23033101	**Acceptable identification**					**23033101**
Propionibacterium propionicum		93.77	0.45	MLT 99		
Actinomyces naeslundii		6.23	0.35	URE 10		
23033102	**Very good identification**					**23033102**
Propionibacterium propionicum		99.67	0.64	MLT 99		
Actinomyces naeslundii		0.33	0.32			
23033103	**Good identification**					**23033103**
Propionibacterium propionicum		95.76	0.48	MLT 99		
Actinomyces naeslundii		4.24	0.35	URE 10		
23033121	**Good identification**					**23033121**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23033123	**Low discrimination**					**23033123**
Actinomyces naeslundii		81.42	0.35	MNS 88		
Propionibacterium propionicum		18.58	0.14	MLT 94		
23033140	**Very good identification**					**23033140**
Propionibacterium propionicum		100.0	0.61	MLT 99		
23033141	**Good identification**					**23033141**
Propionibacterium propionicum		100.0	0.45	MLT 99		
23033142	**Very good identification**					**23033142**
Propionibacterium propionicum		100.0	0.64	MLT 99		
23033143	**Good identification**					**23033143**
Propionibacterium propionicum		100.0	0.48	MLT 99		
23050100	**Low discrimination**					**23050100**
					GRA	
Fusobacterium mortiferum		61.79	0.37	NIT 1	1	
Actinomyces meyeri		32.71	0.23		99	
Actinomyces odontolyticus		5.51	0.13		99	
23051100	**Low discrimination**					**23051100**
					GRA	
Fusobacterium mortiferum		60.60	0.40	NIT 1	1	
Actinomyces meyeri		32.08	0.26		99	
Lactobacillus fermentum		3.35	0.20		99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
23073101	**Good identification**							**23073101**
Actinomyces naeslundii		100.0	0.37	MNS 88				
23073103	**Low discrimination**							**23073103**
Actinomyces naeslundii		85.31	0.37	MNS 88				
Propionibacterium propionicum		14.69	0.14	MLT 94				
23073121	**Good identification**							**23073121**
Actinomyces naeslundii		100.0	0.37	MNS 88				
23073123	**Good identification**							**23073123**
Actinomyces naeslundii		100.0	0.37	MNS 88				
23100000	**Low discrimination**							**23100000**
					COC	SPO	GRA	
Actinomyces odontolyticus		89.61	0.72	FRU 92	1	1	99	
Clostridium botulinum biovar A		3.56	0.41		1	99	99	
Megasphaera elsdenii		3.18	0.44		99	1	1	
23100001	**Low discrimination**							**23100001**
					SPO			
Actinomyces odontolyticus		70.89	0.38	FRU 92	1			
Clostridium novyi biovar A		17.39	0.32	URE 1	99			
Clostridium botulinum biovar A		2.82	0.06		99			
23100002	**Very good identification**							**23100002**
					COC	GRA		
Actinomyces odontolyticus		97.65	0.68	FRU 92	1	99		
Actinomyces viscosus		1.38	0.30		1	99		
Megasphaera elsdenii		0.63	0.28		99	1		
23100010	**Low discrimination**							**23100010**
					SPO			
Actinomyces odontolyticus		74.87	0.38	FRU 92	1			
Clostridium novyi biovar A		18.37	0.32	MLZ 1	99			
Clostridium botulinum biovar A		2.98	0.06		99			
23100020	**Very good identification**							**23100020**
					COC	SPO		
Actinomyces odontolyticus		98.20	0.59	FRU 92	1	1		
Gemella morbillorum		0.45	0.14	NAG 14	99	1		
Clostridium novyi biovar A		0.37	0.21		1	99		
23100022	**Good identification**							**23100022**
Actinomyces odontolyticus		100.0	0.55	FRU 92				
23100040	**Low discrimination**							**23100040**
					COC	SPO	GRA	
Megasphaera elsdenii		48.34	0.41	NIT 1	99	1	1	
Clostridium botulinum biovar A		24.30	0.32	FRU 95	1	99	99	
Actinomyces odontolyticus		20.66	0.38		1	1	99	
23100100	**Very good identification**							**23100100**
Actinomyces odontolyticus		95.33	0.63	FRU 92				
Actinomyces meyeri		3.91	0.36					
Propionibacterium propionicum		0.35	0.29					
23100101	**Low discrimination**							**23100101**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Clostridium chauvoei		89.83	0.40	MNS 90	99			
Actinomyces odontolyticus		9.41	0.29	GAL 99	1			
Actinomyces meyeri		0.39	0.02		1			
23100102	**Good identification**							**23100102**
Actinomyces odontolyticus		91.68	0.60	FRU 92				
Actinomyces viscosus		7.31	0.34					
Propionibacterium propionicum		0.82	0.32					
23100120	**Good identification**							**23100120**
					GRA			
Actinomyces odontolyticus		95.70	0.50	FRU 92	99			
Actinomyces meyeri		3.28	0.21	NAG 14	99			
Prevotella bivia		1.02	-.03		1			
23100122	**Good identification**							**23100122**
Actinomyces odontolyticus		99.51	0.46	FRU 92				
Actinomyces viscosus		0.49	0.00	NAG 14				
23100200	**Very good identification**							**23100200**
					COC	SPO	GRA	
Actinomyces odontolyticus		99.95	0.77	FRU 92	1	1	99	
Clostridium botulinum biovar A		0.02	0.06		1	99	99	
Megasphaera elsdenii		0.02	0.10		99	1	1	
23100201	**Good identification**							**23100201**
Actinomyces odontolyticus		100.0	0.43	FRU 92				
23100202	**Very good identification**							**23100202**
Actinomyces odontolyticus		100.0	0.73	FRU 92				
23100203	**Good identification**							**23100203**
Actinomyces odontolyticus		100.0	0.39	FRU 92				
23100210	**Good identification**							**23100210**
Actinomyces odontolyticus		100.0	0.43	FRU 92				
23100212	**Good identification**							**23100212**
Actinomyces odontolyticus		100.0	0.39	FRU 92				
23100220	**Very good identification**							**23100220**
Actinomyces odontolyticus		100.0	0.63	FRU 92				
23100222	**Very good identification**							**23100222**
Actinomyces odontolyticus		100.0	0.60	FRU 92				
23100240	**Good identification**							**23100240**
					COC	GRA		
Actinomyces odontolyticus		98.80	0.43	FRU 92	1	99		
Megasphaera elsdenii		1.20	0.07	SOR 1	99	1		
23100242	**Good identification**							**23100242**
Actinomyces odontolyticus		100.0	0.39	FRU 92				
23100300	**Very good identification**							**23100300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.98	0.68	FRU 92		
Actinomyces meyeri		0.02	0.02			
23100302	**Very good identification**					**23100302**
Actinomyces odontolyticus		99.96	0.65	FRU 92		
Actinomyces viscosus		0.04	0.00			
23100320	**Good identification**					**23100320**
Actinomyces odontolyticus		100.0	0.55	FRU 92		
23100322	**Good identification**					**23100322**
Actinomyces odontolyticus		100.0	0.51	FRU 92		
23100400	**Low discrimination**					**23100400**
					COC	
Actinomyces odontolyticus		80.35	0.38	FRU 92	1	
Actinomyces meyeri		6.43	0.15	ARA 1	1	
Gemella morbillorum		5.96	0.14		99	
23100600	**Good identification**					**23100600**
Actinomyces odontolyticus		100.0	0.43	FRU 92		
23100602	**Good identification**					**23100602**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23101000	**Very good identification**					**23101000**
					COC	
Actinomyces odontolyticus		98.41	0.69	FRU 92	1	
Actinomyces meyeri		0.72	0.29		1	
Gemella morbillorum		0.54	0.26		99	
23101001	**Low discrimination**					**23101001**
					SPO	
Clostridium chauvoei		81.29	0.40	MNS 90	99	
Actinomyces odontolyticus		17.98	0.35	LAC 99	1	
Clostridium novyi biovar A		0.74	0.15		99	
23101002	**Very good identification**					**23101002**
Actinomyces odontolyticus		98.53	0.65	FRU 92		
Actinomyces viscosus		1.28	0.26			
Lactobacillus fermentum		0.15	0.24			
23101020	**Good identification**					**23101020**
					COC	
Actinomyces odontolyticus		98.66	0.56	FRU 92	1	
Actinomyces meyeri		0.60	0.14	NAG 14	1	
Gemella morbillorum		0.37	0.09		99	
23101022	**Good identification**					**23101022**
Actinomyces odontolyticus		99.68	0.52	FRU 92		
Lactobacillus fermentum		0.32	0.16	NAG 14		
23101100	**Low discrimination**					**23101100**
					SPO	
Actinomyces odontolyticus		80.12	0.60	FRU 92	1	
Clostridium chauvoei		11.59	0.40		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		7.38	0.39		1		
23101101	**Very good identification**						**23101101**
					SPO		
Clostridium chauvoei		99.91	0.74	MNS 90	99		
Actinomyces odontolyticus		0.07	0.26		1		
Actinomyces meyeri		0.01	0.05		1		
23101102	**Low discrimination**						**23101102**
					SPO		
Actinomyces odontolyticus		87.64	0.57	FRU 92	1		
Actinomyces viscosus		6.42	0.30		1		
Clostridium chauvoei		3.65	0.27		99		
23101103	**Very good identification**						**23101103**
					SPO		
Clostridium chauvoei		99.11	0.62	MNS 90	99		
Actinomyces viscosus		0.59	0.22		1		
Actinomyces odontolyticus		0.24	0.23		1		
23101111	**Good identification**						**23101111**
Clostridium chauvoei		100.0	0.40	MNS 90			
23101120	**Low discrimination**						**23101120**
					GRA	SPO	
Actinomyces odontolyticus		66.02	0.47	FRU 92	99	1	
Clostridium chauvoei		19.55	0.32	NAG 14	99	99	
Prevotella melaninogenica		5.80	0.17		1	1	
23101121	**Very good identification**						**23101121**
					SPO		
Clostridium chauvoei		99.95	0.66	MNS 90	99		
Actinomyces odontolyticus		0.03	0.13		1		
Actinomyces naeslundii		0.02	0.24		1		
23101122	**Low discrimination**						**23101122**
					SPO		
Actinomyces odontolyticus		85.46	0.43	FRU 92	1		
Clostridium chauvoei		7.29	0.19	NAG 14	99		
Clostridium perfringens		2.85	0.13		84		
23101123	**Good identification**						**23101123**
					SPO		
Clostridium chauvoei		99.80	0.53	MNS 90	99		
Actinomyces odontolyticus		0.12	0.09		1		
Actinomyces naeslundii		0.09	0.24		1		
23101141	**Good identification**						**23101141**
					SPO		
Clostridium chauvoei		99.58	0.40	MNS 90	99		
Propionibacterium propionicum		0.42	0.12	SOR 1	1		
23101200	**Very good identification**						**23101200**
Actinomyces odontolyticus		100.0	0.74	FRU 92			
23101201	**Good identification**						**23101201**
					SPO		
Actinomyces odontolyticus		97.70	0.40	FRU 92	1		
Clostridium chauvoei		2.30	0.06	URE 1	99		
23101202	**Very good identification**						**23101202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.70	FRU 92		
23101203	**Good identification**					**23101203**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23101210	**Good identification**					**23101210**
Actinomyces odontolyticus		100.0	0.40	FRU 92		
23101212	**Good identification**					**23101212**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23101220	**Very good identification**					**23101220**
Actinomyces odontolyticus		100.0	0.60	FRU 92		
23101222	**Very good identification**					**23101222**
Actinomyces odontolyticus		100.0	0.57	FRU 92		
23101240	**Good identification**					**23101240**
Actinomyces odontolyticus		100.0	0.40	FRU 92		
23101242	**Good identification**					**23101242**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23101300	**Very good identification**					**23101300**
					SPO	
Actinomyces odontolyticus		99.88	0.65	FRU 92	1	
Clostridium chauvoei		0.08	0.06		99	
Actinomyces meyeri		0.05	0.05		1	
23101301	**Low discrimination**					**23101301**
					SPO	
Clostridium chauvoei		88.06	0.40	MNS 90	99	
Actinomyces odontolyticus		11.94	0.31	RHA 1	1	
23101302	**Very good identification**					**23101302**
Actinomyces odontolyticus		100.0	0.62	FRU 92		
23101320	**Good identification**					**23101320**
Actinomyces odontolyticus		100.0	0.52	FRU 92		
23101322	**Good identification**					**23101322**
Actinomyces odontolyticus		100.0	0.48	FRU 92		
23101501	**Good identification**					**23101501**
Clostridium chauvoei		100.0	0.40	MNS 90		
23101600	**Good identification**					**23101600**
Actinomyces odontolyticus		100.0	0.40	FRU 92		
23101602	**Good identification**					**23101602**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23102000	**Low discrimination**					**23102000**
					COC GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Megasphaera elsdenii		75.02	0.44	NIT 1	99	1	
Actinomyces odontolyticus		21.37	0.38	FRU 95	1	99	
Propionibacterium propionicum		2.44	0.29		1	99	
23102040	**Good identification**						**23102040**
					COC	GRA	
Megasphaera elsdenii		95.35	0.41	NIT 1	99	1	
Propionibacterium propionicum		4.65	0.29	FRU 95	1	99	
23102100	**Very good identification**						**23102100**
					COC	GRA	
Propionibacterium propionicum		96.88	0.63	FRU 99	1	99	
Actinomyces odontolyticus		2.71	0.29		1	99	
Megasphaera elsdenii		0.30	0.10		99	1	
23102101	**Good identification**						**23102101**
					SPO		
Propionibacterium propionicum		97.63	0.47	FRU 99	1		
Clostridium chauvoei		2.37	0.06	URE 10	99		
23102102	**Very good identification**						**23102102**
Propionibacterium propionicum		98.78	0.66	FRU 99			
Actinomyces odontolyticus		1.13	0.26				
Actinomyces viscosus		0.09	0.00				
23102103	**Good identification**						**23102103**
Propionibacterium propionicum		100.0	0.50	FRU 99			
23102140	**Very good identification**						**23102140**
					COC	GRA	
Propionibacterium propionicum		99.79	0.63	FRU 99	1	99	
Megasphaera elsdenii		0.21	0.07		99	1	
23102141	**Good identification**						**23102141**
Propionibacterium propionicum		100.0	0.47	FRU 99			
23102142	**Very good identification**						**23102142**
Propionibacterium propionicum		100.0	0.66	FRU 99			
23102143	**Good identification**						**23102143**
Propionibacterium propionicum		100.0	0.50	FRU 99			
23102200	**Good identification**						**23102200**
					COC	GRA	
Actinomyces odontolyticus		98.21	0.43	FRU 92	1	99	
Megasphaera elsdenii		1.79	0.10	MAN 1	99	1	
23102202	**Good identification**						**23102202**
Actinomyces odontolyticus		100.0	0.39	FRU 92			
23103100	**Very good identification**						**23103100**
					SPO		
Propionibacterium propionicum		97.58	0.63	FRU 99	1		
Actinomyces odontolyticus		1.82	0.26		1		
Clostridium chauvoei		0.26	0.06		99		
23103101	**Low discrimination**						**23103101**
					SPO		
Clostridium chauvoei		70.13	0.40	MNS 90	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		29.20	0.47	MAN 1	1	
Actinomyces naeslundii		0.67	0.29		1	
23103102	**Very good identification**					**23103102**
Propionibacterium propionicum		99.13	0.66	FRU 99		
Actinomyces odontolyticus		0.76	0.23			
Actinomyces naeslundii		0.11	0.26			
23103103	**Low discrimination**					**23103103**
					SPO	
Propionibacterium propionicum		77.05	0.50	FRU 99	1	
Clostridium chauvoei		21.77	0.27	URE 10	99	
Actinomyces naeslundii		1.19	0.29		1	
23103140	**Very good identification**					**23103140**
Propionibacterium propionicum		100.0	0.63	FRU 99		
23103141	**Good identification**					**23103141**
					SPO	
Propionibacterium propionicum		97.63	0.47	FRU 99	1	
Clostridium chauvoei		2.37	0.06	URE 10	99	
23103142	**Very good identification**					**23103142**
Propionibacterium propionicum		100.0	0.66	FRU 99		
23103143	**Good identification**					**23103143**
Propionibacterium propionicum		100.0	0.50	FRU 99		
23103200	**Good identification**					**23103200**
Actinomyces odontolyticus		100.0	0.40	FRU 92		
23103202	**Good identification**					**23103202**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23104000	**Very good identification**					**23104000**
					COC	
Actinomyces odontolyticus		98.41	0.67	FRU 92	1	
Actinomyces meyeri		1.25	0.31		1	
Gemella morbillorum		0.15	0.14		99	
23104002	**Very good identification**					**23104002**
Actinomyces odontolyticus		99.94	0.63	FRU 92		
Actinomyces meyeri		0.06	0.05			
23104020	**Good identification**					**23104020**
Actinomyces odontolyticus		98.94	0.53	FRU 92		
Actinomyces meyeri		1.06	0.16	NAG 14		
23104022	**Good identification**					**23104022**
Actinomyces odontolyticus		100.0	0.50	FRU 92		
23104100	**Low discrimination**					**23104100**
Actinomyces odontolyticus		86.10	0.58	FRU 92		
Actinomyces meyeri		13.90	0.41			
23104102	**Good identification**					**23104102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.03	0.55	FRU 92		
Actinomyces meyeri		0.81	0.15			
Actinomyces viscosus		0.16	0.00			
23104120	**Low discrimination**					**23104120**
Actinomyces odontolyticus		88.09	0.45	FRU 92		
Actinomyces meyeri		11.91	0.26	NAG 14		
23104122	**Good identification**					**23104122**
Actinomyces odontolyticus		100.0	0.41	FRU 92		
23104200	**Very good identification**					**23104200**
Actinomyces odontolyticus		100.0	0.72	FRU 92		
23104201	**Good identification**					**23104201**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23104202	**Very good identification**					**23104202**
Actinomyces odontolyticus		100.0	0.68	FRU 92		
23104210	**Good identification**					**23104210**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23104220	**Very good identification**					**23104220**
Actinomyces odontolyticus		100.0	0.58	FRU 92		
23104222	**Good identification**					**23104222**
Actinomyces odontolyticus		100.0	0.55	FRU 92		
23104240	**Good identification**					**23104240**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23104300	**Very good identification**					**23104300**
Actinomyces odontolyticus		99.92	0.63	FRU 92		
Actinomyces meyeri		0.08	0.07			
23104302	**Very good identification**					**23104302**
Actinomyces odontolyticus		100.0	0.59	FRU 92		
23104320	**Good identification**					**23104320**
Actinomyces odontolyticus		100.0	0.50	FRU 92		
23104322	**Good identification**					**23104322**
Actinomyces odontolyticus		100.0	0.46	FRU 92		
23104600	**Good identification**					**23104600**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23105000	**Very good identification**					**23105000**
Actinomyces odontolyticus		96.99	0.64	FRU 92	COC 1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		2.78	0.34		1	
Gemella morbillorum		0.12	0.09		99	
23105002	**Very good identification**					**23105002**
Actinomyces odontolyticus		99.67	0.60	FRU 92		
Lactobacillus fermentum		0.18	0.20			
Actinomyces meyeri		0.14	0.08			
23105020	**Good identification**					**23105020**
Actinomyces odontolyticus		97.43	0.50	FRU 92		
Actinomyces meyeri		2.34	0.19	NAG 14		
Lactobacillus fermentum		0.23	0.12			
23105022	**Good identification**					**23105022**
Actinomyces odontolyticus		99.62	0.47	FRU 92		
Lactobacillus fermentum		0.38	0.12	NAG 14		
23105100	**Low discrimination**					**23105100**
Actinomyces odontolyticus		72.88	0.55	FRU 92		
Actinomyces meyeri		26.47	0.44			
Lactobacillus fermentum		0.43	0.24			
23105101	**Good identification**					**23105101**
					SPO	
Clostridium chauvoei		95.52	0.40	MNS 90	99	
Actinomyces odontolyticus		3.29	0.21	XYL 1	1	
Actinomyces meyeri		1.19	0.10		1	
23105102	**Good identification**					**23105102**
Actinomyces odontolyticus		97.29	0.52	FRU 92		
Actinomyces meyeri		1.78	0.18			
Lactobacillus fermentum		0.93	0.24			
23105120	**Low discrimination**					**23105120**
Actinomyces odontolyticus		75.97	0.42	FRU 92		
Actinomyces meyeri		23.12	0.29	NAG 14		
Lactobacillus fermentum		0.91	0.16			
23105122	**Good identification**					**23105122**
Actinomyces odontolyticus		96.63	0.38	FRU 92		
Lactobacillus fermentum		1.89	0.16	NAG 14		
Actinomyces meyeri		1.48	0.03			
23105200	**Very good identification**					**23105200**
Actinomyces odontolyticus		100.0	0.69	FRU 92		
23105202	**Very good identification**					**23105202**
Actinomyces odontolyticus		100.0	0.65	FRU 92		
23105220	**Good identification**					**23105220**
Actinomyces odontolyticus		100.0	0.55	FRU 92		
23105222	**Good identification**					**23105222**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces odontolyticus		100.0	0.52	FRU 92				
23105300	**Very good identification**							**23105300**
Actinomyces odontolyticus		99.81	0.60	FRU 92				
Actinomyces meyeri		0.19	0.10					
23105302	**Very good identification**							**23105302**
Actinomyces odontolyticus		100.0	0.56	FRU 92				
23105320	**Good identification**							**23105320**
Actinomyces odontolyticus		100.0	0.47	FRU 92				
23105322	**Good identification**							**23105322**
Actinomyces odontolyticus		100.0	0.43	FRU 92				
23106200	**Good identification**							**23106200**
Actinomyces odontolyticus		100.0	0.38	FRU 92				
23110000	**Good identification**							**23110000**
					COC	GRA		
Actinomyces odontolyticus		93.71	0.90		1	99		
Megasphaera elsdenii		5.49	0.66		99	1		
Actinomyces meyeri		0.50	0.48		1	99		
23110001	**Acceptable identification**							**23110001**
					COC	SPO	GRA	
Actinomyces odontolyticus		92.56	0.56	URE 1	1	1	99	
Megasphaera elsdenii		5.42	0.32		99	1	1	
Clostridium chauvoei		0.57	0.12		1	99	99	
23110002	**Very good identification**							**23110002**
					COC	GRA		
Actinomyces odontolyticus		96.65	0.87		1	99		
Actinomyces viscosus		2.26	0.52		1	99		
Megasphaera elsdenii		1.03	0.50		99	1		
23110003	**Low discrimination**							**23110003**
					COC	GRA		
Actinomyces odontolyticus		55.95	0.52	URE 1	1	99		
Actinomyces viscosus		43.20	0.44		1	99		
Megasphaera elsdenii		0.59	0.15		99	1		
23110010	**Acceptable identification**							**23110010**
					COC	GRA		
Actinomyces odontolyticus		93.78	0.56	MLZ 1	1	99		
Megasphaera elsdenii		5.49	0.32		99	1		
Actinomyces meyeri		0.50	0.13		1	99		
23110012	**Good identification**							**23110012**
					COC	GRA		
Actinomyces odontolyticus		96.71	0.52	MLZ 1	1	99		
Actinomyces viscosus		2.26	0.18		1	99		
Megasphaera elsdenii		1.03	0.15		99	1		
23110020	**Very good identification**							**23110020**
					COC	GRA		
Actinomyces odontolyticus		99.09	0.77	NAG 14	1	99		
Actinomyces meyeri		0.44	0.33		1	99		
Megasphaera elsdenii		0.36	0.32		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
23110021	**Good identification**							**23110021**
					SPO			
Actinomyces odontolyticus		98.75	0.43	NAG 14	1			
Clostridium chauvoei		1.25	0.04	URE 1	99			
23110022	**Very good identification**							**23110022**
					COC	GRA		
Actinomyces odontolyticus		99.75	0.73	NAG 14	1	99		
Actinomyces viscosus		0.15	0.18		1	99		
Megasphaera elsdenii		0.07	0.15		99	1		
23110023	**Good identification**							**23110023**
Actinomyces odontolyticus		95.43	0.39	NAG 14				
Actinomyces viscosus		4.57	0.09	URE 1				
23110030	**Good identification**							**23110030**
Actinomyces odontolyticus		100.0	0.43	MLZ 1				
23110032	**Good identification**							**23110032**
Actinomyces odontolyticus		100.0	0.39	MLZ 1				
23110040	**Low discrimination**							**23110040**
					COC	SPO	GRA	
Megasphaera elsdenii		78.16	0.63	NIT 1	99	1	1	
Actinomyces odontolyticus		20.21	0.56		1	1	99	
Clostridium botulinum biovar B		1.03	0.24		1	99	99	
23110042	**Low discrimination**							**23110042**
					COC	GRA		
Actinomyces odontolyticus		57.19	0.52	SOR 1	1	99		
Megasphaera elsdenii		40.09	0.47		99	1		
Propionibacterium propionicum		1.39	0.32		1	99		
23110060	**Low discrimination**							**23110060**
					COC	SPO	GRA	
Actinomyces odontolyticus		71.39	0.43	NAG 14	1	1	99	
Megasphaera elsdenii		17.13	0.29	SOR 1	99	1	1	
Clostridium botulinum biovar B		10.96	0.18		1	99	99	
23110062	**Good identification**							**23110062**
					COC	GRA		
Actinomyces odontolyticus		95.83	0.39	NAG 14	1	99		
Megasphaera elsdenii		4.17	0.12	SOR 1	99	1		
23110100	**Good identification**							**23110100**
Actinomyces odontolyticus		90.69	0.82					
Actinomyces meyeri		6.14	0.58					
Propionibacterium propionicum		2.85	0.63					
23110101	**Low discrimination**							**23110101**
					SPO			
Clostridium chauvoei		57.35	0.47	MNS 90	99			
Actinomyces odontolyticus		29.61	0.47	GAL 99	1			
Propionibacterium propionicum		10.23	0.47		1			
23110102	**Low discrimination**							**23110102**
Actinomyces odontolyticus		82.44	0.78					
Actinomyces viscosus		10.86	0.56					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		6.34	0.66				
23110103	**Low discrimination**						**23110103**
Actinomyces viscosus		64.40	0.48	MNS 96			
Actinomyces odontolyticus		14.81	0.44	RAF 94			
Propionibacterium propionicum		12.53	0.50				
23110110	**Acceptable identification**						**23110110**
Actinomyces odontolyticus		90.99	0.47	MLZ 1			
Actinomyces meyeri		6.16	0.24				
Propionibacterium propionicum		2.86	0.29				
23110112	**Low discrimination**						**23110112**
Actinomyces odontolyticus		82.74	0.44	MLZ 1			
Actinomyces viscosus		10.90	0.22				
Propionibacterium propionicum		6.36	0.32				
23110120	**Good identification**						**23110120**
Actinomyces odontolyticus		94.11	0.68	NAG 14			
Actinomyces meyeri		5.34	0.43				
Clostridium perfringens		0.24	0.19				
23110121	**Low discrimination**						**23110121**
					SPO		
Clostridium chauvoei		78.62	0.38	MNS 90	99		
Actinomyces odontolyticus		19.83	0.34	GAL 99	1		
Actinomyces meyeri		1.12	0.09		1		
23110122	**Very good identification**						**23110122**
Actinomyces odontolyticus		98.12	0.64	NAG 14			
Actinomyces viscosus		0.80	0.22				
Propionibacterium propionicum		0.47	0.32				
23110140	**Low discrimination**						**23110140**
					COC	GRA	
Propionibacterium propionicum		72.30	0.63	MAN 99	1	99	
Actinomyces odontolyticus		23.25	0.47		1	99	
Megasphaera elsdenii		2.88	0.29		99	1	
23110141	**Acceptable identification**						**23110141**
					SPO		
Propionibacterium propionicum		92.10	0.47	MAN 99	1		
Clostridium chauvoei		5.21	0.12	URE 10	99		
Actinomyces odontolyticus		2.69	0.13		1		
23110142	**Low discrimination**						**23110142**
Propionibacterium propionicum		86.83	0.66	MAN 99			
Actinomyces odontolyticus		11.41	0.44				
Actinomyces viscosus		1.50	0.22				
23110143	**Acceptable identification**						**23110143**
Propionibacterium propionicum		93.41	0.50	MAN 99			
Actinomyces viscosus		4.85	0.14	URE 10			
Actinomyces odontolyticus		1.12	0.10				
23110200	**Excellent identification**						**23110200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Actinomyces odontolyticus		99.97	0.95		1	99	
Megasphaera elsdenii		0.03	0.32		99	1	
Actinomyces meyeri		0.00	0.13		1	99	
23110201	**Very good identification**						**23110201**
Actinomyces odontolyticus		100.0	0.61	URE 1			
23110202	**Excellent identification**						**23110202**
					COC	GRA	
Actinomyces odontolyticus		99.98	0.91		1	99	
Actinomyces viscosus		0.01	0.18		1	99	
Megasphaera elsdenii		0.01	0.15		99	1	
23110203	**Very good identification**						**23110203**
Actinomyces odontolyticus		99.60	0.57	URE 1			
Actinomyces viscosus		0.40	0.09				
23110210	**Very good identification**						**23110210**
Actinomyces odontolyticus		100.0	0.61	MLZ 1			
23110212	**Very good identification**						**23110212**
Actinomyces odontolyticus		100.0	0.57	MLZ 1			
23110220	**Very good identification**						**23110220**
Actinomyces odontolyticus		100.0	0.82	NAG 14			
23110221	**Good identification**						**23110221**
Actinomyces odontolyticus		100.0	0.47	NAG 14			
23110222	**Very good identification**						**23110222**
Actinomyces odontolyticus		100.0	0.78	NAG 14			
23110223	**Good identification**						**23110223**
Actinomyces odontolyticus		100.0	0.44	NAG 14			
23110230	**Good identification**						**23110230**
Actinomyces odontolyticus		100.0	0.47	MLZ 1			
23110232	**Good identification**						**23110232**
Actinomyces odontolyticus		100.0	0.44	MLZ 1			
23110240	**Very good identification**						**23110240**
					COC	GRA	
Actinomyces odontolyticus		98.03	0.61	SOR 1	1	99	
Megasphaera elsdenii		1.97	0.29		99	1	
23110242	**Very good identification**						**23110242**
					COC	GRA	
Actinomyces odontolyticus		99.64	0.57	SOR 1	1	99	
Megasphaera elsdenii		0.36	0.12		99	1	
23110260	**Good identification**						**23110260**
Actinomyces odontolyticus		100.0	0.47	NAG 14			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23110262	**Good identification**					**23110262**
Actinomyces odontolyticus		100.0	0.44	NAG 14		
23110300	**Very good identification**					**23110300**
Actinomyces odontolyticus		99.95	0.87			
Actinomyces meyeri		0.04	0.24			
Propionibacterium propionicum		0.02	0.29			
23110301	**Good identification**					**23110301**
					SPO	
Actinomyces odontolyticus		98.83	0.52	URE 1	1	
Clostridium chauvoei		1.00	0.12		99	
Propionibacterium propionicum		0.18	0.12		1	
23110302	**Very good identification**					**23110302**
Actinomyces odontolyticus		99.89	0.83			
Actinomyces viscosus		0.07	0.22			
Propionibacterium propionicum		0.04	0.32			
23110303	**Good identification**					**23110303**
Actinomyces odontolyticus		97.10	0.49	URE 1		
Actinomyces viscosus		2.20	0.14			
Propionibacterium propionicum		0.43	0.15			
23110310	**Good identification**					**23110310**
Actinomyces odontolyticus		100.0	0.52	MLZ 1		
23110312	**Good identification**					**23110312**
Actinomyces odontolyticus		100.0	0.49	MLZ 1		
23110320	**Very good identification**					**23110320**
Actinomyces odontolyticus		99.97	0.73	NAG 14		
Actinomyces meyeri		0.03	0.09			
23110321	**Good identification**					**23110321**
					SPO	
Actinomyces odontolyticus		97.98	0.39	NAG 14	1	
Clostridium chauvoei		2.02	0.04	URE 1	99	
23110322	**Very good identification**					**23110322**
Actinomyces odontolyticus		100.0	0.69	NAG 14		
23110323	**Good identification**					**23110323**
Actinomyces odontolyticus		100.0	0.35	NAG 14		
23110330	**Good identification**					**23110330**
Actinomyces odontolyticus		100.0	0.39	MLZ 1		
23110332	**Good identification**					**23110332**
Actinomyces odontolyticus		100.0	0.35	MLZ 1		
23110340	**Good identification**					**23110340**
Actinomyces odontolyticus		98.41	0.52	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		1.59	0.29				
23110342	**Good identification**						**23110342**
Actinomyces odontolyticus		96.19	0.49	SOR 1			
Propionibacterium propionicum		3.81	0.32				
23110360	**Good identification**						**23110360**
Actinomyces odontolyticus		100.0	0.39	NAG 14			
23110362	**Good identification**						**23110362**
Actinomyces odontolyticus		100.0	0.35	NAG 14			
23110400	**Low discrimination**						**23110400**
					COC	GRA	
Actinomyces odontolyticus		83.05	0.56	ARA 1	1	99	
Actinomyces meyeri		10.98	0.37		1	99	
Megasphaera elsdenii		4.87	0.32		99	1	
23110402	**Good identification**						**23110402**
					COC	GRA	
Actinomyces odontolyticus		95.28	0.52	ARA 1	1	99	
Actinomyces viscosus		2.23	0.18		1	99	
Megasphaera elsdenii		1.01	0.15		99	1	
23110420	**Low discrimination**						**23110420**
Actinomyces odontolyticus		89.13	0.43	ARA 1			
Actinomyces meyeri		9.87	0.23	NAG 14			
Lactobacillus fermentum		1.00	0.16				
23110422	**Good identification**						**23110422**
Actinomyces odontolyticus		98.21	0.39	ARA 1			
Lactobacillus fermentum		1.79	0.16	NAG 14			
23110500	**Low discrimination**						**23110500**
Actinomyces meyeri		61.25	0.48	NIT 1			
Actinomyces odontolyticus		36.57	0.47				
Propionibacterium propionicum		1.15	0.29				
23110502	**Low discrimination**						**23110502**
Actinomyces odontolyticus		74.68	0.44	ARA 1			
Actinomyces viscosus		9.84	0.22				
Actinomyces meyeri		6.31	0.22				
23110600	**Very good identification**						**23110600**
Actinomyces odontolyticus		99.93	0.61	ARA 1			
Actinomyces meyeri		0.07	0.03				
23110602	**Very good identification**						**23110602**
Actinomyces odontolyticus		100.0	0.57	ARA 1			
23110620	**Good identification**						**23110620**
Actinomyces odontolyticus		100.0	0.47	ARA 1			
23110622	**Good identification**						**23110622**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.44	ARA 1		
23110700	**Good identification**					**23110700**
Actinomyces odontolyticus		99.14	0.52	ARA 1		
Actinomyces meyeri		0.86	0.13			
23110702	**Good identification**					**23110702**
Actinomyces odontolyticus		100.0	0.49	ARA 1		
23110720	**Good identification**					**23110720**
Actinomyces odontolyticus		100.0	0.39	ARA 1		
23110722	**Good identification**					**23110722**
Actinomyces odontolyticus		100.0	0.35	ARA 1		
23111000	**Very good identification**					**23111000**
Actinomyces odontolyticus		97.66	0.87			
Actinomyces meyeri		1.17	0.51			
Lactobacillus fermentum		0.80	0.59			
23111001	**Low discrimination**					**23111001**
					SPO	
Actinomyces odontolyticus		51.11	0.53	URE 1	1	
Clostridium chauvoei		46.89	0.47		99	
Actinomyces naeslundii		0.66	0.38		1	
23111002	**Very good identification**					**23111002**
Actinomyces odontolyticus		96.50	0.84			
Actinomyces viscosus		2.08	0.48			
Lactobacillus fermentum		1.29	0.59			
23111003	**Low discrimination**					**23111003**
					SPO	
Actinomyces odontolyticus		49.69	0.49	URE 1	1	
Actinomyces viscosus		35.27	0.40		1	
Clostridium chauvoei		13.12	0.34		99	
23111010	**Good identification**					**23111010**
Actinomyces odontolyticus		98.02	0.53	MLZ 1		
Actinomyces meyeri		1.18	0.16			
Lactobacillus fermentum		0.80	0.24			
23111012	**Good identification**					**23111012**
Actinomyces odontolyticus		96.63	0.49	MLZ 1		
Actinomyces viscosus		2.08	0.14			
Lactobacillus fermentum		1.29	0.24			
23111020	**Very good identification**					**23111020**
Actinomyces odontolyticus		95.71	0.74	NAG 14		
Lactobacillus fermentum		1.61	0.50			
Clostridium perfringens		1.54	0.38			
23111021	**Low discrimination**					**23111021**
					SPO	
Clostridium chauvoei		62.59	0.38	MNS 90	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		33.32	0.40	LAC 99	1		
Actinomyces naeslundii		2.66	0.38		1		
23111022	**Very good identification**						**23111022**
Actinomyces odontolyticus		95.91	0.70	NAG 14			
Lactobacillus fermentum		2.62	0.50				
Clostridium perfringens		1.19	0.32				
23111023	**Low discrimination**						**23111023**
					SPO		
Actinomyces odontolyticus		57.39	0.36	NAG 14	1		
Clostridium chauvoei		31.04	0.25	URE 1	99		
Actinomyces naeslundii		7.47	0.38		1		
23111030	**Good identification**						**23111030**
Actinomyces odontolyticus		95.88	0.40	MLZ 1			
Lactobacillus fermentum		1.61	0.16	NAG 14			
Clostridium perfringens		1.55	0.04				
23111032	**Good identification**						**23111032**
Actinomyces odontolyticus		97.34	0.36	MLZ 1			
Lactobacillus fermentum		2.66	0.16	NAG 14			
23111040	**Acceptable identification**						**23111040**
					COC	GRA	
Actinomyces odontolyticus		91.44	0.53	SOR 1	1	99	
Megasphaera elsdenii		5.36	0.29		99	1	
Propionibacterium propionicum		1.36	0.29		1	99	
23111042	**Acceptable identification**						**23111042**
Actinomyces odontolyticus		92.43	0.49	SOR 1			
Propionibacterium propionicum		3.37	0.32				
Actinomyces viscosus		1.99	0.14				
23111060	**Low discrimination**						**23111060**
Actinomyces odontolyticus		74.53	0.40	NAG 14			
Clostridium perfringens		22.66	0.26	SOR 1			
Lactobacillus fermentum		1.25	0.16				
23111062	**Low discrimination**						**23111062**
Actinomyces odontolyticus		79.31	0.36	NAG 14			
Clostridium perfringens		18.52	0.20	SOR 1			
Lactobacillus fermentum		2.17	0.16				
23111100	**Low discrimination**						**23111100**
Actinomyces odontolyticus		78.39	0.79				
Actinomyces meyeri		11.94	0.61				
Propionibacterium propionicum		3.69	0.63				
23111101	**Very good identification**						**23111101**
					SPO		
Clostridium chauvoei		99.36	0.81	MNS 90	99		
Actinomyces odontolyticus		0.35	0.44		1		
Propionibacterium propionicum		0.18	0.47		1		
23111102	**Low discrimination**						**23111102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		75.43	0.75			
Actinomyces viscosus		9.14	0.52			
Propionibacterium propionicum		8.70	0.66			
23111103	**Good identification**					**23111103**
					SPO	
Clostridium chauvoei		92.70	0.68	MNS 90	99	
Actinomyces viscosus		4.47	0.44		1	
Propionibacterium propionicum		1.42	0.50		1	
23111110	**Low discrimination**					**23111110**
Actinomyces odontolyticus		78.68	0.44	MLZ 1		
Actinomyces meyeri		11.98	0.27			
Propionibacterium propionicum		3.71	0.29			
23111111	**Good identification**					**23111111**
					SPO	
Clostridium chauvoei		99.27	0.47	MNS 90	99	
Actinomyces odontolyticus		0.35	0.10	MLZ 1	1	
Actinomyces naeslundii		0.21	0.24		1	
23111112	**Low discrimination**					**23111112**
Actinomyces odontolyticus		75.34	0.41	MLZ 1		
Actinomyces viscosus		9.12	0.18			
Propionibacterium propionicum		8.69	0.32			
23111120	**Low discrimination**					**23111120**
Actinomyces odontolyticus		57.81	0.65	NAG 14		
Clostridium perfringens		21.65	0.53			
Actinomyces meyeri		7.37	0.46			
23111121	**Very good identification**					**23111121**
					SPO	
Clostridium chauvoei		99.59	0.73	MNS 90	99	
Actinomyces odontolyticus		0.17	0.31		1	
Actinomyces naeslundii		0.12	0.46		1	
23111122	**Low discrimination**					**23111122**
Actinomyces odontolyticus		63.95	0.61	NAG 14		
Clostridium perfringens		18.39	0.47			
Lactobacillus fermentum		9.04	0.54			
23111123	**Very good identification**					**23111123**
					SPO	
Clostridium chauvoei		98.26	0.60	MNS 90	99	
Actinomyces naeslundii		0.68	0.46		1	
Actinomyces odontolyticus		0.58	0.27		1	
23111131	**Good identification**					**23111131**
					SPO	
Clostridium chauvoei		99.37	0.38	MNS 90	99	
Actinomyces naeslundii		0.63	0.24	MLZ 1	1	
23111140	**Low discrimination**					**23111140**
Propionibacterium propionicum		78.56	0.63	MAN 99		
Actinomyces odontolyticus		16.84	0.44			

Code / Taxon		ID score	T index	tests against		Differentiation tests	Code
Actinomyces meyeri		2.56	0.27				
23111141	**Low discrimination**						**23111141**
						SPO	
Clostridium chauvoei		84.10	0.47	MNS	90	99	
Propionibacterium propionicum		15.01	0.47	SOR	1	1	
Actinomyces naeslundii		0.60	0.33			1	
23111142	**Good identification**						**23111142**
Propionibacterium propionicum		90.05	0.66	MAN	99		
Actinomyces odontolyticus		7.89	0.41				
Actinomyces viscosus		0.96	0.18				
23111143	**Low discrimination**						**23111143**
						SPO	
Propionibacterium propionicum		57.95	0.50	MAN	99	1	
Clostridium chauvoei		38.21	0.34	URE	10	99	
Actinomyces viscosus		1.84	0.10			1	
23111160	**Low discrimination**						**23111160**
Clostridium perfringens		79.98	0.40	MNS	99		
Actinomyces odontolyticus		11.33	0.31				
Propionibacterium propionicum		3.28	0.29				
23111161	**Good identification**						**23111161**
						SPO	
Clostridium chauvoei		96.27	0.38	MNS	90	99	
Actinomyces naeslundii		2.05	0.33	SOR	1	1	
Clostridium perfringens		1.15	0.06			84	
23111200	**Excellent identification**						**23111200**
Actinomyces odontolyticus		99.99	0.92				
Actinomyces meyeri		0.01	0.16				
Lactobacillus fermentum		0.00	0.24				
23111201	**Very good identification**						**23111201**
						SPO	
Actinomyces odontolyticus		99.53	0.58	URE	1	1	
Clostridium chauvoei		0.48	0.12			99	
23111202	**Very good identification**						**23111202**
Actinomyces odontolyticus		99.98	0.88				
Actinomyces viscosus		0.01	0.14				
Lactobacillus fermentum		0.01	0.24				
23111203	**Good identification**						**23111203**
						SPO	
Actinomyces odontolyticus		99.50	0.54	URE	1	1	
Actinomyces viscosus		0.37	0.06			1	
Clostridium chauvoei		0.14	-.01			99	
23111210	**Very good identification**						**23111210**
Actinomyces odontolyticus		100.0	0.58	MLZ	1		
23111212	**Good identification**						**23111212**
Actinomyces odontolyticus		100.0	0.54	MLZ	1		
23111220	**Very good identification**						**23111220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.98	0.79	NAG 14		
Lactobacillus fermentum		0.01	0.16			
Clostridium perfringens		0.01	0.04			
23111221	**Good identification**					**23111221**
					SPO	
Actinomyces odontolyticus		99.03	0.44	NAG 14	1	
Clostridium chauvoei		0.97	0.04	URE 1	99	
23111222	**Very good identification**					**23111222**
Actinomyces odontolyticus		99.99	0.75	NAG 14		
Lactobacillus fermentum		0.01	0.16			
23111223	**Good identification**					**23111223**
Actinomyces odontolyticus		100.0	0.41	NAG 14		
23111230	**Good identification**					**23111230**
Actinomyces odontolyticus		100.0	0.44	MLZ 1		
23111232	**Good identification**					**23111232**
Actinomyces odontolyticus		100.0	0.41	MLZ 1		
23111240	**Very good identification**					**23111240**
Actinomyces odontolyticus		100.0	0.58	SOR 1		
23111242	**Good identification**					**23111242**
Actinomyces odontolyticus		100.0	0.54	SOR 1		
23111260	**Good identification**					**23111260**
Actinomyces odontolyticus		100.0	0.44	NAG 14		
23111262	**Good identification**					**23111262**
Actinomyces odontolyticus		100.0	0.41	NAG 14		
23111300	**Very good identification**					**23111300**
Actinomyces odontolyticus		99.86	0.84			
Actinomyces meyeri		0.08	0.27			
Propionibacterium propionicum		0.02	0.29			
23111301	**Low discrimination**					**23111301**
					SPO	
Clostridium chauvoei		59.88	0.47	MNS 90	99	
Actinomyces odontolyticus		40.01	0.49	RHA 1	1	
Propionibacterium propionicum		0.11	0.12		1	
23111302	**Very good identification**					**23111302**
Actinomyces odontolyticus		99.84	0.80			
Actinomyces viscosus		0.06	0.18			
Propionibacterium propionicum		0.06	0.32			
23111303	**Low discrimination**					**23111303**
					SPO	
Actinomyces odontolyticus		68.57	0.46	URE 1	1	
Clostridium chauvoei		29.55	0.34		99	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		1.43	0.10		1	
23111310	**Good identification**					**23111310**
Actinomyces odontolyticus		100.0	0.49	MLZ 1		
23111312	**Good identification**					**23111312**
Actinomyces odontolyticus		100.0	0.46	MLZ 1		
23111320	**Very good identification**					**23111320**
Actinomyces odontolyticus		99.63	0.70	NAG 14		
Clostridium perfringens		0.19	0.19			
Actinomyces meyeri		0.07	0.12			
23111321	**Low discrimination**					**23111321**
					SPO	
Clostridium chauvoei		75.40	0.38	MNS 90	99	
Actinomyces odontolyticus		24.61	0.36	RHA 1	1	
23111322	**Very good identification**					**23111322**
Actinomyces odontolyticus		99.74	0.66	NAG 14		
Clostridium perfringens		0.15	0.13			
Lactobacillus fermentum		0.07	0.20			
23111330	**Good identification**					**23111330**
Actinomyces odontolyticus		100.0	0.36	MLZ 1		
23111340	**Good identification**					**23111340**
Actinomyces odontolyticus		97.63	0.49	SOR 1		
Propionibacterium propionicum		2.37	0.29			
23111342	**Acceptable identification**					**23111342**
Actinomyces odontolyticus		94.39	0.46	SOR 1		
Propionibacterium propionicum		5.61	0.32			
23111360	**Good identification**					**23111360**
Actinomyces odontolyticus		96.46	0.36	NAG 14		
Clostridium perfringens		3.54	0.06	SOR 1		
23111400	**Low discrimination**					**23111400**
Actinomyces odontolyticus		47.29	0.53	ARA 1		
Lactobacillus fermentum		38.34	0.59			
Actinomyces meyeri		14.07	0.40			
23111402	**Low discrimination**					**23111402**
Lactobacillus fermentum		56.06	0.59	NIT 1		
Actinomyces odontolyticus		42.39	0.49			
Actinomyces viscosus		0.91	0.14			
23111420	**Low discrimination**					**23111420**
Lactobacillus fermentum		56.74	0.50	NIT 1		
Actinomyces odontolyticus		34.19	0.40			
Actinomyces meyeri		8.52	0.26			
23111422	**Low discrimination**					**23111422**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		73.03	0.50	NIT 1			
Actinomyces odontolyticus		26.97	0.36				
23111500	**Low discrimination**						**23111500**
Lactobacillus fermentum		46.36	0.62	NIT 1			
Actinomyces meyeri		41.72	0.51				
Actinomyces odontolyticus		11.07	0.44				
23111501	**Good identification**						**23111501**
					SPO		
Clostridium chauvoei		96.81	0.47	MNS 90	99		
Lactobacillus fermentum		1.41	0.28	ARA 1	1		
Actinomyces meyeri		1.27	0.16		1		
23111502	**Low discrimination**						**23111502**
Lactobacillus fermentum		82.43	0.62	NIT 1			
Actinomyces odontolyticus		12.06	0.41				
Actinomyces meyeri		2.29	0.25				
23111520	**Low discrimination**						**23111520**
Lactobacillus fermentum		63.93	0.54	NIT 1			
Actinomyces meyeri		23.53	0.36				
Actinomyces odontolyticus		7.46	0.31				
23111521	**Good identification**						**23111521**
					SPO		
Clostridium chauvoei		98.05	0.38	MNS 90	99		
Lactobacillus fermentum		1.43	0.20	ARA 1	1		
Actinomyces meyeri		0.53	0.02		1		
23111522	**Low discrimination**						**23111522**
					GRA	CAT	
Lactobacillus fermentum		83.41	0.54	NIT 1	99	1	
Capnocytophaga ochracea		7.01	0.20		1	90	
Actinomyces odontolyticus		5.96	0.27		99	10	
23111600	**Very good identification**						**23111600**
Actinomyces odontolyticus		99.43	0.58	ARA 1			
Lactobacillus fermentum		0.42	0.24				
Actinomyces meyeri		0.15	0.06				
23111602	**Good identification**						**23111602**
Actinomyces odontolyticus		99.32	0.54	ARA 1			
Lactobacillus fermentum		0.68	0.24				
23111620	**Good identification**						**23111620**
Actinomyces odontolyticus		99.14	0.44	ARA 1			
Lactobacillus fermentum		0.86	0.16	NAG 14			
23111622	**Good identification**						**23111622**
Actinomyces odontolyticus		98.61	0.41	ARA 1			
Lactobacillus fermentum		1.39	0.16	NAG 14			
23111700	**Good identification**						**23111700**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		96.02	0.49	ARA 1			
Lactobacillus fermentum		2.09	0.28				
Actinomyces meyeri		1.88	0.16				
23111702	**Good identification**						**23111702**
Actinomyces odontolyticus		96.57	0.46	ARA 1			
Lactobacillus fermentum		3.43	0.28				
23111720	**Acceptable identification**						**23111720**
Actinomyces odontolyticus		94.25	0.36	ARA 1			
Lactobacillus fermentum		4.21	0.20	NAG 14			
Actinomyces meyeri		1.55	0.02				
23112000	**Low discrimination**						**23112000**
					COC	GRA	
Megasphaera elsdenii		74.44	0.66	NIT 1	99	1	
Actinomyces odontolyticus		12.83	0.56		1	99	
Propionibacterium propionicum		12.61	0.63		1	99	
23112001	**Low discrimination**						**23112001**
					COC	GRA	
Propionibacterium propionicum		61.37	0.47	LAC 99	1	99	
Megasphaera elsdenii		32.95	0.32	URE 10	99	1	
Actinomyces odontolyticus		5.68	0.22		1	99	
23112002	**Low discrimination**						**23112002**
					COC	GRA	
Propionibacterium propionicum		53.67	0.66	LAC 99	1	99	
Megasphaera elsdenii		23.48	0.50		99	1	
Actinomyces odontolyticus		22.33	0.52		1	99	
23112003	**Acceptable identification**						**23112003**
					COC	GRA	
Propionibacterium propionicum		90.35	0.50	LAC 99	1	99	
Megasphaera elsdenii		3.59	0.15	URE 10	99	1	
Actinomyces odontolyticus		3.42	0.18		1	99	
23112020	**Low discrimination**						**23112020**
					COC	GRA	
Actinomyces odontolyticus		69.47	0.43	MAN 1	1	99	
Megasphaera elsdenii		25.00	0.32	NAG 14	99	1	
Propionibacterium propionicum		4.23	0.29		1	99	
23112022	**Low discrimination**						**23112022**
					COC	GRA	
Actinomyces odontolyticus		82.35	0.39	MAN 1	1	99	
Propionibacterium propionicum		12.28	0.32	NAG 14	1	99	
Megasphaera elsdenii		5.37	0.15		99	1	
23112040	**Low discrimination**						**23112040**
					COC	GRA	
Megasphaera elsdenii		79.58	0.63	NIT 1	99	1	
Propionibacterium propionicum		20.21	0.63		1	99	
Actinomyces odontolyticus		0.21	0.22		1	99	
23112041	**Low discrimination**						**23112041**
					COC	GRA	
Propionibacterium propionicum		73.64	0.47	LAC 99	1	99	
Megasphaera elsdenii		26.36	0.29	URE 10	99	1	
23112042	**Low discrimination**						**23112042**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Propionibacterium propionicum		77.17	0.66	LAC 99	1	99	
Megasphaera elsdenii		22.51	0.47		99	1	
Actinomyces odontolyticus		0.32	0.18		1	99	
23112043	**Good identification**						**23112043**
					COC	GRA	
Propionibacterium propionicum		97.42	0.50	LAC 99	1	99	
Megasphaera elsdenii		2.58	0.12	URE 10	99	1	
23112100	**Excellent identification**						**23112100**
					COC	GRA	
Propionibacterium propionicum		99.59	0.97		1	99	
Actinomyces odontolyticus		0.32	0.47		1	99	
Megasphaera elsdenii		0.06	0.32		99	1	
23112101	**Very good identification**						**23112101**
					SPO		
Propionibacterium propionicum		99.90	0.81	URE 10	1		
Clostridium chauvoei		0.06	0.12		99		
Actinomyces odontolyticus		0.03	0.13		1		
23112102	**Excellent identification**						**23112102**
Propionibacterium propionicum		99.85	1.00				
Actinomyces odontolyticus		0.13	0.44				
Actinomyces viscosus		0.02	0.22				
23112103	**Very good identification**						**23112103**
Propionibacterium propionicum		99.92	0.84	URE 10			
Actinomyces viscosus		0.05	0.14				
Actinomyces naeslundii		0.01	0.27				
23112110	**Very good identification**						**23112110**
Propionibacterium propionicum		99.68	0.63	MLZ 1			
Actinomyces odontolyticus		0.32	0.13				
23112111	**Good identification**						**23112111**
Propionibacterium propionicum		100.0	0.47	MLZ 1			
23112112	**Very good identification**						**23112112**
Propionibacterium propionicum		99.87	0.66	MLZ 1			
Actinomyces odontolyticus		0.13	0.10				
23112113	**Good identification**						**23112113**
Propionibacterium propionicum		100.0	0.50	MLZ 1			
23112120	**Good identification**						**23112120**
Propionibacterium propionicum		94.64	0.63	NAG 1			
Actinomyces odontolyticus		4.95	0.34				
Actinomyces meyeri		0.28	0.09				
23112121	**Good identification**						**23112121**
					SPO		
Propionibacterium propionicum		96.42	0.47	NAG 1	1		
Clostridium chauvoei		1.82	0.04	URE 10	99		
Actinomyces naeslundii		1.76	0.27		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
23112122	**Very good identification**						**23112122**
Propionibacterium propionicum		97.82	0.66	NAG 1			
Actinomyces odontolyticus		2.09	0.30				
Actinomyces naeslundii		0.09	0.24				
23112123	**Good identification**						**23112123**
Propionibacterium propionicum		98.80	0.50	NAG 1			
Actinomyces naeslundii		1.20	0.27	URE 10			
23112140	**Excellent identification**						**23112140**
					COC	GRA	
Propionibacterium propionicum		99.96	0.97		1	99	
Megasphaera elsdenii		0.04	0.29		99	1	
Actinomyces odontolyticus		0.00	0.13		1	99	
23112141	**Very good identification**						**23112141**
Propionibacterium propionicum		100.0	0.81	URE 10			
23112142	**Excellent identification**						**23112142**
					COC	GRA	
Propionibacterium propionicum		100.0	1.00		1	99	
Megasphaera elsdenii		0.00	0.12		99	1	
Actinomyces odontolyticus		0.00	0.10		1	99	
23112143	**Very good identification**						**23112143**
Propionibacterium propionicum		100.0	0.84	URE 10			
23112150	**Very good identification**						**23112150**
Propionibacterium propionicum		100.0	0.63	MLZ 1			
23112151	**Good identification**						**23112151**
Propionibacterium propionicum		100.0	0.47	MLZ 1			
23112152	**Very good identification**						**23112152**
Propionibacterium propionicum		100.0	0.66	MLZ 1			
23112153	**Good identification**						**23112153**
Propionibacterium propionicum		100.0	0.50	MLZ 1			
23112160	**Very good identification**						**23112160**
Propionibacterium propionicum		100.0	0.63	NAG 1			
23112161	**Good identification**						**23112161**
Propionibacterium propionicum		100.0	0.47	NAG 1			
23112162	**Very good identification**						**23112162**
Propionibacterium propionicum		100.0	0.66	NAG 1			
23112163	**Good identification**						**23112163**
Propionibacterium propionicum		100.0	0.50	NAG 1			
23112200	**Very good identification**						**23112200**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		96.59	0.61	MAN 1	1	99	
Megasphaera elsdenii		2.92	0.32		99	1	
Propionibacterium propionicum		0.49	0.29		1	99	
23112202	**Very good identification**						**23112202**
					COC	GRA	
Actinomyces odontolyticus		98.23	0.57	MAN 1	1	99	
Propionibacterium propionicum		1.23	0.32		1	99	
Megasphaera elsdenii		0.54	0.15		99	1	
23112220	**Good identification**						**23112220**
Actinomyces odontolyticus		100.0	0.47	MAN 1			
23112222	**Good identification**						**23112222**
Actinomyces odontolyticus		100.0	0.44	MAN 1			
23112300	**Low discrimination**						**23112300**
Propionibacterium propionicum		61.57	0.63	RHA 1			
Actinomyces odontolyticus		38.43	0.52				
23112301	**Acceptable identification**						**23112301**
Propionibacterium propionicum		94.63	0.47	RHA 1			
Actinomyces odontolyticus		5.37	0.18	URE 10			
23112302	**Low discrimination**						**23112302**
Propionibacterium propionicum		79.68	0.66	RHA 1			
Actinomyces odontolyticus		20.32	0.49				
23112303	**Good identification**						**23112303**
Propionibacterium propionicum		97.73	0.50	RHA 1			
Actinomyces odontolyticus		2.27	0.15	URE 10			
23112320	**Acceptable identification**						**23112320**
Actinomyces odontolyticus		90.96	0.39	MAN 1			
Propionibacterium propionicum		9.04	0.29	NAG 14			
23112322	**Low discrimination**						**23112322**
Actinomyces odontolyticus		80.43	0.35	MAN 1			
Propionibacterium propionicum		19.57	0.32	NAG 14			
23112340	**Very good identification**						**23112340**
Propionibacterium propionicum		99.37	0.63	RHA 1			
Actinomyces odontolyticus		0.63	0.18				
23112341	**Good identification**						**23112341**
Propionibacterium propionicum		100.0	0.47	RHA 1			
23112342	**Very good identification**						**23112342**
Propionibacterium propionicum		99.74	0.66	RHA 1			
Actinomyces odontolyticus		0.26	0.15				
23112343	**Good identification**						**23112343**
Propionibacterium propionicum		100.0	0.50	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23112500	**Very good identification**						**23112500**
Propionibacterium propionicum		99.14	0.63	ARA 1			
Actinomyces meyeri		0.54	0.13				
Actinomyces odontolyticus		0.32	0.13				
23112501	**Good identification**						**23112501**
Propionibacterium propionicum		100.0	0.47	ARA 1			
23112502	**Very good identification**						**23112502**
Propionibacterium propionicum		99.87	0.66	ARA 1			
Actinomyces odontolyticus		0.13	0.10				
23112503	**Good identification**						**23112503**
Propionibacterium propionicum		100.0	0.50	ARA 1			
23112540	**Very good identification**						**23112540**
Propionibacterium propionicum		100.0	0.63	ARA 1			
23112541	**Good identification**						**23112541**
Propionibacterium propionicum		100.0	0.47	ARA 1			
23112542	**Very good identification**						**23112542**
Propionibacterium propionicum		100.0	0.66	ARA 1			
23112543	**Good identification**						**23112543**
Propionibacterium propionicum		100.0	0.50	ARA 1			
23113000	**Low discrimination**						**23113000**
					COC	GRA	
Propionibacterium propionicum		56.62	0.63	LAC 99	1	99	
Actinomyces odontolyticus		38.43	0.53		1	99	
Megasphaera elsdenii		3.38	0.32		99	1	
23113001	**Low discrimination**						**23113001**
Propionibacterium propionicum		78.59	0.47	LAC 99			
Actinomyces naeslundii		12.11	0.43	URE 10			
Actinomyces odontolyticus		4.85	0.19				
23113002	**Low discrimination**						**23113002**
Propionibacterium propionicum		76.98	0.66	LAC 99			
Actinomyces odontolyticus		21.35	0.49				
Actinomyces naeslundii		0.59	0.40				
23113003	**Low discrimination**						**23113003**
Propionibacterium propionicum		86.77	0.50	LAC 99			
Actinomyces naeslundii		8.91	0.43	URE 10			
Actinomyces odontolyticus		2.19	0.15				
23113020	**Low discrimination**						**23113020**
Actinomyces odontolyticus		79.02	0.40	MAN 1			
Actinomyces naeslundii		8.16	0.40	NAG 14			
Propionibacterium propionicum		7.22	0.29				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
23113021	**Low discrimination**						**23113021**
					SPO		
Actinomyces naeslundii		79.80	0.43	MNS 88	1		
Clostridium chauvoei		9.77	0.04	RAF 92	99		
Propionibacterium propionicum		5.23	0.12		1		
23113022	**Low discrimination**						**23113022**
Actinomyces odontolyticus		70.44	0.36	MAN 1			
Propionibacterium propionicum		15.76	0.32	NAG 14			
Actinomyces naeslundii		11.87	0.40				
23113023	**Acceptable identification**						**23113023**
Actinomyces naeslundii		91.05	0.43	MNS 88			
Propionibacterium propionicum		8.95	0.15	RAF 92			
23113040	**Very good identification**						**23113040**
					COC	GRA	
Propionibacterium propionicum		95.36	0.63	LAC 99	1	99	
Megasphaera elsdenii		3.79	0.29		99	1	
Actinomyces odontolyticus		0.65	0.19		1	99	
23113041	**Good identification**						**23113041**
Propionibacterium propionicum		97.35	0.47	LAC 99			
Actinomyces naeslundii		2.65	0.30	URE 10			
23113042	**Very good identification**						**23113042**
					COC	GRA	
Propionibacterium propionicum		99.30	0.66	LAC 99	1	99	
Megasphaera elsdenii		0.29	0.12		99	1	
Actinomyces odontolyticus		0.28	0.15		1	99	
23113043	**Good identification**						**23113043**
Propionibacterium propionicum		98.22	0.50	LAC 99			
Actinomyces naeslundii		1.78	0.30	URE 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23113100	**Excellent identification**					**23113100**
Propionibacterium propionicum		99.70	0.97			
Actinomyces odontolyticus		0.22	0.44			
Actinomyces naeslundii		0.03	0.48			
23113101	**Good identification**					**23113101**
					SPO	
Propionibacterium propionicum		94.23	0.81	URE 10	1	
Clostridium chauvoei		5.33	0.47		99	
Actinomyces naeslundii		0.42	0.51		1	
23113102	**Excellent identification**					**23113102**
Propionibacterium propionicum		99.87	1.00			
Actinomyces odontolyticus		0.09	0.41			
Actinomyces naeslundii		0.02	0.48			
23113103	**Very good identification**					**23113103**
					SPO	
Propionibacterium propionicum		99.01	0.84	URE 10	1	
Clostridium chauvoei		0.66	0.34		99	
Actinomyces naeslundii		0.29	0.51		1	
23113110	**Very good identification**					**23113110**
Propionibacterium propionicum		99.61	0.63	MLZ 1		
Actinomyces odontolyticus		0.22	0.10			
Actinomyces naeslundii		0.17	0.26			
23113111	**Acceptable identification**					**23113111**
					SPO	
Propionibacterium propionicum		92.62	0.47	MLZ 1	1	
Clostridium chauvoei		5.24	0.12	URE 10	99	
Actinomyces naeslundii		2.14	0.29		1	
23113112	**Very good identification**					**23113112**
Propionibacterium propionicum		99.80	0.66	MLZ 1		
Actinomyces naeslundii		0.11	0.26			
Actinomyces odontolyticus		0.09	0.07			
23113113	**Good identification**					**23113113**
					SPO	
Propionibacterium propionicum		97.84	0.50	MLZ 1	1	
Actinomyces naeslundii		1.51	0.29	URE 10	1	
Clostridium chauvoei		0.65	-.01		99	
23113120	**Good identification**					**23113120**
Propionibacterium propionicum		91.54	0.63	NAG 1		
Actinomyces odontolyticus		3.19	0.31			
Actinomyces naeslundii		2.97	0.48			
23113121	**Low discrimination**					**23113121**
					SPO	
Clostridium chauvoei		56.50	0.38	MNS 90	99	
Propionibacterium propionicum		30.24	0.47	MAN 1	1	
Actinomyces naeslundii		13.26	0.51		1	
23113122	**Very good identification**					**23113122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		95.88	0.66	NAG 1		
Actinomyces naeslundii		2.08	0.48			
Actinomyces odontolyticus		1.37	0.27			
23113123	**Low discrimination**					**23113123**
					SPO	
Propionibacterium propionicum		66.13	0.50	NAG 1	1	
Actinomyces naeslundii		19.33	0.51	URE 10	1	
Clostridium chauvoei		14.53	0.25		99	
23113140	**Excellent identification**					**23113140**
Propionibacterium propionicum		99.99	0.97			
Actinomyces naeslundii		0.01	0.35			
Actinomyces odontolyticus		0.00	0.10			
23113141	**Very good identification**					**23113141**
					SPO	
Propionibacterium propionicum		99.87	0.81	URE 10	1	
Actinomyces naeslundii		0.08	0.38		1	
Clostridium chauvoei		0.06	0.12		99	
23113142	**Excellent identification**					**23113142**
Propionibacterium propionicum		100.0	1.00			
Actinomyces naeslundii		0.00	0.35			
Actinomyces odontolyticus		0.00	0.07			
23113143	**Very good identification**					**23113143**
					SPO	
Propionibacterium propionicum		99.94	0.84	URE 10	1	
Actinomyces naeslundii		0.05	0.38		1	
Clostridium chauvoei		0.01	-.01		99	
23113150	**Very good identification**					**23113150**
Propionibacterium propionicum		100.0	0.63	MLZ 1		
23113151	**Good identification**					**23113151**
Propionibacterium propionicum		100.0	0.47	MLZ 1		
23113152	**Very good identification**					**23113152**
Propionibacterium propionicum		100.0	0.66	MLZ 1		
23113153	**Good identification**					**23113153**
Propionibacterium propionicum		100.0	0.50	MLZ 1		
23113160	**Very good identification**					**23113160**
Propionibacterium propionicum		99.19	0.63	NAG 1		
Actinomyces naeslundii		0.57	0.35			
Clostridium perfringens		0.25	0.06			
23113161	**Acceptable identification**					**23113161**
					SPO	
Propionibacterium propionicum		91.22	0.47	NAG 1	1	
Actinomyces naeslundii		7.06	0.38	URE 10	1	
Clostridium chauvoei		1.72	0.04		99	
23113162	**Very good identification**					**23113162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.54	0.66	NAG 1		
Actinomyces naeslundii		0.38	0.35			
Clostridium perfringens		0.08	0.01			
23113163	**Good identification**					**23113163**
Propionibacterium propionicum		95.09	0.50	NAG 1		
Actinomyces naeslundii		4.91	0.38	URE 10		
23113200	**Very good identification**					**23113200**
Actinomyces odontolyticus		99.24	0.58	MAN 1		
Propionibacterium propionicum		0.76	0.29			
23113202	**Good identification**					**23113202**
Actinomyces odontolyticus		98.16	0.54	MAN 1		
Propionibacterium propionicum		1.84	0.32			
23113220	**Good identification**					**23113220**
Actinomyces odontolyticus		100.0	0.44	MAN 1		
23113222	**Good identification**					**23113222**
Actinomyces odontolyticus		100.0	0.41	MAN 1		
23113300	**Low discrimination**					**23113300**
Propionibacterium propionicum		70.62	0.63	RHA 1		
Actinomyces odontolyticus		29.38	0.49			
23113301	**Acceptable identification**					**23113301**
					SPO	
Propionibacterium propionicum		91.37	0.47	RHA 1	1	
Clostridium chauvoei		5.17	0.12	URE 10	99	
Actinomyces odontolyticus		3.46	0.15		1	
23113302	**Low discrimination**					**23113302**
Propionibacterium propionicum		85.47	0.66	RHA 1		
Actinomyces odontolyticus		14.53	0.46			
23113303	**Good identification**					**23113303**
					SPO	
Propionibacterium propionicum		97.84	0.50	RHA 1	1	
Actinomyces odontolyticus		1.51	0.12	URE 10	1	
Clostridium chauvoei		0.65	-.01		99	
23113320	**Low discrimination**					**23113320**
Actinomyces odontolyticus		87.02	0.36	MAN 1		
Propionibacterium propionicum		12.98	0.29	NAG 14		
23113340	**Very good identification**					**23113340**
Propionibacterium propionicum		99.58	0.63	RHA 1		
Actinomyces odontolyticus		0.42	0.15			
23113341	**Good identification**					**23113341**
Propionibacterium propionicum		100.0	0.47	RHA 1		
23113342	**Very good identification**					**23113342**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		99.83	0.66	RHA 1			
Actinomyces odontolyticus		0.17	0.12				
23113343	**Good identification**						**23113343**
Propionibacterium propionicum		100.0	0.50	RHA 1			
23113500	**Very good identification**						**23113500**
Propionibacterium propionicum		98.10	0.63	ARA 1			
Lactobacillus fermentum		0.89	0.28				
Actinomyces meyeri		0.80	0.16				
23113501	**Acceptable identification**						**23113501**
					SPO		
Propionibacterium propionicum		94.64	0.47	ARA 1	1		
Clostridium chauvoei		5.36	0.12	URE 10	99		
23113502	**Very good identification**						**23113502**
Propionibacterium propionicum		99.31	0.66	ARA 1			
Lactobacillus fermentum		0.60	0.28				
Actinomyces odontolyticus		0.09	0.07				
23113503	**Good identification**						**23113503**
					SPO		
Propionibacterium propionicum		99.34	0.50	ARA 1	1		
Clostridium chauvoei		0.66	-.01	URE 10	99		
23113540	**Very good identification**						**23113540**
Propionibacterium propionicum		100.0	0.63	ARA 1			
23113541	**Good identification**						**23113541**
Propionibacterium propionicum		100.0	0.47	ARA 1			
23113542	**Very good identification**						**23113542**
Propionibacterium propionicum		100.0	0.66	ARA 1			
23113543	**Good identification**						**23113543**
Propionibacterium propionicum		100.0	0.50	ARA 1			
23114000	**Very good identification**						**23114000**
					COC	GRA	
Actinomyces odontolyticus		97.79	0.85		1	99	
Actinomyces meyeri		2.06	0.53		1	99	
Megasphaera elsdenii		0.12	0.32		99	1	
23114001	**Good identification**						**23114001**
Actinomyces odontolyticus		97.94	0.51	URE 1			
Actinomyces meyeri		2.06	0.18				
23114002	**Very good identification**						**23114002**
Actinomyces odontolyticus		99.81	0.81				
Actinomyces meyeri		0.11	0.27				
Actinomyces viscosus		0.05	0.18				
23114003	**Good identification**						**23114003**
Actinomyces odontolyticus		98.44	0.47	URE 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces viscosus		1.56	0.09					
23114010	**Good identification**							**23114010**
Actinomyces odontolyticus		97.94	0.51	MLZ 1				
Actinomyces meyeri		2.06	0.18					
23114012	**Good identification**							**23114012**
Actinomyces odontolyticus		100.0	0.47	MLZ 1				
23114020	**Very good identification**							**23114020**
Actinomyces odontolyticus		98.25	0.71	NAG 14				
Actinomyces meyeri		1.73	0.38					
Lactobacillus fermentum		0.01	0.12					
23114021	**Good identification**							**23114021**
Actinomyces odontolyticus		98.27	0.37	NAG 14				
Actinomyces meyeri		1.73	0.04	URE 1				
23114022	**Very good identification**							**23114022**
Actinomyces odontolyticus		99.89	0.68	NAG 14				
Actinomyces meyeri		0.09	0.12					
Lactobacillus fermentum		0.02	0.12					
23114030	**Good identification**							**23114030**
Actinomyces odontolyticus		98.27	0.37	MLZ 1				
Actinomyces meyeri		1.73	0.04	NAG 14				
23114040	**Low discrimination**							**23114040**
					COC	SPO	GRA	
Actinomyces odontolyticus		74.34	0.51	SOR 1	1	1	99	
Clostridium glycolicum		18.20	0.32		1	99	99	
Megasphaera elsdenii		5.90	0.29		99	1	1	
23114042	**Good identification**							**23114042**
					COC	SPO	GRA	
Actinomyces odontolyticus		98.19	0.47	SOR 1	1	1	99	
Megasphaera elsdenii		1.41	0.12		99	1	1	
Clostridium glycolicum		0.40	-.03		1	99	99	
23114060	**Good identification**							**23114060**
					SPO			
Actinomyces odontolyticus		96.82	0.37	NAG 14	1			
Actinomyces meyeri		1.71	0.04	SOR 1	1			
Clostridium glycolicum		1.47	-.03		99			
23114100	**Low discrimination**							**23114100**
Actinomyces odontolyticus		78.88	0.76					
Actinomyces meyeri		21.04	0.63					
Propionibacterium propionicum		0.05	0.29					
23114101	**Low discrimination**							**23114101**
					SPO			
Actinomyces odontolyticus		76.13	0.42	URE 1	1			
Actinomyces meyeri		20.31	0.29		1			
Clostridium chauvoei		3.02	0.12		99			
23114102	**Very good identification**							**23114102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		98.20	0.73			
Actinomyces meyeri		1.32	0.37			
Actinomyces viscosus		0.27	0.22			
23114103	**Low discrimination**					**23114103**
Actinomyces odontolyticus		88.39	0.39	URE 1		
Actinomyces viscosus		7.88	0.14			
Propionibacterium propionicum		1.53	0.15			
23114110	**Low discrimination**					**23114110**
Actinomyces odontolyticus		78.95	0.42	MLZ 1		
Actinomyces meyeri		21.06	0.29			
23114112	**Good identification**					**23114112**
Actinomyces odontolyticus		98.67	0.39	MLZ 1		
Actinomyces meyeri		1.33	0.03			
23114120	**Low discrimination**					**23114120**
Actinomyces odontolyticus		81.69	0.63	NAG 14		
Actinomyces meyeri		18.25	0.48			
Lactobacillus fermentum		0.06	0.16			
23114122	**Very good identification**					**23114122**
Actinomyces odontolyticus		98.77	0.59	NAG 14		
Actinomyces meyeri		1.11	0.22			
Lactobacillus fermentum		0.11	0.16			
23114140	**Low discrimination**					**23114140**
Actinomyces odontolyticus		74.72	0.42	SOR 1		
Actinomyces meyeri		19.93	0.29			
Propionibacterium propionicum		4.77	0.29			
23114142	**Low discrimination**					**23114142**
Actinomyces odontolyticus		85.50	0.39	SOR 1		
Propionibacterium propionicum		13.35	0.32			
Actinomyces meyeri		1.15	0.03			
23114200	**Very good identification**					**23114200**
Actinomyces odontolyticus		99.99	0.90			
Actinomyces meyeri		0.01	0.18			
23114201	**Good identification**					**23114201**
Actinomyces odontolyticus		100.0	0.56	URE 1		
23114202	**Very good identification**					**23114202**
Actinomyces odontolyticus		100.0	0.86			
23114203	**Good identification**					**23114203**
Actinomyces odontolyticus		100.0	0.52	URE 1		
23114210	**Good identification**					**23114210**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces odontolyticus	100.0	0.56	MLZ 1		
23114212	**Good identification**					**23114212**
	Actinomyces odontolyticus	100.0	0.52	MLZ 1		
23114220	**Very good identification**					**23114220**
	Actinomyces odontolyticus	99.99	0.76	NAG 14		
	Actinomyces meyeri	0.01	0.04			
23114221	**Good identification**					**23114221**
	Actinomyces odontolyticus	100.0	0.42	NAG 14		
23114222	**Very good identification**					**23114222**
	Actinomyces odontolyticus	100.0	0.73	NAG 14		
23114223	**Good identification**					**23114223**
	Actinomyces odontolyticus	100.0	0.39	NAG 14		
23114230	**Good identification**					**23114230**
	Actinomyces odontolyticus	100.0	0.42	MLZ 1		
23114232	**Good identification**					**23114232**
	Actinomyces odontolyticus	100.0	0.39	MLZ 1		
23114240	**Good identification**					**23114240**
					SPO	
	Actinomyces odontolyticus	99.87	0.56	SOR 1	1	
	Clostridium glycolicum	0.13	-.03		99	
23114242	**Good identification**					**23114242**
	Actinomyces odontolyticus	100.0	0.52	SOR 1		
23114260	**Good identification**					**23114260**
	Actinomyces odontolyticus	100.0	0.42	NAG 14		
23114262	**Good identification**					**23114262**
	Actinomyces odontolyticus	100.0	0.39	NAG 14		
23114300	**Very good identification**					**23114300**
	Actinomyces odontolyticus	99.86	0.81			
	Actinomyces meyeri	0.14	0.29			
23114301	**Good identification**					**23114301**
	Actinomyces odontolyticus	100.0	0.47	URE 1		
23114302	**Very good identification**					**23114302**
	Actinomyces odontolyticus	99.99	0.78			
	Actinomyces meyeri	0.01	0.03			
23114303	**Good identification**					**23114303**
	Actinomyces odontolyticus	100.0	0.43	URE 1		
23114310	**Good identification**					**23114310**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.47	MLZ 1		
23114312	**Good identification**					**23114312**
Actinomyces odontolyticus		100.0	0.43	MLZ 1		
23114320	**Very good identification**					**23114320**
Actinomyces odontolyticus		99.88	0.68	NAG 14		
Actinomyces meyeri		0.12	0.14			
23114322	**Very good identification**					**23114322**
Actinomyces odontolyticus		100.0	0.64	NAG 14		
23114340	**Good identification**					**23114340**
Actinomyces odontolyticus		100.0	0.47	SOR 1		
23114342	**Good identification**					**23114342**
Actinomyces odontolyticus		100.0	0.43	SOR 1		
23114400	**Low discrimination**					**23114400**
Actinomyces odontolyticus		65.46	0.51	ARA 1		
Actinomyces meyeri		34.11	0.42			
Lactobacillus fermentum		0.43	0.20			
23114402	**Good identification**					**23114402**
Actinomyces odontolyticus		96.44	0.47	ARA 1		
Actinomyces meyeri		2.54	0.16			
Lactobacillus fermentum		1.02	0.20			
23114420	**Low discrimination**					**23114420**
Actinomyces odontolyticus		68.97	0.37	ARA 1		
Actinomyces meyeri		30.11	0.27	NAG 14		
Lactobacillus fermentum		0.92	0.12			
23114500	**Low discrimination**					**23114500**
Actinomyces meyeri		86.46	0.53	NIT 1		
Actinomyces odontolyticus		13.10	0.42			
Lactobacillus fermentum		0.44	0.24			
23114502	**Low discrimination**					**23114502**
Actinomyces odontolyticus		72.05	0.39	ARA 1		
Actinomyces meyeri		24.00	0.27			
Lactobacillus fermentum		3.95	0.24			
23114520	**Low discrimination**					**23114520**
Actinomyces meyeri		83.80	0.38	NIT 1		
Actinomyces odontolyticus		15.16	0.29	NAG 12		
Lactobacillus fermentum		1.04	0.16			
23114600	**Good identification**					**23114600**
Actinomyces odontolyticus		99.73	0.56	ARA 1		
Actinomyces meyeri		0.27	0.08			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23114602	**Good identification**					**23114602**
Actinomyces odontolyticus		100.0	0.52	ARA 1		
23114620	**Good identification**					**23114620**
Actinomyces odontolyticus		100.0	0.42	ARA 1		
23114622	**Good identification**					**23114622**
Actinomyces odontolyticus		100.0	0.39	ARA 1		
23114700	**Good identification**					**23114700**
Actinomyces odontolyticus		96.68	0.47	ARA 1		
Actinomyces meyeri		3.32	0.18			
23114702	**Good identification**					**23114702**
Actinomyces odontolyticus		100.0	0.43	ARA 1		
23115000	**Good identification**					**23115000**
Actinomyces odontolyticus		94.58	0.82			
Actinomyces meyeri		4.48	0.56			
Lactobacillus fermentum		0.92	0.55			
23115001	**Acceptable identification**					**23115001**
					SPO	
Actinomyces odontolyticus		92.94	0.48	URE 1	1	
Actinomyces meyeri		4.40	0.21		1	
Clostridium chauvoei		1.75	0.12		99	
23115002	**Very good identification**					**23115002**
Actinomyces odontolyticus		98.16	0.78			
Lactobacillus fermentum		1.56	0.55			
Actinomyces meyeri		0.24	0.30			
23115003	**Good identification**					**23115003**
Actinomyces odontolyticus		96.53	0.44	URE 1		
Lactobacillus fermentum		1.54	0.20			
Actinomyces viscosus		1.41	0.06			
23115010	**Acceptable identification**					**23115010**
Actinomyces odontolyticus		94.60	0.48	MLZ 1		
Actinomyces meyeri		4.48	0.21			
Lactobacillus fermentum		0.92	0.20			
23115012	**Good identification**					**23115012**
Actinomyces odontolyticus		98.43	0.44	MLZ 1		
Lactobacillus fermentum		1.57	0.20			
23115020	**Good identification**					**23115020**
Actinomyces odontolyticus		94.34	0.68	NAG 14		
Actinomyces meyeri		3.74	0.41			
Lactobacillus fermentum		1.89	0.46			
23115022	**Very good identification**					**23115022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		96.65	0.65	NAG 14			
Lactobacillus fermentum		3.15	0.46				
Actinomyces meyeri		0.19	0.15				
23115040	**Acceptable identification**						**23115040**
					SPO		
Actinomyces odontolyticus		91.08	0.48	SOR 1	1		
Actinomyces meyeri		4.32	0.21		1		
Clostridium glycolicum		3.72	0.15		99		
23115042	**Good identification**						**23115042**
Actinomyces odontolyticus		98.43	0.44	SOR 1			
Lactobacillus fermentum		1.57	0.20				
23115100	**Low discrimination**						**23115100**
Actinomyces odontolyticus		60.52	0.73				
Actinomyces meyeri		36.32	0.66				
Lactobacillus fermentum		3.05	0.58				
23115101	**Low discrimination**						**23115101**
					SPO		
Clostridium chauvoei		78.02	0.47	MNS 90	99		
Actinomyces odontolyticus		13.23	0.39	XYL 1	1		
Actinomyces meyeri		7.94	0.32		1		
23115102	**Low discrimination**						**23115102**
Actinomyces odontolyticus		89.41	0.70				
Lactobacillus fermentum		7.36	0.58				
Actinomyces meyeri		2.71	0.40				
23115110	**Low discrimination**						**23115110**
Actinomyces odontolyticus		60.59	0.39	MLZ 1			
Actinomyces meyeri		36.36	0.32				
Lactobacillus fermentum		3.06	0.24				
23115112	**Low discrimination**						**23115112**
Actinomyces odontolyticus		89.88	0.36	MLZ 1			
Lactobacillus fermentum		7.40	0.24				
Actinomyces meyeri		2.72	0.06				
23115120	**Low discrimination**						**23115120**
Actinomyces odontolyticus		61.78	0.60	NAG 14			
Actinomyces meyeri		31.06	0.51				
Lactobacillus fermentum		6.38	0.50				
23115121	**Low discrimination**						**23115121**
					SPO		
Clostridium chauvoei		88.26	0.38	MNS 90	99		
Actinomyces odontolyticus		7.31	0.26	XYL 1	1		
Actinomyces meyeri		3.67	0.17		1		
23115122	**Low discrimination**						**23115122**
					GRA	CAT	
Actinomyces odontolyticus		81.04	0.56	NAG 14	99	10	
Lactobacillus fermentum		13.66	0.50		99	1	
Capnocytophaga ochracea		2.77	0.23		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23115140	**Low discrimination**					**23115140**
Actinomyces odontolyticus		57.27	0.39	SOR 1		
Actinomyces meyeri		34.37	0.32			
Propionibacterium propionicum		5.48	0.29			
23115142	**Low discrimination**					**23115142**
Actinomyces odontolyticus		74.25	0.36	SOR 1		
Propionibacterium propionicum		17.39	0.32			
Lactobacillus fermentum		6.11	0.24			
23115200	**Very good identification**					**23115200**
Actinomyces odontolyticus		99.97	0.87			
Actinomyces meyeri		0.03	0.21			
Lactobacillus fermentum		0.01	0.20			
23115201	**Good identification**					**23115201**
Actinomyces odontolyticus		100.0	0.53	URE 1		
23115202	**Very good identification**					**23115202**
Actinomyces odontolyticus		99.99	0.83			
Lactobacillus fermentum		0.01	0.20			
23115203	**Good identification**					**23115203**
Actinomyces odontolyticus		100.0	0.49	URE 1		
23115210	**Good identification**					**23115210**
Actinomyces odontolyticus		100.0	0.53	MLZ 1		
23115212	**Good identification**					**23115212**
Actinomyces odontolyticus		100.0	0.49	MLZ 1		
23115220	**Very good identification**					**23115220**
Actinomyces odontolyticus		99.97	0.73	NAG 14		
Actinomyces meyeri		0.02	0.07			
Lactobacillus fermentum		0.01	0.12			
23115221	**Good identification**					**23115221**
Actinomyces odontolyticus		100.0	0.39	NAG 14		
23115222	**Very good identification**					**23115222**
Actinomyces odontolyticus		99.98	0.70	NAG 14		
Lactobacillus fermentum		0.02	0.12			
23115223	**Good identification**					**23115223**
Actinomyces odontolyticus		100.0	0.36	NAG 14		
23115230	**Good identification**					**23115230**
Actinomyces odontolyticus		100.0	0.39	MLZ 1		
23115232	**Good identification**					**23115232**
Actinomyces odontolyticus		100.0	0.36	MLZ 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23115240	**Good identification**					**23115240**
Actinomyces odontolyticus		100.0	0.53	SOR 1		
23115242	**Good identification**					**23115242**
Actinomyces odontolyticus		100.0	0.49	SOR 1		
23115260	**Good identification**					**23115260**
Actinomyces odontolyticus		100.0	0.39	NAG 14		
23115262	**Good identification**					**23115262**
Actinomyces odontolyticus		100.0	0.36	NAG 14		
23115300	**Very good identification**					**23115300**
Actinomyces odontolyticus		99.66	0.78			
Actinomyces meyeri		0.31	0.32			
Lactobacillus fermentum		0.03	0.24			
23115301	**Good identification**					**23115301**
					SPO	
Actinomyces odontolyticus		97.02	0.44	URE 1	1	
Clostridium chauvoei		2.98	0.12		99	
23115302	**Very good identification**					**23115302**
Actinomyces odontolyticus		99.94	0.75			
Lactobacillus fermentum		0.04	0.24			
Actinomyces meyeri		0.02	0.06			
23115303	**Good identification**					**23115303**
					SPO	
Actinomyces odontolyticus		99.12	0.40	URE 1	1	
Clostridium chauvoei		0.88	-.01		99	
23115310	**Good identification**					**23115310**
Actinomyces odontolyticus		100.0	0.44	MLZ 1		
23115312	**Good identification**					**23115312**
Actinomyces odontolyticus		100.0	0.40	MLZ 1		
23115320	**Very good identification**					**23115320**
Actinomyces odontolyticus		99.69	0.65	NAG 14		
Actinomyces meyeri		0.26	0.17			
Lactobacillus fermentum		0.05	0.16			
23115322	**Very good identification**					**23115322**
Actinomyces odontolyticus		99.91	0.61	NAG 14		
Lactobacillus fermentum		0.09	0.16			
23115340	**Good identification**					**23115340**
Actinomyces odontolyticus		100.0	0.44	SOR 1		
23115342	**Good identification**					**23115342**
Actinomyces odontolyticus		100.0	0.40	SOR 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23115400	**Low discrimination**					**23115400**
Actinomyces meyeri		37.35	0.45	NIT 1		
Actinomyces odontolyticus		31.86	0.48			
Lactobacillus fermentum		30.79	0.55			
23115402	**Low discrimination**					**23115402**
Lactobacillus fermentum		59.82	0.55	NIT 1		
Actinomyces odontolyticus		37.93	0.44			
Actinomyces meyeri		2.25	0.19			
23115420	**Low discrimination**					**23115420**
Lactobacillus fermentum		49.96	0.46	NIT 1		
Actinomyces odontolyticus		25.24	0.34			
Actinomyces meyeri		24.80	0.30			
23115422	**Low discrimination**					**23115422**
Lactobacillus fermentum		75.47	0.46	NIT 1		
Actinomyces odontolyticus		23.37	0.31			
Actinomyces meyeri		1.16	0.05			
23115500	**Low discrimination**					**23115500**
Actinomyces meyeri		71.17	0.56	NIT 1		
Lactobacillus fermentum		23.93	0.58			
Actinomyces odontolyticus		4.79	0.39			
23115502	**Low discrimination**					**23115502**
Lactobacillus fermentum		81.99	0.58	NIT 1		
Actinomyces odontolyticus		10.06	0.36			
Actinomyces meyeri		7.54	0.30			
23115520	**Low discrimination**					**23115520**
Actinomyces meyeri		51.28	0.41	NIT 1		
Lactobacillus fermentum		42.14	0.50	NAG 12		
Actinomyces odontolyticus		4.12	0.26			
23115522	**Low discrimination**					**23115522**
Lactobacillus fermentum		83.91	0.50	NIT 1		
Actinomyces odontolyticus		5.03	0.22			
Bifidobacterium longum subsp. longum		4.90	0.22			
23115600	**Good identification**					**23115600**
Actinomyces odontolyticus		98.90	0.53	ARA 1		
Actinomyces meyeri		0.60	0.11			
Lactobacillus fermentum		0.50	0.20			
23115602	**Good identification**					**23115602**
Actinomyces odontolyticus		99.19	0.49	ARA 1		
Lactobacillus fermentum		0.81	0.20			
23115620	**Good identification**					**23115620**
Actinomyces odontolyticus		98.98	0.39	ARA 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		1.02	0.12	NAG 14			
23115622	**Good identification**						**23115622**
Actinomyces odontolyticus		98.35	0.36	ARA 1			
Lactobacillus fermentum		1.65	0.12	NAG 14			
23115700	**Acceptable identification**						**23115700**
Actinomyces odontolyticus		90.64	0.44	ARA 1			
Actinomyces meyeri		7.01	0.21				
Lactobacillus fermentum		2.36	0.24				
23115702	**Good identification**						**23115702**
Actinomyces odontolyticus		95.93	0.40	ARA 1			
Lactobacillus fermentum		4.07	0.24				
23116000	**Low discrimination**						**23116000**
					COC	GRA	
Actinomyces odontolyticus		86.20	0.51	MAN 1	1	99	
Megasphaera elsdenii		10.25	0.32		99	1	
Actinomyces meyeri		1.82	0.18		1	99	
23116002	**Acceptable identification**						**23116002**
					COC	GRA	
Actinomyces odontolyticus		93.38	0.47	MAN 1	1	99	
Propionibacterium propionicum		4.60	0.32		1	99	
Megasphaera elsdenii		2.01	0.15		99	1	
23116020	**Good identification**						**23116020**
Actinomyces odontolyticus		98.27	0.37	MAN 1			
Actinomyces meyeri		1.73	0.04	NAG 14			
23116100	**Low discrimination**						**23116100**
Propionibacterium propionicum		83.29	0.63	XYL 1			
Actinomyces odontolyticus		13.19	0.42				
Actinomyces meyeri		3.52	0.29				
23116101	**Good identification**						**23116101**
Propionibacterium propionicum		98.58	0.47	XYL 1			
Actinomyces odontolyticus		1.42	0.08	URE 10			
23116102	**Good identification**						**23116102**
Propionibacterium propionicum		93.85	0.66	XYL 1			
Actinomyces odontolyticus		6.07	0.39				
Actinomyces meyeri		0.08	0.03				
23116103	**Good identification**						**23116103**
Propionibacterium propionicum		100.0	0.50	XYL 1			
23116140	**Very good identification**						**23116140**
Propionibacterium propionicum		99.84	0.63	XYL 1			
Actinomyces odontolyticus		0.16	0.08				
23116141	**Good identification**						**23116141**
Propionibacterium propionicum		100.0	0.47	XYL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23116142	**Very good identification**					**23116142**
Propionibacterium propionicum		100.0	0.66	XYL 1		
23116143	**Good identification**					**23116143**
Propionibacterium propionicum		100.0	0.50	XYL 1		
23116200	**Good identification**					**23116200**
Actinomyces odontolyticus		100.0	0.56	MAN 1		
23116202	**Good identification**					**23116202**
Actinomyces odontolyticus		100.0	0.52	MAN 1		
23116220	**Good identification**					**23116220**
Actinomyces odontolyticus		100.0	0.42	MAN 1		
23116222	**Good identification**					**23116222**
Actinomyces odontolyticus		100.0	0.39	MAN 1		
23116300	**Good identification**					**23116300**
Actinomyces odontolyticus		96.82	0.47	MAN 1		
Propionibacterium propionicum		3.18	0.29			
23116302	**Acceptable identification**					**23116302**
Actinomyces odontolyticus		92.56	0.43	MAN 1		
Propionibacterium propionicum		7.44	0.32			
23117000	**Acceptable identification**					**23117000**
Actinomyces odontolyticus		91.97	0.48	MAN 1		
Actinomyces meyeri		4.36	0.21			
Propionibacterium propionicum		2.78	0.29			
23117002	**Acceptable identification**					**23117002**
Actinomyces odontolyticus		91.09	0.44	MAN 1		
Propionibacterium propionicum		6.74	0.32			
Lactobacillus fermentum		1.45	0.20			
23117100	**Low discrimination**					**23117100**
Propionibacterium propionicum		84.85	0.63	XYL 1		
Actinomyces odontolyticus		8.96	0.39			
Actinomyces meyeri		5.38	0.32			
23117101	**Acceptable identification**					**23117101**
					SPO	
Propionibacterium propionicum		94.64	0.47	XYL 1	1	
Clostridium chauvoei		5.36	0.12	URE 10	99	
23117102	**Very good identification**					**23117102**
Propionibacterium propionicum		95.03	0.66	XYL 1		
Actinomyces odontolyticus		4.10	0.36			
Actinomyces israelii		0.41	0.26			
23117103	**Good identification**					**23117103**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.34	0.50	XYL 1	1	
Clostridium chauvoei		0.66	-.01	URE 10	99	
23117140	**Very good identification**					**23117140**
Propionibacterium propionicum		99.83	0.63	XYL 1		
Actinomyces israelii		0.17	0.16			
23117141	**Good identification**					**23117141**
Propionibacterium propionicum		100.0	0.47	XYL 1		
23117142	**Very good identification**					**23117142**
Propionibacterium propionicum		99.83	0.66	XYL 1		
Actinomyces israelii		0.17	0.19			
23117143	**Good identification**					**23117143**
Propionibacterium propionicum		100.0	0.50	XYL 1		
23117200	**Good identification**					**23117200**
Actinomyces odontolyticus		100.0	0.53	MAN 1		
23117202	**Good identification**					**23117202**
Actinomyces odontolyticus		100.0	0.49	MAN 1		
23117220	**Good identification**					**23117220**
Actinomyces odontolyticus		100.0	0.39	MAN 1		
23117222	**Good identification**					**23117222**
Actinomyces odontolyticus		100.0	0.36	MAN 1		
23117300	**Good identification**					**23117300**
Actinomyces odontolyticus		95.30	0.44	MAN 1		
Propionibacterium propionicum		4.70	0.29			
23117302	**Low discrimination**					**23117302**
Actinomyces odontolyticus		89.24	0.40	MAN 1		
Propionibacterium propionicum		10.76	0.32			
23120000	**Low discrimination**					**23120000**
					SPO	
Actinomyces odontolyticus		46.92	0.38	FRU 92	1	
Clostridium botulinum biovar A		46.19	0.30	TRE 1	99	
Clostridium sporogenes		4.52	0.11		99	
23120200	**Good identification**					**23120200**
Actinomyces odontolyticus		100.0	0.43	FRU 92		
23120202	**Good identification**					**23120202**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23121101	**Very good identification**					**23121101**
					SPO	
Clostridium chauvoei		99.91	0.58	MNS 90	99	
Actinomyces naeslundii		0.09	0.29	TRE 10	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23121103	**Good identification**					**23121103**
					SPO	
Clostridium chauvoei		98.85	0.45	MNS 90	99	
Actinomyces naeslundii		0.49	0.29	TRE 10	1	
Actinomyces viscosus		0.40	0.03		1	
23121121	**Good identification**					**23121121**
					SPO	
Clostridium chauvoei		99.74	0.50	MNS 90	99	
Actinomyces naeslundii		0.26	0.29	TRE 10	1	
23121123	**Good identification**					**23121123**
					SPO	
Clostridium chauvoei		98.54	0.37	MNS 90	99	
Actinomyces naeslundii		1.46	0.29	TRE 10	1	
23121200	**Good identification**					**23121200**
Actinomyces odontolyticus		100.0	0.40	FRU 92		
23121202	**Good identification**					**23121202**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23122100	**Very good identification**					**23122100**
Propionibacterium propionicum		100.0	0.61	FRU 99		
23122101	**Good identification**					**23122101**
Propionibacterium propionicum		100.0	0.45	FRU 99		
23122102	**Very good identification**					**23122102**
Propionibacterium propionicum		100.0	0.64	FRU 99		
23122103	**Good identification**					**23122103**
Propionibacterium propionicum		100.0	0.48	FRU 99		
23122140	**Very good identification**					**23122140**
Propionibacterium propionicum		100.0	0.61	FRU 99		
23122141	**Good identification**					**23122141**
Propionibacterium propionicum		100.0	0.45	FRU 99		
23122142	**Very good identification**					**23122142**
Propionibacterium propionicum		100.0	0.64	FRU 99		
23122143	**Good identification**					**23122143**
Propionibacterium propionicum		100.0	0.48	FRU 99		
23123100	**Very good identification**					**23123100**
Propionibacterium propionicum		99.60	0.61	FRU 99		
Actinomyces naeslundii		0.40	0.31			
23123101	**Low discrimination**					**23123101**
					SPO	
Propionibacterium propionicum		72.42	0.45	FRU 99	1	
Clostridium chauvoei		23.62	0.24	URE 10	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		3.97	0.34		1			
23123102	**Very good identification**							**23123102**
Propionibacterium propionicum		99.73	0.64	FRU 99				
Actinomyces naeslundii		0.27	0.31					
23123103	**Acceptable identification**							**23123103**
					SPO			
Propionibacterium propionicum		93.03	0.48	FRU 99	1			
Clostridium chauvoei		3.57	0.11	URE 10	99			
Actinomyces naeslundii		3.40	0.34		1			
23123140	**Very good identification**							**23123140**
Propionibacterium propionicum		100.0	0.61	FRU 99				
23123141	**Good identification**							**23123141**
Propionibacterium propionicum		100.0	0.45	FRU 99				
23123142	**Very good identification**							**23123142**
Propionibacterium propionicum		100.0	0.64	FRU 99				
23123143	**Good identification**							**23123143**
Propionibacterium propionicum		100.0	0.48	FRU 99				
23124200	**Good identification**							**23124200**
Actinomyces odontolyticus		100.0	0.38	FRU 92				
23130000	**Acceptable identification**							**23130000**
					COC	CAT	GRA	
Actinomyces odontolyticus		91.48	0.56	TRE 1	1	10	99	
Megasphaera elsdenii		5.36	0.32		99	1	1	
Propionibacterium granulosum		0.90	0.24		1	90	99	
23130002	**Low discrimination**							**23130002**
Actinomyces odontolyticus		82.72	0.52	TRE 1				
Actinomyces viscosus		14.42	0.32					
Propionibacterium propionicum		1.64	0.30					
23130020	**Good identification**							**23130020**
					CAT			
Actinomyces odontolyticus		96.62	0.43	TRE 1	10			
Clostridium perfringens		1.93	0.08	NAG 14	1			
Propionibacterium granulosum		1.45	0.14		90			
23130022	**Good identification**							**23130022**
Actinomyces odontolyticus		97.45	0.39	TRE 1				
Clostridium perfringens		1.49	0.03	NAG 14				
Actinomyces viscosus		1.05	-.02					
23130100	**Low discrimination**							**23130100**
Propionibacterium propionicum		70.00	0.61	MAN 99				
Actinomyces odontolyticus		27.51	0.47					
Actinomyces meyeri		1.86	0.24					
23130101	**Low discrimination**							**23130101**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		54.97	0.45	MAN 99	1	
Clostridium chauvoei		41.84	0.30	URE 10	99	
Actinomyces odontolyticus		1.96	0.13		1	
23130102	**Low discrimination**					**23130102**
Propionibacterium propionicum		75.80	0.64	MAN 99		
Actinomyces odontolyticus		12.17	0.44			
Actinomyces viscosus		11.95	0.37			
23130103	**Low discrimination**					**23130103**
					SPO	
Propionibacterium propionicum		62.82	0.48	MAN 99	1	
Actinomyces viscosus		29.70	0.29	URE 10	1	
Clostridium chauvoei		5.63	0.17		99	
23130140	**Very good identification**					**23130140**
					SPO	
Propionibacterium propionicum		99.42	0.61	MAN 99	1	
Actinomyces odontolyticus		0.40	0.13		1	
Clostridium botulinum biovar B		0.19	-.03		99	
23130141	**Good identification**					**23130141**
Propionibacterium propionicum		100.0	0.45	MAN 99		
23130142	**Very good identification**					**23130142**
Propionibacterium propionicum		99.68	0.64	MAN 99		
Actinomyces odontolyticus		0.16	0.10			
Actinomyces viscosus		0.16	0.03			
23130143	**Good identification**					**23130143**
Propionibacterium propionicum		100.0	0.48	MAN 99		
23130200	**Very good identification**					**23130200**
Actinomyces odontolyticus		100.0	0.61	TRE 1		
23130202	**Very good identification**					**23130202**
Actinomyces odontolyticus		99.91	0.57	TRE 1		
Actinomyces viscosus		0.09	-.02			
23130220	**Good identification**					**23130220**
Actinomyces odontolyticus		100.0	0.47	TRE 1		
23130222	**Good identification**					**23130222**
Actinomyces odontolyticus		100.0	0.44	TRE 1		
23130300	**Good identification**					**23130300**
Actinomyces odontolyticus		98.69	0.52	TRE 1		
Propionibacterium propionicum		1.31	0.27			
23130302	**Good identification**					**23130302**
Actinomyces odontolyticus		96.38	0.49	TRE 1		
Propionibacterium propionicum		3.12	0.30			
Actinomyces viscosus		0.49	0.03			
23130320	**Good identification**					**23130320**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.39	TRE 1		
23130322	**Good identification**					**23130322**
Actinomyces odontolyticus		100.0	0.35	TRE 1		
23131000	**Acceptable identification**					**23131000**
Actinomyces odontolyticus		92.37	0.53	TRE 1		
Actinomyces naeslundii		1.55	0.40			
Clostridium perfringens		1.38	0.17			
23131002	**Low discrimination**					**23131002**
Actinomyces odontolyticus		80.11	0.49	TRE 1		
Actinomyces viscosus		12.84	0.29			
Propionibacterium propionicum		2.39	0.30			
23131020	**Low discrimination**					**23131020**
Clostridium perfringens		72.01	0.43	MNS 99		
Actinomyces odontolyticus		24.29	0.40	LAC 88		
Actinomyces naeslundii		2.51	0.40			
23131022	**Low discrimination**					**23131022**
Clostridium perfringens		65.56	0.37	MNS 99		
Actinomyces odontolyticus		28.80	0.36	LAC 88		
Actinomyces naeslundii		4.85	0.40			
23131023	**Low discrimination**					**23131023**
					SPO	
Actinomyces naeslundii		75.26	0.43	MNS 88	1	
Clostridium chauvoei		17.89	0.09	RAF 92	99	
Clostridium perfringens		6.85	0.03		84	
23131100	**Low discrimination**					**23131100**
Propionibacterium propionicum		64.77	0.61	MAN 99		
Actinomyces odontolyticus		16.97	0.44			
Clostridium perfringens		5.88	0.31			
23131101	**Very good identification**					**23131101**
					SPO	
Clostridium chauvoei		97.97	0.64	MNS 90	99	
Propionibacterium propionicum		1.30	0.45	TRE 10	1	
Actinomyces naeslundii		0.70	0.51		1	
23131102	**Low discrimination**					**23131102**
Propionibacterium propionicum		78.37	0.64	MAN 99		
Actinomyces odontolyticus		8.39	0.41			
Actinomyces viscosus		7.57	0.33			
23131103	**Low discrimination**					**23131103**
					SPO	
Clostridium chauvoei		84.25	0.51	MNS 90	99	
Propionibacterium propionicum		9.51	0.48	TRE 10	1	
Actinomyces naeslundii		3.40	0.51		1	
23131120	**Very good identification**					**23131120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		95.68	0.57	MNS 99		
Actinomyces odontolyticus		1.39	0.31			
Actinomyces naeslundii		1.29	0.48			
23131121	**Very good identification**					**23131121**
					SPO	
Clostridium chauvoei		96.90	0.56	MNS 90	99	
Actinomyces naeslundii		2.07	0.51	TRE 10	1	
Clostridium perfringens		1.03	0.23		84	
23131122	**Acceptable identification**					**23131122**
Clostridium perfringens		93.70	0.52	MNS 99		
Actinomyces naeslundii		2.69	0.48			
Actinomyces odontolyticus		1.77	0.27			
23131123	**Low discrimination**					**23131123**
					SPO	
Clostridium chauvoei		86.75	0.43	MNS 90	99	
Actinomyces naeslundii		10.49	0.51	TRE 10	1	
Clostridium perfringens		2.46	0.18		84	
23131140	**Very good identification**					**23131140**
Propionibacterium propionicum		97.38	0.61	MAN 99		
Clostridium perfringens		1.68	0.19			
Actinomyces naeslundii		0.68	0.35			
23131141	**Low discrimination**					**23131141**
					SPO	
Propionibacterium propionicum		53.89	0.45	MAN 99	1	
Clostridium chauvoei		41.01	0.30	URE 10	99	
Actinomyces naeslundii		5.10	0.38		1	
23131142	**Very good identification**					**23131142**
Propionibacterium propionicum		98.80	0.64	MAN 99		
Clostridium perfringens		0.54	0.14			
Actinomyces naeslundii		0.46	0.35			
23131143	**Low discrimination**					**23131143**
					SPO	
Propionibacterium propionicum		86.76	0.48	MAN 99	1	
Clostridium chauvoei		7.77	0.17	URE 10	99	
Actinomyces naeslundii		5.47	0.38		1	
23131160	**Good identification**					**23131160**
Clostridium perfringens		97.03	0.45	MNS 99		
Propionibacterium propionicum		1.75	0.27			
Actinomyces naeslundii		1.22	0.35			
23131162	**Acceptable identification**					**23131162**
Clostridium perfringens		92.24	0.39	MNS 99		
Propionibacterium propionicum		5.31	0.30			
Actinomyces naeslundii		2.46	0.35			
23131163	**Low discrimination**					**23131163**
					SPO	
Actinomyces naeslundii		53.01	0.38	MNS 88	1	
Clostridium chauvoei		25.09	0.09	RAF 92	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		13.41	0.05		84			
23131200	**Very good identification**							**23131200**
Actinomyces odontolyticus		100.0	0.58	TRE 1				
23131202	**Good identification**							**23131202**
Actinomyces odontolyticus		100.0	0.54	TRE 1				
23131220	**Good identification**							**23131220**
Actinomyces odontolyticus		98.48	0.44	TRE 1				
Clostridium perfringens		1.52	0.08	NAG 14				
23131222	**Good identification**							**23131222**
Actinomyces odontolyticus		98.83	0.41	TRE 1				
Clostridium perfringens		1.17	0.03	NAG 14				
23131300	**Good identification**							**23131300**
Actinomyces odontolyticus		98.05	0.49	TRE 1				
Propionibacterium propionicum		1.95	0.27					
23131302	**Acceptable identification**							**23131302**
Actinomyces odontolyticus		94.94	0.46	TRE 1				
Propionibacterium propionicum		4.62	0.30					
Actinomyces viscosus		0.45	-.01					
23131320	**Low discrimination**							**23131320**
Actinomyces odontolyticus		73.62	0.36	TRE 1				
Clostridium perfringens		26.38	0.23	NAG 14				
23132000	**Good identification**							**23132000**
					COC	GRA		
Propionibacterium propionicum		91.35	0.61	LAC 99	1	99		
Megasphaera elsdenii		6.66	0.32		99	1		
Actinomyces odontolyticus		1.15	0.22		1	99		
23132001	**Good identification**							**23132001**
Propionibacterium propionicum		98.50	0.45	LAC 99				
Actinomyces naeslundii		1.50	0.24	URE 10				
23132002	**Very good identification**							**23132002**
					COC	GRA		
Propionibacterium propionicum		98.72	0.64	LAC 99	1	99		
Megasphaera elsdenii		0.53	0.15		99	1		
Actinomyces odontolyticus		0.51	0.18		1	99		
23132003	**Good identification**							**23132003**
Propionibacterium propionicum		99.00	0.48	LAC 99				
Actinomyces naeslundii		1.01	0.24	URE 10				
23132040	**Very good identification**							**23132040**
					COC	SPO	GRA	
Propionibacterium propionicum		95.19	0.61	LAC 99	1	1	99	
Megasphaera elsdenii		4.63	0.29		99	1	1	
Clostridium botulinum biovar B		0.18	-.03		1	99	99	
23132041	**Good identification**							**23132041**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		100.0	0.45	LAC 99			
23132042	**Very good identification**						**23132042**
					COC	GRA	
Propionibacterium propionicum		99.64	0.64	LAC 99	1	99	
Megasphaera elsdenii		0.36	0.12		99	1	
23132043	**Good identification**						**23132043**
Propionibacterium propionicum		100.0	0.48	LAC 99			
23132100	**Excellent identification**						**23132100**
Propionibacterium propionicum		99.99	0.96				
Actinomyces odontolyticus		0.00	0.13				
Actinomyces naeslundii		0.00	0.29				
23132101	**Very good identification**						**23132101**
Propionibacterium propionicum		99.96	0.79	URE 10			
Actinomyces naeslundii		0.04	0.32				
23132102	**Excellent identification**						**23132102**
Propionibacterium propionicum		100.0	0.99				
Actinomyces naeslundii		0.00	0.29				
Actinomyces odontolyticus		0.00	0.10				
23132103	**Very good identification**						**23132103**
Propionibacterium propionicum		99.97	0.82	URE 10			
Actinomyces naeslundii		0.03	0.32				
23132110	**Very good identification**						**23132110**
Propionibacterium propionicum		100.0	0.61	MLZ 1			
23132111	**Good identification**						**23132111**
Propionibacterium propionicum		100.0	0.45	MLZ 1			
23132112	**Very good identification**						**23132112**
Propionibacterium propionicum		100.0	0.64	MLZ 1			
23132113	**Good identification**						**23132113**
Propionibacterium propionicum		100.0	0.48	MLZ 1			
23132120	**Very good identification**						**23132120**
Propionibacterium propionicum		99.68	0.61	NAG 1			
Actinomyces naeslundii		0.32	0.29				
23132121	**Good identification**						**23132121**
Propionibacterium propionicum		95.85	0.45	NAG 1			
Actinomyces naeslundii		4.16	0.32	URE 10			
23132122	**Very good identification**						**23132122**
Propionibacterium propionicum		99.79	0.64	NAG 1			
Actinomyces naeslundii		0.21	0.29				
23132123	**Good identification**						**23132123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		97.19	0.48	NAG 1		
Actinomyces naeslundii		2.81	0.32	URE 10		
23132140	**Excellent identification**					**23132140**
Propionibacterium propionicum		100.0	0.96			
23132141	**Very good identification**					**23132141**
Propionibacterium propionicum		100.0	0.79	URE 10		
23132142	**Excellent identification**					**23132142**
Propionibacterium propionicum		100.0	0.99			
23132143	**Very good identification**					**23132143**
Propionibacterium propionicum		100.0	0.82	URE 10		
23132150	**Very good identification**					**23132150**
Propionibacterium propionicum		100.0	0.61	MLZ 1		
23132151	**Good identification**					**23132151**
Propionibacterium propionicum		100.0	0.45	MLZ 1		
23132152	**Very good identification**					**23132152**
Propionibacterium propionicum		100.0	0.64	MLZ 1		
23132153	**Good identification**					**23132153**
Propionibacterium propionicum		100.0	0.48	MLZ 1		
23132160	**Very good identification**					**23132160**
Propionibacterium propionicum		100.0	0.61	NAG 1		
23132161	**Good identification**					**23132161**
Propionibacterium propionicum		100.0	0.45	NAG 1		
23132162	**Very good identification**					**23132162**
Propionibacterium propionicum		100.0	0.64	NAG 1		
23132163	**Good identification**					**23132163**
Propionibacterium propionicum		100.0	0.48	NAG 1		
23132300	**Very good identification**					**23132300**
Propionibacterium propionicum		99.24	0.61	RHA 1		
Actinomyces odontolyticus		0.77	0.18			
23132301	**Good identification**					**23132301**
Propionibacterium propionicum		100.0	0.45	RHA 1		
23132302	**Very good identification**					**23132302**
Propionibacterium propionicum		99.69	0.64	RHA 1		
Actinomyces odontolyticus		0.31	0.15			
23132303	**Good identification**					**23132303**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.48	RHA 1		
23132340	**Very good identification**					**23132340**
Propionibacterium propionicum		100.0	0.61	RHA 1		
23132341	**Good identification**					**23132341**
Propionibacterium propionicum		100.0	0.45	RHA 1		
23132342	**Very good identification**					**23132342**
Propionibacterium propionicum		100.0	0.64	RHA 1		
23132343	**Good identification**					**23132343**
Propionibacterium propionicum		100.0	0.48	RHA 1		
23132500	**Very good identification**					**23132500**
Propionibacterium propionicum		100.0	0.61	ARA 1		
23132501	**Good identification**					**23132501**
Propionibacterium propionicum		100.0	0.45	ARA 1		
23132502	**Very good identification**					**23132502**
Propionibacterium propionicum		100.0	0.64	ARA 1		
23132503	**Good identification**					**23132503**
Propionibacterium propionicum		100.0	0.48	ARA 1		
23132540	**Very good identification**					**23132540**
Propionibacterium propionicum		100.0	0.61	ARA 1		
23132541	**Good identification**					**23132541**
Propionibacterium propionicum		100.0	0.45	ARA 1		
23132542	**Very good identification**					**23132542**
Propionibacterium propionicum		100.0	0.64	ARA 1		
23132543	**Good identification**					**23132543**
Propionibacterium propionicum		100.0	0.48	ARA 1		
23133000	**Very good identification**					**23133000**
					CAT	
Propionibacterium propionicum		95.73	0.61	LAC 99	1	
Actinomyces naeslundii		2.59	0.45		25	
Propionibacterium granulosum		0.88	0.22		90	
23133001	**Low discrimination**					**23133001**
Propionibacterium propionicum		73.23	0.45	LAC 99		
Actinomyces naeslundii		26.77	0.48	URE 10		
23133002	**Very good identification**					**23133002**
Propionibacterium propionicum		97.75	0.64	LAC 99		
Actinomyces naeslundii		1.77	0.45			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces odontolyticus	0.34	0.15			
23133003	**Low discrimination**					**23133003**
	Propionibacterium propionicum	80.41	0.48	LAC 99		
	Actinomyces naeslundii	19.59	0.48	URE 10		
23133020	**Low discrimination**					**23133020**
	Actinomyces naeslundii	60.33	0.45	MNS 88		
	Propionibacterium propionicum	22.51	0.27	RAF 92		
	Clostridium perfringens	9.01	0.08			
23133021	**Good identification**					**23133021**
	Actinomyces naeslundii	100.0	0.48	MNS 88		
23133022	**Low discrimination**					**23133022**
	Actinomyces naeslundii	61.36	0.45	MNS 88		
	Propionibacterium propionicum	34.33	0.30	RAF 92		
	Clostridium perfringens	4.31	0.03			
23133023	**Good identification**					**23133023**
	Actinomyces naeslundii	96.02	0.48	MNS 88		
	Propionibacterium propionicum	3.98	0.14	RAF 92		
23133040	**Very good identification**					**23133040**
	Propionibacterium propionicum	99.52	0.61	LAC 99		
	Actinomyces naeslundii	0.48	0.32			
23133041	**Acceptable identification**					**23133041**
	Propionibacterium propionicum	93.94	0.45	LAC 99		
	Actinomyces naeslundii	6.06	0.35	URE 10		
23133042	**Very good identification**					**23133042**
	Propionibacterium propionicum	99.68	0.64	LAC 99		
	Actinomyces naeslundii	0.32	0.32			
23133043	**Good identification**					**23133043**
	Propionibacterium propionicum	95.88	0.48	LAC 99		
	Actinomyces naeslundii	4.12	0.35	URE 10		
23133100	**Excellent identification**					**23133100**
	Propionibacterium propionicum	99.92	0.96			
	Actinomyces naeslundii	0.08	0.53			
	Actinomyces israelii	0.00	0.15			
23133101	**Very good identification**					**23133101**
					SPO	
	Propionibacterium propionicum	98.21	0.79	URE 10	1	
	Actinomyces naeslundii	1.03	0.56		1	
	Clostridium chauvoei	0.76	0.30		99	
23133102	**Excellent identification**					**23133102**
	Propionibacterium propionicum	99.94	0.99			
	Actinomyces naeslundii	0.05	0.53			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.00	0.18			
23133103	**Very good identification**					**23133103**
					SPO	
Propionibacterium propionicum		99.22	0.82	URE 10	1	
Actinomyces naeslundii		0.70	0.56		1	
Clostridium chauvoei		0.09	0.17		99	
23133110	**Very good identification**					**23133110**
Propionibacterium propionicum		99.60	0.61	MLZ 1		
Actinomyces naeslundii		0.40	0.31			
23133111	**Acceptable identification**					**23133111**
Propionibacterium propionicum		94.81	0.45	MLZ 1		
Actinomyces naeslundii		5.19	0.34	URE 10		
23133112	**Very good identification**					**23133112**
Propionibacterium propionicum		99.73	0.64	MLZ 1		
Actinomyces naeslundii		0.27	0.31			
23133113	**Good identification**					**23133113**
Propionibacterium propionicum		96.48	0.48	MLZ 1		
Actinomyces naeslundii		3.52	0.34	URE 10		
23133120	**Good identification**					**23133120**
Propionibacterium propionicum		90.36	0.61	NAG 1		
Actinomyces naeslundii		6.96	0.53			
Clostridium perfringens		2.68	0.23			
23133121	**Low discrimination**					**23133121**
					SPO	
Actinomyces naeslundii		45.35	0.56	MNS 88	1	
Propionibacterium propionicum		43.59	0.45	RAF 92	1	
Clostridium chauvoei		11.06	0.22		99	
23133122	**Good identification**					**23133122**
Propionibacterium propionicum		94.28	0.64	NAG 1		
Actinomyces naeslundii		4.84	0.53			
Clostridium perfringens		0.88	0.18			
23133123	**Low discrimination**					**23133123**
					SPO	
Propionibacterium propionicum		58.02	0.48	NAG 1	1	
Actinomyces naeslundii		40.25	0.56	URE 10	1	
Clostridium chauvoei		1.73	0.09		99	
23133140	**Excellent identification**					**23133140**
Propionibacterium propionicum		99.99	0.96			
Actinomyces naeslundii		0.01	0.40			
23133141	**Very good identification**					**23133141**
Propionibacterium propionicum		99.82	0.79	URE 10		
Actinomyces naeslundii		0.19	0.43			
23133142	**Excellent identification**					**23133142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.99	0.99			
Actinomyces naeslundii		0.01	0.40			
23133143	**Very good identification**					**23133143**
Propionibacterium propionicum		99.88	0.82	URE 10		
Actinomyces naeslundii		0.12	0.43			
23133150	**Very good identification**					**23133150**
Propionibacterium propionicum		100.0	0.61	MLZ 1		
23133151	**Good identification**					**23133151**
Propionibacterium propionicum		100.0	0.45	MLZ 1		
23133152	**Very good identification**					**23133152**
Propionibacterium propionicum		100.0	0.64	MLZ 1		
23133153	**Good identification**					**23133153**
Propionibacterium propionicum		100.0	0.48	MLZ 1		
23133160	**Very good identification**					**23133160**
Propionibacterium propionicum		98.11	0.61	NAG 1		
Actinomyces naeslundii		1.33	0.40			
Clostridium perfringens		0.55	0.11			
23133161	**Low discrimination**					**23133161**
Propionibacterium propionicum		84.49	0.45	NAG 1		
Actinomyces naeslundii		15.51	0.43	URE 10		
23133162	**Very good identification**					**23133162**
Propionibacterium propionicum		98.93	0.64	NAG 1		
Actinomyces naeslundii		0.90	0.40			
Clostridium perfringens		0.18	0.05			
23133163	**Low discrimination**					**23133163**
Propionibacterium propionicum		89.10	0.48	NAG 1		
Actinomyces naeslundii		10.91	0.43	URE 10		
23133300	**Very good identification**					**23133300**
Propionibacterium propionicum		99.49	0.61	RHA 1		
Actinomyces odontolyticus		0.51	0.15			
23133301	**Good identification**					**23133301**
Propionibacterium propionicum		100.0	0.45	RHA 1		
23133302	**Very good identification**					**23133302**
Propionibacterium propionicum		99.79	0.64	RHA 1		
Actinomyces odontolyticus		0.21	0.12			
23133303	**Good identification**					**23133303**
Propionibacterium propionicum		100.0	0.48	RHA 1		
23133340	**Very good identification**					**23133340**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.61	RHA 1		
23133341	**Good identification**					**23133341**
Propionibacterium propionicum		100.0	0.45	RHA 1		
23133342	**Very good identification**					**23133342**
Propionibacterium propionicum		100.0	0.64	RHA 1		
23133343	**Good identification**					**23133343**
Propionibacterium propionicum		100.0	0.48	RHA 1		
23133500	**Very good identification**					**23133500**
Propionibacterium propionicum		99.86	0.61	ARA 1		
Actinomyces israelii		0.14	0.14			
23133501	**Good identification**					**23133501**
Propionibacterium propionicum		100.0	0.45	ARA 1		
23133502	**Very good identification**					**23133502**
Propionibacterium propionicum		99.86	0.64	ARA 1		
Actinomyces israelii		0.14	0.17			
23133503	**Good identification**					**23133503**
Propionibacterium propionicum		100.0	0.48	ARA 1		
23133540	**Very good identification**					**23133540**
Propionibacterium propionicum		100.0	0.61	ARA 1		
23133541	**Good identification**					**23133541**
Propionibacterium propionicum		100.0	0.45	ARA 1		
23133542	**Very good identification**					**23133542**
Propionibacterium propionicum		100.0	0.64	ARA 1		
23133543	**Good identification**					**23133543**
Propionibacterium propionicum		100.0	0.48	ARA 1		
23134000	**Good identification**					**23134000**
Actinomyces odontolyticus		97.94	0.51	TRE 1		
Actinomyces meyeri		2.06	0.18			
23134002	**Good identification**					**23134002**
Actinomyces odontolyticus		99.64	0.47	TRE 1		
Actinomyces viscosus		0.36	-.02			
23134020	**Good identification**					**23134020**
Actinomyces odontolyticus		98.27	0.37	TRE 1		
Actinomyces meyeri		1.73	0.04	NAG 14		
23134100	**Low discrimination**					**23134100**
Actinomyces odontolyticus		75.82	0.42	TRE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		20.22	0.29			
Propionibacterium propionicum		3.96	0.27			
23134102	**Low discrimination**					**23134102**
Actinomyces odontolyticus		86.11	0.39	TRE 1		
Propionibacterium propionicum		11.00	0.30			
Actinomyces viscosus		1.73	0.03			
23134200	**Good identification**					**23134200**
Actinomyces odontolyticus		100.0	0.56	TRE 1		
23134202	**Good identification**					**23134202**
Actinomyces odontolyticus		100.0	0.52	TRE 1		
23134220	**Good identification**					**23134220**
Actinomyces odontolyticus		100.0	0.42	TRE 1		
23134222	**Good identification**					**23134222**
Actinomyces odontolyticus		100.0	0.39	TRE 1		
23134300	**Good identification**					**23134300**
Actinomyces odontolyticus		100.0	0.47	TRE 1		
23134302	**Good identification**					**23134302**
Actinomyces odontolyticus		100.0	0.43	TRE 1		
23135000	**Acceptable identification**					**23135000**
Actinomyces odontolyticus		94.60	0.48	TRE 1		
Actinomyces meyeri		4.48	0.21			
Lactobacillus fermentum		0.92	0.20			
23135002	**Good identification**					**23135002**
Actinomyces odontolyticus		98.43	0.44	TRE 1		
Lactobacillus fermentum		1.57	0.20			
23135100	**Low discrimination**					**23135100**
					COC	
Actinomyces odontolyticus		53.86	0.39	TRE 1	1	
Actinomyces meyeri		32.32	0.32		1	
Atopobium parvulum		5.42	0.16		99	
23135102	**Low discrimination**					**23135102**
Actinomyces odontolyticus		70.52	0.36	TRE 1		
Propionibacterium propionicum		13.51	0.30			
Lactobacillus fermentum		5.80	0.24			
23135200	**Good identification**					**23135200**
Actinomyces odontolyticus		100.0	0.53	TRE 1		
23135202	**Good identification**					**23135202**
Actinomyces odontolyticus		100.0	0.49	TRE 1		
23135220	**Good identification**					**23135220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.39	TRE 1		
23135222	**Good identification**					**23135222**
Actinomyces odontolyticus		100.0	0.36	TRE 1		
23135300	**Good identification**					**23135300**
Actinomyces odontolyticus		100.0	0.44	TRE 1		
23135302	**Good identification**					**23135302**
Actinomyces odontolyticus		100.0	0.40	TRE 1		
23136100	**Very good identification**					**23136100**
Propionibacterium propionicum		99.81	0.61	XYL 1		
Actinomyces odontolyticus		0.20	0.08			
23136101	**Good identification**					**23136101**
Propionibacterium propionicum		100.0	0.45	XYL 1		
23136102	**Very good identification**					**23136102**
Propionibacterium propionicum		100.0	0.64	XYL 1		
23136103	**Good identification**					**23136103**
Propionibacterium propionicum		100.0	0.48	XYL 1		
23136140	**Very good identification**					**23136140**
Propionibacterium propionicum		100.0	0.61	XYL 1		
23136141	**Good identification**					**23136141**
Propionibacterium propionicum		100.0	0.45	XYL 1		
23136142	**Very good identification**					**23136142**
Propionibacterium propionicum		100.0	0.64	XYL 1		
23136143	**Good identification**					**23136143**
Propionibacterium propionicum		100.0	0.48	XYL 1		
23137100	**Very good identification**					**23137100**
Propionibacterium propionicum		98.05	0.61	XYL 1		
Actinomyces israelii		1.95	0.33			
23137101	**Good identification**					**23137101**
Propionibacterium propionicum		100.0	0.45	XYL 1		
23137102	**Very good identification**					**23137102**
Propionibacterium propionicum		98.05	0.64	XYL 1		
Actinomyces israelii		1.95	0.36			
23137103	**Good identification**					**23137103**
Propionibacterium propionicum		100.0	0.48	XYL 1		
23137140	**Very good identification**					**23137140**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium propionicum		99.23	0.61	XYL 1				
Actinomyces israelii		0.77	0.26					
23137141	**Good identification**							**23137141**
Propionibacterium propionicum		100.0	0.45	XYL 1				
23137142	**Very good identification**							**23137142**
Propionibacterium propionicum		99.23	0.64	XYL 1				
Actinomyces israelii		0.77	0.29					
23137143	**Good identification**							**23137143**
Propionibacterium propionicum		100.0	0.48	XYL 1				
23140000	**Acceptable identification**							**23140000**
					COC	SPO	GRA	
Actinomyces odontolyticus		91.71	0.38	FRU 92	1	1	99	
Clostridium botulinum biovar A		3.65	0.06	CEL 1	1	99	99	
Megasphaera elsdenii		3.25	0.10		99	1	1	
23140200	**Good identification**							**23140200**
Actinomyces odontolyticus		100.0	0.43	FRU 92				
23140202	**Good identification**							**23140202**
Actinomyces odontolyticus		100.0	0.39	FRU 92				
23141101	**Good identification**							**23141101**
					SPO			
Clostridium chauvoei		99.35	0.40	MNS 90	99			
Actinomyces naeslundii		0.65	0.26	CEL 1	1			
23141200	**Good identification**							**23141200**
Actinomyces odontolyticus		100.0	0.40	FRU 92				
23141202	**Good identification**							**23141202**
Actinomyces odontolyticus		100.0	0.36	FRU 92				
23144200	**Good identification**							**23144200**
Actinomyces odontolyticus		100.0	0.38	FRU 92				
23150000	**Acceptable identification**							**23150000**
					COC	GRA		
Actinomyces odontolyticus		93.49	0.56	CEL 1	1	99		
Megasphaera elsdenii		5.48	0.32		99	1		
Actinomyces meyeri		0.50	0.13		1	99		
23150002	**Low discrimination**							**23150002**
					COC	GRA		
Actinomyces odontolyticus		61.49	0.52	CEL 1	1	99		
Actinomyces viscosus		37.86	0.42		1	99		
Megasphaera elsdenii		0.65	0.15		99	1		
23150020	**Good identification**							**23150020**
Actinomyces odontolyticus		100.0	0.43	CEL 1				
23150022	**Good identification**							**23150022**
Actinomyces odontolyticus		96.32	0.39	CEL 1				

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		3.68	0.08	NAG 14		
23150100	**Low discrimination**					**23150100**
Actinomyces odontolyticus		88.14	0.47	CEL 1		
Actinomyces meyeri		5.97	0.24			
Propionibacterium propionicum		2.77	0.29			
23150102	**Low discrimination**					**23150102**
Actinomyces viscosus		76.30	0.46	MNS 96		
Actinomyces odontolyticus		22.01	0.44	RAF 94		
Propionibacterium propionicum		1.69	0.32			
23150103	**Good identification**					**23150103**
Actinomyces viscosus		97.36	0.38	MNS 96		
Actinomyces odontolyticus		0.85	0.10	RAF 94		
Propionibacterium propionicum		0.72	0.15			
23150200	**Very good identification**					**23150200**
Actinomyces odontolyticus		100.0	0.61	CEL 1		
23150202	**Very good identification**					**23150202**
Actinomyces odontolyticus		99.68	0.57	CEL 1		
Actinomyces viscosus		0.32	0.08			
23150220	**Good identification**					**23150220**
Actinomyces odontolyticus		100.0	0.47	CEL 1		
23150222	**Good identification**					**23150222**
Actinomyces odontolyticus		100.0	0.44	CEL 1		
23150300	**Good identification**					**23150300**
Actinomyces odontolyticus		100.0	0.52	CEL 1		
23150302	**Good identification**					**23150302**
Actinomyces odontolyticus		98.23	0.49	CEL 1		
Actinomyces viscosus		1.77	0.12			
23150320	**Good identification**					**23150320**
Actinomyces odontolyticus		100.0	0.39	CEL 1		
23150322	**Good identification**					**23150322**
Actinomyces odontolyticus		100.0	0.35	CEL 1		
23151000	**Good identification**					**23151000**
Actinomyces odontolyticus		96.04	0.53	CEL 1		
Actinomyces meyeri		1.15	0.16			
Actinomyces naeslundii		1.10	0.37			
23151001	**Low discrimination**					**23151001**
					SPO	
Actinomyces naeslundii		47.07	0.40	MNS 88	1	
Actinomyces odontolyticus		27.61	0.19	RAF 92	1	
Clostridium chauvoei		25.33	0.12		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23151002	**Low discrimination**					**23151002**
Actinomyces odontolyticus		62.57	0.49	CEL 1		
Actinomyces viscosus		35.42	0.38			
Actinomyces naeslundii		1.17	0.37			
23151020	**Low discrimination**					**23151020**
Actinomyces odontolyticus		85.92	0.40	CEL 1		
Actinomyces naeslundii		6.06	0.37	NAG 14		
Clostridium perfringens		5.72	0.14			
23151021	**Low discrimination**					**23151021**
					SPO	
Actinomyces naeslundii		78.44	0.40	MNS 88	1	
Clostridium chauvoei		14.07	0.04	RAF 92	99	
Actinomyces odontolyticus		7.49	0.05		1	
23151022	**Low discrimination**					**23151022**
Actinomyces odontolyticus		81.39	0.36	CEL 1		
Actinomyces naeslundii		9.37	0.37	NAG 14		
Clostridium perfringens		4.16	0.09			
23151023	**Good identification**					**23151023**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23151100	**Low discrimination**					**23151100**
Actinomyces odontolyticus		69.76	0.44	CEL 1		
Actinomyces meyeri		10.62	0.27			
Actinomyces naeslundii		7.22	0.45			
23151101	**Acceptable identification**					**23151101**
					SPO	
Clostridium chauvoei		94.25	0.47	MNS 90	99	
Actinomyces naeslundii		5.04	0.48	CEL 1	1	
Actinomyces odontolyticus		0.33	0.10		1	
23151102	**Low discrimination**					**23151102**
Actinomyces viscosus		69.45	0.42	MNS 96		
Actinomyces odontolyticus		21.79	0.41	RAF 94		
Actinomyces naeslundii		3.68	0.45			
23151121	**Low discrimination**					**23151121**
					SPO	
Clostridium chauvoei		86.19	0.38	MNS 90	99	
Actinomyces naeslundii		13.82	0.48	CEL 1	1	
23151200	**Very good identification**					**23151200**
Actinomyces odontolyticus		100.0	0.58	CEL 1		
23151202	**Good identification**					**23151202**
Actinomyces odontolyticus		99.71	0.54	CEL 1		
Actinomyces viscosus		0.29	0.04			
23151220	**Good identification**					**23151220**
Actinomyces odontolyticus		100.0	0.44	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23151222	**Good identification**					**23151222**
Actinomyces odontolyticus		100.0	0.41	CEL 1		
23151300	**Good identification**					**23151300**
Actinomyces odontolyticus		100.0	0.49	CEL 1		
23151302	**Good identification**					**23151302**
Actinomyces odontolyticus		98.37	0.46	CEL 1		
Actinomyces viscosus		1.63	0.08			
23151320	**Good identification**					**23151320**
Actinomyces odontolyticus		100.0	0.36	CEL 1		
23152100	**Very good identification**					**23152100**
Propionibacterium propionicum		99.50	0.63	CEL 1		
Actinomyces odontolyticus		0.32	0.13			
Actinomyces naeslundii		0.18	0.26			
23152101	**Good identification**					**23152101**
Propionibacterium propionicum		97.64	0.47	CEL 1		
Actinomyces naeslundii		2.37	0.29	URE 10		
23152102	**Very good identification**					**23152102**
Propionibacterium propionicum		99.29	0.66	CEL 1		
Actinomyces viscosus		0.46	0.12			
Actinomyces odontolyticus		0.13	0.10			
23152103	**Good identification**					**23152103**
Propionibacterium propionicum		97.09	0.50	CEL 1		
Actinomyces naeslundii		1.57	0.29	URE 10		
Actinomyces viscosus		1.34	0.04			
23152140	**Very good identification**					**23152140**
Propionibacterium propionicum		100.0	0.63	CEL 1		
23152141	**Good identification**					**23152141**
Propionibacterium propionicum		100.0	0.47	CEL 1		
23152142	**Very good identification**					**23152142**
Propionibacterium propionicum		100.0	0.66	CEL 1		
23152143	**Good identification**					**23152143**
Propionibacterium propionicum		100.0	0.50	CEL 1		
23153000	**Low discrimination**					**23153000**
Actinomyces naeslundii		45.39	0.42	MNS 88		
Propionibacterium propionicum		30.30	0.29	RAF 92		
Actinomyces odontolyticus		20.57	0.19			
23153001	**Good identification**					**23153001**
Actinomyces naeslundii		95.29	0.45	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		4.71	0.12	RAF 92		
23153003	**Acceptable identification**					**23153003**
Actinomyces naeslundii		93.10	0.45	MNS 88		
Propionibacterium propionicum		6.91	0.15	RAF 92		
23153020	**Acceptable identification**					**23153020**
Actinomyces naeslundii		93.13	0.42	MNS 88		
Actinomyces odontolyticus		6.87	0.05	RAF 92		
23153021	**Good identification**					**23153021**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23153022	**Good identification**					**23153022**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23153023	**Good identification**					**23153023**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23153100	**Very good identification**					**23153100**
Propionibacterium propionicum		95.22	0.63	CEL 1		
Actinomyces naeslundii		4.10	0.50			
Actinomyces israelii		0.48	0.25			
23153101	**Low discrimination**					**23153101**
					SPO	
Propionibacterium propionicum		61.05	0.47	CEL 1	1	
Actinomyces naeslundii		35.49	0.53	URE 10	1	
Clostridium chauvoei		3.46	0.12		99	
23153102	**Very good identification**					**23153102**
Propionibacterium propionicum		96.40	0.66	CEL 1		
Actinomyces naeslundii		2.77	0.50			
Actinomyces israelii		0.48	0.28			
23153103	**Low discrimination**					**23153103**
Propionibacterium propionicum		71.29	0.50	CEL 1		
Actinomyces naeslundii		27.63	0.53	URE 10		
Actinomyces viscosus		0.60	0.00			
23153120	**Low discrimination**					**23153120**
Actinomyces naeslundii		81.00	0.50	MNS 88		
Propionibacterium propionicum		19.00	0.29	RAF 92		
23153121	**Good identification**					**23153121**
					SPO	
Actinomyces naeslundii		95.25	0.53	MNS 88	1	
Clostridium chauvoei		3.09	0.04	RAF 92	99	
Propionibacterium propionicum		1.66	0.12		1	
23153122	**Low discrimination**					**23153122**
Actinomyces naeslundii		73.97	0.50	MNS 88		
Propionibacterium propionicum		26.03	0.32	RAF 92		
23153123	**Good identification**					**23153123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.46	0.53	MNS 88		
Propionibacterium propionicum		2.54	0.15	RAF 92		
23153140	**Very good identification**					**23153140**
Propionibacterium propionicum		99.06	0.63	CEL 1		
Actinomyces naeslundii		0.75	0.37			
Actinomyces israelii		0.19	0.17			
23153141	**Acceptable identification**					**23153141**
Propionibacterium propionicum		90.70	0.47	CEL 1		
Actinomyces naeslundii		9.30	0.40	URE 10		
23153142	**Very good identification**					**23153142**
Propionibacterium propionicum		99.30	0.66	CEL 1		
Actinomyces naeslundii		0.50	0.37			
Actinomyces israelii		0.19	0.20			
23153143	**Acceptable identification**					**23153143**
Propionibacterium propionicum		93.60	0.50	CEL 1		
Actinomyces naeslundii		6.40	0.40	URE 10		
23153161	**Acceptable identification**					**23153161**
Actinomyces naeslundii		91.04	0.40	MNS 88		
Propionibacterium propionicum		8.96	0.12	RAF 92		
23153163	**Low discrimination**					**23153163**
Actinomyces naeslundii		87.13	0.40	MNS 88		
Propionibacterium propionicum		12.87	0.15	RAF 92		
23154000	**Good identification**					**23154000**
Actinomyces odontolyticus		97.94	0.51	CEL 1		
Actinomyces meyeri		2.06	0.18			
23154002	**Good identification**					**23154002**
Actinomyces odontolyticus		98.75	0.47	CEL 1		
Actinomyces viscosus		1.25	0.08			
23154020	**Good identification**					**23154020**
Actinomyces odontolyticus		98.27	0.37	CEL 1		
Actinomyces meyeri		1.73	0.04	NAG 14		
23154100	**Low discrimination**					**23154100**
Actinomyces odontolyticus		78.95	0.42	CEL 1		
Actinomyces meyeri		21.06	0.29			
23154102	**Acceptable identification**					**23154102**
Actinomyces odontolyticus		92.20	0.39	CEL 1		
Actinomyces viscosus		6.56	0.12			
Actinomyces meyeri		1.24	0.03			
23154200	**Good identification**					**23154200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.56	CEL 1		
23154202	**Good identification**					**23154202**
Actinomyces odontolyticus		100.0	0.52	CEL 1		
23154220	**Good identification**					**23154220**
Actinomyces odontolyticus		100.0	0.42	CEL 1		
23154222	**Good identification**					**23154222**
Actinomyces odontolyticus		100.0	0.39	CEL 1		
23154300	**Good identification**					**23154300**
Actinomyces odontolyticus		100.0	0.47	CEL 1		
23154302	**Good identification**					**23154302**
Actinomyces odontolyticus		100.0	0.43	CEL 1		
23155000	**Acceptable identification**					**23155000**
Actinomyces odontolyticus		93.92	0.48	CEL 1		
Actinomyces meyeri		4.45	0.21			
Lactobacillus fermentum		0.92	0.20			
23155002	**Good identification**					**23155002**
Actinomyces odontolyticus		95.60	0.44	CEL 1		
Actinomyces israelii		1.77	0.22			
Lactobacillus fermentum		1.52	0.20			
23155100	**Low discrimination**					**23155100**
Actinomyces odontolyticus		57.26	0.39	CEL 1		
Actinomyces meyeri		34.36	0.32			
Actinomyces israelii		5.49	0.30			
23155102	**Low discrimination**					**23155102**
Actinomyces odontolyticus		70.78	0.36	CEL 1		
Actinomyces israelii		16.62	0.33			
Lactobacillus fermentum		5.83	0.24			
23155200	**Good identification**					**23155200**
Actinomyces odontolyticus		100.0	0.53	CEL 1		
23155202	**Good identification**					**23155202**
Actinomyces odontolyticus		100.0	0.49	CEL 1		
23155220	**Good identification**					**23155220**
Actinomyces odontolyticus		100.0	0.39	CEL 1		
23155222	**Good identification**					**23155222**
Actinomyces odontolyticus		100.0	0.36	CEL 1		
23155300	**Good identification**					**23155300**
Actinomyces odontolyticus		100.0	0.44	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23155302	**Good identification**					**23155302**
Actinomyces odontolyticus		99.10	0.40	CEL 1		
Actinomyces israelii		0.90	0.14			
23157002	**Acceptable identification**					**23157002**
Actinomyces israelii		91.23	0.35	SAL 92		
Actinomyces odontolyticus		8.77	0.10			
23157100	**Low discrimination**					**23157100**
Actinomyces israelii		85.04	0.43	SAL 92		
Propionibacterium propionicum		14.96	0.29			
23157102	**Low discrimination**					**23157102**
Actinomyces israelii		85.04	0.46	SAL 92		
Propionibacterium propionicum		14.96	0.32			
23157140	**Low discrimination**					**23157140**
Actinomyces israelii		68.85	0.36	SAL 92		
Propionibacterium propionicum		31.15	0.29			
23157142	**Low discrimination**					**23157142**
Actinomyces israelii		68.85	0.39	SAL 92		
Propionibacterium propionicum		31.15	0.32			
23157500	**Good identification**					**23157500**
Actinomyces israelii		100.0	0.41	SAL 92		
23157502	**Good identification**					**23157502**
Actinomyces israelii		98.37	0.44	SAL 92		
Bifidobacterium adolescentis		1.63	0.07			
23157542	**Good identification**					**23157542**
Actinomyces israelii		100.0	0.37	SAL 92		
23163101	**Good identification**					**23163101**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23163103	**Low discrimination**					**23163103**
Actinomyces naeslundii		82.73	0.36	MNS 88		
Propionibacterium propionicum		17.27	0.14	RAF 92		
23163121	**Good identification**					**23163121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23163123	**Good identification**					**23163123**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23171000	**Low discrimination**					**23171000**
Actinomyces naeslundii		59.02	0.42	MNS 88		
Actinomyces odontolyticus		26.75	0.19	RAF 92		
Lactobacillus jensenii		14.24	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23171001	**Good identification**					**23171001**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23171003	**Low discrimination**					**23171003**
Actinomyces naeslundii		79.34	0.45	MNS 88		
Actinomyces viscosus		20.66	0.11	RAF 92		
23171020	**Low discrimination**					**23171020**
Actinomyces naeslundii		50.61	0.42	MNS 88		
Clostridium perfringens		45.66	0.19	RAF 92		
Actinomyces odontolyticus		3.73	0.05			
23171021	**Good identification**					**23171021**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23171022	**Low discrimination**					**23171022**
Actinomyces naeslundii		70.20	0.42	MNS 88		
Clostridium perfringens		29.80	0.13	RAF 92		
23171023	**Good identification**					**23171023**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23171100	**Low discrimination**					**23171100**
					COC	
Actinomyces naeslundii		57.21	0.50	MNS 88	1	
Atopobium parvulum		17.74	0.18	RAF 92	99	
Propionibacterium propionicum		10.98	0.27		1	
23171102	**Low discrimination**					**23171102**
Actinomyces naeslundii		44.35	0.50	MNS 88		
Actinomyces viscosus		32.46	0.23	RAF 92		
Propionibacterium propionicum		12.77	0.30			
23171103	**Low discrimination**					**23171103**
					SPO	
Actinomyces naeslundii		72.83	0.53	MNS 88	1	
Clostridium chauvoei		13.77	0.17	RAF 92	99	
Actinomyces viscosus		11.85	0.15		1	
23171121	**Low discrimination**					**23171121**
					SPO	
Actinomyces naeslundii		73.69	0.53	MNS 88	1	
Clostridium chauvoei		26.31	0.22	RAF 92	99	
23171123	**Acceptable identification**					**23171123**
					SPO	
Actinomyces naeslundii		94.07	0.53	MNS 88	1	
Clostridium chauvoei		5.93	0.09	RAF 92	99	
23171141	**Good identification**					**23171141**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23171143	**Low discrimination**					**23171143**
Actinomyces naeslundii		89.22	0.40	MNS 88		
Propionibacterium propionicum		10.78	0.14	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23171161	**Good identification**					**23171161**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23171163	**Good identification**					**23171163**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23172100	**Very good identification**					**23172100**
Propionibacterium propionicum		99.58	0.61	CEL 1		
Actinomyces naeslundii		0.42	0.31			
23172101	**Acceptable identification**					**23172101**
Propionibacterium propionicum		94.57	0.45	CEL 1		
Actinomyces naeslundii		5.43	0.34	URE 10		
23172102	**Very good identification**					**23172102**
Propionibacterium propionicum		99.72	0.64	CEL 1		
Actinomyces naeslundii		0.28	0.31			
23172103	**Good identification**					**23172103**
Propionibacterium propionicum		96.31	0.48	CEL 1		
Actinomyces naeslundii		3.69	0.34	URE 10		
23172140	**Very good identification**					**23172140**
Propionibacterium propionicum		100.0	0.61	CEL 1		
23172141	**Good identification**					**23172141**
Propionibacterium propionicum		100.0	0.45	CEL 1		
23172142	**Very good identification**					**23172142**
Propionibacterium propionicum		100.0	0.64	CEL 1		
23172143	**Good identification**					**23172143**
Propionibacterium propionicum		100.0	0.48	CEL 1		
23173000	**Low discrimination**					**23173000**
Actinomyces naeslundii		69.38	0.47	MNS 88		
Propionibacterium propionicum		19.53	0.27	RAF 92		
Actinomyces israelii		11.09	0.24			
23173001	**Good identification**					**23173001**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23173002	**Low discrimination**					**23173002**
Actinomyces naeslundii		60.17	0.47	MNS 88		
Propionibacterium propionicum		25.40	0.30	RAF 92		
Actinomyces israelii		14.43	0.27			
23173003	**Good identification**					**23173003**
Actinomyces naeslundii		96.97	0.50	MNS 88		
Propionibacterium propionicum		3.03	0.14	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23173020	**Good identification**					**23173020**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23173021	**Good identification**					**23173021**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23173022	**Good identification**					**23173022**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23173023	**Good identification**					**23173023**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23173041	**Good identification**					**23173041**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23173043	**Low discrimination**					**23173043**
Actinomyces naeslundii		84.95	0.37	MNS 88		
Propionibacterium propionicum		15.05	0.14	RAF 92		
23173061	**Good identification**					**23173061**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23173063	**Good identification**					**23173063**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23173100	**Low discrimination**					**23173100**
Propionibacterium propionicum		88.88	0.61	CEL 1		
Actinomyces naeslundii		9.08	0.55			
Actinomyces israelii		2.04	0.34			
23173101	**Low discrimination**					**23173101**
Actinomyces naeslundii		57.97	0.58	MNS 88		
Propionibacterium propionicum		42.03	0.45	RAF 92		
23173102	**Good identification**					**23173102**
Propionibacterium propionicum		91.65	0.64	CEL 1		
Actinomyces naeslundii		6.24	0.55			
Actinomyces israelii		2.10	0.37			
23173103	**Low discrimination**					**23173103**
Propionibacterium propionicum		52.10	0.48	CEL 1		
Actinomyces naeslundii		47.90	0.58	URE 10		
23173111	**Good identification**					**23173111**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23173113	**Low discrimination**					**23173113**
Actinomyces naeslundii		82.73	0.36	MNS 88		
Propionibacterium propionicum		17.27	0.14	RAF 92		
23173120	**Low discrimination**					**23173120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		89.83	0.55	MNS 88		
Propionibacterium propionicum		8.88	0.27	RAF 92		
Actinomyces israelii		1.29	0.14			
23173121	**Very good identification**					**23173121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23173122	**Low discrimination**					**23173122**
Actinomyces naeslundii		85.48	0.55	MNS 88		
Propionibacterium propionicum		12.68	0.30	RAF 92		
Actinomyces israelii		1.84	0.17			
23173123	**Very good identification**					**23173123**
Actinomyces naeslundii		98.91	0.58	MNS 88		
Propionibacterium propionicum		1.09	0.14	RAF 92		
23173131	**Good identification**					**23173131**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23173133	**Good identification**					**23173133**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23173140	**Very good identification**					**23173140**
Propionibacterium propionicum		97.38	0.61	CEL 1		
Actinomyces naeslundii		1.76	0.42			
Actinomyces israelii		0.87	0.27			
23173141	**Low discrimination**					**23173141**
Propionibacterium propionicum		80.43	0.45	CEL 1		
Actinomyces naeslundii		19.57	0.45	URE 10		
23173142	**Very good identification**					**23173142**
Propionibacterium propionicum		97.95	0.64	CEL 1		
Actinomyces naeslundii		1.18	0.42			
Actinomyces israelii		0.87	0.30			
23173143	**Low discrimination**					**23173143**
Propionibacterium propionicum		86.04	0.48	CEL 1		
Actinomyces naeslundii		13.96	0.45	URE 10		
23173160	**Low discrimination**					**23173160**
Actinomyces naeslundii		64.09	0.42	MNS 88		
Propionibacterium propionicum		35.91	0.27	RAF 92		
23173161	**Good identification**					**23173161**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23173162	**Low discrimination**					**23173162**
Actinomyces naeslundii		54.33	0.42	MNS 88		
Propionibacterium propionicum		45.67	0.30	RAF 92		
23173163	**Acceptable identification**					**23173163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		94.14	0.45	MNS 88		
Propionibacterium propionicum		5.86	0.14	RAF 92		
23173502	**Low discrimination**					**23173502**
Actinomyces israelii		65.02	0.36	SAL 92		
Propionibacterium propionicum		34.98	0.30	XYL 92		
23175100	**Low discrimination**					**23175100**
					COC	
Actinomyces israelii		60.58	0.40	SAL 92	1	
Atopobium parvulum		39.42	0.23		99	
23175102	**Acceptable identification**					**23175102**
					COC	
Actinomyces israelii		92.89	0.43	SAL 92	1	
Atopobium parvulum		7.11	0.10		99	
23175142	**Good identification**					**23175142**
Actinomyces israelii		100.0	0.36	SAL 92		
23175500	**Good identification**					**23175500**
Actinomyces israelii		100.0	0.38	SAL 92		
23175502	**Good identification**					**23175502**
Actinomyces israelii		100.0	0.41	SAL 92		
23177000	**Good identification**					**23177000**
Actinomyces israelii		100.0	0.42	SAL 92		
23177002	**Good identification**					**23177002**
Actinomyces israelii		100.0	0.45	SAL 92		
23177040	**Good identification**					**23177040**
Actinomyces israelii		100.0	0.35	SAL 92		
23177042	**Good identification**					**23177042**
Actinomyces israelii		100.0	0.38	SAL 92		
23177100	**Good identification**					**23177100**
Actinomyces israelii		96.31	0.53	SAL 92		
Propionibacterium propionicum		3.69	0.27			
23177102	**Good identification**					**23177102**
Actinomyces israelii		96.31	0.56	SAL 92		
Propionibacterium propionicum		3.69	0.30			
23177110	**Good identification**					**23177110**
Actinomyces israelii		100.0	0.38	SAL 92		
23177112	**Good identification**					**23177112**
Actinomyces israelii		100.0	0.41	SAL 92		
23177122	**Good identification**					**23177122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.35	SAL 92			
23177140	**Acceptable identification**						**23177140**
Actinomyces israelii		91.04	0.45	SAL 92			
Propionibacterium propionicum		8.96	0.27				
23177142	**Acceptable identification**						**23177142**
Actinomyces israelii		91.04	0.49	SAL 92			
Propionibacterium propionicum		8.96	0.30				
23177302	**Good identification**						**23177302**
Actinomyces israelii		100.0	0.36	SAL 92			
23177400	**Good identification**						**23177400**
Actinomyces israelii		100.0	0.41	SAL 92			
23177402	**Good identification**						**23177402**
Actinomyces israelii		100.0	0.44	SAL 92			
23177442	**Good identification**						**23177442**
Actinomyces israelii		100.0	0.37	SAL 92			
23177500	**Good identification**						**23177500**
Actinomyces israelii		100.0	0.51	SAL 92			
23177502	**Good identification**						**23177502**
Actinomyces israelii		100.0	0.54	SAL 92			
23177510	**Good identification**						**23177510**
Actinomyces israelii		100.0	0.37	SAL 92			
23177512	**Good identification**						**23177512**
Actinomyces israelii		100.0	0.40	SAL 92			
23177540	**Good identification**						**23177540**
Actinomyces israelii		100.0	0.44	SAL 92			
23177542	**Good identification**						**23177542**
Actinomyces israelii		100.0	0.47	SAL 92			
23200000	**Low discrimination**						**23200000**
					COC	GRA	
Actinomyces odontolyticus		51.28	0.43	MLT 99	1	99	
Eubacterium tortuosum		41.88	0.30	FRU 92	1	99	
Veillonella parvula		1.83	-.03		99	1	
23200002	**Low discrimination**						**23200002**
Actinomyces odontolyticus		80.15	0.39	MLT 99			
Eubacterium tortuosum		18.85	0.17	FRU 92			
Atopobium minutum		1.01	0.07				
23200200	**Good identification**						**23200200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.58	0.48	MLT 99		
Eubacterium tortuosum		0.42	-.04	FRU 92		
23200202	**Good identification**					**23200202**
Actinomyces odontolyticus		100.0	0.44	MLT 99		
23200300	**Good identification**					**23200300**
Actinomyces odontolyticus		100.0	0.39	MLT 99		
23200302	**Good identification**					**23200302**
Actinomyces odontolyticus		100.0	0.36	MLT 99		
23201000	**Very good identification**					**23201000**
					GRA	
Eubacterium tortuosum		99.17	0.64	ESL 99	99	
Actinomyces odontolyticus		0.82	0.40		99	
Fusobacterium mortiferum		0.01	0.13		1	
23201002	**Good identification**					**23201002**
Eubacterium tortuosum		97.22	0.52	ESL 99		
Actinomyces odontolyticus		2.78	0.36			
23201020	**Good identification**					**23201020**
Eubacterium tortuosum		99.25	0.52	ESL 99		
Actinomyces odontolyticus		0.76	0.26			
23201022	**Good identification**					**23201022**
Eubacterium tortuosum		97.43	0.39	ESL 99		
Actinomyces odontolyticus		2.57	0.23			
23201200	**Low discrimination**					**23201200**
Actinomyces odontolyticus		61.31	0.45	MLT 99		
Eubacterium tortuosum		38.69	0.30	FRU 92		
23201202	**Low discrimination**					**23201202**
Actinomyces odontolyticus		84.62	0.41	MLT 99		
Eubacterium tortuosum		15.38	0.17	FRU 92		
23201300	**Good identification**					**23201300**
Actinomyces odontolyticus		98.02	0.36	MLT 99		
Eubacterium tortuosum		1.98	-.04	FRU 92		
23204000	**Good identification**					**23204000**
Actinomyces odontolyticus		98.35	0.38	MLT 99		
Eubacterium tortuosum		1.65	-.04	FRU 92		
23204200	**Good identification**					**23204200**
Actinomyces odontolyticus		100.0	0.42	MLT 99		
23204202	**Good identification**					**23204202**
Actinomyces odontolyticus		100.0	0.39	MLT 99		
23205200	**Good identification**					**23205200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		98.72	0.39	MLT 99			
Eubacterium tortuosum		1.28	-.04	FRU 92			
23205202	**Good identification**						**23205202**
Actinomyces odontolyticus		100.0	0.36	MLT 99			
23210000	**Good identification**						**23210000**
					COC	GRA	
Actinomyces odontolyticus		93.33	0.61	MLT 99	1	99	
Fusobacterium mortiferum		4.30	0.44		1	1	
Megasphaera elsdenii		1.88	0.29		99	1	
23210002	**Very good identification**						**23210002**
					COC	GRA	
Actinomyces odontolyticus		98.41	0.57	MLT 99	1	99	
Fusobacterium mortiferum		0.82	0.27		1	1	
Megasphaera elsdenii		0.36	0.12		99	1	
23210020	**Good identification**						**23210020**
					CAT		
Actinomyces odontolyticus		98.33	0.47	MLT 99	10		
Propionibacterium acnes		1.67	0.10	NAG 14	95		
23210022	**Good identification**						**23210022**
Actinomyces odontolyticus		100.0	0.44	MLT 99			
23210100	**Low discrimination**						**23210100**
					GRA		
Actinomyces odontolyticus		72.94	0.52	MLT 99	99		
Fusobacterium mortiferum		24.81	0.50		1		
Actinomyces meyeri		2.26	0.23		99		
23210102	**Acceptable identification**						**23210102**
					GRA		
Actinomyces odontolyticus		92.62	0.49	MLT 99	99		
Fusobacterium mortiferum		5.71	0.34		1		
Actinomyces viscosus		1.67	0.12		99		
23210120	**Good identification**						**23210120**
					GRA		
Actinomyces odontolyticus		95.51	0.39	MLT 99	99		
Actinomyces meyeri		2.47	0.08	NAG 14	99		
Fusobacterium mortiferum		2.02	0.16		1		
23210122	**Good identification**						**23210122**
Actinomyces odontolyticus		100.0	0.35	MLT 99			
23210200	**Very good identification**						**23210200**
Actinomyces odontolyticus		100.0	0.66	MLT 99			
23210202	**Very good identification**						**23210202**
Actinomyces odontolyticus		100.0	0.62	MLT 99			
23210220	**Good identification**						**23210220**
Actinomyces odontolyticus		100.0	0.52	MLT 99			
23210222	**Good identification**						**23210222**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.49	MLT 99		
23210300	**Very good identification**					**23210300**
					GRA	
Actinomyces odontolyticus		99.82	0.57	MLT 99	99	
Fusobacterium mortiferum		0.18	0.16		1	
23210302	**Good identification**					**23210302**
Actinomyces odontolyticus		100.0	0.54	MLT 99		
23210320	**Good identification**					**23210320**
Actinomyces odontolyticus		100.0	0.44	MLT 99		
23210322	**Good identification**					**23210322**
Actinomyces odontolyticus		100.0	0.40	MLT 99		
23211000	**Low discrimination**					**23211000**
					GRA	
Actinomyces odontolyticus		82.10	0.58	MLT 99	99	
Eubacterium tortuosum		8.75	0.30		99	
Fusobacterium mortiferum		8.50	0.47		1	
23211002	**Acceptable identification**					**23211002**
					GRA	
Actinomyces odontolyticus		94.67	0.54	MLT 99	99	
Eubacterium tortuosum		2.90	0.17		99	
Fusobacterium mortiferum		1.78	0.30		1	
23211020	**Low discrimination**					**23211020**
					CAT	
Actinomyces odontolyticus		86.68	0.44	MLT 99	10	
Eubacterium tortuosum		10.01	0.17	NAG 14	1	
Propionibacterium acnes		2.21	0.10		95	
23211022	**Good identification**					**23211022**
Actinomyces odontolyticus		95.84	0.41	MLT 99		
Eubacterium tortuosum		3.19	0.04	NAG 14		
Actinomyces naeslundii		0.97	0.24			
23211100	**Low discrimination**					**23211100**
					GRA	
Actinomyces odontolyticus		53.92	0.49	MLT 99	99	
Fusobacterium mortiferum		41.26	0.53		1	
Actinomyces meyeri		3.75	0.26		99	
23211102	**Low discrimination**					**23211102**
					GRA	
Actinomyces odontolyticus		84.60	0.46	MLT 99	99	
Fusobacterium mortiferum		11.74	0.37		1	
Actinomyces viscosus		1.40	0.08		99	
23211120	**Low discrimination**					**23211120**
Actinomyces odontolyticus		82.65	0.36	MLT 99		
Actinomyces meyeri		4.82	0.11	NAG 14		
Actinomyces naeslundii		4.62	0.32			
23211200	**Very good identification**					**23211200**
					GRA	
Actinomyces odontolyticus		99.89	0.63	MLT 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		0.06	-.04		99	
Fusobacterium mortiferum		0.05	0.13		1	
23211202	**Very good identification**					**23211202**
Actinomyces odontolyticus		100.0	0.59	MLT 99		
23211220	**Good identification**					**23211220**
Actinomyces odontolyticus		100.0	0.49	MLT 99		
23211222	**Good identification**					**23211222**
Actinomyces odontolyticus		100.0	0.46	MLT 99		
23211300	**Good identification**					**23211300**
					GRA	
Actinomyces odontolyticus		99.60	0.54	MLT 99	99	
Fusobacterium mortiferum		0.40	0.19		1	
23211302	**Good identification**					**23211302**
Actinomyces odontolyticus		100.0	0.51	MLT 99		
23211320	**Good identification**					**23211320**
Actinomyces odontolyticus		100.0	0.41	MLT 99		
23211322	**Good identification**					**23211322**
Actinomyces odontolyticus		100.0	0.37	MLT 99		
23213101	**Acceptable identification**					**23213101**
Actinomyces naeslundii		90.76	0.40	MNS 88		
Propionibacterium propionicum		9.24	0.12	MLT 94		
23213103	**Low discrimination**					**23213103**
Actinomyces naeslundii		86.76	0.40	MNS 88		
Propionibacterium propionicum		13.24	0.15	MLT 94		
23213120	**Good identification**					**23213120**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23213121	**Good identification**					**23213121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23213122	**Good identification**					**23213122**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23213123	**Good identification**					**23213123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23214000	**Good identification**					**23214000**
					GRA	
Actinomyces odontolyticus		98.36	0.56	MLT 99	99	
Actinomyces meyeri		0.95	0.18		99	
Fusobacterium mortiferum		0.69	0.24		1	
23214002	**Good identification**					**23214002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.52	MLT 99		
23214020	**Good identification**					**23214020**
Actinomyces odontolyticus		99.20	0.42	MLT 99		
Actinomyces meyeri		0.80	0.03	NAG 14		
23214022	**Good identification**					**23214022**
Actinomyces odontolyticus		100.0	0.39	MLT 99		
23214100	**Low discrimination**					**23214100**
					GRA	
Actinomyces odontolyticus		85.19	0.47	MLT 99	99	
Actinomyces meyeri		10.38	0.28		99	
Fusobacterium mortiferum		4.43	0.31		1	
23214102	**Good identification**					**23214102**
					GRA	
Actinomyces odontolyticus		98.47	0.43	MLT 99	99	
Fusobacterium mortiferum		0.93	0.14		1	
Actinomyces meyeri		0.61	0.02		99	
23214200	**Very good identification**					**23214200**
Actinomyces odontolyticus		100.0	0.61	MLT 99		
23214202	**Very good identification**					**23214202**
Actinomyces odontolyticus		100.0	0.57	MLT 99		
23214220	**Good identification**					**23214220**
Actinomyces odontolyticus		100.0	0.47	MLT 99		
23214222	**Good identification**					**23214222**
Actinomyces odontolyticus		100.0	0.43	MLT 99		
23214300	**Good identification**					**23214300**
Actinomyces odontolyticus		100.0	0.52	MLT 99		
23214302	**Good identification**					**23214302**
Actinomyces odontolyticus		100.0	0.48	MLT 99		
23214320	**Good identification**					**23214320**
Actinomyces odontolyticus		100.0	0.39	MLT 99		
23215000	**Good identification**					**23215000**
					GRA	
Actinomyces odontolyticus		96.03	0.53	MLT 99	99	
Actinomyces meyeri		2.08	0.21		99	
Fusobacterium mortiferum		1.52	0.27		1	
23215002	**Good identification**					**23215002**
					GRA	
Actinomyces odontolyticus		99.44	0.49	MLT 99	99	
Fusobacterium mortiferum		0.29	0.11		1	
Lactobacillus fermentum		0.27	0.12		99	
23215020	**Good identification**					**23215020**
Actinomyces odontolyticus		98.22	0.39	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces meyeri		1.78	0.06	NAG 14		
23215022	**Good identification**					**23215022**
Actinomyces odontolyticus		100.0	0.36	MLT 99		
23215100	**Low discrimination**					**23215100**
					GRA	
Actinomyces odontolyticus		71.44	0.44	MLT 99	99	
Actinomyces meyeri		19.59	0.31		99	
Fusobacterium mortiferum		8.35	0.34		1	
23215102	**Good identification**					**23215102**
					GRA	
Actinomyces odontolyticus		95.31	0.40	MLT 99	99	
Fusobacterium mortiferum		2.02	0.17		1	
Lactobacillus fermentum		1.35	0.16		99	
23215200	**Very good identification**					**23215200**
Actinomyces odontolyticus		100.0	0.58	MLT 99		
23215202	**Good identification**					**23215202**
Actinomyces odontolyticus		100.0	0.54	MLT 99		
23215220	**Good identification**					**23215220**
Actinomyces odontolyticus		100.0	0.44	MLT 99		
23215222	**Good identification**					**23215222**
Actinomyces odontolyticus		100.0	0.40	MLT 99		
23215300	**Good identification**					**23215300**
Actinomyces odontolyticus		100.0	0.49	MLT 99		
23215302	**Good identification**					**23215302**
Actinomyces odontolyticus		100.0	0.45	MLT 99		
23215320	**Good identification**					**23215320**
Actinomyces odontolyticus		100.0	0.36	MLT 99		
23230100	**Acceptable identification**					**23230100**
					GRA	
Fusobacterium mortiferum		90.96	0.41	NIT 1	1	
Actinomyces odontolyticus		9.04	0.18		99	
23231000	**Low discrimination**					**23231000**
					GRA	
Fusobacterium mortiferum		70.19	0.38	NIT 1	1	
Actinomyces odontolyticus		22.92	0.24		99	
Actinomyces naeslundii		4.45	0.29		99	
23231100	**Low discrimination**					**23231100**
					GRA	
Fusobacterium mortiferum		89.17	0.44	NIT 1	1	
Actinomyces naeslundii		6.89	0.37		99	
Actinomyces odontolyticus		3.94	0.15		99	
23231101	**Good identification**					**23231101**
Actinomyces naeslundii		100.0	0.40	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23231103	**Good identification**					**23231103**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23231120	**Low discrimination**					**23231120**
Actinomyces naeslundii		54.79	0.37	MNS 88		
Clostridium perfringens		45.21	0.13	MLT 94		
23231121	**Good identification**					**23231121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23231122	**Low discrimination**					**23231122**
Actinomyces naeslundii		72.03	0.37	MNS 88		
Clostridium perfringens		27.97	0.07	MLT 94		
23231123	**Good identification**					**23231123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23233001	**Good identification**					**23233001**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23233003	**Good identification**					**23233003**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23233021	**Good identification**					**23233021**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23233023	**Good identification**					**23233023**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23233100	**Low discrimination**					**23233100**
Actinomyces naeslundii		63.33	0.42	MNS 88		
Propionibacterium propionicum		36.68	0.27	MLT 94		
23233101	**Good identification**					**23233101**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23233102	**Low discrimination**					**23233102**
Actinomyces naeslundii		53.51	0.42	MNS 88		
Propionibacterium propionicum		46.49	0.30	MLT 94		
23233103	**Acceptable identification**					**23233103**
Actinomyces naeslundii		93.95	0.45	MNS 88		
Propionibacterium propionicum		6.05	0.14	MLT 94		
23233120	**Good identification**					**23233120**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23233121	**Good identification**					**23233121**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23233122	**Good identification**					**23233122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	MNS 88		
23233123	**Good identification**					**23233123**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23241000	**Good identification**					**23241000**
Eubacterium tortuosum		99.93	0.48	ESL 99		
Actinomyces odontolyticus		0.08	0.06	CEL 10		
23241002	**Good identification**					**23241002**
Eubacterium tortuosum		100.0	0.35	ESL 99		
23241020	**Good identification**					**23241020**
Eubacterium tortuosum		100.0	0.35	ESL 99		
23250000	**Low discrimination**					**23250000**
					GRA	
Fusobacterium mortiferum		60.30	0.36	NIT 1	1	
Actinomyces odontolyticus		39.70	0.27		99	
23250100	**Acceptable identification**					**23250100**
					GRA	
Fusobacterium mortiferum		91.82	0.42	NIT 1	1	
Actinomyces odontolyticus		8.18	0.18		99	
23251000	**Low discrimination**					**23251000**
					GRA	
Fusobacterium mortiferum		55.14	0.39	NIT 1	1	
Eubacterium tortuosum		18.91	0.14		99	
Actinomyces odontolyticus		16.14	0.24		99	
23251100	**Low discrimination**					**23251100**
					GRA	
Fusobacterium mortiferum		86.27	0.45	NIT 1	1	
Lactobacillus catenaformis		6.24	0.15		99	
Actinomyces naeslundii		4.08	0.34		99	
23251101	**Good identification**					**23251101**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23251103	**Good identification**					**23251103**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23251121	**Good identification**					**23251121**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23251123	**Good identification**					**23251123**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23253100	**Good identification**					**23253100**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23253101	**Good identification**					**23253101**
Actinomyces naeslundii		100.0	0.42	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23253102	**Good identification**					**23253102**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23253103	**Good identification**					**23253103**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23253120	**Good identification**					**23253120**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23253121	**Good identification**					**23253121**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23253122	**Good identification**					**23253122**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23253123	**Good identification**					**23253123**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23271100	**Low discrimination**					**23271100**
					GRA	
Fusobacterium mortiferum		76.49	0.36	NIT 1	1	
Actinomyces naeslundii		23.51	0.39		99	
23271101	**Good identification**					**23271101**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23271102	**Low discrimination**					**23271102**
					GRA	
Actinomyces naeslundii		49.19	0.39	MNS 88	99	
Lactobacillus acidophilus		33.03	0.11	MLT 94	99	
Fusobacterium mortiferum		17.78	0.20	RAF 92	1	
23271103	**Good identification**					**23271103**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23271120	**Good identification**					**23271120**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23271121	**Good identification**					**23271121**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23271122	**Low discrimination**					**23271122**
Actinomyces naeslundii		59.83	0.39	MNS 88		
Lactobacillus acidophilus		40.17	0.11	MLT 94		
23271123	**Good identification**					**23271123**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23273001	**Good identification**					**23273001**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23273003	**Good identification**					**23273003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	MNS 88		
23273020	**Good identification**					**23273020**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23273021	**Good identification**					**23273021**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23273022	**Good identification**					**23273022**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23273023	**Good identification**					**23273023**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23273100	**Low discrimination**					**23273100**
Actinomyces naeslundii		89.66	0.44	MNS 88		
Actinomyces israelii		10.34	0.18	MLT 94		
23273101	**Good identification**					**23273101**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23273102	**Low discrimination**					**23273102**
Actinomyces naeslundii		85.25	0.44	MNS 88		
Actinomyces israelii		14.75	0.21	MLT 94		
23273103	**Good identification**					**23273103**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23273120	**Good identification**					**23273120**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23273121	**Good identification**					**23273121**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23273122	**Good identification**					**23273122**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23273123	**Good identification**					**23273123**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23277100	**Good identification**					**23277100**
Actinomyces israelii		100.0	0.37	MLT 99		
23277102	**Good identification**					**23277102**
Actinomyces israelii		100.0	0.40	MLT 99		
23277500	**Good identification**					**23277500**
Actinomyces israelii		100.0	0.35	MLT 99		
23277502	**Good identification**					**23277502**
Actinomyces israelii		100.0	0.38	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
23300000	**Very good identification**							**23300000**
					COC	SPO	GRA	
Actinomyces odontolyticus		99.93	0.77	FRU 92	1	1	99	
Clostridium botulinum biovar A		0.02	0.06		1	99	99	
Megasphaera elsdenii		0.02	0.10		99	1	1	
23300001	**Good identification**							**23300001**
					SPO			
Actinomyces odontolyticus		98.91	0.43	FRU 92	1			
Clostridium novyi biovar A		1.09	0.14	URE 1	99			
23300002	**Very good identification**							**23300002**
Actinomyces odontolyticus		99.81	0.73	FRU 92				
Actinomyces viscosus		0.19	0.20					
23300003	**Acceptable identification**							**23300003**
Actinomyces odontolyticus		93.99	0.39	FRU 92				
Actinomyces viscosus		6.02	0.12	URE 1				
23300010	**Good identification**							**23300010**
					SPO			
Actinomyces odontolyticus		98.91	0.43	FRU 92	1			
Clostridium novyi biovar A		1.09	0.14	MLZ 1	99			
23300012	**Good identification**							**23300012**
Actinomyces odontolyticus		100.0	0.39	FRU 92				
23300020	**Very good identification**							**23300020**
Actinomyces odontolyticus		100.0	0.63	FRU 92				
23300022	**Very good identification**							**23300022**
Actinomyces odontolyticus		100.0	0.60	FRU 92				
23300040	**Good identification**							**23300040**
					COC	SPO	GRA	
Actinomyces odontolyticus		98.09	0.43	FRU 92	1	1	99	
Megasphaera elsdenii		1.20	0.07	SOR 1	99	1	1	
Clostridium novyi biovar A		0.72	0.11		1	99	99	
23300042	**Good identification**							**23300042**
Actinomyces odontolyticus		100.0	0.39	FRU 92				
23300100	**Very good identification**							**23300100**
Actinomyces odontolyticus		99.98	0.68	FRU 92				
Actinomyces meyeri		0.02	0.02					
23300102	**Very good identification**							**23300102**
Actinomyces odontolyticus		98.92	0.65	FRU 92				
Actinomyces viscosus		1.08	0.24					
23300120	**Good identification**							**23300120**
Actinomyces odontolyticus		100.0	0.55	FRU 92				
23300122	**Good identification**							**23300122**
Actinomyces odontolyticus		100.0	0.51	FRU 92				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23300200	**Very good identification**					**23300200**
Actinomyces odontolyticus		100.0	0.82	FRU 92		
23300201	**Good identification**					**23300201**
Actinomyces odontolyticus		100.0	0.48	FRU 92		
23300202	**Very good identification**					**23300202**
Actinomyces odontolyticus		100.0	0.78	FRU 92		
23300203	**Good identification**					**23300203**
Actinomyces odontolyticus		100.0	0.44	FRU 92		
23300210	**Good identification**					**23300210**
Actinomyces odontolyticus		100.0	0.48	FRU 92		
23300212	**Good identification**					**23300212**
Actinomyces odontolyticus		100.0	0.44	FRU 92		
23300220	**Very good identification**					**23300220**
Actinomyces odontolyticus		100.0	0.68	FRU 92		
23300222	**Very good identification**					**23300222**
Actinomyces odontolyticus		100.0	0.65	FRU 92		
23300240	**Good identification**					**23300240**
Actinomyces odontolyticus		100.0	0.48	FRU 92		
23300242	**Good identification**					**23300242**
Actinomyces odontolyticus		100.0	0.44	FRU 92		
23300300	**Very good identification**					**23300300**
Actinomyces odontolyticus		100.0	0.73	FRU 92		
23300301	**Good identification**					**23300301**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23300302	**Very good identification**					**23300302**
Actinomyces odontolyticus		100.0	0.70	FRU 92		
23300303	**Good identification**					**23300303**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23300310	**Good identification**					**23300310**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23300312	**Good identification**					**23300312**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23300320	**Very good identification**					**23300320**
Actinomyces odontolyticus		100.0	0.60	FRU 92		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23300322	**Very good identification**					**23300322**
Actinomyces odontolyticus		100.0	0.56	FRU 92		
23300340	**Good identification**					**23300340**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23300342	**Good identification**					**23300342**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23300400	**Good identification**					**23300400**
Actinomyces odontolyticus		100.0	0.43	FRU 92		
23300402	**Good identification**					**23300402**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23300600	**Good identification**					**23300600**
Actinomyces odontolyticus		100.0	0.48	FRU 92		
23300602	**Good identification**					**23300602**
Actinomyces odontolyticus		100.0	0.44	FRU 92		
23300700	**Good identification**					**23300700**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23300702	**Good identification**					**23300702**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23301000	**Very good identification**					**23301000**
Actinomyces odontolyticus		98.77	0.74	FRU 92		
Eubacterium tortuosum		1.22	0.30			
Actinomyces naeslundii		0.01	0.22			
23301001	**Good identification**					**23301001**
					SPO	
Actinomyces odontolyticus		95.21	0.40	FRU 92	1	
Clostridium chauvoei		2.24	0.06	URE 1	99	
Actinomyces naeslundii		1.37	0.25		1	
23301002	**Very good identification**					**23301002**
Actinomyces odontolyticus		99.45	0.70	FRU 92		
Eubacterium tortuosum		0.35	0.17			
Actinomyces viscosus		0.18	0.16			
23301003	**Acceptable identification**					**23301003**
Actinomyces odontolyticus		92.40	0.36	FRU 92		
Actinomyces viscosus		5.44	0.08	URE 1		
Actinomyces naeslundii		2.17	0.25			
23301010	**Good identification**					**23301010**
Actinomyces odontolyticus		98.78	0.40	FRU 92		
Eubacterium tortuosum		1.22	-.04	MLZ 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23301012	**Good identification**					**23301012**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23301020	**Very good identification**					**23301020**
Actinomyces odontolyticus		98.62	0.60	FRU 92		
Eubacterium tortuosum		1.32	0.17	NAG 14		
Actinomyces naeslundii		0.06	0.22			
23301022	**Very good identification**					**23301022**
Actinomyces odontolyticus		99.52	0.57	FRU 92		
Eubacterium tortuosum		0.38	0.04	NAG 14		
Actinomyces naeslundii		0.10	0.22			
23301040	**Good identification**					**23301040**
Actinomyces odontolyticus		98.78	0.40	FRU 92		
Eubacterium tortuosum		1.22	-.04	SOR 1		
23301042	**Good identification**					**23301042**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23301100	**Very good identification**					**23301100**
					SPO	
Actinomyces odontolyticus		99.75	0.65	FRU 92	1	
Actinomyces naeslundii		0.09	0.30		1	
Clostridium chauvoei		0.08	0.06		99	
23301101	**Low discrimination**					**23301101**
					SPO	
Clostridium chauvoei		86.72	0.40	MNS 90	99	
Actinomyces odontolyticus		11.76	0.31	SAL 1	1	
Actinomyces naeslundii		1.52	0.33		1	
23301102	**Very good identification**					**23301102**
Actinomyces odontolyticus		98.87	0.62	FRU 92		
Actinomyces viscosus		0.99	0.21			
Actinomyces naeslundii		0.14	0.30			
23301120	**Good identification**					**23301120**
Actinomyces odontolyticus		98.92	0.52	FRU 92		
Clostridium perfringens		0.55	0.08	NAG 14		
Actinomyces naeslundii		0.53	0.30			
23301122	**Good identification**					**23301122**
Actinomyces odontolyticus		98.71	0.48	FRU 92		
Actinomyces naeslundii		0.86	0.30	NAG 14		
Clostridium perfringens		0.43	0.03			
23301200	**Very good identification**					**23301200**
Actinomyces odontolyticus		99.99	0.79	FRU 92		
Eubacterium tortuosum		0.01	-.04			
23301201	**Good identification**					**23301201**
Actinomyces odontolyticus		100.0	0.45	FRU 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23301202	**Very good identification**					**23301202**
Actinomyces odontolyticus		100.0	0.75	FRU 92		
23301203	**Good identification**					**23301203**
Actinomyces odontolyticus		100.0	0.41	FRU 92		
23301210	**Good identification**					**23301210**
Actinomyces odontolyticus		100.0	0.45	FRU 92		
23301212	**Good identification**					**23301212**
Actinomyces odontolyticus		100.0	0.41	FRU 92		
23301220	**Very good identification**					**23301220**
Actinomyces odontolyticus		100.0	0.65	FRU 92		
23301222	**Very good identification**					**23301222**
Actinomyces odontolyticus		100.0	0.62	FRU 92		
23301240	**Good identification**					**23301240**
Actinomyces odontolyticus		100.0	0.45	FRU 92		
23301242	**Good identification**					**23301242**
Actinomyces odontolyticus		100.0	0.41	FRU 92		
23301300	**Very good identification**					**23301300**
Actinomyces odontolyticus		100.0	0.70	FRU 92		
23301301	**Good identification**					**23301301**
					SPO	
Actinomyces odontolyticus		96.30	0.36	FRU 92	1	
Clostridium chauvoei		3.70	0.06	URE 1	99	
23301302	**Very good identification**					**23301302**
Actinomyces odontolyticus		100.0	0.67	FRU 92		
23301310	**Good identification**					**23301310**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23301320	**Very good identification**					**23301320**
Actinomyces odontolyticus		100.0	0.57	FRU 92		
23301322	**Good identification**					**23301322**
Actinomyces odontolyticus		100.0	0.53	FRU 92		
23301340	**Good identification**					**23301340**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23301400	**Good identification**					**23301400**
Actinomyces odontolyticus		98.78	0.40	FRU 92		
Eubacterium tortuosum		1.22	-.04	ARA 1		
23301402	**Good identification**					**23301402**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.36	FRU 92			
23301600	**Good identification**						**23301600**
Actinomyces odontolyticus		100.0	0.45	FRU 92			
23301602	**Good identification**						**23301602**
Actinomyces odontolyticus		100.0	0.41	FRU 92			
23301700	**Good identification**						**23301700**
Actinomyces odontolyticus		100.0	0.36	FRU 92			
23302000	**Good identification**						**23302000**
					COC	GRA	
Actinomyces odontolyticus		98.21	0.43	FRU 92	1	99	
Megasphaera elsdenii		1.79	0.10	MAN 1	99	1	
23302002	**Good identification**						**23302002**
Actinomyces odontolyticus		100.0	0.39	FRU 92			
23302200	**Good identification**						**23302200**
Actinomyces odontolyticus		100.0	0.48	FRU 92			
23302202	**Good identification**						**23302202**
Actinomyces odontolyticus		100.0	0.44	FRU 92			
23302300	**Good identification**						**23302300**
Actinomyces odontolyticus		100.0	0.39	FRU 92			
23302302	**Good identification**						**23302302**
Actinomyces odontolyticus		100.0	0.36	FRU 92			
23303000	**Good identification**						**23303000**
Actinomyces odontolyticus		97.00	0.40	FRU 92			
Actinomyces naeslundii		1.80	0.27	MAN 1			
Eubacterium tortuosum		1.20	-.04				
23303002	**Good identification**						**23303002**
Actinomyces odontolyticus		97.06	0.36	FRU 92			
Actinomyces naeslundii		2.94	0.27	MAN 1			
23303101	**Low discrimination**						**23303101**
					SPO		
Actinomyces naeslundii		70.43	0.38	MNS 88	1		
Clostridium chauvoei		20.88	0.06	RAF 92	99		
Propionibacterium propionicum		8.69	0.12	FRU 95	1		
23303103	**Low discrimination**						**23303103**
Actinomyces naeslundii		84.38	0.38	MNS 88			
Propionibacterium propionicum		15.62	0.15	RAF 92			
23303120	**Low discrimination**						**23303120**
Actinomyces naeslundii		50.72	0.35	MNS 88			
Actinomyces odontolyticus		49.28	0.18	RAF 92			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23303121	**Good identification**					**23303121**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23303122	**Low discrimination**					**23303122**
Actinomyces naeslundii		62.68	0.35	MNS 88		
Actinomyces odontolyticus		37.33	0.14	RAF 92		
23303123	**Good identification**					**23303123**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23303200	**Good identification**					**23303200**
Actinomyces odontolyticus		100.0	0.45	FRU 92		
23303202	**Good identification**					**23303202**
Actinomyces odontolyticus		100.0	0.41	FRU 92		
23303300	**Good identification**					**23303300**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23304000	**Very good identification**					**23304000**
Actinomyces odontolyticus		100.0	0.72	FRU 92		
23304001	**Good identification**					**23304001**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23304002	**Very good identification**					**23304002**
Actinomyces odontolyticus		100.0	0.68	FRU 92		
23304010	**Good identification**					**23304010**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23304020	**Very good identification**					**23304020**
Actinomyces odontolyticus		100.0	0.58	FRU 92		
23304022	**Good identification**					**23304022**
Actinomyces odontolyticus		100.0	0.55	FRU 92		
23304040	**Good identification**					**23304040**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23304100	**Very good identification**					**23304100**
Actinomyces odontolyticus		99.92	0.63	FRU 92		
Actinomyces meyeri		0.08	0.07			
23304102	**Very good identification**					**23304102**
Actinomyces odontolyticus		100.0	0.59	FRU 92		
23304120	**Good identification**					**23304120**
Actinomyces odontolyticus		100.0	0.50	FRU 92		
23304122	**Good identification**					**23304122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.46	FRU 92		
23304200	**Very good identification**					**23304200**
Actinomyces odontolyticus		100.0	0.77	FRU 92		
23304201	**Good identification**					**23304201**
Actinomyces odontolyticus		100.0	0.42	FRU 92		
23304202	**Very good identification**					**23304202**
Actinomyces odontolyticus		100.0	0.73	FRU 92		
23304203	**Good identification**					**23304203**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23304210	**Good identification**					**23304210**
Actinomyces odontolyticus		100.0	0.42	FRU 92		
23304212	**Good identification**					**23304212**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23304220	**Very good identification**					**23304220**
Actinomyces odontolyticus		100.0	0.63	FRU 92		
23304222	**Very good identification**					**23304222**
Actinomyces odontolyticus		100.0	0.59	FRU 92		
23304240	**Good identification**					**23304240**
Actinomyces odontolyticus		100.0	0.42	FRU 92		
23304242	**Good identification**					**23304242**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23304300	**Very good identification**					**23304300**
Actinomyces odontolyticus		100.0	0.68	FRU 92		
23304302	**Very good identification**					**23304302**
Actinomyces odontolyticus		100.0	0.64	FRU 92		
23304320	**Good identification**					**23304320**
Actinomyces odontolyticus		100.0	0.55	FRU 92		
23304322	**Good identification**					**23304322**
Actinomyces odontolyticus		100.0	0.51	FRU 92		
23304400	**Good identification**					**23304400**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23304600	**Good identification**					**23304600**
Actinomyces odontolyticus		100.0	0.42	FRU 92		
23304602	**Good identification**					**23304602**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23305000	**Very good identification**					**23305000**
Actinomyces odontolyticus		99.98	0.69	FRU 92		
Eubacterium tortuosum		0.03	-.04			
23305002	**Very good identification**					**23305002**
Actinomyces odontolyticus		100.0	0.65	FRU 92		
23305020	**Good identification**					**23305020**
Actinomyces odontolyticus		100.0	0.55	FRU 92		
23305022	**Good identification**					**23305022**
Actinomyces odontolyticus		100.0	0.52	FRU 92		
23305100	**Very good identification**					**23305100**
Actinomyces odontolyticus		99.81	0.60	FRU 92		
Actinomyces meyeri		0.19	0.10			
23305102	**Very good identification**					**23305102**
Actinomyces odontolyticus		100.0	0.56	FRU 92		
23305120	**Good identification**					**23305120**
Actinomyces odontolyticus		100.0	0.47	FRU 92		
23305122	**Good identification**					**23305122**
Actinomyces odontolyticus		100.0	0.43	FRU 92		
23305200	**Very good identification**					**23305200**
Actinomyces odontolyticus		100.0	0.74	FRU 92		
23305201	**Good identification**					**23305201**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23305202	**Very good identification**					**23305202**
Actinomyces odontolyticus		100.0	0.70	FRU 92		
23305203	**Good identification**					**23305203**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23305210	**Good identification**					**23305210**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23305212	**Good identification**					**23305212**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23305220	**Very good identification**					**23305220**
Actinomyces odontolyticus		100.0	0.60	FRU 92		
23305222	**Very good identification**					**23305222**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.56	FRU 92			
23305240	**Good identification**						**23305240**
Actinomyces odontolyticus		100.0	0.39	FRU 92			
23305242	**Good identification**						**23305242**
Actinomyces odontolyticus		100.0	0.36	FRU 92			
23305300	**Very good identification**						**23305300**
Actinomyces odontolyticus		100.0	0.65	FRU 92			
23305302	**Very good identification**						**23305302**
Actinomyces odontolyticus		100.0	0.61	FRU 92			
23305320	**Good identification**						**23305320**
Actinomyces odontolyticus		100.0	0.52	FRU 92			
23305322	**Good identification**						**23305322**
Actinomyces odontolyticus		100.0	0.48	FRU 92			
23305600	**Good identification**						**23305600**
Actinomyces odontolyticus		100.0	0.39	FRU 92			
23305602	**Good identification**						**23305602**
Actinomyces odontolyticus		100.0	0.36	FRU 92			
23306000	**Good identification**						**23306000**
Actinomyces odontolyticus		100.0	0.38	FRU 92			
23306200	**Good identification**						**23306200**
Actinomyces odontolyticus		100.0	0.42	FRU 92			
23306202	**Good identification**						**23306202**
Actinomyces odontolyticus		100.0	0.39	FRU 92			
23307200	**Good identification**						**23307200**
Actinomyces odontolyticus		100.0	0.39	FRU 92			
23307202	**Good identification**						**23307202**
Actinomyces odontolyticus		100.0	0.36	FRU 92			
23310000	**Excellent identification**						**23310000**
					COC	GRA	
Actinomyces odontolyticus		99.96	0.95		1	99	
Megasphaera elsdenii		0.03	0.32		99	1	
Fusobacterium mortiferum		0.01	0.28		1	1	
23310001	**Very good identification**						**23310001**
Actinomyces odontolyticus		99.87	0.61	URE 1			
Actinomyces naeslundii		0.07	0.24				
Actinomyces viscosus		0.07	-.01				
23310002	**Excellent identification**						**23310002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Actinomyces odontolyticus		99.67	0.91		1	99	
Actinomyces viscosus		0.32	0.42		1	99	
Megasphaera elsdenii		0.01	0.15		99	1	
23310003	**Good identification**						**23310003**
Actinomyces odontolyticus		90.35	0.57	URE 1			
Actinomyces viscosus		9.55	0.34				
Actinomyces naeslundii		0.10	0.24				
23310010	**Very good identification**						**23310010**
Actinomyces odontolyticus		100.0	0.61	MLZ 1			
23310012	**Very good identification**						**23310012**
Actinomyces odontolyticus		99.68	0.57	MLZ 1			
Actinomyces viscosus		0.32	0.08				
23310020	**Very good identification**						**23310020**
Actinomyces odontolyticus		100.0	0.82	NAG 14			
23310021	**Good identification**						**23310021**
Actinomyces odontolyticus		99.60	0.47	NAG 14			
Actinomyces naeslundii		0.40	0.24	URE 1			
23310022	**Very good identification**						**23310022**
Actinomyces odontolyticus		99.98	0.78	NAG 14			
Actinomyces viscosus		0.02	0.08				
23310023	**Good identification**						**23310023**
Actinomyces odontolyticus		98.70	0.44	NAG 14			
Actinomyces naeslundii		0.65	0.24	URE 1			
Actinomyces viscosus		0.65	-.01				
23310030	**Good identification**						**23310030**
Actinomyces odontolyticus		100.0	0.47	MLZ 1			
23310032	**Good identification**						**23310032**
Actinomyces odontolyticus		100.0	0.44	MLZ 1			
23310040	**Very good identification**						**23310040**
					COC	GRA	
Actinomyces odontolyticus		98.03	0.61	SOR 1	1	99	
Megasphaera elsdenii		1.97	0.29		99	1	
23310042	**Very good identification**						**23310042**
					COC	GRA	
Actinomyces odontolyticus		99.32	0.57	SOR 1	1	99	
Megasphaera elsdenii		0.36	0.12		99	1	
Actinomyces viscosus		0.32	0.08		1	99	
23310060	**Good identification**						**23310060**
Actinomyces odontolyticus		100.0	0.47	NAG 14			
23310062	**Good identification**						**23310062**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.44	NAG 14		
23310100	**Very good identification**					**23310100**
					GRA	
Actinomyces odontolyticus		99.89	0.87		99	
Fusobacterium mortiferum		0.04	0.34		1	
Actinomyces meyeri		0.04	0.24		99	
23310101	**Good identification**					**23310101**
					SPO	
Actinomyces odontolyticus		97.90	0.52	URE 1	1	
Clostridium chauvoei		0.99	0.12		99	
Actinomyces naeslundii		0.58	0.32		1	
23310102	**Very good identification**					**23310102**
Actinomyces odontolyticus		98.18	0.83			
Actinomyces viscosus		1.77	0.46			
Propionibacterium propionicum		0.04	0.32			
23310103	**Low discrimination**					**23310103**
Actinomyces odontolyticus		62.03	0.49	URE 1		
Actinomyces viscosus		36.92	0.38			
Actinomyces naeslundii		0.60	0.32			
23310110	**Good identification**					**23310110**
Actinomyces odontolyticus		100.0	0.52	MLZ 1		
23310112	**Good identification**					**23310112**
Actinomyces odontolyticus		98.23	0.49	MLZ 1		
Actinomyces viscosus		1.77	0.12			
23310120	**Very good identification**					**23310120**
Actinomyces odontolyticus		99.91	0.73	NAG 14		
Clostridium perfringens		0.03	0.08			
Actinomyces meyeri		0.03	0.09			
23310121	**Acceptable identification**					**23310121**
					SPO	
Actinomyces odontolyticus		94.60	0.39	NAG 14	1	
Actinomyces naeslundii		3.45	0.32	URE 1	1	
Clostridium chauvoei		1.95	0.04		99	
23310122	**Very good identification**					**23310122**
Actinomyces odontolyticus		99.82	0.69	NAG 14		
Actinomyces viscosus		0.11	0.12			
Actinomyces naeslundii		0.04	0.28			
23310123	**Acceptable identification**					**23310123**
Actinomyces odontolyticus		91.20	0.35	NAG 14		
Actinomyces naeslundii		5.43	0.32	URE 1		
Actinomyces viscosus		3.37	0.04			
23310130	**Good identification**					**23310130**
Actinomyces odontolyticus		100.0	0.39	MLZ 1		
23310132	**Good identification**					**23310132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.35	MLZ 1		
23310140	**Good identification**					**23310140**
Actinomyces odontolyticus		98.41	0.52	SOR 1		
Propionibacterium propionicum		1.59	0.29			
23310142	**Acceptable identification**					**23310142**
Actinomyces odontolyticus		94.55	0.49	SOR 1		
Propionibacterium propionicum		3.75	0.32			
Actinomyces viscosus		1.71	0.12			
23310160	**Good identification**					**23310160**
Actinomyces odontolyticus		100.0	0.39	NAG 14		
23310162	**Good identification**					**23310162**
Actinomyces odontolyticus		100.0	0.35	NAG 14		
23310200	**Excellent identification**					**23310200**
Actinomyces odontolyticus		100.0	1.00			
23310201	**Very good identification**					**23310201**
Actinomyces odontolyticus		100.0	0.66	URE 1		
23310202	**Excellent identification**					**23310202**
Actinomyces odontolyticus		100.0	0.96			
Actinomyces viscosus		0.00	0.08			
23310203	**Very good identification**					**23310203**
Actinomyces odontolyticus		99.95	0.62	URE 1		
Actinomyces viscosus		0.06	-.01			
23310210	**Very good identification**					**23310210**
Actinomyces odontolyticus		100.0	0.66	MLZ 1		
23310212	**Very good identification**					**23310212**
Actinomyces odontolyticus		100.0	0.62	MLZ 1		
23310220	**Very good identification**					**23310220**
Actinomyces odontolyticus		100.0	0.87	NAG 14		
23310221	**Good identification**					**23310221**
Actinomyces odontolyticus		100.0	0.52	NAG 14		
23310222	**Very good identification**					**23310222**
Actinomyces odontolyticus		100.0	0.83	NAG 14		
23310223	**Good identification**					**23310223**
Actinomyces odontolyticus		100.0	0.49	NAG 14		
23310230	**Good identification**					**23310230**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.52	MLZ 1		
23310232	**Good identification**					**23310232**
Actinomyces odontolyticus		100.0	0.49	MLZ 1		
23310240	**Very good identification**					**23310240**
Actinomyces odontolyticus		100.0	0.66	SOR 1		
23310242	**Very good identification**					**23310242**
Actinomyces odontolyticus		100.0	0.62	SOR 1		
23310260	**Good identification**					**23310260**
Actinomyces odontolyticus		100.0	0.52	NAG 14		
23310262	**Good identification**					**23310262**
Actinomyces odontolyticus		100.0	0.49	NAG 14		
23310300	**Excellent identification**					**23310300**
Actinomyces odontolyticus		100.0	0.91			
23310301	**Very good identification**					**23310301**
Actinomyces odontolyticus		100.0	0.57	URE 1		
23310302	**Very good identification**					**23310302**
Actinomyces odontolyticus		99.99	0.88			
Actinomyces viscosus		0.01	0.12			
23310303	**Good identification**					**23310303**
Actinomyces odontolyticus		99.69	0.54	URE 1		
Actinomyces viscosus		0.31	0.04			
23310310	**Very good identification**					**23310310**
Actinomyces odontolyticus		100.0	0.57	MLZ 1		
23310312	**Good identification**					**23310312**
Actinomyces odontolyticus		100.0	0.54	MLZ 1		
23310320	**Very good identification**					**23310320**
Actinomyces odontolyticus		100.0	0.78	NAG 14		
23310321	**Good identification**					**23310321**
Actinomyces odontolyticus		100.0	0.44	NAG 14		
23310322	**Very good identification**					**23310322**
Actinomyces odontolyticus		100.0	0.74	NAG 14		
23310323	**Good identification**					**23310323**
Actinomyces odontolyticus		100.0	0.40	NAG 14		
23310330	**Good identification**					**23310330**
Actinomyces odontolyticus		100.0	0.44	MLZ 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23310332	**Good identification**					**23310332**
Actinomyces odontolyticus		100.0	0.40	MLZ 1		
23310340	**Very good identification**					**23310340**
Actinomyces odontolyticus		100.0	0.57	SOR 1		
23310342	**Good identification**					**23310342**
Actinomyces odontolyticus		100.0	0.54	SOR 1		
23310360	**Good identification**					**23310360**
Actinomyces odontolyticus		100.0	0.44	NAG 14		
23310362	**Good identification**					**23310362**
Actinomyces odontolyticus		100.0	0.40	NAG 14		
23310400	**Very good identification**					**23310400**
Actinomyces odontolyticus		99.93	0.61	ARA 1		
Actinomyces meyeri		0.07	0.03			
23310402	**Very good identification**					**23310402**
Actinomyces odontolyticus		99.68	0.57	ARA 1		
Actinomyces viscosus		0.32	0.08			
23310420	**Good identification**					**23310420**
Actinomyces odontolyticus		100.0	0.47	ARA 1		
23310422	**Good identification**					**23310422**
Actinomyces odontolyticus		100.0	0.44	ARA 1		
23310500	**Good identification**					**23310500**
Actinomyces odontolyticus		99.14	0.52	ARA 1		
Actinomyces meyeri		0.86	0.13			
23310502	**Good identification**					**23310502**
Actinomyces odontolyticus		98.23	0.49	ARA 1		
Actinomyces viscosus		1.77	0.12			
23310520	**Good identification**					**23310520**
Actinomyces odontolyticus		100.0	0.39	ARA 1		
23310522	**Good identification**					**23310522**
Actinomyces odontolyticus		100.0	0.35	ARA 1		
23310600	**Very good identification**					**23310600**
Actinomyces odontolyticus		100.0	0.66	ARA 1		
23310602	**Very good identification**					**23310602**
Actinomyces odontolyticus		100.0	0.62	ARA 1		
23310620	**Good identification**					**23310620**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.52	ARA 1		
23310622	**Good identification**					**23310622**
Actinomyces odontolyticus		100.0	0.49	ARA 1		
23310700	**Very good identification**					**23310700**
Actinomyces odontolyticus		100.0	0.57	ARA 1		
23310702	**Good identification**					**23310702**
Actinomyces odontolyticus		100.0	0.54	ARA 1		
23310720	**Good identification**					**23310720**
Actinomyces odontolyticus		100.0	0.44	ARA 1		
23310722	**Good identification**					**23310722**
Actinomyces odontolyticus		100.0	0.40	ARA 1		
23311000	**Excellent identification**					**23311000**
					GRA	
Actinomyces odontolyticus		99.96	0.92		99	
Actinomyces naeslundii		0.02	0.44		99	
Fusobacterium mortiferum		0.01	0.31		1	
23311001	**Very good identification**					**23311001**
					SPO	
Actinomyces odontolyticus		97.23	0.58	URE 1	1	
Actinomyces naeslundii		2.31	0.47		1	
Clostridium chauvoei		0.46	0.12		99	
23311002	**Very good identification**					**23311002**
Actinomyces odontolyticus		99.67	0.88			
Actinomyces viscosus		0.29	0.38			
Actinomyces naeslundii		0.03	0.44			
23311003	**Low discrimination**					**23311003**
Actinomyces odontolyticus		87.93	0.54	URE 1		
Actinomyces viscosus		8.55	0.30			
Actinomyces naeslundii		3.40	0.47			
23311010	**Very good identification**					**23311010**
Actinomyces odontolyticus		99.92	0.58	MLZ 1		
Actinomyces naeslundii		0.08	0.22			
23311012	**Good identification**					**23311012**
Actinomyces odontolyticus		99.57	0.54	MLZ 1		
Actinomyces viscosus		0.29	0.04			
Actinomyces naeslundii		0.14	0.22			
23311020	**Very good identification**					**23311020**
Actinomyces odontolyticus		99.68	0.79	NAG 14		
Clostridium perfringens		0.21	0.28			
Actinomyces naeslundii		0.10	0.44			
23311021	**Low discrimination**					**23311021**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		86.54	0.44	NAG 14	1	
Actinomyces naeslundii		12.62	0.47	URE 1	1	
Clostridium chauvoei		0.85	0.04		99	
23311022	**Very good identification**					**23311022**
Actinomyces odontolyticus		99.65	0.75	NAG 14		
Actinomyces naeslundii		0.16	0.44			
Clostridium perfringens		0.16	0.22			
23311023	**Low discrimination**					**23311023**
Actinomyces odontolyticus		80.79	0.41	NAG 14		
Actinomyces naeslundii		19.21	0.47	URE 1		
23311030	**Good identification**					**23311030**
Actinomyces odontolyticus		99.49	0.44	MLZ 1		
Actinomyces naeslundii		0.51	0.22	NAG 14		
23311032	**Good identification**					**23311032**
Actinomyces odontolyticus		99.17	0.41	MLZ 1		
Actinomyces naeslundii		0.83	0.22	NAG 14		
23311040	**Very good identification**					**23311040**
Actinomyces odontolyticus		99.72	0.58	SOR 1		
Actinomyces naeslundii		0.28	0.31			
23311042	**Good identification**					**23311042**
Actinomyces odontolyticus		99.26	0.54	SOR 1		
Actinomyces naeslundii		0.45	0.31			
Actinomyces viscosus		0.29	0.04			
23311060	**Acceptable identification**					**23311060**
Actinomyces odontolyticus		94.67	0.44	NAG 14		
Clostridium perfringens		3.71	0.15	SOR 1		
Actinomyces naeslundii		1.62	0.31			
23311062	**Acceptable identification**					**23311062**
Actinomyces odontolyticus		94.51	0.41	NAG 14		
Clostridium perfringens		2.84	0.10	SOR 1		
Actinomyces naeslundii		2.64	0.31			
23311100	**Very good identification**					**23311100**
					GRA	
Actinomyces odontolyticus		99.59	0.84		99	
Actinomyces naeslundii		0.14	0.52		99	
Fusobacterium mortiferum		0.10	0.37		1	
23311101	**Low discrimination**					**23311101**
					SPO	
Clostridium chauvoei		55.09	0.47	MNS 90	99	
Actinomyces odontolyticus		36.81	0.49	SAL 1	1	
Actinomyces naeslundii		7.87	0.55		1	
23311102	**Very good identification**					**23311102**
Actinomyces odontolyticus		98.01	0.80			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		1.63	0.42			
Actinomyces naeslundii		0.23	0.52			
23311103	**Low discrimination**					**23311103**
					SPO	
Actinomyces odontolyticus		42.85	0.46	URE 1	1	
Actinomyces viscosus		23.45	0.34		1	
Clostridium chauvoei		18.46	0.34		99	
23311110	**Good identification**					**23311110**
Actinomyces odontolyticus		99.26	0.49	MLZ 1		
Actinomyces naeslundii		0.75	0.30			
23311112	**Good identification**					**23311112**
Actinomyces odontolyticus		97.20	0.46	MLZ 1		
Actinomyces viscosus		1.61	0.08			
Actinomyces naeslundii		1.19	0.30			
23311120	**Good identification**					**23311120**
Actinomyces odontolyticus		94.44	0.70	NAG 14		
Clostridium perfringens		4.55	0.42			
Actinomyces naeslundii		0.84	0.52			
23311121	**Low discrimination**					**23311121**
					SPO	
Clostridium chauvoei		56.47	0.38	MNS 90	99	
Actinomyces naeslundii		24.21	0.55	SAL 1	1	
Actinomyces odontolyticus		18.43	0.36		1	
23311122	**Good identification**					**23311122**
Actinomyces odontolyticus		94.92	0.66	NAG 14		
Clostridium perfringens		3.51	0.37			
Actinomyces naeslundii		1.37	0.52			
23311123	**Low discrimination**					**23311123**
					SPO	
Actinomyces naeslundii		52.32	0.55	MNS 88	1	
Actinomyces odontolyticus		24.41	0.32	RAF 92	1	
Clostridium chauvoei		21.54	0.25		99	
23311130	**Acceptable identification**					**23311130**
Actinomyces odontolyticus		91.38	0.36	MLZ 1		
Clostridium perfringens		4.41	0.08	NAG 14		
Actinomyces naeslundii		4.21	0.30			
23311140	**Acceptable identification**					**23311140**
Actinomyces odontolyticus		94.88	0.49	SOR 1		
Actinomyces naeslundii		2.39	0.39			
Propionibacterium propionicum		2.30	0.29			
23311141	**Low discrimination**					**23311141**
					SPO	
Actinomyces naeslundii		57.48	0.42	MNS 88	1	
Clostridium chauvoei		23.02	0.12	RAF 92	99	
Actinomyces odontolyticus		15.39	0.15		1	
23311142	**Low discrimination**					**23311142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		89.52	0.46	SOR 1		
Propionibacterium propionicum		5.32	0.32			
Actinomyces naeslundii		3.68	0.39			
23311143	**Low discrimination**					**23311143**
Actinomyces naeslundii		69.85	0.42	MNS 88		
Actinomyces odontolyticus		11.46	0.12	RAF 92		
Propionibacterium propionicum		7.49	0.15			
23311160	**Low discrimination**					**23311160**
Actinomyces odontolyticus		48.45	0.36	NAG 14		
Clostridium perfringens		44.06	0.30	SOR 1		
Actinomyces naeslundii		7.49	0.39			
23311161	**Low discrimination**					**23311161**
					SPO	
Actinomyces naeslundii		88.22	0.42	MNS 88	1	
Clostridium chauvoei		11.78	0.04	RAF 92	99	
23311163	**Good identification**					**23311163**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23311200	**Excellent identification**					**23311200**
Actinomyces odontolyticus		100.0	0.97			
23311201	**Very good identification**					**23311201**
Actinomyces odontolyticus		100.0	0.63	URE 1		
23311202	**Excellent identification**					**23311202**
Actinomyces odontolyticus		100.0	0.93			
Actinomyces viscosus		0.00	0.04			
23311203	**Very good identification**					**23311203**
Actinomyces odontolyticus		100.0	0.59	URE 1		
23311210	**Very good identification**					**23311210**
Actinomyces odontolyticus		100.0	0.63	MLZ 1		
23311212	**Very good identification**					**23311212**
Actinomyces odontolyticus		100.0	0.59	MLZ 1		
23311220	**Very good identification**					**23311220**
Actinomyces odontolyticus		100.0	0.84	NAG 14		
23311221	**Good identification**					**23311221**
Actinomyces odontolyticus		100.0	0.49	NAG 14		
23311222	**Very good identification**					**23311222**
Actinomyces odontolyticus		100.0	0.80	NAG 14		
23311223	**Good identification**					**23311223**
Actinomyces odontolyticus		100.0	0.46	NAG 14		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23311230	**Good identification**					**23311230**
Actinomyces odontolyticus		100.0	0.49	MLZ 1		
23311232	**Good identification**					**23311232**
Actinomyces odontolyticus		100.0	0.46	MLZ 1		
23311240	**Very good identification**					**23311240**
Actinomyces odontolyticus		100.0	0.63	SOR 1		
23311242	**Very good identification**					**23311242**
Actinomyces odontolyticus		100.0	0.59	SOR 1		
23311260	**Good identification**					**23311260**
Actinomyces odontolyticus		100.0	0.49	NAG 14		
23311262	**Good identification**					**23311262**
Actinomyces odontolyticus		100.0	0.46	NAG 14		
23311300	**Very good identification**					**23311300**
Actinomyces odontolyticus		100.0	0.88			
23311301	**Good identification**					**23311301**
					SPO	
Actinomyces odontolyticus		99.23	0.54	URE 1	1	
Clostridium chauvoei		0.77	0.12		99	
23311302	**Very good identification**					**23311302**
Actinomyces odontolyticus		99.99	0.85			
Actinomyces viscosus		0.01	0.08			
23311303	**Good identification**					**23311303**
					SPO	
Actinomyces odontolyticus		99.49	0.51	URE 1	1	
Actinomyces viscosus		0.28	0.00		1	
Clostridium chauvoei		0.22	-.01		99	
23311310	**Good identification**					**23311310**
Actinomyces odontolyticus		100.0	0.54	MLZ 1		
23311312	**Good identification**					**23311312**
Actinomyces odontolyticus		100.0	0.51	MLZ 1		
23311320	**Very good identification**					**23311320**
Actinomyces odontolyticus		99.98	0.75	NAG 14		
Clostridium perfringens		0.03	0.08			
23311321	**Good identification**					**23311321**
					SPO	
Actinomyces odontolyticus		98.43	0.41	NAG 14	1	
Clostridium chauvoei		1.57	0.04	URE 1	99	
23311322	**Very good identification**					**23311322**
Actinomyces odontolyticus		99.98	0.71	NAG 14		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		0.02	0.03			
23311323	**Good identification**					**23311323**
Actinomyces odontolyticus		100.0	0.37	NAG 14		
23311330	**Good identification**					**23311330**
Actinomyces odontolyticus		100.0	0.41	MLZ 1		
23311332	**Good identification**					**23311332**
Actinomyces odontolyticus		100.0	0.37	MLZ 1		
23311340	**Good identification**					**23311340**
Actinomyces odontolyticus		100.0	0.54	SOR 1		
23311342	**Good identification**					**23311342**
Actinomyces odontolyticus		100.0	0.51	SOR 1		
23311360	**Good identification**					**23311360**
Actinomyces odontolyticus		100.0	0.41	NAG 14		
23311362	**Good identification**					**23311362**
Actinomyces odontolyticus		100.0	0.37	NAG 14		
23311400	**Very good identification**					**23311400**
Actinomyces odontolyticus		99.43	0.58	ARA 1		
Lactobacillus fermentum		0.42	0.24			
Actinomyces meyeri		0.15	0.06			
23311402	**Good identification**					**23311402**
Actinomyces odontolyticus		99.03	0.54	ARA 1		
Lactobacillus fermentum		0.68	0.24			
Actinomyces viscosus		0.29	0.04			
23311420	**Good identification**					**23311420**
Actinomyces odontolyticus		99.14	0.44	ARA 1		
Lactobacillus fermentum		0.86	0.16	NAG 14		
23311422	**Good identification**					**23311422**
Actinomyces odontolyticus		98.61	0.41	ARA 1		
Lactobacillus fermentum		1.39	0.16	NAG 14		
23311500	**Good identification**					**23311500**
Actinomyces odontolyticus		96.02	0.49	ARA 1		
Lactobacillus fermentum		2.09	0.28			
Actinomyces meyeri		1.88	0.16			
23311502	**Good identification**					**23311502**
Actinomyces odontolyticus		95.05	0.46	ARA 1		
Lactobacillus fermentum		3.38	0.28			
Actinomyces viscosus		1.58	0.08			
23311520	**Acceptable identification**					**23311520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		90.15	0.36	ARA 1			
Clostridium perfringens		4.35	0.08	NAG 14			
Lactobacillus fermentum		4.02	0.20				
23311600	**Very good identification**						**23311600**
Actinomyces odontolyticus		100.0	0.63	ARA 1			
23311602	**Very good identification**						**23311602**
Actinomyces odontolyticus		100.0	0.59	ARA 1			
23311620	**Good identification**						**23311620**
Actinomyces odontolyticus		100.0	0.49	ARA 1			
23311622	**Good identification**						**23311622**
Actinomyces odontolyticus		100.0	0.46	ARA 1			
23311700	**Good identification**						**23311700**
Actinomyces odontolyticus		100.0	0.54	ARA 1			
23311702	**Good identification**						**23311702**
Actinomyces odontolyticus		100.0	0.51	ARA 1			
23311720	**Good identification**						**23311720**
Actinomyces odontolyticus		100.0	0.41	ARA 1			
23311722	**Good identification**						**23311722**
Actinomyces odontolyticus		100.0	0.37	ARA 1			
23312000	**Very good identification**						**23312000**
					COC	GRA	
Actinomyces odontolyticus		96.51	0.61	MAN 1	1	99	
Megasphaera elsdenii		2.91	0.32		99	1	
Propionibacterium propionicum		0.49	0.29		1	99	
23312002	**Very good identification**						**23312002**
					COC	GRA	
Actinomyces odontolyticus		97.79	0.57	MAN 1	1	99	
Propionibacterium propionicum		1.22	0.32		1	99	
Megasphaera elsdenii		0.54	0.15		99	1	
23312020	**Good identification**						**23312020**
Actinomyces odontolyticus		99.48	0.47	MAN 1			
Actinomyces naeslundii		0.52	0.26	NAG 14			
23312022	**Good identification**						**23312022**
Actinomyces odontolyticus		99.15	0.44	MAN 1			
Actinomyces naeslundii		0.85	0.26	NAG 14			
23312100	**Low discrimination**						**23312100**
Propionibacterium propionicum		61.39	0.63	SAL 1			
Actinomyces odontolyticus		38.32	0.52				
Actinomyces naeslundii		0.30	0.33				
23312101	**Low discrimination**						**23312101**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		89.16	0.47	SAL 1		
Actinomyces naeslundii		5.78	0.36	URE 10		
Actinomyces odontolyticus		5.06	0.18			
23312102	**Low discrimination**					**23312102**
Propionibacterium propionicum		79.19	0.66	SAL 1		
Actinomyces odontolyticus		20.20	0.49			
Actinomyces viscosus		0.36	0.12			
23312103	**Acceptable identification**					**23312103**
Propionibacterium propionicum		92.58	0.50	SAL 1		
Actinomyces naeslundii		4.00	0.36	URE 10		
Actinomyces odontolyticus		2.15	0.15			
23312120	**Low discrimination**					**23312120**
Actinomyces odontolyticus		87.21	0.39	MAN 1		
Propionibacterium propionicum		8.67	0.29	NAG 14		
Actinomyces naeslundii		4.12	0.33			
23312121	**Low discrimination**					**23312121**
Actinomyces naeslundii		86.51	0.36	MNS 88		
Propionibacterium propionicum		13.49	0.12	RAF 92		
23312122	**Low discrimination**					**23312122**
Actinomyces odontolyticus		75.74	0.35	MAN 1		
Propionibacterium propionicum		18.43	0.32	NAG 14		
Actinomyces naeslundii		5.84	0.33			
23312123	**Low discrimination**					**23312123**
Actinomyces naeslundii		81.05	0.36	MNS 88		
Propionibacterium propionicum		18.95	0.15	RAF 92		
23312140	**Very good identification**					**23312140**
Propionibacterium propionicum		99.37	0.63	SAL 1		
Actinomyces odontolyticus		0.63	0.18			
23312141	**Good identification**					**23312141**
Propionibacterium propionicum		98.87	0.47	SAL 1		
Actinomyces naeslundii		1.13	0.24	URE 10		
23312142	**Very good identification**					**23312142**
Propionibacterium propionicum		99.74	0.66	SAL 1		
Actinomyces odontolyticus		0.26	0.15			
23312143	**Good identification**					**23312143**
Propionibacterium propionicum		99.24	0.50	SAL 1		
Actinomyces naeslundii		0.76	0.24	URE 10		
23312200	**Very good identification**					**23312200**
Actinomyces odontolyticus		100.0	0.66	MAN 1		
23312202	**Very good identification**					**23312202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.62	MAN 1		
23312220	**Good identification**					**23312220**
Actinomyces odontolyticus		100.0	0.52	MAN 1		
23312222	**Good identification**					**23312222**
Actinomyces odontolyticus		100.0	0.49	MAN 1		
23312300	**Very good identification**					**23312300**
Actinomyces odontolyticus		99.17	0.57	MAN 1		
Propionibacterium propionicum		0.83	0.29			
23312302	**Good identification**					**23312302**
Actinomyces odontolyticus		98.00	0.54	MAN 1		
Propionibacterium propionicum		2.00	0.32			
23312320	**Good identification**					**23312320**
Actinomyces odontolyticus		100.0	0.44	MAN 1		
23312322	**Good identification**					**23312322**
Actinomyces odontolyticus		100.0	0.40	MAN 1		
23313000	**Very good identification**					**23313000**
Actinomyces odontolyticus		96.30	0.58	MAN 1		
Actinomyces naeslundii		2.96	0.49			
Propionibacterium propionicum		0.74	0.29			
23313001	**Low discrimination**					**23313001**
Actinomyces naeslundii		80.79	0.52	MNS 88		
Actinomyces odontolyticus		17.72	0.24	RAF 92		
Propionibacterium propionicum		1.49	0.12			
23313002	**Acceptable identification**					**23313002**
Actinomyces odontolyticus		93.13	0.54	MAN 1		
Actinomyces naeslundii		4.67	0.49			
Propionibacterium propionicum		1.75	0.32			
23313003	**Low discrimination**					**23313003**
Actinomyces naeslundii		86.05	0.52	MNS 88		
Actinomyces odontolyticus		11.57	0.20	RAF 92		
Propionibacterium propionicum		2.39	0.15			
23313020	**Low discrimination**					**23313020**
Actinomyces odontolyticus		84.13	0.44	MAN 1		
Actinomyces naeslundii		15.87	0.49	NAG 14		
23313021	**Good identification**					**23313021**
Actinomyces naeslundii		96.55	0.52	MNS 88		
Actinomyces odontolyticus		3.45	0.10	RAF 92		
23313022	**Low discrimination**					**23313022**
Actinomyces odontolyticus		76.47	0.41	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		23.54	0.49	NAG 14		
23313023	**Good identification**					**23313023**
Actinomyces naeslundii		97.86	0.52	MNS 88		
Actinomyces odontolyticus		2.14	0.07	RAF 92		
23313041	**Acceptable identification**					**23313041**
Actinomyces naeslundii		90.52	0.39	MNS 88		
Propionibacterium propionicum		9.48	0.12	RAF 92		
23313043	**Low discrimination**					**23313043**
Actinomyces naeslundii		86.42	0.39	MNS 88		
Propionibacterium propionicum		13.58	0.15	RAF 92		
23313060	**Low discrimination**					**23313060**
Actinomyces naeslundii		76.72	0.36	MNS 88		
Actinomyces odontolyticus		23.28	0.10	RAF 92		
23313061	**Good identification**					**23313061**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23313062	**Low discrimination**					**23313062**
Actinomyces naeslundii		84.32	0.36	MNS 88		
Actinomyces odontolyticus		15.68	0.07	RAF 92		
23313063	**Good identification**					**23313063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23313100	**Low discrimination**					**23313100**
Propionibacterium propionicum		65.12	0.63	SAL 1		
Actinomyces odontolyticus		27.10	0.49			
Actinomyces naeslundii		7.50	0.57			
23313101	**Low discrimination**					**23313101**
					SPO	
Actinomyces naeslundii		58.69	0.60	MNS 88	1	
Propionibacterium propionicum		37.75	0.47	RAF 92	1	
Clostridium chauvoei		2.14	0.12		99	
23313102	**Low discrimination**					**23313102**
Propionibacterium propionicum		79.75	0.66	SAL 1		
Actinomyces odontolyticus		13.56	0.46			
Actinomyces naeslundii		6.12	0.57			
23313103	**Low discrimination**					**23313103**
Actinomyces naeslundii		50.14	0.60	MNS 88		
Propionibacterium propionicum		48.38	0.50	RAF 92		
Actinomyces odontolyticus		0.75	0.12			
23313111	**Low discrimination**					**23313111**
Actinomyces naeslundii		89.01	0.38	MNS 88		
Propionibacterium propionicum		10.99	0.12	RAF 92		
23313113	**Low discrimination**					**23313113**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.38	0.38	MNS 88		
Propionibacterium propionicum		15.62	0.15	RAF 92		
23313120	**Low discrimination**					**23313120**
Actinomyces naeslundii		58.68	0.57	MNS 88		
Actinomyces odontolyticus		34.51	0.36	RAF 92		
Propionibacterium propionicum		5.15	0.29			
23313121	**Very good identification**					**23313121**
					SPO	
Actinomyces naeslundii		98.17	0.60	MNS 88	1	
Clostridium chauvoei		1.19	0.04	RAF 92	99	
Propionibacterium propionicum		0.64	0.12		1	
23313122	**Low discrimination**					**23313122**
Actinomyces naeslundii		66.43	0.57	MNS 88		
Actinomyces odontolyticus		23.95	0.32	RAF 92		
Propionibacterium propionicum		8.74	0.32			
23313123	**Very good identification**					**23313123**
Actinomyces naeslundii		99.04	0.60	MNS 88		
Propionibacterium propionicum		0.97	0.15	RAF 92		
23313130	**Good identification**					**23313130**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23313131	**Good identification**					**23313131**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23313132	**Good identification**					**23313132**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23313133	**Good identification**					**23313133**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23313140	**Very good identification**					**23313140**
Propionibacterium propionicum		97.45	0.63	SAL 1		
Actinomyces naeslundii		1.98	0.44			
Actinomyces odontolyticus		0.41	0.15			
23313141	**Low discrimination**					**23313141**
Propionibacterium propionicum		78.47	0.47	SAL 1		
Actinomyces naeslundii		21.53	0.47	URE 10		
23313142	**Very good identification**					**23313142**
Propionibacterium propionicum		98.33	0.66	SAL 1		
Actinomyces naeslundii		1.33	0.44			
Actinomyces odontolyticus		0.17	0.12			
23313143	**Low discrimination**					**23313143**
Propionibacterium propionicum		84.54	0.50	SAL 1		
Actinomyces naeslundii		15.46	0.47	URE 10		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23313160	**Low discrimination**					**23313160**
Actinomyces naeslundii		66.80	0.44	MNS 88		
Propionibacterium propionicum		33.20	0.29	RAF 92		
23313161	**Good identification**					**23313161**
Actinomyces naeslundii		96.45	0.47	MNS 88		
Propionibacterium propionicum		3.55	0.12	RAF 92		
23313162	**Low discrimination**					**23313162**
Actinomyces naeslundii		57.29	0.44	MNS 88		
Propionibacterium propionicum		42.71	0.32	RAF 92		
23313163	**Acceptable identification**					**23313163**
Actinomyces naeslundii		94.77	0.47	MNS 88		
Propionibacterium propionicum		5.23	0.15	RAF 92		
23313200	**Very good identification**					**23313200**
Actinomyces odontolyticus		100.0	0.63	MAN 1		
23313202	**Very good identification**					**23313202**
Actinomyces odontolyticus		100.0	0.59	MAN 1		
23313220	**Good identification**					**23313220**
Actinomyces odontolyticus		100.0	0.49	MAN 1		
23313222	**Good identification**					**23313222**
Actinomyces odontolyticus		100.0	0.46	MAN 1		
23313300	**Good identification**					**23313300**
Actinomyces odontolyticus		98.63	0.54	MAN 1		
Propionibacterium propionicum		1.23	0.29			
Actinomyces naeslundii		0.14	0.23			
23313302	**Good identification**					**23313302**
Actinomyces odontolyticus		96.81	0.51	MAN 1		
Propionibacterium propionicum		2.96	0.32			
Actinomyces naeslundii		0.23	0.23			
23313320	**Good identification**					**23313320**
Actinomyces odontolyticus		99.12	0.41	MAN 1		
Actinomyces naeslundii		0.88	0.23	NAG 14		
23313322	**Good identification**					**23313322**
Actinomyces odontolyticus		98.58	0.37	MAN 1		
Actinomyces naeslundii		1.42	0.23	NAG 14		
23314000	**Very good identification**					**23314000**
Actinomyces odontolyticus		99.99	0.90			
Actinomyces meyeri		0.01	0.18			
23314001	**Good identification**					**23314001**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces odontolyticus	100.0	0.56	URE 1		
23314002	**Very good identification**					**23314002**
	Actinomyces odontolyticus	99.99	0.86			
	Actinomyces viscosus	0.01	0.08			
23314003	**Good identification**					**23314003**
	Actinomyces odontolyticus	99.78	0.52	URE 1		
	Actinomyces viscosus	0.22	-.01			
23314010	**Good identification**					**23314010**
	Actinomyces odontolyticus	100.0	0.56	MLZ 1		
23314012	**Good identification**					**23314012**
	Actinomyces odontolyticus	100.0	0.52	MLZ 1		
23314020	**Very good identification**					**23314020**
	Actinomyces odontolyticus	99.99	0.76	NAG 14		
	Actinomyces meyeri	0.01	0.04			
23314021	**Good identification**					**23314021**
	Actinomyces odontolyticus	100.0	0.42	NAG 14		
23314022	**Very good identification**					**23314022**
	Actinomyces odontolyticus	100.0	0.73	NAG 14		
23314023	**Good identification**					**23314023**
	Actinomyces odontolyticus	100.0	0.39	NAG 14		
23314030	**Good identification**					**23314030**
	Actinomyces odontolyticus	100.0	0.42	MLZ 1		
23314032	**Good identification**					**23314032**
	Actinomyces odontolyticus	100.0	0.39	MLZ 1		
23314040	**Good identification**					**23314040**
					SPO	
	Actinomyces odontolyticus	99.87	0.56	SOR 1	1	
	Clostridium glycolicum	0.13	-.03		99	
23314042	**Good identification**					**23314042**
	Actinomyces odontolyticus	100.0	0.52	SOR 1		
23314060	**Good identification**					**23314060**
	Actinomyces odontolyticus	100.0	0.42	NAG 14		
23314062	**Good identification**					**23314062**
	Actinomyces odontolyticus	100.0	0.39	NAG 14		
23314100	**Very good identification**					**23314100**
					GRA	
	Actinomyces odontolyticus	99.86	0.81		99	
	Actinomyces meyeri	0.14	0.29		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		0.01	0.15		1	
23314101	**Good identification**					**23314101**
Actinomyces odontolyticus		100.0	0.47	URE	1	
23314102	**Very good identification**					**23314102**
Actinomyces odontolyticus		99.96	0.78			
Actinomyces viscosus		0.04	0.12			
Actinomyces meyeri		0.01	0.03			
23314103	**Good identification**					**23314103**
Actinomyces odontolyticus		98.79	0.43	URE	1	
Actinomyces viscosus		1.21	0.04			
23314110	**Good identification**					**23314110**
Actinomyces odontolyticus		100.0	0.47	MLZ	1	
23314112	**Good identification**					**23314112**
Actinomyces odontolyticus		100.0	0.43	MLZ	1	
23314120	**Very good identification**					**23314120**
Actinomyces odontolyticus		99.88	0.68	NAG	14	
Actinomyces meyeri		0.12	0.14			
23314122	**Very good identification**					**23314122**
Actinomyces odontolyticus		100.0	0.64	NAG	14	
23314140	**Good identification**					**23314140**
Actinomyces odontolyticus		100.0	0.47	SOR	1	
23314142	**Good identification**					**23314142**
Actinomyces odontolyticus		100.0	0.43	SOR	1	
23314200	**Excellent identification**					**23314200**
Actinomyces odontolyticus		100.0	0.95			
23314201	**Very good identification**					**23314201**
Actinomyces odontolyticus		100.0	0.61	URE	1	
23314202	**Excellent identification**					**23314202**
Actinomyces odontolyticus		100.0	0.91			
23314203	**Very good identification**					**23314203**
Actinomyces odontolyticus		100.0	0.57	URE	1	
23314210	**Very good identification**					**23314210**
Actinomyces odontolyticus		100.0	0.61	MLZ	1	
23314212	**Very good identification**					**23314212**
Actinomyces odontolyticus		100.0	0.57	MLZ	1	
23314220	**Very good identification**					**23314220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.81	NAG 14		
23314221	**Good identification**					**23314221**
Actinomyces odontolyticus		100.0	0.47	NAG 14		
23314222	**Very good identification**					**23314222**
Actinomyces odontolyticus		100.0	0.78	NAG 14		
23314223	**Good identification**					**23314223**
Actinomyces odontolyticus		100.0	0.43	NAG 14		
23314230	**Good identification**					**23314230**
Actinomyces odontolyticus		100.0	0.47	MLZ 1		
23314232	**Good identification**					**23314232**
Actinomyces odontolyticus		100.0	0.43	MLZ 1		
23314240	**Very good identification**					**23314240**
Actinomyces odontolyticus		100.0	0.61	SOR 1		
23314242	**Very good identification**					**23314242**
Actinomyces odontolyticus		100.0	0.57	SOR 1		
23314260	**Good identification**					**23314260**
Actinomyces odontolyticus		100.0	0.47	NAG 14		
23314262	**Good identification**					**23314262**
Actinomyces odontolyticus		100.0	0.43	NAG 14		
23314300	**Very good identification**					**23314300**
Actinomyces odontolyticus		100.0	0.86			
23314301	**Good identification**					**23314301**
Actinomyces odontolyticus		100.0	0.52	URE 1		
23314302	**Very good identification**					**23314302**
Actinomyces odontolyticus		100.0	0.83			
23314303	**Good identification**					**23314303**
Actinomyces odontolyticus		100.0	0.48	URE 1		
23314310	**Good identification**					**23314310**
Actinomyces odontolyticus		100.0	0.52	MLZ 1		
23314312	**Good identification**					**23314312**
Actinomyces odontolyticus		100.0	0.48	MLZ 1		
23314320	**Very good identification**					**23314320**
Actinomyces odontolyticus		100.0	0.73	NAG 14		
23314321	**Good identification**					**23314321**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.39	NAG 14		
23314322	**Very good identification**					**23314322**
Actinomyces odontolyticus		100.0	0.69	NAG 14		
23314330	**Good identification**					**23314330**
Actinomyces odontolyticus		100.0	0.39	MLZ 1		
23314340	**Good identification**					**23314340**
Actinomyces odontolyticus		100.0	0.52	SOR 1		
23314342	**Good identification**					**23314342**
Actinomyces odontolyticus		100.0	0.48	SOR 1		
23314360	**Good identification**					**23314360**
Actinomyces odontolyticus		100.0	0.39	NAG 14		
23314400	**Good identification**					**23314400**
Actinomyces odontolyticus		99.73	0.56	ARA 1		
Actinomyces meyeri		0.27	0.08			
23314402	**Good identification**					**23314402**
Actinomyces odontolyticus		100.0	0.52	ARA 1		
23314420	**Good identification**					**23314420**
Actinomyces odontolyticus		100.0	0.42	ARA 1		
23314422	**Good identification**					**23314422**
Actinomyces odontolyticus		100.0	0.39	ARA 1		
23314500	**Good identification**					**23314500**
Actinomyces odontolyticus		96.68	0.47	ARA 1		
Actinomyces meyeri		3.32	0.18			
23314502	**Good identification**					**23314502**
Actinomyces odontolyticus		100.0	0.43	ARA 1		
23314600	**Very good identification**					**23314600**
Actinomyces odontolyticus		100.0	0.61	ARA 1		
23314602	**Very good identification**					**23314602**
Actinomyces odontolyticus		100.0	0.57	ARA 1		
23314620	**Good identification**					**23314620**
Actinomyces odontolyticus		100.0	0.47	ARA 1		
23314622	**Good identification**					**23314622**
Actinomyces odontolyticus		100.0	0.43	ARA 1		
23314700	**Good identification**					**23314700**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces odontolyticus	100.0	0.52	ARA 1		
23314702	**Good identification**					**23314702**
	Actinomyces odontolyticus	100.0	0.48	ARA 1		
23314720	**Good identification**					**23314720**
	Actinomyces odontolyticus	100.0	0.39	ARA 1		
23315000	**Very good identification**					**23315000**
	Actinomyces odontolyticus	99.97	0.87			
	Actinomyces meyeri	0.03	0.21			
	Lactobacillus fermentum	0.01	0.20			
23315001	**Good identification**					**23315001**
	Actinomyces odontolyticus	100.0	0.53	URE 1		
23315002	**Very good identification**					**23315002**
	Actinomyces odontolyticus	99.98	0.83			
	Actinomyces israelii	0.01	0.21			
	Lactobacillus fermentum	0.01	0.20			
23315003	**Good identification**					**23315003**
	Actinomyces odontolyticus	100.0	0.49	URE 1		
23315010	**Good identification**					**23315010**
	Actinomyces odontolyticus	100.0	0.53	MLZ 1		
23315012	**Good identification**					**23315012**
	Actinomyces odontolyticus	100.0	0.49	MLZ 1		
23315020	**Very good identification**					**23315020**
	Actinomyces odontolyticus	99.97	0.73	NAG 14		
	Actinomyces meyeri	0.02	0.07			
	Lactobacillus fermentum	0.01	0.12			
23315021	**Good identification**					**23315021**
	Actinomyces odontolyticus	100.0	0.39	NAG 14		
23315022	**Very good identification**					**23315022**
	Actinomyces odontolyticus	99.98	0.70	NAG 14		
	Lactobacillus fermentum	0.02	0.12			
23315023	**Good identification**					**23315023**
	Actinomyces odontolyticus	100.0	0.36	NAG 14		
23315030	**Good identification**					**23315030**
	Actinomyces odontolyticus	100.0	0.39	MLZ 1		
23315032	**Good identification**					**23315032**
	Actinomyces odontolyticus	100.0	0.36	MLZ 1		
23315040	**Good identification**					**23315040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.53	SOR 1		
23315042	**Good identification**					**23315042**
Actinomyces odontolyticus		99.68	0.49	SOR 1		
Actinomyces israelii		0.32	0.14			
23315060	**Good identification**					**23315060**
Actinomyces odontolyticus		100.0	0.39	NAG 14		
23315062	**Good identification**					**23315062**
Actinomyces odontolyticus		100.0	0.36	NAG 14		
23315100	**Very good identification**					**23315100**
Actinomyces odontolyticus		99.61	0.78			
Actinomyces meyeri		0.31	0.32			
Actinomyces israelii		0.04	0.29			
23315101	**Good identification**					**23315101**
					SPO	
Actinomyces odontolyticus		97.02	0.44	URE 1	1	
Clostridium chauvoei		2.98	0.12		99	
23315102	**Very good identification**					**23315102**
Actinomyces odontolyticus		99.80	0.75			
Actinomyces israelii		0.11	0.32			
Lactobacillus fermentum		0.04	0.24			
23315103	**Good identification**					**23315103**
					SPO	
Actinomyces odontolyticus		98.03	0.40	URE 1	1	
Actinomyces viscosus		1.10	0.00		1	
Clostridium chauvoei		0.87	-.01		99	
23315110	**Good identification**					**23315110**
Actinomyces odontolyticus		99.37	0.44	MLZ 1		
Actinomyces israelii		0.64	0.15			
23315112	**Good identification**					**23315112**
Actinomyces odontolyticus		98.46	0.40	MLZ 1		
Actinomyces israelii		1.54	0.18			
23315120	**Very good identification**					**23315120**
Actinomyces odontolyticus		99.59	0.65	NAG 14		
Actinomyces meyeri		0.26	0.17			
Clostridium perfringens		0.10	0.08			
23315122	**Very good identification**					**23315122**
Actinomyces odontolyticus		99.84	0.61	NAG 14		
Lactobacillus fermentum		0.09	0.16			
Clostridium perfringens		0.08	0.03			
23315140	**Good identification**					**23315140**
Actinomyces odontolyticus		98.36	0.44	SOR 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	1.64	0.22			
23315142	**Good identification**					**23315142**
	Actinomyces odontolyticus	96.09	0.40	SOR 1		
	Actinomyces israelii	3.91	0.25			
23315200	**Excellent identification**					**23315200**
	Actinomyces odontolyticus	100.0	0.92			
23315201	**Very good identification**					**23315201**
	Actinomyces odontolyticus	100.0	0.58	URE 1		
23315202	**Very good identification**					**23315202**
	Actinomyces odontolyticus	100.0	0.88			
23315203	**Good identification**					**23315203**
	Actinomyces odontolyticus	100.0	0.54	URE 1		
23315210	**Very good identification**					**23315210**
	Actinomyces odontolyticus	100.0	0.58	MLZ 1		
23315212	**Good identification**					**23315212**
	Actinomyces odontolyticus	100.0	0.54	MLZ 1		
23315220	**Very good identification**					**23315220**
	Actinomyces odontolyticus	100.0	0.78	NAG 14		
23315221	**Good identification**					**23315221**
	Actinomyces odontolyticus	100.0	0.44	NAG 14		
23315222	**Very good identification**					**23315222**
	Actinomyces odontolyticus	100.0	0.75	NAG 14		
23315223	**Good identification**					**23315223**
	Actinomyces odontolyticus	100.0	0.40	NAG 14		
23315230	**Good identification**					**23315230**
	Actinomyces odontolyticus	100.0	0.44	MLZ 1		
23315232	**Good identification**					**23315232**
	Actinomyces odontolyticus	100.0	0.40	MLZ 1		
23315240	**Very good identification**					**23315240**
	Actinomyces odontolyticus	100.0	0.58	SOR 1		
23315242	**Good identification**					**23315242**
	Actinomyces odontolyticus	100.0	0.54	SOR 1		
23315260	**Good identification**					**23315260**
	Actinomyces odontolyticus	100.0	0.44	NAG 14		
23315262	**Good identification**					**23315262**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.40	NAG 14		
23315300	**Very good identification**					**23315300**
Actinomyces odontolyticus		99.75	0.83			
Eubacterium contortum		0.25	0.32			
23315301	**Good identification**					**23315301**
Actinomyces odontolyticus		100.0	0.49	URE 1		
23315302	**Very good identification**					**23315302**
Actinomyces odontolyticus		99.93	0.80			
Eubacterium contortum		0.07	0.19			
23315303	**Good identification**					**23315303**
Actinomyces odontolyticus		100.0	0.45	URE 1		
23315310	**Good identification**					**23315310**
Actinomyces odontolyticus		100.0	0.49	MLZ 1		
23315312	**Good identification**					**23315312**
Actinomyces odontolyticus		100.0	0.45	MLZ 1		
23315320	**Very good identification**					**23315320**
Actinomyces odontolyticus		99.73	0.70	NAG 14		
Eubacterium contortum		0.27	0.19			
23315321	**Good identification**					**23315321**
Actinomyces odontolyticus		100.0	0.36	NAG 14		
23315322	**Very good identification**					**23315322**
Actinomyces odontolyticus		99.92	0.66	NAG 14		
Eubacterium contortum		0.08	0.06			
23315330	**Good identification**					**23315330**
Actinomyces odontolyticus		100.0	0.36	MLZ 1		
23315340	**Good identification**					**23315340**
Actinomyces odontolyticus		100.0	0.49	SOR 1		
23315342	**Good identification**					**23315342**
Actinomyces odontolyticus		100.0	0.45	SOR 1		
23315360	**Good identification**					**23315360**
Actinomyces odontolyticus		100.0	0.36	NAG 14		
23315400	**Good identification**					**23315400**
Actinomyces odontolyticus		98.63	0.53	ARA 1		
Actinomyces meyeri		0.60	0.11			
Lactobacillus fermentum		0.50	0.20			
23315402	**Good identification**					**23315402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		98.53	0.49	ARA 1		
Lactobacillus fermentum		0.81	0.20			
Actinomyces israelii		0.67	0.20			
23315420	**Good identification**					**23315420**
Actinomyces odontolyticus		98.98	0.39	ARA 1		
Lactobacillus fermentum		1.02	0.12	NAG 14		
23315422	**Good identification**					**23315422**
Actinomyces odontolyticus		98.35	0.36	ARA 1		
Lactobacillus fermentum		1.65	0.12	NAG 14		
23315500	**Low discrimination**					**23315500**
Actinomyces odontolyticus		87.85	0.44	ARA 1		
Actinomyces meyeri		6.79	0.21			
Actinomyces israelii		3.08	0.27			
23315502	**Low discrimination**					**23315502**
Actinomyces odontolyticus		86.33	0.40	ARA 1		
Actinomyces israelii		7.40	0.30			
Lactobacillus fermentum		3.66	0.24			
23315600	**Very good identification**					**23315600**
Actinomyces odontolyticus		100.0	0.58	ARA 1		
23315602	**Good identification**					**23315602**
Actinomyces odontolyticus		100.0	0.54	ARA 1		
23315620	**Good identification**					**23315620**
Actinomyces odontolyticus		100.0	0.44	ARA 1		
23315622	**Good identification**					**23315622**
Actinomyces odontolyticus		100.0	0.40	ARA 1		
23315700	**Low discrimination**					**23315700**
Actinomyces odontolyticus		83.07	0.49	ARA 1		
Eubacterium contortum		16.93	0.30			
23315702	**Acceptable identification**					**23315702**
Actinomyces odontolyticus		94.46	0.45	ARA 1		
Eubacterium contortum		5.54	0.17			
23315720	**Low discrimination**					**23315720**
Actinomyces odontolyticus		81.90	0.36	ARA 1		
Eubacterium contortum		18.10	0.17	NAG 14		
23316000	**Good identification**					**23316000**
Actinomyces odontolyticus		100.0	0.56	MAN 1		
23316002	**Good identification**					**23316002**
Actinomyces odontolyticus		100.0	0.52	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23316020	**Good identification**					**23316020**
Actinomyces odontolyticus		100.0	0.42	MAN 1		
23316022	**Good identification**					**23316022**
Actinomyces odontolyticus		100.0	0.39	MAN 1		
23316100	**Good identification**					**23316100**
Actinomyces odontolyticus		96.82	0.47	MAN 1		
Propionibacterium propionicum		3.18	0.29			
23316102	**Acceptable identification**					**23316102**
Actinomyces odontolyticus		92.56	0.43	MAN 1		
Propionibacterium propionicum		7.44	0.32			
23316200	**Very good identification**					**23316200**
Actinomyces odontolyticus		100.0	0.61	MAN 1		
23316202	**Very good identification**					**23316202**
Actinomyces odontolyticus		100.0	0.57	MAN 1		
23316220	**Good identification**					**23316220**
Actinomyces odontolyticus		100.0	0.47	MAN 1		
23316222	**Good identification**					**23316222**
Actinomyces odontolyticus		100.0	0.43	MAN 1		
23316300	**Good identification**					**23316300**
Actinomyces odontolyticus		100.0	0.52	MAN 1		
23316302	**Good identification**					**23316302**
Actinomyces odontolyticus		100.0	0.48	MAN 1		
23316320	**Good identification**					**23316320**
Actinomyces odontolyticus		100.0	0.39	MAN 1		
23317000	**Good identification**					**23317000**
Actinomyces odontolyticus		98.12	0.53	MAN 1		
Actinomyces israelii		1.88	0.31			
23317002	**Good identification**					**23317002**
Actinomyces odontolyticus		95.53	0.49	MAN 1		
Actinomyces israelii		4.48	0.34			
23317020	**Good identification**					**23317020**
Actinomyces odontolyticus		100.0	0.39	MAN 1		
23317022	**Good identification**					**23317022**
Actinomyces odontolyticus		98.20	0.36	MAN 1		
Actinomyces israelii		1.80	0.14	NAG 14		
23317100	**Low discrimination**					**23317100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		77.08	0.44	MAN 1		
Actinomyces israelii		18.69	0.42			
Propionibacterium propionicum		3.80	0.29			
23317102	**Low discrimination**					**23317102**
Actinomyces odontolyticus		58.03	0.40	MAN 1		
Actinomyces israelii		34.43	0.45			
Propionibacterium propionicum		7.00	0.32			
23317142	**Low discrimination**					**23317142**
Actinomyces israelii		63.84	0.38	CEL 93		
Propionibacterium propionicum		33.37	0.32			
Actinomyces odontolyticus		2.79	0.06			
23317200	**Very good identification**					**23317200**
Actinomyces odontolyticus		100.0	0.58	MAN 1		
23317202	**Good identification**					**23317202**
Actinomyces odontolyticus		99.82	0.54	MAN 1		
Actinomyces israelii		0.18	0.15			
23317220	**Good identification**					**23317220**
Actinomyces odontolyticus		100.0	0.44	MAN 1		
23317222	**Good identification**					**23317222**
Actinomyces odontolyticus		100.0	0.40	MAN 1		
23317300	**Good identification**					**23317300**
Actinomyces odontolyticus		99.07	0.49	MAN 1		
Actinomyces israelii		0.93	0.22			
23317302	**Good identification**					**23317302**
Actinomyces odontolyticus		97.75	0.45	MAN 1		
Actinomyces israelii		2.25	0.25			
23317320	**Good identification**					**23317320**
Actinomyces odontolyticus		100.0	0.36	MAN 1		
23317500	**Good identification**					**23317500**
Actinomyces israelii		95.16	0.40	CEL 93		
Actinomyces odontolyticus		4.85	0.10			
23317502	**Acceptable identification**					**23317502**
Actinomyces israelii		93.29	0.43	CEL 93		
Bifidobacterium adolescentis		4.77	0.15			
Actinomyces odontolyticus		1.94	0.06			
23317542	**Good identification**					**23317542**
Actinomyces israelii		100.0	0.36	CEL 93		
23320000	**Good identification**					**23320000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.43	FRU 92		
23320002	**Good identification**					**23320002**
Actinomyces odontolyticus		98.58	0.39	FRU 92		
Actinomyces viscosus		1.43	0.01	TRE 1		
23320200	**Good identification**					**23320200**
Actinomyces odontolyticus		100.0	0.48	FRU 92		
23320202	**Good identification**					**23320202**
Actinomyces odontolyticus		100.0	0.44	FRU 92		
23320300	**Good identification**					**23320300**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23320302	**Good identification**					**23320302**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23321000	**Good identification**					**23321000**
Actinomyces odontolyticus		97.00	0.40	FRU 92		
Actinomyces naeslundii		1.80	0.27	TRE 1		
Eubacterium tortuosum		1.20	-.04			
23321002	**Good identification**					**23321002**
Actinomyces odontolyticus		97.06	0.36	FRU 92		
Actinomyces naeslundii		2.94	0.27	TRE 1		
23321103	**Low discrimination**					**23321103**
					SPO	
Actinomyces naeslundii		63.47	0.38	MNS 88	1	
Clostridium chauvoei		36.53	0.11	RAF 92	99	
23321122	**Low discrimination**					**23321122**
Actinomyces naeslundii		48.39	0.35	MNS 88		
Actinomyces odontolyticus		28.82	0.14	RAF 92		
Clostridium perfringens		22.79	0.07	FRU 95		
23321123	**Low discrimination**					**23321123**
					SPO	
Actinomyces naeslundii		83.91	0.38	MNS 88	1	
Clostridium chauvoei		16.10	0.03	RAF 92	99	
23321200	**Good identification**					**23321200**
Actinomyces odontolyticus		100.0	0.45	FRU 92		
23321202	**Good identification**					**23321202**
Actinomyces odontolyticus		100.0	0.41	FRU 92		
23321300	**Good identification**					**23321300**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23323001	**Good identification**					**23323001**
Actinomyces naeslundii		100.0	0.35	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23323003	**Good identification**					**23323003**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23323021	**Good identification**					**23323021**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23323023	**Good identification**					**23323023**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23323100	**Low discrimination**					**23323100**
Actinomyces naeslundii		58.74	0.40	MNS 88		
Propionibacterium propionicum		41.26	0.27	RAF 92		
23323101	**Good identification**					**23323101**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23323103	**Acceptable identification**					**23323103**
Actinomyces naeslundii		92.76	0.43	MNS 88		
Propionibacterium propionicum		7.24	0.14	RAF 92		
23323120	**Good identification**					**23323120**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23323121	**Good identification**					**23323121**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23323122	**Good identification**					**23323122**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23323123	**Good identification**					**23323123**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23324000	**Good identification**					**23324000**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23324200	**Good identification**					**23324200**
Actinomyces odontolyticus		100.0	0.42	FRU 92		
23324202	**Good identification**					**23324202**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23325200	**Good identification**					**23325200**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23325202	**Good identification**					**23325202**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23330000	**Very good identification**					**23330000**
					GRA	
Actinomyces odontolyticus		99.75	0.61	TRE 1	99	
Fusobacterium mortiferum		0.17	0.19		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	0.09	0.26		99	
23330002	**Very good identification**					**23330002**
	Actinomyces odontolyticus	97.54	0.57	TRE 1		
	Actinomyces viscosus	2.33	0.23			
	Actinomyces naeslundii	0.14	0.26			
23330020	**Good identification**					**23330020**
	Actinomyces odontolyticus	99.48	0.47	TRE 1		
	Actinomyces naeslundii	0.52	0.26	NAG 14		
23330022	**Good identification**					**23330022**
	Actinomyces odontolyticus	99.15	0.44	TRE 1		
	Actinomyces naeslundii	0.85	0.26	NAG 14		
23330100	**Good identification**					**23330100**
					GRA	
	Actinomyces odontolyticus	96.76	0.52	TRE 1	99	
	Propionibacterium propionicum	1.28	0.27		99	
	Fusobacterium mortiferum	1.22	0.25		1	
23330101	**Low discrimination**					**23330101**
	Actinomyces naeslundii	53.31	0.36	MNS 88		
	Actinomyces odontolyticus	46.69	0.18	RAF 92		
23330102	**Low discrimination**					**23330102**
	Actinomyces odontolyticus	84.79	0.49	TRE 1		
	Actinomyces viscosus	11.40	0.27			
	Propionibacterium propionicum	2.75	0.30			
23330120	**Acceptable identification**					**23330120**
	Actinomyces odontolyticus	90.34	0.39	TRE 1		
	Clostridium perfringens	5.39	0.13	NAG 14		
	Actinomyces naeslundii	4.27	0.33			
23330121	**Good identification**					**23330121**
	Actinomyces naeslundii	100.0	0.36	MNS 88		
23330122	**Low discrimination**					**23330122**
	Actinomyces odontolyticus	89.06	0.35	TRE 1		
	Actinomyces naeslundii	6.86	0.33	NAG 14		
	Clostridium perfringens	4.08	0.07			
23330123	**Good identification**					**23330123**
	Actinomyces naeslundii	100.0	0.36	MNS 88		
23330200	**Very good identification**					**23330200**
	Actinomyces odontolyticus	100.0	0.66	TRE 1		
23330202	**Very good identification**					**23330202**
	Actinomyces odontolyticus	100.0	0.62	TRE 1		
23330220	**Good identification**					**23330220**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces odontolyticus	100.0	0.52	TRE 1		
23330222	**Good identification**					**23330222**
	Actinomyces odontolyticus	100.0	0.49	TRE 1		
23330300	**Very good identification**					**23330300**
	Actinomyces odontolyticus	100.0	0.57	TRE 1		
23330302	**Good identification**					**23330302**
	Actinomyces odontolyticus	100.0	0.54	TRE 1		
23330320	**Good identification**					**23330320**
	Actinomyces odontolyticus	100.0	0.44	TRE 1		
23330322	**Good identification**					**23330322**
	Actinomyces odontolyticus	100.0	0.40	TRE 1		
23331000	**Very good identification**					**23331000**
					GRA	
	Actinomyces odontolyticus	96.23	0.58	TRE 1	99	
	Actinomyces naeslundii	2.96	0.49		99	
	Fusobacterium mortiferum	0.37	0.22		1	
23331001	**Low discrimination**					**23331001**
	Actinomyces naeslundii	82.02	0.52	MNS 88		
	Actinomyces odontolyticus	17.98	0.24	RAF 92		
23331002	**Acceptable identification**					**23331002**
	Actinomyces odontolyticus	92.89	0.54	TRE 1		
	Actinomyces naeslundii	4.65	0.49			
	Actinomyces viscosus	2.04	0.19			
23331003	**Low discrimination**					**23331003**
	Actinomyces naeslundii	81.19	0.52	MNS 88		
	Actinomyces odontolyticus	10.91	0.20	RAF 92		
	Actinomyces viscosus	7.90	0.11			
23331020	**Low discrimination**					**23331020**
	Actinomyces odontolyticus	63.68	0.44	TRE 1		
	Clostridium perfringens	24.31	0.32	NAG 14		
	Actinomyces naeslundii	12.01	0.49			
23331021	**Good identification**					**23331021**
	Actinomyces naeslundii	96.55	0.52	MNS 88		
	Actinomyces odontolyticus	3.45	0.10	RAF 92		
23331022	**Low discrimination**					**23331022**
	Actinomyces odontolyticus	62.46	0.41	TRE 1		
	Actinomyces naeslundii	19.23	0.49	NAG 14		
	Clostridium perfringens	18.31	0.27			
23331023	**Good identification**					**23331023**
	Actinomyces naeslundii	97.86	0.52	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		2.14	0.07	RAF 92		
23331041	**Good identification**					**23331041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23331043	**Good identification**					**23331043**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23331061	**Good identification**					**23331061**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23331063	**Good identification**					**23331063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23331100	**Low discrimination**					**23331100**
Actinomyces odontolyticus		71.70	0.49	TRE 1		
Actinomyces naeslundii		19.85	0.57			
Clostridium perfringens		3.20	0.21			
23331101	**Low discrimination**					**23331101**
					SPO	
Actinomyces naeslundii		70.18	0.60	MNS 88	1	
Clostridium chauvoei		28.11	0.30	RAF 92	99	
Actinomyces odontolyticus		1.71	0.15		1	
23331102	**Low discrimination**					**23331102**
Actinomyces odontolyticus		59.34	0.46	TRE 1		
Actinomyces naeslundii		26.80	0.57			
Actinomyces viscosus		7.33	0.23			
23331103	**Low discrimination**					**23331103**
					SPO	
Actinomyces naeslundii		86.63	0.60	MNS 88	1	
Clostridium chauvoei		6.12	0.17	RAF 92	99	
Actinomyces viscosus		5.27	0.15		1	
23331110	**Low discrimination**					**23331110**
Actinomyces naeslundii		59.05	0.35	MNS 88		
Actinomyces odontolyticus		40.95	0.15	RAF 92		
23331111	**Good identification**					**23331111**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23331112	**Low discrimination**					**23331112**
Actinomyces naeslundii		70.17	0.35	MNS 88		
Actinomyces odontolyticus		29.83	0.12	RAF 92		
23331113	**Good identification**					**23331113**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23331120	**Low discrimination**					**23331120**
Clostridium perfringens		76.53	0.47	MNS 99		
Actinomyces naeslundii		14.67	0.57			
Actinomyces odontolyticus		8.63	0.36			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23331121	**Low discrimination**					**23331121**
					SPO	
Actinomyces naeslundii		85.57	0.60	MNS 88	1	
Clostridium chauvoei		11.42	0.22	RAF 92	99	
Clostridium perfringens		3.01	0.13		84	
23331122	**Low discrimination**					**23331122**
Clostridium perfringens		64.33	0.41	MNS 99		
Actinomyces naeslundii		26.22	0.57			
Actinomyces odontolyticus		9.45	0.32			
23331123	**Very good identification**					**23331123**
					SPO	
Actinomyces naeslundii		96.15	0.60	MNS 88	1	
Clostridium chauvoei		2.27	0.09	RAF 92	99	
Clostridium perfringens		1.59	0.07		84	
23331131	**Good identification**					**23331131**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23331132	**Low discrimination**					**23331132**
Actinomyces naeslundii		67.98	0.35	MNS 88		
Clostridium perfringens		32.02	0.07	RAF 92		
23331133	**Good identification**					**23331133**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23331140	**Low discrimination**					**23331140**
Actinomyces naeslundii		54.28	0.44	MNS 88		
Propionibacterium propionicum		22.07	0.27	RAF 92		
Actinomyces odontolyticus		11.23	0.15			
23331141	**Good identification**					**23331141**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23331142	**Low discrimination**					**23331142**
Actinomyces naeslundii		52.60	0.44	MNS 88		
Propionibacterium propionicum		32.09	0.30	RAF 92		
Actinomyces odontolyticus		6.67	0.12			
23331143	**Good identification**					**23331143**
Actinomyces naeslundii		95.68	0.47	MNS 88		
Propionibacterium propionicum		4.32	0.14	RAF 92		
23331161	**Good identification**					**23331161**
Actinomyces naeslundii		96.35	0.47	MNS 88		
Clostridium perfringens		3.65	0.01	RAF 92		
23331163	**Good identification**					**23331163**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23331200	**Very good identification**					**23331200**
Actinomyces odontolyticus		100.0	0.63	TRE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23331202	**Very good identification**					**23331202**
Actinomyces odontolyticus		100.0	0.59	TRE 1		
23331220	**Good identification**					**23331220**
Actinomyces odontolyticus		100.0	0.49	TRE 1		
23331222	**Good identification**					**23331222**
Actinomyces odontolyticus		100.0	0.46	TRE 1		
23331300	**Good identification**					**23331300**
Actinomyces odontolyticus		99.86	0.54	TRE 1		
Actinomyces naeslundii		0.14	0.23			
23331302	**Good identification**					**23331302**
Actinomyces odontolyticus		99.77	0.51	TRE 1		
Actinomyces naeslundii		0.23	0.23			
23331320	**Acceptable identification**					**23331320**
Actinomyces odontolyticus		94.79	0.41	TRE 1		
Clostridium perfringens		4.37	0.13	NAG 14		
Actinomyces naeslundii		0.84	0.23			
23331322	**Good identification**					**23331322**
Actinomyces odontolyticus		95.25	0.37	TRE 1		
Clostridium perfringens		3.37	0.07	NAG 14		
Actinomyces naeslundii		1.38	0.23			
23332100	**Very good identification**					**23332100**
Propionibacterium propionicum		98.13	0.61	SAL 1		
Actinomyces naeslundii		1.12	0.38			
Actinomyces odontolyticus		0.76	0.18			
23332101	**Low discrimination**					**23332101**
Propionibacterium propionicum		86.68	0.45	SAL 1		
Actinomyces naeslundii		13.32	0.41	URE 10		
23332102	**Very good identification**					**23332102**
Propionibacterium propionicum		98.94	0.64	SAL 1		
Actinomyces naeslundii		0.75	0.38			
Actinomyces odontolyticus		0.31	0.15			
23332103	**Acceptable identification**					**23332103**
Propionibacterium propionicum		90.71	0.48	SAL 1		
Actinomyces naeslundii		9.29	0.41	URE 10		
23332120	**Low discrimination**					**23332120**
Actinomyces naeslundii		52.99	0.38	MNS 88		
Propionibacterium propionicum		47.01	0.27	RAF 92		
23332121	**Good identification**					**23332121**
Actinomyces naeslundii		100.0	0.41	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23332123	**Acceptable identification**					**23332123**
Actinomyces naeslundii		91.03	0.41	MNS 88		
Propionibacterium propionicum		8.97	0.14	RAF 92		
23332140	**Very good identification**					**23332140**
Propionibacterium propionicum		99.80	0.61	SAL 1		
Actinomyces naeslundii		0.20	0.25			
23332141	**Good identification**					**23332141**
Propionibacterium propionicum		97.36	0.45	SAL 1		
Actinomyces naeslundii		2.64	0.28	URE 10		
23332142	**Very good identification**					**23332142**
Propionibacterium propionicum		99.87	0.64	SAL 1		
Actinomyces naeslundii		0.13	0.25			
23332143	**Good identification**					**23332143**
Propionibacterium propionicum		98.22	0.48	SAL 1		
Actinomyces naeslundii		1.78	0.28	URE 10		
23333000	**Low discrimination**					**23333000**
Actinomyces naeslundii		75.39	0.54	MNS 88		
Actinomyces odontolyticus		12.78	0.24	RAF 92		
Propionibacterium propionicum		7.93	0.27			
23333001	**Very good identification**					**23333001**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23333002	**Low discrimination**					**23333002**
Actinomyces naeslundii		74.67	0.54	MNS 88		
Propionibacterium propionicum		11.79	0.30	RAF 92		
Actinomyces odontolyticus		7.75	0.20			
23333003	**Very good identification**					**23333003**
Actinomyces naeslundii		98.84	0.57	MNS 88		
Propionibacterium propionicum		1.16	0.14	RAF 92		
23333011	**Good identification**					**23333011**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23333013	**Good identification**					**23333013**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23333020	**Good identification**					**23333020**
Actinomyces naeslundii		97.32	0.54	MNS 88		
Actinomyces odontolyticus		2.69	0.10	RAF 92		
23333021	**Very good identification**					**23333021**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23333022	**Good identification**					**23333022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.34	0.54	MNS 88		
Actinomyces odontolyticus		1.66	0.07	RAF 92		
23333023	**Very good identification**					**23333023**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23333031	**Good identification**					**23333031**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23333033	**Good identification**					**23333033**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23333040	**Low discrimination**					**23333040**
Actinomyces naeslundii		58.47	0.41	MNS 88		
Propionibacterium propionicum		34.86	0.27	RAF 92		
Actinomyces israelii		6.67	0.16			
23333041	**Good identification**					**23333041**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23333042	**Low discrimination**					**23333042**
Actinomyces naeslundii		48.42	0.41	MNS 88		
Propionibacterium propionicum		43.30	0.30	RAF 92		
Actinomyces israelii		8.28	0.19			
23333043	**Acceptable identification**					**23333043**
Actinomyces naeslundii		93.79	0.44	MNS 88		
Propionibacterium propionicum		6.21	0.14	RAF 92		
23333060	**Good identification**					**23333060**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23333061	**Good identification**					**23333061**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23333062	**Good identification**					**23333062**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23333063	**Good identification**					**23333063**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23333100	**Low discrimination**					**23333100**
Propionibacterium propionicum		77.03	0.61	SAL 1		
Actinomyces naeslundii		21.05	0.62			
Actinomyces israelii		1.53	0.33			
23333101	**Low discrimination**					**23333101**
Actinomyces naeslundii		78.67	0.65	MNS 88		
Propionibacterium propionicum		21.33	0.45	RAF 92		
23333102	**Low discrimination**					**23333102**
Propionibacterium propionicum		83.05	0.64	SAL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		15.13	0.62			
Actinomyces israelii		1.65	0.36			
23333103	**Low discrimination**					**23333103**
Actinomyces naeslundii		71.09	0.65	MNS 88		
Propionibacterium propionicum		28.91	0.48	RAF 92		
23333110	**Low discrimination**					**23333110**
Actinomyces naeslundii		52.39	0.40	MNS 88		
Propionibacterium propionicum		36.80	0.27	RAF 92		
Actinomyces israelii		10.81	0.19	MLZ 5		
23333111	**Good identification**					**23333111**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23333113	**Acceptable identification**					**23333113**
Actinomyces naeslundii		92.76	0.43	MNS 88		
Propionibacterium propionicum		7.24	0.14	RAF 92		
23333120	**Good identification**					**23333120**
Actinomyces naeslundii		93.98	0.62	MNS 88		
Propionibacterium propionicum		3.47	0.27	RAF 92		
Clostridium perfringens		2.55	0.13			
23333121	**Very good identification**					**23333121**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23333122	**Good identification**					**23333122**
Actinomyces naeslundii		93.01	0.62	MNS 88		
Propionibacterium propionicum		5.16	0.30	RAF 92		
Clostridium perfringens		1.19	0.07			
23333123	**Very good identification**					**23333123**
Actinomyces naeslundii		99.59	0.65	MNS 88		
Propionibacterium propionicum		0.41	0.14	RAF 92		
23333130	**Good identification**					**23333130**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23333131	**Good identification**					**23333131**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23333132	**Good identification**					**23333132**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23333133	**Good identification**					**23333133**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23333140	**Good identification**					**23333140**
Propionibacterium propionicum		94.70	0.61	SAL 1		
Actinomyces naeslundii		4.57	0.49			
Actinomyces israelii		0.73	0.26			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23333141	**Low discrimination**					**23333141**
Propionibacterium propionicum		60.57	0.45	SAL 1		
Actinomyces naeslundii		39.43	0.52	URE 10		
23333142	**Very good identification**					**23333142**
Propionibacterium propionicum		96.17	0.64	SAL 1		
Actinomyces naeslundii		3.09	0.49			
Actinomyces israelii		0.74	0.29			
23333143	**Low discrimination**					**23333143**
Propionibacterium propionicum		69.74	0.48	SAL 1		
Actinomyces naeslundii		30.26	0.52	URE 10		
23333160	**Low discrimination**					**23333160**
Actinomyces naeslundii		80.73	0.49	MNS 88		
Propionibacterium propionicum		16.91	0.27	RAF 92		
Clostridium perfringens		2.36	0.01			
23333161	**Good identification**					**23333161**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23333162	**Low discrimination**					**23333162**
Actinomyces naeslundii		76.09	0.49	MNS 88		
Propionibacterium propionicum		23.91	0.30	RAF 92		
23333163	**Good identification**					**23333163**
Actinomyces naeslundii		97.73	0.52	MNS 88		
Propionibacterium propionicum		2.28	0.14	RAF 92		
23334000	**Good identification**					**23334000**
Actinomyces odontolyticus		100.0	0.56	TRE 1		
23334002	**Good identification**					**23334002**
Actinomyces odontolyticus		100.0	0.52	TRE 1		
23334020	**Good identification**					**23334020**
Actinomyces odontolyticus		100.0	0.42	TRE 1		
23334022	**Good identification**					**23334022**
Actinomyces odontolyticus		100.0	0.39	TRE 1		
23334100	**Good identification**					**23334100**
Actinomyces odontolyticus		100.0	0.47	TRE 1		
23334102	**Good identification**					**23334102**
Actinomyces odontolyticus		100.0	0.43	TRE 1		
23334200	**Very good identification**					**23334200**
Actinomyces odontolyticus		100.0	0.61	TRE 1		
23334202	**Very good identification**					**23334202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.57	TRE 1		
23334220	**Good identification**					**23334220**
Actinomyces odontolyticus		100.0	0.47	TRE 1		
23334222	**Good identification**					**23334222**
Actinomyces odontolyticus		100.0	0.43	TRE 1		
23334300	**Good identification**					**23334300**
Actinomyces odontolyticus		100.0	0.52	TRE 1		
23334302	**Good identification**					**23334302**
Actinomyces odontolyticus		100.0	0.48	TRE 1		
23334320	**Good identification**					**23334320**
Actinomyces odontolyticus		100.0	0.39	TRE 1		
23335000	**Good identification**					**23335000**
Actinomyces odontolyticus		98.75	0.53	TRE 1		
Actinomyces israelii		1.26	0.28			
23335002	**Good identification**					**23335002**
Actinomyces odontolyticus		96.98	0.49	TRE 1		
Actinomyces israelii		3.02	0.31			
23335020	**Good identification**					**23335020**
Actinomyces odontolyticus		100.0	0.39	TRE 1		
23335022	**Good identification**					**23335022**
Actinomyces odontolyticus		100.0	0.36	TRE 1		
23335100	**Low discrimination**					**23335100**
					COC	
Actinomyces odontolyticus		80.95	0.44	TRE 1	1	
Actinomyces israelii		13.03	0.39		1	
Atopobium parvulum		5.56	0.18		99	
23335102	**Low discrimination**					**23335102**
					COC	
Actinomyces odontolyticus		70.28	0.40	TRE 1	1	
Actinomyces israelii		27.68	0.42		1	
Atopobium parvulum		1.39	0.06		99	
23335200	**Very good identification**					**23335200**
Actinomyces odontolyticus		100.0	0.58	TRE 1		
23335202	**Good identification**					**23335202**
Actinomyces odontolyticus		100.0	0.54	TRE 1		
23335220	**Good identification**					**23335220**
Actinomyces odontolyticus		100.0	0.44	TRE 1		
23335222	**Good identification**					**23335222**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.40	TRE 1		
23335300	**Good identification**					**23335300**
Actinomyces odontolyticus		99.38	0.49	TRE 1		
Actinomyces israelii		0.62	0.19			
23335302	**Good identification**					**23335302**
Actinomyces odontolyticus		98.50	0.45	TRE 1		
Actinomyces israelii		1.50	0.22			
23335320	**Good identification**					**23335320**
Actinomyces odontolyticus		100.0	0.36	TRE 1		
23335500	**Acceptable identification**					**23335500**
Actinomyces israelii		92.88	0.37	CEL 93		
Actinomyces odontolyticus		7.12	0.10			
23335502	**Acceptable identification**					**23335502**
Actinomyces israelii		91.41	0.40	CEL 93		
Bifidobacterium adolescentis		5.73	0.13			
Actinomyces odontolyticus		2.87	0.06			
23337000	**Low discrimination**					**23337000**
Actinomyces israelii		87.70	0.41	CEL 93		
Actinomyces odontolyticus		12.30	0.19			
23337002	**Acceptable identification**					**23337002**
Actinomyces israelii		94.58	0.44	CEL 93		
Actinomyces odontolyticus		5.42	0.15			
23337042	**Good identification**					**23337042**
Actinomyces israelii		100.0	0.37	CEL 93		
23337100	**Acceptable identification**					**23337100**
Actinomyces israelii		93.69	0.51	CEL 93		
Propionibacterium propionicum		4.14	0.27			
Actinomyces naeslundii		1.13	0.28			
23337102	**Acceptable identification**					**23337102**
Actinomyces israelii		94.63	0.54	CEL 93		
Propionibacterium propionicum		4.18	0.30			
Actinomyces naeslundii		0.76	0.28			
23337110	**Good identification**					**23337110**
Actinomyces israelii		100.0	0.37	CEL 93		
23337112	**Good identification**					**23337112**
Actinomyces israelii		100.0	0.40	CEL 93		
23337140	**Low discrimination**					**23337140**
Actinomyces israelii		89.79	0.44	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		10.21	0.27			
23337142	**Low discrimination**					**23337142**
Actinomyces israelii		89.79	0.47	CEL 93		
Propionibacterium propionicum		10.21	0.30			
23337302	**Low discrimination**					**23337302**
Actinomyces israelii		89.55	0.35	CEL 93		
Actinomyces odontolyticus		10.45	0.11	RHA 7		
23337400	**Good identification**					**23337400**
Actinomyces israelii		100.0	0.40	CEL 93		
23337402	**Good identification**					**23337402**
Actinomyces israelii		100.0	0.43	CEL 93		
23337442	**Good identification**					**23337442**
Actinomyces israelii		100.0	0.36	CEL 93		
23337500	**Good identification**					**23337500**
Actinomyces israelii		100.0	0.50	CEL 93		
23337502	**Good identification**					**23337502**
Actinomyces israelii		99.10	0.53	CEL 93		
Bifidobacterium adolescentis		0.90	0.12			
23337510	**Good identification**					**23337510**
Actinomyces israelii		100.0	0.36	CEL 93		
23337512	**Acceptable identification**					**23337512**
Actinomyces israelii		93.11	0.39	CEL 93		
Bifidobacterium adolescentis		6.89	0.13	MLZ 13		
23337540	**Good identification**					**23337540**
Actinomyces israelii		100.0	0.43	CEL 93		
23337542	**Good identification**					**23337542**
Actinomyces israelii		100.0	0.46	CEL 93		
23340000	**Good identification**					**23340000**
Actinomyces odontolyticus		100.0	0.43	FRU 92		
23340002	**Good identification**					**23340002**
Actinomyces odontolyticus		95.14	0.39	FRU 92		
Actinomyces viscosus		4.86	0.10	CEL 1		
23340200	**Good identification**					**23340200**
Actinomyces odontolyticus		100.0	0.48	FRU 92		
23340202	**Good identification**					**23340202**
Actinomyces odontolyticus		100.0	0.44	FRU 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23340300	**Good identification**					**23340300**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23340302	**Good identification**					**23340302**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23341000	**Low discrimination**					**23341000**
Actinomyces odontolyticus		87.05	0.40	FRU 92		
Eubacterium tortuosum		11.85	0.14	CEL 1		
Actinomyces naeslundii		1.11	0.25			
23341002	**Acceptable identification**					**23341002**
Actinomyces odontolyticus		90.35	0.36	FRU 92		
Actinomyces viscosus		4.24	0.06	CEL 1		
Eubacterium tortuosum		3.54	0.01			
23341101	**Low discrimination**					**23341101**
					SPO	
Actinomyces naeslundii		69.73	0.35	MNS 88	1	
Clostridium chauvoei		30.27	0.06	RAF 92	99	
23341103	**Low discrimination**					**23341103**
Actinomyces naeslundii		76.07	0.35	MNS 88		
Actinomyces viscosus		23.93	0.03	RAF 92		
23341121	**Good identification**					**23341121**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23341123	**Good identification**					**23341123**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23341200	**Good identification**					**23341200**
Actinomyces odontolyticus		100.0	0.45	FRU 92		
23341202	**Good identification**					**23341202**
Actinomyces odontolyticus		100.0	0.41	FRU 92		
23341300	**Good identification**					**23341300**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23343100	**Good identification**					**23343100**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23343101	**Good identification**					**23343101**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23343102	**Good identification**					**23343102**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23343103	**Good identification**					**23343103**
Actinomyces naeslundii		100.0	0.40	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23343120	**Good identification**					**23343120**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23343121	**Good identification**					**23343121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23343122	**Good identification**					**23343122**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23343123	**Good identification**					**23343123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23344000	**Good identification**					**23344000**
Actinomyces odontolyticus		100.0	0.38	FRU 92		
23344200	**Good identification**					**23344200**
Actinomyces odontolyticus		100.0	0.42	FRU 92		
23344202	**Good identification**					**23344202**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23345200	**Good identification**					**23345200**
Actinomyces odontolyticus		100.0	0.39	FRU 92		
23345202	**Good identification**					**23345202**
Actinomyces odontolyticus		100.0	0.36	FRU 92		
23350000	**Very good identification**					**23350000**
					GRA	
Actinomyces odontolyticus		99.75	0.61	CEL 1	99	
Fusobacterium mortiferum		0.19	0.20		1	
Actinomyces naeslundii		0.06	0.23		99	
23350002	**Good identification**					**23350002**
Actinomyces odontolyticus		92.14	0.57	CEL 1		
Actinomyces viscosus		7.77	0.32			
Actinomyces naeslundii		0.09	0.23			
23350020	**Good identification**					**23350020**
Actinomyces odontolyticus		99.64	0.47	CEL 1		
Actinomyces naeslundii		0.36	0.23	NAG 14		
23350022	**Good identification**					**23350022**
Actinomyces odontolyticus		99.42	0.44	CEL 1		
Actinomyces naeslundii		0.58	0.23	NAG 14		
23350100	**Good identification**					**23350100**
					GRA	
Actinomyces odontolyticus		97.83	0.52	CEL 1	99	
Fusobacterium mortiferum		1.37	0.26		1	
Actinomyces naeslundii		0.51	0.31		99	
23350102	**Low discrimination**					**23350102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		67.42	0.49	CEL 1		
Actinomyces viscosus		32.00	0.36			
Actinomyces naeslundii		0.58	0.31			
23350120	**Good identification**					**23350120**
Actinomyces odontolyticus		96.88	0.39	CEL 1		
Actinomyces naeslundii		3.13	0.31	NAG 14		
23350122	**Acceptable identification**					**23350122**
Actinomyces odontolyticus		90.94	0.35	CEL 1		
Actinomyces naeslundii		4.79	0.31	NAG 14		
Actinomyces viscosus		2.68	0.02			
23350200	**Very good identification**					**23350200**
Actinomyces odontolyticus		100.0	0.66	CEL 1		
23350202	**Very good identification**					**23350202**
Actinomyces odontolyticus		100.0	0.62	CEL 1		
23350220	**Good identification**					**23350220**
Actinomyces odontolyticus		100.0	0.52	CEL 1		
23350222	**Good identification**					**23350222**
Actinomyces odontolyticus		100.0	0.49	CEL 1		
23350300	**Very good identification**					**23350300**
Actinomyces odontolyticus		100.0	0.57	CEL 1		
23350302	**Good identification**					**23350302**
Actinomyces odontolyticus		99.75	0.54	CEL 1		
Actinomyces viscosus		0.25	0.02			
23350320	**Good identification**					**23350320**
Actinomyces odontolyticus		100.0	0.44	CEL 1		
23350322	**Good identification**					**23350322**
Actinomyces odontolyticus		100.0	0.40	CEL 1		
23351000	**Good identification**					**23351000**
Actinomyces odontolyticus		92.62	0.58	CEL 1		
Lactobacillus catenaformis		3.24	0.29			
Actinomyces naeslundii		1.94	0.46			
23351001	**Low discrimination**					**23351001**
Actinomyces naeslundii		75.69	0.49	MNS 88		
Actinomyces odontolyticus		24.31	0.24	RAF 92		
23351002	**Low discrimination**					**23351002**
Actinomyces odontolyticus		83.42	0.54	CEL 1		
Actinomyces viscosus		6.47	0.28			
Lactobacillus catenaformis		4.76	0.29			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23351003	**Low discrimination**					**23351003**
Actinomyces naeslundii		58.81	0.49	MNS 88		
Actinomyces viscosus		29.61	0.20	RAF 92		
Actinomyces odontolyticus		11.58	0.20			
23351020	**Low discrimination**					**23351020**
Actinomyces odontolyticus		81.38	0.44	CEL 1		
Actinomyces naeslundii		10.48	0.46	NAG 14		
Lactobacillus catenaformis		4.37	0.18			
23351021	**Good identification**					**23351021**
Actinomyces naeslundii		95.03	0.49	MNS 88		
Actinomyces odontolyticus		4.97	0.10	RAF 92		
23351022	**Low discrimination**					**23351022**
Actinomyces odontolyticus		71.42	0.41	CEL 1		
Actinomyces naeslundii		15.01	0.46	NAG 14		
Lactobacillus catenaformis		6.25	0.18			
23351023	**Good identification**					**23351023**
Actinomyces naeslundii		96.90	0.49	MNS 88		
Actinomyces odontolyticus		3.11	0.07	RAF 92		
23351041	**Good identification**					**23351041**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23351043	**Good identification**					**23351043**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23351061	**Good identification**					**23351061**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23351063	**Good identification**					**23351063**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23351100	**Low discrimination**					**23351100**
Actinomyces odontolyticus		69.86	0.49	CEL 1		
Actinomyces naeslundii		13.21	0.54			
Lactobacillus catenaformis		11.60	0.32			
23351101	**Good identification**					**23351101**
					SPO	
Actinomyces naeslundii		91.83	0.57	MNS 88	1	
Clostridium chauvoei		4.90	0.12	RAF 92	99	
Actinomyces odontolyticus		3.27	0.15		1	
23351102	**Low discrimination**					**23351102**
Actinomyces odontolyticus		44.88	0.46	CEL 1		
Actinomyces viscosus		19.59	0.33			
Actinomyces naeslundii		13.84	0.54			
23351103	**Low discrimination**					**23351103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		74.31	0.57	MNS 88		
Actinomyces viscosus		23.37	0.24	RAF 92		
Actinomyces odontolyticus		1.62	0.12			
23351111	**Good identification**					**23351111**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23351113	**Good identification**					**23351113**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23351120	**Low discrimination**					**23351120**
Actinomyces naeslundii		42.08	0.54	MNS 88		
Actinomyces odontolyticus		36.24	0.36	RAF 92		
Lactobacillus catenaformis		9.24	0.21			
23351121	**Very good identification**					**23351121**
					SPO	
Actinomyces naeslundii		98.25	0.57	MNS 88	1	
Clostridium chauvoei		1.75	0.04	RAF 92	99	
23351122	**Low discrimination**					**23351122**
Actinomyces naeslundii		39.44	0.54	MNS 88		
Actinomyces odontolyticus		20.82	0.32	RAF 92		
Lactobacillus acidophilus		15.35	0.23			
23351123	**Very good identification**					**23351123**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23351131	**Good identification**					**23351131**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23351133	**Good identification**					**23351133**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23351140	**Low discrimination**					**23351140**
Actinomyces naeslundii		63.02	0.41	MNS 88		
Actinomyces odontolyticus		19.09	0.15	RAF 92		
Actinomyces israelii		17.89	0.23			
23351141	**Good identification**					**23351141**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23351142	**Low discrimination**					**23351142**
Actinomyces naeslundii		50.89	0.41	MNS 88		
Actinomyces israelii		21.67	0.26	RAF 92		
Lactobacillus acidophilus		13.87	0.07			
23351143	**Good identification**					**23351143**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23351160	**Low discrimination**					**23351160**
Actinomyces naeslundii		84.39	0.41	MNS 88		
Clostridium perfringens		15.61	0.06	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23351161	**Good identification**					**23351161**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23351162	**Low discrimination**					**23351162**
					SPO	
Actinomyces naeslundii		67.11	0.41	MNS 88	1	
Lactobacillus acidophilus		18.29	0.07	RAF 92	1	
Clostridium barati		8.76	-.01		99	
23351163	**Good identification**					**23351163**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23351200	**Very good identification**					**23351200**
Actinomyces odontolyticus		100.0	0.63	CEL 1		
23351202	**Very good identification**					**23351202**
Actinomyces odontolyticus		100.0	0.59	CEL 1		
23351220	**Good identification**					**23351220**
Actinomyces odontolyticus		100.0	0.49	CEL 1		
23351222	**Good identification**					**23351222**
Actinomyces odontolyticus		100.0	0.46	CEL 1		
23351300	**Good identification**					**23351300**
Actinomyces odontolyticus		100.0	0.54	CEL 1		
23351302	**Good identification**					**23351302**
Actinomyces odontolyticus		99.09	0.51	CEL 1		
Lactobacillus acidophilus		0.46	0.04			
Actinomyces israelii		0.23	0.14			
23351320	**Good identification**					**23351320**
Actinomyces odontolyticus		100.0	0.41	CEL 1		
23351322	**Good identification**					**23351322**
					SPO	
Actinomyces odontolyticus		95.14	0.37	CEL 1	1	
Lactobacillus acidophilus		2.72	0.04	NAG 14	1	
Clostridium barati		2.14	-.01		99	
23352101	**Low discrimination**					**23352101**
Actinomyces naeslundii		89.48	0.39	MNS 88		
Propionibacterium propionicum		10.52	0.12	RAF 92		
23352103	**Low discrimination**					**23352103**
Actinomyces naeslundii		85.00	0.39	MNS 88		
Propionibacterium propionicum		15.00	0.15	RAF 92		
23352120	**Good identification**					**23352120**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23352121	**Good identification**					**23352121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	MNS 88		
23352122	**Good identification**					**23352122**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23352123	**Good identification**					**23352123**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23353000	**Low discrimination**					**23353000**
Actinomyces naeslundii		65.98	0.51	MNS 88		
Actinomyces israelii		17.65	0.32	RAF 92		
Actinomyces odontolyticus		16.37	0.24			
23353001	**Good identification**					**23353001**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23353002	**Low discrimination**					**23353002**
Actinomyces naeslundii		64.38	0.51	MNS 88		
Actinomyces israelii		25.84	0.35	RAF 92		
Actinomyces odontolyticus		9.79	0.20			
23353003	**Good identification**					**23353003**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23353020	**Good identification**					**23353020**
Actinomyces naeslundii		96.12	0.51	MNS 88		
Actinomyces odontolyticus		3.88	0.10	RAF 92		
23353021	**Good identification**					**23353021**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23353022	**Good identification**					**23353022**
Actinomyces naeslundii		95.21	0.51	MNS 88		
Actinomyces israelii		2.44	0.15	RAF 92		
Actinomyces odontolyticus		2.36	0.07			
23353023	**Good identification**					**23353023**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23353040	**Low discrimination**					**23353040**
Actinomyces naeslundii		62.91	0.38	MNS 88		
Actinomyces israelii		37.09	0.25	RAF 92		
23353041	**Good identification**					**23353041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23353042	**Low discrimination**					**23353042**
Actinomyces naeslundii		53.07	0.38	MNS 88		
Actinomyces israelii		46.93	0.28	RAF 92		
23353043	**Good identification**					**23353043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MNS 88		
23353060	**Good identification**					**23353060**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23353061	**Good identification**					**23353061**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23353062	**Good identification**					**23353062**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23353063	**Good identification**					**23353063**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23353100	**Low discrimination**					**23353100**
Actinomyces naeslundii		68.04	0.59	MNS 88		
Actinomyces israelii		25.58	0.43	RAF 92		
Propionibacterium propionicum		4.50	0.29			
23353101	**Very good identification**					**23353101**
Actinomyces naeslundii		99.51	0.62	MNS 88		
Propionibacterium propionicum		0.49	0.12	RAF 92		
23353102	**Low discrimination**					**23353102**
Actinomyces naeslundii		58.89	0.59	MNS 88		
Actinomyces israelii		33.22	0.46	RAF 92		
Propionibacterium propionicum		5.85	0.32			
23353103	**Very good identification**					**23353103**
Actinomyces naeslundii		99.27	0.62	MNS 88		
Propionibacterium propionicum		0.73	0.15	RAF 92		
23353111	**Good identification**					**23353111**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23353113	**Good identification**					**23353113**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23353120	**Very good identification**					**23353120**
Actinomyces naeslundii		97.66	0.59	MNS 88		
Actinomyces israelii		2.34	0.22	RAF 92		
23353121	**Very good identification**					**23353121**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23353122	**Good identification**					**23353122**
					SPO	
Actinomyces naeslundii		93.68	0.59	MNS 88	1	
Actinomyces israelii		3.37	0.25	RAF 92	1	
Clostridium barati		1.11	-.01		99	
23353123	**Very good identification**					**23353123**
Actinomyces naeslundii		100.0	0.62	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23353130	**Good identification**					**23353130**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23353131	**Good identification**					**23353131**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23353132	**Low discrimination**					**23353132**
					SPO	
Actinomyces naeslundii		76.69	0.37	MNS 88	1	
Clostridium tertium		23.31	0.05	RAF 92	99	
23353133	**Good identification**					**23353133**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23353140	**Low discrimination**					**23353140**
Actinomyces naeslundii		45.38	0.46	MNS 88		
Actinomyces israelii		37.60	0.36	RAF 92		
Propionibacterium propionicum		17.02	0.29			
23353141	**Good identification**					**23353141**
Actinomyces naeslundii		97.30	0.49	MNS 88		
Propionibacterium propionicum		2.70	0.12	RAF 92		
23353142	**Low discrimination**					**23353142**
Actinomyces israelii		44.31	0.39	XYL 92		
Actinomyces naeslundii		35.65	0.46			
Propionibacterium propionicum		20.05	0.32			
23353143	**Good identification**					**23353143**
Actinomyces naeslundii		96.00	0.49	MNS 88		
Propionibacterium propionicum		4.00	0.15	RAF 92		
23353160	**Acceptable identification**					**23353160**
Actinomyces naeslundii		94.98	0.46	MNS 88		
Actinomyces israelii		5.02	0.15	RAF 92		
23353161	**Good identification**					**23353161**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23353162	**Acceptable identification**					**23353162**
Actinomyces naeslundii		92.65	0.46	MNS 88		
Actinomyces israelii		7.35	0.18	RAF 92		
23353163	**Good identification**					**23353163**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23353500	**Good identification**					**23353500**
Actinomyces israelii		96.82	0.41	XYL 92		
Actinomyces naeslundii		3.18	0.25			
23353502	**Good identification**					**23353502**
Actinomyces israelii		97.86	0.44	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		2.14	0.25			
23353542	**Good identification**					**23353542**
Actinomyces israelii		100.0	0.37	XYL 92		
23354000	**Good identification**					**23354000**
Actinomyces odontolyticus		100.0	0.56	CEL 1		
23354002	**Good identification**					**23354002**
Actinomyces odontolyticus		100.0	0.52	CEL 1		
23354020	**Good identification**					**23354020**
Actinomyces odontolyticus		100.0	0.42	CEL 1		
23354022	**Good identification**					**23354022**
Actinomyces odontolyticus		100.0	0.39	CEL 1		
23354100	**Good identification**					**23354100**
Actinomyces odontolyticus		99.62	0.47	CEL 1		
Actinomyces israelii		0.38	0.14			
23354102	**Good identification**					**23354102**
Actinomyces odontolyticus		98.13	0.43	CEL 1		
Actinomyces viscosus		0.96	0.02			
Actinomyces israelii		0.92	0.17			
23354200	**Very good identification**					**23354200**
Actinomyces odontolyticus		100.0	0.61	CEL 1		
23354202	**Very good identification**					**23354202**
Actinomyces odontolyticus		100.0	0.57	CEL 1		
23354220	**Good identification**					**23354220**
Actinomyces odontolyticus		100.0	0.47	CEL 1		
23354222	**Good identification**					**23354222**
Actinomyces odontolyticus		100.0	0.43	CEL 1		
23354300	**Good identification**					**23354300**
Actinomyces odontolyticus		100.0	0.52	CEL 1		
23354302	**Good identification**					**23354302**
Actinomyces odontolyticus		100.0	0.48	CEL 1		
23354320	**Good identification**					**23354320**
Actinomyces odontolyticus		100.0	0.39	CEL 1		
23355000	**Good identification**					**23355000**
Actinomyces odontolyticus		95.71	0.53	CEL 1		
Actinomyces israelii		4.30	0.38			
23355002	**Acceptable identification**					**23355002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		90.10	0.49	CEL 1		
Actinomyces israelii		9.90	0.41			
23355020	**Good identification**					**23355020**
Actinomyces odontolyticus		98.27	0.39	CEL 1		
Actinomyces israelii		1.73	0.17	NAG 14		
23355022	**Good identification**					**23355022**
Actinomyces odontolyticus		95.87	0.36	CEL 1		
Actinomyces israelii		4.13	0.20	NAG 14		
23355100	**Low discrimination**					**23355100**
Actinomyces odontolyticus		63.76	0.44	CEL 1		
Actinomyces israelii		36.24	0.48			
23355102	**Low discrimination**					**23355102**
Actinomyces israelii		57.73	0.51			
Actinomyces odontolyticus		41.50	0.40			
Bifidobacterium adolescentis		0.40	0.08			
23355112	**Acceptable identification**					**23355112**
Actinomyces israelii		90.48	0.37	MLZ 13		
Bifidobacterium adolescentis		5.12	0.09			
Actinomyces odontolyticus		4.40	0.06			
23355140	**Good identification**					**23355140**
Actinomyces israelii		95.63	0.41			
Actinomyces odontolyticus		4.37	0.10			
23355142	**Good identification**					**23355142**
Actinomyces israelii		98.17	0.44			
Actinomyces odontolyticus		1.83	0.06			
23355200	**Very good identification**					**23355200**
Actinomyces odontolyticus		99.83	0.58	CEL 1		
Actinomyces israelii		0.17	0.18			
23355202	**Good identification**					**23355202**
Actinomyces odontolyticus		99.58	0.54	CEL 1		
Actinomyces israelii		0.42	0.21			
23355220	**Good identification**					**23355220**
Actinomyces odontolyticus		100.0	0.44	CEL 1		
23355222	**Good identification**					**23355222**
Actinomyces odontolyticus		100.0	0.40	CEL 1		
23355300	**Good identification**					**23355300**
Actinomyces odontolyticus		97.84	0.49	CEL 1		
Actinomyces israelii		2.16	0.29			
23355302	**Acceptable identification**					**23355302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		94.88	0.45	CEL 1		
Actinomyces israelii		5.12	0.32			
23355320	**Good identification**					**23355320**
Actinomyces odontolyticus		100.0	0.36	CEL 1		
23355400	**Low discrimination**					**23355400**
Actinomyces israelii		78.42	0.36			
Actinomyces odontolyticus		21.58	0.19			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23355402	**Low discrimination**					**23355402**
Actinomyces israelii		89.90	0.39			
Actinomyces odontolyticus		10.11	0.15			
23355500	**Good identification**					**23355500**
Actinomyces israelii		97.87	0.46			
Actinomyces odontolyticus		2.13	0.10			
23355502	**Acceptable identification**					**23355502**
Actinomyces israelii		93.86	0.49			
Bifidobacterium adolescentis		5.31	0.22			
Actinomyces odontolyticus		0.83	0.06			
23355512	**Low discrimination**					**23355512**
Actinomyces israelii		68.37	0.35	MLZ 13		
Bifidobacterium adolescentis		31.64	0.23			
23355540	**Good identification**					**23355540**
Actinomyces israelii		100.0	0.39			
23355542	**Good identification**					**23355542**
Actinomyces israelii		100.0	0.42			
23357000	**Good identification**					**23357000**
Actinomyces israelii		96.18	0.51			
Actinomyces odontolyticus		3.82	0.19			
23357002	**Good identification**					**23357002**
Actinomyces israelii		98.40	0.54			
Actinomyces odontolyticus		1.60	0.15			
23357010	**Good identification**					**23357010**
Actinomyces israelii		100.0	0.36	MLZ 13		
23357012	**Good identification**					**23357012**
Actinomyces israelii		100.0	0.39	MLZ 13		
23357040	**Good identification**					**23357040**
Actinomyces israelii		100.0	0.43			
23357042	**Good identification**					**23357042**
Actinomyces israelii		100.0	0.47			
23357100	**Very good identification**					**23357100**
Actinomyces israelii		99.46	0.61			
Actinomyces odontolyticus		0.31	0.10			
Actinomyces naeslundii		0.23	0.25			
23357102	**Very good identification**					**23357102**
Actinomyces israelii		99.72	0.64			
Actinomyces naeslundii		0.16	0.25			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		0.13	0.06			
23357110	**Good identification**					**23357110**
Actinomyces israelii		100.0	0.47	MLZ 13		
23357112	**Good identification**					**23357112**
					SPO	
Actinomyces israelii		98.45	0.50	MLZ 13	1	
Bifidobacterium adolescentis		0.80	0.08		1	
Clostridium tertium		0.75	-.01		99	
23357120	**Good identification**					**23357120**
Actinomyces israelii		96.47	0.40	NAG 6		
Actinomyces naeslundii		3.53	0.25			
23357122	**Acceptable identification**					**23357122**
					SPO	
Actinomyces israelii		94.84	0.43	NAG 6	1	
Clostridium tertium		2.85	0.03		99	
Actinomyces naeslundii		2.31	0.25		1	
23357140	**Good identification**					**23357140**
Actinomyces israelii		100.0	0.54			
23357142	**Very good identification**					**23357142**
Actinomyces israelii		100.0	0.57			
23357150	**Good identification**					**23357150**
Actinomyces israelii		100.0	0.40	MLZ 13		
23357152	**Good identification**					**23357152**
Actinomyces israelii		100.0	0.43	MLZ 13		
23357162	**Good identification**					**23357162**
Actinomyces israelii		100.0	0.36	NAG 6		
23357300	**Acceptable identification**					**23357300**
Actinomyces israelii		92.52	0.42	RHA 7		
Actinomyces odontolyticus		7.48	0.15			
23357302	**Good identification**					**23357302**
Actinomyces israelii		96.80	0.45	RHA 7		
Actinomyces odontolyticus		3.20	0.11			
23357342	**Good identification**					**23357342**
Actinomyces israelii		100.0	0.38	RHA 7		
23357400	**Good identification**					**23357400**
Actinomyces israelii		100.0	0.49			
23357402	**Good identification**					**23357402**
Actinomyces israelii		100.0	0.52			
23357412	**Good identification**					**23357412**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.38	MLZ 13		
23357440	**Good identification**					**23357440**
Actinomyces israelii		100.0	0.42			
23357442	**Good identification**					**23357442**
Actinomyces israelii		100.0	0.45			
23357500	**Very good identification**					**23357500**
Actinomyces israelii		100.0	0.59			
23357502	**Very good identification**					**23357502**
Actinomyces israelii		99.19	0.62			
Bifidobacterium adolescentis		0.81	0.20			
23357510	**Good identification**					**23357510**
Actinomyces israelii		100.0	0.45	MLZ 13		
23357512	**Acceptable identification**					**23357512**
Actinomyces israelii		93.74	0.48	MLZ 13		
Bifidobacterium adolescentis		6.26	0.22			
23357520	**Good identification**					**23357520**
Actinomyces israelii		100.0	0.39	NAG 6		
23357522	**Good identification**					**23357522**
Actinomyces israelii		100.0	0.42	NAG 6		
23357540	**Good identification**					**23357540**
Actinomyces israelii		100.0	0.52			
23357542	**Good identification**					**23357542**
Actinomyces israelii		100.0	0.55			
23357550	**Good identification**					**23357550**
Actinomyces israelii		100.0	0.38	MLZ 13		
23357552	**Good identification**					**23357552**
Actinomyces israelii		100.0	0.41	MLZ 13		
23357700	**Good identification**					**23357700**
Actinomyces israelii		100.0	0.40	RHA 7		
23357702	**Good identification**					**23357702**
Actinomyces israelii		100.0	0.43	RHA 7		
23357742	**Good identification**					**23357742**
Actinomyces israelii		100.0	0.36	RHA 7		
23361100	**Good identification**					**23361100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	MNS 88		
23361101	**Good identification**					**23361101**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23361102	**Good identification**					**23361102**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23361103	**Good identification**					**23361103**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23361120	**Good identification**					**23361120**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23361121	**Good identification**					**23361121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23361122	**Good identification**					**23361122**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23361123	**Good identification**					**23361123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23363001	**Good identification**					**23363001**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23363003	**Good identification**					**23363003**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23363021	**Good identification**					**23363021**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23363023	**Good identification**					**23363023**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23363100	**Low discrimination**					**23363100**
Actinomyces naeslundii		87.73	0.42	MNS 88		
Actinomyces israelii		12.27	0.18	RAF 92		
23363101	**Good identification**					**23363101**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23363102	**Low discrimination**					**23363102**
Actinomyces naeslundii		82.66	0.42	MNS 88		
Actinomyces israelii		17.34	0.21	RAF 92		
23363103	**Good identification**					**23363103**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23363120	**Good identification**					**23363120**
Actinomyces naeslundii		100.0	0.42	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23363121	**Good identification**					**23363121**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23363122	**Good identification**					**23363122**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23363123	**Good identification**					**23363123**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23367100	**Good identification**					**23367100**
Actinomyces israelii		100.0	0.37	FRU 99		
23367102	**Good identification**					**23367102**
Actinomyces israelii		100.0	0.40	FRU 99		
23367500	**Good identification**					**23367500**
Actinomyces israelii		100.0	0.35	FRU 99		
23367502	**Good identification**					**23367502**
Actinomyces israelii		100.0	0.38	FRU 99		
23370100	**Low discrimination**					**23370100**
					GRA	
Actinomyces naeslundii		41.64	0.36	MNS 88	99	
Actinomyces odontolyticus		41.27	0.18	RAF 92	99	
Fusobacterium mortiferum		17.10	0.17	GAL 96	1	
23370101	**Good identification**					**23370101**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23370103	**Low discrimination**					**23370103**
Actinomyces naeslundii		67.68	0.39	MNS 88		
Actinomyces viscosus		32.32	0.09	RAF 92		
23370120	**Good identification**					**23370120**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23370121	**Good identification**					**23370121**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23370122	**Good identification**					**23370122**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23370123	**Good identification**					**23370123**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23371001	**Good identification**					**23371001**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23371002	**Low discrimination**					**23371002**
Actinomyces naeslundii		39.46	0.51	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		34.01	0.29	RAF 92		
Actinomyces israelii		10.51	0.32			
23371003	**Good identification**					**23371003**
Actinomyces naeslundii		98.09	0.54	MNS 88		
Actinomyces viscosus		1.92	0.01	RAF 92		
23371020	**Low discrimination**					**23371020**
Actinomyces naeslundii		70.06	0.51	MNS 88		
Lactobacillus jensenii		22.65	0.21	RAF 92		
Clostridium perfringens		4.46	0.09			
23371021	**Good identification**					**23371021**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23371022	**Low discrimination**					**23371022**
Actinomyces naeslundii		69.62	0.51	MNS 88		
Lactobacillus jensenii		15.00	0.18	RAF 92		
Lactobacillus acidophilus		11.57	0.14			
23371023	**Good identification**					**23371023**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23371040	**Low discrimination**					**23371040**
Actinomyces naeslundii		71.87	0.38	MNS 88		
Actinomyces israelii		28.13	0.22	RAF 92		
23371041	**Good identification**					**23371041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23371042	**Low discrimination**					**23371042**
Actinomyces naeslundii		58.70	0.38	MNS 88		
Actinomyces israelii		34.47	0.25	RAF 92		
Lactobacillus acidophilus		6.83	-.02			
23371043	**Good identification**					**23371043**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23371060	**Good identification**					**23371060**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23371061	**Good identification**					**23371061**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23371062	**Low discrimination**					**23371062**
Actinomyces naeslundii		89.58	0.38	MNS 88		
Lactobacillus acidophilus		10.42	-.02	RAF 92		
23371063	**Good identification**					**23371063**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23371100	**Low discrimination**					**23371100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		70.49	0.59	MNS 88	1	
Actinomyces israelii		17.59	0.40	RAF 92	1	
Atopobium parvulum		8.17	0.20		99	
23371101	**Very good identification**					**23371101**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23371102	**Low discrimination**					**23371102**
Actinomyces naeslundii		46.77	0.59	MNS 88		
Lactobacillus acidophilus		31.40	0.32	RAF 92		
Actinomyces israelii		17.51	0.43			
23371103	**Very good identification**					**23371103**
Actinomyces naeslundii		98.80	0.62	MNS 88		
Actinomyces viscosus		1.21	0.05	RAF 92		
23371110	**Low discrimination**					**23371110**
Actinomyces naeslundii		58.52	0.37	MNS 88		
Actinomyces israelii		41.48	0.25	RAF 92		
23371111	**Good identification**					**23371111**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23371113	**Good identification**					**23371113**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23371120	**Low discrimination**					**23371120**
					COC	
Actinomyces naeslundii		83.31	0.59	MNS 88	1	
Clostridium perfringens		13.66	0.23	RAF 92	1	
Atopobium parvulum		1.70	0.07		99	
23371121	**Very good identification**					**23371121**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23371122	**Low discrimination**					**23371122**
Actinomyces naeslundii		55.49	0.59	MNS 88		
Lactobacillus acidophilus		37.25	0.32	RAF 92		
Clostridium perfringens		4.28	0.18			
23371123	**Very good identification**					**23371123**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23371130	**Good identification**					**23371130**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23371131	**Good identification**					**23371131**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23371132	**Low discrimination**					**23371132**
					GRA	
Actinomyces naeslundii		63.09	0.37	MNS 88	99	
Leptotrichia buccalis		36.91	-.01	RAF 92	1	
23371133	**Good identification**					**23371133**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MNS 88		
23371140	**Low discrimination**					**23371140**
Actinomyces naeslundii		64.51	0.46	MNS 88		
Actinomyces israelii		35.49	0.33	RAF 92		
23371141	**Good identification**					**23371141**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23371142	**Low discrimination**					**23371142**
Actinomyces naeslundii		43.57	0.46	MNS 88		
Actinomyces israelii		35.95	0.36	RAF 92		
Lactobacillus acidophilus		20.49	0.16			
23371143	**Good identification**					**23371143**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23371160	**Low discrimination**					**23371160**
Actinomyces naeslundii		84.97	0.46	MNS 88		
Clostridium perfringens		15.03	0.11	RAF 92		
23371161	**Good identification**					**23371161**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23371162	**Low discrimination**					**23371162**
Actinomyces naeslundii		62.26	0.46	MNS 88		
Lactobacillus acidophilus		29.28	0.16	RAF 92		
Clostridium perfringens		5.18	0.05			
23371163	**Good identification**					**23371163**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23371500	**Good identification**					**23371500**
Actinomyces israelii		95.29	0.38	XYL 92		
Actinomyces naeslundii		4.71	0.25			
23371502	**Good identification**					**23371502**
Actinomyces israelii		96.81	0.41	XYL 92		
Actinomyces naeslundii		3.19	0.25			
23372001	**Good identification**					**23372001**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23372003	**Good identification**					**23372003**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23372021	**Good identification**					**23372021**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23372023	**Good identification**					**23372023**
Actinomyces naeslundii		100.0	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23372100	**Low discrimination**					**23372100**
Actinomyces naeslundii		54.17	0.40	MNS 88		
Propionibacterium propionicum		36.26	0.27	RAF 92		
Actinomyces israelii		9.57	0.18	GAL 96		
23372101	**Good identification**					**23372101**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23372103	**Acceptable identification**					**23372103**
Actinomyces naeslundii		93.08	0.43	MNS 88		
Propionibacterium propionicum		6.92	0.14	RAF 92		
23372120	**Good identification**					**23372120**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23372121	**Good identification**					**23372121**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23372122	**Good identification**					**23372122**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23372123	**Good identification**					**23372123**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23373000	**Low discrimination**					**23373000**
Actinomyces naeslundii		65.86	0.56	MNS 88		
Actinomyces israelii		34.14	0.42	RAF 92		
23373001	**Very good identification**					**23373001**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23373002	**Low discrimination**					**23373002**
Actinomyces naeslundii		56.25	0.56	MNS 88		
Actinomyces israelii		43.75	0.45	RAF 92		
23373003	**Very good identification**					**23373003**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23373011	**Good identification**					**23373011**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23373013	**Good identification**					**23373013**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23373020	**Very good identification**					**23373020**
Actinomyces naeslundii		96.80	0.56	MNS 88		
Actinomyces israelii		3.20	0.22	RAF 92		
23373021	**Very good identification**					**23373021**
Actinomyces naeslundii		100.0	0.59	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23373022	**Very good identification**					**23373022**
Actinomyces naeslundii		95.27	0.56	MNS 88		
Actinomyces israelii		4.73	0.25	RAF 92		
23373023	**Very good identification**					**23373023**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23373031	**Good identification**					**23373031**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23373033	**Good identification**					**23373033**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23373040	**Low discrimination**					**23373040**
Actinomyces israelii		53.33	0.35	XYL 92		
Actinomyces naeslundii		46.67	0.43			
23373041	**Good identification**					**23373041**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23373042	**Low discrimination**					**23373042**
Actinomyces israelii		63.15	0.38	XYL 92		
Actinomyces naeslundii		36.85	0.43			
23373043	**Good identification**					**23373043**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23373060	**Acceptable identification**					**23373060**
Actinomyces naeslundii		93.20	0.43	MNS 88		
Actinomyces israelii		6.80	0.15	RAF 92		
23373061	**Good identification**					**23373061**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23373062	**Acceptable identification**					**23373062**
Actinomyces naeslundii		90.14	0.43	MNS 88		
Actinomyces israelii		9.86	0.18	RAF 92		
23373063	**Good identification**					**23373063**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23373100	**Low discrimination**					**23373100**
Actinomyces naeslundii		56.78	0.64	MNS 88		
Actinomyces israelii		41.37	0.53	RAF 92		
Propionibacterium propionicum		1.58	0.27			
23373101	**Very good identification**					**23373101**
Actinomyces naeslundii		99.51	0.67	MNS 88		
Actinomyces israelii		0.49	0.18	RAF 92		
23373102	**Low discrimination**					**23373102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		50.44	0.56	XYL 92		
Actinomyces naeslundii		46.15	0.64			
Propionibacterium propionicum		1.93	0.30			
23373103	**Very good identification**					**23373103**
Actinomyces naeslundii		98.97	0.67	MNS 88		
Actinomyces israelii		0.73	0.21	RAF 92		
Propionibacterium propionicum		0.31	0.14			
23373110	**Low discrimination**					**23373110**
					SPO	
Actinomyces israelii		64.14	0.38	XYL 92	1	
Actinomyces naeslundii		31.01	0.42	MLZ 13	1	
Clostridium tertium		4.85	0.05		99	
23373111	**Good identification**					**23373111**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23373112	**Low discrimination**					**23373112**
					SPO	
Actinomyces israelii		67.86	0.41	XYL 92	1	
Actinomyces naeslundii		21.87	0.42	MLZ 13	1	
Clostridium tertium		10.27	0.13		99	
23373113	**Good identification**					**23373113**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23373120	**Good identification**					**23373120**
					SPO	
Actinomyces naeslundii		93.15	0.64	MNS 88	1	
Actinomyces israelii		4.33	0.32	RAF 92	1	
Clostridium tertium		1.30	0.09		99	
23373121	**Very good identification**					**23373121**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23373122	**Low discrimination**					**23373122**
					SPO	
Actinomyces naeslundii		84.60	0.64	MNS 88	1	
Actinomyces israelii		5.90	0.35	RAF 92	1	
Clostridium ramosum		4.44	0.17		92	
23373123	**Very good identification**					**23373123**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23373130	**Low discrimination**					**23373130**
					SPO	
Actinomyces naeslundii		62.43	0.42	MNS 88	1	
Clostridium tertium		29.32	0.13	RAF 92	99	
Actinomyces israelii		8.24	0.18	MLZ 5	1	
23373131	**Good identification**					**23373131**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23373133	**Good identification**					**23373133**
Actinomyces naeslundii		100.0	0.45	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23373140	**Low discrimination**					**23373140**
Actinomyces israelii		58.10	0.45	XYL 92		
Actinomyces naeslundii		36.18	0.51			
Propionibacterium propionicum		5.72	0.27			
23373141	**Good identification**					**23373141**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23373142	**Low discrimination**					**23373142**
Actinomyces israelii		66.07	0.49	XYL 92		
Actinomyces naeslundii		27.43	0.51			
Propionibacterium propionicum		6.50	0.30			
23373143	**Good identification**					**23373143**
Actinomyces naeslundii		96.73	0.54	MNS 88		
Propionibacterium propionicum		1.70	0.14	RAF 92		
Actinomyces israelii		1.57	0.14			
23373160	**Acceptable identification**					**23373160**
Actinomyces naeslundii		90.70	0.51	MNS 88		
Actinomyces israelii		9.30	0.25	RAF 92		
23373161	**Good identification**					**23373161**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23373162	**Low discrimination**					**23373162**
Actinomyces naeslundii		86.68	0.51	MNS 88		
Actinomyces israelii		13.33	0.28	RAF 92		
23373163	**Good identification**					**23373163**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23373302	**Low discrimination**					**23373302**
Actinomyces israelii		89.07	0.36	XYL 92		
Actinomyces naeslundii		10.94	0.30	RHA 7		
23373400	**Good identification**					**23373400**
Actinomyces israelii		100.0	0.41	XYL 92		
23373402	**Good identification**					**23373402**
Actinomyces israelii		100.0	0.44	XYL 92		
23373442	**Good identification**					**23373442**
Actinomyces israelii		100.0	0.37	XYL 92		
23373500	**Good identification**					**23373500**
Actinomyces israelii		98.33	0.51	XYL 92		
Actinomyces naeslundii		1.67	0.30			
23373502	**Good identification**					**23373502**
Actinomyces israelii		98.88	0.54	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		1.12	0.30			
23373510	**Good identification**					**23373510**
Actinomyces israelii		100.0	0.37	XYL 92		
23373512	**Good identification**					**23373512**
Actinomyces israelii		100.0	0.40	XYL 92		
23373540	**Good identification**					**23373540**
Actinomyces israelii		100.0	0.44	XYL 92		
23373542	**Good identification**					**23373542**
Actinomyces israelii		100.0	0.47	XYL 92		
23375000	**Acceptable identification**					**23375000**
Actinomyces israelii		94.35	0.47			
Actinomyces odontolyticus		5.65	0.19			
23375002	**Good identification**					**23375002**
Actinomyces israelii		97.61	0.50			
Actinomyces odontolyticus		2.39	0.15			
23375012	**Good identification**					**23375012**
Actinomyces israelii		100.0	0.36	MLZ 13		
23375040	**Good identification**					**23375040**
Actinomyces israelii		100.0	0.40			
23375042	**Good identification**					**23375042**
Actinomyces israelii		100.0	0.43			
23375100	**Good identification**					**23375100**
					COC	
Actinomyces israelii		92.32	0.58		1	
Atopobium parvulum		6.92	0.25		99	
Actinomyces odontolyticus		0.44	0.10		1	
23375102	**Very good identification**					**23375102**
					COC	
Actinomyces israelii		98.71	0.61		1	
Atopobium parvulum		0.87	0.12		99	
Actinomyces naeslundii		0.23	0.25		1	
23375110	**Good identification**					**23375110**
Actinomyces israelii		100.0	0.44	MLZ 13		
23375112	**Good identification**					**23375112**
Actinomyces israelii		100.0	0.47	MLZ 13		
23375120	**Low discrimination**					**23375120**
					COC	
Actinomyces israelii		79.21	0.37	NAG 6	1	
Atopobium parvulum		16.43	0.12		99	
Actinomyces naeslundii		4.37	0.25		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23375122	**Good identification**					**23375122**
Actinomyces israelii		96.46	0.40	NAG 6		
Actinomyces naeslundii		3.55	0.25			
23375140	**Good identification**					**23375140**
Actinomyces israelii		100.0	0.51			
23375142	**Good identification**					**23375142**
Actinomyces israelii		100.0	0.54			
23375150	**Good identification**					**23375150**
Actinomyces israelii		100.0	0.37	MLZ 13		
23375152	**Good identification**					**23375152**
Actinomyces israelii		100.0	0.40	MLZ 13		
23375300	**Low discrimination**					**23375300**
Actinomyces israelii		89.14	0.39	RHA 7		
Actinomyces odontolyticus		10.86	0.15			
23375302	**Good identification**					**23375302**
Actinomyces israelii		95.26	0.42	RHA 7		
Actinomyces odontolyticus		4.74	0.11			
23375400	**Good identification**					**23375400**
Actinomyces israelii		100.0	0.46			
23375402	**Good identification**					**23375402**
Actinomyces israelii		100.0	0.49			
23375440	**Good identification**					**23375440**
Actinomyces israelii		100.0	0.39			
23375442	**Good identification**					**23375442**
Actinomyces israelii		100.0	0.42			
23375500	**Very good identification**					**23375500**
					SPO	
Actinomyces israelii		99.55	0.56		1	
Clostridium butyricum		0.45	-.05		99	
23375502	**Very good identification**					**23375502**
Actinomyces israelii		99.01	0.59			
Bifidobacterium adolescentis		0.99	0.19			
23375510	**Good identification**					**23375510**
Actinomyces israelii		100.0	0.42	MLZ 13		
23375512	**Acceptable identification**					**23375512**
Actinomyces israelii		92.42	0.45	MLZ 13		
Bifidobacterium adolescentis		7.58	0.20			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23375520	**Good identification**					**23375520**
Actinomyces israelii		100.0	0.36	NAG 6		
23375522	**Good identification**					**23375522**
Actinomyces israelii		100.0	0.39	NAG 6		
23375540	**Good identification**					**23375540**
Actinomyces israelii		100.0	0.49			
23375542	**Good identification**					**23375542**
Actinomyces israelii		100.0	0.52			
23375550	**Good identification**					**23375550**
Actinomyces israelii		100.0	0.35	MLZ 13		
23375552	**Good identification**					**23375552**
Actinomyces israelii		100.0	0.38	MLZ 13		
23375700	**Good identification**					**23375700**
Actinomyces israelii		100.0	0.37	RHA 7		
23375702	**Good identification**					**23375702**
Actinomyces israelii		100.0	0.40	RHA 7		
23376100	**Good identification**					**23376100**
Actinomyces israelii		100.0	0.37	GAL 99		
23376102	**Good identification**					**23376102**
Actinomyces israelii		100.0	0.40	GAL 99		
23376500	**Good identification**					**23376500**
Actinomyces israelii		100.0	0.35	GAL 99		
23376502	**Good identification**					**23376502**
Actinomyces israelii		100.0	0.38	GAL 99		
23377000	**Very good identification**					**23377000**
Actinomyces israelii		100.0	0.60			
23377002	**Very good identification**					**23377002**
Actinomyces israelii		100.0	0.63			
23377010	**Good identification**					**23377010**
Actinomyces israelii		100.0	0.46	MLZ 13		
23377012	**Good identification**					**23377012**
Actinomyces israelii		100.0	0.49	MLZ 13		
23377020	**Good identification**					**23377020**
Actinomyces israelii		100.0	0.40	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23377022	**Good identification**					**23377022**
Actinomyces israelii		100.0	0.43	NAG 6		
23377040	**Good identification**					**23377040**
Actinomyces israelii		100.0	0.53			
23377042	**Very good identification**					**23377042**
Actinomyces israelii		100.0	0.56			
23377050	**Good identification**					**23377050**
Actinomyces israelii		100.0	0.39	MLZ 13		
23377052	**Good identification**					**23377052**
Actinomyces israelii		100.0	0.42	MLZ 13		
23377062	**Good identification**					**23377062**
Actinomyces israelii		100.0	0.36	NAG 6		
23377100	**Very good identification**					**23377100**
					SPO	
Actinomyces israelii		99.80	0.71		1	
Actinomyces naeslundii		0.12	0.30		1	
Clostridium tertium		0.08	0.03		99	
23377101	**Low discrimination**					**23377101**
Actinomyces israelii		84.82	0.37	URE 1		
Actinomyces naeslundii		15.18	0.33			
23377102	**Very good identification**					**23377102**
					SPO	
Actinomyces israelii		99.77	0.74		1	
Clostridium tertium		0.15	0.11		99	
Actinomyces naeslundii		0.08	0.30		1	
23377103	**Low discrimination**					**23377103**
Actinomyces israelii		89.34	0.40	URE 1		
Actinomyces naeslundii		10.66	0.33			
23377110	**Very good identification**					**23377110**
					SPO	
Actinomyces israelii		99.10	0.57	MLZ 13	1	
Clostridium tertium		0.90	0.07		99	
23377112	**Very good identification**					**23377112**
					SPO	
Actinomyces israelii		98.22	0.60	MLZ 13	1	
Clostridium tertium		1.79	0.15		99	
23377120	**Acceptable identification**					**23377120**
					SPO	
Actinomyces israelii		94.81	0.50	NAG 6	1	
Clostridium tertium		3.40	0.11		99	
Actinomyces naeslundii		1.79	0.30		1	
23377122	**Acceptable identification**					**23377122**
					SPO	
Actinomyces israelii		92.22	0.53	NAG 6	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		6.62	0.19		99	
Actinomyces naeslundii		1.16	0.30		1	
23377130	**Low discrimination**					**23377130**
					SPO	
Actinomyces israelii		70.07	0.36	MLZ 13	1	
Clostridium tertium		29.93	0.15	NAG 6	99	
23377132	**Low discrimination**					**23377132**
					SPO	
Actinomyces israelii		53.93	0.39	MLZ 13	1	
Clostridium tertium		46.07	0.24	NAG 6	99	
23377140	**Very good identification**					**23377140**
Actinomyces israelii		100.0	0.64			
23377142	**Very good identification**					**23377142**
Actinomyces israelii		100.0	0.67			
23377150	**Good identification**					**23377150**
Actinomyces israelii		100.0	0.50	MLZ 13		
23377152	**Good identification**					**23377152**
Actinomyces israelii		100.0	0.53	MLZ 13		
23377160	**Good identification**					**23377160**
Actinomyces israelii		100.0	0.43	NAG 6		
23377162	**Good identification**					**23377162**
Actinomyces israelii		100.0	0.46	NAG 6		
23377200	**Good identification**					**23377200**
Actinomyces israelii		100.0	0.41	RHA 7		
23377202	**Good identification**					**23377202**
Actinomyces israelii		100.0	0.44	RHA 7		
23377242	**Good identification**					**23377242**
Actinomyces israelii		100.0	0.37	RHA 7		
23377300	**Good identification**					**23377300**
Actinomyces israelii		100.0	0.51	RHA 7		
23377302	**Good identification**					**23377302**
Actinomyces israelii		100.0	0.54	RHA 7		
23377310	**Good identification**					**23377310**
Actinomyces israelii		100.0	0.37	RHA 7		
23377312	**Good identification**					**23377312**
Actinomyces israelii		100.0	0.40	RHA 7		
23377340	**Good identification**					**23377340**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.44	RHA 7		
23377342	**Good identification**					**23377342**
Actinomyces israelii		100.0	0.47	RHA 7		
23377400	**Very good identification**					**23377400**
Actinomyces israelii		100.0	0.59			
23377402	**Very good identification**					**23377402**
Actinomyces israelii		100.0	0.62			
23377410	**Good identification**					**23377410**
Actinomyces israelii		100.0	0.45	MLZ 13		
23377412	**Good identification**					**23377412**
Actinomyces israelii		100.0	0.48	MLZ 13		
23377420	**Good identification**					**23377420**
Actinomyces israelii		100.0	0.38	NAG 6		
23377422	**Good identification**					**23377422**
Actinomyces israelii		100.0	0.41	NAG 6		
23377440	**Good identification**					**23377440**
Actinomyces israelii		100.0	0.52			
23377442	**Good identification**					**23377442**
Actinomyces israelii		100.0	0.55			
23377450	**Good identification**					**23377450**
Actinomyces israelii		100.0	0.38	MLZ 13		
23377452	**Good identification**					**23377452**
Actinomyces israelii		100.0	0.41	MLZ 13		
23377500	**Very good identification**					**23377500**
Actinomyces israelii		100.0	0.69			
23377501	**Good identification**					**23377501**
Actinomyces israelii		100.0	0.35	URE 1		
23377502	**Very good identification**					**23377502**
Actinomyces israelii		99.86	0.72			
Bifidobacterium adolescentis		0.15	0.17			
23377503	**Good identification**					**23377503**
Actinomyces israelii		100.0	0.38	URE 1		
23377510	**Good identification**					**23377510**
Actinomyces israelii		100.0	0.55	MLZ 13		
23377512	**Very good identification**					**23377512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		98.83	0.58	MLZ 13		
Bifidobacterium adolescentis		1.17	0.19			
23377520	**Good identification**					**23377520**
Actinomyces israelii		100.0	0.49	NAG 6		
23377522	**Good identification**					**23377522**
Actinomyces israelii		100.0	0.52	NAG 6		
23377532	**Good identification**					**23377532**
Actinomyces israelii		100.0	0.38	MLZ 13		
23377540	**Very good identification**					**23377540**
Actinomyces israelii		100.0	0.62			
23377542	**Very good identification**					**23377542**
Actinomyces israelii		100.0	0.65			
23377550	**Good identification**					**23377550**
Actinomyces israelii		100.0	0.48	MLZ 13		
23377552	**Good identification**					**23377552**
Actinomyces israelii		100.0	0.51	MLZ 13		
23377560	**Good identification**					**23377560**
Actinomyces israelii		100.0	0.42	NAG 6		
23377562	**Good identification**					**23377562**
Actinomyces israelii		100.0	0.45	NAG 6		
23377600	**Good identification**					**23377600**
Actinomyces israelii		100.0	0.40	RHA 7		
23377602	**Good identification**					**23377602**
Actinomyces israelii		100.0	0.43	RHA 7		
23377642	**Good identification**					**23377642**
Actinomyces israelii		100.0	0.36	RHA 7		
23377700	**Good identification**					**23377700**
Actinomyces israelii		100.0	0.50	RHA 7		
23377702	**Good identification**					**23377702**
Actinomyces israelii		100.0	0.53	RHA 7		
23377710	**Good identification**					**23377710**
Actinomyces israelii		100.0	0.36	RHA 7		
23377712	**Good identification**					**23377712**
Actinomyces israelii		100.0	0.39	RHA 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23377740	**Good identification**						**23377740**
Actinomyces israelii		100.0	0.43	RHA 7			
23377742	**Good identification**						**23377742**
Actinomyces israelii		100.0	0.46	RHA 7			
23410000	**Low discrimination**						**23410000**
					COC	GRA	
Fusobacterium mortiferum		48.46	0.44	NIT 1	1	1	
Megasphaera elsdenii		21.18	0.29		99	1	
Actinomyces odontolyticus		16.78	0.30		1	99	
23410002	**Low discrimination**						**23410002**
					GRA		
Actinomyces viscosus		84.98	0.38	MNS 96	99		
Actinomyces odontolyticus		7.17	0.27	MLT 99	99		
Fusobacterium mortiferum		3.76	0.27		1		
23410100	**Low discrimination**						**23410100**
					GRA		
Fusobacterium mortiferum		82.50	0.50	NIT 1	1		
Actinomyces meyeri		7.50	0.23		99		
Actinomyces odontolyticus		3.87	0.22		99		
23410102	**Acceptable identification**						**23410102**
					GRA		
Actinomyces viscosus		90.10	0.42	MNS 96	99		
Fusobacterium mortiferum		5.23	0.34	MLT 99	1		
Propionibacterium propionicum		2.24	0.29		99		
23410200	**Good identification**						**23410200**
Actinomyces odontolyticus		100.0	0.35	MLT 99			
23411000	**Low discrimination**						**23411000**
					GRA		
Lactobacillus fermentum		62.26	0.54	NIT 1	99		
Fusobacterium mortiferum		28.39	0.47		1		
Actinomyces odontolyticus		4.37	0.27		99		
23411001	**Low discrimination**						**23411001**
					GRA		
Actinomyces naeslundii		59.53	0.36	MNS 88	99		
Lactobacillus fermentum		27.79	0.20	MLT 94	99		
Fusobacterium mortiferum		12.67	0.13		1		
23411002	**Low discrimination**						**23411002**
					GRA		
Lactobacillus fermentum		63.11	0.54	NIT 1	99		
Actinomyces viscosus		29.57	0.34		99		
Fusobacterium mortiferum		3.20	0.30		1		
23411020	**Low discrimination**						**23411020**
					CAT		
Lactobacillus fermentum		85.48	0.46	NIT 1	1		
Propionibacterium acnes		4.69	0.10		95		
Actinomyces naeslundii		3.70	0.33		25		
23411021	**Good identification**						**23411021**
Actinomyces naeslundii		100.0	0.36	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23411022	**Acceptable identification**						**23411022**
Lactobacillus fermentum		92.72	0.46	NIT 1			
Actinomyces naeslundii		4.01	0.33				
Actinomyces odontolyticus		1.95	0.10				
23411023	**Good identification**						**23411023**
Actinomyces naeslundii		100.0	0.36	MNS 88			
23411100	**Low discrimination**						**23411100**
					GRA		
Lactobacillus fermentum		56.53	0.58	NIT 1	99		
Fusobacterium mortiferum		36.86	0.53		1		
Actinomyces meyeri		3.35	0.26		99		
23411101	**Low discrimination**						**23411101**
					SPO		
Actinomyces naeslundii		56.07	0.43	MNS 88	1		
Clostridium chauvoei		19.14	0.12	MLT 94	99		
Lactobacillus fermentum		15.01	0.24		1		
23411102	**Low discrimination**						**23411102**
					GRA		
Lactobacillus fermentum		60.97	0.58	NIT 1	99		
Actinomyces viscosus		31.13	0.39		99		
Fusobacterium mortiferum		4.42	0.37		1		
23411120	**Low discrimination**						**23411120**
					GRA		
Lactobacillus fermentum		78.36	0.50	NIT 1	99		
Prevotella melaninogenica		10.39	0.15		1		
Actinomyces naeslundii		5.91	0.40		99		
23411121	**Low discrimination**						**23411121**
					SPO		
Actinomyces naeslundii		83.12	0.43	MNS 88	1		
Clostridium chauvoei		9.46	0.04	MLT 94	99		
Lactobacillus fermentum		7.42	0.15		1		
23411122	**Low discrimination**						**23411122**
					GRA	CAT	
Lactobacillus fermentum		83.12	0.50	NIT 1	99	1	
Capnocytophaga ochracea		6.62	0.15		1	90	
Actinomyces naeslundii		6.27	0.40		99	25	
23411123	**Acceptable identification**						**23411123**
Actinomyces naeslundii		91.81	0.43	MNS 88			
Lactobacillus fermentum		8.19	0.15	MLT 94			
23411400	**Good identification**						**23411400**
					GRA		
Lactobacillus fermentum		98.95	0.54	NIT 1	99		
Actinomyces meyeri		0.60	0.05		99		
Fusobacterium mortiferum		0.46	0.13		1		
23411402	**Good identification**						**23411402**
Lactobacillus fermentum		99.53	0.54	NIT 1			
Actinomyces viscosus		0.47	0.00				
23411420	**Good identification**						**23411420**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		100.0	0.46	NIT 1			
23411422	**Good identification**						**23411422**
Lactobacillus fermentum		100.0	0.46	NIT 1			
23411500	**Very good identification**						**23411500**
					GRA		
Lactobacillus fermentum		97.90	0.58	NIT 1	99		
Actinomyces meyeri		1.45	0.16		99		
Fusobacterium mortiferum		0.65	0.19		1		
23411502	**Very good identification**						**23411502**
Lactobacillus fermentum		99.49	0.58	NIT 1			
Actinomyces viscosus		0.51	0.04				
23411520	**Good identification**						**23411520**
Lactobacillus fermentum		99.32	0.50	NIT 1			
Bifidobacterium longum subsp. longum		0.68	0.05				
23411522	**Good identification**						**23411522**
					GRA	CAT	
Lactobacillus fermentum		97.95	0.50	NIT 1	99	1	
Capnocytophaga ochracea		1.38	0.02		1	90	
Bifidobacterium longum subsp. longum		0.67	0.05		99	1	
23412100	**Very good identification**						**23412100**
					GRA		
Propionibacterium propionicum		99.68	0.60	MLT 99	99		
Fusobacterium mortiferum		0.32	0.16		1		
23412101	**Good identification**						**23412101**
Propionibacterium propionicum		98.03	0.44	MLT 99			
Actinomyces naeslundii		1.97	0.25	URE 10			
23412102	**Very good identification**						**23412102**
Propionibacterium propionicum		99.59	0.63	MLT 99			
Actinomyces viscosus		0.41	0.08				
23412103	**Good identification**						**23412103**
Propionibacterium propionicum		97.49	0.47	MLT 99			
Actinomyces naeslundii		1.31	0.25	URE 10			
Actinomyces viscosus		1.20	0.00				
23412140	**Very good identification**						**23412140**
Propionibacterium propionicum		100.0	0.60	MLT 99			
23412141	**Good identification**						**23412141**
Propionibacterium propionicum		100.0	0.44	MLT 99			
23412142	**Very good identification**						**23412142**
Propionibacterium propionicum		100.0	0.63	MLT 99			
23412143	**Good identification**						**23412143**
Propionibacterium propionicum		100.0	0.47	MLT 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23413000	**Low discrimination**					**23413000**
					CAT	
Actinomyces naeslundii		31.48	0.38	MNS 88	25	
Propionibacterium granulosum		26.67	0.21	MLT 94	90	
Propionibacterium propionicum		25.31	0.26		1	
23413001	**Good identification**					**23413001**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23413003	**Acceptable identification**					**23413003**
Actinomyces naeslundii		91.80	0.41	MNS 88		
Propionibacterium propionicum		8.20	0.12	MLT 94		
23413020	**Low discrimination**					**23413020**
Actinomyces naeslundii		63.32	0.38	MNS 88		
Propionibacterium acnes		23.27	0.06	MLT 94		
Propionibacterium granulosum		13.41	0.11			
23413021	**Good identification**					**23413021**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23413022	**Good identification**					**23413022**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23413023	**Good identification**					**23413023**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23413100	**Very good identification**					**23413100**
Propionibacterium propionicum		95.42	0.60	MLT 99		
Actinomyces naeslundii		3.41	0.45			
Lactobacillus fermentum		0.71	0.24			
23413101	**Low discrimination**					**23413101**
Propionibacterium propionicum		67.44	0.44	MLT 99		
Actinomyces naeslundii		32.56	0.48	URE 10		
23413102	**Very good identification**					**23413102**
Propionibacterium propionicum		96.97	0.63	MLT 99		
Actinomyces naeslundii		2.31	0.45			
Lactobacillus fermentum		0.48	0.24			
23413103	**Low discrimination**					**23413103**
Propionibacterium propionicum		75.65	0.47	MLT 99		
Actinomyces naeslundii		24.35	0.48	URE 10		
23413120	**Low discrimination**					**23413120**
Actinomyces naeslundii		74.00	0.45	MNS 88		
Propionibacterium propionicum		20.90	0.26	MLT 94		
Lactobacillus fermentum		5.10	0.15			
23413121	**Good identification**					**23413121**
Actinomyces naeslundii		100.0	0.48	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23413122	**Low discrimination**					**23413122**
Actinomyces naeslundii		67.00	0.45	MNS 88		
Propionibacterium propionicum		28.38	0.29	MLT 94		
Lactobacillus fermentum		4.62	0.15			
23413123	**Good identification**					**23413123**
Actinomyces naeslundii		96.96	0.48	MNS 88		
Propionibacterium propionicum		3.04	0.12	MLT 94		
23413140	**Very good identification**					**23413140**
Propionibacterium propionicum		99.37	0.60	MLT 99		
Actinomyces naeslundii		0.63	0.32			
23413141	**Acceptable identification**					**23413141**
Propionibacterium propionicum		92.15	0.44	MLT 99		
Actinomyces naeslundii		7.85	0.35	URE 10		
23413142	**Very good identification**					**23413142**
Propionibacterium propionicum		99.58	0.63	MLT 99		
Actinomyces naeslundii		0.42	0.32			
23413143	**Acceptable identification**					**23413143**
Propionibacterium propionicum		94.63	0.47	MLT 99		
Actinomyces naeslundii		5.38	0.35	URE 10		
23413161	**Good identification**					**23413161**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23413163	**Low discrimination**					**23413163**
Actinomyces naeslundii		84.91	0.35	MNS 88		
Propionibacterium propionicum		15.10	0.12	MLT 94		
23415000	**Low discrimination**					**23415000**
Lactobacillus fermentum		83.53	0.50	NIT 1		
Actinomyces meyeri		6.68	0.21			
Actinomyces odontolyticus		4.92	0.22			
23415002	**Good identification**					**23415002**
Lactobacillus fermentum		95.18	0.50	NIT 1		
Actinomyces odontolyticus		3.43	0.18			
Actinomyces viscosus		0.77	0.00			
23415020	**Acceptable identification**					**23415020**
Lactobacillus fermentum		94.21	0.42	NIT 1		
Actinomyces meyeri		3.08	0.06			
Actinomyces odontolyticus		2.71	0.08			
23415022	**Good identification**					**23415022**
Lactobacillus fermentum		100.0	0.42	NIT 1		
23415100	**Low discrimination**					**23415100**

GRA

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		77.47	0.54	NIT 1	99		
Actinomyces meyeri		15.18	0.31		99		
Fusobacterium mortiferum		6.47	0.34		1		
23415102	**Good identification**						**23415102**
					GRA		
Lactobacillus fermentum		96.98	0.54	NIT 1	99		
Fusobacterium mortiferum		0.90	0.17		1		
Actinomyces viscosus		0.85	0.04		99		
23415120	**Acceptable identification**						**23415120**
Lactobacillus fermentum		92.58	0.46	NIT 1			
Actinomyces meyeri		7.42	0.16				
23415122	**Good identification**						**23415122**
					GRA	CAT	
Lactobacillus fermentum		96.72	0.46	NIT 1	99	1	
Capnocytophaga ochracea		3.28	0.05		1	90	
23415400	**Good identification**						**23415400**
Lactobacillus fermentum		98.04	0.50	NIT 1			
Actinomyces meyeri		1.96	0.10				
23415402	**Good identification**						**23415402**
Lactobacillus fermentum		100.0	0.50	NIT 1			
23415420	**Good identification**						**23415420**
Lactobacillus fermentum		100.0	0.42	NIT 1			
23415422	**Good identification**						**23415422**
Lactobacillus fermentum		100.0	0.42	NIT 1			
23415500	**Good identification**						**23415500**
Lactobacillus fermentum		95.33	0.54	NIT 1			
Actinomyces meyeri		4.67	0.21				
23415502	**Good identification**						**23415502**
Lactobacillus fermentum		100.0	0.54	NIT 1			
23415520	**Acceptable identification**						**23415520**
Lactobacillus fermentum		94.77	0.46	NIT 1			
Bifidobacterium longum subsp. longum		3.33	0.14				
Actinomyces meyeri		1.90	0.06				
23415522	**Good identification**						**23415522**
Lactobacillus fermentum		96.61	0.46	NIT 1			
Bifidobacterium longum subsp. longum		3.39	0.14				
23430100	**Acceptable identification**						**23430100**
					GRA		
Fusobacterium mortiferum		92.01	0.41	NIT 1	1		
Propionibacterium propionicum		8.00	0.24		99		
23431000	**Low discrimination**						**23431000**
					GRA	CAT	
Fusobacterium mortiferum		50.71	0.38	NIT 1	1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		35.11	0.32		99	90	
Actinomyces naeslundii		10.42	0.38		99	25	
23431001	**Good identification**						**23431001**
Actinomyces naeslundii		100.0	0.41	MNS 88			
23431003	**Low discrimination**						**23431003**
Actinomyces naeslundii		78.13	0.41	MNS 88			
Actinomyces viscosus		21.87	0.07	MLT 94			
23431020	**Low discrimination**						**23431020**
Actinomyces naeslundii		54.28	0.38	MNS 88			
Propionibacterium granulosum		45.72	0.21	MLT 94			
23431021	**Good identification**						**23431021**
Actinomyces naeslundii		100.0	0.41	MNS 88			
23431022	**Low discrimination**						**23431022**
Actinomyces naeslundii		82.61	0.38	MNS 88			
Propionibacterium granulosum		17.39	0.11	MLT 94			
23431023	**Good identification**						**23431023**
Actinomyces naeslundii		100.0	0.41	MNS 88			
23431100	**Low discrimination**						**23431100**
					GRA		
Fusobacterium mortiferum		73.50	0.44	NIT 1	1		
Actinomyces naeslundii		18.43	0.45		99		
Propionibacterium propionicum		4.26	0.24		99		
23431101	**Good identification**						**23431101**
Actinomyces naeslundii		100.0	0.48	MNS 88			
23431102	**Low discrimination**						**23431102**
					GRA		
Actinomyces naeslundii		35.93	0.45	MNS 88	99		
Actinomyces viscosus		28.27	0.19	MLT 94	99		
Fusobacterium mortiferum		15.92	0.28		1		
23431103	**Low discrimination**						**23431103**
Actinomyces naeslundii		85.12	0.48	MNS 88			
Actinomyces viscosus		14.88	0.11	MLT 94			
23431120	**Low discrimination**						**23431120**
Actinomyces naeslundii		79.19	0.45	MNS 88			
Clostridium perfringens		15.35	0.11	MLT 94			
Lactobacillus fermentum		5.46	0.15				
23431121	**Good identification**						**23431121**
Actinomyces naeslundii		100.0	0.48	MNS 88			
23431122	**Low discrimination**						**23431122**
Actinomyces naeslundii		86.19	0.45	MNS 88			
Clostridium perfringens		7.86	0.05	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		5.94	0.15			
23431123	**Good identification**					**23431123**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23431141	**Good identification**					**23431141**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23431143	**Good identification**					**23431143**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23431161	**Good identification**					**23431161**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23431163	**Good identification**					**23431163**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23432100	**Very good identification**					**23432100**
Propionibacterium propionicum		99.65	0.58	MLT 99		
Actinomyces naeslundii		0.35	0.27			
23432101	**Good identification**					**23432101**
Propionibacterium propionicum		95.45	0.42	MLT 99		
Actinomyces naeslundii		4.56	0.30	URE 10		
23432102	**Very good identification**					**23432102**
Propionibacterium propionicum		99.77	0.61	MLT 99		
Actinomyces naeslundii		0.24	0.27			
23432103	**Good identification**					**23432103**
Propionibacterium propionicum		96.92	0.45	MLT 99		
Actinomyces naeslundii		3.08	0.30	URE 10		
23432140	**Very good identification**					**23432140**
Propionibacterium propionicum		100.0	0.58	MLT 99		
23432141	**Good identification**					**23432141**
Propionibacterium propionicum		100.0	0.42	MLT 99		
23432142	**Very good identification**					**23432142**
Propionibacterium propionicum		100.0	0.61	MLT 99		
23432143	**Good identification**					**23432143**
Propionibacterium propionicum		100.0	0.45	MLT 99		
23433001	**Good identification**					**23433001**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23433002	**Low discrimination**					**23433002**
					CAT	
Actinomyces naeslundii		54.48	0.42	MNS 88	25	
Propionibacterium propionicum		27.69	0.27	MLT 94	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		17.83	0.19		90	
23433003	**Good identification**					**23433003**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23433020	**Low discrimination**					**23433020**
Actinomyces naeslundii		75.34	0.42	MNS 88		
Propionibacterium granulosum		24.66	0.19	MLT 94		
23433021	**Good identification**					**23433021**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23433022	**Acceptable identification**					**23433022**
Actinomyces naeslundii		92.44	0.42	MNS 88		
Propionibacterium granulosum		7.56	0.09	MLT 94		
23433023	**Good identification**					**23433023**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23433100	**Good identification**					**23433100**
Propionibacterium propionicum		92.18	0.58	MLT 99		
Actinomyces naeslundii		7.82	0.50			
23433101	**Low discrimination**					**23433101**
Actinomyces naeslundii		53.39	0.53	MNS 88		
Propionibacterium propionicum		46.61	0.42	MLT 94		
23433102	**Good identification**					**23433102**
Propionibacterium propionicum		94.65	0.61	MLT 99		
Actinomyces naeslundii		5.35	0.50			
23433103	**Low discrimination**					**23433103**
Propionibacterium propionicum		56.70	0.45	MLT 99		
Actinomyces naeslundii		43.30	0.53	URE 10		
23433120	**Low discrimination**					**23433120**
Actinomyces naeslundii		89.36	0.50	MNS 88		
Propionibacterium propionicum		10.64	0.24	MLT 94		
23433121	**Good identification**					**23433121**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23433122	**Low discrimination**					**23433122**
Actinomyces naeslundii		84.85	0.50	MNS 88		
Propionibacterium propionicum		15.15	0.27	MLT 94		
23433123	**Good identification**					**23433123**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23433140	**Very good identification**					**23433140**
Propionibacterium propionicum		98.53	0.58	MLT 99		
Actinomyces naeslundii		1.48	0.37			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23433141	**Low discrimination**					**23433141**
Propionibacterium propionicum		83.18	0.42	MLT 99		
Actinomyces naeslundii		16.82	0.40	URE 10		
23433142	**Very good identification**					**23433142**
Propionibacterium propionicum		99.01	0.61	MLT 99		
Actinomyces naeslundii		0.99	0.37			
23433143	**Low discrimination**					**23433143**
Propionibacterium propionicum		88.12	0.45	MLT 99		
Actinomyces naeslundii		11.88	0.40	URE 10		
23433160	**Low discrimination**					**23433160**
Actinomyces naeslundii		59.72	0.37	MNS 88		
Propionibacterium propionicum		40.28	0.24	MLT 94		
23433161	**Good identification**					**23433161**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23433163	**Good identification**					**23433163**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23450000	**Good identification**					**23450000**
Fusobacterium mortiferum		100.0	0.36	NIT 1		
23450100	**Good identification**					**23450100**
					GRA	
Fusobacterium mortiferum		98.48	0.42	NIT 1	1	
Actinomyces viscosus		1.52	-.02		99	
23451000	**Low discrimination**					**23451000**
					GRA	
Fusobacterium mortiferum		83.88	0.39	NIT 1	1	
Actinomyces naeslundii		10.55	0.35		99	
Lactobacillus fermentum		5.57	0.20		99	
23451001	**Good identification**					**23451001**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23451021	**Good identification**					**23451021**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23451023	**Good identification**					**23451023**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23451100	**Low discrimination**					**23451100**
					GRA	
Fusobacterium mortiferum		83.34	0.45	NIT 1	1	
Actinomyces naeslundii		12.79	0.42		99	
Lactobacillus fermentum		3.87	0.24		99	
23451101	**Good identification**					**23451101**
Actinomyces naeslundii		100.0	0.46	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23451103	**Low discrimination**						**23451103**
Actinomyces naeslundii		52.51	0.46	MNS 88			
Actinomyces viscosus		47.49	0.21	MLT 94			
23451120	**Low discrimination**						**23451120**
					GRA		
Actinomyces naeslundii		80.11	0.42	MNS 88	99		
Prevotella melaninogenica		11.80	-.01	MLT 94	1		
Lactobacillus fermentum		8.09	0.15		99		
23451121	**Good identification**						**23451121**
Actinomyces naeslundii		100.0	0.46	MNS 88			
23451122	**Low discrimination**						**23451122**
					GRA	CAT	
Actinomyces naeslundii		76.92	0.42	MNS 88	99	25	
Capnocytophaga ochracea		15.31	0.05	MLT 94	1	90	
Lactobacillus fermentum		7.77	0.15		99	1	
23451123	**Good identification**						**23451123**
Actinomyces naeslundii		100.0	0.46	MNS 88			
23453000	**Good identification**						**23453000**
Actinomyces naeslundii		100.0	0.40	MNS 88			
23453001	**Good identification**						**23453001**
Actinomyces naeslundii		100.0	0.43	MNS 88			
23453002	**Good identification**						**23453002**
Actinomyces naeslundii		100.0	0.40	MNS 88			
23453003	**Good identification**						**23453003**
Actinomyces naeslundii		100.0	0.43	MNS 88			
23453020	**Good identification**						**23453020**
Actinomyces naeslundii		100.0	0.40	MNS 88			
23453021	**Good identification**						**23453021**
Actinomyces naeslundii		100.0	0.43	MNS 88			
23453022	**Good identification**						**23453022**
Actinomyces naeslundii		100.0	0.40	MNS 88			
23453023	**Good identification**						**23453023**
Actinomyces naeslundii		100.0	0.43	MNS 88			
23453100	**Low discrimination**						**23453100**
Actinomyces naeslundii		82.44	0.47	MNS 88			
Propionibacterium propionicum		17.56	0.26	MLT 94			
23453101	**Good identification**						**23453101**
Actinomyces naeslundii		100.0	0.50	MNS 88			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23453102	**Low discrimination**					**23453102**
Actinomyces naeslundii		75.78	0.47	MNS 88		
Propionibacterium propionicum		24.22	0.29	MLT 94		
23453103	**Good identification**					**23453103**
Actinomyces naeslundii		97.69	0.50	MNS 88		
Propionibacterium propionicum		2.31	0.12	MLT 94		
23453120	**Good identification**					**23453120**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23453121	**Good identification**					**23453121**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23453122	**Good identification**					**23453122**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23453123	**Good identification**					**23453123**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23453141	**Good identification**					**23453141**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23453143	**Low discrimination**					**23453143**
Actinomyces naeslundii		88.17	0.38	MNS 88		
Propionibacterium propionicum		11.83	0.12	MLT 94		
23453161	**Good identification**					**23453161**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23453163	**Good identification**					**23453163**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23471001	**Good identification**					**23471001**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23471002	**Low discrimination**					**23471002**
					GRA	
Actinomyces naeslundii		72.02	0.40	MNS 88	99	
Actinomyces viscosus		18.19	0.05	MLT 94	99	
Fusobacterium mortiferum		9.79	0.13		1	
23471003	**Good identification**					**23471003**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23471020	**Good identification**					**23471020**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23471021	**Good identification**					**23471021**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23471022	**Good identification**					**23471022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MNS 88		
23471023	**Good identification**					**23471023**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23471100	**Low discrimination**					**23471100**
					GRA	
Fusobacterium mortiferum		50.07	0.36	NIT 1	1	
Actinomyces naeslundii		49.93	0.47		99	
23471101	**Good identification**					**23471101**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23471102	**Low discrimination**					**23471102**
					GRA	
Actinomyces naeslundii		78.79	0.47	MNS 88	99	
Actinomyces viscosus		12.43	0.10	MLT 94	99	
Fusobacterium mortiferum		8.78	0.20		1	
23471103	**Good identification**					**23471103**
Actinomyces naeslundii		96.61	0.50	MNS 88		
Actinomyces viscosus		3.39	0.01	MLT 94		
23471120	**Good identification**					**23471120**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23471121	**Good identification**					**23471121**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23471122	**Good identification**					**23471122**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23471123	**Good identification**					**23471123**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23471141	**Good identification**					**23471141**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23471143	**Good identification**					**23471143**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23471161	**Good identification**					**23471161**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23471163	**Good identification**					**23471163**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23473000	**Good identification**					**23473000**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23473001	**Good identification**					**23473001**
Actinomyces naeslundii		100.0	0.48	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23473002	**Good identification**					**23473002**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23473003	**Good identification**					**23473003**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23473020	**Good identification**					**23473020**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23473021	**Good identification**					**23473021**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23473022	**Good identification**					**23473022**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23473023	**Good identification**					**23473023**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23473100	**Acceptable identification**					**23473100**
Actinomyces naeslundii		91.76	0.52	MNS 88		
Propionibacterium propionicum		8.24	0.24	MLT 94		
23473101	**Good identification**					**23473101**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23473102	**Low discrimination**					**23473102**
Actinomyces naeslundii		88.13	0.52	MNS 88		
Propionibacterium propionicum		11.87	0.27	MLT 94		
23473103	**Good identification**					**23473103**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23473120	**Good identification**					**23473120**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23473121	**Good identification**					**23473121**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23473122	**Good identification**					**23473122**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23473123	**Good identification**					**23473123**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23473140	**Low discrimination**					**23473140**
Actinomyces naeslundii		66.28	0.39	MNS 88		
Propionibacterium propionicum		33.72	0.24	MLT 94		
23473141	**Good identification**					**23473141**
Actinomyces naeslundii		100.0	0.42	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23473142	**Low discrimination**					**23473142**
Actinomyces naeslundii		56.71	0.39	MNS 88		
Propionibacterium propionicum		43.29	0.27	MLT 94		
23473143	**Good identification**					**23473143**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23473160	**Good identification**					**23473160**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23473161	**Good identification**					**23473161**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23473162	**Good identification**					**23473162**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23473163	**Good identification**					**23473163**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23500000	**Acceptable identification**					**23500000**
					SPO	
Actinomyces odontolyticus		92.62	0.46	RAF 3	1	
Actinomyces viscosus		4.11	0.16	FRU 92	1	
Clostridium botulinum biovar A		1.20	0.06		99	
23500002	**Low discrimination**					**23500002**
Actinomyces viscosus		87.77	0.50	MNS 96		
Actinomyces odontolyticus		12.23	0.43	FRU 95		
23500003	**Good identification**					**23500003**
Actinomyces viscosus		99.58	0.42	MNS 96		
Actinomyces odontolyticus		0.42	0.08	FRU 95		
23500100	**Low discrimination**					**23500100**
Actinomyces odontolyticus		73.31	0.38	RAF 3		
Actinomyces viscosus		18.33	0.20	FRU 92		
Propionibacterium propionicum		5.77	0.26			
23500102	**Good identification**					**23500102**
Actinomyces viscosus		97.13	0.55	MNS 96		
Actinomyces odontolyticus		2.41	0.34	FRU 95		
Propionibacterium propionicum		0.46	0.29			
23500103	**Good identification**					**23500103**
Actinomyces viscosus		99.84	0.46	MNS 96		
Propionibacterium propionicum		0.16	0.12	FRU 95		
23500200	**Good identification**					**23500200**
Actinomyces odontolyticus		100.0	0.51	RAF 3		
23500202	**Good identification**					**23500202**
Actinomyces odontolyticus		96.40	0.47	RAF 3		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		3.60	0.16	FRU 92		
23500220	**Good identification**					**23500220**
Actinomyces odontolyticus		100.0	0.38	RAF 3		
23500300	**Good identification**					**23500300**
Actinomyces odontolyticus		100.0	0.43	RAF 3		
23500302	**Low discrimination**					**23500302**
Actinomyces odontolyticus		82.63	0.39	RAF 3		
Actinomyces viscosus		17.37	0.20	FRU 92		
23501000	**Low discrimination**					**23501000**
Actinomyces odontolyticus		88.52	0.43	RAF 3		
Lactobacillus fermentum		4.44	0.28	FRU 92		
Actinomyces viscosus		3.61	0.12			
23501002	**Low discrimination**					**23501002**
Actinomyces viscosus		85.55	0.47	MNS 96		
Actinomyces odontolyticus		12.97	0.40	FRU 95		
Lactobacillus fermentum		1.06	0.28			
23501003	**Good identification**					**23501003**
Actinomyces viscosus		97.42	0.38	MNS 96		
Actinomyces naeslundii		2.14	0.34	FRU 95		
Actinomyces odontolyticus		0.45	0.05			
23501101	**Acceptable identification**					**23501101**
					SPO	
Clostridium chauvoei		92.56	0.40	MNS 90	99	
Actinomyces naeslundii		5.27	0.42	RAF 1	1	
Actinomyces viscosus		1.52	0.09		1	
23501102	**Acceptable identification**					**23501102**
					COC	
Actinomyces viscosus		93.37	0.51	MNS 96	1	
Actinomyces odontolyticus		2.52	0.31	FRU 95	1	
Blautia hansenii		1.50	0.15		99	
23501103	**Low discrimination**					**23501103**
					SPO	
Actinomyces viscosus		87.22	0.43	MNS 96	1	
Clostridium chauvoei		9.49	0.27	FRU 95	99	
Actinomyces naeslundii		3.06	0.42		1	
23501200	**Good identification**					**23501200**
Actinomyces odontolyticus		100.0	0.48	RAF 3		
23501202	**Good identification**					**23501202**
Actinomyces odontolyticus		96.68	0.44	RAF 3		
Actinomyces viscosus		3.32	0.12	FRU 92		
23501300	**Good identification**					**23501300**
					COC	
Actinomyces odontolyticus		98.32	0.40	RAF 3	1	
Blautia hansenii		1.68	-.03	FRU 92	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23501302	**Low discrimination**					**23501302**
Actinomyces odontolyticus		83.81	0.36	RAF 3		
Actinomyces viscosus		16.19	0.17	FRU 92		
23502100	**Very good identification**					**23502100**
Propionibacterium propionicum		100.0	0.60	FRU 99		
23502101	**Good identification**					**23502101**
Propionibacterium propionicum		98.37	0.44	FRU 99		
Actinomyces naeslundii		1.63	0.23	URE 10		
23502102	**Very good identification**					**23502102**
Propionibacterium propionicum		97.90	0.63	FRU 99		
Actinomyces viscosus		2.10	0.20			
23502103	**Acceptable identification**					**23502103**
Propionibacterium propionicum		93.00	0.47	FRU 99		
Actinomyces viscosus		5.97	0.12	URE 10		
Actinomyces naeslundii		1.03	0.23			
23502140	**Very good identification**					**23502140**
Propionibacterium propionicum		100.0	0.60	FRU 99		
23502141	**Good identification**					**23502141**
Propionibacterium propionicum		100.0	0.44	FRU 99		
23502142	**Very good identification**					**23502142**
Propionibacterium propionicum		100.0	0.63	FRU 99		
23502143	**Good identification**					**23502143**
Propionibacterium propionicum		100.0	0.47	FRU 99		
23503000	**Low discrimination**					**23503000**
Actinomyces naeslundii		44.66	0.36	MNS 88		
Propionibacterium propionicum		43.54	0.26	FRU 95		
Actinomyces odontolyticus		11.80	0.09			
23503001	**Good identification**					**23503001**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23503003	**Low discrimination**					**23503003**
Actinomyces naeslundii		74.31	0.39	MNS 88		
Actinomyces viscosus		17.64	0.04	FRU 95		
Propionibacterium propionicum		8.05	0.12			
23503020	**Good identification**					**23503020**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23503021	**Good identification**					**23503021**
Actinomyces naeslundii		100.0	0.39	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23503022	**Good identification**					**23503022**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23503023	**Good identification**					**23503023**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23503100	**Very good identification**					**23503100**
					COC	
Propionibacterium propionicum		96.87	0.60	FRU 99	1	
Actinomyces naeslundii		2.86	0.44		1	
Blautia hansenii		0.28	-.03		99	
23503101	**Low discrimination**					**23503101**
					SPO	
Propionibacterium propionicum		69.71	0.44	FRU 99	1	
Actinomyces naeslundii		27.75	0.47	URE 10	1	
Clostridium chauvoei		2.54	0.06		99	
23503102	**Very good identification**					**23503102**
Propionibacterium propionicum		96.83	0.63	FRU 99		
Actinomyces naeslundii		1.90	0.44			
Actinomyces viscosus		1.27	0.17			
23503103	**Low discrimination**					**23503103**
Propionibacterium propionicum		76.64	0.47	FRU 99		
Actinomyces naeslundii		20.34	0.47	URE 10		
Actinomyces viscosus		3.02	0.09			
23503120	**Low discrimination**					**23503120**
					GRA	
Actinomyces naeslundii		68.29	0.44	MNS 88	99	
Propionibacterium propionicum		23.39	0.26	FRU 95	99	
Prevotella melaninogenica		8.33	-.01		1	
23503121	**Good identification**					**23503121**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23503122	**Low discrimination**					**23503122**
Actinomyces naeslundii		66.06	0.44	MNS 88		
Propionibacterium propionicum		33.94	0.29	FRU 95		
23503123	**Good identification**					**23503123**
Actinomyces naeslundii		96.33	0.47	MNS 88		
Propionibacterium propionicum		3.67	0.12	FRU 95		
23503140	**Very good identification**					**23503140**
Propionibacterium propionicum		99.48	0.60	FRU 99		
Actinomyces naeslundii		0.52	0.31			
23503141	**Acceptable identification**					**23503141**
Propionibacterium propionicum		93.44	0.44	FRU 99		
Actinomyces naeslundii		6.57	0.34	URE 10		
23503142	**Very good identification**					**23503142**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.65	0.63	FRU 99		
Actinomyces naeslundii		0.35	0.31			
23503143	**Good identification**					**23503143**
Propionibacterium propionicum		95.53	0.47	FRU 99		
Actinomyces naeslundii		4.47	0.34	URE 10		
23504000	**Good identification**					**23504000**
Actinomyces odontolyticus		100.0	0.41	RAF 3		
23504002	**Low discrimination**					**23504002**
Actinomyces odontolyticus		87.18	0.37	RAF 3		
Actinomyces viscosus		12.83	0.16	FRU 92		
23504200	**Good identification**					**23504200**
Actinomyces odontolyticus		100.0	0.46	RAF 3		
23504202	**Good identification**					**23504202**
Actinomyces odontolyticus		100.0	0.42	RAF 3		
23504300	**Good identification**					**23504300**
Actinomyces odontolyticus		100.0	0.37	RAF 3		
23505000	**Acceptable identification**					**23505000**
Actinomyces odontolyticus		94.36	0.38	RAF 3		
Lactobacillus fermentum		5.65	0.24	FRU 92		
23505200	**Good identification**					**23505200**
Actinomyces odontolyticus		100.0	0.43	RAF 3		
23505202	**Good identification**					**23505202**
Actinomyces odontolyticus		100.0	0.39	RAF 3		
23505530	**Good identification**					**23505530**
Bifidobacterium longum subsp. longum		100.0	0.35	NIT 1		
23505532	**Good identification**					**23505532**
Bifidobacterium longum subsp. longum		100.0	0.35	NIT 1		
23510000	**Low discrimination**					**23510000**
					GRA	
Actinomyces odontolyticus		86.61	0.64	RAF 3	99	
Actinomyces viscosus		6.36	0.38		99	
Prevotella intermedia		4.28	0.36		1	
23510002	**Good identification**					**23510002**
Actinomyces viscosus		92.10	0.72	MNS 96		
Actinomyces odontolyticus		7.77	0.61			
Propionibacterium propionicum		0.04	0.29			
23510003	**Very good identification**					**23510003**
Actinomyces viscosus		99.68	0.64	MNS 96		
Actinomyces odontolyticus		0.26	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.06	0.32			
23510012	**Acceptable identification**					**23510012**
Actinomyces viscosus		92.22	0.38	MNS 96		
Actinomyces odontolyticus		7.78	0.27	MLZ 1		
23510020	**Good identification**					**23510020**
Actinomyces odontolyticus		98.12	0.51	RAF 3		
Lactobacillus fermentum		0.58	0.20	NAG 14		
Actinomyces naeslundii		0.55	0.29			
23510022	**Low discrimination**					**23510022**
Actinomyces odontolyticus		57.01	0.47	RAF 3		
Actinomyces viscosus		41.92	0.38	NAG 14		
Lactobacillus fermentum		0.55	0.20			
23510042	**Low discrimination**					**23510042**
Actinomyces viscosus		87.12	0.38	MNS 96		
Actinomyces odontolyticus		7.35	0.27	SOR 1		
Propionibacterium propionicum		3.85	0.29			
23510100	**Low discrimination**					**23510100**
Actinomyces odontolyticus		46.22	0.56	RAF 3		
Propionibacterium propionicum		31.30	0.60			
Actinomyces viscosus		19.09	0.42			
23510102	**Very good identification**					**23510102**
Actinomyces viscosus		96.11	0.76	MNS 96		
Propionibacterium propionicum		2.39	0.63			
Actinomyces odontolyticus		1.44	0.52			
23510103	**Very good identification**					**23510103**
Actinomyces viscosus		99.04	0.68	MNS 96		
Propionibacterium propionicum		0.82	0.47			
Actinomyces naeslundii		0.09	0.40			
23510112	**Good identification**					**23510112**
Actinomyces viscosus		96.17	0.42	MNS 96		
Propionibacterium propionicum		2.39	0.29	MLZ 1		
Actinomyces odontolyticus		1.44	0.18			
23510120	**Low discrimination**					**23510120**
					GRA	
Actinomyces odontolyticus		75.41	0.42	RAF 3	99	
Prevotella melaninogenica		10.15	0.15	NAG 14	1	
Actinomyces naeslundii		3.77	0.37		99	
23510121	**Low discrimination**					**23510121**
					SPO	
Actinomyces naeslundii		70.29	0.40	MNS 88	1	
Clostridium chauvoei		12.25	0.04	GAL 96	99	
Actinomyces odontolyticus		9.46	0.08		1	
23510122	**Low discrimination**					**23510122**
					GRA CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		74.46	0.42	MNS 96	99	76	
Actinomyces odontolyticus		17.98	0.39	NAG 1	99	10	
Capnocytophaga ochracea		2.37	0.15		1	90	
23510140	**Very good identification**						**23510140**
Propionibacterium propionicum		97.73	0.60	MAN 99			
Actinomyces odontolyticus		1.46	0.22				
Actinomyces viscosus		0.60	0.08				
23510141	**Good identification**						**23510141**
Propionibacterium propionicum		95.50	0.44	MAN 99			
Actinomyces naeslundii		2.74	0.27	URE 10			
Actinomyces viscosus		1.77	0.00				
23510142	**Low discrimination**						**23510142**
Propionibacterium propionicum		70.71	0.63	MAN 99			
Actinomyces viscosus		28.76	0.42				
Actinomyces odontolyticus		0.43	0.18				
23510200	**Very good identification**						**23510200**
					GRA		
Actinomyces odontolyticus		99.94	0.69	RAF 3	99		
Actinomyces viscosus		0.04	0.04		99		
Prevotella intermedia		0.03	0.02		1		
23510201	**Good identification**						**23510201**
Actinomyces odontolyticus		100.0	0.35	RAF 3			
23510202	**Good identification**						**23510202**
Actinomyces odontolyticus		94.19	0.66	RAF 3			
Actinomyces viscosus		5.81	0.38				
23510210	**Good identification**						**23510210**
Actinomyces odontolyticus		100.0	0.35	RAF 3			
23510220	**Good identification**						**23510220**
Actinomyces odontolyticus		100.0	0.56	RAF 3			
23510222	**Good identification**						**23510222**
Actinomyces odontolyticus		99.62	0.52	RAF 3			
Actinomyces viscosus		0.38	0.04	NAG 14			
23510240	**Good identification**						**23510240**
Actinomyces odontolyticus		100.0	0.35	RAF 3			
23510300	**Very good identification**						**23510300**
Actinomyces odontolyticus		99.44	0.61	RAF 3			
Propionibacterium propionicum		0.35	0.26				
Actinomyces viscosus		0.21	0.08				
23510302	**Low discrimination**						**23510302**
Actinomyces odontolyticus		73.76	0.57	RAF 3			
Actinomyces viscosus		25.61	0.42				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		0.64	0.29			
23510320	**Good identification**					**23510320**
Actinomyces odontolyticus		100.0	0.47	RAF 3		
23510322	**Good identification**					**23510322**
Actinomyces odontolyticus		97.89	0.44	RAF 3		
Actinomyces viscosus		2.11	0.08	NAG 14		
23510402	**Low discrimination**					**23510402**
Actinomyces viscosus		88.97	0.38	MNS 96		
Actinomyces odontolyticus		7.51	0.27	ARA 1		
Lactobacillus fermentum		3.53	0.28			
23510502	**Acceptable identification**					**23510502**
Actinomyces viscosus		92.92	0.42	MNS 96		
Lactobacillus fermentum		3.38	0.32	ARA 1		
Propionibacterium propionicum		2.31	0.29			
23510600	**Good identification**					**23510600**
Actinomyces odontolyticus		100.0	0.35	RAF 3		
23511000	**Low discrimination**					**23511000**
					GRA	
Actinomyces odontolyticus		61.60	0.61	RAF 3	99	
Lactobacillus fermentum		26.61	0.62		99	
Prevotella intermedia		4.56	0.36		1	
23511001	**Low discrimination**					**23511001**
Actinomyces naeslundii		54.51	0.56	MNS 88		
Actinomyces viscosus		25.12	0.26			
Actinomyces odontolyticus		11.28	0.27			
23511002	**Low discrimination**					**23511002**
Actinomyces viscosus		86.03	0.69	MNS 96		
Actinomyces odontolyticus		7.90	0.58			
Lactobacillus fermentum		5.56	0.62			
23511003	**Very good identification**					**23511003**
Actinomyces viscosus		97.36	0.60	MNS 96		
Actinomyces naeslundii		2.13	0.56			
Actinomyces odontolyticus		0.27	0.24			
23511020	**Low discrimination**					**23511020**
Actinomyces odontolyticus		44.18	0.48	RAF 3		
Lactobacillus fermentum		39.07	0.54	NAG 14		
Actinomyces naeslundii		8.83	0.53			
23511021	**Good identification**					**23511021**
Actinomyces naeslundii		92.24	0.56	MNS 88		
Actinomyces odontolyticus		3.11	0.14			
Lactobacillus fermentum		2.75	0.20			
23511022	**Low discrimination**					**23511022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		40.04	0.54	NIT 1		
Actinomyces odontolyticus		27.75	0.44			
Actinomyces viscosus		18.76	0.34			
23511023	**Low discrimination**					**23511023**
Actinomyces naeslundii		66.17	0.56	MNS 88		
Actinomyces viscosus		30.49	0.26			
Lactobacillus fermentum		1.97	0.20			
23511041	**Good identification**					**23511041**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23511060	**Low discrimination**					**23511060**
Actinomyces naeslundii		48.17	0.40	MNS 88		
Clostridium perfringens		25.84	0.13			
Actinomyces odontolyticus		13.80	0.14			
23511061	**Good identification**					**23511061**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23511062	**Low discrimination**					**23511062**
Actinomyces naeslundii		52.07	0.40	MNS 88		
Lactobacillus fermentum		13.19	0.20			
Clostridium perfringens		13.14	0.08			
23511063	**Good identification**					**23511063**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23511100	**Low discrimination**					**23511100**
Lactobacillus fermentum		43.64	0.66	NIT 1		
Propionibacterium propionicum		19.87	0.60			
Actinomyces odontolyticus		19.56	0.53			
23511101	**Low discrimination**					**23511101**
					SPO	
Clostridium chauvoei		57.13	0.47	MNS 90	99	
Actinomyces naeslundii		26.48	0.64	RAF 1	1	
Actinomyces viscosus		7.62	0.30		1	
23511102	**Low discrimination**					**23511102**
Actinomyces viscosus		88.47	0.73	MNS 96		
Lactobacillus fermentum		5.25	0.66			
Propionibacterium propionicum		3.59	0.63			
23511103	**Good identification**					**23511103**
Actinomyces viscosus		93.95	0.65	MNS 96		
Actinomyces naeslundii		3.30	0.64			
Propionibacterium propionicum		1.27	0.47			
23511111	**Low discrimination**					**23511111**
					SPO	
Actinomyces naeslundii		70.72	0.42	MNS 88	1	
Clostridium chauvoei		29.28	0.12	MLZ 5	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23511112	**Low discrimination**						**23511112**
					GRA	CAT	
Actinomyces viscosus		81.94	0.39	MNS 96	99	76	
Capnocytophaga ochracea		5.21	0.17	MLZ 1	1	90	
Lactobacillus fermentum		4.86	0.32		99	1	
23511120	**Low discrimination**						**23511120**
					GRA		
Prevotella melaninogenica		64.16	0.50	NIT 1	1		
Lactobacillus fermentum		14.66	0.58	MNS 90	99		
Clostridium perfringens		7.40	0.40		99		
23511121	**Low discrimination**						**23511121**
					GRA	SPO	
Actinomyces naeslundii		54.81	0.64	MNS 88	99	1	
Clostridium chauvoei		39.41	0.38		99	99	
Prevotella melaninogenica		4.10	0.15		1	1	
23511122	**Low discrimination**						**23511122**
					GRA	CAT	
Capnocytophaga ochracea		45.57	0.50	ESL 99	1	90	
Lactobacillus fermentum		17.34	0.58	MNS 90	99	1	
Prevotella melaninogenica		14.45	0.37		1	1	
23511123	**Low discrimination**						**23511123**
					SPO		
Actinomyces naeslundii		66.72	0.64	MNS 88	1		
Actinomyces viscosus		19.20	0.30		1		
Clostridium chauvoei		8.47	0.25		99		
23511131	**Low discrimination**						**23511131**
					SPO		
Actinomyces naeslundii		87.87	0.42	MNS 88	1		
Clostridium chauvoei		12.13	0.04	MLZ 5	99		
23511133	**Acceptable identification**						**23511133**
					GRA		
Actinomyces naeslundii		91.34	0.42	MNS 88	99		
Capnocytophaga ochracea		8.66	-.01	MLZ 5	1		
23511140	**Good identification**						**23511140**
Propionibacterium propionicum		91.60	0.60	MAN 99			
Actinomyces naeslundii		4.67	0.48				
Lactobacillus fermentum		2.03	0.32				
23511141	**Low discrimination**						**23511141**
					SPO		
Propionibacterium propionicum		56.42	0.44	MAN 99	1		
Actinomyces naeslundii		38.80	0.51	URE 10	1		
Clostridium chauvoei		4.79	0.12		99		
23511142	**Low discrimination**						**23511142**
Propionibacterium propionicum		75.88	0.63	MAN 99			
Actinomyces viscosus		18.91	0.39				
Actinomyces naeslundii		2.58	0.48				
23511143	**Low discrimination**						**23511143**
Propionibacterium propionicum		45.12	0.47	MAN 99			
Actinomyces viscosus		33.74	0.30	URE 10			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		20.69	0.51				
23511161	**Good identification**						**23511161**
					SPO		
Actinomyces naeslundii		96.05	0.51	MNS 88	1		
Clostridium chauvoei		3.95	0.04		99		
23511163	**Good identification**						**23511163**
					GRA	CAT	
Actinomyces naeslundii		95.21	0.51	MNS 88	99	25	
Capnocytophaga ochracea		2.69	-.01		1	90	
Propionibacterium propionicum		2.10	0.12		99	1	
23511200	**Very good identification**						**23511200**
					GRA		
Actinomyces odontolyticus		99.70	0.66	RAF 3	99		
Lactobacillus fermentum		0.22	0.28		99		
Prevotella intermedia		0.04	0.02		1		
23511202	**Good identification**						**23511202**
Actinomyces odontolyticus		94.31	0.63	RAF 3			
Actinomyces viscosus		5.35	0.34				
Lactobacillus fermentum		0.35	0.28				
23511220	**Good identification**						**23511220**
Actinomyces odontolyticus		99.54	0.53	RAF 3			
Lactobacillus fermentum		0.46	0.20	NAG 14			
23511222	**Good identification**						**23511222**
Actinomyces odontolyticus		98.91	0.49	RAF 3			
Lactobacillus fermentum		0.74	0.20	NAG 14			
Actinomyces viscosus		0.35	0.00				
23511300	**Very good identification**						**23511300**
Actinomyces odontolyticus		98.00	0.58	RAF 3			
Lactobacillus fermentum		1.14	0.32				
Propionibacterium propionicum		0.52	0.26				
23511302	**Low discrimination**						**23511302**
Actinomyces odontolyticus		73.78	0.54	RAF 3			
Actinomyces viscosus		23.55	0.39				
Lactobacillus fermentum		1.40	0.32				
23511320	**Low discrimination**						**23511320**
					GRA		
Actinomyces odontolyticus		86.50	0.44	RAF 3	99		
Prevotella melaninogenica		9.00	0.15	NAG 14	1		
Lactobacillus fermentum		2.06	0.24		99		
23511322	**Low discrimination**						**23511322**
					GRA	CAT	
Actinomyces odontolyticus		81.38	0.41	RAF 3	99	10	
Capnocytophaga ochracea		8.30	0.15	NAG 14	1	90	
Lactobacillus fermentum		3.16	0.24		99	1	
23511400	**Very good identification**						**23511400**
Lactobacillus fermentum		97.19	0.62	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		2.27	0.27				
Actinomyces meyeri		0.22	0.06				
23511402	**Low discrimination**						**23511402**
Lactobacillus fermentum		85.43	0.62	NIT 1			
Actinomyces viscosus		13.34	0.34				
Actinomyces odontolyticus		1.23	0.24				
23511420	**Good identification**						**23511420**
Lactobacillus fermentum		98.51	0.54	NIT 1			
Actinomyces odontolyticus		1.13	0.14				
Bifidobacterium longum subsp. longum		0.37	0.05				
23511422	**Good identification**						**23511422**
					GRA	CAT	
Lactobacillus fermentum		97.74	0.54	NIT 1	99	1	
Capnocytophaga ochracea		0.75	0.02		1	90	
Actinomyces odontolyticus		0.68	0.10		99	10	
23511500	**Very good identification**						**23511500**
Lactobacillus fermentum		96.29	0.66	NIT 1			
Bifidobacterium longum subsp. longum		1.82	0.29				
Actinomyces meyeri		0.54	0.16				
23511502	**Low discrimination**						**23511502**
Lactobacillus fermentum		82.10	0.66	NIT 1			
Actinomyces viscosus		13.97	0.39				
Bifidobacterium longum subsp. longum		1.55	0.29				
23511520	**Low discrimination**						**23511520**
					GRA	CAT	
Lactobacillus fermentum		74.97	0.58	NIT 1	99	1	
Bifidobacterium longum subsp. longum		17.00	0.40		99	1	
Capnocytophaga ochracea		3.86	0.20		1	90	
23511522	**Low discrimination**						**23511522**
					GRA	CAT	
Lactobacillus fermentum		58.46	0.58	NIT 1	99	1	
Capnocytophaga ochracea		27.12	0.37		1	90	
Bifidobacterium longum subsp. longum		13.26	0.40		99	1	
23511530	**Good identification**						**23511530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.06	0.44	NIT 1	99	1	
Lactobacillus fermentum		2.51	0.24	MNS 87	99	1	
Capnocytophaga ochracea		1.43	0.04		1	90	
23511532	**Low discrimination**						**23511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		86.23	0.44	NIT 1	99	1	
Capnocytophaga ochracea		11.51	0.20	MNS 87	1	90	
Lactobacillus fermentum		2.26	0.24		99	1	
23512000	**Low discrimination**						**23512000**
					COC	GRA	
Propionibacterium propionicum		86.54	0.60	LAC 99	1	99	
Megasphaera elsdenii		7.74	0.32		99	1	
Actinomyces odontolyticus		4.09	0.30		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23512001	**Acceptable identification**						**23512001**
Propionibacterium propionicum		90.03	0.44	LAC 99			
Actinomyces naeslundii		9.97	0.37	URE 10			
23512002	**Low discrimination**						**23512002**
Propionibacterium propionicum		79.36	0.63	LAC 99			
Actinomyces viscosus		18.15	0.38				
Actinomyces odontolyticus		1.53	0.27				
23512003	**Low discrimination**						**23512003**
Propionibacterium propionicum		56.82	0.47	LAC 99			
Actinomyces viscosus		38.99	0.30	URE 10			
Actinomyces naeslundii		4.19	0.37				
23512021	**Good identification**						**23512021**
Actinomyces naeslundii		100.0	0.37	MNS 88			
23512023	**Low discrimination**						**23512023**
Actinomyces naeslundii		87.97	0.37	MNS 88			
Propionibacterium propionicum		12.03	0.12	GAL 96			
23512040	**Good identification**						**23512040**
					COC	GRA	
Propionibacterium propionicum		94.37	0.60	LAC 99	1	99	
Megasphaera elsdenii		5.63	0.29		99	1	
23512041	**Good identification**						**23512041**
Propionibacterium propionicum		98.08	0.44	LAC 99			
Actinomyces naeslundii		1.92	0.25	URE 10			
23512042	**Very good identification**						**23512042**
					COC	GRA	
Propionibacterium propionicum		99.33	0.63	LAC 99	1	99	
Megasphaera elsdenii		0.44	0.12		99	1	
Actinomyces viscosus		0.23	0.04		1	99	
23512043	**Good identification**						**23512043**
Propionibacterium propionicum		98.71	0.47	LAC 99			
Actinomyces naeslundii		1.29	0.25	URE 10			
23512100	**Excellent identification**						**23512100**
Propionibacterium propionicum		99.96	0.94				
Actinomyces naeslundii		0.02	0.42				
Actinomyces odontolyticus		0.02	0.22				
23512101	**Very good identification**						**23512101**
Propionibacterium propionicum		99.66	0.78	URE 10			
Actinomyces naeslundii		0.32	0.45				
Actinomyces viscosus		0.02	0.00				
23512102	**Excellent identification**						**23512102**
Propionibacterium propionicum		99.57	0.97				
Actinomyces viscosus		0.41	0.42				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.02	0.42			
23512103	**Very good identification**					**23512103**
Propionibacterium propionicum		98.58	0.81	URE 10		
Actinomyces viscosus		1.22	0.34			
Actinomyces naeslundii		0.21	0.45			
23512110	**Very good identification**					**23512110**
Propionibacterium propionicum		100.0	0.60	MLZ 1		
23512111	**Good identification**					**23512111**
Propionibacterium propionicum		98.37	0.44	MLZ 1		
Actinomyces naeslundii		1.63	0.23	URE 10		
23512112	**Very good identification**					**23512112**
Propionibacterium propionicum		99.59	0.63	MLZ 1		
Actinomyces viscosus		0.41	0.08			
23512113	**Good identification**					**23512113**
Propionibacterium propionicum		97.72	0.47	MLZ 1		
Actinomyces viscosus		1.20	0.00	URE 10		
Actinomyces naeslundii		1.08	0.23			
23512120	**Very good identification**					**23512120**
Propionibacterium propionicum		97.49	0.60	NAG 1		
Actinomyces naeslundii		2.28	0.42			
Actinomyces odontolyticus		0.24	0.08			
23512121	**Low discrimination**					**23512121**
Propionibacterium propionicum		76.04	0.44	NAG 1		
Actinomyces naeslundii		23.97	0.45	URE 10		
23512122	**Very good identification**					**23512122**
Propionibacterium propionicum		98.07	0.63	NAG 1		
Actinomyces naeslundii		1.53	0.42			
Actinomyces viscosus		0.40	0.08			
23512123	**Low discrimination**					**23512123**
Propionibacterium propionicum		81.80	0.47	NAG 1		
Actinomyces naeslundii		17.19	0.45	URE 10		
Actinomyces viscosus		1.01	0.00			
23512140	**Excellent identification**					**23512140**
Propionibacterium propionicum		100.0	0.94			
Actinomyces naeslundii		0.00	0.29			
23512141	**Very good identification**					**23512141**
Propionibacterium propionicum		99.94	0.78	URE 10		
Actinomyces naeslundii		0.06	0.32			
23512142	**Excellent identification**					**23512142**
Propionibacterium propionicum		99.99	0.97			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.00	0.08			
Actinomyces naeslundii		0.00	0.29			
23512143	**Very good identification**					**23512143**
Propionibacterium propionicum		99.95	0.81	URE 10		
Actinomyces naeslundii		0.04	0.32			
Actinomyces viscosus		0.01	0.00			
23512150	**Very good identification**					**23512150**
Propionibacterium propionicum		100.0	0.60	MLZ 1		
23512151	**Good identification**					**23512151**
Propionibacterium propionicum		100.0	0.44	MLZ 1		
23512152	**Very good identification**					**23512152**
Propionibacterium propionicum		100.0	0.63	MLZ 1		
23512153	**Good identification**					**23512153**
Propionibacterium propionicum		100.0	0.47	MLZ 1		
23512160	**Very good identification**					**23512160**
Propionibacterium propionicum		99.59	0.60	NAG 1		
Actinomyces naeslundii		0.41	0.29			
23512161	**Acceptable identification**					**23512161**
Propionibacterium propionicum		94.73	0.44	NAG 1		
Actinomyces naeslundii		5.27	0.32	URE 10		
23512162	**Very good identification**					**23512162**
Propionibacterium propionicum		99.73	0.63	NAG 1		
Actinomyces naeslundii		0.27	0.29			
23512163	**Good identification**					**23512163**
Propionibacterium propionicum		96.43	0.47	NAG 1		
Actinomyces naeslundii		3.58	0.32	URE 10		
23512200	**Acceptable identification**					**23512200**
Actinomyces odontolyticus		90.08	0.35	RAF 3		
Propionibacterium propionicum		9.92	0.26	MAN 1		
23512300	**Very good identification**					**23512300**
Propionibacterium propionicum		97.19	0.60	RHA 1		
Actinomyces odontolyticus		2.81	0.27			
23512301	**Good identification**					**23512301**
Propionibacterium propionicum		100.0	0.44	RHA 1		
23512302	**Very good identification**					**23512302**
Propionibacterium propionicum		98.43	0.63	RHA 1		
Actinomyces odontolyticus		1.17	0.23			
Actinomyces viscosus		0.40	0.08			
23512303	**Good identification**					**23512303**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		98.78	0.47	RHA 1		
Actinomyces viscosus		1.22	0.00	URE 10		
23512340	**Very good identification**					**23512340**
Propionibacterium propionicum		100.0	0.60	RHA 1		
23512341	**Good identification**					**23512341**
Propionibacterium propionicum		100.0	0.44	RHA 1		
23512342	**Very good identification**					**23512342**
Propionibacterium propionicum		100.0	0.63	RHA 1		
23512343	**Good identification**					**23512343**
Propionibacterium propionicum		100.0	0.47	RHA 1		
23512500	**Very good identification**					**23512500**
Propionibacterium propionicum		100.0	0.60	ARA 1		
23512501	**Good identification**					**23512501**
Propionibacterium propionicum		100.0	0.44	ARA 1		
23512502	**Very good identification**					**23512502**
Propionibacterium propionicum		99.59	0.63	ARA 1		
Actinomyces viscosus		0.41	0.08			
23512503	**Good identification**					**23512503**
Propionibacterium propionicum		98.78	0.47	ARA 1		
Actinomyces viscosus		1.22	0.00	URE 10		
23512540	**Very good identification**					**23512540**
Propionibacterium propionicum		100.0	0.60	ARA 1		
23512541	**Good identification**					**23512541**
Propionibacterium propionicum		100.0	0.44	ARA 1		
23512542	**Very good identification**					**23512542**
Propionibacterium propionicum		100.0	0.63	ARA 1		
23512543	**Good identification**					**23512543**
Propionibacterium propionicum		100.0	0.47	ARA 1		
23513000	**Low discrimination**					**23513000**
Propionibacterium propionicum		79.92	0.60	LAC 99		
Actinomyces naeslundii		15.73	0.58			
Actinomyces odontolyticus		2.52	0.27			
23513001	**Low discrimination**					**23513001**
Actinomyces naeslundii		72.66	0.61	MNS 88		
Propionibacterium propionicum		27.34	0.44			
23513002	**Low discrimination**					**23513002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		77.32	0.63	LAC 99		
Actinomyces viscosus		10.84	0.34			
Actinomyces naeslundii		10.15	0.58			
23513003	**Low discrimination**					**23513003**
Actinomyces naeslundii		55.50	0.61	MNS 88		
Propionibacterium propionicum		31.32	0.47			
Actinomyces viscosus		13.18	0.26			
23513010	**Low discrimination**					**23513010**
Actinomyces naeslundii		50.64	0.36	MNS 88		
Propionibacterium propionicum		49.36	0.26	MLZ 5		
23513011	**Good identification**					**23513011**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23513013	**Acceptable identification**					**23513013**
Actinomyces naeslundii		90.23	0.39	MNS 88		
Propionibacterium propionicum		9.77	0.12	MLZ 5		
23513020	**Good identification**					**23513020**
Actinomyces naeslundii		90.88	0.58	MNS 88		
Propionibacterium propionicum		4.66	0.26			
Actinomyces odontolyticus		2.37	0.14			
23513021	**Very good identification**					**23513021**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23513022	**Low discrimination**					**23513022**
Actinomyces naeslundii		88.75	0.58	MNS 88		
Propionibacterium propionicum		6.83	0.29			
Lactobacillus fermentum		2.04	0.20			
23513023	**Very good identification**					**23513023**
Actinomyces naeslundii		99.43	0.61	MNS 88		
Propionibacterium propionicum		0.57	0.12			
23513030	**Good identification**					**23513030**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23513031	**Good identification**					**23513031**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23513032	**Good identification**					**23513032**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23513033	**Good identification**					**23513033**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23513040	**Very good identification**					**23513040**
Propionibacterium propionicum		96.64	0.60	LAC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		3.36	0.45			
23513041	**Low discrimination**					**23513041**
Propionibacterium propionicum		68.07	0.44	LAC 99		
Actinomyces naeslundii		31.93	0.48	URE 10		
23513042	**Very good identification**					**23513042**
Propionibacterium propionicum		97.60	0.63	LAC 99		
Actinomyces naeslundii		2.26	0.45			
Actinomyces viscosus		0.14	0.00			
23513043	**Low discrimination**					**23513043**
Propionibacterium propionicum		76.18	0.47	LAC 99		
Actinomyces naeslundii		23.82	0.48	URE 10		
23513060	**Low discrimination**					**23513060**
Actinomyces naeslundii		77.48	0.45	MNS 88		
Propionibacterium propionicum		22.53	0.26			
23513061	**Good identification**					**23513061**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23513062	**Low discrimination**					**23513062**
Actinomyces naeslundii		69.63	0.45	MNS 88		
Propionibacterium propionicum		30.37	0.29			
23513063	**Good identification**					**23513063**
Actinomyces naeslundii		96.87	0.48	MNS 88		
Propionibacterium propionicum		3.13	0.12			
23513100	**Excellent identification**					**23513100**
Propionibacterium propionicum		99.40	0.94			
Actinomyces naeslundii		0.56	0.66			
Lactobacillus fermentum		0.02	0.32			
23513101	**Good identification**					**23513101**
					SPO	
Propionibacterium propionicum		92.83	0.78	URE 10	1	
Actinomyces naeslundii		7.09	0.69		1	
Clostridium chauvoei		0.08	0.12		99	
23513102	**Excellent identification**					**23513102**
Propionibacterium propionicum		99.35	0.97			
Actinomyces naeslundii		0.38	0.66			
Actinomyces viscosus		0.25	0.39			
23513103	**Good identification**					**23513103**
Propionibacterium propionicum		94.47	0.81	URE 10		
Actinomyces naeslundii		4.81	0.69			
Actinomyces viscosus		0.71	0.30			
23513110	**Very good identification**					**23513110**
Propionibacterium propionicum		97.14	0.60	MLZ 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		2.87	0.44				
23513111	**Low discrimination**						**23513111**
Propionibacterium propionicum		71.52	0.44	MLZ 1			
Actinomyces naeslundii		28.48	0.47	URE 10			
23513112	**Very good identification**						**23513112**
Propionibacterium propionicum		97.83	0.63	MLZ 1			
Actinomyces naeslundii		1.92	0.44				
Actinomyces viscosus		0.25	0.04				
23513113	**Low discrimination**						**23513113**
Propionibacterium propionicum		79.03	0.47	MLZ 1			
Actinomyces naeslundii		20.98	0.47	URE 10			
23513120	**Low discrimination**						**23513120**
					GRA		
Propionibacterium propionicum		62.27	0.60	NAG 1	99		
Actinomyces naeslundii		34.89	0.66		99		
Prevotella melaninogenica		2.02	0.15		1		
23513121	**Low discrimination**						**23513121**
					SPO		
Actinomyces naeslundii		88.03	0.69	MNS 88	1		
Propionibacterium propionicum		11.64	0.44		1		
Clostridium chauvoei		0.33	0.04		99		
23513122	**Low discrimination**						**23513122**
					GRA	CAT	
Propionibacterium propionicum		71.47	0.63	NAG 1	99	1	
Actinomyces naeslundii		26.70	0.66		99	25	
Capnocytophaga ochracea		0.93	0.15		1	90	
23513123	**Low discrimination**						**23513123**
Actinomyces naeslundii		83.45	0.69	MNS 88			
Propionibacterium propionicum		16.55	0.47				
23513130	**Low discrimination**						**23513130**
Actinomyces naeslundii		74.49	0.44	MNS 88			
Propionibacterium propionicum		25.51	0.26	MLZ 5			
23513131	**Good identification**						**23513131**
Actinomyces naeslundii		100.0	0.47	MNS 88			
23513132	**Low discrimination**						**23513132**
					GRA	CAT	
Actinomyces naeslundii		63.01	0.44	MNS 88	99	25	
Propionibacterium propionicum		32.37	0.29	MLZ 5	99	1	
Capnocytophaga ochracea		4.62	-.01		1	90	
23513133	**Good identification**						**23513133**
Actinomyces naeslundii		96.33	0.47	MNS 88			
Propionibacterium propionicum		3.67	0.12	MLZ 5			
23513140	**Excellent identification**						**23513140**
Propionibacterium propionicum		99.90	0.94				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.10	0.53				
23513141	**Very good identification**						**23513141**
Propionibacterium propionicum		98.67	0.78	URE 10			
Actinomyces naeslundii		1.33	0.56				
23513142	**Excellent identification**						**23513142**
Propionibacterium propionicum		99.93	0.97				
Actinomyces naeslundii		0.07	0.53				
Actinomyces viscosus		0.00	0.04				
23513143	**Very good identification**						**23513143**
Propionibacterium propionicum		99.11	0.81	URE 10			
Actinomyces naeslundii		0.89	0.56				
23513150	**Very good identification**						**23513150**
Propionibacterium propionicum		99.48	0.60	MLZ 1			
Actinomyces naeslundii		0.52	0.31				
23513151	**Acceptable identification**						**23513151**
Propionibacterium propionicum		93.44	0.44	MLZ 1			
Actinomyces naeslundii		6.57	0.34	URE 10			
23513152	**Very good identification**						**23513152**
Propionibacterium propionicum		99.65	0.63	MLZ 1			
Actinomyces naeslundii		0.35	0.31				
23513153	**Good identification**						**23513153**
Propionibacterium propionicum		95.53	0.47	MLZ 1			
Actinomyces naeslundii		4.47	0.34	URE 10			
23513160	**Good identification**						**23513160**
Propionibacterium propionicum		91.00	0.60	NAG 1			
Actinomyces naeslundii		9.00	0.53				
23513161	**Low discrimination**						**23513161**
Actinomyces naeslundii		57.17	0.56	MNS 88			
Propionibacterium propionicum		42.83	0.44				
23513162	**Good identification**						**23513162**
					GRA	CAT	
Propionibacterium propionicum		93.69	0.63	NAG 1	99	1	
Actinomyces naeslundii		6.18	0.53		99	25	
Capnocytophaga ochracea		0.14	-.01		1	90	
23513163	**Low discrimination**						**23513163**
Propionibacterium propionicum		52.91	0.47	NAG 1			
Actinomyces naeslundii		47.09	0.56	URE 10			
23513300	**Very good identification**						**23513300**
Propionibacterium propionicum		97.56	0.60	RHA 1			
Actinomyces odontolyticus		1.88	0.24				
Actinomyces naeslundii		0.55	0.32				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23513301	**Acceptable identification**					**23513301**
Propionibacterium propionicum		92.90	0.44	RHA 1		
Actinomyces naeslundii		7.10	0.35	URE 10		
23513302	**Very good identification**					**23513302**
Propionibacterium propionicum		98.60	0.63	RHA 1		
Actinomyces odontolyticus		0.78	0.20			
Actinomyces naeslundii		0.37	0.32			
23513303	**Good identification**					**23513303**
Propionibacterium propionicum		95.15	0.47	RHA 1		
Actinomyces naeslundii		4.85	0.35	URE 10		
23513340	**Very good identification**					**23513340**
Propionibacterium propionicum		100.0	0.60	RHA 1		
23513341	**Good identification**					**23513341**
Propionibacterium propionicum		100.0	0.44	RHA 1		
23513342	**Very good identification**					**23513342**
Propionibacterium propionicum		100.0	0.63	RHA 1		
23513343	**Good identification**					**23513343**
Propionibacterium propionicum		100.0	0.47	RHA 1		
23513500	**Very good identification**					**23513500**
Propionibacterium propionicum		97.13	0.60	ARA 1		
Lactobacillus fermentum		2.16	0.32			
Actinomyces naeslundii		0.55	0.32			
23513501	**Acceptable identification**					**23513501**
Propionibacterium propionicum		92.90	0.44	ARA 1		
Actinomyces naeslundii		7.10	0.35	URE 10		
23513502	**Very good identification**					**23513502**
Propionibacterium propionicum		97.77	0.63	ARA 1		
Lactobacillus fermentum		1.45	0.32			
Actinomyces naeslundii		0.37	0.32			
23513503	**Good identification**					**23513503**
Propionibacterium propionicum		95.15	0.47	ARA 1		
Actinomyces naeslundii		4.85	0.35	URE 10		
23513540	**Very good identification**					**23513540**
Propionibacterium propionicum		100.0	0.60	ARA 1		
23513541	**Good identification**					**23513541**
Propionibacterium propionicum		100.0	0.44	ARA 1		
23513542	**Very good identification**					**23513542**
Propionibacterium propionicum		100.0	0.63	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23513543	**Good identification**					**23513543**
Propionibacterium propionicum		100.0	0.47	ARA 1		
23514000	**Very good identification**					**23514000**
Actinomyces odontolyticus		98.73	0.59	RAF 3		
Actinomyces meyeri		0.68	0.18			
Lactobacillus fermentum		0.34	0.24			
23514002	**Low discrimination**					**23514002**
Actinomyces odontolyticus		80.08	0.55	RAF 3		
Actinomyces viscosus		19.47	0.38			
Lactobacillus fermentum		0.45	0.24			
23514020	**Good identification**					**23514020**
Actinomyces odontolyticus		98.73	0.46	RAF 3		
Lactobacillus fermentum		0.70	0.16	NAG 14		
Actinomyces meyeri		0.57	0.04			
23514022	**Good identification**					**23514022**
Actinomyces odontolyticus		97.40	0.42	RAF 3		
Actinomyces viscosus		1.47	0.04	NAG 14		
Lactobacillus fermentum		1.13	0.16			
23514100	**Low discrimination**					**23514100**
Actinomyces odontolyticus		88.38	0.51	RAF 3		
Actinomyces meyeri		7.70	0.29			
Lactobacillus fermentum		1.58	0.28			
23514102	**Low discrimination**					**23514102**
Actinomyces viscosus		56.17	0.42	MNS 96		
Actinomyces odontolyticus		41.05	0.47	XYL 1		
Propionibacterium propionicum		1.40	0.29			
23514120	**Acceptable identification**					**23514120**
Actinomyces odontolyticus		90.12	0.37	RAF 3		
Actinomyces meyeri		6.58	0.14	NAG 14		
Lactobacillus fermentum		3.31	0.20			
23514200	**Very good identification**					**23514200**
Actinomyces odontolyticus		100.0	0.64	RAF 3		
23514202	**Very good identification**					**23514202**
Actinomyces odontolyticus		99.87	0.60	RAF 3		
Actinomyces viscosus		0.13	0.04			
23514220	**Good identification**					**23514220**
Actinomyces odontolyticus		100.0	0.51	RAF 3		
23514222	**Good identification**					**23514222**
Actinomyces odontolyticus		100.0	0.47	RAF 3		
23514300	**Good identification**					**23514300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.55	RAF 3			
23514302	**Good identification**						**23514302**
Actinomyces odontolyticus		99.29	0.52	RAF 3			
Actinomyces viscosus		0.71	0.08				
23514320	**Good identification**						**23514320**
Actinomyces odontolyticus		100.0	0.42	RAF 3			
23514322	**Good identification**						**23514322**
Actinomyces odontolyticus		100.0	0.38	RAF 3			
23515000	**Low discrimination**						**23515000**
Actinomyces odontolyticus		65.16	0.56	RAF 3			
Lactobacillus fermentum		33.56	0.58				
Actinomyces meyeri		1.01	0.21				
23515002	**Low discrimination**						**23515002**
Actinomyces odontolyticus		48.46	0.52	RAF 3			
Lactobacillus fermentum		40.72	0.58				
Actinomyces viscosus		10.83	0.34				
23515020	**Low discrimination**						**23515020**
Lactobacillus fermentum		51.01	0.50	NIT 1			
Actinomyces odontolyticus		48.37	0.43				
Actinomyces meyeri		0.63	0.07				
23515022	**Low discrimination**						**23515022**
					GRA	CAT	
Lactobacillus fermentum		62.20	0.50	NIT 1	99	1	
Actinomyces odontolyticus		36.15	0.39		99	10	
Capnocytophaga ochracea		1.15	0.05		1	90	
23515100	**Low discrimination**						**23515100**
Lactobacillus fermentum		67.85	0.62	NIT 1			
Actinomyces odontolyticus		25.50	0.47				
Actinomyces meyeri		5.00	0.32				
23515102	**Low discrimination**						**23515102**
Lactobacillus fermentum		63.13	0.62	NIT 1			
Actinomyces viscosus		18.29	0.39				
Actinomyces odontolyticus		14.54	0.44				
23515120	**Low discrimination**						**23515120**
					GRA	CAT	
Lactobacillus fermentum		69.43	0.54	NIT 1	99	1	
Actinomyces odontolyticus		12.74	0.34		99	10	
Capnocytophaga ochracea		8.63	0.23		1	90	
23515122	**Low discrimination**						**23515122**
					GRA	CAT	
Capnocytophaga ochracea		48.12	0.39	ESL 99	1	90	
Lactobacillus fermentum		43.01	0.54	MNS 90	99	1	
Actinomyces odontolyticus		4.84	0.30		99	10	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23515200	**Very good identification**						**23515200**
Actinomyces odontolyticus		99.73	0.61	RAF 3			
Lactobacillus fermentum		0.27	0.24				
23515202	**Very good identification**						**23515202**
Actinomyces odontolyticus		99.45	0.57	RAF 3			
Lactobacillus fermentum		0.44	0.24				
Actinomyces viscosus		0.12	0.00				
23515220	**Good identification**						**23515220**
Actinomyces odontolyticus		99.45	0.48	RAF 3			
Lactobacillus fermentum		0.55	0.16	NAG 14			
23515222	**Good identification**						**23515222**
Actinomyces odontolyticus		99.11	0.44	RAF 3			
Lactobacillus fermentum		0.89	0.16	NAG 14			
23515300	**Good identification**						**23515300**
Actinomyces odontolyticus		98.63	0.52	RAF 3			
Lactobacillus fermentum		1.37	0.28				
23515302	**Good identification**						**23515302**
Actinomyces odontolyticus		97.17	0.49	RAF 3			
Lactobacillus fermentum		2.20	0.28				
Actinomyces viscosus		0.64	0.04				
23515320	**Acceptable identification**						**23515320**
					GRA		
Actinomyces odontolyticus		90.06	0.39	RAF 3	99		
Bacteroides vulgatus		7.39	0.06	NAG 14	1		
Lactobacillus fermentum		2.55	0.20		99		
23515322	**Low discrimination**						**23515322**
					GRA	CAT	
Actinomyces odontolyticus		86.25	0.35	RAF 3	99	10	
Parabacteroides distasonis		5.30	0.19	NAG 14	1	76	
Capnocytophaga ochracea		4.46	0.05		1	90	
23515400	**Very good identification**						**23515400**
Lactobacillus fermentum		97.36	0.58	NIT 1			
Actinomyces odontolyticus		1.91	0.22				
Actinomyces meyeri		0.73	0.11				
23515402	**Very good identification**						**23515402**
Lactobacillus fermentum		98.55	0.58	NIT 1			
Actinomyces odontolyticus		1.19	0.18				
Actinomyces viscosus		0.27	0.00				
23515420	**Good identification**						**23515420**
Lactobacillus fermentum		97.21	0.50	NIT 1			
Bifidobacterium longum subsp. longum		1.86	0.14				
Actinomyces odontolyticus		0.93	0.08				
23515422	**Good identification**						**23515422**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		98.13	0.50	NIT 1			
Bifidobacterium longum subsp. longum		1.87	0.14				
23515500	**Low discrimination**						**23515500**
Lactobacillus fermentum		89.17	0.62	NIT 1			
Bifidobacterium longum subsp. longum		8.61	0.37				
Actinomyces meyeri		1.64	0.21				
23515502	**Low discrimination**						**23515502**
Lactobacillus fermentum		89.11	0.62	NIT 1			
Bifidobacterium longum subsp. longum		8.60	0.37				
Bifidobacterium adolescentis		0.80	0.21				
23515510	**Acceptable identification**						**23515510**
Bifidobacterium longum subsp. longum		94.21	0.41	NIT 1			
Lactobacillus fermentum		5.79	0.28	MNS 87			
23515512	**Low discrimination**						**23515512**
Bifidobacterium longum subsp. longum		88.63	0.41	NIT 1			
Bifidobacterium adolescentis		5.92	0.23	MNS 87			
Lactobacillus fermentum		5.45	0.28				
23515520	**Low discrimination**						**23515520**
					GRA		
Bifidobacterium longum subsp. longum		51.92	0.48	NIT 1	99		
Lactobacillus fermentum		44.82	0.54	MNS 87	99		
Bacteroides vulgatus		1.93	0.09		1		
23515522	**Low discrimination**						**23515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		49.17	0.48	NIT 1	99	1	
Lactobacillus fermentum		42.45	0.54	MNS 87	99	1	
Capnocytophaga ochracea		8.38	0.26		1	90	
23515530	**Good identification**						**23515530**
Bifidobacterium longum subsp. longum		99.49	0.52	NIT 1			
Lactobacillus fermentum		0.51	0.20	MNS 87			
23515532	**Good identification**						**23515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.40	0.52	NIT 1	99	1	
Capnocytophaga ochracea		1.09	0.10	MNS 87	1	90	
Lactobacillus fermentum		0.50	0.20		99	1	
23516100	**Very good identification**						**23516100**
Propionibacterium propionicum		99.27	0.60	XYL 1			
Actinomyces odontolyticus		0.73	0.16				
23516101	**Good identification**						**23516101**
Propionibacterium propionicum		100.0	0.44	XYL 1			
23516102	**Very good identification**						**23516102**
Propionibacterium propionicum		99.29	0.63	XYL 1			
Actinomyces viscosus		0.41	0.08				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		0.30	0.13			
23516103	**Good identification**					**23516103**
Propionibacterium propionicum		98.78	0.47	XYL 1		
Actinomyces viscosus		1.22	0.00	URE 10		
23516140	**Very good identification**					**23516140**
Propionibacterium propionicum		100.0	0.60	XYL 1		
23516141	**Good identification**					**23516141**
Propionibacterium propionicum		100.0	0.44	XYL 1		
23516142	**Very good identification**					**23516142**
Propionibacterium propionicum		100.0	0.63	XYL 1		
23516143	**Good identification**					**23516143**
Propionibacterium propionicum		100.0	0.47	XYL 1		
23517100	**Very good identification**					**23517100**
Propionibacterium propionicum		95.42	0.60	XYL 1		
Actinomyces israelii		2.33	0.33			
Lactobacillus fermentum		1.24	0.28			
23517101	**Acceptable identification**					**23517101**
Propionibacterium propionicum		92.90	0.44	XYL 1		
Actinomyces naeslundii		7.10	0.35	URE 10		
23517102	**Very good identification**					**23517102**
Propionibacterium propionicum		96.03	0.63	XYL 1		
Actinomyces israelii		2.34	0.36			
Lactobacillus fermentum		0.83	0.28			
23517103	**Good identification**					**23517103**
Propionibacterium propionicum		95.15	0.47	XYL 1		
Actinomyces naeslundii		4.85	0.35	URE 10		
23517140	**Very good identification**					**23517140**
Propionibacterium propionicum		99.06	0.60	XYL 1		
Actinomyces israelii		0.94	0.26			
23517141	**Good identification**					**23517141**
Propionibacterium propionicum		100.0	0.44	XYL 1		
23517142	**Very good identification**					**23517142**
Propionibacterium propionicum		99.06	0.63	XYL 1		
Actinomyces israelii		0.94	0.29			
23517143	**Good identification**					**23517143**
Propionibacterium propionicum		100.0	0.47	XYL 1		
23520102	**Good identification**					**23520102**
Actinomyces viscosus		95.07	0.35	MNS 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		4.93	0.27	FRU 95		
23521000	**Low discrimination**					**23521000**
Actinomyces naeslundii		79.10	0.36	MNS 88		
Actinomyces odontolyticus		20.90	0.09	FRU 95		
23521001	**Good identification**					**23521001**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23521020	**Good identification**					**23521020**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23521021	**Good identification**					**23521021**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23521022	**Good identification**					**23521022**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23521023	**Good identification**					**23521023**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23521100	**Low discrimination**					**23521100**
					COC	
Actinomyces naeslundii		72.65	0.44	MNS 88	1	
Propionibacterium propionicum		20.36	0.24	FRU 95	1	
Blautia hansenii		6.99	-.03		99	
23521120	**Low discrimination**					**23521120**
					GRA	
Actinomyces naeslundii		73.69	0.44	MNS 88	99	
Clostridium perfringens		17.33	0.11	FRU 95	99	
Prevotella melaninogenica		8.99	-.01		1	
23521121	**Low discrimination**					**23521121**
					SPO	
Actinomyces naeslundii		74.90	0.47	MNS 88	1	
Clostridium chauvoei		25.10	0.16	FRU 95	99	
23521122	**Acceptable identification**					**23521122**
Actinomyces naeslundii		90.04	0.44	MNS 88		
Clostridium perfringens		9.96	0.05	FRU 95		
23521123	**Acceptable identification**					**23521123**
					SPO	
Actinomyces naeslundii		94.42	0.47	MNS 88	1	
Clostridium chauvoei		5.58	0.03	FRU 95	99	
23522100	**Very good identification**					**23522100**
Propionibacterium propionicum		99.71	0.58	FRU 99		
Actinomyces naeslundii		0.29	0.25			
23522101	**Good identification**					**23522101**
Propionibacterium propionicum		96.21	0.42	FRU 99		
Actinomyces naeslundii		3.79	0.28	URE 10		
23522102	**Very good identification**					**23522102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.61	0.61	FRU 99		
Actinomyces viscosus		0.20	0.01			
Actinomyces naeslundii		0.19	0.25			
23522103	**Good identification**					**23522103**
Propionibacterium propionicum		97.44	0.45	FRU 99		
Actinomyces naeslundii		2.56	0.28	URE 10		
23522140	**Very good identification**					**23522140**
Propionibacterium propionicum		100.0	0.58	FRU 99		
23522141	**Good identification**					**23522141**
Propionibacterium propionicum		100.0	0.42	FRU 99		
23522142	**Very good identification**					**23522142**
Propionibacterium propionicum		100.0	0.61	FRU 99		
23522143	**Good identification**					**23522143**
Propionibacterium propionicum		100.0	0.45	FRU 99		
23523000	**Low discrimination**					**23523000**
Actinomyces naeslundii		70.88	0.41	MNS 88		
Propionibacterium propionicum		29.12	0.24	FRU 95		
23523001	**Good identification**					**23523001**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23523002	**Low discrimination**					**23523002**
Actinomyces naeslundii		61.87	0.41	MNS 88		
Propionibacterium propionicum		38.13	0.27	FRU 95		
23523003	**Good identification**					**23523003**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23523020	**Good identification**					**23523020**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23523021	**Good identification**					**23523021**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23523022	**Good identification**					**23523022**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23523023	**Good identification**					**23523023**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23523100	**Good identification**					**23523100**
Propionibacterium propionicum		93.46	0.58	FRU 99		
Actinomyces naeslundii		6.54	0.49			
23523101	**Low discrimination**					**23523101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		51.43	0.42	FRU 99		
Actinomyces naeslundii		48.58	0.52	URE 10		
23523102	**Very good identification**					**23523102**
Propionibacterium propionicum		95.54	0.61	FRU 99		
Actinomyces naeslundii		4.46	0.49			
23523103	**Low discrimination**					**23523103**
Propionibacterium propionicum		61.36	0.45	FRU 99		
Actinomyces naeslundii		38.64	0.52	URE 10		
23523120	**Low discrimination**					**23523120**
Actinomyces naeslundii		87.39	0.49	MNS 88		
Propionibacterium propionicum		12.62	0.24	FRU 95		
23523121	**Good identification**					**23523121**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23523122	**Low discrimination**					**23523122**
Actinomyces naeslundii		82.20	0.49	MNS 88		
Propionibacterium propionicum		17.80	0.27	FRU 95		
23523123	**Good identification**					**23523123**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23523140	**Very good identification**					**23523140**
Propionibacterium propionicum		98.78	0.58	FRU 99		
Actinomyces naeslundii		1.22	0.36			
23523141	**Low discrimination**					**23523141**
Propionibacterium propionicum		85.71	0.42	FRU 99		
Actinomyces naeslundii		14.29	0.39	URE 10		
23523142	**Very good identification**					**23523142**
Propionibacterium propionicum		99.18	0.61	FRU 99		
Actinomyces naeslundii		0.82	0.36			
23523143	**Low discrimination**					**23523143**
Propionibacterium propionicum		90.00	0.45	FRU 99		
Actinomyces naeslundii		10.00	0.39	URE 10		
23523160	**Low discrimination**					**23523160**
Actinomyces naeslundii		55.01	0.36	MNS 88		
Propionibacterium propionicum		44.99	0.24	FRU 95		
23523161	**Good identification**					**23523161**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23523163	**Good identification**					**23523163**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23530001	**Low discrimination**					**23530001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		58.84	0.37	MNS 88		
Actinomyces viscosus		41.17	0.11	GAL 96		
23530002	**Good identification**					**23530002**
Actinomyces viscosus		97.92	0.53	MNS 96		
Actinomyces odontolyticus		1.11	0.27	TRE 7		
Propionibacterium propionicum		0.48	0.27			
23530003	**Good identification**					**23530003**
Actinomyces viscosus		98.58	0.45	MNS 96		
Actinomyces naeslundii		1.42	0.37	TRE 7		
23530021	**Good identification**					**23530021**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23530023	**Low discrimination**					**23530023**
Actinomyces naeslundii		58.84	0.37	MNS 88		
Actinomyces viscosus		41.17	0.11	GAL 96		
23530100	**Low discrimination**					**23530100**
Propionibacterium propionicum		89.51	0.58	MAN 99		
Actinomyces viscosus		5.02	0.23			
Actinomyces naeslundii		2.55	0.42			
23530101	**Low discrimination**					**23530101**
Propionibacterium propionicum		64.37	0.42	MAN 99		
Actinomyces naeslundii		24.80	0.45	URE 10		
Actinomyces viscosus		10.84	0.15			
23530102	**Low discrimination**					**23530102**
Actinomyces viscosus		78.30	0.57	MNS 96		
Propionibacterium propionicum		21.14	0.61	TRE 7		
Actinomyces naeslundii		0.40	0.42			
23530103	**Low discrimination**					**23530103**
Actinomyces viscosus		89.84	0.49	MNS 96		
Propionibacterium propionicum		8.09	0.45	TRE 7		
Actinomyces naeslundii		2.08	0.45			
23530120	**Low discrimination**					**23530120**
Actinomyces naeslundii		56.98	0.42	MNS 88		
Propionibacterium propionicum		20.17	0.24	GAL 96		
Clostridium perfringens		16.92	0.11			
23530121	**Good identification**					**23530121**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23530123	**Low discrimination**					**23530123**
Actinomyces naeslundii		69.59	0.45	MNS 88		
Actinomyces viscosus		30.41	0.15	GAL 96		
23530140	**Very good identification**					**23530140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.50	0.58	MAN 99		
Actinomyces naeslundii		0.50	0.29			
23530141	**Acceptable identification**					**23530141**
Propionibacterium propionicum		93.64	0.42	MAN 99		
Actinomyces naeslundii		6.37	0.32	URE 10		
23530142	**Very good identification**					**23530142**
Propionibacterium propionicum		96.08	0.61	MAN 99		
Actinomyces viscosus		3.60	0.23			
Actinomyces naeslundii		0.32	0.29			
23530143	**Low discrimination**					**23530143**
Propionibacterium propionicum		86.39	0.45	MAN 99		
Actinomyces viscosus		9.70	0.15	URE 10		
Actinomyces naeslundii		3.92	0.32			
23530200	**Good identification**					**23530200**
Actinomyces odontolyticus		100.0	0.35	RAF 3		
23531000	**Low discrimination**					**23531000**
					CAT	
Actinomyces naeslundii		66.10	0.58	MNS 88	25	
Actinomyces odontolyticus		10.57	0.27		10	
Propionibacterium granulosum		6.39	0.26		90	
23531001	**Very good identification**					**23531001**
Actinomyces naeslundii		98.25	0.61	MNS 88		
Actinomyces viscosus		1.76	0.07			
23531002	**Low discrimination**					**23531002**
Actinomyces viscosus		86.39	0.49	MNS 96		
Actinomyces naeslundii		10.85	0.58	TRE 7		
Actinomyces odontolyticus		1.06	0.24			
23531003	**Low discrimination**					**23531003**
Actinomyces viscosus		63.89	0.41	MNS 96		
Actinomyces naeslundii		36.12	0.61	TRE 7		
23531010	**Low discrimination**					**23531010**
Actinomyces naeslundii		79.98	0.36	MNS 88		
Propionibacterium granulosum		20.02	0.11	MLZ 5		
23531011	**Good identification**					**23531011**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23531013	**Low discrimination**					**23531013**
Actinomyces naeslundii		74.66	0.39	MNS 88		
Actinomyces viscosus		25.34	0.07	MLZ 5		
23531020	**Low discrimination**					**23531020**
Actinomyces naeslundii		64.57	0.58	MNS 88		
Clostridium perfringens		30.70	0.30			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		1.68	0.14			
23531021	**Very good identification**					**23531021**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23531022	**Low discrimination**					**23531022**
Actinomyces naeslundii		74.45	0.58	MNS 88		
Clostridium perfringens		16.66	0.25			
Actinomyces viscosus		5.99	0.15			
23531023	**Very good identification**					**23531023**
Actinomyces naeslundii		98.25	0.61	MNS 88		
Actinomyces viscosus		1.76	0.07			
23531030	**Good identification**					**23531030**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23531031	**Good identification**					**23531031**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23531032	**Good identification**					**23531032**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23531033	**Good identification**					**23531033**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23531040	**Low discrimination**					**23531040**
Actinomyces naeslundii		80.78	0.45	MNS 88		
Propionibacterium propionicum		19.22	0.24			
23531041	**Good identification**					**23531041**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23531042	**Low discrimination**					**23531042**
Actinomyces naeslundii		55.17	0.45	MNS 88		
Actinomyces viscosus		25.14	0.15			
Propionibacterium propionicum		19.69	0.27			
23531043	**Acceptable identification**					**23531043**
Actinomyces naeslundii		90.81	0.48	MNS 88		
Actinomyces viscosus		9.19	0.07			
23531060	**Low discrimination**					**23531060**
Actinomyces naeslundii		66.09	0.45	MNS 88		
Clostridium perfringens		33.91	0.18			
23531061	**Good identification**					**23531061**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23531062	**Low discrimination**					**23531062**
Actinomyces naeslundii		80.55	0.45	MNS 88		
Clostridium perfringens		19.45	0.12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23531063	**Good identification**					**23531063**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23531100	**Low discrimination**					**23531100**
Propionibacterium propionicum		55.37	0.58	MAN 99		
Actinomyces naeslundii		37.92	0.66			
Actinomyces viscosus		1.90	0.19			
23531101	**Low discrimination**					**23531101**
					SPO	
Actinomyces naeslundii		80.46	0.69	MNS 88	1	
Clostridium chauvoei		9.94	0.30		99	
Propionibacterium propionicum		8.70	0.42		1	
23531102	**Low discrimination**					**23531102**
Actinomyces viscosus		60.33	0.54	MNS 96		
Propionibacterium propionicum		26.58	0.61	TRE 7		
Actinomyces naeslundii		12.13	0.66			
23531103	**Low discrimination**					**23531103**
Actinomyces viscosus		48.27	0.45	MNS 96		
Actinomyces naeslundii		43.69	0.69	TRE 7		
Propionibacterium propionicum		7.09	0.45			
23531110	**Low discrimination**					**23531110**
Actinomyces naeslundii		78.11	0.44	MNS 88		
Propionibacterium propionicum		21.89	0.24	MLZ 5		
23531111	**Good identification**					**23531111**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23531112	**Low discrimination**					**23531112**
Actinomyces naeslundii		42.11	0.44	MNS 88		
Actinomyces viscosus		40.19	0.19	MLZ 5		
Propionibacterium propionicum		17.70	0.27			
23531113	**Low discrimination**					**23531113**
Actinomyces naeslundii		82.50	0.47	MNS 88		
Actinomyces viscosus		17.50	0.11	MLZ 5		
23531120	**Low discrimination**					**23531120**
					GRA	
Clostridium perfringens		52.93	0.45	MNS 99	99	
Actinomyces naeslundii		43.21	0.66		99	
Prevotella melaninogenica		2.50	0.15		1	
23531121	**Very good identification**					**23531121**
					SPO	
Actinomyces naeslundii		95.29	0.69	MNS 88	1	
Clostridium chauvoei		3.92	0.22		99	
Clostridium perfringens		0.79	0.11		84	
23531122	**Low discrimination**					**23531122**
Actinomyces naeslundii		58.56	0.66	MNS 88		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		33.76	0.39				
Actinomyces viscosus		2.94	0.19				
23531123	**Very good identification**						**23531123**
					SPO		
Actinomyces naeslundii		97.82	0.69	MNS 88	1		
Actinomyces viscosus		1.09	0.11		1		
Clostridium chauvoei		0.71	0.09		99		
23531130	**Low discrimination**						**23531130**
Actinomyces naeslundii		80.96	0.44	MNS 88			
Clostridium perfringens		19.04	0.11	MLZ 5			
23531131	**Good identification**						**23531131**
Actinomyces naeslundii		100.0	0.47	MNS 88			
23531132	**Low discrimination**						**23531132**
					GRA	CAT	
Actinomyces naeslundii		84.47	0.44	MNS 88	99	25	
Clostridium perfringens		9.35	0.05	MLZ 5	99	1	
Capnocytophaga ochracea		6.19	-.01		1	90	
23531133	**Good identification**						**23531133**
Actinomyces naeslundii		100.0	0.47	MNS 88			
23531140	**Low discrimination**						**23531140**
Propionibacterium propionicum		88.83	0.58	MAN 99			
Actinomyces naeslundii		10.74	0.53				
Clostridium perfringens		0.44	0.07				
23531141	**Low discrimination**						**23531141**
Actinomyces naeslundii		62.00	0.56	MNS 88			
Propionibacterium propionicum		38.00	0.42				
23531142	**Good identification**						**23531142**
Propionibacterium propionicum		90.49	0.61	MAN 99			
Actinomyces naeslundii		7.29	0.53				
Actinomyces viscosus		2.08	0.19				
23531143	**Low discrimination**						**23531143**
Actinomyces naeslundii		50.44	0.56	MNS 88			
Propionibacterium propionicum		46.37	0.45				
Actinomyces viscosus		3.19	0.11				
23531161	**Good identification**						**23531161**
Actinomyces naeslundii		100.0	0.56	MNS 88			
23531162	**Low discrimination**						**23531162**
Actinomyces naeslundii		56.52	0.53	MNS 88			
Clostridium perfringens		35.16	0.27				
Propionibacterium propionicum		7.09	0.27				
23531163	**Good identification**						**23531163**
Actinomyces naeslundii		100.0	0.56	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23532000	**Very good identification**					**23532000**
					CAT	
Propionibacterium propionicum		96.43	0.58	LAC 99	1	
Actinomyces naeslundii		1.88	0.39		25	
Propionibacterium granulosum		1.69	0.24		90	
23532001	**Low discrimination**					**23532001**
Propionibacterium propionicum		79.19	0.42	LAC 99		
Actinomyces naeslundii		20.81	0.42	URE 10		
23532002	**Very good identification**					**23532002**
Propionibacterium propionicum		96.44	0.61	LAC 99		
Actinomyces viscosus		2.03	0.19			
Actinomyces naeslundii		1.25	0.39			
23532003	**Low discrimination**					**23532003**
Propionibacterium propionicum		80.76	0.45	LAC 99		
Actinomyces naeslundii		14.15	0.42	URE 10		
Actinomyces viscosus		5.10	0.11			
23532020	**Low discrimination**					**23532020**
					CAT	
Actinomyces naeslundii		57.33	0.39	MNS 88	25	
Propionibacterium propionicum		29.76	0.24	GAL 96	1	
Propionibacterium granulosum		12.91	0.13		90	
23532021	**Good identification**					**23532021**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23532022	**Low discrimination**					**23532022**
Actinomyces naeslundii		56.23	0.39	MNS 88		
Propionibacterium propionicum		43.77	0.27	GAL 96		
23532023	**Good identification**					**23532023**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23532040	**Very good identification**					**23532040**
Propionibacterium propionicum		99.66	0.58	LAC 99		
Actinomyces naeslundii		0.34	0.26			
23532041	**Good identification**					**23532041**
Propionibacterium propionicum		95.57	0.42	LAC 99		
Actinomyces naeslundii		4.43	0.29	URE 10		
23532042	**Very good identification**					**23532042**
Propionibacterium propionicum		99.77	0.61	LAC 99		
Actinomyces naeslundii		0.23	0.26			
23532043	**Good identification**					**23532043**
Propionibacterium propionicum		97.00	0.45	LAC 99		
Actinomyces naeslundii		3.00	0.29	URE 10		
23532100	**Excellent identification**					**23532100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.94	0.93			
Actinomyces naeslundii		0.06	0.47			
23532101	**Very good identification**					**23532101**
Propionibacterium propionicum		99.25	0.76	URE 10		
Actinomyces naeslundii		0.75	0.50			
23532102	**Excellent identification**					**23532102**
Propionibacterium propionicum		99.93	0.96			
Actinomyces viscosus		0.04	0.23			
Actinomyces naeslundii		0.04	0.47			
23532103	**Very good identification**					**23532103**
Propionibacterium propionicum		99.39	0.79	URE 10		
Actinomyces naeslundii		0.50	0.50			
Actinomyces viscosus		0.11	0.15			
23532110	**Very good identification**					**23532110**
Propionibacterium propionicum		99.71	0.58	MLZ 1		
Actinomyces naeslundii		0.29	0.25			
23532111	**Good identification**					**23532111**
Propionibacterium propionicum		96.21	0.42	MLZ 1		
Actinomyces naeslundii		3.79	0.28	URE 10		
23532112	**Very good identification**					**23532112**
Propionibacterium propionicum		99.81	0.61	MLZ 1		
Actinomyces naeslundii		0.19	0.25			
23532113	**Good identification**					**23532113**
Propionibacterium propionicum		97.44	0.45	MLZ 1		
Actinomyces naeslundii		2.56	0.28	URE 10		
23532120	**Good identification**					**23532120**
Propionibacterium propionicum		94.75	0.58	NAG 1		
Actinomyces naeslundii		5.25	0.47			
23532121	**Low discrimination**					**23532121**
Propionibacterium propionicum		57.22	0.42	NAG 1		
Actinomyces naeslundii		42.79	0.50	URE 10		
23532122	**Very good identification**					**23532122**
Propionibacterium propionicum		96.44	0.61	NAG 1		
Actinomyces naeslundii		3.56	0.47			
23532123	**Low discrimination**					**23532123**
Propionibacterium propionicum		66.73	0.45	NAG 1		
Actinomyces naeslundii		33.27	0.50	URE 10		
23532140	**Excellent identification**					**23532140**
Propionibacterium propionicum		99.99	0.93			
Actinomyces naeslundii		0.01	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23532141	**Very good identification**					**23532141**
Propionibacterium propionicum		99.87	0.76	URE 10		
Actinomyces naeslundii		0.13	0.37			
23532142	**Excellent identification**					**23532142**
Propionibacterium propionicum		99.99	0.96			
Actinomyces naeslundii		0.01	0.34			
23532143	**Very good identification**					**23532143**
Propionibacterium propionicum		99.91	0.79	URE 10		
Actinomyces naeslundii		0.09	0.37			
23532150	**Very good identification**					**23532150**
Propionibacterium propionicum		100.0	0.58	MLZ 1		
23532151	**Good identification**					**23532151**
Propionibacterium propionicum		100.0	0.42	MLZ 1		
23532152	**Very good identification**					**23532152**
Propionibacterium propionicum		100.0	0.61	MLZ 1		
23532153	**Good identification**					**23532153**
Propionibacterium propionicum		100.0	0.45	MLZ 1		
23532160	**Very good identification**					**23532160**
Propionibacterium propionicum		99.03	0.58	NAG 1		
Actinomyces naeslundii		0.97	0.34			
23532161	**Low discrimination**					**23532161**
Propionibacterium propionicum		88.34	0.42	NAG 1		
Actinomyces naeslundii		11.66	0.37	URE 10		
23532162	**Very good identification**					**23532162**
Propionibacterium propionicum		99.35	0.61	NAG 1		
Actinomyces naeslundii		0.65	0.34			
23532163	**Acceptable identification**					**23532163**
Propionibacterium propionicum		91.91	0.45	NAG 1		
Actinomyces naeslundii		8.09	0.37	URE 10		
23532300	**Very good identification**					**23532300**
Propionibacterium propionicum		100.0	0.58	RHA 1		
23532301	**Good identification**					**23532301**
Propionibacterium propionicum		100.0	0.42	RHA 1		
23532302	**Very good identification**					**23532302**
Propionibacterium propionicum		100.0	0.61	RHA 1		
23532303	**Good identification**					**23532303**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.45	RHA 1		
23532340	**Very good identification**					**23532340**
Propionibacterium propionicum		100.0	0.58	RHA 1		
23532341	**Good identification**					**23532341**
Propionibacterium propionicum		100.0	0.42	RHA 1		
23532342	**Very good identification**					**23532342**
Propionibacterium propionicum		100.0	0.61	RHA 1		
23532343	**Good identification**					**23532343**
Propionibacterium propionicum		100.0	0.45	RHA 1		
23532500	**Very good identification**					**23532500**
Propionibacterium propionicum		100.0	0.58	ARA 1		
23532501	**Good identification**					**23532501**
Propionibacterium propionicum		100.0	0.42	ARA 1		
23532502	**Very good identification**					**23532502**
Propionibacterium propionicum		100.0	0.61	ARA 1		
23532503	**Good identification**					**23532503**
Propionibacterium propionicum		100.0	0.45	ARA 1		
23532540	**Very good identification**					**23532540**
Propionibacterium propionicum		100.0	0.58	ARA 1		
23532541	**Good identification**					**23532541**
Propionibacterium propionicum		100.0	0.42	ARA 1		
23532542	**Very good identification**					**23532542**
Propionibacterium propionicum		100.0	0.61	ARA 1		
23532543	**Good identification**					**23532543**
Propionibacterium propionicum		100.0	0.45	ARA 1		
23533000	**Low discrimination**					**23533000**
					CAT	
Propionibacterium propionicum		67.25	0.58	LAC 99	1	
Actinomyces naeslundii		31.41	0.63		25	
Propionibacterium granulosum		1.18	0.24		90	
23533001	**Low discrimination**					**23533001**
Actinomyces naeslundii		86.31	0.66	MNS 88		
Propionibacterium propionicum		13.69	0.42			
23533002	**Low discrimination**					**23533002**
Propionibacterium propionicum		75.21	0.61	LAC 99		
Actinomyces naeslundii		23.42	0.63			
Actinomyces viscosus		0.97	0.15			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23533003	**Low discrimination**					**23533003**
Actinomyces naeslundii		80.19	0.66	MNS 88		
Propionibacterium propionicum		19.08	0.45			
Actinomyces viscosus		0.74	0.07			
23533010	**Low discrimination**					**23533010**
					CAT	
Actinomyces naeslundii		66.31	0.41	MNS 88	25	
Propionibacterium propionicum		27.24	0.24	MLZ 5	1	
Propionibacterium granulosum		6.45	0.09		90	
23533011	**Good identification**					**23533011**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23533012	**Low discrimination**					**23533012**
Actinomyces naeslundii		61.87	0.41	MNS 88		
Propionibacterium propionicum		38.13	0.27	MLZ 5		
23533013	**Good identification**					**23533013**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23533020	**Very good identification**					**23533020**
					CAT	
Actinomyces naeslundii		96.99	0.63	MNS 88	25	
Propionibacterium propionicum		2.10	0.24		1	
Propionibacterium granulosum		0.91	0.13		90	
23533021	**Very good identification**					**23533021**
Actinomyces naeslundii		100.0	0.66	MNS 88		
23533022	**Very good identification**					**23533022**
Actinomyces naeslundii		96.86	0.63	MNS 88		
Propionibacterium propionicum		3.14	0.27			
23533023	**Very good identification**					**23533023**
Actinomyces naeslundii		100.0	0.66	MNS 88		
23533030	**Good identification**					**23533030**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23533031	**Good identification**					**23533031**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23533032	**Good identification**					**23533032**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23533033	**Good identification**					**23533033**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23533040	**Good identification**					**23533040**
Propionibacterium propionicum		92.39	0.58	LAC 99		
Actinomyces naeslundii		7.62	0.50			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23533041	**Low discrimination**					**23533041**
Actinomyces naeslundii		52.67	0.53	MNS 88		
Propionibacterium propionicum		47.33	0.42			
23533042	**Good identification**					**23533042**
Propionibacterium propionicum		94.79	0.61	LAC 99		
Actinomyces naeslundii		5.21	0.50			
23533043	**Low discrimination**					**23533043**
Propionibacterium propionicum		57.41	0.45	LAC 99		
Actinomyces naeslundii		42.59	0.53	URE 10		
23533060	**Low discrimination**					**23533060**
Actinomyces naeslundii		89.08	0.50	MNS 88		
Propionibacterium propionicum		10.92	0.24			
23533061	**Good identification**					**23533061**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23533062	**Low discrimination**					**23533062**
Actinomyces naeslundii		84.47	0.50	MNS 88		
Propionibacterium propionicum		15.53	0.27			
23533063	**Good identification**					**23533063**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23533100	**Very good identification**					**23533100**
Propionibacterium propionicum		98.67	0.93			
Actinomyces naeslundii		1.33	0.71			
Actinomyces israelii		0.01	0.25			
23533101	**Low discrimination**					**23533101**
Propionibacterium propionicum		84.65	0.76	URE 10		
Actinomyces naeslundii		15.35	0.74			
23533102	**Excellent identification**					**23533102**
Propionibacterium propionicum		99.08	0.96			
Actinomyces naeslundii		0.89	0.71			
Actinomyces viscosus		0.02	0.19			
23533103	**Low discrimination**					**23533103**
Propionibacterium propionicum		89.16	0.79	URE 10		
Actinomyces naeslundii		10.78	0.74			
Actinomyces viscosus		0.06	0.11			
23533110	**Good identification**					**23533110**
Propionibacterium propionicum		93.46	0.58	MLZ 1		
Actinomyces naeslundii		6.54	0.49			
23533111	**Low discrimination**					**23533111**
Propionibacterium propionicum		51.43	0.42	MLZ 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		48.58	0.52	URE 10		
23533112	**Very good identification**					**23533112**
Propionibacterium propionicum		95.41	0.61	MLZ 1		
Actinomyces naeslundii		4.45	0.49			
Actinomyces israelii		0.14	0.14			
23533113	**Low discrimination**					**23533113**
Propionibacterium propionicum		61.36	0.45	MLZ 1		
Actinomyces naeslundii		38.64	0.52	URE 10		
23533120	**Low discrimination**					**23533120**
Actinomyces naeslundii		56.86	0.71	MNS 88		
Propionibacterium propionicum		42.77	0.58			
Clostridium perfringens		0.36	0.11			
23533121	**Good identification**					**23533121**
Actinomyces naeslundii		94.72	0.74	MNS 88		
Propionibacterium propionicum		5.28	0.42			
23533122	**Low discrimination**					**23533122**
Propionibacterium propionicum		52.94	0.61	NAG 1		
Actinomyces naeslundii		46.92	0.71			
Clostridium perfringens		0.14	0.05			
23533123	**Good identification**					**23533123**
Actinomyces naeslundii		92.29	0.74	MNS 88		
Propionibacterium propionicum		7.71	0.45			
23533130	**Low discrimination**					**23533130**
Actinomyces naeslundii		87.39	0.49	MNS 88		
Propionibacterium propionicum		12.62	0.24	MLZ 5		
23533131	**Good identification**					**23533131**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23533132	**Low discrimination**					**23533132**
Actinomyces naeslundii		82.20	0.49	MNS 88		
Propionibacterium propionicum		17.80	0.27	MLZ 5		
23533133	**Good identification**					**23533133**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23533140	**Excellent identification**					**23533140**
Propionibacterium propionicum		99.76	0.93			
Actinomyces naeslundii		0.24	0.58			
Actinomyces israelii		0.00	0.18			
23533141	**Very good identification**					**23533141**
Propionibacterium propionicum		96.90	0.76	URE 10		
Actinomyces naeslundii		3.10	0.61			
23533142	**Excellent identification**					**23533142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.84	0.96			
Actinomyces naeslundii		0.16	0.58			
Actinomyces israelii		0.00	0.21			
23533143	**Very good identification**					**23533143**
Propionibacterium propionicum		97.91	0.79	URE 10		
Actinomyces naeslundii		2.09	0.61			
23533150	**Very good identification**					**23533150**
Propionibacterium propionicum		98.78	0.58	MLZ 1		
Actinomyces naeslundii		1.22	0.36			
23533151	**Low discrimination**					**23533151**
Propionibacterium propionicum		85.71	0.42	MLZ 1		
Actinomyces naeslundii		14.29	0.39	URE 10		
23533152	**Very good identification**					**23533152**
Propionibacterium propionicum		99.18	0.61	MLZ 1		
Actinomyces naeslundii		0.82	0.36			
23533153	**Low discrimination**					**23533153**
Propionibacterium propionicum		90.00	0.45	MLZ 1		
Actinomyces naeslundii		10.00	0.39	URE 10		
23533160	**Low discrimination**					**23533160**
Propionibacterium propionicum		81.00	0.58	NAG 1		
Actinomyces naeslundii		19.00	0.58			
23533161	**Low discrimination**					**23533161**
Actinomyces naeslundii		76.00	0.61	MNS 88		
Propionibacterium propionicum		24.00	0.42			
23533162	**Low discrimination**					**23533162**
Propionibacterium propionicum		86.48	0.61	NAG 1		
Actinomyces naeslundii		13.53	0.58			
23533163	**Low discrimination**					**23533163**
Actinomyces naeslundii		67.86	0.61	MNS 88		
Propionibacterium propionicum		32.14	0.45			
23533170	**Low discrimination**					**23533170**
Actinomyces naeslundii		55.01	0.36	MNS 88		
Propionibacterium propionicum		44.99	0.24	MLZ 5		
23533171	**Good identification**					**23533171**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23533173	**Good identification**					**23533173**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23533300	**Very good identification**					**23533300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		98.68	0.58	RHA 1		
Actinomyces naeslundii		1.33	0.37			
23533301	**Low discrimination**					**23533301**
Propionibacterium propionicum		84.65	0.42	RHA 1		
Actinomyces naeslundii		15.35	0.40	URE 10		
23533302	**Very good identification**					**23533302**
Propionibacterium propionicum		99.11	0.61	RHA 1		
Actinomyces naeslundii		0.89	0.37			
23533303	**Low discrimination**					**23533303**
Propionibacterium propionicum		89.22	0.45	RHA 1		
Actinomyces naeslundii		10.78	0.40	URE 10		
23533320	**Low discrimination**					**23533320**
Actinomyces naeslundii		57.07	0.37	MNS 88		
Propionibacterium propionicum		42.93	0.24	RHA 1		
23533321	**Good identification**					**23533321**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23533323	**Good identification**					**23533323**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23533340	**Very good identification**					**23533340**
Propionibacterium propionicum		99.76	0.58	RHA 1		
Actinomyces naeslundii		0.24	0.24			
23533341	**Good identification**					**23533341**
Propionibacterium propionicum		96.90	0.42	RHA 1		
Actinomyces naeslundii		3.10	0.27	URE 10		
23533342	**Very good identification**					**23533342**
Propionibacterium propionicum		99.84	0.61	RHA 1		
Actinomyces naeslundii		0.16	0.24			
23533343	**Good identification**					**23533343**
Propionibacterium propionicum		97.91	0.45	RHA 1		
Actinomyces naeslundii		2.09	0.27	URE 10		
23533500	**Very good identification**					**23533500**
Propionibacterium propionicum		97.90	0.58	ARA 1		
Actinomyces naeslundii		1.32	0.37			
Actinomyces israelii		0.78	0.23			
23533501	**Low discrimination**					**23533501**
Propionibacterium propionicum		84.65	0.42	ARA 1		
Actinomyces naeslundii		15.35	0.40	URE 10		
23533502	**Very good identification**					**23533502**
Propionibacterium propionicum		98.34	0.61	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.88	0.37			
Actinomyces israelii		0.78	0.26			
23533503	**Low discrimination**					**23533503**
Propionibacterium propionicum		89.22	0.45	ARA 1		
Actinomyces naeslundii		10.78	0.40	URE 10		
23533520	**Low discrimination**					**23533520**
Actinomyces naeslundii		57.07	0.37	MNS 88		
Propionibacterium propionicum		42.93	0.24	ARA 1		
23533521	**Good identification**					**23533521**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23533523	**Good identification**					**23533523**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23533540	**Very good identification**					**23533540**
Propionibacterium propionicum		99.46	0.58	ARA 1		
Actinomyces israelii		0.31	0.16			
Actinomyces naeslundii		0.24	0.24			
23533541	**Good identification**					**23533541**
Propionibacterium propionicum		96.90	0.42	ARA 1		
Actinomyces naeslundii		3.10	0.27	URE 10		
23533542	**Very good identification**					**23533542**
Propionibacterium propionicum		99.53	0.61	ARA 1		
Actinomyces israelii		0.31	0.19			
Actinomyces naeslundii		0.16	0.24			
23533543	**Good identification**					**23533543**
Propionibacterium propionicum		97.91	0.45	ARA 1		
Actinomyces naeslundii		2.09	0.27	URE 10		
23535520	**Good identification**					**23535520**
Bifidobacterium longum subsp. longum		96.63	0.37	NIT 1		
Lactobacillus fermentum		3.37	0.20	MNS 87		
23535522	**Acceptable identification**					**23535522**
					GRA	
Bifidobacterium longum subsp. longum		92.77	0.37	NIT 1	99	
Parabacteroides distasonis		4.00	0.18	MNS 87	1	
Lactobacillus fermentum		3.24	0.20		99	
23535530	**Good identification**					**23535530**
Bifidobacterium longum subsp. longum		100.0	0.41	NIT 1		
23535532	**Good identification**					**23535532**
					GRA	
Bifidobacterium longum subsp. longum		98.66	0.41	NIT 1	99	
Parabacteroides distasonis		1.35	0.14	MNS 87	1	
23536100	**Very good identification**					**23536100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.58	XYL 1		
23536101	**Good identification**					**23536101**
Propionibacterium propionicum		100.0	0.42	XYL 1		
23536102	**Very good identification**					**23536102**
Propionibacterium propionicum		100.0	0.61	XYL 1		
23536103	**Good identification**					**23536103**
Propionibacterium propionicum		100.0	0.45	XYL 1		
23536140	**Very good identification**					**23536140**
Propionibacterium propionicum		100.0	0.58	XYL 1		
23536141	**Good identification**					**23536141**
Propionibacterium propionicum		100.0	0.42	XYL 1		
23536142	**Very good identification**					**23536142**
Propionibacterium propionicum		100.0	0.61	XYL 1		
23536143	**Good identification**					**23536143**
Propionibacterium propionicum		100.0	0.45	XYL 1		
23537002	**Low discrimination**					**23537002**
Actinomyces israelii		67.90	0.36	SAL 92		
Propionibacterium propionicum		24.48	0.27	CEL 93		
Actinomyces naeslundii		7.62	0.29			
23537100	**Low discrimination**					**23537100**
Propionibacterium propionicum		88.85	0.58	XYL 1		
Actinomyces israelii		9.96	0.43			
Actinomyces naeslundii		1.19	0.37			
23537101	**Low discrimination**					**23537101**
Propionibacterium propionicum		84.65	0.42	XYL 1		
Actinomyces naeslundii		15.35	0.40	URE 10		
23537102	**Low discrimination**					**23537102**
Propionibacterium propionicum		89.20	0.61	XYL 1		
Actinomyces israelii		10.00	0.46			
Actinomyces naeslundii		0.80	0.37			
23537103	**Low discrimination**					**23537103**
Propionibacterium propionicum		89.22	0.45	XYL 1		
Actinomyces naeslundii		10.78	0.40	URE 10		
23537120	**Low discrimination**					**23537120**
Actinomyces naeslundii		43.76	0.37	MNS 88		
Propionibacterium propionicum		32.92	0.24	XYL 1		
Actinomyces israelii		23.32	0.23			
23537121	**Good identification**					**23537121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MNS 88		
23537123	**Good identification**					**23537123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23537140	**Very good identification**					**23537140**
Propionibacterium propionicum		95.61	0.58	XYL 1		
Actinomyces israelii		4.17	0.36			
Actinomyces naeslundii		0.23	0.24			
23537141	**Good identification**					**23537141**
Propionibacterium propionicum		96.90	0.42	XYL 1		
Actinomyces naeslundii		3.10	0.27	URE 10		
23537142	**Very good identification**					**23537142**
Propionibacterium propionicum		95.68	0.61	XYL 1		
Actinomyces israelii		4.17	0.39			
Actinomyces naeslundii		0.15	0.24			
23537143	**Good identification**					**23537143**
Propionibacterium propionicum		97.91	0.45	XYL 1		
Actinomyces naeslundii		2.09	0.27	URE 10		
23537500	**Acceptable identification**					**23537500**
Actinomyces israelii		90.08	0.42	SAL 92		
Propionibacterium propionicum		9.92	0.24	CEL 93		
23537502	**Low discrimination**					**23537502**
Actinomyces israelii		85.74	0.45	SAL 92		
Propionibacterium propionicum		9.44	0.27	CEL 93		
Bifidobacterium adolescentis		4.81	0.17			
23537542	**Low discrimination**					**23537542**
Actinomyces israelii		77.93	0.38	SAL 92		
Propionibacterium propionicum		22.07	0.27	CEL 93		
23540002	**Good identification**					**23540002**
Actinomyces viscosus		99.47	0.40	MNS 96		
Actinomyces odontolyticus		0.53	0.08	FRU 95		
23540102	**Good identification**					**23540102**
Actinomyces viscosus		100.0	0.45	MNS 96		
23540103	**Good identification**					**23540103**
Actinomyces viscosus		100.0	0.37	MNS 96		
23541001	**Good identification**					**23541001**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23541002	**Good identification**					**23541002**
Actinomyces viscosus		97.08	0.37	MNS 96		
Actinomyces naeslundii		2.36	0.33	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		0.56	0.05			
23541021	**Good identification**					**23541021**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23541023	**Good identification**					**23541023**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23541100	**Low discrimination**					**23541100**
					COC	
Actinomyces naeslundii		71.40	0.41	MNS 88	1	
Actinomyces viscosus		18.55	0.07	FRU 95	1	
Blautia hansenii		10.05	-.03		99	
23541101	**Low discrimination**					**23541101**
					SPO	
Actinomyces naeslundii		83.92	0.44	MNS 88	1	
Clostridium chauvoei		11.23	0.06	FRU 95	99	
Actinomyces viscosus		4.85	-.01		1	
23541102	**Good identification**					**23541102**
Actinomyces viscosus		96.26	0.41	MNS 96		
Actinomyces naeslundii		3.74	0.41	FRU 95		
23541121	**Good identification**					**23541121**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23541122	**Low discrimination**					**23541122**
					GRA	
Actinomyces naeslundii		53.33	0.41	MNS 88	99	
Prevotella melaninogenica		19.95	0.05	FRU 95	1	
Actinomyces viscosus		13.85	0.07		99	
23541123	**Acceptable identification**					**23541123**
Actinomyces naeslundii		94.54	0.44	MNS 88		
Actinomyces viscosus		5.46	-.01	FRU 95		
23543000	**Good identification**					**23543000**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23543001	**Good identification**					**23543001**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23543002	**Low discrimination**					**23543002**
Actinomyces naeslundii		82.36	0.38	MNS 88		
Actinomyces viscosus		17.65	0.03	FRU 95		
23543003	**Good identification**					**23543003**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23543020	**Good identification**					**23543020**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23543021	**Good identification**					**23543021**
Actinomyces naeslundii		100.0	0.41	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23543022	**Good identification**					**23543022**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23543023	**Good identification**					**23543023**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23543100	**Low discrimination**					**23543100**
Actinomyces naeslundii		79.47	0.46	MNS 88		
Propionibacterium propionicum		20.53	0.26	FRU 95		
23543101	**Good identification**					**23543101**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23543102	**Low discrimination**					**23543102**
Actinomyces naeslundii		65.73	0.46	MNS 88		
Propionibacterium propionicum		25.48	0.29	FRU 95		
Actinomyces viscosus		8.80	0.07			
23543103	**Acceptable identification**					**23543103**
Actinomyces naeslundii		94.48	0.49	MNS 88		
Actinomyces viscosus		2.81	-.01	FRU 95		
Propionibacterium propionicum		2.71	0.12			
23543120	**Good identification**					**23543120**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23543121	**Good identification**					**23543121**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23543122	**Good identification**					**23543122**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23543123	**Good identification**					**23543123**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23543141	**Good identification**					**23543141**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23543143	**Low discrimination**					**23543143**
Actinomyces naeslundii		86.01	0.36	MNS 88		
Propionibacterium propionicum		13.99	0.12	FRU 95		
23543161	**Good identification**					**23543161**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23543163	**Good identification**					**23543163**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23550002	**Very good identification**					**23550002**
Actinomyces viscosus		99.62	0.62	MNS 96		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces odontolyticus	0.32	0.27			
	Actinomyces naeslundii	0.06	0.32			
23550003	**Good identification**					**23550003**
	Actinomyces viscosus	99.72	0.54	MNS 96		
	Actinomyces naeslundii	0.28	0.35			
23550102	**Very good identification**					**23550102**
	Actinomyces viscosus	99.75	0.67	MNS 96		
	Actinomyces naeslundii	0.10	0.39			
	Propionibacterium propionicum	0.09	0.29			
23550103	**Very good identification**					**23550103**
	Actinomyces viscosus	99.52	0.58	MNS 96		
	Actinomyces naeslundii	0.45	0.42			
	Propionibacterium propionicum	0.03	0.12			
23550120	**Low discrimination**					**23550120**
					GRA	
	Actinomyces naeslundii	67.54	0.39	MNS 88	99	
	Prevotella melaninogenica	15.24	-.01	GAL 96	1	
	Actinomyces odontolyticus	10.29	0.08		99	
23550121	**Good identification**					**23550121**
	Actinomyces naeslundii	100.0	0.42	MNS 88		
23550200	**Good identification**					**23550200**
	Actinomyces odontolyticus	100.0	0.35	RAF 3		
23551000	**Low discrimination**					**23551000**
	Actinomyces naeslundii	54.61	0.55	MNS 88		
	Actinomyces viscosus	22.71	0.24			
	Actinomyces odontolyticus	12.79	0.27			
23551001	**Good identification**					**23551001**
	Actinomyces naeslundii	91.54	0.58	MNS 88		
	Actinomyces viscosus	8.46	0.16			
23551002	**Very good identification**					**23551002**
	Actinomyces viscosus	97.03	0.59	MNS 96		
	Actinomyces naeslundii	2.36	0.55			
	Actinomyces odontolyticus	0.34	0.24			
23551003	**Acceptable identification**					**23551003**
	Actinomyces viscosus	90.15	0.50	MNS 96		
	Actinomyces naeslundii	9.85	0.58			
23551011	**Good identification**					**23551011**
	Actinomyces naeslundii	100.0	0.36	MNS 88		
23551020	**Low discrimination**					**23551020**
	Actinomyces naeslundii	89.75	0.55	MNS 88		
	Actinomyces odontolyticus	3.42	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		3.03	0.20			
23551021	**Very good identification**					**23551021**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23551022	**Low discrimination**					**23551022**
					GRA	
Actinomyces naeslundii		66.27	0.55	MNS 88	99	
Actinomyces viscosus		27.56	0.24		99	
Capnocytophaga ochracea		2.40	0.05		1	
23551023	**Good identification**					**23551023**
Actinomyces naeslundii		91.54	0.58	MNS 88		
Actinomyces viscosus		8.46	0.16			
23551031	**Good identification**					**23551031**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23551033	**Good identification**					**23551033**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23551040	**Good identification**					**23551040**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23551041	**Good identification**					**23551041**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23551043	**Low discrimination**					**23551043**
Actinomyces naeslundii		65.63	0.45	MNS 88		
Actinomyces viscosus		34.37	0.16			
23551060	**Good identification**					**23551060**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23551061	**Good identification**					**23551061**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23551062	**Good identification**					**23551062**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23551063	**Good identification**					**23551063**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23551100	**Low discrimination**					**23551100**
Actinomyces naeslundii		68.52	0.63	MNS 88		
Actinomyces viscosus		17.80	0.29			
Lactobacillus fermentum		3.97	0.32			
23551101	**Good identification**					**23551101**
					SPO	
Actinomyces naeslundii		93.10	0.66	MNS 88	1	
Actinomyces viscosus		5.37	0.21		1	
Clostridium chauvoei		1.53	0.12		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
23551102	**Very good identification**						**23551102**
					GRA		
Actinomyces viscosus		95.26	0.63	MNS 96	99		
Actinomyces naeslundii		3.71	0.63		99		
Capnocytophaga ochracea		0.52	0.23		1		
23551103	**Low discrimination**						**23551103**
Actinomyces viscosus		85.03	0.55	MNS 96			
Actinomyces naeslundii		14.88	0.66				
Propionibacterium propionicum		0.04	0.12				
23551110	**Good identification**						**23551110**
Actinomyces naeslundii		100.0	0.41	MNS 88			
23551111	**Good identification**						**23551111**
Actinomyces naeslundii		100.0	0.44	MNS 88			
23551120	**Low discrimination**						**23551120**
					GRA	CAT	
Actinomyces naeslundii		46.95	0.63	MNS 88	99	25	
Prevotella melaninogenica		43.69	0.33		1	1	
Capnocytophaga ochracea		6.56	0.23		1	90	
23551121	**Very good identification**						**23551121**
					GRA	SPO	
Actinomyces naeslundii		98.84	0.66	MNS 88	99	1	
Prevotella melaninogenica		0.62	-.01		1	1	
Clostridium chauvoei		0.54	0.04		99	99	
23551122	**Low discrimination**						**23551122**
					GRA		
Capnocytophaga ochracea		45.24	0.39	ESL 99	1		
Actinomyces naeslundii		35.97	0.63	MNS 90	99		
Actinomyces viscosus		9.34	0.29		99		
23551123	**Good identification**						**23551123**
					GRA		
Actinomyces naeslundii		93.79	0.66	MNS 88	99		
Actinomyces viscosus		5.41	0.21		99		
Capnocytophaga ochracea		0.79	0.05		1		
23551130	**Low discrimination**						**23551130**
					GRA	CAT	
Actinomyces naeslundii		67.86	0.41	MNS 88	99	25	
Capnocytophaga ochracea		20.02	0.07	MLZ 5	1	90	
Prevotella melaninogenica		12.12	-.01		1	1	
23551131	**Good identification**						**23551131**
Actinomyces naeslundii		100.0	0.44	MNS 88			
23551133	**Good identification**						**23551133**
Actinomyces naeslundii		100.0	0.44	MNS 88			
23551140	**Low discrimination**						**23551140**
Actinomyces naeslundii		84.87	0.50	MNS 88			
Propionibacterium propionicum		12.70	0.26				
Actinomyces israelii		2.43	0.14				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
23551141	**Good identification**						**23551141**
Actinomyces naeslundii		100.0	0.53	MNS 88			
23551143	**Low discrimination**						**23551143**
Actinomyces naeslundii		74.42	0.53	MNS 88			
Actinomyces viscosus		24.34	0.21				
Propionibacterium propionicum		1.24	0.12				
23551160	**Low discrimination**						**23551160**
					GRA	CAT	
Actinomyces naeslundii		84.41	0.50	MNS 88	99	25	
Capnocytophaga ochracea		7.43	0.07		1	90	
Prevotella melaninogenica		4.50	-.01		1	1	
23551161	**Good identification**						**23551161**
Actinomyces naeslundii		100.0	0.53	MNS 88			
23551162	**Low discrimination**						**23551162**
					GRA		
Actinomyces naeslundii		55.81	0.50	MNS 88	99		
Capnocytophaga ochracea		44.19	0.23		1		
23551163	**Good identification**						**23551163**
Actinomyces naeslundii		100.0	0.53	MNS 88			
23552000	**Low discrimination**						**23552000**
Actinomyces naeslundii		51.84	0.36	MNS 88			
Propionibacterium propionicum		48.16	0.26	GAL 96			
23552001	**Good identification**						**23552001**
Actinomyces naeslundii		100.0	0.39	MNS 88			
23552020	**Good identification**						**23552020**
Actinomyces naeslundii		100.0	0.36	MNS 88			
23552021	**Good identification**						**23552021**
Actinomyces naeslundii		100.0	0.39	MNS 88			
23552022	**Good identification**						**23552022**
Actinomyces naeslundii		100.0	0.36	MNS 88			
23552023	**Good identification**						**23552023**
Actinomyces naeslundii		100.0	0.39	MNS 88			
23552100	**Very good identification**						**23552100**
Propionibacterium propionicum		96.84	0.60	CEL 1			
Actinomyces naeslundii		3.00	0.44				
Actinomyces viscosus		0.16	-.02				
23552101	**Low discrimination**						**23552101**
Propionibacterium propionicum		70.53	0.44	CEL 1			
Actinomyces naeslundii		29.47	0.47	URE 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23552102	**Low discrimination**					**23552102**
Propionibacterium propionicum		88.60	0.63	CEL 1		
Actinomyces viscosus		9.58	0.32			
Actinomyces naeslundii		1.83	0.44			
23552103	**Low discrimination**					**23552103**
Propionibacterium propionicum		62.39	0.47	CEL 1		
Actinomyces viscosus		20.23	0.24	URE 10		
Actinomyces naeslundii		17.38	0.47			
23552120	**Low discrimination**					**23552120**
Actinomyces naeslundii		75.39	0.44	MNS 88		
Propionibacterium propionicum		24.61	0.26	GAL 96		
23552121	**Good identification**					**23552121**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23552122	**Low discrimination**					**23552122**
Actinomyces naeslundii		64.83	0.44	MNS 88		
Propionibacterium propionicum		31.74	0.29	GAL 96		
Actinomyces viscosus		3.43	-.02			
23552123	**Good identification**					**23552123**
Actinomyces naeslundii		96.50	0.47	MNS 88		
Propionibacterium propionicum		3.50	0.12	GAL 96		
23552140	**Very good identification**					**23552140**
Propionibacterium propionicum		99.46	0.60	CEL 1		
Actinomyces naeslundii		0.54	0.31			
23552141	**Acceptable identification**					**23552141**
Propionibacterium propionicum		93.13	0.44	CEL 1		
Actinomyces naeslundii		6.87	0.34	URE 10		
23552142	**Very good identification**					**23552142**
Propionibacterium propionicum		99.53	0.63	CEL 1		
Actinomyces naeslundii		0.36	0.31			
Actinomyces viscosus		0.11	-.02			
23552143	**Good identification**					**23552143**
Propionibacterium propionicum		95.32	0.47	CEL 1		
Actinomyces naeslundii		4.69	0.34	URE 10		
23553000	**Good identification**					**23553000**
Actinomyces naeslundii		93.84	0.60	MNS 88		
Propionibacterium propionicum		3.63	0.26			
Actinomyces israelii		2.53	0.24			
23553001	**Very good identification**					**23553001**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23553002	**Low discrimination**					**23553002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		76.18	0.60	MNS 88		
Actinomyces viscosus		16.32	0.24			
Propionibacterium propionicum		4.42	0.29			
23553003	**Very good identification**					**23553003**
Actinomyces naeslundii		95.07	0.63	MNS 88		
Actinomyces viscosus		4.53	0.16			
Propionibacterium propionicum		0.41	0.12			
23553010	**Good identification**					**23553010**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23553011	**Good identification**					**23553011**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23553012	**Good identification**					**23553012**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23553013	**Good identification**					**23553013**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23553020	**Very good identification**					**23553020**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23553021	**Very good identification**					**23553021**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23553022	**Very good identification**					**23553022**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23553023	**Very good identification**					**23553023**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23553030	**Good identification**					**23553030**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23553031	**Good identification**					**23553031**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23553032	**Good identification**					**23553032**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23553033	**Good identification**					**23553033**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23553040	**Low discrimination**					**23553040**
Actinomyces naeslundii		78.20	0.47	MNS 88		
Propionibacterium propionicum		17.15	0.26			
Actinomyces israelii		4.65	0.17			
23553041	**Good identification**					**23553041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MNS 88		
23553042	**Low discrimination**					**23553042**
Actinomyces naeslundii		70.51	0.47	MNS 88		
Propionibacterium propionicum		23.20	0.29			
Actinomyces israelii		6.29	0.20			
23553043	**Good identification**					**23553043**
Actinomyces naeslundii		97.62	0.50	MNS 88		
Propionibacterium propionicum		2.38	0.12			
23553060	**Good identification**					**23553060**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23553061	**Good identification**					**23553061**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23553062	**Good identification**					**23553062**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23553063	**Good identification**					**23553063**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23553100	**Low discrimination**					**23553100**
Propionibacterium propionicum		56.47	0.60	CEL 1		
Actinomyces naeslundii		41.94	0.68			
Actinomyces israelii		1.59	0.34			
23553101	**Good identification**					**23553101**
Actinomyces naeslundii		90.93	0.71	MNS 88		
Propionibacterium propionicum		9.07	0.44			
23553102	**Low discrimination**					**23553102**
Propionibacterium propionicum		62.91	0.63	CEL 1		
Actinomyces naeslundii		31.15	0.68			
Actinomyces viscosus		4.17	0.29			
23553103	**Low discrimination**					**23553103**
Actinomyces naeslundii		84.79	0.71	MNS 88		
Propionibacterium propionicum		12.69	0.47			
Actinomyces viscosus		2.52	0.21			
23553110	**Low discrimination**					**23553110**
Actinomyces naeslundii		73.21	0.46	MNS 88		
Propionibacterium propionicum		18.92	0.26	MLZ 5		
Actinomyces israelii		7.88	0.20			
23553111	**Good identification**					**23553111**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23553112	**Low discrimination**					**23553112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		64.56	0.46	MNS 88			
Propionibacterium propionicum		25.02	0.29	MLZ 5			
Actinomyces israelii		10.43	0.23				
23553113	**Good identification**						**23553113**
Actinomyces naeslundii		97.21	0.49	MNS 88			
Propionibacterium propionicum		2.79	0.12	MLZ 5			
23553120	**Very good identification**						**23553120**
					GRA		
Actinomyces naeslundii		97.96	0.68	MNS 88	99		
Propionibacterium propionicum		1.33	0.26		99		
Prevotella melaninogenica		0.47	-.01		1		
23553121	**Very good identification**						**23553121**
Actinomyces naeslundii		100.0	0.71	MNS 88			
23553122	**Very good identification**						**23553122**
					GRA	CAT	
Actinomyces naeslundii		97.03	0.68	MNS 88	99	25	
Propionibacterium propionicum		1.98	0.29		99	1	
Capnocytophaga ochracea		0.64	0.05		1	90	
23553123	**Very good identification**						**23553123**
Actinomyces naeslundii		99.85	0.71	MNS 88			
Propionibacterium propionicum		0.15	0.12				
23553130	**Good identification**						**23553130**
Actinomyces naeslundii		100.0	0.46	MNS 88			
23553131	**Good identification**						**23553131**
Actinomyces naeslundii		100.0	0.49	MNS 88			
23553132	**Good identification**						**23553132**
Actinomyces naeslundii		100.0	0.46	MNS 88			
23553133	**Good identification**						**23553133**
Actinomyces naeslundii		100.0	0.49	MNS 88			
23553140	**Low discrimination**						**23553140**
Propionibacterium propionicum		87.56	0.60	CEL 1			
Actinomyces naeslundii		11.48	0.55				
Actinomyces israelii		0.96	0.27				
23553141	**Low discrimination**						**23553141**
Actinomyces naeslundii		63.89	0.58	MNS 88			
Propionibacterium propionicum		36.11	0.44				
23553142	**Good identification**						**23553142**
Propionibacterium propionicum		91.05	0.63	CEL 1			
Actinomyces naeslundii		7.96	0.55				
Actinomyces israelii		1.00	0.30				
23553143	**Low discrimination**						**23553143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		54.12	0.58	MNS 88		
Propionibacterium propionicum		45.88	0.47			
23553151	**Good identification**					**23553151**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23553153	**Low discrimination**					**23553153**
Actinomyces naeslundii		86.01	0.36	MNS 88		
Propionibacterium propionicum		13.99	0.12	MLZ 5		
23553160	**Acceptable identification**					**23553160**
Actinomyces naeslundii		92.85	0.55	MNS 88		
Propionibacterium propionicum		7.16	0.26			
23553161	**Very good identification**					**23553161**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23553162	**Low discrimination**					**23553162**
Actinomyces naeslundii		89.64	0.55	MNS 88		
Propionibacterium propionicum		10.36	0.29			
23553163	**Very good identification**					**23553163**
Actinomyces naeslundii		99.15	0.58	MNS 88		
Propionibacterium propionicum		0.85	0.12			
23553171	**Good identification**					**23553171**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23553173	**Good identification**					**23553173**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23553301	**Good identification**					**23553301**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23553303	**Low discrimination**					**23553303**
Actinomyces naeslundii		86.99	0.37	MNS 88		
Propionibacterium propionicum		13.01	0.12	RHA 1		
23553321	**Good identification**					**23553321**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23553323	**Good identification**					**23553323**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23553501	**Good identification**					**23553501**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23553502	**Low discrimination**					**23553502**
Actinomyces israelii		57.39	0.36	SAL 92		
Propionibacterium propionicum		25.16	0.29	XYL 92		
Actinomyces naeslundii		12.46	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23553503	**Low discrimination**					**23553503**
Actinomyces naeslundii		86.99	0.37	MNS 88		
Propionibacterium propionicum		13.01	0.12	ARA 1		
23553521	**Good identification**					**23553521**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23553523	**Good identification**					**23553523**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23555100	**Low discrimination**					**23555100**
Actinomyces israelii		72.40	0.40	SAL 92		
Lactobacillus fermentum		16.51	0.28			
Actinomyces odontolyticus		6.20	0.13			
23555142	**Good identification**					**23555142**
Actinomyces israelii		100.0	0.36	SAL 92		
23555500	**Low discrimination**					**23555500**
Actinomyces israelii		78.21	0.38	SAL 92		
Lactobacillus fermentum		21.80	0.28			
23555502	**Low discrimination**					**23555502**
Actinomyces israelii		63.68	0.41	SAL 92		
Bifidobacterium adolescentis		22.35	0.27			
Lactobacillus fermentum		11.83	0.28			
23557000	**Good identification**					**23557000**
Actinomyces israelii		96.85	0.42	SAL 92		
Actinomyces naeslundii		3.15	0.26			
23557002	**Good identification**					**23557002**
Actinomyces israelii		97.88	0.45	SAL 92		
Actinomyces naeslundii		2.12	0.26			
23557040	**Good identification**					**23557040**
Actinomyces israelii		100.0	0.35	SAL 92		
23557042	**Good identification**					**23557042**
Actinomyces israelii		100.0	0.38	SAL 92		
23557100	**Acceptable identification**					**23557100**
Actinomyces israelii		94.85	0.53	SAL 92		
Propionibacterium propionicum		2.96	0.26			
Actinomyces naeslundii		2.20	0.34			
23557101	**Low discrimination**					**23557101**
Actinomyces naeslundii		77.48	0.37	MNS 88		
Actinomyces israelii		22.52	0.18	XYL 1		
23557102	**Acceptable identification**					**23557102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		94.98	0.56	SAL 92		
Propionibacterium propionicum		2.96	0.29			
Actinomyces naeslundii		1.47	0.34			
23557103	**Low discrimination**					**23557103**
Actinomyces naeslundii		63.07	0.37	MNS 88		
Actinomyces israelii		27.50	0.21	XYL 1		
Propionibacterium propionicum		9.43	0.12			
23557110	**Good identification**					**23557110**
Actinomyces israelii		100.0	0.38	SAL 92		
23557112	**Good identification**					**23557112**
Actinomyces israelii		95.18	0.41	SAL 92		
Bifidobacterium adolescentis		4.82	0.13	MLZ 13		
23557121	**Good identification**					**23557121**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23557122	**Low discrimination**					**23557122**
Actinomyces israelii		80.52	0.35	SAL 92		
Actinomyces naeslundii		19.48	0.34	NAG 6		
23557123	**Good identification**					**23557123**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23557140	**Acceptable identification**					**23557140**
Actinomyces israelii		92.58	0.45	SAL 92		
Propionibacterium propionicum		7.42	0.26			
23557142	**Acceptable identification**					**23557142**
Actinomyces israelii		92.58	0.49	SAL 92		
Propionibacterium propionicum		7.42	0.29			
23557302	**Good identification**					**23557302**
Actinomyces israelii		100.0	0.36	SAL 92		
23557400	**Good identification**					**23557400**
Actinomyces israelii		100.0	0.41	SAL 92		
23557402	**Good identification**					**23557402**
Actinomyces israelii		100.0	0.44	SAL 92		
23557442	**Good identification**					**23557442**
Actinomyces israelii		100.0	0.37	SAL 92		
23557500	**Good identification**					**23557500**
Actinomyces israelii		100.0	0.51	SAL 92		
23557502	**Good identification**					**23557502**
Actinomyces israelii		95.18	0.54	SAL 92		
Bifidobacterium adolescentis		4.82	0.26			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23557510	**Good identification**					**23557510**
Actinomyces israelii		100.0	0.37	SAL 92		
23557512	**Low discrimination**					**23557512**
Actinomyces israelii		70.70	0.40	SAL 92		
Bifidobacterium adolescentis		29.30	0.27	MLZ 13		
23557540	**Good identification**					**23557540**
Actinomyces israelii		100.0	0.44	SAL 92		
23557542	**Good identification**					**23557542**
Actinomyces israelii		98.42	0.47	SAL 92		
Bifidobacterium adolescentis		1.59	0.10			
23561000	**Good identification**					**23561000**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23561001	**Good identification**					**23561001**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23561003	**Low discrimination**					**23561003**
Actinomyces naeslundii		73.81	0.41	MNS 88		
Actinomyces viscosus		26.19	0.09	FRU 95		
23561020	**Good identification**					**23561020**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23561021	**Good identification**					**23561021**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23561022	**Good identification**					**23561022**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23561023	**Good identification**					**23561023**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23561100	**Good identification**					**23561100**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23561101	**Good identification**					**23561101**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23561102	**Low discrimination**					**23561102**
Actinomyces naeslundii		50.07	0.46	MNS 88		
Actinomyces viscosus		49.93	0.22	FRU 95		
23561103	**Low discrimination**					**23561103**
Actinomyces naeslundii		81.86	0.49	MNS 88		
Actinomyces viscosus		18.14	0.14	FRU 95		
23561120	**Good identification**					**23561120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MNS 88		
23561121	**Good identification**					**23561121**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23561122	**Good identification**					**23561122**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23561123	**Good identification**					**23561123**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23561141	**Good identification**					**23561141**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23561143	**Good identification**					**23561143**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23561161	**Good identification**					**23561161**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23561163	**Good identification**					**23561163**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23563000	**Good identification**					**23563000**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23563001	**Good identification**					**23563001**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23563002	**Good identification**					**23563002**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23563003	**Good identification**					**23563003**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23563020	**Good identification**					**23563020**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23563021	**Good identification**					**23563021**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23563022	**Good identification**					**23563022**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23563023	**Good identification**					**23563023**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23563100	**Acceptable identification**					**23563100**
Actinomyces naeslundii		90.18	0.51	MNS 88		
Propionibacterium propionicum		9.82	0.24	FRU 95		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23563101	**Good identification**					**23563101**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23563102	**Low discrimination**					**23563102**
Actinomyces naeslundii		85.96	0.51	MNS 88		
Propionibacterium propionicum		14.04	0.27	FRU 95		
23563103	**Good identification**					**23563103**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23563120	**Good identification**					**23563120**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23563121	**Good identification**					**23563121**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23563122	**Good identification**					**23563122**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23563123	**Good identification**					**23563123**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23563140	**Low discrimination**					**23563140**
Actinomyces naeslundii		61.84	0.38	MNS 88		
Propionibacterium propionicum		38.16	0.24	FRU 95		
23563141	**Good identification**					**23563141**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23563142	**Low discrimination**					**23563142**
Actinomyces naeslundii		51.93	0.38	MNS 88		
Propionibacterium propionicum		48.07	0.27	FRU 95		
23563143	**Good identification**					**23563143**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23563160	**Good identification**					**23563160**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23563161	**Good identification**					**23563161**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23563162	**Good identification**					**23563162**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23563163	**Good identification**					**23563163**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23570000	**Low discrimination**					**23570000**
Actinomyces naeslundii		56.00	0.36	MNS 88	GRA 99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		35.36	0.09	GAL 96	99	
Fusobacterium mortiferum		8.65	0.11		1	
23570001	**Low discrimination**					**23570001**
Actinomyces naeslundii		87.70	0.39	MNS 88		
Actinomyces viscosus		12.30	0.01	GAL 96		
23570002	**Good identification**					**23570002**
Actinomyces viscosus		98.43	0.43	MNS 96		
Actinomyces naeslundii		1.58	0.36	TRE 7		
23570020	**Good identification**					**23570020**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23570021	**Good identification**					**23570021**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23570022	**Low discrimination**					**23570022**
Actinomyces naeslundii		61.30	0.36	MNS 88		
Actinomyces viscosus		38.70	0.09	GAL 96		
23570023	**Low discrimination**					**23570023**
Actinomyces naeslundii		87.70	0.39	MNS 88		
Actinomyces viscosus		12.30	0.01	GAL 96		
23570100	**Low discrimination**					**23570100**
Actinomyces naeslundii		55.93	0.44	MNS 88		
Actinomyces viscosus		22.06	0.13	GAL 96		
Propionibacterium propionicum		14.93	0.24			
23570101	**Acceptable identification**					**23570101**
Actinomyces naeslundii		91.94	0.47	MNS 88		
Actinomyces viscosus		8.06	0.05	GAL 96		
23570102	**Good identification**					**23570102**
Actinomyces viscosus		96.54	0.47	MNS 96		
Actinomyces naeslundii		2.47	0.44	TRE 7		
Propionibacterium propionicum		0.99	0.27			
23570103	**Low discrimination**					**23570103**
Actinomyces viscosus		89.67	0.39	MNS 96		
Actinomyces naeslundii		10.34	0.47	TRE 7		
23570120	**Good identification**					**23570120**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23570121	**Good identification**					**23570121**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23570122	**Low discrimination**					**23570122**
Actinomyces naeslundii		71.72	0.44	MNS 88		
Actinomyces viscosus		28.28	0.13	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23570123	**Acceptable identification**					**23570123**
Actinomyces naeslundii		91.94	0.47	MNS 88		
Actinomyces viscosus		8.06	0.05	GAL 96		
23571000	**Very good identification**					**23571000**
Actinomyces naeslundii		95.81	0.60	MNS 88		
Actinomyces israelii		1.72	0.21			
Actinomyces viscosus		1.55	0.05			
23571001	**Very good identification**					**23571001**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23571002	**Low discrimination**					**23571002**
Actinomyces viscosus		60.85	0.39	MNS 96		
Actinomyces naeslundii		38.12	0.60	TRE 7		
Actinomyces israelii		1.02	0.24			
23571003	**Low discrimination**					**23571003**
Actinomyces naeslundii		73.82	0.63	MNS 88		
Actinomyces viscosus		26.18	0.31			
23571010	**Good identification**					**23571010**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23571011	**Good identification**					**23571011**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23571012	**Low discrimination**					**23571012**
Actinomyces naeslundii		76.55	0.38	MNS 88		
Actinomyces viscosus		23.45	0.05	MLZ 5		
23571013	**Good identification**					**23571013**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23571020	**Very good identification**					**23571020**
Actinomyces naeslundii		98.53	0.60	MNS 88		
Clostridium perfringens		1.47	0.07			
23571021	**Very good identification**					**23571021**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23571022	**Very good identification**					**23571022**
Actinomyces naeslundii		97.74	0.60	MNS 88		
Actinomyces viscosus		1.58	0.05			
Clostridium perfringens		0.69	0.01			
23571023	**Very good identification**					**23571023**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23571030	**Good identification**					**23571030**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23571031	**Good identification**					**23571031**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23571032	**Good identification**					**23571032**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23571033	**Good identification**					**23571033**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23571040	**Good identification**					**23571040**
Actinomyces naeslundii		96.20	0.47	MNS 88		
Actinomyces israelii		3.80	0.14			
23571041	**Good identification**					**23571041**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23571042	**Low discrimination**					**23571042**
Actinomyces naeslundii		86.91	0.47	MNS 88		
Actinomyces viscosus		7.94	0.05			
Actinomyces israelii		5.15	0.17			
23571043	**Good identification**					**23571043**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23571060	**Good identification**					**23571060**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23571061	**Good identification**					**23571061**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23571062	**Good identification**					**23571062**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23571063	**Good identification**					**23571063**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23571100	**Good identification**					**23571100**
Actinomyces naeslundii		94.85	0.68	MNS 88		
Actinomyces israelii		2.39	0.31			
Propionibacterium propionicum		1.06	0.24			
23571101	**Very good identification**					**23571101**
Actinomyces naeslundii		99.78	0.71	MNS 88		
Actinomyces viscosus		0.22	0.01			
23571102	**Low discrimination**					**23571102**
Actinomyces naeslundii		48.74	0.68	MNS 88		
Actinomyces viscosus		48.60	0.44			
Actinomyces israelii		1.84	0.34			
23571103	**Low discrimination**					**23571103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		81.86	0.71	MNS 88		
Actinomyces viscosus		18.14	0.36			
23571110	**Acceptable identification**					**23571110**
Actinomyces naeslundii		93.33	0.46	MNS 88		
Actinomyces israelii		6.67	0.17	MLZ 5		
23571111	**Good identification**					**23571111**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23571112	**Low discrimination**					**23571112**
Actinomyces naeslundii		77.01	0.46	MNS 88		
Actinomyces viscosus		14.74	0.10	MLZ 5		
Actinomyces israelii		8.26	0.20			
23571113	**Good identification**					**23571113**
Actinomyces naeslundii		95.92	0.49	MNS 88		
Actinomyces viscosus		4.08	0.01	MLZ 5		
23571120	**Very good identification**					**23571120**
					GRA	
Actinomyces naeslundii		95.85	0.68	MNS 88	99	
Clostridium perfringens		3.69	0.21		99	
Prevotella melaninogenica		0.46	-.01		1	
23571121	**Very good identification**					**23571121**
Actinomyces naeslundii		100.0	0.71	MNS 88		
23571122	**Very good identification**					**23571122**
Actinomyces naeslundii		96.42	0.68	MNS 88		
Clostridium perfringens		1.75	0.16			
Actinomyces viscosus		0.97	0.10			
23571123	**Very good identification**					**23571123**
Actinomyces naeslundii		99.78	0.71	MNS 88		
Actinomyces viscosus		0.22	0.01			
23571130	**Good identification**					**23571130**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23571131	**Good identification**					**23571131**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23571132	**Good identification**					**23571132**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23571133	**Good identification**					**23571133**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23571140	**Low discrimination**					**23571140**
Actinomyces naeslundii		89.40	0.55	MNS 88		
Propionibacterium propionicum		5.64	0.24			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		4.96	0.24			
23571141	**Very good identification**					**23571141**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23571142	**Low discrimination**					**23571142**
Actinomyces naeslundii		80.98	0.55	MNS 88		
Propionibacterium propionicum		7.66	0.27			
Actinomyces israelii		6.74	0.27			
23571143	**Very good identification**					**23571143**
Actinomyces naeslundii		98.75	0.58	MNS 88		
Actinomyces viscosus		1.25	0.01			
23571151	**Good identification**					**23571151**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23571153	**Good identification**					**23571153**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23571160	**Good identification**					**23571160**
Actinomyces naeslundii		96.01	0.55	MNS 88		
Clostridium perfringens		3.99	0.09			
23571161	**Very good identification**					**23571161**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23571162	**Good identification**					**23571162**
Actinomyces naeslundii		98.08	0.55	MNS 88		
Clostridium perfringens		1.92	0.03			
23571163	**Very good identification**					**23571163**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23571171	**Good identification**					**23571171**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23571173	**Good identification**					**23571173**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23571301	**Good identification**					**23571301**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23571303	**Low discrimination**					**23571303**
Actinomyces naeslundii		81.86	0.37	MNS 88		
Actinomyces viscosus		18.14	0.01	RHA 1		
23571321	**Good identification**					**23571321**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23571323	**Good identification**					**23571323**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.37	MNS 88		
23571501	**Good identification**					**23571501**
	Actinomyces naeslundii	100.0	0.37	MNS 88		
23571503	**Low discrimination**					**23571503**
	Actinomyces naeslundii	81.86	0.37	MNS 88		
	Actinomyces viscosus	18.14	0.01	ARA 1		
23571521	**Good identification**					**23571521**
	Actinomyces naeslundii	100.0	0.37	MNS 88		
23571523	**Good identification**					**23571523**
	Actinomyces naeslundii	100.0	0.37	MNS 88		
23572000	**Low discrimination**					**23572000**
	Actinomyces naeslundii	71.86	0.41	MNS 88		
	Propionibacterium propionicum	28.14	0.24	GAL 96		
23572001	**Good identification**					**23572001**
	Actinomyces naeslundii	100.0	0.44	MNS 88		
23572002	**Low discrimination**					**23572002**
	Actinomyces naeslundii	52.29	0.41	MNS 88		
	Propionibacterium propionicum	30.71	0.27	GAL 96		
	Actinomyces viscosus	17.01	0.09			
23572003	**Acceptable identification**					**23572003**
	Actinomyces naeslundii	93.26	0.44	MNS 88		
	Actinomyces viscosus	6.74	0.01	GAL 96		
23572020	**Good identification**					**23572020**
	Actinomyces naeslundii	100.0	0.41	MNS 88		
23572021	**Good identification**					**23572021**
	Actinomyces naeslundii	100.0	0.44	MNS 88		
23572022	**Good identification**					**23572022**
	Actinomyces naeslundii	100.0	0.41	MNS 88		
23572023	**Good identification**					**23572023**
	Actinomyces naeslundii	100.0	0.44	MNS 88		
23572100	**Good identification**					**23572100**
	Propionibacterium propionicum	93.16	0.58	CEL 1		
	Actinomyces naeslundii	6.84	0.49			
23572101	**Low discrimination**					**23572101**
	Propionibacterium propionicum	50.22	0.42	CEL 1		
	Actinomyces naeslundii	49.78	0.52	URE 10		
23572102	**Good identification**					**23572102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		94.44	0.61	CEL 1		
Actinomyces naeslundii		4.62	0.49			
Actinomyces viscosus		0.94	0.13			
23572103	**Low discrimination**					**23572103**
Propionibacterium propionicum		59.15	0.45	CEL 1		
Actinomyces naeslundii		39.09	0.52	URE 10		
Actinomyces viscosus		1.77	0.05			
23572120	**Low discrimination**					**23572120**
Actinomyces naeslundii		87.91	0.49	MNS 88		
Propionibacterium propionicum		12.09	0.24	GAL 96		
23572121	**Good identification**					**23572121**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23572122	**Low discrimination**					**23572122**
Actinomyces naeslundii		82.90	0.49	MNS 88		
Propionibacterium propionicum		17.11	0.27	GAL 96		
23572123	**Good identification**					**23572123**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23572140	**Very good identification**					**23572140**
Propionibacterium propionicum		98.72	0.58	CEL 1		
Actinomyces naeslundii		1.28	0.36			
23572141	**Low discrimination**					**23572141**
Propionibacterium propionicum		85.11	0.42	CEL 1		
Actinomyces naeslundii		14.89	0.39	URE 10		
23572142	**Very good identification**					**23572142**
Propionibacterium propionicum		99.14	0.61	CEL 1		
Actinomyces naeslundii		0.86	0.36			
23572143	**Low discrimination**					**23572143**
Propionibacterium propionicum		89.56	0.45	CEL 1		
Actinomyces naeslundii		10.44	0.39	URE 10		
23572160	**Low discrimination**					**23572160**
Actinomyces naeslundii		56.20	0.36	MNS 88		
Propionibacterium propionicum		43.80	0.24	GAL 96		
23572161	**Good identification**					**23572161**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23572163	**Good identification**					**23572163**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23573000	**Good identification**					**23573000**
Actinomyces naeslundii		93.58	0.65	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		4.89	0.34			
Propionibacterium propionicum		1.53	0.24			
23573001	**Very good identification**					**23573001**
Actinomyces naeslundii		100.0	0.68	MNS 88		
23573002	**Low discrimination**					**23573002**
Actinomyces naeslundii		89.99	0.65	MNS 88		
Actinomyces israelii		7.06	0.37			
Propionibacterium propionicum		2.20	0.27			
23573003	**Very good identification**					**23573003**
Actinomyces naeslundii		100.0	0.68	MNS 88		
23573010	**Low discrimination**					**23573010**
Actinomyces naeslundii		87.07	0.43	MNS 88		
Actinomyces israelii		12.93	0.20	MLZ 5		
23573011	**Good identification**					**23573011**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23573012	**Low discrimination**					**23573012**
Actinomyces naeslundii		81.79	0.43	MNS 88		
Actinomyces israelii		18.21	0.23	MLZ 5		
23573013	**Good identification**					**23573013**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23573020	**Very good identification**					**23573020**
Actinomyces naeslundii		99.67	0.65	MNS 88		
Actinomyces israelii		0.33	0.13			
23573021	**Very good identification**					**23573021**
Actinomyces naeslundii		100.0	0.68	MNS 88		
23573022	**Very good identification**					**23573022**
Actinomyces naeslundii		99.50	0.65	MNS 88		
Actinomyces israelii		0.50	0.16			
23573023	**Very good identification**					**23573023**
Actinomyces naeslundii		100.0	0.68	MNS 88		
23573030	**Good identification**					**23573030**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23573031	**Good identification**					**23573031**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23573032	**Good identification**					**23573032**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23573033	**Good identification**					**23573033**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MNS 88		
23573040	**Low discrimination**					**23573040**
Actinomyces naeslundii		82.80	0.52	MNS 88		
Actinomyces israelii		9.54	0.27			
Propionibacterium propionicum		7.66	0.24			
23573041	**Good identification**					**23573041**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23573042	**Low discrimination**					**23573042**
Actinomyces naeslundii		76.25	0.52	MNS 88		
Actinomyces israelii		13.18	0.30			
Propionibacterium propionicum		10.57	0.27			
23573043	**Good identification**					**23573043**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23573060	**Good identification**					**23573060**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23573061	**Good identification**					**23573061**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23573062	**Good identification**					**23573062**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23573063	**Good identification**					**23573063**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23573100	**Low discrimination**					**23573100**
Actinomyces naeslundii		60.94	0.73	MNS 88		
Propionibacterium propionicum		34.58	0.58			
Actinomyces israelii		4.48	0.44			
23573101	**Very good identification**					**23573101**
Actinomyces naeslundii		95.97	0.76	MNS 88		
Propionibacterium propionicum		4.03	0.42			
23573102	**Low discrimination**					**23573102**
Actinomyces naeslundii		50.85	0.73	MNS 88		
Propionibacterium propionicum		43.28	0.61			
Actinomyces israelii		5.61	0.47			
23573103	**Good identification**					**23573103**
Actinomyces naeslundii		93.97	0.76	MNS 88		
Propionibacterium propionicum		5.93	0.45			
Actinomyces viscosus		0.11	0.01			
23573110	**Low discrimination**					**23573110**
Actinomyces naeslundii		75.90	0.51	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		15.84	0.30	MLZ 5		
Propionibacterium propionicum		8.27	0.24			
23573111	**Good identification**					**23573111**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23573112	**Low discrimination**					**23573112**
Actinomyces naeslundii		67.74	0.51	MNS 88		
Actinomyces israelii		21.20	0.33	MLZ 5		
Propionibacterium propionicum		11.07	0.27			
23573113	**Good identification**					**23573113**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23573120	**Very good identification**					**23573120**
Actinomyces naeslundii		98.97	0.73	MNS 88		
Propionibacterium propionicum		0.57	0.24			
Actinomyces israelii		0.46	0.24			
23573121	**Very good identification**					**23573121**
Actinomyces naeslundii		100.0	0.76	MNS 88		
23573122	**Very good identification**					**23573122**
Actinomyces naeslundii		98.46	0.73	MNS 88		
Propionibacterium propionicum		0.85	0.27			
Actinomyces israelii		0.69	0.27			
23573123	**Very good identification**					**23573123**
Actinomyces naeslundii		100.0	0.76	MNS 88		
23573130	**Good identification**					**23573130**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23573131	**Good identification**					**23573131**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23573132	**Good identification**					**23573132**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23573133	**Good identification**					**23573133**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23573140	**Low discrimination**					**23573140**
Propionibacterium propionicum		73.46	0.58	CEL 1		
Actinomyces naeslundii		22.84	0.60			
Actinomyces israelii		3.70	0.37			
23573141	**Low discrimination**					**23573141**
Actinomyces naeslundii		80.76	0.63	MNS 88		
Propionibacterium propionicum		19.24	0.42			
23573142	**Low discrimination**					**23573142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		79.51	0.61	CEL 1		
Actinomyces naeslundii		16.48	0.60			
Actinomyces israelii		4.00	0.40			
23573143	**Low discrimination**					**23573143**
Actinomyces naeslundii		73.68	0.63	MNS 88		
Propionibacterium propionicum		26.32	0.45			
23573150	**Low discrimination**					**23573150**
Actinomyces naeslundii		48.15	0.38	MNS 88		
Propionibacterium propionicum		29.71	0.24	MLZ 5		
Actinomyces israelii		22.14	0.23			
23573151	**Good identification**					**23573151**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23573152	**Low discrimination**					**23573152**
Actinomyces naeslundii		38.24	0.38	MNS 88		
Propionibacterium propionicum		35.40	0.27	MLZ 5		
Actinomyces israelii		26.37	0.26			
23573153	**Good identification**					**23573153**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23573160	**Very good identification**					**23573160**
Actinomyces naeslundii		95.89	0.60	MNS 88		
Propionibacterium propionicum		3.12	0.24			
Actinomyces israelii		0.99	0.17			
23573161	**Very good identification**					**23573161**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23573162	**Good identification**					**23573162**
Actinomyces naeslundii		93.97	0.60	MNS 88		
Propionibacterium propionicum		4.58	0.27			
Actinomyces israelii		1.46	0.20			
23573163	**Very good identification**					**23573163**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23573170	**Good identification**					**23573170**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23573171	**Good identification**					**23573171**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23573172	**Good identification**					**23573172**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23573173	**Good identification**					**23573173**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23573300	**Low discrimination**					**23573300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		47.28	0.39	MNS 88		
Propionibacterium propionicum		26.83	0.24	RHA 1		
Actinomyces israelii		25.89	0.25			
23573301	**Good identification**					**23573301**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573302	**Low discrimination**					**23573302**
Actinomyces naeslundii		37.42	0.39	MNS 88		
Propionibacterium propionicum		31.85	0.27	RHA 1		
Actinomyces israelii		30.74	0.28			
23573303	**Good identification**					**23573303**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573320	**Good identification**					**23573320**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23573321	**Good identification**					**23573321**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573322	**Good identification**					**23573322**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23573323	**Good identification**					**23573323**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573402	**Low discrimination**					**23573402**
Actinomyces israelii		86.40	0.35	SAL 92		
Actinomyces naeslundii		13.60	0.31	XYL 92		
23573500	**Low discrimination**					**23573500**
Actinomyces israelii		79.16	0.43	SAL 92		
Actinomyces naeslundii		13.30	0.39	XYL 92		
Propionibacterium propionicum		7.55	0.24			
23573501	**Good identification**					**23573501**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573502	**Low discrimination**					**23573502**
Actinomyces israelii		81.78	0.46	SAL 92		
Actinomyces naeslundii		9.16	0.39	XYL 92		
Propionibacterium propionicum		7.80	0.27			
23573503	**Good identification**					**23573503**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573520	**Low discrimination**					**23573520**
Actinomyces naeslundii		72.47	0.39	MNS 88		
Actinomyces israelii		27.53	0.22	ARA 1		
23573521	**Good identification**					**23573521**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573522	**Low discrimination**					**23573522**
Actinomyces naeslundii		63.70	0.39	MNS 88		
Actinomyces israelii		36.30	0.25	ARA 1		
23573523	**Good identification**					**23573523**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23573540	**Low discrimination**					**23573540**
Actinomyces israelii		75.68	0.36	SAL 92		
Propionibacterium propionicum		18.55	0.24	XYL 92		
Actinomyces naeslundii		5.77	0.26			
23573542	**Low discrimination**					**23573542**
Actinomyces israelii		77.16	0.39	SAL 92		
Propionibacterium propionicum		18.92	0.27	XYL 92		
Actinomyces naeslundii		3.92	0.26			
23575000	**Good identification**					**23575000**
Actinomyces israelii		95.33	0.39	SAL 92		
Actinomyces naeslundii		4.68	0.26			
23575002	**Acceptable identification**					**23575002**
Actinomyces israelii		92.18	0.42	SAL 92		
Actinomyces viscosus		4.81	0.05			
Actinomyces naeslundii		3.01	0.26			
23575042	**Good identification**					**23575042**
Actinomyces israelii		100.0	0.35	SAL 92		
23575100	**Good identification**					**23575100**
Actinomyces israelii		96.63	0.49	SAL 92		
Actinomyces naeslundii		3.37	0.34			
23575101	**Low discrimination**					**23575101**
Actinomyces naeslundii		83.82	0.37	MNS 88		
Actinomyces israelii		16.18	0.15	XYL 1		
23575102	**Acceptable identification**					**23575102**
Actinomyces israelii		94.87	0.52	SAL 92		
Actinomyces naeslundii		2.21	0.34			
Actinomyces viscosus		2.20	0.10			
23575103	**Low discrimination**					**23575103**
Actinomyces naeslundii		66.18	0.37	MNS 88		
Actinomyces israelii		19.16	0.18	XYL 1		
Actinomyces viscosus		14.66	0.01			
23575110	**Good identification**					**23575110**
Actinomyces israelii		100.0	0.35	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23575112	**Acceptable identification**					**23575112**
Actinomyces israelii		94.14	0.38	SAL 92		
Bifidobacterium adolescentis		5.86	0.11	MLZ 13		
23575121	**Good identification**					**23575121**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23575123	**Good identification**					**23575123**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23575140	**Good identification**					**23575140**
Actinomyces israelii		100.0	0.42	SAL 92		
23575142	**Good identification**					**23575142**
Actinomyces israelii		100.0	0.45	SAL 92		
23575400	**Good identification**					**23575400**
Actinomyces israelii		100.0	0.38	SAL 92		
23575402	**Good identification**					**23575402**
Actinomyces israelii		100.0	0.41	SAL 92		
23575500	**Good identification**					**23575500**
Actinomyces israelii		100.0	0.48	SAL 92		
23575502	**Acceptable identification**					**23575502**
Actinomyces israelii		94.15	0.51	SAL 92		
Bifidobacterium adolescentis		5.86	0.24			
23575512	**Low discrimination**					**23575512**
Actinomyces israelii		66.28	0.37	SAL 92		
Bifidobacterium adolescentis		33.72	0.26	MLZ 13		
23575540	**Good identification**					**23575540**
Actinomyces israelii		100.0	0.41	SAL 92		
23575542	**Good identification**					**23575542**
Actinomyces israelii		98.06	0.44	SAL 92		
Bifidobacterium adolescentis		1.94	0.09			
23577000	**Good identification**					**23577000**
Actinomyces israelii		98.35	0.52	SAL 92		
Actinomyces naeslundii		1.65	0.31			
23577002	**Good identification**					**23577002**
Actinomyces israelii		98.89	0.55	SAL 92		
Actinomyces naeslundii		1.11	0.31			
23577010	**Good identification**					**23577010**
Actinomyces israelii		100.0	0.38	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23577012	**Good identification**					**23577012**
Actinomyces israelii		100.0	0.41	SAL 92		
23577040	**Good identification**					**23577040**
Actinomyces israelii		100.0	0.45	SAL 92		
23577042	**Good identification**					**23577042**
Actinomyces israelii		100.0	0.48	SAL 92		
23577100	**Very good identification**					**23577100**
Actinomyces israelii		98.16	0.62	SAL 92		
Actinomyces naeslundii		1.17	0.39			
Propionibacterium propionicum		0.67	0.24			
23577101	**Low discrimination**					**23577101**
Actinomyces naeslundii		63.96	0.42	MNS 88		
Actinomyces israelii		36.04	0.28	XYL 1		
23577102	**Very good identification**					**23577102**
Actinomyces israelii		98.44	0.65	SAL 92		
Actinomyces naeslundii		0.78	0.39			
Propionibacterium propionicum		0.67	0.27			
23577103	**Low discrimination**					**23577103**
Actinomyces naeslundii		54.20	0.42	MNS 88		
Actinomyces israelii		45.80	0.31	XYL 1		
23577110	**Good identification**					**23577110**
Actinomyces israelii		100.0	0.48	SAL 92		
23577112	**Good identification**					**23577112**
Actinomyces israelii		99.11	0.51	SAL 92		
Bifidobacterium adolescentis		0.89	0.10	MLZ 13		
23577120	**Low discrimination**					**23577120**
Actinomyces israelii		84.23	0.42	SAL 92		
Actinomyces naeslundii		15.77	0.39	NAG 6		
23577121	**Good identification**					**23577121**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23577122	**Low discrimination**					**23577122**
Actinomyces israelii		88.90	0.45	SAL 92		
Actinomyces naeslundii		11.10	0.39	NAG 6		
23577123	**Good identification**					**23577123**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23577140	**Good identification**					**23577140**
Actinomyces israelii		97.77	0.55	SAL 92		
Propionibacterium propionicum		1.71	0.24			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.53	0.26			
23577142	**Very good identification**					**23577142**
Actinomyces israelii		97.94	0.58	SAL 92		
Propionibacterium propionicum		1.71	0.27			
Actinomyces naeslundii		0.35	0.26			
23577150	**Good identification**					**23577150**
Actinomyces israelii		100.0	0.41	SAL 92		
23577152	**Good identification**					**23577152**
Actinomyces israelii		100.0	0.44	SAL 92		
23577162	**Acceptable identification**					**23577162**
Actinomyces israelii		94.64	0.38	SAL 92		
Actinomyces naeslundii		5.36	0.26	NAG 6		
23577202	**Good identification**					**23577202**
Actinomyces israelii		100.0	0.36	SAL 92		
23577300	**Good identification**					**23577300**
Actinomyces israelii		100.0	0.43	SAL 92		
23577302	**Good identification**					**23577302**
Actinomyces israelii		100.0	0.46	SAL 92		
23577340	**Good identification**					**23577340**
Actinomyces israelii		100.0	0.36	SAL 92		
23577342	**Good identification**					**23577342**
Actinomyces israelii		100.0	0.39	SAL 92		
23577400	**Good identification**					**23577400**
Actinomyces israelii		100.0	0.51	SAL 92		
23577402	**Good identification**					**23577402**
Actinomyces israelii		100.0	0.54	SAL 92		
23577410	**Good identification**					**23577410**
Actinomyces israelii		100.0	0.36	SAL 92		
23577412	**Good identification**					**23577412**
Actinomyces israelii		100.0	0.39	SAL 92		
23577440	**Good identification**					**23577440**
Actinomyces israelii		100.0	0.44	SAL 92		
23577442	**Good identification**					**23577442**
Actinomyces israelii		100.0	0.47	SAL 92		
23577500	**Very good identification**					**23577500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.61	SAL 92		
23577502	**Very good identification**					**23577502**
Actinomyces israelii		99.11	0.64	SAL 92		
Bifidobacterium adolescentis		0.89	0.23			
23577510	**Good identification**					**23577510**
Actinomyces israelii		100.0	0.47	SAL 92		
23577512	**Acceptable identification**					**23577512**
Actinomyces israelii		93.16	0.50	SAL 92		
Bifidobacterium adolescentis		6.84	0.24	MLZ 13		
23577520	**Good identification**					**23577520**
Actinomyces israelii		100.0	0.40	SAL 92		
23577522	**Good identification**					**23577522**
Actinomyces israelii		100.0	0.43	SAL 92		
23577540	**Good identification**					**23577540**
Actinomyces israelii		100.0	0.54	SAL 92		
23577542	**Very good identification**					**23577542**
Actinomyces israelii		99.72	0.57	SAL 92		
Bifidobacterium adolescentis		0.29	0.07			
23577550	**Good identification**					**23577550**
Actinomyces israelii		100.0	0.40	SAL 92		
23577552	**Good identification**					**23577552**
Actinomyces israelii		97.72	0.43	SAL 92		
Bifidobacterium adolescentis		2.28	0.09	MLZ 13		
23577562	**Good identification**					**23577562**
Actinomyces israelii		100.0	0.36	SAL 92		
23577700	**Good identification**					**23577700**
Actinomyces israelii		100.0	0.42	SAL 92		
23577702	**Good identification**					**23577702**
Actinomyces israelii		100.0	0.45	SAL 92		
23577742	**Good identification**					**23577742**
Actinomyces israelii		100.0	0.38	SAL 92		
23601000	**Very good identification**					**23601000**
					GRA	
Eubacterium tortuosum		99.87	0.56	ESL 99	99	
Actinomyces odontolyticus		0.08	0.14		99	
Fusobacterium mortiferum		0.06	0.17		1	
23601002	**Good identification**					**23601002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		99.50	0.43	ESL 99		
Actinomyces odontolyticus		0.26	0.10			
Actinomyces viscosus		0.24	0.03			
23601020	**Good identification**					**23601020**
Eubacterium tortuosum		100.0	0.43	ESL 99		
23603101	**Good identification**					**23603101**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23603103	**Good identification**					**23603103**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23603121	**Good identification**					**23603121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23603123	**Good identification**					**23603123**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23610000	**Low discrimination**					**23610000**
					GRA	
Fusobacterium mortiferum		74.29	0.49	NIT 1	1	
Actinomyces odontolyticus		25.72	0.35		99	
23610100	**Acceptable identification**					**23610100**
					GRA	
Fusobacterium mortiferum		94.89	0.55	NIT 1	1	
Actinomyces odontolyticus		4.45	0.27		99	
Actinomyces naeslundii		0.42	0.26		99	
23610200	**Good identification**					**23610200**
					GRA	
Actinomyces odontolyticus		98.52	0.40	MLT 99	99	
Fusobacterium mortiferum		1.48	0.14	RAF 3	1	
23610202	**Good identification**					**23610202**
Actinomyces odontolyticus		100.0	0.36	MLT 99		
23611000	**Low discrimination**					**23611000**
					GRA	
Fusobacterium mortiferum		71.45	0.52	NIT 1	1	
Eubacterium tortuosum		12.62	0.22		99	
Actinomyces odontolyticus		10.99	0.32		99	
23611001	**Low discrimination**					**23611001**
					GRA	
Actinomyces naeslundii		89.56	0.45	MNS 88	99	
Fusobacterium mortiferum		10.44	0.17	MLT 94	1	
23611003	**Low discrimination**					**23611003**
Actinomyces naeslundii		64.88	0.45	MNS 88		
Actinomyces viscosus		35.12	0.16	MLT 94		
23611020	**Low discrimination**					**23611020**
Actinomyces naeslundii		46.56	0.42	MNS 88		
Eubacterium tortuosum		25.12	0.09	MLT 94		
Actinomyces odontolyticus		20.18	0.19			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23611021	**Good identification**					**23611021**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23611022	**Low discrimination**					**23611022**
Actinomyces naeslundii		73.48	0.42	MNS 88		
Actinomyces odontolyticus		19.53	0.15	MLT 94		
Eubacterium tortuosum		7.00	-.04			
23611023	**Good identification**					**23611023**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23611100	**Good identification**					**23611100**
					GRA	
Fusobacterium mortiferum		90.97	0.58	NIT 1	1	
Actinomyces naeslundii		6.41	0.50		99	
Actinomyces odontolyticus		1.89	0.24		99	
23611101	**Acceptable identification**					**23611101**
					GRA	
Actinomyces naeslundii		91.28	0.53	MNS 88	99	
Fusobacterium mortiferum		8.72	0.24	MLT 94	1	
23611102	**Low discrimination**					**23611102**
					GRA	
Fusobacterium mortiferum		35.90	0.42	NIT 1	1	
Actinomyces viscosus		34.64	0.29	BGL 10	99	
Actinomyces naeslundii		22.77	0.50		99	
23611103	**Low discrimination**					**23611103**
Actinomyces naeslundii		74.73	0.53	MNS 88		
Actinomyces viscosus		25.27	0.21	MLT 94		
23611120	**Low discrimination**					**23611120**
					GRA	
Actinomyces naeslundii		81.35	0.50	MNS 88	99	
Fusobacterium mortiferum		11.66	0.24	MLT 94	1	
Actinomyces odontolyticus		3.91	0.10		99	
23611121	**Good identification**					**23611121**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23611122	**Acceptable identification**					**23611122**
Actinomyces naeslundii		93.70	0.50	MNS 88		
Lactobacillus fermentum		3.54	0.15	MLT 94		
Actinomyces odontolyticus		2.76	0.06			
23611123	**Good identification**					**23611123**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23611140	**Low discrimination**					**23611140**
					GRA	
Actinomyces naeslundii		55.18	0.37	MNS 88	99	
Fusobacterium mortiferum		44.82	0.24	MLT 94	1	
23611141	**Good identification**					**23611141**
Actinomyces naeslundii		100.0	0.40	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23611142	**Good identification**					**23611142**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23611143	**Good identification**					**23611143**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23611160	**Good identification**					**23611160**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23611161	**Good identification**					**23611161**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23611162	**Good identification**					**23611162**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23611163	**Good identification**					**23611163**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23611200	**Good identification**					**23611200**
					GRA	
Actinomyces odontolyticus		96.73	0.37	MLT 99	99	
Fusobacterium mortiferum		3.27	0.17	RAF 3	1	
23613000	**Acceptable identification**					**23613000**
					GRA	
Actinomyces naeslundii		91.74	0.47	MNS 88	99	
Fusobacterium mortiferum		8.26	0.17	MLT 94	1	
23613001	**Good identification**					**23613001**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23613002	**Good identification**					**23613002**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23613003	**Good identification**					**23613003**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23613020	**Good identification**					**23613020**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23613021	**Good identification**					**23613021**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23613022	**Good identification**					**23613022**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23613023	**Good identification**					**23613023**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23613041	**Good identification**					**23613041**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23613043	**Good identification**					**23613043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	MNS 88		
23613061	**Good identification**					**23613061**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23613063	**Good identification**					**23613063**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23613100	**Low discrimination**					**23613100**
					GRA	
Actinomyces naeslundii		86.69	0.55	MNS 88	99	
Propionibacterium propionicum		6.91	0.26	MLT 94	99	
Fusobacterium mortiferum		6.40	0.24		1	
23613101	**Very good identification**					**23613101**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23613102	**Low discrimination**					**23613102**
Actinomyces naeslundii		89.33	0.55	MNS 88		
Propionibacterium propionicum		10.67	0.29	MLT 94		
23613103	**Very good identification**					**23613103**
Actinomyces naeslundii		99.12	0.58	MNS 88		
Propionibacterium propionicum		0.88	0.12	MLT 94		
23613111	**Good identification**					**23613111**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23613113	**Good identification**					**23613113**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23613120	**Good identification**					**23613120**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23613121	**Very good identification**					**23613121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23613122	**Good identification**					**23613122**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23613123	**Very good identification**					**23613123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23613131	**Good identification**					**23613131**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23613133	**Good identification**					**23613133**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23613140	**Low discrimination**					**23613140**
Actinomyces naeslundii		68.90	0.42	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		31.10	0.26	MLT 94		
23613141	**Good identification**					**23613141**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23613142	**Low discrimination**					**23613142**
Actinomyces naeslundii		59.63	0.42	MNS 88		
Propionibacterium propionicum		40.37	0.29	MLT 94		
23613143	**Good identification**					**23613143**
Actinomyces naeslundii		95.23	0.45	MNS 88		
Propionibacterium propionicum		4.78	0.12	MLT 94		
23613160	**Good identification**					**23613160**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23613161	**Good identification**					**23613161**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23613162	**Good identification**					**23613162**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23613163	**Good identification**					**23613163**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23614100	**Low discrimination**					**23614100**
					GRA	
Fusobacterium mortiferum		76.54	0.36	NIT 1	1	
Actinomyces odontolyticus		23.46	0.21	XYL 7	99	
23615100	**Low discrimination**					**23615100**
					GRA	
Fusobacterium mortiferum		83.44	0.39	NIT 1	1	
Actinomyces odontolyticus		11.37	0.18	XYL 7	99	
Lactobacillus fermentum		5.20	0.20		99	
23621101	**Good identification**					**23621101**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23621103	**Good identification**					**23621103**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23621121	**Good identification**					**23621121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23621123	**Good identification**					**23621123**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23623100	**Good identification**					**23623100**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23623101	**Good identification**					**23623101**
Actinomyces naeslundii		100.0	0.41	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23623102	**Good identification**					**23623102**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23623103	**Good identification**					**23623103**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23623120	**Good identification**					**23623120**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23623121	**Good identification**					**23623121**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23623122	**Good identification**					**23623122**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23623123	**Good identification**					**23623123**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23630000	**Good identification**					**23630000**
					GRA	
Fusobacterium mortiferum		97.71	0.40	NIT 1	1	
Actinomyces naeslundii		2.29	0.23		99	
23630100	**Good identification**					**23630100**
					GRA	
Fusobacterium mortiferum		97.22	0.46	NIT 1	1	
Actinomyces naeslundii		2.78	0.31		99	
23631000	**Low discrimination**					**23631000**
					GRA	
Fusobacterium mortiferum		72.71	0.43	NIT 1	1	
Actinomyces naeslundii		27.30	0.47		99	
23631001	**Good identification**					**23631001**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23631002	**Low discrimination**					**23631002**
					GRA	
Actinomyces naeslundii		71.92	0.47	MNS 88	99	
Fusobacterium mortiferum		21.29	0.26	MLT 94	1	
Actinomyces viscosus		6.79	0.05		99	
23631003	**Good identification**					**23631003**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23631020	**Good identification**					**23631020**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23631021	**Good identification**					**23631021**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23631022	**Good identification**					**23631022**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23631023	**Good identification**					**23631023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MNS 88		
23631041	**Good identification**					**23631041**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23631043	**Good identification**					**23631043**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23631061	**Good identification**					**23631061**
Actinomyces naeslundii		100.0	0.37	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23631063	**Good identification**					**23631063**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23631100	**Low discrimination**					**23631100**
					GRA	
Fusobacterium mortiferum		68.59	0.49	NIT 1	1	
Actinomyces naeslundii		31.41	0.55		99	
23631101	**Very good identification**					**23631101**
					GRA	
Actinomyces naeslundii		98.55	0.58	MNS 88	99	
Fusobacterium mortiferum		1.45	0.15	MLT 94	1	
23631102	**Low discrimination**					**23631102**
					GRA	
Actinomyces naeslundii		76.83	0.55	MNS 88	99	
Fusobacterium mortiferum		18.64	0.33	MLT 94	1	
Actinomyces viscosus		4.53	0.10		99	
23631103	**Very good identification**					**23631103**
Actinomyces naeslundii		98.71	0.58	MNS 88		
Actinomyces viscosus		1.29	0.01	MLT 94		
23631111	**Good identification**					**23631111**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23631113	**Good identification**					**23631113**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23631120	**Good identification**					**23631120**
					GRA	
Actinomyces naeslundii		96.55	0.55	MNS 88	99	
Fusobacterium mortiferum		2.13	0.15	MLT 94	1	
Clostridium perfringens		1.32	0.01		99	
23631121	**Very good identification**					**23631121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23631122	**Good identification**					**23631122**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23631123	**Very good identification**					**23631123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23631131	**Good identification**					**23631131**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23631133	**Good identification**					**23631133**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23631140	**Low discrimination**					**23631140**
					GRA	
Actinomyces naeslundii		88.89	0.42	MNS 88	99	
Fusobacterium mortiferum		11.11	0.15	MLT 94	1	
23631141	**Good identification**					**23631141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	MNS 88		
23631142	**Good identification**					**23631142**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23631143	**Good identification**					**23631143**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23631160	**Good identification**					**23631160**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23631161	**Good identification**					**23631161**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23631162	**Good identification**					**23631162**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23631163	**Good identification**					**23631163**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23632100	**Low discrimination**					**23632100**
					GRA	
Actinomyces naeslundii		50.31	0.36	MNS 88	99	
Propionibacterium propionicum		40.54	0.24	MLT 94	99	
Fusobacterium mortiferum		9.15	0.12	GAL 96	1	
23632101	**Good identification**					**23632101**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23632103	**Good identification**					**23632103**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23632120	**Good identification**					**23632120**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23632121	**Good identification**					**23632121**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23632122	**Good identification**					**23632122**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23632123	**Good identification**					**23632123**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23633000	**Good identification**					**23633000**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23633001	**Good identification**					**23633001**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23633002	**Good identification**					**23633002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	MNS 88		
23633003	**Good identification**					**23633003**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23633020	**Good identification**					**23633020**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23633021	**Good identification**					**23633021**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23633022	**Good identification**					**23633022**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23633023	**Good identification**					**23633023**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23633040	**Good identification**					**23633040**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23633041	**Good identification**					**23633041**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23633042	**Good identification**					**23633042**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23633043	**Good identification**					**23633043**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23633060	**Good identification**					**23633060**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23633061	**Good identification**					**23633061**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23633062	**Good identification**					**23633062**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23633063	**Good identification**					**23633063**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23633100	**Very good identification**					**23633100**
					GRA	
Actinomyces naeslundii		95.70	0.60	MNS 88	99	
Propionibacterium propionicum		3.21	0.24	MLT 94	99	
Fusobacterium mortiferum		1.09	0.15		1	
23633101	**Very good identification**					**23633101**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23633102	**Very good identification**					**23633102**
Actinomyces naeslundii		95.21	0.60	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		4.80	0.27	MLT 94		
23633103	**Very good identification**					**23633103**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23633110	**Good identification**					**23633110**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23633111	**Good identification**					**23633111**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23633112	**Good identification**					**23633112**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23633113	**Good identification**					**23633113**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23633120	**Very good identification**					**23633120**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23633121	**Very good identification**					**23633121**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23633122	**Very good identification**					**23633122**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23633123	**Very good identification**					**23633123**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23633130	**Good identification**					**23633130**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23633131	**Good identification**					**23633131**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23633132	**Good identification**					**23633132**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23633133	**Good identification**					**23633133**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23633140	**Low discrimination**					**23633140**
Actinomyces naeslundii		84.02	0.47	MNS 88		
Propionibacterium propionicum		15.98	0.24	MLT 94		
23633141	**Good identification**					**23633141**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23633142	**Low discrimination**					**23633142**
Actinomyces naeslundii		77.80	0.47	MNS 88		
Propionibacterium propionicum		22.20	0.27	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23633143	**Good identification**					**23633143**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23633160	**Good identification**					**23633160**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23633161	**Good identification**					**23633161**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23633162	**Good identification**					**23633162**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23633163	**Good identification**					**23633163**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23641000	**Good identification**					**23641000**
Eubacterium tortuosum		100.0	0.40	ESL 99		
23643101	**Good identification**					**23643101**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23643103	**Good identification**					**23643103**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23643121	**Good identification**					**23643121**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23643123	**Good identification**					**23643123**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23650000	**Good identification**					**23650000**
Fusobacterium mortiferum		100.0	0.40	NIT 1		
23650100	**Good identification**					**23650100**
					GRA	
Fusobacterium mortiferum		98.28	0.47	NIT 1	1	
Actinomyces naeslundii		1.72	0.28		99	
23651000	**Low discrimination**					**23651000**
					GRA	
Fusobacterium mortiferum		77.60	0.43	NIT 1	1	
Actinomyces naeslundii		17.83	0.44		99	
Eubacterium tortuosum		4.57	0.06		99	
23651001	**Good identification**					**23651001**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23651002	**Low discrimination**					**23651002**
					GRA	
Actinomyces naeslundii		50.71	0.44	MNS 88	99	
Actinomyces viscosus		24.77	0.15	MLT 94	99	
Fusobacterium mortiferum		24.53	0.27		1	
23651003	**Acceptable identification**					**23651003**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		90.21	0.47	MNS 88		
Actinomyces viscosus		9.79	0.06	MLT 94		
23651020	**Good identification**					**23651020**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23651021	**Good identification**					**23651021**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23651022	**Good identification**					**23651022**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23651023	**Good identification**					**23651023**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23651100	**Low discrimination**					**23651100**
					GRA	
Fusobacterium mortiferum		78.11	0.50	NIT 1	1	
Actinomyces naeslundii		21.89	0.52		99	
23651101	**Good identification**					**23651101**
					GRA	
Actinomyces naeslundii		97.65	0.55	MNS 88	99	
Fusobacterium mortiferum		2.35	0.16	MLT 94	1	
23651102	**Low discrimination**					**23651102**
					GRA	
Actinomyces naeslundii		58.77	0.52	MNS 88	99	
Fusobacterium mortiferum		23.30	0.33	MLT 94	1	
Actinomyces viscosus		17.93	0.19		99	
23651103	**Acceptable identification**					**23651103**
Actinomyces naeslundii		93.65	0.55	MNS 88		
Actinomyces viscosus		6.35	0.11	MLT 94		
23651120	**Good identification**					**23651120**
					GRA	
Actinomyces naeslundii		96.52	0.52	MNS 88	99	
Fusobacterium mortiferum		3.48	0.16	MLT 94	1	
23651121	**Good identification**					**23651121**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23651122	**Good identification**					**23651122**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23651123	**Good identification**					**23651123**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23651140	**Low discrimination**					**23651140**
					GRA	
Actinomyces naeslundii		83.04	0.39	MNS 88	99	
Fusobacterium mortiferum		16.96	0.16	MLT 94	1	
23651141	**Good identification**					**23651141**
Actinomyces naeslundii		100.0	0.42	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23651142	**Good identification**					**23651142**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23651143	**Good identification**					**23651143**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23651160	**Good identification**					**23651160**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23651161	**Good identification**					**23651161**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23651162	**Good identification**					**23651162**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23651163	**Good identification**					**23651163**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23652101	**Good identification**					**23652101**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23652103	**Good identification**					**23652103**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23652121	**Good identification**					**23652121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23652123	**Good identification**					**23652123**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23653000	**Good identification**					**23653000**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23653001	**Good identification**					**23653001**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23653002	**Good identification**					**23653002**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23653003	**Good identification**					**23653003**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23653020	**Good identification**					**23653020**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23653021	**Good identification**					**23653021**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23653022	**Good identification**					**23653022**
Actinomyces naeslundii		100.0	0.49	MNS 88		

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
23653023	**Good identification**					**23653023**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23653040	**Good identification**					**23653040**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23653041	**Good identification**					**23653041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23653042	**Good identification**					**23653042**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23653043	**Good identification**					**23653043**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23653060	**Good identification**					**23653060**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23653061	**Good identification**					**23653061**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23653062	**Good identification**					**23653062**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23653063	**Good identification**					**23653063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23653100	**Very good identification**					**23653100**
					GRA	
Actinomyces naeslundii		96.34	0.57	MNS 88	99	
Actinomyces israelii		1.88	0.18	MLT 94	99	
Fusobacterium mortiferum		1.79	0.16		1	
23653101	**Very good identification**					**23653101**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23653102	**Very good identification**					**23653102**
Actinomyces naeslundii		97.16	0.57	MNS 88		
Actinomyces israelii		2.84	0.21	MLT 94		
23653103	**Very good identification**					**23653103**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23653111	**Good identification**					**23653111**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23653113	**Good identification**					**23653113**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23653120	**Very good identification**					**23653120**
Actinomyces naeslundii		100.0	0.57	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23653121	**Very good identification**					**23653121**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23653122	**Very good identification**					**23653122**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23653123	**Very good identification**					**23653123**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23653131	**Good identification**					**23653131**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23653133	**Good identification**					**23653133**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23653140	**Good identification**					**23653140**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23653141	**Good identification**					**23653141**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23653142	**Acceptable identification**					**23653142**
Actinomyces naeslundii		93.96	0.44	MNS 88		
Actinomyces israelii		6.05	0.14	MLT 94		
23653143	**Good identification**					**23653143**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23653160	**Good identification**					**23653160**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23653161	**Good identification**					**23653161**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23653162	**Good identification**					**23653162**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23653163	**Good identification**					**23653163**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23657100	**Good identification**					**23657100**
Actinomyces israelii		95.68	0.37	MLT 99		
Actinomyces naeslundii		4.32	0.23			
23657102	**Good identification**					**23657102**
Actinomyces israelii		97.08	0.40	MLT 99		
Actinomyces naeslundii		2.92	0.23			
23657500	**Good identification**					**23657500**
Actinomyces israelii		100.0	0.35	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23657502	**Good identification**					**23657502**
Actinomyces israelii		100.0	0.38	MLT 99		
23661101	**Good identification**					**23661101**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23661103	**Good identification**					**23661103**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23661121	**Good identification**					**23661121**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23661123	**Good identification**					**23661123**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23663001	**Good identification**					**23663001**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23663003	**Good identification**					**23663003**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23663021	**Good identification**					**23663021**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23663023	**Good identification**					**23663023**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23663100	**Good identification**					**23663100**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23663101	**Good identification**					**23663101**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23663102	**Good identification**					**23663102**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23663103	**Good identification**					**23663103**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23663120	**Good identification**					**23663120**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23663121	**Good identification**					**23663121**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23663122	**Good identification**					**23663122**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23663123	**Good identification**					**23663123**
Actinomyces naeslundii		100.0	0.43	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23670100	**Low discrimination**					**23670100**
					GRA	
Fusobacterium mortiferum		89.78	0.38	NIT 1	1	
Actinomyces naeslundii		10.22	0.33		99	
23670101	**Good identification**					**23670101**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23670103	**Good identification**					**23670103**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23670121	**Good identification**					**23670121**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23670123	**Good identification**					**23670123**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23671000	**Low discrimination**					**23671000**
					GRA	
Actinomyces naeslundii		59.89	0.49	MNS 88	99	
Fusobacterium mortiferum		40.11	0.34	MLT 94	1	
23671001	**Good identification**					**23671001**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23671002	**Acceptable identification**					**23671002**
					GRA	
Actinomyces naeslundii		93.07	0.49	MNS 88	99	
Fusobacterium mortiferum		6.93	0.18	MLT 94	1	
23671003	**Good identification**					**23671003**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23671020	**Good identification**					**23671020**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23671021	**Good identification**					**23671021**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23671022	**Good identification**					**23671022**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23671023	**Good identification**					**23671023**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23671040	**Good identification**					**23671040**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23671041	**Good identification**					**23671041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23671042	**Good identification**					**23671042**
Actinomyces naeslundii		100.0	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23671043	**Good identification**					**23671043**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23671060	**Good identification**					**23671060**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23671061	**Good identification**					**23671061**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23671062	**Good identification**					**23671062**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23671063	**Good identification**					**23671063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23671100	**Low discrimination**					**23671100**
					GRA	
Actinomyces naeslundii		64.02	0.57	MNS 88	99	
Fusobacterium mortiferum		35.16	0.41	MLT 94	1	
Actinomyces israelii		0.83	0.15		99	
23671101	**Very good identification**					**23671101**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23671102	**Good identification**					**23671102**
					GRA	
Actinomyces naeslundii		90.63	0.57	MNS 88	99	
Fusobacterium mortiferum		5.53	0.24	MLT 94	1	
Actinomyces israelii		1.76	0.18		99	
23671103	**Very good identification**					**23671103**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23671111	**Good identification**					**23671111**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23671113	**Good identification**					**23671113**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23671120	**Very good identification**					**23671120**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23671121	**Very good identification**					**23671121**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23671122	**Very good identification**					**23671122**
Actinomyces naeslundii		98.90	0.57	MNS 88		
Lactobacillus acidophilus		1.10	-.01	MLT 94		
23671123	**Very good identification**					**23671123**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23671131	**Good identification**					**23671131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
23671133	**Good identification**					**23671133**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23671140	**Good identification**					**23671140**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23671141	**Good identification**					**23671141**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23671142	**Good identification**					**23671142**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23671143	**Good identification**					**23671143**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23671160	**Good identification**					**23671160**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23671161	**Good identification**					**23671161**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23671162	**Good identification**					**23671162**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23671163	**Good identification**					**23671163**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23672100	**Good identification**					**23672100**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23672101	**Good identification**					**23672101**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23672102	**Good identification**					**23672102**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23672103	**Good identification**					**23672103**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23672120	**Good identification**					**23672120**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23672121	**Good identification**					**23672121**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23672122	**Good identification**					**23672122**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23672123	**Good identification**					**23672123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MNS 88		
23673000	**Good identification**					**23673000**
Actinomyces naeslundii		97.39	0.54	MNS 88		
Actinomyces israelii		2.61	0.18	MLT 94		
23673001	**Very good identification**					**23673001**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23673002	**Good identification**					**23673002**
Actinomyces naeslundii		96.13	0.54	MNS 88		
Actinomyces israelii		3.87	0.21	MLT 94		
23673003	**Very good identification**					**23673003**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23673011	**Good identification**					**23673011**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23673013	**Good identification**					**23673013**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23673020	**Good identification**					**23673020**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23673021	**Very good identification**					**23673021**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23673022	**Good identification**					**23673022**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23673023	**Very good identification**					**23673023**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23673031	**Good identification**					**23673031**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23673033	**Good identification**					**23673033**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23673040	**Good identification**					**23673040**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23673041	**Good identification**					**23673041**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23673042	**Acceptable identification**					**23673042**
Actinomyces naeslundii		91.85	0.41	MNS 88		
Actinomyces israelii		8.15	0.14	MLT 94		
23673043	**Good identification**					**23673043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	MNS 88		
23673060	**Good identification**					**23673060**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23673061	**Good identification**					**23673061**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23673062	**Good identification**					**23673062**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23673063	**Good identification**					**23673063**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23673100	**Very good identification**					**23673100**
Actinomyces naeslundii		96.37	0.62	MNS 88		
Actinomyces israelii		3.64	0.28	MLT 94		
23673101	**Very good identification**					**23673101**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23673102	**Good identification**					**23673102**
Actinomyces naeslundii		94.65	0.62	MNS 88		
Actinomyces israelii		5.36	0.31	MLT 94		
23673103	**Very good identification**					**23673103**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23673110	**Acceptable identification**					**23673110**
Actinomyces naeslundii		90.33	0.40	MNS 88		
Actinomyces israelii		9.67	0.14	MLT 94		
23673111	**Good identification**					**23673111**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23673112	**Low discrimination**					**23673112**
Actinomyces naeslundii		86.16	0.40	MNS 88		
Actinomyces israelii		13.84	0.17	MLT 94		
23673113	**Good identification**					**23673113**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23673120	**Very good identification**					**23673120**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23673121	**Very good identification**					**23673121**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23673122	**Very good identification**					**23673122**
Actinomyces naeslundii		100.0	0.62	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23673123	**Very good identification**					**23673123**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23673130	**Good identification**					**23673130**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23673131	**Good identification**					**23673131**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23673132	**Good identification**					**23673132**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23673133	**Good identification**					**23673133**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23673140	**Acceptable identification**					**23673140**
Actinomyces naeslundii		92.33	0.49	MNS 88		
Actinomyces israelii		7.67	0.21	MLT 94		
23673141	**Good identification**					**23673141**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23673142	**Low discrimination**					**23673142**
Actinomyces naeslundii		88.91	0.49	MNS 88		
Actinomyces israelii		11.09	0.24	MLT 94		
23673143	**Good identification**					**23673143**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23673160	**Good identification**					**23673160**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23673161	**Good identification**					**23673161**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23673162	**Good identification**					**23673162**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23673163	**Good identification**					**23673163**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23675102	**Good identification**					**23675102**
Actinomyces israelii		95.67	0.36	MLT 99		
Actinomyces naeslundii		4.34	0.23			
23677000	**Good identification**					**23677000**
Actinomyces israelii		100.0	0.36	MLT 99		
23677002	**Good identification**					**23677002**
Actinomyces israelii		100.0	0.39	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23677100	**Good identification**					**23677100**
Actinomyces israelii		97.73	0.46	MLT 99		
Actinomyces naeslundii		2.28	0.28			
23677102	**Good identification**					**23677102**
Actinomyces israelii		98.47	0.49	MLT 99		
Actinomyces naeslundii		1.53	0.28			
23677112	**Good identification**					**23677112**
Actinomyces israelii		100.0	0.35	MLT 99		
23677140	**Good identification**					**23677140**
Actinomyces israelii		100.0	0.39	MLT 99		
23677142	**Good identification**					**23677142**
Actinomyces israelii		100.0	0.42	MLT 99		
23677402	**Good identification**					**23677402**
Actinomyces israelii		100.0	0.38	MLT 99		
23677500	**Good identification**					**23677500**
Actinomyces israelii		100.0	0.45	MLT 99		
23677502	**Good identification**					**23677502**
Actinomyces israelii		100.0	0.48	MLT 99		
23677540	**Good identification**					**23677540**
Actinomyces israelii		100.0	0.38	MLT 99		
23677542	**Good identification**					**23677542**
Actinomyces israelii		100.0	0.41	MLT 99		
23700000	**Good identification**					**23700000**
Actinomyces odontolyticus		99.40	0.51	RAF 3		
Actinomyces viscosus		0.61	0.06	FRU 92		
23700002	**Low discrimination**					**23700002**
Actinomyces odontolyticus		50.44	0.47	RAF 3		
Actinomyces viscosus		49.56	0.40	FRU 92		
23700020	**Good identification**					**23700020**
Actinomyces odontolyticus		100.0	0.38	RAF 3		
23700100	**Good identification**					**23700100**
Actinomyces odontolyticus		95.86	0.43	RAF 3		
Actinomyces viscosus		3.28	0.11	FRU 92		
Actinomyces naeslundii		0.86	0.25			
23700102	**Low discrimination**					**23700102**
Actinomyces viscosus		84.50	0.45	MNS 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		15.28	0.39	FRU 95		
Actinomyces naeslundii		0.22	0.25			
23700103	**Good identification**					**23700103**
Actinomyces viscosus		98.82	0.37	MNS 96		
Actinomyces naeslundii		1.18	0.28	FRU 95		
23700200	**Good identification**					**23700200**
Actinomyces odontolyticus		100.0	0.56	RAF 3		
23700202	**Good identification**					**23700202**
Actinomyces odontolyticus		99.49	0.52	RAF 3		
Actinomyces viscosus		0.51	0.06	FRU 92		
23700220	**Good identification**					**23700220**
Actinomyces odontolyticus		100.0	0.43	RAF 3		
23700222	**Good identification**					**23700222**
Actinomyces odontolyticus		100.0	0.39	RAF 3		
23700300	**Good identification**					**23700300**
Actinomyces odontolyticus		100.0	0.47	RAF 3		
23700302	**Good identification**					**23700302**
Actinomyces odontolyticus		97.20	0.44	RAF 3		
Actinomyces viscosus		2.80	0.11	FRU 92		
23701000	**Low discrimination**					**23701000**
Actinomyces odontolyticus		85.11	0.48	RAF 3		
Eubacterium tortuosum		11.35	0.22	FRU 92		
Actinomyces naeslundii		3.06	0.41			
23701001	**Low discrimination**					**23701001**
Actinomyces naeslundii		84.23	0.44	MNS 88		
Actinomyces odontolyticus		15.77	0.14	FRU 95		
23701002	**Low discrimination**					**23701002**
Actinomyces odontolyticus		49.99	0.44	RAF 3		
Actinomyces viscosus		45.15	0.37	FRU 92		
Actinomyces naeslundii		2.93	0.41			
23701021	**Good identification**					**23701021**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23701023	**Good identification**					**23701023**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23701100	**Low discrimination**					**23701100**
Actinomyces odontolyticus		72.85	0.40	RAF 3		
Actinomyces naeslundii		23.61	0.48	FRU 92		
Actinomyces viscosus		2.29	0.07			
23701101	**Acceptable identification**					**23701101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		91.54	0.51	MNS 88	1	
Clostridium chauvoei		4.58	0.06	FRU 95	99	
Actinomyces viscosus		1.98	-.01		1	
23701102	**Low discrimination**					**23701102**
Actinomyces viscosus		76.88	0.41	MNS 96		
Actinomyces odontolyticus		15.12	0.36	FRU 95		
Actinomyces naeslundii		8.00	0.48			
23701120	**Low discrimination**					**23701120**
					GRA	
Actinomyces naeslundii		63.74	0.48	MNS 88	99	
Actinomyces odontolyticus		32.01	0.26	FRU 95	99	
Prevotella melaninogenica		4.26	-.01		1	
23701121	**Good identification**					**23701121**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23701122	**Low discrimination**					**23701122**
Actinomyces naeslundii		71.18	0.48	MNS 88		
Actinomyces odontolyticus		21.91	0.22	FRU 95		
Actinomyces viscosus		6.91	0.07			
23701123	**Good identification**					**23701123**
Actinomyces naeslundii		97.89	0.51	MNS 88		
Actinomyces viscosus		2.11	-.01	FRU 95		
23701140	**Low discrimination**					**23701140**
Actinomyces naeslundii		84.99	0.35	MNS 88		
Actinomyces odontolyticus		15.01	0.05	FRU 95		
23701141	**Good identification**					**23701141**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23701142	**Low discrimination**					**23701142**
Actinomyces naeslundii		64.50	0.35	MNS 88		
Actinomyces viscosus		35.50	0.07	FRU 95		
23701143	**Low discrimination**					**23701143**
Actinomyces naeslundii		89.10	0.38	MNS 88		
Actinomyces viscosus		10.90	-.01	FRU 95		
23701160	**Good identification**					**23701160**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23701161	**Good identification**					**23701161**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23701162	**Good identification**					**23701162**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23701163	**Good identification**					**23701163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
23701200	**Good identification**					**23701200**
Actinomyces odontolyticus		100.0	0.53	RAF 3		
23701202	**Good identification**					**23701202**
Actinomyces odontolyticus		99.53	0.49	RAF 3		
Actinomyces viscosus		0.47	0.03	FRU 92		
23701220	**Good identification**					**23701220**
Actinomyces odontolyticus		100.0	0.40	RAF 3		
23701222	**Good identification**					**23701222**
Actinomyces odontolyticus		100.0	0.36	RAF 3		
23701300	**Good identification**					**23701300**
Actinomyces odontolyticus		100.0	0.44	RAF 3		
23701302	**Good identification**					**23701302**
Actinomyces odontolyticus		97.42	0.41	RAF 3		
Actinomyces viscosus		2.58	0.07	FRU 92		
23703000	**Low discrimination**					**23703000**
Actinomyces naeslundii		87.36	0.46	MNS 88		
Actinomyces odontolyticus		12.64	0.14	FRU 95		
23703001	**Good identification**					**23703001**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23703002	**Low discrimination**					**23703002**
Actinomyces naeslundii		85.56	0.46	MNS 88		
Actinomyces odontolyticus		7.59	0.10	FRU 95		
Actinomyces viscosus		6.85	0.03			
23703003	**Good identification**					**23703003**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23703020	**Good identification**					**23703020**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23703021	**Good identification**					**23703021**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23703022	**Good identification**					**23703022**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23703023	**Good identification**					**23703023**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23703041	**Good identification**					**23703041**
Actinomyces naeslundii		100.0	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23703043	**Good identification**					**23703043**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23703061	**Good identification**					**23703061**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23703063	**Good identification**					**23703063**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23703100	**Low discrimination**					**23703100**
Actinomyces naeslundii		89.88	0.53	MNS 88		
Propionibacterium propionicum		8.68	0.26	FRU 95		
Actinomyces odontolyticus		1.44	0.05			
23703101	**Very good identification**					**23703101**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23703102	**Low discrimination**					**23703102**
Actinomyces naeslundii		83.69	0.53	MNS 88		
Propionibacterium propionicum		12.13	0.29	FRU 95		
Actinomyces viscosus		4.19	0.07			
23703103	**Very good identification**					**23703103**
Actinomyces naeslundii		97.86	0.56	MNS 88		
Actinomyces viscosus		1.09	-.01	FRU 95		
Propionibacterium propionicum		1.05	0.12			
23703120	**Good identification**					**23703120**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23703121	**Very good identification**					**23703121**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23703122	**Good identification**					**23703122**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23703123	**Very good identification**					**23703123**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23703140	**Low discrimination**					**23703140**
Actinomyces naeslundii		64.62	0.40	MNS 88		
Propionibacterium propionicum		35.38	0.26	FRU 95		
23703141	**Good identification**					**23703141**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23703142	**Low discrimination**					**23703142**
Actinomyces naeslundii		54.91	0.40	MNS 88		
Propionibacterium propionicum		45.09	0.29	FRU 95		
23703143	**Acceptable identification**					**23703143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		94.27	0.43	MNS 88		
Propionibacterium propionicum		5.73	0.12	FRU 95		
23703160	**Good identification**					**23703160**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23703161	**Good identification**					**23703161**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23703162	**Good identification**					**23703162**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23703163	**Good identification**					**23703163**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23704000	**Good identification**					**23704000**
Actinomyces odontolyticus		100.0	0.46	RAF 3		
23704002	**Good identification**					**23704002**
Actinomyces odontolyticus		98.03	0.42	RAF 3		
Actinomyces viscosus		1.98	0.06	FRU 92		
23704100	**Good identification**					**23704100**
Actinomyces odontolyticus		100.0	0.37	RAF 3		
23704200	**Good identification**					**23704200**
Actinomyces odontolyticus		100.0	0.51	RAF 3		
23704202	**Good identification**					**23704202**
Actinomyces odontolyticus		100.0	0.47	RAF 3		
23704220	**Good identification**					**23704220**
Actinomyces odontolyticus		100.0	0.37	RAF 3		
23704300	**Good identification**					**23704300**
Actinomyces odontolyticus		100.0	0.42	RAF 3		
23704302	**Good identification**					**23704302**
Actinomyces odontolyticus		100.0	0.39	RAF 3		
23705000	**Good identification**					**23705000**
Actinomyces odontolyticus		100.0	0.43	RAF 3		
23705002	**Good identification**					**23705002**
Actinomyces odontolyticus		98.18	0.39	RAF 3		
Actinomyces viscosus		1.82	0.03	FRU 92		
23705200	**Good identification**					**23705200**
Actinomyces odontolyticus		100.0	0.48	RAF 3		
23705202	**Good identification**					**23705202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.44	RAF 3		
23705300	**Good identification**					**23705300**
Actinomyces odontolyticus		100.0	0.39	RAF 3		
23705302	**Good identification**					**23705302**
Actinomyces odontolyticus		100.0	0.36	RAF 3		
23710000	**Very good identification**					**23710000**
					GRA	
Actinomyces odontolyticus		98.47	0.69	RAF 3	99	
Actinomyces viscosus		0.99	0.28		99	
Fusobacterium mortiferum		0.36	0.33		1	
23710001	**Low discrimination**					**23710001**
Actinomyces odontolyticus		63.42	0.35	RAF 3		
Actinomyces viscosus		21.03	0.20	URE 1		
Actinomyces naeslundii		15.55	0.42			
23710002	**Low discrimination**					**23710002**
Actinomyces viscosus		61.80	0.62	MNS 96		
Actinomyces odontolyticus		38.08	0.66			
Actinomyces naeslundii		0.10	0.39			
23710003	**Good identification**					**23710003**
Actinomyces viscosus		97.45	0.54	MNS 96		
Actinomyces odontolyticus		1.82	0.31			
Actinomyces naeslundii		0.73	0.42			
23710010	**Good identification**					**23710010**
Actinomyces odontolyticus		100.0	0.35	RAF 3		
23710020	**Good identification**					**23710020**
Actinomyces odontolyticus		99.00	0.56	RAF 3		
Actinomyces naeslundii		1.00	0.39	NAG 14		
23710021	**Low discrimination**					**23710021**
Actinomyces naeslundii		60.09	0.42	MNS 88		
Actinomyces odontolyticus		39.91	0.22	GAL 96		
23710022	**Low discrimination**					**23710022**
Actinomyces odontolyticus		89.51	0.52	RAF 3		
Actinomyces viscosus		9.01	0.28	NAG 14		
Actinomyces naeslundii		1.48	0.39			
23710040	**Good identification**					**23710040**
Actinomyces odontolyticus		97.20	0.35	RAF 3		
Actinomyces naeslundii		2.80	0.26	SOR 1		
23710100	**Good identification**					**23710100**
					GRA	
Actinomyces odontolyticus		90.78	0.61	RAF 3	99	
Actinomyces viscosus		5.14	0.32		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Fusobacterium mortiferum	2.42	0.39		1	
23710101	**Low discrimination**					**23710101**
	Actinomyces naeslundii	43.52	0.50	MNS 88		
	Actinomyces viscosus	36.78	0.24	GAL 96		
	Actinomyces odontolyticus	19.70	0.27			
23710102	**Low discrimination**					**23710102**
	Actinomyces viscosus	89.80	0.67	MNS 96		
	Actinomyces odontolyticus	9.83	0.57			
	Actinomyces naeslundii	0.24	0.47			
23710103	**Very good identification**					**23710103**
	Actinomyces viscosus	98.47	0.58	MNS 96		
	Actinomyces naeslundii	1.18	0.50			
	Actinomyces odontolyticus	0.33	0.23			
23710120	**Acceptable identification**					**23710120**
	Actinomyces odontolyticus	91.33	0.47	RAF 3		
	Actinomyces naeslundii	8.35	0.47	NAG 14		
	Actinomyces viscosus	0.32	-.02			
23710121	**Acceptable identification**					**23710121**
	Actinomyces naeslundii	93.14	0.50	MNS 88		
	Actinomyces odontolyticus	6.86	0.13	GAL 96		
23710122	**Low discrimination**					**23710122**
	Actinomyces odontolyticus	58.27	0.44	RAF 3		
	Actinomyces viscosus	33.04	0.32	NAG 14		
	Actinomyces naeslundii	8.69	0.47			
23710123	**Low discrimination**					**23710123**
	Actinomyces naeslundii	52.91	0.50	MNS 88		
	Actinomyces viscosus	44.71	0.24	GAL 96		
	Actinomyces odontolyticus	2.39	0.09			
23710141	**Good identification**					**23710141**
	Actinomyces naeslundii	100.0	0.37	MNS 88		
23710161	**Good identification**					**23710161**
	Actinomyces naeslundii	100.0	0.37	MNS 88		
23710163	**Good identification**					**23710163**
	Actinomyces naeslundii	100.0	0.37	MNS 88		
23710200	**Very good identification**					**23710200**
	Actinomyces odontolyticus	100.0	0.74	RAF 3		
23710201	**Good identification**					**23710201**
	Actinomyces odontolyticus	100.0	0.40	RAF 3		
23710202	**Very good identification**					**23710202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.16	0.71	RAF 3		
Actinomyces viscosus		0.84	0.28			
23710203	**Low discrimination**					**23710203**
Actinomyces odontolyticus		78.20	0.36	RAF 3		
Actinomyces viscosus		21.80	0.20	URE 1		
23710210	**Good identification**					**23710210**
Actinomyces odontolyticus		100.0	0.40	RAF 3		
23710212	**Good identification**					**23710212**
Actinomyces odontolyticus		100.0	0.36	RAF 3		
23710220	**Very good identification**					**23710220**
Actinomyces odontolyticus		100.0	0.61	RAF 3		
23710222	**Very good identification**					**23710222**
Actinomyces odontolyticus		100.0	0.57	RAF 3		
23710240	**Good identification**					**23710240**
Actinomyces odontolyticus		100.0	0.40	RAF 3		
23710242	**Good identification**					**23710242**
Actinomyces odontolyticus		100.0	0.36	RAF 3		
23710300	**Very good identification**					**23710300**
Actinomyces odontolyticus		99.97	0.66	RAF 3		
Actinomyces viscosus		0.03	-.02			
23710302	**Very good identification**					**23710302**
Actinomyces odontolyticus		95.46	0.62	RAF 3		
Actinomyces viscosus		4.54	0.32			
23710320	**Good identification**					**23710320**
Actinomyces odontolyticus		100.0	0.52	RAF 3		
23710322	**Good identification**					**23710322**
Actinomyces odontolyticus		99.71	0.49	RAF 3		
Actinomyces viscosus		0.29	-.02	NAG 14		
23710400	**Good identification**					**23710400**
Actinomyces odontolyticus		100.0	0.35	RAF 3		
23710600	**Good identification**					**23710600**
Actinomyces odontolyticus		100.0	0.40	RAF 3		
23710602	**Good identification**					**23710602**
Actinomyces odontolyticus		100.0	0.36	RAF 3		
23711000	**Good identification**					**23711000**
Actinomyces odontolyticus		92.64	0.66	RAF 3		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		5.51	0.62			
Actinomyces viscosus		0.86	0.24			
23711001	**Low discrimination**					**23711001**
Actinomyces naeslundii		87.12	0.66	MNS 88		
Actinomyces odontolyticus		9.87	0.32			
Actinomyces viscosus		3.01	0.16			
23711002	**Low discrimination**					**23711002**
Actinomyces viscosus		57.52	0.59	MNS 96		
Actinomyces odontolyticus		38.55	0.63			
Actinomyces naeslundii		3.74	0.62			
23711003	**Low discrimination**					**23711003**
Actinomyces viscosus		76.18	0.50	MNS 96		
Actinomyces naeslundii		22.27	0.66			
Actinomyces odontolyticus		1.55	0.28			
23711011	**Good identification**					**23711011**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23711013	**Low discrimination**					**23711013**
Actinomyces naeslundii		60.37	0.44	MNS 88		
Actinomyces viscosus		39.63	0.16	MLZ 5		
23711020	**Low discrimination**					**23711020**
Actinomyces odontolyticus		72.34	0.53	RAF 3		
Actinomyces naeslundii		26.40	0.62	NAG 14		
Clostridium perfringens		0.93	0.15			
23711021	**Very good identification**					**23711021**
Actinomyces naeslundii		98.19	0.66	MNS 88		
Actinomyces odontolyticus		1.81	0.19			
23711022	**Low discrimination**					**23711022**
Actinomyces odontolyticus		58.64	0.49	RAF 3		
Actinomyces naeslundii		34.92	0.62	NAG 14		
Actinomyces viscosus		5.43	0.24			
23711023	**Very good identification**					**23711023**
Actinomyces naeslundii		95.62	0.66	MNS 88		
Actinomyces viscosus		3.30	0.16			
Actinomyces odontolyticus		1.08	0.15			
23711030	**Low discrimination**					**23711030**
Actinomyces naeslundii		65.54	0.41	MNS 88		
Actinomyces odontolyticus		34.46	0.19	MLZ 5		
23711031	**Good identification**					**23711031**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23711032	**Low discrimination**					**23711032**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		75.63	0.41	MNS 88		
Actinomyces odontolyticus		24.37	0.15	MLZ 5		
23711033	**Good identification**					**23711033**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23711040	**Low discrimination**					**23711040**
Actinomyces naeslundii		50.93	0.50	MNS 88		
Actinomyces odontolyticus		49.07	0.32			
23711041	**Good identification**					**23711041**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23711042	**Low discrimination**					**23711042**
Actinomyces naeslundii		40.46	0.50	MNS 88		
Actinomyces viscosus		35.65	0.24			
Actinomyces odontolyticus		23.89	0.28			
23711043	**Low discrimination**					**23711043**
Actinomyces naeslundii		83.63	0.53	MNS 88		
Actinomyces viscosus		16.37	0.16			
23711060	**Low discrimination**					**23711060**
Actinomyces naeslundii		83.71	0.50	MNS 88		
Actinomyces odontolyticus		13.13	0.19			
Clostridium perfringens		3.17	0.03			
23711061	**Good identification**					**23711061**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23711062	**Acceptable identification**					**23711062**
Actinomyces naeslundii		91.23	0.50	MNS 88		
Actinomyces odontolyticus		8.77	0.15			
23711063	**Good identification**					**23711063**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23711100	**Low discrimination**					**23711100**
					GRA	
Actinomyces odontolyticus		59.95	0.58	RAF 3	99	
Actinomyces naeslundii		32.10	0.70		99	
Fusobacterium mortiferum		3.59	0.42		1	
23711101	**Very good identification**					**23711101**
Actinomyces naeslundii		96.13	0.73	MNS 88		
Actinomyces viscosus		2.07	0.21			
Actinomyces odontolyticus		1.21	0.24			
23711102	**Low discrimination**					**23711102**
Actinomyces viscosus		81.36	0.63	MNS 96		
Actinomyces odontolyticus		9.69	0.54			
Actinomyces naeslundii		8.46	0.70			
23711103	**Low discrimination**					**23711103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		67.90	0.55	MNS 96			
Actinomyces naeslundii		31.78	0.73				
Actinomyces odontolyticus		0.25	0.20				
23711110	**Low discrimination**						**23711110**
Actinomyces naeslundii		73.62	0.48	MNS 88			
Actinomyces odontolyticus		26.38	0.24	MLZ 5			
23711111	**Good identification**						**23711111**
Actinomyces naeslundii		100.0	0.51	MNS 88			
23711113	**Low discrimination**						**23711113**
Actinomyces naeslundii		70.92	0.51	MNS 88			
Actinomyces viscosus		29.08	0.21	MLZ 5			
23711120	**Low discrimination**						**23711120**
Actinomyces naeslundii		69.68	0.70	MNS 88			
Actinomyces odontolyticus		21.18	0.44				
Clostridium perfringens		6.29	0.30				
23711121	**Very good identification**						**23711121**
					SPO		
Actinomyces naeslundii		99.59	0.73	MNS 88	1		
Actinomyces odontolyticus		0.20	0.10		1		
Clostridium chauvoei		0.20	0.04		99		
23711122	**Low discrimination**						**23711122**
Actinomyces naeslundii		73.00	0.70	MNS 88			
Actinomyces odontolyticus		13.60	0.41				
Actinomyces viscosus		7.09	0.29				
23711123	**Very good identification**						**23711123**
Actinomyces naeslundii		97.77	0.73	MNS 88			
Actinomyces viscosus		2.11	0.21				
Actinomyces odontolyticus		0.12	0.06				
23711130	**Acceptable identification**						**23711130**
Actinomyces naeslundii		94.49	0.48	MNS 88			
Actinomyces odontolyticus		5.51	0.10	MLZ 5			
23711131	**Good identification**						**23711131**
Actinomyces naeslundii		100.0	0.51	MNS 88			
23711132	**Acceptable identification**						**23711132**
					GRA	CAT	
Actinomyces naeslundii		92.95	0.48	MNS 88	99	25	
Capnocytophaga ochracea		3.73	-.01	MLZ 5	1	90	
Actinomyces odontolyticus		3.32	0.06		99	10	
23711133	**Good identification**						**23711133**
Actinomyces naeslundii		100.0	0.51	MNS 88			
23711140	**Low discrimination**						**23711140**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		85.32	0.57	MNS 88			
Actinomyces odontolyticus		9.12	0.24				
Propionibacterium propionicum		4.77	0.26				
23711141	**Very good identification**						**23711141**
Actinomyces naeslundii		100.0	0.60	MNS 88			
23711142	**Low discrimination**						**23711142**
Actinomyces naeslundii		58.36	0.57	MNS 88			
Actinomyces viscosus		32.11	0.29				
Propionibacterium propionicum		4.90	0.29				
23711143	**Low discrimination**						**23711143**
Actinomyces naeslundii		88.61	0.60	MNS 88			
Actinomyces viscosus		10.84	0.21				
Propionibacterium propionicum		0.55	0.12				
23711150	**Good identification**						**23711150**
Actinomyces naeslundii		100.0	0.35	MNS 88			
23711151	**Good identification**						**23711151**
Actinomyces naeslundii		100.0	0.38	MNS 88			
23711152	**Good identification**						**23711152**
Actinomyces naeslundii		100.0	0.35	MNS 88			
23711153	**Good identification**						**23711153**
Actinomyces naeslundii		100.0	0.38	MNS 88			
23711160	**Low discrimination**						**23711160**
Actinomyces naeslundii		89.70	0.57	MNS 88			
Clostridium perfringens		8.74	0.18				
Actinomyces odontolyticus		1.56	0.10				
23711161	**Very good identification**						**23711161**
Actinomyces naeslundii		100.0	0.60	MNS 88			
23711162	**Good identification**						**23711162**
					GRA	CAT	
Actinomyces naeslundii		93.59	0.57	MNS 88	99	25	
Clostridium perfringens		4.29	0.12		99	1	
Capnocytophaga ochracea		1.12	-.01		1	90	
23711163	**Very good identification**						**23711163**
Actinomyces naeslundii		100.0	0.60	MNS 88			
23711170	**Good identification**						**23711170**
Actinomyces naeslundii		100.0	0.35	MNS 88			
23711171	**Good identification**						**23711171**
Actinomyces naeslundii		100.0	0.38	MNS 88			
23711172	**Good identification**						**23711172**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	MNS 88		
23711173	**Good identification**					**23711173**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23711200	**Very good identification**					**23711200**
Actinomyces odontolyticus		99.97	0.71	RAF 3		
Actinomyces naeslundii		0.03	0.28			
23711201	**Good identification**					**23711201**
Actinomyces odontolyticus		95.61	0.37	RAF 3		
Actinomyces naeslundii		4.39	0.31	URE 1		
23711202	**Very good identification**					**23711202**
Actinomyces odontolyticus		99.18	0.68	RAF 3		
Actinomyces viscosus		0.77	0.24			
Actinomyces naeslundii		0.05	0.28			
23711210	**Good identification**					**23711210**
Actinomyces odontolyticus		100.0	0.37	RAF 3		
23711220	**Very good identification**					**23711220**
Actinomyces odontolyticus		99.81	0.58	RAF 3		
Actinomyces naeslundii		0.19	0.28	NAG 14		
23711222	**Good identification**					**23711222**
Actinomyces odontolyticus		99.69	0.54	RAF 3		
Actinomyces naeslundii		0.31	0.28	NAG 14		
23711240	**Good identification**					**23711240**
Actinomyces odontolyticus		100.0	0.37	RAF 3		
23711300	**Very good identification**					**23711300**
Actinomyces odontolyticus		99.72	0.63	RAF 3		
Actinomyces naeslundii		0.28	0.36			
23711302	**Very good identification**					**23711302**
Actinomyces odontolyticus		95.40	0.59	RAF 3		
Actinomyces viscosus		4.17	0.29			
Actinomyces naeslundii		0.43	0.36			
23711320	**Good identification**					**23711320**
Actinomyces odontolyticus		98.32	0.49	RAF 3		
Actinomyces naeslundii		1.68	0.36	NAG 14		
23711321	**Low discrimination**					**23711321**
Actinomyces naeslundii		71.77	0.39	MNS 88		
Actinomyces odontolyticus		28.23	0.15	RHA 1		
23711322	**Good identification**					**23711322**
					GRA	
Actinomyces odontolyticus		96.30	0.45	RAF 3	99	
Actinomyces naeslundii		2.69	0.36	NAG 14	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		1.01	0.03		1		
23711323	**Low discrimination**						**23711323**
Actinomyces naeslundii		80.58	0.39	MNS 88			
Actinomyces odontolyticus		19.43	0.11	RHA 1			
23711501	**Good identification**						**23711501**
Actinomyces naeslundii		100.0	0.39	MNS 88			
23711520	**Low discrimination**						**23711520**
Actinomyces naeslundii		45.83	0.36	MNS 88			
Lactobacillus fermentum		32.80	0.24	ARA 1			
Actinomyces odontolyticus		13.93	0.10				
23711521	**Good identification**						**23711521**
Actinomyces naeslundii		100.0	0.39	MNS 88			
23711522	**Low discrimination**						**23711522**
					GRA	CAT	
Actinomyces naeslundii		41.74	0.36	MNS 88	99	25	
Lactobacillus fermentum		29.87	0.24	ARA 1	99	1	
Capnocytophaga ochracea		13.85	0.02		1	90	
23711523	**Good identification**						**23711523**
Actinomyces naeslundii		100.0	0.39	MNS 88			
23711600	**Good identification**						**23711600**
Actinomyces odontolyticus		100.0	0.37	RAF 3			
23712000	**Low discrimination**						**23712000**
Actinomyces odontolyticus		70.06	0.35	RAF 3			
Actinomyces naeslundii		22.22	0.44	MAN 1			
Propionibacterium propionicum		7.72	0.26				
23712001	**Good identification**						**23712001**
Actinomyces naeslundii		100.0	0.47	MNS 88			
23712003	**Low discrimination**						**23712003**
Actinomyces naeslundii		57.62	0.47	MNS 88			
Actinomyces viscosus		40.16	0.20	GAL 96			
Propionibacterium propionicum		2.22	0.12				
23712020	**Low discrimination**						**23712020**
Actinomyces naeslundii		66.08	0.44	MNS 88			
Actinomyces odontolyticus		33.92	0.22	GAL 96			
23712021	**Good identification**						**23712021**
Actinomyces naeslundii		100.0	0.47	MNS 88			
23712022	**Low discrimination**						**23712022**
Actinomyces naeslundii		76.07	0.44	MNS 88			
Actinomyces odontolyticus		23.93	0.18	GAL 96			
23712023	**Good identification**						**23712023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	MNS 88		
23712100	**Low discrimination**					**23712100**
Propionibacterium propionicum		89.82	0.60	SAL 1		
Actinomyces naeslundii		7.44	0.52			
Actinomyces odontolyticus		2.60	0.27			
23712101	**Low discrimination**					**23712101**
Actinomyces naeslundii		52.77	0.55	MNS 88		
Propionibacterium propionicum		47.23	0.44	GAL 96		
23712102	**Low discrimination**					**23712102**
Propionibacterium propionicum		85.10	0.63	SAL 1		
Actinomyces viscosus		9.20	0.32			
Actinomyces naeslundii		4.70	0.52			
23712103	**Low discrimination**					**23712103**
Propionibacterium propionicum		48.33	0.47	SAL 1		
Actinomyces naeslundii		36.00	0.55	URE 10		
Actinomyces viscosus		15.67	0.24			
23712120	**Low discrimination**					**23712120**
Actinomyces naeslundii		84.82	0.52	MNS 88		
Propionibacterium propionicum		10.35	0.26	GAL 96		
Actinomyces odontolyticus		4.83	0.13			
23712121	**Good identification**					**23712121**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23712122	**Low discrimination**					**23712122**
Actinomyces naeslundii		80.79	0.52	MNS 88		
Propionibacterium propionicum		14.79	0.29	GAL 96		
Actinomyces odontolyticus		2.82	0.09			
23712123	**Good identification**					**23712123**
Actinomyces naeslundii		98.66	0.55	MNS 88		
Propionibacterium propionicum		1.34	0.12	GAL 96		
23712140	**Very good identification**					**23712140**
Propionibacterium propionicum		98.56	0.60	SAL 1		
Actinomyces naeslundii		1.44	0.39			
23712141	**Low discrimination**					**23712141**
Propionibacterium propionicum		83.53	0.44	SAL 1		
Actinomyces naeslundii		16.47	0.42	URE 10		
23712142	**Very good identification**					**23712142**
Propionibacterium propionicum		98.93	0.63	SAL 1		
Actinomyces naeslundii		0.96	0.39			
Actinomyces viscosus		0.11	-.02			
23712143	**Low discrimination**					**23712143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		88.38	0.47	SAL 1		
Actinomyces naeslundii		11.62	0.42	URE 10		
23712160	**Low discrimination**					**23712160**
Actinomyces naeslundii		59.12	0.39	MNS 88		
Propionibacterium propionicum		40.88	0.26	GAL 96		
23712161	**Good identification**					**23712161**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23712163	**Acceptable identification**					**23712163**
Actinomyces naeslundii		92.87	0.42	MNS 88		
Propionibacterium propionicum		7.13	0.12	GAL 96		
23712200	**Good identification**					**23712200**
Actinomyces odontolyticus		100.0	0.40	RAF 3		
23712202	**Good identification**					**23712202**
Actinomyces odontolyticus		100.0	0.36	RAF 3		
23713000	**Good identification**					**23713000**
Actinomyces naeslundii		90.03	0.67	MNS 88		
Actinomyces odontolyticus		7.88	0.32			
Propionibacterium propionicum		1.30	0.26			
23713001	**Very good identification**					**23713001**
Actinomyces naeslundii		100.0	0.70	MNS 88		
23713002	**Low discrimination**					**23713002**
Actinomyces naeslundii		85.57	0.67	MNS 88		
Actinomyces viscosus		6.85	0.24			
Actinomyces odontolyticus		4.59	0.28			
23713003	**Very good identification**					**23713003**
Actinomyces naeslundii		98.10	0.70	MNS 88		
Actinomyces viscosus		1.75	0.16			
Propionibacterium propionicum		0.16	0.12			
23713010	**Good identification**					**23713010**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23713011	**Good identification**					**23713011**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23713012	**Good identification**					**23713012**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23713013	**Good identification**					**23713013**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23713020	**Very good identification**					**23713020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.59	0.67	MNS 88		
Actinomyces odontolyticus		1.41	0.19			
23713021	**Very good identification**					**23713021**
Actinomyces naeslundii		100.0	0.70	MNS 88		
23713022	**Very good identification**					**23713022**
Actinomyces naeslundii		99.13	0.67	MNS 88		
Actinomyces odontolyticus		0.87	0.15			
23713023	**Very good identification**					**23713023**
Actinomyces naeslundii		100.0	0.70	MNS 88		
23713030	**Good identification**					**23713030**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23713031	**Good identification**					**23713031**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23713032	**Good identification**					**23713032**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23713033	**Good identification**					**23713033**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23713040	**Acceptable identification**					**23713040**
Actinomyces naeslundii		90.81	0.55	MNS 88		
Propionibacterium propionicum		7.45	0.26			
Actinomyces israelii		1.75	0.16			
23713041	**Very good identification**					**23713041**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23713042	**Low discrimination**					**23713042**
Actinomyces naeslundii		86.82	0.55	MNS 88		
Propionibacterium propionicum		10.68	0.29			
Actinomyces israelii		2.51	0.19			
23713043	**Very good identification**					**23713043**
Actinomyces naeslundii		99.10	0.58	MNS 88		
Propionibacterium propionicum		0.90	0.12			
23713051	**Good identification**					**23713051**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713053	**Good identification**					**23713053**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713060	**Good identification**					**23713060**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23713061	**Very good identification**					**23713061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.58	MNS 88		
23713062	**Good identification**					**23713062**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23713063	**Very good identification**					**23713063**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23713071	**Good identification**					**23713071**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713073	**Good identification**					**23713073**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713100	**Low discrimination**					**23713100**
Actinomyces naeslundii		65.56	0.75	MNS 88		
Propionibacterium propionicum		33.00	0.60			
Actinomyces israelii		0.81	0.33			
23713101	**Very good identification**					**23713101**
Actinomyces naeslundii		96.41	0.78	MNS 88		
Propionibacterium propionicum		3.60	0.44			
23713102	**Low discrimination**					**23713102**
Actinomyces naeslundii		54.66	0.75	MNS 88		
Propionibacterium propionicum		41.27	0.63			
Actinomyces viscosus		2.74	0.29			
23713103	**Good identification**					**23713103**
Actinomyces naeslundii		93.72	0.78	MNS 88		
Propionibacterium propionicum		5.24	0.47			
Actinomyces viscosus		1.04	0.21			
23713110	**Low discrimination**					**23713110**
Actinomyces naeslundii		88.38	0.53	MNS 88		
Propionibacterium propionicum		8.54	0.26	MLZ 5		
Actinomyces israelii		3.08	0.19			
23713111	**Very good identification**					**23713111**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23713112	**Low discrimination**					**23713112**
Actinomyces naeslundii		83.53	0.53	MNS 88		
Propionibacterium propionicum		12.10	0.29	MLZ 5		
Actinomyces israelii		4.37	0.22			
23713113	**Very good identification**					**23713113**
Actinomyces naeslundii		98.94	0.56	MNS 88		
Propionibacterium propionicum		1.06	0.12	MLZ 5		
23713120	**Very good identification**					**23713120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.34	0.75	MNS 88		
Propionibacterium propionicum		0.51	0.26			
Actinomyces odontolyticus		0.16	0.10			
23713121	**Very good identification**					**23713121**
Actinomyces naeslundii		100.0	0.78	MNS 88		
23713122	**Very good identification**					**23713122**
Actinomyces naeslundii		99.03	0.75	MNS 88		
Propionibacterium propionicum		0.76	0.29			
Actinomyces israelii		0.12	0.16			
23713123	**Very good identification**					**23713123**
Actinomyces naeslundii		99.94	0.78	MNS 88		
Propionibacterium propionicum		0.06	0.12			
23713130	**Good identification**					**23713130**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23713131	**Very good identification**					**23713131**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23713132	**Good identification**					**23713132**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23713133	**Very good identification**					**23713133**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23713140	**Low discrimination**					**23713140**
Propionibacterium propionicum		73.53	0.60	SAL 1		
Actinomyces naeslundii		25.78	0.62			
Actinomyces israelii		0.70	0.26			
23713141	**Low discrimination**					**23713141**
Actinomyces naeslundii		82.56	0.65	MNS 88		
Propionibacterium propionicum		17.44	0.44			
23713142	**Low discrimination**					**23713142**
Propionibacterium propionicum		80.44	0.63	SAL 1		
Actinomyces naeslundii		18.80	0.62			
Actinomyces israelii		0.76	0.29			
23713143	**Low discrimination**					**23713143**
Actinomyces naeslundii		75.93	0.65	MNS 88		
Propionibacterium propionicum		24.07	0.47			
23713150	**Low discrimination**					**23713150**
Actinomyces naeslundii		64.62	0.40	MNS 88		
Propionibacterium propionicum		35.38	0.26	MLZ 5		
23713151	**Good identification**					**23713151**
Actinomyces naeslundii		100.0	0.43	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23713152	**Low discrimination**					**23713152**
Actinomyces naeslundii		51.65	0.40	MNS 88		
Propionibacterium propionicum		42.41	0.29	MLZ 5		
Actinomyces israelii		5.95	0.15			
23713153	**Acceptable identification**					**23713153**
Actinomyces naeslundii		94.27	0.43	MNS 88		
Propionibacterium propionicum		5.73	0.12	MLZ 5		
23713160	**Very good identification**					**23713160**
Actinomyces naeslundii		97.20	0.62	MNS 88		
Propionibacterium propionicum		2.80	0.26			
23713161	**Very good identification**					**23713161**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23713162	**Very good identification**					**23713162**
Actinomyces naeslundii		95.86	0.62	MNS 88		
Propionibacterium propionicum		4.14	0.29			
23713163	**Very good identification**					**23713163**
Actinomyces naeslundii		99.68	0.65	MNS 88		
Propionibacterium propionicum		0.32	0.12			
23713170	**Good identification**					**23713170**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23713171	**Good identification**					**23713171**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23713172	**Good identification**					**23713172**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23713173	**Good identification**					**23713173**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23713200	**Acceptable identification**					**23713200**
Actinomyces odontolyticus		94.39	0.37	RAF 3		
Actinomyces naeslundii		5.61	0.33	MAN 1		
23713201	**Good identification**					**23713201**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713203	**Good identification**					**23713203**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713221	**Good identification**					**23713221**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713223	**Good identification**					**23713223**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713301	**Good identification**					**23713301**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23713303	**Acceptable identification**					**23713303**
Actinomyces naeslundii		94.70	0.44	MNS 88		
Propionibacterium propionicum		5.30	0.12	RHA 1		
23713320	**Low discrimination**					**23713320**
Actinomyces naeslundii		76.69	0.41	MNS 88		
Actinomyces odontolyticus		23.31	0.15	RHA 1		
23713321	**Good identification**					**23713321**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23713322	**Low discrimination**					**23713322**
Actinomyces naeslundii		84.30	0.41	MNS 88		
Actinomyces odontolyticus		15.70	0.11	RHA 1		
23713323	**Good identification**					**23713323**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23713401	**Good identification**					**23713401**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713403	**Good identification**					**23713403**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713421	**Good identification**					**23713421**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713423	**Good identification**					**23713423**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23713500	**Low discrimination**					**23713500**
Actinomyces naeslundii		40.04	0.41	MNS 88		
Actinomyces israelii		39.80	0.32	ARA 1		
Propionibacterium propionicum		20.16	0.26			
23713501	**Good identification**					**23713501**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23713503	**Acceptable identification**					**23713503**
Actinomyces naeslundii		94.70	0.44	MNS 88		
Propionibacterium propionicum		5.30	0.12	ARA 1		
23713520	**Good identification**					**23713520**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23713521	**Good identification**					**23713521**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	MNS 88		
23713522	**Acceptable identification**					**23713522**
Actinomyces naeslundii		91.31	0.41	MNS 88		
Actinomyces israelii		8.69	0.14	ARA 1		
23713523	**Good identification**					**23713523**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23714000	**Very good identification**					**23714000**
					GRA	
Actinomyces odontolyticus		99.95	0.64	RAF 3	99	
Fusobacterium mortiferum		0.06	0.14		1	
23714002	**Very good identification**					**23714002**
Actinomyces odontolyticus		96.78	0.60	RAF 3		
Actinomyces viscosus		3.22	0.28			
23714020	**Good identification**					**23714020**
Actinomyces odontolyticus		100.0	0.51	RAF 3		
23714022	**Good identification**					**23714022**
Actinomyces odontolyticus		100.0	0.47	RAF 3		
23714100	**Good identification**					**23714100**
					GRA	
Actinomyces odontolyticus		99.48	0.55	RAF 3	99	
Fusobacterium mortiferum		0.41	0.20		1	
Actinomyces viscosus		0.12	-.02		99	
23714102	**Low discrimination**					**23714102**
Actinomyces odontolyticus		84.22	0.52	RAF 3		
Actinomyces viscosus		15.78	0.32			
23714120	**Good identification**					**23714120**
Actinomyces odontolyticus		100.0	0.42	RAF 3		
23714122	**Good identification**					**23714122**
Actinomyces odontolyticus		98.85	0.38	RAF 3		
Actinomyces viscosus		1.15	-.02	NAG 14		
23714200	**Very good identification**					**23714200**
Actinomyces odontolyticus		100.0	0.69	RAF 3		
23714202	**Very good identification**					**23714202**
Actinomyces odontolyticus		100.0	0.65	RAF 3		
23714220	**Good identification**					**23714220**
Actinomyces odontolyticus		100.0	0.55	RAF 3		
23714222	**Good identification**					**23714222**
Actinomyces odontolyticus		100.0	0.52	RAF 3		
23714300	**Very good identification**					**23714300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.60	RAF 3		
23714302	**Very good identification**					**23714302**
Actinomyces odontolyticus		99.90	0.57	RAF 3		
Actinomyces viscosus		0.10	-.02			
23714320	**Good identification**					**23714320**
Actinomyces odontolyticus		100.0	0.47	RAF 3		
23714322	**Good identification**					**23714322**
Actinomyces odontolyticus		100.0	0.43	RAF 3		
23715000	**Very good identification**					**23715000**
Actinomyces odontolyticus		99.08	0.61	RAF 3		
Actinomyces israelii		0.41	0.28			
Lactobacillus fermentum		0.27	0.24			
23715002	**Very good identification**					**23715002**
Actinomyces odontolyticus		95.50	0.57	RAF 3		
Actinomyces viscosus		2.92	0.24			
Actinomyces israelii		0.97	0.31			
23715020	**Good identification**					**23715020**
Actinomyces odontolyticus		98.72	0.48	RAF 3		
Actinomyces naeslundii		0.74	0.28	NAG 14		
Lactobacillus fermentum		0.54	0.16			
23715022	**Good identification**					**23715022**
Actinomyces odontolyticus		97.93	0.44	RAF 3		
Actinomyces naeslundii		1.20	0.28	NAG 14		
Lactobacillus fermentum		0.88	0.16			
23715100	**Acceptable identification**					**23715100**
Actinomyces odontolyticus		92.03	0.52	RAF 3		
Actinomyces israelii		4.84	0.39			
Lactobacillus fermentum		1.27	0.28			
23715101	**Low discrimination**					**23715101**
Actinomyces naeslundii		61.99	0.39	MNS 88		
Actinomyces odontolyticus		38.01	0.18	XYL 1		
23715102	**Low discrimination**					**23715102**
Actinomyces odontolyticus		73.61	0.49	RAF 3		
Actinomyces viscosus		12.68	0.29			
Actinomyces israelii		9.47	0.42			
23715120	**Low discrimination**					**23715120**
Actinomyces odontolyticus		89.57	0.39	RAF 3		
Actinomyces naeslundii		6.04	0.36	NAG 14		
Lactobacillus fermentum		2.54	0.20			
23715121	**Good identification**					**23715121**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	MNS 88			
23715122	**Low discrimination**						**23715122**
					GRA	CAT	
Actinomyces odontolyticus		76.56	0.35	RAF 3	99	10	
Actinomyces naeslundii		8.43	0.36	NAG 14	99	25	
Capnocytophaga ochracea		3.96	0.05		1	90	
23715123	**Good identification**						**23715123**
Actinomyces naeslundii		100.0	0.39	MNS 88			
23715200	**Very good identification**						**23715200**
Actinomyces odontolyticus		100.0	0.66	RAF 3			
23715202	**Very good identification**						**23715202**
Actinomyces odontolyticus		100.0	0.62	RAF 3			
23715220	**Good identification**						**23715220**
Actinomyces odontolyticus		100.0	0.52	RAF 3			
23715222	**Good identification**						**23715222**
Actinomyces odontolyticus		100.0	0.49	RAF 3			
23715300	**Very good identification**						**23715300**
Actinomyces odontolyticus		95.62	0.57	RAF 3			
Eubacterium contortum		4.19	0.27				
Actinomyces israelii		0.20	0.19				
23715302	**Good identification**						**23715302**
Actinomyces odontolyticus		98.27	0.54	RAF 3			
Eubacterium contortum		1.24	0.14				
Actinomyces israelii		0.49	0.22				
23715320	**Acceptable identification**						**23715320**
					GRA		
Actinomyces odontolyticus		94.51	0.44	RAF 3	99		
Eubacterium contortum		4.49	0.14	NAG 14	99		
Bacteroides vulgatus		1.00	-.04		1		
23715322	**Good identification**						**23715322**
					GRA		
Actinomyces odontolyticus		95.42	0.40	RAF 3	99		
Parabacteroides distasonis		3.27	0.19	NAG 14	1		
Eubacterium contortum		1.31	0.01		99		
23715500	**Low discrimination**						**23715500**
Actinomyces israelii		64.23	0.37	CEL 93			
Lactobacillus fermentum		20.68	0.28				
Actinomyces odontolyticus		15.09	0.18				
23715512	**Low discrimination**						**23715512**
Bifidobacterium adolescentis		87.74	0.36	NIT 1			
Actinomyces israelii		9.92	0.26	MNS 87			
Bifidobacterium longum subsp. longum		2.34	0.07				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23717000	**Low discrimination**					**23717000**
Actinomyces israelii		65.35	0.41	CEL 93		
Actinomyces odontolyticus		28.07	0.27			
Actinomyces naeslundii		6.57	0.33			
23717001	**Good identification**					**23717001**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23717002	**Low discrimination**					**23717002**
Actinomyces israelii		80.48	0.44	CEL 93		
Actinomyces odontolyticus		14.13	0.23			
Actinomyces naeslundii		5.40	0.33			
23717003	**Good identification**					**23717003**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23717021	**Good identification**					**23717021**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23717023	**Good identification**					**23717023**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23717042	**Good identification**					**23717042**
Actinomyces israelii		100.0	0.37	CEL 93		
23717100	**Low discrimination**					**23717100**
Actinomyces israelii		87.60	0.51	CEL 93		
Actinomyces naeslundii		6.27	0.41			
Propionibacterium propionicum		3.16	0.26			
23717101	**Acceptable identification**					**23717101**
Actinomyces naeslundii		91.40	0.44	MNS 88		
Actinomyces israelii		8.60	0.17	XYL 1		
23717102	**Low discrimination**					**23717102**
Actinomyces israelii		89.55	0.54	CEL 93		
Actinomyces naeslundii		4.27	0.41			
Propionibacterium propionicum		3.23	0.29			
23717103	**Low discrimination**					**23717103**
Actinomyces naeslundii		83.54	0.44	MNS 88		
Actinomyces israelii		11.79	0.20	XYL 1		
Propionibacterium propionicum		4.67	0.12			
23717110	**Good identification**					**23717110**
Actinomyces israelii		100.0	0.37	CEL 93		
23717112	**Low discrimination**					**23717112**
Actinomyces israelii		86.50	0.40	CEL 93		
Bifidobacterium adolescentis		13.50	0.20	MLZ 13		
23717120	**Low discrimination**					**23717120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		52.86	0.41	MNS 88		
Actinomyces israelii		47.14	0.31	XYL 1		
23717121	**Good identification**					**23717121**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23717123	**Good identification**					**23717123**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23717140	**Low discrimination**					**23717140**
Actinomyces israelii		88.88	0.44	CEL 93		
Propionibacterium propionicum		8.23	0.26			
Actinomyces naeslundii		2.89	0.28			
23717142	**Low discrimination**					**23717142**
Actinomyces israelii		89.74	0.47	CEL 93		
Propionibacterium propionicum		8.31	0.29			
Actinomyces naeslundii		1.94	0.28			
23717302	**Low discrimination**					**23717302**
Actinomyces israelii		73.67	0.35	CEL 93		
Actinomyces odontolyticus		26.33	0.20	RHA 7		
23717400	**Good identification**					**23717400**
Actinomyces israelii		100.0	0.40	CEL 93		
23717402	**Good identification**					**23717402**
Actinomyces israelii		100.0	0.43	CEL 93		
23717442	**Good identification**					**23717442**
Actinomyces israelii		100.0	0.36	CEL 93		
23717500	**Good identification**					**23717500**
Actinomyces israelii		100.0	0.50	CEL 93		
23717502	**Low discrimination**					**23717502**
Actinomyces israelii		86.50	0.53	CEL 93		
Bifidobacterium adolescentis		13.50	0.33			
23717510	**Good identification**					**23717510**
Actinomyces israelii		100.0	0.36	CEL 93		
23717540	**Good identification**					**23717540**
Actinomyces israelii		100.0	0.43	CEL 93		
23717542	**Good identification**					**23717542**
Actinomyces israelii		95.27	0.46	CEL 93		
Bifidobacterium adolescentis		4.73	0.17			
23721000	**Low discrimination**					**23721000**
Actinomyces naeslundii		87.36	0.46	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		12.64	0.14	FRU 95		
23721001	**Good identification**					**23721001**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23721002	**Low discrimination**					**23721002**
Actinomyces naeslundii		59.33	0.46	MNS 88		
Actinomyces viscosus		35.41	0.18	FRU 95		
Actinomyces odontolyticus		5.26	0.10			
23721003	**Low discrimination**					**23721003**
Actinomyces naeslundii		88.29	0.49	MNS 88		
Actinomyces viscosus		11.71	0.09	FRU 95		
23721020	**Good identification**					**23721020**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23721021	**Good identification**					**23721021**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23721022	**Good identification**					**23721022**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23721023	**Good identification**					**23721023**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23721041	**Good identification**					**23721041**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23721043	**Good identification**					**23721043**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23721061	**Good identification**					**23721061**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23721063	**Good identification**					**23721063**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23721100	**Good identification**					**23721100**
Actinomyces naeslundii		98.42	0.53	MNS 88		
Actinomyces odontolyticus		1.58	0.05	FRU 95		
23721101	**Very good identification**					**23721101**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23721102	**Low discrimination**					**23721102**
Actinomyces naeslundii		72.84	0.53	MNS 88		
Actinomyces viscosus		27.16	0.22	FRU 95		
23721103	**Good identification**					**23721103**
Actinomyces naeslundii		92.35	0.56	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		7.65	0.14	FRU 95		
23721120	**Good identification**					**23721120**
Actinomyces naeslundii		98.37	0.53	MNS 88		
Clostridium perfringens		1.63	0.01	FRU 95		
23721121	**Very good identification**					**23721121**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23721122	**Good identification**					**23721122**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23721123	**Very good identification**					**23721123**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23721140	**Good identification**					**23721140**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23721141	**Good identification**					**23721141**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23721142	**Good identification**					**23721142**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23721143	**Good identification**					**23721143**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23721160	**Good identification**					**23721160**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23721161	**Good identification**					**23721161**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23721162	**Good identification**					**23721162**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23721163	**Good identification**					**23721163**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23722101	**Good identification**					**23722101**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23722103	**Good identification**					**23722103**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23722121	**Good identification**					**23722121**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23722123	**Good identification**					**23722123**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23723000	**Good identification**					**23723000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MNS 88		
23723001	**Good identification**					**23723001**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23723002	**Good identification**					**23723002**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23723003	**Good identification**					**23723003**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23723020	**Good identification**					**23723020**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23723021	**Good identification**					**23723021**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23723022	**Good identification**					**23723022**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23723023	**Good identification**					**23723023**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23723040	**Good identification**					**23723040**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23723041	**Good identification**					**23723041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23723042	**Good identification**					**23723042**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23723043	**Good identification**					**23723043**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23723060	**Good identification**					**23723060**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23723061	**Good identification**					**23723061**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23723062	**Good identification**					**23723062**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23723063	**Good identification**					**23723063**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23723100	**Very good identification**					**23723100**
Actinomyces naeslundii		96.09	0.58	MNS 88		
Propionibacterium propionicum		3.91	0.24	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23723101	**Very good identification**					**23723101**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23723102	**Good identification**					**23723102**
Actinomyces naeslundii		94.24	0.58	MNS 88		
Propionibacterium propionicum		5.76	0.27	FRU 95		
23723103	**Very good identification**					**23723103**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23723110	**Good identification**					**23723110**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23723111	**Good identification**					**23723111**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23723112	**Good identification**					**23723112**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23723113	**Good identification**					**23723113**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23723120	**Very good identification**					**23723120**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23723121	**Very good identification**					**23723121**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23723122	**Very good identification**					**23723122**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23723123	**Very good identification**					**23723123**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23723130	**Good identification**					**23723130**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23723131	**Good identification**					**23723131**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23723132	**Good identification**					**23723132**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23723133	**Good identification**					**23723133**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23723140	**Low discrimination**					**23723140**
Actinomyces naeslundii		81.25	0.45	MNS 88		
Propionibacterium propionicum		18.75	0.24	FRU 95		
23723141	**Good identification**					**23723141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	MNS 88		
23723142	**Low discrimination**					**23723142**
Actinomyces naeslundii		74.29	0.45	MNS 88		
Propionibacterium propionicum		25.71	0.27	FRU 95		
23723143	**Good identification**					**23723143**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23723160	**Good identification**					**23723160**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23723161	**Good identification**					**23723161**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23723162	**Good identification**					**23723162**
Actinomyces naeslundii		100.0	0.45	MNS 88		
23723163	**Good identification**					**23723163**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23730000	**Low discrimination**					**23730000**
					GRA	
Actinomyces odontolyticus		66.72	0.35	RAF 3	99	
Actinomyces naeslundii		21.16	0.44	TRE 1	99	
Fusobacterium mortiferum		7.12	0.24		1	
23730001	**Good identification**					**23730001**
Actinomyces naeslundii		95.02	0.47	MNS 88		
Actinomyces viscosus		4.98	0.01	GAL 96		
23730002	**Low discrimination**					**23730002**
Actinomyces viscosus		88.85	0.43	MNS 96		
Actinomyces odontolyticus		7.35	0.31	TRE 7		
Actinomyces naeslundii		3.80	0.44			
23730020	**Low discrimination**					**23730020**
Actinomyces naeslundii		66.08	0.44	MNS 88		
Actinomyces odontolyticus		33.92	0.22	GAL 96		
23730021	**Good identification**					**23730021**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23730022	**Low discrimination**					**23730022**
Actinomyces naeslundii		64.49	0.44	MNS 88		
Actinomyces odontolyticus		20.29	0.18	GAL 96		
Actinomyces viscosus		15.23	0.09			
23730023	**Good identification**					**23730023**
Actinomyces naeslundii		95.02	0.47	MNS 88		
Actinomyces viscosus		4.98	0.01	GAL 96		
23730100	**Low discrimination**					**23730100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		53.40	0.52	MNS 88	99	
Actinomyces odontolyticus		18.68	0.27	GAL 96	99	
Fusobacterium mortiferum		14.72	0.30		1	
23730101	**Good identification**					**23730101**
Actinomyces naeslundii		96.83	0.55	MNS 88		
Actinomyces viscosus		3.17	0.05	GAL 96		
23730102	**Acceptable identification**					**23730102**
Actinomyces viscosus		91.28	0.47	MNS 96		
Actinomyces naeslundii		6.25	0.52	TRE 7		
Actinomyces odontolyticus		1.34	0.23			
23730103	**Low discrimination**					**23730103**
Actinomyces viscosus		76.44	0.39	MNS 96		
Actinomyces naeslundii		23.56	0.55	TRE 7		
23730120	**Acceptable identification**					**23730120**
Actinomyces naeslundii		92.77	0.52	MNS 88		
Actinomyces odontolyticus		5.28	0.13	GAL 96		
Clostridium perfringens		1.94	0.01			
23730121	**Good identification**					**23730121**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23730122	**Low discrimination**					**23730122**
Actinomyces naeslundii		84.58	0.52	MNS 88		
Actinomyces viscosus		12.47	0.13	GAL 96		
Actinomyces odontolyticus		2.95	0.09			
23730123	**Good identification**					**23730123**
Actinomyces naeslundii		96.83	0.55	MNS 88		
Actinomyces viscosus		3.17	0.05	GAL 96		
23730140	**Low discrimination**					**23730140**
Actinomyces naeslundii		63.87	0.39	MNS 88		
Propionibacterium propionicum		36.13	0.24	GAL 96		
23730141	**Good identification**					**23730141**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23730142	**Low discrimination**					**23730142**
Actinomyces naeslundii		37.26	0.39	MNS 88		
Propionibacterium propionicum		31.62	0.27	GAL 96		
Actinomyces viscosus		31.13	0.13			
23730143	**Low discrimination**					**23730143**
Actinomyces naeslundii		84.34	0.42	MNS 88		
Actinomyces viscosus		15.66	0.05	GAL 96		
23730160	**Good identification**					**23730160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	MNS 88		
23730161	**Good identification**					**23730161**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23730162	**Good identification**					**23730162**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23730163	**Good identification**					**23730163**
Actinomyces naeslundii		100.0	0.42	MNS 88		
23730200	**Good identification**					**23730200**
Actinomyces odontolyticus		100.0	0.40	RAF 3		
23730202	**Acceptable identification**					**23730202**
Actinomyces odontolyticus		94.08	0.36	RAF 3		
Actinomyces viscosus		5.92	0.09	TRE 1		
23731000	**Low discrimination**					**23731000**
					GRA	
Actinomyces naeslundii		89.25	0.67	MNS 88	99	
Actinomyces odontolyticus		7.82	0.32		99	
Fusobacterium mortiferum		1.88	0.27		1	
23731001	**Very good identification**					**23731001**
Actinomyces naeslundii		100.0	0.70	MNS 88		
23731002	**Low discrimination**					**23731002**
Actinomyces naeslundii		60.19	0.67	MNS 88		
Actinomyces viscosus		35.92	0.39			
Actinomyces odontolyticus		3.23	0.28			
23731003	**Low discrimination**					**23731003**
Actinomyces naeslundii		88.29	0.70	MNS 88		
Actinomyces viscosus		11.71	0.31			
23731010	**Good identification**					**23731010**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23731011	**Good identification**					**23731011**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23731012	**Low discrimination**					**23731012**
Actinomyces naeslundii		89.72	0.46	MNS 88		
Actinomyces viscosus		10.28	0.05	MLZ 5		
23731013	**Good identification**					**23731013**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23731020	**Very good identification**					**23731020**
Actinomyces naeslundii		95.44	0.67	MNS 88		
Clostridium perfringens		3.20	0.20			
Actinomyces odontolyticus		1.36	0.19			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23731021	**Very good identification**					**23731021**
Actinomyces naeslundii		100.0	0.70	MNS 88		
23731022	**Very good identification**					**23731022**
Actinomyces naeslundii		97.04	0.67	MNS 88		
Clostridium perfringens		1.53	0.14			
Actinomyces odontolyticus		0.85	0.15			
23731023	**Very good identification**					**23731023**
Actinomyces naeslundii		100.0	0.70	MNS 88		
23731030	**Good identification**					**23731030**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23731031	**Good identification**					**23731031**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23731032	**Good identification**					**23731032**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23731033	**Good identification**					**23731033**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23731040	**Good identification**					**23731040**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23731041	**Very good identification**					**23731041**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23731042	**Acceptable identification**					**23731042**
Actinomyces naeslundii		94.94	0.55	MNS 88		
Actinomyces viscosus		3.24	0.05			
Actinomyces israelii		1.82	0.16			
23731043	**Very good identification**					**23731043**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23731051	**Good identification**					**23731051**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731053	**Good identification**					**23731053**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731060	**Good identification**					**23731060**
Actinomyces naeslundii		96.51	0.55	MNS 88		
Clostridium perfringens		3.49	0.08			
23731061	**Very good identification**					**23731061**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23731062	**Good identification**					**23731062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.33	0.55	MNS 88		
Clostridium perfringens		1.68	0.02			
23731063	**Very good identification**					**23731063**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23731071	**Good identification**					**23731071**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731073	**Good identification**					**23731073**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731100	**Very good identification**					**23731100**
					GRA	
Actinomyces naeslundii		95.63	0.75	MNS 88	99	
Fusobacterium mortiferum		1.65	0.33		1	
Actinomyces odontolyticus		0.93	0.24		99	
23731101	**Very good identification**					**23731101**
Actinomyces naeslundii		99.92	0.78	MNS 88		
Actinomyces viscosus		0.08	0.01			
23731102	**Low discrimination**					**23731102**
Actinomyces naeslundii		71.41	0.75	MNS 88		
Actinomyces viscosus		26.62	0.44			
Actinomyces israelii		0.87	0.33			
23731103	**Good identification**					**23731103**
Actinomyces naeslundii		92.35	0.78	MNS 88		
Actinomyces viscosus		7.65	0.36			
23731110	**Good identification**					**23731110**
Actinomyces naeslundii		97.74	0.53	MNS 88		
Actinomyces israelii		2.26	0.16	MLZ 5		
23731111	**Very good identification**					**23731111**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23731112	**Acceptable identification**					**23731112**
Actinomyces naeslundii		90.40	0.53	MNS 88		
Actinomyces viscosus		6.47	0.10	MLZ 5		
Actinomyces israelii		3.14	0.19			
23731113	**Very good identification**					**23731113**
Actinomyces naeslundii		98.44	0.56	MNS 88		
Actinomyces viscosus		1.57	0.01	MLZ 5		
23731120	**Good identification**					**23731120**
Actinomyces naeslundii		91.92	0.75	MNS 88		
Clostridium perfringens		7.94	0.35			
Actinomyces odontolyticus		0.15	0.10			
23731121	**Very good identification**					**23731121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.94	0.78	MNS 88		
Clostridium perfringens		0.06	0.01			
23731122	**Very good identification**					**23731122**
Actinomyces naeslundii		95.66	0.75	MNS 88		
Clostridium perfringens		3.89	0.29			
Actinomyces viscosus		0.36	0.10			
23731123	**Very good identification**					**23731123**
Actinomyces naeslundii		99.92	0.78	MNS 88		
Actinomyces viscosus		0.08	0.01			
23731130	**Good identification**					**23731130**
Actinomyces naeslundii		98.37	0.53	MNS 88		
Clostridium perfringens		1.63	0.01	MLZ 5		
23731131	**Very good identification**					**23731131**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23731132	**Good identification**					**23731132**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23731133	**Very good identification**					**23731133**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23731140	**Very good identification**					**23731140**
Actinomyces naeslundii		96.01	0.62	MNS 88		
Propionibacterium propionicum		2.26	0.24			
Actinomyces israelii		1.72	0.23			
23731141	**Very good identification**					**23731141**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23731142	**Good identification**					**23731142**
Actinomyces naeslundii		92.28	0.62	MNS 88		
Propionibacterium propionicum		3.26	0.27			
Actinomyces israelii		2.49	0.26			
23731143	**Very good identification**					**23731143**
Actinomyces naeslundii		99.53	0.65	MNS 88		
Actinomyces viscosus		0.47	0.01			
23731150	**Good identification**					**23731150**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23731151	**Good identification**					**23731151**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23731152	**Good identification**					**23731152**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23731153	**Good identification**					**23731153**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	MNS 88		
23731160	**Good identification**					**23731160**
Actinomyces naeslundii		91.47	0.62	MNS 88		
Clostridium perfringens		8.53	0.22			
23731161	**Very good identification**					**23731161**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23731162	**Very good identification**					**23731162**
Actinomyces naeslundii		95.80	0.62	MNS 88		
Clostridium perfringens		4.20	0.17			
23731163	**Very good identification**					**23731163**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23731170	**Good identification**					**23731170**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23731171	**Good identification**					**23731171**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23731172	**Good identification**					**23731172**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23731173	**Good identification**					**23731173**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23731200	**Acceptable identification**					**23731200**
Actinomyces odontolyticus		94.39	0.37	RAF 3		
Actinomyces naeslundii		5.61	0.33	TRE 1		
23731201	**Good identification**					**23731201**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731203	**Good identification**					**23731203**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731221	**Good identification**					**23731221**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731223	**Good identification**					**23731223**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731301	**Good identification**					**23731301**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23731303	**Acceptable identification**					**23731303**
Actinomyces naeslundii		92.35	0.44	MNS 88		
Actinomyces viscosus		7.65	0.01	RHA 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23731320	**Low discrimination**					**23731320**
Actinomyces naeslundii		71.93	0.41	MNS 88		
Actinomyces odontolyticus		21.86	0.15	RHA 1		
Clostridium perfringens		6.21	0.01			
23731321	**Good identification**					**23731321**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23731322	**Low discrimination**					**23731322**
					GRA	
Actinomyces naeslundii		74.41	0.41	MNS 88	99	
Actinomyces odontolyticus		13.86	0.11	RHA 1	99	
Parabacteroides distasonis		11.73	0.14		1	
23731323	**Good identification**					**23731323**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23731401	**Good identification**					**23731401**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731403	**Good identification**					**23731403**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731421	**Good identification**					**23731421**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731423	**Good identification**					**23731423**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23731500	**Low discrimination**					**23731500**
Actinomyces naeslundii		60.25	0.41	MNS 88		
Actinomyces israelii		39.75	0.29	ARA 1		
23731501	**Good identification**					**23731501**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23731502	**Low discrimination**					**23731502**
Actinomyces naeslundii		37.20	0.41	MNS 88		
Actinomyces israelii		36.82	0.32	ARA 1		
Actinomyces viscosus		13.87	0.10			
23731503	**Acceptable identification**					**23731503**
Actinomyces naeslundii		92.35	0.44	MNS 88		
Actinomyces viscosus		7.65	0.01	ARA 1		
23731520	**Acceptable identification**					**23731520**
Actinomyces naeslundii		92.05	0.41	MNS 88		
Clostridium perfringens		7.95	0.01	ARA 1		
23731521	**Good identification**					**23731521**
Actinomyces naeslundii		100.0	0.44	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23731522	**Good identification**					**23731522**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23731523	**Good identification**					**23731523**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23732000	**Low discrimination**					**23732000**
Actinomyces naeslundii		87.23	0.49	MNS 88		
Propionibacterium propionicum		12.77	0.24	GAL 96		
23732001	**Good identification**					**23732001**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23732002	**Low discrimination**					**23732002**
Actinomyces naeslundii		74.56	0.49	MNS 88		
Propionibacterium propionicum		16.37	0.27	GAL 96		
Actinomyces viscosus		9.07	0.09			
23732003	**Good identification**					**23732003**
Actinomyces naeslundii		97.37	0.52	MNS 88		
Actinomyces viscosus		2.63	0.01	GAL 96		
23732020	**Good identification**					**23732020**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23732021	**Good identification**					**23732021**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23732022	**Good identification**					**23732022**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23732023	**Good identification**					**23732023**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23732040	**Low discrimination**					**23732040**
Actinomyces naeslundii		54.66	0.36	MNS 88		
Propionibacterium propionicum		45.34	0.24	GAL 96		
23732041	**Good identification**					**23732041**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23732043	**Good identification**					**23732043**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23732060	**Good identification**					**23732060**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23732061	**Good identification**					**23732061**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23732062	**Good identification**					**23732062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	MNS 88		
23732063	**Good identification**					**23732063**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23732100	**Low discrimination**					**23732100**
Propionibacterium propionicum		83.59	0.58	SAL 1		
Actinomyces naeslundii		16.42	0.57			
23732101	**Low discrimination**					**23732101**
Actinomyces naeslundii		72.61	0.60	MNS 88		
Propionibacterium propionicum		27.39	0.42	GAL 96		
23732102	**Low discrimination**					**23732102**
Propionibacterium propionicum		87.65	0.61	SAL 1		
Actinomyces naeslundii		11.48	0.57			
Actinomyces viscosus		0.87	0.13			
23732103	**Low discrimination**					**23732103**
Actinomyces naeslundii		63.19	0.60	MNS 88		
Propionibacterium propionicum		35.75	0.45	GAL 96		
Actinomyces viscosus		1.07	0.05			
23732111	**Good identification**					**23732111**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23732113	**Good identification**					**23732113**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23732120	**Very good identification**					**23732120**
Actinomyces naeslundii		95.11	0.57	MNS 88		
Propionibacterium propionicum		4.89	0.24	GAL 96		
23732121	**Very good identification**					**23732121**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23732122	**Good identification**					**23732122**
Actinomyces naeslundii		92.84	0.57	MNS 88		
Propionibacterium propionicum		7.16	0.27	GAL 96		
23732123	**Very good identification**					**23732123**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23732131	**Good identification**					**23732131**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23732133	**Good identification**					**23732133**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23732140	**Very good identification**					**23732140**
Propionibacterium propionicum		96.65	0.58	SAL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		3.35	0.44			
23732141	**Low discrimination**					**23732141**
Propionibacterium propionicum		68.13	0.42	SAL 1		
Actinomyces naeslundii		31.87	0.47	URE 10		
23732142	**Very good identification**					**23732142**
Propionibacterium propionicum		97.74	0.61	SAL 1		
Actinomyces naeslundii		2.26	0.44			
23732143	**Low discrimination**					**23732143**
Propionibacterium propionicum		76.23	0.45	SAL 1		
Actinomyces naeslundii		23.78	0.47	URE 10		
23732160	**Low discrimination**					**23732160**
Actinomyces naeslundii		77.43	0.44	MNS 88		
Propionibacterium propionicum		22.57	0.24	GAL 96		
23732161	**Good identification**					**23732161**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23732162	**Low discrimination**					**23732162**
Actinomyces naeslundii		69.58	0.44	MNS 88		
Propionibacterium propionicum		30.42	0.27	GAL 96		
23732163	**Good identification**					**23732163**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23733000	**Very good identification**					**23733000**
Actinomyces naeslundii		97.75	0.72	MNS 88		
Actinomyces israelii		1.65	0.33			
Propionibacterium propionicum		0.60	0.24			
23733001	**Very good identification**					**23733001**
Actinomyces naeslundii		100.0	0.75	MNS 88		
23733002	**Very good identification**					**23733002**
Actinomyces naeslundii		96.37	0.72	MNS 88		
Actinomyces israelii		2.45	0.36			
Propionibacterium propionicum		0.88	0.27			
23733003	**Very good identification**					**23733003**
Actinomyces naeslundii		100.0	0.75	MNS 88		
23733010	**Good identification**					**23733010**
Actinomyces naeslundii		95.42	0.50	MNS 88		
Actinomyces israelii		4.58	0.19	MLZ 5		
23733011	**Good identification**					**23733011**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23733012	**Acceptable identification**					**23733012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		93.28	0.50	MNS 88		
Actinomyces israelii		6.72	0.22	MLZ 5		
23733013	**Good identification**					**23733013**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23733020	**Very good identification**					**23733020**
Actinomyces naeslundii		100.0	0.72	MNS 88		
23733021	**Very good identification**					**23733021**
Actinomyces naeslundii		100.0	0.75	MNS 88		
23733022	**Very good identification**					**23733022**
Actinomyces naeslundii		99.84	0.72	MNS 88		
Actinomyces israelii		0.16	0.15			
23733023	**Very good identification**					**23733023**
Actinomyces naeslundii		100.0	0.75	MNS 88		
23733030	**Good identification**					**23733030**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23733031	**Good identification**					**23733031**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23733032	**Good identification**					**23733032**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23733033	**Good identification**					**23733033**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23733040	**Good identification**					**23733040**
Actinomyces naeslundii		93.30	0.59	MNS 88		
Actinomyces israelii		3.48	0.26			
Propionibacterium propionicum		3.23	0.24			
23733041	**Very good identification**					**23733041**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23733042	**Good identification**					**23733042**
Actinomyces naeslundii		90.27	0.59	MNS 88		
Actinomyces israelii		5.05	0.29			
Propionibacterium propionicum		4.68	0.27			
23733043	**Very good identification**					**23733043**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23733050	**Good identification**					**23733050**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23733051	**Good identification**					**23733051**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733052	**Low discrimination**					**23733052**
Actinomyces naeslundii		86.30	0.38	MNS 88		
Actinomyces israelii		13.71	0.15	MLZ 5		
23733053	**Good identification**					**23733053**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733060	**Very good identification**					**23733060**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23733061	**Very good identification**					**23733061**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23733062	**Very good identification**					**23733062**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23733063	**Very good identification**					**23733063**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23733070	**Good identification**					**23733070**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23733071	**Good identification**					**23733071**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733072	**Good identification**					**23733072**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23733073	**Good identification**					**23733073**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733100	**Low discrimination**					**23733100**
Actinomyces naeslundii		80.91	0.80	MNS 88		
Propionibacterium propionicum		17.17	0.58			
Actinomyces israelii		1.92	0.43			
23733101	**Very good identification**					**23733101**
Actinomyces naeslundii		98.45	0.83	MNS 88		
Propionibacterium propionicum		1.55	0.42			
23733102	**Low discrimination**					**23733102**
Actinomyces naeslundii		73.75	0.80	MNS 88		
Propionibacterium propionicum		23.47	0.61			
Actinomyces israelii		2.63	0.46			
23733103	**Very good identification**					**23733103**
Actinomyces naeslundii		97.66	0.83	MNS 88		
Propionibacterium propionicum		2.30	0.45			
Actinomyces viscosus		0.04	0.01			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23733110	**Good identification**					**23733110**
Actinomyces naeslundii		90.23	0.58	MNS 88		
Actinomyces israelii		6.09	0.29	MLZ 5		
Propionibacterium propionicum		3.67	0.24			
23733111	**Very good identification**					**23733111**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23733112	**Low discrimination**					**23733112**
Actinomyces naeslundii		86.03	0.58	MNS 88		
Actinomyces israelii		8.71	0.32	MLZ 5		
Propionibacterium propionicum		5.26	0.27			
23733113	**Very good identification**					**23733113**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23733120	**Very good identification**					**23733120**
Actinomyces naeslundii		99.59	0.80	MNS 88		
Propionibacterium propionicum		0.21	0.24			
Actinomyces israelii		0.15	0.23			
23733121	**Very good identification**					**23733121**
Actinomyces naeslundii		100.0	0.83	MNS 88		
23733122	**Very good identification**					**23733122**
Actinomyces naeslundii		99.45	0.80	MNS 88		
Propionibacterium propionicum		0.32	0.27			
Actinomyces israelii		0.23	0.26			
23733123	**Very good identification**					**23733123**
Actinomyces naeslundii		100.0	0.83	MNS 88		
23733130	**Very good identification**					**23733130**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23733131	**Very good identification**					**23733131**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23733132	**Very good identification**					**23733132**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23733133	**Very good identification**					**23733133**
Actinomyces naeslundii		100.0	0.61	MNS 88		
23733140	**Low discrimination**					**23733140**
Propionibacterium propionicum		53.32	0.58	SAL 1		
Actinomyces naeslundii		44.35	0.67			
Actinomyces israelii		2.33	0.36			
23733141	**Good identification**					**23733141**
Actinomyces naeslundii		91.82	0.70	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Propionibacterium propionicum	8.18	0.42			
23733142	**Low discrimination**					**23733142**
	Propionibacterium propionicum	62.57	0.61	SAL 1		
	Actinomyces naeslundii	34.70	0.67			
	Actinomyces israelii	2.73	0.39			
23733143	**Low discrimination**					**23733143**
	Actinomyces naeslundii	88.22	0.70	MNS 88		
	Propionibacterium propionicum	11.78	0.45			
23733150	**Low discrimination**					**23733150**
	Actinomyces naeslundii	72.49	0.45	MNS 88		
	Propionibacterium propionicum	16.73	0.24	MLZ 5		
	Actinomyces israelii	10.79	0.22			
23733151	**Good identification**					**23733151**
	Actinomyces naeslundii	100.0	0.48	MNS 88		
23733152	**Low discrimination**					**23733152**
	Actinomyces naeslundii	63.72	0.45	MNS 88		
	Propionibacterium propionicum	22.06	0.27	MLZ 5		
	Actinomyces israelii	14.22	0.25			
23733153	**Good identification**					**23733153**
	Actinomyces naeslundii	100.0	0.48	MNS 88		
23733160	**Very good identification**					**23733160**
	Actinomyces naeslundii	98.48	0.67	MNS 88		
	Propionibacterium propionicum	1.20	0.24			
	Actinomyces israelii	0.33	0.16			
23733161	**Very good identification**					**23733161**
	Actinomyces naeslundii	100.0	0.70	MNS 88		
23733162	**Very good identification**					**23733162**
	Actinomyces naeslundii	97.73	0.67	MNS 88		
	Propionibacterium propionicum	1.78	0.27			
	Actinomyces israelii	0.49	0.19			
23733163	**Very good identification**					**23733163**
	Actinomyces naeslundii	100.0	0.70	MNS 88		
23733170	**Good identification**					**23733170**
	Actinomyces naeslundii	100.0	0.45	MNS 88		
23733171	**Good identification**					**23733171**
	Actinomyces naeslundii	100.0	0.48	MNS 88		
23733172	**Good identification**					**23733172**
	Actinomyces naeslundii	100.0	0.45	MNS 88		
23733173	**Good identification**					**23733173**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	MNS 88		
23733200	**Low discrimination**					**23733200**
Actinomyces naeslundii		88.80	0.38	MNS 88		
Actinomyces israelii		11.20	0.14	RHA 1		
23733201	**Good identification**					**23733201**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733202	**Low discrimination**					**23733202**
Actinomyces naeslundii		84.09	0.38	MNS 88		
Actinomyces israelii		15.91	0.17	RHA 1		
23733203	**Good identification**					**23733203**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733220	**Good identification**					**23733220**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23733221	**Good identification**					**23733221**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733222	**Good identification**					**23733222**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23733223	**Good identification**					**23733223**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733300	**Low discrimination**					**23733300**
Actinomyces naeslundii		71.98	0.46	MNS 88		
Propionibacterium propionicum		15.27	0.24	RHA 1		
Actinomyces israelii		12.76	0.24			
23733301	**Good identification**					**23733301**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733302	**Low discrimination**					**23733302**
Actinomyces naeslundii		63.13	0.46	MNS 88		
Propionibacterium propionicum		20.09	0.27	RHA 1		
Actinomyces israelii		16.78	0.27			
23733303	**Good identification**					**23733303**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733320	**Good identification**					**23733320**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23733321	**Good identification**					**23733321**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733322	**Good identification**					**23733322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MNS 88		
23733323	**Good identification**					**23733323**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733341	**Good identification**					**23733341**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23733343	**Good identification**					**23733343**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23733361	**Good identification**					**23733361**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23733363	**Good identification**					**23733363**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23733401	**Good identification**					**23733401**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733403	**Good identification**					**23733403**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733420	**Good identification**					**23733420**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23733421	**Good identification**					**23733421**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733422	**Low discrimination**					**23733422**
Actinomyces naeslundii		88.40	0.38	MNS 88		
Actinomyces israelii		11.60	0.14	ARA 1		
23733423	**Good identification**					**23733423**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23733500	**Low discrimination**					**23733500**
Actinomyces israelii		61.38	0.42	CEL 93		
Actinomyces naeslundii		31.86	0.46	XYL 92		
Propionibacterium propionicum		6.76	0.24			
23733501	**Good identification**					**23733501**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733502	**Low discrimination**					**23733502**
Actinomyces israelii		66.50	0.45	CEL 93		
Actinomyces naeslundii		23.01	0.46	XYL 92		
Propionibacterium propionicum		7.32	0.27			
23733503	**Good identification**					**23733503**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733520	**Low discrimination**					**23733520**
Actinomyces naeslundii		89.05	0.46	MNS 88		
Actinomyces israelii		10.95	0.21	ARA 1		
23733521	**Good identification**					**23733521**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733522	**Low discrimination**					**23733522**
Actinomyces naeslundii		84.43	0.46	MNS 88		
Actinomyces israelii		15.57	0.24	ARA 1		
23733523	**Good identification**					**23733523**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23733541	**Good identification**					**23733541**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23733542	**Low discrimination**					**23733542**
Actinomyces israelii		69.43	0.38	CEL 93		
Propionibacterium propionicum		19.66	0.27	XYL 92		
Actinomyces naeslundii		10.90	0.33			
23733543	**Good identification**					**23733543**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23733561	**Good identification**					**23733561**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23733563	**Good identification**					**23733563**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23735000	**Low discrimination**					**23735000**
Actinomyces israelii		55.60	0.38	CEL 93		
Actinomyces odontolyticus		35.98	0.27			
Actinomyces naeslundii		8.43	0.33			
23735001	**Good identification**					**23735001**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23735002	**Low discrimination**					**23735002**
Actinomyces israelii		70.14	0.41	CEL 93		
Actinomyces odontolyticus		18.54	0.23			
Actinomyces naeslundii		7.09	0.33			
23735003	**Good identification**					**23735003**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23735021	**Good identification**					**23735021**
Actinomyces naeslundii		100.0	0.36	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
23735023	**Good identification**					**23735023**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23735100	**Low discrimination**					**23735100**
Actinomyces israelii		85.27	0.48	CEL 93		
Actinomyces naeslundii		9.19	0.41			
Actinomyces odontolyticus		4.36	0.18			
23735101	**Acceptable identification**					**23735101**
Actinomyces naeslundii		94.12	0.44	MNS 88		
Actinomyces israelii		5.88	0.14	XYL 1		
23735102	**Low discrimination**					**23735102**
Actinomyces israelii		87.49	0.51	CEL 93		
Actinomyces naeslundii		6.29	0.41			
Actinomyces viscosus		2.34	0.10			
23735103	**Low discrimination**					**23735103**
Actinomyces naeslundii		85.00	0.44	MNS 88		
Actinomyces israelii		7.96	0.17	XYL 1		
Actinomyces viscosus		7.04	0.01			
23735112	**Low discrimination**					**23735112**
Actinomyces israelii		83.92	0.37	CEL 93		
Bifidobacterium adolescentis		16.08	0.19	MLZ 13		
23735120	**Low discrimination**					**23735120**
Actinomyces naeslundii		59.58	0.41	MNS 88		
Actinomyces israelii		35.28	0.28	XYL 1		
Clostridium perfringens		5.15	0.01			
23735121	**Good identification**					**23735121**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23735122	**Low discrimination**					**23735122**
					GRA	
Actinomyces naeslundii		42.36	0.41	MNS 88	99	
Actinomyces israelii		37.61	0.31	XYL 1	99	
Parabacteroides distasonis		20.03	0.22		1	
23735123	**Good identification**					**23735123**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23735140	**Good identification**					**23735140**
Actinomyces israelii		95.34	0.41	CEL 93		
Actinomyces naeslundii		4.66	0.28			
23735142	**Good identification**					**23735142**
Actinomyces israelii		96.84	0.44	CEL 93		
Actinomyces naeslundii		3.16	0.28			
23735400	**Good identification**					**23735400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.37	CEL 93		
23735402	**Good identification**					**23735402**
Actinomyces israelii		100.0	0.40	CEL 93		
23735500	**Good identification**					**23735500**
Actinomyces israelii		100.0	0.47	CEL 93		
23735502	**Low discrimination**					**23735502**
Actinomyces israelii		83.92	0.50	CEL 93		
Bifidobacterium adolescentis		16.08	0.31			
23735540	**Good identification**					**23735540**
Actinomyces israelii		100.0	0.40	CEL 93		
23735542	**Acceptable identification**					**23735542**
Actinomyces israelii		94.26	0.43	CEL 93		
Bifidobacterium adolescentis		5.74	0.16			
23737000	**Good identification**					**23737000**
Actinomyces israelii		95.07	0.51	CEL 93		
Actinomyces naeslundii		4.94	0.38			
23737001	**Low discrimination**					**23737001**
Actinomyces naeslundii		88.52	0.41	MNS 88		
Actinomyces israelii		11.48	0.17	XYL 1		
23737002	**Good identification**					**23737002**
Actinomyces israelii		96.66	0.54	CEL 93		
Actinomyces naeslundii		3.35	0.38			
23737003	**Low discrimination**					**23737003**
Actinomyces naeslundii		83.71	0.41	MNS 88		
Actinomyces israelii		16.29	0.20	XYL 1		
23737010	**Good identification**					**23737010**
Actinomyces israelii		100.0	0.37	CEL 93		
23737012	**Good identification**					**23737012**
Actinomyces israelii		100.0	0.40	CEL 93		
23737021	**Good identification**					**23737021**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23737023	**Good identification**					**23737023**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23737040	**Good identification**					**23737040**
Actinomyces israelii		97.70	0.44	CEL 93		
Actinomyces naeslundii		2.30	0.25			
23737042	**Good identification**					**23737042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		98.45	0.47	CEL 93		
Actinomyces naeslundii		1.55	0.25			
23737100	**Very good identification**					**23737100**
Actinomyces israelii		95.72	0.61	CEL 93		
Actinomyces naeslundii		3.54	0.46			
Propionibacterium propionicum		0.75	0.24			
23737101	**Low discrimination**					**23737101**
Actinomyces naeslundii		84.58	0.49	MNS 88		
Actinomyces israelii		15.42	0.27	XYL 1		
23737102	**Very good identification**					**23737102**
Actinomyces israelii		96.54	0.64	CEL 93		
Actinomyces naeslundii		2.38	0.46			
Propionibacterium propionicum		0.76	0.27			
23737103	**Low discrimination**					**23737103**
Actinomyces naeslundii		78.52	0.49	MNS 88		
Actinomyces israelii		21.48	0.30	XYL 1		
23737110	**Good identification**					**23737110**
Actinomyces israelii		98.72	0.47	CEL 93		
Actinomyces naeslundii		1.28	0.24	MLZ 13		
23737112	**Good identification**					**23737112**
Actinomyces israelii		96.49	0.50	CEL 93		
Bifidobacterium adolescentis		2.67	0.17	MLZ 13		
Actinomyces naeslundii		0.84	0.24			
23737120	**Low discrimination**					**23737120**
Actinomyces israelii		63.35	0.41	CEL 93		
Actinomyces naeslundii		36.65	0.46	NAG 6		
23737121	**Good identification**					**23737121**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23737122	**Low discrimination**					**23737122**
Actinomyces israelii		72.16	0.44	CEL 93		
Actinomyces naeslundii		27.84	0.46	NAG 6		
23737123	**Good identification**					**23737123**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23737140	**Good identification**					**23737140**
Actinomyces israelii		96.44	0.54	CEL 93		
Propionibacterium propionicum		1.94	0.24			
Actinomyces naeslundii		1.62	0.33			
23737141	**Low discrimination**					**23737141**
Actinomyces naeslundii		71.34	0.36	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		28.66	0.20	XYL 1		
23737142	**Very good identification**					**23737142**
Actinomyces israelii		96.96	0.57	CEL 93		
Propionibacterium propionicum		1.95	0.27			
Actinomyces naeslundii		1.08	0.33			
23737143	**Low discrimination**					**23737143**
Actinomyces naeslundii		62.39	0.36	MNS 88		
Actinomyces israelii		37.61	0.23	XYL 1		
23737150	**Good identification**					**23737150**
Actinomyces israelii		100.0	0.40	CEL 93		
23737152	**Good identification**					**23737152**
Actinomyces israelii		100.0	0.43	CEL 93		
23737161	**Good identification**					**23737161**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23737162	**Low discrimination**					**23737162**
Actinomyces israelii		85.10	0.37	CEL 93		
Actinomyces naeslundii		14.90	0.33	NAG 6		
23737163	**Good identification**					**23737163**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23737300	**Good identification**					**23737300**
Actinomyces israelii		100.0	0.42	CEL 93		
23737302	**Good identification**					**23737302**
Actinomyces israelii		100.0	0.45	CEL 93		
23737340	**Good identification**					**23737340**
Actinomyces israelii		100.0	0.35	CEL 93		
23737342	**Good identification**					**23737342**
Actinomyces israelii		100.0	0.38	CEL 93		
23737400	**Good identification**					**23737400**
Actinomyces israelii		100.0	0.49	CEL 93		
23737402	**Good identification**					**23737402**
Actinomyces israelii		100.0	0.53	CEL 93		
23737410	**Good identification**					**23737410**
Actinomyces israelii		100.0	0.35	CEL 93		
23737412	**Good identification**					**23737412**
Actinomyces israelii		100.0	0.38	CEL 93		
23737440	**Good identification**					**23737440**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.42	CEL 93		
23737442	**Good identification**					**23737442**
Actinomyces israelii		100.0	0.45	CEL 93		
23737500	**Very good identification**					**23737500**
Actinomyces israelii		100.0	0.60	CEL 93		
23737502	**Very good identification**					**23737502**
Actinomyces israelii		97.31	0.63	CEL 93		
Bifidobacterium adolescentis		2.69	0.30			
23737510	**Good identification**					**23737510**
Actinomyces israelii		100.0	0.46	CEL 93		
23737512	**Low discrimination**					**23737512**
Actinomyces israelii		81.55	0.49	CEL 93		
Bifidobacterium adolescentis		18.45	0.31	MLZ 13		
23737520	**Good identification**					**23737520**
Actinomyces israelii		100.0	0.39	CEL 93		
23737522	**Good identification**					**23737522**
Actinomyces israelii		100.0	0.42	CEL 93		
23737540	**Good identification**					**23737540**
Actinomyces israelii		100.0	0.53	CEL 93		
23737542	**Good identification**					**23737542**
Actinomyces israelii		99.13	0.56	CEL 93		
Bifidobacterium adolescentis		0.87	0.14			
23737550	**Good identification**					**23737550**
Actinomyces israelii		100.0	0.39	CEL 93		
23737552	**Acceptable identification**					**23737552**
Actinomyces israelii		93.29	0.42	CEL 93		
Bifidobacterium adolescentis		6.71	0.16	MLZ 13		
23737562	**Good identification**					**23737562**
Actinomyces israelii		100.0	0.35	CEL 93		
23737700	**Good identification**					**23737700**
Actinomyces israelii		100.0	0.41	CEL 93		
23737702	**Good identification**					**23737702**
Actinomyces israelii		100.0	0.44	CEL 93		
23737742	**Good identification**					**23737742**
Actinomyces israelii		100.0	0.37	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23741000	**Low discrimination**					**23741000**
Actinomyces naeslundii		65.67	0.43	MNS 88		
Eubacterium tortuosum		20.41	0.06	FRU 95		
Actinomyces odontolyticus		13.92	0.14			
23741001	**Good identification**					**23741001**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23741003	**Low discrimination**					**23741003**
Actinomyces naeslundii		59.32	0.46	MNS 88		
Actinomyces viscosus		40.69	0.19	FRU 95		
23741020	**Good identification**					**23741020**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23741021	**Good identification**					**23741021**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23741022	**Good identification**					**23741022**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23741023	**Good identification**					**23741023**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23741100	**Good identification**					**23741100**
Actinomyces naeslundii		97.70	0.50	MNS 88		
Actinomyces odontolyticus		2.30	0.05	FRU 95		
23741101	**Good identification**					**23741101**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23741103	**Low discrimination**					**23741103**
Actinomyces naeslundii		70.01	0.53	MNS 88		
Actinomyces viscosus		29.99	0.23	FRU 95		
23741120	**Good identification**					**23741120**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23741121	**Good identification**					**23741121**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23741122	**Good identification**					**23741122**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23741123	**Good identification**					**23741123**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23741140	**Good identification**					**23741140**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23741141	**Good identification**					**23741141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MNS 88		
23741142	**Good identification**					**23741142**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23741143	**Good identification**					**23741143**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23741160	**Good identification**					**23741160**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23741161	**Good identification**					**23741161**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23741162	**Good identification**					**23741162**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23741163	**Good identification**					**23741163**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23743000	**Good identification**					**23743000**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23743001	**Good identification**					**23743001**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23743002	**Good identification**					**23743002**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23743003	**Good identification**					**23743003**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23743020	**Good identification**					**23743020**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23743021	**Good identification**					**23743021**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23743022	**Good identification**					**23743022**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23743023	**Good identification**					**23743023**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23743041	**Good identification**					**23743041**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23743043	**Good identification**					**23743043**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23743061	**Good identification**					**23743061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
23743063	**Good identification**					**23743063**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23743100	**Good identification**					**23743100**
Actinomyces naeslundii		97.69	0.55	MNS 88		
Actinomyces israelii		2.31	0.18	FRU 95		
23743101	**Very good identification**					**23743101**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23743102	**Good identification**					**23743102**
Actinomyces naeslundii		96.58	0.55	MNS 88		
Actinomyces israelii		3.42	0.21	FRU 95		
23743103	**Very good identification**					**23743103**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23743111	**Good identification**					**23743111**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23743113	**Good identification**					**23743113**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23743120	**Good identification**					**23743120**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23743121	**Very good identification**					**23743121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23743122	**Good identification**					**23743122**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23743123	**Very good identification**					**23743123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23743131	**Good identification**					**23743131**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23743133	**Good identification**					**23743133**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23743140	**Good identification**					**23743140**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23743141	**Good identification**					**23743141**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23743142	**Acceptable identification**					**23743142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.76	0.43	MNS 88		
Actinomyces israelii		7.24	0.14	FRU 95		
23743143	**Good identification**					**23743143**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23743160	**Good identification**					**23743160**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23743161	**Good identification**					**23743161**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23743162	**Good identification**					**23743162**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23743163	**Good identification**					**23743163**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23747100	**Good identification**					**23747100**
Actinomyces israelii		100.0	0.37	FRU 99		
23747102	**Good identification**					**23747102**
Actinomyces israelii		100.0	0.40	FRU 99		
23747500	**Good identification**					**23747500**
Actinomyces israelii		100.0	0.35	FRU 99		
23747502	**Good identification**					**23747502**
Actinomyces israelii		100.0	0.38	FRU 99		
23750000	**Low discrimination**					**23750000**
Actinomyces odontolyticus		62.50	0.35	RAF 3		
Actinomyces viscosus		16.53	0.18	CEL 1		
Actinomyces naeslundii		13.54	0.41			
23750001	**Low discrimination**					**23750001**
Actinomyces naeslundii		78.66	0.44	MNS 88		
Actinomyces viscosus		21.34	0.10	GAL 96		
23750002	**Good identification**					**23750002**
Actinomyces viscosus		96.93	0.52	MNS 96		
Actinomyces odontolyticus		2.27	0.31			
Actinomyces naeslundii		0.80	0.41			
23750003	**Good identification**					**23750003**
Actinomyces viscosus		96.41	0.44	MNS 96		
Actinomyces naeslundii		3.59	0.44			
23750020	**Low discrimination**					**23750020**
Actinomyces naeslundii		57.09	0.41	MNS 88		
Actinomyces odontolyticus		42.91	0.22	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23750021	**Good identification**					**23750021**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23750023	**Low discrimination**					**23750023**
Actinomyces naeslundii		78.66	0.44	MNS 88		
Actinomyces viscosus		21.34	0.10	GAL 96		
23750100	**Low discrimination**					**23750100**
Actinomyces naeslundii		36.69	0.49	MNS 88		
Actinomyces viscosus		27.98	0.23	GAL 96		
Actinomyces odontolyticus		18.80	0.27			
23750101	**Low discrimination**					**23750101**
Actinomyces naeslundii		85.51	0.52	MNS 88		
Actinomyces viscosus		14.49	0.14	GAL 96		
23750102	**Very good identification**					**23750102**
Actinomyces viscosus		98.23	0.57	MNS 96		
Actinomyces naeslundii		1.30	0.49			
Actinomyces odontolyticus		0.41	0.23			
23750103	**Acceptable identification**					**23750103**
Actinomyces viscosus		94.38	0.49	MNS 96		
Actinomyces naeslundii		5.63	0.52			
23750120	**Acceptable identification**					**23750120**
Actinomyces naeslundii		92.30	0.49	MNS 88		
Actinomyces odontolyticus		7.70	0.13	GAL 96		
23750121	**Good identification**					**23750121**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23750122	**Low discrimination**					**23750122**
Actinomyces naeslundii		55.14	0.49	MNS 88		
Actinomyces viscosus		42.05	0.23	GAL 96		
Actinomyces odontolyticus		2.82	0.09			
23750123	**Low discrimination**					**23750123**
Actinomyces naeslundii		85.51	0.52	MNS 88		
Actinomyces viscosus		14.49	0.14	GAL 96		
23750140	**Good identification**					**23750140**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23750141	**Good identification**					**23750141**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23750143	**Low discrimination**					**23750143**
Actinomyces naeslundii		51.01	0.39	MNS 88		
Actinomyces viscosus		48.99	0.14	GAL 96		
23750160	**Good identification**					**23750160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	MNS 88		
23750161	**Good identification**					**23750161**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23750162	**Good identification**					**23750162**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23750163	**Good identification**					**23750163**
Actinomyces naeslundii		100.0	0.39	MNS 88		
23750200	**Good identification**					**23750200**
Actinomyces odontolyticus		100.0	0.40	RAF 3		
23750202	**Low discrimination**					**23750202**
Actinomyces odontolyticus		81.82	0.36	RAF 3		
Actinomyces viscosus		18.18	0.18	CEL 1		
23751000	**Low discrimination**					**23751000**
					GRA	
Actinomyces naeslundii		81.72	0.65	MNS 88	99	
Actinomyces odontolyticus		10.48	0.32		99	
Fusobacterium mortiferum		2.81	0.28		1	
23751001	**Very good identification**					**23751001**
Actinomyces naeslundii		99.31	0.68	MNS 88		
Actinomyces viscosus		0.69	0.06			
23751002	**Low discrimination**					**23751002**
Actinomyces viscosus		73.25	0.49	MNS 96		
Actinomyces naeslundii		23.73	0.65			
Actinomyces odontolyticus		1.87	0.28			
23751003	**Low discrimination**					**23751003**
Actinomyces naeslundii		59.32	0.68	MNS 88		
Actinomyces viscosus		40.69	0.41			
23751010	**Acceptable identification**					**23751010**
Actinomyces naeslundii		92.16	0.43	MNS 88		
Actinomyces israelii		7.84	0.15	MLZ 5		
23751011	**Good identification**					**23751011**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23751012	**Low discrimination**					**23751012**
Actinomyces naeslundii		58.14	0.43	MNS 88		
Actinomyces viscosus		34.44	0.15	MLZ 5		
Actinomyces israelii		7.42	0.18			
23751013	**Low discrimination**					**23751013**
Actinomyces naeslundii		88.37	0.46	MNS 88		
Actinomyces viscosus		11.63	0.06	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23751020	**Very good identification**					**23751020**
Actinomyces naeslundii		97.96	0.65	MNS 88		
Actinomyces odontolyticus		2.05	0.19			
23751021	**Very good identification**					**23751021**
Actinomyces naeslundii		100.0	0.68	MNS 88		
23751022	**Very good identification**					**23751022**
Actinomyces naeslundii		95.40	0.65	MNS 88		
Actinomyces viscosus		2.97	0.15			
Actinomyces odontolyticus		1.22	0.15			
23751023	**Very good identification**					**23751023**
Actinomyces naeslundii		99.31	0.68	MNS 88		
Actinomyces viscosus		0.69	0.06			
23751030	**Good identification**					**23751030**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23751031	**Good identification**					**23751031**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23751032	**Good identification**					**23751032**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23751033	**Good identification**					**23751033**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23751040	**Acceptable identification**					**23751040**
Actinomyces naeslundii		93.80	0.52	MNS 88		
Actinomyces israelii		6.20	0.22			
23751041	**Good identification**					**23751041**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23751042	**Low discrimination**					**23751042**
Actinomyces naeslundii		78.38	0.52	MNS 88		
Actinomyces viscosus		13.85	0.15			
Actinomyces israelii		7.77	0.25			
23751043	**Good identification**					**23751043**
Actinomyces naeslundii		96.22	0.55	MNS 88		
Actinomyces viscosus		3.78	0.06			
23751060	**Good identification**					**23751060**
Actinomyces naeslundii		100.0	0.52	MNS 88		
23751061	**Good identification**					**23751061**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23751062	**Good identification**					**23751062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	MNS 88		
23751063	**Good identification**					**23751063**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23751100	**Good identification**					**23751100**
					GRA	
Actinomyces naeslundii		90.58	0.72	MNS 88	99	
Actinomyces israelii		3.82	0.40		99	
Fusobacterium mortiferum		2.55	0.34		1	
23751101	**Very good identification**					**23751101**
Actinomyces naeslundii		99.57	0.75	MNS 88		
Actinomyces viscosus		0.43	0.11			
23751102	**Low discrimination**					**23751102**
Actinomyces viscosus		63.99	0.53	MNS 96		
Actinomyces naeslundii		33.19	0.72			
Actinomyces israelii		2.10	0.43			
23751103	**Low discrimination**					**23751103**
Actinomyces naeslundii		70.01	0.75	MNS 88		
Actinomyces viscosus		29.99	0.45			
23751110	**Low discrimination**					**23751110**
Actinomyces naeslundii		89.32	0.50	MNS 88		
Actinomyces israelii		10.68	0.25	MLZ 5		
23751111	**Good identification**					**23751111**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23751112	**Low discrimination**					**23751112**
Actinomyces naeslundii		62.39	0.50	MNS 88		
Actinomyces viscosus		23.09	0.19	MLZ 5		
Actinomyces israelii		11.20	0.28			
23751113	**Acceptable identification**					**23751113**
Actinomyces naeslundii		92.40	0.53	MNS 88		
Actinomyces viscosus		7.60	0.11	MLZ 5		
23751120	**Very good identification**					**23751120**
Actinomyces naeslundii		98.84	0.72	MNS 88		
Clostridium perfringens		0.28	0.06			
Actinomyces israelii		0.27	0.19			
23751121	**Very good identification**					**23751121**
Actinomyces naeslundii		100.0	0.75	MNS 88		
23751122	**Good identification**					**23751122**
					GRA	
Actinomyces naeslundii		94.35	0.72	MNS 88	99	
Prevotella oralis		2.23	0.19		1	
Actinomyces viscosus		1.84	0.19		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23751123	**Very good identification**					**23751123**
Actinomyces naeslundii		99.57	0.75	MNS 88		
Actinomyces viscosus		0.43	0.11			
23751130	**Good identification**					**23751130**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23751131	**Good identification**					**23751131**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23751132	**Good identification**					**23751132**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23751133	**Good identification**					**23751133**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23751140	**Good identification**					**23751140**
Actinomyces naeslundii		91.50	0.59	MNS 88		
Actinomyces israelii		8.50	0.33			
23751141	**Very good identification**					**23751141**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23751142	**Low discrimination**					**23751142**
Actinomyces naeslundii		80.02	0.59	MNS 88		
Actinomyces israelii		11.15	0.36			
Actinomyces viscosus		8.83	0.19			
23751143	**Very good identification**					**23751143**
Actinomyces naeslundii		97.61	0.62	MNS 88		
Actinomyces viscosus		2.39	0.11			
23751150	**Low discrimination**					**23751150**
Actinomyces naeslundii		79.14	0.38	MNS 88		
Actinomyces israelii		20.86	0.18	MLZ 5		
23751151	**Good identification**					**23751151**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751152	**Low discrimination**					**23751152**
Actinomyces naeslundii		71.66	0.38	MNS 88		
Actinomyces israelii		28.34	0.21	MLZ 5		
23751153	**Good identification**					**23751153**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751160	**Very good identification**					**23751160**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23751161	**Very good identification**					**23751161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.62	MNS 88		
23751162	**Very good identification**					**23751162**
Actinomyces naeslundii		99.12	0.59	MNS 88		
Actinomyces israelii		0.88	0.15			
23751163	**Very good identification**					**23751163**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23751170	**Good identification**					**23751170**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23751171	**Good identification**					**23751171**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751172	**Good identification**					**23751172**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23751173	**Good identification**					**23751173**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751200	**Good identification**					**23751200**
Actinomyces odontolyticus		96.10	0.37	RAF 3		
Actinomyces naeslundii		3.90	0.30	CEL 1		
23751301	**Good identification**					**23751301**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751303	**Low discrimination**					**23751303**
Actinomyces naeslundii		70.01	0.41	MNS 88		
Actinomyces viscosus		29.99	0.11	RHA 1		
23751320	**Low discrimination**					**23751320**
Actinomyces naeslundii		69.20	0.38	MNS 88		
Actinomyces odontolyticus		30.80	0.15	RHA 1		
23751321	**Good identification**					**23751321**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751323	**Good identification**					**23751323**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751500	**Low discrimination**					**23751500**
Actinomyces israelii		77.34	0.38	XYL 92		
Actinomyces naeslundii		22.66	0.38			
23751501	**Good identification**					**23751501**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751502	**Low discrimination**					**23751502**
Actinomyces israelii		53.50	0.41	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		20.15	0.19			
Bifidobacterium adolescentis		15.90	0.26			
23751503	**Low discrimination**					**23751503**
Actinomyces naeslundii		70.01	0.41	MNS 88		
Actinomyces viscosus		29.99	0.11	ARA 1		
23751520	**Low discrimination**					**23751520**
Actinomyces naeslundii		82.11	0.38	MNS 88		
Actinomyces israelii		17.89	0.18	ARA 1		
23751521	**Good identification**					**23751521**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23751522	**Low discrimination**					**23751522**
Actinomyces naeslundii		75.37	0.38	MNS 88		
Actinomyces israelii		24.63	0.21	ARA 1		
23751523	**Good identification**					**23751523**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23752000	**Good identification**					**23752000**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23752001	**Good identification**					**23752001**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23752002	**Low discrimination**					**23752002**
Actinomyces naeslundii		61.39	0.46	MNS 88		
Actinomyces viscosus		38.61	0.18	GAL 96		
23752003	**Low discrimination**					**23752003**
Actinomyces naeslundii		87.74	0.49	MNS 88		
Actinomyces viscosus		12.26	0.10	GAL 96		
23752020	**Good identification**					**23752020**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23752021	**Good identification**					**23752021**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23752022	**Good identification**					**23752022**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23752023	**Good identification**					**23752023**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23752041	**Good identification**					**23752041**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23752043	**Good identification**					**23752043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	MNS 88		
23752061	**Good identification**					**23752061**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23752063	**Good identification**					**23752063**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23752100	**Low discrimination**					**23752100**
Actinomyces naeslundii		89.14	0.54	MNS 88		
Propionibacterium propionicum		8.21	0.26	GAL 96		
Actinomyces israelii		2.66	0.18			
23752101	**Very good identification**					**23752101**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23752102	**Low discrimination**					**23752102**
Actinomyces naeslundii		63.47	0.54	MNS 88		
Actinomyces viscosus		24.93	0.23	GAL 96		
Propionibacterium propionicum		8.76	0.29			
23752103	**Good identification**					**23752103**
Actinomyces naeslundii		91.11	0.57	MNS 88		
Actinomyces viscosus		7.95	0.14	GAL 96		
Propionibacterium propionicum		0.93	0.12			
23752120	**Good identification**					**23752120**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23752121	**Very good identification**					**23752121**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23752122	**Good identification**					**23752122**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23752123	**Very good identification**					**23752123**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23752140	**Low discrimination**					**23752140**
Actinomyces naeslundii		65.72	0.41	MNS 88		
Propionibacterium propionicum		34.28	0.26	GAL 96		
23752141	**Good identification**					**23752141**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23752142	**Low discrimination**					**23752142**
Actinomyces naeslundii		53.16	0.41	MNS 88		
Propionibacterium propionicum		41.60	0.29	GAL 96		
Actinomyces israelii		5.24	0.14			
23752143	**Acceptable identification**					**23752143**
Actinomyces naeslundii		94.52	0.44	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Propionibacterium propionicum	5.48	0.12	GAL 96		
23752160	**Good identification**					**23752160**
	Actinomyces naeslundii	100.0	0.41	MNS 88		
23752161	**Good identification**					**23752161**
	Actinomyces naeslundii	100.0	0.44	MNS 88		
23752162	**Good identification**					**23752162**
	Actinomyces naeslundii	100.0	0.41	MNS 88		
23752163	**Good identification**					**23752163**
	Actinomyces naeslundii	100.0	0.44	MNS 88		
23753000	**Good identification**					**23753000**
	Actinomyces naeslundii	91.95	0.70	MNS 88		
	Actinomyces israelii	8.05	0.42			
23753001	**Very good identification**					**23753001**
	Actinomyces naeslundii	100.0	0.73	MNS 88		
23753002	**Low discrimination**					**23753002**
	Actinomyces naeslundii	87.16	0.70	MNS 88		
	Actinomyces israelii	11.44	0.45			
	Actinomyces viscosus	1.40	0.15			
23753003	**Very good identification**					**23753003**
	Actinomyces naeslundii	99.64	0.73	MNS 88		
	Actinomyces viscosus	0.36	0.06			
23753010	**Low discrimination**					**23753010**
	Actinomyces naeslundii	80.10	0.48	MNS 88		
	Actinomyces israelii	19.90	0.28	MLZ 5		
23753011	**Good identification**					**23753011**
	Actinomyces naeslundii	100.0	0.51	MNS 88		
23753012	**Low discrimination**					**23753012**
	Actinomyces naeslundii	72.85	0.48	MNS 88		
	Actinomyces israelii	27.15	0.31	MLZ 5		
23753013	**Good identification**					**23753013**
	Actinomyces naeslundii	100.0	0.51	MNS 88		
23753020	**Very good identification**					**23753020**
	Actinomyces naeslundii	99.44	0.70	MNS 88		
	Actinomyces israelii	0.56	0.22			
23753021	**Very good identification**					**23753021**
	Actinomyces naeslundii	100.0	0.73	MNS 88		
23753022	**Very good identification**					**23753022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.17	0.70	MNS 88		
Actinomyces israelii		0.83	0.25			
23753023	**Very good identification**					**23753023**
Actinomyces naeslundii		100.0	0.73	MNS 88		
23753030	**Good identification**					**23753030**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23753031	**Good identification**					**23753031**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23753032	**Good identification**					**23753032**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23753033	**Good identification**					**23753033**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23753040	**Low discrimination**					**23753040**
Actinomyces naeslundii		83.83	0.57	MNS 88		
Actinomyces israelii		16.17	0.35			
23753041	**Very good identification**					**23753041**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23753042	**Low discrimination**					**23753042**
Actinomyces naeslundii		77.56	0.57	MNS 88		
Actinomyces israelii		22.44	0.38			
23753043	**Very good identification**					**23753043**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23753051	**Good identification**					**23753051**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753053	**Good identification**					**23753053**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753060	**Very good identification**					**23753060**
Actinomyces naeslundii		98.78	0.57	MNS 88		
Actinomyces israelii		1.22	0.15			
23753061	**Very good identification**					**23753061**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23753062	**Very good identification**					**23753062**
Actinomyces naeslundii		98.19	0.57	MNS 88		
Actinomyces israelii		1.81	0.18			
23753063	**Very good identification**					**23753063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.60	MNS 88		
23753071	**Good identification**					**23753071**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753073	**Good identification**					**23753073**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753100	**Low discrimination**					**23753100**
Actinomyces naeslundii		88.74	0.77	MNS 88		
Actinomyces israelii		10.92	0.53			
Propionibacterium propionicum		0.34	0.26			
23753101	**Very good identification**					**23753101**
Actinomyces naeslundii		99.92	0.80	MNS 88		
Actinomyces israelii		0.08	0.18			
23753102	**Low discrimination**					**23753102**
Actinomyces naeslundii		83.25	0.77	MNS 88		
Actinomyces israelii		15.36	0.56			
Actinomyces viscosus		0.84	0.19			
23753103	**Very good identification**					**23753103**
Actinomyces naeslundii		99.61	0.80	MNS 88		
Actinomyces viscosus		0.22	0.11			
Actinomyces israelii		0.12	0.21			
23753110	**Low discrimination**					**23753110**
Actinomyces naeslundii		74.12	0.55	MNS 88		
Actinomyces israelii		25.88	0.38	MLZ 5		
23753111	**Very good identification**					**23753111**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23753112	**Low discrimination**					**23753112**
Actinomyces naeslundii		64.67	0.55	MNS 88		
Actinomyces israelii		33.88	0.41	MLZ 5		
Bifidobacterium adolescentis		1.45	0.12			
23753113	**Very good identification**					**23753113**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23753120	**Very good identification**					**23753120**
Actinomyces naeslundii		99.22	0.77	MNS 88		
Actinomyces israelii		0.78	0.32			
23753121	**Very good identification**					**23753121**
Actinomyces naeslundii		100.0	0.80	MNS 88		
23753122	**Very good identification**					**23753122**
					SPO	
Actinomyces naeslundii		98.67	0.77	MNS 88	1	
Actinomyces israelii		1.16	0.35		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		0.17	0.05		92	
23753123	**Very good identification**					**23753123**
Actinomyces naeslundii		100.0	0.80	MNS 88		
23753130	**Good identification**					**23753130**
Actinomyces naeslundii		97.82	0.55	MNS 88		
Actinomyces israelii		2.18	0.18	MLZ 5		
23753131	**Very good identification**					**23753131**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23753132	**Good identification**					**23753132**
Actinomyces naeslundii		96.77	0.55	MNS 88		
Actinomyces israelii		3.24	0.21	MLZ 5		
23753133	**Very good identification**					**23753133**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23753140	**Low discrimination**					**23753140**
Actinomyces naeslundii		77.35	0.64	MNS 88		
Actinomyces israelii		20.97	0.45			
Propionibacterium propionicum		1.68	0.26			
23753141	**Very good identification**					**23753141**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23753142	**Low discrimination**					**23753142**
Actinomyces naeslundii		69.48	0.64	MNS 88		
Actinomyces israelii		28.25	0.49			
Propionibacterium propionicum		2.27	0.29			
23753143	**Very good identification**					**23753143**
Actinomyces naeslundii		99.49	0.67	MNS 88		
Actinomyces israelii		0.27	0.14			
Propionibacterium propionicum		0.24	0.12			
23753150	**Low discrimination**					**23753150**
Actinomyces naeslundii		56.51	0.43	MNS 88		
Actinomyces israelii		43.49	0.31	MLZ 5		
23753151	**Good identification**					**23753151**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753153	**Good identification**					**23753153**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753160	**Very good identification**					**23753160**
Actinomyces naeslundii		98.30	0.64	MNS 88		
Actinomyces israelii		1.70	0.25			
23753161	**Very good identification**					**23753161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.67	MNS 88		
23753162	**Very good identification**					**23753162**
Actinomyces naeslundii		97.47	0.64	MNS 88		
Actinomyces israelii		2.53	0.28			
23753163	**Very good identification**					**23753163**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23753170	**Good identification**					**23753170**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23753171	**Good identification**					**23753171**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753172	**Acceptable identification**					**23753172**
Actinomyces naeslundii		93.14	0.43	MNS 88		
Actinomyces israelii		6.86	0.14	MLZ 5		
23753173	**Good identification**					**23753173**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753200	**Low discrimination**					**23753200**
Actinomyces naeslundii		60.53	0.35	MNS 88		
Actinomyces israelii		39.48	0.23	RHA 1		
23753201	**Good identification**					**23753201**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753202	**Low discrimination**					**23753202**
Actinomyces naeslundii		50.55	0.35	MNS 88		
Actinomyces israelii		49.45	0.26	RHA 1		
23753203	**Good identification**					**23753203**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753220	**Good identification**					**23753220**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23753221	**Good identification**					**23753221**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753222	**Good identification**					**23753222**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23753223	**Good identification**					**23753223**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753300	**Low discrimination**					**23753300**
Actinomyces naeslundii		52.18	0.43	MNS 88		
Actinomyces israelii		47.82	0.33	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23753301	**Good identification**					**23753301**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753302	**Low discrimination**					**23753302**
Actinomyces israelii		57.89	0.36	XYL 92		
Actinomyces naeslundii		42.11	0.43	RHA 7		
23753303	**Good identification**					**23753303**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753320	**Good identification**					**23753320**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23753321	**Good identification**					**23753321**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753322	**Low discrimination**					**23753322**
					SPO	
Actinomyces naeslundii		76.33	0.43	MNS 88	1	
Clostridium ramosum		16.97	0.07	RHA 1	92	
Actinomyces israelii		6.70	0.16		1	
23753323	**Good identification**					**23753323**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753400	**Low discrimination**					**23753400**
Actinomyces israelii		87.64	0.41	XYL 92		
Actinomyces naeslundii		12.36	0.35			
23753401	**Good identification**					**23753401**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753402	**Acceptable identification**					**23753402**
Actinomyces israelii		91.40	0.44	XYL 92		
Actinomyces naeslundii		8.60	0.35			
23753403	**Good identification**					**23753403**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753420	**Low discrimination**					**23753420**
Actinomyces naeslundii		68.85	0.35	MNS 88		
Actinomyces israelii		31.15	0.20	ARA 1		
23753421	**Good identification**					**23753421**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753422	**Low discrimination**					**23753422**
Actinomyces naeslundii		59.57	0.35	MNS 88		
Actinomyces israelii		40.43	0.23	ARA 1		
23753423	**Good identification**					**23753423**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MNS 88		
23753442	**Good identification**					**23753442**
Actinomyces israelii		95.91	0.37	XYL 92		
Actinomyces naeslundii		4.09	0.22			
23753500	**Acceptable identification**					**23753500**
Actinomyces israelii		90.88	0.51	XYL 92		
Actinomyces naeslundii		9.12	0.43			
23753501	**Acceptable identification**					**23753501**
Actinomyces naeslundii		93.71	0.46	MNS 88		
Actinomyces israelii		6.29	0.17	ARA 1		
23753502	**Acceptable identification**					**23753502**
Actinomyces israelii		90.10	0.54	XYL 92		
Actinomyces naeslundii		6.03	0.43			
Bifidobacterium adolescentis		3.87	0.24			
23753503	**Acceptable identification**					**23753503**
Actinomyces naeslundii		90.86	0.46	MNS 88		
Actinomyces israelii		9.14	0.20	ARA 1		
23753510	**Good identification**					**23753510**
Actinomyces israelii		100.0	0.37	XYL 92		
23753512	**Low discrimination**					**23753512**
Actinomyces israelii		74.02	0.40	XYL 92		
Bifidobacterium adolescentis		25.98	0.26	MLZ 13		
23753520	**Low discrimination**					**23753520**
Actinomyces naeslundii		61.13	0.43	MNS 88		
Actinomyces israelii		38.87	0.31	ARA 1		
23753521	**Good identification**					**23753521**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753522	**Low discrimination**					**23753522**
Actinomyces naeslundii		51.18	0.43	MNS 88		
Actinomyces israelii		48.82	0.34	ARA 1		
23753523	**Good identification**					**23753523**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23753540	**Good identification**					**23753540**
Actinomyces israelii		95.64	0.44	XYL 92		
Actinomyces naeslundii		4.36	0.30			
23753542	**Good identification**					**23753542**
Actinomyces israelii		95.79	0.47	XYL 92		
Actinomyces naeslundii		2.91	0.30			
Bifidobacterium adolescentis		1.31	0.09			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23755000	**Low discrimination**					**23755000**
Actinomyces israelii		82.47	0.47			
Actinomyces odontolyticus		15.11	0.27			
Actinomyces naeslundii		2.42	0.30			
23755002	**Low discrimination**					**23755002**
Actinomyces israelii		86.60	0.50			
Actinomyces odontolyticus		6.48	0.23			
Actinomyces viscosus		5.22	0.15			
23755012	**Good identification**					**23755012**
Actinomyces israelii		100.0	0.36	MLZ 13		
23755040	**Good identification**					**23755040**
Actinomyces israelii		100.0	0.40			
23755042	**Good identification**					**23755042**
Actinomyces israelii		100.0	0.43			
23755100	**Very good identification**					**23755100**
Actinomyces israelii		96.18	0.58			
Actinomyces naeslundii		2.01	0.38			
Actinomyces odontolyticus		1.39	0.18			
23755101	**Low discrimination**					**23755101**
Actinomyces naeslundii		75.59	0.41	MNS 88		
Actinomyces israelii		24.42	0.24	XYL 1		
23755102	**Good identification**					**23755102**
Actinomyces israelii		93.66	0.61			
Actinomyces viscosus		2.51	0.19			
Bifidobacterium adolescentis		1.98	0.26			
23755103	**Low discrimination**					**23755103**
Actinomyces naeslundii		52.28	0.41	MNS 88		
Actinomyces israelii		25.33	0.27	XYL 1		
Actinomyces viscosus		22.40	0.11			
23755110	**Good identification**					**23755110**
Actinomyces israelii		100.0	0.44	MLZ 13		
23755112	**Low discrimination**					**23755112**
Actinomyces israelii		85.26	0.47	MLZ 13		
Bifidobacterium adolescentis		14.74	0.27			
23755120	**Low discrimination**					**23755120**
Actinomyces israelii		75.38	0.37	NAG 6		
Actinomyces naeslundii		24.62	0.38			
23755121	**Good identification**					**23755121**
Actinomyces naeslundii		100.0	0.41	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23755122	**Low discrimination**					**23755122**
					GRA	
Actinomyces israelii		78.08	0.40	NAG 6	99	
Actinomyces naeslundii		17.00	0.38		99	
Prevotella oralis		4.92	0.03		1	
23755123	**Good identification**					**23755123**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23755140	**Good identification**					**23755140**
Actinomyces israelii		99.06	0.51			
Actinomyces naeslundii		0.94	0.25			
23755142	**Good identification**					**23755142**
Actinomyces israelii		98.71	0.54			
Bifidobacterium adolescentis		0.66	0.10			
Actinomyces naeslundii		0.62	0.25			
23755150	**Good identification**					**23755150**
Actinomyces israelii		100.0	0.37	MLZ 13		
23755152	**Acceptable identification**					**23755152**
Actinomyces israelii		94.79	0.40	MLZ 13		
Bifidobacterium adolescentis		5.21	0.12			
23755300	**Low discrimination**					**23755300**
Actinomyces israelii		72.83	0.39	RHA 7		
Actinomyces odontolyticus		27.17	0.23			
23755302	**Low discrimination**					**23755302**
Actinomyces israelii		86.77	0.42	RHA 7		
Actinomyces odontolyticus		13.23	0.20			
23755400	**Good identification**					**23755400**
Actinomyces israelii		100.0	0.46			
23755402	**Good identification**					**23755402**
Actinomyces israelii		100.0	0.49			
23755440	**Good identification**					**23755440**
Actinomyces israelii		100.0	0.39			
23755442	**Good identification**					**23755442**
Actinomyces israelii		100.0	0.42			
23755500	**Very good identification**					**23755500**
					SPO	
Actinomyces israelii		99.46	0.56		1	
Clostridium butyricum		0.54	-.04		99	
23755502	**Low discrimination**					**23755502**
Actinomyces israelii		85.26	0.59			
Bifidobacterium adolescentis		14.74	0.40			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23755510	**Good identification**					**23755510**
Actinomyces israelii		97.90	0.42	MLZ 13		
Bifidobacterium adolescentis		2.10	0.07			
23755512	**Low discrimination**					**23755512**
Bifidobacterium adolescentis		58.59	0.41	NIT 1		
Actinomyces israelii		41.42	0.45	MNS 87		
23755520	**Good identification**					**23755520**
Actinomyces israelii		100.0	0.36	NAG 6		
23755522	**Good identification**					**23755522**
Actinomyces israelii		100.0	0.39	NAG 6		
23755540	**Good identification**					**23755540**
Actinomyces israelii		100.0	0.49			
23755542	**Acceptable identification**					**23755542**
Actinomyces israelii		94.79	0.52			
Bifidobacterium adolescentis		5.21	0.24			
23755550	**Good identification**					**23755550**
Actinomyces israelii		100.0	0.35	MLZ 13		
23755552	**Low discrimination**					**23755552**
Actinomyces israelii		68.99	0.38	MLZ 13		
Bifidobacterium adolescentis		31.01	0.26			
23755700	**Good identification**					**23755700**
Actinomyces israelii		100.0	0.37	RHA 7		
23755702	**Good identification**					**23755702**
Actinomyces israelii		100.0	0.40	RHA 7		
23756100	**Good identification**					**23756100**
Actinomyces israelii		100.0	0.37	GAL 99		
23756102	**Good identification**					**23756102**
Actinomyces israelii		100.0	0.40	GAL 99		
23756500	**Good identification**					**23756500**
Actinomyces israelii		100.0	0.35	GAL 99		
23756502	**Good identification**					**23756502**
Actinomyces israelii		100.0	0.38	GAL 99		
23757000	**Very good identification**					**23757000**
Actinomyces israelii		99.01	0.60			
Actinomyces naeslundii		0.99	0.35			
23757001	**Low discrimination**					**23757001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		59.85	0.38	MNS 88		
Actinomyces israelii		40.16	0.26	XYL 1		
23757002	**Very good identification**					**23757002**
Actinomyces israelii		99.34	0.63			
Actinomyces naeslundii		0.67	0.35			
23757010	**Good identification**					**23757010**
Actinomyces israelii		100.0	0.46	MLZ 13		
23757012	**Good identification**					**23757012**
Actinomyces israelii		100.0	0.49	MLZ 13		
23757020	**Low discrimination**					**23757020**
Actinomyces israelii		86.41	0.40	NAG 6		
Actinomyces naeslundii		13.59	0.35			
23757021	**Good identification**					**23757021**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23757022	**Acceptable identification**					**23757022**
Actinomyces israelii		90.51	0.43	NAG 6		
Actinomyces naeslundii		9.49	0.35			
23757023	**Good identification**					**23757023**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23757040	**Good identification**					**23757040**
Actinomyces israelii		99.55	0.53			
Actinomyces naeslundii		0.45	0.22			
23757042	**Very good identification**					**23757042**
Actinomyces israelii		99.70	0.56			
Actinomyces naeslundii		0.30	0.22			
23757050	**Good identification**					**23757050**
Actinomyces israelii		100.0	0.39	MLZ 13		
23757052	**Good identification**					**23757052**
Actinomyces israelii		100.0	0.42	MLZ 13		
23757062	**Good identification**					**23757062**
Actinomyces israelii		95.46	0.36	NAG 6		
Actinomyces naeslundii		4.54	0.22			
23757100	**Very good identification**					**23757100**
Actinomyces israelii		99.29	0.71			
Actinomyces naeslundii		0.71	0.43			
23757101	**Low discrimination**					**23757101**
Actinomyces naeslundii		51.47	0.46	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		48.53	0.37	XYL 1		
23757102	**Very good identification**					**23757102**
Actinomyces israelii		99.23	0.74			
Actinomyces naeslundii		0.47	0.43			
Bifidobacterium adolescentis		0.30	0.24			
23757103	**Low discrimination**					**23757103**
Actinomyces israelii		58.58	0.40	URE 1		
Actinomyces naeslundii		41.42	0.46			
23757110	**Very good identification**					**23757110**
Actinomyces israelii		100.0	0.57	MLZ 13		
23757112	**Very good identification**					**23757112**
Actinomyces israelii		97.56	0.60	MLZ 13		
Bifidobacterium adolescentis		2.44	0.26			
23757120	**Low discrimination**					**23757120**
Actinomyces israelii		89.94	0.50	NAG 6		
Actinomyces naeslundii		10.06	0.43			
23757121	**Acceptable identification**					**23757121**
Actinomyces naeslundii		94.32	0.46	MNS 88		
Actinomyces israelii		5.68	0.16	XYL 1		
23757122	**Acceptable identification**					**23757122**
Actinomyces israelii		93.06	0.53	NAG 6		
Actinomyces naeslundii		6.94	0.43			
23757123	**Acceptable identification**					**23757123**
Actinomyces naeslundii		91.72	0.46	MNS 88		
Actinomyces israelii		8.28	0.19	XYL 1		
23757130	**Good identification**					**23757130**
Actinomyces israelii		100.0	0.36	MLZ 13		
23757132	**Good identification**					**23757132**
Actinomyces israelii		100.0	0.39	MLZ 13		
23757140	**Very good identification**					**23757140**
Actinomyces israelii		99.68	0.64			
Actinomyces naeslundii		0.32	0.30			
23757142	**Very good identification**					**23757142**
Actinomyces israelii		99.69	0.67			
Actinomyces naeslundii		0.22	0.30			
Bifidobacterium adolescentis		0.10	0.09			
23757150	**Good identification**					**23757150**
Actinomyces israelii		100.0	0.50	MLZ 13		
23757152	**Good identification**					**23757152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		99.21	0.53	MLZ 13		
Bifidobacterium adolescentis		0.79	0.10			
23757160	**Good identification**					**23757160**
Actinomyces israelii		95.17	0.43	NAG 6		
Actinomyces naeslundii		4.83	0.30			
23757162	**Good identification**					**23757162**
Actinomyces israelii		96.73	0.46	NAG 6		
Actinomyces naeslundii		3.27	0.30			
23757200	**Good identification**					**23757200**
Actinomyces israelii		100.0	0.41	RHA 7		
23757202	**Good identification**					**23757202**
Actinomyces israelii		100.0	0.44	RHA 7		
23757242	**Good identification**					**23757242**
Actinomyces israelii		100.0	0.37	RHA 7		
23757300	**Good identification**					**23757300**
Actinomyces israelii		100.0	0.51	RHA 7		
23757302	**Good identification**					**23757302**
Actinomyces israelii		100.0	0.54	RHA 7		
23757310	**Good identification**					**23757310**
Actinomyces israelii		100.0	0.37	RHA 7		
23757312	**Good identification**					**23757312**
Actinomyces israelii		100.0	0.40	RHA 7		
23757340	**Good identification**					**23757340**
Actinomyces israelii		100.0	0.44	RHA 7		
23757342	**Good identification**					**23757342**
Actinomyces israelii		100.0	0.47	RHA 7		
23757400	**Very good identification**					**23757400**
Actinomyces israelii		100.0	0.59			
23757402	**Very good identification**					**23757402**
Actinomyces israelii		100.0	0.62			
23757410	**Good identification**					**23757410**
Actinomyces israelii		100.0	0.45	MLZ 13		
23757412	**Good identification**					**23757412**
Actinomyces israelii		100.0	0.48	MLZ 13		
23757420	**Good identification**					**23757420**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.38	NAG 6		
23757422	**Good identification**					**23757422**
Actinomyces israelii		100.0	0.41	NAG 6		
23757440	**Good identification**					**23757440**
Actinomyces israelii		100.0	0.52			
23757442	**Good identification**					**23757442**
Actinomyces israelii		100.0	0.55			
23757450	**Good identification**					**23757450**
Actinomyces israelii		100.0	0.38	MLZ 13		
23757452	**Good identification**					**23757452**
Actinomyces israelii		100.0	0.41	MLZ 13		
23757500	**Very good identification**					**23757500**
Actinomyces israelii		100.0	0.69			
23757501	**Good identification**					**23757501**
Actinomyces israelii		100.0	0.35	URE 1		
23757502	**Very good identification**					**23757502**
Actinomyces israelii		97.56	0.72			
Bifidobacterium adolescentis		2.44	0.38			
23757503	**Good identification**					**23757503**
Actinomyces israelii		100.0	0.38	URE 1		
23757510	**Good identification**					**23757510**
Actinomyces israelii		100.0	0.55	MLZ 13		
23757512	**Low discrimination**					**23757512**
Actinomyces israelii		83.04	0.58	MLZ 13		
Bifidobacterium adolescentis		16.96	0.40			
23757520	**Good identification**					**23757520**
Actinomyces israelii		100.0	0.49	NAG 6		
23757522	**Good identification**					**23757522**
Actinomyces israelii		100.0	0.52	NAG 6		
23757532	**Good identification**					**23757532**
Actinomyces israelii		100.0	0.38	MLZ 13		
23757540	**Very good identification**					**23757540**
Actinomyces israelii		100.0	0.62			
23757542	**Very good identification**					**23757542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		99.21	0.65			
Bifidobacterium adolescentis		0.79	0.23			
23757550	**Good identification**					**23757550**
Actinomyces israelii		100.0	0.48	MLZ 13		
23757552	**Acceptable identification**					**23757552**
Actinomyces israelii		93.90	0.51	MLZ 13		
Bifidobacterium adolescentis		6.10	0.24			
23757560	**Good identification**					**23757560**
Actinomyces israelii		100.0	0.42	NAG 6		
23757562	**Good identification**					**23757562**
Actinomyces israelii		100.0	0.45	NAG 6		
23757600	**Good identification**					**23757600**
Actinomyces israelii		100.0	0.40	RHA 7		
23757602	**Good identification**					**23757602**
Actinomyces israelii		100.0	0.43	RHA 7		
23757642	**Good identification**					**23757642**
Actinomyces israelii		100.0	0.36	RHA 7		
23757700	**Good identification**					**23757700**
Actinomyces israelii		100.0	0.50	RHA 7		
23757702	**Good identification**					**23757702**
Actinomyces israelii		100.0	0.53	RHA 7		
23757710	**Good identification**					**23757710**
Actinomyces israelii		100.0	0.36	RHA 7		
23757712	**Good identification**					**23757712**
Actinomyces israelii		100.0	0.39	RHA 7		
23757740	**Good identification**					**23757740**
Actinomyces israelii		100.0	0.43	RHA 7		
23757742	**Good identification**					**23757742**
Actinomyces israelii		100.0	0.46	RHA 7		
23761000	**Good identification**					**23761000**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23761001	**Good identification**					**23761001**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23761002	**Low discrimination**					**23761002**
Actinomyces naeslundii		89.31	0.48	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		10.69	0.08	FRU 95		
23761003	**Good identification**					**23761003**
Actinomyces naeslundii		97.41	0.51	MNS 88		
Actinomyces viscosus		2.59	-.01	FRU 95		
23761020	**Good identification**					**23761020**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23761021	**Good identification**					**23761021**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23761022	**Good identification**					**23761022**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23761023	**Good identification**					**23761023**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23761041	**Good identification**					**23761041**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23761043	**Good identification**					**23761043**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23761061	**Good identification**					**23761061**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23761063	**Good identification**					**23761063**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23761100	**Good identification**					**23761100**
Actinomyces naeslundii		98.46	0.55	MNS 88		
Actinomyces israelii		1.54	0.15	FRU 95		
23761101	**Very good identification**					**23761101**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23761102	**Acceptable identification**					**23761102**
Actinomyces naeslundii		91.05	0.55	MNS 88		
Actinomyces viscosus		6.81	0.12	FRU 95		
Actinomyces israelii		2.14	0.18			
23761103	**Very good identification**					**23761103**
Actinomyces naeslundii		98.37	0.58	MNS 88		
Actinomyces viscosus		1.63	0.04	FRU 95		
23761111	**Good identification**					**23761111**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23761113	**Good identification**					**23761113**
Actinomyces naeslundii		100.0	0.37	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23761120	**Good identification**					**23761120**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23761121	**Very good identification**					**23761121**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23761122	**Good identification**					**23761122**
Actinomyces naeslundii		100.0	0.55	MNS 88		
23761123	**Very good identification**					**23761123**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23761131	**Good identification**					**23761131**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23761133	**Good identification**					**23761133**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23761140	**Good identification**					**23761140**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23761141	**Good identification**					**23761141**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23761142	**Good identification**					**23761142**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23761143	**Good identification**					**23761143**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23761160	**Good identification**					**23761160**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23761161	**Good identification**					**23761161**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23761162	**Good identification**					**23761162**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23761163	**Good identification**					**23761163**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23762100	**Good identification**					**23762100**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23762101	**Good identification**					**23762101**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23762102	**Good identification**					**23762102**
Actinomyces naeslundii		100.0	0.37	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23762103	**Good identification**					**23762103**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23762120	**Good identification**					**23762120**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23762121	**Good identification**					**23762121**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23762122	**Good identification**					**23762122**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23762123	**Good identification**					**23762123**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23763000	**Good identification**					**23763000**
Actinomyces naeslundii		96.85	0.53	MNS 88		
Actinomyces israelii		3.15	0.18	FRU 95		
23763001	**Good identification**					**23763001**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23763002	**Good identification**					**23763002**
Actinomyces naeslundii		95.34	0.53	MNS 88		
Actinomyces israelii		4.66	0.21	FRU 95		
23763003	**Good identification**					**23763003**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23763020	**Good identification**					**23763020**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23763021	**Good identification**					**23763021**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23763022	**Good identification**					**23763022**
Actinomyces naeslundii		100.0	0.53	MNS 88		
23763023	**Good identification**					**23763023**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23763040	**Good identification**					**23763040**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23763041	**Good identification**					**23763041**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23763042	**Acceptable identification**					**23763042**
Actinomyces naeslundii		90.29	0.40	MNS 88		
Actinomyces israelii		9.71	0.14	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23763043	**Good identification**					**23763043**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23763060	**Good identification**					**23763060**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23763061	**Good identification**					**23763061**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23763062	**Good identification**					**23763062**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23763063	**Good identification**					**23763063**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23763100	**Very good identification**					**23763100**
Actinomyces naeslundii		95.63	0.60	MNS 88		
Actinomyces israelii		4.37	0.28	FRU 95		
23763101	**Very good identification**					**23763101**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23763102	**Good identification**					**23763102**
Actinomyces naeslundii		93.58	0.60	MNS 88		
Actinomyces israelii		6.42	0.31	FRU 95		
23763103	**Very good identification**					**23763103**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23763110	**Low discrimination**					**23763110**
Actinomyces naeslundii		88.51	0.38	MNS 88		
Actinomyces israelii		11.49	0.14	FRU 95		
23763111	**Good identification**					**23763111**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23763112	**Low discrimination**					**23763112**
Actinomyces naeslundii		83.69	0.38	MNS 88		
Actinomyces israelii		16.31	0.17	FRU 95		
23763113	**Good identification**					**23763113**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23763120	**Very good identification**					**23763120**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23763121	**Very good identification**					**23763121**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23763122	**Very good identification**					**23763122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.60	MNS 88		
23763123	**Very good identification**					**23763123**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23763130	**Good identification**					**23763130**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23763131	**Good identification**					**23763131**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23763132	**Good identification**					**23763132**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23763133	**Good identification**					**23763133**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23763140	**Acceptable identification**					**23763140**
Actinomyces naeslundii		90.84	0.47	MNS 88		
Actinomyces israelii		9.16	0.21	FRU 95		
23763141	**Good identification**					**23763141**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23763142	**Low discrimination**					**23763142**
Actinomyces naeslundii		86.87	0.47	MNS 88		
Actinomyces israelii		13.14	0.24	FRU 95		
23763143	**Good identification**					**23763143**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23763160	**Good identification**					**23763160**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23763161	**Good identification**					**23763161**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23763162	**Good identification**					**23763162**
Actinomyces naeslundii		100.0	0.47	MNS 88		
23763163	**Good identification**					**23763163**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23765102	**Good identification**					**23765102**
Actinomyces israelii		100.0	0.36	FRU 99		
23767000	**Good identification**					**23767000**
Actinomyces israelii		100.0	0.36	FRU 99		
23767002	**Good identification**					**23767002**
Actinomyces israelii		100.0	0.39	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23767100	**Good identification**					**23767100**
Actinomyces israelii		98.12	0.46	FRU 99		
Actinomyces naeslundii		1.88	0.26			
23767102	**Good identification**					**23767102**
Actinomyces israelii		98.74	0.49	FRU 99		
Actinomyces naeslundii		1.26	0.26			
23767112	**Good identification**					**23767112**
Actinomyces israelii		100.0	0.35	FRU 99		
23767140	**Good identification**					**23767140**
Actinomyces israelii		100.0	0.39	FRU 99		
23767142	**Good identification**					**23767142**
Actinomyces israelii		100.0	0.42	FRU 99		
23767402	**Good identification**					**23767402**
Actinomyces israelii		100.0	0.38	FRU 99		
23767500	**Good identification**					**23767500**
Actinomyces israelii		100.0	0.45	FRU 99		
23767502	**Good identification**					**23767502**
Actinomyces israelii		100.0	0.48	FRU 99		
23767540	**Good identification**					**23767540**
Actinomyces israelii		100.0	0.38	FRU 99		
23767542	**Good identification**					**23767542**
Actinomyces israelii		100.0	0.41	FRU 99		
23770000	**Low discrimination**					**23770000**
					GRA	
Actinomyces naeslundii		88.35	0.46	MNS 88	99	
Fusobacterium mortiferum		7.47	0.16	GAL 96	1	
Actinomyces viscosus		4.18	-.01		99	
23770001	**Good identification**					**23770001**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23770020	**Good identification**					**23770020**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23770021	**Good identification**					**23770021**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23770022	**Good identification**					**23770022**
Actinomyces naeslundii		95.48	0.46	MNS 88		
Actinomyces viscosus		4.52	-.01	GAL 96		
23770023	**Good identification**					**23770023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	MNS 88		
23770041	**Good identification**					**23770041**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23770043	**Good identification**					**23770043**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23770061	**Good identification**					**23770061**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23770063	**Good identification**					**23770063**
Actinomyces naeslundii		100.0	0.36	MNS 88		
23770100	**Low discrimination**					**23770100**
					GRA	
Actinomyces naeslundii		89.39	0.54	MNS 88	99	
Fusobacterium mortiferum		6.20	0.22	GAL 96	1	
Actinomyces viscosus		2.64	0.03		99	
23770101	**Very good identification**					**23770101**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23770102	**Low discrimination**					**23770102**
Actinomyces viscosus		73.98	0.37	MNS 96		
Actinomyces naeslundii		25.27	0.54	TRE 7		
Actinomyces israelii		0.75	0.18			
23770103	**Low discrimination**					**23770103**
Actinomyces naeslundii		60.59	0.57	MNS 88		
Actinomyces viscosus		39.41	0.29	GAL 96		
23770120	**Good identification**					**23770120**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23770121	**Very good identification**					**23770121**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23770122	**Good identification**					**23770122**
Actinomyces naeslundii		97.13	0.54	MNS 88		
Actinomyces viscosus		2.87	0.03	GAL 96		
23770123	**Very good identification**					**23770123**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23770140	**Good identification**					**23770140**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23770141	**Good identification**					**23770141**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23770142	**Low discrimination**					**23770142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		85.65	0.41	MNS 88		
Actinomyces viscosus		14.35	0.03	GAL 96		
23770143	**Good identification**					**23770143**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23770160	**Good identification**					**23770160**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23770161	**Good identification**					**23770161**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23770162	**Good identification**					**23770162**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23770163	**Good identification**					**23770163**
Actinomyces naeslundii		100.0	0.44	MNS 88		
23771000	**Good identification**					**23771000**
					GRA	
Actinomyces naeslundii		94.04	0.70	MNS 88	99	
Actinomyces israelii		5.46	0.39		99	
Fusobacterium mortiferum		0.50	0.19		1	
23771001	**Very good identification**					**23771001**
Actinomyces naeslundii		100.0	0.73	MNS 88		
23771002	**Low discrimination**					**23771002**
Actinomyces naeslundii		82.67	0.70	MNS 88		
Actinomyces viscosus		9.89	0.30			
Actinomyces israelii		7.21	0.42			
23771003	**Very good identification**					**23771003**
Actinomyces naeslundii		97.41	0.73	MNS 88		
Actinomyces viscosus		2.59	0.21			
23771010	**Low discrimination**					**23771010**
Actinomyces naeslundii		85.84	0.48	MNS 88		
Actinomyces israelii		14.16	0.25	MLZ 5		
23771011	**Good identification**					**23771011**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23771012	**Low discrimination**					**23771012**
Actinomyces naeslundii		80.17	0.48	MNS 88		
Actinomyces israelii		19.83	0.28	MLZ 5		
23771013	**Good identification**					**23771013**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23771020	**Very good identification**					**23771020**
Actinomyces naeslundii		99.63	0.70	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.37	0.19			
23771021	**Very good identification**					**23771021**
Actinomyces naeslundii		100.0	0.73	MNS 88		
23771022	**Very good identification**					**23771022**
Actinomyces naeslundii		99.18	0.70	MNS 88		
Actinomyces israelii		0.55	0.22			
Lactobacillus acidophilus		0.27	0.01			
23771023	**Very good identification**					**23771023**
Actinomyces naeslundii		100.0	0.73	MNS 88		
23771030	**Good identification**					**23771030**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23771031	**Good identification**					**23771031**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23771032	**Good identification**					**23771032**
Actinomyces naeslundii		100.0	0.48	MNS 88		
23771033	**Good identification**					**23771033**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23771040	**Low discrimination**					**23771040**
Actinomyces naeslundii		88.65	0.57	MNS 88		
Actinomyces israelii		11.35	0.32			
23771041	**Very good identification**					**23771041**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23771042	**Low discrimination**					**23771042**
Actinomyces naeslundii		83.89	0.57	MNS 88		
Actinomyces israelii		16.11	0.35			
23771043	**Very good identification**					**23771043**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23771051	**Good identification**					**23771051**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771053	**Good identification**					**23771053**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771060	**Very good identification**					**23771060**
Actinomyces naeslundii		100.0	0.57	MNS 88		
23771061	**Very good identification**					**23771061**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23771062	**Very good identification**					**23771062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.79	0.57	MNS 88		
Actinomyces israelii		1.21	0.15			
23771063	**Very good identification**					**23771063**
Actinomyces naeslundii		100.0	0.60	MNS 88		
23771071	**Good identification**					**23771071**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771073	**Good identification**					**23771073**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771100	**Good identification**					**23771100**
					GRA	
Actinomyces naeslundii		92.02	0.77	MNS 88	99	
Actinomyces israelii		7.51	0.49		99	
Fusobacterium mortiferum		0.40	0.25		1	
23771101	**Very good identification**					**23771101**
Actinomyces naeslundii		99.95	0.80	MNS 88		
Actinomyces israelii		0.06	0.15			
23771102	**Low discrimination**					**23771102**
Actinomyces naeslundii		82.70	0.77	MNS 88		
Actinomyces israelii		10.13	0.52			
Actinomyces viscosus		6.18	0.34			
23771103	**Very good identification**					**23771103**
Actinomyces naeslundii		98.29	0.80	MNS 88		
Actinomyces viscosus		1.63	0.26			
Actinomyces israelii		0.08	0.18			
23771110	**Low discrimination**					**23771110**
Actinomyces naeslundii		81.18	0.55	MNS 88		
Actinomyces israelii		18.82	0.35	MLZ 5		
23771111	**Very good identification**					**23771111**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23771112	**Low discrimination**					**23771112**
					GRA	
Actinomyces naeslundii		65.23	0.55	MNS 88	99	
Actinomyces israelii		22.68	0.38	MLZ 5	99	
Leptotrichia buccalis		9.96	0.07		1	
23771113	**Very good identification**					**23771113**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23771120	**Very good identification**					**23771120**
Actinomyces naeslundii		99.10	0.77	MNS 88		
Actinomyces israelii		0.52	0.29			
Clostridium perfringens		0.27	0.11			
23771121	**Very good identification**					**23771121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.80	MNS 88		
23771122	**Very good identification**					**23771122**
Actinomyces naeslundii		97.43	0.77	MNS 88		
Lactobacillus acidophilus		1.08	0.19			
Actinomyces israelii		0.76	0.32			
23771123	**Very good identification**					**23771123**
Actinomyces naeslundii		100.0	0.80	MNS 88		
23771130	**Good identification**					**23771130**
Actinomyces naeslundii		98.54	0.55	MNS 88		
Actinomyces israelii		1.46	0.15	MLZ 5		
23771131	**Very good identification**					**23771131**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23771132	**Good identification**					**23771132**
					GRA	
Actinomyces naeslundii		96.23	0.55	MNS 88	99	
Actinomyces israelii		2.14	0.18	MLZ 5	99	
Leptotrichia buccalis		1.63	-.09		1	
23771133	**Very good identification**					**23771133**
Actinomyces naeslundii		100.0	0.58	MNS 88		
23771140	**Low discrimination**					**23771140**
Actinomyces naeslundii		84.75	0.64	MNS 88		
Actinomyces israelii		15.25	0.42			
23771141	**Very good identification**					**23771141**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23771142	**Low discrimination**					**23771142**
Actinomyces naeslundii		78.00	0.64	MNS 88		
Actinomyces israelii		21.05	0.45			
Lactobacillus acidophilus		0.61	0.04			
23771143	**Very good identification**					**23771143**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23771150	**Low discrimination**					**23771150**
Actinomyces naeslundii		66.19	0.43	MNS 88		
Actinomyces israelii		33.81	0.28	MLZ 5		
23771151	**Good identification**					**23771151**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771152	**Low discrimination**					**23771152**
Actinomyces naeslundii		56.62	0.43	MNS 88		
Actinomyces israelii		43.39	0.31	MLZ 5		
23771153	**Good identification**					**23771153**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771160	**Very good identification**					**23771160**
Actinomyces naeslundii		98.87	0.64	MNS 88		
Actinomyces israelii		1.14	0.22			
23771161	**Very good identification**					**23771161**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23771162	**Very good identification**					**23771162**
Actinomyces naeslundii		97.56	0.64	MNS 88		
Actinomyces israelii		1.68	0.25			
Lactobacillus acidophilus		0.76	0.04			
23771163	**Very good identification**					**23771163**
Actinomyces naeslundii		100.0	0.67	MNS 88		
23771170	**Good identification**					**23771170**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23771171	**Good identification**					**23771171**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771172	**Good identification**					**23771172**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23771173	**Good identification**					**23771173**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771200	**Low discrimination**					**23771200**
Actinomyces naeslundii		69.79	0.35	MNS 88		
Actinomyces israelii		30.21	0.20	RHA 1		
23771201	**Good identification**					**23771201**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771202	**Low discrimination**					**23771202**
Actinomyces naeslundii		60.63	0.35	MNS 88		
Actinomyces israelii		39.37	0.23	RHA 1		
23771203	**Good identification**					**23771203**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771220	**Good identification**					**23771220**
Actinomyces naeslundii		100.0	0.35	MNS 88		
23771221	**Good identification**					**23771221**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771222	**Good identification**					**23771222**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	MNS 88		
23771223	**Good identification**					**23771223**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771300	**Low discrimination**					**23771300**
Actinomyces naeslundii		62.17	0.43	MNS 88		
Actinomyces israelii		37.83	0.30	RHA 1		
23771301	**Good identification**					**23771301**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771302	**Low discrimination**					**23771302**
Actinomyces naeslundii		48.30	0.43	MNS 88		
Actinomyces israelii		44.08	0.33	RHA 1		
Lactobacillus acidophilus		4.00	0.00			
23771303	**Good identification**					**23771303**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771320	**Low discrimination**					**23771320**
					SPO	
Actinomyces naeslundii		86.68	0.43	MNS 88	1	
Clostridium ramosum		13.32	0.04	RHA 1	92	
23771321	**Good identification**					**23771321**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771322	**Low discrimination**					**23771322**
					SPO	
Actinomyces naeslundii		56.76	0.43	MNS 88	1	
Clostridium ramosum		34.90	0.14	RHA 1	92	
Lactobacillus acidophilus		4.70	0.00		1	
23771323	**Good identification**					**23771323**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771400	**Low discrimination**					**23771400**
Actinomyces israelii		82.48	0.38	XYL 92		
Actinomyces naeslundii		17.53	0.35			
23771401	**Good identification**					**23771401**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771402	**Low discrimination**					**23771402**
Actinomyces israelii		87.59	0.41	XYL 92		
Actinomyces naeslundii		12.41	0.35			
23771403	**Good identification**					**23771403**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771420	**Low discrimination**					**23771420**
Actinomyces naeslundii		76.90	0.35	MNS 88		
Actinomyces israelii		23.10	0.17	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23771421	**Good identification**					**23771421**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771422	**Low discrimination**					**23771422**
Actinomyces naeslundii		68.94	0.35	MNS 88		
Actinomyces israelii		31.07	0.20	ARA 1		
23771423	**Good identification**					**23771423**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23771500	**Low discrimination**					**23771500**
Actinomyces israelii		86.87	0.48	XYL 92		
Actinomyces naeslundii		13.13	0.43			
23771501	**Good identification**					**23771501**
Actinomyces naeslundii		95.74	0.46	MNS 88		
Actinomyces israelii		4.26	0.14	ARA 1		
23771502	**Low discrimination**					**23771502**
Actinomyces israelii		86.13	0.51	XYL 92		
Actinomyces naeslundii		8.68	0.43			
Bifidobacterium adolescentis		4.54	0.23			
23771503	**Acceptable identification**					**23771503**
Actinomyces naeslundii		93.74	0.46	MNS 88		
Actinomyces israelii		6.26	0.17	ARA 1		
23771512	**Low discrimination**					**23771512**
Actinomyces israelii		69.89	0.37	XYL 92		
Bifidobacterium adolescentis		30.11	0.24	MLZ 13		
23771520	**Low discrimination**					**23771520**
Actinomyces naeslundii		70.32	0.43	MNS 88		
Actinomyces israelii		29.68	0.28	ARA 1		
23771521	**Good identification**					**23771521**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771522	**Low discrimination**					**23771522**
Actinomyces naeslundii		61.23	0.43	MNS 88		
Actinomyces israelii		38.77	0.31	ARA 1		
23771523	**Good identification**					**23771523**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23771540	**Acceptable identification**					**23771540**
Actinomyces israelii		93.58	0.41	XYL 92		
Actinomyces naeslundii		6.42	0.30			
23771542	**Acceptable identification**					**23771542**
Actinomyces israelii		94.12	0.44	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		4.31	0.30			
Bifidobacterium adolescentis		1.58	0.07			
23772000	**Good identification**					**23772000**
Actinomyces naeslundii		96.05	0.51	MNS 88		
Actinomyces israelii		3.95	0.18	GAL 96		
23772001	**Good identification**					**23772001**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23772002	**Acceptable identification**					**23772002**
Actinomyces naeslundii		92.08	0.51	MNS 88		
Actinomyces israelii		5.68	0.21	GAL 96		
Actinomyces viscosus		2.25	-.01			
23772003	**Good identification**					**23772003**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23772020	**Good identification**					**23772020**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23772021	**Good identification**					**23772021**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23772022	**Good identification**					**23772022**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23772023	**Good identification**					**23772023**
Actinomyces naeslundii		100.0	0.54	MNS 88		
23772040	**Good identification**					**23772040**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23772041	**Good identification**					**23772041**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23772042	**Low discrimination**					**23772042**
Actinomyces naeslundii		88.04	0.38	MNS 88		
Actinomyces israelii		11.96	0.14	GAL 96		
23772043	**Good identification**					**23772043**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23772060	**Good identification**					**23772060**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23772061	**Good identification**					**23772061**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23772062	**Good identification**					**23772062**
Actinomyces naeslundii		100.0	0.38	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23772063	**Good identification**					**23772063**
Actinomyces naeslundii		100.0	0.41	MNS 88		
23772100	**Good identification**					**23772100**
Actinomyces naeslundii		91.19	0.59	MNS 88		
Actinomyces israelii		5.27	0.28	GAL 96		
Propionibacterium propionicum		3.54	0.24			
23772101	**Very good identification**					**23772101**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23772102	**Low discrimination**					**23772102**
Actinomyces naeslundii		86.20	0.59	MNS 88		
Actinomyces israelii		7.47	0.31	GAL 96		
Propionibacterium propionicum		5.02	0.27			
23772103	**Very good identification**					**23772103**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23772110	**Low discrimination**					**23772110**
Actinomyces naeslundii		85.91	0.37	MNS 88		
Actinomyces israelii		14.09	0.14	GAL 96		
23772111	**Good identification**					**23772111**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23772112	**Low discrimination**					**23772112**
Actinomyces naeslundii		80.25	0.37	MNS 88		
Actinomyces israelii		19.75	0.17	GAL 96		
23772113	**Good identification**					**23772113**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23772120	**Very good identification**					**23772120**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23772121	**Very good identification**					**23772121**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23772122	**Very good identification**					**23772122**
Actinomyces naeslundii		100.0	0.59	MNS 88		
23772123	**Very good identification**					**23772123**
Actinomyces naeslundii		100.0	0.62	MNS 88		
23772130	**Good identification**					**23772130**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23772131	**Good identification**					**23772131**
Actinomyces naeslundii		100.0	0.40	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23772132	**Good identification**					**23772132**
Actinomyces naeslundii		100.0	0.37	MNS 88		
23772133	**Good identification**					**23772133**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23772140	**Low discrimination**					**23772140**
Actinomyces naeslundii		74.23	0.46	MNS 88		
Propionibacterium propionicum		16.32	0.24	GAL 96		
Actinomyces israelii		9.45	0.21			
23772141	**Good identification**					**23772141**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23772142	**Low discrimination**					**23772142**
Actinomyces naeslundii		65.76	0.46	MNS 88		
Propionibacterium propionicum		21.69	0.27	GAL 96		
Actinomyces israelii		12.56	0.24			
23772143	**Good identification**					**23772143**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23772160	**Good identification**					**23772160**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23772161	**Good identification**					**23772161**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23772162	**Good identification**					**23772162**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23772163	**Good identification**					**23772163**
Actinomyces naeslundii		100.0	0.49	MNS 88		
23773000	**Low discrimination**					**23773000**
Actinomyces naeslundii		85.50	0.74	MNS 88		
Actinomyces israelii		14.50	0.52			
23773001	**Very good identification**					**23773001**
Actinomyces naeslundii		99.89	0.77	MNS 88		
Actinomyces israelii		0.11	0.18			
23773002	**Low discrimination**					**23773002**
Actinomyces naeslundii		79.72	0.74	MNS 88		
Actinomyces israelii		20.28	0.55			
23773003	**Very good identification**					**23773003**
Actinomyces naeslundii		99.83	0.77	MNS 88		
Actinomyces israelii		0.17	0.21			
23773010	**Low discrimination**					**23773010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		67.50	0.53	MNS 88		
Actinomyces israelii		32.50	0.38	MLZ 5		
23773011	**Good identification**					**23773011**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23773012	**Low discrimination**					**23773012**
Actinomyces naeslundii		58.06	0.53	MNS 88		
Actinomyces israelii		41.94	0.41	MLZ 5		
23773013	**Good identification**					**23773013**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23773020	**Very good identification**					**23773020**
Actinomyces naeslundii		98.93	0.74	MNS 88		
Actinomyces israelii		1.07	0.32			
23773021	**Very good identification**					**23773021**
Actinomyces naeslundii		100.0	0.77	MNS 88		
23773022	**Very good identification**					**23773022**
Actinomyces naeslundii		98.40	0.74	MNS 88		
Actinomyces israelii		1.60	0.35			
23773023	**Very good identification**					**23773023**
Actinomyces naeslundii		100.0	0.77	MNS 88		
23773030	**Good identification**					**23773030**
Actinomyces naeslundii		97.02	0.53	MNS 88		
Actinomyces israelii		2.98	0.17	MLZ 5		
23773031	**Good identification**					**23773031**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23773032	**Good identification**					**23773032**
Actinomyces naeslundii		95.59	0.53	MNS 88		
Actinomyces israelii		4.41	0.20	MLZ 5		
23773033	**Good identification**					**23773033**
Actinomyces naeslundii		100.0	0.56	MNS 88		
23773040	**Low discrimination**					**23773040**
Actinomyces naeslundii		72.79	0.62	MNS 88		
Actinomyces israelii		27.21	0.45			
23773041	**Very good identification**					**23773041**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23773042	**Low discrimination**					**23773042**
Actinomyces naeslundii		64.08	0.62	MNS 88		
Actinomyces israelii		35.92	0.48			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23773043	**Very good identification**					**23773043**
Actinomyces naeslundii		99.62	0.65	MNS 88		
Actinomyces israelii		0.38	0.14			
23773051	**Good identification**					**23773051**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773053	**Good identification**					**23773053**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773060	**Very good identification**					**23773060**
Actinomyces naeslundii		97.67	0.62	MNS 88		
Actinomyces israelii		2.33	0.25			
23773061	**Very good identification**					**23773061**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23773062	**Very good identification**					**23773062**
Actinomyces naeslundii		96.55	0.62	MNS 88		
Actinomyces israelii		3.46	0.28			
23773063	**Very good identification**					**23773063**
Actinomyces naeslundii		100.0	0.65	MNS 88		
23773070	**Good identification**					**23773070**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23773071	**Good identification**					**23773071**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773072	**Acceptable identification**					**23773072**
Actinomyces naeslundii		90.78	0.40	MNS 88		
Actinomyces israelii		9.22	0.13	MLZ 5		
23773073	**Good identification**					**23773073**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773100	**Low discrimination**					**23773100**
Actinomyces naeslundii		80.65	0.82	MNS 88		
Actinomyces israelii		19.22	0.62			
Propionibacterium propionicum		0.13	0.24			
23773101	**Very good identification**					**23773101**
Actinomyces naeslundii		99.84	0.85	MNS 88		
Actinomyces israelii		0.16	0.28			
23773102	**Low discrimination**					**23773102**
Actinomyces naeslundii		73.49	0.82	MNS 88		
Actinomyces israelii		26.28	0.65			
Propionibacterium propionicum		0.18	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23773103	**Very good identification**					**23773103**
Actinomyces naeslundii		99.76	0.85	MNS 88		
Actinomyces israelii		0.24	0.31			
23773110	**Low discrimination**					**23773110**
Actinomyces naeslundii		59.64	0.60	MNS 88		
Actinomyces israelii		40.36	0.48	MLZ 5		
23773111	**Very good identification**					**23773111**
Actinomyces naeslundii		99.55	0.63	MNS 88		
Actinomyces israelii		0.45	0.14	MLZ 5		
23773112	**Low discrimination**					**23773112**
Actinomyces israelii		50.18	0.51	XYL 92		
Actinomyces naeslundii		49.44	0.60	MLZ 13		
Bifidobacterium adolescentis		0.38	0.09			
23773113	**Very good identification**					**23773113**
Actinomyces naeslundii		99.32	0.63	MNS 88		
Actinomyces israelii		0.68	0.17	MLZ 5		
23773120	**Very good identification**					**23773120**
					SPO	
Actinomyces naeslundii		98.17	0.82	MNS 88	1	
Actinomyces israelii		1.49	0.42		1	
Clostridium ramosum		0.34	0.15		92	
23773121	**Very good identification**					**23773121**
Actinomyces naeslundii		100.0	0.85	MNS 88		
23773122	**Very good identification**					**23773122**
					SPO	
Actinomyces naeslundii		96.48	0.82	MNS 88	1	
Actinomyces israelii		2.20	0.45		1	
Clostridium ramosum		1.32	0.25		92	
23773123	**Very good identification**					**23773123**
Actinomyces naeslundii		100.0	0.85	MNS 88		
23773130	**Very good identification**					**23773130**
Actinomyces naeslundii		95.86	0.60	MNS 88		
Actinomyces israelii		4.14	0.28	MLZ 5		
23773131	**Very good identification**					**23773131**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23773132	**Good identification**					**23773132**
Actinomyces naeslundii		93.92	0.60	MNS 88		
Actinomyces israelii		6.09	0.31	MLZ 5		
23773133	**Very good identification**					**23773133**
Actinomyces naeslundii		100.0	0.63	MNS 88		
23773140	**Low discrimination**					**23773140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		65.17	0.69	MNS 88		
Actinomyces israelii		34.23	0.55			
Propionibacterium propionicum		0.60	0.24			
23773141	**Very good identification**					**23773141**
Actinomyces naeslundii		99.65	0.72	MNS 88		
Actinomyces israelii		0.35	0.21			
23773142	**Low discrimination**					**23773142**
Actinomyces naeslundii		55.50	0.69	MNS 88		
Actinomyces israelii		43.73	0.58			
Propionibacterium propionicum		0.76	0.27			
23773143	**Very good identification**					**23773143**
Actinomyces naeslundii		99.47	0.72	MNS 88		
Actinomyces israelii		0.53	0.24			
23773150	**Low discrimination**					**23773150**
Actinomyces israelii		59.86	0.41	XYL 92		
Actinomyces naeslundii		40.14	0.47	MLZ 13		
23773151	**Good identification**					**23773151**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23773152	**Low discrimination**					**23773152**
Actinomyces israelii		69.11	0.44	XYL 92		
Actinomyces naeslundii		30.89	0.47	MLZ 13		
23773153	**Good identification**					**23773153**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23773160	**Very good identification**					**23773160**
Actinomyces naeslundii		96.76	0.69	MNS 88		
Actinomyces israelii		3.24	0.35			
23773161	**Very good identification**					**23773161**
Actinomyces naeslundii		100.0	0.72	MNS 88		
23773162	**Very good identification**					**23773162**
Actinomyces naeslundii		95.21	0.69	MNS 88		
Actinomyces israelii		4.79	0.38			
23773163	**Very good identification**					**23773163**
Actinomyces naeslundii		100.0	0.72	MNS 88		
23773170	**Acceptable identification**					**23773170**
Actinomyces naeslundii		91.31	0.47	MNS 88		
Actinomyces israelii		8.69	0.21	MLZ 5		
23773171	**Good identification**					**23773171**
Actinomyces naeslundii		100.0	0.50	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23773172	**Low discrimination**					**23773172**
Actinomyces naeslundii		87.51	0.47	MNS 88		
Actinomyces israelii		12.49	0.24	MLZ 5		
23773173	**Good identification**					**23773173**
Actinomyces naeslundii		100.0	0.50	MNS 88		
23773201	**Good identification**					**23773201**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773202	**Low discrimination**					**23773202**
Actinomyces israelii		65.47	0.36	XYL 92		
Actinomyces naeslundii		34.53	0.40	RHA 7		
23773203	**Good identification**					**23773203**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773220	**Good identification**					**23773220**
Actinomyces naeslundii		100.0	0.40	MNS 88		
23773221	**Good identification**					**23773221**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773222	**Low discrimination**					**23773222**
Actinomyces naeslundii		89.21	0.40	MNS 88		
Actinomyces israelii		10.79	0.15	RHA 1		
23773223	**Good identification**					**23773223**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773300	**Low discrimination**					**23773300**
Actinomyces israelii		63.98	0.43	XYL 92		
Actinomyces naeslundii		36.02	0.48	RHA 7		
23773301	**Good identification**					**23773301**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23773302	**Low discrimination**					**23773302**
Actinomyces israelii		72.71	0.46	XYL 92		
Actinomyces naeslundii		27.29	0.48	RHA 7		
23773303	**Good identification**					**23773303**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23773320	**Low discrimination**					**23773320**
					SPO	
Actinomyces naeslundii		64.02	0.48	MNS 88	1	
Clostridium ramosum		28.73	0.17	RHA 1	92	
Actinomyces israelii		7.26	0.23		1	
23773321	**Good identification**					**23773321**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	MNS 88		
23773323	**Good identification**					**23773323**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23773340	**Low discrimination**					**23773340**
Actinomyces israelii		79.65	0.36	XYL 92		
Actinomyces naeslundii		20.35	0.35	RHA 7		
23773341	**Good identification**					**23773341**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23773342	**Low discrimination**					**23773342**
Actinomyces israelii		85.45	0.39	XYL 92		
Actinomyces naeslundii		14.55	0.35	RHA 7		
23773343	**Good identification**					**23773343**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23773360	**Low discrimination**					**23773360**
Actinomyces naeslundii		80.01	0.35	MNS 88		
Actinomyces israelii		19.99	0.16	RHA 1		
23773361	**Good identification**					**23773361**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23773362	**Low discrimination**					**23773362**
Actinomyces naeslundii		72.74	0.35	MNS 88		
Actinomyces israelii		27.26	0.19	RHA 1		
23773363	**Good identification**					**23773363**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23773400	**Acceptable identification**					**23773400**
Actinomyces israelii		93.22	0.51	XYL 92		
Actinomyces naeslundii		6.79	0.40			
23773401	**Acceptable identification**					**23773401**
Actinomyces naeslundii		91.53	0.43	MNS 88		
Actinomyces israelii		8.47	0.16	ARA 1		
23773402	**Good identification**					**23773402**
Actinomyces israelii		95.37	0.54	XYL 92		
Actinomyces naeslundii		4.63	0.40			
23773403	**Low discrimination**					**23773403**
Actinomyces naeslundii		87.81	0.43	MNS 88		
Actinomyces israelii		12.19	0.19	ARA 1		
23773410	**Good identification**					**23773410**
Actinomyces israelii		100.0	0.36	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23773412	**Good identification**					**23773412**
Actinomyces israelii		100.0	0.39	XYL 92		
23773420	**Low discrimination**					**23773420**
Actinomyces naeslundii		53.28	0.40	MNS 88		
Actinomyces israelii		46.72	0.30	ARA 1		
23773421	**Good identification**					**23773421**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773423	**Good identification**					**23773423**
Actinomyces naeslundii		100.0	0.43	MNS 88		
23773440	**Good identification**					**23773440**
Actinomyces israelii		96.80	0.44	XYL 92		
Actinomyces naeslundii		3.20	0.27			
23773442	**Good identification**					**23773442**
Actinomyces israelii		97.85	0.47	XYL 92		
Actinomyces naeslundii		2.16	0.27			
23773500	**Very good identification**					**23773500**
Actinomyces israelii		95.08	0.61	XYL 92		
Actinomyces naeslundii		4.92	0.48			
23773501	**Low discrimination**					**23773501**
Actinomyces naeslundii		88.49	0.51	MNS 88		
Actinomyces israelii		11.51	0.27	ARA 1		
23773502	**Very good identification**					**23773502**
Actinomyces israelii		95.96	0.64	XYL 92		
Actinomyces naeslundii		3.31	0.48			
Bifidobacterium adolescentis		0.73	0.21			
23773503	**Low discrimination**					**23773503**
Actinomyces naeslundii		83.68	0.51	MNS 88		
Actinomyces israelii		16.32	0.30	ARA 1		
23773510	**Good identification**					**23773510**
Actinomyces israelii		98.21	0.47	XYL 92		
Actinomyces naeslundii		1.79	0.26	MLZ 13		
23773512	**Acceptable identification**					**23773512**
Actinomyces israelii		93.08	0.50	XYL 92		
Bifidobacterium adolescentis		5.79	0.23	MLZ 13		
Actinomyces naeslundii		1.13	0.26			
23773520	**Low discrimination**					**23773520**
Actinomyces israelii		55.21	0.40	XYL 92		
Actinomyces naeslundii		44.80	0.48	NAG 6		
23773521	**Good identification**					**23773521**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	MNS 88		
23773522	**Low discrimination**					**23773522**
Actinomyces israelii		64.90	0.43	XYL 92		
Actinomyces naeslundii		35.11	0.48	NAG 6		
23773523	**Good identification**					**23773523**
Actinomyces naeslundii		100.0	0.51	MNS 88		
23773540	**Good identification**					**23773540**
Actinomyces israelii		97.70	0.54	XYL 92		
Actinomyces naeslundii		2.30	0.35			
23773541	**Low discrimination**					**23773541**
Actinomyces naeslundii		77.73	0.38	MNS 88		
Actinomyces israelii		22.27	0.20	ARA 1		
23773542	**Very good identification**					**23773542**
Actinomyces israelii		98.46	0.57	XYL 92		
Actinomyces naeslundii		1.54	0.35			
23773543	**Low discrimination**					**23773543**
Actinomyces naeslundii		69.94	0.38	MNS 88		
Actinomyces israelii		30.06	0.23	ARA 1		
23773550	**Good identification**					**23773550**
Actinomyces israelii		100.0	0.40	XYL 92		
23773552	**Good identification**					**23773552**
Actinomyces israelii		98.06	0.43	XYL 92		
Bifidobacterium adolescentis		1.94	0.07	MLZ 13		
23773561	**Good identification**					**23773561**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23773562	**Low discrimination**					**23773562**
Actinomyces israelii		80.29	0.36	XYL 92		
Actinomyces naeslundii		19.71	0.35	NAG 6		
23773563	**Good identification**					**23773563**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23773700	**Good identification**					**23773700**
Actinomyces israelii		100.0	0.42	XYL 92		
23773702	**Good identification**					**23773702**
Actinomyces israelii		100.0	0.45	XYL 92		
23773742	**Good identification**					**23773742**
Actinomyces israelii		100.0	0.38	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23774102	**Acceptable identification**					**23774102**
Actinomyces israelii		92.03	0.36	GAL 99		
Actinomyces viscosus		7.97	0.03			
23775000	**Very good identification**					**23775000**
Actinomyces israelii		98.51	0.57			
Actinomyces naeslundii		1.49	0.35			
23775001	**Low discrimination**					**23775001**
Actinomyces naeslundii		69.19	0.38	MNS 88		
Actinomyces israelii		30.82	0.23	XYL 1		
23775002	**Very good identification**					**23775002**
Actinomyces israelii		99.00	0.60			
Actinomyces naeslundii		1.00	0.35			

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
23775003	**Low discrimination**					**23775003**
Actinomyces naeslundii		59.95	0.38	MNS 88		
Actinomyces israelii		40.05	0.26	XYL 1		
23775010	**Good identification**					**23775010**
Actinomyces israelii		100.0	0.43	MLZ 13		
23775012	**Good identification**					**23775012**
Actinomyces israelii		100.0	0.46	MLZ 13		
23775020	**Low discrimination**					**23775020**
Actinomyces israelii		80.85	0.37	NAG 6		
Actinomyces naeslundii		19.15	0.35			
23775021	**Good identification**					**23775021**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23775022	**Low discrimination**					**23775022**
Actinomyces israelii		86.36	0.40	NAG 6		
Actinomyces naeslundii		13.64	0.35			
23775023	**Good identification**					**23775023**
Actinomyces naeslundii		100.0	0.38	MNS 88		
23775040	**Good identification**					**23775040**
Actinomyces israelii		99.32	0.50			
Actinomyces naeslundii		0.68	0.22			
23775042	**Good identification**					**23775042**
Actinomyces israelii		99.55	0.53			
Actinomyces naeslundii		0.46	0.22			
23775050	**Good identification**					**23775050**
Actinomyces israelii		100.0	0.36	MLZ 13		
23775052	**Good identification**					**23775052**
Actinomyces israelii		100.0	0.39	MLZ 13		
23775100	**Very good identification**					**23775100**
					SPO	
Actinomyces israelii		98.74	0.68		1	
Actinomyces naeslundii		1.06	0.43		1	
Clostridium butyricum		0.20	0.01		99	
23775101	**Low discrimination**					**23775101**
Actinomyces naeslundii		61.50	0.46	MNS 88		
Actinomyces israelii		38.50	0.33	XYL 1		
23775102	**Very good identification**					**23775102**
Actinomyces israelii		98.82	0.71			
Actinomyces naeslundii		0.71	0.43			
Bifidobacterium adolescentis		0.37	0.23			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23775103	**Low discrimination**					**23775103**
Actinomyces naeslundii		51.57	0.46	MNS 88		
Actinomyces israelii		48.43	0.36	XYL 1		
23775110	**Good identification**					**23775110**
Actinomyces israelii		100.0	0.53	MLZ 13		
23775112	**Very good identification**					**23775112**
Actinomyces israelii		97.03	0.57	MLZ 13		
Bifidobacterium adolescentis		2.97	0.24			
23775120	**Low discrimination**					**23775120**
Actinomyces israelii		85.58	0.47	NAG 6		
Actinomyces naeslundii		14.42	0.43			
23775121	**Good identification**					**23775121**
Actinomyces naeslundii		100.0	0.46	MNS 88		
23775122	**Low discrimination**					**23775122**
					GRA	
Actinomyces israelii		88.19	0.50	NAG 6	99	
Actinomyces naeslundii		9.91	0.43		99	
Parabacteroides distasonis		1.90	0.18		1	
23775123	**Acceptable identification**					**23775123**
Actinomyces naeslundii		94.34	0.46	MNS 88		
Actinomyces israelii		5.66	0.16	XYL 1		
23775132	**Acceptable identification**					**23775132**
					GRA	
Actinomyces israelii		92.78	0.36	MLZ 13	99	
Parabacteroides distasonis		7.22	0.13	NAG 6	1	
23775140	**Very good identification**					**23775140**
Actinomyces israelii		99.51	0.61			
Actinomyces naeslundii		0.49	0.30			
23775142	**Very good identification**					**23775142**
Actinomyces israelii		99.56	0.64			
Actinomyces naeslundii		0.32	0.30			
Bifidobacterium adolescentis		0.12	0.07			
23775150	**Good identification**					**23775150**
Actinomyces israelii		100.0	0.46	MLZ 13		
23775152	**Good identification**					**23775152**
Actinomyces israelii		99.04	0.49	MLZ 13		
Bifidobacterium adolescentis		0.96	0.09			
23775160	**Acceptable identification**					**23775160**
Actinomyces israelii		92.90	0.40	NAG 6		
Actinomyces naeslundii		7.10	0.30			
23775162	**Good identification**					**23775162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		95.15	0.43	NAG 6		
Actinomyces naeslundii		4.85	0.30			
23775200	**Good identification**					**23775200**
Actinomyces israelii		100.0	0.38	RHA 7		
23775202	**Good identification**					**23775202**
Actinomyces israelii		100.0	0.41	RHA 7		
23775300	**Good identification**					**23775300**
Actinomyces israelii		100.0	0.48	RHA 7		
23775302	**Good identification**					**23775302**
Actinomyces israelii		100.0	0.51	RHA 7		
23775312	**Good identification**					**23775312**
Actinomyces israelii		100.0	0.37	RHA 7		
23775340	**Good identification**					**23775340**
Actinomyces israelii		100.0	0.41	RHA 7		
23775342	**Good identification**					**23775342**
Actinomyces israelii		100.0	0.44	RHA 7		
23775400	**Good identification**					**23775400**
					SPO	
Actinomyces israelii		99.62	0.56		1	
Clostridium butyricum		0.38	-.07		99	
23775402	**Very good identification**					**23775402**
Actinomyces israelii		100.0	0.59			
23775410	**Good identification**					**23775410**
Actinomyces israelii		100.0	0.42	MLZ 13		
23775412	**Good identification**					**23775412**
Actinomyces israelii		100.0	0.45	MLZ 13		
23775420	**Good identification**					**23775420**
Actinomyces israelii		100.0	0.35	NAG 6		
23775422	**Good identification**					**23775422**
Actinomyces israelii		100.0	0.38	NAG 6		
23775440	**Good identification**					**23775440**
Actinomyces israelii		100.0	0.49			
23775442	**Good identification**					**23775442**
Actinomyces israelii		100.0	0.52			
23775452	**Good identification**					**23775452**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.38	MLZ 13			
23775500	**Very good identification**						**23775500**
					GRA	SPO	
Actinomyces israelii		96.62	0.66		99	1	
Clostridium butyricum		2.19	0.17		99	99	
Mitsuokella multacida		1.19	0.21		1	1	
23775502	**Very good identification**						**23775502**
					SPO		
Actinomyces israelii		96.39	0.69		1		
Bifidobacterium adolescentis		2.95	0.37		1		
Clostridium butyricum		0.57	0.10		99		
23775510	**Good identification**						**23775510**
					GRA	SPO	
Actinomyces israelii		96.57	0.52	MLZ 13	99	1	
Mitsuokella multacida		2.66	0.13		1	1	
Clostridium butyricum		0.77	-.05		99	99	
23775512	**Low discrimination**						**23775512**
Actinomyces israelii		79.55	0.55	MLZ 13			
Bifidobacterium adolescentis		19.94	0.38				
Clostridium clostridioforme		0.50	0.08				
23775520	**Acceptable identification**						**23775520**
					GRA	SPO	
Actinomyces israelii		92.66	0.46	NAG 6	99	1	
Mitsuokella multacida		4.48	0.11		1	1	
Clostridium butyricum		2.86	-.01		99	99	
23775522	**Good identification**						**23775522**
					GRA		
Actinomyces israelii		98.41	0.49	NAG 6	99		
Parabacteroides distasonis		1.59	0.14		1		
23775540	**Very good identification**						**23775540**
					GRA	SPO	
Actinomyces israelii		96.45	0.59		99	1	
Mitsuokella multacida		3.06	0.21		1	1	
Clostridium butyricum		0.49	-.01		99	99	
23775542	**Very good identification**						**23775542**
					GRA		
Actinomyces israelii		98.81	0.62		99		
Bifidobacterium adolescentis		0.96	0.21		99		
Mitsuokella multacida		0.23	0.04		1		
23775550	**Acceptable identification**						**23775550**
					GRA		
Actinomyces israelii		93.39	0.45	MLZ 13	99		
Mitsuokella multacida		6.61	0.13		1		
23775552	**Acceptable identification**						**23775552**
Actinomyces israelii		92.62	0.48	MLZ 13			
Bifidobacterium adolescentis		7.38	0.23				
23775560	**Low discrimination**						**23775560**
					GRA		
Actinomyces israelii		88.94	0.39	NAG 6	99		
Mitsuokella multacida		11.06	0.11		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23775562	**Good identification**					**23775562**
Actinomyces israelii		100.0	0.42	NAG 6		
23775600	**Good identification**					**23775600**
Actinomyces israelii		100.0	0.37	RHA 7		
23775602	**Good identification**					**23775602**
Actinomyces israelii		100.0	0.40	RHA 7		
23775700	**Low discrimination**					**23775700**
					GRA	
Actinomyces israelii		76.65	0.47	RHA 7	99	
Mitsuokella multacida		23.35	0.25		1	
23775702	**Good identification**					**23775702**
					GRA	
Actinomyces israelii		95.90	0.50	RHA 7	99	
Mitsuokella multacida		2.16	0.09		1	
Clostridium clostridioforme		1.94	0.11		50	
23775712	**Low discrimination**					**23775712**
					GRA	
Actinomyces israelii		60.02	0.36	RHA 7	99	
Clostridium clostridioforme		36.96	0.23	MLZ 13	50	
Mitsuokella multacida		3.02	0.01		1	
23775740	**Low discrimination**					**23775740**
					GRA	
Actinomyces israelii		56.08	0.40	RHA 7	99	
Mitsuokella multacida		43.92	0.25		1	
23775742	**Acceptable identification**					**23775742**
					GRA	
Actinomyces israelii		94.52	0.43	RHA 7	99	
Mitsuokella multacida		5.48	0.09		1	
23776000	**Good identification**					**23776000**
Actinomyces israelii		100.0	0.36	GAL 99		
23776002	**Good identification**					**23776002**
Actinomyces israelii		100.0	0.39	GAL 99		
23776100	**Good identification**					**23776100**
Actinomyces israelii		98.50	0.46	GAL 99		
Actinomyces naeslundii		1.50	0.24			
23776102	**Good identification**					**23776102**
Actinomyces israelii		99.00	0.49	GAL 99		
Actinomyces naeslundii		1.00	0.24			
23776112	**Good identification**					**23776112**
Actinomyces israelii		100.0	0.35	GAL 99		
23776140	**Good identification**					**23776140**
Actinomyces israelii		100.0	0.39	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
23776142	**Good identification**					**23776142**
Actinomyces israelii		100.0	0.42	GAL 99		
23776402	**Good identification**					**23776402**
Actinomyces israelii		100.0	0.38	GAL 99		
23776500	**Good identification**					**23776500**
Actinomyces israelii		100.0	0.45	GAL 99		
23776502	**Good identification**					**23776502**
Actinomyces israelii		100.0	0.48	GAL 99		
23776540	**Good identification**					**23776540**
Actinomyces israelii		100.0	0.38	GAL 99		
23776542	**Good identification**					**23776542**
Actinomyces israelii		100.0	0.41	GAL 99		
23777000	**Very good identification**					**23777000**
Actinomyces israelii		99.49	0.70			
Actinomyces naeslundii		0.52	0.40			
23777001	**Low discrimination**					**23777001**
Actinomyces israelii		56.53	0.36	URE 1		
Actinomyces naeslundii		43.47	0.43			
23777002	**Very good identification**					**23777002**
Actinomyces israelii		99.66	0.73			
Actinomyces naeslundii		0.34	0.40			
23777003	**Low discrimination**					**23777003**
Actinomyces israelii		66.11	0.39	URE 1		
Actinomyces naeslundii		33.89	0.43			
23777010	**Good identification**					**23777010**
Actinomyces israelii		100.0	0.56	MLZ 13		
23777012	**Very good identification**					**23777012**
Actinomyces israelii		100.0	0.59	MLZ 13		
23777020	**Acceptable identification**					**23777020**
Actinomyces israelii		92.50	0.50	NAG 6		
Actinomyces naeslundii		7.51	0.40			
23777021	**Acceptable identification**					**23777021**
Actinomyces naeslundii		92.34	0.43	MNS 88		
Actinomyces israelii		7.67	0.16	XYL 1		
23777022	**Acceptable identification**					**23777022**
Actinomyces israelii		94.87	0.53	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		5.13	0.40			
23777023	**Low discrimination**					**23777023**
Actinomyces naeslundii		88.93	0.43	MNS 88		
Actinomyces israelii		11.07	0.19	XYL 1		
23777030	**Good identification**					**23777030**
Actinomyces israelii		100.0	0.36	MLZ 13		
23777032	**Good identification**					**23777032**
Actinomyces israelii		100.0	0.39	MLZ 13		
23777040	**Very good identification**					**23777040**
Actinomyces israelii		99.77	0.63			
Actinomyces naeslundii		0.23	0.27			
23777042	**Very good identification**					**23777042**
Actinomyces israelii		99.84	0.66			
Actinomyces naeslundii		0.16	0.27			
23777050	**Good identification**					**23777050**
Actinomyces israelii		100.0	0.49	MLZ 13		
23777052	**Good identification**					**23777052**
Actinomyces israelii		100.0	0.52	MLZ 13		
23777060	**Good identification**					**23777060**
Actinomyces israelii		96.45	0.43	NAG 6		
Actinomyces naeslundii		3.55	0.27			
23777062	**Good identification**					**23777062**
Actinomyces israelii		97.60	0.46	NAG 6		
Actinomyces naeslundii		2.40	0.27			
23777100	**Very good identification**					**23777100**
Actinomyces israelii		99.63	0.81			
Actinomyces naeslundii		0.37	0.48			
23777101	**Low discrimination**					**23777101**
Actinomyces israelii		64.63	0.46	URE 1		
Actinomyces naeslundii		35.37	0.51			
23777102	**Very good identification**					**23777102**
Actinomyces israelii		99.70	0.84			
Actinomyces naeslundii		0.25	0.48			
Bifidobacterium adolescentis		0.05	0.21			
23777103	**Low discrimination**					**23777103**
Actinomyces israelii		73.27	0.49	URE 1		
Actinomyces naeslundii		26.73	0.51			
23777110	**Very good identification**					**23777110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		99.87	0.66	MLZ 13		
Actinomyces naeslundii		0.13	0.26			
23777112	**Very good identification**					**23777112**
Actinomyces israelii		99.47	0.69	MLZ 13		
Bifidobacterium adolescentis		0.44	0.23			
Actinomyces naeslundii		0.09	0.26			
23777113	**Low discrimination**					**23777113**
Actinomyces israelii		88.61	0.35	MLZ 13		
Actinomyces naeslundii		11.39	0.29	URE 1		
23777120	**Good identification**					**23777120**
Actinomyces israelii		94.54	0.60	NAG 6		
Actinomyces naeslundii		5.46	0.48			
23777121	**Low discrimination**					**23777121**
Actinomyces naeslundii		89.55	0.51	MNS 88		
Actinomyces israelii		10.45	0.26	XYL 1		
23777122	**Very good identification**					**23777122**
Actinomyces israelii		96.29	0.63	NAG 6		
Actinomyces naeslundii		3.71	0.48			
23777123	**Low discrimination**					**23777123**
Actinomyces naeslundii		85.11	0.51	MNS 88		
Actinomyces israelii		14.89	0.29	XYL 1		
23777130	**Good identification**					**23777130**
Actinomyces israelii		98.01	0.46	MLZ 13		
Actinomyces naeslundii		1.99	0.26	NAG 6		
23777132	**Good identification**					**23777132**
Actinomyces israelii		98.66	0.49	MLZ 13		
Actinomyces naeslundii		1.34	0.26	NAG 6		
23777140	**Very good identification**					**23777140**
Actinomyces israelii		99.83	0.73			
Actinomyces naeslundii		0.17	0.35			
23777141	**Low discrimination**					**23777141**
Actinomyces israelii		80.11	0.39	URE 1		
Actinomyces naeslundii		19.89	0.38			
23777142	**Very good identification**					**23777142**
Actinomyces israelii		99.89	0.77			
Actinomyces naeslundii		0.11	0.35			
23777143	**Low discrimination**					**23777143**
Actinomyces israelii		85.80	0.42	URE 1		
Actinomyces naeslundii		14.20	0.38			
23777150	**Very good identification**					**23777150**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.59	MLZ 13		
23777152	**Very good identification**					**23777152**
Actinomyces israelii		99.86	0.62	MLZ 13		
Bifidobacterium adolescentis		0.14	0.07			
23777160	**Good identification**					**23777160**
Actinomyces israelii		97.45	0.53	NAG 6		
Actinomyces naeslundii		2.55	0.35			
23777161	**Low discrimination**					**23777161**
Actinomyces naeslundii		79.55	0.38	MNS 88		
Actinomyces israelii		20.45	0.19	XYL 1		
23777162	**Good identification**					**23777162**
Actinomyces israelii		98.28	0.56	NAG 6		
Actinomyces naeslundii		1.72	0.35			
23777163	**Low discrimination**					**23777163**
Actinomyces naeslundii		72.17	0.38	MNS 88		
Actinomyces israelii		27.83	0.22	XYL 1		
23777170	**Good identification**					**23777170**
Actinomyces israelii		100.0	0.39	MLZ 13		
23777172	**Good identification**					**23777172**
Actinomyces israelii		100.0	0.42	MLZ 13		
23777200	**Good identification**					**23777200**
Actinomyces israelii		100.0	0.51	RHA 7		
23777202	**Good identification**					**23777202**
Actinomyces israelii		100.0	0.54	RHA 7		
23777210	**Good identification**					**23777210**
Actinomyces israelii		100.0	0.37	RHA 7		
23777212	**Good identification**					**23777212**
Actinomyces israelii		100.0	0.40	RHA 7		
23777240	**Good identification**					**23777240**
Actinomyces israelii		100.0	0.44	RHA 7		
23777242	**Good identification**					**23777242**
Actinomyces israelii		100.0	0.47	RHA 7		
23777300	**Very good identification**					**23777300**
Actinomyces israelii		100.0	0.61	RHA 7		
23777302	**Very good identification**					**23777302**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	100.0	0.64	RHA 7		
23777310	**Good identification**					**23777310**
	Actinomyces israelii	100.0	0.47	RHA 7		
23777312	**Good identification**					**23777312**
	Actinomyces israelii	100.0	0.50	RHA 7		
23777320	**Good identification**					**23777320**
	Actinomyces israelii	100.0	0.41	RHA 7		
23777322	**Good identification**					**23777322**
	Actinomyces israelii	100.0	0.44	RHA 7		
23777340	**Good identification**					**23777340**
	Actinomyces israelii	100.0	0.54	RHA 7		
23777342	**Very good identification**					**23777342**
	Actinomyces israelii	100.0	0.57	RHA 7		
23777350	**Good identification**					**23777350**
	Actinomyces israelii	100.0	0.40	RHA 7		
23777352	**Good identification**					**23777352**
	Actinomyces israelii	100.0	0.43	RHA 7		
23777362	**Good identification**					**23777362**
	Actinomyces israelii	100.0	0.37	RHA 7		
23777400	**Very good identification**					**23777400**
	Actinomyces israelii	100.0	0.69			
23777402	**Very good identification**					**23777402**
	Actinomyces israelii	100.0	0.72			
23777403	**Good identification**					**23777403**
	Actinomyces israelii	100.0	0.38	URE 1		
23777410	**Good identification**					**23777410**
	Actinomyces israelii	100.0	0.55	MLZ 13		
23777412	**Very good identification**					**23777412**
	Actinomyces israelii	100.0	0.58	MLZ 13		
23777420	**Good identification**					**23777420**
	Actinomyces israelii	100.0	0.48	NAG 6		
23777422	**Good identification**					**23777422**
	Actinomyces israelii	100.0	0.51	NAG 6		
23777432	**Good identification**					**23777432**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.37	MLZ 13			
23777440	**Very good identification**						**23777440**
Actinomyces israelii		100.0	0.62				
23777442	**Very good identification**						**23777442**
Actinomyces israelii		100.0	0.65				
23777450	**Good identification**						**23777450**
Actinomyces israelii		100.0	0.48	MLZ 13			
23777452	**Good identification**						**23777452**
Actinomyces israelii		100.0	0.51	MLZ 13			
23777460	**Good identification**						**23777460**
Actinomyces israelii		100.0	0.41	NAG 6			
23777462	**Good identification**						**23777462**
Actinomyces israelii		100.0	0.44	NAG 6			
23777500	**Very good identification**						**23777500**
					GRA	SPO	
Actinomyces israelii		99.45	0.79		99	1	
Mitsuokella multacida		0.51	0.27		1	1	
Clostridium butyricum		0.04	0.01		99	99	
23777501	**Acceptable identification**						**23777501**
					GRA		
Actinomyces israelii		94.71	0.45	URE 1	99		
Mitsuokella multacida		5.30	0.11		1		
23777502	**Very good identification**						**23777502**
					GRA		
Actinomyces israelii		99.51	0.82		99		
Bifidobacterium adolescentis		0.44	0.35		99		
Mitsuokella multacida		0.04	0.11		1		
23777503	**Good identification**						**23777503**
Actinomyces israelii		100.0	0.48	URE 1			
23777510	**Very good identification**						**23777510**
					GRA		
Actinomyces israelii		98.88	0.65	MLZ 13	99		
Mitsuokella multacida		1.12	0.19		1		
23777512	**Very good identification**						**23777512**
Actinomyces israelii		96.32	0.68	MLZ 13			
Bifidobacterium adolescentis		3.49	0.37				
Bifidobacterium dentium		0.11	-.07				
23777520	**Very good identification**						**23777520**
					GRA		
Actinomyces israelii		98.05	0.59	NAG 6	99		
Mitsuokella multacida		1.95	0.17		1		
23777522	**Very good identification**						**23777522**
Actinomyces israelii		100.0	0.62	NAG 6			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
23777530	**Good identification**						**23777530**
					GRA		
Actinomyces israelii		95.75	0.44	MLZ 13	99		
Mitsuokella multacida		4.25	0.09	NAG 6	1		
23777532	**Good identification**						**23777532**
Actinomyces israelii		96.90	0.47	MLZ 13			
Bifidobacterium dentium		3.10	-.03	NAG 6			
23777540	**Very good identification**						**23777540**
					GRA		
Actinomyces israelii		98.71	0.72		99		
Mitsuokella multacida		1.29	0.27		1		
23777541	**Low discrimination**						**23777541**
					GRA		
Actinomyces israelii		87.43	0.38	URE 1	99		
Mitsuokella multacida		12.57	0.11		1		
23777542	**Very good identification**						**23777542**
					GRA		
Actinomyces israelii		99.76	0.75		99		
Bifidobacterium adolescentis		0.14	0.20		99		
Mitsuokella multacida		0.10	0.11		1		
23777543	**Good identification**						**23777543**
Actinomyces israelii		100.0	0.41	URE 1			
23777550	**Very good identification**						**23777550**
					COC	GRA	
Actinomyces israelii		96.76	0.58	MLZ 13	1	99	
Mitsuokella multacida		2.82	0.19		1	1	
Blautia producta		0.42	-.03		99	99	
23777552	**Very good identification**						**23777552**
					GRA		
Actinomyces israelii		98.65	0.61	MLZ 13	99		
Bifidobacterium adolescentis		1.14	0.21		99		
Mitsuokella multacida		0.21	0.03		1		
23777560	**Good identification**						**23777560**
					GRA		
Actinomyces israelii		95.13	0.52	NAG 6	99		
Mitsuokella multacida		4.87	0.17		1		
23777562	**Good identification**						**23777562**
Actinomyces israelii		100.0	0.55	NAG 6			
23777570	**Low discrimination**						**23777570**
					GRA		
Actinomyces israelii		89.75	0.37	MLZ 13	99		
Mitsuokella multacida		10.25	0.09	NAG 6	1		
23777572	**Good identification**						**23777572**
Actinomyces israelii		100.0	0.40	MLZ 13			
23777600	**Good identification**						**23777600**
Actinomyces israelii		100.0	0.49	RHA 7			
23777602	**Good identification**						**23777602**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.53	RHA 7			
23777610	**Good identification**						**23777610**
Actinomyces israelii		100.0	0.35	RHA 7			
23777612	**Good identification**						**23777612**
Actinomyces israelii		100.0	0.38	RHA 7			
23777640	**Good identification**						**23777640**
Actinomyces israelii		100.0	0.42	RHA 7			
23777642	**Good identification**						**23777642**
Actinomyces israelii		100.0	0.45	RHA 7			
23777700	**Low discrimination**						**23777700**
					GRA		
Actinomyces israelii		88.86	0.60	RHA 7	99		
Mitsuokella multacida		11.14	0.32		1		
23777702	**Very good identification**						**23777702**
					GRA		
Actinomyces israelii		99.08	0.63	RHA 7	99		
Mitsuokella multacida		0.92	0.15		1		
23777710	**Low discrimination**						**23777710**
					GRA		
Actinomyces israelii		78.14	0.46	RHA 7	99		
Mitsuokella multacida		21.86	0.24	MLZ 13	1		
23777712	**Good identification**						**23777712**
					GRA		
Actinomyces israelii		97.97	0.49	RHA 7	99		
Mitsuokella multacida		2.03	0.07	MLZ 13	1		
23777720	**Low discrimination**						**23777720**
					GRA		
Actinomyces israelii		67.06	0.39	RHA 7	99		
Mitsuokella multacida		32.94	0.21	NAG 6	1		
23777722	**Good identification**						**23777722**
					GRA		
Actinomyces israelii		96.49	0.42	RHA 7	99		
Mitsuokella multacida		3.51	0.05	NAG 6	1		
23777740	**Low discrimination**						**23777740**
					GRA		
Actinomyces israelii		75.62	0.53	RHA 7	99		
Mitsuokella multacida		24.38	0.32		1		
23777742	**Good identification**						**23777742**
					GRA		
Actinomyces israelii		97.67	0.56	RHA 7	99		
Mitsuokella multacida		2.33	0.15		1		
23777750	**Low discrimination**						**23777750**
					COC	GRA	
Actinomyces israelii		56.26	0.39	RHA 7	1	99	
Mitsuokella multacida		40.47	0.24	MLZ 13	1	1	
Blautia producta		3.27	-.03		99	99	
23777752	**Acceptable identification**						**23777752**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		94.94	0.42	RHA 7	99		
Mitsuokella multacida		5.06	0.07	MLZ 13	1		
23777762	**Acceptable identification**						**23777762**
					GRA		
Actinomyces israelii		91.45	0.35	RHA 7	99		
Mitsuokella multacida		8.56	0.05	NAG 6	1		
24000000	**Low discrimination**						**24000000**
					COC	GRA	
Veillonella parvula		47.47	0.66	MNS 1	99	1	
Campylobacter gracilis		34.87	0.66		1	1	
Eggerthella lenta		5.75	0.51		1	99	
24000001	**Low discrimination**						**24000001**
					COC	GRA	
Anaerococcus prevotii		54.55	0.66	NIT 1	99	99	
Campylobacter ureolyticus		42.28	0.60		1	1	
Veillonella parvula		1.42	0.32		99	1	
24000002	**Low discrimination**						**24000002**
					COC	GRA	
Campylobacter gracilis		64.57	0.50	MNS 1	1	1	
Anaerococcus prevotii		8.52	0.39	BGL 10	99	99	
Parvimonas micra		8.07	0.33		99	99	
24000003	**Good identification**						**24000003**
					COC	GRA	
Anaerococcus prevotii		95.55	0.56	NIT 1	99	99	
Campylobacter ureolyticus		2.99	0.26		1	1	
Campylobacter gracilis		0.81	0.15		1	1	
24000020	**Acceptable identification**						**24000020**
					COC	GRA	
Campylobacter gracilis		90.24	0.56	MNS 1	1	1	
Veillonella parvula		4.96	0.32		99	1	
Atopobium minutum		1.69	0.44		1	99	
24000022	**Low discrimination**						**24000022**
					GRA		
Campylobacter gracilis		73.80	0.39	MNS 1	1		
Pseudoflavonifractor capillosus		15.25	0.32	BGL 10	1		
Atopobium minutum		8.27	0.41		99		
24001002	**Low discrimination**						**24001002**
					COC	GRA	
Atopobium minutum		66.03	0.41	NIT 1	1	99	
Campylobacter gracilis		23.80	0.15		1	1	
Anaerococcus prevotii		3.14	0.05		99	99	
24010000	**Low discrimination**						**24010000**
					COC	GRA	
Fusobacterium varium		58.49	0.50	NIT 1	1	1	
Veillonella parvula		13.22	0.32		99	1	
Atopobium minutum		12.44	0.52		1	99	
24010002	**Low discrimination**						**24010002**
					GRA		
Atopobium minutum		79.55	0.49	NIT 1	99		
Campylobacter gracilis		10.35	0.15		1		
Fusobacterium varium		5.67	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
24010020	**Low discrimination**						**24010020**
					GRA	CAT	
Propionibacterium acnes		80.74	0.52	IND 87	99	95	
Atopobium minutum		10.51	0.44	GLU 99	99	1	
Campylobacter gracilis		6.16	0.21		1	1	
24010022	**Low discrimination**						**24010022**
					GRA	CAT	
Atopobium minutum		69.65	0.41	NIT 1	99	1	
Pseudoflavonifractor capillosus		15.45	0.15		1	1	
Propionibacterium acnes		8.11	0.18		99	95	
24010060	**Good identification**						**24010060**
					GRA	CAT	
Propionibacterium acnes		99.06	0.40	IND 87	99	95	
Atopobium minutum		0.59	0.10	GLU 99	99	1	
Campylobacter gracilis		0.35	-.13		1	1	
24011000	**Low discrimination**						**24011000**
					GRA	CAT	
Atopobium minutum		75.82	0.44	NIT 1	99	1	
Fusobacterium varium		10.80	0.15		1	1	
Propionibacterium acnes		5.88	0.18		99	95	
24011002	**Good identification**						**24011002**
Atopobium minutum		100.0	0.41	NIT 1			
24011020	**Good identification**						**24011020**
					GRA	CAT	
Propionibacterium acnes		95.77	0.52	IND 87	99	95	
Atopobium minutum		4.16	0.36	GLU 99	99	1	
Campylobacter gracilis		0.07	-.13		1	1	
24011060	**Good identification**						**24011060**
Propionibacterium acnes		100.0	0.40	IND 87			
24012020	**Good identification**						**24012020**
					CAT		
Propionibacterium acnes		99.47	0.47	IND 87	95		
Atopobium minutum		0.23	0.10	GLU 99	1		
Eubacterium limosum		0.16	-.06		1		
24012060	**Good identification**						**24012060**
Propionibacterium acnes		100.0	0.36	IND 87			
24013020	**Good identification**						**24013020**
Propionibacterium acnes		100.0	0.47	IND 87			
24013060	**Good identification**						**24013060**
Propionibacterium acnes		100.0	0.36	IND 87			
24111000	**Low discrimination**						**24111000**
					COC	GRA	
Prevotella intermedia		75.23	0.35	NIT 1	1	1	
Gemella morbillorum		22.33	0.27	GLU 90	99	99	
Prevotella disiens		1.34	-.03		1	1	
24111120	**Acceptable identification**						**24111120**
					GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		90.23	0.36	GLU 99	99		
Prevotella bivia		6.57	0.03	SUC 91	1		
Prevotella melaninogenica		1.69	-.01		1		
24131120	**Good identification**						**24131120**
Clostridium perfringens		97.37	0.40	GLU 99			
Clostridium septicum		2.63	0.07	SUC 91			
24400000	**Low discrimination**						**24400000**
					COC	GRA	
Anaerococcus prevotii		40.81	0.41	NIT 1	99	99	
Veillonella parvula		28.72	0.32	URE 90	99	1	
Campylobacter gracilis		21.10	0.32		1	1	
24400001	**Very good identification**						**24400001**
					COC	GRA	
Anaerococcus prevotii		97.56	0.58	NIT 1	99	99	
Campylobacter ureolyticus		2.29	0.26		1	1	
Veillonella parvula		0.08	-.03		99	1	
24400003	**Good identification**						**24400003**
					COC	GRA	
Anaerococcus prevotii		99.91	0.47	NIT 1	99	99	
Campylobacter ureolyticus		0.10	-.08		1	1	
24510000	**Good identification**						**24510000**
Prevotella intermedia		98.83	0.38	NIT 1			
Prevotella disiens		1.17	-.03	GLU 90			
24511000	**Good identification**						**24511000**
Prevotella intermedia		100.0	0.38	NIT 1			
25000000	**Low discrimination**						**25000000**
					COC		
Atopobium minutum		18.49	0.56	NIT 1	1		
Gemella morbillorum		18.10	0.52		99		
Peptostreptococcus anaerobius		14.35	0.42		99		
25000002	**Low discrimination**						**25000002**
					COC	GRA	
Atopobium minutum		86.29	0.53	NIT 1	1	99	
Campylobacter gracilis		6.09	0.15		1	1	
Gemella morbillorum		1.28	0.17		99	99	
25000020	**Low discrimination**						**25000020**
					SPO	CAT	
Propionibacterium acnes		63.65	0.52	IND 87	1	95	
Atopobium minutum		15.29	0.48	FRU 99	1	1	
Clostridium cadaveris		8.95	0.34		99	1	
25000022	**Low discrimination**						**25000022**
					GRA	CAT	
Atopobium minutum		87.66	0.45	NIT 1	99	1	
Propionibacterium acnes		5.53	0.18		99	95	
Campylobacter gracilis		4.64	0.05		1	1	
25000060	**Good identification**						**25000060**
					SPO	CAT	
Propionibacterium acnes		96.16	0.40	IND 87	1	95	
Clostridium novyi biovar A		1.08	0.18	FRU 99	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		1.06	0.14		1	1	
25000400	**Low discrimination**						**25000400**
					COC		
Gemella morbillorum		72.57	0.35	NIT 1	99		
Atopobium minutum		6.74	0.22	ARA 10	1		
Peptostreptococcus anaerobius		5.23	0.08		99		
25001000	**Low discrimination**						**25001000**
					COC	CAT	
Gemella morbillorum		58.06	0.47	NIT 1	99	1	
Atopobium minutum		36.72	0.48		1	1	
Propionibacterium acnes		1.55	0.18		1	95	
25001002	**Good identification**						**25001002**
					COC		
Atopobium minutum		97.66	0.45	NIT 1	1		
Gemella morbillorum		2.34	0.13		99		
25001020	**Low discrimination**						**25001020**
					COC	CAT	
Propionibacterium acnes		88.94	0.52	IND 87	1	95	
Atopobium minutum		7.12	0.40	FRU 99	1	1	
Gemella morbillorum		3.75	0.31		99	1	
25001022	**Low discrimination**						**25001022**
					CAT		
Atopobium minutum		84.08	0.37	NIT 1	1		
Propionibacterium acnes		15.92	0.18		95		
25001060	**Good identification**						**25001060**
					CAT		
Propionibacterium acnes		99.63	0.40	IND 87	95		
Atopobium minutum		0.37	0.06	FRU 99	1		
25002020	**Good identification**						**25002020**
					SPO	CAT	
Propionibacterium acnes		98.89	0.47	IND 87	1	95	
Atopobium minutum		0.43	0.14	FRU 99	1	1	
Clostridium cadaveris		0.25	0.00		99	1	
25002060	**Good identification**						**25002060**
Propionibacterium acnes		100.0	0.36	IND 87			
25003020	**Good identification**						**25003020**
					CAT		
Propionibacterium acnes		99.86	0.47	IND 87	95		
Atopobium minutum		0.14	0.06	FRU 99	1		
25003060	**Good identification**						**25003060**
Propionibacterium acnes		100.0	0.36	IND 87			
25004000	**Low discrimination**						**25004000**
					COC		
Gemella morbillorum		70.88	0.35	NIT 1	99		
Atopobium minutum		6.58	0.22	XYL 10	1		
Peptostreptococcus anaerobius		5.11	0.08		99		
25010000	**Low discrimination**						**25010000**
					GRA	CAT	
Fusobacterium varium		56.14	0.58	NIT 1	1	1	
Eubacterium limosum		13.00	0.49		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		9.13	0.52		99	95	
25010002	**Low discrimination**						**25010002**
					GRA		
Atopobium minutum		76.65	0.53	NIT 1	99		
Fusobacterium varium		10.74	0.24		1		
Pseudoramibacter alactolyticus		3.30	0.15		99		
25010010	**Low discrimination**						**25010010**
					GRA	SPO	
Clostridium difficile		76.86	0.41	NIT 1	99	99	
Fusobacterium varium		12.46	0.24	MAN 90	1	1	
Eubacterium limosum		2.88	0.15		99	1	
25010020	**Very good identification**						**25010020**
					SPO	CAT	
Propionibacterium acnes		99.42	0.86	IND 87	1	95	
Atopobium minutum		0.22	0.48		1	1	
Clostridium cadaveris		0.14	0.34		99	1	
25010021	**Good identification**						**25010021**
					SPO	CAT	
Propionibacterium acnes		99.54	0.52	IND 87	1	95	
Atopobium minutum		0.22	0.14	URE 1	1	1	
Clostridium cadaveris		0.14	0.00		99	1	
25010022	**Low discrimination**						**25010022**
					CAT		
Propionibacterium acnes		86.51	0.52	IND 87	95		
Atopobium minutum		12.79	0.45	BGL 1	1		
Pseudoramibacter alactolyticus		0.29	0.03		1		
25010030	**Good identification**						**25010030**
					SPO	CAT	
Propionibacterium acnes		99.03	0.52	IND 87	1	95	
Clostridium difficile		0.38	0.07	MLZ 1	99	1	
Atopobium minutum		0.22	0.14		1	1	
25010040	**Low discrimination**						**25010040**
					GRA	CAT	
Propionibacterium acnes		56.27	0.40	IND 87	99	95	
Fusobacterium varium		15.92	0.24	NAG 99	1	1	
Pseudoramibacter alactolyticus		11.02	0.21		99	1	
25010060	**Very good identification**						**25010060**
					CAT		
Propionibacterium acnes		99.94	0.75	IND 87	95		
Pseudoramibacter alactolyticus		0.04	0.09		1		
Atopobium minutum		0.01	0.14		1		
25010061	**Good identification**						**25010061**
Propionibacterium acnes		100.0	0.40	IND 87			
25010062	**Good identification**						**25010062**
					CAT		
Propionibacterium acnes		98.95	0.40	IND 87	95		
Atopobium minutum		0.67	0.11	BGL 1	1		
Pseudoramibacter alactolyticus		0.38	-.08		1		
25010070	**Good identification**						**25010070**
					SPO	CAT	
Propionibacterium acnes		99.57	0.40	IND 87	1	95	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar B		0.43	-.03	MLZ 1	99	1	
25010120	**Good identification**						**25010120**
					GRA	CAT	
Propionibacterium acnes		96.02	0.52	IND 87	99	95	
Prevotella bivia		3.46	0.15	LAC 1	1	1	
Atopobium minutum		0.22	0.14		99	1	
25010160	**Good identification**						**25010160**
Propionibacterium acnes		100.0	0.40	IND 87			
25010220	**Good identification**						**25010220**
					SPO	CAT	
Propionibacterium acnes		99.54	0.52	IND 87	1	95	
Atopobium minutum		0.22	0.14	RHA 1	1	1	
Clostridium cadaveris		0.14	0.00		99	1	
25010260	**Good identification**						**25010260**
Propionibacterium acnes		100.0	0.40	IND 87			
25010400	**Low discrimination**						**25010400**
					COC	GRA	
Gemella morbillorum		48.74	0.37	NIT 1	99	99	
Fusobacterium varium		31.61	0.24	ARA 10	1	1	
Eubacterium limosum		7.32	0.15		1	99	
25010420	**Good identification**						**25010420**
					COC	CAT	
Propionibacterium acnes		98.59	0.52	IND 87	1	95	
Gemella morbillorum		1.05	0.20	ARA 1	99	1	
Atopobium minutum		0.22	0.14		1	1	
25010460	**Good identification**						**25010460**
Propionibacterium acnes		100.0	0.40	IND 87			
25011000	**Low discrimination**						**25011000**
					COC	CAT	
Propionibacterium acnes		54.41	0.52	IND 87	1	95	
Gemella morbillorum		25.25	0.48	NAG 99	99	1	
Atopobium minutum		12.06	0.48		1	1	
25011002	**Low discrimination**						**25011002**
					CAT		
Atopobium minutum		81.42	0.45	NIT 1	1		
Propionibacterium acnes		5.57	0.18		95		
Propionibacterium granulosum		5.18	0.26		90		
25011020	**Very good identification**						**25011020**
					COC	CAT	
Propionibacterium acnes		99.85	0.86	IND 87	1	95	
Atopobium minutum		0.08	0.40		1	1	
Gemella morbillorum		0.05	0.32		99	1	
25011021	**Good identification**						**25011021**
					CAT		
Propionibacterium acnes		99.93	0.52	IND 87	95		
Atopobium minutum		0.08	0.05	URE 1	1		
25011022	**Acceptable identification**						**25011022**
					CAT		
Propionibacterium acnes		94.40	0.52	IND 87	95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		4.65	0.37	BGL 1	1		
Clostridium perfringens		0.53	0.15		1		
25011030	**Good identification**						**25011030**
					CAT		
Propionibacterium acnes		99.80	0.52	IND 87	95		
Propionibacterium granulosum		0.13	0.12	MLZ 1	90		
Atopobium minutum		0.07	0.05		1		
25011040	**Good identification**						**25011040**
					COC	CAT	
Propionibacterium acnes		96.94	0.40	IND 87	1	95	
Gemella morbillorum		2.07	0.14	NAG 99	99	1	
Atopobium minutum		0.99	0.14		1	1	
25011060	**Very good identification**						**25011060**
					CAT		
Propionibacterium acnes		99.99	0.75	IND 87	95		
Clostridium perfringens		0.01	0.08		1		
Atopobium minutum		0.00	0.05		1		
25011061	**Good identification**						**25011061**
Propionibacterium acnes		100.0	0.40	IND 87			
25011062	**Good identification**						**25011062**
					CAT		
Propionibacterium acnes		99.52	0.40	IND 87	95		
Clostridium perfringens		0.48	0.03	BGL 1	1		
25011070	**Good identification**						**25011070**
Propionibacterium acnes		100.0	0.40	IND 87			
25011100	**Low discrimination**						**25011100**
					GRA	CAT	
Fusobacterium mortiferum		54.99	0.40	NIT 1	1	5	
Lactobacillus fermentum		14.59	0.32		99	1	
Propionibacterium acnes		14.49	0.18		99	95	
25011120	**Acceptable identification**						**25011120**
					CAT	SPO	
Propionibacterium acnes		90.33	0.52	IND 87	95	1	
Clostridium perfringens		7.89	0.36	LAC 1	1	84	
Clostridium septicum		0.84	0.12		1	99	
25011160	**Acceptable identification**						**25011160**
					CAT		
Propionibacterium acnes		92.96	0.40	IND 87	95		
Clostridium perfringens		7.04	0.23	LAC 1	1		
25011220	**Good identification**						**25011220**
					CAT		
Propionibacterium acnes		99.93	0.52	IND 87	95		
Atopobium minutum		0.08	0.05	RHA 1	1		
25011260	**Good identification**						**25011260**
Propionibacterium acnes		100.0	0.40	IND 87			
25011420	**Good identification**						**25011420**
					COC	CAT	
Propionibacterium acnes		99.15	0.52	IND 87	1	95	
Gemella morbillorum		0.57	0.16	ARA 1	99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Lactobacillus fermentum	0.21	0.20		1	1	
25011460	**Good identification**						**25011460**
	Propionibacterium acnes	100.0	0.40	IND 87			
25012000	**Low discrimination**						**25012000**
					SPO	CAT	
	Eubacterium limosum	85.44	0.63	NIT 1	1	1	
	Clostridium difficile	6.90	0.46		99	1	
	Propionibacterium acnes	5.50	0.47		1	95	
25012010	**Very good identification**						**25012010**
					SPO	CAT	
	Clostridium difficile	97.22	0.58	NIT 1	99	1	
	Eubacterium limosum	2.49	0.29		1	1	
	Propionibacterium acnes	0.16	0.13		1	95	
25012020	**Very good identification**						**25012020**
					CAT		
	Propionibacterium acnes	99.78	0.82	IND 87	95		
	Eubacterium limosum	0.16	0.29		1		
	Pseudoramibacter alactolyticus	0.03	0.14		1		
25012021	**Good identification**						**25012021**
					CAT		
	Propionibacterium acnes	99.84	0.47	IND 87	95		
	Eubacterium limosum	0.16	-.06	URE 1	1		
25012022	**Good identification**						**25012022**
					CAT		
	Propionibacterium acnes	98.95	0.47	IND 87	95		
	Pseudoramibacter alactolyticus	0.32	-.02	BGL 1	1		
	Propionibacterium granulosum	0.31	0.14		90		
25012030	**Acceptable identification**						**25012030**
					SPO	CAT	
	Propionibacterium acnes	93.88	0.47	IND 87	1	95	
	Clostridium difficile	5.81	0.24	MLZ 1	99	1	
	Propionibacterium granulosum	0.16	0.09		1	90	
25012040	**Low discrimination**						**25012040**
					SPO	CAT	
	Propionibacterium acnes	43.45	0.36	IND 87	1	95	
	Eubacterium limosum	31.08	0.29	NAG 99	1	1	
	Clostridium difficile	13.08	0.24		99	1	
25012050	**Good identification**						**25012050**
					SPO	CAT	
	Clostridium difficile	98.84	0.36	NIT 1	99	1	
	Propionibacterium acnes	0.68	0.02	SOR 5	1	95	
	Eubacterium limosum	0.49	-.06		1	1	
25012060	**Very good identification**						**25012060**
					CAT		
	Propionibacterium acnes	99.96	0.70	IND 87	95		
	Pseudoramibacter alactolyticus	0.03	0.04		1		
	Eubacterium limosum	0.01	-.06		1		
25012061	**Good identification**						**25012061**
	Propionibacterium acnes	100.0	0.36	IND 87			
25012062	**Good identification**						**25012062**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		100.0	0.36	IND 87			
25012070	**Good identification**						**25012070**
					SPO	CAT	
Propionibacterium acnes		98.54	0.36	IND 87	1	95	
Clostridium difficile		1.46	0.02	MLZ 1	99	1	
25012120	**Good identification**						**25012120**
					CAT		
Propionibacterium acnes		99.84	0.47	IND 87	95		
Eubacterium limosum		0.16	-.06	LAC 1	1		
25012160	**Good identification**						**25012160**
Propionibacterium acnes		100.0	0.36	IND 87			
25012220	**Good identification**						**25012220**
					CAT		
Propionibacterium acnes		99.84	0.47	IND 87	95		
Eubacterium limosum		0.16	-.06	RHA 1	1		
25012260	**Good identification**						**25012260**
Propionibacterium acnes		100.0	0.36	IND 87			
25012420	**Good identification**						**25012420**
					CAT		
Propionibacterium acnes		99.84	0.47	IND 87	95		
Eubacterium limosum		0.16	-.06	ARA 1	1		
25012460	**Good identification**						**25012460**
Propionibacterium acnes		100.0	0.36	IND 87			
25013000	**Low discrimination**						**25013000**
					CAT		
Propionibacterium acnes		81.03	0.47	IND 87	95		
Eubacterium limosum		12.72	0.29	NAG 99	1		
Propionibacterium granulosum		4.09	0.35		90		
25013020	**Very good identification**						**25013020**
					CAT		
Propionibacterium acnes		99.98	0.82	IND 87	95		
Propionibacterium granulosum		0.01	0.24		90		
Atopobium minutum		0.00	0.05		1		
25013021	**Good identification**						**25013021**
Propionibacterium acnes		100.0	0.47	IND 87			
25013022	**Good identification**						**25013022**
Propionibacterium acnes		99.69	0.47	IND 87			
Propionibacterium granulosum		0.31	0.14	BGL 1			
25013030	**Good identification**						**25013030**
Propionibacterium acnes		99.83	0.47	IND 87			
Propionibacterium granulosum		0.17	0.09	MLZ 1			
25013040	**Good identification**						**25013040**
					CAT		
Propionibacterium acnes		99.28	0.36	IND 87	95		
Eubacterium limosum		0.72	-.06	NAG 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
25013060	**Very good identification**							**25013060**
Propionibacterium acnes		100.0	0.70	IND 87				
25013061	**Good identification**							**25013061**
Propionibacterium acnes		100.0	0.36	IND 87				
25013062	**Good identification**							**25013062**
Propionibacterium acnes		100.0	0.36	IND 87				
25013070	**Good identification**							**25013070**
Propionibacterium acnes		100.0	0.36	IND 87				
25013120	**Good identification**							**25013120**
					CAT			
Propionibacterium acnes		99.84	0.47	IND 87	95			
Clostridium perfringens		0.16	0.01	LAC 1	1			
25013160	**Good identification**							**25013160**
Propionibacterium acnes		100.0	0.36	IND 87				
25013220	**Good identification**							**25013220**
Propionibacterium acnes		100.0	0.47	IND 87				
25013260	**Good identification**							**25013260**
Propionibacterium acnes		100.0	0.36	IND 87				
25013420	**Good identification**							**25013420**
Propionibacterium acnes		100.0	0.47	IND 87				
25013460	**Good identification**							**25013460**
Propionibacterium acnes		100.0	0.36	IND 87				
25014020	**Good identification**							**25014020**
					COC	GRA	CAT	
Propionibacterium acnes		98.34	0.52	IND 87	1	99	95	
Gemella morbillorum		1.05	0.20	XYL 1	99	99	1	
Fusobacterium varium		0.25	0.01		1	1	1	
25014060	**Good identification**							**25014060**
					SPO	CAT		
Propionibacterium acnes		99.65	0.40	IND 87	1	95		
Clostridium glycolicum		0.36	-.03	XYL 1	99	1		
25015020	**Good identification**							**25015020**
					COC	CAT		
Propionibacterium acnes		99.24	0.52	IND 87	1	95		
Gemella morbillorum		0.57	0.16	XYL 1	99	1		
Lactobacillus fermentum		0.12	0.16		1	1		
25015060	**Good identification**							**25015060**
Propionibacterium acnes		100.0	0.40	IND 87				
25016010	**Good identification**							**25016010**
					SPO			
Clostridium difficile		99.51	0.36	NIT 1	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		0.49	-.06	XYL 5	1		
25016020	**Good identification**						**25016020**
					CAT		
Propionibacterium acnes		99.84	0.47	IND 87	95		
Eubacterium limosum		0.16	-.06	XYL 1	1		
25016060	**Good identification**						**25016060**
Propionibacterium acnes		100.0	0.36	IND 87			
25017020	**Good identification**						**25017020**
Propionibacterium acnes		100.0	0.47	IND 87			
25017060	**Good identification**						**25017060**
Propionibacterium acnes		100.0	0.36	IND 87			
25030000	**Low discrimination**						**25030000**
					GRA	CAT	
Propionibacterium granulosum		47.95	0.45	NIT 1	99	90	
Fusobacterium varium		26.40	0.24	SUC 93	1	1	
Eubacterium limosum		6.11	0.15		99	1	
25030020	**Good identification**						**25030020**
					SPO	CAT	
Propionibacterium acnes		96.27	0.52	IND 87	1	95	
Propionibacterium granulosum		2.71	0.35	TRE 1	1	90	
Clostridium cadaveris		0.71	0.12		99	1	
25030060	**Good identification**						**25030060**
					SPO	CAT	
Propionibacterium acnes		97.43	0.40	IND 87	1	95	
Clostridium botulinum biovar B		2.57	0.10	TRE 1	99	1	
25031000	**Low discrimination**						**25031000**
					GRA	CAT	
Propionibacterium granulosum		83.72	0.45	NIT 1	99	90	
Propionibacterium acnes		7.50	0.18	SUC 93	99	95	
Fusobacterium mortiferum		3.64	0.25		1	5	
25031020	**Good identification**						**25031020**
					CAT		
Propionibacterium acnes		95.14	0.52	IND 87	95		
Propionibacterium granulosum		2.68	0.35	TRE 1	90		
Clostridium perfringens		2.10	0.25		1		
25031060	**Good identification**						**25031060**
					CAT		
Propionibacterium acnes		98.12	0.40	IND 87	95		
Clostridium perfringens		1.88	0.13	TRE 1	1		
25031120	**Low discrimination**						**25031120**
					CAT	SPO	
Clostridium perfringens		81.09	0.40	SUC 91	1	84	
Clostridium septicum		13.86	0.20	MLT 99	1	99	
Propionibacterium acnes		5.05	0.18		95	1	
25032010	**Good identification**						**25032010**
					SPO	CAT	
Clostridium difficile		95.41	0.38	NIT 1	99	1	
Propionibacterium granulosum		4.26	0.28	TRE 7	1	90	
Eubacterium limosum		0.33	-.06		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
25032020	**Good identification**						**25032020**
					CAT		
Propionibacterium acnes		96.21	0.47	IND 87	95		
Propionibacterium granulosum		3.64	0.32	TRE 1	90		
Eubacterium limosum		0.15	-.06		1		
25032060	**Good identification**						**25032060**
Propionibacterium acnes		100.0	0.36	IND 87			
25033000	**Acceptable identification**						**25033000**
					CAT		
Propionibacterium granulosum		92.83	0.43	NIT 1	90		
Propionibacterium acnes		6.20	0.13	SUC 93	95		
Eubacterium limosum		0.97	-.06		1		
25033020	**Good identification**						**25033020**
Propionibacterium acnes		96.36	0.47	IND 87			
Propionibacterium granulosum		3.64	0.32	TRE 1			
25033060	**Good identification**						**25033060**
Propionibacterium acnes		100.0	0.36	IND 87			
25050020	**Good identification**						**25050020**
					SPO	CAT	
Propionibacterium acnes		99.54	0.52	IND 87	1	95	
Atopobium minutum		0.22	0.14	CEL 1	1	1	
Clostridium cadaveris		0.14	0.00		99	1	
25050060	**Good identification**						**25050060**
Propionibacterium acnes		100.0	0.40	IND 87			
25051020	**Good identification**						**25051020**
					CAT		
Propionibacterium acnes		99.93	0.52	IND 87	95		
Atopobium minutum		0.08	0.05	CEL 1	1		
25051060	**Good identification**						**25051060**
Propionibacterium acnes		100.0	0.40	IND 87			
25052010	**Good identification**						**25052010**
					SPO		
Clostridium difficile		99.51	0.36	NIT 1	99		
Eubacterium limosum		0.49	-.06	CEL 5	1		
25052020	**Good identification**						**25052020**
					CAT		
Propionibacterium acnes		99.84	0.47	IND 87	95		
Eubacterium limosum		0.16	-.06	CEL 1	1		
25052060	**Good identification**						**25052060**
Propionibacterium acnes		100.0	0.36	IND 87			
25053020	**Good identification**						**25053020**
Propionibacterium acnes		100.0	0.47	IND 87			
25053060	**Good identification**						**25053060**
Propionibacterium acnes		100.0	0.36	IND 87			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
25100000	**Low discrimination**						**25100000**
					COC	GRA	
Prevotella disiens		75.59	0.66	NIT 1	1	1	
Gemella morbillorum		19.35	0.64		99	99	
Peptostreptococcus anaerobius		1.81	0.39		99	99	
25100020	**Low discrimination**						**25100020**
					COC	GRA	
Gemella morbillorum		45.27	0.48	NIT 1	99	99	
Prevotella bivia		33.04	0.32	NAG 10	1	1	
Prevotella disiens		16.08	0.32		1	1	
25100120	**Very good identification**						**25100120**
					COC	GRA	
Prevotella bivia		99.98	0.66	NIT 1	1	1	
Gemella morbillorum		0.01	0.14		99	99	
Clostridium perfringens		0.00	0.01		1	99	
25100400	**Low discrimination**						**25100400**
					COC	GRA	
Gemella morbillorum		72.60	0.48	NIT 1	99	99	
Prevotella disiens		25.79	0.32	ARA 10	1	1	
Peptostreptococcus anaerobius		0.62	0.05		99	99	
25101000	**Good identification**						**25101000**
					COC	GRA	
Gemella morbillorum		92.13	0.60	NIT 1	99	99	
Prevotella disiens		6.75	0.32		1	1	
Prevotella intermedia		0.35	0.17		1	1	
25101020	**Low discrimination**						**25101020**
					COC	GRA	
Gemella morbillorum		74.56	0.43	NIT 1	99	99	
Prevotella bivia		17.83	0.19	NAG 10	1	1	
Clostridium perfringens		3.71	0.21		1	99	
25101101	**Very good identification**						**25101101**
Clostridium chauvoei		100.0	0.57	SUC 99			
25101103	**Good identification**						**25101103**
					SPO		
Clostridium chauvoei		98.44	0.44	SUC 99	99		
Actinomyces viscosus		1.56	0.12		1		
25101120	**Good identification**						**25101120**
					GRA	SPO	
Prevotella bivia		98.29	0.53	NIT 1	1	1	
Clostridium perfringens		1.52	0.36		99	84	
Clostridium chauvoei		0.10	0.14		99	99	
25101121	**Acceptable identification**						**25101121**
					GRA	SPO	
Clostridium chauvoei		90.40	0.48	SUC 99	99	99	
Prevotella bivia		9.46	0.19		1	1	
Clostridium perfringens		0.15	0.01		99	84	
25101123	**Good identification**						**25101123**
Clostridium chauvoei		100.0	0.36	SUC 99			
25101400	**Good identification**						**25101400**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		99.34	0.43	NIT 1	99	99		
Prevotella disiens		0.66	-.03	ARA 10	1	1		
25104000	**Low discrimination**							**25104000**
					COC	GRA	SPO	
Gemella morbillorum		72.25	0.48	NIT 1	99	99	1	
Prevotella disiens		25.66	0.32	XYL 10	1	1	1	
Clostridium bifermentans		0.61	0.09		1	99	99	
25105000	**Good identification**							**25105000**
					COC	GRA		
Gemella morbillorum		98.54	0.43	NIT 1	99	99		
Actinomyces odontolyticus		0.81	0.12	XYL 10	1	99		
Prevotella disiens		0.66	-.03		1	1		
25110000	**Low discrimination**							**25110000**
					COC	GRA		
Prevotella disiens		72.41	0.66	NIT 1	1	1		
Gemella morbillorum		22.65	0.66		99	99		
Prevotella intermedia		3.74	0.52		1	1		
25110002	**Low discrimination**							**25110002**
					COC	GRA		
Actinomyces viscosus		46.97	0.41	SUC 99	1	99		
Prevotella disiens		30.97	0.32	RAF 94	1	1		
Gemella morbillorum		9.69	0.32		99	99		
25110020	**Low discrimination**							**25110020**
					COC	GRA	CAT	
Propionibacterium acnes		56.04	0.52	IND 87	1	99	95	
Gemella morbillorum		30.41	0.50	MLT 1	99	99	1	
Prevotella disiens		8.84	0.32		1	1	1	
25110040	**Low discrimination**							**25110040**
					COC	GRA	SPO	
Clostridium botulinum biovar B		61.17	0.41	NIT 1	1	99	99	
Prevotella disiens		24.86	0.32	SUC 90	1	1	1	
Gemella morbillorum		7.78	0.32		99	99	1	
25110050	**Good identification**							**25110050**
					GRA	SPO		
Clostridium botulinum biovar B		98.61	0.36	NIT 1	99	99		
Prevotella disiens		0.82	-.03	SUC 90	1	1		
Clostridium novyi biovar A		0.57	0.12		99	99		
25110060	**Low discrimination**							**25110060**
					COC	SPO	CAT	
Propionibacterium acnes		52.27	0.40	IND 87	1	1	95	
Clostridium botulinum biovar B		45.50	0.36	MLT 1	1	99	1	
Gemella morbillorum		1.31	0.15		99	1	1	
25110102	**Acceptable identification**							**25110102**
Actinomyces viscosus		94.86	0.45	SUC 99				
Actinomyces odontolyticus		2.97	0.27	RAF 94				
Propionibacterium propionicum		1.54	0.29					
25110103	**Acceptable identification**							**25110103**
					SPO			
Actinomyces viscosus		94.92	0.37	SUC 99	1			
Clostridium chauvoei		4.57	0.16	RAF 94	99			
Propionibacterium propionicum		0.51	0.12		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
25110120	**Good identification**						**25110120**
					GRA	CAT	
Prevotella bivia		97.02	0.50	NIT 1	1	1	
Clostridium perfringens		2.38	0.36	FRU 10	99	1	
Propionibacterium acnes		0.28	0.18		99	95	
25110202	**Good identification**						**25110202**
					GRA		
Actinomyces odontolyticus		95.33	0.40	SUC 99	99		
Actinomyces viscosus		2.82	0.07	MNS 9	99		
Prevotella disiens		1.86	-.03		1		
25110300	**Good identification**						**25110300**
					GRA		
Actinomyces odontolyticus		96.36	0.35	SUC 99	99		
Prevotella disiens		3.64	-.03	MNS 9	1		
25110400	**Low discrimination**						**25110400**
					COC	GRA	
Gemella morbillorum		76.59	0.50	NIT 1	99	99	
Prevotella disiens		22.26	0.32	ARA 10	1	1	
Prevotella intermedia		1.15	0.17		1	1	
25111000	**Low discrimination**						**25111000**
					COC	GRA	
Gemella morbillorum		70.57	0.61	NIT 1	99	99	
Prevotella intermedia		21.61	0.52		1	1	
Prevotella disiens		4.23	0.32		1	1	
25111002	**Low discrimination**						**25111002**
					COC		
Actinomyces viscosus		58.05	0.38	SUC 99	1		
Actinomyces odontolyticus		11.12	0.32	RAF 94	1		
Gemella morbillorum		10.52	0.27		99		
25111020	**Low discrimination**						**25111020**
					COC	CAT	
Clostridium perfringens		47.02	0.55	SUC 91	1	1	
Propionibacterium acnes		39.89	0.52	LAC 88	1	95	
Gemella morbillorum		11.65	0.45		99	1	
25111022	**Good identification**						**25111022**
					CAT		
Clostridium perfringens		95.46	0.49	SUC 91	1		
Propionibacterium acnes		1.74	0.18	LAC 88	95		
Lactobacillus fermentum		1.07	0.28		1		
25111060	**Low discrimination**						**25111060**
					COC	CAT	
Clostridium perfringens		50.02	0.43	SUC 91	1	1	
Propionibacterium acnes		48.90	0.40	LAC 88	1	95	
Gemella morbillorum		0.66	0.11		99	1	
25111062	**Good identification**						**25111062**
					CAT		
Clostridium perfringens		97.95	0.37	SUC 91	1		
Propionibacterium acnes		2.06	0.06	LAC 88	95		
25111100	**Low discrimination**						**25111100**
					SPO		
Clostridium perfringens		65.28	0.44	SUC 91	84		
Clostridium chauvoei		9.43	0.29	NAG 97	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		7.45	0.40		1		
25111101	**Very good identification**						**25111101**
					SPO		
Clostridium chauvoei		99.87	0.63	SUC 99	99		
Clostridium perfringens		0.07	0.10		84		
Actinomyces naeslundii		0.03	0.26		1		
25111102	**Low discrimination**						**25111102**
Actinomyces viscosus		59.05	0.42	SUC 99			
Clostridium perfringens		28.51	0.38	RAF 94			
Lactobacillus fermentum		6.92	0.40				
25111103	**Low discrimination**						**25111103**
					SPO		
Clostridium chauvoei		88.22	0.50	SUC 99	99		
Actinomyces viscosus		11.35	0.34		1		
Actinomyces naeslundii		0.17	0.26		1		
25111120	**Good identification**						**25111120**
					GRA	SPO	
Clostridium perfringens		91.25	0.70	SUC 91	99	84	
Prevotella bivia		6.64	0.37		1	1	
Clostridium septicum		1.53	0.33		99	99	
25111121	**Acceptable identification**						**25111121**
					GRA	SPO	
Clostridium chauvoei		92.97	0.55	SUC 99	99	99	
Clostridium perfringens		6.37	0.36		99	84	
Prevotella bivia		0.46	0.03		1	1	
25111122	**Very good identification**						**25111122**
					GRA	CAT	
Clostridium perfringens		99.03	0.64	SUC 91	99	1	
Lactobacillus fermentum		0.25	0.32		99	1	
Capnocytophaga ochracea		0.22	0.15		1	90	
25111123	**Low discrimination**						**25111123**
					SPO		
Clostridium chauvoei		83.88	0.42	SUC 99	99		
Clostridium perfringens		15.32	0.30		84		
Actinomyces naeslundii		0.47	0.26		1		
25111130	**Acceptable identification**						**25111130**
					GRA	SPO	
Clostridium perfringens		91.79	0.36	SUC 91	99	84	
Prevotella bivia		6.68	0.03	MLZ 1	1	1	
Clostridium septicum		1.54	-.02		99	99	
25111160	**Very good identification**						**25111160**
					GRA	CAT	
Clostridium perfringens		99.39	0.57	SUC 91	99	1	
Prevotella bivia		0.38	0.03		1	1	
Propionibacterium acnes		0.13	0.06		99	95	
25111162	**Good identification**						**25111162**
					GRA	CAT	
Clostridium perfringens		99.87	0.52	SUC 91	99	1	
Capnocytophaga ochracea		0.13	-.01		1	90	
25111200	**Low discrimination**						**25111200**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		71.02	0.41	SUC 99	1	99	
Gemella morbillorum		21.21	0.27	MNS 9	99	99	
Prevotella intermedia		6.50	0.17		1	1	
25111202	**Good identification**						**25111202**
Actinomyces odontolyticus		97.36	0.37	SUC 99			
Actinomyces viscosus		2.64	0.03	MNS 9			
25111320	**Low discrimination**						**25111320**
					GRA		
Clostridium perfringens		87.58	0.36	SUC 91	99		
Prevotella bivia		6.37	0.03	RHA 1	1		
Actinomyces odontolyticus		4.59	0.19		99		
25111400	**Acceptable identification**						**25111400**
					COC	GRA	
Gemella morbillorum		91.13	0.45	NIT 1	99	99	
Lactobacillus fermentum		5.83	0.36	ARA 10	1	99	
Prevotella intermedia		2.54	0.17		1	1	
25111402	**Low discrimination**						**25111402**
					COC		
Lactobacillus fermentum		80.84	0.36	NIT 1	1		
Gemella morbillorum		12.76	0.11	SUC 87	99		
Actinomyces viscosus		6.40	0.03		1		
25111500	**Low discrimination**						**25111500**
					COC		
Lactobacillus fermentum		84.37	0.40	NIT 1	1		
Gemella morbillorum		8.16	0.11	SUC 87	99		
Clostridium perfringens		7.47	0.10		1		
25111502	**Low discrimination**						**25111502**
Lactobacillus fermentum		88.66	0.40	NIT 1			
Actinomyces viscosus		7.65	0.08	SUC 87			
Clostridium perfringens		3.69	0.04				
25111520	**Low discrimination**						**25111520**
					GRA		
Clostridium perfringens		82.92	0.36	SUC 91	99		
Lactobacillus fermentum		9.66	0.32	ARA 1	99		
Prevotella bivia		6.03	0.03		1		
25112020	**Good identification**						**25112020**
					COC	CAT	
Propionibacterium acnes		98.06	0.47	IND 87	1	95	
Gemella morbillorum		0.96	0.15	MLT 1	99	1	
Propionibacterium granulosum		0.56	0.18		1	90	
25112060	**Good identification**						**25112060**
					SPO	CAT	
Propionibacterium acnes		98.46	0.36	IND 87	1	95	
Clostridium botulinum biovar B		1.54	0.02	MLT 1	99	1	
25112100	**Very good identification**						**25112100**
					GRA		
Propionibacterium propionicum		99.46	0.60	SUC 99	99		
Prevotella disiens		0.35	-.03		1		
Eubacterium limosum		0.20	-.06		99		
25112101	**Good identification**						**25112101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.44	SUC 99		
25112102	**Very good identification**					**25112102**
Propionibacterium propionicum		99.38	0.63	SUC 99		
Actinomyces viscosus		0.63	0.11			
25112103	**Good identification**					**25112103**
Propionibacterium propionicum		98.15	0.47	SUC 99		
Actinomyces viscosus		1.85	0.03	URE 10		
25112140	**Very good identification**					**25112140**
Propionibacterium propionicum		100.0	0.60	SUC 99		
25112141	**Good identification**					**25112141**
Propionibacterium propionicum		100.0	0.44	SUC 99		
25112142	**Very good identification**					**25112142**
Propionibacterium propionicum		100.0	0.63	SUC 99		
25112143	**Good identification**					**25112143**
Propionibacterium propionicum		100.0	0.47	SUC 99		
25113020	**Good identification**					**25113020**
					CAT	
Propionibacterium acnes		96.89	0.47	IND 87	95	
Clostridium perfringens		2.05	0.21	MLT 1	1	
Propionibacterium granulosum		0.55	0.18		90	
25113060	**Good identification**					**25113060**
					CAT	
Propionibacterium acnes		98.20	0.36	IND 87	95	
Clostridium perfringens		1.80	0.08	MLT 1	1	
25113100	**Very good identification**					**25113100**
Propionibacterium propionicum		99.06	0.60	SUC 99		
Clostridium perfringens		0.59	0.10			
Actinomyces naeslundii		0.36	0.28			
25113101	**Low discrimination**					**25113101**
					SPO	
Propionibacterium propionicum		55.16	0.44	SUC 99	1	
Clostridium chauvoei		42.15	0.29	URE 10	99	
Actinomyces naeslundii		2.69	0.31		1	
25113102	**Very good identification**					**25113102**
Propionibacterium propionicum		99.19	0.63	SUC 99		
Actinomyces viscosus		0.38	0.08			
Actinomyces naeslundii		0.24	0.28			
25113103	**Low discrimination**					**25113103**
					SPO	
Propionibacterium propionicum		88.19	0.47	SUC 99	1	
Clostridium chauvoei		7.93	0.16	URE 10	99	
Actinomyces naeslundii		2.86	0.31		1	
25113120	**Low discrimination**					**25113120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Clostridium perfringens		81.57	0.36	SUC 91	99	1	
Prevotella bivia		5.94	0.03	MAN 1	1	1	
Propionibacterium acnes		5.31	0.13		99	95	
25113140	**Very good identification**						**25113140**
Propionibacterium propionicum		100.0	0.60	SUC 99			
25113141	**Good identification**						**25113141**
Propionibacterium propionicum		100.0	0.44	SUC 99			
25113142	**Very good identification**						**25113142**
Propionibacterium propionicum		100.0	0.63	SUC 99			
25113143	**Good identification**						**25113143**
Propionibacterium propionicum		100.0	0.47	SUC 99			
25114000	**Low discrimination**						**25114000**
					COC	GRA	
Gemella morbillorum		72.42	0.50	NIT 1	99	99	
Prevotella disiens		21.05	0.32	XYL 10	1	1	
Actinomyces odontolyticus		4.52	0.34		1	99	
25114200	**Acceptable identification**						**25114200**
					COC	GRA	
Actinomyces odontolyticus		90.28	0.38	SUC 99	1	99	
Gemella morbillorum		7.53	0.15	MNS 9	99	99	
Prevotella disiens		2.19	-.03		1	1	
25115000	**Low discrimination**						**25115000**
					COC		
Gemella morbillorum		86.89	0.45	NIT 1	99		
Actinomyces odontolyticus		6.71	0.31	XYL 10	1		
Lactobacillus fermentum		3.27	0.32		1		
25115100	**Low discrimination**						**25115100**
Lactobacillus fermentum		52.78	0.36	NIT 1			
Actinomyces odontolyticus		20.99	0.22	SUC 87			
Actinomyces meyeri		9.59	0.13				
25115102	**Low discrimination**						**25115102**
Lactobacillus fermentum		68.42	0.36	NIT 1			
Actinomyces odontolyticus		16.68	0.18	SUC 87			
Actinomyces viscosus		10.05	0.08				
25115120	**Low discrimination**						**25115120**
					GRA		
Clostridium perfringens		85.38	0.36	SUC 91	99		
Prevotella bivia		6.21	0.03	XYL 1	1		
Lactobacillus fermentum		5.84	0.28		99		
25115200	**Acceptable identification**						**25115200**
					COC		
Actinomyces odontolyticus		93.69	0.35	SUC 99	1		
Gemella morbillorum		6.31	0.11	MNS 9	99		
25115500	**Good identification**						**25115500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		95.66	0.36	NIT 1			
Actinomyces meyeri		4.34	0.02	SUC 87			
25115502	**Good identification**						**25115502**
Lactobacillus fermentum		100.0	0.36	NIT 1			
25120040	**Acceptable identification**						**25120040**
					GRA	SPO	
Clostridium botulinum biovar B		93.76	0.39	NIT 1	99	99	
Clostridium botulinum biovar A		5.73	0.21	SUC 90	99	99	
Prevotella disiens		0.51	-.03		1	1	
25121101	**Good identification**						**25121101**
Clostridium chauvoei		100.0	0.40	SUC 99			
25121120	**Low discrimination**						**25121120**
					GRA	SPO	
Clostridium perfringens		72.48	0.40	SUC 91	99	84	
Prevotella bivia		25.56	0.19	FRU 99	1	1	
Clostridium septicum		1.96	0.07		99	99	
25130040	**Good identification**						**25130040**
					GRA	SPO	
Clostridium botulinum biovar B		99.77	0.50	NIT 1	99	99	
Prevotella disiens		0.14	-.03	SUC 90	1	1	
Clostridium botulinum biovar A		0.10	0.01		99	99	
25130050	**Good identification**						**25130050**
Clostridium botulinum biovar B		100.0	0.44	NIT 1			
25130060	**Good identification**						**25130060**
					CAT	SPO	
Clostridium botulinum biovar B		98.89	0.44	NIT 1	1	99	
Clostridium perfringens		0.73	0.13	SUC 90	1	84	
Propionibacterium acnes		0.38	0.06		95	1	
25130070	**Good identification**						**25130070**
Clostridium botulinum biovar B		100.0	0.39	NIT 1			
25130120	**Low discrimination**						**25130120**
					GRA	SPO	
Clostridium perfringens		80.06	0.40	SUC 91	99	84	
Prevotella bivia		17.77	0.15	GAL 99	1	1	
Clostridium septicum		2.17	0.07		99	99	
25131020	**Very good identification**						**25131020**
					CAT		
Clostridium perfringens		98.64	0.59	SUC 91	1		
Propionibacterium granulosum		0.57	0.29	LAC 88	90		
Propionibacterium acnes		0.46	0.18		95		
25131022	**Good identification**						**25131022**
					CAT		
Clostridium perfringens		99.69	0.54	SUC 91	1		
Propionibacterium granulosum		0.31	0.18	LAC 88	90		
25131060	**Good identification**						**25131060**
					CAT	SPO	
Clostridium perfringens		98.12	0.47	SUC 91	1	84	
Clostridium botulinum biovar B		1.36	0.10	LAC 88	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		0.52	0.06		95	1	
25131062	**Good identification**						**25131062**
Clostridium perfringens		100.0	0.42	SUC 91			
25131100	**Acceptable identification**						**25131100**
					COC	SPO	
Clostridium perfringens		94.02	0.48	SUC 91	1	84	
Atopobium parvulum		3.05	0.22	NAG 97	99	1	
Clostridium septicum		0.83	0.07		1	99	
25131101	**Good identification**						**25131101**
					SPO		
Clostridium chauvoei		98.27	0.47	SUC 99	99		
Clostridium perfringens		1.16	0.14	TRE 10	84		
Actinomyces naeslundii		0.57	0.31		1		
25131102	**Low discrimination**						**25131102**
Clostridium perfringens		88.45	0.43	SUC 91			
Actinomyces viscosus		7.43	0.23	NAG 97			
Propionibacterium propionicum		2.14	0.27				
25131120	**Very good identification**						**25131120**
					GRA	SPO	
Clostridium perfringens		97.29	0.74	SUC 91	99	84	
Clostridium septicum		2.63	0.41		99	99	
Prevotella bivia		0.04	0.03		1	1	
25131121	**Low discrimination**						**25131121**
Clostridium perfringens		52.20	0.40	SUC 91			
Clostridium chauvoei		45.60	0.38	URE 1			
Clostridium septicum		1.41	0.07				
25131122	**Very good identification**						**25131122**
					SPO		
Clostridium perfringens		99.86	0.69	SUC 91	84		
Clostridium septicum		0.12	0.12		99		
Actinomyces naeslundii		0.02	0.28		1		
25131130	**Good identification**						**25131130**
Clostridium perfringens		97.37	0.40	SUC 91			
Clostridium septicum		2.63	0.07	MLZ 1			
25131140	**Good identification**						**25131140**
Clostridium perfringens		96.18	0.36	SUC 91			
Propionibacterium propionicum		3.82	0.24	NAG 97			
25131160	**Very good identification**						**25131160**
Clostridium perfringens		99.86	0.62	SUC 91			
Clostridium septicum		0.14	0.07				
25131162	**Very good identification**						**25131162**
Clostridium perfringens		100.0	0.56	SUC 91			
25131320	**Good identification**						**25131320**
Clostridium perfringens		97.37	0.40	SUC 91			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		2.63	0.07	RHA 1			
25131520	**Good identification**						**25131520**
Clostridium perfringens		97.37	0.40	SUC 91			
Clostridium septicum		2.63	0.07	ARA 1			
25132000	**Low discrimination**						**25132000**
					GRA	CAT	
Propionibacterium granulosum		85.96	0.37	NIT 1	99	90	
Propionibacterium propionicum		8.46	0.24	SUC 93	99	1	
Prevotella disiens		3.58	-.03		1	1	
25132100	**Very good identification**						**25132100**
Propionibacterium propionicum		100.0	0.58	SUC 99			
25132101	**Good identification**						**25132101**
Propionibacterium propionicum		100.0	0.42	SUC 99			
25132102	**Very good identification**						**25132102**
Propionibacterium propionicum		100.0	0.61	SUC 99			
25132103	**Good identification**						**25132103**
Propionibacterium propionicum		100.0	0.45	SUC 99			
25132140	**Very good identification**						**25132140**
Propionibacterium propionicum		100.0	0.58	SUC 99			
25132141	**Good identification**						**25132141**
Propionibacterium propionicum		100.0	0.42	SUC 99			
25132142	**Very good identification**						**25132142**
Propionibacterium propionicum		100.0	0.61	SUC 99			
25132143	**Good identification**						**25132143**
Propionibacterium propionicum		100.0	0.45	SUC 99			
25133000	**Low discrimination**						**25133000**
					CAT		
Propionibacterium granulosum		88.68	0.37	NIT 1	90		
Propionibacterium propionicum		8.72	0.24	SUC 93	1		
Actinomyces naeslundii		2.60	0.25		25		
25133100	**Very good identification**						**25133100**
Propionibacterium propionicum		97.84	0.58	SUC 99			
Clostridium perfringens		1.32	0.14				
Actinomyces naeslundii		0.84	0.33				
25133101	**Low discrimination**						**25133101**
					SPO		
Propionibacterium propionicum		82.01	0.42	SUC 99	1		
Actinomyces naeslundii		9.48	0.36	URE 10	1		
Clostridium chauvoei		8.51	0.12		99		
25133102	**Very good identification**						**25133102**
Propionibacterium propionicum		99.02	0.61	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.57	0.33				
Clostridium perfringens		0.42	0.09				
25133103	**Acceptable identification**						**25133103**
					SPO		
Propionibacterium propionicum		91.80	0.45	SUC 99	1		
Actinomyces naeslundii		7.08	0.36	URE 10	1		
Clostridium chauvoei		1.12	-.01		99		
25133120	**Acceptable identification**						**25133120**
					SPO		
Clostridium perfringens		93.47	0.40	SUC 91	84		
Clostridium septicum		2.53	0.07	MAN 1	99		
Propionibacterium propionicum		2.17	0.24		1		
25133121	**Low discrimination**						**25133121**
					SPO		
Actinomyces naeslundii		60.91	0.36	SUC 99	1		
Clostridium perfringens		20.86	0.06	RAF 92	84		
Clostridium chauvoei		18.23	0.04		99		
25133123	**Low discrimination**						**25133123**
Actinomyces naeslundii		86.12	0.36	SUC 99			
Clostridium perfringens		13.89	0.00	RAF 92			
25133140	**Very good identification**						**25133140**
Propionibacterium propionicum		99.74	0.58	SUC 99			
Clostridium perfringens		0.26	0.02				
25133141	**Good identification**						**25133141**
Propionibacterium propionicum		98.00	0.42	SUC 99			
Actinomyces naeslundii		2.00	0.23	URE 10			
25133142	**Very good identification**						**25133142**
Propionibacterium propionicum		100.0	0.61	SUC 99			
25133143	**Good identification**						**25133143**
Propionibacterium propionicum		98.66	0.45	SUC 99			
Actinomyces naeslundii		1.34	0.23	URE 10			
25135120	**Acceptable identification**						**25135120**
					COC	SPO	
Clostridium perfringens		94.38	0.40	SUC 91	1	84	
Atopobium parvulum		3.07	0.14	XYL 1	99	1	
Clostridium septicum		2.55	0.07		1	99	
25150102	**Good identification**						**25150102**
Actinomyces viscosus		100.0	0.36	SUC 99			
25151120	**Low discrimination**						**25151120**
					GRA	SPO	
Clostridium septicum		55.38	0.41	NIT 1	99	99	
Clostridium perfringens		43.52	0.46		99	84	
Prevotella bivia		0.77	0.03		1	1	
25151122	**Low discrimination**						**25151122**
					GRA	SPO	
Clostridium perfringens		90.00	0.40	SUC 91	99	84	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		4.97	0.12	CEL 4	99	99	
Prevotella oralis		3.26	0.12		1	1	
25171020	**Low discrimination**						**25171020**
					SPO		
Clostridium perfringens		80.92	0.36	SUC 91	84		
Clostridium septicum		12.32	0.15	CEL 4	99		
Lactobacillus jensenii		6.08	0.17	LAC 88	1		
25171120	**Low discrimination**						**25171120**
					COC	SPO	
Clostridium septicum		66.99	0.49	NIT 1	1	99	
Clostridium perfringens		32.58	0.51		1	84	
Atopobium parvulum		0.32	0.16		99	1	
25171122	**Acceptable identification**						**25171122**
					SPO		
Clostridium perfringens		90.93	0.45	SUC 91	84		
Clostridium septicum		8.11	0.20	CEL 4	99		
Actinomyces naeslundii		0.63	0.30		1		
25171160	**Acceptable identification**						**25171160**
Clostridium perfringens		90.17	0.38	SUC 91			
Clostridium septicum		9.83	0.15	CEL 4			
25173100	**Low discrimination**						**25173100**
Actinomyces naeslundii		52.91	0.35	SUC 99			
Propionibacterium propionicum		47.09	0.24	RAF 92			
25173101	**Good identification**						**25173101**
Actinomyces naeslundii		100.0	0.38	SUC 99			
25173103	**Good identification**						**25173103**
Actinomyces naeslundii		100.0	0.38	SUC 99			
25173121	**Good identification**						**25173121**
Actinomyces naeslundii		100.0	0.38	SUC 99			
25173122	**Low discrimination**						**25173122**
Actinomyces naeslundii		57.07	0.35	SUC 99			
Clostridium perfringens		42.94	0.11	RAF 92			
25173123	**Good identification**						**25173123**
Actinomyces naeslundii		100.0	0.38	SUC 99			
25210020	**Good identification**						**25210020**
					SPO	CAT	
Propionibacterium acnes		99.54	0.52	IND 87	1	95	
Atopobium minutum		0.22	0.14	SAL 1	1	1	
Clostridium cadaveris		0.14	0.00		99	1	
25210060	**Good identification**						**25210060**
Propionibacterium acnes		100.0	0.40	IND 87			
25210100	**Good identification**						**25210100**
Fusobacterium mortiferum		100.0	0.42	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
25211000	**Low discrimination**							**25211000**
					COC	GRA	CAT	
Fusobacterium mortiferum		64.03	0.39	NIT 1	1	1	5	
Propionibacterium acnes		20.27	0.18		1	99	95	
Gemella morbillorum		9.41	0.14		99	99	1	
25211020	**Good identification**							**25211020**
					CAT			
Propionibacterium acnes		99.63	0.52	IND 87	95			
Clostridium perfringens		0.30	0.10	SAL 1	1			
Atopobium minutum		0.07	0.05		1			
25211060	**Good identification**							**25211060**
Propionibacterium acnes		100.0	0.40	IND 87				
25211100	**Good identification**							**25211100**
Fusobacterium mortiferum		100.0	0.45	NIT 1				
25212000	**Low discrimination**							**25212000**
					SPO	CAT		
Clostridium difficile		67.42	0.37	NIT 1	99	1		
Eubacterium limosum		29.88	0.29		1	1		
Propionibacterium acnes		1.92	0.13		1	95		
25212010	**Good identification**							**25212010**
					SPO			
Clostridium difficile		99.91	0.48	NIT 1	99			
Eubacterium limosum		0.09	-.06		1			
25212020	**Good identification**							**25212020**
					SPO	CAT		
Propionibacterium acnes		99.49	0.47	IND 87	1	95		
Clostridium difficile		0.36	0.02	SAL 1	99	1		
Eubacterium limosum		0.16	-.06		1	1		
25212060	**Good identification**							**25212060**
Propionibacterium acnes		100.0	0.36	IND 87				
25213020	**Good identification**							**25213020**
Propionibacterium acnes		100.0	0.47	IND 87				
25213060	**Good identification**							**25213060**
Propionibacterium acnes		100.0	0.36	IND 87				
25231100	**Acceptable identification**							**25231100**
					GRA	SPO		
Fusobacterium mortiferum		90.35	0.36	NIT 1	1	1		
Clostridium perfringens		5.10	0.04		99	84		
Clostridium septicum		4.55	-.04		99	99		
25243000	**Good identification**							**25243000**
Clostridium innocuum		100.0	0.37	NIT 1				
25243002	**Good identification**							**25243002**
Clostridium innocuum		100.0	0.37	NIT 1				
25251100	**Low discrimination**							**25251100**
					GRA	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		79.44	0.37	NIT 1	1	1	
Lactobacillus catenaformis		16.77	0.15		99	1	
Clostridium septicum		3.79	-.03		99	99	
25252000	**Acceptable identification**						**25252000**
					SPO		
Clostridium innocuum		92.26	0.37	NIT 1	99		
Clostridium difficile		7.14	0.15	GAL 95	99		
Eubacterium limosum		0.61	-.06		1		
25252002	**Good identification**						**25252002**
Clostridium innocuum		100.0	0.37	NIT 1			
25253000	**Very good identification**						**25253000**
Clostridium innocuum		100.0	0.59	NIT 1			
25253002	**Very good identification**						**25253002**
Clostridium innocuum		100.0	0.59	NIT 1			
25253010	**Good identification**						**25253010**
Clostridium innocuum		100.0	0.35	NIT 1			
25253012	**Good identification**						**25253012**
Clostridium innocuum		100.0	0.35	NIT 1			
25253400	**Good identification**						**25253400**
Clostridium innocuum		100.0	0.46	NIT 1			
25253402	**Good identification**						**25253402**
Clostridium innocuum		100.0	0.46	NIT 1			
25257000	**Good identification**						**25257000**
Clostridium innocuum		100.0	0.37	NIT 1			
25257002	**Good identification**						**25257002**
Clostridium innocuum		100.0	0.37	NIT 1			
25271120	**Good identification**						**25271120**
Clostridium septicum		99.52	0.39	NIT 1			
Clostridium perfringens		0.48	0.06	MLT 94			
25273000	**Good identification**						**25273000**
Clostridium innocuum		100.0	0.51	NIT 1			
25273002	**Good identification**						**25273002**
Clostridium innocuum		100.0	0.51	NIT 1			
25273400	**Good identification**						**25273400**
Clostridium innocuum		100.0	0.38	NIT 1			
25273402	**Good identification**						**25273402**
Clostridium innocuum		100.0	0.38	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25310002	**Low discrimination**					**25310002**
					GRA	
Actinomyces odontolyticus		55.65	0.40	SUC 99	99	
Actinomyces viscosus		43.27	0.31	MNS 9	99	
Prevotella disiens		1.08	-.03		1	
25310100	**Low discrimination**					**25310100**
					GRA	
Actinomyces odontolyticus		82.61	0.35	SUC 99	99	
Fusobacterium mortiferum		12.03	0.26	MNS 9	1	
Prevotella disiens		3.12	-.03		1	
25310102	**Low discrimination**					**25310102**
Actinomyces viscosus		81.40	0.36	SUC 99		
Actinomyces odontolyticus		18.60	0.32	RAF 94		
25310200	**Good identification**					**25310200**
					GRA	
Actinomyces odontolyticus		99.39	0.49	SUC 99	99	
Prevotella disiens		0.61	-.03	MNS 9	1	
25310202	**Good identification**					**25310202**
Actinomyces odontolyticus		100.0	0.45	SUC 99		
25310220	**Good identification**					**25310220**
Actinomyces odontolyticus		100.0	0.35	SUC 99		
25310300	**Good identification**					**25310300**
Actinomyces odontolyticus		100.0	0.40	SUC 99		
25310302	**Good identification**					**25310302**
Actinomyces odontolyticus		97.77	0.36	SUC 99		
Actinomyces viscosus		2.23	0.01	MNS 9		
25311000	**Low discrimination**					**25311000**
					COC	
Actinomyces odontolyticus		59.90	0.41	SUC 99	1	
Gemella morbillorum		17.89	0.27	MNS 9	99	
Lactobacillus jensenii		6.15	0.19		1	
25311002	**Low discrimination**					**25311002**
Actinomyces odontolyticus		51.96	0.37	SUC 99		
Actinomyces viscosus		37.14	0.28	MNS 9		
Lactobacillus jensenii		5.81	0.16			
25311020	**Low discrimination**					**25311020**
					CAT	
Clostridium perfringens		88.92	0.45	SUC 91	1	
Actinomyces odontolyticus		4.37	0.27	LAC 88	10	
Propionibacterium acnes		3.05	0.18		95	
25311022	**Acceptable identification**					**25311022**
Clostridium perfringens		92.37	0.39	SUC 91		
Actinomyces odontolyticus		5.91	0.24	LAC 88		
Lactobacillus jensenii		1.01	0.06			
25311120	**Low discrimination**					**25311120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium perfringens		78.47	0.59	SUC 91	99	84	
Clostridium septicum		21.02	0.43		99	99	
Prevotella bivia		0.23	0.03		1	1	
25311122	**Good identification**						**25311122**
					GRA	SPO	
Clostridium perfringens		98.01	0.54	SUC 91	99	84	
Clostridium septicum		1.14	0.14		99	99	
Prevotella oralis		0.30	0.07		1	1	
25311160	**Good identification**						**25311160**
Clostridium perfringens		98.60	0.47	SUC 91			
Clostridium septicum		1.40	0.09				
25311162	**Good identification**						**25311162**
Clostridium perfringens		100.0	0.41	SUC 91			
25311200	**Good identification**						**25311200**
Actinomyces odontolyticus		100.0	0.46	SUC 99			
25311202	**Good identification**						**25311202**
Actinomyces odontolyticus		100.0	0.42	SUC 99			
25311300	**Good identification**						**25311300**
Actinomyces odontolyticus		100.0	0.37	SUC 99			
25313100	**Low discrimination**						**25313100**
Actinomyces naeslundii		55.89	0.38	SUC 99			
Propionibacterium propionicum		44.11	0.26	RAF 92			
25313101	**Good identification**						**25313101**
Actinomyces naeslundii		100.0	0.41	SUC 99			
25313103	**Acceptable identification**						**25313103**
Actinomyces naeslundii		91.94	0.41	SUC 99			
Propionibacterium propionicum		8.06	0.12	RAF 92			
25313121	**Good identification**						**25313121**
Actinomyces naeslundii		100.0	0.41	SUC 99			
25313123	**Good identification**						**25313123**
Actinomyces naeslundii		100.0	0.41	SUC 99			
25314000	**Acceptable identification**						**25314000**
					COC	GRA	
Actinomyces odontolyticus		90.28	0.38	SUC 99	1	99	
Gemella morbillorum		7.53	0.15	MNS 9	99	99	
Prevotella disiens		2.19	-.03		1	1	
25314200	**Good identification**						**25314200**
Actinomyces odontolyticus		100.0	0.43	SUC 99			
25314202	**Good identification**						**25314202**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.40	SUC 99			
25315000	**Acceptable identification**						**25315000**
					COC		
Actinomyces odontolyticus		93.69	0.35	SUC 99	1		
Gemella morbillorum		6.31	0.11	MNS 9	99		
25315200	**Good identification**						**25315200**
Actinomyces odontolyticus		100.0	0.40	SUC 99			
25315202	**Good identification**						**25315202**
Actinomyces odontolyticus		100.0	0.37	SUC 99			
25331020	**Good identification**						**25331020**
					SPO		
Clostridium perfringens		95.39	0.49	SUC 91	84		
Clostridium septicum		3.06	0.17	LAC 88	99		
Lactobacillus jensenii		1.19	0.17		1		
25331022	**Good identification**						**25331022**
Clostridium perfringens		97.50	0.44	SUC 91			
Lactobacillus jensenii		1.73	0.14	LAC 88			
Actinomyces naeslundii		0.77	0.30				
25331060	**Good identification**						**25331060**
Clostridium perfringens		100.0	0.37	SUC 91			
25331100	**Low discrimination**						**25331100**
					COC	SPO	
Clostridium perfringens		72.34	0.38	SUC 91	1	84	
Atopobium parvulum		12.44	0.24	NAG 97	99	1	
Clostridium septicum		10.23	0.17		1	99	
25331101	**Low discrimination**						**25331101**
					SPO		
Actinomyces naeslundii		61.16	0.41	SUC 99	1		
Clostridium chauvoei		30.06	0.12	RAF 92	99		
Clostridium perfringens		8.78	0.04		84		
25331103	**Low discrimination**						**25331103**
					SPO		
Actinomyces naeslundii		77.78	0.41	SUC 99	1		
Actinomyces viscosus		15.48	0.05	RAF 92	1		
Clostridium chauvoei		6.75	-.01		99		
25331120	**Low discrimination**						**25331120**
					SPO		
Clostridium perfringens		69.68	0.64	SUC 91	84		
Clostridium septicum		30.16	0.51		99		
Actinomyces naeslundii		0.10	0.38		1		
25331122	**Very good identification**						**25331122**
					SPO		
Clostridium perfringens		97.86	0.58	SUC 91	84		
Clostridium septicum		1.84	0.22		99		
Actinomyces naeslundii		0.30	0.38		1		
25331160	**Good identification**						**25331160**
					SPO		
Clostridium perfringens		97.63	0.52	SUC 91	84		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium septicum		2.24	0.17		99	
Actinomyces naeslundii		0.13	0.25		1	
25331162	**Good identification**					**25331162**
Clostridium perfringens		99.72	0.46	SUC 91		
Actinomyces naeslundii		0.29	0.25			
25333001	**Good identification**					**25333001**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25333003	**Good identification**					**25333003**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25333021	**Good identification**					**25333021**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25333023	**Good identification**					**25333023**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25333100	**Low discrimination**					**25333100**
Actinomyces naeslundii		69.27	0.43	SUC 99		
Propionibacterium propionicum		23.05	0.24	RAF 92		
Clostridium perfringens		7.69	0.04			
25333101	**Good identification**					**25333101**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25333102	**Low discrimination**					**25333102**
Actinomyces naeslundii		63.59	0.43	SUC 99		
Propionibacterium propionicum		31.74	0.27	RAF 92		
Actinomyces israelii		4.68	0.14			
25333103	**Good identification**					**25333103**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25333121	**Good identification**					**25333121**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25333123	**Good identification**					**25333123**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25351000	**Low discrimination**					**25351000**
Lactobacillus jensenii		83.20	0.41	NIT 1		
Lactobacillus catenaformis		15.07	0.29	SUC 90		
Actinomyces naeslundii		0.67	0.27			
25351002	**Low discrimination**					**25351002**
Lactobacillus jensenii		71.30	0.38	NIT 1		
Lactobacillus catenaformis		19.37	0.29	SUC 90		
Actinomyces viscosus		6.38	0.18			
25351101	**Good identification**					**25351101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	SUC 99		
25351120	**Acceptable identification**					**25351120**
Clostridium septicum		93.51	0.51	NIT 1		
Clostridium perfringens		4.59	0.36			
Clostridium paraputrificum		0.85	0.14			
25351123	**Good identification**					**25351123**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25353001	**Good identification**					**25353001**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25353003	**Good identification**					**25353003**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25353021	**Good identification**					**25353021**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25353023	**Good identification**					**25353023**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25353100	**Low discrimination**					**25353100**
Actinomyces naeslundii		79.77	0.40	SUC 99		
Actinomyces israelii		20.23	0.20	RAF 92		
25353101	**Good identification**					**25353101**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25353102	**Low discrimination**					**25353102**
Actinomyces naeslundii		65.66	0.40	SUC 99		
Actinomyces israelii		24.98	0.23	RAF 92		
Lactobacillus acidophilus		9.36	0.01			
25353103	**Good identification**					**25353103**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25353121	**Good identification**					**25353121**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25353122	**Low discrimination**					**25353122**
					SPO	
Actinomyces naeslundii		70.22	0.40	SUC 99	1	
Clostridium barati		11.25	-.01	RAF 92	99	
Lactobacillus acidophilus		10.00	0.01		1	
25353123	**Good identification**					**25353123**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25357100	**Good identification**					**25357100**
Actinomyces israelii		100.0	0.39	SUC 98		
25357102	**Good identification**					**25357102**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii	100.0	0.42	SUC 98		
25357500 Good identification					**25357500**
Actinomyces israelii	100.0	0.37	SUC 98		
25357502 Good identification					**25357502**
Actinomyces israelii	100.0	0.40	SUC 98		
25371000 Good identification					**25371000**
Lactobacillus jensenii	99.48	0.50	NIT 1		
Actinomyces naeslundii	0.52	0.32	SUC 90		
25371002 Good identification					**25371002**
Lactobacillus jensenii	97.32	0.47	NIT 1		
Lactobacillus acidophilus	1.71	0.14	SUC 90		
Actinomyces naeslundii	0.76	0.32			
25371003 Low discrimination					**25371003**
Actinomyces naeslundii	53.71	0.35	SUC 99		
Lactobacillus jensenii	46.29	0.12	RAF 92		
25371010 Good identification					**25371010**
Lactobacillus jensenii	100.0	0.41	NIT 1		
25371012 Good identification					**25371012**
Lactobacillus jensenii	100.0	0.38	NIT 1		
25371020 Low discrimination					**25371020**
				SPO	
Lactobacillus jensenii	61.62	0.39	NIT 1	1	
Clostridium septicum	26.30	0.25	SUC 90	99	
Clostridium perfringens	10.80	0.26		84	
25371021 Low discrimination					**25371021**
Actinomyces naeslundii	75.57	0.35	SUC 99		
Lactobacillus jensenii	24.43	0.05	RAF 92		
25371022 Low discrimination					**25371022**
Lactobacillus jensenii	80.77	0.36	NIT 1		
Clostridium perfringens	9.99	0.20	SUC 90		
Lactobacillus acidophilus	5.66	0.14			
25371023 Low discrimination					**25371023**
Actinomyces naeslundii	82.27	0.35	SUC 99		
Lactobacillus jensenii	17.73	0.02	RAF 92		
25371101 Good identification					**25371101**
Actinomyces naeslundii	100.0	0.43	SUC 99		
25371103 Good identification					**25371103**
Actinomyces naeslundii	100.0	0.43	SUC 99		
25371120 Very good identification					**25371120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium septicum		96.67	0.60	NIT 1	1	99	
Clostridium perfringens		2.94	0.40		1	84	
Atopobium parvulum		0.16	0.18		99	1	
25371123	**Low discrimination**						**25371123**
					SPO		
Actinomyces naeslundii		85.72	0.43	SUC 99	1		
Clostridium septicum		8.40	-.04	RAF 92	99		
Clostridium perfringens		5.89	0.01		84		
25373001	**Good identification**						**25373001**
Actinomyces naeslundii		100.0	0.40	SUC 99			
25373003	**Good identification**						**25373003**
Actinomyces naeslundii		100.0	0.40	SUC 99			
25373020	**Low discrimination**						**25373020**
Actinomyces naeslundii		80.01	0.37	SUC 99			
Lactobacillus jensenii		19.99	0.05	RAF 92			
25373021	**Good identification**						**25373021**
Actinomyces naeslundii		100.0	0.40	SUC 99			
25373022	**Low discrimination**						**25373022**
Actinomyces naeslundii		85.72	0.37	SUC 99			
Lactobacillus jensenii		14.28	0.02	RAF 92			
25373023	**Good identification**						**25373023**
Actinomyces naeslundii		100.0	0.40	SUC 99			
25373100	**Low discrimination**						**25373100**
					SPO		
Actinomyces naeslundii		64.35	0.45	SUC 99	1		
Actinomyces israelii		31.63	0.30	RAF 92	1		
Clostridium tertium		4.03	0.01		99		
25373101	**Good identification**						**25373101**
Actinomyces naeslundii		100.0	0.48	SUC 99			
25373102	**Low discrimination**						**25373102**
Actinomyces naeslundii		46.07	0.45	SUC 99			
Actinomyces israelii		33.96	0.33	RAF 92			
Lactobacillus acidophilus		11.32	0.10				
25373103	**Good identification**						**25373103**
Actinomyces naeslundii		100.0	0.48	SUC 99			
25373121	**Good identification**						**25373121**
Actinomyces naeslundii		100.0	0.48	SUC 99			
25373122	**Low discrimination**						**25373122**
					SPO		
Actinomyces naeslundii		37.80	0.45	SUC 99	1		
Clostridium ramosum		26.75	0.17	RAF 92	92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		21.29	0.17		99		
25373123	**Good identification**						**25373123**
Actinomyces naeslundii		100.0	0.48	SUC 99			
25375100	**Low discrimination**						**25375100**
					COC		
Atopobium parvulum		86.10	0.35	NIT 1	99		
Actinomyces israelii		13.91	0.35	SUC 96	1		
25375102	**Low discrimination**						**25375102**
					COC		
Actinomyces israelii		57.86	0.39	SUC 98	1		
Atopobium parvulum		42.15	0.22		99		
25375502	**Good identification**						**25375502**
Actinomyces israelii		100.0	0.37	SUC 98			
25377000	**Good identification**						**25377000**
Actinomyces israelii		100.0	0.38	SUC 98			
25377002	**Good identification**						**25377002**
Actinomyces israelii		100.0	0.41	SUC 98			
25377100	**Good identification**						**25377100**
					COC	SPO	
Actinomyces israelii		97.44	0.48	SUC 98	1	1	
Clostridium tertium		1.49	0.03		1	99	
Atopobium parvulum		1.08	0.01		99	1	
25377102	**Good identification**						**25377102**
					SPO		
Actinomyces israelii		97.03	0.51	SUC 98	1		
Clostridium tertium		2.97	0.11		99		
25377112	**Low discrimination**						**25377112**
					SPO		
Actinomyces israelii		73.33	0.37	SUC 98	1		
Clostridium tertium		26.67	0.15	MLZ 13	99		
25377140	**Good identification**						**25377140**
Actinomyces israelii		100.0	0.41	SUC 98			
25377142	**Good identification**						**25377142**
Actinomyces israelii		100.0	0.44	SUC 98			
25377400	**Good identification**						**25377400**
Actinomyces israelii		100.0	0.37	SUC 98			
25377402	**Good identification**						**25377402**
Actinomyces israelii		100.0	0.40	SUC 98			
25377500	**Good identification**						**25377500**
Actinomyces israelii		100.0	0.47	SUC 98			
25377502	**Good identification**						**25377502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.50	SUC 98			
25377512	**Good identification**						**25377512**
Actinomyces israelii		100.0	0.36	SUC 98			
25377540	**Good identification**						**25377540**
Actinomyces israelii		100.0	0.40	SUC 98			
25377542	**Good identification**						**25377542**
Actinomyces israelii		100.0	0.43	SUC 98			
25410020	**Good identification**						**25410020**
					CAT		
Propionibacterium acnes		98.37	0.52	IND 87	95		
Propionibacterium granulosum		1.18	0.28	RAF 1	90		
Atopobium minutum		0.22	0.14		1		
25410060	**Good identification**						**25410060**
Propionibacterium acnes		100.0	0.40	IND 87			
25410100	**Good identification**						**25410100**
Fusobacterium mortiferum		100.0	0.42	NIT 1			
25411000	**Low discrimination**						**25411000**
					GRA	CAT	
Propionibacterium granulosum		40.15	0.38	NIT 1	99	90	
Fusobacterium mortiferum		26.78	0.39	SUC 93	1	5	
Prevotella intermedia		10.24	0.20		1	1	
25411020	**Good identification**						**25411020**
					CAT		
Propionibacterium acnes		98.20	0.52	IND 87	95		
Propionibacterium granulosum		1.17	0.28	RAF 1	90		
Lactobacillus fermentum		0.33	0.24		1		
25411060	**Good identification**						**25411060**
Propionibacterium acnes		100.0	0.40	IND 87			
25411100	**Low discrimination**						**25411100**
					GRA		
Fusobacterium mortiferum		81.77	0.45	NIT 1	1		
Lactobacillus fermentum		18.23	0.35		99		
25411500	**Good identification**						**25411500**
					GRA		
Lactobacillus fermentum		95.67	0.35	NIT 1	99		
Fusobacterium mortiferum		4.34	0.11	SUC 87	1		
25411502	**Good identification**						**25411502**
Lactobacillus fermentum		100.0	0.35	NIT 1			
25412020	**Good identification**						**25412020**
					CAT		
Propionibacterium acnes		98.27	0.47	IND 87	95		
Propionibacterium granulosum		1.58	0.26	RAF 1	90		
Eubacterium limosum		0.16	-.06		1		
25412060	**Good identification**						**25412060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		100.0	0.36	IND 87			
25413000	**Low discrimination**						**25413000**
					CAT		
Propionibacterium granulosum		84.59	0.36	NIT 1	90		
Propionibacterium acnes		13.32	0.13	SUC 93	95		
Eubacterium limosum		2.09	-.06		1		
25413020	**Good identification**						**25413020**
Propionibacterium acnes		98.42	0.47	IND 87			
Propionibacterium granulosum		1.58	0.26	RAF 1			
25413060	**Good identification**						**25413060**
Propionibacterium acnes		100.0	0.36	IND 87			
25430000	**Good identification**						**25430000**
					GRA	CAT	
Propionibacterium granulosum		95.76	0.47	NIT 1	99	90	
Fusobacterium mortiferum		4.24	0.27	SUC 93	1	5	
25430002	**Acceptable identification**						**25430002**
Propionibacterium granulosum		92.64	0.36	NIT 1			
Actinomyces viscosus		7.36	0.08	SUC 93			
25430020	**Low discrimination**						**25430020**
Propionibacterium granulosum		78.03	0.36	NIT 1			
Propionibacterium acnes		21.97	0.18	SUC 93			
25431000	**Acceptable identification**						**25431000**
					GRA	CAT	
Propionibacterium granulosum		93.77	0.47	NIT 1	99	90	
Fusobacterium mortiferum		6.23	0.30	SUC 93	1	5	
25431002	**Acceptable identification**						**25431002**
					GRA	CAT	
Propionibacterium granulosum		92.75	0.36	NIT 1	99	90	
Actinomyces viscosus		4.51	0.05	SUC 93	99	76	
Fusobacterium mortiferum		2.74	0.13		1	5	
25431020	**Low discrimination**						**25431020**
					CAT		
Propionibacterium granulosum		71.48	0.36	NIT 1	90		
Propionibacterium acnes		20.13	0.18	SUC 93	95		
Clostridium perfringens		8.39	0.13		1		
25431100	**Low discrimination**						**25431100**
					GRA	CAT	
Fusobacterium mortiferum		86.03	0.36	NIT 1	1	5	
Propionibacterium granulosum		5.61	0.13		99	90	
Actinomyces naeslundii		4.66	0.26		99	25	
25432000	**Good identification**						**25432000**
					CAT		
Propionibacterium granulosum		99.18	0.45	NIT 1	90		
Eubacterium limosum		0.82	-.06	SUC 93	1		
25433000	**Good identification**						**25433000**
Propionibacterium granulosum		99.25	0.45	NIT 1			
Actinomyces naeslundii		0.75	0.23	SUC 93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
25435322	**Good identification**						**25435322**
Parabacteroides distasonis		100.0	0.39	NIT 1			
25435722	**Good identification**						**25435722**
Parabacteroides distasonis		100.0	0.36	NIT 1			
25451100	**Good identification**						**25451100**
					GRA		
Fusobacterium mortiferum		96.79	0.37	NIT 1	1		
Actinomyces naeslundii		3.21	0.23		99		
25473101	**Good identification**						**25473101**
Actinomyces naeslundii		100.0	0.36	SUC 99			
25473103	**Good identification**						**25473103**
Actinomyces naeslundii		100.0	0.36	SUC 99			
25473121	**Good identification**						**25473121**
Actinomyces naeslundii		100.0	0.36	SUC 99			
25473123	**Good identification**						**25473123**
Actinomyces naeslundii		100.0	0.36	SUC 99			
25500002	**Good identification**						**25500002**
					GRA		
Actinomyces viscosus		99.20	0.40	SUC 99	99		
Prevotella disiens		0.80	-.03	FRU 95	1		
25500102	**Good identification**						**25500102**
Actinomyces viscosus		100.0	0.44	SUC 99			
25500103	**Good identification**						**25500103**
Actinomyces viscosus		100.0	0.36	SUC 99			
25501002	**Good identification**						**25501002**
Actinomyces viscosus		100.0	0.36	SUC 99			
25501102	**Acceptable identification**						**25501102**
					COC		
Actinomyces viscosus		93.81	0.40	SUC 99	1		
Blautia hansenii		6.20	0.15	FRU 95	99		
25510000	**Low discrimination**						**25510000**
					COC	GRA	
Prevotella intermedia		81.67	0.55	NIT 1	1	1	
Prevotella disiens		10.66	0.32		1	1	
Gemella morbillorum		3.34	0.32		99	99	
25510002	**Very good identification**						**25510002**
					GRA	CAT	
Actinomyces viscosus		99.46	0.62	SUC 99	99	76	
Prevotella intermedia		0.32	0.20		1	1	
Propionibacterium granulosum		0.14	0.22		99	90	
25510003	**Good identification**						**25510003**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		100.0	0.53	SUC 99			
25510102	**Very good identification**						**25510102**
Actinomyces viscosus		99.93	0.66	SUC 99			
Propionibacterium propionicum		0.07	0.26				
25510103	**Very good identification**						**25510103**
Actinomyces viscosus		100.0	0.58	SUC 99			
25511000	**Acceptable identification**						**25511000**
					COC	GRA	
Prevotella intermedia		91.03	0.55	NIT 1	1	1	
Lactobacillus fermentum		2.30	0.40		1	99	
Gemella morbillorum		2.00	0.27		99	99	
25511002	**Very good identification**						**25511002**
					GRA		
Actinomyces viscosus		97.63	0.58	SUC 99	99		
Lactobacillus fermentum		1.30	0.40		99		
Prevotella intermedia		0.52	0.20		1		
25511003	**Good identification**						**25511003**
Actinomyces viscosus		99.34	0.50	SUC 99			
Actinomyces naeslundii		0.67	0.37				
25511020	**Low discrimination**						**25511020**
					GRA		
Clostridium perfringens		83.13	0.43	SUC 91	99		
Prevotella intermedia		4.52	0.20	LAC 88	1		
Lactobacillus fermentum		3.77	0.32		99		
25511021	**Low discrimination**						**25511021**
Actinomyces naeslundii		69.33	0.37	SUC 99			
Clostridium perfringens		30.67	0.08				
25511022	**Low discrimination**						**25511022**
Clostridium perfringens		72.44	0.37	SUC 91			
Actinomyces viscosus		15.90	0.24	LAC 88			
Lactobacillus fermentum		6.97	0.32				
25511102	**Very good identification**						**25511102**
Actinomyces viscosus		97.63	0.62	SUC 99			
Lactobacillus fermentum		1.19	0.44				
Clostridium perfringens		0.57	0.26				
25511103	**Good identification**						**25511103**
					SPO		
Actinomyces viscosus		98.45	0.54	SUC 99	1		
Actinomyces naeslundii		1.06	0.45		1		
Clostridium chauvoei		0.49	0.16		99		
25511120	**Good identification**						**25511120**
					GRA		
Clostridium perfringens		90.32	0.57	SUC 91	99		
Prevotella melaninogenica		7.27	0.32		1		
Lactobacillus fermentum		0.91	0.35		99		
25511122	**Low discrimination**						**25511122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Clostridium perfringens		80.01	0.52	SUC 91	99	1	
Capnocytophaga ochracea		8.22	0.32		1	90	
Actinomyces viscosus		4.26	0.28		99	76	
25511123	**Low discrimination**						**25511123**
Actinomyces naeslundii		37.19	0.45	SUC 99			
Actinomyces viscosus		35.03	0.20				
Clostridium perfringens		19.94	0.18				
25511160	**Good identification**						**25511160**
					GRA		
Clostridium perfringens		98.73	0.45	SUC 91	99		
Actinomyces naeslundii		0.54	0.29		99		
Prevotella melaninogenica		0.42	-.03		1		
25511162	**Acceptable identification**						**25511162**
					GRA	CAT	
Clostridium perfringens		93.32	0.39	SUC 91	99	1	
Capnocytophaga ochracea		5.59	0.15		1	90	
Actinomyces naeslundii		1.09	0.29		99	25	
25511400	**Low discrimination**						**25511400**
					COC	GRA	
Lactobacillus fermentum		66.82	0.40	NIT 1	1	99	
Prevotella intermedia		26.72	0.20	SUC 87	1	1	
Gemella morbillorum		6.46	0.11		99	99	
25511402	**Low discrimination**						**25511402**
Lactobacillus fermentum		56.81	0.40	NIT 1			
Actinomyces viscosus		43.19	0.24	SUC 87			
25511500	**Good identification**						**25511500**
Lactobacillus fermentum		97.50	0.44	NIT 1			
Bifidobacterium longum subsp. longum		2.50	0.09	SUC 87			
25511502	**Low discrimination**						**25511502**
					GRA	CAT	
Lactobacillus fermentum		53.04	0.44	NIT 1	99	1	
Actinomyces viscosus		43.93	0.28	SUC 87	99	76	
Capnocytophaga ochracea		1.66	0.02		1	90	
25511522	**Low discrimination**						**25511522**
					GRA	CAT	
Lactobacillus fermentum		38.08	0.35	NIT 1	99	1	
Capnocytophaga ochracea		32.24	0.19	SUC 87	1	90	
Clostridium perfringens		17.96	0.18		99	1	
25512100	**Very good identification**						**25512100**
Propionibacterium propionicum		99.74	0.57	SUC 99			
Actinomyces naeslundii		0.26	0.23				
25512101	**Good identification**						**25512101**
Propionibacterium propionicum		96.62	0.41	SUC 99			
Actinomyces naeslundii		3.38	0.26	URE 10			
25512102	**Low discrimination**						**25512102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		86.99	0.60	SUC 99		
Actinomyces viscosus		12.86	0.32			
Actinomyces naeslundii		0.15	0.23			
25512103	**Low discrimination**					**25512103**
Propionibacterium propionicum		68.17	0.44	SUC 99		
Actinomyces viscosus		30.24	0.24	URE 10		
Actinomyces naeslundii		1.59	0.26			
25512140	**Very good identification**					**25512140**
Propionibacterium propionicum		100.0	0.57	SUC 99		
25512141	**Good identification**					**25512141**
Propionibacterium propionicum		100.0	0.41	SUC 99		
25512142	**Very good identification**					**25512142**
Propionibacterium propionicum		100.0	0.60	SUC 99		
25512143	**Good identification**					**25512143**
Propionibacterium propionicum		100.0	0.44	SUC 99		
25513001	**Good identification**					**25513001**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25513003	**Low discrimination**					**25513003**
Actinomyces naeslundii		56.28	0.42	SUC 99		
Actinomyces viscosus		43.72	0.16			
25513020	**Low discrimination**					**25513020**
Actinomyces naeslundii		35.60	0.39	SUC 99		
Propionibacterium acnes		30.48	0.13			
Propionibacterium granulosum		21.75	0.20			
25513021	**Good identification**					**25513021**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25513022	**Low discrimination**					**25513022**
					CAT	
Actinomyces naeslundii		76.13	0.39	SUC 99	25	
Clostridium perfringens		12.25	0.03		1	
Propionibacterium granulosum		11.63	0.10		90	
25513023	**Good identification**					**25513023**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25513100	**Good identification**					**25513100**
Propionibacterium propionicum		94.14	0.57	SUC 99		
Actinomyces naeslundii		5.86	0.46			
25513101	**Low discrimination**					**25513101**
Propionibacterium propionicum		54.33	0.41	SUC 99		
Actinomyces naeslundii		45.67	0.49	URE 10		
25513102	**Low discrimination**					**25513102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		88.33	0.60	SUC 99			
Actinomyces viscosus		8.01	0.28				
Actinomyces naeslundii		3.67	0.46				
25513103	**Low discrimination**						**25513103**
Propionibacterium propionicum		54.58	0.44	SUC 99			
Actinomyces naeslundii		30.58	0.49	URE 10			
Actinomyces viscosus		14.84	0.20				
25513120	**Low discrimination**						**25513120**
Actinomyces naeslundii		47.31	0.46	SUC 99			
Clostridium perfringens		41.66	0.23				
Propionibacterium propionicum		7.68	0.23				
25513121	**Good identification**						**25513121**
Actinomyces naeslundii		100.0	0.49	SUC 99			
25513122	**Low discrimination**						**25513122**
Actinomyces naeslundii		58.81	0.46	SUC 99			
Clostridium perfringens		24.37	0.18				
Propionibacterium propionicum		14.31	0.26				
25513123	**Good identification**						**25513123**
Actinomyces naeslundii		100.0	0.49	SUC 99			
25513140	**Very good identification**						**25513140**
Propionibacterium propionicum		98.91	0.57	SUC 99			
Actinomyces naeslundii		1.09	0.34				
25513141	**Low discrimination**						**25513141**
Propionibacterium propionicum		87.08	0.41	SUC 99			
Actinomyces naeslundii		12.92	0.37	URE 10			
25513142	**Very good identification**						**25513142**
Propionibacterium propionicum		99.27	0.60	SUC 99			
Actinomyces naeslundii		0.73	0.34				
25513143	**Acceptable identification**						**25513143**
Propionibacterium propionicum		91.00	0.44	SUC 99			
Actinomyces naeslundii		9.00	0.37	URE 10			
25513161	**Good identification**						**25513161**
Actinomyces naeslundii		100.0	0.37	SUC 99			
25513163	**Good identification**						**25513163**
Actinomyces naeslundii		100.0	0.37	SUC 99			
25515000	**Low discrimination**						**25515000**
					COC	GRA	
Lactobacillus fermentum		54.19	0.36	NIT 1	1	99	
Prevotella intermedia		36.89	0.20	SUC 87	1	1	
Gemella morbillorum		8.92	0.11		99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
25515100	**Good identification**						**25515100**
Lactobacillus fermentum		100.0	0.40	NIT 1			
25515322	**Good identification**						**25515322**
Parabacteroides distasonis		96.82	0.37	NIT 1			
Prevotella oralis		3.18	-.01				
25515400	**Good identification**						**25515400**
Lactobacillus fermentum		100.0	0.36	NIT 1			
25515402	**Good identification**						**25515402**
Lactobacillus fermentum		100.0	0.36	NIT 1			
25515500	**Low discrimination**						**25515500**
Lactobacillus fermentum		88.42	0.40	NIT 1			
Bifidobacterium longum subsp. longum		11.58	0.17	SUC 87			
25515502	**Low discrimination**						**25515502**
Lactobacillus fermentum		88.42	0.40	NIT 1			
Bifidobacterium longum subsp. longum		11.58	0.17	SUC 87			
25515720	**Good identification**						**25515720**
Bacteroides vulgatus		100.0	0.36	NIT 1			
25530000	**Low discrimination**						**25530000**
					GRA	CAT	
Propionibacterium granulosum		78.72	0.41	NIT 1	99	90	
Prevotella intermedia		14.90	0.20	SUC 93	1	1	
Actinomyces viscosus		3.48	0.08		99	76	
25530002	**Acceptable identification**						**25530002**
Actinomyces viscosus		94.59	0.42	SUC 99			
Propionibacterium granulosum		5.41	0.30	TRE 7			
25530102	**Good identification**						**25530102**
Actinomyces viscosus		99.10	0.47	SUC 99			
Propionibacterium propionicum		0.74	0.24	TRE 7			
Actinomyces naeslundii		0.16	0.23				
25530103	**Good identification**						**25530103**
Actinomyces viscosus		99.30	0.38	SUC 99			
Actinomyces naeslundii		0.70	0.26	TRE 7			
25531000	**Low discrimination**						**25531000**
					GRA	CAT	
Propionibacterium granulosum		65.04	0.41	NIT 1	99	90	
Clostridium perfringens		13.01	0.21	SUC 93	99	1	
Prevotella intermedia		12.31	0.20		1	1	
25531001	**Good identification**						**25531001**
Actinomyces naeslundii		100.0	0.42	SUC 99			
25531002	**Low discrimination**						**25531002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		85.71	0.39	SUC 99		
Propionibacterium granulosum		7.99	0.30	TRE 7		
Actinomyces naeslundii		3.29	0.39			
25531020	**Acceptable identification**					**25531020**
					CAT	
Clostridium perfringens		94.83	0.47	SUC 91	1	
Propionibacterium granulosum		3.67	0.30	LAC 88	90	
Actinomyces naeslundii		1.51	0.39		25	
25531021	**Low discrimination**					**25531021**
Actinomyces naeslundii		70.27	0.42	SUC 99		
Clostridium perfringens		29.74	0.13			
25531022	**Acceptable identification**					**25531022**
					CAT	
Clostridium perfringens		94.05	0.42	SUC 91	1	
Actinomyces naeslundii		3.18	0.39	LAC 88	25	
Propionibacterium granulosum		1.93	0.20		90	
25531023	**Low discrimination**					**25531023**
Actinomyces naeslundii		83.39	0.42	SUC 99		
Clostridium perfringens		16.61	0.07			
25531100	**Low discrimination**					**25531100**
Clostridium perfringens		77.64	0.36	SUC 91		
Actinomyces naeslundii		15.50	0.46	NAG 97		
Actinomyces viscosus		2.55	0.09			
25531101	**Low discrimination**					**25531101**
					SPO	
Actinomyces naeslundii		81.85	0.49	SUC 99	1	
Clostridium chauvoei		12.40	0.12		99	
Actinomyces viscosus		2.99	0.01		1	
25531102	**Low discrimination**					**25531102**
Actinomyces viscosus		82.07	0.43	SUC 99		
Clostridium perfringens		11.89	0.31	TRE 7		
Actinomyces naeslundii		5.04	0.46			
25531120	**Very good identification**					**25531120**
					SPO	
Clostridium perfringens		99.20	0.62	SUC 91	84	
Actinomyces naeslundii		0.61	0.46		1	
Clostridium septicum		0.14	0.07		99	
25531122	**Very good identification**					**25531122**
					GRA	
Clostridium perfringens		97.99	0.56	SUC 91	99	
Actinomyces naeslundii		1.29	0.46		99	
Parabacteroides distasonis		0.37	0.24		1	
25531123	**Low discrimination**					**25531123**
Actinomyces naeslundii		64.53	0.49	SUC 99		
Clostridium perfringens		33.11	0.22			
Actinomyces viscosus		2.36	0.01			
25531141	**Good identification**					**25531141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	SUC 99		
25531143	**Low discrimination**					**25531143**
Actinomyces naeslundii		82.86	0.37	SUC 99		
Actinomyces viscosus		17.15	0.01			
25531160	**Good identification**					**25531160**
Clostridium perfringens		99.43	0.50	SUC 91		
Actinomyces naeslundii		0.57	0.34			
25531162	**Good identification**					**25531162**
Clostridium perfringens		98.80	0.44	SUC 91		
Actinomyces naeslundii		1.20	0.34			
25531163	**Low discrimination**					**25531163**
Actinomyces naeslundii		64.36	0.37	SUC 99		
Clostridium perfringens		35.64	0.10			
25532000	**Good identification**					**25532000**
					CAT	
Propionibacterium granulosum		95.10	0.39	NIT 1	90	
Propionibacterium propionicum		4.90	0.21	SUC 93	1	
25532100	**Good identification**					**25532100**
Propionibacterium propionicum		99.39	0.55	SUC 99		
Actinomyces naeslundii		0.61	0.28			
25532101	**Acceptable identification**					**25532101**
Propionibacterium propionicum		92.33	0.39	SUC 99		
Actinomyces naeslundii		7.67	0.31	URE 10		
25532102	**Very good identification**					**25532102**
Propionibacterium propionicum		98.26	0.58	SUC 99		
Actinomyces viscosus		1.34	0.12			
Actinomyces naeslundii		0.40	0.28			
25532103	**Acceptable identification**					**25532103**
Propionibacterium propionicum		91.23	0.42	SUC 99		
Actinomyces naeslundii		5.05	0.31	URE 10		
Actinomyces viscosus		3.72	0.04			
25532140	**Good identification**					**25532140**
Propionibacterium propionicum		100.0	0.55	SUC 99		
25532141	**Good identification**					**25532141**
Propionibacterium propionicum		100.0	0.39	SUC 99		
25532142	**Very good identification**					**25532142**
Propionibacterium propionicum		100.0	0.58	SUC 99		
25532143	**Good identification**					**25532143**
Propionibacterium propionicum		100.0	0.42	SUC 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
25533000	**Low discrimination**					**25533000**
					CAT	
Propionibacterium granulosum		75.97	0.39	NIT 1	90	
Actinomyces naeslundii		20.12	0.44	SUC 93	25	
Propionibacterium propionicum		3.92	0.21		1	
25533001	**Good identification**					**25533001**
Actinomyces naeslundii		100.0	0.47	SUC 99		
25533002	**Low discrimination**					**25533002**
					CAT	
Actinomyces naeslundii		42.17	0.44	SUC 99	25	
Propionibacterium granulosum		39.81	0.28		90	
Propionibacterium propionicum		12.31	0.24		1	
25533003	**Good identification**					**25533003**
Actinomyces naeslundii		100.0	0.47	SUC 99		
25533020	**Low discrimination**					**25533020**
					CAT	
Actinomyces naeslundii		44.03	0.44	SUC 99	25	
Propionibacterium granulosum		41.57	0.28		90	
Clostridium perfringens		14.40	0.13		1	
25533021	**Good identification**					**25533021**
Actinomyces naeslundii		100.0	0.47	SUC 99		
25533022	**Low discrimination**					**25533022**
					CAT	
Actinomyces naeslundii		71.95	0.44	SUC 99	25	
Propionibacterium granulosum		16.98	0.18		90	
Clostridium perfringens		11.07	0.07		1	
25533023	**Good identification**					**25533023**
Actinomyces naeslundii		100.0	0.47	SUC 99		
25533100	**Low discrimination**					**25533100**
Propionibacterium propionicum		86.84	0.55	SUC 99		
Actinomyces naeslundii		12.83	0.51			
Clostridium perfringens		0.33	0.02			
25533101	**Low discrimination**					**25533101**
Actinomyces naeslundii		66.60	0.54	SUC 99		
Propionibacterium propionicum		33.40	0.39			
25533102	**Good identification**					**25533102**
Propionibacterium propionicum		90.35	0.58	SUC 99		
Actinomyces naeslundii		8.90	0.51			
Actinomyces viscosus		0.75	0.09			
25533103	**Low discrimination**					**25533103**
Actinomyces naeslundii		56.46	0.54	SUC 99		
Propionibacterium propionicum		42.47	0.42			
Actinomyces viscosus		1.06	0.01			
25533120	**Low discrimination**					**25533120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		52.33	0.51	SUC 99		
Clostridium perfringens		44.09	0.28			
Propionibacterium propionicum		3.58	0.21			
25533121	**Good identification**					**25533121**
Actinomyces naeslundii		100.0	0.54	SUC 99		
25533122	**Low discrimination**					**25533122**
Actinomyces naeslundii		66.71	0.51	SUC 99		
Clostridium perfringens		26.45	0.22			
Propionibacterium propionicum		6.84	0.24			
25533123	**Good identification**					**25533123**
Actinomyces naeslundii		100.0	0.54	SUC 99		
25533140	**Good identification**					**25533140**
Propionibacterium propionicum		97.46	0.55	SUC 99		
Actinomyces naeslundii		2.54	0.38			
25533141	**Low discrimination**					**25533141**
Propionibacterium propionicum		73.97	0.39	SUC 99		
Actinomyces naeslundii		26.03	0.41	URE 10		
25533142	**Very good identification**					**25533142**
Propionibacterium propionicum		98.29	0.58	SUC 99		
Actinomyces naeslundii		1.71	0.38			
25533143	**Low discrimination**					**25533143**
Propionibacterium propionicum		81.00	0.42	SUC 99		
Actinomyces naeslundii		19.00	0.41	URE 10		
25533160	**Low discrimination**					**25533160**
Actinomyces naeslundii		43.54	0.38	SUC 99		
Clostridium perfringens		39.59	0.15			
Propionibacterium propionicum		16.87	0.21			
25533161	**Good identification**					**25533161**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25533162	**Low discrimination**					**25533162**
Actinomyces naeslundii		49.77	0.38	SUC 99		
Propionibacterium propionicum		28.93	0.24			
Clostridium perfringens		21.30	0.10			
25533163	**Good identification**					**25533163**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25535122	**Low discrimination**					**25535122**
					GRA	
Parabacteroides distasonis		77.06	0.41	NIT 1	1	
Clostridium perfringens		22.94	0.22		99	
25535132	**Good identification**					**25535132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		100.0	0.36	NIT 1		
25535322	**Good identification**					**25535322**
Parabacteroides distasonis		100.0	0.49	NIT 1		
25535332	**Good identification**					**25535332**
Parabacteroides distasonis		100.0	0.44	NIT 1		
25535522	**Low discrimination**					**25535522**
					GRA	
Parabacteroides distasonis		88.65	0.37	NIT 1	1	
Bifidobacterium longum subsp. longum		11.35	0.17		99	
25535722	**Good identification**					**25535722**
Parabacteroides distasonis		100.0	0.45	NIT 1		
25535732	**Good identification**					**25535732**
Parabacteroides distasonis		100.0	0.40	NIT 1		
25550002	**Good identification**					**25550002**
Actinomyces viscosus		100.0	0.52	SUC 99		
25550003	**Good identification**					**25550003**
Actinomyces viscosus		100.0	0.44	SUC 99		
25550102	**Good identification**					**25550102**
Actinomyces viscosus		100.0	0.56	SUC 99		
25550103	**Good identification**					**25550103**
Actinomyces viscosus		99.86	0.48	SUC 99		
Actinomyces naeslundii		0.14	0.23			
25551001	**Low discrimination**					**25551001**
Actinomyces naeslundii		76.78	0.39	SUC 99		
Actinomyces viscosus		23.22	0.06			
25551002	**Good identification**					**25551002**
Actinomyces viscosus		99.26	0.48	SUC 99		
Actinomyces naeslundii		0.74	0.36			
25551003	**Good identification**					**25551003**
Actinomyces viscosus		96.77	0.40	SUC 99		
Actinomyces naeslundii		3.23	0.39			
25551021	**Good identification**					**25551021**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25551023	**Low discrimination**					**25551023**
Actinomyces naeslundii		76.78	0.39	SUC 99		
Actinomyces viscosus		23.22	0.06			
25551100	**Low discrimination**					**25551100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		44.40	0.44	SUC 99	99	
Actinomyces viscosus		37.74	0.18		99	
Fusobacterium mortiferum		10.56	0.21		1	
25551101	**Low discrimination**					**25551101**
Actinomyces naeslundii		84.11	0.47	SUC 99		
Actinomyces viscosus		15.89	0.10			
25551102	**Good identification**					**25551102**
					GRA	
Actinomyces viscosus		98.21	0.52	SUC 99	99	
Actinomyces naeslundii		1.17	0.44		99	
Prevotella oralis		0.34	0.09		1	
25551103	**Acceptable identification**					**25551103**
Actinomyces viscosus		94.92	0.44	SUC 99		
Actinomyces naeslundii		5.08	0.47			
25551121	**Good identification**					**25551121**
Actinomyces naeslundii		100.0	0.47	SUC 99		
25551123	**Low discrimination**					**25551123**
Actinomyces naeslundii		84.11	0.47	SUC 99		
Actinomyces viscosus		15.89	0.10			
25553000	**Good identification**					**25553000**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25553001	**Good identification**					**25553001**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25553002	**Low discrimination**					**25553002**
Actinomyces naeslundii		58.79	0.41	SUC 99		
Actinomyces viscosus		41.21	0.14			
25553003	**Low discrimination**					**25553003**
Actinomyces naeslundii		86.52	0.44	SUC 99		
Actinomyces viscosus		13.48	0.06			
25553020	**Good identification**					**25553020**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25553021	**Good identification**					**25553021**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25553022	**Good identification**					**25553022**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25553023	**Good identification**					**25553023**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25553100	**Low discrimination**					**25553100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		89.10	0.49	SUC 99		
Propionibacterium propionicum		10.91	0.23			
25553101	**Good identification**					**25553101**
Actinomyces naeslundii		100.0	0.52	SUC 99		
25553102	**Low discrimination**					**25553102**
Actinomyces naeslundii		60.25	0.49	SUC 99		
Actinomyces viscosus		26.38	0.18			
Propionibacterium propionicum		11.06	0.26			
25553103	**Acceptable identification**					**25553103**
Actinomyces naeslundii		91.13	0.52	SUC 99		
Actinomyces viscosus		8.87	0.10			
25553120	**Good identification**					**25553120**
Actinomyces naeslundii		100.0	0.49	SUC 99		
25553121	**Good identification**					**25553121**
Actinomyces naeslundii		100.0	0.52	SUC 99		
25553122	**Good identification**					**25553122**
Actinomyces naeslundii		100.0	0.49	SUC 99		
25553123	**Good identification**					**25553123**
Actinomyces naeslundii		100.0	0.52	SUC 99		
25553140	**Low discrimination**					**25553140**
Actinomyces naeslundii		59.05	0.36	SUC 99		
Propionibacterium propionicum		40.95	0.23			
25553141	**Good identification**					**25553141**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25553143	**Good identification**					**25553143**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25553160	**Good identification**					**25553160**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25553161	**Good identification**					**25553161**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25553162	**Good identification**					**25553162**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25553163	**Good identification**					**25553163**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25570102	**Good identification**					**25570102**
Actinomyces viscosus		99.22	0.37	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.78	0.25	TRE 7		
25571000	**Good identification**					**25571000**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25571001	**Good identification**					**25571001**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25571021	**Good identification**					**25571021**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25571022	**Low discrimination**					**25571022**
Actinomyces naeslundii		51.83	0.41	SUC 99		
Clostridium perfringens		48.17	0.18			
25571023	**Good identification**					**25571023**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25571100	**Low discrimination**					**25571100**
					GRA	
Actinomyces naeslundii		81.50	0.49	SUC 99	99	
Clostridium perfringens		12.83	0.12		99	
Fusobacterium mortiferum		2.98	0.12		1	
25571101	**Good identification**					**25571101**
Actinomyces naeslundii		100.0	0.52	SUC 99		
25571103	**Low discrimination**					**25571103**
Actinomyces naeslundii		57.97	0.52	SUC 99		
Actinomyces viscosus		42.03	0.25			
25571120	**Low discrimination**					**25571120**
					SPO	
Clostridium perfringens		76.60	0.38	SUC 91	84	
Actinomyces naeslundii		15.05	0.49	CEL 4	1	
Clostridium septicum		8.35	0.15		99	
25571121	**Good identification**					**25571121**
Actinomyces naeslundii		96.69	0.52	SUC 99		
Clostridium perfringens		3.31	0.04			
25571123	**Good identification**					**25571123**
Actinomyces naeslundii		100.0	0.52	SUC 99		
25571140	**Low discrimination**					**25571140**
Actinomyces naeslundii		85.48	0.36	SUC 99		
Clostridium perfringens		14.52	0.00			
25571141	**Good identification**					**25571141**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25571142	**Low discrimination**					**25571142**
Actinomyces naeslundii		84.26	0.36	SUC 99		
Actinomyces viscosus		15.74	-.01			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25571143	**Good identification**					**25571143**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25571161	**Good identification**					**25571161**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25571163	**Good identification**					**25571163**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25573000	**Good identification**					**25573000**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25573001	**Good identification**					**25573001**
Actinomyces naeslundii		100.0	0.49	SUC 99		
25573002	**Acceptable identification**					**25573002**
Actinomyces naeslundii		94.98	0.46	SUC 99		
Actinomyces israelii		5.03	0.15			
25573003	**Good identification**					**25573003**
Actinomyces naeslundii		100.0	0.49	SUC 99		
25573020	**Good identification**					**25573020**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25573021	**Good identification**					**25573021**
Actinomyces naeslundii		100.0	0.49	SUC 99		
25573022	**Good identification**					**25573022**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25573023	**Good identification**					**25573023**
Actinomyces naeslundii		100.0	0.49	SUC 99		
25573041	**Good identification**					**25573041**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25573043	**Good identification**					**25573043**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25573061	**Good identification**					**25573061**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25573063	**Good identification**					**25573063**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25573100	**Acceptable identification**					**25573100**
Actinomyces naeslundii		90.81	0.53	SUC 99		
Propionibacterium propionicum		4.69	0.21			
Actinomyces israelii		4.50	0.22			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
25573101	**Very good identification**					**25573101**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25573102	**Low discrimination**					**25573102**
Actinomyces naeslundii		85.56	0.53	SUC 99		
Propionibacterium propionicum		6.62	0.24			
Actinomyces israelii		6.36	0.25			
25573103	**Very good identification**					**25573103**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25573120	**Good identification**					**25573120**
Actinomyces naeslundii		97.42	0.53	SUC 99		
Clostridium perfringens		2.58	0.04			
25573121	**Very good identification**					**25573121**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25573122	**Good identification**					**25573122**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25573123	**Very good identification**					**25573123**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25573140	**Low discrimination**					**25573140**
Actinomyces naeslundii		71.35	0.41	SUC 99		
Propionibacterium propionicum		20.86	0.21			
Actinomyces israelii		7.79	0.15			
25573141	**Good identification**					**25573141**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25573142	**Low discrimination**					**25573142**
Actinomyces naeslundii		62.41	0.41	SUC 99		
Propionibacterium propionicum		27.37	0.24			
Actinomyces israelii		10.23	0.18			
25573143	**Good identification**					**25573143**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25573160	**Good identification**					**25573160**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25573161	**Good identification**					**25573161**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25573162	**Good identification**					**25573162**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25573163	**Good identification**					**25573163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	SUC 99		
25575122	**Good identification**					**25575122**
Parabacteroides distasonis		100.0	0.36	NIT 1		
25575322	**Good identification**					**25575322**
Parabacteroides distasonis		100.0	0.44	NIT 1		
25575332	**Good identification**					**25575332**
Parabacteroides distasonis		100.0	0.39	NIT 1		
25575722	**Good identification**					**25575722**
Parabacteroides distasonis		100.0	0.40	NIT 1		
25575732	**Good identification**					**25575732**
Parabacteroides distasonis		95.23	0.36	NIT 1		
Clostridium clostridioforme		4.77	0.08			
25577100	**Good identification**					**25577100**
Actinomyces israelii		100.0	0.40	SUC 98		
25577102	**Good identification**					**25577102**
Actinomyces israelii		100.0	0.43	SUC 98		
25577142	**Good identification**					**25577142**
Actinomyces israelii		100.0	0.36	SUC 98		
25577500	**Good identification**					**25577500**
Actinomyces israelii		100.0	0.39	SUC 98		
25577502	**Good identification**					**25577502**
Actinomyces israelii		100.0	0.42	SUC 98		
25610000	**Good identification**					**25610000**
Fusobacterium mortiferum		100.0	0.41	NIT 1		
25610100	**Good identification**					**25610100**
Fusobacterium mortiferum		100.0	0.47	NIT 1		
25611000	**Good identification**					**25611000**
					GRA	
Fusobacterium mortiferum		98.77	0.44	NIT 1	1	
Actinomyces naeslundii		1.23	0.23		99	
25611100	**Good identification**					**25611100**
					GRA	
Fusobacterium mortiferum		98.50	0.50	NIT 1	1	
Actinomyces naeslundii		1.50	0.30		99	
25613100	**Low discrimination**					**25613100**
					GRA	
Actinomyces naeslundii		74.54	0.35	SUC 99	99	
Fusobacterium mortiferum		25.46	0.16	MLT 94	1	
25613101	**Good identification**					**25613101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	SUC 99		
25613102	**Good identification**					**25613102**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25613103	**Good identification**					**25613103**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25613120	**Good identification**					**25613120**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25613121	**Good identification**					**25613121**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25613122	**Good identification**					**25613122**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25613123	**Good identification**					**25613123**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25630100	**Good identification**					**25630100**
Fusobacterium mortiferum		100.0	0.38	NIT 1		
25631100	**Acceptable identification**					**25631100**
					GRA	
Fusobacterium mortiferum		90.99	0.41	NIT 1	1	
Actinomyces naeslundii		9.01	0.35		99	
25631101	**Good identification**					**25631101**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25631103	**Good identification**					**25631103**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25631121	**Good identification**					**25631121**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25631122	**Low discrimination**					**25631122**
					GRA	
Actinomyces naeslundii		44.68	0.35	SUC 99	99	
Clostridium perfringens		37.99	0.12	MLT 94	99	
Parabacteroides distasonis		17.33	0.15		1	
25631123	**Good identification**					**25631123**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25633001	**Good identification**					**25633001**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25633003	**Good identification**					**25633003**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25633021	**Good identification**					**25633021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	SUC 99		
25633023	**Good identification**					**25633023**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25633100	**Good identification**					**25633100**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25633101	**Good identification**					**25633101**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25633102	**Good identification**					**25633102**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25633103	**Good identification**					**25633103**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25633120	**Good identification**					**25633120**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25633121	**Good identification**					**25633121**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25633122	**Good identification**					**25633122**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25633123	**Good identification**					**25633123**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25635322	**Good identification**					**25635322**
Parabacteroides distasonis		100.0	0.40	NIT 1		
25635332	**Good identification**					**25635332**
Parabacteroides distasonis		100.0	0.35	NIT 1		
25635722	**Good identification**					**25635722**
Parabacteroides distasonis		100.0	0.36	NIT 1		
25650100	**Good identification**					**25650100**
Fusobacterium mortiferum		100.0	0.39	NIT 1		
25651000	**Good identification**					**25651000**
					GRA	
Fusobacterium mortiferum		95.27	0.36	NIT 1	1	
Actinomyces naeslundii		4.73	0.25		99	
25651100	**Acceptable identification**					**25651100**
					GRA	
Fusobacterium mortiferum		94.29	0.42	NIT 1	1	
Actinomyces naeslundii		5.71	0.33		99	
25651101	**Good identification**					**25651101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	SUC 99		
25651103	**Low discrimination**					**25651103**
Actinomyces naeslundii		81.84	0.36	SUC 99		
Actinomyces viscosus		18.16	0.00	MLT 94		
25651121	**Good identification**					**25651121**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25651123	**Good identification**					**25651123**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25653100	**Good identification**					**25653100**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25653101	**Good identification**					**25653101**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25653102	**Good identification**					**25653102**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25653103	**Good identification**					**25653103**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25653120	**Good identification**					**25653120**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25653121	**Good identification**					**25653121**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25653122	**Good identification**					**25653122**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25653123	**Good identification**					**25653123**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25671101	**Good identification**					**25671101**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25671102	**Low discrimination**					**25671102**
					GRA	
Actinomyces naeslundii		69.82	0.37	SUC 99	99	
Fusobacterium mortiferum		19.70	0.16	MLT 94	1	
Lactobacillus acidophilus		10.48	-.01		99	
25671103	**Good identification**					**25671103**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25671120	**Low discrimination**					**25671120**
					SPO	
Actinomyces naeslundii		61.50	0.37	SUC 99	1	
Clostridium septicum		38.50	0.05	MLT 94	99	
25671121	**Good identification**					**25671121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	SUC 99		
25671122	**Low discrimination**					**25671122**
					GRA	
Actinomyces naeslundii		76.47	0.37	SUC 99	99	
Parabacteroides distasonis		12.05	0.11	MLT 94	1	
Lactobacillus acidophilus		11.48	-.01		99	
25671123	**Good identification**					**25671123**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25673001	**Good identification**					**25673001**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25673003	**Good identification**					**25673003**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25673021	**Good identification**					**25673021**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25673023	**Good identification**					**25673023**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25673100	**Good identification**					**25673100**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25673101	**Good identification**					**25673101**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25673102	**Good identification**					**25673102**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25673103	**Good identification**					**25673103**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25673120	**Good identification**					**25673120**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25673121	**Good identification**					**25673121**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25673122	**Good identification**					**25673122**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25673123	**Good identification**					**25673123**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25675322	**Good identification**					**25675322**
Parabacteroides distasonis		100.0	0.35	NIT 1		
25703101	**Good identification**					**25703101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	SUC 99		
25703103	**Good identification**					**25703103**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25703121	**Good identification**					**25703121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25703123	**Good identification**					**25703123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25710002	**Good identification**					**25710002**
Actinomyces viscosus		99.75	0.52	SUC 99		
Actinomyces odontolyticus		0.25	0.14			
25710003	**Good identification**					**25710003**
Actinomyces viscosus		99.77	0.44	SUC 99		
Actinomyces naeslundii		0.23	0.23			
25710102	**Good identification**					**25710102**
					GRA	
Actinomyces viscosus		99.80	0.56	SUC 99	99	
Actinomyces naeslundii		0.08	0.27		99	
Fusobacterium mortiferum		0.08	0.15		1	
25710103	**Good identification**					**25710103**
Actinomyces viscosus		99.64	0.48	SUC 99		
Actinomyces naeslundii		0.36	0.30			
25711001	**Low discrimination**					**25711001**
Actinomyces naeslundii		89.84	0.46	SUC 99		
Actinomyces viscosus		10.16	0.06			
25711002	**Good identification**					**25711002**
Actinomyces viscosus		97.53	0.48	SUC 99		
Actinomyces naeslundii		1.94	0.43			
Actinomyces odontolyticus		0.27	0.11			
25711003	**Acceptable identification**					**25711003**
Actinomyces viscosus		91.80	0.40	SUC 99		
Actinomyces naeslundii		8.20	0.46			
25711021	**Good identification**					**25711021**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25711023	**Low discrimination**					**25711023**
Actinomyces naeslundii		89.84	0.46	SUC 99		
Actinomyces viscosus		10.16	0.06			
25711100	**Low discrimination**					**25711100**
					GRA	
Actinomyces naeslundii		44.85	0.51	SUC 99	99	
Fusobacterium mortiferum		23.22	0.34		1	
Clostridium perfringens		16.55	0.21		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25711101	**Acceptable identification**					**25711101**
Actinomyces naeslundii		93.40	0.54	SUC 99		
Actinomyces viscosus		6.60	0.10			
25711102	**Good identification**					**25711102**
Actinomyces viscosus		96.01	0.52	SUC 99		
Actinomyces naeslundii		3.05	0.51			
Clostridium perfringens		0.53	0.16			
25711103	**Low discrimination**					**25711103**
Actinomyces viscosus		87.49	0.44	SUC 99		
Actinomyces naeslundii		12.51	0.54			
25711120	**Acceptable identification**					**25711120**
					SPO	
Clostridium perfringens		90.43	0.47	SUC 91	84	
Actinomyces naeslundii		7.58	0.51		1	
Clostridium septicum		1.29	0.09		99	
25711121	**Acceptable identification**					**25711121**
Actinomyces naeslundii		92.56	0.54	SUC 99		
Clostridium perfringens		7.44	0.13			
25711122	**Low discrimination**					**25711122**
					GRA	
Clostridium perfringens		69.67	0.41	SUC 91	99	
Prevotella oralis		12.75	0.25		1	
Actinomyces naeslundii		12.41	0.51		99	
25711123	**Acceptable identification**					**25711123**
Actinomyces naeslundii		90.22	0.54	SUC 99		
Actinomyces viscosus		6.37	0.10			
Clostridium perfringens		3.41	0.07			
25711140	**Low discrimination**					**25711140**
Actinomyces naeslundii		71.51	0.38	SUC 99		
Clostridium perfringens		28.49	0.09			
25711141	**Good identification**					**25711141**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25711143	**Low discrimination**					**25711143**
Actinomyces naeslundii		71.42	0.41	SUC 99		
Actinomyces viscosus		28.58	0.10			
25711161	**Acceptable identification**					**25711161**
Actinomyces naeslundii		92.02	0.41	SUC 99		
Clostridium perfringens		7.98	0.01			
25711163	**Good identification**					**25711163**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25712101	**Good identification**					**25712101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	SUC 99		
25712121	**Good identification**					**25712121**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25712123	**Good identification**					**25712123**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25713000	**Good identification**					**25713000**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25713001	**Good identification**					**25713001**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25713002	**Low discrimination**					**25713002**
Actinomyces naeslundii		79.23	0.48	SUC 99		
Actinomyces viscosus		20.77	0.14			
25713003	**Acceptable identification**					**25713003**
Actinomyces naeslundii		94.50	0.51	SUC 99		
Actinomyces viscosus		5.51	0.06			
25713020	**Good identification**					**25713020**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25713021	**Good identification**					**25713021**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25713022	**Good identification**					**25713022**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25713023	**Good identification**					**25713023**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25713040	**Good identification**					**25713040**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25713041	**Good identification**					**25713041**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25713042	**Good identification**					**25713042**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25713043	**Good identification**					**25713043**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25713060	**Good identification**					**25713060**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25713061	**Good identification**					**25713061**
Actinomyces naeslundii		100.0	0.38	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25713062	**Good identification**					**25713062**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25713063	**Good identification**					**25713063**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25713100	**Good identification**					**25713100**
Actinomyces naeslundii		95.62	0.56	SUC 99		
Propionibacterium propionicum		4.38	0.23			
25713101	**Very good identification**					**25713101**
Actinomyces naeslundii		100.0	0.59	SUC 99		
25713102	**Low discrimination**					**25713102**
Actinomyces naeslundii		80.34	0.56	SUC 99		
Actinomyces viscosus		13.15	0.18			
Propionibacterium propionicum		5.51	0.26			
25713103	**Very good identification**					**25713103**
Actinomyces naeslundii		96.49	0.59	SUC 99		
Actinomyces viscosus		3.51	0.10			
25713111	**Good identification**					**25713111**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25713113	**Good identification**					**25713113**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25713120	**Acceptable identification**					**25713120**
Actinomyces naeslundii		94.15	0.56	SUC 99		
Clostridium perfringens		5.85	0.13			
25713121	**Very good identification**					**25713121**
Actinomyces naeslundii		100.0	0.59	SUC 99		
25713122	**Good identification**					**25713122**
Actinomyces naeslundii		97.16	0.56	SUC 99		
Clostridium perfringens		2.84	0.07			
25713123	**Very good identification**					**25713123**
Actinomyces naeslundii		100.0	0.59	SUC 99		
25713131	**Good identification**					**25713131**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25713133	**Good identification**					**25713133**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25713140	**Low discrimination**					**25713140**
Actinomyces naeslundii		79.41	0.43	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		20.59	0.23			
25713141	**Good identification**					**25713141**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25713142	**Low discrimination**					**25713142**
Actinomyces naeslundii		72.00	0.43	SUC 99		
Propionibacterium propionicum		28.00	0.26			
25713143	**Good identification**					**25713143**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25713160	**Acceptable identification**					**25713160**
Actinomyces naeslundii		93.72	0.43	SUC 99		
Clostridium perfringens		6.28	0.01			
25713161	**Good identification**					**25713161**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25713162	**Good identification**					**25713162**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25713163	**Good identification**					**25713163**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25715322	**Acceptable identification**					**25715322**
Parabacteroides distasonis		94.89	0.38	NIT 1		
Prevotella oralis		5.11	0.04			
25721101	**Good identification**					**25721101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25721103	**Low discrimination**					**25721103**
Actinomyces naeslundii		78.67	0.37	SUC 99		
Actinomyces viscosus		21.33	0.03	FRU 95		
25721121	**Good identification**					**25721121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25721123	**Good identification**					**25721123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25723100	**Good identification**					**25723100**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25723101	**Good identification**					**25723101**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25723102	**Good identification**					**25723102**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25723103	**Good identification**					**25723103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	SUC 99		
25723120	**Good identification**					**25723120**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25723121	**Good identification**					**25723121**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25723122	**Good identification**					**25723122**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25723123	**Good identification**					**25723123**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25730101	**Good identification**					**25730101**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25730102	**Good identification**					**25730102**
Actinomyces viscosus		97.95	0.37	SUC 99		
Actinomyces naeslundii		2.05	0.32	TRE 7		
25730121	**Good identification**					**25730121**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25730123	**Good identification**					**25730123**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25731000	**Low discrimination**					**25731000**
					GRA	
Actinomyces naeslundii		81.02	0.48	SUC 99	99	
Clostridium perfringens		11.10	0.11		99	
Fusobacterium mortiferum		7.87	0.19		1	
25731001	**Good identification**					**25731001**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25731003	**Low discrimination**					**25731003**
Actinomyces naeslundii		69.73	0.51	SUC 99		
Actinomyces viscosus		30.27	0.21			
25731020	**Low discrimination**					**25731020**
Clostridium perfringens		81.59	0.37	SUC 91		
Actinomyces naeslundii		18.41	0.48	LAC 88		
25731021	**Good identification**					**25731021**
Actinomyces naeslundii		97.10	0.51	SUC 99		
Clostridium perfringens		2.90	0.03			
25731023	**Good identification**					**25731023**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25731040	**Good identification**					**25731040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	SUC 99		
25731041	**Good identification**					**25731041**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25731042	**Good identification**					**25731042**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25731043	**Good identification**					**25731043**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25731061	**Good identification**					**25731061**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25731063	**Good identification**					**25731063**
Actinomyces naeslundii		100.0	0.38	SUC 99		
25731100	**Low discrimination**					**25731100**
					GRA	
Actinomyces naeslundii		69.20	0.56	SUC 99	99	
Clostridium perfringens		24.44	0.26		99	
Fusobacterium mortiferum		5.51	0.25		1	
25731101	**Very good identification**					**25731101**
Actinomyces naeslundii		100.0	0.59	SUC 99		
25731103	**Low discrimination**					**25731103**
Actinomyces naeslundii		78.67	0.59	SUC 99		
Actinomyces viscosus		21.33	0.25			
25731111	**Good identification**					**25731111**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25731113	**Good identification**					**25731113**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25731120	**Acceptable identification**					**25731120**
					SPO	
Clostridium perfringens		90.05	0.52	SUC 91	84	
Actinomyces naeslundii		7.89	0.56		1	
Clostridium septicum		2.07	0.17		99	
25731121	**Good identification**					**25731121**
Actinomyces naeslundii		92.86	0.59	SUC 99		
Clostridium perfringens		7.14	0.17			
25731122	**Low discrimination**					**25731122**
					GRA	
Clostridium perfringens		82.57	0.46	SUC 91	99	
Actinomyces naeslundii		15.37	0.56		99	
Parabacteroides distasonis		1.35	0.25		1	
25731123	**Very good identification**					**25731123**
Actinomyces naeslundii		96.51	0.59	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		3.49	0.12			
25731131	**Good identification**					**25731131**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25731133	**Good identification**					**25731133**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25731140	**Low discrimination**					**25731140**
Actinomyces naeslundii		72.40	0.43	SUC 99		
Clostridium perfringens		27.60	0.13			
25731141	**Good identification**					**25731141**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25731142	**Low discrimination**					**25731142**
Actinomyces naeslundii		80.05	0.43	SUC 99		
Clostridium perfringens		14.36	0.08			
Actinomyces viscosus		5.59	-.01			
25731143	**Good identification**					**25731143**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25731160	**Acceptable identification**					**25731160**
Clostridium perfringens		92.49	0.39	SUC 91		
Actinomyces naeslundii		7.51	0.43			
25731161	**Acceptable identification**					**25731161**
Actinomyces naeslundii		92.34	0.46	SUC 99		
Clostridium perfringens		7.66	0.05			
25731163	**Good identification**					**25731163**
Actinomyces naeslundii		100.0	0.46	SUC 99		
25732100	**Low discrimination**					**25732100**
Actinomyces naeslundii		68.36	0.37	SUC 99		
Propionibacterium propionicum		31.64	0.21	GAL 96		
25732101	**Good identification**					**25732101**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25732102	**Low discrimination**					**25732102**
Actinomyces naeslundii		51.47	0.37	SUC 99		
Propionibacterium propionicum		35.74	0.24	GAL 96		
Actinomyces viscosus		12.79	0.03			
25732103	**Good identification**					**25732103**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25732120	**Good identification**					**25732120**
Actinomyces naeslundii		100.0	0.37	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25732121	**Good identification**					**25732121**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25732122	**Good identification**					**25732122**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25732123	**Good identification**					**25732123**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25733000	**Good identification**					**25733000**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25733001	**Good identification**					**25733001**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25733002	**Good identification**					**25733002**
Actinomyces naeslundii		98.32	0.53	SUC 99		
Actinomyces israelii		1.68	0.14			
25733003	**Good identification**					**25733003**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25733020	**Good identification**					**25733020**
Actinomyces naeslundii		97.75	0.53	SUC 99		
Clostridium perfringens		2.25	0.03			
25733021	**Good identification**					**25733021**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25733022	**Good identification**					**25733022**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25733023	**Good identification**					**25733023**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25733040	**Good identification**					**25733040**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25733041	**Good identification**					**25733041**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25733042	**Good identification**					**25733042**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25733043	**Good identification**					**25733043**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25733060	**Good identification**					**25733060**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25733061	**Good identification**					**25733061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	SUC 99		
25733062	**Good identification**					**25733062**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25733063	**Good identification**					**25733063**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25733100	**Very good identification**					**25733100**
Actinomyces naeslundii		96.59	0.61	SUC 99		
Propionibacterium propionicum		1.86	0.21			
Actinomyces israelii		1.55	0.21			
25733101	**Very good identification**					**25733101**
Actinomyces naeslundii		100.0	0.64	SUC 99		
25733102	**Good identification**					**25733102**
Actinomyces naeslundii		94.40	0.61	SUC 99		
Propionibacterium propionicum		2.73	0.24			
Actinomyces israelii		2.27	0.24			
25733103	**Very good identification**					**25733103**
Actinomyces naeslundii		100.0	0.64	SUC 99		
25733110	**Good identification**					**25733110**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25733111	**Good identification**					**25733111**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25733112	**Good identification**					**25733112**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25733113	**Good identification**					**25733113**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25733120	**Good identification**					**25733120**
Actinomyces naeslundii		94.39	0.61	SUC 99		
Clostridium perfringens		5.61	0.17			
25733121	**Very good identification**					**25733121**
Actinomyces naeslundii		100.0	0.64	SUC 99		
25733122	**Very good identification**					**25733122**
Actinomyces naeslundii		97.28	0.61	SUC 99		
Clostridium perfringens		2.72	0.12			
25733123	**Very good identification**					**25733123**
Actinomyces naeslundii		100.0	0.64	SUC 99		
25733130	**Good identification**					**25733130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	SUC 99		
25733131	**Good identification**					**25733131**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25733132	**Good identification**					**25733132**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25733133	**Good identification**					**25733133**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25733140	**Low discrimination**					**25733140**
Actinomyces naeslundii		87.36	0.48	SUC 99		
Propionibacterium propionicum		9.55	0.21			
Actinomyces israelii		3.09	0.14			
25733141	**Good identification**					**25733141**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25733142	**Low discrimination**					**25733142**
Actinomyces naeslundii		82.17	0.48	SUC 99		
Propionibacterium propionicum		13.47	0.24			
Actinomyces israelii		4.36	0.17			
25733143	**Good identification**					**25733143**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25733160	**Acceptable identification**					**25733160**
Actinomyces naeslundii		93.97	0.48	SUC 99		
Clostridium perfringens		6.03	0.05			
25733161	**Good identification**					**25733161**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25733162	**Good identification**					**25733162**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25733163	**Good identification**					**25733163**
Actinomyces naeslundii		100.0	0.51	SUC 99		
25735122	**Acceptable identification**					**25735122**
					GRA	
Parabacteroides distasonis		93.57	0.41	NIT 1	1	
Clostridium perfringens		6.43	0.12		99	
25735132	**Good identification**					**25735132**
Parabacteroides distasonis		100.0	0.37	NIT 1		
25735322	**Good identification**					**25735322**
Parabacteroides distasonis		100.0	0.49	NIT 1		
25735332	**Good identification**					**25735332**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		100.0	0.45	NIT 1		
25735522	**Good identification**					**25735522**
Parabacteroides distasonis		100.0	0.37	NIT 1		
25735722	**Good identification**					**25735722**
Parabacteroides distasonis		100.0	0.46	NIT 1		
25735732	**Acceptable identification**					**25735732**
Parabacteroides distasonis		94.83	0.41	NIT 1		
Clostridium clostridioforme		5.17	0.14			
25737100	**Acceptable identification**					**25737100**
Actinomyces israelii		94.81	0.39	SUC 98		
Actinomyces naeslundii		5.19	0.27	CEL 93		
25737102	**Good identification**					**25737102**
Actinomyces israelii		96.48	0.42	SUC 98		
Actinomyces naeslundii		3.52	0.27	CEL 93		
25737142	**Good identification**					**25737142**
Actinomyces israelii		100.0	0.35	SUC 98		
25737500	**Good identification**					**25737500**
Actinomyces israelii		100.0	0.38	SUC 98		
25737502	**Good identification**					**25737502**
Actinomyces israelii		96.39	0.41	SUC 98		
Bifidobacterium adolescentis		3.61	0.10	CEL 93		
25743100	**Good identification**					**25743100**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25743101	**Good identification**					**25743101**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25743102	**Good identification**					**25743102**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25743103	**Good identification**					**25743103**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25743120	**Good identification**					**25743120**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25743121	**Good identification**					**25743121**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25743122	**Good identification**					**25743122**
Actinomyces naeslundii		100.0	0.36	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25743123	**Good identification**					**25743123**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25750002	**Good identification**					**25750002**
Actinomyces viscosus		100.0	0.42	SUC 99		
25750102	**Good identification**					**25750102**
Actinomyces viscosus		99.60	0.46	SUC 99		
Actinomyces naeslundii		0.40	0.29			
25750103	**Good identification**					**25750103**
Actinomyces viscosus		98.21	0.38	SUC 99		
Actinomyces naeslundii		1.79	0.32			
25751000	**Low discrimination**					**25751000**
					GRA	
Actinomyces naeslundii		72.99	0.45	SUC 99	99	
Fusobacterium mortiferum		11.59	0.20		1	
Lactobacillus jensenii		7.97	0.07		99	
25751001	**Good identification**					**25751001**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25751002	**Acceptable identification**					**25751002**
Actinomyces viscosus		90.40	0.38	SUC 99		
Actinomyces naeslundii		8.95	0.45			
Lactobacillus jensenii		0.65	0.04			
25751020	**Low discrimination**					**25751020**
Actinomyces naeslundii		87.29	0.45	SUC 99		
Clostridium perfringens		12.71	0.09			
25751021	**Good identification**					**25751021**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25751022	**Low discrimination**					**25751022**
					GRA	
Actinomyces naeslundii		59.88	0.45	SUC 99	99	
Prevotella oralis		29.90	0.14		1	
Actinomyces viscosus		6.11	0.04		99	
25751023	**Good identification**					**25751023**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25751041	**Good identification**					**25751041**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25751043	**Good identification**					**25751043**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25751061	**Good identification**					**25751061**
Actinomyces naeslundii		100.0	0.35	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25751063	**Good identification**					**25751063**
Actinomyces naeslundii		100.0	0.35	SUC 99		
25751100	**Low discrimination**					**25751100**
					GRA	
Actinomyces naeslundii		81.81	0.53	SUC 99	99	
Fusobacterium mortiferum		10.65	0.26		1	
Actinomyces viscosus		5.21	0.08		99	
25751101	**Good identification**					**25751101**
Actinomyces naeslundii		98.60	0.56	SUC 99		
Actinomyces viscosus		1.40	0.00			
25751102	**Low discrimination**					**25751102**
					GRA	
Actinomyces viscosus		83.20	0.43	SUC 99	99	
Actinomyces naeslundii		13.19	0.53		99	
Prevotella oralis		1.91	0.13		1	
25751120	**Low discrimination**					**25751120**
					SPO	
Actinomyces naeslundii		51.22	0.53	SUC 99	1	
Clostridium septicum		20.75	0.17		99	
Clostridium perfringens		19.21	0.23		84	
25751121	**Good identification**					**25751121**
Actinomyces naeslundii		100.0	0.56	SUC 99		
25751123	**Good identification**					**25751123**
					GRA	
Actinomyces naeslundii		96.84	0.56	SUC 99	99	
Prevotella oralis		1.79	0.01		1	
Actinomyces viscosus		1.37	0.00		99	
25751140	**Good identification**					**25751140**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25751141	**Good identification**					**25751141**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25751142	**Low discrimination**					**25751142**
Actinomyces naeslundii		68.73	0.40	SUC 99		
Actinomyces viscosus		24.82	0.08			
Actinomyces israelii		6.46	0.13			
25751143	**Acceptable identification**					**25751143**
Actinomyces naeslundii		92.57	0.43	SUC 99		
Actinomyces viscosus		7.43	0.00			
25751160	**Low discrimination**					**25751160**
Actinomyces naeslundii		71.18	0.40	SUC 99		
Clostridium perfringens		28.82	0.11			
25751161	**Good identification**					**25751161**
Actinomyces naeslundii		100.0	0.43	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25751162	**Low discrimination**					**25751162**
					GRA	
Actinomyces naeslundii		74.19	0.40	SUC 99	99	
Clostridium perfringens		14.14	0.05		99	
Prevotella oralis		11.67	0.01		1	
25751163	**Good identification**					**25751163**
Actinomyces naeslundii		100.0	0.43	SUC 99		
25752101	**Good identification**					**25752101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25752103	**Low discrimination**					**25752103**
Actinomyces naeslundii		77.78	0.37	SUC 99		
Actinomyces viscosus		22.22	0.04	GAL 96		
25752121	**Good identification**					**25752121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25752123	**Good identification**					**25752123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25753000	**Acceptable identification**					**25753000**
Actinomyces naeslundii		94.43	0.50	SUC 99		
Actinomyces israelii		5.57	0.20			
25753001	**Good identification**					**25753001**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25753002	**Low discrimination**					**25753002**
Actinomyces naeslundii		87.63	0.50	SUC 99		
Actinomyces israelii		7.76	0.23			
Actinomyces viscosus		4.61	0.04			
25753003	**Good identification**					**25753003**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25753020	**Good identification**					**25753020**
Actinomyces naeslundii		100.0	0.50	SUC 99		
25753021	**Good identification**					**25753021**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25753022	**Good identification**					**25753022**
Actinomyces naeslundii		100.0	0.50	SUC 99		
25753023	**Good identification**					**25753023**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25753040	**Good identification**					**25753040**
Actinomyces naeslundii		100.0	0.37	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25753041	**Good identification**					**25753041**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25753042	**Low discrimination**					**25753042**
Actinomyces naeslundii		83.67	0.37	SUC 99		
Actinomyces israelii		16.33	0.16			
25753043	**Good identification**					**25753043**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25753060	**Good identification**					**25753060**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25753061	**Good identification**					**25753061**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25753062	**Good identification**					**25753062**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25753063	**Good identification**					**25753063**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25753100	**Good identification**					**25753100**
Actinomyces naeslundii		92.34	0.58	SUC 99		
Actinomyces israelii		7.66	0.30			
25753101	**Very good identification**					**25753101**
Actinomyces naeslundii		100.0	0.61	SUC 99		
25753102	**Low discrimination**					**25753102**
Actinomyces naeslundii		86.41	0.58	SUC 99		
Actinomyces israelii		10.75	0.33			
Actinomyces viscosus		2.84	0.08			
25753103	**Very good identification**					**25753103**
Actinomyces naeslundii		99.28	0.61	SUC 99		
Actinomyces viscosus		0.72	0.00			
25753110	**Low discrimination**					**25753110**
Actinomyces naeslundii		80.94	0.36	SUC 99		
Actinomyces israelii		19.06	0.16	MLZ 5		
25753111	**Good identification**					**25753111**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25753112	**Low discrimination**					**25753112**
Actinomyces naeslundii		73.89	0.36	SUC 99		
Actinomyces israelii		26.11	0.19	MLZ 5		
25753113	**Good identification**					**25753113**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	SUC 99			
25753120	**Very good identification**						**25753120**
Actinomyces naeslundii		100.0	0.58	SUC 99			
25753121	**Very good identification**						**25753121**
Actinomyces naeslundii		100.0	0.61	SUC 99			
25753122	**Very good identification**						**25753122**
					SPO	GRA	
Actinomyces naeslundii		96.42	0.58	SUC 99	1	99	
Clostridium ramosum		2.21	0.05		92	50	
Prevotella oralis		1.38	0.01		1	1	
25753123	**Very good identification**						**25753123**
Actinomyces naeslundii		100.0	0.61	SUC 99			
25753130	**Good identification**						**25753130**
Actinomyces naeslundii		100.0	0.36	SUC 99			
25753131	**Good identification**						**25753131**
Actinomyces naeslundii		100.0	0.39	SUC 99			
25753132	**Good identification**						**25753132**
Actinomyces naeslundii		100.0	0.36	SUC 99			
25753133	**Good identification**						**25753133**
Actinomyces naeslundii		100.0	0.39	SUC 99			
25753140	**Low discrimination**						**25753140**
Actinomyces naeslundii		84.54	0.45	SUC 99			
Actinomyces israelii		15.46	0.23				
25753141	**Good identification**						**25753141**
Actinomyces naeslundii		100.0	0.48	SUC 99			
25753142	**Low discrimination**						**25753142**
Actinomyces naeslundii		78.48	0.45	SUC 99			
Actinomyces israelii		21.52	0.26				
25753143	**Good identification**						**25753143**
Actinomyces naeslundii		100.0	0.48	SUC 99			
25753160	**Good identification**						**25753160**
Actinomyces naeslundii		100.0	0.45	SUC 99			
25753161	**Good identification**						**25753161**
Actinomyces naeslundii		100.0	0.48	SUC 99			
25753162	**Good identification**						**25753162**
Actinomyces naeslundii		100.0	0.45	SUC 99			
25753163	**Good identification**						**25753163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	SUC 99		
25755100	**Good identification**					**25755100**
Actinomyces israelii		100.0	0.35	SUC 98		
25755102	**Low discrimination**					**25755102**
					GRA	
Actinomyces israelii		85.73	0.39	SUC 98	99	
Actinomyces viscosus		11.14	0.08		99	
Prevotella oralis		3.13	-.03		1	
25755502	**Low discrimination**					**25755502**
Actinomyces israelii		81.04	0.37	SUC 98		
Bifidobacterium adolescentis		18.96	0.20			
25757000	**Good identification**					**25757000**
Actinomyces israelii		100.0	0.38	SUC 98		
25757002	**Good identification**					**25757002**
Actinomyces israelii		100.0	0.41	SUC 98		
25757100	**Good identification**					**25757100**
Actinomyces israelii		98.95	0.48	SUC 98		
Actinomyces naeslundii		1.05	0.24			
25757102	**Good identification**					**25757102**
Actinomyces israelii		99.30	0.51	SUC 98		
Actinomyces naeslundii		0.70	0.24			
25757112	**Good identification**					**25757112**
Actinomyces israelii		100.0	0.37	SUC 98		
25757140	**Good identification**					**25757140**
Actinomyces israelii		100.0	0.41	SUC 98		
25757142	**Good identification**					**25757142**
Actinomyces israelii		100.0	0.44	SUC 98		
25757400	**Good identification**					**25757400**
Actinomyces israelii		100.0	0.37	SUC 98		
25757402	**Good identification**					**25757402**
Actinomyces israelii		100.0	0.40	SUC 98		
25757500	**Good identification**					**25757500**
Actinomyces israelii		100.0	0.47	SUC 98		
25757502	**Good identification**					**25757502**
Actinomyces israelii		96.73	0.50	SUC 98		
Bifidobacterium adolescentis		3.27	0.18			
25757512	**Low discrimination**					**25757512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		78.35	0.36	SUC 98		
Bifidobacterium adolescentis		21.65	0.20	MLZ 13		
25757540	**Good identification**					**25757540**
Actinomyces israelii		100.0	0.40	SUC 98		
25757542	**Good identification**					**25757542**
Actinomyces israelii		100.0	0.43	SUC 98		
25761100	**Good identification**					**25761100**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25761101	**Good identification**					**25761101**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25761102	**Low discrimination**					**25761102**
Actinomyces naeslundii		80.35	0.36	SUC 99		
Actinomyces viscosus		19.65	0.01	FRU 95		
25761103	**Good identification**					**25761103**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25761120	**Good identification**					**25761120**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25761121	**Good identification**					**25761121**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25761122	**Good identification**					**25761122**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25761123	**Good identification**					**25761123**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25763001	**Good identification**					**25763001**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25763003	**Good identification**					**25763003**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25763021	**Good identification**					**25763021**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25763023	**Good identification**					**25763023**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25763100	**Good identification**					**25763100**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25763101	**Good identification**					**25763101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	SUC 99		
25763102	**Good identification**					**25763102**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25763103	**Good identification**					**25763103**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25763120	**Good identification**					**25763120**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25763121	**Good identification**					**25763121**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25763122	**Good identification**					**25763122**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25763123	**Good identification**					**25763123**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25770101	**Good identification**					**25770101**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25770121	**Good identification**					**25770121**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25770123	**Good identification**					**25770123**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25771000	**Low discrimination**					**25771000**
Actinomyces naeslundii		81.15	0.50	SUC 99		
Lactobacillus jensenii		13.69	0.15			
Actinomyces israelii		3.18	0.17			
25771001	**Good identification**					**25771001**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25771002	**Low discrimination**					**25771002**
Actinomyces naeslundii		62.50	0.50	SUC 99		
Actinomyces viscosus		24.48	0.19			
Lactobacillus jensenii		7.03	0.12			
25771003	**Acceptable identification**					**25771003**
Actinomyces naeslundii		91.99	0.53	SUC 99		
Actinomyces viscosus		8.01	0.11			
25771020	**Low discrimination**					**25771020**
Actinomyces naeslundii		84.64	0.50	SUC 99		
Clostridium perfringens		11.79	0.13			
Lactobacillus jensenii		3.57	0.05			
25771021	**Good identification**					**25771021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	SUC 99		
25771022	**Low discrimination**					**25771022**
Actinomyces naeslundii		88.43	0.50	SUC 99		
Clostridium perfringens		5.80	0.08			
Lactobacillus acidophilus		3.29	0.01			
25771023	**Good identification**					**25771023**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25771040	**Good identification**					**25771040**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25771041	**Good identification**					**25771041**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25771042	**Good identification**					**25771042**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25771043	**Good identification**					**25771043**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25771060	**Low discrimination**					**25771060**
Actinomyces naeslundii		86.93	0.37	SUC 99		
Clostridium perfringens		13.07	0.01			
25771061	**Good identification**					**25771061**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25771062	**Good identification**					**25771062**
Actinomyces naeslundii		100.0	0.37	SUC 99		
25771063	**Good identification**					**25771063**
Actinomyces naeslundii		100.0	0.40	SUC 99		
25771100	**Good identification**					**25771100**
					GRA	
Actinomyces naeslundii		92.06	0.58	SUC 99	99	
Actinomyces israelii		5.07	0.27		99	
Fusobacterium mortiferum		1.85	0.17		1	
25771101	**Very good identification**					**25771101**
Actinomyces naeslundii		100.0	0.61	SUC 99		
25771102	**Low discrimination**					**25771102**
Actinomyces naeslundii		67.69	0.58	SUC 99		
Actinomyces viscosus		16.56	0.23			
Lactobacillus acidophilus		10.16	0.19			
25771103	**Good identification**					**25771103**
Actinomyces naeslundii		94.84	0.61	SUC 99		
Actinomyces viscosus		5.16	0.15			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25771110	**Good identification**					**25771110**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25771111	**Good identification**					**25771111**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25771113	**Good identification**					**25771113**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25771120	**Low discrimination**					**25771120**
					SPO	
Actinomyces naeslundii		49.98	0.58	SUC 99	1	
Clostridium septicum		31.29	0.25		99	
Clostridium perfringens		17.94	0.28		84	
25771121	**Very good identification**					**25771121**
Actinomyces naeslundii		100.0	0.61	SUC 99		
25771122	**Low discrimination**					**25771122**
Actinomyces naeslundii		65.74	0.58	SUC 99		
Clostridium perfringens		11.10	0.22			
Lactobacillus acidophilus		9.87	0.19			
25771123	**Very good identification**					**25771123**
Actinomyces naeslundii		100.0	0.61	SUC 99		
25771130	**Good identification**					**25771130**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25771131	**Good identification**					**25771131**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25771132	**Low discrimination**					**25771132**
					GRA	
Actinomyces naeslundii		62.74	0.36	SUC 99	99	
Parabacteroides distasonis		22.90	0.16	MLZ 5	1	
Leptotrichia buccalis		14.36	-.09		1	
25771133	**Good identification**					**25771133**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25771140	**Low discrimination**					**25771140**
Actinomyces naeslundii		89.18	0.45	SUC 99		
Actinomyces israelii		10.82	0.20			
25771141	**Good identification**					**25771141**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25771142	**Low discrimination**					**25771142**
Actinomyces naeslundii		77.69	0.45	SUC 99		
Actinomyces israelii		14.15	0.23			
Lactobacillus acidophilus		8.17	0.04			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25771143	**Good identification**					**25771143**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25771160	**Low discrimination**					**25771160**
Actinomyces naeslundii		72.08	0.45	SUC 99		
Clostridium perfringens		27.92	0.16			
25771161	**Good identification**					**25771161**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25771162	**Low discrimination**					**25771162**
Actinomyces naeslundii		77.68	0.45	SUC 99		
Clostridium perfringens		14.16	0.10			
Lactobacillus acidophilus		8.17	0.04			
25771163	**Good identification**					**25771163**
Actinomyces naeslundii		100.0	0.48	SUC 99		
25772100	**Good identification**					**25772100**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25772101	**Good identification**					**25772101**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25772102	**Good identification**					**25772102**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25772103	**Good identification**					**25772103**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25772120	**Good identification**					**25772120**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25772121	**Good identification**					**25772121**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25772122	**Good identification**					**25772122**
Actinomyces naeslundii		100.0	0.39	SUC 99		
25772123	**Good identification**					**25772123**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25773000	**Low discrimination**					**25773000**
Actinomyces naeslundii		89.73	0.55	SUC 99		
Actinomyces israelii		10.27	0.30			
25773001	**Very good identification**					**25773001**
Actinomyces naeslundii		100.0	0.58	SUC 99		
25773002	**Low discrimination**					**25773002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		85.35	0.55	SUC 99		
Actinomyces israelii		14.65	0.33			
25773003	**Very good identification**					**25773003**
Actinomyces naeslundii		100.0	0.58	SUC 99		
25773011	**Good identification**					**25773011**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25773013	**Good identification**					**25773013**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25773020	**Good identification**					**25773020**
Actinomyces naeslundii		100.0	0.55	SUC 99		
25773021	**Very good identification**					**25773021**
Actinomyces naeslundii		100.0	0.58	SUC 99		
25773022	**Good identification**					**25773022**
Actinomyces naeslundii		100.0	0.55	SUC 99		
25773023	**Very good identification**					**25773023**
Actinomyces naeslundii		100.0	0.58	SUC 99		
25773031	**Good identification**					**25773031**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25773033	**Good identification**					**25773033**
Actinomyces naeslundii		100.0	0.36	SUC 99		
25773040	**Low discrimination**					**25773040**
Actinomyces naeslundii		79.86	0.42	SUC 99		
Actinomyces israelii		20.14	0.23			
25773041	**Good identification**					**25773041**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25773042	**Low discrimination**					**25773042**
Actinomyces naeslundii		72.56	0.42	SUC 99		
Actinomyces israelii		27.44	0.26			
25773043	**Good identification**					**25773043**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25773060	**Good identification**					**25773060**
Actinomyces naeslundii		100.0	0.42	SUC 99		
25773061	**Good identification**					**25773061**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25773062	**Good identification**					**25773062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	SUC 99		
25773063	**Good identification**					**25773063**
Actinomyces naeslundii		100.0	0.45	SUC 99		
25773100	**Low discrimination**					**25773100**
Actinomyces naeslundii		86.15	0.63	SUC 99		
Actinomyces israelii		13.85	0.40			
25773101	**Very good identification**					**25773101**
Actinomyces naeslundii		100.0	0.66	SUC 99		
25773102	**Low discrimination**					**25773102**
Actinomyces naeslundii		80.57	0.63	SUC 99		
Actinomyces israelii		19.43	0.43			
25773103	**Very good identification**					**25773103**
Actinomyces naeslundii		100.0	0.66	SUC 99		
25773110	**Low discrimination**					**25773110**
Actinomyces naeslundii		68.66	0.41	SUC 99		
Actinomyces israelii		31.34	0.26	MLZ 5		
25773111	**Good identification**					**25773111**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25773112	**Low discrimination**					**25773112**
Actinomyces naeslundii		59.36	0.41	SUC 99		
Actinomyces israelii		40.64	0.29	MLZ 5		
25773113	**Good identification**					**25773113**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25773120	**Good identification**					**25773120**
					SPO	
Actinomyces naeslundii		94.66	0.63	SUC 99	1	
Clostridium ramosum		4.37	0.15		92	
Actinomyces israelii		0.97	0.20		1	
25773121	**Very good identification**					**25773121**
Actinomyces naeslundii		100.0	0.66	SUC 99		
25773122	**Low discrimination**					**25773122**
					SPO	
Actinomyces naeslundii		83.33	0.63	SUC 99	1	
Clostridium ramosum		15.39	0.25		92	
Actinomyces israelii		1.28	0.23		1	
25773123	**Very good identification**					**25773123**
Actinomyces naeslundii		100.0	0.66	SUC 99		
25773130	**Good identification**					**25773130**
Actinomyces naeslundii		100.0	0.41	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
25773131	**Good identification**					**25773131**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25773132	**Good identification**					**25773132**
Actinomyces naeslundii		100.0	0.41	SUC 99		
25773133	**Good identification**					**25773133**
Actinomyces naeslundii		100.0	0.44	SUC 99		
25773140	**Low discrimination**					**25773140**
Actinomyces naeslundii		73.84	0.50	SUC 99		
Actinomyces israelii		26.16	0.33			
25773141	**Good identification**					**25773141**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25773142	**Low discrimination**					**25773142**
Actinomyces naeslundii		65.30	0.50	SUC 99		
Actinomyces israelii		34.70	0.36			
25773143	**Good identification**					**25773143**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25773160	**Good identification**					**25773160**
Actinomyces naeslundii		100.0	0.50	SUC 99		
25773161	**Good identification**					**25773161**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25773162	**Good identification**					**25773162**
Actinomyces naeslundii		96.72	0.50	SUC 99		
Actinomyces israelii		3.28	0.16			
25773163	**Good identification**					**25773163**
Actinomyces naeslundii		100.0	0.53	SUC 99		
25773500	**Acceptable identification**					**25773500**
Actinomyces israelii		92.87	0.39	SUC 98		
Actinomyces naeslundii		7.13	0.29	XYL 92		
25773502	**Good identification**					**25773502**
Actinomyces israelii		95.13	0.42	SUC 98		
Actinomyces naeslundii		4.87	0.29	XYL 92		
25775000	**Good identification**					**25775000**
Actinomyces israelii		100.0	0.35	SUC 98		
25775002	**Good identification**					**25775002**
Actinomyces israelii		100.0	0.38	SUC 98		
25775100	**Acceptable identification**					**25775100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		93.13	0.45	SUC 98	1	1	
Clostridium butyricum		3.83	0.01		1	99	
Atopobium parvulum		1.55	0.01		99	1	
25775102	**Good identification**						**25775102**
					SPO		
Actinomyces israelii		97.91	0.48	SUC 98	1		
Clostridium butyricum		1.05	-.07		99		
Actinomyces naeslundii		1.04	0.24		1		
25775122	**Low discrimination**						**25775122**
					GRA		
Parabacteroides distasonis		79.67	0.37	NIT 1	1		
Actinomyces israelii		15.05	0.28		99		
Prevotella oralis		2.77	-.01		1		
25775140	**Good identification**						**25775140**
Actinomyces israelii		100.0	0.38	SUC 98			
25775142	**Good identification**						**25775142**
Actinomyces israelii		100.0	0.41	SUC 98			
25775322	**Good identification**						**25775322**
Parabacteroides distasonis		100.0	0.45	NIT 1			
25775332	**Good identification**						**25775332**
Parabacteroides distasonis		100.0	0.40	NIT 1			
25775402	**Good identification**						**25775402**
Actinomyces israelii		100.0	0.37	SUC 98			
25775500	**Low discrimination**						**25775500**
					GRA	SPO	
Actinomyces israelii		58.83	0.44	SUC 98	99	1	
Clostridium butyricum		26.64	0.17		99	99	
Mitsuokella multacida		14.53	0.21		1	1	
25775502	**Low discrimination**						**25775502**
					SPO		
Actinomyces israelii		84.95	0.47	SUC 98	1		
Clostridium butyricum		9.97	0.10		99		
Bifidobacterium adolescentis		3.52	0.17		1		
25775540	**Low discrimination**						**25775540**
					GRA	SPO	
Actinomyces israelii		57.59	0.37	SUC 98	99	1	
Mitsuokella multacida		36.58	0.21		1	1	
Clostridium butyricum		5.83	-.01		99	99	
25775542	**Good identification**						**25775542**
					GRA		
Actinomyces israelii		95.51	0.40	SUC 98	99		
Mitsuokella multacida		4.49	0.04		1		
25775722	**Good identification**						**25775722**
Parabacteroides distasonis		96.15	0.41	NIT 1			
Clostridium clostridioforme		3.85	0.11				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
25775732	**Low discrimination**					**25775732**
Parabacteroides distasonis		74.68	0.36	NIT 1		
Clostridium clostridioforme		25.33	0.23			
25777000	**Good identification**					**25777000**
Actinomyces israelii		100.0	0.48	SUC 98		
25777002	**Good identification**					**25777002**
Actinomyces israelii		100.0	0.51	SUC 98		
25777012	**Good identification**					**25777012**
Actinomyces israelii		100.0	0.37	SUC 98		
25777040	**Good identification**					**25777040**
Actinomyces israelii		100.0	0.41	SUC 98		
25777042	**Good identification**					**25777042**
Actinomyces israelii		100.0	0.44	SUC 98		
25777100	**Very good identification**					**25777100**
Actinomyces israelii		99.46	0.58	SUC 98		
Actinomyces naeslundii		0.54	0.29			
25777102	**Very good identification**					**25777102**
Actinomyces israelii		99.64	0.61	SUC 98		
Actinomyces naeslundii		0.36	0.29			
25777110	**Good identification**					**25777110**
Actinomyces israelii		100.0	0.44	SUC 98		
25777112	**Good identification**					**25777112**
Actinomyces israelii		100.0	0.47	SUC 98		
25777120	**Acceptable identification**					**25777120**
Actinomyces israelii		92.12	0.38	SUC 98		
Actinomyces naeslundii		7.88	0.29	NAG 6		
25777122	**Acceptable identification**					**25777122**
Actinomyces israelii		94.60	0.41	SUC 98		
Actinomyces naeslundii		5.40	0.29	NAG 6		
25777140	**Good identification**					**25777140**
Actinomyces israelii		100.0	0.51	SUC 98		
25777142	**Good identification**					**25777142**
Actinomyces israelii		100.0	0.54	SUC 98		
25777150	**Good identification**					**25777150**
Actinomyces israelii		100.0	0.37	SUC 98		
25777152	**Good identification**					**25777152**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	SUC 98			
25777300	**Good identification**						**25777300**
Actinomyces israelii		100.0	0.39	SUC 98			
25777302	**Good identification**						**25777302**
Actinomyces israelii		100.0	0.42	SUC 98			
25777342	**Good identification**						**25777342**
Actinomyces israelii		100.0	0.35	SUC 98			
25777400	**Good identification**						**25777400**
Actinomyces israelii		100.0	0.46	SUC 98			
25777402	**Good identification**						**25777402**
Actinomyces israelii		100.0	0.49	SUC 98			
25777412	**Good identification**						**25777412**
Actinomyces israelii		100.0	0.35	SUC 98			
25777440	**Good identification**						**25777440**
Actinomyces israelii		100.0	0.39	SUC 98			
25777442	**Good identification**						**25777442**
Actinomyces israelii		100.0	0.42	SUC 98			
25777500	**Good identification**						**25777500**
					GRA	SPO	
Actinomyces israelii		90.04	0.57	SUC 98	99	1	
Mitsuokella multacida		9.16	0.27		1	1	
Clostridium butyricum		0.80	0.01		99	99	
25777502	**Very good identification**						**25777502**
					GRA		
Actinomyces israelii		98.44	0.60	SUC 98	99		
Mitsuokella multacida		0.74	0.11		1		
Bifidobacterium adolescentis		0.59	0.15		99		
25777510	**Low discrimination**						**25777510**
					GRA		
Actinomyces israelii		81.51	0.43	SUC 98	99		
Mitsuokella multacida		18.50	0.19	MLZ 13	1		
25777512	**Acceptable identification**						**25777512**
Actinomyces israelii		91.84	0.46	SUC 98			
Bifidobacterium adolescentis		4.50	0.17	MLZ 13			
Bifidobacterium dentium		2.11	-.07				
25777520	**Low discrimination**						**25777520**
					GRA		
Actinomyces israelii		71.51	0.36	SUC 98	99		
Mitsuokella multacida		28.49	0.17	NAG 6	1		
25777522	**Good identification**						**25777522**
Actinomyces israelii		100.0	0.39	SUC 98			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
25777540	**Low discrimination**						**25777540**
					GRA		
Actinomyces israelii		79.27	0.50	SUC 98	99		
Mitsuokella multacida		20.73	0.27		1		
25777542	**Good identification**						**25777542**
					GRA		
Actinomyces israelii		98.10	0.53	SUC 98	99		
Mitsuokella multacida		1.90	0.11		1		
25777550	**Low discrimination**						**25777550**
					COC	GRA	
Actinomyces israelii		59.84	0.36	SUC 98	1	99	
Mitsuokella multacida		34.92	0.19	MLZ 13	1	1	
Blautia producta		5.24	-.03		99	99	
25777552	**Good identification**						**25777552**
					GRA		
Actinomyces israelii		95.86	0.39	SUC 98	99		
Mitsuokella multacida		4.14	0.03	MLZ 13	1		
25777702	**Low discrimination**						**25777702**
					GRA		
Actinomyces israelii		84.32	0.41	SUC 98	99		
Mitsuokella multacida		15.68	0.15	RHA 7	1		
26101101	**Very good identification**						**26101101**
Clostridium chauvoei		100.0	0.57	GLU 99			
26101103	**Good identification**						**26101103**
					SPO		
Clostridium chauvoei		98.44	0.44	GLU 99	99		
Actinomyces viscosus		1.56	0.12		1		
26101121	**Good identification**						**26101121**
Clostridium chauvoei		100.0	0.48	GLU 99			
26101123	**Good identification**						**26101123**
Clostridium chauvoei		100.0	0.36	GLU 99			
26110000	**Low discrimination**						**26110000**
					COC	GRA	
Actinomyces odontolyticus		40.81	0.39	GLU 99	1	99	
Gemella morbillorum		29.29	0.32	MNS 9	99	99	
Prevotella intermedia		26.17	0.30		1	1	
26110002	**Low discrimination**						**26110002**
Actinomyces viscosus		85.03	0.41	GLU 99			
Actinomyces odontolyticus		14.98	0.35	RAF 94			
26110102	**Good identification**						**26110102**
Actinomyces viscosus		95.47	0.45	GLU 99			
Actinomyces odontolyticus		2.99	0.27	RAF 94			
Propionibacterium propionicum		1.55	0.29				
26110103	**Acceptable identification**						**26110103**
					SPO		
Actinomyces viscosus		94.92	0.37	GLU 99	1		
Clostridium chauvoei		4.57	0.16	RAF 94	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		0.51	0.12		1		
26110200	**Good identification**						**26110200**
Actinomyces odontolyticus		100.0	0.44	GLU 99			
26110202	**Good identification**						**26110202**
Actinomyces odontolyticus		97.13	0.40	GLU 99			
Actinomyces viscosus		2.87	0.07	MNS 9			
26110300	**Good identification**						**26110300**
Actinomyces odontolyticus		100.0	0.35	GLU 99			
26111000	**Low discrimination**						**26111000**
					COC	GRA	
Actinomyces odontolyticus		36.60	0.36	GLU 99	1	99	
Prevotella intermedia		35.21	0.30	MNS 9	1	1	
Gemella morbillorum		21.22	0.27		99	99	
26111002	**Low discrimination**						**26111002**
Actinomyces viscosus		82.50	0.38	GLU 99			
Actinomyces odontolyticus		15.80	0.32	RAF 94			
Clostridium perfringens		0.99	0.06				
26111020	**Acceptable identification**						**26111020**
					COC		
Clostridium perfringens		90.90	0.38	GLU 99	1		
Actinomyces odontolyticus		5.63	0.22	LAC 88	1		
Gemella morbillorum		2.23	0.11		99		
26111101	**Very good identification**						**26111101**
					SPO		
Clostridium chauvoei		99.94	0.63	GLU 99	99		
Actinomyces naeslundii		0.03	0.26		1		
Actinomyces viscosus		0.02	-.01		1		
26111102	**Low discrimination**						**26111102**
Actinomyces viscosus		86.89	0.42	GLU 99			
Clostridium perfringens		4.28	0.21	RAF 94			
Actinomyces odontolyticus		2.96	0.24				
26111103	**Low discrimination**						**26111103**
					SPO		
Clostridium chauvoei		88.36	0.50	GLU 99	99		
Actinomyces viscosus		11.37	0.34		1		
Actinomyces naeslundii		0.17	0.26		1		
26111120	**Low discrimination**						**26111120**
					GRA	SPO	
Clostridium perfringens		83.14	0.53	GLU 99	99	84	
Prevotella melaninogenica		15.26	0.33		1	1	
Clostridium chauvoei		1.21	0.21		99	99	
26111121	**Good identification**						**26111121**
					GRA	SPO	
Clostridium chauvoei		99.08	0.55	GLU 99	99	99	
Clostridium perfringens		0.69	0.19		99	84	
Prevotella melaninogenica		0.13	-.01		1	1	
26111122	**Low discrimination**						**26111122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Clostridium perfringens		89.86	0.47	GLU 99	99	1	
Prevotella melaninogenica		6.67	0.21		1	1	
Capnocytophaga ochracea		1.91	0.15		1	90	
26111123	**Good identification**						**26111123**
					SPO		
Clostridium chauvoei		97.26	0.42	GLU 99	99		
Clostridium perfringens		1.81	0.13		84		
Actinomyces naeslundii		0.55	0.26		1		
26111160	**Good identification**						**26111160**
					GRA		
Clostridium perfringens		99.04	0.40	GLU 99	99		
Prevotella melaninogenica		0.96	-.01		1		
26111200	**Good identification**						**26111200**
Actinomyces odontolyticus		100.0	0.41	GLU 99			
26111202	**Good identification**						**26111202**
Actinomyces odontolyticus		97.36	0.37	GLU 99			
Actinomyces viscosus		2.64	0.03	MNS 9			
26112100	**Very good identification**						**26112100**
Propionibacterium propionicum		100.0	0.60	GLU 99			
26112101	**Good identification**						**26112101**
Propionibacterium propionicum		100.0	0.44	GLU 99			
26112102	**Very good identification**						**26112102**
Propionibacterium propionicum		99.38	0.63	GLU 99			
Actinomyces viscosus		0.63	0.11				
26112103	**Good identification**						**26112103**
Propionibacterium propionicum		98.15	0.47	GLU 99			
Actinomyces viscosus		1.85	0.03	URE 10			
26112140	**Very good identification**						**26112140**
Propionibacterium propionicum		100.0	0.60	GLU 99			
26112141	**Good identification**						**26112141**
Propionibacterium propionicum		100.0	0.44	GLU 99			
26112142	**Very good identification**						**26112142**
Propionibacterium propionicum		100.0	0.63	GLU 99			
26112143	**Good identification**						**26112143**
Propionibacterium propionicum		100.0	0.47	GLU 99			
26113100	**Very good identification**						**26113100**
Propionibacterium propionicum		99.64	0.60	GLU 99			
Actinomyces naeslundii		0.36	0.28				
26113101	**Low discrimination**						**26113101**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		55.16	0.44	GLU 99	1	
Clostridium chauvoei		42.15	0.29	URE 10	99	
Actinomyces naeslundii		2.69	0.31		1	
26113102	**Very good identification**					**26113102**
Propionibacterium propionicum		99.38	0.63	GLU 99		
Actinomyces viscosus		0.38	0.08			
Actinomyces naeslundii		0.24	0.28			
26113103	**Low discrimination**					**26113103**
					SPO	
Propionibacterium propionicum		88.19	0.47	GLU 99	1	
Clostridium chauvoei		7.93	0.16	URE 10	99	
Actinomyces naeslundii		2.86	0.31		1	
26113140	**Very good identification**					**26113140**
Propionibacterium propionicum		100.0	0.60	GLU 99		
26113141	**Good identification**					**26113141**
Propionibacterium propionicum		100.0	0.44	GLU 99		
26113142	**Very good identification**					**26113142**
Propionibacterium propionicum		100.0	0.63	GLU 99		
26113143	**Good identification**					**26113143**
Propionibacterium propionicum		100.0	0.47	GLU 99		
26114200	**Good identification**					**26114200**
Actinomyces odontolyticus		100.0	0.38	GLU 99		
26115200	**Good identification**					**26115200**
Actinomyces odontolyticus		100.0	0.35	GLU 99		
26121101	**Good identification**					**26121101**
Clostridium chauvoei		100.0	0.40	GLU 99		
26131020	**Good identification**					**26131020**
					CAT	
Clostridium perfringens		99.25	0.43	GLU 99	1	
Propionibacterium granulosum		0.76	0.14	LAC 88	90	
26131022	**Good identification**					**26131022**
Clostridium perfringens		100.0	0.37	GLU 99		
26131101	**Good identification**					**26131101**
					SPO	
Clostridium chauvoei		99.42	0.47	GLU 99	99	
Actinomyces naeslundii		0.58	0.31	TRE 10	1	
26131120	**Very good identification**					**26131120**
					GRA	
Clostridium perfringens		99.71	0.57	GLU 99	99	
Actinomyces naeslundii		0.10	0.28		99	
Prevotella melaninogenica		0.10	-.01		1	
26131121	**Low discrimination**					**26131121**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		88.16	0.38	GLU 99	99	
Clostridium perfringens		10.31	0.23	TRE 10	84	
Actinomyces naeslundii		1.53	0.31		1	
26131122	**Good identification**					**26131122**
Clostridium perfringens		99.79	0.52	GLU 99		
Actinomyces naeslundii		0.21	0.28			
26131160	**Good identification**					**26131160**
Clostridium perfringens		100.0	0.45	GLU 99		
26131162	**Good identification**					**26131162**
Clostridium perfringens		100.0	0.39	GLU 99		
26132100	**Very good identification**					**26132100**
Propionibacterium propionicum		100.0	0.58	GLU 99		
26132101	**Good identification**					**26132101**
Propionibacterium propionicum		100.0	0.42	GLU 99		
26132102	**Very good identification**					**26132102**
Propionibacterium propionicum		100.0	0.61	GLU 99		
26132103	**Good identification**					**26132103**
Propionibacterium propionicum		100.0	0.45	GLU 99		
26132140	**Very good identification**					**26132140**
Propionibacterium propionicum		100.0	0.58	GLU 99		
26132141	**Good identification**					**26132141**
Propionibacterium propionicum		100.0	0.42	GLU 99		
26132142	**Very good identification**					**26132142**
Propionibacterium propionicum		100.0	0.61	GLU 99		
26132143	**Good identification**					**26132143**
Propionibacterium propionicum		100.0	0.45	GLU 99		
26133100	**Very good identification**					**26133100**
Propionibacterium propionicum		99.15	0.58	GLU 99		
Actinomyces naeslundii		0.85	0.33			
26133101	**Low discrimination**					**26133101**
					SPO	
Propionibacterium propionicum		82.01	0.42	GLU 99	1	
Actinomyces naeslundii		9.48	0.36	URE 10	1	
Clostridium chauvoei		8.51	0.12		99	
26133102	**Very good identification**					**26133102**
Propionibacterium propionicum		99.43	0.61	GLU 99		
Actinomyces naeslundii		0.57	0.33			
26133103	**Acceptable identification**					**26133103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Propionibacterium propionicum		91.80	0.45	GLU 99	1	
Actinomyces naeslundii		7.08	0.36	URE 10	1	
Clostridium chauvoei		1.12	-.01		99	
26133121	**Low discrimination**					**26133121**
					SPO	
Actinomyces naeslundii		76.97	0.36	GLU 99	1	
Clostridium chauvoei		23.03	0.04	RAF 92	99	
26133123	**Good identification**					**26133123**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26133140	**Very good identification**					**26133140**
Propionibacterium propionicum		100.0	0.58	GLU 99		
26133141	**Good identification**					**26133141**
Propionibacterium propionicum		98.00	0.42	GLU 99		
Actinomyces naeslundii		2.00	0.23	URE 10		
26133142	**Very good identification**					**26133142**
Propionibacterium propionicum		100.0	0.61	GLU 99		
26133143	**Good identification**					**26133143**
Propionibacterium propionicum		98.66	0.45	GLU 99		
Actinomyces naeslundii		1.34	0.23	URE 10		
26150102	**Good identification**					**26150102**
Actinomyces viscosus		100.0	0.36	GLU 99		
26173100	**Low discrimination**					**26173100**
Actinomyces naeslundii		52.91	0.35	GLU 99		
Propionibacterium propionicum		47.09	0.24	RAF 92		
26173101	**Good identification**					**26173101**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26173103	**Good identification**					**26173103**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26173120	**Good identification**					**26173120**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26173121	**Good identification**					**26173121**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26173122	**Good identification**					**26173122**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26173123	**Good identification**					**26173123**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26310000	**Good identification**					**26310000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.44	GLU 99		
26310002	**Low discrimination**					**26310002**
Actinomyces odontolyticus		56.26	0.40	GLU 99		
Actinomyces viscosus		43.74	0.31	MNS 9		
26310100	**Good identification**					**26310100**
Actinomyces odontolyticus		97.36	0.35	GLU 99		
Actinomyces viscosus		2.64	0.01	MNS 9		
26310102	**Low discrimination**					**26310102**
Actinomyces viscosus		81.40	0.36	GLU 99		
Actinomyces odontolyticus		18.60	0.32	RAF 94		
26310200	**Good identification**					**26310200**
Actinomyces odontolyticus		100.0	0.49	GLU 99		
26310202	**Good identification**					**26310202**
Actinomyces odontolyticus		100.0	0.45	GLU 99		
26310220	**Good identification**					**26310220**
Actinomyces odontolyticus		100.0	0.35	GLU 99		
26310300	**Good identification**					**26310300**
Actinomyces odontolyticus		100.0	0.40	GLU 99		
26310302	**Good identification**					**26310302**
Actinomyces odontolyticus		97.77	0.36	GLU 99		
Actinomyces viscosus		2.23	0.01	MNS 9		
26311000	**Good identification**					**26311000**
Actinomyces odontolyticus		96.93	0.41	GLU 99		
Actinomyces naeslundii		1.15	0.25	MNS 9		
Clostridium perfringens		1.02	0.02			
26311002	**Low discrimination**					**26311002**
Actinomyces odontolyticus		57.66	0.37	GLU 99		
Actinomyces viscosus		41.22	0.28	MNS 9		
Actinomyces naeslundii		1.12	0.25			
26311120	**Good identification**					**26311120**
Clostridium perfringens		96.00	0.42	GLU 99		
Actinomyces odontolyticus		1.99	0.19			
Actinomyces naeslundii		1.31	0.33			
26311122	**Acceptable identification**					**26311122**
Clostridium perfringens		94.71	0.37	GLU 99		
Actinomyces naeslundii		2.74	0.33			
Actinomyces odontolyticus		2.56	0.15			
26311123	**Low discrimination**					**26311123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		57.53	0.36	GLU 99	1	
Clostridium chauvoei		29.06	0.08	RAF 92	99	
Clostridium perfringens		13.42	0.03		84	
26311200	**Good identification**					**26311200**
Actinomyces odontolyticus		100.0	0.46	GLU 99		
26311202	**Good identification**					**26311202**
Actinomyces odontolyticus		100.0	0.42	GLU 99		
26311300	**Good identification**					**26311300**
Actinomyces odontolyticus		100.0	0.37	GLU 99		
26313100	**Low discrimination**					**26313100**
Actinomyces naeslundii		55.89	0.38	GLU 99		
Propionibacterium propionicum		44.11	0.26	RAF 92		
26313101	**Good identification**					**26313101**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26313103	**Acceptable identification**					**26313103**
Actinomyces naeslundii		91.94	0.41	GLU 99		
Propionibacterium propionicum		8.06	0.12	RAF 92		
26313120	**Low discrimination**					**26313120**
Actinomyces naeslundii		72.31	0.38	GLU 99		
Clostridium perfringens		27.69	0.08	RAF 92		
26313121	**Good identification**					**26313121**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26313122	**Low discrimination**					**26313122**
Actinomyces naeslundii		84.73	0.38	GLU 99		
Clostridium perfringens		15.27	0.03	RAF 92		
26313123	**Good identification**					**26313123**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26314000	**Good identification**					**26314000**
Actinomyces odontolyticus		100.0	0.38	GLU 99		
26314200	**Good identification**					**26314200**
Actinomyces odontolyticus		100.0	0.43	GLU 99		
26314202	**Good identification**					**26314202**
Actinomyces odontolyticus		100.0	0.40	GLU 99		
26315000	**Good identification**					**26315000**
Actinomyces odontolyticus		100.0	0.35	GLU 99		
26315200	**Good identification**					**26315200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.40	GLU 99		
26315202	**Good identification**					**26315202**
Actinomyces odontolyticus		100.0	0.37	GLU 99		
26331101	**Low discrimination**					**26331101**
					SPO	
Actinomyces naeslundii		67.04	0.41	GLU 99	1	
Clostridium chauvoei		32.96	0.12	RAF 92	99	
26331103	**Low discrimination**					**26331103**
					SPO	
Actinomyces naeslundii		77.78	0.41	GLU 99	1	
Actinomyces viscosus		15.48	0.05	RAF 92	1	
Clostridium chauvoei		6.75	-.01		99	
26331120	**Good identification**					**26331120**
					COC	
Clostridium perfringens		98.38	0.47	GLU 99	1	
Actinomyces naeslundii		1.40	0.38		1	
Atopobium parvulum		0.22	0.01		99	
26331121	**Low discrimination**					**26331121**
					SPO	
Actinomyces naeslundii		61.05	0.41	GLU 99	1	
Clostridium perfringens		28.95	0.13	RAF 92	84	
Clostridium chauvoei		10.00	0.04		99	
26331122	**Good identification**					**26331122**
Clostridium perfringens		97.07	0.41	GLU 99		
Actinomyces naeslundii		2.93	0.38			
26331123	**Low discrimination**					**26331123**
Actinomyces naeslundii		81.76	0.41	GLU 99		
Clostridium perfringens		18.24	0.07	RAF 92		
26333001	**Good identification**					**26333001**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26333003	**Good identification**					**26333003**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26333021	**Good identification**					**26333021**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26333023	**Good identification**					**26333023**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26333100	**Low discrimination**					**26333100**
Actinomyces naeslundii		75.04	0.43	GLU 99		
Propionibacterium propionicum		24.96	0.24	RAF 92		
26333101	**Good identification**					**26333101**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26333102	**Low discrimination**					**26333102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		66.71	0.43	GLU 99		
Propionibacterium propionicum		33.29	0.27	RAF 92		
26333103	**Good identification**					**26333103**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26333120	**Low discrimination**					**26333120**
Actinomyces naeslundii		73.19	0.43	GLU 99		
Clostridium perfringens		26.81	0.13	RAF 92		
26333121	**Good identification**					**26333121**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26333122	**Low discrimination**					**26333122**
Actinomyces naeslundii		85.30	0.43	GLU 99		
Clostridium perfringens		14.71	0.07	RAF 92		
26333123	**Good identification**					**26333123**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26351101	**Good identification**					**26351101**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26351121	**Good identification**					**26351121**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26351123	**Good identification**					**26351123**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26353001	**Good identification**					**26353001**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26353003	**Good identification**					**26353003**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26353021	**Good identification**					**26353021**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26353023	**Good identification**					**26353023**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26353100	**Low discrimination**					**26353100**
Actinomyces naeslundii		88.85	0.40	GLU 99		
Actinomyces israelii		11.15	0.15	RAF 92		
26353101	**Good identification**					**26353101**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26353102	**Low discrimination**					**26353102**
Actinomyces naeslundii		75.14	0.40	GLU 99		
Actinomyces israelii		14.15	0.18	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		10.71	0.01			
26353103	**Good identification**					**26353103**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26353120	**Good identification**					**26353120**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26353121	**Good identification**					**26353121**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26353122	**Low discrimination**					**26353122**
					SPO	
Actinomyces naeslundii		70.22	0.40	GLU 99	1	
Clostridium barati		11.25	-.01	RAF 92	99	
Lactobacillus acidophilus		10.00	0.01		1	
26353123	**Good identification**					**26353123**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26357102	**Good identification**					**26357102**
Actinomyces israelii		100.0	0.36	GLU 99		
26371001	**Good identification**					**26371001**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26371003	**Good identification**					**26371003**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26371021	**Good identification**					**26371021**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26371023	**Good identification**					**26371023**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26371101	**Good identification**					**26371101**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26371103	**Good identification**					**26371103**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26371121	**Good identification**					**26371121**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26371123	**Good identification**					**26371123**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26373000	**Low discrimination**					**26373000**
Actinomyces naeslundii		85.25	0.37	GLU 99		
Actinomyces israelii		14.75	0.15	RAF 92		
26373001	**Good identification**					**26373001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	GLU 99		
26373002	**Low discrimination**					**26373002**
Actinomyces naeslundii		79.39	0.37	GLU 99		
Actinomyces israelii		20.61	0.18	RAF 92		
26373003	**Good identification**					**26373003**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26373020	**Good identification**					**26373020**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26373021	**Good identification**					**26373021**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26373022	**Good identification**					**26373022**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26373023	**Good identification**					**26373023**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26373100	**Low discrimination**					**26373100**
					SPO	
Actinomyces naeslundii		76.58	0.45	GLU 99	1	
Actinomyces israelii		18.63	0.25	RAF 92	1	
Clostridium tertium		4.79	0.01		99	
26373101	**Good identification**					**26373101**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26373102	**Low discrimination**					**26373102**
Actinomyces naeslundii		55.61	0.45	GLU 99		
Actinomyces israelii		20.29	0.28	RAF 92		
Lactobacillus acidophilus		13.67	0.10			
26373103	**Good identification**					**26373103**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26373120	**Low discrimination**					**26373120**
					SPO	
Actinomyces naeslundii		73.28	0.45	GLU 99	1	
Clostridium tertium		13.76	0.09	RAF 92	99	
Clostridium ramosum		12.97	0.07		92	
26373121	**Good identification**					**26373121**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26373122	**Low discrimination**					**26373122**
					SPO	
Actinomyces naeslundii		39.74	0.45	GLU 99	1	
Clostridium ramosum		28.12	0.17	RAF 92	92	
Clostridium tertium		22.38	0.17		99	
26373123	**Good identification**					**26373123**
Actinomyces naeslundii		100.0	0.48	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26377002	**Good identification**					**26377002**
Actinomyces israelii		100.0	0.36	GLU 99		
26377100	**Good identification**					**26377100**
					SPO	
Actinomyces israelii		97.00	0.43	GLU 99	1	
Clostridium tertium		3.00	0.03		99	
26377102	**Acceptable identification**					**26377102**
					SPO	
Actinomyces israelii		94.18	0.46	GLU 99	1	
Clostridium tertium		5.82	0.11		99	
26377140	**Good identification**					**26377140**
Actinomyces israelii		100.0	0.36	GLU 99		
26377142	**Good identification**					**26377142**
Actinomyces israelii		100.0	0.39	GLU 99		
26377500	**Good identification**					**26377500**
Actinomyces israelii		100.0	0.42	GLU 99		
26377502	**Good identification**					**26377502**
Actinomyces israelii		100.0	0.45	GLU 99		
26377542	**Good identification**					**26377542**
Actinomyces israelii		100.0	0.38	GLU 99		
26473101	**Good identification**					**26473101**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26473103	**Good identification**					**26473103**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26473121	**Good identification**					**26473121**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26473123	**Good identification**					**26473123**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26500002	**Good identification**					**26500002**
Actinomyces viscosus		100.0	0.40	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26500102	**Good identification**					**26500102**
Actinomyces viscosus		100.0	0.44	GLU 99		
26500103	**Good identification**					**26500103**
Actinomyces viscosus		100.0	0.36	GLU 99		
26501002	**Good identification**					**26501002**
Actinomyces viscosus		100.0	0.36	GLU 99		
26501102	**Good identification**					**26501102**
Actinomyces viscosus		100.0	0.40	GLU 99		
26510002	**Very good identification**					**26510002**
Actinomyces viscosus		99.97	0.62	GLU 99		
Actinomyces odontolyticus		0.04	0.09			
26510003	**Good identification**					**26510003**
Actinomyces viscosus		100.0	0.53	GLU 99		
26510102	**Very good identification**					**26510102**
Actinomyces viscosus		99.93	0.66	GLU 99		
Propionibacterium propionicum		0.07	0.26			
26510103	**Very good identification**					**26510103**
Actinomyces viscosus		100.0	0.58	GLU 99		
26511002	**Very good identification**					**26511002**
Actinomyces viscosus		99.72	0.58	GLU 99		
Actinomyces naeslundii		0.15	0.34			
Lactobacillus fermentum		0.09	0.20			
26511003	**Good identification**					**26511003**
Actinomyces viscosus		99.34	0.50	GLU 99		
Actinomyces naeslundii		0.67	0.37			
26511021	**Good identification**					**26511021**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26511102	**Very good identification**					**26511102**
					GRA	
Actinomyces viscosus		99.23	0.62	GLU 99	99	
Actinomyces naeslundii		0.24	0.42		99	
Capnocytophaga ochracea		0.22	0.15		1	
26511103	**Good identification**					**26511103**
					SPO	
Actinomyces viscosus		98.45	0.54	GLU 99	1	
Actinomyces naeslundii		1.06	0.45		1	
Clostridium chauvoei		0.49	0.16		99	
26511120	**Low discrimination**					**26511120**
					GRA	
Prevotella melaninogenica		88.59	0.50	NIT 1	1	
Clostridium perfringens		10.22	0.40	GLU 90	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.59	0.42		99		
26511122	**Low discrimination**						**26511122**
					GRA	CAT	
Prevotella melaninogenica		56.57	0.37	NIT 1	1	1	
Capnocytophaga ochracea		16.22	0.32	GLU 90	1	90	
Clostridium perfringens		16.12	0.35		99	1	
26511123	**Low discrimination**						**26511123**
					GRA		
Actinomyces naeslundii		41.67	0.45	GLU 99	99		
Actinomyces viscosus		39.25	0.20		99		
Prevotella melaninogenica		8.01	0.03		1		
26512100	**Very good identification**						**26512100**
Propionibacterium propionicum		99.74	0.57	GLU 99			
Actinomyces naeslundii		0.26	0.23				
26512101	**Good identification**						**26512101**
Propionibacterium propionicum		96.62	0.41	GLU 99			
Actinomyces naeslundii		3.38	0.26	URE 10			
26512102	**Low discrimination**						**26512102**
Propionibacterium propionicum		86.99	0.60	GLU 99			
Actinomyces viscosus		12.86	0.32				
Actinomyces naeslundii		0.15	0.23				
26512103	**Low discrimination**						**26512103**
Propionibacterium propionicum		68.17	0.44	GLU 99			
Actinomyces viscosus		30.24	0.24	URE 10			
Actinomyces naeslundii		1.59	0.26				
26512140	**Very good identification**						**26512140**
Propionibacterium propionicum		100.0	0.57	GLU 99			
26512141	**Good identification**						**26512141**
Propionibacterium propionicum		100.0	0.41	GLU 99			
26512142	**Very good identification**						**26512142**
Propionibacterium propionicum		100.0	0.60	GLU 99			
26512143	**Good identification**						**26512143**
Propionibacterium propionicum		100.0	0.44	GLU 99			
26513000	**Low discrimination**						**26513000**
					CAT		
Actinomyces naeslundii		55.88	0.39	GLU 99	25		
Propionibacterium propionicum		25.80	0.23		1		
Propionibacterium granulosum		18.32	0.16		90		
26513001	**Good identification**						**26513001**
Actinomyces naeslundii		100.0	0.42	GLU 99			
26513003	**Low discrimination**						**26513003**
Actinomyces naeslundii		56.28	0.42	GLU 99			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		43.72	0.16			
26513020	**Good identification**					**26513020**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26513021	**Good identification**					**26513021**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26513022	**Good identification**					**26513022**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26513023	**Good identification**					**26513023**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26513100	**Good identification**					**26513100**
Propionibacterium propionicum		94.14	0.57	GLU 99		
Actinomyces naeslundii		5.86	0.46			
26513101	**Low discrimination**					**26513101**
Propionibacterium propionicum		54.33	0.41	GLU 99		
Actinomyces naeslundii		45.67	0.49	URE 10		
26513102	**Low discrimination**					**26513102**
Propionibacterium propionicum		88.33	0.60	GLU 99		
Actinomyces viscosus		8.01	0.28			
Actinomyces naeslundii		3.67	0.46			
26513103	**Low discrimination**					**26513103**
Propionibacterium propionicum		54.58	0.44	GLU 99		
Actinomyces naeslundii		30.58	0.49	URE 10		
Actinomyces viscosus		14.84	0.20			
26513120	**Low discrimination**					**26513120**
					GRA	
Actinomyces naeslundii		49.21	0.46	GLU 99	99	
Prevotella melaninogenica		38.38	0.15		1	
Propionibacterium propionicum		7.99	0.23		99	
26513121	**Good identification**					**26513121**
Actinomyces naeslundii		100.0	0.49	GLU 99		
26513122	**Low discrimination**					**26513122**
					GRA	
Actinomyces naeslundii		67.71	0.46	GLU 99	99	
Propionibacterium propionicum		16.48	0.26		99	
Prevotella melaninogenica		10.06	0.03		1	
26513123	**Good identification**					**26513123**
Actinomyces naeslundii		100.0	0.49	GLU 99		
26513140	**Very good identification**					**26513140**
Propionibacterium propionicum		98.91	0.57	GLU 99		
Actinomyces naeslundii		1.09	0.34			
26513141	**Low discrimination**					**26513141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		87.08	0.41	GLU 99		
Actinomyces naeslundii		12.92	0.37	URE 10		
26513142	**Very good identification**					**26513142**
Propionibacterium propionicum		99.27	0.60	GLU 99		
Actinomyces naeslundii		0.73	0.34			
26513143	**Acceptable identification**					**26513143**
Propionibacterium propionicum		91.00	0.44	GLU 99		
Actinomyces naeslundii		9.00	0.37	URE 10		
26513161	**Good identification**					**26513161**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26513163	**Good identification**					**26513163**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26530002	**Good identification**					**26530002**
Actinomyces viscosus		99.24	0.42	GLU 99		
Propionibacterium granulosum		0.76	0.15	TRE 7		
26530102	**Good identification**					**26530102**
Actinomyces viscosus		99.10	0.47	GLU 99		
Propionibacterium propionicum		0.74	0.24	TRE 7		
Actinomyces naeslundii		0.16	0.23			
26530103	**Good identification**					**26530103**
Actinomyces viscosus		99.30	0.38	GLU 99		
Actinomyces naeslundii		0.70	0.26	TRE 7		
26531001	**Good identification**					**26531001**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26531002	**Good identification**					**26531002**
Actinomyces viscosus		95.16	0.39	GLU 99		
Actinomyces naeslundii		3.65	0.39	TRE 7		
Propionibacterium granulosum		1.19	0.15			
26531021	**Good identification**					**26531021**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26531023	**Good identification**					**26531023**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26531100	**Low discrimination**					**26531100**
Actinomyces naeslundii		55.29	0.46	GLU 99		
Clostridium perfringens		28.28	0.19			
Actinomyces viscosus		9.09	0.09			
26531101	**Low discrimination**					**26531101**
					SPO	
Actinomyces naeslundii		84.17	0.49	GLU 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		12.76	0.12		99	
Actinomyces viscosus		3.07	0.01		1	
26531102	**Acceptable identification**					**26531102**
Actinomyces viscosus		91.87	0.43	GLU 99		
Actinomyces naeslundii		5.65	0.46	TRE 7		
Clostridium perfringens		1.36	0.14			
26531120	**Acceptable identification**					**26531120**
					GRA	
Clostridium perfringens		90.28	0.45	GLU 99	99	
Actinomyces naeslundii		5.46	0.46		99	
Prevotella melaninogenica		4.26	0.15		1	
26531121	**Low discrimination**					**26531121**
					SPO	
Actinomyces naeslundii		86.07	0.49	GLU 99	1	
Clostridium perfringens		9.59	0.11		84	
Clostridium chauvoei		4.35	0.04		99	
26531122	**Low discrimination**					**26531122**
Clostridium perfringens		85.17	0.39	GLU 99		
Actinomyces naeslundii		10.94	0.46			
Actinomyces viscosus		1.80	0.09			
26531123	**Acceptable identification**					**26531123**
Actinomyces naeslundii		91.83	0.49	GLU 99		
Clostridium perfringens		4.81	0.05			
Actinomyces viscosus		3.35	0.01			
26531141	**Good identification**					**26531141**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26531143	**Low discrimination**					**26531143**
Actinomyces naeslundii		82.86	0.37	GLU 99		
Actinomyces viscosus		17.15	0.01			
26531161	**Good identification**					**26531161**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26531163	**Good identification**					**26531163**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26532100	**Good identification**					**26532100**
Propionibacterium propionicum		99.39	0.55	GLU 99		
Actinomyces naeslundii		0.61	0.28			
26532101	**Acceptable identification**					**26532101**
Propionibacterium propionicum		92.33	0.39	GLU 99		
Actinomyces naeslundii		7.67	0.31	URE 10		
26532102	**Very good identification**					**26532102**
Propionibacterium propionicum		98.26	0.58	GLU 99		
Actinomyces viscosus		1.34	0.12			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.40	0.28			
26532103	**Acceptable identification**					**26532103**
Propionibacterium propionicum		91.23	0.42	GLU 99		
Actinomyces naeslundii		5.05	0.31	URE 10		
Actinomyces viscosus		3.72	0.04			
26532140	**Good identification**					**26532140**
Propionibacterium propionicum		100.0	0.55	GLU 99		
26532141	**Good identification**					**26532141**
Propionibacterium propionicum		100.0	0.39	GLU 99		
26532142	**Very good identification**					**26532142**
Propionibacterium propionicum		100.0	0.58	GLU 99		
26532143	**Good identification**					**26532143**
Propionibacterium propionicum		100.0	0.42	GLU 99		
26533000	**Low discrimination**					**26533000**
					CAT	
Actinomyces naeslundii		58.77	0.44	GLU 99	25	
Propionibacterium granulosum		29.79	0.24		90	
Propionibacterium propionicum		11.44	0.21		1	
26533001	**Good identification**					**26533001**
Actinomyces naeslundii		100.0	0.47	GLU 99		
26533002	**Low discrimination**					**26533002**
Actinomyces naeslundii		64.34	0.44	GLU 99		
Propionibacterium propionicum		18.78	0.24			
Actinomyces viscosus		8.72	0.05			
26533003	**Good identification**					**26533003**
Actinomyces naeslundii		100.0	0.47	GLU 99		
26533020	**Low discrimination**					**26533020**
Actinomyces naeslundii		88.76	0.44	GLU 99		
Propionibacterium granulosum		11.24	0.13			
26533021	**Good identification**					**26533021**
Actinomyces naeslundii		100.0	0.47	GLU 99		
26533022	**Good identification**					**26533022**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26533023	**Good identification**					**26533023**
Actinomyces naeslundii		100.0	0.47	GLU 99		
26533100	**Low discrimination**					**26533100**
Propionibacterium propionicum		87.13	0.55	GLU 99		
Actinomyces naeslundii		12.87	0.51			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
26533101	**Low discrimination**					**26533101**
Actinomyces naeslundii		66.60	0.54	GLU 99		
Propionibacterium propionicum		33.40	0.39			
26533102	**Good identification**					**26533102**
Propionibacterium propionicum		90.35	0.58	GLU 99		
Actinomyces naeslundii		8.90	0.51			
Actinomyces viscosus		0.75	0.09			
26533103	**Low discrimination**					**26533103**
Actinomyces naeslundii		56.46	0.54	GLU 99		
Propionibacterium propionicum		42.47	0.42			
Actinomyces viscosus		1.06	0.01			
26533120	**Low discrimination**					**26533120**
Actinomyces naeslundii		86.62	0.51	GLU 99		
Clostridium perfringens		7.45	0.11			
Propionibacterium propionicum		5.92	0.21			
26533121	**Good identification**					**26533121**
Actinomyces naeslundii		100.0	0.54	GLU 99		
26533122	**Low discrimination**					**26533122**
Actinomyces naeslundii		87.48	0.51	GLU 99		
Propionibacterium propionicum		8.97	0.24			
Clostridium perfringens		3.54	0.05			
26533123	**Good identification**					**26533123**
Actinomyces naeslundii		100.0	0.54	GLU 99		
26533140	**Good identification**					**26533140**
Propionibacterium propionicum		97.46	0.55	GLU 99		
Actinomyces naeslundii		2.54	0.38			
26533141	**Low discrimination**					**26533141**
Propionibacterium propionicum		73.97	0.39	GLU 99		
Actinomyces naeslundii		26.03	0.41	URE 10		
26533142	**Very good identification**					**26533142**
Propionibacterium propionicum		98.29	0.58	GLU 99		
Actinomyces naeslundii		1.71	0.38			
26533143	**Low discrimination**					**26533143**
Propionibacterium propionicum		81.00	0.42	GLU 99		
Actinomyces naeslundii		19.00	0.41	URE 10		
26533160	**Low discrimination**					**26533160**
Actinomyces naeslundii		72.07	0.38	GLU 99		
Propionibacterium propionicum		27.93	0.21			
26533161	**Good identification**					**26533161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	GLU 99		
26533162	**Low discrimination**					**26533162**
Actinomyces naeslundii		63.24	0.38	GLU 99		
Propionibacterium propionicum		36.76	0.24			
26533163	**Good identification**					**26533163**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26550002	**Good identification**					**26550002**
Actinomyces viscosus		100.0	0.52	GLU 99		
26550003	**Good identification**					**26550003**
Actinomyces viscosus		100.0	0.44	GLU 99		
26550102	**Good identification**					**26550102**
Actinomyces viscosus		100.0	0.56	GLU 99		
26550103	**Good identification**					**26550103**
Actinomyces viscosus		99.86	0.48	GLU 99		
Actinomyces naeslundii		0.14	0.23			
26551001	**Low discrimination**					**26551001**
Actinomyces naeslundii		76.78	0.39	GLU 99		
Actinomyces viscosus		23.22	0.06			
26551002	**Good identification**					**26551002**
Actinomyces viscosus		99.26	0.48	GLU 99		
Actinomyces naeslundii		0.74	0.36			
26551003	**Good identification**					**26551003**
Actinomyces viscosus		96.77	0.40	GLU 99		
Actinomyces naeslundii		3.23	0.39			
26551020	**Low discrimination**					**26551020**
					GRA	
Actinomyces naeslundii		63.61	0.36	GLU 99	99	
Prevotella melaninogenica		22.97	-.01		1	
Clostridium perfringens		13.42	0.02		99	
26551021	**Good identification**					**26551021**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26551023	**Low discrimination**					**26551023**
Actinomyces naeslundii		76.78	0.39	GLU 99		
Actinomyces viscosus		23.22	0.06			
26551100	**Low discrimination**					**26551100**
					GRA	
Actinomyces naeslundii		50.58	0.44	GLU 99	99	
Actinomyces viscosus		43.00	0.18		99	
Prevotella melaninogenica		6.42	-.01		1	
26551101	**Low discrimination**					**26551101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.11	0.47	GLU 99		
Actinomyces viscosus		15.89	0.10			
26551102	**Good identification**					**26551102**
					GRA	
Actinomyces viscosus		98.63	0.52	GLU 99	99	
Actinomyces naeslundii		1.17	0.44		99	
Capnocytophaga ochracea		0.20	0.05		1	
26551103	**Acceptable identification**					**26551103**
Actinomyces viscosus		94.92	0.44	GLU 99		
Actinomyces naeslundii		5.08	0.47			
26551121	**Acceptable identification**					**26551121**
					GRA	
Actinomyces naeslundii		92.20	0.47	GLU 99	99	
Prevotella melaninogenica		7.80	-.01		1	
26551123	**Low discrimination**					**26551123**
Actinomyces naeslundii		84.11	0.47	GLU 99		
Actinomyces viscosus		15.89	0.10			
26553000	**Good identification**					**26553000**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26553001	**Good identification**					**26553001**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26553002	**Low discrimination**					**26553002**
Actinomyces naeslundii		58.79	0.41	GLU 99		
Actinomyces viscosus		41.21	0.14			
26553003	**Low discrimination**					**26553003**
Actinomyces naeslundii		86.52	0.44	GLU 99		
Actinomyces viscosus		13.48	0.06			
26553020	**Good identification**					**26553020**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26553021	**Good identification**					**26553021**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26553022	**Good identification**					**26553022**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26553023	**Good identification**					**26553023**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26553100	**Low discrimination**					**26553100**
Actinomyces naeslundii		89.10	0.49	GLU 99		
Propionibacterium propionicum		10.91	0.23			
26553101	**Good identification**					**26553101**
Actinomyces naeslundii		100.0	0.52	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26553102	**Low discrimination**					**26553102**
Actinomyces naeslundii		61.67	0.49	GLU 99		
Actinomyces viscosus		27.01	0.18			
Propionibacterium propionicum		11.32	0.26			
26553103	**Acceptable identification**					**26553103**
Actinomyces naeslundii		91.13	0.52	GLU 99		
Actinomyces viscosus		8.87	0.10			
26553120	**Acceptable identification**					**26553120**
					GRA	
Actinomyces naeslundii		93.86	0.49	GLU 99	99	
Prevotella melaninogenica		6.14	-.01		1	
26553121	**Good identification**					**26553121**
Actinomyces naeslundii		100.0	0.52	GLU 99		
26553122	**Good identification**					**26553122**
Actinomyces naeslundii		100.0	0.49	GLU 99		
26553123	**Good identification**					**26553123**
Actinomyces naeslundii		100.0	0.52	GLU 99		
26553140	**Low discrimination**					**26553140**
Actinomyces naeslundii		59.05	0.36	GLU 99		
Propionibacterium propionicum		40.95	0.23			
26553141	**Good identification**					**26553141**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26553143	**Good identification**					**26553143**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26553160	**Good identification**					**26553160**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26553161	**Good identification**					**26553161**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26553162	**Good identification**					**26553162**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26553163	**Good identification**					**26553163**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26570102	**Good identification**					**26570102**
Actinomyces viscosus		99.22	0.37	GLU 99		
Actinomyces naeslundii		0.78	0.25	TRE 7		
26571000	**Good identification**					**26571000**
Actinomyces naeslundii		100.0	0.41	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26571001	**Good identification**					**26571001**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26571020	**Low discrimination**					**26571020**
Actinomyces naeslundii		83.22	0.41	GLU 99		
Clostridium perfringens		16.79	0.07			
26571021	**Good identification**					**26571021**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26571022	**Acceptable identification**					**26571022**
Actinomyces naeslundii		91.33	0.41	GLU 99		
Clostridium perfringens		8.67	0.01			
26571023	**Good identification**					**26571023**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26571100	**Good identification**					**26571100**
Actinomyces naeslundii		96.81	0.49	GLU 99		
Actinomyces viscosus		3.19	-.01			
26571101	**Good identification**					**26571101**
Actinomyces naeslundii		100.0	0.52	GLU 99		
26571103	**Low discrimination**					**26571103**
Actinomyces naeslundii		57.97	0.52	GLU 99		
Actinomyces viscosus		42.03	0.25			
26571120	**Low discrimination**					**26571120**
					GRA	
Actinomyces naeslundii		63.08	0.49	GLU 99	99	
Clostridium perfringens		32.79	0.21		99	
Prevotella melaninogenica		4.12	-.01		1	
26571121	**Good identification**					**26571121**
Actinomyces naeslundii		100.0	0.52	GLU 99		
26571122	**Low discrimination**					**26571122**
Actinomyces naeslundii		78.27	0.49	GLU 99		
Clostridium perfringens		19.15	0.16			
Actinomyces viscosus		2.58	-.01			
26571123	**Good identification**					**26571123**
Actinomyces naeslundii		100.0	0.52	GLU 99		
26571140	**Good identification**					**26571140**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26571141	**Good identification**					**26571141**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26571142	**Low discrimination**					**26571142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.26	0.36	GLU 99		
Actinomyces viscosus		15.74	-.01			
26571143	**Good identification**					**26571143**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26571160	**Low discrimination**					**26571160**
Actinomyces naeslundii		64.06	0.36	GLU 99		
Clostridium perfringens		35.94	0.09			
26571161	**Good identification**					**26571161**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26571162	**Low discrimination**					**26571162**
Actinomyces naeslundii		79.12	0.36	GLU 99		
Clostridium perfringens		20.89	0.03			
26571163	**Good identification**					**26571163**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26573000	**Good identification**					**26573000**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26573001	**Good identification**					**26573001**
Actinomyces naeslundii		100.0	0.49	GLU 99		
26573002	**Good identification**					**26573002**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26573003	**Good identification**					**26573003**
Actinomyces naeslundii		100.0	0.49	GLU 99		
26573020	**Good identification**					**26573020**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26573021	**Good identification**					**26573021**
Actinomyces naeslundii		100.0	0.49	GLU 99		
26573022	**Good identification**					**26573022**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26573023	**Good identification**					**26573023**
Actinomyces naeslundii		100.0	0.49	GLU 99		
26573041	**Good identification**					**26573041**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26573043	**Good identification**					**26573043**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26573061	**Good identification**					**26573061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	GLU 99		
26573063	**Good identification**					**26573063**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26573100	**Acceptable identification**					**26573100**
Actinomyces naeslundii		92.93	0.53	GLU 99		
Propionibacterium propionicum		4.79	0.21			
Actinomyces israelii		2.28	0.17			
26573101	**Very good identification**					**26573101**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26573102	**Low discrimination**					**26573102**
Actinomyces naeslundii		88.40	0.53	GLU 99		
Propionibacterium propionicum		6.84	0.24			
Actinomyces israelii		3.25	0.20			
26573103	**Very good identification**					**26573103**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26573120	**Good identification**					**26573120**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26573121	**Very good identification**					**26573121**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26573122	**Good identification**					**26573122**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26573123	**Very good identification**					**26573123**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26573140	**Low discrimination**					**26573140**
Actinomyces naeslundii		77.38	0.41	GLU 99		
Propionibacterium propionicum		22.62	0.21			
26573141	**Good identification**					**26573141**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26573142	**Low discrimination**					**26573142**
Actinomyces naeslundii		69.52	0.41	GLU 99		
Propionibacterium propionicum		30.48	0.24			
26573143	**Good identification**					**26573143**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26573160	**Good identification**					**26573160**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26573161	**Good identification**					**26573161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	GLU 99		
26573162	**Good identification**					**26573162**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26573163	**Good identification**					**26573163**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26577102	**Good identification**					**26577102**
Actinomyces israelii		100.0	0.38	GLU 99		
26577502	**Good identification**					**26577502**
Actinomyces israelii		100.0	0.36	GLU 99		
26613100	**Good identification**					**26613100**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26613101	**Good identification**					**26613101**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26613102	**Good identification**					**26613102**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26613103	**Good identification**					**26613103**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26613120	**Good identification**					**26613120**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26613121	**Good identification**					**26613121**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26613122	**Good identification**					**26613122**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26613123	**Good identification**					**26613123**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26631100	**Low discrimination**					**26631100**
					GRA	
Actinomyces naeslundii		63.37	0.35	GLU 99	99	
Fusobacterium mortiferum		36.63	0.20	MLT 94	1	
26631101	**Good identification**					**26631101**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26631102	**Low discrimination**					**26631102**
Actinomyces naeslundii		83.81	0.35	GLU 99		
Actinomyces viscosus		16.19	-.01	MLT 94		
26631103	**Good identification**					**26631103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	GLU 99		
26631120	**Low discrimination**					**26631120**
Actinomyces naeslundii		84.43	0.35	GLU 99		
Clostridium perfringens		15.58	0.01	MLT 94		
26631121	**Good identification**					**26631121**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26631122	**Good identification**					**26631122**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26631123	**Good identification**					**26631123**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26633001	**Good identification**					**26633001**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26633003	**Good identification**					**26633003**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26633021	**Good identification**					**26633021**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26633023	**Good identification**					**26633023**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26633100	**Good identification**					**26633100**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26633101	**Good identification**					**26633101**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26633102	**Good identification**					**26633102**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26633103	**Good identification**					**26633103**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26633120	**Good identification**					**26633120**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26633121	**Good identification**					**26633121**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26633122	**Good identification**					**26633122**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26633123	**Good identification**					**26633123**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26651101	**Good identification**					**26651101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	GLU 99		
26651103	**Low discrimination**					**26651103**
Actinomyces naeslundii		81.84	0.36	GLU 99		
Actinomyces viscosus		18.16	0.00	MLT 94		
26651121	**Good identification**					**26651121**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26651123	**Good identification**					**26651123**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26653100	**Good identification**					**26653100**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26653101	**Good identification**					**26653101**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26653102	**Good identification**					**26653102**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26653103	**Good identification**					**26653103**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26653120	**Good identification**					**26653120**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26653121	**Good identification**					**26653121**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26653122	**Good identification**					**26653122**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26653123	**Good identification**					**26653123**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26671100	**Low discrimination**					**26671100**
					GRA	
Actinomyces naeslundii		87.31	0.37	GLU 99	99	
Fusobacterium mortiferum		12.69	0.12	MLT 94	1	
26671101	**Good identification**					**26671101**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26671102	**Low discrimination**					**26671102**
Actinomyces naeslundii		86.95	0.37	GLU 99		
Lactobacillus acidophilus		13.05	-.01	MLT 94		
26671103	**Good identification**					**26671103**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26671120	**Good identification**					**26671120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GLU 99		
26671121	**Good identification**					**26671121**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26671122	**Low discrimination**					**26671122**
Actinomyces naeslundii		86.95	0.37	GLU 99		
Lactobacillus acidophilus		13.05	-.01	MLT 94		
26671123	**Good identification**					**26671123**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26673001	**Good identification**					**26673001**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26673003	**Good identification**					**26673003**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26673021	**Good identification**					**26673021**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26673023	**Good identification**					**26673023**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26673100	**Good identification**					**26673100**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26673101	**Good identification**					**26673101**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26673102	**Good identification**					**26673102**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26673103	**Good identification**					**26673103**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26673120	**Good identification**					**26673120**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26673121	**Good identification**					**26673121**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26673122	**Good identification**					**26673122**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26673123	**Good identification**					**26673123**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26703101	**Good identification**					**26703101**
Actinomyces naeslundii		100.0	0.37	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26703103	**Good identification**					**26703103**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26703121	**Good identification**					**26703121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26703123	**Good identification**					**26703123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26710002	**Good identification**					**26710002**
Actinomyces viscosus		99.75	0.52	GLU 99		
Actinomyces odontolyticus		0.25	0.14			
26710003	**Good identification**					**26710003**
Actinomyces viscosus		99.77	0.44	GLU 99		
Actinomyces naeslundii		0.23	0.23			
26710102	**Good identification**					**26710102**
Actinomyces viscosus		99.87	0.56	GLU 99		
Actinomyces naeslundii		0.08	0.27			
Actinomyces odontolyticus		0.05	0.06			
26710103	**Good identification**					**26710103**
Actinomyces viscosus		99.64	0.48	GLU 99		
Actinomyces naeslundii		0.36	0.30			
26711000	**Low discrimination**					**26711000**
Actinomyces naeslundii		57.61	0.43	GLU 99		
Actinomyces viscosus		29.31	0.14			
Actinomyces odontolyticus		13.08	0.15			
26711001	**Low discrimination**					**26711001**
Actinomyces naeslundii		89.84	0.46	GLU 99		
Actinomyces viscosus		10.16	0.06			
26711002	**Good identification**					**26711002**
Actinomyces viscosus		97.79	0.48	GLU 99		
Actinomyces naeslundii		1.94	0.43			
Actinomyces odontolyticus		0.27	0.11			
26711003	**Acceptable identification**					**26711003**
Actinomyces viscosus		91.80	0.40	GLU 99		
Actinomyces naeslundii		8.20	0.46			
26711020	**Low discrimination**					**26711020**
Actinomyces naeslundii		67.89	0.43	GLU 99		
Clostridium perfringens		32.11	0.15			
26711021	**Good identification**					**26711021**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26711022	**Low discrimination**					**26711022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		57.76	0.43	GLU 99		
Actinomyces viscosus		29.39	0.14			
Clostridium perfringens		12.86	0.10			
26711023	**Low discrimination**					**26711023**
Actinomyces naeslundii		89.84	0.46	GLU 99		
Actinomyces viscosus		10.16	0.06			
26711100	**Low discrimination**					**26711100**
Actinomyces naeslundii		70.91	0.51	GLU 99		
Actinomyces viscosus		22.54	0.18			
Clostridium perfringens		2.67	0.04			
26711101	**Acceptable identification**					**26711101**
Actinomyces naeslundii		93.40	0.54	GLU 99		
Actinomyces viscosus		6.60	0.10			
26711102	**Good identification**					**26711102**
Actinomyces viscosus		96.86	0.52	GLU 99		
Actinomyces naeslundii		3.08	0.51			
Bifidobacterium longum subsp. infantis		0.07	-.07			
26711103	**Low discrimination**					**26711103**
Actinomyces viscosus		87.49	0.44	GLU 99		
Actinomyces naeslundii		12.51	0.54			
26711121	**Good identification**					**26711121**
Actinomyces naeslundii		100.0	0.54	GLU 99		
26711122	**Low discrimination**					**26711122**
Actinomyces naeslundii		46.43	0.51	GLU 99		
Clostridium perfringens		26.63	0.24			
Actinomyces viscosus		14.76	0.18			
26711123	**Acceptable identification**					**26711123**
Actinomyces naeslundii		93.40	0.54	GLU 99		
Actinomyces viscosus		6.60	0.10			
26711140	**Good identification**					**26711140**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26711141	**Good identification**					**26711141**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26711143	**Low discrimination**					**26711143**
Actinomyces naeslundii		71.42	0.41	GLU 99		
Actinomyces viscosus		28.58	0.10			
26711161	**Good identification**					**26711161**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26711162	**Low discrimination**					**26711162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		61.76	0.38	GLU 99		
Clostridium perfringens		38.24	0.12			
26711163	**Good identification**					**26711163**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26712101	**Good identification**					**26712101**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26712121	**Good identification**					**26712121**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26712123	**Good identification**					**26712123**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26713000	**Good identification**					**26713000**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26713001	**Good identification**					**26713001**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26713002	**Low discrimination**					**26713002**
Actinomyces naeslundii		79.23	0.48	GLU 99		
Actinomyces viscosus		20.77	0.14			
26713003	**Acceptable identification**					**26713003**
Actinomyces naeslundii		94.50	0.51	GLU 99		
Actinomyces viscosus		5.51	0.06			
26713020	**Good identification**					**26713020**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26713021	**Good identification**					**26713021**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26713022	**Good identification**					**26713022**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26713023	**Good identification**					**26713023**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26713040	**Good identification**					**26713040**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26713041	**Good identification**					**26713041**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26713042	**Good identification**					**26713042**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26713043	**Good identification**					**26713043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	GLU 99		
26713060	**Good identification**					**26713060**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26713061	**Good identification**					**26713061**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26713062	**Good identification**					**26713062**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26713063	**Good identification**					**26713063**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26713100	**Good identification**					**26713100**
Actinomyces naeslundii		95.62	0.56	GLU 99		
Propionibacterium propionicum		4.38	0.23			
26713101	**Very good identification**					**26713101**
Actinomyces naeslundii		100.0	0.59	GLU 99		
26713102	**Low discrimination**					**26713102**
Actinomyces naeslundii		81.15	0.56	GLU 99		
Actinomyces viscosus		13.29	0.18			
Propionibacterium propionicum		5.57	0.26			
26713103	**Very good identification**					**26713103**
Actinomyces naeslundii		96.49	0.59	GLU 99		
Actinomyces viscosus		3.51	0.10			
26713111	**Good identification**					**26713111**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26713113	**Good identification**					**26713113**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26713120	**Good identification**					**26713120**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26713121	**Very good identification**					**26713121**
Actinomyces naeslundii		100.0	0.59	GLU 99		
26713122	**Good identification**					**26713122**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26713123	**Very good identification**					**26713123**
Actinomyces naeslundii		100.0	0.59	GLU 99		
26713131	**Good identification**					**26713131**
Actinomyces naeslundii		100.0	0.37	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26713133	**Good identification**					**26713133**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26713140	**Low discrimination**					**26713140**
Actinomyces naeslundii		79.41	0.43	GLU 99		
Propionibacterium propionicum		20.59	0.23			
26713141	**Good identification**					**26713141**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26713142	**Low discrimination**					**26713142**
Actinomyces naeslundii		72.00	0.43	GLU 99		
Propionibacterium propionicum		28.00	0.26			
26713143	**Good identification**					**26713143**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26713160	**Good identification**					**26713160**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26713161	**Good identification**					**26713161**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26713162	**Good identification**					**26713162**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26713163	**Good identification**					**26713163**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26721101	**Good identification**					**26721101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26721103	**Low discrimination**					**26721103**
Actinomyces naeslundii		78.67	0.37	GLU 99		
Actinomyces viscosus		21.33	0.03	FRU 95		
26721121	**Good identification**					**26721121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26721123	**Good identification**					**26721123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26723100	**Good identification**					**26723100**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26723101	**Good identification**					**26723101**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26723102	**Good identification**					**26723102**
Actinomyces naeslundii		100.0	0.39	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26723103	**Good identification**					**26723103**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26723120	**Good identification**					**26723120**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26723121	**Good identification**					**26723121**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26723122	**Good identification**					**26723122**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26723123	**Good identification**					**26723123**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26730101	**Good identification**					**26730101**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26730102	**Good identification**					**26730102**
Actinomyces viscosus		97.95	0.37	GLU 99		
Actinomyces naeslundii		2.05	0.32	TRE 7		
26730121	**Good identification**					**26730121**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26730123	**Good identification**					**26730123**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26731000	**Good identification**					**26731000**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26731001	**Good identification**					**26731001**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26731003	**Low discrimination**					**26731003**
Actinomyces naeslundii		69.73	0.51	GLU 99		
Actinomyces viscosus		30.27	0.21			
26731020	**Low discrimination**					**26731020**
Actinomyces naeslundii		68.84	0.48	GLU 99		
Clostridium perfringens		31.16	0.20			
26731021	**Good identification**					**26731021**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26731022	**Low discrimination**					**26731022**
Actinomyces naeslundii		82.44	0.48	GLU 99		
Clostridium perfringens		17.56	0.14			
26731023	**Good identification**					**26731023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	GLU 99		
26731040	**Good identification**					**26731040**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26731041	**Good identification**					**26731041**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26731042	**Good identification**					**26731042**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26731043	**Good identification**					**26731043**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26731060	**Low discrimination**					**26731060**
Actinomyces naeslundii		67.18	0.35	GLU 99		
Clostridium perfringens		32.82	0.08			
26731061	**Good identification**					**26731061**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26731062	**Low discrimination**					**26731062**
Actinomyces naeslundii		81.31	0.35	GLU 99		
Clostridium perfringens		18.70	0.02			
26731063	**Good identification**					**26731063**
Actinomyces naeslundii		100.0	0.38	GLU 99		
26731100	**Good identification**					**26731100**
Actinomyces naeslundii		95.39	0.56	GLU 99		
Clostridium perfringens		3.44	0.09			
Actinomyces viscosus		1.18	-.01			
26731101	**Very good identification**					**26731101**
Actinomyces naeslundii		100.0	0.59	GLU 99		
26731103	**Low discrimination**					**26731103**
Actinomyces naeslundii		78.67	0.59	GLU 99		
Actinomyces viscosus		21.33	0.25			
26731111	**Good identification**					**26731111**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26731113	**Good identification**					**26731113**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26731121	**Very good identification**					**26731121**
Actinomyces naeslundii		99.22	0.59	GLU 99		
Clostridium perfringens		0.78	0.01			
26731122	**Low discrimination**					**26731122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		64.06	0.56	GLU 99		
Clostridium perfringens		35.15	0.29			
Actinomyces viscosus		0.79	-.01			
26731123	**Very good identification**					**26731123**
Actinomyces naeslundii		100.0	0.59	GLU 99		
26731131	**Good identification**					**26731131**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26731133	**Good identification**					**26731133**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26731140	**Good identification**					**26731140**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26731141	**Good identification**					**26731141**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26731142	**Acceptable identification**					**26731142**
Actinomyces naeslundii		93.47	0.43	GLU 99		
Actinomyces viscosus		6.53	-.01			
26731143	**Good identification**					**26731143**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26731161	**Good identification**					**26731161**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26731162	**Low discrimination**					**26731162**
Actinomyces naeslundii		62.80	0.43	GLU 99		
Clostridium perfringens		37.20	0.17			
26731163	**Good identification**					**26731163**
Actinomyces naeslundii		100.0	0.46	GLU 99		
26732100	**Low discrimination**					**26732100**
Actinomyces naeslundii		68.36	0.37	GLU 99		
Propionibacterium propionicum		31.64	0.21	GAL 96		
26732101	**Good identification**					**26732101**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26732102	**Low discrimination**					**26732102**
Actinomyces naeslundii		51.47	0.37	GLU 99		
Propionibacterium propionicum		35.74	0.24	GAL 96		
Actinomyces viscosus		12.79	0.03			
26732103	**Good identification**					**26732103**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26732120	**Good identification**					**26732120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GLU 99		
26732121	**Good identification**					**26732121**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26732122	**Good identification**					**26732122**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26732123	**Good identification**					**26732123**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26733000	**Good identification**					**26733000**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26733001	**Good identification**					**26733001**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26733002	**Good identification**					**26733002**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26733003	**Good identification**					**26733003**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26733020	**Good identification**					**26733020**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26733021	**Good identification**					**26733021**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26733022	**Good identification**					**26733022**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26733023	**Good identification**					**26733023**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26733040	**Good identification**					**26733040**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26733041	**Good identification**					**26733041**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26733042	**Good identification**					**26733042**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26733043	**Good identification**					**26733043**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26733060	**Good identification**					**26733060**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26733061	**Good identification**					**26733061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	GLU 99		
26733062	**Good identification**					**26733062**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26733063	**Good identification**					**26733063**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26733100	**Very good identification**					**26733100**
Actinomyces naeslundii		97.35	0.61	GLU 99		
Propionibacterium propionicum		1.88	0.21			
Actinomyces israelii		0.77	0.16			
26733101	**Very good identification**					**26733101**
Actinomyces naeslundii		100.0	0.64	GLU 99		
26733102	**Very good identification**					**26733102**
Actinomyces naeslundii		95.49	0.61	GLU 99		
Propionibacterium propionicum		2.76	0.24			
Actinomyces israelii		1.14	0.19			
26733103	**Very good identification**					**26733103**
Actinomyces naeslundii		100.0	0.64	GLU 99		
26733110	**Good identification**					**26733110**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26733111	**Good identification**					**26733111**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26733112	**Good identification**					**26733112**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26733113	**Good identification**					**26733113**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26733120	**Very good identification**					**26733120**
Actinomyces naeslundii		99.40	0.61	GLU 99		
Clostridium perfringens		0.60	0.01			
26733121	**Very good identification**					**26733121**
Actinomyces naeslundii		100.0	0.64	GLU 99		
26733122	**Very good identification**					**26733122**
Actinomyces naeslundii		100.0	0.61	GLU 99		
26733123	**Very good identification**					**26733123**
Actinomyces naeslundii		100.0	0.64	GLU 99		
26733130	**Good identification**					**26733130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	GLU 99		
26733131	**Good identification**					**26733131**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26733132	**Good identification**					**26733132**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26733133	**Good identification**					**26733133**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26733140	**Acceptable identification**					**26733140**
Actinomyces naeslundii		90.15	0.48	GLU 99		
Propionibacterium propionicum		9.85	0.21			
26733141	**Good identification**					**26733141**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26733142	**Low discrimination**					**26733142**
Actinomyces naeslundii		85.92	0.48	GLU 99		
Propionibacterium propionicum		14.09	0.24			
26733143	**Good identification**					**26733143**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26733160	**Good identification**					**26733160**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26733161	**Good identification**					**26733161**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26733162	**Good identification**					**26733162**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26733163	**Good identification**					**26733163**
Actinomyces naeslundii		100.0	0.51	GLU 99		
26737102	**Acceptable identification**					**26737102**
Actinomyces israelii		93.13	0.37	GLU 99		
Actinomyces naeslundii		6.87	0.27	CEL 93		
26737502	**Acceptable identification**					**26737502**
Actinomyces israelii		92.97	0.35	GLU 99		
Bifidobacterium adolescentis		7.03	0.10	CEL 93		
26743100	**Good identification**					**26743100**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26743101	**Good identification**					**26743101**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26743102	**Good identification**					**26743102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	GLU 99		
26743103	**Good identification**					**26743103**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26743120	**Good identification**					**26743120**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26743121	**Good identification**					**26743121**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26743122	**Good identification**					**26743122**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26743123	**Good identification**					**26743123**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26750002	**Good identification**					**26750002**
Actinomyces viscosus		100.0	0.42	GLU 99		
26750102	**Good identification**					**26750102**
Actinomyces viscosus		99.60	0.46	GLU 99		
Actinomyces naeslundii		0.40	0.29			
26750103	**Good identification**					**26750103**
Actinomyces viscosus		98.21	0.38	GLU 99		
Actinomyces naeslundii		1.79	0.32			
26751000	**Acceptable identification**					**26751000**
Actinomyces naeslundii		90.74	0.45	GLU 99		
Actinomyces viscosus		9.26	0.04			
26751001	**Good identification**					**26751001**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26751002	**Acceptable identification**					**26751002**
Actinomyces viscosus		90.99	0.38	GLU 99		
Actinomyces naeslundii		9.01	0.45			
26751020	**Good identification**					**26751020**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26751021	**Good identification**					**26751021**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26751022	**Low discrimination**					**26751022**
					GRA	
Actinomyces naeslundii		86.20	0.45	GLU 99	99	
Actinomyces viscosus		8.80	0.04		99	
Prevotella oralis		5.00	-.02		1	
26751023	**Good identification**					**26751023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	GLU 99		
26751041	**Good identification**					**26751041**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26751043	**Good identification**					**26751043**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26751061	**Good identification**					**26751061**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26751063	**Good identification**					**26751063**
Actinomyces naeslundii		100.0	0.35	GLU 99		
26751100	**Acceptable identification**					**26751100**
Actinomyces naeslundii		94.01	0.53	GLU 99		
Actinomyces viscosus		5.99	0.08			
26751101	**Good identification**					**26751101**
Actinomyces naeslundii		98.60	0.56	GLU 99		
Actinomyces viscosus		1.40	0.00			
26751102	**Low discrimination**					**26751102**
Actinomyces viscosus		84.87	0.43	GLU 99		
Actinomyces naeslundii		13.45	0.53			
Lactobacillus acidophilus		1.17	0.10			
26751120	**Acceptable identification**					**26751120**
					GRA	
Actinomyces naeslundii		91.67	0.53	GLU 99	99	
Clostridium perfringens		3.51	0.06		99	
Prevotella melaninogenica		3.28	-.01		1	
26751121	**Good identification**					**26751121**
Actinomyces naeslundii		100.0	0.56	GLU 99		
26751122	**Low discrimination**					**26751122**
					GRA	
Actinomyces naeslundii		67.20	0.53	GLU 99	99	
Prevotella oralis		21.46	0.19		1	
Lactobacillus acidophilus		5.85	0.10		99	
26751123	**Good identification**					**26751123**
Actinomyces naeslundii		98.60	0.56	GLU 99		
Actinomyces viscosus		1.40	0.00			
26751140	**Good identification**					**26751140**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26751141	**Good identification**					**26751141**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26751142	**Low discrimination**					**26751142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		73.47	0.40	GLU 99		
Actinomyces viscosus		26.53	0.08			
26751143	**Acceptable identification**					**26751143**
Actinomyces naeslundii		92.57	0.43	GLU 99		
Actinomyces viscosus		7.43	0.00			
26751160	**Good identification**					**26751160**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26751161	**Good identification**					**26751161**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26751162	**Good identification**					**26751162**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26751163	**Good identification**					**26751163**
Actinomyces naeslundii		100.0	0.43	GLU 99		
26752101	**Good identification**					**26752101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26752103	**Low discrimination**					**26752103**
Actinomyces naeslundii		77.78	0.37	GLU 99		
Actinomyces viscosus		22.22	0.04	GAL 96		
26752121	**Good identification**					**26752121**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26752123	**Good identification**					**26752123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26753000	**Good identification**					**26753000**
Actinomyces naeslundii		97.16	0.50	GLU 99		
Actinomyces israelii		2.84	0.15			
26753001	**Good identification**					**26753001**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26753002	**Acceptable identification**					**26753002**
Actinomyces naeslundii		91.21	0.50	GLU 99		
Actinomyces viscosus		4.80	0.04			
Actinomyces israelii		4.00	0.18			
26753003	**Good identification**					**26753003**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26753020	**Good identification**					**26753020**
Actinomyces naeslundii		100.0	0.50	GLU 99		
26753021	**Good identification**					**26753021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	GLU 99		
26753022	**Good identification**					**26753022**
Actinomyces naeslundii		100.0	0.50	GLU 99		
26753023	**Good identification**					**26753023**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26753040	**Good identification**					**26753040**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26753041	**Good identification**					**26753041**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26753042	**Good identification**					**26753042**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26753043	**Good identification**					**26753043**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26753060	**Good identification**					**26753060**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26753061	**Good identification**					**26753061**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26753062	**Good identification**					**26753062**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26753063	**Good identification**					**26753063**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26753100	**Very good identification**					**26753100**
Actinomyces naeslundii		96.06	0.58	GLU 99		
Actinomyces israelii		3.95	0.25			
26753101	**Very good identification**					**26753101**
Actinomyces naeslundii		100.0	0.61	GLU 99		
26753102	**Good identification**					**26753102**
Actinomyces naeslundii		91.37	0.58	GLU 99		
Actinomyces israelii		5.63	0.28			
Actinomyces viscosus		3.00	0.08			
26753103	**Very good identification**					**26753103**
Actinomyces naeslundii		99.28	0.61	GLU 99		
Actinomyces viscosus		0.72	0.00			
26753110	**Good identification**					**26753110**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26753111	**Good identification**					**26753111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	GLU 99		
26753112	**Low discrimination**					**26753112**
Actinomyces naeslundii		85.11	0.36	GLU 99		
Actinomyces israelii		14.89	0.14	MLZ 5		
26753113	**Good identification**					**26753113**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26753120	**Very good identification**					**26753120**
Actinomyces naeslundii		100.0	0.58	GLU 99		
26753121	**Very good identification**					**26753121**
Actinomyces naeslundii		100.0	0.61	GLU 99		
26753122	**Very good identification**					**26753122**
					SPO	
Actinomyces naeslundii		97.76	0.58	GLU 99	1	
Clostridium ramosum		2.24	0.05		92	
26753123	**Very good identification**					**26753123**
Actinomyces naeslundii		100.0	0.61	GLU 99		
26753130	**Good identification**					**26753130**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26753131	**Good identification**					**26753131**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26753132	**Good identification**					**26753132**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26753133	**Good identification**					**26753133**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26753140	**Acceptable identification**					**26753140**
Actinomyces naeslundii		91.70	0.45	GLU 99		
Actinomyces israelii		8.30	0.18			
26753141	**Good identification**					**26753141**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26753142	**Low discrimination**					**26753142**
Actinomyces naeslundii		88.05	0.45	GLU 99		
Actinomyces israelii		11.95	0.21			
26753143	**Good identification**					**26753143**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26753160	**Good identification**					**26753160**
Actinomyces naeslundii		100.0	0.45	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26753161	**Good identification**					**26753161**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26753162	**Good identification**					**26753162**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26753163	**Good identification**					**26753163**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26757002	**Good identification**					**26757002**
Actinomyces israelii		100.0	0.36	GLU 99		
26757100	**Good identification**					**26757100**
Actinomyces israelii		97.91	0.43	GLU 99		
Actinomyces naeslundii		2.09	0.24			
26757102	**Good identification**					**26757102**
Actinomyces israelii		98.59	0.46	GLU 99		
Actinomyces naeslundii		1.41	0.24			
26757140	**Good identification**					**26757140**
Actinomyces israelii		100.0	0.36	GLU 99		
26757142	**Good identification**					**26757142**
Actinomyces israelii		100.0	0.39	GLU 99		
26757500	**Good identification**					**26757500**
Actinomyces israelii		100.0	0.42	GLU 99		
26757502	**Acceptable identification**					**26757502**
Actinomyces israelii		93.61	0.45	GLU 99		
Bifidobacterium adolescentis		6.39	0.18			
26757542	**Good identification**					**26757542**
Actinomyces israelii		100.0	0.38	GLU 99		
26761100	**Good identification**					**26761100**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26761101	**Good identification**					**26761101**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26761102	**Low discrimination**					**26761102**
Actinomyces naeslundii		80.35	0.36	GLU 99		
Actinomyces viscosus		19.65	0.01	FRU 95		
26761103	**Good identification**					**26761103**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26761120	**Good identification**					**26761120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	GLU 99		
26761121	**Good identification**					**26761121**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26761122	**Good identification**					**26761122**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26761123	**Good identification**					**26761123**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26763001	**Good identification**					**26763001**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26763003	**Good identification**					**26763003**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26763021	**Good identification**					**26763021**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26763023	**Good identification**					**26763023**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26763100	**Good identification**					**26763100**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26763101	**Good identification**					**26763101**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26763102	**Good identification**					**26763102**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26763103	**Good identification**					**26763103**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26763120	**Good identification**					**26763120**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26763121	**Good identification**					**26763121**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26763122	**Good identification**					**26763122**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26763123	**Good identification**					**26763123**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26770101	**Good identification**					**26770101**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26770121	**Good identification**					**26770121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GLU 99		
26770123	**Good identification**					**26770123**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26771000	**Good identification**					**26771000**
Actinomyces naeslundii		100.0	0.50	GLU 99		
26771001	**Good identification**					**26771001**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26771002	**Low discrimination**					**26771002**
Actinomyces naeslundii		68.59	0.50	GLU 99		
Actinomyces viscosus		26.87	0.19			
Lactobacillus acidophilus		2.55	0.01			
26771003	**Acceptable identification**					**26771003**
Actinomyces naeslundii		91.99	0.53	GLU 99		
Actinomyces viscosus		8.01	0.11			
26771020	**Good identification**					**26771020**
Actinomyces naeslundii		100.0	0.50	GLU 99		
26771021	**Good identification**					**26771021**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26771022	**Good identification**					**26771022**
Actinomyces naeslundii		96.42	0.50	GLU 99		
Lactobacillus acidophilus		3.58	0.01			
26771023	**Good identification**					**26771023**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26771040	**Good identification**					**26771040**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26771041	**Good identification**					**26771041**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26771042	**Good identification**					**26771042**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26771043	**Good identification**					**26771043**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26771060	**Good identification**					**26771060**
Actinomyces naeslundii		100.0	0.37	GLU 99		
26771061	**Good identification**					**26771061**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26771062	**Good identification**					**26771062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GLU 99		
26771063	**Good identification**					**26771063**
Actinomyces naeslundii		100.0	0.40	GLU 99		
26771100	**Very good identification**					**26771100**
Actinomyces naeslundii		97.35	0.58	GLU 99		
Actinomyces israelii		2.65	0.22			
26771101	**Very good identification**					**26771101**
Actinomyces naeslundii		100.0	0.61	GLU 99		
26771102	**Low discrimination**					**26771102**
Actinomyces naeslundii		69.65	0.58	GLU 99		
Actinomyces viscosus		17.04	0.23			
Lactobacillus acidophilus		10.46	0.19			
26771103	**Good identification**					**26771103**
Actinomyces naeslundii		94.84	0.61	GLU 99		
Actinomyces viscosus		5.16	0.15			
26771110	**Good identification**					**26771110**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26771111	**Good identification**					**26771111**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26771113	**Good identification**					**26771113**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26771120	**Very good identification**					**26771120**
					SPO	
Actinomyces naeslundii		95.02	0.58	GLU 99	1	
Clostridium perfringens		3.48	0.11		84	
Clostridium ramosum		1.50	0.02		92	
26771121	**Very good identification**					**26771121**
Actinomyces naeslundii		100.0	0.61	GLU 99		
26771122	**Low discrimination**					**26771122**
					SPO	
Actinomyces naeslundii		80.57	0.58	GLU 99	1	
Lactobacillus acidophilus		12.10	0.19		1	
Clostridium ramosum		5.10	0.12		92	
26771123	**Very good identification**					**26771123**
Actinomyces naeslundii		100.0	0.61	GLU 99		
26771130	**Good identification**					**26771130**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26771131	**Good identification**					**26771131**
Actinomyces naeslundii		100.0	0.39	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
26771132	**Low discrimination**					**26771132**
					GRA	
Actinomyces naeslundii		81.37	0.36	GLU 99	99	
Leptotrichia buccalis		18.63	-.09	MLZ 5	1	
26771133	**Good identification**					**26771133**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26771140	**Acceptable identification**					**26771140**
Actinomyces naeslundii		94.33	0.45	GLU 99		
Actinomyces israelii		5.67	0.15			
26771141	**Good identification**					**26771141**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26771142	**Low discrimination**					**26771142**
Actinomyces naeslundii		83.67	0.45	GLU 99		
Lactobacillus acidophilus		8.80	0.04			
Actinomyces israelii		7.54	0.18			
26771143	**Good identification**					**26771143**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26771160	**Good identification**					**26771160**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26771161	**Good identification**					**26771161**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26771162	**Acceptable identification**					**26771162**
Actinomyces naeslundii		90.49	0.45	GLU 99		
Lactobacillus acidophilus		9.51	0.04			
26771163	**Good identification**					**26771163**
Actinomyces naeslundii		100.0	0.48	GLU 99		
26772100	**Good identification**					**26772100**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26772101	**Good identification**					**26772101**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26772102	**Good identification**					**26772102**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26772103	**Good identification**					**26772103**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26772120	**Good identification**					**26772120**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26772121	**Good identification**					**26772121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	GLU 99		
26772122	**Good identification**					**26772122**
Actinomyces naeslundii		100.0	0.39	GLU 99		
26772123	**Good identification**					**26772123**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26773000	**Acceptable identification**					**26773000**
Actinomyces naeslundii		94.64	0.55	GLU 99		
Actinomyces israelii		5.36	0.25			
26773001	**Very good identification**					**26773001**
Actinomyces naeslundii		100.0	0.58	GLU 99		
26773002	**Acceptable identification**					**26773002**
Actinomyces naeslundii		92.17	0.55	GLU 99		
Actinomyces israelii		7.83	0.28			
26773003	**Very good identification**					**26773003**
Actinomyces naeslundii		100.0	0.58	GLU 99		
26773011	**Good identification**					**26773011**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26773013	**Good identification**					**26773013**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26773020	**Good identification**					**26773020**
Actinomyces naeslundii		100.0	0.55	GLU 99		
26773021	**Very good identification**					**26773021**
Actinomyces naeslundii		100.0	0.58	GLU 99		
26773022	**Good identification**					**26773022**
Actinomyces naeslundii		100.0	0.55	GLU 99		
26773023	**Very good identification**					**26773023**
Actinomyces naeslundii		100.0	0.58	GLU 99		
26773031	**Good identification**					**26773031**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26773033	**Good identification**					**26773033**
Actinomyces naeslundii		100.0	0.36	GLU 99		
26773040	**Low discrimination**					**26773040**
Actinomyces naeslundii		88.91	0.42	GLU 99		
Actinomyces israelii		11.09	0.18			
26773041	**Good identification**					**26773041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	GLU 99		
26773042	**Low discrimination**					**26773042**
Actinomyces naeslundii		84.24	0.42	GLU 99		
Actinomyces israelii		15.77	0.21			
26773043	**Good identification**					**26773043**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26773060	**Good identification**					**26773060**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26773061	**Good identification**					**26773061**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26773062	**Good identification**					**26773062**
Actinomyces naeslundii		100.0	0.42	GLU 99		
26773063	**Good identification**					**26773063**
Actinomyces naeslundii		100.0	0.45	GLU 99		
26773100	**Good identification**					**26773100**
Actinomyces naeslundii		92.63	0.63	GLU 99		
Actinomyces israelii		7.37	0.35			
26773101	**Very good identification**					**26773101**
Actinomyces naeslundii		100.0	0.66	GLU 99		
26773102	**Low discrimination**					**26773102**
Actinomyces naeslundii		89.34	0.63	GLU 99		
Actinomyces israelii		10.66	0.38			
26773103	**Very good identification**					**26773103**
Actinomyces naeslundii		100.0	0.66	GLU 99		
26773110	**Low discrimination**					**26773110**
Actinomyces naeslundii		81.57	0.41	GLU 99		
Actinomyces israelii		18.43	0.21	MLZ 5		
26773111	**Good identification**					**26773111**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26773112	**Low discrimination**					**26773112**
Actinomyces naeslundii		74.69	0.41	GLU 99		
Actinomyces israelii		25.31	0.24	MLZ 5		
26773113	**Good identification**					**26773113**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26773120	**Very good identification**					**26773120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.13	0.63	GLU 99	1	
Clostridium ramosum		4.39	0.15		92	
Actinomyces israelii		0.48	0.14		1	
26773121	**Very good identification**					**26773121**
Actinomyces naeslundii		100.0	0.66	GLU 99		
26773122	**Low discrimination**					**26773122**
					SPO	
Actinomyces naeslundii		83.88	0.63	GLU 99	1	
Clostridium ramosum		15.49	0.25		92	
Actinomyces israelii		0.64	0.17		1	
26773123	**Very good identification**					**26773123**
Actinomyces naeslundii		100.0	0.66	GLU 99		
26773130	**Good identification**					**26773130**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26773131	**Good identification**					**26773131**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26773132	**Good identification**					**26773132**
Actinomyces naeslundii		100.0	0.41	GLU 99		
26773133	**Good identification**					**26773133**
Actinomyces naeslundii		100.0	0.44	GLU 99		
26773140	**Low discrimination**					**26773140**
Actinomyces naeslundii		85.08	0.50	GLU 99		
Actinomyces israelii		14.92	0.28			
26773141	**Good identification**					**26773141**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26773142	**Low discrimination**					**26773142**
Actinomyces naeslundii		79.17	0.50	GLU 99		
Actinomyces israelii		20.83	0.31			
26773143	**Good identification**					**26773143**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26773160	**Good identification**					**26773160**
Actinomyces naeslundii		100.0	0.50	GLU 99		
26773161	**Good identification**					**26773161**
Actinomyces naeslundii		100.0	0.53	GLU 99		
26773162	**Good identification**					**26773162**
Actinomyces naeslundii		100.0	0.50	GLU 99		
26773163	**Good identification**					**26773163**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.53	GLU 99			
26773502	**Acceptable identification**						**26773502**
Actinomyces israelii		90.63	0.36	GLU 99			
Actinomyces naeslundii		9.37	0.29	XYL 92			
26775100	**Low discrimination**						**26775100**
					SPO		
Actinomyces israelii		89.66	0.40	GLU 99	1		
Clostridium butyricum		7.46	0.01		99		
Actinomyces naeslundii		2.89	0.24		1		
26775102	**Good identification**						**26775102**
					SPO		
Actinomyces israelii		95.87	0.43	GLU 99	1		
Clostridium butyricum		2.07	-.07		99		
Actinomyces naeslundii		2.06	0.24		1		
26775142	**Good identification**						**26775142**
Actinomyces israelii		100.0	0.36	GLU 99			
26775500	**Low discrimination**						**26775500**
					GRA	SPO	
Actinomyces israelii		41.43	0.39	GLU 99	99	1	
Clostridium butyricum		37.90	0.17		99	99	
Mitsuokella multacida		20.68	0.21		1	1	
26775502	**Low discrimination**						**26775502**
					SPO		
Actinomyces israelii		67.98	0.42	GLU 99	1		
Clostridium butyricum		16.12	0.10		99		
Clostridium clostridioforme		7.69	0.16		66		
26775712	**Good identification**						**26775712**
Clostridium clostridioforme		99.40	0.42	NIT 1			
Mitsuokella multacida		0.60	0.01	GLU 88			
26775732	**Good identification**						**26775732**
Clostridium clostridioforme		99.13	0.42	NIT 1			
Parabacteroides distasonis		0.88	0.13	GLU 88			
26777000	**Good identification**						**26777000**
Actinomyces israelii		100.0	0.43	GLU 99			
26777002	**Good identification**						**26777002**
Actinomyces israelii		100.0	0.46	GLU 99			
26777040	**Good identification**						**26777040**
Actinomyces israelii		100.0	0.36	GLU 99			
26777042	**Good identification**						**26777042**
Actinomyces israelii		100.0	0.39	GLU 99			
26777100	**Good identification**						**26777100**
Actinomyces israelii		98.91	0.53	GLU 99			
Actinomyces naeslundii		1.09	0.29				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
26777102	**Good identification**						**26777102**
Actinomyces israelii		99.27	0.56	GLU 99			
Actinomyces naeslundii		0.73	0.29				
26777110	**Good identification**						**26777110**
Actinomyces israelii		100.0	0.39	GLU 99			
26777112	**Good identification**						**26777112**
Actinomyces israelii		100.0	0.42	GLU 99			
26777122	**Low discrimination**						**26777122**
Actinomyces israelii		89.66	0.36	GLU 99			
Actinomyces naeslundii		10.34	0.29	NAG 6			
26777140	**Good identification**						**26777140**
Actinomyces israelii		100.0	0.46	GLU 99			
26777142	**Good identification**						**26777142**
Actinomyces israelii		100.0	0.49	GLU 99			
26777302	**Good identification**						**26777302**
Actinomyces israelii		100.0	0.37	GLU 99			
26777400	**Good identification**						**26777400**
Actinomyces israelii		100.0	0.41	GLU 99			
26777402	**Good identification**						**26777402**
Actinomyces israelii		100.0	0.44	GLU 99			
26777442	**Good identification**						**26777442**
Actinomyces israelii		100.0	0.37	GLU 99			
26777500	**Low discrimination**						**26777500**
					GRA	SPO	
Actinomyces israelii		81.74	0.52	GLU 99	99	1	
Mitsuokella multacida		16.80	0.27		1	1	
Clostridium butyricum		1.47	0.01		99	99	
26777502	**Good identification**						**26777502**
					GRA		
Actinomyces israelii		96.90	0.55	GLU 99	99		
Mitsuokella multacida		1.48	0.11		1		
Bifidobacterium adolescentis		1.17	0.15		99		
26777510	**Low discrimination**						**26777510**
					GRA		
Actinomyces israelii		68.57	0.37	GLU 99	99		
Mitsuokella multacida		31.44	0.19	MLZ 13	1		
26777512	**Low discrimination**						**26777512**
Actinomyces israelii		84.79	0.40	GLU 99			
Bifidobacterium adolescentis		8.39	0.17	MLZ 13			
Bifidobacterium dentium		3.94	-.07				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
26777540	**Low discrimination**						**26777540**
					GRA		
Actinomyces israelii		65.42	0.44	GLU 99	99		
Mitsuokella multacida		34.58	0.27		1		
26777542	**Good identification**						**26777542**
					GRA		
Actinomyces israelii		96.23	0.48	GLU 99	99		
Mitsuokella multacida		3.77	0.11		1		
26777702	**Low discrimination**						**26777702**
					GRA		
Actinomyces israelii		72.69	0.35	GLU 99	99		
Mitsuokella multacida		27.31	0.15	RHA 7	1		
27000000	**Low discrimination**						**27000000**
					COC	SPO	
Gemella morbillorum		88.50	0.52	NIT 1	99	1	
Atopobium minutum		4.76	0.35		1	1	
Clostridium botulinum biovar A		2.75	0.23		1	99	
27000020	**Low discrimination**						**27000020**
					COC	SPO	
Gemella morbillorum		68.32	0.35	NIT 1	99	1	
Atopobium minutum		11.02	0.26	NAG 10	1	1	
Clostridium cadaveris		9.23	0.15		1	99	
27000400	**Good identification**						**27000400**
Gemella morbillorum		100.0	0.35	NIT 1			
27001000	**Acceptable identification**						**27001000**
					COC	CAT	
Gemella morbillorum		94.15	0.47	NIT 1	99	1	
Atopobium minutum		3.13	0.26		1	1	
Propionibacterium granulosum		1.25	0.22		1	90	
27001101	**Very good identification**						**27001101**
Clostridium chauvoei		100.0	0.57	MLT 99			
27001103	**Good identification**						**27001103**
					SPO		
Clostridium chauvoei		98.44	0.44	MLT 99	99		
Actinomyces viscosus		1.56	0.12		1		
27001121	**Good identification**						**27001121**
Clostridium chauvoei		100.0	0.48	MLT 99			
27001123	**Good identification**						**27001123**
Clostridium chauvoei		100.0	0.36	MLT 99			
27004000	**Good identification**						**27004000**
					COC		
Gemella morbillorum		96.38	0.35	NIT 1	99		
Actinomyces odontolyticus		3.62	0.15	XYL 10	1		
27010000	**Low discrimination**						**27010000**
					COC	CAT	
Gemella morbillorum		51.94	0.53	NIT 1	99	1	
Propionibacterium granulosum		30.10	0.56		1	90	
Actinomyces odontolyticus		4.14	0.39		1	10	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
27010002	**Low discrimination**							**27010002**
					CAT			
Actinomyces viscosus		53.38	0.41	MLT 99	76			
Propionibacterium granulosum		27.86	0.46	RAF 94	90			
Actinomyces odontolyticus		9.40	0.35		10			
27010010	**Low discrimination**							**27010010**
					COC	SPO	CAT	
Propionibacterium granulosum		84.41	0.41	NIT 1	1	1	90	
Gemella morbillorum		10.79	0.19	MLZ 12	99	1	1	
Clostridium difficile		4.80	0.07		1	99	1	
27010020	**Low discrimination**							**27010020**
					COC	CAT		
Propionibacterium acnes		79.67	0.52	IND 87	1	95		
Propionibacterium granulosum		9.95	0.46	SUC 1	1	90		
Gemella morbillorum		7.63	0.37		99	1		
27010022	**Low discrimination**							**27010022**
					CAT			
Propionibacterium granulosum		52.61	0.35	NIT 1	90			
Propionibacterium acnes		17.03	0.18		95			
Atopobium minutum		13.11	0.23		1			
27010060	**Acceptable identification**							**27010060**
					SPO	CAT		
Propionibacterium acnes		91.80	0.40	IND 87	1	95		
Clostridium botulinum biovar B		7.27	0.18	SUC 1	99	1		
Propionibacterium granulosum		0.53	0.11		1	90		
27010100	**Low discrimination**							**27010100**
					GRA			
Fusobacterium mortiferum		71.58	0.50	NIT 1	1			
Actinomyces odontolyticus		10.72	0.30		99			
Actinomyces meyeri		6.51	0.23		99			
27010102	**Acceptable identification**							**27010102**
					GRA			
Actinomyces viscosus		91.89	0.45	MLT 99	99			
Fusobacterium mortiferum		3.48	0.34	RAF 94	1			
Actinomyces odontolyticus		2.88	0.27		99			
27010103	**Acceptable identification**							**27010103**
					SPO			
Actinomyces viscosus		94.92	0.37	MLT 99	1			
Clostridium chauvoei		4.57	0.16	RAF 94	99			
Propionibacterium propionicum		0.51	0.12		1			
27010200	**Low discrimination**							**27010200**
					COC	CAT		
Actinomyces odontolyticus		84.67	0.44	MNS 9	1	10		
Propionibacterium granulosum		9.80	0.30	MLT 99	1	90		
Gemella morbillorum		5.53	0.19		99	1		
27010202	**Acceptable identification**							**27010202**
					CAT			
Actinomyces odontolyticus		92.87	0.40	MNS 9	10			
Propionibacterium granulosum		4.39	0.20	MLT 99	90			
Actinomyces viscosus		2.74	0.07		76			
27010300	**Good identification**							**27010300**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces odontolyticus		96.64	0.35	MNS 9	99			
Fusobacterium mortiferum		3.36	0.16	MLT 99	1			
27010400	**Acceptable identification**							**27010400**
					COC	CAT		
Gemella morbillorum		94.23	0.37	NIT 1	99	1		
Propionibacterium granulosum		4.96	0.22	ARA 10	1	90		
Actinomyces meyeri		0.81	0.02		1	1		
27011000	**Low discrimination**							**27011000**
					COC	GRA	CAT	
Propionibacterium granulosum		42.25	0.56	NIT 1	1	99	90	
Gemella morbillorum		39.26	0.48		99	99	1	
Fusobacterium mortiferum		7.88	0.47		1	1	5	
27011002	**Low discrimination**							**27011002**
					CAT			
Actinomyces viscosus		39.87	0.38	MLT 99	76			
Propionibacterium granulosum		33.96	0.46	RAF 94	90			
Lactobacillus fermentum		11.35	0.42		1			
27011010	**Acceptable identification**							**27011010**
					COC	GRA	CAT	
Propionibacterium granulosum		92.37	0.41	NIT 1	1	99	90	
Gemella morbillorum		6.36	0.14	MLZ 12	99	99	1	
Fusobacterium mortiferum		1.28	0.13		1	1	5	
27011020	**Low discrimination**							**27011020**
					CAT			
Propionibacterium acnes		75.48	0.52	IND 87	95			
Propionibacterium granulosum		9.42	0.46	SUC 1	90			
Clostridium perfringens		9.09	0.38		1			
27011060	**Acceptable identification**							**27011060**
					CAT			
Propionibacterium acnes		90.07	0.40	IND 87	95			
Clostridium perfringens		9.41	0.26	SUC 1	1			
Propionibacterium granulosum		0.52	0.11		90			
27011100	**Low discrimination**							**27011100**
					GRA	SPO		
Fusobacterium mortiferum		53.28	0.53	NIT 1	1	1		
Lactobacillus fermentum		16.69	0.46		99	1		
Clostridium chauvoei		9.47	0.29		99	99		
27011101	**Very good identification**							**27011101**
					GRA	SPO		
Clostridium chauvoei		99.71	0.63	MLT 99	99	99		
Actinomyces naeslundii		0.21	0.40		99	1		
Fusobacterium mortiferum		0.06	0.19		1	1		
27011102	**Low discrimination**							**27011102**
					GRA			
Actinomyces viscosus		65.84	0.42	MLT 99	99			
Lactobacillus fermentum		17.20	0.46	RAF 94	99			
Fusobacterium mortiferum		6.10	0.37		1			
27011103	**Low discrimination**							**27011103**
					SPO			
Clostridium chauvoei		87.59	0.50	MLT 99	99			
Actinomyces viscosus		11.27	0.34		1			
Actinomyces naeslundii		1.04	0.40		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27011120	**Acceptable identification**						**27011120**
					GRA		
Clostridium perfringens		92.15	0.53	MLT 99	99		
Lactobacillus fermentum		2.37	0.38		99		
Prevotella melaninogenica		1.54	0.15		1		
27011121	**Good identification**						**27011121**
					SPO		
Clostridium chauvoei		98.69	0.55	MLT 99	99		
Clostridium perfringens		0.69	0.19		84		
Actinomyces naeslundii		0.62	0.40		1		
27011122	**Acceptable identification**						**27011122**
					GRA	CAT	
Clostridium perfringens		90.03	0.47	MLT 99	99	1	
Lactobacillus fermentum		4.92	0.38		99	1	
Capnocytophaga ochracea		1.92	0.15		1	90	
27011123	**Acceptable identification**						**27011123**
					SPO		
Clostridium chauvoei		94.50	0.42	MLT 99	99		
Actinomyces naeslundii		3.37	0.40		1		
Clostridium perfringens		1.76	0.13		84		
27011160	**Good identification**						**27011160**
					CAT		
Clostridium perfringens		98.16	0.40	MLT 99	1		
Propionibacterium acnes		1.29	0.06		95		
Actinomyces naeslundii		0.55	0.24		25		
27011200	**Low discrimination**						**27011200**
					COC	CAT	
Actinomyces odontolyticus		80.85	0.41	MNS 9	1	10	
Propionibacterium granulosum		14.04	0.30	MLT 99	1	90	
Gemella morbillorum		4.26	0.14		99	1	
27011202	**Acceptable identification**						**27011202**
					CAT		
Actinomyces odontolyticus		91.08	0.37	MNS 9	10		
Propionibacterium granulosum		6.45	0.20	MLT 99	90		
Actinomyces viscosus		2.47	0.03		76		
27011402	**Acceptable identification**						**27011402**
					CAT		
Lactobacillus fermentum		93.83	0.42	NIT 1	1		
Actinomyces viscosus		3.33	0.03		76		
Propionibacterium granulosum		2.84	0.11		90		
27011500	**Acceptable identification**						**27011500**
					GRA		
Lactobacillus fermentum		90.52	0.46	NIT 1	99		
Actinomyces meyeri		6.57	0.16		99		
Fusobacterium mortiferum		2.92	0.19		1		
27011502	**Good identification**						**27011502**
Lactobacillus fermentum		96.28	0.46	NIT 1			
Actinomyces viscosus		3.72	0.08				
27011520	**Low discrimination**						**27011520**
Lactobacillus fermentum		71.79	0.38	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		28.21	0.19					
27011522	**Low discrimination**							**27011522**
					GRA	CAT		
Lactobacillus fermentum		79.76	0.38	NIT 1	99	1		
Clostridium perfringens		14.75	0.13		99	1		
Capnocytophaga ochracea		5.49	0.02		1	90		
27012000	**Low discrimination**							**27012000**
					COC	CAT	GRA	
Propionibacterium granulosum		72.56	0.54	NIT 1	1	90	99	
Eubacterium limosum		17.01	0.29		1	1	99	
Megasphaera elsdenii		5.23	0.29		99	1	1	
27012002	**Low discrimination**							**27012002**
					COC	CAT	GRA	
Propionibacterium granulosum		87.69	0.44	NIT 1	1	90	99	
Propionibacterium propionicum		6.42	0.29		1	1	99	
Megasphaera elsdenii		2.81	0.12		99	1	1	
27012010	**Low discrimination**							**27012010**
					SPO	CAT		
Propionibacterium granulosum		58.98	0.39	NIT 1	1	90		
Clostridium difficile		39.99	0.24	MLZ 12	99	1		
Eubacterium limosum		1.02	-.06		1	1		
27012020	**Low discrimination**							**27012020**
					COC	CAT		
Propionibacterium acnes		85.42	0.47	IND 87	1	95		
Propionibacterium granulosum		14.30	0.44	SUC 1	1	90		
Gemella morbillorum		0.15	0.03		99	1		
27012060	**Good identification**							**27012060**
Propionibacterium acnes		99.24	0.36	IND 87				
Propionibacterium granulosum		0.77	0.09	SUC 1				
27012100	**Very good identification**							**27012100**
					GRA	CAT		
Propionibacterium propionicum		98.66	0.60	MLT 99	99	1		
Propionibacterium granulosum		0.83	0.20		99	90		
Fusobacterium mortiferum		0.32	0.16		1	5		
27012101	**Good identification**							**27012101**
Propionibacterium propionicum		100.0	0.44	MLT 99				
27012102	**Very good identification**							**27012102**
					CAT			
Propionibacterium propionicum		99.24	0.63	MLT 99	1			
Actinomyces viscosus		0.62	0.11		76			
Propionibacterium granulosum		0.14	0.09		90			
27012103	**Good identification**							**27012103**
Propionibacterium propionicum		98.15	0.47	MLT 99				
Actinomyces viscosus		1.85	0.03	URE 10				
27012140	**Very good identification**							**27012140**
Propionibacterium propionicum		100.0	0.60	MLT 99				
27012141	**Good identification**							**27012141**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		100.0	0.44	MLT 99			
27012142	**Very good identification**						**27012142**
Propionibacterium propionicum		100.0	0.63	MLT 99			
27012143	**Good identification**						**27012143**
Propionibacterium propionicum		100.0	0.47	MLT 99			
27013000	**Acceptable identification**						**27013000**
					COC	CAT	
Propionibacterium granulosum		94.85	0.54	NIT 1	1	90	
Propionibacterium acnes		1.43	0.13		1	95	
Gemella morbillorum		1.18	0.14		99	1	
27013001	**Low discrimination**						**27013001**
Actinomyces naeslundii		58.97	0.37	MLT 94			
Propionibacterium granulosum		41.03	0.20	RAF 92			
27013002	**Low discrimination**						**27013002**
					CAT		
Propionibacterium granulosum		88.68	0.44	NIT 1	90		
Propionibacterium propionicum		6.49	0.29		1		
Actinomyces naeslundii		3.43	0.34		25		
27013003	**Low discrimination**						**27013003**
					CAT		
Actinomyces naeslundii		76.10	0.37	MLT 94	25		
Propionibacterium granulosum		13.24	0.09	RAF 92	90		
Propionibacterium propionicum		10.66	0.12		1		
27013010	**Good identification**						**27013010**
Propionibacterium granulosum		100.0	0.39	NIT 1			
27013020	**Low discrimination**						**27013020**
Propionibacterium acnes		85.03	0.47	IND 87			
Propionibacterium granulosum		14.24	0.44	SUC 1			
Actinomyces naeslundii		0.55	0.34				
27013023	**Good identification**						**27013023**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27013060	**Good identification**						**27013060**
Propionibacterium acnes		99.24	0.36	IND 87			
Propionibacterium granulosum		0.77	0.09	SUC 1			
27013100	**Very good identification**						**27013100**
					CAT		
Propionibacterium propionicum		96.53	0.60	MLT 99	1		
Actinomyces naeslundii		2.20	0.42		25		
Propionibacterium granulosum		0.81	0.20		90		
27013101	**Low discrimination**						**27013101**
					SPO		
Propionibacterium propionicum		48.26	0.44	MLT 99	1		
Clostridium chauvoei		36.88	0.29	URE 10	99		
Actinomyces naeslundii		14.86	0.45		1		
27013102	**Very good identification**						**27013102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		98.00	0.63	MLT 99		
Actinomyces naeslundii		1.49	0.42			
Actinomyces viscosus		0.38	0.08			
27013103	**Low discrimination**					**27013103**
					SPO	
Propionibacterium propionicum		76.53	0.47	MLT 99	1	
Actinomyces naeslundii		15.71	0.45	URE 10	1	
Clostridium chauvoei		6.88	0.16		99	
27013120	**Low discrimination**					**27013120**
					CAT	
Actinomyces naeslundii		34.00	0.42	MLT 94	25	
Clostridium perfringens		29.22	0.19	RAF 92	1	
Propionibacterium acnes		18.61	0.13		95	
27013121	**Low discrimination**					**27013121**
					SPO	
Actinomyces naeslundii		54.72	0.45	MLT 94	1	
Clostridium chauvoei		45.28	0.21	RAF 92	99	
27013122	**Low discrimination**					**27013122**
Actinomyces naeslundii		48.34	0.42	MLT 94		
Propionibacterium propionicum		32.11	0.29	RAF 92		
Clostridium perfringens		19.55	0.13			
27013123	**Low discrimination**					**27013123**
					SPO	
Actinomyces naeslundii		83.67	0.45	MLT 94	1	
Clostridium chauvoei		12.22	0.08	RAF 92	99	
Propionibacterium propionicum		4.12	0.12		1	
27013140	**Very good identification**					**27013140**
Propionibacterium propionicum		99.60	0.60	MLT 99		
Actinomyces naeslundii		0.40	0.29			
27013141	**Acceptable identification**					**27013141**
Propionibacterium propionicum		94.85	0.44	MLT 99		
Actinomyces naeslundii		5.15	0.32	URE 10		
27013142	**Very good identification**					**27013142**
Propionibacterium propionicum		99.73	0.63	MLT 99		
Actinomyces naeslundii		0.27	0.29			
27013143	**Good identification**					**27013143**
Propionibacterium propionicum		96.51	0.47	MLT 99		
Actinomyces naeslundii		3.50	0.32	URE 10		
27014000	**Low discrimination**					**27014000**
					COC	
Gemella morbillorum		64.57	0.37	NIT 1	99	
Actinomyces odontolyticus		22.83	0.34	XYL 10	1	
Actinomyces meyeri		4.31	0.18		1	
27014200	**Good identification**					**27014200**
					COC	
Actinomyces odontolyticus		98.55	0.38	MNS 9	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		1.45	0.03	MLT 99	99			
27015002	**Low discrimination**							**27015002**
Lactobacillus fermentum		58.54	0.38	NIT 1				
Actinomyces odontolyticus		33.05	0.27					
Actinomyces viscosus		3.54	0.03					
27015100	**Low discrimination**							**27015100**
					GRA			
Lactobacillus fermentum		39.27	0.42	NIT 1	99			
Actinomyces meyeri		37.67	0.31		99			
Fusobacterium mortiferum		16.07	0.34		1			
27015102	**Low discrimination**							**27015102**
Lactobacillus fermentum		79.99	0.42	NIT 1				
Actinomyces odontolyticus		8.74	0.18					
Actinomyces viscosus		5.27	0.08					
27015200	**Good identification**							**27015200**
Actinomyces odontolyticus		100.0	0.35	MNS 9				
27015400	**Low discrimination**							**27015400**
					COC			
Lactobacillus fermentum		75.10	0.38	NIT 1	1			
Gemella morbillorum		17.56	0.16		99			
Actinomyces meyeri		7.35	0.10		1			
27015402	**Good identification**							**27015402**
Lactobacillus fermentum		100.0	0.38	NIT 1				
27015500	**Low discrimination**							**27015500**
Lactobacillus fermentum		80.66	0.42	NIT 1				
Actinomyces meyeri		19.34	0.21					
27015502	**Good identification**							**27015502**
Lactobacillus fermentum		100.0	0.42	NIT 1				
27021101	**Good identification**							**27021101**
					SPO			
Clostridium chauvoei		99.55	0.40	MLT 99	99			
Actinomyces naeslundii		0.45	0.23	TRE 10	1			
27030000	**Very good identification**							**27030000**
					COC	GRA	CAT	
Propionibacterium granulosum		98.15	0.64	NIT 1	1	99	90	
Fusobacterium mortiferum		1.22	0.35		1	1	5	
Gemella morbillorum		0.57	0.19		99	99	1	
27030002	**Acceptable identification**							**27030002**
					GRA	CAT		
Propionibacterium granulosum		94.92	0.54	NIT 1	99	90		
Actinomyces viscosus		4.56	0.22		99	76		
Fusobacterium mortiferum		0.52	0.18		1	5		
27030010	**Good identification**							**27030010**
Propionibacterium granulosum		100.0	0.49	NIT 1				
27030012	**Good identification**							**27030012**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		100.0	0.39	NIT 1			
27030020	**Good identification**						**27030020**
					CAT		
Propionibacterium granulosum		96.56	0.54	NIT 1	90		
Propionibacterium acnes		2.60	0.18		95		
Clostridium perfringens		0.58	0.08		1		
27030022	**Good identification**						**27030022**
					CAT		
Propionibacterium granulosum		98.88	0.43	NIT 1	90		
Clostridium perfringens		1.12	0.03		1		
27030030	**Good identification**						**27030030**
Propionibacterium granulosum		100.0	0.39	NIT 1			
27030100	**Low discrimination**						**27030100**
					GRA	CAT	
Fusobacterium mortiferum		69.61	0.41	NIT 1	1	5	
Propionibacterium granulosum		24.34	0.30		99	90	
Propionibacterium propionicum		6.05	0.24		99	1	
27030200	**Good identification**						**27030200**
					CAT		
Propionibacterium granulosum		97.17	0.38	NIT 1	90		
Actinomyces odontolyticus		2.83	0.10	RHA 3	10		
27031000	**Very good identification**						**27031000**
					GRA	CAT	
Propionibacterium granulosum		97.21	0.64	NIT 1	99	90	
Fusobacterium mortiferum		1.81	0.38		1	5	
Clostridium perfringens		0.45	0.17		99	1	
27031002	**Acceptable identification**						**27031002**
Propionibacterium granulosum		94.69	0.54	NIT 1			
Actinomyces viscosus		2.79	0.18				
Actinomyces naeslundii		0.92	0.34				
27031003	**Low discrimination**						**27031003**
Actinomyces naeslundii		39.96	0.37	MLT 94			
Propionibacterium granulosum		27.64	0.20	RAF 92			
Actinomyces viscosus		26.87	0.10				
27031010	**Good identification**						**27031010**
Propionibacterium granulosum		100.0	0.49	NIT 1			
27031012	**Good identification**						**27031012**
Propionibacterium granulosum		100.0	0.39	NIT 1			
27031020	**Low discrimination**						**27031020**
					CAT		
Propionibacterium granulosum		61.21	0.54	NIT 1	90		
Clostridium perfringens		36.54	0.43		1		
Propionibacterium acnes		1.65	0.18		95		
27031021	**Low discrimination**						**27031021**
					CAT		
Actinomyces naeslundii		42.26	0.37	MLT 94	25		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		29.23	0.20	RAF 92	90		
Clostridium perfringens		17.45	0.08		1		
27031022	**Low discrimination**						**27031022**
					CAT		
Clostridium perfringens		51.96	0.37	MLT 99	1		
Propionibacterium granulosum		46.24	0.43	LAC 88	90		
Actinomyces naeslundii		1.80	0.34		25		
27031023	**Low discrimination**						**27031023**
					CAT		
Actinomyces naeslundii		73.14	0.37	MLT 94	25		
Clostridium perfringens		14.21	0.03	RAF 92	1		
Propionibacterium granulosum		12.65	0.09		90		
27031030	**Good identification**						**27031030**
					CAT		
Propionibacterium granulosum		95.77	0.39	NIT 1	90		
Clostridium perfringens		4.23	0.08	MLZ 12	1		
27031100	**Low discrimination**						**27031100**
					GRA	CAT	
Fusobacterium mortiferum		42.40	0.44	NIT 1	1	5	
Clostridium perfringens		33.13	0.31		99	1	
Propionibacterium granulosum		9.89	0.30		99	90	
27031101	**Good identification**						**27031101**
					SPO		
Clostridium chauvoei		96.47	0.47	MLT 99	99		
Actinomyces naeslundii		3.53	0.45	TRE 10	1		
27031120	**Very good identification**						**27031120**
					CAT		
Clostridium perfringens		99.06	0.57	MLT 99	1		
Actinomyces naeslundii		0.63	0.42		25		
Propionibacterium granulosum		0.23	0.20		90		
27031121	**Low discrimination**						**27031121**
					SPO		
Clostridium chauvoei		81.51	0.38	MLT 99	99		
Clostridium perfringens		9.53	0.23	TRE 10	84		
Actinomyces naeslundii		8.96	0.45		1		
27031122	**Good identification**						**27031122**
					CAT		
Clostridium perfringens		98.55	0.52	MLT 99	1		
Actinomyces naeslundii		1.33	0.42		25		
Propionibacterium granulosum		0.12	0.09		90		
27031160	**Good identification**						**27031160**
Clostridium perfringens		99.42	0.45	MLT 99			
Actinomyces naeslundii		0.58	0.29				
27031162	**Good identification**						**27031162**
Clostridium perfringens		98.77	0.39	MLT 99			
Actinomyces naeslundii		1.23	0.29				
27031200	**Good identification**						**27031200**
					CAT		
Propionibacterium granulosum		98.10	0.38	NIT 1	90		
Actinomyces odontolyticus		1.90	0.07	RHA 3	10		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27032000	**Very good identification**						**27032000**
					CAT		
Propionibacterium granulosum		99.59	0.62	NIT 1	90		
Propionibacterium propionicum		0.33	0.24		1		
Eubacterium limosum		0.08	-.06		1		
27032002	**Good identification**						**27032002**
					CAT		
Propionibacterium granulosum		98.04	0.52	NIT 1	90		
Propionibacterium propionicum		1.96	0.27		1		
27032010	**Good identification**						**27032010**
					SPO	CAT	
Propionibacterium granulosum		98.33	0.47	NIT 1	1	90	
Clostridium difficile		1.67	0.04	MLZ 12	99	1	
27032012	**Good identification**						**27032012**
Propionibacterium granulosum		100.0	0.37	NIT 1			
27032020	**Good identification**						**27032020**
Propionibacterium granulosum		98.03	0.52	NIT 1			
Propionibacterium acnes		1.97	0.13				
27032022	**Good identification**						**27032022**
Propionibacterium granulosum		100.0	0.41	NIT 1			
27032030	**Good identification**						**27032030**
Propionibacterium granulosum		100.0	0.37	NIT 1			
27032100	**Very good identification**						**27032100**
					CAT		
Propionibacterium propionicum		96.81	0.58	MLT 99	1		
Propionibacterium granulosum		2.97	0.28		90		
Actinomyces naeslundii		0.22	0.23		25		
27032101	**Good identification**						**27032101**
Propionibacterium propionicum		97.05	0.42	MLT 99			
Actinomyces naeslundii		2.95	0.26	URE 10			
27032102	**Very good identification**						**27032102**
					CAT		
Propionibacterium propionicum		99.34	0.61	MLT 99	1		
Propionibacterium granulosum		0.51	0.18		90		
Actinomyces naeslundii		0.15	0.23		25		
27032103	**Good identification**						**27032103**
Propionibacterium propionicum		98.01	0.45	MLT 99			
Actinomyces naeslundii		1.99	0.26	URE 10			
27032140	**Very good identification**						**27032140**
Propionibacterium propionicum		100.0	0.58	MLT 99			
27032141	**Good identification**						**27032141**
Propionibacterium propionicum		100.0	0.42	MLT 99			
27032142	**Very good identification**						**27032142**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.61	MLT 99		
27032143	**Good identification**					**27032143**
Propionibacterium propionicum		100.0	0.45	MLT 99		
27032200	**Good identification**					**27032200**
Propionibacterium granulosum		100.0	0.36	NIT 1		
27033000	**Very good identification**					**27033000**
					CAT	
Propionibacterium granulosum		99.05	0.62	NIT 1	90	
Actinomyces naeslundii		0.62	0.39		25	
Propionibacterium propionicum		0.33	0.24		1	
27033002	**Good identification**					**27033002**
					CAT	
Propionibacterium granulosum		95.69	0.52	NIT 1	90	
Actinomyces naeslundii		2.40	0.39		25	
Propionibacterium propionicum		1.91	0.27		1	
27033003	**Low discrimination**					**27033003**
Actinomyces naeslundii		78.81	0.42	MLT 94		
Propionibacterium granulosum		21.19	0.18	RAF 92		
27033010	**Good identification**					**27033010**
Propionibacterium granulosum		100.0	0.47	NIT 1		
27033012	**Good identification**					**27033012**
Propionibacterium granulosum		100.0	0.37	NIT 1		
27033020	**Acceptable identification**					**27033020**
Propionibacterium granulosum		94.95	0.52	NIT 1		
Actinomyces naeslundii		2.38	0.39			
Propionibacterium acnes		1.91	0.13			
27033021	**Low discrimination**					**27033021**
Actinomyces naeslundii		78.81	0.42	MLT 94		
Propionibacterium granulosum		21.19	0.18	RAF 92		
27033022	**Low discrimination**					**27033022**
					CAT	
Propionibacterium granulosum		89.67	0.41	NIT 1	90	
Actinomyces naeslundii		8.98	0.39		25	
Clostridium perfringens		1.35	0.03		1	
27033023	**Good identification**					**27033023**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27033030	**Good identification**					**27033030**
Propionibacterium granulosum		100.0	0.37	NIT 1		
27033100	**Good identification**					**27033100**
					CAT	
Propionibacterium propionicum		92.19	0.58	MLT 99	1	
Actinomyces naeslundii		4.99	0.47		25	
Propionibacterium granulosum		2.83	0.28		90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27033101	**Low discrimination**					**27033101**
					SPO	
Propionibacterium propionicum		54.52	0.42	MLT 99	1	
Actinomyces naeslundii		39.82	0.50	URE 10	1	
Clostridium chauvoei		5.66	0.12		99	
27033102	**Very good identification**					**27033102**
					CAT	
Propionibacterium propionicum		96.04	0.61	MLT 99	1	
Actinomyces naeslundii		3.47	0.47		25	
Propionibacterium granulosum		0.49	0.18		90	
27033103	**Low discrimination**					**27033103**
					SPO	
Propionibacterium propionicum		66.70	0.45	MLT 99	1	
Actinomyces naeslundii		32.48	0.50	URE 10	1	
Clostridium chauvoei		0.82	-.01		99	
27033120	**Low discrimination**					**27033120**
Actinomyces naeslundii		46.50	0.47	MLT 94		
Clostridium perfringens		38.23	0.23	RAF 92		
Propionibacterium propionicum		8.68	0.24			
27033121	**Good identification**					**27033121**
					SPO	
Actinomyces naeslundii		95.48	0.50	MLT 94	1	
Clostridium chauvoei		4.52	0.04	RAF 92	99	
27033122	**Low discrimination**					**27033122**
Actinomyces naeslundii		59.99	0.47	MLT 94		
Clostridium perfringens		23.21	0.18	RAF 92		
Propionibacterium propionicum		16.80	0.27			
27033123	**Good identification**					**27033123**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27033140	**Very good identification**					**27033140**
Propionibacterium propionicum		99.05	0.58	MLT 99		
Actinomyces naeslundii		0.95	0.34			
27033141	**Low discrimination**					**27033141**
Propionibacterium propionicum		88.58	0.42	MLT 99		
Actinomyces naeslundii		11.42	0.37	URE 10		
27033142	**Very good identification**					**27033142**
Propionibacterium propionicum		99.37	0.61	MLT 99		
Actinomyces naeslundii		0.63	0.34			
27033143	**Acceptable identification**					**27033143**
Propionibacterium propionicum		92.09	0.45	MLT 99		
Actinomyces naeslundii		7.91	0.37	URE 10		
27033161	**Good identification**					**27033161**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27033163	**Good identification**					**27033163**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		100.0	0.37	MLT 94				
27033200	**Good identification**							**27033200**
Propionibacterium granulosum		100.0	0.36	NIT 1				
27050000	**Low discrimination**							**27050000**
					COC	GRA	CAT	
Fusobacterium mortiferum		60.08	0.36	NIT 1	1	1	5	
Gemella morbillorum		25.27	0.19		99	99	1	
Propionibacterium granulosum		14.65	0.22		1	99	90	
27050100	**Good identification**							**27050100**
					GRA			
Fusobacterium mortiferum		97.69	0.42	NIT 1	1			
Actinomyces viscosus		2.31	0.01		99			
27050102	**Good identification**							**27050102**
					GRA			
Actinomyces viscosus		95.47	0.36	MLT 99	99			
Fusobacterium mortiferum		4.53	0.25	RAF 94	1			
27051000	**Low discrimination**							**27051000**
					COC	GRA	CAT	
Fusobacterium mortiferum		60.77	0.39	NIT 1	1	1	5	
Propionibacterium granulosum		9.88	0.22		1	99	90	
Gemella morbillorum		9.18	0.14		99	99	1	
27051100	**Low discrimination**							**27051100**
					GRA			
Fusobacterium mortiferum		84.33	0.45	NIT 1	1			
Actinomyces naeslundii		8.25	0.39		99			
Lactobacillus catenaformis		6.10	0.15		99			
27051123	**Low discrimination**							**27051123**
					SPO			
Actinomyces naeslundii		82.39	0.42	MLT 94	1			
Clostridium chauvoei		17.61	0.08	RAF 92	99			
27053001	**Good identification**							**27053001**
Actinomyces naeslundii		100.0	0.39	MLT 94				
27053003	**Good identification**							**27053003**
Actinomyces naeslundii		100.0	0.39	MLT 94				
27053021	**Good identification**							**27053021**
Actinomyces naeslundii		100.0	0.39	MLT 94				
27053022	**Good identification**							**27053022**
Actinomyces naeslundii		100.0	0.36	MLT 94				
27053023	**Good identification**							**27053023**
Actinomyces naeslundii		100.0	0.39	MLT 94				
27053100	**Low discrimination**							**27053100**
Actinomyces naeslundii		74.96	0.44	MLT 94				
Propionibacterium propionicum		25.04	0.26	RAF 92				
27053101	**Good identification**							**27053101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.47	MLT 94			
27053102	**Low discrimination**						**27053102**
Actinomyces naeslundii		66.62	0.44	MLT 94			
Propionibacterium propionicum		33.39	0.29	RAF 92			
27053103	**Good identification**						**27053103**
Actinomyces naeslundii		96.42	0.47	MLT 94			
Propionibacterium propionicum		3.58	0.12	RAF 92			
27053120	**Good identification**						**27053120**
Actinomyces naeslundii		100.0	0.44	MLT 94			
27053121	**Good identification**						**27053121**
Actinomyces naeslundii		100.0	0.47	MLT 94			
27053122	**Good identification**						**27053122**
Actinomyces naeslundii		100.0	0.44	MLT 94			
27053123	**Good identification**						**27053123**
Actinomyces naeslundii		100.0	0.47	MLT 94			
27071001	**Good identification**						**27071001**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27071003	**Low discrimination**						**27071003**
Actinomyces naeslundii		88.12	0.39	MLT 94			
Actinomyces viscosus		11.88	0.00	RAF 92			
27071021	**Good identification**						**27071021**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27071022	**Low discrimination**						**27071022**
Actinomyces naeslundii		39.21	0.36	MLT 94			
Clostridium perfringens		35.57	0.13	RAF 92			
Lactobacillus jensenii		17.55	0.09				
27071023	**Good identification**						**27071023**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27071100	**Low discrimination**						**27071100**
					COC	GRA	
Fusobacterium mortiferum		46.41	0.36	NIT 1	1	1	
Actinomyces naeslundii		29.51	0.44		1	99	
Atopobium parvulum		19.55	0.18		99	99	
27071101	**Low discrimination**						**27071101**
					SPO		
Actinomyces naeslundii		82.78	0.47	MLT 94	1		
Clostridium chauvoei		17.22	0.12	RAF 92	99		
27071102	**Low discrimination**						**27071102**
Actinomyces naeslundii		48.37	0.44	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		18.33	0.13	RAF 92		
Lactobacillus acidophilus		15.70	0.11			
27071103	**Low discrimination**					**27071103**
					SPO	
Actinomyces naeslundii		89.21	0.47	MLT 94	1	
Actinomyces viscosus		7.51	0.05	RAF 92	1	
Clostridium chauvoei		3.28	-.01		99	
27071121	**Acceptable identification**					**27071121**
					SPO	
Actinomyces naeslundii		93.52	0.47	MLT 94	1	
Clostridium chauvoei		6.49	0.04	RAF 92	99	
27071123	**Good identification**					**27071123**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27073001	**Good identification**					**27073001**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27073003	**Good identification**					**27073003**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27073020	**Low discrimination**					**27073020**
Actinomyces naeslundii		76.68	0.41	MLT 94		
Propionibacterium granulosum		23.32	0.18	RAF 92		
27073021	**Good identification**					**27073021**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27073022	**Good identification**					**27073022**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27073023	**Good identification**					**27073023**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27073100	**Low discrimination**					**27073100**
Actinomyces naeslundii		87.66	0.49	MLT 94		
Propionibacterium propionicum		12.34	0.24	RAF 92		
27073101	**Good identification**					**27073101**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27073102	**Low discrimination**					**27073102**
Actinomyces naeslundii		82.56	0.49	MLT 94		
Propionibacterium propionicum		17.44	0.27	RAF 92		
27073103	**Good identification**					**27073103**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27073120	**Good identification**					**27073120**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27073121	**Good identification**					**27073121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.52	MLT 94			
27073122	**Good identification**						**27073122**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27073123	**Good identification**						**27073123**
Actinomyces naeslundii		100.0	0.52	MLT 94			
27073140	**Low discrimination**						**27073140**
Actinomyces naeslundii		55.62	0.36	MLT 94			
Propionibacterium propionicum		44.38	0.24	RAF 92			
27073141	**Good identification**						**27073141**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27073143	**Good identification**						**27073143**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27073160	**Good identification**						**27073160**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27073161	**Good identification**						**27073161**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27073162	**Good identification**						**27073162**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27073163	**Good identification**						**27073163**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27100000	**Low discrimination**						**27100000**
					COC	GRA	
Gemella morbillorum		83.24	0.64	NIT 1	99	99	
Actinomyces odontolyticus		12.33	0.55		1	99	
Prevotella disiens		3.29	0.32		1	1	
27100002	**Low discrimination**						**27100002**
					COC		
Actinomyces viscosus		75.49	0.53	RAF 94	1		
Actinomyces odontolyticus		21.97	0.51	FRU 95	1		
Gemella morbillorum		2.45	0.30		99		
27100003	**Good identification**						**27100003**
					SPO		
Actinomyces viscosus		98.44	0.45	RAF 94	1		
Actinomyces odontolyticus		0.87	0.17	FRU 95	1		
Clostridium chauvoei		0.69	0.10		99		
27100020	**Low discrimination**						**27100020**
					COC	SPO	
Gemella morbillorum		80.62	0.48	NIT 1	99	1	
Actinomyces odontolyticus		17.49	0.41	NAG 10	1	1	
Clostridium botulinum biovar B		0.79	0.08		1	99	
27100022	**Low discrimination**						**27100022**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		77.57	0.38	MNS 9	1		
Actinomyces viscosus		16.54	0.19	FRU 92	1		
Gemella morbillorum		5.89	0.14	NAG 14	99		
27100040	**Acceptable identification**						**27100040**
					COC	SPO	
Clostridium botulinum biovar B		94.10	0.47	NIT 1	1	99	
Gemella morbillorum		4.35	0.30		99	1	
Actinomyces odontolyticus		0.64	0.21		1	1	
27100050	**Good identification**						**27100050**
Clostridium botulinum biovar B		100.0	0.42	NIT 1			
27100060	**Good identification**						**27100060**
					COC	SPO	
Clostridium botulinum biovar B		98.75	0.42	NIT 1	1	99	
Gemella morbillorum		1.03	0.14		99	1	
Actinomyces odontolyticus		0.22	0.07		1	1	
27100070	**Good identification**						**27100070**
Clostridium botulinum biovar B		100.0	0.37	NIT 1			
27100100	**Low discrimination**						**27100100**
					COC		
Actinomyces odontolyticus		67.60	0.46	MNS 9	1		
Gemella morbillorum		14.60	0.30	FRU 92	99		
Actinomyces viscosus		8.10	0.24		1		
27100101	**Very good identification**						**27100101**
					SPO		
Clostridium chauvoei		99.43	0.57	GAL 99	99		
Actinomyces viscosus		0.45	0.15		1		
Actinomyces odontolyticus		0.11	0.12		1		
27100102	**Good identification**						**27100102**
Actinomyces viscosus		94.69	0.58	RAF 94			
Actinomyces odontolyticus		4.89	0.43	FRU 95			
Propionibacterium propionicum		0.30	0.29				
27100103	**Low discrimination**						**27100103**
					SPO		
Actinomyces viscosus		71.72	0.50	RAF 94	1		
Clostridium chauvoei		28.09	0.44	FRU 95	99		
Actinomyces odontolyticus		0.11	0.08		1		
27100121	**Good identification**						**27100121**
					GRA	SPO	
Clostridium chauvoei		99.41	0.48	GAL 99	99	99	
Prevotella bivia		0.60	-.03		1	1	
27100123	**Acceptable identification**						**27100123**
					SPO		
Clostridium chauvoei		92.82	0.36	GAL 99	99		
Actinomyces viscosus		7.18	0.15		1		
27100200	**Very good identification**						**27100200**
					COC	GRA	
Actinomyces odontolyticus		96.48	0.60	MNS 9	1	99	
Gemella morbillorum		3.39	0.30	FRU 92	99	99	
Prevotella disiens		0.13	-.03		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27100202	**Very good identification**						**27100202**
Actinomyces odontolyticus		98.24	0.56	MNS 9			
Actinomyces viscosus		1.76	0.19	FRU 92			
27100220	**Good identification**						**27100220**
					COC		
Actinomyces odontolyticus		97.66	0.46	MNS 9	1		
Gemella morbillorum		2.34	0.14	FRU 92	99		
27100222	**Good identification**						**27100222**
Actinomyces odontolyticus		100.0	0.43	MNS 9			
27100300	**Good identification**						**27100300**
Actinomyces odontolyticus		100.0	0.51	MNS 9			
27100302	**Acceptable identification**						**27100302**
Actinomyces odontolyticus		90.85	0.48	MNS 9			
Actinomyces viscosus		9.15	0.24	FRU 92			
27100320	**Acceptable identification**						**27100320**
					GRA		
Actinomyces odontolyticus		94.75	0.38	MNS 9	99		
Prevotella bivia		5.25	-.03	FRU 92	1		
27100400	**Good identification**						**27100400**
					COC	GRA	
Gemella morbillorum		98.32	0.48	NIT 1	99	99	
Actinomyces odontolyticus		1.32	0.21	ARA 10	1	99	
Prevotella disiens		0.35	-.03		1	1	
27101000	**Low discrimination**						**27101000**
					COC	SPO	
Gemella morbillorum		83.24	0.60	NIT 1	99	1	
Actinomyces odontolyticus		15.26	0.52		1	1	
Clostridium chauvoei		0.64	0.22		1	99	
27101001	**Very good identification**						**27101001**
					COC	SPO	
Clostridium chauvoei		98.17	0.57	LAC 99	1	99	
Gemella morbillorum		1.30	0.26		99	1	
Actinomyces odontolyticus		0.24	0.18		1	1	
27101002	**Low discrimination**						**27101002**
					COC		
Actinomyces viscosus		73.81	0.50	RAF 94	1		
Actinomyces odontolyticus		23.36	0.48	FRU 95	1		
Gemella morbillorum		2.10	0.26		99		
27101003	**Low discrimination**						**27101003**
					SPO		
Clostridium chauvoei		52.73	0.44	LAC 99	99		
Actinomyces viscosus		46.41	0.42		1		
Actinomyces odontolyticus		0.45	0.14		1		
27101020	**Low discrimination**						**27101020**
					COC		
Gemella morbillorum		55.29	0.43	NIT 1	99		
Clostridium perfringens		27.85	0.38	NAG 10	1		
Actinomyces odontolyticus		14.85	0.38		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27101021	**Good identification**						**27101021**
					COC	SPO	
Clostridium chauvoei		98.92	0.48	LAC 99	1	99	
Gemella morbillorum		0.44	0.09		99	1	
Actinomyces naeslundii		0.42	0.31		1	1	
27101023	**Good identification**						**27101023**
					SPO		
Clostridium chauvoei		95.17	0.36	LAC 99	99		
Actinomyces viscosus		2.54	0.07		1		
Actinomyces naeslundii		2.29	0.31		1		
27101100	**Low discrimination**						**27101100**
					SPO		
Clostridium chauvoei		89.07	0.57	URE 99	99		
Actinomyces odontolyticus		6.77	0.43		1		
Clostridium perfringens		1.48	0.27		84		
27101101	**Very good identification**						**27101101**
					SPO		
Clostridium chauvoei		99.99	0.91		99		
Actinomyces naeslundii		0.00	0.39		1		
Actinomyces viscosus		0.00	0.12		1		
27101102	**Low discrimination**						**27101102**
					SPO		
Actinomyces viscosus		77.56	0.54	RAF 94	1		
Clostridium chauvoei		16.52	0.44	FRU 95	99		
Actinomyces odontolyticus		4.36	0.40		1		
27101103	**Very good identification**						**27101103**
					SPO		
Clostridium chauvoei		98.42	0.78		99		
Actinomyces viscosus		1.56	0.46		1		
Actinomyces naeslundii		0.02	0.39		1		
27101111	**Very good identification**						**27101111**
Clostridium chauvoei		100.0	0.57	MLZ 1			
27101113	**Good identification**						**27101113**
					SPO		
Clostridium chauvoei		98.44	0.44	MLZ 1	99		
Actinomyces viscosus		1.56	0.12		1		
27101120	**Low discrimination**						**27101120**
					GRA	SPO	
Clostridium perfringens		52.66	0.53	FRU 99	99	84	
Clostridium chauvoei		32.59	0.48		99	99	
Prevotella melaninogenica		9.66	0.33		1	1	
27101121	**Very good identification**						**27101121**
					SPO		
Clostridium chauvoei		99.97	0.83		99		
Clostridium perfringens		0.02	0.19		84		
Actinomyces naeslundii		0.01	0.39		1		
27101122	**Low discrimination**						**27101122**
					GRA	SPO	
Clostridium perfringens		71.37	0.47	FRU 99	99	84	
Clostridium chauvoei		16.56	0.36		99	99	
Prevotella melaninogenica		5.30	0.21		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27101123	**Very good identification**						**27101123**
					SPO		
Clostridium chauvoei		99.84	0.70		99		
Actinomyces naeslundii		0.07	0.39		1		
Actinomyces viscosus		0.05	0.12		1		
27101131	**Good identification**						**27101131**
Clostridium chauvoei		100.0	0.48	MLZ 1			
27101133	**Good identification**						**27101133**
Clostridium chauvoei		100.0	0.36	MLZ 1			
27101141	**Very good identification**						**27101141**
					SPO		
Clostridium chauvoei		99.93	0.57	SOR 1	99		
Actinomyces naeslundii		0.07	0.26		1		
27101143	**Good identification**						**27101143**
					SPO		
Clostridium chauvoei		97.80	0.44	SOR 1	99		
Actinomyces viscosus		1.55	0.12		1		
Actinomyces naeslundii		0.39	0.26		1		
27101160	**Good identification**						**27101160**
					GRA	SPO	
Clostridium perfringens		95.50	0.40	FRU 99	99	84	
Clostridium chauvoei		3.13	0.14		99	99	
Prevotella melaninogenica		0.93	-.01		1	1	
27101161	**Good identification**						**27101161**
					SPO		
Clostridium chauvoei		99.48	0.48	SOR 1	99		
Clostridium perfringens		0.31	0.06		84		
Actinomyces naeslundii		0.21	0.26		1		
27101163	**Good identification**						**27101163**
					SPO		
Clostridium chauvoei		98.00	0.36	SOR 1	99		
Actinomyces naeslundii		1.19	0.26		1		
Clostridium perfringens		0.81	0.01		84		
27101200	**Very good identification**						**27101200**
					COC		
Actinomyces odontolyticus		97.24	0.57	MNS 9	1		
Gemella morbillorum		2.76	0.26	FRU 92	99		
27101202	**Good identification**						**27101202**
Actinomyces odontolyticus		98.38	0.53	MNS 9			
Actinomyces viscosus		1.62	0.16	FRU 92			
27101220	**Good identification**						**27101220**
					COC		
Actinomyces odontolyticus		97.17	0.43	MNS 9	1		
Gemella morbillorum		1.88	0.09	FRU 92	99		
Clostridium perfringens		0.95	0.04	NAG 14	1		
27101222	**Good identification**						**27101222**
Actinomyces odontolyticus		100.0	0.40	MNS 9			
27101300	**Acceptable identification**						**27101300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces odontolyticus		93.59	0.48	MNS 9	1	
Clostridium chauvoei		6.41	0.22	FRU 92	99	
27101301	**Very good identification**					**27101301**
					SPO	
Clostridium chauvoei		99.85	0.57	RHA 1	99	
Actinomyces odontolyticus		0.15	0.14		1	
27101302	**Low discrimination**					**27101302**
					SPO	
Actinomyces odontolyticus		89.91	0.45	MNS 9	1	
Actinomyces viscosus		8.32	0.20	FRU 92	1	
Clostridium chauvoei		1.77	0.10		99	
27101303	**Good identification**					**27101303**
					SPO	
Clostridium chauvoei		97.95	0.44	RHA 1	99	
Actinomyces viscosus		1.55	0.12		1	
Actinomyces odontolyticus		0.51	0.10		1	
27101321	**Good identification**					**27101321**
Clostridium chauvoei		100.0	0.48	RHA 1		
27101323	**Good identification**					**27101323**
Clostridium chauvoei		100.0	0.36	RHA 1		
27101400	**Good identification**					**27101400**
					COC	
Gemella morbillorum		97.85	0.43	NIT 1	99	
Actinomyces odontolyticus		1.63	0.18	ARA 10	1	
Lactobacillus fermentum		0.52	0.16		1	
27101501	**Very good identification**					**27101501**
Clostridium chauvoei		100.0	0.57	ARA 1		
27101503	**Good identification**					**27101503**
					SPO	
Clostridium chauvoei		98.44	0.44	ARA 1	99	
Actinomyces viscosus		1.56	0.12		1	
27101521	**Good identification**					**27101521**
Clostridium chauvoei		100.0	0.48	ARA 1		
27101523	**Good identification**					**27101523**
Clostridium chauvoei		100.0	0.36	ARA 1		
27102100	**Very good identification**					**27102100**
Propionibacterium propionicum		99.59	0.60	FRU 99		
Actinomyces odontolyticus		0.41	0.12			
27102101	**Low discrimination**					**27102101**
					SPO	
Propionibacterium propionicum		75.33	0.44	FRU 99	1	
Clostridium chauvoei		24.67	0.22	URE 10	99	
27102102	**Very good identification**					**27102102**
Propionibacterium propionicum		96.67	0.63	FRU 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		3.17	0.24			
Actinomyces odontolyticus		0.16	0.08			
27102103	**Low discrimination**					**27102103**
					SPO	
Propionibacterium propionicum		87.96	0.47	FRU 99	1	
Actinomyces viscosus		8.65	0.15	URE 10	1	
Clostridium chauvoei		3.39	0.10		99	
27102140	**Very good identification**					**27102140**
Propionibacterium propionicum		100.0	0.60	FRU 99		
27102141	**Good identification**					**27102141**
Propionibacterium propionicum		100.0	0.44	FRU 99		
27102142	**Very good identification**					**27102142**
Propionibacterium propionicum		100.0	0.63	FRU 99		
27102143	**Good identification**					**27102143**
Propionibacterium propionicum		100.0	0.47	FRU 99		
27103003	**Low discrimination**					**27103003**
					SPO	
Actinomyces naeslundii		41.88	0.36	RAF 92	1	
Clostridium chauvoei		27.13	0.10	FRU 95	99	
Actinomyces viscosus		23.88	0.07		1	
27103023	**Low discrimination**					**27103023**
					SPO	
Actinomyces naeslundii		82.24	0.36	RAF 92	1	
Clostridium chauvoei		17.76	0.01	FRU 95	99	
27103100	**Good identification**					**27103100**
					SPO	
Propionibacterium propionicum		94.52	0.60	FRU 99	1	
Clostridium chauvoei		3.44	0.22		99	
Actinomyces naeslundii		1.78	0.41		1	
27103101	**Very good identification**					**27103101**
					SPO	
Clostridium chauvoei		96.28	0.57	MAN 1	99	
Propionibacterium propionicum		2.97	0.44		1	
Actinomyces naeslundii		0.75	0.44		1	
27103102	**Very good identification**					**27103102**
Propionibacterium propionicum		96.34	0.63	FRU 99		
Actinomyces viscosus		1.94	0.20			
Actinomyces naeslundii		1.21	0.41			
27103103	**Low discrimination**					**27103103**
					SPO	
Clostridium chauvoei		75.62	0.44	MAN 1	99	
Propionibacterium propionicum		19.83	0.47		1	
Actinomyces naeslundii		3.36	0.44		1	
27103121	**Good identification**					**27103121**
					SPO	
Clostridium chauvoei		97.71	0.48	MAN 1	99	
Actinomyces naeslundii		2.30	0.44		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27103122	**Low discrimination**						**27103122**
Actinomyces naeslundii		41.49	0.41	RAF 92			
Propionibacterium propionicum		33.43	0.29	FRU 95			
Clostridium perfringens		20.35	0.13				
27103123	**Low discrimination**						**27103123**
					SPO		
Clostridium chauvoei		87.64	0.36	MAN 1	99		
Actinomyces naeslundii		11.67	0.44		1		
Propionibacterium propionicum		0.70	0.12		1		
27103140	**Very good identification**						**27103140**
Propionibacterium propionicum		99.67	0.60	FRU 99			
Actinomyces naeslundii		0.33	0.28				
27103141	**Low discrimination**						**27103141**
					SPO		
Propionibacterium propionicum		72.87	0.44	FRU 99	1		
Clostridium chauvoei		23.87	0.22	URE 10	99		
Actinomyces naeslundii		3.26	0.31		1		
27103142	**Very good identification**						**27103142**
Propionibacterium propionicum		99.78	0.63	FRU 99			
Actinomyces naeslundii		0.22	0.28				
27103143	**Acceptable identification**						**27103143**
					SPO		
Propionibacterium propionicum		93.60	0.47	FRU 99	1		
Clostridium chauvoei		3.61	0.10	URE 10	99		
Actinomyces naeslundii		2.80	0.31		1		
27104000	**Low discrimination**						**27104000**
					COC	GRA	
Gemella morbillorum		60.24	0.48	NIT 1	99	99	
Actinomyces odontolyticus		39.54	0.50	XYL 10	1	99	
Prevotella disiens		0.22	-.03		1	1	
27104002	**Acceptable identification**						**27104002**
					COC		
Actinomyces odontolyticus		91.28	0.46	MNS 9	1		
Actinomyces viscosus		6.43	0.19	FRU 92	1		
Gemella morbillorum		2.29	0.14		99		
27104100	**Acceptable identification**						**27104100**
					COC		
Actinomyces odontolyticus		93.88	0.41	MNS 9	1		
Gemella morbillorum		4.58	0.14	FRU 92	99		
Actinomyces meyeri		1.55	0.07		1		
27104102	**Low discrimination**						**27104102**
Actinomyces odontolyticus		71.59	0.37	MNS 9			
Actinomyces viscosus		28.41	0.24	FRU 92			
27104200	**Good identification**						**27104200**
					COC		
Actinomyces odontolyticus		99.21	0.54	MNS 9	1		
Gemella morbillorum		0.79	0.14	FRU 92	99		
27104202	**Good identification**						**27104202**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.51	MNS 9			
27104220	**Good identification**						**27104220**
Actinomyces odontolyticus		100.0	0.41	MNS 9			
27104222	**Good identification**						**27104222**
Actinomyces odontolyticus		100.0	0.37	MNS 9			
27104300	**Good identification**						**27104300**
Actinomyces odontolyticus		100.0	0.46	MNS 9			
27104302	**Good identification**						**27104302**
Actinomyces odontolyticus		100.0	0.42	MNS 9			
27105000	**Low discrimination**						**27105000**
					COC		
Gemella morbillorum		55.07	0.43	NIT 1	99		
Actinomyces odontolyticus		44.76	0.47	XYL 10	1		
Lactobacillus fermentum		0.17	0.12		1		
27105002	**Acceptable identification**						**27105002**
					COC		
Actinomyces odontolyticus		91.63	0.43	MNS 9	1		
Actinomyces viscosus		5.94	0.16	FRU 92	1		
Gemella morbillorum		1.86	0.09		99		
27105100	**Low discrimination**						**27105100**
					COC	SPO	
Actinomyces odontolyticus		73.20	0.38	MNS 9	1	1	
Clostridium chauvoei		19.76	0.22	FRU 92	1	99	
Gemella morbillorum		2.88	0.09		99	1	
27105101	**Very good identification**						**27105101**
Clostridium chauvoei		100.0	0.57	XYL 1			
27105103	**Good identification**						**27105103**
					SPO		
Clostridium chauvoei		98.44	0.44	XYL 1	99		
Actinomyces viscosus		1.56	0.12		1		
27105121	**Good identification**						**27105121**
Clostridium chauvoei		100.0	0.48	XYL 1			
27105123	**Good identification**						**27105123**
Clostridium chauvoei		100.0	0.36	XYL 1			
27105200	**Good identification**						**27105200**
					COC		
Actinomyces odontolyticus		99.36	0.51	MNS 9	1		
Gemella morbillorum		0.64	0.09	FRU 92	99		
27105202	**Good identification**						**27105202**
Actinomyces odontolyticus		100.0	0.48	MNS 9			
27105220	**Good identification**						**27105220**
Actinomyces odontolyticus		100.0	0.38	MNS 9			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27105300	**Good identification**						**27105300**
Actinomyces odontolyticus		100.0	0.43	MNS 9			
27105302	**Good identification**						**27105302**
Actinomyces odontolyticus		100.0	0.39	MNS 9			
27110000	**Low discrimination**						**27110000**
					COC	GRA	
Actinomyces odontolyticus		53.14	0.73	MNS 9	1	99	
Gemella morbillorum		38.14	0.66		99	99	
Prevotella intermedia		3.10	0.46		1	1	
27110002	**Low discrimination**						**27110002**
					CAT		
Actinomyces viscosus		84.64	0.75	RAF 94	76		
Actinomyces odontolyticus		14.91	0.69		10		
Propionibacterium granulosum		0.20	0.40		90		
27110003	**Very good identification**						**27110003**
					SPO		
Actinomyces viscosus		99.35	0.67	RAF 94	1		
Actinomyces odontolyticus		0.53	0.35		1		
Clostridium chauvoei		0.09	0.16		99		
27110010	**Low discrimination**						**27110010**
					COC	CAT	
Actinomyces odontolyticus		39.49	0.39	MNS 9	1	10	
Gemella morbillorum		28.34	0.32	MLZ 1	99	1	
Propionibacterium granulosum		17.58	0.35		1	90	
27110012	**Low discrimination**						**27110012**
					CAT		
Actinomyces viscosus		82.78	0.41	RAF 94	76		
Actinomyces odontolyticus		14.58	0.35	MLZ 1	10		
Propionibacterium granulosum		2.65	0.25		90		
27110020	**Low discrimination**						**27110020**
					COC		
Actinomyces odontolyticus		59.44	0.59	MNS 9	1		
Gemella morbillorum		29.12	0.50	NAG 14	99		
Clostridium perfringens		6.46	0.38		1		
27110022	**Low discrimination**						**27110022**
Actinomyces odontolyticus		68.20	0.56	MNS 9			
Actinomyces viscosus		24.03	0.41	NAG 14			
Clostridium perfringens		5.69	0.32				
27110040	**Very good identification**						**27110040**
					COC	SPO	
Clostridium botulinum biovar B		95.62	0.58	NIT 1	1	99	
Actinomyces odontolyticus		1.88	0.39		1	1	
Gemella morbillorum		1.35	0.32		99	1	
27110042	**Low discrimination**						**27110042**
					SPO		
Actinomyces viscosus		73.26	0.41	RAF 94	1		
Actinomyces odontolyticus		12.90	0.35	SOR 1	1		
Clostridium botulinum biovar B		10.80	0.24		99		
27110050	**Good identification**						**27110050**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium botulinum biovar B		100.0	0.52	NIT 1		
27110060	**Good identification**					**27110060**
					SPO	
Clostridium botulinum biovar B		97.63	0.52	NIT 1	99	
Clostridium perfringens		1.30	0.26		84	
Actinomyces odontolyticus		0.64	0.25		1	
27110070	**Good identification**					**27110070**
Clostridium botulinum biovar B		100.0	0.47	NIT 1		
27110100	**Low discrimination**					**27110100**
Actinomyces odontolyticus		67.35	0.64	MNS 9		
Propionibacterium propionicum		14.26	0.60			
Actinomyces viscosus		13.33	0.45			
27110101	**Good identification**					**27110101**
					SPO	
Clostridium chauvoei		94.58	0.63	GAL 99	99	
Actinomyces viscosus		3.51	0.37		1	
Propionibacterium propionicum		1.25	0.44		1	
27110102	**Very good identification**					**27110102**
Actinomyces viscosus		95.41	0.80	RAF 94		
Actinomyces odontolyticus		2.99	0.61			
Propionibacterium propionicum		1.55	0.63			
27110103	**Good identification**					**27110103**
					SPO	
Actinomyces viscosus		94.80	0.71	RAF 94	1	
Clostridium chauvoei		4.56	0.50		99	
Propionibacterium propionicum		0.51	0.47		1	
27110112	**Good identification**					**27110112**
Actinomyces viscosus		95.47	0.45	RAF 94		
Actinomyces odontolyticus		2.99	0.27	MLZ 1		
Propionibacterium propionicum		1.55	0.29			
27110113	**Acceptable identification**					**27110113**
					SPO	
Actinomyces viscosus		94.92	0.37	RAF 94	1	
Clostridium chauvoei		4.57	0.16	MLZ 1	99	
Propionibacterium propionicum		0.51	0.12		1	
27110120	**Low discrimination**					**27110120**
					GRA	
Clostridium perfringens		67.03	0.53	GAL 99	99	
Actinomyces odontolyticus		26.56	0.51		99	
Prevotella bivia		2.73	0.15		1	
27110121	**Good identification**					**27110121**
					SPO	
Clostridium chauvoei		98.52	0.55	GAL 99	99	
Clostridium perfringens		0.69	0.19		84	
Actinomyces naeslundii		0.41	0.37		1	
27110122	**Low discrimination**					**27110122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		39.39	0.45	RAF 94			
Clostridium perfringens		38.49	0.47	NAG 1			
Actinomyces odontolyticus		19.86	0.47				
27110123	**Low discrimination**						**27110123**
					SPO		
Clostridium chauvoei		59.35	0.42	GAL 99	99		
Actinomyces viscosus		37.39	0.37		1		
Actinomyces naeslundii		1.38	0.37		1		
27110140	**Low discrimination**						**27110140**
					SPO		
Propionibacterium propionicum		87.25	0.60	MAN 99	1		
Clostridium botulinum biovar B		6.76	0.24		99		
Actinomyces odontolyticus		4.16	0.30		1		
27110141	**Low discrimination**						**27110141**
					SPO		
Propionibacterium propionicum		55.23	0.44	MAN 99	1		
Clostridium chauvoei		42.20	0.29	URE 10	99		
Actinomyces viscosus		1.56	0.03		1		
27110142	**Low discrimination**						**27110142**
Propionibacterium propionicum		60.75	0.63	MAN 99			
Actinomyces viscosus		37.85	0.45				
Actinomyces odontolyticus		1.18	0.27				
27110143	**Low discrimination**						**27110143**
					SPO		
Actinomyces viscosus		62.91	0.37	RAF 94	1		
Propionibacterium propionicum		33.66	0.47	SOR 1	1		
Clostridium chauvoei		3.03	0.16		99		
27110160	**Low discrimination**						**27110160**
					SPO		
Clostridium perfringens		86.83	0.40	GAL 99	84		
Clostridium botulinum biovar B		8.97	0.18		99		
Propionibacterium propionicum		2.37	0.26		1		
27110200	**Very good identification**						**27110200**
					COC	CAT	
Actinomyces odontolyticus		99.52	0.78	MNS 9	1	10	
Gemella morbillorum		0.37	0.32		99	1	
Propionibacterium granulosum		0.05	0.24		1	90	
27110201	**Good identification**						**27110201**
Actinomyces odontolyticus		99.40	0.44	MNS 9			
Actinomyces viscosus		0.60	-.01	URE 1			
27110202	**Very good identification**						**27110202**
					CAT		
Actinomyces odontolyticus		97.11	0.74	MNS 9	10		
Actinomyces viscosus		2.87	0.41		76		
Propionibacterium granulosum		0.02	0.14		90		
27110203	**Low discrimination**						**27110203**
Actinomyces odontolyticus		50.63	0.40	MNS 9			
Actinomyces viscosus		49.37	0.33	URE 1			
27110210	**Good identification**						**27110210**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Actinomyces odontolyticus		99.30	0.44	MNS 9	10		
Propionibacterium granulosum		0.70	0.09	MLZ 1	90		
27110212	**Good identification**						**27110212**
Actinomyces odontolyticus		97.13	0.40	MNS 9			
Actinomyces viscosus		2.87	0.07	MLZ 1			
27110220	**Very good identification**						**27110220**
					COC	CAT	
Actinomyces odontolyticus		99.61	0.64	MNS 9	1	10	
Gemella morbillorum		0.25	0.15	NAG 14	99	1	
Propionibacterium granulosum		0.08	0.14		1	90	
27110222	**Very good identification**						**27110222**
Actinomyces odontolyticus		99.82	0.61	MNS 9			
Actinomyces viscosus		0.18	0.07	NAG 14			
27110240	**Low discrimination**						**27110240**
					SPO		
Actinomyces odontolyticus		79.09	0.44	MNS 9	1		
Clostridium botulinum biovar B		20.91	0.24	SOR 1	99		
27110242	**Good identification**						**27110242**
Actinomyces odontolyticus		97.13	0.40	MNS 9			
Actinomyces viscosus		2.87	0.07	SOR 1			
27110300	**Very good identification**						**27110300**
Actinomyces odontolyticus		99.79	0.69	MNS 9			
Propionibacterium propionicum		0.11	0.26				
Actinomyces viscosus		0.10	0.11				
27110301	**Low discrimination**						**27110301**
					SPO		
Actinomyces odontolyticus		51.26	0.35	MNS 9	1		
Clostridium chauvoei		47.00	0.29	URE 1	99		
Actinomyces viscosus		1.74	0.03		1		
27110302	**Low discrimination**						**27110302**
Actinomyces odontolyticus		85.54	0.66	MNS 9			
Actinomyces viscosus		14.23	0.45				
Propionibacterium propionicum		0.23	0.29				
27110303	**Low discrimination**						**27110303**
					SPO		
Actinomyces viscosus		80.93	0.37	RAF 94	1		
Actinomyces odontolyticus		14.74	0.32	RHA 1	1		
Clostridium chauvoei		3.89	0.16		99		
27110310	**Good identification**						**27110310**
Actinomyces odontolyticus		100.0	0.35	MNS 9			
27110320	**Good identification**						**27110320**
Actinomyces odontolyticus		98.70	0.56	MNS 9			
Clostridium perfringens		1.30	0.19	NAG 14			
27110322	**Good identification**						**27110322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		98.00	0.52	MNS 9			
Actinomyces viscosus		1.01	0.11	NAG 14			
Clostridium perfringens		0.99	0.13				
27110340	**Acceptable identification**						**27110340**
Actinomyces odontolyticus		90.16	0.35	MNS 9			
Propionibacterium propionicum		9.84	0.26	SOR 1			
27110400	**Low discrimination**						**27110400**
					COC	GRA	
Gemella morbillorum		86.98	0.50	NIT 1	99	99	
Actinomyces odontolyticus		11.02	0.39	ARA 10	1	99	
Prevotella intermedia		0.64	0.12		1	1	
27110402	**Low discrimination**						**27110402**
					COC		
Actinomyces viscosus		83.03	0.41	RAF 94	1		
Actinomyces odontolyticus		14.62	0.35	ARA 1	1		
Gemella morbillorum		1.90	0.15		99		
27110502	**Good identification**						**27110502**
Actinomyces viscosus		95.03	0.45	RAF 94			
Actinomyces odontolyticus		2.97	0.27	ARA 1			
Propionibacterium propionicum		1.54	0.29				
27110503	**Acceptable identification**						**27110503**
					SPO		
Actinomyces viscosus		94.92	0.37	RAF 94	1		
Clostridium chauvoei		4.57	0.16	ARA 1	99		
Propionibacterium propionicum		0.51	0.12		1		
27110600	**Good identification**						**27110600**
					COC		
Actinomyces odontolyticus		96.05	0.44	MNS 9	1		
Gemella morbillorum		3.95	0.15	ARA 1	99		
27110602	**Good identification**						**27110602**
Actinomyces odontolyticus		97.13	0.40	MNS 9			
Actinomyces viscosus		2.87	0.07	ARA 1			
27110700	**Good identification**						**27110700**
Actinomyces odontolyticus		100.0	0.35	MNS 9			
27111000	**Low discrimination**						**27111000**
					COC	GRA	
Actinomyces odontolyticus		53.22	0.70	MNS 9	1	99	
Gemella morbillorum		30.85	0.61		99	99	
Prevotella intermedia		4.66	0.46		1	1	
27111001	**Very good identification**						**27111001**
					SPO		
Clostridium chauvoei		95.59	0.63	LAC 99	99		
Actinomyces viscosus		1.22	0.29		1		
Actinomyces odontolyticus		1.15	0.36		1		
27111002	**Low discrimination**						**27111002**
Actinomyces viscosus		81.87	0.72	RAF 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		15.68	0.66				
Clostridium perfringens		0.98	0.41				
27111003	**Low discrimination**						**27111003**
					SPO		
Actinomyces viscosus		86.56	0.64	RAF 94	1		
Clostridium chauvoei		12.08	0.50		99		
Actinomyces naeslundii		0.79	0.53		1		
27111010	**Low discrimination**						**27111010**
					COC	CAT	
Actinomyces odontolyticus		39.35	0.36	MNS 9	1	10	
Propionibacterium granulosum		26.28	0.35	MLZ 1	1	90	
Gemella morbillorum		22.81	0.27		99	1	
27111012	**Low discrimination**						**27111012**
					CAT		
Actinomyces viscosus		78.04	0.38	RAF 94	76		
Actinomyces odontolyticus		14.95	0.32	MLZ 1	10		
Propionibacterium granulosum		4.07	0.25		90		
27111020	**Good identification**						**27111020**
					COC		
Clostridium perfringens		90.83	0.72	LAC 88	1		
Actinomyces odontolyticus		5.63	0.56		1		
Gemella morbillorum		2.23	0.45		99		
27111021	**Low discrimination**						**27111021**
					SPO		
Clostridium chauvoei		87.93	0.55	LAC 99	99		
Clostridium perfringens		8.30	0.38		84		
Actinomyces naeslundii		3.04	0.53		1		
27111022	**Low discrimination**						**27111022**
Clostridium perfringens		88.87	0.66	LAC 88			
Actinomyces odontolyticus		7.17	0.53				
Actinomyces viscosus		2.32	0.38				
27111023	**Low discrimination**						**27111023**
					SPO		
Clostridium chauvoei		59.33	0.42	LAC 99	99		
Clostridium perfringens		14.94	0.32		84		
Actinomyces viscosus		12.89	0.29		1		
27111030	**Low discrimination**						**27111030**
					CAT		
Clostridium perfringens		85.99	0.38	LAC 88	1		
Propionibacterium granulosum		5.46	0.25	MLZ 1	90		
Actinomyces odontolyticus		5.33	0.22		10		
27111042	**Low discrimination**						**27111042**
Actinomyces viscosus		66.31	0.38	RAF 94			
Clostridium perfringens		14.95	0.28	SOR 1			
Actinomyces odontolyticus		12.70	0.32				
27111060	**Very good identification**						**27111060**
					SPO		
Clostridium perfringens		98.47	0.60	LAC 88	84		
Clostridium botulinum biovar B		0.75	0.18		99		
Actinomyces odontolyticus		0.32	0.22		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27111062	**Good identification**						**27111062**
Clostridium perfringens		98.87	0.54	LAC 88			
Actinomyces naeslundii		0.48	0.37				
Actinomyces odontolyticus		0.42	0.19				
27111100	**Low discrimination**						**27111100**
					SPO		
Clostridium chauvoei		42.15	0.63	URE 99	99		
Clostridium perfringens		29.79	0.61		84		
Actinomyces odontolyticus		15.78	0.61		1		
27111101	**Excellent identification**						**27111101**
					SPO		
Clostridium chauvoei		99.92	0.97		99		
Actinomyces naeslundii		0.03	0.61		1		
Actinomyces viscosus		0.02	0.34		1		
27111102	**Low discrimination**						**27111102**
Actinomyces viscosus		86.87	0.76	RAF 94			
Clostridium perfringens		4.28	0.55				
Actinomyces odontolyticus		2.96	0.58				
27111103	**Low discrimination**						**27111103**
					SPO		
Clostridium chauvoei		88.33	0.84		99		
Actinomyces viscosus		11.37	0.68		1		
Actinomyces naeslundii		0.17	0.61		1		
27111111	**Very good identification**						**27111111**
					SPO		
Clostridium chauvoei		99.81	0.63	MLZ 1	99		
Actinomyces naeslundii		0.17	0.39		1		
Actinomyces viscosus		0.02	-.01		1		
27111112	**Low discrimination**						**27111112**
					GRA	CAT	
Actinomyces viscosus		83.18	0.42	RAF 94	99	76	
Clostridium perfringens		4.10	0.21	MLZ 1	99	1	
Capnocytophaga ochracea		3.45	0.17		1	90	
27111113	**Low discrimination**						**27111113**
					SPO		
Clostridium chauvoei		87.75	0.50	MLZ 1	99		
Actinomyces viscosus		11.29	0.34		1		
Actinomyces naeslundii		0.86	0.39		1		
27111120	**Very good identification**						**27111120**
					GRA	SPO	
Clostridium perfringens		96.43	0.87		99	84	
Prevotella melaninogenica		1.61	0.50		1	1	
Clostridium chauvoei		1.41	0.55		99	99	
27111121	**Very good identification**						**27111121**
					SPO		
Clostridium chauvoei		99.19	0.89		99		
Clostridium perfringens		0.69	0.53		84		
Actinomyces naeslundii		0.10	0.61		1		
27111122	**Very good identification**						**27111122**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		95.45	0.81		99	1	
Capnocytophaga ochracea		2.03	0.50		1	90	
Prevotella melaninogenica		0.64	0.37		1	1	
27111123	**Very good identification**						**27111123**
					SPO		
Clostridium chauvoei		97.20	0.76		99		
Clostridium perfringens		1.81	0.47		84		
Actinomyces naeslundii		0.55	0.61		1		
27111130	**Good identification**						**27111130**
					GRA	SPO	
Clostridium perfringens		95.13	0.53	MLZ 1	99	84	
Prevotella melaninogenica		1.59	0.15		1	1	
Clostridium chauvoei		1.39	0.21		99	99	
27111131	**Good identification**						**27111131**
					SPO		
Clostridium chauvoei		98.80	0.55	MLZ 1	99		
Clostridium perfringens		0.69	0.19		84		
Actinomyces naeslundii		0.51	0.39		1		
27111132	**Low discrimination**						**27111132**
					GRA	CAT	
Clostridium perfringens		78.94	0.47	MLZ 1	99	1	
Capnocytophaga ochracea		18.49	0.33		1	90	
Actinomyces naeslundii		0.84	0.36		99	25	
27111133	**Acceptable identification**						**27111133**
					SPO		
Clostridium chauvoei		94.67	0.42	MLZ 1	99		
Actinomyces naeslundii		2.78	0.39		1		
Clostridium perfringens		1.77	0.13		84		
27111140	**Low discrimination**						**27111140**
					SPO		
Clostridium perfringens		49.33	0.49	NAG 97	84		
Propionibacterium propionicum		43.60	0.60		1		
Clostridium chauvoei		3.70	0.29		99		
27111141	**Very good identification**						**27111141**
					SPO		
Clostridium chauvoei		97.98	0.63	SOR 1	99		
Propionibacterium propionicum		1.30	0.44		1		
Actinomyces naeslundii		0.57	0.48		1		
27111142	**Low discrimination**						**27111142**
Propionibacterium propionicum		55.55	0.63	MAN 99			
Actinomyces viscosus		21.21	0.42				
Clostridium perfringens		19.72	0.43				
27111143	**Low discrimination**						**27111143**
					SPO		
Clostridium chauvoei		78.28	0.50	SOR 1	99		
Actinomyces viscosus		10.07	0.34		1		
Propionibacterium propionicum		8.80	0.47		1		
27111160	**Very good identification**						**27111160**
					GRA		
Clostridium perfringens		99.64	0.75		99		
Actinomyces naeslundii		0.09	0.45		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		0.09	0.15		1		
27111161	**Low discrimination**						**27111161**
					SPO		
Clostridium chauvoei		87.01	0.55	SOR 1	99		
Clostridium perfringens		11.48	0.40		84		
Actinomyces naeslundii		1.51	0.48		1		
27111162	**Very good identification**						**27111162**
					GRA	CAT	
Clostridium perfringens		98.39	0.69		99	1	
Capnocytophaga ochracea		1.22	0.33		1	90	
Actinomyces naeslundii		0.19	0.45		99	25	
27111163	**Low discrimination**						**27111163**
					SPO		
Clostridium chauvoei		68.40	0.42	SOR 1	99		
Clostridium perfringens		24.06	0.35		84		
Actinomyces naeslundii		6.74	0.48		1		
27111170	**Good identification**						**27111170**
					GRA	CAT	
Clostridium perfringens		98.84	0.40	MLZ 1	99	1	
Capnocytophaga ochracea		0.71	0.01		1	90	
Actinomyces naeslundii		0.46	0.23		99	25	
27111200	**Very good identification**						**27111200**
					COC	CAT	
Actinomyces odontolyticus		99.50	0.75	MNS 9	1	10	
Gemella morbillorum		0.30	0.27		99	1	
Propionibacterium granulosum		0.08	0.24		1	90	
27111201	**Low discrimination**						**27111201**
					SPO		
Actinomyces odontolyticus		69.72	0.41	MNS 9	1		
Clostridium chauvoei		30.28	0.29	URE 1	99		
27111202	**Very good identification**						**27111202**
					CAT		
Actinomyces odontolyticus		97.27	0.71	MNS 9	10		
Actinomyces viscosus		2.64	0.38		76		
Propionibacterium granulosum		0.03	0.14		90		
27111203	**Low discrimination**						**27111203**
					SPO		
Actinomyces odontolyticus		49.47	0.37	MNS 9	1		
Actinomyces viscosus		44.34	0.29	URE 1	1		
Clostridium chauvoei		6.19	0.16		99		
27111210	**Good identification**						**27111210**
					CAT		
Actinomyces odontolyticus		98.95	0.41	MNS 9	10		
Propionibacterium granulosum		1.05	0.09	MLZ 1	90		
27111212	**Good identification**						**27111212**
Actinomyces odontolyticus		97.36	0.37	MNS 9			
Actinomyces viscosus		2.64	0.03	MLZ 1			
27111220	**Good identification**						**27111220**
					COC		
Actinomyces odontolyticus		91.97	0.61	MNS 9	1		
Clostridium perfringens		7.73	0.38	NAG 14	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		0.19	0.11		99		
27111222	**Good identification**						**27111222**
Actinomyces odontolyticus		93.79	0.58	MNS 9			
Clostridium perfringens		6.05	0.32	NAG 14			
Actinomyces viscosus		0.16	0.03				
27111240	**Good identification**						**27111240**
Actinomyces odontolyticus		100.0	0.41	MNS 9			
27111242	**Good identification**						**27111242**
Actinomyces odontolyticus		97.36	0.37	MNS 9			
Actinomyces viscosus		2.64	0.03	SOR 1			
27111300	**Very good identification**						**27111300**
					SPO		
Actinomyces odontolyticus		97.34	0.66	MNS 9	1		
Clostridium chauvoei		1.35	0.29		99		
Clostridium perfringens		0.96	0.27		84		
27111301	**Very good identification**						**27111301**
					SPO		
Clostridium chauvoei		99.22	0.63	RHA 1	99		
Actinomyces odontolyticus		0.73	0.32		1		
Actinomyces naeslundii		0.03	0.26		1		
27111302	**Low discrimination**						**27111302**
Actinomyces odontolyticus		85.41	0.63	MNS 9			
Actinomyces viscosus		13.06	0.42				
Clostridium perfringens		0.64	0.21				
27111303	**Low discrimination**						**27111303**
					SPO		
Clostridium chauvoei		86.41	0.50	RHA 1	99		
Actinomyces viscosus		11.12	0.34		1		
Actinomyces odontolyticus		2.20	0.29		1		
27111320	**Low discrimination**						**27111320**
					GRA		
Clostridium perfringens		64.68	0.53	RHA 1	99		
Actinomyces odontolyticus		33.16	0.53		99		
Prevotella melaninogenica		1.08	0.15		1		
27111321	**Good identification**						**27111321**
					SPO		
Clostridium chauvoei		98.86	0.55	RHA 1	99		
Clostridium perfringens		0.69	0.19		84		
Actinomyces odontolyticus		0.36	0.19		1		
27111322	**Low discrimination**						**27111322**
					GRA	CAT	
Clostridium perfringens		57.80	0.47	RHA 1	99	1	
Actinomyces odontolyticus		38.59	0.49		99	10	
Capnocytophaga ochracea		1.23	0.15		1	90	
27111323	**Good identification**						**27111323**
					SPO		
Clostridium chauvoei		96.09	0.42	RHA 1	99		
Clostridium perfringens		1.79	0.13		84		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		1.20	0.15		1		
27111360	**Good identification**						**27111360**
Clostridium perfringens		97.35	0.40	RHA 1			
Actinomyces odontolyticus		2.65	0.19				
27111400	**Low discrimination**						**27111400**
					COC		
Gemella morbillorum		61.38	0.45	NIT 1	99		
Lactobacillus fermentum		26.29	0.50	ARA 10	1		
Actinomyces odontolyticus		9.63	0.36		1		
27111402	**Low discrimination**						**27111402**
Actinomyces viscosus		48.13	0.38	RAF 94			
Lactobacillus fermentum		41.10	0.50	ARA 1			
Actinomyces odontolyticus		9.22	0.32				
27111420	**Low discrimination**						**27111420**
					COC		
Clostridium perfringens		59.59	0.38	LAC 88	1		
Lactobacillus fermentum		20.65	0.42	ARA 1	1		
Gemella morbillorum		16.07	0.29		99		
27111500	**Low discrimination**						**27111500**
					SPO		
Lactobacillus fermentum		66.51	0.54	NIT 1	1		
Clostridium chauvoei		12.58	0.29		99		
Clostridium perfringens		8.89	0.27		84		
27111501	**Very good identification**						**27111501**
					SPO		
Clostridium chauvoei		99.89	0.63	ARA 1	99		
Lactobacillus fermentum		0.05	0.20		1		
Actinomyces naeslundii		0.03	0.26		1		
27111502	**Low discrimination**						**27111502**
					GRA	CAT	
Actinomyces viscosus		49.91	0.42	RAF 94	99	76	
Lactobacillus fermentum		39.11	0.54	ARA 1	99	1	
Capnocytophaga ochracea		3.29	0.20		1	90	
27111503	**Low discrimination**						**27111503**
					SPO		
Clostridium chauvoei		88.12	0.50	ARA 1	99		
Actinomyces viscosus		11.34	0.34		1		
Lactobacillus fermentum		0.27	0.20		1		
27111520	**Low discrimination**						**27111520**
					GRA	CAT	
Clostridium perfringens		86.96	0.53	ARA 1	99	1	
Lactobacillus fermentum		6.71	0.46		99	1	
Capnocytophaga ochracea		1.69	0.20		1	90	
27111521	**Good identification**						**27111521**
					SPO		
Clostridium chauvoei		99.21	0.55	ARA 1	99		
Clostridium perfringens		0.69	0.19		84		
Actinomyces naeslundii		0.10	0.26		1		
27111522	**Low discrimination**						**27111522**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		62.60	0.47	ARA 1	99	1		
Capnocytophaga ochracea		23.29	0.37		1	90		
Lactobacillus fermentum		10.26	0.46		99	1		
27111523	**Good identification**							**27111523**
					GRA	CAT		
Clostridium chauvoei		96.61	0.42	ARA 1	99	1		
Clostridium perfringens		1.80	0.13		99	1		
Capnocytophaga ochracea		0.67	0.02		1	90		
27111560	**Good identification**							**27111560**
					GRA	CAT		
Clostridium perfringens		98.88	0.40	ARA 1	99	1		
Capnocytophaga ochracea		1.12	0.04		1	90		
27111600	**Good identification**							**27111600**
					COC			
Actinomyces odontolyticus		95.48	0.41	MNS 9	1			
Gemella morbillorum		3.17	0.11	ARA 1	99			
Lactobacillus fermentum		1.36	0.16		1			
27111602	**Good identification**							**27111602**
Actinomyces odontolyticus		95.21	0.37	MNS 9				
Actinomyces viscosus		2.59	0.03	ARA 1				
Lactobacillus fermentum		2.21	0.16					
27112000	**Low discrimination**							**27112000**
					CAT			
Propionibacterium propionicum		57.28	0.60	LAC 99	1			
Propionibacterium granulosum		21.31	0.48		90			
Actinomyces odontolyticus		8.65	0.39		10			
27112001	**Low discrimination**							**27112001**
					CAT			
Propionibacterium propionicum		89.26	0.44	LAC 99	1			
Actinomyces naeslundii		6.30	0.34	URE 10	25			
Propionibacterium granulosum		3.02	0.14		90			
27112002	**Low discrimination**							**27112002**
					CAT			
Propionibacterium propionicum		67.37	0.63	LAC 99	1			
Actinomyces viscosus		23.61	0.41		76			
Propionibacterium granulosum		4.18	0.38		90			
27112020	**Low discrimination**							**27112020**
					CAT			
Propionibacterium granulosum		58.08	0.38	NIT 1	90			
Actinomyces odontolyticus		15.36	0.25		10			
Propionibacterium acnes		7.80	0.13		95			
27112040	**Low discrimination**							**27112040**
					COC	SPO	GRA	
Propionibacterium propionicum		87.65	0.60	LAC 99	1	1	99	
Clostridium botulinum biovar B		6.79	0.24		1	99	99	
Megasphaera elsdenii		5.23	0.29		99	1	1	
27112041	**Good identification**							**27112041**
Propionibacterium propionicum		100.0	0.44	LAC 99				
27112042	**Very good identification**							**27112042**
					COC	GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		99.21	0.63	LAC 99	1	99	
Megasphaera elsdenii		0.44	0.12		99	1	
Actinomyces viscosus		0.35	0.07		1	99	
27112043	**Good identification**						**27112043**
Propionibacterium propionicum		98.95	0.47	LAC 99			
Actinomyces viscosus		1.05	-.01	URE 10			
27112100	**Excellent identification**						**27112100**
Propionibacterium propionicum		99.92	0.94				
Actinomyces odontolyticus		0.05	0.30				
Actinomyces naeslundii		0.02	0.39				
27112101	**Very good identification**						**27112101**
					SPO		
Propionibacterium propionicum		99.01	0.78	URE 10	1		
Clostridium chauvoei		0.76	0.29		99		
Actinomyces naeslundii		0.20	0.42		1		
27112102	**Excellent identification**						**27112102**
Propionibacterium propionicum		99.35	0.97				
Actinomyces viscosus		0.63	0.45				
Actinomyces odontolyticus		0.02	0.27				
27112103	**Very good identification**						**27112103**
Propionibacterium propionicum		97.93	0.81	URE 10			
Actinomyces viscosus		1.85	0.37				
Actinomyces naeslundii		0.13	0.42				
27112110	**Very good identification**						**27112110**
Propionibacterium propionicum		100.0	0.60	MLZ 1			
27112111	**Good identification**						**27112111**
Propionibacterium propionicum		100.0	0.44	MLZ 1			
27112112	**Very good identification**						**27112112**
Propionibacterium propionicum		99.38	0.63	MLZ 1			
Actinomyces viscosus		0.63	0.11				
27112113	**Good identification**						**27112113**
Propionibacterium propionicum		98.15	0.47	MLZ 1			
Actinomyces viscosus		1.85	0.03	URE 10			
27112120	**Very good identification**						**27112120**
Propionibacterium propionicum		95.95	0.60	NAG 1			
Clostridium perfringens		1.88	0.19				
Actinomyces naeslundii		1.43	0.39				
27112121	**Low discrimination**						**27112121**
					SPO		
Propionibacterium propionicum		68.70	0.44	NAG 1	1		
Clostridium chauvoei		17.50	0.21	URE 10	99		
Actinomyces naeslundii		13.81	0.42		1		
27112122	**Very good identification**						**27112122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		97.51	0.63	NAG 1		
Actinomyces naeslundii		0.97	0.39			
Actinomyces viscosus		0.61	0.11			
27112123	**Low discrimination**					**27112123**
					SPO	
Propionibacterium propionicum		84.54	0.47	NAG 1	1	
Actinomyces naeslundii		11.33	0.42	URE 10	1	
Clostridium chauvoei		2.53	0.08		99	
27112140	**Excellent identification**					**27112140**
Propionibacterium propionicum		100.0	0.94			
Actinomyces naeslundii		0.00	0.26			
27112141	**Very good identification**					**27112141**
Propionibacterium propionicum		99.96	0.78	URE 10		
Actinomyces naeslundii		0.04	0.29			
27112142	**Excellent identification**					**27112142**
Propionibacterium propionicum		99.99	0.97			
Actinomyces viscosus		0.01	0.11			
Actinomyces naeslundii		0.00	0.26			
27112143	**Very good identification**					**27112143**
Propionibacterium propionicum		99.96	0.81	URE 10		
Actinomyces naeslundii		0.02	0.29			
Actinomyces viscosus		0.02	0.03			
27112150	**Very good identification**					**27112150**
Propionibacterium propionicum		100.0	0.60	MLZ 1		
27112151	**Good identification**					**27112151**
Propionibacterium propionicum		100.0	0.44	MLZ 1		
27112152	**Very good identification**					**27112152**
Propionibacterium propionicum		100.0	0.63	MLZ 1		
27112153	**Good identification**					**27112153**
Propionibacterium propionicum		100.0	0.47	MLZ 1		
27112160	**Very good identification**					**27112160**
Propionibacterium propionicum		99.37	0.60	NAG 1		
Clostridium perfringens		0.37	0.06			
Actinomyces naeslundii		0.26	0.26			
27112161	**Good identification**					**27112161**
Propionibacterium propionicum		96.58	0.44	NAG 1		
Actinomyces naeslundii		3.43	0.29	URE 10		
27112162	**Very good identification**					**27112162**
Propionibacterium propionicum		99.71	0.63	NAG 1		
Actinomyces naeslundii		0.18	0.26			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		0.12	0.01			
27112163	**Good identification**					**27112163**
Propionibacterium propionicum		97.69	0.47	NAG 1		
Actinomyces naeslundii		2.31	0.29	URE 10		
27112200	**Acceptable identification**					**27112200**
					CAT	
Actinomyces odontolyticus		93.14	0.44	MNS 9	10	
Propionibacterium granulosum		3.65	0.22	MAN 1	90	
Propionibacterium propionicum		3.21	0.26		1	
27112202	**Low discrimination**					**27112202**
Actinomyces odontolyticus		88.51	0.40	MNS 9		
Propionibacterium propionicum		7.46	0.29	MAN 1		
Actinomyces viscosus		2.62	0.07			
27112300	**Good identification**					**27112300**
Propionibacterium propionicum		91.53	0.60	RHA 1		
Actinomyces odontolyticus		8.47	0.35			
27112301	**Good identification**					**27112301**
Propionibacterium propionicum		100.0	0.44	RHA 1		
27112302	**Very good identification**					**27112302**
Propionibacterium propionicum		95.77	0.63	RHA 1		
Actinomyces odontolyticus		3.62	0.32			
Actinomyces viscosus		0.60	0.11			
27112303	**Good identification**					**27112303**
Propionibacterium propionicum		98.15	0.47	RHA 1		
Actinomyces viscosus		1.85	0.03	URE 10		
27112340	**Very good identification**					**27112340**
Propionibacterium propionicum		100.0	0.60	RHA 1		
27112341	**Good identification**					**27112341**
Propionibacterium propionicum		100.0	0.44	RHA 1		
27112342	**Very good identification**					**27112342**
Propionibacterium propionicum		100.0	0.63	RHA 1		
27112343	**Good identification**					**27112343**
Propionibacterium propionicum		100.0	0.47	RHA 1		
27112500	**Very good identification**					**27112500**
Propionibacterium propionicum		100.0	0.60	ARA 1		
27112501	**Good identification**					**27112501**
Propionibacterium propionicum		100.0	0.44	ARA 1		
27112502	**Very good identification**					**27112502**
Propionibacterium propionicum		99.38	0.63	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.63	0.11			
27112503	**Good identification**					**27112503**
Propionibacterium propionicum		98.15	0.47	ARA 1		
Actinomyces viscosus		1.85	0.03	URE 10		
27112540	**Very good identification**					**27112540**
Propionibacterium propionicum		100.0	0.60	ARA 1		
27112541	**Good identification**					**27112541**
Propionibacterium propionicum		100.0	0.44	ARA 1		
27112542	**Very good identification**					**27112542**
Propionibacterium propionicum		100.0	0.63	ARA 1		
27112543	**Good identification**					**27112543**
Propionibacterium propionicum		100.0	0.47	ARA 1		
27113000	**Low discrimination**					**27113000**
					CAT	
Propionibacterium propionicum		59.54	0.60	LAC 99	1	
Propionibacterium granulosum		22.15	0.48		90	
Actinomyces naeslundii		7.47	0.55		25	
27113001	**Low discrimination**					**27113001**
					SPO	
Actinomyces naeslundii		48.52	0.58	RAF 92	1	
Propionibacterium propionicum		28.63	0.44		1	
Clostridium chauvoei		21.88	0.29		99	
27113002	**Low discrimination**					**27113002**
Propionibacterium propionicum		71.12	0.63	LAC 99		
Actinomyces viscosus		15.28	0.38			
Actinomyces naeslundii		5.95	0.55			
27113003	**Low discrimination**					**27113003**
Actinomyces naeslundii		39.45	0.58	RAF 92		
Propionibacterium propionicum		34.92	0.47			
Actinomyces viscosus		22.50	0.29			
27113011	**Good identification**					**27113011**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27113013	**Low discrimination**					**27113013**
Actinomyces naeslundii		85.48	0.36	RAF 92		
Propionibacterium propionicum		14.52	0.12	MLZ 5		
27113020	**Low discrimination**					**27113020**
					CAT	
Clostridium perfringens		50.06	0.38	MAN 1	1	
Actinomyces naeslundii		23.75	0.55	LAC 88	25	
Propionibacterium granulosum		17.59	0.38		90	
27113021	**Low discrimination**					**27113021**
					SPO	
Actinomyces naeslundii		85.87	0.58	RAF 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		12.91	0.21		99	
Clostridium perfringens		1.22	0.04		84	
27113022	**Low discrimination**					**27113022**
					CAT	
Actinomyces naeslundii		41.60	0.55	RAF 92	25	
Clostridium perfringens		41.26	0.32		1	
Propionibacterium granulosum		7.71	0.27		90	
27113023	**Very good identification**					**27113023**
					SPO	
Actinomyces naeslundii		96.58	0.58	RAF 92	1	
Clostridium chauvoei		2.56	0.08		99	
Propionibacterium propionicum		0.86	0.12		1	
27113031	**Good identification**					**27113031**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27113033	**Good identification**					**27113033**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27113040	**Very good identification**					**27113040**
					CAT	
Propionibacterium propionicum		97.47	0.60	LAC 99	1	
Actinomyces naeslundii		2.16	0.42		25	
Propionibacterium granulosum		0.37	0.14		90	
27113041	**Low discrimination**					**27113041**
Propionibacterium propionicum		76.98	0.44	LAC 99		
Actinomyces naeslundii		23.02	0.45	URE 10		
27113042	**Very good identification**					**27113042**
Propionibacterium propionicum		98.33	0.63	LAC 99		
Actinomyces naeslundii		1.45	0.42			
Actinomyces viscosus		0.21	0.03			
27113043	**Low discrimination**					**27113043**
Propionibacterium propionicum		83.38	0.47	LAC 99		
Actinomyces naeslundii		16.62	0.45	URE 10		
27113061	**Good identification**					**27113061**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27113063	**Good identification**					**27113063**
Actinomyces naeslundii		95.18	0.45	RAF 92		
Propionibacterium propionicum		4.82	0.12			
27113100	**Excellent identification**					**27113100**
					SPO	
Propionibacterium propionicum		99.45	0.94		1	
Actinomyces naeslundii		0.36	0.62		1	
Clostridium chauvoei		0.09	0.29		99	
27113101	**Low discrimination**					**27113101**
					SPO	
Propionibacterium propionicum		55.15	0.78	URE 10	1	
Clostridium chauvoei		42.15	0.63		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	2.69	0.65		1	
27113102	**Excellent identification**					**27113102**
	Propionibacterium propionicum	99.33	0.97			
	Actinomyces viscosus	0.38	0.42			
	Actinomyces naeslundii	0.24	0.62			
27113103	**Low discrimination**					**27113103**
					SPO	
	Propionibacterium propionicum	88.19	0.81	URE 10	1	
	Clostridium chauvoei	7.93	0.50		99	
	Actinomyces naeslundii	2.87	0.65		1	
27113110	**Very good identification**					**27113110**
	Propionibacterium propionicum	98.15	0.60	MLZ 1		
	Actinomyces naeslundii	1.85	0.41			
27113111	**Low discrimination**					**27113111**
					SPO	
	Propionibacterium propionicum	49.55	0.44	MLZ 1	1	
	Clostridium chauvoei	37.87	0.29	URE 10	99	
	Actinomyces naeslundii	12.58	0.44		1	
27113112	**Very good identification**					**27113112**
	Propionibacterium propionicum	98.39	0.63	MLZ 1		
	Actinomyces naeslundii	1.23	0.41			
	Actinomyces viscosus	0.38	0.08			
27113113	**Low discrimination**					**27113113**
					SPO	
	Propionibacterium propionicum	78.70	0.47	MLZ 1	1	
	Actinomyces naeslundii	13.32	0.44	URE 10	1	
	Clostridium chauvoei	7.08	0.16		99	
27113120	**Low discrimination**					**27113120**
	Clostridium perfringens	57.65	0.53	MAN 1		
	Propionibacterium propionicum	29.71	0.60			
	Actinomyces naeslundii	10.62	0.62			
27113121	**Low discrimination**					**27113121**
					SPO	
	Clostridium chauvoei	80.78	0.55	MAN 1	99	
	Actinomyces naeslundii	15.45	0.65		1	
	Propionibacterium propionicum	3.20	0.44		1	
27113122	**Low discrimination**					**27113122**
	Propionibacterium propionicum	53.38	0.63	NAG 1		
	Clostridium perfringens	32.50	0.47			
	Actinomyces naeslundii	12.72	0.62			
27113123	**Low discrimination**					**27113123**
					SPO	
	Actinomyces naeslundii	44.36	0.65	RAF 92	1	
	Clostridium chauvoei	40.93	0.42		99	
	Propionibacterium propionicum	13.79	0.47		1	
27113132	**Low discrimination**					**27113132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		41.48	0.41	RAF 92		
Propionibacterium propionicum		33.42	0.29	MLZ 5		
Clostridium perfringens		20.34	0.13			
27113133	**Low discrimination**					**27113133**
					SPO	
Actinomyces naeslundii		80.86	0.44	RAF 92	1	
Clostridium chauvoei		14.32	0.08	MLZ 5	99	
Propionibacterium propionicum		4.83	0.12		1	
27113140	**Excellent identification**					**27113140**
Propionibacterium propionicum		99.93	0.94			
Actinomyces naeslundii		0.06	0.50			
Clostridium perfringens		0.01	0.15			
27113141	**Very good identification**					**27113141**
					SPO	
Propionibacterium propionicum		98.39	0.78	URE 10	1	
Actinomyces naeslundii		0.85	0.53		1	
Clostridium chauvoei		0.76	0.29		99	
27113142	**Excellent identification**					**27113142**
Propionibacterium propionicum		99.95	0.97			
Actinomyces naeslundii		0.04	0.50			
Clostridium perfringens		0.00	0.09			
27113143	**Very good identification**					**27113143**
					SPO	
Propionibacterium propionicum		99.33	0.81	URE 10	1	
Actinomyces naeslundii		0.57	0.53		1	
Clostridium chauvoei		0.09	0.16		99	
27113150	**Very good identification**					**27113150**
Propionibacterium propionicum		99.67	0.60	MLZ 1		
Actinomyces naeslundii		0.33	0.28			
27113151	**Good identification**					**27113151**
Propionibacterium propionicum		95.71	0.44	MLZ 1		
Actinomyces naeslundii		4.29	0.31	URE 10		
27113152	**Very good identification**					**27113152**
Propionibacterium propionicum		99.78	0.63	MLZ 1		
Actinomyces naeslundii		0.22	0.28			
27113153	**Good identification**					**27113153**
Propionibacterium propionicum		97.10	0.47	MLZ 1		
Actinomyces naeslundii		2.90	0.31	URE 10		
27113160	**Low discrimination**					**27113160**
Propionibacterium propionicum		69.80	0.60	NAG 1		
Clostridium perfringens		25.80	0.40			
Actinomyces naeslundii		4.40	0.50			
27113161	**Low discrimination**					**27113161**
					SPO	
Propionibacterium propionicum		46.74	0.44	NAG 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		39.79	0.53	URE 10	1	
Clostridium chauvoei		11.91	0.21		99	
27113162	**Low discrimination**					**27113162**
Propionibacterium propionicum		86.25	0.63	NAG 1		
Clostridium perfringens		10.00	0.35			
Actinomyces naeslundii		3.63	0.50			
27113163	**Low discrimination**					**27113163**
					SPO	
Propionibacterium propionicum		62.19	0.47	NAG 1	1	
Actinomyces naeslundii		35.29	0.53	URE 10	1	
Clostridium chauvoei		1.86	0.08		99	
27113200	**Acceptable identification**					**27113200**
					CAT	
Actinomyces odontolyticus		90.05	0.41	MNS 9	10	
Propionibacterium granulosum		5.30	0.22	MAN 1	90	
Propionibacterium propionicum		4.65	0.26		1	
27113202	**Low discrimination**					**27113202**
Actinomyces odontolyticus		84.92	0.37	MNS 9		
Propionibacterium propionicum		10.74	0.29	MAN 1		
Actinomyces viscosus		2.31	0.03			
27113300	**Good identification**					**27113300**
Propionibacterium propionicum		93.87	0.60	RHA 1		
Actinomyces odontolyticus		5.79	0.32			
Actinomyces naeslundii		0.34	0.28			
27113301	**Low discrimination**					**27113301**
					SPO	
Propionibacterium propionicum		55.16	0.44	RHA 1	1	
Clostridium chauvoei		42.15	0.29	URE 10	99	
Actinomyces naeslundii		2.69	0.31		1	
27113302	**Very good identification**					**27113302**
Propionibacterium propionicum		96.95	0.63	RHA 1		
Actinomyces odontolyticus		2.45	0.29			
Actinomyces viscosus		0.37	0.08			
27113303	**Low discrimination**					**27113303**
					SPO	
Propionibacterium propionicum		88.19	0.47	RHA 1	1	
Clostridium chauvoei		7.93	0.16	URE 10	99	
Actinomyces naeslundii		2.86	0.31		1	
27113340	**Very good identification**					**27113340**
Propionibacterium propionicum		100.0	0.60	RHA 1		
27113341	**Good identification**					**27113341**
Propionibacterium propionicum		100.0	0.44	RHA 1		
27113342	**Very good identification**					**27113342**
Propionibacterium propionicum		100.0	0.63	RHA 1		
27113343	**Good identification**					**27113343**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.47	RHA 1		
27113500	**Very good identification**					**27113500**
Propionibacterium propionicum		99.19	0.60	ARA 1		
Lactobacillus fermentum		0.45	0.20			
Actinomyces naeslundii		0.36	0.28			
27113501	**Low discrimination**					**27113501**
					SPO	
Propionibacterium propionicum		55.16	0.44	ARA 1	1	
Clostridium chauvoei		42.15	0.29	URE 10	99	
Actinomyces naeslundii		2.69	0.31		1	
27113502	**Very good identification**					**27113502**
Propionibacterium propionicum		98.97	0.63	ARA 1		
Actinomyces viscosus		0.38	0.08			
Lactobacillus fermentum		0.30	0.20			
27113503	**Low discrimination**					**27113503**
					SPO	
Propionibacterium propionicum		88.19	0.47	ARA 1	1	
Clostridium chauvoei		7.93	0.16	URE 10	99	
Actinomyces naeslundii		2.86	0.31		1	
27113540	**Very good identification**					**27113540**
Propionibacterium propionicum		100.0	0.60	ARA 1		
27113541	**Good identification**					**27113541**
Propionibacterium propionicum		100.0	0.44	ARA 1		
27113542	**Very good identification**					**27113542**
Propionibacterium propionicum		100.0	0.63	ARA 1		
27113543	**Good identification**					**27113543**
Propionibacterium propionicum		100.0	0.47	ARA 1		
27114000	**Low discrimination**					**27114000**
					COC	
Actinomyces odontolyticus		85.66	0.68	MNS 9	1	
Gemella morbillorum		13.87	0.50		99	
Actinomyces meyeri		0.18	0.18		1	
27114002	**Low discrimination**					**27114002**
					COC	
Actinomyces odontolyticus		89.33	0.64	MNS 9	1	
Actinomyces viscosus		10.40	0.41		1	
Gemella morbillorum		0.24	0.15		99	
27114020	**Low discrimination**					**27114020**
					COC	
Actinomyces odontolyticus		89.72	0.54	MNS 9	1	
Gemella morbillorum		9.92	0.33	NAG 14	99	
Clostridium perfringens		0.20	0.04		1	
27114022	**Good identification**					**27114022**
Actinomyces odontolyticus		99.28	0.51	MNS 9		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.72	0.07	NAG 14		
27114100	**Very good identification**					**27114100**
					COC	
Actinomyces odontolyticus		95.86	0.59	MNS 9	1	
Actinomyces meyeri		2.61	0.29		1	
Gemella morbillorum		0.50	0.15		99	
27114102	**Low discrimination**					**27114102**
Actinomyces odontolyticus		59.90	0.56	MNS 9		
Actinomyces viscosus		39.27	0.45			
Propionibacterium propionicum		0.64	0.29			
27114103	**Acceptable identification**					**27114103**
					SPO	
Actinomyces viscosus		90.93	0.37	RAF 94	1	
Clostridium chauvoei		4.37	0.16	XYL 1	99	
Actinomyces odontolyticus		4.20	0.21		1	
27114120	**Acceptable identification**					**27114120**
Actinomyces odontolyticus		93.06	0.46	MNS 9		
Clostridium perfringens		4.82	0.19	NAG 14		
Actinomyces meyeri		2.12	0.14			
27114122	**Acceptable identification**					**27114122**
Actinomyces odontolyticus		90.80	0.42	MNS 9		
Actinomyces viscosus		3.69	0.11	NAG 14		
Clostridium perfringens		3.61	0.13			
27114200	**Very good identification**					**27114200**
					COC	
Actinomyces odontolyticus		99.92	0.73	MNS 9	1	
Gemella morbillorum		0.08	0.15		99	
27114201	**Good identification**					**27114201**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27114202	**Very good identification**					**27114202**
Actinomyces odontolyticus		99.94	0.69	MNS 9		
Actinomyces viscosus		0.06	0.07			
27114210	**Good identification**					**27114210**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27114220	**Very good identification**					**27114220**
Actinomyces odontolyticus		100.0	0.59	MNS 9		
27114222	**Good identification**					**27114222**
Actinomyces odontolyticus		100.0	0.56	MNS 9		
27114240	**Good identification**					**27114240**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27114300	**Very good identification**					**27114300**
Actinomyces odontolyticus		100.0	0.64	MNS 9		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27114302	**Very good identification**					**27114302**
Actinomyces odontolyticus		99.66	0.60	MNS 9		
Actinomyces viscosus		0.34	0.11			
27114320	**Good identification**					**27114320**
Actinomyces odontolyticus		100.0	0.51	MNS 9		
27114322	**Good identification**					**27114322**
Actinomyces odontolyticus		100.0	0.47	MNS 9		
27114600	**Good identification**					**27114600**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27115000	**Low discrimination**					**27115000**
					COC	
Actinomyces odontolyticus		85.18	0.65	MNS 9	1	
Gemella morbillorum		11.14	0.45		99	
Lactobacillus fermentum		2.80	0.46		1	
27115002	**Low discrimination**					**27115002**
Actinomyces odontolyticus		85.90	0.61	MNS 9		
Actinomyces viscosus		9.20	0.38			
Lactobacillus fermentum		4.61	0.46			
27115020	**Low discrimination**					**27115020**
					COC	
Actinomyces odontolyticus		67.04	0.51	MNS 9	1	
Clostridium perfringens		22.19	0.38	NAG 14	1	
Gemella morbillorum		5.98	0.29		99	
27115022	**Low discrimination**					**27115022**
Actinomyces odontolyticus		72.43	0.48	MNS 9		
Clostridium perfringens		18.41	0.32	NAG 14		
Lactobacillus fermentum		7.96	0.38			
27115100	**Low discrimination**					**27115100**
Actinomyces odontolyticus		73.57	0.56	MNS 9		
Lactobacillus fermentum		12.51	0.50			
Actinomyces meyeri		4.51	0.32			
27115101	**Very good identification**					**27115101**
					SPO	
Clostridium chauvoei		99.73	0.63	XYL 1	99	
Actinomyces odontolyticus		0.19	0.22		1	
Actinomyces naeslundii		0.03	0.26		1	
27115102	**Low discrimination**					**27115102**
Actinomyces odontolyticus		49.75	0.53	MNS 9		
Actinomyces viscosus		29.98	0.42			
Lactobacillus fermentum		13.80	0.50			
27115103	**Low discrimination**					**27115103**
					SPO	
Clostridium chauvoei		87.72	0.50	XYL 1	99	
Actinomyces viscosus		11.29	0.34		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		0.57	0.18		1		
27115120	**Low discrimination**						**27115120**
Clostridium perfringens		80.29	0.53	XYL 1			
Actinomyces odontolyticus		10.45	0.43				
Lactobacillus fermentum		3.64	0.42				
27115121	**Good identification**						**27115121**
					SPO		
Clostridium chauvoei		99.12	0.55	XYL 1	99		
Clostridium perfringens		0.69	0.19		84		
Actinomyces naeslundii		0.10	0.26		1		
27115122	**Low discrimination**						**27115122**
					GRA	CAT	
Clostridium perfringens		54.36	0.47	XYL 1	99	1	
Capnocytophaga ochracea		28.65	0.39		1	90	
Actinomyces odontolyticus		9.21	0.39		99	10	
27115123	**Good identification**						**27115123**
					GRA	CAT	
Clostridium chauvoei		96.34	0.42	XYL 1	99	1	
Clostridium perfringens		1.80	0.13		99	1	
Capnocytophaga ochracea		0.95	0.05		1	90	
27115160	**Good identification**						**27115160**
					GRA	CAT	
Clostridium perfringens		97.75	0.40	XYL 1	99	1	
Capnocytophaga ochracea		1.57	0.07		1	90	
Actinomyces odontolyticus		0.67	0.08		99	10	
27115200	**Very good identification**						**27115200**
					COC		
Actinomyces odontolyticus		99.92	0.70	MNS 9	1		
Gemella morbillorum		0.07	0.11		99		
Lactobacillus fermentum		0.02	0.12		1		
27115201	**Good identification**						**27115201**
Actinomyces odontolyticus		100.0	0.35	MNS 9			
27115202	**Very good identification**						**27115202**
Actinomyces odontolyticus		99.92	0.66	MNS 9			
Actinomyces viscosus		0.06	0.03				
Lactobacillus fermentum		0.03	0.12				
27115210	**Good identification**						**27115210**
Actinomyces odontolyticus		100.0	0.35	MNS 9			
27115220	**Very good identification**						**27115220**
Actinomyces odontolyticus		99.83	0.56	MNS 9			
Clostridium perfringens		0.17	0.04	NAG 14			
27115222	**Good identification**						**27115222**
Actinomyces odontolyticus		100.0	0.53	MNS 9			
27115240	**Good identification**						**27115240**
Actinomyces odontolyticus		100.0	0.35	MNS 9			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27115300	**Very good identification**						**27115300**
Actinomyces odontolyticus		99.91	0.61	MNS 9			
Lactobacillus fermentum		0.09	0.16				
27115302	**Very good identification**						**27115302**
Actinomyces odontolyticus		99.54	0.57	MNS 9			
Actinomyces viscosus		0.31	0.08				
Lactobacillus fermentum		0.14	0.16				
27115320	**Low discrimination**						**27115320**
					GRA		
Actinomyces odontolyticus		89.40	0.48	MNS 9	99		
Bacteroides vulgatus		7.02	0.15	NAG 14	1		
Clostridium perfringens		3.58	0.19		99		
27115322	**Acceptable identification**						**27115322**
					GRA	CAT	
Actinomyces odontolyticus		93.97	0.44	MNS 9	99	10	
Clostridium perfringens		2.89	0.13	NAG 14	99	1	
Capnocytophaga ochracea		1.52	0.05		1	90	
27115400	**Low discrimination**						**27115400**
					COC		
Lactobacillus fermentum		56.01	0.46	NIT 1	1		
Gemella morbillorum		24.74	0.29		99		
Actinomyces odontolyticus		17.19	0.31		1		
27115402	**Low discrimination**						**27115402**
Lactobacillus fermentum		82.76	0.46	NIT 1			
Actinomyces odontolyticus		15.57	0.27				
Actinomyces viscosus		1.67	0.03				
27115420	**Low discrimination**						**27115420**
					COC		
Lactobacillus fermentum		74.25	0.38	NIT 1	1		
Actinomyces odontolyticus		11.13	0.17		1		
Gemella morbillorum		10.93	0.12		99		
27115422	**Acceptable identification**						**27115422**
Lactobacillus fermentum		91.59	0.38	NIT 1			
Actinomyces odontolyticus		8.42	0.13				
27115500	**Low discrimination**						**27115500**
Lactobacillus fermentum		79.62	0.50	NIT 1			
Bifidobacterium longum subsp. longum		7.79	0.26				
Actinomyces meyeri		7.18	0.21				
27115502	**Low discrimination**						**27115502**
Lactobacillus fermentum		81.26	0.50	NIT 1			
Bifidobacterium longum subsp. longum		7.95	0.26				
Actinomyces odontolyticus		2.96	0.18				
27115520	**Low discrimination**						**27115520**
					GRA		
Bifidobacterium longum subsp. longum		36.49	0.36	NIT 1	99		
Lactobacillus fermentum		31.09	0.42	RAF 97	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		20.10	0.17		1		
27115522	**Low discrimination**						**27115522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		36.17	0.36	NIT 1	99	1	
Lactobacillus fermentum		30.82	0.42	RAF 97	99	1	
Capnocytophaga ochracea		29.78	0.26		1	90	
27115530	**Good identification**						**27115530**
Bifidobacterium longum subsp. longum		100.0	0.40	NIT 1			
27115532	**Acceptable identification**						**27115532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		94.90	0.40	NIT 1	99	1	
Capnocytophaga ochracea		5.10	0.10	RAF 97	1	90	
27115600	**Good identification**						**27115600**
Actinomyces odontolyticus		98.33	0.35	MNS 9			
Lactobacillus fermentum		1.67	0.12	ARA 1			
27116100	**Very good identification**						**27116100**
Propionibacterium propionicum		97.71	0.60	XYL 1			
Actinomyces odontolyticus		2.30	0.25				
27116101	**Good identification**						**27116101**
Propionibacterium propionicum		100.0	0.44	XYL 1			
27116102	**Very good identification**						**27116102**
Propionibacterium propionicum		98.44	0.63	XYL 1			
Actinomyces odontolyticus		0.95	0.21				
Actinomyces viscosus		0.62	0.11				
27116103	**Good identification**						**27116103**
Propionibacterium propionicum		98.15	0.47	XYL 1			
Actinomyces viscosus		1.85	0.03	URE 10			
27116140	**Very good identification**						**27116140**
Propionibacterium propionicum		100.0	0.60	XYL 1			
27116141	**Good identification**						**27116141**
Propionibacterium propionicum		100.0	0.44	XYL 1			
27116142	**Very good identification**						**27116142**
Propionibacterium propionicum		100.0	0.63	XYL 1			
27116143	**Good identification**						**27116143**
Propionibacterium propionicum		100.0	0.47	XYL 1			
27116200	**Good identification**						**27116200**
Actinomyces odontolyticus		100.0	0.38	MNS 9			
27117100	**Very good identification**						**27117100**
Propionibacterium propionicum		96.36	0.60	XYL 1			
Actinomyces israelii		1.53	0.30				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		1.51	0.22			
27117101	**Low discrimination**					**27117101**
					SPO	
Propionibacterium propionicum		55.16	0.44	XYL 1	1	
Clostridium chauvoei		42.15	0.29	URE 10	99	
Actinomyces naeslundii		2.69	0.31		1	
27117102	**Very good identification**					**27117102**
Propionibacterium propionicum		97.06	0.63	XYL 1		
Actinomyces israelii		1.54	0.33			
Actinomyces odontolyticus		0.62	0.18			
27117103	**Low discrimination**					**27117103**
					SPO	
Propionibacterium propionicum		88.19	0.47	XYL 1	1	
Clostridium chauvoei		7.93	0.16	URE 10	99	
Actinomyces naeslundii		2.86	0.31		1	
27117140	**Very good identification**					**27117140**
Propionibacterium propionicum		99.39	0.60	XYL 1		
Actinomyces israelii		0.61	0.23			
27117141	**Good identification**					**27117141**
Propionibacterium propionicum		100.0	0.44	XYL 1		
27117142	**Very good identification**					**27117142**
Propionibacterium propionicum		99.39	0.63	XYL 1		
Actinomyces israelii		0.61	0.26			
27117143	**Good identification**					**27117143**
Propionibacterium propionicum		100.0	0.47	XYL 1		
27117200	**Good identification**					**27117200**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27120040	**Good identification**					**27120040**
Clostridium botulinum biovar B		99.95	0.56	NIT 1		
Clostridium botulinum biovar A		0.05	0.02			
27120050	**Good identification**					**27120050**
Clostridium botulinum biovar B		100.0	0.50	NIT 1		
27120060	**Good identification**					**27120060**
Clostridium botulinum biovar B		100.0	0.50	NIT 1		
27120070	**Good identification**					**27120070**
Clostridium botulinum biovar B		100.0	0.45	NIT 1		
27120101	**Good identification**					**27120101**
Clostridium chauvoei		100.0	0.40	TRE 10		
27120102	**Good identification**					**27120102**
Actinomyces viscosus		96.08	0.38	RAF 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		3.25	0.27	FRU 95		
Actinomyces odontolyticus		0.67	0.08	TRE 7		
27121001	**Good identification**					**27121001**
					SPO	
Clostridium chauvoei		97.58	0.40	TRE 10	99	
Actinomyces naeslundii		2.42	0.36	LAC 99	1	
27121020	**Good identification**					**27121020**
					COC	
Clostridium perfringens		96.91	0.43	FRU 99	1	
Actinomyces naeslundii		1.30	0.33	LAC 88	1	
Gemella morbillorum		1.05	0.09		99	
27121022	**Good identification**					**27121022**
Clostridium perfringens		97.22	0.37	FRU 99		
Actinomyces naeslundii		2.78	0.33	LAC 88		
27121100	**Low discrimination**					**27121100**
					SPO	
Clostridium chauvoei		72.46	0.40	TRE 10	99	
Clostridium perfringens		20.17	0.31	URE 99	84	
Actinomyces naeslundii		3.40	0.41		1	
27121101	**Very good identification**					**27121101**
					SPO	
Clostridium chauvoei		99.93	0.74	TRE 10	99	
Actinomyces naeslundii		0.07	0.44		1	
27121103	**Very good identification**					**27121103**
					SPO	
Clostridium chauvoei		98.55	0.62	TRE 10	99	
Actinomyces viscosus		1.06	0.27		1	
Actinomyces naeslundii		0.40	0.44		1	
27121111	**Good identification**					**27121111**
Clostridium chauvoei		100.0	0.40	TRE 10		
27121120	**Very good identification**					**27121120**
					SPO	
Clostridium perfringens		95.86	0.57	FRU 99	84	
Clostridium chauvoei		3.55	0.32		99	
Actinomyces naeslundii		0.50	0.41		1	
27121121	**Very good identification**					**27121121**
					SPO	
Clostridium chauvoei		99.51	0.66	TRE 10	99	
Clostridium perfringens		0.27	0.23		84	
Actinomyces naeslundii		0.21	0.44		1	
27121122	**Good identification**					**27121122**
					SPO	
Clostridium perfringens		97.44	0.52	FRU 99	84	
Clostridium chauvoei		1.35	0.19		99	
Actinomyces naeslundii		1.08	0.41		1	
27121123	**Good identification**					**27121123**
					SPO	
Clostridium chauvoei		98.09	0.53	TRE 10	99	
Actinomyces naeslundii		1.19	0.44		1	
Clostridium perfringens		0.72	0.18		84	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27121141	**Good identification**					**27121141**
					SPO	
Clostridium chauvoei		98.77	0.40	TRE 10	99	
Actinomyces naeslundii		1.23	0.31	SOR 1	1	
27121160	**Good identification**					**27121160**
Clostridium perfringens		99.52	0.45	FRU 99		
Actinomyces naeslundii		0.48	0.28			
27121162	**Good identification**					**27121162**
Clostridium perfringens		98.98	0.39	FRU 99		
Actinomyces naeslundii		1.02	0.28			
27121301	**Good identification**					**27121301**
Clostridium chauvoei		100.0	0.40	TRE 10		
27121501	**Good identification**					**27121501**
Clostridium chauvoei		100.0	0.40	TRE 10		
27122100	**Very good identification**					**27122100**
Propionibacterium propionicum		100.0	0.58	FRU 99		
27122101	**Acceptable identification**					**27122101**
					SPO	
Propionibacterium propionicum		93.50	0.42	FRU 99	1	
Clostridium chauvoei		4.16	0.06	URE 10	99	
Actinomyces naeslundii		2.35	0.25		1	
27122102	**Very good identification**					**27122102**
Propionibacterium propionicum		99.70	0.61	FRU 99		
Actinomyces viscosus		0.30	0.04			
27122103	**Good identification**					**27122103**
Propionibacterium propionicum		98.35	0.45	FRU 99		
Actinomyces naeslundii		1.65	0.25	URE 10		
27122140	**Very good identification**					**27122140**
Propionibacterium propionicum		100.0	0.58	FRU 99		
27122141	**Good identification**					**27122141**
Propionibacterium propionicum		100.0	0.42	FRU 99		
27122142	**Very good identification**					**27122142**
Propionibacterium propionicum		100.0	0.61	FRU 99		
27122143	**Good identification**					**27122143**
Propionibacterium propionicum		100.0	0.45	FRU 99		
27123000	**Low discrimination**					**27123000**
					CAT	
Actinomyces naeslundii		39.63	0.38	RAF 92	25	
Propionibacterium granulosum		34.83	0.22	FRU 95	90	
Propionibacterium propionicum		25.54	0.24		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27123001	**Low discrimination**					**27123001**
					SPO	
Actinomyces naeslundii		82.63	0.41	RAF 92	1	
Clostridium chauvoei		17.37	0.06	FRU 95	99	
27123002	**Low discrimination**					**27123002**
					CAT	
Actinomyces naeslundii		45.74	0.38	RAF 92	25	
Propionibacterium propionicum		44.21	0.27	FRU 95	1	
Propionibacterium granulosum		10.05	0.12		90	
27123003	**Good identification**					**27123003**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27123020	**Low discrimination**					**27123020**
					CAT	
Actinomyces naeslundii		62.24	0.38	RAF 92	25	
Clostridium perfringens		24.08	0.08	FRU 95	1	
Propionibacterium granulosum		13.67	0.12		90	
27123021	**Good identification**					**27123021**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27123022	**Low discrimination**					**27123022**
Actinomyces naeslundii		84.60	0.38	RAF 92		
Clostridium perfringens		15.41	0.03	FRU 95		
27123023	**Good identification**					**27123023**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27123100	**Very good identification**					**27123100**
					SPO	
Propionibacterium propionicum		95.28	0.58	FRU 99	1	
Actinomyces naeslundii		4.25	0.45		1	
Clostridium chauvoei		0.47	0.06		99	
27123101	**Low discrimination**					**27123101**
					SPO	
Clostridium chauvoei		73.32	0.40	TRE 10	99	
Propionibacterium propionicum		16.65	0.42	MAN 1	1	
Actinomyces naeslundii		10.03	0.49		1	
27123102	**Very good identification**					**27123102**
Propionibacterium propionicum		96.94	0.61	FRU 99		
Actinomyces naeslundii		2.88	0.45			
Actinomyces viscosus		0.18	0.01			
27123103	**Low discrimination**					**27123103**
					SPO	
Propionibacterium propionicum		52.10	0.45	FRU 99	1	
Clostridium chauvoei		26.99	0.27	URE 10	99	
Actinomyces naeslundii		20.92	0.49		1	
27123120	**Low discrimination**					**27123120**
Actinomyces naeslundii		44.98	0.45	RAF 92		
Clostridium perfringens		44.84	0.23	FRU 95		
Propionibacterium propionicum		10.18	0.24			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27123122	**Low discrimination**						**27123122**
Actinomyces naeslundii		55.29	0.45	RAF 92			
Clostridium perfringens		25.94	0.18	FRU 95			
Propionibacterium propionicum		18.77	0.27				
27123123	**Low discrimination**						**27123123**
					SPO		
Actinomyces naeslundii		69.93	0.49	RAF 92	1		
Clostridium chauvoei		30.07	0.19	FRU 95	99		
27123140	**Very good identification**						**27123140**
Propionibacterium propionicum		99.22	0.58	FRU 99			
Actinomyces naeslundii		0.78	0.33				
27123141	**Low discrimination**						**27123141**
					SPO		
Propionibacterium propionicum		86.90	0.42	FRU 99	1		
Actinomyces naeslundii		9.24	0.36	URE 10	1		
Clostridium chauvoei		3.87	0.06		99		
27123142	**Very good identification**						**27123142**
Propionibacterium propionicum		99.48	0.61	FRU 99			
Actinomyces naeslundii		0.52	0.33				
27123143	**Acceptable identification**						**27123143**
Propionibacterium propionicum		93.38	0.45	FRU 99			
Actinomyces naeslundii		6.62	0.36	URE 10			
27123161	**Good identification**						**27123161**
Actinomyces naeslundii		100.0	0.36	RAF 92			
27123163	**Good identification**						**27123163**
Actinomyces naeslundii		100.0	0.36	RAF 92			
27125101	**Good identification**						**27125101**
Clostridium chauvoei		100.0	0.40	TRE 10			
27130000	**Low discrimination**						**27130000**
					SPO	CAT	
Propionibacterium granulosum		71.92	0.58	NIT 1	1	90	
Clostridium botulinum biovar B		11.31	0.32		99	1	
Actinomyces odontolyticus		7.34	0.39		1	10	
27130002	**Low discrimination**						**27130002**
					CAT		
Actinomyces viscosus		88.89	0.56	RAF 94	76		
Propionibacterium granulosum		8.39	0.48	TRE 7	90		
Actinomyces odontolyticus		2.10	0.35		10		
27130003	**Good identification**						**27130003**
Actinomyces viscosus		99.00	0.48	RAF 94			
Actinomyces naeslundii		0.60	0.34	TRE 7			
Propionibacterium granulosum		0.28	0.14				
27130010	**Low discrimination**						**27130010**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		63.78	0.43	NIT 1	1	90	
Clostridium botulinum biovar B		36.22	0.26	MLZ 12	99	1	
27130020	**Low discrimination**						**27130020**
					CAT	SPO	
Clostridium perfringens		48.29	0.43	GAL 99	1	84	
Propionibacterium granulosum		36.34	0.48	LAC 88	90	1	
Clostridium botulinum biovar B		11.26	0.26		1	99	
27130022	**Low discrimination**						**27130022**
					CAT		
Clostridium perfringens		60.29	0.37	GAL 99	1		
Propionibacterium granulosum		24.11	0.38	LAC 88	90		
Actinomyces viscosus		10.32	0.22		76		
27130040	**Very good identification**						**27130040**
					SPO	CAT	
Clostridium botulinum biovar B		99.89	0.66	NIT 1	99	1	
Propionibacterium granulosum		0.07	0.24		1	90	
Propionibacterium propionicum		0.04	0.24		1	1	
27130050	**Very good identification**						**27130050**
					SPO	CAT	
Clostridium botulinum biovar B		99.98	0.61	NIT 1	99	1	
Propionibacterium granulosum		0.02	0.09		1	90	
27130060	**Very good identification**						**27130060**
					CAT	SPO	
Clostridium botulinum biovar B		99.15	0.61	NIT 1	1	99	
Clostridium perfringens		0.82	0.30		1	84	
Propionibacterium granulosum		0.03	0.14		90	1	
27130070	**Good identification**						**27130070**
Clostridium botulinum biovar B		100.0	0.55	NIT 1			
27130100	**Low discrimination**						**27130100**
Propionibacterium propionicum		73.74	0.58	MAN 99			
Clostridium perfringens		10.04	0.31				
Actinomyces viscosus		6.34	0.26				
27130101	**Low discrimination**						**27130101**
					SPO		
Clostridium chauvoei		87.23	0.47	TRE 10	99		
Propionibacterium propionicum		8.49	0.42	GAL 99	1		
Actinomyces viscosus		2.19	0.18		1		
27130102	**Low discrimination**						**27130102**
Actinomyces viscosus		83.97	0.60	RAF 94			
Propionibacterium propionicum		14.80	0.61	TRE 7			
Clostridium perfringens		0.63	0.26				
27130103	**Low discrimination**						**27130103**
					SPO		
Actinomyces viscosus		87.77	0.52	RAF 94	1		
Clostridium chauvoei		6.23	0.34	TRE 7	99		
Propionibacterium propionicum		5.16	0.45		1		
27130120	**Very good identification**						**27130120**
Clostridium perfringens		98.96	0.57	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.41	0.39				
Propionibacterium propionicum		0.23	0.24				
27130121	**Low discrimination**						**27130121**
					SPO		
Clostridium chauvoei		84.13	0.38	TRE 10	99		
Clostridium perfringens		9.84	0.23	GAL 99	84		
Actinomyces naeslundii		6.03	0.42		1		
27130122	**Acceptable identification**						**27130122**
Clostridium perfringens		94.31	0.52	GAL 99			
Actinomyces viscosus		3.91	0.26				
Actinomyces naeslundii		0.83	0.39				
27130140	**Low discrimination**						**27130140**
					SPO		
Propionibacterium propionicum		76.14	0.58	MAN 99	1		
Clostridium botulinum biovar B		21.64	0.32		99		
Clostridium perfringens		1.98	0.19		84		
27130141	**Low discrimination**						**27130141**
					SPO		
Propionibacterium propionicum		85.25	0.42	MAN 99	1		
Clostridium chauvoei		8.85	0.12	URE 10	99		
Actinomyces naeslundii		3.70	0.29		1		
27130142	**Good identification**						**27130142**
Propionibacterium propionicum		93.50	0.61	MAN 99			
Actinomyces viscosus		5.36	0.26				
Clostridium perfringens		0.76	0.14				
27130143	**Low discrimination**						**27130143**
Propionibacterium propionicum		82.44	0.45	MAN 99			
Actinomyces viscosus		14.17	0.18	URE 10			
Actinomyces naeslundii		2.38	0.29				
27130160	**Low discrimination**						**27130160**
					SPO		
Clostridium perfringens		84.55	0.45	GAL 99	84		
Clostridium botulinum biovar B		14.11	0.26		99		
Propionibacterium propionicum		1.02	0.24		1		
27130162	**Good identification**						**27130162**
Clostridium perfringens		95.56	0.39	GAL 99			
Propionibacterium propionicum		3.67	0.27				
Actinomyces naeslundii		0.78	0.26				
27130200	**Low discrimination**						**27130200**
					SPO	CAT	
Actinomyces odontolyticus		85.91	0.44	MNS 9	1	10	
Propionibacterium granulosum		13.41	0.32	TRE 1	1	90	
Clostridium botulinum biovar B		0.69	-.03		99	1	
27130202	**Low discrimination**						**27130202**
					CAT		
Actinomyces odontolyticus		77.89	0.40	MNS 9	10		
Actinomyces viscosus		17.15	0.22	TRE 1	76		
Propionibacterium granulosum		4.96	0.22		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27130300	**Acceptable identification**						**27130300**
Actinomyces odontolyticus		91.81	0.35	MNS 9			
Propionibacterium propionicum		8.19	0.24	TRE 1			
27131000	**Low discrimination**						**27131000**
					CAT		
Clostridium perfringens		44.66	0.51	LAC 88	1		
Propionibacterium granulosum		43.91	0.58	NAG 97	90		
Actinomyces naeslundii		3.73	0.55		25		
27131001	**Low discrimination**						**27131001**
					SPO		
Clostridium chauvoei		80.48	0.47	TRE 10	99		
Actinomyces naeslundii		16.23	0.58	LAC 99	1		
Clostridium perfringens		1.31	0.17		84		
27131002	**Low discrimination**						**27131002**
					CAT		
Actinomyces viscosus		64.85	0.52	RAF 94	76		
Clostridium perfringens		19.13	0.45	TRE 7	1		
Propionibacterium granulosum		9.99	0.48		90		
27131003	**Low discrimination**						**27131003**
					SPO		
Actinomyces viscosus		68.73	0.44	RAF 94	1		
Actinomyces naeslundii		16.17	0.58	TRE 7	1		
Clostridium chauvoei		14.16	0.34		99		
27131010	**Low discrimination**						**27131010**
					CAT		
Propionibacterium granulosum		86.56	0.43	NIT 1	90		
Clostridium perfringens		6.52	0.17	MLZ 12	1		
Lactobacillus jensenii		4.08	0.14		1		
27131011	**Low discrimination**						**27131011**
					SPO	CAT	
Actinomyces naeslundii		46.35	0.36	RAF 92	1	25	
Clostridium chauvoei		44.12	0.12	MLZ 5	99	1	
Propionibacterium granulosum		9.53	0.09		1	90	
27131013	**Low discrimination**						**27131013**
					SPO		
Actinomyces naeslundii		50.42	0.36	RAF 92	1		
Actinomyces viscosus		41.11	0.10	MLZ 5	1		
Clostridium chauvoei		8.47	-.01		99		
27131020	**Very good identification**						**27131020**
					CAT		
Clostridium perfringens		98.93	0.77	LAC 88	1		
Propionibacterium granulosum		0.75	0.48		90		
Actinomyces naeslundii		0.26	0.55		25		
27131021	**Low discrimination**						**27131021**
					SPO		
Clostridium perfringens		49.40	0.43	LAC 88	84		
Clostridium chauvoei		31.30	0.38	URE 1	99		
Actinomyces naeslundii		18.93	0.58		1		
27131022	**Very good identification**						**27131022**
					CAT		
Clostridium perfringens		98.89	0.71	LAC 88	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.54	0.55		25	
Propionibacterium granulosum		0.40	0.38		90	
27131023	**Low discrimination**					**27131023**
					SPO	
Clostridium perfringens		47.92	0.37	LAC 88	84	
Actinomyces naeslundii		39.02	0.58	URE 1	1	
Clostridium chauvoei		11.39	0.25		99	
27131030	**Low discrimination**					**27131030**
					CAT	
Clostridium perfringens		89.21	0.43	LAC 88	1	
Propionibacterium granulosum		9.16	0.33	MLZ 1	90	
Actinomyces naeslundii		1.20	0.33		25	
27131031	**Low discrimination**					**27131031**
					SPO	
Actinomyces naeslundii		55.01	0.36	RAF 92	1	
Clostridium perfringens		27.54	0.08	MLZ 5	84	
Clostridium chauvoei		17.45	0.04		99	
27131032	**Acceptable identification**					**27131032**
					CAT	
Clostridium perfringens		92.33	0.37	LAC 88	1	
Propionibacterium granulosum		5.03	0.23	MLZ 1	90	
Actinomyces naeslundii		2.64	0.33		25	
27131033	**Low discrimination**					**27131033**
Actinomyces naeslundii		80.93	0.36	RAF 92		
Clostridium perfringens		19.07	0.03	MLZ 5		
27131040	**Low discrimination**					**27131040**
					SPO	
Clostridium perfringens		50.77	0.39	LAC 88	84	
Clostridium botulinum biovar B		41.20	0.32	NAG 97	99	
Actinomyces naeslundii		3.93	0.42		1	
27131041	**Low discrimination**					**27131041**
					SPO	
Actinomyces naeslundii		69.36	0.45	RAF 92	1	
Clostridium chauvoei		19.69	0.12		99	
Clostridium perfringens		6.04	0.04		84	
27131043	**Low discrimination**					**27131043**
					SPO	
Actinomyces naeslundii		77.32	0.45	RAF 92	1	
Actinomyces viscosus		18.81	0.10		1	
Clostridium chauvoei		3.87	-.01		99	
27131060	**Very good identification**					**27131060**
					SPO	
Clostridium perfringens		98.51	0.64	LAC 88	84	
Clostridium botulinum biovar B		1.22	0.26		99	
Actinomyces naeslundii		0.24	0.42		1	
27131062	**Very good identification**					**27131062**
Clostridium perfringens		99.49	0.59	LAC 88		
Actinomyces naeslundii		0.51	0.42			
27131100	**Low discrimination**					**27131100**
					COC SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		78.23	0.66	NAG 97	1	84	
Clostridium chauvoei		6.62	0.47		1	99	
Atopobium parvulum		6.02	0.46		99	1	
27131101	**Very good identification**						**27131101**
					SPO		
Clostridium chauvoei		99.18	0.81	TRE 10	99		
Actinomyces naeslundii		0.58	0.65		1		
Clostridium perfringens		0.12	0.31		84		
27131102	**Low discrimination**						**27131102**
Clostridium perfringens		45.58	0.60	NAG 97			
Actinomyces viscosus		37.47	0.57				
Propionibacterium propionicum		10.77	0.61				
27131103	**Low discrimination**						**27131103**
					SPO		
Clostridium chauvoei		88.37	0.68	TRE 10	99		
Actinomyces viscosus		7.70	0.49		1		
Actinomyces naeslundii		2.90	0.65		1		
27131111	**Good identification**						**27131111**
					SPO		
Clostridium chauvoei		97.07	0.47	TRE 10	99		
Actinomyces naeslundii		2.93	0.44	MLZ 1	1		
27131120	**Excellent identification**						**27131120**
					SPO		
Clostridium perfringens		99.75	0.91		84		
Actinomyces naeslundii		0.10	0.62		1		
Clostridium chauvoei		0.09	0.38		99		
27131121	**Low discrimination**						**27131121**
					SPO		
Clostridium chauvoei		88.16	0.73	TRE 10	99		
Clostridium perfringens		10.31	0.57		84		
Actinomyces naeslundii		1.53	0.65		1		
27131122	**Very good identification**						**27131122**
					SPO		
Clostridium perfringens		99.69	0.86		84		
Actinomyces naeslundii		0.21	0.62		1		
Clostridium chauvoei		0.03	0.25		99		
27131123	**Low discrimination**						**27131123**
					SPO		
Clostridium chauvoei		70.78	0.60	TRE 10	99		
Clostridium perfringens		22.07	0.52		84		
Actinomyces naeslundii		6.97	0.65		1		
27131130	**Very good identification**						**27131130**
					SPO		
Clostridium perfringens		99.40	0.57	MLZ 1	84		
Actinomyces naeslundii		0.52	0.41		1		
Clostridium chauvoei		0.09	0.04		99		
27131131	**Low discrimination**						**27131131**
					SPO		
Clostridium chauvoei		82.82	0.38	TRE 10	99		
Clostridium perfringens		9.68	0.23	MLZ 1	84		
Actinomyces naeslundii		7.50	0.44		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27131132	**Good identification**						**27131132**
					GRA	CAT	
Clostridium perfringens		98.78	0.52	MLZ 1	99	1	
Actinomyces naeslundii		1.10	0.41		99	25	
Capnocytophaga ochracea		0.13	-.01		1	90	
27131140	**Low discrimination**						**27131140**
Clostridium perfringens		69.98	0.53	NAG 97			
Propionibacterium propionicum		27.24	0.58				
Actinomyces naeslundii		2.10	0.50				
27131141	**Low discrimination**						**27131141**
					SPO		
Clostridium chauvoei		81.88	0.47	TRE 10	99		
Actinomyces naeslundii		8.29	0.53	SOR 1	1		
Propionibacterium propionicum		7.97	0.42		1		
27131142	**Low discrimination**						**27131142**
Propionibacterium propionicum		52.81	0.61	MAN 99			
Clostridium perfringens		42.55	0.48				
Actinomyces naeslundii		2.71	0.50				
27131160	**Very good identification**						**27131160**
Clostridium perfringens		99.89	0.79				
Actinomyces naeslundii		0.09	0.50				
Propionibacterium propionicum		0.01	0.24				
27131161	**Low discrimination**						**27131161**
					SPO		
Clostridium perfringens		62.84	0.45	URE 1	84		
Clostridium chauvoei		28.50	0.38		99		
Actinomyces naeslundii		8.66	0.53		1		
27131162	**Very good identification**						**27131162**
Clostridium perfringens		99.76	0.74				
Actinomyces naeslundii		0.20	0.50				
Propionibacterium propionicum		0.04	0.27				
27131163	**Low discrimination**						**27131163**
					SPO		
Clostridium perfringens		68.36	0.39	URE 1	84		
Actinomyces naeslundii		20.02	0.53		1		
Clostridium chauvoei		11.63	0.25		99		
27131170	**Good identification**						**27131170**
Clostridium perfringens		99.52	0.45	MLZ 1			
Actinomyces naeslundii		0.48	0.28				
27131172	**Good identification**						**27131172**
Clostridium perfringens		98.98	0.39	MLZ 1			
Actinomyces naeslundii		1.02	0.28				
27131200	**Low discrimination**						**27131200**
					CAT		
Actinomyces odontolyticus		76.23	0.41	MNS 9	10		
Propionibacterium granulosum		17.84	0.32	TRE 1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		5.93	0.17		1	
27131202	**Low discrimination**					**27131202**
					CAT	
Actinomyces odontolyticus		73.66	0.37	MNS 9	10	
Actinomyces viscosus		14.91	0.18	TRE 1	76	
Propionibacterium granulosum		7.03	0.22		90	
27131220	**Acceptable identification**					**27131220**
					CAT	
Clostridium perfringens		91.91	0.43	LAC 88	1	
Actinomyces odontolyticus		5.95	0.27	RHA 1	10	
Propionibacterium granulosum		2.14	0.22		90	
27131222	**Acceptable identification**					**27131222**
					CAT	
Clostridium perfringens		91.18	0.37	LAC 88	1	
Actinomyces odontolyticus		7.69	0.24	RHA 1	10	
Propionibacterium granulosum		1.13	0.12		90	
27131301	**Good identification**					**27131301**
					SPO	
Clostridium chauvoei		99.42	0.47	TRE 10	99	
Actinomyces naeslundii		0.58	0.31	RHA 1	1	
27131320	**Very good identification**					**27131320**
Clostridium perfringens		99.54	0.57	RHA 1		
Actinomyces odontolyticus		0.28	0.19			
Actinomyces naeslundii		0.10	0.28			
27131321	**Low discrimination**					**27131321**
					SPO	
Clostridium chauvoei		88.16	0.38	TRE 10	99	
Clostridium perfringens		10.31	0.23	RHA 1	84	
Actinomyces naeslundii		1.53	0.31		1	
27131322	**Good identification**					**27131322**
Clostridium perfringens		99.43	0.52	RHA 1		
Actinomyces odontolyticus		0.36	0.15			
Actinomyces naeslundii		0.21	0.28			
27131360	**Good identification**					**27131360**
Clostridium perfringens		100.0	0.45	RHA 1		
27131362	**Good identification**					**27131362**
Clostridium perfringens		100.0	0.39	RHA 1		
27131420	**Good identification**					**27131420**
					CAT	
Clostridium perfringens		99.25	0.43	LAC 88	1	
Propionibacterium granulosum		0.76	0.14	ARA 1	90	
27131422	**Good identification**					**27131422**
Clostridium perfringens		100.0	0.37	LAC 88		
27131501	**Good identification**					**27131501**
					SPO	
Clostridium chauvoei		99.42	0.47	TRE 10	99	
Actinomyces naeslundii		0.58	0.31	ARA 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27131520	**Very good identification**					**27131520**
Clostridium perfringens		99.58	0.57	ARA 1		
Bifidobacterium longum subsp. longum		0.24	0.18			
Actinomyces naeslundii		0.10	0.28			
27131521	**Low discrimination**					**27131521**
					SPO	
Clostridium chauvoei		88.16	0.38	TRE 10	99	
Clostridium perfringens		10.31	0.23	ARA 1	84	
Actinomyces naeslundii		1.53	0.31		1	
27131522	**Good identification**					**27131522**
Clostridium perfringens		99.09	0.52	ARA 1		
Bifidobacterium longum subsp. longum		0.50	0.18			
Actinomyces naeslundii		0.21	0.28			
27131560	**Good identification**					**27131560**
Clostridium perfringens		100.0	0.45	ARA 1		
27131562	**Good identification**					**27131562**
Clostridium perfringens		100.0	0.39	ARA 1		
27132000	**Low discrimination**					**27132000**
					CAT	
Propionibacterium granulosum		57.33	0.56	NIT 1	90	
Propionibacterium propionicum		42.03	0.58		1	
Actinomyces naeslundii		0.52	0.36		25	
27132001	**Low discrimination**					**27132001**
					CAT	
Propionibacterium propionicum		77.43	0.42	LAC 99	1	
Actinomyces naeslundii		12.97	0.39	URE 10	25	
Propionibacterium granulosum		9.60	0.22		90	
27132002	**Low discrimination**					**27132002**
					CAT	
Propionibacterium propionicum		78.88	0.61	LAC 99	1	
Propionibacterium granulosum		17.93	0.46		90	
Actinomyces viscosus		2.54	0.22		76	
27132003	**Low discrimination**					**27132003**
Propionibacterium propionicum		81.36	0.45	LAC 99		
Actinomyces naeslundii		9.09	0.39	URE 10		
Actinomyces viscosus		7.87	0.14			
27132010	**Acceptable identification**					**27132010**
					CAT	
Propionibacterium granulosum		94.85	0.41	NIT 1	90	
Propionibacterium propionicum		5.15	0.24	MLZ 12	1	
27132020	**Acceptable identification**					**27132020**
					CAT	
Propionibacterium granulosum		92.27	0.46	NIT 1	90	
Actinomyces naeslundii		3.36	0.36		25	
Propionibacterium propionicum		2.73	0.24		1	
27132021	**Low discrimination**					**27132021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		84.39	0.39	RAF 92			
Propionibacterium granulosum		15.61	0.12	GAL 96			
27132022	**Low discrimination**						**27132022**
					CAT		
Propionibacterium granulosum		73.70	0.35	NIT 1	90		
Propionibacterium propionicum		13.10	0.27		1		
Actinomyces naeslundii		10.73	0.36		25		
27132023	**Good identification**						**27132023**
Actinomyces naeslundii		100.0	0.39	RAF 92			
27132040	**Low discrimination**						**27132040**
					SPO	CAT	
Propionibacterium propionicum		76.91	0.58	LAC 99	1	1	
Clostridium botulinum biovar B		21.86	0.32		99	1	
Propionibacterium granulosum		1.06	0.22		1	90	
27132041	**Acceptable identification**						**27132041**
					SPO		
Propionibacterium propionicum		94.75	0.42	LAC 99	1		
Actinomyces naeslundii		2.80	0.26	URE 10	1		
Clostridium botulinum biovar B		2.45	-.03		99		
27132042	**Very good identification**						**27132042**
					SPO	CAT	
Propionibacterium propionicum		99.44	0.61	LAC 99	1	1	
Propionibacterium granulosum		0.23	0.12		1	90	
Clostridium botulinum biovar B		0.19	-.03		99	1	
27132043	**Good identification**						**27132043**
Propionibacterium propionicum		98.07	0.45	LAC 99			
Actinomyces naeslundii		1.93	0.26	URE 10			
27132100	**Excellent identification**						**27132100**
					CAT		
Propionibacterium propionicum		99.95	0.93		1		
Actinomyces naeslundii		0.04	0.44		25		
Propionibacterium granulosum		0.01	0.22		90		
27132101	**Very good identification**						**27132101**
					SPO		
Propionibacterium propionicum		99.42	0.76	URE 10	1		
Actinomyces naeslundii		0.48	0.47		1		
Clostridium chauvoei		0.10	0.12		99		
27132102	**Excellent identification**						**27132102**
Propionibacterium propionicum		99.92	0.96				
Actinomyces viscosus		0.06	0.26				
Actinomyces naeslundii		0.02	0.44				
27132103	**Very good identification**						**27132103**
Propionibacterium propionicum		99.50	0.79	URE 10			
Actinomyces naeslundii		0.32	0.47				
Actinomyces viscosus		0.17	0.18				
27132110	**Very good identification**						**27132110**
Propionibacterium propionicum		100.0	0.58	MLZ 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27132111	**Good identification**					**27132111**
Propionibacterium propionicum		97.55	0.42	MLZ 1		
Actinomyces naeslundii		2.45	0.25	URE 10		
27132112	**Very good identification**					**27132112**
Propionibacterium propionicum		100.0	0.61	MLZ 1		
27132113	**Good identification**					**27132113**
Propionibacterium propionicum		98.35	0.45	MLZ 1		
Actinomyces naeslundii		1.65	0.25	URE 10		
27132120	**Good identification**					**27132120**
Propionibacterium propionicum		92.31	0.58	NAG 1		
Clostridium perfringens		4.11	0.23			
Actinomyces naeslundii		3.26	0.44			
27132121	**Low discrimination**					**27132121**
					SPO	
Propionibacterium propionicum		66.16	0.42	NAG 1	1	
Actinomyces naeslundii		31.55	0.47	URE 10	1	
Clostridium chauvoei		2.29	0.04		99	
27132122	**Very good identification**					**27132122**
Propionibacterium propionicum		96.39	0.61	NAG 1		
Actinomyces naeslundii		2.27	0.44			
Clostridium perfringens		1.35	0.18			
27132123	**Low discrimination**					**27132123**
Propionibacterium propionicum		75.88	0.45	NAG 1		
Actinomyces naeslundii		24.12	0.47	URE 10		
27132140	**Excellent identification**					**27132140**
					SPO	
Propionibacterium propionicum		99.99	0.93		1	
Actinomyces naeslundii		0.01	0.31		1	
Clostridium botulinum biovar B		0.00	-.03		99	
27132141	**Very good identification**					**27132141**
Propionibacterium propionicum		99.92	0.76	URE 10		
Actinomyces naeslundii		0.09	0.34			
27132142	**Excellent identification**					**27132142**
Propionibacterium propionicum		100.0	0.96			
Actinomyces naeslundii		0.00	0.31			
27132143	**Very good identification**					**27132143**
Propionibacterium propionicum		99.94	0.79	URE 10		
Actinomyces naeslundii		0.06	0.34			
27132150	**Very good identification**					**27132150**
Propionibacterium propionicum		100.0	0.58	MLZ 1		
27132151	**Good identification**					**27132151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.42	MLZ 1		
27132152	**Very good identification**					**27132152**
Propionibacterium propionicum		100.0	0.61	MLZ 1		
27132153	**Good identification**					**27132153**
Propionibacterium propionicum		100.0	0.45	MLZ 1		
27132160	**Very good identification**					**27132160**
Propionibacterium propionicum		98.55	0.58	NAG 1		
Clostridium perfringens		0.84	0.11			
Actinomyces naeslundii		0.61	0.31			
27132161	**Acceptable identification**					**27132161**
Propionibacterium propionicum		92.24	0.42	NAG 1		
Actinomyces naeslundii		7.76	0.34	URE 10		
27132162	**Very good identification**					**27132162**
Propionibacterium propionicum		99.32	0.61	NAG 1		
Actinomyces naeslundii		0.41	0.31			
Clostridium perfringens		0.26	0.05			
27132163	**Acceptable identification**					**27132163**
Propionibacterium propionicum		94.69	0.45	NAG 1		
Actinomyces naeslundii		5.31	0.34	URE 10		
27132300	**Very good identification**					**27132300**
Propionibacterium propionicum		100.0	0.58	RHA 1		
27132301	**Good identification**					**27132301**
Propionibacterium propionicum		100.0	0.42	RHA 1		
27132302	**Very good identification**					**27132302**
Propionibacterium propionicum		100.0	0.61	RHA 1		
27132303	**Good identification**					**27132303**
Propionibacterium propionicum		100.0	0.45	RHA 1		
27132340	**Very good identification**					**27132340**
Propionibacterium propionicum		100.0	0.58	RHA 1		
27132341	**Good identification**					**27132341**
Propionibacterium propionicum		100.0	0.42	RHA 1		
27132342	**Very good identification**					**27132342**
Propionibacterium propionicum		100.0	0.61	RHA 1		
27132343	**Good identification**					**27132343**
Propionibacterium propionicum		100.0	0.45	RHA 1		
27132500	**Very good identification**					**27132500**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.58	ARA 1		
27132501	**Good identification**					**27132501**
Propionibacterium propionicum		100.0	0.42	ARA 1		
27132502	**Very good identification**					**27132502**
Propionibacterium propionicum		100.0	0.61	ARA 1		
27132503	**Good identification**					**27132503**
Propionibacterium propionicum		100.0	0.45	ARA 1		
27132540	**Very good identification**					**27132540**
Propionibacterium propionicum		100.0	0.58	ARA 1		
27132541	**Good identification**					**27132541**
Propionibacterium propionicum		100.0	0.42	ARA 1		
27132542	**Very good identification**					**27132542**
Propionibacterium propionicum		100.0	0.61	ARA 1		
27132543	**Good identification**					**27132543**
Propionibacterium propionicum		100.0	0.45	ARA 1		
27133000	**Low discrimination**					**27133000**
					CAT	
Propionibacterium granulosum		50.89	0.56	NIT 1	90	
Propionibacterium propionicum		37.31	0.58		1	
Actinomyces naeslundii		11.11	0.60		25	
27133001	**Low discrimination**					**27133001**
					CAT	
Actinomyces naeslundii		76.61	0.63	RAF 92	25	
Propionibacterium propionicum		19.05	0.42		1	
Propionibacterium granulosum		2.36	0.22		90	
27133002	**Low discrimination**					**27133002**
					CAT	
Propionibacterium propionicum		68.82	0.61	LAC 99	1	
Propionibacterium granulosum		15.64	0.46		90	
Actinomyces naeslundii		13.67	0.60		25	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27133003	**Low discrimination**					**27133003**
Actinomyces naeslundii		71.05	0.63	RAF 92		
Propionibacterium propionicum		26.51	0.45			
Actinomyces viscosus		1.57	0.10			
27133010	**Low discrimination**					**27133010**
					CAT	
Propionibacterium granulosum		87.83	0.41	NIT 1	90	
Actinomyces naeslundii		7.40	0.38	MLZ 12	25	
Propionibacterium propionicum		4.77	0.24		1	
27133011	**Good identification**					**27133011**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27133013	**Good identification**					**27133013**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27133020	**Low discrimination**					**27133020**
					CAT	
Clostridium perfringens		48.07	0.43	MAN 1	1	
Propionibacterium granulosum		27.29	0.46	LAC 88	90	
Actinomyces naeslundii		23.84	0.60		25	
27133021	**Very good identification**					**27133021**
					SPO	
Actinomyces naeslundii		97.10	0.63	RAF 92	1	
Clostridium perfringens		1.32	0.08		84	
Clostridium chauvoei		0.84	0.04		99	
27133022	**Low discrimination**					**27133022**
					CAT	
Actinomyces naeslundii		43.75	0.60	RAF 92	25	
Clostridium perfringens		41.51	0.37		1	
Propionibacterium granulosum		12.52	0.35		90	
27133023	**Very good identification**					**27133023**
Actinomyces naeslundii		99.37	0.63	RAF 92		
Clostridium perfringens		0.64	0.03			
27133031	**Good identification**					**27133031**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27133032	**Low discrimination**					**27133032**
					CAT	
Actinomyces naeslundii		51.98	0.38	RAF 92	25	
Propionibacterium granulosum		38.55	0.21	MLZ 5	90	
Clostridium perfringens		9.47	0.03		1	
27133033	**Good identification**					**27133033**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27133040	**Good identification**					**27133040**
					CAT	
Propionibacterium propionicum		93.22	0.58	LAC 99	1	
Actinomyces naeslundii		4.90	0.47		25	
Propionibacterium granulosum		1.28	0.22		90	
27133041	**Low discrimination**					**27133041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		58.49	0.42	LAC 99		
Actinomyces naeslundii		41.51	0.50	URE 10		
27133042	**Very good identification**					**27133042**
					CAT	
Propionibacterium propionicum		96.40	0.61	LAC 99	1	
Actinomyces naeslundii		3.38	0.47		25	
Propionibacterium granulosum		0.22	0.12		90	
27133043	**Low discrimination**					**27133043**
Propionibacterium propionicum		67.89	0.45	LAC 99		
Actinomyces naeslundii		32.11	0.50	URE 10		
27133061	**Good identification**					**27133061**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27133063	**Good identification**					**27133063**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27133100	**Very good identification**					**27133100**
Propionibacterium propionicum		98.97	0.93			
Actinomyces naeslundii		0.85	0.67			
Clostridium perfringens		0.14	0.31			
27133101	**Low discrimination**					**27133101**
					SPO	
Propionibacterium propionicum		82.01	0.76	URE 10	1	
Actinomyces naeslundii		9.48	0.70		1	
Clostridium chauvoei		8.51	0.47		99	
27133102	**Excellent identification**					**27133102**
Propionibacterium propionicum		99.35	0.96			
Actinomyces naeslundii		0.57	0.67			
Clostridium perfringens		0.04	0.26			
27133103	**Good identification**					**27133103**
					SPO	
Propionibacterium propionicum		91.72	0.79	URE 10	1	
Actinomyces naeslundii		7.07	0.70		1	
Clostridium chauvoei		1.12	0.34		99	
27133110	**Very good identification**					**27133110**
Propionibacterium propionicum		95.73	0.58	MLZ 1		
Actinomyces naeslundii		4.27	0.45			
27133111	**Low discrimination**					**27133111**
					SPO	
Propionibacterium propionicum		58.61	0.42	MLZ 1	1	
Actinomyces naeslundii		35.31	0.49	URE 10	1	
Clostridium chauvoei		6.08	0.12		99	
27133112	**Very good identification**					**27133112**
Propionibacterium propionicum		97.11	0.61	MLZ 1		
Actinomyces naeslundii		2.89	0.45			
27133113	**Low discrimination**					**27133113**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Propionibacterium propionicum		70.73	0.45	MLZ 1	1	
Actinomyces naeslundii		28.40	0.49	URE 10	1	
Clostridium chauvoei		0.86	-.01		99	
27133120	**Low discrimination**					**27133120**
Clostridium perfringens		70.36	0.57	MAN 1		
Propionibacterium propionicum		15.98	0.58			
Actinomyces naeslundii		13.54	0.67			
27133121	**Low discrimination**					**27133121**
					SPO	
Actinomyces naeslundii		70.34	0.70	RAF 92	1	
Clostridium chauvoei		21.05	0.38		99	
Propionibacterium propionicum		6.15	0.42		1	
27133122	**Low discrimination**					**27133122**
Clostridium perfringens		46.89	0.52	MAN 1		
Propionibacterium propionicum		33.93	0.61			
Actinomyces naeslundii		19.18	0.67			
27133123	**Low discrimination**					**27133123**
					SPO	
Actinomyces naeslundii		83.31	0.70	RAF 92	1	
Propionibacterium propionicum		10.92	0.45		1	
Clostridium chauvoei		4.40	0.25		99	
27133130	**Low discrimination**					**27133130**
Actinomyces naeslundii		44.98	0.45	RAF 92		
Clostridium perfringens		44.84	0.23	MLZ 5		
Propionibacterium propionicum		10.18	0.24			
27133131	**Acceptable identification**					**27133131**
					SPO	
Actinomyces naeslundii		94.57	0.49	RAF 92	1	
Clostridium chauvoei		5.43	0.04	MLZ 5	99	
27133132	**Low discrimination**					**27133132**
Actinomyces naeslundii		55.29	0.45	RAF 92		
Clostridium perfringens		25.94	0.18	MLZ 5		
Propionibacterium propionicum		18.77	0.27			
27133133	**Good identification**					**27133133**
Actinomyces naeslundii		100.0	0.49	RAF 92		
27133140	**Excellent identification**					**27133140**
Propionibacterium propionicum		99.82	0.93			
Actinomyces naeslundii		0.15	0.54			
Clostridium perfringens		0.03	0.19			
27133141	**Very good identification**					**27133141**
					SPO	
Propionibacterium propionicum		97.90	0.76	URE 10	1	
Actinomyces naeslundii		2.00	0.57		1	
Clostridium chauvoei		0.10	0.12		99	
27133142	**Excellent identification**					**27133142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.89	0.96			
Actinomyces naeslundii		0.10	0.54			
Clostridium perfringens		0.01	0.14			
27133143	**Very good identification**					**27133143**
					SPO	
Propionibacterium propionicum		98.65	0.79	URE 10	1	
Actinomyces naeslundii		1.34	0.57		1	
Clostridium chauvoei		0.01	-.01		99	
27133150	**Very good identification**					**27133150**
Propionibacterium propionicum		99.22	0.58	MLZ 1		
Actinomyces naeslundii		0.78	0.33			
27133151	**Acceptable identification**					**27133151**
Propionibacterium propionicum		90.39	0.42	MLZ 1		
Actinomyces naeslundii		9.61	0.36	URE 10		
27133152	**Very good identification**					**27133152**
Propionibacterium propionicum		99.48	0.61	MLZ 1		
Actinomyces naeslundii		0.52	0.33			
27133153	**Acceptable identification**					**27133153**
Propionibacterium propionicum		93.38	0.45	MLZ 1		
Actinomyces naeslundii		6.62	0.36	URE 10		
27133160	**Low discrimination**					**27133160**
Propionibacterium propionicum		50.29	0.58	NAG 1		
Clostridium perfringens		42.19	0.45			
Actinomyces naeslundii		7.52	0.54			
27133161	**Low discrimination**					**27133161**
Actinomyces naeslundii		64.51	0.57	RAF 92		
Propionibacterium propionicum		31.94	0.42			
Clostridium perfringens		2.44	0.11			
27133162	**Low discrimination**					**27133162**
Propionibacterium propionicum		73.37	0.61	NAG 1		
Clostridium perfringens		19.31	0.39			
Actinomyces naeslundii		7.32	0.54			
27133163	**Low discrimination**					**27133163**
Actinomyces naeslundii		56.80	0.57	RAF 92		
Propionibacterium propionicum		42.19	0.45			
Clostridium perfringens		1.01	0.05			
27133171	**Good identification**					**27133171**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27133173	**Good identification**					**27133173**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27133300	**Very good identification**					**27133300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.15	0.58	RHA 1		
Actinomyces naeslundii		0.85	0.33			
27133301	**Low discrimination**					**27133301**
					SPO	
Propionibacterium propionicum		82.01	0.42	RHA 1	1	
Actinomyces naeslundii		9.48	0.36	URE 10	1	
Clostridium chauvoei		8.51	0.12		99	
27133302	**Very good identification**					**27133302**
Propionibacterium propionicum		99.43	0.61	RHA 1		
Actinomyces naeslundii		0.57	0.33			
27133303	**Acceptable identification**					**27133303**
					SPO	
Propionibacterium propionicum		91.80	0.45	RHA 1	1	
Actinomyces naeslundii		7.08	0.36	URE 10	1	
Clostridium chauvoei		1.12	-.01		99	
27133321	**Low discrimination**					**27133321**
					SPO	
Actinomyces naeslundii		76.97	0.36	RAF 92	1	
Clostridium chauvoei		23.03	0.04	RHA 1	99	
27133323	**Good identification**					**27133323**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27133340	**Very good identification**					**27133340**
Propionibacterium propionicum		100.0	0.58	RHA 1		
27133341	**Good identification**					**27133341**
Propionibacterium propionicum		98.00	0.42	RHA 1		
Actinomyces naeslundii		2.00	0.23	URE 10		
27133342	**Very good identification**					**27133342**
Propionibacterium propionicum		100.0	0.61	RHA 1		
27133343	**Good identification**					**27133343**
Propionibacterium propionicum		98.66	0.45	RHA 1		
Actinomyces naeslundii		1.34	0.23	URE 10		
27133500	**Very good identification**					**27133500**
Propionibacterium propionicum		98.64	0.58	ARA 1		
Actinomyces naeslundii		0.85	0.33			
Actinomyces israelii		0.51	0.20			
27133501	**Low discrimination**					**27133501**
					SPO	
Propionibacterium propionicum		82.01	0.42	ARA 1	1	
Actinomyces naeslundii		9.48	0.36	URE 10	1	
Clostridium chauvoei		8.51	0.12		99	
27133502	**Very good identification**					**27133502**
Propionibacterium propionicum		98.92	0.61	ARA 1		
Actinomyces naeslundii		0.57	0.33			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.51	0.23				
27133503	**Acceptable identification**						**27133503**
					SPO		
Propionibacterium propionicum		91.80	0.45	ARA 1	1		
Actinomyces naeslundii		7.08	0.36	URE 10	1		
Clostridium chauvoei		1.12	-.01		99		
27133521	**Low discrimination**						**27133521**
					SPO		
Actinomyces naeslundii		76.97	0.36	RAF 92	1		
Clostridium chauvoei		23.03	0.04	ARA 1	99		
27133523	**Good identification**						**27133523**
Actinomyces naeslundii		100.0	0.36	RAF 92			
27133540	**Very good identification**						**27133540**
Propionibacterium propionicum		99.80	0.58	ARA 1			
Actinomyces israelii		0.20	0.13				
27133541	**Good identification**						**27133541**
Propionibacterium propionicum		98.00	0.42	ARA 1			
Actinomyces naeslundii		2.00	0.23	URE 10			
27133542	**Very good identification**						**27133542**
Propionibacterium propionicum		99.80	0.61	ARA 1			
Actinomyces israelii		0.20	0.16				
27133543	**Good identification**						**27133543**
Propionibacterium propionicum		98.66	0.45	ARA 1			
Actinomyces naeslundii		1.34	0.23	URE 10			
27134200	**Good identification**						**27134200**
Actinomyces odontolyticus		100.0	0.38	MNS 9			
27135020	**Good identification**						**27135020**
					CAT		
Clostridium perfringens		97.11	0.43	XYL 1	1		
Actinomyces odontolyticus		1.60	0.17	LAC 88	10		
Propionibacterium granulosum		0.74	0.14		90		
27135022	**Good identification**						**27135022**
Clostridium perfringens		97.91	0.37	XYL 1			
Actinomyces odontolyticus		2.10	0.13	LAC 88			
27135100	**Acceptable identification**						**27135100**
					COC		
Atopobium parvulum		90.71	0.50	NIT 1	99		
Clostridium perfringens		6.41	0.31		1		
Actinomyces odontolyticus		0.90	0.22		1		
27135101	**Good identification**						**27135101**
					COC	SPO	
Clostridium chauvoei		97.77	0.47	TRE 10	1	99	
Atopobium parvulum		1.67	0.16	XYL 1	99	1	
Actinomyces naeslundii		0.57	0.31		1	1	
27135102	**Low discrimination**						**27135102**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC			
Atopobium parvulum		66.34	0.38	NIT 1	99			
Clostridium perfringens		12.50	0.26		1			
Actinomyces viscosus		10.27	0.23		1			
27135120	**Good identification**							**27135120**
					COC			
Clostridium perfringens		92.61	0.57	XYL 1	1			
Atopobium parvulum		7.16	0.38		99			
Actinomyces naeslundii		0.09	0.28		1			
27135121	**Low discrimination**							**27135121**
					SPO			
Clostridium chauvoei		87.46	0.38	TRE 10	99			
Clostridium perfringens		10.23	0.23	XYL 1	84			
Actinomyces naeslundii		1.52	0.31		1			
27135122	**Good identification**							**27135122**
					COC	GRA	CAT	
Clostridium perfringens		96.48	0.52	XYL 1	1	99	1	
Atopobium parvulum		2.80	0.25		99	99	1	
Capnocytophaga ochracea		0.28	0.05		1	1	90	
27135160	**Good identification**							**27135160**
					COC			
Clostridium perfringens		99.59	0.45	XYL 1	1			
Atopobium parvulum		0.41	0.03		99			
27135162	**Good identification**							**27135162**
Clostridium perfringens		100.0	0.39	XYL 1				
27135200	**Good identification**							**27135200**
Actinomyces odontolyticus		100.0	0.35	MNS 9				
27136100	**Very good identification**							**27136100**
Propionibacterium propionicum		100.0	0.58	XYL 1				
27136101	**Good identification**							**27136101**
Propionibacterium propionicum		100.0	0.42	XYL 1				
27136102	**Very good identification**							**27136102**
Propionibacterium propionicum		100.0	0.61	XYL 1				
27136103	**Good identification**							**27136103**
Propionibacterium propionicum		100.0	0.45	XYL 1				
27136140	**Very good identification**							**27136140**
Propionibacterium propionicum		100.0	0.58	XYL 1				
27136141	**Good identification**							**27136141**
Propionibacterium propionicum		100.0	0.42	XYL 1				
27136142	**Very good identification**							**27136142**
Propionibacterium propionicum		100.0	0.61	XYL 1				
27136143	**Good identification**							**27136143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.45	XYL 1		
27137100	**Good identification**					**27137100**
					COC	
Propionibacterium propionicum		90.83	0.58	XYL 1	1	
Actinomyces israelii		6.63	0.40		1	
Atopobium parvulum		1.77	0.16		99	
27137101	**Low discrimination**					**27137101**
					SPO	
Propionibacterium propionicum		82.01	0.42	XYL 1	1	
Actinomyces naeslundii		9.48	0.36	URE 10	1	
Clostridium chauvoei		8.51	0.12		99	
27137102	**Good identification**					**27137102**
Propionibacterium propionicum		92.51	0.61	XYL 1		
Actinomyces israelii		6.75	0.43			
Actinomyces naeslundii		0.53	0.33			
27137103	**Acceptable identification**					**27137103**
					SPO	
Propionibacterium propionicum		91.80	0.45	XYL 1	1	
Actinomyces naeslundii		7.08	0.36	URE 10	1	
Clostridium chauvoei		1.12	-.01		99	
27137121	**Low discrimination**					**27137121**
					SPO	
Actinomyces naeslundii		76.97	0.36	RAF 92	1	
Clostridium chauvoei		23.03	0.04	XYL 1	99	
27137123	**Good identification**					**27137123**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27137140	**Very good identification**					**27137140**
Propionibacterium propionicum		97.24	0.58	XYL 1		
Actinomyces israelii		2.76	0.33			
27137141	**Good identification**					**27137141**
Propionibacterium propionicum		98.00	0.42	XYL 1		
Actinomyces naeslundii		2.00	0.23	URE 10		
27137142	**Very good identification**					**27137142**
Propionibacterium propionicum		97.24	0.61	XYL 1		
Actinomyces israelii		2.76	0.36			
27137143	**Good identification**					**27137143**
Propionibacterium propionicum		98.66	0.45	XYL 1		
Actinomyces naeslundii		1.34	0.23	URE 10		
27137500	**Low discrimination**					**27137500**
Actinomyces israelii		85.53	0.38	SAL 92		
Propionibacterium propionicum		14.47	0.24	CEL 93		
27137502	**Low discrimination**					**27137502**
Actinomyces israelii		82.01	0.41	SAL 92		
Propionibacterium propionicum		13.88	0.27	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		4.12	0.13			
27140002	**Good identification**					**27140002**
Actinomyces viscosus		98.91	0.44	RAF 94		
Actinomyces odontolyticus		1.09	0.17	FRU 95		
27140003	**Good identification**					**27140003**
Actinomyces viscosus		100.0	0.35	RAF 94		
27140102	**Good identification**					**27140102**
Actinomyces viscosus		99.80	0.48	RAF 94		
Actinomyces odontolyticus		0.20	0.08	FRU 95		
27140103	**Good identification**					**27140103**
					SPO	
Actinomyces viscosus		98.53	0.40	RAF 94	1	
Clostridium chauvoei		1.47	0.10	FRU 95	99	
27141002	**Good identification**					**27141002**
Actinomyces viscosus		97.83	0.40	RAF 94		
Actinomyces odontolyticus		1.18	0.14	FRU 95		
Actinomyces naeslundii		0.99	0.30			
27141101	**Very good identification**					**27141101**
					SPO	
Clostridium chauvoei		99.40	0.57	CEL 1	99	
Actinomyces naeslundii		0.53	0.41		1	
Actinomyces viscosus		0.07	0.02		1	
27141102	**Good identification**					**27141102**
					SPO	
Actinomyces viscosus		97.43	0.44	RAF 94	1	
Actinomyces naeslundii		1.58	0.38	FRU 95	1	
Clostridium chauvoei		0.79	0.10		99	
27141103	**Low discrimination**					**27141103**
					SPO	
Clostridium chauvoei		69.15	0.44	CEL 1	99	
Actinomyces viscosus		28.76	0.36		1	
Actinomyces naeslundii		2.10	0.41		1	
27141121	**Good identification**					**27141121**
					SPO	
Clostridium chauvoei		98.42	0.48	CEL 1	99	
Actinomyces naeslundii		1.58	0.41		1	
27141123	**Acceptable identification**					**27141123**
					SPO	
Clostridium chauvoei		90.62	0.36	CEL 1	99	
Actinomyces naeslundii		8.24	0.41		1	
Actinomyces viscosus		1.14	0.02		1	
27143001	**Good identification**					**27143001**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27143003	**Good identification**					**27143003**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27143021	**Good identification**					**27143021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	RAF 92		
27143023	**Good identification**					**27143023**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27143100	**Low discrimination**					**27143100**
Actinomyces naeslundii		71.17	0.43	RAF 92		
Propionibacterium propionicum		28.84	0.26	FRU 95		
27143101	**Low discrimination**					**27143101**
					SPO	
Actinomyces naeslundii		50.68	0.46	RAF 92	1	
Clostridium chauvoei		49.32	0.22	FRU 95	99	
27143102	**Low discrimination**					**27143102**
Actinomyces naeslundii		51.83	0.43	RAF 92		
Propionibacterium propionicum		31.50	0.29	FRU 95		
Actinomyces viscosus		16.66	0.10			
27143103	**Low discrimination**					**27143103**
					SPO	
Actinomyces naeslundii		77.63	0.46	RAF 92	1	
Clostridium chauvoei		13.33	0.10	FRU 95	99	
Actinomyces viscosus		5.55	0.02		1	
27143120	**Good identification**					**27143120**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27143121	**Low discrimination**					**27143121**
					SPO	
Actinomyces naeslundii		75.51	0.46	RAF 92	1	
Clostridium chauvoei		24.49	0.14	FRU 95	99	
27143122	**Good identification**					**27143122**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27143123	**Acceptable identification**					**27143123**
					SPO	
Actinomyces naeslundii		94.59	0.46	RAF 92	1	
Clostridium chauvoei		5.42	0.01	FRU 95	99	
27150000	**Low discrimination**					**27150000**
					COC	
Actinomyces odontolyticus		35.33	0.39	MNS 9	1	
Actinomyces viscosus		32.69	0.31	CEL 1	1	
Gemella morbillorum		25.36	0.32		99	
27150002	**Very good identification**					**27150002**
Actinomyces viscosus		99.31	0.65	RAF 94		
Actinomyces odontolyticus		0.67	0.35			
Actinomyces naeslundii		0.03	0.28			
27150003	**Very good identification**					**27150003**
Actinomyces viscosus		99.88	0.57	RAF 94		
Actinomyces naeslundii		0.12	0.31			
27150100	**Low discrimination**					**27150100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces viscosus		74.82	0.36	RAF 94	99	
Actinomyces odontolyticus		14.37	0.30	BGL 99	99	
Fusobacterium mortiferum		3.95	0.26		1	
27150102	**Very good identification**					**27150102**
Actinomyces viscosus		99.78	0.70	RAF 94		
Actinomyces odontolyticus		0.12	0.27			
Propionibacterium propionicum		0.06	0.29			
27150103	**Very good identification**					**27150103**
					SPO	
Actinomyces viscosus		99.61	0.62	RAF 94	1	
Actinomyces naeslundii		0.19	0.39		1	
Clostridium chauvoei		0.18	0.16		99	
27150112	**Good identification**					**27150112**
Actinomyces viscosus		100.0	0.36	RAF 94		
27150122	**Low discrimination**					**27150122**
Actinomyces viscosus		80.68	0.36	RAF 94		
Clostridium perfringens		12.36	0.23	NAG 1		
Actinomyces naeslundii		3.30	0.36			
27150142	**Acceptable identification**					**27150142**
Actinomyces viscosus		93.61	0.36	RAF 94		
Propionibacterium propionicum		5.71	0.29	SOR 1		
Actinomyces naeslundii		0.68	0.23			
27150200	**Good identification**					**27150200**
Actinomyces odontolyticus		100.0	0.44	MNS 9		
27150202	**Low discrimination**					**27150202**
Actinomyces odontolyticus		56.26	0.40	MNS 9		
Actinomyces viscosus		43.74	0.31	CEL 1		
27150300	**Good identification**					**27150300**
Actinomyces odontolyticus		97.36	0.35	MNS 9		
Actinomyces viscosus		2.64	0.01	CEL 1		
27150302	**Low discrimination**					**27150302**
Actinomyces viscosus		81.40	0.36	RAF 94		
Actinomyces odontolyticus		18.60	0.32	RHA 1		
27150502	**Good identification**					**27150502**
Actinomyces viscosus		100.0	0.36	RAF 94		
27151000	**Low discrimination**					**27151000**
Lactobacillus jensenii		28.28	0.36	NIT 1		
Actinomyces odontolyticus		15.76	0.36	SAL 95		
Actinomyces naeslundii		13.42	0.52			
27151001	**Low discrimination**					**27151001**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		52.73	0.55	RAF 92	1	
Clostridium chauvoei		34.82	0.29		99	
Actinomyces viscosus		11.71	0.20		1	
27151002	**Very good identification**					**27151002**
Actinomyces viscosus		95.89	0.62	RAF 94		
Lactobacillus jensenii		1.36	0.33			
Actinomyces naeslundii		0.97	0.52			
27151003	**Good identification**					**27151003**
					SPO	
Actinomyces viscosus		95.17	0.54	RAF 94	1	
Actinomyces naeslundii		4.33	0.55		1	
Clostridium chauvoei		0.51	0.16		99	
27151020	**Low discrimination**					**27151020**
Clostridium perfringens		85.97	0.48	CEL 4		
Actinomyces naeslundii		6.75	0.52	LAC 88		
Lactobacillus jensenii		3.56	0.26			
27151021	**Low discrimination**					**27151021**
					SPO	
Actinomyces naeslundii		76.58	0.55	RAF 92	1	
Clostridium chauvoei		16.86	0.21		99	
Clostridium perfringens		6.57	0.14		84	
27151022	**Low discrimination**					**27151022**
Clostridium perfringens		66.57	0.43	CEL 4		
Actinomyces naeslundii		11.11	0.52	LAC 88		
Actinomyces viscosus		11.10	0.28			
27151023	**Low discrimination**					**27151023**
Actinomyces naeslundii		76.85	0.55	RAF 92		
Actinomyces viscosus		17.06	0.20			
Clostridium perfringens		3.10	0.09			
27151040	**Low discrimination**					**27151040**
Actinomyces naeslundii		64.70	0.39	RAF 92		
Clostridium perfringens		27.50	0.10			
Lactobacillus jensenii		7.81	0.02			
27151041	**Good identification**					**27151041**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27151060	**Acceptable identification**					**27151060**
Clostridium perfringens		93.22	0.36	CEL 4		
Actinomyces naeslundii		6.78	0.39	LAC 88		
27151061	**Acceptable identification**					**27151061**
Actinomyces naeslundii		91.53	0.42	RAF 92		
Clostridium perfringens		8.47	0.02			
27151063	**Good identification**					**27151063**
Actinomyces naeslundii		100.0	0.42	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27151100	**Low discrimination**						**27151100**
Clostridium perfringens		26.22	0.37	CEL 4			
Actinomyces naeslundii		25.84	0.60	NAG 97			
Actinomyces viscosus		16.13	0.32				
27151101	**Very good identification**						**27151101**
					SPO		
Clostridium chauvoei		95.25	0.63	CEL 1	99		
Actinomyces naeslundii		4.15	0.63		1		
Actinomyces viscosus		0.58	0.24		1		
27151102	**Very good identification**						**27151102**
Actinomyces viscosus		95.89	0.66	RAF 94			
Actinomyces naeslundii		1.55	0.60				
Clostridium perfringens		0.74	0.32				
27151103	**Low discrimination**						**27151103**
					SPO		
Actinomyces viscosus		73.08	0.58	RAF 94	1		
Clostridium chauvoei		21.58	0.50		99		
Actinomyces naeslundii		5.32	0.63		1		
27151110	**Low discrimination**						**27151110**
Actinomyces naeslundii		83.70	0.38	RAF 92			
Clostridium perfringens		16.30	0.03	MLZ 5			
27151120	**Good identification**						**27151120**
					GRA		
Clostridium perfringens		91.44	0.63	CEL 4	99		
Prevotella melaninogenica		4.07	0.33		1		
Actinomyces naeslundii		2.79	0.60		99		
27151121	**Low discrimination**						**27151121**
					SPO		
Clostridium chauvoei		86.15	0.55	CEL 1	99		
Actinomyces naeslundii		11.25	0.63		1		
Clostridium perfringens		2.49	0.29		84		
27151122	**Low discrimination**						**27151122**
					GRA	CAT	
Clostridium perfringens		75.63	0.58	CEL 4	99	1	
Capnocytophaga ochracea		9.66	0.39		1	90	
Actinomyces naeslundii		4.90	0.60		99	25	
27151123	**Low discrimination**						**27151123**
					SPO		
Clostridium chauvoei		51.83	0.42	CEL 1	99		
Actinomyces naeslundii		38.36	0.63		1		
Actinomyces viscosus		5.32	0.24		1		
27151133	**Low discrimination**						**27151133**
					SPO		
Actinomyces naeslundii		79.41	0.41	RAF 92	1		
Clostridium chauvoei		20.59	0.08	MLZ 5	99		
27151143	**Low discrimination**						**27151143**
					SPO		
Actinomyces naeslundii		48.93	0.50	RAF 92	1		
Actinomyces viscosus		38.45	0.24		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		11.35	0.16		99		
27151160	**Good identification**						**27151160**
					GRA	CAT	
Clostridium perfringens		96.67	0.51	CEL 4	99	1	
Actinomyces naeslundii		2.73	0.47		99	25	
Capnocytophaga ochracea		0.38	0.07		1	90	
27151161	**Low discrimination**						**27151161**
					SPO		
Actinomyces naeslundii		59.64	0.50	RAF 92	1		
Clostridium chauvoei		26.14	0.21		99		
Clostridium perfringens		14.22	0.17		84		
27151162	**Low discrimination**						**27151162**
					GRA	CAT	
Clostridium perfringens		87.24	0.45	CEL 4	99	1	
Capnocytophaga ochracea		6.50	0.23		1	90	
Actinomyces naeslundii		5.24	0.47		99	25	
27151163	**Low discrimination**						**27151163**
					SPO		
Actinomyces naeslundii		84.07	0.50	RAF 92	1		
Clostridium perfringens		9.43	0.11		84		
Clostridium chauvoei		6.50	0.08		99		
27151200	**Good identification**						**27151200**
Actinomyces odontolyticus		99.08	0.41	MNS 9			
Lactobacillus jensenii		0.93	0.02	CEL 1			
27151202	**Low discrimination**						**27151202**
Actinomyces odontolyticus		58.32	0.37	MNS 9			
Actinomyces viscosus		41.69	0.28	CEL 1			
27152001	**Good identification**						**27152001**
Actinomyces naeslundii		100.0	0.36	RAF 92			
27152021	**Good identification**						**27152021**
Actinomyces naeslundii		100.0	0.36	RAF 92			
27152023	**Good identification**						**27152023**
Actinomyces naeslundii		100.0	0.36	RAF 92			
27152100	**Very good identification**						**27152100**
Propionibacterium propionicum		97.82	0.60	CEL 1			
Actinomyces naeslundii		1.93	0.41				
Actinomyces viscosus		0.25	0.01				
27152101	**Low discrimination**						**27152101**
Propionibacterium propionicum		78.96	0.44	CEL 1			
Actinomyces naeslundii		21.04	0.44	URE 10			
27152102	**Low discrimination**						**27152102**
Propionibacterium propionicum		84.84	0.63	CEL 1			
Actinomyces viscosus		14.05	0.36				
Actinomyces naeslundii		1.12	0.41				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27152103	**Low discrimination**					**27152103**
Propionibacterium propionicum		59.72	0.47	CEL 1		
Actinomyces viscosus		29.67	0.27	URE 10		
Actinomyces naeslundii		10.61	0.44			
27152120	**Low discrimination**					**27152120**
Actinomyces naeslundii		66.15	0.41	RAF 92		
Propionibacterium propionicum		33.85	0.26	GAL 96		
27152121	**Good identification**					**27152121**
Actinomyces naeslundii		100.0	0.44	RAF 92		
27152122	**Low discrimination**					**27152122**
Actinomyces naeslundii		52.77	0.41	RAF 92		
Propionibacterium propionicum		40.52	0.29	GAL 96		
Actinomyces viscosus		6.71	0.01			
27152123	**Acceptable identification**					**27152123**
Actinomyces naeslundii		94.62	0.44	RAF 92		
Propionibacterium propionicum		5.38	0.12	GAL 96		
27152140	**Very good identification**					**27152140**
Propionibacterium propionicum		99.65	0.60	CEL 1		
Actinomyces naeslundii		0.35	0.28			
27152141	**Good identification**					**27152141**
Propionibacterium propionicum		95.51	0.44	CEL 1		
Actinomyces naeslundii		4.49	0.31	URE 10		
27152142	**Very good identification**					**27152142**
Propionibacterium propionicum		99.60	0.63	CEL 1		
Actinomyces naeslundii		0.23	0.28			
Actinomyces viscosus		0.17	0.01			
27152143	**Good identification**					**27152143**
Propionibacterium propionicum		96.96	0.47	CEL 1		
Actinomyces naeslundii		3.04	0.31	URE 10		
27153000	**Low discrimination**					**27153000**
Actinomyces naeslundii		89.14	0.57	RAF 92		
Propionibacterium propionicum		5.41	0.26			
Actinomyces israelii		2.46	0.21			
27153001	**Very good identification**					**27153001**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27153002	**Low discrimination**					**27153002**
Actinomyces naeslundii		60.72	0.57	RAF 92		
Actinomyces viscosus		31.25	0.28			
Propionibacterium propionicum		5.53	0.29			
27153003	**Low discrimination**					**27153003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		89.20	0.60	RAF 92		
Actinomyces viscosus		10.20	0.20			
Propionibacterium propionicum		0.60	0.12			
27153011	**Good identification**					**27153011**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27153013	**Good identification**					**27153013**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27153020	**Good identification**					**27153020**
Actinomyces naeslundii		93.79	0.57	RAF 92		
Clostridium perfringens		6.21	0.14			
27153021	**Very good identification**					**27153021**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27153022	**Very good identification**					**27153022**
Actinomyces naeslundii		96.98	0.57	RAF 92		
Clostridium perfringens		3.02	0.09			
27153023	**Very good identification**					**27153023**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27153031	**Good identification**					**27153031**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27153033	**Good identification**					**27153033**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27153040	**Low discrimination**					**27153040**
Actinomyces naeslundii		71.19	0.44	RAF 92		
Propionibacterium propionicum		24.49	0.26			
Actinomyces israelii		4.32	0.14			
27153041	**Good identification**					**27153041**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27153042	**Low discrimination**					**27153042**
Actinomyces naeslundii		62.23	0.44	RAF 92		
Propionibacterium propionicum		32.11	0.29			
Actinomyces israelii		5.67	0.17			
27153043	**Good identification**					**27153043**
Actinomyces naeslundii		96.32	0.47	RAF 92		
Propionibacterium propionicum		3.68	0.12			
27153060	**Acceptable identification**					**27153060**
Actinomyces naeslundii		93.33	0.44	RAF 92		
Clostridium perfringens		6.67	0.02			
27153061	**Good identification**					**27153061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	RAF 92		
27153062	**Good identification**					**27153062**
Actinomyces naeslundii		100.0	0.44	RAF 92		
27153063	**Good identification**					**27153063**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27153100	**Low discrimination**					**27153100**
Propionibacterium propionicum		66.91	0.60	CEL 1		
Actinomyces naeslundii		31.69	0.65			
Actinomyces israelii		1.23	0.31			
27153101	**Low discrimination**					**27153101**
					SPO	
Actinomyces naeslundii		78.38	0.68	RAF 92	1	
Propionibacterium propionicum		12.26	0.44		1	
Clostridium chauvoei		9.37	0.29		99	
27153102	**Low discrimination**					**27153102**
Propionibacterium propionicum		69.58	0.63	CEL 1		
Actinomyces naeslundii		21.97	0.65			
Actinomyces viscosus		7.06	0.32			
27153103	**Low discrimination**					**27153103**
Actinomyces naeslundii		75.35	0.68	RAF 92		
Propionibacterium propionicum		17.68	0.47			
Actinomyces viscosus		5.38	0.24			
27153110	**Low discrimination**					**27153110**
Actinomyces naeslundii		66.00	0.43	RAF 92		
Propionibacterium propionicum		26.74	0.26	MLZ 5		
Actinomyces israelii		7.25	0.17			
27153111	**Good identification**					**27153111**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27153112	**Low discrimination**					**27153112**
Actinomyces naeslundii		56.41	0.43	RAF 92		
Propionibacterium propionicum		34.29	0.29	MLZ 5		
Actinomyces israelii		9.30	0.20			
27153113	**Good identification**					**27153113**
Actinomyces naeslundii		95.69	0.46	RAF 92		
Propionibacterium propionicum		4.31	0.12	MLZ 5		
27153120	**Low discrimination**					**27153120**
Actinomyces naeslundii		83.36	0.65	RAF 92		
Clostridium perfringens		14.23	0.29			
Propionibacterium propionicum		1.78	0.26			
27153121	**Very good identification**					**27153121**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.17	0.68	RAF 92	1	
Clostridium chauvoei		3.83	0.21		99	
27153122	**Low discrimination**					**27153122**
Actinomyces naeslundii		87.61	0.65	RAF 92		
Clostridium perfringens		7.04	0.23			
Propionibacterium propionicum		2.80	0.29			
27153123	**Very good identification**					**27153123**
					SPO	
Actinomyces naeslundii		99.07	0.68	RAF 92	1	
Clostridium chauvoei		0.70	0.08		99	
Propionibacterium propionicum		0.24	0.12		1	
27153130	**Good identification**					**27153130**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27153131	**Good identification**					**27153131**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27153132	**Low discrimination**					**27153132**
					SPO	
Actinomyces naeslundii		87.19	0.43	RAF 92	1	
Clostridium tertium		12.81	0.05	MLZ 5	99	
27153133	**Good identification**					**27153133**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27153140	**Good identification**					**27153140**
Propionibacterium propionicum		91.68	0.60	CEL 1		
Actinomyces naeslundii		7.66	0.52			
Actinomyces israelii		0.65	0.24			
27153141	**Low discrimination**					**27153141**
Actinomyces naeslundii		53.02	0.55	RAF 92		
Propionibacterium propionicum		46.98	0.44			
27153142	**Good identification**					**27153142**
Propionibacterium propionicum		94.09	0.63	CEL 1		
Actinomyces naeslundii		5.24	0.52			
Actinomyces israelii		0.67	0.27			
27153143	**Low discrimination**					**27153143**
Propionibacterium propionicum		57.07	0.47	CEL 1		
Actinomyces naeslundii		42.93	0.55	URE 10		
27153160	**Low discrimination**					**27153160**
Actinomyces naeslundii		76.62	0.52	RAF 92		
Clostridium perfringens		14.12	0.17			
Propionibacterium propionicum		9.26	0.26			
27153161	**Good identification**					**27153161**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27153162	**Low discrimination**					**27153162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		78.87	0.52	RAF 92		
Propionibacterium propionicum		14.30	0.29			
Clostridium perfringens		6.84	0.11			
27153163	**Good identification**					**27153163**
Actinomyces naeslundii		98.68	0.55	RAF 92		
Propionibacterium propionicum		1.33	0.12			
27154102	**Acceptable identification**					**27154102**
Actinomyces viscosus		94.52	0.36	RAF 94		
Actinomyces odontolyticus		5.48	0.21	XYL 1		
27154200	**Good identification**					**27154200**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27155100	**Low discrimination**					**27155100**
					COC	
Actinomyces israelii		38.08	0.36	SAL 92	1	
Atopobium parvulum		38.08	0.23		99	
Actinomyces odontolyticus		16.03	0.22		1	
27155200	**Good identification**					**27155200**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27155502	**Low discrimination**					**27155502**
Actinomyces israelii		70.23	0.38	SAL 92		
Bifidobacterium adolescentis		22.04	0.23			
Lactobacillus fermentum		4.10	0.16			
27157000	**Good identification**					**27157000**
Actinomyces israelii		96.91	0.39	SAL 92		
Actinomyces naeslundii		3.09	0.23			
27157002	**Good identification**					**27157002**
Actinomyces israelii		97.92	0.42	SAL 92		
Actinomyces naeslundii		2.08	0.23			
27157042	**Good identification**					**27157042**
Actinomyces israelii		100.0	0.35	SAL 92		
27157100	**Acceptable identification**					**27157100**
Actinomyces israelii		93.41	0.49	SAL 92		
Propionibacterium propionicum		4.48	0.26			
Actinomyces naeslundii		2.12	0.30			
27157102	**Acceptable identification**					**27157102**
Actinomyces israelii		93.58	0.52	SAL 92		
Propionibacterium propionicum		4.48	0.29			
Actinomyces naeslundii		1.42	0.30			
27157110	**Good identification**					**27157110**
Actinomyces israelii		100.0	0.35	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27157112	**Acceptable identification**					**27157112**
					SPO	
Actinomyces israelii		92.51	0.38	SAL 92	1	
Bifidobacterium adolescentis		4.19	0.09	MLZ 13	1	
Clostridium tertium		3.30	-.01		99	
27157140	**Low discrimination**					**27157140**
Actinomyces israelii		89.03	0.42	SAL 92		
Propionibacterium propionicum		10.97	0.26			
27157142	**Low discrimination**					**27157142**
Actinomyces israelii		89.03	0.45	SAL 92		
Propionibacterium propionicum		10.97	0.29			
27157400	**Good identification**					**27157400**
Actinomyces israelii		100.0	0.38	SAL 92		
27157402	**Good identification**					**27157402**
Actinomyces israelii		100.0	0.41	SAL 92		
27157500	**Good identification**					**27157500**
Actinomyces israelii		100.0	0.48	SAL 92		
27157502	**Good identification**					**27157502**
Actinomyces israelii		95.67	0.51	SAL 92		
Bifidobacterium adolescentis		4.34	0.22			
27157512	**Low discrimination**					**27157512**
Actinomyces israelii		72.96	0.37	SAL 92		
Bifidobacterium adolescentis		27.04	0.23	MLZ 13		
27157540	**Good identification**					**27157540**
Actinomyces israelii		100.0	0.41	SAL 92		
27157542	**Good identification**					**27157542**
Actinomyces israelii		100.0	0.44	SAL 92		
27161001	**Low discrimination**					**27161001**
					SPO	
Actinomyces naeslundii		76.47	0.38	RAF 92	1	
Clostridium chauvoei		23.53	0.06	FRU 95	99	
27161003	**Low discrimination**					**27161003**
Actinomyces naeslundii		53.99	0.38	RAF 92		
Actinomyces viscosus		46.01	0.13	FRU 95		
27161021	**Good identification**					**27161021**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27161023	**Good identification**					**27161023**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27161100	**Low discrimination**					**27161100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Actinomyces naeslundii		46.47	0.43	RAF 92	1	
Atopobium parvulum		37.33	0.18	FRU 95	99	
Clostridium perfringens		8.66	0.08		1	
27161101	**Acceptable identification**					**27161101**
					SPO	
Clostridium chauvoei		91.46	0.40	TRE 10	99	
Actinomyces naeslundii		8.54	0.46	CEL 1	1	
27161123	**Low discrimination**					**27161123**
					SPO	
Actinomyces naeslundii		61.36	0.46	RAF 92	1	
Clostridium chauvoei		38.64	0.19	FRU 95	99	
27163000	**Good identification**					**27163000**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27163001	**Good identification**					**27163001**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27163002	**Good identification**					**27163002**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27163003	**Good identification**					**27163003**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27163020	**Good identification**					**27163020**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27163021	**Good identification**					**27163021**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27163022	**Good identification**					**27163022**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27163023	**Good identification**					**27163023**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27163100	**Low discrimination**					**27163100**
Actinomyces naeslundii		85.41	0.48	RAF 92		
Propionibacterium propionicum		14.59	0.24	FRU 95		
27163101	**Acceptable identification**					**27163101**
					SPO	
Actinomyces naeslundii		94.72	0.51	RAF 92	1	
Clostridium chauvoei		5.28	0.06	FRU 95	99	
27163102	**Low discrimination**					**27163102**
Actinomyces naeslundii		79.61	0.48	RAF 92		
Propionibacterium propionicum		20.39	0.27	FRU 95		
27163103	**Good identification**					**27163103**
Actinomyces naeslundii		100.0	0.51	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27163120	**Good identification**					**27163120**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27163121	**Good identification**					**27163121**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27163122	**Good identification**					**27163122**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27163123	**Good identification**					**27163123**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27163141	**Good identification**					**27163141**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27163143	**Good identification**					**27163143**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27163161	**Good identification**					**27163161**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27163163	**Good identification**					**27163163**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27170001	**Low discrimination**					**27170001**
Actinomyces naeslundii		74.79	0.36	RAF 92		
Actinomyces viscosus		25.21	0.04	GAL 96		
27170002	**Good identification**					**27170002**
Actinomyces viscosus		98.64	0.46	RAF 94		
Actinomyces naeslundii		0.66	0.33	TRE 7		
Propionibacterium granulosum		0.35	0.14			
27170003	**Good identification**					**27170003**
Actinomyces viscosus		97.09	0.38	RAF 94		
Actinomyces naeslundii		2.91	0.36	TRE 7		
27170021	**Good identification**					**27170021**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27170023	**Low discrimination**					**27170023**
Actinomyces naeslundii		74.79	0.36	RAF 92		
Actinomyces viscosus		25.21	0.04	GAL 96		
27170101	**Low discrimination**					**27170101**
					SPO	
Actinomyces naeslundii		65.39	0.44	RAF 92	1	
Clostridium chauvoei		20.84	0.12	GAL 96	99	
Actinomyces viscosus		13.77	0.08		1	
27170102	**Good identification**					**27170102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		98.00	0.51	RAF 94		
Actinomyces naeslundii		1.05	0.41	TRE 7		
Propionibacterium propionicum		0.66	0.27			
27170103	**Good identification**					**27170103**
					SPO	
Actinomyces viscosus		95.18	0.42	RAF 94	1	
Actinomyces naeslundii		4.57	0.44	TRE 7	1	
Clostridium chauvoei		0.26	-.01		99	
27170121	**Acceptable identification**					**27170121**
					SPO	
Actinomyces naeslundii		90.40	0.44	RAF 92	1	
Clostridium chauvoei		9.60	0.04	GAL 96	99	
27170123	**Low discrimination**					**27170123**
Actinomyces naeslundii		82.61	0.44	RAF 92		
Actinomyces viscosus		17.39	0.08	GAL 96		
27171000	**Low discrimination**					**27171000**
Lactobacillus jensenii		66.26	0.44	NIT 1		
Actinomyces naeslundii		20.34	0.57	SAL 95		
Clostridium perfringens		7.66	0.27			
27171001	**Good identification**					**27171001**
					SPO	
Actinomyces naeslundii		93.60	0.60	RAF 92	1	
Clostridium chauvoei		3.54	0.12		99	
Lactobacillus jensenii		2.05	0.10		1	
27171002	**Low discrimination**					**27171002**
Actinomyces viscosus		52.28	0.43	RAF 94		
Lactobacillus jensenii		29.61	0.41	TRE 7		
Actinomyces naeslundii		13.63	0.57			
27171003	**Low discrimination**					**27171003**
Actinomyces naeslundii		53.38	0.60	RAF 92		
Actinomyces viscosus		45.49	0.34			
Lactobacillus jensenii		0.78	0.07			
27171010	**Acceptable identification**					**27171010**
					CAT	
Lactobacillus jensenii		94.36	0.36	NIT 1	1	
Actinomyces naeslundii		4.57	0.35	SAL 95	25	
Propionibacterium granulosum		1.06	0.09		90	
27171011	**Low discrimination**					**27171011**
Actinomyces naeslundii		87.80	0.38	RAF 92		
Lactobacillus jensenii		12.20	0.02	MLZ 5		
27171013	**Low discrimination**					**27171013**
Actinomyces naeslundii		85.95	0.38	RAF 92		
Actinomyces viscosus		14.06	0.00	MLZ 5		
27171020	**Low discrimination**					**27171020**
Clostridium perfringens		86.77	0.53	CEL 4		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		7.12	0.57	LAC 88		
Lactobacillus jensenii		5.80	0.34			
27171021	**Good identification**					**27171021**
					SPO	
Actinomyces naeslundii		91.36	0.60	RAF 92	1	
Clostridium perfringens		7.49	0.19		84	
Clostridium chauvoei		1.15	0.04		99	
27171022	**Low discrimination**					**27171022**
Clostridium perfringens		77.52	0.47	CEL 4		
Actinomyces naeslundii		13.53	0.57	LAC 88		
Lactobacillus jensenii		7.34	0.31			
27171023	**Very good identification**					**27171023**
Actinomyces naeslundii		95.49	0.60	RAF 92		
Clostridium perfringens		3.69	0.13			
Actinomyces viscosus		0.82	0.00			
27171031	**Good identification**					**27171031**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27171033	**Good identification**					**27171033**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27171040	**Low discrimination**					**27171040**
Actinomyces naeslundii		59.77	0.44	RAF 92		
Clostridium perfringens		24.30	0.15			
Lactobacillus jensenii		11.14	0.10			
27171041	**Good identification**					**27171041**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27171042	**Low discrimination**					**27171042**
Actinomyces naeslundii		60.54	0.44	RAF 92		
Actinomyces viscosus		13.29	0.08			
Clostridium perfringens		11.58	0.09			
27171043	**Good identification**					**27171043**
Actinomyces naeslundii		95.35	0.47	RAF 92		
Actinomyces viscosus		4.65	0.00			
27171060	**Acceptable identification**					**27171060**
Clostridium perfringens		92.93	0.41	CEL 4		
Actinomyces naeslundii		7.07	0.44	LAC 88		
27171061	**Acceptable identification**					**27171061**
Actinomyces naeslundii		91.87	0.47	RAF 92		
Clostridium perfringens		8.13	0.07			
27171062	**Low discrimination**					**27171062**
Clostridium perfringens		85.42	0.35	CEL 4		
Actinomyces naeslundii		13.81	0.44	LAC 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		0.78	-.02				
27171063	**Good identification**						**27171063**
Actinomyces naeslundii		96.00	0.47	RAF 92			
Clostridium perfringens		4.00	0.01				
27171100	**Low discrimination**						**27171100**
					COC		
Atopobium parvulum		66.78	0.52	NIT 1	99		
Actinomyces naeslundii		15.96	0.65		1		
Clostridium perfringens		15.49	0.42		1		
27171101	**Low discrimination**						**27171101**
					COC	SPO	
Clostridium chauvoei		55.83	0.47	TRE 10	1	99	
Actinomyces naeslundii		42.47	0.68	CEL 1	1	1	
Atopobium parvulum		1.20	0.18		99	1	
27171102	**Low discrimination**						**27171102**
					COC		
Actinomyces viscosus		47.99	0.47	RAF 94	1		
Actinomyces naeslundii		20.04	0.65	TRE 7	1		
Atopobium parvulum		14.80	0.39		99		
27171103	**Low discrimination**						**27171103**
					SPO		
Actinomyces naeslundii		56.42	0.68	RAF 92	1		
Actinomyces viscosus		30.03	0.39		1		
Clostridium chauvoei		13.09	0.34		99		
27171110	**Low discrimination**						**27171110**
					COC		
Actinomyces naeslundii		46.82	0.43	RAF 92	1		
Atopobium parvulum		37.61	0.18	MLZ 5	99		
Clostridium perfringens		8.72	0.08		1		
27171111	**Low discrimination**						**27171111**
					SPO		
Actinomyces naeslundii		79.85	0.46	RAF 92	1		
Clostridium chauvoei		20.15	0.12	MLZ 5	99		
27171113	**Low discrimination**						**27171113**
					SPO		
Actinomyces naeslundii		87.21	0.46	RAF 92	1		
Actinomyces viscosus		8.91	0.05	MLZ 5	1		
Clostridium chauvoei		3.88	-.01		99		
27171120	**Good identification**						**27171120**
					COC		
Clostridium perfringens		94.53	0.68	CEL 4	1		
Actinomyces naeslundii		3.01	0.65		1		
Atopobium parvulum		2.23	0.39		99		
27171121	**Low discrimination**						**27171121**
					SPO		
Actinomyces naeslundii		60.44	0.68	RAF 92	1		
Clostridium chauvoei		26.49	0.38		99		
Clostridium perfringens		12.77	0.34		84		
27171122	**Good identification**						**27171122**
Clostridium perfringens		90.64	0.62	CEL 4			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		6.14	0.65			
Lactobacillus acidophilus		1.99	0.32			
27171123	**Low discrimination**					**27171123**
					SPO	
Actinomyces naeslundii		84.59	0.68	RAF 92	1	
Clostridium perfringens		8.41	0.28		84	
Clostridium chauvoei		6.54	0.25		99	
27171131	**Acceptable identification**					**27171131**
					SPO	
Actinomyces naeslundii		92.24	0.46	RAF 92	1	
Clostridium chauvoei		7.76	0.04	MLZ 5	99	
27171133	**Good identification**					**27171133**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27171141	**Acceptable identification**					**27171141**
					SPO	
Actinomyces naeslundii		93.00	0.55	RAF 92	1	
Clostridium chauvoei		7.00	0.12		99	
27171142	**Low discrimination**					**27171142**
Actinomyces naeslundii		46.88	0.52	RAF 92		
Clostridium perfringens		23.11	0.24			
Lactobacillus acidophilus		10.66	0.16			
27171143	**Good identification**					**27171143**
					SPO	
Actinomyces naeslundii		95.81	0.55	RAF 92	1	
Actinomyces viscosus		2.92	0.05		1	
Clostridium chauvoei		1.27	-.01		99	
27171160	**Good identification**					**27171160**
					COC	
Clostridium perfringens		97.02	0.56	CEL 4	1	
Actinomyces naeslundii		2.86	0.52		1	
Atopobium parvulum		0.12	0.05		99	
27171161	**Low discrimination**					**27171161**
					SPO	
Actinomyces naeslundii		79.80	0.55	RAF 92	1	
Clostridium perfringens		18.20	0.21		84	
Clostridium chauvoei		2.00	0.04		99	
27171162	**Acceptable identification**					**27171162**
Clostridium perfringens		92.85	0.50	CEL 4		
Actinomyces naeslundii		5.83	0.52			
Lactobacillus acidophilus		1.32	0.16			
27171163	**Acceptable identification**					**27171163**
Actinomyces naeslundii		90.31	0.55	RAF 92		
Clostridium perfringens		9.69	0.16			
27172000	**Low discrimination**					**27172000**
					CAT	
Actinomyces naeslundii		40.79	0.38	RAF 92	25	
Propionibacterium granulosum		34.16	0.22	GAL 96	90	
Propionibacterium propionicum		25.05	0.24		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27172001	**Good identification**					**27172001**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27172002	**Low discrimination**					**27172002**
Actinomyces naeslundii		34.34	0.38	RAF 92		
Propionibacterium propionicum		31.63	0.27	GAL 96		
Actinomyces viscosus		26.84	0.12			
27172003	**Low discrimination**					**27172003**
Actinomyces naeslundii		85.20	0.41	RAF 92		
Actinomyces viscosus		14.80	0.04	GAL 96		
27172020	**Low discrimination**					**27172020**
Actinomyces naeslundii		82.69	0.38	RAF 92		
Propionibacterium granulosum		17.31	0.12	GAL 96		
27172021	**Good identification**					**27172021**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27172022	**Good identification**					**27172022**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27172023	**Good identification**					**27172023**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27172100	**Very good identification**					**27172100**
Propionibacterium propionicum		95.53	0.58	CEL 1		
Actinomyces naeslundii		4.47	0.46			
27172101	**Low discrimination**					**27172101**
Propionibacterium propionicum		61.27	0.42	CEL 1		
Actinomyces naeslundii		38.73	0.49	URE 10		
27172102	**Very good identification**					**27172102**
Propionibacterium propionicum		95.56	0.61	CEL 1		
Actinomyces naeslundii		2.98	0.46			
Actinomyces viscosus		1.46	0.16			
27172103	**Low discrimination**					**27172103**
Propionibacterium propionicum		68.16	0.45	CEL 1		
Actinomyces naeslundii		28.72	0.49	URE 10		
Actinomyces viscosus		3.12	0.08			
27172120	**Low discrimination**					**27172120**
Actinomyces naeslundii		82.26	0.46	RAF 92		
Propionibacterium propionicum		17.74	0.24	GAL 96		
27172121	**Good identification**					**27172121**
Actinomyces naeslundii		100.0	0.49	RAF 92		
27172122	**Low discrimination**					**27172122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		75.55	0.46	RAF 92		
Propionibacterium propionicum		24.45	0.27	GAL 96		
27172123	**Good identification**					**27172123**
Actinomyces naeslundii		100.0	0.49	RAF 92		
27172140	**Very good identification**					**27172140**
Propionibacterium propionicum		99.18	0.58	CEL 1		
Actinomyces naeslundii		0.82	0.33			
27172141	**Low discrimination**					**27172141**
Propionibacterium propionicum		89.96	0.42	CEL 1		
Actinomyces naeslundii		10.04	0.36	URE 10		
27172142	**Very good identification**					**27172142**
Propionibacterium propionicum		99.45	0.61	CEL 1		
Actinomyces naeslundii		0.55	0.33			
27172143	**Acceptable identification**					**27172143**
Propionibacterium propionicum		93.08	0.45	CEL 1		
Actinomyces naeslundii		6.92	0.36	URE 10		
27172161	**Good identification**					**27172161**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27172163	**Good identification**					**27172163**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27173000	**Low discrimination**					**27173000**
					CAT	
Actinomyces naeslundii		88.43	0.62	RAF 92	25	
Actinomyces israelii		4.72	0.31		1	
Propionibacterium granulosum		3.09	0.22		90	
27173001	**Very good identification**					**27173001**
Actinomyces naeslundii		100.0	0.65	RAF 92		
27173002	**Low discrimination**					**27173002**
Actinomyces naeslundii		86.33	0.62	RAF 92		
Actinomyces israelii		6.91	0.34			
Propionibacterium propionicum		3.31	0.27			
27173003	**Very good identification**					**27173003**
Actinomyces naeslundii		99.56	0.65	RAF 92		
Actinomyces viscosus		0.44	0.00			
27173010	**Low discrimination**					**27173010**
Actinomyces naeslundii		79.44	0.40	RAF 92		
Actinomyces israelii		12.03	0.17	MLZ 5		
Lactobacillus jensenii		8.53	0.02			
27173011	**Good identification**					**27173011**
Actinomyces naeslundii		100.0	0.43	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27173012	**Low discrimination**					**27173012**
Actinomyces naeslundii		81.48	0.40	RAF 92		
Actinomyces israelii		18.52	0.20	MLZ 5		
27173013	**Good identification**					**27173013**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27173020	**Good identification**					**27173020**
					CAT	
Actinomyces naeslundii		93.28	0.62	RAF 92	25	
Clostridium perfringens		5.91	0.19		1	
Propionibacterium granulosum		0.81	0.12		90	
27173021	**Very good identification**					**27173021**
Actinomyces naeslundii		100.0	0.65	RAF 92		
27173022	**Very good identification**					**27173022**
Actinomyces naeslundii		96.62	0.62	RAF 92		
Clostridium perfringens		2.88	0.13			
Actinomyces israelii		0.49	0.13			
27173023	**Very good identification**					**27173023**
Actinomyces naeslundii		100.0	0.65	RAF 92		
27173030	**Good identification**					**27173030**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27173031	**Good identification**					**27173031**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27173032	**Good identification**					**27173032**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27173033	**Good identification**					**27173033**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27173040	**Low discrimination**					**27173040**
Actinomyces naeslundii		79.20	0.49	RAF 92		
Propionibacterium propionicum		11.48	0.24			
Actinomyces israelii		9.31	0.24			
27173041	**Good identification**					**27173041**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27173042	**Low discrimination**					**27173042**
Actinomyces naeslundii		71.74	0.49	RAF 92		
Propionibacterium propionicum		15.60	0.27			
Actinomyces israelii		12.66	0.27			
27173043	**Good identification**					**27173043**
Actinomyces naeslundii		100.0	0.52	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27173060	**Acceptable identification**					**27173060**
Actinomyces naeslundii		93.60	0.49	RAF 92		
Clostridium perfringens		6.40	0.07			
27173061	**Good identification**					**27173061**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27173062	**Good identification**					**27173062**
Actinomyces naeslundii		96.88	0.49	RAF 92		
Clostridium perfringens		3.12	0.01			
27173063	**Good identification**					**27173063**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27173100	**Low discrimination**					**27173100**
Actinomyces naeslundii		50.15	0.69	RAF 92		
Propionibacterium propionicum		44.63	0.58			
Actinomyces israelii		3.76	0.41			
27173101	**Good identification**					**27173101**
					SPO	
Actinomyces naeslundii		93.22	0.72	RAF 92	1	
Propionibacterium propionicum		6.15	0.42		1	
Clostridium chauvoei		0.64	0.12		99	
27173102	**Low discrimination**					**27173102**
Propionibacterium propionicum		53.79	0.61	CEL 1		
Actinomyces naeslundii		40.30	0.69			
Actinomyces israelii		4.53	0.44			
27173103	**Good identification**					**27173103**
Actinomyces naeslundii		90.67	0.72	RAF 92		
Propionibacterium propionicum		8.97	0.45			
Actinomyces viscosus		0.25	0.05			
27173110	**Low discrimination**					**27173110**
Actinomyces naeslundii		68.52	0.48	RAF 92		
Actinomyces israelii		14.59	0.27	MLZ 5		
Propionibacterium propionicum		11.70	0.24			
27173111	**Good identification**					**27173111**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27173112	**Low discrimination**					**27173112**
Actinomyces naeslundii		55.47	0.48	RAF 92		
Actinomyces israelii		17.72	0.30	MLZ 5		
Propionibacterium propionicum		14.21	0.27			
27173113	**Good identification**					**27173113**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27173120	**Low discrimination**					**27173120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		83.75	0.69	RAF 92		
Clostridium perfringens		13.68	0.34			
Propionibacterium propionicum		0.75	0.24			
27173121	**Very good identification**					**27173121**
					SPO	
Actinomyces naeslundii		99.77	0.72	RAF 92	1	
Clostridium chauvoei		0.23	0.04		99	
27173122	**Low discrimination**					**27173122**
					SPO	
Actinomyces naeslundii		86.82	0.69	RAF 92	1	
Clostridium perfringens		6.67	0.28		84	
Clostridium ramosum		2.20	0.17		92	
27173123	**Very good identification**					**27173123**
Actinomyces naeslundii		100.0	0.72	RAF 92		
27173130	**Low discrimination**					**27173130**
					SPO	
Actinomyces naeslundii		81.49	0.48	RAF 92	1	
Clostridium tertium		18.51	0.13	MLZ 5	99	
27173131	**Good identification**					**27173131**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27173132	**Low discrimination**					**27173132**
					SPO	
Actinomyces naeslundii		59.48	0.48	RAF 92	1	
Clostridium tertium		40.52	0.21	MLZ 5	99	
27173133	**Good identification**					**27173133**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27173140	**Low discrimination**					**27173140**
Propionibacterium propionicum		81.23	0.58	CEL 1		
Actinomyces naeslundii		16.11	0.57			
Actinomyces israelii		2.66	0.34			
27173141	**Low discrimination**					**27173141**
Actinomyces naeslundii		72.81	0.60	RAF 92		
Propionibacterium propionicum		27.20	0.42			
27173142	**Low discrimination**					**27173142**
Propionibacterium propionicum		85.84	0.61	CEL 1		
Actinomyces naeslundii		11.35	0.57			
Actinomyces israelii		2.81	0.37			
27173143	**Low discrimination**					**27173143**
Actinomyces naeslundii		64.09	0.60	RAF 92		
Propionibacterium propionicum		35.91	0.45			
27173151	**Good identification**					**27173151**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173153	**Good identification**					**27173153**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173160	**Low discrimination**					**27173160**
Actinomyces naeslundii		80.79	0.57	RAF 92		
Clostridium perfringens		14.24	0.21			
Propionibacterium propionicum		4.12	0.24			
27173161	**Very good identification**					**27173161**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27173162	**Low discrimination**					**27173162**
Actinomyces naeslundii		85.09	0.57	RAF 92		
Clostridium perfringens		7.06	0.16			
Propionibacterium propionicum		6.50	0.27			
27173163	**Very good identification**					**27173163**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27173171	**Good identification**					**27173171**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173173	**Good identification**					**27173173**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173300	**Low discrimination**					**27173300**
Actinomyces naeslundii		40.84	0.35	RAF 92		
Propionibacterium propionicum		36.34	0.24	RHA 1		
Actinomyces israelii		22.82	0.22			
27173301	**Good identification**					**27173301**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173303	**Good identification**					**27173303**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173320	**Low discrimination**					**27173320**
					SPO	
Actinomyces naeslundii		54.60	0.35	RAF 92	1	
Clostridium ramosum		45.40	0.09	RHA 1	92	
27173321	**Good identification**					**27173321**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173323	**Good identification**					**27173323**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173500	**Low discrimination**					**27173500**
Actinomyces israelii		76.27	0.40	SAL 92		
Actinomyces naeslundii		12.56	0.35	XYL 92		
Propionibacterium propionicum		11.17	0.24			
27173501	**Good identification**					**27173501**
Actinomyces naeslundii		100.0	0.38	RAF 92		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27173502	**Low discrimination**					**27173502**
Actinomyces israelii		79.60	0.43	SAL 92		
Propionibacterium propionicum		11.66	0.27	XYL 92		
Actinomyces naeslundii		8.74	0.35			
27173503	**Good identification**					**27173503**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173520	**Low discrimination**					**27173520**
Actinomyces naeslundii		72.06	0.35	RAF 92		
Actinomyces israelii		27.94	0.19	ARA 1		
27173521	**Good identification**					**27173521**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173522	**Low discrimination**					**27173522**
Actinomyces naeslundii		63.22	0.35	RAF 92		
Actinomyces israelii		36.78	0.22	ARA 1		
27173523	**Good identification**					**27173523**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27173542	**Low discrimination**					**27173542**
Actinomyces israelii		70.10	0.36	SAL 92		
Propionibacterium propionicum		26.41	0.27	XYL 92		
Actinomyces naeslundii		3.49	0.22			
27175002	**Low discrimination**					**27175002**
					COC	
Actinomyces israelii		74.10	0.39	SAL 92	1	
Atopobium parvulum		9.27	0.10		99	
Actinomyces viscosus		9.10	0.08		1	
27175100	**Very good identification**					**27175100**
					COC	
Atopobium parvulum		96.10	0.57	NIT 1	99	
Actinomyces israelii		3.65	0.46		1	
Actinomyces naeslundii		0.13	0.30		1	
27175102	**Low discrimination**					**27175102**
					COC	
Atopobium parvulum		74.00	0.44	NIT 1	99	
Actinomyces israelii		23.91	0.49		1	
Actinomyces viscosus		1.31	0.13		1	
27175112	**Low discrimination**					**27175112**
					COC	
Actinomyces israelii		79.06	0.35	SAL 92	1	
Atopobium parvulum		16.54	0.10	MLZ 13	99	
Bifidobacterium adolescentis		4.40	0.07		1	
27175120	**Low discrimination**					**27175120**
					COC	
Atopobium parvulum		79.86	0.44	NIT 1	99	
Clostridium perfringens		18.45	0.34		1	
Actinomyces israelii		1.10	0.26		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27175140	**Low discrimination**						**27175140**
					COC		
Actinomyces israelii		59.40	0.39	SAL 92	1		
Atopobium parvulum		40.60	0.23		99		
27175142	**Acceptable identification**						**27175142**
					COC		
Actinomyces israelii		92.56	0.42	SAL 92	1		
Atopobium parvulum		7.44	0.10		99		
27175402	**Good identification**						**27175402**
Actinomyces israelii		100.0	0.37	SAL 92			
27175500	**Low discrimination**						**27175500**
					COC	SPO	
Actinomyces israelii		74.30	0.45	SAL 92	1	1	
Atopobium parvulum		24.14	0.23		99	1	
Clostridium butyricum		1.56	-.05		1	99	
27175502	**Acceptable identification**						**27175502**
					COC		
Actinomyces israelii		91.42	0.48	SAL 92	1		
Bifidobacterium adolescentis		5.09	0.20		1		
Atopobium parvulum		3.49	0.10		99		
27175540	**Good identification**						**27175540**
Actinomyces israelii		100.0	0.38	SAL 92			
27175542	**Good identification**						**27175542**
Actinomyces israelii		100.0	0.41	SAL 92			
27177000	**Good identification**						**27177000**
Actinomyces israelii		98.38	0.49	SAL 92			
Actinomyces naeslundii		1.62	0.28				
27177002	**Good identification**						**27177002**
Actinomyces israelii		98.92	0.52	SAL 92			
Actinomyces naeslundii		1.09	0.28				
27177012	**Good identification**						**27177012**
Actinomyces israelii		100.0	0.38	SAL 92			
27177040	**Good identification**						**27177040**
Actinomyces israelii		100.0	0.42	SAL 92			
27177042	**Good identification**						**27177042**
Actinomyces israelii		100.0	0.45	SAL 92			
27177100	**Good identification**						**27177100**
					COC		
Actinomyces israelii		93.23	0.59	SAL 92	1		
Atopobium parvulum		4.37	0.23		99		
Actinomyces naeslundii		1.09	0.35		1		
27177101	**Low discrimination**						**27177101**
Actinomyces naeslundii		63.49	0.38	RAF 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		36.51	0.25	XYL 1		
27177102	**Very good identification**					**27177102**
Actinomyces israelii		97.00	0.62	SAL 92		
Propionibacterium propionicum		1.01	0.27			
Actinomyces naeslundii		0.76	0.35			
27177103	**Low discrimination**					**27177103**
Actinomyces naeslundii		53.69	0.38	RAF 92		
Actinomyces israelii		46.31	0.28	XYL 1		
27177110	**Good identification**					**27177110**
					SPO	
Actinomyces israelii		95.91	0.45	SAL 92	1	
Clostridium tertium		4.09	0.07	MLZ 13	99	
27177112	**Acceptable identification**					**27177112**
					SPO	
Actinomyces israelii		92.13	0.48	SAL 92	1	
Clostridium tertium		7.87	0.15	MLZ 13	99	
27177120	**Low discrimination**					**27177120**
					SPO	
Actinomyces israelii		67.49	0.39	SAL 92	1	
Actinomyces naeslundii		12.38	0.35	NAG 6	1	
Clostridium tertium		11.38	0.11		99	
27177121	**Good identification**					**27177121**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27177122	**Low discrimination**					**27177122**
					SPO	
Actinomyces israelii		68.51	0.42	SAL 92	1	
Clostridium tertium		23.11	0.19	NAG 6	99	
Actinomyces naeslundii		8.38	0.35		1	
27177123	**Good identification**					**27177123**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27177140	**Good identification**					**27177140**
Actinomyces israelii		96.89	0.52	SAL 92		
Propionibacterium propionicum		2.60	0.24			
Actinomyces naeslundii		0.52	0.22			
27177142	**Good identification**					**27177142**
Actinomyces israelii		97.06	0.55	SAL 92		
Propionibacterium propionicum		2.60	0.27			
Actinomyces naeslundii		0.34	0.22			
27177150	**Good identification**					**27177150**
Actinomyces israelii		100.0	0.38	SAL 92		
27177152	**Good identification**					**27177152**
Actinomyces israelii		100.0	0.41	SAL 92		
27177300	**Good identification**					**27177300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.40	SAL 92		
27177302	**Good identification**					**27177302**
Actinomyces israelii		100.0	0.43	SAL 92		
27177342	**Good identification**					**27177342**
Actinomyces israelii		100.0	0.36	SAL 92		
27177400	**Good identification**					**27177400**
Actinomyces israelii		100.0	0.47	SAL 92		
27177402	**Good identification**					**27177402**
Actinomyces israelii		100.0	0.50	SAL 92		
27177412	**Good identification**					**27177412**
Actinomyces israelii		100.0	0.36	SAL 92		
27177440	**Good identification**					**27177440**
Actinomyces israelii		100.0	0.40	SAL 92		
27177442	**Good identification**					**27177442**
Actinomyces israelii		100.0	0.43	SAL 92		
27177500	**Very good identification**					**27177500**
Actinomyces israelii		100.0	0.58	SAL 92		
27177502	**Very good identification**					**27177502**
Actinomyces israelii		99.20	0.61	SAL 92		
Bifidobacterium adolescentis		0.80	0.19			
27177510	**Good identification**					**27177510**
Actinomyces israelii		100.0	0.44	SAL 92		
27177512	**Acceptable identification**					**27177512**
Actinomyces israelii		93.84	0.47	SAL 92		
Bifidobacterium adolescentis		6.16	0.20	MLZ 13		
27177520	**Good identification**					**27177520**
Actinomyces israelii		100.0	0.37	SAL 92		
27177522	**Good identification**					**27177522**
Actinomyces israelii		100.0	0.40	SAL 92		
27177540	**Good identification**					**27177540**
Actinomyces israelii		100.0	0.51	SAL 92		
27177542	**Good identification**					**27177542**
Actinomyces israelii		100.0	0.54	SAL 92		
27177550	**Good identification**					**27177550**
Actinomyces israelii		100.0	0.37	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27177552	**Good identification**						**27177552**
Actinomyces israelii		100.0	0.40	SAL 92			
27177700	**Good identification**						**27177700**
Actinomyces israelii		100.0	0.38	SAL 92			
27177702	**Good identification**						**27177702**
Actinomyces israelii		100.0	0.41	SAL 92			
27210000	**Low discrimination**						**27210000**
					COC	GRA	
Actinomyces odontolyticus		49.84	0.44	MNS 9	1	99	
Fusobacterium mortiferum		45.02	0.49	MLT 99	1	1	
Gemella morbillorum		3.25	0.19		99	99	
27210002	**Low discrimination**						**27210002**
					GRA		
Actinomyces odontolyticus		51.11	0.40	MNS 9	99		
Actinomyces viscosus		39.74	0.31	MLT 99	99		
Fusobacterium mortiferum		8.37	0.32		1		
27210100	**Low discrimination**						**27210100**
					GRA		
Fusobacterium mortiferum		86.45	0.55	NIT 1	1		
Actinomyces odontolyticus		12.95	0.35		99		
Actinomyces viscosus		0.35	0.01		99		
27210102	**Low discrimination**						**27210102**
					GRA		
Actinomyces viscosus		66.14	0.36	MLT 99	99		
Fusobacterium mortiferum		18.29	0.39	RAF 94	1		
Actinomyces odontolyticus		15.11	0.32		99		
27210200	**Good identification**						**27210200**
					GRA		
Actinomyces odontolyticus		99.53	0.49	MNS 9	99		
Fusobacterium mortiferum		0.47	0.14	MLT 99	1		
27210202	**Good identification**						**27210202**
Actinomyces odontolyticus		100.0	0.45	MNS 9			
27210220	**Good identification**						**27210220**
Actinomyces odontolyticus		100.0	0.35	MNS 9			
27210300	**Good identification**						**27210300**
					GRA		
Actinomyces odontolyticus		96.64	0.40	MNS 9	99		
Fusobacterium mortiferum		3.36	0.21	MLT 99	1		
27210302	**Good identification**						**27210302**
Actinomyces odontolyticus		97.77	0.36	MNS 9			
Actinomyces viscosus		2.23	0.01	MLT 99			
27211000	**Low discrimination**						**27211000**
					GRA		
Fusobacterium mortiferum		61.11	0.52	NIT 1	1		
Actinomyces odontolyticus		30.07	0.41		99		
Actinomyces naeslundii		2.25	0.39		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27211001	**Low discrimination**					**27211001**
					GRA	
Actinomyces naeslundii		78.57	0.42	MLT 94	99	
Fusobacterium mortiferum		14.36	0.17	RAF 92	1	
Actinomyces odontolyticus		7.07	0.07		99	
27211002	**Low discrimination**					**27211002**
					GRA	
Actinomyces odontolyticus		42.63	0.37	MNS 9	99	
Actinomyces viscosus		30.48	0.28	MLT 99	99	
Fusobacterium mortiferum		15.71	0.35		1	
27211021	**Good identification**					**27211021**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27211023	**Good identification**					**27211023**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27211100	**Low discrimination**					**27211100**
					GRA	
Fusobacterium mortiferum		87.71	0.58	NIT 1	1	
Actinomyces odontolyticus		5.84	0.32		99	
Actinomyces naeslundii		3.94	0.46		99	
27211103	**Low discrimination**					**27211103**
					SPO	
Actinomyces naeslundii		48.73	0.49	MLT 94	1	
Actinomyces viscosus		39.58	0.24	RAF 92	1	
Clostridium chauvoei		11.69	0.16		99	
27211120	**Low discrimination**					**27211120**
Clostridium perfringens		88.01	0.42	MLT 99		
Actinomyces naeslundii		7.56	0.46			
Actinomyces odontolyticus		1.82	0.19			
27211121	**Low discrimination**					**27211121**
					SPO	
Actinomyces naeslundii		65.30	0.49	MLT 94	1	
Clostridium chauvoei		29.58	0.21	RAF 92	99	
Clostridium perfringens		5.12	0.08		84	
27211122	**Low discrimination**					**27211122**
					SPO	
Clostridium perfringens		79.18	0.37	MLT 99	84	
Actinomyces naeslundii		14.45	0.46		1	
Clostridium barati		2.22	0.06		99	
27211123	**Low discrimination**					**27211123**
					SPO	
Actinomyces naeslundii		89.54	0.49	MLT 94	1	
Clostridium chauvoei		7.16	0.08	RAF 92	99	
Clostridium perfringens		3.31	0.03		84	
27211141	**Good identification**					**27211141**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27211143	**Good identification**					**27211143**
Actinomyces naeslundii		100.0	0.37	MLT 94		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27211161	**Good identification**					**27211161**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27211163	**Good identification**					**27211163**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27211200	**Good identification**					**27211200**
					GRA	
Actinomyces odontolyticus		98.95	0.46	MNS 9	99	
Fusobacterium mortiferum		1.05	0.17	MLT 99	1	
27211202	**Good identification**					**27211202**
Actinomyces odontolyticus		100.0	0.42	MNS 9		
27211300	**Acceptable identification**					**27211300**
					GRA	
Actinomyces odontolyticus		92.75	0.37	MNS 9	99	
Fusobacterium mortiferum		7.25	0.24	MLT 99	1	
27213001	**Good identification**					**27213001**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27213003	**Good identification**					**27213003**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27213020	**Low discrimination**					**27213020**
Actinomyces naeslundii		66.09	0.44	MLT 94		
Propionibacterium acnes		29.05	0.13	RAF 92		
Propionibacterium granulosum		4.86	0.09			
27213021	**Good identification**					**27213021**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27213022	**Good identification**					**27213022**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27213023	**Good identification**					**27213023**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27213100	**Low discrimination**					**27213100**
					GRA	
Actinomyces naeslundii		80.60	0.51	MLT 94	99	
Propionibacterium propionicum		10.07	0.26	RAF 92	99	
Fusobacterium mortiferum		9.33	0.24		1	
27213101	**Good identification**					**27213101**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27213102	**Low discrimination**					**27213102**
Actinomyces naeslundii		84.22	0.51	MLT 94		
Propionibacterium propionicum		15.78	0.29	RAF 92		
27213103	**Good identification**					**27213103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		98.63	0.54	MLT 94			
Propionibacterium propionicum		1.37	0.12	RAF 92			
27213120	**Acceptable identification**						**27213120**
Actinomyces naeslundii		94.29	0.51	MLT 94			
Clostridium perfringens		5.71	0.08	RAF 92			
27213121	**Good identification**						**27213121**
Actinomyces naeslundii		100.0	0.54	MLT 94			
27213122	**Good identification**						**27213122**
Actinomyces naeslundii		97.23	0.51	MLT 94			
Clostridium perfringens		2.77	0.03	RAF 92			
27213123	**Good identification**						**27213123**
Actinomyces naeslundii		100.0	0.54	MLT 94			
27213140	**Low discrimination**						**27213140**
Actinomyces naeslundii		58.55	0.39	MLT 94			
Propionibacterium propionicum		41.45	0.26	RAF 92			
27213141	**Good identification**						**27213141**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27213143	**Acceptable identification**						**27213143**
Actinomyces naeslundii		92.71	0.42	MLT 94			
Propionibacterium propionicum		7.29	0.12	RAF 92			
27213160	**Good identification**						**27213160**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27213161	**Good identification**						**27213161**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27213162	**Good identification**						**27213162**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27213163	**Good identification**						**27213163**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27214000	**Low discrimination**						**27214000**
					COC	GRA	
Actinomyces odontolyticus		86.75	0.38	MNS 9	1	99	
Fusobacterium mortiferum		11.98	0.29	MLT 99	1	1	
Gemella morbillorum		1.28	0.03		99	99	
27214100	**Low discrimination**						**27214100**
					GRA		
Fusobacterium mortiferum		50.50	0.36	NIT 1	1		
Actinomyces odontolyticus		49.50	0.30	XYL 7	99		
27214200	**Good identification**						**27214200**
Actinomyces odontolyticus		100.0	0.43	MNS 9			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
27214202	**Good identification**						**27214202**
Actinomyces odontolyticus		100.0	0.40	MNS 9			
27215000	**Low discrimination**						**27215000**
					GRA		
Actinomyces odontolyticus		76.30	0.35	MNS 9	99		
Fusobacterium mortiferum		23.70	0.32	MLT 99	1		
27215100	**Low discrimination**						**27215100**
					GRA		
Fusobacterium mortiferum		69.65	0.39	NIT 1	1		
Actinomyces odontolyticus		30.35	0.27	XYL 7	99		
27215200	**Good identification**						**27215200**
Actinomyces odontolyticus		100.0	0.40	MNS 9			
27215202	**Good identification**						**27215202**
Actinomyces odontolyticus		100.0	0.37	MNS 9			
27223101	**Good identification**						**27223101**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27223103	**Good identification**						**27223103**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27223121	**Good identification**						**27223121**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27223123	**Good identification**						**27223123**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27230000	**Low discrimination**						**27230000**
					GRA	CAT	
Fusobacterium mortiferum		68.60	0.40	NIT 1	1	5	
Propionibacterium granulosum		28.84	0.30		99	90	
Actinomyces odontolyticus		2.57	0.10		99	10	
27230100	**Good identification**						**27230100**
					GRA		
Fusobacterium mortiferum		98.21	0.46	NIT 1	1		
Actinomyces naeslundii		1.79	0.28		99		
27231000	**Low discrimination**						**27231000**
					GRA	CAT	
Fusobacterium mortiferum		55.29	0.43	NIT 1	1	5	
Propionibacterium granulosum		15.50	0.30		99	90	
Lactobacillus jensenii		13.29	0.22		99	1	
27231001	**Good identification**						**27231001**
Actinomyces naeslundii		100.0	0.47	MLT 94			
27231002	**Low discrimination**						**27231002**
					GRA		
Actinomyces naeslundii		36.82	0.44	MLT 94	99		
Lactobacillus jensenii		24.65	0.19	RAF 92	99		
Fusobacterium mortiferum		17.09	0.26		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27231003	**Good identification**						**27231003**
Actinomyces naeslundii		95.20	0.47	MLT 94			
Actinomyces viscosus		4.80	0.00	RAF 92			
27231021	**Good identification**						**27231021**
Actinomyces naeslundii		100.0	0.47	MLT 94			
27231023	**Good identification**						**27231023**
Actinomyces naeslundii		100.0	0.47	MLT 94			
27231100	**Low discrimination**						**27231100**
					GRA		
Fusobacterium mortiferum		69.75	0.49	NIT 1	1		
Actinomyces naeslundii		20.37	0.51		99		
Clostridium perfringens		7.02	0.21		99		
27231101	**Acceptable identification**						**27231101**
					GRA	SPO	
Actinomyces naeslundii		90.84	0.54	MLT 94	99	1	
Clostridium chauvoei		7.07	0.12	RAF 92	99	99	
Fusobacterium mortiferum		2.10	0.15		1	1	
27231102	**Low discrimination**						**27231102**
					GRA		
Actinomyces naeslundii		54.63	0.51	MLT 94	99		
Fusobacterium mortiferum		20.79	0.33	RAF 92	1		
Clostridium perfringens		8.86	0.16		99		
27231103	**Good identification**						**27231103**
					SPO		
Actinomyces naeslundii		95.67	0.54	MLT 94	1		
Actinomyces viscosus		3.01	0.05	RAF 92	1		
Clostridium chauvoei		1.31	-.01		99		
27231120	**Acceptable identification**						**27231120**
					GRA		
Clostridium perfringens		91.09	0.47	MLT 99	99		
Actinomyces naeslundii		8.18	0.51		99		
Fusobacterium mortiferum		0.28	0.15		1		
27231121	**Acceptable identification**						**27231121**
					SPO		
Actinomyces naeslundii		90.83	0.54	MLT 94	1		
Clostridium perfringens		6.82	0.13	RAF 92	84		
Clostridium chauvoei		2.36	0.04		99		
27231122	**Low discrimination**						**27231122**
Clostridium perfringens		82.38	0.41	MLT 99			
Actinomyces naeslundii		15.71	0.51				
Lactobacillus acidophilus		1.91	0.11				
27231123	**Good identification**						**27231123**
Actinomyces naeslundii		96.59	0.54	MLT 94			
Clostridium perfringens		3.41	0.07	RAF 92			
27231140	**Low discrimination**						**27231140**
					GRA		
Actinomyces naeslundii		63.78	0.39	MLT 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		23.72	0.09	RAF 92	99		
Fusobacterium mortiferum		12.50	0.15		1		
27231141	**Good identification**						**27231141**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27231142	**Low discrimination**						**27231142**
Actinomyces naeslundii		85.10	0.39	MLT 94			
Clostridium perfringens		14.90	0.03	RAF 92			
27231143	**Good identification**						**27231143**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27231161	**Acceptable identification**						**27231161**
Actinomyces naeslundii		92.51	0.42	MLT 94			
Clostridium perfringens		7.49	0.01	RAF 92			
27231163	**Good identification**						**27231163**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27232101	**Good identification**						**27232101**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27232103	**Good identification**						**27232103**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27232121	**Good identification**						**27232121**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27232123	**Good identification**						**27232123**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27233000	**Low discrimination**						**27233000**
					SPO	CAT	
Actinomyces naeslundii		41.81	0.49	MLT 94	1	25	
Clostridium innocuum		39.17	0.24	RAF 92	99	1	
Propionibacterium granulosum		19.02	0.28		1	90	
27233001	**Good identification**						**27233001**
Actinomyces naeslundii		100.0	0.52	MLT 94			
27233002	**Low discrimination**						**27233002**
					SPO	CAT	
Actinomyces naeslundii		48.77	0.49	MLT 94	1	25	
Clostridium innocuum		45.69	0.24	RAF 92	99	1	
Propionibacterium granulosum		5.55	0.18		1	90	
27233003	**Good identification**						**27233003**
Actinomyces naeslundii		100.0	0.52	MLT 94			
27233020	**Low discrimination**						**27233020**
Actinomyces naeslundii		89.79	0.49	MLT 94			
Propionibacterium granulosum		10.21	0.18	RAF 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27233021	**Good identification**					**27233021**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27233022	**Good identification**					**27233022**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27233023	**Good identification**					**27233023**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27233040	**Good identification**					**27233040**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27233041	**Good identification**					**27233041**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27233042	**Good identification**					**27233042**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27233043	**Good identification**					**27233043**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27233060	**Good identification**					**27233060**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27233061	**Good identification**					**27233061**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27233062	**Good identification**					**27233062**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27233063	**Good identification**					**27233063**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27233100	**Good identification**					**27233100**
					GRA	
Actinomyces naeslundii		93.42	0.56	MLT 94	99	
Propionibacterium propionicum		4.92	0.24	RAF 92	99	
Fusobacterium mortiferum		1.67	0.15		1	
27233101	**Very good identification**					**27233101**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27233102	**Good identification**					**27233102**
Actinomyces naeslundii		92.68	0.56	MLT 94		
Propionibacterium propionicum		7.32	0.27	RAF 92		
27233103	**Very good identification**					**27233103**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27233111	**Good identification**					**27233111**
Actinomyces naeslundii		100.0	0.37	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27233113	**Good identification**					**27233113**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27233120	**Good identification**					**27233120**
Actinomyces naeslundii		94.52	0.56	MLT 94		
Clostridium perfringens		5.48	0.13	RAF 92		
27233121	**Very good identification**					**27233121**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27233122	**Very good identification**					**27233122**
Actinomyces naeslundii		97.34	0.56	MLT 94		
Clostridium perfringens		2.66	0.07	RAF 92		
27233123	**Very good identification**					**27233123**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27233131	**Good identification**					**27233131**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27233133	**Good identification**					**27233133**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27233140	**Low discrimination**					**27233140**
Actinomyces naeslundii		77.02	0.43	MLT 94		
Propionibacterium propionicum		22.98	0.24	RAF 92		
27233141	**Good identification**					**27233141**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27233142	**Low discrimination**					**27233142**
Actinomyces naeslundii		69.08	0.43	MLT 94		
Propionibacterium propionicum		30.92	0.27	RAF 92		
27233143	**Good identification**					**27233143**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27233160	**Acceptable identification**					**27233160**
Actinomyces naeslundii		94.11	0.43	MLT 94		
Clostridium perfringens		5.89	0.01	RAF 92		
27233161	**Good identification**					**27233161**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27233162	**Good identification**					**27233162**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27233163	**Good identification**					**27233163**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27243000	**Good identification**					**27243000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium innocuum		99.74	0.44	NIT 1	99		
Actinomyces naeslundii		0.26	0.24	FRU 95	1		
27243002	**Good identification**						**27243002**
					SPO		
Clostridium innocuum		99.74	0.44	NIT 1	99		
Actinomyces naeslundii		0.26	0.24	FRU 95	1		
27250000	**Low discrimination**						**27250000**
					GRA	SPO	
Fusobacterium mortiferum		81.38	0.40	NIT 1	1	1	
Clostridium innocuum		7.13	0.10		99	99	
Lactobacillus catenaformis		7.07	0.12		99	1	
27250100	**Acceptable identification**						**27250100**
					GRA	SPO	
Fusobacterium mortiferum		92.16	0.47	NIT 1	1	1	
Lactobacillus catenaformis		5.15	0.15		99	1	
Clostridium barati		1.66	0.02		99	99	
27251000	**Low discrimination**						**27251000**
					SPO		
Lactobacillus catenaformis		60.07	0.47	NIT 1	1		
Lactobacillus jensenii		14.29	0.36	MLT 90	1		
Clostridium innocuum		11.63	0.32		99		
27251001	**Low discrimination**						**27251001**
Actinomyces naeslundii		75.39	0.44	MLT 94			
Lactobacillus catenaformis		19.88	0.12	RAF 92			
Lactobacillus jensenii		4.73	0.02				
27251002	**Low discrimination**						**27251002**
					SPO		
Lactobacillus catenaformis		64.40	0.47	NIT 1	1		
Clostridium innocuum		12.46	0.32	MLT 90	99		
Lactobacillus jensenii		10.21	0.33		1		
27251003	**Low discrimination**						**27251003**
Actinomyces naeslundii		65.60	0.44	MLT 94			
Lactobacillus catenaformis		17.30	0.12	RAF 92			
Actinomyces viscosus		17.10	0.10				
27251020	**Low discrimination**						**27251020**
					SPO		
Lactobacillus catenaformis		57.97	0.36	NIT 1	1		
Clostridium barati		21.49	0.24	MLT 90	99		
Lactobacillus jensenii		13.79	0.26		1		
27251021	**Acceptable identification**						**27251021**
Actinomyces naeslundii		93.82	0.44	MLT 94			
Lactobacillus catenaformis		6.19	0.02	RAF 92			
27251023	**Low discrimination**						**27251023**
					SPO		
Actinomyces naeslundii		87.78	0.44	MLT 94	1		
Clostridium barati		6.44	-.02	RAF 92	99		
Lactobacillus catenaformis		5.79	0.02		1		
27251100	**Low discrimination**						**27251100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Lactobacillus catenaformis		71.83	0.50	NIT 1	99	1	
Fusobacterium mortiferum		19.48	0.50	MLT 90	1	1	
Clostridium barati		4.44	0.24		99	99	
27251101	**Low discrimination**						**27251101**
					GRA		
Actinomyces naeslundii		84.99	0.52	MLT 94	99		
Lactobacillus catenaformis		11.81	0.15	RAF 92	99		
Fusobacterium mortiferum		3.20	0.16		1		
27251102	**Low discrimination**						**27251102**
					SPO		
Lactobacillus catenaformis		64.35	0.50	NIT 1	1		
Lactobacillus acidophilus		16.38	0.36	MLT 90	1		
Clostridium barati		11.93	0.32		99		
27251103	**Low discrimination**						**27251103**
Actinomyces naeslundii		73.37	0.52	MLT 94			
Actinomyces viscosus		11.95	0.14	RAF 92			
Lactobacillus catenaformis		10.20	0.15				
27251120	**Low discrimination**						**27251120**
					SPO		
Lactobacillus catenaformis		48.75	0.39	NIT 1	1		
Clostridium barati		36.14	0.32	MLT 90	99		
Actinomyces naeslundii		9.45	0.49		1		
27251121	**Acceptable identification**						**27251121**
					SPO		
Actinomyces naeslundii		94.30	0.52	MLT 94	1		
Lactobacillus catenaformis		3.28	0.05	RAF 92	1		
Clostridium barati		2.43	-.02		99		
27251122	**Low discrimination**						**27251122**
					SPO		
Clostridium barati		49.58	0.40	NIT 1	99		
Lactobacillus acidophilus		22.70	0.36	ESL 87	1		
Lactobacillus catenaformis		22.29	0.39		1		
27251123	**Low discrimination**						**27251123**
					SPO		
Actinomyces naeslundii		87.15	0.52	MLT 94	1		
Clostridium barati		6.74	0.06	RAF 92	99		
Lactobacillus acidophilus		3.09	0.02		1		
27251141	**Good identification**						**27251141**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27251143	**Good identification**						**27251143**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27251161	**Good identification**						**27251161**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27251163	**Good identification**						**27251163**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27252000	**Good identification**						**27252000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium innocuum		100.0	0.44	NIT 1		
27252002	**Good identification**					**27252002**
Clostridium innocuum		100.0	0.44	NIT 1		
27253000	**Very good identification**					**27253000**
					SPO	
Clostridium innocuum		99.66	0.66	NIT 1	99	
Actinomyces naeslundii		0.26	0.46		1	
Lactobacillus catenaformis		0.05	0.12		1	
27253002	**Very good identification**					**27253002**
					SPO	
Clostridium innocuum		99.65	0.66	NIT 1	99	
Actinomyces naeslundii		0.26	0.46		1	
Lactobacillus catenaformis		0.05	0.12		1	
27253010	**Good identification**					**27253010**
					SPO	
Clostridium innocuum		99.67	0.42	NIT 1	99	
Actinomyces naeslundii		0.33	0.24	MLZ 4	1	
27253012	**Good identification**					**27253012**
					SPO	
Clostridium innocuum		99.67	0.42	NIT 1	99	
Actinomyces naeslundii		0.33	0.24	MLZ 4	1	
27253021	**Good identification**					**27253021**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27253023	**Good identification**					**27253023**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27253041	**Good identification**					**27253041**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27253043	**Good identification**					**27253043**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27253061	**Good identification**					**27253061**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27253063	**Good identification**					**27253063**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27253101	**Very good identification**					**27253101**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27253103	**Very good identification**					**27253103**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27253120	**Low discrimination**					**27253120**
					SPO	
Actinomyces naeslundii		83.46	0.53	MLT 94	1	
Clostridium barati		14.30	0.14	RAF 92	99	
Lactobacillus catenaformis		2.24	0.05		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27253121	**Very good identification**					**27253121**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27253122	**Low discrimination**					**27253122**
					SPO	
Actinomyces naeslundii		58.73	0.53	MLT 94	1	
Clostridium barati		30.19	0.22	RAF 92	99	
Lactobacillus acidophilus		8.37	0.15		1	
27253123	**Very good identification**					**27253123**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27253140	**Good identification**					**27253140**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27253141	**Good identification**					**27253141**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27253142	**Acceptable identification**					**27253142**
Actinomyces naeslundii		90.93	0.41	MLT 94		
Lactobacillus acidophilus		9.07	-.01	RAF 92		
27253143	**Good identification**					**27253143**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27253160	**Good identification**					**27253160**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27253161	**Good identification**					**27253161**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27253162	**Low discrimination**					**27253162**
					SPO	
Actinomyces naeslundii		73.91	0.41	MLT 94	1	
Clostridium barati		18.72	0.04	RAF 92	99	
Lactobacillus acidophilus		7.37	-.01		1	
27253163	**Good identification**					**27253163**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27253400	**Good identification**					**27253400**
Clostridium innocuum		100.0	0.53	NIT 1		
27253402	**Good identification**					**27253402**
Clostridium innocuum		100.0	0.53	NIT 1		
27257000	**Good identification**					**27257000**
					SPO	
Clostridium innocuum		99.21	0.44	NIT 1	99	
Actinomyces israelii		0.79	0.23	XYL 5	1	
27257002	**Good identification**					**27257002**
					SPO	
Clostridium innocuum		98.82	0.44	NIT 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.18	0.26	XYL 5	1		
27257102	**Low discrimination**						**27257102**
					SPO		
Actinomyces israelii		82.60	0.36	MLT 99	1		
Clostridium innocuum		17.40	0.10		99		
27263000	**Good identification**						**27263000**
					SPO		
Clostridium innocuum		98.59	0.36	NIT 1	99		
Actinomyces naeslundii		1.41	0.29	FRU 95	1		
27263002	**Good identification**						**27263002**
					SPO		
Clostridium innocuum		98.59	0.36	NIT 1	99		
Actinomyces naeslundii		1.41	0.29	FRU 95	1		
27263100	**Low discrimination**						**27263100**
					SPO		
Actinomyces naeslundii		80.11	0.37	MLT 94	1		
Clostridium innocuum		19.89	0.02	RAF 92	99		
27263101	**Good identification**						**27263101**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27263102	**Low discrimination**						**27263102**
					SPO		
Actinomyces naeslundii		80.11	0.37	MLT 94	1		
Clostridium innocuum		19.89	0.02	RAF 92	99		
27263103	**Good identification**						**27263103**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27263120	**Good identification**						**27263120**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27263121	**Good identification**						**27263121**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27263122	**Good identification**						**27263122**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27263123	**Good identification**						**27263123**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27270100	**Acceptable identification**						**27270100**
					GRA		
Fusobacterium mortiferum		93.23	0.38	NIT 1	1		
Actinomyces naeslundii		6.77	0.30		99		
27271000	**Low discrimination**						**27271000**
					GRA	SPO	
Lactobacillus jensenii		79.64	0.44	NIT 1	99	1	
Clostridium innocuum		7.59	0.24	MLT 95	99	99	
Fusobacterium mortiferum		5.81	0.34		1	1	
27271001	**Acceptable identification**						**27271001**
Actinomyces naeslundii		91.16	0.49	MLT 94			
Lactobacillus jensenii		8.84	0.10	RAF 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27271002	**Low discrimination**						**27271002**
					SPO		
Lactobacillus jensenii		65.10	0.41	NIT 1	1		
Lactobacillus acidophilus		15.22	0.27	MLT 95	1		
Clostridium innocuum		9.31	0.24		99		
27271003	**Acceptable identification**						**27271003**
Actinomyces naeslundii		93.93	0.49	MLT 94			
Lactobacillus jensenii		6.07	0.07	RAF 92			
27271010	**Good identification**						**27271010**
					SPO		
Lactobacillus jensenii		97.76	0.36	NIT 1	1		
Clostridium innocuum		1.17	0.00	MLT 95	99		
Actinomyces naeslundii		1.07	0.24		1		
27271021	**Good identification**						**27271021**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27271023	**Good identification**						**27271023**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27271041	**Good identification**						**27271041**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27271043	**Good identification**						**27271043**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27271061	**Good identification**						**27271061**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27271063	**Good identification**						**27271063**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27271100	**Low discrimination**						**27271100**
					COC	GRA	
Actinomyces naeslundii		41.39	0.53	MLT 94	1	99	
Fusobacterium mortiferum		35.65	0.41	RAF 92	1	1	
Atopobium parvulum		10.25	0.20		99	99	
27271101	**Very good identification**						**27271101**
Actinomyces naeslundii		100.0	0.57	MLT 94			
27271102	**Low discrimination**						**27271102**
					SPO		
Lactobacillus acidophilus		85.80	0.45	NIT 1	1		
Actinomyces naeslundii		9.47	0.53	MLT 94	1		
Clostridium barati		1.62	0.14		99		
27271103	**Good identification**						**27271103**
Actinomyces naeslundii		94.25	0.57	MLT 94			
Lactobacillus acidophilus		5.75	0.11	RAF 92			
27271120	**Low discrimination**						**27271120**
					SPO		
Actinomyces naeslundii		56.94	0.53	MLT 94	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		19.94	0.23	RAF 92	84	
Clostridium barati		9.76	0.14		99	
27271121	**Very good identification**					**27271121**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27271122	**Low discrimination**					**27271122**
					SPO	
Lactobacillus acidophilus		83.86	0.45	NIT 1	1	
Actinomyces naeslundii		9.25	0.53	MLT 94	1	
Clostridium barati		4.76	0.22		99	
27271123	**Good identification**					**27271123**
Actinomyces naeslundii		94.25	0.57	MLT 94		
Lactobacillus acidophilus		5.75	0.11	RAF 92		
27271140	**Good identification**					**27271140**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27271141	**Good identification**					**27271141**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27271143	**Good identification**					**27271143**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27271160	**Low discrimination**					**27271160**
Actinomyces naeslundii		72.57	0.41	MLT 94		
Clostridium perfringens		27.43	0.11	RAF 92		
27271161	**Good identification**					**27271161**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27271163	**Good identification**					**27271163**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27272000	**Good identification**					**27272000**
					SPO	
Clostridium innocuum		98.88	0.36	NIT 1	99	
Actinomyces naeslundii		1.12	0.27	GAL 95	1	
27272002	**Good identification**					**27272002**
					SPO	
Clostridium innocuum		98.88	0.36	NIT 1	99	
Actinomyces naeslundii		1.12	0.27	GAL 95	1	
27272101	**Good identification**					**27272101**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27272103	**Good identification**					**27272103**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27272121	**Good identification**					**27272121**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27272123	**Good identification**					**27272123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MLT 94		
27273000	**Very good identification**					**27273000**
					SPO	
Clostridium innocuum		98.45	0.58	NIT 1	99	
Actinomyces naeslundii		1.41	0.51		1	
Lactobacillus jensenii		0.11	0.10		1	
27273001	**Low discrimination**					**27273001**
					SPO	
Actinomyces naeslundii		67.98	0.54	MLT 94	1	
Clostridium innocuum		32.03	0.24	RAF 92	99	
27273002	**Very good identification**					**27273002**
					SPO	
Clostridium innocuum		98.38	0.58	NIT 1	99	
Actinomyces naeslundii		1.41	0.51		1	
Lactobacillus acidophilus		0.09	0.05		1	
27273003	**Low discrimination**					**27273003**
					SPO	
Actinomyces naeslundii		67.98	0.54	MLT 94	1	
Clostridium innocuum		32.03	0.24	RAF 92	99	
27273020	**Low discrimination**					**27273020**
					SPO	
Actinomyces naeslundii		58.59	0.51	MLT 94	1	
Clostridium innocuum		41.41	0.24	RAF 92	99	
27273021	**Good identification**					**27273021**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27273022	**Low discrimination**					**27273022**
					SPO	
Actinomyces naeslundii		56.58	0.51	MLT 94	1	
Clostridium innocuum		39.98	0.24	RAF 92	99	
Lactobacillus acidophilus		3.44	0.05		1	
27273023	**Good identification**					**27273023**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27273041	**Good identification**					**27273041**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27273043	**Good identification**					**27273043**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27273060	**Good identification**					**27273060**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27273061	**Good identification**					**27273061**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27273062	**Good identification**					**27273062**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27273063	**Good identification**					**27273063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MLT 94		
27273100	**Low discrimination**					**27273100**
					SPO	
Actinomyces naeslundii		77.12	0.58	MLT 94	1	
Clostridium innocuum		19.15	0.24	RAF 92	99	
Actinomyces israelii		2.97	0.25		1	
27273101	**Very good identification**					**27273101**
Actinomyces naeslundii		100.0	0.61	MLT 94		
27273102	**Low discrimination**					**27273102**
					SPO	
Actinomyces naeslundii		63.23	0.58	MLT 94	1	
Clostridium innocuum		15.70	0.24	RAF 92	99	
Lactobacillus acidophilus		15.54	0.24		1	
27273103	**Very good identification**					**27273103**
Actinomyces naeslundii		100.0	0.61	MLT 94		
27273110	**Low discrimination**					**27273110**
					SPO	
Actinomyces naeslundii		65.31	0.37	MLT 94	1	
Clostridium tertium		21.85	0.05	RAF 92	99	
Clostridium innocuum		12.84	0.00	MLZ 5	99	
27273111	**Good identification**					**27273111**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27273113	**Good identification**					**27273113**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27273120	**Good identification**					**27273120**
					SPO	
Actinomyces naeslundii		94.54	0.58	MLT 94	1	
Clostridium tertium		2.81	0.09	RAF 92	99	
Clostridium ramosum		2.65	0.07		92	
27273121	**Very good identification**					**27273121**
Actinomyces naeslundii		100.0	0.61	MLT 94		
27273122	**Low discrimination**					**27273122**
					SPO	
Actinomyces naeslundii		68.03	0.58	MLT 94	1	
Lactobacillus acidophilus		16.72	0.24	RAF 92	1	
Clostridium ramosum		7.62	0.17		92	
27273123	**Very good identification**					**27273123**
Actinomyces naeslundii		100.0	0.61	MLT 94		
27273131	**Good identification**					**27273131**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27273133	**Good identification**					**27273133**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27273140	**Acceptable identification**					**27273140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.18	0.46	MLT 94		
Actinomyces israelii		7.82	0.18	RAF 92		
27273141	**Good identification**					**27273141**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27273142	**Low discrimination**					**27273142**
Actinomyces naeslundii		76.96	0.46	MLT 94		
Lactobacillus acidophilus		13.25	0.08	RAF 92		
Actinomyces israelii		9.79	0.21			
27273143	**Good identification**					**27273143**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27273160	**Good identification**					**27273160**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27273161	**Good identification**					**27273161**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27273162	**Low discrimination**					**27273162**
Actinomyces naeslundii		85.32	0.46	MLT 94		
Lactobacillus acidophilus		14.68	0.08	RAF 92		
27273163	**Good identification**					**27273163**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27273400	**Good identification**					**27273400**
					SPO	
Clostridium innocuum		99.82	0.45	NIT 1	99	
Actinomyces israelii		0.18	0.13		1	
27273402	**Good identification**					**27273402**
					SPO	
Clostridium innocuum		99.73	0.45	NIT 1	99	
Actinomyces israelii		0.27	0.16		1	
27277000	**Acceptable identification**					**27277000**
					SPO	
Clostridium innocuum		92.12	0.36	NIT 1	99	
Actinomyces israelii		7.88	0.33	XYL 5	1	
27277002	**Low discrimination**					**27277002**
					SPO	
Clostridium innocuum		88.63	0.36	NIT 1	99	
Actinomyces israelii		11.37	0.36	XYL 5	1	
27277100	**Acceptable identification**					**27277100**
					SPO	
Actinomyces israelii		92.32	0.43	MLT 99	1	
Clostridium tertium		2.85	0.03		99	
Clostridium innocuum		2.73	0.02		99	
27277102	**Acceptable identification**					**27277102**
					SPO	
Actinomyces israelii		91.19	0.46	MLT 99	1	
Clostridium tertium		5.63	0.11		99	
Clostridium innocuum		1.79	0.02		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27277140	**Good identification**					**27277140**
Actinomyces israelii		100.0	0.36	MLT 99		
27277142	**Good identification**					**27277142**
Actinomyces israelii		100.0	0.39	MLT 99		
27277500	**Good identification**					**27277500**
Actinomyces israelii		100.0	0.42	MLT 99		
27277502	**Good identification**					**27277502**
Actinomyces israelii		100.0	0.45	MLT 99		
27277542	**Good identification**					**27277542**
Actinomyces israelii		100.0	0.38	MLT 99		
27300000	**Very good identification**					**27300000**
					COC	
Actinomyces odontolyticus		96.21	0.60	MNS 9	1	
Gemella morbillorum		3.38	0.30	FRU 92	99	
Actinomyces viscosus		0.28	0.09		1	
27300002	**Low discrimination**					**27300002**
Actinomyces odontolyticus		68.00	0.56	MNS 9		
Actinomyces viscosus		32.00	0.44	FRU 92		
27300003	**Acceptable identification**					**27300003**
Actinomyces viscosus		93.95	0.35	RAF 94		
Actinomyces odontolyticus		6.05	0.22	FRU 95		
27300020	**Good identification**					**27300020**
					COC	
Actinomyces odontolyticus		97.66	0.46	MNS 9	1	
Gemella morbillorum		2.34	0.14	FRU 92	99	
27300022	**Good identification**					**27300022**
Actinomyces odontolyticus		97.16	0.43	MNS 9		
Actinomyces viscosus		2.84	0.09	FRU 92		
27300100	**Good identification**					**27300100**
Actinomyces odontolyticus		98.39	0.51	MNS 9		
Actinomyces viscosus		1.61	0.14	FRU 92		
27300102	**Low discrimination**					**27300102**
Actinomyces viscosus		72.60	0.48	RAF 94		
Actinomyces odontolyticus		27.40	0.48	FRU 95		
27300103	**Good identification**					**27300103**
					SPO	
Actinomyces viscosus		96.97	0.40	RAF 94	1	
Clostridium chauvoei		1.44	0.10	FRU 95	99	
Actinomyces odontolyticus		1.11	0.13		1	
27300120	**Acceptable identification**					**27300120**
					GRA	
Actinomyces odontolyticus		91.48	0.38	MNS 9	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella bivia		5.07	-.03	FRU 92	1	
Clostridium perfringens		3.45	0.08	NAG 14	99	
27300200	**Very good identification**					**27300200**
Actinomyces odontolyticus		100.0	0.65	MNS 9		
27300202	**Very good identification**					**27300202**
Actinomyces odontolyticus		99.76	0.61	MNS 9		
Actinomyces viscosus		0.24	0.09	FRU 92		
27300220	**Good identification**					**27300220**
Actinomyces odontolyticus		100.0	0.51	MNS 9		
27300222	**Good identification**					**27300222**
Actinomyces odontolyticus		100.0	0.48	MNS 9		
27300300	**Very good identification**					**27300300**
Actinomyces odontolyticus		100.0	0.56	MNS 9		
27300302	**Good identification**					**27300302**
Actinomyces odontolyticus		98.64	0.52	MNS 9		
Actinomyces viscosus		1.36	0.14	FRU 92		
27300320	**Good identification**					**27300320**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27300322	**Good identification**					**27300322**
Actinomyces odontolyticus		100.0	0.39	MNS 9		
27301000	**Very good identification**					**27301000**
					COC	
Actinomyces odontolyticus		95.99	0.57	MNS 9	1	
Gemella morbillorum		2.73	0.26	FRU 92	99	
Actinomyces naeslundii		0.69	0.37		1	
27301002	**Low discrimination**					**27301002**
Actinomyces odontolyticus		69.24	0.53	MNS 9		
Actinomyces viscosus		29.96	0.40	FRU 92		
Actinomyces naeslundii		0.81	0.37			
27301020	**Low discrimination**					**27301020**
Actinomyces odontolyticus		76.63	0.43	MNS 9		
Clostridium perfringens		18.51	0.28	FRU 92		
Actinomyces naeslundii		3.38	0.37	NAG 14		
27301021	**Low discrimination**					**27301021**
					SPO	
Actinomyces naeslundii		54.86	0.40	RAF 92	1	
Clostridium chauvoei		36.76	0.14	FRU 95	99	
Actinomyces odontolyticus		8.39	0.09		1	
27301022	**Low discrimination**					**27301022**
Actinomyces odontolyticus		77.87	0.40	MNS 9		
Clostridium perfringens		14.44	0.22	FRU 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		5.60	0.37	NAG 14		
27301023	**Low discrimination**					**27301023**
					SPO	
Actinomyces naeslundii		82.51	0.40	RAF 92	1	
Clostridium chauvoei		9.76	0.01	FRU 95	99	
Actinomyces odontolyticus		7.73	0.05		1	
27301100	**Low discrimination**					**27301100**
					SPO	
Actinomyces odontolyticus		85.00	0.48	MNS 9	1	
Clostridium chauvoei		5.82	0.22	FRU 92	99	
Actinomyces naeslundii		5.49	0.45		1	
27301101	**Very good identification**					**27301101**
					SPO	
Clostridium chauvoei		98.37	0.57	SAL 1	99	
Actinomyces naeslundii		1.41	0.48		1	
Actinomyces odontolyticus		0.15	0.14		1	
27301102	**Low discrimination**					**27301102**
Actinomyces viscosus		67.99	0.44	RAF 94		
Actinomyces odontolyticus		27.91	0.45	FRU 95		
Actinomyces naeslundii		2.94	0.45			
27301103	**Low discrimination**					**27301103**
					SPO	
Clostridium chauvoei		66.57	0.44	SAL 1	99	
Actinomyces viscosus		27.69	0.36		1	
Actinomyces naeslundii		5.40	0.48		1	
27301120	**Low discrimination**					**27301120**
Clostridium perfringens		78.03	0.42	FRU 99		
Actinomyces odontolyticus		13.92	0.35			
Actinomyces naeslundii		5.53	0.45			
27301121	**Good identification**					**27301121**
					SPO	
Clostridium chauvoei		95.51	0.48	SAL 1	99	
Actinomyces naeslundii		4.10	0.48		1	
Clostridium perfringens		0.39	0.08		84	
27301122	**Low discrimination**					**27301122**
Clostridium perfringens		70.07	0.37	FRU 99		
Actinomyces odontolyticus		16.27	0.31			
Actinomyces naeslundii		10.54	0.45			
27301123	**Low discrimination**					**27301123**
					SPO	
Clostridium chauvoei		78.95	0.36	SAL 1	99	
Actinomyces naeslundii		19.19	0.48		1	
Actinomyces viscosus		1.00	0.02		1	
27301143	**Low discrimination**					**27301143**
					SPO	
Actinomyces naeslundii		50.00	0.35	RAF 92	1	
Clostridium chauvoei		35.31	0.10	FRU 95	99	
Actinomyces viscosus		14.69	0.02		1	
27301163	**Low discrimination**					**27301163**

Code Taxon		ID score	T index	tests against	Differentiation tests SPO	Code
Actinomyces naeslundii		80.94	0.35	RAF 92	1	
Clostridium chauvoei		19.06	0.01	FRU 95	99	
27301200	**Very good identification**					**27301200**
Actinomyces odontolyticus		100.0	0.62	MNS 9		
27301202	**Very good identification**					**27301202**
Actinomyces odontolyticus		99.78	0.58	MNS 9		
Actinomyces viscosus		0.23	0.06	FRU 92		
27301220	**Good identification**					**27301220**
Actinomyces odontolyticus		100.0	0.48	MNS 9		
27301222	**Good identification**					**27301222**
Actinomyces odontolyticus		100.0	0.45	MNS 9		
27301300	**Good identification**					**27301300**
Actinomyces odontolyticus		100.0	0.53	MNS 9		
27301302	**Good identification**					**27301302**
Actinomyces odontolyticus		98.75	0.49	MNS 9		
Actinomyces viscosus		1.25	0.10	FRU 92		
27301320	**Good identification**					**27301320**
Actinomyces odontolyticus		97.17	0.40	MNS 9		
Clostridium perfringens		2.84	0.08	FRU 92		
27301322	**Good identification**					**27301322**
Actinomyces odontolyticus		97.81	0.36	MNS 9		
Clostridium perfringens		2.19	0.03	FRU 92		
27303000	**Low discrimination**					**27303000**
Actinomyces naeslundii		57.95	0.42	RAF 92		
Actinomyces odontolyticus		42.05	0.23	FRU 95		
27303001	**Good identification**					**27303001**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27303002	**Low discrimination**					**27303002**
Actinomyces naeslundii		61.08	0.42	RAF 92		
Actinomyces odontolyticus		27.16	0.19	FRU 95		
Actinomyces viscosus		11.75	0.06			
27303003	**Good identification**					**27303003**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27303020	**Low discrimination**					**27303020**
Actinomyces naeslundii		89.44	0.42	RAF 92		
Actinomyces odontolyticus		10.56	0.09	FRU 95		
27303021	**Good identification**					**27303021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	RAF 92		
27303022	**Acceptable identification**					**27303022**
Actinomyces naeslundii		93.25	0.42	RAF 92		
Actinomyces odontolyticus		6.75	0.05	FRU 95		
27303023	**Good identification**					**27303023**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27303100	**Low discrimination**					**27303100**
Actinomyces naeslundii		81.17	0.50	RAF 92		
Propionibacterium propionicum		12.30	0.26	FRU 95		
Actinomyces odontolyticus		6.53	0.14			
27303101	**Low discrimination**					**27303101**
					SPO	
Actinomyces naeslundii		73.32	0.53	RAF 92	1	
Clostridium chauvoei		26.68	0.22	FRU 95	99	
27303102	**Low discrimination**					**27303102**
Actinomyces naeslundii		71.59	0.50	RAF 92		
Propionibacterium propionicum		16.27	0.29	FRU 95		
Actinomyces viscosus		8.60	0.10			
27303103	**Acceptable identification**					**27303103**
					SPO	
Actinomyces naeslundii		90.27	0.53	RAF 92	1	
Clostridium chauvoei		5.80	0.10	FRU 95	99	
Actinomyces viscosus		2.41	0.02		1	
27303120	**Acceptable identification**					**27303120**
Actinomyces naeslundii		93.15	0.50	RAF 92		
Clostridium perfringens		6.85	0.08	FRU 95		
27303121	**Low discrimination**					**27303121**
					SPO	
Actinomyces naeslundii		89.18	0.53	RAF 92	1	
Clostridium chauvoei		10.82	0.14	FRU 95	99	
27303122	**Good identification**					**27303122**
Actinomyces naeslundii		96.66	0.50	RAF 92		
Clostridium perfringens		3.34	0.03	FRU 95		
27303123	**Good identification**					**27303123**
					SPO	
Actinomyces naeslundii		97.90	0.53	RAF 92	1	
Clostridium chauvoei		2.10	0.01	FRU 95	99	
27303140	**Low discrimination**					**27303140**
Actinomyces naeslundii		53.81	0.37	RAF 92		
Propionibacterium propionicum		46.19	0.26	FRU 95		
27303141	**Good identification**					**27303141**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27303143	**Acceptable identification**					**27303143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.29	0.40	RAF 92		
Propionibacterium propionicum		8.71	0.12	FRU 95		
27303160	**Good identification**					**27303160**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27303161	**Good identification**					**27303161**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27303162	**Good identification**					**27303162**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27303163	**Good identification**					**27303163**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27304000	**Good identification**					**27304000**
					COC	
Actinomyces odontolyticus		99.21	0.54	MNS 9	1	
Gemella morbillorum		0.79	0.14	FRU 92	99	
27304002	**Good identification**					**27304002**
Actinomyces odontolyticus		99.04	0.51	MNS 9		
Actinomyces viscosus		0.96	0.09	FRU 92		
27304020	**Good identification**					**27304020**
Actinomyces odontolyticus		100.0	0.41	MNS 9		
27304022	**Good identification**					**27304022**
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27304100	**Good identification**					**27304100**
Actinomyces odontolyticus		100.0	0.46	MNS 9		
27304102	**Acceptable identification**					**27304102**
Actinomyces odontolyticus		94.85	0.42	MNS 9		
Actinomyces viscosus		5.15	0.14	FRU 92		
27304200	**Very good identification**					**27304200**
Actinomyces odontolyticus		100.0	0.59	MNS 9		
27304202	**Good identification**					**27304202**
Actinomyces odontolyticus		100.0	0.56	MNS 9		
27304220	**Good identification**					**27304220**
Actinomyces odontolyticus		100.0	0.46	MNS 9		
27304222	**Good identification**					**27304222**
Actinomyces odontolyticus		100.0	0.42	MNS 9		
27304300	**Good identification**					**27304300**
Actinomyces odontolyticus		100.0	0.51	MNS 9		
27304302	**Good identification**					**27304302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.47	MNS 9		
27304320	**Good identification**					**27304320**
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27305000	**Good identification**					**27305000**
					COC	
Actinomyces odontolyticus		99.36	0.51	MNS 9	1	
Gemella morbillorum		0.64	0.09	FRU 92	99	
27305002	**Good identification**					**27305002**
Actinomyces odontolyticus		99.12	0.48	MNS 9		
Actinomyces viscosus		0.88	0.06	FRU 92		
27305020	**Good identification**					**27305020**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27305100	**Good identification**					**27305100**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27305102	**Good identification**					**27305102**
Actinomyces odontolyticus		95.24	0.39	MNS 9		
Actinomyces viscosus		4.76	0.10	FRU 92		
27305200	**Very good identification**					**27305200**
Actinomyces odontolyticus		100.0	0.56	MNS 9		
27305202	**Good identification**					**27305202**
Actinomyces odontolyticus		100.0	0.53	MNS 9		
27305220	**Good identification**					**27305220**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27305222	**Good identification**					**27305222**
Actinomyces odontolyticus		100.0	0.39	MNS 9		
27305300	**Good identification**					**27305300**
Actinomyces odontolyticus		100.0	0.48	MNS 9		
27305302	**Good identification**					**27305302**
Actinomyces odontolyticus		100.0	0.44	MNS 9		
27310000	**Very good identification**					**27310000**
					COC	
Actinomyces odontolyticus		98.94	0.78	MNS 9	1	
Actinomyces viscosus		0.48	0.31		1	
Gemella morbillorum		0.37	0.32		99	
27310001	**Low discrimination**					**27310001**
Actinomyces odontolyticus		82.80	0.44	MNS 9		
Actinomyces viscosus		13.15	0.23	URE 1		
Actinomyces naeslundii		4.05	0.39			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27310002	**Low discrimination**					**27310002**
Actinomyces odontolyticus		56.23	0.74	MNS 9		
Actinomyces viscosus		43.72	0.65			
Actinomyces naeslundii		0.03	0.36			
27310003	**Very good identification**					**27310003**
Actinomyces viscosus		95.96	0.57	RAF 94		
Actinomyces odontolyticus		3.74	0.40			
Actinomyces naeslundii		0.30	0.39			
27310010	**Good identification**					**27310010**
Actinomyces odontolyticus		100.0	0.44	MNS 9		
27310012	**Low discrimination**					**27310012**
Actinomyces odontolyticus		56.26	0.40	MNS 9		
Actinomyces viscosus		43.74	0.31	MLZ 1		
27310020	**Very good identification**					**27310020**
					COC	
Actinomyces odontolyticus		98.18	0.64	MNS 9	1	
Clostridium perfringens		1.37	0.28	NAG 14	1	
Gemella morbillorum		0.25	0.15		99	
27310022	**Good identification**					**27310022**
Actinomyces odontolyticus		94.14	0.61	MNS 9		
Actinomyces viscosus		4.54	0.31	NAG 14		
Clostridium perfringens		1.01	0.22			
27310040	**Low discrimination**					**27310040**
					SPO	
Actinomyces odontolyticus		78.73	0.44	MNS 9	1	
Clostridium botulinum biovar B		20.81	0.24	SOR 1	99	
Actinomyces naeslundii		0.45	0.23		1	
27310042	**Low discrimination**					**27310042**
Actinomyces odontolyticus		55.96	0.40	MNS 9		
Actinomyces viscosus		43.51	0.31	SOR 1		
Actinomyces naeslundii		0.53	0.23			
27310100	**Very good identification**					**27310100**
					GRA	
Actinomyces odontolyticus		96.05	0.69	MNS 9	99	
Actinomyces viscosus		2.60	0.36		99	
Fusobacterium mortiferum		0.80	0.39		1	
27310101	**Low discrimination**					**27310101**
					SPO	
Actinomyces odontolyticus		30.75	0.35	MNS 9	1	
Clostridium chauvoei		28.20	0.29	URE 1	99	
Actinomyces viscosus		27.50	0.27		1	
27310102	**Low discrimination**					**27310102**
Actinomyces viscosus		81.25	0.70	RAF 94		
Actinomyces odontolyticus		18.56	0.66			
Actinomyces naeslundii		0.09	0.43			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27310103	**Very good identification**					**27310103**
Actinomyces viscosus		98.63	0.62	RAF 94		
Actinomyces odontolyticus		0.68	0.32			
Actinomyces naeslundii		0.49	0.46			
27310110	**Good identification**					**27310110**
Actinomyces odontolyticus		97.36	0.35	MNS 9		
Actinomyces viscosus		2.64	0.01	MLZ 1		
27310112	**Low discrimination**					**27310112**
Actinomyces viscosus		81.40	0.36	RAF 94		
Actinomyces odontolyticus		18.60	0.32	MLZ 1		
27310120	**Low discrimination**					**27310120**
Actinomyces odontolyticus		74.35	0.56	MNS 9		
Clostridium perfringens		24.17	0.42	NAG 14		
Actinomyces naeslundii		1.36	0.43			
27310121	**Low discrimination**					**27310121**
					SPO	
Actinomyces naeslundii		45.80	0.46	RAF 92	1	
Clostridium chauvoei		31.78	0.21	GAL 96	99	
Actinomyces odontolyticus		16.92	0.22		1	
27310122	**Low discrimination**					**27310122**
Actinomyces odontolyticus		64.40	0.52	MNS 9		
Actinomyces viscosus		17.49	0.36	NAG 14		
Clostridium perfringens		16.07	0.37			
27310140	**Low discrimination**					**27310140**
Actinomyces odontolyticus		82.04	0.35	MNS 9		
Propionibacterium propionicum		8.95	0.26	SOR 1		
Actinomyces naeslundii		4.25	0.30			
27310142	**Low discrimination**					**27310142**
Actinomyces viscosus		76.41	0.36	RAF 94		
Actinomyces odontolyticus		17.46	0.32	SOR 1		
Propionibacterium propionicum		4.66	0.29			
27310200	**Very good identification**					**27310200**
Actinomyces odontolyticus		100.0	0.83	MNS 9		
27310201	**Good identification**					**27310201**
Actinomyces odontolyticus		100.0	0.49	MNS 9		
27310202	**Very good identification**					**27310202**
Actinomyces odontolyticus		99.60	0.79	MNS 9		
Actinomyces viscosus		0.40	0.31			
27310203	**Low discrimination**					**27310203**
Actinomyces odontolyticus		88.22	0.45	MNS 9		
Actinomyces viscosus		11.78	0.23	URE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27310210	**Good identification**					**27310210**
Actinomyces odontolyticus		100.0	0.49	MNS 9		
27310212	**Good identification**					**27310212**
Actinomyces odontolyticus		100.0	0.45	MNS 9		
27310220	**Very good identification**					**27310220**
Actinomyces odontolyticus		100.0	0.69	MNS 9		
27310221	**Good identification**					**27310221**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27310222	**Very good identification**					**27310222**
Actinomyces odontolyticus		100.0	0.66	MNS 9		
27310230	**Good identification**					**27310230**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27310240	**Good identification**					**27310240**
Actinomyces odontolyticus		100.0	0.49	MNS 9		
27310242	**Good identification**					**27310242**
Actinomyces odontolyticus		100.0	0.45	MNS 9		
27310260	**Good identification**					**27310260**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27310300	**Very good identification**					**27310300**
Actinomyces odontolyticus		99.99	0.74	MNS 9		
Actinomyces viscosus		0.01	0.01			
27310301	**Good identification**					**27310301**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27310302	**Very good identification**					**27310302**
Actinomyces odontolyticus		97.77	0.71	MNS 9		
Actinomyces viscosus		2.23	0.36			
27310303	**Low discrimination**					**27310303**
Actinomyces odontolyticus		57.09	0.36	MNS 9		
Actinomyces viscosus		42.91	0.27	URE 1		
27310310	**Good identification**					**27310310**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27310312	**Good identification**					**27310312**
Actinomyces odontolyticus		97.77	0.36	MNS 9		
Actinomyces viscosus		2.23	0.01	MLZ 1		
27310320	**Very good identification**					**27310320**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.83	0.61	MNS 9		
Clostridium perfringens		0.17	0.08	NAG 14		
27310322	**Very good identification**					**27310322**
Actinomyces odontolyticus		99.73	0.57	MNS 9		
Actinomyces viscosus		0.14	0.01	NAG 14		
Clostridium perfringens		0.13	0.03			
27310340	**Good identification**					**27310340**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27310342	**Good identification**					**27310342**
Actinomyces odontolyticus		97.77	0.36	MNS 9		
Actinomyces viscosus		2.23	0.01	SOR 1		
27310400	**Good identification**					**27310400**
					COC	
Actinomyces odontolyticus		96.05	0.44	MNS 9	1	
Gemella morbillorum		3.95	0.15	ARA 1	99	
27310402	**Low discrimination**					**27310402**
Actinomyces odontolyticus		56.26	0.40	MNS 9		
Actinomyces viscosus		43.74	0.31	ARA 1		
27310500	**Good identification**					**27310500**
Actinomyces odontolyticus		97.36	0.35	MNS 9		
Actinomyces viscosus		2.64	0.01	ARA 1		
27310502	**Low discrimination**					**27310502**
Actinomyces viscosus		81.40	0.36	RAF 94		
Actinomyces odontolyticus		18.60	0.32	ARA 1		
27310600	**Good identification**					**27310600**
Actinomyces odontolyticus		100.0	0.49	MNS 9		
27310602	**Good identification**					**27310602**
Actinomyces odontolyticus		100.0	0.45	MNS 9		
27310620	**Good identification**					**27310620**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27310700	**Good identification**					**27310700**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27310702	**Good identification**					**27310702**
Actinomyces odontolyticus		97.77	0.36	MNS 9		
Actinomyces viscosus		2.23	0.01	ARA 1		
27311000	**Very good identification**					**27311000**
Actinomyces odontolyticus		95.60	0.75	MNS 9		
Actinomyces naeslundii		1.13	0.59			
Clostridium perfringens		1.00	0.36			
27311001	**Low discrimination**					**27311001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		52.37	0.62	RAF 92	1	
Actinomyces odontolyticus		29.76	0.41		1	
Clostridium chauvoei		12.93	0.29		99	
27311002	**Low discrimination**					**27311002**
Actinomyces odontolyticus		56.83	0.71	MNS 9		
Actinomyces viscosus		40.63	0.62			
Actinomyces naeslundii		1.10	0.59			
27311003	**Low discrimination**					**27311003**
Actinomyces viscosus		85.51	0.54	RAF 94		
Actinomyces naeslundii		10.41	0.62			
Actinomyces odontolyticus		3.63	0.37			
27311010	**Low discrimination**					**27311010**
Actinomyces odontolyticus		72.45	0.41	MNS 9		
Lactobacillus jensenii		22.32	0.28	MLZ 1		
Actinomyces naeslundii		4.47	0.37			
27311011	**Acceptable identification**					**27311011**
Actinomyces naeslundii		90.17	0.40	RAF 92		
Actinomyces odontolyticus		9.83	0.07	MLZ 5		
27311012	**Low discrimination**					**27311012**
Actinomyces odontolyticus		46.50	0.37	MNS 9		
Actinomyces viscosus		33.24	0.28	MLZ 1		
Lactobacillus jensenii		15.58	0.25			
27311020	**Low discrimination**					**27311020**
Clostridium perfringens		65.48	0.62	LAC 88		
Actinomyces odontolyticus		31.50	0.61			
Actinomyces naeslundii		2.29	0.59			
27311021	**Low discrimination**					**27311021**
Actinomyces naeslundii		73.15	0.62	RAF 92		
Clostridium perfringens		14.07	0.28			
Actinomyces odontolyticus		6.77	0.27			
27311022	**Low discrimination**					**27311022**
Clostridium perfringens		57.13	0.56	LAC 88		
Actinomyces odontolyticus		35.79	0.58			
Actinomyces naeslundii		4.25	0.59			
27311023	**Low discrimination**					**27311023**
Actinomyces naeslundii		80.34	0.62	RAF 92		
Clostridium perfringens		7.27	0.22			
Actinomyces viscosus		6.67	0.20			
27311031	**Good identification**					**27311031**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27311033	**Good identification**					**27311033**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	RAF 92		
27311040	**Low discrimination**					**27311040**
Actinomyces odontolyticus		70.74	0.41	MNS 9		
Actinomyces naeslundii		14.64	0.46	SOR 1		
Clostridium perfringens		13.96	0.24			
27311041	**Good identification**					**27311041**
Actinomyces naeslundii		96.85	0.49	RAF 92		
Actinomyces odontolyticus		3.15	0.07			
27311042	**Low discrimination**					**27311042**
Actinomyces odontolyticus		45.37	0.37	MNS 9		
Actinomyces viscosus		32.43	0.28	SOR 1		
Actinomyces naeslundii		15.32	0.46			
27311043	**Low discrimination**					**27311043**
Actinomyces naeslundii		68.01	0.49	RAF 92		
Actinomyces viscosus		31.99	0.20			
27311060	**Acceptable identification**					**27311060**
Clostridium perfringens		94.52	0.49	LAC 88		
Actinomyces naeslundii		3.07	0.46			
Actinomyces odontolyticus		2.41	0.27			
27311061	**Low discrimination**					**27311061**
Actinomyces naeslundii		82.81	0.49	RAF 92		
Clostridium perfringens		17.19	0.15			
27311062	**Acceptable identification**					**27311062**
Clostridium perfringens		90.73	0.44	LAC 88		
Actinomyces naeslundii		6.25	0.46			
Actinomyces odontolyticus		3.02	0.24			
27311063	**Acceptable identification**					**27311063**
Actinomyces naeslundii		91.10	0.49	RAF 92		
Clostridium perfringens		8.90	0.10			
27311100	**Low discrimination**					**27311100**
Actinomyces odontolyticus		69.87	0.66	MNS 9		
Clostridium perfringens		16.98	0.51			
Actinomyces naeslundii		7.46	0.67			
27311101	**Low discrimination**					**27311101**
					SPO	
Clostridium chauvoei		88.37	0.63	SAL 1	99	
Actinomyces naeslundii		10.29	0.70		1	
Actinomyces odontolyticus		0.65	0.32		1	
27311102	**Low discrimination**					**27311102**
Actinomyces viscosus		73.89	0.66	RAF 94		
Actinomyces odontolyticus		18.36	0.63			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		3.43	0.45			
27311103	**Low discrimination**					**27311103**
					SPO	
Actinomyces viscosus		66.70	0.58	RAF 94	1	
Clostridium chauvoei		19.69	0.50		99	
Actinomyces naeslundii		13.00	0.70		1	
27311113	**Low discrimination**					**27311113**
					SPO	
Actinomyces naeslundii		43.94	0.48	RAF 92	1	
Actinomyces viscosus		43.28	0.24	MLZ 5	1	
Clostridium chauvoei		12.78	0.16		99	
27311120	**Very good identification**					**27311120**
Clostridium perfringens		96.45	0.77			
Actinomyces odontolyticus		2.00	0.53			
Actinomyces naeslundii		1.31	0.67			
27311121	**Low discrimination**					**27311121**
					SPO	
Clostridium chauvoei		65.53	0.55	SAL 1	99	
Actinomyces naeslundii		22.89	0.70		1	
Clostridium perfringens		11.34	0.42		84	
27311122	**Good identification**					**27311122**
Clostridium perfringens		93.22	0.71			
Actinomyces naeslundii		2.69	0.67			
Actinomyces odontolyticus		2.52	0.49			
27311123	**Low discrimination**					**27311123**
					SPO	
Actinomyces naeslundii		55.67	0.70	RAF 92	1	
Clostridium chauvoei		28.12	0.42		99	
Clostridium perfringens		12.98	0.37		84	
27311130	**Acceptable identification**					**27311130**
Clostridium perfringens		91.62	0.42	MLZ 1		
Actinomyces naeslundii		6.49	0.45			
Actinomyces odontolyticus		1.90	0.19			
27311131	**Low discrimination**					**27311131**
					SPO	
Actinomyces naeslundii		60.81	0.48	RAF 92	1	
Clostridium chauvoei		33.41	0.21	MLZ 5	99	
Clostridium perfringens		5.78	0.08		84	
27311132	**Low discrimination**					**27311132**
Clostridium perfringens		84.25	0.37	MLZ 1		
Actinomyces naeslundii		12.68	0.45			
Actinomyces odontolyticus		2.27	0.15			
27311133	**Low discrimination**					**27311133**
					SPO	
Actinomyces naeslundii		87.59	0.48	RAF 92	1	
Clostridium chauvoei		8.49	0.08	MLZ 5	99	
Clostridium perfringens		3.92	0.03		84	
27311140	**Low discrimination**					**27311140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		60.21	0.38	NAG 97		
Actinomyces naeslundii		24.51	0.54			
Actinomyces odontolyticus		13.14	0.32			
27311141	**Low discrimination**					**27311141**
					SPO	
Actinomyces naeslundii		66.31	0.57	RAF 92	1	
Clostridium chauvoei		32.59	0.29		99	
Clostridium perfringens		1.10	0.04		84	
27311143	**Low discrimination**					**27311143**
					SPO	
Actinomyces naeslundii		71.93	0.57	RAF 92	1	
Actinomyces viscosus		21.13	0.24		1	
Clostridium chauvoei		6.24	0.16		99	
27311151	**Good identification**					**27311151**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27311153	**Good identification**					**27311153**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27311160	**Very good identification**					**27311160**
Clostridium perfringens		98.65	0.64			
Actinomyces naeslundii		1.24	0.54			
Actinomyces odontolyticus		0.11	0.19			
27311161	**Low discrimination**					**27311161**
					SPO	
Actinomyces naeslundii		58.87	0.57	RAF 92	1	
Clostridium perfringens		31.49	0.30		84	
Clostridium chauvoei		9.65	0.21		99	
27311162	**Very good identification**					**27311162**
Clostridium perfringens		97.03	0.59			
Actinomyces naeslundii		2.60	0.54			
Actinomyces odontolyticus		0.14	0.15			
27311163	**Low discrimination**					**27311163**
					SPO	
Actinomyces naeslundii		78.09	0.57	RAF 92	1	
Clostridium perfringens		19.65	0.24		84	
Clostridium chauvoei		2.26	0.08		99	
27311171	**Good identification**					**27311171**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27311173	**Good identification**					**27311173**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27311200	**Very good identification**					**27311200**
Actinomyces odontolyticus		99.98	0.80	MNS 9		
Actinomyces naeslundii		0.01	0.25			
Clostridium perfringens		0.01	0.02			
27311201	**Good identification**					**27311201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		99.09	0.46	MNS 9		
Actinomyces naeslundii		0.91	0.28	URE 1		
27311202	**Very good identification**					**27311202**
Actinomyces odontolyticus		99.62	0.76	MNS 9		
Actinomyces viscosus		0.37	0.28			
Actinomyces naeslundii		0.01	0.25			
27311203	**Low discrimination**					**27311203**
Actinomyces odontolyticus		87.90	0.42	MNS 9		
Actinomyces viscosus		10.79	0.20	URE 1		
Actinomyces naeslundii		1.31	0.28			
27311210	**Good identification**					**27311210**
Actinomyces odontolyticus		100.0	0.46	MNS 9		
27311212	**Good identification**					**27311212**
Actinomyces odontolyticus		100.0	0.42	MNS 9		
27311220	**Very good identification**					**27311220**
Actinomyces odontolyticus		98.89	0.66	MNS 9		
Clostridium perfringens		1.07	0.28	NAG 14		
Actinomyces naeslundii		0.04	0.25			
27311222	**Very good identification**					**27311222**
Actinomyces odontolyticus		99.06	0.63	MNS 9		
Clostridium perfringens		0.82	0.22	NAG 14		
Actinomyces naeslundii		0.06	0.25			
27311240	**Good identification**					**27311240**
Actinomyces odontolyticus		100.0	0.46	MNS 9		
27311242	**Good identification**					**27311242**
Actinomyces odontolyticus		100.0	0.42	MNS 9		
27311300	**Very good identification**					**27311300**
Actinomyces odontolyticus		99.82	0.71	MNS 9		
Clostridium perfringens		0.13	0.17			
Actinomyces naeslundii		0.06	0.33			
27311301	**Low discrimination**					**27311301**
					SPO	
Actinomyces odontolyticus		55.83	0.37	MNS 9	1	
Clostridium chauvoei		39.56	0.29	URE 1	99	
Actinomyces naeslundii		4.61	0.36		1	
27311302	**Very good identification**					**27311302**
Actinomyces odontolyticus		97.70	0.68	MNS 9		
Actinomyces viscosus		2.05	0.32			
Actinomyces naeslundii		0.09	0.33			
27311310	**Good identification**					**27311310**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27311320	**Low discrimination**					**27311320**
Actinomyces odontolyticus		79.65	0.58	MNS 9		
Clostridium perfringens		20.01	0.42	NAG 14		
Actinomyces naeslundii		0.27	0.33			
27311322	**Low discrimination**					**27311322**
					GRA	
Actinomyces odontolyticus		81.18	0.54	MNS 9	99	
Clostridium perfringens		15.66	0.37	NAG 14	99	
Prevotella oralis		2.29	0.19		1	
27311340	**Good identification**					**27311340**
Actinomyces odontolyticus		97.67	0.37	MNS 9		
Clostridium perfringens		2.33	0.04	SOR 1		
27311400	**Acceptable identification**					**27311400**
					COC	
Actinomyces odontolyticus		92.67	0.41	MNS 9	1	
Gemella morbillorum		3.08	0.11	ARA 1	99	
Lactobacillus fermentum		1.32	0.16		1	
27311402	**Low discrimination**					**27311402**
Actinomyces odontolyticus		56.90	0.37	MNS 9		
Actinomyces viscosus		40.68	0.28	ARA 1		
Lactobacillus fermentum		1.32	0.16			
27311520	**Good identification**					**27311520**
Clostridium perfringens		96.68	0.42	ARA 1		
Actinomyces odontolyticus		2.00	0.19			
Actinomyces naeslundii		1.31	0.33			
27311522	**Acceptable identification**					**27311522**
Clostridium perfringens		93.38	0.37	ARA 1		
Actinomyces naeslundii		2.70	0.33			
Actinomyces odontolyticus		2.52	0.15			
27311523	**Low discrimination**					**27311523**
					SPO	
Actinomyces naeslundii		57.53	0.36	RAF 92	1	
Clostridium chauvoei		29.06	0.08	ARA 1	99	
Clostridium perfringens		13.42	0.03		84	
27311600	**Good identification**					**27311600**
Actinomyces odontolyticus		100.0	0.46	MNS 9		
27311602	**Good identification**					**27311602**
Actinomyces odontolyticus		100.0	0.42	MNS 9		
27311700	**Good identification**					**27311700**
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27312000	**Acceptable identification**					**27312000**
Actinomyces odontolyticus		90.05	0.44	MNS 9		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		5.70	0.40	MAN 1		
Propionibacterium propionicum		3.10	0.26			
27312001	**Acceptable identification**					**27312001**
Actinomyces naeslundii		90.38	0.43	RAF 92		
Actinomyces odontolyticus		9.62	0.10	GAL 96		
27312002	**Low discrimination**					**27312002**
Actinomyces odontolyticus		50.89	0.40	MNS 9		
Actinomyces viscosus		39.57	0.31	MAN 1		
Actinomyces naeslundii		5.25	0.40			
27312021	**Good identification**					**27312021**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27312023	**Good identification**					**27312023**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27312100	**Low discrimination**					**27312100**
Propionibacterium propionicum		87.12	0.60	SAL 1		
Actinomyces odontolyticus		8.07	0.35			
Actinomyces naeslundii		4.60	0.48			
27312101	**Low discrimination**					**27312101**
Propionibacterium propionicum		58.39	0.44	SAL 1		
Actinomyces naeslundii		41.61	0.51	URE 10		
27312102	**Low discrimination**					**27312102**
Propionibacterium propionicum		80.74	0.63	SAL 1		
Actinomyces viscosus		13.37	0.36			
Actinomyces odontolyticus		3.05	0.32			
27312103	**Low discrimination**					**27312103**
Propionibacterium propionicum		50.71	0.47	SAL 1		
Actinomyces viscosus		25.20	0.27	URE 10		
Actinomyces naeslundii		24.09	0.51			
27312120	**Low discrimination**					**27312120**
Actinomyces naeslundii		63.71	0.48	RAF 92		
Actinomyces odontolyticus		18.19	0.22	GAL 96		
Propionibacterium propionicum		12.19	0.26			
27312121	**Good identification**					**27312121**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27312122	**Low discrimination**					**27312122**
Actinomyces naeslundii		64.38	0.48	RAF 92		
Propionibacterium propionicum		18.48	0.29	GAL 96		
Actinomyces odontolyticus		11.27	0.18			
27312123	**Good identification**					**27312123**
Actinomyces naeslundii		97.92	0.51	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		2.08	0.12	GAL 96		
27312140	**Very good identification**					**27312140**
Propionibacterium propionicum		99.08	0.60	SAL 1		
Actinomyces naeslundii		0.92	0.35			
27312141	**Low discrimination**					**27312141**
Propionibacterium propionicum		88.83	0.44	SAL 1		
Actinomyces naeslundii		11.17	0.38	URE 10		
27312142	**Very good identification**					**27312142**
Propionibacterium propionicum		99.22	0.63	SAL 1		
Actinomyces naeslundii		0.62	0.35			
Actinomyces viscosus		0.17	0.01			
27312143	**Acceptable identification**					**27312143**
Propionibacterium propionicum		92.27	0.47	SAL 1		
Actinomyces naeslundii		7.74	0.38	URE 10		
27312161	**Good identification**					**27312161**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27312163	**Low discrimination**					**27312163**
Actinomyces naeslundii		89.25	0.38	RAF 92		
Propionibacterium propionicum		10.75	0.12	GAL 96		
27312200	**Good identification**					**27312200**
Actinomyces odontolyticus		100.0	0.49	MNS 9		
27312202	**Good identification**					**27312202**
Actinomyces odontolyticus		100.0	0.45	MNS 9		
27312220	**Good identification**					**27312220**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27312300	**Acceptable identification**					**27312300**
Actinomyces odontolyticus		94.68	0.40	MNS 9		
Propionibacterium propionicum		5.32	0.26	MAN 1		
27312302	**Low discrimination**					**27312302**
Actinomyces odontolyticus		86.18	0.36	MNS 9		
Propionibacterium propionicum		11.85	0.29	MAN 1		
Actinomyces viscosus		1.96	0.01			
27313000	**Low discrimination**					**27313000**
Actinomyces naeslundii		67.21	0.64	RAF 92		
Actinomyces odontolyticus		29.52	0.41			
Propionibacterium propionicum		1.53	0.26			
27313001	**Very good identification**					**27313001**
Actinomyces naeslundii		99.71	0.67	RAF 92		
Actinomyces odontolyticus		0.30	0.07			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27313002	**Low discrimination**					**27313002**
Actinomyces naeslundii		66.27	0.64	RAF 92		
Actinomyces odontolyticus		17.84	0.37			
Actinomyces viscosus		12.75	0.28			
27313003	**Very good identification**					**27313003**
Actinomyces naeslundii		95.67	0.67	RAF 92		
Actinomyces viscosus		4.09	0.20			
Propionibacterium propionicum		0.24	0.12			
27313010	**Acceptable identification**					**27313010**
Actinomyces naeslundii		92.23	0.42	RAF 92		
Actinomyces odontolyticus		7.77	0.07	MLZ 5		
27313011	**Good identification**					**27313011**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27313012	**Good identification**					**27313012**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27313013	**Good identification**					**27313013**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27313020	**Low discrimination**					**27313020**
Actinomyces naeslundii		81.96	0.64	RAF 92		
Clostridium perfringens		12.18	0.28			
Actinomyces odontolyticus		5.86	0.27			
27313021	**Very good identification**					**27313021**
Actinomyces naeslundii		100.0	0.67	RAF 92		
27313022	**Low discrimination**					**27313022**
Actinomyces naeslundii		89.79	0.64	RAF 92		
Clostridium perfringens		6.28	0.22			
Actinomyces odontolyticus		3.93	0.24			
27313023	**Very good identification**					**27313023**
Actinomyces naeslundii		100.0	0.67	RAF 92		
27313030	**Good identification**					**27313030**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27313031	**Good identification**					**27313031**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27313032	**Good identification**					**27313032**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27313033	**Good identification**					**27313033**
Actinomyces naeslundii		100.0	0.45	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27313040	**Low discrimination**					**27313040**
Actinomyces naeslundii		86.68	0.51	RAF 92		
Propionibacterium propionicum		11.15	0.26			
Actinomyces odontolyticus		2.18	0.07			
27313041	**Good identification**					**27313041**
Actinomyces naeslundii		100.0	0.54	RAF 92		
27313042	**Low discrimination**					**27313042**
Actinomyces naeslundii		81.81	0.51	RAF 92		
Propionibacterium propionicum		15.78	0.29			
Actinomyces israelii		2.41	0.16			
27313043	**Good identification**					**27313043**
Actinomyces naeslundii		98.59	0.54	RAF 92		
Propionibacterium propionicum		1.41	0.12			
27313060	**Low discrimination**					**27313060**
Actinomyces naeslundii		86.18	0.51	RAF 92		
Clostridium perfringens		13.82	0.15			
27313061	**Good identification**					**27313061**
Actinomyces naeslundii		100.0	0.54	RAF 92		
27313062	**Acceptable identification**					**27313062**
Actinomyces naeslundii		92.98	0.51	RAF 92		
Clostridium perfringens		7.02	0.10			
27313063	**Good identification**					**27313063**
Actinomyces naeslundii		100.0	0.54	RAF 92		
27313100	**Low discrimination**					**27313100**
Actinomyces naeslundii		53.69	0.72	RAF 92		
Propionibacterium propionicum		42.38	0.60			
Actinomyces odontolyticus		2.62	0.32			
27313101	**Good identification**					**27313101**
					SPO	
Actinomyces naeslundii		90.65	0.75	RAF 92	1	
Propionibacterium propionicum		5.30	0.44		1	
Clostridium chauvoei		4.05	0.29		99	
27313102	**Low discrimination**					**27313102**
Propionibacterium propionicum		50.16	0.63	SAL 1		
Actinomyces naeslundii		42.37	0.72			
Actinomyces viscosus		5.09	0.32			
27313103	**Low discrimination**					**27313103**
Actinomyces naeslundii		89.10	0.75	RAF 92		
Propionibacterium propionicum		7.82	0.47			
Actinomyces viscosus		2.38	0.24			
27313110	**Low discrimination**					**27313110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.24	0.50	RAF 92		
Propionibacterium propionicum		12.76	0.26	MLZ 5		
Actinomyces israelii		3.00	0.16			
27313111	**Good identification**					**27313111**
Actinomyces naeslundii		100.0	0.53	RAF 92		
27313112	**Low discrimination**					**27313112**
Actinomyces naeslundii		78.09	0.50	RAF 92		
Propionibacterium propionicum		17.75	0.29	MLZ 5		
Actinomyces israelii		4.17	0.19			
27313113	**Good identification**					**27313113**
Actinomyces naeslundii		98.35	0.53	RAF 92		
Propionibacterium propionicum		1.66	0.12	MLZ 5		
27313120	**Low discrimination**					**27313120**
Actinomyces naeslundii		71.49	0.72	RAF 92		
Clostridium perfringens		27.37	0.42			
Propionibacterium propionicum		0.57	0.26			
27313121	**Very good identification**					**27313121**
					SPO	
Actinomyces naeslundii		98.28	0.75	RAF 92	1	
Clostridium chauvoei		1.46	0.21		99	
Clostridium perfringens		0.25	0.08		84	
27313122	**Low discrimination**					**27313122**
Actinomyces naeslundii		83.15	0.72	RAF 92		
Clostridium perfringens		14.98	0.37			
Propionibacterium propionicum		0.99	0.29			
27313123	**Very good identification**					**27313123**
					SPO	
Actinomyces naeslundii		99.53	0.75	RAF 92	1	
Clostridium chauvoei		0.26	0.08		99	
Clostridium perfringens		0.12	0.03		84	
27313130	**Acceptable identification**					**27313130**
Actinomyces naeslundii		93.15	0.50	RAF 92		
Clostridium perfringens		6.85	0.08	MLZ 5		
27313131	**Good identification**					**27313131**
Actinomyces naeslundii		100.0	0.53	RAF 92		
27313132	**Acceptable identification**					**27313132**
					SPO	
Actinomyces naeslundii		91.78	0.50	RAF 92	1	
Clostridium tertium		5.04	0.05	MLZ 5	99	
Clostridium perfringens		3.18	0.03		84	
27313133	**Good identification**					**27313133**
Actinomyces naeslundii		100.0	0.53	RAF 92		
27313140	**Low discrimination**					**27313140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		81.13	0.60	SAL 1		
Actinomyces naeslundii		18.14	0.59			
Actinomyces israelii		0.50	0.23			
27313141	**Low discrimination**					**27313141**
Actinomyces naeslundii		75.11	0.62	RAF 92		
Propionibacterium propionicum		24.89	0.44			
27313142	**Low discrimination**					**27313142**
Propionibacterium propionicum		86.57	0.63	SAL 1		
Actinomyces naeslundii		12.90	0.59			
Actinomyces israelii		0.53	0.26			
27313143	**Low discrimination**					**27313143**
Actinomyces naeslundii		66.80	0.62	RAF 92		
Propionibacterium propionicum		33.20	0.47			
27313150	**Low discrimination**					**27313150**
Actinomyces naeslundii		53.81	0.37	RAF 92		
Propionibacterium propionicum		46.19	0.26	MLZ 5		
27313151	**Good identification**					**27313151**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27313153	**Acceptable identification**					**27313153**
Actinomyces naeslundii		91.29	0.40	RAF 92		
Propionibacterium propionicum		8.71	0.12	MLZ 5		
27313160	**Low discrimination**					**27313160**
Actinomyces naeslundii		68.56	0.59	RAF 92		
Clostridium perfringens		28.34	0.30			
Propionibacterium propionicum		3.10	0.26			
27313161	**Very good identification**					**27313161**
Actinomyces naeslundii		100.0	0.62	RAF 92		
27313162	**Low discrimination**					**27313162**
Actinomyces naeslundii		79.22	0.59	RAF 92		
Clostridium perfringens		15.41	0.24			
Propionibacterium propionicum		5.37	0.29			
27313163	**Very good identification**					**27313163**
Actinomyces naeslundii		99.50	0.62	RAF 92		
Propionibacterium propionicum		0.50	0.12			
27313170	**Good identification**					**27313170**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27313171	**Good identification**					**27313171**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27313172	**Good identification**					**27313172**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	RAF 92		
27313173	**Good identification**					**27313173**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27313200	**Good identification**					**27313200**
Actinomyces odontolyticus		98.83	0.46	MNS 9		
Actinomyces naeslundii		1.17	0.30	MAN 1		
27313202	**Good identification**					**27313202**
Actinomyces odontolyticus		98.10	0.42	MNS 9		
Actinomyces naeslundii		1.90	0.30	MAN 1		
27313300	**Low discrimination**					**27313300**
Actinomyces odontolyticus		83.96	0.37	MNS 9		
Actinomyces naeslundii		8.97	0.38	MAN 1		
Propionibacterium propionicum		7.08	0.26			
27313301	**Good identification**					**27313301**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27313303	**Acceptable identification**					**27313303**
Actinomyces naeslundii		91.94	0.41	RAF 92		
Propionibacterium propionicum		8.06	0.12	RHA 1		
27313321	**Good identification**					**27313321**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27313322	**Low discrimination**					**27313322**
Actinomyces naeslundii		44.08	0.38	RAF 92		
Actinomyces odontolyticus		41.18	0.20	RHA 1		
Clostridium perfringens		7.95	0.03			
27313323	**Good identification**					**27313323**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27313501	**Good identification**					**27313501**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27313503	**Acceptable identification**					**27313503**
Actinomyces naeslundii		91.94	0.41	RAF 92		
Propionibacterium propionicum		8.06	0.12	ARA 1		
27313520	**Low discrimination**					**27313520**
Actinomyces naeslundii		72.31	0.38	RAF 92		
Clostridium perfringens		27.69	0.08	ARA 1		
27313521	**Good identification**					**27313521**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27313522	**Low discrimination**					**27313522**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		84.73	0.38	RAF 92			
Clostridium perfringens		15.27	0.03	ARA 1			
27313523	**Good identification**						**27313523**
Actinomyces naeslundii		100.0	0.41	RAF 92			
27314000	**Very good identification**						**27314000**
					COC	GRA	
Actinomyces odontolyticus		99.90	0.73	MNS 9	1	99	
Gemella morbillorum		0.08	0.15		99	99	
Fusobacterium mortiferum		0.02	0.14		1	1	
27314001	**Good identification**						**27314001**
Actinomyces odontolyticus		100.0	0.38	MNS 9			
27314002	**Very good identification**						**27314002**
Actinomyces odontolyticus		98.43	0.69	MNS 9			
Actinomyces viscosus		1.57	0.31				
27314010	**Good identification**						**27314010**
Actinomyces odontolyticus		100.0	0.38	MNS 9			
27314020	**Very good identification**						**27314020**
Actinomyces odontolyticus		100.0	0.59	MNS 9			
27314022	**Good identification**						**27314022**
Actinomyces odontolyticus		100.0	0.56	MNS 9			
27314040	**Good identification**						**27314040**
Actinomyces odontolyticus		100.0	0.38	MNS 9			
27314100	**Very good identification**						**27314100**
					GRA		
Actinomyces odontolyticus		99.82	0.64	MNS 9	99		
Fusobacterium mortiferum		0.13	0.20		1		
Actinomyces viscosus		0.06	0.01		99		
27314102	**Good identification**						**27314102**
Actinomyces odontolyticus		91.76	0.60	MNS 9			
Actinomyces viscosus		8.24	0.36				
27314120	**Good identification**						**27314120**
Actinomyces odontolyticus		99.34	0.51	MNS 9			
Clostridium perfringens		0.66	0.08	NAG 14			
27314122	**Good identification**						**27314122**
Actinomyces odontolyticus		98.94	0.47	MNS 9			
Actinomyces viscosus		0.55	0.01	NAG 14			
Clostridium perfringens		0.51	0.03				
27314200	**Very good identification**						**27314200**
Actinomyces odontolyticus		100.0	0.78	MNS 9			
27314201	**Good identification**						**27314201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27314202	**Very good identification**					**27314202**
Actinomyces odontolyticus		100.0	0.74	MNS 9		
27314203	**Good identification**					**27314203**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27314210	**Good identification**					**27314210**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27314212	**Good identification**					**27314212**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27314220	**Very good identification**					**27314220**
Actinomyces odontolyticus		100.0	0.64	MNS 9		
27314222	**Very good identification**					**27314222**
Actinomyces odontolyticus		100.0	0.60	MNS 9		
27314240	**Good identification**					**27314240**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27314242	**Good identification**					**27314242**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27314300	**Very good identification**					**27314300**
Actinomyces odontolyticus		100.0	0.69	MNS 9		
27314302	**Very good identification**					**27314302**
Actinomyces odontolyticus		99.95	0.65	MNS 9		
Actinomyces viscosus		0.05	0.01			
27314320	**Good identification**					**27314320**
Actinomyces odontolyticus		100.0	0.56	MNS 9		
27314322	**Good identification**					**27314322**
Actinomyces odontolyticus		100.0	0.52	MNS 9		
27314400	**Good identification**					**27314400**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27314600	**Good identification**					**27314600**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27314602	**Good identification**					**27314602**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27315000	**Very good identification**					**27315000**
					COC	
Actinomyces odontolyticus		99.73	0.70	MNS 9	1	
Actinomyces israelii		0.08	0.25		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Gemella morbillorum		0.07	0.11		99	
27315001	**Good identification**					**27315001**
Actinomyces odontolyticus		96.52	0.35	MNS 9		
Actinomyces naeslundii		3.48	0.28	URE 1		
27315002	**Very good identification**					**27315002**
Actinomyces odontolyticus		98.29	0.66	MNS 9		
Actinomyces viscosus		1.44	0.28			
Actinomyces israelii		0.20	0.28			
27315010	**Good identification**					**27315010**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27315020	**Very good identification**					**27315020**
Actinomyces odontolyticus		95.77	0.56	MNS 9		
Clostridium perfringens		4.08	0.28	NAG 14		
Actinomyces naeslundii		0.14	0.25			
27315022	**Good identification**					**27315022**
Actinomyces odontolyticus		96.60	0.53	MNS 9		
Clostridium perfringens		3.16	0.22	NAG 14		
Actinomyces naeslundii		0.24	0.25			
27315040	**Good identification**					**27315040**
Actinomyces odontolyticus		96.85	0.35	MNS 9		
Actinomyces israelii		3.15	0.18	SOR 1		
27315100	**Very good identification**					**27315100**
					COC	
Actinomyces odontolyticus		97.22	0.61	MNS 9	1	
Actinomyces israelii		1.04	0.35		1	
Atopobium parvulum		0.68	0.18		99	
27315102	**Low discrimination**					**27315102**
Actinomyces odontolyticus		88.77	0.57	MNS 9		
Actinomyces viscosus		7.33	0.32			
Actinomyces israelii		2.32	0.38			
27315120	**Low discrimination**					**27315120**
Actinomyces odontolyticus		49.62	0.48	MNS 9		
Clostridium perfringens		49.13	0.42	NAG 14		
Actinomyces naeslundii		0.67	0.33			
27315122	**Low discrimination**					**27315122**
Actinomyces odontolyticus		53.13	0.44	MNS 9		
Clostridium perfringens		40.39	0.37	NAG 14		
Bifidobacterium longum subsp. infantis		2.20	0.08			
27315123	**Low discrimination**					**27315123**
					SPO	
Actinomyces naeslundii		48.90	0.36	RAF 92	1	
Clostridium chauvoei		24.70	0.08	XYL 1	99	
Actinomyces odontolyticus		15.00	0.10		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27315200	**Very good identification**					**27315200**
Actinomyces odontolyticus		100.0	0.75	MNS 9		
27315201	**Good identification**					**27315201**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27315202	**Very good identification**					**27315202**
Actinomyces odontolyticus		100.0	0.71	MNS 9		
27315203	**Good identification**					**27315203**
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27315210	**Good identification**					**27315210**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27315212	**Good identification**					**27315212**
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27315220	**Very good identification**					**27315220**
Actinomyces odontolyticus		100.0	0.61	MNS 9		
27315222	**Very good identification**					**27315222**
Actinomyces odontolyticus		100.0	0.57	MNS 9		
27315240	**Good identification**					**27315240**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27315242	**Good identification**					**27315242**
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27315300	**Very good identification**					**27315300**
Actinomyces odontolyticus		98.29	0.66	MNS 9		
Eubacterium contortum		1.67	0.29			
Actinomyces israelii		0.04	0.16			
27315302	**Very good identification**					**27315302**
Actinomyces odontolyticus		99.41	0.62	MNS 9		
Eubacterium contortum		0.49	0.16			
Actinomyces israelii		0.10	0.19			
27315320	**Good identification**					**27315320**
					GRA	
Actinomyces odontolyticus		96.74	0.53	MNS 9	99	
Eubacterium contortum		1.78	0.16	NAG 14	99	
Bacteroides vulgatus		0.98	0.04		1	
27315322	**Good identification**					**27315322**
					GRA	
Actinomyces odontolyticus		97.88	0.49	MNS 9	99	
Prevotella oralis		0.69	0.04	NAG 14	1	
Parabacteroides distasonis		0.52	0.14		1	
27315400	**Acceptable identification**					**27315400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		92.14	0.35	MNS 9		
Actinomyces israelii		6.30	0.24	ARA 1		
Lactobacillus fermentum		1.56	0.12			
27315502	**Low discrimination**					**27315502**
Actinomyces israelii		39.91	0.37	CEL 93		
Bifidobacterium adolescentis		38.59	0.30			
Actinomyces odontolyticus		18.82	0.23			
27315600	**Good identification**					**27315600**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27315602	**Good identification**					**27315602**
Actinomyces odontolyticus		100.0	0.37	MNS 9		
27316000	**Good identification**					**27316000**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27316200	**Good identification**					**27316200**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27316202	**Good identification**					**27316202**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27317000	**Low discrimination**					**27317000**
Actinomyces odontolyticus		65.77	0.35	MNS 9		
Actinomyces israelii		31.16	0.38	MAN 1		
Actinomyces naeslundii		3.07	0.30			
27317002	**Low discrimination**					**27317002**
Actinomyces israelii		51.87	0.41	CEL 93		
Actinomyces odontolyticus		44.73	0.32			
Actinomyces naeslundii		3.41	0.30			
27317100	**Low discrimination**					**27317100**
Actinomyces israelii		77.39	0.48	CEL 93		
Actinomyces odontolyticus		12.90	0.27			
Actinomyces naeslundii		5.43	0.38			
27317101	**Acceptable identification**					**27317101**
Actinomyces naeslundii		91.24	0.41	RAF 92		
Actinomyces israelii		8.76	0.14	XYL 1		
27317102	**Low discrimination**					**27317102**
Actinomyces israelii		84.23	0.51	CEL 93		
Actinomyces odontolyticus		5.73	0.23			
Propionibacterium propionicum		4.66	0.29			
27317103	**Low discrimination**					**27317103**
Actinomyces naeslundii		81.19	0.41	RAF 92		
Actinomyces israelii		11.70	0.17	XYL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		7.12	0.12			
27317112	**Low discrimination**					**27317112**
					SPO	
Actinomyces israelii		84.69	0.37	CEL 93	1	
Bifidobacterium adolescentis		11.82	0.16	MLZ 13	1	
Clostridium tertium		3.49	-.01		99	
27317120	**Low discrimination**					**27317120**
Actinomyces naeslundii		37.32	0.38	RAF 92		
Actinomyces israelii		33.96	0.28	XYL 1		
Actinomyces odontolyticus		14.43	0.13			
27317121	**Good identification**					**27317121**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27317123	**Good identification**					**27317123**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27317140	**Low discrimination**					**27317140**
Actinomyces israelii		85.17	0.41	CEL 93		
Propionibacterium propionicum		12.12	0.26			
Actinomyces naeslundii		2.71	0.25			
27317142	**Low discrimination**					**27317142**
Actinomyces israelii		85.94	0.44	CEL 93		
Propionibacterium propionicum		12.23	0.29			
Actinomyces naeslundii		1.82	0.25			
27317200	**Good identification**					**27317200**
Actinomyces odontolyticus		98.20	0.40	MNS 9		
Actinomyces israelii		1.80	0.19	MAN 1		
27317202	**Good identification**					**27317202**
Actinomyces odontolyticus		95.70	0.37	MNS 9		
Actinomyces israelii		4.30	0.22	MAN 1		
27317400	**Good identification**					**27317400**
Actinomyces israelii		100.0	0.36	CEL 93		
27317402	**Good identification**					**27317402**
Actinomyces israelii		100.0	0.39	CEL 93		
27317500	**Good identification**					**27317500**
Actinomyces israelii		100.0	0.47	CEL 93		
27317502	**Low discrimination**					**27317502**
Actinomyces israelii		87.75	0.50	CEL 93		
Bifidobacterium adolescentis		12.25	0.29			
27317540	**Good identification**					**27317540**
Actinomyces israelii		100.0	0.40	CEL 93		
27317542	**Good identification**					**27317542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		95.75	0.43	CEL 93		
Bifidobacterium adolescentis		4.25	0.13			
27321000	**Low discrimination**					**27321000**
Actinomyces naeslundii		47.14	0.42	RAF 92		
Actinomyces odontolyticus		34.20	0.23	FRU 95		
Lactobacillus jensenii		11.02	0.10			
27321001	**Low discrimination**					**27321001**
					SPO	
Actinomyces naeslundii		89.68	0.45	RAF 92	1	
Clostridium chauvoei		10.32	0.06	FRU 95	99	
27321003	**Low discrimination**					**27321003**
Actinomyces naeslundii		75.84	0.45	RAF 92		
Actinomyces viscosus		24.16	0.13	FRU 95		
27321021	**Good identification**					**27321021**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27321023	**Good identification**					**27321023**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27321100	**Low discrimination**					**27321100**
					COC	
Actinomyces naeslundii		57.82	0.50	RAF 92	1	
Clostridium perfringens		24.16	0.21	FRU 95	1	
Atopobium parvulum		9.87	0.14		99	
27321101	**Low discrimination**					**27321101**
					SPO	
Clostridium chauvoei		80.01	0.40	SAL 1	99	
Actinomyces naeslundii		19.99	0.53	TRE 10	1	
27321102	**Low discrimination**					**27321102**
Actinomyces naeslundii		46.05	0.50	RAF 92		
Actinomyces viscosus		41.24	0.25	FRU 95		
Clostridium perfringens		9.06	0.16			
27321103	**Low discrimination**					**27321103**
					SPO	
Actinomyces naeslundii		52.48	0.53	RAF 92	1	
Clostridium chauvoei		37.07	0.27	FRU 95	99	
Actinomyces viscosus		10.45	0.17		1	
27321120	**Acceptable identification**					**27321120**
					COC	
Clostridium perfringens		92.92	0.47	FRU 99	1	
Actinomyces naeslundii		6.88	0.50		1	
Atopobium parvulum		0.21	0.01		99	
27321122	**Low discrimination**					**27321122**
Clostridium perfringens		86.41	0.41	FRU 99		
Actinomyces naeslundii		13.59	0.50			
27321123	**Low discrimination**					**27321123**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		78.23	0.53	RAF 92	1	
Clostridium chauvoei		18.42	0.19	FRU 95	99	
Clostridium perfringens		3.35	0.07		84	
27321140	**Low discrimination**					**27321140**
Actinomyces naeslundii		68.92	0.37	RAF 92		
Clostridium perfringens		31.08	0.09	FRU 95		
27321141	**Low discrimination**					**27321141**
					SPO	
Actinomyces naeslundii		81.36	0.40	RAF 92	1	
Clostridium chauvoei		18.64	0.06	FRU 95	99	
27321142	**Low discrimination**					**27321142**
Actinomyces naeslundii		82.49	0.37	RAF 92		
Clostridium perfringens		17.51	0.03	FRU 95		
27321143	**Good identification**					**27321143**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27321161	**Acceptable identification**					**27321161**
Actinomyces naeslundii		91.06	0.40	RAF 92		
Clostridium perfringens		8.94	0.01	FRU 95		
27321163	**Good identification**					**27321163**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27323000	**Good identification**					**27323000**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27323001	**Good identification**					**27323001**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27323002	**Good identification**					**27323002**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27323003	**Good identification**					**27323003**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27323020	**Good identification**					**27323020**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27323021	**Good identification**					**27323021**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27323022	**Good identification**					**27323022**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27323023	**Good identification**					**27323023**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27323041	**Good identification**					**27323041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	RAF 92		
27323043	**Good identification**					**27323043**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27323061	**Good identification**					**27323061**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27323063	**Good identification**					**27323063**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27323100	**Acceptable identification**					**27323100**
Actinomyces naeslundii		94.00	0.55	RAF 92		
Propionibacterium propionicum		6.00	0.24	FRU 95		
27323101	**Very good identification**					**27323101**
					SPO	
Actinomyces naeslundii		97.96	0.58	RAF 92	1	
Clostridium chauvoei		2.04	0.06	FRU 95	99	
27323102	**Acceptable identification**					**27323102**
Actinomyces naeslundii		91.26	0.55	RAF 92		
Propionibacterium propionicum		8.74	0.27	FRU 95		
27323103	**Very good identification**					**27323103**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27323111	**Good identification**					**27323111**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27323113	**Good identification**					**27323113**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27323120	**Acceptable identification**					**27323120**
Actinomyces naeslundii		93.43	0.55	RAF 92		
Clostridium perfringens		6.57	0.13	FRU 95		
27323121	**Very good identification**					**27323121**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27323122	**Good identification**					**27323122**
Actinomyces naeslundii		96.80	0.55	RAF 92		
Clostridium perfringens		3.20	0.07	FRU 95		
27323123	**Very good identification**					**27323123**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27323131	**Good identification**					**27323131**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27323133	**Good identification**					**27323133**
Actinomyces naeslundii		100.0	0.36	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27323140	**Low discrimination**					**27323140**
Actinomyces naeslundii		73.43	0.42	RAF 92		
Propionibacterium propionicum		26.57	0.24	FRU 95		
27323141	**Good identification**					**27323141**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27323142	**Low discrimination**					**27323142**
Actinomyces naeslundii		64.82	0.42	RAF 92		
Propionibacterium propionicum		35.18	0.27	FRU 95		
27323143	**Good identification**					**27323143**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27323160	**Acceptable identification**					**27323160**
Actinomyces naeslundii		92.95	0.42	RAF 92		
Clostridium perfringens		7.05	0.01	FRU 95		
27323161	**Good identification**					**27323161**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27323162	**Good identification**					**27323162**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27323163	**Good identification**					**27323163**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27330000	**Low discrimination**					**27330000**
					CAT	
Actinomyces odontolyticus		81.76	0.44	MNS 9	10	
Actinomyces naeslundii		5.17	0.40	TRE 1	25	
Propionibacterium granulosum		4.17	0.24		90	
27330001	**Low discrimination**					**27330001**
Actinomyces naeslundii		81.14	0.43	RAF 92		
Actinomyces viscosus		10.23	0.04	GAL 96		
Actinomyces odontolyticus		8.64	0.10			
27330002	**Low discrimination**					**27330002**
Actinomyces viscosus		83.39	0.46	RAF 94		
Actinomyces odontolyticus		14.39	0.40	TRE 7		
Actinomyces naeslundii		1.49	0.40			
27330003	**Acceptable identification**					**27330003**
Actinomyces viscosus		92.13	0.38	RAF 94		
Actinomyces naeslundii		7.39	0.43	TRE 7		
Actinomyces odontolyticus		0.48	0.06			
27330021	**Good identification**					**27330021**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27330023	**Low discrimination**					**27330023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		88.81	0.43	RAF 92		
Actinomyces viscosus		11.19	0.04	GAL 96		
27330100	**Low discrimination**					**27330100**
Actinomyces odontolyticus		39.50	0.35	MNS 9		
Actinomyces naeslundii		22.52	0.48	TRE 1		
Clostridium perfringens		11.89	0.21			
27330101	**Low discrimination**					**27330101**
					SPO	
Actinomyces naeslundii		83.48	0.51	RAF 92	1	
Clostridium chauvoei		9.95	0.12	GAL 96	99	
Actinomyces viscosus		6.57	0.08		1	
27330102	**Acceptable identification**					**27330102**
Actinomyces viscosus		92.99	0.51	RAF 94		
Actinomyces odontolyticus		2.85	0.32	TRE 7		
Actinomyces naeslundii		2.65	0.48			
27330103	**Low discrimination**					**27330103**
					SPO	
Actinomyces viscosus		88.42	0.42	RAF 94	1	
Actinomyces naeslundii		11.35	0.51	TRE 7	1	
Clostridium chauvoei		0.24	-.01		99	
27330120	**Acceptable identification**					**27330120**
Clostridium perfringens		92.80	0.47	GAL 99		
Actinomyces naeslundii		5.44	0.48			
Actinomyces odontolyticus		1.55	0.22			
27330121	**Low discrimination**					**27330121**
					SPO	
Actinomyces naeslundii		86.61	0.51	RAF 92	1	
Clostridium perfringens		9.95	0.13	GAL 96	84	
Clostridium chauvoei		3.44	0.04		99	
27330122	**Low discrimination**					**27330122**
Clostridium perfringens		84.01	0.41	GAL 99		
Actinomyces naeslundii		10.46	0.48			
Actinomyces viscosus		3.71	0.16			
27330123	**Low discrimination**					**27330123**
Actinomyces naeslundii		88.28	0.51	RAF 92		
Actinomyces viscosus		6.95	0.08	GAL 96		
Clostridium perfringens		4.78	0.07			
27330140	**Low discrimination**					**27330140**
Actinomyces naeslundii		36.91	0.35	RAF 92		
Propionibacterium propionicum		32.75	0.24	GAL 96		
Clostridium perfringens		21.03	0.09			
27330141	**Good identification**					**27330141**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27330143	**Low discrimination**					**27330143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		69.15	0.38	RAF 92		
Actinomyces viscosus		30.85	0.08	GAL 96		
27330161	**Low discrimination**					**27330161**
Actinomyces naeslundii		88.97	0.38	RAF 92		
Clostridium perfringens		11.03	0.01	GAL 96		
27330163	**Good identification**					**27330163**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27330200	**Good identification**					**27330200**
Actinomyces odontolyticus		100.0	0.49	MNS 9		
27330202	**Good identification**					**27330202**
Actinomyces odontolyticus		97.07	0.45	MNS 9		
Actinomyces viscosus		2.93	0.12	TRE 1		
27330220	**Good identification**					**27330220**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27330300	**Good identification**					**27330300**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27330302	**Low discrimination**					**27330302**
Actinomyces odontolyticus		85.49	0.36	MNS 9		
Actinomyces viscosus		14.51	0.16	TRE 1		
27331000	**Low discrimination**					**27331000**
Lactobacillus jensenii		33.69	0.44	NIT 1		
Actinomyces naeslundii		27.67	0.64	CEL 95		
Clostridium perfringens		23.38	0.41			
27331001	**Very good identification**					**27331001**
					SPO	
Actinomyces naeslundii		96.69	0.67	RAF 92	1	
Clostridium chauvoei		1.37	0.12		99	
Lactobacillus jensenii		0.79	0.10		1	
27331002	**Low discrimination**					**27331002**
Actinomyces viscosus		36.13	0.43	RAF 94		
Actinomyces naeslundii		25.20	0.64	TRE 7		
Lactobacillus jensenii		20.46	0.41			
27331003	**Low discrimination**					**27331003**
Actinomyces naeslundii		75.23	0.67	RAF 92		
Actinomyces viscosus		23.97	0.34			
Lactobacillus jensenii		0.41	0.07			
27331010	**Low discrimination**					**27331010**
Lactobacillus jensenii		85.26	0.36	NIT 1		
Actinomyces naeslundii		11.05	0.42	CEL 95		
Clostridium perfringens		1.79	0.06			
27331011	**Good identification**					**27331011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.06	0.45	RAF 92		
Lactobacillus jensenii		4.94	0.02	MLZ 5		
27331013	**Acceptable identification**					**27331013**
Actinomyces naeslundii		94.24	0.45	RAF 92		
Actinomyces viscosus		5.76	0.00	MLZ 5		
27331020	**Very good identification**					**27331020**
Clostridium perfringens		95.16	0.66	LAC 88		
Actinomyces naeslundii		3.48	0.64			
Lactobacillus jensenii		1.06	0.34			
27331021	**Low discrimination**					**27331021**
					SPO	
Actinomyces naeslundii		84.13	0.67	RAF 92	1	
Clostridium perfringens		15.48	0.32		84	
Clostridium chauvoei		0.40	0.04		99	
27331022	**Good identification**					**27331022**
Clostridium perfringens		90.87	0.61	LAC 88		
Actinomyces naeslundii		7.07	0.64			
Lactobacillus jensenii		1.43	0.31			
27331023	**Good identification**					**27331023**
Actinomyces naeslundii		91.76	0.67	RAF 92		
Clostridium perfringens		7.95	0.27			
Actinomyces viscosus		0.30	0.00			
27331031	**Good identification**					**27331031**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27331033	**Good identification**					**27331033**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27331040	**Low discrimination**					**27331040**
Actinomyces naeslundii		49.10	0.51	RAF 92		
Clostridium perfringens		44.78	0.28			
Lactobacillus jensenii		3.42	0.10			
27331041	**Good identification**					**27331041**
Actinomyces naeslundii		100.0	0.54	RAF 92		
27331042	**Low discrimination**					**27331042**
Actinomyces naeslundii		63.34	0.51	RAF 92		
Clostridium perfringens		27.19	0.23			
Actinomyces viscosus		5.20	0.08			
27331043	**Good identification**					**27331043**
Actinomyces naeslundii		98.21	0.54	RAF 92		
Actinomyces viscosus		1.79	0.00			
27331060	**Good identification**					**27331060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		96.72	0.54	LAC 88		
Actinomyces naeslundii		3.28	0.51			
27331061	**Low discrimination**					**27331061**
Actinomyces naeslundii		83.43	0.54	RAF 92		
Clostridium perfringens		16.57	0.20			
27331062	**Acceptable identification**					**27331062**
Clostridium perfringens		93.15	0.48	LAC 88		
Actinomyces naeslundii		6.71	0.51			
Lactobacillus acidophilus		0.14	-.02			
27331063	**Acceptable identification**					**27331063**
Actinomyces naeslundii		91.45	0.54	RAF 92		
Clostridium perfringens		8.55	0.14			
27331100	**Low discrimination**					**27331100**
					COC	
Clostridium perfringens		52.12	0.55	NAG 97	1	
Actinomyces naeslundii		23.94	0.72		1	
Atopobium parvulum		21.28	0.48		99	
27331101	**Low discrimination**					**27331101**
					SPO	
Actinomyces naeslundii		66.04	0.75	RAF 92	1	
Clostridium chauvoei		32.46	0.47		99	
Clostridium perfringens		0.97	0.21		84	
27331102	**Low discrimination**					**27331102**
Clostridium perfringens		31.45	0.50	NAG 97		
Actinomyces naeslundii		30.70	0.72			
Actinomyces viscosus		27.49	0.47			
27331103	**Low discrimination**					**27331103**
					SPO	
Actinomyces naeslundii		77.30	0.75	RAF 92	1	
Actinomyces viscosus		15.38	0.39		1	
Clostridium chauvoei		6.71	0.34		99	
27331110	**Low discrimination**					**27331110**
					COC	
Actinomyces naeslundii		61.89	0.50	RAF 92	1	
Clostridium perfringens		25.86	0.21	MLZ 5	1	
Atopobium parvulum		10.56	0.14		99	
27331111	**Acceptable identification**					**27331111**
					SPO	
Actinomyces naeslundii		91.38	0.53	RAF 92	1	
Clostridium chauvoei		8.62	0.12	MLZ 5	99	
27331112	**Low discrimination**					**27331112**
					GRA	
Actinomyces naeslundii		41.91	0.50	RAF 92	99	
Leptotrichia buccalis		39.90	0.15	MLZ 5	1	
Clostridium perfringens		8.24	0.16		99	
27331113	**Acceptable identification**					**27331113**
					SPO	
Actinomyces naeslundii		94.80	0.53	RAF 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		3.62	0.05	MLZ 5	1	
Clostridium chauvoei		1.58	-.01		99	
27331120	**Very good identification**					**27331120**
					COC	
Clostridium perfringens		98.33	0.81		1	
Actinomyces naeslundii		1.40	0.72		1	
Atopobium parvulum		0.22	0.35		99	
27331121	**Low discrimination**					**27331121**
					SPO	
Actinomyces naeslundii		61.01	0.75	RAF 92	1	
Clostridium perfringens		28.93	0.47		84	
Clostridium chauvoei		10.00	0.38		99	
27331122	**Very good identification**					**27331122**
Clostridium perfringens		96.58	0.76			
Actinomyces naeslundii		2.92	0.72			
Lactobacillus acidophilus		0.35	0.32			
27331123	**Low discrimination**					**27331123**
					SPO	
Actinomyces naeslundii		79.74	0.75	RAF 92	1	
Clostridium perfringens		17.79	0.41		84	
Clostridium chauvoei		2.31	0.25		99	
27331130	**Acceptable identification**					**27331130**
					COC	
Clostridium perfringens		92.92	0.47	MLZ 1	1	
Actinomyces naeslundii		6.88	0.50		1	
Atopobium parvulum		0.21	0.01		99	
27331131	**Low discrimination**					**27331131**
					SPO	
Actinomyces naeslundii		89.09	0.53	RAF 92	1	
Clostridium perfringens		8.11	0.13	MLZ 5	84	
Clostridium chauvoei		2.80	0.04		99	
27331132	**Low discrimination**					**27331132**
					GRA	
Clostridium perfringens		85.19	0.41	MLZ 1	99	
Actinomyces naeslundii		13.40	0.50		99	
Leptotrichia buccalis		1.42	-.01		1	
27331133	**Good identification**					**27331133**
Actinomyces naeslundii		95.89	0.53	RAF 92		
Clostridium perfringens		4.11	0.07	MLZ 5		
27331140	**Low discrimination**					**27331140**
					COC	
Clostridium perfringens		68.00	0.43	NAG 97	1	
Actinomyces naeslundii		28.93	0.59		1	
Atopobium parvulum		1.47	0.14		99	
27331141	**Very good identification**					**27331141**
					SPO	
Actinomyces naeslundii		95.79	0.62	RAF 92	1	
Clostridium chauvoei		2.70	0.12		99	
Clostridium perfringens		1.52	0.09		84	
27331142	**Low discrimination**					**27331142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		47.38	0.37	NAG 97		
Actinomyces naeslundii		42.84	0.59			
Lactobacillus acidophilus		3.64	0.16			
27331143	**Very good identification**					**27331143**
Actinomyces naeslundii		97.68	0.62	RAF 92		
Actinomyces viscosus		1.11	0.05			
Clostridium perfringens		0.73	0.03			
27331150	**Low discrimination**					**27331150**
Actinomyces naeslundii		68.92	0.37	RAF 92		
Clostridium perfringens		31.08	0.09	MLZ 5		
27331151	**Good identification**					**27331151**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27331152	**Low discrimination**					**27331152**
Actinomyces naeslundii		82.49	0.37	RAF 92		
Clostridium perfringens		17.51	0.03	MLZ 5		
27331153	**Good identification**					**27331153**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27331160	**Very good identification**					**27331160**
					COC	
Clostridium perfringens		98.69	0.69		1	
Actinomyces naeslundii		1.30	0.59		1	
Atopobium parvulum		0.01	0.01		99	
27331161	**Low discrimination**					**27331161**
					SPO	
Actinomyces naeslundii		65.74	0.62	RAF 92	1	
Clostridium perfringens		33.64	0.35		84	
Clostridium chauvoei		0.62	0.04		99	
27331162	**Very good identification**					**27331162**
Clostridium perfringens		97.06	0.63			
Actinomyces naeslundii		2.71	0.59			
Lactobacillus acidophilus		0.23	0.16			
27331163	**Low discrimination**					**27331163**
Actinomyces naeslundii		80.59	0.62	RAF 92		
Clostridium perfringens		19.41	0.29			
27331171	**Acceptable identification**					**27331171**
Actinomyces naeslundii		91.06	0.40	RAF 92		
Clostridium perfringens		8.94	0.01	MLZ 5		
27331173	**Good identification**					**27331173**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27331200	**Good identification**					**27331200**
Actinomyces odontolyticus		96.50	0.46	MNS 9		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		1.39	0.10	TRE 1		
Actinomyces naeslundii		1.14	0.30			
27331202	**Acceptable identification**					**27331202**
Actinomyces odontolyticus		93.42	0.42	MNS 9		
Actinomyces viscosus		2.59	0.08	TRE 1		
Actinomyces naeslundii		1.81	0.30			
27331300	**Low discrimination**					**27331300**
Actinomyces odontolyticus		69.72	0.37	MNS 9		
Clostridium perfringens		16.21	0.21	TRE 1		
Actinomyces naeslundii		7.45	0.38			
27331301	**Low discrimination**					**27331301**
					SPO	
Actinomyces naeslundii		67.04	0.41	RAF 92	1	
Clostridium chauvoei		32.96	0.12	RHA 1	99	
27331303	**Low discrimination**					**27331303**
					SPO	
Actinomyces naeslundii		77.78	0.41	RAF 92	1	
Actinomyces viscosus		15.48	0.05	RHA 1	1	
Clostridium chauvoei		6.75	-.01		99	
27331320	**Good identification**					**27331320**
Clostridium perfringens		96.33	0.47	RHA 1		
Actinomyces odontolyticus		2.09	0.24			
Actinomyces naeslundii		1.37	0.38			
27331321	**Low discrimination**					**27331321**
					SPO	
Actinomyces naeslundii		61.05	0.41	RAF 92	1	
Clostridium perfringens		28.95	0.13	RHA 1	84	
Clostridium chauvoei		10.00	0.04		99	
27331322	**Acceptable identification**					**27331322**
Clostridium perfringens		90.62	0.41	RHA 1		
Actinomyces naeslundii		2.74	0.38			
Actinomyces odontolyticus		2.56	0.20			
27331323	**Low discrimination**					**27331323**
Actinomyces naeslundii		81.76	0.41	RAF 92		
Clostridium perfringens		18.24	0.07	RHA 1		
27331501	**Low discrimination**					**27331501**
					SPO	
Actinomyces naeslundii		67.04	0.41	RAF 92	1	
Clostridium chauvoei		32.96	0.12	ARA 1	99	
27331503	**Low discrimination**					**27331503**
					SPO	
Actinomyces naeslundii		77.78	0.41	RAF 92	1	
Actinomyces viscosus		15.48	0.05	ARA 1	1	
Clostridium chauvoei		6.75	-.01		99	
27331520	**Good identification**					**27331520**
					COC	
Clostridium perfringens		98.38	0.47	ARA 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		1.40	0.38		1	
Atopobium parvulum		0.22	0.01		99	
27331521	**Low discrimination**					**27331521**
					SPO	
Actinomyces naeslundii		61.05	0.41	RAF 92	1	
Clostridium perfringens		28.95	0.13	ARA 1	84	
Clostridium chauvoei		10.00	0.04		99	
27331522	**Good identification**					**27331522**
Clostridium perfringens		97.07	0.41	ARA 1		
Actinomyces naeslundii		2.93	0.38			
27331523	**Low discrimination**					**27331523**
Actinomyces naeslundii		81.76	0.41	RAF 92		
Clostridium perfringens		18.24	0.07	ARA 1		
27332000	**Low discrimination**					**27332000**
					CAT	
Actinomyces naeslundii		61.54	0.45	RAF 92	25	
Propionibacterium granulosum		19.27	0.22	GAL 96	90	
Propionibacterium propionicum		14.13	0.24		1	
27332001	**Good identification**					**27332001**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27332002	**Low discrimination**					**27332002**
Actinomyces naeslundii		56.65	0.45	RAF 92		
Propionibacterium propionicum		19.51	0.27	GAL 96		
Actinomyces viscosus		16.55	0.12			
27332003	**Acceptable identification**					**27332003**
Actinomyces naeslundii		93.90	0.48	RAF 92		
Actinomyces viscosus		6.10	0.04	GAL 96		
27332020	**Acceptable identification**					**27332020**
Actinomyces naeslundii		92.74	0.45	RAF 92		
Propionibacterium granulosum		7.26	0.12	GAL 96		
27332021	**Good identification**					**27332021**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27332022	**Good identification**					**27332022**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27332023	**Good identification**					**27332023**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27332041	**Good identification**					**27332041**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27332043	**Good identification**					**27332043**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27332061	**Good identification**					**27332061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	RAF 92		
27332063	**Good identification**					**27332063**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27332100	**Low discrimination**					**27332100**
Propionibacterium propionicum		88.87	0.58	SAL 1		
Actinomyces naeslundii		11.13	0.53			
27332101	**Low discrimination**					**27332101**
Actinomyces naeslundii		62.83	0.56	RAF 92		
Propionibacterium propionicum		37.17	0.42	GAL 96		
27332102	**Good identification**					**27332102**
Propionibacterium propionicum		91.02	0.61	SAL 1		
Actinomyces naeslundii		7.60	0.53			
Actinomyces viscosus		1.39	0.16			
27332103	**Low discrimination**					**27332103**
Actinomyces naeslundii		51.87	0.56	RAF 92		
Propionibacterium propionicum		46.02	0.45	GAL 96		
Actinomyces viscosus		2.10	0.08			
27332120	**Low discrimination**					**27332120**
Actinomyces naeslundii		85.51	0.53	RAF 92		
Clostridium perfringens		7.59	0.13	GAL 96		
Propionibacterium propionicum		6.90	0.24			
27332121	**Very good identification**					**27332121**
Actinomyces naeslundii		100.0	0.56	RAF 92		
27332122	**Low discrimination**					**27332122**
Actinomyces naeslundii		86.00	0.53	RAF 92		
Propionibacterium propionicum		10.41	0.27	GAL 96		
Clostridium perfringens		3.59	0.07			
27332123	**Very good identification**					**27332123**
Actinomyces naeslundii		100.0	0.56	RAF 92		
27332140	**Very good identification**					**27332140**
Propionibacterium propionicum		97.84	0.58	SAL 1		
Actinomyces naeslundii		2.16	0.40			
27332141	**Low discrimination**					**27332141**
Propionibacterium propionicum		77.02	0.42	SAL 1		
Actinomyces naeslundii		22.98	0.43	URE 10		
27332142	**Very good identification**					**27332142**
Propionibacterium propionicum		98.55	0.61	SAL 1		
Actinomyces naeslundii		1.45	0.40			
27332143	**Low discrimination**					**27332143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		83.41	0.45	SAL 1		
Actinomyces naeslundii		16.59	0.43	URE 10		
27332160	**Low discrimination**					**27332160**
Actinomyces naeslundii		64.40	0.40	RAF 92		
Propionibacterium propionicum		29.43	0.24	GAL 96		
Clostridium perfringens		6.17	0.01			
27332161	**Good identification**					**27332161**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27332162	**Low discrimination**					**27332162**
Actinomyces naeslundii		59.33	0.40	RAF 92		
Propionibacterium propionicum		40.67	0.27	GAL 96		
27332163	**Good identification**					**27332163**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27333000	**Good identification**					**27333000**
					CAT	
Actinomyces naeslundii		94.98	0.69	RAF 92	25	
Actinomyces israelii		1.64	0.30		1	
Propionibacterium granulosum		1.24	0.22		90	
27333001	**Very good identification**					**27333001**
Actinomyces naeslundii		100.0	0.72	RAF 92		
27333002	**Good identification**					**27333002**
Actinomyces naeslundii		94.58	0.69	RAF 92		
Actinomyces israelii		2.45	0.33			
Propionibacterium propionicum		1.36	0.27			
27333003	**Very good identification**					**27333003**
Actinomyces naeslundii		99.83	0.72	RAF 92		
Actinomyces viscosus		0.17	0.00			
27333010	**Acceptable identification**					**27333010**
Actinomyces naeslundii		91.81	0.47	RAF 92		
Actinomyces israelii		4.50	0.15	MLZ 5		
Lactobacillus jensenii		3.69	0.02			
27333011	**Good identification**					**27333011**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27333012	**Acceptable identification**					**27333012**
Actinomyces naeslundii		93.15	0.47	RAF 92		
Actinomyces israelii		6.85	0.19	MLZ 5		
27333013	**Good identification**					**27333013**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27333020	**Low discrimination**					**27333020**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		87.30	0.69	RAF 92	25	
Clostridium perfringens		12.41	0.32		1	
Propionibacterium granulosum		0.29	0.12		90	
27333021	**Very good identification**					**27333021**
Actinomyces naeslundii		100.0	0.72	RAF 92		
27333022	**Good identification**					**27333022**
Actinomyces naeslundii		93.73	0.69	RAF 92		
Clostridium perfringens		6.27	0.27			
27333023	**Very good identification**					**27333023**
Actinomyces naeslundii		100.0	0.72	RAF 92		
27333030	**Good identification**					**27333030**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27333031	**Good identification**					**27333031**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27333032	**Good identification**					**27333032**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27333033	**Good identification**					**27333033**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27333040	**Good identification**					**27333040**
Actinomyces naeslundii		91.55	0.56	RAF 92		
Propionibacterium propionicum		4.96	0.24			
Actinomyces israelii		3.49	0.23			
27333041	**Very good identification**					**27333041**
Actinomyces naeslundii		100.0	0.59	RAF 92		
27333042	**Low discrimination**					**27333042**
Actinomyces naeslundii		87.84	0.56	RAF 92		
Propionibacterium propionicum		7.14	0.27			
Actinomyces israelii		5.02	0.26			
27333043	**Very good identification**					**27333043**
Actinomyces naeslundii		100.0	0.59	RAF 92		
27333051	**Good identification**					**27333051**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27333053	**Good identification**					**27333053**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27333060	**Low discrimination**					**27333060**
Actinomyces naeslundii		86.70	0.56	RAF 92		
Clostridium perfringens		13.31	0.20			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27333061	**Very good identification**					**27333061**
Actinomyces naeslundii		100.0	0.59	RAF 92		
27333062	**Good identification**					**27333062**
Actinomyces naeslundii		93.27	0.56	RAF 92		
Clostridium perfringens		6.74	0.14			
27333063	**Very good identification**					**27333063**
Actinomyces naeslundii		100.0	0.59	RAF 92		
27333071	**Good identification**					**27333071**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27333073	**Good identification**					**27333073**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27333100	**Low discrimination**					**27333100**
Actinomyces naeslundii		72.79	0.77	RAF 92		
Propionibacterium propionicum		24.22	0.58			
Actinomyces israelii		1.77	0.40			
27333101	**Very good identification**					**27333101**
					SPO	
Actinomyces naeslundii		97.35	0.80	RAF 92	1	
Propionibacterium propionicum		2.40	0.42		1	
Clostridium chauvoei		0.25	0.12		99	
27333102	**Low discrimination**					**27333102**
Actinomyces naeslundii		64.43	0.77	RAF 92		
Propionibacterium propionicum		32.16	0.61			
Actinomyces israelii		2.35	0.43			
27333103	**Very good identification**					**27333103**
Actinomyces naeslundii		96.30	0.80	RAF 92		
Propionibacterium propionicum		3.56	0.45			
Actinomyces viscosus		0.10	0.05			
27333110	**Low discrimination**					**27333110**
Actinomyces naeslundii		86.13	0.55	RAF 92		
Actinomyces israelii		5.94	0.26	MLZ 5		
Propionibacterium propionicum		5.50	0.24			
27333111	**Very good identification**					**27333111**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27333112	**Low discrimination**					**27333112**
Actinomyces naeslundii		77.88	0.55	RAF 92		
Actinomyces israelii		8.05	0.29	MLZ 5		
Propionibacterium propionicum		7.46	0.27			
27333113	**Very good identification**					**27333113**
Actinomyces naeslundii		100.0	0.58	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27333120	**Low discrimination**					**27333120**
Actinomyces naeslundii		72.62	0.77	RAF 92		
Clostridium perfringens		26.61	0.47			
Propionibacterium propionicum		0.24	0.24			
27333121	**Very good identification**					**27333121**
					SPO	
Actinomyces naeslundii		99.67	0.80	RAF 92	1	
Clostridium perfringens		0.25	0.13		84	
Clostridium chauvoei		0.09	0.04		99	
27333122	**Low discrimination**					**27333122**
					SPO	
Actinomyces naeslundii		83.33	0.77	RAF 92	1	
Clostridium perfringens		14.37	0.41		84	
Clostridium ramosum		0.79	0.17		92	
27333123	**Very good identification**					**27333123**
Actinomyces naeslundii		99.88	0.80	RAF 92		
Clostridium perfringens		0.12	0.07			
27333130	**Low discrimination**					**27333130**
					SPO	
Actinomyces naeslundii		86.56	0.55	RAF 92	1	
Clostridium tertium		7.35	0.13	MLZ 5	99	
Clostridium perfringens		6.09	0.13		84	
27333131	**Very good identification**					**27333131**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27333132	**Low discrimination**					**27333132**
					SPO	
Actinomyces naeslundii		77.65	0.55	RAF 92	1	
Clostridium tertium		19.78	0.21	MLZ 5	99	
Clostridium perfringens		2.57	0.07		84	
27333133	**Very good identification**					**27333133**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27333140	**Low discrimination**					**27333140**
Propionibacterium propionicum		63.89	0.58	SAL 1		
Actinomyces naeslundii		33.89	0.64			
Actinomyces israelii		1.81	0.33			
27333141	**Low discrimination**					**27333141**
Actinomyces naeslundii		87.75	0.67	RAF 92		
Propionibacterium propionicum		12.25	0.42			
27333142	**Low discrimination**					**27333142**
Propionibacterium propionicum		72.25	0.61	SAL 1		
Actinomyces naeslundii		25.55	0.64			
Actinomyces israelii		2.05	0.36			
27333143	**Low discrimination**					**27333143**
Actinomyces naeslundii		82.68	0.67	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		17.32	0.45			
27333150	**Low discrimination**					**27333150**
Actinomyces naeslundii		66.06	0.42	RAF 92		
Propionibacterium propionicum		23.90	0.24	MLZ 5		
Actinomyces israelii		10.03	0.19			
27333151	**Good identification**					**27333151**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27333152	**Low discrimination**					**27333152**
Actinomyces naeslundii		56.48	0.42	RAF 92		
Propionibacterium propionicum		30.66	0.27	MLZ 5		
Actinomyces israelii		12.87	0.22			
27333153	**Good identification**					**27333153**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27333160	**Low discrimination**					**27333160**
Actinomyces naeslundii		70.70	0.64	RAF 92		
Clostridium perfringens		27.96	0.35			
Propionibacterium propionicum		1.35	0.24			
27333161	**Very good identification**					**27333161**
Actinomyces naeslundii		99.74	0.67	RAF 92		
Clostridium perfringens		0.27	0.01			
27333162	**Low discrimination**					**27333162**
Actinomyces naeslundii		81.98	0.64	RAF 92		
Clostridium perfringens		15.26	0.29			
Propionibacterium propionicum		2.34	0.27			
27333163	**Very good identification**					**27333163**
Actinomyces naeslundii		100.0	0.67	RAF 92		
27333170	**Acceptable identification**					**27333170**
Actinomyces naeslundii		92.95	0.42	RAF 92		
Clostridium perfringens		7.05	0.01	MLZ 5		
27333171	**Good identification**					**27333171**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27333172	**Good identification**					**27333172**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27333173	**Good identification**					**27333173**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27333201	**Good identification**					**27333201**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333203	**Good identification**					**27333203**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333221	**Good identification**					**27333221**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333223	**Good identification**					**27333223**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333300	**Low discrimination**					**27333300**
Actinomyces naeslundii		66.07	0.43	RAF 92		
Propionibacterium propionicum		21.98	0.24	RHA 1		
Actinomyces israelii		11.95	0.21			
27333301	**Good identification**					**27333301**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27333302	**Low discrimination**					**27333302**
Actinomyces naeslundii		56.49	0.43	RAF 92		
Propionibacterium propionicum		28.19	0.27	RHA 1		
Actinomyces israelii		15.32	0.24			
27333303	**Good identification**					**27333303**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27333320	**Low discrimination**					**27333320**
					SPO	
Actinomyces naeslundii		59.62	0.43	RAF 92	1	
Clostridium perfringens		21.84	0.13	RHA 1	84	
Clostridium ramosum		18.54	0.09		92	
27333321	**Good identification**					**27333321**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27333323	**Good identification**					**27333323**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27333401	**Good identification**					**27333401**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333403	**Good identification**					**27333403**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333421	**Good identification**					**27333421**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333423	**Good identification**					**27333423**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27333500	**Low discrimination**					**27333500**
Actinomyces israelii		59.60	0.38	CEL 93		
Actinomyces naeslundii		30.32	0.43	XYL 92		
Propionibacterium propionicum		10.09	0.24			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27333501	**Good identification**					**27333501**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27333502	**Low discrimination**					**27333502**
Actinomyces israelii		64.48	0.41	CEL 93		
Actinomyces naeslundii		21.87	0.43	XYL 92		
Propionibacterium propionicum		10.91	0.27			
27333503	**Good identification**					**27333503**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27333520	**Low discrimination**					**27333520**
Actinomyces naeslundii		67.03	0.43	RAF 92		
Clostridium perfringens		24.56	0.13	ARA 1		
Actinomyces israelii		8.41	0.18			
27333521	**Good identification**					**27333521**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27333522	**Low discrimination**					**27333522**
Actinomyces naeslundii		73.50	0.43	RAF 92		
Actinomyces israelii		13.83	0.21	ARA 1		
Clostridium perfringens		12.67	0.07			
27333523	**Good identification**					**27333523**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27334000	**Good identification**					**27334000**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27334200	**Good identification**					**27334200**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27334202	**Good identification**					**27334202**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27335000	**Low discrimination**					**27335000**
					COC	
Actinomyces odontolyticus		59.63	0.35	MNS 9	1	
Actinomyces israelii		18.76	0.35	TRE 1	1	
Atopobium parvulum		13.09	0.18		99	
27335100	**Acceptable identification**					**27335100**
					COC	
Atopobium parvulum		91.48	0.53	NIT 1	99	
Actinomyces israelii		5.30	0.45		1	
Actinomyces odontolyticus		1.33	0.27		1	
27335102	**Low discrimination**					**27335102**
					COC	
Atopobium parvulum		60.45	0.40	NIT 1	99	
Actinomyces israelii		29.76	0.48		1	
Actinomyces odontolyticus		3.05	0.23		1	
27335103	**Low discrimination**					**27335103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces naeslundii		63.47	0.41	RAF 92	1		
Actinomyces viscosus		12.63	0.05	XYL 1	1		
Atopobium parvulum		12.33	0.06		99		
27335120	**Low discrimination**						**27335120**
					COC		
Clostridium perfringens		69.54	0.47	XYL 1	1		
Atopobium parvulum		28.49	0.40		99		
Actinomyces naeslundii		0.99	0.38		1		
27335121	**Low discrimination**						**27335121**
					COC		
Actinomyces naeslundii		54.58	0.41	RAF 92	1		
Clostridium perfringens		25.88	0.13	XYL 1	1		
Atopobium parvulum		10.60	0.06		99		
27335122	**Low discrimination**						**27335122**
					COC		
Clostridium perfringens		81.34	0.41	XYL 1	1		
Atopobium parvulum		12.50	0.27		99		
Actinomyces naeslundii		2.46	0.38		1		
27335123	**Low discrimination**						**27335123**
Actinomyces naeslundii		81.76	0.41	RAF 92			
Clostridium perfringens		18.24	0.07	XYL 1			
27335140	**Low discrimination**						**27335140**
					COC		
Actinomyces israelii		62.15	0.38	CEL 93	1		
Atopobium parvulum		27.87	0.18		99		
Clostridium perfringens		7.00	0.09		1		
27335142	**Low discrimination**						**27335142**
					COC		
Actinomyces israelii		89.28	0.41	CEL 93	1		
Atopobium parvulum		4.71	0.06		99		
Clostridium perfringens		3.16	0.03		1		
27335200	**Good identification**						**27335200**
Actinomyces odontolyticus		98.80	0.40	MNS 9			
Actinomyces israelii		1.21	0.16	TRE 1			
27335202	**Good identification**						**27335202**
Actinomyces odontolyticus		97.10	0.37	MNS 9			
Actinomyces israelii		2.90	0.19	TRE 1			
27335402	**Good identification**						**27335402**
Actinomyces israelii		100.0	0.36	CEL 93			
27335500	**Low discrimination**						**27335500**
					COC	SPO	
Actinomyces israelii		80.81	0.44	CEL 93	1	1	
Atopobium parvulum		17.23	0.18		99	1	
Clostridium butyricum		1.96	-.05		1	99	
27335502	**Low discrimination**						**27335502**
					COC		
Actinomyces israelii		83.58	0.47	CEL 93	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		14.32	0.27		1	
Atopobium parvulum		2.10	0.06		99	
27335540	**Good identification**					**27335540**
Actinomyces israelii		100.0	0.37	CEL 93		
27335542	**Acceptable identification**					**27335542**
Actinomyces israelii		94.84	0.40	CEL 93		
Bifidobacterium adolescentis		5.16	0.12			
27337000	**Good identification**					**27337000**
Actinomyces israelii		95.16	0.48	CEL 93		
Actinomyces naeslundii		4.84	0.35			
27337001	**Low discrimination**					**27337001**
Actinomyces naeslundii		88.30	0.38	RAF 92		
Actinomyces israelii		11.70	0.14	XYL 1		
27337002	**Good identification**					**27337002**
Actinomyces israelii		96.72	0.51	CEL 93		
Actinomyces naeslundii		3.28	0.35			
27337003	**Low discrimination**					**27337003**
Actinomyces naeslundii		83.43	0.38	RAF 92		
Actinomyces israelii		16.57	0.17	XYL 1		
27337012	**Good identification**					**27337012**
Actinomyces israelii		100.0	0.37	CEL 93		
27337021	**Good identification**					**27337021**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27337023	**Good identification**					**27337023**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27337040	**Good identification**					**27337040**
Actinomyces israelii		100.0	0.41	CEL 93		
27337042	**Good identification**					**27337042**
Actinomyces israelii		100.0	0.44	CEL 93		
27337100	**Good identification**					**27337100**
					COC	
Actinomyces israelii		92.32	0.58	CEL 93	1	
Actinomyces naeslundii		3.34	0.43		1	
Atopobium parvulum		2.84	0.18		99	
27337101	**Low discrimination**					**27337101**
Actinomyces naeslundii		84.31	0.46	RAF 92		
Actinomyces israelii		15.69	0.24	XYL 1		
27337102	**Very good identification**					**27337102**
Actinomyces israelii		95.14	0.61	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		2.30	0.43			
Propionibacterium propionicum		1.15	0.27			
27337103	**Low discrimination**					**27337103**
Actinomyces naeslundii		78.18	0.46	RAF 92		
Actinomyces israelii		21.82	0.27	XYL 1		
27337110	**Good identification**					**27337110**
					SPO	
Actinomyces israelii		95.30	0.44	CEL 93	1	
Clostridium tertium		4.70	0.07	MLZ 13	99	
27337112	**Low discrimination**					**27337112**
					SPO	
Actinomyces israelii		89.02	0.47	CEL 93	1	
Clostridium tertium		8.78	0.15	MLZ 13	99	
Bifidobacterium adolescentis		2.20	0.13		1	
27337120	**Low discrimination**					**27337120**
Actinomyces israelii		48.67	0.38	CEL 93		
Actinomyces naeslundii		27.59	0.43	NAG 6		
Clostridium perfringens		10.11	0.13			
27337121	**Good identification**					**27337121**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27337122	**Low discrimination**					**27337122**
					SPO	
Actinomyces israelii		54.56	0.41	CEL 93	1	
Clostridium tertium		21.26	0.19	NAG 6	99	
Actinomyces naeslundii		20.62	0.43		1	
27337123	**Good identification**					**27337123**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27337140	**Good identification**					**27337140**
Actinomyces israelii		95.48	0.51	CEL 93		
Propionibacterium propionicum		2.96	0.24			
Actinomyces naeslundii		1.57	0.30			
27337142	**Good identification**					**27337142**
Actinomyces israelii		95.98	0.54	CEL 93		
Propionibacterium propionicum		2.97	0.27			
Actinomyces naeslundii		1.05	0.30			
27337150	**Good identification**					**27337150**
Actinomyces israelii		100.0	0.37	CEL 93		
27337152	**Good identification**					**27337152**
Actinomyces israelii		100.0	0.40	CEL 93		
27337300	**Good identification**					**27337300**
Actinomyces israelii		100.0	0.39	CEL 93		
27337302	**Good identification**					**27337302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.42	CEL 93		
27337400	**Good identification**					**27337400**
Actinomyces israelii		100.0	0.46	CEL 93		
27337402	**Good identification**					**27337402**
Actinomyces israelii		100.0	0.49	CEL 93		
27337412	**Good identification**					**27337412**
Actinomyces israelii		100.0	0.35	CEL 93		
27337440	**Good identification**					**27337440**
Actinomyces israelii		100.0	0.39	CEL 93		
27337442	**Good identification**					**27337442**
Actinomyces israelii		100.0	0.42	CEL 93		
27337500	**Very good identification**					**27337500**
Actinomyces israelii		100.0	0.57	CEL 93		
27337502	**Very good identification**					**27337502**
Actinomyces israelii		97.59	0.60	CEL 93		
Bifidobacterium adolescentis		2.41	0.26			
27337510	**Good identification**					**27337510**
Actinomyces israelii		100.0	0.42	CEL 93		
27337512	**Low discrimination**					**27337512**
Actinomyces israelii		83.17	0.45	CEL 93		
Bifidobacterium adolescentis		16.83	0.27	MLZ 13		
27337520	**Good identification**					**27337520**
Actinomyces israelii		100.0	0.36	CEL 93		
27337522	**Good identification**					**27337522**
Actinomyces israelii		100.0	0.39	CEL 93		
27337540	**Good identification**					**27337540**
Actinomyces israelii		100.0	0.50	CEL 93		
27337542	**Good identification**					**27337542**
Actinomyces israelii		99.22	0.53	CEL 93		
Bifidobacterium adolescentis		0.78	0.10			
27337550	**Good identification**					**27337550**
Actinomyces israelii		100.0	0.35	CEL 93		
27337552	**Acceptable identification**					**27337552**
Actinomyces israelii		93.96	0.38	CEL 93		
Bifidobacterium adolescentis		6.04	0.12	MLZ 13		
27337700	**Good identification**					**27337700**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.37	CEL 93		
27337702	**Good identification**					**27337702**
Actinomyces israelii		100.0	0.40	CEL 93		
27340102	**Good identification**					**27340102**
Actinomyces viscosus		98.05	0.38	RAF 94		
Actinomyces odontolyticus		1.41	0.13	FRU 95		
Actinomyces naeslundii		0.54	0.24			
27341001	**Good identification**					**27341001**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27341021	**Good identification**					**27341021**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27341023	**Good identification**					**27341023**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27341101	**Low discrimination**					**27341101**
					SPO	
Actinomyces naeslundii		65.24	0.50	RAF 92	1	
Clostridium chauvoei		34.76	0.22	FRU 95	99	
27341120	**Low discrimination**					**27341120**
Actinomyces naeslundii		42.82	0.47	RAF 92		
Lactobacillus catenaformis		24.36	0.21	FRU 95		
Clostridium perfringens		19.01	0.19			
27341121	**Low discrimination**					**27341121**
					SPO	
Actinomyces naeslundii		84.92	0.50	RAF 92	1	
Clostridium chauvoei		15.08	0.14	FRU 95	99	
27341123	**Good identification**					**27341123**
					SPO	
Actinomyces naeslundii		96.96	0.50	RAF 92	1	
Clostridium chauvoei		3.04	0.01	FRU 95	99	
27341141	**Good identification**					**27341141**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27341143	**Good identification**					**27341143**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27341161	**Good identification**					**27341161**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27341163	**Good identification**					**27341163**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27343000	**Low discrimination**					**27343000**
					SPO	
Actinomyces naeslundii		80.05	0.44	RAF 92	1	
Clostridium innocuum		19.95	0.10	FRU 95	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27343001	**Good identification**					**27343001**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27343002	**Low discrimination**					**27343002**
					SPO	
Actinomyces naeslundii		80.05	0.44	RAF 92	1	
Clostridium innocuum		19.95	0.10	FRU 95	99	
27343003	**Good identification**					**27343003**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27343020	**Good identification**					**27343020**
Actinomyces naeslundii		100.0	0.44	RAF 92		
27343021	**Good identification**					**27343021**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27343022	**Good identification**					**27343022**
Actinomyces naeslundii		100.0	0.44	RAF 92		
27343023	**Good identification**					**27343023**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27343100	**Good identification**					**27343100**
Actinomyces naeslundii		97.65	0.52	RAF 92		
Actinomyces israelii		2.35	0.15	FRU 95		
27343101	**Good identification**					**27343101**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27343102	**Acceptable identification**					**27343102**
Actinomyces naeslundii		91.95	0.52	RAF 92		
Actinomyces israelii		3.32	0.18	FRU 95		
Lactobacillus acidophilus		2.51	0.01			
27343103	**Good identification**					**27343103**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27343120	**Good identification**					**27343120**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27343121	**Good identification**					**27343121**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27343122	**Acceptable identification**					**27343122**
					SPO	
Actinomyces naeslundii		92.47	0.52	RAF 92	1	
Clostridium barati		2.84	-.01	FRU 95	99	
Lactobacillus acidophilus		2.53	0.01		1	
27343123	**Good identification**					**27343123**
Actinomyces naeslundii		100.0	0.55	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27343140	**Good identification**					**27343140**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27343141	**Good identification**					**27343141**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27343142	**Good identification**					**27343142**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27343143	**Good identification**					**27343143**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27343160	**Good identification**					**27343160**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27343161	**Good identification**					**27343161**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27343162	**Good identification**					**27343162**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27343163	**Good identification**					**27343163**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27347102	**Good identification**					**27347102**
Actinomyces israelii		100.0	0.36	FRU 99		
27350000	**Low discrimination**					**27350000**
Actinomyces odontolyticus		63.23	0.44	MNS 9		
Lactobacillus catenaformis		14.90	0.29	CEL 1		
Actinomyces viscosus		8.01	0.21			
27350001	**Low discrimination**					**27350001**
Actinomyces naeslundii		55.32	0.41	RAF 92		
Actinomyces viscosus		36.05	0.13	GAL 96		
Actinomyces odontolyticus		8.63	0.10			
27350002	**Acceptable identification**					**27350002**
Actinomyces viscosus		92.30	0.56	RAF 94		
Actinomyces odontolyticus		4.51	0.40			
Lactobacillus catenaformis		1.73	0.29			
27350003	**Good identification**					**27350003**
Actinomyces viscosus		98.33	0.47	RAF 94		
Actinomyces naeslundii		1.52	0.41			
Actinomyces odontolyticus		0.15	0.06			
27350021	**Good identification**					**27350021**
Actinomyces naeslundii		100.0	0.41	RAF 92		
27350023	**Low discrimination**					**27350023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		60.54	0.41	RAF 92		
Actinomyces viscosus		39.46	0.13	GAL 96		
27350101	**Low discrimination**					**27350101**
Actinomyces naeslundii		71.07	0.48	RAF 92		
Actinomyces viscosus		28.93	0.18	GAL 96		
27350102	**Very good identification**					**27350102**
Actinomyces viscosus		95.62	0.60	RAF 94		
Lactobacillus catenaformis		1.52	0.32			
Actinomyces odontolyticus		0.83	0.32			
27350103	**Good identification**					**27350103**
Actinomyces viscosus		97.58	0.52	RAF 94		
Actinomyces naeslundii		2.42	0.48			
27350121	**Good identification**					**27350121**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27350123	**Low discrimination**					**27350123**
					SPO	
Actinomyces naeslundii		69.56	0.48	RAF 92	1	
Actinomyces viscosus		28.32	0.18	GAL 96	1	
Clostridium barati		2.12	-.04		99	
27350141	**Good identification**					**27350141**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27350161	**Good identification**					**27350161**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27350163	**Good identification**					**27350163**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27350200	**Good identification**					**27350200**
Actinomyces odontolyticus		100.0	0.49	MNS 9		
27350202	**Acceptable identification**					**27350202**
Actinomyces odontolyticus		90.38	0.45	MNS 9		
Actinomyces viscosus		9.62	0.21	CEL 1		
27350220	**Good identification**					**27350220**
Actinomyces odontolyticus		100.0	0.35	MNS 9		
27350300	**Good identification**					**27350300**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27350302	**Low discrimination**					**27350302**
					SPO	
Actinomyces odontolyticus		59.75	0.36	MNS 9	1	
Actinomyces viscosus		35.82	0.26	CEL 1	1	
Clostridium barati		2.56	0.04		99	
27351000	**Low discrimination**					**27351000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		62.43	0.63	NIT 1		
Lactobacillus jensenii		31.34	0.58			
Actinomyces naeslundii		2.78	0.61			
27351001	**Low discrimination**					**27351001**
Actinomyces naeslundii		80.09	0.64	RAF 92		
Lactobacillus catenaformis		12.13	0.29			
Lactobacillus jensenii		6.09	0.24			
27351002	**Low discrimination**					**27351002**
Lactobacillus catenaformis		54.43	0.63	NIT 1		
Lactobacillus jensenii		18.22	0.55			
Actinomyces viscosus		17.94	0.52			
27351003	**Low discrimination**					**27351003**
Actinomyces viscosus		57.49	0.44	RAF 94		
Actinomyces naeslundii		34.89	0.64			
Lactobacillus catenaformis		5.29	0.29			
27351010	**Acceptable identification**					**27351010**
Lactobacillus jensenii		92.93	0.50	NIT 1		
Lactobacillus catenaformis		5.61	0.29			
Actinomyces naeslundii		1.30	0.39			
27351011	**Low discrimination**					**27351011**
Actinomyces naeslundii		67.49	0.42	RAF 92		
Lactobacillus jensenii		32.51	0.15	MLZ 5		
27351012	**Low discrimination**					**27351012**
Lactobacillus jensenii		86.67	0.47	NIT 1		
Lactobacillus catenaformis		7.85	0.29			
Actinomyces viscosus		2.59	0.18			
27351013	**Low discrimination**					**27351013**
Actinomyces naeslundii		61.07	0.42	RAF 92		
Lactobacillus jensenii		19.61	0.12	MLZ 5		
Actinomyces viscosus		19.31	0.10			
27351020	**Low discrimination**					**27351020**
					SPO	
Lactobacillus catenaformis		48.45	0.53	NIT 1	1	
Lactobacillus jensenii		24.32	0.47		1	
Clostridium barati		9.74	0.35		99	
27351021	**Good identification**					**27351021**
Actinomyces naeslundii		93.41	0.64	RAF 92		
Lactobacillus catenaformis		3.54	0.18			
Lactobacillus jensenii		1.78	0.13			
27351022	**Low discrimination**					**27351022**
					SPO	
Lactobacillus catenaformis		40.68	0.53	NIT 1	1	
Clostridium barati		24.54	0.44		99	

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii	13.62	0.44		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27351023	**Good identification**					**27351023**
					SPO	
Actinomyces naeslundii		91.01	0.64	RAF 92	1	
Lactobacillus catenaformis		3.45	0.18		1	
Clostridium barati		2.08	0.09		99	
27351030	**Low discrimination**					**27351030**
Lactobacillus jensenii		87.87	0.39	NIT 1		
Lactobacillus catenaformis		5.30	0.18			
Actinomyces naeslundii		4.91	0.39			
27351031	**Low discrimination**					**27351031**
Actinomyces naeslundii		89.25	0.42	RAF 92		
Lactobacillus jensenii		10.75	0.05	MLZ 5		
27351032	**Low discrimination**					**27351032**
Lactobacillus jensenii		80.00	0.36	NIT 1		
Lactobacillus catenaformis		7.24	0.18			
Actinomyces naeslundii		6.71	0.39			
27351033	**Acceptable identification**					**27351033**
Actinomyces naeslundii		92.57	0.42	RAF 92		
Lactobacillus jensenii		7.43	0.02	MLZ 5		
27351041	**Good identification**					**27351041**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27351043	**Acceptable identification**					**27351043**
Actinomyces naeslundii		91.38	0.51	RAF 92		
Actinomyces viscosus		8.62	0.10			
27351060	**Low discrimination**					**27351060**
					SPO	
Actinomyces naeslundii		33.22	0.48	RAF 92	1	
Clostridium perfringens		32.19	0.26		84	
Clostridium barati		18.52	0.17		99	
27351061	**Good identification**					**27351061**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27351063	**Good identification**					**27351063**
Actinomyces naeslundii		100.0	0.51	RAF 92		
27351100	**Low discrimination**					**27351100**
Lactobacillus catenaformis		81.60	0.66	NIT 1		
Actinomyces naeslundii		6.88	0.69			
Collinsella aerofaciens		6.00	0.32			
27351101	**Low discrimination**					**27351101**
					SPO	
Actinomyces naeslundii		85.89	0.72	RAF 92	1	
Lactobacillus catenaformis		6.86	0.32		1	
Clostridium chauvoei		5.62	0.29		99	
27351102	**Low discrimination**					**27351102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		48.93	0.66	NIT 1		
Lactobacillus acidophilus		21.69	0.57			
Actinomyces viscosus		19.11	0.56			
27351103	**Low discrimination**					**27351103**
Actinomyces viscosus		47.56	0.48	RAF 94		
Actinomyces naeslundii		46.21	0.72			
Lactobacillus catenaformis		3.69	0.32			
27351111	**Good identification**					**27351111**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27351113	**Low discrimination**					**27351113**
Actinomyces naeslundii		83.51	0.50	RAF 92		
Actinomyces viscosus		16.50	0.14	MLZ 5		
27351120	**Low discrimination**					**27351120**
					SPO	
Lactobacillus catenaformis		36.09	0.56	NIT 1	1	
Clostridium perfringens		28.17	0.53		84	
Clostridium barati		14.52	0.44		99	
27351121	**Good identification**					**27351121**
					SPO	
Actinomyces naeslundii		93.58	0.72	RAF 92	1	
Clostridium chauvoei		2.04	0.21		99	
Lactobacillus catenaformis		1.87	0.21		1	
27351122	**Low discrimination**					**27351122**
					SPO	
Lactobacillus acidophilus		36.17	0.57	NIT 1	1	
Clostridium barati		24.62	0.52		99	
Lactobacillus catenaformis		20.40	0.56		1	
27351123	**Good identification**					**27351123**
					SPO	
Actinomyces naeslundii		90.53	0.72	RAF 92	1	
Lactobacillus acidophilus		3.20	0.23		1	
Clostridium barati		2.18	0.18		99	
27351130	**Low discrimination**					**27351130**
Actinomyces naeslundii		41.69	0.47	RAF 92		
Lactobacillus catenaformis		23.72	0.21	MLZ 5		
Clostridium perfringens		18.51	0.19			
27351131	**Good identification**					**27351131**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27351133	**Good identification**					**27351133**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27351140	**Low discrimination**					**27351140**
					SPO	
Actinomyces naeslundii		46.87	0.56	RAF 92	1	
Lactobacillus catenaformis		31.81	0.32		1	
Clostridium barati		9.18	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27351141	**Very good identification**					**27351141**
Actinomyces naeslundii		100.0	0.59	RAF 92		
27351142	**Low discrimination**					**27351142**
Lactobacillus acidophilus		57.60	0.41	NIT 1		
Actinomyces naeslundii		15.65	0.56	SOR 11		
Lactobacillus catenaformis		10.62	0.32			
27351143	**Good identification**					**27351143**
Actinomyces naeslundii		92.28	0.59	RAF 92		
Actinomyces viscosus		5.44	0.14			
Lactobacillus acidophilus		2.29	0.07			
27351151	**Good identification**					**27351151**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27351153	**Good identification**					**27351153**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27351160	**Low discrimination**					**27351160**
					SPO	
Clostridium perfringens		55.49	0.41	CEL 4	84	
Actinomyces naeslundii		22.23	0.56		1	
Clostridium barati		13.06	0.25		99	
27351161	**Very good identification**					**27351161**
Actinomyces naeslundii		98.35	0.59	RAF 92		
Clostridium perfringens		1.65	0.06			
27351162	**Low discrimination**					**27351162**
					SPO	
Lactobacillus acidophilus		46.54	0.41	NIT 1	1	
Clostridium barati		22.28	0.34	SOR 11	99	
Clostridium perfringens		14.86	0.35		84	
27351163	**Very good identification**					**27351163**
					SPO	
Actinomyces naeslundii		95.73	0.59	RAF 92	1	
Lactobacillus acidophilus		2.37	0.07		1	
Clostridium barati		1.14	-.01		99	
27351171	**Good identification**					**27351171**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27351173	**Good identification**					**27351173**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27351200	**Low discrimination**					**27351200**
Actinomyces odontolyticus		76.47	0.46	MNS 9		
Lactobacillus catenaformis		13.92	0.29	CEL 1		
Lactobacillus jensenii		6.99	0.24			
27351202	**Low discrimination**					**27351202**
					SPO	
Actinomyces odontolyticus		56.69	0.42	MNS 9	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		16.84	0.29	CEL 1	1		
Clostridium barati		7.29	0.17		99		
27351300	**Low discrimination**						**27351300**
					SPO		
Actinomyces odontolyticus		43.43	0.37	MNS 9	1		
Lactobacillus catenaformis		37.55	0.32	CEL 1	1		
Clostridium barati		10.83	0.17		99		
27351301	**Good identification**						**27351301**
Actinomyces naeslundii		100.0	0.38	RAF 92			
27351302	**Low discrimination**						**27351302**
					SPO		
Lactobacillus acidophilus		51.18	0.38	NIT 1	1		
Lactobacillus catenaformis		15.50	0.32	RHA 7	1		
Clostridium barati		13.41	0.25		99		
27351321	**Good identification**						**27351321**
Actinomyces naeslundii		100.0	0.38	RAF 92			
27351322	**Low discrimination**						**27351322**
					GRA	SPO	
Lactobacillus acidophilus		41.63	0.38	NIT 1	99	1	
Clostridium barati		32.73	0.34	RHA 7	99	99	
Prevotella oralis		14.40	0.29		1	1	
27351323	**Low discrimination**						**27351323**
					SPO		
Actinomyces naeslundii		67.98	0.38	RAF 92	1		
Lactobacillus acidophilus		17.93	0.04	RHA 1	1		
Clostridium barati		14.09	-.01		99		
27351501	**Good identification**						**27351501**
Actinomyces naeslundii		100.0	0.38	RAF 92			
27351521	**Good identification**						**27351521**
Actinomyces naeslundii		100.0	0.38	RAF 92			
27351523	**Good identification**						**27351523**
Actinomyces naeslundii		100.0	0.38	RAF 92			
27352000	**Low discrimination**						**27352000**
					SPO		
Actinomyces naeslundii		69.67	0.43	RAF 92	1		
Clostridium innocuum		21.93	0.10	GAL 96	99		
Actinomyces odontolyticus		8.40	0.10		1		
27352001	**Good identification**						**27352001**
Actinomyces naeslundii		100.0	0.46	RAF 92			
27352003	**Low discrimination**						**27352003**
Actinomyces naeslundii		74.86	0.46	RAF 92			
Actinomyces viscosus		25.14	0.13	GAL 96			
27352020	**Good identification**						**27352020**
Actinomyces naeslundii		100.0	0.43	RAF 92			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27352021	**Good identification**					**27352021**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27352022	**Low discrimination**					**27352022**
					SPO	
Actinomyces naeslundii		83.89	0.43	RAF 92	1	
Clostridium barati		16.11	0.04	GAL 96	99	
27352023	**Good identification**					**27352023**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27352100	**Low discrimination**					**27352100**
Actinomyces naeslundii		85.12	0.50	RAF 92		
Propionibacterium propionicum		12.29	0.26	GAL 96		
Actinomyces israelii		2.59	0.15			
27352101	**Good identification**					**27352101**
Actinomyces naeslundii		100.0	0.53	RAF 92		
27352102	**Low discrimination**					**27352102**
Actinomyces naeslundii		43.33	0.50	RAF 92		
Actinomyces viscosus		40.89	0.26	GAL 96		
Propionibacterium propionicum		9.38	0.29			
27352103	**Low discrimination**					**27352103**
Actinomyces naeslundii		81.58	0.53	RAF 92		
Actinomyces viscosus		17.11	0.18	GAL 96		
Propionibacterium propionicum		1.31	0.12			
27352120	**Acceptable identification**					**27352120**
					SPO	
Actinomyces naeslundii		93.68	0.50	RAF 92	1	
Clostridium barati		6.32	0.04	GAL 96	99	
27352121	**Good identification**					**27352121**
Actinomyces naeslundii		100.0	0.53	RAF 92		
27352122	**Low discrimination**					**27352122**
					SPO	
Actinomyces naeslundii		78.97	0.50	RAF 92	1	
Clostridium barati		15.98	0.12	GAL 96	99	
Lactobacillus acidophilus		2.73	0.01		1	
27352123	**Good identification**					**27352123**
Actinomyces naeslundii		100.0	0.53	RAF 92		
27352140	**Low discrimination**					**27352140**
Actinomyces naeslundii		55.00	0.37	RAF 92		
Propionibacterium propionicum		45.00	0.26	GAL 96		
27352141	**Good identification**					**27352141**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27352143	**Acceptable identification**					**27352143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.67	0.40	RAF 92		
Propionibacterium propionicum		8.33	0.12	GAL 96		
27352160	**Good identification**					**27352160**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27352161	**Good identification**					**27352161**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27352162	**Good identification**					**27352162**
Actinomyces naeslundii		100.0	0.37	RAF 92		
27352163	**Good identification**					**27352163**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27353000	**Low discrimination**					**27353000**
					SPO	
Actinomyces naeslundii		65.16	0.66	RAF 92	1	
Clostridium innocuum		16.24	0.32		99	
Lactobacillus catenaformis		7.63	0.29		1	
27353001	**Very good identification**					**27353001**
Actinomyces naeslundii		100.0	0.69	RAF 92		
27353002	**Low discrimination**					**27353002**
					SPO	
Actinomyces naeslundii		60.10	0.66	RAF 92	1	
Clostridium innocuum		14.98	0.32		99	
Actinomyces israelii		8.05	0.42		1	
27353003	**Very good identification**					**27353003**
Actinomyces naeslundii		99.15	0.69	RAF 92		
Actinomyces viscosus		0.85	0.10			
27353010	**Low discrimination**					**27353010**
Actinomyces naeslundii		54.85	0.44	RAF 92		
Lactobacillus jensenii		20.42	0.15	MLZ 5		
Actinomyces israelii		13.91	0.25			
27353011	**Good identification**					**27353011**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27353012	**Low discrimination**					**27353012**
Actinomyces naeslundii		54.77	0.44	RAF 92		
Actinomyces israelii		20.83	0.28	MLZ 5		
Lactobacillus jensenii		13.59	0.12			
27353013	**Good identification**					**27353013**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27353020	**Good identification**					**27353020**
					SPO	
Actinomyces naeslundii		90.50	0.66	RAF 92	1	
Clostridium barati		4.59	0.17		99	
Lactobacillus catenaformis		2.65	0.18		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27353021	**Very good identification**					**27353021**
Actinomyces naeslundii		100.0	0.69	RAF 92		
27353022	**Low discrimination**					**27353022**
					SPO	
Actinomyces naeslundii		80.75	0.66	RAF 92	1	
Clostridium barati		12.27	0.25		99	
Lactobacillus acidophilus		2.85	0.17		1	
27353023	**Very good identification**					**27353023**
Actinomyces naeslundii		100.0	0.69	RAF 92		
27353030	**Acceptable identification**					**27353030**
Actinomyces naeslundii		91.49	0.44	RAF 92		
Lactobacillus jensenii		8.51	0.05	MLZ 5		
27353031	**Good identification**					**27353031**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27353032	**Low discrimination**					**27353032**
					SPO	
Actinomyces naeslundii		84.75	0.44	RAF 92	1	
Clostridium tertium		10.00	0.05	MLZ 5	99	
Lactobacillus jensenii		5.26	0.02		1	
27353033	**Good identification**					**27353033**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27353040	**Low discrimination**					**27353040**
Actinomyces naeslundii		83.55	0.53	RAF 92		
Actinomyces israelii		16.45	0.32			
27353041	**Very good identification**					**27353041**
Actinomyces naeslundii		100.0	0.56	RAF 92		
27353042	**Low discrimination**					**27353042**
					SPO	
Actinomyces naeslundii		74.36	0.53	RAF 92	1	
Actinomyces israelii		21.95	0.35		1	
Clostridium barati		1.86	-.01		99	
27353043	**Very good identification**					**27353043**
Actinomyces naeslundii		100.0	0.56	RAF 92		
27353060	**Good identification**					**27353060**
					SPO	
Actinomyces naeslundii		97.56	0.53	RAF 92	1	
Clostridium barati		2.44	-.01		99	
27353061	**Very good identification**					**27353061**
Actinomyces naeslundii		100.0	0.56	RAF 92		
27353062	**Low discrimination**					**27353062**
					SPO	
Actinomyces naeslundii		89.41	0.53	RAF 92	1	
Clostridium barati		6.70	0.07		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		2.21	0.01		1	
27353063	**Very good identification**					**27353063**
Actinomyces naeslundii		100.0	0.56	RAF 92		
27353100	**Low discrimination**					**27353100**
Actinomyces naeslundii		81.27	0.74	RAF 92		
Actinomyces israelii		10.20	0.49			
Lactobacillus catenaformis		5.01	0.32			
27353101	**Very good identification**					**27353101**
Actinomyces naeslundii		99.92	0.77	RAF 92		
Actinomyces israelii		0.08	0.15			
27353102	**Low discrimination**					**27353102**
Actinomyces naeslundii		65.81	0.74	RAF 92		
Actinomyces israelii		12.39	0.52			
Lactobacillus acidophilus		9.38	0.35			
27353103	**Very good identification**					**27353103**
Actinomyces naeslundii		99.18	0.77	RAF 92		
Actinomyces viscosus		0.53	0.14			
Actinomyces israelii		0.13	0.18			
27353110	**Low discrimination**					**27353110**
					SPO	
Actinomyces naeslundii		55.18	0.52	RAF 92	1	
Clostridium tertium		25.16	0.23	MLZ 5	99	
Actinomyces israelii		19.67	0.35		1	
27353111	**Good identification**					**27353111**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27353113	**Good identification**					**27353113**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27353120	**Low discrimination**					**27353120**
					SPO	
Actinomyces naeslundii		86.56	0.74	RAF 92	1	
Clostridium barati		4.62	0.25		99	
Clostridium tertium		3.51	0.27		99	
27353121	**Very good identification**					**27353121**
Actinomyces naeslundii		100.0	0.77	RAF 92		
27353122	**Low discrimination**					**27353122**
					SPO	
Actinomyces naeslundii		64.56	0.74	RAF 92	1	
Clostridium barati		10.34	0.34		99	
Lactobacillus acidophilus		9.20	0.35		1	
27353123	**Very good identification**					**27353123**
					SPO	
Actinomyces naeslundii		99.72	0.77	RAF 92	1	
Clostridium barati		0.11	-.01		99	
Lactobacillus acidophilus		0.10	0.01		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27353131	**Good identification**					**27353131**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27353132	**Low discrimination**					**27353132**
					SPO	
Clostridium tertium		78.61	0.39	NIT 1	99	
Actinomyces naeslundii		19.16	0.52	TRE 90	1	
Actinomyces israelii		0.65	0.18		1	
27353133	**Good identification**					**27353133**
					SPO	
Actinomyces naeslundii		97.31	0.55	RAF 92	1	
Clostridium tertium		2.69	0.05	MLZ 5	99	
27353140	**Low discrimination**					**27353140**
Actinomyces naeslundii		75.79	0.61	RAF 92		
Actinomyces israelii		20.97	0.42			
Propionibacterium propionicum		2.58	0.26			
27353141	**Very good identification**					**27353141**
Actinomyces naeslundii		100.0	0.64	RAF 92		
27353142	**Low discrimination**					**27353142**
Actinomyces naeslundii		62.81	0.61	RAF 92		
Actinomyces israelii		26.06	0.45			
Lactobacillus acidophilus		6.27	0.19			
27353143	**Very good identification**					**27353143**
Actinomyces naeslundii		99.62	0.64	RAF 92		
Propionibacterium propionicum		0.38	0.12			
27353150	**Low discrimination**					**27353150**
Actinomyces naeslundii		56.01	0.39	RAF 92		
Actinomyces israelii		43.99	0.28	MLZ 5		
27353151	**Good identification**					**27353151**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27353153	**Good identification**					**27353153**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27353160	**Good identification**					**27353160**
					SPO	
Actinomyces naeslundii		94.61	0.61	RAF 92	1	
Clostridium barati		2.49	0.07		99	
Actinomyces israelii		1.67	0.22		1	
27353161	**Very good identification**					**27353161**
Actinomyces naeslundii		100.0	0.64	RAF 92		
27353162	**Low discrimination**					**27353162**
					SPO	
Actinomyces naeslundii		81.60	0.61	RAF 92	1	
Lactobacillus acidophilus		8.14	0.19		1	
Clostridium barati		6.44	0.15		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27353163	**Very good identification**					**27353163**
Actinomyces naeslundii		100.0	0.64	RAF 92		
27353170	**Good identification**					**27353170**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27353171	**Good identification**					**27353171**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27353172	**Low discrimination**					**27353172**
					SPO	
Actinomyces naeslundii		72.89	0.39	RAF 92	1	
Clostridium tertium		17.12	0.05	MLZ 5	99	
Bifidobacterium breve		9.99	0.10		1	
27353173	**Good identification**					**27353173**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27353201	**Good identification**					**27353201**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353203	**Good identification**					**27353203**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353221	**Good identification**					**27353221**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353223	**Good identification**					**27353223**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353300	**Low discrimination**					**27353300**
					SPO	
Actinomyces naeslundii		47.88	0.40	RAF 92	1	
Actinomyces israelii		44.79	0.30	RHA 1	1	
Clostridium barati		7.34	-.01		99	
27353301	**Good identification**					**27353301**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27353303	**Good identification**					**27353303**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27353321	**Good identification**					**27353321**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27353323	**Acceptable identification**					**27353323**
					SPO	
Actinomyces naeslundii		92.81	0.43	RAF 92	1	
Clostridium ramosum		7.19	-.02	RHA 1	92	
27353400	**Low discrimination**					**27353400**
					SPO	
Actinomyces israelii		57.47	0.38	XYL 92	1	
Clostridium innocuum		34.59	0.19		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		7.94	0.32		1	
27353401	**Good identification**					**27353401**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353402	**Low discrimination**					**27353402**
					SPO	
Actinomyces israelii		66.96	0.41	XYL 92	1	
Clostridium innocuum		26.87	0.19		99	
Actinomyces naeslundii		6.17	0.32		1	
27353403	**Good identification**					**27353403**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353421	**Good identification**					**27353421**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353423	**Good identification**					**27353423**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27353500	**Acceptable identification**					**27353500**
Actinomyces israelii		91.05	0.48	XYL 92		
Actinomyces naeslundii		8.95	0.40			
27353501	**Acceptable identification**					**27353501**
Actinomyces naeslundii		93.59	0.43	RAF 92		
Actinomyces israelii		6.41	0.14	ARA 1		
27353502	**Low discrimination**					**27353502**
Actinomyces israelii		89.83	0.51	XYL 92		
Actinomyces naeslundii		5.89	0.40			
Bifidobacterium adolescentis		3.45	0.20			
27353503	**Acceptable identification**					**27353503**
Actinomyces naeslundii		90.69	0.43	RAF 92		
Actinomyces israelii		9.31	0.17	ARA 1		
27353512	**Low discrimination**					**27353512**
Actinomyces israelii		76.11	0.37	XYL 92		
Bifidobacterium adolescentis		23.89	0.22	MLZ 13		
27353520	**Low discrimination**					**27353520**
Actinomyces naeslundii		60.64	0.40	RAF 92		
Actinomyces israelii		39.36	0.27	ARA 1		
27353521	**Good identification**					**27353521**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27353522	**Low discrimination**					**27353522**
					SPO	
Actinomyces naeslundii		41.71	0.40	RAF 92	1	
Actinomyces israelii		40.60	0.30	ARA 1	1	
Clostridium barati		6.68	-.01		99	
27353523	**Good identification**					**27353523**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	RAF 92		
27353540	**Good identification**					**27353540**
Actinomyces israelii		95.73	0.41	XYL 92		
Actinomyces naeslundii		4.27	0.27			
27353542	**Good identification**					**27353542**
Actinomyces israelii		97.11	0.44	XYL 92		
Actinomyces naeslundii		2.89	0.27			
27354000	**Good identification**					**27354000**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27354200	**Good identification**					**27354200**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27354202	**Good identification**					**27354202**
Actinomyces odontolyticus		100.0	0.40	MNS 9		
27355000	**Low discrimination**					**27355000**
Actinomyces israelii		34.50	0.44			
Actinomyces odontolyticus		31.05	0.35			
Lactobacillus catenaformis		22.28	0.29			
27355002	**Low discrimination**					**27355002**
Actinomyces israelii		46.51	0.47			
Lactobacillus catenaformis		20.03	0.29			
Actinomyces odontolyticus		17.11	0.32			
27355040	**Good identification**					**27355040**
Actinomyces israelii		100.0	0.37			
27355042	**Good identification**					**27355042**
Actinomyces israelii		100.0	0.40			
27355100	**Low discrimination**					**27355100**
					COC	
Actinomyces israelii		67.32	0.55		1	
Lactobacillus catenaformis		16.30	0.32		1	
Atopobium parvulum		7.76	0.25		99	
27355101	**Low discrimination**					**27355101**
Actinomyces naeslundii		75.21	0.38	RAF 92		
Actinomyces israelii		24.79	0.20	XYL 1		
27355102	**Low discrimination**					**27355102**
Actinomyces israelii		71.93	0.58			
Lactobacillus catenaformis		11.61	0.32			
Lactobacillus acidophilus		5.15	0.23			
27355110	**Good identification**					**27355110**
Actinomyces israelii		100.0	0.40	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27355112	**Low discrimination**					**27355112**
					SPO	
Actinomyces israelii		85.31	0.43	MLZ 13	1	
Bifidobacterium adolescentis		13.19	0.23		1	
Clostridium tertium		1.50	-.01		99	
27355121	**Good identification**					**27355121**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27355123	**Good identification**					**27355123**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27355140	**Good identification**					**27355140**
Actinomyces israelii		100.0	0.48			
27355142	**Good identification**					**27355142**
Actinomyces israelii		97.78	0.51			
Lactobacillus acidophilus		2.22	0.07			
27355152	**Good identification**					**27355152**
Actinomyces israelii		95.31	0.36	MLZ 13		
Bifidobacterium adolescentis		4.69	0.08			
27355200	**Good identification**					**27355200**
Actinomyces odontolyticus		95.87	0.40	MNS 9		
Actinomyces israelii		4.13	0.25	CEL 1		
27355202	**Acceptable identification**					**27355202**
Actinomyces odontolyticus		90.46	0.37	MNS 9		
Actinomyces israelii		9.54	0.28	CEL 1		
27355302	**Low discrimination**					**27355302**
Actinomyces israelii		54.93	0.38	RHA 7		
Actinomyces odontolyticus		41.14	0.28			
Lactobacillus acidophilus		3.93	0.04			
27355400	**Good identification**					**27355400**
Actinomyces israelii		100.0	0.43			
27355402	**Good identification**					**27355402**
Actinomyces israelii		100.0	0.46			
27355440	**Good identification**					**27355440**
Actinomyces israelii		100.0	0.36			
27355442	**Good identification**					**27355442**
Actinomyces israelii		100.0	0.39			
27355500	**Good identification**					**27355500**
					SPO	
Actinomyces israelii		95.83	0.53		1	
Clostridium butyricum		4.17	0.09		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27355502	**Low discrimination**					**27355502**
					SPO	
Actinomyces israelii		85.77	0.56		1	
Bifidobacterium adolescentis		13.26	0.36		1	
Clostridium butyricum		0.97	0.02		99	
27355510	**Good identification**					**27355510**
Actinomyces israelii		100.0	0.39	MLZ 13		
27355512	**Low discrimination**					**27355512**
Bifidobacterium adolescentis		55.08	0.37	NIT 1		
Actinomyces israelii		43.55	0.42	RAF 92		
Clostridium clostridioforme		1.37	0.07			
27355522	**Good identification**					**27355522**
Actinomyces israelii		100.0	0.36	NAG 6		
27355540	**Good identification**					**27355540**
Actinomyces israelii		100.0	0.46			
27355542	**Good identification**					**27355542**
Actinomyces israelii		95.32	0.49			
Bifidobacterium adolescentis		4.69	0.20			
27355702	**Acceptable identification**					**27355702**
Actinomyces israelii		90.86	0.37	RHA 7		
Clostridium clostridioforme		9.14	0.10			
27356102	**Good identification**					**27356102**
Actinomyces israelii		100.0	0.36	GAL 99		
27357000	**Very good identification**					**27357000**
					SPO	
Actinomyces israelii		97.79	0.57		1	
Clostridium innocuum		1.25	0.10		99	
Actinomyces naeslundii		0.96	0.32		1	
27357001	**Low discrimination**					**27357001**
Actinomyces naeslundii		59.36	0.35	RAF 92		
Actinomyces israelii		40.64	0.23	XYL 1		
27357002	**Very good identification**					**27357002**
					SPO	
Actinomyces israelii		98.52	0.60		1	
Clostridium innocuum		0.84	0.10		99	
Actinomyces naeslundii		0.65	0.32		1	
27357010	**Good identification**					**27357010**
Actinomyces israelii		100.0	0.43	MLZ 13		
27357012	**Good identification**					**27357012**
					SPO	
Actinomyces israelii		98.77	0.46	MLZ 13	1	
Clostridium tertium		1.23	-.01		99	
27357020	**Low discrimination**					**27357020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		86.65	0.37	NAG 6		
Actinomyces naeslundii		13.35	0.32			
27357021	**Good identification**					**27357021**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27357022	**Low discrimination**					**27357022**
					SPO	
Actinomyces israelii		86.83	0.40	NAG 6	1	
Actinomyces naeslundii		8.92	0.32		1	
Clostridium tertium		4.26	0.03		99	
27357023	**Good identification**					**27357023**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27357040	**Good identification**					**27357040**
Actinomyces israelii		100.0	0.50			
27357042	**Good identification**					**27357042**
Actinomyces israelii		100.0	0.53			
27357050	**Good identification**					**27357050**
Actinomyces israelii		100.0	0.36	MLZ 13		
27357052	**Good identification**					**27357052**
Actinomyces israelii		100.0	0.39	MLZ 13		
27357100	**Very good identification**					**27357100**
					SPO	
Actinomyces israelii		98.05	0.67		1	
Clostridium tertium		1.27	0.21		99	
Actinomyces naeslundii		0.69	0.40		1	
27357101	**Low discrimination**					**27357101**
Actinomyces naeslundii		50.96	0.43	RAF 92		
Actinomyces israelii		49.04	0.33	XYL 1		
27357102	**Very good identification**					**27357102**
					SPO	
Actinomyces israelii		96.72	0.70		1	
Clostridium tertium		2.50	0.29		99	
Actinomyces naeslundii		0.45	0.40		1	
27357103	**Low discrimination**					**27357103**
Actinomyces israelii		59.08	0.36	URE 1		
Actinomyces naeslundii		40.92	0.43			
27357110	**Low discrimination**					**27357110**
					SPO	
Actinomyces israelii		86.69	0.53	MLZ 13	1	
Clostridium tertium		13.32	0.25		99	
27357112	**Low discrimination**					**27357112**
					SPO	
Actinomyces israelii		75.21	0.56	MLZ 13	1	
Clostridium tertium		23.11	0.33		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		1.68	0.22		1	
27357120	**Low discrimination**					**27357120**
					SPO	
Actinomyces israelii		58.26	0.47	NAG 6	1	
Clostridium tertium		35.35	0.29		99	
Actinomyces naeslundii		6.39	0.40		1	
27357121	**Good identification**					**27357121**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27357122	**Low discrimination**					**27357122**
					SPO	
Clostridium tertium		52.56	0.37	NIT 1	99	
Actinomyces israelii		43.31	0.50	TRE 90	1	
Actinomyces naeslundii		3.17	0.40		1	
27357123	**Low discrimination**					**27357123**
					SPO	
Actinomyces naeslundii		83.06	0.43	RAF 92	1	
Clostridium tertium		9.29	0.03	XYL 1	99	
Actinomyces israelii		7.65	0.16		1	
27357132	**Acceptable identification**					**27357132**
					SPO	
Clostridium tertium		93.52	0.41	NIT 1	99	
Actinomyces israelii		6.48	0.36	TRE 90	1	
27357140	**Very good identification**					**27357140**
Actinomyces israelii		99.68	0.60			
Actinomyces naeslundii		0.32	0.27			
27357142	**Very good identification**					**27357142**
Actinomyces israelii		99.79	0.63			
Actinomyces naeslundii		0.21	0.27			
27357150	**Good identification**					**27357150**
Actinomyces israelii		100.0	0.46	MLZ 13		
27357152	**Good identification**					**27357152**
					SPO	
Actinomyces israelii		99.21	0.49	MLZ 13	1	
Clostridium tertium		0.79	-.01		99	
27357160	**Good identification**					**27357160**
Actinomyces israelii		95.26	0.40	NAG 6		
Actinomyces naeslundii		4.74	0.27			
27357162	**Acceptable identification**					**27357162**
					SPO	
Actinomyces israelii		93.93	0.43	NAG 6	1	
Actinomyces naeslundii		3.12	0.27		1	
Clostridium tertium		2.96	0.03		99	
27357200	**Good identification**					**27357200**
Actinomyces israelii		96.03	0.38	RHA 7		
Actinomyces odontolyticus		3.98	0.06			
27357202	**Good identification**					**27357202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.41	RHA 7		
27357300	**Good identification**					**27357300**
Actinomyces israelii		100.0	0.48	RHA 7		
27357302	**Good identification**					**27357302**
Actinomyces israelii		100.0	0.51	RHA 7		
27357312	**Good identification**					**27357312**
					SPO	
Actinomyces israelii		96.04	0.37	RHA 7	1	
Clostridium tertium		3.96	-.01	MLZ 13	99	
27357340	**Good identification**					**27357340**
Actinomyces israelii		100.0	0.41	RHA 7		
27357342	**Good identification**					**27357342**
Actinomyces israelii		100.0	0.44	RHA 7		
27357400	**Good identification**					**27357400**
Actinomyces israelii		100.0	0.56			
27357402	**Very good identification**					**27357402**
Actinomyces israelii		100.0	0.59			
27357410	**Good identification**					**27357410**
Actinomyces israelii		100.0	0.42	MLZ 13		
27357412	**Good identification**					**27357412**
Actinomyces israelii		100.0	0.45	MLZ 13		
27357420	**Good identification**					**27357420**
Actinomyces israelii		100.0	0.35	NAG 6		
27357422	**Good identification**					**27357422**
Actinomyces israelii		100.0	0.38	NAG 6		
27357440	**Good identification**					**27357440**
Actinomyces israelii		100.0	0.49			
27357442	**Good identification**					**27357442**
Actinomyces israelii		100.0	0.52			
27357452	**Good identification**					**27357452**
Actinomyces israelii		100.0	0.38	MLZ 13		
27357500	**Very good identification**					**27357500**
					SPO	
Actinomyces israelii		99.92	0.66		1	
Clostridium butyricum		0.09	-.08		99	
27357502	**Very good identification**					**27357502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		97.82	0.69			
Bifidobacterium adolescentis		2.18	0.34			
27357510	**Good identification**					**27357510**
Actinomyces israelii		100.0	0.52	MLZ 13		
27357512	**Low discrimination**					**27357512**
					SPO	
Actinomyces israelii		84.29	0.55	MLZ 13	1	
Bifidobacterium adolescentis		15.39	0.36		1	
Clostridium tertium		0.32	-.01		99	
27357520	**Good identification**					**27357520**
Actinomyces israelii		100.0	0.46	NAG 6		
27357522	**Good identification**					**27357522**
					SPO	
Actinomyces israelii		98.52	0.49	NAG 6	1	
Clostridium tertium		1.48	0.03		99	
27357540	**Very good identification**					**27357540**
Actinomyces israelii		100.0	0.59			
27357542	**Very good identification**					**27357542**
Actinomyces israelii		99.30	0.62			
Bifidobacterium adolescentis		0.71	0.19			
27357550	**Good identification**					**27357550**
Actinomyces israelii		100.0	0.45	MLZ 13		
27357552	**Acceptable identification**					**27357552**
Actinomyces israelii		94.51	0.48	MLZ 13		
Bifidobacterium adolescentis		5.49	0.20			
27357560	**Good identification**					**27357560**
Actinomyces israelii		100.0	0.39	NAG 6		
27357562	**Good identification**					**27357562**
Actinomyces israelii		100.0	0.42	NAG 6		
27357600	**Good identification**					**27357600**
Actinomyces israelii		100.0	0.36	RHA 7		
27357602	**Good identification**					**27357602**
Actinomyces israelii		100.0	0.39	RHA 7		
27357700	**Good identification**					**27357700**
Actinomyces israelii		100.0	0.47	RHA 7		
27357702	**Good identification**					**27357702**
Actinomyces israelii		100.0	0.50	RHA 7		
27357712	**Good identification**					**27357712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	RHA 7		
27357740	**Good identification**					**27357740**
Actinomyces israelii		100.0	0.40	RHA 7		
27357742	**Good identification**					**27357742**
Actinomyces israelii		100.0	0.43	RHA 7		
27361001	**Good identification**					**27361001**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27361003	**Acceptable identification**					**27361003**
Actinomyces naeslundii		94.00	0.47	RAF 92		
Actinomyces viscosus		6.00	0.03	FRU 95		
27361020	**Low discrimination**					**27361020**
Actinomyces naeslundii		49.94	0.44	RAF 92		
Lactobacillus jensenii		41.83	0.21	FRU 95		
Clostridium perfringens		8.23	0.09			
27361021	**Good identification**					**27361021**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27361022	**Low discrimination**					**27361022**
Actinomyces naeslundii		48.39	0.44	RAF 92		
Lactobacillus jensenii		27.02	0.18	FRU 95		
Lactobacillus acidophilus		20.83	0.14			
27361023	**Good identification**					**27361023**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27361100	**Low discrimination**					**27361100**
					COC	
Actinomyces naeslundii		76.90	0.52	RAF 92	1	
Atopobium parvulum		23.10	0.20	FRU 95	99	
27361101	**Good identification**					**27361101**
					SPO	
Actinomyces naeslundii		97.04	0.55	RAF 92	1	
Clostridium chauvoei		2.96	0.06	FRU 95	99	
27361103	**Good identification**					**27361103**
Actinomyces naeslundii		96.16	0.55	RAF 92		
Actinomyces viscosus		3.84	0.07	FRU 95		
27361120	**Low discrimination**					**27361120**
					COC	
Actinomyces naeslundii		67.67	0.52	RAF 92	1	
Clostridium perfringens		28.74	0.23	FRU 95	1	
Atopobium parvulum		3.59	0.07		99	
27361121	**Good identification**					**27361121**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27361123	**Good identification**					**27361123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	RAF 92		
27361140	**Good identification**					**27361140**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27361141	**Good identification**					**27361141**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27361143	**Good identification**					**27361143**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27361160	**Low discrimination**					**27361160**
Actinomyces naeslundii		68.57	0.39	RAF 92		
Clostridium perfringens		31.43	0.11	FRU 95		
27361161	**Good identification**					**27361161**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27361163	**Good identification**					**27361163**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27362101	**Good identification**					**27362101**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27362103	**Good identification**					**27362103**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27362121	**Good identification**					**27362121**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27362123	**Good identification**					**27362123**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27363000	**Acceptable identification**					**27363000**
					SPO	
Actinomyces naeslundii		92.74	0.49	RAF 92	1	
Clostridium innocuum		4.18	0.02	FRU 95	99	
Actinomyces israelii		3.08	0.15		1	
27363001	**Good identification**					**27363001**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27363002	**Acceptable identification**					**27363002**
					SPO	
Actinomyces naeslundii		91.33	0.49	RAF 92	1	
Actinomyces israelii		4.55	0.18	FRU 95	1	
Clostridium innocuum		4.12	0.02		99	
27363003	**Good identification**					**27363003**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27363020	**Good identification**					**27363020**
Actinomyces naeslundii		100.0	0.49	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27363021	**Good identification**					**27363021**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27363022	**Good identification**					**27363022**
Actinomyces naeslundii		100.0	0.49	RAF 92		
27363023	**Good identification**					**27363023**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27363040	**Good identification**					**27363040**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27363041	**Good identification**					**27363041**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27363042	**Good identification**					**27363042**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27363043	**Good identification**					**27363043**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27363060	**Good identification**					**27363060**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27363061	**Good identification**					**27363061**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27363062	**Good identification**					**27363062**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27363063	**Good identification**					**27363063**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27363100	**Good identification**					**27363100**
					SPO	
Actinomyces naeslundii		94.46	0.57	RAF 92	1	
Actinomyces israelii		4.41	0.25	FRU 95	1	
Clostridium tertium		1.13	0.01		99	
27363101	**Very good identification**					**27363101**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27363102	**Low discrimination**					**27363102**
Actinomyces naeslundii		86.71	0.57	RAF 92		
Actinomyces israelii		6.07	0.28	FRU 95		
Lactobacillus acidophilus		4.09	0.10			
27363103	**Very good identification**					**27363103**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27363110	**Low discrimination**					**27363110**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		71.14	0.35	RAF 92	1	
Clostridium tertium		28.86	0.05	FRU 95	99	
27363111	**Good identification**					**27363111**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27363113	**Good identification**					**27363113**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27363120	**Good identification**					**27363120**
					SPO	
Actinomyces naeslundii		93.46	0.57	RAF 92	1	
Clostridium tertium		3.37	0.09	FRU 95	99	
Clostridium ramosum		3.17	0.07		92	
27363121	**Very good identification**					**27363121**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27363122	**Low discrimination**					**27363122**
					SPO	
Actinomyces naeslundii		77.46	0.57	RAF 92	1	
Clostridium ramosum		10.52	0.17	FRU 95	92	
Clostridium tertium		8.37	0.17		99	
27363123	**Very good identification**					**27363123**
Actinomyces naeslundii		100.0	0.60	RAF 92		
27363131	**Good identification**					**27363131**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27363133	**Good identification**					**27363133**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27363140	**Acceptable identification**					**27363140**
Actinomyces naeslundii		90.67	0.44	RAF 92		
Actinomyces israelii		9.33	0.18	FRU 95		
27363141	**Good identification**					**27363141**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27363142	**Low discrimination**					**27363142**
Actinomyces naeslundii		86.63	0.44	RAF 92		
Actinomyces israelii		13.37	0.21	FRU 95		
27363143	**Good identification**					**27363143**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27363160	**Good identification**					**27363160**
Actinomyces naeslundii		100.0	0.44	RAF 92		
27363161	**Good identification**					**27363161**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27363162	**Good identification**					**27363162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	RAF 92		
27363163	**Good identification**					**27363163**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27367002	**Good identification**					**27367002**
Actinomyces israelii		100.0	0.36	FRU 99		
27367100	**Good identification**					**27367100**
					SPO	
Actinomyces israelii		95.27	0.43	FRU 99	1	
Clostridium tertium		2.94	0.03		99	
Actinomyces naeslundii		1.79	0.23		1	
27367102	**Acceptable identification**					**27367102**
					SPO	
Actinomyces israelii		93.08	0.46	FRU 99	1	
Clostridium tertium		5.75	0.11		99	
Actinomyces naeslundii		1.17	0.23		1	
27367140	**Good identification**					**27367140**
Actinomyces israelii		100.0	0.36	FRU 99		
27367142	**Good identification**					**27367142**
Actinomyces israelii		100.0	0.39	FRU 99		
27367500	**Good identification**					**27367500**
Actinomyces israelii		100.0	0.42	FRU 99		
27367502	**Good identification**					**27367502**
Actinomyces israelii		100.0	0.45	FRU 99		
27367542	**Good identification**					**27367542**
Actinomyces israelii		100.0	0.38	FRU 99		
27370001	**Good identification**					**27370001**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27370002	**Low discrimination**					**27370002**
Actinomyces viscosus		71.72	0.36	RAF 94		
Lactobacillus jensenii		17.97	0.29	TRE 7		
Actinomyces naeslundii		6.37	0.43			
27370021	**Good identification**					**27370021**
Actinomyces naeslundii		100.0	0.46	RAF 92		
27370022	**Low discrimination**					**27370022**
Actinomyces naeslundii		37.70	0.43	RAF 92		
Lactobacillus jensenii		26.59	0.18	GAL 96		
Lactobacillus acidophilus		20.50	0.14			
27370023	**Good identification**					**27370023**
Actinomyces naeslundii		100.0	0.46	RAF 92		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
27370100	**Low discrimination**						**27370100**
					COC	GRA	
Actinomyces naeslundii		64.14	0.50	RAF 92	1	99	
Atopobium parvulum		24.33	0.20	GAL 96	99	99	
Fusobacterium mortiferum		6.97	0.22		1	1	
27370101	**Good identification**						**27370101**
Actinomyces naeslundii		98.45	0.53	RAF 92			
Actinomyces viscosus		1.55	-.02	GAL 96			
27370102	**Low discrimination**						**27370102**
Actinomyces viscosus		67.66	0.41	RAF 94			
Lactobacillus acidophilus		21.14	0.32	TRE 7			
Actinomyces naeslundii		9.62	0.50				
27370120	**Low discrimination**						**27370120**
					SPO		
Actinomyces naeslundii		59.85	0.50	RAF 92	1		
Clostridium perfringens		32.11	0.23	GAL 96	84		
Clostridium barati		4.04	0.04		99		
27370121	**Good identification**						**27370121**
Actinomyces naeslundii		100.0	0.53	RAF 92			
27370123	**Good identification**						**27370123**
Actinomyces naeslundii		98.45	0.53	RAF 92			
Actinomyces viscosus		1.55	-.02	GAL 96			
27370140	**Good identification**						**27370140**
Actinomyces naeslundii		100.0	0.37	RAF 92			
27370141	**Good identification**						**27370141**
Actinomyces naeslundii		100.0	0.40	RAF 92			
27370143	**Acceptable identification**						**27370143**
Actinomyces naeslundii		91.79	0.40	RAF 92			
Actinomyces viscosus		8.21	-.02	GAL 96			
27370160	**Low discrimination**						**27370160**
Actinomyces naeslundii		63.33	0.37	RAF 92			
Clostridium perfringens		36.67	0.11	GAL 96			
27370161	**Good identification**						**27370161**
Actinomyces naeslundii		100.0	0.40	RAF 92			
27370163	**Good identification**						**27370163**
Actinomyces naeslundii		100.0	0.40	RAF 92			
27371000	**Good identification**						**27371000**
Lactobacillus jensenii		93.06	0.66	NIT 1			
Actinomyces naeslundii		5.33	0.66				
Lactobacillus catenaformis		0.62	0.29				
27371001	**Low discrimination**						**27371001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		89.48	0.69	RAF 92		
Lactobacillus jensenii		10.52	0.32			
27371002	**Low discrimination**					**27371002**
Lactobacillus jensenii		75.35	0.63	NIT 1		
Lactobacillus acidophilus		14.52	0.48			
Actinomyces naeslundii		6.48	0.66			
27371003	**Low discrimination**					**27371003**
Actinomyces naeslundii		86.40	0.69	RAF 92		
Lactobacillus jensenii		6.77	0.29			
Actinomyces viscosus		5.52	0.25			
27371010	**Very good identification**					**27371010**
Lactobacillus jensenii		98.95	0.58	NIT 1		
Actinomyces naeslundii		0.90	0.44			
Actinomyces israelii		0.15	0.22			
27371011	**Low discrimination**					**27371011**
Actinomyces naeslundii		57.32	0.47	RAF 92		
Lactobacillus jensenii		42.68	0.24	MLZ 5		
27371012	**Good identification**					**27371012**
					GRA	
Lactobacillus jensenii		95.12	0.55	NIT 1	99	
Leptotrichia buccalis		2.64	0.15		1	
Actinomyces naeslundii		1.29	0.44		99	
27371013	**Low discrimination**					**27371013**
Actinomyces naeslundii		66.83	0.47	RAF 92		
Lactobacillus jensenii		33.17	0.21	MLZ 5		
27371020	**Low discrimination**					**27371020**
Lactobacillus jensenii		68.86	0.56	NIT 1		
Actinomyces naeslundii		15.78	0.66			
Clostridium perfringens		13.55	0.43			
27371021	**Very good identification**					**27371021**
Actinomyces naeslundii		96.60	0.69	RAF 92		
Lactobacillus jensenii		2.84	0.21			
Clostridium perfringens		0.56	0.09			
27371022	**Low discrimination**					**27371022**
Lactobacillus jensenii		43.08	0.53	NIT 1		
Lactobacillus acidophilus		33.21	0.48			
Actinomyces naeslundii		14.81	0.66			
27371023	**Very good identification**					**27371023**
Actinomyces naeslundii		96.39	0.69	RAF 92		
Lactobacillus jensenii		1.89	0.18			
Lactobacillus acidophilus		1.46	0.14			
27371030	**Good identification**					**27371030**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		95.96	0.47	NIT 1			
Actinomyces naeslundii		3.47	0.44				
Clostridium perfringens		0.57	0.09				
27371031	**Low discrimination**						**27371031**
Actinomyces naeslundii		84.31	0.47	RAF 92			
Lactobacillus jensenii		15.69	0.13	MLZ 5			
27371032	**Acceptable identification**						**27371032**
Lactobacillus jensenii		91.39	0.44	NIT 1			
Actinomyces naeslundii		4.96	0.44				
Lactobacillus acidophilus		2.14	0.14				
27371033	**Low discrimination**						**27371033**
Actinomyces naeslundii		88.96	0.47	RAF 92			
Lactobacillus jensenii		11.04	0.10	MLZ 5			
27371040	**Low discrimination**						**27371040**
Actinomyces naeslundii		45.99	0.53	RAF 92			
Lactobacillus jensenii		45.95	0.32				
Actinomyces israelii		6.01	0.29				
27371041	**Very good identification**						**27371041**
Actinomyces naeslundii		100.0	0.56	RAF 92			
27371043	**Very good identification**						**27371043**
Actinomyces naeslundii		98.95	0.56	RAF 92			
Lactobacillus acidophilus		1.05	-.02				
27371060	**Low discrimination**						**27371060**
Actinomyces naeslundii		45.09	0.53	RAF 92			
Clostridium perfringens		41.80	0.30				
Lactobacillus jensenii		11.26	0.21				
27371061	**Very good identification**						**27371061**
Actinomyces naeslundii		100.0	0.56	RAF 92			
27371063	**Very good identification**						**27371063**
Actinomyces naeslundii		98.95	0.56	RAF 92			
Lactobacillus acidophilus		1.05	-.02				
27371100	**Low discrimination**						**27371100**
					COC		
Atopobium parvulum		52.80	0.54	NIT 1	99		
Actinomyces naeslundii		33.74	0.74		1		
Lactobacillus acidophilus		3.09	0.32		1		
27371101	**Very good identification**						**27371101**
					COC	SPO	
Actinomyces naeslundii		98.59	0.77	RAF 92	1	1	
Atopobium parvulum		1.04	0.20		99	1	
Clostridium chauvoei		0.37	0.12		1	99	
27371102	**Low discrimination**						**27371102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Lactobacillus acidophilus		83.33	0.66	NIT 1	1	
Actinomyces naeslundii		9.19	0.74		1	
Atopobium parvulum		2.54	0.41		99	
27371103	**Good identification**					**27371103**
Actinomyces naeslundii		90.56	0.77	RAF 92		
Lactobacillus acidophilus		5.53	0.32			
Actinomyces viscosus		3.61	0.29			
27371110	**Low discrimination**					**27371110**
					GRA	
Actinomyces naeslundii		37.19	0.52	RAF 92	99	
Leptotrichia buccalis		26.71	0.15	MLZ 5	1	
Lactobacillus jensenii		14.59	0.24		99	
27371111	**Good identification**					**27371111**
Actinomyces naeslundii		100.0	0.55	RAF 92		
27371112	**Good identification**					**27371112**
					GRA	
Leptotrichia buccalis		95.22	0.50	NIT 10	1	
Lactobacillus acidophilus		2.33	0.32	ESL 99	99	
Actinomyces naeslundii		1.34	0.52		99	
27371113	**Low discrimination**					**27371113**
					GRA	
Actinomyces naeslundii		67.63	0.55	RAF 92	99	
Leptotrichia buccalis		32.37	0.15	MLZ 5	1	
27371120	**Low discrimination**					**27371120**
					COC	
Clostridium perfringens		57.12	0.58	CEL 4	1	
Actinomyces naeslundii		25.81	0.74		1	
Atopobium parvulum		7.13	0.41		99	
27371121	**Very good identification**					**27371121**
					COC	
Actinomyces naeslundii		98.23	0.77	RAF 92	1	
Clostridium perfringens		1.46	0.23		1	
Atopobium parvulum		0.18	0.07		99	
27371122	**Low discrimination**					**27371122**
Lactobacillus acidophilus		77.81	0.66	NIT 1		
Clostridium perfringens		8.94	0.52			
Actinomyces naeslundii		8.59	0.74			
27371123	**Good identification**					**27371123**
Actinomyces naeslundii		93.39	0.77	RAF 92		
Lactobacillus acidophilus		5.70	0.32			
Clostridium perfringens		0.66	0.18			
27371130	**Low discrimination**					**27371130**
					SPO	
Actinomyces naeslundii		56.18	0.52	RAF 92	1	
Clostridium perfringens		23.86	0.23	MLZ 5	84	
Clostridium tertium		6.99	0.13		99	
27371131	**Good identification**					**27371131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	RAF 92		
27371133	**Acceptable identification**					**27371133**
					GRA	
Actinomyces naeslundii		94.95	0.55	RAF 92	99	
Leptotrichia buccalis		5.05	-.01	MLZ 5	1	
27371140	**Low discrimination**					**27371140**
					COC	
Actinomyces naeslundii		70.42	0.61	RAF 92	1	
Actinomyces israelii		12.93	0.39		1	
Atopobium parvulum		6.31	0.20		99	
27371141	**Very good identification**					**27371141**
Actinomyces naeslundii		100.0	0.64	RAF 92		
27371142	**Low discrimination**					**27371142**
Lactobacillus acidophilus		82.33	0.50	NIT 1		
Actinomyces naeslundii		12.97	0.61	SOR 11		
Actinomyces israelii		3.57	0.42			
27371143	**Very good identification**					**27371143**
Actinomyces naeslundii		95.90	0.64	RAF 92		
Lactobacillus acidophilus		4.10	0.16			
27371150	**Low discrimination**					**27371150**
Actinomyces naeslundii		65.73	0.39	RAF 92		
Actinomyces israelii		34.27	0.25	MLZ 5		
27371151	**Good identification**					**27371151**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27371153	**Good identification**					**27371153**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27371160	**Low discrimination**					**27371160**
Clostridium perfringens		67.57	0.45	CEL 4		
Actinomyces naeslundii		28.29	0.61			
Lactobacillus acidophilus		1.81	0.16			
27371161	**Very good identification**					**27371161**
Actinomyces naeslundii		98.42	0.64	RAF 92		
Clostridium perfringens		1.58	0.11			
27371162	**Low discrimination**					**27371162**
Lactobacillus acidophilus		73.86	0.50	NIT 1		
Clostridium perfringens		13.08	0.40	SOR 11		
Actinomyces naeslundii		11.64	0.61			
27371163	**Very good identification**					**27371163**
Actinomyces naeslundii		95.21	0.64	RAF 92		
Lactobacillus acidophilus		4.07	0.16			
Clostridium perfringens		0.72	0.05			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27371170	**Low discrimination**					**27371170**
Actinomyces naeslundii		68.57	0.39	RAF 92		
Clostridium perfringens		31.43	0.11	MLZ 5		
27371171	**Good identification**					**27371171**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27371173	**Good identification**					**27371173**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27371201	**Good identification**					**27371201**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371203	**Good identification**					**27371203**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371221	**Good identification**					**27371221**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371223	**Good identification**					**27371223**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371301	**Good identification**					**27371301**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27371302	**Good identification**					**27371302**
Lactobacillus acidophilus		95.52	0.47	NIT 1		
Actinomyces naeslundii		1.41	0.40	RHA 7		
Actinomyces israelii		1.32	0.30			
27371303	**Low discrimination**					**27371303**
Actinomyces naeslundii		68.74	0.43	RAF 92		
Lactobacillus acidophilus		31.26	0.12	RHA 1		
27371321	**Good identification**					**27371321**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27371322	**Low discrimination**					**27371322**
					SPO	
Lactobacillus acidophilus		65.43	0.47	NIT 1	1	
Clostridium ramosum		30.83	0.40	RHA 7	92	
Clostridium barati		1.34	0.15		99	
27371323	**Low discrimination**					**27371323**
					SPO	
Actinomyces naeslundii		59.92	0.43	RAF 92	1	
Lactobacillus acidophilus		27.25	0.12	RHA 1	1	
Clostridium ramosum		12.84	0.06		92	
27371401	**Good identification**					**27371401**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371403	**Good identification**					**27371403**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371421	**Good identification**					**27371421**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371423	**Good identification**					**27371423**
Actinomyces naeslundii		100.0	0.35	RAF 92		
27371500	**Low discrimination**					**27371500**
					COC	
Actinomyces israelii		69.23	0.45	XYL 92	1	
Atopobium parvulum		16.05	0.20		99	
Actinomyces naeslundii		10.26	0.40		1	
27371501	**Good identification**					**27371501**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27371502	**Low discrimination**					**27371502**
Actinomyces israelii		47.56	0.48	XYL 92		
Lactobacillus acidophilus		42.57	0.32			
Actinomyces naeslundii		4.70	0.40			
27371503	**Acceptable identification**					**27371503**
Actinomyces naeslundii		93.62	0.43	RAF 92		
Actinomyces israelii		6.38	0.14	ARA 1		
27371521	**Good identification**					**27371521**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27371523	**Good identification**					**27371523**
Actinomyces naeslundii		100.0	0.43	RAF 92		
27371540	**Acceptable identification**					**27371540**
Actinomyces israelii		93.70	0.38	XYL 92		
Actinomyces naeslundii		6.30	0.27			
27371542	**Low discrimination**					**27371542**
Actinomyces israelii		75.23	0.41	XYL 92		
Lactobacillus acidophilus		21.40	0.16			
Actinomyces naeslundii		3.37	0.27			
27372000	**Acceptable identification**					**27372000**
					SPO	
Actinomyces naeslundii		91.00	0.47	RAF 92	1	
Clostridium innocuum		5.18	0.02	GAL 96	99	
Actinomyces israelii		3.82	0.15		1	
27372001	**Good identification**					**27372001**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27372002	**Low discrimination**					**27372002**
Actinomyces naeslundii		84.86	0.47	RAF 92		
Actinomyces israelii		5.34	0.18	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		4.97	0.02			
27372003	**Good identification**					**27372003**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27372020	**Good identification**					**27372020**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27372021	**Good identification**					**27372021**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27372022	**Good identification**					**27372022**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27372023	**Good identification**					**27372023**
Actinomyces naeslundii		100.0	0.50	RAF 92		
27372041	**Good identification**					**27372041**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27372043	**Good identification**					**27372043**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27372061	**Good identification**					**27372061**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27372063	**Good identification**					**27372063**
Actinomyces naeslundii		100.0	0.38	RAF 92		
27372100	**Low discrimination**					**27372100**
Actinomyces naeslundii		88.11	0.55	RAF 92		
Propionibacterium propionicum		5.36	0.24	GAL 96		
Actinomyces israelii		5.20	0.25			
27372101	**Very good identification**					**27372101**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27372102	**Low discrimination**					**27372102**
Actinomyces naeslundii		75.68	0.55	RAF 92		
Propionibacterium propionicum		6.91	0.27	GAL 96		
Actinomyces israelii		6.69	0.28			
27372103	**Very good identification**					**27372103**
Actinomyces naeslundii		99.19	0.58	RAF 92		
Actinomyces viscosus		0.81	-.02	GAL 96		
27372111	**Good identification**					**27372111**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27372113	**Good identification**					**27372113**
Actinomyces naeslundii		100.0	0.36	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27372120	**Acceptable identification**					**27372120**
					SPO	
Actinomyces naeslundii		91.88	0.55	RAF 92	1	
Clostridium tertium		4.18	0.09	GAL 96	99	
Clostridium ramosum		3.94	0.07		92	
27372121	**Very good identification**					**27372121**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27372122	**Low discrimination**					**27372122**
					SPO	
Actinomyces naeslundii		73.12	0.55	RAF 92	1	
Clostridium ramosum		12.55	0.17	GAL 96	92	
Clostridium tertium		9.98	0.17		99	
27372123	**Very good identification**					**27372123**
Actinomyces naeslundii		100.0	0.58	RAF 92		
27372131	**Good identification**					**27372131**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27372133	**Good identification**					**27372133**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27372140	**Low discrimination**					**27372140**
Actinomyces naeslundii		67.81	0.42	RAF 92		
Propionibacterium propionicum		23.38	0.24	GAL 96		
Actinomyces israelii		8.81	0.18			
27372141	**Good identification**					**27372141**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27372142	**Low discrimination**					**27372142**
Actinomyces naeslundii		58.41	0.42	RAF 92		
Propionibacterium propionicum		30.21	0.27	GAL 96		
Actinomyces israelii		11.39	0.21			
27372143	**Good identification**					**27372143**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27372160	**Good identification**					**27372160**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27372161	**Good identification**					**27372161**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27372162	**Good identification**					**27372162**
Actinomyces naeslundii		100.0	0.42	RAF 92		
27372163	**Good identification**					**27372163**
Actinomyces naeslundii		100.0	0.45	RAF 92		
27373000	**Low discrimination**					**27373000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		76.29	0.71	RAF 92		
Actinomyces israelii		13.21	0.49			
Lactobacillus jensenii		6.93	0.32			
27373001	**Very good identification**					**27373001**
Actinomyces naeslundii		99.88	0.74	RAF 92		
Actinomyces israelii		0.12	0.15			
27373002	**Low discrimination**					**27373002**
Actinomyces naeslundii		69.67	0.71	RAF 92		
Actinomyces israelii		18.09	0.52			
Lactobacillus acidophilus		4.24	0.26			
27373003	**Very good identification**					**27373003**
Actinomyces naeslundii		99.83	0.74	RAF 92		
Actinomyces israelii		0.18	0.18			
27373010	**Low discrimination**					**27373010**
Actinomyces naeslundii		46.23	0.49	RAF 92		
Lactobacillus jensenii		26.59	0.24	MLZ 5		
Actinomyces israelii		22.72	0.35			
27373011	**Good identification**					**27373011**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27373012	**Low discrimination**					**27373012**
Actinomyces naeslundii		42.76	0.49	RAF 92		
Actinomyces israelii		31.52	0.38	MLZ 5		
Lactobacillus jensenii		16.40	0.21			
27373013	**Good identification**					**27373013**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27373020	**Very good identification**					**27373020**
Actinomyces naeslundii		95.15	0.71	RAF 92		
Lactobacillus jensenii		2.16	0.21			
Actinomyces israelii		1.05	0.28			
27373021	**Very good identification**					**27373021**
Actinomyces naeslundii		100.0	0.74	RAF 92		
27373022	**Low discrimination**					**27373022**
					SPO	
Actinomyces naeslundii		87.88	0.71	RAF 92	1	
Lactobacillus acidophilus		5.34	0.26		1	
Clostridium ramosum		1.79	0.17		92	
27373023	**Very good identification**					**27373023**
Actinomyces naeslundii		100.0	0.74	RAF 92		
27373030	**Low discrimination**					**27373030**
					SPO	
Actinomyces naeslundii		73.67	0.49	RAF 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		13.43	0.13	MLZ 5	99	
Lactobacillus jensenii		10.60	0.13		1	
27373031	**Good identification**					**27373031**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27373032	**Low discrimination**					**27373032**
					SPO	
Actinomyces naeslundii		59.18	0.49	RAF 92	1	
Clostridium tertium		32.36	0.21	MLZ 5	99	
Lactobacillus jensenii		5.67	0.10		1	
27373033	**Good identification**					**27373033**
Actinomyces naeslundii		100.0	0.52	RAF 92		
27373040	**Low discrimination**					**27373040**
Actinomyces naeslundii		72.39	0.58	RAF 92		
Actinomyces israelii		27.61	0.42			
27373041	**Very good identification**					**27373041**
Actinomyces naeslundii		100.0	0.61	RAF 92		
27373042	**Low discrimination**					**27373042**
Actinomyces naeslundii		61.93	0.58	RAF 92		
Actinomyces israelii		35.43	0.45			
Lactobacillus acidophilus		2.64	0.10			
27373043	**Very good identification**					**27373043**
Actinomyces naeslundii		100.0	0.61	RAF 92		
27373051	**Good identification**					**27373051**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27373053	**Good identification**					**27373053**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27373060	**Very good identification**					**27373060**
Actinomyces naeslundii		97.62	0.58	RAF 92		
Actinomyces israelii		2.38	0.21			
27373061	**Very good identification**					**27373061**
Actinomyces naeslundii		100.0	0.61	RAF 92		
27373062	**Good identification**					**27373062**
Actinomyces naeslundii		92.67	0.58	RAF 92		
Lactobacillus acidophilus		3.95	0.10			
Actinomyces israelii		3.39	0.24			
27373063	**Very good identification**					**27373063**
Actinomyces naeslundii		100.0	0.61	RAF 92		
27373070	**Good identification**					**27373070**
Actinomyces naeslundii		100.0	0.36	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27373071	**Good identification**					**27373071**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27373072	**Good identification**					**27373072**
Actinomyces naeslundii		100.0	0.36	RAF 92		
27373073	**Good identification**					**27373073**
Actinomyces naeslundii		100.0	0.39	RAF 92		
27373100	**Low discrimination**					**27373100**
					SPO	
Actinomyces naeslundii		75.67	0.79	RAF 92	1	
Actinomyces israelii		18.41	0.59		1	
Clostridium tertium		4.74	0.35		99	
27373101	**Very good identification**					**27373101**
					SPO	
Actinomyces naeslundii		99.79	0.82	RAF 92	1	
Actinomyces israelii		0.16	0.25		1	
Clostridium tertium		0.04	0.01		99	
27373102	**Low discrimination**					**27373102**
Actinomyces naeslundii		55.13	0.79	RAF 92		
Actinomyces israelii		20.12	0.62			
Lactobacillus acidophilus		13.55	0.44			
27373103	**Very good identification**					**27373103**
Actinomyces naeslundii		99.47	0.82	RAF 92		
Actinomyces israelii		0.24	0.28			
Lactobacillus acidophilus		0.17	0.10			
27373110	**Low discrimination**					**27373110**
					SPO	
Clostridium tertium		55.56	0.39	NIT 1	99	
Actinomyces naeslundii		26.29	0.57		1	
Actinomyces israelii		18.16	0.45		1	
27373111	**Very good identification**					**27373111**
					SPO	
Actinomyces naeslundii		98.60	0.60	RAF 92	1	
Clostridium tertium		1.40	0.05	MLZ 5	99	
27373112	**Low discrimination**					**27373112**
					SPO	
Clostridium tertium		72.11	0.47	NIT 1	99	
Actinomyces israelii		11.78	0.48		1	
Actinomyces naeslundii		11.37	0.57		1	
27373113	**Very good identification**					**27373113**
					SPO	
Actinomyces naeslundii		95.27	0.60	RAF 92	1	
Clostridium tertium		4.07	0.13	MLZ 5	99	
Actinomyces israelii		0.67	0.14		1	
27373120	**Low discrimination**					**27373120**
					SPO	
Actinomyces naeslundii		71.53	0.79	RAF 92	1	
Clostridium tertium		13.43	0.43		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		12.66	0.41		92	
27373121	**Very good identification**					**27373121**
					SPO	
Actinomyces naeslundii		99.76	0.82	RAF 92	1	
Clostridium tertium		0.13	0.09		99	
Clostridium ramosum		0.12	0.07		92	
27373122	**Low discrimination**					**27373122**
					SPO	
Actinomyces naeslundii		39.18	0.79	RAF 92	1	
Clostridium ramosum		27.73	0.51		92	
Clostridium tertium		22.06	0.51		99	
27373123	**Very good identification**					**27373123**
					SPO	
Actinomyces naeslundii		98.99	0.82	RAF 92	1	
Clostridium ramosum		0.47	0.17		92	
Clostridium tertium		0.38	0.17		99	
27373130	**Low discrimination**					**27373130**
					SPO	
Clostridium tertium		85.47	0.47	NIT 1	99	
Actinomyces naeslundii		13.48	0.57		1	
Actinomyces israelii		0.59	0.25		1	
27373131	**Very good identification**					**27373131**
					SPO	
Actinomyces naeslundii		95.91	0.60	RAF 92	1	
Clostridium tertium		4.10	0.13	MLZ 5	99	
27373132	**Acceptable identification**					**27373132**
					SPO	
Clostridium tertium		93.65	0.55	NIT 1	99	
Actinomyces naeslundii		4.92	0.57		1	
Clostridium ramosum		0.67	0.17		92	
27373133	**Low discrimination**					**27373133**
					SPO	
Actinomyces naeslundii		88.65	0.60	RAF 92	1	
Clostridium tertium		11.36	0.21	MLZ 5	99	
27373140	**Low discrimination**					**27373140**
Actinomyces naeslundii		64.35	0.66	RAF 92		
Actinomyces israelii		34.50	0.52			
Propionibacterium propionicum		0.92	0.24			
27373141	**Very good identification**					**27373141**
Actinomyces naeslundii		99.64	0.69	RAF 92		
Actinomyces israelii		0.36	0.18			
27373142	**Low discrimination**					**27373142**
Actinomyces naeslundii		49.79	0.66	RAF 92		
Actinomyces israelii		40.04	0.55			
Lactobacillus acidophilus		8.57	0.28			
27373143	**Very good identification**					**27373143**
Actinomyces naeslundii		99.46	0.69	RAF 92		
Actinomyces israelii		0.54	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27373150	**Low discrimination**					**27373150**
					SPO	
Actinomyces israelii		57.59	0.38	XYL 92	1	
Actinomyces naeslundii		37.84	0.44	MLZ 13	1	
Clostridium tertium		4.58	0.05		99	
27373151	**Good identification**					**27373151**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27373152	**Low discrimination**					**27373152**
					SPO	
Actinomyces israelii		62.62	0.41	XYL 92	1	
Actinomyces naeslundii		27.43	0.44	MLZ 13	1	
Clostridium tertium		9.95	0.13		99	
27373153	**Good identification**					**27373153**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27373160	**Good identification**					**27373160**
Actinomyces naeslundii		93.68	0.66	RAF 92		
Actinomyces israelii		3.21	0.32			
Clostridium perfringens		1.16	0.11			
27373161	**Very good identification**					**27373161**
Actinomyces naeslundii		100.0	0.69	RAF 92		
27373162	**Low discrimination**					**27373162**
Actinomyces naeslundii		76.60	0.66	RAF 92		
Lactobacillus acidophilus		13.18	0.28			
Actinomyces israelii		3.93	0.35			
27373163	**Very good identification**					**27373163**
Actinomyces naeslundii		100.0	0.69	RAF 92		
27373170	**Low discrimination**					**27373170**
					SPO	
Actinomyces naeslundii		68.49	0.44	RAF 92	1	
Clostridium tertium		24.86	0.13	MLZ 5	99	
Actinomyces israelii		6.65	0.18		1	
27373171	**Good identification**					**27373171**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27373173	**Good identification**					**27373173**
Actinomyces naeslundii		100.0	0.47	RAF 92		
27373201	**Good identification**					**27373201**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27373203	**Good identification**					**27373203**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27373220	**Low discrimination**					**27373220**
					SPO	
Actinomyces naeslundii		59.97	0.37	RAF 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		40.03	0.09	RHA 1	92	
27373221	**Good identification**					**27373221**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27373223	**Good identification**					**27373223**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27373300	**Low discrimination**					**27373300**
					SPO	
Actinomyces israelii		58.29	0.40	XYL 92	1	
Actinomyces naeslundii		32.16	0.45	RHA 7	1	
Clostridium ramosum		7.54	0.09		92	
27373301	**Good identification**					**27373301**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27373302	**Low discrimination**					**27373302**
Actinomyces israelii		40.73	0.43	XYL 92		
Lactobacillus acidophilus		27.43	0.25	RHA 7		
Actinomyces naeslundii		14.98	0.45			
27373303	**Good identification**					**27373303**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27373320	**Acceptable identification**					**27373320**
					SPO	
Clostridium ramosum		94.69	0.43	NIT 1	92	
Actinomyces naeslundii		4.08	0.45		1	
Clostridium tertium		0.77	0.09		99	
27373321	**Low discrimination**					**27373321**
					SPO	
Actinomyces naeslundii		86.48	0.48	RAF 92	1	
Clostridium ramosum		13.52	0.09	RHA 1	92	
27373322	**Good identification**					**27373322**
					SPO	
Clostridium ramosum		96.24	0.53	NIT 1	92	
Lactobacillus acidophilus		1.90	0.25		1	
Actinomyces naeslundii		1.04	0.45		1	
27373323	**Low discrimination**					**27373323**
					SPO	
Actinomyces naeslundii		61.52	0.48	RAF 92	1	
Clostridium ramosum		38.48	0.19	RHA 1	92	
27373342	**Low discrimination**					**27373342**
Actinomyces israelii		72.42	0.36	XYL 92		
Lactobacillus acidophilus		15.50	0.09	RHA 7		
Actinomyces naeslundii		12.08	0.32			
27373400	**Low discrimination**					**27373400**
					SPO	
Actinomyces israelii		88.69	0.47	XYL 92	1	
Actinomyces naeslundii		6.33	0.37		1	
Clostridium innocuum		4.99	0.11		99	
27373401	**Acceptable identification**					**27373401**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.37	0.40	RAF 92		
Actinomyces israelii		8.63	0.13	ARA 1		
27373402	**Acceptable identification**					**27373402**
					SPO	
Actinomyces israelii		92.16	0.50	XYL 92	1	
Actinomyces naeslundii		4.38	0.37		1	
Clostridium innocuum		3.45	0.11		99	
27373403	**Low discrimination**					**27373403**
Actinomyces naeslundii		87.60	0.40	RAF 92		
Actinomyces israelii		12.40	0.16	ARA 1		
27373412	**Good identification**					**27373412**
Actinomyces israelii		100.0	0.36	XYL 92		
27373420	**Low discrimination**					**27373420**
Actinomyces naeslundii		52.77	0.37	RAF 92		
Actinomyces israelii		47.23	0.27	ARA 1		
27373421	**Good identification**					**27373421**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27373423	**Good identification**					**27373423**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27373440	**Good identification**					**27373440**
Actinomyces israelii		96.87	0.40	XYL 92		
Actinomyces naeslundii		3.14	0.24			
27373442	**Good identification**					**27373442**
Actinomyces israelii		97.89	0.43	XYL 92		
Actinomyces naeslundii		2.11	0.24			
27373500	**Good identification**					**27373500**
					SPO	
Actinomyces israelii		94.88	0.58	XYL 92	1	
Actinomyces naeslundii		4.82	0.45		1	
Clostridium tertium		0.30	0.01		99	
27373501	**Low discrimination**					**27373501**
Actinomyces naeslundii		88.29	0.48	RAF 92		
Actinomyces israelii		11.72	0.24	ARA 1		
27373502	**Good identification**					**27373502**
Actinomyces israelii		94.76	0.61	XYL 92		
Actinomyces naeslundii		3.21	0.45			
Lactobacillus acidophilus		0.79	0.10			
27373503	**Low discrimination**					**27373503**
Actinomyces naeslundii		83.40	0.48	RAF 92		
Actinomyces israelii		16.60	0.27	ARA 1		
27373510	**Acceptable identification**					**27373510**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		94.73	0.44	XYL 92	1	
Clostridium tertium		3.58	0.05	MLZ 13	99	
Actinomyces naeslundii		1.69	0.23		1	
27373512	**Low discrimination**					**27373512**
					SPO	
Actinomyces israelii		87.48	0.47	XYL 92	1	
Clostridium tertium		6.61	0.13	MLZ 13	99	
Bifidobacterium adolescentis		4.87	0.19		1	
27373520	**Low discrimination**					**27373520**
					SPO	
Actinomyces israelii		47.96	0.37	XYL 92	1	
Actinomyces naeslundii		38.13	0.45	NAG 6	1	
Clostridium tertium		7.16	0.09		99	
27373521	**Good identification**					**27373521**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27373522	**Low discrimination**					**27373522**
					SPO	
Actinomyces israelii		42.85	0.40	XYL 92	1	
Actinomyces naeslundii		22.71	0.45	NAG 6	1	
Clostridium ramosum		16.07	0.17		92	
27373523	**Good identification**					**27373523**
Actinomyces naeslundii		100.0	0.48	RAF 92		
27373540	**Good identification**					**27373540**
Actinomyces israelii		97.75	0.51	XYL 92		
Actinomyces naeslundii		2.25	0.32			
27373542	**Good identification**					**27373542**
Actinomyces israelii		98.49	0.54	XYL 92		
Actinomyces naeslundii		1.51	0.32			
27373550	**Good identification**					**27373550**
Actinomyces israelii		100.0	0.37	XYL 92		
27373552	**Good identification**					**27373552**
Actinomyces israelii		100.0	0.40	XYL 92		
27373700	**Good identification**					**27373700**
Actinomyces israelii		100.0	0.38	XYL 92		
27373702	**Good identification**					**27373702**
Actinomyces israelii		100.0	0.41	XYL 92		
27375000	**Low discrimination**					**27375000**
					COC	
Actinomyces israelii		71.64	0.54		1	
Lactobacillus jensenii		18.52	0.32		1	
Atopobium parvulum		8.78	0.25		99	
27375001	**Low discrimination**					**27375001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		68.75	0.35	RAF 92			
Actinomyces israelii		31.25	0.20	XYL 1			
27375002	**Low discrimination**						**27375002**
Actinomyces israelii		85.90	0.57				
Lactobacillus jensenii		9.87	0.29				
Lactobacillus acidophilus		1.90	0.14				
27375003	**Low discrimination**						**27375003**
Actinomyces naeslundii		59.46	0.35	RAF 92			
Actinomyces israelii		40.54	0.23	XYL 1			
27375010	**Low discrimination**						**27375010**
Actinomyces israelii		63.42	0.40	MLZ 13			
Lactobacillus jensenii		36.58	0.24				
27375012	**Low discrimination**						**27375012**
Actinomyces israelii		79.60	0.43	MLZ 13			
Lactobacillus jensenii		20.41	0.21				
27375021	**Good identification**						**27375021**
Actinomyces naeslundii		100.0	0.35	RAF 92			
27375022	**Low discrimination**						**27375022**
Actinomyces israelii		49.64	0.37	NAG 6			
Lactobacillus jensenii		22.34	0.18				
Lactobacillus acidophilus		17.23	0.14				
27375023	**Good identification**						**27375023**
Actinomyces naeslundii		100.0	0.35	RAF 92			
27375040	**Good identification**						**27375040**
Actinomyces israelii		100.0	0.47				
27375042	**Good identification**						**27375042**
Actinomyces israelii		99.30	0.50				
Lactobacillus acidophilus		0.70	-.02				
27375052	**Good identification**						**27375052**
Actinomyces israelii		100.0	0.36	MLZ 13			
27375100	**Low discrimination**						**27375100**
					COC	SPO	
Atopobium parvulum		74.67	0.59	NIT 1	99	1	
Actinomyces israelii		24.62	0.64		1	1	
Clostridium butyricum		0.41	0.13		1	99	
27375102	**Low discrimination**						**27375102**
					COC		
Actinomyces israelii		69.45	0.67		1		
Atopobium parvulum		24.78	0.46		99		
Lactobacillus acidophilus		4.42	0.32		1		
27375103	**Low discrimination**						**27375103**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		43.47	0.43	RAF 92	1		
Actinomyces israelii		41.66	0.33	XYL 1	1		
Atopobium parvulum		14.87	0.12		99		
27375110	**Low discrimination**						**27375110**
					COC	SPO	
Actinomyces israelii		81.56	0.50	MLZ 13	1	1	
Atopobium parvulum		16.72	0.25		99	1	
Clostridium tertium		1.72	0.07		1	99	
27375112	**Low discrimination**						**27375112**
					GRA	SPO	
Actinomyces israelii		78.77	0.53	MLZ 13	99	1	
Leptotrichia buccalis		13.86	0.15		1	1	
Clostridium tertium		3.31	0.15		99	99	
27375120	**Low discrimination**						**27375120**
					COC		
Atopobium parvulum		83.67	0.46	NIT 1	99		
Actinomyces israelii		9.98	0.44		1		
Clostridium perfringens		3.65	0.23		1		
27375121	**Low discrimination**						**27375121**
					COC		
Actinomyces naeslundii		74.52	0.43	RAF 92	1		
Atopobium parvulum		25.48	0.12	XYL 1	99		
27375122	**Low discrimination**						**27375122**
					COC		
Actinomyces israelii		28.99	0.47	NAG 6	1		
Lactobacillus acidophilus		28.92	0.32		1		
Atopobium parvulum		28.59	0.33		99		
27375123	**Good identification**						**27375123**
Actinomyces naeslundii		100.0	0.43	RAF 92			
27375140	**Good identification**						**27375140**
					COC		
Actinomyces israelii		91.98	0.57		1		
Atopobium parvulum		7.25	0.25		99		
Actinomyces naeslundii		0.44	0.27		1		
27375142	**Very good identification**						**27375142**
					COC		
Actinomyces israelii		96.83	0.60		1		
Lactobacillus acidophilus		1.96	0.16		1		
Atopobium parvulum		0.90	0.12		99		
27375150	**Good identification**						**27375150**
Actinomyces israelii		100.0	0.43	MLZ 13			
27375152	**Good identification**						**27375152**
Actinomyces israelii		100.0	0.46	MLZ 13			
27375160	**Low discrimination**						**27375160**
					COC		
Actinomyces israelii		67.95	0.37	NAG 6	1		
Atopobium parvulum		14.80	0.12		99		
Clostridium perfringens		12.16	0.11		1		
27375162	**Low discrimination**						**27375162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		70.26	0.40	NAG 6			
Lactobacillus acidophilus		22.28	0.16				
Clostridium perfringens		3.95	0.05				
27375202	**Good identification**						**27375202**
Actinomyces israelii		100.0	0.38	RHA 7			
27375300	**Low discrimination**						**27375300**
					COC		
Actinomyces israelii		71.07	0.45	RHA 7	1		
Atopobium parvulum		28.93	0.25		99		
27375302	**Low discrimination**						**27375302**
					COC		
Actinomyces israelii		89.14	0.48	RHA 7	1		
Lactobacillus acidophilus		5.68	0.12		1		
Atopobium parvulum		4.27	0.12		99		
27375340	**Good identification**						**27375340**
Actinomyces israelii		100.0	0.38	RHA 7			
27375342	**Good identification**						**27375342**
Actinomyces israelii		100.0	0.41	RHA 7			
27375400	**Good identification**						**27375400**
					SPO		
Actinomyces israelii		97.07	0.53		1		
Clostridium butyricum		2.93	0.06		99		
27375402	**Good identification**						**27375402**
					SPO		
Actinomyces israelii		99.22	0.56		1		
Clostridium butyricum		0.78	-.02		99		
27375410	**Good identification**						**27375410**
Actinomyces israelii		100.0	0.38	MLZ 13			
27375412	**Acceptable identification**						**27375412**
Actinomyces israelii		93.17	0.42	MLZ 13			
Clostridium clostridioforme		6.83	0.13				
27375422	**Good identification**						**27375422**
Actinomyces israelii		100.0	0.35	NAG 6			
27375440	**Good identification**						**27375440**
Actinomyces israelii		100.0	0.46				
27375442	**Good identification**						**27375442**
Actinomyces israelii		100.0	0.49				
27375500	**Low discrimination**						**27375500**
					COC	SPO	
Actinomyces israelii		80.81	0.63		1	1	
Clostridium butyricum		14.64	0.29		1	99	
Atopobium parvulum		3.03	0.25		99	1	
27375502	**Good identification**						**27375502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		92.41	0.66		1		
Clostridium butyricum		4.34	0.22		99		
Bifidobacterium adolescentis		2.53	0.33		1		
27375510	**Acceptable identification**						**27375510**
					GRA	SPO	
Actinomyces israelii		90.41	0.49	MLZ 13	99	1	
Clostridium butyricum		5.77	0.07		99	99	
Mitsuokella multacida		3.83	0.13		1	1	
27375512	**Low discrimination**						**27375512**
Actinomyces israelii		76.80	0.52	MLZ 13			
Bifidobacterium adolescentis		17.22	0.34				
Clostridium clostridioforme		4.71	0.22				
27375520	**Low discrimination**						**27375520**
					COC	SPO	
Actinomyces israelii		70.20	0.42	NAG 6	1	1	
Clostridium butyricum		17.32	0.11		1	99	
Atopobium parvulum		7.27	0.12		99	1	
27375522	**Acceptable identification**						**27375522**
					SPO		
Actinomyces israelii		91.28	0.45	NAG 6	1		
Clostridium butyricum		5.84	0.04		99		
Clostridium clostridioforme		2.88	0.10		66		
27375540	**Acceptable identification**						**27375540**
					GRA	SPO	
Actinomyces israelii		91.80	0.56		99	1	
Mitsuokella multacida		4.48	0.21		1	1	
Clostridium butyricum		3.72	0.11		99	99	
27375542	**Very good identification**						**27375542**
					SPO		
Actinomyces israelii		97.77	0.59		1		
Clostridium butyricum		1.03	0.04		99		
Bifidobacterium adolescentis		0.85	0.17		1		
27375550	**Acceptable identification**						**27375550**
					GRA		
Actinomyces israelii		90.19	0.42	MLZ 13	99		
Mitsuokella multacida		9.81	0.13		1		
27375552	**Acceptable identification**						**27375552**
Actinomyces israelii		93.35	0.45	MLZ 13			
Bifidobacterium adolescentis		6.65	0.19				
27375560	**Low discrimination**						**27375560**
					GRA	SPO	
Actinomyces israelii		80.24	0.35	NAG 6	99	1	
Mitsuokella multacida		15.33	0.11		1	1	
Clostridium butyricum		4.43	-.07		99	99	
27375562	**Good identification**						**27375562**
Actinomyces israelii		100.0	0.38	NAG 6			
27375602	**Low discrimination**						**27375602**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		81.01	0.36	RHA 7			
Clostridium clostridioforme		18.99	0.16				
27375700	**Low discrimination**						**27375700**
					GRA	SPO	
Actinomyces israelii		64.83	0.44	RHA 7	99	1	
Mitsuokella multacida		30.34	0.25		1	1	
Clostridium butyricum		4.83	0.03		99	99	
27375702	**Low discrimination**						**27375702**
					GRA		
Actinomyces israelii		79.99	0.47	RHA 7	99		
Clostridium clostridioforme		15.69	0.25		50		
Mitsuokella multacida		2.77	0.09		1		
27375712	**Low discrimination**						**27375712**
					GRA		
Clostridium clostridioforme		84.73	0.36	NIT 1	50		
Actinomyces israelii		14.17	0.33	RAF 94	99		
Mitsuokella multacida		1.10	0.01		1		
27375732	**Good identification**						**27375732**
Clostridium clostridioforme		98.15	0.36	NIT 1			
Parabacteroides distasonis		1.85	0.13	RAF 94			
27375742	**Acceptable identification**						**27375742**
					GRA		
Actinomyces israelii		91.82	0.40	RHA 7	99		
Mitsuokella multacida		8.18	0.09		1		
27376002	**Good identification**						**27376002**
Actinomyces israelii		100.0	0.36	GAL 99			
27376100	**Good identification**						**27376100**
					SPO		
Actinomyces israelii		97.00	0.43	GAL 99	1		
Clostridium tertium		3.00	0.03		99		
27376102	**Acceptable identification**						**27376102**
					SPO		
Actinomyces israelii		94.18	0.46	GAL 99	1		
Clostridium tertium		5.82	0.11		99		
27376140	**Good identification**						**27376140**
Actinomyces israelii		100.0	0.36	GAL 99			
27376142	**Good identification**						**27376142**
Actinomyces israelii		100.0	0.39	GAL 99			
27376500	**Good identification**						**27376500**
Actinomyces israelii		100.0	0.42	GAL 99			
27376502	**Good identification**						**27376502**
Actinomyces israelii		100.0	0.45	GAL 99			
27376542	**Good identification**						**27376542**
Actinomyces israelii		100.0	0.38	GAL 99			
27377000	**Very good identification**						**27377000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		99.25	0.67		1	
Actinomyces naeslundii		0.50	0.37		1	
Clostridium tertium		0.12	0.03		99	
27377002	**Very good identification**					**27377002**
					SPO	
Actinomyces israelii		99.34	0.70		1	
Actinomyces naeslundii		0.34	0.37		1	
Clostridium tertium		0.25	0.11		99	
27377003	**Low discrimination**					**27377003**
Actinomyces israelii		66.56	0.36	URE 1		
Actinomyces naeslundii		33.44	0.40			
27377010	**Good identification**					**27377010**
					SPO	
Actinomyces israelii		98.54	0.53	MLZ 13	1	
Clostridium tertium		1.46	0.07		99	
27377012	**Good identification**					**27377012**
					SPO	
Actinomyces israelii		97.12	0.56	MLZ 13	1	
Clostridium tertium		2.88	0.15		99	
27377020	**Low discrimination**					**27377020**
					SPO	
Actinomyces israelii		87.86	0.47	NAG 6	1	
Actinomyces naeslundii		6.99	0.37		1	
Clostridium tertium		5.15	0.11		99	
27377021	**Good identification**					**27377021**
Actinomyces naeslundii		100.0	0.40	RAF 92		
27377022	**Low discrimination**					**27377022**
					SPO	
Actinomyces israelii		85.45	0.50	NAG 6	1	
Clostridium tertium		10.02	0.19		99	
Actinomyces naeslundii		4.53	0.37		1	
27377023	**Low discrimination**					**27377023**
Actinomyces naeslundii		88.73	0.40	RAF 92		
Actinomyces israelii		11.27	0.15	XYL 1		
27377040	**Very good identification**					**27377040**
Actinomyces israelii		99.77	0.60			
Actinomyces naeslundii		0.23	0.24			
27377042	**Very good identification**					**27377042**
Actinomyces israelii		99.85	0.63			
Actinomyces naeslundii		0.15	0.24			
27377050	**Good identification**					**27377050**
Actinomyces israelii		100.0	0.46	MLZ 13		
27377052	**Good identification**					**27377052**
Actinomyces israelii		100.0	0.49	MLZ 13		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
27377060	**Good identification**						**27377060**
Actinomyces israelii		96.52	0.40	NAG 6			
Actinomyces naeslundii		3.48	0.24				
27377062	**Good identification**						**27377062**
Actinomyces israelii		97.65	0.43	NAG 6			
Actinomyces naeslundii		2.35	0.24				
27377100	**Very good identification**						**27377100**
					COC	SPO	
Actinomyces israelii		96.13	0.77		1	1	
Clostridium tertium		2.97	0.37		1	99	
Atopobium parvulum		0.52	0.25		99	1	
27377101	**Low discrimination**						**27377101**
					SPO		
Actinomyces israelii		63.81	0.43	URE 1	1		
Actinomyces naeslundii		34.22	0.48		1		
Clostridium tertium		1.97	0.03		99		
27377102	**Good identification**						**27377102**
					SPO		
Actinomyces israelii		93.82	0.80		1		
Clostridium tertium		5.80	0.45		99		
Actinomyces naeslundii		0.23	0.45		1		
27377103	**Low discrimination**						**27377103**
					SPO		
Actinomyces israelii		70.46	0.46	URE 1	1		
Actinomyces naeslundii		25.19	0.48		1		
Clostridium tertium		4.35	0.11		99		
27377110	**Low discrimination**						**27377110**
					SPO		
Actinomyces israelii		73.06	0.63	MLZ 13	1		
Clostridium tertium		26.85	0.41		99		
Actinomyces naeslundii		0.09	0.23		1		
27377112	**Low discrimination**						**27377112**
					SPO		
Actinomyces israelii		57.48	0.66	MLZ 13	1		
Clostridium tertium		42.25	0.50		99		
Bifidobacterium adolescentis		0.23	0.19		1		
27377120	**Low discrimination**						**27377120**
					SPO		
Clostridium tertium		57.31	0.45	NIT 1	99		
Actinomyces israelii		39.48	0.57		1		
Actinomyces naeslundii		2.23	0.45		1		
27377121	**Low discrimination**						**27377121**
					SPO		
Actinomyces naeslundii		77.41	0.48	RAF 92	1		
Clostridium tertium		13.38	0.11	XYL 1	99		
Actinomyces israelii		9.22	0.23		1		
27377122	**Low discrimination**						**27377122**
					SPO		
Clostridium tertium		73.00	0.53	NIT 1	99		
Actinomyces israelii		25.14	0.60		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.95	0.45		1	
27377123	**Low discrimination**					**27377123**
					SPO	
Actinomyces naeslundii		58.93	0.48	RAF 92	1	
Clostridium tertium		30.55	0.19	XYL 1	99	
Actinomyces israelii		10.52	0.26		1	
27377130	**Acceptable identification**					**27377130**
					SPO	
Clostridium tertium		94.42	0.50	NIT 1	99	
Actinomyces israelii		5.47	0.43		1	
Actinomyces naeslundii		0.11	0.23		1	
27377132	**Very good identification**					**27377132**
					SPO	
Clostridium tertium		97.15	0.58	NIT 1	99	
Actinomyces israelii		2.81	0.46		1	
Actinomyces naeslundii		0.04	0.23		1	
27377140	**Very good identification**					**27377140**
					SPO	
Actinomyces israelii		99.76	0.70		1	
Actinomyces naeslundii		0.16	0.32		1	
Clostridium tertium		0.08	0.03		99	
27377141	**Low discrimination**					**27377141**
Actinomyces israelii		80.43	0.36	URE 1		
Actinomyces naeslundii		19.57	0.35			
27377142	**Very good identification**					**27377142**
					SPO	
Actinomyces israelii		99.73	0.73		1	
Clostridium tertium		0.16	0.11		99	
Actinomyces naeslundii		0.11	0.32		1	
27377143	**Low discrimination**					**27377143**
Actinomyces israelii		86.04	0.39	URE 1		
Actinomyces naeslundii		13.96	0.35			
27377150	**Very good identification**					**27377150**
					SPO	
Actinomyces israelii		99.05	0.56	MLZ 13	1	
Clostridium tertium		0.95	0.07		99	
27377152	**Very good identification**					**27377152**
					SPO	
Actinomyces israelii		98.13	0.59	MLZ 13	1	
Clostridium tertium		1.87	0.15		99	
27377160	**Acceptable identification**					**27377160**
					SPO	
Actinomyces israelii		94.04	0.50	NAG 6	1	
Clostridium tertium		3.55	0.11		99	
Actinomyces naeslundii		2.41	0.32		1	
27377162	**Acceptable identification**					**27377162**
					SPO	
Actinomyces israelii		91.53	0.53	NAG 6	1	
Clostridium tertium		6.90	0.19		99	
Actinomyces naeslundii		1.57	0.32		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27377170	**Low discrimination**					**27377170**
					SPO	
Actinomyces israelii		69.03	0.36	MLZ 13	1	
Clostridium tertium		30.97	0.15	NAG 6	99	
27377172	**Low discrimination**					**27377172**
					SPO	
Actinomyces israelii		52.71	0.39	MLZ 13	1	
Clostridium tertium		47.29	0.24	NAG 6	99	
27377200	**Good identification**					**27377200**
Actinomyces israelii		100.0	0.48	RHA 7		
27377202	**Good identification**					**27377202**
Actinomyces israelii		100.0	0.51	RHA 7		
27377212	**Good identification**					**27377212**
Actinomyces israelii		100.0	0.37	RHA 7		
27377240	**Good identification**					**27377240**
Actinomyces israelii		100.0	0.41	RHA 7		
27377242	**Good identification**					**27377242**
Actinomyces israelii		100.0	0.44	RHA 7		
27377300	**Very good identification**					**27377300**
					SPO	
Actinomyces israelii		99.59	0.58	RHA 7	1	
Clostridium tertium		0.41	0.03		99	
27377302	**Very good identification**					**27377302**
					SPO	
Actinomyces israelii		99.18	0.61	RHA 7	1	
Clostridium tertium		0.82	0.11		99	
27377310	**Good identification**					**27377310**
					SPO	
Actinomyces israelii		95.30	0.44	RHA 7	1	
Clostridium tertium		4.70	0.07	MLZ 13	99	
27377312	**Acceptable identification**					**27377312**
					SPO	
Actinomyces israelii		91.02	0.47	RHA 7	1	
Clostridium tertium		8.98	0.15	MLZ 13	99	
27377320	**Low discrimination**					**27377320**
					SPO	
Actinomyces israelii		72.94	0.38	RHA 7	1	
Clostridium tertium		14.21	0.11	NAG 6	99	
Clostridium ramosum		12.86	0.09		92	
27377322	**Low discrimination**					**27377322**
					SPO	
Actinomyces israelii		53.77	0.41	RHA 7	1	
Clostridium ramosum		25.28	0.19	NAG 6	92	
Clostridium tertium		20.95	0.19		99	
27377340	**Good identification**					**27377340**
Actinomyces israelii		100.0	0.51	RHA 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27377342	**Good identification**						**27377342**
Actinomyces israelii		100.0	0.54	RHA 7			
27377350	**Good identification**						**27377350**
Actinomyces israelii		100.0	0.37	RHA 7			
27377352	**Good identification**						**27377352**
Actinomyces israelii		100.0	0.40	RHA 7			
27377400	**Very good identification**						**27377400**
Actinomyces israelii		100.0	0.66				
27377402	**Very good identification**						**27377402**
Actinomyces israelii		100.0	0.69				
27377410	**Good identification**						**27377410**
Actinomyces israelii		100.0	0.51	MLZ 13			
27377412	**Good identification**						**27377412**
Actinomyces israelii		100.0	0.54	MLZ 13			
27377420	**Good identification**						**27377420**
Actinomyces israelii		100.0	0.45	NAG 6			
27377422	**Good identification**						**27377422**
Actinomyces israelii		100.0	0.48	NAG 6			
27377440	**Very good identification**						**27377440**
Actinomyces israelii		100.0	0.59				
27377442	**Very good identification**						**27377442**
Actinomyces israelii		100.0	0.62				
27377450	**Good identification**						**27377450**
Actinomyces israelii		100.0	0.44	MLZ 13			
27377452	**Good identification**						**27377452**
Actinomyces israelii		100.0	0.47	MLZ 13			
27377460	**Good identification**						**27377460**
Actinomyces israelii		100.0	0.38	NAG 6			
27377462	**Good identification**						**27377462**
Actinomyces israelii		100.0	0.41	NAG 6			
27377500	**Very good identification**						**27377500**
					GRA	SPO	
Actinomyces israelii		98.84	0.76		99	1	
Mitsuokella multacida		0.77	0.27		1	1	
Clostridium butyricum		0.35	0.13		99	99	
27377501	**Acceptable identification**						**27377501**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		92.09	0.42	URE 1	99		
Mitsuokella multacida		7.91	0.11		1		
27377502	**Very good identification**						**27377502**
					SPO		
Actinomyces israelii		99.38	0.79		1		
Bifidobacterium adolescentis		0.39	0.31		1		
Clostridium butyricum		0.09	0.06		99		
27377503	**Good identification**						**27377503**
Actinomyces israelii		100.0	0.45	URE 1			
27377510	**Very good identification**						**27377510**
					GRA	SPO	
Actinomyces israelii		97.85	0.62	MLZ 13	99	1	
Mitsuokella multacida		1.71	0.19		1	1	
Clostridium tertium		0.44	0.07		99	99	
27377512	**Very good identification**						**27377512**
Actinomyces israelii		95.06	0.65	MLZ 13			
Bifidobacterium adolescentis		3.08	0.33				
Bifidobacterium dentium		0.88	0.06				
27377520	**Acceptable identification**						**27377520**
					GRA	SPO	
Actinomyces israelii		94.94	0.55	NAG 6	99	1	
Mitsuokella multacida		2.90	0.17		1	1	
Clostridium tertium		1.70	0.11		99	99	
27377522	**Very good identification**						**27377522**
					SPO		
Actinomyces israelii		96.54	0.58	NAG 6	1		
Clostridium tertium		3.46	0.19		99		
27377530	**Low discrimination**						**27377530**
					GRA	SPO	
Actinomyces israelii		75.79	0.41	MLZ 13	99	1	
Clostridium tertium		16.16	0.15	NAG 6	99	99	
Mitsuokella multacida		5.17	0.09		1	1	
27377532	**Low discrimination**						**27377532**
					SPO		
Actinomyces israelii		59.42	0.44	MLZ 13	1		
Clostridium tertium		25.34	0.24	NAG 6	99		
Bifidobacterium dentium		15.24	0.10		1		
27377540	**Very good identification**						**27377540**
					GRA	SPO	
Actinomyces israelii		97.96	0.69		99	1	
Mitsuokella multacida		1.97	0.27		1	1	
Clostridium butyricum		0.08	-.05		99	99	
27377542	**Very good identification**						**27377542**
					GRA		
Actinomyces israelii		99.73	0.72		99		
Mitsuokella multacida		0.15	0.11		1		
Bifidobacterium adolescentis		0.13	0.16		99		
27377543	**Good identification**						**27377543**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.38	URE 1			
27377550	**Good identification**						**27377550**
					COC	GRA	
Actinomyces israelii		95.10	0.55	MLZ 13	1	99	
Mitsuokella multacida		4.26	0.19		1	1	
Blautia producta		0.64	-.03		99	99	
27377552	**Very good identification**						**27377552**
					GRA		
Actinomyces israelii		98.66	0.58	MLZ 13	99		
Bifidobacterium adolescentis		1.02	0.17		99		
Mitsuokella multacida		0.33	0.03		1		
27377560	**Acceptable identification**						**27377560**
					GRA		
Actinomyces israelii		92.71	0.48	NAG 6	99		
Mitsuokella multacida		7.29	0.17		1		
27377562	**Good identification**						**27377562**
Actinomyces israelii		100.0	0.51	NAG 6			
27377572	**Good identification**						**27377572**
Actinomyces israelii		100.0	0.37	MLZ 13			
27377600	**Good identification**						**27377600**
Actinomyces israelii		100.0	0.46	RHA 7			
27377602	**Good identification**						**27377602**
Actinomyces israelii		100.0	0.49	RHA 7			
27377612	**Good identification**						**27377612**
Actinomyces israelii		100.0	0.35	RHA 7			
27377640	**Good identification**						**27377640**
Actinomyces israelii		100.0	0.39	RHA 7			
27377642	**Good identification**						**27377642**
Actinomyces israelii		100.0	0.42	RHA 7			
27377700	**Low discrimination**						**27377700**
					GRA		
Actinomyces israelii		83.84	0.57	RHA 7	99		
Mitsuokella multacida		16.16	0.32		1		
27377702	**Very good identification**						**27377702**
					GRA		
Actinomyces israelii		98.59	0.60	RHA 7	99		
Mitsuokella multacida		1.41	0.15		1		
27377710	**Low discrimination**						**27377710**
					GRA		
Actinomyces israelii		69.94	0.42	RHA 7	99		
Mitsuokella multacida		30.06	0.24	MLZ 13	1		
27377712	**Good identification**						**27377712**
					GRA		
Actinomyces israelii		96.91	0.45	RHA 7	99		
Mitsuokella multacida		3.09	0.07	MLZ 13	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27377720	**Low discrimination**					**27377720**
					GRA	
Actinomyces israelii		56.99	0.36	RHA 7	99	
Mitsuokella multacida		43.01	0.21	NAG 6	1	
27377722	**Acceptable identification**					**27377722**
					GRA	
Actinomyces israelii		94.71	0.39	RHA 7	99	
Mitsuokella multacida		5.29	0.05	NAG 6	1	
27377740	**Low discrimination**					**27377740**
					GRA	
Actinomyces israelii		66.87	0.50	RHA 7	99	
Mitsuokella multacida		33.13	0.32		1	
27377742	**Good identification**					**27377742**
					GRA	
Actinomyces israelii		96.46	0.53	RHA 7	99	
Mitsuokella multacida		3.54	0.15		1	
27377752	**Acceptable identification**					**27377752**
					GRA	
Actinomyces israelii		92.43	0.38	RHA 7	99	
Mitsuokella multacida		7.57	0.07	MLZ 13	1	
27400002	**Good identification**					**27400002**
Actinomyces viscosus		99.19	0.40	MLT 99		
Propionibacterium granulosum		0.81	0.13	FRU 95		
27400102	**Good identification**					**27400102**
Actinomyces viscosus		100.0	0.44	MLT 99		
27400103	**Good identification**					**27400103**
Actinomyces viscosus		100.0	0.36	MLT 99		
27401002	**Good identification**					**27401002**
Actinomyces viscosus		97.78	0.36	MLT 99		
Propionibacterium granulosum		1.30	0.13	FRU 95		
Actinomyces naeslundii		0.92	0.26			
27401102	**Good identification**					**27401102**
Actinomyces viscosus		98.01	0.40	MLT 99		
Actinomyces naeslundii		1.48	0.33	FRU 95		
Lactobacillus fermentum		0.51	0.15			
27401123	**Low discrimination**					**27401123**
					SPO	
Actinomyces naeslundii		83.29	0.36	MLT 94	1	
Clostridium chauvoei		16.71	0.01	FRU 95	99	
27403100	**Low discrimination**					**27403100**
Actinomyces naeslundii		67.21	0.38	MLT 94		
Propionibacterium propionicum		32.79	0.23	FRU 95		
27403101	**Good identification**					**27403101**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27403102	**Low discrimination**					**27403102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		48.14	0.38	MLT 94			
Propionibacterium propionicum		35.23	0.26	FRU 95			
Actinomyces viscosus		16.64	0.06				
27403103	**Good identification**						**27403103**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27403120	**Good identification**						**27403120**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27403121	**Good identification**						**27403121**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27403122	**Good identification**						**27403122**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27403123	**Good identification**						**27403123**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27410000	**Low discrimination**						**27410000**
					GRA	CAT	
Propionibacterium granulosum		78.30	0.58	NIT 1	99	90	
Fusobacterium mortiferum		14.85	0.49		1	5	
Actinomyces viscosus		4.66	0.27		99	76	
27410002	**Very good identification**						**27410002**
					GRA	CAT	
Actinomyces viscosus		95.55	0.62	MLT 99	99	76	
Propionibacterium granulosum		4.05	0.47		99	90	
Fusobacterium mortiferum		0.34	0.32		1	5	
27410003	**Good identification**						**27410003**
Actinomyces viscosus		99.76	0.53	MLT 99			
Propionibacterium granulosum		0.13	0.13				
Actinomyces naeslundii		0.11	0.27				
27410010	**Good identification**						**27410010**
					GRA	CAT	
Propionibacterium granulosum		98.62	0.43	NIT 1	99	90	
Fusobacterium mortiferum		1.39	0.14	MLZ 12	1	5	
27410020	**Acceptable identification**						**27410020**
					GRA	CAT	
Propionibacterium granulosum		92.34	0.47	NIT 1	99	90	
Propionibacterium acnes		5.87	0.18		99	95	
Fusobacterium mortiferum		0.71	0.14		1	5	
27410022	**Low discrimination**						**27410022**
Propionibacterium granulosum		50.62	0.37	NIT 1			
Actinomyces viscosus		48.23	0.27				
Actinomyces naeslundii		1.15	0.24				
27410100	**Low discrimination**						**27410100**
					GRA	CAT	
Fusobacterium mortiferum		77.76	0.55	NIT 1	1	5	
Actinomyces viscosus		18.61	0.32		99	76	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		1.78	0.24		99	90	
27410102	**Very good identification**						**27410102**
					GRA		
Actinomyces viscosus		99.39	0.66	MLT 99	99		
Fusobacterium mortiferum		0.47	0.39		1		
Propionibacterium propionicum		0.07	0.26		99		
27410103	**Very good identification**						**27410103**
Actinomyces viscosus		99.83	0.58	MLT 99			
Actinomyces naeslundii		0.17	0.35				
27411000	**Low discrimination**						**27411000**
					GRA	CAT	
Propionibacterium granulosum		67.52	0.58	NIT 1	99	90	
Fusobacterium mortiferum		19.20	0.52		1	5	
Lactobacillus fermentum		7.23	0.46		99	1	
27411001	**Low discrimination**						**27411001**
Actinomyces naeslundii		66.97	0.50	MLT 94			
Actinomyces viscosus		15.98	0.16				
Propionibacterium granulosum		13.27	0.24				
27411002	**Low discrimination**						**27411002**
					CAT		
Actinomyces viscosus		89.50	0.58	MLT 99	76		
Propionibacterium granulosum		6.19	0.47		90		
Lactobacillus fermentum		2.65	0.46		1		
27411003	**Good identification**						**27411003**
Actinomyces viscosus		95.75	0.50	MLT 99			
Actinomyces naeslundii		4.05	0.50				
Propionibacterium granulosum		0.20	0.13				
27411010	**Good identification**						**27411010**
					GRA	CAT	
Propionibacterium granulosum		96.69	0.43	NIT 1	99	90	
Fusobacterium mortiferum		2.04	0.17	MLZ 12	1	5	
Actinomyces naeslundii		1.27	0.26		99	25	
27411020	**Low discrimination**						**27411020**
					CAT		
Propionibacterium granulosum		66.38	0.47	NIT 1	90		
Clostridium perfringens		9.58	0.26		1		
Lactobacillus fermentum		9.48	0.38		1		
27411021	**Good identification**						**27411021**
Actinomyces naeslundii		95.28	0.50	MLT 94			
Propionibacterium granulosum		4.72	0.13				
27411022	**Low discrimination**						**27411022**
					CAT		
Propionibacterium granulosum		33.44	0.37	NIT 1	90		
Actinomyces viscosus		19.53	0.24		76		
Lactobacillus fermentum		19.10	0.38		1		
27411023	**Low discrimination**						**27411023**
Actinomyces naeslundii		80.73	0.50	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		19.27	0.16			
27411041	**Good identification**					**27411041**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27411061	**Good identification**					**27411061**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27411063	**Good identification**					**27411063**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27411100	**Low discrimination**					**27411100**
					GRA	
Fusobacterium mortiferum		64.64	0.58	NIT 1	1	
Lactobacillus fermentum		17.02	0.50		99	
Actinomyces naeslundii		9.42	0.55		99	
27411101	**Low discrimination**					**27411101**
					SPO	
Actinomyces naeslundii		61.49	0.58	MLT 94	1	
Clostridium chauvoei		25.76	0.29		99	
Actinomyces viscosus		9.17	0.20		1	
27411102	**Good identification**					**27411102**
Actinomyces viscosus		94.52	0.62	MLT 99		
Lactobacillus fermentum		2.57	0.50			
Actinomyces naeslundii		1.42	0.55			
27411103	**Acceptable identification**					**27411103**
					SPO	
Actinomyces viscosus		93.15	0.54	MLT 99	1	
Actinomyces naeslundii		6.31	0.58		1	
Clostridium chauvoei		0.47	0.16		99	
27411111	**Good identification**					**27411111**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27411120	**Low discrimination**					**27411120**
					GRA	
Clostridium perfringens		40.63	0.40	MLT 99	99	
Prevotella melaninogenica		32.02	0.32		1	
Actinomyces naeslundii		14.85	0.55		99	
27411121	**Low discrimination**					**27411121**
					SPO	
Actinomyces naeslundii		85.28	0.58	MLT 94	1	
Clostridium chauvoei		11.91	0.21		99	
Clostridium perfringens		1.57	0.06		84	
27411123	**Low discrimination**					**27411123**
					SPO	
Actinomyces naeslundii		83.96	0.58	MLT 94	1	
Actinomyces viscosus		12.52	0.20		1	
Clostridium chauvoei		2.07	0.08		99	
27411131	**Good identification**					**27411131**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27411133	**Good identification**					**27411133**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.36	MLT 94			
27411140	**Low discrimination**						**27411140**
					GRA		
Actinomyces naeslundii		53.48	0.42	MLT 94	99		
Fusobacterium mortiferum		21.00	0.24		1		
Propionibacterium propionicum		15.10	0.23		99		
27411141	**Good identification**						**27411141**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27411143	**Low discrimination**						**27411143**
Actinomyces naeslundii		54.21	0.45	MLT 94			
Actinomyces viscosus		45.79	0.20				
27411161	**Good identification**						**27411161**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27411163	**Good identification**						**27411163**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27411400	**Low discrimination**						**27411400**
					GRA	CAT	
Lactobacillus fermentum		89.20	0.46	NIT 1	99	1	
Propionibacterium granulosum		8.41	0.24		99	90	
Fusobacterium mortiferum		2.39	0.17		1	5	
27411402	**Low discrimination**						**27411402**
					CAT		
Lactobacillus fermentum		73.30	0.46	NIT 1	1		
Actinomyces viscosus		24.98	0.24		76		
Propionibacterium granulosum		1.73	0.13		90		
27411420	**Acceptable identification**						**27411420**
					CAT		
Lactobacillus fermentum		93.39	0.38	NIT 1	1		
Propionibacterium granulosum		6.61	0.13		90		
27411422	**Good identification**						**27411422**
Lactobacillus fermentum		100.0	0.38	NIT 1			
27411500	**Good identification**						**27411500**
					GRA		
Lactobacillus fermentum		95.25	0.50	NIT 1	99		
Fusobacterium mortiferum		3.65	0.24		1		
Bifidobacterium longum subsp. longum		1.10	0.09		99		
27411502	**Low discrimination**						**27411502**
					GRA	CAT	
Lactobacillus fermentum		71.59	0.50	NIT 1	99	1	
Actinomyces viscosus		26.58	0.28		99	76	
Capnocytophaga ochracea		1.01	0.02		1	90	
27411520	**Low discrimination**						**27411520**
Lactobacillus fermentum		79.23	0.41	NIT 1			
Bifidobacterium longum subsp. longum		10.93	0.20				
Clostridium perfringens		3.63	0.06				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27411522	**Low discrimination**						**27411522**
					GRA	CAT	
Lactobacillus fermentum		64.05	0.41	NIT 1	99	1	
Capnocytophaga ochracea		24.31	0.19		1	90	
Bifidobacterium longum subsp. longum		8.84	0.20		99	1	
27412000	**Good identification**						**27412000**
					CAT		
Propionibacterium granulosum		98.58	0.56	NIT 1	90		
Propionibacterium propionicum		0.63	0.23		1		
Actinomyces naeslundii		0.36	0.29		25		
27412002	**Low discrimination**						**27412002**
					CAT		
Propionibacterium granulosum		73.06	0.45	NIT 1	90		
Actinomyces viscosus		23.07	0.27		76		
Propionibacterium propionicum		2.80	0.26		1		
27412010	**Good identification**						**27412010**
Propionibacterium granulosum		100.0	0.41	NIT 1			
27412020	**Acceptable identification**						**27412020**
Propionibacterium granulosum		94.16	0.45	NIT 1			
Propionibacterium acnes		4.46	0.13				
Actinomyces naeslundii		1.37	0.29				
27412022	**Acceptable identification**						**27412022**
Propionibacterium granulosum		94.49	0.35	NIT 1			
Actinomyces naeslundii		5.51	0.29				
27412100	**Very good identification**						**27412100**
					CAT		
Propionibacterium propionicum		96.00	0.57	MLT 99	1		
Actinomyces naeslundii		1.57	0.37		25		
Propionibacterium granulosum		1.53	0.21		90		
27412101	**Low discrimination**						**27412101**
Propionibacterium propionicum		81.88	0.41	MLT 99			
Actinomyces naeslundii		18.12	0.40	URE 10			
27412102	**Low discrimination**						**27412102**
Propionibacterium propionicum		86.10	0.60	MLT 99			
Actinomyces viscosus		12.73	0.32				
Actinomyces naeslundii		0.94	0.37				
27412103	**Low discrimination**						**27412103**
Propionibacterium propionicum		62.85	0.44	MLT 99			
Actinomyces viscosus		27.88	0.24	URE 10			
Actinomyces naeslundii		9.27	0.40				
27412120	**Low discrimination**						**27412120**
					CAT		
Actinomyces naeslundii		53.77	0.37	MLT 94	25		
Propionibacterium propionicum		33.13	0.23	GAL 96	1		
Propionibacterium granulosum		13.09	0.11		90		
27412121	**Good identification**						**27412121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MLT 94		
27412122	**Low discrimination**					**27412122**
Actinomyces naeslundii		51.97	0.37	MLT 94		
Propionibacterium propionicum		48.03	0.26	GAL 96		
27412123	**Good identification**					**27412123**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27412140	**Very good identification**					**27412140**
Propionibacterium propionicum		99.71	0.57	MLT 99		
Actinomyces naeslundii		0.29	0.24			
27412141	**Good identification**					**27412141**
Propionibacterium propionicum		96.24	0.41	MLT 99		
Actinomyces naeslundii		3.76	0.27	URE 10		
27412142	**Very good identification**					**27412142**
Propionibacterium propionicum		99.81	0.60	MLT 99		
Actinomyces naeslundii		0.19	0.24			
27412143	**Good identification**					**27412143**
Propionibacterium propionicum		97.46	0.44	MLT 99		
Actinomyces naeslundii		2.54	0.27	URE 10		
27413000	**Acceptable identification**					**27413000**
					CAT	
Propionibacterium granulosum		91.10	0.56	NIT 1	90	
Actinomyces naeslundii		7.97	0.52		25	
Propionibacterium propionicum		0.58	0.23		1	
27413001	**Acceptable identification**					**27413001**
Actinomyces naeslundii		92.85	0.55	MLT 94		
Propionibacterium granulosum		7.15	0.21			
27413002	**Low discrimination**					**27413002**
Propionibacterium granulosum		63.23	0.45	NIT 1		
Actinomyces naeslundii		22.12	0.52			
Actinomyces viscosus		12.24	0.24			
27413003	**Low discrimination**					**27413003**
Actinomyces naeslundii		87.55	0.55	MLT 94		
Actinomyces viscosus		10.76	0.16			
Propionibacterium granulosum		1.69	0.11			
27413010	**Good identification**					**27413010**
Propionibacterium granulosum		96.74	0.41	NIT 1		
Actinomyces naeslundii		3.27	0.30	MLZ 12		
27413020	**Low discrimination**					**27413020**
Propionibacterium granulosum		71.57	0.45	NIT 1		
Actinomyces naeslundii		25.04	0.52			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium acnes		3.39	0.13			
27413021	**Good identification**					**27413021**
Actinomyces naeslundii		98.11	0.55	MLT 94		
Propionibacterium granulosum		1.89	0.11			
27413022	**Low discrimination**					**27413022**
Actinomyces naeslundii		58.32	0.52	MLT 94		
Propionibacterium granulosum		41.68	0.35			
27413023	**Good identification**					**27413023**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27413040	**Low discrimination**					**27413040**
					CAT	
Actinomyces naeslundii		48.34	0.39	MLT 94	25	
Propionibacterium granulosum		31.64	0.21		90	
Propionibacterium propionicum		20.02	0.23		1	
27413041	**Good identification**					**27413041**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27413042	**Low discrimination**					**27413042**
					CAT	
Actinomyces naeslundii		56.03	0.39	MLT 94	25	
Propionibacterium propionicum		34.80	0.26		1	
Propionibacterium granulosum		9.17	0.11		90	
27413043	**Good identification**					**27413043**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27413060	**Low discrimination**					**27413060**
Actinomyces naeslundii		75.06	0.39	MLT 94		
Propionibacterium acnes		12.65	0.02			
Propionibacterium granulosum		12.28	0.11			
27413061	**Good identification**					**27413061**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27413062	**Good identification**					**27413062**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27413063	**Good identification**					**27413063**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27413100	**Low discrimination**					**27413100**
					CAT	
Propionibacterium propionicum		70.07	0.57	MLT 99	1	
Actinomyces naeslundii		27.57	0.60		25	
Propionibacterium granulosum		1.12	0.21		90	
27413101	**Low discrimination**					**27413101**
Actinomyces naeslundii		84.16	0.63	MLT 94		
Propionibacterium propionicum		15.85	0.41			
27413102	**Low discrimination**					**27413102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		73.64	0.60	MLT 99		
Actinomyces naeslundii		19.31	0.60			
Actinomyces viscosus		6.67	0.28			
27413103	**Low discrimination**					**27413103**
Actinomyces naeslundii		73.57	0.63	MLT 94		
Propionibacterium propionicum		20.78	0.44			
Actinomyces viscosus		5.65	0.20			
27413110	**Low discrimination**					**27413110**
Actinomyces naeslundii		67.21	0.38	MLT 94		
Propionibacterium propionicum		32.79	0.23	MLZ 5		
27413111	**Good identification**					**27413111**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27413112	**Low discrimination**					**27413112**
Actinomyces naeslundii		57.75	0.38	MLT 94		
Propionibacterium propionicum		42.25	0.26	MLZ 5		
27413113	**Good identification**					**27413113**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27413120	**Good identification**					**27413120**
Actinomyces naeslundii		94.23	0.60	MLT 94		
Propionibacterium propionicum		2.42	0.23			
Clostridium perfringens		1.34	0.06			
27413121	**Very good identification**					**27413121**
Actinomyces naeslundii		100.0	0.63	MLT 94		
27413122	**Very good identification**					**27413122**
Actinomyces naeslundii		95.06	0.60	MLT 94		
Propionibacterium propionicum		3.66	0.26			
Clostridium perfringens		0.64	0.01			
27413123	**Very good identification**					**27413123**
Actinomyces naeslundii		100.0	0.63	MLT 94		
27413130	**Good identification**					**27413130**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27413131	**Good identification**					**27413131**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27413132	**Good identification**					**27413132**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27413133	**Good identification**					**27413133**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27413140	**Good identification**					**27413140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		93.51	0.57	MLT 99			
Actinomyces naeslundii		6.49	0.47				
27413141	**Low discrimination**						**27413141**
Propionibacterium propionicum		51.62	0.41	MLT 99			
Actinomyces naeslundii		48.38	0.50	URE 10			
27413142	**Very good identification**						**27413142**
Propionibacterium propionicum		95.58	0.60	MLT 99			
Actinomyces naeslundii		4.42	0.47				
27413143	**Low discrimination**						**27413143**
Propionibacterium propionicum		61.55	0.44	MLT 99			
Actinomyces naeslundii		38.46	0.50	URE 10			
27413160	**Low discrimination**						**27413160**
Actinomyces naeslundii		87.30	0.47	MLT 94			
Propionibacterium propionicum		12.70	0.23				
27413161	**Good identification**						**27413161**
Actinomyces naeslundii		100.0	0.50	MLT 94			
27413162	**Low discrimination**						**27413162**
Actinomyces naeslundii		82.09	0.47	MLT 94			
Propionibacterium propionicum		17.91	0.26				
27413163	**Good identification**						**27413163**
Actinomyces naeslundii		100.0	0.50	MLT 94			
27414100	**Good identification**						**27414100**
Fusobacterium mortiferum		100.0	0.36	NIT 1			
27415000	**Low discrimination**						**27415000**
					GRA	CAT	
Lactobacillus fermentum		66.63	0.42	NIT 1	99	1	
Fusobacterium mortiferum		22.68	0.32		1	5	
Propionibacterium granulosum		10.70	0.24		99	90	
27415002	**Low discrimination**						**27415002**
					CAT		
Lactobacillus fermentum		60.31	0.42	NIT 1	1		
Actinomyces viscosus		34.99	0.24		76		
Propionibacterium granulosum		2.42	0.13		90		
27415100	**Low discrimination**						**27415100**
					GRA		
Lactobacillus fermentum		67.26	0.46	NIT 1	99		
Fusobacterium mortiferum		32.74	0.39		1		
27415102	**Low discrimination**						**27415102**
					GRA		
Lactobacillus fermentum		58.13	0.46	NIT 1	99		
Actinomyces viscosus		36.75	0.28		99		
Fusobacterium mortiferum		3.14	0.22		1		
27415120	**Low discrimination**						**27415120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Lactobacillus fermentum		80.55	0.37	NIT 1	99	1	
Capnocytophaga ochracea		8.19	0.05		1	90	
Clostridium perfringens		6.29	0.06		99	1	
27415122	**Low discrimination**						**27415122**
					GRA	CAT	
Lactobacillus fermentum		34.47	0.37	NIT 1	99	1	
Capnocytophaga ochracea		31.56	0.21		1	90	
Parabacteroides distasonis		19.17	0.30		1	76	
27415322	**Good identification**						**27415322**
Parabacteroides distasonis		100.0	0.38	NIT 1			
27415400	**Good identification**						**27415400**
Lactobacillus fermentum		100.0	0.42	NIT 1			
27415402	**Good identification**						**27415402**
Lactobacillus fermentum		100.0	0.42	NIT 1			
27415500	**Acceptable identification**						**27415500**
Lactobacillus fermentum		94.45	0.46	NIT 1			
Bifidobacterium longum subsp. longum		5.55	0.17				
27415502	**Acceptable identification**						**27415502**
Lactobacillus fermentum		94.45	0.46	NIT 1			
Bifidobacterium longum subsp. longum		5.55	0.17				
27415520	**Low discrimination**						**27415520**
					GRA		
Lactobacillus fermentum		47.78	0.37	NIT 1	99		
Bifidobacterium longum subsp. longum		33.67	0.28		99		
Bacteroides vulgatus		18.55	0.09		1		
27415522	**Low discrimination**						**27415522**
					GRA		
Lactobacillus fermentum		45.31	0.37	NIT 1	99		
Bifidobacterium longum subsp. longum		31.93	0.28		99		
Parabacteroides distasonis		15.45	0.27		1		
27421100	**Low discrimination**						**27421100**
					GRA		
Actinomyces naeslundii		83.15	0.38	MLT 94	99		
Fusobacterium mortiferum		16.85	0.15	FRU 95	1		
27421101	**Low discrimination**						**27421101**
					SPO		
Actinomyces naeslundii		83.66	0.41	MLT 94	1		
Clostridium chauvoei		16.34	0.06	FRU 95	99		
27421103	**Low discrimination**						**27421103**
Actinomyces naeslundii		63.60	0.41	MLT 94			
Actinomyces viscosus		36.40	0.13	FRU 95			
27421120	**Low discrimination**						**27421120**
Actinomyces naeslundii		66.57	0.38	MLT 94			
Clostridium perfringens		33.44	0.11	FRU 95			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27421121	**Good identification**					**27421121**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27421122	**Low discrimination**					**27421122**
Actinomyces naeslundii		80.88	0.38	MLT 94		
Clostridium perfringens		19.12	0.05	FRU 95		
27421123	**Good identification**					**27421123**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27423001	**Good identification**					**27423001**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27423002	**Low discrimination**					**27423002**
Actinomyces naeslundii		54.11	0.35	MLT 94		
Propionibacterium granulosum		45.89	0.19	FRU 95		
27423003	**Good identification**					**27423003**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27423020	**Low discrimination**					**27423020**
Actinomyces naeslundii		54.11	0.35	MLT 94		
Propionibacterium granulosum		45.89	0.19	FRU 95		
27423021	**Good identification**					**27423021**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27423022	**Low discrimination**					**27423022**
Actinomyces naeslundii		82.51	0.35	MLT 94		
Propionibacterium granulosum		17.49	0.09	FRU 95		
27423023	**Good identification**					**27423023**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27423100	**Low discrimination**					**27423100**
Actinomyces naeslundii		82.95	0.43	MLT 94		
Propionibacterium propionicum		17.05	0.21	FRU 95		
27423101	**Good identification**					**27423101**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27423102	**Low discrimination**					**27423102**
Actinomyces naeslundii		76.43	0.43	MLT 94		
Propionibacterium propionicum		23.57	0.24	FRU 95		
27423103	**Good identification**					**27423103**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27423120	**Good identification**					**27423120**
Actinomyces naeslundii		100.0	0.43	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27423121	**Good identification**						**27423121**
Actinomyces naeslundii		100.0	0.46	MLT 94			
27423122	**Good identification**						**27423122**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27423123	**Good identification**						**27423123**
Actinomyces naeslundii		100.0	0.46	MLT 94			
27430000	**Very good identification**						**27430000**
					GRA	CAT	
Propionibacterium granulosum		97.92	0.66	NIT 1	99	90	
Fusobacterium mortiferum		1.85	0.40		1	5	
Actinomyces viscosus		0.15	0.08		99	76	
27430002	**Low discrimination**						**27430002**
					GRA	CAT	
Propionibacterium granulosum		62.36	0.56	NIT 1	99	90	
Actinomyces viscosus		36.89	0.42		99	76	
Fusobacterium mortiferum		0.52	0.23		1	5	
27430010	**Good identification**						**27430010**
Propionibacterium granulosum		100.0	0.51	NIT 1			
27430012	**Good identification**						**27430012**
Propionibacterium granulosum		95.80	0.41	NIT 1			
Actinomyces viscosus		4.20	0.08	MLZ 12			
27430020	**Good identification**						**27430020**
Propionibacterium granulosum		99.64	0.56	NIT 1			
Actinomyces naeslundii		0.37	0.29				
27430022	**Good identification**						**27430022**
Propionibacterium granulosum		96.29	0.45	NIT 1			
Actinomyces viscosus		2.30	0.08				
Actinomyces naeslundii		1.41	0.29				
27430030	**Good identification**						**27430030**
Propionibacterium granulosum		100.0	0.41	NIT 1			
27430100	**Low discrimination**						**27430100**
					GRA	CAT	
Fusobacterium mortiferum		72.54	0.46	NIT 1	1	5	
Propionibacterium granulosum		16.63	0.32		99	90	
Actinomyces viscosus		4.37	0.12		99	76	
27430101	**Low discrimination**						**27430101**
					GRA		
Actinomyces naeslundii		74.62	0.40	MLT 94	99		
Actinomyces viscosus		16.89	0.04	GAL 96	99		
Fusobacterium mortiferum		8.49	0.12		1		
27430102	**Good identification**						**27430102**
					GRA		
Actinomyces viscosus		95.64	0.47	MLT 99	99		
Fusobacterium mortiferum		1.78	0.30	TRE 7	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.95	0.37		99		
27430103	**Good identification**						**27430103**
Actinomyces viscosus		95.73	0.38	MLT 99			
Actinomyces naeslundii		4.27	0.40	TRE 7			
27430120	**Low discrimination**						**27430120**
					CAT		
Actinomyces naeslundii		36.06	0.37	MLT 94	25		
Propionibacterium granulosum		34.92	0.21	GAL 96	90		
Clostridium perfringens		22.88	0.11		1		
27430121	**Good identification**						**27430121**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27430123	**Low discrimination**						**27430123**
Actinomyces naeslundii		81.55	0.40	MLT 94			
Actinomyces viscosus		18.46	0.04	GAL 96			
27430200	**Good identification**						**27430200**
Propionibacterium granulosum		100.0	0.40	NIT 1			
27431000	**Very good identification**						**27431000**
					GRA	CAT	
Propionibacterium granulosum		95.07	0.66	NIT 1	99	90	
Fusobacterium mortiferum		2.69	0.43		1	5	
Actinomyces naeslundii		2.09	0.52		99	25	
27431001	**Low discrimination**						**27431001**
Actinomyces naeslundii		76.56	0.55	MLT 94			
Propionibacterium granulosum		23.44	0.32				
27431002	**Low discrimination**						**27431002**
Propionibacterium granulosum		68.35	0.56	NIT 1			
Actinomyces viscosus		24.78	0.39				
Actinomyces naeslundii		6.01	0.52				
27431003	**Low discrimination**						**27431003**
Actinomyces naeslundii		50.18	0.55	MLT 94			
Actinomyces viscosus		45.98	0.31				
Propionibacterium granulosum		3.84	0.21				
27431010	**Good identification**						**27431010**
Propionibacterium granulosum		99.16	0.51	NIT 1			
Actinomyces naeslundii		0.84	0.30	MLZ 12			
27431012	**Acceptable identification**						**27431012**
Propionibacterium granulosum		94.27	0.41	NIT 1			
Actinomyces naeslundii		3.20	0.30	MLZ 12			
Actinomyces viscosus		2.53	0.05				
27431020	**Low discrimination**						**27431020**
					CAT		
Propionibacterium granulosum		84.94	0.56	NIT 1	90		
Clostridium perfringens		7.59	0.30		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		7.47	0.52		25		
27431021	**Acceptable identification**						**27431021**
Actinomyces naeslundii		92.89	0.55	MLT 94			
Propionibacterium granulosum		7.11	0.21				
27431022	**Low discrimination**						**27431022**
					CAT		
Propionibacterium granulosum		65.16	0.45	NIT 1	90		
Actinomyces naeslundii		22.93	0.52		25		
Clostridium perfringens		10.96	0.25		1		
27431023	**Good identification**						**27431023**
Actinomyces naeslundii		98.12	0.55	MLT 94			
Propionibacterium granulosum		1.88	0.11				
27431030	**Good identification**						**27431030**
Propionibacterium granulosum		96.72	0.41	NIT 1			
Actinomyces naeslundii		3.28	0.30	MLZ 12			
27431041	**Good identification**						**27431041**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27431042	**Low discrimination**						**27431042**
Actinomyces naeslundii		53.00	0.39	MLT 94			
Propionibacterium granulosum		34.49	0.21				
Actinomyces viscosus		12.51	0.05				
27431043	**Good identification**						**27431043**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27431061	**Good identification**						**27431061**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27431062	**Low discrimination**						**27431062**
					CAT		
Actinomyces naeslundii		59.58	0.39	MLT 94	25		
Clostridium perfringens		30.73	0.12		1		
Propionibacterium granulosum		9.69	0.11		90		
27431063	**Good identification**						**27431063**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27431100	**Low discrimination**						**27431100**
					GRA	CAT	
Fusobacterium mortiferum		44.84	0.49	NIT 1	1	5	
Actinomyces naeslundii		42.47	0.60		99	25	
Propionibacterium granulosum		6.85	0.32		99	90	
27431101	**Very good identification**						**27431101**
					GRA	SPO	
Actinomyces naeslundii		96.44	0.63	MLT 94	99	1	
Clostridium chauvoei		2.31	0.12		99	99	
Fusobacterium mortiferum		0.69	0.15		1	1	
27431102	**Low discrimination**						**27431102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		67.56	0.43	MLT 99	99		
Actinomyces naeslundii		26.24	0.60	TRE 7	99		
Fusobacterium mortiferum		3.08	0.33		1		
27431103	**Low discrimination**						**27431103**
					SPO		
Actinomyces naeslundii		63.43	0.63	MLT 94	1		
Actinomyces viscosus		36.30	0.35		1		
Clostridium chauvoei		0.27	-.01		99		
27431110	**Low discrimination**						**27431110**
					GRA	CAT	
Actinomyces naeslundii		61.70	0.38	MLT 94	99	25	
Propionibacterium granulosum		25.80	0.17	MLZ 5	99	90	
Fusobacterium mortiferum		12.50	0.15		1	5	
27431111	**Good identification**						**27431111**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27431112	**Low discrimination**						**27431112**
Actinomyces naeslundii		66.93	0.38	MLT 94			
Actinomyces viscosus		33.08	0.09	MLZ 5			
27431113	**Acceptable identification**						**27431113**
Actinomyces naeslundii		90.11	0.41	MLT 94			
Actinomyces viscosus		9.89	0.01	MLZ 5			
27431120	**Low discrimination**						**27431120**
					CAT		
Clostridium perfringens		71.13	0.45	MLT 99	1		
Actinomyces naeslundii		27.18	0.60		25		
Propionibacterium granulosum		1.10	0.21		90		
27431121	**Very good identification**						**27431121**
					SPO		
Actinomyces naeslundii		97.50	0.63	MLT 94	1		
Clostridium perfringens		1.72	0.11		84		
Clostridium chauvoei		0.78	0.04		99		
27431122	**Low discrimination**						**27431122**
					GRA		
Clostridium perfringens		52.95	0.39	MLT 99	99		
Actinomyces naeslundii		43.00	0.60		99		
Parabacteroides distasonis		2.21	0.25		1		
27431123	**Very good identification**						**27431123**
Actinomyces naeslundii		98.61	0.63	MLT 94			
Clostridium perfringens		0.82	0.05				
Actinomyces viscosus		0.57	0.01				
27431130	**Low discrimination**						**27431130**
Actinomyces naeslundii		66.57	0.38	MLT 94			
Clostridium perfringens		33.44	0.11	MLZ 5			
27431131	**Good identification**						**27431131**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27431132	**Low discrimination**						**27431132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		56.78	0.38	MLT 94	99	
Parabacteroides distasonis		29.80	0.20	MLZ 5	1	
Clostridium perfringens		13.42	0.05		99	
27431133	**Good identification**					**27431133**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27431140	**Low discrimination**					**27431140**
Actinomyces naeslundii		78.94	0.47	MLT 94		
Propionibacterium propionicum		9.40	0.21			
Clostridium perfringens		6.90	0.07			
27431141	**Good identification**					**27431141**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27431142	**Low discrimination**					**27431142**
Actinomyces naeslundii		73.15	0.47	MLT 94		
Propionibacterium propionicum		13.06	0.24			
Actinomyces viscosus		10.78	0.09			
27431143	**Good identification**					**27431143**
Actinomyces naeslundii		96.83	0.50	MLT 94		
Actinomyces viscosus		3.17	0.01			
27431161	**Good identification**					**27431161**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27431163	**Good identification**					**27431163**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27431200	**Good identification**					**27431200**
Propionibacterium granulosum		100.0	0.40	NIT 1		
27432000	**Very good identification**					**27432000**
					CAT	
Propionibacterium granulosum		99.59	0.64	NIT 1	90	
Actinomyces naeslundii		0.24	0.34		25	
Propionibacterium propionicum		0.17	0.21		1	
27432002	**Good identification**					**27432002**
					CAT	
Propionibacterium granulosum		97.29	0.53	NIT 1	90	
Propionibacterium propionicum		1.02	0.24		1	
Actinomyces naeslundii		0.92	0.34		25	
27432003	**Low discrimination**					**27432003**
Actinomyces naeslundii		52.61	0.37	MLT 94		
Propionibacterium granulosum		37.57	0.19	GAL 96		
Actinomyces viscosus		9.82	0.00			
27432010	**Good identification**					**27432010**
Propionibacterium granulosum		100.0	0.49	NIT 1		
27432012	**Good identification**					**27432012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		100.0	0.39	NIT 1		
27432020	**Good identification**					**27432020**
Propionibacterium granulosum		99.07	0.53	NIT 1		
Actinomyces naeslundii		0.93	0.34			
27432021	**Low discrimination**					**27432021**
Actinomyces naeslundii		58.34	0.37	MLT 94		
Propionibacterium granulosum		41.66	0.19	GAL 96		
27432022	**Good identification**					**27432022**
Propionibacterium granulosum		96.37	0.43	NIT 1		
Actinomyces naeslundii		3.64	0.34			
27432023	**Low discrimination**					**27432023**
Actinomyces naeslundii		84.86	0.37	MLT 94		
Propionibacterium granulosum		15.15	0.09	GAL 96		
27432030	**Good identification**					**27432030**
Propionibacterium granulosum		100.0	0.39	NIT 1		
27432100	**Acceptable identification**					**27432100**
					CAT	
Propionibacterium propionicum		90.84	0.55	MLT 99	1	
Propionibacterium granulosum		5.32	0.30		90	
Actinomyces naeslundii		3.53	0.41		25	
27432101	**Low discrimination**					**27432101**
Propionibacterium propionicum		65.57	0.39	MLT 99		
Actinomyces naeslundii		34.43	0.44	URE 10		
27432102	**Very good identification**					**27432102**
Propionibacterium propionicum		95.30	0.58	MLT 99		
Actinomyces naeslundii		2.47	0.41			
Actinomyces viscosus		1.30	0.12			
27432103	**Low discrimination**					**27432103**
Propionibacterium propionicum		71.90	0.42	MLT 99		
Actinomyces naeslundii		25.17	0.44	URE 10		
Actinomyces viscosus		2.93	0.04			
27432120	**Low discrimination**					**27432120**
					CAT	
Actinomyces naeslundii		61.12	0.41	MLT 94	25	
Propionibacterium granulosum		23.00	0.19	GAL 96	90	
Propionibacterium propionicum		15.87	0.21		1	
27432121	**Good identification**					**27432121**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27432122	**Low discrimination**					**27432122**
					CAT	
Actinomyces naeslundii		67.40	0.41	MLT 94	25	
Propionibacterium propionicum		26.26	0.24	GAL 96	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		6.34	0.09		90	
27432123	**Good identification**					**27432123**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27432140	**Good identification**					**27432140**
Propionibacterium propionicum		99.32	0.55	MLT 99		
Actinomyces naeslundii		0.68	0.29			
27432141	**Acceptable identification**					**27432141**
Propionibacterium propionicum		91.52	0.39	MLT 99		
Actinomyces naeslundii		8.48	0.32	URE 10		
27432142	**Very good identification**					**27432142**
Propionibacterium propionicum		99.54	0.58	MLT 99		
Actinomyces naeslundii		0.46	0.29			
27432143	**Acceptable identification**					**27432143**
Propionibacterium propionicum		94.18	0.42	MLT 99		
Actinomyces naeslundii		5.82	0.32	URE 10		
27432200	**Good identification**					**27432200**
Propionibacterium granulosum		100.0	0.38	NIT 1		
27433000	**Good identification**					**27433000**
					CAT	
Propionibacterium granulosum		94.49	0.64	NIT 1	90	
Actinomyces naeslundii		5.35	0.57		25	
Propionibacterium propionicum		0.17	0.21		1	
27433001	**Low discrimination**					**27433001**
Actinomyces naeslundii		89.37	0.60	MLT 94		
Propionibacterium granulosum		10.64	0.30			
27433002	**Low discrimination**					**27433002**
					CAT	
Propionibacterium granulosum		80.54	0.53	NIT 1	90	
Actinomyces naeslundii		18.23	0.57		25	
Propionibacterium propionicum		0.84	0.24		1	
27433003	**Very good identification**					**27433003**
Actinomyces naeslundii		97.11	0.60	MLT 94		
Propionibacterium granulosum		2.89	0.19			
27433010	**Good identification**					**27433010**
Propionibacterium granulosum		97.86	0.49	NIT 1		
Actinomyces naeslundii		2.14	0.35	MLZ 12		
27433011	**Low discrimination**					**27433011**
Actinomyces naeslundii		76.43	0.38	MLT 94		
Propionibacterium granulosum		23.57	0.15	MLZ 5		
27433012	**Acceptable identification**					**27433012**
Propionibacterium granulosum		91.97	0.39	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		8.03	0.35	MLZ 12		
27433013	**Good identification**					**27433013**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27433020	**Low discrimination**					**27433020**
Propionibacterium granulosum		81.54	0.53	NIT 1		
Actinomyces naeslundii		18.46	0.57			
27433021	**Very good identification**					**27433021**
Actinomyces naeslundii		97.11	0.60	MLT 94		
Propionibacterium granulosum		2.89	0.19			
27433022	**Low discrimination**					**27433022**
Propionibacterium granulosum		52.48	0.43	NIT 1		
Actinomyces naeslundii		47.52	0.57			
27433023	**Very good identification**					**27433023**
Actinomyces naeslundii		99.26	0.60	MLT 94		
Propionibacterium granulosum		0.74	0.09			
27433030	**Acceptable identification**					**27433030**
Propionibacterium granulosum		91.97	0.39	NIT 1		
Actinomyces naeslundii		8.03	0.35	MLZ 12		
27433031	**Good identification**					**27433031**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27433033	**Good identification**					**27433033**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27433041	**Good identification**					**27433041**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27433042	**Low discrimination**					**27433042**
					CAT	
Actinomyces naeslundii		66.02	0.44	MLT 94	25	
Propionibacterium propionicum		17.28	0.24		1	
Propionibacterium granulosum		16.70	0.19		90	
27433043	**Good identification**					**27433043**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27433060	**Low discrimination**					**27433060**
Actinomyces naeslundii		79.82	0.44	MLT 94		
Propionibacterium granulosum		20.19	0.19			
27433061	**Good identification**					**27433061**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27433062	**Acceptable identification**					**27433062**
Actinomyces naeslundii		94.06	0.44	MLT 94		
Propionibacterium granulosum		5.95	0.09			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27433063	**Good identification**					**27433063**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27433100	**Low discrimination**					**27433100**
					CAT	
Propionibacterium propionicum		50.07	0.55	MLT 99	1	
Actinomyces naeslundii		46.74	0.65		25	
Propionibacterium granulosum		2.93	0.30		90	
27433101	**Good identification**					**27433101**
Actinomyces naeslundii		92.65	0.68	MLT 94		
Propionibacterium propionicum		7.35	0.39			
27433102	**Low discrimination**					**27433102**
					CAT	
Propionibacterium propionicum		60.96	0.58	MLT 99	1	
Actinomyces naeslundii		37.94	0.65		25	
Propionibacterium granulosum		0.60	0.19		90	
27433103	**Low discrimination**					**27433103**
Actinomyces naeslundii		89.13	0.68	MLT 94		
Propionibacterium propionicum		10.61	0.42			
Actinomyces viscosus		0.27	0.01			
27433110	**Low discrimination**					**27433110**
					CAT	
Actinomyces naeslundii		73.09	0.43	MLT 94	25	
Propionibacterium propionicum		15.03	0.21	MLZ 5	1	
Propionibacterium granulosum		11.88	0.15		90	
27433111	**Good identification**					**27433111**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27433112	**Low discrimination**					**27433112**
Actinomyces naeslundii		76.43	0.43	MLT 94		
Propionibacterium propionicum		23.57	0.24	MLZ 5		
27433113	**Good identification**					**27433113**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27433120	**Very good identification**					**27433120**
					CAT	
Actinomyces naeslundii		96.14	0.65	MLT 94	25	
Propionibacterium granulosum		1.51	0.19		90	
Clostridium perfringens		1.31	0.11		1	
27433121	**Very good identification**					**27433121**
Actinomyces naeslundii		100.0	0.68	MLT 94		
27433122	**Very good identification**					**27433122**
Actinomyces naeslundii		97.41	0.65	MLT 94		
Propionibacterium propionicum		1.58	0.24			
Clostridium perfringens		0.62	0.05			
27433123	**Very good identification**					**27433123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.68	MLT 94		
27433130	**Good identification**					**27433130**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27433131	**Good identification**					**27433131**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27433132	**Good identification**					**27433132**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27433133	**Good identification**					**27433133**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27433140	**Low discrimination**					**27433140**
Propionibacterium propionicum		85.86	0.55	MLT 99		
Actinomyces naeslundii		14.14	0.52			
27433141	**Low discrimination**					**27433141**
Actinomyces naeslundii		68.98	0.55	MLT 94		
Propionibacterium propionicum		31.02	0.39			
27433142	**Good identification**					**27433142**
Propionibacterium propionicum		90.11	0.58	MLT 99		
Actinomyces naeslundii		9.90	0.52			
27433143	**Low discrimination**					**27433143**
Actinomyces naeslundii		59.72	0.55	MLT 94		
Propionibacterium propionicum		40.28	0.42			
27433160	**Acceptable identification**					**27433160**
Actinomyces naeslundii		94.22	0.52	MLT 94		
Propionibacterium propionicum		5.78	0.21			
27433161	**Good identification**					**27433161**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27433162	**Acceptable identification**					**27433162**
Actinomyces naeslundii		91.58	0.52	MLT 94		
Propionibacterium propionicum		8.42	0.24			
27433163	**Good identification**					**27433163**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27433200	**Good identification**					**27433200**
Propionibacterium granulosum		98.19	0.38	NIT 1		
Actinomyces naeslundii		1.82	0.23	RHA 3		
27435122	**Good identification**					**27435122**
					GRA	
Parabacteroides distasonis		95.35	0.41	NIT 1	1	
Clostridium perfringens		2.57	0.05		99	
Actinomyces naeslundii		2.09	0.26		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27435132	**Good identification**					**27435132**
Parabacteroides distasonis		100.0	0.37	NIT 1		
27435322	**Good identification**					**27435322**
Parabacteroides distasonis		100.0	0.50	NIT 1		
27435332	**Good identification**					**27435332**
Parabacteroides distasonis		100.0	0.45	NIT 1		
27435522	**Low discrimination**					**27435522**
					GRA	
Parabacteroides distasonis		89.81	0.38	NIT 1	1	
Bifidobacterium longum subsp. longum		10.19	0.17		99	
27435722	**Good identification**					**27435722**
Parabacteroides distasonis		100.0	0.46	NIT 1		
27435732	**Good identification**					**27435732**
Parabacteroides distasonis		100.0	0.41	NIT 1		
27441100	**Low discrimination**					**27441100**
					GRA	
Actinomyces naeslundii		75.13	0.35	MLT 94	99	
Fusobacterium mortiferum		24.87	0.16	FRU 95	1	
27441101	**Good identification**					**27441101**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27441120	**Low discrimination**					**27441120**
					GRA	
Actinomyces naeslundii		72.39	0.35	MLT 94	99	
Prevotella melaninogenica		27.62	-.01	FRU 95	1	
27441121	**Good identification**					**27441121**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27441122	**Good identification**					**27441122**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27441123	**Good identification**					**27441123**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27443001	**Good identification**					**27443001**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27443003	**Good identification**					**27443003**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27443021	**Good identification**					**27443021**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27443023	**Good identification**					**27443023**
Actinomyces naeslundii		100.0	0.36	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27443100	**Good identification**						**27443100**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27443101	**Good identification**						**27443101**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27443102	**Good identification**						**27443102**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27443103	**Good identification**						**27443103**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27443120	**Good identification**						**27443120**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27443121	**Good identification**						**27443121**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27443122	**Good identification**						**27443122**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27443123	**Good identification**						**27443123**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27450000	**Low discrimination**						**27450000**
					GRA	CAT	
Fusobacterium mortiferum		69.45	0.40	NIT 1	1	5	
Actinomyces viscosus		17.39	0.18		99	76	
Propionibacterium granulosum		11.10	0.24		99	90	
27450002	**Good identification**						**27450002**
					GRA	CAT	
Actinomyces viscosus		99.28	0.52	MLT 99	99	76	
Fusobacterium mortiferum		0.45	0.24		1	5	
Propionibacterium granulosum		0.16	0.13		99	90	
27450003	**Good identification**						**27450003**
Actinomyces viscosus		99.46	0.44	MLT 99			
Actinomyces naeslundii		0.54	0.29				
27450100	**Low discrimination**						**27450100**
					GRA		
Fusobacterium mortiferum		81.50	0.47	NIT 1	1		
Actinomyces viscosus		15.55	0.22		99		
Actinomyces naeslundii		2.95	0.34		99		
27450102	**Good identification**						**27450102**
					GRA		
Actinomyces viscosus		99.23	0.56	MLT 99	99		
Fusobacterium mortiferum		0.58	0.30		1		
Actinomyces naeslundii		0.19	0.34		99		
27450103	**Good identification**						**27450103**
Actinomyces viscosus		99.14	0.48	MLT 99			
Actinomyces naeslundii		0.86	0.37				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27450121	**Good identification**						**27450121**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27451000	**Low discrimination**						**27451000**
					GRA	CAT	
Fusobacterium mortiferum		59.38	0.43	NIT 1	1	5	
Actinomyces naeslundii		28.22	0.50		99	25	
Propionibacterium granulosum		6.33	0.24		99	90	
27451001	**Good identification**						**27451001**
Actinomyces naeslundii		95.43	0.53	MLT 94			
Actinomyces viscosus		4.57	0.06				
27451002	**Acceptable identification**						**27451002**
					GRA		
Actinomyces viscosus		94.30	0.48	MLT 99	99		
Actinomyces naeslundii		4.42	0.50		99		
Fusobacterium mortiferum		1.03	0.27		1		
27451003	**Low discrimination**						**27451003**
Actinomyces viscosus		82.57	0.40	MLT 99			
Actinomyces naeslundii		17.43	0.53				
27451020	**Acceptable identification**						**27451020**
					CAT		
Actinomyces naeslundii		91.79	0.50	MLT 94	25		
Propionibacterium granulosum		5.15	0.13		90		
Clostridium perfringens		3.06	0.02		1		
27451021	**Good identification**						**27451021**
Actinomyces naeslundii		100.0	0.53	MLT 94			
27451022	**Low discrimination**						**27451022**
Actinomyces naeslundii		82.28	0.50	MLT 94			
Actinomyces viscosus		17.72	0.14				
27451023	**Good identification**						**27451023**
Actinomyces naeslundii		95.43	0.53	MLT 94			
Actinomyces viscosus		4.57	0.06				
27451040	**Good identification**						**27451040**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27451041	**Good identification**						**27451041**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27451043	**Low discrimination**						**27451043**
Actinomyces naeslundii		78.67	0.40	MLT 94			
Actinomyces viscosus		21.34	0.06				
27451060	**Good identification**						**27451060**
Actinomyces naeslundii		100.0	0.37	MLT 94			
27451061	**Good identification**						**27451061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MLT 94		
27451062	**Good identification**					**27451062**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27451063	**Good identification**					**27451063**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27451100	**Low discrimination**					**27451100**
					GRA	
Fusobacterium mortiferum		60.04	0.50	NIT 1	1	
Actinomyces naeslundii		34.80	0.57		99	
Actinomyces viscosus		4.68	0.18		99	
27451101	**Very good identification**					**27451101**
					GRA	
Actinomyces naeslundii		96.01	0.60	MLT 94	99	
Actinomyces viscosus		2.87	0.10		99	
Fusobacterium mortiferum		1.12	0.16		1	
27451102	**Acceptable identification**					**27451102**
					GRA	
Actinomyces viscosus		91.53	0.52	MLT 99	99	
Actinomyces naeslundii		6.87	0.57		99	
Fusobacterium mortiferum		1.32	0.33		1	
27451103	**Low discrimination**					**27451103**
Actinomyces viscosus		74.74	0.44	MLT 99		
Actinomyces naeslundii		25.26	0.60			
27451110	**Low discrimination**					**27451110**
					GRA	
Actinomyces naeslundii		75.13	0.35	MLT 94	99	
Fusobacterium mortiferum		24.87	0.16	MLZ 5	1	
27451111	**Good identification**					**27451111**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27451113	**Low discrimination**					**27451113**
Actinomyces naeslundii		63.78	0.38	MLT 94		
Actinomyces viscosus		36.22	0.10	MLZ 5		
27451120	**Low discrimination**					**27451120**
					GRA	
Actinomyces naeslundii		76.26	0.57	MLT 94	99	
Prevotella melaninogenica		13.78	0.15		1	
Clostridium perfringens		6.56	0.17		99	
27451121	**Very good identification**					**27451121**
Actinomyces naeslundii		100.0	0.60	MLT 94		
27451122	**Low discrimination**					**27451122**
					GRA	
Actinomyces naeslundii		64.33	0.57	MLT 94	99	
Capnocytophaga ochracea		15.71	0.21		1	
Actinomyces viscosus		8.65	0.18		99	
27451123	**Very good identification**					**27451123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.10	0.60	MLT 94		
Actinomyces viscosus		2.90	0.10			
27451130	**Good identification**					**27451130**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27451131	**Good identification**					**27451131**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27451132	**Low discrimination**					**27451132**
					GRA	
Actinomyces naeslundii		65.98	0.35	MLT 94	99	
Capnocytophaga ochracea		34.02	0.05	MLZ 5	1	
27451133	**Good identification**					**27451133**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27451140	**Acceptable identification**					**27451140**
					GRA	
Actinomyces naeslundii		91.01	0.44	MLT 94	99	
Fusobacterium mortiferum		8.99	0.16		1	
27451141	**Good identification**					**27451141**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27451142	**Low discrimination**					**27451142**
Actinomyces naeslundii		56.74	0.44	MLT 94		
Actinomyces viscosus		43.26	0.18			
27451143	**Low discrimination**					**27451143**
Actinomyces naeslundii		85.52	0.47	MLT 94		
Actinomyces viscosus		14.49	0.10			
27451160	**Acceptable identification**					**27451160**
Actinomyces naeslundii		91.51	0.44	MLT 94		
Clostridium perfringens		8.49	0.04			
27451161	**Good identification**					**27451161**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27451162	**Low discrimination**					**27451162**
					GRA	
Actinomyces naeslundii		86.67	0.44	MLT 94	99	
Capnocytophaga ochracea		13.33	0.05		1	
27451163	**Good identification**					**27451163**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27452100	**Low discrimination**					**27452100**
					GRA	
Actinomyces naeslundii		62.17	0.39	MLT 94	99	
Propionibacterium propionicum		28.90	0.23	GAL 96	99	
Fusobacterium mortiferum		8.93	0.13		1	
27452101	**Good identification**					**27452101**
Actinomyces naeslundii		100.0	0.42	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27452103	**Low discrimination**					**27452103**
Actinomyces naeslundii		62.38	0.42	MLT 94		
Actinomyces viscosus		37.62	0.14	GAL 96		
27452120	**Good identification**					**27452120**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27452121	**Good identification**					**27452121**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27452122	**Good identification**					**27452122**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27452123	**Good identification**					**27452123**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27453000	**Acceptable identification**					**27453000**
Actinomyces naeslundii		91.98	0.54	MLT 94		
Propionibacterium granulosum		8.02	0.21			
27453001	**Very good identification**					**27453001**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27453002	**Low discrimination**					**27453002**
Actinomyces naeslundii		88.28	0.54	MLT 94		
Actinomyces viscosus		9.80	0.14			
Propionibacterium granulosum		1.92	0.11			
27453003	**Very good identification**					**27453003**
Actinomyces naeslundii		97.59	0.57	MLT 94		
Actinomyces viscosus		2.41	0.06			
27453011	**Good identification**					**27453011**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27453013	**Good identification**					**27453013**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27453020	**Good identification**					**27453020**
Actinomyces naeslundii		97.87	0.54	MLT 94		
Propionibacterium granulosum		2.13	0.11			
27453021	**Very good identification**					**27453021**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27453022	**Good identification**					**27453022**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27453023	**Very good identification**					**27453023**
Actinomyces naeslundii		100.0	0.57	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27453031	**Good identification**					**27453031**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27453033	**Good identification**					**27453033**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27453040	**Good identification**					**27453040**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27453041	**Good identification**					**27453041**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27453042	**Good identification**					**27453042**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27453043	**Good identification**					**27453043**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27453060	**Good identification**					**27453060**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27453061	**Good identification**					**27453061**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27453062	**Good identification**					**27453062**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27453063	**Good identification**					**27453063**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27453100	**Very good identification**					**27453100**
					GRA	
Actinomyces naeslundii		97.24	0.62	MLT 94	99	
Propionibacterium propionicum		1.88	0.23		99	
Fusobacterium mortiferum		0.87	0.16		1	
27453101	**Very good identification**					**27453101**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27453102	**Good identification**					**27453102**
Actinomyces naeslundii		91.05	0.62	MLT 94		
Actinomyces viscosus		6.31	0.18			
Propionibacterium propionicum		2.65	0.26			
27453103	**Very good identification**					**27453103**
Actinomyces naeslundii		98.48	0.65	MLT 94		
Actinomyces viscosus		1.52	0.10			
27453110	**Good identification**					**27453110**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27453111	**Good identification**					**27453111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	MLT 94		
27453112	**Good identification**					**27453112**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27453113	**Good identification**					**27453113**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27453120	**Very good identification**					**27453120**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27453121	**Very good identification**					**27453121**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27453122	**Very good identification**					**27453122**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27453123	**Very good identification**					**27453123**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27453130	**Good identification**					**27453130**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27453131	**Good identification**					**27453131**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27453132	**Good identification**					**27453132**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27453133	**Good identification**					**27453133**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27453140	**Acceptable identification**					**27453140**
Actinomyces naeslundii		90.11	0.49	MLT 94		
Propionibacterium propionicum		9.89	0.23			
27453141	**Good identification**					**27453141**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27453142	**Low discrimination**					**27453142**
Actinomyces naeslundii		85.86	0.49	MLT 94		
Propionibacterium propionicum		14.14	0.26			
27453143	**Good identification**					**27453143**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27453160	**Good identification**					**27453160**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27453161	**Good identification**					**27453161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	MLT 94		
27453162	**Good identification**					**27453162**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27453163	**Good identification**					**27453163**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27461001	**Good identification**					**27461001**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27461003	**Low discrimination**					**27461003**
Actinomyces naeslundii		84.48	0.36	MLT 94		
Actinomyces viscosus		15.52	-.01	FRU 95		
27461021	**Good identification**					**27461021**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27461023	**Good identification**					**27461023**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27461100	**Good identification**					**27461100**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27461101	**Good identification**					**27461101**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27461102	**Low discrimination**					**27461102**
Actinomyces naeslundii		65.95	0.40	MLT 94		
Actinomyces viscosus		34.06	0.11	FRU 95		
27461103	**Low discrimination**					**27461103**
Actinomyces naeslundii		89.71	0.43	MLT 94		
Actinomyces viscosus		10.30	0.03	FRU 95		
27461120	**Good identification**					**27461120**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27461121	**Good identification**					**27461121**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27461122	**Good identification**					**27461122**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27461123	**Good identification**					**27461123**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27463000	**Good identification**					**27463000**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27463001	**Good identification**					**27463001**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.41	MLT 94		
27463002	**Good identification**					**27463002**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27463003	**Good identification**					**27463003**
	Actinomyces naeslundii	100.0	0.41	MLT 94		
27463020	**Good identification**					**27463020**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27463021	**Good identification**					**27463021**
	Actinomyces naeslundii	100.0	0.41	MLT 94		
27463022	**Good identification**					**27463022**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27463023	**Good identification**					**27463023**
	Actinomyces naeslundii	100.0	0.41	MLT 94		
27463100	**Good identification**					**27463100**
	Actinomyces naeslundii	100.0	0.45	MLT 94		
27463101	**Good identification**					**27463101**
	Actinomyces naeslundii	100.0	0.48	MLT 94		
27463102	**Good identification**					**27463102**
	Actinomyces naeslundii	100.0	0.45	MLT 94		
27463103	**Good identification**					**27463103**
	Actinomyces naeslundii	100.0	0.48	MLT 94		
27463120	**Good identification**					**27463120**
	Actinomyces naeslundii	100.0	0.45	MLT 94		
27463121	**Good identification**					**27463121**
	Actinomyces naeslundii	100.0	0.48	MLT 94		
27463122	**Good identification**					**27463122**
	Actinomyces naeslundii	100.0	0.45	MLT 94		
27463123	**Good identification**					**27463123**
	Actinomyces naeslundii	100.0	0.48	MLT 94		
27463141	**Good identification**					**27463141**
	Actinomyces naeslundii	100.0	0.35	MLT 94		
27463143	**Good identification**					**27463143**
	Actinomyces naeslundii	100.0	0.35	MLT 94		
27463161	**Good identification**					**27463161**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.35	MLT 94			
27463163	**Good identification**						**27463163**
Actinomyces naeslundii		100.0	0.35	MLT 94			
27470100	**Low discrimination**						**27470100**
					GRA		
Fusobacterium mortiferum		77.91	0.38	NIT 1	1		
Actinomyces naeslundii		18.34	0.39		99		
Actinomyces viscosus		3.75	0.03		99		
27470101	**Good identification**						**27470101**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27470102	**Acceptable identification**						**27470102**
					GRA		
Actinomyces viscosus		93.21	0.37	MLT 99	99		
Actinomyces naeslundii		4.61	0.39	TRE 7	99		
Fusobacterium mortiferum		2.18	0.21		1		
27470120	**Good identification**						**27470120**
Actinomyces naeslundii		100.0	0.39	MLT 94			
27470121	**Good identification**						**27470121**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27470122	**Low discrimination**						**27470122**
Actinomyces naeslundii		83.04	0.39	MLT 94			
Actinomyces viscosus		16.96	0.03	GAL 96			
27470123	**Good identification**						**27470123**
Actinomyces naeslundii		100.0	0.42	MLT 94			
27471000	**Low discrimination**						**27471000**
					GRA	CAT	
Actinomyces naeslundii		59.87	0.54	MLT 94	99	25	
Propionibacterium granulosum		20.75	0.32		99	90	
Fusobacterium mortiferum		19.39	0.34		1	5	
27471001	**Very good identification**						**27471001**
Actinomyces naeslundii		100.0	0.57	MLT 94			
27471002	**Low discrimination**						**27471002**
Actinomyces naeslundii		51.30	0.54	MLT 94			
Actinomyces viscosus		42.41	0.29				
Propionibacterium granulosum		4.44	0.21				
27471003	**Low discrimination**						**27471003**
Actinomyces naeslundii		84.48	0.57	MLT 94			
Actinomyces viscosus		15.52	0.21				
27471011	**Good identification**						**27471011**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27471013	**Good identification**						**27471013**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	MLT 94		
27471020	**Low discrimination**					**27471020**
					CAT	
Actinomyces naeslundii		89.40	0.54	MLT 94	25	
Propionibacterium granulosum		7.75	0.21		90	
Clostridium perfringens		2.85	0.07		1	
27471021	**Very good identification**					**27471021**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27471022	**Good identification**					**27471022**
					CAT	
Actinomyces naeslundii		96.46	0.54	MLT 94	25	
Propionibacterium granulosum		2.09	0.11		90	
Clostridium perfringens		1.45	0.01		1	
27471023	**Very good identification**					**27471023**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27471031	**Good identification**					**27471031**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27471033	**Good identification**					**27471033**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27471040	**Good identification**					**27471040**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27471041	**Good identification**					**27471041**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27471042	**Good identification**					**27471042**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27471043	**Good identification**					**27471043**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27471060	**Good identification**					**27471060**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27471061	**Good identification**					**27471061**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27471062	**Good identification**					**27471062**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27471063	**Good identification**					**27471063**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27471100	**Low discrimination**					**27471100**
					GRA	
Actinomyces naeslundii		78.70	0.62	MLT 94	99	
Fusobacterium mortiferum		20.89	0.41		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.41	-.01		99	
27471101	**Very good identification**					**27471101**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27471102	**Low discrimination**					**27471102**
					GRA	
Actinomyces naeslundii		64.46	0.62	MLT 94	99	
Actinomyces viscosus		33.29	0.33		99	
Fusobacterium mortiferum		1.90	0.24		1	
27471103	**Low discrimination**					**27471103**
Actinomyces naeslundii		89.71	0.65	MLT 94		
Actinomyces viscosus		10.29	0.25			
27471110	**Good identification**					**27471110**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27471111	**Good identification**					**27471111**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27471112	**Acceptable identification**					**27471112**
Actinomyces naeslundii		90.98	0.40	MLT 94		
Actinomyces viscosus		9.02	-.01	MLZ 5		
27471113	**Good identification**					**27471113**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27471120	**Good identification**					**27471120**
Actinomyces naeslundii		92.40	0.62	MLT 94		
Clostridium perfringens		7.60	0.21			
27471121	**Very good identification**					**27471121**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27471122	**Good identification**					**27471122**
					GRA	
Actinomyces naeslundii		93.45	0.62	MLT 94	99	
Clostridium perfringens		3.62	0.16		99	
Parabacteroides distasonis		1.95	0.20		1	
27471123	**Very good identification**					**27471123**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27471130	**Good identification**					**27471130**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27471131	**Good identification**					**27471131**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27471132	**Low discrimination**					**27471132**
					GRA	
Actinomyces naeslundii		82.43	0.40	MLT 94	99	
Parabacteroides distasonis		17.57	0.16	MLZ 5	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27471133	**Good identification**					**27471133**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27471140	**Good identification**					**27471140**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27471141	**Good identification**					**27471141**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27471142	**Good identification**					**27471142**
Actinomyces naeslundii		97.13	0.49	MLT 94		
Actinomyces viscosus		2.87	-.01			
27471143	**Good identification**					**27471143**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27471160	**Acceptable identification**					**27471160**
Actinomyces naeslundii		91.85	0.49	MLT 94		
Clostridium perfringens		8.16	0.09			
27471161	**Good identification**					**27471161**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27471162	**Good identification**					**27471162**
Actinomyces naeslundii		95.99	0.49	MLT 94		
Clostridium perfringens		4.01	0.03			
27471163	**Good identification**					**27471163**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27472001	**Good identification**					**27472001**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27472002	**Low discrimination**					**27472002**
Actinomyces naeslundii		50.59	0.36	MLT 94		
Propionibacterium granulosum		40.88	0.19	GAL 96		
Actinomyces viscosus		8.52	-.02			
27472003	**Good identification**					**27472003**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27472020	**Low discrimination**					**27472020**
Actinomyces naeslundii		55.31	0.36	MLT 94		
Propionibacterium granulosum		44.69	0.19	GAL 96		
27472021	**Good identification**					**27472021**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27472022	**Low discrimination**					**27472022**
Actinomyces naeslundii		83.20	0.36	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		16.80	0.09	GAL 96		
27472023	**Good identification**					**27472023**
Actinomyces naeslundii		100.0	0.39	MLT 94		
27472100	**Low discrimination**					**27472100**
Actinomyces naeslundii		83.62	0.44	MLT 94		
Propionibacterium propionicum		16.38	0.21	GAL 96		
27472101	**Good identification**					**27472101**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27472102	**Low discrimination**					**27472102**
Actinomyces naeslundii		71.48	0.44	MLT 94		
Propionibacterium propionicum		21.01	0.24	GAL 96		
Actinomyces viscosus		7.52	0.03			
27472103	**Good identification**					**27472103**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27472120	**Good identification**					**27472120**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27472121	**Good identification**					**27472121**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27472122	**Good identification**					**27472122**
Actinomyces naeslundii		100.0	0.44	MLT 94		
27472123	**Good identification**					**27472123**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27473000	**Low discrimination**					**27473000**
Actinomyces naeslundii		88.13	0.59	MLT 94		
Propionibacterium granulosum		11.87	0.30			
27473001	**Very good identification**					**27473001**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27473002	**Very good identification**					**27473002**
Actinomyces naeslundii		96.74	0.59	MLT 94		
Propionibacterium granulosum		3.26	0.19			
27473003	**Very good identification**					**27473003**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27473010	**Low discrimination**					**27473010**
Actinomyces naeslundii		74.13	0.38	MLT 94		
Propionibacterium granulosum		25.87	0.15	MLZ 5		
27473011	**Good identification**					**27473011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MLT 94		
27473012	**Good identification**					**27473012**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27473013	**Good identification**					**27473013**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27473020	**Very good identification**					**27473020**
Actinomyces naeslundii		96.74	0.59	MLT 94		
Propionibacterium granulosum		3.26	0.19			
27473021	**Very good identification**					**27473021**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27473022	**Very good identification**					**27473022**
Actinomyces naeslundii		99.17	0.59	MLT 94		
Propionibacterium granulosum		0.83	0.09			
27473023	**Very good identification**					**27473023**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27473030	**Good identification**					**27473030**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27473031	**Good identification**					**27473031**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27473032	**Good identification**					**27473032**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27473033	**Good identification**					**27473033**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27473040	**Good identification**					**27473040**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27473041	**Good identification**					**27473041**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27473042	**Good identification**					**27473042**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27473043	**Good identification**					**27473043**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27473060	**Good identification**					**27473060**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27473061	**Good identification**					**27473061**
Actinomyces naeslundii		100.0	0.50	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27473062	**Good identification**					**27473062**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27473063	**Good identification**					**27473063**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27473100	**Very good identification**					**27473100**
Actinomyces naeslundii		98.81	0.67	MLT 94		
Propionibacterium propionicum		0.81	0.21			
Actinomyces israelii		0.38	0.17			
27473101	**Very good identification**					**27473101**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27473102	**Very good identification**					**27473102**
Actinomyces naeslundii		97.97	0.67	MLT 94		
Propionibacterium propionicum		1.20	0.24			
Actinomyces israelii		0.57	0.20			
27473103	**Very good identification**					**27473103**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27473110	**Good identification**					**27473110**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27473111	**Good identification**					**27473111**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27473112	**Good identification**					**27473112**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27473113	**Good identification**					**27473113**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27473120	**Very good identification**					**27473120**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27473121	**Very good identification**					**27473121**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27473122	**Very good identification**					**27473122**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27473123	**Very good identification**					**27473123**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27473130	**Good identification**					**27473130**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27473131	**Good identification**					**27473131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	MLT 94		
27473132	**Good identification**					**27473132**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27473133	**Good identification**					**27473133**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27473140	**Good identification**					**27473140**
Actinomyces naeslundii		95.58	0.54	MLT 94		
Propionibacterium propionicum		4.42	0.21			
27473141	**Very good identification**					**27473141**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27473142	**Acceptable identification**					**27473142**
Actinomyces naeslundii		93.51	0.54	MLT 94		
Propionibacterium propionicum		6.49	0.24			
27473143	**Very good identification**					**27473143**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27473151	**Good identification**					**27473151**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27473153	**Good identification**					**27473153**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27473160	**Good identification**					**27473160**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27473161	**Very good identification**					**27473161**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27473162	**Good identification**					**27473162**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27473163	**Very good identification**					**27473163**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27473171	**Good identification**					**27473171**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27473173	**Good identification**					**27473173**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27473301	**Good identification**					**27473301**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27473303	**Good identification**					**27473303**
Actinomyces naeslundii		100.0	0.36	MLT 94		

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
27473321	**Good identification**					**27473321**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27473323	**Good identification**					**27473323**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27473501	**Good identification**					**27473501**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27473503	**Good identification**					**27473503**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27473521	**Good identification**					**27473521**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27473523	**Good identification**					**27473523**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27475122	**Acceptable identification**					**27475122**
					GRA	
Parabacteroides distasonis		94.89	0.37	NIT 1	1	
Actinomyces naeslundii		5.11	0.28		99	
27475322	**Good identification**					**27475322**
Parabacteroides distasonis		100.0	0.45	NIT 1		
27475332	**Good identification**					**27475332**
Parabacteroides distasonis		100.0	0.40	NIT 1		
27475722	**Good identification**					**27475722**
Parabacteroides distasonis		100.0	0.41	NIT 1		
27475732	**Good identification**					**27475732**
Parabacteroides distasonis		95.75	0.37	NIT 1		
Clostridium clostridioforme		4.25	0.08			
27477101	**Good identification**					**27477101**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27477102	**Low discrimination**					**27477102**
Actinomyces israelii		86.89	0.38	MLT 99		
Actinomyces naeslundii		13.11	0.33	SAL 92		
27477103	**Good identification**					**27477103**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27477121	**Good identification**					**27477121**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27477123	**Good identification**					**27477123**
Actinomyces naeslundii		100.0	0.36	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27477502	**Good identification**					**27477502**
Actinomyces israelii		100.0	0.36	MLT 99		
27500000	**Low discrimination**					**27500000**
					COC	
Actinomyces viscosus		73.83	0.40	FRU 95	1	
Gemella morbillorum		15.11	0.30	BGL 99	99	
Actinomyces odontolyticus		6.85	0.29		1	
27500002	**Very good identification**					**27500002**
Actinomyces viscosus		99.94	0.74	FRU 95		
Actinomyces odontolyticus		0.06	0.25			
Actinomyces naeslundii		0.00	0.22			
27500003	**Very good identification**					**27500003**
Actinomyces viscosus		99.98	0.66	FRU 95		
Actinomyces naeslundii		0.02	0.25			
27500012	**Good identification**					**27500012**
Actinomyces viscosus		100.0	0.40	FRU 95		
27500022	**Good identification**					**27500022**
Actinomyces viscosus		98.72	0.40	FRU 95		
Actinomyces odontolyticus		0.91	0.12	NAG 1		
Actinomyces naeslundii		0.37	0.22			
27500042	**Good identification**					**27500042**
Actinomyces viscosus		100.0	0.40	FRU 95		
27500100	**Good identification**					**27500100**
Actinomyces viscosus		96.97	0.44	FRU 95		
Actinomyces odontolyticus		1.60	0.20	BGL 99		
Propionibacterium propionicum		0.85	0.23			
27500101	**Low discrimination**					**27500101**
					SPO	
Actinomyces viscosus		85.56	0.36	FRU 95	1	
Clostridium chauvoei		12.12	0.22	BGL 99	99	
Actinomyces naeslundii		2.32	0.33		1	
27500102	**Very good identification**					**27500102**
Actinomyces viscosus		99.97	0.78	FRU 95		
Propionibacterium propionicum		0.01	0.26			
Actinomyces naeslundii		0.01	0.30			
27500103	**Very good identification**					**27500103**
					SPO	
Actinomyces viscosus		99.95	0.70	FRU 95	1	
Actinomyces naeslundii		0.03	0.33		1	
Clostridium chauvoei		0.03	0.10		99	
27500112	**Good identification**					**27500112**
Actinomyces viscosus		100.0	0.44	FRU 95		
27500113	**Good identification**					**27500113**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		100.0	0.36	FRU 95		
27500122	**Good identification**					**27500122**
					GRA	
Actinomyces viscosus		98.17	0.44	FRU 95	99	
Prevotella melaninogenica		0.79	0.03	NAG 1	1	
Actinomyces naeslundii		0.59	0.30		99	
27500123	**Good identification**					**27500123**
					SPO	
Actinomyces viscosus		96.58	0.36	FRU 95	1	
Actinomyces naeslundii		2.62	0.33	NAG 1	1	
Clostridium chauvoei		0.81	0.01		99	
27500142	**Good identification**					**27500142**
Actinomyces viscosus		98.71	0.44	FRU 95		
Propionibacterium propionicum		1.29	0.26	SOR 1		
27500143	**Good identification**					**27500143**
Actinomyces viscosus		100.0	0.36	FRU 95		
27500202	**Acceptable identification**					**27500202**
Actinomyces viscosus		90.06	0.40	FRU 95		
Actinomyces odontolyticus		9.94	0.30	RHA 1		
27500302	**Good identification**					**27500302**
Actinomyces viscosus		98.08	0.44	FRU 95		
Actinomyces odontolyticus		1.92	0.22	RHA 1		
27500303	**Good identification**					**27500303**
Actinomyces viscosus		100.0	0.36	FRU 95		
27500402	**Good identification**					**27500402**
Actinomyces viscosus		100.0	0.40	FRU 95		
27500502	**Good identification**					**27500502**
Actinomyces viscosus		100.0	0.44	FRU 95		
27500503	**Good identification**					**27500503**
Actinomyces viscosus		100.0	0.36	FRU 95		
27501000	**Low discrimination**					**27501000**
					COC	
Actinomyces viscosus		65.84	0.36	FRU 95	1	
Gemella morbillorum		11.84	0.26	BGL 99	99	
Actinomyces naeslundii		9.70	0.46		1	
27501002	**Very good identification**					**27501002**
Actinomyces viscosus		99.77	0.70	FRU 95		
Actinomyces naeslundii		0.15	0.46			
Actinomyces odontolyticus		0.06	0.22			
27501003	**Very good identification**					**27501003**
					SPO	
Actinomyces viscosus		99.26	0.62	FRU 95	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.67	0.49		1	
Clostridium chauvoei		0.07	0.10		99	
27501012	**Good identification**					**27501012**
Actinomyces viscosus		99.23	0.36	FRU 95		
Actinomyces naeslundii		0.77	0.24	MLZ 1		
27501021	**Low discrimination**					**27501021**
					SPO	
Actinomyces naeslundii		82.88	0.49	FRU 95	1	
Clostridium chauvoei		17.12	0.14		99	
27501022	**Low discrimination**					**27501022**
Actinomyces viscosus		78.34	0.36	FRU 95		
Actinomyces naeslundii		11.54	0.46	NAG 1		
Clostridium perfringens		6.99	0.20			
27501041	**Good identification**					**27501041**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27501042	**Good identification**					**27501042**
Actinomyces viscosus		97.47	0.36	FRU 95		
Actinomyces naeslundii		2.53	0.33	SOR 1		
27501061	**Good identification**					**27501061**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27501063	**Good identification**					**27501063**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27501100	**Low discrimination**					**27501100**
					SPO	
Actinomyces viscosus		68.30	0.40	FRU 95	1	
Actinomyces naeslundii		16.11	0.54	BGL 99	1	
Clostridium chauvoei		5.26	0.22		99	
27501101	**Good identification**					**27501101**
					SPO	
Clostridium chauvoei		91.74	0.57	RAF 1	99	
Actinomyces naeslundii		4.26	0.57		1	
Actinomyces viscosus		4.01	0.32		1	
27501102	**Very good identification**					**27501102**
					GRA	
Actinomyces viscosus		99.64	0.75	FRU 95	99	
Actinomyces naeslundii		0.24	0.54		99	
Capnocytophaga ochracea		0.04	0.15		1	
27501103	**Very good identification**					**27501103**
					SPO	
Actinomyces viscosus		95.10	0.66	FRU 95	1	
Clostridium chauvoei		3.88	0.44		99	
Actinomyces naeslundii		1.02	0.57		1	
27501112	**Good identification**					**27501112**
					GRA	
Actinomyces viscosus		98.33	0.40	FRU 95	99	
Actinomyces naeslundii		1.22	0.32	MLZ 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		0.45	-.01		1		
27501120	**Low discrimination**						**27501120**
					GRA		
Prevotella melaninogenica		86.04	0.50	NIT 1	1		
Clostridium perfringens		9.92	0.40	FRU 90	99		
Actinomyces naeslundii		2.99	0.54		99		
27501121	**Low discrimination**						**27501121**
					GRA	SPO	
Clostridium chauvoei		85.53	0.48	RAF 1	99	99	
Actinomyces naeslundii		11.90	0.57		99	1	
Prevotella melaninogenica		2.31	0.15		1	1	
27501122	**Low discrimination**						**27501122**
					GRA	CAT	
Prevotella melaninogenica		39.31	0.37	NIT 1	1	1	
Actinomyces viscosus		30.42	0.40	FRU 90	99	76	
Capnocytophaga ochracea		11.27	0.32		1	90	
27501123	**Low discrimination**						**27501123**
					SPO		
Clostridium chauvoei		38.80	0.36	RAF 1	99		
Actinomyces naeslundii		30.60	0.57		1		
Actinomyces viscosus		28.82	0.32		1		
27501140	**Low discrimination**						**27501140**
Actinomyces naeslundii		58.97	0.41	FRU 95			
Propionibacterium propionicum		20.20	0.23				
Actinomyces viscosus		14.31	0.06				
27501142	**Acceptable identification**						**27501142**
Actinomyces viscosus		93.67	0.40	FRU 95			
Actinomyces naeslundii		3.90	0.41	SOR 1			
Propionibacterium propionicum		2.00	0.26				
27501161	**Low discrimination**						**27501161**
					SPO		
Actinomyces naeslundii		70.86	0.44	FRU 95	1		
Clostridium chauvoei		29.14	0.14		99		
27501163	**Acceptable identification**						**27501163**
					SPO		
Actinomyces naeslundii		93.23	0.44	FRU 95	1		
Clostridium chauvoei		6.77	0.01		99		
27501202	**Low discrimination**						**27501202**
Actinomyces viscosus		89.28	0.36	FRU 95			
Actinomyces odontolyticus		10.72	0.27	RHA 1			
27501302	**Good identification**						**27501302**
Actinomyces viscosus		97.91	0.40	FRU 95			
Actinomyces odontolyticus		2.09	0.19	RHA 1			
27501402	**Good identification**						**27501402**
Actinomyces viscosus		98.32	0.36	FRU 95			
Lactobacillus fermentum		1.68	0.20	ARA 1			
27501502	**Acceptable identification**						**27501502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		91.89	0.40	FRU 95			
Bifidobacterium longum subsp. longum		6.01	0.27	ARA 1			
Lactobacillus fermentum		1.44	0.24				
27501520	**Low discrimination**						**27501520**
					GRA		
Bifidobacterium longum subsp. longum		74.62	0.37	NIT 1	99		
Prevotella melaninogenica		19.56	0.15	FRU 90	1		
Clostridium perfringens		2.26	0.06		99		
27501522	**Low discrimination**						**27501522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		72.84	0.37	NIT 1	99	1	
Capnocytophaga ochracea		18.22	0.19	FRU 90	1	90	
Prevotella melaninogenica		3.64	0.03		1	1	
27501530	**Good identification**						**27501530**
Bifidobacterium longum subsp. longum		100.0	0.41	NIT 1			
27501532	**Good identification**						**27501532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.40	0.41	NIT 1	99	1	
Capnocytophaga ochracea		1.61	0.02	FRU 90	1	90	
27502002	**Good identification**						**27502002**
Actinomyces viscosus		97.03	0.40	FRU 95			
Propionibacterium propionicum		2.26	0.26	MAN 1			
Actinomyces naeslundii		0.71	0.27				
27502100	**Very good identification**						**27502100**
Propionibacterium propionicum		97.54	0.57	FRU 99			
Actinomyces naeslundii		1.32	0.35				
Actinomyces viscosus		1.14	0.10				
27502101	**Low discrimination**						**27502101**
Propionibacterium propionicum		82.14	0.41	FRU 99			
Actinomyces naeslundii		14.99	0.38	URE 10			
Actinomyces viscosus		2.88	0.02				
27502102	**Low discrimination**						**27502102**
Propionibacterium propionicum		56.19	0.60	FRU 99			
Actinomyces viscosus		43.30	0.44				
Actinomyces naeslundii		0.51	0.35				
27502103	**Low discrimination**						**27502103**
Actinomyces viscosus		67.33	0.36	FRU 95			
Propionibacterium propionicum		29.13	0.44	MAN 1			
Actinomyces naeslundii		3.54	0.38				
27502120	**Low discrimination**						**27502120**
Actinomyces naeslundii		57.23	0.35	FRU 95			
Propionibacterium propionicum		42.77	0.23	GAL 96			
27502121	**Good identification**						**27502121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	FRU 95		
27502123	**Low discrimination**					**27502123**
Actinomyces naeslundii		83.90	0.38	FRU 95		
Actinomyces viscosus		16.11	0.02	GAL 96		
27502140	**Very good identification**					**27502140**
Propionibacterium propionicum		100.0	0.57	FRU 99		
27502141	**Good identification**					**27502141**
Propionibacterium propionicum		96.88	0.41	FRU 99		
Actinomyces naeslundii		3.12	0.25	URE 10		
27502142	**Very good identification**					**27502142**
Propionibacterium propionicum		99.23	0.60	FRU 99		
Actinomyces viscosus		0.77	0.10			
27502143	**Good identification**					**27502143**
Propionibacterium propionicum		95.71	0.44	FRU 99		
Actinomyces viscosus		2.24	0.02	URE 10		
Actinomyces naeslundii		2.06	0.25			
27503000	**Low discrimination**					**27503000**
					CAT	
Actinomyces naeslundii		84.26	0.51	FRU 95	25	
Propionibacterium propionicum		7.47	0.23		1	
Propionibacterium granulosum		5.30	0.16		90	
27503001	**Good identification**					**27503001**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27503002	**Low discrimination**					**27503002**
Actinomyces viscosus		75.53	0.36	FRU 95		
Actinomyces naeslundii		21.60	0.51	MAN 1		
Propionibacterium propionicum		2.87	0.26			
27503003	**Low discrimination**					**27503003**
Actinomyces naeslundii		56.28	0.54	FRU 95		
Actinomyces viscosus		43.72	0.28			
27503020	**Good identification**					**27503020**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27503021	**Good identification**					**27503021**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27503022	**Good identification**					**27503022**
Actinomyces naeslundii		96.59	0.51	FRU 95		
Actinomyces viscosus		3.41	0.02			
27503023	**Good identification**					**27503023**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27503040	**Low discrimination**					**27503040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		66.57	0.38	FRU 95		
Propionibacterium propionicum		33.43	0.23			
27503041	**Good identification**					**27503041**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27503042	**Low discrimination**					**27503042**
Actinomyces naeslundii		51.20	0.38	FRU 95		
Propionibacterium propionicum		38.56	0.26			
Actinomyces viscosus		10.24	0.02			
27503043	**Good identification**					**27503043**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27503060	**Good identification**					**27503060**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27503061	**Good identification**					**27503061**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27503062	**Good identification**					**27503062**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27503063	**Good identification**					**27503063**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27503100	**Low discrimination**					**27503100**
Propionibacterium propionicum		75.10	0.57	FRU 99		
Actinomyces naeslundii		24.36	0.59			
Actinomyces viscosus		0.54	0.06			
27503101	**Low discrimination**					**27503101**
					SPO	
Actinomyces naeslundii		74.60	0.62	FRU 95	1	
Propionibacterium propionicum		17.03	0.41		1	
Clostridium chauvoei		8.37	0.22		99	
27503102	**Low discrimination**					**27503102**
Propionibacterium propionicum		59.22	0.60	FRU 99		
Actinomyces viscosus		27.97	0.40			
Actinomyces naeslundii		12.81	0.59			
27503103	**Low discrimination**					**27503103**
Actinomyces naeslundii		54.13	0.62	FRU 95		
Actinomyces viscosus		26.27	0.32			
Propionibacterium propionicum		18.54	0.44			
27503110	**Low discrimination**					**27503110**
Actinomyces naeslundii		62.83	0.37	FRU 95		
Propionibacterium propionicum		37.17	0.23	MLZ 5		
27503111	**Good identification**					**27503111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	FRU 95		
27503112	**Low discrimination**					**27503112**
Actinomyces naeslundii		43.36	0.37	FRU 95		
Propionibacterium propionicum		38.47	0.26	MLZ 5		
Actinomyces viscosus		18.17	0.06			
27503113	**Good identification**					**27503113**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27503120	**Low discrimination**					**27503120**
					GRA	
Actinomyces naeslundii		83.47	0.59	FRU 95	99	
Prevotella melaninogenica		12.49	0.15		1	
Propionibacterium propionicum		2.60	0.23		99	
27503121	**Very good identification**					**27503121**
					SPO	
Actinomyces naeslundii		96.40	0.62	FRU 95	1	
Clostridium chauvoei		3.60	0.14		99	
27503122	**Low discrimination**					**27503122**
					GRA	
Actinomyces naeslundii		89.80	0.59	FRU 95	99	
Propionibacterium propionicum		4.19	0.26		99	
Prevotella melaninogenica		2.56	0.03		1	
27503123	**Very good identification**					**27503123**
					SPO	
Actinomyces naeslundii		99.34	0.62	FRU 95	1	
Clostridium chauvoei		0.66	0.01		99	
27503130	**Good identification**					**27503130**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27503131	**Good identification**					**27503131**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27503132	**Good identification**					**27503132**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27503133	**Good identification**					**27503133**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27503140	**Good identification**					**27503140**
Propionibacterium propionicum		94.59	0.57	FRU 99		
Actinomyces naeslundii		5.42	0.46			
27503141	**Low discrimination**					**27503141**
Propionibacterium propionicum		56.41	0.41	FRU 99		
Actinomyces naeslundii		43.59	0.49	URE 10		
27503142	**Very good identification**					**27503142**
Propionibacterium propionicum		95.88	0.60	FRU 99		
Actinomyces naeslundii		3.66	0.46			
Actinomyces viscosus		0.46	0.06			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27503143	**Low discrimination**						**27503143**
Propionibacterium propionicum		66.00	0.44	FRU 99			
Actinomyces naeslundii		34.00	0.49	URE 10			
27503160	**Low discrimination**						**27503160**
Actinomyces naeslundii		85.00	0.46	FRU 95			
Propionibacterium propionicum		15.00	0.23				
27503161	**Good identification**						**27503161**
Actinomyces naeslundii		100.0	0.49	FRU 95			
27503162	**Low discrimination**						**27503162**
Actinomyces naeslundii		79.07	0.46	FRU 95			
Propionibacterium propionicum		20.93	0.26				
27503163	**Good identification**						**27503163**
Actinomyces naeslundii		100.0	0.49	FRU 95			
27504002	**Good identification**						**27504002**
Actinomyces viscosus		97.28	0.40	FRU 95			
Actinomyces odontolyticus		2.73	0.20	XYL 1			
27504102	**Good identification**						**27504102**
Actinomyces viscosus		99.51	0.44	FRU 95			
Actinomyces odontolyticus		0.50	0.12	XYL 1			
27504103	**Good identification**						**27504103**
Actinomyces viscosus		100.0	0.36	FRU 95			
27505002	**Good identification**						**27505002**
Actinomyces viscosus		96.11	0.36	FRU 95			
Actinomyces odontolyticus		2.93	0.17	XYL 1			
Lactobacillus fermentum		0.96	0.16				
27505102	**Good identification**						**27505102**
					GRA	CAT	
Actinomyces viscosus		97.57	0.40	FRU 95	99	76	
Capnocytophaga ochracea		1.00	0.05	XYL 1	1	90	
Lactobacillus fermentum		0.90	0.20		99	1	
27505500	**Good identification**						**27505500**
Bifidobacterium longum subsp. longum		95.52	0.35	NIT 1			
Lactobacillus fermentum		4.48	0.20	FRU 90			
27505502	**Acceptable identification**						**27505502**
Bifidobacterium longum subsp. longum		91.04	0.35	NIT 1			
Actinomyces viscosus		4.69	0.06	FRU 90			
Lactobacillus fermentum		4.27	0.20				
27505510	**Good identification**						**27505510**
Bifidobacterium longum subsp. longum		100.0	0.39	NIT 1			
27505512	**Good identification**						**27505512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		100.0	0.39	NIT 1			
27505520	**Good identification**						**27505520**
					GRA		
Bifidobacterium longum subsp. longum		95.23	0.46	NIT 1	99		
Bacteroides vulgatus		4.77	0.09	FRU 90	1		
27505522	**Good identification**						**27505522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.96	0.46	NIT 1	99	1	
Capnocytophaga ochracea		2.04	0.09	FRU 90	1	90	
27505530	**Good identification**						**27505530**
Bifidobacterium longum subsp. longum		100.0	0.50	NIT 1			
27505532	**Good identification**						**27505532**
Bifidobacterium longum subsp. longum		100.0	0.50	NIT 1			
27510000	**Low discrimination**						**27510000**
					GRA	CAT	
Actinomyces viscosus		76.11	0.62	BGL 99	99	76	
Prevotella intermedia		12.09	0.49		1	1	
Propionibacterium granulosum		5.80	0.52		99	90	
27510001	**Good identification**						**27510001**
					GRA		
Actinomyces viscosus		97.49	0.53	BGL 99	99		
Actinomyces naeslundii		1.65	0.47		99		
Prevotella intermedia		0.47	0.15		1		
27510002	**Excellent identification**						**27510002**
					CAT		
Actinomyces viscosus		99.94	0.96		76		
Actinomyces odontolyticus		0.04	0.44		10		
Propionibacterium granulosum		0.02	0.41		90		
27510003	**Very good identification**						**27510003**
Actinomyces viscosus		99.98	0.88				
Actinomyces naeslundii		0.02	0.47				
Actinomyces odontolyticus		0.00	0.09				
27510010	**Low discrimination**						**27510010**
					GRA	CAT	
Propionibacterium granulosum		45.45	0.37	NIT 1	99	90	
Actinomyces viscosus		44.18	0.27	MLZ 12	99	76	
Prevotella intermedia		7.02	0.15		1	1	
27510012	**Very good identification**						**27510012**
					CAT		
Actinomyces viscosus		99.69	0.62	MLZ 1	76		
Propionibacterium granulosum		0.26	0.27		90		
Actinomyces odontolyticus		0.04	0.09		10		
27510013	**Good identification**						**27510013**
Actinomyces viscosus		99.91	0.53	MLZ 1			
Actinomyces naeslundii		0.09	0.25				
27510020	**Low discrimination**						**27510020**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		36.92	0.41	NIT 1	90	
Actinomyces viscosus		19.58	0.27		76	
Actinomyces odontolyticus		17.72	0.34		10	
27510021	**Low discrimination**					**27510021**
Actinomyces naeslundii		62.64	0.47	GAL 96		
Actinomyces viscosus		37.36	0.19			
27510022	**Very good identification**					**27510022**
					CAT	
Actinomyces viscosus		98.31	0.62	NAG 1	76	
Actinomyces odontolyticus		0.55	0.30		10	
Propionibacterium granulosum		0.47	0.31		90	
27510023	**Good identification**					**27510023**
Actinomyces viscosus		98.34	0.53	NAG 1		
Actinomyces naeslundii		1.67	0.47			
27510042	**Very good identification**					**27510042**
Actinomyces viscosus		99.78	0.62	SOR 1		
Propionibacterium propionicum		0.12	0.26			
Actinomyces naeslundii		0.07	0.31			
27510043	**Good identification**					**27510043**
Actinomyces viscosus		99.70	0.53	SOR 1		
Actinomyces naeslundii		0.30	0.34			
27510100	**Good identification**					**27510100**
Actinomyces viscosus		93.48	0.66	BGL 99		
Propionibacterium propionicum		4.26	0.57			
Actinomyces odontolyticus		0.93	0.39			
27510101	**Good identification**					**27510101**
					SPO	
Actinomyces viscosus		94.37	0.58	BGL 99	1	
Actinomyces naeslundii		2.56	0.55		1	
Clostridium chauvoei		1.64	0.29		99	
27510102	**Excellent identification**					**27510102**
Actinomyces viscosus		99.92	1.00			
Propionibacterium propionicum		0.07	0.60			
Actinomyces naeslundii		0.01	0.52			
27510103	**Excellent identification**					**27510103**
Actinomyces viscosus		99.95	0.92			
Actinomyces naeslundii		0.03	0.55			
Propionibacterium propionicum		0.02	0.44			
27510112	**Very good identification**					**27510112**
Actinomyces viscosus		99.89	0.66	MLZ 1		
Propionibacterium propionicum		0.07	0.26			
Actinomyces naeslundii		0.03	0.30			
27510113	**Very good identification**					**27510113**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		99.86	0.58	MLZ 1			
Actinomyces naeslundii		0.14	0.33				
27510120	**Low discrimination**						**27510120**
					GRA		
Clostridium perfringens		38.93	0.40	GAL 99	99		
Prevotella melaninogenica		30.68	0.32		1		
Actinomyces viscosus		15.58	0.32		99		
27510121	**Low discrimination**						**27510121**
					SPO		
Actinomyces naeslundii		61.09	0.55	GAL 96	1		
Actinomyces viscosus		22.76	0.24		1		
Clostridium chauvoei		13.07	0.21		99		
27510122	**Very good identification**						**27510122**
					GRA	CAT	
Actinomyces viscosus		96.54	0.66	NAG 1	99	76	
Capnocytophaga ochracea		1.15	0.32		1	90	
Clostridium perfringens		1.15	0.35		99	1	
27510123	**Very good identification**						**27510123**
					SPO		
Actinomyces viscosus		97.20	0.58	NAG 1	1		
Actinomyces naeslundii		2.64	0.55		1		
Clostridium chauvoei		0.10	0.08		99		
27510140	**Low discrimination**						**27510140**
Propionibacterium propionicum		80.10	0.57	MAN 99			
Actinomyces viscosus		17.77	0.32				
Actinomyces naeslundii		1.87	0.39				
27510141	**Low discrimination**						**27510141**
Propionibacterium propionicum		50.49	0.41	MAN 99			
Actinomyces viscosus		33.60	0.24	URE 10			
Actinomyces naeslundii		15.91	0.42				
27510142	**Good identification**						**27510142**
Actinomyces viscosus		93.50	0.66	SOR 1			
Propionibacterium propionicum		6.39	0.60				
Actinomyces naeslundii		0.10	0.39				
27510143	**Very good identification**						**27510143**
Actinomyces viscosus		97.32	0.58	SOR 1			
Propionibacterium propionicum		2.22	0.44				
Actinomyces naeslundii		0.47	0.42				
27510161	**Good identification**						**27510161**
Actinomyces naeslundii		100.0	0.42	GAL 96			
27510200	**Low discrimination**						**27510200**
					CAT		
Actinomyces odontolyticus		88.58	0.52	MNS 9	10		
Actinomyces viscosus		8.21	0.27	RAF 3	76		
Propionibacterium granulosum		1.92	0.26		90		
27510202	**Good identification**						**27510202**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		93.69	0.62	RHA 1	76	
Actinomyces odontolyticus		6.26	0.48		10	
Propionibacterium granulosum		0.06	0.16		90	
27510203	**Good identification**					**27510203**
Actinomyces viscosus		99.80	0.53	RHA 1		
Actinomyces odontolyticus		0.20	0.14			
27510220	**Good identification**					**27510220**
					CAT	
Actinomyces odontolyticus		96.79	0.39	MNS 9	10	
Propionibacterium granulosum		3.21	0.16	RAF 3	90	
27510300	**Low discrimination**					**27510300**
Actinomyces odontolyticus		64.71	0.44	MNS 9		
Actinomyces viscosus		33.75	0.32	RAF 3		
Propionibacterium propionicum		1.54	0.23			
27510302	**Very good identification**					**27510302**
Actinomyces viscosus		98.76	0.66	RHA 1		
Actinomyces odontolyticus		1.17	0.40			
Propionibacterium propionicum		0.07	0.26			
27510303	**Very good identification**					**27510303**
Actinomyces viscosus		99.96	0.58	RHA 1		
Actinomyces odontolyticus		0.04	0.06			
27510402	**Very good identification**					**27510402**
Actinomyces viscosus		99.91	0.62	ARA 1		
Lactobacillus fermentum		0.06	0.20			
Actinomyces odontolyticus		0.04	0.09			
27510403	**Good identification**					**27510403**
Actinomyces viscosus		100.0	0.53	ARA 1		
27510502	**Very good identification**					**27510502**
Actinomyces viscosus		99.84	0.66	ARA 1		
Propionibacterium propionicum		0.07	0.26			
Lactobacillus fermentum		0.05	0.24			
27510503	**Very good identification**					**27510503**
Actinomyces viscosus		100.0	0.58	ARA 1		
27511000	**Low discrimination**					**27511000**
					GRA	
Actinomyces viscosus		57.64	0.58	BGL 99	99	
Prevotella intermedia		14.93	0.49		1	
Actinomyces naeslundii		8.49	0.68		99	
27511001	**Low discrimination**					**27511001**
					SPO	
Actinomyces viscosus		57.81	0.50	BGL 99	1	
Actinomyces naeslundii		38.34	0.71		1	
Clostridium chauvoei		2.92	0.29		99	
27511002	**Excellent identification**					**27511002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		99.67	0.92			
Actinomyces naeslundii		0.15	0.68			
Lactobacillus fermentum		0.09	0.54			
27511003	**Very good identification**					**27511003**
					SPO	
Actinomyces viscosus		99.32	0.84		1	
Actinomyces naeslundii		0.67	0.71		1	
Clostridium chauvoei		0.01	0.16		99	
27511010	**Low discrimination**					**27511010**
Propionibacterium granulosum		43.54	0.37	NIT 1		
Actinomyces viscosus		25.94	0.24	MLZ 12		
Actinomyces naeslundii		19.91	0.46			
27511011	**Low discrimination**					**27511011**
Actinomyces naeslundii		77.55	0.49	MLZ 5		
Actinomyces viscosus		22.45	0.16			
27511012	**Very good identification**					**27511012**
Actinomyces viscosus		98.65	0.58	MLZ 1		
Actinomyces naeslundii		0.77	0.46			
Propionibacterium granulosum		0.42	0.27			
27511013	**Good identification**					**27511013**
Actinomyces viscosus		96.63	0.50	MLZ 1		
Actinomyces naeslundii		3.37	0.49			
27511020	**Low discrimination**					**27511020**
					GRA	
Clostridium perfringens		77.08	0.60	LAC 88	99	
Actinomyces naeslundii		11.49	0.68		99	
Prevotella melaninogenica		4.50	0.32		1	
27511021	**Good identification**					**27511021**
					SPO	
Actinomyces naeslundii		91.88	0.71		1	
Clostridium perfringens		4.15	0.26		84	
Clostridium chauvoei		2.33	0.21		99	
27511022	**Low discrimination**					**27511022**
Actinomyces viscosus		58.55	0.58	NAG 1		
Clostridium perfringens		27.24	0.54			
Actinomyces naeslundii		8.63	0.68			
27511023	**Low discrimination**					**27511023**
Actinomyces viscosus		59.48	0.50	NAG 1		
Actinomyces naeslundii		39.44	0.71			
Clostridium perfringens		0.84	0.20			
27511031	**Good identification**					**27511031**
Actinomyces naeslundii		100.0	0.49	MLZ 5		
27511033	**Low discrimination**					**27511033**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		77.55	0.49	MLZ 5		
Actinomyces viscosus		22.45	0.16			
27511040	**Low discrimination**					**27511040**
Actinomyces naeslundii		53.45	0.55			
Actinomyces viscosus		20.76	0.24			
Clostridium perfringens		11.97	0.22			
27511041	**Good identification**					**27511041**
Actinomyces naeslundii		92.05	0.58			
Actinomyces viscosus		7.95	0.16			
27511042	**Very good identification**					**27511042**
Actinomyces viscosus		96.86	0.58	SOR 1		
Actinomyces naeslundii		2.52	0.55			
Clostridium perfringens		0.27	0.16			
27511043	**Low discrimination**					**27511043**
Actinomyces viscosus		89.52	0.50	SOR 1		
Actinomyces naeslundii		10.48	0.58			
27511051	**Good identification**					**27511051**
Actinomyces naeslundii		100.0	0.36	MLZ 5		
27511060	**Low discrimination**					**27511060**
					GRA	
Clostridium perfringens		87.45	0.47	LAC 88	99	
Actinomyces naeslundii		12.08	0.55		99	
Prevotella melaninogenica		0.27	-.03		1	
27511061	**Very good identification**					**27511061**
Actinomyces naeslundii		95.35	0.58			
Clostridium perfringens		4.65	0.13			
27511062	**Low discrimination**					**27511062**
Clostridium perfringens		68.92	0.42	LAC 88		
Actinomyces naeslundii		20.23	0.55			
Actinomyces viscosus		7.86	0.24			
27511063	**Good identification**					**27511063**
Actinomyces naeslundii		90.15	0.58			
Actinomyces viscosus		7.78	0.16			
Clostridium perfringens		2.07	0.08			
27511071	**Good identification**					**27511071**
Actinomyces naeslundii		100.0	0.36	MLZ 5		
27511073	**Good identification**					**27511073**
Actinomyces naeslundii		100.0	0.36	MLZ 5		
27511100	**Low discrimination**					**27511100**
Actinomyces viscosus		62.12	0.62	BGL 99		
Actinomyces naeslundii		14.65	0.76			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		7.83	0.49				
27511101	**Low discrimination**						**27511101**
					SPO		
Clostridium chauvoei		57.30	0.63	RAF 1	99		
Actinomyces naeslundii		21.64	0.79		1		
Actinomyces viscosus		20.39	0.54		1		
27511102	**Excellent identification**						**27511102**
					GRA		
Actinomyces viscosus		99.28	0.96		99		
Actinomyces naeslundii		0.24	0.76		99		
Capnocytophaga ochracea		0.22	0.50		1		
27511103	**Very good identification**						**27511103**
					SPO		
Actinomyces viscosus		98.40	0.88		1		
Actinomyces naeslundii		1.06	0.79		1		
Clostridium chauvoei		0.49	0.50		99		
27511110	**Low discrimination**						**27511110**
					GRA		
Actinomyces naeslundii		43.00	0.54	MLZ 5	99		
Actinomyces viscosus		34.98	0.28		99		
Capnocytophaga ochracea		9.17	0.17		1		
27511111	**Low discrimination**						**27511111**
					SPO		
Actinomyces naeslundii		59.21	0.57	MLZ 5	1		
Clostridium chauvoei		30.09	0.29		99		
Actinomyces viscosus		10.71	0.20		1		
27511112	**Very good identification**						**27511112**
					GRA		
Actinomyces viscosus		96.19	0.62	MLZ 1	99		
Capnocytophaga ochracea		2.29	0.33		1		
Actinomyces naeslundii		1.19	0.54		99		
27511113	**Acceptable identification**						**27511113**
					SPO		
Actinomyces viscosus		94.20	0.54	MLZ 1	1		
Actinomyces naeslundii		5.26	0.57		1		
Clostridium chauvoei		0.47	0.16		99		
27511120	**Low discrimination**						**27511120**
					GRA		
Clostridium perfringens		52.35	0.75		99		
Prevotella melaninogenica		41.26	0.66		1		
Actinomyces naeslundii		3.03	0.76		99		
27511121	**Low discrimination**						**27511121**
					SPO		
Actinomyces naeslundii		47.46	0.79		1		
Clostridium chauvoei		41.88	0.55		99		
Clostridium perfringens		5.52	0.40		84		
27511122	**Low discrimination**						**27511122**
					GRA	CAT	
Capnocytophaga ochracea		33.35	0.66	ESL 99	1	90	
Clostridium perfringens		33.15	0.69		99	1	
Actinomyces viscosus		17.28	0.62		99	76	
27511123	**Low discrimination**						**27511123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		44.89	0.79		1		
Actinomyces viscosus		42.29	0.54		1		
Clostridium chauvoei		6.99	0.42		99		
27511130	**Low discrimination**						**27511130**
					GRA	CAT	
Clostridium perfringens		36.68	0.40	MLZ 1	99	1	
Prevotella melaninogenica		28.91	0.32		1	1	
Capnocytophaga ochracea		21.22	0.33		1	90	
27511131	**Low discrimination**						**27511131**
					SPO		
Actinomyces naeslundii		81.82	0.57	MLZ 5	1		
Clostridium chauvoei		13.86	0.21		99		
Clostridium perfringens		1.83	0.06		84		
27511132	**Low discrimination**						**27511132**
					GRA	CAT	
Capnocytophaga ochracea		80.58	0.50	ESL 99	1	90	
Clostridium perfringens		7.28	0.35	MLZ 10	99	1	
Actinomyces naeslundii		4.66	0.54		99	25	
27511133	**Low discrimination**						**27511133**
					GRA		
Actinomyces naeslundii		74.77	0.57	MLZ 5	99		
Actinomyces viscosus		13.52	0.20		99		
Capnocytophaga ochracea		8.70	0.15		1		
27511140	**Low discrimination**						**27511140**
Propionibacterium propionicum		48.29	0.57	MAN 99			
Actinomyces naeslundii		27.06	0.63				
Clostridium perfringens		15.61	0.36				
27511141	**Low discrimination**						**27511141**
					SPO		
Actinomyces naeslundii		74.54	0.66		1		
Clostridium chauvoei		11.30	0.29		99		
Propionibacterium propionicum		9.85	0.41		1		
27511142	**Low discrimination**						**27511142**
Actinomyces viscosus		84.10	0.62	SOR 1			
Propionibacterium propionicum		9.37	0.60				
Actinomyces naeslundii		3.50	0.63				
27511143	**Low discrimination**						**27511143**
Actinomyces viscosus		81.29	0.54	SOR 1			
Actinomyces naeslundii		15.23	0.66				
Propionibacterium propionicum		3.02	0.44				
27511150	**Low discrimination**						**27511150**
					GRA	CAT	
Actinomyces naeslundii		62.99	0.41	MLZ 5	99	25	
Propionibacterium propionicum		21.58	0.23		99	1	
Capnocytophaga ochracea		8.46	0.01		1	90	
27511151	**Good identification**						**27511151**
Actinomyces naeslundii		100.0	0.44	MLZ 5			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27511160	**Low discrimination**						**27511160**
					GRA		
Clostridium perfringens		88.70	0.62		99		
Actinomyces naeslundii		4.75	0.63		99		
Prevotella melaninogenica		3.71	0.32		1		
27511161	**Low discrimination**						**27511161**
					SPO		
Actinomyces naeslundii		84.37	0.66		1		
Clostridium perfringens		10.60	0.28		84		
Clostridium chauvoei		4.26	0.21		99		
27511162	**Low discrimination**						**27511162**
					GRA	CAT	
Clostridium perfringens		57.12	0.57		99	1	
Capnocytophaga ochracea		33.52	0.50		1	90	
Actinomyces naeslundii		6.51	0.63		99	25	
27511163	**Low discrimination**						**27511163**
Actinomyces naeslundii		86.46	0.66				
Clostridium perfringens		5.11	0.22				
Actinomyces viscosus		4.66	0.20				
27511171	**Good identification**						**27511171**
Actinomyces naeslundii		100.0	0.44	MLZ 5			
27511173	**Acceptable identification**						**27511173**
					GRA		
Actinomyces naeslundii		93.18	0.44	MLZ 5	99		
Capnocytophaga ochracea		6.82	-.01		1		
27511200	**Low discrimination**						**27511200**
					CAT		
Actinomyces odontolyticus		86.22	0.49	MNS 9	10		
Actinomyces viscosus		7.35	0.24	RAF 3	76		
Propionibacterium granulosum		2.80	0.26		90		
27511202	**Good identification**						**27511202**
Actinomyces viscosus		92.94	0.58	RHA 1			
Actinomyces odontolyticus		6.75	0.45				
Actinomyces naeslundii		0.14	0.34				
27511203	**Good identification**						**27511203**
Actinomyces viscosus		99.12	0.50	RHA 1			
Actinomyces naeslundii		0.66	0.37				
Actinomyces odontolyticus		0.22	0.11				
27511220	**Low discrimination**						**27511220**
Actinomyces odontolyticus		59.72	0.36	MNS 9			
Clostridium perfringens		30.89	0.26	RAF 3			
Actinomyces naeslundii		4.61	0.34	NAG 14			
27511221	**Good identification**						**27511221**
Actinomyces naeslundii		100.0	0.37	RHA 1			
27511300	**Low discrimination**						**27511300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		56.98	0.40	MNS 9		
Actinomyces viscosus		27.33	0.28	RAF 3		
Actinomyces naeslundii		6.45	0.42			
27511302	**Very good identification**					**27511302**
Actinomyces viscosus		97.92	0.62	RHA 1		
Actinomyces odontolyticus		1.26	0.37			
Actinomyces naeslundii		0.23	0.42			
27511303	**Good identification**					**27511303**
					SPO	
Actinomyces viscosus		98.45	0.54	RHA 1	1	
Actinomyces naeslundii		1.06	0.45		1	
Clostridium chauvoei		0.49	0.16		99	
27511320	**Low discrimination**					**27511320**
					GRA	
Clostridium perfringens		47.89	0.40	RHA 1	99	
Prevotella melaninogenica		37.75	0.32		1	
Actinomyces odontolyticus		3.99	0.27		99	
27511321	**Low discrimination**					**27511321**
					SPO	
Actinomyces naeslundii		47.83	0.45	RHA 1	1	
Clostridium chauvoei		42.21	0.21		99	
Clostridium perfringens		5.57	0.06		84	
27511323	**Low discrimination**					**27511323**
					SPO	
Actinomyces naeslundii		43.94	0.45	RHA 1	1	
Actinomyces viscosus		41.40	0.20		1	
Clostridium chauvoei		6.84	0.08		99	
27511400	**Low discrimination**					**27511400**
					GRA	
Lactobacillus fermentum		83.23	0.54	NIT 1	99	
Actinomyces viscosus		9.45	0.24		99	
Prevotella intermedia		2.45	0.15		1	
27511402	**Good identification**					**27511402**
Actinomyces viscosus		91.57	0.58	ARA 1		
Lactobacillus fermentum		8.14	0.54			
Actinomyces naeslundii		0.14	0.34			
27511403	**Good identification**					**27511403**
Actinomyces viscosus		99.07	0.50	ARA 1		
Actinomyces naeslundii		0.66	0.37			
Lactobacillus fermentum		0.27	0.20			
27511420	**Low discrimination**					**27511420**
Lactobacillus fermentum		66.46	0.46	NIT 1		
Clostridium perfringens		22.39	0.26			
Bifidobacterium longum subsp. longum		4.99	0.20			
27511421	**Good identification**					**27511421**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27511422	**Low discrimination**					**27511422**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Lactobacillus fermentum		54.62	0.46	NIT 1	99	1	
Actinomyces viscosus		18.61	0.24		99	76	
Capnocytophaga ochracea		11.27	0.19		1	90	
27511500	**Low discrimination**						**27511500**
Lactobacillus fermentum		62.01	0.58	NIT 1			
Bifidobacterium longum subsp. longum		23.53	0.43				
Actinomyces viscosus		7.67	0.28				
27511502	**Low discrimination**						**27511502**
					GRA	CAT	
Actinomyces viscosus		86.52	0.62	ARA 1	99	76	
Lactobacillus fermentum		7.06	0.58		99	1	
Capnocytophaga ochracea		3.28	0.37		1	90	
27511503	**Good identification**						**27511503**
					SPO		
Actinomyces viscosus		98.01	0.54	ARA 1	1		
Actinomyces naeslundii		1.05	0.45		1		
Clostridium chauvoei		0.49	0.16		99		
27511510	**Good identification**						**27511510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.61	0.47	NIT 1	99	1	
Lactobacillus fermentum		1.53	0.24		99	1	
Capnocytophaga ochracea		0.87	0.04		1	90	
27511512	**Low discrimination**						**27511512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		75.93	0.47	NIT 1	99	1	
Actinomyces viscosus		14.55	0.28		99	76	
Capnocytophaga ochracea		6.06	0.20		1	90	
27511520	**Low discrimination**						**27511520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		46.02	0.54	NIT 1	99	1	
Clostridium perfringens		15.31	0.40		99	1	
Capnocytophaga ochracea		14.07	0.37		1	90	
27511521	**Low discrimination**						**27511521**
					SPO		
Actinomyces naeslundii		38.10	0.45	ARA 1	1		
Clostridium chauvoei		33.63	0.21		99		
Bifidobacterium longum subsp. longum		13.34	0.20		1		
27511522	**Low discrimination**						**27511522**
					GRA	CAT	
Capnocytophaga ochracea		64.15	0.53	ESL 99	1	90	
Bifidobacterium longum subsp. longum		23.32	0.54		99	1	
Lactobacillus fermentum		5.12	0.50		99	1	
27511523	**Low discrimination**						**27511523**
					GRA		
Actinomyces naeslundii		28.23	0.45	ARA 1	99		
Capnocytophaga ochracea		27.18	0.19		1		
Actinomyces viscosus		26.59	0.20		99		
27511530	**Very good identification**						**27511530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.58	0.58	NIT 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		1.95	0.20		1	90	
Clostridium perfringens		0.19	0.06		99	1	
27511532	**Low discrimination**						**27511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		84.49	0.58	NIT 1	99	1	
Capnocytophaga ochracea		15.17	0.37		1	90	
Parabacteroides distasonis		0.15	0.15		1	76	
27511562	**Low discrimination**						**27511562**
					GRA	CAT	
Capnocytophaga ochracea		87.04	0.37	ESL 99	1	90	
Clostridium perfringens		8.49	0.22	SOR 10	99	1	
Bifidobacterium longum subsp. longum		2.88	0.20		99	1	
27512000	**Low discrimination**						**27512000**
					CAT		
Propionibacterium propionicum		51.23	0.57	LAC 99	1		
Propionibacterium granulosum		36.38	0.50		90		
Actinomyces viscosus		6.39	0.27		76		
27512001	**Low discrimination**						**27512001**
Actinomyces naeslundii		45.85	0.52	GAL 96			
Propionibacterium propionicum		37.64	0.41				
Actinomyces viscosus		14.09	0.19				
27512002	**Low discrimination**						**27512002**
					CAT		
Actinomyces viscosus		87.45	0.62	MAN 1	76		
Propionibacterium propionicum		10.62	0.60		1		
Propionibacterium granulosum		1.26	0.39		90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27512003	**Acceptable identification**					**27512003**
Actinomyces viscosus		93.17	0.53	MAN 1		
Propionibacterium propionicum		3.77	0.44			
Actinomyces naeslundii		3.06	0.52			
27512020	**Low discrimination**					**27512020**
					CAT	
Propionibacterium granulosum		63.89	0.39	NIT 1	90	
Actinomyces naeslundii		32.47	0.49		25	
Propionibacterium propionicum		3.64	0.23		1	
27512021	**Good identification**					**27512021**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27512023	**Low discrimination**					**27512023**
Actinomyces naeslundii		76.49	0.52	GAL 96		
Actinomyces viscosus		23.51	0.19			
27512040	**Very good identification**					**27512040**
					CAT	
Propionibacterium propionicum		97.74	0.57	LAC 99	1	
Actinomyces naeslundii		1.56	0.36		25	
Propionibacterium granulosum		0.70	0.16		90	
27512041	**Low discrimination**					**27512041**
Propionibacterium propionicum		82.31	0.41	LAC 99		
Actinomyces naeslundii		17.69	0.39	URE 10		
27512042	**Good identification**					**27512042**
Propionibacterium propionicum		91.43	0.60	LAC 99		
Actinomyces viscosus		7.61	0.27			
Actinomyces naeslundii		0.97	0.36			
27512043	**Low discrimination**					**27512043**
Propionibacterium propionicum		71.80	0.44	LAC 99		
Actinomyces viscosus		17.92	0.19	URE 10		
Actinomyces naeslundii		10.29	0.39			
27512060	**Low discrimination**					**27512060**
Actinomyces naeslundii		61.19	0.36	GAL 96		
Propionibacterium propionicum		38.81	0.23			
27512061	**Good identification**					**27512061**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27512062	**Low discrimination**					**27512062**
Actinomyces naeslundii		51.24	0.36	GAL 96		
Propionibacterium propionicum		48.76	0.26			
27512063	**Good identification**					**27512063**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27512100	**Very good identification**					**27512100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		99.51	0.91			
Actinomyces naeslundii		0.26	0.57			
Actinomyces viscosus		0.22	0.32			
27512101	**Very good identification**					**27512101**
Propionibacterium propionicum		95.99	0.75	URE 10		
Actinomyces naeslundii		3.36	0.60			
Actinomyces viscosus		0.65	0.24			
27512102	**Low discrimination**					**27512102**
Propionibacterium propionicum		86.99	0.94			
Actinomyces viscosus		12.86	0.66			
Actinomyces naeslundii		0.15	0.57			
27512103	**Low discrimination**					**27512103**
Propionibacterium propionicum		68.17	0.78	URE 10		
Actinomyces viscosus		30.24	0.58			
Actinomyces naeslundii		1.59	0.60			
27512110	**Very good identification**					**27512110**
Propionibacterium propionicum		98.67	0.57	MLZ 1		
Actinomyces naeslundii		1.33	0.35			
27512111	**Low discrimination**					**27512111**
Propionibacterium propionicum		84.57	0.41	MLZ 1		
Actinomyces naeslundii		15.43	0.38	URE 10		
27512112	**Low discrimination**					**27512112**
Propionibacterium propionicum		86.44	0.60	MLZ 1		
Actinomyces viscosus		12.78	0.32			
Actinomyces naeslundii		0.78	0.35			
27512113	**Low discrimination**					**27512113**
Propionibacterium propionicum		63.89	0.44	MLZ 1		
Actinomyces viscosus		28.34	0.24	URE 10		
Actinomyces naeslundii		7.77	0.38			
27512120	**Low discrimination**					**27512120**
Propionibacterium propionicum		78.94	0.57	NAG 1		
Actinomyces naeslundii		20.27	0.57			
Clostridium perfringens		0.44	0.06			
27512121	**Low discrimination**					**27512121**
Actinomyces naeslundii		77.61	0.60	GAL 96		
Propionibacterium propionicum		22.39	0.41			
27512122	**Low discrimination**					**27512122**
Propionibacterium propionicum		75.61	0.60	NAG 1		
Actinomyces naeslundii		12.95	0.57			
Actinomyces viscosus		11.18	0.32			
27512123	**Low discrimination**					**27512123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		61.55	0.60	GAL 96		
Propionibacterium propionicum		26.63	0.44			
Actinomyces viscosus		11.82	0.24			
27512130	**Low discrimination**					**27512130**
Actinomyces naeslundii		57.23	0.35	GAL 96		
Propionibacterium propionicum		42.77	0.23	MLZ 5		
27512131	**Good identification**					**27512131**
Actinomyces naeslundii		100.0	0.38	GAL 96		
27512133	**Good identification**					**27512133**
Actinomyces naeslundii		100.0	0.38	GAL 96		
27512140	**Very good identification**					**27512140**
Propionibacterium propionicum		99.95	0.91			
Actinomyces naeslundii		0.05	0.44			
27512141	**Very good identification**					**27512141**
Propionibacterium propionicum		99.39	0.75	URE 10		
Actinomyces naeslundii		0.61	0.47			
27512142	**Excellent identification**					**27512142**
Propionibacterium propionicum		99.82	0.94			
Actinomyces viscosus		0.15	0.32			
Actinomyces naeslundii		0.03	0.44			
27512143	**Very good identification**					**27512143**
Propionibacterium propionicum		99.15	0.78	URE 10		
Actinomyces viscosus		0.44	0.24			
Actinomyces naeslundii		0.41	0.47			
27512150	**Very good identification**					**27512150**
Propionibacterium propionicum		100.0	0.57	MLZ 1		
27512151	**Good identification**					**27512151**
Propionibacterium propionicum		96.88	0.41	MLZ 1		
Actinomyces naeslundii		3.12	0.25	URE 10		
27512152	**Very good identification**					**27512152**
Propionibacterium propionicum		100.0	0.60	MLZ 1		
27512153	**Good identification**					**27512153**
Propionibacterium propionicum		97.90	0.44	MLZ 1		
Actinomyces naeslundii		2.10	0.25	URE 10		
27512160	**Very good identification**					**27512160**
Propionibacterium propionicum		95.66	0.57	NAG 1		
Actinomyces naeslundii		4.34	0.44			
27512161	**Low discrimination**					**27512161**
Propionibacterium propionicum		62.04	0.41	NAG 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		37.96	0.47	URE 10		
27512162	**Very good identification**					**27512162**
Propionibacterium propionicum		97.07	0.60	NAG 1		
Actinomyces naeslundii		2.93	0.44			
27512163	**Low discrimination**					**27512163**
Propionibacterium propionicum		71.03	0.44	NAG 1		
Actinomyces naeslundii		28.97	0.47	URE 10		
27512300	**Very good identification**					**27512300**
Propionibacterium propionicum		99.32	0.57	RHA 1		
Actinomyces odontolyticus		0.43	0.09			
Actinomyces naeslundii		0.26	0.23			
27512301	**Good identification**					**27512301**
Propionibacterium propionicum		96.62	0.41	RHA 1		
Actinomyces naeslundii		3.38	0.26	URE 10		
27512302	**Low discrimination**					**27512302**
Propionibacterium propionicum		86.85	0.60	RHA 1		
Actinomyces viscosus		12.84	0.32			
Actinomyces naeslundii		0.15	0.23			
27512303	**Low discrimination**					**27512303**
Propionibacterium propionicum		68.17	0.44	RHA 1		
Actinomyces viscosus		30.24	0.24	URE 10		
Actinomyces naeslundii		1.59	0.26			
27512340	**Very good identification**					**27512340**
Propionibacterium propionicum		100.0	0.57	RHA 1		
27512341	**Good identification**					**27512341**
Propionibacterium propionicum		100.0	0.41	RHA 1		
27512342	**Very good identification**					**27512342**
Propionibacterium propionicum		100.0	0.60	RHA 1		
27512343	**Good identification**					**27512343**
Propionibacterium propionicum		100.0	0.44	RHA 1		
27512500	**Very good identification**					**27512500**
Propionibacterium propionicum		99.74	0.57	ARA 1		
Actinomyces naeslundii		0.26	0.23			
27512501	**Good identification**					**27512501**
Propionibacterium propionicum		96.62	0.41	ARA 1		
Actinomyces naeslundii		3.38	0.26	URE 10		
27512502	**Low discrimination**					**27512502**
Propionibacterium propionicum		86.99	0.60	ARA 1		
Actinomyces viscosus		12.86	0.32			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.15	0.23			
27512503	**Low discrimination**					**27512503**
Propionibacterium propionicum		68.17	0.44	ARA 1		
Actinomyces viscosus		30.24	0.24	URE 10		
Actinomyces naeslundii		1.59	0.26			
27512540	**Very good identification**					**27512540**
Propionibacterium propionicum		100.0	0.57	ARA 1		
27512541	**Good identification**					**27512541**
Propionibacterium propionicum		100.0	0.41	ARA 1		
27512542	**Very good identification**					**27512542**
Propionibacterium propionicum		100.0	0.60	ARA 1		
27512543	**Good identification**					**27512543**
Propionibacterium propionicum		100.0	0.44	ARA 1		
27513000	**Low discrimination**					**27513000**
					CAT	
Actinomyces naeslundii		54.36	0.73		25	
Propionibacterium propionicum		25.10	0.57		1	
Propionibacterium granulosum		17.83	0.50		90	
27513001	**Very good identification**					**27513001**
Actinomyces naeslundii		95.76	0.76			
Propionibacterium propionicum		3.28	0.41			
Actinomyces viscosus		0.75	0.16			
27513002	**Low discrimination**					**27513002**
Actinomyces viscosus		66.26	0.58	MAN 1		
Actinomyces naeslundii		18.95	0.73			
Propionibacterium propionicum		13.13	0.60			
27513003	**Low discrimination**					**27513003**
Actinomyces naeslundii		54.70	0.76			
Actinomyces viscosus		42.50	0.50			
Propionibacterium propionicum		2.81	0.44			
27513010	**Low discrimination**					**27513010**
					CAT	
Actinomyces naeslundii		51.59	0.51	MLZ 5	25	
Propionibacterium granulosum		43.84	0.35		90	
Propionibacterium propionicum		4.57	0.23		1	
27513011	**Good identification**					**27513011**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
27513012	**Low discrimination**					**27513012**
Actinomyces naeslundii		49.60	0.51	MLZ 5		
Actinomyces viscosus		33.28	0.24			
Propionibacterium granulosum		10.53	0.25			
27513013	**Low discrimination**					**27513013**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		87.02	0.54	MLZ 5		
Actinomyces viscosus		12.98	0.16			
27513020	**Low discrimination**					**27513020**
					CAT	
Actinomyces naeslundii		89.00	0.73		25	
Propionibacterium granulosum		7.30	0.39		90	
Clostridium perfringens		3.11	0.26		1	
27513021	**Very good identification**					**27513021**
Actinomyces naeslundii		100.0	0.76			
27513022	**Good identification**					**27513022**
Actinomyces naeslundii		92.55	0.73			
Actinomyces viscosus		3.27	0.24			
Propionibacterium granulosum		1.90	0.29			
27513023	**Very good identification**					**27513023**
Actinomyces naeslundii		99.22	0.76			
Actinomyces viscosus		0.78	0.16			
27513030	**Low discrimination**					**27513030**
Actinomyces naeslundii		82.48	0.51	MLZ 5		
Propionibacterium granulosum		17.52	0.25			
27513031	**Good identification**					**27513031**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
27513032	**Acceptable identification**					**27513032**
Actinomyces naeslundii		94.96	0.51	MLZ 5		
Propionibacterium granulosum		5.04	0.14			
27513033	**Good identification**					**27513033**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
27513040	**Low discrimination**					**27513040**
					CAT	
Propionibacterium propionicum		71.98	0.57	LAC 99	1	
Actinomyces naeslundii		27.51	0.60		25	
Propionibacterium granulosum		0.52	0.16		90	
27513041	**Low discrimination**					**27513041**
Actinomyces naeslundii		83.76	0.63			
Propionibacterium propionicum		16.24	0.41			
27513042	**Low discrimination**					**27513042**
Propionibacterium propionicum		76.58	0.60	LAC 99		
Actinomyces naeslundii		19.51	0.60			
Actinomyces viscosus		3.90	0.24			
27513043	**Low discrimination**					**27513043**
Actinomyces naeslundii		74.89	0.63			
Propionibacterium propionicum		21.78	0.44			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		3.33	0.16			
27513050	**Low discrimination**					**27513050**
Actinomyces naeslundii		66.57	0.38	MLZ 5		
Propionibacterium propionicum		33.43	0.23			
27513051	**Good identification**					**27513051**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27513052	**Low discrimination**					**27513052**
Actinomyces naeslundii		57.04	0.38	MLZ 5		
Propionibacterium propionicum		42.96	0.26			
27513053	**Good identification**					**27513053**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27513060	**Good identification**					**27513060**
Actinomyces naeslundii		93.98	0.60			
Clostridium perfringens		3.54	0.13			
Propionibacterium propionicum		2.48	0.23			
27513061	**Very good identification**					**27513061**
Actinomyces naeslundii		100.0	0.63			
27513062	**Good identification**					**27513062**
Actinomyces naeslundii		94.57	0.60			
Propionibacterium propionicum		3.75	0.26			
Clostridium perfringens		1.68	0.08			
27513063	**Very good identification**					**27513063**
Actinomyces naeslundii		100.0	0.63			
27513070	**Good identification**					**27513070**
Actinomyces naeslundii		100.0	0.38	MLZ 5		
27513071	**Good identification**					**27513071**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27513072	**Good identification**					**27513072**
Actinomyces naeslundii		100.0	0.38	MLZ 5		
27513073	**Good identification**					**27513073**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27513100	**Good identification**					**27513100**
Propionibacterium propionicum		93.97	0.91			
Actinomyces naeslundii		5.85	0.81			
Actinomyces viscosus		0.13	0.28			
27513101	**Low discrimination**					**27513101**
					SPO	
Propionibacterium propionicum		53.87	0.75	URE 10	1	
Actinomyces naeslundii		45.28	0.84		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		0.62	0.29		99		
27513102	**Low discrimination**						**27513102**
Propionibacterium propionicum		88.30	0.94				
Actinomyces viscosus		8.00	0.62				
Actinomyces naeslundii		3.67	0.81				
27513103	**Low discrimination**						**27513103**
Propionibacterium propionicum		54.54	0.78	URE 10			
Actinomyces naeslundii		30.56	0.84				
Actinomyces viscosus		14.83	0.54				
27513110	**Low discrimination**						**27513110**
Propionibacterium propionicum		75.51	0.57	MLZ 1			
Actinomyces naeslundii		24.49	0.59				
27513111	**Low discrimination**						**27513111**
Actinomyces naeslundii		81.41	0.62	MLZ 5			
Propionibacterium propionicum		18.59	0.41				
27513112	**Low discrimination**						**27513112**
Propionibacterium propionicum		76.29	0.60	MLZ 1			
Actinomyces naeslundii		16.50	0.59				
Actinomyces viscosus		6.92	0.28				
27513113	**Low discrimination**						**27513113**
Actinomyces naeslundii		69.66	0.62	MLZ 5			
Propionibacterium propionicum		23.86	0.44				
Actinomyces viscosus		6.49	0.20				
27513120	**Low discrimination**						**27513120**
Actinomyces naeslundii		75.28	0.81				
Propionibacterium propionicum		12.21	0.57				
Clostridium perfringens		6.77	0.40				
27513121	**Very good identification**						**27513121**
					SPO		
Actinomyces naeslundii		98.26	0.84		1		
Propionibacterium propionicum		1.18	0.41		1		
Clostridium chauvoei		0.45	0.21		99		
27513122	**Low discrimination**						**27513122**
					GRA	CAT	
Actinomyces naeslundii		73.22	0.81		99	25	
Propionibacterium propionicum		17.82	0.60		99	1	
Capnocytophaga ochracea		3.12	0.32		1	90	
27513123	**Very good identification**						**27513123**
Actinomyces naeslundii		97.63	0.84				
Propionibacterium propionicum		1.76	0.44				
Actinomyces viscosus		0.48	0.20				
27513130	**Good identification**						**27513130**
Actinomyces naeslundii		93.28	0.59	MLZ 5			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		2.91	0.23				
Clostridium perfringens		1.61	0.06				
27513131	**Very good identification**						**27513131**
Actinomyces naeslundii		100.0	0.62	MLZ 5			
27513132	**Low discrimination**						**27513132**
					GRA	CAT	
Actinomyces naeslundii		87.36	0.59	MLZ 5	99	25	
Capnocytophaga ochracea		7.85	0.15		1	90	
Propionibacterium propionicum		4.08	0.26		99	1	
27513133	**Very good identification**						**27513133**
Actinomyces naeslundii		100.0	0.62	MLZ 5			
27513140	**Very good identification**						**27513140**
Propionibacterium propionicum		98.91	0.91				
Actinomyces naeslundii		1.09	0.68				
Actinomyces israelii		0.00	0.15				
27513141	**Low discrimination**						**27513141**
Propionibacterium propionicum		87.08	0.75	URE 10			
Actinomyces naeslundii		12.92	0.71				
27513142	**Excellent identification**						**27513142**
Propionibacterium propionicum		99.18	0.94				
Actinomyces naeslundii		0.73	0.68				
Actinomyces viscosus		0.09	0.28				
27513143	**Good identification**						**27513143**
Propionibacterium propionicum		90.78	0.78	URE 10			
Actinomyces naeslundii		8.98	0.71				
Actinomyces viscosus		0.25	0.20				
27513150	**Good identification**						**27513150**
Propionibacterium propionicum		94.59	0.57	MLZ 1			
Actinomyces naeslundii		5.42	0.46				
27513151	**Low discrimination**						**27513151**
Propionibacterium propionicum		56.41	0.41	MLZ 1			
Actinomyces naeslundii		43.59	0.49	URE 10			
27513152	**Very good identification**						**27513152**
Propionibacterium propionicum		96.32	0.60	MLZ 1			
Actinomyces naeslundii		3.68	0.46				
27513153	**Low discrimination**						**27513153**
Propionibacterium propionicum		66.00	0.44	MLZ 1			
Actinomyces naeslundii		34.00	0.49	URE 10			
27513160	**Low discrimination**						**27513160**
Actinomyces naeslundii		49.42	0.68				
Propionibacterium propionicum		45.44	0.57				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		4.80	0.28				
27513161	**Good identification**						**27513161**
Actinomyces naeslundii		93.62	0.71				
Propionibacterium propionicum		6.38	0.41				
27513162	**Low discrimination**						**27513162**
Propionibacterium propionicum		56.25	0.60	NAG 1			
Actinomyces naeslundii		40.79	0.68				
Clostridium perfringens		1.86	0.22				
27513163	**Good identification**						**27513163**
Actinomyces naeslundii		90.73	0.71				
Propionibacterium propionicum		9.27	0.44				
27513170	**Low discrimination**						**27513170**
Actinomyces naeslundii		85.00	0.46	MLZ 5			
Propionibacterium propionicum		15.00	0.23				
27513171	**Good identification**						**27513171**
Actinomyces naeslundii		100.0	0.49	MLZ 5			
27513172	**Low discrimination**						**27513172**
					GRA	CAT	
Actinomyces naeslundii		75.69	0.46	MLZ 5	99	25	
Propionibacterium propionicum		20.03	0.26		99	1	
Capnocytophaga ochracea		4.28	-.01		1	90	
27513173	**Good identification**						**27513173**
Actinomyces naeslundii		100.0	0.49	MLZ 5			
27513201	**Good identification**						**27513201**
Actinomyces naeslundii		100.0	0.42	RHA 1			
27513203	**Low discrimination**						**27513203**
Actinomyces naeslundii		56.28	0.42	RHA 1			
Actinomyces viscosus		43.72	0.16				
27513220	**Low discrimination**						**27513220**
Actinomyces naeslundii		79.94	0.39	RHA 1			
Propionibacterium granulosum		20.06	0.14				
27513221	**Good identification**						**27513221**
Actinomyces naeslundii		100.0	0.42	RHA 1			
27513222	**Good identification**						**27513222**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27513223	**Good identification**						**27513223**
Actinomyces naeslundii		100.0	0.42	RHA 1			
27513300	**Good identification**						**27513300**
Propionibacterium propionicum		93.89	0.57	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		5.85	0.46			
Actinomyces odontolyticus		0.27	0.06			
27513301	**Low discrimination**					**27513301**
Propionibacterium propionicum		54.33	0.41	RHA 1		
Actinomyces naeslundii		45.67	0.49	URE 10		
27513302	**Low discrimination**					**27513302**
Propionibacterium propionicum		88.33	0.60	RHA 1		
Actinomyces viscosus		8.01	0.28			
Actinomyces naeslundii		3.67	0.46			
27513303	**Low discrimination**					**27513303**
Propionibacterium propionicum		54.58	0.44	RHA 1		
Actinomyces naeslundii		30.58	0.49	URE 10		
Actinomyces viscosus		14.84	0.20			
27513320	**Low discrimination**					**27513320**
Actinomyces naeslundii		75.58	0.46	RHA 1		
Propionibacterium propionicum		12.26	0.23			
Clostridium perfringens		6.80	0.06			
27513321	**Good identification**					**27513321**
Actinomyces naeslundii		100.0	0.49	RHA 1		
27513322	**Low discrimination**					**27513322**
					GRA	
Actinomyces naeslundii		72.42	0.46	RHA 1	99	
Propionibacterium propionicum		17.62	0.26		99	
Prevotella oralis		3.81	-.01		1	
27513323	**Good identification**					**27513323**
Actinomyces naeslundii		100.0	0.49	RHA 1		
27513340	**Very good identification**					**27513340**
Propionibacterium propionicum		98.91	0.57	RHA 1		
Actinomyces naeslundii		1.09	0.34			
27513341	**Low discrimination**					**27513341**
Propionibacterium propionicum		87.08	0.41	RHA 1		
Actinomyces naeslundii		12.92	0.37	URE 10		
27513342	**Very good identification**					**27513342**
Propionibacterium propionicum		99.27	0.60	RHA 1		
Actinomyces naeslundii		0.73	0.34			
27513343	**Acceptable identification**					**27513343**
Propionibacterium propionicum		91.00	0.44	RHA 1		
Actinomyces naeslundii		9.00	0.37	URE 10		
27513361	**Good identification**					**27513361**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27513363	**Good identification**					**27513363**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	RHA 1		
27513400	**Low discrimination**					**27513400**
					CAT	
Actinomyces naeslundii		47.61	0.39	ARA 1	25	
Propionibacterium propionicum		21.98	0.23		1	
Propionibacterium granulosum		15.61	0.16		90	
27513401	**Good identification**					**27513401**
Actinomyces naeslundii		100.0	0.42	ARA 1		
27513403	**Low discrimination**					**27513403**
Actinomyces naeslundii		56.28	0.42	ARA 1		
Actinomyces viscosus		43.72	0.16			
27513420	**Good identification**					**27513420**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27513421	**Good identification**					**27513421**
Actinomyces naeslundii		100.0	0.42	ARA 1		
27513422	**Good identification**					**27513422**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27513423	**Good identification**					**27513423**
Actinomyces naeslundii		100.0	0.42	ARA 1		
27513500	**Good identification**					**27513500**
Propionibacterium propionicum		92.26	0.57	ARA 1		
Actinomyces naeslundii		5.74	0.46			
Lactobacillus fermentum		1.02	0.24			
27513501	**Low discrimination**					**27513501**
Propionibacterium propionicum		54.33	0.41	ARA 1		
Actinomyces naeslundii		45.67	0.49	URE 10		
27513502	**Low discrimination**					**27513502**
Propionibacterium propionicum		86.09	0.60	ARA 1		
Actinomyces viscosus		7.80	0.28			
Actinomyces naeslundii		3.57	0.46			
27513503	**Low discrimination**					**27513503**
Propionibacterium propionicum		54.58	0.44	ARA 1		
Actinomyces naeslundii		30.58	0.49	URE 10		
Actinomyces viscosus		14.84	0.20			
27513520	**Low discrimination**					**27513520**
Actinomyces naeslundii		57.62	0.46	ARA 1		
Bifidobacterium longum subsp. longum		15.58	0.20			
Propionibacterium propionicum		9.35	0.23			
27513521	**Good identification**					**27513521**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.49	ARA 1			
27513522	**Low discrimination**						**27513522**
					GRA	CAT	
Actinomyces naeslundii		42.38	0.46	ARA 1	99	25	
Capnocytophaga ochracea		31.53	0.19		1	90	
Bifidobacterium longum subsp. longum		11.46	0.20		99	1	
27513523	**Good identification**						**27513523**
Actinomyces naeslundii		100.0	0.49	ARA 1			
27513540	**Very good identification**						**27513540**
Propionibacterium propionicum		98.67	0.57	ARA 1			
Actinomyces naeslundii		1.08	0.34				
Actinomyces israelii		0.24	0.13				
27513541	**Low discrimination**						**27513541**
Propionibacterium propionicum		87.08	0.41	ARA 1			
Actinomyces naeslundii		12.92	0.37	URE 10			
27513542	**Very good identification**						**27513542**
Propionibacterium propionicum		99.03	0.60	ARA 1			
Actinomyces naeslundii		0.73	0.34				
Actinomyces israelii		0.25	0.16				
27513543	**Acceptable identification**						**27513543**
Propionibacterium propionicum		91.00	0.44	ARA 1			
Actinomyces naeslundii		9.00	0.37	URE 10			
27513561	**Good identification**						**27513561**
Actinomyces naeslundii		100.0	0.37	ARA 1			
27513563	**Good identification**						**27513563**
Actinomyces naeslundii		100.0	0.37	ARA 1			
27514000	**Low discrimination**						**27514000**
					GRA		
Actinomyces odontolyticus		65.51	0.42	MNS 9	99		
Actinomyces viscosus		23.92	0.27	RAF 3	99		
Prevotella intermedia		3.80	0.15		1		
27514002	**Very good identification**						**27514002**
Actinomyces viscosus		98.30	0.62	XYL 1			
Actinomyces odontolyticus		1.67	0.38				
Lactobacillus fermentum		0.03	0.16				
27514003	**Good identification**						**27514003**
Actinomyces viscosus		100.0	0.53	XYL 1			
27514102	**Very good identification**						**27514102**
Actinomyces viscosus		99.57	0.66	XYL 1			
Actinomyces odontolyticus		0.30	0.30				
Propionibacterium propionicum		0.07	0.26				
27514103	**Very good identification**						**27514103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		100.0	0.58	XYL 1			
27514200	**Good identification**						**27514200**
Actinomyces odontolyticus		100.0	0.47	MNS 9			
27514202	**Low discrimination**						**27514202**
Actinomyces odontolyticus		76.52	0.43	MNS 9			
Actinomyces viscosus		23.49	0.27	RAF 3			
27514300	**Good identification**						**27514300**
Actinomyces odontolyticus		100.0	0.38	MNS 9			
27515000	**Low discrimination**						**27515000**
Lactobacillus fermentum		52.06	0.50	NIT 1			
Actinomyces odontolyticus		29.99	0.39				
Actinomyces viscosus		10.07	0.24				
27515002	**Good identification**						**27515002**
Actinomyces viscosus		93.12	0.58	XYL 1			
Lactobacillus fermentum		4.86	0.50				
Actinomyces odontolyticus		1.72	0.35				
27515003	**Good identification**						**27515003**
Actinomyces viscosus		99.18	0.50	XYL 1			
Actinomyces naeslundii		0.66	0.37				
Lactobacillus fermentum		0.16	0.16				
27515020	**Low discrimination**						**27515020**
Lactobacillus fermentum		49.28	0.42	NIT 1			
Clostridium perfringens		28.27	0.26				
Actinomyces odontolyticus		13.87	0.25				
27515021	**Good identification**						**27515021**
Actinomyces naeslundii		100.0	0.37	XYL 1			
27515022	**Low discrimination**						**27515022**
					GRA	CAT	
Lactobacillus fermentum		34.91	0.42	NIT 1	99	1	
Actinomyces viscosus		20.26	0.24		99	76	
Capnocytophaga ochracea		17.38	0.21		1	90	
27515100	**Low discrimination**						**27515100**
					GRA	CAT	
Lactobacillus fermentum		62.13	0.54	NIT 1	99	1	
Actinomyces viscosus		13.09	0.28		99	76	
Capnocytophaga ochracea		7.72	0.23		1	90	
27515102	**Low discrimination**						**27515102**
					GRA	CAT	
Actinomyces viscosus		89.66	0.62	XYL 1	99	76	
Capnocytophaga ochracea		4.81	0.39		1	90	
Lactobacillus fermentum		4.30	0.54		99	1	
27515103	**Good identification**						**27515103**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		98.15	0.54	XYL 1	1		
Actinomyces naeslundii		1.05	0.45		1		
Clostridium chauvoei		0.49	0.16		99		
27515120	**Low discrimination**						**27515120**
					GRA	CAT	
Capnocytophaga ochracea		30.31	0.39	ESL 99	1	90	
Clostridium perfringens		23.28	0.40	BGL 90	99	1	
Prevotella melaninogenica		18.35	0.32		1	1	
27515121	**Low discrimination**						**27515121**
					GRA	SPO	
Actinomyces naeslundii		44.60	0.45	XYL 1	99	1	
Clostridium chauvoei		39.36	0.21		99	99	
Capnocytophaga ochracea		6.76	0.05		1	1	
27515122	**Low discrimination**						**27515122**
					GRA	CAT	
Capnocytophaga ochracea		63.04	0.56	ESL 99	1	90	
Bacteroides fragilis		27.03	0.40		1	99	
Clostridium perfringens		2.53	0.35		99	1	
27515132	**Low discrimination**						**27515132**
					GRA	CAT	
Capnocytophaga ochracea		78.91	0.39	ESL 99	1	90	
Bifidobacterium longum subsp. longum		9.40	0.32	MLZ 10	99	1	
Parabacteroides distasonis		8.32	0.35		1	76	
27515162	**Low discrimination**						**27515162**
					GRA	CAT	
Capnocytophaga ochracea		89.63	0.39	ESL 99	1	90	
Clostridium perfringens		6.17	0.22	SOR 10	99	1	
Bacteroides fragilis		3.49	0.06		1	99	
27515200	**Good identification**						**27515200**
Actinomyces odontolyticus		99.11	0.44	MNS 9			
Lactobacillus fermentum		0.90	0.16	RAF 3			
27515202	**Low discrimination**						**27515202**
Actinomyces odontolyticus		77.11	0.40	MNS 9			
Actinomyces viscosus		21.76	0.24	RAF 3			
Lactobacillus fermentum		1.14	0.16				
27515300	**Low discrimination**						**27515300**
					GRA		
Actinomyces odontolyticus		84.63	0.35	MNS 9	99		
Bacteroides vulgatus		11.42	0.06	RAF 3	1		
Lactobacillus fermentum		3.95	0.20		99		
27515320	**Good identification**						**27515320**
					GRA	CAT	
Bacteroides vulgatus		97.68	0.40	NIT 1	1	3	
Actinomyces odontolyticus		1.19	0.22	ARA 93	99	10	
Capnocytophaga ochracea		0.38	0.05		1	90	
27515322	**Low discrimination**						**27515322**
					CAT		
Parabacteroides distasonis		71.48	0.48	NIT 1	76		
Capnocytophaga ochracea		11.06	0.21		90		
Prevotella oralis		6.75	0.17		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27515332	**Good identification**						**27515332**
Parabacteroides distasonis		96.91	0.43	NIT 1			
Capnocytophaga ochracea		3.09	0.05				
27515400	**Good identification**						**27515400**
Lactobacillus fermentum		96.14	0.50	NIT 1			
Bifidobacterium longum subsp. longum		3.07	0.17				
Actinomyces israelii		0.79	0.15				
27515402	**Low discrimination**						**27515402**
Lactobacillus fermentum		80.80	0.50	NIT 1			
Actinomyces viscosus		15.63	0.24				
Bifidobacterium longum subsp. longum		2.58	0.17				
27515420	**Low discrimination**						**27515420**
					GRA		
Lactobacillus fermentum		62.72	0.42	NIT 1	99		
Bifidobacterium longum subsp. longum		24.04	0.28		99		
Bacteroides vulgatus		13.24	0.09		1		
27515422	**Low discrimination**						**27515422**
					GRA	CAT	
Lactobacillus fermentum		67.98	0.42	NIT 1	99	1	
Bifidobacterium longum subsp. longum		26.05	0.28		99	1	
Capnocytophaga ochracea		5.97	0.09		1	90	
27515500	**Low discrimination**						**27515500**
					GRA		
Bifidobacterium longum subsp. longum		64.11	0.52	NIT 1	99		
Lactobacillus fermentum		33.08	0.54		99		
Bacteroides vulgatus		1.43	0.09		1		
27515502	**Low discrimination**						**27515502**
Bifidobacterium longum subsp. longum		54.49	0.52	NIT 1			
Lactobacillus fermentum		28.12	0.54				
Actinomyces viscosus		5.93	0.28				
27515510	**Good identification**						**27515510**
Bifidobacterium longum subsp. longum		99.70	0.56	NIT 1			
Lactobacillus fermentum		0.31	0.20				
27515512	**Acceptable identification**						**27515512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.78	0.56	NIT 1	99	1	
Bifidobacterium adolescentis		6.20	0.37		99	1	
Capnocytophaga ochracea		0.62	0.10		1	90	
27515520	**Low discrimination**						**27515520**
					GRA		
Bifidobacterium longum subsp. longum		61.68	0.62	NIT 1	99		
Bacteroides vulgatus		33.98	0.43		1		
Lactobacillus fermentum		2.65	0.46		99		
27515522	**Low discrimination**						**27515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		76.85	0.62	NIT 1	99	1	
Capnocytophaga ochracea		17.62	0.43		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		3.30	0.46		99	1	
27515530	**Very good identification**						**27515530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.51	0.66	NIT 1	99	1	
Bacteroides vulgatus		0.33	0.09		1	3	
Capnocytophaga ochracea		0.17	0.10		1	90	
27515532	**Very good identification**						**27515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.00	0.66	NIT 1	99	1	
Capnocytophaga ochracea		1.47	0.26		1	90	
Parabacteroides distasonis		0.54	0.31		1	76	
27515720	**Very good identification**						**27515720**
					GRA		
Bacteroides vulgatus		99.79	0.60	NIT 1	1		
Bifidobacterium longum subsp. longum		0.20	0.28		99		
Parabacteroides distasonis		0.01	0.10		1		
27515722	**Low discrimination**						**27515722**
					GRA		
Parabacteroides distasonis		45.10	0.44	NIT 1	1		
Bacteroides vulgatus		43.46	0.25		1		
Bifidobacterium longum subsp. longum		8.77	0.28		99		
27515732	**Low discrimination**						**27515732**
					GRA		
Parabacteroides distasonis		56.93	0.39	NIT 1	1		
Bifidobacterium longum subsp. longum		34.99	0.32		99		
Clostridium clostridioforme		7.05	0.18		50		
27516100	**Very good identification**						**27516100**
Propionibacterium propionicum		99.74	0.57	XYL 1			
Actinomyces naeslundii		0.26	0.23				
27516101	**Good identification**						**27516101**
Propionibacterium propionicum		96.62	0.41	XYL 1			
Actinomyces naeslundii		3.38	0.26	URE 10			
27516102	**Low discrimination**						**27516102**
Propionibacterium propionicum		86.99	0.60	XYL 1			
Actinomyces viscosus		12.86	0.32				
Actinomyces naeslundii		0.15	0.23				
27516103	**Low discrimination**						**27516103**
Propionibacterium propionicum		68.17	0.44	XYL 1			
Actinomyces viscosus		30.24	0.24	URE 10			
Actinomyces naeslundii		1.59	0.26				
27516140	**Very good identification**						**27516140**
Propionibacterium propionicum		100.0	0.57	XYL 1			
27516141	**Good identification**						**27516141**
Propionibacterium propionicum		100.0	0.41	XYL 1			
27516142	**Very good identification**						**27516142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		100.0	0.60	XYL 1			
27516143	**Good identification**						**27516143**
Propionibacterium propionicum		100.0	0.44	XYL 1			
27517001	**Good identification**						**27517001**
Actinomyces naeslundii		100.0	0.42	XYL 1			
27517003	**Low discrimination**						**27517003**
Actinomyces naeslundii		56.28	0.42	XYL 1			
Actinomyces viscosus		43.72	0.16				
27517020	**Good identification**						**27517020**
Actinomyces naeslundii		100.0	0.39	XYL 1			
27517021	**Good identification**						**27517021**
Actinomyces naeslundii		100.0	0.42	XYL 1			
27517022	**Good identification**						**27517022**
Actinomyces naeslundii		100.0	0.39	XYL 1			
27517023	**Good identification**						**27517023**
Actinomyces naeslundii		100.0	0.42	XYL 1			
27517100	**Low discrimination**						**27517100**
Propionibacterium propionicum		86.33	0.57	XYL 1			
Actinomyces israelii		7.73	0.40				
Actinomyces naeslundii		5.38	0.46				
27517101	**Low discrimination**						**27517101**
Propionibacterium propionicum		54.33	0.41	XYL 1			
Actinomyces naeslundii		45.67	0.49	URE 10			
27517102	**Low discrimination**						**27517102**
Propionibacterium propionicum		80.62	0.60	XYL 1			
Actinomyces viscosus		7.31	0.28				
Actinomyces israelii		7.22	0.43				
27517103	**Low discrimination**						**27517103**
Propionibacterium propionicum		54.58	0.44	XYL 1			
Actinomyces naeslundii		30.58	0.49	URE 10			
Actinomyces viscosus		14.84	0.20				
27517120	**Low discrimination**						**27517120**
					GRA	CAT	
Actinomyces naeslundii		65.27	0.46	XYL 1	99	25	
Propionibacterium propionicum		10.59	0.23		99	1	
Capnocytophaga ochracea		7.64	0.05		1	90	
27517121	**Good identification**						**27517121**
Actinomyces naeslundii		100.0	0.49	XYL 1			
27517123	**Good identification**						**27517123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	XYL 1		
27517140	**Very good identification**					**27517140**
Propionibacterium propionicum		95.62	0.57	XYL 1		
Actinomyces israelii		3.33	0.33			
Actinomyces naeslundii		1.05	0.34			
27517141	**Low discrimination**					**27517141**
Propionibacterium propionicum		87.08	0.41	XYL 1		
Actinomyces naeslundii		12.92	0.37	URE 10		
27517142	**Very good identification**					**27517142**
Propionibacterium propionicum		95.96	0.60	XYL 1		
Actinomyces israelii		3.34	0.36			
Actinomyces naeslundii		0.70	0.34			
27517143	**Acceptable identification**					**27517143**
Propionibacterium propionicum		91.00	0.44	XYL 1		
Actinomyces naeslundii		9.00	0.37	URE 10		
27517161	**Good identification**					**27517161**
Actinomyces naeslundii		100.0	0.37	XYL 1		
27517163	**Good identification**					**27517163**
Actinomyces naeslundii		100.0	0.37	XYL 1		
27517500	**Low discrimination**					**27517500**
Actinomyces israelii		71.47	0.38	SAL 92		
Bifidobacterium longum subsp. longum		12.32	0.17	CEL 93		
Propionibacterium propionicum		9.86	0.23			
27517502	**Low discrimination**					**27517502**
Actinomyces israelii		45.91	0.41	SAL 92		
Bifidobacterium adolescentis		39.76	0.34	CEL 93		
Propionibacterium propionicum		6.33	0.26			
27517512	**Low discrimination**					**27517512**
Bifidobacterium adolescentis		75.41	0.36	NIT 1		
Bifidobacterium longum subsp. longum		13.94	0.21			
Actinomyces israelii		10.65	0.27			
27520002	**Good identification**					**27520002**
Actinomyces viscosus		99.76	0.55	FRU 95		
Propionibacterium granulosum		0.15	0.15	TRE 7		
Actinomyces naeslundii		0.10	0.27			
27520003	**Good identification**					**27520003**
Actinomyces viscosus		99.56	0.46	FRU 95		
Actinomyces naeslundii		0.44	0.30	TRE 7		
27520102	**Very good identification**					**27520102**
Actinomyces viscosus		99.70	0.59	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.16	0.35	TRE 7		
Propionibacterium propionicum		0.14	0.24			
27520103	**Good identification**					**27520103**
Actinomyces viscosus		99.30	0.51	FRU 95		
Actinomyces naeslundii		0.70	0.38	TRE 7		
27520120	**Low discrimination**					**27520120**
Actinomyces naeslundii		56.52	0.35	FRU 95		
Clostridium perfringens		43.49	0.11	GAL 96		
27520121	**Good identification**					**27520121**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27521000	**Low discrimination**					**27521000**
Actinomyces naeslundii		64.45	0.51	FRU 95		
Actinomyces viscosus		16.96	0.17			
Propionibacterium granulosum		16.13	0.26			
27521001	**Acceptable identification**					**27521001**
					SPO	
Actinomyces naeslundii		91.41	0.54	FRU 95	1	
Actinomyces viscosus		5.35	0.09		1	
Clostridium chauvoei		3.24	0.06		99	
27521002	**Good identification**					**27521002**
Actinomyces viscosus		96.08	0.51	FRU 95		
Actinomyces naeslundii		3.69	0.51	TRE 7		
Propionibacterium granulosum		0.23	0.15			
27521003	**Low discrimination**					**27521003**
Actinomyces viscosus		85.27	0.43	FRU 95		
Actinomyces naeslundii		14.73	0.54	TRE 7		
27521021	**Good identification**					**27521021**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27521022	**Low discrimination**					**27521022**
Actinomyces naeslundii		54.27	0.51	FRU 95		
Clostridium perfringens		31.46	0.25			
Actinomyces viscosus		14.28	0.17			
27521023	**Acceptable identification**					**27521023**
Actinomyces naeslundii		94.48	0.54	FRU 95		
Actinomyces viscosus		5.53	0.09			
27521040	**Good identification**					**27521040**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27521041	**Good identification**					**27521041**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27521043	**Low discrimination**					**27521043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		75.11	0.41	FRU 95		
Actinomyces viscosus		24.89	0.09			
27521061	**Good identification**					**27521061**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27521062	**Low discrimination**					**27521062**
Actinomyces naeslundii		61.52	0.38	FRU 95		
Clostridium perfringens		38.49	0.12			
27521063	**Good identification**					**27521063**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27521100	**Low discrimination**					**27521100**
Actinomyces naeslundii		76.53	0.59	FRU 95		
Actinomyces viscosus		12.58	0.21			
Clostridium perfringens		7.51	0.19			
27521101	**Low discrimination**					**27521101**
					SPO	
Clostridium chauvoei		54.35	0.40	RAF 1	99	
Actinomyces naeslundii		44.04	0.62	TRE 10	1	
Actinomyces viscosus		1.61	0.13		1	
27521102	**Acceptable identification**					**27521102**
Actinomyces viscosus		93.75	0.55	FRU 95		
Actinomyces naeslundii		5.76	0.59	TRE 7		
Clostridium perfringens		0.27	0.14			
27521103	**Low discrimination**					**27521103**
					SPO	
Actinomyces viscosus		74.81	0.47	FRU 95	1	
Actinomyces naeslundii		20.69	0.62	TRE 7	1	
Clostridium chauvoei		4.51	0.27		99	
27521110	**Good identification**					**27521110**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27521111	**Low discrimination**					**27521111**
					SPO	
Actinomyces naeslundii		80.85	0.40	FRU 95	1	
Clostridium chauvoei		19.15	0.06	MLZ 5	99	
27521113	**Low discrimination**					**27521113**
Actinomyces naeslundii		59.03	0.40	FRU 95		
Actinomyces viscosus		40.97	0.13	MLZ 5		
27521120	**Low discrimination**					**27521120**
					GRA	
Clostridium perfringens		73.41	0.45	FRU 99	99	
Actinomyces naeslundii		23.13	0.59		99	
Prevotella melaninogenica		3.46	0.15		1	
27521121	**Low discrimination**					**27521121**
					SPO	
Actinomyces naeslundii		69.80	0.62	FRU 95	1	
Clostridium chauvoei		28.71	0.32		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		1.49	0.11		84	
27521122	**Low discrimination**					**27521122**
Clostridium perfringens		55.43	0.39	FRU 99		
Actinomyces naeslundii		37.11	0.59			
Actinomyces viscosus		6.10	0.21			
27521123	**Low discrimination**					**27521123**
					SPO	
Actinomyces naeslundii		89.35	0.62	FRU 95	1	
Clostridium chauvoei		6.49	0.19		99	
Actinomyces viscosus		3.26	0.13		1	
27521130	**Low discrimination**					**27521130**
Actinomyces naeslundii		62.15	0.37	FRU 95		
Clostridium perfringens		37.85	0.11	MLZ 5		
27521131	**Good identification**					**27521131**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27521132	**Low discrimination**					**27521132**
Actinomyces naeslundii		77.72	0.37	FRU 95		
Clostridium perfringens		22.28	0.05	MLZ 5		
27521133	**Good identification**					**27521133**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27521140	**Low discrimination**					**27521140**
Actinomyces naeslundii		79.98	0.46	FRU 95		
Propionibacterium propionicum		11.55	0.21			
Clostridium perfringens		8.47	0.07			
27521141	**Acceptable identification**					**27521141**
					SPO	
Actinomyces naeslundii		93.40	0.49	FRU 95	1	
Clostridium chauvoei		6.60	0.06		99	
27521142	**Low discrimination**					**27521142**
Actinomyces naeslundii		45.50	0.46	FRU 95		
Actinomyces viscosus		42.38	0.21			
Propionibacterium propionicum		9.85	0.24			
27521143	**Low discrimination**					**27521143**
Actinomyces naeslundii		82.85	0.49	FRU 95		
Actinomyces viscosus		17.15	0.13			
27521161	**Good identification**					**27521161**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27521163	**Good identification**					**27521163**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27522001	**Good identification**					**27522001**
Actinomyces naeslundii		100.0	0.35	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27522021	**Good identification**					**27522021**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27522023	**Good identification**					**27522023**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27522100	**Good identification**					**27522100**
Propionibacterium propionicum		96.89	0.55	FRU 99		
Actinomyces naeslundii		3.11	0.40			
27522101	**Low discrimination**					**27522101**
Propionibacterium propionicum		69.79	0.39	FRU 99		
Actinomyces naeslundii		30.21	0.43	URE 10		
27522102	**Good identification**					**27522102**
Propionibacterium propionicum		91.55	0.58	FRU 99		
Actinomyces viscosus		6.49	0.25			
Actinomyces naeslundii		1.96	0.40			
27522103	**Low discrimination**					**27522103**
Propionibacterium propionicum		66.61	0.42	FRU 99		
Actinomyces naeslundii		19.23	0.43	URE 10		
Actinomyces viscosus		14.17	0.17			
27522120	**Low discrimination**					**27522120**
Actinomyces naeslundii		76.05	0.40	FRU 95		
Propionibacterium propionicum		23.95	0.21	GAL 96		
27522121	**Good identification**					**27522121**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27522122	**Low discrimination**					**27522122**
Actinomyces naeslundii		67.91	0.40	FRU 95		
Propionibacterium propionicum		32.09	0.24	GAL 96		
27522123	**Good identification**					**27522123**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27522140	**Good identification**					**27522140**
Propionibacterium propionicum		99.44	0.55	FRU 99		
Actinomyces naeslundii		0.56	0.27			
27522141	**Acceptable identification**					**27522141**
Propionibacterium propionicum		92.90	0.39	FRU 99		
Actinomyces naeslundii		7.10	0.30	URE 10		
27522142	**Very good identification**					**27522142**
Propionibacterium propionicum		99.62	0.58	FRU 99		
Actinomyces naeslundii		0.38	0.27			
27522143	**Good identification**					**27522143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		95.15	0.42	FRU 99		
Actinomyces naeslundii		4.85	0.30	URE 10		
27523000	**Low discrimination**					**27523000**
					CAT	
Actinomyces naeslundii		88.14	0.56	FRU 95	25	
Propionibacterium granulosum		8.57	0.24		90	
Propionibacterium propionicum		3.29	0.21		1	
27523001	**Very good identification**					**27523001**
Actinomyces naeslundii		100.0	0.59	FRU 95		
27523002	**Low discrimination**					**27523002**
Actinomyces naeslundii		82.24	0.56	FRU 95		
Actinomyces viscosus		11.15	0.17			
Propionibacterium propionicum		4.61	0.24			
27523003	**Very good identification**					**27523003**
Actinomyces naeslundii		97.08	0.59	FRU 95		
Actinomyces viscosus		2.92	0.09			
27523011	**Good identification**					**27523011**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27523013	**Good identification**					**27523013**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27523020	**Good identification**					**27523020**
Actinomyces naeslundii		97.63	0.56	FRU 95		
Propionibacterium granulosum		2.37	0.13			
27523021	**Very good identification**					**27523021**
Actinomyces naeslundii		100.0	0.59	FRU 95		
27523022	**Good identification**					**27523022**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27523023	**Very good identification**					**27523023**
Actinomyces naeslundii		100.0	0.59	FRU 95		
27523031	**Good identification**					**27523031**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27523033	**Good identification**					**27523033**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27523040	**Low discrimination**					**27523040**
Actinomyces naeslundii		82.53	0.43	FRU 95		
Propionibacterium propionicum		17.47	0.21			
27523041	**Good identification**					**27523041**
Actinomyces naeslundii		100.0	0.46	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27523042	**Low discrimination**					**27523042**
Actinomyces naeslundii		75.90	0.43	FRU 95		
Propionibacterium propionicum		24.10	0.24			
27523043	**Good identification**					**27523043**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27523060	**Good identification**					**27523060**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27523061	**Good identification**					**27523061**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27523062	**Good identification**					**27523062**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27523063	**Good identification**					**27523063**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27523100	**Low discrimination**					**27523100**
Propionibacterium propionicum		56.51	0.55	FRU 99		
Actinomyces naeslundii		43.49	0.64			
27523101	**Good identification**					**27523101**
					SPO	
Actinomyces naeslundii		90.69	0.67	FRU 95	1	
Propionibacterium propionicum		8.73	0.39		1	
Clostridium chauvoei		0.58	0.06		99	
27523102	**Low discrimination**					**27523102**
Propionibacterium propionicum		64.25	0.58	FRU 99		
Actinomyces naeslundii		32.96	0.64			
Actinomyces viscosus		2.79	0.21			
27523103	**Low discrimination**					**27523103**
Actinomyces naeslundii		85.97	0.67	FRU 95		
Propionibacterium propionicum		12.41	0.42			
Actinomyces viscosus		1.62	0.13			
27523110	**Low discrimination**					**27523110**
Actinomyces naeslundii		80.04	0.42	FRU 95		
Propionibacterium propionicum		19.96	0.21	MLZ 5		
27523111	**Good identification**					**27523111**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27523112	**Low discrimination**					**27523112**
Actinomyces naeslundii		72.78	0.42	FRU 95		
Propionibacterium propionicum		27.22	0.24	MLZ 5		
27523113	**Good identification**					**27523113**
Actinomyces naeslundii		100.0	0.45	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27523120	**Very good identification**					**27523120**
Actinomyces naeslundii		97.12	0.64	FRU 95		
Clostridium perfringens		1.60	0.11			
Propionibacterium propionicum		1.28	0.21			
27523121	**Very good identification**					**27523121**
Actinomyces naeslundii		100.0	0.67	FRU 95		
27523122	**Very good identification**					**27523122**
Actinomyces naeslundii		97.33	0.64	FRU 95		
Propionibacterium propionicum		1.92	0.24			
Clostridium perfringens		0.76	0.05			
27523123	**Very good identification**					**27523123**
Actinomyces naeslundii		100.0	0.67	FRU 95		
27523130	**Good identification**					**27523130**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27523131	**Good identification**					**27523131**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27523132	**Good identification**					**27523132**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27523133	**Good identification**					**27523133**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27523140	**Low discrimination**					**27523140**
Propionibacterium propionicum		88.04	0.55	FRU 99		
Actinomyces naeslundii		11.96	0.51			
27523141	**Low discrimination**					**27523141**
Actinomyces naeslundii		64.71	0.54	FRU 95		
Propionibacterium propionicum		35.29	0.39			
27523142	**Good identification**					**27523142**
Propionibacterium propionicum		91.70	0.58	FRU 99		
Actinomyces naeslundii		8.30	0.51			
27523143	**Low discrimination**					**27523143**
Actinomyces naeslundii		55.00	0.54	FRU 95		
Propionibacterium propionicum		45.00	0.42			
27523160	**Acceptable identification**					**27523160**
Actinomyces naeslundii		93.08	0.51	FRU 95		
Propionibacterium propionicum		6.92	0.21			
27523161	**Good identification**					**27523161**
Actinomyces naeslundii		100.0	0.54	FRU 95		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27523162	**Low discrimination**					**27523162**
Actinomyces naeslundii		89.96	0.51	FRU 95		
Propionibacterium propionicum		10.04	0.24			
27523163	**Good identification**					**27523163**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27525520	**Good identification**					**27525520**
Bifidobacterium longum subsp. longum		100.0	0.35	NIT 1		
27525522	**Good identification**					**27525522**
					GRA	
Bifidobacterium longum subsp. longum		97.21	0.35	NIT 1	99	
Parabacteroides distasonis		2.79	0.13	FRU 90	1	
27525530	**Good identification**					**27525530**
Bifidobacterium longum subsp. longum		100.0	0.39	NIT 1		
27525532	**Good identification**					**27525532**
Bifidobacterium longum subsp. longum		100.0	0.39	NIT 1		
27530000	**Low discrimination**					**27530000**
Propionibacterium granulosum		72.38	0.60	NIT 1		
Actinomyces viscosus		23.83	0.42			
Actinomyces naeslundii		2.31	0.49			
27530002	**Very good identification**					**27530002**
Actinomyces viscosus		99.13	0.77	TRE 7		
Propionibacterium granulosum		0.76	0.50			
Actinomyces naeslundii		0.10	0.49			
27530003	**Very good identification**					**27530003**
Actinomyces viscosus		99.54	0.68	TRE 7		
Actinomyces naeslundii		0.44	0.52			
Propionibacterium granulosum		0.02	0.15			
27530010	**Good identification**					**27530010**
Propionibacterium granulosum		96.46	0.45	NIT 1		
Actinomyces viscosus		2.35	0.08	MLZ 12		
Actinomyces naeslundii		1.19	0.27			
27530012	**Acceptable identification**					**27530012**
Actinomyces viscosus		90.20	0.42	TRE 7		
Propionibacterium granulosum		9.34	0.35	MLZ 1		
Actinomyces naeslundii		0.46	0.27			
27530020	**Low discrimination**					**27530020**
					CAT	
Propionibacterium granulosum		74.63	0.50	NIT 1	90	
Clostridium perfringens		14.84	0.30		1	
Actinomyces naeslundii		9.54	0.49		25	
27530021	**Acceptable identification**					**27530021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.95	0.52	GAL 96		
Propionibacterium granulosum		4.90	0.15			
Actinomyces viscosus		2.15	0.00			
27530022	**Low discrimination**					**27530022**
Actinomyces viscosus		73.64	0.42	TRE 7		
Propionibacterium granulosum		13.98	0.39	NAG 1		
Actinomyces naeslundii		7.15	0.49			
27530041	**Low discrimination**					**27530041**
					SPO	
Actinomyces naeslundii		78.21	0.39	GAL 96	1	
Clostridium botulinum biovar B		11.54	-.03		99	
Actinomyces viscosus		10.25	0.00		1	
27530042	**Good identification**					**27530042**
Actinomyces viscosus		95.98	0.42	TRE 7		
Actinomyces naeslundii		1.64	0.36	SOR 1		
Propionibacterium propionicum		1.28	0.24			
27530061	**Good identification**					**27530061**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27530062	**Low discrimination**					**27530062**
Actinomyces naeslundii		42.02	0.36	GAL 96		
Clostridium perfringens		33.20	0.12			
Actinomyces viscosus		24.78	0.08			
27530063	**Low discrimination**					**27530063**
Actinomyces naeslundii		88.41	0.39	GAL 96		
Actinomyces viscosus		11.59	0.00			
27530100	**Low discrimination**					**27530100**
Actinomyces viscosus		58.56	0.47	TRE 7		
Propionibacterium propionicum		28.99	0.55	BGL 99		
Actinomyces naeslundii		9.10	0.57			
27530101	**Low discrimination**					**27530101**
Actinomyces viscosus		52.91	0.38	TRE 7		
Actinomyces naeslundii		37.00	0.60	BGL 99		
Propionibacterium propionicum		8.73	0.39			
27530102	**Very good identification**					**27530102**
Actinomyces viscosus		99.09	0.81	TRE 7		
Propionibacterium propionicum		0.74	0.58			
Actinomyces naeslundii		0.16	0.57			
27530103	**Very good identification**					**27530103**
Actinomyces viscosus		99.05	0.73	TRE 7		
Actinomyces naeslundii		0.70	0.60			
Propionibacterium propionicum		0.25	0.42			
27530111	**Low discrimination**					**27530111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		78.46	0.38	GAL 96		
Actinomyces viscosus		21.54	0.04	MLZ 5		
27530112	**Good identification**					**27530112**
Actinomyces viscosus		98.46	0.47	TRE 7		
Actinomyces naeslundii		0.81	0.35	MLZ 1		
Propionibacterium propionicum		0.74	0.24			
27530113	**Good identification**					**27530113**
Actinomyces viscosus		96.45	0.38	TRE 7		
Actinomyces naeslundii		3.55	0.38	MLZ 1		
27530120	**Low discrimination**					**27530120**
Clostridium perfringens		77.83	0.45	GAL 99		
Actinomyces naeslundii		19.41	0.57			
Actinomyces viscosus		1.26	0.12			
27530121	**Good identification**					**27530121**
Actinomyces naeslundii		94.91	0.60	GAL 96		
Clostridium perfringens		2.56	0.11			
Actinomyces viscosus		1.37	0.04			
27530122	**Low discrimination**					**27530122**
Actinomyces viscosus		68.61	0.47	TRE 7		
Clostridium perfringens		20.11	0.39	NAG 1		
Actinomyces naeslundii		10.66	0.57			
27530123	**Low discrimination**					**27530123**
Actinomyces viscosus		58.55	0.38	TRE 7		
Actinomyces naeslundii		40.93	0.60	NAG 1		
Clostridium perfringens		0.52	0.05			
27530130	**Low discrimination**					**27530130**
Actinomyces naeslundii		56.52	0.35	GAL 96		
Clostridium perfringens		43.49	0.11	MLZ 5		
27530131	**Good identification**					**27530131**
Actinomyces naeslundii		100.0	0.38	GAL 96		
27530133	**Low discrimination**					**27530133**
Actinomyces naeslundii		78.46	0.38	GAL 96		
Actinomyces viscosus		21.54	0.04	MLZ 5		
27530140	**Acceptable identification**					**27530140**
Propionibacterium propionicum		91.95	0.55	MAN 99		
Actinomyces naeslundii		5.09	0.44			
Actinomyces viscosus		1.88	0.12			
27530141	**Low discrimination**					**27530141**
Propionibacterium propionicum		55.28	0.39	MAN 99		
Actinomyces naeslundii		41.34	0.47	URE 10		
Actinomyces viscosus		3.38	0.04			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27530142	**Low discrimination**					**27530142**
Actinomyces viscosus		56.45	0.47	TRE 7		
Propionibacterium propionicum		41.91	0.58	SOR 1		
Actinomyces naeslundii		1.55	0.44			
27530143	**Low discrimination**					**27530143**
Actinomyces viscosus		72.95	0.38	TRE 7		
Propionibacterium propionicum		18.05	0.42	SOR 1		
Actinomyces naeslundii		9.00	0.47			
27530161	**Good identification**					**27530161**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27530163	**Acceptable identification**					**27530163**
Actinomyces naeslundii		92.43	0.47	GAL 96		
Actinomyces viscosus		7.57	0.04			
27530202	**Good identification**					**27530202**
					CAT	
Actinomyces viscosus		96.86	0.42	TRE 7	76	
Propionibacterium granulosum		2.28	0.24	RHA 1	90	
Actinomyces odontolyticus		0.87	0.14		10	
27530302	**Good identification**					**27530302**
Actinomyces viscosus		98.95	0.47	TRE 7		
Propionibacterium propionicum		0.74	0.24	RHA 1		
Actinomyces odontolyticus		0.16	0.06			
27530303	**Good identification**					**27530303**
Actinomyces viscosus		99.30	0.38	TRE 7		
Actinomyces naeslundii		0.70	0.26	RHA 1		
27530402	**Good identification**					**27530402**
Actinomyces viscosus		99.24	0.42	TRE 7		
Propionibacterium granulosum		0.76	0.15	ARA 1		
27530502	**Good identification**					**27530502**
Actinomyces viscosus		99.10	0.47	TRE 7		
Propionibacterium propionicum		0.74	0.24	ARA 1		
Actinomyces naeslundii		0.16	0.23			
27530503	**Good identification**					**27530503**
Actinomyces viscosus		99.30	0.38	TRE 7		
Actinomyces naeslundii		0.70	0.26	ARA 1		
27531000	**Low discrimination**					**27531000**
Propionibacterium granulosum		46.67	0.60	NIT 1		
Actinomyces naeslundii		35.79	0.73			
Actinomyces viscosus		9.42	0.39			
27531001	**Good identification**					**27531001**
Actinomyces naeslundii		93.20	0.76			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		5.45	0.31			
Propionibacterium granulosum		0.82	0.26			
27531002	**Good identification**					**27531002**
Actinomyces viscosus		94.79	0.73	TRE 7		
Actinomyces naeslundii		3.64	0.73			
Propionibacterium granulosum		1.19	0.50			
27531003	**Low discrimination**					**27531003**
Actinomyces viscosus		85.23	0.65	TRE 7		
Actinomyces naeslundii		14.72	0.76			
Propionibacterium granulosum		0.03	0.15			
27531010	**Low discrimination**					**27531010**
Propionibacterium granulosum		75.63	0.45	NIT 1		
Actinomyces naeslundii		22.38	0.51	MLZ 12		
Actinomyces viscosus		1.13	0.05			
27531011	**Good identification**					**27531011**
Actinomyces naeslundii		97.78	0.54	MLZ 5		
Propionibacterium granulosum		2.23	0.11			
27531012	**Low discrimination**					**27531012**
Actinomyces viscosus		73.05	0.39	TRE 7		
Actinomyces naeslundii		14.61	0.51	MLZ 1		
Propionibacterium granulosum		12.34	0.35			
27531020	**Low discrimination**					**27531020**
					CAT	
Clostridium perfringens		82.83	0.64	LAC 88	1	
Actinomyces naeslundii		12.91	0.73		25	
Propionibacterium granulosum		4.21	0.50		90	
27531021	**Very good identification**					**27531021**
					CAT	
Actinomyces naeslundii		95.52	0.76		25	
Clostridium perfringens		4.13	0.30		1	
Propionibacterium granulosum		0.21	0.15		90	
27531022	**Low discrimination**					**27531022**
Clostridium perfringens		69.15	0.59	LAC 88		
Actinomyces naeslundii		22.90	0.73			
Actinomyces viscosus		6.03	0.39			
27531023	**Good identification**					**27531023**
Actinomyces naeslundii		92.69	0.76			
Actinomyces viscosus		5.42	0.31			
Clostridium perfringens		1.89	0.25			
27531031	**Good identification**					**27531031**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
27531032	**Low discrimination**					**27531032**
					CAT	
Actinomyces naeslundii		54.31	0.51	MLZ 5	25	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		31.48	0.25		1	
Propionibacterium granulosum		11.47	0.25		90	
27531033	**Good identification**					**27531033**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
27531040	**Low discrimination**					**27531040**
					CAT	
Actinomyces naeslundii		74.92	0.60		25	
Clostridium perfringens		16.05	0.26		1	
Propionibacterium granulosum		5.59	0.26		90	
27531041	**Very good identification**					**27531041**
Actinomyces naeslundii		100.0	0.63			
27531042	**Low discrimination**					**27531042**
Actinomyces viscosus		56.42	0.39	TRE 7		
Actinomyces naeslundii		37.84	0.60	SOR 1		
Clostridium perfringens		3.82	0.21			
27531043	**Low discrimination**					**27531043**
Actinomyces naeslundii		75.11	0.63			
Actinomyces viscosus		24.89	0.31			
27531050	**Low discrimination**					**27531050**
Actinomyces naeslundii		83.79	0.38	MLZ 5		
Propionibacterium granulosum		16.21	0.11			
27531051	**Good identification**					**27531051**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27531052	**Low discrimination**					**27531052**
Actinomyces naeslundii		77.75	0.38	MLZ 5		
Actinomyces viscosus		22.25	0.05			
27531053	**Good identification**					**27531053**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27531060	**Low discrimination**					**27531060**
					CAT	
Clostridium perfringens		87.18	0.52	LAC 88	1	
Actinomyces naeslundii		12.59	0.60		25	
Propionibacterium granulosum		0.24	0.15		90	
27531061	**Very good identification**					**27531061**
Actinomyces naeslundii		95.54	0.63			
Clostridium perfringens		4.46	0.18			
27531062	**Low discrimination**					**27531062**
Clostridium perfringens		76.26	0.46	LAC 88		
Actinomyces naeslundii		23.39	0.60			
Actinomyces viscosus		0.35	0.05			
27531063	**Very good identification**					**27531063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.85	0.63			
Clostridium perfringens		2.15	0.12			
27531071	**Good identification**					**27531071**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27531072	**Low discrimination**					**27531072**
Actinomyces naeslundii		61.52	0.38	MLZ 5		
Clostridium perfringens		38.49	0.12			
27531073	**Good identification**					**27531073**
Actinomyces naeslundii		100.0	0.41	MLZ 5		
27531100	**Low discrimination**					**27531100**
Actinomyces naeslundii		54.65	0.81			
Clostridium perfringens		27.95	0.53			
Actinomyces viscosus		8.98	0.43			
27531101	**Low discrimination**					**27531101**
					SPO	
Actinomyces naeslundii		83.24	0.84		1	
Clostridium chauvoei		12.61	0.47		99	
Actinomyces viscosus		3.04	0.35		1	
27531102	**Good identification**					**27531102**
Actinomyces viscosus		91.82	0.77	TRE 7		
Actinomyces naeslundii		5.64	0.81			
Clostridium perfringens		1.36	0.48			
27531103	**Low discrimination**					**27531103**
					SPO	
Actinomyces viscosus		77.61	0.69	TRE 7	1	
Actinomyces naeslundii		21.46	0.84		1	
Clostridium chauvoei		0.57	0.34		99	
27531110	**Low discrimination**					**27531110**
Actinomyces naeslundii		85.68	0.59	MLZ 5		
Clostridium perfringens		8.41	0.19			
Actinomyces viscosus		2.70	0.09			
27531111	**Very good identification**					**27531111**
					SPO	
Actinomyces naeslundii		96.52	0.62	MLZ 5	1	
Clostridium chauvoei		2.81	0.12		99	
Actinomyces viscosus		0.68	0.01		1	
27531112	**Low discrimination**					**27531112**
Actinomyces viscosus		74.05	0.43	TRE 7		
Actinomyces naeslundii		23.71	0.59	MLZ 1		
Clostridium perfringens		1.10	0.14			
27531113	**Low discrimination**					**27531113**
					SPO	
Actinomyces naeslundii		58.85	0.62	MLZ 5	1	
Actinomyces viscosus		40.84	0.35		1	
Clostridium chauvoei		0.30	-.01		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27531120	**Good identification**					**27531120**
					GRA	
Clostridium perfringens		93.86	0.79		99	
Actinomyces naeslundii		5.68	0.81		99	
Prevotella melaninogenica		0.40	0.32		1	
27531121	**Low discrimination**					**27531121**
					SPO	
Actinomyces naeslundii		86.01	0.84		1	
Clostridium perfringens		9.58	0.45		84	
Clostridium chauvoei		4.34	0.38		99	
27531122	**Low discrimination**					**27531122**
Clostridium perfringens		86.23	0.74			
Actinomyces naeslundii		11.08	0.81			
Actinomyces viscosus		1.82	0.43			
27531123	**Good identification**					**27531123**
Actinomyces naeslundii		91.06	0.84			
Clostridium perfringens		4.77	0.39			
Actinomyces viscosus		3.33	0.35			
27531130	**Low discrimination**					**27531130**
Clostridium perfringens		75.22	0.45	MLZ 1		
Actinomyces naeslundii		23.70	0.59			
Bifidobacterium longum subsp. longum		0.52	0.13			
27531131	**Very good identification**					**27531131**
					SPO	
Actinomyces naeslundii		96.99	0.62	MLZ 5	1	
Clostridium perfringens		2.07	0.11		84	
Clostridium chauvoei		0.94	0.04		99	
27531132	**Low discrimination**					**27531132**
					GRA	
Clostridium perfringens		54.20	0.39	MLZ 1	99	
Actinomyces naeslundii		36.29	0.59		99	
Parabacteroides distasonis		4.31	0.30		1	
27531133	**Very good identification**					**27531133**
Actinomyces naeslundii		98.32	0.62	MLZ 5		
Clostridium perfringens		0.99	0.05			
Actinomyces viscosus		0.69	0.01			
27531140	**Low discrimination**					**27531140**
Actinomyces naeslundii		43.22	0.68			
Propionibacterium propionicum		32.51	0.55			
Clostridium perfringens		23.86	0.41			
27531141	**Good identification**					**27531141**
					SPO	
Actinomyces naeslundii		93.44	0.71		1	
Propionibacterium propionicum		5.21	0.39		1	
Clostridium chauvoei		0.81	0.12		99	
27531142	**Low discrimination**					**27531142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		33.93	0.58	MAN 99			
Actinomyces naeslundii		30.07	0.68				
Actinomyces viscosus		28.01	0.43				
27531143	**Low discrimination**						**27531143**
Actinomyces naeslundii		77.29	0.71				
Actinomyces viscosus		16.00	0.35				
Propionibacterium propionicum		6.46	0.42				
27531150	**Low discrimination**						**27531150**
Actinomyces naeslundii		79.98	0.46	MLZ 5			
Propionibacterium propionicum		11.55	0.21				
Clostridium perfringens		8.47	0.07				
27531151	**Good identification**						**27531151**
Actinomyces naeslundii		100.0	0.49	MLZ 5			
27531152	**Low discrimination**						**27531152**
Actinomyces naeslundii		69.20	0.46	MLZ 5			
Propionibacterium propionicum		14.98	0.24				
Actinomyces viscosus		12.37	0.09				
27531153	**Good identification**						**27531153**
Actinomyces naeslundii		96.18	0.49	MLZ 5			
Actinomyces viscosus		3.82	0.01				
27531160	**Good identification**						**27531160**
Clostridium perfringens		94.62	0.67				
Actinomyces naeslundii		5.30	0.68				
Propionibacterium propionicum		0.04	0.21				
27531161	**Low discrimination**						**27531161**
					SPO		
Actinomyces naeslundii		89.04	0.71		1		
Clostridium perfringens		10.70	0.33		84		
Clostridium chauvoei		0.26	0.04		99		
27531162	**Low discrimination**						**27531162**
					GRA	CAT	
Clostridium perfringens		88.91	0.61		99	1	
Actinomyces naeslundii		10.59	0.68		99	25	
Capnocytophaga ochracea		0.28	0.15		1	90	
27531163	**Good identification**						**27531163**
Actinomyces naeslundii		94.46	0.71				
Clostridium perfringens		5.34	0.27				
Actinomyces viscosus		0.20	0.01				
27531171	**Good identification**						**27531171**
Actinomyces naeslundii		100.0	0.49	MLZ 5			
27531173	**Good identification**						**27531173**
Actinomyces naeslundii		100.0	0.49	MLZ 5			
27531201	**Good identification**						**27531201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	RHA 1		
27531202	**Acceptable identification**					**27531202**
Actinomyces viscosus		92.04	0.39	TRE 7		
Actinomyces naeslundii		3.53	0.39	RHA 1		
Propionibacterium granulosum		3.53	0.24			
27531221	**Good identification**					**27531221**
Actinomyces naeslundii		100.0	0.42	RHA 1		
27531223	**Good identification**					**27531223**
Actinomyces naeslundii		100.0	0.42	RHA 1		
27531300	**Low discrimination**					**27531300**
Actinomyces naeslundii		53.92	0.46	RHA 1		
Clostridium perfringens		27.58	0.19			
Actinomyces viscosus		8.86	0.09			
27531301	**Low discrimination**					**27531301**
					SPO	
Actinomyces naeslundii		84.17	0.49	RHA 1	1	
Clostridium chauvoei		12.76	0.12		99	
Actinomyces viscosus		3.07	0.01		1	
27531302	**Acceptable identification**					**27531302**
Actinomyces viscosus		91.87	0.43	TRE 7		
Actinomyces naeslundii		5.65	0.46	RHA 1		
Clostridium perfringens		1.36	0.14			
27531320	**Acceptable identification**					**27531320**
					GRA	
Clostridium perfringens		93.92	0.45	RHA 1	99	
Actinomyces naeslundii		5.68	0.46		99	
Prevotella melaninogenica		0.40	-.03		1	
27531321	**Low discrimination**					**27531321**
					SPO	
Actinomyces naeslundii		86.07	0.49	RHA 1	1	
Clostridium perfringens		9.59	0.11		84	
Clostridium chauvoei		4.35	0.04		99	
27531322	**Low discrimination**					**27531322**
					GRA	
Clostridium perfringens		60.95	0.39	RHA 1	99	
Parabacteroides distasonis		27.00	0.43		1	
Actinomyces naeslundii		7.83	0.46		99	
27531323	**Acceptable identification**					**27531323**
Actinomyces naeslundii		91.83	0.49	RHA 1		
Clostridium perfringens		4.81	0.05			
Actinomyces viscosus		3.35	0.01			
27531332	**Acceptable identification**					**27531332**
					GRA	
Parabacteroides distasonis		93.40	0.38	NIT 1	1	
Clostridium perfringens		3.96	0.05	XYL 90	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	2.65	0.25		99	
27531341	**Good identification**					**27531341**
	Actinomyces naeslundii	100.0	0.37	RHA 1		
27531343	**Low discrimination**					**27531343**
	Actinomyces naeslundii	82.86	0.37	RHA 1		
	Actinomyces viscosus	17.15	0.01			
27531361	**Good identification**					**27531361**
	Actinomyces naeslundii	100.0	0.37	RHA 1		
27531363	**Good identification**					**27531363**
	Actinomyces naeslundii	100.0	0.37	RHA 1		
27531401	**Good identification**					**27531401**
	Actinomyces naeslundii	100.0	0.42	ARA 1		
27531402	**Acceptable identification**					**27531402**
	Actinomyces viscosus	94.09	0.39	TRE 7		
	Actinomyces naeslundii	3.61	0.39	ARA 1		
	Propionibacterium granulosum	1.18	0.15			
27531421	**Good identification**					**27531421**
	Actinomyces naeslundii	100.0	0.42	ARA 1		
27531423	**Good identification**					**27531423**
	Actinomyces naeslundii	100.0	0.42	ARA 1		
27531501	**Low discrimination**					**27531501**
					SPO	
	Actinomyces naeslundii	84.17	0.49	ARA 1	1	
	Clostridium chauvoei	12.76	0.12		99	
	Actinomyces viscosus	3.07	0.01		1	
27531502	**Low discrimination**					**27531502**
	Actinomyces viscosus	81.63	0.43	TRE 7		
	Bifidobacterium longum subsp. longum	8.40	0.33	ARA 1		
	Actinomyces naeslundii	5.02	0.46			
27531510	**Good identification**					**27531510**
	Bifidobacterium longum subsp. longum	98.19	0.37	NIT 1		
	Actinomyces naeslundii	1.81	0.25			
27531512	**Low discrimination**					**27531512**
	Bifidobacterium longum subsp. longum	86.51	0.37	NIT 1		
	Bifidobacterium adolescentis	6.91	0.20			
	Actinomyces viscosus	4.99	0.09			
27531520	**Low discrimination**					**27531520**
	Clostridium perfringens	67.65	0.45	ARA 1		
	Bifidobacterium longum subsp. longum	27.39	0.43			
	Actinomyces naeslundii	4.09	0.46			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27531521	**Low discrimination**						**27531521**
					SPO		
Actinomyces naeslundii		82.85	0.49	ARA 1	1		
Clostridium perfringens		9.23	0.11		84		
Clostridium chauvoei		4.19	0.04		99		
27531522	**Low discrimination**						**27531522**
Clostridium perfringens		45.38	0.39	ARA 1			
Bifidobacterium longum subsp. longum		39.04	0.43				
Actinomyces naeslundii		5.83	0.46				
27531523	**Low discrimination**						**27531523**
Actinomyces naeslundii		88.18	0.49	ARA 1			
Clostridium perfringens		4.62	0.05				
Bifidobacterium longum subsp. longum		3.98	0.09				
27531530	**Good identification**						**27531530**
Bifidobacterium longum subsp. longum		98.11	0.47	NIT 1			
Clostridium perfringens		1.44	0.11				
Actinomyces naeslundii		0.45	0.25				
27531532	**Good identification**						**27531532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		95.05	0.47	NIT 1	99	1	
Parabacteroides distasonis		3.16	0.26		1	76	
Capnocytophaga ochracea		0.69	0.02		1	90	
27531541	**Good identification**						**27531541**
Actinomyces naeslundii		100.0	0.37	ARA 1			
27531543	**Low discrimination**						**27531543**
Actinomyces naeslundii		82.86	0.37	ARA 1			
Actinomyces viscosus		17.15	0.01				
27531561	**Good identification**						**27531561**
Actinomyces naeslundii		100.0	0.37	ARA 1			
27531563	**Good identification**						**27531563**
Actinomyces naeslundii		100.0	0.37	ARA 1			
27531722	**Acceptable identification**						**27531722**
					GRA		
Parabacteroides distasonis		91.73	0.39	NIT 1	1		
Clostridium perfringens		3.41	0.05	XYL 90	99		
Bifidobacterium longum subsp. longum		2.94	0.09		99		
27532000	**Low discrimination**						**27532000**
					CAT		
Propionibacterium granulosum		68.00	0.58	NIT 1	90		
Propionibacterium propionicum		26.11	0.55		1		
Actinomyces naeslundii		5.59	0.54		25		
27532001	**Low discrimination**						**27532001**
					CAT		
Actinomyces naeslundii		69.45	0.57	GAL 96	25		
Propionibacterium propionicum		24.03	0.39		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		5.69	0.24		90	
27532002	**Low discrimination**					**27532002**
					CAT	
Propionibacterium propionicum		42.84	0.58	LAC 99	1	
Actinomyces viscosus		32.46	0.42		76	
Propionibacterium granulosum		18.59	0.48		90	
27532010	**Acceptable identification**					**27532010**
					CAT	
Propionibacterium granulosum		94.32	0.43	NIT 1	90	
Actinomyces naeslundii		2.99	0.32	MLZ 12	25	
Propionibacterium propionicum		2.68	0.21		1	
27532011	**Low discrimination**					**27532011**
Actinomyces naeslundii		82.50	0.35	GAL 96		
Propionibacterium granulosum		17.50	0.09	MLZ 5		
27532013	**Low discrimination**					**27532013**
Actinomyces naeslundii		81.54	0.35	GAL 96		
Actinomyces viscosus		18.46	0.00	MLZ 5		
27532020	**Low discrimination**					**27532020**
					CAT	
Propionibacterium granulosum		74.38	0.48	NIT 1	90	
Actinomyces naeslundii		24.46	0.54		25	
Propionibacterium propionicum		1.15	0.21		1	
27532021	**Very good identification**					**27532021**
Actinomyces naeslundii		97.99	0.57	GAL 96		
Propionibacterium granulosum		2.01	0.13			
27532022	**Low discrimination**					**27532022**
					CAT	
Actinomyces naeslundii		53.06	0.54	GAL 96	25	
Propionibacterium granulosum		40.34	0.37		90	
Propionibacterium propionicum		3.76	0.24		1	
27532023	**Very good identification**					**27532023**
Actinomyces naeslundii		98.82	0.57	GAL 96		
Actinomyces viscosus		1.18	0.00			
27532031	**Good identification**					**27532031**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27532033	**Good identification**					**27532033**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27532040	**Acceptable identification**					**27532040**
					CAT	
Propionibacterium propionicum		93.60	0.55	LAC 99	1	
Actinomyces naeslundii		3.54	0.41		25	
Propionibacterium granulosum		2.46	0.24		90	
27532041	**Low discrimination**					**27532041**
Propionibacterium propionicum		66.23	0.39	LAC 99		
Actinomyces naeslundii		33.78	0.44	URE 10		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27532042	**Very good identification**					**27532042**
Propionibacterium propionicum		96.41	0.58	LAC 99		
Actinomyces naeslundii		2.43	0.41			
Actinomyces viscosus		0.74	0.08			
27532043	**Low discrimination**					**27532043**
Propionibacterium propionicum		73.37	0.42	LAC 99		
Actinomyces naeslundii		24.95	0.44	URE 10		
Actinomyces viscosus		1.68	0.00			
27532060	**Low discrimination**					**27532060**
					CAT	
Actinomyces naeslundii		69.38	0.41	GAL 96	25	
Propionibacterium propionicum		18.55	0.21		1	
Propionibacterium granulosum		12.07	0.13		90	
27532061	**Good identification**					**27532061**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27532062	**Low discrimination**					**27532062**
Actinomyces naeslundii		71.38	0.41	GAL 96		
Propionibacterium propionicum		28.62	0.24			
27532063	**Good identification**					**27532063**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27532100	**Very good identification**					**27532100**
					CAT	
Propionibacterium propionicum		99.34	0.90		1	
Actinomyces naeslundii		0.61	0.62		25	
Propionibacterium granulosum		0.03	0.24		90	
27532101	**Good identification**					**27532101**
Propionibacterium propionicum		92.28	0.73	URE 10		
Actinomyces naeslundii		7.67	0.65			
Actinomyces viscosus		0.06	0.04			
27532102	**Very good identification**					**27532102**
Propionibacterium propionicum		98.26	0.93			
Actinomyces viscosus		1.34	0.47			
Actinomyces naeslundii		0.40	0.62			
27532103	**Good identification**					**27532103**
Propionibacterium propionicum		91.22	0.76	URE 10		
Actinomyces naeslundii		5.05	0.65			
Actinomyces viscosus		3.72	0.38			
27532110	**Good identification**					**27532110**
					CAT	
Propionibacterium propionicum		96.56	0.55	MLZ 1	1	
Actinomyces naeslundii		3.10	0.40		25	
Propionibacterium granulosum		0.35	0.09		90	
27532111	**Low discrimination**					**27532111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		69.79	0.39	MLZ 1		
Actinomyces naeslundii		30.21	0.43	URE 10		
27532112	**Very good identification**					**27532112**
Propionibacterium propionicum		96.62	0.58	MLZ 1		
Actinomyces naeslundii		2.07	0.40			
Actinomyces viscosus		1.31	0.12			
27532113	**Low discrimination**					**27532113**
Propionibacterium propionicum		75.22	0.42	MLZ 1		
Actinomyces naeslundii		21.71	0.43	URE 10		
Actinomyces viscosus		3.07	0.04			
27532120	**Low discrimination**					**27532120**
Propionibacterium propionicum		61.40	0.55	NAG 1		
Actinomyces naeslundii		37.41	0.62			
Clostridium perfringens		0.78	0.11			
27532121	**Low discrimination**					**27532121**
Actinomyces naeslundii		89.16	0.65	GAL 96		
Propionibacterium propionicum		10.84	0.39			
27532122	**Low discrimination**					**27532122**
Propionibacterium propionicum		70.24	0.58	NAG 1		
Actinomyces naeslundii		28.53	0.62			
Actinomyces viscosus		0.96	0.12			
27532123	**Low discrimination**					**27532123**
Actinomyces naeslundii		84.05	0.65	GAL 96		
Propionibacterium propionicum		15.33	0.42			
Actinomyces viscosus		0.63	0.04			
27532130	**Low discrimination**					**27532130**
Actinomyces naeslundii		76.05	0.40	GAL 96		
Propionibacterium propionicum		23.95	0.21	MLZ 5		
27532131	**Good identification**					**27532131**
Actinomyces naeslundii		100.0	0.43	GAL 96		
27532132	**Low discrimination**					**27532132**
Actinomyces naeslundii		67.91	0.40	GAL 96		
Propionibacterium propionicum		32.09	0.24	MLZ 5		
27532133	**Good identification**					**27532133**
Actinomyces naeslundii		100.0	0.43	GAL 96		
27532140	**Very good identification**					**27532140**
Propionibacterium propionicum		99.89	0.90			
Actinomyces naeslundii		0.11	0.49			
27532141	**Very good identification**					**27532141**
Propionibacterium propionicum		98.56	0.73	URE 10		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		1.45	0.52			
27532142	**Excellent identification**					**27532142**
Propionibacterium propionicum		99.91	0.93			
Actinomyces naeslundii		0.07	0.49			
Actinomyces viscosus		0.01	0.12			
27532143	**Very good identification**					**27532143**
Propionibacterium propionicum		98.99	0.76	URE 10		
Actinomyces naeslundii		0.97	0.52			
Actinomyces viscosus		0.04	0.04			
27532150	**Good identification**					**27532150**
Propionibacterium propionicum		99.44	0.55	MLZ 1		
Actinomyces naeslundii		0.56	0.27			
27532151	**Acceptable identification**					**27532151**
Propionibacterium propionicum		92.90	0.39	MLZ 1		
Actinomyces naeslundii		7.10	0.30	URE 10		
27532152	**Very good identification**					**27532152**
Propionibacterium propionicum		99.62	0.58	MLZ 1		
Actinomyces naeslundii		0.38	0.27			
27532153	**Good identification**					**27532153**
Propionibacterium propionicum		95.15	0.42	MLZ 1		
Actinomyces naeslundii		4.85	0.30	URE 10		
27532160	**Acceptable identification**					**27532160**
Propionibacterium propionicum		90.29	0.55	NAG 1		
Actinomyces naeslundii		9.71	0.49			
27532161	**Low discrimination**					**27532161**
Actinomyces naeslundii		59.21	0.52	GAL 96		
Propionibacterium propionicum		40.79	0.39			
27532162	**Good identification**					**27532162**
Propionibacterium propionicum		93.31	0.58	NAG 1		
Actinomyces naeslundii		6.69	0.49			
27532163	**Low discrimination**					**27532163**
Propionibacterium propionicum		50.82	0.42	NAG 1		
Actinomyces naeslundii		49.18	0.52	URE 10		
27532300	**Good identification**					**27532300**
Propionibacterium propionicum		99.39	0.55	RHA 1		
Actinomyces naeslundii		0.61	0.28			
27532301	**Acceptable identification**					**27532301**
Propionibacterium propionicum		92.33	0.39	RHA 1		
Actinomyces naeslundii		7.67	0.31	URE 10		
27532302	**Very good identification**					**27532302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		98.26	0.58	RHA 1		
Actinomyces viscosus		1.34	0.12			
Actinomyces naeslundii		0.40	0.28			
27532303	**Acceptable identification**					**27532303**
Propionibacterium propionicum		91.23	0.42	RHA 1		
Actinomyces naeslundii		5.05	0.31	URE 10		
Actinomyces viscosus		3.72	0.04			
27532340	**Good identification**					**27532340**
Propionibacterium propionicum		100.0	0.55	RHA 1		
27532341	**Good identification**					**27532341**
Propionibacterium propionicum		100.0	0.39	RHA 1		
27532342	**Very good identification**					**27532342**
Propionibacterium propionicum		100.0	0.58	RHA 1		
27532343	**Good identification**					**27532343**
Propionibacterium propionicum		100.0	0.42	RHA 1		
27532500	**Good identification**					**27532500**
Propionibacterium propionicum		99.39	0.55	ARA 1		
Actinomyces naeslundii		0.61	0.28			
27532501	**Acceptable identification**					**27532501**
Propionibacterium propionicum		92.33	0.39	ARA 1		
Actinomyces naeslundii		7.67	0.31	URE 10		
27532502	**Very good identification**					**27532502**
Propionibacterium propionicum		98.26	0.58	ARA 1		
Actinomyces viscosus		1.34	0.12			
Actinomyces naeslundii		0.40	0.28			
27532503	**Acceptable identification**					**27532503**
Propionibacterium propionicum		91.23	0.42	ARA 1		
Actinomyces naeslundii		5.05	0.31	URE 10		
Actinomyces viscosus		3.72	0.04			
27532540	**Good identification**					**27532540**
Propionibacterium propionicum		100.0	0.55	ARA 1		
27532541	**Good identification**					**27532541**
Propionibacterium propionicum		100.0	0.39	ARA 1		
27532542	**Very good identification**					**27532542**
Propionibacterium propionicum		100.0	0.58	ARA 1		
27532543	**Good identification**					**27532543**
Propionibacterium propionicum		100.0	0.42	ARA 1		
27533000	**Low discrimination**					**27533000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Actinomyces naeslundii		58.63	0.78		25	
Propionibacterium granulosum		29.71	0.58		90	
Propionibacterium propionicum		11.41	0.55		1	
27533001	**Very good identification**					**27533001**
					CAT	
Actinomyces naeslundii		98.25	0.81		25	
Propionibacterium propionicum		1.42	0.39		1	
Propionibacterium granulosum		0.34	0.24		90	
27533002	**Low discrimination**					**27533002**
Actinomyces naeslundii		64.23	0.78			
Propionibacterium propionicum		18.75	0.58			
Actinomyces viscosus		8.71	0.39			
27533003	**Very good identification**					**27533003**
Actinomyces naeslundii		95.00	0.81			
Actinomyces viscosus		2.86	0.31			
Propionibacterium propionicum		2.06	0.42			
27533010	**Low discrimination**					**27533010**
					CAT	
Propionibacterium granulosum		55.87	0.43	NIT 1	90	
Actinomyces naeslundii		42.55	0.56	MLZ 12	25	
Propionibacterium propionicum		1.59	0.21		1	
27533011	**Very good identification**					**27533011**
Actinomyces naeslundii		99.12	0.59	MLZ 5		
Propionibacterium granulosum		0.88	0.09			
27533012	**Low discrimination**					**27533012**
					CAT	
Actinomyces naeslundii		70.91	0.56	MLZ 5	25	
Propionibacterium granulosum		23.28	0.33		90	
Propionibacterium propionicum		3.97	0.24		1	
27533013	**Very good identification**					**27533013**
Actinomyces naeslundii		100.0	0.59	MLZ 5		
27533020	**Low discrimination**					**27533020**
					CAT	
Actinomyces naeslundii		86.05	0.78		25	
Propionibacterium granulosum		10.90	0.48		90	
Clostridium perfringens		2.87	0.30		1	
27533021	**Very good identification**					**27533021**
Actinomyces naeslundii		99.92	0.81			
Propionibacterium granulosum		0.09	0.13			
27533022	**Very good identification**					**27533022**
					CAT	
Actinomyces naeslundii		95.08	0.78		25	
Propionibacterium granulosum		3.01	0.37		90	
Clostridium perfringens		1.49	0.25		1	
27533023	**Very good identification**					**27533023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.81			
27533030	**Low discrimination**					**27533030**
Actinomyces naeslundii		75.29	0.56	MLZ 5		
Propionibacterium granulosum		24.71	0.33			
27533031	**Very good identification**					**27533031**
Actinomyces naeslundii		100.0	0.59	MLZ 5		
27533032	**Acceptable identification**					**27533032**
Actinomyces naeslundii		92.42	0.56	MLZ 5		
Propionibacterium granulosum		7.58	0.22			
27533033	**Very good identification**					**27533033**
Actinomyces naeslundii		100.0	0.59	MLZ 5		
27533040	**Low discrimination**					**27533040**
					CAT	
Propionibacterium propionicum		51.64	0.55	LAC 99	1	
Actinomyces naeslundii		46.82	0.65		25	
Propionibacterium granulosum		1.36	0.24		90	
27533041	**Good identification**					**27533041**
Actinomyces naeslundii		92.45	0.68			
Propionibacterium propionicum		7.55	0.39			
27533042	**Low discrimination**					**27533042**
Propionibacterium propionicum		61.84	0.58	LAC 99		
Actinomyces naeslundii		37.38	0.65			
Actinomyces viscosus		0.29	0.05			
27533043	**Low discrimination**					**27533043**
Actinomyces naeslundii		89.08	0.68			
Propionibacterium propionicum		10.92	0.42			
27533050	**Low discrimination**					**27533050**
					CAT	
Actinomyces naeslundii		77.71	0.43	MLZ 5	25	
Propionibacterium propionicum		16.45	0.21		1	
Propionibacterium granulosum		5.84	0.09		90	
27533051	**Good identification**					**27533051**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
27533052	**Low discrimination**					**27533052**
Actinomyces naeslundii		75.90	0.43	MLZ 5		
Propionibacterium propionicum		24.10	0.24			
27533053	**Good identification**					**27533053**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
27533060	**Good identification**					**27533060**
Actinomyces naeslundii		94.84	0.65			
Clostridium perfringens		3.42	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		1.06	0.21			
27533061	**Very good identification**					**27533061**
Actinomyces naeslundii		100.0	0.68			
27533062	**Very good identification**					**27533062**
Actinomyces naeslundii		96.74	0.65			
Clostridium perfringens		1.64	0.12			
Propionibacterium propionicum		1.62	0.24			
27533063	**Very good identification**					**27533063**
Actinomyces naeslundii		100.0	0.68			
27533070	**Good identification**					**27533070**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27533071	**Good identification**					**27533071**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
27533072	**Good identification**					**27533072**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27533073	**Good identification**					**27533073**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
27533100	**Low discrimination**					**27533100**
Propionibacterium propionicum		87.04	0.90			
Actinomyces naeslundii		12.86	0.86			
Actinomyces israelii		0.03	0.32			
27533101	**Low discrimination**					**27533101**
					SPO	
Actinomyces naeslundii		66.56	0.89		1	
Propionibacterium propionicum		33.38	0.73		1	
Clostridium chauvoei		0.05	0.12		99	
27533102	**Good identification**					**27533102**
Propionibacterium propionicum		90.31	0.93			
Actinomyces naeslundii		8.89	0.86			
Actinomyces viscosus		0.75	0.43			
27533103	**Low discrimination**					**27533103**
Actinomyces naeslundii		56.46	0.89			
Propionibacterium propionicum		42.47	0.76			
Actinomyces viscosus		1.06	0.35			
27533110	**Low discrimination**					**27533110**
Propionibacterium propionicum		56.23	0.55	MLZ 1		
Actinomyces naeslundii		43.27	0.64			
Actinomyces israelii		0.30	0.17			
27533111	**Good identification**					**27533111**
Actinomyces naeslundii		91.22	0.67	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		8.78	0.39			
27533112	**Low discrimination**					**27533112**
Propionibacterium propionicum		65.50	0.58	MLZ 1		
Actinomyces naeslundii		33.61	0.64			
Actinomyces viscosus		0.55	0.09			
27533113	**Low discrimination**					**27533113**
Actinomyces naeslundii		87.11	0.67	MLZ 5		
Propionibacterium propionicum		12.58	0.42			
Actinomyces viscosus		0.32	0.01			
27533120	**Low discrimination**					**27533120**
Actinomyces naeslundii		86.56	0.86			
Clostridium perfringens		7.45	0.45			
Propionibacterium propionicum		5.92	0.55			
27533121	**Very good identification**					**27533121**
Actinomyces naeslundii		99.41	0.89			
Propionibacterium propionicum		0.50	0.39			
Clostridium perfringens		0.06	0.11			
27533122	**Low discrimination**					**27533122**
Actinomyces naeslundii		87.38	0.86			
Propionibacterium propionicum		8.96	0.58			
Clostridium perfringens		3.54	0.39			
27533123	**Very good identification**					**27533123**
Actinomyces naeslundii		99.20	0.89			
Propionibacterium propionicum		0.75	0.42			
Clostridium perfringens		0.03	0.05			
27533130	**Very good identification**					**27533130**
Actinomyces naeslundii		97.12	0.64	MLZ 5		
Clostridium perfringens		1.60	0.11			
Propionibacterium propionicum		1.28	0.21			
27533131	**Very good identification**					**27533131**
Actinomyces naeslundii		100.0	0.67	MLZ 5		
27533132	**Very good identification**					**27533132**
Actinomyces naeslundii		97.33	0.64	MLZ 5		
Propionibacterium propionicum		1.92	0.24			
Clostridium perfringens		0.76	0.05			
27533133	**Very good identification**					**27533133**
Actinomyces naeslundii		100.0	0.67	MLZ 5		
27533140	**Very good identification**					**27533140**
Propionibacterium propionicum		97.44	0.90			
Actinomyces naeslundii		2.54	0.73			
Actinomyces israelii		0.01	0.25			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27533141	**Low discrimination**					**27533141**
Propionibacterium propionicum		73.97	0.73	URE 10		
Actinomyces naeslundii		26.03	0.76			
27533142	**Very good identification**					**27533142**
Propionibacterium propionicum		98.27	0.93			
Actinomyces naeslundii		1.71	0.73			
Actinomyces israelii		0.01	0.28			
27533143	**Low discrimination**					**27533143**
Propionibacterium propionicum		80.98	0.76	URE 10		
Actinomyces naeslundii		19.00	0.76			
Actinomyces viscosus		0.02	0.01			
27533150	**Low discrimination**					**27533150**
Propionibacterium propionicum		88.04	0.55	MLZ 1		
Actinomyces naeslundii		11.96	0.51			
27533151	**Low discrimination**					**27533151**
Actinomyces naeslundii		64.71	0.54	MLZ 5		
Propionibacterium propionicum		35.29	0.39			
27533152	**Good identification**					**27533152**
Propionibacterium propionicum		91.52	0.58	MLZ 1		
Actinomyces naeslundii		8.29	0.51			
Actinomyces israelii		0.19	0.13			
27533153	**Low discrimination**					**27533153**
Actinomyces naeslundii		55.00	0.54	MLZ 5		
Propionibacterium propionicum		45.00	0.42			
27533160	**Low discrimination**					**27533160**
Actinomyces naeslundii		67.55	0.73			
Propionibacterium propionicum		26.18	0.55			
Clostridium perfringens		6.27	0.33			
27533161	**Very good identification**					**27533161**
Actinomyces naeslundii		97.21	0.76			
Propionibacterium propionicum		2.79	0.39			
27533162	**Low discrimination**					**27533162**
Actinomyces naeslundii		61.54	0.73			
Propionibacterium propionicum		35.77	0.58			
Clostridium perfringens		2.69	0.27			
27533163	**Very good identification**					**27533163**
Actinomyces naeslundii		95.87	0.76			
Propionibacterium propionicum		4.13	0.42			
27533170	**Acceptable identification**					**27533170**
Actinomyces naeslundii		93.08	0.51	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		6.92	0.21			
27533171	**Good identification**					**27533171**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
27533172	**Low discrimination**					**27533172**
Actinomyces naeslundii		89.96	0.51	MLZ 5		
Propionibacterium propionicum		10.04	0.24			
27533173	**Good identification**					**27533173**
Actinomyces naeslundii		100.0	0.54	MLZ 5		
27533201	**Good identification**					**27533201**
Actinomyces naeslundii		100.0	0.47	RHA 1		
27533202	**Low discrimination**					**27533202**
					CAT	
Actinomyces naeslundii		55.08	0.44	RHA 1	25	
Propionibacterium granulosum		21.37	0.22		90	
Propionibacterium propionicum		16.08	0.24		1	
27533203	**Good identification**					**27533203**
Actinomyces naeslundii		100.0	0.47	RHA 1		
27533220	**Low discrimination**					**27533220**
Actinomyces naeslundii		72.05	0.44	RHA 1		
Propionibacterium granulosum		27.95	0.22			
27533221	**Good identification**					**27533221**
Actinomyces naeslundii		100.0	0.47	RHA 1		
27533222	**Acceptable identification**					**27533222**
Actinomyces naeslundii		91.16	0.44	RHA 1		
Propionibacterium granulosum		8.84	0.11			
27533223	**Good identification**					**27533223**
Actinomyces naeslundii		100.0	0.47	RHA 1		
27533300	**Low discrimination**					**27533300**
Propionibacterium propionicum		87.13	0.55	RHA 1		
Actinomyces naeslundii		12.87	0.51			
27533301	**Low discrimination**					**27533301**
Actinomyces naeslundii		66.60	0.54	RHA 1		
Propionibacterium propionicum		33.40	0.39			
27533302	**Good identification**					**27533302**
Propionibacterium propionicum		90.13	0.58	RHA 1		
Actinomyces naeslundii		8.88	0.51			
Actinomyces viscosus		0.75	0.09			
27533303	**Low discrimination**					**27533303**
Actinomyces naeslundii		56.46	0.54	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		42.47	0.42			
Actinomyces viscosus		1.06	0.01			
27533320	**Low discrimination**					**27533320**
Actinomyces naeslundii		86.62	0.51	RHA 1		
Clostridium perfringens		7.45	0.11			
Propionibacterium propionicum		5.92	0.21			
27533321	**Good identification**					**27533321**
Actinomyces naeslundii		100.0	0.54	RHA 1		
27533322	**Low discrimination**					**27533322**
Actinomyces naeslundii		87.48	0.51	RHA 1		
Propionibacterium propionicum		8.97	0.24			
Clostridium perfringens		3.54	0.05			
27533323	**Good identification**					**27533323**
Actinomyces naeslundii		100.0	0.54	RHA 1		
27533340	**Good identification**					**27533340**
Propionibacterium propionicum		97.46	0.55	RHA 1		
Actinomyces naeslundii		2.54	0.38			
27533341	**Low discrimination**					**27533341**
Propionibacterium propionicum		73.97	0.39	RHA 1		
Actinomyces naeslundii		26.03	0.41	URE 10		
27533342	**Very good identification**					**27533342**
Propionibacterium propionicum		98.29	0.58	RHA 1		
Actinomyces naeslundii		1.71	0.38			
27533343	**Low discrimination**					**27533343**
Propionibacterium propionicum		81.00	0.42	RHA 1		
Actinomyces naeslundii		19.00	0.41	URE 10		
27533360	**Low discrimination**					**27533360**
Actinomyces naeslundii		72.07	0.38	RHA 1		
Propionibacterium propionicum		27.93	0.21			
27533361	**Good identification**					**27533361**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27533362	**Low discrimination**					**27533362**
Actinomyces naeslundii		63.24	0.38	RHA 1		
Propionibacterium propionicum		36.76	0.24			
27533363	**Good identification**					**27533363**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27533400	**Low discrimination**					**27533400**
					CAT	
Actinomyces naeslundii		54.27	0.44	ARA 1	25	
Propionibacterium granulosum		27.51	0.24		90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		10.56	0.21		1	
27533401	**Good identification**					**27533401**
Actinomyces naeslundii		100.0	0.47	ARA	1	
27533402	**Low discrimination**					**27533402**
Actinomyces naeslundii		56.63	0.44	ARA	1	
Propionibacterium propionicum		16.53	0.24			
Actinomyces israelii		11.98	0.23			
27533403	**Good identification**					**27533403**
Actinomyces naeslundii		100.0	0.47	ARA	1	
27533420	**Low discrimination**					**27533420**
Actinomyces naeslundii		88.76	0.44	ARA	1	
Propionibacterium granulosum		11.24	0.13			
27533421	**Good identification**					**27533421**
Actinomyces naeslundii		100.0	0.47	ARA	1	
27533422	**Good identification**					**27533422**
Actinomyces naeslundii		100.0	0.44	ARA	1	
27533423	**Good identification**					**27533423**
Actinomyces naeslundii		100.0	0.47	ARA	1	
27533500	**Low discrimination**					**27533500**
Propionibacterium propionicum		84.96	0.55	ARA	1	
Actinomyces naeslundii		12.55	0.51			
Actinomyces israelii		2.49	0.30			
27533501	**Low discrimination**					**27533501**
Actinomyces naeslundii		66.60	0.54	ARA	1	
Propionibacterium propionicum		33.40	0.39			
27533502	**Low discrimination**					**27533502**
Propionibacterium propionicum		87.43	0.58	ARA	1	
Actinomyces naeslundii		8.61	0.51			
Actinomyces israelii		2.56	0.33			
27533503	**Low discrimination**					**27533503**
Actinomyces naeslundii		56.46	0.54	ARA	1	
Propionibacterium propionicum		42.47	0.42			
Actinomyces viscosus		1.06	0.01			
27533520	**Low discrimination**					**27533520**
Actinomyces naeslundii		84.09	0.51	ARA	1	
Clostridium perfringens		7.24	0.11			
Propionibacterium propionicum		5.75	0.21			
27533521	**Good identification**					**27533521**
Actinomyces naeslundii		100.0	0.54	ARA	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27533522	**Low discrimination**					**27533522**
Actinomyces naeslundii		84.90	0.51	ARA 1		
Propionibacterium propionicum		8.71	0.24			
Clostridium perfringens		3.44	0.05			
27533523	**Good identification**					**27533523**
Actinomyces naeslundii		100.0	0.54	ARA 1		
27533540	**Good identification**					**27533540**
Propionibacterium propionicum		96.39	0.55	ARA 1		
Actinomyces naeslundii		2.51	0.38			
Actinomyces israelii		1.10	0.23			
27533541	**Low discrimination**					**27533541**
Propionibacterium propionicum		73.97	0.39	ARA 1		
Actinomyces naeslundii		26.03	0.41	URE 10		
27533542	**Very good identification**					**27533542**
Propionibacterium propionicum		97.20	0.58	ARA 1		
Actinomyces naeslundii		1.69	0.38			
Actinomyces israelii		1.11	0.26			
27533543	**Low discrimination**					**27533543**
Propionibacterium propionicum		81.00	0.42	ARA 1		
Actinomyces naeslundii		19.00	0.41	URE 10		
27533560	**Low discrimination**					**27533560**
Actinomyces naeslundii		72.07	0.38	ARA 1		
Propionibacterium propionicum		27.93	0.21			
27533561	**Good identification**					**27533561**
Actinomyces naeslundii		100.0	0.41	ARA 1		
27533562	**Low discrimination**					**27533562**
Actinomyces naeslundii		63.24	0.38	ARA 1		
Propionibacterium propionicum		36.76	0.24			
27533563	**Good identification**					**27533563**
Actinomyces naeslundii		100.0	0.41	ARA 1		
27534002	**Good identification**					**27534002**
Actinomyces viscosus		99.24	0.42	TRE 7		
Propionibacterium granulosum		0.76	0.15	XYL 1		
27534102	**Good identification**					**27534102**
Actinomyces viscosus		99.10	0.47	TRE 7		
Propionibacterium propionicum		0.74	0.24	XYL 1		
Actinomyces naeslundii		0.16	0.23			
27534103	**Good identification**					**27534103**
Actinomyces viscosus		99.30	0.38	TRE 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.70	0.26	XYL 1			
27535001	**Good identification**						**27535001**
Actinomyces naeslundii		100.0	0.42	XYL 1			
27535002	**Acceptable identification**						**27535002**
Actinomyces viscosus		91.16	0.39	TRE 7			
Actinomyces israelii		3.57	0.30	XYL 1			
Actinomyces naeslundii		3.50	0.39				
27535021	**Good identification**						**27535021**
Actinomyces naeslundii		100.0	0.42	XYL 1			
27535023	**Good identification**						**27535023**
Actinomyces naeslundii		100.0	0.42	XYL 1			
27535100	**Low discrimination**						**27535100**
Actinomyces naeslundii		29.72	0.46	XYL 1			
Actinomyces israelii		28.37	0.37				
Clostridium perfringens		15.20	0.19				
27535101	**Low discrimination**						**27535101**
					SPO		
Actinomyces naeslundii		84.17	0.49	XYL 1	1		
Clostridium chauvoei		12.76	0.12		99		
Actinomyces viscosus		3.07	0.01		1		
27535102	**Low discrimination**						**27535102**
Actinomyces viscosus		82.29	0.43	TRE 7			
Actinomyces israelii		7.24	0.40	XYL 1			
Actinomyces naeslundii		5.06	0.46				
27535120	**Acceptable identification**						**27535120**
Clostridium perfringens		90.37	0.45	XYL 1			
Actinomyces naeslundii		5.47	0.46				
Bifidobacterium longum subsp. longum		1.11	0.17				
27535121	**Low discrimination**						**27535121**
					SPO		
Actinomyces naeslundii		86.07	0.49	XYL 1	1		
Clostridium perfringens		9.59	0.11		84		
Clostridium chauvoei		4.35	0.04		99		
27535122	**Low discrimination**						**27535122**
					GRA	CAT	
Parabacteroides distasonis		48.82	0.51	NIT 1	1	76	
Clostridium perfringens		36.73	0.39		99	1	
Capnocytophaga ochracea		4.98	0.21		1	90	
27535123	**Low discrimination**						**27535123**
					GRA		
Actinomyces naeslundii		86.31	0.49	XYL 1	99		
Parabacteroides distasonis		6.01	0.17		1		
Clostridium perfringens		4.52	0.05		99		
27535132	**Acceptable identification**						**27535132**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		90.37	0.46	NIT 1	1	76	
Bifidobacterium longum subsp. longum		5.60	0.21		99	1	
Capnocytophaga ochracea		1.90	0.05		1	90	
27535141	**Good identification**						**27535141**
Actinomyces naeslundii		100.0	0.37	XYL 1			
27535143	**Low discrimination**						**27535143**
Actinomyces naeslundii		82.86	0.37	XYL 1			
Actinomyces viscosus		17.15	0.01				
27535161	**Good identification**						**27535161**
Actinomyces naeslundii		100.0	0.37	XYL 1			
27535163	**Good identification**						**27535163**
Actinomyces naeslundii		100.0	0.37	XYL 1			
27535322	**Very good identification**						**27535322**
					GRA		
Parabacteroides distasonis		99.62	0.59	NIT 1	1		
Clostridium perfringens		0.25	0.05		99		
Prevotella oralis		0.13	-.03		1		
27535332	**Good identification**						**27535332**
Parabacteroides distasonis		99.23	0.54	NIT 1			
Clostridium clostridioforme		0.77	0.13				
27535500	**Low discrimination**						**27535500**
Bifidobacterium longum subsp. longum		85.00	0.41	NIT 1			
Actinomyces israelii		13.23	0.35				
Lactobacillus fermentum		1.77	0.20				
27535502	**Low discrimination**						**27535502**
Bifidobacterium longum subsp. longum		64.10	0.41	NIT 1			
Bifidobacterium adolescentis		15.90	0.33				
Actinomyces israelii		14.96	0.38				
27535510	**Good identification**						**27535510**
Bifidobacterium longum subsp. longum		98.65	0.45	NIT 1			
Actinomyces israelii		1.35	0.21				
27535512	**Low discrimination**						**27535512**
Bifidobacterium longum subsp. longum		81.23	0.45	NIT 1			
Bifidobacterium adolescentis		14.46	0.34				
Clostridium clostridioforme		2.65	0.18				
27535520	**Acceptable identification**						**27535520**
					GRA		
Bifidobacterium longum subsp. longum		90.43	0.52	NIT 1	99		
Bacteroides vulgatus		8.30	0.20		1		
Clostridium perfringens		0.75	0.11		99		
27535522	**Low discrimination**						**27535522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.87	0.52	NIT 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		23.64	0.47		1	76	
Capnocytophaga ochracea		0.69	0.09		1	90	
27535530	**Good identification**						**27535530**
Bifidobacterium longum subsp. longum		100.0	0.56	NIT 1			
27535532	**Acceptable identification**						**27535532**
					GRA		
Bifidobacterium longum subsp. longum		90.26	0.56	NIT 1	99		
Parabacteroides distasonis		9.01	0.43		1		
Clostridium clostridioforme		0.74	0.18		50		
27535720	**Good identification**						**27535720**
					GRA		
Bacteroides vulgatus		97.68	0.36	NIT 1	1		
Bifidobacterium longum subsp. longum		1.19	0.17	TRE 4	99		
Parabacteroides distasonis		1.13	0.21		1		
27535722	**Acceptable identification**						**27535722**
Parabacteroides distasonis		94.54	0.55	NIT 1			
Clostridium clostridioforme		2.23	0.22				
Bacteroides thetaiotaomicron		1.39	0.08				
27535732	**Low discrimination**						**27535732**
					GRA		
Parabacteroides distasonis		80.40	0.51	NIT 1	1		
Clostridium clostridioforme		16.03	0.33		50		
Bifidobacterium longum subsp. longum		2.71	0.21		99		
27536100	**Good identification**						**27536100**
Propionibacterium propionicum		98.98	0.55	XYL 1			
Actinomyces naeslundii		0.61	0.28				
Actinomyces israelii		0.41	0.16				
27536101	**Acceptable identification**						**27536101**
Propionibacterium propionicum		92.33	0.39	XYL 1			
Actinomyces naeslundii		7.67	0.31	URE 10			
27536102	**Very good identification**						**27536102**
Propionibacterium propionicum		97.86	0.58	XYL 1			
Actinomyces viscosus		1.33	0.12				
Actinomyces israelii		0.41	0.19				
27536103	**Acceptable identification**						**27536103**
Propionibacterium propionicum		91.23	0.42	XYL 1			
Actinomyces naeslundii		5.05	0.31	URE 10			
Actinomyces viscosus		3.72	0.04				
27536140	**Good identification**						**27536140**
Propionibacterium propionicum		100.0	0.55	XYL 1			
27536141	**Good identification**						**27536141**
Propionibacterium propionicum		100.0	0.39	XYL 1			
27536142	**Very good identification**						**27536142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.58	XYL 1		
27536143	**Good identification**					**27536143**
Propionibacterium propionicum		100.0	0.42	XYL 1		
27537000	**Low discrimination**					**27537000**
					CAT	
Actinomyces israelii		53.82	0.39	SAL 92	1	
Actinomyces naeslundii		27.14	0.44	CEL 93	25	
Propionibacterium granulosum		13.76	0.24		90	
27537001	**Good identification**					**27537001**
Actinomyces naeslundii		100.0	0.47	XYL 1		
27537002	**Low discrimination**					**27537002**
Actinomyces israelii		65.68	0.42	SAL 92		
Actinomyces naeslundii		22.08	0.44	CEL 93		
Propionibacterium propionicum		6.45	0.24			
27537003	**Good identification**					**27537003**
Actinomyces naeslundii		100.0	0.47	XYL 1		
27537020	**Low discrimination**					**27537020**
					CAT	
Actinomyces naeslundii		79.79	0.44	XYL 1	25	
Propionibacterium granulosum		10.11	0.13		90	
Actinomyces israelii		10.10	0.19		1	
27537021	**Good identification**					**27537021**
Actinomyces naeslundii		100.0	0.47	XYL 1		
27537022	**Low discrimination**					**27537022**
Actinomyces naeslundii		84.05	0.44	XYL 1		
Actinomyces israelii		15.96	0.22			
27537023	**Good identification**					**27537023**
Actinomyces naeslundii		100.0	0.47	XYL 1		
27537042	**Low discrimination**					**27537042**
Actinomyces israelii		71.18	0.35	SAL 92		
Propionibacterium propionicum		17.96	0.24	CEL 93		
Actinomyces naeslundii		10.86	0.31			
27537100	**Low discrimination**					**27537100**
Propionibacterium propionicum		64.13	0.55	XYL 1		
Actinomyces israelii		26.40	0.50			
Actinomyces naeslundii		9.47	0.51			
27537101	**Low discrimination**					**27537101**
Actinomyces naeslundii		65.78	0.54	XYL 1		
Propionibacterium propionicum		32.99	0.39			
Actinomyces israelii		1.23	0.16			
27537102	**Low discrimination**					**27537102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		65.52	0.58	XYL 1		
Actinomyces israelii		26.97	0.53			
Actinomyces naeslundii		6.45	0.51			
27537103	**Low discrimination**					**27537103**
Actinomyces naeslundii		55.58	0.54	XYL 1		
Propionibacterium propionicum		41.81	0.42			
Actinomyces israelii		1.57	0.19			
27537110	**Low discrimination**					**27537110**
Actinomyces israelii		77.48	0.36	SAL 92		
Propionibacterium propionicum		12.72	0.21	CEL 93		
Actinomyces naeslundii		9.79	0.29	MLZ 13		
27537112	**Low discrimination**					**27537112**
Actinomyces israelii		71.33	0.39	SAL 92		
Propionibacterium propionicum		11.71	0.24	CEL 93		
Bifidobacterium adolescentis		10.95	0.18	MLZ 13		
27537120	**Low discrimination**					**27537120**
Actinomyces naeslundii		75.06	0.51	XYL 1		
Actinomyces israelii		13.35	0.29			
Clostridium perfringens		6.46	0.11			
27537121	**Good identification**					**27537121**
Actinomyces naeslundii		100.0	0.54	XYL 1		
27537122	**Low discrimination**					**27537122**
Actinomyces naeslundii		68.32	0.51	XYL 1		
Actinomyces israelii		18.23	0.32			
Propionibacterium propionicum		7.01	0.24			
27537123	**Good identification**					**27537123**
Actinomyces naeslundii		100.0	0.54	XYL 1		
27537140	**Low discrimination**					**27537140**
Propionibacterium propionicum		84.31	0.55	XYL 1		
Actinomyces israelii		13.50	0.43			
Actinomyces naeslundii		2.20	0.38			
27537141	**Low discrimination**					**27537141**
Propionibacterium propionicum		73.97	0.39	XYL 1		
Actinomyces naeslundii		26.03	0.41	URE 10		
27537142	**Low discrimination**					**27537142**
Propionibacterium propionicum		84.93	0.58	XYL 1		
Actinomyces israelii		13.60	0.46			
Actinomyces naeslundii		1.48	0.38			
27537143	**Low discrimination**					**27537143**
Propionibacterium propionicum		81.00	0.42	XYL 1		
Actinomyces naeslundii		19.00	0.41	URE 10		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27537160	**Low discrimination**					**27537160**
Actinomyces naeslundii		56.20	0.38	XYL 1		
Actinomyces israelii		22.03	0.22			
Propionibacterium propionicum		21.78	0.21			
27537161	**Good identification**					**27537161**
Actinomyces naeslundii		100.0	0.41	XYL 1		
27537162	**Low discrimination**					**27537162**
Actinomyces naeslundii		46.10	0.38	XYL 1		
Actinomyces israelii		27.11	0.25			
Propionibacterium propionicum		26.79	0.24			
27537163	**Good identification**					**27537163**
Actinomyces naeslundii		100.0	0.41	XYL 1		
27537400	**Good identification**					**27537400**
Actinomyces israelii		100.0	0.38	SAL 92		
27537402	**Good identification**					**27537402**
Actinomyces israelii		100.0	0.41	SAL 92		
27537500	**Good identification**					**27537500**
Actinomyces israelii		96.02	0.48	SAL 92		
Propionibacterium propionicum		2.88	0.21	CEL 93		
Bifidobacterium longum subsp. longum		1.10	0.07			
27537502	**Low discrimination**					**27537502**
Actinomyces israelii		83.96	0.51	SAL 92		
Bifidobacterium adolescentis		12.89	0.31	CEL 93		
Propionibacterium propionicum		2.52	0.24			
27537540	**Acceptable identification**					**27537540**
Actinomyces israelii		92.84	0.41	SAL 92		
Propionibacterium propionicum		7.16	0.21	CEL 93		
27537542	**Low discrimination**					**27537542**
Actinomyces israelii		88.82	0.44	SAL 92		
Propionibacterium propionicum		6.85	0.24	CEL 93		
Bifidobacterium adolescentis		4.33	0.16			
27540002	**Very good identification**					**27540002**
Actinomyces viscosus		99.98	0.64	FRU 95		
Actinomyces naeslundii		0.02	0.24			
27540003	**Good identification**					**27540003**
Actinomyces viscosus		99.92	0.56	FRU 95		
Actinomyces naeslundii		0.09	0.27			
27540102	**Very good identification**					**27540102**
Actinomyces viscosus		99.97	0.68	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.03	0.32			
27540103	**Very good identification**					**27540103**
Actinomyces viscosus		99.86	0.60	FRU 95		
Actinomyces naeslundii		0.14	0.35			
27540121	**Good identification**					**27540121**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27541001	**Low discrimination**					**27541001**
Actinomyces naeslundii		76.78	0.51	FRU 95		
Actinomyces viscosus		23.22	0.18			
27541002	**Very good identification**					**27541002**
Actinomyces viscosus		99.26	0.60	FRU 95		
Actinomyces naeslundii		0.74	0.48			
27541003	**Good identification**					**27541003**
Actinomyces viscosus		96.77	0.52	FRU 95		
Actinomyces naeslundii		3.23	0.51			
27541020	**Acceptable identification**					**27541020**
					GRA	
Actinomyces naeslundii		90.11	0.48	FRU 95	99	
Prevotella melaninogenica		6.25	-.01		1	
Clostridium perfringens		3.65	0.02		99	
27541021	**Good identification**					**27541021**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27541023	**Low discrimination**					**27541023**
Actinomyces naeslundii		76.78	0.51	FRU 95		
Actinomyces viscosus		23.22	0.18			
27541040	**Good identification**					**27541040**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27541041	**Good identification**					**27541041**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27541060	**Good identification**					**27541060**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27541061	**Good identification**					**27541061**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27541062	**Good identification**					**27541062**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27541063	**Good identification**					**27541063**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27541100	**Low discrimination**					**27541100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		53.35	0.56	FRU 95	99		
Actinomyces viscosus		45.35	0.31		99		
Prevotella melaninogenica		1.30	-.01		1		
27541101	**Low discrimination**						**27541101**
					SPO		
Actinomyces naeslundii		73.90	0.59	FRU 95	1		
Actinomyces viscosus		13.96	0.22		1		
Clostridium chauvoei		12.14	0.22		99		
27541102	**Very good identification**						**27541102**
					GRA		
Actinomyces viscosus		98.79	0.65	FRU 95	99		
Actinomyces naeslundii		1.17	0.56		99		
Capnocytophaga ochracea		0.04	0.05		1		
27541103	**Good identification**						**27541103**
					SPO		
Actinomyces viscosus		94.79	0.57	FRU 95	1		
Actinomyces naeslundii		5.07	0.59		1		
Clostridium chauvoei		0.15	0.10		99		
27541111	**Good identification**						**27541111**
Actinomyces naeslundii		100.0	0.37	FRU 95			
27541121	**Good identification**						**27541121**
					GRA	SPO	
Actinomyces naeslundii		93.37	0.59	FRU 95	99	1	
Clostridium chauvoei		5.11	0.14		99	99	
Prevotella melaninogenica		1.52	-.01		1	1	
27541122	**Low discrimination**						**27541122**
					GRA		
Actinomyces naeslundii		36.01	0.56	FRU 95	99		
Actinomyces viscosus		30.61	0.31		99		
Prevotella melaninogenica		16.54	0.21		1		
27541123	**Low discrimination**						**27541123**
					SPO		
Actinomyces naeslundii		83.43	0.59	FRU 95	1		
Actinomyces viscosus		15.76	0.22		1		
Clostridium chauvoei		0.81	0.01		99		
27541131	**Good identification**						**27541131**
Actinomyces naeslundii		100.0	0.37	FRU 95			
27541133	**Good identification**						**27541133**
Actinomyces naeslundii		100.0	0.37	FRU 95			
27541140	**Good identification**						**27541140**
Actinomyces naeslundii		100.0	0.43	FRU 95			
27541141	**Good identification**						**27541141**
Actinomyces naeslundii		100.0	0.46	FRU 95			
27541160	**Low discrimination**						**27541160**
					GRA		
Actinomyces naeslundii		79.97	0.43	FRU 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		11.04	-.01		1	
Clostridium perfringens		9.00	0.04		99	
27541161	**Good identification**					**27541161**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27541162	**Low discrimination**					**27541162**
					GRA	
Actinomyces naeslundii		84.28	0.43	FRU 95	99	
Capnocytophaga ochracea		15.72	0.05		1	
27541163	**Good identification**					**27541163**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27542100	**Low discrimination**					**27542100**
Actinomyces naeslundii		57.57	0.37	FRU 95		
Propionibacterium propionicum		32.46	0.23	GAL 96		
Actinomyces viscosus		9.97	0.00			
27542101	**Good identification**					**27542101**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27542120	**Good identification**					**27542120**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27542121	**Good identification**					**27542121**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27542122	**Low discrimination**					**27542122**
Actinomyces naeslundii		85.24	0.37	FRU 95		
Actinomyces viscosus		14.76	0.00	GAL 96		
27542123	**Good identification**					**27542123**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27543000	**Good identification**					**27543000**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27543001	**Good identification**					**27543001**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27543002	**Low discrimination**					**27543002**
Actinomyces naeslundii		58.79	0.53	FRU 95		
Actinomyces viscosus		41.21	0.26			
27543003	**Low discrimination**					**27543003**
Actinomyces naeslundii		86.52	0.56	FRU 95		
Actinomyces viscosus		13.48	0.18			
27543020	**Good identification**					**27543020**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27543021	**Good identification**					**27543021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.56	FRU 95		
27543022	**Good identification**					**27543022**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27543023	**Good identification**					**27543023**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27543040	**Good identification**					**27543040**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27543041	**Good identification**					**27543041**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27543042	**Good identification**					**27543042**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27543043	**Good identification**					**27543043**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27543060	**Good identification**					**27543060**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27543061	**Good identification**					**27543061**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27543062	**Good identification**					**27543062**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27543063	**Good identification**					**27543063**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27543100	**Very good identification**					**27543100**
Actinomyces naeslundii		97.71	0.61	FRU 95		
Propionibacterium propionicum		2.30	0.23			
27543101	**Very good identification**					**27543101**
Actinomyces naeslundii		100.0	0.64	FRU 95		
27543102	**Low discrimination**					**27543102**
Actinomyces naeslundii		67.88	0.61	FRU 95		
Actinomyces viscosus		29.73	0.31			
Propionibacterium propionicum		2.39	0.26			
27543103	**Good identification**					**27543103**
Actinomyces naeslundii		91.13	0.64	FRU 95		
Actinomyces viscosus		8.87	0.22			
27543110	**Good identification**					**27543110**
Actinomyces naeslundii		100.0	0.39	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27543111	**Good identification**					**27543111**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27543112	**Good identification**					**27543112**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27543113	**Good identification**					**27543113**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27543120	**Very good identification**					**27543120**
					GRA	
Actinomyces naeslundii		98.76	0.61	FRU 95	99	
Prevotella melaninogenica		1.24	-.01		1	
27543121	**Very good identification**					**27543121**
Actinomyces naeslundii		100.0	0.64	FRU 95		
27543122	**Very good identification**					**27543122**
Actinomyces naeslundii		100.0	0.61	FRU 95		
27543123	**Very good identification**					**27543123**
Actinomyces naeslundii		100.0	0.64	FRU 95		
27543130	**Good identification**					**27543130**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27543131	**Good identification**					**27543131**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27543132	**Good identification**					**27543132**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27543133	**Good identification**					**27543133**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27543140	**Low discrimination**					**27543140**
Actinomyces naeslundii		88.25	0.48	FRU 95		
Propionibacterium propionicum		11.75	0.23			
27543141	**Good identification**					**27543141**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27543142	**Low discrimination**					**27543142**
Actinomyces naeslundii		83.36	0.48	FRU 95		
Propionibacterium propionicum		16.64	0.26			
27543143	**Good identification**					**27543143**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27543160	**Good identification**					**27543160**
Actinomyces naeslundii		100.0	0.48	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27543161	**Good identification**					**27543161**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27543162	**Good identification**					**27543162**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27543163	**Good identification**					**27543163**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27550000	**Good identification**					**27550000**
					GRA	
Actinomyces viscosus		96.64	0.52	BGL 99	99	
Actinomyces naeslundii		1.81	0.46		99	
Prevotella intermedia		0.58	0.15		1	
27550001	**Acceptable identification**					**27550001**
Actinomyces viscosus		92.21	0.44	BGL 99		
Actinomyces naeslundii		7.79	0.49			
27550002	**Very good identification**					**27550002**
Actinomyces viscosus		99.98	0.86			
Actinomyces naeslundii		0.02	0.46			
Actinomyces odontolyticus		0.00	0.09			
27550003	**Very good identification**					**27550003**
Actinomyces viscosus		99.92	0.78			
Actinomyces naeslundii		0.09	0.49			
27550012	**Good identification**					**27550012**
Actinomyces viscosus		99.90	0.52	MLZ 1		
Actinomyces naeslundii		0.10	0.24			
27550013	**Good identification**					**27550013**
Actinomyces viscosus		99.56	0.44	MLZ 1		
Actinomyces naeslundii		0.44	0.27			
27550020	**Low discrimination**					**27550020**
Actinomyces naeslundii		62.92	0.46	GAL 96		
Actinomyces viscosus		33.87	0.18			
Clostridium perfringens		3.22	0.02			
27550021	**Low discrimination**					**27550021**
Actinomyces naeslundii		89.32	0.49	GAL 96		
Actinomyces viscosus		10.68	0.09			
27550022	**Good identification**					**27550022**
Actinomyces viscosus		98.16	0.52	NAG 1		
Actinomyces naeslundii		1.84	0.46			
27550023	**Acceptable identification**					**27550023**
Actinomyces viscosus		92.21	0.44	NAG 1		
Actinomyces naeslundii		7.79	0.49			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27550041	**Low discrimination**					**27550041**
Actinomyces naeslundii		59.60	0.36	GAL 96		
Actinomyces viscosus		40.40	0.09			
27550042	**Good identification**					**27550042**
Actinomyces viscosus		99.67	0.52	SOR 1		
Actinomyces naeslundii		0.33	0.33			
27550043	**Good identification**					**27550043**
Actinomyces viscosus		98.53	0.44	SOR 1		
Actinomyces naeslundii		1.47	0.36			
27550061	**Good identification**					**27550061**
Actinomyces naeslundii		100.0	0.36	GAL 96		
27550063	**Low discrimination**					**27550063**
Actinomyces naeslundii		59.60	0.36	GAL 96		
Actinomyces viscosus		40.40	0.09			
27550100	**Good identification**					**27550100**
					GRA	
Actinomyces viscosus		96.31	0.56	BGL 99	99	
Actinomyces naeslundii		2.89	0.54		99	
Fusobacterium mortiferum		0.63	0.31		1	
27550101	**Low discrimination**					**27550101**
Actinomyces viscosus		88.09	0.48	BGL 99		
Actinomyces naeslundii		11.91	0.57			
27550102	**Very good identification**					**27550102**
Actinomyces viscosus		99.97	0.90			
Actinomyces naeslundii		0.03	0.54			
Propionibacterium propionicum		0.00	0.26			
27550103	**Very good identification**					**27550103**
Actinomyces viscosus		99.86	0.82			
Actinomyces naeslundii		0.14	0.57			
27550112	**Good identification**					**27550112**
Actinomyces viscosus		99.84	0.56	MLZ 1		
Actinomyces naeslundii		0.16	0.32			
27550113	**Good identification**					**27550113**
Actinomyces viscosus		99.29	0.48	MLZ 1		
Actinomyces naeslundii		0.71	0.35			
27550120	**Low discrimination**					**27550120**
					GRA	
Actinomyces naeslundii		55.98	0.54	GAL 96	99	
Actinomyces viscosus		18.82	0.22		99	
Prevotella melaninogenica		15.50	0.15		1	
27550121	**Good identification**					**27550121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		93.05	0.57	GAL 96		
Actinomyces viscosus		6.95	0.14			
27550122	**Good identification**					**27550122**
					GRA	
Actinomyces viscosus		95.30	0.56	NAG 1	99	
Actinomyces naeslundii		2.86	0.54		99	
Capnocytophaga ochracea		1.07	0.21		1	
27550123	**Low discrimination**					**27550123**
Actinomyces viscosus		88.09	0.48	NAG 1		
Actinomyces naeslundii		11.91	0.57			
27550131	**Good identification**					**27550131**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27550141	**Low discrimination**					**27550141**
Actinomyces naeslundii		70.26	0.44	GAL 96		
Actinomyces viscosus		29.75	0.14			
27550142	**Good identification**					**27550142**
Actinomyces viscosus		99.22	0.56	SOR 1		
Actinomyces naeslundii		0.53	0.41			
Propionibacterium propionicum		0.26	0.26			
27550143	**Good identification**					**27550143**
Actinomyces viscosus		97.67	0.48	SOR 1		
Actinomyces naeslundii		2.33	0.44			
27550160	**Low discrimination**					**27550160**
Actinomyces naeslundii		87.56	0.41	GAL 96		
Clostridium perfringens		12.45	0.04			
27550161	**Good identification**					**27550161**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27550163	**Low discrimination**					**27550163**
Actinomyces naeslundii		70.26	0.44	GAL 96		
Actinomyces viscosus		29.75	0.14			
27550202	**Good identification**					**27550202**
Actinomyces viscosus		99.75	0.52	RHA 1		
Actinomyces odontolyticus		0.25	0.14			
27550203	**Good identification**					**27550203**
Actinomyces viscosus		100.0	0.44	RHA 1		
27550302	**Good identification**					**27550302**
Actinomyces viscosus		99.96	0.56	RHA 1		
Actinomyces odontolyticus		0.05	0.06			
27550303	**Good identification**					**27550303**
Actinomyces viscosus		99.86	0.48	RHA 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.14	0.23			
27550402	**Good identification**					**27550402**
Actinomyces viscosus		100.0	0.52	ARA 1		
27550403	**Good identification**					**27550403**
Actinomyces viscosus		100.0	0.44	ARA 1		
27550502	**Good identification**					**27550502**
Actinomyces viscosus		100.0	0.56	ARA 1		
27550503	**Good identification**					**27550503**
Actinomyces viscosus		99.86	0.48	ARA 1		
Actinomyces naeslundii		0.14	0.23			
27551000	**Low discrimination**					**27551000**
					GRA	
Actinomyces viscosus		56.32	0.48	BGL 99	99	
Actinomyces naeslundii		41.38	0.70		99	
Fusobacterium mortiferum		0.69	0.28		1	
27551001	**Low discrimination**					**27551001**
Actinomyces naeslundii		76.78	0.73			
Actinomyces viscosus		23.22	0.40			
27551002	**Very good identification**					**27551002**
Actinomyces viscosus		99.24	0.82			
Actinomyces naeslundii		0.74	0.70			
Actinomyces israelii		0.00	0.21			
27551003	**Very good identification**					**27551003**
Actinomyces viscosus		96.77	0.74			
Actinomyces naeslundii		3.23	0.73			
27551010	**Low discrimination**					**27551010**
Actinomyces naeslundii		79.29	0.48	MLZ 5		
Actinomyces viscosus		20.71	0.14			
27551011	**Acceptable identification**					**27551011**
Actinomyces naeslundii		94.51	0.51	MLZ 5		
Actinomyces viscosus		5.49	0.06			
27551012	**Good identification**					**27551012**
Actinomyces viscosus		96.28	0.48	MLZ 1		
Actinomyces naeslundii		3.72	0.48			
27551013	**Low discrimination**					**27551013**
Actinomyces viscosus		85.18	0.40	MLZ 1		
Actinomyces naeslundii		14.82	0.51			
27551020	**Low discrimination**					**27551020**
					GRA	
Actinomyces naeslundii		78.84	0.70		99	
Clostridium perfringens		16.62	0.36		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		2.59	0.15		1	
27551021	**Very good identification**					**27551021**
Actinomyces naeslundii		99.55	0.73			
Actinomyces viscosus		0.30	0.06			
Clostridium perfringens		0.14	0.02			
27551022	**Low discrimination**					**27551022**
					GRA	
Actinomyces viscosus		51.82	0.48	NAG 1	99	
Actinomyces naeslundii		38.07	0.70		99	
Prevotella oralis		4.40	0.28		1	
27551023	**Low discrimination**					**27551023**
Actinomyces naeslundii		76.78	0.73			
Actinomyces viscosus		23.22	0.40			
27551030	**Good identification**					**27551030**
Actinomyces naeslundii		96.11	0.48	MLZ 5		
Clostridium perfringens		3.89	0.02			
27551031	**Good identification**					**27551031**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
27551032	**Low discrimination**					**27551032**
					GRA	
Actinomyces naeslundii		73.81	0.48	MLZ 5	99	
Actinomyces viscosus		19.28	0.14		99	
Capnocytophaga ochracea		6.91	0.05		1	
27551033	**Acceptable identification**					**27551033**
Actinomyces naeslundii		94.51	0.51	MLZ 5		
Actinomyces viscosus		5.49	0.06			
27551040	**Good identification**					**27551040**
Actinomyces naeslundii		92.77	0.57			
Actinomyces viscosus		7.23	0.14			
27551041	**Very good identification**					**27551041**
Actinomyces naeslundii		98.30	0.60			
Actinomyces viscosus		1.70	0.06			
27551042	**Low discrimination**					**27551042**
Actinomyces viscosus		88.42	0.48	SOR 1		
Actinomyces naeslundii		11.47	0.57			
Actinomyces israelii		0.12	0.14			
27551043	**Low discrimination**					**27551043**
Actinomyces viscosus		63.15	0.40	SOR 1		
Actinomyces naeslundii		36.85	0.60			
27551050	**Good identification**					**27551050**
Actinomyces naeslundii		100.0	0.35	MLZ 5		
27551051	**Good identification**					**27551051**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MLZ 5		
27551053	**Low discrimination**					**27551053**
Actinomyces naeslundii		75.25	0.38	MLZ 5		
Actinomyces viscosus		24.75	0.06			
27551060	**Low discrimination**					**27551060**
Actinomyces naeslundii		81.46	0.57			
Clostridium perfringens		18.54	0.24			
27551061	**Very good identification**					**27551061**
Actinomyces naeslundii		100.0	0.60			
27551062	**Low discrimination**					**27551062**
Actinomyces naeslundii		82.44	0.57			
Clostridium perfringens		8.83	0.18			
Actinomyces viscosus		6.42	0.14			
27551063	**Very good identification**					**27551063**
Actinomyces naeslundii		98.30	0.60			
Actinomyces viscosus		1.70	0.06			
27551070	**Good identification**					**27551070**
Actinomyces naeslundii		100.0	0.35	MLZ 5		
27551071	**Good identification**					**27551071**
Actinomyces naeslundii		100.0	0.38	MLZ 5		
27551072	**Good identification**					**27551072**
Actinomyces naeslundii		100.0	0.35	MLZ 5		
27551073	**Good identification**					**27551073**
Actinomyces naeslundii		100.0	0.38	MLZ 5		
27551100	**Low discrimination**					**27551100**
					GRA	
Actinomyces naeslundii		52.02	0.78		99	
Actinomyces viscosus		44.22	0.52		99	
Capnocytophaga ochracea		0.99	0.23		1	
27551101	**Low discrimination**					**27551101**
					SPO	
Actinomyces naeslundii		82.71	0.81		1	
Actinomyces viscosus		15.62	0.44		1	
Clostridium chauvoei		1.67	0.29		99	
27551102	**Very good identification**					**27551102**
					GRA	
Actinomyces viscosus		98.56	0.87		99	
Actinomyces naeslundii		1.17	0.78		99	
Capnocytophaga ochracea		0.20	0.39		1	
27551103	**Good identification**					**27551103**
					SPO	
Actinomyces viscosus		94.90	0.78		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		5.08	0.81		1		
Clostridium chauvoei		0.02	0.16		99		
27551110	**Low discrimination**						**27551110**
					GRA		
Actinomyces naeslundii		82.26	0.56	MLZ 5	99		
Actinomyces viscosus		13.42	0.18		99		
Capnocytophaga ochracea		3.31	0.07		1		
27551111	**Very good identification**						**27551111**
Actinomyces naeslundii		96.50	0.59	MLZ 5			
Actinomyces viscosus		3.50	0.10				
27551112	**Acceptable identification**						**27551112**
					GRA		
Actinomyces viscosus		91.97	0.52	MLZ 1	99		
Actinomyces naeslundii		5.69	0.56		99		
Capnocytophaga ochracea		2.06	0.23		1		
27551113	**Low discrimination**						**27551113**
Actinomyces viscosus		78.21	0.44	MLZ 1			
Actinomyces naeslundii		21.79	0.59				
27551120	**Low discrimination**						**27551120**
					GRA		
Prevotella melaninogenica		39.38	0.50	NIT 1	1		
Actinomyces naeslundii		34.49	0.78	CEL 10	99		
Clostridium perfringens		18.74	0.51		99		
27551121	**Very good identification**						**27551121**
					GRA	SPO	
Actinomyces naeslundii		97.93	0.81		99	1	
Prevotella melaninogenica		0.75	0.15		1	1	
Clostridium chauvoei		0.66	0.21		99	99	
27551122	**Low discrimination**						**27551122**
					GRA		
Capnocytophaga ochracea		34.21	0.56	ESL 99	1		
Actinomyces naeslundii		22.16	0.78		99		
Actinomyces viscosus		18.84	0.52		99		
27551123	**Low discrimination**						**27551123**
					GRA		
Actinomyces naeslundii		82.78	0.81		99		
Actinomyces viscosus		15.64	0.44		99		
Capnocytophaga ochracea		0.86	0.21		1		
27551130	**Low discrimination**						**27551130**
					GRA	CAT	
Actinomyces naeslundii		59.33	0.56	MLZ 5	99	25	
Capnocytophaga ochracea		21.48	0.23		1	90	
Prevotella melaninogenica		13.00	0.15		1	1	
27551131	**Very good identification**						**27551131**
Actinomyces naeslundii		100.0	0.59	MLZ 5			
27551132	**Low discrimination**						**27551132**
					GRA		
Capnocytophaga ochracea		70.00	0.39	ESL 99	1		
Actinomyces naeslundii		21.48	0.56	MLZ 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		3.51	0.18		99		
27551133	**Good identification**						**27551133**
					GRA		
Actinomyces naeslundii		94.50	0.59	MLZ 5	99		
Actinomyces viscosus		3.43	0.10		99		
Capnocytophaga ochracea		2.07	0.05		1		
27551140	**Good identification**						**27551140**
Actinomyces naeslundii		90.75	0.65				
Actinomyces viscosus		4.42	0.18				
Clostridium perfringens		1.65	0.13				
27551141	**Very good identification**						**27551141**
Actinomyces naeslundii		98.93	0.68				
Actinomyces viscosus		1.07	0.10				
27551142	**Low discrimination**						**27551142**
					GRA		
Actinomyces viscosus		80.71	0.52	SOR 1	99		
Actinomyces naeslundii		16.76	0.65		99		
Capnocytophaga ochracea		1.81	0.23		1		
27551143	**Low discrimination**						**27551143**
Actinomyces viscosus		51.70	0.44	SOR 1			
Actinomyces naeslundii		48.30	0.68				
27551150	**Good identification**						**27551150**
Actinomyces naeslundii		100.0	0.43	MLZ 5			
27551151	**Good identification**						**27551151**
Actinomyces naeslundii		100.0	0.46	MLZ 5			
27551152	**Low discrimination**						**27551152**
					GRA		
Actinomyces naeslundii		46.46	0.43	MLZ 5	99		
Actinomyces viscosus		42.95	0.18		99		
Capnocytophaga ochracea		10.59	0.07		1		
27551153	**Low discrimination**						**27551153**
Actinomyces naeslundii		82.96	0.46	MLZ 5			
Actinomyces viscosus		17.04	0.10				
27551160	**Low discrimination**						**27551160**
					GRA	CAT	
Actinomyces naeslundii		56.83	0.65		99	25	
Clostridium perfringens		33.33	0.39		99	1	
Capnocytophaga ochracea		6.14	0.23		1	90	
27551161	**Very good identification**						**27551161**
Actinomyces naeslundii		99.61	0.68				
Clostridium perfringens		0.39	0.04				
27551162	**Low discrimination**						**27551162**
					GRA	CAT	
Actinomyces naeslundii		42.64	0.65		99	25	
Capnocytophaga ochracea		41.44	0.39		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		11.77	0.33		99	1	
27551163	**Very good identification**						**27551163**
					GRA		
Actinomyces naeslundii		98.29	0.68		99		
Actinomyces viscosus		1.06	0.10		99		
Capnocytophaga ochracea		0.64	0.05		1		
27551170	**Low discrimination**						**27551170**
					GRA	CAT	
Actinomyces naeslundii		74.60	0.43	MLZ 5	99	25	
Capnocytophaga ochracea		17.01	0.07		1	90	
Clostridium perfringens		8.40	0.04		99	1	
27551171	**Good identification**						**27551171**
Actinomyces naeslundii		100.0	0.46	MLZ 5			
27551173	**Good identification**						**27551173**
Actinomyces naeslundii		100.0	0.46	MLZ 5			
27551201	**Low discrimination**						**27551201**
Actinomyces naeslundii		76.78	0.39	RHA 1			
Actinomyces viscosus		23.22	0.06				
27551202	**Good identification**						**27551202**
Actinomyces viscosus		98.80	0.48	RHA 1			
Actinomyces naeslundii		0.73	0.36				
Actinomyces odontolyticus		0.27	0.11				
27551203	**Good identification**						**27551203**
Actinomyces viscosus		96.77	0.40	RHA 1			
Actinomyces naeslundii		3.23	0.39				
27551220	**Low discrimination**						**27551220**
					GRA		
Actinomyces naeslundii		68.07	0.36	RHA 1	99		
Prevotella oralis		17.58	0.00		1		
Clostridium perfringens		14.35	0.02		99		
27551221	**Good identification**						**27551221**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27551223	**Low discrimination**						**27551223**
Actinomyces naeslundii		76.78	0.39	RHA 1			
Actinomyces viscosus		23.22	0.06				
27551300	**Low discrimination**						**27551300**
					GRA		
Actinomyces naeslundii		50.20	0.44	RHA 1	99		
Actinomyces viscosus		42.67	0.18		99		
Prevotella oralis		3.76	-.01		1		
27551301	**Low discrimination**						**27551301**
Actinomyces naeslundii		84.11	0.47	RHA 1			
Actinomyces viscosus		15.89	0.10				
27551302	**Good identification**						**27551302**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				GRA		
Actinomyces viscosus	97.01	0.52	RHA 1	99		
Prevotella oralis	1.64	0.21		1		
Actinomyces naeslundii	1.15	0.44		99		
27551303 Acceptable identification						**27551303**
Actinomyces viscosus	94.92	0.44	RHA 1			
Actinomyces naeslundii	5.08	0.47				
27551321 Good identification						**27551321**
Actinomyces naeslundii	100.0	0.47	RHA 1			
27551322 Low discrimination						**27551322**
				GRA	CAT	
Prevotella oralis	85.82	0.42	NIT 1	1	1	
Capnocytophaga ochracea	4.90	0.21		1	90	
Actinomyces naeslundii	3.18	0.44		99	25	
27551323 Low discrimination						**27551323**
				GRA		
Actinomyces naeslundii	72.95	0.47	RHA 1	99		
Actinomyces viscosus	13.78	0.10		99		
Prevotella oralis	13.27	0.08		1		
27551401 Low discrimination						**27551401**
Actinomyces naeslundii	76.78	0.39	ARA 1			
Actinomyces viscosus	23.22	0.06				
27551402 Good identification						**27551402**
Actinomyces viscosus	98.66	0.48	ARA 1			
Actinomyces naeslundii	0.73	0.36				
Lactobacillus fermentum	0.33	0.20				
27551403 Good identification						**27551403**
Actinomyces viscosus	96.77	0.40	ARA 1			
Actinomyces naeslundii	3.23	0.39				
27551420 Low discrimination						**27551420**
Actinomyces naeslundii	82.58	0.36	ARA 1			
Clostridium perfringens	17.42	0.02				
27551421 Good identification						**27551421**
Actinomyces naeslundii	100.0	0.39	ARA 1			
27551423 Low discrimination						**27551423**
Actinomyces naeslundii	76.78	0.39	ARA 1			
Actinomyces viscosus	23.22	0.06				
27551500 Low discrimination						**27551500**
Actinomyces naeslundii	34.55	0.44	ARA 1			
Actinomyces viscosus	29.37	0.18				
Actinomyces israelii	12.14	0.27				
27551501 Low discrimination						**27551501**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.11	0.47	ARA 1		
Actinomyces viscosus		15.89	0.10			
27551502	**Acceptable identification**					**27551502**
					GRA	
Actinomyces viscosus		93.60	0.52	ARA 1	99	
Capnocytophaga ochracea		3.33	0.26		1	
Actinomyces naeslundii		1.11	0.44		99	
27551503	**Acceptable identification**					**27551503**
Actinomyces viscosus		94.92	0.44	ARA 1		
Actinomyces naeslundii		5.08	0.47			
27551521	**Good identification**					**27551521**
Actinomyces naeslundii		100.0	0.47	ARA 1		
27551522	**Low discrimination**					**27551522**
					GRA	
Capnocytophaga ochracea		88.32	0.43	ESL 99	1	
Actinomyces naeslundii		3.28	0.44		99	
Actinomyces viscosus		2.78	0.18		99	
27551523	**Low discrimination**					**27551523**
					GRA	
Actinomyces naeslundii		72.97	0.47	ARA 1	99	
Actinomyces viscosus		13.78	0.10		99	
Capnocytophaga ochracea		13.25	0.09		1	
27552000	**Low discrimination**					**27552000**
Actinomyces naeslundii		70.34	0.51	GAL 96		
Actinomyces viscosus		19.50	0.18			
Propionibacterium propionicum		5.94	0.23			
27552001	**Acceptable identification**					**27552001**
Actinomyces naeslundii		94.20	0.54	GAL 96		
Actinomyces viscosus		5.81	0.09			
27552002	**Good identification**					**27552002**
Actinomyces viscosus		96.06	0.52	MAN 1		
Actinomyces naeslundii		3.50	0.51			
Propionibacterium propionicum		0.44	0.26			
27552003	**Low discrimination**					**27552003**
Actinomyces viscosus		85.92	0.44	MAN 1		
Actinomyces naeslundii		14.08	0.54			
27552020	**Good identification**					**27552020**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27552021	**Good identification**					**27552021**
Actinomyces naeslundii		100.0	0.54	GAL 96		
27552022	**Low discrimination**					**27552022**
Actinomyces naeslundii		78.29	0.51	GAL 96		
Actinomyces viscosus		21.71	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27552023	**Acceptable identification**					**27552023**
Actinomyces naeslundii		94.20	0.54	GAL 96		
Actinomyces viscosus		5.81	0.09			
27552040	**Low discrimination**					**27552040**
Actinomyces naeslundii		67.63	0.38	GAL 96		
Propionibacterium propionicum		32.37	0.23			
27552041	**Good identification**					**27552041**
Actinomyces naeslundii		100.0	0.41	GAL 96		
27552043	**Low discrimination**					**27552043**
Actinomyces naeslundii		74.12	0.41	GAL 96		
Actinomyces viscosus		25.88	0.09			
27552060	**Good identification**					**27552060**
Actinomyces naeslundii		100.0	0.38	GAL 96		
27552061	**Good identification**					**27552061**
Actinomyces naeslundii		100.0	0.41	GAL 96		
27552062	**Good identification**					**27552062**
Actinomyces naeslundii		100.0	0.38	GAL 96		
27552063	**Good identification**					**27552063**
Actinomyces naeslundii		100.0	0.41	GAL 96		
27552100	**Low discrimination**					**27552100**
Propionibacterium propionicum		71.46	0.57	CEL 1		
Actinomyces naeslundii		24.33	0.59			
Actinomyces viscosus		4.21	0.22			
27552101	**Low discrimination**					**27552101**
Actinomyces naeslundii		79.61	0.62	GAL 96		
Propionibacterium propionicum		17.32	0.41			
Actinomyces viscosus		3.06	0.14			
27552102	**Low discrimination**					**27552102**
Actinomyces viscosus		76.03	0.56	MAN 1		
Propionibacterium propionicum		19.54	0.60			
Actinomyces naeslundii		4.43	0.59			
27552103	**Low discrimination**					**27552103**
Actinomyces viscosus		74.18	0.48	MAN 1		
Actinomyces naeslundii		19.47	0.62			
Propionibacterium propionicum		6.35	0.44			
27552110	**Low discrimination**					**27552110**
Actinomyces naeslundii		63.95	0.37	GAL 96		
Propionibacterium propionicum		36.05	0.23	MLZ 5		
27552111	**Good identification**					**27552111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	GAL 96		
27552113	**Low discrimination**					**27552113**
Actinomyces naeslundii		57.76	0.40	GAL 96		
Actinomyces viscosus		42.24	0.14	MLZ 5		
27552120	**Very good identification**					**27552120**
Actinomyces naeslundii		97.12	0.59	GAL 96		
Propionibacterium propionicum		2.88	0.23			
27552121	**Very good identification**					**27552121**
Actinomyces naeslundii		100.0	0.62	GAL 96		
27552122	**Low discrimination**					**27552122**
Actinomyces naeslundii		82.12	0.59	GAL 96		
Actinomyces viscosus		14.22	0.22			
Propionibacterium propionicum		3.66	0.26			
27552123	**Very good identification**					**27552123**
Actinomyces naeslundii		96.29	0.62	GAL 96		
Actinomyces viscosus		3.71	0.14			
27552130	**Good identification**					**27552130**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27552131	**Good identification**					**27552131**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27552132	**Good identification**					**27552132**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27552133	**Good identification**					**27552133**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27552140	**Good identification**					**27552140**
Propionibacterium propionicum		94.33	0.57	CEL 1		
Actinomyces naeslundii		5.67	0.46			
27552141	**Low discrimination**					**27552141**
Propionibacterium propionicum		55.22	0.41	CEL 1		
Actinomyces naeslundii		44.78	0.49	URE 10		
27552142	**Good identification**					**27552142**
Propionibacterium propionicum		92.65	0.60	CEL 1		
Actinomyces naeslundii		3.71	0.46			
Actinomyces viscosus		3.64	0.22			
27552143	**Low discrimination**					**27552143**
Propionibacterium propionicum		60.29	0.44	CEL 1		
Actinomyces naeslundii		32.60	0.49	URE 10		
Actinomyces viscosus		7.11	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27552160	**Low discrimination**					**27552160**
Actinomyces naeslundii		85.61	0.46	GAL 96		
Propionibacterium propionicum		14.39	0.23			
27552161	**Good identification**					**27552161**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27552162	**Low discrimination**					**27552162**
Actinomyces naeslundii		79.86	0.46	GAL 96		
Propionibacterium propionicum		20.14	0.26			
27552163	**Good identification**					**27552163**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27553000	**Very good identification**					**27553000**
Actinomyces naeslundii		97.84	0.75			
Actinomyces israelii		0.88	0.31			
Actinomyces viscosus		0.69	0.14			
27553001	**Very good identification**					**27553001**
Actinomyces naeslundii		99.84	0.78			
Actinomyces viscosus		0.16	0.06			
27553002	**Low discrimination**					**27553002**
Actinomyces naeslundii		58.14	0.75			
Actinomyces viscosus		40.76	0.48			
Actinomyces israelii		0.79	0.34			
27553003	**Low discrimination**					**27553003**
Actinomyces naeslundii		86.52	0.78			
Actinomyces viscosus		13.48	0.40			
27553010	**Good identification**					**27553010**
Actinomyces naeslundii		97.51	0.53	MLZ 5		
Actinomyces israelii		2.49	0.17			
27553011	**Good identification**					**27553011**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27553012	**Low discrimination**					**27553012**
Actinomyces naeslundii		85.26	0.53	MLZ 5		
Actinomyces viscosus		11.47	0.14			
Actinomyces israelii		3.27	0.20			
27553013	**Good identification**					**27553013**
Actinomyces naeslundii		97.10	0.56	MLZ 5		
Actinomyces viscosus		2.90	0.06			
27553020	**Very good identification**					**27553020**
Actinomyces naeslundii		99.89	0.75			
Clostridium perfringens		0.11	0.02			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27553021	**Very good identification**					**27553021**
Actinomyces naeslundii		100.0	0.78			
27553022	**Very good identification**					**27553022**
Actinomyces naeslundii		99.21	0.75			
Actinomyces viscosus		0.70	0.14			
Actinomyces israelii		0.09	0.13			
27553023	**Very good identification**					**27553023**
Actinomyces naeslundii		99.84	0.78			
Actinomyces viscosus		0.16	0.06			
27553030	**Good identification**					**27553030**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27553031	**Good identification**					**27553031**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27553032	**Good identification**					**27553032**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27553033	**Good identification**					**27553033**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27553040	**Very good identification**					**27553040**
Actinomyces naeslundii		96.17	0.62			
Propionibacterium propionicum		1.92	0.23			
Actinomyces israelii		1.91	0.24			
27553041	**Very good identification**					**27553041**
Actinomyces naeslundii		100.0	0.65			
27553042	**Good identification**					**27553042**
Actinomyces naeslundii		90.92	0.62			
Actinomyces viscosus		3.65	0.14			
Propionibacterium propionicum		2.72	0.26			
27553043	**Very good identification**					**27553043**
Actinomyces naeslundii		99.12	0.65			
Actinomyces viscosus		0.88	0.06			
27553050	**Good identification**					**27553050**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27553051	**Good identification**					**27553051**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27553052	**Good identification**					**27553052**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27553053	**Good identification**					**27553053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27553060	**Very good identification**					**27553060**
Actinomyces naeslundii		100.0	0.62			
27553061	**Very good identification**					**27553061**
Actinomyces naeslundii		100.0	0.65			
27553062	**Very good identification**					**27553062**
Actinomyces naeslundii		100.0	0.62			
27553063	**Very good identification**					**27553063**
Actinomyces naeslundii		100.0	0.65			
27553070	**Good identification**					**27553070**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27553071	**Good identification**					**27553071**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27553072	**Good identification**					**27553072**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27553073	**Good identification**					**27553073**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27553100	**Low discrimination**					**27553100**
Actinomyces naeslundii		87.76	0.83			
Propionibacterium propionicum		10.74	0.57			
Actinomyces israelii		1.11	0.41			
27553101	**Very good identification**					**27553101**
Actinomyces naeslundii		99.01	0.86			
Propionibacterium propionicum		0.90	0.41			
Actinomyces viscosus		0.10	0.10			
27553102	**Low discrimination**					**27553102**
Actinomyces naeslundii		60.90	0.83			
Actinomyces viscosus		26.67	0.52			
Propionibacterium propionicum		11.18	0.60			
27553103	**Good identification**					**27553103**
Actinomyces naeslundii		90.02	0.86			
Actinomyces viscosus		8.76	0.44			
Propionibacterium propionicum		1.22	0.44			
27553110	**Good identification**					**27553110**
Actinomyces naeslundii		94.39	0.61	MLZ 5		
Actinomyces israelii		3.39	0.27			
Propionibacterium propionicum		2.22	0.23			
27553111	**Very good identification**					**27553111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.64	MLZ 5		
27553112	**Low discrimination**					**27553112**
Actinomyces naeslundii		84.32	0.61	MLZ 5		
Actinomyces viscosus		7.09	0.18			
Actinomyces israelii		4.55	0.30			
27553113	**Very good identification**					**27553113**
Actinomyces naeslundii		98.17	0.64	MLZ 5		
Actinomyces viscosus		1.83	0.10			
27553120	**Very good identification**					**27553120**
					GRA	
Actinomyces naeslundii		98.84	0.83		99	
Prevotella melaninogenica		0.59	0.15		1	
Clostridium perfringens		0.28	0.17		99	
27553121	**Very good identification**					**27553121**
Actinomyces naeslundii		100.0	0.86			
27553122	**Very good identification**					**27553122**
					GRA	
Actinomyces naeslundii		97.84	0.83		99	
Capnocytophaga ochracea		0.79	0.21		1	
Actinomyces viscosus		0.43	0.18		99	
27553123	**Very good identification**					**27553123**
Actinomyces naeslundii		99.90	0.86			
Actinomyces viscosus		0.10	0.10			
27553130	**Very good identification**					**27553130**
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27553131	**Very good identification**					**27553131**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
27553132	**Very good identification**					**27553132**
					GRA	
Actinomyces naeslundii		98.33	0.61	MLZ 5	99	
Capnocytophaga ochracea		1.67	0.05		1	
27553133	**Very good identification**					**27553133**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
27553140	**Low discrimination**					**27553140**
Actinomyces naeslundii		58.09	0.70			
Propionibacterium propionicum		40.29	0.57			
Actinomyces israelii		1.62	0.34			
27553141	**Very good identification**					**27553141**
Actinomyces naeslundii		95.11	0.73			
Propionibacterium propionicum		4.89	0.41			
27553142	**Low discrimination**					**27553142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		49.37	0.60	CEL 1			
Actinomyces naeslundii		47.46	0.70				
Actinomyces israelii		1.99	0.37				
27553143	**Good identification**						**27553143**
Actinomyces naeslundii		92.37	0.73				
Propionibacterium propionicum		7.12	0.44				
Actinomyces viscosus		0.51	0.10				
27553150	**Low discrimination**						**27553150**
Actinomyces naeslundii		82.49	0.48	MLZ 5			
Propionibacterium propionicum		10.98	0.23				
Actinomyces israelii		6.53	0.20				
27553151	**Good identification**						**27553151**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
27553152	**Low discrimination**						**27553152**
Actinomyces naeslundii		75.85	0.48	MLZ 5			
Propionibacterium propionicum		15.14	0.26				
Actinomyces israelii		9.01	0.23				
27553153	**Good identification**						**27553153**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
27553160	**Very good identification**						**27553160**
Actinomyces naeslundii		98.83	0.70				
Propionibacterium propionicum		0.69	0.23				
Clostridium perfringens		0.30	0.04				
27553161	**Very good identification**						**27553161**
Actinomyces naeslundii		100.0	0.73				
27553162	**Very good identification**						**27553162**
					GRA	CAT	
Actinomyces naeslundii		98.21	0.70		99	25	
Propionibacterium propionicum		1.03	0.26		99	1	
Capnocytophaga ochracea		0.50	0.05		1	90	
27553163	**Very good identification**						**27553163**
Actinomyces naeslundii		100.0	0.73				
27553170	**Good identification**						**27553170**
Actinomyces naeslundii		100.0	0.48	MLZ 5			
27553171	**Good identification**						**27553171**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
27553172	**Good identification**						**27553172**
Actinomyces naeslundii		100.0	0.48	MLZ 5			
27553173	**Good identification**						**27553173**
Actinomyces naeslundii		100.0	0.51	MLZ 5			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27553200	**Good identification**					**27553200**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27553201	**Good identification**					**27553201**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27553202	**Low discrimination**					**27553202**
Actinomyces naeslundii		55.50	0.41	RHA 1		
Actinomyces viscosus		38.91	0.14			
Actinomyces israelii		5.59	0.14			
27553203	**Low discrimination**					**27553203**
Actinomyces naeslundii		86.52	0.44	RHA 1		
Actinomyces viscosus		13.48	0.06			
27553220	**Good identification**					**27553220**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27553221	**Good identification**					**27553221**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27553222	**Good identification**					**27553222**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27553223	**Good identification**					**27553223**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27553300	**Low discrimination**					**27553300**
Actinomyces naeslundii		82.19	0.49	RHA 1		
Propionibacterium propionicum		10.06	0.23			
Actinomyces israelii		7.75	0.22			
27553301	**Good identification**					**27553301**
Actinomyces naeslundii		100.0	0.52	RHA 1		
27553302	**Low discrimination**					**27553302**
Actinomyces naeslundii		56.72	0.49	RHA 1		
Actinomyces viscosus		24.84	0.18			
Propionibacterium propionicum		10.41	0.26			
27553303	**Acceptable identification**					**27553303**
Actinomyces naeslundii		91.13	0.52	RHA 1		
Actinomyces viscosus		8.87	0.10			
27553320	**Good identification**					**27553320**
Actinomyces naeslundii		100.0	0.49	RHA 1		
27553321	**Good identification**					**27553321**
Actinomyces naeslundii		100.0	0.52	RHA 1		
27553322	**Low discrimination**					**27553322**

Code Taxon		ID score	T index	tests against	Differentiation tests SPO	 GRA	Code
Actinomyces naeslundii		80.12	0.49	RHA 1	1	99	
Prevotella oralis		11.26	0.08		1	1	
Clostridium ramosum		8.61	0.07		92	50	
27553323	**Good identification**						**27553323**
Actinomyces naeslundii		100.0	0.52	RHA 1			
27553340	**Low discrimination**						**27553340**
Actinomyces naeslundii		52.59	0.36	RHA 1			
Propionibacterium propionicum		36.47	0.23				
Actinomyces israelii		10.94	0.15				
27553341	**Good identification**						**27553341**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27553343	**Good identification**						**27553343**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27553360	**Good identification**						**27553360**
Actinomyces naeslundii		100.0	0.36	RHA 1			
27553361	**Good identification**						**27553361**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27553362	**Good identification**						**27553362**
Actinomyces naeslundii		100.0	0.36	RHA 1			
27553363	**Good identification**						**27553363**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27553400	**Low discrimination**						**27553400**
Actinomyces naeslundii		57.82	0.41	ARA 1			
Actinomyces israelii		42.18	0.29				
27553401	**Good identification**						**27553401**
Actinomyces naeslundii		100.0	0.44	ARA 1			
27553403	**Low discrimination**						**27553403**
Actinomyces naeslundii		86.52	0.44	ARA 1			
Actinomyces viscosus		13.48	0.06				
27553420	**Good identification**						**27553420**
Actinomyces naeslundii		100.0	0.41	ARA 1			
27553421	**Good identification**						**27553421**
Actinomyces naeslundii		100.0	0.44	ARA 1			
27553422	**Good identification**						**27553422**
Actinomyces naeslundii		100.0	0.41	ARA 1			
27553423	**Good identification**						**27553423**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	ARA 1			
27553500	**Low discrimination**						**27553500**
Actinomyces israelii		47.74	0.40	SAL 92			
Actinomyces naeslundii		46.56	0.49	XYL 92			
Propionibacterium propionicum		5.70	0.23				
27553501	**Good identification**						**27553501**
Actinomyces naeslundii		100.0	0.52	ARA 1			
27553502	**Low discrimination**						**27553502**
Actinomyces israelii		43.63	0.43	SAL 92			
Actinomyces naeslundii		28.36	0.49	XYL 92			
Actinomyces viscosus		12.42	0.18				
27553503	**Acceptable identification**						**27553503**
Actinomyces naeslundii		91.13	0.52	ARA 1			
Actinomyces viscosus		8.87	0.10				
27553520	**Acceptable identification**						**27553520**
Actinomyces naeslundii		93.86	0.49	ARA 1			
Actinomyces israelii		6.14	0.19				
27553521	**Good identification**						**27553521**
Actinomyces naeslundii		100.0	0.52	ARA 1			
27553522	**Low discrimination**						**27553522**
					GRA	CAT	
Actinomyces naeslundii		80.74	0.49	ARA 1	99	25	
Capnocytophaga ochracea		11.33	0.09		1	90	
Actinomyces israelii		7.93	0.22		99	1	
27553523	**Good identification**						**27553523**
Actinomyces naeslundii		100.0	0.52	ARA 1			
27553541	**Good identification**						**27553541**
Actinomyces naeslundii		100.0	0.39	ARA 1			
27553542	**Low discrimination**						**27553542**
Actinomyces israelii		59.61	0.36	SAL 92			
Propionibacterium propionicum		18.29	0.26	XYL 92			
Actinomyces naeslundii		17.59	0.36				
27553543	**Good identification**						**27553543**
Actinomyces naeslundii		100.0	0.39	ARA 1			
27553560	**Good identification**						**27553560**
Actinomyces naeslundii		100.0	0.36	ARA 1			
27553561	**Good identification**						**27553561**
Actinomyces naeslundii		100.0	0.39	ARA 1			
27553562	**Low discrimination**						**27553562**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		82.21	0.36	ARA 1		
Actinomyces israelii		17.79	0.15			
27553563	**Good identification**					**27553563**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27554002	**Good identification**					**27554002**
Actinomyces viscosus		100.0	0.52	XYL 1		
27554003	**Good identification**					**27554003**
Actinomyces viscosus		100.0	0.44	XYL 1		
27554102	**Good identification**					**27554102**
Actinomyces viscosus		99.95	0.56	XYL 1		
Actinomyces israelii		0.05	0.15			
27554103	**Good identification**					**27554103**
Actinomyces viscosus		99.86	0.48	XYL 1		
Actinomyces naeslundii		0.14	0.23			
27555000	**Low discrimination**					**27555000**
Actinomyces israelii		57.20	0.36	SAL 92		
Actinomyces viscosus		22.16	0.14			
Actinomyces naeslundii		16.28	0.36			
27555001	**Low discrimination**					**27555001**
Actinomyces naeslundii		76.78	0.39	XYL 1		
Actinomyces viscosus		23.22	0.06			
27555002	**Good identification**					**27555002**
Actinomyces viscosus		95.37	0.48	XYL 1		
Actinomyces israelii		3.73	0.39			
Actinomyces naeslundii		0.71	0.36			
27555003	**Good identification**					**27555003**
Actinomyces viscosus		96.77	0.40	XYL 1		
Actinomyces naeslundii		3.23	0.39			
27555020	**Low discrimination**					**27555020**
Actinomyces naeslundii		69.68	0.36	XYL 1		
Actinomyces israelii		15.63	0.16			
Clostridium perfringens		14.70	0.02			
27555021	**Good identification**					**27555021**
Actinomyces naeslundii		100.0	0.39	XYL 1		
27555023	**Low discrimination**					**27555023**
Actinomyces naeslundii		76.78	0.39	XYL 1		
Actinomyces viscosus		23.22	0.06			
27555100	**Low discrimination**					**27555100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		65.71	0.46	SAL 92			
Actinomyces naeslundii		13.31	0.44				
Actinomyces viscosus		11.31	0.18				
27555101	**Low discrimination**						**27555101**
Actinomyces naeslundii		84.11	0.47	XYL 1			
Actinomyces viscosus		15.89	0.10				
27555102	**Low discrimination**						**27555102**
					GRA	CAT	
Actinomyces viscosus		85.42	0.52	XYL 1	99	76	
Actinomyces israelii		7.52	0.49		99	1	
Capnocytophaga ochracea		4.31	0.29		1	90	
27555103	**Acceptable identification**						**27555103**
Actinomyces viscosus		94.68	0.44	XYL 1			
Actinomyces naeslundii		5.06	0.47				
Actinomyces israelii		0.25	0.15				
27555112	**Low discrimination**						**27555112**
Actinomyces israelii		31.71	0.35	SAL 92			
Bifidobacterium adolescentis		30.42	0.29	MLZ 13			
Actinomyces viscosus		24.36	0.18				
27555121	**Good identification**						**27555121**
Actinomyces naeslundii		100.0	0.47	XYL 1			
27555122	**Low discrimination**						**27555122**
					CAT		
Capnocytophaga ochracea		48.57	0.45	ESL 99	90		
Bacteroides fragilis		35.70	0.34		99		
Prevotella oralis		9.91	0.33		1		
27555123	**Low discrimination**						**27555123**
					GRA		
Actinomyces naeslundii		59.25	0.47	XYL 1	99		
Capnocytophaga ochracea		15.24	0.11		1		
Bacteroides fragilis		11.20	-.01		1		
27555140	**Acceptable identification**						**27555140**
Actinomyces israelii		91.58	0.39	SAL 92			
Actinomyces naeslundii		8.42	0.31				
27555142	**Low discrimination**						**27555142**
					GRA	CAT	
Actinomyces israelii		64.21	0.42	SAL 92	99	1	
Actinomyces viscosus		18.95	0.18		99	76	
Capnocytophaga ochracea		10.51	0.13		1	90	
27555322	**Low discrimination**						**27555322**
					CAT		
Parabacteroides distasonis		57.26	0.43	NIT 1	76		
Prevotella oralis		35.60	0.27		1		
Capnocytophaga ochracea		4.11	0.11		90		
27555332	**Good identification**						**27555332**
Parabacteroides distasonis		97.08	0.39	NIT 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		2.93	0.07				
27555402	**Low discrimination**						**27555402**
Actinomyces israelii		70.60	0.37	SAL 92			
Actinomyces viscosus		22.29	0.14				
Lactobacillus fermentum		4.38	0.16				
27555500	**Low discrimination**						**27555500**
Actinomyces israelii		85.32	0.45	SAL 92			
Bifidobacterium longum subsp. longum		6.27	0.17				
Lactobacillus fermentum		3.24	0.20				
27555502	**Low discrimination**						**27555502**
Actinomyces israelii		43.52	0.48	SAL 92			
Bifidobacterium adolescentis		41.76	0.41				
Actinomyces viscosus		6.11	0.18				
27555512	**Low discrimination**						**27555512**
Bifidobacterium adolescentis		80.87	0.43	NIT 1			
Actinomyces israelii		10.31	0.34				
Bifidobacterium longum subsp. longum		5.76	0.21				
27555540	**Good identification**						**27555540**
Actinomyces israelii		100.0	0.38	SAL 92			
27555542	**Low discrimination**						**27555542**
					GRA	CAT	
Actinomyces israelii		74.61	0.41	SAL 92	99	1	
Bifidobacterium adolescentis		22.75	0.26		99	1	
Capnocytophaga ochracea		2.64	0.00		1	90	
27555720	**Good identification**						**27555720**
Bacteroides vulgatus		100.0	0.38	NIT 1			
27555722	**Low discrimination**						**27555722**
Parabacteroides distasonis		81.58	0.39	NIT 1			
Bacteroides vulgatus		7.68	0.04				
Clostridium clostridioforme		7.43	0.16				
27557000	**Acceptable identification**						**27557000**
Actinomyces israelii		91.12	0.49	SAL 92			
Actinomyces naeslundii		8.89	0.41				
27557001	**Acceptable identification**						**27557001**
Actinomyces naeslundii		93.54	0.44	XYL 1			
Actinomyces israelii		6.46	0.15				
27557002	**Acceptable identification**						**27557002**
Actinomyces israelii		90.04	0.52	SAL 92			
Actinomyces naeslundii		5.85	0.41				
Actinomyces viscosus		4.10	0.14				
27557003	**Low discrimination**						**27557003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		79.40	0.44	XYL 1		
Actinomyces viscosus		12.37	0.06			
Actinomyces israelii		8.23	0.18			
27557012	**Good identification**					**27557012**
Actinomyces israelii		100.0	0.38	SAL 92		
27557020	**Low discrimination**					**27557020**
Actinomyces naeslundii		60.44	0.41	XYL 1		
Actinomyces israelii		39.56	0.28			
27557021	**Good identification**					**27557021**
Actinomyces naeslundii		100.0	0.44	XYL 1		
27557022	**Low discrimination**					**27557022**
Actinomyces naeslundii		50.46	0.41	XYL 1		
Actinomyces israelii		49.54	0.31			
27557023	**Good identification**					**27557023**
Actinomyces naeslundii		100.0	0.44	XYL 1		
27557040	**Good identification**					**27557040**
Actinomyces israelii		95.76	0.42	SAL 92		
Actinomyces naeslundii		4.24	0.28			
27557042	**Good identification**					**27557042**
Actinomyces israelii		97.14	0.45	SAL 92		
Actinomyces naeslundii		2.87	0.28			
27557100	**Good identification**					**27557100**
Actinomyces israelii		92.78	0.59	SAL 92		
Actinomyces naeslundii		6.44	0.49			
Propionibacterium propionicum		0.79	0.23			
27557101	**Acceptable identification**					**27557101**
Actinomyces naeslundii		91.15	0.52	XYL 1		
Actinomyces israelii		8.85	0.25			
27557102	**Good identification**					**27557102**
Actinomyces israelii		91.58	0.62	SAL 92		
Actinomyces naeslundii		4.24	0.49			
Actinomyces viscosus		1.86	0.18			
27557103	**Low discrimination**					**27557103**
Actinomyces naeslundii		80.46	0.52	XYL 1		
Actinomyces israelii		11.71	0.28			
Actinomyces viscosus		7.83	0.10			
27557110	**Good identification**					**27557110**
Actinomyces israelii		97.62	0.45	SAL 92		
Actinomyces naeslundii		2.39	0.27	MLZ 13		
27557112	**Low discrimination**					**27557112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		86.59	0.48	SAL 92			
Bifidobacterium adolescentis		12.00	0.27	MLZ 13			
Actinomyces naeslundii		1.41	0.27				
27557120	**Low discrimination**						**27557120**
Actinomyces naeslundii		52.08	0.49	XYL 1			
Actinomyces israelii		47.92	0.39				
27557121	**Good identification**						**27557121**
Actinomyces naeslundii		100.0	0.52	XYL 1			
27557122	**Low discrimination**						**27557122**
					GRA	CAT	
Actinomyces israelii		49.90	0.42	SAL 92	99	1	
Actinomyces naeslundii		36.16	0.49	NAG 6	99	25	
Capnocytophaga ochracea		7.19	0.11		1	90	
27557123	**Good identification**						**27557123**
Actinomyces naeslundii		100.0	0.52	XYL 1			
27557140	**Acceptable identification**						**27557140**
Actinomyces israelii		94.94	0.52	SAL 92			
Actinomyces naeslundii		2.99	0.36				
Propionibacterium propionicum		2.07	0.23				
27557141	**Low discrimination**						**27557141**
Actinomyces naeslundii		82.38	0.39	XYL 1			
Actinomyces israelii		17.62	0.18				
27557142	**Good identification**						**27557142**
Actinomyces israelii		95.40	0.55	SAL 92			
Propionibacterium propionicum		2.08	0.26				
Actinomyces naeslundii		2.00	0.36				
27557143	**Low discrimination**						**27557143**
Actinomyces naeslundii		75.71	0.39	XYL 1			
Actinomyces israelii		24.29	0.21				
27557150	**Good identification**						**27557150**
Actinomyces israelii		100.0	0.38	SAL 92			
27557152	**Good identification**						**27557152**
Actinomyces israelii		95.78	0.41	SAL 92			
Bifidobacterium adolescentis		4.22	0.12	MLZ 13			
27557161	**Good identification**						**27557161**
Actinomyces naeslundii		100.0	0.39	XYL 1			
27557163	**Good identification**						**27557163**
Actinomyces naeslundii		100.0	0.39	XYL 1			
27557300	**Good identification**						**27557300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	SAL 92			
27557302	**Good identification**						**27557302**
Actinomyces israelii		100.0	0.43	SAL 92			
27557342	**Good identification**						**27557342**
Actinomyces israelii		100.0	0.36	SAL 92			
27557400	**Good identification**						**27557400**
Actinomyces israelii		100.0	0.47	SAL 92			
27557402	**Good identification**						**27557402**
Actinomyces israelii		100.0	0.50	SAL 92			
27557412	**Good identification**						**27557412**
Actinomyces israelii		95.62	0.36	SAL 92			
Bifidobacterium adolescentis		4.38	0.07	MLZ 13			
27557440	**Good identification**						**27557440**
Actinomyces israelii		100.0	0.40	SAL 92			
27557442	**Good identification**						**27557442**
Actinomyces israelii		100.0	0.43	SAL 92			
27557500	**Very good identification**						**27557500**
Actinomyces israelii		100.0	0.58	SAL 92			
27557502	**Low discrimination**						**27557502**
Actinomyces israelii		87.83	0.61	SAL 92			
Bifidobacterium adolescentis		12.17	0.40				
27557510	**Good identification**						**27557510**
Actinomyces israelii		98.31	0.44	SAL 92			
Bifidobacterium adolescentis		1.69	0.07	MLZ 13			
27557512	**Low discrimination**						**27557512**
Bifidobacterium adolescentis		53.12	0.41	NIT 1			
Actinomyces israelii		46.88	0.47				
27557520	**Good identification**						**27557520**
Actinomyces israelii		100.0	0.37	SAL 92			
27557522	**Good identification**						**27557522**
					GRA	CAT	
Actinomyces israelii		96.99	0.40	SAL 92	99	1	
Capnocytophaga ochracea		3.01	-.02	NAG 6	1	90	
27557540	**Good identification**						**27557540**
Actinomyces israelii		100.0	0.51	SAL 92			
27557542	**Good identification**						**27557542**
Actinomyces israelii		95.78	0.54	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		4.22	0.24			
27557550	**Good identification**					**27557550**
Actinomyces israelii		100.0	0.37	SAL 92		
27557552	**Low discrimination**					**27557552**
Actinomyces israelii		73.53	0.40	SAL 92		
Bifidobacterium adolescentis		26.48	0.26	MLZ 13		
27557700	**Good identification**					**27557700**
Actinomyces israelii		100.0	0.38	SAL 92		
27557702	**Good identification**					**27557702**
Actinomyces israelii		100.0	0.41	SAL 92		
27560002	**Good identification**					**27560002**
Actinomyces viscosus		99.51	0.45	FRU 95		
Actinomyces naeslundii		0.49	0.29	TRE 7		
27560003	**Good identification**					**27560003**
Actinomyces viscosus		97.85	0.37	FRU 95		
Actinomyces naeslundii		2.15	0.32	TRE 7		
27560101	**Low discrimination**					**27560101**
Actinomyces naeslundii		77.71	0.40	FRU 95		
Actinomyces viscosus		22.29	0.07	GAL 96		
27560102	**Good identification**					**27560102**
Actinomyces viscosus		99.22	0.49	FRU 95		
Actinomyces naeslundii		0.78	0.37	TRE 7		
27560103	**Good identification**					**27560103**
Actinomyces viscosus		96.60	0.41	FRU 95		
Actinomyces naeslundii		3.40	0.40	TRE 7		
27560120	**Good identification**					**27560120**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27560121	**Good identification**					**27560121**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27560123	**Low discrimination**					**27560123**
Actinomyces naeslundii		77.71	0.40	FRU 95		
Actinomyces viscosus		22.29	0.07	GAL 96		
27561000	**Acceptable identification**					**27561000**
Actinomyces naeslundii		94.99	0.53	FRU 95		
Actinomyces viscosus		5.01	0.07			
27561001	**Good identification**					**27561001**
Actinomyces naeslundii		98.84	0.56	FRU 95		
Actinomyces viscosus		1.16	-.01			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27561002	**Low discrimination**					**27561002**
Actinomyces viscosus		83.93	0.41	FRU 95		
Actinomyces naeslundii		16.07	0.53	TRE 7		
27561020	**Good identification**					**27561020**
Actinomyces naeslundii		96.27	0.53	FRU 95		
Clostridium perfringens		3.73	0.07			
27561021	**Good identification**					**27561021**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27561022	**Acceptable identification**					**27561022**
Actinomyces naeslundii		93.37	0.53	FRU 95		
Actinomyces viscosus		4.93	0.07			
Clostridium perfringens		1.70	0.01			
27561023	**Good identification**					**27561023**
Actinomyces naeslundii		98.84	0.56	FRU 95		
Actinomyces viscosus		1.16	-.01			
27561040	**Good identification**					**27561040**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27561041	**Good identification**					**27561041**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27561042	**Low discrimination**					**27561042**
Actinomyces naeslundii		76.98	0.40	FRU 95		
Actinomyces viscosus		23.02	0.07			
27561043	**Acceptable identification**					**27561043**
Actinomyces naeslundii		93.77	0.43	FRU 95		
Actinomyces viscosus		6.23	-.01			
27561060	**Good identification**					**27561060**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27561061	**Good identification**					**27561061**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27561062	**Good identification**					**27561062**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27561063	**Good identification**					**27561063**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27561100	**Very good identification**					**27561100**
Actinomyces naeslundii		96.81	0.61	FRU 95		
Actinomyces viscosus		3.19	0.11			
27561101	**Very good identification**					**27561101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		98.36	0.64	FRU 95	1	
Clostridium chauvoei		0.93	0.06		99	
Actinomyces viscosus		0.72	0.03		1	
27561102	**Low discrimination**					**27561102**
Actinomyces viscosus		76.54	0.45	FRU 95		
Actinomyces naeslundii		23.46	0.61	TRE 7		
27561103	**Low discrimination**					**27561103**
Actinomyces naeslundii		57.97	0.64	FRU 95		
Actinomyces viscosus		42.03	0.37			
27561110	**Good identification**					**27561110**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27561111	**Good identification**					**27561111**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27561112	**Low discrimination**					**27561112**
Actinomyces naeslundii		61.49	0.39	FRU 95		
Actinomyces viscosus		38.51	0.11	MLZ 5		
27561113	**Low discrimination**					**27561113**
Actinomyces naeslundii		87.79	0.42	FRU 95		
Actinomyces viscosus		12.21	0.03	MLZ 5		
27561120	**Low discrimination**					**27561120**
					GRA	
Actinomyces naeslundii		89.90	0.61	FRU 95	99	
Clostridium perfringens		8.97	0.21		99	
Prevotella melaninogenica		1.13	-.01		1	
27561121	**Very good identification**					**27561121**
Actinomyces naeslundii		100.0	0.64	FRU 95		
27561122	**Good identification**					**27561122**
Actinomyces naeslundii		92.60	0.61	FRU 95		
Clostridium perfringens		4.35	0.16			
Actinomyces viscosus		3.05	0.11			
27561123	**Very good identification**					**27561123**
Actinomyces naeslundii		99.27	0.64	FRU 95		
Actinomyces viscosus		0.73	0.03			
27561130	**Good identification**					**27561130**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27561131	**Good identification**					**27561131**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27561132	**Good identification**					**27561132**
Actinomyces naeslundii		100.0	0.39	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27561133	**Good identification**					**27561133**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27561140	**Good identification**					**27561140**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27561141	**Good identification**					**27561141**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27561142	**Low discrimination**					**27561142**
Actinomyces naeslundii		84.26	0.48	FRU 95		
Actinomyces viscosus		15.74	0.11			
27561143	**Good identification**					**27561143**
Actinomyces naeslundii		96.02	0.51	FRU 95		
Actinomyces viscosus		3.98	0.03			
27561160	**Acceptable identification**					**27561160**
Actinomyces naeslundii		90.28	0.48	FRU 95		
Clostridium perfringens		9.72	0.09			
27561161	**Good identification**					**27561161**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27561162	**Good identification**					**27561162**
Actinomyces naeslundii		95.18	0.48	FRU 95		
Clostridium perfringens		4.82	0.03			
27561163	**Good identification**					**27561163**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27562001	**Good identification**					**27562001**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27562003	**Low discrimination**					**27562003**
Actinomyces naeslundii		80.87	0.37	FRU 95		
Actinomyces viscosus		19.13	0.02	GAL 96		
27562021	**Good identification**					**27562021**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27562023	**Good identification**					**27562023**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27562100	**Low discrimination**					**27562100**
Actinomyces naeslundii		80.80	0.42	FRU 95		
Propionibacterium propionicum		19.20	0.21	GAL 96		
27562101	**Good identification**					**27562101**
Actinomyces naeslundii		100.0	0.45	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27562102	**Low discrimination**					**27562102**
Actinomyces naeslundii		49.47	0.42	FRU 95		
Actinomyces viscosus		32.90	0.15	GAL 96		
Propionibacterium propionicum		17.63	0.24			
27562103	**Low discrimination**					**27562103**
Actinomyces naeslundii		87.13	0.45	FRU 95		
Actinomyces viscosus		12.87	0.07	GAL 96		
27562120	**Good identification**					**27562120**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27562121	**Good identification**					**27562121**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27562122	**Good identification**					**27562122**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27562123	**Good identification**					**27562123**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27563000	**Very good identification**					**27563000**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27563001	**Very good identification**					**27563001**
Actinomyces naeslundii		100.0	0.61	FRU 95		
27563002	**Very good identification**					**27563002**
Actinomyces naeslundii		97.35	0.58	FRU 95		
Actinomyces viscosus		2.65	0.07			
27563003	**Very good identification**					**27563003**
Actinomyces naeslundii		99.40	0.61	FRU 95		
Actinomyces viscosus		0.60	-.01			
27563010	**Good identification**					**27563010**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27563011	**Good identification**					**27563011**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27563012	**Good identification**					**27563012**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27563013	**Good identification**					**27563013**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27563020	**Very good identification**					**27563020**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27563021	**Very good identification**					**27563021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.61	FRU 95		
27563022	**Very good identification**					**27563022**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27563023	**Very good identification**					**27563023**
Actinomyces naeslundii		100.0	0.61	FRU 95		
27563030	**Good identification**					**27563030**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27563031	**Good identification**					**27563031**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27563032	**Good identification**					**27563032**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27563033	**Good identification**					**27563033**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27563040	**Good identification**					**27563040**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27563041	**Good identification**					**27563041**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27563042	**Good identification**					**27563042**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27563043	**Good identification**					**27563043**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27563060	**Good identification**					**27563060**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27563061	**Good identification**					**27563061**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27563062	**Good identification**					**27563062**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27563063	**Good identification**					**27563063**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27563100	**Very good identification**					**27563100**
Actinomyces naeslundii		98.56	0.66	FRU 95		
Propionibacterium propionicum		0.98	0.21			
Actinomyces israelii		0.46	0.17			
27563101	**Very good identification**					**27563101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.69	FRU 95		
27563102	**Very good identification**					**27563102**
Actinomyces naeslundii		96.26	0.66	FRU 95		
Actinomyces viscosus		1.63	0.11			
Propionibacterium propionicum		1.43	0.24			
27563103	**Very good identification**					**27563103**
Actinomyces naeslundii		99.62	0.69	FRU 95		
Actinomyces viscosus		0.38	0.03			
27563110	**Good identification**					**27563110**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27563111	**Good identification**					**27563111**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27563112	**Good identification**					**27563112**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27563113	**Good identification**					**27563113**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27563120	**Very good identification**					**27563120**
Actinomyces naeslundii		100.0	0.66	FRU 95		
27563121	**Very good identification**					**27563121**
Actinomyces naeslundii		100.0	0.69	FRU 95		
27563122	**Very good identification**					**27563122**
Actinomyces naeslundii		100.0	0.66	FRU 95		
27563123	**Very good identification**					**27563123**
Actinomyces naeslundii		100.0	0.69	FRU 95		
27563130	**Good identification**					**27563130**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27563131	**Good identification**					**27563131**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27563132	**Good identification**					**27563132**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27563133	**Good identification**					**27563133**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27563140	**Acceptable identification**					**27563140**
Actinomyces naeslundii		94.69	0.53	FRU 95		
Propionibacterium propionicum		5.31	0.21			
27563141	**Good identification**					**27563141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.56	FRU 95		
27563142	**Acceptable identification**					**27563142**
Actinomyces naeslundii		92.24	0.53	FRU 95		
Propionibacterium propionicum		7.76	0.24			
27563143	**Good identification**					**27563143**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27563160	**Good identification**					**27563160**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27563161	**Good identification**					**27563161**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27563162	**Good identification**					**27563162**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27563163	**Good identification**					**27563163**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27567102	**Low discrimination**					**27567102**
Actinomyces israelii		88.94	0.38	SAL 92		
Actinomyces naeslundii		11.06	0.32	FRU 99		
27567502	**Good identification**					**27567502**
Actinomyces israelii		100.0	0.36	SAL 92		
27570001	**Low discrimination**					**27570001**
Actinomyces naeslundii		68.53	0.54	GAL 96		
Actinomyces viscosus		31.47	0.24			
27570002	**Very good identification**					**27570002**
Actinomyces viscosus		99.49	0.67	TRE 7		
Actinomyces naeslundii		0.49	0.51			
Propionibacterium granulosum		0.03	0.15			
27570003	**Very good identification**					**27570003**
Actinomyces viscosus		97.85	0.59	TRE 7		
Actinomyces naeslundii		2.15	0.54			
27570020	**Low discrimination**					**27570020**
					CAT	
Actinomyces naeslundii		88.54	0.51	GAL 96	25	
Propionibacterium granulosum		5.28	0.15		90	
Clostridium perfringens		4.33	0.07		1	
27570021	**Good identification**					**27570021**
Actinomyces naeslundii		100.0	0.54	GAL 96		
27570023	**Low discrimination**					**27570023**
Actinomyces naeslundii		68.53	0.54	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		31.47	0.24			
27570040	**Low discrimination**					**27570040**
					SPO	
Actinomyces naeslundii		77.81	0.38	GAL 96	1	
Clostridium botulinum biovar B		12.99	-.03		99	
Actinomyces viscosus		9.20	-.02		1	
27570041	**Good identification**					**27570041**
Actinomyces naeslundii		100.0	0.41	GAL 96		
27570060	**Good identification**					**27570060**
Actinomyces naeslundii		100.0	0.38	GAL 96		
27570061	**Good identification**					**27570061**
Actinomyces naeslundii		100.0	0.41	GAL 96		
27570062	**Low discrimination**					**27570062**
Actinomyces naeslundii		89.43	0.38	GAL 96		
Actinomyces viscosus		10.58	-.02			
27570063	**Good identification**					**27570063**
Actinomyces naeslundii		100.0	0.41	GAL 96		
27570100	**Low discrimination**					**27570100**
					GRA	
Actinomyces viscosus		54.96	0.37	TRE 7	99	
Actinomyces naeslundii		42.58	0.59	BGL 99	99	
Fusobacterium mortiferum		1.43	0.22		1	
27570101	**Low discrimination**					**27570101**
Actinomyces naeslundii		77.71	0.62	GAL 96		
Actinomyces viscosus		22.29	0.29			
27570102	**Very good identification**					**27570102**
Actinomyces viscosus		99.20	0.71	TRE 7		
Actinomyces naeslundii		0.78	0.59			
Propionibacterium propionicum		0.03	0.24			
27570103	**Very good identification**					**27570103**
Actinomyces viscosus		96.60	0.63	TRE 7		
Actinomyces naeslundii		3.40	0.62			
27570110	**Low discrimination**					**27570110**
Actinomyces naeslundii		80.15	0.37	GAL 96		
Actinomyces viscosus		19.85	0.03	MLZ 5		
27570111	**Good identification**					**27570111**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27570112	**Good identification**					**27570112**
Actinomyces viscosus		96.08	0.37	TRE 7		
Actinomyces naeslundii		3.92	0.37	MLZ 1		
27570120	**Low discrimination**					**27570120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		87.79	0.59	GAL 96		
Clostridium perfringens		11.06	0.21			
Actinomyces viscosus		1.14	0.03			
27570121	**Very good identification**					**27570121**
Actinomyces naeslundii		100.0	0.62	GAL 96		
27570122	**Low discrimination**					**27570122**
Actinomyces viscosus		54.92	0.37	TRE 7		
Actinomyces naeslundii		42.55	0.59	NAG 1		
Clostridium perfringens		2.52	0.16			
27570123	**Low discrimination**					**27570123**
Actinomyces naeslundii		77.71	0.62	GAL 96		
Actinomyces viscosus		22.29	0.29			
27570130	**Good identification**					**27570130**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27570131	**Good identification**					**27570131**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27570132	**Low discrimination**					**27570132**
Actinomyces naeslundii		80.15	0.37	GAL 96		
Actinomyces viscosus		19.85	0.03	MLZ 5		
27570133	**Good identification**					**27570133**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27570140	**Low discrimination**					**27570140**
Actinomyces naeslundii		82.55	0.46	GAL 96		
Propionibacterium propionicum		11.36	0.21			
Actinomyces viscosus		6.10	0.03			
27570141	**Good identification**					**27570141**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27570142	**Low discrimination**					**27570142**
Actinomyces viscosus		85.84	0.37	TRE 7		
Actinomyces naeslundii		11.74	0.46	SOR 1		
Propionibacterium propionicum		2.42	0.24			
27570160	**Low discrimination**					**27570160**
Actinomyces naeslundii		88.03	0.46	GAL 96		
Clostridium perfringens		11.97	0.09			
27570161	**Good identification**					**27570161**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27570162	**Low discrimination**					**27570162**
Actinomyces naeslundii		87.88	0.46	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		6.49	0.03			
Clostridium perfringens		5.62	0.03			
27570163	**Good identification**					**27570163**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27570302	**Good identification**					**27570302**
Actinomyces viscosus		99.22	0.37	TRE 7		
Actinomyces naeslundii		0.78	0.25	RHA 1		
27570502	**Good identification**					**27570502**
Actinomyces viscosus		99.22	0.37	TRE 7		
Actinomyces naeslundii		0.78	0.25	ARA 1		
27571000	**Good identification**					**27571000**
Actinomyces naeslundii		92.56	0.75			
Actinomyces viscosus		4.88	0.29			
Propionibacterium granulosum		0.92	0.26			
27571001	**Very good identification**					**27571001**
Actinomyces naeslundii		98.84	0.78			
Actinomyces viscosus		1.16	0.21			
27571002	**Low discrimination**					**27571002**
Actinomyces viscosus		83.70	0.63	TRE 7		
Actinomyces naeslundii		16.02	0.75			
Actinomyces israelii		0.14	0.31			
27571003	**Low discrimination**					**27571003**
Actinomyces viscosus		53.72	0.55	TRE 7		
Actinomyces naeslundii		46.28	0.78			
27571010	**Acceptable identification**					**27571010**
					CAT	
Actinomyces naeslundii		94.26	0.53	MLZ 5	25	
Propionibacterium granulosum		2.43	0.11		90	
Lactobacillus jensenii		1.71	0.02		1	
27571011	**Good identification**					**27571011**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27571013	**Low discrimination**					**27571013**
Actinomyces naeslundii		81.78	0.56	MLZ 5		
Actinomyces viscosus		18.22	0.21			
27571020	**Low discrimination**					**27571020**
					CAT	
Actinomyces naeslundii		83.04	0.75		25	
Clostridium perfringens		16.75	0.41		1	
Propionibacterium granulosum		0.21	0.15		90	
27571021	**Very good identification**					**27571021**
Actinomyces naeslundii		99.86	0.78			
Clostridium perfringens		0.14	0.07			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27571022	**Low discrimination**					**27571022**
Actinomyces naeslundii		86.74	0.75			
Clostridium perfringens		8.24	0.35			
Actinomyces viscosus		4.58	0.29			
27571023	**Very good identification**					**27571023**
Actinomyces naeslundii		98.78	0.78			
Actinomyces viscosus		1.16	0.21			
Clostridium perfringens		0.06	0.01			
27571030	**Good identification**					**27571030**
Actinomyces naeslundii		96.27	0.53	MLZ 5		
Clostridium perfringens		3.73	0.07			
27571031	**Good identification**					**27571031**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27571032	**Good identification**					**27571032**
Actinomyces naeslundii		98.21	0.53	MLZ 5		
Clostridium perfringens		1.79	0.01			
27571033	**Good identification**					**27571033**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27571040	**Very good identification**					**27571040**
Actinomyces naeslundii		98.05	0.62			
Actinomyces israelii		1.29	0.21			
Clostridium perfringens		0.66	0.03			
27571041	**Very good identification**					**27571041**
Actinomyces naeslundii		100.0	0.65			
27571042	**Low discrimination**					**27571042**
Actinomyces naeslundii		75.83	0.62			
Actinomyces viscosus		22.67	0.29			
Actinomyces israelii		1.50	0.24			
27571043	**Good identification**					**27571043**
Actinomyces naeslundii		93.77	0.65			
Actinomyces viscosus		6.23	0.21			
27571050	**Good identification**					**27571050**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27571051	**Good identification**					**27571051**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27571052	**Good identification**					**27571052**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27571053	**Good identification**					**27571053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27571060	**Low discrimination**					**27571060**
Actinomyces naeslundii		82.12	0.62			
Clostridium perfringens		17.88	0.28			
27571061	**Very good identification**					**27571061**
Actinomyces naeslundii		100.0	0.65			
27571062	**Good identification**					**27571062**
Actinomyces naeslundii		90.71	0.62			
Clostridium perfringens		9.30	0.23			
27571063	**Very good identification**					**27571063**
Actinomyces naeslundii		100.0	0.65			
27571070	**Good identification**					**27571070**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27571071	**Good identification**					**27571071**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27571072	**Good identification**					**27571072**
Actinomyces naeslundii		100.0	0.40	MLZ 5		
27571073	**Good identification**					**27571073**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27571100	**Good identification**					**27571100**
Actinomyces naeslundii		93.97	0.83			
Actinomyces viscosus		3.10	0.33			
Clostridium perfringens		1.51	0.30			
27571101	**Very good identification**					**27571101**
					SPO	
Actinomyces naeslundii		99.16	0.86		1	
Actinomyces viscosus		0.73	0.25		1	
Clostridium chauvoei		0.12	0.12		99	
27571102	**Low discrimination**					**27571102**
Actinomyces viscosus		76.01	0.67	TRE 7		
Actinomyces naeslundii		23.29	0.83			
Actinomyces israelii		0.29	0.41			
27571103	**Low discrimination**					**27571103**
					SPO	
Actinomyces naeslundii		57.96	0.86		1	
Actinomyces viscosus		42.03	0.59		1	
Clostridium chauvoei		0.01	-.01		99	
27571110	**Very good identification**					**27571110**
Actinomyces naeslundii		97.07	0.61	MLZ 5		
Actinomyces israelii		2.32	0.24			
Actinomyces viscosus		0.61	-.01			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27571111	**Very good identification**					**27571111**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
27571112	**Low discrimination**					**27571112**
					GRA	
Actinomyces naeslundii		57.26	0.61	MLZ 5	99	
Actinomyces viscosus		35.86	0.33		99	
Leptotrichia buccalis		4.23	0.07		1	
27571113	**Low discrimination**					**27571113**
Actinomyces naeslundii		87.78	0.64	MLZ 5		
Actinomyces viscosus		12.22	0.25			
27571120	**Low discrimination**					**27571120**
					GRA	
Actinomyces naeslundii		65.37	0.83		99	
Clostridium perfringens		33.98	0.56		99	
Prevotella melaninogenica		0.39	0.15		1	
27571121	**Very good identification**					**27571121**
					SPO	
Actinomyces naeslundii		99.61	0.86		1	
Clostridium perfringens		0.35	0.21		84	
Clostridium chauvoei		0.04	0.04		99	
27571122	**Low discrimination**					**27571122**
Actinomyces naeslundii		75.71	0.83			
Clostridium perfringens		18.52	0.50			
Actinomyces viscosus		2.50	0.33			
27571123	**Very good identification**					**27571123**
Actinomyces naeslundii		99.11	0.86			
Actinomyces viscosus		0.73	0.25			
Clostridium perfringens		0.16	0.16			
27571130	**Good identification**					**27571130**
Actinomyces naeslundii		90.93	0.61	MLZ 5		
Clostridium perfringens		9.07	0.21			
27571131	**Very good identification**					**27571131**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
27571132	**Low discrimination**					**27571132**
					GRA	
Actinomyces naeslundii		88.76	0.61	MLZ 5	99	
Parabacteroides distasonis		4.28	0.25		1	
Clostridium perfringens		4.17	0.16		99	
27571133	**Very good identification**					**27571133**
Actinomyces naeslundii		100.0	0.64	MLZ 5		
27571140	**Very good identification**					**27571140**
Actinomyces naeslundii		95.83	0.70			
Actinomyces israelii		1.78	0.31			
Clostridium perfringens		1.66	0.17			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27571141	**Very good identification**						**27571141**
Actinomyces naeslundii		100.0	0.73				
27571142	**Low discrimination**						**27571142**
Actinomyces naeslundii		80.97	0.70				
Actinomyces viscosus		15.12	0.33				
Actinomyces israelii		2.25	0.34				
27571143	**Very good identification**						**27571143**
Actinomyces naeslundii		96.02	0.73				
Actinomyces viscosus		3.99	0.25				
27571150	**Good identification**						**27571150**
Actinomyces naeslundii		95.01	0.48	MLZ 5			
Actinomyces israelii		5.00	0.17				
27571151	**Good identification**						**27571151**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
27571152	**Low discrimination**						**27571152**
Actinomyces naeslundii		89.71	0.48	MLZ 5			
Actinomyces israelii		7.08	0.20				
Actinomyces viscosus		3.22	-.01				
27571153	**Good identification**						**27571153**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
27571160	**Low discrimination**						**27571160**
Actinomyces naeslundii		64.06	0.70				
Clostridium perfringens		35.94	0.43				
27571161	**Very good identification**						**27571161**
Actinomyces naeslundii		99.62	0.73				
Clostridium perfringens		0.38	0.09				
27571162	**Low discrimination**						**27571162**
					GRA	CAT	
Actinomyces naeslundii		78.34	0.70		99	25	
Clostridium perfringens		20.68	0.38		99	1	
Capnocytophaga ochracea		0.40	0.05		1	90	
27571163	**Very good identification**						**27571163**
Actinomyces naeslundii		99.82	0.73				
Clostridium perfringens		0.18	0.03				
27571170	**Acceptable identification**						**27571170**
Actinomyces naeslundii		90.28	0.48	MLZ 5			
Clostridium perfringens		9.72	0.09				
27571171	**Good identification**						**27571171**
Actinomyces naeslundii		100.0	0.51	MLZ 5			
27571172	**Good identification**						**27571172**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.18	0.48	MLZ 5		
Clostridium perfringens		4.82	0.03			
27571173	**Good identification**					**27571173**
Actinomyces naeslundii		100.0	0.51	MLZ 5		
27571200	**Good identification**					**27571200**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27571201	**Good identification**					**27571201**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27571220	**Low discrimination**					**27571220**
Actinomyces naeslundii		83.22	0.41	RHA 1		
Clostridium perfringens		16.79	0.07			
27571221	**Good identification**					**27571221**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27571222	**Low discrimination**					**27571222**
					GRA	
Actinomyces naeslundii		79.60	0.41	RHA 1	99	
Prevotella oralis		12.85	0.02		1	
Clostridium perfringens		7.56	0.01		99	
27571223	**Good identification**					**27571223**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27571300	**Acceptable identification**					**27571300**
Actinomyces naeslundii		91.28	0.49	RHA 1		
Actinomyces israelii		5.72	0.19			
Actinomyces viscosus		3.01	-.01			
27571301	**Good identification**					**27571301**
Actinomyces naeslundii		100.0	0.52	RHA 1		
27571303	**Low discrimination**					**27571303**
Actinomyces naeslundii		57.97	0.52	RHA 1		
Actinomyces viscosus		42.03	0.25			
27571320	**Low discrimination**					**27571320**
					SPO	
Actinomyces naeslundii		60.94	0.49	RHA 1	1	
Clostridium perfringens		31.68	0.21		84	
Clostridium ramosum		4.53	0.04		92	
27571321	**Good identification**					**27571321**
Actinomyces naeslundii		100.0	0.52	RHA 1		
27571322	**Low discrimination**					**27571322**
					GRA	
Parabacteroides distasonis		35.88	0.38	NIT 1	1	
Actinomyces naeslundii		25.62	0.49	XYL 90	99	
Prevotella oralis		22.76	0.22		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27571323	**Good identification**					**27571323**
Actinomyces naeslundii		100.0	0.52	RHA 1		
27571340	**Good identification**					**27571340**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27571341	**Good identification**					**27571341**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27571342	**Low discrimination**					**27571342**
Actinomyces naeslundii		71.75	0.36	RHA 1		
Actinomyces israelii		14.85	0.15			
Actinomyces viscosus		13.40	-.01			
27571343	**Good identification**					**27571343**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27571360	**Low discrimination**					**27571360**
Actinomyces naeslundii		64.06	0.36	RHA 1		
Clostridium perfringens		35.94	0.09			
27571361	**Good identification**					**27571361**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27571362	**Low discrimination**					**27571362**
Actinomyces naeslundii		79.12	0.36	RHA 1		
Clostridium perfringens		20.89	0.03			
27571363	**Good identification**					**27571363**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27571400	**Low discrimination**					**27571400**
Actinomyces naeslundii		67.37	0.41	ARA 1		
Actinomyces israelii		32.63	0.26			
27571401	**Good identification**					**27571401**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27571420	**Low discrimination**					**27571420**
Actinomyces naeslundii		83.22	0.41	ARA 1		
Clostridium perfringens		16.79	0.07			
27571421	**Good identification**					**27571421**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27571422	**Acceptable identification**					**27571422**
Actinomyces naeslundii		91.33	0.41	ARA 1		
Clostridium perfringens		8.67	0.01			
27571423	**Good identification**					**27571423**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	ARA 1		
27571500	**Low discrimination**					**27571500**
Actinomyces naeslundii		58.35	0.49	ARA 1		
Actinomyces israelii		39.73	0.36			
Actinomyces viscosus		1.92	-.01			
27571501	**Good identification**					**27571501**
Actinomyces naeslundii		100.0	0.52	ARA 1		
27571503	**Low discrimination**					**27571503**
Actinomyces naeslundii		57.97	0.52	ARA 1		
Actinomyces viscosus		42.03	0.25			
27571520	**Low discrimination**					**27571520**
Actinomyces naeslundii		61.95	0.49	ARA 1		
Clostridium perfringens		32.20	0.21			
Bifidobacterium longum subsp. longum		3.16	0.09			
27571521	**Good identification**					**27571521**
Actinomyces naeslundii		100.0	0.52	ARA 1		
27571522	**Low discrimination**					**27571522**
					GRA	
Actinomyces naeslundii		54.94	0.49	ARA 1	99	
Parabacteroides distasonis		15.72	0.26		1	
Clostridium perfringens		13.44	0.16		99	
27571523	**Good identification**					**27571523**
Actinomyces naeslundii		100.0	0.52	ARA 1		
27571541	**Good identification**					**27571541**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27571543	**Good identification**					**27571543**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27571560	**Low discrimination**					**27571560**
Actinomyces naeslundii		64.06	0.36	ARA 1		
Clostridium perfringens		35.94	0.09			
27571561	**Good identification**					**27571561**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27571562	**Low discrimination**					**27571562**
Actinomyces naeslundii		79.12	0.36	ARA 1		
Clostridium perfringens		20.89	0.03			
27571563	**Good identification**					**27571563**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27572000	**Low discrimination**					**27572000**
					CAT	
Actinomyces naeslundii		87.79	0.56	GAL 96	25	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		8.14	0.24		90	
Propionibacterium propionicum		3.13	0.21		1	
27572001	**Very good identification**					**27572001**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27572003	**Low discrimination**					**27572003**
Actinomyces naeslundii		80.87	0.59	GAL 96		
Actinomyces viscosus		19.13	0.24			
27572011	**Good identification**					**27572011**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27572013	**Good identification**					**27572013**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27572020	**Very good identification**					**27572020**
Actinomyces naeslundii		97.74	0.56	GAL 96		
Propionibacterium granulosum		2.26	0.13			
27572021	**Very good identification**					**27572021**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27572022	**Very good identification**					**27572022**
Actinomyces naeslundii		98.94	0.56	GAL 96		
Actinomyces viscosus		1.06	-.02			
27572023	**Very good identification**					**27572023**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27572031	**Good identification**					**27572031**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27572033	**Good identification**					**27572033**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27572040	**Low discrimination**					**27572040**
Actinomyces naeslundii		83.22	0.43	GAL 96		
Propionibacterium propionicum		16.78	0.21			
27572041	**Good identification**					**27572041**
Actinomyces naeslundii		100.0	0.46	GAL 96		
27572042	**Low discrimination**					**27572042**
Actinomyces naeslundii		73.34	0.43	GAL 96		
Propionibacterium propionicum		22.19	0.24			
Actinomyces viscosus		4.47	-.02			
27572043	**Good identification**					**27572043**
Actinomyces naeslundii		100.0	0.46	GAL 96		
27572060	**Good identification**					**27572060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	GAL 96		
27572061	**Good identification**					**27572061**
Actinomyces naeslundii		100.0	0.46	GAL 96		
27572062	**Good identification**					**27572062**
Actinomyces naeslundii		100.0	0.43	GAL 96		
27572063	**Good identification**					**27572063**
Actinomyces naeslundii		100.0	0.46	GAL 96		
27572100	**Low discrimination**					**27572100**
Propionibacterium propionicum		55.01	0.55	CEL 1		
Actinomyces naeslundii		44.43	0.64			
Actinomyces viscosus		0.30	0.03			
27572101	**Good identification**					**27572101**
Actinomyces naeslundii		91.60	0.67	GAL 96		
Propionibacterium propionicum		8.40	0.39			
27572102	**Low discrimination**					**27572102**
Propionibacterium propionicum		52.60	0.58	CEL 1		
Actinomyces naeslundii		28.32	0.64			
Actinomyces viscosus		18.83	0.37			
27572103	**Low discrimination**					**27572103**
Actinomyces naeslundii		77.80	0.67	GAL 96		
Actinomyces viscosus		11.50	0.29			
Propionibacterium propionicum		10.70	0.42			
27572110	**Low discrimination**					**27572110**
Actinomyces naeslundii		80.80	0.42	GAL 96		
Propionibacterium propionicum		19.20	0.21	MLZ 5		
27572111	**Good identification**					**27572111**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27572112	**Low discrimination**					**27572112**
Actinomyces naeslundii		67.39	0.42	GAL 96		
Propionibacterium propionicum		24.02	0.24	MLZ 5		
Actinomyces viscosus		8.60	0.03			
27572113	**Good identification**					**27572113**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27572120	**Very good identification**					**27572120**
Actinomyces naeslundii		98.77	0.64	GAL 96		
Propionibacterium propionicum		1.24	0.21			
27572121	**Very good identification**					**27572121**
Actinomyces naeslundii		100.0	0.67	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27572122	**Very good identification**					**27572122**
Actinomyces naeslundii		97.52	0.64	GAL 96		
Propionibacterium propionicum		1.83	0.24			
Actinomyces viscosus		0.66	0.03			
27572123	**Very good identification**					**27572123**
Actinomyces naeslundii		100.0	0.67	GAL 96		
27572130	**Good identification**					**27572130**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27572131	**Good identification**					**27572131**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27572132	**Good identification**					**27572132**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27572133	**Good identification**					**27572133**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27572140	**Low discrimination**					**27572140**
Propionibacterium propionicum		87.53	0.55	CEL 1		
Actinomyces naeslundii		12.48	0.51			
27572141	**Low discrimination**					**27572141**
Actinomyces naeslundii		65.80	0.54	GAL 96		
Propionibacterium propionicum		34.20	0.39			
27572142	**Good identification**					**27572142**
Propionibacterium propionicum		91.02	0.58	CEL 1		
Actinomyces naeslundii		8.65	0.51			
Actinomyces viscosus		0.33	0.03			
27572143	**Low discrimination**					**27572143**
Actinomyces naeslundii		56.19	0.54	GAL 96		
Propionibacterium propionicum		43.81	0.42			
27572160	**Acceptable identification**					**27572160**
Actinomyces naeslundii		93.38	0.51	GAL 96		
Propionibacterium propionicum		6.62	0.21			
27572161	**Good identification**					**27572161**
Actinomyces naeslundii		100.0	0.54	GAL 96		
27572162	**Acceptable identification**					**27572162**
Actinomyces naeslundii		90.39	0.51	GAL 96		
Propionibacterium propionicum		9.61	0.24			
27572163	**Good identification**					**27572163**
Actinomyces naeslundii		100.0	0.54	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27573000	**Very good identification**					**27573000**
					CAT	
Actinomyces naeslundii		97.77	0.80		25	
Actinomyces israelii		1.71	0.41		1	
Propionibacterium granulosum		0.38	0.24		90	
27573001	**Very good identification**					**27573001**
Actinomyces naeslundii		100.0	0.83			
27573002	**Good identification**					**27573002**
Actinomyces naeslundii		94.65	0.80			
Actinomyces viscosus		2.57	0.29			
Actinomyces israelii		2.48	0.44			
27573003	**Very good identification**					**27573003**
Actinomyces naeslundii		99.40	0.83			
Actinomyces viscosus		0.60	0.21			
27573010	**Good identification**					**27573010**
					CAT	
Actinomyces naeslundii		94.38	0.58	MLZ 5	25	
Actinomyces israelii		4.68	0.26		1	
Propionibacterium granulosum		0.94	0.09		90	
27573011	**Very good identification**					**27573011**
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27573012	**Good identification**					**27573012**
Actinomyces naeslundii		93.08	0.58	MLZ 5		
Actinomyces israelii		6.92	0.29			
27573013	**Very good identification**					**27573013**
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27573020	**Very good identification**					**27573020**
					CAT	
Actinomyces naeslundii		99.69	0.80		25	
Actinomyces israelii		0.11	0.20		1	
Propionibacterium granulosum		0.10	0.13		90	
27573021	**Very good identification**					**27573021**
Actinomyces naeslundii		100.0	0.83			
27573022	**Very good identification**					**27573022**
Actinomyces naeslundii		99.78	0.80			
Actinomyces israelii		0.17	0.23			
Clostridium perfringens		0.05	0.01			
27573023	**Very good identification**					**27573023**
Actinomyces naeslundii		100.0	0.83			
27573030	**Very good identification**					**27573030**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27573031	**Very good identification**					**27573031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27573032	**Very good identification**					**27573032**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27573033	**Very good identification**					**27573033**
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27573040	**Very good identification**					**27573040**
Actinomyces naeslundii		95.52	0.67			
Actinomyces israelii		3.68	0.34			
Propionibacterium propionicum		0.80	0.21			
27573041	**Very good identification**					**27573041**
Actinomyces naeslundii		100.0	0.70			
27573042	**Good identification**					**27573042**
Actinomyces naeslundii		93.43	0.67			
Actinomyces israelii		5.39	0.37			
Propionibacterium propionicum		1.18	0.24			
27573043	**Very good identification**					**27573043**
Actinomyces naeslundii		100.0	0.70			
27573050	**Acceptable identification**					**27573050**
Actinomyces naeslundii		90.15	0.45	MLZ 5		
Actinomyces israelii		9.85	0.19			
27573051	**Good identification**					**27573051**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27573052	**Low discrimination**					**27573052**
Actinomyces naeslundii		85.92	0.45	MLZ 5		
Actinomyces israelii		14.08	0.22			
27573053	**Good identification**					**27573053**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27573060	**Very good identification**					**27573060**
Actinomyces naeslundii		100.0	0.67			
27573061	**Very good identification**					**27573061**
Actinomyces naeslundii		100.0	0.70			
27573062	**Very good identification**					**27573062**
Actinomyces naeslundii		99.63	0.67			
Actinomyces israelii		0.37	0.16			
27573063	**Very good identification**					**27573063**
Actinomyces naeslundii		100.0	0.70			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27573070	**Good identification**					**27573070**
Actinomyces naeslundii		100.0	0.45	MLZ 5		
27573071	**Good identification**					**27573071**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27573072	**Good identification**					**27573072**
Actinomyces naeslundii		100.0	0.45	MLZ 5		
27573073	**Good identification**					**27573073**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27573100	**Good identification**					**27573100**
Actinomyces naeslundii		92.91	0.88			
Propionibacterium propionicum		4.79	0.55			
Actinomyces israelii		2.28	0.51			
27573101	**Very good identification**					**27573101**
Actinomyces naeslundii		99.60	0.91			
Propionibacterium propionicum		0.38	0.39			
Actinomyces israelii		0.02	0.17			
27573102	**Low discrimination**					**27573102**
Actinomyces naeslundii		88.39	0.88			
Propionibacterium propionicum		6.84	0.58			
Actinomyces israelii		3.25	0.54			
27573103	**Very good identification**					**27573103**
Actinomyces naeslundii		99.03	0.91			
Propionibacterium propionicum		0.57	0.42			
Actinomyces viscosus		0.37	0.25			
27573110	**Good identification**					**27573110**
Actinomyces naeslundii		92.63	0.66	MLZ 5		
Actinomyces israelii		6.45	0.37			
Propionibacterium propionicum		0.92	0.21			
27573111	**Very good identification**					**27573111**
Actinomyces naeslundii		100.0	0.69	MLZ 5		
27573112	**Low discrimination**					**27573112**
Actinomyces naeslundii		88.73	0.66	MLZ 5		
Actinomyces israelii		9.27	0.40			
Propionibacterium propionicum		1.32	0.24			
27573113	**Very good identification**					**27573113**
Actinomyces naeslundii		100.0	0.69	MLZ 5		
27573120	**Very good identification**					**27573120**
Actinomyces naeslundii		99.36	0.88			
Clostridium perfringens		0.27	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.16	0.30			
27573121	**Very good identification**					**27573121**
Actinomyces naeslundii		100.0	0.91			
27573122	**Very good identification**					**27573122**
					SPO	
Actinomyces naeslundii		98.89	0.88		1	
Clostridium ramosum		0.65	0.25		92	
Actinomyces israelii		0.23	0.33		1	
27573123	**Very good identification**					**27573123**
Actinomyces naeslundii		100.0	0.91			
27573130	**Very good identification**					**27573130**
Actinomyces naeslundii		99.56	0.66	MLZ 5		
Actinomyces israelii		0.44	0.16			
27573131	**Very good identification**					**27573131**
Actinomyces naeslundii		100.0	0.69	MLZ 5		
27573132	**Very good identification**					**27573132**
Actinomyces naeslundii		99.34	0.66	MLZ 5		
Actinomyces israelii		0.66	0.19			
27573133	**Very good identification**					**27573133**
Actinomyces naeslundii		100.0	0.69	MLZ 5		
27573140	**Low discrimination**					**27573140**
Actinomyces naeslundii		74.27	0.75			
Propionibacterium propionicum		21.71	0.55			
Actinomyces israelii		4.02	0.44			
27573141	**Very good identification**					**27573141**
Actinomyces naeslundii		97.88	0.78			
Propionibacterium propionicum		2.12	0.39			
27573142	**Low discrimination**					**27573142**
Actinomyces naeslundii		65.77	0.75			
Propionibacterium propionicum		28.84	0.58			
Actinomyces israelii		5.33	0.47			
27573143	**Very good identification**					**27573143**
Actinomyces naeslundii		96.85	0.78			
Propionibacterium propionicum		3.15	0.42			
27573150	**Low discrimination**					**27573150**
Actinomyces naeslundii		82.67	0.53	MLZ 5		
Actinomyces israelii		12.69	0.30			
Propionibacterium propionicum		4.64	0.21			
27573151	**Good identification**					**27573151**
Actinomyces naeslundii		100.0	0.56	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27573152	**Low discrimination**					**27573152**
Actinomyces naeslundii		76.08	0.53	MLZ 5		
Actinomyces israelii		17.52	0.33			
Propionibacterium propionicum		6.40	0.24			
27573153	**Good identification**					**27573153**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27573160	**Very good identification**					**27573160**
Actinomyces naeslundii		99.08	0.75			
Actinomyces israelii		0.34	0.23			
Propionibacterium propionicum		0.29	0.21			
27573161	**Very good identification**					**27573161**
Actinomyces naeslundii		100.0	0.78			
27573162	**Very good identification**					**27573162**
Actinomyces naeslundii		98.91	0.75			
Actinomyces israelii		0.51	0.26			
Propionibacterium propionicum		0.44	0.24			
27573163	**Very good identification**					**27573163**
Actinomyces naeslundii		100.0	0.78			
27573170	**Good identification**					**27573170**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27573171	**Good identification**					**27573171**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27573172	**Good identification**					**27573172**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27573173	**Good identification**					**27573173**
Actinomyces naeslundii		100.0	0.56	MLZ 5		
27573200	**Low discrimination**					**27573200**
Actinomyces naeslundii		88.49	0.46	RHA 1		
Actinomyces israelii		11.51	0.21			
27573201	**Good identification**					**27573201**
Actinomyces naeslundii		100.0	0.49	RHA 1		
27573202	**Low discrimination**					**27573202**
Actinomyces naeslundii		83.67	0.46	RHA 1		
Actinomyces israelii		16.33	0.24			
27573203	**Good identification**					**27573203**
Actinomyces naeslundii		100.0	0.49	RHA 1		
27573220	**Good identification**					**27573220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	RHA 1		
27573221	**Good identification**					**27573221**
Actinomyces naeslundii		100.0	0.49	RHA 1		
27573222	**Good identification**					**27573222**
Actinomyces naeslundii		100.0	0.46	RHA 1		
27573223	**Good identification**					**27573223**
Actinomyces naeslundii		100.0	0.49	RHA 1		
27573241	**Good identification**					**27573241**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27573243	**Good identification**					**27573243**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27573261	**Good identification**					**27573261**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27573263	**Good identification**					**27573263**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27573300	**Low discrimination**					**27573300**
Actinomyces naeslundii		81.01	0.53	RHA 1		
Actinomyces israelii		14.81	0.32			
Propionibacterium propionicum		4.18	0.21			
27573301	**Very good identification**					**27573301**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27573302	**Low discrimination**					**27573302**
Actinomyces naeslundii		73.07	0.53	RHA 1		
Actinomyces israelii		20.04	0.35			
Propionibacterium propionicum		5.65	0.24			
27573303	**Very good identification**					**27573303**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27573320	**Low discrimination**					**27573320**
					SPO	
Actinomyces naeslundii		82.17	0.53	RHA 1	1	
Clostridium ramosum		17.83	0.17		92	
27573321	**Very good identification**					**27573321**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27573322	**Low discrimination**					**27573322**
					SPO	
Actinomyces naeslundii		53.04	0.53	RHA 1	1	
Clostridium ramosum		46.03	0.27		92	
Actinomyces israelii		0.93	0.14		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27573323	**Very good identification**					**27573323**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27573340	**Low discrimination**					**27573340**
Actinomyces naeslundii		58.99	0.41	RHA 1		
Actinomyces israelii		23.77	0.25			
Propionibacterium propionicum		17.24	0.21			
27573341	**Good identification**					**27573341**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27573342	**Low discrimination**					**27573342**
Actinomyces naeslundii		48.95	0.41	RHA 1		
Actinomyces israelii		29.58	0.28			
Propionibacterium propionicum		21.47	0.24			
27573343	**Good identification**					**27573343**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27573360	**Good identification**					**27573360**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27573361	**Good identification**					**27573361**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27573362	**Good identification**					**27573362**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27573363	**Good identification**					**27573363**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27573400	**Low discrimination**					**27573400**
Actinomyces israelii		58.58	0.39	SAL 92		
Actinomyces naeslundii		41.43	0.46	XYL 92		
27573401	**Good identification**					**27573401**
Actinomyces naeslundii		100.0	0.49	ARA 1		
27573402	**Low discrimination**					**27573402**
Actinomyces israelii		67.96	0.42	SAL 92		
Actinomyces naeslundii		32.04	0.46	XYL 92		
27573403	**Good identification**					**27573403**
Actinomyces naeslundii		100.0	0.49	ARA 1		
27573420	**Acceptable identification**					**27573420**
Actinomyces naeslundii		91.72	0.46	ARA 1		
Actinomyces israelii		8.28	0.19			
27573421	**Good identification**					**27573421**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	ARA 1		
27573422	**Low discrimination**					**27573422**
Actinomyces naeslundii		88.08	0.46	ARA 1		
Actinomyces israelii		11.92	0.22			
27573423	**Good identification**					**27573423**
Actinomyces naeslundii		100.0	0.49	ARA 1		
27573441	**Good identification**					**27573441**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27573442	**Low discrimination**					**27573442**
Actinomyces israelii		82.38	0.35	SAL 92		
Actinomyces naeslundii		17.63	0.33	XYL 92		
27573443	**Good identification**					**27573443**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27573461	**Good identification**					**27573461**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27573463	**Good identification**					**27573463**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27573500	**Low discrimination**					**27573500**
Actinomyces israelii		65.40	0.49	SAL 92		
Actinomyces naeslundii		32.91	0.53	XYL 92		
Propionibacterium propionicum		1.70	0.21			
27573501	**Very good identification**					**27573501**
Actinomyces naeslundii		98.68	0.56	ARA 1		
Actinomyces israelii		1.32	0.15			
27573502	**Low discrimination**					**27573502**
Actinomyces israelii		70.96	0.52	SAL 92		
Actinomyces naeslundii		23.80	0.53	XYL 92		
Bifidobacterium adolescentis		2.99	0.23			
27573503	**Very good identification**					**27573503**
Actinomyces naeslundii		98.03	0.56	ARA 1		
Actinomyces israelii		1.97	0.18			
27573510	**Low discrimination**					**27573510**
Actinomyces israelii		84.95	0.35	SAL 92		
Actinomyces naeslundii		15.06	0.32	XYL 92		
27573512	**Low discrimination**					**27573512**
Actinomyces israelii		68.34	0.38	SAL 92		
Bifidobacterium adolescentis		23.58	0.24	XYL 92		
Actinomyces naeslundii		8.08	0.32	MLZ 13		
27573520	**Low discrimination**					**27573520**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		88.74	0.53	ARA 1		
Actinomyces israelii		11.26	0.29			
27573521	**Very good identification**					**27573521**
Actinomyces naeslundii		100.0	0.56	ARA 1		
27573522	**Low discrimination**					**27573522**
Actinomyces naeslundii		84.01	0.53	ARA 1		
Actinomyces israelii		15.99	0.32			
27573523	**Very good identification**					**27573523**
Actinomyces naeslundii		100.0	0.56	ARA 1		
27573540	**Low discrimination**					**27573540**
Actinomyces israelii		77.22	0.42	SAL 92		
Actinomyces naeslundii		17.63	0.41	XYL 92		
Propionibacterium propionicum		5.15	0.21			
27573541	**Good identification**					**27573541**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27573542	**Low discrimination**					**27573542**
Actinomyces israelii		81.14	0.45	SAL 92		
Actinomyces naeslundii		12.35	0.41	XYL 92		
Propionibacterium propionicum		5.42	0.24			
27573543	**Good identification**					**27573543**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27573560	**Low discrimination**					**27573560**
Actinomyces naeslundii		78.15	0.41	ARA 1		
Actinomyces israelii		21.85	0.22			
27573561	**Good identification**					**27573561**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27573562	**Low discrimination**					**27573562**
Actinomyces naeslundii		70.46	0.41	ARA 1		
Actinomyces israelii		29.54	0.25			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27573563	**Good identification**					**27573563**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27574102	**Good identification**					**27574102**
Actinomyces viscosus		96.61	0.37	TRE 7		
Actinomyces israelii		2.63	0.25	XYL 1		
Actinomyces naeslundii		0.76	0.25			
27575000	**Low discrimination**					**27575000**
Actinomyces israelii		87.19	0.46	SAL 92		
Actinomyces naeslundii		12.81	0.41			
27575001	**Good identification**					**27575001**
Actinomyces naeslundii		100.0	0.44	XYL 1		
27575002	**Low discrimination**					**27575002**
Actinomyces israelii		62.13	0.49	SAL 92		
Actinomyces viscosus		31.78	0.29			
Actinomyces naeslundii		6.08	0.41			
27575020	**Low discrimination**					**27575020**
Actinomyces naeslundii		61.12	0.41	XYL 1		
Actinomyces israelii		26.56	0.25			
Clostridium perfringens		12.33	0.07			
27575021	**Good identification**					**27575021**
Actinomyces naeslundii		100.0	0.44	XYL 1		
27575022	**Low discrimination**					**27575022**
					GRA	
Actinomyces naeslundii		53.55	0.41	XYL 1	99	
Actinomyces israelii		34.90	0.28		99	
Parabacteroides distasonis		6.47	0.12		1	
27575023	**Good identification**					**27575023**
Actinomyces naeslundii		100.0	0.44	XYL 1		
27575040	**Acceptable identification**					**27575040**
Actinomyces israelii		93.75	0.39	SAL 92		
Actinomyces naeslundii		6.25	0.28			
27575042	**Good identification**					**27575042**
Actinomyces israelii		95.75	0.42	SAL 92		
Actinomyces naeslundii		4.26	0.28			
27575100	**Low discrimination**					**27575100**
					COC	
Actinomyces israelii		84.16	0.56	SAL 92	1	
Actinomyces naeslundii		8.80	0.49		1	
Atopobium parvulum		5.95	0.23		99	
27575101	**Acceptable identification**					**27575101**
Actinomyces naeslundii		93.95	0.52	XYL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		6.05	0.22			
27575102	**Low discrimination**					**27575102**
Actinomyces israelii		75.10	0.59	SAL 92		
Actinomyces viscosus		17.07	0.33			
Actinomyces naeslundii		5.23	0.49			
27575103	**Low discrimination**					**27575103**
Actinomyces naeslundii		54.89	0.52	XYL 1		
Actinomyces viscosus		39.80	0.25			
Actinomyces israelii		5.31	0.25			
27575110	**Good identification**					**27575110**
Actinomyces israelii		96.45	0.42	SAL 92		
Actinomyces naeslundii		3.55	0.27	MLZ 13		
27575112	**Low discrimination**					**27575112**
Actinomyces israelii		81.25	0.45	SAL 92		
Bifidobacterium adolescentis		13.81	0.26	MLZ 13		
Actinomyces naeslundii		1.99	0.27			
27575120	**Low discrimination**					**27575120**
Actinomyces naeslundii		43.62	0.49	XYL 1		
Actinomyces israelii		26.64	0.36			
Clostridium perfringens		22.68	0.21			
27575121	**Good identification**					**27575121**
Actinomyces naeslundii		100.0	0.52	XYL 1		
27575122	**Low discrimination**					**27575122**
					GRA	
Parabacteroides distasonis		60.05	0.46	NIT 1	1	
Actinomyces naeslundii		14.29	0.49		99	
Actinomyces israelii		13.09	0.39		99	
27575123	**Good identification**					**27575123**
					GRA	
Actinomyces naeslundii		97.25	0.52	XYL 1	99	
Parabacteroides distasonis		2.75	0.12		1	
27575132	**Acceptable identification**					**27575132**
					GRA	
Parabacteroides distasonis		90.61	0.42	NIT 1	1	
Actinomyces israelii		5.48	0.25		99	
Actinomyces naeslundii		2.11	0.27		99	
27575140	**Good identification**					**27575140**
Actinomyces israelii		95.47	0.49	SAL 92		
Actinomyces naeslundii		4.53	0.36			
27575141	**Low discrimination**					**27575141**
Actinomyces naeslundii		87.57	0.39	XYL 1		
Actinomyces israelii		12.43	0.15			
27575142	**Good identification**					**27575142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		95.77	0.52	SAL 92			
Actinomyces naeslundii		3.03	0.36				
Bifidobacterium adolescentis		0.63	0.09				
27575143	**Low discrimination**						**27575143**
Actinomyces naeslundii		82.44	0.39	XYL 1			
Actinomyces israelii		17.56	0.18				
27575152	**Acceptable identification**						**27575152**
Actinomyces israelii		94.87	0.38	SAL 92			
Bifidobacterium adolescentis		5.13	0.10	MLZ 13			
27575161	**Good identification**						**27575161**
Actinomyces naeslundii		100.0	0.39	XYL 1			
27575163	**Good identification**						**27575163**
Actinomyces naeslundii		100.0	0.39	XYL 1			
27575300	**Good identification**						**27575300**
Actinomyces israelii		100.0	0.37	SAL 92			
27575302	**Low discrimination**						**27575302**
					GRA		
Actinomyces israelii		82.29	0.40	SAL 92	99		
Parabacteroides distasonis		9.70	0.20	RHA 7	1		
Clostridium clostridioforme		5.51	0.10		50		
27575322	**Good identification**						**27575322**
Parabacteroides distasonis		98.05	0.54	NIT 1			
Prevotella oralis		0.85	0.06				
Clostridium clostridioforme		0.56	0.10				
27575332	**Good identification**						**27575332**
Parabacteroides distasonis		95.37	0.50	NIT 1			
Clostridium clostridioforme		4.63	0.22				
27575400	**Good identification**						**27575400**
					SPO		
Actinomyces israelii		98.26	0.44	SAL 92	1		
Clostridium butyricum		1.74	-.07		99		
27575402	**Good identification**						**27575402**
Actinomyces israelii		98.45	0.47	SAL 92			
Clostridium clostridioforme		1.55	0.07				
27575440	**Good identification**						**27575440**
Actinomyces israelii		100.0	0.37	SAL 92			
27575442	**Good identification**						**27575442**
Actinomyces israelii		100.0	0.40	SAL 92			
27575500	**Low discrimination**						**27575500**
					GRA	SPO	
Actinomyces israelii		85.53	0.55	SAL 92	99	1	
Clostridium butyricum		9.09	0.17		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		4.96	0.21		1	1	
27575502	**Low discrimination**						**27575502**
					SPO		
Actinomyces israelii		81.90	0.58	SAL 92	1		
Bifidobacterium adolescentis		13.92	0.38		1		
Clostridium butyricum		2.26	0.10		99		
27575510	**Low discrimination**						**27575510**
					GRA		
Actinomyces israelii		81.80	0.40	SAL 92	99		
Mitsuokella multacida		10.58	0.13	MLZ 13	1		
Bifidobacterium longum subsp. longum		4.56	0.11		99		
27575512	**Low discrimination**						**27575512**
Bifidobacterium adolescentis		49.16	0.40	NIT 1			
Actinomyces israelii		35.35	0.44				
Clostridium clostridioforme		14.17	0.27				
27575522	**Low discrimination**						**27575522**
					GRA		
Parabacteroides distasonis		69.58	0.43	NIT 1	1		
Actinomyces israelii		20.25	0.37		99		
Clostridium clostridioforme		4.17	0.16		50		
27575532	**Low discrimination**						**27575532**
					GRA		
Parabacteroides distasonis		56.31	0.38	NIT 1	1		
Clostridium clostridioforme		28.56	0.27		50		
Bifidobacterium longum subsp. longum		10.58	0.21		99		
27575540	**Low discrimination**						**27575540**
					GRA	SPO	
Actinomyces israelii		85.26	0.48	SAL 92	99	1	
Mitsuokella multacida		12.71	0.21		1	1	
Clostridium butyricum		2.03	-.01		99	99	
27575542	**Acceptable identification**						**27575542**
					GRA		
Actinomyces israelii		93.89	0.51	SAL 92	99		
Bifidobacterium adolescentis		5.07	0.23		99		
Mitsuokella multacida		1.04	0.04		1		
27575552	**Low discrimination**						**27575552**
Actinomyces israelii		69.35	0.36	SAL 92			
Bifidobacterium adolescentis		30.65	0.24	MLZ 13			
27575712	**Good identification**						**27575712**
					GRA		
Clostridium clostridioforme		96.17	0.42	NIT 1	50		
Actinomyces israelii		2.46	0.24	SAL 88	99		
Parabacteroides distasonis		0.78	0.12		1		
27575722	**Low discrimination**						**27575722**
Parabacteroides distasonis		80.70	0.51	NIT 1			
Clostridium clostridioforme		11.83	0.31				
Bacteroides thetaiotaomicron		4.68	0.14				
27575732	**Low discrimination**						**27575732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		53.55	0.42	NIT 1		
Parabacteroides distasonis		43.18	0.46	SAL 88		
Bacteroides thetaiotaomicron		1.81	0.07			
27576102	**Low discrimination**					**27576102**
Actinomyces israelii		74.25	0.38	SAL 92		
Propionibacterium propionicum		13.58	0.24	GAL 99		
Actinomyces naeslundii		7.31	0.30			
27576502	**Good identification**					**27576502**
Actinomyces israelii		100.0	0.36	SAL 92		
27577000	**Very good identification**					**27577000**
Actinomyces israelii		95.21	0.59	SAL 92		
Actinomyces naeslundii		4.79	0.46			
27577001	**Low discrimination**					**27577001**
Actinomyces naeslundii		88.20	0.49	XYL 1		
Actinomyces israelii		11.80	0.25			
27577002	**Very good identification**					**27577002**
Actinomyces israelii		96.76	0.62	SAL 92		
Actinomyces naeslundii		3.25	0.46			
27577003	**Low discrimination**					**27577003**
Actinomyces naeslundii		83.28	0.49	XYL 1		
Actinomyces israelii		16.72	0.28			
27577010	**Good identification**					**27577010**
Actinomyces israelii		98.26	0.45	SAL 92		
Actinomyces naeslundii		1.74	0.24	MLZ 13		
27577012	**Good identification**					**27577012**
Actinomyces israelii		98.83	0.48	SAL 92		
Actinomyces naeslundii		1.17	0.24	MLZ 13		
27577020	**Low discrimination**					**27577020**
Actinomyces israelii		55.92	0.38	SAL 92		
Actinomyces naeslundii		44.08	0.46	NAG 6		
27577021	**Good identification**					**27577021**
Actinomyces naeslundii		100.0	0.49	XYL 1		
27577022	**Low discrimination**					**27577022**
Actinomyces israelii		65.55	0.41	SAL 92		
Actinomyces naeslundii		34.45	0.46	NAG 6		
27577023	**Good identification**					**27577023**
Actinomyces naeslundii		100.0	0.49	XYL 1		
27577040	**Good identification**					**27577040**
Actinomyces israelii		97.77	0.52	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		2.23	0.33			
27577041	**Low discrimination**					**27577041**
Actinomyces naeslundii		77.23	0.36	XYL 1		
Actinomyces israelii		22.78	0.18			
27577042	**Good identification**					**27577042**
Actinomyces israelii		98.50	0.55	SAL 92		
Actinomyces naeslundii		1.50	0.33			
27577043	**Low discrimination**					**27577043**
Actinomyces naeslundii		69.33	0.36	XYL 1		
Actinomyces israelii		30.67	0.21			
27577050	**Good identification**					**27577050**
Actinomyces israelii		100.0	0.38	SAL 92		
27577052	**Good identification**					**27577052**
Actinomyces israelii		100.0	0.41	SAL 92		
27577061	**Good identification**					**27577061**
Actinomyces naeslundii		100.0	0.36	XYL 1		
27577063	**Good identification**					**27577063**
Actinomyces naeslundii		100.0	0.36	XYL 1		
27577100	**Very good identification**					**27577100**
Actinomyces israelii		96.37	0.69	SAL 92		
Actinomyces naeslundii		3.45	0.53			
Propionibacterium propionicum		0.18	0.21			
27577101	**Low discrimination**					**27577101**
Actinomyces naeslundii		84.17	0.56	XYL 1		
Actinomyces israelii		15.83	0.35			
27577102	**Very good identification**					**27577102**
Actinomyces israelii		97.17	0.72	SAL 92		
Actinomyces naeslundii		2.32	0.53			
Bifidobacterium adolescentis		0.29	0.23			
27577103	**Low discrimination**					**27577103**
Actinomyces naeslundii		78.00	0.56	XYL 1		
Actinomyces israelii		22.01	0.38			
27577110	**Good identification**					**27577110**
Actinomyces israelii		98.76	0.55	SAL 92		
Actinomyces naeslundii		1.25	0.32	MLZ 13		
27577112	**Very good identification**					**27577112**
Actinomyces israelii		96.81	0.58	SAL 92		
Bifidobacterium adolescentis		2.38	0.24	MLZ 13		
Actinomyces naeslundii		0.81	0.32			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27577120	**Low discrimination**					**27577120**
Actinomyces israelii		64.07	0.49	SAL 92		
Actinomyces naeslundii		35.93	0.53	NAG 6		
27577121	**Very good identification**					**27577121**
Actinomyces naeslundii		98.81	0.56	XYL 1		
Actinomyces israelii		1.19	0.14			
27577122	**Low discrimination**					**27577122**
					GRA	
Actinomyces israelii		72.35	0.52	SAL 92	99	
Actinomyces naeslundii		27.06	0.53	NAG 6	99	
Parabacteroides distasonis		0.59	0.12		1	
27577123	**Very good identification**					**27577123**
Actinomyces naeslundii		98.23	0.56	XYL 1		
Actinomyces israelii		1.77	0.17			
27577132	**Low discrimination**					**27577132**
Actinomyces israelii		88.36	0.37	SAL 92		
Actinomyces naeslundii		11.64	0.32	MLZ 13		
27577140	**Very good identification**					**27577140**
Actinomyces israelii		97.94	0.62	SAL 92		
Actinomyces naeslundii		1.59	0.41			
Propionibacterium propionicum		0.47	0.21			
27577141	**Low discrimination**					**27577141**
Actinomyces naeslundii		70.70	0.44	XYL 1		
Actinomyces israelii		29.30	0.28			
27577142	**Very good identification**					**27577142**
Actinomyces israelii		98.37	0.65	SAL 92		
Actinomyces naeslundii		1.07	0.41			
Propionibacterium propionicum		0.47	0.24			
27577143	**Low discrimination**					**27577143**
Actinomyces naeslundii		61.66	0.44	XYL 1		
Actinomyces israelii		38.34	0.31			
27577150	**Good identification**					**27577150**
Actinomyces israelii		100.0	0.48	SAL 92		
27577152	**Good identification**					**27577152**
Actinomyces israelii		99.23	0.51	SAL 92		
Bifidobacterium adolescentis		0.77	0.09	MLZ 13		
27577160	**Low discrimination**					**27577160**
Actinomyces israelii		79.71	0.42	SAL 92		
Actinomyces naeslundii		20.29	0.41	NAG 6		
27577161	**Good identification**					**27577161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	XYL 1		
27577162	**Low discrimination**					**27577162**
Actinomyces israelii		85.49	0.45	SAL 92		
Actinomyces naeslundii		14.51	0.41	NAG 6		
27577163	**Good identification**					**27577163**
Actinomyces naeslundii		100.0	0.44	XYL 1		
27577200	**Good identification**					**27577200**
Actinomyces israelii		100.0	0.39	SAL 92		
27577202	**Good identification**					**27577202**
Actinomyces israelii		100.0	0.42	SAL 92		
27577242	**Good identification**					**27577242**
Actinomyces israelii		100.0	0.35	SAL 92		
27577300	**Good identification**					**27577300**
Actinomyces israelii		100.0	0.50	SAL 92		
27577302	**Good identification**					**27577302**
Actinomyces israelii		100.0	0.53	SAL 92		
27577310	**Good identification**					**27577310**
Actinomyces israelii		100.0	0.36	SAL 92		
27577312	**Good identification**					**27577312**
Actinomyces israelii		100.0	0.39	SAL 92		
27577340	**Good identification**					**27577340**
Actinomyces israelii		100.0	0.43	SAL 92		
27577342	**Good identification**					**27577342**
Actinomyces israelii		100.0	0.46	SAL 92		
27577400	**Very good identification**					**27577400**
Actinomyces israelii		100.0	0.57	SAL 92		
27577402	**Very good identification**					**27577402**
Actinomyces israelii		100.0	0.60	SAL 92		
27577410	**Good identification**					**27577410**
Actinomyces israelii		100.0	0.43	SAL 92		
27577412	**Good identification**					**27577412**
Actinomyces israelii		100.0	0.46	SAL 92		
27577420	**Good identification**					**27577420**
Actinomyces israelii		100.0	0.37	SAL 92		
27577422	**Good identification**					**27577422**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	SAL 92			
27577440	**Good identification**						**27577440**
Actinomyces israelii		100.0	0.50	SAL 92			
27577442	**Good identification**						**27577442**
Actinomyces israelii		100.0	0.53	SAL 92			
27577450	**Good identification**						**27577450**
Actinomyces israelii		100.0	0.36	SAL 92			
27577452	**Good identification**						**27577452**
Actinomyces israelii		100.0	0.39	SAL 92			
27577500	**Very good identification**						**27577500**
					GRA	SPO	
Actinomyces israelii		97.47	0.68	SAL 92	99	1	
Mitsuokella multacida		2.33	0.27		1	1	
Clostridium butyricum		0.20	0.01		99	99	
27577502	**Very good identification**						**27577502**
					GRA		
Actinomyces israelii		97.39	0.71	SAL 92	99		
Bifidobacterium adolescentis		2.39	0.37		99		
Mitsuokella multacida		0.17	0.11		1		
27577503	**Good identification**						**27577503**
Actinomyces israelii		100.0	0.36	SAL 92			
27577510	**Acceptable identification**						**27577510**
					GRA		
Actinomyces israelii		94.94	0.53	SAL 92	99		
Mitsuokella multacida		5.06	0.19	MLZ 13	1		
27577512	**Low discrimination**						**27577512**
Actinomyces israelii		82.64	0.56	SAL 92			
Bifidobacterium adolescentis		16.59	0.38	MLZ 13			
Bifidobacterium dentium		0.45	-.07				
27577520	**Acceptable identification**						**27577520**
					GRA		
Actinomyces israelii		91.45	0.47	SAL 92	99		
Mitsuokella multacida		8.55	0.17	NAG 6	1		
27577522	**Good identification**						**27577522**
Actinomyces israelii		100.0	0.50	SAL 92			
27577532	**Low discrimination**						**27577532**
Actinomyces israelii		86.92	0.36	SAL 92			
Bifidobacterium dentium		13.08	-.03	MLZ 13			
27577540	**Good identification**						**27577540**
					GRA		
Actinomyces israelii		94.22	0.61	SAL 92	99		
Mitsuokella multacida		5.78	0.27		1		
27577542	**Very good identification**						**27577542**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				GRA		
Actinomyces israelii	98.78	0.64	SAL 92	99		
Bifidobacterium adolescentis	0.77	0.21		99		
Mitsuokella multacida	0.45	0.11		1		
27577550 Low discrimination						**27577550**
				COC	GRA	
Actinomyces israelii	86.39	0.46	SAL 92	1	99	
Mitsuokella multacida	11.83	0.19	MLZ 13	1	1	
Blautia producta	1.78	-.03		99	99	
27577552 Acceptable identification						**27577552**
				GRA		
Actinomyces israelii	93.11	0.49	SAL 92	99		
Bifidobacterium adolescentis	5.94	0.23	MLZ 13	99		
Mitsuokella multacida	0.95	0.03		1		
27577560 Low discrimination						**27577560**
				GRA		
Actinomyces israelii	80.62	0.40	SAL 92	99		
Mitsuokella multacida	19.38	0.17	NAG 6	1		
27577562 Good identification						**27577562**
Actinomyces israelii	100.0	0.43	SAL 92			
27577600 Good identification						**27577600**
Actinomyces israelii	100.0	0.38	SAL 92			
27577602 Good identification						**27577602**
Actinomyces israelii	100.0	0.41	SAL 92			
27577700 Low discrimination						**27577700**
				GRA		
Actinomyces israelii	62.93	0.48	SAL 92	99		
Mitsuokella multacida	37.07	0.32	RHA 7	1		
27577702 Good identification						**27577702**
				GRA		
Actinomyces israelii	95.82	0.51	SAL 92	99		
Mitsuokella multacida	4.18	0.15	RHA 7	1		
27577712 Low discrimination						**27577712**
				GRA		
Actinomyces israelii	85.69	0.37	SAL 92	99		
Mitsuokella multacida	8.34	0.07	RHA 7	1		
Clostridium clostridioforme	5.97	0.08	MLZ 13	50		
27577742 Low discrimination						**27577742**
				GRA		
Actinomyces israelii	89.91	0.44	SAL 92	99		
Mitsuokella multacida	10.09	0.15	RHA 7	1		
27601001 Good identification						**27601001**
Actinomyces naeslundii	100.0	0.38	MLT 94			
27601021 Good identification						**27601021**
Actinomyces naeslundii	100.0	0.38	MLT 94			
27601023 Good identification						**27601023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MLT 94		
27601100	**Low discrimination**					**27601100**
					GRA	
Actinomyces naeslundii		58.11	0.43	MLT 94	99	
Fusobacterium mortiferum		41.89	0.29	FRU 95	1	
27601101	**Good identification**					**27601101**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27601120	**Good identification**					**27601120**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27601121	**Good identification**					**27601121**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27601122	**Good identification**					**27601122**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27601123	**Good identification**					**27601123**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27603000	**Good identification**					**27603000**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27603001	**Good identification**					**27603001**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27603002	**Good identification**					**27603002**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27603003	**Good identification**					**27603003**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27603020	**Good identification**					**27603020**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27603021	**Good identification**					**27603021**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27603022	**Good identification**					**27603022**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27603023	**Good identification**					**27603023**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27603100	**Good identification**					**27603100**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27603101	**Good identification**					**27603101**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27603102	**Good identification**					**27603102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.48	MLT 94			
27603103	**Good identification**						**27603103**
Actinomyces naeslundii		100.0	0.51	MLT 94			
27603120	**Good identification**						**27603120**
Actinomyces naeslundii		100.0	0.48	MLT 94			
27603121	**Good identification**						**27603121**
Actinomyces naeslundii		100.0	0.51	MLT 94			
27603122	**Good identification**						**27603122**
Actinomyces naeslundii		100.0	0.48	MLT 94			
27603123	**Good identification**						**27603123**
Actinomyces naeslundii		100.0	0.51	MLT 94			
27603141	**Good identification**						**27603141**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27603143	**Good identification**						**27603143**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27603161	**Good identification**						**27603161**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27603163	**Good identification**						**27603163**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27610000	**Acceptable identification**						**27610000**
					GRA	CAT	
Fusobacterium mortiferum		90.77	0.54	NIT 1	1	5	
Actinomyces viscosus		3.90	0.18		99	76	
Propionibacterium granulosum		2.49	0.24		99	90	
27610001	**Low discrimination**						**27610001**
					GRA		
Actinomyces naeslundii		45.57	0.36	MLT 94	99		
Actinomyces viscosus		31.93	0.09	GAL 96	99		
Fusobacterium mortiferum		22.50	0.19		1		
27610002	**Good identification**						**27610002**
					GRA		
Actinomyces viscosus		96.76	0.52	MLT 99	99		
Fusobacterium mortiferum		2.53	0.37		1		
Actinomyces naeslundii		0.31	0.33		99		
27610003	**Good identification**						**27610003**
Actinomyces viscosus		98.58	0.44	MLT 99			
Actinomyces naeslundii		1.42	0.36				
27610021	**Good identification**						**27610021**
Actinomyces naeslundii		100.0	0.36	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27610023	**Low discrimination**						**27610023**
Actinomyces naeslundii		58.80	0.36	MLT 94			
Actinomyces viscosus		41.20	0.09	GAL 96			
27610100	**Very good identification**						**27610100**
					GRA		
Fusobacterium mortiferum		95.08	0.60	NIT 1	1		
Actinomyces viscosus		3.12	0.22		99		
Actinomyces naeslundii		1.58	0.41		99		
27610101	**Low discrimination**						**27610101**
					GRA		
Actinomyces naeslundii		54.28	0.44	MLT 94	99		
Actinomyces viscosus		23.75	0.14	GAL 96	99		
Fusobacterium mortiferum		21.97	0.26		1		
27610102	**Good identification**						**27610102**
					GRA		
Actinomyces viscosus		96.17	0.56	MLT 99	99		
Fusobacterium mortiferum		3.29	0.44		1		
Actinomyces naeslundii		0.49	0.41		99		
27610103	**Good identification**						**27610103**
Actinomyces viscosus		97.74	0.48	MLT 99			
Actinomyces naeslundii		2.26	0.44				
27610120	**Low discrimination**						**27610120**
					GRA		
Actinomyces naeslundii		62.22	0.41	MLT 94	99		
Fusobacterium mortiferum		37.78	0.26	GAL 96	1		
27610121	**Good identification**						**27610121**
Actinomyces naeslundii		100.0	0.44	MLT 94			
27610123	**Low discrimination**						**27610123**
Actinomyces naeslundii		69.56	0.44	MLT 94			
Actinomyces viscosus		30.44	0.14	GAL 96			
27611000	**Low discrimination**						**27611000**
					GRA	CAT	
Fusobacterium mortiferum		79.24	0.57	NIT 1	1	5	
Actinomyces naeslundii		17.29	0.57		99	25	
Propionibacterium granulosum		1.45	0.24		99	90	
27611001	**Very good identification**						**27611001**
					GRA		
Actinomyces naeslundii		95.35	0.60	MLT 94	99		
Fusobacterium mortiferum		2.94	0.22		1		
Actinomyces viscosus		1.71	0.06		99		
27611002	**Low discrimination**						**27611002**
					GRA		
Actinomyces viscosus		83.70	0.48	MLT 99	99		
Actinomyces naeslundii		10.50	0.57		99		
Fusobacterium mortiferum		5.35	0.40		1		
27611003	**Low discrimination**						**27611003**
Actinomyces viscosus		63.92	0.40	MLT 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		36.08	0.60			
27611011	**Good identification**					**27611011**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27611013	**Low discrimination**					**27611013**
Actinomyces naeslundii		74.63	0.38	MLT 94		
Actinomyces viscosus		25.37	0.06	MLZ 5		
27611020	**Low discrimination**					**27611020**
					GRA	
Actinomyces naeslundii		87.56	0.57	MLT 94	99	
Clostridium perfringens		6.55	0.15		99	
Fusobacterium mortiferum		4.05	0.22		1	
27611021	**Very good identification**					**27611021**
Actinomyces naeslundii		100.0	0.60	MLT 94		
27611022	**Low discrimination**					**27611022**
Actinomyces naeslundii		89.63	0.57	MLT 94		
Actinomyces viscosus		7.22	0.14			
Clostridium perfringens		3.16	0.10			
27611023	**Very good identification**					**27611023**
Actinomyces naeslundii		98.24	0.60	MLT 94		
Actinomyces viscosus		1.76	0.06			
27611031	**Good identification**					**27611031**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27611033	**Good identification**					**27611033**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27611040	**Low discrimination**					**27611040**
					GRA	
Actinomyces naeslundii		79.22	0.44	MLT 94	99	
Fusobacterium mortiferum		20.78	0.22		1	
27611041	**Good identification**					**27611041**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27611042	**Low discrimination**					**27611042**
Actinomyces naeslundii		68.67	0.44	MLT 94		
Actinomyces viscosus		31.33	0.14			
27611043	**Acceptable identification**					**27611043**
Actinomyces naeslundii		90.79	0.47	MLT 94		
Actinomyces viscosus		9.21	0.06			
27611060	**Acceptable identification**					**27611060**
Actinomyces naeslundii		92.52	0.44	MLT 94		
Clostridium perfringens		7.48	0.03			
27611061	**Good identification**					**27611061**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.47	MLT 94		
27611062	**Good identification**					**27611062**
	Actinomyces naeslundii	100.0	0.44	MLT 94		
27611063	**Good identification**					**27611063**
	Actinomyces naeslundii	100.0	0.47	MLT 94		
27611100	**Low discrimination**					**27611100**
					GRA	
	Fusobacterium mortiferum	77.91	0.63	NIT 1	1	
	Actinomyces naeslundii	20.74	0.65		99	
	Actinomyces viscosus	1.04	0.18		99	
27611101	**Very good identification**					**27611101**
					GRA	
	Actinomyces naeslundii	96.48	0.68	MLT 94	99	
	Fusobacterium mortiferum	2.44	0.29		1	
	Actinomyces viscosus	1.08	0.10		99	
27611102	**Low discrimination**					**27611102**
					GRA	
	Actinomyces viscosus	77.74	0.52	MLT 99	99	
	Actinomyces naeslundii	15.61	0.65		99	
	Fusobacterium mortiferum	6.52	0.47		1	
27611103	**Low discrimination**					**27611103**
					GRA	
	Actinomyces viscosus	52.46	0.44	MLT 99	99	
	Actinomyces naeslundii	47.41	0.68		99	
	Fusobacterium mortiferum	0.13	0.12		1	
27611110	**Low discrimination**					**27611110**
					GRA	
	Actinomyces naeslundii	58.11	0.43	MLT 94	99	
	Fusobacterium mortiferum	41.89	0.29	MLZ 5	1	
27611111	**Good identification**					**27611111**
	Actinomyces naeslundii	100.0	0.46	MLT 94		
27611112	**Low discrimination**					**27611112**
					GRA	
	Actinomyces naeslundii	49.12	0.43	MLT 94	99	
	Actinomyces viscosus	46.94	0.18	MLZ 5	99	
	Fusobacterium mortiferum	3.94	0.12		1	
27611113	**Low discrimination**					**27611113**
	Actinomyces naeslundii	82.48	0.46	MLT 94		
	Actinomyces viscosus	17.52	0.10	MLZ 5		
27611120	**Low discrimination**					**27611120**
					GRA	
	Actinomyces naeslundii	80.84	0.65	MLT 94	99	
	Clostridium perfringens	15.59	0.30		99	
	Fusobacterium mortiferum	3.07	0.29		1	
27611121	**Very good identification**					**27611121**
	Actinomyces naeslundii	100.0	0.68	MLT 94		
27611122	**Low discrimination**					**27611122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.70	0.65	MLT 94		
Clostridium perfringens		7.69	0.24			
Actinomyces viscosus		4.26	0.18			
27611123	**Very good identification**					**27611123**
Actinomyces naeslundii		98.90	0.68	MLT 94		
Actinomyces viscosus		1.11	0.10			
27611130	**Good identification**					**27611130**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27611131	**Good identification**					**27611131**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27611132	**Acceptable identification**					**27611132**
					GRA	
Actinomyces naeslundii		93.60	0.43	MLT 94	99	
Parabacteroides distasonis		6.40	0.10	MLZ 5	1	
27611133	**Good identification**					**27611133**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27611140	**Low discrimination**					**27611140**
					GRA	
Actinomyces naeslundii		82.30	0.52	MLT 94	99	
Fusobacterium mortiferum		17.70	0.29		1	
27611141	**Good identification**					**27611141**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27611142	**Low discrimination**					**27611142**
					GRA	
Actinomyces naeslundii		76.40	0.52	MLT 94	99	
Actinomyces viscosus		21.78	0.18		99	
Fusobacterium mortiferum		1.83	0.12		1	
27611143	**Acceptable identification**					**27611143**
Actinomyces naeslundii		94.04	0.55	MLT 94		
Actinomyces viscosus		5.96	0.10			
27611160	**Low discrimination**					**27611160**
Actinomyces naeslundii		82.77	0.52	MLT 94		
Clostridium perfringens		17.23	0.18			
27611161	**Good identification**					**27611161**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27611162	**Acceptable identification**					**27611162**
Actinomyces naeslundii		91.08	0.52	MLT 94		
Clostridium perfringens		8.92	0.12			
27611163	**Good identification**					**27611163**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27612000	**Low discrimination**					**27612000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces naeslundii		46.22	0.38	MLT 94	99	25	
Propionibacterium granulosum		36.14	0.21	GAL 96	99	90	
Fusobacterium mortiferum		17.64	0.19		1	5	
27612001	**Good identification**						**27612001**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27612003	**Low discrimination**						**27612003**
Actinomyces naeslundii		73.48	0.41	MLT 94			
Actinomyces viscosus		26.52	0.09	GAL 96			
27612020	**Low discrimination**						**27612020**
Actinomyces naeslundii		83.65	0.38	MLT 94			
Propionibacterium granulosum		16.35	0.11	GAL 96			
27612021	**Good identification**						**27612021**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27612022	**Good identification**						**27612022**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27612023	**Good identification**						**27612023**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27612100	**Low discrimination**						**27612100**
					GRA		
Actinomyces naeslundii		67.27	0.46	MLT 94	99		
Fusobacterium mortiferum		21.04	0.26	GAL 96	1		
Propionibacterium propionicum		11.69	0.23		99		
27612101	**Good identification**						**27612101**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27612103	**Low discrimination**						**27612103**
Actinomyces naeslundii		81.60	0.49	MLT 94			
Actinomyces viscosus		18.40	0.14	GAL 96			
27612120	**Good identification**						**27612120**
Actinomyces naeslundii		100.0	0.46	MLT 94			
27612121	**Good identification**						**27612121**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27612122	**Good identification**						**27612122**
Actinomyces naeslundii		100.0	0.46	MLT 94			
27612123	**Good identification**						**27612123**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27612141	**Good identification**						**27612141**
Actinomyces naeslundii		100.0	0.36	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27612143	**Good identification**						**27612143**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27612161	**Good identification**						**27612161**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27612163	**Good identification**						**27612163**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27613000	**Good identification**						**27613000**
					GRA	CAT	
Actinomyces naeslundii		94.66	0.62	MLT 94	99	25	
Propionibacterium granulosum		3.08	0.21		99	90	
Fusobacterium mortiferum		2.26	0.22		1	5	
27613001	**Very good identification**						**27613001**
Actinomyces naeslundii		100.0	0.65	MLT 94			
27613002	**Very good identification**						**27613002**
Actinomyces naeslundii		95.27	0.62	MLT 94			
Actinomyces viscosus		3.95	0.14				
Propionibacterium granulosum		0.78	0.11				
27613003	**Very good identification**						**27613003**
Actinomyces naeslundii		99.09	0.65	MLT 94			
Actinomyces viscosus		0.91	0.06				
27613010	**Good identification**						**27613010**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27613011	**Good identification**						**27613011**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27613012	**Good identification**						**27613012**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27613013	**Good identification**						**27613013**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27613020	**Very good identification**						**27613020**
Actinomyces naeslundii		99.19	0.62	MLT 94			
Propionibacterium granulosum		0.81	0.11				
27613021	**Very good identification**						**27613021**
Actinomyces naeslundii		100.0	0.65	MLT 94			
27613022	**Very good identification**						**27613022**
Actinomyces naeslundii		100.0	0.62	MLT 94			
27613023	**Very good identification**						**27613023**
Actinomyces naeslundii		100.0	0.65	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27613030	**Good identification**					**27613030**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27613031	**Good identification**					**27613031**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27613032	**Good identification**					**27613032**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27613033	**Good identification**					**27613033**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27613040	**Good identification**					**27613040**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27613041	**Good identification**					**27613041**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27613042	**Good identification**					**27613042**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27613043	**Good identification**					**27613043**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27613060	**Good identification**					**27613060**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27613061	**Good identification**					**27613061**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27613062	**Good identification**					**27613062**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27613063	**Good identification**					**27613063**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27613100	**Very good identification**					**27613100**
					GRA	
Actinomyces naeslundii		97.39	0.70	MLT 94	99	
Fusobacterium mortiferum		1.90	0.29		1	
Propionibacterium propionicum		0.71	0.23		99	
27613101	**Very good identification**					**27613101**
Actinomyces naeslundii		100.0	0.73	MLT 94		
27613102	**Very good identification**					**27613102**
Actinomyces naeslundii		96.25	0.70	MLT 94		
Actinomyces viscosus		2.49	0.18			
Propionibacterium propionicum		1.05	0.26			
27613103	**Very good identification**					**27613103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.43	0.73	MLT 94		
Actinomyces viscosus		0.57	0.10			
27613110	**Good identification**					**27613110**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27613111	**Good identification**					**27613111**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27613112	**Good identification**					**27613112**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27613113	**Good identification**					**27613113**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27613120	**Very good identification**					**27613120**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27613121	**Very good identification**					**27613121**
Actinomyces naeslundii		100.0	0.73	MLT 94		
27613122	**Very good identification**					**27613122**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27613123	**Very good identification**					**27613123**
Actinomyces naeslundii		100.0	0.73	MLT 94		
27613130	**Good identification**					**27613130**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27613131	**Good identification**					**27613131**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27613132	**Good identification**					**27613132**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27613133	**Good identification**					**27613133**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27613140	**Very good identification**					**27613140**
Actinomyces naeslundii		96.06	0.57	MLT 94		
Propionibacterium propionicum		3.94	0.23			
27613141	**Very good identification**					**27613141**
Actinomyces naeslundii		100.0	0.60	MLT 94		
27613142	**Good identification**					**27613142**
Actinomyces naeslundii		94.20	0.57	MLT 94		
Propionibacterium propionicum		5.80	0.26			
27613143	**Very good identification**					**27613143**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.60	MLT 94		
27613151	**Good identification**					**27613151**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613153	**Good identification**					**27613153**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613160	**Very good identification**					**27613160**
	Actinomyces naeslundii	100.0	0.57	MLT 94		
27613161	**Very good identification**					**27613161**
	Actinomyces naeslundii	100.0	0.60	MLT 94		
27613162	**Very good identification**					**27613162**
	Actinomyces naeslundii	100.0	0.57	MLT 94		
27613163	**Very good identification**					**27613163**
	Actinomyces naeslundii	100.0	0.60	MLT 94		
27613171	**Good identification**					**27613171**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613173	**Good identification**					**27613173**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613300	**Good identification**					**27613300**
	Actinomyces naeslundii	100.0	0.35	MLT 94		
27613301	**Good identification**					**27613301**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613302	**Good identification**					**27613302**
	Actinomyces naeslundii	100.0	0.35	MLT 94		
27613303	**Good identification**					**27613303**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613320	**Good identification**					**27613320**
	Actinomyces naeslundii	100.0	0.35	MLT 94		
27613321	**Good identification**					**27613321**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613322	**Good identification**					**27613322**
	Actinomyces naeslundii	100.0	0.35	MLT 94		
27613323	**Good identification**					**27613323**
	Actinomyces naeslundii	100.0	0.38	MLT 94		
27613500	**Good identification**					**27613500**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	MLT 94		
27613501	**Good identification**					**27613501**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27613502	**Good identification**					**27613502**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27613503	**Good identification**					**27613503**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27613520	**Good identification**					**27613520**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27613521	**Good identification**					**27613521**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27613522	**Good identification**					**27613522**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27613523	**Good identification**					**27613523**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27614100	**Good identification**					**27614100**
Fusobacterium mortiferum		100.0	0.41	NIT 1		
27615000	**Acceptable identification**					**27615000**
					GRA	
Fusobacterium mortiferum		92.54	0.37	NIT 1	1	
Actinomyces odontolyticus		4.75	0.10	XYL 7	99	
Actinomyces naeslundii		2.71	0.23		99	
27615100	**Good identification**					**27615100**
					GRA	
Fusobacterium mortiferum		96.55	0.44	NIT 1	1	
Actinomyces naeslundii		3.45	0.30	XYL 7	99	
27615322	**Good identification**					**27615322**
Parabacteroides distasonis		100.0	0.39	NIT 1		
27615722	**Good identification**					**27615722**
Parabacteroides distasonis		100.0	0.35	NIT 1		
27617100	**Low discrimination**					**27617100**
Actinomyces naeslundii		57.53	0.35	MLT 94		
Actinomyces israelii		42.47	0.24	XYL 1		
27617101	**Good identification**					**27617101**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27617103	**Good identification**					**27617103**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27617120	**Good identification**					**27617120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	MLT 94		
27617121	**Good identification**					**27617121**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27617122	**Good identification**					**27617122**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27617123	**Good identification**					**27617123**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27621000	**Low discrimination**					**27621000**
					GRA	
Actinomyces naeslundii		88.08	0.40	MLT 94	99	
Fusobacterium mortiferum		11.92	0.13	FRU 95	1	
27621001	**Good identification**					**27621001**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27621002	**Low discrimination**					**27621002**
Actinomyces naeslundii		76.39	0.40	MLT 94		
Actinomyces viscosus		23.61	0.07	FRU 95		
27621003	**Acceptable identification**					**27621003**
Actinomyces naeslundii		93.57	0.43	MLT 94		
Actinomyces viscosus		6.43	-.01	FRU 95		
27621020	**Good identification**					**27621020**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27621021	**Good identification**					**27621021**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27621022	**Good identification**					**27621022**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27621023	**Good identification**					**27621023**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27621100	**Acceptable identification**					**27621100**
					GRA	
Actinomyces naeslundii		90.01	0.48	MLT 94	99	
Fusobacterium mortiferum		9.99	0.20	FRU 95	1	
27621101	**Good identification**					**27621101**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27621102	**Low discrimination**					**27621102**
Actinomyces naeslundii		83.82	0.48	MLT 94		
Actinomyces viscosus		16.18	0.11	FRU 95		
27621103	**Good identification**					**27621103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.89	0.51	MLT 94		
Actinomyces viscosus		4.11	0.03	FRU 95		
27621120	**Good identification**					**27621120**
Actinomyces naeslundii		96.58	0.48	MLT 94		
Clostridium perfringens		3.42	0.01	FRU 95		
27621121	**Good identification**					**27621121**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27621122	**Good identification**					**27621122**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27621123	**Good identification**					**27621123**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27621141	**Good identification**					**27621141**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27621143	**Good identification**					**27621143**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27621161	**Good identification**					**27621161**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27621163	**Good identification**					**27621163**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27623000	**Good identification**					**27623000**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27623001	**Good identification**					**27623001**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27623002	**Good identification**					**27623002**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27623003	**Good identification**					**27623003**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27623020	**Good identification**					**27623020**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27623021	**Good identification**					**27623021**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27623022	**Good identification**					**27623022**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27623023	**Good identification**					**27623023**
Actinomyces naeslundii		100.0	0.48	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27623100	**Good identification**						**27623100**
Actinomyces naeslundii		100.0	0.53	MLT 94			
27623101	**Good identification**						**27623101**
Actinomyces naeslundii		100.0	0.56	MLT 94			
27623102	**Good identification**						**27623102**
Actinomyces naeslundii		100.0	0.53	MLT 94			
27623103	**Good identification**						**27623103**
Actinomyces naeslundii		100.0	0.56	MLT 94			
27623120	**Good identification**						**27623120**
Actinomyces naeslundii		100.0	0.53	MLT 94			
27623121	**Good identification**						**27623121**
Actinomyces naeslundii		100.0	0.56	MLT 94			
27623122	**Good identification**						**27623122**
Actinomyces naeslundii		100.0	0.53	MLT 94			
27623123	**Good identification**						**27623123**
Actinomyces naeslundii		100.0	0.56	MLT 94			
27623140	**Good identification**						**27623140**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27623141	**Good identification**						**27623141**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27623142	**Good identification**						**27623142**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27623143	**Good identification**						**27623143**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27623160	**Good identification**						**27623160**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27623161	**Good identification**						**27623161**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27623162	**Good identification**						**27623162**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27623163	**Good identification**						**27623163**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27630000	**Low discrimination**						**27630000**
					GRA	CAT	
Fusobacterium mortiferum		72.72	0.45	NIT 1	1	5	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		20.04	0.32		99	90	
Actinomyces naeslundii		6.45	0.38		99	25	
27630001	**Good identification**						**27630001**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27630020	**Low discrimination**						**27630020**
Actinomyces naeslundii		56.26	0.38	MLT 94			
Propionibacterium granulosum		43.74	0.21	GAL 96			
27630021	**Good identification**						**27630021**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27630022	**Low discrimination**						**27630022**
Actinomyces naeslundii		75.96	0.38	MLT 94			
Propionibacterium granulosum		14.76	0.11	GAL 96			
Actinomyces viscosus		9.29	-.02				
27630023	**Good identification**						**27630023**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27630100	**Low discrimination**						**27630100**
					GRA		
Fusobacterium mortiferum		89.58	0.51	NIT 1	1		
Actinomyces naeslundii		9.69	0.46		99		
Actinomyces viscosus		0.74	0.03		99		
27630101	**Acceptable identification**						**27630101**
					GRA		
Actinomyces naeslundii		94.14	0.49	MLT 94	99		
Fusobacterium mortiferum		5.86	0.17	GAL 96	1		
27630102	**Low discrimination**						**27630102**
					GRA		
Actinomyces viscosus		78.85	0.37	MLT 99	99		
Fusobacterium mortiferum		10.72	0.35	TRE 7	1		
Actinomyces naeslundii		10.43	0.46		99		
27630120	**Low discrimination**						**27630120**
					GRA		
Actinomyces naeslundii		87.86	0.46	MLT 94	99		
Fusobacterium mortiferum		8.21	0.17	GAL 96	1		
Clostridium perfringens		3.93	0.01		99		
27630121	**Good identification**						**27630121**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27630122	**Acceptable identification**						**27630122**
Actinomyces naeslundii		92.91	0.46	MLT 94			
Actinomyces viscosus		7.09	0.03	GAL 96			
27630123	**Good identification**						**27630123**
Actinomyces naeslundii		100.0	0.49	MLT 94			
27630141	**Good identification**						**27630141**
Actinomyces naeslundii		100.0	0.36	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27630143	**Good identification**						**27630143**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27630161	**Good identification**						**27630161**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27630163	**Good identification**						**27630163**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27631000	**Low discrimination**						**27631000**
					GRA	CAT	
Actinomyces naeslundii		54.51	0.62	MLT 94	99	25	
Fusobacterium mortiferum		38.43	0.48		1	5	
Propionibacterium granulosum		7.06	0.32		99	90	
27631001	**Very good identification**						**27631001**
					GRA		
Actinomyces naeslundii		99.53	0.65	MLT 94	99		
Fusobacterium mortiferum		0.47	0.13		1		
27631002	**Low discrimination**						**27631002**
					GRA		
Actinomyces naeslundii		70.43	0.62	MLT 94	99		
Actinomyces viscosus		21.77	0.29		99		
Fusobacterium mortiferum		5.52	0.31		1		
27631003	**Good identification**						**27631003**
Actinomyces naeslundii		93.57	0.65	MLT 94			
Actinomyces viscosus		6.43	0.21				
27631010	**Low discrimination**						**27631010**
					GRA	CAT	
Actinomyces naeslundii		67.98	0.40	MLT 94	99	25	
Propionibacterium granulosum		22.82	0.17	MLZ 5	99	90	
Fusobacterium mortiferum		9.20	0.13		1	5	
27631011	**Good identification**						**27631011**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27631012	**Good identification**						**27631012**
Actinomyces naeslundii		100.0	0.40	MLT 94			
27631013	**Good identification**						**27631013**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27631020	**Low discrimination**						**27631020**
					CAT		
Actinomyces naeslundii		90.00	0.62	MLT 94	25		
Clostridium perfringens		6.45	0.20		1		
Propionibacterium granulosum		2.92	0.21		90		
27631021	**Very good identification**						**27631021**
Actinomyces naeslundii		100.0	0.65	MLT 94			
27631022	**Very good identification**						**27631022**
					CAT		
Actinomyces naeslundii		95.99	0.62	MLT 94	25		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		3.24	0.14		1	
Propionibacterium granulosum		0.78	0.11		90	
27631023	**Very good identification**					**27631023**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27631030	**Good identification**					**27631030**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27631031	**Good identification**					**27631031**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27631032	**Good identification**					**27631032**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27631033	**Good identification**					**27631033**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27631040	**Good identification**					**27631040**
					GRA	
Actinomyces naeslundii		96.12	0.49	MLT 94	99	
Fusobacterium mortiferum		3.88	0.13		1	
27631041	**Good identification**					**27631041**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27631042	**Good identification**					**27631042**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27631043	**Good identification**					**27631043**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27631060	**Acceptable identification**					**27631060**
Actinomyces naeslundii		92.82	0.49	MLT 94		
Clostridium perfringens		7.18	0.08			
27631061	**Good identification**					**27631061**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27631062	**Good identification**					**27631062**
Actinomyces naeslundii		96.49	0.49	MLT 94		
Clostridium perfringens		3.51	0.02			
27631063	**Good identification**					**27631063**
Actinomyces naeslundii		100.0	0.52	MLT 94		
27631100	**Low discrimination**					**27631100**
					GRA	
Actinomyces naeslundii		63.06	0.70	MLT 94	99	
Fusobacterium mortiferum		36.45	0.54		1	
Clostridium perfringens		0.36	0.09		99	
27631101	**Very good identification**					**27631101**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.61	0.73	MLT 94	99	
Fusobacterium mortiferum		0.39	0.20		1	
27631102	**Low discrimination**					**27631102**
					GRA	
Actinomyces naeslundii		79.24	0.70	MLT 94	99	
Actinomyces viscosus		15.30	0.33		99	
Fusobacterium mortiferum		5.09	0.38		1	
27631103	**Very good identification**					**27631103**
Actinomyces naeslundii		95.89	0.73	MLT 94		
Actinomyces viscosus		4.11	0.25			
27631110	**Acceptable identification**					**27631110**
					GRA	
Actinomyces naeslundii		90.01	0.48	MLT 94	99	
Fusobacterium mortiferum		9.99	0.20	MLZ 5	1	
27631111	**Good identification**					**27631111**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27631112	**Good identification**					**27631112**
Actinomyces naeslundii		96.43	0.48	MLT 94		
Actinomyces viscosus		3.57	-.01	MLZ 5		
27631113	**Good identification**					**27631113**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27631120	**Low discrimination**					**27631120**
					GRA	
Actinomyces naeslundii		84.01	0.70	MLT 94	99	
Clostridium perfringens		15.50	0.35		99	
Fusobacterium mortiferum		0.49	0.20		1	
27631121	**Very good identification**					**27631121**
Actinomyces naeslundii		99.88	0.73	MLT 94		
Clostridium perfringens		0.12	0.01			
27631122	**Good identification**					**27631122**
					GRA	
Actinomyces naeslundii		90.38	0.70	MLT 94	99	
Clostridium perfringens		7.85	0.29		99	
Parabacteroides distasonis		1.42	0.26		1	
27631123	**Very good identification**					**27631123**
Actinomyces naeslundii		100.0	0.73	MLT 94		
27631130	**Good identification**					**27631130**
Actinomyces naeslundii		96.58	0.48	MLT 94		
Clostridium perfringens		3.42	0.01	MLZ 5		
27631131	**Good identification**					**27631131**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27631132	**Low discrimination**					**27631132**
					GRA	
Actinomyces naeslundii		86.18	0.48	MLT 94	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		13.82	0.21	MLZ 5	1	
27631133	**Good identification**					**27631133**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27631140	**Very good identification**					**27631140**
					GRA	
Actinomyces naeslundii		96.80	0.57	MLT 94	99	
Fusobacterium mortiferum		3.20	0.20		1	
27631141	**Very good identification**					**27631141**
Actinomyces naeslundii		100.0	0.60	MLT 94		
27631142	**Very good identification**					**27631142**
Actinomyces naeslundii		98.91	0.57	MLT 94		
Actinomyces viscosus		1.09	-.01			
27631143	**Very good identification**					**27631143**
Actinomyces naeslundii		100.0	0.60	MLT 94		
27631151	**Good identification**					**27631151**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631153	**Good identification**					**27631153**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631160	**Low discrimination**					**27631160**
Actinomyces naeslundii		83.39	0.57	MLT 94		
Clostridium perfringens		16.61	0.22			
27631161	**Very good identification**					**27631161**
Actinomyces naeslundii		100.0	0.60	MLT 94		
27631162	**Good identification**					**27631162**
Actinomyces naeslundii		91.43	0.57	MLT 94		
Clostridium perfringens		8.57	0.17			
27631163	**Very good identification**					**27631163**
Actinomyces naeslundii		100.0	0.60	MLT 94		
27631171	**Good identification**					**27631171**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631173	**Good identification**					**27631173**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631300	**Low discrimination**					**27631300**
					GRA	
Actinomyces naeslundii		63.37	0.35	MLT 94	99	
Fusobacterium mortiferum		36.63	0.20	RHA 1	1	
27631301	**Good identification**					**27631301**
Actinomyces naeslundii		100.0	0.38	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27631302	**Low discrimination**					**27631302**
Actinomyces naeslundii		83.81	0.35	MLT 94		
Actinomyces viscosus		16.19	-.01	RHA 1		
27631303	**Good identification**					**27631303**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631320	**Low discrimination**					**27631320**
Actinomyces naeslundii		84.43	0.35	MLT 94		
Clostridium perfringens		15.58	0.01	RHA 1		
27631321	**Good identification**					**27631321**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631323	**Good identification**					**27631323**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631500	**Low discrimination**					**27631500**
					GRA	
Actinomyces naeslundii		63.37	0.35	MLT 94	99	
Fusobacterium mortiferum		36.63	0.20	ARA 1	1	
27631501	**Good identification**					**27631501**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631502	**Low discrimination**					**27631502**
Actinomyces naeslundii		83.81	0.35	MLT 94		
Actinomyces viscosus		16.19	-.01	ARA 1		
27631503	**Good identification**					**27631503**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631520	**Low discrimination**					**27631520**
Actinomyces naeslundii		84.43	0.35	MLT 94		
Clostridium perfringens		15.58	0.01	ARA 1		
27631521	**Good identification**					**27631521**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27631522	**Low discrimination**					**27631522**
					GRA	
Actinomyces naeslundii		51.25	0.35	MLT 94	99	
Parabacteroides distasonis		48.75	0.22	ARA 1	1	
27631523	**Good identification**					**27631523**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27632001	**Good identification**					**27632001**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27632002	**Low discrimination**					**27632002**
Actinomyces naeslundii		73.26	0.43	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		22.13	0.19	GAL 96		
Actinomyces viscosus		4.61	-.02			
27632003	**Good identification**					**27632003**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27632020	**Low discrimination**					**27632020**
Actinomyces naeslundii		76.80	0.43	MLT 94		
Propionibacterium granulosum		23.20	0.19	GAL 96		
27632021	**Good identification**					**27632021**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27632022	**Acceptable identification**					**27632022**
Actinomyces naeslundii		92.98	0.43	MLT 94		
Propionibacterium granulosum		7.02	0.09	GAL 96		
27632023	**Good identification**					**27632023**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27632100	**Low discrimination**					**27632100**
					GRA	
Actinomyces naeslundii		89.18	0.51	MLT 94	99	
Propionibacterium propionicum		6.53	0.21	GAL 96	99	
Fusobacterium mortiferum		4.29	0.17		1	
27632101	**Good identification**					**27632101**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27632102	**Low discrimination**					**27632102**
Actinomyces naeslundii		87.02	0.51	MLT 94		
Propionibacterium propionicum		9.56	0.24	GAL 96		
Actinomyces viscosus		3.42	0.03			
27632103	**Good identification**					**27632103**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27632120	**Good identification**					**27632120**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27632121	**Good identification**					**27632121**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27632122	**Good identification**					**27632122**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27632123	**Good identification**					**27632123**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27632140	**Low discrimination**					**27632140**
Actinomyces naeslundii		70.67	0.38	MLT 94		
Propionibacterium propionicum		29.33	0.21	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27632141	**Good identification**						**27632141**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27632142	**Low discrimination**						**27632142**
Actinomyces naeslundii		61.63	0.38	MLT 94			
Propionibacterium propionicum		38.37	0.24	GAL 96			
27632143	**Good identification**						**27632143**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27632160	**Good identification**						**27632160**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27632161	**Good identification**						**27632161**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27632162	**Good identification**						**27632162**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27632163	**Good identification**						**27632163**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27633000	**Good identification**						**27633000**
					GRA	CAT	
Actinomyces naeslundii		94.88	0.67	MLT 94	99	25	
Propionibacterium granulosum		4.78	0.30		99	90	
Fusobacterium mortiferum		0.35	0.13		1	5	
27633001	**Very good identification**						**27633001**
Actinomyces naeslundii		100.0	0.70	MLT 94			
27633002	**Very good identification**						**27633002**
Actinomyces naeslundii		98.76	0.67	MLT 94			
Propionibacterium granulosum		1.24	0.19				
27633003	**Very good identification**						**27633003**
Actinomyces naeslundii		100.0	0.70	MLT 94			
27633010	**Low discrimination**						**27633010**
Actinomyces naeslundii		88.46	0.45	MLT 94			
Propionibacterium granulosum		11.54	0.15	MLZ 5			
27633011	**Good identification**						**27633011**
Actinomyces naeslundii		100.0	0.48	MLT 94			
27633012	**Good identification**						**27633012**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27633013	**Good identification**						**27633013**
Actinomyces naeslundii		100.0	0.48	MLT 94			
27633020	**Very good identification**						**27633020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.76	0.67	MLT 94		
Propionibacterium granulosum		1.24	0.19			
27633021	**Very good identification**					**27633021**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27633022	**Very good identification**					**27633022**
Actinomyces naeslundii		99.69	0.67	MLT 94		
Propionibacterium granulosum		0.31	0.09			
27633023	**Very good identification**					**27633023**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27633030	**Good identification**					**27633030**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27633031	**Good identification**					**27633031**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27633032	**Good identification**					**27633032**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27633033	**Good identification**					**27633033**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27633040	**Good identification**					**27633040**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27633041	**Very good identification**					**27633041**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27633042	**Good identification**					**27633042**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27633043	**Very good identification**					**27633043**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27633060	**Good identification**					**27633060**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27633061	**Very good identification**					**27633061**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27633062	**Good identification**					**27633062**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27633063	**Very good identification**					**27633063**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27633100	**Very good identification**					**27633100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.27	0.74	MLT 94	99	
Propionibacterium propionicum		0.30	0.21		99	
Fusobacterium mortiferum		0.30	0.20		1	
27633101	**Very good identification**					**27633101**
Actinomyces naeslundii		100.0	0.78	MLT 94		
27633102	**Very good identification**					**27633102**
Actinomyces naeslundii		99.26	0.74	MLT 94		
Propionibacterium propionicum		0.45	0.24			
Actinomyces israelii		0.19	0.19			
27633103	**Very good identification**					**27633103**
Actinomyces naeslundii		100.0	0.78	MLT 94		
27633110	**Good identification**					**27633110**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27633111	**Good identification**					**27633111**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27633112	**Good identification**					**27633112**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27633113	**Good identification**					**27633113**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27633120	**Very good identification**					**27633120**
Actinomyces naeslundii		99.90	0.74	MLT 94		
Clostridium perfringens		0.10	0.01			
27633121	**Very good identification**					**27633121**
Actinomyces naeslundii		100.0	0.78	MLT 94		
27633122	**Very good identification**					**27633122**
Actinomyces naeslundii		100.0	0.74	MLT 94		
27633123	**Very good identification**					**27633123**
Actinomyces naeslundii		100.0	0.78	MLT 94		
27633130	**Good identification**					**27633130**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27633131	**Good identification**					**27633131**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27633132	**Good identification**					**27633132**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27633133	**Good identification**					**27633133**
Actinomyces naeslundii		100.0	0.56	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27633140	**Very good identification**					**27633140**
Actinomyces naeslundii		98.30	0.62	MLT 94		
Propionibacterium propionicum		1.70	0.21			
27633141	**Very good identification**					**27633141**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27633142	**Very good identification**					**27633142**
Actinomyces naeslundii		97.47	0.62	MLT 94		
Propionibacterium propionicum		2.53	0.24			
27633143	**Very good identification**					**27633143**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27633150	**Good identification**					**27633150**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633151	**Good identification**					**27633151**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633152	**Good identification**					**27633152**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633153	**Good identification**					**27633153**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633160	**Very good identification**					**27633160**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27633161	**Very good identification**					**27633161**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27633162	**Very good identification**					**27633162**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27633163	**Very good identification**					**27633163**
Actinomyces naeslundii		100.0	0.65	MLT 94		
27633170	**Good identification**					**27633170**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633171	**Good identification**					**27633171**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633172	**Good identification**					**27633172**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633173	**Good identification**					**27633173**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633201	**Good identification**					**27633201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633203	**Good identification**					**27633203**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633221	**Good identification**					**27633221**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633223	**Good identification**					**27633223**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633300	**Good identification**					**27633300**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633301	**Good identification**					**27633301**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633302	**Good identification**					**27633302**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633303	**Good identification**					**27633303**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633320	**Good identification**					**27633320**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633321	**Good identification**					**27633321**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633322	**Good identification**					**27633322**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633323	**Good identification**					**27633323**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633401	**Good identification**					**27633401**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633403	**Good identification**					**27633403**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633421	**Good identification**					**27633421**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633423	**Good identification**					**27633423**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27633500	**Acceptable identification**					**27633500**
Actinomyces naeslundii		90.76	0.40	MLT 94		
Actinomyces israelii		9.24	0.14	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27633501	**Good identification**					**27633501**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633502	**Low discrimination**					**27633502**
Actinomyces naeslundii		86.75	0.40	MLT 94		
Actinomyces israelii		13.25	0.17	ARA 1		
27633503	**Good identification**					**27633503**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633520	**Good identification**					**27633520**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633521	**Good identification**					**27633521**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27633522	**Good identification**					**27633522**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27633523	**Good identification**					**27633523**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27635101	**Good identification**					**27635101**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27635102	**Low discrimination**					**27635102**
					GRA	
Actinomyces naeslundii		41.55	0.35	MLT 94	99	
Actinomyces israelii		30.54	0.24	XYL 1	99	
Fusobacterium mortiferum		19.89	0.18		1	
27635103	**Good identification**					**27635103**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27635120	**Low discrimination**					**27635120**
Actinomyces naeslundii		84.43	0.35	MLT 94		
Clostridium perfringens		15.58	0.01	XYL 1		
27635121	**Good identification**					**27635121**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27635122	**Acceptable identification**					**27635122**
					GRA	
Parabacteroides distasonis		93.32	0.42	NIT 1	1	
Actinomyces naeslundii		6.68	0.35		99	
27635123	**Good identification**					**27635123**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27635132	**Good identification**					**27635132**
Parabacteroides distasonis		100.0	0.37	NIT 1		
27635322	**Good identification**					**27635322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		100.0	0.50	NIT 1		
27635332	**Good identification**					**27635332**
Parabacteroides distasonis		100.0	0.46	NIT 1		
27635522	**Good identification**					**27635522**
Parabacteroides distasonis		100.0	0.38	NIT 1		
27635722	**Good identification**					**27635722**
Parabacteroides distasonis		100.0	0.47	NIT 1		
27635732	**Good identification**					**27635732**
Parabacteroides distasonis		95.39	0.42	NIT 1		
Clostridium clostridioforme		4.61	0.14			
27637001	**Good identification**					**27637001**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27637003	**Good identification**					**27637003**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27637021	**Good identification**					**27637021**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27637023	**Good identification**					**27637023**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27637101	**Good identification**					**27637101**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27637102	**Low discrimination**					**27637102**
Actinomyces israelii		68.21	0.37	MLT 99		
Actinomyces naeslundii		31.79	0.40	CEL 93		
27637103	**Good identification**					**27637103**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27637120	**Acceptable identification**					**27637120**
Actinomyces naeslundii		91.63	0.40	MLT 94		
Actinomyces israelii		8.37	0.13	XYL 1		
27637121	**Good identification**					**27637121**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27637122	**Low discrimination**					**27637122**
Actinomyces naeslundii		87.95	0.40	MLT 94		
Actinomyces israelii		12.05	0.16	XYL 1		
27637123	**Good identification**					**27637123**
Actinomyces naeslundii		100.0	0.43	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27637502	**Acceptable identification**					**27637502**
Actinomyces israelii		92.97	0.35	MLT 99		
Bifidobacterium adolescentis		7.03	0.10	CEL 93		
27641000	**Low discrimination**					**27641000**
					GRA	
Actinomyces naeslundii		53.04	0.37	MLT 94	99	
Eubacterium tortuosum		35.23	0.06	FRU 95	99	
Fusobacterium mortiferum		11.73	0.14		1	
27641001	**Good identification**					**27641001**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27641003	**Low discrimination**					**27641003**
Actinomyces naeslundii		73.79	0.40	MLT 94		
Actinomyces viscosus		26.21	0.08	FRU 95		
27641020	**Good identification**					**27641020**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27641021	**Good identification**					**27641021**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27641022	**Good identification**					**27641022**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27641023	**Good identification**					**27641023**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27641100	**Low discrimination**					**27641100**
					GRA	
Actinomyces naeslundii		84.65	0.45	MLT 94	99	
Fusobacterium mortiferum		15.35	0.21	FRU 95	1	
27641101	**Good identification**					**27641101**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27641102	**Low discrimination**					**27641102**
Actinomyces naeslundii		50.04	0.45	MLT 94		
Actinomyces viscosus		49.96	0.21	FRU 95		
27641103	**Low discrimination**					**27641103**
Actinomyces naeslundii		81.84	0.48	MLT 94		
Actinomyces viscosus		18.16	0.12	FRU 95		
27641120	**Good identification**					**27641120**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27641121	**Good identification**					**27641121**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27641122	**Good identification**					**27641122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	MLT 94		
27641123	**Good identification**					**27641123**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27643000	**Low discrimination**					**27643000**
					SPO	
Actinomyces naeslundii		74.65	0.42	MLT 94	1	
Clostridium innocuum		25.35	0.10	FRU 95	99	
27643001	**Good identification**					**27643001**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27643002	**Low discrimination**					**27643002**
					SPO	
Actinomyces naeslundii		74.65	0.42	MLT 94	1	
Clostridium innocuum		25.35	0.10	FRU 95	99	
27643003	**Good identification**					**27643003**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27643020	**Good identification**					**27643020**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27643021	**Good identification**					**27643021**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27643022	**Good identification**					**27643022**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27643023	**Good identification**					**27643023**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27643100	**Good identification**					**27643100**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27643101	**Good identification**					**27643101**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27643102	**Good identification**					**27643102**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27643103	**Good identification**					**27643103**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27643120	**Good identification**					**27643120**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27643121	**Good identification**					**27643121**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27643122	**Good identification**					**27643122**
Actinomyces naeslundii		100.0	0.50	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27643123	**Good identification**					**27643123**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27643140	**Good identification**					**27643140**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27643141	**Good identification**					**27643141**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27643142	**Good identification**					**27643142**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27643143	**Good identification**					**27643143**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27643160	**Good identification**					**27643160**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27643161	**Good identification**					**27643161**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27643162	**Good identification**					**27643162**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27643163	**Good identification**					**27643163**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27650000	**Acceptable identification**					**27650000**
					GRA	
Fusobacterium mortiferum		91.87	0.45	NIT 1	1	
Actinomyces naeslundii		4.98	0.35		99	
Actinomyces viscosus		3.15	0.08		99	
27650001	**Low discrimination**					**27650001**
					GRA	
Actinomyces naeslundii		79.07	0.38	MLT 94	99	
Actinomyces viscosus		11.11	-.01	GAL 96	99	
Fusobacterium mortiferum		9.82	0.11		1	
27650002	**Good identification**					**27650002**
					GRA	
Actinomyces viscosus		95.36	0.42	MLT 99	99	
Fusobacterium mortiferum		3.12	0.29		1	
Actinomyces naeslundii		1.52	0.35		99	
27650020	**Low discrimination**					**27650020**
					GRA	
Actinomyces naeslundii		84.30	0.35	MLT 94	99	
Fusobacterium mortiferum		15.70	0.11	GAL 96	1	
27650021	**Good identification**					**27650021**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27650022	**Low discrimination**					**27650022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		61.26	0.35	MLT 94		
Actinomyces viscosus		38.74	0.08	GAL 96		
27650023	**Low discrimination**					**27650023**
Actinomyces naeslundii		87.68	0.38	MLT 94		
Actinomyces viscosus		12.32	-.01	GAL 96		
27650100	**Acceptable identification**					**27650100**
					GRA	
Fusobacterium mortiferum		91.55	0.52	NIT 1	1	
Actinomyces naeslundii		6.06	0.43		99	
Actinomyces viscosus		2.39	0.12		99	
27650101	**Low discrimination**					**27650101**
					GRA	
Actinomyces naeslundii		84.07	0.46	MLT 94	99	
Fusobacterium mortiferum		8.56	0.18	GAL 96	1	
Actinomyces viscosus		7.38	0.04		99	
27650102	**Acceptable identification**					**27650102**
					GRA	
Actinomyces viscosus		93.59	0.46	MLT 99	99	
Fusobacterium mortiferum		4.02	0.35		1	
Actinomyces naeslundii		2.39	0.43		99	
27650103	**Low discrimination**					**27650103**
Actinomyces viscosus		89.68	0.38	MLT 99		
Actinomyces naeslundii		10.32	0.46			
27650120	**Low discrimination**					**27650120**
					GRA	
Actinomyces naeslundii		86.76	0.43	MLT 94	99	
Fusobacterium mortiferum		13.24	0.18	GAL 96	1	
27650121	**Good identification**					**27650121**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27650122	**Low discrimination**					**27650122**
Actinomyces naeslundii		71.69	0.43	MLT 94		
Actinomyces viscosus		28.31	0.12	GAL 96		
27650123	**Acceptable identification**					**27650123**
Actinomyces naeslundii		91.93	0.46	MLT 94		
Actinomyces viscosus		8.07	0.04	GAL 96		
27651000	**Low discrimination**					**27651000**
					GRA	
Fusobacterium mortiferum		52.12	0.48	NIT 1	1	
Actinomyces naeslundii		45.23	0.59		99	
Lactobacillus catenaformis		1.56	0.12		99	
27651001	**Very good identification**					**27651001**
					GRA	
Actinomyces naeslundii		99.23	0.62	MLT 94	99	
Fusobacterium mortiferum		0.77	0.14		1	
27651002	**Low discrimination**					**27651002**
					GRA	
Actinomyces viscosus		57.45	0.38	MLT 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		35.94	0.59		99	
Fusobacterium mortiferum		4.60	0.32		1	
27651003	**Low discrimination**					**27651003**
Actinomyces naeslundii		73.79	0.62	MLT 94		
Actinomyces viscosus		26.21	0.30			
27651010	**Low discrimination**					**27651010**
					GRA	
Actinomyces naeslundii		81.89	0.37	MLT 94	99	
Fusobacterium mortiferum		18.11	0.14	MLZ 5	1	
27651011	**Good identification**					**27651011**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27651012	**Low discrimination**					**27651012**
Actinomyces naeslundii		76.52	0.37	MLT 94		
Actinomyces viscosus		23.48	0.04	MLZ 5		
27651013	**Good identification**					**27651013**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27651020	**Very good identification**					**27651020**
					GRA	
Actinomyces naeslundii		98.02	0.59	MLT 94	99	
Fusobacterium mortiferum		1.14	0.14		1	
Lactobacillus catenaformis		0.84	0.02		99	
27651021	**Very good identification**					**27651021**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27651022	**Good identification**					**27651022**
Actinomyces naeslundii		93.89	0.59	MLT 94		
Lactobacillus acidophilus		2.02	0.06			
Actinomyces viscosus		1.52	0.04			
27651023	**Very good identification**					**27651023**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27651030	**Good identification**					**27651030**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27651031	**Good identification**					**27651031**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27651032	**Good identification**					**27651032**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27651033	**Good identification**					**27651033**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27651040	**Acceptable identification**					**27651040**
					GRA	
Actinomyces naeslundii		93.81	0.46	MLT 94	99	
Fusobacterium mortiferum		6.19	0.14		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27651041	**Good identification**					**27651041**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27651042	**Acceptable identification**					**27651042**
Actinomyces naeslundii		91.62	0.46	MLT 94		
Actinomyces viscosus		8.38	0.04			
27651043	**Good identification**					**27651043**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27651060	**Good identification**					**27651060**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27651061	**Good identification**					**27651061**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27651062	**Good identification**					**27651062**
Actinomyces naeslundii		100.0	0.46	MLT 94		
27651063	**Good identification**					**27651063**
Actinomyces naeslundii		100.0	0.49	MLT 94		
27651100	**Low discrimination**					**27651100**
					GRA	
Actinomyces naeslundii		50.69	0.67	MLT 94	99	
Fusobacterium mortiferum		47.88	0.55		1	
Lactobacillus catenaformis		0.92	0.15		99	
27651101	**Very good identification**					**27651101**
					GRA	
Actinomyces naeslundii		99.15	0.70	MLT 94	99	
Fusobacterium mortiferum		0.63	0.21		1	
Actinomyces viscosus		0.22	0.00		99	
27651102	**Low discrimination**					**27651102**
					GRA	
Actinomyces naeslundii		45.09	0.67	MLT 94	99	
Actinomyces viscosus		45.02	0.43		99	
Fusobacterium mortiferum		4.73	0.38		1	
27651103	**Low discrimination**					**27651103**
Actinomyces naeslundii		81.84	0.70	MLT 94		
Actinomyces viscosus		18.16	0.34			
27651110	**Low discrimination**					**27651110**
					GRA	
Actinomyces naeslundii		84.65	0.45	MLT 94	99	
Fusobacterium mortiferum		15.35	0.21	MLZ 5	1	
27651111	**Good identification**					**27651111**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27651112	**Low discrimination**					**27651112**
Actinomyces naeslundii		83.92	0.45	MLT 94		
Actinomyces viscosus		16.08	0.08	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27651113	**Good identification**					**27651113**
Actinomyces naeslundii		95.92	0.48	MLT 94		
Actinomyces viscosus		4.09	0.00	MLZ 5		
27651120	**Very good identification**					**27651120**
					GRA	
Actinomyces naeslundii		97.45	0.67	MLT 94	99	
Fusobacterium mortiferum		0.93	0.21		1	
Clostridium perfringens		0.59	0.06		99	
27651121	**Very good identification**					**27651121**
Actinomyces naeslundii		100.0	0.70	MLT 94		
27651122	**Low discrimination**					**27651122**
					GRA	
Actinomyces naeslundii		85.63	0.67	MLT 94	99	
Lactobacillus acidophilus		7.45	0.24		99	
Prevotella oralis		4.33	0.19		1	
27651123	**Very good identification**					**27651123**
Actinomyces naeslundii		99.78	0.70	MLT 94		
Actinomyces viscosus		0.22	0.00			
27651130	**Good identification**					**27651130**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27651131	**Good identification**					**27651131**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27651132	**Good identification**					**27651132**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27651133	**Good identification**					**27651133**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27651140	**Acceptable identification**					**27651140**
					GRA	
Actinomyces naeslundii		94.87	0.54	MLT 94	99	
Fusobacterium mortiferum		5.13	0.21		1	
27651141	**Very good identification**					**27651141**
Actinomyces naeslundii		100.0	0.57	MLT 94		
27651142	**Low discrimination**					**27651142**
Actinomyces naeslundii		89.44	0.54	MLT 94		
Lactobacillus acidophilus		5.45	0.09			
Actinomyces viscosus		5.11	0.08			
27651143	**Very good identification**					**27651143**
Actinomyces naeslundii		98.75	0.57	MLT 94		
Actinomyces viscosus		1.25	0.00			
27651160	**Good identification**					**27651160**
Actinomyces naeslundii		100.0	0.54	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27651161	**Very good identification**						**27651161**
Actinomyces naeslundii		100.0	0.57	MLT 94			
27651162	**Acceptable identification**						**27651162**
Actinomyces naeslundii		94.25	0.54	MLT 94			
Lactobacillus acidophilus		5.75	0.09				
27651163	**Very good identification**						**27651163**
Actinomyces naeslundii		100.0	0.57	MLT 94			
27651301	**Good identification**						**27651301**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27651303	**Low discrimination**						**27651303**
Actinomyces naeslundii		81.84	0.36	MLT 94			
Actinomyces viscosus		18.16	0.00	RHA 1			
27651321	**Good identification**						**27651321**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27651323	**Good identification**						**27651323**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27651501	**Good identification**						**27651501**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27651503	**Low discrimination**						**27651503**
Actinomyces naeslundii		81.84	0.36	MLT 94			
Actinomyces viscosus		18.16	0.00	ARA 1			
27651521	**Good identification**						**27651521**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27651523	**Good identification**						**27651523**
Actinomyces naeslundii		100.0	0.36	MLT 94			
27652000	**Low discrimination**						**27652000**
					GRA	SPO	
Actinomyces naeslundii		65.58	0.40	MLT 94	99	1	
Clostridium innocuum		28.13	0.10	GAL 96	99	99	
Fusobacterium mortiferum		6.29	0.11		1	1	
27652001	**Good identification**						**27652001**
Actinomyces naeslundii		100.0	0.43	MLT 94			
27652002	**Low discrimination**						**27652002**
					SPO		
Actinomyces naeslundii		56.99	0.40	MLT 94	1		
Clostridium innocuum		24.44	0.10	GAL 96	99		
Actinomyces viscosus		18.56	0.08		1		
27652003	**Acceptable identification**						**27652003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		93.25	0.43	MLT 94		
Actinomyces viscosus		6.75	-.01	GAL 96		
27652020	**Good identification**					**27652020**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27652021	**Good identification**					**27652021**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27652022	**Good identification**					**27652022**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27652023	**Good identification**					**27652023**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27652100	**Acceptable identification**					**27652100**
					GRA	
Actinomyces naeslundii		92.71	0.48	MLT 94	99	
Fusobacterium mortiferum		7.29	0.18	GAL 96	1	
27652101	**Good identification**					**27652101**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27652102	**Low discrimination**					**27652102**
Actinomyces naeslundii		83.09	0.48	MLT 94		
Actinomyces viscosus		16.91	0.12	GAL 96		
27652103	**Good identification**					**27652103**
Actinomyces naeslundii		95.67	0.51	MLT 94		
Actinomyces viscosus		4.33	0.04	GAL 96		
27652120	**Good identification**					**27652120**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27652121	**Good identification**					**27652121**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27652122	**Good identification**					**27652122**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27652123	**Good identification**					**27652123**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27652140	**Good identification**					**27652140**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27652141	**Good identification**					**27652141**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27652142	**Good identification**					**27652142**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27652143	**Good identification**					**27652143**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	MLT 94			
27652160	**Good identification**						**27652160**
Actinomyces naeslundii		100.0	0.35	MLT 94			
27652161	**Good identification**						**27652161**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27652162	**Good identification**						**27652162**
Actinomyces naeslundii		100.0	0.35	MLT 94			
27652163	**Good identification**						**27652163**
Actinomyces naeslundii		100.0	0.38	MLT 94			
27653000	**Low discrimination**						**27653000**
					GRA	SPO	
Actinomyces naeslundii		74.07	0.64	MLT 94	99	1	
Clostridium innocuum		25.15	0.32		99	99	
Fusobacterium mortiferum		0.44	0.14		1	1	
27653001	**Very good identification**						**27653001**
Actinomyces naeslundii		100.0	0.67	MLT 94			
27653002	**Low discrimination**						**27653002**
					SPO		
Actinomyces naeslundii		73.81	0.64	MLT 94	1		
Clostridium innocuum		25.06	0.32		99		
Actinomyces viscosus		0.61	0.04		1		
27653003	**Very good identification**						**27653003**
Actinomyces naeslundii		100.0	0.67	MLT 94			
27653010	**Low discrimination**						**27653010**
					SPO		
Actinomyces naeslundii		78.82	0.42	MLT 94	1		
Clostridium innocuum		21.18	0.08	MLZ 5	99		
27653011	**Good identification**						**27653011**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27653012	**Low discrimination**						**27653012**
					SPO		
Actinomyces naeslundii		78.82	0.42	MLT 94	1		
Clostridium innocuum		21.18	0.08	MLZ 5	99		
27653013	**Good identification**						**27653013**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27653020	**Very good identification**						**27653020**
Actinomyces naeslundii		100.0	0.64	MLT 94			
27653021	**Very good identification**						**27653021**
Actinomyces naeslundii		100.0	0.67	MLT 94			
27653022	**Very good identification**						**27653022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.64	MLT 94		
27653023	**Very good identification**					**27653023**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27653030	**Good identification**					**27653030**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27653031	**Good identification**					**27653031**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27653032	**Good identification**					**27653032**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27653033	**Good identification**					**27653033**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27653040	**Good identification**					**27653040**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27653041	**Good identification**					**27653041**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27653042	**Good identification**					**27653042**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27653043	**Good identification**					**27653043**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27653060	**Good identification**					**27653060**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27653061	**Good identification**					**27653061**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27653062	**Good identification**					**27653062**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27653063	**Good identification**					**27653063**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27653100	**Very good identification**					**27653100**
					GRA	
Actinomyces naeslundii		98.87	0.72	MLT 94	99	
Actinomyces israelii		0.64	0.25		99	
Fusobacterium mortiferum		0.49	0.21		1	
27653101	**Very good identification**					**27653101**
Actinomyces naeslundii		100.0	0.75	MLT 94		
27653102	**Very good identification**					**27653102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.30	0.72	MLT 94		
Actinomyces israelii		0.96	0.28			
Actinomyces viscosus		0.51	0.08			
27653103	**Very good identification**					**27653103**
Actinomyces naeslundii		99.89	0.75	MLT 94		
Actinomyces viscosus		0.12	0.00			
27653110	**Good identification**					**27653110**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27653111	**Good identification**					**27653111**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27653112	**Good identification**					**27653112**
Actinomyces naeslundii		97.31	0.50	MLT 94		
Actinomyces israelii		2.69	0.14	MLZ 5		
27653113	**Good identification**					**27653113**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27653120	**Very good identification**					**27653120**
Actinomyces naeslundii		100.0	0.72	MLT 94		
27653121	**Very good identification**					**27653121**
Actinomyces naeslundii		100.0	0.75	MLT 94		
27653122	**Very good identification**					**27653122**
					SPO	
Actinomyces naeslundii		99.41	0.72	MLT 94	1	
Clostridium ramosum		0.36	0.05		92	
Lactobacillus acidophilus		0.24	0.02		1	
27653123	**Very good identification**					**27653123**
Actinomyces naeslundii		100.0	0.75	MLT 94		
27653130	**Good identification**					**27653130**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27653131	**Good identification**					**27653131**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27653132	**Good identification**					**27653132**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27653133	**Good identification**					**27653133**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27653140	**Very good identification**					**27653140**
Actinomyces naeslundii		98.59	0.59	MLT 94		
Actinomyces israelii		1.41	0.18			
27653141	**Very good identification**					**27653141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.62	MLT 94		
27653142	**Very good identification**					**27653142**
Actinomyces naeslundii		97.90	0.59	MLT 94		
Actinomyces israelii		2.10	0.21			
27653143	**Very good identification**					**27653143**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27653150	**Good identification**					**27653150**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653151	**Good identification**					**27653151**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27653152	**Good identification**					**27653152**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653153	**Good identification**					**27653153**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27653160	**Very good identification**					**27653160**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27653161	**Very good identification**					**27653161**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27653162	**Very good identification**					**27653162**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27653163	**Very good identification**					**27653163**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27653170	**Good identification**					**27653170**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653171	**Good identification**					**27653171**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27653172	**Good identification**					**27653172**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653173	**Good identification**					**27653173**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27653300	**Good identification**					**27653300**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653301	**Good identification**					**27653301**
Actinomyces naeslundii		100.0	0.41	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27653302	**Good identification**					**27653302**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653303	**Good identification**					**27653303**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27653320	**Good identification**					**27653320**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653321	**Good identification**					**27653321**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27653322	**Low discrimination**					**27653322**
					SPO	
Actinomyces naeslundii		67.79	0.37	MLT 94	1	
Clostridium ramosum		32.21	0.07	RHA 1	92	
27653323	**Good identification**					**27653323**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27653500	**Low discrimination**					**27653500**
Actinomyces naeslundii		65.52	0.37	MLT 94		
Actinomyces israelii		34.49	0.24	ARA 1		
27653501	**Good identification**					**27653501**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27653502	**Low discrimination**					**27653502**
Actinomyces naeslundii		55.88	0.37	MLT 94		
Actinomyces israelii		44.12	0.27	ARA 1		
27653503	**Good identification**					**27653503**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27653520	**Good identification**					**27653520**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653521	**Good identification**					**27653521**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27653522	**Good identification**					**27653522**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27653523	**Good identification**					**27653523**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27655100	**Low discrimination**					**27655100**
					GRA	
Fusobacterium mortiferum		66.57	0.35	NIT 1	1	
Actinomyces israelii		23.97	0.30	XYL 7	99	
Actinomyces naeslundii		9.46	0.33		99	
27655101	**Good identification**					**27655101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	MLT 94		
27655103	**Low discrimination**					**27655103**
Actinomyces naeslundii		81.84	0.36	MLT 94		
Actinomyces viscosus		18.16	0.00	XYL 1		
27655121	**Good identification**					**27655121**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27655123	**Good identification**					**27655123**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27657002	**Low discrimination**					**27657002**
					SPO	
Actinomyces israelii		74.04	0.36	MLT 99	1	
Clostridium innocuum		16.59	0.10		99	
Actinomyces naeslundii		9.38	0.30		1	
27657100	**Low discrimination**					**27657100**
Actinomyces israelii		88.09	0.43	MLT 99		
Actinomyces naeslundii		11.91	0.37			
27657101	**Good identification**					**27657101**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27657102	**Acceptable identification**					**27657102**
Actinomyces israelii		91.73	0.46	MLT 99		
Actinomyces naeslundii		8.27	0.37			
27657103	**Good identification**					**27657103**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27657120	**Low discrimination**					**27657120**
Actinomyces naeslundii		67.92	0.37	MLT 94		
Actinomyces israelii		32.08	0.23	XYL 1		
27657121	**Good identification**					**27657121**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27657122	**Low discrimination**					**27657122**
Actinomyces naeslundii		58.54	0.37	MLT 94		
Actinomyces israelii		41.46	0.26	XYL 1		
27657123	**Good identification**					**27657123**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27657140	**Acceptable identification**					**27657140**
Actinomyces israelii		94.22	0.36	MLT 99		
Actinomyces naeslundii		5.78	0.25			
27657142	**Good identification**					**27657142**
Actinomyces israelii		96.07	0.39	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		3.93	0.25			
27657500	**Good identification**					**27657500**
Actinomyces israelii		100.0	0.42	MLT 99		
27657502	**Acceptable identification**					**27657502**
Actinomyces israelii		93.61	0.45	MLT 99		
Bifidobacterium adolescentis		6.39	0.18			
27657542	**Good identification**					**27657542**
Actinomyces israelii		100.0	0.38	MLT 99		
27661000	**Good identification**					**27661000**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27661001	**Good identification**					**27661001**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27661002	**Good identification**					**27661002**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27661003	**Good identification**					**27661003**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27661020	**Good identification**					**27661020**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27661021	**Good identification**					**27661021**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27661022	**Good identification**					**27661022**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27661023	**Good identification**					**27661023**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27661100	**Good identification**					**27661100**
					GRA	
Actinomyces naeslundii		97.29	0.50	MLT 94	99	
Fusobacterium mortiferum		2.71	0.12	FRU 95	1	
27661101	**Good identification**					**27661101**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27661102	**Acceptable identification**					**27661102**
Actinomyces naeslundii		93.68	0.50	MLT 94		
Actinomyces viscosus		3.63	0.01	FRU 95		
Lactobacillus acidophilus		2.70	-.01			
27661103	**Good identification**					**27661103**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27661120	**Good identification**					**27661120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	MLT 94		
27661121	**Good identification**					**27661121**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27661122	**Good identification**					**27661122**
Actinomyces naeslundii		97.20	0.50	MLT 94		
Lactobacillus acidophilus		2.80	-.01	FRU 95		
27661123	**Good identification**					**27661123**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27661140	**Good identification**					**27661140**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27661141	**Good identification**					**27661141**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27661142	**Good identification**					**27661142**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27661143	**Good identification**					**27661143**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27661160	**Good identification**					**27661160**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27661161	**Good identification**					**27661161**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27661162	**Good identification**					**27661162**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27661163	**Good identification**					**27661163**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27663000	**Acceptable identification**					**27663000**
					SPO	
Actinomyces naeslundii		94.21	0.47	MLT 94	1	
Clostridium innocuum		5.79	0.02	FRU 95	99	
27663001	**Good identification**					**27663001**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27663002	**Acceptable identification**					**27663002**
					SPO	
Actinomyces naeslundii		94.21	0.47	MLT 94	1	
Clostridium innocuum		5.79	0.02	FRU 95	99	
27663003	**Good identification**					**27663003**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27663020	**Good identification**					**27663020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	MLT 94		
27663021	**Good identification**					**27663021**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27663022	**Good identification**					**27663022**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27663023	**Good identification**					**27663023**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27663041	**Good identification**					**27663041**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27663043	**Good identification**					**27663043**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27663061	**Good identification**					**27663061**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27663063	**Good identification**					**27663063**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27663100	**Good identification**					**27663100**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27663101	**Very good identification**					**27663101**
Actinomyces naeslundii		100.0	0.58	MLT 94		
27663102	**Good identification**					**27663102**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27663103	**Very good identification**					**27663103**
Actinomyces naeslundii		100.0	0.58	MLT 94		
27663111	**Good identification**					**27663111**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27663113	**Good identification**					**27663113**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27663120	**Good identification**					**27663120**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27663121	**Very good identification**					**27663121**
Actinomyces naeslundii		100.0	0.58	MLT 94		
27663122	**Good identification**					**27663122**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27663123	**Very good identification**					**27663123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.58	MLT 94		
27663131	**Good identification**					**27663131**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27663133	**Good identification**					**27663133**
Actinomyces naeslundii		100.0	0.36	MLT 94		
27663140	**Good identification**					**27663140**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27663141	**Good identification**					**27663141**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27663142	**Good identification**					**27663142**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27663143	**Good identification**					**27663143**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27663160	**Good identification**					**27663160**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27663161	**Good identification**					**27663161**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27663162	**Good identification**					**27663162**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27663163	**Good identification**					**27663163**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27670000	**Low discrimination**					**27670000**
					GRA	
Fusobacterium mortiferum		73.94	0.36	NIT 1	1	
Actinomyces naeslundii		26.06	0.40		99	
27670001	**Good identification**					**27670001**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27670003	**Low discrimination**					**27670003**
Actinomyces naeslundii		64.96	0.43	MLT 94		
Actinomyces viscosus		35.04	0.15	GAL 96		
27670020	**Good identification**					**27670020**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27670021	**Good identification**					**27670021**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27670022	**Good identification**					**27670022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	MLT 94		
27670023	**Good identification**					**27670023**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27670100	**Low discrimination**					**27670100**
					GRA	
Fusobacterium mortiferum		69.93	0.43	NIT 1	1	
Actinomyces naeslundii		30.07	0.48		99	
27670101	**Good identification**					**27670101**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27670103	**Low discrimination**					**27670103**
Actinomyces naeslundii		74.80	0.51	MLT 94		
Actinomyces viscosus		25.20	0.19	GAL 96		
27670120	**Good identification**					**27670120**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27670121	**Good identification**					**27670121**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27670122	**Good identification**					**27670122**
Actinomyces naeslundii		96.49	0.48	MLT 94		
Lactobacillus acidophilus		3.51	-.01	GAL 96		
27670123	**Good identification**					**27670123**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27670140	**Good identification**					**27670140**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27670141	**Good identification**					**27670141**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27670142	**Good identification**					**27670142**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27670143	**Good identification**					**27670143**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27670160	**Good identification**					**27670160**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27670161	**Good identification**					**27670161**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27670162	**Good identification**					**27670162**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27670163	**Good identification**					**27670163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	MLT 94		
27671000	**Low discrimination**					**27671000**
					GRA	
Actinomyces naeslundii		84.04	0.64	MLT 94	99	
Fusobacterium mortiferum		14.90	0.39		1	
Lactobacillus jensenii		1.06	0.10		99	
27671001	**Very good identification**					**27671001**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27671002	**Low discrimination**					**27671002**
Actinomyces naeslundii		88.35	0.64	MLT 94		
Actinomyces viscosus		5.48	0.19			
Lactobacillus acidophilus		3.28	0.15			
27671003	**Very good identification**					**27671003**
Actinomyces naeslundii		98.64	0.67	MLT 94		
Actinomyces viscosus		1.36	0.11			
27671010	**Acceptable identification**					**27671010**
Actinomyces naeslundii		92.59	0.42	MLT 94		
Lactobacillus jensenii		7.41	0.02	MLZ 5		
27671011	**Good identification**					**27671011**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27671012	**Good identification**					**27671012**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27671013	**Good identification**					**27671013**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27671020	**Very good identification**					**27671020**
Actinomyces naeslundii		100.0	0.64	MLT 94		
27671021	**Very good identification**					**27671021**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27671022	**Very good identification**					**27671022**
Actinomyces naeslundii		96.42	0.64	MLT 94		
Lactobacillus acidophilus		3.58	0.15			
27671023	**Very good identification**					**27671023**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27671030	**Good identification**					**27671030**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27671031	**Good identification**					**27671031**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27671032	**Good identification**					**27671032**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	MLT 94		
27671033	**Good identification**					**27671033**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27671040	**Good identification**					**27671040**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27671041	**Good identification**					**27671041**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27671042	**Good identification**					**27671042**
Actinomyces naeslundii		97.46	0.51	MLT 94		
Lactobacillus acidophilus		2.54	-.01			
27671043	**Good identification**					**27671043**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27671060	**Good identification**					**27671060**
Actinomyces naeslundii		100.0	0.51	MLT 94		
27671061	**Good identification**					**27671061**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27671062	**Good identification**					**27671062**
Actinomyces naeslundii		97.46	0.51	MLT 94		
Lactobacillus acidophilus		2.54	-.01			
27671063	**Good identification**					**27671063**
Actinomyces naeslundii		100.0	0.54	MLT 94		
27671100	**Low discrimination**					**27671100**
					GRA	
Actinomyces naeslundii		86.87	0.72	MLT 94	99	
Fusobacterium mortiferum		12.63	0.46		1	
Actinomyces israelii		0.38	0.22		99	
27671101	**Very good identification**					**27671101**
					GRA	
Actinomyces naeslundii		99.90	0.75	MLT 94	99	
Fusobacterium mortiferum		0.10	0.12		1	
27671102	**Low discrimination**					**27671102**
Actinomyces naeslundii		82.55	0.72	MLT 94		
Lactobacillus acidophilus		12.39	0.33			
Actinomyces viscosus		3.20	0.23			
27671103	**Very good identification**					**27671103**
Actinomyces naeslundii		99.05	0.75	MLT 94		
Actinomyces viscosus		0.85	0.15			
Lactobacillus acidophilus		0.10	-.01			
27671110	**Good identification**					**27671110**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.29	0.50	MLT 94	99	
Fusobacterium mortiferum		2.71	0.12	MLZ 5	1	
27671111	**Good identification**					**27671111**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27671112	**Low discrimination**					**27671112**
					GRA	
Actinomyces naeslundii		73.81	0.50	MLT 94	99	
Leptotrichia buccalis		24.07	0.07	MLZ 5	1	
Lactobacillus acidophilus		2.13	-.01		99	
27671113	**Good identification**					**27671113**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27671120	**Very good identification**					**27671120**
					SPO	
Actinomyces naeslundii		98.88	0.72	MLT 94	1	
Clostridium perfringens		0.57	0.11		84	
Clostridium ramosum		0.25	0.02		92	
27671121	**Very good identification**					**27671121**
Actinomyces naeslundii		100.0	0.75	MLT 94		
27671122	**Low discrimination**					**27671122**
					SPO	
Actinomyces naeslundii		85.41	0.72	MLT 94	1	
Lactobacillus acidophilus		12.82	0.33		1	
Clostridium ramosum		0.86	0.12		92	
27671123	**Very good identification**					**27671123**
Actinomyces naeslundii		99.90	0.75	MLT 94		
Lactobacillus acidophilus		0.10	-.01			
27671130	**Good identification**					**27671130**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27671131	**Good identification**					**27671131**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27671132	**Low discrimination**					**27671132**
					GRA	
Actinomyces naeslundii		88.48	0.50	MLT 94	99	
Parabacteroides distasonis		5.76	0.16	MLZ 5	1	
Leptotrichia buccalis		3.21	-.09		1	
27671133	**Good identification**					**27671133**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27671140	**Very good identification**					**27671140**
					GRA	
Actinomyces naeslundii		98.25	0.59	MLT 94	99	
Actinomyces israelii		0.93	0.15		99	
Fusobacterium mortiferum		0.82	0.12		1	
27671141	**Very good identification**					**27671141**
Actinomyces naeslundii		100.0	0.62	MLT 94		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27671142	**Low discrimination**					**27671142**
Actinomyces naeslundii		89.33	0.59	MLT 94		
Lactobacillus acidophilus		9.39	0.18			
Actinomyces israelii		1.27	0.18			
27671143	**Very good identification**					**27671143**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27671150	**Good identification**					**27671150**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27671151	**Good identification**					**27671151**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27671152	**Good identification**					**27671152**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27671153	**Good identification**					**27671153**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27671160	**Very good identification**					**27671160**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27671161	**Very good identification**					**27671161**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27671162	**Good identification**					**27671162**
Actinomyces naeslundii		90.49	0.59	MLT 94		
Lactobacillus acidophilus		9.51	0.18			
27671163	**Very good identification**					**27671163**
Actinomyces naeslundii		100.0	0.62	MLT 94		
27671170	**Good identification**					**27671170**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27671171	**Good identification**					**27671171**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27671172	**Good identification**					**27671172**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27671173	**Good identification**					**27671173**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27671300	**Low discrimination**					**27671300**
					GRA	
Actinomyces naeslundii		87.31	0.37	MLT 94	99	
Fusobacterium mortiferum		12.69	0.12	RHA 1	1	
27671301	**Good identification**					**27671301**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	MLT 94		
27671303	**Good identification**					**27671303**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27671320	**Low discrimination**					**27671320**
					SPO	
Actinomyces naeslundii		75.28	0.37	MLT 94	1	
Clostridium ramosum		24.72	0.04	RHA 1	92	
27671321	**Good identification**					**27671321**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27671323	**Good identification**					**27671323**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27671500	**Low discrimination**					**27671500**
					GRA	
Actinomyces naeslundii		66.90	0.37	MLT 94	99	
Actinomyces israelii		23.38	0.20	ARA 1	99	
Fusobacterium mortiferum		9.72	0.12		1	
27671501	**Good identification**					**27671501**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27671502	**Low discrimination**					**27671502**
Actinomyces naeslundii		59.73	0.37	MLT 94		
Actinomyces israelii		31.31	0.23	ARA 1		
Lactobacillus acidophilus		8.97	-.01			
27671503	**Good identification**					**27671503**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27671520	**Good identification**					**27671520**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27671521	**Good identification**					**27671521**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27671522	**Low discrimination**					**27671522**
					GRA	
Actinomyces naeslundii		65.09	0.37	MLT 94	99	
Parabacteroides distasonis		25.14	0.17	ARA 1	1	
Lactobacillus acidophilus		9.77	-.01		99	
27671523	**Good identification**					**27671523**
Actinomyces naeslundii		100.0	0.41	MLT 94		
27672000	**Acceptable identification**					**27672000**
					SPO	
Actinomyces naeslundii		92.80	0.45	MLT 94	1	
Clostridium innocuum		7.20	0.02	GAL 96	99	
27672001	**Good identification**					**27672001**
Actinomyces naeslundii		100.0	0.48	MLT 94		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27672002	**Acceptable identification**					**27672002**
					SPO	
Actinomyces naeslundii		92.80	0.45	MLT 94	1	
Clostridium innocuum		7.20	0.02	GAL 96	99	
27672003	**Good identification**					**27672003**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27672020	**Good identification**					**27672020**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27672021	**Good identification**					**27672021**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27672022	**Good identification**					**27672022**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27672023	**Good identification**					**27672023**
Actinomyces naeslundii		100.0	0.48	MLT 94		
27672041	**Good identification**					**27672041**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27672043	**Good identification**					**27672043**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27672061	**Good identification**					**27672061**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27672063	**Good identification**					**27672063**
Actinomyces naeslundii		100.0	0.35	MLT 94		
27672100	**Good identification**					**27672100**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27672101	**Good identification**					**27672101**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27672102	**Good identification**					**27672102**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27672103	**Good identification**					**27672103**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27672120	**Good identification**					**27672120**
Actinomyces naeslundii		100.0	0.53	MLT 94		
27672121	**Good identification**					**27672121**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27672122	**Good identification**					**27672122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	MLT 94		
27672123	**Good identification**					**27672123**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27672140	**Good identification**					**27672140**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27672141	**Good identification**					**27672141**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27672142	**Good identification**					**27672142**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27672143	**Good identification**					**27672143**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27672160	**Good identification**					**27672160**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27672161	**Good identification**					**27672161**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27672162	**Good identification**					**27672162**
Actinomyces naeslundii		100.0	0.40	MLT 94		
27672163	**Good identification**					**27672163**
Actinomyces naeslundii		100.0	0.43	MLT 94		
27673000	**Good identification**					**27673000**
					SPO	
Actinomyces naeslundii		93.42	0.69	MLT 94	1	
Clostridium innocuum		5.74	0.24		99	
Actinomyces israelii		0.84	0.25		1	
27673001	**Very good identification**					**27673001**
Actinomyces naeslundii		100.0	0.72	MLT 94		
27673002	**Good identification**					**27673002**
					SPO	
Actinomyces naeslundii		93.03	0.69	MLT 94	1	
Clostridium innocuum		5.72	0.24		99	
Actinomyces israelii		1.25	0.28		1	
27673003	**Very good identification**					**27673003**
Actinomyces naeslundii		100.0	0.72	MLT 94		
27673010	**Good identification**					**27673010**
					SPO	
Actinomyces naeslundii		95.36	0.47	MLT 94	1	
Clostridium innocuum		4.64	0.00	MLZ 5	99	
27673011	**Good identification**					**27673011**
Actinomyces naeslundii		100.0	0.50	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27673012	**Acceptable identification**					**27673012**
					SPO	
Actinomyces naeslundii		92.01	0.47	MLT 94	1	
Clostridium innocuum		4.48	0.00	MLZ 5	99	
Actinomyces israelii		3.51	0.13		1	
27673013	**Good identification**					**27673013**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27673020	**Very good identification**					**27673020**
Actinomyces naeslundii		100.0	0.69	MLT 94		
27673021	**Very good identification**					**27673021**
Actinomyces naeslundii		100.0	0.72	MLT 94		
27673022	**Very good identification**					**27673022**
Actinomyces naeslundii		100.0	0.69	MLT 94		
27673023	**Very good identification**					**27673023**
Actinomyces naeslundii		100.0	0.72	MLT 94		
27673030	**Good identification**					**27673030**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27673031	**Good identification**					**27673031**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27673032	**Good identification**					**27673032**
Actinomyces naeslundii		100.0	0.47	MLT 94		
27673033	**Good identification**					**27673033**
Actinomyces naeslundii		100.0	0.50	MLT 94		
27673040	**Good identification**					**27673040**
Actinomyces naeslundii		98.06	0.56	MLT 94		
Actinomyces israelii		1.94	0.18			
27673041	**Very good identification**					**27673041**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27673042	**Good identification**					**27673042**
Actinomyces naeslundii		97.12	0.56	MLT 94		
Actinomyces israelii		2.88	0.21			
27673043	**Very good identification**					**27673043**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27673051	**Good identification**					**27673051**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27673053	**Good identification**					**27673053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	MLT 94		
27673060	**Good identification**					**27673060**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27673061	**Very good identification**					**27673061**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27673062	**Good identification**					**27673062**
Actinomyces naeslundii		100.0	0.56	MLT 94		
27673063	**Very good identification**					**27673063**
Actinomyces naeslundii		100.0	0.59	MLT 94		
27673071	**Good identification**					**27673071**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27673073	**Good identification**					**27673073**
Actinomyces naeslundii		100.0	0.37	MLT 94		
27673100	**Very good identification**					**27673100**
					GRA	
Actinomyces naeslundii		98.68	0.77	MLT 94	99	
Actinomyces israelii		1.24	0.35		99	
Fusobacterium mortiferum		0.08	0.12		1	
27673101	**Very good identification**					**27673101**
Actinomyces naeslundii		100.0	0.80	MLT 94		
27673102	**Very good identification**					**27673102**
Actinomyces naeslundii		97.76	0.77	MLT 94		
Actinomyces israelii		1.85	0.38			
Lactobacillus acidophilus		0.40	0.11			
27673103	**Very good identification**					**27673103**
Actinomyces naeslundii		100.0	0.80	MLT 94		
27673110	**Good identification**					**27673110**
Actinomyces naeslundii		96.55	0.55	MLT 94		
Actinomyces israelii		3.45	0.21	MLZ 5		
27673111	**Very good identification**					**27673111**
Actinomyces naeslundii		100.0	0.58	MLT 94		
27673112	**Acceptable identification**					**27673112**
Actinomyces naeslundii		94.91	0.55	MLT 94		
Actinomyces israelii		5.09	0.24	MLZ 5		
27673113	**Very good identification**					**27673113**
Actinomyces naeslundii		100.0	0.58	MLT 94		
27673120	**Very good identification**					**27673120**
					SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.20	0.77	MLT 94	1	
Clostridium ramosum		0.73	0.15		92	
Actinomyces israelii		0.08	0.14		1	
27673121	**Very good identification**					**27673121**
Actinomyces naeslundii		100.0	0.80	MLT 94		
27673122	**Very good identification**					**27673122**
					SPO	
Actinomyces naeslundii		96.67	0.77	MLT 94	1	
Clostridium ramosum		2.82	0.25		92	
Lactobacillus acidophilus		0.39	0.11		1	
27673123	**Very good identification**					**27673123**
Actinomyces naeslundii		100.0	0.80	MLT 94		
27673130	**Good identification**					**27673130**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27673131	**Very good identification**					**27673131**
Actinomyces naeslundii		100.0	0.58	MLT 94		
27673132	**Good identification**					**27673132**
Actinomyces naeslundii		100.0	0.55	MLT 94		
27673133	**Very good identification**					**27673133**
Actinomyces naeslundii		100.0	0.58	MLT 94		
27673140	**Very good identification**					**27673140**
Actinomyces naeslundii		97.30	0.64	MLT 94		
Actinomyces israelii		2.70	0.28			
27673141	**Very good identification**					**27673141**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27673142	**Very good identification**					**27673142**
Actinomyces naeslundii		96.00	0.64	MLT 94		
Actinomyces israelii		4.00	0.31			
27673143	**Very good identification**					**27673143**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27673150	**Acceptable identification**					**27673150**
Actinomyces naeslundii		92.70	0.42	MLT 94		
Actinomyces israelii		7.30	0.14	MLZ 5		
27673151	**Good identification**					**27673151**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673152	**Low discrimination**					**27673152**
Actinomyces naeslundii		89.43	0.42	MLT 94		
Actinomyces israelii		10.57	0.17	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27673153	**Good identification**					**27673153**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673160	**Very good identification**					**27673160**
Actinomyces naeslundii		100.0	0.64	MLT 94		
27673161	**Very good identification**					**27673161**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27673162	**Very good identification**					**27673162**
Actinomyces naeslundii		100.0	0.64	MLT 94		
27673163	**Very good identification**					**27673163**
Actinomyces naeslundii		100.0	0.67	MLT 94		
27673170	**Good identification**					**27673170**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27673171	**Good identification**					**27673171**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673172	**Good identification**					**27673172**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27673173	**Good identification**					**27673173**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673201	**Good identification**					**27673201**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673203	**Good identification**					**27673203**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673221	**Good identification**					**27673221**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673223	**Good identification**					**27673223**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673300	**Acceptable identification**					**27673300**
Actinomyces naeslundii		91.42	0.42	MLT 94		
Actinomyces israelii		8.58	0.16	RHA 1		
27673301	**Good identification**					**27673301**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673302	**Low discrimination**					**27673302**
Actinomyces naeslundii		87.66	0.42	MLT 94		
Actinomyces israelii		12.34	0.19	RHA 1		
27673303	**Good identification**					**27673303**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673320	**Low discrimination**					**27673320**
					SPO	
Actinomyces naeslundii		51.06	0.42	MLT 94	1	
Clostridium ramosum		48.94	0.17	RHA 1	92	
27673321	**Good identification**					**27673321**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673323	**Good identification**					**27673323**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673401	**Good identification**					**27673401**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673403	**Good identification**					**27673403**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673421	**Good identification**					**27673421**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673423	**Good identification**					**27673423**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27673501	**Good identification**					**27673501**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673502	**Low discrimination**					**27673502**
Actinomyces israelii		60.48	0.36	MLT 99		
Actinomyces naeslundii		39.52	0.42	XYL 92		
27673503	**Good identification**					**27673503**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673520	**Good identification**					**27673520**
Actinomyces naeslundii		100.0	0.42	MLT 94		
27673521	**Good identification**					**27673521**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27673522	**Acceptable identification**					**27673522**
Actinomyces naeslundii		91.10	0.42	MLT 94		
Actinomyces israelii		8.90	0.16	ARA 1		
27673523	**Good identification**					**27673523**
Actinomyces naeslundii		100.0	0.45	MLT 94		
27675100	**Low discrimination**					**27675100**
					GRA	
Actinomyces israelii		66.34	0.40	MLT 99	99	
Fusobacterium mortiferum		14.63	0.26		1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		13.51	0.37		99		
27675101	**Good identification**						**27675101**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27675102	**Low discrimination**						**27675102**
					SPO		
Actinomyces israelii		84.91	0.43	MLT 99	1		
Actinomyces naeslundii		11.53	0.37		1		
Clostridium butyricum		1.83	-.07		99		
27675103	**Good identification**						**27675103**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27675120	**Low discrimination**						**27675120**
Actinomyces naeslundii		76.13	0.37	MLT 94			
Actinomyces israelii		23.87	0.20	XYL 1			
27675121	**Good identification**						**27675121**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27675122	**Low discrimination**						**27675122**
					GRA		
Parabacteroides distasonis		77.78	0.37	NIT 1	1		
Actinomyces naeslundii		13.71	0.37		99		
Actinomyces israelii		6.45	0.23		99		
27675123	**Good identification**						**27675123**
Actinomyces naeslundii		100.0	0.41	MLT 94			
27675142	**Acceptable identification**						**27675142**
Actinomyces israelii		94.20	0.36	MLT 99			
Actinomyces naeslundii		5.80	0.25				
27675322	**Good identification**						**27675322**
Parabacteroides distasonis		100.0	0.46	NIT 1			
27675332	**Good identification**						**27675332**
Parabacteroides distasonis		100.0	0.41	NIT 1			
27675500	**Low discrimination**						**27675500**
					GRA	SPO	
Actinomyces israelii		41.43	0.39	MLT 99	99	1	
Clostridium butyricum		37.90	0.17		99	99	
Mitsuokella multacida		20.68	0.21		1	1	
27675502	**Low discrimination**						**27675502**
					SPO		
Actinomyces israelii		73.64	0.42	MLT 99	1		
Clostridium butyricum		17.47	0.10		99		
Bifidobacterium adolescentis		6.17	0.17		1		
27675722	**Good identification**						**27675722**
Parabacteroides distasonis		96.57	0.42	NIT 1			
Clostridium clostridioforme		3.43	0.11				
27675732	**Low discrimination**						**27675732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		76.89	0.37	NIT 1		
Clostridium clostridioforme		23.11	0.23			
27677000	**Low discrimination**					**27677000**
					SPO	
Actinomyces israelii		88.54	0.43	MLT 99	1	
Actinomyces naeslundii		8.68	0.35		1	
Clostridium innocuum		2.78	0.02		99	
27677001	**Good identification**					**27677001**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27677002	**Acceptable identification**					**27677002**
					SPO	
Actinomyces israelii		92.06	0.46	MLT 99	1	
Actinomyces naeslundii		6.02	0.35		1	
Clostridium innocuum		1.93	0.02		99	
27677003	**Good identification**					**27677003**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27677021	**Good identification**					**27677021**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27677023	**Good identification**					**27677023**
Actinomyces naeslundii		100.0	0.38	MLT 94		
27677040	**Good identification**					**27677040**
Actinomyces israelii		100.0	0.36	MLT 99		
27677042	**Good identification**					**27677042**
Actinomyces israelii		100.0	0.39	MLT 99		
27677100	**Acceptable identification**					**27677100**
Actinomyces israelii		93.48	0.53	MLT 99		
Actinomyces naeslundii		6.52	0.42			
27677101	**Acceptable identification**					**27677101**
Actinomyces naeslundii		91.20	0.45	MLT 94		
Actinomyces israelii		8.81	0.19	XYL 1		
27677102	**Good identification**					**27677102**
Actinomyces israelii		95.56	0.56	MLT 99		
Actinomyces naeslundii		4.44	0.42			
27677103	**Low discrimination**					**27677103**
Actinomyces naeslundii		87.35	0.45	MLT 94		
Actinomyces israelii		12.65	0.22	XYL 1		
27677110	**Good identification**					**27677110**
Actinomyces israelii		100.0	0.39	MLT 99		
27677112	**Good identification**					**27677112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.42	MLT 99			
27677120	**Low discrimination**						**27677120**
Actinomyces naeslundii		52.22	0.42	MLT 94			
Actinomyces israelii		47.79	0.33	XYL 1			
27677121	**Good identification**						**27677121**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27677122	**Low discrimination**						**27677122**
Actinomyces israelii		57.86	0.36	MLT 99			
Actinomyces naeslundii		42.15	0.42	NAG 6			
27677123	**Good identification**						**27677123**
Actinomyces naeslundii		100.0	0.45	MLT 94			
27677140	**Good identification**						**27677140**
Actinomyces israelii		96.93	0.46	MLT 99			
Actinomyces naeslundii		3.07	0.30				
27677142	**Good identification**						**27677142**
Actinomyces israelii		97.93	0.49	MLT 99			
Actinomyces naeslundii		2.07	0.30				
27677302	**Good identification**						**27677302**
Actinomyces israelii		100.0	0.37	MLT 99			
27677400	**Good identification**						**27677400**
Actinomyces israelii		100.0	0.41	MLT 99			
27677402	**Good identification**						**27677402**
Actinomyces israelii		100.0	0.44	MLT 99			
27677442	**Good identification**						**27677442**
Actinomyces israelii		100.0	0.37	MLT 99			
27677500	**Low discrimination**						**27677500**
					GRA	SPO	
Actinomyces israelii		81.74	0.52	MLT 99	99	1	
Mitsuokella multacida		16.80	0.27		1	1	
Clostridium butyricum		1.47	0.01		99	99	
27677502	**Good identification**						**27677502**
					GRA		
Actinomyces israelii		96.90	0.55	MLT 99	99		
Mitsuokella multacida		1.48	0.11		1		
Bifidobacterium adolescentis		1.17	0.15		99		
27677510	**Low discrimination**						**27677510**
					GRA		
Actinomyces israelii		68.57	0.37	MLT 99	99		
Mitsuokella multacida		31.44	0.19	MLZ 13	1		
27677512	**Low discrimination**						**27677512**
Actinomyces israelii		84.79	0.40	MLT 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		8.39	0.17	MLZ 13		
Bifidobacterium dentium		3.94	-.07			
27677540	**Low discrimination**					**27677540**
					GRA	
Actinomyces israelii		65.42	0.44	MLT 99	99	
Mitsuokella multacida		34.58	0.27		1	
27677542	**Good identification**					**27677542**
					GRA	
Actinomyces israelii		96.23	0.48	MLT 99	99	
Mitsuokella multacida		3.77	0.11		1	
27677702	**Low discrimination**					**27677702**
					GRA	
Actinomyces israelii		72.69	0.35	MLT 99	99	
Mitsuokella multacida		27.31	0.15	RHA 7	1	
27700002	**Very good identification**					**27700002**
Actinomyces viscosus		99.53	0.64	FRU 95		
Actinomyces odontolyticus		0.42	0.30			
Actinomyces naeslundii		0.05	0.32			
27700003	**Good identification**					**27700003**
Actinomyces viscosus		99.77	0.56	FRU 95		
Actinomyces naeslundii		0.23	0.35			
27700102	**Very good identification**					**27700102**
Actinomyces viscosus		99.85	0.68	FRU 95		
Actinomyces naeslundii		0.08	0.40			
Actinomyces odontolyticus		0.07	0.22			
27700103	**Very good identification**					**27700103**
Actinomyces viscosus		99.64	0.60	FRU 95		
Actinomyces naeslundii		0.36	0.43			
27700120	**Low discrimination**					**27700120**
Actinomyces naeslundii		73.02	0.40	FRU 95		
Actinomyces odontolyticus		17.81	0.12	GAL 96		
Actinomyces viscosus		9.18	0.00			
27700121	**Good identification**					**27700121**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27700200	**Good identification**					**27700200**
Actinomyces odontolyticus		100.0	0.39	MNS 9		
27701000	**Low discrimination**					**27701000**
Actinomyces naeslundii		53.08	0.55	FRU 95		
Actinomyces viscosus		27.01	0.26			
Actinomyces odontolyticus		19.91	0.31			
27701001	**Low discrimination**					**27701001**
Actinomyces naeslundii		89.84	0.58	FRU 95		
Actinomyces viscosus		10.16	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27701002	**Very good identification**					**27701002**
Actinomyces viscosus		97.62	0.60	FRU 95		
Actinomyces naeslundii		1.94	0.55			
Actinomyces odontolyticus		0.45	0.27			
27701003	**Acceptable identification**					**27701003**
Actinomyces viscosus		91.80	0.52	FRU 95		
Actinomyces naeslundii		8.20	0.58			
27701011	**Good identification**					**27701011**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27701020	**Low discrimination**					**27701020**
Actinomyces naeslundii		86.82	0.55	FRU 95		
Clostridium perfringens		7.88	0.15			
Actinomyces odontolyticus		5.30	0.17			
27701021	**Very good identification**					**27701021**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27701022	**Low discrimination**					**27701022**
Actinomyces naeslundii		62.93	0.55	FRU 95		
Actinomyces viscosus		32.02	0.26			
Clostridium perfringens		2.69	0.10			
27701023	**Low discrimination**					**27701023**
Actinomyces naeslundii		89.84	0.58	FRU 95		
Actinomyces viscosus		10.16	0.18			
27701031	**Good identification**					**27701031**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27701033	**Good identification**					**27701033**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27701040	**Good identification**					**27701040**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27701041	**Good identification**					**27701041**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27701043	**Low discrimination**					**27701043**
Actinomyces naeslundii		60.95	0.46	FRU 95		
Actinomyces viscosus		39.05	0.18			
27701060	**Acceptable identification**					**27701060**
Actinomyces naeslundii		91.08	0.43	FRU 95		
Clostridium perfringens		8.92	0.03			
27701061	**Good identification**					**27701061**
Actinomyces naeslundii		100.0	0.46	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27701062	**Good identification**					**27701062**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27701063	**Good identification**					**27701063**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27701100	**Low discrimination**					**27701100**
Actinomyces naeslundii		72.87	0.63	FRU 95		
Actinomyces viscosus		23.16	0.31			
Actinomyces odontolyticus		3.03	0.22			
27701101	**Low discrimination**					**27701101**
					SPO	
Actinomyces naeslundii		88.34	0.66	FRU 95	1	
Actinomyces viscosus		6.24	0.22		1	
Clostridium chauvoei		5.43	0.22		99	
27701102	**Very good identification**					**27701102**
Actinomyces viscosus		96.83	0.65	FRU 95		
Actinomyces naeslundii		3.08	0.63			
Actinomyces odontolyticus		0.08	0.19			
27701103	**Low discrimination**					**27701103**
					SPO	
Actinomyces viscosus		87.37	0.57	FRU 95	1	
Actinomyces naeslundii		12.50	0.66		1	
Clostridium chauvoei		0.14	0.10		99	
27701110	**Good identification**					**27701110**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27701111	**Good identification**					**27701111**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27701120	**Low discrimination**					**27701120**
					GRA	
Actinomyces naeslundii		75.61	0.63	FRU 95	99	
Clostridium perfringens		17.69	0.30		99	
Prevotella melaninogenica		6.20	0.15		1	
27701121	**Very good identification**					**27701121**
					SPO	
Actinomyces naeslundii		97.99	0.66	FRU 95	1	
Clostridium chauvoei		2.01	0.14		99	
27701122	**Low discrimination**					**27701122**
Actinomyces naeslundii		67.46	0.63	FRU 95		
Actinomyces viscosus		21.44	0.31			
Clostridium perfringens		7.43	0.24			
27701123	**Good identification**					**27701123**
					SPO	
Actinomyces naeslundii		93.09	0.66	FRU 95	1	
Actinomyces viscosus		6.58	0.22		1	
Clostridium chauvoei		0.34	0.01		99	
27701130	**Good identification**					**27701130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	FRU 95		
27701131	**Good identification**					**27701131**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27701132	**Good identification**					**27701132**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27701133	**Good identification**					**27701133**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27701140	**Good identification**					**27701140**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27701141	**Good identification**					**27701141**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27701143	**Low discrimination**					**27701143**
Actinomyces naeslundii		71.42	0.53	FRU 95		
Actinomyces viscosus		28.58	0.22			
27701160	**Low discrimination**					**27701160**
Actinomyces naeslundii		79.84	0.50	FRU 95		
Clostridium perfringens		20.16	0.18			
27701161	**Good identification**					**27701161**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27701162	**Low discrimination**					**27701162**
Actinomyces naeslundii		89.38	0.50	FRU 95		
Clostridium perfringens		10.62	0.12			
27701163	**Good identification**					**27701163**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27701200	**Good identification**					**27701200**
Actinomyces odontolyticus		100.0	0.36	MNS 9		
27702000	**Good identification**					**27702000**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27702001	**Good identification**					**27702001**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27702020	**Good identification**					**27702020**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27702021	**Good identification**					**27702021**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27702022	**Good identification**					**27702022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	FRU 95		
27702023	**Good identification**					**27702023**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27702100	**Low discrimination**					**27702100**
Actinomyces naeslundii		78.40	0.44	FRU 95		
Propionibacterium propionicum		16.53	0.23	GAL 96		
Actinomyces viscosus		5.08	0.00			
27702101	**Good identification**					**27702101**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27702120	**Good identification**					**27702120**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27702121	**Good identification**					**27702121**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27702122	**Acceptable identification**					**27702122**
Actinomyces naeslundii		93.92	0.44	FRU 95		
Actinomyces viscosus		6.08	0.00	GAL 96		
27702123	**Good identification**					**27702123**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27703000	**Very good identification**					**27703000**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27703001	**Very good identification**					**27703001**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27703002	**Low discrimination**					**27703002**
Actinomyces naeslundii		79.23	0.60	FRU 95		
Actinomyces viscosus		20.77	0.26			
27703003	**Good identification**					**27703003**
Actinomyces naeslundii		94.50	0.63	FRU 95		
Actinomyces viscosus		5.50	0.18			
27703010	**Good identification**					**27703010**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27703011	**Good identification**					**27703011**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27703012	**Good identification**					**27703012**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27703013	**Good identification**					**27703013**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	FRU 95		
27703020	**Very good identification**					**27703020**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27703021	**Very good identification**					**27703021**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27703022	**Very good identification**					**27703022**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27703023	**Very good identification**					**27703023**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27703030	**Good identification**					**27703030**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27703031	**Good identification**					**27703031**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27703032	**Good identification**					**27703032**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27703033	**Good identification**					**27703033**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27703040	**Good identification**					**27703040**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27703041	**Good identification**					**27703041**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27703042	**Good identification**					**27703042**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27703043	**Good identification**					**27703043**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27703060	**Good identification**					**27703060**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27703061	**Good identification**					**27703061**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27703062	**Good identification**					**27703062**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27703063	**Good identification**					**27703063**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27703100	**Very good identification**					**27703100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.13	0.68	FRU 95		
Propionibacterium propionicum		0.87	0.23			
27703101	**Very good identification**					**27703101**
Actinomyces naeslundii		100.0	0.71	FRU 95		
27703102	**Low discrimination**					**27703102**
Actinomyces naeslundii		84.97	0.68	FRU 95		
Actinomyces viscosus		13.91	0.31			
Propionibacterium propionicum		1.12	0.26			
27703103	**Very good identification**					**27703103**
Actinomyces naeslundii		96.49	0.71	FRU 95		
Actinomyces viscosus		3.51	0.22			
27703110	**Good identification**					**27703110**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27703111	**Good identification**					**27703111**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27703112	**Good identification**					**27703112**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27703113	**Good identification**					**27703113**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27703120	**Very good identification**					**27703120**
Actinomyces naeslundii		100.0	0.68	FRU 95		
27703121	**Very good identification**					**27703121**
Actinomyces naeslundii		100.0	0.71	FRU 95		
27703122	**Very good identification**					**27703122**
Actinomyces naeslundii		100.0	0.68	FRU 95		
27703123	**Very good identification**					**27703123**
Actinomyces naeslundii		100.0	0.71	FRU 95		
27703130	**Good identification**					**27703130**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27703131	**Good identification**					**27703131**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27703132	**Good identification**					**27703132**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27703133	**Good identification**					**27703133**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27703140	**Good identification**					**27703140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.26	0.55	FRU 95		
Propionibacterium propionicum		4.74	0.23			
27703141	**Very good identification**					**27703141**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27703142	**Acceptable identification**					**27703142**
Actinomyces naeslundii		93.05	0.55	FRU 95		
Propionibacterium propionicum		6.95	0.26			
27703143	**Very good identification**					**27703143**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27703151	**Good identification**					**27703151**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27703153	**Good identification**					**27703153**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27703160	**Good identification**					**27703160**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27703161	**Very good identification**					**27703161**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27703162	**Good identification**					**27703162**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27703163	**Very good identification**					**27703163**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27703171	**Good identification**					**27703171**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27703173	**Good identification**					**27703173**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27703301	**Good identification**					**27703301**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27703303	**Good identification**					**27703303**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27703321	**Good identification**					**27703321**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27703323	**Good identification**					**27703323**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27703501	**Good identification**					**27703501**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	FRU 95		
27703503	**Good identification**					**27703503**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27703521	**Good identification**					**27703521**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27703523	**Good identification**					**27703523**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27707101	**Good identification**					**27707101**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27707103	**Good identification**					**27707103**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27707121	**Good identification**					**27707121**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27707123	**Good identification**					**27707123**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27710000	**Low discrimination**					**27710000**
Actinomyces viscosus		66.74	0.52	BGL 99		
Actinomyces odontolyticus		27.37	0.52			
Actinomyces naeslundii		3.35	0.54			
27710001	**Low discrimination**					**27710001**
Actinomyces viscosus		80.76	0.44	BGL 99		
Actinomyces naeslundii		18.24	0.57			
Actinomyces odontolyticus		1.00	0.18			
27710002	**Very good identification**					**27710002**
Actinomyces viscosus		99.69	0.86			
Actinomyces odontolyticus		0.25	0.48			
Actinomyces naeslundii		0.05	0.54			
27710003	**Very good identification**					**27710003**
Actinomyces viscosus		99.77	0.78			
Actinomyces naeslundii		0.23	0.57			
Actinomyces odontolyticus		0.01	0.14			
27710012	**Good identification**					**27710012**
Actinomyces viscosus		99.49	0.52	MLZ 1		
Actinomyces naeslundii		0.26	0.32			
Actinomyces odontolyticus		0.25	0.14			
27710013	**Good identification**					**27710013**
Actinomyces viscosus		98.83	0.44	MLZ 1		
Actinomyces naeslundii		1.18	0.35			
27710020	**Low discrimination**					**27710020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		50.27	0.39	MNS 9		
Actinomyces naeslundii		37.79	0.54	RAF 3		
Actinomyces viscosus		7.61	0.18	NAG 14		
27710021	**Very good identification**					**27710021**
Actinomyces naeslundii		95.72	0.57	GAL 96		
Actinomyces viscosus		4.28	0.09			
27710022	**Acceptable identification**					**27710022**
Actinomyces viscosus		91.42	0.52	NAG 1		
Actinomyces naeslundii		4.59	0.54			
Actinomyces odontolyticus		3.74	0.35			
27710023	**Low discrimination**					**27710023**
Actinomyces viscosus		81.58	0.44	NAG 1		
Actinomyces naeslundii		18.42	0.57			
27710041	**Low discrimination**					**27710041**
Actinomyces naeslundii		79.78	0.44	GAL 96		
Actinomyces viscosus		20.22	0.09			
27710042	**Good identification**					**27710042**
Actinomyces viscosus		98.87	0.52	SOR 1		
Actinomyces naeslundii		0.88	0.41			
Actinomyces odontolyticus		0.25	0.14			
27710043	**Good identification**					**27710043**
Actinomyces viscosus		96.17	0.44	SOR 1		
Actinomyces naeslundii		3.83	0.44			
27710060	**Low discrimination**					**27710060**
Actinomyces naeslundii		88.99	0.41	GAL 96		
Clostridium perfringens		11.01	0.03			
27710061	**Good identification**					**27710061**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27710063	**Low discrimination**					**27710063**
Actinomyces naeslundii		79.78	0.44	GAL 96		
Actinomyces viscosus		20.22	0.09			
27710100	**Low discrimination**					**27710100**
Actinomyces viscosus		83.77	0.56	BGL 99		
Actinomyces naeslundii		6.73	0.61			
Actinomyces odontolyticus		6.10	0.44			
27710101	**Low discrimination**					**27710101**
Actinomyces viscosus		73.33	0.48	BGL 99		
Actinomyces naeslundii		26.51	0.64			
Actinomyces odontolyticus		0.16	0.09			
27710102	**Very good identification**					**27710102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		99.87	0.90			
Actinomyces naeslundii		0.08	0.61			
Actinomyces odontolyticus		0.05	0.40			
27710103	**Very good identification**					**27710103**
Actinomyces viscosus		99.64	0.82			
Actinomyces naeslundii		0.36	0.64			
Actinomyces odontolyticus		0.00	0.06			
27710111	**Low discrimination**					**27710111**
Actinomyces naeslundii		65.33	0.43	GAL 96		
Actinomyces viscosus		34.67	0.14	MLZ 5		
27710112	**Good identification**					**27710112**
Actinomyces viscosus		99.53	0.56	MLZ 1		
Actinomyces naeslundii		0.42	0.40			
Actinomyces odontolyticus		0.05	0.06			
27710113	**Good identification**					**27710113**
Actinomyces viscosus		98.13	0.48	MLZ 1		
Actinomyces naeslundii		1.87	0.43			
27710120	**Low discrimination**					**27710120**
Actinomyces naeslundii		63.36	0.61	GAL 96		
Clostridium perfringens		18.72	0.30			
Actinomyces odontolyticus		9.35	0.30			
27710121	**Very good identification**					**27710121**
Actinomyces naeslundii		97.28	0.64	GAL 96		
Actinomyces viscosus		2.72	0.14			
27710122	**Acceptable identification**					**27710122**
Actinomyces viscosus		90.72	0.56	NAG 1		
Actinomyces naeslundii		7.29	0.61			
Clostridium perfringens		1.01	0.24			
27710123	**Low discrimination**					**27710123**
Actinomyces viscosus		73.44	0.48	NAG 1		
Actinomyces naeslundii		26.56	0.64			
27710130	**Good identification**					**27710130**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27710131	**Good identification**					**27710131**
Actinomyces naeslundii		100.0	0.43	GAL 96		
27710133	**Low discrimination**					**27710133**
Actinomyces naeslundii		65.33	0.43	GAL 96		
Actinomyces viscosus		34.67	0.14	MLZ 5		
27710140	**Low discrimination**					**27710140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		53.01	0.49	GAL 96		
Actinomyces viscosus		37.77	0.22			
Propionibacterium propionicum		6.47	0.23			
27710141	**Low discrimination**					**27710141**
Actinomyces naeslundii		86.33	0.52	GAL 96		
Actinomyces viscosus		13.67	0.14			
27710142	**Good identification**					**27710142**
Actinomyces viscosus		98.31	0.56	SOR 1		
Actinomyces naeslundii		1.39	0.49			
Propionibacterium propionicum		0.26	0.26			
27710143	**Acceptable identification**					**27710143**
Actinomyces viscosus		94.00	0.48	SOR 1		
Actinomyces naeslundii		6.00	0.52			
27710160	**Low discrimination**					**27710160**
Actinomyces naeslundii		75.82	0.49	GAL 96		
Clostridium perfringens		24.18	0.18			
27710161	**Good identification**					**27710161**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27710162	**Low discrimination**					**27710162**
Actinomyces naeslundii		53.69	0.49	GAL 96		
Actinomyces viscosus		38.25	0.22			
Clostridium perfringens		8.06	0.12			
27710163	**Low discrimination**					**27710163**
Actinomyces naeslundii		86.33	0.52	GAL 96		
Actinomyces viscosus		13.67	0.14			
27710200	**Very good identification**					**27710200**
Actinomyces odontolyticus		98.75	0.57	MNS 9		
Actinomyces viscosus		1.25	0.18	RAF 3		
27710202	**Low discrimination**					**27710202**
Actinomyces viscosus		67.21	0.52	RHA 1		
Actinomyces odontolyticus		32.79	0.53			
27710203	**Good identification**					**27710203**
Actinomyces viscosus		98.32	0.44	RHA 1		
Actinomyces odontolyticus		1.45	0.19			
Actinomyces naeslundii		0.22	0.23			
27710220	**Good identification**					**27710220**
Actinomyces odontolyticus		100.0	0.44	MNS 9		
27710222	**Low discrimination**					**27710222**
Actinomyces odontolyticus		88.72	0.40	MNS 9		
Actinomyces viscosus		11.28	0.18	RAF 3		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27710300	**Acceptable identification**					**27710300**
Actinomyces odontolyticus		92.84	0.48	MNS 9		
Actinomyces viscosus		6.63	0.22	RAF 3		
Actinomyces naeslundii		0.53	0.27			
27710302	**Acceptable identification**					**27710302**
Actinomyces viscosus		91.96	0.56	RHA 1		
Actinomyces odontolyticus		7.97	0.45			
Actinomyces naeslundii		0.08	0.27			
27710303	**Good identification**					**27710303**
Actinomyces viscosus		99.38	0.48	RHA 1		
Actinomyces naeslundii		0.36	0.30			
Actinomyces odontolyticus		0.26	0.11			
27710402	**Good identification**					**27710402**
Actinomyces viscosus		99.75	0.52	ARA 1		
Actinomyces odontolyticus		0.25	0.14			
27710403	**Good identification**					**27710403**
Actinomyces viscosus		99.77	0.44	ARA 1		
Actinomyces naeslundii		0.23	0.23			
27710502	**Good identification**					**27710502**
Actinomyces viscosus		99.87	0.56	ARA 1		
Actinomyces naeslundii		0.08	0.27			
Actinomyces odontolyticus		0.05	0.06			
27710503	**Good identification**					**27710503**
Actinomyces viscosus		99.64	0.48	ARA 1		
Actinomyces naeslundii		0.36	0.30			
27711000	**Low discrimination**					**27711000**
Actinomyces naeslundii		55.64	0.77			
Actinomyces viscosus		28.31	0.48			
Actinomyces odontolyticus		12.63	0.49			
27711001	**Low discrimination**					**27711001**
Actinomyces naeslundii		89.72	0.80			
Actinomyces viscosus		10.15	0.40			
Actinomyces odontolyticus		0.14	0.15			
27711002	**Very good identification**					**27711002**
Actinomyces viscosus		97.76	0.82			
Actinomyces naeslundii		1.94	0.77			
Actinomyces odontolyticus		0.27	0.45			
27711003	**Good identification**					**27711003**
Actinomyces viscosus		91.79	0.74			
Actinomyces naeslundii		8.20	0.80			
Actinomyces odontolyticus		0.01	0.11			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27711010	**Low discrimination**					**27711010**
Actinomyces naeslundii		87.63	0.55	MLZ 5		
Actinomyces viscosus		8.56	0.14			
Actinomyces odontolyticus		3.82	0.15			
27711011	**Very good identification**					**27711011**
Actinomyces naeslundii		97.88	0.58	MLZ 5		
Actinomyces viscosus		2.12	0.06			
27711012	**Acceptable identification**					**27711012**
Actinomyces viscosus		90.40	0.48	MLZ 1		
Actinomyces naeslundii		9.35	0.55			
Actinomyces odontolyticus		0.25	0.11			
27711013	**Low discrimination**					**27711013**
Actinomyces viscosus		68.24	0.40	MLZ 1		
Actinomyces naeslundii		31.76	0.58			
27711020	**Low discrimination**					**27711020**
Actinomyces naeslundii		65.90	0.77			
Clostridium perfringens		31.18	0.49			
Actinomyces odontolyticus		2.44	0.36			
27711021	**Very good identification**					**27711021**
Actinomyces naeslundii		99.57	0.80			
Clostridium perfringens		0.32	0.15			
Actinomyces viscosus		0.11	0.06			
27711022	**Low discrimination**					**27711022**
Actinomyces naeslundii		56.26	0.77			
Actinomyces viscosus		28.63	0.48			
Clostridium perfringens		12.52	0.44			
27711023	**Low discrimination**					**27711023**
Actinomyces naeslundii		89.72	0.80			
Actinomyces viscosus		10.15	0.40			
Clostridium perfringens		0.14	0.10			
27711030	**Acceptable identification**					**27711030**
Actinomyces naeslundii		91.68	0.55	MLZ 5		
Clostridium perfringens		8.32	0.15			
27711031	**Very good identification**					**27711031**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27711032	**Low discrimination**					**27711032**
Actinomyces naeslundii		87.69	0.55	MLZ 5		
Actinomyces viscosus		8.56	0.14			
Clostridium perfringens		3.75	0.10			
27711033	**Very good identification**					**27711033**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.88	0.58	MLZ 5		
Actinomyces viscosus		2.12	0.06			
27711040	**Good identification**					**27711040**
Actinomyces naeslundii		94.53	0.64			
Actinomyces viscosus		2.75	0.14			
Clostridium perfringens		1.49	0.11			
27711041	**Very good identification**					**27711041**
Actinomyces naeslundii		99.36	0.67			
Actinomyces viscosus		0.64	0.06			
27711042	**Low discrimination**					**27711042**
Actinomyces viscosus		73.96	0.48	SOR 1		
Actinomyces naeslundii		25.65	0.64			
Actinomyces odontolyticus		0.20	0.11			
27711043	**Low discrimination**					**27711043**
Actinomyces naeslundii		60.95	0.67			
Actinomyces viscosus		39.05	0.40			
27711050	**Good identification**					**27711050**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27711051	**Good identification**					**27711051**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
27711052	**Low discrimination**					**27711052**
Actinomyces naeslundii		64.38	0.43	MLZ 5		
Actinomyces viscosus		35.62	0.14			
27711053	**Low discrimination**					**27711053**
Actinomyces naeslundii		89.05	0.46	MLZ 5		
Actinomyces viscosus		10.95	0.06			
27711060	**Low discrimination**					**27711060**
Actinomyces naeslundii		66.20	0.64			
Clostridium perfringens		33.80	0.37			
27711061	**Very good identification**					**27711061**
Actinomyces naeslundii		99.66	0.67			
Clostridium perfringens		0.34	0.03			
27711062	**Low discrimination**					**27711062**
Actinomyces naeslundii		78.78	0.64			
Clostridium perfringens		18.93	0.32			
Actinomyces viscosus		2.29	0.14			
27711063	**Very good identification**					**27711063**
Actinomyces naeslundii		99.36	0.67			
Actinomyces viscosus		0.64	0.06			
27711070	**Acceptable identification**					**27711070**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.08	0.43	MLZ 5		
Clostridium perfringens		8.92	0.03			
27711071	**Good identification**					**27711071**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
27711072	**Good identification**					**27711072**
Actinomyces naeslundii		100.0	0.43	MLZ 5		
27711073	**Good identification**					**27711073**
Actinomyces naeslundii		100.0	0.46	MLZ 5		
27711100	**Low discrimination**					**27711100**
Actinomyces naeslundii		70.58	0.85			
Actinomyces viscosus		22.43	0.52			
Clostridium perfringens		2.66	0.38			
27711101	**Good identification**					**27711101**
					SPO	
Actinomyces naeslundii		92.70	0.88		1	
Actinomyces viscosus		6.55	0.44		1	
Clostridium chauvoei		0.70	0.29		99	
27711102	**Very good identification**					**27711102**
Actinomyces viscosus		96.70	0.87			
Actinomyces naeslundii		3.07	0.85			
Bifidobacterium longum subsp. infantis		0.07	0.27			
27711103	**Low discrimination**					**27711103**
					SPO	
Actinomyces viscosus		87.47	0.78		1	
Actinomyces naeslundii		12.51	0.88		1	
Clostridium chauvoei		0.02	0.16		99	
27711110	**Good identification**					**27711110**
Actinomyces naeslundii		92.70	0.63	MLZ 5		
Actinomyces viscosus		5.65	0.18			
Clostridium perfringens		0.67	0.04			
27711111	**Very good identification**					**27711111**
Actinomyces naeslundii		98.66	0.66	MLZ 5		
Actinomyces viscosus		1.34	0.10			
27711112	**Low discrimination**					**27711112**
Actinomyces viscosus		85.24	0.52	MLZ 1		
Actinomyces naeslundii		14.12	0.63			
Bifidobacterium adolescentis		0.43	0.22			
27711113	**Low discrimination**					**27711113**
Actinomyces viscosus		57.30	0.44	MLZ 1		
Actinomyces naeslundii		42.70	0.66			
27711120	**Low discrimination**					**27711120**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		53.50	0.64		99	
Actinomyces naeslundii		43.89	0.85		99	
Prevotella melaninogenica		1.70	0.32		1	
27711121	**Very good identification**					**27711121**
					SPO	
Actinomyces naeslundii		98.84	0.88		1	
Clostridium perfringens		0.81	0.30		84	
Clostridium chauvoei		0.25	0.21		99	
27711122	**Low discrimination**					**27711122**
Actinomyces naeslundii		47.95	0.85			
Clostridium perfringens		27.50	0.59			
Actinomyces viscosus		15.24	0.52			
27711123	**Good identification**					**27711123**
Actinomyces naeslundii		92.92	0.88			
Actinomyces viscosus		6.56	0.44			
Clostridium perfringens		0.36	0.24			
27711130	**Low discrimination**					**27711130**
					GRA	
Actinomyces naeslundii		80.20	0.63	MLZ 5	99	
Clostridium perfringens		18.76	0.30		99	
Prevotella melaninogenica		0.60	-.03		1	
27711131	**Very good identification**					**27711131**
Actinomyces naeslundii		100.0	0.66	MLZ 5		
27711132	**Low discrimination**					**27711132**
Actinomyces naeslundii		78.99	0.63	MLZ 5		
Clostridium perfringens		8.70	0.24			
Actinomyces viscosus		4.82	0.18			
27711133	**Very good identification**					**27711133**
Actinomyces naeslundii		98.66	0.66	MLZ 5		
Actinomyces viscosus		1.34	0.10			
27711140	**Good identification**					**27711140**
Actinomyces naeslundii		93.44	0.72			
Clostridium perfringens		3.80	0.26			
Actinomyces viscosus		1.70	0.18			
27711141	**Very good identification**					**27711141**
Actinomyces naeslundii		99.60	0.75			
Actinomyces viscosus		0.40	0.10			
27711142	**Low discrimination**					**27711142**
Actinomyces viscosus		63.52	0.52	SOR 1		
Actinomyces naeslundii		35.27	0.72			
Clostridium perfringens		0.68	0.20			
27711143	**Low discrimination**					**27711143**
Actinomyces naeslundii		71.42	0.75			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		28.58	0.44				
27711150	**Good identification**						**27711150**
Actinomyces naeslundii		100.0	0.50	MLZ 5			
27711151	**Good identification**						**27711151**
Actinomyces naeslundii		100.0	0.53	MLZ 5			
27711152	**Low discrimination**						**27711152**
Actinomyces naeslundii		74.31	0.50	MLZ 5			
Actinomyces viscosus		25.69	0.18				
27711153	**Acceptable identification**						**27711153**
Actinomyces naeslundii		92.87	0.53	MLZ 5			
Actinomyces viscosus		7.13	0.10				
27711160	**Low discrimination**						**27711160**
					GRA		
Clostridium perfringens		56.72	0.52		99		
Actinomyces naeslundii		43.11	0.72		99		
Prevotella melaninogenica		0.10	-.03		1		
27711161	**Very good identification**						**27711161**
Actinomyces naeslundii		99.12	0.75				
Clostridium perfringens		0.88	0.18				
27711162	**Low discrimination**						**27711162**
Actinomyces naeslundii		60.21	0.72				
Clostridium perfringens		37.27	0.46				
Actinomyces viscosus		1.10	0.18				
27711163	**Very good identification**						**27711163**
Actinomyces naeslundii		99.19	0.75				
Clostridium perfringens		0.41	0.12				
Actinomyces viscosus		0.40	0.10				
27711170	**Low discrimination**						**27711170**
Actinomyces naeslundii		79.84	0.50	MLZ 5			
Clostridium perfringens		20.16	0.18				
27711171	**Good identification**						**27711171**
Actinomyces naeslundii		100.0	0.53	MLZ 5			
27711172	**Low discrimination**						**27711172**
					GRA	CAT	
Actinomyces naeslundii		86.97	0.50	MLZ 5	99	25	
Clostridium perfringens		10.33	0.12		99	1	
Capnocytophaga ochracea		2.70	-.01		1	90	
27711173	**Good identification**						**27711173**
Actinomyces naeslundii		100.0	0.53	MLZ 5			
27711200	**Good identification**						**27711200**
Actinomyces odontolyticus		96.66	0.54	MNS 9			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		2.22	0.43	RAF 3		
Actinomyces viscosus		1.13	0.14			
27711201	**Low discrimination**					**27711201**
Actinomyces naeslundii		71.08	0.46	RHA 1		
Actinomyces odontolyticus		20.88	0.20			
Actinomyces viscosus		8.04	0.06			
27711202	**Low discrimination**					**27711202**
Actinomyces viscosus		64.49	0.48	RHA 1		
Actinomyces odontolyticus		34.23	0.50			
Actinomyces naeslundii		1.28	0.43			
27711203	**Acceptable identification**					**27711203**
Actinomyces viscosus		90.46	0.40	RHA 1		
Actinomyces naeslundii		8.08	0.46			
Actinomyces odontolyticus		1.46	0.16			
27711220	**Low discrimination**					**27711220**
Actinomyces odontolyticus		82.82	0.40	MNS 9		
Actinomyces naeslundii		11.66	0.43	RAF 3		
Clostridium perfringens		5.52	0.15	NAG 14		
27711221	**Good identification**					**27711221**
Actinomyces naeslundii		95.44	0.46	RHA 1		
Actinomyces odontolyticus		4.56	0.06			
27711222	**Low discrimination**					**27711222**
					GRA	
Actinomyces odontolyticus		62.14	0.37	MNS 9	99	
Actinomyces naeslundii		14.28	0.43	RAF 3	99	
Prevotella oralis		13.14	0.17	NAG 14	1	
27711223	**Low discrimination**					**27711223**
Actinomyces naeslundii		89.84	0.46	RHA 1		
Actinomyces viscosus		10.16	0.06			
27711300	**Low discrimination**					**27711300**
Actinomyces odontolyticus		77.75	0.45	MNS 9		
Actinomyces naeslundii		16.07	0.51	RAF 3		
Actinomyces viscosus		5.11	0.18			
27711301	**Acceptable identification**					**27711301**
Actinomyces naeslundii		90.64	0.54	RHA 1		
Actinomyces viscosus		6.40	0.10			
Actinomyces odontolyticus		2.95	0.11			
27711302	**Low discrimination**					**27711302**
Actinomyces viscosus		88.09	0.52	RHA 1		
Actinomyces odontolyticus		8.30	0.42			
Actinomyces naeslundii		2.80	0.51			
27711303	**Low discrimination**					**27711303**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		87.27	0.44	RHA 1		
Actinomyces naeslundii		12.48	0.54			
Actinomyces odontolyticus		0.25	0.08			
27711321	**Good identification**					**27711321**
Actinomyces naeslundii		100.0	0.54	RHA 1		
27711322	**Low discrimination**					**27711322**
					GRA	
Prevotella oralis		63.71	0.37	NIT 1	1	
Actinomyces naeslundii		12.58	0.51		99	
Clostridium perfringens		7.22	0.24		99	
27711323	**Acceptable identification**					**27711323**
					GRA	
Actinomyces naeslundii		90.52	0.54	RHA 1	99	
Actinomyces viscosus		6.39	0.10		99	
Prevotella oralis		3.09	0.03		1	
27711340	**Low discrimination**					**27711340**
Actinomyces naeslundii		78.31	0.38	RHA 1		
Actinomyces odontolyticus		21.69	0.11			
27711341	**Good identification**					**27711341**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27711343	**Low discrimination**					**27711343**
Actinomyces naeslundii		71.42	0.41	RHA 1		
Actinomyces viscosus		28.58	0.10			
27711361	**Good identification**					**27711361**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27711362	**Low discrimination**					**27711362**
					GRA	
Actinomyces naeslundii		52.39	0.38	RHA 1	99	
Clostridium perfringens		32.43	0.12		99	
Prevotella oralis		15.19	0.03		1	
27711363	**Good identification**					**27711363**
Actinomyces naeslundii		100.0	0.41	RHA 1		
27711400	**Low discrimination**					**27711400**
Actinomyces naeslundii		50.33	0.43	ARA 1		
Actinomyces viscosus		25.61	0.14			
Actinomyces odontolyticus		11.42	0.15			
27711401	**Low discrimination**					**27711401**
Actinomyces naeslundii		89.84	0.46	ARA 1		
Actinomyces viscosus		10.16	0.06			
27711402	**Good identification**					**27711402**
Actinomyces viscosus		97.24	0.48	ARA 1		
Actinomyces naeslundii		1.93	0.43			
Lactobacillus fermentum		0.33	0.20			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27711403	**Acceptable identification**						**27711403**
Actinomyces viscosus		91.80	0.40	ARA 1			
Actinomyces naeslundii		8.20	0.46				
27711420	**Low discrimination**						**27711420**
Actinomyces naeslundii		67.89	0.43	ARA 1			
Clostridium perfringens		32.11	0.15				
27711421	**Good identification**						**27711421**
Actinomyces naeslundii		100.0	0.46	ARA 1			
27711422	**Low discrimination**						**27711422**
Actinomyces naeslundii		57.76	0.43	ARA 1			
Actinomyces viscosus		29.39	0.14				
Clostridium perfringens		12.86	0.10				
27711423	**Low discrimination**						**27711423**
Actinomyces naeslundii		89.84	0.46	ARA 1			
Actinomyces viscosus		10.16	0.06				
27711500	**Low discrimination**						**27711500**
Actinomyces naeslundii		60.29	0.51	ARA 1			
Actinomyces viscosus		19.16	0.18				
Actinomyces israelii		6.85	0.26				
27711501	**Acceptable identification**						**27711501**
Actinomyces naeslundii		93.40	0.54	ARA 1			
Actinomyces viscosus		6.60	0.10				
27711502	**Acceptable identification**						**27711502**
Actinomyces viscosus		93.36	0.52	ARA 1			
Actinomyces naeslundii		2.97	0.51				
Bifidobacterium adolescentis		2.57	0.34				
27711503	**Low discrimination**						**27711503**
Actinomyces viscosus		87.49	0.44	ARA 1			
Actinomyces naeslundii		12.51	0.54				
27711512	**Low discrimination**						**27711512**
Bifidobacterium adolescentis		69.76	0.36	NIT 1			
Actinomyces viscosus		20.94	0.18	XYL 87			
Bifidobacterium longum subsp. longum		4.15	0.13				
27711521	**Good identification**						**27711521**
Actinomyces naeslundii		100.0	0.54	ARA 1			
27711522	**Low discrimination**						**27711522**
					GRA	CAT	
Actinomyces naeslundii		36.64	0.51	ARA 1	99	25	
Clostridium perfringens		21.02	0.24		99	1	
Capnocytophaga ochracea		14.93	0.19		1	90	
27711523	**Acceptable identification**						**27711523**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		93.40	0.54	ARA 1			
Actinomyces viscosus		6.60	0.10				
27711540	**Low discrimination**						**27711540**
Actinomyces naeslundii		79.97	0.38	ARA 1			
Actinomyces israelii		20.04	0.19				
27711541	**Good identification**						**27711541**
Actinomyces naeslundii		100.0	0.41	ARA 1			
27711543	**Low discrimination**						**27711543**
Actinomyces naeslundii		71.42	0.41	ARA 1			
Actinomyces viscosus		28.58	0.10				
27711561	**Good identification**						**27711561**
Actinomyces naeslundii		100.0	0.41	ARA 1			
27711562	**Low discrimination**						**27711562**
					GRA	CAT	
Actinomyces naeslundii		53.32	0.38	ARA 1	99	25	
Clostridium perfringens		33.01	0.12		99	1	
Capnocytophaga ochracea		13.67	0.02		1	90	
27711563	**Good identification**						**27711563**
Actinomyces naeslundii		100.0	0.41	ARA 1			
27712000	**Low discrimination**						**27712000**
Actinomyces naeslundii		83.32	0.59	GAL 96			
Actinomyces viscosus		8.64	0.18				
Actinomyces odontolyticus		3.54	0.18				
27712001	**Very good identification**						**27712001**
Actinomyces naeslundii		97.75	0.62	GAL 96			
Actinomyces viscosus		2.25	0.09				
27712002	**Acceptable identification**						**27712002**
Actinomyces viscosus		90.53	0.52	MAN 1			
Actinomyces naeslundii		8.82	0.59				
Propionibacterium propionicum		0.42	0.26				
27712003	**Low discrimination**						**27712003**
Actinomyces viscosus		69.52	0.44	MAN 1			
Actinomyces naeslundii		30.48	0.62				
27712010	**Good identification**						**27712010**
Actinomyces naeslundii		100.0	0.37	GAL 96			
27712011	**Good identification**						**27712011**
Actinomyces naeslundii		100.0	0.40	GAL 96			
27712013	**Low discrimination**						**27712013**
Actinomyces naeslundii		69.55	0.40	GAL 96			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		30.45	0.09	MLZ 5		
27712020	**Very good identification**					**27712020**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27712021	**Very good identification**					**27712021**
Actinomyces naeslundii		100.0	0.62	GAL 96		
27712022	**Good identification**					**27712022**
Actinomyces naeslundii		90.61	0.59	GAL 96		
Actinomyces viscosus		9.39	0.18			
27712023	**Very good identification**					**27712023**
Actinomyces naeslundii		97.75	0.62	GAL 96		
Actinomyces viscosus		2.25	0.09			
27712030	**Good identification**					**27712030**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27712031	**Good identification**					**27712031**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27712032	**Good identification**					**27712032**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27712033	**Good identification**					**27712033**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27712040	**Low discrimination**					**27712040**
Actinomyces naeslundii		84.82	0.46	GAL 96		
Propionibacterium propionicum		15.18	0.23			
27712041	**Good identification**					**27712041**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27712042	**Low discrimination**					**27712042**
Actinomyces naeslundii		53.88	0.46	GAL 96		
Actinomyces viscosus		31.66	0.18			
Propionibacterium propionicum		14.46	0.26			
27712043	**Low discrimination**					**27712043**
Actinomyces naeslundii		88.45	0.49	GAL 96		
Actinomyces viscosus		11.55	0.09			
27712060	**Good identification**					**27712060**
Actinomyces naeslundii		100.0	0.46	GAL 96		
27712061	**Good identification**					**27712061**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27712062	**Good identification**					**27712062**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	GAL 96		
27712063	**Good identification**					**27712063**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27712100	**Low discrimination**					**27712100**
Propionibacterium propionicum		50.66	0.57	SAL 1		
Actinomyces naeslundii		46.13	0.66			
Actinomyces viscosus		2.99	0.22			
27712101	**Good identification**					**27712101**
Actinomyces naeslundii		91.26	0.69	GAL 96		
Propionibacterium propionicum		7.43	0.41			
Actinomyces viscosus		1.31	0.14			
27712102	**Low discrimination**					**27712102**
Actinomyces viscosus		70.75	0.56	MAN 1		
Propionibacterium propionicum		18.18	0.60			
Actinomyces naeslundii		11.04	0.66			
27712103	**Low discrimination**					**27712103**
Actinomyces viscosus		55.94	0.48	MAN 1		
Actinomyces naeslundii		39.27	0.69			
Propionibacterium propionicum		4.79	0.44			
27712110	**Low discrimination**					**27712110**
Actinomyces naeslundii		82.59	0.44	GAL 96		
Propionibacterium propionicum		17.41	0.23	MLZ 5		
27712111	**Good identification**					**27712111**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27712113	**Low discrimination**					**27712113**
Actinomyces naeslundii		78.53	0.48	GAL 96		
Actinomyces viscosus		21.47	0.14	MLZ 5		
27712120	**Very good identification**					**27712120**
Actinomyces naeslundii		98.90	0.66	GAL 96		
Propionibacterium propionicum		1.10	0.23			
27712121	**Very good identification**					**27712121**
Actinomyces naeslundii		100.0	0.69	GAL 96		
27712122	**Good identification**					**27712122**
Actinomyces naeslundii		92.47	0.66	GAL 96		
Actinomyces viscosus		5.99	0.22			
Propionibacterium propionicum		1.54	0.26			
27712123	**Very good identification**					**27712123**
Actinomyces naeslundii		98.58	0.69	GAL 96		
Actinomyces viscosus		1.42	0.14			
27712130	**Good identification**					**27712130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	GAL 96		
27712131	**Good identification**					**27712131**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27712132	**Good identification**					**27712132**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27712133	**Good identification**					**27712133**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27712140	**Low discrimination**					**27712140**
Propionibacterium propionicum		86.16	0.57	SAL 1		
Actinomyces naeslundii		13.84	0.53			
27712141	**Low discrimination**					**27712141**
Actinomyces naeslundii		68.45	0.57	GAL 96		
Propionibacterium propionicum		31.55	0.41			
27712142	**Low discrimination**					**27712142**
Propionibacterium propionicum		87.23	0.60	SAL 1		
Actinomyces naeslundii		9.34	0.53			
Actinomyces viscosus		3.43	0.22			
27712143	**Low discrimination**					**27712143**
Actinomyces naeslundii		56.40	0.57	GAL 96		
Propionibacterium propionicum		39.00	0.44			
Actinomyces viscosus		4.60	0.14			
27712160	**Acceptable identification**					**27712160**
Actinomyces naeslundii		94.09	0.53	GAL 96		
Propionibacterium propionicum		5.92	0.23			
27712161	**Very good identification**					**27712161**
Actinomyces naeslundii		100.0	0.57	GAL 96		
27712162	**Acceptable identification**					**27712162**
Actinomyces naeslundii		91.38	0.53	GAL 96		
Propionibacterium propionicum		8.62	0.26			
27712163	**Very good identification**					**27712163**
Actinomyces naeslundii		100.0	0.57	GAL 96		
27712301	**Good identification**					**27712301**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27712321	**Good identification**					**27712321**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27712323	**Good identification**					**27712323**
Actinomyces naeslundii		100.0	0.35	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27712501	**Good identification**					**27712501**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27712521	**Good identification**					**27712521**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27712523	**Good identification**					**27712523**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27713000	**Very good identification**					**27713000**
Actinomyces naeslundii		99.11	0.82			
Actinomyces israelii		0.29	0.30			
Actinomyces viscosus		0.26	0.14			
27713001	**Very good identification**					**27713001**
Actinomyces naeslundii		99.94	0.85			
Actinomyces viscosus		0.06	0.06			
27713002	**Low discrimination**					**27713002**
Actinomyces naeslundii		78.79	0.82			
Actinomyces viscosus		20.65	0.48			
Actinomyces israelii		0.35	0.33			
27713003	**Good identification**					**27713003**
Actinomyces naeslundii		94.50	0.85			
Actinomyces viscosus		5.50	0.40			
27713010	**Very good identification**					**27713010**
Actinomyces naeslundii		99.18	0.60	MLZ 5		
Actinomyces israelii		0.82	0.15			
27713011	**Very good identification**					**27713011**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27713012	**Good identification**					**27713012**
Actinomyces naeslundii		94.10	0.60	MLZ 5		
Actinomyces viscosus		4.73	0.14			
Actinomyces israelii		1.17	0.19			
27713013	**Very good identification**					**27713013**
Actinomyces naeslundii		98.89	0.63	MLZ 5		
Actinomyces viscosus		1.11	0.06			
27713020	**Very good identification**					**27713020**
Actinomyces naeslundii		99.75	0.82			
Clostridium perfringens		0.25	0.15			
27713021	**Very good identification**					**27713021**
Actinomyces naeslundii		100.0	0.85			
27713022	**Very good identification**					**27713022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.62	0.82			
Actinomyces viscosus		0.26	0.14			
Clostridium perfringens		0.12	0.10			
27713023	**Very good identification**					**27713023**
Actinomyces naeslundii		99.94	0.85			
Actinomyces viscosus		0.06	0.06			
27713030	**Very good identification**					**27713030**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27713031	**Very good identification**					**27713031**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27713032	**Very good identification**					**27713032**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27713033	**Very good identification**					**27713033**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27713040	**Very good identification**					**27713040**
Actinomyces naeslundii		98.63	0.69			
Propionibacterium propionicum		0.74	0.23			
Actinomyces israelii		0.63	0.23			
27713041	**Very good identification**					**27713041**
Actinomyces naeslundii		100.0	0.72			
27713042	**Very good identification**					**27713042**
Actinomyces naeslundii		96.54	0.69			
Actinomyces viscosus		1.45	0.14			
Propionibacterium propionicum		1.08	0.26			
27713043	**Very good identification**					**27713043**
Actinomyces naeslundii		99.67	0.72			
Actinomyces viscosus		0.33	0.06			
27713050	**Good identification**					**27713050**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
27713051	**Good identification**					**27713051**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27713052	**Good identification**					**27713052**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
27713053	**Good identification**					**27713053**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27713060	**Very good identification**					**27713060**
Actinomyces naeslundii		99.74	0.69			
Clostridium perfringens		0.27	0.03			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27713061	**Very good identification**					**27713061**
Actinomyces naeslundii		100.0	0.72			
27713062	**Very good identification**					**27713062**
Actinomyces naeslundii		100.0	0.69			
27713063	**Very good identification**					**27713063**
Actinomyces naeslundii		100.0	0.72			
27713070	**Good identification**					**27713070**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
27713071	**Good identification**					**27713071**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27713072	**Good identification**					**27713072**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
27713073	**Good identification**					**27713073**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27713100	**Very good identification**					**27713100**
Actinomyces naeslundii		95.06	0.90			
Propionibacterium propionicum		4.35	0.57			
Actinomyces israelii		0.39	0.40			
27713101	**Excellent identification**					**27713101**
Actinomyces naeslundii		99.63	0.93			
Propionibacterium propionicum		0.34	0.41			
Actinomyces viscosus		0.04	0.10			
27713102	**Low discrimination**					**27713102**
Actinomyces naeslundii		80.70	0.90			
Actinomyces viscosus		13.21	0.52			
Propionibacterium propionicum		5.54	0.60			
27713103	**Very good identification**					**27713103**
Actinomyces naeslundii		96.02	0.93			
Actinomyces viscosus		3.49	0.44			
Propionibacterium propionicum		0.49	0.44			
27713110	**Very good identification**					**27713110**
Actinomyces naeslundii		98.00	0.68	MLZ 5		
Actinomyces israelii		1.14	0.26			
Propionibacterium propionicum		0.86	0.23			
27713111	**Very good identification**					**27713111**
Actinomyces naeslundii		100.0	0.71	MLZ 5		
27713112	**Good identification**					**27713112**
Actinomyces naeslundii		93.04	0.68	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		2.92	0.18			
Actinomyces israelii		1.62	0.29			
27713113	**Very good identification**					**27713113**
Actinomyces naeslundii		99.31	0.71	MLZ 5		
Actinomyces viscosus		0.69	0.10			
27713120	**Very good identification**					**27713120**
Actinomyces naeslundii		99.28	0.90			
Clostridium perfringens		0.63	0.30			
Propionibacterium propionicum		0.05	0.23			
27713121	**Excellent identification**					**27713121**
Actinomyces naeslundii		100.0	0.93			
27713122	**Very good identification**					**27713122**
Actinomyces naeslundii		99.31	0.90			
Clostridium perfringens		0.30	0.24			
Actinomyces viscosus		0.16	0.18			
27713123	**Excellent identification**					**27713123**
Actinomyces naeslundii		99.96	0.93			
Actinomyces viscosus		0.04	0.10			
27713130	**Very good identification**					**27713130**
Actinomyces naeslundii		100.0	0.68	MLZ 5		
27713131	**Very good identification**					**27713131**
Actinomyces naeslundii		100.0	0.71	MLZ 5		
27713132	**Very good identification**					**27713132**
Actinomyces naeslundii		100.0	0.68	MLZ 5		
27713133	**Very good identification**					**27713133**
Actinomyces naeslundii		100.0	0.71	MLZ 5		
27713140	**Low discrimination**					**27713140**
Actinomyces naeslundii		78.84	0.77			
Propionibacterium propionicum		20.45	0.57			
Actinomyces israelii		0.71	0.33			
27713141	**Very good identification**					**27713141**
Actinomyces naeslundii		98.12	0.80			
Propionibacterium propionicum		1.89	0.41			
27713142	**Low discrimination**					**27713142**
Actinomyces naeslundii		70.83	0.77			
Propionibacterium propionicum		27.55	0.60			
Actinomyces israelii		0.96	0.36			
27713143	**Very good identification**					**27713143**
Actinomyces naeslundii		97.00	0.80			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		2.80	0.44			
Actinomyces viscosus		0.20	0.10			
27713150	**Acceptable identification**					**27713150**
Actinomyces naeslundii		92.99	0.55	MLZ 5		
Propionibacterium propionicum		4.63	0.23			
Actinomyces israelii		2.38	0.19			
27713151	**Very good identification**					**27713151**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27713152	**Low discrimination**					**27713152**
Actinomyces naeslundii		89.84	0.55	MLZ 5		
Propionibacterium propionicum		6.71	0.26			
Actinomyces israelii		3.45	0.22			
27713153	**Very good identification**					**27713153**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27713160	**Very good identification**					**27713160**
Actinomyces naeslundii		99.06	0.77			
Clostridium perfringens		0.68	0.18			
Propionibacterium propionicum		0.26	0.23			
27713161	**Very good identification**					**27713161**
Actinomyces naeslundii		100.0	0.80			
27713162	**Very good identification**					**27713162**
Actinomyces naeslundii		99.21	0.77			
Propionibacterium propionicum		0.39	0.26			
Clostridium perfringens		0.32	0.12			
27713163	**Very good identification**					**27713163**
Actinomyces naeslundii		100.0	0.80			
27713170	**Good identification**					**27713170**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27713171	**Very good identification**					**27713171**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27713172	**Good identification**					**27713172**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27713173	**Very good identification**					**27713173**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27713200	**Low discrimination**					**27713200**
Actinomyces naeslundii		81.50	0.48	RHA 1		
Actinomyces odontolyticus		18.50	0.20			
27713201	**Good identification**					**27713201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	RHA 1		
27713202	**Low discrimination**					**27713202**
Actinomyces naeslundii		69.74	0.48	RHA 1		
Actinomyces viscosus		18.28	0.14			
Actinomyces odontolyticus		9.70	0.16			
27713203	**Acceptable identification**					**27713203**
Actinomyces naeslundii		94.50	0.51	RHA 1		
Actinomyces viscosus		5.51	0.06			
27713220	**Good identification**					**27713220**
Actinomyces naeslundii		96.44	0.48	RHA 1		
Actinomyces odontolyticus		3.56	0.06			
27713221	**Good identification**					**27713221**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27713222	**Good identification**					**27713222**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27713223	**Good identification**					**27713223**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27713240	**Good identification**					**27713240**
Actinomyces naeslundii		100.0	0.35	RHA 1		
27713241	**Good identification**					**27713241**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27713242	**Good identification**					**27713242**
Actinomyces naeslundii		100.0	0.35	RHA 1		
27713243	**Good identification**					**27713243**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27713260	**Good identification**					**27713260**
Actinomyces naeslundii		100.0	0.35	RHA 1		
27713261	**Good identification**					**27713261**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27713262	**Good identification**					**27713262**
Actinomyces naeslundii		100.0	0.35	RHA 1		
27713263	**Good identification**					**27713263**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27713300	**Acceptable identification**					**27713300**
Actinomyces naeslundii		90.79	0.56	RHA 1		
Propionibacterium propionicum		4.16	0.23			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		2.77	0.21				
27713301	**Very good identification**						**27713301**
Actinomyces naeslundii		100.0	0.59	RHA 1			
27713302	**Low discrimination**						**27713302**
Actinomyces naeslundii		77.30	0.56	RHA 1			
Actinomyces viscosus		12.66	0.18				
Propionibacterium propionicum		5.31	0.26				
27713303	**Very good identification**						**27713303**
Actinomyces naeslundii		96.49	0.59	RHA 1			
Actinomyces viscosus		3.51	0.10				
27713311	**Good identification**						**27713311**
Actinomyces naeslundii		100.0	0.37	RHA 1			
27713313	**Good identification**						**27713313**
Actinomyces naeslundii		100.0	0.37	RHA 1			
27713320	**Good identification**						**27713320**
Actinomyces naeslundii		100.0	0.56	RHA 1			
27713321	**Very good identification**						**27713321**
Actinomyces naeslundii		100.0	0.59	RHA 1			
27713322	**Acceptable identification**						**27713322**
					SPO	GRA	
Actinomyces naeslundii		93.76	0.56	RHA 1	1	99	
Clostridium ramosum		3.77	0.07		92	50	
Prevotella oralis		2.47	0.03		1	1	
27713323	**Very good identification**						**27713323**
Actinomyces naeslundii		100.0	0.59	RHA 1			
27713331	**Good identification**						**27713331**
Actinomyces naeslundii		100.0	0.37	RHA 1			
27713333	**Good identification**						**27713333**
Actinomyces naeslundii		100.0	0.37	RHA 1			
27713340	**Low discrimination**						**27713340**
Actinomyces naeslundii		75.38	0.43	RHA 1			
Propionibacterium propionicum		19.55	0.23				
Actinomyces israelii		5.07	0.14				
27713341	**Good identification**						**27713341**
Actinomyces naeslundii		100.0	0.46	RHA 1			
27713342	**Low discrimination**						**27713342**
Actinomyces naeslundii		67.12	0.43	RHA 1			
Propionibacterium propionicum		26.11	0.26				
Actinomyces israelii		6.77	0.17				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27713343	**Good identification**					**27713343**
Actinomyces naeslundii		100.0	0.46	RHA 1		
27713360	**Good identification**					**27713360**
Actinomyces naeslundii		100.0	0.43	RHA 1		
27713361	**Good identification**					**27713361**
Actinomyces naeslundii		100.0	0.46	RHA 1		
27713362	**Good identification**					**27713362**
Actinomyces naeslundii		100.0	0.43	RHA 1		
27713363	**Good identification**					**27713363**
Actinomyces naeslundii		100.0	0.46	RHA 1		
27713400	**Low discrimination**					**27713400**
Actinomyces naeslundii		80.90	0.48	ARA 1		
Actinomyces israelii		19.10	0.28			
27713401	**Good identification**					**27713401**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27713402	**Low discrimination**					**27713402**
Actinomyces naeslundii		61.87	0.48	ARA 1		
Actinomyces israelii		21.91	0.31			
Actinomyces viscosus		16.22	0.14			
27713403	**Acceptable identification**					**27713403**
Actinomyces naeslundii		94.50	0.51	ARA 1		
Actinomyces viscosus		5.51	0.06			
27713420	**Good identification**					**27713420**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27713421	**Good identification**					**27713421**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27713422	**Good identification**					**27713422**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27713423	**Good identification**					**27713423**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27713440	**Low discrimination**					**27713440**
Actinomyces naeslundii		65.77	0.35	ARA 1		
Actinomyces israelii		34.23	0.21			
27713441	**Good identification**					**27713441**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27713442	**Low discrimination**					**27713442**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		56.16	0.35	ARA 1		
Actinomyces israelii		43.84	0.24			
27713443	**Good identification**					**27713443**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27713460	**Good identification**					**27713460**
Actinomyces naeslundii		100.0	0.35	ARA 1		
27713461	**Good identification**					**27713461**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27713462	**Good identification**					**27713462**
Actinomyces naeslundii		100.0	0.35	ARA 1		
27713463	**Good identification**					**27713463**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27713500	**Low discrimination**					**27713500**
Actinomyces naeslundii		72.59	0.56	ARA 1		
Actinomyces israelii		24.09	0.38			
Propionibacterium propionicum		3.32	0.23			
27713501	**Very good identification**					**27713501**
Actinomyces naeslundii		100.0	0.59	ARA 1		
27713502	**Low discrimination**					**27713502**
Actinomyces naeslundii		47.73	0.56	ARA 1		
Actinomyces israelii		23.76	0.41			
Bifidobacterium adolescentis		17.42	0.33			
27713503	**Very good identification**					**27713503**
Actinomyces naeslundii		96.49	0.59	ARA 1		
Actinomyces viscosus		3.51	0.10			
27713511	**Good identification**					**27713511**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27713513	**Good identification**					**27713513**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27713520	**Good identification**					**27713520**
Actinomyces naeslundii		97.93	0.56	ARA 1		
Actinomyces israelii		2.08	0.18			
27713521	**Very good identification**					**27713521**
Actinomyces naeslundii		100.0	0.59	ARA 1		
27713522	**Good identification**					**27713522**
Actinomyces naeslundii		96.92	0.56	ARA 1		
Actinomyces israelii		3.08	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27713523	**Very good identification**					**27713523**
Actinomyces naeslundii		100.0	0.59	ARA 1		
27713531	**Good identification**					**27713531**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27713533	**Good identification**					**27713533**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27713540	**Low discrimination**					**27713540**
Actinomyces naeslundii		50.23	0.43	ARA 1		
Actinomyces israelii		36.74	0.31			
Propionibacterium propionicum		13.03	0.23			
27713541	**Good identification**					**27713541**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27713543	**Good identification**					**27713543**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27713560	**Good identification**					**27713560**
Actinomyces naeslundii		100.0	0.43	ARA 1		
27713561	**Good identification**					**27713561**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27713562	**Acceptable identification**					**27713562**
Actinomyces naeslundii		93.46	0.43	ARA 1		
Actinomyces israelii		6.54	0.14			
27713563	**Good identification**					**27713563**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27714000	**Acceptable identification**					**27714000**
					GRA	
Actinomyces odontolyticus		94.27	0.47	MNS 9	99	
Actinomyces viscosus		4.71	0.18	RAF 3	99	
Fusobacterium mortiferum		1.02	0.19		1	
27714002	**Low discrimination**					**27714002**
Actinomyces viscosus		88.98	0.52	XYL 1		
Actinomyces odontolyticus		11.02	0.43			
27714003	**Good identification**					**27714003**
Actinomyces viscosus		99.40	0.44	XYL 1		
Actinomyces odontolyticus		0.37	0.09			
Actinomyces naeslundii		0.23	0.23			
27714100	**Low discrimination**					**27714100**
					GRA	
Actinomyces odontolyticus		72.25	0.38	MNS 9	99	
Actinomyces viscosus		20.34	0.22	RAF 3	99	
Fusobacterium mortiferum		5.78	0.25		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27714102	**Good identification**					**27714102**
Actinomyces viscosus		97.73	0.56	XYL 1		
Actinomyces odontolyticus		2.15	0.35			
Actinomyces naeslundii		0.08	0.27			
27714103	**Good identification**					**27714103**
Actinomyces viscosus		99.64	0.48	XYL 1		
Actinomyces naeslundii		0.36	0.30			
27714200	**Good identification**					**27714200**
Actinomyces odontolyticus		100.0	0.52	MNS 9		
27714202	**Good identification**					**27714202**
Actinomyces odontolyticus		95.97	0.48	MNS 9		
Actinomyces viscosus		4.03	0.18	RAF 3		
27714220	**Good identification**					**27714220**
Actinomyces odontolyticus		100.0	0.38	MNS 9		
27714300	**Good identification**					**27714300**
Actinomyces odontolyticus		100.0	0.43	MNS 9		
27714302	**Low discrimination**					**27714302**
Actinomyces odontolyticus		80.87	0.40	MNS 9		
Actinomyces viscosus		19.14	0.22	RAF 3		
27715000	**Low discrimination**					**27715000**
Actinomyces odontolyticus		78.59	0.44	MNS 9		
Actinomyces israelii		8.07	0.35	RAF 3		
Actinomyces naeslundii		7.10	0.43			
27715001	**Low discrimination**					**27715001**
Actinomyces naeslundii		84.20	0.46	XYL 1		
Actinomyces viscosus		9.52	0.06			
Actinomyces odontolyticus		6.28	0.10			
27715002	**Low discrimination**					**27715002**
Actinomyces viscosus		84.01	0.48	XYL 1		
Actinomyces odontolyticus		11.31	0.40			
Actinomyces israelii		2.84	0.38			
27715003	**Acceptable identification**					**27715003**
Actinomyces viscosus		91.46	0.40	XYL 1		
Actinomyces naeslundii		8.17	0.46			
Actinomyces odontolyticus		0.37	0.06			
27715021	**Good identification**					**27715021**
Actinomyces naeslundii		100.0	0.46	XYL 1		
27715023	**Low discrimination**					**27715023**
Actinomyces naeslundii		89.84	0.46	XYL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		10.16	0.06			
27715100	**Low discrimination**					**27715100**
Actinomyces israelii		34.49	0.45	CEL 93		
Actinomyces odontolyticus		26.51	0.35			
Actinomyces naeslundii		21.58	0.51			
27715101	**Acceptable identification**					**27715101**
Actinomyces naeslundii		93.40	0.54	XYL 1		
Actinomyces viscosus		6.60	0.10			
27715102	**Low discrimination**					**27715102**
Actinomyces viscosus		81.82	0.52	XYL 1		
Actinomyces israelii		6.23	0.48			
Bifidobacterium longum subsp. infantis		4.56	0.26			
27715103	**Low discrimination**					**27715103**
Actinomyces viscosus		87.31	0.44	XYL 1		
Actinomyces naeslundii		12.49	0.54			
Actinomyces israelii		0.20	0.14			
27715112	**Low discrimination**					**27715112**
Bifidobacterium adolescentis		59.23	0.36	NIT 1		
Actinomyces israelii		20.04	0.34	ARA 87		
Actinomyces viscosus		17.78	0.18			
27715121	**Good identification**					**27715121**
Actinomyces naeslundii		100.0	0.54	XYL 1		
27715123	**Acceptable identification**					**27715123**
Actinomyces naeslundii		91.61	0.54	XYL 1		
Actinomyces viscosus		6.47	0.10			
Bifidobacterium longum subsp. infantis		1.92	-.04			
27715132	**Low discrimination**					**27715132**
					GRA	
Parabacteroides distasonis		80.91	0.36	NIT 1	1	
Capnocytophaga ochracea		7.15	0.05		1	
Actinomyces naeslundii		5.87	0.29		99	
27715140	**Low discrimination**					**27715140**
Actinomyces israelii		77.88	0.38	CEL 93		
Actinomyces naeslundii		22.12	0.38			
27715141	**Good identification**					**27715141**
Actinomyces naeslundii		100.0	0.41	XYL 1		
27715142	**Low discrimination**					**27715142**
Actinomyces israelii		60.79	0.41	CEL 93		
Actinomyces viscosus		20.73	0.18			
Actinomyces naeslundii		11.51	0.38			
27715143	**Low discrimination**					**27715143**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		71.42	0.41	XYL 1			
Actinomyces viscosus		28.58	0.10				
27715161	**Good identification**						**27715161**
Actinomyces naeslundii		100.0	0.41	XYL 1			
27715162	**Low discrimination**						**27715162**
					GRA	CAT	
Actinomyces naeslundii		40.03	0.38	XYL 1	99	25	
Clostridium perfringens		24.78	0.12		99	1	
Capnocytophaga ochracea		14.54	0.05		1	90	
27715163	**Good identification**						**27715163**
Actinomyces naeslundii		100.0	0.41	XYL 1			
27715200	**Good identification**						**27715200**
Actinomyces odontolyticus		99.60	0.49	MNS 9			
Actinomyces israelii		0.40	0.16	RAF 3			
27715202	**Good identification**						**27715202**
Actinomyces odontolyticus		95.38	0.45	MNS 9			
Actinomyces viscosus		3.69	0.14	RAF 3			
Actinomyces israelii		0.93	0.19				
27715220	**Good identification**						**27715220**
					GRA		
Actinomyces odontolyticus		96.74	0.35	MNS 9	99		
Bacteroides vulgatus		3.26	-.04	RAF 3	1		
27715300	**Low discrimination**						**27715300**
Actinomyces odontolyticus		73.36	0.40	MNS 9			
Eubacterium contortum		21.67	0.24	RAF 3			
Actinomyces israelii		3.70	0.26				
27715302	**Low discrimination**						**27715302**
Actinomyces odontolyticus		67.73	0.37	MNS 9			
Actinomyces viscosus		14.73	0.18	RAF 3			
Actinomyces israelii		8.36	0.29				
27715322	**Low discrimination**						**27715322**
					GRA		
Parabacteroides distasonis		80.82	0.48	NIT 1	1		
Prevotella oralis		12.52	0.22		1		
Actinomyces odontolyticus		4.76	0.23		99		
27715332	**Good identification**						**27715332**
Parabacteroides distasonis		96.82	0.44	NIT 1			
Clostridium clostridioforme		3.18	0.13				
27715402	**Low discrimination**						**27715402**
Actinomyces israelii		62.31	0.36	CEL 93			
Actinomyces viscosus		22.72	0.14				
Bifidobacterium adolescentis		7.44	0.14				
27715500	**Low discrimination**						**27715500**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		80.01	0.44	CEL 93	99	
Bifidobacterium longum subsp. longum		6.80	0.17		99	
Bacteroides vulgatus		3.74	-.01		1	
27715502	**Low discrimination**					**27715502**
Bifidobacterium adolescentis		70.03	0.48	NIT 1		
Actinomyces israelii		23.69	0.47			
Actinomyces viscosus		3.84	0.18			
27715512	**Acceptable identification**					**27715512**
Bifidobacterium adolescentis		91.78	0.50	NIT 1		
Actinomyces israelii		3.80	0.33			
Bifidobacterium longum subsp. longum		2.45	0.21			
27715522	**Low discrimination**					**27715522**
					GRA	
Parabacteroides distasonis		49.09	0.37	NIT 1	1	
Bifidobacterium longum subsp. longum		26.42	0.28		99	
Actinomyces israelii		7.45	0.26		99	
27715540	**Good identification**					**27715540**
Actinomyces israelii		100.0	0.37	CEL 93		
27715542	**Low discrimination**					**27715542**
Actinomyces israelii		51.56	0.40	CEL 93		
Bifidobacterium adolescentis		48.44	0.33			
27715720	**Good identification**					**27715720**
					GRA	
Bacteroides vulgatus		99.77	0.49	NIT 1	1	
Eubacterium contortum		0.19	0.10		99	
Parabacteroides distasonis		0.05	0.11		1	
27715722	**Low discrimination**					**27715722**
Parabacteroides distasonis		75.67	0.45	NIT 1		
Bacteroides vulgatus		16.83	0.15			
Clostridium clostridioforme		7.50	0.22			
27715732	**Low discrimination**					**27715732**
Parabacteroides distasonis		54.39	0.40	NIT 1		
Clostridium clostridioforme		45.61	0.33			
27716101	**Good identification**					**27716101**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27716121	**Good identification**					**27716121**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27716123	**Good identification**					**27716123**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27717000	**Low discrimination**					**27717000**
Actinomyces israelii		75.84	0.48	CEL 93		
Actinomyces naeslundii		22.85	0.48			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		1.32	0.10			
27717001	**Good identification**					**27717001**
Actinomyces naeslundii		97.82	0.51	XYL 1		
Actinomyces israelii		2.19	0.14			
27717002	**Low discrimination**					**27717002**
Actinomyces israelii		79.33	0.51	CEL 93		
Actinomyces naeslundii		15.93	0.48			
Actinomyces viscosus		4.18	0.14			
27717003	**Acceptable identification**					**27717003**
Actinomyces naeslundii		91.59	0.51	XYL 1		
Actinomyces viscosus		5.34	0.06			
Actinomyces israelii		3.07	0.17			
27717012	**Acceptable identification**					**27717012**
Actinomyces israelii		93.39	0.37	CEL 93		
Actinomyces naeslundii		6.61	0.26	MLZ 13		
27717020	**Low discrimination**					**27717020**
Actinomyces naeslundii		82.52	0.48	XYL 1		
Actinomyces israelii		17.48	0.27			
27717021	**Good identification**					**27717021**
Actinomyces naeslundii		100.0	0.51	XYL 1		
27717022	**Low discrimination**					**27717022**
Actinomyces naeslundii		75.89	0.48	XYL 1		
Actinomyces israelii		24.11	0.30			
27717023	**Good identification**					**27717023**
Actinomyces naeslundii		100.0	0.51	XYL 1		
27717040	**Low discrimination**					**27717040**
Actinomyces israelii		87.97	0.41	CEL 93		
Actinomyces naeslundii		12.03	0.35			
27717041	**Good identification**					**27717041**
Actinomyces naeslundii		100.0	0.38	XYL 1		
27717042	**Acceptable identification**					**27717042**
Actinomyces israelii		91.65	0.44	CEL 93		
Actinomyces naeslundii		8.35	0.35			
27717043	**Good identification**					**27717043**
Actinomyces naeslundii		100.0	0.38	XYL 1		
27717060	**Low discrimination**					**27717060**
Actinomyces naeslundii		68.17	0.35	XYL 1		
Actinomyces israelii		31.83	0.20			
27717061	**Good identification**					**27717061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	XYL 1		
27717062	**Low discrimination**					**27717062**
Actinomyces naeslundii		58.81	0.35	XYL 1		
Actinomyces israelii		41.19	0.23			
27717063	**Good identification**					**27717063**
Actinomyces naeslundii		100.0	0.38	XYL 1		
27717100	**Low discrimination**					**27717100**
Actinomyces israelii		81.69	0.58	CEL 93		
Actinomyces naeslundii		17.51	0.56			
Propionibacterium propionicum		0.80	0.23			
27717101	**Very good identification**					**27717101**
Actinomyces naeslundii		96.96	0.59	XYL 1		
Actinomyces israelii		3.05	0.24			
27717102	**Low discrimination**					**27717102**
Actinomyces israelii		81.41	0.61	CEL 93		
Actinomyces naeslundii		11.64	0.56			
Bifidobacterium adolescentis		4.25	0.33			
27717103	**Good identification**					**27717103**
Actinomyces naeslundii		92.29	0.59	XYL 1		
Actinomyces israelii		4.35	0.27			
Actinomyces viscosus		3.36	0.10			
27717110	**Acceptable identification**					**27717110**
Actinomyces israelii		92.98	0.44	CEL 93		
Actinomyces naeslundii		7.02	0.34	MLZ 13		
27717111	**Good identification**					**27717111**
Actinomyces naeslundii		100.0	0.37	XYL 1		
27717112	**Low discrimination**					**27717112**
Actinomyces israelii		67.70	0.47	CEL 93		
Bifidobacterium adolescentis		28.89	0.34	MLZ 13		
Actinomyces naeslundii		3.41	0.34			
27717113	**Good identification**					**27717113**
Actinomyces naeslundii		100.0	0.37	XYL 1		
27717120	**Low discrimination**					**27717120**
Actinomyces naeslundii		77.06	0.56	XYL 1		
Actinomyces israelii		22.94	0.38			
27717121	**Very good identification**					**27717121**
Actinomyces naeslundii		100.0	0.59	XYL 1		
27717122	**Low discrimination**					**27717122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		68.36	0.56	XYL 1		
Actinomyces israelii		30.53	0.41			
Bifidobacterium longum subsp. infantis		1.11	-.04			
27717123	**Very good identification**					**27717123**
Actinomyces naeslundii		100.0	0.59	XYL 1		
27717131	**Good identification**					**27717131**
Actinomyces naeslundii		100.0	0.37	XYL 1		
27717133	**Good identification**					**27717133**
Actinomyces naeslundii		100.0	0.37	XYL 1		
27717140	**Low discrimination**					**27717140**
Actinomyces israelii		89.09	0.51	CEL 93		
Actinomyces naeslundii		8.67	0.43			
Propionibacterium propionicum		2.25	0.23			
27717141	**Acceptable identification**					**27717141**
Actinomyces naeslundii		93.53	0.46	XYL 1		
Actinomyces israelii		6.47	0.17			
27717142	**Acceptable identification**					**27717142**
Actinomyces israelii		90.36	0.54	CEL 93		
Actinomyces naeslundii		5.86	0.43			
Propionibacterium propionicum		2.28	0.26			
27717143	**Acceptable identification**					**27717143**
Actinomyces naeslundii		90.59	0.46	XYL 1		
Actinomyces israelii		9.41	0.20			
27717150	**Good identification**					**27717150**
Actinomyces israelii		100.0	0.37	CEL 93		
27717152	**Low discrimination**					**27717152**
Actinomyces israelii		88.06	0.40	CEL 93		
Bifidobacterium adolescentis		11.94	0.19	MLZ 13		
27717160	**Low discrimination**					**27717160**
Actinomyces naeslundii		60.38	0.43	XYL 1		
Actinomyces israelii		39.62	0.31			
27717161	**Good identification**					**27717161**
Actinomyces naeslundii		100.0	0.46	XYL 1		
27717162	**Low discrimination**					**27717162**
Actinomyces naeslundii		50.40	0.43	XYL 1		
Actinomyces israelii		49.60	0.34			
27717163	**Good identification**					**27717163**
Actinomyces naeslundii		100.0	0.46	XYL 1		
27717300	**Good identification**					**27717300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		96.59	0.39	CEL 93		
Actinomyces odontolyticus		3.41	0.06	RHA 7		
27717302	**Good identification**					**27717302**
Actinomyces israelii		100.0	0.42	CEL 93		
27717400	**Good identification**					**27717400**
Actinomyces israelii		100.0	0.46	CEL 93		
27717402	**Good identification**					**27717402**
Actinomyces israelii		98.31	0.49	CEL 93		
Bifidobacterium adolescentis		1.70	0.13			
27717412	**Low discrimination**					**27717412**
Actinomyces israelii		87.64	0.35	CEL 93		
Bifidobacterium adolescentis		12.36	0.14	MLZ 13		
27717440	**Good identification**					**27717440**
Actinomyces israelii		100.0	0.39	CEL 93		
27717442	**Good identification**					**27717442**
Actinomyces israelii		100.0	0.42	CEL 93		
27717500	**Very good identification**					**27717500**
Actinomyces israelii		99.36	0.57	CEL 93		
Bifidobacterium adolescentis		0.64	0.13			
27717502	**Low discrimination**					**27717502**
Actinomyces israelii		70.09	0.60	CEL 93		
Bifidobacterium adolescentis		29.91	0.47			
27717510	**Acceptable identification**					**27717510**
Actinomyces israelii		94.98	0.42	CEL 93		
Bifidobacterium adolescentis		5.02	0.14	MLZ 13		
27717512	**Low discrimination**					**27717512**
Bifidobacterium adolescentis		77.73	0.48	NIT 1		
Actinomyces israelii		22.27	0.45			
27717520	**Low discrimination**					**27717520**
					GRA	
Actinomyces israelii		88.55	0.36	CEL 93	99	
Bacteroides vulgatus		11.46	-.01	NAG 6	1	
27717522	**Acceptable identification**					**27717522**
Actinomyces israelii		93.67	0.39	CEL 93		
Bifidobacterium adolescentis		6.33	0.13	NAG 6		
27717540	**Good identification**					**27717540**
Actinomyces israelii		100.0	0.50	CEL 93		
27717542	**Low discrimination**					**27717542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		88.06	0.53	CEL 93		
Bifidobacterium adolescentis		11.94	0.31			
27717550	**Good identification**					**27717550**
Actinomyces israelii		100.0	0.35	CEL 93		
27717700	**Good identification**					**27717700**
Actinomyces israelii		100.0	0.37	CEL 93		
27717702	**Acceptable identification**					**27717702**
Actinomyces israelii		94.58	0.40	CEL 93		
Bifidobacterium adolescentis		5.42	0.13	RHA 7		
27720000	**Low discrimination**					**27720000**
Actinomyces naeslundii		56.42	0.37	FRU 95		
Actinomyces viscosus		43.58	0.11	GAL 96		
27720001	**Low discrimination**					**27720001**
Actinomyces naeslundii		85.35	0.40	FRU 95		
Actinomyces viscosus		14.65	0.02	GAL 96		
27720002	**Good identification**					**27720002**
Actinomyces viscosus		98.71	0.45	FRU 95		
Actinomyces naeslundii		1.29	0.37	TRE 7		
27720003	**Acceptable identification**					**27720003**
Actinomyces viscosus		94.44	0.37	FRU 95		
Actinomyces naeslundii		5.56	0.40	TRE 7		
27720020	**Good identification**					**27720020**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27720021	**Good identification**					**27720021**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27720022	**Low discrimination**					**27720022**
Actinomyces naeslundii		56.42	0.37	FRU 95		
Actinomyces viscosus		43.58	0.11	GAL 96		
27720023	**Low discrimination**					**27720023**
Actinomyces naeslundii		85.35	0.40	FRU 95		
Actinomyces viscosus		14.65	0.02	GAL 96		
27720100	**Low discrimination**					**27720100**
Actinomyces naeslundii		67.45	0.44	FRU 95		
Actinomyces viscosus		32.55	0.15	GAL 96		
27720101	**Acceptable identification**					**27720101**
Actinomyces naeslundii		90.32	0.48	FRU 95		
Actinomyces viscosus		9.68	0.07	GAL 96		
27720102	**Good identification**					**27720102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		97.95	0.49	FRU 95		
Actinomyces naeslundii		2.05	0.44	TRE 7		
27720103	**Acceptable identification**					**27720103**
Actinomyces viscosus		91.39	0.41	FRU 95		
Actinomyces naeslundii		8.61	0.48	TRE 7		
27720120	**Acceptable identification**					**27720120**
Actinomyces naeslundii		94.86	0.44	FRU 95		
Clostridium perfringens		5.15	0.01	GAL 96		
27720121	**Good identification**					**27720121**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27720122	**Low discrimination**					**27720122**
Actinomyces naeslundii		67.45	0.44	FRU 95		
Actinomyces viscosus		32.55	0.15	GAL 96		
27720123	**Acceptable identification**					**27720123**
Actinomyces naeslundii		90.32	0.48	FRU 95		
Actinomyces viscosus		9.68	0.07	GAL 96		
27721000	**Very good identification**					**27721000**
Actinomyces naeslundii		98.07	0.60	FRU 95		
Actinomyces viscosus		1.94	0.07			
27721001	**Very good identification**					**27721001**
Actinomyces naeslundii		99.56	0.63	FRU 95		
Actinomyces viscosus		0.44	-.01			
27721002	**Low discrimination**					**27721002**
Actinomyces viscosus		66.14	0.41	FRU 95		
Actinomyces naeslundii		33.86	0.60	TRE 7		
27721003	**Low discrimination**					**27721003**
Actinomyces naeslundii		69.73	0.63	FRU 95		
Actinomyces viscosus		30.27	0.33			
27721010	**Good identification**					**27721010**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27721011	**Good identification**					**27721011**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27721012	**Low discrimination**					**27721012**
Actinomyces naeslundii		72.73	0.38	FRU 95		
Actinomyces viscosus		27.27	0.07	MLZ 5		
27721013	**Acceptable identification**					**27721013**
Actinomyces naeslundii		92.31	0.41	FRU 95		
Actinomyces viscosus		7.69	-.01	MLZ 5		
27721020	**Good identification**					**27721020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.01	0.60	FRU 95		
Clostridium perfringens		7.99	0.20			
27721021	**Very good identification**					**27721021**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27721022	**Good identification**					**27721022**
Actinomyces naeslundii		94.29	0.60	FRU 95		
Clostridium perfringens		3.85	0.14			
Actinomyces viscosus		1.86	0.07			
27721023	**Very good identification**					**27721023**
Actinomyces naeslundii		99.56	0.63	FRU 95		
Actinomyces viscosus		0.44	-.01			
27721030	**Good identification**					**27721030**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27721031	**Good identification**					**27721031**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27721032	**Good identification**					**27721032**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27721033	**Good identification**					**27721033**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27721040	**Good identification**					**27721040**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27721041	**Good identification**					**27721041**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27721042	**Low discrimination**					**27721042**
Actinomyces naeslundii		89.94	0.47	FRU 95		
Actinomyces viscosus		10.06	0.07			
27721043	**Good identification**					**27721043**
Actinomyces naeslundii		97.58	0.50	FRU 95		
Actinomyces viscosus		2.43	-.01			
27721060	**Acceptable identification**					**27721060**
Actinomyces naeslundii		91.43	0.47	FRU 95		
Clostridium perfringens		8.57	0.08			
27721061	**Good identification**					**27721061**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27721062	**Good identification**					**27721062**
Actinomyces naeslundii		95.77	0.47	FRU 95		
Clostridium perfringens		4.23	0.02			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27721063	**Good identification**					**27721063**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27721100	**Very good identification**					**27721100**
Actinomyces naeslundii		98.11	0.68	FRU 95		
Actinomyces viscosus		1.21	0.11			
Clostridium perfringens		0.68	0.09			
27721101	**Very good identification**					**27721101**
					SPO	
Actinomyces naeslundii		99.38	0.71	FRU 95	1	
Clostridium chauvoei		0.35	0.06		99	
Actinomyces viscosus		0.27	0.03		1	
27721102	**Low discrimination**					**27721102**
Actinomyces viscosus		54.88	0.45	FRU 95		
Actinomyces naeslundii		44.98	0.68	TRE 7		
Clostridium perfringens		0.15	0.03			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27721103	**Low discrimination**					**27721103**
Actinomyces naeslundii		78.67	0.71	FRU 95		
Actinomyces viscosus		21.33	0.37			
27721110	**Good identification**					**27721110**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27721111	**Good identification**					**27721111**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27721112	**Low discrimination**					**27721112**
Actinomyces naeslundii		81.03	0.46	FRU 95		
Actinomyces viscosus		18.97	0.11	MLZ 5		
27721113	**Good identification**					**27721113**
Actinomyces naeslundii		95.05	0.49	FRU 95		
Actinomyces viscosus		4.95	0.03	MLZ 5		
27721120	**Low discrimination**					**27721120**
Actinomyces naeslundii		81.71	0.68	FRU 95		
Clostridium perfringens		18.29	0.35			
27721121	**Very good identification**					**27721121**
Actinomyces naeslundii		99.85	0.71	FRU 95		
Clostridium perfringens		0.15	0.01			
27721122	**Low discrimination**					**27721122**
Actinomyces naeslundii		89.47	0.68	FRU 95		
Clostridium perfringens		9.42	0.29			
Actinomyces viscosus		1.10	0.11			
27721123	**Very good identification**					**27721123**
Actinomyces naeslundii		99.73	0.71	FRU 95		
Actinomyces viscosus		0.27	0.03			
27721130	**Good identification**					**27721130**
Actinomyces naeslundii		95.88	0.46	FRU 95		
Clostridium perfringens		4.12	0.01	MLZ 5		
27721131	**Good identification**					**27721131**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27721132	**Good identification**					**27721132**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27721133	**Good identification**					**27721133**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27721140	**Good identification**					**27721140**
Actinomyces naeslundii		100.0	0.55	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27721141	**Very good identification**					**27721141**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27721142	**Acceptable identification**					**27721142**
Actinomyces naeslundii		93.47	0.55	FRU 95		
Actinomyces viscosus		6.53	0.11			
27721143	**Very good identification**					**27721143**
Actinomyces naeslundii		98.47	0.58	FRU 95		
Actinomyces viscosus		1.53	0.03			
27721151	**Good identification**					**27721151**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27721153	**Good identification**					**27721153**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27721160	**Low discrimination**					**27721160**
Actinomyces naeslundii		80.54	0.55	FRU 95		
Clostridium perfringens		19.46	0.22			
27721161	**Very good identification**					**27721161**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27721162	**Low discrimination**					**27721162**
Actinomyces naeslundii		89.79	0.55	FRU 95		
Clostridium perfringens		10.21	0.17			
27721163	**Very good identification**					**27721163**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27721171	**Good identification**					**27721171**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27721173	**Good identification**					**27721173**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27721301	**Good identification**					**27721301**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27721303	**Low discrimination**					**27721303**
Actinomyces naeslundii		78.67	0.37	FRU 95		
Actinomyces viscosus		21.33	0.03	RHA 1		
27721321	**Good identification**					**27721321**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27721323	**Good identification**					**27721323**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27721501	**Good identification**					**27721501**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	FRU 95		
27721503	**Low discrimination**					**27721503**
Actinomyces naeslundii		78.67	0.37	FRU 95		
Actinomyces viscosus		21.33	0.03	ARA 1		
27721521	**Good identification**					**27721521**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27721523	**Good identification**					**27721523**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27722000	**Good identification**					**27722000**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27722001	**Good identification**					**27722001**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27722002	**Low discrimination**					**27722002**
Actinomyces naeslundii		71.53	0.42	FRU 95		
Actinomyces viscosus		28.47	0.11	GAL 96		
27722003	**Acceptable identification**					**27722003**
Actinomyces naeslundii		91.87	0.45	FRU 95		
Actinomyces viscosus		8.13	0.02	GAL 96		
27722020	**Good identification**					**27722020**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27722021	**Good identification**					**27722021**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27722022	**Good identification**					**27722022**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27722023	**Good identification**					**27722023**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27722100	**Acceptable identification**					**27722100**
Actinomyces naeslundii		91.84	0.49	FRU 95		
Propionibacterium propionicum		8.16	0.21	GAL 96		
27722101	**Good identification**					**27722101**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27722102	**Low discrimination**					**27722102**
Actinomyces naeslundii		72.37	0.49	FRU 95		
Actinomyces viscosus		17.99	0.15	GAL 96		
Propionibacterium propionicum		9.64	0.24			
27722103	**Acceptable identification**					**27722103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		94.77	0.52	FRU 95		
Actinomyces viscosus		5.24	0.07	GAL 96		
27722120	**Good identification**					**27722120**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27722121	**Good identification**					**27722121**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27722122	**Good identification**					**27722122**
Actinomyces naeslundii		100.0	0.49	FRU 95		
27722123	**Good identification**					**27722123**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27722140	**Low discrimination**					**27722140**
Actinomyces naeslundii		66.52	0.37	FRU 95		
Propionibacterium propionicum		33.48	0.21	GAL 96		
27722141	**Good identification**					**27722141**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27722142	**Low discrimination**					**27722142**
Actinomyces naeslundii		56.98	0.37	FRU 95		
Propionibacterium propionicum		43.02	0.24	GAL 96		
27722143	**Good identification**					**27722143**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27722160	**Good identification**					**27722160**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27722161	**Good identification**					**27722161**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27722162	**Good identification**					**27722162**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27722163	**Good identification**					**27722163**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27723000	**Very good identification**					**27723000**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27723001	**Very good identification**					**27723001**
Actinomyces naeslundii		100.0	0.68	FRU 95		
27723002	**Very good identification**					**27723002**
Actinomyces naeslundii		98.99	0.65	FRU 95		
Actinomyces viscosus		1.01	0.07			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27723003	**Very good identification**					**27723003**
Actinomyces naeslundii		99.78	0.68	FRU 95		
Actinomyces viscosus		0.23	-.01			
27723010	**Good identification**					**27723010**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27723011	**Good identification**					**27723011**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27723012	**Good identification**					**27723012**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27723013	**Good identification**					**27723013**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27723020	**Very good identification**					**27723020**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27723021	**Very good identification**					**27723021**
Actinomyces naeslundii		100.0	0.68	FRU 95		
27723022	**Very good identification**					**27723022**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27723023	**Very good identification**					**27723023**
Actinomyces naeslundii		100.0	0.68	FRU 95		
27723030	**Good identification**					**27723030**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27723031	**Good identification**					**27723031**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27723032	**Good identification**					**27723032**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27723033	**Good identification**					**27723033**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27723040	**Good identification**					**27723040**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27723041	**Good identification**					**27723041**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27723042	**Good identification**					**27723042**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27723043	**Good identification**					**27723043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	FRU 95		
27723060	**Good identification**					**27723060**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27723061	**Good identification**					**27723061**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27723062	**Good identification**					**27723062**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27723063	**Good identification**					**27723063**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27723100	**Very good identification**					**27723100**
Actinomyces naeslundii		99.48	0.73	FRU 95		
Propionibacterium propionicum		0.37	0.21			
Actinomyces israelii		0.15	0.16			
27723101	**Very good identification**					**27723101**
Actinomyces naeslundii		100.0	0.76	FRU 95		
27723102	**Very good identification**					**27723102**
Actinomyces naeslundii		98.60	0.73	FRU 95		
Actinomyces viscosus		0.63	0.11			
Propionibacterium propionicum		0.55	0.24			
27723103	**Very good identification**					**27723103**
Actinomyces naeslundii		99.86	0.76	FRU 95		
Actinomyces viscosus		0.14	0.03			
27723110	**Good identification**					**27723110**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27723111	**Good identification**					**27723111**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27723112	**Good identification**					**27723112**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27723113	**Good identification**					**27723113**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27723120	**Very good identification**					**27723120**
Actinomyces naeslundii		99.88	0.73	FRU 95		
Clostridium perfringens		0.12	0.01			
27723121	**Very good identification**					**27723121**
Actinomyces naeslundii		100.0	0.76	FRU 95		
27723122	**Very good identification**					**27723122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.73	FRU 95		
27723123	**Very good identification**					**27723123**
Actinomyces naeslundii		100.0	0.76	FRU 95		
27723130	**Good identification**					**27723130**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27723131	**Good identification**					**27723131**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27723132	**Good identification**					**27723132**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27723133	**Good identification**					**27723133**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27723140	**Very good identification**					**27723140**
Actinomyces naeslundii		97.95	0.60	FRU 95		
Propionibacterium propionicum		2.05	0.21			
27723141	**Very good identification**					**27723141**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27723142	**Very good identification**					**27723142**
Actinomyces naeslundii		96.95	0.60	FRU 95		
Propionibacterium propionicum		3.05	0.24			
27723143	**Very good identification**					**27723143**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27723150	**Good identification**					**27723150**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27723151	**Good identification**					**27723151**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27723152	**Good identification**					**27723152**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27723153	**Good identification**					**27723153**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27723160	**Very good identification**					**27723160**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27723161	**Very good identification**					**27723161**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27723162	**Very good identification**					**27723162**
Actinomyces naeslundii		100.0	0.60	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27723163	**Very good identification**					**27723163**
Actinomyces naeslundii		100.0	0.63	FRU 95		
27723170	**Good identification**					**27723170**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27723171	**Good identification**					**27723171**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27723172	**Good identification**					**27723172**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27723173	**Good identification**					**27723173**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27723300	**Good identification**					**27723300**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27723301	**Good identification**					**27723301**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27723302	**Good identification**					**27723302**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27723303	**Good identification**					**27723303**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27723320	**Good identification**					**27723320**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27723321	**Good identification**					**27723321**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27723322	**Good identification**					**27723322**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27723323	**Good identification**					**27723323**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27723500	**Low discrimination**					**27723500**
Actinomyces naeslundii		89.01	0.39	FRU 95		
Actinomyces israelii		10.99	0.14	ARA 1		
27723501	**Good identification**					**27723501**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27723502	**Low discrimination**					**27723502**
Actinomyces naeslundii		84.38	0.39	FRU 95		
Actinomyces israelii		15.62	0.17	ARA 1		
27723503	**Good identification**					**27723503**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	FRU 95		
27723520	**Good identification**					**27723520**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27723521	**Good identification**					**27723521**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27723522	**Good identification**					**27723522**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27723523	**Good identification**					**27723523**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27725101	**Good identification**					**27725101**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27725103	**Low discrimination**					**27725103**
Actinomyces naeslundii		78.67	0.37	FRU 95		
Actinomyces viscosus		21.33	0.03	XYL 1		
27725121	**Good identification**					**27725121**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27725123	**Good identification**					**27725123**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27727101	**Good identification**					**27727101**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27727102	**Low discrimination**					**27727102**
Actinomyces israelii		72.24	0.37	FRU 99		
Actinomyces naeslundii		27.76	0.39	CEL 93		
27727103	**Good identification**					**27727103**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27727120	**Acceptable identification**					**27727120**
Actinomyces naeslundii		90.03	0.39	FRU 95		
Actinomyces israelii		9.97	0.13	XYL 1		
27727121	**Good identification**					**27727121**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27727122	**Low discrimination**					**27727122**
Actinomyces naeslundii		85.75	0.39	FRU 95		
Actinomyces israelii		14.25	0.16	XYL 1		
27727123	**Good identification**					**27727123**
Actinomyces naeslundii		100.0	0.42	FRU 95		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27727502	**Acceptable identification**					**27727502**
Actinomyces israelii		92.97	0.35	FRU 99		
Bifidobacterium adolescentis		7.03	0.10	CEL 93		
27730000	**Low discrimination**					**27730000**
Actinomyces naeslundii		50.17	0.59	GAL 96		
Actinomyces viscosus		38.76	0.33			
Propionibacterium granulosum		4.47	0.26			
27730001	**Low discrimination**					**27730001**
Actinomyces naeslundii		85.35	0.62	GAL 96		
Actinomyces viscosus		14.65	0.24			
27730002	**Very good identification**					**27730002**
Actinomyces viscosus		98.64	0.67	TRE 7		
Actinomyces naeslundii		1.29	0.59			
Propionibacterium granulosum		0.03	0.15			
27730003	**Good identification**					**27730003**
Actinomyces viscosus		94.44	0.59	TRE 7		
Actinomyces naeslundii		5.56	0.62			
27730010	**Low discrimination**					**27730010**
Actinomyces naeslundii		72.50	0.37	GAL 96		
Propionibacterium granulosum		16.75	0.11	MLZ 5		
Actinomyces viscosus		10.75	-.02			
27730011	**Good identification**					**27730011**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27730020	**Low discrimination**					**27730020**
					CAT	
Actinomyces naeslundii		87.73	0.59	GAL 96	25	
Clostridium perfringens		9.63	0.20		1	
Propionibacterium granulosum		1.96	0.15		90	
27730021	**Very good identification**					**27730021**
Actinomyces naeslundii		100.0	0.62	GAL 96		
27730022	**Low discrimination**					**27730022**
Actinomyces naeslundii		54.82	0.59	GAL 96		
Actinomyces viscosus		42.35	0.33			
Clostridium perfringens		2.83	0.14			
27730023	**Low discrimination**					**27730023**
Actinomyces naeslundii		85.35	0.62	GAL 96		
Actinomyces viscosus		14.65	0.24			
27730030	**Good identification**					**27730030**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27730031	**Good identification**					**27730031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	GAL 96		
27730032	**Low discrimination**					**27730032**
Actinomyces naeslundii		87.09	0.37	GAL 96		
Actinomyces viscosus		12.91	-.02	MLZ 5		
27730033	**Good identification**					**27730033**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27730040	**Acceptable identification**					**27730040**
					SPO	
Actinomyces naeslundii		90.37	0.46	GAL 96	1	
Clostridium botulinum biovar B		5.64	-.03		99	
Actinomyces viscosus		4.00	-.02		1	
27730041	**Good identification**					**27730041**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27730043	**Low discrimination**					**27730043**
Actinomyces naeslundii		50.69	0.49	GAL 96		
Actinomyces viscosus		49.31	0.24			
27730060	**Low discrimination**					**27730060**
Actinomyces naeslundii		89.41	0.46	GAL 96		
Clostridium perfringens		10.59	0.08			
27730061	**Good identification**					**27730061**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27730062	**Acceptable identification**					**27730062**
Actinomyces naeslundii		90.91	0.46	GAL 96		
Clostridium perfringens		5.07	0.02			
Actinomyces viscosus		4.02	-.02			
27730063	**Good identification**					**27730063**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27730100	**Low discrimination**					**27730100**
					GRA	
Actinomyces naeslundii		63.37	0.66	GAL 96	99	
Actinomyces viscosus		30.58	0.37		99	
Fusobacterium mortiferum		4.62	0.35		1	
27730101	**Good identification**					**27730101**
Actinomyces naeslundii		90.32	0.69	GAL 96		
Actinomyces viscosus		9.69	0.29			
27730102	**Very good identification**					**27730102**
Actinomyces viscosus		97.89	0.71	TRE 7		
Actinomyces naeslundii		2.05	0.66			
Propionibacterium propionicum		0.03	0.24			
27730103	**Good identification**					**27730103**
Actinomyces viscosus		91.39	0.63	TRE 7		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	8.61	0.69			
27730110	**Acceptable identification**					**27730110**
	Actinomyces naeslundii	91.52	0.44	GAL 96		
	Actinomyces viscosus	8.48	0.03	MLZ 5		
27730111	**Good identification**					**27730111**
	Actinomyces naeslundii	100.0	0.48	GAL 96		
27730112	**Acceptable identification**					**27730112**
	Actinomyces viscosus	90.17	0.37	TRE 7		
	Actinomyces naeslundii	9.83	0.44	MLZ 1		
27730120	**Low discrimination**					**27730120**
	Actinomyces naeslundii	77.67	0.66	GAL 96		
	Clostridium perfringens	21.96	0.35			
	Actinomyces viscosus	0.38	0.03			
27730121	**Very good identification**					**27730121**
	Actinomyces naeslundii	99.81	0.69	GAL 96		
	Clostridium perfringens	0.19	0.01			
27730122	**Low discrimination**					**27730122**
	Actinomyces naeslundii	61.90	0.66	GAL 96		
	Actinomyces viscosus	29.87	0.37			
	Clostridium perfringens	8.23	0.29			
27730123	**Good identification**					**27730123**
	Actinomyces naeslundii	90.32	0.69	GAL 96		
	Actinomyces viscosus	9.69	0.29			
27730130	**Acceptable identification**					**27730130**
	Actinomyces naeslundii	94.86	0.44	GAL 96		
	Clostridium perfringens	5.15	0.01	MLZ 5		
27730131	**Good identification**					**27730131**
	Actinomyces naeslundii	100.0	0.48	GAL 96		
27730132	**Acceptable identification**					**27730132**
	Actinomyces naeslundii	91.52	0.44	GAL 96		
	Actinomyces viscosus	8.48	0.03	MLZ 5		
27730133	**Good identification**					**27730133**
	Actinomyces naeslundii	100.0	0.48	GAL 96		
27730140	**Acceptable identification**					**27730140**
	Actinomyces naeslundii	92.67	0.53	GAL 96		
	Propionibacterium propionicum	4.77	0.21			
	Actinomyces viscosus	2.56	0.03			
27730141	**Very good identification**					**27730141**
	Actinomyces naeslundii	100.0	0.57	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27730142	**Low discrimination**					**27730142**
Actinomyces viscosus		71.74	0.37	TRE 7		
Actinomyces naeslundii		26.24	0.53	SOR 1		
Propionibacterium propionicum		2.02	0.24			
27730143	**Low discrimination**					**27730143**
Actinomyces naeslundii		62.20	0.57	GAL 96		
Actinomyces viscosus		37.80	0.29			
27730160	**Low discrimination**					**27730160**
Actinomyces naeslundii		76.62	0.53	GAL 96		
Clostridium perfringens		23.38	0.22			
27730161	**Very good identification**					**27730161**
Actinomyces naeslundii		100.0	0.57	GAL 96		
27730162	**Low discrimination**					**27730162**
Actinomyces naeslundii		85.38	0.53	GAL 96		
Clostridium perfringens		12.26	0.17			
Actinomyces viscosus		2.36	0.03			
27730163	**Very good identification**					**27730163**
Actinomyces naeslundii		100.0	0.57	GAL 96		
27730301	**Good identification**					**27730301**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27730302	**Good identification**					**27730302**
Actinomyces viscosus		96.85	0.37	TRE 7		
Actinomyces naeslundii		2.03	0.32	RHA 1		
Actinomyces odontolyticus		1.13	0.11			
27730321	**Good identification**					**27730321**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27730323	**Good identification**					**27730323**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27730501	**Good identification**					**27730501**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27730502	**Good identification**					**27730502**
Actinomyces viscosus		97.95	0.37	TRE 7		
Actinomyces naeslundii		2.05	0.32	ARA 1		
27730521	**Good identification**					**27730521**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27730523	**Good identification**					**27730523**
Actinomyces naeslundii		100.0	0.35	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27731000	**Very good identification**					**27731000**
Actinomyces naeslundii		95.49	0.82			
Actinomyces viscosus		1.88	0.29			
Clostridium perfringens		1.34	0.28			
27731001	**Very good identification**					**27731001**
Actinomyces naeslundii		99.56	0.85			
Actinomyces viscosus		0.44	0.21			
27731002	**Low discrimination**					**27731002**
Actinomyces viscosus		65.85	0.63	TRE 7		
Actinomyces naeslundii		33.71	0.82			
Clostridium perfringens		0.22	0.23			
27731003	**Low discrimination**					**27731003**
Actinomyces naeslundii		69.73	0.85			
Actinomyces viscosus		30.27	0.55			
27731010	**Very good identification**					**27731010**
					CAT	
Actinomyces naeslundii		98.39	0.60	MLZ 5	25	
Propionibacterium granulosum		0.95	0.11		90	
Lactobacillus jensenii		0.67	0.02		1	
27731011	**Very good identification**					**27731011**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27731012	**Low discrimination**					**27731012**
Actinomyces naeslundii		72.30	0.60	MLZ 5		
Actinomyces viscosus		27.10	0.29			
Actinomyces israelii		0.60	0.15			
27731013	**Good identification**					**27731013**
Actinomyces naeslundii		92.31	0.63	MLZ 5		
Actinomyces viscosus		7.69	0.21			
27731020	**Low discrimination**					**27731020**
					CAT	
Actinomyces naeslundii		68.80	0.82		25	
Clostridium perfringens		31.14	0.54		1	
Propionibacterium granulosum		0.06	0.15		90	
27731021	**Very good identification**					**27731021**
Actinomyces naeslundii		99.70	0.85			
Clostridium perfringens		0.30	0.20			
27731022	**Low discrimination**					**27731022**
Actinomyces naeslundii		81.04	0.82			
Clostridium perfringens		17.26	0.48			
Actinomyces viscosus		1.60	0.29			
27731023	**Very good identification**					**27731023**
Actinomyces naeslundii		99.42	0.85			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.44	0.21			
Clostridium perfringens		0.14	0.14			
27731030	**Good identification**					**27731030**
Actinomyces naeslundii		92.01	0.60	MLZ 5		
Clostridium perfringens		7.99	0.20			
27731031	**Very good identification**					**27731031**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27731032	**Very good identification**					**27731032**
Actinomyces naeslundii		96.07	0.60	MLZ 5		
Clostridium perfringens		3.93	0.14			
27731033	**Very good identification**					**27731033**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27731040	**Very good identification**					**27731040**
Actinomyces naeslundii		98.10	0.69			
Clostridium perfringens		1.48	0.16			
Actinomyces israelii		0.42	0.20			
27731041	**Very good identification**					**27731041**
Actinomyces naeslundii		100.0	0.72			
27731042	**Low discrimination**					**27731042**
Actinomyces naeslundii		88.87	0.69			
Actinomyces viscosus		9.94	0.29			
Clostridium perfringens		0.63	0.10			
27731043	**Very good identification**					**27731043**
Actinomyces naeslundii		97.58	0.72			
Actinomyces viscosus		2.42	0.21			
27731050	**Good identification**					**27731050**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
27731051	**Good identification**					**27731051**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27731052	**Good identification**					**27731052**
Actinomyces naeslundii		100.0	0.47	MLZ 5		
27731053	**Good identification**					**27731053**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27731060	**Low discrimination**					**27731060**
Actinomyces naeslundii		67.18	0.69			
Clostridium perfringens		32.82	0.42			
27731061	**Very good identification**					**27731061**
Actinomyces naeslundii		99.67	0.72			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		0.33	0.08			
27731062	**Low discrimination**					**27731062**
Actinomyces naeslundii		81.31	0.69			
Clostridium perfringens		18.69	0.36			
27731063	**Very good identification**					**27731063**
Actinomyces naeslundii		99.85	0.72			
Clostridium perfringens		0.16	0.02			
27731070	**Acceptable identification**					**27731070**
Actinomyces naeslundii		91.43	0.47	MLZ 5		
Clostridium perfringens		8.57	0.08			
27731071	**Good identification**					**27731071**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27731072	**Good identification**					**27731072**
Actinomyces naeslundii		95.77	0.47	MLZ 5		
Clostridium perfringens		4.23	0.02			
27731073	**Good identification**					**27731073**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27731100	**Good identification**					**27731100**
Actinomyces naeslundii		94.61	0.90			
Clostridium perfringens		3.41	0.43			
Actinomyces viscosus		1.17	0.33			
27731101	**Excellent identification**					**27731101**
					SPO	
Actinomyces naeslundii		99.66	0.93		1	
Actinomyces viscosus		0.27	0.25		1	
Clostridium chauvoei		0.04	0.12		99	
27731102	**Low discrimination**					**27731102**
Actinomyces viscosus		54.34	0.67	TRE 7		
Actinomyces naeslundii		44.54	0.90			
Clostridium perfringens		0.76	0.37			
27731103	**Low discrimination**					**27731103**
Actinomyces naeslundii		78.66	0.93			
Actinomyces viscosus		21.33	0.59			
Clostridium perfringens		0.01	0.03			
27731110	**Very good identification**					**27731110**
Actinomyces naeslundii		98.33	0.68	MLZ 5		
Actinomyces israelii		0.76	0.23			
Clostridium perfringens		0.68	0.09			
27731111	**Very good identification**					**27731111**
Actinomyces naeslundii		100.0	0.71	MLZ 5		
27731112	**Low discrimination**					**27731112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		77.70	0.68	MLZ 5	99	
Actinomyces viscosus		18.19	0.33		99	
Leptotrichia buccalis		2.14	0.07		1	
27731113	**Very good identification**					**27731113**
Actinomyces naeslundii		95.05	0.71	MLZ 5		
Actinomyces viscosus		4.95	0.25			
27731120	**Low discrimination**					**27731120**
Clostridium perfringens		53.81	0.69			
Actinomyces naeslundii		46.14	0.90			
Actinomyces israelii		0.01	0.16			
27731121	**Excellent identification**					**27731121**
					SPO	
Actinomyces naeslundii		99.21	0.93		1	
Clostridium perfringens		0.78	0.35		84	
Clostridium chauvoei		0.01	0.04		99	
27731122	**Low discrimination**					**27731122**
Actinomyces naeslundii		63.61	0.90			
Clostridium perfringens		34.91	0.63			
Actinomyces viscosus		0.78	0.33			
27731123	**Excellent identification**					**27731123**
Actinomyces naeslundii		99.36	0.93			
Clostridium perfringens		0.37	0.29			
Actinomyces viscosus		0.27	0.25			
27731130	**Low discrimination**					**27731130**
Actinomyces naeslundii		81.71	0.68	MLZ 5		
Clostridium perfringens		18.29	0.35			
27731131	**Very good identification**					**27731131**
Actinomyces naeslundii		99.85	0.71	MLZ 5		
Clostridium perfringens		0.15	0.01			
27731132	**Low discrimination**					**27731132**
					GRA	
Actinomyces naeslundii		87.18	0.68	MLZ 5	99	
Clostridium perfringens		9.18	0.29		99	
Parabacteroides distasonis		3.16	0.30		1	
27731133	**Very good identification**					**27731133**
Actinomyces naeslundii		100.0	0.71	MLZ 5		
27731140	**Very good identification**					**27731140**
Actinomyces naeslundii		95.44	0.77			
Clostridium perfringens		3.72	0.31			
Actinomyces israelii		0.57	0.30			
27731141	**Very good identification**					**27731141**
Actinomyces naeslundii		100.0	0.80			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27731142	**Good identification**					**27731142**
Actinomyces naeslundii		90.76	0.77			
Actinomyces viscosus		6.34	0.33			
Clostridium perfringens		1.66	0.25			
27731143	**Very good identification**					**27731143**
Actinomyces naeslundii		98.47	0.80			
Actinomyces viscosus		1.53	0.25			
27731150	**Good identification**					**27731150**
Actinomyces naeslundii		98.33	0.55	MLZ 5		
Actinomyces israelii		1.67	0.16			
27731151	**Very good identification**					**27731151**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27731152	**Good identification**					**27731152**
Actinomyces naeslundii		96.25	0.55	MLZ 5		
Actinomyces israelii		2.46	0.19			
Actinomyces viscosus		1.29	-.01			
27731153	**Very good identification**					**27731153**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27731160	**Low discrimination**					**27731160**
Clostridium perfringens		55.73	0.57			
Actinomyces naeslundii		44.28	0.77			
27731161	**Very good identification**					**27731161**
Actinomyces naeslundii		99.16	0.80			
Clostridium perfringens		0.84	0.22			
27731162	**Low discrimination**					**27731162**
Actinomyces naeslundii		62.72	0.77			
Clostridium perfringens		37.15	0.51			
Lactobacillus acidophilus		0.09	0.04			
27731163	**Very good identification**					**27731163**
Actinomyces naeslundii		99.60	0.80			
Clostridium perfringens		0.40	0.17			
27731170	**Low discrimination**					**27731170**
Actinomyces naeslundii		80.54	0.55	MLZ 5		
Clostridium perfringens		19.46	0.22			
27731171	**Very good identification**					**27731171**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27731172	**Low discrimination**					**27731172**
Actinomyces naeslundii		89.79	0.55	MLZ 5		
Clostridium perfringens		10.21	0.17			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27731173	**Very good identification**					**27731173**
Actinomyces naeslundii		100.0	0.58	MLZ 5		
27731200	**Low discrimination**					**27731200**
Actinomyces naeslundii		81.50	0.48	RHA 1		
Actinomyces odontolyticus		18.50	0.20			
27731201	**Good identification**					**27731201**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27731203	**Low discrimination**					**27731203**
Actinomyces naeslundii		69.73	0.51	RHA 1		
Actinomyces viscosus		30.27	0.21			
27731220	**Low discrimination**					**27731220**
Actinomyces naeslundii		67.14	0.48	RHA 1		
Clostridium perfringens		30.38	0.20			
Actinomyces odontolyticus		2.48	0.06			
27731221	**Good identification**					**27731221**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27731222	**Low discrimination**					**27731222**
Actinomyces naeslundii		82.44	0.48	RHA 1		
Clostridium perfringens		17.56	0.14			
27731223	**Good identification**					**27731223**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27731240	**Good identification**					**27731240**
Actinomyces naeslundii		100.0	0.35	RHA 1		
27731241	**Good identification**					**27731241**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27731242	**Good identification**					**27731242**
Actinomyces naeslundii		100.0	0.35	RHA 1		
27731243	**Good identification**					**27731243**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27731260	**Low discrimination**					**27731260**
Actinomyces naeslundii		67.18	0.35	RHA 1		
Clostridium perfringens		32.82	0.08			
27731261	**Good identification**					**27731261**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27731262	**Low discrimination**					**27731262**
Actinomyces naeslundii		81.31	0.35	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		18.70	0.02			
27731263	**Good identification**					**27731263**
Actinomyces naeslundii		100.0	0.38	RHA 1		
27731300	**Acceptable identification**					**27731300**
Actinomyces naeslundii		91.42	0.56	RHA 1		
Clostridium perfringens		3.30	0.09			
Actinomyces odontolyticus		2.30	0.11			
27731301	**Very good identification**					**27731301**
Actinomyces naeslundii		100.0	0.59	RHA 1		
27731303	**Low discrimination**					**27731303**
Actinomyces naeslundii		78.67	0.59	RHA 1		
Actinomyces viscosus		21.33	0.25			
27731311	**Good identification**					**27731311**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27731313	**Good identification**					**27731313**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27731321	**Very good identification**					**27731321**
Actinomyces naeslundii		99.22	0.59	RHA 1		
Clostridium perfringens		0.78	0.01			
27731322	**Low discrimination**					**27731322**
					GRA	
Parabacteroides distasonis		36.24	0.43	NIT 1	1	
Actinomyces naeslundii		34.40	0.56	XYL 90	99	
Clostridium perfringens		18.88	0.29		99	
27731323	**Very good identification**					**27731323**
Actinomyces naeslundii		100.0	0.59	RHA 1		
27731331	**Good identification**					**27731331**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27731332	**Low discrimination**					**27731332**
					GRA	
Parabacteroides distasonis		87.74	0.39	NIT 1	1	
Actinomyces naeslundii		8.14	0.34	XYL 90	99	
Clostridium clostridioforme		4.12	0.10		50	
27731333	**Good identification**					**27731333**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27731340	**Good identification**					**27731340**
Actinomyces naeslundii		100.0	0.43	RHA 1		
27731341	**Good identification**					**27731341**
Actinomyces naeslundii		100.0	0.46	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27731342	**Low discrimination**					**27731342**
Actinomyces naeslundii		87.96	0.43	RHA 1		
Actinomyces viscosus		6.14	-.01			
Actinomyces israelii		5.89	0.14			
27731343	**Good identification**					**27731343**
Actinomyces naeslundii		100.0	0.46	RHA 1		
27731361	**Good identification**					**27731361**
Actinomyces naeslundii		100.0	0.46	RHA 1		
27731362	**Low discrimination**					**27731362**
Actinomyces naeslundii		62.80	0.43	RHA 1		
Clostridium perfringens		37.20	0.17			
27731363	**Good identification**					**27731363**
Actinomyces naeslundii		100.0	0.46	RHA 1		
27731400	**Low discrimination**					**27731400**
Actinomyces naeslundii		86.45	0.48	ARA 1		
Actinomyces israelii		13.55	0.25			
27731401	**Good identification**					**27731401**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27731403	**Low discrimination**					**27731403**
Actinomyces naeslundii		69.73	0.51	ARA 1		
Actinomyces viscosus		30.27	0.21			
27731420	**Low discrimination**					**27731420**
Actinomyces naeslundii		68.84	0.48	ARA 1		
Clostridium perfringens		31.16	0.20			
27731421	**Good identification**					**27731421**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27731422	**Low discrimination**					**27731422**
Actinomyces naeslundii		82.44	0.48	ARA 1		
Clostridium perfringens		17.56	0.14			
27731423	**Good identification**					**27731423**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27731440	**Low discrimination**					**27731440**
Actinomyces naeslundii		74.32	0.35	ARA 1		
Actinomyces israelii		25.68	0.18			
27731441	**Good identification**					**27731441**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27731442	**Low discrimination**					**27731442**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		65.87	0.35	ARA 1		
Actinomyces israelii		34.13	0.21			
27731443	**Good identification**					**27731443**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27731460	**Low discrimination**					**27731460**
Actinomyces naeslundii		67.18	0.35	ARA 1		
Clostridium perfringens		32.82	0.08			
27731461	**Good identification**					**27731461**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27731462	**Low discrimination**					**27731462**
Actinomyces naeslundii		81.31	0.35	ARA 1		
Clostridium perfringens		18.70	0.02			
27731463	**Good identification**					**27731463**
Actinomyces naeslundii		100.0	0.38	ARA 1		
27731500	**Low discrimination**					**27731500**
Actinomyces naeslundii		78.82	0.56	ARA 1		
Actinomyces israelii		17.37	0.35			
Clostridium perfringens		2.84	0.09			
27731501	**Very good identification**					**27731501**
Actinomyces naeslundii		100.0	0.59	ARA 1		
27731503	**Low discrimination**					**27731503**
Actinomyces naeslundii		78.67	0.59	ARA 1		
Actinomyces viscosus		21.33	0.25			
27731511	**Good identification**					**27731511**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27731513	**Good identification**					**27731513**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27731521	**Very good identification**					**27731521**
Actinomyces naeslundii		99.22	0.59	ARA 1		
Clostridium perfringens		0.78	0.01			
27731522	**Low discrimination**					**27731522**
					GRA	
Actinomyces naeslundii		54.68	0.56	ARA 1	99	
Clostridium perfringens		30.00	0.29		99	
Parabacteroides distasonis		11.77	0.31		1	
27731523	**Very good identification**					**27731523**
Actinomyces naeslundii		100.0	0.59	ARA 1		
27731531	**Good identification**					**27731531**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ARA 1		
27731533	**Good identification**					**27731533**
Actinomyces naeslundii		100.0	0.37	ARA 1		
27731540	**Low discrimination**					**27731540**
Actinomyces naeslundii		67.32	0.43	ARA 1		
Actinomyces israelii		32.68	0.28			
27731541	**Good identification**					**27731541**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27731542	**Low discrimination**					**27731542**
Actinomyces naeslundii		49.84	0.43	ARA 1		
Actinomyces israelii		36.30	0.31			
Bifidobacterium adolescentis		10.38	0.16			
27731543	**Good identification**					**27731543**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27731561	**Good identification**					**27731561**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27731562	**Low discrimination**					**27731562**
Actinomyces naeslundii		62.80	0.43	ARA 1		
Clostridium perfringens		37.20	0.17			
27731563	**Good identification**					**27731563**
Actinomyces naeslundii		100.0	0.46	ARA 1		
27731722	**Low discrimination**					**27731722**
Parabacteroides distasonis		87.56	0.40	NIT 1		
Clostridium clostridioforme		12.44	0.19	XYL 90		
27732000	**Very good identification**					**27732000**
					CAT	
Actinomyces naeslundii		95.06	0.64	GAL 96	25	
Propionibacterium granulosum		3.29	0.24		90	
Propionibacterium propionicum		1.27	0.21		1	
27732001	**Very good identification**					**27732001**
Actinomyces naeslundii		100.0	0.67	GAL 96		
27732002	**Low discrimination**					**27732002**
Actinomyces naeslundii		70.10	0.64	GAL 96		
Actinomyces viscosus		27.90	0.33			
Propionibacterium propionicum		1.40	0.24			
27732003	**Good identification**					**27732003**
Actinomyces naeslundii		91.88	0.67	GAL 96		
Actinomyces viscosus		8.13	0.24			
27732010	**Acceptable identification**					**27732010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.76	0.42	GAL 96		
Propionibacterium granulosum		8.24	0.09	MLZ 5		
27732011	**Good identification**					**27732011**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27732012	**Acceptable identification**					**27732012**
Actinomyces naeslundii		92.91	0.42	GAL 96		
Actinomyces viscosus		7.10	-.02	MLZ 5		
27732013	**Good identification**					**27732013**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27732020	**Very good identification**					**27732020**
Actinomyces naeslundii		99.14	0.64	GAL 96		
Propionibacterium granulosum		0.86	0.13			
27732021	**Very good identification**					**27732021**
Actinomyces naeslundii		100.0	0.67	GAL 96		
27732022	**Very good identification**					**27732022**
Actinomyces naeslundii		99.60	0.64	GAL 96		
Actinomyces viscosus		0.40	-.02			
27732023	**Very good identification**					**27732023**
Actinomyces naeslundii		100.0	0.67	GAL 96		
27732030	**Good identification**					**27732030**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27732031	**Good identification**					**27732031**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27732032	**Good identification**					**27732032**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27732033	**Good identification**					**27732033**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27732040	**Acceptable identification**					**27732040**
Actinomyces naeslundii		92.99	0.51	GAL 96		
Propionibacterium propionicum		7.01	0.21			
27732041	**Good identification**					**27732041**
Actinomyces naeslundii		100.0	0.54	GAL 96		
27732042	**Low discrimination**					**27732042**
Actinomyces naeslundii		88.04	0.51	GAL 96		
Propionibacterium propionicum		9.96	0.24			
Actinomyces viscosus		2.01	-.02			
27732043	**Good identification**					**27732043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.54	GAL 96		
27732060	**Good identification**					**27732060**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27732061	**Good identification**					**27732061**
Actinomyces naeslundii		100.0	0.54	GAL 96		
27732062	**Good identification**					**27732062**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27732063	**Good identification**					**27732063**
Actinomyces naeslundii		100.0	0.54	GAL 96		
27732100	**Low discrimination**					**27732100**
Actinomyces naeslundii		68.15	0.71	GAL 96		
Propionibacterium propionicum		31.55	0.55			
Actinomyces viscosus		0.17	0.03			
27732101	**Very good identification**					**27732101**
Actinomyces naeslundii		96.69	0.74	GAL 96		
Propionibacterium propionicum		3.32	0.39			
27732102	**Low discrimination**					**27732102**
Actinomyces naeslundii		51.39	0.71	GAL 96		
Propionibacterium propionicum		35.68	0.58			
Actinomyces viscosus		12.78	0.37			
27732103	**Good identification**					**27732103**
Actinomyces naeslundii		90.36	0.74	GAL 96		
Actinomyces viscosus		4.99	0.29			
Propionibacterium propionicum		4.65	0.42			
27732110	**Acceptable identification**					**27732110**
Actinomyces naeslundii		91.84	0.49	GAL 96		
Propionibacterium propionicum		8.16	0.21	MLZ 5		
27732111	**Good identification**					**27732111**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27732112	**Low discrimination**					**27732112**
Actinomyces naeslundii		84.68	0.49	GAL 96		
Propionibacterium propionicum		11.28	0.24	MLZ 5		
Actinomyces viscosus		4.04	0.03			
27732113	**Good identification**					**27732113**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27732120	**Very good identification**					**27732120**
Actinomyces naeslundii		99.39	0.71	GAL 96		
Propionibacterium propionicum		0.47	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		0.15	0.01			
27732121	**Very good identification**					**27732121**
Actinomyces naeslundii		100.0	0.74	GAL 96		
27732122	**Very good identification**					**27732122**
Actinomyces naeslundii		99.06	0.71	GAL 96		
Propionibacterium propionicum		0.70	0.24			
Actinomyces viscosus		0.25	0.03			
27732123	**Very good identification**					**27732123**
Actinomyces naeslundii		100.0	0.74	GAL 96		
27732130	**Good identification**					**27732130**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27732131	**Good identification**					**27732131**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27732132	**Good identification**					**27732132**
Actinomyces naeslundii		100.0	0.49	GAL 96		
27732133	**Good identification**					**27732133**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27732140	**Low discrimination**					**27732140**
Propionibacterium propionicum		72.40	0.55	SAL 1		
Actinomyces naeslundii		27.60	0.58			
27732141	**Low discrimination**					**27732141**
Actinomyces naeslundii		83.73	0.61	GAL 96		
Propionibacterium propionicum		16.27	0.39			
27732142	**Low discrimination**					**27732142**
Propionibacterium propionicum		79.51	0.58	SAL 1		
Actinomyces naeslundii		20.21	0.58			
Actinomyces viscosus		0.29	0.03			
27732143	**Low discrimination**					**27732143**
Actinomyces naeslundii		77.43	0.61	GAL 96		
Propionibacterium propionicum		22.57	0.42			
27732150	**Low discrimination**					**27732150**
Actinomyces naeslundii		66.52	0.37	GAL 96		
Propionibacterium propionicum		33.48	0.21	MLZ 5		
27732151	**Good identification**					**27732151**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732152	**Low discrimination**					**27732152**
Actinomyces naeslundii		56.98	0.37	GAL 96		
Propionibacterium propionicum		43.02	0.24	MLZ 5		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27732153	**Good identification**					**27732153**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732160	**Very good identification**					**27732160**
Actinomyces naeslundii		97.42	0.58	GAL 96		
Propionibacterium propionicum		2.58	0.21			
27732161	**Very good identification**					**27732161**
Actinomyces naeslundii		100.0	0.61	GAL 96		
27732162	**Very good identification**					**27732162**
Actinomyces naeslundii		96.18	0.58	GAL 96		
Propionibacterium propionicum		3.82	0.24			
27732163	**Very good identification**					**27732163**
Actinomyces naeslundii		100.0	0.61	GAL 96		
27732170	**Good identification**					**27732170**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27732171	**Good identification**					**27732171**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732172	**Good identification**					**27732172**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27732173	**Good identification**					**27732173**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732300	**Low discrimination**					**27732300**
Actinomyces naeslundii		68.36	0.37	GAL 96		
Propionibacterium propionicum		31.64	0.21	RHA 1		
27732301	**Good identification**					**27732301**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732302	**Low discrimination**					**27732302**
Actinomyces naeslundii		51.47	0.37	GAL 96		
Propionibacterium propionicum		35.74	0.24	RHA 1		
Actinomyces viscosus		12.79	0.03			
27732303	**Good identification**					**27732303**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732320	**Good identification**					**27732320**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27732321	**Good identification**					**27732321**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732322	**Good identification**					**27732322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GAL 96		
27732323	**Good identification**					**27732323**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732500	**Low discrimination**					**27732500**
Actinomyces naeslundii		61.77	0.37	GAL 96		
Propionibacterium propionicum		28.60	0.21	ARA 1		
Actinomyces israelii		9.63	0.14			
27732501	**Good identification**					**27732501**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732502	**Low discrimination**					**27732502**
Actinomyces naeslundii		45.94	0.37	GAL 96		
Propionibacterium propionicum		31.90	0.24	ARA 1		
Actinomyces viscosus		11.42	0.03			
27732503	**Good identification**					**27732503**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732520	**Good identification**					**27732520**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27732521	**Good identification**					**27732521**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27732522	**Good identification**					**27732522**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27732523	**Good identification**					**27732523**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27733000	**Very good identification**					**27733000**
					CAT	
Actinomyces naeslundii		99.24	0.87		25	
Actinomyces israelii		0.56	0.39		1	
Propionibacterium granulosum		0.14	0.24		90	
27733001	**Very good identification**					**27733001**
Actinomyces naeslundii		100.0	0.90			
27733002	**Very good identification**					**27733002**
Actinomyces naeslundii		98.06	0.87			
Actinomyces viscosus		1.00	0.29			
Actinomyces israelii		0.83	0.42			
27733003	**Very good identification**					**27733003**
Actinomyces naeslundii		99.78	0.90			
Actinomyces viscosus		0.23	0.21			
27733010	**Very good identification**					**27733010**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.06	0.65	MLZ 5	25	
Actinomyces israelii		1.57	0.25		1	
Propionibacterium granulosum		0.37	0.09		90	
27733011	**Very good identification**					**27733011**
Actinomyces naeslundii		100.0	0.68	MLZ 5		
27733012	**Very good identification**					**27733012**
Actinomyces naeslundii		97.65	0.65	MLZ 5		
Actinomyces israelii		2.35	0.28			
27733013	**Very good identification**					**27733013**
Actinomyces naeslundii		100.0	0.68	MLZ 5		
27733020	**Very good identification**					**27733020**
Actinomyces naeslundii		99.69	0.87			
Clostridium perfringens		0.24	0.20			
Actinomyces israelii		0.04	0.19			
27733021	**Very good identification**					**27733021**
Actinomyces naeslundii		100.0	0.90			
27733022	**Very good identification**					**27733022**
Actinomyces naeslundii		99.84	0.87			
Clostridium perfringens		0.11	0.14			
Actinomyces israelii		0.05	0.22			
27733023	**Very good identification**					**27733023**
Actinomyces naeslundii		100.0	0.90			
27733030	**Very good identification**					**27733030**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27733031	**Very good identification**					**27733031**
Actinomyces naeslundii		100.0	0.68	MLZ 5		
27733032	**Very good identification**					**27733032**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27733033	**Very good identification**					**27733033**
Actinomyces naeslundii		100.0	0.68	MLZ 5		
27733040	**Very good identification**					**27733040**
Actinomyces naeslundii		98.47	0.74			
Actinomyces israelii		1.23	0.32			
Propionibacterium propionicum		0.31	0.21			
27733041	**Very good identification**					**27733041**
Actinomyces naeslundii		100.0	0.77			
27733042	**Very good identification**					**27733042**
Actinomyces naeslundii		97.72	0.74			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		1.83	0.35			
Propionibacterium propionicum		0.46	0.24			
27733043	**Very good identification**					**27733043**
Actinomyces naeslundii		100.0	0.77			
27733050	**Good identification**					**27733050**
Actinomyces naeslundii		96.59	0.52	MLZ 5		
Actinomyces israelii		3.41	0.18			
27733051	**Good identification**					**27733051**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27733052	**Acceptable identification**					**27733052**
Actinomyces naeslundii		94.97	0.52	MLZ 5		
Actinomyces israelii		5.04	0.21			
27733053	**Good identification**					**27733053**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27733060	**Very good identification**					**27733060**
Actinomyces naeslundii		99.75	0.74			
Clostridium perfringens		0.25	0.08			
27733061	**Very good identification**					**27733061**
Actinomyces naeslundii		100.0	0.77			
27733062	**Very good identification**					**27733062**
Actinomyces naeslundii		99.76	0.74			
Actinomyces israelii		0.12	0.15			
Clostridium perfringens		0.12	0.02			
27733063	**Very good identification**					**27733063**
Actinomyces naeslundii		100.0	0.77			
27733070	**Good identification**					**27733070**
Actinomyces naeslundii		100.0	0.52	MLZ 5		
27733071	**Good identification**					**27733071**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27733072	**Good identification**					**27733072**
Actinomyces naeslundii		100.0	0.52	MLZ 5		
27733073	**Good identification**					**27733073**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27733100	**Very good identification**					**27733100**
Actinomyces naeslundii		97.33	0.95			
Propionibacterium propionicum		1.88	0.55			
Actinomyces israelii		0.77	0.50			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27733101	**Excellent identification**					**27733101**
Actinomyces naeslundii		99.85	0.98			
Propionibacterium propionicum		0.14	0.39			
Actinomyces israelii		0.01	0.16			
27733102	**Very good identification**					**27733102**
Actinomyces naeslundii		95.47	0.95			
Propionibacterium propionicum		2.76	0.58			
Actinomyces israelii		1.14	0.53			
27733103	**Excellent identification**					**27733103**
Actinomyces naeslundii		99.64	0.98			
Propionibacterium propionicum		0.21	0.42			
Actinomyces viscosus		0.14	0.25			
27733110	**Very good identification**					**27733110**
Actinomyces naeslundii		97.44	0.73	MLZ 5		
Actinomyces israelii		2.20	0.36			
Propionibacterium propionicum		0.36	0.21			
27733111	**Very good identification**					**27733111**
Actinomyces naeslundii		100.0	0.76	MLZ 5		
27733112	**Very good identification**					**27733112**
Actinomyces naeslundii		95.70	0.73	MLZ 5		
Actinomyces israelii		3.24	0.39			
Propionibacterium propionicum		0.53	0.24			
27733113	**Very good identification**					**27733113**
Actinomyces naeslundii		100.0	0.76	MLZ 5		
27733120	**Excellent identification**					**27733120**
					SPO	
Actinomyces naeslundii		99.27	0.95		1	
Clostridium perfringens		0.60	0.35		84	
Clostridium ramosum		0.06	0.15		92	
27733121	**Excellent identification**					**27733121**
Actinomyces naeslundii		100.0	0.98			
Clostridium perfringens		0.00	0.01			
27733122	**Excellent identification**					**27733122**
					SPO	
Actinomyces naeslundii		99.36	0.95		1	
Clostridium perfringens		0.28	0.29		84	
Clostridium ramosum		0.25	0.25		92	
27733123	**Excellent identification**					**27733123**
Actinomyces naeslundii		100.0	0.98			
27733130	**Very good identification**					**27733130**
Actinomyces naeslundii		99.74	0.73	MLZ 5		
Actinomyces israelii		0.14	0.15			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		0.12	0.01			
27733131	**Very good identification**					**27733131**
Actinomyces naeslundii		100.0	0.76	MLZ 5		
27733132	**Very good identification**					**27733132**
Actinomyces naeslundii		99.79	0.73	MLZ 5		
Actinomyces israelii		0.22	0.18			
27733133	**Very good identification**					**27733133**
Actinomyces naeslundii		100.0	0.76	MLZ 5		
27733140	**Low discrimination**					**27733140**
Actinomyces naeslundii		88.75	0.82			
Propionibacterium propionicum		9.70	0.55			
Actinomyces israelii		1.55	0.43			
27733141	**Very good identification**					**27733141**
Actinomyces naeslundii		99.20	0.85			
Propionibacterium propionicum		0.80	0.39			
27733142	**Low discrimination**					**27733142**
Actinomyces naeslundii		83.99	0.82			
Propionibacterium propionicum		13.77	0.58			
Actinomyces israelii		2.21	0.46			
27733143	**Very good identification**					**27733143**
Actinomyces naeslundii		98.80	0.85			
Propionibacterium propionicum		1.20	0.42			
27733150	**Good identification**					**27733150**
Actinomyces naeslundii		93.40	0.60	MLZ 5		
Actinomyces israelii		4.64	0.29			
Propionibacterium propionicum		1.96	0.21			
27733151	**Very good identification**					**27733151**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27733152	**Good identification**					**27733152**
Actinomyces naeslundii		90.42	0.60	MLZ 5		
Actinomyces israelii		6.74	0.32			
Propionibacterium propionicum		2.85	0.24			
27733153	**Very good identification**					**27733153**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27733160	**Very good identification**					**27733160**
Actinomyces naeslundii		99.13	0.82			
Clostridium perfringens		0.65	0.22			
Actinomyces israelii		0.11	0.22			
27733161	**Very good identification**					**27733161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.85			
27733162	**Very good identification**					**27733162**
Actinomyces naeslundii		99.36	0.82			
Clostridium perfringens		0.31	0.17			
Actinomyces israelii		0.17	0.25			
27733163	**Very good identification**					**27733163**
Actinomyces naeslundii		100.0	0.85			
27733170	**Very good identification**					**27733170**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27733171	**Very good identification**					**27733171**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27733172	**Very good identification**					**27733172**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27733173	**Very good identification**					**27733173**
Actinomyces naeslundii		100.0	0.63	MLZ 5		
27733200	**Good identification**					**27733200**
Actinomyces naeslundii		95.96	0.53	RHA 1		
Actinomyces israelii		4.04	0.20			
27733201	**Good identification**					**27733201**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27733202	**Acceptable identification**					**27733202**
Actinomyces naeslundii		94.06	0.53	RHA 1		
Actinomyces israelii		5.94	0.23			
27733203	**Good identification**					**27733203**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27733220	**Good identification**					**27733220**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27733221	**Good identification**					**27733221**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27733222	**Good identification**					**27733222**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27733223	**Good identification**					**27733223**
Actinomyces naeslundii		100.0	0.56	RHA 1		
27733240	**Acceptable identification**					**27733240**
Actinomyces naeslundii		91.51	0.40	RHA 1		
Actinomyces israelii		8.49	0.13			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
27733241	**Good identification**					**27733241**
Actinomyces naeslundii		100.0	0.43	RHA 1		
27733242	**Low discrimination**					**27733242**
Actinomyces naeslundii		87.78	0.40	RHA 1		
Actinomyces israelii		12.22	0.16			
27733243	**Good identification**					**27733243**
Actinomyces naeslundii		100.0	0.43	RHA 1		
27733260	**Good identification**					**27733260**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27733261	**Good identification**					**27733261**
Actinomyces naeslundii		100.0	0.43	RHA 1		
27733262	**Good identification**					**27733262**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27733263	**Good identification**					**27733263**
Actinomyces naeslundii		100.0	0.43	RHA 1		
27733300	**Good identification**					**27733300**
Actinomyces naeslundii		92.73	0.61	RHA 1		
Actinomyces israelii		5.49	0.31			
Propionibacterium propionicum		1.79	0.21			
27733301	**Very good identification**					**27733301**
Actinomyces naeslundii		100.0	0.64	RHA 1		
27733302	**Low discrimination**					**27733302**
Actinomyces naeslundii		88.97	0.61	RHA 1		
Actinomyces israelii		7.90	0.34			
Propionibacterium propionicum		2.57	0.24			
27733303	**Very good identification**					**27733303**
Actinomyces naeslundii		100.0	0.64	RHA 1		
27733310	**Low discrimination**					**27733310**
Actinomyces naeslundii		85.62	0.39	RHA 1		
Actinomyces israelii		14.39	0.16	MLZ 5		
27733311	**Good identification**					**27733311**
Actinomyces naeslundii		100.0	0.42	RHA 1		
27733312	**Low discrimination**					**27733312**
Actinomyces naeslundii		79.88	0.39	RHA 1		
Actinomyces israelii		20.13	0.19	MLZ 5		
27733313	**Good identification**					**27733313**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	RHA 1		
27733320	**Good identification**					**27733320**
					SPO	
Actinomyces naeslundii		91.98	0.61	RHA 1	1	
Clostridium ramosum		7.46	0.17		92	
Clostridium perfringens		0.56	0.01		84	
27733321	**Very good identification**					**27733321**
Actinomyces naeslundii		100.0	0.64	RHA 1		
27733322	**Low discrimination**					**27733322**
					SPO	
Actinomyces naeslundii		75.50	0.61	RHA 1	1	
Clostridium ramosum		24.50	0.27		92	
27733323	**Very good identification**					**27733323**
Actinomyces naeslundii		100.0	0.64	RHA 1		
27733330	**Good identification**					**27733330**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27733331	**Good identification**					**27733331**
Actinomyces naeslundii		100.0	0.42	RHA 1		
27733332	**Good identification**					**27733332**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27733333	**Good identification**					**27733333**
Actinomyces naeslundii		100.0	0.42	RHA 1		
27733340	**Low discrimination**					**27733340**
Actinomyces naeslundii		80.67	0.48	RHA 1		
Actinomyces israelii		10.52	0.24			
Propionibacterium propionicum		8.82	0.21			
27733341	**Good identification**					**27733341**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27733342	**Low discrimination**					**27733342**
Actinomyces naeslundii		73.56	0.48	RHA 1		
Actinomyces israelii		14.39	0.27			
Propionibacterium propionicum		12.06	0.24			
27733343	**Good identification**					**27733343**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27733360	**Good identification**					**27733360**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27733361	**Good identification**					**27733361**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27733362	**Good identification**					**27733362**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	RHA 1		
27733363	**Good identification**					**27733363**
Actinomyces naeslundii		100.0	0.51	RHA 1		
27733400	**Low discrimination**					**27733400**
Actinomyces naeslundii		68.61	0.53	ARA 1		
Actinomyces israelii		31.40	0.38			
27733401	**Good identification**					**27733401**
Actinomyces naeslundii		100.0	0.56	ARA 1		
27733402	**Low discrimination**					**27733402**
Actinomyces naeslundii		59.30	0.53	ARA 1		
Actinomyces israelii		40.70	0.41			
27733403	**Good identification**					**27733403**
Actinomyces naeslundii		100.0	0.56	ARA 1		
27733420	**Good identification**					**27733420**
Actinomyces naeslundii		97.16	0.53	ARA 1		
Actinomyces israelii		2.84	0.18			
27733421	**Good identification**					**27733421**
Actinomyces naeslundii		100.0	0.56	ARA 1		
27733422	**Good identification**					**27733422**
Actinomyces naeslundii		95.80	0.53	ARA 1		
Actinomyces israelii		4.20	0.21			
27733423	**Good identification**					**27733423**
Actinomyces naeslundii		100.0	0.56	ARA 1		
27733441	**Good identification**					**27733441**
Actinomyces naeslundii		100.0	0.43	ARA 1		
27733443	**Good identification**					**27733443**
Actinomyces naeslundii		100.0	0.43	ARA 1		
27733460	**Good identification**					**27733460**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27733461	**Good identification**					**27733461**
Actinomyces naeslundii		100.0	0.43	ARA 1		
27733462	**Acceptable identification**					**27733462**
Actinomyces naeslundii		91.19	0.40	ARA 1		
Actinomyces israelii		8.81	0.14			
27733463	**Good identification**					**27733463**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ARA 1		
27733500	**Low discrimination**					**27733500**
Actinomyces naeslundii		60.15	0.61	ARA 1		
Actinomyces israelii		38.69	0.48			
Propionibacterium propionicum		1.16	0.21			
27733501	**Very good identification**					**27733501**
Actinomyces naeslundii		99.57	0.64	ARA 1		
Actinomyces israelii		0.43	0.14			
27733502	**Low discrimination**					**27733502**
Actinomyces naeslundii		47.05	0.61	ARA 1		
Actinomyces israelii		45.39	0.51			
Bifidobacterium adolescentis		5.90	0.30			
27733503	**Very good identification**					**27733503**
Actinomyces naeslundii		99.35	0.64	ARA 1		
Actinomyces israelii		0.65	0.17			
27733511	**Good identification**					**27733511**
Actinomyces naeslundii		100.0	0.42	ARA 1		
27733513	**Good identification**					**27733513**
Actinomyces naeslundii		100.0	0.42	ARA 1		
27733520	**Very good identification**					**27733520**
Actinomyces naeslundii		95.50	0.61	ARA 1		
Actinomyces israelii		3.92	0.28			
Clostridium perfringens		0.58	0.01			
27733521	**Very good identification**					**27733521**
Actinomyces naeslundii		100.0	0.64	ARA 1		
27733522	**Good identification**					**27733522**
Actinomyces naeslundii		94.20	0.61	ARA 1		
Actinomyces israelii		5.80	0.31			
27733523	**Very good identification**					**27733523**
Actinomyces naeslundii		100.0	0.64	ARA 1		
27733530	**Low discrimination**					**27733530**
Actinomyces naeslundii		89.56	0.39	ARA 1		
Actinomyces israelii		10.44	0.14	MLZ 5		
27733531	**Good identification**					**27733531**
Actinomyces naeslundii		100.0	0.42	ARA 1		
27733532	**Low discrimination**					**27733532**
Actinomyces naeslundii		85.12	0.39	ARA 1		
Actinomyces israelii		14.88	0.17	MLZ 5		
27733533	**Good identification**					**27733533**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	ARA 1		
27733540	**Low discrimination**					**27733540**
Actinomyces israelii		56.10	0.41	CEL 93		
Actinomyces naeslundii		39.58	0.48	XYL 92		
Propionibacterium propionicum		4.33	0.21			
27733541	**Good identification**					**27733541**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27733542	**Low discrimination**					**27733542**
Actinomyces israelii		62.94	0.44	CEL 93		
Actinomyces naeslundii		29.61	0.48	XYL 92		
Propionibacterium propionicum		4.85	0.24			
27733543	**Good identification**					**27733543**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27733560	**Acceptable identification**					**27733560**
Actinomyces naeslundii		91.70	0.48	ARA 1		
Actinomyces israelii		8.30	0.21			
27733561	**Good identification**					**27733561**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27733562	**Low discrimination**					**27733562**
Actinomyces naeslundii		88.05	0.48	ARA 1		
Actinomyces israelii		11.95	0.24			
27733563	**Good identification**					**27733563**
Actinomyces naeslundii		100.0	0.51	ARA 1		
27734101	**Good identification**					**27734101**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27734102	**Good identification**					**27734102**
Actinomyces viscosus		95.74	0.37	TRE 7		
Actinomyces israelii		2.26	0.24	XYL 1		
Actinomyces naeslundii		2.00	0.32			
27734121	**Good identification**					**27734121**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27734123	**Good identification**					**27734123**
Actinomyces naeslundii		100.0	0.35	GAL 96		
27734712	**Good identification**					**27734712**
Clostridium clostridioforme		100.0	0.40	NIT 1		
27734732	**Good identification**					**27734732**
Clostridium clostridioforme		97.81	0.40	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		2.20	0.17			
27735000	**Low discrimination**					**27735000**
Actinomyces israelii		66.72	0.45	CEL 93		
Actinomyces naeslundii		30.28	0.48			
Actinomyces odontolyticus		1.74	0.10			
27735001	**Good identification**					**27735001**
Actinomyces naeslundii		100.0	0.51	XYL 1		
27735002	**Low discrimination**					**27735002**
Actinomyces israelii		52.52	0.48	CEL 93		
Actinomyces viscosus		31.04	0.29			
Actinomyces naeslundii		15.89	0.48			
27735003	**Low discrimination**					**27735003**
Actinomyces naeslundii		68.67	0.51	XYL 1		
Actinomyces viscosus		29.81	0.21			
Actinomyces israelii		1.53	0.14			
27735020	**Low discrimination**					**27735020**
Actinomyces naeslundii		62.77	0.48	XYL 1		
Clostridium perfringens		28.41	0.20			
Actinomyces israelii		8.83	0.24			
27735021	**Good identification**					**27735021**
Actinomyces naeslundii		100.0	0.51	XYL 1		
27735022	**Low discrimination**					**27735022**
Actinomyces naeslundii		66.02	0.48	XYL 1		
Clostridium perfringens		14.06	0.14			
Actinomyces israelii		13.93	0.27			
27735023	**Good identification**					**27735023**
Actinomyces naeslundii		100.0	0.51	XYL 1		
27735040	**Low discrimination**					**27735040**
Actinomyces israelii		82.92	0.38	CEL 93		
Actinomyces naeslundii		17.08	0.35			
27735041	**Good identification**					**27735041**
Actinomyces naeslundii		100.0	0.38	XYL 1		
27735042	**Low discrimination**					**27735042**
Actinomyces israelii		87.93	0.41	CEL 93		
Actinomyces naeslundii		12.07	0.35			
27735043	**Good identification**					**27735043**
Actinomyces naeslundii		100.0	0.38	XYL 1		
27735060	**Low discrimination**					**27735060**
Actinomyces naeslundii		55.60	0.35	XYL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Clostridium perfringens	27.16	0.08			
	Actinomyces israelii	17.23	0.17			
27735061	**Good identification**					**27735061**
	Actinomyces naeslundii	100.0	0.38	XYL 1		
27735062	**Low discrimination**					**27735062**
	Actinomyces naeslundii	59.00	0.35	XYL 1		
	Actinomyces israelii	27.43	0.20			
	Clostridium perfringens	13.57	0.02			
27735063	**Good identification**					**27735063**
	Actinomyces naeslundii	100.0	0.38	XYL 1		
27735100	**Low discrimination**					**27735100**
					COC	
	Actinomyces israelii	71.07	0.55	CEL 93	1	
	Actinomyces naeslundii	22.95	0.56		1	
	Atopobium parvulum	3.30	0.18		99	
27735101	**Very good identification**					**27735101**
	Actinomyces naeslundii	97.96	0.59	XYL 1		
	Actinomyces israelii	2.04	0.21			
27735102	**Low discrimination**					**27735102**
	Actinomyces israelii	64.07	0.58	CEL 93		
	Actinomyces viscosus	16.83	0.33			
	Actinomyces naeslundii	13.79	0.56			
27735103	**Low discrimination**					**27735103**
	Actinomyces naeslundii	76.78	0.59	XYL 1		
	Actinomyces viscosus	20.82	0.25			
	Actinomyces israelii	2.40	0.24			
27735110	**Low discrimination**					**27735110**
	Actinomyces israelii	89.79	0.41	CEL 93		
	Actinomyces naeslundii	10.21	0.34	MLZ 13		
27735111	**Good identification**					**27735111**
	Actinomyces naeslundii	100.0	0.37	XYL 1		
27735112	**Low discrimination**					**27735112**
	Actinomyces israelii	58.99	0.44	CEL 93		
	Bifidobacterium adolescentis	30.90	0.33	MLZ 13		
	Actinomyces naeslundii	4.47	0.34			
27735113	**Good identification**					**27735113**
	Actinomyces naeslundii	100.0	0.37	XYL 1		
27735121	**Very good identification**					**27735121**
	Actinomyces naeslundii	99.22	0.59	XYL 1		
	Clostridium perfringens	0.78	0.01			
27735122	**Low discrimination**					**27735122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		62.19	0.51	NIT 1	1	
Actinomyces naeslundii		19.68	0.56		99	
Clostridium perfringens		10.80	0.29		99	
27735123	**Very good identification**					**27735123**
					GRA	
Actinomyces naeslundii		97.92	0.59	XYL 1	99	
Parabacteroides distasonis		2.08	0.17		1	
27735131	**Good identification**					**27735131**
Actinomyces naeslundii		100.0	0.37	XYL 1		
27735132	**Acceptable identification**					**27735132**
					GRA	
Parabacteroides distasonis		92.72	0.47	NIT 1	1	
Actinomyces naeslundii		2.87	0.34		99	
Actinomyces israelii		2.41	0.23		99	
27735133	**Low discrimination**					**27735133**
					GRA	
Actinomyces naeslundii		82.12	0.37	XYL 1	99	
Parabacteroides distasonis		17.88	0.13	MLZ 5	1	
27735140	**Low discrimination**					**27735140**
Actinomyces israelii		87.22	0.48	CEL 93		
Actinomyces naeslundii		12.78	0.43			
27735141	**Good identification**					**27735141**
Actinomyces naeslundii		95.61	0.46	XYL 1		
Actinomyces israelii		4.39	0.14			
27735142	**Low discrimination**					**27735142**
Actinomyces israelii		88.90	0.51	CEL 93		
Actinomyces naeslundii		8.69	0.43			
Bifidobacterium adolescentis		1.81	0.16			
27735143	**Acceptable identification**					**27735143**
Actinomyces naeslundii		93.55	0.46	XYL 1		
Actinomyces israelii		6.45	0.17			
27735152	**Low discrimination**					**27735152**
Actinomyces israelii		85.73	0.37	CEL 93		
Bifidobacterium adolescentis		14.27	0.17	MLZ 13		
27735161	**Good identification**					**27735161**
Actinomyces naeslundii		100.0	0.46	XYL 1		
27735162	**Low discrimination**					**27735162**
Actinomyces naeslundii		41.21	0.43	XYL 1		
Actinomyces israelii		26.93	0.31			
Clostridium perfringens		24.41	0.17			
27735163	**Good identification**					**27735163**
Actinomyces naeslundii		100.0	0.46	XYL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27735300	**Acceptable identification**						**27735300**
Actinomyces israelii		94.95	0.36	CEL 93			
Actinomyces odontolyticus		5.05	0.06	RHA 7			
27735302	**Low discrimination**						**27735302**
					GRA		
Actinomyces israelii		67.64	0.39	CEL 93	99		
Parabacteroides distasonis		18.53	0.25	RHA 7	1		
Clostridium clostridioforme		11.45	0.16		50		
27735322	**Very good identification**						**27735322**
					GRA		
Parabacteroides distasonis		98.94	0.60	NIT 1	1		
Clostridium clostridioforme		0.62	0.16		50		
Actinomyces israelii		0.23	0.18		99		
27735332	**Acceptable identification**						**27735332**
Parabacteroides distasonis		94.98	0.55	NIT 1			
Clostridium clostridioforme		5.02	0.27				
27735400	**Good identification**						**27735400**
					SPO		
Actinomyces israelii		98.00	0.43	CEL 93	1		
Clostridium butyricum		2.01	-.07		99		
27735402	**Acceptable identification**						**27735402**
Actinomyces israelii		94.26	0.46	CEL 93			
Clostridium clostridioforme		3.75	0.13				
Bifidobacterium adolescentis		2.00	0.11				
27735440	**Good identification**						**27735440**
Actinomyces israelii		100.0	0.36	CEL 93			
27735442	**Good identification**						**27735442**
Actinomyces israelii		100.0	0.39	CEL 93			
27735500	**Low discrimination**						**27735500**
					GRA	SPO	
Actinomyces israelii		83.10	0.54	CEL 93	99	1	
Clostridium butyricum		10.20	0.17		99	99	
Mitsuokella multacida		5.57	0.21		1	1	
27735502	**Low discrimination**						**27735502**
Actinomyces israelii		62.27	0.57	CEL 93			
Bifidobacterium adolescentis		32.61	0.45				
Clostridium clostridioforme		2.07	0.22				
27735510	**Low discrimination**						**27735510**
					GRA		
Actinomyces israelii		75.65	0.39	CEL 93	99		
Mitsuokella multacida		11.30	0.13	MLZ 13	1		
Bifidobacterium adolescentis		4.91	0.13		99		
27735512	**Low discrimination**						**27735512**
Bifidobacterium adolescentis		67.56	0.47	NIT 1			
Clostridium clostridioforme		16.00	0.33				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		15.77	0.42				
27735522	**Low discrimination**						**27735522**
					GRA		
Parabacteroides distasonis		81.27	0.48	NIT 1	1		
Actinomyces israelii		10.18	0.36		99		
Clostridium clostridioforme		5.31	0.22		50		
27735532	**Low discrimination**						**27735532**
					GRA		
Parabacteroides distasonis		58.69	0.43	NIT 1	1		
Clostridium clostridioforme		32.41	0.33		50		
Bifidobacterium longum subsp. longum		5.48	0.21		99		
27735540	**Low discrimination**						**27735540**
					GRA	SPO	
Actinomyces israelii		83.35	0.47	CEL 93	99	1	
Mitsuokella multacida		14.36	0.21		1	1	
Clostridium butyricum		2.29	-.01		99	99	
27735542	**Low discrimination**						**27735542**
					GRA		
Actinomyces israelii		84.80	0.50	CEL 93	99		
Bifidobacterium adolescentis		14.12	0.30		99		
Mitsuokella multacida		1.08	0.04		1		
27735612	**Good identification**						**27735612**
Clostridium clostridioforme		100.0	0.39	NIT 1			
27735632	**Good identification**						**27735632**
Clostridium clostridioforme		97.56	0.39	NIT 1			
Parabacteroides distasonis		2.44	0.17				
27735702	**Low discrimination**						**27735702**
					GRA		
Clostridium clostridioforme		69.87	0.37	NIT 1	50		
Actinomyces israelii		21.55	0.37		99		
Parabacteroides distasonis		4.42	0.22		1		
27735712	**Good identification**						**27735712**
					GRA		
Clostridium clostridioforme		97.45	0.48	NIT 1	50		
Actinomyces israelii		0.99	0.23		99		
Parabacteroides distasonis		0.73	0.17		1		
27735722	**Low discrimination**						**27735722**
					GRA		
Parabacteroides distasonis		85.68	0.56	NIT 1	1		
Clostridium clostridioforme		13.67	0.37		50		
Actinomyces israelii		0.27	0.17		99		
27735732	**Low discrimination**						**27735732**
Clostridium clostridioforme		57.45	0.48	NIT 1			
Parabacteroides distasonis		42.55	0.51				
27736101	**Good identification**						**27736101**
Actinomyces naeslundii		100.0	0.40	GAL 96			
27736102	**Low discrimination**						**27736102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		62.85	0.37	CEL 93		
Actinomyces naeslundii		19.12	0.37	GAL 99		
Propionibacterium propionicum		13.28	0.24			
27736103	**Good identification**					**27736103**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27736120	**Low discrimination**					**27736120**
Actinomyces naeslundii		87.73	0.37	GAL 96		
Actinomyces israelii		12.27	0.13	XYL 1		
27736121	**Good identification**					**27736121**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27736122	**Low discrimination**					**27736122**
Actinomyces naeslundii		82.65	0.37	GAL 96		
Actinomyces israelii		17.35	0.16	XYL 1		
27736123	**Good identification**					**27736123**
Actinomyces naeslundii		100.0	0.40	GAL 96		
27736502	**Acceptable identification**					**27736502**
Actinomyces israelii		92.97	0.35	CEL 93		
Bifidobacterium adolescentis		7.03	0.10	GAL 99		
27737000	**Low discrimination**					**27737000**
Actinomyces israelii		86.55	0.58	CEL 93		
Actinomyces naeslundii		13.46	0.53			
27737001	**Good identification**					**27737001**
Actinomyces naeslundii		95.85	0.56	XYL 1		
Actinomyces israelii		4.15	0.23			
27737002	**Good identification**					**27737002**
Actinomyces israelii		90.61	0.61	CEL 93		
Actinomyces naeslundii		9.39	0.53			
27737003	**Acceptable identification**					**27737003**
Actinomyces naeslundii		93.90	0.56	XYL 1		
Actinomyces israelii		6.10	0.26			
27737010	**Acceptable identification**					**27737010**
Actinomyces israelii		94.81	0.44	CEL 93		
Actinomyces naeslundii		5.19	0.31	MLZ 13		
27737012	**Good identification**					**27737012**
Actinomyces israelii		96.48	0.47	CEL 93		
Actinomyces naeslundii		3.52	0.31	MLZ 13		
27737020	**Low discrimination**					**27737020**
Actinomyces naeslundii		70.89	0.53	XYL 1		
Actinomyces israelii		29.11	0.37			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27737021	**Good identification**					**27737021**
Actinomyces naeslundii		100.0	0.56	XYL 1		
27737022	**Low discrimination**					**27737022**
Actinomyces naeslundii		61.89	0.53	XYL 1		
Actinomyces israelii		38.11	0.40			
27737023	**Good identification**					**27737023**
Actinomyces naeslundii		100.0	0.56	XYL 1		
27737040	**Acceptable identification**					**27737040**
Actinomyces israelii		93.41	0.51	CEL 93		
Actinomyces naeslundii		6.59	0.40			
27737041	**Acceptable identification**					**27737041**
Actinomyces naeslundii		91.29	0.43	XYL 1		
Actinomyces israelii		8.71	0.16			
27737042	**Good identification**					**27737042**
Actinomyces israelii		95.51	0.54	CEL 93		
Actinomyces naeslundii		4.49	0.40			
27737043	**Low discrimination**					**27737043**
Actinomyces naeslundii		87.48	0.43	XYL 1		
Actinomyces israelii		12.52	0.19			
27737050	**Good identification**					**27737050**
Actinomyces israelii		100.0	0.36	CEL 93		
27737052	**Good identification**					**27737052**
Actinomyces israelii		100.0	0.39	CEL 93		
27737060	**Low discrimination**					**27737060**
Actinomyces naeslundii		52.50	0.40	XYL 1		
Actinomyces israelii		47.50	0.30			
27737061	**Good identification**					**27737061**
Actinomyces naeslundii		100.0	0.43	XYL 1		
27737063	**Good identification**					**27737063**
Actinomyces naeslundii		100.0	0.43	XYL 1		
27737100	**Low discrimination**					**27737100**
Actinomyces israelii		89.87	0.68	CEL 93		
Actinomyces naeslundii		9.94	0.61			
Propionibacterium propionicum		0.19	0.21			
27737101	**Good identification**					**27737101**
Actinomyces naeslundii		94.26	0.64	XYL 1		
Actinomyces israelii		5.74	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27737102	**Good identification**					**27737102**
Actinomyces israelii		92.12	0.71	CEL 93		
Actinomyces naeslundii		6.79	0.61			
Bifidobacterium adolescentis		0.85	0.30			
27737103	**Good identification**					**27737103**
Actinomyces naeslundii		91.63	0.64	XYL 1		
Actinomyces israelii		8.37	0.37			
27737110	**Good identification**					**27737110**
Actinomyces israelii		96.25	0.54	CEL 93		
Actinomyces naeslundii		3.75	0.39	MLZ 13		
27737111	**Low discrimination**					**27737111**
Actinomyces naeslundii		85.27	0.42	XYL 1		
Actinomyces israelii		14.74	0.20	MLZ 5		
27737112	**Good identification**					**27737112**
Actinomyces israelii		90.78	0.57	CEL 93		
Bifidobacterium adolescentis		6.87	0.31	MLZ 13		
Actinomyces naeslundii		2.36	0.39			
27737113	**Low discrimination**					**27737113**
Actinomyces naeslundii		79.41	0.42	XYL 1		
Actinomyces israelii		20.59	0.23	MLZ 5		
27737120	**Low discrimination**					**27737120**
Actinomyces naeslundii		63.17	0.61	XYL 1		
Actinomyces israelii		36.45	0.47			
Clostridium perfringens		0.38	0.01			
27737121	**Very good identification**					**27737121**
Actinomyces naeslundii		99.61	0.64	XYL 1		
Actinomyces israelii		0.39	0.13			
27737122	**Low discrimination**					**27737122**
					GRA	
Actinomyces naeslundii		53.14	0.61	XYL 1	99	
Actinomyces israelii		45.99	0.50		99	
Parabacteroides distasonis		0.87	0.17		1	
27737123	**Very good identification**					**27737123**
Actinomyces naeslundii		99.42	0.64	XYL 1		
Actinomyces israelii		0.58	0.16			
27737131	**Good identification**					**27737131**
Actinomyces naeslundii		100.0	0.42	XYL 1		
27737132	**Low discrimination**					**27737132**
					GRA	
Actinomyces israelii		67.78	0.36	CEL 93	99	
Actinomyces naeslundii		27.58	0.39	MLZ 13	99	
Parabacteroides distasonis		4.64	0.13	NAG 6	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27737133	**Good identification**					**27737133**
Actinomyces naeslundii		100.0	0.42	XYL 1		
27737140	**Good identification**					**27737140**
Actinomyces israelii		94.73	0.61	CEL 93		
Actinomyces naeslundii		4.76	0.48			
Propionibacterium propionicum		0.52	0.21			
27737141	**Low discrimination**					**27737141**
Actinomyces naeslundii		88.17	0.51	XYL 1		
Actinomyces israelii		11.83	0.27			
27737142	**Very good identification**					**27737142**
Actinomyces israelii		95.98	0.64	CEL 93		
Actinomyces naeslundii		3.21	0.48			
Propionibacterium propionicum		0.53	0.24			
27737143	**Low discrimination**					**27737143**
Actinomyces naeslundii		83.25	0.51	XYL 1		
Actinomyces israelii		16.75	0.30			
27737150	**Good identification**					**27737150**
Actinomyces israelii		98.26	0.47	CEL 93		
Actinomyces naeslundii		1.74	0.26	MLZ 13		
27737152	**Good identification**					**27737152**
Actinomyces israelii		96.54	0.50	CEL 93		
Bifidobacterium adolescentis		2.32	0.16	MLZ 13		
Actinomyces naeslundii		1.14	0.26			
27737160	**Low discrimination**					**27737160**
Actinomyces israelii		55.98	0.40	CEL 93		
Actinomyces naeslundii		44.02	0.48	NAG 6		
27737161	**Good identification**					**27737161**
Actinomyces naeslundii		100.0	0.51	XYL 1		
27737162	**Low discrimination**					**27737162**
Actinomyces israelii		65.60	0.43	CEL 93		
Actinomyces naeslundii		34.40	0.48	NAG 6		
27737163	**Good identification**					**27737163**
Actinomyces naeslundii		100.0	0.51	XYL 1		
27737200	**Good identification**					**27737200**
Actinomyces israelii		100.0	0.38	CEL 93		
27737202	**Good identification**					**27737202**
Actinomyces israelii		100.0	0.41	CEL 93		
27737300	**Good identification**					**27737300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		98.54	0.49	CEL 93			
Actinomyces naeslundii		1.46	0.27	RHA 7			
27737302	**Good identification**						**27737302**
Actinomyces israelii		99.02	0.52	CEL 93			
Actinomyces naeslundii		0.98	0.27	RHA 7			
27737312	**Good identification**						**27737312**
Actinomyces israelii		100.0	0.38	CEL 93			
27737340	**Good identification**						**27737340**
Actinomyces israelii		100.0	0.42	CEL 93			
27737342	**Good identification**						**27737342**
Actinomyces israelii		100.0	0.45	CEL 93			
27737400	**Very good identification**						**27737400**
Actinomyces israelii		100.0	0.56	CEL 93			
27737402	**Very good identification**						**27737402**
Actinomyces israelii		99.70	0.59	CEL 93			
Bifidobacterium adolescentis		0.31	0.10				
27737410	**Good identification**						**27737410**
Actinomyces israelii		100.0	0.42	CEL 93			
27737412	**Good identification**						**27737412**
Actinomyces israelii		97.56	0.45	CEL 93			
Bifidobacterium adolescentis		2.44	0.11	MLZ 13			
27737420	**Good identification**						**27737420**
Actinomyces israelii		100.0	0.36	CEL 93			
27737422	**Good identification**						**27737422**
Actinomyces israelii		100.0	0.39	CEL 93			
27737440	**Good identification**						**27737440**
Actinomyces israelii		100.0	0.49	CEL 93			
27737442	**Good identification**						**27737442**
Actinomyces israelii		100.0	0.52	CEL 93			
27737452	**Good identification**						**27737452**
Actinomyces israelii		100.0	0.38	CEL 93			
27737500	**Very good identification**						**27737500**
					GRA	SPO	
Actinomyces israelii		96.85	0.66	CEL 93	99	1	
Mitsuokella multacida		2.67	0.27		1	1	
Clostridium butyricum		0.23	0.01		99	99	
27737502	**Good identification**						**27737502**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		92.66	0.69	CEL 93	99		
Bifidobacterium adolescentis		7.01	0.44		99		
Mitsuokella multacida		0.19	0.11		1		
27737503	**Low discrimination**						**27737503**
Actinomyces israelii		82.59	0.35	CEL 93			
Actinomyces naeslundii		11.17	0.30	URE 1			
Bifidobacterium adolescentis		6.25	0.10				
27737510	**Acceptable identification**						**27737510**
					GRA		
Actinomyces israelii		93.38	0.52	CEL 93	99		
Mitsuokella multacida		5.75	0.19	MLZ 13	1		
Bifidobacterium adolescentis		0.88	0.11		99		
27737512	**Low discrimination**						**27737512**
Actinomyces israelii		61.37	0.55	CEL 93			
Bifidobacterium adolescentis		37.96	0.45	MLZ 13			
Bifidobacterium dentium		0.38	-.07				
27737520	**Low discrimination**						**27737520**
					GRA		
Actinomyces israelii		88.54	0.46	CEL 93	99		
Mitsuokella multacida		9.57	0.17	NAG 6	1		
Actinomyces naeslundii		1.89	0.27		99		
27737522	**Good identification**						**27737522**
					GRA		
Actinomyces israelii		96.11	0.49	CEL 93	99		
Actinomyces naeslundii		1.37	0.27	NAG 6	99		
Parabacteroides distasonis		1.37	0.14		1		
27737540	**Good identification**						**27737540**
					GRA		
Actinomyces israelii		93.38	0.59	CEL 93	99		
Mitsuokella multacida		6.62	0.27		1		
27737542	**Very good identification**						**27737542**
					GRA		
Actinomyces israelii		97.15	0.62	CEL 93	99		
Bifidobacterium adolescentis		2.34	0.28		99		
Mitsuokella multacida		0.51	0.11		1		
27737550	**Low discrimination**						**27737550**
					COC	GRA	
Actinomyces israelii		84.61	0.45	CEL 93	1	99	
Mitsuokella multacida		13.39	0.19	MLZ 13	1	1	
Blautia producta		2.01	-.03		99	99	
27737552	**Low discrimination**						**27737552**
					GRA		
Actinomyces israelii		82.76	0.48	CEL 93	99		
Bifidobacterium adolescentis		16.27	0.30	MLZ 13	99		
Mitsuokella multacida		0.97	0.03		1		
27737560	**Low discrimination**						**27737560**
					GRA		
Actinomyces israelii		78.26	0.39	CEL 93	99		
Mitsuokella multacida		21.74	0.17	NAG 6	1		
27737562	**Good identification**						**27737562**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.42	CEL 93		
27737600	**Good identification**					**27737600**
Actinomyces israelii		100.0	0.37	CEL 93		
27737602	**Good identification**					**27737602**
Actinomyces israelii		100.0	0.40	CEL 93		
27737700	**Low discrimination**					**27737700**
					GRA	
Actinomyces israelii		59.51	0.47	CEL 93	99	
Mitsuokella multacida		40.50	0.32	RHA 7	1	
27737702	**Acceptable identification**					**27737702**
					GRA	
Actinomyces israelii		94.29	0.50	CEL 93	99	
Mitsuokella multacida		4.75	0.15	RHA 7	1	
Bifidobacterium adolescentis		0.96	0.10		99	
27737712	**Low discrimination**					**27737712**
					GRA	
Actinomyces israelii		72.91	0.36	CEL 93	99	
Clostridium clostridioforme		12.84	0.14	RHA 7	50	
Mitsuokella multacida		8.20	0.07	MLZ 13	1	
27737742	**Low discrimination**					**27737742**
					GRA	
Actinomyces israelii		88.53	0.43	CEL 93	99	
Mitsuokella multacida		11.48	0.15	RHA 7	1	
27740001	**Low discrimination**					**27740001**
Actinomyces naeslundii		52.97	0.37	FRU 95		
Actinomyces viscosus		47.03	0.12	GAL 96		
27740002	**Good identification**					**27740002**
Actinomyces viscosus		99.75	0.54	FRU 95		
Actinomyces naeslundii		0.25	0.34			
27740003	**Good identification**					**27740003**
Actinomyces viscosus		98.88	0.46	FRU 95		
Actinomyces naeslundii		1.13	0.37			
27740021	**Good identification**					**27740021**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27740023	**Low discrimination**					**27740023**
Actinomyces naeslundii		52.97	0.37	FRU 95		
Actinomyces viscosus		47.03	0.12	GAL 96		
27740101	**Low discrimination**					**27740101**
Actinomyces naeslundii		64.33	0.45	FRU 95		
Actinomyces viscosus		35.68	0.16	GAL 96		
27740102	**Very good identification**					**27740102**
Actinomyces viscosus		99.60	0.58	FRU 95		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.40	0.42			
27740103	**Good identification**					**27740103**
Actinomyces viscosus		98.21	0.50	FRU 95		
Actinomyces naeslundii		1.79	0.45			
27740120	**Good identification**					**27740120**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27740121	**Good identification**					**27740121**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27740123	**Low discrimination**					**27740123**
Actinomyces naeslundii		64.33	0.45	FRU 95		
Actinomyces viscosus		35.68	0.16	GAL 96		
27741000	**Good identification**					**27741000**
Actinomyces naeslundii		90.74	0.58	FRU 95		
Actinomyces viscosus		9.26	0.16			
27741001	**Very good identification**					**27741001**
Actinomyces naeslundii		97.78	0.61	FRU 95		
Actinomyces viscosus		2.22	0.08			
27741002	**Acceptable identification**					**27741002**
Actinomyces viscosus		90.99	0.51	FRU 95		
Actinomyces naeslundii		9.01	0.58			
27741003	**Low discrimination**					**27741003**
Actinomyces viscosus		69.18	0.42	FRU 95		
Actinomyces naeslundii		30.82	0.61			
27741010	**Good identification**					**27741010**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27741011	**Good identification**					**27741011**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27741013	**Low discrimination**					**27741013**
Actinomyces naeslundii		69.89	0.39	FRU 95		
Actinomyces viscosus		30.11	0.08	MLZ 5		
27741020	**Very good identification**					**27741020**
Actinomyces naeslundii		100.0	0.58	FRU 95		
27741021	**Very good identification**					**27741021**
Actinomyces naeslundii		100.0	0.61	FRU 95		
27741022	**Low discrimination**					**27741022**
					GRA	
Actinomyces naeslundii		89.83	0.58	FRU 95	99	
Actinomyces viscosus		9.17	0.16		99	
Prevotella oralis		1.00	-.02		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27741023	**Very good identification**					**27741023**
Actinomyces naeslundii		97.78	0.61	FRU 95		
Actinomyces viscosus		2.22	0.08			
27741030	**Good identification**					**27741030**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27741031	**Good identification**					**27741031**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27741032	**Good identification**					**27741032**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27741033	**Good identification**					**27741033**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27741040	**Good identification**					**27741040**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27741041	**Good identification**					**27741041**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27741042	**Low discrimination**					**27741042**
Actinomyces naeslundii		63.36	0.45	FRU 95		
Actinomyces viscosus		36.64	0.16			
27741043	**Low discrimination**					**27741043**
Actinomyces naeslundii		88.61	0.48	FRU 95		
Actinomyces viscosus		11.39	0.08			
27741060	**Good identification**					**27741060**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27741061	**Good identification**					**27741061**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27741062	**Good identification**					**27741062**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27741063	**Good identification**					**27741063**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27741100	**Good identification**					**27741100**
Actinomyces naeslundii		94.01	0.65	FRU 95		
Actinomyces viscosus		5.99	0.21			
27741101	**Very good identification**					**27741101**
Actinomyces naeslundii		98.60	0.68	FRU 95		
Actinomyces viscosus		1.40	0.12			
27741102	**Low discrimination**					**27741102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		86.04	0.55	FRU 95		
Actinomyces naeslundii		13.64	0.65			
Lactobacillus acidophilus		0.23	0.10			
27741103	**Low discrimination**					**27741103**
Actinomyces viscosus		58.37	0.47	FRU 95		
Actinomyces naeslundii		41.63	0.68			
27741110	**Good identification**					**27741110**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27741111	**Good identification**					**27741111**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27741113	**Low discrimination**					**27741113**
Actinomyces naeslundii		78.80	0.46	FRU 95		
Actinomyces viscosus		21.20	0.12	MLZ 5		
27741120	**Very good identification**					**27741120**
					GRA	
Actinomyces naeslundii		98.29	0.65	FRU 95	99	
Clostridium perfringens		0.72	0.06		99	
Prevotella melaninogenica		0.68	-.01		1	
27741121	**Very good identification**					**27741121**
Actinomyces naeslundii		100.0	0.68	FRU 95		
27741122	**Low discrimination**					**27741122**
					GRA	
Actinomyces naeslundii		87.32	0.65	FRU 95	99	
Actinomyces viscosus		5.57	0.21		99	
Prevotella oralis		5.35	0.19		1	
27741123	**Very good identification**					**27741123**
Actinomyces naeslundii		98.60	0.68	FRU 95		
Actinomyces viscosus		1.40	0.12			
27741130	**Good identification**					**27741130**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27741131	**Good identification**					**27741131**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27741132	**Good identification**					**27741132**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27741133	**Good identification**					**27741133**
Actinomyces naeslundii		100.0	0.46	FRU 95		
27741140	**Good identification**					**27741140**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27741141	**Good identification**					**27741141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	FRU 95		
27741142	**Low discrimination**					**27741142**
Actinomyces naeslundii		73.47	0.52	FRU 95		
Actinomyces viscosus		26.53	0.21			
27741143	**Acceptable identification**					**27741143**
Actinomyces naeslundii		92.57	0.55	FRU 95		
Actinomyces viscosus		7.43	0.12			
27741160	**Good identification**					**27741160**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27741161	**Good identification**					**27741161**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27741162	**Good identification**					**27741162**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27741163	**Good identification**					**27741163**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27742000	**Good identification**					**27742000**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27742001	**Good identification**					**27742001**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27742003	**Low discrimination**					**27742003**
Actinomyces naeslundii		68.62	0.42	FRU 95		
Actinomyces viscosus		31.38	0.12	GAL 96		
27742020	**Good identification**					**27742020**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27742021	**Good identification**					**27742021**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27742022	**Good identification**					**27742022**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27742023	**Good identification**					**27742023**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27742100	**Good identification**					**27742100**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27742101	**Good identification**					**27742101**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27742103	**Low discrimination**					**27742103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		77.78	0.50	FRU 95		
Actinomyces viscosus		22.22	0.16	GAL 96		
27742120	**Good identification**					**27742120**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27742121	**Good identification**					**27742121**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27742122	**Good identification**					**27742122**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27742123	**Good identification**					**27742123**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27742141	**Good identification**					**27742141**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27742143	**Good identification**					**27742143**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27742161	**Good identification**					**27742161**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27742163	**Good identification**					**27742163**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27743000	**Very good identification**					**27743000**
Actinomyces naeslundii		99.44	0.62	FRU 95		
Actinomyces israelii		0.56	0.15			
27743001	**Very good identification**					**27743001**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27743002	**Good identification**					**27743002**
Actinomyces naeslundii		94.25	0.62	FRU 95		
Actinomyces viscosus		4.95	0.16			
Actinomyces israelii		0.79	0.18			
27743003	**Very good identification**					**27743003**
Actinomyces naeslundii		98.85	0.65	FRU 95		
Actinomyces viscosus		1.16	0.08			
27743010	**Good identification**					**27743010**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27743011	**Good identification**					**27743011**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27743012	**Good identification**					**27743012**
Actinomyces naeslundii		100.0	0.41	FRU 95		

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
27743013	**Good identification**					**27743013**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27743020	**Very good identification**					**27743020**
Actinomyces naeslundii		100.0	0.62	FRU 95		
27743021	**Very good identification**					**27743021**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27743022	**Very good identification**					**27743022**
Actinomyces naeslundii		100.0	0.62	FRU 95		
27743023	**Very good identification**					**27743023**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27743030	**Good identification**					**27743030**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27743031	**Good identification**					**27743031**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27743032	**Good identification**					**27743032**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27743033	**Good identification**					**27743033**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27743040	**Good identification**					**27743040**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27743041	**Good identification**					**27743041**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27743042	**Good identification**					**27743042**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27743043	**Good identification**					**27743043**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27743060	**Good identification**					**27743060**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27743061	**Good identification**					**27743061**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27743062	**Good identification**					**27743062**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27743063	**Good identification**					**27743063**
Actinomyces naeslundii		100.0	0.53	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27743100	**Very good identification**					**27743100**
Actinomyces naeslundii		99.22	0.70	FRU 95		
Actinomyces israelii		0.78	0.25			
27743101	**Very good identification**					**27743101**
Actinomyces naeslundii		100.0	0.73	FRU 95		
27743102	**Very good identification**					**27743102**
Actinomyces naeslundii		95.73	0.70	FRU 95		
Actinomyces viscosus		3.14	0.21			
Actinomyces israelii		1.13	0.28			
27743103	**Very good identification**					**27743103**
Actinomyces naeslundii		99.28	0.73	FRU 95		
Actinomyces viscosus		0.72	0.12			
27743110	**Good identification**					**27743110**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27743111	**Good identification**					**27743111**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27743112	**Good identification**					**27743112**
Actinomyces naeslundii		96.75	0.48	FRU 95		
Actinomyces israelii		3.25	0.14	MLZ 5		
27743113	**Good identification**					**27743113**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27743120	**Very good identification**					**27743120**
Actinomyces naeslundii		100.0	0.70	FRU 95		
27743121	**Very good identification**					**27743121**
Actinomyces naeslundii		100.0	0.73	FRU 95		
27743122	**Very good identification**					**27743122**
					SPO	
Actinomyces naeslundii		99.56	0.70	FRU 95	1	
Clostridium ramosum		0.44	0.05		92	
27743123	**Very good identification**					**27743123**
Actinomyces naeslundii		100.0	0.73	FRU 95		
27743130	**Good identification**					**27743130**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27743131	**Good identification**					**27743131**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27743132	**Good identification**					**27743132**
Actinomyces naeslundii		100.0	0.48	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27743133	**Good identification**					**27743133**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27743140	**Very good identification**					**27743140**
Actinomyces naeslundii		98.29	0.57	FRU 95		
Actinomyces israelii		1.71	0.18			
27743141	**Very good identification**					**27743141**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27743142	**Very good identification**					**27743142**
Actinomyces naeslundii		97.46	0.57	FRU 95		
Actinomyces israelii		2.54	0.21			
27743143	**Very good identification**					**27743143**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27743150	**Good identification**					**27743150**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27743151	**Good identification**					**27743151**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27743152	**Good identification**					**27743152**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27743153	**Good identification**					**27743153**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27743160	**Very good identification**					**27743160**
Actinomyces naeslundii		100.0	0.57	FRU 95		
27743161	**Very good identification**					**27743161**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27743162	**Very good identification**					**27743162**
Actinomyces naeslundii		100.0	0.57	FRU 95		
27743163	**Very good identification**					**27743163**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27743170	**Good identification**					**27743170**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27743171	**Good identification**					**27743171**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27743172	**Good identification**					**27743172**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27743173	**Good identification**					**27743173**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	FRU 95		
27743300	**Good identification**					**27743300**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27743301	**Good identification**					**27743301**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27743302	**Good identification**					**27743302**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27743303	**Good identification**					**27743303**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27743320	**Good identification**					**27743320**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27743321	**Good identification**					**27743321**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27743322	**Low discrimination**					**27743322**
					SPO	
Actinomyces naeslundii		63.45	0.36	FRU 95	1	
Clostridium ramosum		36.56	0.07	RHA 1	92	
27743323	**Good identification**					**27743323**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27743500	**Low discrimination**					**27743500**
Actinomyces naeslundii		61.04	0.36	FRU 95		
Actinomyces israelii		38.96	0.24	ARA 1		
27743501	**Good identification**					**27743501**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27743502	**Low discrimination**					**27743502**
Actinomyces naeslundii		51.09	0.36	FRU 95		
Actinomyces israelii		48.92	0.27	ARA 1		
27743503	**Good identification**					**27743503**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27743520	**Good identification**					**27743520**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27743521	**Good identification**					**27743521**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27743522	**Good identification**					**27743522**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27743523	**Good identification**					**27743523**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	FRU 95		
27747002	**Acceptable identification**					**27747002**
Actinomyces israelii		90.54	0.36	FRU 99		
Actinomyces naeslundii		9.46	0.28			
27747100	**Low discrimination**					**27747100**
Actinomyces israelii		89.97	0.43	FRU 99		
Actinomyces naeslundii		10.03	0.36			
27747101	**Good identification**					**27747101**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27747102	**Acceptable identification**					**27747102**
Actinomyces israelii		93.08	0.46	FRU 99		
Actinomyces naeslundii		6.92	0.36			
27747103	**Good identification**					**27747103**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27747120	**Low discrimination**					**27747120**
Actinomyces naeslundii		63.58	0.36	FRU 95		
Actinomyces israelii		36.42	0.23	XYL 1		
27747121	**Good identification**					**27747121**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27747122	**Low discrimination**					**27747122**
Actinomyces naeslundii		53.79	0.36	FRU 95		
Actinomyces israelii		46.21	0.26	XYL 1		
27747123	**Good identification**					**27747123**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27747140	**Good identification**					**27747140**
Actinomyces israelii		95.19	0.36	FRU 99		
Actinomyces naeslundii		4.81	0.23			
27747142	**Good identification**					**27747142**
Actinomyces israelii		96.74	0.39	FRU 99		
Actinomyces naeslundii		3.26	0.23			
27747500	**Good identification**					**27747500**
Actinomyces israelii		100.0	0.42	FRU 99		
27747502	**Acceptable identification**					**27747502**
Actinomyces israelii		93.61	0.45	FRU 99		
Bifidobacterium adolescentis		6.39	0.18			
27747542	**Good identification**					**27747542**
Actinomyces israelii		100.0	0.38	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27750000	**Low discrimination**					**27750000**
					GRA	
Actinomyces viscosus		76.79	0.42	BGL 99	99	
Actinomyces naeslundii		19.22	0.56		99	
Fusobacterium mortiferum		2.80	0.30		1	
27750001	**Low discrimination**					**27750001**
Actinomyces naeslundii		52.97	0.59	GAL 96		
Actinomyces viscosus		47.03	0.34			
27750002	**Very good identification**					**27750002**
Actinomyces viscosus		99.73	0.76			
Actinomyces naeslundii		0.25	0.56			
Actinomyces odontolyticus		0.01	0.14			
27750003	**Very good identification**					**27750003**
Actinomyces viscosus		98.88	0.68			
Actinomyces naeslundii		1.13	0.59			
27750011	**Low discrimination**					**27750011**
Actinomyces naeslundii		85.44	0.37	GAL 96		
Actinomyces viscosus		14.56	-.01	MLZ 5		
27750012	**Good identification**					**27750012**
Actinomyces viscosus		98.70	0.42	MLZ 1		
Actinomyces naeslundii		1.30	0.34			
27750020	**Good identification**					**27750020**
Actinomyces naeslundii		96.12	0.56	GAL 96		
Actinomyces viscosus		3.88	0.08			
27750021	**Very good identification**					**27750021**
Actinomyces naeslundii		99.11	0.59	GAL 96		
Actinomyces viscosus		0.89	-.01			
27750022	**Low discrimination**					**27750022**
					GRA	
Actinomyces viscosus		79.76	0.42	NAG 1	99	
Actinomyces naeslundii		19.96	0.56		99	
Prevotella oralis		0.28	-.02		1	
27750023	**Low discrimination**					**27750023**
Actinomyces naeslundii		52.97	0.59	GAL 96		
Actinomyces viscosus		47.03	0.34			
27750031	**Good identification**					**27750031**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27750033	**Low discrimination**					**27750033**
Actinomyces naeslundii		85.44	0.37	GAL 96		
Actinomyces viscosus		14.56	-.01	MLZ 5		
27750040	**Low discrimination**					**27750040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		81.39	0.43	GAL 96		
Actinomyces viscosus		18.61	0.08			
27750041	**Good identification**					**27750041**
Actinomyces naeslundii		95.16	0.46	GAL 96		
Actinomyces viscosus		4.84	-.01			
27750042	**Good identification**					**27750042**
Actinomyces viscosus		95.77	0.42	SOR 1		
Actinomyces naeslundii		4.23	0.43			
27750060	**Good identification**					**27750060**
Actinomyces naeslundii		100.0	0.43	GAL 96		
27750061	**Good identification**					**27750061**
Actinomyces naeslundii		100.0	0.46	GAL 96		
27750062	**Low discrimination**					**27750062**
Actinomyces naeslundii		81.39	0.43	GAL 96		
Actinomyces viscosus		18.61	0.08			
27750063	**Good identification**					**27750063**
Actinomyces naeslundii		95.16	0.46	GAL 96		
Actinomyces viscosus		4.84	-.01			
27750100	**Low discrimination**					**27750100**
					GRA	
Actinomyces viscosus		68.91	0.46	BGL 99	99	
Actinomyces naeslundii		27.61	0.64		99	
Fusobacterium mortiferum		3.29	0.36		1	
27750101	**Low discrimination**					**27750101**
Actinomyces naeslundii		64.33	0.67	GAL 96		
Actinomyces viscosus		35.68	0.38			
27750102	**Very good identification**					**27750102**
Actinomyces viscosus		99.58	0.80			
Actinomyces naeslundii		0.40	0.64			
Lactobacillus acidophilus		0.01	0.10			
27750103	**Very good identification**					**27750103**
Actinomyces viscosus		98.21	0.72			
Actinomyces naeslundii		1.79	0.67			
27750110	**Low discrimination**					**27750110**
Actinomyces naeslundii		67.61	0.42	GAL 96		
Actinomyces viscosus		32.39	0.12	MLZ 5		
27750111	**Acceptable identification**					**27750111**
Actinomyces naeslundii		90.38	0.45	GAL 96		
Actinomyces viscosus		9.62	0.04	MLZ 5		
27750112	**Good identification**					**27750112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		97.94	0.46	MLZ 1		
Actinomyces naeslundii		2.07	0.42			
27750113	**Acceptable identification**					**27750113**
Actinomyces viscosus		91.33	0.38	MLZ 1		
Actinomyces naeslundii		8.67	0.45			
27750120	**Very good identification**					**27750120**
Actinomyces naeslundii		96.29	0.64	GAL 96		
Actinomyces viscosus		2.43	0.12			
Clostridium perfringens		0.89	0.06			
27750121	**Very good identification**					**27750121**
Actinomyces naeslundii		99.44	0.67	GAL 96		
Actinomyces viscosus		0.56	0.04			
27750122	**Low discrimination**					**27750122**
					GRA	
Actinomyces viscosus		69.28	0.46	NAG 1	99	
Actinomyces naeslundii		27.76	0.64		99	
Prevotella oralis		2.15	0.19		1	
27750123	**Low discrimination**					**27750123**
Actinomyces naeslundii		64.33	0.67	GAL 96		
Actinomyces viscosus		35.68	0.38			
27750130	**Good identification**					**27750130**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27750131	**Good identification**					**27750131**
Actinomyces naeslundii		100.0	0.45	GAL 96		
27750132	**Low discrimination**					**27750132**
Actinomyces naeslundii		67.61	0.42	GAL 96		
Actinomyces viscosus		32.39	0.12	MLZ 5		
27750133	**Acceptable identification**					**27750133**
Actinomyces naeslundii		90.38	0.45	GAL 96		
Actinomyces viscosus		9.62	0.04	MLZ 5		
27750140	**Low discrimination**					**27750140**
Actinomyces naeslundii		87.50	0.51	GAL 96		
Actinomyces viscosus		12.50	0.12			
27750141	**Good identification**					**27750141**
Actinomyces naeslundii		96.92	0.54	GAL 96		
Actinomyces viscosus		3.08	0.04			
27750142	**Acceptable identification**					**27750142**
Actinomyces viscosus		93.40	0.46	SOR 1		
Actinomyces naeslundii		6.60	0.51			
27750143	**Low discrimination**					**27750143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		75.86	0.38	SOR 1		
Actinomyces naeslundii		24.14	0.54			
27750160	**Good identification**					**27750160**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27750161	**Good identification**					**27750161**
Actinomyces naeslundii		100.0	0.54	GAL 96		
27750162	**Low discrimination**					**27750162**
Actinomyces naeslundii		87.50	0.51	GAL 96		
Actinomyces viscosus		12.50	0.12			
27750163	**Good identification**					**27750163**
Actinomyces naeslundii		96.92	0.54	GAL 96		
Actinomyces viscosus		3.08	0.04			
27750202	**Good identification**					**27750202**
Actinomyces viscosus		98.18	0.42	RHA 1		
Actinomyces odontolyticus		1.82	0.19			
27750302	**Good identification**					**27750302**
Actinomyces viscosus		99.27	0.46	RHA 1		
Actinomyces naeslundii		0.40	0.29			
Actinomyces odontolyticus		0.33	0.11			
27750303	**Good identification**					**27750303**
Actinomyces viscosus		98.21	0.38	RHA 1		
Actinomyces naeslundii		1.79	0.32			
27750402	**Good identification**					**27750402**
Actinomyces viscosus		100.0	0.42	ARA 1		
27750502	**Good identification**					**27750502**
Actinomyces viscosus		99.43	0.46	ARA 1		
Actinomyces naeslundii		0.40	0.29			
Actinomyces israelii		0.17	0.14			
27750503	**Good identification**					**27750503**
Actinomyces viscosus		98.21	0.38	ARA 1		
Actinomyces naeslundii		1.79	0.32			
27751000	**Low discrimination**					**27751000**
Actinomyces naeslundii		86.76	0.79			
Actinomyces viscosus		8.85	0.38			
Lactobacillus catenaformis		1.72	0.29			
27751001	**Very good identification**					**27751001**
Actinomyces naeslundii		97.78	0.82			
Actinomyces viscosus		2.22	0.30			
27751002	**Good identification**					**27751002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		90.43	0.72			
Actinomyces naeslundii		8.95	0.79			
Lactobacillus acidophilus		0.19	0.26			
27751003	**Low discrimination**					**27751003**
Actinomyces viscosus		69.18	0.64			
Actinomyces naeslundii		30.82	0.82			
27751010	**Good identification**					**27751010**
Actinomyces naeslundii		90.02	0.58	MLZ 5		
Lactobacillus jensenii		5.66	0.15			
Actinomyces israelii		2.56	0.22			
27751011	**Very good identification**					**27751011**
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27751012	**Low discrimination**					**27751012**
Actinomyces viscosus		64.13	0.38	MLZ 1		
Actinomyces naeslundii		33.08	0.58			
Actinomyces israelii		1.41	0.25			
27751013	**Low discrimination**					**27751013**
Actinomyces naeslundii		69.89	0.61	MLZ 5		
Actinomyces viscosus		30.11	0.30			
27751020	**Very good identification**					**27751020**
Actinomyces naeslundii		97.28	0.79			
Clostridium perfringens		1.45	0.26			
Lactobacillus catenaformis		0.48	0.18			
27751021	**Very good identification**					**27751021**
Actinomyces naeslundii		100.0	0.82			
27751022	**Low discrimination**					**27751022**
					GRA	
Actinomyces naeslundii		83.40	0.79		99	
Actinomyces viscosus		8.51	0.38		99	
Prevotella oralis		4.84	0.33		1	
27751023	**Very good identification**					**27751023**
					GRA	
Actinomyces naeslundii		97.75	0.82		99	
Actinomyces viscosus		2.22	0.30		99	
Prevotella oralis		0.04	-.02		1	
27751030	**Very good identification**					**27751030**
Actinomyces naeslundii		98.45	0.58	MLZ 5		
Lactobacillus jensenii		1.55	0.05			
27751031	**Very good identification**					**27751031**
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27751032	**Very good identification**					**27751032**
					GRA	
Actinomyces naeslundii		96.04	0.58	MLZ 5	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		1.88	0.04		99	
Prevotella oralis		1.07	-.02		1	
27751033	**Very good identification**					**27751033**
Actinomyces naeslundii		100.0	0.61	MLZ 5		
27751040	**Very good identification**					**27751040**
Actinomyces naeslundii		97.29	0.67			
Actinomyces israelii		2.15	0.29			
Actinomyces viscosus		0.57	0.04			
27751041	**Very good identification**					**27751041**
Actinomyces naeslundii		100.0	0.70			
27751042	**Low discrimination**					**27751042**
Actinomyces naeslundii		61.49	0.67			
Actinomyces viscosus		35.55	0.38			
Actinomyces israelii		2.03	0.32			
27751043	**Low discrimination**					**27751043**
Actinomyces naeslundii		88.61	0.70			
Actinomyces viscosus		11.39	0.30			
27751050	**Acceptable identification**					**27751050**
Actinomyces naeslundii		94.11	0.45	MLZ 5		
Actinomyces israelii		5.89	0.15			
27751051	**Good identification**					**27751051**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27751052	**Low discrimination**					**27751052**
Actinomyces naeslundii		82.99	0.45	MLZ 5		
Actinomyces viscosus		9.21	0.04			
Actinomyces israelii		7.80	0.18			
27751053	**Good identification**					**27751053**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27751060	**Very good identification**					**27751060**
Actinomyces naeslundii		98.42	0.67			
Clostridium perfringens		1.58	0.13			
27751061	**Very good identification**					**27751061**
Actinomyces naeslundii		100.0	0.70			
27751062	**Very good identification**					**27751062**
Actinomyces naeslundii		96.92	0.67			
Lactobacillus acidophilus		1.46	0.11			
Clostridium perfringens		0.73	0.08			
27751063	**Very good identification**					**27751063**
Actinomyces naeslundii		100.0	0.70			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27751070	**Good identification**					**27751070**
Actinomyces naeslundii		100.0	0.45	MLZ 5		
27751071	**Good identification**					**27751071**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27751072	**Good identification**					**27751072**
Actinomyces naeslundii		100.0	0.45	MLZ 5		
27751073	**Good identification**					**27751073**
Actinomyces naeslundii		100.0	0.48	MLZ 5		
27751100	**Good identification**					**27751100**
Actinomyces naeslundii		90.90	0.87			
Actinomyces viscosus		5.79	0.43			
Actinomyces israelii		1.28	0.46			
27751101	**Very good identification**					**27751101**
Actinomyces naeslundii		98.60	0.90			
Actinomyces viscosus		1.40	0.34			
27751102	**Low discrimination**					**27751102**
Actinomyces viscosus		84.67	0.77			
Actinomyces naeslundii		13.42	0.87			
Lactobacillus acidophilus		1.17	0.45			
27751103	**Low discrimination**					**27751103**
Actinomyces viscosus		58.35	0.69			
Actinomyces naeslundii		41.62	0.90			
Lactobacillus acidophilus		0.02	0.10			
27751110	**Good identification**					**27751110**
Actinomyces naeslundii		94.65	0.65	MLZ 5		
Actinomyces israelii		3.78	0.32			
Actinomyces viscosus		1.16	0.08			
27751111	**Very good identification**					**27751111**
Actinomyces naeslundii		99.73	0.68	MLZ 5		
Actinomyces viscosus		0.27	0.00			
27751112	**Low discrimination**					**27751112**
Actinomyces viscosus		49.97	0.43	MLZ 1		
Actinomyces naeslundii		41.26	0.65			
Actinomyces israelii		2.47	0.35			
27751113	**Low discrimination**					**27751113**
Actinomyces naeslundii		78.80	0.68	MLZ 5		
Actinomyces viscosus		21.20	0.34			
27751120	**Good identification**					**27751120**
					GRA	
Actinomyces naeslundii		93.42	0.87		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		3.58	0.41		99	
Prevotella oralis		1.57	0.31		1	
27751121	**Very good identification**					**27751121**
Actinomyces naeslundii		99.95	0.90			
Clostridium perfringens		0.03	0.06			
Actinomyces viscosus		0.01	0.00			
27751122	**Low discrimination**					**27751122**
					GRA	
Actinomyces naeslundii		66.03	0.87		99	
Prevotella oralis		21.08	0.53		1	
Lactobacillus acidophilus		5.75	0.45		99	
27751123	**Very good identification**					**27751123**
					GRA	
Actinomyces naeslundii		98.33	0.90		99	
Actinomyces viscosus		1.39	0.34		99	
Prevotella oralis		0.21	0.19		1	
27751130	**Very good identification**					**27751130**
Actinomyces naeslundii		98.21	0.65	MLZ 5		
Bifidobacterium breve		0.75	0.15			
Clostridium perfringens		0.72	0.06			
27751131	**Very good identification**					**27751131**
Actinomyces naeslundii		100.0	0.68	MLZ 5		
27751132	**Low discrimination**					**27751132**
					GRA	
Actinomyces naeslundii		83.67	0.65	MLZ 5	99	
Bifidobacterium breve		6.49	0.32		99	
Prevotella oralis		5.13	0.19		1	
27751133	**Very good identification**					**27751133**
Actinomyces naeslundii		99.73	0.68	MLZ 5		
Actinomyces viscosus		0.27	0.00			
27751140	**Very good identification**					**27751140**
Actinomyces naeslundii		96.41	0.74			
Actinomyces israelii		2.99	0.39			
Actinomyces viscosus		0.35	0.08			
27751141	**Very good identification**					**27751141**
Actinomyces naeslundii		99.92	0.77			
Actinomyces viscosus		0.08	0.00			
27751142	**Low discrimination**					**27751142**
Actinomyces naeslundii		67.38	0.74			
Actinomyces viscosus		24.34	0.43			
Lactobacillus acidophilus		4.11	0.29			
27751143	**Good identification**					**27751143**
Actinomyces naeslundii		92.57	0.77			
Actinomyces viscosus		7.43	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27751150	**Low discrimination**					**27751150**
Actinomyces naeslundii		89.89	0.52	MLZ 5		
Actinomyces israelii		7.91	0.25			
Bifidobacterium breve		2.20	0.10			
27751151	**Good identification**					**27751151**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27751152	**Low discrimination**					**27751152**
Actinomyces naeslundii		67.45	0.52	MLZ 5		
Bifidobacterium breve		16.68	0.27			
Actinomyces israelii		8.90	0.28			
27751153	**Good identification**					**27751153**
Actinomyces naeslundii		98.48	0.55	MLZ 5		
Actinomyces viscosus		1.52	0.00			
27751160	**Very good identification**					**27751160**
Actinomyces naeslundii		95.43	0.74			
Clostridium perfringens		3.95	0.28			
Bifidobacterium breve		0.22	0.15			
27751161	**Very good identification**					**27751161**
Actinomyces naeslundii		100.0	0.77			
27751162	**Low discrimination**					**27751162**
Actinomyces naeslundii		88.28	0.74			
Lactobacillus acidophilus		5.38	0.29			
Bifidobacterium breve		2.04	0.32			
27751163	**Very good identification**					**27751163**
Actinomyces naeslundii		99.92	0.77			
Actinomyces viscosus		0.08	0.00			
27751170	**Good identification**					**27751170**
Actinomyces naeslundii		95.83	0.52	MLZ 5		
Bifidobacterium breve		4.17	0.15			
27751171	**Good identification**					**27751171**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27751172	**Low discrimination**					**27751172**
Actinomyces naeslundii		69.47	0.52	MLZ 5		
Bifidobacterium breve		30.53	0.32			
27751173	**Good identification**					**27751173**
Actinomyces naeslundii		100.0	0.55	MLZ 5		
27751200	**Low discrimination**					**27751200**
Actinomyces naeslundii		66.27	0.45	RHA 1		
Actinomyces odontolyticus		22.03	0.20			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		6.76	0.04				
27751201	**Good identification**						**27751201**
Actinomyces naeslundii		100.0	0.48	RHA 1			
27751202	**Low discrimination**						**27751202**
Actinomyces viscosus		86.29	0.38	RHA 1			
Actinomyces naeslundii		8.54	0.45				
Actinomyces odontolyticus		1.74	0.16				
27751220	**Low discrimination**						**27751220**
					GRA		
Actinomyces naeslundii		84.49	0.45	RHA 1	99		
Prevotella oralis		10.94	0.04		1		
Actinomyces odontolyticus		4.57	0.06		99		
27751221	**Good identification**						**27751221**
Actinomyces naeslundii		100.0	0.48	RHA 1			
27751223	**Good identification**						**27751223**
Actinomyces naeslundii		100.0	0.48	RHA 1			
27751241	**Good identification**						**27751241**
Actinomyces naeslundii		100.0	0.35	RHA 1			
27751243	**Good identification**						**27751243**
Actinomyces naeslundii		100.0	0.35	RHA 1			
27751261	**Good identification**						**27751261**
Actinomyces naeslundii		100.0	0.35	RHA 1			
27751263	**Good identification**						**27751263**
Actinomyces naeslundii		100.0	0.35	RHA 1			
27751300	**Low discrimination**						**27751300**
Actinomyces naeslundii		80.45	0.53	RHA 1			
Actinomyces israelii		8.43	0.27				
Actinomyces viscosus		5.13	0.08				
27751301	**Good identification**						**27751301**
Actinomyces naeslundii		98.60	0.56	RHA 1			
Actinomyces viscosus		1.40	0.00				
27751302	**Low discrimination**						**27751302**
					GRA		
Actinomyces viscosus		71.29	0.43	RHA 1	99		
Actinomyces naeslundii		11.30	0.53		99		
Prevotella oralis		8.06	0.25		1		
27751320	**Low discrimination**						**27751320**
					SPO	GRA	
Actinomyces naeslundii		49.79	0.53	RHA 1	1	99	
Prevotella oralis		35.50	0.25		1	1	
Clostridium ramosum		12.80	0.18		92	50	
27751321	**Good identification**						**27751321**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.56	RHA 1			
27751322	**Low discrimination**						**27751322**
					SPO	GRA	
Prevotella oralis		82.03	0.47	NIT 1	1	1	
Clostridium ramosum		6.23	0.28		92	50	
Actinomyces naeslundii		6.06	0.53		1	99	
27751323	**Acceptable identification**						**27751323**
					GRA		
Actinomyces naeslundii		90.47	0.56	RHA 1	99		
Prevotella oralis		8.25	0.13		1		
Actinomyces viscosus		1.28	0.00		99		
27751340	**Low discrimination**						**27751340**
Actinomyces naeslundii		81.24	0.40	RHA 1			
Actinomyces israelii		18.77	0.20				
27751341	**Good identification**						**27751341**
Actinomyces naeslundii		100.0	0.43	RHA 1			
27751342	**Low discrimination**						**27751342**
Actinomyces naeslundii		46.26	0.40	RHA 1			
Lactobacillus acidophilus		21.01	0.10				
Actinomyces viscosus		16.70	0.08				
27751343	**Acceptable identification**						**27751343**
Actinomyces naeslundii		92.57	0.43	RHA 1			
Actinomyces viscosus		7.43	0.00				
27751360	**Good identification**						**27751360**
Actinomyces naeslundii		100.0	0.40	RHA 1			
27751361	**Good identification**						**27751361**
Actinomyces naeslundii		100.0	0.43	RHA 1			
27751362	**Low discrimination**						**27751362**
					GRA		
Actinomyces naeslundii		44.85	0.40	RHA 1	99		
Prevotella oralis		34.78	0.13		1		
Lactobacillus acidophilus		20.37	0.10		99		
27751363	**Good identification**						**27751363**
Actinomyces naeslundii		100.0	0.43	RHA 1			
27751400	**Low discrimination**						**27751400**
Actinomyces naeslundii		52.28	0.45	ARA 1			
Actinomyces israelii		42.38	0.34				
Actinomyces viscosus		5.34	0.04				
27751401	**Good identification**						**27751401**
Actinomyces naeslundii		100.0	0.48	ARA 1			
27751402	**Low discrimination**						**27751402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		82.01	0.38	ARA 1		
Actinomyces israelii		9.87	0.37			
Actinomyces naeslundii		8.12	0.45			
27751420	**Good identification**					**27751420**
Actinomyces naeslundii		95.08	0.45	ARA 1		
Actinomyces israelii		4.92	0.14			
27751421	**Good identification**					**27751421**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27751422	**Low discrimination**					**27751422**
Actinomyces naeslundii		80.80	0.45	ARA 1		
Actinomyces viscosus		8.25	0.04			
Actinomyces israelii		6.27	0.17			
27751423	**Good identification**					**27751423**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27751441	**Good identification**					**27751441**
Actinomyces naeslundii		100.0	0.35	ARA 1		
27751443	**Good identification**					**27751443**
Actinomyces naeslundii		100.0	0.35	ARA 1		
27751461	**Good identification**					**27751461**
Actinomyces naeslundii		100.0	0.35	ARA 1		
27751463	**Good identification**					**27751463**
Actinomyces naeslundii		100.0	0.35	ARA 1		
27751500	**Low discrimination**					**27751500**
					SPO	
Actinomyces israelii		49.71	0.45	XYL 92	1	
Actinomyces naeslundii		43.63	0.53		1	
Clostridium butyricum		3.88	0.05		99	
27751501	**Good identification**					**27751501**
Actinomyces naeslundii		98.60	0.56	ARA 1		
Actinomyces viscosus		1.40	0.00			
27751502	**Low discrimination**					**27751502**
Actinomyces viscosus		59.83	0.43	ARA 1		
Actinomyces israelii		16.21	0.48			
Bifidobacterium adolescentis		13.17	0.40			
27751512	**Low discrimination**					**27751512**
Bifidobacterium adolescentis		80.55	0.41	NIT 1		
Actinomyces israelii		12.12	0.34	XYL 87		
Actinomyces viscosus		3.03	0.08			
27751520	**Low discrimination**					**27751520**
Actinomyces naeslundii		88.66	0.53	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		6.45	0.24			
Clostridium perfringens		3.40	0.06			
27751521	**Good identification**					**27751521**
Actinomyces naeslundii		100.0	0.56	ARA 1		
27751522	**Low discrimination**					**27751522**
					GRA	
Actinomyces naeslundii		58.44	0.53	ARA 1	99	
Prevotella oralis		18.66	0.19		1	
Actinomyces israelii		6.38	0.27		99	
27751523	**Good identification**					**27751523**
Actinomyces naeslundii		98.60	0.56	ARA 1		
Actinomyces viscosus		1.40	0.00			
27751540	**Low discrimination**					**27751540**
Actinomyces israelii		71.52	0.38	XYL 92		
Actinomyces naeslundii		28.48	0.40			
27751541	**Good identification**					**27751541**
Actinomyces naeslundii		100.0	0.43	ARA 1		
27751542	**Low discrimination**					**27751542**
Actinomyces israelii		61.75	0.41	XYL 92		
Actinomyces naeslundii		16.39	0.40			
Bifidobacterium adolescentis		15.94	0.24			
27751543	**Acceptable identification**					**27751543**
Actinomyces naeslundii		92.57	0.43	ARA 1		
Actinomyces viscosus		7.43	0.00			
27751560	**Low discrimination**					**27751560**
Actinomyces naeslundii		86.19	0.40	ARA 1		
Actinomyces israelii		13.82	0.17			
27751561	**Good identification**					**27751561**
Actinomyces naeslundii		100.0	0.43	ARA 1		
27751562	**Low discrimination**					**27751562**
Actinomyces naeslundii		80.62	0.40	ARA 1		
Actinomyces israelii		19.38	0.20			
27751563	**Good identification**					**27751563**
Actinomyces naeslundii		100.0	0.43	ARA 1		
27752000	**Very good identification**					**27752000**
Actinomyces naeslundii		97.29	0.61	GAL 96		
Actinomyces viscosus		2.02	0.08			
Actinomyces israelii		0.69	0.15			
27752001	**Very good identification**					**27752001**
Actinomyces naeslundii		99.54	0.64	GAL 96		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.46	-.01			
27752002	**Low discrimination**					**27752002**
Actinomyces viscosus		67.07	0.42	MAN 1		
Actinomyces naeslundii		32.59	0.61			
Actinomyces israelii		0.35	0.18			
27752003	**Low discrimination**					**27752003**
Actinomyces naeslundii		68.62	0.64	GAL 96		
Actinomyces viscosus		31.38	0.34			
27752010	**Good identification**					**27752010**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27752011	**Good identification**					**27752011**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27752012	**Low discrimination**					**27752012**
Actinomyces naeslundii		71.68	0.39	GAL 96		
Actinomyces viscosus		28.32	0.08	MLZ 5		
27752013	**Acceptable identification**					**27752013**
Actinomyces naeslundii		91.93	0.42	GAL 96		
Actinomyces viscosus		8.07	-.01	MLZ 5		
27752020	**Very good identification**					**27752020**
Actinomyces naeslundii		100.0	0.61	GAL 96		
27752021	**Very good identification**					**27752021**
Actinomyces naeslundii		100.0	0.64	GAL 96		
27752022	**Very good identification**					**27752022**
Actinomyces naeslundii		97.96	0.61	GAL 96		
Actinomyces viscosus		2.04	0.08			
27752023	**Very good identification**					**27752023**
Actinomyces naeslundii		99.54	0.64	GAL 96		
Actinomyces viscosus		0.46	-.01			
27752030	**Good identification**					**27752030**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27752031	**Good identification**					**27752031**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27752032	**Good identification**					**27752032**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27752033	**Good identification**					**27752033**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27752040	**Good identification**					**27752040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	GAL 96		
27752041	**Good identification**					**27752041**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27752042	**Low discrimination**					**27752042**
Actinomyces naeslundii		89.46	0.48	GAL 96		
Actinomyces viscosus		10.54	0.08			
27752043	**Good identification**					**27752043**
Actinomyces naeslundii		97.45	0.51	GAL 96		
Actinomyces viscosus		2.55	-.01			
27752060	**Good identification**					**27752060**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27752061	**Good identification**					**27752061**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27752062	**Good identification**					**27752062**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27752063	**Good identification**					**27752063**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27752100	**Very good identification**					**27752100**
Actinomyces naeslundii		96.96	0.68	GAL 96		
Actinomyces viscosus		1.26	0.12			
Actinomyces israelii		0.97	0.25			
27752101	**Very good identification**					**27752101**
Actinomyces naeslundii		99.71	0.71	GAL 96		
Actinomyces viscosus		0.29	0.04			
27752102	**Low discrimination**					**27752102**
Actinomyces viscosus		55.58	0.46	MAN 1		
Actinomyces naeslundii		43.23	0.68			
Actinomyces israelii		0.65	0.28			
27752103	**Low discrimination**					**27752103**
Actinomyces naeslundii		77.78	0.71	GAL 96		
Actinomyces viscosus		22.22	0.38			
27752110	**Good identification**					**27752110**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27752111	**Good identification**					**27752111**
Actinomyces naeslundii		100.0	0.50	GAL 96		
27752112	**Low discrimination**					**27752112**
Actinomyces naeslundii		77.57	0.47	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		19.14	0.12	MLZ 5		
Actinomyces israelii		3.29	0.14			
27752113	**Acceptable identification**					**27752113**
Actinomyces naeslundii		94.80	0.50	GAL 96		
Actinomyces viscosus		5.20	0.04	MLZ 5		
27752120	**Very good identification**					**27752120**
Actinomyces naeslundii		100.0	0.68	GAL 96		
27752121	**Very good identification**					**27752121**
Actinomyces naeslundii		100.0	0.71	GAL 96		
27752122	**Very good identification**					**27752122**
					SPO	
Actinomyces naeslundii		98.18	0.68	GAL 96	1	
Actinomyces viscosus		1.28	0.12		1	
Clostridium ramosum		0.55	0.05		92	
27752123	**Very good identification**					**27752123**
Actinomyces naeslundii		99.71	0.71	GAL 96		
Actinomyces viscosus		0.29	0.04			
27752130	**Good identification**					**27752130**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27752131	**Good identification**					**27752131**
Actinomyces naeslundii		100.0	0.50	GAL 96		
27752132	**Good identification**					**27752132**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27752133	**Good identification**					**27752133**
Actinomyces naeslundii		100.0	0.50	GAL 96		
27752140	**Acceptable identification**					**27752140**
Actinomyces naeslundii		93.51	0.56	GAL 96		
Propionibacterium propionicum		4.44	0.23			
Actinomyces israelii		2.05	0.18			
27752141	**Very good identification**					**27752141**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27752142	**Low discrimination**					**27752142**
Actinomyces naeslundii		84.92	0.56	GAL 96		
Actinomyces viscosus		6.25	0.12			
Propionibacterium propionicum		6.04	0.26			
27752143	**Very good identification**					**27752143**
Actinomyces naeslundii		98.39	0.59	GAL 96		
Actinomyces viscosus		1.61	0.04			
27752151	**Good identification**					**27752151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752153	**Good identification**					**27752153**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752160	**Good identification**					**27752160**
Actinomyces naeslundii		100.0	0.56	GAL 96		
27752161	**Very good identification**					**27752161**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27752162	**Good identification**					**27752162**
Actinomyces naeslundii		100.0	0.56	GAL 96		
27752163	**Very good identification**					**27752163**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27752171	**Good identification**					**27752171**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752173	**Good identification**					**27752173**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752301	**Good identification**					**27752301**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752303	**Low discrimination**					**27752303**
Actinomyces naeslundii		77.78	0.37	GAL 96		
Actinomyces viscosus		22.22	0.04	RHA 1		
27752321	**Good identification**					**27752321**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752323	**Good identification**					**27752323**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752501	**Good identification**					**27752501**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752503	**Low discrimination**					**27752503**
Actinomyces naeslundii		77.78	0.37	GAL 96		
Actinomyces viscosus		22.22	0.04	ARA 1		
27752521	**Good identification**					**27752521**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27752523	**Good identification**					**27752523**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27753000	**Very good identification**					**27753000**
Actinomyces naeslundii		97.11	0.84			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		2.84	0.49			
Actinomyces viscosus		0.05	0.04			
27753001	**Very good identification**					**27753001**
Actinomyces naeslundii		99.98	0.87			
Actinomyces israelii		0.02	0.15			
27753002	**Good identification**					**27753002**
Actinomyces naeslundii		91.16	0.84			
Actinomyces viscosus		4.79	0.38			
Actinomyces israelii		4.00	0.52			
27753003	**Very good identification**					**27753003**
Actinomyces naeslundii		98.82	0.87			
Actinomyces viscosus		1.15	0.30			
Actinomyces israelii		0.03	0.18			
27753010	**Good identification**					**27753010**
Actinomyces naeslundii		92.34	0.62	MLZ 5		
Actinomyces israelii		7.66	0.35			
27753011	**Very good identification**					**27753011**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27753012	**Low discrimination**					**27753012**
Actinomyces naeslundii		88.14	0.62	MLZ 5		
Actinomyces israelii		10.97	0.38			
Actinomyces viscosus		0.89	0.04			
27753013	**Very good identification**					**27753013**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27753020	**Very good identification**					**27753020**
Actinomyces naeslundii		99.81	0.84			
Actinomyces israelii		0.19	0.28			
27753021	**Very good identification**					**27753021**
Actinomyces naeslundii		100.0	0.87			
27753022	**Very good identification**					**27753022**
Actinomyces naeslundii		99.52	0.84			
Actinomyces israelii		0.28	0.31			
Lactobacillus acidophilus		0.06	0.05			
27753023	**Very good identification**					**27753023**
Actinomyces naeslundii		100.0	0.87			
27753030	**Very good identification**					**27753030**
Actinomyces naeslundii		99.47	0.62	MLZ 5		
Actinomyces israelii		0.53	0.14			
27753031	**Very good identification**					**27753031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27753032	**Very good identification**					**27753032**
Actinomyces naeslundii		99.21	0.62	MLZ 5		
Actinomyces israelii		0.79	0.17			
27753033	**Very good identification**					**27753033**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27753040	**Good identification**					**27753040**
Actinomyces naeslundii		93.95	0.71			
Actinomyces israelii		6.05	0.42			
27753041	**Very good identification**					**27753041**
Actinomyces naeslundii		100.0	0.74			
27753042	**Good identification**					**27753042**
Actinomyces naeslundii		90.94	0.71			
Actinomyces israelii		8.78	0.45			
Actinomyces viscosus		0.27	0.04			
27753043	**Very good identification**					**27753043**
Actinomyces naeslundii		100.0	0.74			
27753050	**Low discrimination**					**27753050**
Actinomyces naeslundii		84.55	0.50	MLZ 5		
Actinomyces israelii		15.46	0.28			
27753051	**Good identification**					**27753051**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27753052	**Low discrimination**					**27753052**
Actinomyces naeslundii		78.48	0.50	MLZ 5		
Actinomyces israelii		21.52	0.31			
27753053	**Good identification**					**27753053**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27753060	**Very good identification**					**27753060**
Actinomyces naeslundii		99.59	0.71			
Actinomyces israelii		0.41	0.21			
27753061	**Very good identification**					**27753061**
Actinomyces naeslundii		100.0	0.74			
27753062	**Very good identification**					**27753062**
Actinomyces naeslundii		99.39	0.71			
Actinomyces israelii		0.61	0.24			
27753063	**Very good identification**					**27753063**
Actinomyces naeslundii		100.0	0.74			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27753070	**Good identification**					**27753070**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27753071	**Good identification**					**27753071**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27753072	**Good identification**					**27753072**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27753073	**Good identification**					**27753073**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27753100	**Very good identification**					**27753100**
Actinomyces naeslundii		95.98	0.92			
Actinomyces israelii		3.94	0.59			
Propionibacterium propionicum		0.03	0.23			
27753101	**Excellent identification**					**27753101**
Actinomyces naeslundii		99.97	0.95			
Actinomyces israelii		0.03	0.25			
Actinomyces viscosus		0.01	0.00			
27753102	**Good identification**					**27753102**
Actinomyces naeslundii		90.93	0.92			
Actinomyces israelii		5.60	0.62			
Actinomyces viscosus		2.99	0.43			
27753103	**Excellent identification**					**27753103**
Actinomyces naeslundii		99.24	0.95			
Actinomyces viscosus		0.72	0.34			
Actinomyces israelii		0.04	0.28			
27753110	**Low discrimination**					**27753110**
Actinomyces naeslundii		89.38	0.70	MLZ 5		
Actinomyces israelii		10.42	0.45			
Bifidobacterium breve		0.20	0.10			
27753111	**Very good identification**					**27753111**
Actinomyces naeslundii		100.0	0.73	MLZ 5		
27753112	**Low discrimination**					**27753112**
Actinomyces naeslundii		81.69	0.70	MLZ 5		
Actinomyces israelii		14.29	0.48			
Bifidobacterium breve		1.84	0.27			
27753113	**Very good identification**					**27753113**
Actinomyces naeslundii		99.74	0.73	MLZ 5		
Actinomyces viscosus		0.14	0.00			
Actinomyces israelii		0.12	0.14			
27753120	**Excellent identification**					**27753120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.12	0.92		1	
Clostridium ramosum		0.57	0.28		92	
Actinomyces israelii		0.26	0.39		1	
27753121	**Excellent identification**					**27753121**
Actinomyces naeslundii		100.0	0.95			
27753122	**Very good identification**					**27753122**
					SPO	
Actinomyces naeslundii		96.75	0.92		1	
Clostridium ramosum		2.21	0.39		92	
Actinomyces israelii		0.38	0.42		1	
27753123	**Excellent identification**					**27753123**
					SPO	
Actinomyces naeslundii		99.98	0.95		1	
Clostridium ramosum		0.02	0.05		92	
Actinomyces viscosus		0.01	0.00		1	
27753130	**Very good identification**					**27753130**
Actinomyces naeslundii		98.87	0.70	MLZ 5		
Actinomyces israelii		0.74	0.25			
Bifidobacterium breve		0.39	0.15			
27753131	**Very good identification**					**27753131**
Actinomyces naeslundii		100.0	0.73	MLZ 5		
27753132	**Good identification**					**27753132**
Actinomyces naeslundii		94.42	0.70	MLZ 5		
Bifidobacterium breve		3.77	0.32			
Actinomyces israelii		1.05	0.28			
27753133	**Very good identification**					**27753133**
Actinomyces naeslundii		100.0	0.73	MLZ 5		
27753140	**Good identification**					**27753140**
Actinomyces naeslundii		91.48	0.79			
Actinomyces israelii		8.28	0.52			
Propionibacterium propionicum		0.18	0.23			
27753141	**Very good identification**					**27753141**
Actinomyces naeslundii		99.94	0.82			
Actinomyces israelii		0.06	0.18			
27753142	**Low discrimination**					**27753142**
Actinomyces naeslundii		86.99	0.79			
Actinomyces israelii		11.81	0.55			
Bifidobacterium breve		0.58	0.27			
27753143	**Very good identification**					**27753143**
Actinomyces naeslundii		99.87	0.82			
Actinomyces israelii		0.09	0.21			
Actinomyces viscosus		0.04	0.00			
27753150	**Low discrimination**					**27753150**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		78.77	0.57	MLZ 5		
Actinomyces israelii		20.24	0.38			
Bifidobacterium breve		0.99	0.10			
27753151	**Very good identification**					**27753151**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27753152	**Low discrimination**					**27753152**
Actinomyces naeslundii		65.48	0.57	MLZ 5		
Actinomyces israelii		25.24	0.41			
Bifidobacterium breve		8.34	0.27			
27753153	**Very good identification**					**27753153**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27753160	**Very good identification**					**27753160**
Actinomyces naeslundii		99.31	0.79			
Actinomyces israelii		0.57	0.32			
Bifidobacterium breve		0.12	0.15			
27753161	**Very good identification**					**27753161**
Actinomyces naeslundii		100.0	0.82			
27753162	**Very good identification**					**27753162**
Actinomyces naeslundii		97.70	0.79			
Bifidobacterium breve		1.16	0.32			
Actinomyces israelii		0.85	0.35			
27753163	**Very good identification**					**27753163**
Actinomyces naeslundii		100.0	0.82			
27753170	**Very good identification**					**27753170**
Actinomyces naeslundii		96.27	0.57	MLZ 5		
Bifidobacterium breve		2.16	0.15			
Actinomyces israelii		1.58	0.18			
27753171	**Very good identification**					**27753171**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27753172	**Low discrimination**					**27753172**
Actinomyces naeslundii		79.93	0.57	MLZ 5		
Bifidobacterium breve		18.10	0.32			
Actinomyces israelii		1.97	0.21			
27753173	**Very good identification**					**27753173**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27753200	**Low discrimination**					**27753200**
Actinomyces naeslundii		82.12	0.50	RHA 1		
Actinomyces israelii		17.88	0.30			
27753201	**Good identification**					**27753201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	RHA 1		
27753202	**Low discrimination**					**27753202**
Actinomyces naeslundii		72.51	0.50	RHA 1		
Actinomyces israelii		23.68	0.33			
Actinomyces viscosus		3.81	0.04			
27753203	**Good identification**					**27753203**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27753220	**Good identification**					**27753220**
Actinomyces naeslundii		100.0	0.50	RHA 1		
27753221	**Good identification**					**27753221**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27753222	**Acceptable identification**					**27753222**
					SPO	
Actinomyces naeslundii		92.06	0.50	RHA 1	1	
Clostridium ramosum		7.95	0.07		92	
27753223	**Good identification**					**27753223**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27753240	**Low discrimination**					**27753240**
Actinomyces naeslundii		67.58	0.37	RHA 1		
Actinomyces israelii		32.43	0.23			
27753241	**Good identification**					**27753241**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27753242	**Low discrimination**					**27753242**
Actinomyces naeslundii		58.15	0.37	RHA 1		
Actinomyces israelii		41.85	0.26			
27753243	**Good identification**					**27753243**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27753260	**Good identification**					**27753260**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27753261	**Good identification**					**27753261**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27753262	**Good identification**					**27753262**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27753263	**Good identification**					**27753263**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27753300	**Low discrimination**					**27753300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		76.57	0.58	RHA 1			
Actinomyces israelii		23.43	0.40				
27753301	**Very good identification**						**27753301**
Actinomyces naeslundii		100.0	0.61	RHA 1			
27753302	**Low discrimination**						**27753302**
Actinomyces naeslundii		64.95	0.58	RHA 1			
Actinomyces israelii		29.81	0.43				
Actinomyces viscosus		2.13	0.08				
27753303	**Very good identification**						**27753303**
Actinomyces naeslundii		99.28	0.61	RHA 1			
Actinomyces viscosus		0.72	0.00				
27753310	**Low discrimination**						**27753310**
Actinomyces naeslundii		53.51	0.36	RHA 1			
Actinomyces israelii		46.49	0.26	MLZ 5			
27753311	**Good identification**						**27753311**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27753313	**Good identification**						**27753313**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27753320	**Low discrimination**						**27753320**
					SPO		
Actinomyces naeslundii		56.50	0.58	RHA 1	1		
Clostridium ramosum		42.40	0.31		92		
Actinomyces israelii		1.10	0.19		1		
27753321	**Very good identification**						**27753321**
Actinomyces naeslundii		100.0	0.61	RHA 1			
27753322	**Low discrimination**						**27753322**
					SPO	GRA	
Clostridium ramosum		72.88	0.41	NIT 1	92	50	
Actinomyces naeslundii		24.27	0.58	TRE 94	1	99	
Prevotella oralis		1.71	0.13		1	1	
27753323	**Very good identification**						**27753323**
					SPO		
Actinomyces naeslundii		98.02	0.61	RHA 1	1		
Clostridium ramosum		1.98	0.07		92		
27753330	**Good identification**						**27753330**
Actinomyces naeslundii		100.0	0.36	RHA 1			
27753331	**Good identification**						**27753331**
Actinomyces naeslundii		100.0	0.39	RHA 1			
27753332	**Low discrimination**						**27753332**
					SPO		
Actinomyces naeslundii		63.45	0.36	RHA 1	1		
Clostridium ramosum		36.56	0.07	MLZ 5	92		
27753333	**Good identification**						**27753333**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	RHA 1		
27753340	**Low discrimination**					**27753340**
Actinomyces naeslundii		59.73	0.45	RHA 1		
Actinomyces israelii		40.27	0.33			
27753341	**Good identification**					**27753341**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27753342	**Low discrimination**					**27753342**
Actinomyces israelii		50.29	0.36	XYL 92		
Actinomyces naeslundii		49.72	0.45	RHA 7		
27753343	**Good identification**					**27753343**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27753360	**Good identification**					**27753360**
Actinomyces naeslundii		100.0	0.45	RHA 1		
27753361	**Good identification**					**27753361**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27753362	**Low discrimination**					**27753362**
					SPO	
Actinomyces naeslundii		80.88	0.45	RHA 1	1	
Clostridium ramosum		13.90	0.07		92	
Actinomyces israelii		5.22	0.15		1	
27753363	**Good identification**					**27753363**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27753400	**Low discrimination**					**27753400**
Actinomyces israelii		70.30	0.47	XYL 92		
Actinomyces naeslundii		29.70	0.50			
27753401	**Good identification**					**27753401**
Actinomyces naeslundii		98.43	0.53	ARA 1		
Actinomyces israelii		1.57	0.13			
27753402	**Low discrimination**					**27753402**
Actinomyces israelii		77.13	0.50	XYL 92		
Actinomyces naeslundii		21.73	0.50			
Actinomyces viscosus		1.14	0.04			
27753403	**Good identification**					**27753403**
Actinomyces naeslundii		97.67	0.53	ARA 1		
Actinomyces israelii		2.33	0.16			
27753412	**Acceptable identification**					**27753412**
Actinomyces israelii		90.97	0.36	XYL 92		
Actinomyces naeslundii		9.03	0.28	MLZ 13		
27753420	**Low discrimination**					**27753420**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		86.88	0.50	ARA 1		
Actinomyces israelii		13.12	0.27			
27753421	**Good identification**					**27753421**
Actinomyces naeslundii		100.0	0.53	ARA 1		
27753422	**Low discrimination**					**27753422**
Actinomyces naeslundii		81.53	0.50	ARA 1		
Actinomyces israelii		18.47	0.30			
27753423	**Good identification**					**27753423**
Actinomyces naeslundii		100.0	0.53	ARA 1		
27753440	**Low discrimination**					**27753440**
Actinomyces israelii		83.91	0.40	XYL 92		
Actinomyces naeslundii		16.09	0.37			
27753441	**Good identification**					**27753441**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27753442	**Low discrimination**					**27753442**
Actinomyces israelii		88.67	0.43	XYL 92		
Actinomyces naeslundii		11.33	0.37			
27753443	**Good identification**					**27753443**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27753460	**Low discrimination**					**27753460**
Actinomyces naeslundii		75.02	0.37	ARA 1		
Actinomyces israelii		24.98	0.20			
27753461	**Good identification**					**27753461**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27753462	**Low discrimination**					**27753462**
Actinomyces naeslundii		66.69	0.37	ARA 1		
Actinomyces israelii		33.31	0.23			
27753463	**Good identification**					**27753463**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27753500	**Low discrimination**					**27753500**
Actinomyces israelii		76.89	0.58	XYL 92		
Actinomyces naeslundii		23.12	0.58			
27753501	**Very good identification**					**27753501**
Actinomyces naeslundii		97.81	0.61	ARA 1		
Actinomyces israelii		2.19	0.24			
27753502	**Low discrimination**					**27753502**
Actinomyces israelii		75.51	0.61	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		15.14	0.58			
Bifidobacterium adolescentis		8.86	0.38			
27753503	**Very good identification**					**27753503**
Actinomyces naeslundii		96.07	0.61	ARA 1		
Actinomyces israelii		3.23	0.27			
Actinomyces viscosus		0.70	0.00			
27753510	**Acceptable identification**					**27753510**
Actinomyces israelii		90.43	0.44	XYL 92		
Actinomyces naeslundii		9.58	0.36	MLZ 13		
27753511	**Good identification**					**27753511**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27753512	**Low discrimination**					**27753512**
Actinomyces israelii		49.26	0.47	XYL 92		
Bifidobacterium adolescentis		47.26	0.40	MLZ 13		
Actinomyces naeslundii		3.48	0.36			
27753513	**Good identification**					**27753513**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27753520	**Low discrimination**					**27753520**
Actinomyces naeslundii		82.49	0.58	ARA 1		
Actinomyces israelii		17.51	0.37			
27753521	**Very good identification**					**27753521**
Actinomyces naeslundii		100.0	0.61	ARA 1		
27753522	**Low discrimination**					**27753522**
					SPO	
Actinomyces naeslundii		74.55	0.58	ARA 1	1	
Actinomyces israelii		23.74	0.40		1	
Clostridium ramosum		1.71	0.05		92	
27753523	**Very good identification**					**27753523**
Actinomyces naeslundii		100.0	0.61	ARA 1		
27753530	**Low discrimination**					**27753530**
Actinomyces naeslundii		62.39	0.36	ARA 1		
Actinomyces israelii		37.61	0.23	MLZ 5		
27753531	**Good identification**					**27753531**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27753532	**Low discrimination**					**27753532**
Actinomyces naeslundii		52.52	0.36	ARA 1		
Actinomyces israelii		47.48	0.26	MLZ 5		
27753533	**Good identification**					**27753533**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27753540	**Low discrimination**					**27753540**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		88.00	0.51	XYL 92		
Actinomyces naeslundii		12.01	0.45			
27753541	**Good identification**					**27753541**
Actinomyces naeslundii		95.30	0.48	ARA 1		
Actinomyces israelii		4.70	0.16			
27753542	**Low discrimination**					**27753542**
Actinomyces israelii		88.63	0.54	XYL 92		
Actinomyces naeslundii		8.06	0.45			
Bifidobacterium adolescentis		3.30	0.23			
27753543	**Acceptable identification**					**27753543**
Actinomyces naeslundii		93.11	0.48	ARA 1		
Actinomyces israelii		6.89	0.20			
27753550	**Good identification**					**27753550**
Actinomyces israelii		95.42	0.37	XYL 92		
Actinomyces naeslundii		4.59	0.23	MLZ 13		
27753552	**Low discrimination**					**27753552**
Actinomyces israelii		74.80	0.40	XYL 92		
Bifidobacterium adolescentis		22.81	0.24	MLZ 13		
Actinomyces naeslundii		2.40	0.23			
27753560	**Low discrimination**					**27753560**
Actinomyces naeslundii		68.13	0.45	ARA 1		
Actinomyces israelii		31.87	0.30			
27753561	**Good identification**					**27753561**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27753562	**Low discrimination**					**27753562**
Actinomyces naeslundii		58.76	0.45	ARA 1		
Actinomyces israelii		41.24	0.33			
27753563	**Good identification**					**27753563**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27753700	**Good identification**					**27753700**
Actinomyces israelii		96.12	0.38	XYL 92		
Actinomyces naeslundii		3.88	0.24	RHA 7		
27753702	**Good identification**					**27753702**
Actinomyces israelii		97.38	0.41	XYL 92		
Actinomyces naeslundii		2.62	0.24	RHA 7		
27754002	**Good identification**					**27754002**
Actinomyces viscosus		98.51	0.42	XYL 1		
Actinomyces israelii		1.03	0.23			
Actinomyces odontolyticus		0.46	0.09			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27754102	**Good identification**					**27754102**
Actinomyces viscosus		97.31	0.46	XYL 1		
Actinomyces israelii		2.29	0.33			
Actinomyces naeslundii		0.39	0.29			
27754103	**Good identification**					**27754103**
Actinomyces viscosus		98.21	0.38	XYL 1		
Actinomyces naeslundii		1.79	0.32			
27755000	**Acceptable identification**					**27755000**
Actinomyces israelii		90.09	0.54			
Actinomyces naeslundii		7.91	0.45			
Actinomyces viscosus		0.81	0.04			
27755001	**Acceptable identification**					**27755001**
Actinomyces naeslundii		92.87	0.48	XYL 1		
Actinomyces israelii		7.13	0.20			
27755002	**Low discrimination**					**27755002**
Actinomyces israelii		60.51	0.57			
Actinomyces viscosus		35.76	0.38			
Actinomyces naeslundii		3.54	0.45			
27755010	**Good identification**					**27755010**
Actinomyces israelii		97.00	0.40	MLZ 13		
Actinomyces naeslundii		3.00	0.23			
27755012	**Acceptable identification**					**27755012**
Actinomyces israelii		92.62	0.43	MLZ 13		
Actinomyces viscosus		3.70	0.04			
Actinomyces naeslundii		1.91	0.23			
27755020	**Low discrimination**					**27755020**
Actinomyces naeslundii		57.89	0.45	XYL 1		
Actinomyces israelii		42.11	0.34			
27755021	**Good identification**					**27755021**
Actinomyces naeslundii		100.0	0.48	XYL 1		
27755022	**Low discrimination**					**27755022**
					GRA	
Actinomyces israelii		37.59	0.37	NAG 6	99	
Actinomyces naeslundii		34.45	0.45		99	
Prevotella oralis		24.45	0.17		1	
27755023	**Good identification**					**27755023**
Actinomyces naeslundii		100.0	0.48	XYL 1		
27755040	**Good identification**					**27755040**
Actinomyces israelii		96.17	0.47			
Actinomyces naeslundii		3.83	0.32			
27755041	**Good identification**					**27755041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	XYL 1		
27755042	**Good identification**					**27755042**
Actinomyces israelii		95.98	0.50			
Actinomyces naeslundii		2.55	0.32			
Actinomyces viscosus		1.47	0.04			
27755043	**Low discrimination**					**27755043**
Actinomyces naeslundii		79.77	0.35	XYL 1		
Actinomyces israelii		20.23	0.16			
27755052	**Good identification**					**27755052**
Actinomyces israelii		100.0	0.36	MLZ 13		
27755061	**Good identification**					**27755061**
Actinomyces naeslundii		100.0	0.35	XYL 1		
27755063	**Good identification**					**27755063**
Actinomyces naeslundii		100.0	0.35	XYL 1		
27755100	**Good identification**					**27755100**
					SPO	
Actinomyces israelii		91.76	0.64		1	
Actinomyces naeslundii		5.73	0.53		1	
Clostridium butyricum		1.83	0.14		99	
27755101	**Low discrimination**					**27755101**
Actinomyces naeslundii		89.13	0.56	XYL 1		
Actinomyces israelii		9.61	0.30			
Actinomyces viscosus		1.26	0.00			
27755102	**Low discrimination**					**27755102**
Actinomyces israelii		72.49	0.67			
Actinomyces viscosus		19.04	0.43			
Bifidobacterium adolescentis		4.19	0.40			
27755110	**Good identification**					**27755110**
Actinomyces israelii		96.51	0.50	MLZ 13		
Actinomyces naeslundii		2.12	0.31			
Bifidobacterium adolescentis		0.69	0.07			
27755112	**Low discrimination**					**27755112**
Actinomyces israelii		66.14	0.53	MLZ 13		
Bifidobacterium adolescentis		31.27	0.41			
Actinomyces viscosus		1.17	0.08			
27755120	**Low discrimination**					**27755120**
					GRA	
Actinomyces israelii		44.21	0.44	NAG 6	99	
Actinomyces naeslundii		43.25	0.53		99	
Prevotella oralis		8.89	0.16		1	
27755121	**Good identification**					**27755121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.56	XYL 1		
27755122	**Low discrimination**					**27755122**
					GRA	
Prevotella oralis		53.02	0.37	NIT 1	1	
Actinomyces israelii		20.81	0.47	XYL 11	99	
Actinomyces naeslundii		13.57	0.53		99	
27755123	**Good identification**					**27755123**
					GRA	
Actinomyces naeslundii		96.11	0.56	XYL 1	99	
Prevotella oralis		2.53	0.03		1	
Actinomyces viscosus		1.36	0.00		99	
27755140	**Very good identification**					**27755140**
					SPO	
Actinomyces israelii		96.82	0.57		1	
Actinomyces naeslundii		2.74	0.40		1	
Clostridium butyricum		0.43	-.04		99	
27755141	**Low discrimination**					**27755141**
Actinomyces naeslundii		80.80	0.43	XYL 1		
Actinomyces israelii		19.20	0.23			
27755142	**Very good identification**					**27755142**
Actinomyces israelii		95.78	0.60			
Actinomyces naeslundii		1.81	0.40			
Bifidobacterium adolescentis		1.76	0.24			
27755143	**Low discrimination**					**27755143**
Actinomyces naeslundii		69.60	0.43	XYL 1		
Actinomyces israelii		24.81	0.26			
Actinomyces viscosus		5.59	0.00			
27755150	**Good identification**					**27755150**
Actinomyces israelii		100.0	0.43	MLZ 13		
27755152	**Low discrimination**					**27755152**
Actinomyces israelii		86.94	0.46	MLZ 13		
Bifidobacterium adolescentis		13.06	0.26			
27755160	**Low discrimination**					**27755160**
Actinomyces israelii		69.26	0.37	NAG 6		
Actinomyces naeslundii		30.74	0.40			
27755161	**Good identification**					**27755161**
Actinomyces naeslundii		100.0	0.43	XYL 1		
27755162	**Low discrimination**					**27755162**
					GRA	
Actinomyces israelii		73.41	0.40	NAG 6	99	
Actinomyces naeslundii		21.73	0.40		99	
Prevotella oralis		4.86	0.03		1	
27755163	**Good identification**					**27755163**
Actinomyces naeslundii		100.0	0.43	XYL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27755202	**Low discrimination**					**27755202**
Actinomyces israelii		86.41	0.38	RHA 7		
Actinomyces viscosus		6.85	0.04			
Actinomyces odontolyticus		6.74	0.11			
27755300	**Good identification**					**27755300**
Actinomyces israelii		98.52	0.45	RHA 7		
Actinomyces odontolyticus		1.48	0.06			
27755302	**Low discrimination**					**27755302**
					GRA	
Actinomyces israelii		89.68	0.48	RHA 7	99	
Prevotella oralis		4.37	0.09		1	
Actinomyces viscosus		3.16	0.08		99	
27755332	**Low discrimination**					**27755332**
					GRA	
Parabacteroides distasonis		80.47	0.39	NIT 1	1	
Clostridium clostridioforme		16.42	0.22		50	
Actinomyces israelii		1.57	0.14		99	
27755340	**Good identification**					**27755340**
Actinomyces israelii		100.0	0.38	RHA 7		
27755342	**Good identification**					**27755342**
Actinomyces israelii		100.0	0.41	RHA 7		
27755400	**Good identification**					**27755400**
					SPO	
Actinomyces israelii		96.47	0.53		1	
Clostridium butyricum		3.53	0.07		99	
27755402	**Good identification**					**27755402**
					SPO	
Actinomyces israelii		96.06	0.56		1	
Bifidobacterium adolescentis		1.84	0.20		1	
Clostridium butyricum		0.91	0.00		99	
27755410	**Good identification**					**27755410**
Actinomyces israelii		100.0	0.38	MLZ 13		
27755412	**Low discrimination**					**27755412**
Actinomyces israelii		76.17	0.42	MLZ 13		
Clostridium clostridioforme		11.93	0.18			
Bifidobacterium adolescentis		11.90	0.21			
27755422	**Acceptable identification**					**27755422**
Actinomyces israelii		92.55	0.35	NAG 6		
Clostridium clostridioforme		7.45	0.07			
27755440	**Good identification**					**27755440**
Actinomyces israelii		100.0	0.46			
27755442	**Good identification**					**27755442**
Actinomyces israelii		100.0	0.49			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
27755500	**Low discrimination**						**27755500**
					GRA	SPO	
Actinomyces israelii		80.27	0.63		99	1	
Clostridium butyricum		17.63	0.31		99	99	
Mitsuokella multacida		1.52	0.21		1	1	
27755502	**Low discrimination**						**27755502**
					SPO		
Actinomyces israelii		64.99	0.66		1		
Bifidobacterium adolescentis		30.72	0.54		1		
Clostridium butyricum		3.70	0.24		99		
27755510	**Low discrimination**						**27755510**
					SPO		
Actinomyces israelii		84.87	0.49	MLZ 13	1		
Clostridium butyricum		6.57	0.09		99		
Bifidobacterium adolescentis		4.97	0.21		1		
27755512	**Low discrimination**						**27755512**
Bifidobacterium adolescentis		77.06	0.56	NIT 1			
Actinomyces israelii		19.93	0.52				
Clostridium clostridioforme		2.61	0.27				
27755520	**Low discrimination**						**27755520**
					GRA	SPO	
Actinomyces israelii		60.20	0.42	NAG 6	99	1	
Clostridium butyricum		18.02	0.12		99	99	
Bacteroides vulgatus		17.31	0.11		1	1	
27755522	**Low discrimination**						**27755522**
					GRA	SPO	
Actinomyces israelii		63.21	0.45	NAG 6	99	1	
Parabacteroides distasonis		16.89	0.32		1	1	
Clostridium butyricum		4.90	0.05		99	99	
27755540	**Acceptable identification**						**27755540**
					GRA	SPO	
Actinomyces israelii		91.08	0.56		99	1	
Clostridium butyricum		4.47	0.12		99	99	
Mitsuokella multacida		4.44	0.21		1	1	
27755542	**Low discrimination**						**27755542**
					SPO		
Actinomyces israelii		85.72	0.59		1		
Bifidobacterium adolescentis		12.88	0.39		1		
Clostridium butyricum		1.09	0.05		99		
27755550	**Acceptable identification**						**27755550**
					GRA		
Actinomyces israelii		90.19	0.42	MLZ 13	99		
Mitsuokella multacida		9.81	0.13		1		
27755552	**Low discrimination**						**27755552**
Bifidobacterium adolescentis		55.14	0.40	NIT 1			
Actinomyces israelii		44.87	0.45	SOR 11			
27755560	**Low discrimination**						**27755560**
					GRA	SPO	
Actinomyces israelii		79.50	0.35	NAG 6	99	1	
Mitsuokella multacida		15.19	0.11		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		5.32	-.06		99	99	
27755562	**Good identification**						**27755562**
Actinomyces israelii		100.0	0.38	NAG 6			
27755602	**Low discrimination**						**27755602**
Actinomyces israelii		66.64	0.36	RHA 7			
Clostridium clostridioforme		33.36	0.22				
27755700	**Low discrimination**						**27755700**
					GRA	SPO	
Actinomyces israelii		63.28	0.44	RHA 7	99	1	
Mitsuokella multacida		29.61	0.25		1	1	
Clostridium butyricum		5.71	0.05		99	99	
27755702	**Low discrimination**						**27755702**
Actinomyces israelii		64.91	0.47	RHA 7			
Clostridium clostridioforme		27.20	0.31				
Bifidobacterium adolescentis		4.12	0.20				
27755712	**Low discrimination**						**27755712**
Clostridium clostridioforme		88.89	0.42	NIT 1			
Actinomyces israelii		6.96	0.33	TRE 88			
Bifidobacterium adolescentis		3.61	0.21				
27755722	**Low discrimination**						**27755722**
					GRA		
Parabacteroides distasonis		52.22	0.40	NIT 1	1		
Clostridium clostridioforme		32.18	0.31		50		
Actinomyces israelii		4.90	0.26		99		
27755732	**Low discrimination**						**27755732**
Clostridium clostridioforme		82.54	0.42	NIT 1			
Parabacteroides distasonis		15.83	0.35	TRE 88			
Bacteroides ovatus		1.63	0.03				
27755742	**Acceptable identification**						**27755742**
					GRA		
Actinomyces israelii		91.82	0.40	RHA 7	99		
Mitsuokella multacida		8.18	0.09		1		
27756002	**Low discrimination**						**27756002**
Actinomyces israelii		79.82	0.36	GAL 99			
Actinomyces viscosus		13.58	0.08				
Actinomyces naeslundii		6.60	0.27				
27756100	**Acceptable identification**						**27756100**
Actinomyces israelii		91.89	0.43	GAL 99			
Actinomyces naeslundii		8.11	0.34				
27756101	**Good identification**						**27756101**
Actinomyces naeslundii		100.0	0.37	GAL 96			
27756102	**Low discrimination**						**27756102**
Actinomyces israelii		88.15	0.46	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		6.67	0.12			
Actinomyces naeslundii		5.19	0.34			
27756103	**Low discrimination**					**27756103**
Actinomyces naeslundii		77.78	0.37	GAL 96		
Actinomyces viscosus		22.22	0.04	XYL 1		
27756121	**Good identification**					**27756121**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27756123	**Good identification**					**27756123**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27756140	**Good identification**					**27756140**
Actinomyces israelii		100.0	0.36	GAL 99		
27756142	**Good identification**					**27756142**
Actinomyces israelii		100.0	0.39	GAL 99		
27756500	**Good identification**					**27756500**
Actinomyces israelii		100.0	0.42	GAL 99		
27756502	**Acceptable identification**					**27756502**
Actinomyces israelii		93.61	0.45	GAL 99		
Bifidobacterium adolescentis		6.39	0.18			
27756542	**Good identification**					**27756542**
Actinomyces israelii		100.0	0.38	GAL 99		
27757000	**Very good identification**					**27757000**
Actinomyces israelii		97.08	0.67			
Actinomyces naeslundii		2.92	0.50			
27757001	**Low discrimination**					**27757001**
Actinomyces naeslundii		81.70	0.53	XYL 1		
Actinomyces israelii		18.30	0.33			
27757002	**Very good identification**					**27757002**
Actinomyces israelii		97.93	0.70			
Actinomyces naeslundii		1.96	0.50			
Actinomyces viscosus		0.10	0.04			
27757003	**Low discrimination**					**27757003**
Actinomyces naeslundii		74.85	0.53	XYL 1		
Actinomyces israelii		25.15	0.36			
27757010	**Good identification**					**27757010**
Actinomyces israelii		98.95	0.53	MLZ 13		
Actinomyces naeslundii		1.05	0.28			
27757012	**Good identification**					**27757012**
Actinomyces israelii		99.30	0.56	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.70	0.28			
27757020	**Low discrimination**					**27757020**
Actinomyces israelii		67.98	0.47	NAG 6		
Actinomyces naeslundii		32.02	0.50			
27757021	**Good identification**					**27757021**
Actinomyces naeslundii		100.0	0.53	XYL 1		
27757022	**Low discrimination**					**27757022**
Actinomyces israelii		76.11	0.50	NAG 6		
Actinomyces naeslundii		23.90	0.50			
27757023	**Good identification**					**27757023**
Actinomyces naeslundii		97.90	0.53	XYL 1		
Actinomyces israelii		2.10	0.15			
27757032	**Acceptable identification**					**27757032**
Actinomyces israelii		90.04	0.35	MLZ 13		
Actinomyces naeslundii		9.96	0.28	NAG 6		
27757040	**Very good identification**					**27757040**
Actinomyces israelii		98.65	0.60			
Actinomyces naeslundii		1.35	0.37			
27757041	**Low discrimination**					**27757041**
Actinomyces naeslundii		66.95	0.40	XYL 1		
Actinomyces israelii		33.05	0.26			
27757042	**Very good identification**					**27757042**
Actinomyces israelii		99.10	0.63			
Actinomyces naeslundii		0.90	0.37			
27757043	**Low discrimination**					**27757043**
Actinomyces naeslundii		57.46	0.40	XYL 1		
Actinomyces israelii		42.54	0.29			
27757050	**Good identification**					**27757050**
Actinomyces israelii		100.0	0.46	MLZ 13		
27757052	**Good identification**					**27757052**
Actinomyces israelii		100.0	0.49	MLZ 13		
27757060	**Low discrimination**					**27757060**
Actinomyces israelii		82.39	0.40	NAG 6		
Actinomyces naeslundii		17.61	0.37			
27757061	**Good identification**					**27757061**
Actinomyces naeslundii		100.0	0.40	XYL 1		
27757062	**Low discrimination**					**27757062**
Actinomyces israelii		87.53	0.43	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		12.47	0.37			
27757063	**Good identification**					**27757063**
Actinomyces naeslundii		100.0	0.40	XYL 1		
27757100	**Very good identification**					**27757100**
					SPO	
Actinomyces israelii		97.87	0.77		1	
Actinomyces naeslundii		2.09	0.58		1	
Clostridium butyricum		0.04	-.02		99	
27757101	**Low discrimination**					**27757101**
Actinomyces naeslundii		76.06	0.61	XYL 1		
Actinomyces israelii		23.94	0.43			
27757102	**Very good identification**					**27757102**
Actinomyces israelii		97.75	0.80			
Actinomyces naeslundii		1.39	0.58			
Bifidobacterium adolescentis		0.82	0.38			
27757103	**Low discrimination**					**27757103**
Actinomyces naeslundii		67.59	0.61	XYL 1		
Actinomyces israelii		31.92	0.46			
Actinomyces viscosus		0.49	0.00			
27757110	**Very good identification**					**27757110**
Actinomyces israelii		99.25	0.63	MLZ 13		
Actinomyces naeslundii		0.75	0.36			
27757111	**Low discrimination**					**27757111**
Actinomyces naeslundii		52.81	0.39	XYL 1		
Actinomyces israelii		47.20	0.29	MLZ 5		
27757112	**Good identification**					**27757112**
Actinomyces israelii		93.10	0.66	MLZ 13		
Bifidobacterium adolescentis		6.36	0.40			
Actinomyces naeslundii		0.47	0.36			
27757120	**Low discrimination**					**27757120**
Actinomyces israelii		74.90	0.57	NAG 6		
Actinomyces naeslundii		25.10	0.58			
27757121	**Very good identification**					**27757121**
Actinomyces naeslundii		98.03	0.61	XYL 1		
Actinomyces israelii		1.97	0.23			
27757122	**Low discrimination**					**27757122**
					SPO	
Actinomyces israelii		80.89	0.60	NAG 6	1	
Actinomyces naeslundii		18.07	0.58		1	
Clostridium ramosum		0.41	0.05		92	
27757123	**Very good identification**					**27757123**
Actinomyces naeslundii		97.07	0.61	XYL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		2.93	0.26			
27757130	**Low discrimination**					**27757130**
					SPO	
Actinomyces israelii		87.92	0.43	MLZ 13	1	
Actinomyces naeslundii		10.38	0.36	NAG 6	1	
Clostridium tertium		1.71	-.01		99	
27757131	**Good identification**					**27757131**
Actinomyces naeslundii		100.0	0.39	XYL 1		
27757132	**Low discrimination**					**27757132**
					SPO	
Actinomyces israelii		89.49	0.46	MLZ 13	1	
Actinomyces naeslundii		7.04	0.36	NAG 6	1	
Clostridium tertium		3.47	0.07		99	
27757133	**Good identification**					**27757133**
Actinomyces naeslundii		100.0	0.39	XYL 1		
27757140	**Very good identification**					**27757140**
Actinomyces israelii		99.04	0.70			
Actinomyces naeslundii		0.96	0.45			
27757141	**Low discrimination**					**27757141**
Actinomyces naeslundii		59.04	0.48	XYL 1		
Actinomyces israelii		40.96	0.36			
27757142	**Very good identification**					**27757142**
Actinomyces israelii		99.10	0.73			
Actinomyces naeslundii		0.64	0.45			
Bifidobacterium adolescentis		0.26	0.23			
27757143	**Low discrimination**					**27757143**
Actinomyces israelii		51.00	0.39	URE 1		
Actinomyces naeslundii		49.00	0.48			
27757150	**Very good identification**					**27757150**
Actinomyces israelii		99.66	0.56	MLZ 13		
Actinomyces naeslundii		0.34	0.23			
27757152	**Very good identification**					**27757152**
Actinomyces israelii		97.66	0.59	MLZ 13		
Bifidobacterium adolescentis		2.12	0.24			
Actinomyces naeslundii		0.22	0.23			
27757160	**Low discrimination**					**27757160**
Actinomyces israelii		86.80	0.50	NAG 6		
Actinomyces naeslundii		13.20	0.45			
27757161	**Good identification**					**27757161**
Actinomyces naeslundii		95.76	0.48	XYL 1		
Actinomyces israelii		4.24	0.16			
27757162	**Acceptable identification**					**27757162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		90.80	0.53	NAG 6		
Actinomyces naeslundii		9.21	0.45			
27757163	**Acceptable identification**					**27757163**
Actinomyces naeslundii		93.77	0.48	XYL 1		
Actinomyces israelii		6.23	0.19			
27757170	**Acceptable identification**					**27757170**
Actinomyces israelii		94.92	0.36	MLZ 13		
Actinomyces naeslundii		5.08	0.23	NAG 6		
27757172	**Good identification**					**27757172**
Actinomyces israelii		96.55	0.39	MLZ 13		
Actinomyces naeslundii		3.45	0.23	NAG 6		
27757200	**Good identification**					**27757200**
Actinomyces israelii		100.0	0.48	RHA 7		
27757202	**Good identification**					**27757202**
Actinomyces israelii		100.0	0.51	RHA 7		
27757212	**Good identification**					**27757212**
Actinomyces israelii		100.0	0.37	RHA 7		
27757240	**Good identification**					**27757240**
Actinomyces israelii		100.0	0.41	RHA 7		
27757242	**Good identification**					**27757242**
Actinomyces israelii		100.0	0.44	RHA 7		
27757300	**Very good identification**					**27757300**
Actinomyces israelii		99.71	0.58	RHA 7		
Actinomyces naeslundii		0.29	0.24			
27757302	**Very good identification**					**27757302**
Actinomyces israelii		99.81	0.61	RHA 7		
Actinomyces naeslundii		0.19	0.24			
27757310	**Good identification**					**27757310**
Actinomyces israelii		100.0	0.44	RHA 7		
27757312	**Good identification**					**27757312**
Actinomyces israelii		100.0	0.47	RHA 7		
27757320	**Good identification**					**27757320**
Actinomyces israelii		95.70	0.38	RHA 7		
Actinomyces naeslundii		4.30	0.24	NAG 6		
27757322	**Low discrimination**					**27757322**
					SPO	
Actinomyces israelii		85.38	0.41	RHA 7	1	
Clostridium ramosum		7.69	0.07	NAG 6	92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		2.56	0.24		1	
27757340	**Good identification**					**27757340**
Actinomyces israelii		100.0	0.51	RHA 7		
27757342	**Good identification**					**27757342**
Actinomyces israelii		100.0	0.54	RHA 7		
27757350	**Good identification**					**27757350**
Actinomyces israelii		100.0	0.37	RHA 7		
27757352	**Good identification**					**27757352**
Actinomyces israelii		100.0	0.40	RHA 7		
27757400	**Very good identification**					**27757400**
Actinomyces israelii		100.0	0.66			
27757402	**Very good identification**					**27757402**
Actinomyces israelii		99.73	0.69			
Bifidobacterium adolescentis		0.28	0.18			
27757410	**Good identification**					**27757410**
Actinomyces israelii		100.0	0.51	MLZ 13		
27757412	**Good identification**					**27757412**
Actinomyces israelii		97.79	0.54	MLZ 13		
Bifidobacterium adolescentis		2.21	0.20			
27757420	**Good identification**					**27757420**
Actinomyces israelii		100.0	0.45	NAG 6		
27757422	**Good identification**					**27757422**
Actinomyces israelii		100.0	0.48	NAG 6		
27757440	**Very good identification**					**27757440**
Actinomyces israelii		100.0	0.59			
27757442	**Very good identification**					**27757442**
Actinomyces israelii		100.0	0.62			
27757450	**Good identification**					**27757450**
Actinomyces israelii		100.0	0.44	MLZ 13		
27757452	**Good identification**					**27757452**
Actinomyces israelii		100.0	0.47	MLZ 13		
27757460	**Good identification**					**27757460**
Actinomyces israelii		100.0	0.38	NAG 6		
27757462	**Good identification**					**27757462**
Actinomyces israelii		100.0	0.41	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27757500	**Very good identification**						**27757500**
					GRA	SPO	
Actinomyces israelii		98.68	0.76		99	1	
Mitsuokella multacida		0.77	0.27		1	1	
Clostridium butyricum		0.43	0.14		99	99	
27757501	**Low discrimination**						**27757501**
					GRA		
Actinomyces israelii		88.88	0.42	URE 1	99		
Mitsuokella multacida		7.64	0.11		1		
Actinomyces naeslundii		3.49	0.27		99		
27757502	**Good identification**						**27757502**
					SPO		
Actinomyces israelii		93.45	0.79		1		
Bifidobacterium adolescentis		6.38	0.53		1		
Clostridium butyricum		0.10	0.07		99		
27757503	**Acceptable identification**						**27757503**
Actinomyces israelii		91.38	0.45	URE 1			
Bifidobacterium adolescentis		6.24	0.18				
Actinomyces naeslundii		2.39	0.27				
27757510	**Very good identification**						**27757510**
					GRA		
Actinomyces israelii		97.33	0.62	MLZ 13	99		
Mitsuokella multacida		1.70	0.19		1		
Bifidobacterium adolescentis		0.82	0.20		99		
27757512	**Low discrimination**						**27757512**
Actinomyces israelii		64.05	0.65	MLZ 13			
Bifidobacterium adolescentis		35.76	0.54				
Bifidobacterium dentium		0.11	-.07				
27757520	**Good identification**						**27757520**
					GRA	SPO	
Actinomyces israelii		96.10	0.55	NAG 6	99	1	
Mitsuokella multacida		2.94	0.17		1	1	
Clostridium butyricum		0.56	-.04		99	99	
27757522	**Very good identification**						**27757522**
Actinomyces israelii		98.66	0.58	NAG 6			
Bifidobacterium adolescentis		1.07	0.18				
Actinomyces naeslundii		0.27	0.24				
27757530	**Acceptable identification**						**27757530**
					GRA		
Actinomyces israelii		93.61	0.41	MLZ 13	99		
Mitsuokella multacida		6.39	0.09	NAG 6	1		
27757532	**Low discrimination**						**27757532**
Actinomyces israelii		87.91	0.44	MLZ 13			
Bifidobacterium adolescentis		7.77	0.20	NAG 6			
Bifidobacterium dentium		4.33	-.03				
27757540	**Very good identification**						**27757540**
					GRA	SPO	
Actinomyces israelii		97.94	0.69		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		1.97	0.27		1	1	
Clostridium butyricum		0.09	-.04		99	99	
27757542	**Very good identification**						**27757542**
					GRA		
Actinomyces israelii		97.73	0.72		99		
Bifidobacterium adolescentis		2.12	0.37		99		
Mitsuokella multacida		0.15	0.11		1		
27757543	**Good identification**						**27757543**
Actinomyces israelii		100.0	0.38	URE 1			
27757550	**Good identification**						**27757550**
					COC	GRA	
Actinomyces israelii		95.10	0.55	MLZ 13	1	99	
Mitsuokella multacida		4.26	0.19		1	1	
Blautia producta		0.64	-.03		99	99	
27757552	**Low discrimination**						**27757552**
					GRA		
Actinomyces israelii		84.69	0.58	MLZ 13	99		
Bifidobacterium adolescentis		15.03	0.39		99		
Mitsuokella multacida		0.28	0.03		1		
27757560	**Acceptable identification**						**27757560**
					GRA		
Actinomyces israelii		92.71	0.48	NAG 6	99		
Mitsuokella multacida		7.29	0.17		1		
27757562	**Good identification**						**27757562**
Actinomyces israelii		100.0	0.51	NAG 6			
27757572	**Good identification**						**27757572**
Actinomyces israelii		100.0	0.37	MLZ 13			
27757600	**Good identification**						**27757600**
Actinomyces israelii		100.0	0.46	RHA 7			
27757602	**Good identification**						**27757602**
Actinomyces israelii		100.0	0.49	RHA 7			
27757612	**Good identification**						**27757612**
Actinomyces israelii		100.0	0.35	RHA 7			
27757640	**Good identification**						**27757640**
Actinomyces israelii		100.0	0.39	RHA 7			
27757642	**Good identification**						**27757642**
Actinomyces israelii		100.0	0.42	RHA 7			
27757700	**Low discrimination**						**27757700**
					GRA		
Actinomyces israelii		83.84	0.57	RHA 7	99		
Mitsuokella multacida		16.16	0.32		1		
27757702	**Very good identification**						**27757702**
					GRA		
Actinomyces israelii		97.71	0.60	RHA 7	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Mitsuokella multacida		1.40	0.15		1	
Bifidobacterium adolescentis		0.90	0.18		99	
27757710	**Low discrimination**					**27757710**
					GRA	
Actinomyces israelii		69.94	0.42	RHA 7	99	
Mitsuokella multacida		30.06	0.24	MLZ 13	1	
27757712	**Low discrimination**					**27757712**
					GRA	
Actinomyces israelii		88.53	0.45	RHA 7	99	
Bifidobacterium adolescentis		6.63	0.20	MLZ 13	99	
Mitsuokella multacida		2.82	0.07		1	
27757720	**Low discrimination**					**27757720**
					GRA	
Actinomyces israelii		55.07	0.36	RHA 7	99	
Mitsuokella multacida		41.56	0.21	NAG 6	1	
Bacteroides vulgatus		3.38	-.07		1	
27757722	**Acceptable identification**					**27757722**
					GRA	
Actinomyces israelii		90.25	0.39	RHA 7	99	
Mitsuokella multacida		5.05	0.05	NAG 6	1	
Bacteroides ovatus		4.71	-.03		1	
27757740	**Low discrimination**					**27757740**
					GRA	
Actinomyces israelii		66.87	0.50	RHA 7	99	
Mitsuokella multacida		33.13	0.32		1	
27757742	**Good identification**					**27757742**
					GRA	
Actinomyces israelii		96.46	0.53	RHA 7	99	
Mitsuokella multacida		3.54	0.15		1	
27757752	**Acceptable identification**					**27757752**
					GRA	
Actinomyces israelii		92.43	0.38	RHA 7	99	
Mitsuokella multacida		7.57	0.07	MLZ 13	1	
27760000	**Low discrimination**					**27760000**
Actinomyces naeslundii		86.59	0.39	FRU 95		
Actinomyces viscosus		13.41	0.01	GAL 96		
27760001	**Good identification**					**27760001**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27760020	**Good identification**					**27760020**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27760021	**Good identification**					**27760021**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27760022	**Low discrimination**					**27760022**
Actinomyces naeslundii		86.59	0.39	FRU 95		
Actinomyces viscosus		13.41	0.01	GAL 96		
27760023	**Good identification**					**27760023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	FRU 95		
27760100	**Acceptable identification**					**27760100**
Actinomyces naeslundii		91.18	0.47	FRU 95		
Actinomyces viscosus		8.82	0.05	GAL 96		
27760101	**Good identification**					**27760101**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27760102	**Acceptable identification**					**27760102**
Actinomyces viscosus		90.55	0.39	FRU 95		
Actinomyces naeslundii		9.45	0.47	TRE 7		
27760120	**Good identification**					**27760120**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27760121	**Good identification**					**27760121**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27760122	**Acceptable identification**					**27760122**
Actinomyces naeslundii		91.18	0.47	FRU 95		
Actinomyces viscosus		8.82	0.05	GAL 96		
27760123	**Good identification**					**27760123**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27760141	**Good identification**					**27760141**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27760143	**Good identification**					**27760143**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27760161	**Good identification**					**27760161**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27760163	**Good identification**					**27760163**
Actinomyces naeslundii		100.0	0.37	FRU 95		
27761000	**Very good identification**					**27761000**
Actinomyces naeslundii		100.0	0.62	FRU 95		
27761001	**Very good identification**					**27761001**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27761002	**Low discrimination**					**27761002**
Actinomyces naeslundii		71.21	0.62	FRU 95		
Actinomyces viscosus		27.89	0.31			
Lactobacillus acidophilus		0.51	0.01			
27761003	**Good identification**					**27761003**
Actinomyces naeslundii		91.99	0.65	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		8.01	0.23			
27761010	**Good identification**					**27761010**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27761011	**Good identification**					**27761011**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27761012	**Good identification**					**27761012**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27761013	**Good identification**					**27761013**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27761020	**Very good identification**					**27761020**
Actinomyces naeslundii		100.0	0.62	FRU 95		
27761021	**Very good identification**					**27761021**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27761022	**Very good identification**					**27761022**
Actinomyces naeslundii		99.29	0.62	FRU 95		
Lactobacillus acidophilus		0.71	0.01			
27761023	**Very good identification**					**27761023**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27761030	**Good identification**					**27761030**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27761031	**Good identification**					**27761031**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27761032	**Good identification**					**27761032**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27761033	**Good identification**					**27761033**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27761040	**Good identification**					**27761040**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27761041	**Good identification**					**27761041**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27761042	**Good identification**					**27761042**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27761043	**Good identification**					**27761043**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27761060	**Good identification**					**27761060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	FRU 95		
27761061	**Good identification**					**27761061**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27761062	**Good identification**					**27761062**
Actinomyces naeslundii		100.0	0.50	FRU 95		
27761063	**Good identification**					**27761063**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27761100	**Very good identification**					**27761100**
Actinomyces naeslundii		99.24	0.70	FRU 95		
Actinomyces israelii		0.52	0.22			
Actinomyces viscosus		0.25	0.01			
27761101	**Very good identification**					**27761101**
Actinomyces naeslundii		100.0	0.73	FRU 95		
27761102	**Low discrimination**					**27761102**
Actinomyces naeslundii		78.05	0.70	FRU 95		
Actinomyces viscosus		19.09	0.36			
Lactobacillus acidophilus		2.25	0.19			
27761103	**Good identification**					**27761103**
Actinomyces naeslundii		94.84	0.73	FRU 95		
Actinomyces viscosus		5.16	0.27			
27761110	**Good identification**					**27761110**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27761111	**Good identification**					**27761111**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27761112	**Low discrimination**					**27761112**
					GRA	
Actinomyces naeslundii		69.33	0.48	FRU 95	99	
Leptotrichia buccalis		27.42	0.07	MLZ 5	1	
Actinomyces viscosus		3.26	0.01		99	
27761113	**Good identification**					**27761113**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27761120	**Very good identification**					**27761120**
					SPO	
Actinomyces naeslundii		99.00	0.70	FRU 95	1	
Clostridium perfringens		0.70	0.11		84	
Clostridium ramosum		0.30	0.02		92	
27761121	**Very good identification**					**27761121**
Actinomyces naeslundii		100.0	0.73	FRU 95		
27761122	**Very good identification**					**27761122**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.35	0.70	FRU 95	1	
Lactobacillus acidophilus		2.75	0.19		1	
Clostridium ramosum		1.16	0.12		92	
27761123	**Very good identification**					**27761123**
Actinomyces naeslundii		100.0	0.73	FRU 95		
27761130	**Good identification**					**27761130**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27761131	**Good identification**					**27761131**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27761132	**Good identification**					**27761132**
					GRA	
Actinomyces naeslundii		95.79	0.48	FRU 95	99	
Leptotrichia buccalis		4.21	-.09	MLZ 5	1	
27761133	**Good identification**					**27761133**
Actinomyces naeslundii		100.0	0.51	FRU 95		
27761140	**Very good identification**					**27761140**
Actinomyces naeslundii		98.86	0.57	FRU 95		
Actinomyces israelii		1.14	0.15			
27761141	**Very good identification**					**27761141**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27761142	**Very good identification**					**27761142**
Actinomyces naeslundii		95.11	0.57	FRU 95		
Lactobacillus acidophilus		1.92	0.04			
Actinomyces israelii		1.65	0.18			
27761143	**Very good identification**					**27761143**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27761150	**Good identification**					**27761150**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27761151	**Good identification**					**27761151**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27761152	**Good identification**					**27761152**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27761153	**Good identification**					**27761153**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27761160	**Very good identification**					**27761160**
Actinomyces naeslundii		100.0	0.57	FRU 95		
27761161	**Very good identification**					**27761161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.60	FRU 95		
27761162	**Very good identification**					**27761162**
Actinomyces naeslundii		98.02	0.57	FRU 95		
Lactobacillus acidophilus		1.98	0.04			
27761163	**Very good identification**					**27761163**
Actinomyces naeslundii		100.0	0.60	FRU 95		
27761170	**Good identification**					**27761170**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27761171	**Good identification**					**27761171**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27761172	**Good identification**					**27761172**
Actinomyces naeslundii		100.0	0.35	FRU 95		
27761173	**Good identification**					**27761173**
Actinomyces naeslundii		100.0	0.38	FRU 95		
27761300	**Good identification**					**27761300**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27761301	**Good identification**					**27761301**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27761302	**Low discrimination**					**27761302**
Actinomyces naeslundii		68.53	0.36	FRU 95		
Actinomyces viscosus		16.76	0.01	RHA 1		
Lactobacillus acidophilus		14.71	0.00			
27761303	**Good identification**					**27761303**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27761320	**Low discrimination**					**27761320**
					SPO	
Actinomyces naeslundii		71.52	0.36	FRU 95	1	
Clostridium ramosum		28.48	0.04	RHA 1	92	
27761321	**Good identification**					**27761321**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27761323	**Good identification**					**27761323**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27761500	**Low discrimination**					**27761500**
Actinomyces naeslundii		70.23	0.36	FRU 95		
Actinomyces israelii		29.77	0.20	ARA 1		
27761501	**Good identification**					**27761501**
Actinomyces naeslundii		100.0	0.39	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27761502	**Low discrimination**					**27761502**
Actinomyces naeslundii		53.18	0.36	FRU 95		
Actinomyces israelii		33.81	0.23	ARA 1		
Actinomyces viscosus		13.01	0.01			
27761503	**Good identification**					**27761503**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27761520	**Good identification**					**27761520**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27761521	**Good identification**					**27761521**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27761522	**Good identification**					**27761522**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27761523	**Good identification**					**27761523**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27762000	**Good identification**					**27762000**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27762001	**Good identification**					**27762001**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27762002	**Acceptable identification**					**27762002**
Actinomyces naeslundii		92.61	0.44	FRU 95		
Actinomyces viscosus		7.39	0.01	GAL 96		
27762003	**Good identification**					**27762003**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27762020	**Good identification**					**27762020**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27762021	**Good identification**					**27762021**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27762022	**Good identification**					**27762022**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27762023	**Good identification**					**27762023**
Actinomyces naeslundii		100.0	0.47	FRU 95		
27762100	**Good identification**					**27762100**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27762101	**Good identification**					**27762101**
Actinomyces naeslundii		100.0	0.55	FRU 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27762102	**Good identification**					**27762102**
Actinomyces naeslundii		95.25	0.52	FRU 95		
Actinomyces viscosus		4.75	0.05	GAL 96		
27762103	**Good identification**					**27762103**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27762120	**Good identification**					**27762120**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27762121	**Good identification**					**27762121**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27762122	**Good identification**					**27762122**
Actinomyces naeslundii		100.0	0.52	FRU 95		
27762123	**Good identification**					**27762123**
Actinomyces naeslundii		100.0	0.55	FRU 95		
27762140	**Good identification**					**27762140**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27762141	**Good identification**					**27762141**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27762142	**Good identification**					**27762142**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27762143	**Good identification**					**27762143**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27762160	**Good identification**					**27762160**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27762161	**Good identification**					**27762161**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27762162	**Good identification**					**27762162**
Actinomyces naeslundii		100.0	0.39	FRU 95		
27762163	**Good identification**					**27762163**
Actinomyces naeslundii		100.0	0.42	FRU 95		
27763000	**Very good identification**					**27763000**
Actinomyces naeslundii		98.93	0.67	FRU 95		
Actinomyces israelii		1.08	0.25			
27763001	**Very good identification**					**27763001**
Actinomyces naeslundii		100.0	0.70	FRU 95		
27763002	**Very good identification**					**27763002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.40	0.67	FRU 95		
Actinomyces israelii		1.60	0.28			
27763003	**Very good identification**					**27763003**
Actinomyces naeslundii		100.0	0.70	FRU 95		
27763010	**Good identification**					**27763010**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27763011	**Good identification**					**27763011**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27763012	**Good identification**					**27763012**
Actinomyces naeslundii		95.58	0.45	FRU 95		
Actinomyces israelii		4.42	0.13	MLZ 5		
27763013	**Good identification**					**27763013**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27763020	**Very good identification**					**27763020**
Actinomyces naeslundii		100.0	0.67	FRU 95		
27763021	**Very good identification**					**27763021**
Actinomyces naeslundii		100.0	0.70	FRU 95		
27763022	**Very good identification**					**27763022**
Actinomyces naeslundii		100.0	0.67	FRU 95		
27763023	**Very good identification**					**27763023**
Actinomyces naeslundii		100.0	0.70	FRU 95		
27763030	**Good identification**					**27763030**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27763031	**Good identification**					**27763031**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27763032	**Good identification**					**27763032**
Actinomyces naeslundii		100.0	0.45	FRU 95		
27763033	**Good identification**					**27763033**
Actinomyces naeslundii		100.0	0.48	FRU 95		
27763040	**Good identification**					**27763040**
Actinomyces naeslundii		97.66	0.54	FRU 95		
Actinomyces israelii		2.34	0.18			
27763041	**Very good identification**					**27763041**
Actinomyces naeslundii		100.0	0.57	FRU 95		
27763042	**Good identification**					**27763042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.53	0.54	FRU 95		
Actinomyces israelii		3.47	0.21			
27763043	**Very good identification**					**27763043**
Actinomyces naeslundii		100.0	0.57	FRU 95		
27763051	**Good identification**					**27763051**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763053	**Good identification**					**27763053**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763060	**Good identification**					**27763060**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27763061	**Very good identification**					**27763061**
Actinomyces naeslundii		100.0	0.57	FRU 95		
27763062	**Good identification**					**27763062**
Actinomyces naeslundii		100.0	0.54	FRU 95		
27763063	**Very good identification**					**27763063**
Actinomyces naeslundii		100.0	0.57	FRU 95		
27763071	**Good identification**					**27763071**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763073	**Good identification**					**27763073**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763100	**Very good identification**					**27763100**
Actinomyces naeslundii		98.50	0.75	FRU 95		
Actinomyces israelii		1.50	0.35			
27763101	**Very good identification**					**27763101**
Actinomyces naeslundii		100.0	0.78	FRU 95		
27763102	**Very good identification**					**27763102**
Actinomyces naeslundii		97.64	0.75	FRU 95		
Actinomyces israelii		2.24	0.38			
Actinomyces viscosus		0.12	0.01			
27763103	**Very good identification**					**27763103**
Actinomyces naeslundii		100.0	0.78	FRU 95		
27763110	**Good identification**					**27763110**
Actinomyces naeslundii		95.85	0.53	FRU 95		
Actinomyces israelii		4.16	0.21	MLZ 5		
27763111	**Very good identification**					**27763111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.56	FRU 95		
27763112	**Acceptable identification**					**27763112**
Actinomyces naeslundii		93.89	0.53	FRU 95		
Actinomyces israelii		6.11	0.24	MLZ 5		
27763113	**Very good identification**					**27763113**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27763120	**Very good identification**					**27763120**
					SPO	
Actinomyces naeslundii		99.03	0.75	FRU 95	1	
Clostridium ramosum		0.88	0.15		92	
Actinomyces israelii		0.10	0.14		1	
27763121	**Very good identification**					**27763121**
Actinomyces naeslundii		100.0	0.78	FRU 95		
27763122	**Very good identification**					**27763122**
					SPO	
Actinomyces naeslundii		96.44	0.75	FRU 95	1	
Clostridium ramosum		3.42	0.25		92	
Actinomyces israelii		0.14	0.17		1	
27763123	**Very good identification**					**27763123**
Actinomyces naeslundii		100.0	0.78	FRU 95		
27763130	**Good identification**					**27763130**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27763131	**Very good identification**					**27763131**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27763132	**Good identification**					**27763132**
Actinomyces naeslundii		100.0	0.53	FRU 95		
27763133	**Very good identification**					**27763133**
Actinomyces naeslundii		100.0	0.56	FRU 95		
27763140	**Very good identification**					**27763140**
Actinomyces naeslundii		96.74	0.62	FRU 95		
Actinomyces israelii		3.26	0.28			
27763141	**Very good identification**					**27763141**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27763142	**Very good identification**					**27763142**
Actinomyces naeslundii		95.19	0.62	FRU 95		
Actinomyces israelii		4.81	0.31			
27763143	**Very good identification**					**27763143**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27763150	**Acceptable identification**					**27763150**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.28	0.40	FRU 95		
Actinomyces israelii		8.72	0.14	MLZ 5		
27763151	**Good identification**					**27763151**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27763152	**Low discrimination**					**27763152**
Actinomyces naeslundii		87.47	0.40	FRU 95		
Actinomyces israelii		12.54	0.17	MLZ 5		
27763153	**Good identification**					**27763153**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27763160	**Very good identification**					**27763160**
Actinomyces naeslundii		100.0	0.62	FRU 95		
27763161	**Very good identification**					**27763161**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27763162	**Very good identification**					**27763162**
Actinomyces naeslundii		100.0	0.62	FRU 95		
27763163	**Very good identification**					**27763163**
Actinomyces naeslundii		100.0	0.65	FRU 95		
27763170	**Good identification**					**27763170**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27763171	**Good identification**					**27763171**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27763172	**Good identification**					**27763172**
Actinomyces naeslundii		100.0	0.40	FRU 95		
27763173	**Good identification**					**27763173**
Actinomyces naeslundii		100.0	0.43	FRU 95		
27763201	**Good identification**					**27763201**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763203	**Good identification**					**27763203**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763221	**Good identification**					**27763221**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763223	**Good identification**					**27763223**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763300	**Low discrimination**					**27763300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		89.78	0.41	FRU 95		
Actinomyces israelii		10.22	0.16	RHA 1		
27763301	**Good identification**					**27763301**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27763302	**Low discrimination**					**27763302**
Actinomyces naeslundii		85.42	0.41	FRU 95		
Actinomyces israelii		14.58	0.19	RHA 1		
27763303	**Good identification**					**27763303**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27763321	**Good identification**					**27763321**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27763323	**Good identification**					**27763323**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27763401	**Good identification**					**27763401**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763403	**Good identification**					**27763403**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763421	**Good identification**					**27763421**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763423	**Good identification**					**27763423**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27763501	**Good identification**					**27763501**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27763502	**Low discrimination**					**27763502**
Actinomyces israelii		64.98	0.36	FRU 99		
Actinomyces naeslundii		35.02	0.41	XYL 92		
27763503	**Good identification**					**27763503**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27763520	**Good identification**					**27763520**
Actinomyces naeslundii		100.0	0.41	FRU 95		
27763521	**Good identification**					**27763521**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27763522	**Low discrimination**					**27763522**
Actinomyces naeslundii		89.41	0.41	FRU 95		
Actinomyces israelii		10.59	0.16	ARA 1		
27763523	**Good identification**					**27763523**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	FRU 95			
27765100	**Low discrimination**						**27765100**
					SPO		
Actinomyces israelii		79.93	0.40	FRU 99	1		
Actinomyces naeslundii		13.42	0.36		1		
Clostridium butyricum		6.65	0.01		99		
27765101	**Good identification**						**27765101**
Actinomyces naeslundii		100.0	0.39	FRU 95			
27765102	**Low discrimination**						**27765102**
Actinomyces israelii		86.14	0.43	FRU 99			
Actinomyces naeslundii		9.64	0.36				
Actinomyces viscosus		2.36	0.01				
27765103	**Good identification**						**27765103**
Actinomyces naeslundii		100.0	0.39	FRU 95			
27765120	**Low discrimination**						**27765120**
Actinomyces naeslundii		72.45	0.36	FRU 95			
Actinomyces israelii		27.55	0.20	XYL 1			
27765121	**Good identification**						**27765121**
Actinomyces naeslundii		100.0	0.39	FRU 95			
27765122	**Low discrimination**						**27765122**
					GRA		
Actinomyces naeslundii		55.04	0.36	FRU 95	99		
Actinomyces israelii		31.39	0.23	XYL 1	99		
Parabacteroides distasonis		13.56	0.13		1		
27765123	**Good identification**						**27765123**
Actinomyces naeslundii		100.0	0.39	FRU 95			
27765142	**Good identification**						**27765142**
Actinomyces israelii		95.17	0.36	FRU 99			
Actinomyces naeslundii		4.83	0.23				
27765500	**Low discrimination**						**27765500**
					GRA	SPO	
Actinomyces israelii		41.43	0.39	FRU 99	99	1	
Clostridium butyricum		37.90	0.17		99	99	
Mitsuokella multacida		20.68	0.21		1	1	
27765502	**Low discrimination**						**27765502**
					SPO		
Actinomyces israelii		73.64	0.42	FRU 99	1		
Clostridium butyricum		17.47	0.10		99		
Bifidobacterium adolescentis		6.17	0.17		1		
27767000	**Acceptable identification**						**27767000**
Actinomyces israelii		92.52	0.43	FRU 99			
Actinomyces naeslundii		7.48	0.33				
27767001	**Good identification**						**27767001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	FRU 95		
27767002	**Acceptable identification**					**27767002**
Actinomyces israelii		94.89	0.46	FRU 99		
Actinomyces naeslundii		5.11	0.33			
27767003	**Good identification**					**27767003**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27767021	**Good identification**					**27767021**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27767023	**Good identification**					**27767023**
Actinomyces naeslundii		100.0	0.36	FRU 95		
27767040	**Good identification**					**27767040**
Actinomyces israelii		100.0	0.36	FRU 99		
27767042	**Good identification**					**27767042**
Actinomyces israelii		100.0	0.39	FRU 99		
27767100	**Acceptable identification**					**27767100**
Actinomyces israelii		94.56	0.53	FRU 99		
Actinomyces naeslundii		5.44	0.41			
27767101	**Low discrimination**					**27767101**
Actinomyces naeslundii		89.52	0.44	FRU 95		
Actinomyces israelii		10.48	0.19	XYL 1		
27767102	**Good identification**					**27767102**
Actinomyces israelii		96.31	0.56	FRU 99		
Actinomyces naeslundii		3.69	0.41			
27767103	**Low discrimination**					**27767103**
Actinomyces naeslundii		85.06	0.44	FRU 95		
Actinomyces israelii		14.94	0.22	XYL 1		
27767110	**Good identification**					**27767110**
Actinomyces israelii		100.0	0.39	FRU 99		
27767112	**Good identification**					**27767112**
Actinomyces israelii		100.0	0.42	FRU 99		
27767121	**Good identification**					**27767121**
Actinomyces naeslundii		100.0	0.44	FRU 95		
27767122	**Low discrimination**					**27767122**
Actinomyces israelii		62.47	0.36	FRU 99		
Actinomyces naeslundii		37.53	0.41	NAG 6		
27767123	**Good identification**					**27767123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	FRU 95			
27767140	**Good identification**						**27767140**
Actinomyces israelii		97.46	0.46	FRU 99			
Actinomyces naeslundii		2.54	0.28				
27767142	**Good identification**						**27767142**
Actinomyces israelii		98.29	0.49	FRU 99			
Actinomyces naeslundii		1.71	0.28				
27767302	**Good identification**						**27767302**
Actinomyces israelii		100.0	0.37	FRU 99			
27767400	**Good identification**						**27767400**
Actinomyces israelii		100.0	0.41	FRU 99			
27767402	**Good identification**						**27767402**
Actinomyces israelii		100.0	0.44	FRU 99			
27767442	**Good identification**						**27767442**
Actinomyces israelii		100.0	0.37	FRU 99			
27767500	**Low discrimination**						**27767500**
					GRA	SPO	
Actinomyces israelii		81.74	0.52	FRU 99	99	1	
Mitsuokella multacida		16.80	0.27		1	1	
Clostridium butyricum		1.47	0.01		99	99	
27767502	**Good identification**						**27767502**
					GRA		
Actinomyces israelii		96.90	0.55	FRU 99	99		
Mitsuokella multacida		1.48	0.11		1		
Bifidobacterium adolescentis		1.17	0.15		99		
27767510	**Low discrimination**						**27767510**
					GRA		
Actinomyces israelii		68.57	0.37	FRU 99	99		
Mitsuokella multacida		31.44	0.19	MLZ 13	1		
27767512	**Low discrimination**						**27767512**
Actinomyces israelii		84.79	0.40	FRU 99			
Bifidobacterium adolescentis		8.39	0.17	MLZ 13			
Bifidobacterium dentium		3.94	-.07				
27767540	**Low discrimination**						**27767540**
					GRA		
Actinomyces israelii		65.42	0.44	FRU 99	99		
Mitsuokella multacida		34.58	0.27		1		
27767542	**Good identification**						**27767542**
					GRA		
Actinomyces israelii		96.23	0.48	FRU 99	99		
Mitsuokella multacida		3.77	0.11		1		
27767702	**Low discrimination**						**27767702**
					GRA		
Actinomyces israelii		72.69	0.35	FRU 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Mitsuokella multacida		27.31	0.15	RHA 7	1	
27770000	**Low discrimination**					**27770000**
					GRA	
Actinomyces naeslundii		84.94	0.61	GAL 96	99	
Actinomyces viscosus		13.16	0.23		99	
Fusobacterium mortiferum		1.90	0.21		1	
27770001	**Very good identification**					**27770001**
Actinomyces naeslundii		96.67	0.64	GAL 96		
Actinomyces viscosus		3.33	0.15			
27770002	**Good identification**					**27770002**
Actinomyces viscosus		93.79	0.57	TRE 7		
Actinomyces naeslundii		6.12	0.61			
Lactobacillus acidophilus		0.06	0.01			
27770003	**Low discrimination**					**27770003**
Actinomyces viscosus		77.32	0.49	TRE 7		
Actinomyces naeslundii		22.68	0.64			
27770010	**Good identification**					**27770010**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27770011	**Good identification**					**27770011**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27770013	**Low discrimination**					**27770013**
Actinomyces naeslundii		60.46	0.42	GAL 96		
Actinomyces viscosus		39.55	0.15	MLZ 5		
27770020	**Very good identification**					**27770020**
Actinomyces naeslundii		100.0	0.61	GAL 96		
27770021	**Very good identification**					**27770021**
Actinomyces naeslundii		100.0	0.64	GAL 96		
27770022	**Low discrimination**					**27770022**
Actinomyces naeslundii		85.92	0.61	GAL 96		
Actinomyces viscosus		13.31	0.23			
Lactobacillus acidophilus		0.77	0.01			
27770023	**Very good identification**					**27770023**
Actinomyces naeslundii		96.67	0.64	GAL 96		
Actinomyces viscosus		3.33	0.15			
27770030	**Good identification**					**27770030**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27770031	**Good identification**					**27770031**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27770032	**Good identification**					**27770032**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	GAL 96		
27770033	**Good identification**					**27770033**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27770040	**Good identification**					**27770040**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27770041	**Good identification**					**27770041**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27770042	**Low discrimination**					**27770042**
Actinomyces naeslundii		53.25	0.48	GAL 96		
Actinomyces viscosus		46.75	0.23			
27770043	**Low discrimination**					**27770043**
Actinomyces naeslundii		83.68	0.51	GAL 96		
Actinomyces viscosus		16.32	0.15			
27770060	**Good identification**					**27770060**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27770061	**Good identification**					**27770061**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27770062	**Good identification**					**27770062**
Actinomyces naeslundii		100.0	0.48	GAL 96		
27770063	**Good identification**					**27770063**
Actinomyces naeslundii		100.0	0.51	GAL 96		
27770100	**Low discrimination**					**27770100**
					GRA	
Actinomyces naeslundii		89.15	0.68	GAL 96	99	
Actinomyces viscosus		8.63	0.27		99	
Fusobacterium mortiferum		1.64	0.27		1	
27770101	**Very good identification**					**27770101**
Actinomyces naeslundii		97.90	0.71	GAL 96		
Actinomyces viscosus		2.11	0.19			
27770102	**Good identification**					**27770102**
Actinomyces viscosus		90.14	0.61	TRE 7		
Actinomyces naeslundii		9.41	0.68			
Lactobacillus acidophilus		0.34	0.19			
27770103	**Low discrimination**					**27770103**
Actinomyces viscosus		68.04	0.53	TRE 7		
Actinomyces naeslundii		31.96	0.71			
27770110	**Good identification**					**27770110**
Actinomyces naeslundii		100.0	0.47	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27770111	**Good identification**					**27770111**
Actinomyces naeslundii		100.0	0.50	GAL 96		
27770113	**Low discrimination**					**27770113**
Actinomyces naeslundii		70.99	0.50	GAL 96		
Actinomyces viscosus		29.01	0.19	MLZ 5		
27770120	**Very good identification**					**27770120**
					SPO	
Actinomyces naeslundii		98.74	0.68	GAL 96	1	
Clostridium perfringens		0.88	0.11		84	
Clostridium ramosum		0.38	0.02		92	
27770121	**Very good identification**					**27770121**
Actinomyces naeslundii		100.0	0.71	GAL 96		
27770122	**Low discrimination**					**27770122**
Actinomyces naeslundii		86.56	0.68	GAL 96		
Actinomyces viscosus		8.38	0.27			
Lactobacillus acidophilus		3.15	0.19			
27770123	**Very good identification**					**27770123**
Actinomyces naeslundii		97.90	0.71	GAL 96		
Actinomyces viscosus		2.11	0.19			
27770130	**Good identification**					**27770130**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27770131	**Good identification**					**27770131**
Actinomyces naeslundii		100.0	0.50	GAL 96		
27770132	**Acceptable identification**					**27770132**
					GRA	
Actinomyces naeslundii		94.74	0.47	GAL 96	99	
Leptotrichia buccalis		5.26	-.09	MLZ 5	1	
27770133	**Good identification**					**27770133**
Actinomyces naeslundii		100.0	0.50	GAL 96		
27770140	**Good identification**					**27770140**
Actinomyces naeslundii		98.57	0.56	GAL 96		
Actinomyces israelii		1.44	0.15			
27770141	**Very good identification**					**27770141**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27770142	**Low discrimination**					**27770142**
Actinomyces naeslundii		62.67	0.56	GAL 96		
Actinomyces viscosus		34.37	0.27			
Lactobacillus acidophilus		1.60	0.04			
27770143	**Low discrimination**					**27770143**
Actinomyces naeslundii		89.14	0.59	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		10.86	0.19			
27770151	**Good identification**					**27770151**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770153	**Good identification**					**27770153**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770160	**Good identification**					**27770160**
Actinomyces naeslundii		100.0	0.56	GAL 96		
27770161	**Very good identification**					**27770161**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27770162	**Good identification**					**27770162**
Actinomyces naeslundii		97.52	0.56	GAL 96		
Lactobacillus acidophilus		2.49	0.04			
27770163	**Very good identification**					**27770163**
Actinomyces naeslundii		100.0	0.59	GAL 96		
27770171	**Good identification**					**27770171**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770173	**Good identification**					**27770173**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770301	**Good identification**					**27770301**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770321	**Good identification**					**27770321**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770323	**Good identification**					**27770323**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770501	**Good identification**					**27770501**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770521	**Good identification**					**27770521**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27770523	**Good identification**					**27770523**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27771000	**Very good identification**					**27771000**
Actinomyces naeslundii		96.07	0.84			
Actinomyces israelii		1.86	0.46			
Lactobacillus jensenii		1.47	0.32			
27771001	**Very good identification**					**27771001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.91	0.87			
Actinomyces viscosus		0.09	0.11			
27771002	**Low discrimination**					**27771002**
Actinomyces naeslundii		68.11	0.84			
Actinomyces viscosus		26.68	0.53			
Lactobacillus acidophilus		2.53	0.35			
27771003	**Good identification**					**27771003**
Actinomyces naeslundii		91.96	0.87			
Actinomyces viscosus		8.00	0.45			
Actinomyces israelii		0.02	0.15			
27771010	**Low discrimination**					**27771010**
Actinomyces naeslundii		86.79	0.62	MLZ 5		
Lactobacillus jensenii		8.43	0.24			
Actinomyces israelii		4.78	0.32			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27771011	**Very good identification**					**27771011**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27771012	**Low discrimination**					**27771012**
Actinomyces naeslundii		77.59	0.62	MLZ 5		
Actinomyces israelii		6.41	0.35			
Actinomyces viscosus		5.83	0.19			
27771013	**Very good identification**					**27771013**
Actinomyces naeslundii		98.36	0.65	MLZ 5		
Actinomyces viscosus		1.64	0.11			
27771020	**Very good identification**					**27771020**
Actinomyces naeslundii		98.03	0.84			
Clostridium perfringens		1.39	0.30			
Lactobacillus jensenii		0.38	0.21			
27771021	**Very good identification**					**27771021**
Actinomyces naeslundii		100.0	0.87			
27771022	**Good identification**					**27771022**
Actinomyces naeslundii		94.70	0.84			
Lactobacillus acidophilus		3.52	0.35			
Clostridium perfringens		0.63	0.25			
27771023	**Very good identification**					**27771023**
Actinomyces naeslundii		99.89	0.87			
Actinomyces viscosus		0.09	0.11			
Lactobacillus acidophilus		0.03	0.01			
27771030	**Very good identification**					**27771030**
Actinomyces naeslundii		97.63	0.62	MLZ 5		
Lactobacillus jensenii		2.37	0.13			
27771031	**Very good identification**					**27771031**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27771032	**Very good identification**					**27771032**
Actinomyces naeslundii		96.60	0.62	MLZ 5		
Lactobacillus jensenii		1.56	0.10			
Lactobacillus acidophilus		0.69	0.01			
27771033	**Very good identification**					**27771033**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27771040	**Very good identification**					**27771040**
Actinomyces naeslundii		95.90	0.71			
Actinomyces israelii		4.10	0.39			
27771041	**Very good identification**					**27771041**
Actinomyces naeslundii		100.0	0.74			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27771042	**Low discrimination**					**27771042**
Actinomyces naeslundii		89.88	0.71			
Actinomyces israelii		5.76	0.42			
Lactobacillus acidophilus		2.34	0.20			
27771043	**Very good identification**					**27771043**
Actinomyces naeslundii		99.50	0.74			
Actinomyces viscosus		0.50	0.11			
27771050	**Low discrimination**					**27771050**
Actinomyces naeslundii		89.18	0.50	MLZ 5		
Actinomyces israelii		10.82	0.25			
27771051	**Good identification**					**27771051**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27771052	**Low discrimination**					**27771052**
Actinomyces naeslundii		84.60	0.50	MLZ 5		
Actinomyces israelii		15.40	0.28			
27771053	**Good identification**					**27771053**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27771060	**Very good identification**					**27771060**
Actinomyces naeslundii		98.22	0.71			
Clostridium perfringens		1.51	0.18			
Actinomyces israelii		0.27	0.18			
27771061	**Very good identification**					**27771061**
Actinomyces naeslundii		100.0	0.74			
27771062	**Very good identification**					**27771062**
Actinomyces naeslundii		96.40	0.71			
Lactobacillus acidophilus		2.51	0.20			
Clostridium perfringens		0.70	0.12			
27771063	**Very good identification**					**27771063**
Actinomyces naeslundii		100.0	0.74			
27771070	**Good identification**					**27771070**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27771071	**Good identification**					**27771071**
Actinomyces naeslundii		100.0	0.53	MLZ 5		
27771072	**Good identification**					**27771072**
Actinomyces naeslundii		100.0	0.50	MLZ 5		
27771073	**Good identification**					**27771073**
Actinomyces naeslundii		100.0	0.53	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27771100	**Very good identification**					**27771100**
Actinomyces naeslundii		96.54	0.92			
Actinomyces israelii		2.63	0.56			
Actinomyces viscosus		0.24	0.23			
27771101	**Excellent identification**					**27771101**
Actinomyces naeslundii		99.93	0.95			
Actinomyces viscosus		0.06	0.15			
Actinomyces israelii		0.02	0.22			
27771102	**Low discrimination**					**27771102**
Actinomyces naeslundii		69.41	0.92			
Actinomyces viscosus		16.98	0.57			
Lactobacillus acidophilus		10.42	0.54			
27771103	**Good identification**					**27771103**
Actinomyces naeslundii		94.73	0.95			
Actinomyces viscosus		5.15	0.49			
Lactobacillus acidophilus		0.10	0.19			
27771110	**Good identification**					**27771110**
					GRA	
Actinomyces naeslundii		91.06	0.70	MLZ 5	99	
Actinomyces israelii		7.05	0.42		99	
Leptotrichia buccalis		1.90	0.07		1	
27771111	**Very good identification**					**27771111**
Actinomyces naeslundii		100.0	0.73	MLZ 5		
27771112	**Low discrimination**					**27771112**
					GRA	
Leptotrichia buccalis		62.71	0.41	NIT 10	1	
Actinomyces naeslundii		30.43	0.70	ESL 99	99	
Actinomyces israelii		3.53	0.45		99	
27771113	**Very good identification**					**27771113**
					GRA	
Actinomyces naeslundii		97.63	0.73	MLZ 5	99	
Leptotrichia buccalis		1.36	0.07		1	
Actinomyces viscosus		1.02	0.15		99	
27771120	**Good identification**					**27771120**
					SPO	
Actinomyces naeslundii		94.60	0.92		1	
Clostridium perfringens		3.47	0.45		84	
Clostridium ramosum		1.50	0.36		92	
27771121	**Excellent identification**					**27771121**
					SPO	
Actinomyces naeslundii		99.97	0.95		1	
Clostridium perfringens		0.03	0.11		84	
Clostridium ramosum		0.01	0.02		92	
27771122	**Low discrimination**					**27771122**
					SPO	
Actinomyces naeslundii		80.03	0.92		1	
Lactobacillus acidophilus		12.01	0.54		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		5.06	0.46		92	
27771123	**Excellent identification**					**27771123**
Actinomyces naeslundii		99.78	0.95			
Lactobacillus acidophilus		0.10	0.19			
Actinomyces viscosus		0.06	0.15			
27771130	**Very good identification**					**27771130**
Actinomyces naeslundii		98.04	0.70	MLZ 5		
Clostridium perfringens		0.69	0.11			
Actinomyces israelii		0.48	0.22			
27771131	**Very good identification**					**27771131**
Actinomyces naeslundii		100.0	0.73	MLZ 5		
27771132	**Low discrimination**					**27771132**
					GRA	
Actinomyces naeslundii		75.43	0.70	MLZ 5	99	
Leptotrichia buccalis		17.27	0.25		1	
Lactobacillus acidophilus		2.17	0.19		99	
27771133	**Very good identification**					**27771133**
					GRA	
Actinomyces naeslundii		99.85	0.73	MLZ 5	99	
Leptotrichia buccalis		0.15	-.09		1	
27771140	**Good identification**					**27771140**
Actinomyces naeslundii		94.13	0.79			
Actinomyces israelii		5.66	0.49			
Clostridium perfringens		0.12	0.07			
27771141	**Very good identification**					**27771141**
Actinomyces naeslundii		99.96	0.82			
Actinomyces israelii		0.04	0.15			
27771142	**Low discrimination**					**27771142**
Actinomyces naeslundii		82.32	0.79			
Lactobacillus acidophilus		8.66	0.38			
Actinomyces israelii		7.42	0.52			
27771143	**Very good identification**					**27771143**
Actinomyces naeslundii		99.56	0.82			
Actinomyces viscosus		0.31	0.15			
Lactobacillus acidophilus		0.07	0.04			
27771150	**Low discrimination**					**27771150**
Actinomyces naeslundii		85.43	0.57	MLZ 5		
Actinomyces israelii		14.57	0.35			
27771151	**Very good identification**					**27771151**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27771152	**Low discrimination**					**27771152**
					GRA	
Actinomyces naeslundii		67.09	0.57	MLZ 5	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		17.16	0.38		99	
Leptotrichia buccalis		7.91	0.07		1	
27771153	**Very good identification**					**27771153**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27771160	**Very good identification**					**27771160**
Actinomyces naeslundii		95.59	0.79			
Clostridium perfringens		3.78	0.33			
Actinomyces israelii		0.37	0.29			
27771161	**Very good identification**					**27771161**
Actinomyces naeslundii		100.0	0.82			
27771162	**Low discrimination**					**27771162**
Actinomyces naeslundii		87.59	0.79			
Lactobacillus acidophilus		9.21	0.38			
Clostridium perfringens		1.63	0.27			
27771163	**Very good identification**					**27771163**
Actinomyces naeslundii		99.93	0.82			
Lactobacillus acidophilus		0.07	0.04			
27771170	**Very good identification**					**27771170**
Actinomyces naeslundii		97.48	0.57	MLZ 5		
Bifidobacterium breve		1.46	0.12			
Actinomyces israelii		1.06	0.15			
27771171	**Very good identification**					**27771171**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27771172	**Low discrimination**					**27771172**
Actinomyces naeslundii		83.29	0.57	MLZ 5		
Bifidobacterium breve		12.57	0.29			
Lactobacillus acidophilus		1.68	0.04			
27771173	**Very good identification**					**27771173**
Actinomyces naeslundii		100.0	0.60	MLZ 5		
27771200	**Low discrimination**					**27771200**
Actinomyces naeslundii		87.37	0.50	RHA 1		
Actinomyces israelii		12.63	0.27			
27771201	**Good identification**					**27771201**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27771202	**Low discrimination**					**27771202**
Actinomyces naeslundii		53.04	0.50	RHA 1		
Actinomyces viscosus		20.78	0.19			
Lactobacillus acidophilus		14.68	0.16			
27771203	**Acceptable identification**					**27771203**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.99	0.53	RHA 1		
Actinomyces viscosus		8.01	0.11			
27771220	**Acceptable identification**					**27771220**
					SPO	
Actinomyces naeslundii		94.37	0.50	RHA 1	1	
Clostridium ramosum		5.63	0.04		92	
27771221	**Good identification**					**27771221**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27771222	**Low discrimination**					**27771222**
					SPO	
Actinomyces naeslundii		62.64	0.50	RHA 1	1	
Lactobacillus acidophilus		17.34	0.16		1	
Clostridium ramosum		14.95	0.14		92	
27771223	**Good identification**					**27771223**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27771240	**Low discrimination**					**27771240**
Actinomyces naeslundii		75.84	0.37	RHA 1		
Actinomyces israelii		24.16	0.20			
27771241	**Good identification**					**27771241**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27771242	**Low discrimination**					**27771242**
Actinomyces naeslundii		59.82	0.37	RHA 1		
Actinomyces israelii		28.58	0.23			
Lactobacillus acidophilus		11.60	0.01			
27771243	**Good identification**					**27771243**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27771260	**Good identification**					**27771260**
Actinomyces naeslundii		100.0	0.37	RHA 1		
27771261	**Good identification**					**27771261**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27771262	**Low discrimination**					**27771262**
Actinomyces naeslundii		83.76	0.37	RHA 1		
Lactobacillus acidophilus		16.24	0.01			
27771263	**Good identification**					**27771263**
Actinomyces naeslundii		100.0	0.40	RHA 1		
27771300	**Low discrimination**					**27771300**
					SPO	
Actinomyces naeslundii		80.95	0.58	RHA 1	1	
Actinomyces israelii		16.44	0.37		1	
Clostridium ramosum		1.70	0.04		92	
27771301	**Very good identification**					**27771301**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.61	RHA 1		
27771303	**Good identification**					**27771303**
Actinomyces naeslundii		94.17	0.61	RHA 1		
Actinomyces viscosus		5.12	0.15			
Lactobacillus acidophilus		0.71	0.00			
27771310	**Low discrimination**					**27771310**
Actinomyces naeslundii		63.42	0.36	RHA 1		
Actinomyces israelii		36.58	0.23	MLZ 5		
27771311	**Good identification**					**27771311**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27771313	**Good identification**					**27771313**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27771320	**Low discrimination**					**27771320**
					SPO	
Clostridium ramosum		65.68	0.38	NIT 1	92	
Actinomyces naeslundii		31.65	0.58		1	
Clostridium perfringens		1.16	0.11		84	
27771321	**Very good identification**					**27771321**
					SPO	
Actinomyces naeslundii		98.62	0.61	RHA 1	1	
Clostridium ramosum		1.38	0.04		92	
27771322	**Low discrimination**					**27771322**
					SPO	
Clostridium ramosum		73.17	0.48	NIT 1	92	
Lactobacillus acidophilus		9.86	0.34		1	
Actinomyces naeslundii		8.82	0.58		1	
27771323	**Good identification**					**27771323**
					SPO	
Actinomyces naeslundii		94.04	0.61	RHA 1	1	
Clostridium ramosum		5.26	0.14		92	
Lactobacillus acidophilus		0.71	0.00		1	
27771330	**Low discrimination**					**27771330**
					SPO	
Actinomyces naeslundii		71.52	0.36	RHA 1	1	
Clostridium ramosum		28.48	0.04	MLZ 5	92	
27771331	**Good identification**					**27771331**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27771333	**Good identification**					**27771333**
Actinomyces naeslundii		100.0	0.39	RHA 1		
27771340	**Low discrimination**					**27771340**
Actinomyces naeslundii		69.08	0.45	RHA 1		
Actinomyces israelii		30.92	0.30			
27771341	**Good identification**					**27771341**
Actinomyces naeslundii		100.0	0.48	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27771342	**Low discrimination**					**27771342**
Actinomyces naeslundii		40.74	0.45	RHA 1		
Lactobacillus acidophilus		31.91	0.19			
Actinomyces israelii		27.35	0.33			
27771343	**Good identification**					**27771343**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27771360	**Low discrimination**					**27771360**
					SPO	
Actinomyces naeslundii		89.38	0.45	RHA 1	1	
Clostridium ramosum		10.62	0.04		92	
27771361	**Good identification**					**27771361**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27771362	**Low discrimination**					**27771362**
					SPO	
Actinomyces naeslundii		44.28	0.45	RHA 1	1	
Lactobacillus acidophilus		34.69	0.19		1	
Clostridium ramosum		21.04	0.14		92	
27771363	**Good identification**					**27771363**
Actinomyces naeslundii		100.0	0.48	RHA 1		
27771400	**Low discrimination**					**27771400**
					SPO	
Actinomyces israelii		59.15	0.44	XYL 92	1	
Actinomyces naeslundii		37.65	0.50		1	
Clostridium butyricum		3.21	0.02		99	
27771401	**Good identification**					**27771401**
Actinomyces naeslundii		100.0	0.53	ARA 1		
27771402	**Low discrimination**					**27771402**
Actinomyces israelii		61.28	0.47	XYL 92		
Actinomyces naeslundii		26.00	0.50			
Actinomyces viscosus		10.19	0.19			
27771403	**Acceptable identification**					**27771403**
Actinomyces naeslundii		90.67	0.53	ARA 1		
Actinomyces viscosus		7.89	0.11			
Actinomyces israelii		1.44	0.13			
27771420	**Acceptable identification**					**27771420**
Actinomyces naeslundii		90.89	0.50	ARA 1		
Actinomyces israelii		9.12	0.24			
27771421	**Good identification**					**27771421**
Actinomyces naeslundii		100.0	0.53	ARA 1		
27771422	**Low discrimination**					**27771422**
Actinomyces naeslundii		82.34	0.50	ARA 1		
Actinomyces israelii		12.39	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		3.06	0.01			
27771423	**Good identification**					**27771423**
Actinomyces naeslundii		100.0	0.53	ARA 1		
27771440	**Low discrimination**					**27771440**
Actinomyces israelii		77.59	0.37	XYL 92		
Actinomyces naeslundii		22.41	0.37			
27771441	**Good identification**					**27771441**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27771442	**Low discrimination**					**27771442**
Actinomyces israelii		83.85	0.40	XYL 92		
Actinomyces naeslundii		16.15	0.37			
27771443	**Good identification**					**27771443**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27771460	**Low discrimination**					**27771460**
Actinomyces naeslundii		81.90	0.37	ARA 1		
Actinomyces israelii		18.10	0.17			
27771461	**Good identification**					**27771461**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27771462	**Low discrimination**					**27771462**
Actinomyces naeslundii		75.10	0.37	ARA 1		
Actinomyces israelii		24.90	0.20			
27771463	**Good identification**					**27771463**
Actinomyces naeslundii		100.0	0.40	ARA 1		
27771500	**Low discrimination**					**27771500**
					SPO	
Actinomyces israelii		54.46	0.55	XYL 92	1	
Actinomyces naeslundii		24.66	0.58		1	
Clostridium butyricum		17.72	0.25		99	
27771501	**Very good identification**					**27771501**
					GRA	
Actinomyces naeslundii		97.62	0.61	ARA 1	99	
Actinomyces israelii		1.45	0.20		99	
Mitsuokella multacida		0.93	0.04		1	
27771502	**Low discrimination**					**27771502**
Actinomyces israelii		60.12	0.58	XYL 92		
Actinomyces naeslundii		18.15	0.58			
Bifidobacterium adolescentis		8.66	0.37			
27771503	**Good identification**					**27771503**
Actinomyces naeslundii		92.88	0.61	ARA 1		
Actinomyces viscosus		5.05	0.15			
Actinomyces israelii		2.07	0.23			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27771510	**Low discrimination**					**27771510**
					GRA	
Actinomyces israelii		71.25	0.40	XYL 92	99	
Actinomyces naeslundii		11.37	0.36	MLZ 13	99	
Mitsuokella multacida		9.22	0.13		1	
27771511	**Good identification**					**27771511**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27771512	**Low discrimination**					**27771512**
Bifidobacterium adolescentis		41.71	0.38	NIT 1		
Actinomyces israelii		35.42	0.44	XYL 87		
Clostridium clostridioforme		10.30	0.25			
27771513	**Good identification**					**27771513**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27771520	**Low discrimination**					**27771520**
					SPO	
Actinomyces naeslundii		77.65	0.58	ARA 1	1	
Actinomyces israelii		10.94	0.34		1	
Clostridium butyricum		4.85	0.07		99	
27771521	**Very good identification**					**27771521**
Actinomyces naeslundii		100.0	0.61	ARA 1		
27771522	**Low discrimination**					**27771522**
Actinomyces naeslundii		62.66	0.58	ARA 1		
Actinomyces israelii		13.25	0.37			
Lactobacillus acidophilus		9.41	0.19			
27771523	**Very good identification**					**27771523**
Actinomyces naeslundii		100.0	0.61	ARA 1		
27771530	**Low discrimination**					**27771530**
					GRA	
Actinomyces naeslundii		62.39	0.36	ARA 1	99	
Actinomyces israelii		24.96	0.20	MLZ 5	99	
Mitsuokella multacida		12.65	0.02		1	
27771531	**Good identification**					**27771531**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27771533	**Good identification**					**27771533**
Actinomyces naeslundii		100.0	0.39	ARA 1		
27771540	**Low discrimination**					**27771540**
					GRA	
Actinomyces israelii		70.06	0.48	XYL 92	99	
Actinomyces naeslundii		14.40	0.45		99	
Mitsuokella multacida		10.45	0.21		1	
27771541	**Acceptable identification**					**27771541**
					GRA	
Actinomyces naeslundii		92.04	0.48	ARA 1	99	
Mitsuokella multacida		4.95	0.04		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		3.02	0.13		99	
27771542	**Low discrimination**					**27771542**
Actinomyces israelii		81.50	0.51	XYL 92		
Actinomyces naeslundii		11.17	0.45			
Bifidobacterium adolescentis		3.73	0.21			
27771543	**Good identification**					**27771543**
Actinomyces naeslundii		95.31	0.48	ARA 1		
Actinomyces israelii		4.69	0.16			
27771552	**Low discrimination**					**27771552**
Actinomyces israelii		70.30	0.36	XYL 92		
Bifidobacterium adolescentis		26.31	0.23	MLZ 13		
Actinomyces naeslundii		3.39	0.23			
27771560	**Low discrimination**					**27771560**
					GRA	
Actinomyces naeslundii		67.03	0.45	ARA 1	99	
Actinomyces israelii		20.82	0.27		99	
Mitsuokella multacida		12.16	0.11		1	
27771561	**Good identification**					**27771561**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27771562	**Low discrimination**					**27771562**
Actinomyces naeslundii		63.65	0.45	ARA 1		
Actinomyces israelii		29.66	0.30			
Lactobacillus acidophilus		6.69	0.04			
27771563	**Good identification**					**27771563**
Actinomyces naeslundii		100.0	0.48	ARA 1		
27771702	**Low discrimination**					**27771702**
					GRA	
Actinomyces israelii		46.39	0.38	XYL 92	99	
Clostridium clostridioforme		43.10	0.28	RHA 7	50	
Mitsuokella multacida		4.92	0.09		1	
27771712	**Good identification**					**27771712**
					GRA	
Clostridium clostridioforme		95.82	0.40	NIT 1	50	
Actinomyces israelii		3.38	0.24	XYL 91	99	
Mitsuokella multacida		0.80	0.01		1	
27771722	**Low discrimination**					**27771722**
					SPO	
Parabacteroides distasonis		42.64	0.35	NIT 1	1	
Clostridium clostridioforme		37.66	0.28	XYL 90	66	
Clostridium ramosum		13.63	0.14		92	
27771732	**Low discrimination**					**27771732**
Clostridium clostridioforme		88.20	0.40	NIT 1		
Parabacteroides distasonis		11.81	0.30	XYL 91		
27772000	**Very good identification**					**27772000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.65	0.66	GAL 96		
Actinomyces israelii		1.35	0.25			
27772001	**Very good identification**					**27772001**
Actinomyces naeslundii		100.0	0.69	GAL 96		
27772002	**Good identification**					**27772002**
Actinomyces naeslundii		90.88	0.66	GAL 96		
Actinomyces viscosus		7.25	0.23			
Actinomyces israelii		1.87	0.28			
27772003	**Very good identification**					**27772003**
Actinomyces naeslundii		98.26	0.69	GAL 96		
Actinomyces viscosus		1.74	0.15			
27772010	**Good identification**					**27772010**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27772011	**Good identification**					**27772011**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27772012	**Acceptable identification**					**27772012**
Actinomyces naeslundii		94.48	0.44	GAL 96		
Actinomyces israelii		5.52	0.13	MLZ 5		
27772013	**Good identification**					**27772013**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27772020	**Very good identification**					**27772020**
Actinomyces naeslundii		100.0	0.66	GAL 96		
27772021	**Very good identification**					**27772021**
Actinomyces naeslundii		100.0	0.69	GAL 96		
27772022	**Very good identification**					**27772022**
Actinomyces naeslundii		100.0	0.66	GAL 96		
27772023	**Very good identification**					**27772023**
Actinomyces naeslundii		100.0	0.69	GAL 96		
27772030	**Good identification**					**27772030**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27772031	**Good identification**					**27772031**
Actinomyces naeslundii		100.0	0.47	GAL 96		
27772032	**Good identification**					**27772032**
Actinomyces naeslundii		100.0	0.44	GAL 96		
27772033	**Good identification**					**27772033**
Actinomyces naeslundii		100.0	0.47	GAL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27772040	**Good identification**					**27772040**
Actinomyces naeslundii		97.06	0.53	GAL 96		
Actinomyces israelii		2.94	0.18			
27772041	**Good identification**					**27772041**
Actinomyces naeslundii		100.0	0.56	GAL 96		
27772042	**Good identification**					**27772042**
Actinomyces naeslundii		95.66	0.53	GAL 96		
Actinomyces israelii		4.34	0.21			
27772043	**Good identification**					**27772043**
Actinomyces naeslundii		100.0	0.56	GAL 96		
27772060	**Good identification**					**27772060**
Actinomyces naeslundii		100.0	0.53	GAL 96		
27772061	**Good identification**					**27772061**
Actinomyces naeslundii		100.0	0.56	GAL 96		
27772062	**Good identification**					**27772062**
Actinomyces naeslundii		100.0	0.53	GAL 96		
27772063	**Good identification**					**27772063**
Actinomyces naeslundii		100.0	0.56	GAL 96		
27772100	**Very good identification**					**27772100**
Actinomyces naeslundii		97.77	0.73	GAL 96		
Actinomyces israelii		1.89	0.35			
Propionibacterium propionicum		0.35	0.21			
27772101	**Very good identification**					**27772101**
Actinomyces naeslundii		100.0	0.76	GAL 96		
27772102	**Good identification**					**27772102**
Actinomyces naeslundii		92.24	0.73	GAL 96		
Actinomyces viscosus		4.60	0.27			
Actinomyces israelii		2.67	0.38			
27772103	**Very good identification**					**27772103**
Actinomyces naeslundii		98.90	0.76	GAL 96		
Actinomyces viscosus		1.10	0.19			
27772110	**Acceptable identification**					**27772110**
Actinomyces naeslundii		94.81	0.52	GAL 96		
Actinomyces israelii		5.19	0.21	MLZ 5		
27772111	**Good identification**					**27772111**
Actinomyces naeslundii		100.0	0.55	GAL 96		
27772112	**Acceptable identification**					**27772112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.41	0.52	GAL 96		
Actinomyces israelii		7.59	0.24	MLZ 5		
27772113	**Good identification**					**27772113**
Actinomyces naeslundii		100.0	0.55	GAL 96		
27772120	**Very good identification**					**27772120**
					SPO	
Actinomyces naeslundii		98.77	0.73	GAL 96	1	
Clostridium ramosum		1.11	0.15		92	
Actinomyces israelii		0.12	0.14		1	
27772121	**Very good identification**					**27772121**
Actinomyces naeslundii		100.0	0.76	GAL 96		
27772122	**Very good identification**					**27772122**
					SPO	
Actinomyces naeslundii		95.55	0.73	GAL 96	1	
Clostridium ramosum		4.28	0.25		92	
Actinomyces israelii		0.18	0.17		1	
27772123	**Very good identification**					**27772123**
Actinomyces naeslundii		100.0	0.76	GAL 96		
27772130	**Good identification**					**27772130**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27772131	**Good identification**					**27772131**
Actinomyces naeslundii		100.0	0.55	GAL 96		
27772132	**Good identification**					**27772132**
Actinomyces naeslundii		100.0	0.52	GAL 96		
27772133	**Good identification**					**27772133**
Actinomyces naeslundii		100.0	0.55	GAL 96		
27772140	**Good identification**					**27772140**
Actinomyces naeslundii		94.12	0.61	GAL 96		
Actinomyces israelii		4.00	0.28			
Propionibacterium propionicum		1.88	0.21			
27772141	**Very good identification**					**27772141**
Actinomyces naeslundii		100.0	0.64	GAL 96		
27772142	**Good identification**					**27772142**
Actinomyces naeslundii		91.43	0.61	GAL 96		
Actinomyces israelii		5.83	0.31			
Propionibacterium propionicum		2.74	0.24			
27772143	**Very good identification**					**27772143**
Actinomyces naeslundii		100.0	0.64	GAL 96		
27772150	**Low discrimination**					**27772150**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		89.23	0.39	GAL 96		
Actinomyces israelii		10.77	0.14	MLZ 5		
27772151	**Good identification**					**27772151**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772152	**Low discrimination**					**27772152**
Actinomyces naeslundii		84.67	0.39	GAL 96		
Actinomyces israelii		15.33	0.17	MLZ 5		
27772153	**Good identification**					**27772153**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772160	**Very good identification**					**27772160**
Actinomyces naeslundii		100.0	0.61	GAL 96		
27772161	**Very good identification**					**27772161**
Actinomyces naeslundii		100.0	0.64	GAL 96		
27772162	**Very good identification**					**27772162**
Actinomyces naeslundii		100.0	0.61	GAL 96		
27772163	**Very good identification**					**27772163**
Actinomyces naeslundii		100.0	0.64	GAL 96		
27772170	**Good identification**					**27772170**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27772171	**Good identification**					**27772171**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772172	**Good identification**					**27772172**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27772173	**Good identification**					**27772173**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772300	**Low discrimination**					**27772300**
Actinomyces naeslundii		87.43	0.39	GAL 96		
Actinomyces israelii		12.57	0.16	RHA 1		
27772301	**Good identification**					**27772301**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772302	**Low discrimination**					**27772302**
Actinomyces naeslundii		82.26	0.39	GAL 96		
Actinomyces israelii		17.74	0.19	RHA 1		
27772303	**Good identification**					**27772303**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772321	**Good identification**					**27772321**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772323	**Good identification**					**27772323**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772501	**Good identification**					**27772501**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772502	**Low discrimination**					**27772502**
Actinomyces israelii		70.10	0.36	GAL 99		
Actinomyces naeslundii		29.91	0.39	XYL 92		
27772503	**Good identification**					**27772503**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772520	**Good identification**					**27772520**
Actinomyces naeslundii		100.0	0.39	GAL 96		
27772521	**Good identification**					**27772521**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27772522	**Low discrimination**					**27772522**
Actinomyces naeslundii		86.99	0.39	GAL 96		
Actinomyces israelii		13.01	0.16	ARA 1		
27772523	**Good identification**					**27772523**
Actinomyces naeslundii		100.0	0.42	GAL 96		
27773000	**Good identification**					**27773000**
Actinomyces naeslundii		94.64	0.89			
Actinomyces israelii		5.36	0.59			
27773001	**Excellent identification**					**27773001**
Actinomyces naeslundii		99.96	0.92			
Actinomyces israelii		0.04	0.25			
27773002	**Good identification**					**27773002**
Actinomyces naeslundii		91.91	0.89			
Actinomyces israelii		7.81	0.62			
Actinomyces viscosus		0.19	0.19			
27773003	**Excellent identification**					**27773003**
Actinomyces naeslundii		99.90	0.92			
Actinomyces israelii		0.06	0.28			
Actinomyces viscosus		0.05	0.11			
27773010	**Low discrimination**					**27773010**
Actinomyces naeslundii		86.15	0.67	MLZ 5		
Actinomyces israelii		13.85	0.45			
27773011	**Very good identification**					**27773011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.70	MLZ 5		
27773012	**Low discrimination**					**27773012**
Actinomyces naeslundii		80.57	0.67	MLZ 5		
Actinomyces israelii		19.43	0.48			
27773013	**Very good identification**					**27773013**
Actinomyces naeslundii		99.84	0.70	MLZ 5		
Actinomyces israelii		0.16	0.13			
27773020	**Very good identification**					**27773020**
					SPO	
Actinomyces naeslundii		99.51	0.89		1	
Actinomyces israelii		0.36	0.38		1	
Clostridium ramosum		0.13	0.15		92	
27773021	**Excellent identification**					**27773021**
Actinomyces naeslundii		100.0	0.92			
27773022	**Very good identification**					**27773022**
					SPO	
Actinomyces naeslundii		98.84	0.89		1	
Actinomyces israelii		0.54	0.41		1	
Clostridium ramosum		0.53	0.25		92	
27773023	**Excellent identification**					**27773023**
Actinomyces naeslundii		100.0	0.92			
27773030	**Very good identification**					**27773030**
Actinomyces naeslundii		98.98	0.67	MLZ 5		
Actinomyces israelii		1.02	0.24			
27773031	**Very good identification**					**27773031**
Actinomyces naeslundii		100.0	0.70	MLZ 5		
27773032	**Very good identification**					**27773032**
Actinomyces naeslundii		98.48	0.67	MLZ 5		
Actinomyces israelii		1.52	0.27			
27773033	**Very good identification**					**27773033**
Actinomyces naeslundii		100.0	0.70	MLZ 5		
27773040	**Low discrimination**					**27773040**
Actinomyces naeslundii		88.91	0.76			
Actinomyces israelii		11.09	0.52			
27773041	**Very good identification**					**27773041**
Actinomyces naeslundii		99.92	0.79			
Actinomyces israelii		0.08	0.18			
27773042	**Low discrimination**					**27773042**
Actinomyces naeslundii		84.18	0.76			
Actinomyces israelii		15.76	0.55			
Lactobacillus acidophilus		0.06	-.02			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27773043	**Very good identification**					**27773043**
Actinomyces naeslundii		99.87	0.79			
Actinomyces israelii		0.13	0.21			
27773050	**Low discrimination**					**27773050**
Actinomyces naeslundii		73.84	0.54	MLZ 5		
Actinomyces israelii		26.16	0.38			
27773051	**Very good identification**					**27773051**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
27773052	**Low discrimination**					**27773052**
Actinomyces naeslundii		65.30	0.54	MLZ 5		
Actinomyces israelii		34.70	0.41			
27773053	**Very good identification**					**27773053**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
27773060	**Very good identification**					**27773060**
Actinomyces naeslundii		99.21	0.76			
Actinomyces israelii		0.79	0.31			
27773061	**Very good identification**					**27773061**
Actinomyces naeslundii		100.0	0.79			
27773062	**Very good identification**					**27773062**
Actinomyces naeslundii		98.75	0.76			
Actinomyces israelii		1.18	0.34			
Lactobacillus acidophilus		0.07	-.02			
27773063	**Very good identification**					**27773063**
Actinomyces naeslundii		100.0	0.79			
27773070	**Good identification**					**27773070**
Actinomyces naeslundii		97.79	0.54	MLZ 5		
Actinomyces israelii		2.21	0.17			
27773071	**Very good identification**					**27773071**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
27773072	**Good identification**					**27773072**
Actinomyces naeslundii		96.72	0.54	MLZ 5		
Actinomyces israelii		3.28	0.20			
27773073	**Very good identification**					**27773073**
Actinomyces naeslundii		100.0	0.57	MLZ 5		
27773100	**Good identification**					**27773100**
					SPO	
Actinomyces naeslundii		92.57	0.97		1	
Actinomyces israelii		7.37	0.69		1	
Clostridium ramosum		0.04	0.15		92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27773101	**Excellent identification**					**27773101**
Actinomyces naeslundii		99.95	1.00			
Actinomyces israelii		0.05	0.35			
27773102	**Low discrimination**					**27773102**
Actinomyces naeslundii		88.68	0.97			
Actinomyces israelii		10.59	0.72			
Lactobacillus acidophilus		0.36	0.32			
27773103	**Excellent identification**					**27773103**
Actinomyces naeslundii		99.89	1.00			
Actinomyces israelii		0.08	0.38			
Actinomyces viscosus		0.03	0.15			
27773110	**Low discrimination**					**27773110**
					SPO	
Actinomyces naeslundii		81.45	0.75	MLZ 5	1	
Actinomyces israelii		18.40	0.55		1	
Clostridium tertium		0.15	0.05		99	
27773111	**Very good identification**					**27773111**
Actinomyces naeslundii		99.85	0.78	MLZ 5		
Actinomyces israelii		0.15	0.21			
27773112	**Low discrimination**					**27773112**
					GRA	
Actinomyces naeslundii		73.00	0.75	MLZ 5	99	
Actinomyces israelii		24.74	0.58		99	
Leptotrichia buccalis		0.78	0.07		1	
27773113	**Very good identification**					**27773113**
Actinomyces naeslundii		99.77	0.78	MLZ 5		
Actinomyces israelii		0.23	0.24			
27773120	**Very good identification**					**27773120**
					SPO	
Actinomyces naeslundii		95.09	0.97		1	
Clostridium ramosum		4.39	0.49		92	
Actinomyces israelii		0.48	0.49		1	
27773121	**Excellent identification**					**27773121**
					SPO	
Actinomyces naeslundii		99.97	1.00		1	
Clostridium ramosum		0.03	0.15		92	
Actinomyces israelii		0.00	0.14		1	
27773122	**Low discrimination**					**27773122**
					SPO	
Actinomyces naeslundii		83.50	0.97		1	
Clostridium ramosum		15.42	0.59		92	
Actinomyces israelii		0.64	0.52		1	
27773123	**Excellent identification**					**27773123**
					SPO	
Actinomyces naeslundii		99.87	1.00		1	
Clostridium ramosum		0.12	0.25		92	
Actinomyces israelii		0.01	0.17		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27773130	**Very good identification**					**27773130**
					SPO	
Actinomyces naeslundii		97.07	0.75	MLZ 5	1	
Actinomyces israelii		1.40	0.34		1	
Clostridium ramosum		0.86	0.15		92	
27773131	**Very good identification**					**27773131**
Actinomyces naeslundii		100.0	0.78	MLZ 5		
27773132	**Good identification**					**27773132**
					SPO	
Actinomyces naeslundii		91.85	0.75	MLZ 5	1	
Clostridium ramosum		3.26	0.25		92	
Actinomyces israelii		1.99	0.37		1	
27773133	**Very good identification**					**27773133**
Actinomyces naeslundii		100.0	0.78	MLZ 5		
27773140	**Low discrimination**					**27773140**
Actinomyces naeslundii		85.02	0.84			
Actinomyces israelii		14.91	0.62			
Propionibacterium propionicum		0.07	0.21			
27773141	**Very good identification**					**27773141**
Actinomyces naeslundii		99.88	0.87			
Actinomyces israelii		0.12	0.28			
27773142	**Low discrimination**					**27773142**
Actinomyces naeslundii		78.77	0.84			
Actinomyces israelii		20.72	0.65			
Lactobacillus acidophilus		0.23	0.16			
27773143	**Very good identification**					**27773143**
Actinomyces naeslundii		99.82	0.87			
Actinomyces israelii		0.18	0.31			
27773150	**Low discrimination**					**27773150**
Actinomyces naeslundii		66.76	0.62	MLZ 5		
Actinomyces israelii		33.24	0.48			
27773151	**Very good identification**					**27773151**
Actinomyces naeslundii		99.67	0.65	MLZ 5		
Actinomyces israelii		0.33	0.14			
27773152	**Low discrimination**					**27773152**
Actinomyces naeslundii		55.70	0.62	MLZ 5		
Actinomyces israelii		41.59	0.51			
Bifidobacterium breve		2.44	0.24			
27773153	**Very good identification**					**27773153**
Actinomyces naeslundii		99.50	0.65	MLZ 5		
Actinomyces israelii		0.50	0.17			
27773160	**Very good identification**					**27773160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		98.60	0.84		1	
Actinomyces israelii		1.10	0.42		1	
Clostridium ramosum		0.26	0.15		92	
27773161	**Very good identification**					**27773161**
Actinomyces naeslundii		100.0	0.87			
27773162	**Very good identification**					**27773162**
					SPO	
Actinomyces naeslundii		96.68	0.84		1	
Actinomyces israelii		1.62	0.45		1	
Clostridium ramosum		1.02	0.25		92	
27773163	**Very good identification**					**27773163**
Actinomyces naeslundii		100.0	0.87			
27773170	**Very good identification**					**27773170**
Actinomyces naeslundii		96.20	0.62	MLZ 5		
Actinomyces israelii		3.06	0.27			
Bifidobacterium breve		0.74	0.12			
27773171	**Very good identification**					**27773171**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27773172	**Low discrimination**					**27773172**
Actinomyces naeslundii		88.85	0.62	MLZ 5		
Bifidobacterium breve		6.91	0.29			
Actinomyces israelii		4.24	0.30			
27773173	**Very good identification**					**27773173**
Actinomyces naeslundii		100.0	0.65	MLZ 5		
27773200	**Low discrimination**					**27773200**
Actinomyces naeslundii		70.33	0.55	RHA 1		
Actinomyces israelii		29.67	0.39			
27773201	**Very good identification**					**27773201**
Actinomyces naeslundii		100.0	0.58	RHA 1		
27773202	**Low discrimination**					**27773202**
Actinomyces naeslundii		61.24	0.55	RHA 1		
Actinomyces israelii		38.76	0.42			
27773203	**Very good identification**					**27773203**
Actinomyces naeslundii		100.0	0.58	RHA 1		
27773211	**Good identification**					**27773211**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27773213	**Good identification**					**27773213**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27773220	**Low discrimination**					**27773220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		83.26	0.55	RHA 1	1	
Clostridium ramosum		14.50	0.17		92	
Actinomyces israelii		2.24	0.19		1	
27773221	**Very good identification**					**27773221**
Actinomyces naeslundii		100.0	0.58	RHA 1		
27773222	**Low discrimination**					**27773222**
					SPO	
Actinomyces naeslundii		57.57	0.55	RHA 1	1	
Clostridium ramosum		40.10	0.27		92	
Actinomyces israelii		2.33	0.22		1	
27773223	**Very good identification**					**27773223**
Actinomyces naeslundii		100.0	0.58	RHA 1		
27773231	**Good identification**					**27773231**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27773233	**Good identification**					**27773233**
Actinomyces naeslundii		100.0	0.36	RHA 1		
27773240	**Low discrimination**					**27773240**
Actinomyces naeslundii		51.82	0.42	RHA 1		
Actinomyces israelii		48.18	0.32			
27773241	**Good identification**					**27773241**
Actinomyces naeslundii		100.0	0.45	RHA 1		
27773242	**Low discrimination**					**27773242**
Actinomyces israelii		58.24	0.35	XYL 92		
Actinomyces naeslundii		41.76	0.42	RHA 7		
27773243	**Good identification**					**27773243**
Actinomyces naeslundii		100.0	0.45	RHA 1		
27773260	**Good identification**					**27773260**
Actinomyces naeslundii		100.0	0.42	RHA 1		
27773261	**Good identification**					**27773261**
Actinomyces naeslundii		100.0	0.45	RHA 1		
27773262	**Acceptable identification**					**27773262**
Actinomyces naeslundii		91.83	0.42	RHA 1		
Actinomyces israelii		8.17	0.15			
27773263	**Good identification**					**27773263**
Actinomyces naeslundii		100.0	0.45	RHA 1		
27773300	**Low discrimination**					**27773300**
					SPO	
Actinomyces naeslundii		60.45	0.63	RHA 1	1	
Actinomyces israelii		35.85	0.50		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		3.70	0.17		92	
27773301	**Very good identification**					**27773301**
Actinomyces naeslundii		99.60	0.66	RHA 1		
Actinomyces israelii		0.40	0.16			
27773302	**Low discrimination**					**27773302**
					SPO	
Actinomyces naeslundii		46.20	0.63	RHA 1	1	
Actinomyces israelii		41.09	0.53		1	
Clostridium ramosum		11.31	0.27		92	
27773303	**Very good identification**					**27773303**
Actinomyces naeslundii		99.41	0.66	RHA 1		
Actinomyces israelii		0.60	0.19			
27773310	**Low discrimination**					**27773310**
Actinomyces israelii		62.74	0.36	XYL 92		
Actinomyces naeslundii		37.26	0.41	RHA 7		
27773311	**Good identification**					**27773311**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27773312	**Low discrimination**					**27773312**
Actinomyces israelii		71.63	0.39	XYL 92		
Actinomyces naeslundii		28.37	0.41	RHA 7		
27773313	**Good identification**					**27773313**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27773320	**Low discrimination**					**27773320**
					SPO	
Clostridium ramosum		85.37	0.51	NIT 1	92	
Actinomyces naeslundii		14.09	0.63		1	
Actinomyces israelii		0.53	0.29		1	
27773321	**Very good identification**					**27773321**
					SPO	
Actinomyces naeslundii		96.08	0.66	RHA 1	1	
Clostridium ramosum		3.92	0.17		92	
27773322	**Very good identification**					**27773322**
					SPO	
Clostridium ramosum		95.71	0.61	NIT 1	92	
Actinomyces naeslundii		3.95	0.63		1	
Actinomyces israelii		0.22	0.32		1	
27773323	**Low discrimination**					**27773323**
					SPO	
Actinomyces naeslundii		85.97	0.66	RHA 1	1	
Clostridium ramosum		14.03	0.27		92	
27773331	**Good identification**					**27773331**
Actinomyces naeslundii		100.0	0.44	RHA 1		
27773333	**Good identification**					**27773333**
Actinomyces naeslundii		100.0	0.44	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27773340	**Low discrimination**					**27773340**
Actinomyces israelii		56.65	0.43	XYL 92		
Actinomyces naeslundii		43.35	0.50	RHA 7		
27773341	**Good identification**					**27773341**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27773342	**Low discrimination**					**27773342**
Actinomyces israelii		66.22	0.46	XYL 92		
Actinomyces naeslundii		33.78	0.50	RHA 7		
27773343	**Good identification**					**27773343**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27773360	**Low discrimination**					**27773360**
					SPO	
Actinomyces naeslundii		69.92	0.50	RHA 1	1	
Clostridium ramosum		24.24	0.17		92	
Actinomyces israelii		5.83	0.22		1	
27773361	**Good identification**					**27773361**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27773363	**Good identification**					**27773363**
Actinomyces naeslundii		100.0	0.53	RHA 1		
27773400	**Low discrimination**					**27773400**
Actinomyces israelii		82.10	0.57	XYL 92		
Actinomyces naeslundii		17.90	0.55			
27773401	**Very good identification**					**27773401**
Actinomyces naeslundii		97.00	0.58	ARA 1		
Actinomyces israelii		3.00	0.23			
27773402	**Low discrimination**					**27773402**
Actinomyces israelii		87.31	0.60	XYL 92		
Actinomyces naeslundii		12.69	0.55			
27773403	**Very good identification**					**27773403**
Actinomyces naeslundii		95.57	0.58	ARA 1		
Actinomyces israelii		4.43	0.26			
27773410	**Acceptable identification**					**27773410**
Actinomyces israelii		92.87	0.43	XYL 92		
Actinomyces naeslundii		7.13	0.33	MLZ 13		
27773411	**Good identification**					**27773411**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27773412	**Good identification**					**27773412**
Actinomyces israelii		95.13	0.46	XYL 92		
Actinomyces naeslundii		4.87	0.33	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27773413	**Good identification**					**27773413**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27773420	**Low discrimination**					**27773420**
Actinomyces naeslundii		77.35	0.55	ARA 1		
Actinomyces israelii		22.65	0.37			
27773421	**Very good identification**					**27773421**
Actinomyces naeslundii		100.0	0.58	ARA 1		
27773422	**Low discrimination**					**27773422**
Actinomyces naeslundii		69.49	0.55	ARA 1		
Actinomyces israelii		30.51	0.40			
27773423	**Very good identification**					**27773423**
Actinomyces naeslundii		100.0	0.58	ARA 1		
27773431	**Good identification**					**27773431**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27773433	**Good identification**					**27773433**
Actinomyces naeslundii		100.0	0.36	ARA 1		
27773440	**Acceptable identification**					**27773440**
Actinomyces israelii		91.00	0.50	XYL 92		
Actinomyces naeslundii		9.00	0.42			
27773441	**Acceptable identification**					**27773441**
Actinomyces naeslundii		93.63	0.45	ARA 1		
Actinomyces israelii		6.37	0.16			
27773442	**Acceptable identification**					**27773442**
Actinomyces israelii		93.81	0.53	XYL 92		
Actinomyces naeslundii		6.19	0.42			
27773443	**Acceptable identification**					**27773443**
Actinomyces naeslundii		90.74	0.45	ARA 1		
Actinomyces israelii		9.26	0.19			
27773450	**Good identification**					**27773450**
Actinomyces israelii		100.0	0.36	XYL 92		
27773452	**Good identification**					**27773452**
Actinomyces israelii		100.0	0.39	XYL 92		
27773460	**Low discrimination**					**27773460**
Actinomyces naeslundii		60.78	0.42	ARA 1		
Actinomyces israelii		39.22	0.30			
27773461	**Good identification**					**27773461**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	ARA 1		
27773462	**Low discrimination**					**27773462**
Actinomyces naeslundii		50.82	0.42	ARA 1		
Actinomyces israelii		49.18	0.33			
27773463	**Good identification**					**27773463**
Actinomyces naeslundii		100.0	0.45	ARA 1		
27773500	**Low discrimination**					**27773500**
					GRA	
Actinomyces israelii		84.36	0.68	XYL 92	99	
Actinomyces naeslundii		13.09	0.63		99	
Mitsuokella multacida		2.01	0.27		1	
27773501	**Good identification**					**27773501**
					GRA	
Actinomyces naeslundii		94.80	0.66	ARA 1	99	
Actinomyces israelii		4.12	0.33		99	
Mitsuokella multacida		1.08	0.11		1	
27773502	**Low discrimination**					**27773502**
Actinomyces israelii		88.68	0.71	XYL 92		
Actinomyces naeslundii		9.17	0.63			
Bifidobacterium adolescentis		1.84	0.35			
27773503	**Good identification**					**27773503**
Actinomyces naeslundii		93.89	0.66	ARA 1		
Actinomyces israelii		6.11	0.36			
27773510	**Acceptable identification**					**27773510**
					GRA	
Actinomyces israelii		90.26	0.53	XYL 92	99	
Actinomyces naeslundii		4.93	0.41	MLZ 13	99	
Mitsuokella multacida		4.81	0.19		1	
27773511	**Low discrimination**					**27773511**
					GRA	
Actinomyces naeslundii		83.65	0.44	ARA 1	99	
Actinomyces israelii		10.31	0.19	MLZ 5	99	
Mitsuokella multacida		6.04	0.03		1	
27773512	**Low discrimination**					**27773512**
Actinomyces israelii		82.25	0.56	XYL 92		
Bifidobacterium adolescentis		13.98	0.37	MLZ 13		
Actinomyces naeslundii		3.00	0.41			
27773513	**Low discrimination**					**27773513**
Actinomyces naeslundii		84.40	0.44	ARA 1		
Actinomyces israelii		15.60	0.22	MLZ 5		
27773520	**Low discrimination**					**27773520**
					SPO	
Actinomyces naeslundii		66.84	0.63	ARA 1	1	
Actinomyces israelii		27.50	0.47		1	
Clostridium ramosum		3.09	0.15		92	
27773521	**Very good identification**					**27773521**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.66	ARA 1		
27773522	**Low discrimination**					**27773522**
					SPO	
Actinomyces naeslundii		55.50	0.63	ARA 1	1	
Actinomyces israelii		34.25	0.50		1	
Clostridium ramosum		10.25	0.25		92	
27773523	**Very good identification**					**27773523**
Actinomyces naeslundii		99.59	0.66	ARA 1		
Actinomyces israelii		0.41	0.16			
27773531	**Good identification**					**27773531**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27773532	**Low discrimination**					**27773532**
Actinomyces israelii		58.10	0.36	XYL 92		
Actinomyces naeslundii		33.16	0.41	MLZ 13		
Bifidobacterium dentium		8.74	-.03	NAG 6		
27773533	**Good identification**					**27773533**
Actinomyces naeslundii		100.0	0.44	ARA 1		
27773540	**Low discrimination**					**27773540**
					GRA	
Actinomyces israelii		88.36	0.61	XYL 92	99	
Actinomyces naeslundii		6.22	0.50		99	
Mitsuokella multacida		5.42	0.27		1	
27773541	**Low discrimination**					**27773541**
					GRA	
Actinomyces naeslundii		86.19	0.53	ARA 1	99	
Actinomyces israelii		8.24	0.26		99	
Mitsuokella multacida		5.57	0.11		1	
27773542	**Good identification**					**27773542**
Actinomyces israelii		94.51	0.64	XYL 92		
Actinomyces naeslundii		4.44	0.50			
Bifidobacterium adolescentis		0.62	0.20			
27773543	**Low discrimination**					**27773543**
Actinomyces naeslundii		87.45	0.53	ARA 1		
Actinomyces israelii		12.55	0.29			
27773550	**Low discrimination**					**27773550**
					GRA	
Actinomyces israelii		84.58	0.46	XYL 92	99	
Mitsuokella multacida		11.58	0.19	MLZ 13	1	
Actinomyces naeslundii		2.10	0.28		99	
27773552	**Acceptable identification**					**27773552**
Actinomyces israelii		92.53	0.49	XYL 92		
Bifidobacterium adolescentis		5.00	0.21	MLZ 13		
Actinomyces naeslundii		1.53	0.28			
27773560	**Low discrimination**					**27773560**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		47.07	0.50	ARA 1	99	
Actinomyces israelii		42.67	0.40		99	
Mitsuokella multacida		10.26	0.17		1	
27773561	**Good identification**					**27773561**
Actinomyces naeslundii		100.0	0.53	ARA 1		
27773562	**Low discrimination**					**27773562**
Actinomyces israelii		57.63	0.43	XYL 92		
Actinomyces naeslundii		42.37	0.50	NAG 6		
27773563	**Good identification**					**27773563**
Actinomyces naeslundii		100.0	0.53	ARA 1		
27773600	**Good identification**					**27773600**
Actinomyces israelii		100.0	0.38	XYL 92		
27773602	**Good identification**					**27773602**
Actinomyces israelii		100.0	0.41	XYL 92		
27773700	**Low discrimination**					**27773700**
					GRA	
Actinomyces israelii		62.12	0.48	XYL 92	99	
Mitsuokella multacida		36.59	0.32	RHA 7	1	
Actinomyces naeslundii		1.29	0.29		99	
27773702	**Acceptable identification**					**27773702**
					GRA	
Actinomyces israelii		94.56	0.51	XYL 92	99	
Mitsuokella multacida		4.13	0.15	RHA 7	1	
Actinomyces naeslundii		1.31	0.29		99	
27773712	**Low discrimination**					**27773712**
					GRA	
Actinomyces israelii		87.12	0.37	XYL 92	99	
Mitsuokella multacida		8.48	0.07	RHA 7	1	
Clostridium clostridioforme		4.40	0.06	MLZ 13	50	
27773742	**Low discrimination**					**27773742**
					GRA	
Actinomyces israelii		89.91	0.44	XYL 92	99	
Mitsuokella multacida		10.09	0.15	RHA 7	1	
27774100	**Low discrimination**					**27774100**
					SPO	
Actinomyces israelii		82.23	0.40	GAL 99	1	
Actinomyces naeslundii		10.93	0.34		1	
Clostridium butyricum		6.84	0.01		99	
27774101	**Good identification**					**27774101**
Actinomyces naeslundii		100.0	0.37	GAL 96		
27774102	**Low discrimination**					**27774102**
Actinomyces israelii		51.05	0.43	GAL 99		
Actinomyces viscosus		43.33	0.27			
Actinomyces naeslundii		4.52	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27774121	**Good identification**						**27774121**
Actinomyces naeslundii		100.0	0.37	GAL 96			
27774123	**Good identification**						**27774123**
Actinomyces naeslundii		100.0	0.37	GAL 96			
27774142	**Good identification**						**27774142**
Actinomyces israelii		100.0	0.36	GAL 99			
27774500	**Low discrimination**						**27774500**
					GRA	SPO	
Actinomyces israelii		41.43	0.39	GAL 99	99	1	
Clostridium butyricum		37.90	0.17		99	99	
Mitsuokella multacida		20.68	0.21		1	1	
27774502	**Low discrimination**						**27774502**
					SPO		
Actinomyces israelii		61.18	0.42	GAL 99	1		
Clostridium clostridioforme		16.93	0.23		66		
Clostridium butyricum		14.51	0.10		99		
27774612	**Good identification**						**27774612**
Clostridium clostridioforme		100.0	0.40	NIT 1			
27774632	**Good identification**						**27774632**
Clostridium clostridioforme		100.0	0.40	NIT 1			
27774702	**Acceptable identification**						**27774702**
					GRA		
Clostridium clostridioforme		93.38	0.37	NIT 1	50		
Actinomyces israelii		3.46	0.22		99		
Mitsuokella multacida		3.16	0.09		1		
27774712	**Good identification**						**27774712**
Clostridium clostridioforme		99.75	0.49	NIT 1			
Mitsuokella multacida		0.25	0.01				
27774722	**Good identification**						**27774722**
Clostridium clostridioforme		97.04	0.37	NIT 1			
Parabacteroides distasonis		2.96	0.17				
27774732	**Good identification**						**27774732**
Clostridium clostridioforme		99.64	0.49	NIT 1			
Parabacteroides distasonis		0.36	0.13				
27775000	**Good identification**						**27775000**
					SPO		
Actinomyces israelii		94.42	0.64		1		
Actinomyces naeslundii		4.28	0.50		1		
Clostridium butyricum		1.31	0.11		99		
27775001	**Low discrimination**						**27775001**
Actinomyces naeslundii		87.06	0.53	XYL 1			
Actinomyces israelii		12.95	0.30				
27775002	**Very good identification**						**27775002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		95.54	0.67			
Actinomyces naeslundii		2.88	0.50			
Actinomyces viscosus		1.13	0.19			
27775003	**Low discrimination**					**27775003**
Actinomyces naeslundii		76.33	0.53	XYL 1		
Actinomyces israelii		17.03	0.33			
Actinomyces viscosus		6.64	0.11			
27775010	**Good identification**					**27775010**
Actinomyces israelii		98.43	0.50	MLZ 13		
Actinomyces naeslundii		1.57	0.28			
27775012	**Good identification**					**27775012**
Actinomyces israelii		97.41	0.53	MLZ 13		
Clostridium clostridioforme		1.55	0.13			
Actinomyces naeslundii		1.04	0.28			
27775020	**Low discrimination**					**27775020**
					SPO	
Actinomyces israelii		57.86	0.44	NAG 6	1	
Actinomyces naeslundii		41.05	0.50		1	
Clostridium butyricum		1.09	-.07		99	
27775021	**Good identification**					**27775021**
Actinomyces naeslundii		100.0	0.53	XYL 1		
27775022	**Low discrimination**					**27775022**
					GRA	
Actinomyces israelii		66.31	0.47	NAG 6	99	
Actinomyces naeslundii		31.36	0.50		99	
Parabacteroides distasonis		1.16	0.13		1	
27775023	**Good identification**					**27775023**
Actinomyces naeslundii		100.0	0.53	XYL 1		
27775040	**Very good identification**					**27775040**
					SPO	
Actinomyces israelii		97.69	0.57		1	
Actinomyces naeslundii		2.01	0.37		1	
Clostridium butyricum		0.30	-.07		99	
27775041	**Low discrimination**					**27775041**
Actinomyces naeslundii		75.32	0.40	XYL 1		
Actinomyces israelii		24.68	0.23			
27775042	**Very good identification**					**27775042**
Actinomyces israelii		98.65	0.60			
Actinomyces naeslundii		1.35	0.37			
27775043	**Low discrimination**					**27775043**
Actinomyces naeslundii		67.04	0.40	XYL 1		
Actinomyces israelii		32.96	0.26			
27775050	**Good identification**					**27775050**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.43	MLZ 13		
27775052	**Good identification**					**27775052**
Actinomyces israelii		100.0	0.46	MLZ 13		
27775060	**Low discrimination**					**27775060**
Actinomyces israelii		75.65	0.36	NAG 6		
Actinomyces naeslundii		24.35	0.37			
27775061	**Good identification**					**27775061**
Actinomyces naeslundii		100.0	0.40	XYL 1		
27775062	**Low discrimination**					**27775062**
Actinomyces israelii		82.33	0.40	NAG 6		
Actinomyces naeslundii		17.67	0.37			
27775063	**Good identification**					**27775063**
Actinomyces naeslundii		100.0	0.40	XYL 1		
27775100	**Low discrimination**					**27775100**
					SPO	
Actinomyces israelii		88.66	0.74		1	
Clostridium butyricum		7.37	0.35		99	
Actinomyces naeslundii		2.86	0.58		1	
27775101	**Low discrimination**					**27775101**
					SPO	
Actinomyces naeslundii		80.92	0.61	XYL 1	1	
Actinomyces israelii		16.91	0.40		1	
Clostridium butyricum		1.41	0.01		99	
27775102	**Good identification**					**27775102**
					SPO	
Actinomyces israelii		94.01	0.77		1	
Clostridium butyricum		2.03	0.28		99	
Actinomyces naeslundii		2.02	0.58		1	
27775103	**Low discrimination**					**27775103**
Actinomyces naeslundii		72.75	0.61	XYL 1		
Actinomyces israelii		22.81	0.43			
Actinomyces viscosus		3.96	0.15			
27775110	**Very good identification**					**27775110**
					SPO	
Actinomyces israelii		95.25	0.60	MLZ 13	1	
Clostridium butyricum		2.79	0.13		99	
Actinomyces naeslundii		1.08	0.36		1	
27775111	**Low discrimination**					**27775111**
Actinomyces naeslundii		62.76	0.39	XYL 1		
Actinomyces israelii		37.24	0.26	MLZ 5		
27775112	**Low discrimination**					**27775112**
					GRA	
Actinomyces israelii		88.66	0.63	MLZ 13	99	
Bifidobacterium adolescentis		7.43	0.38		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Leptotrichia buccalis		1.38	0.07		1	
27775113	**Low discrimination**					**27775113**
Actinomyces naeslundii		52.91	0.39	XYL 1		
Actinomyces israelii		47.09	0.29	MLZ 5		
27775120	**Low discrimination**					**27775120**
					SPO	
Actinomyces israelii		59.18	0.54	NAG 6	1	
Actinomyces naeslundii		29.87	0.58		1	
Clostridium butyricum		6.71	0.17		99	
27775121	**Very good identification**					**27775121**
Actinomyces naeslundii		98.68	0.61	XYL 1		
Actinomyces israelii		1.32	0.20			
27775122	**Low discrimination**					**27775122**
					GRA	
Actinomyces israelii		51.93	0.57	NAG 6	99	
Parabacteroides distasonis		22.44	0.47		1	
Actinomyces naeslundii		17.48	0.58		99	
27775123	**Very good identification**					**27775123**
					GRA	
Actinomyces naeslundii		97.21	0.61	XYL 1	99	
Actinomyces israelii		1.95	0.23		99	
Parabacteroides distasonis		0.84	0.13		1	
27775130	**Low discrimination**					**27775130**
					SPO	
Actinomyces israelii		79.76	0.40	MLZ 13	1	
Actinomyces naeslundii		14.18	0.36	NAG 6	1	
Clostridium butyricum		3.18	-.05		99	
27775131	**Good identification**					**27775131**
Actinomyces naeslundii		100.0	0.39	XYL 1		
27775132	**Low discrimination**					**27775132**
					GRA	
Parabacteroides distasonis		53.47	0.42	NIT 1	1	
Actinomyces israelii		34.35	0.43		99	
Clostridium clostridioforme		7.18	0.22		50	
27775133	**Good identification**					**27775133**
Actinomyces naeslundii		100.0	0.39	XYL 1		
27775140	**Very good identification**					**27775140**
					SPO	
Actinomyces israelii		95.80	0.67		1	
Clostridium butyricum		1.78	0.17		99	
Actinomyces naeslundii		1.40	0.45		1	
27775141	**Low discrimination**					**27775141**
					GRA	
Actinomyces naeslundii		66.04	0.48	XYL 1	99	
Actinomyces israelii		30.42	0.33		99	
Mitsuokella multacida		3.55	0.04		1	
27775142	**Very good identification**					**27775142**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		98.07	0.70		1		
Actinomyces naeslundii		0.96	0.45		1		
Clostridium butyricum		0.47	0.10		99		
27775143	**Low discrimination**						**27775143**
Actinomyces naeslundii		59.14	0.48	XYL 1			
Actinomyces israelii		40.86	0.36				
27775150	**Good identification**						**27775150**
					GRA	SPO	
Actinomyces israelii		96.59	0.53	MLZ 13	99	1	
Mitsuokella multacida		2.29	0.13		1	1	
Clostridium butyricum		0.63	-.05		99	99	
27775152	**Very good identification**						**27775152**
Actinomyces israelii		97.08	0.56	MLZ 13			
Bifidobacterium adolescentis		2.59	0.23				
Actinomyces naeslundii		0.33	0.23				
27775160	**Low discrimination**						**27775160**
					GRA		
Actinomyces israelii		77.16	0.47	NAG 6	99		
Actinomyces naeslundii		17.68	0.45		99		
Mitsuokella multacida		3.21	0.11		1		
27775161	**Good identification**						**27775161**
Actinomyces naeslundii		100.0	0.48	XYL 1			
27775162	**Low discrimination**						**27775162**
Actinomyces israelii		84.75	0.50	NAG 6			
Actinomyces naeslundii		12.94	0.45				
Lactobacillus acidophilus		1.36	0.04				
27775163	**Good identification**						**27775163**
Actinomyces naeslundii		95.78	0.48	XYL 1			
Actinomyces israelii		4.22	0.16				
27775172	**Acceptable identification**						**27775172**
Actinomyces israelii		94.90	0.36	MLZ 13			
Actinomyces naeslundii		5.10	0.23	NAG 6			
27775200	**Good identification**						**27775200**
Actinomyces israelii		100.0	0.45	RHA 7			
27775202	**Good identification**						**27775202**
Actinomyces israelii		95.15	0.48	RHA 7			
Clostridium clostridioforme		4.85	0.16				
27775240	**Good identification**						**27775240**
Actinomyces israelii		100.0	0.38	RHA 7			
27775242	**Good identification**						**27775242**
Actinomyces israelii		100.0	0.41	RHA 7			
27775300	**Low discrimination**						**27775300**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				GRA	SPO	
Actinomyces israelii	87.70	0.55	RHA 7	99	1	
Mitsuokella multacida	8.93	0.25		1	1	
Clostridium butyricum	3.00	0.09		99	99	
27775302 Good identification						**27775302**
				GRA		
Actinomyces israelii	92.91	0.58	RHA 7	99		
Clostridium clostridioforme	3.96	0.25		50		
Parabacteroides distasonis	1.03	0.21		1		
27775310 Low discrimination						**27775310**
				GRA		
Actinomyces israelii	81.50	0.41	RHA 7	99		
Mitsuokella multacida	18.50	0.17	MLZ 13	1		
27775312 Low discrimination						**27775312**
				GRA		
Clostridium clostridioforme	55.17	0.36	NIT 1	50		
Actinomyces israelii	42.42	0.44	ARA 94	99		
Parabacteroides distasonis	1.70	0.16		1		
27775322 Low discrimination						**27775322**
				GRA		
Parabacteroides distasonis	87.57	0.55	NIT 1	1		
Actinomyces israelii	5.09	0.38		99		
Clostridium clostridioforme	3.40	0.25		50		
27775332 Low discrimination						**27775332**
				GRA		
Parabacteroides distasonis	74.14	0.50	NIT 1	1		
Clostridium clostridioforme	24.35	0.36		50		
Actinomyces israelii	1.20	0.23		99		
27775340 Low discrimination						**27775340**
				GRA		
Actinomyces israelii	79.26	0.48	RHA 7	99		
Mitsuokella multacida	20.74	0.25		1		
27775342 Good identification						**27775342**
				GRA		
Actinomyces israelii	98.10	0.51	RHA 7	99		
Mitsuokella multacida	1.90	0.09		1		
27775352 Good identification						**27775352**
				GRA		
Actinomyces israelii	95.85	0.37	RHA 7	99		
Mitsuokella multacida	4.15	0.01	MLZ 13	1		
27775400 Low discrimination						**27775400**
				GRA	SPO	
Actinomyces israelii	85.28	0.62		99	1	
Clostridium butyricum	13.00	0.27		99	99	
Mitsuokella multacida	1.72	0.21		1	1	
27775402 Good identification						**27775402**
				SPO		
Actinomyces israelii	94.84	0.65		1		
Clostridium butyricum	3.75	0.20		99		
Clostridium clostridioforme	0.95	0.22		66		
27775410 Acceptable identification						**27775410**
				GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		91.02	0.48	MLZ 13	99	1	
Clostridium butyricum		4.89	0.06		99	99	
Mitsuokella multacida		4.10	0.13		1	1	
27775412	**Low discrimination**						**27775412**
Actinomyces israelii		74.24	0.51	MLZ 13			
Clostridium clostridioforme		22.67	0.33				
Bifidobacterium adolescentis		2.06	0.18				
27775420	**Low discrimination**						**27775420**
					GRA	SPO	
Actinomyces israelii		77.72	0.42	NAG 6	99	1	
Clostridium butyricum		16.14	0.09		99	99	
Mitsuokella multacida		6.14	0.11		1	1	
27775422	**Low discrimination**						**27775422**
					SPO		
Actinomyces israelii		82.59	0.45	NAG 6	1		
Clostridium clostridioforme		12.96	0.22		66		
Clostridium butyricum		4.45	0.02		99		
27775440	**Acceptable identification**						**27775440**
					GRA	SPO	
Actinomyces israelii		92.09	0.55		99	1	
Mitsuokella multacida		4.78	0.21		1	1	
Clostridium butyricum		3.14	0.09		99	99	
27775442	**Very good identification**						**27775442**
					GRA	SPO	
Actinomyces israelii		98.75	0.58		99	1	
Clostridium butyricum		0.87	0.02		99	99	
Mitsuokella multacida		0.38	0.04		1	1	
27775450	**Low discrimination**						**27775450**
					GRA		
Actinomyces israelii		89.63	0.41	MLZ 13	99		
Mitsuokella multacida		10.37	0.13		1		
27775452	**Good identification**						**27775452**
Actinomyces israelii		100.0	0.44	MLZ 13			
27775460	**Low discrimination**						**27775460**
					GRA		
Actinomyces israelii		83.12	0.35	NAG 6	99		
Mitsuokella multacida		16.88	0.11		1		
27775462	**Good identification**						**27775462**
Actinomyces israelii		100.0	0.38	NAG 6			
27775500	**Low discrimination**						**27775500**
					GRA	SPO	
Actinomyces israelii		41.40	0.73		99	1	
Clostridium butyricum		37.87	0.51		99	99	
Mitsuokella multacida		20.66	0.55		1	1	
27775501	**Low discrimination**						**27775501**
					GRA	SPO	
Mitsuokella multacida		73.56	0.39	ESL 99	1	1	
Actinomyces israelii		13.40	0.39	URE 10	99	1	
Clostridium butyricum		12.26	0.17		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27775502	**Low discrimination**						**27775502**
					SPO		
Actinomyces israelii		73.18	0.76		1		
Clostridium butyricum		17.36	0.44		99		
Bifidobacterium adolescentis		6.13	0.51		1		
27775503	**Low discrimination**						**27775503**
					GRA	SPO	
Actinomyces israelii		56.59	0.42	URE 1	99	1	
Mitsuokella multacida		23.02	0.22		1	1	
Clostridium butyricum		13.42	0.10		99	99	
27775510	**Low discrimination**						**27775510**
					GRA	SPO	
Mitsuokella multacida		45.45	0.47	ESL 99	1	1	
Actinomyces israelii		40.82	0.59		99	1	
Clostridium butyricum		13.15	0.29		99	99	
27775512	**Low discrimination**						**27775512**
Actinomyces israelii		47.40	0.62	MLZ 13			
Bifidobacterium adolescentis		32.48	0.53				
Clostridium clostridioforme		12.12	0.42				
27775520	**Low discrimination**						**27775520**
					GRA	SPO	
Mitsuokella multacida		46.46	0.45	ESL 99	1	1	
Clostridium butyricum		29.62	0.33		99	99	
Actinomyces israelii		23.77	0.52		99	1	
27775522	**Low discrimination**						**27775522**
					GRA	SPO	
Actinomyces israelii		51.29	0.55	NAG 6	99	1	
Parabacteroides distasonis		16.60	0.43		1	1	
Clostridium butyricum		16.57	0.26		99	99	
27775530	**Low discrimination**						**27775530**
					GRA	SPO	
Mitsuokella multacida		74.41	0.37	ESL 99	1	1	
Actinomyces israelii		17.06	0.38		99	1	
Clostridium butyricum		7.49	0.11		99	99	
27775532	**Low discrimination**						**27775532**
					GRA		
Clostridium clostridioforme		58.61	0.42	NIT 1	50		
Parabacteroides distasonis		17.07	0.39	RHA 88	1		
Actinomyces israelii		14.64	0.41		99		
27775540	**Low discrimination**						**27775540**
					GRA	SPO	
Mitsuokella multacida		51.59	0.55	ESL 99	1	1	
Actinomyces israelii		40.19	0.66		99	1	
Clostridium butyricum		8.22	0.33		99	99	
27775541	**Acceptable identification**						**27775541**
					GRA	SPO	
Mitsuokella multacida		92.14	0.39	ESL 99	1	1	
Actinomyces israelii		6.53	0.32	URE 10	99	1	
Clostridium butyricum		1.34	-.01		99	99	
27775542	**Low discrimination**						**27775542**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		85.13	0.69		99	1	
Mitsuokella multacida		8.09	0.39		1	1	
Clostridium butyricum		4.52	0.26		99	99	
27775550	**Low discrimination**						**27775550**
					GRA	SPO	
Mitsuokella multacida		72.57	0.47	ESL 99	1	1	
Actinomyces israelii		25.35	0.52		99	1	
Clostridium butyricum		1.83	0.11		99	99	
27775552	**Low discrimination**						**27775552**
					GRA		
Actinomyces israelii		68.72	0.55	MLZ 13	99		
Bifidobacterium adolescentis		14.97	0.37		99		
Mitsuokella multacida		14.58	0.30		1		
27775560	**Low discrimination**						**27775560**
					GRA	SPO	
Mitsuokella multacida		79.72	0.45	ESL 99	1	1	
Actinomyces israelii		15.86	0.45		99	1	
Clostridium butyricum		4.42	0.15		99	99	
27775562	**Low discrimination**						**27775562**
					GRA	SPO	
Actinomyces israelii		69.22	0.48	NAG 6	99	1	
Mitsuokella multacida		25.78	0.28		1	1	
Clostridium butyricum		5.00	0.08		99	99	
27775570	**Acceptable identification**						**27775570**
					GRA	SPO	
Mitsuokella multacida		91.08	0.37	ESL 99	1	1	
Actinomyces israelii		8.12	0.31		99	1	
Clostridium butyricum		0.80	-.07		99	99	
27775600	**Low discrimination**						**27775600**
					GRA	SPO	
Actinomyces israelii		64.09	0.43	RHA 7	99	1	
Mitsuokella multacida		31.89	0.25		1	1	
Clostridium butyricum		4.02	0.02		99	99	
27775602	**Low discrimination**						**27775602**
					GRA		
Actinomyces israelii		49.28	0.46	RHA 7	99		
Clostridium clostridioforme		48.10	0.37		50		
Mitsuokella multacida		1.82	0.09		1		
27775612	**Good identification**						**27775612**
					GRA		
Clostridium clostridioforme		96.49	0.48	NIT 1	50		
Actinomyces israelii		3.24	0.32		99		
Mitsuokella multacida		0.27	0.01		1		
27775622	**Acceptable identification**						**27775622**
					GRA		
Clostridium clostridioforme		90.95	0.37	NIT 1	50		
Actinomyces israelii		5.95	0.26		99		
Parabacteroides distasonis		3.10	0.17		1		
27775632	**Good identification**						**27775632**
Clostridium clostridioforme		99.60	0.48	NIT 1			
Parabacteroides distasonis		0.40	0.13				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27775642	**Acceptable identification**						**27775642**
					GRA		
Actinomyces israelii		91.34	0.39	RHA 7	99		
Mitsuokella multacida		8.66	0.09		1		
27775700	**Low discrimination**						**27775700**
					GRA	SPO	
Mitsuokella multacida		89.87	0.60	ESL 99	1	1	
Actinomyces israelii		7.30	0.54		99	1	
Clostridium butyricum		2.74	0.25		99	99	
27775701	**Good identification**						**27775701**
					GRA		
Mitsuokella multacida		99.27	0.43	ESL 99	1		
Actinomyces israelii		0.73	0.19	URE 10	99		
27775702	**Low discrimination**						**27775702**
					GRA		
Actinomyces israelii		35.14	0.57	RHA 7	99		
Mitsuokella multacida		32.05	0.43		1		
Clostridium clostridioforme		28.70	0.46		50		
27775710	**Acceptable identification**						**27775710**
					GRA		
Mitsuokella multacida		94.79	0.51	ESL 99	1		
Actinomyces israelii		3.45	0.39		99		
Clostridium clostridioforme		1.30	0.23		50		
27775711	**Good identification**						**27775711**
Mitsuokella multacida		100.0	0.35	ESL 99			
27775712	**Low discrimination**						**27775712**
					GRA		
Clostridium clostridioforme		88.64	0.57	NIT 1	50		
Mitsuokella multacida		7.24	0.35		1		
Actinomyces israelii		3.56	0.42		99		
27775720	**Good identification**						**27775720**
					GRA	SPO	
Mitsuokella multacida		95.48	0.49	ESL 99	1	1	
Actinomyces israelii		1.98	0.33		99	1	
Clostridium butyricum		1.01	0.07		99	99	
27775722	**Low discrimination**						**27775722**
Parabacteroides distasonis		39.20	0.51	NIT 1			
Clostridium clostridioforme		38.89	0.46				
Mitsuokella multacida		10.86	0.33				
27775730	**Acceptable identification**						**27775730**
					GRA		
Mitsuokella multacida		93.39	0.41	ESL 99	1		
Clostridium clostridioforme		5.13	0.23		50		
Actinomyces israelii		0.87	0.19		99		
27775732	**Low discrimination**						**27775732**
Clostridium clostridioforme		85.46	0.57	NIT 1			
Parabacteroides distasonis		10.18	0.47				
Bacteroides ovatus		2.33	0.20				
27775740	**Very good identification**						**27775740**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Mitsuokella multacida		96.69	0.60	ESL 99	1	1	
Actinomyces israelii		3.05	0.47		99	1	
Clostridium butyricum		0.26	0.07		99	99	
27775741	**Good identification**						**27775741**
Mitsuokella multacida		100.0	0.43	ESL 99			
27775742	**Low discrimination**						**27775742**
					GRA	SPO	
Mitsuokella multacida		69.22	0.43	ESL 99	1	1	
Actinomyces israelii		29.51	0.50	BGL 10	99	1	
Clostridium butyricum		0.64	0.00		99	99	
27775750	**Good identification**						**27775750**
					COC	GRA	
Mitsuokella multacida		98.42	0.51	ESL 99	1	1	
Actinomyces israelii		1.39	0.32		1	99	
Blautia producta		0.19	-.03		99	99	
27775751	**Good identification**						**27775751**
Mitsuokella multacida		100.0	0.35	ESL 99			
27775752	**Low discrimination**						**27775752**
					GRA		
Mitsuokella multacida		76.06	0.35	ESL 99	1		
Actinomyces israelii		14.54	0.35	BGL 10	99		
Clostridium clostridioforme		9.40	0.23		50		
27775760	**Good identification**						**27775760**
					GRA		
Mitsuokella multacida		99.20	0.49	ESL 99	1		
Actinomyces israelii		0.80	0.26		99		
27775770	**Good identification**						**27775770**
Mitsuokella multacida		100.0	0.41	ESL 99			
27776000	**Acceptable identification**						**27776000**
Actinomyces israelii		93.99	0.43	GAL 99			
Actinomyces naeslundii		6.01	0.31				
27776002	**Good identification**						**27776002**
Actinomyces israelii		95.91	0.46	GAL 99			
Actinomyces naeslundii		4.09	0.31				
27776040	**Good identification**						**27776040**
Actinomyces israelii		100.0	0.36	GAL 99			
27776042	**Good identification**						**27776042**
Actinomyces israelii		100.0	0.39	GAL 99			
27776100	**Good identification**						**27776100**
Actinomyces israelii		95.65	0.53	GAL 99			
Actinomyces naeslundii		4.36	0.39				
27776101	**Low discrimination**						**27776101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		87.11	0.42	GAL 96			
Actinomyces israelii		12.89	0.19	XYL 1			
27776102	**Good identification**						**27776102**
Actinomyces israelii		97.05	0.56	GAL 99			
Actinomyces naeslundii		2.95	0.39				
27776103	**Low discrimination**						**27776103**
Actinomyces naeslundii		81.84	0.42	GAL 96			
Actinomyces israelii		18.16	0.22	XYL 1			
27776110	**Good identification**						**27776110**
Actinomyces israelii		100.0	0.39	GAL 99			
27776112	**Good identification**						**27776112**
Actinomyces israelii		100.0	0.42	GAL 99			
27776121	**Good identification**						**27776121**
Actinomyces naeslundii		100.0	0.42	GAL 96			
27776122	**Low discrimination**						**27776122**
Actinomyces israelii		67.77	0.36	GAL 99			
Actinomyces naeslundii		32.23	0.39	NAG 6			
27776123	**Good identification**						**27776123**
Actinomyces naeslundii		100.0	0.42	GAL 96			
27776140	**Good identification**						**27776140**
Actinomyces israelii		97.98	0.46	GAL 99			
Actinomyces naeslundii		2.02	0.26				
27776142	**Good identification**						**27776142**
Actinomyces israelii		98.64	0.49	GAL 99			
Actinomyces naeslundii		1.36	0.26				
27776302	**Good identification**						**27776302**
Actinomyces israelii		100.0	0.37	GAL 99			
27776400	**Good identification**						**27776400**
Actinomyces israelii		100.0	0.41	GAL 99			
27776402	**Good identification**						**27776402**
Actinomyces israelii		100.0	0.44	GAL 99			
27776442	**Good identification**						**27776442**
Actinomyces israelii		100.0	0.37	GAL 99			
27776500	**Low discrimination**						**27776500**
					GRA	SPO	
Actinomyces israelii		81.74	0.52	GAL 99	99	1	
Mitsuokella multacida		16.80	0.27		1	1	
Clostridium butyricum		1.47	0.01		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
27776502	**Good identification**					**27776502**
					GRA	
Actinomyces israelii		96.90	0.55	GAL 99	99	
Mitsuokella multacida		1.48	0.11		1	
Bifidobacterium adolescentis		1.17	0.15		99	
27776510	**Low discrimination**					**27776510**
					GRA	
Actinomyces israelii		68.57	0.37	GAL 99	99	
Mitsuokella multacida		31.44	0.19	MLZ 13	1	
27776512	**Low discrimination**					**27776512**
Actinomyces israelii		84.79	0.40	GAL 99		
Bifidobacterium adolescentis		8.39	0.17	MLZ 13		
Bifidobacterium dentium		3.94	-.07			
27776540	**Low discrimination**					**27776540**
					GRA	
Actinomyces israelii		65.42	0.44	GAL 99	99	
Mitsuokella multacida		34.58	0.27		1	
27776542	**Good identification**					**27776542**
					GRA	
Actinomyces israelii		96.23	0.48	GAL 99	99	
Mitsuokella multacida		3.77	0.11		1	
27776702	**Low discrimination**					**27776702**
					GRA	
Actinomyces israelii		72.69	0.35	GAL 99	99	
Mitsuokella multacida		27.31	0.15	RHA 7	1	
27777000	**Very good identification**					**27777000**
					SPO	
Actinomyces israelii		98.45	0.77		1	
Actinomyces naeslundii		1.53	0.55		1	
Clostridium butyricum		0.03	-.05		99	
27777001	**Low discrimination**					**27777001**
Actinomyces naeslundii		69.73	0.58	XYL 1		
Actinomyces israelii		30.27	0.43			
27777002	**Very good identification**					**27777002**
Actinomyces israelii		98.98	0.80			
Actinomyces naeslundii		1.02	0.55			
27777003	**Low discrimination**					**27777003**
Actinomyces naeslundii		60.56	0.58	XYL 1		
Actinomyces israelii		39.44	0.46			
27777010	**Very good identification**					**27777010**
Actinomyces israelii		99.46	0.63	MLZ 13		
Actinomyces naeslundii		0.54	0.33			
27777012	**Very good identification**					**27777012**
Actinomyces israelii		99.64	0.66	MLZ 13		
Actinomyces naeslundii		0.36	0.33			
27777020	**Low discrimination**					**27777020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		80.45	0.56	NAG 6		
Actinomyces naeslundii		19.55	0.55			
27777021	**Very good identification**					**27777021**
Actinomyces naeslundii		97.30	0.58	XYL 1		
Actinomyces israelii		2.70	0.22			
27777022	**Low discrimination**					**27777022**
Actinomyces israelii		86.06	0.59	NAG 6		
Actinomyces naeslundii		13.94	0.55			
27777023	**Very good identification**					**27777023**
Actinomyces naeslundii		96.01	0.58	XYL 1		
Actinomyces israelii		3.99	0.25			
27777030	**Acceptable identification**					**27777030**
Actinomyces israelii		92.12	0.42	MLZ 13		
Actinomyces naeslundii		7.89	0.33	NAG 6		
27777031	**Good identification**					**27777031**
Actinomyces naeslundii		100.0	0.36	XYL 1		
27777032	**Acceptable identification**					**27777032**
Actinomyces israelii		94.60	0.45	MLZ 13		
Actinomyces naeslundii		5.40	0.33	NAG 6		
27777033	**Good identification**					**27777033**
Actinomyces naeslundii		100.0	0.36	XYL 1		
27777040	**Very good identification**					**27777040**
Actinomyces israelii		99.30	0.70			
Actinomyces naeslundii		0.70	0.42			
27777041	**Low discrimination**					**27777041**
Actinomyces naeslundii		51.11	0.45	XYL 1		
Actinomyces israelii		48.89	0.36			
27777042	**Very good identification**					**27777042**
Actinomyces israelii		99.53	0.73			
Actinomyces naeslundii		0.47	0.42			
27777043	**Low discrimination**					**27777043**
Actinomyces israelii		58.93	0.39	URE 1		
Actinomyces naeslundii		41.07	0.45			
27777050	**Good identification**					**27777050**
Actinomyces israelii		100.0	0.56	MLZ 13		
27777052	**Very good identification**					**27777052**
Actinomyces israelii		100.0	0.59	MLZ 13		
27777060	**Acceptable identification**					**27777060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		90.07	0.49	NAG 6		
Actinomyces naeslundii		9.93	0.42			
27777061	**Acceptable identification**					**27777061**
Actinomyces naeslundii		94.24	0.45	XYL 1		
Actinomyces israelii		5.76	0.15			
27777062	**Acceptable identification**					**27777062**
Actinomyces israelii		93.15	0.52	NAG 6		
Actinomyces naeslundii		6.85	0.42			
27777063	**Acceptable identification**					**27777063**
Actinomyces naeslundii		91.61	0.45	XYL 1		
Actinomyces israelii		8.39	0.18			
27777070	**Good identification**					**27777070**
Actinomyces israelii		100.0	0.35	MLZ 13		
27777072	**Good identification**					**27777072**
Actinomyces israelii		100.0	0.38	MLZ 13		
27777100	**Very good identification**					**27777100**
					GRA	
Actinomyces israelii		98.58	0.87		99	
Actinomyces naeslundii		1.09	0.63		99	
Mitsuokella multacida		0.17	0.27		1	
27777101	**Low discrimination**					**27777101**
					GRA	
Actinomyces naeslundii		61.67	0.66	XYL 1	99	
Actinomyces israelii		37.63	0.53		99	
Mitsuokella multacida		0.70	0.11		1	
27777102	**Very good identification**					**27777102**
Actinomyces israelii		99.05	0.90			
Actinomyces naeslundii		0.73	0.63			
Bifidobacterium adolescentis		0.15	0.35			
27777103	**Low discrimination**					**27777103**
Actinomyces naeslundii		52.22	0.66	XYL 1		
Actinomyces israelii		47.79	0.56			
27777110	**Very good identification**					**27777110**
					GRA	
Actinomyces israelii		99.09	0.73	MLZ 13	99	
Actinomyces naeslundii		0.39	0.41		99	
Mitsuokella multacida		0.38	0.19		1	
27777111	**Low discrimination**					**27777111**
					GRA	
Actinomyces israelii		61.77	0.39	MLZ 13	99	
Actinomyces naeslundii		35.66	0.44	URE 1	99	
Mitsuokella multacida		2.58	0.03		1	
27777112	**Very good identification**					**27777112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		98.30	0.76	MLZ 13		
Bifidobacterium adolescentis		1.19	0.37			
Actinomyces naeslundii		0.26	0.41			
27777113	**Low discrimination**					**27777113**
Actinomyces israelii		72.21	0.42	MLZ 13		
Actinomyces naeslundii		27.79	0.44	URE 1		
27777120	**Low discrimination**					**27777120**
					SPO	
Actinomyces israelii		83.78	0.67	NAG 6	1	
Actinomyces naeslundii		14.49	0.63		1	
Clostridium ramosum		0.67	0.15		92	
27777121	**Very good identification**					**27777121**
Actinomyces naeslundii		96.25	0.66	XYL 1		
Actinomyces israelii		3.75	0.33			
27777122	**Low discrimination**					**27777122**
					SPO	
Actinomyces israelii		87.28	0.70	NAG 6	1	
Actinomyces naeslundii		10.06	0.63		1	
Clostridium ramosum		1.86	0.25		92	
27777123	**Good identification**					**27777123**
Actinomyces naeslundii		94.48	0.66	XYL 1		
Actinomyces israelii		5.52	0.36			
27777130	**Low discrimination**					**27777130**
					SPO	
Actinomyces israelii		89.12	0.53	MLZ 13	1	
Actinomyces naeslundii		5.43	0.41	NAG 6	1	
Clostridium tertium		4.13	0.15		99	
27777131	**Acceptable identification**					**27777131**
Actinomyces naeslundii		90.04	0.44	XYL 1		
Actinomyces israelii		9.96	0.18	MLZ 5		
27777132	**Low discrimination**					**27777132**
					SPO	
Actinomyces israelii		87.41	0.56	MLZ 13	1	
Clostridium tertium		8.11	0.24	NAG 6	99	
Actinomyces naeslundii		3.55	0.41		1	
27777133	**Low discrimination**					**27777133**
Actinomyces naeslundii		85.78	0.44	XYL 1		
Actinomyces israelii		14.22	0.21	MLZ 5		
27777140	**Very good identification**					**27777140**
					GRA	
Actinomyces israelii		99.04	0.80		99	
Actinomyces naeslundii		0.50	0.50		99	
Mitsuokella multacida		0.43	0.27		1	
27777141	**Low discrimination**					**27777141**
					GRA	
Actinomyces israelii		55.81	0.46	URE 1	99	
Actinomyces naeslundii		41.51	0.53		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Mitsuokella multacida		2.68	0.11		1	
27777142	**Very good identification**					**27777142**
Actinomyces israelii		99.58	0.83			
Actinomyces naeslundii		0.33	0.50			
Bifidobacterium adolescentis		0.05	0.20			
27777143	**Low discrimination**					**27777143**
Actinomyces israelii		66.85	0.49	URE 1		
Actinomyces naeslundii		33.15	0.53			
27777150	**Very good identification**					**27777150**
					GRA	
Actinomyces israelii		98.72	0.66	MLZ 13	99	
Mitsuokella multacida		0.96	0.19		1	
Actinomyces naeslundii		0.17	0.28		99	
27777152	**Very good identification**					**27777152**
Actinomyces israelii		99.43	0.69	MLZ 13		
Bifidobacterium adolescentis		0.38	0.21			
Actinomyces naeslundii		0.12	0.28			
27777160	**Good identification**					**27777160**
					GRA	
Actinomyces israelii		91.28	0.60	NAG 6	99	
Actinomyces naeslundii		7.16	0.50		99	
Mitsuokella multacida		1.56	0.17		1	
27777161	**Acceptable identification**					**27777161**
Actinomyces naeslundii		92.10	0.53	XYL 1		
Actinomyces israelii		7.90	0.26			
27777162	**Very good identification**					**27777162**
Actinomyces israelii		95.03	0.63	NAG 6		
Actinomyces naeslundii		4.97	0.50			
27777163	**Low discrimination**					**27777163**
Actinomyces naeslundii		88.60	0.53	XYL 1		
Actinomyces israelii		11.41	0.29			
27777170	**Acceptable identification**					**27777170**
					GRA	
Actinomyces israelii		93.83	0.46	MLZ 13	99	
Mitsuokella multacida		3.58	0.09	NAG 6	1	
Actinomyces naeslundii		2.59	0.28		99	
27777172	**Good identification**					**27777172**
Actinomyces israelii		98.19	0.49	MLZ 13		
Actinomyces naeslundii		1.81	0.28	NAG 6		
27777200	**Very good identification**					**27777200**
Actinomyces israelii		100.0	0.58	RHA 7		
27777202	**Very good identification**					**27777202**
Actinomyces israelii		100.0	0.61	RHA 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27777210	**Good identification**						**27777210**
Actinomyces israelii		100.0	0.44	RHA 7			
27777212	**Good identification**						**27777212**
Actinomyces israelii		100.0	0.47	RHA 7			
27777220	**Good identification**						**27777220**
Actinomyces israelii		100.0	0.37	RHA 7			
27777222	**Good identification**						**27777222**
Actinomyces israelii		100.0	0.40	RHA 7			
27777240	**Good identification**						**27777240**
Actinomyces israelii		100.0	0.51	RHA 7			
27777242	**Good identification**						**27777242**
Actinomyces israelii		100.0	0.54	RHA 7			
27777250	**Good identification**						**27777250**
Actinomyces israelii		100.0	0.36	RHA 7			
27777252	**Good identification**						**27777252**
Actinomyces israelii		100.0	0.39	RHA 7			
27777300	**Very good identification**						**27777300**
					GRA		
Actinomyces israelii		95.78	0.68	RHA 7	99		
Mitsuokella multacida		4.01	0.32		1		
Actinomyces naeslundii		0.14	0.29		99		
27777302	**Very good identification**						**27777302**
					GRA		
Actinomyces israelii		99.59	0.71	RHA 7	99		
Mitsuokella multacida		0.31	0.15		1		
Actinomyces naeslundii		0.10	0.29		99		
27777303	**Low discrimination**						**27777303**
Actinomyces israelii		87.21	0.37	RHA 7			
Actinomyces naeslundii		12.79	0.32	URE 1			
27777310	**Acceptable identification**						**27777310**
					GRA		
Actinomyces israelii		91.45	0.54	RHA 7	99		
Mitsuokella multacida		8.55	0.24	MLZ 13	1		
27777312	**Very good identification**						**27777312**
					GRA		
Actinomyces israelii		99.31	0.57	RHA 7	99		
Mitsuokella multacida		0.69	0.07	MLZ 13	1		
27777320	**Low discrimination**						**27777320**
					SPO	GRA	
Actinomyces israelii		75.31	0.47	RHA 7	1	99	
Mitsuokella multacida		12.36	0.21	NAG 6	1	1	
Clostridium ramosum		10.59	0.17		92	50	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
27777322	**Low discrimination**						**27777322**
					SPO	GRA	
Actinomyces israelii		69.78	0.50	RHA 7	1	99	
Clostridium ramosum		26.16	0.27	NAG 6	92	50	
Parabacteroides distasonis		2.14	0.21		1	1	
27777332	**Acceptable identification**						**27777332**
					GRA		
Actinomyces israelii		90.04	0.36	RHA 7	99		
Parabacteroides distasonis		9.96	0.16	MLZ 13	1		
27777340	**Good identification**						**27777340**
					GRA		
Actinomyces israelii		90.27	0.61	RHA 7	99		
Mitsuokella multacida		9.73	0.32		1		
27777342	**Very good identification**						**27777342**
					GRA		
Actinomyces israelii		99.21	0.64	RHA 7	99		
Mitsuokella multacida		0.79	0.15		1		
27777350	**Low discrimination**						**27777350**
					COC	GRA	
Actinomyces israelii		79.38	0.47	RHA 7	1	99	
Mitsuokella multacida		19.08	0.24	MLZ 13	1	1	
Blautia producta		1.54	-.03		99	99	
27777352	**Good identification**						**27777352**
					GRA		
Actinomyces israelii		98.25	0.50	RHA 7	99		
Mitsuokella multacida		1.75	0.07	MLZ 13	1		
27777360	**Low discrimination**						**27777360**
					GRA		
Actinomyces israelii		70.32	0.40	RHA 7	99		
Mitsuokella multacida		29.68	0.21	NAG 6	1		
27777362	**Good identification**						**27777362**
					GRA		
Actinomyces israelii		96.97	0.43	RHA 7	99		
Mitsuokella multacida		3.03	0.05	NAG 6	1		
27777400	**Very good identification**						**27777400**
					GRA	SPO	
Actinomyces israelii		98.88	0.75		99	1	
Mitsuokella multacida		0.82	0.27		1	1	
Clostridium butyricum		0.30	0.11		99	99	
27777401	**Low discrimination**						**27777401**
					GRA		
Actinomyces israelii		89.30	0.41	URE 1	99		
Mitsuokella multacida		8.16	0.11		1		
Actinomyces naeslundii		2.54	0.24		99		
27777402	**Very good identification**						**27777402**
					GRA	SPO	
Actinomyces israelii		99.81	0.78		99	1	
Clostridium butyricum		0.08	0.04		99	99	
Mitsuokella multacida		0.06	0.11		1	1	
27777403	**Good identification**						**27777403**
Actinomyces israelii		98.14	0.44	URE 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		1.86	0.24				
27777410	**Very good identification**						**27777410**
					GRA		
Actinomyces israelii		98.18	0.61	MLZ 13	99		
Mitsuokella multacida		1.82	0.19		1		
27777412	**Very good identification**						**27777412**
Actinomyces israelii		99.28	0.64	MLZ 13			
Bifidobacterium adolescentis		0.40	0.17				
Bifidobacterium dentium		0.19	-.07				
27777420	**Good identification**						**27777420**
					GRA	SPO	
Actinomyces israelii		96.47	0.55	NAG 6	99	1	
Mitsuokella multacida		3.14	0.17		1	1	
Clostridium butyricum		0.39	-.07		99	99	
27777422	**Very good identification**						**27777422**
Actinomyces israelii		100.0	0.58	NAG 6			
27777430	**Acceptable identification**						**27777430**
					GRA		
Actinomyces israelii		93.24	0.41	MLZ 13	99		
Mitsuokella multacida		6.76	0.09	NAG 6	1		
27777432	**Good identification**						**27777432**
Actinomyces israelii		95.03	0.44	MLZ 13			
Bifidobacterium dentium		4.97	-.03	NAG 6			
27777440	**Very good identification**						**27777440**
					GRA	SPO	
Actinomyces israelii		97.85	0.68		99	1	
Mitsuokella multacida		2.09	0.27		1	1	
Clostridium butyricum		0.07	-.07		99	99	
27777442	**Very good identification**						**27777442**
					GRA		
Actinomyces israelii		99.84	0.71		99		
Mitsuokella multacida		0.16	0.11		1		
27777443	**Good identification**						**27777443**
Actinomyces israelii		100.0	0.37	URE 1			
27777450	**Acceptable identification**						**27777450**
					COC	GRA	
Actinomyces israelii		94.81	0.54	MLZ 13	1	99	
Mitsuokella multacida		4.52	0.19		1	1	
Blautia producta		0.68	-.03		99	99	
27777452	**Very good identification**						**27777452**
					GRA		
Actinomyces israelii		99.65	0.57	MLZ 13	99		
Mitsuokella multacida		0.35	0.03		1		
27777460	**Acceptable identification**						**27777460**
					GRA		
Actinomyces israelii		92.28	0.48	NAG 6	99		
Mitsuokella multacida		7.72	0.17		1		
27777462	**Good identification**						**27777462**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.51	NAG 6			
27777472	**Good identification**						**27777472**
Actinomyces israelii		100.0	0.37	MLZ 13			
27777500	**Low discrimination**						**27777500**
					GRA	SPO	
Actinomyces israelii		81.71	0.86		99	1	
Mitsuokella multacida		16.79	0.61		1	1	
Clostridium butyricum		1.47	0.35		99	99	
27777501	**Low discrimination**						**27777501**
					GRA		
Mitsuokella multacida		68.53	0.45	ESL 99	1		
Actinomyces israelii		30.31	0.52	URE 10	99		
Actinomyces naeslundii		0.61	0.32		99		
27777502	**Very good identification**						**27777502**
					GRA		
Actinomyces israelii		96.89	0.89		99		
Mitsuokella multacida		1.48	0.45		1		
Bifidobacterium adolescentis		1.17	0.50		99		
27777503	**Low discrimination**						**27777503**
					GRA		
Actinomyces israelii		83.49	0.55	URE 1	99		
Mitsuokella multacida		13.98	0.29		1		
Actinomyces naeslundii		1.13	0.32		99		
27777510	**Low discrimination**						**27777510**
					GRA		
Actinomyces israelii		67.85	0.72	MLZ 13	99		
Mitsuokella multacida		31.11	0.53		1		
Bifidobacterium dentium		0.47	0.10		99		
27777511	**Low discrimination**						**27777511**
					GRA		
Mitsuokella multacida		83.45	0.37	ESL 99	1		
Actinomyces israelii		16.55	0.37	URE 10	99		
27777512	**Low discrimination**						**27777512**
Actinomyces israelii		84.64	0.75	MLZ 13			
Bifidobacterium adolescentis		8.37	0.51				
Bifidobacterium dentium		3.94	0.27				
27777513	**Low discrimination**						**27777513**
					GRA		
Actinomyces israelii		65.83	0.40	MLZ 13	99		
Mitsuokella multacida		24.59	0.20	URE 1	1		
Bifidobacterium adolescentis		6.51	0.17		99		
27777520	**Low discrimination**						**27777520**
					GRA	SPO	
Actinomyces israelii		54.60	0.65	NAG 6	99	1	
Mitsuokella multacida		43.95	0.51		1	1	
Clostridium butyricum		1.33	0.17		99	99	
27777522	**Good identification**						**27777522**
					GRA	SPO	
Actinomyces israelii		93.35	0.68	NAG 6	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		5.57	0.35		1	1	
Clostridium butyricum		0.59	0.10		99	99	
27777530	**Low discrimination**						**27777530**
					GRA		
Mitsuokella multacida		59.93	0.43	ESL 99	1		
Actinomyces israelii		33.37	0.51		99		
Bifidobacterium dentium		6.41	0.15		99		
27777540	**Low discrimination**						**27777540**
					COC	GRA	
Actinomyces israelii		64.88	0.79		1	99	
Mitsuokella multacida		34.29	0.61		1	1	
Blautia producta		0.57	0.24		99	99	
27777541	**Low discrimination**						**27777541**
					GRA		
Mitsuokella multacida		85.32	0.45	ESL 99	1		
Actinomyces israelii		14.68	0.44	URE 10	99		
27777542	**Very good identification**						**27777542**
					GRA		
Actinomyces israelii		95.72	0.82		99		
Mitsuokella multacida		3.75	0.45		1		
Bifidobacterium adolescentis		0.37	0.34		99		
27777543	**Low discrimination**						**27777543**
					GRA		
Actinomyces israelii		69.90	0.48	URE 1	99		
Mitsuokella multacida		30.10	0.29		1		
27777550	**Low discrimination**						**27777550**
					COC	GRA	
Mitsuokella multacida		50.01	0.53	ESL 99	1	1	
Actinomyces israelii		42.42	0.65		1	99	
Blautia producta		7.50	0.32		99	99	
27777551	**Acceptable identification**						**27777551**
					COC	GRA	
Mitsuokella multacida		91.68	0.37	ESL 99	1	1	
Actinomyces israelii		7.07	0.30	URE 10	1	99	
Blautia producta		1.25	-.03		99	99	
27777552	**Low discrimination**						**27777552**
					GRA		
Actinomyces israelii		88.26	0.68	MLZ 13	99		
Mitsuokella multacida		7.71	0.37		1		
Bifidobacterium adolescentis		2.78	0.36		99		
27777560	**Low discrimination**						**27777560**
					COC	GRA	
Mitsuokella multacida		66.97	0.51	ESL 99	1	1	
Actinomyces israelii		32.36	0.58		1	99	
Blautia producta		0.50	0.07		99	99	
27777562	**Low discrimination**						**27777562**
					GRA		
Actinomyces israelii		86.71	0.61	NAG 6	99		
Mitsuokella multacida		13.29	0.35		1		
27777570	**Low discrimination**						**27777570**
					COC	GRA	
Mitsuokella multacida		77.93	0.43	ESL 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		16.88	0.44		1	99	
Blautia producta		5.20	0.15		99	99	
27777572	**Low discrimination**						**27777572**
					GRA		
Actinomyces israelii		71.51	0.47	MLZ 13	99		
Mitsuokella multacida		24.46	0.26	NAG 6	1		
Bifidobacterium dentium		2.41	-.03		99		
27777600	**Low discrimination**						**27777600**
					GRA		
Actinomyces israelii		83.00	0.56	RHA 7	99		
Mitsuokella multacida		17.01	0.32		1		
27777602	**Very good identification**						**27777602**
					GRA		
Actinomyces israelii		98.51	0.59	RHA 7	99		
Mitsuokella multacida		1.50	0.15		1		
27777610	**Low discrimination**						**27777610**
					GRA		
Actinomyces israelii		68.63	0.42	RHA 7	99		
Mitsuokella multacida		31.37	0.24	MLZ 13	1		
27777612	**Acceptable identification**						**27777612**
					GRA		
Actinomyces israelii		92.01	0.45	RHA 7	99		
Clostridium clostridioforme		4.88	0.14	MLZ 13	50		
Mitsuokella multacida		3.12	0.07		1		
27777620	**Low discrimination**						**27777620**
					GRA		
Actinomyces israelii		55.48	0.36	RHA 7	99		
Mitsuokella multacida		44.52	0.21	NAG 6	1		
27777622	**Acceptable identification**						**27777622**
					GRA		
Actinomyces israelii		94.39	0.39	RHA 7	99		
Mitsuokella multacida		5.61	0.05	NAG 6	1		
27777640	**Low discrimination**						**27777640**
					GRA		
Actinomyces israelii		65.49	0.49	RHA 7	99		
Mitsuokella multacida		34.51	0.32		1		
27777642	**Good identification**						**27777642**
					GRA		
Actinomyces israelii		96.24	0.52	RHA 7	99		
Mitsuokella multacida		3.76	0.15		1		
27777652	**Acceptable identification**						**27777652**
					GRA		
Actinomyces israelii		91.99	0.38	RHA 7	99		
Mitsuokella multacida		8.01	0.07	MLZ 13	1		
27777700	**Low discrimination**						**27777700**
					GRA	SPO	
Mitsuokella multacida		83.43	0.66	ESL 99	1	1	
Actinomyces israelii		16.45	0.66		99	1	
Clostridium butyricum		0.12	0.09		99	99	
27777701	**Good identification**						**27777701**
					GRA		
Mitsuokella multacida		98.24	0.50	ESL 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.76	0.32	URE 10	99		
27777702	**Low discrimination**						**27777702**
					GRA	SPO	
Actinomyces israelii		72.43	0.69	RHA 7	99	1	
Mitsuokella multacida		27.21	0.50		1	1	
Clostridium butyricum		0.14	0.02		99	99	
27777710	**Good identification**						**27777710**
					COC	GRA	
Mitsuokella multacida		91.81	0.58	ESL 99	1	1	
Actinomyces israelii		8.12	0.52		1	99	
Blautia producta		0.08	-.03		99	99	
27777711	**Good identification**						**27777711**
					GRA		
Mitsuokella multacida		99.20	0.41	ESL 99	1		
Actinomyces israelii		0.80	0.18	URE 10	99		
27777712	**Low discrimination**						**27777712**
					GRA		
Actinomyces israelii		52.58	0.55	RHA 7	99		
Mitsuokella multacida		44.06	0.41	MLZ 13	1		
Clostridium clostridioforme		2.33	0.23		50		
27777720	**Good identification**						**27777720**
					GRA		
Mitsuokella multacida		95.21	0.56	ESL 99	1		
Actinomyces israelii		4.79	0.46		99		
27777721	**Good identification**						**27777721**
Mitsuokella multacida		100.0	0.39	ESL 99			
27777722	**Low discrimination**						**27777722**
					GRA		
Mitsuokella multacida		55.37	0.39	ESL 99	1		
Actinomyces israelii		37.63	0.49	BGL 10	99		
Bacteroides ovatus		5.28	0.15		1		
27777730	**Good identification**						**27777730**
					GRA		
Mitsuokella multacida		97.79	0.47	ESL 99	1		
Actinomyces israelii		2.21	0.32		99		
27777740	**Good identification**						**27777740**
					COC	GRA	
Mitsuokella multacida		92.11	0.66	ESL 99	1	1	
Actinomyces israelii		7.06	0.59		1	99	
Blautia producta		0.83	0.24		99	99	
27777741	**Good identification**						**27777741**
					GRA		
Mitsuokella multacida		99.31	0.50	ESL 99	1		
Actinomyces israelii		0.69	0.25	URE 10	99		
27777742	**Low discrimination**						**27777742**
					COC	GRA	
Actinomyces israelii		50.64	0.62	RHA 7	1	99	
Mitsuokella multacida		48.92	0.50		1	1	
Blautia producta		0.44	0.07		99	99	
27777750	**Low discrimination**						**27777750**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Mitsuokella multacida		89.67	0.58	ESL 99	1	1		
Blautia producta		7.24	0.32		99	99		
Actinomyces israelii		3.08	0.45		1	99		
27777751	**Good identification**							**27777751**
					COC	GRA		
Mitsuokella multacida		99.27	0.41	ESL 99	1	1		
Blautia producta		0.73	-.03	URE 10	99	99		
27777752	**Low discrimination**							**27777752**
					COC	GRA		
Mitsuokella multacida		64.73	0.41	ESL 99	1	1		
Actinomyces israelii		30.04	0.48	BGL 10	1	99		
Blautia producta		5.23	0.15		99	99		
27777760	**Good identification**							**27777760**
					COC	GRA		
Mitsuokella multacida		97.70	0.56	ESL 99	1	1		
Actinomyces israelii		1.91	0.39		1	99		
Blautia producta		0.39	0.07		99	99		
27777761	**Good identification**							**27777761**
Mitsuokella multacida		100.0	0.39	ESL 99				
27777762	**Low discrimination**							**27777762**
					GRA			
Mitsuokella multacida		79.09	0.39	ESL 99	1			
Actinomyces israelii		20.91	0.42	BGL 10	99			
27777770	**Good identification**							**27777770**
					COC	GRA		
Mitsuokella multacida		95.72	0.47	ESL 99	1	1		
Blautia producta		3.44	0.15		99	99		
Actinomyces israelii		0.84	0.25		1	99		
30000000	**Low discrimination**							**30000000**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.98	0.74	NIT 3	99	99		
Veillonella parvula		12.66	0.66		99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.66		1	1		
30000001	**Low discrimination**							**30000001**
					COC	GRA		
Campylobacter ureolyticus		75.99	0.60	IND 1	1	1		
Peptoniphilus asaccharolyticus		7.67	0.40		99	99		
Anaerococcus prevotii		3.23	0.41		99	99		
30000002	**Low discrimination**							**30000002**
					COC	GRA		
Campylobacter gracilis		43.38	0.50	IND 1	1	1		
Porphyromonas asaccharolytica		21.47	0.42	BGL 10	1	1		
Peptoniphilus asaccharolyticus		16.10	0.40		99	99		
30000010	**Low discrimination**							**30000010**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.99	0.40	NIT 3	99	99		
Veillonella parvula		12.67	0.32	MLZ 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30000020	**Low discrimination**							**30000020**
					COC	GRA	SPO	
Campylobacter gracilis		67.81	0.56	IND 1	1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptoniphilus asaccharolyticus		11.19	0.40		99	99	1	
Clostridium cadaveris		4.89	0.44		1	99	99	
30000022	**Low discrimination**							**30000022**
Campylobacter gracilis		78.50	0.39	IND 1				
Pseudoflavonifractor capillosus		16.22	0.32	BGL 10				
Porphyromonas asaccharolytica		1.57	0.08					
30000040	**Low discrimination**							**30000040**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.94	0.40	NIT 3	99	99		
Veillonella parvula		12.65	0.32	SOR 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.39	0.32		1	1		
30000100	**Low discrimination**							**30000100**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.99	0.40	NIT 3	99	99		
Veillonella parvula		12.67	0.32	LAC 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30000200	**Low discrimination**							**30000200**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.99	0.40	NIT 3	99	99		
Veillonella parvula		12.67	0.32	RHA 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30000400	**Low discrimination**							**30000400**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.99	0.40	NIT 3	99	99		
Veillonella parvula		12.67	0.32	ARA 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30001000	**Low discrimination**							**30001000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		85.63	0.66	NIT 1	1	1		
Peptoniphilus asaccharolyticus		5.78	0.40		99	99		
Veillonella parvula		1.93	0.32		99	1		
30002000	**Low discrimination**							**30002000**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.99	0.40	NIT 3	99	99		
Veillonella parvula		12.67	0.32	MAN 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30004000	**Low discrimination**							**30004000**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.98	0.40	NIT 3	99	99		
Veillonella parvula		12.66	0.32	XYL 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30010000	**Low discrimination**							**30010000**
					COC	GRA		
Fusobacterium varium		36.71	0.50	NIT 1	1	1		
Peptoniphilus asaccharolyticus		23.68	0.40		99	99		
Veillonella parvula		7.89	0.32		99	1		
30010020	**Low discrimination**							**30010020**
					GRA	SPO	CAT	
Propionibacterium acnes		78.61	0.58	GLU 99	99	1	95	
Clostridium cadaveris		17.28	0.44		99	99	1	
Campylobacter gracilis		2.42	0.21		1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
30010060	**Good identification**							**30010060**
					GRA	SPO	CAT	
Propionibacterium acnes		98.86	0.47	GLU 99	99	1	95	
Clostridium cadaveris		1.00	0.10		99	99	1	
Campylobacter gracilis		0.14	-.13		1	1	1	
30011020	**Very good identification**							**30011020**
					GRA	SPO	CAT	
Propionibacterium acnes		99.61	0.58	GLU 99	99	1	95	
Clostridium cadaveris		0.22	0.10		99	99	1	
Fusobacterium necrophorum subsp. necrophorum		0.08	-.03		1	1	1	
30011060	**Good identification**							**30011060**
Propionibacterium acnes		100.0	0.47	GLU 99				
30012020	**Good identification**							**30012020**
					GRA	SPO	CAT	
Propionibacterium acnes		99.55	0.54	GLU 99	99	1	95	
Clostridium cadaveris		0.39	0.10		99	99	1	
Campylobacter gracilis		0.06	-.13		1	1	1	
30012060	**Good identification**							**30012060**
Propionibacterium acnes		100.0	0.43	GLU 99				
30013020	**Good identification**							**30013020**
Propionibacterium acnes		100.0	0.54	GLU 99				
30013060	**Good identification**							**30013060**
Propionibacterium acnes		100.0	0.43	GLU 99				
30020000	**Low discrimination**							**30020000**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.98	0.40	NIT 3	99	99		
Veillonella parvula		12.66	0.32	TRE 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30040000	**Low discrimination**							**30040000**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.99	0.40	NIT 3	99	99		
Veillonella parvula		12.67	0.32	CEL 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
30100000	**Low discrimination**							**30100000**
					COC	GRA		
Peptoniphilus asaccharolyticus		34.60	0.40	NIT 3	99	99		
Veillonella parvula		11.53	0.32	MLT 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		11.30	0.32		1	1		
30100001	**Low discrimination**							**30100001**
					COC	GRA	SPO	
Clostridium sordellii		81.70	0.44	NIT 1	1	99	99	
Campylobacter ureolyticus		13.80	0.26	GLU 95	1	1	1	
Peptoniphilus asaccharolyticus		1.39	0.06		99	99	1	
30110001	**Good identification**							**30110001**
					GRA	SPO		
Clostridium sordellii		99.83	0.44	NIT 1	99	99		
Campylobacter ureolyticus		0.17	-.08	GLU 95	1	1		
30200000	**Low discrimination**							**30200000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Peptoniphilus asaccharolyticus		37.88	0.40	NIT 3	99	99		
Veillonella parvula		12.63	0.32	SAL 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.37	0.32		1	1		
30400000	**Low discrimination**							**30400000**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.77	0.40	NIT 3	99	99		
Veillonella parvula		12.59	0.32	RAF 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.34	0.32		1	1		
30510000	**Good identification**							**30510000**
Prevotella intermedia		100.0	0.37	NIT 1				
30511000	**Good identification**							**30511000**
Prevotella intermedia		100.0	0.37	NIT 1				
31000000	**Low discrimination**							**31000000**
					COC	GRA	SPO	
Fusobacterium necrophorum subsp. necrophorum		34.90	0.51	NIT 1	1	1	1	
Peptoniphilus asaccharolyticus		17.28	0.40	GLU 12	99	99	1	
Clostridium bifermentans		16.15	0.52		1	99	99	
31000001	**Low discrimination**							**31000001**
					GRA	SPO		
Clostridium sordellii		87.89	0.50	NIT 1	99	99		
Campylobacter ureolyticus		7.03	0.26	MLT 90	1	1		
Fusobacterium necrophorum subsp. necrophorum		1.43	0.17		1	1		
31000020	**Low discrimination**							**31000020**
					GRA	SPO	CAT	
Clostridium cadaveris		79.34	0.66	NIT 1	99	99	1	
Propionibacterium acnes		19.00	0.58		99	1	95	
Campylobacter gracilis		0.59	0.21		1	1	1	
31000060	**Low discrimination**							**31000060**
					SPO	CAT		
Propionibacterium acnes		83.07	0.47	FRU 99	1	95		
Clostridium cadaveris		15.96	0.32		99	1		
Clostridium bifermentans		0.47	0.08		99	1		
31001000	**Good identification**							**31001000**
					COC	GRA	CAT	
Fusobacterium necrophorum subsp. necrophorum		96.82	0.51	NIT 1	1	1	1	
Propionibacterium acnes		0.96	0.24	GLU 12	1	99	95	
Peptoniphilus asaccharolyticus		0.48	0.06		99	99	15	
31001001	**Acceptable identification**							**31001001**
					GRA	SPO		
Clostridium sordellii		90.67	0.37	NIT 1	99	99		
Fusobacterium necrophorum subsp. necrophorum		8.38	0.17	MLT 90	1	1		
Clostridium novyi biovar A		0.54	0.15		99	99		
31001020	**Good identification**							**31001020**
					GRA	SPO	CAT	
Propionibacterium acnes		94.86	0.58	FRU 99	99	1	95	
Clostridium cadaveris		4.00	0.32		99	99	1	
Fusobacterium necrophorum subsp. necrophorum		0.97	0.17		1	1	1	
31001060	**Good identification**							**31001060**
					SPO	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		99.81	0.47	FRU 99	1	95		
Clostridium cadaveris		0.19	-.03		99	1		
31002020	**Acceptable identification**							**31002020**
					SPO	CAT		
Propionibacterium acnes		92.93	0.54	FRU 99	1	95		
Clostridium cadaveris		6.97	0.32		99	1		
Clostridium bifermentans		0.05	-.03		99	1		
31002060	**Good identification**							**31002060**
					SPO	CAT		
Propionibacterium acnes		99.66	0.43	FRU 99	1	95		
Clostridium cadaveris		0.34	-.03		99	1		
31003020	**Good identification**							**31003020**
					SPO	CAT		
Propionibacterium acnes		99.92	0.54	FRU 99	1	95		
Clostridium cadaveris		0.08	-.03		99	1		
31003060	**Good identification**							**31003060**
Propionibacterium acnes		100.0	0.43	FRU 99				
31010000	**Low discrimination**							**31010000**
					GRA	CAT		
Fusobacterium varium		68.66	0.59	NIT 1	1	1		
Propionibacterium acnes		26.29	0.58		99	95		
Pseudoramibacter alactolyticus		1.83	0.32		99	1		
31010001	**Acceptable identification**							**31010001**
					GRA	SPO	CAT	
Clostridium sordellii		94.20	0.50	NIT 1	99	99	1	
Fusobacterium varium		3.95	0.25	MLT 90	1	1	1	
Propionibacterium acnes		1.51	0.24		99	1	95	
31010020	**Very good identification**							**31010020**
					GRA	SPO	CAT	
Propionibacterium acnes		95.91	0.93		99	1	95	
Clostridium cadaveris		4.05	0.66		99	99	1	
Fusobacterium varium		0.03	0.25		1	1	1	
31010021	**Very good identification**							**31010021**
					SPO	CAT		
Propionibacterium acnes		95.37	0.58	URE 1	1	95		
Clostridium cadaveris		4.02	0.32		99	1		
Clostridium sordellii		0.61	0.15		99	1		
31010022	**Very good identification**							**31010022**
					SPO	CAT		
Propionibacterium acnes		95.63	0.58	BGL 1	1	95		
Clostridium cadaveris		4.03	0.32		99	1		
Atopobium minutum		0.20	0.20		1	1		
31010030	**Very good identification**							**31010030**
					SPO	CAT		
Propionibacterium acnes		95.95	0.58	MLZ 1	1	95		
Clostridium cadaveris		4.05	0.32		99	1		
31010040	**Low discrimination**							**31010040**
					GRA	CAT		
Propionibacterium acnes		80.71	0.47	NAG 99	99	95		
Fusobacterium varium		9.70	0.25		1	1		
Pseudoramibacter alactolyticus		6.38	0.21		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
31010060	**Very good identification**						**31010060**
					SPO	CAT	
Propionibacterium acnes		99.79	0.81		1	95	
Clostridium cadaveris		0.19	0.32		99	1	
Pseudoramibacter alactolyticus		0.01	0.09		1	1	
31010061	**Good identification**						**31010061**
					SPO	CAT	
Propionibacterium acnes		99.81	0.47	URE 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
31010062	**Good identification**						**31010062**
					SPO	CAT	
Propionibacterium acnes		99.65	0.47	BGL 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
Pseudoramibacter alactolyticus		0.16	-.08		1	1	
31010070	**Good identification**						**31010070**
					SPO	CAT	
Propionibacterium acnes		99.81	0.47	MLZ 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
31010120	**Very good identification**						**31010120**
					SPO	CAT	
Propionibacterium acnes		95.95	0.58	LAC 1	1	95	
Clostridium cadaveris		4.05	0.32		99	1	
31010160	**Good identification**						**31010160**
					SPO	CAT	
Propionibacterium acnes		99.81	0.47	LAC 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
31010220	**Very good identification**						**31010220**
					SPO	CAT	
Propionibacterium acnes		95.95	0.58	RHA 1	1	95	
Clostridium cadaveris		4.05	0.32		99	1	
31010260	**Good identification**						**31010260**
					SPO	CAT	
Propionibacterium acnes		99.81	0.47	RHA 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
31010420	**Very good identification**						**31010420**
					SPO	CAT	
Propionibacterium acnes		95.95	0.58	ARA 1	1	95	
Clostridium cadaveris		4.05	0.32		99	1	
31010460	**Good identification**						**31010460**
					SPO	CAT	
Propionibacterium acnes		99.81	0.47	ARA 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
31011000	**Very good identification**						**31011000**
					GRA	CAT	
Propionibacterium acnes		95.72	0.58	NAG 99	99	95	
Fusobacterium varium		2.53	0.25		1	1	
Fusobacterium necrophorum subsp. necrophorum		0.98	0.17		1	1	
31011001	**Acceptable identification**						**31011001**
					SPO	CAT	
Clostridium sordellii		91.67	0.37	NIT 1	99	1	
Propionibacterium acnes		8.33	0.24	MLT 90	1	95	
31011020	**Excellent identification**						**31011020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO	CAT	
Propionibacterium acnes		99.96	0.93		1	95	
Clostridium cadaveris		0.04	0.32		99	1	
Atopobium minutum		0.00	0.15		1	1	
31011021	**Very good identification**						**31011021**
					SPO	CAT	
Propionibacterium acnes		99.85	0.58	URE 1	1	95	
Clostridium sordellii		0.11	0.03		99	1	
Clostridium cadaveris		0.04	-.03		99	1	
31011022	**Very good identification**						**31011022**
					SPO	CAT	
Propionibacterium acnes		99.89	0.58	BGL 1	1	95	
Atopobium minutum		0.07	0.12		1	1	
Clostridium cadaveris		0.04	-.03		99	1	
31011030	**Very good identification**						**31011030**
					SPO	CAT	
Propionibacterium acnes		99.96	0.58	MLZ 1	1	95	
Clostridium cadaveris		0.04	-.03		99	1	
31011040	**Good identification**						**31011040**
Propionibacterium acnes		100.0	0.47	NAG 99			
31011060	**Very good identification**						**31011060**
					SPO	CAT	
Propionibacterium acnes		100.0	0.81		1	95	
Clostridium cadaveris		0.00	-.03		99	1	
31011061	**Good identification**						**31011061**
Propionibacterium acnes		100.0	0.47	URE 1			
31011062	**Good identification**						**31011062**
Propionibacterium acnes		100.0	0.47	BGL 1			
31011070	**Good identification**						**31011070**
Propionibacterium acnes		100.0	0.47	MLZ 1			
31011120	**Very good identification**						**31011120**
					SPO	CAT	
Propionibacterium acnes		99.96	0.58	LAC 1	1	95	
Clostridium cadaveris		0.04	-.03		99	1	
31011160	**Good identification**						**31011160**
Propionibacterium acnes		100.0	0.47	LAC 1			
31011220	**Very good identification**						**31011220**
					SPO	CAT	
Propionibacterium acnes		99.96	0.58	RHA 1	1	95	
Clostridium cadaveris		0.04	-.03		99	1	
31011260	**Good identification**						**31011260**
Propionibacterium acnes		100.0	0.47	RHA 1			
31011420	**Very good identification**						**31011420**
					SPO	CAT	
Propionibacterium acnes		99.96	0.58	ARA 1	1	95	
Clostridium cadaveris		0.04	-.03		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
31011460	**Good identification**							**31011460**
Propionibacterium acnes		100.0	0.47	ARA 1				
31012000	**Low discrimination**							**31012000**
					GRA	CAT		
Propionibacterium acnes		85.32	0.54	NAG 99	99	95		
Pseudoramibacter alactolyticus		5.67	0.27		99	1		
Fusobacterium varium		4.00	0.25		1	1		
31012020	**Very good identification**							**31012020**
					SPO	CAT		
Propionibacterium acnes		99.91	0.88		1	95		
Clostridium cadaveris		0.08	0.32		99	1		
Pseudoramibacter alactolyticus		0.01	0.14		1	1		
31012021	**Good identification**							**31012021**
					SPO	CAT		
Propionibacterium acnes		99.92	0.54	URE 1	1	95		
Clostridium cadaveris		0.08	-.03		99	1		
31012022	**Good identification**							**31012022**
					SPO	CAT		
Propionibacterium acnes		99.79	0.54	BGL 1	1	95		
Pseudoramibacter alactolyticus		0.13	-.02		1	1		
Clostridium cadaveris		0.08	-.03		99	1		
31012030	**Good identification**							**31012030**
					SPO	CAT		
Propionibacterium acnes		99.92	0.54	MLZ 1	1	95		
Clostridium cadaveris		0.08	-.03		99	1		
31012040	**Low discrimination**							**31012040**
					COC	CAT	GRA	
Propionibacterium acnes		89.67	0.43	NAG 99	1	95	99	
Pseudoramibacter alactolyticus		6.79	0.17		1	1	99	
Megasphaera elsdenii		3.54	0.23		99	1	1	
31012060	**Very good identification**							**31012060**
					SPO	CAT		
Propionibacterium acnes		99.98	0.77		1	95		
Pseudoramibacter alactolyticus		0.01	0.04		1	1		
Clostridium cadaveris		0.00	-.03		99	1		
31012061	**Good identification**							**31012061**
Propionibacterium acnes		100.0	0.43	URE 1				
31012062	**Good identification**							**31012062**
Propionibacterium acnes		100.0	0.43	BGL 1				
31012070	**Good identification**							**31012070**
Propionibacterium acnes		100.0	0.43	MLZ 1				
31012120	**Good identification**							**31012120**
					SPO	CAT		
Propionibacterium acnes		99.92	0.54	LAC 1	1	95		
Clostridium cadaveris		0.08	-.03		99	1		
31012160	**Good identification**							**31012160**
Propionibacterium acnes		100.0	0.43	LAC 1				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
31012220	**Good identification**						**31012220**
					SPO	CAT	
Propionibacterium acnes		99.92	0.54	RHA 1	1	95	
Clostridium cadaveris		0.08	-.03		99	1	
31012260	**Good identification**						**31012260**
Propionibacterium acnes		100.0	0.43	RHA 1			
31012420	**Good identification**						**31012420**
					SPO	CAT	
Propionibacterium acnes		99.92	0.54	ARA 1	1	95	
Clostridium cadaveris		0.08	-.03		99	1	
31012460	**Good identification**						**31012460**
Propionibacterium acnes		100.0	0.43	ARA 1			
31013000	**Good identification**						**31013000**
					CAT		
Propionibacterium acnes		99.93	0.54	NAG 99	95		
Pseudoramibacter alactolyticus		0.07	-.07		1		
31013020	**Very good identification**						**31013020**
					SPO	CAT	
Propionibacterium acnes		100.0	0.88		1	95	
Clostridium cadaveris		0.00	-.03		99	1	
31013021	**Good identification**						**31013021**
Propionibacterium acnes		100.0	0.54	URE 1			
31013022	**Good identification**						**31013022**
Propionibacterium acnes		100.0	0.54	BGL 1			
31013030	**Good identification**						**31013030**
Propionibacterium acnes		100.0	0.54	MLZ 1			
31013040	**Good identification**						**31013040**
Propionibacterium acnes		100.0	0.43	NAG 99			
31013060	**Very good identification**						**31013060**
Propionibacterium acnes		100.0	0.77				
31013061	**Good identification**						**31013061**
Propionibacterium acnes		100.0	0.43	URE 1			
31013062	**Good identification**						**31013062**
Propionibacterium acnes		100.0	0.43	BGL 1			
31013070	**Good identification**						**31013070**
Propionibacterium acnes		100.0	0.43	MLZ 1			
31013120	**Good identification**						**31013120**
Propionibacterium acnes		100.0	0.54	LAC 1			
31013160	**Good identification**						**31013160**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Propionibacterium acnes	100.0	0.43	LAC 1				
31013220	**Good identification**							**31013220**
	Propionibacterium acnes	100.0	0.54	RHA 1				
31013260	**Good identification**							**31013260**
	Propionibacterium acnes	100.0	0.43	RHA 1				
31013420	**Good identification**							**31013420**
	Propionibacterium acnes	100.0	0.54	ARA 1				
31013460	**Good identification**							**31013460**
	Propionibacterium acnes	100.0	0.43	ARA 1				
31014000	**Low discrimination**							**31014000**
					GRA	SPO	CAT	
	Fusobacterium varium	89.42	0.35	NIT 1	1	1	1	
	Propionibacterium acnes	8.30	0.24	XYL 4	99	1	95	
	Clostridium glycolicum	1.35	0.09		99	99	1	
31014020	**Very good identification**							**31014020**
					GRA	SPO	CAT	
	Propionibacterium acnes	95.85	0.58	XYL 1	99	1	95	
	Clostridium cadaveris	4.04	0.32		99	99	1	
	Fusobacterium varium	0.11	0.01		1	1	1	
31014060	**Good identification**							**31014060**
					SPO	CAT		
	Propionibacterium acnes	99.66	0.47	XYL 1	1	95		
	Clostridium cadaveris	0.19	-.03		99	1		
	Clostridium glycolicum	0.14	-.03		99	1		
31015020	**Very good identification**							**31015020**
					SPO	CAT		
	Propionibacterium acnes	99.96	0.58	XYL 1	1	95		
	Clostridium cadaveris	0.04	-.03		99	1		
31015060	**Good identification**							**31015060**
	Propionibacterium acnes	100.0	0.47	XYL 1				
31016020	**Good identification**							**31016020**
					SPO	CAT		
	Propionibacterium acnes	99.92	0.54	XYL 1	1	95		
	Clostridium cadaveris	0.08	-.03		99	1		
31016060	**Good identification**							**31016060**
	Propionibacterium acnes	100.0	0.43	XYL 1				
31017020	**Good identification**							**31017020**
	Propionibacterium acnes	100.0	0.54	XYL 1				
31017060	**Good identification**							**31017060**
	Propionibacterium acnes	100.0	0.43	XYL 1				
31020020	**Good identification**							**31020020**
					SPO	CAT		
	Clostridium cadaveris	95.35	0.44	NIT 1	99	1		
	Propionibacterium acnes	4.38	0.24	TRE 5	1	95		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		0.13	-.03		99	1	
31030020	**Low discrimination**						**31030020**
					SPO	CAT	
Propionibacterium acnes		81.98	0.58	TRE 1	1	95	
Clostridium cadaveris		18.02	0.44		99	1	
31030060	**Good identification**						**31030060**
					SPO	CAT	
Propionibacterium acnes		99.00	0.47	TRE 1	1	95	
Clostridium cadaveris		1.00	0.10		99	1	
31031020	**Very good identification**						**31031020**
					SPO	CAT	
Propionibacterium acnes		99.78	0.58	TRE 1	1	95	
Clostridium cadaveris		0.22	0.10		99	1	
31031060	**Good identification**						**31031060**
Propionibacterium acnes		100.0	0.47	TRE 1			
31032020	**Good identification**						**31032020**
					SPO	CAT	
Propionibacterium acnes		99.61	0.54	TRE 1	1	95	
Clostridium cadaveris		0.39	0.10		99	1	
31032060	**Good identification**						**31032060**
Propionibacterium acnes		100.0	0.43	TRE 1			
31033020	**Good identification**						**31033020**
Propionibacterium acnes		100.0	0.54	TRE 1			
31033060	**Good identification**						**31033060**
Propionibacterium acnes		100.0	0.43	TRE 1			
31050020	**Very good identification**						**31050020**
					SPO	CAT	
Propionibacterium acnes		95.95	0.58	CEL 1	1	95	
Clostridium cadaveris		4.05	0.32		99	1	
31050060	**Good identification**						**31050060**
					SPO	CAT	
Propionibacterium acnes		99.81	0.47	CEL 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
31051020	**Very good identification**						**31051020**
					SPO	CAT	
Propionibacterium acnes		99.96	0.58	CEL 1	1	95	
Clostridium cadaveris		0.04	-.03		99	1	
31051060	**Good identification**						**31051060**
Propionibacterium acnes		100.0	0.47	CEL 1			
31052020	**Good identification**						**31052020**
					SPO	CAT	
Propionibacterium acnes		99.92	0.54	CEL 1	1	95	
Clostridium cadaveris		0.08	-.03		99	1	
31052060	**Good identification**						**31052060**
Propionibacterium acnes		100.0	0.43	CEL 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
31053020	**Good identification**						**31053020**
Propionibacterium acnes		100.0	0.54	CEL 1			
31053060	**Good identification**						**31053060**
Propionibacterium acnes		100.0	0.43	CEL 1			
31100000	**Low discrimination**						**31100000**
					COC	SPO	
Clostridium bifermentans		86.74	0.58	NIT 1	1	99	
Clostridium sordellii		4.53	0.32		1	99	
Peptostreptococcus anaerobius		2.89	0.29		99	1	
31100001	**Very good identification**						**31100001**
Clostridium sordellii		99.68	0.66	NIT 1			
Clostridium bifermentans		0.20	0.24				
Clostridium novyi biovar A		0.11	0.32				
31100020	**Low discrimination**						**31100020**
					SPO	CAT	
Clostridium cadaveris		72.61	0.44	NIT 1	99	1	
Clostridium bifermentans		23.14	0.38	MLT 5	99	1	
Propionibacterium acnes		3.34	0.24		1	95	
31100041	**Good identification**						**31100041**
Clostridium sordellii		98.15	0.42	NIT 1			
Clostridium novyi biovar A		1.66	0.29	SOR 4			
Clostridium bifermentans		0.19	0.00				
31101001	**Good identification**						**31101001**
Clostridium sordellii		99.93	0.53	NIT 1			
Clostridium novyi biovar A		0.07	0.15				
31110000	**Low discrimination**						**31110000**
					GRA	SPO	
Prevotella intermedia		57.75	0.50	NIT 1	1	1	
Clostridium bifermentans		12.75	0.36		99	99	
Clostridium sordellii		12.64	0.32		99	99	
31110001	**Very good identification**						**31110001**
					GRA	SPO	
Clostridium sordellii		99.93	0.66	NIT 1	99	99	
Prevotella intermedia		0.05	0.16		1	1	
Clostridium novyi biovar A		0.01	0.15		99	99	
31110020	**Low discrimination**						**31110020**
					SPO	CAT	
Propionibacterium acnes		81.49	0.58	MLT 1	1	95	
Clostridium cadaveris		17.91	0.44		99	1	
Clostridium bifermentans		0.30	0.16		99	1	
31110041	**Good identification**						**31110041**
Clostridium sordellii		99.81	0.42	NIT 1			
Clostridium novyi biovar A		0.19	0.12	SOR 4			
31110060	**Good identification**						**31110060**
					SPO	CAT	
Propionibacterium acnes		99.00	0.47	MLT 1	1	95	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Clostridium cadaveris	1.00	0.10		99	1		
31111000	**Acceptable identification**							**31111000**
					GRA	SPO	CAT	
	Prevotella intermedia	92.02	0.50	NIT 1	1	1	1	
	Clostridium sordellii	3.55	0.19		99	99	1	
	Propionibacterium acnes	3.55	0.24		99	1	95	
31111001	**Good identification**							**31111001**
					GRA	SPO		
	Clostridium sordellii	99.74	0.53	NIT 1	99	99		
	Prevotella intermedia	0.26	0.16		1	1		
31111020	**Very good identification**							**31111020**
					GRA	SPO	CAT	
	Propionibacterium acnes	99.49	0.58	MLT 1	99	1	95	
	Prevotella intermedia	0.26	0.16		1	1	1	
	Clostridium cadaveris	0.22	0.10		99	99	1	
31111060	**Good identification**							**31111060**
	Propionibacterium acnes	100.0	0.47	MLT 1				
31112020	**Good identification**							**31112020**
					SPO	CAT		
	Propionibacterium acnes	99.61	0.54	MLT 1	1	95		
	Clostridium cadaveris	0.39	0.10		99	1		
31112060	**Good identification**							**31112060**
	Propionibacterium acnes	100.0	0.43	MLT 1				
31113020	**Good identification**							**31113020**
	Propionibacterium acnes	100.0	0.54	MLT 1				
31113060	**Good identification**							**31113060**
	Propionibacterium acnes	100.0	0.43	MLT 1				
31200000	**Low discrimination**							**31200000**
					COC	GRA	SPO	
	Clostridium bifermentans	68.50	0.35	NIT 1	1	99	99	
	Fusobacterium necrophorum subsp. necrophorum	13.46	0.17	SAL 10	1	1	1	
	Peptoniphilus asaccharolyticus	6.66	0.06		99	99	1	
31210020	**Very good identification**							**31210020**
					SPO	CAT		
	Propionibacterium acnes	95.95	0.58	SAL 1	1	95		
	Clostridium cadaveris	4.05	0.32		99	1		
31210060	**Good identification**							**31210060**
					SPO	CAT		
	Propionibacterium acnes	99.81	0.47	SAL 1	1	95		
	Clostridium cadaveris	0.19	-.03		99	1		
31211020	**Very good identification**							**31211020**
					SPO	CAT		
	Propionibacterium acnes	99.96	0.58	SAL 1	1	95		
	Clostridium cadaveris	0.04	-.03		99	1		
31211060	**Good identification**							**31211060**
	Propionibacterium acnes	100.0	0.47	SAL 1				
31212020	**Good identification**							**31212020**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO	CAT	
Propionibacterium acnes		99.92	0.54	SAL 1	1	95	
Clostridium cadaveris		0.08	-.03		99	1	
31212060	**Good identification**						**31212060**
Propionibacterium acnes		100.0	0.43	SAL 1			
31213020	**Good identification**						**31213020**
Propionibacterium acnes		100.0	0.54	SAL 1			
31213060	**Good identification**						**31213060**
Propionibacterium acnes		100.0	0.43	SAL 1			
31300000	**Good identification**						**31300000**
Clostridium bifermentans		98.43	0.42	NIT 1			
Clostridium sordellii		0.47	-.03	SAL 10			
Clostridium novyi biovar A		0.42	0.14				
31410020	**Very good identification**						**31410020**
					SPO	CAT	
Propionibacterium acnes		95.95	0.58	RAF 1	1	95	
Clostridium cadaveris		4.05	0.32		99	1	
31410060	**Good identification**						**31410060**
					SPO	CAT	
Propionibacterium acnes		99.81	0.47	RAF 1	1	95	
Clostridium cadaveris		0.19	-.03		99	1	
31411020	**Very good identification**						**31411020**
					SPO	CAT	
Propionibacterium acnes		99.96	0.58	RAF 1	1	95	
Clostridium cadaveris		0.04	-.03		99	1	
31411060	**Good identification**						**31411060**
Propionibacterium acnes		100.0	0.47	RAF 1			
31412020	**Good identification**						**31412020**
					SPO	CAT	
Propionibacterium acnes		99.92	0.54	RAF 1	1	95	
Clostridium cadaveris		0.08	-.03		99	1	
31412060	**Good identification**						**31412060**
Propionibacterium acnes		100.0	0.43	RAF 1			
31413020	**Good identification**						**31413020**
Propionibacterium acnes		100.0	0.54	RAF 1			
31413060	**Good identification**						**31413060**
Propionibacterium acnes		100.0	0.43	RAF 1			
31510000	**Good identification**						**31510000**
					GRA	SPO	
Prevotella intermedia		99.71	0.53	NIT 1	1	1	
Clostridium bifermentans		0.15	0.02		99	99	
Clostridium sordellii		0.15	-.03		99	99	
31511000	**Good identification**						**31511000**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		99.91	0.53	NIT 1	1			
Lactobacillus fermentum		0.10	0.14		99			
32000000	**Low discrimination**							**32000000**
					COC	GRA		
Peptoniphilus asaccharolyticus		37.99	0.40	NIT 3	99	99		
Veillonella parvula		12.66	0.32	SUC 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		12.41	0.32		1	1		
33000020	**Good identification**							**33000020**
					GRA	SPO	CAT	
Clostridium cadaveris		96.70	0.47	NIT 1	99	99	1	
Propionibacterium acnes		3.11	0.24	SUC 7	99	1	95	
Campylobacter gracilis		0.10	-.13		1	1	1	
33010020	**Low discrimination**							**33010020**
					SPO	CAT		
Propionibacterium acnes		76.06	0.58	SUC 1	1	95		
Clostridium cadaveris		23.91	0.47		99	1		
Actinomyces odontolyticus		0.04	0.08		1	10		
33010060	**Good identification**							**33010060**
					SPO	CAT		
Propionibacterium acnes		98.57	0.47	SUC 1	1	95		
Clostridium cadaveris		1.43	0.13		99	1		
33011020	**Very good identification**							**33011020**
					SPO	CAT		
Propionibacterium acnes		99.65	0.58	SUC 1	1	95		
Clostridium cadaveris		0.32	0.13		99	1		
Actinomyces odontolyticus		0.03	0.05		1	10		
33011060	**Good identification**							**33011060**
Propionibacterium acnes		100.0	0.47	SUC 1				
33012020	**Good identification**							**33012020**
					SPO	CAT		
Propionibacterium acnes		99.44	0.54	SUC 1	1	95		
Clostridium cadaveris		0.56	0.13		99	1		
33012060	**Good identification**							**33012060**
Propionibacterium acnes		100.0	0.43	SUC 1				
33013020	**Good identification**							**33013020**
Propionibacterium acnes		100.0	0.54	SUC 1				
33013060	**Good identification**							**33013060**
Propionibacterium acnes		100.0	0.43	SUC 1				
33100000	**Low discrimination**							**33100000**
					GRA	SPO		
Actinomyces odontolyticus		61.68	0.38	IND 1	99	1		
Clostridium bifermentans		28.00	0.24	FRU 92	99	99		
Prevotella intermedia		3.29	0.11		1	1		
33100200	**Good identification**							**33100200**
Actinomyces odontolyticus		100.0	0.43	IND 1				
33100202	**Good identification**							**33100202**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.39	IND 1			
33101101	**Good identification**						**33101101**
Clostridium chauvoei		100.0	0.40	IND 1			
33101200	**Good identification**						**33101200**
Actinomyces odontolyticus		100.0	0.40	IND 1			
33101202	**Good identification**						**33101202**
Actinomyces odontolyticus		100.0	0.36	IND 1			
33104200	**Good identification**						**33104200**
Actinomyces odontolyticus		100.0	0.38	IND 1			
33110000	**Low discrimination**						**33110000**
					COC	GRA	
Actinomyces odontolyticus		65.49	0.56	IND 1	1	99	
Prevotella intermedia		30.05	0.45		1	1	
Megasphaera elsdenii		3.84	0.32		99	1	
33110002	**Good identification**						**33110002**
					COC	GRA	
Actinomyces odontolyticus		96.01	0.52	IND 1	1	99	
Actinomyces viscosus		2.25	0.18		1	99	
Megasphaera elsdenii		1.02	0.15		99	1	
33110020	**Low discrimination**						**33110020**
					SPO	CAT	
Actinomyces odontolyticus		62.06	0.43	IND 1	1	10	
Clostridium cadaveris		22.46	0.25	NAG 14	99	1	
Propionibacterium acnes		13.72	0.24		1	95	
33110022	**Good identification**						**33110022**
Actinomyces odontolyticus		100.0	0.39	IND 1			
33110100	**Low discrimination**						**33110100**
Actinomyces odontolyticus		89.79	0.47	IND 1			
Actinomyces meyeri		6.08	0.24				
Propionibacterium propionicum		2.82	0.29				
33110102	**Low discrimination**						**33110102**
Actinomyces odontolyticus		82.74	0.44	IND 1			
Actinomyces viscosus		10.90	0.22				
Propionibacterium propionicum		6.36	0.32				
33110200	**Very good identification**						**33110200**
					GRA		
Actinomyces odontolyticus		99.76	0.61	IND 1	99		
Prevotella intermedia		0.24	0.11		1		
33110202	**Very good identification**						**33110202**
Actinomyces odontolyticus		100.0	0.57	IND 1			
33110220	**Good identification**						**33110220**
Actinomyces odontolyticus		100.0	0.47	IND 1			
33110222	**Good identification**						**33110222**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		100.0	0.44	IND 1			
33110300	**Good identification**						**33110300**
Actinomyces odontolyticus		100.0	0.52	IND 1			
33110302	**Good identification**						**33110302**
Actinomyces odontolyticus		100.0	0.49	IND 1			
33110320	**Good identification**						**33110320**
Actinomyces odontolyticus		100.0	0.39	IND 1			
33110322	**Good identification**						**33110322**
Actinomyces odontolyticus		100.0	0.35	IND 1			
33111000	**Low discrimination**						**33111000**
					GRA		
Actinomyces odontolyticus		58.04	0.53	IND 1	99		
Prevotella intermedia		39.95	0.45		1		
Eubacterium saburreum		0.83	0.25		85		
33111002	**Good identification**						**33111002**
Actinomyces odontolyticus		95.21	0.49	IND 1			
Actinomyces viscosus		2.05	0.14				
Lactobacillus fermentum		1.27	0.24				
33111020	**Low discrimination**						**33111020**
					GRA	CAT	
Actinomyces odontolyticus		69.80	0.40	IND 1	99	10	
Propionibacterium acnes		23.14	0.24	NAG 14	99	95	
Prevotella intermedia		2.98	0.11		1	1	
33111022	**Good identification**						**33111022**
Actinomyces odontolyticus		97.34	0.36	IND 1			
Lactobacillus fermentum		2.66	0.16	NAG 14			
33111100	**Low discrimination**						**33111100**
Actinomyces odontolyticus		71.47	0.44	IND 1			
Actinomyces meyeri		10.88	0.27				
Eubacterium saburreum		7.59	0.32				
33111101	**Good identification**						**33111101**
					SPO		
Clostridium chauvoei		99.48	0.47	IND 1	99		
Actinomyces odontolyticus		0.35	0.10	MNS 90	1		
Propionibacterium propionicum		0.18	0.12		1		
33111102	**Low discrimination**						**33111102**
Actinomyces odontolyticus		74.38	0.41	IND 1			
Actinomyces viscosus		9.01	0.18				
Propionibacterium propionicum		8.58	0.32				
33111121	**Good identification**						**33111121**
Clostridium chauvoei		100.0	0.38	IND 1			
33111200	**Very good identification**						**33111200**

Code Taxon		ID score	T index	tests against	Differentiation tests GRA	Code
Actinomyces odontolyticus		99.64	0.58	IND 1	99	
Prevotella intermedia		0.36	0.11		1	
33111202	**Good identification**					**33111202**
Actinomyces odontolyticus		100.0	0.54	IND 1		
33111220	**Good identification**					**33111220**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33111222	**Good identification**					**33111222**
Actinomyces odontolyticus		100.0	0.41	IND 1		
33111300	**Good identification**					**33111300**
Actinomyces odontolyticus		100.0	0.49	IND 1		
33111302	**Good identification**					**33111302**
Actinomyces odontolyticus		100.0	0.46	IND 1		
33111320	**Good identification**					**33111320**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33112100	**Very good identification**					**33112100**
Propionibacterium propionicum		99.68	0.63	IND 1		
Actinomyces odontolyticus		0.32	0.13			
33112101	**Good identification**					**33112101**
Propionibacterium propionicum		100.0	0.47	IND 1		
33112102	**Very good identification**					**33112102**
Propionibacterium propionicum		99.87	0.66	IND 1		
Actinomyces odontolyticus		0.13	0.10			
33112103	**Good identification**					**33112103**
Propionibacterium propionicum		100.0	0.50	IND 1		
33112140	**Very good identification**					**33112140**
Propionibacterium propionicum		100.0	0.63	IND 1		
33112141	**Good identification**					**33112141**
Propionibacterium propionicum		100.0	0.47	IND 1		
33112142	**Very good identification**					**33112142**
Propionibacterium propionicum		100.0	0.66	IND 1		
33112143	**Good identification**					**33112143**
Propionibacterium propionicum		100.0	0.50	IND 1		
33113100	**Very good identification**					**33113100**
Propionibacterium propionicum		99.78	0.63	IND 1		
Actinomyces odontolyticus		0.22	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
33113101	**Acceptable identification**					**33113101**
					SPO	
Propionibacterium propionicum		94.64	0.47	IND 1	1	
Clostridium chauvoei		5.36	0.12	URE 10	99	
33113102	**Very good identification**					**33113102**
Propionibacterium propionicum		99.91	0.66	IND 1		
Actinomyces odontolyticus		0.09	0.07			
33113103	**Good identification**					**33113103**
					SPO	
Propionibacterium propionicum		99.34	0.50	IND 1	1	
Clostridium chauvoei		0.66	-.01	URE 10	99	
33113140	**Very good identification**					**33113140**
Propionibacterium propionicum		100.0	0.63	IND 1		
33113141	**Good identification**					**33113141**
Propionibacterium propionicum		100.0	0.47	IND 1		
33113142	**Very good identification**					**33113142**
Propionibacterium propionicum		100.0	0.66	IND 1		
33113143	**Good identification**					**33113143**
Propionibacterium propionicum		100.0	0.50	IND 1		
33114000	**Good identification**					**33114000**
					GRA	
Actinomyces odontolyticus		97.04	0.51	IND 1	99	
Actinomyces meyeri		2.04	0.18		99	
Prevotella intermedia		0.91	0.11		1	
33114002	**Good identification**					**33114002**
Actinomyces odontolyticus		100.0	0.47	IND 1		
33114020	**Good identification**					**33114020**
Actinomyces odontolyticus		98.27	0.37	IND 1		
Actinomyces meyeri		1.73	0.04	NAG 14		
33114100	**Low discrimination**					**33114100**
Actinomyces odontolyticus		78.94	0.42	IND 1		
Actinomyces meyeri		21.06	0.29			
33114102	**Good identification**					**33114102**
Actinomyces odontolyticus		98.67	0.39	IND 1		
Actinomyces meyeri		1.33	0.03			
33114200	**Good identification**					**33114200**
Actinomyces odontolyticus		100.0	0.56	IND 1		
33114202	**Good identification**					**33114202**
Actinomyces odontolyticus		100.0	0.52	IND 1		
33114220	**Good identification**					**33114220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.42	IND 1		
33114222	**Good identification**					**33114222**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33114300	**Good identification**					**33114300**
Actinomyces odontolyticus		100.0	0.47	IND 1		
33114302	**Good identification**					**33114302**
Actinomyces odontolyticus		100.0	0.43	IND 1		
33115000	**Acceptable identification**					**33115000**
Actinomyces odontolyticus		92.00	0.48	IND 1		
Actinomyces meyeri		4.36	0.21			
Eubacterium saburreum		1.45	0.21			
33115002	**Good identification**					**33115002**
Actinomyces odontolyticus		98.43	0.44	IND 1		
Lactobacillus fermentum		1.57	0.20			
33115100	**Low discrimination**					**33115100**
Actinomyces odontolyticus		56.61	0.39	IND 1		
Actinomyces meyeri		33.97	0.32			
Eubacterium saburreum		6.57	0.27			
33115102	**Low discrimination**					**33115102**
Actinomyces odontolyticus		87.26	0.36	IND 1		
Lactobacillus fermentum		7.18	0.24			
Eubacterium saburreum		2.92	0.14			
33115200	**Good identification**					**33115200**
Actinomyces odontolyticus		100.0	0.53	IND 1		
33115202	**Good identification**					**33115202**
Actinomyces odontolyticus		100.0	0.49	IND 1		
33115220	**Good identification**					**33115220**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33115222	**Good identification**					**33115222**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33115300	**Good identification**					**33115300**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33115302	**Good identification**					**33115302**
Actinomyces odontolyticus		100.0	0.40	IND 1		
33132100	**Very good identification**					**33132100**
Propionibacterium propionicum		100.0	0.61	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
33132101	**Good identification**					**33132101**
Propionibacterium propionicum		100.0	0.45	IND 1		
33132102	**Very good identification**					**33132102**
Propionibacterium propionicum		100.0	0.64	IND 1		
33132103	**Good identification**					**33132103**
Propionibacterium propionicum		100.0	0.48	IND 1		
33132140	**Very good identification**					**33132140**
Propionibacterium propionicum		100.0	0.61	IND 1		
33132141	**Good identification**					**33132141**
Propionibacterium propionicum		100.0	0.45	IND 1		
33132142	**Very good identification**					**33132142**
Propionibacterium propionicum		100.0	0.64	IND 1		
33132143	**Good identification**					**33132143**
Propionibacterium propionicum		100.0	0.48	IND 1		
33133100	**Very good identification**					**33133100**
Propionibacterium propionicum		100.0	0.61	IND 1		
33133101	**Good identification**					**33133101**
Propionibacterium propionicum		100.0	0.45	IND 1		
33133102	**Very good identification**					**33133102**
Propionibacterium propionicum		100.0	0.64	IND 1		
33133103	**Good identification**					**33133103**
Propionibacterium propionicum		100.0	0.48	IND 1		
33133140	**Very good identification**					**33133140**
Propionibacterium propionicum		100.0	0.61	IND 1		
33133141	**Good identification**					**33133141**
Propionibacterium propionicum		100.0	0.45	IND 1		
33133142	**Very good identification**					**33133142**
Propionibacterium propionicum		100.0	0.64	IND 1		
33133143	**Good identification**					**33133143**
Propionibacterium propionicum		100.0	0.48	IND 1		
33300000	**Good identification**					**33300000**
					SPO	
Actinomyces odontolyticus		97.47	0.43	IND 1	1	
Clostridium bifermentans		2.53	0.08	FRU 92	99	
33300002	**Good identification**					**33300002**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		100.0	0.39	IND 1		
33300200	**Good identification**					**33300200**
Actinomyces odontolyticus		100.0	0.48	IND 1		
33300202	**Good identification**					**33300202**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33300300	**Good identification**					**33300300**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33300302	**Good identification**					**33300302**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33301000	**Good identification**					**33301000**
Actinomyces odontolyticus		98.78	0.40	IND 1		
Eubacterium tortuosum		1.22	-.04	FRU 92		
33301002	**Good identification**					**33301002**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33301200	**Good identification**					**33301200**
Actinomyces odontolyticus		100.0	0.45	IND 1		
33301202	**Good identification**					**33301202**
Actinomyces odontolyticus		100.0	0.41	IND 1		
33301300	**Good identification**					**33301300**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33304000	**Good identification**					**33304000**
Actinomyces odontolyticus		100.0	0.38	IND 1		
33304200	**Good identification**					**33304200**
Actinomyces odontolyticus		100.0	0.42	IND 1		
33304202	**Good identification**					**33304202**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33305200	**Good identification**					**33305200**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33305202	**Good identification**					**33305202**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33310000	**Very good identification**					**33310000**
					GRA	
Actinomyces odontolyticus		99.76	0.61	IND 1	99	
Prevotella intermedia		0.24	0.11		1	
33310002	**Very good identification**					**33310002**
Actinomyces odontolyticus		99.68	0.57	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.32	0.08			
33310020	**Good identification**					**33310020**
Actinomyces odontolyticus		100.0	0.47	IND 1		
33310022	**Good identification**					**33310022**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33310100	**Good identification**					**33310100**
Actinomyces odontolyticus		100.0	0.52	IND 1		
33310102	**Good identification**					**33310102**
Actinomyces odontolyticus		98.23	0.49	IND 1		
Actinomyces viscosus		1.77	0.12			
33310120	**Good identification**					**33310120**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33310122	**Good identification**					**33310122**
Actinomyces odontolyticus		100.0	0.35	IND 1		
33310200	**Very good identification**					**33310200**
Actinomyces odontolyticus		100.0	0.66	IND 1		
33310202	**Very good identification**					**33310202**
Actinomyces odontolyticus		100.0	0.62	IND 1		
33310220	**Good identification**					**33310220**
Actinomyces odontolyticus		100.0	0.52	IND 1		
33310222	**Good identification**					**33310222**
Actinomyces odontolyticus		100.0	0.49	IND 1		
33310300	**Very good identification**					**33310300**
Actinomyces odontolyticus		100.0	0.57	IND 1		
33310302	**Good identification**					**33310302**
Actinomyces odontolyticus		100.0	0.54	IND 1		
33310320	**Good identification**					**33310320**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33310322	**Good identification**					**33310322**
Actinomyces odontolyticus		100.0	0.40	IND 1		
33311000	**Very good identification**					**33311000**
					GRA	
Actinomyces odontolyticus		99.64	0.58	IND 1	99	
Prevotella intermedia		0.36	0.11		1	
33311002	**Good identification**					**33311002**
Actinomyces odontolyticus		99.71	0.54	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.29	0.04			
33311020	**Good identification**					**33311020**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33311022	**Good identification**					**33311022**
Actinomyces odontolyticus		100.0	0.41	IND 1		
33311100	**Good identification**					**33311100**
Actinomyces odontolyticus		100.0	0.49	IND 1		
33311102	**Good identification**					**33311102**
Actinomyces odontolyticus		98.37	0.46	IND 1		
Actinomyces viscosus		1.63	0.08			
33311120	**Good identification**					**33311120**
Actinomyces odontolyticus		95.40	0.36	IND 1		
Clostridium perfringens		4.60	0.08	NAG 14		
33311200	**Very good identification**					**33311200**
Actinomyces odontolyticus		100.0	0.63	IND 1		
33311202	**Very good identification**					**33311202**
Actinomyces odontolyticus		100.0	0.59	IND 1		
33311220	**Good identification**					**33311220**
Actinomyces odontolyticus		100.0	0.49	IND 1		
33311222	**Good identification**					**33311222**
Actinomyces odontolyticus		100.0	0.46	IND 1		
33311300	**Good identification**					**33311300**
Actinomyces odontolyticus		100.0	0.54	IND 1		
33311302	**Good identification**					**33311302**
Actinomyces odontolyticus		100.0	0.51	IND 1		
33311320	**Good identification**					**33311320**
Actinomyces odontolyticus		100.0	0.41	IND 1		
33311322	**Good identification**					**33311322**
Actinomyces odontolyticus		100.0	0.37	IND 1		
33314000	**Good identification**					**33314000**
Actinomyces odontolyticus		100.0	0.56	IND 1		
33314002	**Good identification**					**33314002**
Actinomyces odontolyticus		100.0	0.52	IND 1		
33314020	**Good identification**					**33314020**
Actinomyces odontolyticus		100.0	0.42	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
33314022	**Good identification**					**33314022**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33314100	**Good identification**					**33314100**
Actinomyces odontolyticus		100.0	0.47	IND 1		
33314102	**Good identification**					**33314102**
Actinomyces odontolyticus		100.0	0.43	IND 1		
33314200	**Very good identification**					**33314200**
Actinomyces odontolyticus		100.0	0.61	IND 1		
33314202	**Very good identification**					**33314202**
Actinomyces odontolyticus		100.0	0.57	IND 1		
33314220	**Good identification**					**33314220**
Actinomyces odontolyticus		100.0	0.47	IND 1		
33314222	**Good identification**					**33314222**
Actinomyces odontolyticus		100.0	0.43	IND 1		
33314300	**Good identification**					**33314300**
Actinomyces odontolyticus		100.0	0.52	IND 1		
33314302	**Good identification**					**33314302**
Actinomyces odontolyticus		100.0	0.48	IND 1		
33314320	**Good identification**					**33314320**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33315000	**Good identification**					**33315000**
Actinomyces odontolyticus		100.0	0.53	IND 1		
33315002	**Good identification**					**33315002**
Actinomyces odontolyticus		100.0	0.49	IND 1		
33315020	**Good identification**					**33315020**
Actinomyces odontolyticus		100.0	0.39	IND 1		
33315022	**Good identification**					**33315022**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33315100	**Good identification**					**33315100**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33315102	**Good identification**					**33315102**
Actinomyces odontolyticus		100.0	0.40	IND 1		
33315200	**Very good identification**					**33315200**
Actinomyces odontolyticus		100.0	0.58	IND 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
33315202	**Good identification**					**33315202**
Actinomyces odontolyticus		100.0	0.54	IND 1		
33315220	**Good identification**					**33315220**
Actinomyces odontolyticus		100.0	0.44	IND 1		
33315222	**Good identification**					**33315222**
Actinomyces odontolyticus		100.0	0.40	IND 1		
33315300	**Good identification**					**33315300**
Actinomyces odontolyticus		100.0	0.49	IND 1		
33315302	**Good identification**					**33315302**
Actinomyces odontolyticus		100.0	0.45	IND 1		
33315320	**Good identification**					**33315320**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33377100	**Good identification**					**33377100**
Actinomyces israelii		100.0	0.37	IND 1		
33377102	**Good identification**					**33377102**
Actinomyces israelii		100.0	0.40	IND 1		
33377500	**Good identification**					**33377500**
Actinomyces israelii		100.0	0.35	IND 1		
33377502	**Good identification**					**33377502**
Actinomyces israelii		100.0	0.38	IND 1		
33510000	**Good identification**					**33510000**
					GRA	
Prevotella intermedia		95.40	0.48	NIT 1	1	
Actinomyces odontolyticus		4.29	0.30		99	
Actinomyces viscosus		0.31	0.04		99	
33510002	**Low discrimination**					**33510002**
					GRA	
Actinomyces viscosus		89.66	0.38	IND 1	99	
Actinomyces odontolyticus		7.57	0.27	MNS 96	99	
Prevotella intermedia		2.78	0.14		1	
33510102	**Good identification**					**33510102**
Actinomyces viscosus		96.17	0.42	IND 1		
Propionibacterium propionicum		2.39	0.29	MNS 96		
Actinomyces odontolyticus		1.44	0.18			
33510200	**Low discrimination**					**33510200**
					GRA	
Actinomyces odontolyticus		89.62	0.35	IND 1	99	
Prevotella intermedia		10.38	0.14	RAF 3	1	
33511000	**Good identification**					**33511000**
					GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella intermedia		95.02	0.48	NIT 1	1	
Actinomyces odontolyticus		2.85	0.27		99	
Lactobacillus fermentum		1.23	0.28		99	
33511102	**Low discrimination**					**33511102**
Actinomyces viscosus		87.84	0.39	IND 1		
Lactobacillus fermentum		5.21	0.32	MNS 96		
Propionibacterium propionicum		3.56	0.29			
33512100	**Very good identification**					**33512100**
Propionibacterium propionicum		100.0	0.60	IND 1		
33512101	**Good identification**					**33512101**
Propionibacterium propionicum		100.0	0.44	IND 1		
33512102	**Very good identification**					**33512102**
Propionibacterium propionicum		99.59	0.63	IND 1		
Actinomyces viscosus		0.41	0.08			
33512103	**Good identification**					**33512103**
Propionibacterium propionicum		98.78	0.47	IND 1		
Actinomyces viscosus		1.22	0.00	URE 10		
33512140	**Very good identification**					**33512140**
Propionibacterium propionicum		100.0	0.60	IND 1		
33512141	**Good identification**					**33512141**
Propionibacterium propionicum		100.0	0.44	IND 1		
33512142	**Very good identification**					**33512142**
Propionibacterium propionicum		100.0	0.63	IND 1		
33512143	**Good identification**					**33512143**
Propionibacterium propionicum		100.0	0.47	IND 1		
33513100	**Very good identification**					**33513100**
Propionibacterium propionicum		99.44	0.60	IND 1		
Actinomyces naeslundii		0.56	0.32			
33513101	**Acceptable identification**					**33513101**
Propionibacterium propionicum		92.90	0.44	IND 1		
Actinomyces naeslundii		7.10	0.35	URE 10		
33513102	**Very good identification**					**33513102**
Propionibacterium propionicum		99.38	0.63	IND 1		
Actinomyces naeslundii		0.38	0.32			
Actinomyces viscosus		0.25	0.04			
33513103	**Good identification**					**33513103**
Propionibacterium propionicum		95.15	0.47	IND 1		
Actinomyces naeslundii		4.85	0.35	URE 10		
33513140	**Very good identification**					**33513140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.60	IND 1		
33513141	**Good identification**					**33513141**
Propionibacterium propionicum		100.0	0.44	IND 1		
33513142	**Very good identification**					**33513142**
Propionibacterium propionicum		100.0	0.63	IND 1		
33513143	**Good identification**					**33513143**
Propionibacterium propionicum		100.0	0.47	IND 1		
33532100	**Very good identification**					**33532100**
Propionibacterium propionicum		100.0	0.58	IND 1		
33532101	**Good identification**					**33532101**
Propionibacterium propionicum		100.0	0.42	IND 1		
33532102	**Very good identification**					**33532102**
Propionibacterium propionicum		100.0	0.61	IND 1		
33532103	**Good identification**					**33532103**
Propionibacterium propionicum		100.0	0.45	IND 1		
33532140	**Very good identification**					**33532140**
Propionibacterium propionicum		100.0	0.58	IND 1		
33532141	**Good identification**					**33532141**
Propionibacterium propionicum		100.0	0.42	IND 1		
33532142	**Very good identification**					**33532142**
Propionibacterium propionicum		100.0	0.61	IND 1		
33532143	**Good identification**					**33532143**
Propionibacterium propionicum		100.0	0.45	IND 1		
33533100	**Very good identification**					**33533100**
Propionibacterium propionicum		98.68	0.58	IND 1		
Actinomyces naeslundii		1.33	0.37			
33533101	**Low discrimination**					**33533101**
Propionibacterium propionicum		84.65	0.42	IND 1		
Actinomyces naeslundii		15.35	0.40	URE 10		
33533102	**Very good identification**					**33533102**
Propionibacterium propionicum		99.11	0.61	IND 1		
Actinomyces naeslundii		0.89	0.37			
33533103	**Low discrimination**					**33533103**
Propionibacterium propionicum		89.22	0.45	IND 1		
Actinomyces naeslundii		10.78	0.40	URE 10		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
33533120	**Low discrimination**					**33533120**
Actinomyces naeslundii		57.07	0.37	IND 1		
Propionibacterium propionicum		42.93	0.24	MNS 88		
33533121	**Good identification**					**33533121**
Actinomyces naeslundii		100.0	0.40	IND 1		
33533123	**Good identification**					**33533123**
Actinomyces naeslundii		100.0	0.40	IND 1		
33533140	**Very good identification**					**33533140**
Propionibacterium propionicum		99.76	0.58	IND 1		
Actinomyces naeslundii		0.24	0.24			
33533141	**Good identification**					**33533141**
Propionibacterium propionicum		96.90	0.42	IND 1		
Actinomyces naeslundii		3.10	0.27	URE 10		
33533142	**Very good identification**					**33533142**
Propionibacterium propionicum		99.84	0.61	IND 1		
Actinomyces naeslundii		0.16	0.24			
33533143	**Good identification**					**33533143**
Propionibacterium propionicum		97.91	0.45	IND 1		
Actinomyces naeslundii		2.09	0.27	URE 10		
33553101	**Good identification**					**33553101**
Actinomyces naeslundii		100.0	0.37	IND 1		
33553103	**Low discrimination**					**33553103**
Actinomyces naeslundii		86.99	0.37	IND 1		
Propionibacterium propionicum		13.01	0.12	MNS 88		
33553121	**Good identification**					**33553121**
Actinomyces naeslundii		100.0	0.37	IND 1		
33553123	**Good identification**					**33553123**
Actinomyces naeslundii		100.0	0.37	IND 1		
33571101	**Good identification**					**33571101**
Actinomyces naeslundii		100.0	0.37	IND 1		
33571103	**Low discrimination**					**33571103**
Actinomyces naeslundii		81.86	0.37	IND 1		
Actinomyces viscosus		18.14	0.01	MNS 88		
33571121	**Good identification**					**33571121**
Actinomyces naeslundii		100.0	0.37	IND 1		
33571123	**Good identification**					**33571123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	IND 1		
33573100	**Low discrimination**					**33573100**
Actinomyces naeslundii		63.80	0.39	IND 1		
Propionibacterium propionicum		36.20	0.24	MNS 88		
33573101	**Good identification**					**33573101**
Actinomyces naeslundii		100.0	0.42	IND 1		
33573102	**Low discrimination**					**33573102**
Actinomyces naeslundii		54.02	0.39	IND 1		
Propionibacterium propionicum		45.98	0.27	MNS 88		
33573103	**Good identification**					**33573103**
Actinomyces naeslundii		100.0	0.42	IND 1		
33573120	**Good identification**					**33573120**
Actinomyces naeslundii		100.0	0.39	IND 1		
33573121	**Good identification**					**33573121**
Actinomyces naeslundii		100.0	0.42	IND 1		
33573122	**Good identification**					**33573122**
Actinomyces naeslundii		100.0	0.39	IND 1		
33573123	**Good identification**					**33573123**
Actinomyces naeslundii		100.0	0.42	IND 1		
33710000	**Low discrimination**					**33710000**
					GRA	
Actinomyces odontolyticus		89.62	0.35	IND 1	99	
Prevotella intermedia		10.38	0.14	RAF 3	1	
33710200	**Good identification**					**33710200**
Actinomyces odontolyticus		100.0	0.40	IND 1		
33710202	**Good identification**					**33710202**
Actinomyces odontolyticus		100.0	0.36	IND 1		
33711101	**Good identification**					**33711101**
Actinomyces naeslundii		100.0	0.39	IND 1		
33711120	**Low discrimination**					**33711120**
Actinomyces naeslundii		76.69	0.36	IND 1		
Actinomyces odontolyticus		23.31	0.10	MNS 88		
33711121	**Good identification**					**33711121**
Actinomyces naeslundii		100.0	0.39	IND 1		
33711122	**Low discrimination**					**33711122**
Actinomyces naeslundii		84.30	0.36	IND 1		
Actinomyces odontolyticus		15.70	0.06	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
33711123	**Good identification**					**33711123**
Actinomyces naeslundii		100.0	0.39	IND 1		
33711200	**Good identification**					**33711200**
Actinomyces odontolyticus		100.0	0.37	IND 1		
33713001	**Good identification**					**33713001**
Actinomyces naeslundii		100.0	0.36	IND 1		
33713003	**Good identification**					**33713003**
Actinomyces naeslundii		100.0	0.36	IND 1		
33713021	**Good identification**					**33713021**
Actinomyces naeslundii		100.0	0.36	IND 1		
33713023	**Good identification**					**33713023**
Actinomyces naeslundii		100.0	0.36	IND 1		
33713100	**Low discrimination**					**33713100**
Actinomyces naeslundii		66.52	0.41	IND 1		
Propionibacterium propionicum		33.48	0.26	MNS 88		
33713101	**Good identification**					**33713101**
Actinomyces naeslundii		100.0	0.44	IND 1		
33713102	**Low discrimination**					**33713102**
Actinomyces naeslundii		56.98	0.41	IND 1		
Propionibacterium propionicum		43.02	0.29	MNS 88		
33713103	**Acceptable identification**					**33713103**
Actinomyces naeslundii		94.70	0.44	IND 1		
Propionibacterium propionicum		5.30	0.12	MNS 88		
33713120	**Good identification**					**33713120**
Actinomyces naeslundii		100.0	0.41	IND 1		
33713121	**Good identification**					**33713121**
Actinomyces naeslundii		100.0	0.44	IND 1		
33713122	**Good identification**					**33713122**
Actinomyces naeslundii		100.0	0.41	IND 1		
33713123	**Good identification**					**33713123**
Actinomyces naeslundii		100.0	0.44	IND 1		
33731001	**Good identification**					**33731001**
Actinomyces naeslundii		100.0	0.36	IND 1		
33731003	**Good identification**					**33731003**
Actinomyces naeslundii		100.0	0.36	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
33731021	**Good identification**					**33731021**
Actinomyces naeslundii		100.0	0.36	IND 1		
33731023	**Good identification**					**33731023**
Actinomyces naeslundii		100.0	0.36	IND 1		
33731100	**Good identification**					**33731100**
Actinomyces naeslundii		100.0	0.41	IND 1		
33731101	**Good identification**					**33731101**
Actinomyces naeslundii		100.0	0.44	IND 1		
33731102	**Low discrimination**					**33731102**
Actinomyces naeslundii		72.84	0.41	IND 1		
Actinomyces viscosus		27.16	0.10	MNS 88		
33731103	**Acceptable identification**					**33731103**
Actinomyces naeslundii		92.35	0.44	IND 1		
Actinomyces viscosus		7.65	0.01	MNS 88		
33731120	**Acceptable identification**					**33731120**
Actinomyces naeslundii		92.05	0.41	IND 1		
Clostridium perfringens		7.95	0.01	MNS 88		
33731121	**Good identification**					**33731121**
Actinomyces naeslundii		100.0	0.44	IND 1		
33731122	**Good identification**					**33731122**
Actinomyces naeslundii		100.0	0.41	IND 1		
33731123	**Good identification**					**33731123**
Actinomyces naeslundii		100.0	0.44	IND 1		
33733000	**Good identification**					**33733000**
Actinomyces naeslundii		100.0	0.38	IND 1		
33733001	**Good identification**					**33733001**
Actinomyces naeslundii		100.0	0.41	IND 1		
33733002	**Good identification**					**33733002**
Actinomyces naeslundii		100.0	0.38	IND 1		
33733003	**Good identification**					**33733003**
Actinomyces naeslundii		100.0	0.41	IND 1		
33733020	**Good identification**					**33733020**
Actinomyces naeslundii		100.0	0.38	IND 1		
33733021	**Good identification**					**33733021**
Actinomyces naeslundii		100.0	0.41	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
33733022	**Good identification**					**33733022**
Actinomyces naeslundii		100.0	0.38	IND 1		
33733023	**Good identification**					**33733023**
Actinomyces naeslundii		100.0	0.41	IND 1		
33733100	**Low discrimination**					**33733100**
Actinomyces naeslundii		82.50	0.46	IND 1		
Propionibacterium propionicum		17.50	0.24	MNS 88		
33733101	**Good identification**					**33733101**
Actinomyces naeslundii		100.0	0.49	IND 1		
33733102	**Low discrimination**					**33733102**
Actinomyces naeslundii		75.86	0.46	IND 1		
Propionibacterium propionicum		24.14	0.27	MNS 88		
33733103	**Good identification**					**33733103**
Actinomyces naeslundii		100.0	0.49	IND 1		
33733120	**Good identification**					**33733120**
Actinomyces naeslundii		100.0	0.46	IND 1		
33733121	**Good identification**					**33733121**
Actinomyces naeslundii		100.0	0.49	IND 1		
33733122	**Good identification**					**33733122**
Actinomyces naeslundii		100.0	0.46	IND 1		
33733123	**Good identification**					**33733123**
Actinomyces naeslundii		100.0	0.49	IND 1		
33733141	**Good identification**					**33733141**
Actinomyces naeslundii		100.0	0.36	IND 1		
33733143	**Good identification**					**33733143**
Actinomyces naeslundii		100.0	0.36	IND 1		
33733161	**Good identification**					**33733161**
Actinomyces naeslundii		100.0	0.36	IND 1		
33733163	**Good identification**					**33733163**
Actinomyces naeslundii		100.0	0.36	IND 1		
33751100	**Good identification**					**33751100**
Actinomyces naeslundii		100.0	0.38	IND 1		
33751101	**Good identification**					**33751101**
Actinomyces naeslundii		100.0	0.41	IND 1		
33751103	**Low discrimination**					**33751103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		70.01	0.41	IND 1		
Actinomyces viscosus		29.99	0.11	MNS 88		
33751120	**Good identification**					**33751120**
Actinomyces naeslundii		100.0	0.38	IND 1		
33751121	**Good identification**					**33751121**
Actinomyces naeslundii		100.0	0.41	IND 1		
33751122	**Good identification**					**33751122**
Actinomyces naeslundii		100.0	0.38	IND 1		
33751123	**Good identification**					**33751123**
Actinomyces naeslundii		100.0	0.41	IND 1		
33753000	**Good identification**					**33753000**
Actinomyces naeslundii		100.0	0.35	IND 1		
33753001	**Good identification**					**33753001**
Actinomyces naeslundii		100.0	0.38	IND 1		
33753002	**Good identification**					**33753002**
Actinomyces naeslundii		100.0	0.35	IND 1		
33753003	**Good identification**					**33753003**
Actinomyces naeslundii		100.0	0.38	IND 1		
33753020	**Good identification**					**33753020**
Actinomyces naeslundii		100.0	0.35	IND 1		
33753021	**Good identification**					**33753021**
Actinomyces naeslundii		100.0	0.38	IND 1		
33753022	**Good identification**					**33753022**
Actinomyces naeslundii		100.0	0.35	IND 1		
33753023	**Good identification**					**33753023**
Actinomyces naeslundii		100.0	0.38	IND 1		
33753100	**Low discrimination**					**33753100**
Actinomyces naeslundii		89.05	0.43	IND 1		
Actinomyces israelii		10.95	0.18	MNS 88		
33753101	**Good identification**					**33753101**
Actinomyces naeslundii		100.0	0.46	IND 1		
33753102	**Low discrimination**					**33753102**
Actinomyces naeslundii		84.42	0.43	IND 1		
Actinomyces israelii		15.58	0.21	MNS 88		
33753103	**Good identification**					**33753103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	IND 1		
33753120	**Good identification**					**33753120**
Actinomyces naeslundii		100.0	0.43	IND 1		
33753121	**Good identification**					**33753121**
Actinomyces naeslundii		100.0	0.46	IND 1		
33753122	**Good identification**					**33753122**
Actinomyces naeslundii		100.0	0.43	IND 1		
33753123	**Good identification**					**33753123**
Actinomyces naeslundii		100.0	0.46	IND 1		
33757100	**Good identification**					**33757100**
Actinomyces israelii		100.0	0.37	IND 1		
33757102	**Good identification**					**33757102**
Actinomyces israelii		100.0	0.40	IND 1		
33757500	**Good identification**					**33757500**
Actinomyces israelii		100.0	0.35	IND 1		
33757502	**Good identification**					**33757502**
Actinomyces israelii		100.0	0.38	IND 1		
33771000	**Good identification**					**33771000**
Actinomyces naeslundii		100.0	0.35	IND 1		
33771001	**Good identification**					**33771001**
Actinomyces naeslundii		100.0	0.38	IND 1		
33771002	**Good identification**					**33771002**
Actinomyces naeslundii		100.0	0.35	IND 1		
33771003	**Good identification**					**33771003**
Actinomyces naeslundii		100.0	0.38	IND 1		
33771020	**Good identification**					**33771020**
Actinomyces naeslundii		100.0	0.35	IND 1		
33771021	**Good identification**					**33771021**
Actinomyces naeslundii		100.0	0.38	IND 1		
33771022	**Good identification**					**33771022**
Actinomyces naeslundii		100.0	0.35	IND 1		
33771023	**Good identification**					**33771023**
Actinomyces naeslundii		100.0	0.38	IND 1		
33771100	**Acceptable identification**					**33771100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		92.45	0.43	IND 1		
Actinomyces israelii		7.55	0.15	MNS 88		
33771101	**Good identification**					**33771101**
Actinomyces naeslundii		100.0	0.46	IND 1		
33771102	**Low discrimination**					**33771102**
Actinomyces naeslundii		83.53	0.43	IND 1		
Actinomyces israelii		10.23	0.18	MNS 88		
Actinomyces viscosus		6.24	0.00			
33771103	**Good identification**					**33771103**
Actinomyces naeslundii		100.0	0.46	IND 1		
33771120	**Good identification**					**33771120**
Actinomyces naeslundii		100.0	0.43	IND 1		
33771121	**Good identification**					**33771121**
Actinomyces naeslundii		100.0	0.46	IND 1		
33771122	**Good identification**					**33771122**
Actinomyces naeslundii		100.0	0.43	IND 1		
33771123	**Good identification**					**33771123**
Actinomyces naeslundii		100.0	0.46	IND 1		
33773000	**Low discrimination**					**33773000**
Actinomyces naeslundii		85.50	0.40	IND 1		
Actinomyces israelii		14.50	0.18	MNS 88		
33773001	**Good identification**					**33773001**
Actinomyces naeslundii		100.0	0.43	IND 1		
33773002	**Low discrimination**					**33773002**
Actinomyces naeslundii		79.72	0.40	IND 1		
Actinomyces israelii		20.28	0.21	MNS 88		
33773003	**Good identification**					**33773003**
Actinomyces naeslundii		100.0	0.43	IND 1		
33773020	**Good identification**					**33773020**
Actinomyces naeslundii		100.0	0.40	IND 1		
33773021	**Good identification**					**33773021**
Actinomyces naeslundii		100.0	0.43	IND 1		
33773022	**Good identification**					**33773022**
Actinomyces naeslundii		100.0	0.40	IND 1		
33773023	**Good identification**					**33773023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	IND 1		
33773100	**Low discrimination**					**33773100**
Actinomyces naeslundii		80.75	0.48	IND 1		
Actinomyces israelii		19.25	0.28	MNS 88		
33773101	**Good identification**					**33773101**
Actinomyces naeslundii		100.0	0.51	IND 1		
33773102	**Low discrimination**					**33773102**
Actinomyces naeslundii		73.66	0.48	IND 1		
Actinomyces israelii		26.34	0.31	MNS 88		
33773103	**Good identification**					**33773103**
Actinomyces naeslundii		100.0	0.51	IND 1		
33773120	**Good identification**					**33773120**
Actinomyces naeslundii		100.0	0.48	IND 1		
33773121	**Good identification**					**33773121**
Actinomyces naeslundii		100.0	0.51	IND 1		
33773122	**Good identification**					**33773122**
Actinomyces naeslundii		100.0	0.48	IND 1		
33773123	**Good identification**					**33773123**
Actinomyces naeslundii		100.0	0.51	IND 1		
33773140	**Low discrimination**					**33773140**
Actinomyces naeslundii		65.56	0.35	IND 1		
Actinomyces israelii		34.44	0.21	MNS 88		
33773141	**Good identification**					**33773141**
Actinomyces naeslundii		100.0	0.38	IND 1		
33773142	**Low discrimination**					**33773142**
Actinomyces naeslundii		55.93	0.35	IND 1		
Actinomyces israelii		44.07	0.24	MNS 88		
33773143	**Good identification**					**33773143**
Actinomyces naeslundii		100.0	0.38	IND 1		
33773160	**Good identification**					**33773160**
Actinomyces naeslundii		100.0	0.35	IND 1		
33773161	**Good identification**					**33773161**
Actinomyces naeslundii		100.0	0.38	IND 1		
33773162	**Good identification**					**33773162**
Actinomyces naeslundii		100.0	0.35	IND 1		
33773163	**Good identification**					**33773163**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	IND 1			
33775102	**Good identification**						**33775102**
Actinomyces israelii		100.0	0.36	IND 1			
33777000	**Good identification**						**33777000**
Actinomyces israelii		100.0	0.36	IND 1			
33777002	**Good identification**						**33777002**
Actinomyces israelii		100.0	0.39	IND 1			
33777100	**Good identification**						**33777100**
Actinomyces israelii		100.0	0.46	IND 1			
33777102	**Good identification**						**33777102**
Actinomyces israelii		100.0	0.49	IND 1			
33777112	**Good identification**						**33777112**
Actinomyces israelii		100.0	0.35	IND 1			
33777140	**Good identification**						**33777140**
Actinomyces israelii		100.0	0.39	IND 1			
33777142	**Good identification**						**33777142**
Actinomyces israelii		100.0	0.42	IND 1			
33777402	**Good identification**						**33777402**
Actinomyces israelii		100.0	0.38	IND 1			
33777500	**Good identification**						**33777500**
Actinomyces israelii		100.0	0.45	IND 1			
33777502	**Good identification**						**33777502**
Actinomyces israelii		100.0	0.48	IND 1			
33777540	**Good identification**						**33777540**
Actinomyces israelii		100.0	0.38	IND 1			
33777542	**Good identification**						**33777542**
Actinomyces israelii		100.0	0.41	IND 1			
34000000	**Low discrimination**						**34000000**
					COC	GRA	
Peptoniphilus asaccharolyticus		33.86	0.40	NIT 3	99	99	
Veillonella parvula		11.29	0.32	MNS 1	99	1	
Fusobacterium nucleatum subsp. nucleatum		11.06	0.32		1	1	
34000001	**Acceptable identification**						**34000001**
					COC	GRA	
Anaerococcus prevotii		90.65	0.49	IND 9	99	99	
Campylobacter ureolyticus		7.18	0.26	NIT 1	1	1	
Peptoniphilus asaccharolyticus		0.72	0.06		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
34000003	**Good identification**							**34000003**
					COC	GRA		
Anaerococcus prevotii		99.68	0.38	IND 9	99	99		
Campylobacter ureolyticus		0.32	-.08	NIT 1	1	1		
34000020	**Low discrimination**							**34000020**
					GRA	SPO	CAT	
Clostridium cadaveris		66.21	0.41	NIT 1	99	99	1	
Campylobacter gracilis		13.92	0.21	GLU 95	1	1	1	
Propionibacterium acnes		12.34	0.32		99	1	95	
34010000	**Very good identification**							**34010000**
					COC	GRA	CAT	
Fusobacterium varium		97.64	0.57	NIT 1	1	1	1	
Propionibacterium acnes		1.33	0.32		1	99	95	
Peptoniphilus asaccharolyticus		0.25	0.06		99	99	15	
34010020	**Good identification**							**34010020**
					GRA	SPO	CAT	
Propionibacterium acnes		94.16	0.66	GLU 99	99	1	95	
Clostridium cadaveris		5.10	0.41		99	99	1	
Fusobacterium varium		0.71	0.23		1	1	1	
34010060	**Good identification**							**34010060**
					SPO	CAT		
Propionibacterium acnes		99.75	0.55	GLU 99	1	95		
Clostridium cadaveris		0.25	0.07		99	1		
34011020	**Very good identification**							**34011020**
					SPO	CAT		
Propionibacterium acnes		99.95	0.66	GLU 99	1	95		
Clostridium cadaveris		0.06	0.07		99	1		
34011060	**Good identification**							**34011060**
Propionibacterium acnes		100.0	0.55	GLU 99				
34012020	**Very good identification**							**34012020**
					SPO	CAT		
Propionibacterium acnes		99.90	0.62	GLU 99	1	95		
Clostridium cadaveris		0.10	0.07		99	1		
34012060	**Good identification**							**34012060**
Propionibacterium acnes		100.0	0.50	GLU 99				
34013020	**Very good identification**							**34013020**
Propionibacterium acnes		100.0	0.62	GLU 99				
34013060	**Good identification**							**34013060**
Propionibacterium acnes		100.0	0.50	GLU 99				
34100000	**Low discrimination**							**34100000**
					COC	GRA	SPO	
Clostridium bifermentans		87.09	0.41	NIT 1	1	99	99	
Peptoniphilus asaccharolyticus		3.80	0.06	GLU 92	99	99	1	
Veillonella parvula		1.27	-.03		99	1	1	
34110000	**Low discrimination**							**34110000**
					GRA	SPO		
Prevotella intermedia		85.90	0.47	NIT 1	1	1		
Fusobacterium varium		10.57	0.23	GLU 90	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		3.20	0.20		99	99	
34111000	**Good identification**						**34111000**
Prevotella intermedia		100.0	0.47	NIT 1			
34400001	**Good identification**						**34400001**
					COC	GRA	
Anaerococcus prevotii		99.76	0.41	IND 9	99	99	
Campylobacter ureolyticus		0.24	-.08	NIT 1	1	1	
34510000	**Good identification**						**34510000**
Prevotella intermedia		100.0	0.50	NIT 1			
34511000	**Good identification**						**34511000**
Prevotella intermedia		100.0	0.50	NIT 1			
35000000	**Low discrimination**						**35000000**
					GRA	SPO	
Clostridium bifermentans		78.46	0.53	NIT 1	99	99	
Fusobacterium varium		9.26	0.32		1	1	
Clostridium cadaveris		3.84	0.29		99	99	
35000020	**Low discrimination**						**35000020**
					SPO	CAT	
Clostridium cadaveris		50.37	0.63	NIT 1	99	1	
Propionibacterium acnes		48.92	0.66		1	95	
Clostridium bifermentans		0.66	0.33		99	1	
35000060	**Good identification**						**35000060**
					SPO	CAT	
Propionibacterium acnes		95.24	0.55	FRU 99	1	95	
Clostridium cadaveris		4.51	0.29		99	1	
Clostridium bifermentans		0.25	0.09		99	1	
35001020	**Very good identification**						**35001020**
					SPO	CAT	
Propionibacterium acnes		98.95	0.66	FRU 99	1	95	
Clostridium cadaveris		1.03	0.29		99	1	
Atopobium minutum		0.01	0.06		1	1	
35001060	**Good identification**						**35001060**
Propionibacterium acnes		100.0	0.55	FRU 99			
35001520	**Good identification**						**35001520**
Odoribacter splanchnicus		100.0	0.51	ESL 99			
35002020	**Very good identification**						**35002020**
					SPO	CAT	
Propionibacterium acnes		98.16	0.62	FRU 99	1	95	
Clostridium cadaveris		1.82	0.29		99	1	
Clostridium bifermentans		0.02	-.01		99	1	
35002060	**Good identification**						**35002060**
Propionibacterium acnes		100.0	0.50	FRU 99			
35003020	**Very good identification**						**35003020**
Propionibacterium acnes		100.0	0.62	FRU 99			
35003060	**Good identification**						**35003060**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		100.0	0.50	FRU 99				
35010000	**Low discrimination**							**35010000**
					GRA	SPO	CAT	
Fusobacterium varium		70.63	0.66	NIT 1	1	1	1	
Propionibacterium acnes		28.44	0.66		99	1	95	
Clostridium bifermentans		0.32	0.31		99	99	1	
35010020	**Very good identification**							**35010020**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	1.00		99	1	95	
Clostridium cadaveris		1.03	0.63		99	99	1	
Fusobacterium varium		0.03	0.32		1	1	1	
35010021	**Very good identification**							**35010021**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	URE 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	
35010022	**Very good identification**							**35010022**
					SPO	CAT		
Propionibacterium acnes		98.92	0.66	BGL 1	1	95		
Clostridium cadaveris		1.03	0.29		99	1		
Atopobium minutum		0.02	0.11		1	1		
35010030	**Very good identification**							**35010030**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	MLZ 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	
35010040	**Low discrimination**							**35010040**
					GRA	SPO	CAT	
Propionibacterium acnes		89.57	0.55	NAG 99	99	1	95	
Fusobacterium varium		10.24	0.32		1	1	1	
Clostridium bifermentans		0.19	0.08		99	99	1	
35010060	**Very good identification**							**35010060**
					GRA	SPO	CAT	
Propionibacterium acnes		99.95	0.89		99	1	95	
Clostridium cadaveris		0.05	0.29		99	99	1	
Fusobacterium varium		0.00	-.03		1	1	1	
35010061	**Good identification**							**35010061**
Propionibacterium acnes		100.0	0.55	URE 1				
35010062	**Good identification**							**35010062**
Propionibacterium acnes		100.0	0.55	BGL 1				
35010070	**Good identification**							**35010070**
Propionibacterium acnes		100.0	0.55	MLZ 1				
35010120	**Very good identification**							**35010120**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	LAC 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	
35010160	**Good identification**							**35010160**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		100.0	0.55	LAC 1				
35010220	**Very good identification**							**35010220**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	RHA 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	
35010260	**Good identification**							**35010260**
Propionibacterium acnes		100.0	0.55	RHA 1				
35010420	**Very good identification**							**35010420**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	ARA 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	
35010460	**Good identification**							**35010460**
Propionibacterium acnes		100.0	0.55	ARA 1				
35011000	**Very good identification**							**35011000**
					GRA	CAT		
Propionibacterium acnes		96.92	0.66	NAG 99	99	95		
Fusobacterium varium		2.43	0.32		1	1		
Prevotella intermedia		0.54	0.29		1	1		
35011020	**Excellent identification**							**35011020**
					SPO	CAT		
Propionibacterium acnes		99.99	1.00		1	95		
Clostridium cadaveris		0.01	0.29		99	1		
Atopobium minutum		0.00	0.05		1	1		
35011021	**Very good identification**							**35011021**
Propionibacterium acnes		100.0	0.66	URE 1				
35011022	**Very good identification**							**35011022**
Propionibacterium acnes		100.0	0.66	BGL 1				
35011030	**Very good identification**							**35011030**
Propionibacterium acnes		100.0	0.66	MLZ 1				
35011040	**Good identification**							**35011040**
					GRA	CAT		
Propionibacterium acnes		99.89	0.55	NAG 99	99	95		
Fusobacterium varium		0.12	-.03		1	1		
35011060	**Very good identification**							**35011060**
Propionibacterium acnes		100.0	0.89					
35011061	**Good identification**							**35011061**
Propionibacterium acnes		100.0	0.55	URE 1				
35011062	**Good identification**							**35011062**
Propionibacterium acnes		100.0	0.55	BGL 1				
35011070	**Good identification**							**35011070**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		100.0	0.55	MLZ 1				
35011120	**Very good identification**							**35011120**
					GRA	CAT		
Propionibacterium acnes		99.25	0.66	LAC 1	99	95		
Odoribacter splanchnicus		0.74	0.21		1	1		
Clostridium perfringens		0.01	0.01		99	1		
35011160	**Good identification**							**35011160**
					GRA	CAT		
Propionibacterium acnes		99.62	0.55	LAC 1	99	95		
Odoribacter splanchnicus		0.38	0.05		1	1		
35011220	**Very good identification**							**35011220**
Propionibacterium acnes		100.0	0.66	RHA 1				
35011260	**Good identification**							**35011260**
Propionibacterium acnes		100.0	0.55	RHA 1				
35011420	**Very good identification**							**35011420**
					GRA	CAT		
Propionibacterium acnes		99.26	0.66	ARA 1	99	95		
Odoribacter splanchnicus		0.74	0.21		1	1		
35011460	**Good identification**							**35011460**
					GRA	CAT		
Propionibacterium acnes		99.62	0.55	ARA 1	99	95		
Odoribacter splanchnicus		0.38	0.05		1	1		
35011520	**Good identification**							**35011520**
					GRA	CAT		
Odoribacter splanchnicus		98.65	0.56	ESL 99	1	1		
Propionibacterium acnes		1.35	0.32		99	95		
35011560	**Good identification**							**35011560**
					GRA	CAT		
Odoribacter splanchnicus		97.38	0.39	ESL 99	1	1		
Propionibacterium acnes		2.62	0.20	SOR 10	99	95		
35012000	**Good identification**							**35012000**
					GRA	CAT		
Propionibacterium acnes		93.42	0.62	NAG 99	99	95		
Fusobacterium varium		4.17	0.32		1	1		
Eubacterium limosum		2.19	0.29		99	1		
35012020	**Excellent identification**							**35012020**
					GRA	SPO	CAT	
Propionibacterium acnes		99.98	0.96		99	1	95	
Clostridium cadaveris		0.02	0.29		99	99	1	
Fusobacterium varium		0.00	-.03		1	1	1	
35012021	**Very good identification**							**35012021**
Propionibacterium acnes		100.0	0.62	URE 1				
35012022	**Very good identification**							**35012022**
Propionibacterium acnes		100.0	0.62	BGL 1				
35012030	**Very good identification**							**35012030**
Propionibacterium acnes		100.0	0.62	MLZ 1				
35012040	**Good identification**							**35012040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Propionibacterium acnes		99.69	0.50	NAG 99	99	95	
Fusobacterium varium		0.21	-.03		1	1	
Eubacterium limosum		0.11	-.06		99	1	
35012060	**Very good identification**						**35012060**
Propionibacterium acnes		100.0	0.85				
35012061	**Good identification**						**35012061**
Propionibacterium acnes		100.0	0.50	URE 1			
35012062	**Good identification**						**35012062**
Propionibacterium acnes		100.0	0.50	BGL 1			
35012070	**Good identification**						**35012070**
Propionibacterium acnes		100.0	0.50	MLZ 1			
35012120	**Very good identification**						**35012120**
Propionibacterium acnes		100.0	0.62	LAC 1			
35012160	**Good identification**						**35012160**
Propionibacterium acnes		100.0	0.50	LAC 1			
35012220	**Very good identification**						**35012220**
Propionibacterium acnes		100.0	0.62	RHA 1			
35012260	**Good identification**						**35012260**
Propionibacterium acnes		100.0	0.50	RHA 1			
35012420	**Very good identification**						**35012420**
Propionibacterium acnes		100.0	0.62	ARA 1			
35012460	**Good identification**						**35012460**
Propionibacterium acnes		100.0	0.50	ARA 1			
35013000	**Very good identification**						**35013000**
					GRA	CAT	
Propionibacterium acnes		99.93	0.62	NAG 99	99	95	
Fusobacterium varium		0.05	-.03		1	1	
Eubacterium limosum		0.02	-.06		99	1	
35013020	**Excellent identification**						**35013020**
Propionibacterium acnes		100.0	0.96				
35013021	**Very good identification**						**35013021**
Propionibacterium acnes		100.0	0.62	URE 1			
35013022	**Very good identification**						**35013022**
Propionibacterium acnes		100.0	0.62	BGL 1			
35013030	**Very good identification**						**35013030**
Propionibacterium acnes		100.0	0.62	MLZ 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
35013040	**Good identification**							**35013040**
Propionibacterium acnes		100.0	0.50	NAG 99				
35013060	**Very good identification**							**35013060**
Propionibacterium acnes		100.0	0.85					
35013061	**Good identification**							**35013061**
Propionibacterium acnes		100.0	0.50	URE 1				
35013062	**Good identification**							**35013062**
Propionibacterium acnes		100.0	0.50	BGL 1				
35013070	**Good identification**							**35013070**
Propionibacterium acnes		100.0	0.50	MLZ 1				
35013120	**Very good identification**							**35013120**
Propionibacterium acnes		100.0	0.62	LAC 1				
35013160	**Good identification**							**35013160**
Propionibacterium acnes		100.0	0.50	LAC 1				
35013220	**Very good identification**							**35013220**
Propionibacterium acnes		100.0	0.62	RHA 1				
35013260	**Good identification**							**35013260**
Propionibacterium acnes		100.0	0.50	RHA 1				
35013420	**Very good identification**							**35013420**
Propionibacterium acnes		100.0	0.62	ARA 1				
35013460	**Good identification**							**35013460**
Propionibacterium acnes		100.0	0.50	ARA 1				
35014000	**Acceptable identification**							**35014000**
					COC	GRA	CAT	
Fusobacterium varium		90.90	0.42	NIT 1	1	1	1	
Propionibacterium acnes		8.87	0.32	XYL 4	1	99	95	
Gemella morbillorum		0.13	0.03		99	99	1	
35014020	**Very good identification**							**35014020**
					GRA	SPO	CAT	
Propionibacterium acnes		98.87	0.66	XYL 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.10	0.08		1	1	1	
35014060	**Good identification**							**35014060**
Propionibacterium acnes		100.0	0.55	XYL 1				
35015020	**Very good identification**							**35015020**
Propionibacterium acnes		100.0	0.66	XYL 1				
35015060	**Good identification**							**35015060**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium acnes		100.0	0.55	XYL 1				
35016020	**Very good identification**							**35016020**
Propionibacterium acnes		100.0	0.62	XYL 1				
35016060	**Good identification**							**35016060**
Propionibacterium acnes		100.0	0.50	XYL 1				
35017020	**Very good identification**							**35017020**
Propionibacterium acnes		100.0	0.62	XYL 1				
35017060	**Good identification**							**35017060**
Propionibacterium acnes		100.0	0.50	XYL 1				
35020020	**Low discrimination**							**35020020**
					SPO	CAT		
Clostridium cadaveris		84.11	0.41	NIT 1	99	1		
Propionibacterium acnes		15.68	0.32	TRE 5	1	95		
Clostridium bifermentans		0.21	-.01		99	1		
35030020	**Good identification**							**35030020**
					GRA	SPO	CAT	
Propionibacterium acnes		94.84	0.66	TRE 1	99	1	95	
Clostridium cadaveris		5.14	0.41		99	99	1	
Fusobacterium varium		0.02	-.03		1	1	1	
35030060	**Good identification**							**35030060**
					SPO	CAT		
Propionibacterium acnes		99.75	0.55	TRE 1	1	95		
Clostridium cadaveris		0.25	0.07		99	1		
35031020	**Very good identification**							**35031020**
					SPO	CAT		
Propionibacterium acnes		99.95	0.66	TRE 1	1	95		
Clostridium cadaveris		0.06	0.07		99	1		
35031060	**Good identification**							**35031060**
Propionibacterium acnes		100.0	0.55	TRE 1				
35032020	**Very good identification**							**35032020**
					SPO	CAT		
Propionibacterium acnes		99.90	0.62	TRE 1	1	95		
Clostridium cadaveris		0.10	0.07		99	1		
35032060	**Good identification**							**35032060**
Propionibacterium acnes		100.0	0.50	TRE 1				
35033020	**Very good identification**							**35033020**
Propionibacterium acnes		100.0	0.62	TRE 1				
35033060	**Good identification**							**35033060**
Propionibacterium acnes		100.0	0.50	TRE 1				
35050020	**Very good identification**							**35050020**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	CEL 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
35050060	**Good identification**							**35050060**
Propionibacterium acnes		100.0	0.55	CEL 1				
35051020	**Very good identification**							**35051020**
Propionibacterium acnes		100.0	0.66	CEL 1				
35051060	**Good identification**							**35051060**
Propionibacterium acnes		100.0	0.55	CEL 1				
35052020	**Very good identification**							**35052020**
Propionibacterium acnes		100.0	0.62	CEL 1				
35052060	**Good identification**							**35052060**
Propionibacterium acnes		100.0	0.50	CEL 1				
35053020	**Very good identification**							**35053020**
Propionibacterium acnes		100.0	0.62	CEL 1				
35053060	**Good identification**							**35053060**
Propionibacterium acnes		100.0	0.50	CEL 1				
35100000	**Good identification**							**35100000**
					COC	GRA	SPO	
Clostridium bifermentans		92.93	0.60	NIT 1	1	99	99	
Prevotella disiens		4.42	0.32		1	1	1	
Gemella morbillorum		1.13	0.30		99	99	1	
35100001	**Acceptable identification**							**35100001**
					GRA	SPO		
Clostridium sordellii		94.33	0.42	NIT 1	99	99		
Clostridium bifermentans		5.41	0.25	MNS 4	99	99		
Prevotella disiens		0.26	-.03		1	1		
35100020	**Low discrimination**							**35100020**
					SPO	CAT		
Clostridium cadaveris		55.67	0.41	NIT 1	99	1		
Clostridium bifermentans		32.52	0.39	MLT 5	99	1		
Propionibacterium acnes		10.38	0.32		1	95		
35100040	**Good identification**							**35100040**
					GRA	SPO		
Clostridium bifermentans		98.17	0.36	NIT 1	99	99		
Prevotella disiens		1.13	-.03	SOR 4	1	1		
Clostridium botulinum biovar B		0.70	-.03		99	99		
35105722	**Good identification**							**35105722**
Bacteroides eggerthii		100.0	0.39	NIT 1				
35110000	**Low discrimination**							**35110000**
					GRA	SPO		
Prevotella intermedia		85.41	0.63	NIT 1	1	1		
Clostridium bifermentans		4.07	0.38		99	99		
Fusobacterium varium		3.91	0.32		1	1		
35110001	**Acceptable identification**							**35110001**
					GRA	SPO		
Clostridium sordellii		93.28	0.42	NIT 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		5.92	0.29	MNS 4	1	1		
Clostridium bifermentans		0.28	0.04		99	99		
35110020	**Good identification**							**35110020**
					GRA	SPO	CAT	
Propionibacterium acnes		94.10	0.66	MLT 1	99	1	95	
Clostridium cadaveris		5.10	0.41		99	99	1	
Prevotella intermedia		0.52	0.29		1	1	1	
35110060	**Good identification**							**35110060**
					SPO	CAT		
Propionibacterium acnes		99.62	0.55	MLT 1	1	95		
Clostridium cadaveris		0.25	0.07		99	1		
Clostridium botulinum biovar B		0.13	0.02		99	1		
35111000	**Very good identification**							**35111000**
					COC	GRA	CAT	
Prevotella intermedia		97.34	0.63	NIT 1	1	1	1	
Propionibacterium acnes		1.79	0.32		1	99	95	
Gemella morbillorum		0.71	0.27		99	99	1	
35111020	**Very good identification**							**35111020**
					GRA	CAT		
Propionibacterium acnes		99.18	0.66	MLT 1	99	95		
Prevotella intermedia		0.55	0.29		1	1		
Clostridium perfringens		0.18	0.21		99	1		
35111060	**Good identification**							**35111060**
					CAT			
Propionibacterium acnes		99.85	0.55	MLT 1	95			
Clostridium perfringens		0.15	0.08		1			
35111120	**Low discrimination**							**35111120**
					GRA	CAT		
Clostridium perfringens		53.66	0.36	IND 1	99	1		
Propionibacterium acnes		41.54	0.32	SUC 91	99	95		
Prevotella bivia		3.90	0.03		1	1		
35112020	**Very good identification**							**35112020**
					SPO	CAT		
Propionibacterium acnes		99.90	0.62	MLT 1	1	95		
Clostridium cadaveris		0.10	0.07		99	1		
35112060	**Good identification**							**35112060**
Propionibacterium acnes		100.0	0.50	MLT 1				
35113020	**Very good identification**							**35113020**
Propionibacterium acnes		100.0	0.62	MLT 1				
35113060	**Good identification**							**35113060**
Propionibacterium acnes		100.0	0.50	MLT 1				
35131120	**Good identification**							**35131120**
Clostridium perfringens		97.37	0.40	IND 1				
Clostridium septicum		2.63	0.07	SUC 91				
35145722	**Good identification**							**35145722**
Bacteroides eggerthii		100.0	0.50	NIT 1				
35200000	**Good identification**							**35200000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Clostridium bifermentans		98.94	0.37	NIT 1	99	99		
Fusobacterium varium		1.06	-.03	SAL 10	1	1		
35210020	**Very good identification**							**35210020**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	SAL 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	
35210060	**Good identification**							**35210060**
Propionibacterium acnes		100.0	0.55	SAL 1				
35211020	**Very good identification**							**35211020**
Propionibacterium acnes		100.0	0.66	SAL 1				
35211060	**Good identification**							**35211060**
Propionibacterium acnes		100.0	0.55	SAL 1				
35212020	**Very good identification**							**35212020**
Propionibacterium acnes		100.0	0.62	SAL 1				
35212060	**Good identification**							**35212060**
Propionibacterium acnes		100.0	0.50	SAL 1				
35213020	**Very good identification**							**35213020**
Propionibacterium acnes		100.0	0.62	SAL 1				
35213060	**Good identification**							**35213060**
Propionibacterium acnes		100.0	0.50	SAL 1				
35300000	**Good identification**							**35300000**
					GRA	SPO		
Clostridium bifermentans		99.57	0.43	NIT 1	99	99		
Prevotella disiens		0.43	-.03	SAL 10	1	1		
35410020	**Very good identification**							**35410020**
					GRA	SPO	CAT	
Propionibacterium acnes		98.95	0.66	RAF 1	99	1	95	
Clostridium cadaveris		1.03	0.29		99	99	1	
Fusobacterium varium		0.03	-.03		1	1	1	
35410060	**Good identification**							**35410060**
Propionibacterium acnes		100.0	0.55	RAF 1				
35411020	**Very good identification**							**35411020**
Propionibacterium acnes		100.0	0.66	RAF 1				
35411060	**Good identification**							**35411060**
Propionibacterium acnes		100.0	0.55	RAF 1				
35412020	**Very good identification**							**35412020**
Propionibacterium acnes		100.0	0.62	RAF 1				
35412060	**Good identification**							**35412060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		100.0	0.50	RAF 1			
35413020	**Very good identification**						**35413020**
Propionibacterium acnes		100.0	0.62	RAF 1			
35413060	**Good identification**						**35413060**
Propionibacterium acnes		100.0	0.50	RAF 1			
35510000	**Very good identification**						**35510000**
					GRA	SPO	
Prevotella intermedia		99.91	0.66	NIT 1	1	1	
Clostridium bifermentans		0.03	0.04		99	99	
Prevotella disiens		0.03	-.03		1	1	
35511000	**Very good identification**						**35511000**
Prevotella intermedia		100.0	0.66	NIT 1			
36110000	**Good identification**						**36110000**
Prevotella intermedia		100.0	0.41	NIT 1			
36111000	**Good identification**						**36111000**
Prevotella intermedia		100.0	0.41	NIT 1			
36510000	**Good identification**						**36510000**
Prevotella intermedia		100.0	0.44	NIT 1			
36511000	**Good identification**						**36511000**
Prevotella intermedia		100.0	0.44	NIT 1			
37000020	**Low discrimination**						**37000020**
					SPO	CAT	
Clostridium cadaveris		88.21	0.44	NIT 1	99	1	
Propionibacterium acnes		11.50	0.32	SUC 7	1	95	
Clostridium bifermentans		0.16	-.01		99	1	
37010020	**Good identification**						**37010020**
					SPO	CAT	
Propionibacterium acnes		92.76	0.66	SUC 1	1	95	
Clostridium cadaveris		7.19	0.44		99	1	
Propionibacterium granulosum		0.02	0.11		1	90	
37010060	**Good identification**						**37010060**
					SPO	CAT	
Propionibacterium acnes		99.65	0.55	SUC 1	1	95	
Clostridium cadaveris		0.36	0.09		99	1	
37011020	**Very good identification**						**37011020**
					SPO	CAT	
Propionibacterium acnes		99.89	0.66	SUC 1	1	95	
Clostridium cadaveris		0.08	0.09		99	1	
Propionibacterium granulosum		0.02	0.11		1	90	
37011060	**Good identification**						**37011060**
Propionibacterium acnes		100.0	0.55	SUC 1			
37012020	**Very good identification**						**37012020**
					SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		99.84	0.62	SUC 1	1	95	
Clostridium cadaveris		0.14	0.09		99	1	
Propionibacterium granulosum		0.03	0.09		1	90	
37012060	**Good identification**						**37012060**
Propionibacterium acnes		100.0	0.50	SUC 1			
37013020	**Very good identification**						**37013020**
Propionibacterium acnes		99.98	0.62	SUC 1			
Propionibacterium granulosum		0.03	0.09				
37013060	**Good identification**						**37013060**
Propionibacterium acnes		100.0	0.50	SUC 1			
37101101	**Very good identification**						**37101101**
Clostridium chauvoei		100.0	0.57	IND 1			
37101103	**Good identification**						**37101103**
					SPO		
Clostridium chauvoei		98.44	0.44	IND 1	99		
Actinomyces viscosus		1.56	0.12		1		
37101121	**Good identification**						**37101121**
Clostridium chauvoei		100.0	0.48	IND 1			
37101123	**Good identification**						**37101123**
Clostridium chauvoei		100.0	0.36	IND 1			
37110000	**Good identification**						**37110000**
					COC	GRA	
Prevotella intermedia		93.40	0.58	NIT 1	1	1	
Actinomyces odontolyticus		3.55	0.39		1	99	
Gemella morbillorum		2.55	0.32		99	99	
37110002	**Low discrimination**						**37110002**
					GRA		
Actinomyces viscosus		79.84	0.41	IND 1	99		
Actinomyces odontolyticus		14.06	0.35	RAF 94	99		
Prevotella intermedia		6.09	0.23		1		
37110102	**Good identification**						**37110102**
Actinomyces viscosus		95.47	0.45	IND 1			
Actinomyces odontolyticus		2.99	0.27	RAF 94			
Propionibacterium propionicum		1.55	0.29				
37110103	**Acceptable identification**						**37110103**
					SPO		
Actinomyces viscosus		94.92	0.37	IND 1	1		
Clostridium chauvoei		4.57	0.16	RAF 94	99		
Propionibacterium propionicum		0.51	0.12		1		
37110200	**Low discrimination**						**37110200**
					GRA		
Actinomyces odontolyticus		87.96	0.44	IND 1	99		
Prevotella intermedia		12.04	0.23	MNS 9	1		
37110202	**Good identification**						**37110202**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		97.13	0.40	IND 1			
Actinomyces viscosus		2.87	0.07	MNS 9			
37110300	**Good identification**						**37110300**
Actinomyces odontolyticus		100.0	0.35	IND 1			
37111000	**Very good identification**						**37111000**
					COC	GRA	
Prevotella intermedia		95.62	0.58	NIT 1	1	1	
Actinomyces odontolyticus		2.42	0.36		1	99	
Gemella morbillorum		1.41	0.27		99	99	
37111002	**Low discrimination**						**37111002**
					GRA		
Actinomyces viscosus		74.81	0.38	IND 1	99		
Actinomyces odontolyticus		14.33	0.32	RAF 94	99		
Prevotella intermedia		9.32	0.23		1		
37111020	**Low discrimination**						**37111020**
					GRA	CAT	
Clostridium perfringens		55.57	0.38	IND 1	99	1	
Propionibacterium acnes		31.20	0.32	LAC 88	99	95	
Prevotella intermedia		8.43	0.23		1	1	
37111101	**Very good identification**						**37111101**
					SPO		
Clostridium chauvoei		99.94	0.63	IND 1	99		
Actinomyces naeslundii		0.03	0.26		1		
Actinomyces viscosus		0.02	-.01		1		
37111102	**Low discrimination**						**37111102**
Actinomyces viscosus		86.89	0.42	IND 1			
Clostridium perfringens		4.28	0.21	RAF 94			
Actinomyces odontolyticus		2.96	0.24				
37111103	**Low discrimination**						**37111103**
					SPO		
Clostridium chauvoei		88.36	0.50	IND 1	99		
Actinomyces viscosus		11.37	0.34		1		
Actinomyces naeslundii		0.17	0.26		1		
37111120	**Good identification**						**37111120**
					GRA	SPO	
Clostridium perfringens		96.52	0.53	IND 1	99	84	
Prevotella melaninogenica		1.61	0.15		1	1	
Clostridium chauvoei		1.41	0.21		99	99	
37111121	**Good identification**						**37111121**
					SPO		
Clostridium chauvoei		99.21	0.55	IND 1	99		
Clostridium perfringens		0.69	0.19		84		
Actinomyces naeslundii		0.10	0.26		1		
37111122	**Good identification**						**37111122**
					GRA	CAT	
Clostridium perfringens		95.66	0.47	IND 1	99	1	
Capnocytophaga ochracea		2.04	0.15		1	90	
Prevotella melaninogenica		0.65	0.03		1	1	
37111123	**Good identification**						**37111123**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		97.26	0.42	IND 1	99	
Clostridium perfringens		1.81	0.13		84	
Actinomyces naeslundii		0.55	0.26		1	
37111160	**Good identification**					**37111160**
Clostridium perfringens		100.0	0.40	IND 1		
37111200	**Low discrimination**					**37111200**
					GRA	
Actinomyces odontolyticus		82.97	0.41	IND 1	99	
Prevotella intermedia		17.03	0.23	MNS 9	1	
37111202	**Good identification**					**37111202**
Actinomyces odontolyticus		97.36	0.37	IND 1		
Actinomyces viscosus		2.64	0.03	MNS 9		
37112100	**Very good identification**					**37112100**
Propionibacterium propionicum		100.0	0.60	IND 1		
37112101	**Good identification**					**37112101**
Propionibacterium propionicum		100.0	0.44	IND 1		
37112102	**Very good identification**					**37112102**
Propionibacterium propionicum		99.38	0.63	IND 1		
Actinomyces viscosus		0.63	0.11			
37112103	**Good identification**					**37112103**
Propionibacterium propionicum		98.15	0.47	IND 1		
Actinomyces viscosus		1.85	0.03	URE 10		
37112140	**Very good identification**					**37112140**
Propionibacterium propionicum		100.0	0.60	IND 1		
37112141	**Good identification**					**37112141**
Propionibacterium propionicum		100.0	0.44	IND 1		
37112142	**Very good identification**					**37112142**
Propionibacterium propionicum		100.0	0.63	IND 1		
37112143	**Good identification**					**37112143**
Propionibacterium propionicum		100.0	0.47	IND 1		
37113100	**Very good identification**					**37113100**
Propionibacterium propionicum		99.64	0.60	IND 1		
Actinomyces naeslundii		0.36	0.28			
37113101	**Low discrimination**					**37113101**
					SPO	
Propionibacterium propionicum		55.16	0.44	IND 1	1	
Clostridium chauvoei		42.15	0.29	URE 10	99	
Actinomyces naeslundii		2.69	0.31		1	
37113102	**Very good identification**					**37113102**
Propionibacterium propionicum		99.38	0.63	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		0.38	0.08			
Actinomyces naeslundii		0.24	0.28			
37113103	**Low discrimination**					**37113103**
					SPO	
Propionibacterium propionicum		88.19	0.47	IND 1	1	
Clostridium chauvoei		7.93	0.16	URE 10	99	
Actinomyces naeslundii		2.86	0.31		1	
37113140	**Very good identification**					**37113140**
Propionibacterium propionicum		100.0	0.60	IND 1		
37113141	**Good identification**					**37113141**
Propionibacterium propionicum		100.0	0.44	IND 1		
37113142	**Very good identification**					**37113142**
Propionibacterium propionicum		100.0	0.63	IND 1		
37113143	**Good identification**					**37113143**
Propionibacterium propionicum		100.0	0.47	IND 1		
37114200	**Good identification**					**37114200**
Actinomyces odontolyticus		100.0	0.38	IND 1		
37115200	**Good identification**					**37115200**
Actinomyces odontolyticus		100.0	0.35	IND 1		
37121101	**Good identification**					**37121101**
Clostridium chauvoei		100.0	0.40	IND 1		
37131020	**Good identification**					**37131020**
					CAT	
Clostridium perfringens		99.25	0.43	IND 1	1	
Propionibacterium granulosum		0.76	0.14	LAC 88	90	
37131022	**Good identification**					**37131022**
Clostridium perfringens		100.0	0.37	IND 1		
37131101	**Good identification**					**37131101**
					SPO	
Clostridium chauvoei		99.42	0.47	IND 1	99	
Actinomyces naeslundii		0.58	0.31	TRE 10	1	
37131120	**Very good identification**					**37131120**
					SPO	
Clostridium perfringens		99.81	0.57	IND 1	84	
Actinomyces naeslundii		0.10	0.28		1	
Clostridium chauvoei		0.09	0.04		99	
37131121	**Low discrimination**					**37131121**
					SPO	
Clostridium chauvoei		88.16	0.38	IND 1	99	
Clostridium perfringens		10.31	0.23	TRE 10	84	
Actinomyces naeslundii		1.53	0.31		1	
37131122	**Good identification**					**37131122**
Clostridium perfringens		99.79	0.52	IND 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	0.21	0.28			
37131160	**Good identification**					**37131160**
	Clostridium perfringens	100.0	0.45	IND 1		
37131162	**Good identification**					**37131162**
	Clostridium perfringens	100.0	0.39	IND 1		
37132100	**Very good identification**					**37132100**
	Propionibacterium propionicum	100.0	0.58	IND 1		
37132101	**Good identification**					**37132101**
	Propionibacterium propionicum	100.0	0.42	IND 1		
37132102	**Very good identification**					**37132102**
	Propionibacterium propionicum	100.0	0.61	IND 1		
37132103	**Good identification**					**37132103**
	Propionibacterium propionicum	100.0	0.45	IND 1		
37132140	**Very good identification**					**37132140**
	Propionibacterium propionicum	100.0	0.58	IND 1		
37132141	**Good identification**					**37132141**
	Propionibacterium propionicum	100.0	0.42	IND 1		
37132142	**Very good identification**					**37132142**
	Propionibacterium propionicum	100.0	0.61	IND 1		
37132143	**Good identification**					**37132143**
	Propionibacterium propionicum	100.0	0.45	IND 1		
37133100	**Very good identification**					**37133100**
	Propionibacterium propionicum	99.15	0.58	IND 1		
	Actinomyces naeslundii	0.85	0.33			
37133101	**Low discrimination**					**37133101**
					SPO	
	Propionibacterium propionicum	82.01	0.42	IND 1	1	
	Actinomyces naeslundii	9.48	0.36	URE 10	1	
	Clostridium chauvoei	8.51	0.12		99	
37133102	**Very good identification**					**37133102**
	Propionibacterium propionicum	99.43	0.61	IND 1		
	Actinomyces naeslundii	0.57	0.33			
37133103	**Acceptable identification**					**37133103**
					SPO	
	Propionibacterium propionicum	91.80	0.45	IND 1	1	
	Actinomyces naeslundii	7.08	0.36	URE 10	1	
	Clostridium chauvoei	1.12	-.01		99	
37133121	**Low discrimination**					**37133121**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		76.97	0.36	IND 1	1	
Clostridium chauvoei		23.03	0.04	RAF 92	99	
37133123	**Good identification**					**37133123**
Actinomyces naeslundii		100.0	0.36	IND 1		
37133140	**Very good identification**					**37133140**
Propionibacterium propionicum		100.0	0.58	IND 1		
37133141	**Good identification**					**37133141**
Propionibacterium propionicum		98.00	0.42	IND 1		
Actinomyces naeslundii		2.00	0.23	URE 10		
37133142	**Very good identification**					**37133142**
Propionibacterium propionicum		100.0	0.61	IND 1		
37133143	**Good identification**					**37133143**
Propionibacterium propionicum		98.66	0.45	IND 1		
Actinomyces naeslundii		1.34	0.23	URE 10		
37150102	**Good identification**					**37150102**
Actinomyces viscosus		100.0	0.36	IND 1		
37173100	**Low discrimination**					**37173100**
Actinomyces naeslundii		52.91	0.35	IND 1		
Propionibacterium propionicum		47.09	0.24	RAF 92		
37173101	**Good identification**					**37173101**
Actinomyces naeslundii		100.0	0.38	IND 1		
37173103	**Good identification**					**37173103**
Actinomyces naeslundii		100.0	0.38	IND 1		
37173120	**Good identification**					**37173120**
Actinomyces naeslundii		100.0	0.35	IND 1		
37173121	**Good identification**					**37173121**
Actinomyces naeslundii		100.0	0.38	IND 1		
37173122	**Good identification**					**37173122**
Actinomyces naeslundii		100.0	0.35	IND 1		
37173123	**Good identification**					**37173123**
Actinomyces naeslundii		100.0	0.38	IND 1		
37310000	**Low discrimination**					**37310000**
					GRA	
Actinomyces odontolyticus		87.96	0.44	IND 1	99	
Prevotella intermedia		12.04	0.23	MNS 9	1	
37310002	**Low discrimination**					**37310002**
Actinomyces odontolyticus		56.26	0.40	IND 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		43.74	0.31	MNS 9		
37310100	**Good identification**					**37310100**
Actinomyces odontolyticus		97.36	0.35	IND 1		
Actinomyces viscosus		2.64	0.01	MNS 9		
37310102	**Low discrimination**					**37310102**
Actinomyces viscosus		81.40	0.36	IND 1		
Actinomyces odontolyticus		18.60	0.32	RAF 94		
37310200	**Good identification**					**37310200**
Actinomyces odontolyticus		100.0	0.49	IND 1		
37310202	**Good identification**					**37310202**
Actinomyces odontolyticus		100.0	0.45	IND 1		
37310220	**Good identification**					**37310220**
Actinomyces odontolyticus		100.0	0.35	IND 1		
37310300	**Good identification**					**37310300**
Actinomyces odontolyticus		100.0	0.40	IND 1		
37310302	**Good identification**					**37310302**
Actinomyces odontolyticus		97.77	0.36	IND 1		
Actinomyces viscosus		2.23	0.01	MNS 9		
37311000	**Low discrimination**					**37311000**
					GRA	
Actinomyces odontolyticus		80.85	0.41	IND 1	99	
Prevotella intermedia		16.59	0.23	MNS 9	1	
Actinomyces naeslundii		0.96	0.25		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
37311002	**Low discrimination**					**37311002**
Actinomyces odontolyticus		57.66	0.37	IND 1		
Actinomyces viscosus		41.22	0.28	MNS 9		
Actinomyces naeslundii		1.12	0.25			
37311120	**Good identification**					**37311120**
Clostridium perfringens		96.68	0.42	IND 1		
Actinomyces odontolyticus		2.00	0.19			
Actinomyces naeslundii		1.31	0.33			
37311122	**Acceptable identification**					**37311122**
Clostridium perfringens		94.71	0.37	IND 1		
Actinomyces naeslundii		2.74	0.33			
Actinomyces odontolyticus		2.56	0.15			
37311123	**Low discrimination**					**37311123**
					SPO	
Actinomyces naeslundii		57.53	0.36	IND 1	1	
Clostridium chauvoei		29.06	0.08	RAF 92	99	
Clostridium perfringens		13.42	0.03		84	
37311200	**Good identification**					**37311200**
Actinomyces odontolyticus		100.0	0.46	IND 1		
37311202	**Good identification**					**37311202**
Actinomyces odontolyticus		100.0	0.42	IND 1		
37311300	**Good identification**					**37311300**
Actinomyces odontolyticus		100.0	0.37	IND 1		
37313100	**Low discrimination**					**37313100**
Actinomyces naeslundii		55.89	0.38	IND 1		
Propionibacterium propionicum		44.11	0.26	RAF 92		
37313101	**Good identification**					**37313101**
Actinomyces naeslundii		100.0	0.41	IND 1		
37313103	**Acceptable identification**					**37313103**
Actinomyces naeslundii		91.94	0.41	IND 1		
Propionibacterium propionicum		8.06	0.12	RAF 92		
37313120	**Low discrimination**					**37313120**
Actinomyces naeslundii		72.31	0.38	IND 1		
Clostridium perfringens		27.69	0.08	RAF 92		
37313121	**Good identification**					**37313121**
Actinomyces naeslundii		100.0	0.41	IND 1		
37313122	**Low discrimination**					**37313122**
Actinomyces naeslundii		84.73	0.38	IND 1		
Clostridium perfringens		15.27	0.03	RAF 92		
37313123	**Good identification**					**37313123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	IND 1		
37314000	**Good identification**					**37314000**
Actinomyces odontolyticus		100.0	0.38	IND 1		
37314200	**Good identification**					**37314200**
Actinomyces odontolyticus		100.0	0.43	IND 1		
37314202	**Good identification**					**37314202**
Actinomyces odontolyticus		100.0	0.40	IND 1		
37315000	**Good identification**					**37315000**
Actinomyces odontolyticus		100.0	0.35	IND 1		
37315200	**Good identification**					**37315200**
Actinomyces odontolyticus		100.0	0.40	IND 1		
37315202	**Good identification**					**37315202**
Actinomyces odontolyticus		100.0	0.37	IND 1		
37331101	**Low discrimination**					**37331101**
					SPO	
Actinomyces naeslundii		67.04	0.41	IND 1	1	
Clostridium chauvoei		32.96	0.12	RAF 92	99	
37331103	**Low discrimination**					**37331103**
					SPO	
Actinomyces naeslundii		77.78	0.41	IND 1	1	
Actinomyces viscosus		15.48	0.05	RAF 92	1	
Clostridium chauvoei		6.75	-.01		99	
37331120	**Good identification**					**37331120**
					COC	
Clostridium perfringens		98.38	0.47	IND 1	1	
Actinomyces naeslundii		1.40	0.38		1	
Atopobium parvulum		0.22	0.01		99	
37331121	**Low discrimination**					**37331121**
					SPO	
Actinomyces naeslundii		61.05	0.41	IND 1	1	
Clostridium perfringens		28.95	0.13	RAF 92	84	
Clostridium chauvoei		10.00	0.04		99	
37331122	**Good identification**					**37331122**
Clostridium perfringens		97.07	0.41	IND 1		
Actinomyces naeslundii		2.93	0.38			
37331123	**Low discrimination**					**37331123**
Actinomyces naeslundii		81.76	0.41	IND 1		
Clostridium perfringens		18.24	0.07	RAF 92		
37333001	**Good identification**					**37333001**
Actinomyces naeslundii		100.0	0.38	IND 1		
37333003	**Good identification**					**37333003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	IND 1		
37333021	**Good identification**					**37333021**
Actinomyces naeslundii		100.0	0.38	IND 1		
37333023	**Good identification**					**37333023**
Actinomyces naeslundii		100.0	0.38	IND 1		
37333100	**Low discrimination**					**37333100**
Actinomyces naeslundii		75.04	0.43	IND 1		
Propionibacterium propionicum		24.96	0.24	RAF 92		
37333101	**Good identification**					**37333101**
Actinomyces naeslundii		100.0	0.46	IND 1		
37333102	**Low discrimination**					**37333102**
Actinomyces naeslundii		66.71	0.43	IND 1		
Propionibacterium propionicum		33.29	0.27	RAF 92		
37333103	**Good identification**					**37333103**
Actinomyces naeslundii		100.0	0.46	IND 1		
37333120	**Low discrimination**					**37333120**
Actinomyces naeslundii		73.19	0.43	IND 1		
Clostridium perfringens		26.81	0.13	RAF 92		
37333121	**Good identification**					**37333121**
Actinomyces naeslundii		100.0	0.46	IND 1		
37333122	**Low discrimination**					**37333122**
Actinomyces naeslundii		85.30	0.43	IND 1		
Clostridium perfringens		14.71	0.07	RAF 92		
37333123	**Good identification**					**37333123**
Actinomyces naeslundii		100.0	0.46	IND 1		
37351101	**Good identification**					**37351101**
Actinomyces naeslundii		100.0	0.38	IND 1		
37351121	**Good identification**					**37351121**
Actinomyces naeslundii		100.0	0.38	IND 1		
37351123	**Good identification**					**37351123**
Actinomyces naeslundii		100.0	0.38	IND 1		
37353001	**Good identification**					**37353001**
Actinomyces naeslundii		100.0	0.35	IND 1		
37353003	**Good identification**					**37353003**
Actinomyces naeslundii		100.0	0.35	IND 1		
37353021	**Good identification**					**37353021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	IND 1		
37353023	**Good identification**					**37353023**
Actinomyces naeslundii		100.0	0.35	IND 1		
37353100	**Low discrimination**					**37353100**
Actinomyces naeslundii		88.85	0.40	IND 1		
Actinomyces israelii		11.15	0.15	RAF 92		
37353101	**Good identification**					**37353101**
Actinomyces naeslundii		100.0	0.43	IND 1		
37353102	**Low discrimination**					**37353102**
Actinomyces naeslundii		75.14	0.40	IND 1		
Actinomyces israelii		14.15	0.18	RAF 92		
Lactobacillus acidophilus		10.71	0.01			
37353103	**Good identification**					**37353103**
Actinomyces naeslundii		100.0	0.43	IND 1		
37353120	**Good identification**					**37353120**
Actinomyces naeslundii		100.0	0.40	IND 1		
37353121	**Good identification**					**37353121**
Actinomyces naeslundii		100.0	0.43	IND 1		
37353122	**Low discrimination**					**37353122**
					SPO	
Actinomyces naeslundii		70.22	0.40	IND 1	1	
Clostridium barati		11.25	-.01	RAF 92	99	
Lactobacillus acidophilus		10.00	0.01		1	
37353123	**Good identification**					**37353123**
Actinomyces naeslundii		100.0	0.43	IND 1		
37357102	**Good identification**					**37357102**
Actinomyces israelii		100.0	0.36	IND 1		
37371001	**Good identification**					**37371001**
Actinomyces naeslundii		100.0	0.35	IND 1		
37371003	**Good identification**					**37371003**
Actinomyces naeslundii		100.0	0.35	IND 1		
37371021	**Good identification**					**37371021**
Actinomyces naeslundii		100.0	0.35	IND 1		
37371023	**Good identification**					**37371023**
Actinomyces naeslundii		100.0	0.35	IND 1		
37371101	**Good identification**					**37371101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	IND 1		
37371103	**Good identification**					**37371103**
Actinomyces naeslundii		100.0	0.43	IND 1		
37371121	**Good identification**					**37371121**
Actinomyces naeslundii		100.0	0.43	IND 1		
37371123	**Good identification**					**37371123**
Actinomyces naeslundii		100.0	0.43	IND 1		
37373000	**Low discrimination**					**37373000**
Actinomyces naeslundii		85.25	0.37	IND 1		
Actinomyces israelii		14.75	0.15	RAF 92		
37373001	**Good identification**					**37373001**
Actinomyces naeslundii		100.0	0.40	IND 1		
37373002	**Low discrimination**					**37373002**
Actinomyces naeslundii		79.39	0.37	IND 1		
Actinomyces israelii		20.61	0.18	RAF 92		
37373003	**Good identification**					**37373003**
Actinomyces naeslundii		100.0	0.40	IND 1		
37373020	**Good identification**					**37373020**
Actinomyces naeslundii		100.0	0.37	IND 1		
37373021	**Good identification**					**37373021**
Actinomyces naeslundii		100.0	0.40	IND 1		
37373022	**Good identification**					**37373022**
Actinomyces naeslundii		100.0	0.37	IND 1		
37373023	**Good identification**					**37373023**
Actinomyces naeslundii		100.0	0.40	IND 1		
37373100	**Low discrimination**					**37373100**
					SPO	
Actinomyces naeslundii		76.58	0.45	IND 1	1	
Actinomyces israelii		18.63	0.25	RAF 92	1	
Clostridium tertium		4.79	0.01		99	
37373101	**Good identification**					**37373101**
Actinomyces naeslundii		100.0	0.48	IND 1		
37373102	**Low discrimination**					**37373102**
Actinomyces naeslundii		55.61	0.45	IND 1		
Actinomyces israelii		20.29	0.28	RAF 92		
Lactobacillus acidophilus		13.67	0.10			
37373103	**Good identification**					**37373103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	IND 1		
37373120	**Low discrimination**					**37373120**
					SPO	
Actinomyces naeslundii		73.28	0.45	IND 1	1	
Clostridium tertium		13.76	0.09	RAF 92	99	
Clostridium ramosum		12.97	0.07		92	
37373121	**Good identification**					**37373121**
Actinomyces naeslundii		100.0	0.48	IND 1		
37373122	**Low discrimination**					**37373122**
					SPO	
Actinomyces naeslundii		39.74	0.45	IND 1	1	
Clostridium ramosum		28.12	0.17	RAF 92	92	
Clostridium tertium		22.38	0.17		99	
37373123	**Good identification**					**37373123**
Actinomyces naeslundii		100.0	0.48	IND 1		
37377002	**Good identification**					**37377002**
Actinomyces israelii		100.0	0.36	IND 1		
37377100	**Good identification**					**37377100**
					SPO	
Actinomyces israelii		97.00	0.43	IND 1	1	
Clostridium tertium		3.00	0.03		99	
37377102	**Acceptable identification**					**37377102**
					SPO	
Actinomyces israelii		94.18	0.46	IND 1	1	
Clostridium tertium		5.82	0.11		99	
37377140	**Good identification**					**37377140**
Actinomyces israelii		100.0	0.36	IND 1		
37377142	**Good identification**					**37377142**
Actinomyces israelii		100.0	0.39	IND 1		
37377500	**Good identification**					**37377500**
Actinomyces israelii		100.0	0.42	IND 1		
37377502	**Good identification**					**37377502**
Actinomyces israelii		100.0	0.45	IND 1		
37377542	**Good identification**					**37377542**
Actinomyces israelii		100.0	0.38	IND 1		
37473101	**Good identification**					**37473101**
Actinomyces naeslundii		100.0	0.36	IND 1		
37473103	**Good identification**					**37473103**
Actinomyces naeslundii		100.0	0.36	IND 1		
37473121	**Good identification**					**37473121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.36	IND 1			
37473123	**Good identification**						**37473123**
Actinomyces naeslundii		100.0	0.36	IND 1			
37500002	**Good identification**						**37500002**
Actinomyces viscosus		100.0	0.40	IND 1			
37500102	**Good identification**						**37500102**
Actinomyces viscosus		100.0	0.44	IND 1			
37500103	**Good identification**						**37500103**
Actinomyces viscosus		100.0	0.36	IND 1			
37501002	**Good identification**						**37501002**
Actinomyces viscosus		100.0	0.36	IND 1			
37501102	**Good identification**						**37501102**
Actinomyces viscosus		100.0	0.40	IND 1			
37510000	**Very good identification**						**37510000**
					GRA	CAT	
Prevotella intermedia		98.44	0.61	NIT 1	1	1	
Actinomyces viscosus		1.38	0.27		99	76	
Propionibacterium granulosum		0.11	0.18		99	90	
37510002	**Very good identification**						**37510002**
					GRA		
Actinomyces viscosus		99.24	0.62	IND 1	99		
Prevotella intermedia		0.73	0.26		1		
Actinomyces odontolyticus		0.04	0.09		99		
37510003	**Good identification**						**37510003**
Actinomyces viscosus		100.0	0.53	IND 1			
37510102	**Very good identification**						**37510102**
Actinomyces viscosus		99.93	0.66	IND 1			
Propionibacterium propionicum		0.07	0.26				
37510103	**Very good identification**						**37510103**
Actinomyces viscosus		100.0	0.58	IND 1			
37511000	**Very good identification**						**37511000**
					GRA		
Prevotella intermedia		98.77	0.61	NIT 1	1		
Actinomyces viscosus		0.85	0.24		99		
Actinomyces naeslundii		0.13	0.34		99		
37511002	**Very good identification**						**37511002**
					GRA		
Actinomyces viscosus		98.55	0.58	IND 1	99		
Prevotella intermedia		1.18	0.26		1		
Actinomyces naeslundii		0.15	0.34		99		
37511003	**Good identification**						**37511003**
Actinomyces viscosus		99.34	0.50	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.67	0.37			
37511021	**Good identification**					**37511021**
Actinomyces naeslundii		100.0	0.37	IND 1		
37511102	**Very good identification**					**37511102**
					GRA	
Actinomyces viscosus		99.30	0.62	IND 1	99	
Actinomyces naeslundii		0.24	0.42		99	
Capnocytophaga ochracea		0.22	0.15		1	
37511103	**Good identification**					**37511103**
					SPO	
Actinomyces viscosus		98.45	0.54	IND 1	1	
Actinomyces naeslundii		1.06	0.45		1	
Clostridium chauvoei		0.49	0.16		99	
37511120	**Low discrimination**					**37511120**
					GRA	
Clostridium perfringens		52.49	0.40	IND 1	99	
Prevotella melaninogenica		41.37	0.32		1	
Actinomyces naeslundii		3.04	0.42		99	
37511121	**Low discrimination**					**37511121**
					SPO	
Actinomyces naeslundii		47.83	0.45	IND 1	1	
Clostridium chauvoei		42.21	0.21		99	
Clostridium perfringens		5.57	0.06		84	
37511123	**Low discrimination**					**37511123**
					SPO	
Actinomyces naeslundii		45.30	0.45	IND 1	1	
Actinomyces viscosus		42.67	0.20		1	
Clostridium chauvoei		7.05	0.08		99	
37512100	**Very good identification**					**37512100**
Propionibacterium propionicum		99.74	0.57	IND 1		
Actinomyces naeslundii		0.26	0.23			
37512101	**Good identification**					**37512101**
Propionibacterium propionicum		96.62	0.41	IND 1		
Actinomyces naeslundii		3.38	0.26	URE 10		
37512102	**Low discrimination**					**37512102**
Propionibacterium propionicum		86.99	0.60	IND 1		
Actinomyces viscosus		12.86	0.32			
Actinomyces naeslundii		0.15	0.23			
37512103	**Low discrimination**					**37512103**
Propionibacterium propionicum		68.17	0.44	IND 1		
Actinomyces viscosus		30.24	0.24	URE 10		
Actinomyces naeslundii		1.59	0.26			
37512140	**Very good identification**					**37512140**
Propionibacterium propionicum		100.0	0.57	IND 1		
37512141	**Good identification**					**37512141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		100.0	0.41	IND 1		
37512142	**Very good identification**					**37512142**
Propionibacterium propionicum		100.0	0.60	IND 1		
37512143	**Good identification**					**37512143**
Propionibacterium propionicum		100.0	0.44	IND 1		
37513001	**Good identification**					**37513001**
Actinomyces naeslundii		100.0	0.42	IND 1		
37513003	**Low discrimination**					**37513003**
Actinomyces naeslundii		56.28	0.42	IND 1		
Actinomyces viscosus		43.72	0.16			
37513020	**Good identification**					**37513020**
Actinomyces naeslundii		100.0	0.39	IND 1		
37513021	**Good identification**					**37513021**
Actinomyces naeslundii		100.0	0.42	IND 1		
37513022	**Good identification**					**37513022**
Actinomyces naeslundii		100.0	0.39	IND 1		
37513023	**Good identification**					**37513023**
Actinomyces naeslundii		100.0	0.42	IND 1		
37513100	**Good identification**					**37513100**
Propionibacterium propionicum		94.14	0.57	IND 1		
Actinomyces naeslundii		5.86	0.46			
37513101	**Low discrimination**					**37513101**
Propionibacterium propionicum		54.33	0.41	IND 1		
Actinomyces naeslundii		45.67	0.49	URE 10		
37513102	**Low discrimination**					**37513102**
Propionibacterium propionicum		88.33	0.60	IND 1		
Actinomyces viscosus		8.01	0.28			
Actinomyces naeslundii		3.67	0.46			
37513103	**Low discrimination**					**37513103**
Propionibacterium propionicum		54.58	0.44	IND 1		
Actinomyces naeslundii		30.58	0.49	URE 10		
Actinomyces viscosus		14.84	0.20			
37513120	**Low discrimination**					**37513120**
Actinomyces naeslundii		75.58	0.46	IND 1		
Propionibacterium propionicum		12.26	0.23			
Clostridium perfringens		6.80	0.06			
37513121	**Good identification**					**37513121**
Actinomyces naeslundii		100.0	0.49	IND 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
37513122	**Low discrimination**						**37513122**
					GRA	CAT	
Actinomyces naeslundii		75.29	0.46	IND 1	99	25	
Propionibacterium propionicum		18.32	0.26		99	1	
Capnocytophaga ochracea		3.21	-.03		1	90	
37513123	**Good identification**						**37513123**
Actinomyces naeslundii		100.0	0.49	IND 1			
37513140	**Very good identification**						**37513140**
Propionibacterium propionicum		98.91	0.57	IND 1			
Actinomyces naeslundii		1.09	0.34				
37513141	**Low discrimination**						**37513141**
Propionibacterium propionicum		87.08	0.41	IND 1			
Actinomyces naeslundii		12.92	0.37	URE 10			
37513142	**Very good identification**						**37513142**
Propionibacterium propionicum		99.27	0.60	IND 1			
Actinomyces naeslundii		0.73	0.34				
37513143	**Acceptable identification**						**37513143**
Propionibacterium propionicum		91.00	0.44	IND 1			
Actinomyces naeslundii		9.00	0.37	URE 10			
37513161	**Good identification**						**37513161**
Actinomyces naeslundii		100.0	0.37	IND 1			
37513163	**Good identification**						**37513163**
Actinomyces naeslundii		100.0	0.37	IND 1			
37530002	**Good identification**						**37530002**
Actinomyces viscosus		99.24	0.42	IND 1			
Propionibacterium granulosum		0.76	0.15	TRE 7			
37530102	**Good identification**						**37530102**
Actinomyces viscosus		99.10	0.47	IND 1			
Propionibacterium propionicum		0.74	0.24	TRE 7			
Actinomyces naeslundii		0.16	0.23				
37530103	**Good identification**						**37530103**
Actinomyces viscosus		99.30	0.38	IND 1			
Actinomyces naeslundii		0.70	0.26	TRE 7			
37531001	**Good identification**						**37531001**
Actinomyces naeslundii		100.0	0.42	IND 1			
37531002	**Good identification**						**37531002**
Actinomyces viscosus		95.16	0.39	IND 1			
Actinomyces naeslundii		3.65	0.39	TRE 7			
Propionibacterium granulosum		1.19	0.15				
37531021	**Good identification**						**37531021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	IND 1		
37531023	**Good identification**					**37531023**
Actinomyces naeslundii		100.0	0.42	IND 1		
37531100	**Low discrimination**					**37531100**
Actinomyces naeslundii		53.12	0.46	IND 1		
Clostridium perfringens		27.17	0.19			
Actinomyces viscosus		8.73	0.09			
37531101	**Low discrimination**					**37531101**
					SPO	
Actinomyces naeslundii		84.17	0.49	IND 1	1	
Clostridium chauvoei		12.76	0.12		99	
Actinomyces viscosus		3.07	0.01		1	
37531102	**Acceptable identification**					**37531102**
Actinomyces viscosus		91.87	0.43	IND 1		
Actinomyces naeslundii		5.65	0.46	TRE 7		
Clostridium perfringens		1.36	0.14			
37531120	**Acceptable identification**					**37531120**
					GRA	
Clostridium perfringens		93.92	0.45	IND 1	99	
Actinomyces naeslundii		5.68	0.46		99	
Prevotella melaninogenica		0.40	-.03		1	
37531121	**Low discrimination**					**37531121**
					SPO	
Actinomyces naeslundii		86.07	0.49	IND 1	1	
Clostridium perfringens		9.59	0.11		84	
Clostridium chauvoei		4.35	0.04		99	
37531122	**Low discrimination**					**37531122**
Clostridium perfringens		86.57	0.39	IND 1		
Actinomyces naeslundii		11.13	0.46			
Actinomyces viscosus		1.83	0.09			
37531123	**Acceptable identification**					**37531123**
Actinomyces naeslundii		91.83	0.49	IND 1		
Clostridium perfringens		4.81	0.05			
Actinomyces viscosus		3.35	0.01			
37531141	**Good identification**					**37531141**
Actinomyces naeslundii		100.0	0.37	IND 1		
37531143	**Low discrimination**					**37531143**
Actinomyces naeslundii		82.86	0.37	IND 1		
Actinomyces viscosus		17.15	0.01			
37531161	**Good identification**					**37531161**
Actinomyces naeslundii		100.0	0.37	IND 1		
37531163	**Good identification**					**37531163**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.37	IND 1		
37532100	**Good identification**					**37532100**
	Propionibacterium propionicum	99.39	0.55	IND 1		
	Actinomyces naeslundii	0.61	0.28			
37532101	**Acceptable identification**					**37532101**
	Propionibacterium propionicum	92.33	0.39	IND 1		
	Actinomyces naeslundii	7.67	0.31	URE 10		
37532102	**Very good identification**					**37532102**
	Propionibacterium propionicum	98.26	0.58	IND 1		
	Actinomyces viscosus	1.34	0.12			
	Actinomyces naeslundii	0.40	0.28			
37532103	**Acceptable identification**					**37532103**
	Propionibacterium propionicum	91.23	0.42	IND 1		
	Actinomyces naeslundii	5.05	0.31	URE 10		
	Actinomyces viscosus	3.72	0.04			
37532140	**Good identification**					**37532140**
	Propionibacterium propionicum	100.0	0.55	IND 1		
37532141	**Good identification**					**37532141**
	Propionibacterium propionicum	100.0	0.39	IND 1		
37532142	**Very good identification**					**37532142**
	Propionibacterium propionicum	100.0	0.58	IND 1		
37532143	**Good identification**					**37532143**
	Propionibacterium propionicum	100.0	0.42	IND 1		
37533000	**Low discrimination**					**37533000**
					CAT	
	Actinomyces naeslundii	58.77	0.44	IND 1	25	
	Propionibacterium granulosum	29.79	0.24		90	
	Propionibacterium propionicum	11.44	0.21		1	
37533001	**Good identification**					**37533001**
	Actinomyces naeslundii	100.0	0.47	IND 1		
37533002	**Low discrimination**					**37533002**
	Actinomyces naeslundii	64.34	0.44	IND 1		
	Propionibacterium propionicum	18.78	0.24			
	Actinomyces viscosus	8.72	0.05			
37533003	**Good identification**					**37533003**
	Actinomyces naeslundii	100.0	0.47	IND 1		
37533020	**Low discrimination**					**37533020**
	Actinomyces naeslundii	88.76	0.44	IND 1		
	Propionibacterium granulosum	11.24	0.13			
37533021	**Good identification**					**37533021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	IND 1		
37533022	**Good identification**					**37533022**
Actinomyces naeslundii		100.0	0.44	IND 1		
37533023	**Good identification**					**37533023**
Actinomyces naeslundii		100.0	0.47	IND 1		
37533100	**Low discrimination**					**37533100**
Propionibacterium propionicum		87.13	0.55	IND 1		
Actinomyces naeslundii		12.87	0.51			
37533101	**Low discrimination**					**37533101**
Actinomyces naeslundii		66.60	0.54	IND 1		
Propionibacterium propionicum		33.40	0.39			
37533102	**Good identification**					**37533102**
Propionibacterium propionicum		90.35	0.58	IND 1		
Actinomyces naeslundii		8.90	0.51			
Actinomyces viscosus		0.75	0.09			
37533103	**Low discrimination**					**37533103**
Actinomyces naeslundii		56.46	0.54	IND 1		
Propionibacterium propionicum		42.47	0.42			
Actinomyces viscosus		1.06	0.01			
37533120	**Low discrimination**					**37533120**
Actinomyces naeslundii		86.62	0.51	IND 1		
Clostridium perfringens		7.45	0.11			
Propionibacterium propionicum		5.92	0.21			
37533121	**Good identification**					**37533121**
Actinomyces naeslundii		100.0	0.54	IND 1		
37533122	**Low discrimination**					**37533122**
Actinomyces naeslundii		87.48	0.51	IND 1		
Propionibacterium propionicum		8.97	0.24			
Clostridium perfringens		3.54	0.05			
37533123	**Good identification**					**37533123**
Actinomyces naeslundii		100.0	0.54	IND 1		
37533140	**Good identification**					**37533140**
Propionibacterium propionicum		97.46	0.55	IND 1		
Actinomyces naeslundii		2.54	0.38			
37533141	**Low discrimination**					**37533141**
Propionibacterium propionicum		73.97	0.39	IND 1		
Actinomyces naeslundii		26.03	0.41	URE 10		
37533142	**Very good identification**					**37533142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium propionicum		98.29	0.58	IND 1		
Actinomyces naeslundii		1.71	0.38			
37533143	**Low discrimination**					**37533143**
Propionibacterium propionicum		81.00	0.42	IND 1		
Actinomyces naeslundii		19.00	0.41	URE 10		
37533160	**Low discrimination**					**37533160**
Actinomyces naeslundii		72.07	0.38	IND 1		
Propionibacterium propionicum		27.93	0.21			
37533161	**Good identification**					**37533161**
Actinomyces naeslundii		100.0	0.41	IND 1		
37533162	**Low discrimination**					**37533162**
Actinomyces naeslundii		63.24	0.38	IND 1		
Propionibacterium propionicum		36.76	0.24			
37533163	**Good identification**					**37533163**
Actinomyces naeslundii		100.0	0.41	IND 1		
37535722	**Good identification**					**37535722**
Bacteroides thetaiotaomicron		98.99	0.42	NIT 1		
Parabacteroides distasonis		0.69	0.21	ESL 92		
Bacteroides ovatus		0.33	0.00			
37535732	**Good identification**					**37535732**
Bacteroides thetaiotaomicron		98.50	0.35	NIT 1		
Parabacteroides distasonis		0.95	0.17	ESL 92		
Bacteroides ovatus		0.56	-.03			
37550002	**Good identification**					**37550002**
Actinomyces viscosus		100.0	0.52	IND 1		
37550003	**Good identification**					**37550003**
Actinomyces viscosus		100.0	0.44	IND 1		
37550102	**Good identification**					**37550102**
Actinomyces viscosus		100.0	0.56	IND 1		
37550103	**Good identification**					**37550103**
Actinomyces viscosus		99.86	0.48	IND 1		
Actinomyces naeslundii		0.14	0.23			
37551001	**Low discrimination**					**37551001**
Actinomyces naeslundii		76.78	0.39	IND 1		
Actinomyces viscosus		23.22	0.06			
37551002	**Good identification**					**37551002**
Actinomyces viscosus		99.26	0.48	IND 1		
Actinomyces naeslundii		0.74	0.36			
37551003	**Good identification**					**37551003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		96.77	0.40	IND 1		
Actinomyces naeslundii		3.23	0.39			
37551020	**Low discrimination**					**37551020**
Actinomyces naeslundii		82.58	0.36	IND 1		
Clostridium perfringens		17.42	0.02			
37551021	**Good identification**					**37551021**
Actinomyces naeslundii		100.0	0.39	IND 1		
37551023	**Low discrimination**					**37551023**
Actinomyces naeslundii		76.78	0.39	IND 1		
Actinomyces viscosus		23.22	0.06			
37551100	**Low discrimination**					**37551100**
Actinomyces naeslundii		54.05	0.44	IND 1		
Actinomyces viscosus		45.95	0.18			
37551101	**Low discrimination**					**37551101**
Actinomyces naeslundii		84.11	0.47	IND 1		
Actinomyces viscosus		15.89	0.10			
37551102	**Good identification**					**37551102**
					GRA	
Actinomyces viscosus		98.63	0.52	IND 1	99	
Actinomyces naeslundii		1.17	0.44		99	
Capnocytophaga ochracea		0.20	0.05		1	
37551103	**Acceptable identification**					**37551103**
Actinomyces viscosus		94.92	0.44	IND 1		
Actinomyces naeslundii		5.08	0.47			
37551121	**Good identification**					**37551121**
Actinomyces naeslundii		100.0	0.47	IND 1		
37551123	**Low discrimination**					**37551123**
Actinomyces naeslundii		84.11	0.47	IND 1		
Actinomyces viscosus		15.89	0.10			
37553000	**Good identification**					**37553000**
Actinomyces naeslundii		100.0	0.41	IND 1		
37553001	**Good identification**					**37553001**
Actinomyces naeslundii		100.0	0.44	IND 1		
37553002	**Low discrimination**					**37553002**
Actinomyces naeslundii		58.79	0.41	IND 1		
Actinomyces viscosus		41.21	0.14			
37553003	**Low discrimination**					**37553003**
Actinomyces naeslundii		86.52	0.44	IND 1		
Actinomyces viscosus		13.48	0.06			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37553020	**Good identification**					**37553020**
Actinomyces naeslundii		100.0	0.41	IND 1		
37553021	**Good identification**					**37553021**
Actinomyces naeslundii		100.0	0.44	IND 1		
37553022	**Good identification**					**37553022**
Actinomyces naeslundii		100.0	0.41	IND 1		
37553023	**Good identification**					**37553023**
Actinomyces naeslundii		100.0	0.44	IND 1		
37553100	**Low discrimination**					**37553100**
Actinomyces naeslundii		89.10	0.49	IND 1		
Propionibacterium propionicum		10.91	0.23			
37553101	**Good identification**					**37553101**
Actinomyces naeslundii		100.0	0.52	IND 1		
37553102	**Low discrimination**					**37553102**
Actinomyces naeslundii		61.67	0.49	IND 1		
Actinomyces viscosus		27.01	0.18			
Propionibacterium propionicum		11.32	0.26			
37553103	**Acceptable identification**					**37553103**
Actinomyces naeslundii		91.13	0.52	IND 1		
Actinomyces viscosus		8.87	0.10			
37553120	**Good identification**					**37553120**
Actinomyces naeslundii		100.0	0.49	IND 1		
37553121	**Good identification**					**37553121**
Actinomyces naeslundii		100.0	0.52	IND 1		
37553122	**Good identification**					**37553122**
Actinomyces naeslundii		100.0	0.49	IND 1		
37553123	**Good identification**					**37553123**
Actinomyces naeslundii		100.0	0.52	IND 1		
37553140	**Low discrimination**					**37553140**
Actinomyces naeslundii		59.05	0.36	IND 1		
Propionibacterium propionicum		40.95	0.23			
37553141	**Good identification**					**37553141**
Actinomyces naeslundii		100.0	0.39	IND 1		
37553143	**Good identification**					**37553143**
Actinomyces naeslundii		100.0	0.39	IND 1		
37553160	**Good identification**					**37553160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	IND 1		
37553161	**Good identification**					**37553161**
Actinomyces naeslundii		100.0	0.39	IND 1		
37553162	**Good identification**					**37553162**
Actinomyces naeslundii		100.0	0.36	IND 1		
37553163	**Good identification**					**37553163**
Actinomyces naeslundii		100.0	0.39	IND 1		
37570102	**Good identification**					**37570102**
Actinomyces viscosus		99.22	0.37	IND 1		
Actinomyces naeslundii		0.78	0.25	TRE 7		
37571000	**Good identification**					**37571000**
Actinomyces naeslundii		100.0	0.41	IND 1		
37571001	**Good identification**					**37571001**
Actinomyces naeslundii		100.0	0.44	IND 1		
37571020	**Low discrimination**					**37571020**
Actinomyces naeslundii		83.22	0.41	IND 1		
Clostridium perfringens		16.79	0.07			
37571021	**Good identification**					**37571021**
Actinomyces naeslundii		100.0	0.44	IND 1		
37571022	**Acceptable identification**					**37571022**
Actinomyces naeslundii		91.33	0.41	IND 1		
Clostridium perfringens		8.67	0.01			
37571023	**Good identification**					**37571023**
Actinomyces naeslundii		100.0	0.44	IND 1		
37571100	**Good identification**					**37571100**
Actinomyces naeslundii		96.81	0.49	IND 1		
Actinomyces viscosus		3.19	-.01			
37571101	**Good identification**					**37571101**
Actinomyces naeslundii		100.0	0.52	IND 1		
37571103	**Low discrimination**					**37571103**
Actinomyces naeslundii		57.97	0.52	IND 1		
Actinomyces viscosus		42.03	0.25			
37571120	**Low discrimination**					**37571120**
Actinomyces naeslundii		65.80	0.49	IND 1		
Clostridium perfringens		34.20	0.21			
37571121	**Good identification**					**37571121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	IND 1		
37571122	**Low discrimination**					**37571122**
Actinomyces naeslundii		78.27	0.49	IND 1		
Clostridium perfringens		19.15	0.16			
Actinomyces viscosus		2.58	-.01			
37571123	**Good identification**					**37571123**
Actinomyces naeslundii		100.0	0.52	IND 1		
37571140	**Good identification**					**37571140**
Actinomyces naeslundii		100.0	0.36	IND 1		
37571141	**Good identification**					**37571141**
Actinomyces naeslundii		100.0	0.39	IND 1		
37571142	**Low discrimination**					**37571142**
Actinomyces naeslundii		84.26	0.36	IND 1		
Actinomyces viscosus		15.74	-.01			
37571143	**Good identification**					**37571143**
Actinomyces naeslundii		100.0	0.39	IND 1		
37571160	**Low discrimination**					**37571160**
Actinomyces naeslundii		64.06	0.36	IND 1		
Clostridium perfringens		35.94	0.09			
37571161	**Good identification**					**37571161**
Actinomyces naeslundii		100.0	0.39	IND 1		
37571162	**Low discrimination**					**37571162**
Actinomyces naeslundii		79.12	0.36	IND 1		
Clostridium perfringens		20.89	0.03			
37571163	**Good identification**					**37571163**
Actinomyces naeslundii		100.0	0.39	IND 1		
37573000	**Good identification**					**37573000**
Actinomyces naeslundii		100.0	0.46	IND 1		
37573001	**Good identification**					**37573001**
Actinomyces naeslundii		100.0	0.49	IND 1		
37573002	**Good identification**					**37573002**
Actinomyces naeslundii		100.0	0.46	IND 1		
37573003	**Good identification**					**37573003**
Actinomyces naeslundii		100.0	0.49	IND 1		
37573020	**Good identification**					**37573020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	IND 1		
37573021	**Good identification**					**37573021**
Actinomyces naeslundii		100.0	0.49	IND 1		
37573022	**Good identification**					**37573022**
Actinomyces naeslundii		100.0	0.46	IND 1		
37573023	**Good identification**					**37573023**
Actinomyces naeslundii		100.0	0.49	IND 1		
37573041	**Good identification**					**37573041**
Actinomyces naeslundii		100.0	0.36	IND 1		
37573043	**Good identification**					**37573043**
Actinomyces naeslundii		100.0	0.36	IND 1		
37573061	**Good identification**					**37573061**
Actinomyces naeslundii		100.0	0.36	IND 1		
37573063	**Good identification**					**37573063**
Actinomyces naeslundii		100.0	0.36	IND 1		
37573100	**Acceptable identification**					**37573100**
Actinomyces naeslundii		92.93	0.53	IND 1		
Propionibacterium propionicum		4.79	0.21			
Actinomyces israelii		2.28	0.17			
37573101	**Very good identification**					**37573101**
Actinomyces naeslundii		100.0	0.56	IND 1		
37573102	**Low discrimination**					**37573102**
Actinomyces naeslundii		88.40	0.53	IND 1		
Propionibacterium propionicum		6.84	0.24			
Actinomyces israelii		3.25	0.20			
37573103	**Very good identification**					**37573103**
Actinomyces naeslundii		100.0	0.56	IND 1		
37573120	**Good identification**					**37573120**
Actinomyces naeslundii		100.0	0.53	IND 1		
37573121	**Very good identification**					**37573121**
Actinomyces naeslundii		100.0	0.56	IND 1		
37573122	**Good identification**					**37573122**
Actinomyces naeslundii		100.0	0.53	IND 1		
37573123	**Very good identification**					**37573123**
Actinomyces naeslundii		100.0	0.56	IND 1		
37573140	**Low discrimination**					**37573140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		77.38	0.41	IND 1		
Propionibacterium propionicum		22.62	0.21			
37573141	**Good identification**					**37573141**
Actinomyces naeslundii		100.0	0.44	IND 1		
37573142	**Low discrimination**					**37573142**
Actinomyces naeslundii		69.52	0.41	IND 1		
Propionibacterium propionicum		30.48	0.24			
37573143	**Good identification**					**37573143**
Actinomyces naeslundii		100.0	0.44	IND 1		
37573160	**Good identification**					**37573160**
Actinomyces naeslundii		100.0	0.41	IND 1		
37573161	**Good identification**					**37573161**
Actinomyces naeslundii		100.0	0.44	IND 1		
37573162	**Good identification**					**37573162**
Actinomyces naeslundii		100.0	0.41	IND 1		
37573163	**Good identification**					**37573163**
Actinomyces naeslundii		100.0	0.44	IND 1		
37575722	**Low discrimination**					**37575722**
Bacteroides thetaiotaomicron		86.55	0.48	NIT 1		
Bacteroides ovatus		13.30	0.34	ESL 92		
Parabacteroides distasonis		0.15	0.17			
37575732	**Low discrimination**					**37575732**
Bacteroides thetaiotaomicron		78.81	0.41	NIT 1		
Bacteroides ovatus		20.76	0.31	ESL 92		
Clostridium clostridioforme		0.24	0.08			
37577102	**Good identification**					**37577102**
Actinomyces israelii		100.0	0.38	IND 1		
37577502	**Good identification**					**37577502**
Actinomyces israelii		100.0	0.36	IND 1		
37613100	**Good identification**					**37613100**
Actinomyces naeslundii		100.0	0.35	IND 1		
37613101	**Good identification**					**37613101**
Actinomyces naeslundii		100.0	0.38	IND 1		
37613102	**Good identification**					**37613102**
Actinomyces naeslundii		100.0	0.35	IND 1		
37613103	**Good identification**					**37613103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	IND 1		
37613120	**Good identification**					**37613120**
Actinomyces naeslundii		100.0	0.35	IND 1		
37613121	**Good identification**					**37613121**
Actinomyces naeslundii		100.0	0.38	IND 1		
37613122	**Good identification**					**37613122**
Actinomyces naeslundii		100.0	0.35	IND 1		
37613123	**Good identification**					**37613123**
Actinomyces naeslundii		100.0	0.38	IND 1		
37631100	**Low discrimination**					**37631100**
					GRA	
Actinomyces naeslundii		63.37	0.35	IND 1	99	
Fusobacterium mortiferum		36.63	0.20	MLT 94	1	
37631101	**Good identification**					**37631101**
Actinomyces naeslundii		100.0	0.38	IND 1		
37631102	**Low discrimination**					**37631102**
Actinomyces naeslundii		83.81	0.35	IND 1		
Actinomyces viscosus		16.19	-.01	MLT 94		
37631103	**Good identification**					**37631103**
Actinomyces naeslundii		100.0	0.38	IND 1		
37631120	**Low discrimination**					**37631120**
Actinomyces naeslundii		84.43	0.35	IND 1		
Clostridium perfringens		15.58	0.01	MLT 94		
37631121	**Good identification**					**37631121**
Actinomyces naeslundii		100.0	0.38	IND 1		
37631122	**Good identification**					**37631122**
Actinomyces naeslundii		100.0	0.35	IND 1		
37631123	**Good identification**					**37631123**
Actinomyces naeslundii		100.0	0.38	IND 1		
37633001	**Good identification**					**37633001**
Actinomyces naeslundii		100.0	0.36	IND 1		
37633003	**Good identification**					**37633003**
Actinomyces naeslundii		100.0	0.36	IND 1		
37633021	**Good identification**					**37633021**
Actinomyces naeslundii		100.0	0.36	IND 1		
37633023	**Good identification**					**37633023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	IND 1		
37633100	**Good identification**					**37633100**
Actinomyces naeslundii		100.0	0.40	IND 1		
37633101	**Good identification**					**37633101**
Actinomyces naeslundii		100.0	0.43	IND 1		
37633102	**Good identification**					**37633102**
Actinomyces naeslundii		100.0	0.40	IND 1		
37633103	**Good identification**					**37633103**
Actinomyces naeslundii		100.0	0.43	IND 1		
37633120	**Good identification**					**37633120**
Actinomyces naeslundii		100.0	0.40	IND 1		
37633121	**Good identification**					**37633121**
Actinomyces naeslundii		100.0	0.43	IND 1		
37633122	**Good identification**					**37633122**
Actinomyces naeslundii		100.0	0.40	IND 1		
37633123	**Good identification**					**37633123**
Actinomyces naeslundii		100.0	0.43	IND 1		
37651101	**Good identification**					**37651101**
Actinomyces naeslundii		100.0	0.36	IND 1		
37651103	**Low discrimination**					**37651103**
Actinomyces naeslundii		81.84	0.36	IND 1		
Actinomyces viscosus		18.16	0.00	MLT 94		
37651121	**Good identification**					**37651121**
Actinomyces naeslundii		100.0	0.36	IND 1		
37651123	**Good identification**					**37651123**
Actinomyces naeslundii		100.0	0.36	IND 1		
37653100	**Good identification**					**37653100**
Actinomyces naeslundii		100.0	0.37	IND 1		
37653101	**Good identification**					**37653101**
Actinomyces naeslundii		100.0	0.41	IND 1		
37653102	**Good identification**					**37653102**
Actinomyces naeslundii		100.0	0.37	IND 1		
37653103	**Good identification**					**37653103**
Actinomyces naeslundii		100.0	0.41	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37653120	**Good identification**					**37653120**
Actinomyces naeslundii		100.0	0.37	IND 1		
37653121	**Good identification**					**37653121**
Actinomyces naeslundii		100.0	0.41	IND 1		
37653122	**Good identification**					**37653122**
Actinomyces naeslundii		100.0	0.37	IND 1		
37653123	**Good identification**					**37653123**
Actinomyces naeslundii		100.0	0.41	IND 1		
37671100	**Low discrimination**					**37671100**
					GRA	
Actinomyces naeslundii		87.31	0.37	IND 1	99	
Fusobacterium mortiferum		12.69	0.12	MLT 94	1	
37671101	**Good identification**					**37671101**
Actinomyces naeslundii		100.0	0.41	IND 1		
37671102	**Low discrimination**					**37671102**
Actinomyces naeslundii		86.95	0.37	IND 1		
Lactobacillus acidophilus		13.05	-.01	MLT 94		
37671103	**Good identification**					**37671103**
Actinomyces naeslundii		100.0	0.41	IND 1		
37671120	**Good identification**					**37671120**
Actinomyces naeslundii		100.0	0.37	IND 1		
37671121	**Good identification**					**37671121**
Actinomyces naeslundii		100.0	0.41	IND 1		
37671122	**Low discrimination**					**37671122**
Actinomyces naeslundii		86.95	0.37	IND 1		
Lactobacillus acidophilus		13.05	-.01	MLT 94		
37671123	**Good identification**					**37671123**
Actinomyces naeslundii		100.0	0.41	IND 1		
37673001	**Good identification**					**37673001**
Actinomyces naeslundii		100.0	0.38	IND 1		
37673003	**Good identification**					**37673003**
Actinomyces naeslundii		100.0	0.38	IND 1		
37673021	**Good identification**					**37673021**
Actinomyces naeslundii		100.0	0.38	IND 1		
37673023	**Good identification**					**37673023**
Actinomyces naeslundii		100.0	0.38	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37673100	**Good identification**					**37673100**
Actinomyces naeslundii		100.0	0.42	IND 1		
37673101	**Good identification**					**37673101**
Actinomyces naeslundii		100.0	0.45	IND 1		
37673102	**Good identification**					**37673102**
Actinomyces naeslundii		100.0	0.42	IND 1		
37673103	**Good identification**					**37673103**
Actinomyces naeslundii		100.0	0.45	IND 1		
37673120	**Good identification**					**37673120**
Actinomyces naeslundii		100.0	0.42	IND 1		
37673121	**Good identification**					**37673121**
Actinomyces naeslundii		100.0	0.45	IND 1		
37673122	**Good identification**					**37673122**
Actinomyces naeslundii		100.0	0.42	IND 1		
37673123	**Good identification**					**37673123**
Actinomyces naeslundii		100.0	0.45	IND 1		
37703101	**Good identification**					**37703101**
Actinomyces naeslundii		100.0	0.37	IND 1		
37703103	**Good identification**					**37703103**
Actinomyces naeslundii		100.0	0.37	IND 1		
37703121	**Good identification**					**37703121**
Actinomyces naeslundii		100.0	0.37	IND 1		
37703123	**Good identification**					**37703123**
Actinomyces naeslundii		100.0	0.37	IND 1		
37710002	**Good identification**					**37710002**
Actinomyces viscosus		99.75	0.52	IND 1		
Actinomyces odontolyticus		0.25	0.14			
37710003	**Good identification**					**37710003**
Actinomyces viscosus		99.77	0.44	IND 1		
Actinomyces naeslundii		0.23	0.23			
37710102	**Good identification**					**37710102**
Actinomyces viscosus		99.87	0.56	IND 1		
Actinomyces naeslundii		0.08	0.27			
Actinomyces odontolyticus		0.05	0.06			
37710103	**Good identification**					**37710103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		99.64	0.48	IND 1		
Actinomyces naeslundii		0.36	0.30			
37711001	**Low discrimination**					**37711001**
Actinomyces naeslundii		89.84	0.46	IND 1		
Actinomyces viscosus		10.16	0.06			
37711002	**Good identification**					**37711002**
Actinomyces viscosus		97.79	0.48	IND 1		
Actinomyces naeslundii		1.94	0.43			
Actinomyces odontolyticus		0.27	0.11			
37711003	**Acceptable identification**					**37711003**
Actinomyces viscosus		91.80	0.40	IND 1		
Actinomyces naeslundii		8.20	0.46			
37711020	**Low discrimination**					**37711020**
Actinomyces naeslundii		67.89	0.43	IND 1		
Clostridium perfringens		32.11	0.15			
37711021	**Good identification**					**37711021**
Actinomyces naeslundii		100.0	0.46	IND 1		
37711022	**Low discrimination**					**37711022**
Actinomyces naeslundii		57.76	0.43	IND 1		
Actinomyces viscosus		29.39	0.14			
Clostridium perfringens		12.86	0.10			
37711023	**Low discrimination**					**37711023**
Actinomyces naeslundii		89.84	0.46	IND 1		
Actinomyces viscosus		10.16	0.06			
37711100	**Low discrimination**					**37711100**
Actinomyces naeslundii		70.91	0.51	IND 1		
Actinomyces viscosus		22.54	0.18			
Clostridium perfringens		2.67	0.04			
37711101	**Acceptable identification**					**37711101**
Actinomyces naeslundii		93.40	0.54	IND 1		
Actinomyces viscosus		6.60	0.10			
37711102	**Good identification**					**37711102**
Actinomyces viscosus		96.86	0.52	IND 1		
Actinomyces naeslundii		3.08	0.51			
Bifidobacterium longum subsp. infantis		0.07	-.07			
37711103	**Low discrimination**					**37711103**
Actinomyces viscosus		87.49	0.44	IND 1		
Actinomyces naeslundii		12.51	0.54			
37711121	**Good identification**					**37711121**
Actinomyces naeslundii		100.0	0.54	IND 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
37711122	**Low discrimination**					**37711122**
Actinomyces naeslundii		48.25	0.51	IND 1		
Clostridium perfringens		27.67	0.24			
Actinomyces viscosus		15.34	0.18			
37711123	**Acceptable identification**					**37711123**
Actinomyces naeslundii		93.40	0.54	IND 1		
Actinomyces viscosus		6.60	0.10			
37711140	**Good identification**					**37711140**
Actinomyces naeslundii		100.0	0.38	IND 1		
37711141	**Good identification**					**37711141**
Actinomyces naeslundii		100.0	0.41	IND 1		
37711143	**Low discrimination**					**37711143**
Actinomyces naeslundii		71.42	0.41	IND 1		
Actinomyces viscosus		28.58	0.10			
37711161	**Good identification**					**37711161**
Actinomyces naeslundii		100.0	0.41	IND 1		
37711162	**Low discrimination**					**37711162**
Actinomyces naeslundii		61.76	0.38	IND 1		
Clostridium perfringens		38.24	0.12			
37711163	**Good identification**					**37711163**
Actinomyces naeslundii		100.0	0.41	IND 1		
37712101	**Good identification**					**37712101**
Actinomyces naeslundii		100.0	0.35	IND 1		
37712121	**Good identification**					**37712121**
Actinomyces naeslundii		100.0	0.35	IND 1		
37712123	**Good identification**					**37712123**
Actinomyces naeslundii		100.0	0.35	IND 1		
37713000	**Good identification**					**37713000**
Actinomyces naeslundii		100.0	0.48	IND 1		
37713001	**Good identification**					**37713001**
Actinomyces naeslundii		100.0	0.51	IND 1		
37713002	**Low discrimination**					**37713002**
Actinomyces naeslundii		79.23	0.48	IND 1		
Actinomyces viscosus		20.77	0.14			
37713003	**Acceptable identification**					**37713003**
Actinomyces naeslundii		94.50	0.51	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		5.51	0.06			
37713020	**Good identification**					**37713020**
Actinomyces naeslundii		100.0	0.48	IND 1		
37713021	**Good identification**					**37713021**
Actinomyces naeslundii		100.0	0.51	IND 1		
37713022	**Good identification**					**37713022**
Actinomyces naeslundii		100.0	0.48	IND 1		
37713023	**Good identification**					**37713023**
Actinomyces naeslundii		100.0	0.51	IND 1		
37713040	**Good identification**					**37713040**
Actinomyces naeslundii		100.0	0.35	IND 1		
37713041	**Good identification**					**37713041**
Actinomyces naeslundii		100.0	0.38	IND 1		
37713042	**Good identification**					**37713042**
Actinomyces naeslundii		100.0	0.35	IND 1		
37713043	**Good identification**					**37713043**
Actinomyces naeslundii		100.0	0.38	IND 1		
37713060	**Good identification**					**37713060**
Actinomyces naeslundii		100.0	0.35	IND 1		
37713061	**Good identification**					**37713061**
Actinomyces naeslundii		100.0	0.38	IND 1		
37713062	**Good identification**					**37713062**
Actinomyces naeslundii		100.0	0.35	IND 1		
37713063	**Good identification**					**37713063**
Actinomyces naeslundii		100.0	0.38	IND 1		
37713100	**Good identification**					**37713100**
Actinomyces naeslundii		95.62	0.56	IND 1		
Propionibacterium propionicum		4.38	0.23			
37713101	**Very good identification**					**37713101**
Actinomyces naeslundii		100.0	0.59	IND 1		
37713102	**Low discrimination**					**37713102**
Actinomyces naeslundii		81.15	0.56	IND 1		
Actinomyces viscosus		13.29	0.18			
Propionibacterium propionicum		5.57	0.26			
37713103	**Very good identification**					**37713103**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.49	0.59	IND 1		
Actinomyces viscosus		3.51	0.10			
37713111	**Good identification**					**37713111**
Actinomyces naeslundii		100.0	0.37	IND 1		
37713113	**Good identification**					**37713113**
Actinomyces naeslundii		100.0	0.37	IND 1		
37713120	**Good identification**					**37713120**
Actinomyces naeslundii		100.0	0.56	IND 1		
37713121	**Very good identification**					**37713121**
Actinomyces naeslundii		100.0	0.59	IND 1		
37713122	**Good identification**					**37713122**
Actinomyces naeslundii		100.0	0.56	IND 1		
37713123	**Very good identification**					**37713123**
Actinomyces naeslundii		100.0	0.59	IND 1		
37713131	**Good identification**					**37713131**
Actinomyces naeslundii		100.0	0.37	IND 1		
37713133	**Good identification**					**37713133**
Actinomyces naeslundii		100.0	0.37	IND 1		
37713140	**Low discrimination**					**37713140**
Actinomyces naeslundii		79.41	0.43	IND 1		
Propionibacterium propionicum		20.59	0.23			
37713141	**Good identification**					**37713141**
Actinomyces naeslundii		100.0	0.46	IND 1		
37713142	**Low discrimination**					**37713142**
Actinomyces naeslundii		72.00	0.43	IND 1		
Propionibacterium propionicum		28.00	0.26			
37713143	**Good identification**					**37713143**
Actinomyces naeslundii		100.0	0.46	IND 1		
37713160	**Good identification**					**37713160**
Actinomyces naeslundii		100.0	0.43	IND 1		
37713161	**Good identification**					**37713161**
Actinomyces naeslundii		100.0	0.46	IND 1		
37713162	**Good identification**					**37713162**
Actinomyces naeslundii		100.0	0.43	IND 1		
37713163	**Good identification**					**37713163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	IND 1		
37721101	**Good identification**					**37721101**
Actinomyces naeslundii		100.0	0.37	IND 1		
37721103	**Low discrimination**					**37721103**
Actinomyces naeslundii		78.67	0.37	IND 1		
Actinomyces viscosus		21.33	0.03	FRU 95		
37721121	**Good identification**					**37721121**
Actinomyces naeslundii		100.0	0.37	IND 1		
37721123	**Good identification**					**37721123**
Actinomyces naeslundii		100.0	0.37	IND 1		
37723100	**Good identification**					**37723100**
Actinomyces naeslundii		100.0	0.39	IND 1		
37723101	**Good identification**					**37723101**
Actinomyces naeslundii		100.0	0.42	IND 1		
37723102	**Good identification**					**37723102**
Actinomyces naeslundii		100.0	0.39	IND 1		
37723103	**Good identification**					**37723103**
Actinomyces naeslundii		100.0	0.42	IND 1		
37723120	**Good identification**					**37723120**
Actinomyces naeslundii		100.0	0.39	IND 1		
37723121	**Good identification**					**37723121**
Actinomyces naeslundii		100.0	0.42	IND 1		
37723122	**Good identification**					**37723122**
Actinomyces naeslundii		100.0	0.39	IND 1		
37723123	**Good identification**					**37723123**
Actinomyces naeslundii		100.0	0.42	IND 1		
37730101	**Good identification**					**37730101**
Actinomyces naeslundii		100.0	0.35	IND 1		
37730102	**Good identification**					**37730102**
Actinomyces viscosus		97.95	0.37	IND 1		
Actinomyces naeslundii		2.05	0.32	TRE 7		
37730121	**Good identification**					**37730121**
Actinomyces naeslundii		100.0	0.35	IND 1		
37730123	**Good identification**					**37730123**
Actinomyces naeslundii		100.0	0.35	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37731000	**Good identification**					**37731000**
Actinomyces naeslundii		100.0	0.48	IND 1		
37731001	**Good identification**					**37731001**
Actinomyces naeslundii		100.0	0.51	IND 1		
37731003	**Low discrimination**					**37731003**
Actinomyces naeslundii		69.73	0.51	IND 1		
Actinomyces viscosus		30.27	0.21			
37731020	**Low discrimination**					**37731020**
Actinomyces naeslundii		68.84	0.48	IND 1		
Clostridium perfringens		31.16	0.20			
37731021	**Good identification**					**37731021**
Actinomyces naeslundii		100.0	0.51	IND 1		
37731022	**Low discrimination**					**37731022**
Actinomyces naeslundii		82.44	0.48	IND 1		
Clostridium perfringens		17.56	0.14			
37731023	**Good identification**					**37731023**
Actinomyces naeslundii		100.0	0.51	IND 1		
37731040	**Good identification**					**37731040**
Actinomyces naeslundii		100.0	0.35	IND 1		
37731041	**Good identification**					**37731041**
Actinomyces naeslundii		100.0	0.38	IND 1		
37731042	**Good identification**					**37731042**
Actinomyces naeslundii		100.0	0.35	IND 1		
37731043	**Good identification**					**37731043**
Actinomyces naeslundii		100.0	0.38	IND 1		
37731060	**Low discrimination**					**37731060**
Actinomyces naeslundii		67.18	0.35	IND 1		
Clostridium perfringens		32.82	0.08			
37731061	**Good identification**					**37731061**
Actinomyces naeslundii		100.0	0.38	IND 1		
37731062	**Low discrimination**					**37731062**
Actinomyces naeslundii		81.31	0.35	IND 1		
Clostridium perfringens		18.70	0.02			
37731063	**Good identification**					**37731063**
Actinomyces naeslundii		100.0	0.38	IND 1		
37731100	**Good identification**					**37731100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		95.39	0.56	IND 1		
Clostridium perfringens		3.44	0.09			
Actinomyces viscosus		1.18	-.01			
37731101	**Very good identification**					**37731101**
Actinomyces naeslundii		100.0	0.59	IND 1		
37731103	**Low discrimination**					**37731103**
Actinomyces naeslundii		78.67	0.59	IND 1		
Actinomyces viscosus		21.33	0.25			
37731111	**Good identification**					**37731111**
Actinomyces naeslundii		100.0	0.37	IND 1		
37731113	**Good identification**					**37731113**
Actinomyces naeslundii		100.0	0.37	IND 1		
37731121	**Very good identification**					**37731121**
Actinomyces naeslundii		99.22	0.59	IND 1		
Clostridium perfringens		0.78	0.01			
37731122	**Low discrimination**					**37731122**
Actinomyces naeslundii		64.06	0.56	IND 1		
Clostridium perfringens		35.15	0.29			
Actinomyces viscosus		0.79	-.01			
37731123	**Very good identification**					**37731123**
Actinomyces naeslundii		100.0	0.59	IND 1		
37731131	**Good identification**					**37731131**
Actinomyces naeslundii		100.0	0.37	IND 1		
37731133	**Good identification**					**37731133**
Actinomyces naeslundii		100.0	0.37	IND 1		
37731140	**Good identification**					**37731140**
Actinomyces naeslundii		100.0	0.43	IND 1		
37731141	**Good identification**					**37731141**
Actinomyces naeslundii		100.0	0.46	IND 1		
37731142	**Acceptable identification**					**37731142**
Actinomyces naeslundii		93.47	0.43	IND 1		
Actinomyces viscosus		6.53	-.01			
37731143	**Good identification**					**37731143**
Actinomyces naeslundii		100.0	0.46	IND 1		
37731161	**Good identification**					**37731161**
Actinomyces naeslundii		100.0	0.46	IND 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
37731162	**Low discrimination**					**37731162**
Actinomyces naeslundii		62.80	0.43	IND 1		
Clostridium perfringens		37.20	0.17			
37731163	**Good identification**					**37731163**
Actinomyces naeslundii		100.0	0.46	IND 1		
37732100	**Low discrimination**					**37732100**
Actinomyces naeslundii		68.36	0.37	IND 1		
Propionibacterium propionicum		31.64	0.21	GAL 96		
37732101	**Good identification**					**37732101**
Actinomyces naeslundii		100.0	0.40	IND 1		
37732102	**Low discrimination**					**37732102**
Actinomyces naeslundii		51.47	0.37	IND 1		
Propionibacterium propionicum		35.74	0.24	GAL 96		
Actinomyces viscosus		12.79	0.03			
37732103	**Good identification**					**37732103**
Actinomyces naeslundii		100.0	0.40	IND 1		
37732120	**Good identification**					**37732120**
Actinomyces naeslundii		100.0	0.37	IND 1		
37732121	**Good identification**					**37732121**
Actinomyces naeslundii		100.0	0.40	IND 1		
37732122	**Good identification**					**37732122**
Actinomyces naeslundii		100.0	0.37	IND 1		
37732123	**Good identification**					**37732123**
Actinomyces naeslundii		100.0	0.40	IND 1		
37733000	**Good identification**					**37733000**
Actinomyces naeslundii		100.0	0.53	IND 1		
37733001	**Good identification**					**37733001**
Actinomyces naeslundii		100.0	0.56	IND 1		
37733002	**Good identification**					**37733002**
Actinomyces naeslundii		100.0	0.53	IND 1		
37733003	**Good identification**					**37733003**
Actinomyces naeslundii		100.0	0.56	IND 1		
37733020	**Good identification**					**37733020**
Actinomyces naeslundii		100.0	0.53	IND 1		
37733021	**Good identification**					**37733021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.56	IND 1		
37733022	**Good identification**					**37733022**
Actinomyces naeslundii		100.0	0.53	IND 1		
37733023	**Good identification**					**37733023**
Actinomyces naeslundii		100.0	0.56	IND 1		
37733040	**Good identification**					**37733040**
Actinomyces naeslundii		100.0	0.40	IND 1		
37733041	**Good identification**					**37733041**
Actinomyces naeslundii		100.0	0.43	IND 1		
37733042	**Good identification**					**37733042**
Actinomyces naeslundii		100.0	0.40	IND 1		
37733043	**Good identification**					**37733043**
Actinomyces naeslundii		100.0	0.43	IND 1		
37733060	**Good identification**					**37733060**
Actinomyces naeslundii		100.0	0.40	IND 1		
37733061	**Good identification**					**37733061**
Actinomyces naeslundii		100.0	0.43	IND 1		
37733062	**Good identification**					**37733062**
Actinomyces naeslundii		100.0	0.40	IND 1		
37733063	**Good identification**					**37733063**
Actinomyces naeslundii		100.0	0.43	IND 1		
37733100	**Very good identification**					**37733100**
Actinomyces naeslundii		97.35	0.61	IND 1		
Propionibacterium propionicum		1.88	0.21			
Actinomyces israelii		0.77	0.16			
37733101	**Very good identification**					**37733101**
Actinomyces naeslundii		100.0	0.64	IND 1		
37733102	**Very good identification**					**37733102**
Actinomyces naeslundii		95.49	0.61	IND 1		
Propionibacterium propionicum		2.76	0.24			
Actinomyces israelii		1.14	0.19			
37733103	**Very good identification**					**37733103**
Actinomyces naeslundii		100.0	0.64	IND 1		
37733110	**Good identification**					**37733110**
Actinomyces naeslundii		100.0	0.39	IND 1		
37733111	**Good identification**					**37733111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	IND 1		
37733112	**Good identification**					**37733112**
Actinomyces naeslundii		100.0	0.39	IND 1		
37733113	**Good identification**					**37733113**
Actinomyces naeslundii		100.0	0.42	IND 1		
37733120	**Very good identification**					**37733120**
Actinomyces naeslundii		99.40	0.61	IND 1		
Clostridium perfringens		0.60	0.01			
37733121	**Very good identification**					**37733121**
Actinomyces naeslundii		100.0	0.64	IND 1		
37733122	**Very good identification**					**37733122**
Actinomyces naeslundii		100.0	0.61	IND 1		
37733123	**Very good identification**					**37733123**
Actinomyces naeslundii		100.0	0.64	IND 1		
37733130	**Good identification**					**37733130**
Actinomyces naeslundii		100.0	0.39	IND 1		
37733131	**Good identification**					**37733131**
Actinomyces naeslundii		100.0	0.42	IND 1		
37733132	**Good identification**					**37733132**
Actinomyces naeslundii		100.0	0.39	IND 1		
37733133	**Good identification**					**37733133**
Actinomyces naeslundii		100.0	0.42	IND 1		
37733140	**Acceptable identification**					**37733140**
Actinomyces naeslundii		90.15	0.48	IND 1		
Propionibacterium propionicum		9.85	0.21			
37733141	**Good identification**					**37733141**
Actinomyces naeslundii		100.0	0.51	IND 1		
37733142	**Low discrimination**					**37733142**
Actinomyces naeslundii		85.92	0.48	IND 1		
Propionibacterium propionicum		14.09	0.24			
37733143	**Good identification**					**37733143**
Actinomyces naeslundii		100.0	0.51	IND 1		
37733160	**Good identification**					**37733160**
Actinomyces naeslundii		100.0	0.48	IND 1		
37733161	**Good identification**					**37733161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	IND 1		
37733162	**Good identification**					**37733162**
Actinomyces naeslundii		100.0	0.48	IND 1		
37733163	**Good identification**					**37733163**
Actinomyces naeslundii		100.0	0.51	IND 1		
37737102	**Acceptable identification**					**37737102**
Actinomyces israelii		93.13	0.37	IND 1		
Actinomyces naeslundii		6.87	0.27	CEL 93		
37737502	**Acceptable identification**					**37737502**
Actinomyces israelii		92.97	0.35	IND 1		
Bifidobacterium adolescentis		7.03	0.10	CEL 93		
37743100	**Good identification**					**37743100**
Actinomyces naeslundii		100.0	0.36	IND 1		
37743101	**Good identification**					**37743101**
Actinomyces naeslundii		100.0	0.39	IND 1		
37743102	**Good identification**					**37743102**
Actinomyces naeslundii		100.0	0.36	IND 1		
37743103	**Good identification**					**37743103**
Actinomyces naeslundii		100.0	0.39	IND 1		
37743120	**Good identification**					**37743120**
Actinomyces naeslundii		100.0	0.36	IND 1		
37743121	**Good identification**					**37743121**
Actinomyces naeslundii		100.0	0.39	IND 1		
37743122	**Good identification**					**37743122**
Actinomyces naeslundii		100.0	0.36	IND 1		
37743123	**Good identification**					**37743123**
Actinomyces naeslundii		100.0	0.39	IND 1		
37750002	**Good identification**					**37750002**
Actinomyces viscosus		100.0	0.42	IND 1		
37750102	**Good identification**					**37750102**
Actinomyces viscosus		99.60	0.46	IND 1		
Actinomyces naeslundii		0.40	0.29			
37750103	**Good identification**					**37750103**
Actinomyces viscosus		98.21	0.38	IND 1		
Actinomyces naeslundii		1.79	0.32			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37751000	**Acceptable identification**					**37751000**
Actinomyces naeslundii		90.74	0.45	IND 1		
Actinomyces viscosus		9.26	0.04			
37751001	**Good identification**					**37751001**
Actinomyces naeslundii		100.0	0.48	IND 1		
37751002	**Acceptable identification**					**37751002**
Actinomyces viscosus		90.99	0.38	IND 1		
Actinomyces naeslundii		9.01	0.45			
37751020	**Good identification**					**37751020**
Actinomyces naeslundii		100.0	0.45	IND 1		
37751021	**Good identification**					**37751021**
Actinomyces naeslundii		100.0	0.48	IND 1		
37751022	**Low discrimination**					**37751022**
					GRA	
Actinomyces naeslundii		86.20	0.45	IND 1	99	
Actinomyces viscosus		8.80	0.04		99	
Prevotella oralis		5.00	-.02		1	
37751023	**Good identification**					**37751023**
Actinomyces naeslundii		100.0	0.48	IND 1		
37751041	**Good identification**					**37751041**
Actinomyces naeslundii		100.0	0.35	IND 1		
37751043	**Good identification**					**37751043**
Actinomyces naeslundii		100.0	0.35	IND 1		
37751061	**Good identification**					**37751061**
Actinomyces naeslundii		100.0	0.35	IND 1		
37751063	**Good identification**					**37751063**
Actinomyces naeslundii		100.0	0.35	IND 1		
37751100	**Acceptable identification**					**37751100**
Actinomyces naeslundii		94.01	0.53	IND 1		
Actinomyces viscosus		5.99	0.08			
37751101	**Good identification**					**37751101**
Actinomyces naeslundii		98.60	0.56	IND 1		
Actinomyces viscosus		1.40	0.00			
37751102	**Low discrimination**					**37751102**
Actinomyces viscosus		84.87	0.43	IND 1		
Actinomyces naeslundii		13.45	0.53			
Lactobacillus acidophilus		1.17	0.10			
37751120	**Acceptable identification**					**37751120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		94.78	0.53	IND 1	99	
Clostridium perfringens		3.63	0.06		99	
Prevotella oralis		1.59	-.03		1	
37751121	**Good identification**					**37751121**
Actinomyces naeslundii		100.0	0.56	IND 1		
37751122	**Low discrimination**					**37751122**
					GRA	
Actinomyces naeslundii		67.20	0.53	IND 1	99	
Prevotella oralis		21.46	0.19		1	
Lactobacillus acidophilus		5.85	0.10		99	
37751123	**Good identification**					**37751123**
Actinomyces naeslundii		98.60	0.56	IND 1		
Actinomyces viscosus		1.40	0.00			
37751140	**Good identification**					**37751140**
Actinomyces naeslundii		100.0	0.40	IND 1		
37751141	**Good identification**					**37751141**
Actinomyces naeslundii		100.0	0.43	IND 1		
37751142	**Low discrimination**					**37751142**
Actinomyces naeslundii		73.47	0.40	IND 1		
Actinomyces viscosus		26.53	0.08			
37751143	**Acceptable identification**					**37751143**
Actinomyces naeslundii		92.57	0.43	IND 1		
Actinomyces viscosus		7.43	0.00			
37751160	**Good identification**					**37751160**
Actinomyces naeslundii		100.0	0.40	IND 1		
37751161	**Good identification**					**37751161**
Actinomyces naeslundii		100.0	0.43	IND 1		
37751162	**Good identification**					**37751162**
Actinomyces naeslundii		100.0	0.40	IND 1		
37751163	**Good identification**					**37751163**
Actinomyces naeslundii		100.0	0.43	IND 1		
37752101	**Good identification**					**37752101**
Actinomyces naeslundii		100.0	0.37	IND 1		
37752103	**Low discrimination**					**37752103**
Actinomyces naeslundii		77.78	0.37	IND 1		
Actinomyces viscosus		22.22	0.04	GAL 96		
37752121	**Good identification**					**37752121**
Actinomyces naeslundii		100.0	0.37	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37752123	**Good identification**					**37752123**
Actinomyces naeslundii		100.0	0.37	IND 1		
37753000	**Good identification**					**37753000**
Actinomyces naeslundii		97.16	0.50	IND 1		
Actinomyces israelii		2.84	0.15			
37753001	**Good identification**					**37753001**
Actinomyces naeslundii		100.0	0.53	IND 1		
37753002	**Acceptable identification**					**37753002**
Actinomyces naeslundii		91.21	0.50	IND 1		
Actinomyces viscosus		4.80	0.04			
Actinomyces israelii		4.00	0.18			
37753003	**Good identification**					**37753003**
Actinomyces naeslundii		100.0	0.53	IND 1		
37753020	**Good identification**					**37753020**
Actinomyces naeslundii		100.0	0.50	IND 1		
37753021	**Good identification**					**37753021**
Actinomyces naeslundii		100.0	0.53	IND 1		
37753022	**Good identification**					**37753022**
Actinomyces naeslundii		100.0	0.50	IND 1		
37753023	**Good identification**					**37753023**
Actinomyces naeslundii		100.0	0.53	IND 1		
37753040	**Good identification**					**37753040**
Actinomyces naeslundii		100.0	0.37	IND 1		
37753041	**Good identification**					**37753041**
Actinomyces naeslundii		100.0	0.40	IND 1		
37753042	**Good identification**					**37753042**
Actinomyces naeslundii		100.0	0.37	IND 1		
37753043	**Good identification**					**37753043**
Actinomyces naeslundii		100.0	0.40	IND 1		
37753060	**Good identification**					**37753060**
Actinomyces naeslundii		100.0	0.37	IND 1		
37753061	**Good identification**					**37753061**
Actinomyces naeslundii		100.0	0.40	IND 1		
37753062	**Good identification**					**37753062**
Actinomyces naeslundii		100.0	0.37	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37753063	**Good identification**					**37753063**
Actinomyces naeslundii		100.0	0.40	IND 1		
37753100	**Very good identification**					**37753100**
Actinomyces naeslundii		96.06	0.58	IND 1		
Actinomyces israelii		3.95	0.25			
37753101	**Very good identification**					**37753101**
Actinomyces naeslundii		100.0	0.61	IND 1		
37753102	**Good identification**					**37753102**
Actinomyces naeslundii		91.37	0.58	IND 1		
Actinomyces israelii		5.63	0.28			
Actinomyces viscosus		3.00	0.08			
37753103	**Very good identification**					**37753103**
Actinomyces naeslundii		99.28	0.61	IND 1		
Actinomyces viscosus		0.72	0.00			
37753110	**Good identification**					**37753110**
Actinomyces naeslundii		100.0	0.36	IND 1		
37753111	**Good identification**					**37753111**
Actinomyces naeslundii		100.0	0.39	IND 1		
37753112	**Low discrimination**					**37753112**
Actinomyces naeslundii		85.11	0.36	IND 1		
Actinomyces israelii		14.89	0.14	MLZ 5		
37753113	**Good identification**					**37753113**
Actinomyces naeslundii		100.0	0.39	IND 1		
37753120	**Very good identification**					**37753120**
Actinomyces naeslundii		100.0	0.58	IND 1		
37753121	**Very good identification**					**37753121**
Actinomyces naeslundii		100.0	0.61	IND 1		
37753122	**Very good identification**					**37753122**
					SPO	
Actinomyces naeslundii		97.76	0.58	IND 1	1	
Clostridium ramosum		2.24	0.05		92	
37753123	**Very good identification**					**37753123**
Actinomyces naeslundii		100.0	0.61	IND 1		
37753130	**Good identification**					**37753130**
Actinomyces naeslundii		100.0	0.36	IND 1		
37753131	**Good identification**					**37753131**
Actinomyces naeslundii		100.0	0.39	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37753132	**Good identification**					**37753132**
Actinomyces naeslundii		100.0	0.36	IND 1		
37753133	**Good identification**					**37753133**
Actinomyces naeslundii		100.0	0.39	IND 1		
37753140	**Acceptable identification**					**37753140**
Actinomyces naeslundii		91.70	0.45	IND 1		
Actinomyces israelii		8.30	0.18			
37753141	**Good identification**					**37753141**
Actinomyces naeslundii		100.0	0.48	IND 1		
37753142	**Low discrimination**					**37753142**
Actinomyces naeslundii		88.05	0.45	IND 1		
Actinomyces israelii		11.95	0.21			
37753143	**Good identification**					**37753143**
Actinomyces naeslundii		100.0	0.48	IND 1		
37753160	**Good identification**					**37753160**
Actinomyces naeslundii		100.0	0.45	IND 1		
37753161	**Good identification**					**37753161**
Actinomyces naeslundii		100.0	0.48	IND 1		
37753162	**Good identification**					**37753162**
Actinomyces naeslundii		100.0	0.45	IND 1		
37753163	**Good identification**					**37753163**
Actinomyces naeslundii		100.0	0.48	IND 1		
37757002	**Good identification**					**37757002**
Actinomyces israelii		100.0	0.36	IND 1		
37757100	**Good identification**					**37757100**
Actinomyces israelii		97.91	0.43	IND 1		
Actinomyces naeslundii		2.09	0.24			
37757102	**Good identification**					**37757102**
Actinomyces israelii		98.59	0.46	IND 1		
Actinomyces naeslundii		1.41	0.24			
37757140	**Good identification**					**37757140**
Actinomyces israelii		100.0	0.36	IND 1		
37757142	**Good identification**					**37757142**
Actinomyces israelii		100.0	0.39	IND 1		
37757500	**Good identification**					**37757500**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	100.0	0.42	IND 1		
37757502	**Acceptable identification**					**37757502**
	Actinomyces israelii	93.61	0.45	IND 1		
	Bifidobacterium adolescentis	6.39	0.18			
37757542	**Good identification**					**37757542**
	Actinomyces israelii	100.0	0.38	IND 1		
37761100	**Good identification**					**37761100**
	Actinomyces naeslundii	100.0	0.36	IND 1		
37761101	**Good identification**					**37761101**
	Actinomyces naeslundii	100.0	0.39	IND 1		
37761102	**Low discrimination**					**37761102**
	Actinomyces naeslundii	80.35	0.36	IND 1		
	Actinomyces viscosus	19.65	0.01	FRU 95		
37761103	**Good identification**					**37761103**
	Actinomyces naeslundii	100.0	0.39	IND 1		
37761120	**Good identification**					**37761120**
	Actinomyces naeslundii	100.0	0.36	IND 1		
37761121	**Good identification**					**37761121**
	Actinomyces naeslundii	100.0	0.39	IND 1		
37761122	**Good identification**					**37761122**
	Actinomyces naeslundii	100.0	0.36	IND 1		
37761123	**Good identification**					**37761123**
	Actinomyces naeslundii	100.0	0.39	IND 1		
37763001	**Good identification**					**37763001**
	Actinomyces naeslundii	100.0	0.36	IND 1		
37763003	**Good identification**					**37763003**
	Actinomyces naeslundii	100.0	0.36	IND 1		
37763021	**Good identification**					**37763021**
	Actinomyces naeslundii	100.0	0.36	IND 1		
37763023	**Good identification**					**37763023**
	Actinomyces naeslundii	100.0	0.36	IND 1		
37763100	**Good identification**					**37763100**
	Actinomyces naeslundii	100.0	0.41	IND 1		
37763101	**Good identification**					**37763101**
	Actinomyces naeslundii	100.0	0.44	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37763102	**Good identification**					**37763102**
Actinomyces naeslundii		100.0	0.41	IND 1		
37763103	**Good identification**					**37763103**
Actinomyces naeslundii		100.0	0.44	IND 1		
37763120	**Good identification**					**37763120**
Actinomyces naeslundii		100.0	0.41	IND 1		
37763121	**Good identification**					**37763121**
Actinomyces naeslundii		100.0	0.44	IND 1		
37763122	**Good identification**					**37763122**
Actinomyces naeslundii		100.0	0.41	IND 1		
37763123	**Good identification**					**37763123**
Actinomyces naeslundii		100.0	0.44	IND 1		
37770101	**Good identification**					**37770101**
Actinomyces naeslundii		100.0	0.37	IND 1		
37770121	**Good identification**					**37770121**
Actinomyces naeslundii		100.0	0.37	IND 1		
37770123	**Good identification**					**37770123**
Actinomyces naeslundii		100.0	0.37	IND 1		
37771000	**Good identification**					**37771000**
Actinomyces naeslundii		100.0	0.50	IND 1		
37771001	**Good identification**					**37771001**
Actinomyces naeslundii		100.0	0.53	IND 1		
37771002	**Low discrimination**					**37771002**
Actinomyces naeslundii		68.59	0.50	IND 1		
Actinomyces viscosus		26.87	0.19			
Lactobacillus acidophilus		2.55	0.01			
37771003	**Acceptable identification**					**37771003**
Actinomyces naeslundii		91.99	0.53	IND 1		
Actinomyces viscosus		8.01	0.11			
37771020	**Good identification**					**37771020**
Actinomyces naeslundii		100.0	0.50	IND 1		
37771021	**Good identification**					**37771021**
Actinomyces naeslundii		100.0	0.53	IND 1		
37771022	**Good identification**					**37771022**
Actinomyces naeslundii		96.42	0.50	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		3.58	0.01			
37771023	**Good identification**					**37771023**
Actinomyces naeslundii		100.0	0.53	IND 1		
37771040	**Good identification**					**37771040**
Actinomyces naeslundii		100.0	0.37	IND 1		
37771041	**Good identification**					**37771041**
Actinomyces naeslundii		100.0	0.40	IND 1		
37771042	**Good identification**					**37771042**
Actinomyces naeslundii		100.0	0.37	IND 1		
37771043	**Good identification**					**37771043**
Actinomyces naeslundii		100.0	0.40	IND 1		
37771060	**Good identification**					**37771060**
Actinomyces naeslundii		100.0	0.37	IND 1		
37771061	**Good identification**					**37771061**
Actinomyces naeslundii		100.0	0.40	IND 1		
37771062	**Good identification**					**37771062**
Actinomyces naeslundii		100.0	0.37	IND 1		
37771063	**Good identification**					**37771063**
Actinomyces naeslundii		100.0	0.40	IND 1		
37771100	**Very good identification**					**37771100**
Actinomyces naeslundii		97.35	0.58	IND 1		
Actinomyces israelii		2.65	0.22			
37771101	**Very good identification**					**37771101**
Actinomyces naeslundii		100.0	0.61	IND 1		
37771102	**Low discrimination**					**37771102**
Actinomyces naeslundii		69.65	0.58	IND 1		
Actinomyces viscosus		17.04	0.23			
Lactobacillus acidophilus		10.46	0.19			
37771103	**Good identification**					**37771103**
Actinomyces naeslundii		94.84	0.61	IND 1		
Actinomyces viscosus		5.16	0.15			
37771110	**Good identification**					**37771110**
Actinomyces naeslundii		100.0	0.36	IND 1		
37771111	**Good identification**					**37771111**
Actinomyces naeslundii		100.0	0.39	IND 1		
37771113	**Good identification**					**37771113**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	IND 1		
37771120	**Very good identification**					**37771120**
					SPO	
Actinomyces naeslundii		95.02	0.58	IND 1	1	
Clostridium perfringens		3.48	0.11		84	
Clostridium ramosum		1.50	0.02		92	
37771121	**Very good identification**					**37771121**
Actinomyces naeslundii		100.0	0.61	IND 1		
37771122	**Low discrimination**					**37771122**
					SPO	
Actinomyces naeslundii		80.57	0.58	IND 1	1	
Lactobacillus acidophilus		12.10	0.19		1	
Clostridium ramosum		5.10	0.12		92	
37771123	**Very good identification**					**37771123**
Actinomyces naeslundii		100.0	0.61	IND 1		
37771130	**Good identification**					**37771130**
Actinomyces naeslundii		100.0	0.36	IND 1		
37771131	**Good identification**					**37771131**
Actinomyces naeslundii		100.0	0.39	IND 1		
37771132	**Low discrimination**					**37771132**
					GRA	
Actinomyces naeslundii		81.37	0.36	IND 1	99	
Leptotrichia buccalis		18.63	-.09	MLZ 5	1	
37771133	**Good identification**					**37771133**
Actinomyces naeslundii		100.0	0.39	IND 1		
37771140	**Acceptable identification**					**37771140**
Actinomyces naeslundii		94.33	0.45	IND 1		
Actinomyces israelii		5.67	0.15			
37771141	**Good identification**					**37771141**
Actinomyces naeslundii		100.0	0.48	IND 1		
37771142	**Low discrimination**					**37771142**
Actinomyces naeslundii		83.67	0.45	IND 1		
Lactobacillus acidophilus		8.80	0.04			
Actinomyces israelii		7.54	0.18			
37771143	**Good identification**					**37771143**
Actinomyces naeslundii		100.0	0.48	IND 1		
37771160	**Good identification**					**37771160**
Actinomyces naeslundii		100.0	0.45	IND 1		
37771161	**Good identification**					**37771161**
Actinomyces naeslundii		100.0	0.48	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37771162	**Acceptable identification**					**37771162**
Actinomyces naeslundii		90.49	0.45	IND 1		
Lactobacillus acidophilus		9.51	0.04			
37771163	**Good identification**					**37771163**
Actinomyces naeslundii		100.0	0.48	IND 1		
37772100	**Good identification**					**37772100**
Actinomyces naeslundii		100.0	0.39	IND 1		
37772101	**Good identification**					**37772101**
Actinomyces naeslundii		100.0	0.42	IND 1		
37772102	**Good identification**					**37772102**
Actinomyces naeslundii		100.0	0.39	IND 1		
37772103	**Good identification**					**37772103**
Actinomyces naeslundii		100.0	0.42	IND 1		
37772120	**Good identification**					**37772120**
Actinomyces naeslundii		100.0	0.39	IND 1		
37772121	**Good identification**					**37772121**
Actinomyces naeslundii		100.0	0.42	IND 1		
37772122	**Good identification**					**37772122**
Actinomyces naeslundii		100.0	0.39	IND 1		
37772123	**Good identification**					**37772123**
Actinomyces naeslundii		100.0	0.42	IND 1		
37773000	**Acceptable identification**					**37773000**
Actinomyces naeslundii		94.64	0.55	IND 1		
Actinomyces israelii		5.36	0.25			
37773001	**Very good identification**					**37773001**
Actinomyces naeslundii		100.0	0.58	IND 1		
37773002	**Acceptable identification**					**37773002**
Actinomyces naeslundii		92.17	0.55	IND 1		
Actinomyces israelii		7.83	0.28			
37773003	**Very good identification**					**37773003**
Actinomyces naeslundii		100.0	0.58	IND 1		
37773011	**Good identification**					**37773011**
Actinomyces naeslundii		100.0	0.36	IND 1		
37773013	**Good identification**					**37773013**
Actinomyces naeslundii		100.0	0.36	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37773020	**Good identification**					**37773020**
Actinomyces naeslundii		100.0	0.55	IND 1		
37773021	**Very good identification**					**37773021**
Actinomyces naeslundii		100.0	0.58	IND 1		
37773022	**Good identification**					**37773022**
Actinomyces naeslundii		100.0	0.55	IND 1		
37773023	**Very good identification**					**37773023**
Actinomyces naeslundii		100.0	0.58	IND 1		
37773031	**Good identification**					**37773031**
Actinomyces naeslundii		100.0	0.36	IND 1		
37773033	**Good identification**					**37773033**
Actinomyces naeslundii		100.0	0.36	IND 1		
37773040	**Low discrimination**					**37773040**
Actinomyces naeslundii		88.91	0.42	IND 1		
Actinomyces israelii		11.09	0.18			
37773041	**Good identification**					**37773041**
Actinomyces naeslundii		100.0	0.45	IND 1		
37773042	**Low discrimination**					**37773042**
Actinomyces naeslundii		84.24	0.42	IND 1		
Actinomyces israelii		15.77	0.21			
37773043	**Good identification**					**37773043**
Actinomyces naeslundii		100.0	0.45	IND 1		
37773060	**Good identification**					**37773060**
Actinomyces naeslundii		100.0	0.42	IND 1		
37773061	**Good identification**					**37773061**
Actinomyces naeslundii		100.0	0.45	IND 1		
37773062	**Good identification**					**37773062**
Actinomyces naeslundii		100.0	0.42	IND 1		
37773063	**Good identification**					**37773063**
Actinomyces naeslundii		100.0	0.45	IND 1		
37773100	**Good identification**					**37773100**
Actinomyces naeslundii		92.63	0.63	IND 1		
Actinomyces israelii		7.37	0.35			
37773101	**Very good identification**					**37773101**
Actinomyces naeslundii		100.0	0.66	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
37773102	**Low discrimination**					**37773102**
Actinomyces naeslundii		89.34	0.63	IND 1		
Actinomyces israelii		10.66	0.38			
37773103	**Very good identification**					**37773103**
Actinomyces naeslundii		100.0	0.66	IND 1		
37773110	**Low discrimination**					**37773110**
Actinomyces naeslundii		81.57	0.41	IND 1		
Actinomyces israelii		18.43	0.21	MLZ 5		
37773111	**Good identification**					**37773111**
Actinomyces naeslundii		100.0	0.44	IND 1		
37773112	**Low discrimination**					**37773112**
Actinomyces naeslundii		74.69	0.41	IND 1		
Actinomyces israelii		25.31	0.24	MLZ 5		
37773113	**Good identification**					**37773113**
Actinomyces naeslundii		100.0	0.44	IND 1		
37773120	**Very good identification**					**37773120**
					SPO	
Actinomyces naeslundii		95.13	0.63	IND 1	1	
Clostridium ramosum		4.39	0.15		92	
Actinomyces israelii		0.48	0.14		1	
37773121	**Very good identification**					**37773121**
Actinomyces naeslundii		100.0	0.66	IND 1		
37773122	**Low discrimination**					**37773122**
					SPO	
Actinomyces naeslundii		83.88	0.63	IND 1	1	
Clostridium ramosum		15.49	0.25		92	
Actinomyces israelii		0.64	0.17		1	
37773123	**Very good identification**					**37773123**
Actinomyces naeslundii		100.0	0.66	IND 1		
37773130	**Good identification**					**37773130**
Actinomyces naeslundii		100.0	0.41	IND 1		
37773131	**Good identification**					**37773131**
Actinomyces naeslundii		100.0	0.44	IND 1		
37773132	**Good identification**					**37773132**
Actinomyces naeslundii		100.0	0.41	IND 1		
37773133	**Good identification**					**37773133**
Actinomyces naeslundii		100.0	0.44	IND 1		
37773140	**Low discrimination**					**37773140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		85.08	0.50	IND 1		
Actinomyces israelii		14.92	0.28			
37773141	**Good identification**					**37773141**
Actinomyces naeslundii		100.0	0.53	IND 1		
37773142	**Low discrimination**					**37773142**
Actinomyces naeslundii		79.17	0.50	IND 1		
Actinomyces israelii		20.83	0.31			
37773143	**Good identification**					**37773143**
Actinomyces naeslundii		100.0	0.53	IND 1		
37773160	**Good identification**					**37773160**
Actinomyces naeslundii		100.0	0.50	IND 1		
37773161	**Good identification**					**37773161**
Actinomyces naeslundii		100.0	0.53	IND 1		
37773162	**Good identification**					**37773162**
Actinomyces naeslundii		100.0	0.50	IND 1		
37773163	**Good identification**					**37773163**
Actinomyces naeslundii		100.0	0.53	IND 1		
37773502	**Acceptable identification**					**37773502**
Actinomyces israelii		90.63	0.36	IND 1		
Actinomyces naeslundii		9.37	0.29	XYL 92		
37775100	**Low discrimination**					**37775100**
					SPO	
Actinomyces israelii		77.71	0.40	IND 1	1	
Clostridium butyricum		19.79	0.09		99	
Actinomyces naeslundii		2.50	0.24		1	
37775102	**Acceptable identification**					**37775102**
					SPO	
Actinomyces israelii		91.95	0.43	IND 1	1	
Clostridium butyricum		6.07	0.02		99	
Actinomyces naeslundii		1.98	0.24		1	
37775142	**Good identification**					**37775142**
Actinomyces israelii		100.0	0.36	IND 1		
37775502	**Low discrimination**					**37775502**
					SPO	
Actinomyces israelii		54.14	0.42	IND 1	1	
Clostridium butyricum		39.32	0.18		99	
Bifidobacterium adolescentis		4.54	0.17		1	
37775722	**Low discrimination**					**37775722**
Bacteroides ovatus		86.10	0.49	NIT 1		
Bacteroides thetaiotaomicron		13.48	0.35	ESL 91		
Parabacteroides distasonis		0.15	0.17			
37775732	**Acceptable identification**					**37775732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		90.57	0.46	NIT 1		
Bacteroides thetaiotaomicron		8.27	0.28	ESL 91		
Clostridium clostridioforme		1.04	0.23			
37777000	**Good identification**					**37777000**
Actinomyces israelii		100.0	0.43	IND 1		
37777002	**Good identification**					**37777002**
Actinomyces israelii		100.0	0.46	IND 1		
37777040	**Good identification**					**37777040**
Actinomyces israelii		100.0	0.36	IND 1		
37777042	**Good identification**					**37777042**
Actinomyces israelii		100.0	0.39	IND 1		
37777100	**Good identification**					**37777100**
					SPO	
Actinomyces israelii		98.42	0.53	IND 1	1	
Actinomyces naeslundii		1.09	0.29		1	
Clostridium butyricum		0.49	-.08		99	
37777102	**Good identification**					**37777102**
Actinomyces israelii		99.27	0.56	IND 1		
Actinomyces naeslundii		0.73	0.29			
37777110	**Good identification**					**37777110**
Actinomyces israelii		100.0	0.39	IND 1		
37777112	**Good identification**					**37777112**
Actinomyces israelii		100.0	0.42	IND 1		
37777122	**Low discrimination**					**37777122**
Actinomyces israelii		89.66	0.36	IND 1		
Actinomyces naeslundii		10.34	0.29	NAG 6		
37777140	**Good identification**					**37777140**
Actinomyces israelii		100.0	0.46	IND 1		
37777142	**Good identification**					**37777142**
Actinomyces israelii		100.0	0.49	IND 1		
37777302	**Good identification**					**37777302**
Actinomyces israelii		100.0	0.37	IND 1		
37777400	**Good identification**					**37777400**
Actinomyces israelii		100.0	0.41	IND 1		
37777402	**Good identification**					**37777402**
Actinomyces israelii		100.0	0.44	IND 1		
37777442	**Good identification**					**37777442**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.37	IND 1			
37777500	**Low discrimination**						**37777500**
					GRA	SPO	
Actinomyces israelii		79.34	0.52	IND 1	99	1	
Mitsuokella multacida		16.31	0.27		1	1	
Clostridium butyricum		4.36	0.09		99	99	
37777502	**Good identification**						**37777502**
					GRA	SPO	
Actinomyces israelii		96.01	0.55	IND 1	99	1	
Mitsuokella multacida		1.46	0.11		1	1	
Clostridium butyricum		1.37	0.02		99	99	
37777510	**Low discrimination**						**37777510**
					GRA		
Actinomyces israelii		68.57	0.37	IND 1	99		
Mitsuokella multacida		31.44	0.19	MLZ 13	1		
37777512	**Low discrimination**						**37777512**
Actinomyces israelii		84.79	0.40	IND 1			
Bifidobacterium adolescentis		8.39	0.17	MLZ 13			
Bifidobacterium dentium		3.94	-.07				
37777540	**Low discrimination**						**37777540**
					GRA		
Actinomyces israelii		65.42	0.44	IND 1	99		
Mitsuokella multacida		34.58	0.27		1		
37777542	**Good identification**						**37777542**
					GRA		
Actinomyces israelii		96.23	0.48	IND 1	99		
Mitsuokella multacida		3.77	0.11		1		
37777702	**Low discrimination**						**37777702**
					GRA		
Actinomyces israelii		60.06	0.35	IND 1	99		
Mitsuokella multacida		22.56	0.15	RHA 7	1		
Bacteroides ovatus		17.38	0.06		1		
37777722	**Good identification**						**37777722**
					CAT		
Bacteroides ovatus		98.99	0.41	NIT 1	68		
Bacteroides thetaiotaomicron		0.47	0.01	ESL 91	89		
Mitsuokella multacida		0.32	0.05		1		
37777732	**Good identification**						**37777732**
Bacteroides ovatus		99.72	0.38	NIT 1			
Bacteroides thetaiotaomicron		0.28	-.06	ESL 91			
40000000	**Low discrimination**						**40000000**
					SPO		
Clostridium sporogenes		17.29	0.83		99		
Clostridium botulinum biovar A		14.80	0.84		99		
Atopobium minutum		11.93	0.90		1		
40000001	**Low discrimination**						**40000001**
					COC	GRA	
Campylobacter ureolyticus		66.03	0.66	ESL 1	1	1	
Anaerococcus prevotii		13.36	0.58		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptococcus niger		7.19	0.49		99	99		
40000002	**Low discrimination**							**40000002**
					GRA	SPO		
Atopobium minutum		39.89	0.87		99	1		
Clostridium sporogenes		28.89	0.74		99	99		
Pseudoflavonifractor capillosus		16.13	0.66		1	1		
40000003	**Low discrimination**							**40000003**
					COC			
Anaerococcus prevotii		41.98	0.47	ESL 1	99			
Peptococcus niger		15.95	0.36		99			
Atopobium minutum		13.56	0.53		1			
40000010	**Low discrimination**							**40000010**
					SPO			
Clostridium sporogenes		16.49	0.48	MLZ 1	99			
Clostridium botulinum biovar A		14.11	0.50		99			
Atopobium minutum		11.38	0.56		1			
40000012	**Low discrimination**							**40000012**
					GRA	SPO		
Atopobium minutum		38.32	0.53	MLZ 1	99	1		
Clostridium sporogenes		27.76	0.40		99	99		
Pseudoflavonifractor capillosus		15.49	0.32		1	1		
40000013	**Low discrimination**							**40000013**
					COC	SPO		
Clostridium novyi biovar A		60.07	0.40	GLU 99	1	99		
Anaerococcus prevotii		17.62	0.13		99	1		
Peptococcus niger		6.69	0.02		99	1		
40000020	**Low discrimination**							**40000020**
					COC	GRA		
Atopobium minutum		45.81	0.82		1	99		
Pseudoflavonifractor capillosus		37.04	0.66		1	1		
Peptococcus niger		6.93	0.50		99	99		
40000022	**Excellent identification**							**40000022**
					COC	GRA		
Pseudoflavonifractor capillosus		99.12	1.00		1	1		
Atopobium minutum		0.83	0.79		1	99		
Peptococcus niger		0.03	0.37		99	99		
40000023	**Very good identification**							**40000023**
					COC	GRA		
Pseudoflavonifractor capillosus		98.68	0.66	URE 1	1	1		
Atopobium minutum		0.82	0.44		1	99		
Peptococcus niger		0.32	0.19		99	99		
40000030	**Low discrimination**							**40000030**
					GRA	SPO		
Atopobium minutum		40.34	0.47	MLZ 1	99	1		
Pseudoflavonifractor capillosus		32.61	0.32		1	1		
Clostridium novyi biovar A		12.19	0.43		99	99		
40000031	**Low discrimination**							**40000031**
					COC	GRA	SPO	
Clostridium novyi biovar A		86.86	0.43	GLU 99	1	99	99	
Peptococcus niger		4.30	-.02		99	99	1	
Campylobacter ureolyticus		3.59	-.03		1	1	1	
40000032	**Very good identification**							**40000032**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Pseudoflavonifractor capillosus		99.06	0.66	MLZ 1	1	1		
Atopobium minutum		0.83	0.44		99	1		
Clostridium novyi biovar A		0.07	0.30		99	99		
40000040	**Low discrimination**							**40000040**
					SPO			
Clostridium botulinum biovar A		82.83	0.75	GLU 90	99			
Clostridium sporogenes		3.27	0.48		99			
Atopobium minutum		2.26	0.56		1			
40000042	**Low discrimination**							**40000042**
					SPO			
Atopobium minutum		32.12	0.53	SOR 1	1			
Clostridium sporogenes		23.26	0.40		99			
Clostridium botulinum biovar A		17.85	0.41		99			
40000043	**Low discrimination**							**40000043**
					COC	SPO		
Clostridium novyi biovar A		48.17	0.37	GLU 99	1	99		
Anaerococcus prevotii		21.19	0.13		99	1		
Peptococcus niger		8.05	0.02		99	1		
40000050	**Low discrimination**							**40000050**
Clostridium botulinum biovar A		52.07	0.41	GLU 90				
Clostridium novyi biovar A		37.75	0.50	MLZ 1				
Clostridium sporogenes		2.06	0.14					
40000051	**Good identification**							**40000051**
					GRA	SPO		
Clostridium novyi biovar A		97.14	0.50	GLU 99	99	99		
Campylobacter ureolyticus		1.51	-.03		1	1		
Clostridium botulinum biovar A		1.35	0.06		99	99		
40000052	**Low discrimination**							**40000052**
					SPO			
Clostridium novyi biovar A		72.39	0.37	GLU 99	99			
Atopobium minutum		10.28	0.18		1			
Clostridium sporogenes		7.45	0.06		99			
40000053	**Good identification**							**40000053**
Clostridium novyi biovar A		100.0	0.37	GLU 99				
40000060	**Low discrimination**							**40000060**
					GRA	SPO		
Clostridium botulinum biovar A		31.97	0.41	GLU 90	99	99		
Atopobium minutum		28.76	0.47	NAG 1	99	1		
Pseudoflavonifractor capillosus		23.26	0.32		1	1		
40000061	**Low discrimination**							**40000061**
					COC	GRA	SPO	
Clostridium novyi biovar A		77.96	0.40	GLU 99	1	99	99	
Peptococcus niger		5.79	-.02		99	99	1	
Campylobacter ureolyticus		4.83	-.03		1	1	1	
40000062	**Very good identification**							**40000062**
					GRA	SPO		
Pseudoflavonifractor capillosus		99.07	0.66	SOR 1	1	1		
Atopobium minutum		0.83	0.44		99	1		
Clostridium novyi biovar A		0.04	0.27		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40000070	**Low discrimination**						**40000070**
					SPO		
Clostridium novyi biovar A		87.23	0.40	GLU 99	99		
Clostridium botulinum biovar A		4.86	0.06		99		
Atopobium minutum		4.37	0.13		1		
40000071	**Good identification**						**40000071**
Clostridium novyi biovar A		100.0	0.40	GLU 99			
40000100	**Low discrimination**						**40000100**
					SPO		
Clostridium sporogenes		17.29	0.48	LAC 1	99		
Clostridium botulinum biovar A		14.80	0.50		99		
Atopobium minutum		11.93	0.56		1		
40000102	**Low discrimination**						**40000102**
					GRA	SPO	
Atopobium minutum		39.95	0.53	LAC 1	99	1	
Clostridium sporogenes		28.94	0.40		99	99	
Pseudoflavonifractor capillosus		16.15	0.32		1	1	
40000120	**Low discrimination**						**40000120**
					COC	GRA	
Atopobium minutum		45.94	0.47	LAC 1	1	99	
Pseudoflavonifractor capillosus		37.14	0.32		1	1	
Peptococcus niger		6.95	0.15		99	99	
40000122	**Very good identification**						**40000122**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	LAC 1	1	1	
Atopobium minutum		0.83	0.44		1	99	
Peptococcus niger		0.03	0.03		99	99	
40000140	**Low discrimination**						**40000140**
					SPO		
Clostridium botulinum biovar A		83.14	0.41	GLU 90	99		
Clostridium sporogenes		3.28	0.14	LAC 1	99		
Atopobium minutum		2.27	0.21		1		
40000200	**Low discrimination**						**40000200**
					SPO		
Clostridium sporogenes		17.29	0.48	RHA 1	99		
Clostridium botulinum biovar A		14.80	0.50		99		
Atopobium minutum		11.93	0.56		1		
40000202	**Low discrimination**						**40000202**
					GRA	SPO	
Atopobium minutum		39.95	0.53	RHA 1	99	1	
Clostridium sporogenes		28.94	0.40		99	99	
Pseudoflavonifractor capillosus		16.15	0.32		1	1	
40000220	**Low discrimination**						**40000220**
					COC	GRA	
Atopobium minutum		45.94	0.47	RHA 1	1	99	
Pseudoflavonifractor capillosus		37.14	0.32		1	1	
Peptococcus niger		6.95	0.15		99	99	
40000222	**Very good identification**						**40000222**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	RHA 1	1	1	
Atopobium minutum		0.83	0.44		1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Peptococcus niger		0.03	0.03		99	99	
40000240	**Low discrimination**						**40000240**
					SPO		
Clostridium botulinum biovar A		83.14	0.41	GLU 90	99		
Clostridium sporogenes		3.28	0.14	RHA 1	99		
Atopobium minutum		2.27	0.21		1		
40000400	**Low discrimination**						**40000400**
					SPO		
Clostridium sporogenes		17.29	0.48	ARA 1	99		
Clostridium botulinum biovar A		14.80	0.50		99		
Atopobium minutum		11.93	0.56		1		
40000402	**Low discrimination**						**40000402**
					GRA	SPO	
Atopobium minutum		39.95	0.53	ARA 1	99	1	
Clostridium sporogenes		28.94	0.40		99	99	
Pseudoflavonifractor capillosus		16.15	0.32		1	1	
40000420	**Low discrimination**						**40000420**
					COC	GRA	
Atopobium minutum		45.94	0.47	ARA 1	1	99	
Pseudoflavonifractor capillosus		37.14	0.32		1	1	
Peptococcus niger		6.95	0.15		99	99	
40000422	**Very good identification**						**40000422**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	ARA 1	1	1	
Atopobium minutum		0.83	0.44		1	99	
Peptococcus niger		0.03	0.03		99	99	
40000440	**Low discrimination**						**40000440**
					SPO		
Clostridium botulinum biovar A		83.14	0.41	GLU 90	99		
Clostridium sporogenes		3.28	0.14	ARA 1	99		
Atopobium minutum		2.27	0.21		1		
40001000	**Low discrimination**						**40001000**
					SPO		
Atopobium minutum		81.01	0.82		1		
Clostridium sporogenes		3.56	0.48		99		
Clostridium botulinum biovar A		3.05	0.50		99		
40001002	**Low discrimination**						**40001002**
					GRA	SPO	
Atopobium minutum		85.58	0.79		99	1	
Pseudoflavonifractor capillosus		11.53	0.50		1	1	
Clostridium sporogenes		1.88	0.40		99	99	
40001003	**Low discrimination**						**40001003**
					COC	GRA	
Atopobium minutum		74.60	0.44	URE 1	1	99	
Pseudoflavonifractor capillosus		10.05	0.15		1	1	
Anaerococcus prevotii		7.00	0.13		99	99	
40001010	**Low discrimination**						**40001010**
					SPO		
Atopobium minutum		73.39	0.47	MLZ 1	1		
Clostridium novyi biovar A		9.86	0.37		99		
Clostridium sporogenes		3.22	0.14		99		
40001011	**Low discrimination**						**40001011**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium novyi biovar A		85.60	0.37	GLU 99	99	99	
Campylobacter ureolyticus		7.96	-.03	GAL 10	1	1	
Atopobium minutum		6.44	0.13		99	1	
40001012	**Low discrimination**						**40001012**
					GRA	SPO	
Atopobium minutum		83.87	0.44	MLZ 1	99	1	
Pseudoflavonifractor capillosus		11.30	0.15		1	1	
Clostridium novyi biovar A		2.98	0.24		99	99	
40001020	**Low discrimination**						**40001020**
					COC	GRA	
Atopobium minutum		78.01	0.73		1	99	
Pseudoflavonifractor capillosus		21.02	0.50		1	1	
Peptococcus niger		0.36	0.15		99	99	
40001021	**Low discrimination**						**40001021**
					GRA	SPO	
Atopobium minutum		68.84	0.39	URE 1	99	1	
Pseudoflavonifractor capillosus		18.55	0.15		1	1	
Clostridium novyi biovar A		6.94	0.26		99	99	
40001022	**Very good identification**						**40001022**
					GRA	SPO	
Pseudoflavonifractor capillosus		97.56	0.84	GAL 10	1	1	
Atopobium minutum		2.44	0.70		99	1	
Clostridium novyi biovar A		0.00	0.13		99	99	
40001023	**Good identification**						**40001023**
					GRA	SPO	
Pseudoflavonifractor capillosus		97.50	0.50	GAL 10	1	1	
Atopobium minutum		2.44	0.36	URE 1	99	1	
Clostridium novyi biovar A		0.07	0.13		99	99	
40001030	**Low discrimination**						**40001030**
					GRA	SPO	
Atopobium minutum		72.98	0.39	MLZ 1	99	1	
Pseudoflavonifractor capillosus		19.67	0.15		1	1	
Clostridium novyi biovar A		7.35	0.26		99	99	
40001032	**Good identification**						**40001032**
					GRA	SPO	
Pseudoflavonifractor capillosus		97.50	0.50	GAL 10	1	1	
Atopobium minutum		2.44	0.36	MLZ 1	99	1	
Clostridium novyi biovar A		0.07	0.13		99	99	
40001040	**Low discrimination**						**40001040**
					SPO		
Clostridium botulinum biovar A		45.93	0.41	GLU 90	99		
Atopobium minutum		41.32	0.47	GAL 1	1		
Clostridium novyi biovar A		3.70	0.34		99		
40001042	**Low discrimination**						**40001042**
					GRA	SPO	
Atopobium minutum		83.53	0.44	SOR 1	99	1	
Pseudoflavonifractor capillosus		11.26	0.15		1	1	
Clostridium novyi biovar A		1.98	0.21		99	99	
40001060	**Low discrimination**						**40001060**
					GRA	SPO	
Atopobium minutum		72.97	0.39	SOR 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		19.67	0.15		1	1	
Clostridium novyi biovar A		4.90	0.23		99	99	
40001062	**Good identification**						**40001062**
					GRA	SPO	
Pseudoflavonifractor capillosus		97.52	0.50	GAL 10	1	1	
Atopobium minutum		2.44	0.36	SOR 1	99	1	
Clostridium novyi biovar A		0.04	0.10		99	99	
40001100	**Low discrimination**						**40001100**
					SPO		
Atopobium minutum		81.02	0.47	LAC 1	1		
Clostridium sporogenes		3.56	0.14		99		
Clostridium botulinum biovar A		3.05	0.15		99		
40001102	**Low discrimination**						**40001102**
					GRA	SPO	
Atopobium minutum		86.45	0.44	LAC 1	99	1	
Pseudoflavonifractor capillosus		11.65	0.15		1	1	
Clostridium sporogenes		1.90	0.06		99	99	
40001120	**Low discrimination**						**40001120**
					GRA		
Atopobium minutum		78.77	0.39	LAC 1	99		
Pseudoflavonifractor capillosus		21.23	0.15		1		
40001122	**Good identification**						**40001122**
					GRA		
Pseudoflavonifractor capillosus		97.56	0.50	GAL 10	1		
Atopobium minutum		2.44	0.36	LAC 1	99		
40001200	**Low discrimination**						**40001200**
					SPO		
Atopobium minutum		81.42	0.47	RHA 1	1		
Clostridium sporogenes		3.57	0.14		99		
Clostridium botulinum biovar A		3.06	0.15		99		
40001202	**Low discrimination**						**40001202**
					GRA	SPO	
Atopobium minutum		86.45	0.44	RHA 1	99	1	
Pseudoflavonifractor capillosus		11.65	0.15		1	1	
Clostridium sporogenes		1.90	0.06		99	99	
40001220	**Low discrimination**						**40001220**
					GRA		
Atopobium minutum		78.77	0.39	RHA 1	99		
Pseudoflavonifractor capillosus		21.23	0.15		1		
40001222	**Good identification**						**40001222**
					GRA		
Pseudoflavonifractor capillosus		97.56	0.50	GAL 10	1		
Atopobium minutum		2.44	0.36	RHA 1	99		
40001400	**Low discrimination**						**40001400**
					SPO		
Atopobium minutum		81.42	0.47	ARA 1	1		
Clostridium sporogenes		3.57	0.14		99		
Clostridium botulinum biovar A		3.06	0.15		99		
40001402	**Low discrimination**						**40001402**
					GRA	SPO	
Atopobium minutum		86.45	0.44	ARA 1	99	1	
Pseudoflavonifractor capillosus		11.65	0.15		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		1.90	0.06		99	99	
40001420	**Low discrimination**						**40001420**
					GRA		
Atopobium minutum		78.77	0.39	ARA 1	99		
Pseudoflavonifractor capillosus		21.23	0.15		1		
40001422	**Good identification**						**40001422**
					GRA		
Pseudoflavonifractor capillosus		97.56	0.50	GAL 10	1		
Atopobium minutum		2.44	0.36	ARA 1	99		
40002000	**Low discrimination**						**40002000**
					SPO		
Clostridium sporogenes		16.97	0.48	MAN 1	99		
Clostridium botulinum biovar A		14.53	0.50		99		
Atopobium minutum		11.71	0.56		1		
40002002	**Low discrimination**						**40002002**
					GRA	SPO	
Atopobium minutum		39.92	0.53	MAN 1	99	1	
Clostridium sporogenes		28.92	0.40		99	99	
Pseudoflavonifractor capillosus		16.14	0.32		1	1	
40002020	**Low discrimination**						**40002020**
					COC	GRA	
Atopobium minutum		45.85	0.47	MAN 1	1	99	
Pseudoflavonifractor capillosus		37.07	0.32		1	1	
Peptococcus niger		6.94	0.15		99	99	
40002022	**Very good identification**						**40002022**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	MAN 1	1	1	
Atopobium minutum		0.83	0.44		1	99	
Peptococcus niger		0.03	0.03		99	99	
40002040	**Low discrimination**						**40002040**
					SPO		
Clostridium botulinum biovar A		82.88	0.41	GLU 90	99		
Clostridium sporogenes		3.27	0.14	MAN 1	99		
Atopobium minutum		2.26	0.21		1		
40003000	**Low discrimination**						**40003000**
					SPO		
Atopobium minutum		81.15	0.47	MAN 1	1		
Clostridium sporogenes		3.56	0.14		99		
Clostridium botulinum biovar A		3.05	0.15		99		
40003002	**Low discrimination**						**40003002**
					GRA	SPO	
Atopobium minutum		86.45	0.44	MAN 1	99	1	
Pseudoflavonifractor capillosus		11.65	0.15		1	1	
Clostridium sporogenes		1.90	0.06		99	99	
40003020	**Low discrimination**						**40003020**
					GRA		
Atopobium minutum		78.77	0.39	MAN 1	99		
Pseudoflavonifractor capillosus		21.23	0.15		1		
40003022	**Good identification**						**40003022**
					GRA		
Pseudoflavonifractor capillosus		97.56	0.50	GAL 10	1		
Atopobium minutum		2.44	0.36	MAN 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40004000	**Low discrimination**						**40004000**
					SPO		
Clostridium sporogenes		17.15	0.48	XYL 1	99		
Clostridium botulinum biovar A		14.68	0.50		99		
Atopobium minutum		11.84	0.56		1		
40004002	**Low discrimination**						**40004002**
					GRA	SPO	
Atopobium minutum		39.95	0.53	XYL 1	99	1	
Clostridium sporogenes		28.94	0.40		99	99	
Pseudoflavonifractor capillosus		16.15	0.32		1	1	
40004020	**Low discrimination**						**40004020**
					COC	GRA	
Atopobium minutum		45.94	0.47	XYL 1	1	99	
Pseudoflavonifractor capillosus		37.14	0.32		1	1	
Peptococcus niger		6.95	0.15		99	99	
40004022	**Very good identification**						**40004022**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	XYL 1	1	1	
Atopobium minutum		0.83	0.44		1	99	
Peptococcus niger		0.03	0.03		99	99	
40004040	**Low discrimination**						**40004040**
Clostridium glycolicum		74.81	0.49	GLU 99			
Clostridium botulinum biovar A		20.94	0.41	FRU 90			
Clostridium sporogenes		0.83	0.14				
40005000	**Low discrimination**						**40005000**
					SPO		
Atopobium minutum		79.91	0.47	XYL 1	1		
Clostridium sporogenes		3.51	0.14		99		
Clostridium botulinum biovar A		3.00	0.15		99		
40005002	**Low discrimination**						**40005002**
					GRA	SPO	
Atopobium minutum		86.45	0.44	XYL 1	99	1	
Pseudoflavonifractor capillosus		11.65	0.15		1	1	
Clostridium sporogenes		1.90	0.06		99	99	
40005020	**Low discrimination**						**40005020**
					GRA		
Atopobium minutum		78.77	0.39	XYL 1	99		
Pseudoflavonifractor capillosus		21.23	0.15		1		
40005022	**Good identification**						**40005022**
					GRA		
Pseudoflavonifractor capillosus		97.56	0.50	GAL 10	1		
Atopobium minutum		2.44	0.36	XYL 1	99		
40010000	**Low discrimination**						**40010000**
					SPO		
Atopobium minutum		68.54	0.89		1		
Clostridium sporogenes		18.98	0.70		99		
Clostridium botulinum biovar A		5.88	0.63		99		
40010001	**Low discrimination**						**40010001**
					GRA	SPO	
Atopobium minutum		47.79	0.55	URE 1	99	1	
Campylobacter ureolyticus		21.34	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		13.24	0.35		99	99	
40010002	**Low discrimination**						**40010002**
					GRA	SPO	
Atopobium minutum		83.78	0.86		99	1	
Clostridium sporogenes		11.60	0.61		99	99	
Pseudoflavonifractor capillosus		4.08	0.50		1	1	
40010003	**Low discrimination**						**40010003**
					GRA	SPO	
Atopobium minutum		79.78	0.52	URE 1	99	1	
Clostridium sporogenes		11.05	0.27		99	99	
Pseudoflavonifractor capillosus		3.88	0.15		1	1	
40010010	**Low discrimination**						**40010010**
					SPO		
Atopobium minutum		39.99	0.55	MLZ 1	1		
Clostridium difficile		39.91	0.45		99		
Clostridium sporogenes		11.08	0.35		99		
40010011	**Low discrimination**						**40010011**
					SPO		
Clostridium novyi biovar A		63.83	0.37	GLU 99	99		
Atopobium minutum		13.29	0.21	FRU 10	1		
Clostridium difficile		13.26	0.10		99		
40010012	**Low discrimination**						**40010012**
					GRA	SPO	
Atopobium minutum		82.30	0.52	MLZ 1	99	1	
Clostridium sporogenes		11.40	0.27		99	99	
Pseudoflavonifractor capillosus		4.01	0.15		1	1	
40010020	**Low discrimination**						**40010020**
					GRA	SPO	
Atopobium minutum		89.65	0.81		99	1	
Pseudoflavonifractor capillosus		8.73	0.50		1	1	
Clostridium sporogenes		0.75	0.35		99	99	
40010021	**Low discrimination**						**40010021**
					GRA	SPO	
Atopobium minutum		85.34	0.47	URE 1	99	1	
Pseudoflavonifractor capillosus		8.31	0.15		1	1	
Clostridium novyi biovar A		3.10	0.26		99	99	
40010022	**Good identification**						**40010022**
					GRA	SPO	
Pseudoflavonifractor capillosus		93.50	0.84	FRU 10	1	1	
Atopobium minutum		6.47	0.78		99	1	
Clostridium sporogenes		0.03	0.27		99	99	
40010023	**Acceptable identification**						**40010023**
					GRA	SPO	
Pseudoflavonifractor capillosus		93.47	0.50	FRU 10	1	1	
Atopobium minutum		6.47	0.44	URE 1	99	1	
Clostridium novyi biovar A		0.06	0.13		99	99	
40010030	**Low discrimination**						**40010030**
					GRA	SPO	
Atopobium minutum		85.30	0.47	MLZ 1	99	1	
Pseudoflavonifractor capillosus		8.30	0.15		1	1	
Clostridium novyi biovar A		3.10	0.26		99	99	
40010032	**Acceptable identification**						**40010032**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Pseudoflavonifractor capillosus		93.47	0.50	FRU 10	1	1	
Atopobium minutum		6.47	0.44	MLZ 1	99	1	
Clostridium novyi biovar A		0.06	0.13		99	99	
40010040	**Low discrimination**						**40010040**
					SPO		
Clostridium botulinum biovar A		53.69	0.54	GLU 90	99		
Atopobium minutum		21.17	0.55	FRU 6	1		
Clostridium glycolicum		14.38	0.43		99		
40010042	**Low discrimination**						**40010042**
					GRA	SPO	
Atopobium minutum		79.37	0.52	SOR 1	99	1	
Clostridium sporogenes		10.99	0.27		99	99	
Pseudoflavonifractor capillosus		3.86	0.15		1	1	
40010060	**Low discrimination**						**40010060**
					GRA	SPO	
Atopobium minutum		79.01	0.47	SOR 1	99	1	
Pseudoflavonifractor capillosus		7.69	0.15		1	1	
Clostridium botulinum biovar A		6.07	0.20		99	99	
40010062	**Acceptable identification**						**40010062**
					GRA	SPO	
Pseudoflavonifractor capillosus		93.45	0.50	FRU 10	1	1	
Atopobium minutum		6.47	0.44	SOR 1	99	1	
Clostridium novyi biovar A		0.04	0.10		99	99	
40010100	**Low discrimination**						**40010100**
					SPO		
Atopobium minutum		66.58	0.55	LAC 1	1		
Clostridium sporogenes		18.44	0.35		99		
Clostridium botulinum biovar A		5.71	0.29		99		
40010102	**Low discrimination**						**40010102**
					GRA	SPO	
Atopobium minutum		83.84	0.52	LAC 1	99	1	
Clostridium sporogenes		11.61	0.27		99	99	
Pseudoflavonifractor capillosus		4.08	0.15		1	1	
40010120	**Acceptable identification**						**40010120**
					GRA	SPO	
Atopobium minutum		90.03	0.47	LAC 1	99	1	
Pseudoflavonifractor capillosus		8.76	0.15		1	1	
Clostridium sporogenes		0.76	0.01		99	99	
40010122	**Acceptable identification**						**40010122**
					GRA		
Pseudoflavonifractor capillosus		93.53	0.50	FRU 10	1		
Atopobium minutum		6.47	0.44	LAC 1	99		
40010200	**Low discrimination**						**40010200**
					SPO		
Atopobium minutum		68.67	0.55	RHA 1	1		
Clostridium sporogenes		19.02	0.35		99		
Clostridium botulinum biovar A		5.89	0.29		99		
40010202	**Low discrimination**						**40010202**
					GRA	SPO	
Atopobium minutum		84.24	0.52	RHA 1	99	1	
Clostridium sporogenes		11.67	0.27		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		4.10	0.15		1	1	
40010220	**Acceptable identification**						**40010220**
					GRA	SPO	
Atopobium minutum		90.44	0.47	RHA 1	99	1	
Pseudoflavonifractor capillosus		8.80	0.15		1	1	
Clostridium sporogenes		0.76	0.01		99	99	
40010222	**Acceptable identification**						**40010222**
					GRA		
Pseudoflavonifractor capillosus		93.53	0.50	FRU 10	1		
Atopobium minutum		6.47	0.44	RHA 1	99		
40010400	**Low discrimination**						**40010400**
					SPO		
Atopobium minutum		68.67	0.55	ARA 1	1		
Clostridium sporogenes		19.02	0.35		99		
Clostridium botulinum biovar A		5.89	0.29		99		
40010402	**Low discrimination**						**40010402**
					GRA	SPO	
Atopobium minutum		84.24	0.52	ARA 1	99	1	
Clostridium sporogenes		11.67	0.27		99	99	
Pseudoflavonifractor capillosus		4.10	0.15		1	1	
40010420	**Acceptable identification**						**40010420**
					GRA	SPO	
Atopobium minutum		90.44	0.47	ARA 1	99	1	
Pseudoflavonifractor capillosus		8.80	0.15		1	1	
Clostridium sporogenes		0.76	0.01		99	99	
40010422	**Acceptable identification**						**40010422**
					GRA		
Pseudoflavonifractor capillosus		93.53	0.50	FRU 10	1		
Atopobium minutum		6.47	0.44	ARA 1	99		
40011000	**Very good identification**						**40011000**
					SPO		
Atopobium minutum		98.51	0.81		1		
Clostridium sporogenes		0.83	0.35		99		
Clostridium botulinum biovar A		0.26	0.29		99		
40011001	**Good identification**						**40011001**
					GRA	SPO	
Atopobium minutum		95.98	0.47	URE 1	99	1	
Clostridium novyi biovar A		1.55	0.20		99	99	
Campylobacter ureolyticus		1.30	-.03		1	1	
40011002	**Very good identification**						**40011002**
					GRA	SPO	
Atopobium minutum		97.99	0.78		99	1	
Pseudoflavonifractor capillosus		1.59	0.33		1	1	
Clostridium sporogenes		0.41	0.27		99	99	
40011003	**Good identification**						**40011003**
					GRA		
Atopobium minutum		98.40	0.44	URE 1	99		
Pseudoflavonifractor capillosus		1.60	-.01		1		
40011010	**Acceptable identification**						**40011010**
					SPO		
Atopobium minutum		94.81	0.47	MLZ 1	1		
Clostridium difficile		2.87	0.10		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		1.53	0.20		99		
40011012	**Good identification**						**40011012**
					GRA		
Atopobium minutum		98.40	0.44	MLZ 1	99		
Pseudoflavonifractor capillosus		1.60	-.01		1		
40011020	**Very good identification**						**40011020**
					GRA	SPO	
Atopobium minutum		96.82	0.73		99	1	
Pseudoflavonifractor capillosus		3.14	0.33		1	1	
Clostridium sporogenes		0.03	0.01		99	99	
40011021	**Good identification**						**40011021**
					GRA	SPO	
Atopobium minutum		95.73	0.39	URE 1	99	1	
Pseudoflavonifractor capillosus		3.11	-.01		1	1	
Clostridium novyi biovar A		1.16	0.10		99	99	
40011022	**Low discrimination**						**40011022**
					GRA		
Pseudoflavonifractor capillosus		82.81	0.67	FRU 10	1		
Atopobium minutum		17.19	0.70	GAL 10	99		
40011030	**Good identification**						**40011030**
					GRA	SPO	
Atopobium minutum		95.73	0.39	MLZ 1	99	1	
Pseudoflavonifractor capillosus		3.11	-.01		1	1	
Clostridium novyi biovar A		1.16	0.10		99	99	
40011040	**Low discrimination**						**40011040**
					SPO		
Atopobium minutum		75.62	0.47	SOR 1	1		
Clostridium glycolicum		17.12	0.27		99		
Clostridium botulinum biovar A		5.81	0.20		99		
40011042	**Good identification**						**40011042**
					GRA		
Atopobium minutum		98.40	0.44	SOR 1	99		
Pseudoflavonifractor capillosus		1.60	-.01		1		
40011060	**Good identification**						**40011060**
					GRA		
Atopobium minutum		96.86	0.39	SOR 1	99		
Pseudoflavonifractor capillosus		3.14	-.01		1		
40011100	**Low discrimination**						**40011100**
					GRA		
Atopobium minutum		82.36	0.47	LAC 1	99		
Fusobacterium mortiferum		12.34	0.38		1		
Actinomyces meyeri		4.61	0.22		99		
40011102	**Good identification**						**40011102**
					GRA		
Atopobium minutum		96.04	0.44	LAC 1	99		
Fusobacterium mortiferum		2.40	0.22		1		
Pseudoflavonifractor capillosus		1.56	-.01		1		
40011120	**Acceptable identification**						**40011120**
					GRA		
Atopobium minutum		94.76	0.39	LAC 1	99		
Pseudoflavonifractor capillosus		3.07	-.01		1		
Actinomyces meyeri		2.17	0.07		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
40011200	**Good identification**					**40011200**
					SPO	
Atopobium minutum		99.17	0.47	RHA 1	1	
Clostridium sporogenes		0.83	0.01		99	
40011202	**Good identification**					**40011202**
					GRA	
Atopobium minutum		98.40	0.44	RHA 1	99	
Pseudoflavonifractor capillosus		1.60	-.01		1	
40011220	**Good identification**					**40011220**
					GRA	
Atopobium minutum		96.86	0.39	RHA 1	99	
Pseudoflavonifractor capillosus		3.14	-.01		1	
40011400	**Good identification**					**40011400**
					SPO	
Atopobium minutum		99.17	0.47	ARA 1	1	
Clostridium sporogenes		0.83	0.01		99	
40011402	**Good identification**					**40011402**
					GRA	
Atopobium minutum		98.40	0.44	ARA 1	99	
Pseudoflavonifractor capillosus		1.60	-.01		1	
40011420	**Good identification**					**40011420**
					GRA	
Atopobium minutum		96.86	0.39	ARA 1	99	
Pseudoflavonifractor capillosus		3.14	-.01		1	
40012000	**Low discrimination**					**40012000**
					SPO	
Eubacterium limosum		81.01	0.63	GLU 99	1	
Clostridium difficile		10.12	0.49		99	
Atopobium minutum		5.50	0.55		1	
40012002	**Low discrimination**					**40012002**
					SPO	
Atopobium minutum		68.06	0.52	MAN 1	1	
Eubacterium limosum		15.19	0.29		1	
Clostridium sporogenes		9.43	0.27		99	
40012010	**Very good identification**					**40012010**
					SPO	
Clostridium difficile		98.21	0.61	GLU 99	99	
Eubacterium limosum		1.63	0.29		1	
Atopobium minutum		0.11	0.21		1	
40012020	**Low discrimination**					**40012020**
					GRA	
Atopobium minutum		59.16	0.47	MAN 1	99	
Eubacterium limosum		26.41	0.29		99	
Pseudoflavonifractor capillosus		5.76	0.15		1	
40012022	**Acceptable identification**					**40012022**
					GRA	
Pseudoflavonifractor capillosus		93.41	0.50	FRU 10	1	
Atopobium minutum		6.46	0.44	MAN 1	99	
Pseudoramibacter alactolyticus		0.09	-.02		99	
40012050	**Good identification**					**40012050**
					SPO	
Clostridium difficile		99.68	0.39	GLU 99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		0.32	-.06	SOR 5	1		
40013000	**Low discrimination**						**40013000**
					SPO		
Atopobium minutum		66.00	0.47	MAN 1	1		
Eubacterium limosum		29.46	0.29		1		
Clostridium difficile		3.68	0.15		99		
40013002	**Good identification**						**40013002**
					GRA		
Atopobium minutum		97.75	0.44	MAN 1	99		
Pseudoflavonifractor capillosus		1.59	-.01		1		
Eubacterium limosum		0.66	-.06		99		
40013020	**Good identification**						**40013020**
					GRA		
Atopobium minutum		95.61	0.39	MAN 1	99		
Pseudoflavonifractor capillosus		3.10	-.01		1		
Eubacterium limosum		1.29	-.06		99		
40014000	**Low discrimination**						**40014000**
					SPO		
Atopobium minutum		45.70	0.55	XYL 1	1		
Clostridium glycolicum		31.04	0.43		99		
Clostridium sporogenes		12.66	0.35		99		
40014002	**Low discrimination**						**40014002**
					GRA	SPO	
Atopobium minutum		83.51	0.52	XYL 1	99	1	
Clostridium sporogenes		11.57	0.27		99	99	
Pseudoflavonifractor capillosus		4.06	0.15		1	1	
40014020	**Low discrimination**						**40014020**
					GRA	SPO	
Atopobium minutum		88.79	0.47	XYL 1	99	1	
Pseudoflavonifractor capillosus		8.64	0.15		1	1	
Clostridium glycolicum		1.83	0.09		99	99	
40014022	**Acceptable identification**						**40014022**
					GRA		
Pseudoflavonifractor capillosus		93.53	0.50	FRU 10	1		
Atopobium minutum		6.47	0.44	XYL 1	99		
40014040	**Very good identification**						**40014040**
					SPO		
Clostridium glycolicum		99.70	0.65	GLU 99	99		
Clostridium botulinum biovar A		0.20	0.20		99		
Atopobium minutum		0.08	0.21		1		
40015000	**Low discrimination**						**40015000**
					SPO		
Atopobium minutum		78.94	0.47	XYL 1	1		
Clostridium glycolicum		17.87	0.27		99		
Actinomyces meyeri		2.15	0.16		1		
40015002	**Good identification**						**40015002**
					GRA		
Atopobium minutum		98.40	0.44	XYL 1	99		
Pseudoflavonifractor capillosus		1.60	-.01		1		
40015020	**Good identification**						**40015020**
					GRA		
Atopobium minutum		96.86	0.39	XYL 1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		3.14	-.01		1		
40015040	**Good identification**						**40015040**
					SPO		
Clostridium glycolicum		99.77	0.49	GLU 99	99		
Atopobium minutum		0.23	0.13	GAL 10	1		
40016010	**Good identification**						**40016010**
					SPO		
Clostridium difficile		99.68	0.39	GLU 99	99		
Eubacterium limosum		0.32	-.06	XYL 5	1		
40020000	**Low discrimination**						**40020000**
					SPO		
Clostridium sporogenes		56.78	0.74		99		
Clostridium botulinum biovar A		36.46	0.73		99		
Atopobium minutum		1.19	0.56		1		
40020001	**Low discrimination**						**40020001**
					GRA	SPO	
Clostridium sporogenes		34.77	0.40	URE 1	99	99	
Campylobacter ureolyticus		29.67	0.32		1	1	
Clostridium botulinum biovar A		22.32	0.39		99	99	
40020002	**Good identification**						**40020002**
					SPO		
Clostridium sporogenes		91.49	0.66		99		
Atopobium minutum		3.83	0.53		1		
Clostridium botulinum biovar A		1.78	0.39		99		
40020010	**Low discrimination**						**40020010**
					SPO		
Clostridium sporogenes		56.60	0.40	MLZ 1	99		
Clostridium botulinum biovar A		36.35	0.39		99		
Atopobium minutum		1.18	0.21		1		
40020020	**Low discrimination**						**40020020**
					SPO		
Clostridium sporogenes		32.38	0.40	NAG 1	99		
Atopobium minutum		22.35	0.47		1		
Clostridium botulinum biovar A		20.79	0.39		99		
40020022	**Very good identification**						**40020022**
					GRA	SPO	
Pseudoflavonifractor capillosus		98.54	0.66	TRE 1	1	1	
Atopobium minutum		0.82	0.44		99	1	
Clostridium sporogenes		0.59	0.32		99	99	
40020040	**Good identification**						**40020040**
					SPO		
Clostridium botulinum biovar A		94.41	0.64	GLU 90	99		
Clostridium sporogenes		4.97	0.40		99		
Atopobium minutum		0.10	0.21		1		
40020100	**Low discrimination**						**40020100**
					SPO		
Clostridium sporogenes		56.88	0.40	LAC 1	99		
Clostridium botulinum biovar A		36.52	0.39		99		
Atopobium minutum		1.19	0.21		1		
40020200	**Low discrimination**						**40020200**
					SPO		
Clostridium sporogenes		56.88	0.40	RHA 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		36.52	0.39		99		
Atopobium minutum		1.19	0.21		1		
40020400	**Low discrimination**						**40020400**
					SPO		
Clostridium sporogenes		56.88	0.40	ARA 1	99		
Clostridium botulinum biovar A		36.52	0.39		99		
Atopobium minutum		1.19	0.21		1		
40021000	**Low discrimination**						**40021000**
					SPO		
Clostridium sporogenes		41.09	0.40	GAL 1	99		
Atopobium minutum		28.36	0.47		1		
Clostridium botulinum biovar A		26.39	0.39		99		
40021002	**Low discrimination**						**40021002**
					GRA	SPO	
Atopobium minutum		53.39	0.44	TRE 1	99	1	
Clostridium sporogenes		38.67	0.32		99	99	
Pseudoflavonifractor capillosus		7.19	0.15		1	1	
40021020	**Low discrimination**						**40021020**
					GRA	SPO	
Atopobium minutum		74.54	0.39	TRE 1	99	1	
Pseudoflavonifractor capillosus		20.09	0.15		1	1	
Clostridium sporogenes		3.27	0.06		99	99	
40021022	**Good identification**						**40021022**
					GRA	SPO	
Pseudoflavonifractor capillosus		97.51	0.50	TRE 1	1	1	
Atopobium minutum		2.44	0.36	GAL 10	99	1	
Clostridium sporogenes		0.05	-.02		99	99	
40022000	**Low discrimination**						**40022000**
					SPO		
Clostridium sporogenes		56.70	0.40	MAN 1	99		
Clostridium botulinum biovar A		36.41	0.39		99		
Atopobium minutum		1.19	0.21		1		
40024000	**Low discrimination**						**40024000**
					SPO		
Clostridium sporogenes		56.88	0.40	XYL 1	99		
Clostridium botulinum biovar A		36.52	0.39		99		
Atopobium minutum		1.19	0.21		1		
40030000	**Low discrimination**						**40030000**
					SPO		
Clostridium sporogenes		73.87	0.61		99		
Clostridium botulinum biovar A		17.16	0.53		99		
Atopobium minutum		8.08	0.55		1		
40030002	**Low discrimination**						**40030002**
					GRA	SPO	
Clostridium sporogenes		80.87	0.53		99	99	
Atopobium minutum		17.70	0.52		99	1	
Pseudoflavonifractor capillosus		0.86	0.15		1	1	
40030020	**Low discrimination**						**40030020**
					GRA	SPO	
Atopobium minutum		69.51	0.47	TRE 1	99	1	
Clostridium sporogenes		19.25	0.27		99	99	
Pseudoflavonifractor capillosus		6.77	0.15		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40030022	**Acceptable identification**						**40030022**
					GRA	SPO	
Pseudoflavonifractor capillosus		92.70	0.50	FRU 10	1	1	
Atopobium minutum		6.41	0.44	TRE 1	99	1	
Clostridium sporogenes		0.89	0.19		99	99	
40030040	**Low discrimination**						**40030040**
					SPO		
Clostridium botulinum biovar A		85.30	0.44	GLU 90	99		
Clostridium sporogenes		12.42	0.27	FRU 6	99		
Atopobium minutum		1.36	0.21		1		
40031000	**Low discrimination**						**40031000**
					SPO		
Atopobium minutum		73.51	0.47	TRE 1	1		
Clostridium sporogenes		20.36	0.27		99		
Clostridium botulinum biovar A		4.73	0.19		99		
40031002	**Low discrimination**						**40031002**
					GRA	SPO	
Atopobium minutum		86.60	0.44	TRE 1	99	1	
Clostridium sporogenes		11.99	0.19		99	99	
Pseudoflavonifractor capillosus		1.41	-.01		1	1	
40031020	**Good identification**						**40031020**
					GRA		
Atopobium minutum		96.86	0.39	TRE 1	99		
Pseudoflavonifractor capillosus		3.14	-.01		1		
40032010	**Good identification**						**40032010**
					SPO		
Clostridium difficile		99.78	0.42	GLU 99	99		
Eubacterium limosum		0.22	-.06	TRE 7	1		
40040000	**Low discrimination**						**40040000**
					SPO		
Clostridium sporogenes		17.29	0.48	CEL 1	99		
Clostridium botulinum biovar A		14.80	0.50		99		
Atopobium minutum		11.93	0.56		1		
40040002	**Low discrimination**						**40040002**
					GRA	SPO	
Atopobium minutum		39.95	0.53	CEL 1	99	1	
Clostridium sporogenes		28.94	0.40		99	99	
Pseudoflavonifractor capillosus		16.15	0.32		1	1	
40040020	**Low discrimination**						**40040020**
					COC	GRA	
Atopobium minutum		45.94	0.47	CEL 1	1	99	
Pseudoflavonifractor capillosus		37.14	0.32		1	1	
Peptococcus niger		6.95	0.15		99	99	
40040022	**Very good identification**						**40040022**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	CEL 1	1	1	
Atopobium minutum		0.83	0.44		1	99	
Peptococcus niger		0.03	0.03		99	99	
40040040	**Low discrimination**						**40040040**
					SPO		
Clostridium botulinum biovar A		83.14	0.41	GLU 90	99		
Clostridium sporogenes		3.28	0.14	CEL 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		2.27	0.21		1		
40041000	**Low discrimination**						**40041000**
					SPO		
Atopobium minutum		81.42	0.47	CEL 1	1		
Clostridium sporogenes		3.57	0.14		99		
Clostridium botulinum biovar A		3.06	0.15		99		
40041002	**Low discrimination**						**40041002**
					GRA	SPO	
Atopobium minutum		86.45	0.44	CEL 1	99	1	
Pseudoflavonifractor capillosus		11.65	0.15		1	1	
Clostridium sporogenes		1.90	0.06		99	99	
40041020	**Low discrimination**						**40041020**
					GRA		
Atopobium minutum		78.77	0.39	CEL 1	99		
Pseudoflavonifractor capillosus		21.23	0.15		1		
40041022	**Good identification**						**40041022**
					GRA		
Pseudoflavonifractor capillosus		97.56	0.50	CEL 1	1		
Atopobium minutum		2.44	0.36	GAL 10	99		
40050000	**Low discrimination**						**40050000**
					SPO		
Atopobium minutum		68.04	0.55	CEL 1	1		
Clostridium sporogenes		18.85	0.35		99		
Clostridium botulinum biovar A		5.84	0.29		99		
40050002	**Low discrimination**						**40050002**
					GRA	SPO	
Atopobium minutum		84.24	0.52	CEL 1	99	1	
Clostridium sporogenes		11.67	0.27		99	99	
Pseudoflavonifractor capillosus		4.10	0.15		1	1	
40050020	**Acceptable identification**						**40050020**
					GRA	SPO	
Atopobium minutum		90.44	0.47	CEL 1	99	1	
Pseudoflavonifractor capillosus		8.80	0.15		1	1	
Clostridium sporogenes		0.76	0.01		99	99	
40050022	**Acceptable identification**						**40050022**
					GRA		
Pseudoflavonifractor capillosus		93.53	0.50	FRU 10	1		
Atopobium minutum		6.47	0.44	CEL 1	99		
40051000	**Good identification**						**40051000**
					GRA	SPO	
Atopobium minutum		97.11	0.47	CEL 1	99	1	
Fusobacterium mortiferum		2.08	0.24		1	1	
Clostridium sporogenes		0.82	0.01		99	99	
40051002	**Good identification**						**40051002**
					GRA		
Atopobium minutum		98.40	0.44	CEL 1	99		
Pseudoflavonifractor capillosus		1.60	-.01		1		
40051020	**Good identification**						**40051020**
					GRA		
Atopobium minutum		96.86	0.39	CEL 1	99		
Pseudoflavonifractor capillosus		3.14	-.01		1		
40052010	**Good identification**						**40052010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium difficile		99.68	0.39	GLU 99	99		
Eubacterium limosum		0.32	-.06	CEL 5	1		
40060000	**Low discrimination**						**40060000**
					SPO		
Clostridium sporogenes		56.88	0.40	CEL 1	99		
Clostridium botulinum biovar A		36.52	0.39		99		
Atopobium minutum		1.19	0.21		1		
40100000	**Low discrimination**						**40100000**
					COC	SPO	
Clostridium sporogenes		67.67	0.87		1	99	
Clostridium botulinum biovar A		25.53	0.82		1	99	
Peptostreptococcus anaerobius		2.89	0.61		99	1	
40100001	**Low discrimination**						**40100001**
					GRA	SPO	
Clostridium sporogenes		54.14	0.53	URE 1	99	99	
Clostridium botulinum biovar A		20.42	0.48		99	99	
Campylobacter ureolyticus		8.29	0.32		1	1	
40100002	**Very good identification**						**40100002**
					SPO		
Clostridium sporogenes		97.30	0.79		99		
Clostridium botulinum biovar A		1.11	0.48		99		
Atopobium minutum		0.73	0.53		1		
40100003	**Low discrimination**						**40100003**
					COC	SPO	
Clostridium sporogenes		87.77	0.45	URE 1	1	99	
Clostridium novyi biovar A		6.96	0.40		1	99	
Anaerococcus prevotii		2.04	0.13		99	1	
40100010	**Low discrimination**						**40100010**
Clostridium sporogenes		61.53	0.53	MLZ 1			
Clostridium botulinum biovar A		23.21	0.48				
Clostridium novyi biovar A		9.22	0.53				
40100011	**Acceptable identification**						**40100011**
Clostridium novyi biovar A		90.41	0.53	GLU 99			
Clostridium sporogenes		6.10	0.19				
Clostridium botulinum biovar A		2.30	0.14				
40100012	**Acceptable identification**						**40100012**
Clostridium sporogenes		90.70	0.45	MLZ 1			
Clostridium novyi biovar A		7.19	0.40				
Clostridium botulinum biovar A		1.04	0.14				
40100013	**Low discrimination**						**40100013**
Clostridium novyi biovar A		88.70	0.40	GLU 99			
Clostridium sporogenes		11.30	0.11				
40100020	**Low discrimination**						**40100020**
Clostridium sporogenes		51.57	0.53	NAG 1			
Clostridium botulinum biovar A		19.45	0.48				
Clostridium bifermentans		11.54	0.44				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
40100021	**Low discrimination**						**40100021**
Clostridium novyi biovar A		62.71	0.43	GLU 99			
Clostridium sporogenes		16.92	0.19				
Clostridium botulinum biovar A		6.38	0.14				
40100022	**Very good identification**						**40100022**
					GRA	SPO	
Pseudoflavonifractor capillosus		95.82	0.66	MLT 1	1	1	
Clostridium sporogenes		3.22	0.45		99	99	
Atopobium minutum		0.80	0.44		99	1	
40100030	**Low discrimination**						**40100030**
Clostridium novyi biovar A		66.50	0.43	GLU 99			
Clostridium sporogenes		17.94	0.19				
Clostridium botulinum biovar A		6.77	0.14				
40100031	**Good identification**						**40100031**
Clostridium novyi biovar A		100.0	0.43	GLU 99			
40100040	**Low discrimination**						**40100040**
Clostridium botulinum biovar A		89.29	0.73	GLU 90			
Clostridium sporogenes		8.01	0.53				
Clostridium bifermentans		1.17	0.40				
40100041	**Low discrimination**						**40100041**
Clostridium botulinum biovar A		49.63	0.39	GLU 90			
Clostridium novyi biovar A		43.98	0.50	URE 1			
Clostridium sporogenes		4.45	0.19				
40100042	**Low discrimination**						**40100042**
Clostridium sporogenes		71.07	0.45	SOR 1			
Clostridium botulinum biovar A		24.02	0.39				
Clostridium novyi biovar A		3.76	0.37				
40100043	**Low discrimination**						**40100043**
Clostridium novyi biovar A		79.64	0.37	GLU 99			
Clostridium sporogenes		15.22	0.11				
Clostridium botulinum biovar A		5.14	0.05				
40100050	**Low discrimination**						**40100050**
Clostridium botulinum biovar A		50.18	0.39	GLU 90			
Clostridium novyi biovar A		44.47	0.50	MLZ 1			
Clostridium sporogenes		4.50	0.19				
40100051	**Good identification**						**40100051**
Clostridium novyi biovar A		98.87	0.50	GLU 99			
Clostridium botulinum biovar A		1.13	0.05				
40100052	**Low discrimination**						**40100052**
Clostridium novyi biovar A		79.64	0.37	GLU 99			
Clostridium sporogenes		15.22	0.11				
Clostridium botulinum biovar A		5.14	0.05				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40100053	**Good identification**						**40100053**
Clostridium novyi biovar A		100.0	0.37	GLU 99			
40100060	**Low discrimination**						**40100060**
Clostridium botulinum biovar A		70.53	0.39	GLU 90			
Clostridium novyi biovar A		15.63	0.40	NAG 1			
Clostridium sporogenes		6.32	0.19				
40100061	**Good identification**						**40100061**
Clostridium novyi biovar A		95.64	0.40	GLU 99			
Clostridium botulinum biovar A		4.36	0.05				
40100070	**Good identification**						**40100070**
Clostridium novyi biovar A		95.64	0.40	GLU 99			
Clostridium botulinum biovar A		4.36	0.05				
40100071	**Good identification**						**40100071**
Clostridium novyi biovar A		100.0	0.40	GLU 99			
40100100	**Low discrimination**						**40100100**
					COC	SPO	
Clostridium sporogenes		67.67	0.53	LAC 1	1	99	
Clostridium botulinum biovar A		25.52	0.48		1	99	
Peptostreptococcus anaerobius		2.89	0.27		99	1	
40100102	**Good identification**						**40100102**
					SPO		
Clostridium sporogenes		97.73	0.45	LAC 1	99		
Clostridium botulinum biovar A		1.12	0.14		99		
Atopobium minutum		0.73	0.18		1		
40100140	**Low discrimination**						**40100140**
Clostridium botulinum biovar A		89.64	0.39	GLU 90			
Clostridium sporogenes		8.04	0.19	LAC 1			
Clostridium bifermentans		1.17	0.06				
40100200	**Low discrimination**						**40100200**
					COC	SPO	
Clostridium sporogenes		67.70	0.53	RHA 1	1	99	
Clostridium botulinum biovar A		25.54	0.48		1	99	
Peptostreptococcus anaerobius		2.89	0.27		99	1	
40100202	**Good identification**						**40100202**
					SPO		
Clostridium sporogenes		97.73	0.45	RHA 1	99		
Clostridium botulinum biovar A		1.12	0.14		99		
Atopobium minutum		0.73	0.18		1		
40100240	**Low discrimination**						**40100240**
Clostridium botulinum biovar A		89.64	0.39	GLU 90			
Clostridium sporogenes		8.04	0.19	RHA 1			
Clostridium bifermentans		1.17	0.06				
40100400	**Low discrimination**						**40100400**
					COC	SPO	
Clostridium sporogenes		67.70	0.53	ARA 1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		25.54	0.48		1	99	
Peptostreptococcus anaerobius		2.89	0.27		99	1	
40100402	**Good identification**						**40100402**
					SPO		
Clostridium sporogenes		97.73	0.45	ARA 1	99		
Clostridium botulinum biovar A		1.12	0.14		99		
Atopobium minutum		0.73	0.18		1		
40100440	**Low discrimination**						**40100440**
Clostridium botulinum biovar A		89.64	0.39	GLU 90			
Clostridium sporogenes		8.04	0.19	ARA 1			
Clostridium bifermentans		1.17	0.06				
40101000	**Low discrimination**						**40101000**
					SPO		
Clostridium sporogenes		61.95	0.53	GAL 1	99		
Clostridium botulinum biovar A		23.37	0.48		99		
Atopobium minutum		7.68	0.47		1		
40101001	**Low discrimination**						**40101001**
Clostridium novyi biovar A		44.09	0.37	GLU 99			
Clostridium sporogenes		26.76	0.19	GAL 10			
Clostridium sordellii		10.50	0.09				
40101002	**Low discrimination**						**40101002**
					GRA	SPO	
Clostridium sporogenes		76.76	0.45	GAL 1	99	99	
Atopobium minutum		19.02	0.44		99	1	
Pseudoflavonifractor capillosus		2.56	0.15		1	1	
40101010	**Low discrimination**						**40101010**
Clostridium novyi biovar A		51.62	0.37	GLU 99			
Clostridium sporogenes		31.34	0.19	GAL 10			
Clostridium botulinum biovar A		11.82	0.14				
40101011	**Good identification**						**40101011**
Clostridium novyi biovar A		100.0	0.37	GLU 99			
40101020	**Low discrimination**						**40101020**
					GRA	SPO	
Atopobium minutum		56.43	0.39	MLT 1	99	1	
Pseudoflavonifractor capillosus		15.21	0.15		1	1	
Clostridium sporogenes		13.80	0.19		99	99	
40101022	**Good identification**						**40101022**
					GRA	SPO	
Pseudoflavonifractor capillosus		97.21	0.50	MLT 1	1	1	
Atopobium minutum		2.43	0.36	GAL 10	99	1	
Clostridium sporogenes		0.30	0.11		99	99	
40101040	**Low discrimination**						**40101040**
Clostridium botulinum biovar A		80.40	0.39	GLU 90			
Clostridium novyi biovar A		7.92	0.34	GAL 1			
Clostridium sporogenes		7.21	0.19				
40102000	**Low discrimination**						**40102000**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		67.64	0.53	MAN 1	1	99	
Clostridium botulinum biovar A		25.51	0.48		1	99	
Peptostreptococcus anaerobius		2.89	0.27		99	1	
40102002	**Good identification**						**40102002**
					SPO		
Clostridium sporogenes		97.73	0.45	MAN 1	99		
Clostridium botulinum biovar A		1.12	0.14		99		
Atopobium minutum		0.73	0.18		1		
40102040	**Low discrimination**						**40102040**
Clostridium botulinum biovar A		89.25	0.39	GLU 90			
Clostridium sporogenes		8.00	0.19	MAN 1			
Clostridium bifermentans		1.17	0.06				
40104000	**Low discrimination**						**40104000**
					COC	SPO	
Clostridium sporogenes		66.44	0.53	XYL 1	1	99	
Clostridium botulinum biovar A		25.06	0.48		1	99	
Peptostreptococcus anaerobius		2.84	0.27		99	1	
40104002	**Good identification**						**40104002**
					SPO		
Clostridium sporogenes		97.73	0.45	XYL 1	99		
Clostridium botulinum biovar A		1.12	0.14		99		
Atopobium minutum		0.73	0.18		1		
40104040	**Low discrimination**						**40104040**
Clostridium glycolicum		80.92	0.50	GLU 99			
Clostridium botulinum biovar A		17.11	0.39	FRU 90			
Clostridium sporogenes		1.53	0.19				
40110000	**Low discrimination**						**40110000**
					SPO		
Clostridium sporogenes		85.06	0.74		99		
Clostridium botulinum biovar A		11.60	0.62		99		
Atopobium minutum		1.67	0.55		1		
40110001	**Low discrimination**						**40110001**
Clostridium sporogenes		71.46	0.40	URE 1			
Clostridium botulinum biovar A		9.75	0.28				
Clostridium sordellii		9.10	0.22				
40110002	**Very good identification**						**40110002**
					SPO		
Clostridium sporogenes		95.56	0.66		99		
Atopobium minutum		3.75	0.52		1		
Clostridium botulinum biovar A		0.40	0.28		99		
40110010	**Low discrimination**						**40110010**
Clostridium sporogenes		77.94	0.40	MLZ 1			
Clostridium botulinum biovar A		10.63	0.28				
Clostridium novyi biovar A		7.35	0.37				
40110011	**Acceptable identification**						**40110011**
Clostridium novyi biovar A		90.33	0.37	GLU 99			
Clostridium sporogenes		9.68	0.06	FRU 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40110020	**Low discrimination**						**40110020**
					SPO		
Clostridium sporogenes		51.54	0.40	NAG 1	99		
Atopobium minutum		33.40	0.47		1		
Clostridium botulinum biovar A		7.03	0.28		99		
40110022	**Low discrimination**						**40110022**
					GRA	SPO	
Pseudoflavonifractor capillosus		89.03	0.50	MLT 1	1	1	
Atopobium minutum		6.16	0.44	FRU 10	99	1	
Clostridium sporogenes		4.75	0.32		99	99	
40110040	**Low discrimination**						**40110040**
Clostridium botulinum biovar A		60.64	0.53	GLU 90			
Clostridium glycolicum		21.50	0.44	FRU 6			
Clostridium sporogenes		15.03	0.40				
40110100	**Low discrimination**						**40110100**
					SPO		
Clostridium sporogenes		83.03	0.40	LAC 1	99		
Clostridium botulinum biovar A		11.33	0.28		99		
Actinomyces meyeri		2.27	0.19		1		
40110200	**Low discrimination**						**40110200**
					SPO		
Clostridium sporogenes		85.54	0.40	RHA 1	99		
Clostridium botulinum biovar A		11.67	0.28		99		
Atopobium minutum		1.68	0.21		1		
40110400	**Low discrimination**						**40110400**
					SPO		
Clostridium sporogenes		85.54	0.40	ARA 1	99		
Clostridium botulinum biovar A		11.67	0.28		99		
Atopobium minutum		1.68	0.21		1		
40111000	**Low discrimination**						**40111000**
					SPO		
Clostridium sporogenes		53.09	0.40	GAL 1	99		
Atopobium minutum		34.41	0.47		1		
Clostridium botulinum biovar A		7.24	0.28		99		
40111002	**Low discrimination**						**40111002**
					GRA	SPO	
Atopobium minutum		55.94	0.44	MLT 1	99	1	
Clostridium sporogenes		43.16	0.32		99	99	
Pseudoflavonifractor capillosus		0.91	-.01		1	1	
40111020	**Acceptable identification**						**40111020**
					GRA	SPO	
Atopobium minutum		91.63	0.39	MLT 1	99	1	
Clostridium sporogenes		4.29	0.06		99	99	
Pseudoflavonifractor capillosus		2.97	-.01		1	1	
40112000	**Low discrimination**						**40112000**
					SPO		
Clostridium sporogenes		63.03	0.40	MAN 1	99		
Eubacterium limosum		18.24	0.29		1		
Clostridium botulinum biovar A		8.60	0.28		99		
40114000	**Low discrimination**						**40114000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium glycolicum		54.75	0.44	GLU 99			
Clostridium sporogenes		38.27	0.40	SOR 95			
Clostridium botulinum biovar A		5.22	0.28				
40114040	**Very good identification**						**40114040**
Clostridium glycolicum		99.81	0.66	GLU 99			
Clostridium botulinum biovar A		0.15	0.19				
Clostridium sporogenes		0.04	0.06				
40115040	**Good identification**						**40115040**
Clostridium glycolicum		100.0	0.50	GLU 99			
40120000	**Low discrimination**						**40120000**
					COC	SPO	
Clostridium sporogenes		77.77	0.79		1	99	
Clostridium botulinum biovar A		22.00	0.72		1	99	
Peptostreptococcus anaerobius		0.10	0.27		99	1	
40120001	**Low discrimination**						**40120001**
					GRA	SPO	
Clostridium sporogenes		77.32	0.45	URE 1	99	99	
Clostridium botulinum biovar A		21.87	0.38		99	99	
Campylobacter ureolyticus		0.36	-.03		1	1	
40120002	**Very good identification**						**40120002**
					SPO		
Clostridium sporogenes		99.12	0.71		99		
Clostridium botulinum biovar A		0.85	0.38		99		
Atopobium minutum		0.02	0.18		1		
40120003	**Good identification**						**40120003**
Clostridium sporogenes		99.15	0.37	URE 1			
Clostridium botulinum biovar A		0.85	0.04				
40120010	**Low discrimination**						**40120010**
Clostridium sporogenes		77.60	0.45	MLZ 1			
Clostridium botulinum biovar A		21.95	0.38				
Clostridium novyi biovar A		0.35	0.19				
40120012	**Good identification**						**40120012**
Clostridium sporogenes		99.15	0.37	MLZ 1			
Clostridium botulinum biovar A		0.85	0.04				
40120020	**Low discrimination**						**40120020**
Clostridium sporogenes		77.06	0.45	NAG 1			
Clostridium botulinum biovar A		21.80	0.38				
Clostridium bifermentans		0.52	0.09				
40120022	**Low discrimination**						**40120022**
					GRA	SPO	
Clostridium sporogenes		52.15	0.37	NAG 1	99	99	
Pseudoflavonifractor capillosus		47.01	0.32		1	1	
Clostridium botulinum biovar A		0.45	0.04		99	99	
40120040	**Low discrimination**						**40120040**
Clostridium botulinum biovar A		89.24	0.63	GLU 90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		10.67	0.45				
Clostridium bifermentans		0.05	0.06				
40120042	**Low discrimination**						**40120042**
Clostridium sporogenes		79.78	0.37	SOR 1			
Clostridium botulinum biovar A		20.22	0.29				
40120100	**Low discrimination**						**40120100**
					COC	SPO	
Clostridium sporogenes		77.87	0.45	LAC 1	1	99	
Clostridium botulinum biovar A		22.03	0.38		1	99	
Peptostreptococcus anaerobius		0.10	-.07		99	1	
40120102	**Good identification**						**40120102**
Clostridium sporogenes		99.15	0.37	LAC 1			
Clostridium botulinum biovar A		0.85	0.04				
40120200	**Low discrimination**						**40120200**
					COC	SPO	
Clostridium sporogenes		77.87	0.45	RHA 1	1	99	
Clostridium botulinum biovar A		22.03	0.38		1	99	
Peptostreptococcus anaerobius		0.10	-.07		99	1	
40120202	**Good identification**						**40120202**
Clostridium sporogenes		99.15	0.37	RHA 1			
Clostridium botulinum biovar A		0.85	0.04				
40120400	**Low discrimination**						**40120400**
					COC	SPO	
Clostridium sporogenes		77.87	0.45	ARA 1	1	99	
Clostridium botulinum biovar A		22.03	0.38		1	99	
Peptostreptococcus anaerobius		0.10	-.07		99	1	
40120402	**Good identification**						**40120402**
Clostridium sporogenes		99.15	0.37	ARA 1			
Clostridium botulinum biovar A		0.85	0.04				
40121000	**Low discrimination**						**40121000**
					SPO		
Clostridium sporogenes		77.64	0.45	GAL 1	99		
Clostridium botulinum biovar A		21.97	0.38		99		
Atopobium minutum		0.29	0.13		1		
40121002	**Good identification**						**40121002**
					SPO		
Clostridium sporogenes		98.42	0.37	GAL 1	99		
Clostridium botulinum biovar A		0.84	0.04		99		
Atopobium minutum		0.74	0.10		1		
40122000	**Low discrimination**						**40122000**
					COC	SPO	
Clostridium sporogenes		77.87	0.45	MAN 1	1	99	
Clostridium botulinum biovar A		22.03	0.38		1	99	
Peptostreptococcus anaerobius		0.10	-.07		99	1	
40122002	**Good identification**						**40122002**
Clostridium sporogenes		99.15	0.37	MAN 1			
Clostridium botulinum biovar A		0.85	0.04				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40124000	**Low discrimination**						**40124000**
					COC	SPO	
Clostridium sporogenes		77.87	0.45	XYL 1	1	99	
Clostridium botulinum biovar A		22.03	0.38		1	99	
Peptostreptococcus anaerobius		0.10	-.07		99	1	
40124002	**Good identification**						**40124002**
Clostridium sporogenes		99.15	0.37	XYL 1			
Clostridium botulinum biovar A		0.85	0.04				
40130000	**Good identification**						**40130000**
					SPO		
Clostridium sporogenes		90.64	0.66		99		
Clostridium botulinum biovar A		9.27	0.51		99		
Atopobium minutum		0.05	0.21		1		
40130002	**Very good identification**						**40130002**
					SPO		
Clostridium sporogenes		99.57	0.58		99		
Clostridium botulinum biovar A		0.31	0.17		99		
Atopobium minutum		0.12	0.18		1		
40130040	**Low discrimination**						**40130040**
Clostridium botulinum biovar A		74.36	0.42	GLU 90			
Clostridium sporogenes		24.58	0.32	FRU 6			
Clostridium glycolicum		1.07	0.10				
40140000	**Low discrimination**						**40140000**
					COC	SPO	
Clostridium sporogenes		67.70	0.53	CEL 1	1	99	
Clostridium botulinum biovar A		25.54	0.48		1	99	
Peptostreptococcus anaerobius		2.89	0.27		99	1	
40140002	**Good identification**						**40140002**
					SPO		
Clostridium sporogenes		97.73	0.45	CEL 1	99		
Clostridium botulinum biovar A		1.12	0.14		99		
Atopobium minutum		0.73	0.18		1		
40140040	**Low discrimination**						**40140040**
Clostridium botulinum biovar A		89.64	0.39	GLU 90			
Clostridium sporogenes		8.04	0.19	CEL 1			
Clostridium bifermentans		1.17	0.06				
40150000	**Low discrimination**						**40150000**
					SPO		
Clostridium sporogenes		85.54	0.40	CEL 1	99		
Clostridium botulinum biovar A		11.67	0.28		99		
Atopobium minutum		1.68	0.21		1		
40160000	**Low discrimination**						**40160000**
					COC	SPO	
Clostridium sporogenes		77.87	0.45	CEL 1	1	99	
Clostridium botulinum biovar A		22.03	0.38		1	99	
Peptostreptococcus anaerobius		0.10	-.07		99	1	
40160002	**Good identification**						**40160002**
Clostridium sporogenes		99.15	0.37	CEL 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		0.85	0.04				
40200000	**Low discrimination**						**40200000**
					SPO		
Clostridium sporogenes		16.42	0.48	SAL 1	99		
Clostridium botulinum biovar A		14.06	0.50		99		
Atopobium minutum		11.34	0.56		1		
40200002	**Low discrimination**						**40200002**
					GRA	SPO	
Atopobium minutum		39.70	0.53	SAL 1	99	1	
Clostridium sporogenes		28.76	0.40		99	99	
Pseudoflavonifractor capillosus		16.05	0.32		1	1	
40200020	**Low discrimination**						**40200020**
					COC	GRA	
Atopobium minutum		43.83	0.47	SAL 1	1	99	
Pseudoflavonifractor capillosus		35.44	0.32		1	1	
Peptococcus niger		6.63	0.15		99	99	
40200022	**Very good identification**						**40200022**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	SAL 1	1	1	
Atopobium minutum		0.83	0.44		1	99	
Peptococcus niger		0.03	0.03		99	99	
40200040	**Low discrimination**						**40200040**
Clostridium botulinum biovar A		76.66	0.41	GLU 90			
Clostridium novyi biovar A		4.83	0.32	SAL 1			
Clostridium bifermentans		3.87	0.18				
40201000	**Low discrimination**						**40201000**
					SPO		
Atopobium minutum		55.56	0.47	SAL 1	1		
Eubacterium tortuosum		30.36	0.32		1		
Clostridium sporogenes		2.44	0.14		99		
40201002	**Low discrimination**						**40201002**
					GRA		
Atopobium minutum		76.84	0.44	SAL 1	99		
Eubacterium tortuosum		11.11	0.19		99		
Pseudoflavonifractor capillosus		10.36	0.15		1		
40201020	**Low discrimination**						**40201020**
					GRA		
Atopobium minutum		64.15	0.39	SAL 1	99		
Eubacterium tortuosum		18.56	0.19		99		
Pseudoflavonifractor capillosus		17.29	0.15		1		
40201022	**Good identification**						**40201022**
					GRA		
Pseudoflavonifractor capillosus		97.38	0.50	SAL 1	1		
Atopobium minutum		2.43	0.36	GAL 10	99		
Eubacterium tortuosum		0.19	0.06		99		
40210000	**Low discrimination**						**40210000**
					SPO		
Atopobium minutum		63.82	0.55	SAL 1	1		
Clostridium sporogenes		17.67	0.35		99		
Clostridium botulinum biovar A		5.47	0.29		99		
40210002	**Low discrimination**						**40210002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Atopobium minutum		83.91	0.52	SAL 1	99	1	
Clostridium sporogenes		11.62	0.27		99	99	
Pseudoflavonifractor capillosus		4.08	0.15		1	1	
40210010	**Acceptable identification**						**40210010**
					SPO		
Clostridium difficile		94.27	0.35	GLU 99	99		
Atopobium minutum		3.38	0.21	MAN 90	1		
Clostridium novyi biovar A		1.41	0.19		99		
40210020	**Acceptable identification**						**40210020**
					GRA	SPO	
Atopobium minutum		90.04	0.47	SAL 1	99	1	
Pseudoflavonifractor capillosus		8.76	0.15		1	1	
Clostridium sporogenes		0.76	0.01		99	99	
40210022	**Acceptable identification**						**40210022**
					GRA		
Pseudoflavonifractor capillosus		93.53	0.50	SAL 1	1		
Atopobium minutum		6.47	0.44	FRU 10	99		
40210100	**Low discrimination**						**40210100**
					GRA	SPO	
Fusobacterium mortiferum		83.36	0.40	GLU 99	1	1	
Atopobium minutum		13.03	0.21		99	1	
Clostridium sporogenes		3.61	0.01		99	99	
40211000	**Low discrimination**						**40211000**
					GRA	SPO	
Atopobium minutum		87.80	0.47	SAL 1	99	1	
Fusobacterium mortiferum		10.94	0.37		1	1	
Clostridium sporogenes		0.74	0.01		99	99	
40211002	**Good identification**						**40211002**
					GRA		
Atopobium minutum		96.43	0.44	SAL 1	99		
Fusobacterium mortiferum		2.00	0.20		1		
Pseudoflavonifractor capillosus		1.56	-.01		1		
40211020	**Good identification**						**40211020**
					GRA		
Atopobium minutum		96.86	0.39	SAL 1	99		
Pseudoflavonifractor capillosus		3.14	-.01		1		
40211100	**Good identification**						**40211100**
					GRA		
Fusobacterium mortiferum		96.64	0.43	GLU 99	1		
Atopobium minutum		3.36	0.13		99		
40212000	**Low discrimination**						**40212000**
					SPO		
Clostridium difficile		76.06	0.40	GLU 99	99		
Eubacterium limosum		21.82	0.29		1		
Atopobium minutum		1.48	0.21		1		
40212010	**Good identification**						**40212010**
					SPO		
Clostridium difficile		99.94	0.52	GLU 99	99		
Eubacterium limosum		0.06	-.06		1		
40220000	**Low discrimination**						**40220000**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		56.57	0.40	SAL 1	99		
Clostridium botulinum biovar A		36.33	0.39		99		
Atopobium minutum		1.18	0.21		1		
40300000	**Low discrimination**						**40300000**
Clostridium sporogenes		54.27	0.53	SAL 1			
Clostridium bifermentans		21.13	0.48				
Clostridium botulinum biovar A		20.47	0.48				
40300002	**Good identification**						**40300002**
Clostridium sporogenes		95.97	0.45	SAL 1			
Clostridium bifermentans		1.13	0.14				
Clostridium botulinum biovar A		1.10	0.14				
40300040	**Low discrimination**						**40300040**
Clostridium botulinum biovar A		76.07	0.39	GLU 90			
Clostridium bifermentans		10.96	0.24	SAL 1			
Clostridium sporogenes		6.82	0.19				
40310000	**Low discrimination**						**40310000**
Clostridium sporogenes		77.55	0.40	SAL 1			
Clostridium botulinum biovar A		10.58	0.28				
Clostridium bifermentans		9.01	0.26				
40320000	**Low discrimination**						**40320000**
Clostridium sporogenes		77.16	0.45	SAL 1			
Clostridium botulinum biovar A		21.83	0.38				
Clostridium bifermentans		0.91	0.14				
40320002	**Good identification**						**40320002**
Clostridium sporogenes		99.15	0.37	SAL 1			
Clostridium botulinum biovar A		0.85	0.04				
40400000	**Low discrimination**						**40400000**
					SPO		
Clostridium sporogenes		16.70	0.48	RAF 1	99		
Clostridium botulinum biovar A		14.30	0.50		99		
Atopobium minutum		11.53	0.56		1		
40400001	**Low discrimination**						**40400001**
					COC	GRA	
Anaerococcus prevotii		83.61	0.50	ESL 1	99	99	
Campylobacter ureolyticus		12.52	0.32		1	1	
Peptococcus niger		1.36	0.15		99	99	
40400002	**Low discrimination**						**40400002**
					GRA	SPO	
Atopobium minutum		38.25	0.53	RAF 1	99	1	
Clostridium sporogenes		27.71	0.40		99	99	
Pseudoflavonifractor capillosus		15.47	0.32		1	1	
40400003	**Good identification**						**40400003**
					COC		
Anaerococcus prevotii		96.30	0.39	ESL 1	99		
Peptococcus niger		1.11	0.02		99		
Atopobium minutum		0.94	0.18		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40400020	**Low discrimination**						**40400020**
					COC	GRA	
Atopobium minutum		45.74	0.47	RAF 1	1	99	
Pseudoflavonifractor capillosus		36.98	0.32		1	1	
Peptococcus niger		6.92	0.15		99	99	
40400022	**Very good identification**						**40400022**
					COC	GRA	
Pseudoflavonifractor capillosus		99.12	0.66	RAF 1	1	1	
Atopobium minutum		0.83	0.44		1	99	
Peptococcus niger		0.03	0.03		99	99	
40400040	**Low discrimination**						**40400040**
					SPO		
Clostridium botulinum biovar A		82.57	0.41	GLU 90	99		
Clostridium sporogenes		3.26	0.14	RAF 1	99		
Atopobium minutum		2.25	0.21		1		
40401000	**Low discrimination**						**40401000**
					SPO		
Atopobium minutum		80.42	0.47	RAF 1	1		
Clostridium sporogenes		3.53	0.14		99		
Clostridium botulinum biovar A		3.02	0.15		99		
40401002	**Low discrimination**						**40401002**
					GRA	SPO	
Atopobium minutum		86.45	0.44	RAF 1	99	1	
Pseudoflavonifractor capillosus		11.65	0.15		1	1	
Clostridium sporogenes		1.90	0.06		99	99	
40401020	**Low discrimination**						**40401020**
					GRA		
Atopobium minutum		78.77	0.39	RAF 1	99		
Pseudoflavonifractor capillosus		21.23	0.15		1		
40401022	**Good identification**						**40401022**
					GRA		
Pseudoflavonifractor capillosus		97.56	0.50	RAF 1	1		
Atopobium minutum		2.44	0.36	GAL 10	99		
40410000	**Low discrimination**						**40410000**
					SPO		
Atopobium minutum		67.24	0.55	RAF 1	1		
Clostridium sporogenes		18.62	0.35		99		
Clostridium botulinum biovar A		5.77	0.29		99		
40410002	**Low discrimination**						**40410002**
					GRA	SPO	
Atopobium minutum		83.91	0.52	RAF 1	99	1	
Clostridium sporogenes		11.62	0.27		99	99	
Pseudoflavonifractor capillosus		4.08	0.15		1	1	
40410020	**Acceptable identification**						**40410020**
					GRA	SPO	
Atopobium minutum		90.44	0.47	RAF 1	99	1	
Pseudoflavonifractor capillosus		8.80	0.15		1	1	
Clostridium sporogenes		0.76	0.01		99	99	
40410022	**Acceptable identification**						**40410022**
					GRA		
Pseudoflavonifractor capillosus		93.53	0.50	RAF 1	1		
Atopobium minutum		6.47	0.44	FRU 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
40410100	**Low discrimination**						**40410100**
					GRA	SPO	
Fusobacterium mortiferum		83.36	0.40	GLU 99	1	1	
Atopobium minutum		13.03	0.21		99	1	
Clostridium sporogenes		3.61	0.01		99	99	
40411000	**Low discrimination**						**40411000**
					GRA	SPO	
Atopobium minutum		88.26	0.47	RAF 1	99	1	
Fusobacterium mortiferum		11.00	0.37		1	1	
Clostridium sporogenes		0.74	0.01		99	99	
40411002	**Good identification**						**40411002**
					GRA		
Atopobium minutum		96.43	0.44	RAF 1	99		
Fusobacterium mortiferum		2.00	0.20		1		
Pseudoflavonifractor capillosus		1.56	-.01		1		
40411020	**Good identification**						**40411020**
					GRA		
Atopobium minutum		96.86	0.39	RAF 1	99		
Pseudoflavonifractor capillosus		3.14	-.01		1		
40411100	**Acceptable identification**						**40411100**
					COC	GRA	
Fusobacterium mortiferum		94.59	0.43	GLU 99	1	1	
Atopobium minutum		3.29	0.13		1	99	
Blautia hansenii		2.12	-.03		99	99	
40420000	**Low discrimination**						**40420000**
					SPO		
Clostridium sporogenes		56.67	0.40	RAF 1	99		
Clostridium botulinum biovar A		36.39	0.39		99		
Atopobium minutum		1.19	0.21		1		
40500000	**Low discrimination**						**40500000**
					COC	SPO	
Clostridium sporogenes		67.62	0.53	RAF 1	1	99	
Clostridium botulinum biovar A		25.50	0.48		1	99	
Peptostreptococcus anaerobius		2.89	0.27		99	1	
40500002	**Good identification**						**40500002**
					SPO		
Clostridium sporogenes		97.73	0.45	RAF 1	99		
Clostridium botulinum biovar A		1.12	0.14		99		
Atopobium minutum		0.73	0.18		1		
40500040	**Low discrimination**						**40500040**
Clostridium botulinum biovar A		89.64	0.39	GLU 90			
Clostridium sporogenes		8.04	0.19	RAF 1			
Clostridium bifermentans		1.17	0.06				
40501100	**Very good identification**						**40501100**
Blautia hansenii		100.0	0.66	GLU 99			
40501102	**Good identification**						**40501102**
Blautia hansenii		100.0	0.50	GLU 99			
40501120	**Good identification**						**40501120**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		99.15	0.50	GLU 99	99	99	
Prevotella melaninogenica		0.85	0.14	NAG 10	1	1	
40501500	**Good identification**						**40501500**
Blautia hansenii		100.0	0.53	GLU 99			
40501502	**Good identification**						**40501502**
Blautia hansenii		100.0	0.37	GLU 99			
40501520	**Good identification**						**40501520**
Blautia hansenii		100.0	0.37	GLU 99			
40505100	**Good identification**						**40505100**
Blautia hansenii		100.0	0.53	GLU 99			
40505102	**Good identification**						**40505102**
Blautia hansenii		100.0	0.37	GLU 99			
40505120	**Good identification**						**40505120**
Blautia hansenii		100.0	0.37	GLU 99			
40505500	**Good identification**						**40505500**
Blautia hansenii		100.0	0.40	GLU 99			
40510000	**Low discrimination**						**40510000**
					GRA	SPO	
Clostridium sporogenes		79.79	0.40	RAF 1	99	99	
Clostridium botulinum biovar A		10.89	0.28		99	99	
Prevotella intermedia		6.19	0.25		1	1	
40520000	**Low discrimination**						**40520000**
					COC	SPO	
Clostridium sporogenes		77.87	0.45	RAF 1	1	99	
Clostridium botulinum biovar A		22.03	0.38		1	99	
Peptostreptococcus anaerobius		0.10	-.07		99	1	
40520002	**Good identification**						**40520002**
Clostridium sporogenes		99.15	0.37	RAF 1			
Clostridium botulinum biovar A		0.85	0.04				
40610000	**Low discrimination**						**40610000**
					GRA	SPO	
Fusobacterium mortiferum		80.65	0.39	GLU 99	1	1	
Atopobium minutum		15.15	0.21		99	1	
Clostridium sporogenes		4.20	0.01		99	99	
40610100	**Good identification**						**40610100**
Fusobacterium mortiferum		100.0	0.45	GLU 99			
40611000	**Good identification**						**40611000**
					GRA		
Fusobacterium mortiferum		95.99	0.42	GLU 99	1		
Atopobium minutum		4.01	0.13		99		
40611100	**Good identification**						**40611100**
Fusobacterium mortiferum		100.0	0.48	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
40630100	**Good identification**					**40630100**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
40631100	**Good identification**					**40631100**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
40650100	**Good identification**					**40650100**
Fusobacterium mortiferum		100.0	0.37	GLU 99		
40651100	**Good identification**					**40651100**
Fusobacterium mortiferum		100.0	0.40	GLU 99		
41000000	**Low discrimination**					**41000000**
					SPO	
Clostridium botulinum biovar A		50.20	1.00		99	
Clostridium sporogenes		36.91	0.95		99	
Atopobium minutum		7.66	0.94		1	
41000001	**Low discrimination**					**41000001**
Clostridium novyi biovar A		63.94	0.87			
Clostridium botulinum biovar A		17.89	0.66			
Clostridium sporogenes		13.16	0.61			
41000002	**Low discrimination**					**41000002**
					SPO	
Clostridium sporogenes		67.26	0.87		99	
Atopobium minutum		27.90	0.91		1	
Clostridium botulinum biovar A		2.77	0.66		99	
41000003	**Low discrimination**					**41000003**
					SPO	
Clostridium novyi biovar A		63.61	0.74		99	
Clostridium sporogenes		24.72	0.53		99	
Atopobium minutum		10.26	0.57		1	
41000010	**Low discrimination**					**41000010**
Clostridium novyi biovar A		63.68	0.87			
Clostridium botulinum biovar A		17.82	0.66			
Clostridium sporogenes		13.10	0.61			
41000011	**Very good identification**					**41000011**
Clostridium novyi biovar A		99.42	0.87			
Clostridium botulinum biovar A		0.28	0.32			
Clostridium sporogenes		0.21	0.27			
41000012	**Low discrimination**					**41000012**
					SPO	
Clostridium novyi biovar A		63.75	0.74		99	
Clostridium sporogenes		24.78	0.53		99	
Atopobium minutum		10.28	0.57		1	
41000013	**Very good identification**					**41000013**
					SPO	
Clostridium novyi biovar A		99.45	0.74		99	
Clostridium sporogenes		0.39	0.19		99	
Atopobium minutum		0.16	0.22		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41000020	**Low discrimination**						**41000020**
					SPO		
Atopobium minutum		62.80	0.86		1		
Clostridium botulinum biovar A		12.48	0.66		99		
Clostridium novyi biovar A		11.15	0.77		99		
41000021	**Good identification**						**41000021**
					SPO		
Clostridium novyi biovar A		92.47	0.77		99		
Atopobium minutum		5.26	0.51		1		
Clostridium botulinum biovar A		1.05	0.32		99		
41000022	**Low discrimination**						**41000022**
					GRA	SPO	
Atopobium minutum		54.45	0.83		99	1	
Pseudoflavonifractor capillosus		38.78	0.66		1	1	
Clostridium sporogenes		3.98	0.53		99	99	
41000023	**Low discrimination**						**41000023**
					GRA	SPO	
Clostridium novyi biovar A		72.26	0.64		99	99	
Atopobium minutum		15.54	0.48		99	1	
Pseudoflavonifractor capillosus		11.07	0.32		1	1	
41000030	**Good identification**						**41000030**
					SPO		
Clostridium novyi biovar A		92.48	0.77		99		
Atopobium minutum		5.26	0.51		1		
Clostridium botulinum biovar A		1.05	0.32		99		
41000031	**Very good identification**						**41000031**
					SPO		
Clostridium novyi biovar A		99.94	0.77		99		
Atopobium minutum		0.06	0.17		1		
41000032	**Low discrimination**						**41000032**
					GRA	SPO	
Clostridium novyi biovar A		72.26	0.64		99	99	
Atopobium minutum		15.54	0.48		99	1	
Pseudoflavonifractor capillosus		11.07	0.32		1	1	
41000033	**Very good identification**						**41000033**
					GRA	SPO	
Clostridium novyi biovar A		99.63	0.64		99	99	
Atopobium minutum		0.22	0.14		99	1	
Pseudoflavonifractor capillosus		0.15	-.03		1	1	
41000040	**Low discrimination**						**41000040**
Clostridium botulinum biovar A		87.82	0.91				
Clostridium novyi biovar A		7.08	0.84				
Clostridium sporogenes		2.18	0.61				
41000041	**Low discrimination**						**41000041**
Clostridium novyi biovar A		88.27	0.84				
Clostridium botulinum biovar A		11.07	0.57				
Clostridium sporogenes		0.28	0.27				
41000042	**Low discrimination**						**41000042**
Clostridium novyi biovar A		39.07	0.71				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		27.76	0.57				
Clostridium sporogenes		22.78	0.53				
41000043	**Very good identification**						**41000043**
Clostridium novyi biovar A		98.47	0.71				
Clostridium botulinum biovar A		0.71	0.23				
Clostridium sporogenes		0.58	0.19				
41000050	**Low discrimination**						**41000050**
Clostridium novyi biovar A		88.15	0.84				
Clostridium botulinum biovar A		11.05	0.57				
Clostridium sporogenes		0.28	0.27				
41000051	**Very good identification**						**41000051**
Clostridium novyi biovar A		99.87	0.84				
Clostridium botulinum biovar A		0.13	0.23				
41000052	**Very good identification**						**41000052**
Clostridium novyi biovar A		98.47	0.71				
Clostridium botulinum biovar A		0.71	0.23				
Clostridium sporogenes		0.58	0.19				
41000053	**Very good identification**						**41000053**
Clostridium novyi biovar A		100.0	0.71				
41000060	**Low discrimination**						**41000060**
					SPO		
Clostridium novyi biovar A		61.30	0.74		99		
Clostridium botulinum biovar A		30.74	0.57		99		
Atopobium minutum		5.23	0.51		1		
41000061	**Very good identification**						**41000061**
					SPO		
Clostridium novyi biovar A		99.39	0.74		99		
Clostridium botulinum biovar A		0.50	0.23		99		
Atopobium minutum		0.09	0.17		1		
41000062	**Low discrimination**						**41000062**
					GRA	SPO	
Clostridium novyi biovar A		62.24	0.61		99	99	
Atopobium minutum		20.07	0.48		99	1	
Pseudoflavonifractor capillosus		14.30	0.32		1	1	
41000063	**Very good identification**						**41000063**
					GRA	SPO	
Clostridium novyi biovar A		99.45	0.61		99	99	
Atopobium minutum		0.32	0.14		99	1	
Pseudoflavonifractor capillosus		0.23	-.03		1	1	
41000070	**Very good identification**						**41000070**
					SPO		
Clostridium novyi biovar A		99.39	0.74		99		
Clostridium botulinum biovar A		0.50	0.23		99		
Atopobium minutum		0.09	0.17		1		
41000071	**Very good identification**						**41000071**
Clostridium novyi biovar A		100.0	0.74				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41000072	**Very good identification**						**41000072**
					GRA	SPO	
Clostridium novyi biovar A		99.45	0.61		99	99	
Atopobium minutum		0.32	0.14		99	1	
Pseudoflavonifractor capillosus		0.23	-.03		1	1	
41000073	**Very good identification**						**41000073**
Clostridium novyi biovar A		100.0	0.61				
41000100	**Low discrimination**						**41000100**
					SPO		
Clostridium botulinum biovar A		50.11	0.66	LAC 1	99		
Clostridium sporogenes		36.85	0.61		99		
Atopobium minutum		7.64	0.60		1		
41000101	**Low discrimination**						**41000101**
Clostridium novyi biovar A		64.15	0.53	LAC 1			
Clostridium botulinum biovar A		17.95	0.32				
Clostridium sporogenes		13.20	0.27				
41000102	**Low discrimination**						**41000102**
					SPO		
Clostridium sporogenes		67.26	0.53	LAC 1	99		
Atopobium minutum		27.90	0.57		1		
Clostridium botulinum biovar A		2.77	0.32		99		
41000103	**Low discrimination**						**41000103**
					SPO		
Clostridium novyi biovar A		64.52	0.40	LAC 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41000110	**Low discrimination**						**41000110**
Clostridium novyi biovar A		63.76	0.53	LAC 1			
Clostridium botulinum biovar A		17.84	0.32				
Clostridium sporogenes		13.12	0.27				
41000111	**Good identification**						**41000111**
Clostridium novyi biovar A		100.0	0.53	LAC 1			
41000112	**Low discrimination**						**41000112**
					SPO		
Clostridium novyi biovar A		64.52	0.40	LAC 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41000113	**Good identification**						**41000113**
Clostridium novyi biovar A		100.0	0.40	LAC 1			
41000120	**Low discrimination**						**41000120**
					SPO		
Atopobium minutum		62.56	0.51	LAC 1	1		
Clostridium botulinum biovar A		12.43	0.32		99		
Clostridium novyi biovar A		11.10	0.43		99		
41000121	**Acceptable identification**						**41000121**
					SPO		
Clostridium novyi biovar A		94.62	0.43	LAC 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		5.38	0.17		1		
41000122	**Low discrimination**						**41000122**
					GRA	SPO	
Atopobium minutum		54.58	0.48	LAC 1	99	1	
Pseudoflavonifractor capillosus		38.87	0.32		1	1	
Clostridium sporogenes		3.99	0.19		99	99	
41000130	**Acceptable identification**						**41000130**
					SPO		
Clostridium novyi biovar A		94.62	0.43	LAC 1	99		
Atopobium minutum		5.38	0.17		1		
41000131	**Good identification**						**41000131**
Clostridium novyi biovar A		100.0	0.43	LAC 1			
41000140	**Low discrimination**						**41000140**
Clostridium botulinum biovar A		87.87	0.57	LAC 1			
Clostridium novyi biovar A		7.08	0.50				
Clostridium sporogenes		2.19	0.27				
41000141	**Low discrimination**						**41000141**
Clostridium novyi biovar A		88.86	0.50	LAC 1			
Clostridium botulinum biovar A		11.14	0.23				
41000142	**Low discrimination**						**41000142**
Clostridium novyi biovar A		39.44	0.37	LAC 1			
Clostridium botulinum biovar A		28.02	0.23				
Clostridium sporogenes		23.00	0.19				
41000143	**Good identification**						**41000143**
Clostridium novyi biovar A		100.0	0.37	LAC 1			
41000150	**Low discrimination**						**41000150**
Clostridium novyi biovar A		88.86	0.50	LAC 1			
Clostridium botulinum biovar A		11.14	0.23				
41000151	**Good identification**						**41000151**
Clostridium novyi biovar A		100.0	0.50	LAC 1			
41000152	**Good identification**						**41000152**
Clostridium novyi biovar A		100.0	0.37	LAC 1			
41000153	**Good identification**						**41000153**
Clostridium novyi biovar A		100.0	0.37	LAC 1			
41000160	**Low discrimination**						**41000160**
					SPO		
Clostridium novyi biovar A		62.29	0.40	LAC 1	99		
Clostridium botulinum biovar A		31.24	0.23		99		
Atopobium minutum		5.32	0.17		1		
41000161	**Good identification**						**41000161**
Clostridium novyi biovar A		100.0	0.40	LAC 1			
41000170	**Good identification**						**41000170**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.40	LAC 1		
41000171	**Good identification**					**41000171**
Clostridium novyi biovar A		100.0	0.40	LAC 1		
41000200	**Low discrimination**					**41000200**
					SPO	
Clostridium botulinum biovar A		50.20	0.66	RHA 1	99	
Clostridium sporogenes		36.92	0.61		99	
Atopobium minutum		7.66	0.60		1	
41000201	**Low discrimination**					**41000201**
Clostridium novyi biovar A		64.15	0.53	RHA 1		
Clostridium botulinum biovar A		17.95	0.32			
Clostridium sporogenes		13.20	0.27			
41000202	**Low discrimination**					**41000202**
					SPO	
Clostridium sporogenes		67.32	0.53	RHA 1	99	
Atopobium minutum		27.93	0.57		1	
Clostridium botulinum biovar A		2.77	0.32		99	
41000203	**Low discrimination**					**41000203**
					SPO	
Clostridium novyi biovar A		64.52	0.40	RHA 1	99	
Clostridium sporogenes		25.08	0.19		99	
Atopobium minutum		10.40	0.22		1	
41000210	**Low discrimination**					**41000210**
Clostridium novyi biovar A		63.76	0.53	RHA 1		
Clostridium botulinum biovar A		17.84	0.32			
Clostridium sporogenes		13.12	0.27			
41000211	**Good identification**					**41000211**
Clostridium novyi biovar A		100.0	0.53	RHA 1		
41000212	**Low discrimination**					**41000212**
					SPO	
Clostridium novyi biovar A		64.52	0.40	RHA 1	99	
Clostridium sporogenes		25.08	0.19		99	
Atopobium minutum		10.40	0.22		1	
41000213	**Good identification**					**41000213**
Clostridium novyi biovar A		100.0	0.40	RHA 1		
41000220	**Low discrimination**					**41000220**
					SPO	
Atopobium minutum		62.92	0.51	RHA 1	1	
Clostridium botulinum biovar A		12.50	0.32		99	
Clostridium novyi biovar A		11.17	0.43		99	
41000221	**Acceptable identification**					**41000221**
					SPO	
Clostridium novyi biovar A		94.62	0.43	RHA 1	99	
Atopobium minutum		5.38	0.17		1	
41000222	**Low discrimination**					**41000222**
					GRA SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		54.58	0.48	RHA 1	99	1	
Pseudoflavonifractor capillosus		38.87	0.32		1	1	
Clostridium sporogenes		3.99	0.19		99	99	
41000230	**Acceptable identification**						**41000230**
					SPO		
Clostridium novyi biovar A		94.62	0.43	RHA 1	99		
Atopobium minutum		5.38	0.17		1		
41000231	**Good identification**						**41000231**
Clostridium novyi biovar A		100.0	0.43	RHA 1			
41000240	**Low discrimination**						**41000240**
Clostridium botulinum biovar A		87.87	0.57	RHA 1			
Clostridium novyi biovar A		7.08	0.50				
Clostridium sporogenes		2.19	0.27				
41000241	**Low discrimination**						**41000241**
Clostridium novyi biovar A		88.86	0.50	RHA 1			
Clostridium botulinum biovar A		11.14	0.23				
41000242	**Low discrimination**						**41000242**
Clostridium novyi biovar A		39.44	0.37	RHA 1			
Clostridium botulinum biovar A		28.02	0.23				
Clostridium sporogenes		23.00	0.19				
41000243	**Good identification**						**41000243**
Clostridium novyi biovar A		100.0	0.37	RHA 1			
41000250	**Low discrimination**						**41000250**
Clostridium novyi biovar A		88.86	0.50	RHA 1			
Clostridium botulinum biovar A		11.14	0.23				
41000251	**Good identification**						**41000251**
Clostridium novyi biovar A		100.0	0.50	RHA 1			
41000252	**Good identification**						**41000252**
Clostridium novyi biovar A		100.0	0.37	RHA 1			
41000253	**Good identification**						**41000253**
Clostridium novyi biovar A		100.0	0.37	RHA 1			
41000260	**Low discrimination**						**41000260**
					SPO		
Clostridium novyi biovar A		62.29	0.40	RHA 1	99		
Clostridium botulinum biovar A		31.24	0.23		99		
Atopobium minutum		5.32	0.17		1		
41000261	**Good identification**						**41000261**
Clostridium novyi biovar A		100.0	0.40	RHA 1			
41000270	**Good identification**						**41000270**
Clostridium novyi biovar A		100.0	0.40	RHA 1			
41000271	**Good identification**						**41000271**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.40	RHA 1			
41000400	**Low discrimination**						**41000400**
					SPO		
Clostridium botulinum biovar A		50.20	0.66	ARA 1	99		
Clostridium sporogenes		36.92	0.61		99		
Atopobium minutum		7.66	0.60		1		
41000401	**Low discrimination**						**41000401**
Clostridium novyi biovar A		64.15	0.53	ARA 1			
Clostridium botulinum biovar A		17.95	0.32				
Clostridium sporogenes		13.20	0.27				
41000402	**Low discrimination**						**41000402**
					SPO		
Clostridium sporogenes		67.32	0.53	ARA 1	99		
Atopobium minutum		27.93	0.57		1		
Clostridium botulinum biovar A		2.77	0.32		99		
41000403	**Low discrimination**						**41000403**
					SPO		
Clostridium novyi biovar A		64.52	0.40	ARA 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41000410	**Low discrimination**						**41000410**
Clostridium novyi biovar A		63.76	0.53	ARA 1			
Clostridium botulinum biovar A		17.84	0.32				
Clostridium sporogenes		13.12	0.27				
41000411	**Good identification**						**41000411**
Clostridium novyi biovar A		100.0	0.53	ARA 1			
41000412	**Low discrimination**						**41000412**
					SPO		
Clostridium novyi biovar A		64.52	0.40	ARA 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41000413	**Good identification**						**41000413**
Clostridium novyi biovar A		100.0	0.40	ARA 1			
41000420	**Low discrimination**						**41000420**
					SPO		
Atopobium minutum		62.92	0.51	ARA 1	1		
Clostridium botulinum biovar A		12.50	0.32		99		
Clostridium novyi biovar A		11.17	0.43		99		
41000421	**Acceptable identification**						**41000421**
					SPO		
Clostridium novyi biovar A		94.62	0.43	ARA 1	99		
Atopobium minutum		5.38	0.17		1		
41000422	**Low discrimination**						**41000422**
					GRA	SPO	
Atopobium minutum		54.58	0.48	ARA 1	99	1	
Pseudoflavonifractor capillosus		38.87	0.32		1	1	
Clostridium sporogenes		3.99	0.19		99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
41000430	**Acceptable identification**					**41000430**
					SPO	
Clostridium novyi biovar A		94.62	0.43	ARA 1	99	
Atopobium minutum		5.38	0.17		1	
41000431	**Good identification**					**41000431**
Clostridium novyi biovar A		100.0	0.43	ARA 1		
41000440	**Low discrimination**					**41000440**
Clostridium botulinum biovar A		87.87	0.57	ARA 1		
Clostridium novyi biovar A		7.08	0.50			
Clostridium sporogenes		2.19	0.27			
41000441	**Low discrimination**					**41000441**
Clostridium novyi biovar A		88.86	0.50	ARA 1		
Clostridium botulinum biovar A		11.14	0.23			
41000442	**Low discrimination**					**41000442**
Clostridium novyi biovar A		39.44	0.37	ARA 1		
Clostridium botulinum biovar A		28.02	0.23			
Clostridium sporogenes		23.00	0.19			
41000443	**Good identification**					**41000443**
Clostridium novyi biovar A		100.0	0.37	ARA 1		
41000450	**Low discrimination**					**41000450**
Clostridium novyi biovar A		88.86	0.50	ARA 1		
Clostridium botulinum biovar A		11.14	0.23			
41000451	**Good identification**					**41000451**
Clostridium novyi biovar A		100.0	0.50	ARA 1		
41000452	**Good identification**					**41000452**
Clostridium novyi biovar A		100.0	0.37	ARA 1		
41000453	**Good identification**					**41000453**
Clostridium novyi biovar A		100.0	0.37	ARA 1		
41000460	**Low discrimination**					**41000460**
					SPO	
Clostridium novyi biovar A		62.29	0.40	ARA 1	99	
Clostridium botulinum biovar A		31.24	0.23		99	
Atopobium minutum		5.32	0.17		1	
41000461	**Good identification**					**41000461**
Clostridium novyi biovar A		100.0	0.40	ARA 1		
41000470	**Good identification**					**41000470**
Clostridium novyi biovar A		100.0	0.40	ARA 1		
41000471	**Good identification**					**41000471**
Clostridium novyi biovar A		100.0	0.40	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41001000	**Low discrimination**						**41001000**
					SPO		
Atopobium minutum		69.33	0.86		1		
Clostridium botulinum biovar A		13.77	0.66		99		
Clostridium sporogenes		10.13	0.61		99		
41001001	**Low discrimination**						**41001001**
					SPO		
Clostridium novyi biovar A		84.73	0.71	GAL 10	99		
Atopobium minutum		10.85	0.51		1		
Clostridium botulinum biovar A		2.16	0.32		99		
41001002	**Good identification**						**41001002**
					SPO		
Atopobium minutum		91.01	0.83		1		
Clostridium sporogenes		6.65	0.53		99		
Clostridium novyi biovar A		1.90	0.58		99		
41001003	**Low discrimination**						**41001003**
					SPO		
Clostridium novyi biovar A		65.83	0.58	GAL 10	99		
Atopobium minutum		31.85	0.48		1		
Clostridium sporogenes		2.33	0.19		99		
41001010	**Low discrimination**						**41001010**
					SPO		
Clostridium novyi biovar A		85.02	0.71	GAL 10	99		
Atopobium minutum		10.89	0.51		1		
Clostridium botulinum biovar A		2.16	0.32		99		
41001011	**Very good identification**						**41001011**
					SPO		
Clostridium novyi biovar A		99.87	0.71	GAL 10	99		
Atopobium minutum		0.13	0.17		1		
41001012	**Low discrimination**						**41001012**
					SPO		
Clostridium novyi biovar A		65.83	0.58	GAL 10	99		
Atopobium minutum		31.85	0.48		1		
Clostridium sporogenes		2.33	0.19		99		
41001013	**Very good identification**						**41001013**
					SPO		
Clostridium novyi biovar A		99.51	0.58	GAL 10	99		
Atopobium minutum		0.49	0.14		1		
41001020	**Good identification**						**41001020**
					SPO		
Atopobium minutum		93.04	0.77		1		
Clostridium novyi biovar A		5.50	0.61		99		
Clostridium botulinum biovar A		0.56	0.32		99		
41001021	**Low discrimination**						**41001021**
					SPO		
Clostridium novyi biovar A		85.42	0.61	GAL 10	99		
Atopobium minutum		14.58	0.43		1		
41001022	**Low discrimination**						**41001022**
					GRA	SPO	
Atopobium minutum		79.63	0.74		99	1	
Pseudoflavonifractor capillosus		18.91	0.50		1	1	
Clostridium novyi biovar A		1.25	0.48		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41001023	**Low discrimination**						**41001023**
					GRA	SPO	
Clostridium novyi biovar A		55.61	0.48	GAL 10	99	99	
Atopobium minutum		35.87	0.40		99	1	
Pseudoflavonifractor capillosus		8.52	0.15		1	1	
41001030	**Low discrimination**						**41001030**
					SPO		
Clostridium novyi biovar A		85.42	0.61	GAL 10	99		
Atopobium minutum		14.58	0.43		1		
41001031	**Very good identification**						**41001031**
					SPO		
Clostridium novyi biovar A		99.83	0.61	GAL 10	99		
Atopobium minutum		0.17	0.09		1		
41001032	**Low discrimination**						**41001032**
					GRA	SPO	
Clostridium novyi biovar A		55.61	0.48	GAL 10	99	99	
Atopobium minutum		35.87	0.40		99	1	
Pseudoflavonifractor capillosus		8.52	0.15		1	1	
41001033	**Good identification**						**41001033**
					SPO		
Clostridium novyi biovar A		99.35	0.48	GAL 10	99		
Atopobium minutum		0.65	0.06		1		
41001040	**Low discrimination**						**41001040**
Clostridium botulinum biovar A		43.14	0.57	GAL 1			
Clostridium novyi biovar A		38.23	0.68				
Clostridium glycolicum		9.91	0.45				
41001041	**Very good identification**						**41001041**
Clostridium novyi biovar A		98.43	0.68	GAL 10			
Clostridium botulinum biovar A		1.12	0.23				
Clostridium glycolicum		0.26	0.11				
41001042	**Low discrimination**						**41001042**
					SPO		
Clostridium novyi biovar A		53.82	0.55	GAL 10	99		
Atopobium minutum		39.05	0.48		1		
Clostridium botulinum biovar A		3.48	0.23		99		
41001043	**Good identification**						**41001043**
					SPO		
Clostridium novyi biovar A		99.27	0.55	GAL 10	99		
Atopobium minutum		0.73	0.14		1		
41001050	**Very good identification**						**41001050**
Clostridium novyi biovar A		98.43	0.68	GAL 10			
Clostridium botulinum biovar A		1.12	0.23				
Clostridium glycolicum		0.26	0.11				
41001051	**Very good identification**						**41001051**
Clostridium novyi biovar A		100.0	0.68	GAL 10			
41001052	**Good identification**						**41001052**
					SPO		
Clostridium novyi biovar A		99.27	0.55	GAL 10	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Atopobium minutum	0.73	0.14		1		
41001053	**Good identification**						**41001053**
	Clostridium novyi biovar A	100.0	0.55	GAL 10			
41001060	**Low discrimination**						**41001060**
					SPO		
	Clostridium novyi biovar A	76.11	0.58	GAL 10	99		
	Atopobium minutum	19.49	0.43		1		
	Clostridium botulinum biovar A	3.47	0.23		99		
41001061	**Very good identification**						**41001061**
					SPO		
	Clostridium novyi biovar A	99.74	0.58	GAL 10	99		
	Atopobium minutum	0.26	0.09		1		
41001062	**Low discrimination**						**41001062**
					GRA	SPO	
	Clostridium novyi biovar A	45.51	0.45	GAL 10	99	99	
	Atopobium minutum	44.03	0.40		99	1	
	Pseudoflavonifractor capillosus	10.46	0.15		1	1	
41001063	**Good identification**						**41001063**
					SPO		
	Clostridium novyi biovar A	99.03	0.45	GAL 10	99		
	Atopobium minutum	0.97	0.06		1		
41001070	**Very good identification**						**41001070**
					SPO		
	Clostridium novyi biovar A	99.74	0.58	GAL 10	99		
	Atopobium minutum	0.26	0.09		1		
41001071	**Very good identification**						**41001071**
	Clostridium novyi biovar A	100.0	0.58	GAL 10			
41001072	**Good identification**						**41001072**
					SPO		
	Clostridium novyi biovar A	99.03	0.45	GAL 10	99		
	Atopobium minutum	0.97	0.06		1		
41001073	**Good identification**						**41001073**
	Clostridium novyi biovar A	100.0	0.45	GAL 10			
41001100	**Low discrimination**						**41001100**
					COC	SPO	
	Atopobium minutum	52.77	0.51	LAC 1	1	1	
	Blautia hansenii	18.47	0.32		99	1	
	Clostridium botulinum biovar A	10.48	0.32		1	99	
41001101	**Low discrimination**						**41001101**
					SPO		
	Clostridium novyi biovar A	80.85	0.37	GAL 10	99		
	Atopobium minutum	10.35	0.17	LAC 1	1		
	Clostridium chauvoei	5.17	0.06		99		
41001102	**Low discrimination**						**41001102**
					COC	SPO	
	Atopobium minutum	85.78	0.48	LAC 1	1	1	
	Clostridium sporogenes	6.27	0.19		1	99	
	Blautia hansenii	5.00	0.15		99	1	
41001110	**Low discrimination**						**41001110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium novyi biovar A		82.67	0.37	GAL 10	1	99	
Atopobium minutum		10.59	0.17	LAC 1	1	1	
Blautia hansenii		3.71	-.03		99	1	
41001111	**Good identification**						**41001111**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
41001120	**Low discrimination**						**41001120**
					COC	SPO	
Atopobium minutum		83.79	0.43	LAC 1	1	1	
Blautia hansenii		9.78	0.15		99	1	
Clostridium novyi biovar A		4.96	0.26		1	99	
41001122	**Low discrimination**						**41001122**
					COC	GRA	
Atopobium minutum		78.58	0.40	LAC 1	1	99	
Pseudoflavonifractor capillosus		18.66	0.15		1	1	
Blautia hansenii		1.53	-.01		99	99	
41001200	**Low discrimination**						**41001200**
					SPO		
Atopobium minutum		69.49	0.51	RHA 1	1		
Clostridium botulinum biovar A		13.81	0.32		99		
Clostridium sporogenes		10.15	0.27		99		
41001201	**Low discrimination**						**41001201**
					SPO		
Clostridium novyi biovar A		88.65	0.37	GAL 10	99		
Atopobium minutum		11.35	0.17	RHA 1	1		
41001202	**Acceptable identification**						**41001202**
					SPO		
Atopobium minutum		91.41	0.48	RHA 1	1		
Clostridium sporogenes		6.68	0.19		99		
Clostridium novyi biovar A		1.91	0.24		99		
41001210	**Low discrimination**						**41001210**
					SPO		
Clostridium novyi biovar A		88.65	0.37	GAL 10	99		
Atopobium minutum		11.35	0.17	RHA 1	1		
41001211	**Good identification**						**41001211**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
41001220	**Acceptable identification**						**41001220**
					SPO		
Atopobium minutum		94.41	0.43	RHA 1	1		
Clostridium novyi biovar A		5.59	0.26		99		
41001222	**Low discrimination**						**41001222**
					GRA	SPO	
Atopobium minutum		79.80	0.40	RHA 1	99	1	
Pseudoflavonifractor capillosus		18.95	0.15		1	1	
Clostridium novyi biovar A		1.25	0.13		99	99	
41001400	**Low discrimination**						**41001400**
					SPO		
Atopobium minutum		69.49	0.51	ARA 1	1		
Clostridium botulinum biovar A		13.81	0.32		99		
Clostridium sporogenes		10.15	0.27		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41001401	**Low discrimination**						**41001401**
					SPO		
Clostridium novyi biovar A		88.65	0.37	GAL 10	99		
Atopobium minutum		11.35	0.17	ARA 1	1		
41001402	**Acceptable identification**						**41001402**
					SPO		
Atopobium minutum		91.41	0.48	ARA 1	1		
Clostridium sporogenes		6.68	0.19		99		
Clostridium novyi biovar A		1.91	0.24		99		
41001410	**Low discrimination**						**41001410**
					SPO		
Clostridium novyi biovar A		88.65	0.37	GAL 10	99		
Atopobium minutum		11.35	0.17	ARA 1	1		
41001411	**Good identification**						**41001411**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
41001420	**Acceptable identification**						**41001420**
					GRA	SPO	
Atopobium minutum		93.62	0.43	ARA 1	99	1	
Clostridium novyi biovar A		5.54	0.26		99	99	
Odoribacter splanchnicus		0.84	-.07		1	1	
41001422	**Low discrimination**						**41001422**
					GRA	SPO	
Atopobium minutum		79.80	0.40	ARA 1	99	1	
Pseudoflavonifractor capillosus		18.95	0.15		1	1	
Clostridium novyi biovar A		1.25	0.13		99	99	
41002000	**Low discrimination**						**41002000**
					SPO		
Eubacterium limosum		35.98	0.63	FRU 99	1		
Clostridium botulinum biovar A		29.54	0.66		99		
Clostridium sporogenes		21.72	0.61		99		
41002001	**Low discrimination**						**41002001**
					SPO		
Clostridium novyi biovar A		51.29	0.53	MAN 1	99		
Eubacterium limosum		17.48	0.29		1		
Clostridium botulinum biovar A		14.35	0.32		99		
41002002	**Low discrimination**						**41002002**
					SPO		
Clostridium sporogenes		64.36	0.53	MAN 1	99		
Atopobium minutum		26.70	0.57		1		
Eubacterium limosum		3.23	0.29		1		
41002003	**Low discrimination**						**41002003**
					SPO		
Clostridium novyi biovar A		63.72	0.40	MAN 1	99		
Clostridium sporogenes		24.77	0.19		99		
Atopobium minutum		10.27	0.22		1		
41002010	**Good identification**						**41002010**
					SPO		
Clostridium difficile		91.57	0.61	FRU 99	99		
Clostridium novyi biovar A		4.45	0.53		99		
Eubacterium limosum		1.52	0.29		1		
41002011	**Low discrimination**						**41002011**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium novyi biovar A		82.56	0.53	MAN 1	99		
Clostridium difficile		17.16	0.27		99		
Eubacterium limosum		0.28	-.06		1		
41002013	**Good identification**						**41002013**
Clostridium novyi biovar A		100.0	0.40	MAN 1			
41002020	**Low discrimination**						**41002020**
					SPO		
Atopobium minutum		52.38	0.51	MAN 1	1		
Eubacterium limosum		12.68	0.29		1		
Clostridium botulinum biovar A		10.41	0.32		99		
41002021	**Acceptable identification**						**41002021**
					SPO		
Clostridium novyi biovar A		93.40	0.43	MAN 1	99		
Atopobium minutum		5.32	0.17		1		
Eubacterium limosum		1.29	-.06		1		
41002022	**Low discrimination**						**41002022**
					GRA	SPO	
Atopobium minutum		54.25	0.48	MAN 1	99	1	
Pseudoflavonifractor capillosus		38.64	0.32		1	1	
Clostridium sporogenes		3.96	0.19		99	99	
41002030	**Low discrimination**						**41002030**
					SPO		
Clostridium novyi biovar A		52.58	0.43	MAN 1	99		
Clostridium difficile		43.71	0.27		99		
Atopobium minutum		2.99	0.17		1		
41002031	**Good identification**						**41002031**
Clostridium novyi biovar A		100.0	0.43	MAN 1			
41002040	**Low discrimination**						**41002040**
					SPO		
Clostridium botulinum biovar A		80.25	0.57	MAN 1	99		
Clostridium novyi biovar A		6.47	0.50		99		
Eubacterium limosum		3.31	0.29		1		
41002041	**Low discrimination**						**41002041**
					SPO		
Clostridium novyi biovar A		88.45	0.50	MAN 1	99		
Clostridium botulinum biovar A		11.09	0.23		99		
Eubacterium limosum		0.46	-.06		1		
41002042	**Low discrimination**						**41002042**
Clostridium novyi biovar A		34.74	0.37	MAN 1			
Clostridium botulinum biovar A		24.68	0.23				
Clostridium sporogenes		20.26	0.19				
41002043	**Good identification**						**41002043**
Clostridium novyi biovar A		100.0	0.37	MAN 1			
41002050	**Low discrimination**						**41002050**
Clostridium difficile		58.96	0.39	FRU 99			
Clostridium novyi biovar A		36.30	0.50	SOR 5			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium botulinum biovar A		4.55	0.23			
41002051	**Good identification**					**41002051**
Clostridium novyi biovar A		98.39	0.50	MAN 1		
Clostridium difficile		1.61	0.05			
41002052	**Acceptable identification**					**41002052**
Clostridium novyi biovar A		91.49	0.37	MAN 1		
Clostridium difficile		8.51	0.05			
41002053	**Good identification**					**41002053**
Clostridium novyi biovar A		100.0	0.37	MAN 1		
41002060	**Low discrimination**					**41002060**
					SPO	
Clostridium novyi biovar A		58.12	0.40	MAN 1	99	
Clostridium botulinum biovar A		29.15	0.23		99	
Pseudoramibacter alactolyticus		5.50	0.04		1	
41002061	**Good identification**					**41002061**
Clostridium novyi biovar A		100.0	0.40	MAN 1		
41002070	**Acceptable identification**					**41002070**
Clostridium novyi biovar A		93.84	0.40	MAN 1		
Clostridium difficile		6.16	0.05			
41002071	**Good identification**					**41002071**
Clostridium novyi biovar A		100.0	0.40	MAN 1		
41003000	**Low discrimination**					**41003000**
					SPO	
Atopobium minutum		58.26	0.51	MAN 1	1	
Eubacterium limosum		14.10	0.29		1	
Clostridium botulinum biovar A		11.58	0.32		99	
41003001	**Low discrimination**					**41003001**
					SPO	
Clostridium novyi biovar A		86.28	0.37	GAL 10	99	
Atopobium minutum		11.05	0.17	MAN 1	1	
Eubacterium limosum		2.67	-.06		1	
41003002	**Acceptable identification**					**41003002**
					SPO	
Atopobium minutum		91.11	0.48	MAN 1	1	
Clostridium sporogenes		6.66	0.19		99	
Clostridium novyi biovar A		1.90	0.24		99	
41003011	**Good identification**					**41003011**
Clostridium novyi biovar A		100.0	0.37	GAL 10		
41003020	**Acceptable identification**					**41003020**
					SPO	
Atopobium minutum		93.77	0.43	MAN 1	1	
Clostridium novyi biovar A		5.55	0.26		99	
Eubacterium limosum		0.69	-.06		1	
41003022	**Low discrimination**					**41003022**
					GRA SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		79.80	0.40	MAN 1	99	1	
Pseudoflavonifractor capillosus		18.95	0.15		1	1	
Clostridium novyi biovar A		1.25	0.13		99	99	
41004000	**Low discrimination**						**41004000**
Clostridium botulinum biovar A		38.30	0.66	XYL 1			
Clostridium sporogenes		28.17	0.61				
Clostridium glycolicum		23.66	0.61				
41004001	**Low discrimination**						**41004001**
Clostridium novyi biovar A		57.75	0.53	XYL 1			
Clostridium botulinum biovar A		16.16	0.32				
Clostridium sporogenes		11.88	0.27				
41004002	**Low discrimination**						**41004002**
					SPO		
Clostridium sporogenes		66.18	0.53	XYL 1	99		
Atopobium minutum		27.45	0.57		1		
Clostridium botulinum biovar A		2.73	0.32		99		
41004003	**Low discrimination**						**41004003**
					SPO		
Clostridium novyi biovar A		64.52	0.40	XYL 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41004010	**Low discrimination**						**41004010**
Clostridium novyi biovar A		54.39	0.53	XYL 1			
Clostridium botulinum biovar A		15.22	0.32				
Clostridium sporogenes		11.19	0.27				
41004011	**Good identification**						**41004011**
Clostridium novyi biovar A		100.0	0.53	XYL 1			
41004012	**Low discrimination**						**41004012**
					SPO		
Clostridium novyi biovar A		64.52	0.40	XYL 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41004013	**Good identification**						**41004013**
Clostridium novyi biovar A		100.0	0.40	XYL 1			
41004020	**Low discrimination**						**41004020**
					SPO		
Atopobium minutum		58.41	0.51	XYL 1	1		
Clostridium botulinum biovar A		11.60	0.32		99		
Clostridium novyi biovar A		10.37	0.43		99		
41004021	**Acceptable identification**						**41004021**
					SPO		
Clostridium novyi biovar A		94.62	0.43	XYL 1	99		
Atopobium minutum		5.38	0.17		1		
41004022	**Low discrimination**						**41004022**
					GRA	SPO	
Atopobium minutum		54.58	0.48	XYL 1	99	1	
Pseudoflavonifractor capillosus		38.87	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sporogenes		3.99	0.19		99 99	
41004030	**Acceptable identification**					**41004030**
					SPO	
Clostridium novyi biovar A		94.62	0.43	XYL 1	99	
Atopobium minutum		5.38	0.17		1	
41004031	**Good identification**					**41004031**
Clostridium novyi biovar A		100.0	0.43	XYL 1		
41004040	**Very good identification**					**41004040**
Clostridium glycolicum		97.24	0.83	FRU 90		
Clostridium botulinum biovar A		2.47	0.57			
Clostridium novyi biovar A		0.20	0.50			
41004041	**Low discrimination**					**41004041**
Clostridium glycolicum		81.41	0.49	FRU 90		
Clostridium novyi biovar A		16.52	0.50	URE 1		
Clostridium botulinum biovar A		2.07	0.23			
41004042	**Acceptable identification**					**41004042**
Clostridium glycolicum		91.68	0.49	FRU 90		
Clostridium novyi biovar A		3.28	0.37	BGL 1		
Clostridium botulinum biovar A		2.33	0.23			
41004043	**Low discrimination**					**41004043**
Clostridium novyi biovar A		78.00	0.37	XYL 1		
Clostridium glycolicum		22.00	0.15			
41004050	**Low discrimination**					**41004050**
Clostridium glycolicum		81.28	0.49	FRU 90		
Clostridium novyi biovar A		16.50	0.50	MLZ 1		
Clostridium botulinum biovar A		2.07	0.23			
41004051	**Good identification**					**41004051**
Clostridium novyi biovar A		95.26	0.50	XYL 1		
Clostridium glycolicum		4.74	0.15			
41004052	**Low discrimination**					**41004052**
Clostridium novyi biovar A		78.00	0.37	XYL 1		
Clostridium glycolicum		22.00	0.15			
41004053	**Good identification**					**41004053**
Clostridium novyi biovar A		100.0	0.37	XYL 1		
41004060	**Acceptable identification**					**41004060**
Clostridium glycolicum		92.47	0.49	FRU 90		
Clostridium novyi biovar A		4.69	0.40	NAG 1		
Clostridium botulinum biovar A		2.35	0.23			
41004061	**Low discrimination**					**41004061**
Clostridium novyi biovar A		83.40	0.40	XYL 1		
Clostridium glycolicum		16.60	0.15			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
41004070	**Low discrimination**					**41004070**
Clostridium novyi biovar A		83.40	0.40	XYL 1		
Clostridium glycolicum		16.60	0.15			
41004071	**Good identification**					**41004071**
Clostridium novyi biovar A		100.0	0.40	XYL 1		
41004140	**Good identification**					**41004140**
Clostridium glycolicum		97.32	0.49	FRU 90		
Clostridium botulinum biovar A		2.48	0.23	LAC 1		
Clostridium novyi biovar A		0.20	0.16			
41004240	**Good identification**					**41004240**
Clostridium glycolicum		97.32	0.49	FRU 90		
Clostridium botulinum biovar A		2.48	0.23	RHA 1		
Clostridium novyi biovar A		0.20	0.16			
41004440	**Good identification**					**41004440**
Clostridium glycolicum		97.32	0.49	FRU 90		
Clostridium botulinum biovar A		2.48	0.23	ARA 1		
Clostridium novyi biovar A		0.20	0.16			
41005000	**Low discrimination**					**41005000**
					SPO	
Clostridium glycolicum		48.18	0.45	FRU 90	99	
Atopobium minutum		35.69	0.51	GAL 10	1	
Clostridium botulinum biovar A		7.09	0.32	SOR 95	99	
41005001	**Low discrimination**					**41005001**
					SPO	
Clostridium novyi biovar A		76.87	0.37	GAL 10	99	
Clostridium glycolicum		13.29	0.11	XYL 1	99	
Atopobium minutum		9.84	0.17		1	
41005002	**Low discrimination**					**41005002**
					SPO	
Atopobium minutum		89.74	0.48	XYL 1	1	
Clostridium sporogenes		6.56	0.19		99	
Clostridium novyi biovar A		1.87	0.24		99	
41005010	**Low discrimination**					**41005010**
					SPO	
Clostridium novyi biovar A		76.87	0.37	GAL 10	99	
Clostridium glycolicum		13.29	0.11	XYL 1	99	
Atopobium minutum		9.84	0.17		1	
41005011	**Good identification**					**41005011**
Clostridium novyi biovar A		100.0	0.37	GAL 10		
41005020	**Acceptable identification**					**41005020**
					SPO	
Atopobium minutum		90.90	0.43	XYL 1	1	
Clostridium novyi biovar A		5.38	0.26		99	
Clostridium glycolicum		3.72	0.11		99	
41005022	**Low discrimination**					**41005022**
					GRA SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		79.80	0.40	XYL 1	99	1	
Pseudoflavonifractor capillosus		18.95	0.15		1	1	
Clostridium novyi biovar A		1.25	0.13		99	99	
41005040	**Very good identification**						**41005040**
Clostridium glycolicum		99.53	0.67	FRU 90			
Clostridium botulinum biovar A		0.23	0.23	GAL 10			
Clostridium novyi biovar A		0.20	0.34				
41006010	**Good identification**						**41006010**
					SPO		
Clostridium difficile		98.77	0.39	FRU 99	99		
Clostridium novyi biovar A		0.92	0.19	XYL 5	99		
Eubacterium limosum		0.31	-.06		1		
41006040	**Good identification**						**41006040**
Clostridium glycolicum		96.89	0.49	FRU 90			
Clostridium botulinum biovar A		2.47	0.23	MAN 1			
Clostridium difficile		0.34	0.05				
41010000	**Low discrimination**						**41010000**
					SPO		
Eubacterium limosum		33.59	0.84	MAN 86	1		
Atopobium minutum		23.85	0.93		1		
Clostridium sporogenes		21.99	0.83		99		
41010001	**Low discrimination**						**41010001**
					SPO		
Clostridium novyi biovar A		39.90	0.71	FRU 10	99		
Eubacterium limosum		19.93	0.49		1		
Atopobium minutum		14.15	0.59		1		
41010002	**Low discrimination**						**41010002**
					SPO		
Atopobium minutum		65.26	0.90		1		
Clostridium sporogenes		30.07	0.74		99		
Pseudoramibacter alactolyticus		1.85	0.50		1		
41010003	**Low discrimination**						**41010003**
					SPO		
Atopobium minutum		44.03	0.56	URE 1	1		
Clostridium novyi biovar A		32.87	0.58		99		
Clostridium sporogenes		20.29	0.40		99		
41010010	**Low discrimination**						**41010010**
					SPO		
Clostridium difficile		89.42	0.79	MAN 90	99		
Clostridium novyi biovar A		4.35	0.71		99		
Eubacterium limosum		2.17	0.49		1		
41010011	**Low discrimination**						**41010011**
					SPO		
Clostridium novyi biovar A		81.82	0.71	FRU 10	99		
Clostridium difficile		17.00	0.45		99		
Eubacterium limosum		0.41	0.15		1		
41010012	**Low discrimination**						**41010012**
					SPO		
Atopobium minutum		31.77	0.56	MLZ 1	1		
Clostridium difficile		27.93	0.45		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		23.72	0.58		99		
41010013	**Very good identification**						**41010013**
					SPO		
Clostridium novyi biovar A		96.93	0.58	FRU 10	99		
Atopobium minutum		1.31	0.22		1		
Clostridium difficile		1.15	0.10		99		
41010020	**Low discrimination**						**41010020**
Atopobium minutum		82.46	0.85				
Pseudoramibacter alactolyticus		7.43	0.53				
Eubacterium limosum		3.52	0.49				
41010021	**Low discrimination**						**41010021**
					SPO		
Clostridium novyi biovar A		64.00	0.61	FRU 10	99		
Atopobium minutum		30.26	0.51		1		
Pseudoramibacter alactolyticus		2.73	0.19		1		
41010022	**Low discrimination**						**41010022**
					GRA		
Atopobium minutum		89.16	0.82		99		
Pseudoflavonifractor capillosus		7.65	0.50		1		
Pseudoramibacter alactolyticus		1.34	0.37		99		
41010023	**Low discrimination**						**41010023**
					GRA	SPO	
Atopobium minutum		59.72	0.48	URE 1	99	1	
Clostridium novyi biovar A		33.43	0.48		99	99	
Pseudoflavonifractor capillosus		5.12	0.15		1	1	
41010030	**Low discrimination**						**41010030**
					SPO		
Clostridium novyi biovar A		41.83	0.61	FRU 10	99		
Clostridium difficile		34.77	0.45		99		
Atopobium minutum		19.78	0.51		1		
41010031	**Very good identification**						**41010031**
					SPO		
Clostridium novyi biovar A		98.70	0.61	FRU 10	99		
Clostridium difficile		0.83	0.10		99		
Atopobium minutum		0.47	0.17		1		
41010032	**Low discrimination**						**41010032**
					GRA	SPO	
Atopobium minutum		58.78	0.48	MLZ 1	99	1	
Clostridium novyi biovar A		32.91	0.48		99	99	
Pseudoflavonifractor capillosus		5.04	0.15		1	1	
41010033	**Good identification**						**41010033**
					SPO		
Clostridium novyi biovar A		98.23	0.48	FRU 10	99		
Atopobium minutum		1.77	0.14		1		
41010040	**Low discrimination**						**41010040**
					SPO		
Clostridium glycolicum		61.49	0.78	XYL 95	99		
Clostridium botulinum biovar A		20.87	0.71		99		
Pseudoramibacter alactolyticus		6.56	0.56		1		
41010041	**Low discrimination**						**41010041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		74.84	0.68	FRU 10		
Clostridium glycolicum		15.88	0.43			
Clostridium botulinum biovar A		5.39	0.36			
41010042	**Low discrimination**					**41010042**
					SPO	
Atopobium minutum		27.09	0.56	SOR 1	1	
Pseudoramibacter alactolyticus		19.01	0.39		1	
Clostridium glycolicum		16.21	0.43		99	
41010043	**Acceptable identification**					**41010043**
					SPO	
Clostridium novyi biovar A		93.97	0.55	FRU 10	99	
Atopobium minutum		1.91	0.22		1	
Pseudoramibacter alactolyticus		1.34	0.05		1	
41010050	**Low discrimination**					**41010050**
Clostridium difficile		54.96	0.57	MAN 90		
Clostridium novyi biovar A		33.84	0.68	SOR 5		
Clostridium glycolicum		7.18	0.43			
41010051	**Very good identification**					**41010051**
Clostridium novyi biovar A		98.11	0.68	FRU 10		
Clostridium difficile		1.61	0.23			
Clostridium glycolicum		0.21	0.09			
41010052	**Low discrimination**					**41010052**
					SPO	
Clostridium novyi biovar A		86.42	0.55	FRU 10	99	
Clostridium difficile		8.03	0.23		99	
Atopobium minutum		1.75	0.22		1	
41010053	**Good identification**					**41010053**
Clostridium novyi biovar A		100.0	0.55	FRU 10		
41010060	**Low discrimination**					**41010060**
					SPO	
Pseudoramibacter alactolyticus		34.79	0.43	ESL 1	1	
Clostridium novyi biovar A		22.01	0.58		99	
Clostridium glycolicum		18.68	0.43		99	
41010061	**Very good identification**					**41010061**
					SPO	
Clostridium novyi biovar A		96.65	0.58	FRU 10	99	
Pseudoramibacter alactolyticus		1.54	0.09		1	
Clostridium glycolicum		0.83	0.09		99	
41010062	**Low discrimination**					**41010062**
					SPO	
Atopobium minutum		53.34	0.48	SOR 1	1	
Clostridium novyi biovar A		19.91	0.45		99	
Pseudoramibacter alactolyticus		19.82	0.26		1	
41010063	**Good identification**					**41010063**
					SPO	
Clostridium novyi biovar A		96.42	0.45	FRU 10	99	
Atopobium minutum		2.61	0.14		1	
Pseudoramibacter alactolyticus		0.97	-.08		1	
41010070	**Good identification**					**41010070**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium novyi biovar A		90.89	0.58	FRU 10	99		
Clostridium difficile		5.97	0.23		99		
Pseudoramibacter alactolyticus		1.45	0.09		1		
41010071	**Very good identification**						**41010071**
Clostridium novyi biovar A		100.0	0.58	FRU 10			
41010072	**Good identification**						**41010072**
					SPO		
Clostridium novyi biovar A		96.42	0.45	FRU 10	99		
Atopobium minutum		2.61	0.14		1		
Pseudoramibacter alactolyticus		0.97	-.08		1		
41010073	**Good identification**						**41010073**
Clostridium novyi biovar A		100.0	0.45	FRU 10			
41010100	**Low discrimination**						**41010100**
					GRA		
Fusobacterium mortiferum		31.14	0.69		1		
Eubacterium limosum		22.66	0.49		99		
Atopobium minutum		16.09	0.59		99		
41010101	**Low discrimination**						**41010101**
					GRA	SPO	
Clostridium novyi biovar A		31.31	0.37	FRU 10	99	99	
Fusobacterium mortiferum		21.49	0.35	LAC 1	1	1	
Eubacterium limosum		15.64	0.15		99	1	
41010102	**Low discrimination**						**41010102**
					GRA	SPO	
Atopobium minutum		53.85	0.56	LAC 1	99	1	
Clostridium sporogenes		24.82	0.40		99	99	
Fusobacterium mortiferum		17.37	0.53		1	1	
41010110	**Low discrimination**						**41010110**
					GRA	SPO	
Clostridium difficile		86.82	0.45	MAN 90	99	99	
Clostridium novyi biovar A		4.22	0.37	LAC 1	99	99	
Fusobacterium mortiferum		2.90	0.35		1	1	
41010111	**Low discrimination**						**41010111**
Clostridium novyi biovar A		82.79	0.37	FRU 10			
Clostridium difficile		17.21	0.10	LAC 1			
41010120	**Low discrimination**						**41010120**
					GRA		
Atopobium minutum		76.50	0.51	LAC 1	99		
Pseudoramibacter alactolyticus		6.89	0.19		99		
Fusobacterium mortiferum		4.49	0.35		1		
41010122	**Low discrimination**						**41010122**
					GRA		
Atopobium minutum		88.48	0.48	LAC 1	99		
Pseudoflavonifractor capillosus		7.59	0.15		1		
Pseudoramibacter alactolyticus		1.33	0.03		99		
41010140	**Low discrimination**						**41010140**
					SPO		
Clostridium glycolicum		59.69	0.43	XYL 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium botulinum biovar A		20.26	0.36	LAC 1	99	
Pseudoramibacter alactolyticus		6.37	0.21		1	
41010200	**Low discrimination**					**41010200**
					SPO	
Eubacterium limosum		33.60	0.49	MAN 86	1	
Atopobium minutum		23.86	0.59	RHA 1	1	
Clostridium sporogenes		21.99	0.48		99	
41010201	**Low discrimination**					**41010201**
					SPO	
Clostridium novyi biovar A		40.43	0.37	FRU 10	99	
Eubacterium limosum		20.19	0.15	RHA 1	1	
Atopobium minutum		14.34	0.25		1	
41010202	**Low discrimination**					**41010202**
					SPO	
Atopobium minutum		65.31	0.56	RHA 1	1	
Clostridium sporogenes		30.10	0.40		99	
Pseudoramibacter alactolyticus		1.85	0.15		1	
41010210	**Low discrimination**					**41010210**
					SPO	
Clostridium difficile		89.55	0.45	MAN 90	99	
Clostridium novyi biovar A		4.35	0.37	RHA 1	99	
Eubacterium limosum		2.17	0.15		1	
41010211	**Low discrimination**					**41010211**
Clostridium novyi biovar A		82.79	0.37	FRU 10		
Clostridium difficile		17.21	0.10	RHA 1		
41010220	**Low discrimination**					**41010220**
Atopobium minutum		82.73	0.51	RHA 1		
Pseudoramibacter alactolyticus		7.45	0.19			
Eubacterium limosum		3.53	0.15			
41010222	**Low discrimination**					**41010222**
					GRA	
Atopobium minutum		89.26	0.48	RHA 1	99	
Pseudoflavonifractor capillosus		7.65	0.15		1	
Pseudoramibacter alactolyticus		1.34	0.03		99	
41010240	**Low discrimination**					**41010240**
					SPO	
Clostridium glycolicum		61.59	0.43	XYL 95	99	
Clostridium botulinum biovar A		20.91	0.36	RHA 1	99	
Pseudoramibacter alactolyticus		6.57	0.21		1	
41010300	**Low discrimination**					**41010300**
					GRA	
Fusobacterium mortiferum		31.70	0.35	RHA 1	1	
Eubacterium limosum		23.07	0.15		99	
Atopobium minutum		16.38	0.25		99	
41010400	**Low discrimination**					**41010400**
					SPO	
Eubacterium limosum		33.54	0.49	MAN 86	1	
Atopobium minutum		23.82	0.59	ARA 1	1	
Clostridium sporogenes		21.95	0.48		99	
41010401	**Low discrimination**					**41010401**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium novyi biovar A		40.43	0.37	FRU 10	99		
Eubacterium limosum		20.19	0.15	ARA 1	1		
Atopobium minutum		14.34	0.25		1		
41010402	**Low discrimination**						**41010402**
					SPO		
Atopobium minutum		65.31	0.56	ARA 1	1		
Clostridium sporogenes		30.10	0.40		99		
Pseudoramibacter alactolyticus		1.85	0.15		1		
41010410	**Low discrimination**						**41010410**
					SPO		
Clostridium difficile		89.55	0.45	MAN 90	99		
Clostridium novyi biovar A		4.35	0.37	ARA 1	99		
Eubacterium limosum		2.17	0.15		1		
41010411	**Low discrimination**						**41010411**
Clostridium novyi biovar A		82.79	0.37	FRU 10			
Clostridium difficile		17.21	0.10	ARA 1			
41010420	**Low discrimination**						**41010420**
Atopobium minutum		82.56	0.51	ARA 1			
Pseudoramibacter alactolyticus		7.44	0.19				
Eubacterium limosum		3.52	0.15				
41010422	**Low discrimination**						**41010422**
					GRA		
Atopobium minutum		89.26	0.48	ARA 1	99		
Pseudoflavonifractor capillosus		7.65	0.15		1		
Pseudoramibacter alactolyticus		1.34	0.03		99		
41010440	**Low discrimination**						**41010440**
					SPO		
Clostridium glycolicum		61.59	0.43	XYL 95	99		
Clostridium botulinum biovar A		20.91	0.36	ARA 1	99		
Pseudoramibacter alactolyticus		6.57	0.21		1		
41011000	**Low discrimination**						**41011000**
					GRA		
Atopobium minutum		86.87	0.85		99		
Eubacterium limosum		3.71	0.49		99		
Fusobacterium mortiferum		3.28	0.66		1		
41011001	**Low discrimination**						**41011001**
					SPO		
Atopobium minutum		47.15	0.51	URE 1	1		
Clostridium novyi biovar A		44.32	0.54		99		
Clostridium sordellii		2.03	0.15		99		
41011002	**Very good identification**						**41011002**
					GRA	SPO	
Atopobium minutum		97.49	0.82		99	1	
Clostridium sporogenes		1.36	0.40		99	99	
Fusobacterium mortiferum		0.61	0.50		1	1	
41011003	**Low discrimination**						**41011003**
					SPO		
Atopobium minutum		78.80	0.48	URE 1	1		
Clostridium novyi biovar A		19.61	0.42		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		1.10	0.06		99		
41011010	**Low discrimination**						**41011010**
					SPO		
Clostridium difficile		45.83	0.45	GAL 1	99		
Atopobium minutum		26.07	0.51	MAN 90	1		
Clostridium novyi biovar A		24.50	0.54		99		
41011011	**Good identification**						**41011011**
					SPO		
Clostridium novyi biovar A		97.12	0.54	FRU 10	99		
Clostridium difficile		1.84	0.10	GAL 10	99		
Atopobium minutum		1.04	0.17		1		
41011012	**Low discrimination**						**41011012**
					SPO		
Atopobium minutum		77.18	0.48	MLZ 1	1		
Clostridium novyi biovar A		19.20	0.42		99		
Clostridium difficile		2.06	0.10		99		
41011013	**Good identification**						**41011013**
					SPO		
Clostridium novyi biovar A		96.10	0.42	FRU 10	99		
Atopobium minutum		3.90	0.14	GAL 10	1		
41011020	**Very good identification**						**41011020**
					SPO	CAT	
Atopobium minutum		98.21	0.77		1	1	
Clostridium novyi biovar A		0.70	0.44		99	1	
Propionibacterium acnes		0.28	0.26		1	95	
41011021	**Low discrimination**						**41011021**
					SPO		
Atopobium minutum		58.65	0.43	URE 1	1		
Clostridium novyi biovar A		41.35	0.44		99		
41011022	**Very good identification**						**41011022**
					GRA	SPO	
Atopobium minutum		96.86	0.74		99	1	
Pseudoflavonifractor capillosus		2.77	0.33		1	1	
Clostridium novyi biovar A		0.18	0.31		99	99	
41011023	**Low discrimination**						**41011023**
					GRA	SPO	
Atopobium minutum		82.29	0.40	URE 1	99	1	
Clostridium novyi biovar A		15.36	0.31		99	99	
Pseudoflavonifractor capillosus		2.35	-.01		1	1	
41011030	**Low discrimination**						**41011030**
					SPO		
Atopobium minutum		56.87	0.43	MLZ 1	1		
Clostridium novyi biovar A		40.10	0.44		99		
Clostridium difficile		3.03	0.10		99		
41011031	**Good identification**						**41011031**
					SPO		
Clostridium novyi biovar A		98.59	0.44	FRU 10	99		
Atopobium minutum		1.41	0.09	GAL 10	1		
41011032	**Low discrimination**						**41011032**
					GRA	SPO	
Atopobium minutum		82.29	0.40	MLZ 1	99	1	
Clostridium novyi biovar A		15.36	0.31		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Pseudoflavonifractor capillosus		2.35	-.01		1 1	
41011040	**Low discrimination**					**41011040**
					SPO	
Clostridium glycolicum		85.00	0.61	GAL 10	99	
Atopobium minutum		6.46	0.51	XYL 95	1	
Clostridium novyi biovar A		4.05	0.51		99	
41011041	**Low discrimination**					**41011041**
					SPO	
Clostridium novyi biovar A		80.98	0.51	FRU 10	99	
Clostridium glycolicum		17.18	0.27	GAL 10	99	
Atopobium minutum		1.31	0.17		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
41011042	**Low discrimination**						**41011042**
					SPO		
Atopobium minutum		70.47	0.48	SOR 1	1		
Clostridium glycolicum		14.06	0.27		99		
Clostridium novyi biovar A		11.69	0.39		99		
41011043	**Acceptable identification**						**41011043**
					SPO		
Clostridium novyi biovar A		94.26	0.39	FRU 10	99		
Atopobium minutum		5.74	0.14	GAL 10	1		
41011050	**Low discrimination**						**41011050**
Clostridium novyi biovar A		72.33	0.51	FRU 10			
Clostridium glycolicum		15.35	0.27	GAL 10			
Clostridium difficile		10.68	0.23				
41011051	**Good identification**						**41011051**
Clostridium novyi biovar A		100.0	0.51	FRU 10			
41011052	**Acceptable identification**						**41011052**
					SPO		
Clostridium novyi biovar A		94.26	0.39	FRU 10	99		
Atopobium minutum		5.74	0.14	GAL 10	1		
41011053	**Good identification**						**41011053**
Clostridium novyi biovar A		100.0	0.39	FRU 10			
41011060	**Low discrimination**						**41011060**
					SPO		
Atopobium minutum		49.71	0.43	SOR 1	1		
Clostridium novyi biovar A		23.36	0.41		99		
Clostridium glycolicum		19.83	0.27		99		
41011061	**Good identification**						**41011061**
					SPO		
Clostridium novyi biovar A		97.90	0.41	FRU 10	99		
Atopobium minutum		2.10	0.09	GAL 10	1		
41011062	**Low discrimination**						**41011062**
					GRA	SPO	
Atopobium minutum		85.89	0.40	SOR 1	99	1	
Clostridium novyi biovar A		10.69	0.28		99	99	
Pseudoflavonifractor capillosus		2.46	-.01		1	1	
41011070	**Good identification**						**41011070**
					SPO		
Clostridium novyi biovar A		97.90	0.41	FRU 10	99		
Atopobium minutum		2.10	0.09	GAL 10	1		
41011071	**Good identification**						**41011071**
Clostridium novyi biovar A		100.0	0.41	FRU 10			
41011100	**Low discrimination**						**41011100**
					GRA		
Fusobacterium mortiferum		84.05	0.72		1		
Atopobium minutum		9.65	0.51		99		
Actinomyces meyeri		4.22	0.41		99		
41011101	**Low discrimination**						**41011101**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		70.53	0.38	URE 1	1	1	
Clostridium chauvoei		10.22	0.12		99	99	
Atopobium minutum		8.10	0.17		99	1	
41011102	**Low discrimination**						**41011102**
					GRA		
Fusobacterium mortiferum		55.37	0.56	BGL 10	1		
Atopobium minutum		38.14	0.48		99		
Lactobacillus fermentum		5.18	0.40		99		
41011110	**Low discrimination**						**41011110**
					GRA	SPO	
Fusobacterium mortiferum		63.21	0.38	MLZ 1	1	1	
Clostridium difficile		12.76	0.10		99	99	
Atopobium minutum		7.26	0.17		99	1	
41011120	**Low discrimination**						**41011120**
					GRA	SPO	
Atopobium minutum		56.81	0.43	LAC 1	99	1	
Fusobacterium mortiferum		15.00	0.38		1	1	
Clostridium septicum		10.91	0.18		99	99	
41011122	**Low discrimination**						**41011122**
					GRA		
Atopobium minutum		82.75	0.40	LAC 1	99		
Lactobacillus fermentum		11.25	0.32		99		
Fusobacterium mortiferum		3.64	0.22		1		
41011200	**Low discrimination**						**41011200**
					GRA		
Atopobium minutum		87.16	0.51	RHA 1	99		
Eubacterium limosum		3.72	0.15		99		
Fusobacterium mortiferum		3.29	0.32		1		
41011202	**Good identification**						**41011202**
					GRA	SPO	
Atopobium minutum		98.02	0.48	RHA 1	99	1	
Clostridium sporogenes		1.37	0.06		99	99	
Fusobacterium mortiferum		0.62	0.16		1	1	
41011220	**Good identification**						**41011220**
					SPO		
Atopobium minutum		99.29	0.43	RHA 1	1		
Clostridium novyi biovar A		0.71	0.10		99		
41011222	**Good identification**						**41011222**
					GRA		
Atopobium minutum		97.22	0.40	RHA 1	99		
Pseudoflavonifractor capillosus		2.78	-.01		1		
41011300	**Low discrimination**						**41011300**
					GRA		
Fusobacterium mortiferum		85.84	0.38	RHA 1	1		
Atopobium minutum		9.86	0.17		99		
Actinomyces meyeri		4.31	0.07		99		
41011400	**Low discrimination**						**41011400**
Atopobium minutum		81.29	0.51	ARA 1			
Lactobacillus fermentum		4.51	0.36				
Eubacterium limosum		3.47	0.15				
41011402	**Acceptable identification**						**41011402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Atopobium minutum		90.62	0.48	ARA 1	1	
Lactobacillus fermentum		7.55	0.36		1	
Clostridium sporogenes		1.27	0.06		99	
41011420	**Acceptable identification**					**41011420**
					GRA	
Atopobium minutum		91.59	0.43	ARA 1	99	
Lactobacillus fermentum		5.09	0.28		99	
Odoribacter splanchnicus		1.65	-.03		1	
41011422	**Low discrimination**					**41011422**
					GRA	
Atopobium minutum		89.94	0.40	ARA 1	99	
Lactobacillus fermentum		7.49	0.28		99	
Pseudoflavonifractor capillosus		2.57	-.01		1	
41011502	**Low discrimination**					**41011502**
					GRA	
Lactobacillus fermentum		82.00	0.40	ESL 1	99	
Fusobacterium mortiferum		8.85	0.22	SUC 87	1	
Atopobium minutum		6.09	0.14		99	
41012000	**Low discrimination**					**41012000**
					SPO	
Eubacterium limosum		87.69	0.97		1	
Clostridium difficile		10.95	0.83		99	
Pseudoramibacter alactolyticus		0.93	0.61		1	
41012001	**Low discrimination**					**41012001**
					SPO	
Eubacterium limosum		87.44	0.63	URE 1	1	
Clostridium difficile		10.92	0.49		99	
Pseudoramibacter alactolyticus		0.93	0.27		1	
41012002	**Low discrimination**					**41012002**
					SPO	
Eubacterium limosum		72.57	0.63	BGL 1	1	
Clostridium difficile		9.06	0.49		99	
Pseudoramibacter alactolyticus		8.45	0.45		1	
41012010	**Very good identification**					**41012010**
					SPO	
Clostridium difficile		98.34	0.95		99	
Eubacterium limosum		1.63	0.63		1	
Pseudoramibacter alactolyticus		0.02	0.27		1	
41012011	**Very good identification**					**41012011**
					SPO	
Clostridium difficile		97.83	0.61	URE 1	99	
Eubacterium limosum		1.62	0.29		1	
Clostridium novyi biovar A		0.53	0.37		99	
41012012	**Very good identification**					**41012012**
					SPO	
Clostridium difficile		97.88	0.61	BGL 1	99	
Eubacterium limosum		1.62	0.29		1	
Pseudoramibacter alactolyticus		0.19	0.11		1	
41012020	**Low discrimination**					**41012020**
					SPO	
Eubacterium limosum		73.78	0.63	NAG 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Pseudoramibacter alactolyticus		13.65	0.48		1	
Clostridium difficile		9.21	0.49		99	
41012022	**Low discrimination**					**41012022**
Atopobium minutum		42.43	0.48	MAN 1		
Pseudoramibacter alactolyticus		33.95	0.32			
Eubacterium limosum		16.69	0.29			
41012030	**Very good identification**					**41012030**
					SPO	
Clostridium difficile		97.88	0.61	NAG 1	99	
Eubacterium limosum		1.62	0.29		1	
Pseudoramibacter alactolyticus		0.30	0.14		1	
41012040	**Low discrimination**					**41012040**
					SPO	
Eubacterium limosum		47.27	0.63	SOR 1	1	
Clostridium difficile		30.76	0.61		99	
Pseudoramibacter alactolyticus		12.39	0.51		1	
41012050	**Very good identification**					**41012050**
					SPO	
Clostridium difficile		99.47	0.73	SOR 5	99	
Eubacterium limosum		0.32	0.29		1	
Pseudoramibacter alactolyticus		0.08	0.17		1	
41012051	**Acceptable identification**					**41012051**
					SPO	
Clostridium difficile		93.32	0.39	SOR 5	99	
Clostridium novyi biovar A		6.38	0.34	URE 1	99	
Eubacterium limosum		0.30	-.06		1	
41012052	**Good identification**					**41012052**
					SPO	
Clostridium difficile		97.16	0.39	SOR 5	99	
Clostridium novyi biovar A		1.17	0.21	BGL 1	99	
Pseudoramibacter alactolyticus		0.89	0.00		1	
41012060	**Low discrimination**					**41012060**
					SPO	
Pseudoramibacter alactolyticus		69.74	0.38	ESL 1	1	
Eubacterium limosum		15.23	0.29		1	
Clostridium difficile		9.91	0.27		99	
41012070	**Good identification**					**41012070**
					SPO	
Clostridium difficile		96.63	0.39	NAG 1	99	
Clostridium novyi biovar A		1.65	0.23	SOR 5	99	
Pseudoramibacter alactolyticus		1.41	0.04		1	
41012100	**Low discrimination**					**41012100**
					SPO	
Eubacterium limosum		87.51	0.63	LAC 1	1	
Clostridium difficile		10.93	0.49		99	
Pseudoramibacter alactolyticus		0.93	0.27		1	
41012110	**Very good identification**					**41012110**
					SPO	
Clostridium difficile		98.35	0.61	LAC 1	99	
Eubacterium limosum		1.63	0.29		1	
Pseudoramibacter alactolyticus		0.02	-.07		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
41012150	**Good identification**					**41012150**
					SPO	
Clostridium difficile		99.68	0.39	LAC 1	99	
Eubacterium limosum		0.32	-.06	SOR 5	1	
41012200	**Low discrimination**					**41012200**
					SPO	
Eubacterium limosum		87.70	0.63	RHA 1	1	
Clostridium difficile		10.95	0.49		99	
Pseudoramibacter alactolyticus		0.93	0.27		1	
41012210	**Very good identification**					**41012210**
					SPO	
Clostridium difficile		98.35	0.61	RHA 1	99	
Eubacterium limosum		1.63	0.29		1	
Pseudoramibacter alactolyticus		0.02	-.07		1	
41012250	**Good identification**					**41012250**
					SPO	
Clostridium difficile		99.68	0.39	RHA 1	99	
Eubacterium limosum		0.32	-.06	SOR 5	1	
41012400	**Low discrimination**					**41012400**
					SPO	
Eubacterium limosum		87.70	0.63	ARA 1	1	
Clostridium difficile		10.95	0.49		99	
Pseudoramibacter alactolyticus		0.93	0.27		1	
41012410	**Very good identification**					**41012410**
					SPO	
Clostridium difficile		98.35	0.61	ARA 1	99	
Eubacterium limosum		1.63	0.29		1	
Pseudoramibacter alactolyticus		0.02	-.07		1	
41012450	**Good identification**					**41012450**
					SPO	
Clostridium difficile		99.68	0.39	ARA 1	99	
Eubacterium limosum		0.32	-.06	SOR 5	1	
41013000	**Low discrimination**					**41013000**
					SPO	
Eubacterium limosum		84.76	0.63	GAL 1	1	
Clostridium difficile		10.58	0.49		99	
Atopobium minutum		3.27	0.51		1	
41013002	**Low discrimination**					**41013002**
					SPO	
Atopobium minutum		65.90	0.48	MAN 1	1	
Eubacterium limosum		25.92	0.29		1	
Clostridium difficile		3.24	0.15		99	
41013010	**Very good identification**					**41013010**
					SPO	
Clostridium difficile		98.23	0.61	GAL 1	99	
Eubacterium limosum		1.63	0.29		1	
Clostridium novyi biovar A		0.06	0.20		99	
41013020	**Low discrimination**					**41013020**
					CAT	
Atopobium minutum		45.49	0.43	MAN 1	1	
Eubacterium limosum		35.78	0.29		1	
Propionibacterium acnes		7.33	0.22		95	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41013022	**Acceptable identification**						**41013022**
					GRA		
Atopobium minutum		93.92	0.40	MAN 1	99		
Pseudoflavonifractor capillosus		2.68	-.01		1		
Pseudoramibacter alactolyticus		2.28	-.02		99		
41013050	**Good identification**						**41013050**
					SPO		
Clostridium difficile		98.93	0.39	GAL 1	99		
Clostridium novyi biovar A		0.75	0.17	SOR 5	99		
Eubacterium limosum		0.32	-.06		1		
41014000	**Low discrimination**						**41014000**
					SPO		
Clostridium glycolicum		89.06	0.78	SOR 95	99		
Eubacterium limosum		3.18	0.49		1		
Atopobium minutum		2.26	0.59		1		
41014001	**Low discrimination**						**41014001**
					SPO		
Clostridium glycolicum		83.90	0.43	SOR 95	99		
Clostridium novyi biovar A		5.99	0.37	URE 1	99		
Eubacterium limosum		2.99	0.15		1		
41014002	**Low discrimination**						**41014002**
					SPO		
Atopobium minutum		46.52	0.56	XYL 1	1		
Clostridium glycolicum		27.83	0.43		99		
Clostridium sporogenes		21.44	0.40		99		
41014010	**Low discrimination**						**41014010**
Clostridium difficile		86.70	0.57	MAN 90			
Clostridium glycolicum		11.33	0.43	XYL 5			
Clostridium novyi biovar A		0.81	0.37				
41014020	**Low discrimination**						**41014020**
					SPO		
Clostridium glycolicum		48.98	0.43	NAG 1	99		
Atopobium minutum		40.93	0.51	SOR 95	1		
Pseudoramibacter alactolyticus		3.69	0.19		1		
41014022	**Low discrimination**						**41014022**
					GRA	SPO	
Atopobium minutum		87.84	0.48	XYL 1	99	1	
Pseudoflavonifractor capillosus		7.53	0.15		1	1	
Clostridium glycolicum		1.59	0.09		99	99	
41014040	**Excellent identification**						**41014040**
					SPO		
Clostridium glycolicum		99.96	0.99		99		
Clostridium botulinum biovar A		0.02	0.36		99		
Pseudoramibacter alactolyticus		0.01	0.21		1		
41014041	**Very good identification**						**41014041**
Clostridium glycolicum		99.73	0.65	URE 1			
Clostridium novyi biovar A		0.25	0.34				
Clostridium botulinum biovar A		0.02	0.02				
41014042	**Very good identification**						**41014042**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium glycolicum		99.73	0.65	BGL 1	99		
Atopobium minutum		0.09	0.22		1		
Pseudoramibacter alactolyticus		0.06	0.05		1		
41014050	**Very good identification**						**41014050**
Clostridium glycolicum		97.67	0.65	MLZ 1			
Clostridium difficile		2.07	0.35				
Clostridium novyi biovar A		0.25	0.34				
41014060	**Very good identification**						**41014060**
					SPO		
Clostridium glycolicum		99.78	0.65	NAG 1	99		
Pseudoramibacter alactolyticus		0.10	0.09		1		
Clostridium novyi biovar A		0.06	0.23		99		
41014100	**Low discrimination**						**41014100**
					GRA	SPO	
Clostridium glycolicum		51.12	0.43	LAC 1	99	99	
Actinomyces meyeri		24.16	0.43	SOR 95	99	1	
Fusobacterium mortiferum		18.67	0.50		1	1	
41014140	**Very good identification**						**41014140**
					GRA	SPO	
Clostridium glycolicum		99.94	0.65	LAC 1	99	99	
Actinomyces meyeri		0.03	0.09		99	1	
Fusobacterium mortiferum		0.02	0.16		1	1	
41014200	**Low discrimination**						**41014200**
					SPO		
Clostridium glycolicum		89.41	0.43	RHA 1	99		
Eubacterium limosum		3.19	0.15	SOR 95	1		
Atopobium minutum		2.26	0.25		1		
41014240	**Very good identification**						**41014240**
Clostridium glycolicum		99.98	0.65	RHA 1			
Clostridium botulinum biovar A		0.02	0.02				
41014400	**Low discrimination**						**41014400**
					SPO		
Clostridium glycolicum		87.11	0.43	ARA 1	99		
Eubacterium limosum		3.11	0.15	SOR 95	1		
Actinomyces meyeri		2.57	0.22		1		
41014440	**Very good identification**						**41014440**
Clostridium glycolicum		99.98	0.65	ARA 1			
Clostridium botulinum biovar A		0.02	0.02				
41015000	**Low discrimination**						**41015000**
					GRA	SPO	
Clostridium glycolicum		88.87	0.61	GAL 10	99	99	
Atopobium minutum		6.75	0.51	SOR 95	99	1	
Fusobacterium mortiferum		1.90	0.47		1	1	
41015002	**Low discrimination**						**41015002**
					SPO		
Atopobium minutum		75.82	0.48	XYL 1	1		
Clostridium glycolicum		15.12	0.27		99		
Lactobacillus fermentum		3.71	0.32		1		
41015020	**Low discrimination**						**41015020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Atopobium minutum		65.19	0.43	XYL 1	1		
Clostridium glycolicum		26.01	0.27		99		
Actinomyces meyeri		5.66	0.21		1		
41015022	**Acceptable identification**						**41015022**
					GRA		
Atopobium minutum		92.81	0.40	XYL 1	99		
Lactobacillus fermentum		4.54	0.24		99		
Pseudoflavonifractor capillosus		2.65	-.01		1		
41015040	**Very good identification**						**41015040**
					SPO		
Clostridium glycolicum		99.99	0.83	GAL 10	99		
Atopobium minutum		0.00	0.17		1		
Clostridium novyi biovar A		0.00	0.17		99		
41015041	**Good identification**						**41015041**
Clostridium glycolicum		99.75	0.49	GAL 10			
Clostridium novyi biovar A		0.25	0.17	URE 1			
41015042	**Good identification**						**41015042**
					SPO		
Clostridium glycolicum		99.73	0.49	GAL 10	99		
Atopobium minutum		0.27	0.14	BGL 1	1		
41015050	**Good identification**						**41015050**
Clostridium glycolicum		99.56	0.49	GAL 10			
Clostridium novyi biovar A		0.25	0.17	MLZ 1			
Clostridium difficile		0.19	0.01				
41015060	**Good identification**						**41015060**
					SPO		
Clostridium glycolicum		99.87	0.49	GAL 10	99		
Atopobium minutum		0.13	0.09	NAG 1	1		
41015100	**Low discrimination**						**41015100**
					GRA	SPO	
Actinomyces meyeri		49.72	0.46	ESL 1	99	1	
Fusobacterium mortiferum		38.42	0.53	SUC 93	1	1	
Clostridium glycolicum		7.79	0.27		99	99	
41015102	**Low discrimination**						**41015102**
					GRA		
Fusobacterium mortiferum		45.81	0.37	XYL 7	1		
Lactobacillus fermentum		33.46	0.36	BGL 10	99		
Actinomyces meyeri		16.50	0.20		99		
41015140	**Good identification**						**41015140**
					GRA	SPO	
Clostridium glycolicum		99.40	0.49	GAL 10	99	99	
Actinomyces meyeri		0.34	0.12	LAC 1	99	1	
Fusobacterium mortiferum		0.26	0.19		1	1	
41015240	**Good identification**						**41015240**
Clostridium glycolicum		100.0	0.49	GAL 10			
41015440	**Good identification**						**41015440**
Clostridium glycolicum		100.0	0.49	GAL 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
41015500	**Low discrimination**					**41015500**
					GRA	
Actinomyces meyeri		78.00	0.35	ESL 1	99	
Lactobacillus fermentum		19.57	0.36	SUC 93	99	
Fusobacterium mortiferum		2.44	0.19		1	
41015502	**Low discrimination**					**41015502**
Lactobacillus fermentum		89.03	0.36	ESL 1		
Actinomyces meyeri		10.98	0.10	SUC 87		
41016000	**Low discrimination**					**41016000**
					SPO	
Eubacterium limosum		58.40	0.63	XYL 1	1	
Clostridium difficile		38.00	0.61		99	
Clostridium glycolicum		2.69	0.43		99	
41016010	**Very good identification**					**41016010**
					SPO	
Clostridium difficile		99.67	0.73	XYL 5	99	
Eubacterium limosum		0.32	0.29		1	
Clostridium glycolicum		0.02	0.09		99	
41016011	**Good identification**					**41016011**
					SPO	
Clostridium difficile		99.68	0.39	XYL 5	99	
Eubacterium limosum		0.32	-.06	URE 1	1	
41016012	**Good identification**					**41016012**
					SPO	
Clostridium difficile		99.68	0.39	XYL 5	99	
Eubacterium limosum		0.32	-.06	BGL 1	1	
41016030	**Good identification**					**41016030**
					SPO	
Clostridium difficile		99.68	0.39	XYL 5	99	
Eubacterium limosum		0.32	-.06	NAG 1	1	
41016040	**Good identification**					**41016040**
					SPO	
Clostridium glycolicum		94.75	0.65	MAN 1	99	
Clostridium difficile		3.70	0.40		99	
Eubacterium limosum		1.09	0.29		1	
41016050	**Acceptable identification**					**41016050**
					SPO	
Clostridium difficile		94.92	0.51	XYL 5	99	
Clostridium glycolicum		5.02	0.31	SOR 5	99	
Eubacterium limosum		0.06	-.06		1	
41016110	**Good identification**					**41016110**
					SPO	
Clostridium difficile		99.68	0.39	XYL 5	99	
Eubacterium limosum		0.32	-.06	LAC 1	1	
41016210	**Good identification**					**41016210**
					SPO	
Clostridium difficile		99.68	0.39	XYL 5	99	
Eubacterium limosum		0.32	-.06	RHA 1	1	
41016410	**Good identification**					**41016410**
					SPO	
Clostridium difficile		99.68	0.39	XYL 5	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium limosum		0.32	-.06	ARA 1	1	
41017010	**Good identification**					**41017010**
					SPO	
Clostridium difficile		99.68	0.39	GAL 1	99	
Eubacterium limosum		0.32	-.06	XYL 5	1	
41017040	**Good identification**					**41017040**
					SPO	
Clostridium glycolicum		99.54	0.49	GAL 10	99	
Clostridium difficile		0.35	0.05	MAN 1	99	
Eubacterium limosum		0.10	-.06		1	
41020000	**Low discrimination**					**41020000**
					SPO	
Clostridium botulinum biovar A		50.23	0.90		99	
Clostridium sporogenes		49.25	0.87		99	
Atopobium minutum		0.31	0.60		1	
41020001	**Low discrimination**					**41020001**
Clostridium botulinum biovar A		46.83	0.56	URE 1		
Clostridium sporogenes		45.92	0.53			
Clostridium novyi biovar A		6.76	0.53			
41020002	**Very good identification**					**41020002**
					SPO	
Clostridium sporogenes		95.75	0.79		99	
Clostridium botulinum biovar A		2.96	0.56		99	
Atopobium minutum		1.20	0.57		1	
41020003	**Low discrimination**					**41020003**
Clostridium sporogenes		89.17	0.45	URE 1		
Clostridium novyi biovar A		6.95	0.40			
Clostridium botulinum biovar A		2.76	0.21			
41020010	**Low discrimination**					**41020010**
Clostridium botulinum biovar A		46.33	0.56	MLZ 1		
Clostridium sporogenes		45.43	0.53			
Clostridium novyi biovar A		6.69	0.53			
41020011	**Low discrimination**					**41020011**
Clostridium novyi biovar A		87.83	0.53	TRE 1		
Clostridium botulinum biovar A		6.14	0.21			
Clostridium sporogenes		6.03	0.19			
41020012	**Low discrimination**					**41020012**
Clostridium sporogenes		89.17	0.45	MLZ 1		
Clostridium novyi biovar A		6.95	0.40			
Clostridium botulinum biovar A		2.76	0.21			
41020013	**Low discrimination**					**41020013**
Clostridium novyi biovar A		88.53	0.40	TRE 1		
Clostridium sporogenes		11.47	0.11			
41020020	**Low discrimination**					**41020020**
					SPO	
Clostridium botulinum biovar A		44.67	0.56	NAG 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		43.80	0.53		99		
Atopobium minutum		9.09	0.51		1		
41020021	**Low discrimination**						**41020021**
Clostridium novyi biovar A		62.07	0.43	TRE 1			
Clostridium botulinum biovar A		17.37	0.21				
Clostridium sporogenes		17.03	0.19				
41020022	**Low discrimination**						**41020022**
					GRA	SPO	
Clostridium sporogenes		56.80	0.45	NAG 1	99	99	
Atopobium minutum		23.56	0.48		99	1	
Pseudoflavonifractor capillosus		16.78	0.32		1	1	
41020030	**Low discrimination**						**41020030**
Clostridium novyi biovar A		62.07	0.43	TRE 1			
Clostridium botulinum biovar A		17.37	0.21				
Clostridium sporogenes		17.03	0.19				
41020031	**Good identification**						**41020031**
Clostridium novyi biovar A		100.0	0.43	TRE 1			
41020040	**Very good identification**						**41020040**
Clostridium botulinum biovar A		96.36	0.81				
Clostridium sporogenes		3.20	0.53				
Clostridium novyi biovar A		0.31	0.50				
41020041	**Low discrimination**						**41020041**
Clostridium botulinum biovar A		73.78	0.47	URE 1			
Clostridium novyi biovar A		23.78	0.50				
Clostridium sporogenes		2.45	0.19				
41020042	**Low discrimination**						**41020042**
Clostridium sporogenes		50.55	0.45	SOR 1			
Clostridium botulinum biovar A		46.19	0.47				
Clostridium novyi biovar A		2.63	0.37				
41020043	**Low discrimination**						**41020043**
Clostridium novyi biovar A		72.89	0.37	TRE 1			
Clostridium sporogenes		14.17	0.11				
Clostridium botulinum biovar A		12.95	0.12				
41020050	**Low discrimination**						**41020050**
Clostridium botulinum biovar A		73.54	0.47	MLZ 1			
Clostridium novyi biovar A		23.70	0.50				
Clostridium sporogenes		2.44	0.19				
41020051	**Good identification**						**41020051**
Clostridium novyi biovar A		96.96	0.50	TRE 1			
Clostridium botulinum biovar A		3.04	0.12				
41020052	**Low discrimination**						**41020052**
Clostridium novyi biovar A		72.89	0.37	TRE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sporogenes		14.17	0.11			
Clostridium botulinum biovar A		12.95	0.12			
41020053	**Good identification**					**41020053**
Clostridium novyi biovar A		100.0	0.37	TRE 1		
41020060	**Low discrimination**					**41020060**
Clostridium botulinum biovar A		89.12	0.47	NAG 1		
Clostridium novyi biovar A		7.18	0.40			
Clostridium sporogenes		2.96	0.19			
41020061	**Low discrimination**					**41020061**
Clostridium novyi biovar A		88.86	0.40	TRE 1		
Clostridium botulinum biovar A		11.14	0.12			
41020070	**Low discrimination**					**41020070**
Clostridium novyi biovar A		88.86	0.40	TRE 1		
Clostridium botulinum biovar A		11.14	0.12			
41020071	**Good identification**					**41020071**
Clostridium novyi biovar A		100.0	0.40	TRE 1		
41020100	**Low discrimination**					**41020100**
					SPO	
Clostridium botulinum biovar A		50.15	0.56	LAC 1	99	
Clostridium sporogenes		49.18	0.53		99	
Atopobium minutum		0.31	0.25		1	
41020102	**Good identification**					**41020102**
					SPO	
Clostridium sporogenes		95.83	0.45	LAC 1	99	
Clostridium botulinum biovar A		2.96	0.21		99	
Atopobium minutum		1.21	0.22		1	
41020140	**Good identification**					**41020140**
Clostridium botulinum biovar A		96.49	0.47	LAC 1		
Clostridium sporogenes		3.20	0.19			
Clostridium novyi biovar A		0.31	0.16			
41020200	**Low discrimination**					**41020200**
					SPO	
Clostridium botulinum biovar A		50.24	0.56	RHA 1	99	
Clostridium sporogenes		49.26	0.53		99	
Atopobium minutum		0.31	0.25		1	
41020202	**Good identification**					**41020202**
					SPO	
Clostridium sporogenes		95.83	0.45	RHA 1	99	
Clostridium botulinum biovar A		2.96	0.21		99	
Atopobium minutum		1.21	0.22		1	
41020240	**Good identification**					**41020240**
Clostridium botulinum biovar A		96.49	0.47	RHA 1		
Clostridium sporogenes		3.20	0.19			
Clostridium novyi biovar A		0.31	0.16			
41020400	**Low discrimination**					**41020400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium botulinum biovar A		50.24	0.56	ARA 1	99		
Clostridium sporogenes		49.26	0.53		99		
Atopobium minutum		0.31	0.25		1		
41020402	**Good identification**						**41020402**
					SPO		
Clostridium sporogenes		95.83	0.45	ARA 1	99		
Clostridium botulinum biovar A		2.96	0.21		99		
Atopobium minutum		1.21	0.22		1		
41020440	**Good identification**						**41020440**
Clostridium botulinum biovar A		96.49	0.47	ARA 1			
Clostridium sporogenes		3.20	0.19				
Clostridium novyi biovar A		0.31	0.16				
41021000	**Low discrimination**						**41021000**
					SPO		
Clostridium botulinum biovar A		45.35	0.56	GAL 1	99		
Clostridium sporogenes		44.47	0.53		99		
Atopobium minutum		9.22	0.51		1		
41021001	**Low discrimination**						**41021001**
Clostridium novyi biovar A		42.11	0.37	TRE 1			
Clostridium botulinum biovar A		26.51	0.21	GAL 10			
Clostridium sporogenes		25.99	0.19				
41021002	**Low discrimination**						**41021002**
					SPO		
Clostridium sporogenes		68.76	0.45	GAL 1	99		
Atopobium minutum		28.52	0.48		1		
Clostridium botulinum biovar A		2.13	0.21		99		
41021010	**Low discrimination**						**41021010**
Clostridium novyi biovar A		42.11	0.37	TRE 1			
Clostridium botulinum biovar A		26.51	0.21	GAL 10			
Clostridium sporogenes		25.99	0.19				
41021011	**Good identification**						**41021011**
Clostridium novyi biovar A		100.0	0.37	TRE 1			
41021020	**Low discrimination**						**41021020**
					SPO		
Atopobium minutum		73.84	0.43	TRE 1	1		
Clostridium botulinum biovar A		11.00	0.21		99		
Clostridium sporogenes		10.79	0.19		99		
41021022	**Low discrimination**						**41021022**
					GRA	SPO	
Atopobium minutum		75.41	0.40	TRE 1	99	1	
Pseudoflavonifractor capillosus		17.90	0.15		1	1	
Clostridium sporogenes		5.51	0.11		99	99	
41021040	**Acceptable identification**						**41021040**
Clostridium botulinum biovar A		92.16	0.47	GAL 1			
Clostridium novyi biovar A		3.30	0.34				
Clostridium sporogenes		3.06	0.19				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
41022000	**Low discrimination**					**41022000**
					SPO	
Clostridium botulinum biovar A		47.92	0.56	MAN 1	99	
Clostridium sporogenes		46.99	0.53		99	
Eubacterium limosum		2.36	0.29		1	
41022002	**Good identification**					**41022002**
					SPO	
Clostridium sporogenes		95.57	0.45	MAN 1	99	
Clostridium botulinum biovar A		2.95	0.21		99	
Atopobium minutum		1.20	0.22		1	
41022010	**Acceptable identification**					**41022010**
Clostridium difficile		91.06	0.42	FRU 99		
Clostridium botulinum biovar A		4.11	0.21	TRE 7		
Clostridium sporogenes		4.03	0.19			
41022040	**Good identification**					**41022040**
Clostridium botulinum biovar A		95.59	0.47	MAN 1		
Clostridium sporogenes		3.17	0.19			
Clostridium difficile		0.77	0.08			
41024000	**Low discrimination**					**41024000**
Clostridium botulinum biovar A		49.61	0.56	XYL 1		
Clostridium sporogenes		48.65	0.53			
Clostridium glycolicum		1.24	0.27			
41024002	**Good identification**					**41024002**
					SPO	
Clostridium sporogenes		95.83	0.45	XYL 1	99	
Clostridium botulinum biovar A		2.96	0.21		99	
Atopobium minutum		1.21	0.22		1	
41024040	**Low discrimination**					**41024040**
Clostridium glycolicum		60.51	0.49	FRU 90		
Clostridium botulinum biovar A		38.11	0.47	TRE 1		
Clostridium sporogenes		1.26	0.19			
41030000	**Low discrimination**					**41030000**
					SPO	
Clostridium sporogenes		66.90	0.74		99	
Clostridium botulinum biovar A		24.68	0.69		99	
Eubacterium limosum		3.10	0.49		1	
41030001	**Low discrimination**					**41030001**
Clostridium sporogenes		63.03	0.40	URE 1		
Clostridium botulinum biovar A		23.25	0.35			
Clostridium novyi biovar A		5.84	0.37			
41030002	**Good identification**					**41030002**
					SPO	
Clostridium sporogenes		92.21	0.66		99	
Atopobium minutum		6.06	0.56		1	
Clostridium botulinum biovar A		1.03	0.35		99	
41030010	**Good identification**					**41030010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		90.14	0.60	TRE 7			
Clostridium sporogenes		6.34	0.40	MAN 90			
Clostridium botulinum biovar A		2.34	0.35				
41030020	**Low discrimination**						**41030020**
					SPO		
Atopobium minutum		40.22	0.51	TRE 1	1		
Clostridium sporogenes		37.07	0.40		99		
Clostridium botulinum biovar A		13.67	0.35		99		
41030022	**Low discrimination**						**41030022**
					GRA	SPO	
Atopobium minutum		63.60	0.48	TRE 1	99	1	
Clostridium sporogenes		29.31	0.32		99	99	
Pseudoflavonifractor capillosus		5.45	0.15		1	1	
41030040	**Low discrimination**						**41030040**
Clostridium botulinum biovar A		79.70	0.60	FRU 6			
Clostridium glycolicum		9.49	0.43				
Clostridium sporogenes		7.31	0.40				
41030050	**Low discrimination**						**41030050**
Clostridium difficile		79.31	0.38	TRE 7			
Clostridium botulinum biovar A		11.68	0.26	MAN 90			
Clostridium novyi biovar A		6.55	0.34	SOR 5			
41030100	**Low discrimination**						**41030100**
					GRA	SPO	
Fusobacterium mortiferum		55.87	0.60		1	1	
Clostridium sporogenes		29.69	0.40		99	99	
Clostridium botulinum biovar A		10.95	0.35		99	99	
41030200	**Low discrimination**						**41030200**
					SPO		
Clostridium sporogenes		67.06	0.40	RHA 1	99		
Clostridium botulinum biovar A		24.74	0.35		99		
Eubacterium limosum		3.11	0.15		1		
41030400	**Low discrimination**						**41030400**
					SPO		
Clostridium sporogenes		67.06	0.40	ARA 1	99		
Clostridium botulinum biovar A		24.74	0.35		99		
Eubacterium limosum		3.11	0.15		1		
41031000	**Low discrimination**						**41031000**
					GRA	SPO	
Fusobacterium mortiferum		32.09	0.57		1	1	
Atopobium minutum		28.78	0.51		99	1	
Clostridium sporogenes		26.53	0.40		99	99	
41031002	**Low discrimination**						**41031002**
					GRA	SPO	
Atopobium minutum		60.54	0.48	TRE 1	99	1	
Clostridium sporogenes		27.90	0.32		99	99	
Fusobacterium mortiferum		11.25	0.41		1	1	
41031020	**Acceptable identification**						**41031020**
					GRA	SPO	
Atopobium minutum		92.67	0.43	TRE 1	99	1	
Fusobacterium mortiferum		3.13	0.23		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		2.59	0.06		99	99	
41031022	**Good identification**						**41031022**
					GRA	SPO	
Atopobium minutum		95.92	0.40	TRE 1	99	1	
Pseudoflavonifractor capillosus		2.74	-.01		1	1	
Clostridium sporogenes		1.34	-.02		99	99	
41031100	**Very good identification**						**41031100**
					GRA	SPO	
Fusobacterium mortiferum		98.68	0.63		1	1	
Atopobium minutum		0.38	0.17		99	1	
Clostridium sporogenes		0.35	0.06		99	99	
41031102	**Good identification**						**41031102**
					GRA	SPO	
Fusobacterium mortiferum		96.71	0.47	BGL 10	1	1	
Atopobium minutum		2.25	0.14		99	1	
Clostridium sporogenes		1.04	-.02		99	99	
41032000	**Low discrimination**						**41032000**
					SPO		
Eubacterium limosum		50.17	0.63	TRE 1	1		
Clostridium difficile		46.68	0.64		99		
Clostridium sporogenes		1.78	0.40		99		
41032010	**Very good identification**						**41032010**
					SPO		
Clostridium difficile		99.77	0.76	TRE 7	99		
Eubacterium limosum		0.22	0.29		1		
Clostridium sporogenes		0.01	0.06		99		
41032011	**Good identification**						**41032011**
					SPO		
Clostridium difficile		99.78	0.42	TRE 7	99		
Eubacterium limosum		0.22	-.06	URE 1	1		
41032012	**Good identification**						**41032012**
					SPO		
Clostridium difficile		99.52	0.42	TRE 7	99		
Clostridium sporogenes		0.26	-.02	BGL 1	99		
Eubacterium limosum		0.22	-.06		1		
41032030	**Good identification**						**41032030**
					SPO		
Clostridium difficile		99.78	0.42	TRE 7	99		
Eubacterium limosum		0.22	-.06	NAG 1	1		
41032040	**Low discrimination**						**41032040**
					SPO		
Clostridium difficile		72.19	0.42	TRE 7	99		
Eubacterium limosum		14.89	0.29	SOR 5	1		
Clostridium botulinum biovar A		5.77	0.26		99		
41032050	**Good identification**						**41032050**
					SPO		
Clostridium difficile		99.96	0.54	TRE 7	99		
Eubacterium limosum		0.04	-.06	SOR 5	1		
41032110	**Good identification**						**41032110**
					SPO		
Clostridium difficile		99.78	0.42	TRE 7	99		
Eubacterium limosum		0.22	-.06	LAC 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
41032210	**Good identification**						**41032210**
					SPO		
Clostridium difficile		99.78	0.42	TRE 7	99		
Eubacterium limosum		0.22	-.06	RHA 1	1		
41032410	**Good identification**						**41032410**
					SPO		
Clostridium difficile		99.78	0.42	TRE 7	99		
Eubacterium limosum		0.22	-.06	ARA 1	1		
41033010	**Good identification**						**41033010**
					SPO		
Clostridium difficile		99.78	0.42	TRE 7	99		
Eubacterium limosum		0.22	-.06	GAL 1	1		
41034000	**Low discrimination**						**41034000**
Clostridium glycolicum		43.73	0.43	TRE 1			
Clostridium sporogenes		33.68	0.40	SOR 95			
Clostridium botulinum biovar A		12.42	0.35				
41034010	**Good identification**						**41034010**
Clostridium difficile		96.52	0.38	TRE 7			
Clostridium glycolicum		1.69	0.09	MAN 90			
Clostridium sporogenes		1.30	0.06	XYL 5			
41034040	**Very good identification**						**41034040**
Clostridium glycolicum		99.48	0.65	TRE 1			
Clostridium botulinum biovar A		0.44	0.26				
Clostridium sporogenes		0.04	0.06				
41034100	**Low discrimination**						**41034100**
					GRA	SPO	
Fusobacterium mortiferum		81.04	0.41	XYL 7	1	1	
Clostridium glycolicum		7.50	0.09		99	99	
Clostridium sporogenes		5.78	0.06		99	99	
41035040	**Good identification**						**41035040**
Clostridium glycolicum		100.0	0.49	TRE 1			
41035100	**Acceptable identification**						**41035100**
					GRA		
Fusobacterium mortiferum		92.88	0.44	XYL 7	1		
Actinomyces meyeri		4.06	0.12		99		
Eubacterium saburreum		1.96	0.14		85		
41036000	**Low discrimination**						**41036000**
					SPO		
Clostridium difficile		81.44	0.42	TRE 7	99		
Eubacterium limosum		16.80	0.29	XYL 5	1		
Clostridium glycolicum		0.77	0.09		99		
41036010	**Good identification**						**41036010**
					SPO		
Clostridium difficile		99.96	0.54	TRE 7	99		
Eubacterium limosum		0.04	-.06	XYL 5	1		
41040000	**Low discrimination**						**41040000**
					SPO		
Clostridium botulinum biovar A		50.17	0.66	CEL 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		36.90	0.61		99		
Atopobium minutum		7.65	0.60		1		
41040001	**Low discrimination**						**41040001**
Clostridium novyi biovar A		64.15	0.53	CEL 1			
Clostridium botulinum biovar A		17.95	0.32				
Clostridium sporogenes		13.20	0.27				
41040002	**Low discrimination**						**41040002**
					SPO		
Clostridium sporogenes		67.32	0.53	CEL 1	99		
Atopobium minutum		27.93	0.57		1		
Clostridium botulinum biovar A		2.77	0.32		99		
41040003	**Low discrimination**						**41040003**
					SPO		
Clostridium novyi biovar A		64.52	0.40	CEL 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41040010	**Low discrimination**						**41040010**
Clostridium novyi biovar A		60.04	0.53	CEL 1			
Clostridium botulinum biovar A		16.80	0.32				
Clostridium sporogenes		12.36	0.27				
41040011	**Good identification**						**41040011**
Clostridium novyi biovar A		100.0	0.53	CEL 1			
41040012	**Low discrimination**						**41040012**
					SPO		
Clostridium novyi biovar A		64.52	0.40	CEL 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41040013	**Good identification**						**41040013**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
41040020	**Low discrimination**						**41040020**
					SPO		
Atopobium minutum		62.92	0.51	CEL 1	1		
Clostridium botulinum biovar A		12.50	0.32		99		
Clostridium novyi biovar A		11.17	0.43		99		
41040021	**Acceptable identification**						**41040021**
					SPO		
Clostridium novyi biovar A		94.62	0.43	CEL 1	99		
Atopobium minutum		5.38	0.17		1		
41040022	**Low discrimination**						**41040022**
					GRA	SPO	
Atopobium minutum		54.58	0.48	CEL 1	99	1	
Pseudoflavonifractor capillosus		38.87	0.32		1	1	
Clostridium sporogenes		3.99	0.19		99	99	
41040030	**Acceptable identification**						**41040030**
					SPO		
Clostridium novyi biovar A		94.62	0.43	CEL 1	99		
Atopobium minutum		5.38	0.17		1		
41040031	**Good identification**						**41040031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.43	CEL 1		
41040040	**Low discrimination**					**41040040**
Clostridium botulinum biovar A		87.87	0.57	CEL 1		
Clostridium novyi biovar A		7.08	0.50			
Clostridium sporogenes		2.19	0.27			
41040041	**Low discrimination**					**41040041**
Clostridium novyi biovar A		88.86	0.50	CEL 1		
Clostridium botulinum biovar A		11.14	0.23			
41040042	**Low discrimination**					**41040042**
Clostridium novyi biovar A		39.44	0.37	CEL 1		
Clostridium botulinum biovar A		28.02	0.23			
Clostridium sporogenes		23.00	0.19			
41040043	**Good identification**					**41040043**
Clostridium novyi biovar A		100.0	0.37	CEL 1		
41040050	**Low discrimination**					**41040050**
Clostridium novyi biovar A		88.12	0.50	CEL 1		
Clostridium botulinum biovar A		11.05	0.23			
Clostridium difficile		0.84	0.01			
41040051	**Good identification**					**41040051**
Clostridium novyi biovar A		100.0	0.50	CEL 1		
41040052	**Good identification**					**41040052**
Clostridium novyi biovar A		100.0	0.37	CEL 1		
41040053	**Good identification**					**41040053**
Clostridium novyi biovar A		100.0	0.37	CEL 1		
41040060	**Low discrimination**					**41040060**
					SPO	
Clostridium novyi biovar A		62.29	0.40	CEL 1	99	
Clostridium botulinum biovar A		31.24	0.23		99	
Atopobium minutum		5.32	0.17		1	
41040061	**Good identification**					**41040061**
Clostridium novyi biovar A		100.0	0.40	CEL 1		
41040070	**Good identification**					**41040070**
Clostridium novyi biovar A		100.0	0.40	CEL 1		
41040071	**Good identification**					**41040071**
Clostridium novyi biovar A		100.0	0.40	CEL 1		
41041000	**Low discrimination**					**41041000**
					SPO	
Atopobium minutum		67.43	0.51	CEL 1	1	
Clostridium botulinum biovar A		13.40	0.32		99	
Clostridium sporogenes		9.85	0.27		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41041001	**Low discrimination**						**41041001**
					SPO		
Clostridium novyi biovar A		88.65	0.37	CEL 1	99		
Atopobium minutum		11.35	0.17	GAL 10	1		
41041002	**Acceptable identification**						**41041002**
					SPO		
Atopobium minutum		90.63	0.48	CEL 1	1		
Clostridium sporogenes		6.62	0.19		99		
Clostridium novyi biovar A		1.89	0.24		99		
41041010	**Low discrimination**						**41041010**
					SPO		
Clostridium novyi biovar A		88.65	0.37	CEL 1	99		
Atopobium minutum		11.35	0.17	GAL 10	1		
41041011	**Good identification**						**41041011**
Clostridium novyi biovar A		100.0	0.37	CEL 1			
41041020	**Acceptable identification**						**41041020**
					SPO		
Atopobium minutum		92.76	0.43	CEL 1	1		
Clostridium novyi biovar A		5.49	0.26		99		
Eubacterium tortuosum		1.75	0.03		1		
41041022	**Low discrimination**						**41041022**
					GRA	SPO	
Atopobium minutum		79.80	0.40	CEL 1	99	1	
Pseudoflavonifractor capillosus		18.95	0.15		1	1	
Clostridium novyi biovar A		1.25	0.13		99	99	
41042010	**Good identification**						**41042010**
					SPO		
Clostridium difficile		98.77	0.39	FRU 99	99		
Clostridium novyi biovar A		0.92	0.19	CEL 5	99		
Eubacterium limosum		0.31	-.06		1		
41044040	**Good identification**						**41044040**
Clostridium glycolicum		97.32	0.49	FRU 90			
Clostridium botulinum biovar A		2.48	0.23	CEL 1			
Clostridium novyi biovar A		0.20	0.16				
41050000	**Low discrimination**						**41050000**
					SPO		
Eubacterium limosum		28.36	0.49	CEL 1	1		
Atopobium minutum		20.14	0.59	MAN 86	1		
Clostridium sporogenes		18.56	0.48		99		
41050001	**Low discrimination**						**41050001**
					SPO		
Clostridium novyi biovar A		36.36	0.37	FRU 10	99		
Eubacterium limosum		18.16	0.15	CEL 1	1		
Atopobium minutum		12.89	0.25		1		
41050002	**Low discrimination**						**41050002**
					GRA	SPO	
Atopobium minutum		63.15	0.56	CEL 1	99	1	
Clostridium sporogenes		29.10	0.40		99	99	
Fusobacterium mortiferum		2.91	0.39		1	1	
41050010	**Very good identification**						**41050010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium difficile		97.72	0.57	CEL 5	99		
Clostridium novyi biovar A		0.91	0.37	MAN 90	99		
Eubacterium limosum		0.46	0.15		1		
41050020	**Low discrimination**						**41050020**
Atopobium minutum		81.14	0.51	CEL 1			
Pseudoramibacter alactolyticus		7.31	0.19				
Eubacterium limosum		3.46	0.15				
41050022	**Low discrimination**						**41050022**
					GRA		
Atopobium minutum		89.26	0.48	CEL 1	99		
Pseudoflavonifractor capillosus		7.65	0.15		1		
Pseudoramibacter alactolyticus		1.34	0.03		99		
41050040	**Low discrimination**						**41050040**
					SPO		
Clostridium glycolicum		58.91	0.43	CEL 1	99		
Clostridium botulinum biovar A		20.00	0.36	XYL 95	99		
Pseudoramibacter alactolyticus		6.28	0.21		1		
41050050	**Low discrimination**						**41050050**
Clostridium difficile		86.83	0.35	CEL 5			
Clostridium novyi biovar A		10.26	0.34	MAN 90			
Clostridium glycolicum		2.18	0.09	SOR 5			
41050100	**Good identification**						**41050100**
					GRA		
Fusobacterium mortiferum		93.18	0.61		1		
Eubacterium limosum		2.06	0.15		99		
Atopobium minutum		1.46	0.25		99		
41050102	**Low discrimination**						**41050102**
					GRA	SPO	
Fusobacterium mortiferum		87.93	0.45	BGL 10	1	1	
Atopobium minutum		8.26	0.22		99	1	
Clostridium sporogenes		3.81	0.06		99	99	
41051000	**Low discrimination**						**41051000**
					GRA		
Fusobacterium mortiferum		52.52	0.58		1		
Atopobium minutum		42.21	0.51		99		
Eubacterium limosum		1.80	0.15		99		
41051002	**Low discrimination**						**41051002**
					GRA	SPO	
Atopobium minutum		81.88	0.48	CEL 1	99	1	
Fusobacterium mortiferum		16.98	0.42		1	1	
Clostridium sporogenes		1.14	0.06		99	99	
41051020	**Good identification**						**41051020**
					GRA	SPO	
Atopobium minutum		95.71	0.43	CEL 1	99	1	
Fusobacterium mortiferum		3.61	0.24		1	1	
Clostridium novyi biovar A		0.68	0.10		99	99	
41051022	**Good identification**						**41051022**
					GRA		
Atopobium minutum		97.22	0.40	CEL 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Pseudoflavonifractor capillosus		2.78	-.01		1	
41051100	**Very good identification**					**41051100**
					GRA	
Fusobacterium mortiferum		99.50	0.64		1	
Atopobium minutum		0.35	0.17		99	
Actinomyces meyeri		0.15	0.07		99	
41051102	**Good identification**					**41051102**
					GRA	
Fusobacterium mortiferum		97.96	0.48	BGL 10	1	
Atopobium minutum		2.05	0.14		99	
41052000	**Low discrimination**					**41052000**
					SPO	
Eubacterium limosum		60.01	0.63	CEL 1	1	
Clostridium difficile		39.05	0.61		99	
Pseudoramibacter alactolyticus		0.64	0.27		1	
41052010	**Very good identification**					**41052010**
					SPO	
Clostridium difficile		99.68	0.73	CEL 5	99	
Eubacterium limosum		0.32	0.29		1	
Pseudoramibacter alactolyticus		0.00	-.07		1	
41052011	**Good identification**					**41052011**
					SPO	
Clostridium difficile		99.68	0.39	CEL 5	99	
Eubacterium limosum		0.32	-.06	URE 1	1	
41052012	**Good identification**					**41052012**
					SPO	
Clostridium difficile		99.68	0.39	CEL 5	99	
Eubacterium limosum		0.32	-.06	BGL 1	1	
41052030	**Good identification**					**41052030**
					SPO	
Clostridium difficile		99.68	0.39	CEL 5	99	
Eubacterium limosum		0.32	-.06	NAG 1	1	
41052040	**Low discrimination**					**41052040**
					SPO	
Clostridium difficile		69.89	0.40	CEL 5	99	
Eubacterium limosum		20.62	0.29	SOR 5	1	
Pseudoramibacter alactolyticus		5.40	0.17		1	
41052050	**Good identification**					**41052050**
					SPO	
Clostridium difficile		99.94	0.51	CEL 5	99	
Eubacterium limosum		0.06	-.06	SOR 5	1	
41052110	**Good identification**					**41052110**
					SPO	
Clostridium difficile		99.68	0.39	CEL 5	99	
Eubacterium limosum		0.32	-.06	LAC 1	1	
41052210	**Good identification**					**41052210**
					SPO	
Clostridium difficile		99.68	0.39	CEL 5	99	
Eubacterium limosum		0.32	-.06	RHA 1	1	
41052410	**Good identification**					**41052410**
					SPO	
Clostridium difficile		99.68	0.39	CEL 5	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		0.32	-.06	ARA 1	1		
41053010	**Good identification**						**41053010**
					SPO		
Clostridium difficile		99.68	0.39	CEL 5	99		
Eubacterium limosum		0.32	-.06	GAL 1	1		
41054000	**Low discrimination**						**41054000**
					GRA	SPO	
Clostridium glycolicum		80.82	0.43	CEL 1	99	99	
Clostridium difficile		6.67	0.23	SOR 95	99	99	
Fusobacterium mortiferum		4.22	0.36		1	1	
41054010	**Good identification**						**41054010**
Clostridium difficile		97.55	0.35	CEL 5			
Clostridium glycolicum		2.45	0.09	MAN 90			
41054040	**Very good identification**						**41054040**
Clostridium glycolicum		99.96	0.65	CEL 1			
Clostridium difficile		0.02	0.01				
Clostridium botulinum biovar A		0.02	0.02				
41054100	**Low discrimination**						**41054100**
					GRA	SPO	
Fusobacterium mortiferum		89.11	0.42	XYL 7	1	1	
Clostridium glycolicum		7.39	0.09		99	99	
Actinomyces meyeri		3.49	0.09		99	1	
41055040	**Good identification**						**41055040**
Clostridium glycolicum		100.0	0.49	CEL 1			
41055100	**Good identification**						**41055100**
					GRA		
Fusobacterium mortiferum		96.23	0.45	XYL 7	1		
Actinomyces meyeri		3.77	0.12		99		
41056000	**Low discrimination**						**41056000**
					SPO		
Clostridium difficile		76.24	0.40	CEL 5	99		
Eubacterium limosum		22.49	0.29	XYL 5	1		
Clostridium glycolicum		1.04	0.09		99		
41056010	**Good identification**						**41056010**
					SPO		
Clostridium difficile		99.94	0.51	CEL 5	99		
Eubacterium limosum		0.06	-.06	XYL 5	1		
41060000	**Low discrimination**						**41060000**
					SPO		
Clostridium botulinum biovar A		50.22	0.56	CEL 1	99		
Clostridium sporogenes		49.25	0.53		99		
Atopobium minutum		0.31	0.25		1		
41060002	**Good identification**						**41060002**
					SPO		
Clostridium sporogenes		95.83	0.45	CEL 1	99		
Clostridium botulinum biovar A		2.96	0.21		99		
Atopobium minutum		1.21	0.22		1		
41060040	**Good identification**						**41060040**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		96.49	0.47	CEL 1				
Clostridium sporogenes		3.20	0.19					
Clostridium novyi biovar A		0.31	0.16					
41070000	**Low discrimination**							**41070000**
					GRA	SPO		
Clostridium sporogenes		53.31	0.40	CEL 1	99	99		
Clostridium botulinum biovar A		19.67	0.35		99	99		
Fusobacterium mortiferum		14.33	0.46		1	1		
41070010	**Good identification**							**41070010**
Clostridium difficile		97.84	0.38	TRE 7				
Clostridium sporogenes		1.32	0.06	CEL 5				
Clostridium botulinum biovar A		0.49	0.01	MAN 90				
41070100	**Good identification**							**41070100**
					GRA	SPO		
Fusobacterium mortiferum		97.84	0.52		1	1		
Clostridium sporogenes		1.58	0.06		99	99		
Clostridium botulinum biovar A		0.58	0.01		99	99		
41070102	**Good identification**							**41070102**
					GRA	SPO		
Fusobacterium mortiferum		95.39	0.36	BGL 10	1	1		
Clostridium sporogenes		4.61	-.02		99	99		
41071000	**Acceptable identification**							**41071000**
					GRA	SPO		
Fusobacterium mortiferum		93.20	0.49		1	1		
Atopobium minutum		2.53	0.17		99	1		
Clostridium sporogenes		2.34	0.06		99	99		
41071100	**Good identification**							**41071100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.97	0.55		1	1	1	
Clostridium septicum		0.70	0.01		1	99	99	
Atopobium parvulum		0.33	0.01		99	99	1	
41071102	**Good identification**							**41071102**
Fusobacterium mortiferum		100.0	0.39	BGL 10				
41071120	**Good identification**							**41071120**
					GRA	SPO		
Clostridium septicum		98.58	0.35	MNS 99	99	99		
Fusobacterium mortiferum		1.42	0.21	MLT 94	1	1		
41072000	**Low discrimination**							**41072000**
					SPO			
Clostridium difficile		81.94	0.42	TRE 7	99			
Eubacterium limosum		16.90	0.29	CEL 5	1			
Clostridium sporogenes		0.60	0.06		99			
41072010	**Good identification**							**41072010**
					SPO			
Clostridium difficile		99.96	0.54	TRE 7	99			
Eubacterium limosum		0.04	-.06	CEL 5	1			
41075100	**Acceptable identification**							**41075100**
					COC	GRA		
Fusobacterium mortiferum		92.38	0.36	XYL 7	1	1		
Atopobium parvulum		7.62	0.06		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
41100000	**Low discrimination**					**41100000**
Clostridium sporogenes		58.37	1.00			
Clostridium botulinum biovar A		34.97	0.99			
Clostridium bifermentans		4.19	0.82			
41100001	**Low discrimination**					**41100001**
Clostridium novyi biovar A		59.75	0.87			
Clostridium sporogenes		22.84	0.66			
Clostridium botulinum biovar A		13.68	0.64			
41100002	**Very good identification**					**41100002**
Clostridium sporogenes		96.43	0.92			
Clostridium botulinum biovar A		1.75	0.64			
Clostridium novyi biovar A		1.35	0.74			
41100003	**Low discrimination**					**41100003**
Clostridium novyi biovar A		57.47	0.74			
Clostridium sporogenes		41.49	0.58			
Clostridium botulinum biovar A		0.75	0.30			
41100010	**Low discrimination**					**41100010**
Clostridium novyi biovar A		60.80	0.87			
Clostridium sporogenes		23.23	0.66			
Clostridium botulinum biovar A		13.92	0.64			
41100011	**Very good identification**					**41100011**
Clostridium novyi biovar A		99.33	0.87			
Clostridium sporogenes		0.38	0.32			
Clostridium botulinum biovar A		0.23	0.30			
41100012	**Low discrimination**					**41100012**
Clostridium novyi biovar A		57.52	0.74			
Clostridium sporogenes		41.53	0.58			
Clostridium botulinum biovar A		0.75	0.30			
41100013	**Very good identification**					**41100013**
Clostridium novyi biovar A		99.28	0.74			
Clostridium sporogenes		0.72	0.24			
41100020	**Low discrimination**					**41100020**
Clostridium sporogenes		36.22	0.66	NAG 1		
Clostridium novyi biovar A		23.69	0.77			
Clostridium botulinum biovar A		21.70	0.64			
41100021	**Very good identification**					**41100021**
Clostridium novyi biovar A		96.75	0.77			
Clostridium sporogenes		1.49	0.32			
Clostridium botulinum biovar A		0.90	0.30			
41100022	**Low discrimination**					**41100022**
					SPO	
Clostridium sporogenes		66.40	0.58	NAG 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		23.00	0.64		99	
Atopobium minutum		4.94	0.48		1	
41100023	**Very good identification**					**41100023**
					SPO	
Clostridium novyi biovar A		96.82	0.64		99	
Clostridium sporogenes		2.82	0.24		99	
Atopobium minutum		0.21	0.14		1	
41100030	**Very good identification**					**41100030**
Clostridium novyi biovar A		96.86	0.77			
Clostridium sporogenes		1.50	0.32			
Clostridium botulinum biovar A		0.90	0.30			
41100031	**Very good identification**					**41100031**
Clostridium novyi biovar A		99.98	0.77			
Clostridium sporogenes		0.02	-.03			
41100032	**Very good identification**					**41100032**
					SPO	
Clostridium novyi biovar A		96.82	0.64		99	
Clostridium sporogenes		2.82	0.24		99	
Atopobium minutum		0.21	0.14		1	
41100033	**Very good identification**					**41100033**
Clostridium novyi biovar A		100.0	0.64			
41100040	**Low discrimination**					**41100040**
Clostridium botulinum biovar A		83.28	0.90			
Clostridium novyi biovar A		8.20	0.84			
Clostridium sporogenes		4.70	0.66			
41100041	**Low discrimination**					**41100041**
Clostridium novyi biovar A		89.70	0.84			
Clostridium botulinum biovar A		9.20	0.55			
Clostridium sporogenes		0.52	0.32			
41100042	**Low discrimination**					**41100042**
Clostridium sporogenes		40.16	0.58	SOR 1		
Clostridium novyi biovar A		37.09	0.71			
Clostridium botulinum biovar A		21.56	0.55			
41100043	**Very good identification**					**41100043**
Clostridium novyi biovar A		98.35	0.71			
Clostridium sporogenes		1.08	0.24			
Clostridium botulinum biovar A		0.58	0.21			
41100050	**Low discrimination**					**41100050**
Clostridium novyi biovar A		89.84	0.84			
Clostridium botulinum biovar A		9.22	0.55			
Clostridium sporogenes		0.52	0.32			
41100051	**Very good identification**					**41100051**
Clostridium novyi biovar A		99.89	0.84			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium botulinum biovar A		0.10	0.21			
Clostridium sporogenes		0.01	-.03			
41100052	**Very good identification**					**41100052**
Clostridium novyi biovar A		98.35	0.71			
Clostridium sporogenes		1.08	0.24			
Clostridium botulinum biovar A		0.58	0.21			
41100053	**Very good identification**					**41100053**
Clostridium novyi biovar A		100.0	0.71			
41100060	**Low discrimination**					**41100060**
Clostridium novyi biovar A		67.14	0.74			
Clostridium botulinum biovar A		27.55	0.55			
Clostridium bifermentans		2.91	0.38			
41100061	**Very good identification**					**41100061**
Clostridium novyi biovar A		99.52	0.74			
Clostridium botulinum biovar A		0.41	0.21			
Clostridium bifermentans		0.04	0.04			
41100062	**Good identification**					**41100062**
Clostridium novyi biovar A		93.00	0.61			
Clostridium sporogenes		4.07	0.24			
Clostridium botulinum biovar A		2.18	0.21			
41100063	**Very good identification**					**41100063**
Clostridium novyi biovar A		100.0	0.61			
41100070	**Very good identification**					**41100070**
Clostridium novyi biovar A		99.52	0.74			
Clostridium botulinum biovar A		0.41	0.21			
Clostridium bifermentans		0.04	0.04			
41100071	**Very good identification**					**41100071**
Clostridium novyi biovar A		100.0	0.74			
41100072	**Very good identification**					**41100072**
Clostridium novyi biovar A		100.0	0.61			
41100073	**Very good identification**					**41100073**
Clostridium novyi biovar A		100.0	0.61			
41100100	**Low discrimination**					**41100100**
Clostridium sporogenes		57.89	0.66	LAC 1		
Clostridium botulinum biovar A		34.68	0.64			
Clostridium bifermentans		4.16	0.48			
41100101	**Low discrimination**					**41100101**
Clostridium novyi biovar A		59.42	0.53	LAC 1		
Clostridium sporogenes		22.71	0.32			
Clostridium botulinum biovar A		13.60	0.30			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41100102	**Very good identification**						**41100102**
Clostridium sporogenes		96.04	0.58	LAC 1			
Clostridium botulinum biovar A		1.74	0.30				
Clostridium novyi biovar A		1.34	0.40				
41100103	**Low discrimination**						**41100103**
Clostridium novyi biovar A		58.08	0.40	LAC 1			
Clostridium sporogenes		41.92	0.24				
41100110	**Low discrimination**						**41100110**
Clostridium novyi biovar A		60.71	0.53	LAC 1			
Clostridium sporogenes		23.20	0.32				
Clostridium botulinum biovar A		13.90	0.30				
41100111	**Good identification**						**41100111**
Clostridium novyi biovar A		99.62	0.53	LAC 1			
Clostridium sporogenes		0.39	-.03				
41100112	**Low discrimination**						**41100112**
Clostridium novyi biovar A		58.08	0.40	LAC 1			
Clostridium sporogenes		41.92	0.24				
41100113	**Good identification**						**41100113**
Clostridium novyi biovar A		100.0	0.40	LAC 1			
41100121	**Acceptable identification**						**41100121**
					GRA	SPO	
Clostridium novyi biovar A		93.81	0.43	LAC 1	99	99	
Prevotella bivia		4.74	-.03		1	1	
Clostridium sporogenes		1.45	-.03		99	99	
41100130	**Acceptable identification**						**41100130**
					GRA	SPO	
Clostridium novyi biovar A		93.81	0.43	LAC 1	99	99	
Prevotella bivia		4.74	-.03		1	1	
Clostridium sporogenes		1.45	-.03		99	99	
41100131	**Good identification**						**41100131**
Clostridium novyi biovar A		100.0	0.43	LAC 1			
41100140	**Low discrimination**						**41100140**
Clostridium botulinum biovar A		83.26	0.55	LAC 1			
Clostridium novyi biovar A		8.20	0.50				
Clostridium sporogenes		4.70	0.32				
41100141	**Acceptable identification**						**41100141**
Clostridium novyi biovar A		90.22	0.50	LAC 1			
Clostridium botulinum biovar A		9.26	0.21				
Clostridium sporogenes		0.52	-.03				
41100143	**Good identification**						**41100143**
Clostridium novyi biovar A		100.0	0.37	LAC 1			
41100150	**Acceptable identification**						**41100150**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		90.22	0.50	LAC 1			
Clostridium botulinum biovar A		9.26	0.21				
Clostridium sporogenes		0.52	-.03				
41100151	**Good identification**						**41100151**
Clostridium novyi biovar A		100.0	0.50	LAC 1			
41100152	**Good identification**						**41100152**
Clostridium novyi biovar A		100.0	0.37	LAC 1			
41100153	**Good identification**						**41100153**
Clostridium novyi biovar A		100.0	0.37	LAC 1			
41100160	**Low discrimination**						**41100160**
					GRA	SPO	
Clostridium novyi biovar A		64.41	0.40	LAC 1	99	99	
Clostridium botulinum biovar A		26.43	0.21		99	99	
Prevotella bivia		4.88	-.03		1	1	
41100161	**Good identification**						**41100161**
Clostridium novyi biovar A		100.0	0.40	LAC 1			
41100170	**Good identification**						**41100170**
Clostridium novyi biovar A		100.0	0.40	LAC 1			
41100171	**Good identification**						**41100171**
Clostridium novyi biovar A		100.0	0.40	LAC 1			
41100200	**Low discrimination**						**41100200**
Clostridium sporogenes		58.37	0.66	RHA 1			
Clostridium botulinum biovar A		34.97	0.64				
Clostridium bifermentans		4.19	0.48				
41100201	**Low discrimination**						**41100201**
Clostridium novyi biovar A		59.85	0.53	RHA 1			
Clostridium sporogenes		22.87	0.32				
Clostridium botulinum biovar A		13.70	0.30				
41100202	**Very good identification**						**41100202**
Clostridium sporogenes		96.45	0.58	RHA 1			
Clostridium botulinum biovar A		1.75	0.30				
Clostridium novyi biovar A		1.35	0.40				
41100203	**Low discrimination**						**41100203**
Clostridium novyi biovar A		58.08	0.40	RHA 1			
Clostridium sporogenes		41.92	0.24				
41100210	**Low discrimination**						**41100210**
Clostridium novyi biovar A		60.90	0.53	RHA 1			
Clostridium sporogenes		23.27	0.32				
Clostridium botulinum biovar A		13.94	0.30				
41100211	**Good identification**						**41100211**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		99.62	0.53	RHA 1		
Clostridium sporogenes		0.39	-.03			
41100212	**Low discrimination**					**41100212**
Clostridium novyi biovar A		58.08	0.40	RHA 1		
Clostridium sporogenes		41.92	0.24			
41100213	**Good identification**					**41100213**
Clostridium novyi biovar A		100.0	0.40	RHA 1		
41100221	**Good identification**					**41100221**
Clostridium novyi biovar A		98.48	0.43	RHA 1		
Clostridium sporogenes		1.52	-.03			
41100230	**Good identification**					**41100230**
Clostridium novyi biovar A		98.48	0.43	RHA 1		
Clostridium sporogenes		1.52	-.03			
41100231	**Good identification**					**41100231**
Clostridium novyi biovar A		100.0	0.43	RHA 1		
41100240	**Low discrimination**					**41100240**
Clostridium botulinum biovar A		83.31	0.55	RHA 1		
Clostridium novyi biovar A		8.20	0.50			
Clostridium sporogenes		4.70	0.32			
41100241	**Acceptable identification**					**41100241**
Clostridium novyi biovar A		90.22	0.50	RHA 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41100243	**Good identification**					**41100243**
Clostridium novyi biovar A		100.0	0.37	RHA 1		
41100250	**Acceptable identification**					**41100250**
Clostridium novyi biovar A		90.22	0.50	RHA 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41100251	**Good identification**					**41100251**
Clostridium novyi biovar A		100.0	0.50	RHA 1		
41100252	**Good identification**					**41100252**
Clostridium novyi biovar A		100.0	0.37	RHA 1		
41100253	**Good identification**					**41100253**
Clostridium novyi biovar A		100.0	0.37	RHA 1		
41100260	**Low discrimination**					**41100260**
Clostridium novyi biovar A		67.71	0.40	RHA 1		
Clostridium botulinum biovar A		27.78	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		2.94	0.04			
41100261	**Good identification**					**41100261**
Clostridium novyi biovar A		100.0	0.40	RHA 1		
41100270	**Good identification**					**41100270**
Clostridium novyi biovar A		100.0	0.40	RHA 1		
41100271	**Good identification**					**41100271**
Clostridium novyi biovar A		100.0	0.40	RHA 1		
41100400	**Low discrimination**					**41100400**
Clostridium sporogenes		58.36	0.66	ARA 1		
Clostridium botulinum biovar A		34.96	0.64			
Clostridium bifermentans		4.19	0.48			
41100401	**Low discrimination**					**41100401**
Clostridium novyi biovar A		59.85	0.53	ARA 1		
Clostridium sporogenes		22.87	0.32			
Clostridium botulinum biovar A		13.70	0.30			
41100402	**Very good identification**					**41100402**
Clostridium sporogenes		96.45	0.58	ARA 1		
Clostridium botulinum biovar A		1.75	0.30			
Clostridium novyi biovar A		1.35	0.40			
41100403	**Low discrimination**					**41100403**
Clostridium novyi biovar A		58.08	0.40	ARA 1		
Clostridium sporogenes		41.92	0.24			
41100410	**Low discrimination**					**41100410**
Clostridium novyi biovar A		60.90	0.53	ARA 1		
Clostridium sporogenes		23.27	0.32			
Clostridium botulinum biovar A		13.94	0.30			
41100411	**Good identification**					**41100411**
Clostridium novyi biovar A		99.62	0.53	ARA 1		
Clostridium sporogenes		0.39	-.03			
41100412	**Low discrimination**					**41100412**
Clostridium novyi biovar A		58.08	0.40	ARA 1		
Clostridium sporogenes		41.92	0.24			
41100413	**Good identification**					**41100413**
Clostridium novyi biovar A		100.0	0.40	ARA 1		
41100421	**Good identification**					**41100421**
Clostridium novyi biovar A		98.48	0.43	ARA 1		
Clostridium sporogenes		1.52	-.03			
41100430	**Good identification**					**41100430**
Clostridium novyi biovar A		98.48	0.43	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sporogenes		1.52	-.03			
41100431	**Good identification**					**41100431**
Clostridium novyi biovar A		100.0	0.43	ARA 1		
41100440	**Low discrimination**					**41100440**
Clostridium botulinum biovar A		83.31	0.55	ARA 1		
Clostridium novyi biovar A		8.20	0.50			
Clostridium sporogenes		4.70	0.32			
41100441	**Acceptable identification**					**41100441**
Clostridium novyi biovar A		90.22	0.50	ARA 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41100443	**Good identification**					**41100443**
Clostridium novyi biovar A		100.0	0.37	ARA 1		
41100450	**Acceptable identification**					**41100450**
Clostridium novyi biovar A		90.22	0.50	ARA 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41100451	**Good identification**					**41100451**
Clostridium novyi biovar A		100.0	0.50	ARA 1		
41100452	**Good identification**					**41100452**
Clostridium novyi biovar A		100.0	0.37	ARA 1		
41100453	**Good identification**					**41100453**
Clostridium novyi biovar A		100.0	0.37	ARA 1		
41100460	**Low discrimination**					**41100460**
Clostridium novyi biovar A		67.71	0.40	ARA 1		
Clostridium botulinum biovar A		27.78	0.21			
Clostridium bifermentans		2.94	0.04			
41100461	**Good identification**					**41100461**
Clostridium novyi biovar A		100.0	0.40	ARA 1		
41100470	**Good identification**					**41100470**
Clostridium novyi biovar A		100.0	0.40	ARA 1		
41100471	**Good identification**					**41100471**
Clostridium novyi biovar A		100.0	0.40	ARA 1		
41101000	**Low discrimination**					**41101000**
Clostridium sporogenes		49.20	0.66	GAL 1		
Clostridium botulinum biovar A		29.48	0.64			
Clostridium novyi biovar A		14.31	0.71			
41101001	**Good identification**					**41101001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		90.38	0.71	GAL 10		
Clostridium sordellii		4.13	0.31			
Clostridium sporogenes		3.14	0.32			
41101002	**Low discrimination**					**41101002**
					SPO	
Clostridium sporogenes		79.79	0.58	GAL 1	99	
Clostridium novyi biovar A		12.28	0.58		99	
Atopobium minutum		5.94	0.48		1	
41101003	**Good identification**					**41101003**
					SPO	
Clostridium novyi biovar A		93.19	0.58	GAL 10	99	
Clostridium sporogenes		6.12	0.24		99	
Atopobium minutum		0.46	0.14		1	
41101010	**Good identification**					**41101010**
Clostridium novyi biovar A		94.27	0.71	GAL 10		
Clostridium sporogenes		3.28	0.32			
Clostridium botulinum biovar A		1.96	0.30			
41101011	**Very good identification**					**41101011**
Clostridium novyi biovar A		99.92	0.71	GAL 10		
Clostridium sordellii		0.05	-.03			
Clostridium sporogenes		0.04	-.03			
41101012	**Good identification**					**41101012**
					SPO	
Clostridium novyi biovar A		93.41	0.58	GAL 10	99	
Clostridium sporogenes		6.13	0.24		99	
Atopobium minutum		0.46	0.14		1	
41101013	**Very good identification**					**41101013**
Clostridium novyi biovar A		100.0	0.58	GAL 10		
41101020	**Low discrimination**					**41101020**
					SPO	
Clostridium novyi biovar A		67.54	0.61	GAL 10	99	
Atopobium minutum		11.53	0.43		1	
Clostridium sporogenes		9.39	0.32		99	
41101021	**Very good identification**					**41101021**
					SPO	
Clostridium novyi biovar A		99.51	0.61	GAL 10	99	
Clostridium sordellii		0.18	-.03		99	
Atopobium minutum		0.17	0.09		1	
41101022	**Low discrimination**					**41101022**
					SPO	
Clostridium novyi biovar A		48.24	0.48	GAL 10	99	
Atopobium minutum		31.11	0.40		1	
Clostridium sporogenes		12.66	0.24		99	
41101023	**Good identification**					**41101023**
					SPO	
Clostridium novyi biovar A		99.35	0.48	GAL 10	99	
Atopobium minutum		0.65	0.06		1	
41101030	**Very good identification**					**41101030**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		99.69	0.61	GAL 10	99	
Atopobium minutum		0.17	0.09		1	
Clostridium sporogenes		0.14	-.03		99	
41101031	**Very good identification**					**41101031**
Clostridium novyi biovar A		100.0	0.61	GAL 10		
41101032	**Good identification**					**41101032**
					SPO	
Clostridium novyi biovar A		99.35	0.48	GAL 10	99	
Atopobium minutum		0.65	0.06		1	
41101033	**Good identification**					**41101033**
Clostridium novyi biovar A		100.0	0.48	GAL 10		
41101040	**Low discrimination**					**41101040**
Clostridium novyi biovar A		43.95	0.68	GAL 10		
Clostridium botulinum biovar A		40.58	0.55			
Clostridium glycolicum		12.35	0.45			
41101041	**Very good identification**					**41101041**
Clostridium novyi biovar A		98.47	0.68	GAL 10		
Clostridium botulinum biovar A		0.92	0.21			
Clostridium glycolicum		0.28	0.11			
41101042	**Low discrimination**					**41101042**
Clostridium novyi biovar A		85.13	0.55	GAL 10		
Clostridium sporogenes		8.38	0.24			
Clostridium botulinum biovar A		4.50	0.21			
41101043	**Good identification**					**41101043**
Clostridium novyi biovar A		100.0	0.55	GAL 10		
41101050	**Very good identification**					**41101050**
Clostridium novyi biovar A		98.75	0.68	GAL 10		
Clostridium botulinum biovar A		0.92	0.21			
Clostridium glycolicum		0.28	0.11			
41101051	**Very good identification**					**41101051**
Clostridium novyi biovar A		100.0	0.68	GAL 10		
41101052	**Good identification**					**41101052**
Clostridium novyi biovar A		100.0	0.55	GAL 10		
41101053	**Good identification**					**41101053**
Clostridium novyi biovar A		100.0	0.55	GAL 10		
41101060	**Good identification**					**41101060**
Clostridium novyi biovar A		94.58	0.58	GAL 10		
Clostridium botulinum biovar A		3.53	0.21			
Clostridium glycolicum		1.07	0.11			
41101061	**Very good identification**					**41101061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.58	GAL 10			
41101062	**Good identification**						**41101062**
					SPO		
Clostridium novyi biovar A		99.03	0.45	GAL 10	99		
Atopobium minutum		0.97	0.06		1		
41101063	**Good identification**						**41101063**
Clostridium novyi biovar A		100.0	0.45	GAL 10			
41101070	**Very good identification**						**41101070**
Clostridium novyi biovar A		100.0	0.58	GAL 10			
41101071	**Very good identification**						**41101071**
Clostridium novyi biovar A		100.0	0.58	GAL 10			
41101072	**Good identification**						**41101072**
Clostridium novyi biovar A		100.0	0.45	GAL 10			
41101073	**Good identification**						**41101073**
Clostridium novyi biovar A		100.0	0.45	GAL 10			
41101100	**Very good identification**						**41101100**
					COC	SPO	
Blautia hansenii		98.30	0.66	RAF 99	99	1	
Clostridium sporogenes		0.77	0.32		1	99	
Clostridium botulinum biovar A		0.46	0.30		1	99	
41101101	**Low discrimination**						**41101101**
					COC	SPO	
Clostridium chauvoei		53.42	0.40	SUC 99	1	99	
Blautia hansenii		37.46	0.32	MNS 90	99	1	
Clostridium novyi biovar A		8.44	0.37		1	99	
41101102	**Good identification**						**41101102**
					COC	SPO	
Blautia hansenii		97.07	0.50	RAF 99	99	1	
Clostridium sporogenes		2.28	0.24	BGL 10	1	99	
Clostridium novyi biovar A		0.35	0.24		1	99	
41101111	**Acceptable identification**						**41101111**
					COC	SPO	
Clostridium novyi biovar A		90.19	0.37	GAL 10	1	99	
Clostridium chauvoei		5.77	0.06	LAC 1	1	99	
Blautia hansenii		4.04	-.03		99	1	
41101120	**Acceptable identification**						**41101120**
					COC	GRA	
Blautia hansenii		94.35	0.50	RAF 99	99	99	
Prevotella bivia		3.84	0.19	NAG 10	1	1	
Prevotella melaninogenica		0.81	0.14		1	1	
41101201	**Acceptable identification**						**41101201**
Clostridium novyi biovar A		92.55	0.37	GAL 10			
Clostridium sordellii		4.23	-.03	RHA 1			
Clostridium sporogenes		3.22	-.03				
41101210	**Good identification**						**41101210**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		96.64	0.37	GAL 10			
Clostridium sporogenes		3.36	-.03	RHA 1			
41101211	**Good identification**						**41101211**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
41101401	**Acceptable identification**						**41101401**
Clostridium novyi biovar A		92.55	0.37	GAL 10			
Clostridium sordellii		4.23	-.03	ARA 1			
Clostridium sporogenes		3.22	-.03				
41101410	**Good identification**						**41101410**
Clostridium novyi biovar A		96.64	0.37	GAL 10			
Clostridium sporogenes		3.36	-.03	ARA 1			
41101411	**Good identification**						**41101411**
Clostridium novyi biovar A		100.0	0.37	GAL 10			
41101500	**Good identification**						**41101500**
					COC		
Blautia hansenii		99.82	0.53	RAF 99	99		
Actinomyces meyeri		0.10	0.10		1		
Lactobacillus fermentum		0.04	0.14		1		
41101502	**Good identification**						**41101502**
					COC		
Blautia hansenii		99.63	0.37	RAF 99	99		
Lactobacillus fermentum		0.37	0.14	BGL 10	1		
41101520	**Good identification**						**41101520**
					COC	GRA	
Blautia hansenii		99.56	0.37	RAF 99	99	99	
Odoribacter splanchnicus		0.44	-.07	NAG 10	1	1	
41102000	**Low discrimination**						**41102000**
Clostridium sporogenes		57.55	0.66	MAN 1			
Clostridium botulinum biovar A		34.48	0.64				
Clostridium bifermentans		4.13	0.48				
41102001	**Low discrimination**						**41102001**
Clostridium novyi biovar A		59.53	0.53	MAN 1			
Clostridium sporogenes		22.75	0.32				
Clostridium botulinum biovar A		13.63	0.30				
41102002	**Very good identification**						**41102002**
Clostridium sporogenes		95.98	0.58	MAN 1			
Clostridium botulinum biovar A		1.74	0.30				
Clostridium novyi biovar A		1.34	0.40				
41102003	**Low discrimination**						**41102003**
Clostridium novyi biovar A		58.08	0.40	MAN 1			
Clostridium sporogenes		41.92	0.24				
41102010	**Low discrimination**						**41102010**
Clostridium novyi biovar A		53.80	0.53	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sporogenes		20.56	0.32			
Clostridium botulinum biovar A		12.32	0.30			
41102011	**Good identification**					**41102011**
Clostridium novyi biovar A		99.62	0.53	MAN 1		
Clostridium sporogenes		0.39	-.03			
41102012	**Low discrimination**					**41102012**
Clostridium novyi biovar A		58.08	0.40	MAN 1		
Clostridium sporogenes		41.92	0.24			
41102013	**Good identification**					**41102013**
Clostridium novyi biovar A		100.0	0.40	MAN 1		
41102021	**Good identification**					**41102021**
Clostridium novyi biovar A		98.48	0.43	MAN 1		
Clostridium sporogenes		1.52	-.03			
41102030	**Good identification**					**41102030**
Clostridium novyi biovar A		98.48	0.43	MAN 1		
Clostridium sporogenes		1.52	-.03			
41102031	**Good identification**					**41102031**
Clostridium novyi biovar A		100.0	0.43	MAN 1		
41102040	**Low discrimination**					**41102040**
Clostridium botulinum biovar A		79.74	0.55	MAN 1		
Clostridium novyi biovar A		7.85	0.50			
Clostridium sporogenes		4.50	0.32			
41102041	**Low discrimination**					**41102041**
Clostridium novyi biovar A		89.78	0.50	MAN 1		
Clostridium botulinum biovar A		9.21	0.21			
Clostridium sporogenes		0.52	-.03			
41102043	**Good identification**					**41102043**
Clostridium novyi biovar A		100.0	0.37	MAN 1		
41102050	**Low discrimination**					**41102050**
Clostridium novyi biovar A		88.47	0.50	MAN 1		
Clostridium botulinum biovar A		9.08	0.21			
Clostridium difficile		1.45	0.05			
41102051	**Good identification**					**41102051**
Clostridium novyi biovar A		100.0	0.50	MAN 1		
41102052	**Good identification**					**41102052**
Clostridium novyi biovar A		100.0	0.37	MAN 1		
41102053	**Good identification**					**41102053**
Clostridium novyi biovar A		100.0	0.37	MAN 1		
41102060	**Low discrimination**					**41102060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		66.71	0.40	MAN 1		
Clostridium botulinum biovar A		27.38	0.21			
Clostridium bifermentans		2.89	0.04			
41102061	**Good identification**					**41102061**
Clostridium novyi biovar A		100.0	0.40	MAN 1		
41102070	**Good identification**					**41102070**
Clostridium novyi biovar A		100.0	0.40	MAN 1		
41102071	**Good identification**					**41102071**
Clostridium novyi biovar A		100.0	0.40	MAN 1		
41103001	**Acceptable identification**					**41103001**
Clostridium novyi biovar A		92.55	0.37	GAL 10		
Clostridium sordellii		4.23	-.03	MAN 1		
Clostridium sporogenes		3.22	-.03			
41103010	**Good identification**					**41103010**
Clostridium novyi biovar A		96.64	0.37	GAL 10		
Clostridium sporogenes		3.36	-.03	MAN 1		
41103011	**Good identification**					**41103011**
Clostridium novyi biovar A		100.0	0.37	GAL 10		
41104000	**Low discrimination**					**41104000**
Clostridium sporogenes		45.38	0.66	XYL 1		
Clostridium botulinum biovar A		27.19	0.64			
Clostridium glycolicum		22.24	0.62			
41104001	**Low discrimination**					**41104001**
Clostridium novyi biovar A		53.82	0.53	XYL 1		
Clostridium sporogenes		20.57	0.32			
Clostridium botulinum biovar A		12.32	0.30			
41104002	**Very good identification**					**41104002**
Clostridium sporogenes		95.08	0.58	XYL 1		
Clostridium botulinum biovar A		1.73	0.30			
Clostridium glycolicum		1.41	0.28			
41104003	**Low discrimination**					**41104003**
Clostridium novyi biovar A		58.08	0.40	XYL 1		
Clostridium sporogenes		41.92	0.24			
41104010	**Low discrimination**					**41104010**
Clostridium novyi biovar A		54.66	0.53	XYL 1		
Clostridium sporogenes		20.89	0.32			
Clostridium botulinum biovar A		12.52	0.30			
41104011	**Good identification**					**41104011**
Clostridium novyi biovar A		99.62	0.53	XYL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sporogenes		0.39	-.03			
41104012	**Low discrimination**					**41104012**
Clostridium novyi biovar A		58.08	0.40	XYL 1		
Clostridium sporogenes		41.92	0.24			
41104013	**Good identification**					**41104013**
Clostridium novyi biovar A		100.0	0.40	XYL 1		
41104021	**Good identification**					**41104021**
Clostridium novyi biovar A		98.48	0.43	XYL 1		
Clostridium sporogenes		1.52	-.03			
41104030	**Good identification**					**41104030**
Clostridium novyi biovar A		98.48	0.43	XYL 1		
Clostridium sporogenes		1.52	-.03			
41104031	**Good identification**					**41104031**
Clostridium novyi biovar A		100.0	0.43	XYL 1		
41104040	**Very good identification**					**41104040**
Clostridium glycolicum		97.80	0.84	FRU 90		
Clostridium botulinum biovar A		1.88	0.55			
Clostridium novyi biovar A		0.19	0.50			
41104041	**Low discrimination**					**41104041**
Clostridium glycolicum		82.81	0.50	FRU 90		
Clostridium novyi biovar A		15.51	0.50	URE 1		
Clostridium botulinum biovar A		1.59	0.21			
41104042	**Acceptable identification**					**41104042**
Clostridium glycolicum		91.91	0.50	FRU 90		
Clostridium sporogenes		3.29	0.24	BGL 1		
Clostridium novyi biovar A		3.04	0.37			
41104043	**Low discrimination**					**41104043**
Clostridium novyi biovar A		76.60	0.37	XYL 1		
Clostridium glycolicum		23.40	0.15			
41104050	**Low discrimination**					**41104050**
Clostridium glycolicum		82.81	0.50	FRU 90		
Clostridium novyi biovar A		15.51	0.50	MLZ 1		
Clostridium botulinum biovar A		1.59	0.21			
41104051	**Acceptable identification**					**41104051**
Clostridium novyi biovar A		94.88	0.50	XYL 1		
Clostridium glycolicum		5.12	0.15			
41104052	**Low discrimination**					**41104052**
Clostridium novyi biovar A		76.60	0.37	XYL 1		
Clostridium glycolicum		23.40	0.15			
41104053	**Good identification**					**41104053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.37	XYL 1		
41104060	**Acceptable identification**					**41104060**
Clostridium glycolicum		93.53	0.50	FRU 90		
Clostridium novyi biovar A		4.38	0.40	NAG 1		
Clostridium botulinum biovar A		1.80	0.21			
41104061	**Low discrimination**					**41104061**
Clostridium novyi biovar A		82.26	0.40	XYL 1		
Clostridium glycolicum		17.74	0.15			
41104070	**Low discrimination**					**41104070**
Clostridium novyi biovar A		82.26	0.40	XYL 1		
Clostridium glycolicum		17.74	0.15			
41104071	**Good identification**					**41104071**
Clostridium novyi biovar A		100.0	0.40	XYL 1		
41104140	**Good identification**					**41104140**
Clostridium glycolicum		97.83	0.50	FRU 90		
Clostridium botulinum biovar A		1.88	0.21	LAC 1		
Clostridium novyi biovar A		0.19	0.16			
41104240	**Good identification**					**41104240**
Clostridium glycolicum		97.83	0.50	FRU 90		
Clostridium botulinum biovar A		1.88	0.21	RHA 1		
Clostridium novyi biovar A		0.19	0.16			
41104440	**Good identification**					**41104440**
Clostridium glycolicum		97.83	0.50	FRU 90		
Clostridium botulinum biovar A		1.88	0.21	ARA 1		
Clostridium novyi biovar A		0.19	0.16			
41105000	**Low discrimination**					**41105000**
Clostridium glycolicum		69.96	0.45	FRU 90		
Clostridium sporogenes		12.98	0.32	GAL 10		
Clostridium botulinum biovar A		7.77	0.30	SOR 95		
41105001	**Low discrimination**					**41105001**
Clostridium novyi biovar A		78.88	0.37	GAL 10		
Clostridium glycolicum		14.77	0.11	XYL 1		
Clostridium sordellii		3.61	-.03			
41105010	**Low discrimination**					**41105010**
Clostridium novyi biovar A		79.62	0.37	GAL 10		
Clostridium glycolicum		14.91	0.11	XYL 1		
Clostridium sporogenes		2.77	-.03			
41105011	**Good identification**					**41105011**
Clostridium novyi biovar A		100.0	0.37	GAL 10		
41105040	**Very good identification**					**41105040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium glycolicum		99.63	0.67	FRU 90			
Clostridium novyi biovar A		0.19	0.34	GAL 10			
Clostridium botulinum biovar A		0.17	0.21				
41105100	**Good identification**						**41105100**
					COC	SPO	
Blautia hansenii		98.84	0.53	RAF 99	99	1	
Actinomyces meyeri		0.75	0.25		1	1	
Clostridium glycolicum		0.24	0.11		1	99	
41105102	**Good identification**						**41105102**
Blautia hansenii		100.0	0.37	RAF 99			
41105120	**Good identification**						**41105120**
					COC		
Blautia hansenii		99.08	0.37	RAF 99	99		
Actinomyces meyeri		0.92	0.10	NAG 10	1		
41105500	**Good identification**						**41105500**
					COC		
Blautia hansenii		98.94	0.40	RAF 99	99		
Actinomyces meyeri		1.07	0.14		1		
41106040	**Good identification**						**41106040**
Clostridium glycolicum		97.73	0.50	FRU 90			
Clostridium botulinum biovar A		1.88	0.21	MAN 1			
Clostridium novyi biovar A		0.19	0.16				
41110000	**Low discrimination**						**41110000**
					GRA	SPO	
Clostridium sporogenes		75.13	0.87		99	99	
Clostridium botulinum biovar A		16.28	0.78		99	99	
Prevotella disiens		2.16	0.57		1	1	
41110001	**Low discrimination**						**41110001**
Clostridium novyi biovar A		48.63	0.71	FRU 10			
Clostridium sporogenes		29.52	0.53				
Clostridium sordellii		12.60	0.44				
41110002	**Very good identification**						**41110002**
					SPO		
Clostridium sporogenes		96.60	0.79		99		
Atopobium minutum		1.14	0.56		1		
Clostridium novyi biovar A		0.85	0.58		99		
41110003	**Low discrimination**						**41110003**
Clostridium sporogenes		52.39	0.45	URE 1			
Clostridium novyi biovar A		45.70	0.58				
Clostridium sordellii		0.68	0.10				
41110010	**Low discrimination**						**41110010**
Clostridium novyi biovar A		49.88	0.71	FRU 10			
Clostridium sporogenes		30.28	0.53				
Clostridium difficile		10.37	0.45				
41110011	**Very good identification**						**41110011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		98.80	0.71	FRU 10		
Clostridium sporogenes		0.61	0.19			
Clostridium sordellii		0.26	0.10			
41110012	**Low discrimination**					**41110012**
					SPO	
Clostridium sporogenes		52.46	0.45	MLZ 1	99	
Clostridium novyi biovar A		45.76	0.58		99	
Atopobium minutum		0.62	0.22		1	
41110013	**Very good identification**					**41110013**
Clostridium novyi biovar A		98.86	0.58	FRU 10		
Clostridium sporogenes		1.15	0.11			
41110020	**Low discrimination**					**41110020**
Clostridium sporogenes		48.62	0.53	NAG 1		
Clostridium novyi biovar A		20.03	0.61			
Clostridium botulinum biovar A		10.54	0.44			
41110021	**Very good identification**					**41110021**
Clostridium novyi biovar A		95.38	0.61	FRU 10		
Clostridium sporogenes		2.34	0.19			
Clostridium sordellii		1.00	0.10			
41110022	**Low discrimination**					**41110022**
					SPO	
Clostridium sporogenes		60.15	0.45	NAG 1	99	
Atopobium minutum		23.42	0.48		1	
Clostridium novyi biovar A		13.12	0.48		99	
41110023	**Acceptable identification**					**41110023**
					SPO	
Clostridium novyi biovar A		93.95	0.48	FRU 10	99	
Clostridium sporogenes		4.35	0.11		99	
Atopobium minutum		1.70	0.14		1	
41110030	**Very good identification**					**41110030**
Clostridium novyi biovar A		95.57	0.61	FRU 10		
Clostridium sporogenes		2.34	0.19			
Clostridium difficile		0.80	0.10			
41110031	**Very good identification**					**41110031**
Clostridium novyi biovar A		100.0	0.61	FRU 10		
41110032	**Acceptable identification**					**41110032**
					SPO	
Clostridium novyi biovar A		93.95	0.48	FRU 10	99	
Clostridium sporogenes		4.35	0.11		99	
Atopobium minutum		1.70	0.14		1	
41110033	**Good identification**					**41110033**
Clostridium novyi biovar A		100.0	0.48	FRU 10		
41110040	**Low discrimination**					**41110040**
Clostridium glycolicum		72.03	0.78	XYL 95		
Clostridium botulinum biovar A		18.47	0.69			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		3.17	0.68			
41110041	**Low discrimination**					**41110041**
Clostridium novyi biovar A		75.50	0.68	FRU 10		
Clostridium glycolicum		17.35	0.44			
Clostridium botulinum biovar A		4.45	0.35			
41110042	**Low discrimination**					**41110042**
Clostridium sporogenes		34.47	0.45	SOR 1		
Clostridium glycolicum		26.11	0.44			
Clostridium novyi biovar A		20.04	0.55			
41110043	**Good identification**					**41110043**
Clostridium novyi biovar A		96.16	0.55	FRU 10		
Clostridium sporogenes		1.67	0.11			
Clostridium glycolicum		1.27	0.10			
41110050	**Low discrimination**					**41110050**
Clostridium novyi biovar A		75.43	0.68	FRU 10		
Clostridium glycolicum		17.34	0.44			
Clostridium botulinum biovar A		4.45	0.35			
41110051	**Very good identification**					**41110051**
Clostridium novyi biovar A		99.71	0.68	FRU 10		
Clostridium glycolicum		0.23	0.10			
Clostridium botulinum biovar A		0.06	0.01			
41110052	**Good identification**					**41110052**
Clostridium novyi biovar A		96.16	0.55	FRU 10		
Clostridium sporogenes		1.67	0.11			
Clostridium glycolicum		1.27	0.10			
41110053	**Good identification**					**41110053**
Clostridium novyi biovar A		100.0	0.55	FRU 10		
41110060	**Low discrimination**					**41110060**
Clostridium novyi biovar A		43.90	0.58	FRU 10		
Clostridium glycolicum		40.37	0.44			
Clostridium botulinum biovar A		10.35	0.35			
41110061	**Very good identification**					**41110061**
Clostridium novyi biovar A		98.85	0.58	FRU 10		
Clostridium glycolicum		0.92	0.10			
Clostridium botulinum biovar A		0.24	0.01			
41110062	**Low discrimination**					**41110062**
Clostridium novyi biovar A		83.56	0.45	FRU 10		
Clostridium sporogenes		5.81	0.11			
Clostridium glycolicum		4.40	0.10			
41110063	**Good identification**					**41110063**
Clostridium novyi biovar A		100.0	0.45	FRU 10		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41110070	**Very good identification**						**41110070**
Clostridium novyi biovar A		98.65	0.58	FRU 10			
Clostridium glycolicum		0.92	0.10				
Clostridium botulinum biovar A		0.24	0.01				
41110071	**Very good identification**						**41110071**
Clostridium novyi biovar A		100.0	0.58	FRU 10			
41110072	**Good identification**						**41110072**
Clostridium novyi biovar A		100.0	0.45	FRU 10			
41110073	**Good identification**						**41110073**
Clostridium novyi biovar A		100.0	0.45	FRU 10			
41110100	**Low discrimination**						**41110100**
					GRA	SPO	
Clostridium sporogenes		65.15	0.53	LAC 1	99	99	
Clostridium botulinum biovar A		14.12	0.44		99	99	
Fusobacterium mortiferum		9.11	0.54		1	1	
41110101	**Low discrimination**						**41110101**
Clostridium novyi biovar A		44.05	0.37	FRU 10			
Clostridium sporogenes		26.73	0.19	LAC 1			
Clostridium sordellii		11.41	0.10				
41110102	**Acceptable identification**						**41110102**
					GRA	SPO	
Clostridium sporogenes		92.04	0.45	LAC 1	99	99	
Fusobacterium mortiferum		4.29	0.37		1	1	
Atopobium minutum		1.09	0.22		99	1	
41110110	**Low discrimination**						**41110110**
Clostridium novyi biovar A		48.31	0.37	FRU 10			
Clostridium sporogenes		29.32	0.19	LAC 1			
Clostridium difficile		10.04	0.10				
41110111	**Good identification**						**41110111**
Clostridium novyi biovar A		100.0	0.37	FRU 10			
41110140	**Low discrimination**						**41110140**
Clostridium glycolicum		71.76	0.44	XYL 95			
Clostridium botulinum biovar A		18.40	0.35	LAC 1			
Clostridium novyi biovar A		3.15	0.34				
41110200	**Low discrimination**						**41110200**
					GRA	SPO	
Clostridium sporogenes		75.17	0.53	RHA 1	99	99	
Clostridium botulinum biovar A		16.29	0.44		99	99	
Prevotella disiens		2.16	0.23		1	1	
41110201	**Low discrimination**						**41110201**
Clostridium novyi biovar A		50.06	0.37	FRU 10			
Clostridium sporogenes		30.39	0.19	RHA 1			
Clostridium sordellii		12.97	0.10				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41110202	**Good identification**						**41110202**
					SPO		
Clostridium sporogenes		96.86	0.45	RHA 1	99		
Atopobium minutum		1.14	0.22		1		
Clostridium novyi biovar A		0.85	0.24		99		
41110210	**Low discrimination**						**41110210**
Clostridium novyi biovar A		51.38	0.37	FRU 10			
Clostridium sporogenes		31.19	0.19	RHA 1			
Clostridium difficile		10.68	0.10				
41110211	**Good identification**						**41110211**
Clostridium novyi biovar A		100.0	0.37	FRU 10			
41110240	**Low discrimination**						**41110240**
Clostridium glycolicum		72.19	0.44	XYL 95			
Clostridium botulinum biovar A		18.51	0.35	RHA 1			
Clostridium novyi biovar A		3.17	0.34				
41110400	**Low discrimination**						**41110400**
					GRA	SPO	
Clostridium sporogenes		74.76	0.53	ARA 1	99	99	
Clostridium botulinum biovar A		16.20	0.44		99	99	
Prevotella disiens		2.15	0.23		1	1	
41110401	**Low discrimination**						**41110401**
Clostridium novyi biovar A		50.06	0.37	FRU 10			
Clostridium sporogenes		30.39	0.19	ARA 1			
Clostridium sordellii		12.97	0.10				
41110402	**Good identification**						**41110402**
					SPO		
Clostridium sporogenes		96.86	0.45	ARA 1	99		
Atopobium minutum		1.14	0.22		1		
Clostridium novyi biovar A		0.85	0.24		99		
41110410	**Low discrimination**						**41110410**
Clostridium novyi biovar A		51.38	0.37	FRU 10			
Clostridium sporogenes		31.19	0.19	ARA 1			
Clostridium difficile		10.68	0.10				
41110411	**Good identification**						**41110411**
Clostridium novyi biovar A		100.0	0.37	FRU 10			
41110440	**Low discrimination**						**41110440**
Clostridium glycolicum		72.19	0.44	XYL 95			
Clostridium botulinum biovar A		18.51	0.35	ARA 1			
Clostridium novyi biovar A		3.17	0.34				
41111000	**Low discrimination**						**41111000**
					SPO		
Clostridium sporogenes		48.40	0.53	GAL 1	99		
Clostridium botulinum biovar A		10.49	0.44		99		
Atopobium minutum		9.42	0.51		1		
41111001	**Low discrimination**						**41111001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		65.77	0.54	FRU 10		
Clostridium sordellii		27.06	0.31	GAL 10		
Clostridium sporogenes		3.63	0.19			
41111002	**Low discrimination**					**41111002**
					SPO	
Clostridium sporogenes		61.10	0.45	GAL 1	99	
Atopobium minutum		23.80	0.48		1	
Lactobacillus fermentum		5.95	0.44		1	
41111003	**Low discrimination**					**41111003**
					SPO	
Clostridium novyi biovar A		84.61	0.42	FRU 10	99	
Clostridium sporogenes		8.82	0.11	GAL 10	99	
Atopobium minutum		3.43	0.14		1	
41111010	**Low discrimination**					**41111010**
Clostridium novyi biovar A		88.38	0.54	FRU 10		
Clostridium sporogenes		4.88	0.19	GAL 10		
Clostridium difficile		1.67	0.10			
41111011	**Good identification**					**41111011**
Clostridium novyi biovar A		99.59	0.54	FRU 10		
Clostridium sordellii		0.41	-.03	GAL 10		
41111012	**Low discrimination**					**41111012**
					SPO	
Clostridium novyi biovar A		86.06	0.42	FRU 10	99	
Clostridium sporogenes		8.97	0.11	GAL 10	99	
Atopobium minutum		3.49	0.14		1	
41111013	**Good identification**					**41111013**
Clostridium novyi biovar A		100.0	0.42	FRU 10		
41111020	**Low discrimination**					**41111020**
					SPO	
Atopobium minutum		42.69	0.43	MLT 1	1	
Clostridium novyi biovar A		30.09	0.44		99	
Lactobacillus fermentum		7.11	0.36		1	
41111021	**Good identification**					**41111021**
Clostridium novyi biovar A		95.33	0.44	FRU 10		
Clostridium chauvoei		1.72	0.04	GAL 10		
Clostridium sordellii		1.59	-.03			
41111022	**Low discrimination**					**41111022**
					SPO	
Atopobium minutum		62.42	0.40	MLT 1	1	
Lactobacillus fermentum		15.60	0.36		1	
Clostridium novyi biovar A		11.65	0.31		99	
41111030	**Good identification**					**41111030**
					SPO	
Clostridium novyi biovar A		97.42	0.44	FRU 10	99	
Atopobium minutum		1.40	0.09	GAL 10	1	
Eubacterium saburreum		1.18	0.11		1	
41111031	**Good identification**					**41111031**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.44	FRU 10			
41111040	**Good identification**						**41111040**
Clostridium glycolicum		92.91	0.62	GAL 10			
Clostridium novyi biovar A		4.08	0.51	XYL 95			
Clostridium botulinum biovar A		2.17	0.35				
41111041	**Low discrimination**						**41111041**
Clostridium novyi biovar A		79.31	0.51	FRU 10			
Clostridium glycolicum		18.23	0.28	GAL 10			
Clostridium sordellii		2.04	0.07				
41111043	**Good identification**						**41111043**
Clostridium novyi biovar A		100.0	0.39	FRU 10			
41111050	**Low discrimination**						**41111050**
Clostridium novyi biovar A		80.96	0.51	FRU 10			
Clostridium glycolicum		18.61	0.28	GAL 10			
Clostridium botulinum biovar A		0.43	0.01				
41111051	**Good identification**						**41111051**
Clostridium novyi biovar A		100.0	0.51	FRU 10			
41111052	**Good identification**						**41111052**
Clostridium novyi biovar A		100.0	0.39	FRU 10			
41111053	**Good identification**						**41111053**
Clostridium novyi biovar A		100.0	0.39	FRU 10			
41111060	**Low discrimination**						**41111060**
Clostridium novyi biovar A		50.09	0.41	FRU 10			
Clostridium glycolicum		46.06	0.28	GAL 10			
Clostridium perfringens		1.70	0.07				
41111061	**Good identification**						**41111061**
Clostridium novyi biovar A		100.0	0.41	FRU 10			
41111070	**Good identification**						**41111070**
Clostridium novyi biovar A		100.0	0.41	FRU 10			
41111071	**Good identification**						**41111071**
Clostridium novyi biovar A		100.0	0.41	FRU 10			
41111100	**Low discrimination**						**41111100**
					COC	GRA	
Fusobacterium mortiferum		44.93	0.57	MLT 11	1	1	
Actinomyces meyeri		20.57	0.42		1	99	
Blautia hansenii		15.83	0.32		99	99	
41111101	**Good identification**						**41111101**
					GRA	SPO	
Clostridium chauvoei		97.24	0.47	SUC 99	99	99	
Fusobacterium mortiferum		0.84	0.23	MNS 90	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		0.73	0.20		99	99	
41111102	**Low discrimination**						**41111102**
					COC	GRA	
Lactobacillus fermentum		53.55	0.48	ESL 1	1	99	
Fusobacterium mortiferum		23.56	0.40	SUC 87	1	1	
Blautia hansenii		8.30	0.15		99	99	
41111120	**Low discrimination**						**41111120**
					GRA	SPO	
Clostridium septicum		55.78	0.39	MNS 99	99	99	
Prevotella melaninogenica		17.83	0.30		1	1	
Clostridium perfringens		11.42	0.34		99	84	
41111121	**Good identification**						**41111121**
					GRA	SPO	
Clostridium chauvoei		96.39	0.38	SUC 99	99	99	
Clostridium septicum		2.32	0.05	MNS 90	99	99	
Prevotella melaninogenica		0.74	-.04		1	1	
41111400	**Low discrimination**						**41111400**
					SPO		
Lactobacillus fermentum		54.56	0.44	ESL 1	1		
Clostridium sporogenes		16.99	0.19	SUC 87	99		
Actinomyces meyeri		10.09	0.21		1		
41111402	**Low discrimination**						**41111402**
					SPO		
Lactobacillus fermentum		87.40	0.44	ESL 1	1		
Clostridium sporogenes		9.07	0.11	SUC 87	99		
Atopobium minutum		3.53	0.14		1		
41111420	**Low discrimination**						**41111420**
Lactobacillus fermentum		84.82	0.36	ESL 1			
Actinomyces meyeri		6.42	0.06	SUC 87			
Atopobium minutum		5.14	0.09				
41111422	**Good identification**						**41111422**
Lactobacillus fermentum		96.11	0.36	ESL 1			
Atopobium minutum		3.89	0.05	SUC 87			
41111500	**Low discrimination**						**41111500**
					COC		
Lactobacillus fermentum		57.49	0.48	ESL 1	1		
Actinomyces meyeri		26.06	0.31	SUC 87	1		
Blautia hansenii		14.15	0.19		99		
41111502	**Acceptable identification**						**41111502**
					COC		
Lactobacillus fermentum		94.76	0.48	ESL 1	1		
Blautia hansenii		2.59	0.03	SUC 87	99		
Actinomyces meyeri		1.33	0.06		1		
41111520	**Low discrimination**						**41111520**
					SPO		
Lactobacillus fermentum		62.52	0.40	ESL 1	1		
Actinomyces meyeri		11.59	0.17	SUC 87	1		
Clostridium septicum		6.99	0.05		99		
41111522	**Low discrimination**						**41111522**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		68.88	0.40	ESL 1	99	1		
Capnocytophaga ochracea		25.84	0.17	SUC 87	1	90		
Bifidobacterium longum subsp. longum		5.28	0.14		99	1		
41112000	**Low discrimination**							**41112000**
					COC	SPO	GRA	
Eubacterium limosum		58.79	0.63	MLT 1	1	1	99	
Megasphaera elsdenii		18.07	0.63		99	1	1	
Clostridium sporogenes		11.63	0.53		1	99	99	
41112002	**Low discrimination**							**41112002**
					COC	SPO	GRA	
Clostridium sporogenes		57.63	0.45	MAN 1	1	99	99	
Megasphaera elsdenii		29.85	0.47		99	1	1	
Eubacterium limosum		8.83	0.29		1	1	99	
41112010	**Very good identification**							**41112010**
					SPO			
Clostridium difficile		96.97	0.61	MLT 1	99			
Eubacterium limosum		1.61	0.29		1			
Clostridium novyi biovar A		0.52	0.37		99			
41112040	**Low discrimination**							**41112040**
					COC	GRA		
Megasphaera elsdenii		70.33	0.60	ESL 1	99	1		
Clostridium glycolicum		17.14	0.44		1	99		
Clostridium botulinum biovar A		4.40	0.35		1	99		
41112042	**Acceptable identification**							**41112042**
					COC	GRA		
Megasphaera elsdenii		91.62	0.44	ESL 1	99	1		
Clostridium sporogenes		2.68	0.11	BGL 10	1	99		
Clostridium glycolicum		2.03	0.10		1	99		
41112050	**Low discrimination**							**41112050**
					COC	GRA		
Clostridium difficile		86.52	0.39	MLT 1	1	99		
Clostridium novyi biovar A		5.92	0.34	SOR 5	1	99		
Megasphaera elsdenii		5.58	0.26		99	1		
41114000	**Good identification**							**41114000**
Clostridium glycolicum		94.84	0.78	SOR 95				
Clostridium sporogenes		3.80	0.53					
Clostridium botulinum biovar A		0.82	0.44					
41114001	**Low discrimination**							**41114001**
Clostridium glycolicum		88.36	0.44	SOR 95				
Clostridium novyi biovar A		5.83	0.37	URE 1				
Clostridium sporogenes		3.54	0.19					
41114002	**Low discrimination**							**41114002**
					SPO			
Clostridium sporogenes		55.74	0.45	XYL 1	99			
Clostridium glycolicum		42.21	0.44		99			
Atopobium minutum		0.66	0.22		1			
41114010	**Low discrimination**							**41114010**
Clostridium glycolicum		84.32	0.44	MLZ 1				
Clostridium difficile		6.02	0.23	SOR 95				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		5.56	0.37				
41114011	**Low discrimination**						**41114011**
Clostridium novyi biovar A		86.71	0.37	FRU 10			
Clostridium glycolicum		13.29	0.10	XYL 1			
41114020	**Acceptable identification**						**41114020**
					SPO		
Clostridium glycolicum		91.31	0.44	NAG 1	99		
Clostridium sporogenes		3.65	0.19	SOR 95	99		
Actinomyces meyeri		1.53	0.19		1		
41114040	**Excellent identification**						**41114040**
Clostridium glycolicum		99.98	1.00				
Clostridium botulinum biovar A		0.01	0.35				
Clostridium novyi biovar A		0.00	0.34				
41114041	**Very good identification**						**41114041**
Clostridium glycolicum		99.76	0.66	URE 1			
Clostridium novyi biovar A		0.23	0.34				
Clostridium botulinum biovar A		0.01	0.01				
41114042	**Very good identification**						**41114042**
Clostridium glycolicum		99.85	0.66	BGL 1			
Clostridium sporogenes		0.07	0.11				
Clostridium novyi biovar A		0.04	0.21				
41114050	**Very good identification**						**41114050**
Clostridium glycolicum		99.74	0.66	MLZ 1			
Clostridium novyi biovar A		0.23	0.34				
Clostridium difficile		0.02	0.01				
41114060	**Very good identification**						**41114060**
Clostridium glycolicum		99.93	0.66	NAG 1			
Clostridium novyi biovar A		0.06	0.23				
Clostridium botulinum biovar A		0.01	0.01				
41114100	**Low discrimination**						**41114100**
					GRA	SPO	
Clostridium glycolicum		63.20	0.44	LAC 1	99	99	
Actinomyces meyeri		31.09	0.44	SOR 95	99	1	
Fusobacterium mortiferum		2.63	0.35		1	1	
41114140	**Very good identification**						**41114140**
					SPO		
Clostridium glycolicum		99.96	0.66	LAC 1	99		
Actinomyces meyeri		0.03	0.09		1		
Clostridium botulinum biovar A		0.01	0.01		99		
41114200	**Good identification**						**41114200**
Clostridium glycolicum		95.36	0.44	RHA 1			
Clostridium sporogenes		3.82	0.19	SOR 95			
Clostridium botulinum biovar A		0.83	0.10				
41114240	**Very good identification**						**41114240**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium glycolicum		99.99	0.66	RHA 1			
Clostridium botulinum biovar A		0.01	0.01				
41114400	**Acceptable identification**						**41114400**
					SPO		
Clostridium glycolicum		92.64	0.44	ARA 1	99		
Clostridium sporogenes		3.71	0.19	SOR 95	99		
Actinomyces meyeri		2.85	0.23		1		
41114440	**Very good identification**						**41114440**
Clostridium glycolicum		99.99	0.66	ARA 1			
Clostridium botulinum biovar A		0.01	0.01				
41115000	**Very good identification**						**41115000**
					SPO		
Clostridium glycolicum		96.76	0.62	GAL 10	99		
Actinomyces meyeri		1.62	0.36	SOR 95	1		
Lactobacillus fermentum		0.66	0.41		1		
41115040	**Very good identification**						**41115040**
					SPO		
Clostridium glycolicum		100.0	0.84	GAL 10	99		
Clostridium novyi biovar A		0.00	0.17		99		
Actinomyces meyeri		0.00	0.02		1		
41115041	**Good identification**						**41115041**
Clostridium glycolicum		99.77	0.50	GAL 10			
Clostridium novyi biovar A		0.23	0.17	URE 1			
41115042	**Good identification**						**41115042**
Clostridium glycolicum		100.0	0.50	GAL 10			
41115050	**Good identification**						**41115050**
Clostridium glycolicum		99.77	0.50	GAL 10			
Clostridium novyi biovar A		0.23	0.17	MLZ 1			
41115060	**Good identification**						**41115060**
Clostridium glycolicum		100.0	0.50	GAL 10			
41115100	**Low discrimination**						**41115100**
					SPO		
Actinomyces meyeri		66.65	0.47	ESL 1	1		
Lactobacillus fermentum		11.12	0.44	SUC 93	1		
Clostridium glycolicum		10.04	0.28		99		
41115102	**Low discrimination**						**41115102**
					GRA		
Lactobacillus fermentum		74.11	0.44	ESL 1	99		
Actinomyces meyeri		13.74	0.21	SUC 87	99		
Fusobacterium mortiferum		4.18	0.21		1		
41115122	**Low discrimination**						**41115122**
					GRA	CAT	
Lactobacillus fermentum		45.37	0.36	ESL 1	99	1	
Capnocytophaga ochracea		41.06	0.20	SUC 87	1	90	
Bacteroides fragilis		7.06	-.03		1	99	
41115140	**Good identification**						**41115140**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium glycolicum		99.65	0.50	GAL 10	99		
Actinomyces meyeri		0.35	0.13	LAC 1	1		
41115240	**Good identification**						**41115240**
Clostridium glycolicum		100.0	0.50	GAL 10			
41115402	**Good identification**						**41115402**
Lactobacillus fermentum		100.0	0.41	ESL 1			
41115440	**Good identification**						**41115440**
Clostridium glycolicum		100.0	0.50	GAL 10			
41115500	**Low discrimination**						**41115500**
					COC		
Actinomyces meyeri		57.52	0.36	ESL 1	1		
Lactobacillus fermentum		38.39	0.44	SUC 93	1		
Blautia hansenii		2.84	0.06		99		
41115502	**Low discrimination**						**41115502**
Lactobacillus fermentum		89.84	0.44	ESL 1			
Actinomyces meyeri		4.16	0.11	SUC 87			
Bifidobacterium adolescentis		3.06	0.14				
41115520	**Low discrimination**						**41115520**
Lactobacillus fermentum		46.64	0.36	ESL 1			
Actinomyces meyeri		28.58	0.22	SUC 87			
Bifidobacterium longum subsp. longum		18.24	0.22				
41115522	**Low discrimination**						**41115522**
					GRA	CAT	
Lactobacillus fermentum		64.48	0.36	ESL 1	99	1	
Bifidobacterium longum subsp. longum		25.22	0.22	SUC 87	99	1	
Capnocytophaga ochracea		10.30	0.07		1	90	
41116000	**Low discrimination**						**41116000**
					SPO		
Clostridium glycolicum		69.12	0.44	MAN 1	99		
Eubacterium limosum		13.98	0.29	SOR 95	1		
Clostridium difficile		9.09	0.27		99		
41116010	**Good identification**						**41116010**
					SPO		
Clostridium difficile		98.15	0.39	MLT 1	99		
Clostridium glycolicum		1.54	0.10	XYL 5	99		
Eubacterium limosum		0.31	-.06		1		
41116040	**Very good identification**						**41116040**
					COC	GRA	
Clostridium glycolicum		99.72	0.66	MAN 1	1	99	
Megasphaera elsdenii		0.22	0.26		99	1	
Clostridium difficile		0.04	0.05		1	99	
41117040	**Good identification**						**41117040**
Clostridium glycolicum		100.0	0.50	GAL 10			
41120000	**Low discrimination**						**41120000**
Clostridium sporogenes		68.83	0.92				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium botulinum biovar A		30.93	0.88			
Clostridium bifermentans		0.15	0.48			
41120001	**Low discrimination**					**41120001**
Clostridium sporogenes		65.22	0.58	URE 1		
Clostridium botulinum biovar A		29.30	0.54			
Clostridium novyi biovar A		5.17	0.53			
41120002	**Very good identification**					**41120002**
Clostridium sporogenes		98.60	0.84			
Clostridium botulinum biovar A		1.34	0.54			
Clostridium novyi biovar A		0.04	0.40			
41120003	**Acceptable identification**					**41120003**
Clostridium sporogenes		94.73	0.50	URE 1		
Clostridium novyi biovar A		3.98	0.40			
Clostridium botulinum biovar A		1.29	0.20			
41120010	**Low discrimination**					**41120010**
Clostridium sporogenes		65.32	0.58	MLZ 1		
Clostridium botulinum biovar A		29.35	0.54			
Clostridium novyi biovar A		5.18	0.53			
41120011	**Low discrimination**					**41120011**
Clostridium novyi biovar A		84.42	0.53	TRE 1		
Clostridium sporogenes		10.75	0.24			
Clostridium botulinum biovar A		4.83	0.20			
41120012	**Acceptable identification**					**41120012**
Clostridium sporogenes		94.73	0.50	MLZ 1		
Clostridium novyi biovar A		3.98	0.40			
Clostridium botulinum biovar A		1.29	0.20			
41120013	**Low discrimination**					**41120013**
Clostridium novyi biovar A		80.60	0.40	TRE 1		
Clostridium sporogenes		19.40	0.15			
41120020	**Low discrimination**					**41120020**
Clostridium sporogenes		67.37	0.58	NAG 1		
Clostridium botulinum biovar A		30.27	0.54			
Clostridium novyi biovar A		1.34	0.43			
41120021	**Low discrimination**					**41120021**
Clostridium novyi biovar A		57.52	0.43	TRE 1		
Clostridium sporogenes		29.31	0.24			
Clostridium botulinum biovar A		13.17	0.20			
41120022	**Good identification**					**41120022**
Clostridium sporogenes		97.28	0.50	NAG 1		
Clostridium botulinum biovar A		1.32	0.20			
Clostridium novyi biovar A		1.02	0.30			
41120030	**Low discrimination**					**41120030**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		57.52	0.43	TRE 1		
Clostridium sporogenes		29.31	0.24			
Clostridium botulinum biovar A		13.17	0.20			
41120031	**Good identification**					**41120031**
Clostridium novyi biovar A		100.0	0.43	TRE 1		
41120040	**Good identification**					**41120040**
Clostridium botulinum biovar A		92.49	0.79			
Clostridium sporogenes		6.96	0.58			
Clostridium novyi biovar A		0.37	0.50			
41120041	**Low discrimination**					**41120041**
Clostridium botulinum biovar A		68.07	0.45	URE 1		
Clostridium novyi biovar A		26.81	0.50			
Clostridium sporogenes		5.12	0.24			
41120042	**Low discrimination**					**41120042**
Clostridium sporogenes		69.90	0.50	SOR 1		
Clostridium botulinum biovar A		28.14	0.45			
Clostridium novyi biovar A		1.96	0.37			
41120043	**Low discrimination**					**41120043**
Clostridium novyi biovar A		66.39	0.37	TRE 1		
Clostridium sporogenes		23.96	0.15			
Clostridium botulinum biovar A		9.65	0.11			
41120050	**Low discrimination**					**41120050**
Clostridium botulinum biovar A		67.92	0.45	MLZ 1		
Clostridium novyi biovar A		26.75	0.50			
Clostridium sporogenes		5.11	0.24			
41120051	**Good identification**					**41120051**
Clostridium novyi biovar A		97.50	0.50	TRE 1		
Clostridium botulinum biovar A		2.50	0.11			
41120052	**Low discrimination**					**41120052**
Clostridium novyi biovar A		66.39	0.37	TRE 1		
Clostridium sporogenes		23.96	0.15			
Clostridium botulinum biovar A		9.65	0.11			
41120053	**Good identification**					**41120053**
Clostridium novyi biovar A		100.0	0.37	TRE 1		
41120060	**Low discrimination**					**41120060**
Clostridium botulinum biovar A		84.65	0.45	NAG 1		
Clostridium novyi biovar A		8.34	0.40			
Clostridium sporogenes		6.37	0.24			
41120061	**Acceptable identification**					**41120061**
Clostridium novyi biovar A		90.70	0.40	TRE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium botulinum biovar A		9.30	0.11			
41120070	**Acceptable identification**					**41120070**
Clostridium novyi biovar A		90.70	0.40	TRE 1		
Clostridium botulinum biovar A		9.30	0.11			
41120071	**Good identification**					**41120071**
Clostridium novyi biovar A		100.0	0.40	TRE 1		
41120100	**Low discrimination**					**41120100**
Clostridium sporogenes		68.83	0.58	LAC 1		
Clostridium botulinum biovar A		30.92	0.54			
Clostridium bifermentans		0.15	0.14			
41120102	**Good identification**					**41120102**
Clostridium sporogenes		98.66	0.50	LAC 1		
Clostridium botulinum biovar A		1.34	0.20			
41120140	**Acceptable identification**					**41120140**
Clostridium botulinum biovar A		92.66	0.45	LAC 1		
Clostridium sporogenes		6.97	0.24			
Clostridium novyi biovar A		0.37	0.16			
41120200	**Low discrimination**					**41120200**
Clostridium sporogenes		68.84	0.58	RHA 1		
Clostridium botulinum biovar A		30.93	0.54			
Clostridium bifermentans		0.15	0.14			
41120202	**Good identification**					**41120202**
Clostridium sporogenes		98.66	0.50	RHA 1		
Clostridium botulinum biovar A		1.34	0.20			
41120240	**Acceptable identification**					**41120240**
Clostridium botulinum biovar A		92.66	0.45	RHA 1		
Clostridium sporogenes		6.97	0.24			
Clostridium novyi biovar A		0.37	0.16			
41120400	**Low discrimination**					**41120400**
Clostridium sporogenes		68.84	0.58	ARA 1		
Clostridium botulinum biovar A		30.93	0.54			
Clostridium bifermentans		0.15	0.14			
41120402	**Good identification**					**41120402**
Clostridium sporogenes		98.66	0.50	ARA 1		
Clostridium botulinum biovar A		1.34	0.20			
41120440	**Acceptable identification**					**41120440**
Clostridium botulinum biovar A		92.66	0.45	ARA 1		
Clostridium sporogenes		6.97	0.24			
Clostridium novyi biovar A		0.37	0.16			
41121000	**Low discrimination**					**41121000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		68.37	0.58	GAL 1			
Clostridium botulinum biovar A		30.72	0.54				
Clostridium novyi biovar A		0.60	0.37				
41121002	**Good identification**						**41121002**
Clostridium sporogenes		97.99	0.50	GAL 1			
Clostridium botulinum biovar A		1.33	0.20				
Clostridium novyi biovar A		0.46	0.24				
41121011	**Good identification**						**41121011**
Clostridium novyi biovar A		100.0	0.37	TRE 1			
41121040	**Low discrimination**						**41121040**
Clostridium botulinum biovar A		88.39	0.45	GAL 1			
Clostridium sporogenes		6.65	0.24				
Clostridium novyi biovar A		3.87	0.34				
41122000	**Low discrimination**						**41122000**
Clostridium sporogenes		68.80	0.58	MAN 1			
Clostridium botulinum biovar A		30.91	0.54				
Clostridium bifermentans		0.15	0.14				
41122002	**Good identification**						**41122002**
Clostridium sporogenes		98.66	0.50	MAN 1			
Clostridium botulinum biovar A		1.34	0.20				
41122040	**Acceptable identification**						**41122040**
Clostridium botulinum biovar A		92.47	0.45	MAN 1			
Clostridium sporogenes		6.96	0.24				
Clostridium novyi biovar A		0.37	0.16				
41124000	**Low discrimination**						**41124000**
Clostridium sporogenes		68.15	0.58	XYL 1			
Clostridium botulinum biovar A		30.62	0.54				
Clostridium glycolicum		1.01	0.28				
41124002	**Good identification**						**41124002**
Clostridium sporogenes		98.66	0.50	XYL 1			
Clostridium botulinum biovar A		1.34	0.20				
41124040	**Low discrimination**						**41124040**
Clostridium glycolicum		66.08	0.50	FRU 90			
Clostridium botulinum biovar A		31.43	0.45	TRE 1			
Clostridium sporogenes		2.37	0.24				
41130000	**Low discrimination**						**41130000**
					GRA	SPO	
Clostridium sporogenes		85.72	0.79		99	99	
Clostridium botulinum biovar A		13.93	0.68		99	99	
Prevotella disiens		0.08	0.23		1	1	
41130001	**Low discrimination**						**41130001**
Clostridium sporogenes		81.61	0.45	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		13.26	0.34				
Clostridium novyi biovar A		4.07	0.37				
41130002	**Very good identification**						**41130002**
					SPO		
Clostridium sporogenes		99.42	0.71		99		
Clostridium botulinum biovar A		0.49	0.34		99		
Atopobium minutum		0.04	0.22		1		
41130003	**Good identification**						**41130003**
Clostridium sporogenes		96.96	0.37	URE 1			
Clostridium novyi biovar A		2.56	0.24				
Clostridium botulinum biovar A		0.48	-.01				
41130010	**Low discrimination**						**41130010**
Clostridium sporogenes		77.54	0.45	MLZ 1			
Clostridium botulinum biovar A		12.60	0.34				
Clostridium difficile		5.99	0.25				
41130011	**Low discrimination**						**41130011**
Clostridium novyi biovar A		80.96	0.37	FRU 10			
Clostridium sporogenes		16.38	0.11	TRE 1			
Clostridium botulinum biovar A		2.66	-.01				
41130012	**Good identification**						**41130012**
Clostridium sporogenes		96.96	0.37	MLZ 1			
Clostridium novyi biovar A		2.56	0.24				
Clostridium botulinum biovar A		0.48	-.01				
41130020	**Low discrimination**						**41130020**
Clostridium sporogenes		84.39	0.45	NAG 1			
Clostridium botulinum biovar A		13.71	0.34				
Clostridium novyi biovar A		1.05	0.26				
41130022	**Good identification**						**41130022**
					SPO		
Clostridium sporogenes		97.72	0.37	NAG 1	99		
Atopobium minutum		1.15	0.14		1		
Clostridium novyi biovar A		0.65	0.13		99		
41130040	**Low discrimination**						**41130040**
Clostridium botulinum biovar A		72.26	0.59	FRU 6			
Clostridium sporogenes		15.04	0.45				
Clostridium glycolicum		11.39	0.44				
41130042	**Low discrimination**						**41130042**
Clostridium sporogenes		83.57	0.37	SOR 1			
Clostridium botulinum biovar A		12.17	0.25				
Clostridium glycolicum		1.92	0.10				
41130100	**Low discrimination**						**41130100**
					GRA	SPO	
Clostridium sporogenes		77.54	0.45	LAC 1	99	99	
Clostridium botulinum biovar A		12.60	0.34		99	99	
Fusobacterium mortiferum		9.71	0.45		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41130102	**Good identification**						**41130102**
					GRA	SPO	
Clostridium sporogenes		95.54	0.37	LAC 1	99	99	
Fusobacterium mortiferum		3.99	0.28		1	1	
Clostridium botulinum biovar A		0.47	-.01		99	99	
41130200	**Low discrimination**						**41130200**
Clostridium sporogenes		86.02	0.45	RHA 1			
Clostridium botulinum biovar A		13.98	0.34				
41130202	**Good identification**						**41130202**
Clostridium sporogenes		99.51	0.37	RHA 1			
Clostridium botulinum biovar A		0.49	-.01				
41130400	**Low discrimination**						**41130400**
Clostridium sporogenes		86.02	0.45	ARA 1			
Clostridium botulinum biovar A		13.98	0.34				
41130402	**Good identification**						**41130402**
Clostridium sporogenes		99.51	0.37	ARA 1			
Clostridium botulinum biovar A		0.49	-.01				
41131000	**Low discrimination**						**41131000**
					GRA	SPO	
Clostridium sporogenes		78.64	0.45	GAL 1	99	99	
Clostridium botulinum biovar A		12.78	0.34		99	99	
Fusobacterium mortiferum		6.33	0.42		1	1	
41131002	**Acceptable identification**						**41131002**
					GRA	SPO	
Clostridium sporogenes		95.00	0.37	GAL 1	99	99	
Fusobacterium mortiferum		2.55	0.25		1	1	
Atopobium minutum		1.12	0.14		99	1	
41131100	**Low discrimination**						**41131100**
					COC	GRA	
Fusobacterium mortiferum		58.56	0.48	MLT 11	1	1	
Atopobium parvulum		22.40	0.29		99	99	
Eubacterium saburreum		5.60	0.29		1	85	
41131120	**Low discrimination**						**41131120**
					COC	SPO	
Clostridium septicum		87.88	0.47	MNS 99	1	99	
Clostridium perfringens		11.13	0.38		1	84	
Atopobium parvulum		0.63	0.16		99	1	
41132000	**Low discrimination**						**41132000**
					SPO		
Clostridium sporogenes		66.36	0.45	MAN 1	99		
Clostridium botulinum biovar A		10.79	0.34		99		
Eubacterium limosum		10.16	0.29		1		
41132002	**Good identification**						**41132002**
					COC	GRA	
Clostridium sporogenes		97.13	0.37	MAN 1	1	99	
Megasphaera elsdenii		1.52	0.12		99	1	
Clostridium botulinum biovar A		0.48	-.01		1	99	
41132010	**Good identification**						**41132010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium difficile		98.13	0.42	MLT 1		
Clostridium sporogenes		1.43	0.11	TRE 7		
Clostridium botulinum biovar A		0.23	-.01			
41134000	**Low discrimination**					**41134000**
Clostridium sporogenes		51.98	0.45	XYL 1		
Clostridium glycolicum		39.37	0.44			
Clostridium botulinum biovar A		8.45	0.34			
41134002	**Good identification**					**41134002**
Clostridium sporogenes		97.29	0.37	XYL 1		
Clostridium glycolicum		2.23	0.10			
Clostridium botulinum biovar A		0.48	-.01			
41134040	**Very good identification**					**41134040**
Clostridium glycolicum		99.59	0.66	TRE 1		
Clostridium botulinum biovar A		0.34	0.25			
Clostridium sporogenes		0.07	0.11			
41135040	**Good identification**					**41135040**
Clostridium glycolicum		100.0	0.50	TRE 1		
41140000	**Low discrimination**					**41140000**
Clostridium sporogenes		58.37	0.66	CEL 1		
Clostridium botulinum biovar A		34.97	0.64			
Clostridium bifermentans		4.19	0.48			
41140001	**Low discrimination**					**41140001**
Clostridium novyi biovar A		59.85	0.53	CEL 1		
Clostridium sporogenes		22.87	0.32			
Clostridium botulinum biovar A		13.70	0.30			
41140002	**Very good identification**					**41140002**
Clostridium sporogenes		96.45	0.58	CEL 1		
Clostridium botulinum biovar A		1.75	0.30			
Clostridium novyi biovar A		1.35	0.40			
41140003	**Low discrimination**					**41140003**
Clostridium novyi biovar A		58.08	0.40	CEL 1		
Clostridium sporogenes		41.92	0.24			
41140010	**Low discrimination**					**41140010**
Clostridium novyi biovar A		60.90	0.53	CEL 1		
Clostridium sporogenes		23.27	0.32			
Clostridium botulinum biovar A		13.94	0.30			
41140011	**Good identification**					**41140011**
Clostridium novyi biovar A		99.62	0.53	CEL 1		
Clostridium sporogenes		0.39	-.03			
41140012	**Low discrimination**					**41140012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		58.08	0.40	CEL 1		
Clostridium sporogenes		41.92	0.24			
41140013	**Good identification**					**41140013**
Clostridium novyi biovar A		100.0	0.40	CEL 1		
41140021	**Good identification**					**41140021**
Clostridium novyi biovar A		98.48	0.43	CEL 1		
Clostridium sporogenes		1.52	-.03			
41140030	**Good identification**					**41140030**
Clostridium novyi biovar A		98.48	0.43	CEL 1		
Clostridium sporogenes		1.52	-.03			
41140031	**Good identification**					**41140031**
Clostridium novyi biovar A		100.0	0.43	CEL 1		
41140040	**Low discrimination**					**41140040**
Clostridium botulinum biovar A		83.31	0.55	CEL 1		
Clostridium novyi biovar A		8.20	0.50			
Clostridium sporogenes		4.70	0.32			
41140041	**Acceptable identification**					**41140041**
Clostridium novyi biovar A		90.22	0.50	CEL 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41140043	**Good identification**					**41140043**
Clostridium novyi biovar A		100.0	0.37	CEL 1		
41140050	**Acceptable identification**					**41140050**
Clostridium novyi biovar A		90.22	0.50	CEL 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41140051	**Good identification**					**41140051**
Clostridium novyi biovar A		100.0	0.50	CEL 1		
41140052	**Good identification**					**41140052**
Clostridium novyi biovar A		100.0	0.37	CEL 1		
41140053	**Good identification**					**41140053**
Clostridium novyi biovar A		100.0	0.37	CEL 1		
41140060	**Low discrimination**					**41140060**
Clostridium novyi biovar A		67.71	0.40	CEL 1		
Clostridium botulinum biovar A		27.78	0.21			
Clostridium bifermentans		2.94	0.04			
41140061	**Good identification**					**41140061**
Clostridium novyi biovar A		100.0	0.40	CEL 1		
41140070	**Good identification**					**41140070**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.40	CEL 1			
41140071	**Good identification**						**41140071**
Clostridium novyi biovar A		100.0	0.40	CEL 1			
41141001	**Acceptable identification**						**41141001**
Clostridium novyi biovar A		92.55	0.37	CEL 1			
Clostridium sordellii		4.23	-.03	GAL 10			
Clostridium sporogenes		3.22	-.03				
41141010	**Good identification**						**41141010**
Clostridium novyi biovar A		96.64	0.37	CEL 1			
Clostridium sporogenes		3.36	-.03	GAL 10			
41141011	**Good identification**						**41141011**
Clostridium novyi biovar A		100.0	0.37	CEL 1			
41144040	**Good identification**						**41144040**
Clostridium glycolicum		97.83	0.50	FRU 90			
Clostridium botulinum biovar A		1.88	0.21	CEL 1			
Clostridium novyi biovar A		0.19	0.16				
41150000	**Low discrimination**						**41150000**
					GRA	SPO	
Clostridium sporogenes		73.92	0.53	CEL 1	99	99	
Clostridium botulinum biovar A		16.02	0.44		99	99	
Prevotella disiens		2.13	0.23		1	1	
41150001	**Low discrimination**						**41150001**
Clostridium novyi biovar A		50.06	0.37	FRU 10			
Clostridium sporogenes		30.39	0.19	CEL 1			
Clostridium sordellii		12.97	0.10				
41150002	**Good identification**						**41150002**
					SPO		
Clostridium sporogenes		96.24	0.45	CEL 1	99		
Atopobium minutum		1.14	0.22		1		
Clostridium novyi biovar A		0.85	0.24		99		
41150011	**Good identification**						**41150011**
Clostridium novyi biovar A		100.0	0.37	FRU 10			
41150040	**Low discrimination**						**41150040**
Clostridium glycolicum		72.19	0.44	CEL 1			
Clostridium botulinum biovar A		18.51	0.35	XYL 95			
Clostridium novyi biovar A		3.17	0.34				
41150100	**Low discrimination**						**41150100**
					GRA	SPO	
Fusobacterium mortiferum		78.27	0.46	MLT 11	1	1	
Clostridium sporogenes		16.97	0.19		99	99	
Clostridium botulinum biovar A		3.68	0.10		99	99	
41151000	**Low discrimination**						**41151000**
					GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		55.83	0.42	MLT 11	1	1	
Clostridium sporogenes		18.83	0.19		99	99	
Lactobacillus jensenii		9.85	0.19		99	1	
41151100	**Low discrimination**						**41151100**
					GRA	SPO	
Fusobacterium mortiferum		86.33	0.49	MLT 11	1	1	
Clostridium septicum		7.72	0.13		99	99	
Collinsella aerofaciens		2.48	-.03		99	1	
41151120	**Good identification**						**41151120**
					GRA	SPO	
Clostridium septicum		96.71	0.47	MNS 99	99	99	
Clostridium paraputrificum		1.51	0.14		99	99	
Prevotella melaninogenica		1.09	0.14		1	1	
41152010	**Good identification**						**41152010**
					SPO		
Clostridium difficile		99.68	0.39	MLT 1	99		
Eubacterium limosum		0.32	-.06	CEL 5	1		
41154000	**Acceptable identification**						**41154000**
Clostridium glycolicum		94.82	0.44	CEL 1			
Clostridium sporogenes		3.79	0.19	SOR 95			
Clostridium botulinum biovar A		0.82	0.10				
41154040	**Very good identification**						**41154040**
Clostridium glycolicum		99.99	0.66	CEL 1			
Clostridium botulinum biovar A		0.01	0.01				
41155040	**Good identification**						**41155040**
Clostridium glycolicum		100.0	0.50	CEL 1			
41160000	**Low discrimination**						**41160000**
Clostridium sporogenes		68.84	0.58	CEL 1			
Clostridium botulinum biovar A		30.93	0.54				
Clostridium bifermentans		0.15	0.14				
41160002	**Good identification**						**41160002**
Clostridium sporogenes		98.66	0.50	CEL 1			
Clostridium botulinum biovar A		1.34	0.20				
41160040	**Acceptable identification**						**41160040**
Clostridium botulinum biovar A		92.66	0.45	CEL 1			
Clostridium sporogenes		6.97	0.24				
Clostridium novyi biovar A		0.37	0.16				
41170000	**Low discrimination**						**41170000**
					GRA	SPO	
Clostridium sporogenes		84.72	0.45	CEL 1	99	99	
Clostridium botulinum biovar A		13.77	0.34		99	99	
Fusobacterium mortiferum		1.52	0.30		1	1	
41170002	**Good identification**						**41170002**
					GRA	SPO	
Clostridium sporogenes		98.92	0.37	CEL 1	99	99	
Fusobacterium mortiferum		0.59	0.14		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		0.49	-.01		99	99		
41170100	**Low discrimination**							**41170100**
					COC	GRA	SPO	
Fusobacterium mortiferum		75.65	0.37	MLT 11	1	1	1	
Clostridium sporogenes		18.31	0.11		1	99	99	
Atopobium parvulum		3.07	0.01		99	99	1	
41171100	**Low discrimination**							**41171100**
					COC	GRA	SPO	
Atopobium parvulum		58.52	0.36	SUC 96	99	99	1	
Fusobacterium mortiferum		21.86	0.40	MNS 99	1	1	1	
Clostridium septicum		19.62	0.21		1	99	99	
41171120	**Good identification**							**41171120**
					COC	SPO		
Clostridium septicum		99.17	0.56	MNS 99	1	99		
Atopobium parvulum		0.53	0.23		99	1		
Clostridium perfringens		0.17	0.15		1	84		
41175100	**Good identification**							**41175100**
					COC	GRA		
Atopobium parvulum		97.44	0.40	SUC 96	99	99		
Fusobacterium mortiferum		1.48	0.20	MNS 99	1	1		
Actinomyces israelii		1.09	0.21		1	99		
41177102	**Good identification**							**41177102**
Actinomyces israelii		100.0	0.36	SUC 98				
41200000	**Low discrimination**							**41200000**
Clostridium botulinum biovar A		36.90	0.66	SAL 1				
Clostridium sporogenes		27.13	0.61					
Clostridium bifermentans		17.07	0.60					
41200001	**Good identification**							**41200001**
Clostridium novyi biovar A		92.67	0.69	SAL 8				
Clostridium botulinum biovar A		3.01	0.32					
Clostridium sporogenes		2.22	0.27					
41200002	**Low discrimination**							**41200002**
					SPO			
Clostridium sporogenes		58.37	0.53	SAL 1	99			
Atopobium minutum		24.21	0.57		1			
Clostridium novyi biovar A		13.06	0.56		99			
41200003	**Good identification**							**41200003**
					SPO			
Clostridium novyi biovar A		94.00	0.56	SAL 8	99			
Clostridium sporogenes		4.24	0.19		99			
Atopobium minutum		1.76	0.22		1			
41200010	**Low discrimination**							**41200010**
Clostridium novyi biovar A		86.77	0.69	SAL 8				
Clostridium difficile		6.50	0.35					
Clostridium botulinum biovar A		2.82	0.32					
41200011	**Very good identification**							**41200011**
Clostridium novyi biovar A		99.92	0.69	SAL 8				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		0.08	0.01				
41200012	**Good identification**						**41200012**
					SPO		
Clostridium novyi biovar A		93.62	0.56	SAL 8	99		
Clostridium sporogenes		4.23	0.19		99		
Atopobium minutum		1.75	0.22		1		
41200013	**Very good identification**						**41200013**
Clostridium novyi biovar A		100.0	0.56	SAL 8			
41200020	**Low discrimination**						**41200020**
					SPO		
Clostridium novyi biovar A		43.35	0.59	SAL 8	99		
Atopobium minutum		28.37	0.51		1		
Clostridium bifermentans		16.48	0.39		99		
41200021	**Very good identification**						**41200021**
					SPO		
Clostridium novyi biovar A		98.97	0.59	SAL 8	99		
Atopobium minutum		0.65	0.17		1		
Clostridium bifermentans		0.38	0.05		99		
41200022	**Low discrimination**						**41200022**
					GRA	SPO	
Atopobium minutum		45.18	0.48	SAL 1	99	1	
Pseudoflavonifractor capillosus		32.18	0.32		1	1	
Clostridium novyi biovar A		18.27	0.46		99	99	
41200023	**Good identification**						**41200023**
					GRA	SPO	
Clostridium novyi biovar A		95.90	0.46	SAL 8	99	99	
Atopobium minutum		2.40	0.14		99	1	
Pseudoflavonifractor capillosus		1.71	-.03		1	1	
41200030	**Very good identification**						**41200030**
					SPO		
Clostridium novyi biovar A		98.67	0.59	SAL 8	99		
Atopobium minutum		0.65	0.17		1		
Clostridium bifermentans		0.38	0.05		99		
41200031	**Very good identification**						**41200031**
Clostridium novyi biovar A		100.0	0.59	SAL 8			
41200032	**Good identification**						**41200032**
					GRA	SPO	
Clostridium novyi biovar A		95.90	0.46	SAL 8	99	99	
Atopobium minutum		2.40	0.14		99	1	
Pseudoflavonifractor capillosus		1.71	-.03		1	1	
41200033	**Good identification**						**41200033**
Clostridium novyi biovar A		100.0	0.46	SAL 8			
41200040	**Low discrimination**						**41200040**
Clostridium botulinum biovar A		55.22	0.57	SAL 1			
Clostridium novyi biovar A		38.30	0.66				
Clostridium bifermentans		3.56	0.36				
41200041	**Very good identification**						**41200041**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		98.47	0.66	SAL 8			
Clostridium botulinum biovar A		1.43	0.23				
Clostridium bifermentans		0.09	0.02				
41200042	**Low discrimination**						**41200042**
Clostridium novyi biovar A		84.48	0.53	SAL 8			
Clostridium botulinum biovar A		6.97	0.23				
Clostridium sporogenes		5.72	0.19				
41200043	**Good identification**						**41200043**
Clostridium novyi biovar A		100.0	0.53	SAL 8			
41200050	**Very good identification**						**41200050**
Clostridium novyi biovar A		97.90	0.66	SAL 8			
Clostridium botulinum biovar A		1.43	0.23				
Clostridium difficile		0.58	0.13				
41200051	**Very good identification**						**41200051**
Clostridium novyi biovar A		100.0	0.66	SAL 8			
41200052	**Good identification**						**41200052**
Clostridium novyi biovar A		100.0	0.53	SAL 8			
41200053	**Good identification**						**41200053**
Clostridium novyi biovar A		100.0	0.53	SAL 8			
41200060	**Acceptable identification**						**41200060**
Clostridium novyi biovar A		91.58	0.56	SAL 8			
Clostridium botulinum biovar A		5.34	0.23				
Clostridium bifermentans		2.18	0.16				
41200061	**Good identification**						**41200061**
Clostridium novyi biovar A		100.0	0.56	SAL 8			
41200062	**Acceptable identification**						**41200062**
					GRA	SPO	
Clostridium novyi biovar A		93.97	0.43	SAL 8	99	99	
Atopobium minutum		3.52	0.14		99	1	
Pseudoflavonifractor capillosus		2.51	-.03		1	1	
41200063	**Good identification**						**41200063**
Clostridium novyi biovar A		100.0	0.43	SAL 8			
41200070	**Good identification**						**41200070**
Clostridium novyi biovar A		100.0	0.56	SAL 8			
41200071	**Good identification**						**41200071**
Clostridium novyi biovar A		100.0	0.56	SAL 8			
41200072	**Good identification**						**41200072**
Clostridium novyi biovar A		100.0	0.43	SAL 8			
41200073	**Good identification**						**41200073**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.43	SAL 8		
41201000	**Good identification**					**41201000**
					SPO	
Eubacterium tortuosum		93.56	0.66	SUC 99	1	
Atopobium minutum		2.95	0.51		1	
Clostridium novyi biovar A		2.00	0.53		99	
41201001	**Low discrimination**					**41201001**
					SPO	
Clostridium novyi biovar A		67.23	0.53	SAL 8	99	
Eubacterium tortuosum		31.77	0.32	GAL 10	1	
Atopobium minutum		1.00	0.17		1	
41201002	**Low discrimination**					**41201002**
					SPO	
Eubacterium tortuosum		86.93	0.53	SUC 99	1	
Atopobium minutum		10.34	0.48		1	
Clostridium novyi biovar A		1.86	0.40		99	
41201003	**Low discrimination**					**41201003**
					SPO	
Clostridium novyi biovar A		65.42	0.40	SAL 8	99	
Eubacterium tortuosum		30.91	0.19	GAL 10	1	
Atopobium minutum		3.68	0.14		1	
41201010	**Low discrimination**					**41201010**
					SPO	
Clostridium novyi biovar A		66.93	0.53	SAL 8	99	
Eubacterium tortuosum		31.62	0.32	GAL 10	1	
Atopobium minutum		1.00	0.17		1	
41201011	**Good identification**					**41201011**
					SPO	
Clostridium novyi biovar A		99.53	0.53	SAL 8	99	
Eubacterium tortuosum		0.48	-.03	GAL 10	1	
41201012	**Low discrimination**					**41201012**
					SPO	
Clostridium novyi biovar A		65.42	0.40	SAL 8	99	
Eubacterium tortuosum		30.91	0.19	GAL 10	1	
Atopobium minutum		3.68	0.14		1	
41201013	**Good identification**					**41201013**
Clostridium novyi biovar A		100.0	0.40	SAL 8		
41201020	**Acceptable identification**					**41201020**
					SPO	
Eubacterium tortuosum		91.68	0.53	SUC 99	1	
Atopobium minutum		5.45	0.43		1	
Clostridium novyi biovar A		2.78	0.42		99	
41201021	**Low discrimination**					**41201021**
					SPO	
Clostridium novyi biovar A		73.89	0.42	SAL 8	99	
Eubacterium tortuosum		24.65	0.19	GAL 10	1	
Atopobium minutum		1.47	0.09		1	
41201022	**Low discrimination**					**41201022**
					GRA	
Eubacterium tortuosum		76.44	0.40	SUC 99	99	
Atopobium minutum		17.17	0.40		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Pseudoflavonifractor capillosus		4.08	0.15		1	
41201030	**Low discrimination**					**41201030**
					SPO	
Clostridium novyi biovar A		73.89	0.42	SAL 8	99	
Eubacterium tortuosum		24.65	0.19	GAL 10	1	
Atopobium minutum		1.47	0.09		1	
41201031	**Good identification**					**41201031**
Clostridium novyi biovar A		100.0	0.42	SAL 8		
41201040	**Low discrimination**					**41201040**
					SPO	
Clostridium novyi biovar A		52.61	0.50	SAL 8	99	
Eubacterium tortuosum		37.29	0.32	GAL 10	1	
Clostridium botulinum biovar A		6.90	0.23		99	
41201041	**Good identification**					**41201041**
					SPO	
Clostridium novyi biovar A		99.29	0.50	SAL 8	99	
Eubacterium tortuosum		0.71	-.03	GAL 10	1	
41201042	**Low discrimination**					**41201042**
					SPO	
Clostridium novyi biovar A		55.77	0.37	SAL 8	99	
Eubacterium tortuosum		39.53	0.19	GAL 10	1	
Atopobium minutum		4.70	0.14		1	
41201043	**Good identification**					**41201043**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41201050	**Good identification**					**41201050**
					SPO	
Clostridium novyi biovar A		99.29	0.50	SAL 8	99	
Eubacterium tortuosum		0.71	-.03	GAL 10	1	
41201051	**Good identification**					**41201051**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
41201052	**Good identification**					**41201052**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41201053	**Good identification**					**41201053**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41201060	**Low discrimination**					**41201060**
					SPO	
Clostridium novyi biovar A		65.36	0.39	SAL 8	99	
Eubacterium tortuosum		32.70	0.19	GAL 10	1	
Atopobium minutum		1.94	0.09		1	
41201061	**Good identification**					**41201061**
Clostridium novyi biovar A		100.0	0.39	SAL 8		
41201070	**Good identification**					**41201070**
Clostridium novyi biovar A		100.0	0.39	SAL 8		
41201071	**Good identification**					**41201071**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41202000	**Low discrimination**						**41202000**
					SPO		
Clostridium difficile		51.87	0.40	FRU 99	99		
Eubacterium limosum		14.88	0.29		1		
Clostridium botulinum biovar A		12.21	0.32		99		
41202010	**Good identification**						**41202010**
					SPO		
Clostridium difficile		98.47	0.52	FRU 99	99		
Clostridium novyi biovar A		1.48	0.35		99		
Eubacterium limosum		0.06	-.06		1		
41204040	**Good identification**						**41204040**
Clostridium glycolicum		95.72	0.49	SAL 1			
Clostridium botulinum biovar A		2.44	0.23	FRU 90			
Clostridium novyi biovar A		1.69	0.32				
41210000	**Low discrimination**						**41210000**
					GRA	SPO	
Clostridium difficile		35.66	0.58	MAN 90	99	99	
Fusobacterium mortiferum		17.13	0.68		1	1	
Eubacterium limosum		14.98	0.49		99	1	
41210001	**Low discrimination**						**41210001**
					GRA	SPO	
Clostridium novyi biovar A		72.59	0.53	SAL 8	99	99	
Clostridium difficile		10.02	0.23	FRU 10	99	99	
Fusobacterium mortiferum		4.81	0.34		1	1	
41210002	**Low discrimination**						**41210002**
					GRA	SPO	
Atopobium minutum		52.44	0.56	SAL 1	99	1	
Clostridium sporogenes		24.17	0.40		99	99	
Fusobacterium mortiferum		14.07	0.52		1	1	
41210003	**Low discrimination**						**41210003**
					SPO		
Clostridium novyi biovar A		78.80	0.40	SAL 8	99		
Atopobium minutum		12.26	0.22	FRU 10	1		
Clostridium sporogenes		5.65	0.06		99		
41210010	**Very good identification**						**41210010**
					GRA	SPO	
Clostridium difficile		98.18	0.69	MAN 90	99	99	
Clostridium novyi biovar A		1.47	0.53		99	99	
Fusobacterium mortiferum		0.10	0.34		1	1	
41210011	**Low discrimination**						**41210011**
Clostridium novyi biovar A		59.74	0.53	SAL 8			
Clostridium difficile		40.27	0.35	FRU 10			
41210012	**Low discrimination**						**41210012**
					SPO		
Clostridium difficile		75.06	0.35	MAN 90	99		
Clostridium novyi biovar A		19.65	0.40	BGL 1	99		
Atopobium minutum		3.06	0.22		1		
41210013	**Good identification**						**41210013**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		96.29	0.40	SAL 8			
Clostridium difficile		3.72	0.01	FRU 10			
41210020	**Low discrimination**						**41210020**
					SPO		
Atopobium minutum		64.59	0.51	SAL 1	1		
Clostridium novyi biovar A		11.88	0.42		99		
Clostridium difficile		6.56	0.23		99		
41210021	**Acceptable identification**						**41210021**
					SPO		
Clostridium novyi biovar A		94.79	0.42	SAL 8	99		
Atopobium minutum		5.21	0.17	FRU 10	1		
41210022	**Low discrimination**						**41210022**
					GRA	SPO	
Atopobium minutum		85.36	0.48	SAL 1	99	1	
Pseudoflavonifractor capillosus		7.32	0.15		1	1	
Clostridium novyi biovar A		4.16	0.29		99	99	
41210030	**Low discrimination**						**41210030**
					SPO		
Clostridium difficile		71.88	0.35	MAN 90	99		
Clostridium novyi biovar A		26.66	0.42	NAG 1	99		
Atopobium minutum		1.46	0.17		1		
41210031	**Good identification**						**41210031**
Clostridium novyi biovar A		97.35	0.42	SAL 8			
Clostridium difficile		2.65	0.01	FRU 10			
41210040	**Low discrimination**						**41210040**
Clostridium glycolicum		40.47	0.43	SAL 1			
Clostridium difficile		17.90	0.36	XYL 95			
Clostridium novyi biovar A		16.58	0.50				
41210041	**Good identification**						**41210041**
Clostridium novyi biovar A		95.79	0.50	SAL 8			
Clostridium glycolicum		2.36	0.09	FRU 10			
Clostridium difficile		1.04	0.02				
41210042	**Low discrimination**						**41210042**
					SPO		
Clostridium novyi biovar A		53.70	0.37	SAL 8	99		
Atopobium minutum		12.53	0.22	FRU 10	1		
Pseudoramibacter alactolyticus		8.80	0.05		1		
41210043	**Good identification**						**41210043**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
41210050	**Low discrimination**						**41210050**
Clostridium difficile		83.61	0.47	MAN 90			
Clostridium novyi biovar A		15.87	0.50	SOR 5			
Clostridium glycolicum		0.39	0.09				
41210051	**Acceptable identification**						**41210051**
Clostridium novyi biovar A		94.95	0.50	SAL 8			
Clostridium difficile		5.05	0.13	FRU 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41210052	**Low discrimination**						**41210052**
Clostridium novyi biovar A		76.83	0.37	SAL 8			
Clostridium difficile		23.17	0.13	FRU 10			
41210053	**Good identification**						**41210053**
Clostridium novyi biovar A		100.0	0.37	SAL 8			
41210060	**Low discrimination**						**41210060**
					SPO		
Clostridium novyi biovar A		69.36	0.39	SAL 8	99		
Pseudoramibacter alactolyticus		12.74	0.09	FRU 10	1		
Clostridium glycolicum		6.84	0.09		99		
41210061	**Good identification**						**41210061**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41210070	**Low discrimination**						**41210070**
Clostridium novyi biovar A		82.45	0.39	SAL 8			
Clostridium difficile		17.55	0.13	FRU 10			
41210071	**Good identification**						**41210071**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41210100	**Very good identification**						**41210100**
					GRA	SPO	
Fusobacterium mortiferum		97.94	0.74		1	1	
Clostridium difficile		0.88	0.23		99	99	
Eubacterium limosum		0.37	0.15		99	1	
41210101	**Acceptable identification**						**41210101**
					GRA	SPO	
Fusobacterium mortiferum		93.87	0.40	URE 1	1	1	
Clostridium novyi biovar A		6.13	0.19		99	99	
41210102	**Very good identification**						**41210102**
					GRA	SPO	
Fusobacterium mortiferum		97.70	0.58	BGL 10	1	1	
Atopobium minutum		1.58	0.22		99	1	
Clostridium sporogenes		0.73	0.06		99	99	
41210110	**Low discrimination**						**41210110**
					GRA	SPO	
Clostridium difficile		80.35	0.35	MAN 90	99	99	
Fusobacterium mortiferum		18.45	0.40	LAC 1	1	1	
Clostridium novyi biovar A		1.20	0.19		99	99	
41210120	**Low discrimination**						**41210120**
					GRA	SPO	
Fusobacterium mortiferum		89.98	0.40	NAG 1	1	1	
Atopobium minutum		7.98	0.17		99	1	
Clostridium septicum		2.04	-.06		99	99	
41210140	**Low discrimination**						**41210140**
					GRA	SPO	
Fusobacterium mortiferum		81.12	0.40	SOR 1	1	1	
Clostridium glycolicum		8.61	0.09		99	99	
Clostridium difficile		3.81	0.02		99	99	
41210210	**Good identification**						**41210210**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		98.52	0.35	MAN 90			
Clostridium novyi biovar A		1.48	0.19	RHA 1			
41210300	**Good identification**						**41210300**
Fusobacterium mortiferum		100.0	0.40	RHA 1			
41210410	**Good identification**						**41210410**
Clostridium difficile		98.52	0.35	MAN 90			
Clostridium novyi biovar A		1.48	0.19	ARA 1			
41210500	**Good identification**						**41210500**
Fusobacterium mortiferum		100.0	0.40	ARA 1			
41211000	**Low discrimination**						**41211000**
					GRA		
Fusobacterium mortiferum		81.61	0.71		1		
Atopobium minutum		11.26	0.51		99		
Eubacterium tortuosum		3.92	0.32		99		
41211001	**Low discrimination**						**41211001**
					GRA	SPO	
Clostridium novyi biovar A		48.50	0.36	SAL 8	99	99	
Fusobacterium mortiferum		43.43	0.37	FRU 10	1	1	
Atopobium minutum		5.99	0.17	GAL 10	99	1	
41211002	**Low discrimination**						**41211002**
					GRA		
Fusobacterium mortiferum		51.71	0.55	BGL 10	1		
Atopobium minutum		42.82	0.48		99		
Eubacterium tortuosum		3.94	0.19		99		
41211010	**Low discrimination**						**41211010**
					GRA	SPO	
Clostridium difficile		74.63	0.35	GAL 1	99	99	
Clostridium novyi biovar A		12.30	0.36	MAN 90	99	99	
Fusobacterium mortiferum		11.02	0.37		1	1	
41211011	**Acceptable identification**						**41211011**
Clostridium novyi biovar A		94.23	0.36	SAL 8			
Clostridium difficile		5.77	0.01	FRU 10			
41211020	**Low discrimination**						**41211020**
					GRA		
Atopobium minutum		67.90	0.43	SAL 1	99		
Fusobacterium mortiferum		14.91	0.37		1		
Eubacterium tortuosum		12.50	0.19		99		
41211022	**Low discrimination**						**41211022**
					GRA		
Atopobium minutum		88.49	0.40	SAL 1	99		
Eubacterium tortuosum		4.31	0.06		99		
Fusobacterium mortiferum		3.24	0.20		1		
41211100	**Very good identification**						**41211100**
					GRA		
Fusobacterium mortiferum		99.88	0.77		1		
Atopobium minutum		0.06	0.17		99		
Actinomyces meyeri		0.03	0.07		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
41211101	**Good identification**						**41211101**
Fusobacterium mortiferum		100.0	0.43	URE 1			
41211102	**Very good identification**						**41211102**
					GRA		
Fusobacterium mortiferum		99.64	0.61	BGL 10	1		
Atopobium minutum		0.36	0.14		99		
41211110	**Good identification**						**41211110**
					GRA	SPO	
Fusobacterium mortiferum		97.15	0.43	MLZ 1	1	1	
Clostridium difficile		2.85	0.01		99	99	
41211140	**Good identification**						**41211140**
Fusobacterium mortiferum		100.0	0.43	SOR 1			
41211200	**Low discrimination**						**41211200**
					GRA		
Fusobacterium mortiferum		84.32	0.37	RHA 1	1		
Atopobium minutum		11.64	0.17		99		
Eubacterium tortuosum		4.05	-.03		99		
41211300	**Good identification**						**41211300**
Fusobacterium mortiferum		100.0	0.43	RHA 1			
41211400	**Low discrimination**						**41211400**
					GRA		
Fusobacterium mortiferum		84.32	0.37	ARA 1	1		
Atopobium minutum		11.64	0.17		99		
Eubacterium tortuosum		4.05	-.03		99		
41211500	**Good identification**						**41211500**
Fusobacterium mortiferum		100.0	0.43	ARA 1			
41212000	**Low discrimination**						**41212000**
					SPO		
Clostridium difficile		77.41	0.74		99		
Eubacterium limosum		22.20	0.63		1		
Pseudoramibacter alactolyticus		0.24	0.27		1		
41212001	**Low discrimination**						**41212001**
					SPO		
Clostridium difficile		77.04	0.40	URE 1	99		
Eubacterium limosum		22.10	0.29		1		
Clostridium novyi biovar A		0.63	0.19		99		
41212002	**Low discrimination**						**41212002**
					SPO		
Clostridium difficile		73.26	0.40	BGL 1	99		
Eubacterium limosum		21.01	0.29		1		
Pseudoramibacter alactolyticus		2.45	0.11		1		
41212010	**Very good identification**						**41212010**
					SPO		
Clostridium difficile		99.94	0.86		99		
Eubacterium limosum		0.06	0.29		1		
Clostridium novyi biovar A		0.00	0.19		99		
41212011	**Good identification**						**41212011**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		99.78	0.52	URE 1	99		
Clostridium novyi biovar A		0.17	0.19		99		
Eubacterium limosum		0.06	-.06		1		
41212012	**Good identification**						**41212012**
					SPO		
Clostridium difficile		99.94	0.52	BGL 1	99		
Eubacterium limosum		0.06	-.06		1		
41212020	**Low discrimination**						**41212020**
					SPO		
Clostridium difficile		74.03	0.40	NAG 1	99		
Eubacterium limosum		21.23	0.29		1		
Pseudoramibacter alactolyticus		3.93	0.14		1		
41212030	**Good identification**						**41212030**
					SPO		
Clostridium difficile		99.94	0.52	NAG 1	99		
Eubacterium limosum		0.06	-.06		1		
41212040	**Acceptable identification**						**41212040**
					SPO		
Clostridium difficile		92.47	0.52	SOR 5	99		
Eubacterium limosum		5.09	0.29		1		
Pseudoramibacter alactolyticus		1.33	0.17		1		
41212050	**Very good identification**						**41212050**
					SPO		
Clostridium difficile		99.97	0.64	SOR 5	99		
Clostridium novyi biovar A		0.02	0.15		99		
Eubacterium limosum		0.01	-.06		1		
41212100	**Low discrimination**						**41212100**
					GRA	SPO	
Clostridium difficile		70.70	0.40	LAC 1	99	99	
Eubacterium limosum		20.28	0.29		99	1	
Fusobacterium mortiferum		8.81	0.40		1	1	
41212110	**Good identification**						**41212110**
					SPO		
Clostridium difficile		99.94	0.52	LAC 1	99		
Eubacterium limosum		0.06	-.06		1		
41212200	**Low discrimination**						**41212200**
					SPO		
Clostridium difficile		77.53	0.40	RHA 1	99		
Eubacterium limosum		22.24	0.29		1		
Pseudoramibacter alactolyticus		0.24	-.07		1		
41212210	**Good identification**						**41212210**
					SPO		
Clostridium difficile		99.94	0.52	RHA 1	99		
Eubacterium limosum		0.06	-.06		1		
41212400	**Low discrimination**						**41212400**
					SPO		
Clostridium difficile		77.53	0.40	ARA 1	99		
Eubacterium limosum		22.24	0.29		1		
Pseudoramibacter alactolyticus		0.24	-.07		1		
41212410	**Good identification**						**41212410**
					SPO		
Clostridium difficile		99.94	0.52	ARA 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		0.06	-.06		1		
41213000	**Low discrimination**						**41213000**
					GRA	SPO	
Clostridium difficile		71.06	0.40	GAL 1	99	99	
Eubacterium limosum		20.38	0.29		99	1	
Fusobacterium mortiferum		5.69	0.37		1	1	
41213010	**Good identification**						**41213010**
					SPO		
Clostridium difficile		99.94	0.52	GAL 1	99		
Eubacterium limosum		0.06	-.06		1		
41213100	**Acceptable identification**						**41213100**
					GRA	SPO	
Fusobacterium mortiferum		93.50	0.43	MAN 1	1	1	
Clostridium difficile		5.06	0.06		99	99	
Eubacterium limosum		1.45	-.06		99	1	
41214000	**Low discrimination**						**41214000**
					GRA	SPO	
Clostridium glycolicum		53.79	0.43	SAL 1	99	99	
Clostridium difficile		23.79	0.36	SOR 95	99	99	
Fusobacterium mortiferum		16.34	0.49		1	1	
41214010	**Good identification**						**41214010**
Clostridium difficile		99.11	0.47	MAN 90			
Clostridium glycolicum		0.46	0.09	XYL 5			
Clostridium novyi biovar A		0.29	0.19				
41214040	**Very good identification**						**41214040**
Clostridium glycolicum		99.82	0.65	SAL 1			
Clostridium difficile		0.12	0.14				
Clostridium novyi biovar A		0.02	0.15				
41214100	**Good identification**						**41214100**
					GRA	SPO	
Fusobacterium mortiferum		97.34	0.55	XYL 7	1	1	
Clostridium glycolicum		1.39	0.09		99	99	
Actinomyces meyeri		0.66	0.09		99	1	
41214102	**Good identification**						**41214102**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
41215000	**Low discrimination**						**41215000**
					GRA	SPO	
Fusobacterium mortiferum		78.21	0.52	XYL 7	1	1	
Clostridium glycolicum		19.07	0.27		99	99	
Atopobium minutum		1.45	0.17		99	1	
41215002	**Acceptable identification**						**41215002**
					GRA		
Fusobacterium mortiferum		90.00	0.35	XYL 7	1		
Atopobium minutum		10.00	0.14	BGL 10	99		
41215040	**Good identification**						**41215040**
					GRA	SPO	
Clostridium glycolicum		99.78	0.49	SAL 1	99	99	
Fusobacterium mortiferum		0.22	0.18	GAL 10	1	1	
41215100	**Very good identification**						**41215100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		99.33	0.58	XYL 7	1		
Actinomyces meyeri		0.67	0.12		99		
41215102	**Good identification**						**41215102**
Fusobacterium mortiferum		100.0	0.42	XYL 7			
41216000	**Acceptable identification**						**41216000**
					SPO		
Clostridium difficile		94.44	0.52	XYL 5	99		
Eubacterium limosum		5.20	0.29		1		
Clostridium glycolicum		0.24	0.09		99		
41216010	**Very good identification**						**41216010**
					SPO		
Clostridium difficile		99.99	0.64	XYL 5	99		
Eubacterium limosum		0.01	-.06		1		
41216050	**Good identification**						**41216050**
Clostridium difficile		99.81	0.42	XYL 5			
Clostridium glycolicum		0.19	-.03	SOR 5			
41220000	**Low discrimination**						**41220000**
Clostridium botulinum biovar A		49.42	0.56	SAL 1			
Clostridium sporogenes		48.46	0.53				
Clostridium bifermentans		0.92	0.25				
41220002	**Good identification**						**41220002**
					SPO		
Clostridium sporogenes		95.22	0.45	SAL 1	99		
Clostridium botulinum biovar A		2.94	0.21		99		
Atopobium minutum		1.20	0.22		1		
41220040	**Acceptable identification**						**41220040**
Clostridium botulinum biovar A		94.00	0.47	SAL 1			
Clostridium sporogenes		3.12	0.19				
Clostridium novyi biovar A		2.63	0.32				
41230000	**Low discrimination**						**41230000**
					GRA	SPO	
Fusobacterium mortiferum		40.62	0.59		1	1	
Clostridium sporogenes		25.95	0.40		99	99	
Clostridium difficile		21.31	0.38		99	99	
41230010	**Good identification**						**41230010**
					GRA	SPO	
Clostridium difficile		99.07	0.50	TRE 7	99	99	
Fusobacterium mortiferum		0.39	0.25	MAN 90	1	1	
Clostridium sporogenes		0.25	0.06		99	99	
41230100	**Very good identification**						**41230100**
					GRA	SPO	
Fusobacterium mortiferum		99.40	0.65		1	1	
Clostridium sporogenes		0.28	0.06		99	99	
Clostridium difficile		0.23	0.04		99	99	
41230102	**Good identification**						**41230102**
					GRA	SPO	
Fusobacterium mortiferum		99.18	0.49	BGL 10	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		0.82	-.02		99	99	
41231000	**Very good identification**						**41231000**
					GRA	SPO	
Fusobacterium mortiferum		98.26	0.62		1	1	
Atopobium minutum		0.46	0.17		99	1	
Clostridium sporogenes		0.42	0.06		99	99	
41231002	**Acceptable identification**						**41231002**
					GRA	SPO	
Fusobacterium mortiferum		94.99	0.46	BGL 10	1	1	
Atopobium minutum		2.66	0.14		99	1	
Clostridium sporogenes		1.23	-.02		99	99	
41231100	**Very good identification**						**41231100**
					GRA	SPO	
Fusobacterium mortiferum		99.85	0.68		1	1	
Clostridium septicum		0.15	0.03		99	99	
41231102	**Good identification**						**41231102**
Fusobacterium mortiferum		100.0	0.52	BGL 10			
41231120	**Acceptable identification**						**41231120**
					GRA	SPO	
Clostridium septicum		93.77	0.37	MNS 99	99	99	
Fusobacterium mortiferum		6.23	0.34	MLT 94	1	1	
41232000	**Good identification**						**41232000**
					GRA	SPO	
Clostridium difficile		95.89	0.55	TRE 7	99	99	
Eubacterium limosum		3.69	0.29		99	1	
Fusobacterium mortiferum		0.21	0.25		1	1	
41232010	**Very good identification**						**41232010**
					SPO		
Clostridium difficile		99.99	0.66	TRE 7	99		
Eubacterium limosum		0.01	-.06		1		
41232050	**Good identification**						**41232050**
Clostridium difficile		100.0	0.45	TRE 7			
41234000	**Low discrimination**						**41234000**
					GRA	SPO	
Fusobacterium mortiferum		62.68	0.40	XYL 7	1	1	
Clostridium difficile		22.99	0.16		99	99	
Clostridium glycolicum		6.98	0.09		99	99	
41234100	**Good identification**						**41234100**
Fusobacterium mortiferum		100.0	0.46	XYL 7			
41235000	**Good identification**						**41235000**
Fusobacterium mortiferum		100.0	0.43	XYL 7			
41235100	**Good identification**						**41235100**
					COC	GRA	
Fusobacterium mortiferum		99.20	0.49	XYL 7	1	1	
Atopobium parvulum		0.80	0.02		99	99	
41236010	**Good identification**						**41236010**
Clostridium difficile		100.0	0.45	TRE 7			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41241000	**Good identification**						**41241000**
					GRA		
Eubacterium tortuosum		98.89	0.50	SUC 99	99		
Fusobacterium mortiferum		0.63	0.29	CEL 10	1		
Atopobium minutum		0.28	0.17		99		
41241002	**Good identification**						**41241002**
					GRA		
Eubacterium tortuosum		98.54	0.37	SUC 99	99		
Atopobium minutum		1.07	0.14	CEL 10	99		
Fusobacterium mortiferum		0.40	0.12		1		
41241020	**Good identification**						**41241020**
Eubacterium tortuosum		99.46	0.37	SUC 99			
Atopobium minutum		0.54	0.09	CEL 10			
41250000	**Low discrimination**						**41250000**
					GRA	SPO	
Fusobacterium mortiferum		70.88	0.60		1	1	
Clostridium difficile		23.29	0.36		99	99	
Eubacterium limosum		1.88	0.15		99	1	
41250002	**Low discrimination**						**41250002**
					GRA	SPO	
Fusobacterium mortiferum		83.69	0.43	BGL 10	1	1	
Atopobium minutum		9.45	0.22		99	1	
Clostridium sporogenes		4.36	0.06		99	99	
41250010	**Good identification**						**41250010**
					GRA	SPO	
Clostridium difficile		99.09	0.47	CEL 5	99	99	
Fusobacterium mortiferum		0.62	0.26	MAN 90	1	1	
Clostridium novyi biovar A		0.29	0.19		99	99	
41250100	**Very good identification**						**41250100**
					GRA	SPO	
Fusobacterium mortiferum		99.86	0.66		1	1	
Clostridium difficile		0.14	0.02		99	99	
41250102	**Good identification**						**41250102**
Fusobacterium mortiferum		100.0	0.50	BGL 10			
41251000	**Very good identification**						**41251000**
					GRA		
Fusobacterium mortiferum		95.57	0.63		1		
Eubacterium tortuosum		1.53	0.15		99		
Lactobacillus jensenii		1.07	0.19		99		
41251002	**Low discrimination**						**41251002**
					GRA	SPO	
Fusobacterium mortiferum		79.56	0.46	BGL 10	1	1	
Clostridium innocuum		6.09	0.15		99	99	
Lactobacillus jensenii		5.36	0.16		99	1	
41251100	**Very good identification**						**41251100**
					GRA		
Fusobacterium mortiferum		99.29	0.69		1		
Lactobacillus catenaformis		0.27	0.15		99		
Collinsella aerofaciens		0.18	-.03		99		
41251102	**Acceptable identification**						**41251102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		94.73	0.53	BGL 10	1	1	
Clostridium barati		2.92	0.12		99	99	
Lactobacillus catenaformis		2.35	0.15		99	1	
41251120	**Low discrimination**						**41251120**
					GRA	SPO	
Clostridium septicum		86.21	0.37	MNS 99	99	99	
Fusobacterium mortiferum		6.05	0.35	MLT 94	1	1	
Clostridium paraputrificum		5.27	0.14		99	99	
41252000	**Acceptable identification**						**41252000**
					SPO		
Clostridium difficile		92.46	0.52	CEL 5	99		
Eubacterium limosum		5.09	0.29		1		
Clostridium innocuum		2.08	0.27		99		
41252010	**Very good identification**						**41252010**
					SPO		
Clostridium difficile		99.97	0.64	CEL 5	99		
Clostridium innocuum		0.02	0.04		99		
Eubacterium limosum		0.01	-.06		1		
41252050	**Good identification**						**41252050**
Clostridium difficile		100.0	0.42	CEL 5			
41253000	**Good identification**						**41253000**
					GRA	SPO	
Clostridium innocuum		96.44	0.49	MNS 99	99	99	
Clostridium difficile		2.28	0.18		99	99	
Fusobacterium mortiferum		1.16	0.29		1	1	
41253002	**Good identification**						**41253002**
					GRA	SPO	
Clostridium innocuum		99.87	0.49	MNS 99	99	99	
Fusobacterium mortiferum		0.13	0.12		1	1	
41253400	**Good identification**						**41253400**
Clostridium innocuum		100.0	0.36	MNS 99			
41253402	**Good identification**						**41253402**
Clostridium innocuum		100.0	0.36	MNS 99			
41254000	**Low discrimination**						**41254000**
					GRA	SPO	
Fusobacterium mortiferum		75.21	0.41	XYL 7	1	1	
Clostridium difficile		17.29	0.14		99	99	
Clostridium glycolicum		7.50	0.09		99	99	
41254100	**Good identification**						**41254100**
Fusobacterium mortiferum		100.0	0.47	XYL 7			
41255000	**Good identification**						**41255000**
Fusobacterium mortiferum		100.0	0.44	XYL 7			
41255100	**Good identification**						**41255100**
Fusobacterium mortiferum		100.0	0.50	XYL 7			
41256010	**Good identification**						**41256010**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		100.0	0.42	CEL 5				
41270000	**Acceptable identification**							**41270000**
					GRA	SPO		
Fusobacterium mortiferum		90.15	0.51		1	1		
Clostridium difficile		7.47	0.16		99	99		
Clostridium sporogenes		1.75	0.06		99	99		
41270100	**Very good identification**							**41270100**
Fusobacterium mortiferum		100.0	0.57					
41270102	**Good identification**							**41270102**
Fusobacterium mortiferum		100.0	0.41	BGL 10				
41271000	**Low discrimination**							**41271000**
					GRA	SPO		
Fusobacterium mortiferum		89.07	0.54		1	1		
Lactobacillus jensenii		10.05	0.28		99	1		
Clostridium innocuum		0.89	0.07		99	99		
41271002	**Low discrimination**							**41271002**
					GRA	SPO		
Fusobacterium mortiferum		56.60	0.38	BGL 10	1	1		
Lactobacillus jensenii		38.30	0.25		99	1		
Clostridium innocuum		5.09	0.07		99	99		
41271100	**Very good identification**							**41271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.34	0.60		1	1	1	
Clostridium septicum		1.44	0.11		1	99	99	
Atopobium parvulum		0.22	0.04		99	99	1	
41271102	**Good identification**							**41271102**
					GRA			
Fusobacterium mortiferum		97.05	0.44	BGL 10	1			
Lactobacillus acidophilus		2.95	0.05		99			
41271120	**Good identification**							**41271120**
					GRA	SPO		
Clostridium septicum		98.77	0.45	MNS 99	99	99		
Fusobacterium mortiferum		0.69	0.26	MLT 94	1	1		
Clostridium paraputrificum		0.54	0.04		99	99		
41272010	**Good identification**							**41272010**
Clostridium difficile		100.0	0.45	TRE 7				
41273000	**Good identification**							**41273000**
					GRA	SPO		
Clostridium innocuum		98.50	0.41	MNS 99	99	99		
Fusobacterium mortiferum		1.01	0.20		1	1		
Clostridium difficile		0.50	-.01		99	99		
41273002	**Good identification**							**41273002**
Clostridium innocuum		100.0	0.41	MNS 99				
41274100	**Good identification**							**41274100**
Fusobacterium mortiferum		100.0	0.38	XYL 7				
41275100	**Acceptable identification**							**41275100**

Code Taxon		ID score	T index	tests against	Differentiation tests COC	 GRA	Code
Fusobacterium mortiferum		94.67	0.41	XYL 7	1	1	
Atopobium parvulum		5.33	0.08		99	99	
41300000	**Low discrimination**						**41300000**
Clostridium sporogenes		37.98	0.66	SAL 1			
Clostridium bifermentans		30.01	0.66				
Clostridium botulinum biovar A		22.75	0.64				
41300001	**Good identification**						**41300001**
Clostridium novyi biovar A		90.10	0.69	SAL 8			
Clostridium sporogenes		4.00	0.32				
Clostridium bifermentans		3.16	0.32				
41300002	**Low discrimination**						**41300002**
Clostridium sporogenes		85.83	0.58	SAL 1			
Clostridium novyi biovar A		10.34	0.56				
Clostridium bifermentans		2.06	0.32				
41300003	**Good identification**						**41300003**
Clostridium novyi biovar A		92.09	0.56	SAL 8			
Clostridium sporogenes		7.72	0.24				
Clostridium bifermentans		0.19	-.03				
41300010	**Good identification**						**41300010**
Clostridium novyi biovar A		90.32	0.69	SAL 8			
Clostridium sporogenes		4.01	0.32				
Clostridium bifermentans		3.17	0.32				
41300011	**Very good identification**						**41300011**
Clostridium novyi biovar A		99.92	0.69	SAL 8			
Clostridium sporogenes		0.05	-.03				
Clostridium bifermentans		0.04	-.03				
41300012	**Good identification**						**41300012**
Clostridium novyi biovar A		92.09	0.56	SAL 8			
Clostridium sporogenes		7.72	0.24				
Clostridium bifermentans		0.19	-.03				
41300013	**Very good identification**						**41300013**
Clostridium novyi biovar A		100.0	0.56	SAL 8			
41300020	**Low discrimination**						**41300020**
Clostridium novyi biovar A		45.89	0.59	SAL 8			
Clostridium bifermentans		40.70	0.45				
Clostridium sporogenes		8.15	0.32				
41300021	**Very good identification**						**41300021**
Clostridium novyi biovar A		98.94	0.59	SAL 8			
Clostridium bifermentans		0.89	0.11				
Clostridium sporogenes		0.18	-.03				
41300022	**Low discrimination**						**41300022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		69.98	0.46	SAL 8		
Clostridium sporogenes		23.47	0.24			
Clostridium bifermentans		3.55	0.11			
41300023	**Good identification**					**41300023**
Clostridium novyi biovar A		100.0	0.46	SAL 8		
41300030	**Very good identification**					**41300030**
Clostridium novyi biovar A		98.94	0.59	SAL 8		
Clostridium bifermentans		0.89	0.11			
Clostridium sporogenes		0.18	-.03			
41300031	**Very good identification**					**41300031**
Clostridium novyi biovar A		100.0	0.59	SAL 8		
41300032	**Good identification**					**41300032**
Clostridium novyi biovar A		100.0	0.46	SAL 8		
41300033	**Good identification**					**41300033**
Clostridium novyi biovar A		100.0	0.46	SAL 8		
41300040	**Low discrimination**					**41300040**
Clostridium botulinum biovar A		47.24	0.55	SAL 1		
Clostridium novyi biovar A		40.04	0.66			
Clostridium bifermentans		8.69	0.42			
41300041	**Very good identification**					**41300041**
Clostridium novyi biovar A		98.54	0.66	SAL 8		
Clostridium botulinum biovar A		1.17	0.21			
Clostridium bifermentans		0.22	0.08			
41300042	**Low discrimination**					**41300042**
Clostridium novyi biovar A		82.94	0.53	SAL 8		
Clostridium sporogenes		10.43	0.24			
Clostridium botulinum biovar A		5.60	0.21			
41300043	**Good identification**					**41300043**
Clostridium novyi biovar A		100.0	0.53	SAL 8		
41300050	**Very good identification**					**41300050**
Clostridium novyi biovar A		98.54	0.66	SAL 8		
Clostridium botulinum biovar A		1.17	0.21			
Clostridium bifermentans		0.22	0.08			
41300051	**Very good identification**					**41300051**
Clostridium novyi biovar A		100.0	0.66	SAL 8		
41300052	**Good identification**					**41300052**
Clostridium novyi biovar A		100.0	0.53	SAL 8		
41300053	**Good identification**					**41300053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.53	SAL 8		
41300060	**Acceptable identification**					**41300060**
Clostridium novyi biovar A		90.43	0.56	SAL 8		
Clostridium bifermentans		5.01	0.22			
Clostridium botulinum biovar A		4.31	0.21			
41300061	**Good identification**					**41300061**
Clostridium novyi biovar A		100.0	0.56	SAL 8		
41300062	**Good identification**					**41300062**
Clostridium novyi biovar A		100.0	0.43	SAL 8		
41300063	**Good identification**					**41300063**
Clostridium novyi biovar A		100.0	0.43	SAL 8		
41300070	**Good identification**					**41300070**
Clostridium novyi biovar A		100.0	0.56	SAL 8		
41300071	**Good identification**					**41300071**
Clostridium novyi biovar A		100.0	0.56	SAL 8		
41300072	**Good identification**					**41300072**
Clostridium novyi biovar A		100.0	0.43	SAL 8		
41300073	**Good identification**					**41300073**
Clostridium novyi biovar A		100.0	0.43	SAL 8		
41301000	**Low discrimination**					**41301000**
					SPO	
Clostridium novyi biovar A		40.60	0.53	SAL 8	99	
Eubacterium tortuosum		19.18	0.32	GAL 10	1	
Clostridium sporogenes		16.22	0.32		99	
41301001	**Good identification**					**41301001**
					SPO	
Clostridium novyi biovar A		98.30	0.53	SAL 8	99	
Clostridium sordellii		0.52	-.03	GAL 10	99	
Eubacterium tortuosum		0.47	-.03		1	
41301002	**Low discrimination**					**41301002**
					SPO	
Clostridium novyi biovar A		43.45	0.40	SAL 8	99	
Clostridium sporogenes		32.79	0.24	GAL 10	99	
Eubacterium tortuosum		20.53	0.19		1	
41301003	**Good identification**					**41301003**
Clostridium novyi biovar A		100.0	0.40	SAL 8		
41301010	**Good identification**					**41301010**
					SPO	
Clostridium novyi biovar A		98.81	0.53	SAL 8	99	
Eubacterium tortuosum		0.47	-.03	GAL 10	1	
Clostridium sporogenes		0.40	-.03		99	
41301011	**Good identification**					**41301011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.53	SAL 8		
41301012	**Good identification**					**41301012**
Clostridium novyi biovar A		100.0	0.40	SAL 8		
41301013	**Good identification**					**41301013**
Clostridium novyi biovar A		100.0	0.40	SAL 8		
41301020	**Low discrimination**					**41301020**
					SPO	
Clostridium novyi biovar A		68.96	0.42	SAL 8	99	
Eubacterium tortuosum		23.00	0.19	GAL 10	1	
Clostridium bifermentans		5.56	0.11		99	
41301021	**Good identification**					**41301021**
Clostridium novyi biovar A		100.0	0.42	SAL 8		
41301030	**Good identification**					**41301030**
Clostridium novyi biovar A		100.0	0.42	SAL 8		
41301031	**Good identification**					**41301031**
Clostridium novyi biovar A		100.0	0.42	SAL 8		
41301040	**Low discrimination**					**41301040**
Clostridium novyi biovar A		85.26	0.50	SAL 8		
Clostridium botulinum biovar A		9.14	0.21	GAL 10		
Clostridium glycolicum		2.78	0.11			
41301041	**Good identification**					**41301041**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
41301042	**Good identification**					**41301042**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41301043	**Good identification**					**41301043**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41301050	**Good identification**					**41301050**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
41301051	**Good identification**					**41301051**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
41301052	**Good identification**					**41301052**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41301053	**Good identification**					**41301053**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41301060	**Good identification**					**41301060**
Clostridium novyi biovar A		100.0	0.39	SAL 8		
41301061	**Good identification**					**41301061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41301070	**Good identification**						**41301070**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41301071	**Good identification**						**41301071**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41304040	**Good identification**						**41304040**
Clostridium glycolicum		96.14	0.50	SAL 1			
Clostridium botulinum biovar A		1.85	0.21	FRU 90			
Clostridium novyi biovar A		1.57	0.32				
41310000	**Low discrimination**						**41310000**
Clostridium sporogenes		55.37	0.53	SAL 1			
Clostridium bifermentans		13.05	0.44				
Clostridium botulinum biovar A		12.00	0.44				
41310001	**Low discrimination**						**41310001**
Clostridium novyi biovar A		87.67	0.53	SAL 8			
Clostridium sporogenes		6.18	0.19	FRU 10			
Clostridium sordellii		2.64	0.10				
41310002	**Low discrimination**						**41310002**
					GRA	SPO	
Clostridium sporogenes		87.31	0.45	SAL 1	99	99	
Clostridium novyi biovar A		6.62	0.40		99	99	
Fusobacterium mortiferum		3.38	0.36		1	1	
41310003	**Low discrimination**						**41310003**
Clostridium novyi biovar A		88.25	0.40	SAL 8			
Clostridium sporogenes		11.75	0.11	FRU 10			
41310010	**Low discrimination**						**41310010**
Clostridium novyi biovar A		56.03	0.53	SAL 8			
Clostridium difficile		37.77	0.35	FRU 10			
Clostridium sporogenes		3.95	0.19				
41310011	**Good identification**						**41310011**
Clostridium novyi biovar A		99.32	0.53	SAL 8			
Clostridium difficile		0.68	0.01	FRU 10			
41310012	**Low discrimination**						**41310012**
Clostridium novyi biovar A		85.34	0.40	SAL 8			
Clostridium sporogenes		11.37	0.11	FRU 10			
Clostridium difficile		3.29	0.01				
41310013	**Good identification**						**41310013**
Clostridium novyi biovar A		100.0	0.40	SAL 8			
41310020	**Low discrimination**						**41310020**
Clostridium novyi biovar A		54.03	0.42	SAL 8			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		22.69	0.24	FRU 10		
Clostridium sporogenes		15.24	0.19			
41310021	**Good identification**					**41310021**
Clostridium novyi biovar A		100.0	0.42	SAL 8		
41310030	**Good identification**					**41310030**
Clostridium novyi biovar A		97.35	0.42	SAL 8		
Clostridium difficile		2.65	0.01	FRU 10		
41310031	**Good identification**					**41310031**
Clostridium novyi biovar A		100.0	0.42	SAL 8		
41310040	**Low discrimination**					**41310040**
Clostridium glycolicum		56.70	0.44	SAL 1		
Clostridium novyi biovar A		21.45	0.50	XYL 95		
Clostridium botulinum biovar A		14.54	0.35			
41310041	**Good identification**					**41310041**
Clostridium novyi biovar A		96.75	0.50	SAL 8		
Clostridium glycolicum		2.58	0.10	FRU 10		
Clostridium botulinum biovar A		0.66	0.01			
41310042	**Low discrimination**					**41310042**
Clostridium novyi biovar A		68.55	0.37	SAL 8		
Clostridium sporogenes		13.69	0.11	FRU 10		
Clostridium glycolicum		10.37	0.10			
41310043	**Good identification**					**41310043**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41310050	**Acceptable identification**					**41310050**
Clostridium novyi biovar A		92.02	0.50	SAL 8		
Clostridium difficile		4.90	0.13	FRU 10		
Clostridium glycolicum		2.46	0.10			
41310051	**Good identification**					**41310051**
Clostridium novyi biovar A		100.0	0.50	SAL 8		
41310052	**Good identification**					**41310052**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41310053	**Good identification**					**41310053**
Clostridium novyi biovar A		100.0	0.37	SAL 8		
41310060	**Low discrimination**					**41310060**
Clostridium novyi biovar A		86.17	0.39	SAL 8		
Clostridium glycolicum		9.20	0.10	FRU 10		
Clostridium botulinum biovar A		2.36	0.01			
41310061	**Good identification**					**41310061**
Clostridium novyi biovar A		100.0	0.39	SAL 8		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41310070	**Good identification**						**41310070**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41310071	**Good identification**						**41310071**
Clostridium novyi biovar A		100.0	0.39	SAL 8			
41310100	**Good identification**						**41310100**
					GRA	SPO	
Fusobacterium mortiferum		94.18	0.59	MLT 11	1	1	
Clostridium sporogenes		3.51	0.19		99	99	
Clostridium bifermentans		0.83	0.10		99	99	
41310102	**Low discrimination**						**41310102**
					GRA	SPO	
Fusobacterium mortiferum		89.95	0.42	MLT 11	1	1	
Clostridium sporogenes		10.05	0.11	BGL 10	99	99	
41311000	**Low discrimination**						**41311000**
					GRA	SPO	
Fusobacterium mortiferum		81.03	0.55	MLT 11	1	1	
Clostridium novyi biovar A		7.40	0.36		99	99	
Clostridium sporogenes		4.69	0.19		99	99	
41311001	**Low discrimination**						**41311001**
					GRA	SPO	
Clostridium novyi biovar A		86.32	0.36	SAL 8	99	99	
Fusobacterium mortiferum		9.55	0.21	FRU 10	1	1	
Clostridium sordellii		4.13	-.03	GAL 10	99	99	
41311002	**Low discrimination**						**41311002**
					GRA	SPO	
Fusobacterium mortiferum		63.77	0.39	MLT 11	1	1	
Lactobacillus jensenii		11.59	0.16	BGL 10	99	1	
Clostridium sporogenes		11.08	0.11		99	99	
41311010	**Low discrimination**						**41311010**
					GRA	SPO	
Clostridium novyi biovar A		77.97	0.36	SAL 8	99	99	
Fusobacterium mortiferum		8.63	0.21	FRU 10	1	1	
Lactobacillus jensenii		8.62	0.11	GAL 10	99	1	
41311011	**Good identification**						**41311011**
Clostridium novyi biovar A		100.0	0.36	SAL 8			
41311100	**Very good identification**						**41311100**
					GRA	SPO	
Fusobacterium mortiferum		97.10	0.62	MLT 11	1	1	
Clostridium septicum		1.88	0.15		99	99	
Collinsella aerofaciens		0.48	-.03		99	1	
41311102	**Acceptable identification**						**41311102**
					GRA		
Fusobacterium mortiferum		93.17	0.45	MLT 11	1		
Bifidobacterium longum subsp. infantis		4.64	0.06	BGL 10	99		
Lactobacillus fermentum		1.10	0.14		99		
41311120	**Good identification**						**41311120**
Clostridium septicum		96.91	0.49	MNS 99			
Clostridium perfringens		1.24	0.23				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium paraputrificum		1.20	0.14					
41312000	**Low discrimination**							**41312000**
					COC	SPO	GRA	
Clostridium difficile		67.79	0.40	MLT 1	1	99	99	
Eubacterium limosum		19.44	0.29		1	1	99	
Megasphaera elsdenii		5.98	0.29		99	1	1	
41312010	**Good identification**							**41312010**
					SPO			
Clostridium difficile		99.78	0.52	MLT 1	99			
Clostridium novyi biovar A		0.17	0.19		99			
Eubacterium limosum		0.06	-.06		1			
41314000	**Acceptable identification**							**41314000**
					GRA	SPO		
Clostridium glycolicum		90.69	0.44	SAL 1	99	99		
Clostridium sporogenes		3.63	0.19	SOR 95	99	99		
Fusobacterium mortiferum		3.14	0.33		1	1		
41314040	**Very good identification**							**41314040**
Clostridium glycolicum		99.97	0.66	SAL 1				
Clostridium novyi biovar A		0.02	0.15					
Clostridium botulinum biovar A		0.01	0.01					
41314100	**Low discrimination**							**41314100**
					GRA	SPO		
Fusobacterium mortiferum		84.29	0.40	MLT 11	1	1		
Clostridium glycolicum		10.53	0.10	XYL 7	99	99		
Actinomyces meyeri		5.18	0.09		99	1		
41315040	**Good identification**							**41315040**
Clostridium glycolicum		100.0	0.50	SAL 1				
41315300	**Very good identification**							**41315300**
Eubacterium contortum		100.0	0.66	SUC 99				
41315302	**Good identification**							**41315302**
Eubacterium contortum		100.0	0.53	SUC 99				
41315320	**Good identification**							**41315320**
Eubacterium contortum		100.0	0.53	SUC 99				
41315322	**Good identification**							**41315322**
					GRA			
Eubacterium contortum		98.78	0.40	SUC 99	99			
Parabacteroides distasonis		0.79	0.15		1			
Prevotella oralis		0.44	-.02		1			
41315700	**Very good identification**							**41315700**
Eubacterium contortum		100.0	0.64	SUC 99				
41315702	**Good identification**							**41315702**
Eubacterium contortum		100.0	0.52	SUC 99				
41315720	**Good identification**							**41315720**
					GRA			
Eubacterium contortum		99.42	0.52	SUC 99	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides vulgatus		0.58	0.06		1			
41315722	**Good identification**							**41315722**
					GRA			
Eubacterium contortum		99.41	0.39	SUC 99	99			
Parabacteroides distasonis		0.59	0.12		1			
41320000	**Low discrimination**							**41320000**
Clostridium sporogenes		67.55	0.58	SAL 1				
Clostridium botulinum biovar A		30.35	0.54					
Clostridium bifermentans		1.62	0.32					
41320002	**Good identification**							**41320002**
Clostridium sporogenes		98.23	0.50	SAL 1				
Clostridium botulinum biovar A		1.34	0.20					
Clostridium novyi biovar A		0.36	0.22					
41320040	**Low discrimination**							**41320040**
Clostridium botulinum biovar A		89.53	0.45	SAL 1				
Clostridium sporogenes		6.74	0.24					
Clostridium novyi biovar A		3.07	0.32					
41330000	**Low discrimination**							**41330000**
					GRA	SPO		
Clostridium sporogenes		77.96	0.45	SAL 1	99	99		
Clostridium botulinum biovar A		12.67	0.34		99	99		
Fusobacterium mortiferum		8.12	0.43		1	1		
41330002	**Good identification**							**41330002**
					GRA	SPO		
Clostridium sporogenes		96.19	0.37	SAL 1	99	99		
Fusobacterium mortiferum		3.34	0.27		1	1		
Clostridium botulinum biovar A		0.47	-.01		99	99		
41330100	**Good identification**							**41330100**
					GRA	SPO		
Fusobacterium mortiferum		95.39	0.50	MLT 11	1	1		
Clostridium sporogenes		3.96	0.11		99	99		
Clostridium botulinum biovar A		0.64	-.01		99	99		
41331000	**Low discrimination**							**41331000**
					GRA	SPO		
Fusobacterium mortiferum		72.50	0.46	MLT 11	1	1		
Lactobacillus jensenii		22.05	0.28		99	1		
Clostridium sporogenes		4.69	0.11		99	99		
41331100	**Low discrimination**							**41331100**
					COC	GRA	SPO	
Fusobacterium mortiferum		68.46	0.53	MLT 11	1	1	1	
Atopobium parvulum		17.88	0.31		99	99	1	
Clostridium septicum		13.33	0.23		1	99	99	
41331102	**Low discrimination**							**41331102**
					COC	GRA	SPO	
Fusobacterium mortiferum		68.94	0.36	MLT 11	1	1	1	
Atopobium parvulum		28.60	0.18	BGL 10	99	99	1	
Clostridium septicum		2.47	-.06		1	99	99	
41331120	**Very good identification**							**41331120**
					COC	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		98.82	0.57	MNS 99	1	99	
Clostridium perfringens		0.78	0.28		1	84	
Atopobium parvulum		0.24	0.18		99	1	
41335100	**Low discrimination**						**41335100**
					COC	GRA	
Atopobium parvulum		85.31	0.36	SUC 96	99	99	
Fusobacterium mortiferum		13.24	0.34	MNS 99	1	1	
Actinomyces israelii		1.45	0.19		1	99	
41337102	**Acceptable identification**						**41337102**
Actinomyces israelii		94.56	0.35	SUC 98			
Bifidobacterium adolescentis		5.44	0.08	CEL 93			
41350000	**Low discrimination**						**41350000**
					GRA	SPO	
Fusobacterium mortiferum		68.08	0.44	MLT 11	1	1	
Clostridium sporogenes		17.74	0.19		99	99	
Clostridium bifermentans		4.18	0.10		99	99	
41350100	**Good identification**						**41350100**
					GRA		
Fusobacterium mortiferum		97.83	0.51	MLT 11	1		
Collinsella aerofaciens		2.17	-.03		99		
41351000	**Low discrimination**						**41351000**
					GRA		
Lactobacillus jensenii		55.54	0.41	SUC 90	99		
Fusobacterium mortiferum		32.17	0.47	MNS 99	1		
Lactobacillus catenaformis		10.06	0.29		99		
41351002	**Low discrimination**						**41351002**
					GRA		
Lactobacillus jensenii		68.20	0.38	SUC 90	99		
Lactobacillus catenaformis		18.52	0.29	MNS 99	99		
Fusobacterium mortiferum		6.59	0.31		1		
41351120	**Low discrimination**						**41351120**
					SPO		
Clostridium septicum		71.08	0.58	MNS 99	99		
Clostridium paraputrificum		27.47	0.48		99		
Collinsella aerofaciens		0.63	0.15		1		
41351122	**Low discrimination**						**41351122**
Clostridium paraputrificum		72.39	0.41	SUC 92			
Clostridium barati		9.72	0.32	MNS 99			
Clostridium septicum		8.92	0.29				
41357100	**Good identification**						**41357100**
Actinomyces israelii		100.0	0.42	SUC 98			
41357102	**Acceptable identification**						**41357102**
					SPO		
Actinomyces israelii		92.87	0.45	SUC 98	1		
Bifidobacterium adolescentis		4.82	0.16		1		
Clostridium tertium		2.31	0.03		99		
41357142	**Good identification**						**41357142**
Actinomyces israelii		100.0	0.38	SUC 98			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
41357500	**Good identification**							**41357500**
Actinomyces israelii		100.0	0.40	SUC 98				
41357502	**Low discrimination**							**41357502**
Actinomyces israelii		70.20	0.43	SUC 98				
Bifidobacterium adolescentis		29.81	0.31					
41357542	**Low discrimination**							**41357542**
Actinomyces israelii		88.11	0.36	SUC 98				
Bifidobacterium adolescentis		11.89	0.15					
41370000	**Low discrimination**							**41370000**
					GRA	SPO		
Fusobacterium mortiferum		62.47	0.35	MLT 11	1	1		
Clostridium sporogenes		18.17	0.11		99	99		
Lactobacillus jensenii		16.41	0.15		99	1		
41370100	**Acceptable identification**							**41370100**
					COC	GRA	SPO	
Fusobacterium mortiferum		94.72	0.42	MLT 11	1	1	1	
Clostridium septicum		2.66	-.03		1	99	99	
Atopobium parvulum		2.62	0.04		99	99	1	
41371000	**Acceptable identification**							**41371000**
					GRA	SPO		
Lactobacillus jensenii		94.10	0.50	SUC 90	99	1		
Fusobacterium mortiferum		5.43	0.38	MNS 99	1	1		
Clostridium septicum		0.24	-.03		99	99		
41371002	**Good identification**							**41371002**
					GRA			
Lactobacillus jensenii		97.85	0.47	SUC 90	99			
Fusobacterium mortiferum		0.94	0.22	MNS 99	1			
Lactobacillus acidophilus		0.74	0.07		99			
41371010	**Good identification**							**41371010**
Lactobacillus jensenii		100.0	0.41	SUC 90				
41371012	**Good identification**							**41371012**
					GRA			
Lactobacillus jensenii		97.78	0.38	SUC 90	99			
Leptotrichia buccalis		2.22	-.03	MNS 99	1			
41371020	**Low discrimination**							**41371020**
					SPO			
Lactobacillus jensenii		49.09	0.39	SUC 90	1			
Clostridium septicum		48.89	0.32	MNS 99	99			
Clostridium paraputrificum		1.67	0.04		99			
41371022	**Low discrimination**							**41371022**
					SPO			
Lactobacillus jensenii		89.75	0.36	SUC 90	1			
Clostridium barati		2.85	0.05	MNS 99	99			
Clostridium septicum		2.74	0.03		99			
41371120	**Very good identification**							**41371120**
					COC	SPO		
Clostridium septicum		96.48	0.66	MNS 99	1	99		
Clostridium paraputrificum		3.30	0.38		1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		0.17	0.25		99	1	
41371122	**Low discrimination**						**41371122**
					SPO		
Clostridium septicum		52.30	0.37	MNS 99	99		
Clostridium paraputrificum		37.61	0.32	BGL 2	99		
Lactobacillus acidophilus		5.99	0.25		1		
41373102	**Low discrimination**						**41373102**
					SPO		
Actinomyces israelii		62.94	0.36	SUC 98	1		
Clostridium tertium		31.22	0.17	XYL 92	99		
Lactobacillus acidophilus		5.84	0.04		1		
41375100	**Low discrimination**						**41375100**
					COC	GRA	
Atopobium parvulum		88.64	0.42	SUC 96	99	99	
Actinomyces israelii		8.57	0.39	MNS 99	1	99	
Fusobacterium mortiferum		1.97	0.25		1	1	
41375500	**Low discrimination**						**41375500**
					COC	SPO	
Actinomyces israelii		64.33	0.37	SUC 98	1	1	
Clostridium butyricum		27.46	0.10		1	99	
Atopobium parvulum		8.21	0.08		99	1	
41375502	**Low discrimination**						**41375502**
					SPO		
Actinomyces israelii		61.28	0.40	SUC 98	1		
Bifidobacterium adolescentis		31.94	0.29		1		
Clostridium butyricum		6.78	0.03		99		
41377000	**Good identification**						**41377000**
Actinomyces israelii		100.0	0.41	SUC 98			
41377002	**Good identification**						**41377002**
Actinomyces israelii		100.0	0.44	SUC 98			
41377042	**Good identification**						**41377042**
Actinomyces israelii		100.0	0.37	SUC 98			
41377100	**Good identification**						**41377100**
					COC	SPO	
Actinomyces israelii		95.40	0.52	SUC 98	1	1	
Clostridium tertium		2.84	0.11		1	99	
Atopobium parvulum		1.76	0.08		99	1	
41377102	**Acceptable identification**						**41377102**
					SPO		
Actinomyces israelii		93.57	0.55	SUC 98	1		
Clostridium tertium		5.57	0.19		99		
Bifidobacterium adolescentis		0.86	0.13		1		
41377110	**Low discrimination**						**41377110**
					SPO		
Actinomyces israelii		73.84	0.37	SUC 98	1		
Clostridium tertium		26.16	0.15	MLZ 13	99		
41377112	**Low discrimination**						**41377112**
					SPO		
Actinomyces israelii		56.06	0.40	SUC 98	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		39.72	0.24	MLZ 13	99	
Bifidobacterium adolescentis		4.22	0.15		1	
41377140	**Good identification**					**41377140**
Actinomyces israelii		100.0	0.45	SUC 98		
41377142	**Good identification**					**41377142**
Actinomyces israelii		100.0	0.48	SUC 98		
41377302	**Good identification**					**41377302**
Actinomyces israelii		100.0	0.35	SUC 98		
41377400	**Good identification**					**41377400**
Actinomyces israelii		100.0	0.40	SUC 98		
41377402	**Good identification**					**41377402**
Actinomyces israelii		100.0	0.43	SUC 98		
41377442	**Good identification**					**41377442**
Actinomyces israelii		100.0	0.36	SUC 98		
41377500	**Good identification**					**41377500**
					SPO	
Actinomyces israelii		99.17	0.50	SUC 98	1	
Clostridium butyricum		0.83	-.07		99	
41377502	**Acceptable identification**					**41377502**
Actinomyces israelii		93.00	0.53	SUC 98		
Bifidobacterium adolescentis		7.00	0.28			
41377510	**Good identification**					**41377510**
Actinomyces israelii		100.0	0.36	SUC 98		
41377512	**Low discrimination**					**41377512**
Actinomyces israelii		52.40	0.39	SUC 98		
Bifidobacterium adolescentis		32.25	0.29	MLZ 13		
Bifidobacterium dentium		15.35	0.06			
41377540	**Good identification**					**41377540**
Actinomyces israelii		100.0	0.43	SUC 98		
41377542	**Good identification**					**41377542**
Actinomyces israelii		97.66	0.46	SUC 98		
Bifidobacterium adolescentis		2.34	0.12			
41400000	**Low discrimination**					**41400000**
					SPO	
Clostridium botulinum biovar A		50.13	0.66	RAF 1	99	
Clostridium sporogenes		36.87	0.61		99	
Atopobium minutum		7.65	0.60		1	
41400001	**Low discrimination**					**41400001**
Clostridium novyi biovar A		61.43	0.53	RAF 1		
Clostridium botulinum biovar A		17.19	0.32			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		12.64	0.27				
41400002	**Low discrimination**						**41400002**
					SPO		
Clostridium sporogenes		67.27	0.53	RAF 1	99		
Atopobium minutum		27.91	0.57		1		
Clostridium botulinum biovar A		2.77	0.32		99		
41400003	**Low discrimination**						**41400003**
					SPO		
Clostridium novyi biovar A		60.69	0.40	RAF 1	99		
Clostridium sporogenes		23.59	0.19		99		
Atopobium minutum		9.79	0.22		1		
41400010	**Low discrimination**						**41400010**
Clostridium novyi biovar A		63.76	0.53	RAF 1			
Clostridium botulinum biovar A		17.84	0.32				
Clostridium sporogenes		13.12	0.27				
41400011	**Good identification**						**41400011**
Clostridium novyi biovar A		100.0	0.53	RAF 1			
41400012	**Low discrimination**						**41400012**
					SPO		
Clostridium novyi biovar A		64.52	0.40	RAF 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
41400013	**Good identification**						**41400013**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
41400020	**Low discrimination**						**41400020**
					SPO		
Atopobium minutum		62.92	0.51	RAF 1	1		
Clostridium botulinum biovar A		12.50	0.32		99		
Clostridium novyi biovar A		11.17	0.43		99		
41400021	**Acceptable identification**						**41400021**
					SPO		
Clostridium novyi biovar A		94.62	0.43	RAF 1	99		
Atopobium minutum		5.38	0.17		1		
41400022	**Low discrimination**						**41400022**
					GRA	SPO	
Atopobium minutum		54.58	0.48	RAF 1	99	1	
Pseudoflavonifractor capillosus		38.87	0.32		1	1	
Clostridium sporogenes		3.99	0.19		99	99	
41400030	**Acceptable identification**						**41400030**
					SPO		
Clostridium novyi biovar A		94.62	0.43	RAF 1	99		
Atopobium minutum		5.38	0.17		1		
41400031	**Good identification**						**41400031**
Clostridium novyi biovar A		100.0	0.43	RAF 1			
41400040	**Low discrimination**						**41400040**
Clostridium botulinum biovar A		87.87	0.57	RAF 1			
Clostridium novyi biovar A		7.08	0.50				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		2.19	0.27				
41400041	**Low discrimination**						**41400041**
Clostridium novyi biovar A		88.86	0.50	RAF 1			
Clostridium botulinum biovar A		11.14	0.23				
41400042	**Low discrimination**						**41400042**
Clostridium novyi biovar A		39.44	0.37	RAF 1			
Clostridium botulinum biovar A		28.02	0.23				
Clostridium sporogenes		23.00	0.19				
41400043	**Good identification**						**41400043**
Clostridium novyi biovar A		100.0	0.37	RAF 1			
41400050	**Low discrimination**						**41400050**
Clostridium novyi biovar A		88.86	0.50	RAF 1			
Clostridium botulinum biovar A		11.14	0.23				
41400051	**Good identification**						**41400051**
Clostridium novyi biovar A		100.0	0.50	RAF 1			
41400052	**Good identification**						**41400052**
Clostridium novyi biovar A		100.0	0.37	RAF 1			
41400053	**Good identification**						**41400053**
Clostridium novyi biovar A		100.0	0.37	RAF 1			
41400060	**Low discrimination**						**41400060**
					SPO		
Clostridium novyi biovar A		62.29	0.40	RAF 1	99		
Clostridium botulinum biovar A		31.24	0.23		99		
Atopobium minutum		5.32	0.17		1		
41400061	**Good identification**						**41400061**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
41400070	**Good identification**						**41400070**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
41400071	**Good identification**						**41400071**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
41401000	**Low discrimination**						**41401000**
					COC	SPO	
Atopobium minutum		51.01	0.51	RAF 1	1	1	
Blautia hansenii		17.86	0.32		99	1	
Clostridium botulinum biovar A		10.13	0.32		1	99	
41401001	**Low discrimination**						**41401001**
					COC	SPO	
Clostridium novyi biovar A		85.26	0.37	RAF 1	1	99	
Atopobium minutum		10.92	0.17	GAL 10	1	1	
Blautia hansenii		3.82	-.03		99	1	
41401002	**Low discrimination**						**41401002**
					COC	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		83.90	0.48	RAF 1	1	1	
Clostridium sporogenes		6.13	0.19		1	99	
Blautia hansenii		4.90	0.15		99	1	
41401010	**Low discrimination**						**41401010**
					COC	SPO	
Clostridium novyi biovar A		85.26	0.37	RAF 1	1	99	
Atopobium minutum		10.92	0.17	GAL 10	1	1	
Blautia hansenii		3.82	-.03		99	1	
41401011	**Good identification**						**41401011**
Clostridium novyi biovar A		100.0	0.37	RAF 1			
41401020	**Low discrimination**						**41401020**
					COC	SPO	
Atopobium minutum		81.14	0.43	RAF 1	1	1	
Blautia hansenii		9.47	0.15		99	1	
Clostridium novyi biovar A		4.80	0.26		1	99	
41401022	**Low discrimination**						**41401022**
					COC	GRA	
Atopobium minutum		77.67	0.40	RAF 1	1	99	
Pseudoflavonifractor capillosus		18.44	0.15		1	1	
Blautia hansenii		1.51	-.01		99	99	
41401100	**Very good identification**						**41401100**
					COC	GRA	
Blautia hansenii		99.52	0.66	MLT 99	99	99	
Fusobacterium mortiferum		0.45	0.43		1	1	
Atopobium minutum		0.03	0.17		1	99	
41401102	**Good identification**						**41401102**
					COC	GRA	
Blautia hansenii		99.37	0.50	MLT 99	99	99	
Fusobacterium mortiferum		0.45	0.27	BGL 10	1	1	
Atopobium minutum		0.17	0.14		1	99	
41401120	**Good identification**						**41401120**
					COC	GRA	
Blautia hansenii		99.84	0.50	MLT 99	99	99	
Atopobium minutum		0.09	0.09	NAG 10	1	99	
Prevotella melaninogenica		0.08	-.04		1	1	
41401500	**Good identification**						**41401500**
Blautia hansenii		100.0	0.53	MLT 99			
41401502	**Good identification**						**41401502**
Blautia hansenii		100.0	0.37	MLT 99			
41401520	**Good identification**						**41401520**
					COC	GRA	
Blautia hansenii		99.56	0.37	MLT 99	99	99	
Odoribacter splanchnicus		0.44	-.07	NAG 10	1	1	
41404040	**Good identification**						**41404040**
Clostridium glycolicum		97.32	0.49	RAF 1			
Clostridium botulinum biovar A		2.48	0.23	FRU 90			
Clostridium novyi biovar A		0.20	0.16				
41405100	**Good identification**						**41405100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Blautia hansenii		99.81	0.53	MLT 99	99	99	
Fusobacterium mortiferum		0.19	0.24		1	1	
41405102	**Good identification**						**41405102**
Blautia hansenii		100.0	0.37	MLT 99			
41405120	**Good identification**						**41405120**
Blautia hansenii		100.0	0.37	MLT 99			
41405500	**Good identification**						**41405500**
Blautia hansenii		100.0	0.40	MLT 99			
41410000	**Low discrimination**						**41410000**
					GRA		
Fusobacterium mortiferum		27.78	0.68		1		
Eubacterium limosum		24.30	0.49		99		
Atopobium minutum		17.25	0.59		99		
41410001	**Low discrimination**						**41410001**
					GRA	SPO	
Clostridium novyi biovar A		32.85	0.37	RAF 1	99	99	
Fusobacterium mortiferum		18.76	0.34	FRU 10	1	1	
Eubacterium limosum		16.41	0.15		99	1	
41410002	**Low discrimination**						**41410002**
					GRA	SPO	
Atopobium minutum		55.61	0.56	RAF 1	99	1	
Clostridium sporogenes		25.63	0.40		99	99	
Fusobacterium mortiferum		14.93	0.52		1	1	
41410010	**Low discrimination**						**41410010**
					GRA	SPO	
Clostridium difficile		87.37	0.45	RAF 1	99	99	
Clostridium novyi biovar A		4.25	0.37	MAN 90	99	99	
Fusobacterium mortiferum		2.43	0.34		1	1	
41410011	**Low discrimination**						**41410011**
Clostridium novyi biovar A		82.79	0.37	RAF 1			
Clostridium difficile		17.21	0.10	FRU 10			
41410020	**Low discrimination**						**41410020**
					GRA		
Atopobium minutum		79.52	0.51	RAF 1	99		
Pseudoramibacter alactolyticus		7.16	0.19		99		
Fusobacterium mortiferum		3.88	0.34		1		
41410022	**Low discrimination**						**41410022**
					GRA		
Atopobium minutum		88.61	0.48	RAF 1	99		
Pseudoflavonifractor capillosus		7.60	0.15		1		
Pseudoramibacter alactolyticus		1.33	0.03		99		
41410040	**Low discrimination**						**41410040**
					SPO		
Clostridium glycolicum		60.08	0.43	RAF 1	99		
Clostridium botulinum biovar A		20.39	0.36	XYL 95	99		
Pseudoramibacter alactolyticus		6.41	0.21		1		
41410100	**Very good identification**						**41410100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		98.86	0.74		1		
Eubacterium limosum		0.37	0.15		99		
Atopobium minutum		0.27	0.25		99		
41410101	**Good identification**						**41410101**
Fusobacterium mortiferum		100.0	0.40	URE 1			
41410102	**Very good identification**						**41410102**
					GRA	SPO	
Fusobacterium mortiferum		97.70	0.58	BGL 10	1	1	
Atopobium minutum		1.58	0.22		99	1	
Clostridium sporogenes		0.73	0.06		99	99	
41410110	**Low discrimination**						**41410110**
					GRA	SPO	
Fusobacterium mortiferum		86.51	0.40	MLZ 1	1	1	
Clostridium difficile		13.49	0.10		99	99	
41410120	**Acceptable identification**						**41410120**
					GRA		
Fusobacterium mortiferum		91.85	0.40	NAG 1	1		
Atopobium minutum		8.15	0.17		99		
41410140	**Low discrimination**						**41410140**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.40	SOR 1	1	1	
Clostridium glycolicum		9.30	0.09		99	99	
Clostridium botulinum biovar A		3.16	0.02		99	99	
41410300	**Good identification**						**41410300**
Fusobacterium mortiferum		100.0	0.40	RHA 1			
41410500	**Good identification**						**41410500**
Fusobacterium mortiferum		100.0	0.40	ARA 1			
41411000	**Low discrimination**						**41411000**
					GRA		
Fusobacterium mortiferum		85.66	0.71		1		
Atopobium minutum		11.82	0.51		99		
Lactobacillus fermentum		1.07	0.40		99		
41411001	**Low discrimination**						**41411001**
					GRA	SPO	
Fusobacterium mortiferum		78.88	0.37	URE 1	1	1	
Atopobium minutum		10.89	0.17		99	1	
Clostridium novyi biovar A		10.23	0.20		99	99	
41411002	**Low discrimination**						**41411002**
					GRA		
Fusobacterium mortiferum		51.23	0.55	BGL 10	1		
Atopobium minutum		42.42	0.48		99		
Lactobacillus fermentum		5.77	0.40		99		
41411010	**Low discrimination**						**41411010**
					GRA	SPO	
Fusobacterium mortiferum		66.21	0.37	MLZ 1	1	1	
Clostridium difficile		16.07	0.10		99	99	
Atopobium minutum		9.14	0.17		99	1	
41411020	**Low discrimination**						**41411020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Atopobium minutum		75.91	0.43	RAF 1	99		
Fusobacterium mortiferum		16.67	0.37		1		
Lactobacillus fermentum		6.88	0.32		99		
41411022	**Low discrimination**						**41411022**
					GRA		
Atopobium minutum		83.26	0.40	RAF 1	99		
Lactobacillus fermentum		11.32	0.32		99		
Fusobacterium mortiferum		3.05	0.20		1		
41411100	**Very good identification**						**41411100**
					COC	GRA	
Fusobacterium mortiferum		96.86	0.77		1	1	
Blautia hansenii		2.17	0.32		99	99	
Lactobacillus fermentum		0.85	0.44		1	99	
41411101	**Good identification**						**41411101**
					COC	GRA	
Fusobacterium mortiferum		97.81	0.43	URE 1	1	1	
Blautia hansenii		2.19	-.03		99	99	
41411102	**Good identification**						**41411102**
					COC	GRA	
Fusobacterium mortiferum		90.52	0.61	BGL 10	1	1	
Lactobacillus fermentum		7.12	0.44		1	99	
Blautia hansenii		2.03	0.15		99	99	
41411110	**Good identification**						**41411110**
					COC	GRA	
Fusobacterium mortiferum		95.04	0.43	MLZ 1	1	1	
Eubacterium saburreum		2.83	0.15		1	85	
Blautia hansenii		2.13	-.03		99	99	
41411120	**Low discrimination**						**41411120**
					COC	GRA	
Fusobacterium mortiferum		57.95	0.43	NAG 1	1	1	
Lactobacillus fermentum		16.72	0.35		1	99	
Blautia hansenii		14.30	0.15		99	99	
41411122	**Low discrimination**						**41411122**
					GRA	CAT	
Lactobacillus fermentum		46.34	0.35	ESL 1	99	1	
Fusobacterium mortiferum		17.85	0.27	SUC 87	1	5	
Capnocytophaga ochracea		16.47	0.12		1	90	
41411140	**Good identification**						**41411140**
					COC	GRA	
Fusobacterium mortiferum		97.81	0.43	SOR 1	1	1	
Blautia hansenii		2.19	-.03		99	99	
41411200	**Low discrimination**						**41411200**
					GRA		
Fusobacterium mortiferum		87.87	0.37	RHA 1	1		
Atopobium minutum		12.13	0.17		99		
41411300	**Good identification**						**41411300**
					COC	GRA	
Fusobacterium mortiferum		97.81	0.43	RHA 1	1	1	
Blautia hansenii		2.19	-.03		99	99	
41411400	**Low discrimination**						**41411400**
					GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		52.10	0.40	ESL 1	99		
Fusobacterium mortiferum		42.09	0.37	SUC 87	1		
Atopobium minutum		5.81	0.17		99		
41411402	**Low discrimination**						**41411402**
					GRA		
Lactobacillus fermentum		85.91	0.40	ESL 1	99		
Fusobacterium mortiferum		7.71	0.20	SUC 87	1		
Atopobium minutum		6.38	0.14		99		
41411500	**Low discrimination**						**41411500**
					COC	GRA	
Fusobacterium mortiferum		44.30	0.43	ARA 1	1	1	
Lactobacillus fermentum		38.34	0.44		1	99	
Blautia hansenii		17.36	0.19		99	99	
41411502	**Low discrimination**						**41411502**
					COC	GRA	
Lactobacillus fermentum		84.84	0.44	ESL 1	1	99	
Fusobacterium mortiferum		10.89	0.27	SUC 87	1	1	
Blautia hansenii		4.27	0.03		99	99	
41411520	**Low discrimination**						**41411520**
					COC	GRA	
Lactobacillus fermentum		75.83	0.35	ESL 1	1	99	
Blautia hansenii		11.44	0.03	SUC 87	99	99	
Odoribacter splanchnicus		9.24	-.03		1	1	
41411522	**Low discrimination**						**41411522**
					GRA	CAT	
Lactobacillus fermentum		85.68	0.35	ESL 1	99	1	
Capnocytophaga ochracea		5.37	-.01	SUC 87	1	90	
Parabacteroides distasonis		5.01	0.11		1	76	
41412000	**Low discrimination**						**41412000**
					SPO		
Eubacterium limosum		87.55	0.63	RAF 1	1		
Clostridium difficile		10.93	0.49		99		
Pseudoramibacter alactolyticus		0.93	0.27		1		
41412010	**Very good identification**						**41412010**
					SPO		
Clostridium difficile		98.35	0.61	RAF 1	99		
Eubacterium limosum		1.63	0.29		1		
Pseudoramibacter alactolyticus		0.02	-.07		1		
41412050	**Good identification**						**41412050**
					SPO		
Clostridium difficile		99.68	0.39	RAF 1	99		
Eubacterium limosum		0.32	-.06	SOR 5	1		
41413100	**Good identification**						**41413100**
					COC	GRA	
Fusobacterium mortiferum		96.35	0.43	MAN 1	1	1	
Blautia hansenii		2.16	-.03		99	99	
Eubacterium limosum		1.49	-.06		1	99	
41414000	**Low discrimination**						**41414000**
					GRA	SPO	
Clostridium glycolicum		70.31	0.43	RAF 1	99	99	
Fusobacterium mortiferum		21.36	0.49	SOR 95	1	1	
Eubacterium limosum		2.51	0.15		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41414040	**Very good identification**						**41414040**
					GRA	SPO	
Clostridium glycolicum		99.97	0.65	RAF 1	99	99	
Clostridium botulinum biovar A		0.02	0.02		99	99	
Fusobacterium mortiferum		0.02	0.15		1	1	
41414100	**Good identification**						**41414100**
					GRA	SPO	
Fusobacterium mortiferum		97.95	0.55	XYL 7	1	1	
Clostridium glycolicum		1.40	0.09		99	99	
Actinomyces meyeri		0.66	0.09		99	1	
41414102	**Good identification**						**41414102**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
41415000	**Low discrimination**						**41415000**
					GRA	SPO	
Fusobacterium mortiferum		73.54	0.52	XYL 7	1	1	
Clostridium glycolicum		17.93	0.27		99	99	
Lactobacillus fermentum		7.18	0.36		99	1	
41415002	**Low discrimination**						**41415002**
					GRA		
Fusobacterium mortiferum		50.27	0.35	XYL 7	1		
Lactobacillus fermentum		44.14	0.36	BGL 10	99		
Atopobium minutum		5.59	0.14		99		
41415040	**Good identification**						**41415040**
					GRA	SPO	
Clostridium glycolicum		99.78	0.49	RAF 1	99	99	
Fusobacterium mortiferum		0.22	0.18	GAL 10	1	1	
41415100	**Low discrimination**						**41415100**
					COC	GRA	
Fusobacterium mortiferum		88.48	0.58	XYL 7	1	1	
Lactobacillus fermentum		6.04	0.40		1	99	
Blautia hansenii		4.65	0.19		99	99	
41415102	**Low discrimination**						**41415102**
					COC	GRA	
Fusobacterium mortiferum		60.00	0.42	XYL 7	1	1	
Lactobacillus fermentum		36.84	0.40	BGL 10	1	99	
Blautia hansenii		3.16	0.03		99	99	
41415322	**Good identification**						**41415322**
Parabacteroides distasonis		100.0	0.40	MNS 98			
41415400	**Acceptable identification**						**41415400**
					GRA		
Lactobacillus fermentum		90.62	0.36	ESL 1	99		
Fusobacterium mortiferum		9.38	0.18	SUC 87	1		
41415402	**Good identification**						**41415402**
Lactobacillus fermentum		100.0	0.36	ESL 1			
41415500	**Low discrimination**						**41415500**
					COC	GRA	
Lactobacillus fermentum		77.88	0.40	ESL 1	1	99	
Fusobacterium mortiferum		11.53	0.24	SUC 87	1	1	
Blautia hansenii		10.59	0.06		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41415502	**Good identification**						**41415502**
Lactobacillus fermentum		100.0	0.40	ESL 1			
41415722	**Good identification**						**41415722**
Parabacteroides distasonis		100.0	0.36	MNS 98			
41416010	**Good identification**						**41416010**
					SPO		
Clostridium difficile		99.68	0.39	RAF 1	99		
Eubacterium limosum		0.32	-.06	XYL 5	1		
41420000	**Low discrimination**						**41420000**
					SPO		
Clostridium botulinum biovar A		50.17	0.56	RAF 1	99		
Clostridium sporogenes		49.19	0.53		99		
Atopobium minutum		0.31	0.25		1		
41420002	**Good identification**						**41420002**
					SPO		
Clostridium sporogenes		95.83	0.45	RAF 1	99		
Clostridium botulinum biovar A		2.96	0.21		99		
Atopobium minutum		1.21	0.22		1		
41420040	**Good identification**						**41420040**
Clostridium botulinum biovar A		96.49	0.47	RAF 1			
Clostridium sporogenes		3.20	0.19				
Clostridium novyi biovar A		0.31	0.16				
41430000	**Low discrimination**						**41430000**
					GRA	SPO	
Fusobacterium mortiferum		51.27	0.59		1	1	
Clostridium sporogenes		32.75	0.40		99	99	
Clostridium botulinum biovar A		12.08	0.35		99	99	
41430100	**Very good identification**						**41430100**
					GRA	SPO	
Fusobacterium mortiferum		99.62	0.65		1	1	
Clostridium sporogenes		0.28	0.06		99	99	
Clostridium botulinum biovar A		0.10	0.01		99	99	
41430102	**Good identification**						**41430102**
					GRA	SPO	
Fusobacterium mortiferum		99.18	0.49	BGL 10	1	1	
Clostridium sporogenes		0.82	-.02		99	99	
41431000	**Very good identification**						**41431000**
					GRA	SPO	
Fusobacterium mortiferum		98.76	0.62		1	1	
Atopobium minutum		0.46	0.17		99	1	
Clostridium sporogenes		0.43	0.06		99	99	
41431002	**Good identification**						**41431002**
					GRA	SPO	
Fusobacterium mortiferum		96.07	0.46	BGL 10	1	1	
Atopobium minutum		2.69	0.14		99	1	
Clostridium sporogenes		1.24	-.02		99	99	
41431100	**Very good identification**						**41431100**
					COC	GRA	
Fusobacterium mortiferum		99.84	0.68		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		0.08	0.14		1	85	
Blautia hansenii		0.08	-.03		99	99	
41431102	**Good identification**						**41431102**
Fusobacterium mortiferum		100.0	0.52	BGL 10			
41432010	**Good identification**						**41432010**
					SPO		
Clostridium difficile		99.78	0.42	RAF 1	99		
Eubacterium limosum		0.22	-.06	TRE 7	1		
41434000	**Low discrimination**						**41434000**
					GRA	SPO	
Fusobacterium mortiferum		81.39	0.40	XYL 7	1	1	
Clostridium glycolicum		9.06	0.09		99	99	
Clostridium sporogenes		6.98	0.06		99	99	
41434100	**Good identification**						**41434100**
Fusobacterium mortiferum		100.0	0.46	XYL 7			
41435000	**Good identification**						**41435000**
Fusobacterium mortiferum		100.0	0.43	XYL 7			
41435100	**Good identification**						**41435100**
Fusobacterium mortiferum		99.42	0.49	XYL 7			
Eubacterium saburreum		0.58	0.09				
41435122	**Good identification**						**41435122**
Parabacteroides distasonis		100.0	0.43	MNS 98			
41435132	**Good identification**						**41435132**
Parabacteroides distasonis		100.0	0.38	MNS 98			
41435322	**Good identification**						**41435322**
Parabacteroides distasonis		100.0	0.51	MNS 98			
41435332	**Good identification**						**41435332**
Parabacteroides distasonis		100.0	0.46	MNS 98			
41435522	**Good identification**						**41435522**
Parabacteroides distasonis		100.0	0.39	MNS 98			
41435722	**Good identification**						**41435722**
Parabacteroides distasonis		100.0	0.47	MNS 98			
41435732	**Good identification**						**41435732**
Parabacteroides distasonis		100.0	0.43	MNS 98			
41450000	**Good identification**						**41450000**
					GRA		
Fusobacterium mortiferum		92.07	0.60		1		
Eubacterium limosum		2.44	0.15		99		
Atopobium minutum		1.73	0.25		99		
41450002	**Low discrimination**						**41450002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		85.84	0.43	BGL 10	1	1	
Atopobium minutum		9.69	0.22		99	1	
Clostridium sporogenes		4.47	0.06		99	99	
41450100	**Very good identification**						**41450100**
Fusobacterium mortiferum		100.0	0.66				
41450102	**Good identification**						**41450102**
Fusobacterium mortiferum		100.0	0.50	BGL 10			
41451000	**Very good identification**						**41451000**
					GRA		
Fusobacterium mortiferum		99.58	0.63		1		
Atopobium minutum		0.42	0.17		99		
41451002	**Good identification**						**41451002**
					GRA		
Fusobacterium mortiferum		97.55	0.46	BGL 10	1		
Atopobium minutum		2.45	0.14		99		
41451100	**Very good identification**						**41451100**
					COC	GRA	
Fusobacterium mortiferum		99.93	0.69		1	1	
Blautia hansenii		0.07	-.03		99	99	
41451102	**Good identification**						**41451102**
Fusobacterium mortiferum		100.0	0.53	BGL 10			
41452010	**Good identification**						**41452010**
					SPO		
Clostridium difficile		99.68	0.39	RAF 1	99		
Eubacterium limosum		0.32	-.06	CEL 5	1		
41454000	**Acceptable identification**						**41454000**
					GRA	SPO	
Fusobacterium mortiferum		90.93	0.41	XYL 7	1	1	
Clostridium glycolicum		9.07	0.09		99	99	
41454100	**Good identification**						**41454100**
Fusobacterium mortiferum		100.0	0.47	XYL 7			
41455000	**Good identification**						**41455000**
Fusobacterium mortiferum		100.0	0.44	XYL 7			
41455100	**Good identification**						**41455100**
Fusobacterium mortiferum		100.0	0.50	XYL 7			
41470000	**Good identification**						**41470000**
					GRA	SPO	
Fusobacterium mortiferum		97.42	0.51		1	1	
Clostridium sporogenes		1.89	0.06		99	99	
Clostridium botulinum biovar A		0.70	0.01		99	99	
41470100	**Very good identification**						**41470100**
Fusobacterium mortiferum		100.0	0.57				
41470102	**Good identification**						**41470102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.41	BGL 10		
41471000	**Good identification**					**41471000**
Fusobacterium mortiferum		100.0	0.54			
41471002	**Good identification**					**41471002**
Fusobacterium mortiferum		100.0	0.38	BGL 10		
41471100	**Very good identification**					**41471100**
Fusobacterium mortiferum		100.0	0.60			
41471102	**Good identification**					**41471102**
Fusobacterium mortiferum		100.0	0.44	BGL 10		
41474100	**Good identification**					**41474100**
Fusobacterium mortiferum		100.0	0.38	XYL 7		
41475100	**Good identification**					**41475100**
Fusobacterium mortiferum		100.0	0.41	XYL 7		
41475122	**Good identification**					**41475122**
Parabacteroides distasonis		100.0	0.38	MNS 98		
41475322	**Good identification**					**41475322**
Parabacteroides distasonis		100.0	0.46	MNS 98		
41475332	**Good identification**					**41475332**
Parabacteroides distasonis		100.0	0.42	MNS 98		
41475722	**Good identification**					**41475722**
Parabacteroides distasonis		100.0	0.43	MNS 98		
41475732	**Good identification**					**41475732**
Parabacteroides distasonis		100.0	0.38	MNS 98		
41500000	**Low discrimination**					**41500000**
Clostridium sporogenes		57.92	0.66	RAF 1		
Clostridium botulinum biovar A		34.70	0.64			
Clostridium bifermentans		4.16	0.48			
41500001	**Low discrimination**					**41500001**
Clostridium novyi biovar A		59.67	0.53	RAF 1		
Clostridium sporogenes		22.81	0.32			
Clostridium botulinum biovar A		13.66	0.30			
41500002	**Very good identification**					**41500002**
Clostridium sporogenes		96.04	0.58	RAF 1		
Clostridium botulinum biovar A		1.74	0.30			
Clostridium novyi biovar A		1.34	0.40			
41500003	**Low discrimination**					**41500003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		58.08	0.40	RAF 1		
Clostridium sporogenes		41.92	0.24			
41500010	**Low discrimination**					**41500010**
Clostridium novyi biovar A		60.71	0.53	RAF 1		
Clostridium sporogenes		23.20	0.32			
Clostridium botulinum biovar A		13.90	0.30			
41500011	**Good identification**					**41500011**
Clostridium novyi biovar A		99.62	0.53	RAF 1		
Clostridium sporogenes		0.39	-.03			
41500012	**Low discrimination**					**41500012**
Clostridium novyi biovar A		58.08	0.40	RAF 1		
Clostridium sporogenes		41.92	0.24			
41500013	**Good identification**					**41500013**
Clostridium novyi biovar A		100.0	0.40	RAF 1		
41500021	**Good identification**					**41500021**
Clostridium novyi biovar A		98.48	0.43	RAF 1		
Clostridium sporogenes		1.52	-.03			
41500030	**Good identification**					**41500030**
Clostridium novyi biovar A		98.48	0.43	RAF 1		
Clostridium sporogenes		1.52	-.03			
41500031	**Good identification**					**41500031**
Clostridium novyi biovar A		100.0	0.43	RAF 1		
41500040	**Low discrimination**					**41500040**
Clostridium botulinum biovar A		83.26	0.55	RAF 1		
Clostridium novyi biovar A		8.20	0.50			
Clostridium sporogenes		4.70	0.32			
41500041	**Acceptable identification**					**41500041**
Clostridium novyi biovar A		90.22	0.50	RAF 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41500043	**Good identification**					**41500043**
Clostridium novyi biovar A		100.0	0.37	RAF 1		
41500050	**Acceptable identification**					**41500050**
Clostridium novyi biovar A		90.22	0.50	RAF 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
41500051	**Good identification**					**41500051**
Clostridium novyi biovar A		100.0	0.50	RAF 1		
41500052	**Good identification**					**41500052**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.37	RAF 1			
41500053	**Good identification**						**41500053**
Clostridium novyi biovar A		100.0	0.37	RAF 1			
41500060	**Low discrimination**						**41500060**
Clostridium novyi biovar A		67.71	0.40	RAF 1			
Clostridium botulinum biovar A		27.78	0.21				
Clostridium bifermentans		2.94	0.04				
41500061	**Good identification**						**41500061**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
41500070	**Good identification**						**41500070**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
41500071	**Good identification**						**41500071**
Clostridium novyi biovar A		100.0	0.40	RAF 1			
41500100	**Very good identification**						**41500100**
					COC	SPO	
Blautia hansenii		98.65	0.66	GAL 99	99	1	
Clostridium sporogenes		0.77	0.32		1	99	
Clostridium botulinum biovar A		0.46	0.30		1	99	
41500102	**Good identification**						**41500102**
					COC	SPO	
Blautia hansenii		97.71	0.50	GAL 99	99	1	
Clostridium sporogenes		2.30	0.24	BGL 10	1	99	
41500120	**Good identification**						**41500120**
					COC	GRA	
Blautia hansenii		99.62	0.50	GAL 99	99	99	
Prevotella bivia		0.23	-.03	NAG 10	1	1	
Prevotella melaninogenica		0.08	-.04		1	1	
41500500	**Good identification**						**41500500**
					COC	SPO	
Blautia hansenii		99.96	0.53	GAL 99	99	1	
Clostridium sporogenes		0.05	-.03		1	99	
41500502	**Good identification**						**41500502**
Blautia hansenii		100.0	0.37	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41500520	**Good identification**						**41500520**
Blautia hansenii		100.0	0.37	GAL 99			
41501000	**Very good identification**						**41501000**
					COC	SPO	
Blautia hansenii		98.40	0.66	LAC 99	99	1	
Clostridium sporogenes		0.77	0.32		1	99	
Clostridium botulinum biovar A		0.46	0.30		1	99	
41501002	**Good identification**						**41501002**
					COC	SPO	
Blautia hansenii		97.13	0.50	LAC 99	99	1	
Clostridium sporogenes		2.28	0.24	BGL 10	1	99	
Clostridium novyi biovar A		0.35	0.24		1	99	
41501011	**Good identification**						**41501011**
					COC	SPO	
Clostridium novyi biovar A		95.71	0.37	RAF 1	1	99	
Blautia hansenii		4.29	-.03	GAL 10	99	1	
41501020	**Good identification**						**41501020**
					COC	SPO	
Blautia hansenii		99.26	0.50	LAC 99	99	1	
Clostridium novyi biovar A		0.51	0.26	NAG 10	1	99	
Atopobium minutum		0.09	0.09		1	1	
41501100	**Excellent identification**						**41501100**
					COC	GRA	
Blautia hansenii		100.0	1.00		99	99	
Fusobacterium mortiferum		0.00	0.28		1	1	
Eubacterium saburreum		0.00	0.19		1	85	
41501101	**Very good identification**						**41501101**
					COC	SPO	
Blautia hansenii		99.99	0.66	URE 1	99	1	
Clostridium chauvoei		0.02	0.06		1	99	
41501102	**Very good identification**						**41501102**
					COC	GRA	
Blautia hansenii		100.0	0.84	BGL 10	99	99	
Lactobacillus fermentum		0.00	0.18		1	99	
Fusobacterium mortiferum		0.00	0.11		1	1	
41501103	**Good identification**						**41501103**
Blautia hansenii		100.0	0.50	URE 1			
41501110	**Very good identification**						**41501110**
					COC		
Blautia hansenii		99.98	0.66	MLZ 1	99		
Eubacterium saburreum		0.02	0.17		1		
41501112	**Good identification**						**41501112**
Blautia hansenii		100.0	0.50	MLZ 1			
41501120	**Very good identification**						**41501120**
					COC	GRA	
Blautia hansenii		99.92	0.84	NAG 10	99	99	
Prevotella melaninogenica		0.08	0.30		1	1	
41501121	**Good identification**						**41501121**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		99.92	0.50	NAG 10	99	99		
Prevotella melaninogenica		0.08	-.04	URE 1	1	1		
41501122	**Very good identification**							**41501122**
					COC	CAT	GRA	
Blautia hansenii		99.82	0.67	NAG 10	99	1	99	
Prevotella melaninogenica		0.13	0.18	BGL 10	1	1	1	
Capnocytophaga ochracea		0.04	0.12		1	90	1	
41501130	**Good identification**							**41501130**
					COC	GRA		
Blautia hansenii		99.92	0.50	MLZ 1	99	99		
Prevotella melaninogenica		0.08	-.04	NAG 10	1	1		
41501140	**Very good identification**							**41501140**
Blautia hansenii		100.0	0.66	SOR 1				
41501142	**Good identification**							**41501142**
Blautia hansenii		100.0	0.50	SOR 1				
41501160	**Good identification**							**41501160**
					COC	GRA		
Blautia hansenii		99.92	0.50	NAG 10	99	99		
Prevotella melaninogenica		0.08	-.04	SOR 1	1	1		
41501300	**Very good identification**							**41501300**
Blautia hansenii		100.0	0.66	RHA 1				
41501302	**Good identification**							**41501302**
Blautia hansenii		100.0	0.50	RHA 1				
41501320	**Good identification**							**41501320**
					COC	GRA		
Blautia hansenii		99.92	0.50	RHA 1	99	99		
Prevotella melaninogenica		0.08	-.04	NAG 10	1	1		
41501400	**Good identification**							**41501400**
					COC	SPO		
Blautia hansenii		99.91	0.53	LAC 99	99	1		
Clostridium sporogenes		0.05	-.03		1	99		
Lactobacillus fermentum		0.04	0.14		1	1		
41501402	**Good identification**							**41501402**
					COC			
Blautia hansenii		99.63	0.37	LAC 99	99			
Lactobacillus fermentum		0.37	0.14	BGL 10	1			
41501420	**Good identification**							**41501420**
Blautia hansenii		100.0	0.37	LAC 99				
41501500	**Very good identification**							**41501500**
					COC			
Blautia hansenii		100.0	0.87		99			
Lactobacillus fermentum		0.00	0.18		1			
Bifidobacterium longum subsp. longum		0.00	0.13		1			
41501501	**Good identification**							**41501501**
Blautia hansenii		100.0	0.53	URE 1				
41501502	**Very good identification**							**41501502**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC			
Blautia hansenii		99.99	0.71	BGL 10	99			
Bifidobacterium longum subsp. longum		0.01	0.13		1			
Lactobacillus fermentum		0.01	0.18		1			
41501503	**Good identification**							**41501503**
Blautia hansenii		100.0	0.37	URE 1				
41501510	**Good identification**							**41501510**
					COC			
Blautia hansenii		99.84	0.53	MLZ 1	99			
Bifidobacterium longum subsp. longum		0.16	0.17		1			
41501512	**Good identification**							**41501512**
					COC			
Blautia hansenii		98.58	0.37	MLZ 1	99			
Bifidobacterium longum subsp. longum		1.42	0.17	BGL 10	1			
41501520	**Very good identification**							**41501520**
					COC	GRA		
Blautia hansenii		99.96	0.71	NAG 10	99	99		
Bifidobacterium longum subsp. longum		0.03	0.23		1	99		
Prevotella melaninogenica		0.00	-.04		1	1		
41501521	**Good identification**							**41501521**
Blautia hansenii		100.0	0.37	NAG 10				
41501522	**Good identification**							**41501522**
					COC	GRA	CAT	
Blautia hansenii		99.66	0.54	NAG 10	99	99	1	
Bifidobacterium longum subsp. longum		0.31	0.23	BGL 10	1	99	1	
Capnocytophaga ochracea		0.04	-.01		1	1	90	
41501530	**Acceptable identification**							**41501530**
					COC			
Blautia hansenii		94.55	0.37	MLZ 1	99			
Bifidobacterium longum subsp. longum		5.45	0.27	NAG 10	1			
41501540	**Good identification**							**41501540**
Blautia hansenii		100.0	0.53	SOR 1				
41501542	**Good identification**							**41501542**
Blautia hansenii		100.0	0.37	SOR 1				
41501560	**Good identification**							**41501560**
Blautia hansenii		100.0	0.37	NAG 10				
41501700	**Good identification**							**41501700**
Blautia hansenii		100.0	0.53	RHA 1				
41501702	**Good identification**							**41501702**
Blautia hansenii		100.0	0.37	RHA 1				
41501720	**Good identification**							**41501720**
Blautia hansenii		100.0	0.37	RHA 1				
41503100	**Very good identification**							**41503100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		100.0	0.66	MAN 1			
41503102	**Good identification**						**41503102**
Blautia hansenii		100.0	0.50	MAN 1			
41503120	**Good identification**						**41503120**
					COC	GRA	
Blautia hansenii		99.92	0.50	MAN 1	99	99	
Prevotella melaninogenica		0.08	-.04	NAG 10	1	1	
41503500	**Good identification**						**41503500**
Blautia hansenii		100.0	0.53	MAN 1			
41503502	**Good identification**						**41503502**
Blautia hansenii		100.0	0.37	MAN 1			
41503520	**Good identification**						**41503520**
Blautia hansenii		100.0	0.37	MAN 1			
41504040	**Good identification**						**41504040**
Clostridium glycolicum		97.83	0.50	RAF 1			
Clostridium botulinum biovar A		1.88	0.21	FRU 90			
Clostridium novyi biovar A		0.19	0.16				
41504100	**Good identification**						**41504100**
					COC	SPO	
Blautia hansenii		99.96	0.53	GAL 99	99	1	
Clostridium sporogenes		0.05	-.03		1	99	
41504102	**Good identification**						**41504102**
Blautia hansenii		100.0	0.37	GAL 99			
41504120	**Good identification**						**41504120**
Blautia hansenii		100.0	0.37	GAL 99			
41504500	**Good identification**						**41504500**
Blautia hansenii		100.0	0.40	GAL 99			
41505000	**Good identification**						**41505000**
					COC	SPO	
Blautia hansenii		99.71	0.53	LAC 99	99	1	
Clostridium glycolicum		0.24	0.11		1	99	
Clostridium sporogenes		0.05	-.03		1	99	
41505002	**Good identification**						**41505002**
Blautia hansenii		100.0	0.37	LAC 99			
41505020	**Good identification**						**41505020**
Blautia hansenii		100.0	0.37	LAC 99			
41505100	**Very good identification**						**41505100**
					COC		
Blautia hansenii		100.0	0.87		99		
Eubacterium saburreum		0.00	0.14		1		
Lactobacillus fermentum		0.00	0.14		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41505101	**Good identification**						**41505101**
Blautia hansenii		100.0	0.53	URE 1			
41505102	**Very good identification**						**41505102**
					COC		
Blautia hansenii		100.0	0.71	BGL 10	99		
Lactobacillus fermentum		0.00	0.14		1		
41505103	**Good identification**						**41505103**
Blautia hansenii		100.0	0.37	URE 1			
41505110	**Good identification**						**41505110**
					COC		
Blautia hansenii		99.95	0.53	MLZ 1	99		
Eubacterium saburreum		0.06	0.13		1		
41505112	**Good identification**						**41505112**
Blautia hansenii		100.0	0.37	MLZ 1			
41505120	**Very good identification**						**41505120**
					COC	GRA	
Blautia hansenii		100.0	0.71	NAG 10	99	99	
Prevotella melaninogenica		0.00	-.04		1	1	
41505121	**Good identification**						**41505121**
Blautia hansenii		100.0	0.37	NAG 10			
41505122	**Good identification**						**41505122**
					COC	GRA	
Blautia hansenii		99.84	0.54	NAG 10	99	99	
Bacteroides fragilis		0.10	-.03	BGL 10	1	1	
Capnocytophaga ochracea		0.05	0.02		1	1	
41505130	**Good identification**						**41505130**
Blautia hansenii		100.0	0.37	MLZ 1			
41505140	**Good identification**						**41505140**
					COC	SPO	
Blautia hansenii		99.95	0.53	SOR 1	99	1	
Clostridium glycolicum		0.05	-.01		1	99	
41505142	**Good identification**						**41505142**
Blautia hansenii		100.0	0.37	SOR 1			
41505160	**Good identification**						**41505160**
Blautia hansenii		100.0	0.37	NAG 10			
41505300	**Good identification**						**41505300**
Blautia hansenii		100.0	0.53	RHA 1			
41505302	**Good identification**						**41505302**
Blautia hansenii		100.0	0.37	RHA 1			
41505320	**Good identification**						**41505320**
Blautia hansenii		100.0	0.37	RHA 1			

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
41505400	**Good identification**						**41505400**
Blautia hansenii		100.0	0.40	LAC 99			
41505500	**Very good identification**						**41505500**
					COC		
Blautia hansenii		99.98	0.74		99		
Bifidobacterium longum subsp. longum		0.02	0.21		1		
Lactobacillus fermentum		0.00	0.14		1		
41505501	**Good identification**						**41505501**
Blautia hansenii		100.0	0.40	URE 1			
41505502	**Very good identification**						**41505502**
					COC		
Blautia hansenii		99.83	0.58	BGL 10	99		
Bifidobacterium longum subsp. longum		0.15	0.21		1		
Lactobacillus fermentum		0.02	0.14		1		
41505510	**Good identification**						**41505510**
					COC		
Blautia hansenii		97.35	0.40	MLZ 1	99		
Bifidobacterium longum subsp. longum		2.65	0.25		1		
41505520	**Very good identification**						**41505520**
					COC	GRA	
Blautia hansenii		99.40	0.58	NAG 10	99	99	
Bifidobacterium longum subsp. longum		0.58	0.31		1	99	
Bacteroides vulgatus		0.02	-.08		1	1	
41505522	**Good identification**						**41505522**
					COC		
Blautia hansenii		95.03	0.42	NAG 10	99		
Bifidobacterium longum subsp. longum		4.97	0.31	BGL 10	1		
41505532	**Low discrimination**						**41505532**
					COC		
Bifidobacterium longum subsp. longum		89.82	0.35	SUC 99	1		
Blautia hansenii		10.18	0.07	MNS 87	99		
41505540	**Good identification**						**41505540**
Blautia hansenii		100.0	0.40	SOR 1			
41505700	**Good identification**						**41505700**
Blautia hansenii		100.0	0.40	RHA 1			
41507100	**Good identification**						**41507100**
Blautia hansenii		100.0	0.53	MAN 1			
41507102	**Good identification**						**41507102**
Blautia hansenii		100.0	0.37	MAN 1			
41507120	**Good identification**						**41507120**
Blautia hansenii		100.0	0.37	MAN 1			
41507500	**Good identification**						**41507500**
Blautia hansenii		100.0	0.40	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
41510000	**Low discrimination**							**41510000**
					GRA	SPO		
Clostridium sporogenes		63.70	0.53	RAF 1	99	99		
Clostridium botulinum biovar A		13.80	0.44		99	99		
Prevotella intermedia		7.85	0.42		1	1		
41510001	**Low discrimination**							**41510001**
Clostridium novyi biovar A		46.66	0.37	RAF 1				
Clostridium sporogenes		28.32	0.19	FRU 10				
Clostridium sordellii		12.09	0.10					
41510002	**Acceptable identification**							**41510002**
					GRA	SPO		
Clostridium sporogenes		92.89	0.45	RAF 1	99	99		
Fusobacterium mortiferum		3.60	0.36		1	1		
Atopobium minutum		1.10	0.22		99	1		
41510010	**Low discrimination**							**41510010**
Clostridium novyi biovar A		47.81	0.37	RAF 1				
Clostridium sporogenes		29.02	0.19	FRU 10				
Clostridium difficile		9.94	0.10					
41510011	**Good identification**							**41510011**
Clostridium novyi biovar A		100.0	0.37	RAF 1				
41510040	**Low discrimination**							**41510040**
Clostridium glycolicum		71.69	0.44	RAF 1				
Clostridium botulinum biovar A		18.38	0.35	XYL 95				
Clostridium novyi biovar A		3.15	0.34					
41510100	**Low discrimination**							**41510100**
					COC	GRA	SPO	
Fusobacterium mortiferum		75.32	0.59	MLT 11	1	1	1	
Blautia hansenii		20.50	0.32		99	99	1	
Clostridium sporogenes		2.80	0.19		1	99	99	
41510102	**Low discrimination**							**41510102**
					COC	GRA	SPO	
Fusobacterium mortiferum		70.78	0.42	MLT 11	1	1	1	
Blautia hansenii		19.27	0.15	BGL 10	99	99	1	
Clostridium sporogenes		7.91	0.11		1	99	99	
41511000	**Low discrimination**							**41511000**
					COC	GRA		
Fusobacterium mortiferum		39.27	0.55	MLT 11	1	1		
Prevotella intermedia		27.76	0.42		1	1		
Blautia hansenii		16.63	0.32		99	99		
41511002	**Low discrimination**							**41511002**
					COC	GRA		
Lactobacillus fermentum		61.24	0.48	ESL 1	1	99		
Fusobacterium mortiferum		22.42	0.39	SUC 87	1	1		
Blautia hansenii		9.49	0.15		99	99		
41511020	**Low discrimination**							**41511020**
					COC	GRA		
Lactobacillus fermentum		48.09	0.40	ESL 1	1	99		
Blautia hansenii		22.37	0.15	SUC 87	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella melaninogenica		15.47	0.12		1	1		
41511022	**Low discrimination**							**41511022**
					GRA	CAT		
Lactobacillus fermentum		62.70	0.40	ESL 1	99	1		
Prevotella oralis		14.82	0.14	SUC 87	1	1		
Capnocytophaga ochracea		12.12	0.12		1	90		
41511100	**Good identification**							**41511100**
					COC	GRA		
Blautia hansenii		93.48	0.66	FRU 1	99	99		
Fusobacterium mortiferum		5.20	0.62		1	1		
Lactobacillus fermentum		1.10	0.52		1	99		
41511102	**Low discrimination**							**41511102**
					COC	GRA		
Blautia hansenii		82.28	0.50	FRU 1	99	99		
Lactobacillus fermentum		8.75	0.52	BGL 10	1	99		
Fusobacterium mortiferum		4.58	0.45		1	1		
41511120	**Low discrimination**							**41511120**
					COC	CAT	GRA	
Blautia hansenii		55.68	0.50	FRU 1	99	1	99	
Prevotella melaninogenica		38.52	0.47	NAG 10	1	1	1	
Capnocytophaga ochracea		2.57	0.30		1	90	1	
41511122	**Low discrimination**							**41511122**
					COC	CAT	GRA	
Capnocytophaga ochracea		52.82	0.47	SUC 99	1	90	1	
Prevotella melaninogenica		16.75	0.34	MNS 90	1	1	1	
Blautia hansenii		14.12	0.33		99	1	99	
41511400	**Low discrimination**							**41511400**
					COC	GRA		
Lactobacillus fermentum		76.75	0.48	ESL 1	1	99		
Blautia hansenii		18.90	0.19	SUC 87	99	99		
Fusobacterium mortiferum		2.55	0.21		1	1		
41511402	**Good identification**							**41511402**
					COC			
Lactobacillus fermentum		97.34	0.48	ESL 1	1			
Blautia hansenii		2.66	0.03	SUC 87	99			
41511420	**Acceptable identification**							**41511420**
					COC			
Lactobacillus fermentum		90.33	0.40	ESL 1	1			
Blautia hansenii		7.41	0.03	SUC 87	99			
Bifidobacterium longum subsp. longum		2.26	0.05		1			
41511422	**Acceptable identification**							**41511422**
					GRA	CAT		
Lactobacillus fermentum		94.42	0.40	ESL 1	99	1		
Capnocytophaga ochracea		3.22	-.01	SUC 87	1	90		
Bifidobacterium longum subsp. longum		2.36	0.05		99	1		
41511500	**Acceptable identification**							**41511500**
					COC			
Blautia hansenii		92.67	0.53	FRU 1	99			
Lactobacillus fermentum		6.20	0.52		1			
Bifidobacterium longum subsp. longum		0.78	0.29		1			
41511502	**Low discrimination**							**41511502**
					COC			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		57.45	0.37	FRU 1	99			
Lactobacillus fermentum		34.61	0.52	BGL 10	1			
Bifidobacterium longum subsp. longum		4.38	0.29		1			
41511520	**Low discrimination**							**41511520**
					COC			
Blautia hansenii		62.58	0.37	FRU 1	99			
Bifidobacterium longum subsp. longum		19.07	0.40	NAG 10	1			
Lactobacillus fermentum		12.57	0.44		1			
41511530	**Good identification**							**41511530**
					COC	GRA	CAT	
Bifidobacterium longum subsp. longum		97.15	0.44	SUC 99	1	99	1	
Blautia hansenii		1.89	0.03	MNS 87	99	99	1	
Capnocytophaga ochracea		0.96	0.01		1	1	90	
41511532	**Acceptable identification**							**41511532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		90.63	0.44	SUC 99	99	1		
Capnocytophaga ochracea		8.07	0.17	MNS 87	1	90		
Parabacteroides distasonis		1.31	0.16		1	76		
41514000	**Acceptable identification**							**41514000**
					GRA	SPO		
Clostridium glycolicum		91.89	0.44	RAF 1	99	99		
Clostridium sporogenes		3.68	0.19	SOR 95	99	99		
Fusobacterium mortiferum		3.19	0.33		1	1		
41514040	**Very good identification**							**41514040**
Clostridium glycolicum		99.99	0.66	RAF 1				
Clostridium botulinum biovar A		0.01	0.01					
41514100	**Low discrimination**							**41514100**
					COC	GRA	SPO	
Fusobacterium mortiferum		54.07	0.40	MLT 11	1	1	1	
Blautia hansenii		34.50	0.19	XYL 7	99	99	1	
Clostridium glycolicum		6.75	0.10		1	99	99	
41515000	**Low discrimination**							**41515000**
					GRA	SPO		
Lactobacillus fermentum		35.27	0.44	ESL 1	99	1		
Clostridium glycolicum		31.83	0.28	SUC 87	99	99		
Fusobacterium mortiferum		14.90	0.36		1	1		
41515002	**Acceptable identification**							**41515002**
					COC	GRA		
Lactobacillus fermentum		91.45	0.44	ESL 1	1	99		
Fusobacterium mortiferum		4.29	0.20	SUC 87	1	1		
Blautia hansenii		4.26	0.03		99	99		
41515020	**Low discrimination**							**41515020**
					COC			
Lactobacillus fermentum		87.74	0.36	ESL 1	1			
Blautia hansenii		12.26	0.03	SUC 87	99			
41515022	**Low discrimination**							**41515022**
					GRA	CAT		
Lactobacillus fermentum		77.67	0.36	ESL 1	99	1		
Bacteroides fragilis		12.08	-.03	SUC 87	1	99		
Capnocytophaga ochracea		6.39	0.02		1	90		
41515040	**Good identification**							**41515040**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium glycolicum		100.0	0.50	RAF 1				
41515100	**Acceptable identification**							**41515100**
					COC	GRA		
Blautia hansenii		93.69	0.53	FRU 1	99	99		
Lactobacillus fermentum		3.68	0.48		1	99		
Fusobacterium mortiferum		2.22	0.43		1	1		
41515102	**Low discrimination**							**41515102**
					COC	GRA	CAT	
Blautia hansenii		68.02	0.37	FRU 1	99	99	1	
Lactobacillus fermentum		24.07	0.48	BGL 10	1	99	1	
Capnocytophaga ochracea		4.45	0.20		1	1	90	
41515120	**Low discrimination**							**41515120**
					COC	GRA	CAT	
Blautia hansenii		75.84	0.37	FRU 1	99	99	1	
Lactobacillus fermentum		8.94	0.40	NAG 10	1	99	1	
Capnocytophaga ochracea		4.96	0.20		1	1	90	
41515132	**Low discrimination**							**41515132**
					GRA	CAT		
Parabacteroides distasonis		54.30	0.36	MNS 98	1	76		
Capnocytophaga ochracea		32.36	0.20		1	90		
Bifidobacterium longum subsp. longum		7.78	0.18		99	1		
41515322	**Acceptable identification**							**41515322**
					CAT			
Parabacteroides distasonis		93.02	0.49	MNS 98	76			
Prevotella oralis		3.67	0.12		1			
Bacteroides fragilis		1.71	-.03		99			
41515332	**Good identification**							**41515332**
Parabacteroides distasonis		100.0	0.44	MNS 98				
41515400	**Acceptable identification**							**41515400**
					COC			
Lactobacillus fermentum		93.11	0.44	ESL 1	1			
Blautia hansenii		6.89	0.06	SUC 87	99			
41515402	**Good identification**							**41515402**
Lactobacillus fermentum		100.0	0.44	ESL 1				
41515420	**Low discrimination**							**41515420**
					GRA			
Lactobacillus fermentum		85.21	0.36	ESL 1	99			
Bifidobacterium longum subsp. longum		10.89	0.14	SUC 87	99			
Bacteroides vulgatus		3.90	-.08		1			
41515422	**Low discrimination**							**41515422**
Lactobacillus fermentum		88.67	0.36	ESL 1				
Bifidobacterium longum subsp. longum		11.33	0.14	SUC 87				
41515500	**Low discrimination**							**41515500**
					COC			
Blautia hansenii		72.95	0.40	FRU 1	99			
Lactobacillus fermentum		16.25	0.48		1			
Bifidobacterium longum subsp. longum		10.50	0.37		1			
41515502	**Low discrimination**							**41515502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Lactobacillus fermentum		41.36	0.48	ESL 1	1		
Bifidobacterium longum subsp. longum		26.72	0.37	SUC 87	1		
Blautia hansenii		20.63	0.24		99		
41515510	**Good identification**						**41515510**
					COC		
Bifidobacterium longum subsp. longum		95.20	0.41	SUC 99	1		
Blautia hansenii		3.92	0.06	MNS 87	99		
Lactobacillus fermentum		0.88	0.14		1		
41515512	**Low discrimination**						**41515512**
Bifidobacterium longum subsp. longum		78.35	0.41	SUC 99			
Bifidobacterium adolescentis		20.93	0.33	MNS 87			
Lactobacillus fermentum		0.72	0.14				
41515520	**Low discrimination**						**41515520**
					COC	GRA	
Bifidobacterium longum subsp. longum		59.07	0.48	SUC 99	1	99	
Bacteroides vulgatus		21.17	0.26	MNS 87	1	1	
Blautia hansenii		11.40	0.24		99	99	
41515522	**Low discrimination**						**41515522**
					GRA		
Bifidobacterium longum subsp. longum		70.59	0.48	SUC 99	99		
Parabacteroides distasonis		9.65	0.37	MNS 87	1		
Lactobacillus fermentum		9.11	0.40		99		
41515530	**Good identification**						**41515530**
					GRA		
Bifidobacterium longum subsp. longum		99.79	0.52	SUC 99	99		
Bacteroides vulgatus		0.21	-.08	MNS 87	1		
41515532	**Good identification**						**41515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		95.18	0.52	SUC 99	99	1	
Parabacteroides distasonis		4.11	0.33	MNS 87	1	76	
Capnocytophaga ochracea		0.71	0.07		1	90	
41515720	**Good identification**						**41515720**
					GRA		
Bacteroides vulgatus		98.85	0.42	SUC 96	1		
Eubacterium contortum		0.71	0.13	MNS 99	99		
Bifidobacterium longum subsp. longum		0.31	0.14		99		
41515722	**Low discrimination**						**41515722**
					GRA		
Parabacteroides distasonis		89.76	0.45	MNS 98	1		
Bacteroides vulgatus		7.13	0.08		1		
Bifidobacterium longum subsp. longum		2.21	0.14		99		
41515732	**Acceptable identification**						**41515732**
					GRA		
Parabacteroides distasonis		92.77	0.41	MNS 98	1		
Bifidobacterium longum subsp. longum		7.23	0.18		99		
41520000	**Low discrimination**						**41520000**
Clostridium sporogenes		68.83	0.58	RAF 1			
Clostridium botulinum biovar A		30.92	0.54				
Clostridium bifermentans		0.15	0.14				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41520002	**Good identification**						**41520002**
Clostridium sporogenes		98.66	0.50	RAF 1			
Clostridium botulinum biovar A		1.34	0.20				
41520040	**Acceptable identification**						**41520040**
Clostridium botulinum biovar A		92.66	0.45	RAF 1			
Clostridium sporogenes		6.97	0.24				
Clostridium novyi biovar A		0.37	0.16				
41521100	**Very good identification**						**41521100**
					COC	GRA	
Blautia hansenii		99.97	0.66	TRE 1	99	99	
Fusobacterium mortiferum		0.02	0.19		1	1	
Eubacterium saburreum		0.02	0.16		1	85	
41521102	**Good identification**						**41521102**
Blautia hansenii		100.0	0.50	TRE 1			
41521120	**Good identification**						**41521120**
					COC	GRA	
Blautia hansenii		99.92	0.50	TRE 1	99	99	
Prevotella melaninogenica		0.08	-.04	NAG 10	1	1	
41521500	**Good identification**						**41521500**
Blautia hansenii		100.0	0.53	TRE 1			
41521502	**Good identification**						**41521502**
Blautia hansenii		100.0	0.37	TRE 1			
41521520	**Good identification**						**41521520**
					COC		
Blautia hansenii		99.16	0.37	TRE 1	99		
Bifidobacterium longum subsp. longum		0.84	0.13	NAG 10	1		
41525100	**Good identification**						**41525100**
					COC		
Blautia hansenii		99.96	0.53	TRE 1	99		
Eubacterium saburreum		0.05	0.11		1		
41525102	**Good identification**						**41525102**
Blautia hansenii		100.0	0.37	TRE 1			
41525120	**Good identification**						**41525120**
Blautia hansenii		100.0	0.37	TRE 1			
41525500	**Good identification**						**41525500**
					COC		
Blautia hansenii		99.60	0.40	TRE 1	99		
Bifidobacterium longum subsp. longum		0.40	0.11		1		
41530000	**Low discrimination**						**41530000**
					GRA	SPO	
Clostridium sporogenes		78.71	0.45	RAF 1	99	99	
Clostridium botulinum biovar A		12.79	0.34		99	99	
Fusobacterium mortiferum		8.20	0.43		1	1	
41530002	**Good identification**						**41530002**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sporogenes		96.19	0.37	RAF 1	99	99		
Fusobacterium mortiferum		3.34	0.27		1	1		
Clostridium botulinum biovar A		0.47	-.01		99	99		
41530100	**Acceptable identification**							**41530100**
					COC	GRA	SPO	
Fusobacterium mortiferum		94.56	0.50	MLT 11	1	1	1	
Clostridium sporogenes		3.93	0.11		1	99	99	
Blautia hansenii		0.87	-.03		99	99	1	
41531000	**Low discrimination**							**41531000**
					SPO	GRA		
Fusobacterium mortiferum		86.95	0.46	MLT 11	1	1		
Clostridium sporogenes		5.62	0.11		99	99		
Eubacterium saburreum		2.31	0.18		1	85		
41531100	**Low discrimination**							**41531100**
					COC	GRA		
Fusobacterium mortiferum		60.80	0.53	MLT 11	1	1		
Blautia hansenii		36.94	0.32		99	99		
Eubacterium saburreum		1.61	0.24		1	85		
41531102	**Low discrimination**							**41531102**
					COC	GRA		
Fusobacterium mortiferum		58.33	0.36	MLT 11	1	1		
Blautia hansenii		35.44	0.15	BGL 10	99	99		
Lactobacillus fermentum		3.77	0.18		1	99		
41531122	**Low discrimination**							**41531122**
					GRA			
Parabacteroides distasonis		47.69	0.36	MNS 98	1			
Clostridium perfringens		21.45	0.20	XYL 90	99			
Prevotella oralis		13.44	0.14		1			
41531322	**Good identification**							**41531322**
Parabacteroides distasonis		96.13	0.44	MNS 98				
Prevotella oralis		3.87	0.07	XYL 90				
41531332	**Good identification**							**41531332**
Parabacteroides distasonis		100.0	0.39	MNS 98				
41531722	**Good identification**							**41531722**
Parabacteroides distasonis		100.0	0.40	MNS 98				
41531732	**Good identification**							**41531732**
Parabacteroides distasonis		100.0	0.36	MNS 98				
41535122	**Good identification**							**41535122**
Parabacteroides distasonis		97.81	0.52	MNS 98				
Bacteroides fragilis		1.18	-.03					
Capnocytophaga ochracea		0.63	0.02					
41535132	**Good identification**							**41535132**
					GRA			
Parabacteroides distasonis		99.22	0.47	MNS 98	1			
Bifidobacterium longum subsp. longum		0.78	0.07		99			
41535322	**Very good identification**							**41535322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.60	MNS 98			
41535332	**Good identification**						**41535332**
Parabacteroides distasonis		100.0	0.56	MNS 98			
41535520	**Acceptable identification**						**41535520**
					GRA		
Bifidobacterium longum subsp. longum		92.18	0.37	SUC 99	99		
Bacteroides vulgatus		5.51	0.02	MNS 87	1		
Parabacteroides distasonis		2.32	0.14		1		
41535522	**Low discrimination**						**41535522**
					GRA		
Parabacteroides distasonis		71.35	0.48	MNS 98	1		
Bifidobacterium longum subsp. longum		28.65	0.37		99		
41535530	**Good identification**						**41535530**
					GRA		
Bifidobacterium longum subsp. longum		99.21	0.41	SUC 99	99		
Parabacteroides distasonis		0.79	0.10	MNS 87	1		
41535532	**Low discrimination**						**41535532**
					GRA		
Bifidobacterium longum subsp. longum		55.95	0.41	SUC 99	99		
Parabacteroides distasonis		44.05	0.44	MNS 87	1		
41535722	**Very good identification**						**41535722**
Parabacteroides distasonis		99.71	0.57	MNS 98			
Bacteroides thetaiotaomicron		0.29	-.03				
41535732	**Good identification**						**41535732**
					GRA		
Parabacteroides distasonis		99.01	0.52	MNS 98	1		
Clostridium clostridioforme		0.57	0.08		50		
Bifidobacterium longum subsp. longum		0.42	0.07		99		
41541100	**Very good identification**						**41541100**
					COC	GRA	
Blautia hansenii		99.98	0.66	CEL 1	99	99	
Fusobacterium mortiferum		0.02	0.19		1	1	
41541102	**Good identification**						**41541102**
Blautia hansenii		100.0	0.50	CEL 1			
41541120	**Good identification**						**41541120**
					COC	GRA	
Blautia hansenii		99.15	0.50	CEL 1	99	99	
Prevotella melaninogenica		0.85	0.14	NAG 10	1	1	
41541500	**Good identification**						**41541500**
Blautia hansenii		100.0	0.53	CEL 1			
41541502	**Good identification**						**41541502**
Blautia hansenii		100.0	0.37	CEL 1			
41541520	**Good identification**						**41541520**
Blautia hansenii		100.0	0.37	CEL 1			
41545100	**Good identification**						**41545100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Blautia hansenii		100.0	0.53	CEL 1				
41545102	**Good identification**							**41545102**
Blautia hansenii		100.0	0.37	CEL 1				
41545120	**Good identification**							**41545120**
Blautia hansenii		100.0	0.37	CEL 1				
41545500	**Good identification**							**41545500**
Blautia hansenii		100.0	0.40	CEL 1				
41550000	**Low discrimination**							**41550000**
					GRA	SPO		
Fusobacterium mortiferum		74.12	0.44	MLT 11	1	1		
Clostridium sporogenes		19.32	0.19		99	99		
Clostridium botulinum biovar A		4.19	0.10		99	99		
41550100	**Good identification**							**41550100**
					COC	GRA		
Fusobacterium mortiferum		99.18	0.51	MLT 11	1	1		
Blautia hansenii		0.82	-.03		99	99		
41551000	**Good identification**							**41551000**
					COC	GRA		
Fusobacterium mortiferum		95.84	0.47	MLT 11	1	1		
Prevotella intermedia		2.05	0.08		1	1		
Blautia hansenii		1.23	-.03		99	99		
41551100	**Low discrimination**							**41551100**
					COC	CAT	GRA	
Fusobacterium mortiferum		63.78	0.54	MLT 11	1	5	1	
Blautia hansenii		34.72	0.32		99	1	99	
Capnocytophaga ochracea		0.49	0.04		1	90	1	
41551102	**Low discrimination**							**41551102**
					CAT			
Fusobacterium mortiferum		32.56	0.37	MLT 11	5			
Prevotella oralis		26.33	0.22	BGL 10	1			
Capnocytophaga ochracea		20.26	0.20		90			
41551122	**Low discrimination**							**41551122**
					CAT			
Prevotella oralis		69.92	0.43	SUC 92	1			
Capnocytophaga ochracea		25.48	0.36	MNS 99	90			
Prevotella melaninogenica		3.59	0.18		1			
41551322	**Acceptable identification**							**41551322**
					CAT			
Prevotella oralis		94.21	0.37	SUC 92	1			
Parabacteroides distasonis		4.98	0.28	MNS 99	76			
Capnocytophaga ochracea		0.81	0.02		90			
41555322	**Low discrimination**							**41555322**
					CAT			
Parabacteroides distasonis		78.46	0.44	MNS 98	76			
Prevotella oralis		20.39	0.22		1			
Bacteroides fragilis		1.15	-.09		99			
41555332	**Good identification**							**41555332**
Parabacteroides distasonis		100.0	0.40	MNS 98				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41555502	**Acceptable identification**						**41555502**
Bifidobacterium adolescentis		93.06	0.37	SUC 99			
Actinomyces israelii		5.10	0.22	MNS 87			
Lactobacillus fermentum		1.84	0.14				
41555512	**Good identification**						**41555512**
Bifidobacterium adolescentis		98.25	0.39	SUC 99			
Bifidobacterium longum subsp. longum		1.75	0.07	MNS 87			
41555722	**Good identification**						**41555722**
Parabacteroides distasonis		100.0	0.41	MNS 98			
41555732	**Good identification**						**41555732**
Parabacteroides distasonis		100.0	0.36	MNS 98			
41557102	**Low discrimination**						**41557102**
Actinomyces israelii		75.64	0.36	SUC 98			
Bifidobacterium adolescentis		24.36	0.22	SAL 92			
41557502	**Low discrimination**						**41557502**
Bifidobacterium adolescentis		72.48	0.36	SUC 99			
Actinomyces israelii		27.52	0.35	MNS 87			
41557512	**Good identification**						**41557512**
Bifidobacterium adolescentis		95.57	0.37	SUC 99			
Actinomyces israelii		4.44	0.21	MNS 87			
41570000	**Low discrimination**						**41570000**
					GRA	SPO	
Fusobacterium mortiferum		74.73	0.35	MLT 11	1	1	
Clostridium sporogenes		21.74	0.11		99	99	
Clostridium botulinum biovar A		3.53	-.01		99	99	
41570100	**Good identification**						**41570100**
Fusobacterium mortiferum		100.0	0.42	MLT 11			
41571000	**Good identification**						**41571000**
Fusobacterium mortiferum		100.0	0.38	MLT 11			
41571100	**Good identification**						**41571100**
					COC	GRA	
Fusobacterium mortiferum		96.87	0.45	MLT 11	1	1	
Blautia hansenii		1.78	-.03		99	99	
Atopobium parvulum		1.35	0.01		99	99	
41571322	**Low discrimination**						**41571322**
Parabacteroides distasonis		79.04	0.39	MNS 98			
Prevotella oralis		20.96	0.17	XYL 90			
41571722	**Good identification**						**41571722**
Parabacteroides distasonis		100.0	0.36	MNS 98			
41575122	**Good identification**						**41575122**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		96.61	0.47	MNS 98	76	
Prevotella oralis		2.46	0.07		1	
Bacteroides fragilis		0.93	-.09		99	
41575132	**Good identification**					**41575132**
Parabacteroides distasonis		100.0	0.43	MNS 98		
41575322	**Good identification**					**41575322**
Parabacteroides distasonis		99.64	0.56	MNS 98		
Prevotella oralis		0.36	0.01			
41575332	**Good identification**					**41575332**
Parabacteroides distasonis		100.0	0.51	MNS 98		
41575512	**Good identification**					**41575512**
Bifidobacterium adolescentis		96.36	0.36	SUC 99		
Actinomyces israelii		3.64	0.18	MNS 87		
41575522	**Good identification**					**41575522**
Parabacteroides distasonis		100.0	0.44	MNS 98		
41575532	**Good identification**					**41575532**
					GRA	
Parabacteroides distasonis		97.67	0.39	MNS 98	1	
Bifidobacterium longum subsp. longum		2.33	0.07		99	
41575722	**Good identification**					**41575722**
Parabacteroides distasonis		98.45	0.52	MNS 98		
Bacteroides thetaiotaomicron		1.14	0.03			
Clostridium clostridioforme		0.41	0.06			
41575732	**Good identification**					**41575732**
Parabacteroides distasonis		95.79	0.47	MNS 98		
Clostridium clostridioforme		3.41	0.17			
Bacteroides thetaiotaomicron		0.80	-.04			
41577002	**Good identification**					**41577002**
Actinomyces israelii		100.0	0.36	SUC 98		
41577100	**Good identification**					**41577100**
Actinomyces israelii		100.0	0.43	SUC 98		
41577102	**Acceptable identification**					**41577102**
Actinomyces israelii		94.60	0.46	SUC 98		
Bifidobacterium adolescentis		5.40	0.19	SAL 92		
41577140	**Good identification**					**41577140**
Actinomyces israelii		100.0	0.36	SUC 98		
41577142	**Good identification**					**41577142**
Actinomyces israelii		100.0	0.39	SUC 98		
41577500	**Good identification**					**41577500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.42	SUC 98			
41577502	**Low discrimination**						**41577502**
Actinomyces israelii		68.18	0.45	SUC 98			
Bifidobacterium adolescentis		31.83	0.33	SAL 92			
41577542	**Low discrimination**						**41577542**
Actinomyces israelii		87.08	0.38	SUC 98			
Bifidobacterium adolescentis		12.92	0.17	SAL 92			
41600100	**Good identification**						**41600100**
					COC	GRA	
Fusobacterium mortiferum		98.30	0.45	FRU 99	1	1	
Blautia hansenii		1.70	-.03		99	99	
41601000	**Very good identification**						**41601000**
					GRA		
Eubacterium tortuosum		98.59	0.58	SUC 99	99		
Fusobacterium mortiferum		1.22	0.42		1		
Atopobium minutum		0.09	0.17		99		
41601002	**Good identification**						**41601002**
					GRA		
Eubacterium tortuosum		98.87	0.45	SUC 99	99		
Fusobacterium mortiferum		0.77	0.25		1		
Atopobium minutum		0.36	0.14		99		
41601020	**Good identification**						**41601020**
Eubacterium tortuosum		99.82	0.45	SUC 99			
Atopobium minutum		0.18	0.09				
41610000	**Very good identification**						**41610000**
					GRA	SPO	
Fusobacterium mortiferum		97.57	0.73		1	1	
Clostridium difficile		1.06	0.23		99	99	
Eubacterium limosum		0.44	0.15		99	1	
41610001	**Acceptable identification**						**41610001**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	URE 1	1	1	
Clostridium novyi biovar A		7.27	0.19		99	99	
41610002	**Very good identification**						**41610002**
					GRA	SPO	
Fusobacterium mortiferum		97.25	0.57	BGL 10	1	1	
Atopobium minutum		1.89	0.22		99	1	
Clostridium sporogenes		0.87	0.06		99	99	
41610010	**Low discrimination**						**41610010**
					GRA	SPO	
Clostridium difficile		82.92	0.35	RAF 1	99	99	
Fusobacterium mortiferum		15.84	0.39	MAN 90	1	1	
Clostridium novyi biovar A		1.24	0.19		99	99	
41610020	**Acceptable identification**						**41610020**
					GRA		
Fusobacterium mortiferum		90.36	0.39	NAG 1	1		
Atopobium minutum		9.64	0.17		99		
41610040	**Low discrimination**						**41610040**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		78.14	0.39	SOR 1	1	1	
Clostridium glycolicum		9.97	0.09		99	99	
Clostridium difficile		4.41	0.02		99	99	
41610100	**Very good identification**						**41610100**
Fusobacterium mortiferum		100.0	0.79				
41610101	**Good identification**						**41610101**
Fusobacterium mortiferum		100.0	0.45	URE 1			
41610102	**Very good identification**						**41610102**
Fusobacterium mortiferum		100.0	0.63	BGL 10			
41610110	**Good identification**						**41610110**
					GRA	SPO	
Fusobacterium mortiferum		97.78	0.45	MLZ 1	1	1	
Clostridium difficile		2.22	0.01		99	99	
41610120	**Good identification**						**41610120**
Fusobacterium mortiferum		100.0	0.45	NAG 1			
41610140	**Good identification**						**41610140**
Fusobacterium mortiferum		100.0	0.45	SOR 1			
41610200	**Good identification**						**41610200**
Fusobacterium mortiferum		100.0	0.39	RHA 1			
41610300	**Good identification**						**41610300**
Fusobacterium mortiferum		100.0	0.45	RHA 1			
41610400	**Good identification**						**41610400**
Fusobacterium mortiferum		100.0	0.39	ARA 1			
41610500	**Good identification**						**41610500**
Fusobacterium mortiferum		100.0	0.45	ARA 1			
41611000	**Very good identification**						**41611000**
					GRA		
Fusobacterium mortiferum		99.11	0.76		1		
Eubacterium tortuosum		0.82	0.24		99		
Atopobium minutum		0.07	0.17		99		
41611001	**Good identification**						**41611001**
Fusobacterium mortiferum		100.0	0.42	URE 1			
41611002	**Very good identification**						**41611002**
					GRA		
Fusobacterium mortiferum		98.29	0.60	BGL 10	1		
Eubacterium tortuosum		1.29	0.11		99		
Atopobium minutum		0.42	0.14		99		
41611010	**Good identification**						**41611010**
					GRA	SPO	
Fusobacterium mortiferum		96.60	0.42	MLZ 1	1	1	
Clostridium difficile		3.41	0.01		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41611020	**Low discrimination**						**41611020**
					GRA		
Fusobacterium mortiferum		85.65	0.42	NAG 1	1		
Eubacterium tortuosum		12.33	0.11		99		
Atopobium minutum		2.03	0.09		99		
41611040	**Good identification**						**41611040**
Fusobacterium mortiferum		100.0	0.42	SOR 1			
41611100	**Very good identification**						**41611100**
					COC	GRA	
Fusobacterium mortiferum		99.99	0.82		1	1	
Blautia hansenii		0.01	-.03		99	99	
41611101	**Good identification**						**41611101**
Fusobacterium mortiferum		100.0	0.48	URE 1			
41611102	**Very good identification**						**41611102**
					GRA		
Fusobacterium mortiferum		99.94	0.66	BGL 10	1		
Bifidobacterium longum subsp. infantis		0.06	-.07		99		
41611110	**Good identification**						**41611110**
Fusobacterium mortiferum		100.0	0.48	MLZ 1			
41611120	**Good identification**						**41611120**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.48	NAG 1	1	1	
Clostridium septicum		0.77	-.06		99	99	
41611140	**Good identification**						**41611140**
Fusobacterium mortiferum		100.0	0.48	SOR 1			
41611200	**Good identification**						**41611200**
Fusobacterium mortiferum		100.0	0.42	RHA 1			
41611300	**Good identification**						**41611300**
Fusobacterium mortiferum		100.0	0.48	RHA 1			
41611400	**Good identification**						**41611400**
Fusobacterium mortiferum		100.0	0.42	ARA 1			
41611500	**Good identification**						**41611500**
Fusobacterium mortiferum		100.0	0.48	ARA 1			
41612000	**Low discrimination**						**41612000**
					GRA	SPO	
Clostridium difficile		71.76	0.40	RAF 1	99	99	
Eubacterium limosum		20.58	0.29		99	1	
Fusobacterium mortiferum		7.44	0.39		1	1	
41612010	**Good identification**						**41612010**
					SPO		
Clostridium difficile		99.94	0.52	RAF 1	99		
Eubacterium limosum		0.06	-.06		1		
41612100	**Acceptable identification**						**41612100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		94.90	0.45	MAN 1	1	1	
Clostridium difficile		3.96	0.06		99	99	
Eubacterium limosum		1.14	-.06		99	1	
41613000	**Acceptable identification**						**41613000**
					GRA	SPO	
Fusobacterium mortiferum		92.28	0.42	MAN 1	1	1	
Clostridium difficile		6.00	0.06		99	99	
Eubacterium limosum		1.72	-.06		99	1	
41613100	**Good identification**						**41613100**
Fusobacterium mortiferum		100.0	0.48	MAN 1			
41614000	**Good identification**						**41614000**
					GRA	SPO	
Fusobacterium mortiferum		97.59	0.54	XYL 7	1	1	
Clostridium glycolicum		1.67	0.09		99	99	
Clostridium difficile		0.74	0.02		99	99	
41614002	**Good identification**						**41614002**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
41614100	**Very good identification**						**41614100**
Fusobacterium mortiferum		100.0	0.60	XYL 7			
41614102	**Good identification**						**41614102**
Fusobacterium mortiferum		100.0	0.44	XYL 7			
41615000	**Very good identification**						**41615000**
Fusobacterium mortiferum		100.0	0.57	XYL 7			
41615002	**Good identification**						**41615002**
Fusobacterium mortiferum		100.0	0.40	XYL 7			
41615100	**Very good identification**						**41615100**
Fusobacterium mortiferum		100.0	0.63	XYL 7			
41615102	**Good identification**						**41615102**
Fusobacterium mortiferum		100.0	0.47	XYL 7			
41615322	**Good identification**						**41615322**
					GRA		
Parabacteroides distasonis		97.61	0.40	MNS 98	1		
Eubacterium contortum		2.39	0.01		99		
41615332	**Good identification**						**41615332**
Parabacteroides distasonis		100.0	0.36	MNS 98			
41615722	**Good identification**						**41615722**
					GRA		
Parabacteroides distasonis		96.84	0.36	MNS 98	1		
Eubacterium contortum		3.17	0.00		99		
41620100	**Good identification**						**41620100**
Fusobacterium mortiferum		100.0	0.36	FRU 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
41621100	**Good identification**						**41621100**
					COC	GRA	
Fusobacterium mortiferum		96.28	0.39	FRU 99	1	1	
Blautia hansenii		3.73	-.03		99	99	
41630000	**Very good identification**						**41630000**
					GRA	SPO	
Fusobacterium mortiferum		99.28	0.64		1	1	
Clostridium sporogenes		0.33	0.06		99	99	
Clostridium difficile		0.27	0.04		99	99	
41630002	**Good identification**						**41630002**
					GRA	SPO	
Fusobacterium mortiferum		99.01	0.48	BGL 10	1	1	
Clostridium sporogenes		0.99	-.02		99	99	
41630100	**Very good identification**						**41630100**
Fusobacterium mortiferum		100.0	0.70				
41630101	**Good identification**						**41630101**
Fusobacterium mortiferum		100.0	0.36	URE 1			
41630102	**Good identification**						**41630102**
Fusobacterium mortiferum		100.0	0.54	BGL 10			
41630110	**Good identification**						**41630110**
Fusobacterium mortiferum		100.0	0.36	MLZ 1			
41630120	**Good identification**						**41630120**
Fusobacterium mortiferum		100.0	0.36	NAG 1			
41630140	**Good identification**						**41630140**
Fusobacterium mortiferum		100.0	0.36	SOR 1			
41630300	**Good identification**						**41630300**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
41630500	**Good identification**						**41630500**
Fusobacterium mortiferum		100.0	0.36	ARA 1			
41631000	**Very good identification**						**41631000**
Fusobacterium mortiferum		100.0	0.67				
41631002	**Good identification**						**41631002**
Fusobacterium mortiferum		100.0	0.51	BGL 10			
41631100	**Very good identification**						**41631100**
Fusobacterium mortiferum		100.0	0.73				
41631101	**Good identification**						**41631101**
Fusobacterium mortiferum		100.0	0.39	URE 1			
41631102	**Very good identification**						**41631102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.57	BGL 10			
41631110	**Good identification**						**41631110**
Fusobacterium mortiferum		100.0	0.39	MLZ 1			
41631120	**Acceptable identification**						**41631120**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	NAG 1	1	1	
Clostridium septicum		7.27	0.03		99	99	
41631140	**Good identification**						**41631140**
Fusobacterium mortiferum		100.0	0.39	SOR 1			
41631300	**Good identification**						**41631300**
Fusobacterium mortiferum		100.0	0.39	RHA 1			
41631322	**Good identification**						**41631322**
Parabacteroides distasonis		100.0	0.35	MNS 98			
41631500	**Good identification**						**41631500**
Fusobacterium mortiferum		100.0	0.39	ARA 1			
41632100	**Good identification**						**41632100**
Fusobacterium mortiferum		100.0	0.36	MAN 1			
41633100	**Good identification**						**41633100**
Fusobacterium mortiferum		100.0	0.39	MAN 1			
41634000	**Good identification**						**41634000**
Fusobacterium mortiferum		100.0	0.45	XYL 7			
41634100	**Good identification**						**41634100**
Fusobacterium mortiferum		100.0	0.51	XYL 7			
41635000	**Good identification**						**41635000**
Fusobacterium mortiferum		100.0	0.48	XYL 7			
41635100	**Good identification**						**41635100**
Fusobacterium mortiferum		100.0	0.54	XYL 7			
41635102	**Good identification**						**41635102**
Fusobacterium mortiferum		100.0	0.38	XYL 7			
41635122	**Good identification**						**41635122**
Parabacteroides distasonis		100.0	0.43	MNS 98			
41635132	**Good identification**						**41635132**
Parabacteroides distasonis		100.0	0.39	MNS 98			
41635322	**Good identification**						**41635322**
Parabacteroides distasonis		100.0	0.51	MNS 98			
41635332	**Good identification**						**41635332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.47	MNS 98			
41635522	**Good identification**						**41635522**
Parabacteroides distasonis		100.0	0.40	MNS 98			
41635722	**Good identification**						**41635722**
Parabacteroides distasonis		100.0	0.48	MNS 98			
41635732	**Good identification**						**41635732**
Parabacteroides distasonis		100.0	0.43	MNS 98			
41640100	**Good identification**						**41640100**
Fusobacterium mortiferum		100.0	0.37	FRU 99			
41641000	**Good identification**						**41641000**
					GRA		
Eubacterium tortuosum		96.42	0.41	SUC 99	99		
Fusobacterium mortiferum		3.58	0.34	CEL 10	1		
41641100	**Low discrimination**						**41641100**
					COC	GRA	
Fusobacterium mortiferum		86.87	0.40	FRU 99	1	1	
Eubacterium tortuosum		10.12	0.07		1	99	
Blautia hansenii		3.01	-.03		99	99	
41650000	**Very good identification**						**41650000**
					GRA	SPO	
Fusobacterium mortiferum		99.83	0.65		1	1	
Clostridium difficile		0.17	0.02		99	99	
41650002	**Good identification**						**41650002**
Fusobacterium mortiferum		100.0	0.48	BGL 10			
41650100	**Very good identification**						**41650100**
Fusobacterium mortiferum		100.0	0.71				
41650101	**Good identification**						**41650101**
Fusobacterium mortiferum		100.0	0.37	URE 1			
41650102	**Good identification**						**41650102**
Fusobacterium mortiferum		100.0	0.55	BGL 10			
41650110	**Good identification**						**41650110**
Fusobacterium mortiferum		100.0	0.37	MLZ 1			
41650120	**Good identification**						**41650120**
Fusobacterium mortiferum		100.0	0.37	NAG 1			
41650140	**Good identification**						**41650140**
Fusobacterium mortiferum		100.0	0.37	SOR 1			
41650300	**Good identification**						**41650300**
Fusobacterium mortiferum		100.0	0.37	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41650500	**Good identification**						**41650500**
Fusobacterium mortiferum		100.0	0.37	ARA 1			
41651000	**Very good identification**						**41651000**
					GRA		
Fusobacterium mortiferum		99.73	0.68		1		
Eubacterium tortuosum		0.27	0.07		99		
41651002	**Good identification**						**41651002**
					GRA		
Fusobacterium mortiferum		99.57	0.51	BGL 10	1		
Eubacterium tortuosum		0.43	-.06		99		
41651100	**Very good identification**						**41651100**
Fusobacterium mortiferum		100.0	0.74				
41651101	**Good identification**						**41651101**
Fusobacterium mortiferum		100.0	0.40	URE 1			
41651102	**Very good identification**						**41651102**
Fusobacterium mortiferum		100.0	0.58	BGL 10			
41651110	**Good identification**						**41651110**
Fusobacterium mortiferum		100.0	0.40	MLZ 1			
41651120	**Acceptable identification**						**41651120**
					GRA	SPO	
Fusobacterium mortiferum		90.26	0.40	NAG 1	1	1	
Clostridium septicum		6.69	0.03		99	99	
Clostridium paraputrificum		3.05	-.06		99	99	
41651140	**Good identification**						**41651140**
Fusobacterium mortiferum		100.0	0.40	SOR 1			
41651300	**Good identification**						**41651300**
Fusobacterium mortiferum		100.0	0.40	RHA 1			
41651500	**Good identification**						**41651500**
Fusobacterium mortiferum		100.0	0.40	ARA 1			
41652100	**Good identification**						**41652100**
Fusobacterium mortiferum		100.0	0.37	MAN 1			
41653100	**Good identification**						**41653100**
Fusobacterium mortiferum		100.0	0.40	MAN 1			
41654000	**Good identification**						**41654000**
Fusobacterium mortiferum		100.0	0.46	XYL 7			
41654100	**Good identification**						**41654100**
Fusobacterium mortiferum		100.0	0.52	XYL 7			
41654102	**Good identification**						**41654102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.35	XYL 7		
41655000	**Good identification**					**41655000**
Fusobacterium mortiferum		100.0	0.49	XYL 7		
41655100	**Good identification**					**41655100**
Fusobacterium mortiferum		100.0	0.55	XYL 7		
41655102	**Good identification**					**41655102**
Fusobacterium mortiferum		100.0	0.38	XYL 7		
41655322	**Good identification**					**41655322**
Parabacteroides distasonis		100.0	0.36	MNS 98		
41670000	**Good identification**					**41670000**
Fusobacterium mortiferum		100.0	0.56			
41670002	**Good identification**					**41670002**
Fusobacterium mortiferum		100.0	0.39	BGL 10		
41670100	**Very good identification**					**41670100**
Fusobacterium mortiferum		100.0	0.62			
41670102	**Good identification**					**41670102**
Fusobacterium mortiferum		100.0	0.46	BGL 10		
41671000	**Very good identification**					**41671000**
Fusobacterium mortiferum		100.0	0.59			
41671002	**Good identification**					**41671002**
Fusobacterium mortiferum		100.0	0.42	BGL 10		
41671100	**Very good identification**					**41671100**
Fusobacterium mortiferum		100.0	0.65			
41671102	**Good identification**					**41671102**
Fusobacterium mortiferum		100.0	0.49	BGL 10		
41674000	**Good identification**					**41674000**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
41674100	**Good identification**					**41674100**
Fusobacterium mortiferum		100.0	0.43	XYL 7		
41675000	**Good identification**					**41675000**
Fusobacterium mortiferum		100.0	0.40	XYL 7		
41675100	**Good identification**					**41675100**
					GRA	
Fusobacterium mortiferum		98.96	0.46	XYL 7	1	
Actinomyces israelii		1.05	0.14		99	
41675122	**Good identification**					**41675122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.39	MNS 98			
41675322	**Good identification**						**41675322**
Parabacteroides distasonis		100.0	0.47	MNS 98			
41675332	**Good identification**						**41675332**
Parabacteroides distasonis		100.0	0.42	MNS 98			
41675722	**Good identification**						**41675722**
Parabacteroides distasonis		100.0	0.43	MNS 98			
41675732	**Good identification**						**41675732**
Parabacteroides distasonis		100.0	0.39	MNS 98			
41701100	**Very good identification**						**41701100**
					COC	GRA	
Blautia hansenii		99.89	0.66	SAL 1	99	99	
Fusobacterium mortiferum		0.11	0.33		1	1	
41701102	**Good identification**						**41701102**
					COC	GRA	
Blautia hansenii		99.84	0.50	SAL 1	99	99	
Fusobacterium mortiferum		0.11	0.16	BGL 10	1	1	
Bifidobacterium longum subsp. infantis		0.05	-.07		1	99	
41701120	**Good identification**						**41701120**
					COC	GRA	
Blautia hansenii		99.92	0.50	SAL 1	99	99	
Prevotella melaninogenica		0.08	-.04	NAG 10	1	1	
41701500	**Good identification**						**41701500**
Blautia hansenii		100.0	0.53	SAL 1			
41701502	**Good identification**						**41701502**
Blautia hansenii		100.0	0.37	SAL 1			
41701520	**Good identification**						**41701520**
Blautia hansenii		100.0	0.37	SAL 1			
41705100	**Good identification**						**41705100**
					COC	GRA	
Blautia hansenii		99.95	0.53	SAL 1	99	99	
Fusobacterium mortiferum		0.05	0.13		1	1	
41705102	**Good identification**						**41705102**
Blautia hansenii		100.0	0.37	SAL 1			
41705120	**Good identification**						**41705120**
Blautia hansenii		100.0	0.37	SAL 1			
41705500	**Good identification**						**41705500**
Blautia hansenii		100.0	0.40	SAL 1			
41710000	**Good identification**						**41710000**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		92.86	0.57	MLT 11	1	1	
Clostridium sporogenes		4.16	0.19		99	99	
Clostridium bifermentans		0.98	0.10		99	99	
41710002	**Low discrimination**						**41710002**
					GRA	SPO	
Fusobacterium mortiferum		88.16	0.41	MLT 11	1	1	
Clostridium sporogenes		11.84	0.11	BGL 10	99	99	
41710100	**Very good identification**						**41710100**
					COC	GRA	
Fusobacterium mortiferum		99.86	0.64	MLT 11	1	1	
Blautia hansenii		0.14	-.03		99	99	
41710102	**Good identification**						**41710102**
					GRA		
Fusobacterium mortiferum		99.27	0.47	MLT 11	1		
Bifidobacterium longum subsp. infantis		0.73	-.07	BGL 10	99		
41711000	**Very good identification**						**41711000**
					COC	GRA	
Fusobacterium mortiferum		99.26	0.60	MLT 11	1	1	
Prevotella intermedia		0.37	0.08		1	1	
Blautia hansenii		0.22	-.03		99	99	
41711002	**Good identification**						**41711002**
					GRA		
Fusobacterium mortiferum		97.50	0.44	MLT 11	1		
Lactobacillus fermentum		1.39	0.14	BGL 10	99		
Bifidobacterium longum subsp. infantis		1.12	-.07		99		
41711100	**Good identification**						**41711100**
					COC	GRA	
Fusobacterium mortiferum		90.91	0.67	MLT 11	1	1	
Blautia hansenii		8.50	0.32		99	99	
Bifidobacterium longum subsp. infantis		0.49	0.10		1	99	
41711102	**Low discrimination**						**41711102**
					COC	GRA	
Fusobacterium mortiferum		59.60	0.50	MLT 11	1	1	
Bifidobacterium longum subsp. infantis		29.04	0.27	BGL 10	1	99	
Blautia hansenii		5.57	0.15		99	99	
41711122	**Low discrimination**						**41711122**
					GRA		
Prevotella oralis		57.25	0.38	SUC 92	1		
Bifidobacterium longum subsp. infantis		37.48	0.32	MNS 99	99		
Parabacteroides distasonis		1.73	0.25		1		
41714000	**Low discrimination**						**41714000**
					GRA	SPO	
Fusobacterium mortiferum		86.95	0.38	MLT 11	1	1	
Clostridium glycolicum		13.05	0.10	XYL 7	99	99	
41714100	**Good identification**						**41714100**
Fusobacterium mortiferum		100.0	0.44	MLT 11			
41715000	**Good identification**						**41715000**
Fusobacterium mortiferum		100.0	0.41	MLT 11			
41715100	**Low discrimination**						**41715100**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		60.35	0.47	MLT 11	1	1	
Eubacterium contortum		21.38	0.27	XYL 7	1	99	
Blautia hansenii		13.23	0.19		99	99	
41715132	**Good identification**						**41715132**
					GRA		
Parabacteroides distasonis		96.44	0.37	MNS 98	1		
Bifidobacterium longum subsp. infantis		3.56	-.04		99		
41715300	**Very good identification**						**41715300**
					GRA		
Eubacterium contortum		99.97	0.61	SUC 99	99		
Fusobacterium mortiferum		0.03	0.13		1		
41715302	**Good identification**						**41715302**
					GRA		
Eubacterium contortum		99.74	0.48	SUC 99	99		
Parabacteroides distasonis		0.26	0.15		1		
41715320	**Good identification**						**41715320**
					GRA		
Eubacterium contortum		97.63	0.48	SUC 99	99		
Bacteroides vulgatus		2.11	0.13		1		
Parabacteroides distasonis		0.26	0.15		1		
41715322	**Low discrimination**						**41715322**
					GRA		
Parabacteroides distasonis		56.98	0.50	MNS 98	1		
Eubacterium contortum		38.96	0.35		99		
Prevotella oralis		3.69	0.16		1		
41715332	**Good identification**						**41715332**
					GRA		
Parabacteroides distasonis		98.73	0.45	MNS 98	1		
Eubacterium contortum		1.27	0.01		99		
41715502	**Acceptable identification**						**41715502**
Bifidobacterium adolescentis		93.53	0.45	SUC 99			
Eubacterium contortum		4.12	0.13	MNS 87			
Actinomyces israelii		1.66	0.21				
41715512	**Good identification**						**41715512**
Bifidobacterium adolescentis		99.34	0.46	SUC 99			
Bifidobacterium longum subsp. longum		0.66	0.07	MNS 87			
41715522	**Low discrimination**						**41715522**
					GRA		
Parabacteroides distasonis		85.89	0.38	MNS 98	1		
Bifidobacterium longum subsp. longum		5.86	0.14		99		
Bifidobacterium adolescentis		3.09	0.11		99		
41715700	**Very good identification**						**41715700**
					GRA		
Eubacterium contortum		99.94	0.60	SUC 99	99		
Bacteroides vulgatus		0.06	-.02		1		
41715702	**Good identification**						**41715702**
					GRA		
Eubacterium contortum		99.58	0.47	SUC 99	99		
Bifidobacterium adolescentis		0.23	0.11		99		
Parabacteroides distasonis		0.19	0.12		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
41715720	**Low discrimination**						**41715720**
					GRA		
Eubacterium contortum		73.91	0.47	SUC 99	99		
Bacteroides vulgatus		25.95	0.32		1		
Parabacteroides distasonis		0.14	0.12		1		
41715722	**Low discrimination**						**41715722**
					GRA		
Parabacteroides distasonis		51.78	0.46	MNS 98	1		
Eubacterium contortum		47.27	0.34		99		
Bacteroides vulgatus		0.95	-.02		1		
41715732	**Good identification**						**41715732**
					GRA		
Parabacteroides distasonis		96.05	0.41	MNS 98	1		
Clostridium clostridioforme		2.31	0.08		50		
Eubacterium contortum		1.64	0.00		99		
41717102	**Low discrimination**						**41717102**
Actinomyces israelii		50.20	0.35	SUC 98			
Bifidobacterium adolescentis		49.80	0.29	CEL 93			
41717502	**Low discrimination**						**41717502**
Bifidobacterium adolescentis		89.03	0.43	SUC 99			
Actinomyces israelii		10.97	0.34	MNS 87			
41717512	**Good identification**						**41717512**
Bifidobacterium adolescentis		98.52	0.45	SUC 99			
Actinomyces israelii		1.48	0.20	MNS 87			
41730000	**Acceptable identification**						**41730000**
					GRA	SPO	
Fusobacterium mortiferum		94.51	0.48	MLT 11	1	1	
Clostridium sporogenes		4.72	0.11		99	99	
Clostridium botulinum biovar A		0.77	-.01		99	99	
41730100	**Good identification**						**41730100**
Fusobacterium mortiferum		100.0	0.55	MLT 11			
41730102	**Good identification**						**41730102**
Fusobacterium mortiferum		100.0	0.38	MLT 11			
41731000	**Good identification**						**41731000**
Fusobacterium mortiferum		100.0	0.51	MLT 11			
41731002	**Good identification**						**41731002**
Fusobacterium mortiferum		100.0	0.35	MLT 11			
41731100	**Very good identification**						**41731100**
					COC	GRA	
Fusobacterium mortiferum		99.38	0.58	MLT 11	1	1	
Blautia hansenii		0.31	-.03		99	99	
Actinomyces naeslundii		0.30	0.26		1	99	
41731102	**Acceptable identification**						**41731102**
					GRA		
Fusobacterium mortiferum		90.75	0.41	MLT 11	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		3.56	0.13	BGL 10	99		
Actinomyces naeslundii		2.49	0.26		99		
41731122	**Low discrimination**						**41731122**
					GRA		
Parabacteroides distasonis		59.19	0.36	MNS 98	1		
Prevotella oralis		27.38	0.18	XYL 90	1		
Clostridium perfringens		6.15	0.10		99		
41731322	**Acceptable identification**						**41731322**
Parabacteroides distasonis		93.80	0.44	MNS 98			
Prevotella oralis		6.20	0.12	XYL 90			
41731332	**Good identification**						**41731332**
Parabacteroides distasonis		100.0	0.40	MNS 98			
41731722	**Good identification**						**41731722**
Parabacteroides distasonis		100.0	0.41	MNS 98			
41731732	**Good identification**						**41731732**
Parabacteroides distasonis		96.67	0.36	MNS 98			
Clostridium clostridioforme		3.33	0.06	XYL 90			
41734100	**Good identification**						**41734100**
Fusobacterium mortiferum		100.0	0.35	MLT 11			
41735100	**Low discrimination**						**41735100**
					COC	GRA	
Fusobacterium mortiferum		80.28	0.38	MLT 11	1	1	
Actinomyces israelii		17.03	0.29	XYL 7	1	99	
Atopobium parvulum		2.69	0.02		99	99	
41735122	**Good identification**						**41735122**
					GRA		
Parabacteroides distasonis		98.94	0.53	MNS 98	1		
Prevotella oralis		0.63	0.02		1		
Bifidobacterium longum subsp. infantis		0.43	-.04		99		
41735132	**Good identification**						**41735132**
Parabacteroides distasonis		100.0	0.48	MNS 98			
41735322	**Very good identification**						**41735322**
					GRA		
Parabacteroides distasonis		99.85	0.61	MNS 98	1		
Eubacterium contortum		0.15	0.01		99		
41735332	**Very good identification**						**41735332**
Parabacteroides distasonis		100.0	0.56	MNS 98			
41735502	**Low discrimination**						**41735502**
					GRA		
Bifidobacterium adolescentis		89.30	0.42	SUC 99	99		
Actinomyces israelii		8.97	0.31	MNS 87	99		
Parabacteroides distasonis		1.73	0.15		1		
41735512	**Good identification**						**41735512**
Bifidobacterium adolescentis		96.69	0.43	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		1.30	0.08	MNS 87		
Actinomyces israelii		1.19	0.17			
41735522	**Good identification**					**41735522**
					GRA	
Parabacteroides distasonis		99.48	0.49	MNS 98	1	
Bifidobacterium adolescentis		0.52	0.07		99	
41735532	**Good identification**					**41735532**
					GRA	
Parabacteroides distasonis		96.19	0.44	MNS 98	1	
Clostridium clostridioforme		1.52	0.08		50	
Bifidobacterium adolescentis		1.15	0.09		99	
41735722	**Very good identification**					**41735722**
					GRA	
Parabacteroides distasonis		99.34	0.57	MNS 98	1	
Clostridium clostridioforme		0.46	0.11		50	
Eubacterium contortum		0.20	0.00		99	
41735732	**Good identification**					**41735732**
Parabacteroides distasonis		96.27	0.53	MNS 98		
Clostridium clostridioforme		3.73	0.23			
41737100	**Good identification**					**41737100**
Actinomyces israelii		100.0	0.42	SUC 98		
41737102	**Low discrimination**					**41737102**
Actinomyces israelii		85.05	0.45	SUC 98		
Bifidobacterium adolescentis		14.95	0.26	CEL 93		
41737140	**Good identification**					**41737140**
Actinomyces israelii		100.0	0.35	SUC 98		
41737142	**Acceptable identification**					**41737142**
Actinomyces israelii		94.71	0.38	SUC 98		
Bifidobacterium adolescentis		5.29	0.10	CEL 93		
41737500	**Good identification**					**41737500**
Actinomyces israelii		100.0	0.41	SUC 98		
41737502	**Low discrimination**					**41737502**
Bifidobacterium adolescentis		58.99	0.40	SUC 99		
Actinomyces israelii		41.02	0.44	MNS 87		
41737512	**Acceptable identification**					**41737512**
Bifidobacterium adolescentis		90.75	0.42	SUC 99		
Actinomyces israelii		7.72	0.30	MNS 87		
Bifidobacterium dentium		1.53	-.07			
41737542	**Low discrimination**					**41737542**
Actinomyces israelii		68.63	0.37	SUC 98		
Bifidobacterium adolescentis		31.37	0.25	CEL 93		
41750000	**Good identification**					**41750000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.49	MLT 11			
41750100	**Good identification**						**41750100**
Fusobacterium mortiferum		100.0	0.56	MLT 11			
41750102	**Good identification**						**41750102**
Fusobacterium mortiferum		100.0	0.39	MLT 11			
41751000	**Good identification**						**41751000**
					GRA		
Fusobacterium mortiferum		99.11	0.52	MLT 11	1		
Lactobacillus jensenii		0.89	0.07		99		
41751002	**Low discrimination**						**41751002**
					GRA		
Fusobacterium mortiferum		83.42	0.36	MLT 11	1		
Prevotella oralis		9.20	0.05	BGL 10	1		
Lactobacillus jensenii		4.50	0.04		99		
41751100	**Very good identification**						**41751100**
					GRA		
Fusobacterium mortiferum		97.46	0.59	MLT 11	1		
Collinsella aerofaciens		0.74	-.03		99		
Actinomyces israelii		0.43	0.21		99		
41751102	**Low discrimination**						**41751102**
					GRA		
Fusobacterium mortiferum		45.81	0.42	MLT 11	1		
Prevotella oralis		33.93	0.26	BGL 10	1		
Bifidobacterium breve		9.41	0.27		99		
41751122	**Low discrimination**						**41751122**
					GRA	SPO	
Prevotella oralis		89.64	0.48	SUC 92	1	1	
Clostridium paraputrificum		6.71	0.22	MNS 99	99	99	
Bifidobacterium breve		2.33	0.32		99	1	
41751322	**Good identification**						**41751322**
					SPO		
Prevotella oralis		96.61	0.41	SUC 92	1		
Parabacteroides distasonis		3.11	0.29	MNS 99	1		
Clostridium ramosum		0.28	-.02		92		
41751502	**Acceptable identification**						**41751502**
Bifidobacterium adolescentis		93.92	0.36	SUC 99			
Actinomyces israelii		6.08	0.22	MNS 87			
41751512	**Good identification**						**41751512**
Bifidobacterium adolescentis		100.0	0.38	SUC 99			
41753102	**Low discrimination**						**41753102**
Actinomyces israelii		50.49	0.36	SUC 98			
Bifidobacterium breve		30.85	0.27	XYL 92			
Bifidobacterium adolescentis		13.77	0.21				
41753512	**Acceptable identification**						**41753512**
Bifidobacterium adolescentis		94.80	0.36	SUC 99			
Actinomyces israelii		5.20	0.21	MNS 87			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41754100	**Good identification**						**41754100**
Fusobacterium mortiferum		100.0	0.36	MLT 11			
41755100	**Low discrimination**						**41755100**
					GRA	SPO	
Fusobacterium mortiferum		58.72	0.39	MLT 11	1	1	
Actinomyces israelii		39.43	0.39	XYL 7	99	1	
Clostridium butyricum		1.86	-.05		99	99	
41755102	**Low discrimination**						**41755102**
					GRA		
Bifidobacterium adolescentis		46.90	0.36	SUC 99	99		
Actinomyces israelii		42.67	0.42	MNS 87	99		
Prevotella oralis		5.73	0.10	ARA 87	1		
41755112	**Low discrimination**						**41755112**
Bifidobacterium adolescentis		89.99	0.38	SUC 99			
Actinomyces israelii		10.01	0.28	MNS 87			
41755322	**Low discrimination**						**41755322**
					GRA		
Parabacteroides distasonis		69.48	0.45	MNS 98	1		
Prevotella oralis		29.63	0.26		1		
Eubacterium contortum		0.89	0.01		99		
41755332	**Good identification**						**41755332**
Parabacteroides distasonis		100.0	0.40	MNS 98			
41755500	**Low discrimination**						**41755500**
					SPO		
Actinomyces israelii		60.46	0.37	SUC 98	1		
Clostridium butyricum		31.30	0.11		99		
Bifidobacterium adolescentis		8.24	0.16		1		
41755502	**Low discrimination**						**41755502**
					SPO		
Bifidobacterium adolescentis		88.80	0.50	SUC 99	1		
Actinomyces israelii		9.88	0.40	MNS 87	1		
Clostridium butyricum		1.33	0.04		99		
41755512	**Good identification**						**41755512**
Bifidobacterium adolescentis		98.66	0.52	SUC 99			
Actinomyces israelii		1.34	0.26	MNS 87			
41755552	**Good identification**						**41755552**
Bifidobacterium adolescentis		95.90	0.36	SUC 99			
Actinomyces israelii		4.10	0.19	MNS 87			
41755722	**Acceptable identification**						**41755722**
Parabacteroides distasonis		90.47	0.41	MNS 98			
Bacteroides uniformis		6.38	0.03				
Clostridium clostridioforme		1.60	0.06				
41755732	**Low discrimination**						**41755732**
Parabacteroides distasonis		86.99	0.37	MNS 98			
Clostridium clostridioforme		13.01	0.17				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
41757000	**Good identification**					**41757000**
Actinomyces israelii		100.0	0.41	SUC 98		
41757002	**Good identification**					**41757002**
Actinomyces israelii		100.0	0.44	SUC 98		
41757042	**Good identification**					**41757042**
Actinomyces israelii		100.0	0.37	SUC 98		
41757100	**Good identification**					**41757100**
Actinomyces israelii		100.0	0.52	SUC 98		
41757102	**Low discrimination**					**41757102**
Actinomyces israelii		86.30	0.55	SUC 98		
Bifidobacterium adolescentis		13.70	0.35			
41757110	**Good identification**					**41757110**
Actinomyces israelii		100.0	0.37	SUC 98		
41757112	**Low discrimination**					**41757112**
Bifidobacterium adolescentis		56.48	0.36	SUC 99		
Actinomyces israelii		43.52	0.40	MNS 87		
41757140	**Good identification**					**41757140**
Actinomyces israelii		100.0	0.45	SUC 98		
41757142	**Good identification**					**41757142**
Actinomyces israelii		95.20	0.48	SUC 98		
Bifidobacterium adolescentis		4.80	0.19			
41757302	**Good identification**					**41757302**
Actinomyces israelii		100.0	0.35	SUC 98		
41757400	**Good identification**					**41757400**
Actinomyces israelii		100.0	0.40	SUC 98		
41757402	**Good identification**					**41757402**
Actinomyces israelii		95.02	0.43	SUC 98		
Bifidobacterium adolescentis		4.98	0.15			
41757442	**Good identification**					**41757442**
Actinomyces israelii		100.0	0.36	SUC 98		
41757500	**Good identification**					**41757500**
					SPO	
Actinomyces israelii		97.10	0.50	SUC 98	1	
Bifidobacterium adolescentis		1.91	0.15		1	
Clostridium butyricum		0.99	-.05		99	
41757502	**Low discrimination**					**41757502**
Bifidobacterium adolescentis		56.48	0.49	SUC 99		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		43.52	0.53	MNS 87		
41757510	**Low discrimination**					**41757510**
Actinomyces israelii		86.14	0.36	SUC 98		
Bifidobacterium adolescentis		13.86	0.16	MLZ 13		
41757512	**Acceptable identification**					**41757512**
Bifidobacterium adolescentis		90.95	0.50	SUC 99		
Actinomyces israelii		8.57	0.39	MNS 87		
Bifidobacterium dentium		0.48	-.07			
41757540	**Good identification**					**41757540**
Actinomyces israelii		100.0	0.43	SUC 98		
41757542	**Low discrimination**					**41757542**
Actinomyces israelii		70.80	0.46	SUC 98		
Bifidobacterium adolescentis		29.21	0.33			
41770000	**Good identification**					**41770000**
Fusobacterium mortiferum		100.0	0.40	MLT 11		
41770100	**Good identification**					**41770100**
Fusobacterium mortiferum		100.0	0.47	MLT 11		
41771000	**Low discrimination**					**41771000**
					GRA	
Fusobacterium mortiferum		89.09	0.43	MLT 11	1	
Lactobacillus jensenii		8.04	0.15		99	
Actinomyces israelii		2.88	0.20		99	
41771100	**Acceptable identification**					**41771100**
					GRA	
Fusobacterium mortiferum		92.03	0.50	MLT 11	1	
Actinomyces israelii		5.09	0.30		99	
Actinomyces naeslundii		1.12	0.28		99	
41771322	**Low discrimination**					**41771322**
					SPO	
Parabacteroides distasonis		57.34	0.40	MNS 98	1	
Prevotella oralis		24.95	0.21	XYL 90	1	
Clostridium ramosum		17.70	0.19		92	
41771332	**Good identification**					**41771332**
Parabacteroides distasonis		100.0	0.35	MNS 98		
41771722	**Good identification**					**41771722**
Parabacteroides distasonis		100.0	0.36	MNS 98		
41773002	**Acceptable identification**					**41773002**
Actinomyces israelii		93.43	0.36	SUC 98		
Actinomyces naeslundii		6.57	0.25	XYL 92		
41773100	**Acceptable identification**					**41773100**
					GRA	
Actinomyces israelii		90.31	0.43	SUC 98	99	
Actinomyces naeslundii		6.78	0.33	XYL 92	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		2.91	0.15		1		
41773101	**Good identification**						**41773101**
Actinomyces naeslundii		100.0	0.36	ESL 12			
41773102	**Low discrimination**						**41773102**
Actinomyces israelii		82.87	0.46	SUC 98			
Bifidobacterium breve		8.98	0.24	XYL 92			
Actinomyces naeslundii		4.15	0.33				
41773103	**Good identification**						**41773103**
Actinomyces naeslundii		100.0	0.36	ESL 12			
41773121	**Good identification**						**41773121**
Actinomyces naeslundii		100.0	0.36	ESL 12			
41773123	**Good identification**						**41773123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
41773140	**Good identification**						**41773140**
Actinomyces israelii		100.0	0.36	SUC 98			
41773142	**Low discrimination**						**41773142**
Actinomyces israelii		78.22	0.39	SUC 98			
Bifidobacterium breve		21.78	0.24	XYL 92			
41773500	**Good identification**						**41773500**
Actinomyces israelii		100.0	0.42	SUC 98			
41773502	**Low discrimination**						**41773502**
Actinomyces israelii		71.67	0.45	SUC 98			
Bifidobacterium adolescentis		28.33	0.32	XYL 92			
41773542	**Low discrimination**						**41773542**
Actinomyces israelii		88.84	0.38	SUC 98			
Bifidobacterium adolescentis		11.16	0.16	XYL 92			
41775000	**Low discrimination**						**41775000**
					GRA		
Actinomyces israelii		83.14	0.38	SUC 98	99		
Fusobacterium mortiferum		16.86	0.24		1		
41775002	**Good identification**						**41775002**
Actinomyces israelii		100.0	0.41	SUC 98			
41775100	**Low discrimination**						**41775100**
					GRA	SPO	
Actinomyces israelii		74.51	0.49	SUC 98	99	1	
Clostridium butyricum		14.61	0.15		99	99	
Fusobacterium mortiferum		8.81	0.30		1	1	
41775102	**Low discrimination**						**41775102**
					SPO		
Actinomyces israelii		79.06	0.52	SUC 98	1		
Bifidobacterium adolescentis		15.40	0.33		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		4.02	0.08		99		
41775122	**Acceptable identification**						**41775122**
					GRA		
Parabacteroides distasonis		90.62	0.48	MNS 98	1		
Actinomyces israelii		5.59	0.31		99		
Prevotella oralis		3.79	0.12		1		
41775132	**Good identification**						**41775132**
					GRA		
Parabacteroides distasonis		98.32	0.43	MNS 98	1		
Actinomyces israelii		1.68	0.17		99		
41775140	**Good identification**						**41775140**
					SPO		
Actinomyces israelii		95.80	0.42	SUC 98	1		
Clostridium butyricum		4.20	-.03		99		
41775142	**Acceptable identification**						**41775142**
Actinomyces israelii		94.17	0.45	SUC 98			
Bifidobacterium adolescentis		5.83	0.18				
41775322	**Very good identification**						**41775322**
Parabacteroides distasonis		99.41	0.56	MNS 98			
Prevotella oralis		0.59	0.05				
41775332	**Good identification**						**41775332**
Parabacteroides distasonis		99.07	0.52	MNS 98			
Clostridium clostridioforme		0.93	0.11				
41775400	**Low discrimination**						**41775400**
					SPO		
Actinomyces israelii		73.57	0.37	SUC 98	1		
Clostridium butyricum		26.44	0.08		99		
41775402	**Low discrimination**						**41775402**
					SPO		
Actinomyces israelii		86.39	0.40	SUC 98	1		
Clostridium butyricum		8.05	0.01		99		
Bifidobacterium adolescentis		5.56	0.13		1		
41775502	**Low discrimination**						**41775502**
					SPO		
Bifidobacterium adolescentis		50.27	0.47	SUC 99	1		
Actinomyces israelii		31.56	0.50	MNS 87	1		
Clostridium butyricum		17.64	0.25		99		
41775512	**Low discrimination**						**41775512**
Bifidobacterium adolescentis		89.33	0.49	SUC 99			
Actinomyces israelii		6.86	0.36	MNS 87			
Clostridium clostridioforme		1.89	0.17				
41775522	**Low discrimination**						**41775522**
					GRA	SPO	
Parabacteroides distasonis		84.92	0.44	MNS 98	1	1	
Actinomyces israelii		7.00	0.30		99	1	
Clostridium butyricum		5.33	0.07		99	99	
41775532	**Low discrimination**						**41775532**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		81.80	0.40	MNS 98	1	
Clostridium clostridioforme		8.06	0.17		50	
Bifidobacterium dentium		4.41	-.03		99	
41775540	**Low discrimination**					**41775540**
					SPO	
Actinomyces israelii		67.47	0.40	SUC 98	1	
Clostridium butyricum		32.53	0.14		99	
41775542	**Low discrimination**					**41775542**
					SPO	
Actinomyces israelii		61.30	0.43	SUC 98	1	
Bifidobacterium adolescentis		31.04	0.32		1	
Clostridium butyricum		7.66	0.07		99	
41775722	**Good identification**					**41775722**
Parabacteroides distasonis		95.74	0.53	MNS 98		
Clostridium clostridioforme		2.73	0.20			
Bacteroides ovatus		1.53	0.06			
41775732	**Low discrimination**					**41775732**
Parabacteroides distasonis		79.32	0.48	MNS 98		
Clostridium clostridioforme		19.10	0.32			
Bacteroides ovatus		1.57	0.03			
41777000	**Good identification**					**41777000**
Actinomyces israelii		100.0	0.51	SUC 98		
41777002	**Good identification**					**41777002**
Actinomyces israelii		100.0	0.54	SUC 98		
41777010	**Good identification**					**41777010**
Actinomyces israelii		100.0	0.37	SUC 98		
41777012	**Good identification**					**41777012**
Actinomyces israelii		100.0	0.40	SUC 98		
41777040	**Good identification**					**41777040**
Actinomyces israelii		100.0	0.44	SUC 98		
41777042	**Good identification**					**41777042**
Actinomyces israelii		100.0	0.47	SUC 98		
41777100	**Very good identification**					**41777100**
					SPO	
Actinomyces israelii		99.62	0.61	SUC 98	1	
Clostridium butyricum		0.38	-.01		99	
41777102	**Very good identification**					**41777102**
Actinomyces israelii		97.27	0.64	SUC 98		
Bifidobacterium adolescentis		2.74	0.32			
41777110	**Good identification**					**41777110**
Actinomyces israelii		100.0	0.47	SUC 98		
41777112	**Low discrimination**					**41777112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		80.50	0.50	SUC 98		
Bifidobacterium adolescentis		18.52	0.33	MLZ 13		
Bifidobacterium dentium		0.98	-.07			
41777120	**Good identification**					**41777120**
Actinomyces israelii		100.0	0.41	SUC 98		
41777122	**Good identification**					**41777122**
					GRA	
Actinomyces israelii		97.19	0.44	SUC 98	99	
Parabacteroides distasonis		2.81	0.14	NAG 6	1	
41777140	**Good identification**					**41777140**
Actinomyces israelii		100.0	0.54	SUC 98		
41777142	**Very good identification**					**41777142**
Actinomyces israelii		99.11	0.57	SUC 98		
Bifidobacterium adolescentis		0.89	0.16			
41777150	**Good identification**					**41777150**
					COC	
Actinomyces israelii		95.56	0.40	SUC 98	1	
Blautia producta		4.44	-.03	MLZ 13	99	
41777152	**Acceptable identification**					**41777152**
Actinomyces israelii		93.19	0.43	SUC 98		
Bifidobacterium adolescentis		6.81	0.18	MLZ 13		
41777162	**Good identification**					**41777162**
Actinomyces israelii		100.0	0.37	SUC 98		
41777300	**Good identification**					**41777300**
Actinomyces israelii		100.0	0.42	SUC 98		
41777302	**Good identification**					**41777302**
Actinomyces israelii		100.0	0.45	SUC 98		
41777340	**Good identification**					**41777340**
Actinomyces israelii		100.0	0.35	SUC 98		
41777342	**Good identification**					**41777342**
Actinomyces israelii		100.0	0.38	SUC 98		
41777400	**Good identification**					**41777400**
Actinomyces israelii		100.0	0.50	SUC 98		
41777402	**Good identification**					**41777402**
Actinomyces israelii		99.08	0.53	SUC 98		
Bifidobacterium adolescentis		0.92	0.12			
41777410	**Good identification**					**41777410**
Actinomyces israelii		100.0	0.36	SUC 98		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
41777412	**Low discrimination**						**41777412**
Actinomyces israelii		88.04	0.39	SUC 98			
Bifidobacterium adolescentis		6.69	0.13	MLZ 13			
Bifidobacterium dentium		5.26	-.07				
41777440	**Good identification**						**41777440**
Actinomyces israelii		100.0	0.43	SUC 98			
41777442	**Good identification**						**41777442**
Actinomyces israelii		100.0	0.46	SUC 98			
41777500	**Very good identification**						**41777500**
					SPO		
Actinomyces israelii		95.62	0.60	SUC 98	1		
Clostridium butyricum		4.04	0.15		99		
Bifidobacterium adolescentis		0.33	0.12		1		
41777502	**Low discrimination**						**41777502**
					SPO		
Actinomyces israelii		80.44	0.63	SUC 98	1		
Bifidobacterium adolescentis		18.50	0.46		1		
Clostridium butyricum		0.88	0.08		99		
41777510	**Low discrimination**						**41777510**
Actinomyces israelii		77.82	0.46	SUC 98			
Bifidobacterium dentium		17.08	0.10	MLZ 13			
Bifidobacterium adolescentis		2.22	0.13				
41777512	**Low discrimination**						**41777512**
Bifidobacterium adolescentis		43.14	0.47	SUC 99			
Bifidobacterium dentium		33.93	0.27	MNS 87			
Actinomyces israelii		22.93	0.49				
41777520	**Acceptable identification**						**41777520**
					SPO		
Actinomyces israelii		94.55	0.40	SUC 98	1		
Clostridium butyricum		5.45	-.03	NAG 6	99		
41777522	**Low discrimination**						**41777522**
Actinomyces israelii		89.27	0.43	SUC 98			
Bifidobacterium dentium		5.55	-.03	NAG 6			
Bifidobacterium adolescentis		3.25	0.12				
41777540	**Low discrimination**						**41777540**
					COC	SPO	
Actinomyces israelii		77.54	0.53	SUC 98	1	1	
Blautia producta		21.73	0.24		99	1	
Clostridium butyricum		0.73	-.03		1	99	
41777542	**Acceptable identification**						**41777542**
					COC		
Actinomyces israelii		91.42	0.56	SUC 98	1		
Bifidobacterium adolescentis		6.68	0.30		1		
Blautia producta		1.90	0.07		99		
41777552	**Low discrimination**						**41777552**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		48.71	0.42	SUC 98	1		
Bifidobacterium adolescentis		29.12	0.32	MLZ 13	1		
Blautia producta		20.30	0.15		99		
41777562	**Good identification**						**41777562**
Actinomyces israelii		100.0	0.35	SUC 98			
41777700	**Good identification**						**41777700**
Actinomyces israelii		100.0	0.41	SUC 98			
41777702	**Good identification**						**41777702**
Actinomyces israelii		97.01	0.44	SUC 98			
Bifidobacterium adolescentis		2.99	0.12	RHA 7			
41777742	**Low discrimination**						**41777742**
					COC		
Actinomyces israelii		78.38	0.37	SUC 98	1		
Blautia producta		21.62	0.07	RHA 7	99		
42000000	**Low discrimination**						**42000000**
					SPO		
Clostridium botulinum biovar A		44.87	0.64	GLU 90	99		
Atopobium minutum		25.30	0.68	SUC 7	1		
Clostridium sporogenes		7.03	0.48		99		
42000002	**Low discrimination**						**42000002**
					GRA	SPO	
Atopobium minutum		76.24	0.65	SUC 5	99	1	
Clostridium sporogenes		10.60	0.40		99	99	
Pseudoflavonifractor capillosus		5.92	0.32		1	1	
42000020	**Low discrimination**						**42000020**
					GRA	SPO	
Atopobium minutum		78.55	0.60	SUC 5	99	1	
Pseudoflavonifractor capillosus		12.19	0.32		1	1	
Clostridium botulinum biovar A		4.22	0.30		99	99	
42000022	**Very good identification**						**42000022**
					COC	GRA	
Pseudoflavonifractor capillosus		95.79	0.66	SUC 1	1	1	
Atopobium minutum		4.16	0.57		1	99	
Peptococcus niger		0.03	0.03		99	99	
42000040	**Good identification**						**42000040**
					SPO		
Clostridium botulinum biovar A		95.86	0.56	GLU 90	99		
Atopobium minutum		1.83	0.34	SUC 7	1		
Clostridium sporogenes		0.51	0.14		99		
42001000	**Low discrimination**						**42001000**
					SPO		
Atopobium minutum		83.55	0.60	SUC 5	1		
Eubacterium tortuosum		8.76	0.32		1		
Clostridium botulinum biovar A		4.49	0.30		99		
42001002	**Good identification**						**42001002**
					GRA		
Atopobium minutum		94.53	0.57	SUC 5	99		
Eubacterium tortuosum		2.62	0.19		99		
Pseudoflavonifractor capillosus		2.45	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
42001020	**Acceptable identification**							**42001020**
					GRA			
Atopobium minutum		90.32	0.52	SUC 5	99			
Eubacterium tortuosum		5.01	0.19		99			
Pseudoflavonifractor capillosus		4.67	0.15		1			
42001022	**Low discrimination**							**42001022**
					GRA			
Pseudoflavonifractor capillosus		88.33	0.50	SUC 1	1			
Atopobium minutum		11.50	0.49	GAL 10	99			
Eubacterium tortuosum		0.17	0.06		99			
42010000	**Low discrimination**							**42010000**
					SPO			
Atopobium minutum		81.57	0.67	SUC 5	1			
Clostridium botulinum biovar A		10.01	0.44		99			
Clostridium sporogenes		4.34	0.35		99			
42010002	**Very good identification**							**42010002**
					GRA	SPO		
Atopobium minutum		95.95	0.64	SUC 5	99	1		
Clostridium sporogenes		2.55	0.27		99	99		
Pseudoflavonifractor capillosus		0.90	0.15		1	1		
42010020	**Very good identification**							**42010020**
					GRA	SPO		
Atopobium minutum		97.27	0.59	SUC 5	99	1		
Pseudoflavonifractor capillosus		1.82	0.15		1	1		
Clostridium botulinum biovar A		0.36	0.10		99	99		
42010022	**Low discrimination**							**42010022**
					GRA			
Pseudoflavonifractor capillosus		73.50	0.50	SUC 1	1			
Atopobium minutum		26.50	0.56	FRU 10	99			
42010040	**Low discrimination**							**42010040**
					COC	SPO	GRA	
Clostridium botulinum biovar A		67.29	0.35	GLU 90	1	99	99	
Atopobium minutum		18.55	0.33	SUC 7	1	1	99	
Megasphaera elsdenii		10.75	0.26	FRU 6	99	1	1	
42010100	**Low discrimination**							**42010100**
					GRA			
Fusobacterium mortiferum		45.36	0.48	GLU 99	1			
Actinomyces meyeri		39.76	0.38		99			
Atopobium minutum		12.66	0.33		99			
42011000	**Low discrimination**							**42011000**
					GRA			
Atopobium minutum		89.70	0.59	SUC 5	99			
Fusobacterium mortiferum		6.26	0.45		1			
Actinomyces meyeri		3.20	0.31		99			
42011002	**Very good identification**							**42011002**
					GRA			
Atopobium minutum		98.21	0.56	SUC 5	99			
Fusobacterium mortiferum		1.14	0.28		1			
Pseudoflavonifractor capillosus		0.31	-.01		1			
42011020	**Good identification**							**42011020**
					GRA			
Atopobium minutum		97.96	0.51	SUC 5	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		1.43	0.16		99		
Pseudoflavonifractor capillosus		0.61	-.01		1		
42011022	**Low discrimination**						**42011022**
					GRA		
Atopobium minutum		51.96	0.48	SUC 5	99		
Pseudoflavonifractor capillosus		48.04	0.33		1		
42011100	**Low discrimination**						**42011100**
					GRA		
Fusobacterium mortiferum		50.66	0.51	GLU 99	1		
Actinomyces meyeri		44.41	0.41		99		
Atopobium minutum		3.14	0.25		99		
42014100	**Good identification**						**42014100**
					GRA		
Actinomyces meyeri		95.76	0.43	ESL 1	99		
Fusobacterium mortiferum		4.24	0.29	GLU 93	1		
42015000	**Low discrimination**						**42015000**
					GRA		
Actinomyces meyeri		80.35	0.35	ESL 1	99		
Atopobium minutum		11.71	0.25	GLU 93	99		
Fusobacterium mortiferum		6.09	0.26		1		
42015100	**Acceptable identification**						**42015100**
					GRA		
Actinomyces meyeri		94.72	0.46	ESL 1	99		
Fusobacterium mortiferum		4.19	0.32	GLU 93	1		
Eubacterium saburreum		0.69	0.17		85		
42015500	**Good identification**						**42015500**
Actinomyces meyeri		98.33	0.35	ESL 1			
Lactobacillus fermentum		1.67	0.16	GLU 93			
42020000	**Low discrimination**						**42020000**
					SPO		
Clostridium botulinum biovar A		79.89	0.54	GLU 90	99		
Clostridium sporogenes		16.70	0.40	SUC 7	99		
Atopobium minutum		1.82	0.34		1		
42020040	**Good identification**						**42020040**
Clostridium botulinum biovar A		99.30	0.45	GLU 90			
Clostridium sporogenes		0.70	0.06	SUC 7			
42030100	**Acceptable identification**						**42030100**
					GRA	SPO	
Fusobacterium mortiferum		94.49	0.39	GLU 99	1	1	
Actinomyces meyeri		2.80	0.04		99	1	
Clostridium botulinum biovar A		2.71	-.01		99	99	
42031000	**Low discrimination**						**42031000**
					GRA		
Fusobacterium mortiferum		63.27	0.36	GLU 99	1		
Atopobium minutum		30.64	0.25		99		
Eubacterium saburreum		3.27	0.12		85		
42031100	**Acceptable identification**						**42031100**
					GRA		
Fusobacterium mortiferum		92.48	0.42	GLU 99	1		
Eubacterium saburreum		4.78	0.18		85		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		2.74	0.07		99		
42050100	**Good identification**						**42050100**
					GRA		
Fusobacterium mortiferum		97.41	0.40	GLU 99	1		
Actinomyces meyeri		2.59	0.04		99		
42051000	**Low discrimination**						**42051000**
					GRA		
Fusobacterium mortiferum		69.74	0.37	GLU 99	1		
Atopobium minutum		30.27	0.25		99		
42051100	**Good identification**						**42051100**
					GRA		
Fusobacterium mortiferum		97.41	0.43	GLU 99	1		
Actinomyces meyeri		2.59	0.07		99		
42100000	**Low discrimination**						**42100000**
					COC	SPO	
Clostridium botulinum biovar A		71.55	0.63	GLU 90	1	99	
Clostridium sporogenes		25.46	0.53	SUC 7	1	99	
Peptostreptococcus anaerobius		1.09	0.27		99	1	
42100002	**Low discrimination**						**42100002**
					SPO		
Clostridium sporogenes		88.60	0.45	SUC 1	99		
Clostridium botulinum biovar A		7.55	0.29		99		
Atopobium minutum		3.47	0.31		1		
42100040	**Good identification**						**42100040**
Clostridium botulinum biovar A		98.38	0.54	GLU 90			
Clostridium sporogenes		1.18	0.19	SUC 7			
Clostridium bifermentans		0.17	0.06				
42101101	**Good identification**						**42101101**
Clostridium chauvoei		100.0	0.40	GLU 99			
42110000	**Low discrimination**						**42110000**
					COC	GRA	
Clostridium botulinum biovar A		42.34	0.43	GLU 90	1	99	
Clostridium sporogenes		41.65	0.40	SUC 7	1	99	
Megasphaera elsdenii		5.56	0.32	FRU 6	99	1	
42110100	**Low discrimination**						**42110100**
					GRA	SPO	
Actinomyces meyeri		84.66	0.39	ESL 1	99	1	
Fusobacterium mortiferum		10.59	0.33	GLU 93	1	1	
Clostridium botulinum biovar A		2.40	0.08		99	99	
42111100	**Low discrimination**						**42111100**
					GRA		
Actinomyces meyeri		73.59	0.42	ESL 1	99		
Fusobacterium mortiferum		9.20	0.36	GLU 93	1		
Eubacterium saburreum		6.89	0.32		85		
42111101	**Good identification**						**42111101**
					SPO		
Clostridium chauvoei		99.61	0.47	GLU 99	99		
Actinomyces meyeri		0.39	0.08	MNS 90	1		
42111120	**Good identification**						**42111120**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		96.50	0.48	GLU 90	1	1	
Actinomyces meyeri		1.84	0.27	MNS 90	99	1	
Capnocytophaga ochracea		0.59	0.14	RAF 90	1	90	
42111121	**Acceptable identification**						**42111121**
					GRA	SPO	
Clostridium chauvoei		92.19	0.38	GLU 99	99	99	
Prevotella melaninogenica		7.81	0.14	MNS 90	1	1	
42111122	**Low discrimination**						**42111122**
					GRA	CAT	
Prevotella melaninogenica		76.01	0.36	GLU 90	1	1	
Capnocytophaga ochracea		21.79	0.30	MNS 90	1	90	
Clostridium perfringens		1.12	0.11	RAF 90	99	1	
42114100	**Good identification**						**42114100**
					GRA		
Actinomyces meyeri		99.52	0.44	ESL 1	99		
Fusobacterium mortiferum		0.48	0.13	GLU 93	1		
42115000	**Acceptable identification**						**42115000**
Actinomyces meyeri		93.27	0.36	ESL 1			
Eubacterium saburreum		4.15	0.21	GLU 93			
Lactobacillus fermentum		2.58	0.20				
42115100	**Good identification**						**42115100**
Actinomyces meyeri		95.96	0.47	ESL 1			
Eubacterium saburreum		2.49	0.27	GLU 93			
Lactobacillus fermentum		1.08	0.24				
42115500	**Acceptable identification**						**42115500**
Actinomyces meyeri		93.73	0.36	ESL 1			
Lactobacillus fermentum		4.23	0.24	GLU 93			
Bifidobacterium longum subsp. longum		2.04	0.12				
42120000	**Low discrimination**						**42120000**
					COC	SPO	
Clostridium botulinum biovar A		67.80	0.53	GLU 90	1	99	
Clostridium sporogenes		32.16	0.45	SUC 7	1	99	
Peptostreptococcus anaerobius		0.04	-.07		99	1	
42120002	**Acceptable identification**						**42120002**
Clostridium sporogenes		94.00	0.37	SUC 1			
Clostridium botulinum biovar A		6.00	0.19				
42120040	**Good identification**						**42120040**
Clostridium botulinum biovar A		98.42	0.44	GLU 90			
Clostridium sporogenes		1.58	0.11	SUC 7			
42201000	**Very good identification**						**42201000**
					GRA		
Eubacterium tortuosum		99.89	0.66	GLU 99	99		
Atopobium minutum		0.10	0.26		99		
Fusobacterium mortiferum		0.01	0.16		1		
42201002	**Good identification**						**42201002**
Eubacterium tortuosum		99.63	0.53	GLU 99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		0.37	0.23				
42201020	**Good identification**						**42201020**
Eubacterium tortuosum		99.82	0.53	GLU 99			
Atopobium minutum		0.18	0.17				
42201022	**Acceptable identification**						**42201022**
					GRA		
Eubacterium tortuosum		94.32	0.40	GLU 99	99		
Pseudoflavonifractor capillosus		5.03	0.15		1		
Atopobium minutum		0.66	0.14		99		
42210000	**Low discrimination**						**42210000**
					GRA	SPO	
Fusobacterium mortiferum		71.07	0.47	GLU 99	1	1	
Atopobium minutum		23.84	0.33		99	1	
Clostridium botulinum biovar A		2.92	0.10		99	99	
42210100	**Good identification**						**42210100**
					GRA		
Fusobacterium mortiferum		99.55	0.53	GLU 99	1		
Actinomyces meyeri		0.45	0.04		99		
42210102	**Good identification**						**42210102**
Fusobacterium mortiferum		100.0	0.37	GLU 99			
42211000	**Low discrimination**						**42211000**
					GRA		
Fusobacterium mortiferum		52.28	0.50	GLU 99	1		
Eubacterium tortuosum		43.82	0.32		99		
Atopobium minutum		3.90	0.25		99		
42211100	**Good identification**						**42211100**
					GRA		
Fusobacterium mortiferum		99.19	0.56	GLU 99	1		
Actinomyces meyeri		0.45	0.07		99		
Eubacterium tortuosum		0.36	-.03		99		
42211102	**Good identification**						**42211102**
Fusobacterium mortiferum		100.0	0.40	GLU 99			
42215100	**Low discrimination**						**42215100**
					GRA		
Fusobacterium mortiferum		89.47	0.37	GLU 99	1		
Actinomyces meyeri		10.53	0.12	XYL 7	99		
42230000	**Good identification**						**42230000**
					GRA	SPO	
Fusobacterium mortiferum		96.67	0.38	GLU 99	1	1	
Clostridium botulinum biovar A		3.33	-.01		99	99	
42230100	**Good identification**						**42230100**
Fusobacterium mortiferum		100.0	0.44	GLU 99			
42231000	**Good identification**						**42231000**
					GRA		
Fusobacterium mortiferum		97.24	0.41	GLU 99	1		
Eubacterium tortuosum		2.76	-.03		99		
42231100	**Good identification**						**42231100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.47	GLU 99		
42241000	**Good identification**					**42241000**
Eubacterium tortuosum		100.0	0.50	GLU 99		
42241002	**Good identification**					**42241002**
Eubacterium tortuosum		100.0	0.37	GLU 99		
42241020	**Good identification**					**42241020**
Eubacterium tortuosum		100.0	0.37	GLU 99		
42250000	**Good identification**					**42250000**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
42250100	**Good identification**					**42250100**
Fusobacterium mortiferum		100.0	0.45	GLU 99		
42251000	**Low discrimination**					**42251000**
					GRA	
Fusobacterium mortiferum		72.88	0.42	GLU 99	1	
Eubacterium tortuosum		20.36	0.15		99	
Lactobacillus catenaformis		5.47	0.12		99	
42251100	**Acceptable identification**					**42251100**
					GRA	
Fusobacterium mortiferum		90.91	0.48	GLU 99	1	
Lactobacillus catenaformis		4.38	0.15		99	
Collinsella aerofaciens		2.90	-.03		99	
42270100	**Good identification**					**42270100**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
42271100	**Good identification**					**42271100**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
42310100	**Good identification**					**42310100**
					GRA	
Fusobacterium mortiferum		96.01	0.37	GLU 99	1	
Actinomyces meyeri		4.00	0.05	MLT 11	99	
42311100	**Low discrimination**					**42311100**
					GRA	
Fusobacterium mortiferum		88.68	0.40	GLU 99	1	
Collinsella aerofaciens		7.63	-.03	MLT 11	99	
Actinomyces meyeri		3.69	0.08		99	
42315300	**Very good identification**					**42315300**
Eubacterium contortum		100.0	0.66	GLU 99		
42315302	**Good identification**					**42315302**
Eubacterium contortum		100.0	0.53	GLU 99		
42315320	**Good identification**					**42315320**
Eubacterium contortum		100.0	0.53	GLU 99		
42315322	**Good identification**					**42315322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		100.0	0.40	GLU 99		
42315700	**Very good identification**					**42315700**
Eubacterium contortum		100.0	0.64	GLU 99		
42315702	**Good identification**					**42315702**
Eubacterium contortum		100.0	0.52	GLU 99		
42315720	**Good identification**					**42315720**
					GRA	
Eubacterium contortum		99.86	0.52	GLU 99	99	
Bacteroides vulgatus		0.14	-.05		1	
42315722	**Good identification**					**42315722**
Eubacterium contortum		100.0	0.39	GLU 99		
42357100	**Good identification**					**42357100**
Actinomyces israelii		100.0	0.37	GLU 99		
42357102	**Low discrimination**					**42357102**
					SPO	
Actinomyces israelii		86.57	0.40	GLU 99	1	
Bifidobacterium adolescentis		9.08	0.16		1	
Clostridium tertium		4.35	0.03		99	
42357500	**Good identification**					**42357500**
Actinomyces israelii		100.0	0.35	GLU 99		
42357502	**Low discrimination**					**42357502**
Actinomyces israelii		53.83	0.38	GLU 99		
Bifidobacterium adolescentis		46.18	0.31			
42375102	**Low discrimination**					**42375102**
					COC	
Actinomyces israelii		57.99	0.36	GLU 99	1	
Atopobium parvulum		34.55	0.19		99	
Bifidobacterium adolescentis		7.47	0.15		1	
42377000	**Good identification**					**42377000**
Actinomyces israelii		100.0	0.36	GLU 99		
42377002	**Good identification**					**42377002**
Actinomyces israelii		100.0	0.39	GLU 99		
42377100	**Acceptable identification**					**42377100**
					SPO	
Actinomyces israelii		94.33	0.46	GLU 99	1	
Clostridium tertium		5.68	0.11		99	
42377102	**Low discrimination**					**42377102**
					SPO	
Actinomyces israelii		87.80	0.49	GLU 99	1	
Clostridium tertium		10.57	0.19		99	
Bifidobacterium adolescentis		1.63	0.13		1	
42377140	**Good identification**					**42377140**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.39	GLU 99			
42377142	**Good identification**						**42377142**
Actinomyces israelii		100.0	0.42	GLU 99			
42377402	**Good identification**						**42377402**
Actinomyces israelii		100.0	0.38	GLU 99			
42377500	**Good identification**						**42377500**
					SPO		
Actinomyces israelii		98.34	0.45	GLU 99	1		
Clostridium butyricum		1.66	-.07		99		
42377502	**Low discrimination**						**42377502**
Actinomyces israelii		86.80	0.48	GLU 99			
Bifidobacterium adolescentis		13.20	0.28				
42377540	**Good identification**						**42377540**
Actinomyces israelii		100.0	0.38	GLU 99			
42377542	**Good identification**						**42377542**
Actinomyces israelii		95.39	0.41	GLU 99			
Bifidobacterium adolescentis		4.61	0.12				
42410000	**Low discrimination**						**42410000**
					GRA	SPO	
Fusobacterium mortiferum		71.72	0.47	GLU 99	1	1	
Atopobium minutum		24.05	0.33		99	1	
Clostridium botulinum biovar A		2.95	0.10		99	99	
42410100	**Good identification**						**42410100**
					GRA		
Fusobacterium mortiferum		99.55	0.53	GLU 99	1		
Actinomyces meyeri		0.45	0.04		99		
42410102	**Good identification**						**42410102**
Fusobacterium mortiferum		100.0	0.37	GLU 99			
42411000	**Acceptable identification**						**42411000**
					GRA		
Fusobacterium mortiferum		91.27	0.50	GLU 99	1		
Atopobium minutum		6.80	0.25		99		
Lactobacillus fermentum		1.35	0.20		99		
42411100	**Good identification**						**42411100**
					GRA		
Fusobacterium mortiferum		97.59	0.56	GLU 99	1		
Lactobacillus fermentum		1.01	0.24		99		
Eubacterium saburreum		0.63	0.17		85		
42411102	**Acceptable identification**						**42411102**
					GRA		
Fusobacterium mortiferum		91.50	0.40	GLU 99	1		
Lactobacillus fermentum		8.50	0.24	BGL 10	99		
42415100	**Low discrimination**						**42415100**
					GRA		
Fusobacterium mortiferum		80.38	0.37	GLU 99	1		
Actinomyces meyeri		9.46	0.12	XYL 7	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		6.48	0.20		99			
42430000	**Good identification**							**42430000**
					GRA	SPO		
Fusobacterium mortiferum		96.67	0.38	GLU 99	1	1		
Clostridium botulinum biovar A		3.33	-.01		99	99		
42430100	**Good identification**							**42430100**
Fusobacterium mortiferum		100.0	0.44	GLU 99				
42431000	**Good identification**							**42431000**
Fusobacterium mortiferum		100.0	0.41	GLU 99				
42431100	**Good identification**							**42431100**
Fusobacterium mortiferum		98.59	0.47	GLU 99				
Eubacterium saburreum		1.41	0.14					
42450000	**Good identification**							**42450000**
Fusobacterium mortiferum		100.0	0.39	GLU 99				
42450100	**Good identification**							**42450100**
Fusobacterium mortiferum		100.0	0.45	GLU 99				
42451000	**Good identification**							**42451000**
Fusobacterium mortiferum		100.0	0.42	GLU 99				
42451100	**Good identification**							**42451100**
Fusobacterium mortiferum		100.0	0.48	GLU 99				
42470100	**Good identification**							**42470100**
Fusobacterium mortiferum		100.0	0.36	GLU 99				
42471100	**Good identification**							**42471100**
Fusobacterium mortiferum		100.0	0.39	GLU 99				
42501120	**Good identification**							**42501120**
					COC	GRA		
Prevotella melaninogenica		98.82	0.48	GLU 90	1	1		
Blautia hansenii		1.18	0.15	MNS 90	99	99		
42501122	**Good identification**							**42501122**
					COC	CAT	GRA	
Prevotella melaninogenica		96.80	0.36	GLU 90	1	1	1	
Capnocytophaga ochracea		2.52	0.12	MNS 90	1	90	1	
Blautia hansenii		0.68	-.01	FRU 90	99	1	99	
42505530	**Good identification**							**42505530**
Bifidobacterium longum subsp. longum		100.0	0.35	GLU 99				
42505532	**Good identification**							**42505532**
Bifidobacterium longum subsp. longum		100.0	0.35	GLU 99				
42510100	**Acceptable identification**							**42510100**
					GRA			
Fusobacterium mortiferum		92.40	0.37	GLU 99	1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		3.85	0.05	MLT 11	99		
Prevotella melaninogenica		3.75	-.04		1		
42511120	**Very good identification**						**42511120**
					GRA	CAT	
Prevotella melaninogenica		99.33	0.64	GLU 90	1	1	
Capnocytophaga ochracea		0.60	0.30	MNS 90	1	90	
Lactobacillus fermentum		0.03	0.24		99	1	
42511122	**Low discrimination**						**42511122**
					CAT		
Prevotella melaninogenica		77.09	0.52	GLU 90	1		
Capnocytophaga ochracea		22.10	0.47	MNS 90	90		
Prevotella oralis		0.50	0.18		1		
42511530	**Good identification**						**42511530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.20	0.44	GLU 99	99	1	
Capnocytophaga ochracea		0.97	0.01	MNS 87	1	90	
Prevotella melaninogenica		0.83	-.04		1	1	
42511532	**Acceptable identification**						**42511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.83	0.44	GLU 99	99	1	
Capnocytophaga ochracea		8.17	0.17	MNS 87	1	90	
42515500	**Low discrimination**						**42515500**
Bifidobacterium longum subsp. longum		89.37	0.37	GLU 99			
Lactobacillus fermentum		9.35	0.28	MNS 87			
Actinomyces meyeri		1.28	0.02				
42515502	**Low discrimination**						**42515502**
Bifidobacterium longum subsp. longum		65.48	0.37	GLU 99			
Bifidobacterium adolescentis		24.36	0.32	MNS 87			
Lactobacillus fermentum		6.85	0.28				
42515510	**Good identification**						**42515510**
Bifidobacterium longum subsp. longum		100.0	0.41	GLU 99			
42515512	**Low discrimination**						**42515512**
Bifidobacterium longum subsp. longum		78.92	0.41	GLU 99			
Bifidobacterium adolescentis		21.08	0.33	MNS 87			
42515520	**Low discrimination**						**42515520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.84	0.48	GLU 99	99	1	
Bacteroides vulgatus		7.81	0.15	MNS 87	1	3	
Capnocytophaga ochracea		1.13	0.07		1	90	
42515522	**Low discrimination**						**42515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.71	0.48	GLU 99	99	1	
Capnocytophaga ochracea		9.97	0.23	MNS 87	1	90	
Bacteroides fragilis		1.08	-.03		1	99	
42515530	**Good identification**						**42515530**
Bifidobacterium longum subsp. longum		100.0	0.52	GLU 99			
42515532	**Good identification**						**42515532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.26	0.52	GLU 99	99	1	
Capnocytophaga ochracea		0.74	0.07	MNS 87	1	90	
42535322	**Good identification**						**42535322**
Parabacteroides distasonis		100.0	0.36	GLU 99			
42535520	**Good identification**						**42535520**
					GRA		
Bifidobacterium longum subsp. longum		98.57	0.37	GLU 99	99		
Bacteroides vulgatus		1.43	-.08	MNS 87	1		
42535522	**Acceptable identification**						**42535522**
					GRA		
Bifidobacterium longum subsp. longum		90.86	0.37	GLU 99	99		
Parabacteroides distasonis		9.14	0.25	MNS 87	1		
42535530	**Good identification**						**42535530**
Bifidobacterium longum subsp. longum		100.0	0.41	GLU 99			
42535532	**Acceptable identification**						**42535532**
					GRA		
Bifidobacterium longum subsp. longum		94.63	0.41	GLU 99	99		
Parabacteroides distasonis		3.01	0.20	MNS 87	1		
Clostridium clostridioforme		2.36	0.12		50		
42551120	**Good identification**						**42551120**
					CAT		
Prevotella melaninogenica		98.45	0.48	GLU 90	1		
Capnocytophaga ochracea		1.34	0.20	MNS 90	90		
Prevotella oralis		0.20	0.06	CEL 10	1		
42551122	**Low discrimination**						**42551122**
					CAT		
Prevotella melaninogenica		53.88	0.36	GLU 90	1		
Capnocytophaga ochracea		34.76	0.36	MNS 90	90		
Prevotella oralis		11.08	0.27	CEL 10	1		
42555502	**Good identification**						**42555502**
Bifidobacterium adolescentis		97.36	0.37	GLU 99			
Actinomyces israelii		2.64	0.17	MNS 87			
42555512	**Good identification**						**42555512**
Bifidobacterium adolescentis		96.36	0.39	GLU 99			
Clostridium clostridioforme		1.93	0.07	MNS 87			
Bifidobacterium longum subsp. longum		1.72	0.07				
42557502	**Low discrimination**						**42557502**
Bifidobacterium adolescentis		84.18	0.36	GLU 99			
Actinomyces israelii		15.82	0.30	MNS 87			
42557512	**Good identification**						**42557512**
Bifidobacterium adolescentis		97.75	0.37	GLU 99			
Actinomyces israelii		2.25	0.16	MNS 87			
42575512	**Low discrimination**						**42575512**
Bifidobacterium adolescentis		81.96	0.36	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		18.04	0.21	MNS 87		
42575712	**Good identification**					**42575712**
Clostridium clostridioforme		100.0	0.36	GLU 88		
42575732	**Acceptable identification**					**42575732**
Clostridium clostridioforme		90.05	0.36	GLU 88		
Parabacteroides distasonis		7.58	0.24	MNS 99		
Bacteroides thetaiotaomicron		2.37	-.01	SAL 88		
42577100	**Good identification**					**42577100**
Actinomyces israelii		100.0	0.38	GLU 99		
42577102	**Low discrimination**					**42577102**
Actinomyces israelii		89.66	0.41	GLU 99		
Bifidobacterium adolescentis		10.34	0.19	SAL 92		
42577500	**Good identification**					**42577500**
Actinomyces israelii		100.0	0.37	GLU 99		
42577502	**Low discrimination**					**42577502**
Actinomyces israelii		51.46	0.40	GLU 99		
Bifidobacterium adolescentis		48.54	0.33	SAL 92		
42601000	**Very good identification**					**42601000**
					GRA	
Eubacterium tortuosum		99.93	0.58	GLU 99	99	
Fusobacterium mortiferum		0.07	0.20		1	
42601002	**Good identification**					**42601002**
Eubacterium tortuosum		100.0	0.45	GLU 99		
42601020	**Good identification**					**42601020**
Eubacterium tortuosum		100.0	0.45	GLU 99		
42610000	**Good identification**					**42610000**
Fusobacterium mortiferum		100.0	0.52	GLU 99		
42610002	**Good identification**					**42610002**
Fusobacterium mortiferum		100.0	0.35	GLU 99		
42610100	**Very good identification**					**42610100**
Fusobacterium mortiferum		100.0	0.58	GLU 99		
42610102	**Good identification**					**42610102**
Fusobacterium mortiferum		100.0	0.42	GLU 99		
42611000	**Low discrimination**					**42611000**
					GRA	
Fusobacterium mortiferum		87.42	0.55	GLU 99	1	
Eubacterium tortuosum		12.58	0.24		99	
42611002	**Low discrimination**					**42611002**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		81.40	0.38	GLU 99	1	
Eubacterium tortuosum		18.61	0.11	BGL 10	99	
42611100	**Very good identification**					**42611100**
Fusobacterium mortiferum		100.0	0.61	GLU 99		
42611102	**Good identification**					**42611102**
					GRA	
Fusobacterium mortiferum		98.95	0.45	GLU 99	1	
Bifidobacterium longum subsp. infantis		1.05	-.07	BGL 10	99	
42614100	**Good identification**					**42614100**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
42615000	**Good identification**					**42615000**
Fusobacterium mortiferum		100.0	0.35	GLU 99		
42615100	**Good identification**					**42615100**
Fusobacterium mortiferum		100.0	0.42	GLU 99		
42630000	**Good identification**					**42630000**
Fusobacterium mortiferum		100.0	0.43	GLU 99		
42630100	**Good identification**					**42630100**
Fusobacterium mortiferum		100.0	0.49	GLU 99		
42631000	**Good identification**					**42631000**
Fusobacterium mortiferum		100.0	0.46	GLU 99		
42631100	**Good identification**					**42631100**
Fusobacterium mortiferum		100.0	0.52	GLU 99		
42631102	**Good identification**					**42631102**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
42641000	**Good identification**					**42641000**
					GRA	
Eubacterium tortuosum		99.79	0.41	GLU 99	99	
Fusobacterium mortiferum		0.21	0.12	CEL 10	1	
42650000	**Good identification**					**42650000**
Fusobacterium mortiferum		100.0	0.43	GLU 99		
42650100	**Good identification**					**42650100**
Fusobacterium mortiferum		100.0	0.50	GLU 99		
42651000	**Good identification**					**42651000**
					GRA	
Fusobacterium mortiferum		95.42	0.46	GLU 99	1	
Eubacterium tortuosum		4.58	0.07		99	
42651100	**Good identification**					**42651100**
Fusobacterium mortiferum		100.0	0.53	GLU 99		
42651102	**Good identification**					**42651102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.36	GLU 99		
42670100	**Good identification**					**42670100**
Fusobacterium mortiferum		100.0	0.41	GLU 99		
42671000	**Good identification**					**42671000**
Fusobacterium mortiferum		100.0	0.38	GLU 99		
42671100	**Good identification**					**42671100**
Fusobacterium mortiferum		100.0	0.44	GLU 99		
42710000	**Good identification**					**42710000**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
42710100	**Good identification**					**42710100**
Fusobacterium mortiferum		100.0	0.42	GLU 99		
42711000	**Good identification**					**42711000**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
42711100	**Acceptable identification**					**42711100**
					GRA	
Fusobacterium mortiferum		90.29	0.45	GLU 99	1	
Bifidobacterium longum subsp. infantis		8.45	0.10	MLT 11	99	
Prevotella melaninogenica		1.26	-.04		1	
42715300	**Very good identification**					**42715300**
Eubacterium contortum		100.0	0.61	GLU 99		
42715302	**Good identification**					**42715302**
Eubacterium contortum		100.0	0.48	GLU 99		
42715320	**Good identification**					**42715320**
					GRA	
Eubacterium contortum		99.48	0.48	GLU 99	99	
Bacteroides vulgatus		0.52	0.02		1	
42715322	**Acceptable identification**					**42715322**
					GRA	
Eubacterium contortum		92.61	0.35	GLU 99	99	
Parabacteroides distasonis		5.47	0.26		1	
Prevotella oralis		1.02	0.00		1	
42715502	**Acceptable identification**					**42715502**
Bifidobacterium adolescentis		94.98	0.45	GLU 99		
Eubacterium contortum		4.18	0.13	MNS 87		
Actinomyces israelii		0.84	0.16			
42715512	**Good identification**					**42715512**
Bifidobacterium adolescentis		97.74	0.46	GLU 99		
Clostridium clostridioforme		1.61	0.12	MNS 87		
Bifidobacterium longum subsp. longum		0.65	0.07			
42715700	**Very good identification**					**42715700**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		100.0	0.60	GLU 99		
42715702	**Good identification**					**42715702**
Eubacterium contortum		99.04	0.47	GLU 99		
Clostridium clostridioforme		0.74	0.16			
Bifidobacterium adolescentis		0.23	0.11			
42715720	**Acceptable identification**					**42715720**
					GRA	
Eubacterium contortum		92.16	0.47	GLU 99	99	
Bacteroides vulgatus		7.84	0.21		1	
42717502	**Acceptable identification**					**42717502**
Bifidobacterium adolescentis		94.25	0.43	GLU 99		
Actinomyces israelii		5.75	0.29	MNS 87		
42717512	**Good identification**					**42717512**
Bifidobacterium adolescentis		99.26	0.45	GLU 99		
Actinomyces israelii		0.74	0.15	MNS 87		
42731100	**Acceptable identification**					**42731100**
					GRA	
Fusobacterium mortiferum		94.94	0.36	GLU 99	1	
Actinomyces naeslundii		5.06	0.26	MLT 11	99	
42735322	**Acceptable identification**					**42735322**
					GRA	
Parabacteroides distasonis		91.13	0.37	GLU 99	1	
Clostridium clostridioforme		5.45	0.10	MNS 98	50	
Eubacterium contortum		3.42	0.01		99	
42735502	**Acceptable identification**					**42735502**
Bifidobacterium adolescentis		91.06	0.42	GLU 99		
Actinomyces israelii		4.53	0.26	MNS 87		
Clostridium clostridioforme		4.42	0.16			
42735512	**Low discrimination**					**42735512**
Bifidobacterium adolescentis		84.69	0.43	GLU 99		
Clostridium clostridioforme		15.32	0.27	MNS 87		
42735712	**Good identification**					**42735712**
Clostridium clostridioforme		99.24	0.42	GLU 88		
Bifidobacterium adolescentis		0.76	0.09	MNS 99		
42735732	**Acceptable identification**					**42735732**
Clostridium clostridioforme		92.83	0.42	GLU 88		
Parabacteroides distasonis		7.17	0.29	MNS 99		
42737100	**Good identification**					**42737100**
Actinomyces israelii		100.0	0.37	GLU 99		
42737102	**Low discrimination**					**42737102**
Actinomyces israelii		73.79	0.40	GLU 99		
Bifidobacterium adolescentis		26.21	0.26	CEL 93		
42737500	**Good identification**					**42737500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.35	GLU 99		
42737502	**Low discrimination**					**42737502**
Bifidobacterium adolescentis		74.40	0.40	GLU 99		
Actinomyces israelii		25.60	0.39	MNS 87		
42737512	**Acceptable identification**					**42737512**
Bifidobacterium adolescentis		94.43	0.42	GLU 99		
Actinomyces israelii		3.97	0.24	MNS 87		
Bifidobacterium dentium		1.59	-.07			
42751100	**Low discrimination**					**42751100**
					GRA	
Fusobacterium mortiferum		80.45	0.37	GLU 99	1	
Collinsella aerofaciens		10.70	-.03	MLT 11	99	
Bifidobacterium breve		3.17	0.10		99	
42751502	**Good identification**					**42751502**
Bifidobacterium adolescentis		96.90	0.36	GLU 99		
Actinomyces israelii		3.10	0.17	MNS 87		
42751512	**Good identification**					**42751512**
Bifidobacterium adolescentis		98.31	0.38	GLU 99		
Clostridium clostridioforme		1.69	0.04	MNS 87		
42753512	**Good identification**					**42753512**
Bifidobacterium adolescentis		97.36	0.36	GLU 99		
Actinomyces israelii		2.64	0.16	MNS 87		
42755102	**Low discrimination**					**42755102**
Bifidobacterium adolescentis		68.95	0.36	GLU 99		
Actinomyces israelii		31.05	0.36	MNS 87		
42755112	**Acceptable identification**					**42755112**
Bifidobacterium adolescentis		94.78	0.38	GLU 99		
Actinomyces israelii		5.22	0.22	MNS 87		
42755502	**Acceptable identification**					**42755502**
					SPO	
Bifidobacterium adolescentis		92.86	0.50	GLU 99	1	
Actinomyces israelii		5.11	0.35	MNS 87	1	
Clostridium butyricum		1.39	0.04		99	
42755512	**Good identification**					**42755512**
Bifidobacterium adolescentis		96.84	0.52	GLU 99		
Clostridium clostridioforme		2.51	0.21	MNS 87		
Actinomyces israelii		0.65	0.21			
42755552	**Good identification**					**42755552**
Bifidobacterium adolescentis		97.93	0.36	GLU 99		
Actinomyces israelii		2.07	0.14	MNS 87		
42755712	**Acceptable identification**					**42755712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		94.95	0.36	GLU 88		
Bifidobacterium adolescentis		5.05	0.18	MNS 99		
42755732	**Good identification**					**42755732**
Clostridium clostridioforme		98.04	0.36	GLU 88		
Parabacteroides distasonis		1.96	0.13	MNS 99		
42757000	**Good identification**					**42757000**
Actinomyces israelii		100.0	0.36	GLU 99		
42757002	**Good identification**					**42757002**
Actinomyces israelii		100.0	0.39	GLU 99		
42757100	**Good identification**					**42757100**
Actinomyces israelii		100.0	0.46	GLU 99		
42757102	**Low discrimination**					**42757102**
Actinomyces israelii		75.72	0.49	GLU 99		
Bifidobacterium adolescentis		24.28	0.35			
42757112	**Low discrimination**					**42757112**
Bifidobacterium adolescentis		72.40	0.36	GLU 99		
Actinomyces israelii		27.61	0.35	MNS 87		
42757140	**Good identification**					**42757140**
Actinomyces israelii		100.0	0.39	GLU 99		
42757142	**Acceptable identification**					**42757142**
Actinomyces israelii		90.75	0.42	GLU 99		
Bifidobacterium adolescentis		9.25	0.19			
42757402	**Acceptable identification**					**42757402**
Actinomyces israelii		90.42	0.38	GLU 99		
Bifidobacterium adolescentis		9.58	0.15			
42757500	**Acceptable identification**					**42757500**
					SPO	
Actinomyces israelii		94.32	0.45	GLU 99	1	
Bifidobacterium adolescentis		3.75	0.15		1	
Clostridium butyricum		1.93	-.05		99	
42757502	**Low discrimination**					**42757502**
Bifidobacterium adolescentis		72.40	0.49	GLU 99		
Actinomyces israelii		27.61	0.48	MNS 87		
42757512	**Good identification**					**42757512**
Bifidobacterium adolescentis		95.07	0.50	GLU 99		
Actinomyces israelii		4.43	0.34	MNS 87		
Bifidobacterium dentium		0.50	-.07			
42757540	**Good identification**					**42757540**
Actinomyces israelii		100.0	0.38	GLU 99		
42757542	**Low discrimination**					**42757542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		54.54	0.41	GLU 99		
Bifidobacterium adolescentis		45.46	0.33			
42773100	**Low discrimination**					**42773100**
Actinomyces israelii		86.83	0.38	GLU 99		
Actinomyces naeslundii		13.17	0.33	XYL 92		
42773101	**Good identification**					**42773101**
Actinomyces naeslundii		100.0	0.36	ESL 12		
42773102	**Low discrimination**					**42773102**
Actinomyces israelii		70.54	0.41	GLU 99		
Bifidobacterium breve		15.44	0.24	XYL 92		
Actinomyces naeslundii		7.14	0.33			
42773103	**Good identification**					**42773103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
42773121	**Good identification**					**42773121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
42773123	**Good identification**					**42773123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
42773500	**Good identification**					**42773500**
Actinomyces israelii		100.0	0.37	GLU 99		
42773502	**Low discrimination**					**42773502**
Actinomyces israelii		55.60	0.40	GLU 99		
Bifidobacterium adolescentis		44.40	0.32	XYL 92		
42774712	**Good identification**					**42774712**
Clostridium clostridioforme		100.0	0.43	GLU 88		
42774732	**Good identification**					**42774732**
Clostridium clostridioforme		100.0	0.43	GLU 88		
42775002	**Good identification**					**42775002**
Actinomyces israelii		100.0	0.36	GLU 99		
42775100	**Low discrimination**					**42775100**
					SPO	
Actinomyces israelii		71.63	0.43	GLU 99	1	
Clostridium butyricum		28.37	0.15		99	
42775102	**Low discrimination**					**42775102**
					SPO	
Actinomyces israelii		66.27	0.46	GLU 99	1	
Bifidobacterium adolescentis		26.08	0.33		1	
Clostridium butyricum		6.80	0.08		99	
42775140	**Acceptable identification**					**42775140**
					SPO	
Actinomyces israelii		91.86	0.36	GLU 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		8.14	-.03		99	
42775142	**Low discrimination**					**42775142**
Actinomyces israelii		88.88	0.39	GLU 99		
Bifidobacterium adolescentis		11.12	0.18			
42775502	**Low discrimination**					**42775502**
					SPO	
Bifidobacterium adolescentis		57.54	0.47	GLU 99	1	
Clostridium butyricum		20.19	0.25	MNS 87	99	
Actinomyces israelii		17.88	0.45		1	
42775512	**Low discrimination**					**42775512**
Bifidobacterium adolescentis		74.38	0.49	GLU 99		
Clostridium clostridioforme		21.19	0.36	MNS 87		
Actinomyces israelii		2.83	0.31			
42775532	**Acceptable identification**					**42775532**
Clostridium clostridioforme		90.38	0.36	GLU 88		
Bifidobacterium dentium		3.67	-.03	MNS 99		
Bifidobacterium adolescentis		3.21	0.15	RHA 88		
42775612	**Good identification**					**42775612**
Clostridium clostridioforme		100.0	0.42	GLU 88		
42775632	**Good identification**					**42775632**
Clostridium clostridioforme		100.0	0.42	GLU 88		
42775702	**Acceptable identification**					**42775702**
					SPO	
Clostridium clostridioforme		92.67	0.40	GLU 88	66	
Actinomyces israelii		3.87	0.26	MNS 99	1	
Clostridium butyricum		1.79	-.01		99	
42775712	**Good identification**					**42775712**
Clostridium clostridioforme		99.52	0.51	GLU 88		
Bifidobacterium adolescentis		0.48	0.15	MNS 99		
42775722	**Low discrimination**					**42775722**
Clostridium clostridioforme		87.20	0.40	GLU 88		
Parabacteroides distasonis		9.17	0.29	MNS 99		
Bacteroides ovatus		3.63	0.06			
42775732	**Good identification**					**42775732**
Clostridium clostridioforme		98.18	0.51	GLU 88		
Parabacteroides distasonis		1.22	0.24	MNS 99		
Bacteroides ovatus		0.60	0.03			
42777000	**Good identification**					**42777000**
Actinomyces israelii		100.0	0.46	GLU 99		
42777002	**Good identification**					**42777002**
Actinomyces israelii		100.0	0.49	GLU 99		
42777040	**Good identification**					**42777040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.39	GLU 99		
42777042	**Good identification**					**42777042**
Actinomyces israelii		100.0	0.42	GLU 99		
42777100	**Very good identification**					**42777100**
					SPO	
Actinomyces israelii		99.23	0.56	GLU 99	1	
Clostridium butyricum		0.77	-.01		99	
42777102	**Good identification**					**42777102**
Actinomyces israelii		94.62	0.59	GLU 99		
Bifidobacterium adolescentis		5.38	0.32			
42777110	**Good identification**					**42777110**
Actinomyces israelii		100.0	0.42	GLU 99		
42777112	**Low discrimination**					**42777112**
Actinomyces israelii		67.14	0.45	GLU 99		
Bifidobacterium adolescentis		31.20	0.33	MLZ 13		
Bifidobacterium dentium		1.66	-.07			
42777120	**Good identification**					**42777120**
Actinomyces israelii		100.0	0.36	GLU 99		
42777122	**Good identification**					**42777122**
Actinomyces israelii		100.0	0.39	GLU 99		
42777140	**Good identification**					**42777140**
Actinomyces israelii		100.0	0.49	GLU 99		
42777142	**Good identification**					**42777142**
Actinomyces israelii		98.23	0.52	GLU 99		
Bifidobacterium adolescentis		1.77	0.16			
42777150	**Acceptable identification**					**42777150**
					COC	
Actinomyces israelii		91.41	0.35	GLU 99	1	
Blautia producta		8.59	-.03	MLZ 13	99	
42777152	**Low discrimination**					**42777152**
Actinomyces israelii		87.13	0.38	GLU 99		
Bifidobacterium adolescentis		12.87	0.18	MLZ 13		
42777300	**Good identification**					**42777300**
Actinomyces israelii		100.0	0.37	GLU 99		
42777302	**Good identification**					**42777302**
Actinomyces israelii		100.0	0.40	GLU 99		
42777400	**Good identification**					**42777400**
Actinomyces israelii		100.0	0.44	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
42777402	**Good identification**						**42777402**
Actinomyces israelii		98.16	0.47	GLU 99			
Bifidobacterium adolescentis		1.84	0.12				
42777440	**Good identification**						**42777440**
Actinomyces israelii		100.0	0.37	GLU 99			
42777442	**Good identification**						**42777442**
Actinomyces israelii		100.0	0.40	GLU 99			
42777500	**Acceptable identification**						**42777500**
					SPO		
Actinomyces israelii		91.54	0.55	GLU 99	1		
Clostridium butyricum		7.82	0.15		99		
Bifidobacterium adolescentis		0.65	0.12		1		
42777502	**Low discrimination**						**42777502**
					SPO		
Actinomyces israelii		67.05	0.58	GLU 99	1		
Bifidobacterium adolescentis		31.16	0.46		1		
Clostridium butyricum		1.49	0.08		99		
42777510	**Low discrimination**						**42777510**
Actinomyces israelii		63.46	0.41	GLU 99			
Bifidobacterium dentium		28.15	0.10	MLZ 13			
Bifidobacterium adolescentis		3.66	0.13				
42777512	**Low discrimination**						**42777512**
Bifidobacterium adolescentis		48.79	0.47	GLU 99			
Bifidobacterium dentium		38.37	0.27	MNS 87			
Actinomyces israelii		12.84	0.44				
42777522	**Low discrimination**						**42777522**
Actinomyces israelii		83.39	0.37	GLU 99			
Bifidobacterium dentium		10.48	-.03	NAG 6			
Bifidobacterium adolescentis		6.13	0.12				
42777540	**Low discrimination**						**42777540**
					COC	SPO	
Actinomyces israelii		63.08	0.48	GLU 99	1	1	
Blautia producta		35.72	0.24		99	1	
Clostridium butyricum		1.21	-.03		1	99	
42777542	**Low discrimination**						**42777542**
					COC		
Actinomyces israelii		84.06	0.51	GLU 99	1		
Bifidobacterium adolescentis		12.42	0.30		1		
Blautia producta		3.53	0.07		99		
42777700	**Good identification**						**42777700**
Actinomyces israelii		100.0	0.35	GLU 99			
42777702	**Acceptable identification**						**42777702**
Actinomyces israelii		90.49	0.39	GLU 99			
Bifidobacterium adolescentis		5.64	0.12	RHA 7			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		3.87	0.06				
43000000	**Low discrimination**						**43000000**
					SPO		
Clostridium botulinum biovar A		81.60	0.81	SUC 7	99		
Atopobium minutum		8.70	0.72		1		
Clostridium sporogenes		8.05	0.61		99		
43000001	**Low discrimination**						**43000001**
					SPO		
Clostridium botulinum biovar A		58.67	0.47	SUC 7	99		
Clostridium novyi biovar A		28.14	0.53	URE 1	99		
Atopobium minutum		6.26	0.38		1		
43000002	**Low discrimination**						**43000002**
					SPO		
Atopobium minutum		61.38	0.69	SUC 5	1		
Clostridium sporogenes		28.40	0.53		99		
Clostridium botulinum biovar A		8.72	0.47		99		
43000003	**Low discrimination**						**43000003**
					SPO		
Clostridium novyi biovar A		42.59	0.40	SUC 1	99		
Atopobium minutum		35.78	0.35		1		
Clostridium sporogenes		16.55	0.19		99		
43000010	**Low discrimination**						**43000010**
					SPO		
Clostridium botulinum biovar A		58.51	0.47	SUC 7	99		
Clostridium novyi biovar A		28.06	0.53	MLZ 1	99		
Atopobium minutum		6.24	0.38		1		
43000011	**Good identification**						**43000011**
Clostridium novyi biovar A		97.94	0.53	SUC 1			
Clostridium botulinum biovar A		2.06	0.13				
43000012	**Low discrimination**						**43000012**
					SPO		
Clostridium novyi biovar A		42.59	0.40	SUC 1	99		
Atopobium minutum		35.78	0.35		1		
Clostridium sporogenes		16.55	0.19		99		
43000013	**Good identification**						**43000013**
Clostridium novyi biovar A		100.0	0.40	SUC 1			
43000020	**Low discrimination**						**43000020**
					SPO		
Atopobium minutum		72.49	0.64	SUC 5	1		
Clostridium botulinum biovar A		20.59	0.47		99		
Clostridium novyi biovar A		2.47	0.43		99		
43000021	**Low discrimination**						**43000021**
					SPO		
Clostridium novyi biovar A		72.42	0.43	SUC 1	99		
Atopobium minutum		21.48	0.30		1		
Clostridium botulinum biovar A		6.10	0.13		99		
43000022	**Low discrimination**						**43000022**
					GRA	SPO	
Atopobium minutum		85.70	0.61	SUC 5	99	1	
Pseudoflavonifractor capillosus		11.72	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		1.20	0.19		99	99	
43000030	**Low discrimination**						**43000030**
					SPO		
Clostridium novyi biovar A		72.42	0.43	SUC 1	99		
Atopobium minutum		21.48	0.30		1		
Clostridium botulinum biovar A		6.10	0.13		99		
43000031	**Good identification**						**43000031**
Clostridium novyi biovar A		100.0	0.43	SUC 1			
43000040	**Very good identification**						**43000040**
					COC	GRA	
Clostridium botulinum biovar A		97.33	0.72	SUC 7	1	99	
Clostridium novyi biovar A		1.05	0.50		1	99	
Megasphaera elsdenii		0.58	0.38		99	1	
43000041	**Low discrimination**						**43000041**
					COC	GRA	
Clostridium novyi biovar A		51.55	0.50	SUC 1	1	99	
Clostridium botulinum biovar A		48.16	0.38		1	99	
Megasphaera elsdenii		0.28	0.04		99	1	
43000042	**Low discrimination**						**43000042**
					SPO		
Clostridium botulinum biovar A		62.42	0.38	SUC 7	99		
Atopobium minutum		14.86	0.35	BGL 1	1		
Clostridium novyi biovar A		11.79	0.37		99		
43000043	**Acceptable identification**						**43000043**
Clostridium novyi biovar A		94.92	0.37	SUC 1			
Clostridium botulinum biovar A		5.08	0.04				
43000050	**Low discrimination**						**43000050**
					COC	GRA	
Clostridium novyi biovar A		51.55	0.50	SUC 1	1	99	
Clostridium botulinum biovar A		48.16	0.38		1	99	
Megasphaera elsdenii		0.28	0.04		99	1	
43000051	**Good identification**						**43000051**
Clostridium novyi biovar A		99.07	0.50	SUC 1			
Clostridium botulinum biovar A		0.94	0.04				
43000052	**Acceptable identification**						**43000052**
Clostridium novyi biovar A		94.92	0.37	SUC 1			
Clostridium botulinum biovar A		5.08	0.04				
43000053	**Good identification**						**43000053**
Clostridium novyi biovar A		100.0	0.37	SUC 1			
43000060	**Low discrimination**						**43000060**
					SPO		
Clostridium botulinum biovar A		71.56	0.38	SUC 7	99		
Clostridium novyi biovar A		19.15	0.40	NAG 1	99		
Atopobium minutum		8.52	0.30		1		
43000061	**Good identification**						**43000061**
Clostridium novyi biovar A		96.36	0.40	SUC 1			

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Clostridium botulinum biovar A		3.64	0.04				
43000070	**Good identification**						**43000070**
Clostridium novyi biovar A		96.36	0.40	SUC 1			
Clostridium botulinum biovar A		3.64	0.04				
43000071	**Good identification**						**43000071**
Clostridium novyi biovar A		100.0	0.40	SUC 1			
43000100	**Low discrimination**						**43000100**
					GRA	SPO	
Clostridium botulinum biovar A		64.11	0.47	SUC 7	99	99	
Fusobacterium mortiferum		13.29	0.48	LAC 1	1	1	
Actinomyces meyeri		8.14	0.35		99	1	
43000140	**Good identification**						**43000140**
					GRA	SPO	
Clostridium botulinum biovar A		97.68	0.38	SUC 7	99	99	
Clostridium novyi biovar A		1.06	0.16	LAC 1	99	99	
Fusobacterium mortiferum		0.69	0.14		1	1	
43000200	**Low discrimination**						**43000200**
					SPO		
Clostridium botulinum biovar A		81.75	0.47	SUC 7	99		
Atopobium minutum		8.72	0.38	RHA 1	1		
Clostridium sporogenes		8.07	0.27		99		
43000240	**Good identification**						**43000240**
					COC	GRA	
Clostridium botulinum biovar A		98.36	0.38	SUC 7	1	99	
Clostridium novyi biovar A		1.06	0.16	RHA 1	1	99	
Megasphaera elsdenii		0.58	0.04		99	1	
43000400	**Low discrimination**						**43000400**
					SPO		
Clostridium botulinum biovar A		81.13	0.47	SUC 7	99		
Atopobium minutum		8.65	0.38	ARA 1	1		
Clostridium sporogenes		8.01	0.27		99		
43000440	**Good identification**						**43000440**
					COC	GRA	
Clostridium botulinum biovar A		98.36	0.38	SUC 7	1	99	
Clostridium novyi biovar A		1.06	0.16	ARA 1	1	99	
Megasphaera elsdenii		0.58	0.04		99	1	
43001000	**Low discrimination**						**43001000**
					SPO		
Eubacterium tortuosum		81.18	0.66	SAL 99	1		
Atopobium minutum		13.32	0.64		1		
Clostridium botulinum biovar A		3.78	0.47		99		
43001002	**Low discrimination**						**43001002**
					SPO		
Eubacterium tortuosum		60.86	0.53	SAL 99	1		
Atopobium minutum		37.71	0.61		1		
Clostridium sporogenes		0.53	0.19		99		
43001011	**Good identification**						**43001011**
					SPO		
Clostridium novyi biovar A		96.05	0.37	SUC 1	99		
Eubacterium tortuosum		3.95	-.03	GAL 10	1		

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43001020	**Low discrimination**							**43001020**
Eubacterium tortuosum		75.62	0.53	SAL 99				
Atopobium minutum		23.43	0.56					
Eubacterium saburreum		0.36	0.28					
43001022	**Low discrimination**							**43001022**
					GRA			
Atopobium minutum		52.44	0.53	SUC 5	99			
Eubacterium tortuosum		44.80	0.40		99			
Pseudoflavonifractor capillosus		2.39	0.15		1			
43001040	**Low discrimination**							**43001040**
					SPO			
Clostridium botulinum biovar A		49.99	0.38	SUC 7	99			
Eubacterium tortuosum		36.28	0.32	GAL 1	1			
Atopobium minutum		5.95	0.30		1			
43001100	**Low discrimination**							**43001100**
					GRA			
Fusobacterium mortiferum		30.76	0.51	FRU 99	1			
Eubacterium saburreum		23.48	0.47		85			
Eubacterium tortuosum		21.45	0.32		99			
43001101	**Good identification**							**43001101**
					SPO	GRA		
Clostridium chauvoei		97.22	0.40	MNS 90	99	99		
Fusobacterium mortiferum		0.91	0.17	MLT 99	1	1		
Eubacterium saburreum		0.69	0.13		1	85		
43001110	**Good identification**							**43001110**
					GRA			
Eubacterium saburreum		96.40	0.46	IND 99	85			
Fusobacterium mortiferum		1.56	0.17		1			
Eubacterium tortuosum		1.09	-.03		99			
43002000	**Low discrimination**							**43002000**
					COC	SPO	GRA	
Clostridium botulinum biovar A		59.95	0.47	SUC 7	1	99	99	
Megasphaera elsdenii		15.70	0.41	MAN 1	99	1	1	
Eubacterium limosum		9.80	0.29		1	1	99	
43002040	**Low discrimination**							**43002040**
					COC	GRA		
Clostridium botulinum biovar A		62.47	0.38	SUC 7	1	99		
Megasphaera elsdenii		36.51	0.38	MAN 1	99	1		
Clostridium novyi biovar A		0.68	0.16		1	99		
43004000	**Low discrimination**							**43004000**
					SPO			
Clostridium botulinum biovar A		72.68	0.47	SUC 7	99			
Atopobium minutum		7.75	0.38	XYL 1	1			
Clostridium sporogenes		7.17	0.27		99			
43004040	**Low discrimination**							**43004040**
Clostridium glycolicum		83.84	0.49	SUC 1				
Clostridium botulinum biovar A		15.90	0.38	FRU 90				
Clostridium novyi biovar A		0.17	0.16					
43004100	**Acceptable identification**							**43004100**
					GRA	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		90.56	0.40	ESL 1	99	1	
Fusobacterium mortiferum		5.73	0.29	FRU 95	1	1	
Clostridium botulinum biovar A		3.71	0.13		99	99	
43005010	**Good identification**						**43005010**
Eubacterium saburreum		95.34	0.35	IND 99			
Eubacterium tortuosum		4.66	-.03				
43005100	**Low discrimination**						**43005100**
					GRA		
Actinomyces meyeri		70.39	0.43	ESL 1	99		
Eubacterium saburreum		24.31	0.43	FRU 95	85		
Fusobacterium mortiferum		4.45	0.32		1		
43005110	**Good identification**						**43005110**
Eubacterium saburreum		96.55	0.41	IND 99			
Actinomyces meyeri		3.45	0.09				
43010000	**Low discrimination**						**43010000**
					GRA	SPO	
Atopobium minutum		27.42	0.71	SUC 5	99	1	
Fusobacterium mortiferum		24.74	0.76		1	1	
Clostridium botulinum biovar A		17.78	0.60		99	99	
43010001	**Low discrimination**						**43010001**
					GRA	SPO	
Atopobium minutum		23.91	0.37	SUC 5	99	1	
Fusobacterium mortiferum		21.57	0.42	URE 1	1	1	
Clostridium botulinum biovar A		15.50	0.26		99	99	
43010002	**Low discrimination**						**43010002**
					GRA	SPO	
Atopobium minutum		74.15	0.68	SUC 5	99	1	
Fusobacterium mortiferum		11.15	0.60		1	1	
Clostridium sporogenes		6.56	0.40		99	99	
43010010	**Low discrimination**						**43010010**
					GRA	SPO	
Clostridium difficile		72.76	0.45	SUC 1	99	99	
Atopobium minutum		6.53	0.37	MAN 90	99	1	
Fusobacterium mortiferum		5.90	0.42		1	1	
43010011	**Low discrimination**						**43010011**
Clostridium novyi biovar A		82.79	0.37	SUC 1			
Clostridium difficile		17.21	0.10	FRU 10			
43010020	**Low discrimination**						**43010020**
					GRA		
Atopobium minutum		88.16	0.63	SUC 5	99		
Fusobacterium mortiferum		2.41	0.42		1		
Actinomyces meyeri		2.23	0.32		99		
43010022	**Very good identification**						**43010022**
					GRA		
Atopobium minutum		96.96	0.60	SUC 5	99		
Pseudoflavonifractor capillosus		1.60	0.15		1		
Fusobacterium mortiferum		0.44	0.25		1		
43010040	**Low discrimination**						**43010040**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		52.52	0.60	ESL 1	99	1		
Clostridium botulinum biovar A		29.89	0.51		1	99		
Clostridium glycolicum		11.82	0.43		1	99		
43010042	**Low discrimination**							**43010042**
					COC	SPO	GRA	
Megasphaera elsdenii		74.97	0.44	ESL 1	99	1	1	
Atopobium minutum		13.35	0.34	BGL 10	1	1	99	
Clostridium botulinum biovar A		3.88	0.17		1	99	99	
43010100	**Low discrimination**							**43010100**
					GRA			
Fusobacterium mortiferum		88.42	0.82		1			
Actinomyces meyeri		10.40	0.57		99			
Atopobium minutum		0.42	0.37		99			
43010101	**Low discrimination**							**43010101**
					GRA	SPO		
Fusobacterium mortiferum		86.51	0.48	URE 1	1	1		
Actinomyces meyeri		10.18	0.23		99	1		
Clostridium chauvoei		3.32	0.12		99	99		
43010102	**Good identification**							**43010102**
					GRA			
Fusobacterium mortiferum		93.62	0.66	BGL 10	1			
Actinomyces meyeri		3.07	0.31		99			
Atopobium minutum		2.70	0.34		99			
43010110	**Low discrimination**							**43010110**
					GRA	SPO		
Fusobacterium mortiferum		83.37	0.48	MLZ 1	1	1		
Actinomyces meyeri		9.81	0.23		99	1		
Clostridium difficile		4.46	0.10		99	99		
43010120	**Low discrimination**							**43010120**
					GRA			
Actinomyces meyeri		57.62	0.42	ESL 1	99			
Fusobacterium mortiferum		36.28	0.48	NAG 12	1			
Atopobium minutum		5.75	0.29		99			
43010140	**Low discrimination**							**43010140**
					COC	GRA		
Fusobacterium mortiferum		71.04	0.48	SOR 1	1	1		
Megasphaera elsdenii		11.48	0.26		99	1		
Actinomyces meyeri		8.36	0.23		1	99		
43010200	**Low discrimination**							**43010200**
					GRA	SPO		
Atopobium minutum		27.58	0.37	SUC 5	99	1		
Fusobacterium mortiferum		24.88	0.42	RHA 1	1	1		
Clostridium botulinum biovar A		17.88	0.26		99	99		
43010300	**Low discrimination**							**43010300**
					GRA			
Fusobacterium mortiferum		89.47	0.48	RHA 1	1			
Actinomyces meyeri		10.53	0.23		99			
43010400	**Low discrimination**							**43010400**
					GRA			
Actinomyces meyeri		29.82	0.36	ESL 1	99			
Atopobium minutum		19.46	0.37		99			
Fusobacterium mortiferum		17.56	0.42		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43010500	**Low discrimination**							**43010500**
					GRA			
Actinomyces meyeri		74.09	0.47	ESL 1	99			
Fusobacterium mortiferum		25.45	0.48		1			
Lactobacillus fermentum		0.47	0.20		99			
43011000	**Low discrimination**							**43011000**
					GRA			
Fusobacterium mortiferum		72.12	0.79		1			
Atopobium minutum		17.76	0.63		99			
Actinomyces meyeri		4.95	0.50		99			
43011001	**Low discrimination**							**43011001**
					GRA			
Fusobacterium mortiferum		67.74	0.45	URE 1	1			
Atopobium minutum		16.69	0.29		99			
Actinomyces meyeri		4.65	0.16		99			
43011002	**Low discrimination**							**43011002**
					GRA			
Atopobium minutum		54.19	0.60	SUC 5	99			
Fusobacterium mortiferum		36.67	0.63		1			
Lactobacillus fermentum		5.80	0.50		99			
43011010	**Low discrimination**							**43011010**
					GRA			
Eubacterium saburreum		55.93	0.48	IND 99	85			
Fusobacterium mortiferum		29.85	0.45		1			
Atopobium minutum		7.35	0.29		99			
43011020	**Low discrimination**							**43011020**
					GRA			
Atopobium minutum		71.41	0.55	SUC 5	99			
Fusobacterium mortiferum		8.79	0.45		1			
Actinomyces meyeri		8.14	0.35		99			
43011022	**Low discrimination**							**43011022**
					GRA			
Atopobium minutum		86.15	0.52	SUC 5	99			
Lactobacillus fermentum		9.22	0.42		99			
Fusobacterium mortiferum		1.77	0.28		1			
43011040	**Low discrimination**							**43011040**
					COC	GRA	SPO	
Fusobacterium mortiferum		41.35	0.45	SOR 1	1	1	1	
Clostridium glycolicum		25.73	0.27		1	99	99	
Megasphaera elsdenii		10.40	0.26		99	1	1	
43011100	**Low discrimination**							**43011100**
					GRA			
Fusobacterium mortiferum		86.61	0.85		1			
Actinomyces meyeri		10.19	0.60		99			
Eubacterium saburreum		2.00	0.56		85			
43011101	**Low discrimination**							**43011101**
					GRA	SPO		
Clostridium chauvoei		68.70	0.47	MNS 90	99	99		
Fusobacterium mortiferum		27.14	0.51	MLT 99	1	1		
Actinomyces meyeri		3.19	0.26		99	1		
43011102	**Low discrimination**							**43011102**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		83.98	0.69	BGL 10	1		
Lactobacillus fermentum		9.29	0.54		99		
Eubacterium saburreum		3.09	0.43		85		
43011110	**Low discrimination**						**43011110**
					GRA		
Eubacterium saburreum		62.38	0.54	IND 99	85		
Fusobacterium mortiferum		33.30	0.51		1		
Actinomyces meyeri		3.92	0.26		99		
43011112	**Low discrimination**						**43011112**
					GRA		
Eubacterium saburreum		72.06	0.41	IND 99	85		
Fusobacterium mortiferum		24.22	0.35		1		
Lactobacillus fermentum		2.68	0.20		99		
43011120	**Low discrimination**						**43011120**
					GRA		
Actinomyces meyeri		38.09	0.45	ESL 1	99		
Fusobacterium mortiferum		23.98	0.51	NAG 12	1		
Prevotella melaninogenica		15.43	0.30		1		
43011121	**Good identification**						**43011121**
					GRA	SPO	
Clostridium chauvoei		96.28	0.38	MNS 90	99	99	
Actinomyces meyeri		1.83	0.11	MLT 99	99	1	
Fusobacterium mortiferum		1.15	0.17		1	1	
43011122	**Low discrimination**						**43011122**
					GRA	CAT	
Lactobacillus fermentum		33.90	0.46	ESL 1	99	1	
Capnocytophaga ochracea		32.31	0.30		1	90	
Prevotella melaninogenica		10.24	0.18		1	1	
43011130	**Low discrimination**						**43011130**
					GRA		
Eubacterium saburreum		89.84	0.41	IND 99	85		
Actinomyces meyeri		4.36	0.11		99		
Fusobacterium mortiferum		2.75	0.17		1		
43011140	**Low discrimination**						**43011140**
					GRA		
Fusobacterium mortiferum		86.73	0.51	SOR 1	1		
Actinomyces meyeri		10.20	0.26		99		
Eubacterium saburreum		2.01	0.21		85		
43011200	**Low discrimination**						**43011200**
					GRA		
Fusobacterium mortiferum		72.88	0.45	RHA 1	1		
Atopobium minutum		17.95	0.29		99		
Actinomyces meyeri		5.00	0.16		99		
43011300	**Low discrimination**						**43011300**
					GRA		
Fusobacterium mortiferum		86.73	0.51	RHA 1	1		
Actinomyces meyeri		10.20	0.26		99		
Eubacterium saburreum		2.01	0.21		85		
43011400	**Low discrimination**						**43011400**
					GRA		
Lactobacillus fermentum		36.82	0.50	ESL 1	99		
Actinomyces meyeri		35.96	0.39		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		21.17	0.45		1			
43011402	**Low discrimination**							**43011402**
					GRA			
Lactobacillus fermentum		84.14	0.50	ESL 1	99			
Atopobium minutum		7.94	0.26		99			
Fusobacterium mortiferum		5.38	0.28		1			
43011420	**Low discrimination**							**43011420**
Lactobacillus fermentum		64.89	0.42	ESL 1				
Actinomyces meyeri		25.92	0.25					
Atopobium minutum		9.19	0.21					
43011422	**Acceptable identification**							**43011422**
Lactobacillus fermentum		91.37	0.42	ESL 1				
Atopobium minutum		8.63	0.18					
43011500	**Low discrimination**							**43011500**
					GRA			
Actinomyces meyeri		56.53	0.50	ESL 1	99			
Lactobacillus fermentum		23.61	0.54		99			
Fusobacterium mortiferum		19.41	0.51		1			
43011502	**Low discrimination**							**43011502**
					GRA			
Lactobacillus fermentum		85.35	0.54	ESL 1	99			
Fusobacterium mortiferum		7.80	0.35		1			
Actinomyces meyeri		6.32	0.24		99			
43011520	**Low discrimination**							**43011520**
Lactobacillus fermentum		47.55	0.46	ESL 1				
Actinomyces meyeri		46.57	0.35					
Bifidobacterium longum subsp. longum		1.63	0.14					
43011522	**Low discrimination**							**43011522**
					GRA	CAT		
Lactobacillus fermentum		81.12	0.46	ESL 1	99	1		
Capnocytophaga ochracea		13.64	0.17		1	90		
Bifidobacterium longum subsp. longum		2.79	0.14		99	1		
43012000	**Low discrimination**							**43012000**
					COC	SPO	GRA	
Eubacterium limosum		68.48	0.63	SUC 1	1	1	99	
Megasphaera elsdenii		21.05	0.63		99	1	1	
Clostridium difficile		8.55	0.49		1	99	99	
43012002	**Low discrimination**							**43012002**
					COC	GRA		
Megasphaera elsdenii		65.90	0.47	ESL 1	99	1		
Eubacterium limosum		19.49	0.29	BGL 10	1	99		
Atopobium minutum		7.83	0.34		1	99		
43012010	**Very good identification**							**43012010**
					COC	SPO	GRA	
Clostridium difficile		97.86	0.61	SUC 1	1	99	99	
Eubacterium limosum		1.62	0.29		1	1	99	
Megasphaera elsdenii		0.50	0.29		99	1	1	
43012040	**Good identification**							**43012040**
					COC	SPO	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		90.72	0.60	ESL 1	99	1	1	
Eubacterium limosum		4.47	0.29		1	1	99	
Clostridium difficile		2.91	0.27		1	99	99	
43012042	**Good identification**							**43012042**
					COC	GRA		
Megasphaera elsdenii		98.29	0.44	ESL 1	99	1		
Pseudoramibacter alactolyticus		1.27	0.00	BGL 10	1	99		
Eubacterium limosum		0.44	-.06		1	99		
43012050	**Acceptable identification**							**43012050**
					COC	SPO	GRA	
Clostridium difficile		93.66	0.39	SUC 1	1	99	99	
Megasphaera elsdenii		6.04	0.26	SOR 5	99	1	1	
Eubacterium limosum		0.30	-.06		1	1	99	
43012100	**Low discrimination**							**43012100**
					COC	GRA		
Fusobacterium mortiferum		44.14	0.48	MAN 1	1	1		
Eubacterium limosum		34.81	0.29		1	99		
Megasphaera elsdenii		10.70	0.29		99	1		
43013100	**Low discrimination**							**43013100**
					GRA			
Fusobacterium mortiferum		85.83	0.51	MAN 1	1			
Actinomyces meyeri		10.10	0.26		99			
Eubacterium saburreum		1.99	0.21		85			
43014000	**Low discrimination**							**43014000**
					GRA	SPO		
Actinomyces meyeri		40.99	0.52	ESL 1	99	1		
Clostridium glycolicum		26.11	0.43		99	99		
Fusobacterium mortiferum		23.16	0.57		1	1		
43014002	**Low discrimination**							**43014002**
					GRA			
Fusobacterium mortiferum		37.58	0.40	XYL 7	1			
Atopobium minutum		33.54	0.34	BGL 10	99			
Actinomyces meyeri		18.51	0.26		99			
43014020	**Low discrimination**							**43014020**
					SPO			
Actinomyces meyeri		77.25	0.37	ESL 1	1			
Atopobium minutum		15.87	0.29	NAG 12	1			
Clostridium glycolicum		3.65	0.09		99			
43014040	**Very good identification**							**43014040**
					COC	GRA		
Clostridium glycolicum		99.50	0.65	SUC 1	1	99		
Megasphaera elsdenii		0.24	0.26		99	1		
Clostridium botulinum biovar A		0.13	0.17		1	99		
43014100	**Low discrimination**							**43014100**
					GRA	SPO		
Actinomyces meyeri		75.05	0.62	ESL 1	99	1		
Fusobacterium mortiferum		24.74	0.63		1	1		
Clostridium glycolicum		0.12	0.09		99	99		
43014102	**Low discrimination**							**43014102**
					GRA			
Fusobacterium mortiferum		53.83	0.47	XYL 7	1			
Actinomyces meyeri		45.46	0.36	BGL 10	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		0.70	0.16		99		
43014120	**Good identification**						**43014120**
					GRA		
Actinomyces meyeri		97.62	0.47	ESL 1	99		
Fusobacterium mortiferum		2.38	0.29	NAG 12	1		
43014400	**Acceptable identification**						**43014400**
					GRA	SPO	
Actinomyces meyeri		94.94	0.41	ESL 1	99	1	
Clostridium glycolicum		2.44	0.09		99	99	
Fusobacterium mortiferum		2.17	0.23		1	1	
43014500	**Good identification**						**43014500**
					GRA		
Actinomyces meyeri		98.50	0.52	ESL 1	99		
Fusobacterium mortiferum		1.31	0.29		1		
Lactobacillus fermentum		0.19	0.16		99		
43014520	**Good identification**						**43014520**
Actinomyces meyeri		100.0	0.37	ESL 1			
43015000	**Low discrimination**						**43015000**
					GRA		
Actinomyces meyeri		55.46	0.55	ESL 1	99		
Fusobacterium mortiferum		31.34	0.60		1		
Eubacterium saburreum		5.19	0.45		85		
43015002	**Low discrimination**						**43015002**
					GRA		
Lactobacillus fermentum		38.71	0.46	ESL 1	99		
Fusobacterium mortiferum		31.37	0.43		1		
Actinomyces meyeri		15.46	0.29		99		
43015010	**Low discrimination**						**43015010**
					GRA		
Eubacterium saburreum		82.18	0.43	IND 99	85		
Actinomyces meyeri		10.85	0.21		99		
Fusobacterium mortiferum		6.13	0.26		1		
43015020	**Low discrimination**						**43015020**
Actinomyces meyeri		71.54	0.40	ESL 1			
Lactobacillus fermentum		13.55	0.38	NAG 12			
Eubacterium saburreum		8.66	0.32				
43015022	**Low discrimination**						**43015022**
Lactobacillus fermentum		69.60	0.38	ESL 1			
Actinomyces meyeri		11.37	0.14				
Atopobium minutum		11.19	0.18				
43015040	**Good identification**						**43015040**
					GRA	SPO	
Clostridium glycolicum		98.08	0.49	SUC 1	99	99	
Actinomyces meyeri		1.11	0.21	GAL 10	99	1	
Fusobacterium mortiferum		0.62	0.26		1	1	
43015100	**Low discrimination**						**43015100**
					GRA		
Actinomyces meyeri		70.64	0.65	ESL 1	99		
Fusobacterium mortiferum		23.28	0.66		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		3.85	0.51		85		
43015102	**Low discrimination**						**43015102**
					GRA		
Fusobacterium mortiferum		33.52	0.50	XYL 7	1		
Lactobacillus fermentum		28.92	0.50	BGL 10	99		
Actinomyces meyeri		28.31	0.39		99		
43015110	**Low discrimination**						**43015110**
					GRA		
Eubacterium saburreum		76.45	0.49	IND 99	85		
Actinomyces meyeri		17.31	0.31		99		
Fusobacterium mortiferum		5.70	0.32		1		
43015112	**Low discrimination**						**43015112**
					GRA		
Eubacterium saburreum		88.72	0.37	IND 99	85		
Fusobacterium mortiferum		4.17	0.16		1		
Lactobacillus fermentum		3.60	0.16		99		
43015120	**Low discrimination**						**43015120**
Actinomyces meyeri		84.90	0.50	ESL 1			
Lactobacillus fermentum		6.56	0.42	NAG 12			
Eubacterium saburreum		5.99	0.38				
43015122	**Low discrimination**						**43015122**
					GRA	CAT	
Lactobacillus fermentum		48.26	0.42	ESL 1	99	1	
Capnocytophaga ochracea		19.58	0.20		1	90	
Actinomyces meyeri		19.33	0.24		99	1	
43015130	**Low discrimination**						**43015130**
Eubacterium saburreum		85.11	0.37	IND 99			
Actinomyces meyeri		14.89	0.16				
43015400	**Low discrimination**						**43015400**
					GRA		
Actinomyces meyeri		74.83	0.44	ESL 1	99		
Lactobacillus fermentum		23.18	0.46		99		
Fusobacterium mortiferum		1.71	0.26		1		
43015402	**Acceptable identification**						**43015402**
Lactobacillus fermentum		90.92	0.46	ESL 1			
Actinomyces meyeri		9.08	0.19				
43015422	**Good identification**						**43015422**
Lactobacillus fermentum		96.08	0.38	ESL 1			
Actinomyces meyeri		3.92	0.04				
43015500	**Low discrimination**						**43015500**
					GRA		
Actinomyces meyeri		87.43	0.55	ESL 1	99		
Lactobacillus fermentum		11.05	0.50		99		
Fusobacterium mortiferum		1.16	0.32		1		
43015502	**Low discrimination**						**43015502**
Lactobacillus fermentum		77.74	0.50	ESL 1			
Actinomyces meyeri		19.03	0.29				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		1.19	0.14				
43015520	**Low discrimination**						**43015520**
Actinomyces meyeri		73.36	0.40	ESL 1			
Lactobacillus fermentum		22.66	0.42	NAG 12			
Bifidobacterium longum subsp. longum		3.97	0.22				
43015522	**Low discrimination**						**43015522**
Lactobacillus fermentum		74.24	0.42	ESL 1			
Bifidobacterium longum subsp. longum		13.02	0.22				
Actinomyces meyeri		7.43	0.14				
43016010	**Good identification**						**43016010**
					SPO		
Clostridium difficile		99.68	0.39	SUC 1	99		
Eubacterium limosum		0.32	-.06	XYL 5	1		
43020000	**Low discrimination**						**43020000**
					SPO		
Clostridium botulinum biovar A		87.90	0.71	SUC 7	99		
Clostridium sporogenes		11.57	0.53		99		
Atopobium minutum		0.38	0.38		1		
43020001	**Low discrimination**						**43020001**
Clostridium botulinum biovar A		86.88	0.36	SUC 7			
Clostridium sporogenes		11.43	0.19	URE 1			
Clostridium novyi biovar A		1.68	0.19				
43020002	**Low discrimination**						**43020002**
					SPO		
Clostridium sporogenes		77.01	0.45	SUC 1	99		
Clostridium botulinum biovar A		17.74	0.36		99		
Atopobium minutum		5.04	0.35		1		
43020010	**Low discrimination**						**43020010**
Clostridium botulinum biovar A		86.88	0.36	SUC 7			
Clostridium sporogenes		11.43	0.19	MLZ 1			
Clostridium novyi biovar A		1.68	0.19				
43020020	**Low discrimination**						**43020020**
					SPO		
Clostridium botulinum biovar A		78.13	0.36	SUC 7	99		
Atopobium minutum		11.11	0.30	NAG 1	1		
Clostridium sporogenes		10.28	0.19		99		
43020040	**Very good identification**						**43020040**
Clostridium botulinum biovar A		99.49	0.62	SUC 7			
Clostridium sporogenes		0.44	0.19				
Clostridium novyi biovar A		0.04	0.16				
43020100	**Low discrimination**						**43020100**
					GRA	SPO	
Clostridium botulinum biovar A		72.17	0.36	SUC 7	99	99	
Fusobacterium mortiferum		17.87	0.39	LAC 1	1	1	
Clostridium sporogenes		9.50	0.19		99	99	
43020200	**Low discrimination**						**43020200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		88.37	0.36	SUC 7			
Clostridium sporogenes		11.63	0.19	RHA 1			
43020400	**Low discrimination**						**43020400**
Clostridium botulinum biovar A		88.37	0.36	SUC 7			
Clostridium sporogenes		11.63	0.19	ARA 1			
43021000	**Low discrimination**						**43021000**
					SPO		
Clostridium botulinum biovar A		38.89	0.36	SUC 7	99		
Eubacterium tortuosum		33.72	0.32	GAL 1	1		
Eubacterium saburreum		10.55	0.38		1		
43021010	**Acceptable identification**						**43021010**
					SPO		
Eubacterium saburreum		92.17	0.37	IND 99	1		
Clostridium botulinum biovar A		4.20	0.02		99		
Eubacterium tortuosum		3.64	-.03		1		
43021100	**Low discrimination**						**43021100**
					SPO	GRA	
Eubacterium saburreum		61.39	0.44	IND 99	1	85	
Fusobacterium mortiferum		36.04	0.42		1	1	
Clostridium botulinum biovar A		0.98	0.02		99	99	
43021110	**Good identification**						**43021110**
Eubacterium saburreum		100.0	0.43	IND 99			
43022000	**Low discrimination**						**43022000**
					COC	GRA	
Clostridium botulinum biovar A		86.59	0.36	SUC 7	1	99	
Clostridium sporogenes		11.39	0.19	MAN 1	1	99	
Megasphaera elsdenii		0.92	0.07		99	1	
43024000	**Low discrimination**						**43024000**
					GRA	SPO	
Clostridium botulinum biovar A		87.75	0.36	SUC 7	99	99	
Clostridium sporogenes		11.55	0.19	XYL 1	99	99	
Fusobacterium mortiferum		0.70	0.14		1	1	
43025100	**Low discrimination**						**43025100**
					GRA		
Eubacterium saburreum		88.82	0.40	IND 99	85		
Fusobacterium mortiferum		7.29	0.23		1		
Actinomyces meyeri		3.90	0.09		99		
43025110	**Good identification**						**43025110**
Eubacterium saburreum		100.0	0.38	IND 99			
43030000	**Low discrimination**						**43030000**
					GRA	SPO	
Fusobacterium mortiferum		52.40	0.67		1	1	
Clostridium botulinum biovar A		31.52	0.50		99	99	
Clostridium sporogenes		11.47	0.40		99	99	
43030002	**Low discrimination**						**43030002**
					GRA	SPO	
Fusobacterium mortiferum		50.53	0.51	BGL 10	1	1	
Clostridium sporogenes		33.17	0.32		99	99	
Atopobium minutum		11.36	0.34		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43030040	**Low discrimination**							**43030040**
					COC	GRA	SPO	
Clostridium botulinum biovar A		86.23	0.41	SUC 7	1	99	99	
Megasphaera elsdenii		6.12	0.26	FRU 6	99	1	1	
Fusobacterium mortiferum		4.85	0.33		1	1	1	
43030100	**Very good identification**							**43030100**
					GRA	SPO		
Fusobacterium mortiferum		99.18	0.73		1	1		
Actinomyces meyeri		0.40	0.23		99	1		
Clostridium botulinum biovar A		0.26	0.16		99	99		
43030101	**Good identification**							**43030101**
Fusobacterium mortiferum		100.0	0.39	URE 1				
43030102	**Very good identification**							**43030102**
					GRA	SPO		
Fusobacterium mortiferum		99.72	0.57	BGL 10	1	1		
Clostridium sporogenes		0.28	-.02		99	99		
43030110	**Acceptable identification**							**43030110**
Fusobacterium mortiferum		94.04	0.39	MLZ 1				
Eubacterium saburreum		5.96	0.17					
43030120	**Acceptable identification**							**43030120**
					GRA			
Fusobacterium mortiferum		94.90	0.39	NAG 1	1			
Actinomyces meyeri		5.10	0.08		99			
43030140	**Acceptable identification**							**43030140**
					GRA	SPO		
Fusobacterium mortiferum		92.85	0.39	SOR 1	1	1		
Clostridium botulinum biovar A		7.15	0.07		99	99		
43030300	**Good identification**							**43030300**
Fusobacterium mortiferum		100.0	0.39	RHA 1				
43030500	**Acceptable identification**							**43030500**
					GRA			
Fusobacterium mortiferum		91.04	0.39	ARA 1	1			
Actinomyces meyeri		8.96	0.13		99			
43031000	**Good identification**							**43031000**
					GRA			
Fusobacterium mortiferum		92.93	0.70		1			
Eubacterium saburreum		4.80	0.46		85			
Atopobium minutum		0.77	0.29		99			
43031001	**Good identification**							**43031001**
Fusobacterium mortiferum		95.09	0.36	URE 1				
Eubacterium saburreum		4.91	0.12					
43031002	**Low discrimination**							**43031002**
					GRA			
Fusobacterium mortiferum		86.38	0.54	BGL 10	1			
Eubacterium saburreum		7.08	0.33		85			
Atopobium minutum		4.32	0.26		99			
43031010	**Low discrimination**							**43031010**
					CAT	GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		79.23	0.45	IND 99	1	85	
Fusobacterium mortiferum		18.95	0.36		5	1	
Propionibacterium granulosum		1.82	0.15		90	99	
43031020	**Low discrimination**						**43031020**
					GRA		
Fusobacterium mortiferum		41.05	0.36	NAG 1	1		
Eubacterium saburreum		37.02	0.33		85		
Atopobium minutum		11.28	0.21		99		
43031040	**Low discrimination**						**43031040**
					SPO	GRA	
Fusobacterium mortiferum		85.37	0.36	SOR 1	1	1	
Clostridium botulinum biovar A		10.22	0.07		99	99	
Eubacterium saburreum		4.41	0.12		1	85	
43031100	**Good identification**						**43031100**
					GRA		
Fusobacterium mortiferum		94.51	0.76		1		
Eubacterium saburreum		4.88	0.53		85		
Actinomyces meyeri		0.38	0.26		99		
43031101	**Low discrimination**						**43031101**
					SPO	GRA	
Fusobacterium mortiferum		50.17	0.42	URE 1	1	1	
Clostridium chauvoei		47.24	0.30		99	99	
Eubacterium saburreum		2.59	0.18		1	85	
43031102	**Good identification**						**43031102**
					GRA		
Fusobacterium mortiferum		91.83	0.60	BGL 10	1		
Eubacterium saburreum		7.53	0.40		85		
Lactobacillus fermentum		0.34	0.20		99		
43031110	**Low discrimination**						**43031110**
Eubacterium saburreum		80.70	0.51	IND 99			
Fusobacterium mortiferum		19.30	0.42				
43031112	**Low discrimination**						**43031112**
Eubacterium saburreum		86.91	0.38	IND 99			
Fusobacterium mortiferum		13.09	0.26				
43031120	**Low discrimination**						**43031120**
					GRA		
Fusobacterium mortiferum		43.97	0.42	NAG 1	1		
Eubacterium saburreum		39.66	0.40		85		
Clostridium perfringens		11.50	0.21		99		
43031130	**Good identification**						**43031130**
Eubacterium saburreum		100.0	0.38	IND 99			
43031140	**Good identification**						**43031140**
Fusobacterium mortiferum		95.09	0.42	SOR 1			
Eubacterium saburreum		4.91	0.18				
43031200	**Good identification**						**43031200**
Fusobacterium mortiferum		95.09	0.36	RHA 1			
Eubacterium saburreum		4.91	0.12				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
43031300	**Good identification**						**43031300**
Fusobacterium mortiferum		95.09	0.42	RHA 1			
Eubacterium saburreum		4.91	0.18				
43031400	**Low discrimination**						**43031400**
					GRA		
Fusobacterium mortiferum		85.63	0.36	ARA 1	1		
Lactobacillus fermentum		5.04	0.16		99		
Actinomyces meyeri		4.92	0.05		99		
43031500	**Low discrimination**						**43031500**
					GRA		
Fusobacterium mortiferum		83.95	0.42	ARA 1	1		
Actinomyces meyeri		8.27	0.16		99		
Eubacterium saburreum		4.33	0.18		85		
43032010	**Good identification**						**43032010**
					SPO	CAT	
Clostridium difficile		99.39	0.42	SUC 1	99	1	
Propionibacterium granulosum		0.39	0.13	TRE 7	1	90	
Eubacterium limosum		0.22	-.06		1	1	
43032100	**Acceptable identification**						**43032100**
					GRA		
Fusobacterium mortiferum		92.68	0.39	MAN 1	1		
Propionibacterium propionicum		2.55	0.14		99		
Eubacterium limosum		2.47	-.06		99		
43033000	**Low discrimination**						**43033000**
					CAT	GRA	
Fusobacterium mortiferum		60.14	0.36	MAN 1	5	1	
Propionibacterium granulosum		31.94	0.28		90	99	
Eubacterium saburreum		3.11	0.12		1	85	
43033100	**Acceptable identification**						**43033100**
					GRA		
Fusobacterium mortiferum		93.46	0.42	MAN 1	1		
Eubacterium saburreum		4.83	0.18		85		
Propionibacterium propionicum		1.71	0.14		99		
43034000	**Low discrimination**						**43034000**
					GRA	SPO	
Fusobacterium mortiferum		82.78	0.48	XYL 7	1	1	
Clostridium botulinum biovar A		6.68	0.16		99	99	
Actinomyces meyeri		4.95	0.18		99	1	
43034100	**Acceptable identification**						**43034100**
					GRA		
Fusobacterium mortiferum		90.24	0.54	XYL 7	1		
Actinomyces meyeri		9.26	0.28		99		
Eubacterium saburreum		0.51	0.14		85		
43034102	**Good identification**						**43034102**
					GRA		
Fusobacterium mortiferum		97.22	0.38	XYL 7	1		
Actinomyces meyeri		2.78	0.02	BGL 10	99		
43035000	**Low discrimination**						**43035000**
					GRA		
Fusobacterium mortiferum		69.75	0.51	XYL 7	1		
Eubacterium saburreum		25.76	0.42		85		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		4.17	0.21		99		
43035010	**Good identification**						**43035010**
Eubacterium saburreum		96.77	0.40	IND 99			
Fusobacterium mortiferum		3.24	0.17				
43035100	**Low discrimination**						**43035100**
					GRA		
Fusobacterium mortiferum		65.57	0.57	XYL 7	1		
Eubacterium saburreum		24.22	0.48		85		
Actinomyces meyeri		6.73	0.31		99		
43035102	**Low discrimination**						**43035102**
					COC	GRA	
Fusobacterium mortiferum		58.02	0.41	XYL 7	1	1	
Eubacterium saburreum		34.03	0.35	BGL 10	1	85	
Atopobium parvulum		4.60	0.10		99	99	
43035110	**Good identification**						**43035110**
Eubacterium saburreum		96.77	0.46	IND 99			
Fusobacterium mortiferum		3.24	0.23				
43035120	**Low discrimination**						**43035120**
					GRA		
Eubacterium saburreum		66.46	0.35	IND 99	85		
Actinomyces meyeri		14.27	0.16		99		
Fusobacterium mortiferum		10.30	0.23		1		
43040000	**Low discrimination**						**43040000**
					SPO		
Clostridium botulinum biovar A		78.45	0.47	SUC 7	99		
Atopobium minutum		8.37	0.38	CEL 1	1		
Clostridium sporogenes		7.74	0.27		99		
43040040	**Good identification**						**43040040**
					COC	GRA	
Clostridium botulinum biovar A		98.36	0.38	SUC 7	1	99	
Clostridium novyi biovar A		1.06	0.16	CEL 1	1	99	
Megasphaera elsdenii		0.58	0.04		99	1	
43040100	**Low discrimination**						**43040100**
					GRA	SPO	
Fusobacterium mortiferum		87.24	0.40	FRU 99	1	1	
Clostridium botulinum biovar A		12.76	0.13		99	99	
43041000	**Good identification**						**43041000**
					GRA		
Eubacterium tortuosum		96.36	0.50	SAL 99	99		
Fusobacterium mortiferum		1.80	0.37	CEL 10	1		
Atopobium minutum		1.44	0.30		99		
43041002	**Acceptable identification**						**43041002**
					GRA		
Eubacterium tortuosum		93.63	0.37	SAL 99	99		
Atopobium minutum		5.27	0.27	CEL 10	99		
Fusobacterium mortiferum		1.10	0.20		1		
43041020	**Good identification**						**43041020**
Eubacterium tortuosum		97.26	0.37	SAL 99			
Atopobium minutum		2.74	0.21	CEL 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43041100	**Low discrimination**						**43041100**
					GRA		
Fusobacterium mortiferum		78.49	0.43	FRU 99	1		
Eubacterium tortuosum		18.24	0.15		99		
Eubacterium saburreum		1.82	0.13		85		
43050000	**Good identification**						**43050000**
					GRA	SPO	
Fusobacterium mortiferum		91.35	0.68		1	1	
Atopobium minutum		3.07	0.37		99	1	
Clostridium botulinum biovar A		1.99	0.26		99	99	
43050002	**Low discrimination**						**43050002**
					GRA	SPO	
Fusobacterium mortiferum		80.87	0.51	BGL 10	1	1	
Atopobium minutum		16.30	0.34		99	1	
Clostridium sporogenes		1.44	0.06		99	99	
43050100	**Very good identification**						**43050100**
					GRA		
Fusobacterium mortiferum		99.65	0.74		1		
Actinomyces meyeri		0.36	0.23		99		
43050101	**Good identification**						**43050101**
Fusobacterium mortiferum		100.0	0.40	URE 1			
43050102	**Very good identification**						**43050102**
Fusobacterium mortiferum		100.0	0.58	BGL 10			
43050110	**Good identification**						**43050110**
Fusobacterium mortiferum		100.0	0.40	MLZ 1			
43050120	**Good identification**						**43050120**
					GRA		
Fusobacterium mortiferum		95.41	0.40	NAG 1	1		
Actinomyces meyeri		4.59	0.08		99		
43050140	**Good identification**						**43050140**
Fusobacterium mortiferum		100.0	0.40	SOR 1			
43050300	**Good identification**						**43050300**
Fusobacterium mortiferum		100.0	0.40	RHA 1			
43050500	**Acceptable identification**						**43050500**
					GRA		
Fusobacterium mortiferum		91.89	0.40	ARA 1	1		
Actinomyces meyeri		8.11	0.13		99		
43051000	**Very good identification**						**43051000**
					GRA		
Fusobacterium mortiferum		98.08	0.71		1		
Atopobium minutum		0.73	0.29		99		
Eubacterium tortuosum		0.54	0.15		99		
43051001	**Good identification**						**43051001**
Fusobacterium mortiferum		100.0	0.37	URE 1			
43051002	**Acceptable identification**						**43051002**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		91.69	0.54	BGL 10	1		
Atopobium minutum		4.11	0.26		99		
Lactobacillus catenaformis		1.21	0.12		99		
43051010	**Low discrimination**						**43051010**
					GRA		
Fusobacterium mortiferum		89.53	0.37	MLZ 1	1		
Lactobacillus jensenii		5.38	0.06		99		
Eubacterium saburreum		5.08	0.14		85		
43051020	**Low discrimination**						**43051020**
					GRA		
Fusobacterium mortiferum		66.06	0.37	NAG 1	1		
Atopobium minutum		16.27	0.21		99		
Eubacterium tortuosum		6.32	0.03		99		
43051040	**Good identification**						**43051040**
Fusobacterium mortiferum		100.0	0.37	SOR 1			
43051100	**Very good identification**						**43051100**
					GRA		
Fusobacterium mortiferum		99.34	0.77		1		
Actinomyces meyeri		0.35	0.26		99		
Lactobacillus catenaformis		0.09	0.15		99		
43051101	**Acceptable identification**						**43051101**
					GRA	SPO	
Fusobacterium mortiferum		92.88	0.43	URE 1	1	1	
Clostridium chauvoei		7.12	0.12		99	99	
43051102	**Very good identification**						**43051102**
					GRA	SPO	
Fusobacterium mortiferum		97.53	0.61	BGL 10	1	1	
Clostridium barati		1.03	0.12		99	99	
Lactobacillus catenaformis		0.83	0.15		99	1	
43051110	**Acceptable identification**						**43051110**
Fusobacterium mortiferum		94.63	0.43	MLZ 1			
Eubacterium saburreum		5.37	0.20				
43051120	**Low discrimination**						**43051120**
					GRA	SPO	
Fusobacterium mortiferum		68.08	0.43	NAG 1	1	1	
Prevotella melaninogenica		14.60	0.14		1	1	
Clostridium barati		7.90	0.12		99	99	
43051140	**Good identification**						**43051140**
Fusobacterium mortiferum		100.0	0.43	SOR 1			
43051200	**Good identification**						**43051200**
Fusobacterium mortiferum		100.0	0.37	RHA 1			
43051300	**Good identification**						**43051300**
Fusobacterium mortiferum		100.0	0.43	RHA 1			
43051400	**Acceptable identification**						**43051400**
					GRA		
Fusobacterium mortiferum		90.57	0.37	ARA 1	1		
Lactobacillus fermentum		4.77	0.16		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		4.66	0.05		99		
43051500	**Low discrimination**						**43051500**
					GRA		
Fusobacterium mortiferum		88.88	0.43	ARA 1	1		
Actinomyces meyeri		7.84	0.16		99		
Lactobacillus fermentum		3.28	0.20		99		
43052010	**Good identification**						**43052010**
					SPO		
Clostridium difficile		99.68	0.39	SUC 1	99		
Eubacterium limosum		0.32	-.06	CEL 5	1		
43052100	**Good identification**						**43052100**
					GRA		
Fusobacterium mortiferum		97.67	0.40	MAN 1	1		
Eubacterium limosum		2.34	-.06		99		
43053000	**Low discrimination**						**43053000**
					GRA	SPO	
Fusobacterium mortiferum		56.22	0.37	MAN 1	1	1	
Clostridium innocuum		41.69	0.22		99	99	
Eubacterium limosum		2.09	-.06		99	1	
43053100	**Good identification**						**43053100**
Fusobacterium mortiferum		100.0	0.43	MAN 1			
43054000	**Acceptable identification**						**43054000**
					GRA	SPO	
Fusobacterium mortiferum		91.93	0.49	XYL 7	1	1	
Actinomyces meyeri		4.93	0.18		99	1	
Clostridium glycolicum		3.14	0.09		99	99	
43054100	**Acceptable identification**						**43054100**
					GRA		
Fusobacterium mortiferum		91.58	0.55	XYL 7	1		
Actinomyces meyeri		8.42	0.28		99		
43054102	**Good identification**						**43054102**
					GRA		
Fusobacterium mortiferum		97.51	0.39	XYL 7	1		
Actinomyces meyeri		2.50	0.02	BGL 10	99		
43055000	**Acceptable identification**						**43055000**
					GRA		
Fusobacterium mortiferum		94.09	0.52	XYL 7	1		
Actinomyces meyeri		5.05	0.21		99		
Eubacterium saburreum		0.47	0.11		85		
43055002	**Good identification**						**43055002**
					GRA		
Fusobacterium mortiferum		96.40	0.35	XYL 7	1		
Lactobacillus fermentum		3.60	0.12	BGL 10	99		
43055100	**Good identification**						**43055100**
					GRA		
Fusobacterium mortiferum		90.92	0.58	XYL 7	1		
Actinomyces meyeri		8.36	0.31		99		
Eubacterium saburreum		0.46	0.17		85		
43055102	**Good identification**						**43055102**
					GRA		
Fusobacterium mortiferum		95.08	0.42	XYL 7	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		2.49	0.16	BGL 10	99		
Actinomyces meyeri		2.43	0.05		99		
43060000	**Low discrimination**						**43060000**
					GRA	SPO	
Clostridium botulinum biovar A		85.69	0.36	SUC 7	99	99	
Clostridium sporogenes		11.28	0.19	CEL 1	99	99	
Fusobacterium mortiferum		3.03	0.25		1	1	
43070000	**Very good identification**						**43070000**
					GRA	SPO	
Fusobacterium mortiferum		97.58	0.59		1	1	
Clostridium botulinum biovar A		1.78	0.16		99	99	
Clostridium sporogenes		0.65	0.06		99	99	
43070002	**Good identification**						**43070002**
					GRA	SPO	
Fusobacterium mortiferum		98.05	0.42	BGL 10	1	1	
Clostridium sporogenes		1.95	-.02		99	99	
43070100	**Very good identification**						**43070100**
Fusobacterium mortiferum		100.0	0.65				
43070102	**Good identification**						**43070102**
Fusobacterium mortiferum		100.0	0.49	BGL 10			
43071000	**Very good identification**						**43071000**
					GRA		
Fusobacterium mortiferum		98.05	0.62		1		
Lactobacillus jensenii		1.79	0.22		99		
Eubacterium saburreum		0.15	0.12		85		
43071002	**Acceptable identification**						**43071002**
					GRA		
Fusobacterium mortiferum		90.11	0.45	BGL 10	1		
Lactobacillus jensenii		9.89	0.19		99		
43071100	**Very good identification**						**43071100**
					COC	GRA	
Fusobacterium mortiferum		98.45	0.68		1	1	
Atopobium parvulum		1.39	0.25		99	99	
Eubacterium saburreum		0.15	0.18		1	85	
43071102	**Good identification**						**43071102**
					COC	GRA	
Fusobacterium mortiferum		96.82	0.52	BGL 10	1	1	
Atopobium parvulum		2.18	0.12		99	99	
Lactobacillus acidophilus		1.01	0.05		1	99	
43074000	**Good identification**						**43074000**
Fusobacterium mortiferum		100.0	0.40	XYL 7			
43074100	**Good identification**						**43074100**
Fusobacterium mortiferum		100.0	0.46	XYL 7			
43075000	**Good identification**						**43075000**
Fusobacterium mortiferum		100.0	0.43	XYL 7			
43075100	**Low discrimination**						**43075100**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		73.08	0.49	XYL 7	1	1	
Atopobium parvulum		25.52	0.30		99	99	
Eubacterium saburreum		0.82	0.14		1	85	
43100000	**Low discrimination**						**43100000**
Clostridium botulinum biovar A		79.55	0.79	SUC 7			
Clostridium sporogenes		17.82	0.66				
Clostridium bifermentans		1.28	0.48				
43100001	**Low discrimination**						**43100001**
Clostridium botulinum biovar A		53.93	0.45	SUC 7			
Clostridium novyi biovar A		31.61	0.53	URE 1			
Clostridium sporogenes		12.08	0.32				
43100002	**Low discrimination**						**43100002**
Clostridium sporogenes		85.42	0.58	SUC 1			
Clostridium botulinum biovar A		11.56	0.45				
Clostridium novyi biovar A		1.20	0.40				
43100003	**Low discrimination**						**43100003**
Clostridium novyi biovar A		54.96	0.40	SUC 1			
Clostridium sporogenes		39.67	0.24				
Clostridium botulinum biovar A		5.37	0.11				
43100010	**Low discrimination**						**43100010**
Clostridium botulinum biovar A		54.43	0.45	SUC 7			
Clostridium novyi biovar A		31.90	0.53	MLZ 1			
Clostridium sporogenes		12.19	0.32				
43100011	**Good identification**						**43100011**
Clostridium novyi biovar A		97.93	0.53	SUC 1			
Clostridium botulinum biovar A		1.69	0.11				
Clostridium sporogenes		0.38	-.03				
43100012	**Low discrimination**						**43100012**
Clostridium novyi biovar A		54.96	0.40	SUC 1			
Clostridium sporogenes		39.67	0.24				
Clostridium botulinum biovar A		5.37	0.11				
43100013	**Good identification**						**43100013**
Clostridium novyi biovar A		100.0	0.40	SUC 1			
43100020	**Low discrimination**						**43100020**
Clostridium botulinum biovar A		64.91	0.45	SUC 7			
Clostridium sporogenes		14.54	0.32	NAG 1			
Clostridium novyi biovar A		9.51	0.43				
43100021	**Acceptable identification**						**43100021**
Clostridium novyi biovar A		92.22	0.43	SUC 1			
Clostridium botulinum biovar A		6.36	0.11				
Clostridium sporogenes		1.42	-.03				
43100030	**Acceptable identification**						**43100030**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		92.22	0.43	SUC 1			
Clostridium botulinum biovar A		6.36	0.11				
Clostridium sporogenes		1.42	-.03				
43100031	**Good identification**						**43100031**
Clostridium novyi biovar A		100.0	0.43	SUC 1			
43100040	**Very good identification**						**43100040**
					COC	GRA	
Clostridium botulinum biovar A		96.31	0.70	SUC 7	1	99	
Clostridium novyi biovar A		1.27	0.50		1	99	
Megasphaera elsdenii		1.04	0.41		99	1	
43100041	**Low discrimination**						**43100041**
					COC	GRA	
Clostridium novyi biovar A		56.23	0.50	SUC 1	1	99	
Clostridium botulinum biovar A		42.98	0.36		1	99	
Megasphaera elsdenii		0.47	0.07		99	1	
43100042	**Low discrimination**						**43100042**
Clostridium botulinum biovar A		62.51	0.36	SUC 7			
Clostridium sporogenes		15.63	0.24	BGL 1			
Clostridium novyi biovar A		14.43	0.37				
43100043	**Good identification**						**43100043**
Clostridium novyi biovar A		95.81	0.37	SUC 1			
Clostridium botulinum biovar A		4.19	0.02				
43100050	**Low discrimination**						**43100050**
Clostridium novyi biovar A		55.44	0.50	SUC 1			
Clostridium botulinum biovar A		42.38	0.36				
Clostridium botulinum biovar B		1.39	0.08				
43100051	**Good identification**						**43100051**
Clostridium novyi biovar A		99.23	0.50	SUC 1			
Clostridium botulinum biovar A		0.77	0.02				
43100052	**Good identification**						**43100052**
Clostridium novyi biovar A		95.81	0.37	SUC 1			
Clostridium botulinum biovar A		4.19	0.02				
43100053	**Good identification**						**43100053**
Clostridium novyi biovar A		100.0	0.37	SUC 1			
43100060	**Low discrimination**						**43100060**
Clostridium botulinum biovar A		71.81	0.36	SUC 7			
Clostridium novyi biovar A		23.49	0.40	NAG 1			
Clostridium botulinum biovar B		2.36	0.08				
43100061	**Good identification**						**43100061**
Clostridium novyi biovar A		97.00	0.40	SUC 1			
Clostridium botulinum biovar A		3.00	0.02				
43100070	**Acceptable identification**						**43100070**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		92.56	0.40	SUC 1			
Clostridium botulinum biovar B		4.59	0.03				
Clostridium botulinum biovar A		2.86	0.02				
43100071	**Good identification**						**43100071**
Clostridium novyi biovar A		100.0	0.40	SUC 1			
43100100	**Low discrimination**						**43100100**
					SPO		
Clostridium botulinum biovar A		67.65	0.45	SUC 7	99		
Clostridium sporogenes		15.16	0.32	LAC 1	99		
Actinomyces meyeri		11.85	0.36		1		
43100101	**Good identification**						**43100101**
Clostridium chauvoei		95.36	0.40	MNS 90			
Clostridium botulinum biovar A		2.34	0.11	GAL 99			
Clostridium novyi biovar A		1.37	0.19				
43100140	**Good identification**						**43100140**
					COC	GRA	
Clostridium botulinum biovar A		96.38	0.36	SUC 7	1	99	
Clostridium novyi biovar A		1.27	0.16	LAC 1	1	99	
Megasphaera elsdenii		1.04	0.07		99	1	
43100200	**Low discrimination**						**43100200**
Clostridium botulinum biovar A		79.64	0.45	SUC 7			
Clostridium sporogenes		17.84	0.32	RHA 1			
Clostridium bifermentans		1.28	0.14				
43100240	**Good identification**						**43100240**
					COC	GRA	
Clostridium botulinum biovar A		96.94	0.36	SUC 7	1	99	
Clostridium novyi biovar A		1.28	0.16	RHA 1	1	99	
Megasphaera elsdenii		1.05	0.07		99	1	
43100400	**Low discrimination**						**43100400**
Clostridium botulinum biovar A		78.49	0.45	SUC 7			
Clostridium sporogenes		17.58	0.32	ARA 1			
Clostridium bifermentans		1.26	0.14				
43100440	**Good identification**						**43100440**
					COC	GRA	
Clostridium botulinum biovar A		96.94	0.36	SUC 7	1	99	
Clostridium novyi biovar A		1.28	0.16	ARA 1	1	99	
Megasphaera elsdenii		1.05	0.07		99	1	
43101000	**Low discrimination**						**43101000**
					SPO		
Clostridium botulinum biovar A		44.09	0.45	SUC 7	99		
Eubacterium saburreum		21.92	0.51	GAL 1	1		
Eubacterium tortuosum		11.68	0.32		1		
43101001	**Low discrimination**						**43101001**
Clostridium chauvoei		82.38	0.40	MNS 90			
Clostridium novyi biovar A		13.02	0.37	LAC 99			
Clostridium botulinum biovar A		2.02	0.11				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43101002	**Low discrimination**						**43101002**
					SPO		
Eubacterium saburreum		32.19	0.39	IND 99	1		
Clostridium sporogenes		27.40	0.24		99		
Eubacterium tortuosum		17.15	0.19		1		
43101010	**Low discrimination**						**43101010**
					SPO		
Eubacterium saburreum		83.54	0.50	IND 99	1		
Clostridium novyi biovar A		13.38	0.37		99		
Clostridium botulinum biovar A		2.07	0.11		99		
43101011	**Low discrimination**						**43101011**
					SPO		
Clostridium novyi biovar A		88.73	0.37	SUC 1	99		
Clostridium chauvoei		5.67	0.06	GAL 10	99		
Eubacterium saburreum		5.60	0.16		1		
43101012	**Low discrimination**						**43101012**
					SPO		
Eubacterium saburreum		86.20	0.37	IND 99	1		
Clostridium novyi biovar A		13.80	0.24		99		
43101020	**Low discrimination**						**43101020**
					GRA		
Eubacterium saburreum		39.13	0.39	IND 99	85		
Eubacterium tortuosum		20.85	0.19		99		
Prevotella melaninogenica		12.11	0.14		1		
43101030	**Low discrimination**						**43101030**
					SPO		
Eubacterium saburreum		81.51	0.37	IND 99	1		
Clostridium novyi biovar A		18.49	0.26		99		
43101040	**Low discrimination**						**43101040**
Clostridium botulinum biovar A		81.28	0.36	SUC 7			
Clostridium novyi biovar A		11.81	0.34	GAL 1			
Clostridium glycolicum		3.32	0.11				
43101100	**Low discrimination**						**43101100**
					COC	SPO	
Eubacterium saburreum		52.05	0.58	IND 99	1	1	
Clostridium chauvoei		18.49	0.40		1	99	
Blautia hansenii		12.97	0.32		99	1	
43101101	**Very good identification**						**43101101**
					SPO		
Clostridium chauvoei		99.96	0.74	MNS 90	99		
Eubacterium saburreum		0.03	0.24		1		
Actinomyces meyeri		0.01	0.05		1		
43101102	**Low discrimination**						**43101102**
					COC	SPO	
Eubacterium saburreum		57.12	0.45	IND 99	1	1	
Clostridium chauvoei		20.29	0.27		1	99	
Blautia hansenii		8.96	0.15		99	1	
43101103	**Very good identification**						**43101103**
					SPO		
Clostridium chauvoei		99.97	0.62	MNS 90	99		
Eubacterium saburreum		0.03	0.11		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43101110	**Very good identification**						**43101110**
					COC	SPO	
Eubacterium saburreum		98.99	0.56	IND 99	1	1	
Clostridium chauvoei		0.43	0.06		1	99	
Blautia hansenii		0.30	-.03		99	1	
43101111	**Good identification**						**43101111**
					SPO		
Clostridium chauvoei		97.73	0.40	MNS 90	99		
Eubacterium saburreum		2.27	0.22	MLZ 1	1		
43101112	**Good identification**						**43101112**
					CAT		
Eubacterium saburreum		98.93	0.43	IND 99	1		
Capnocytophaga ochracea		1.07	-.03		90		
43101120	**Low discrimination**						**43101120**
					SPO	GRA	
Prevotella melaninogenica		85.74	0.48	MNS 90	1	1	
Eubacterium saburreum		6.53	0.45	RAF 90	1	85	
Clostridium chauvoei		4.38	0.32	FRU 90	99	99	
43101121	**Very good identification**						**43101121**
					SPO	GRA	
Clostridium chauvoei		99.79	0.66	MNS 90	99	99	
Prevotella melaninogenica		0.20	0.14		1	1	
Eubacterium saburreum		0.02	0.11		1	85	
43101122	**Low discrimination**						**43101122**
					CAT		
Prevotella melaninogenica		69.43	0.36	MNS 90	1		
Capnocytophaga ochracea		19.90	0.30	RAF 90	90		
Eubacterium saburreum		4.90	0.32	FRU 90	1		
43101123	**Good identification**						**43101123**
					GRA	SPO	
Clostridium chauvoei		99.79	0.53	MNS 90	99	99	
Prevotella melaninogenica		0.22	0.02		1	1	
43101130	**Low discrimination**						**43101130**
					CAT		
Eubacterium saburreum		85.26	0.43	IND 99	1		
Prevotella melaninogenica		13.82	0.14		1		
Capnocytophaga ochracea		0.92	-.03		90		
43101141	**Good identification**						**43101141**
Clostridium chauvoei		100.0	0.40	MNS 90			
43101301	**Good identification**						**43101301**
Clostridium chauvoei		100.0	0.40	MNS 90			
43101501	**Good identification**						**43101501**
Clostridium chauvoei		100.0	0.40	MNS 90			
43101530	**Good identification**						**43101530**
Bifidobacterium longum subsp. longum		98.32	0.36	MNS 87			
Eubacterium saburreum		1.68	0.09	RAF 97			
43101532	**Good identification**						**43101532**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		97.18	0.36	MNS 87	99	1	
Capnocytophaga ochracea		2.82	0.01	RAF 97	1	90	
43102000	**Low discrimination**						**43102000**
					COC	GRA	
Clostridium botulinum biovar A		57.80	0.45	SUC 7	1	99	
Megasphaera elsdenii		27.74	0.44	MAN 1	99	1	
Clostridium sporogenes		12.95	0.32		1	99	
43102040	**Low discrimination**						**43102040**
					COC	GRA	
Megasphaera elsdenii		51.21	0.41	ESL 1	99	1	
Clostridium botulinum biovar A		47.80	0.36	FRU 95	1	99	
Clostridium novyi biovar A		0.63	0.16		1	99	
43103101	**Good identification**						**43103101**
Clostridium chauvoei		100.0	0.40	MNS 90			
43104000	**Low discrimination**						**43104000**
Clostridium botulinum biovar A		68.47	0.45	SUC 7			
Clostridium sporogenes		15.34	0.32	XYL 1			
Clostridium glycolicum		7.52	0.28				
43104040	**Low discrimination**						**43104040**
Clostridium glycolicum		87.13	0.50	SUC 1			
Clostridium botulinum biovar A		12.48	0.36	FRU 90			
Clostridium novyi biovar A		0.17	0.16				
43104100	**Acceptable identification**						**43104100**
					SPO		
Actinomyces meyeri		94.17	0.41	ESL 1	1		
Clostridium botulinum biovar A		2.80	0.11	FRU 95	99		
Eubacterium saburreum		1.75	0.19		1		
43105000	**Low discrimination**						**43105000**
					SPO		
Eubacterium saburreum		61.84	0.47	IND 99	1		
Actinomyces meyeri		29.44	0.34		1		
Clostridium glycolicum		2.82	0.11		99		
43105010	**Good identification**						**43105010**
Eubacterium saburreum		100.0	0.45	IND 99			
43105100	**Low discrimination**						**43105100**
					COC		
Eubacterium saburreum		51.99	0.53	IND 99	1		
Actinomyces meyeri		42.44	0.44		1		
Blautia hansenii		4.25	0.19		99		
43105101	**Good identification**						**43105101**
					SPO		
Clostridium chauvoei		97.30	0.40	MNS 90	99		
Eubacterium saburreum		1.49	0.19	XYL 1	1		
Actinomyces meyeri		1.22	0.10		1		
43105102	**Low discrimination**						**43105102**
Eubacterium saburreum		76.57	0.40	IND 99			
Actinomyces meyeri		10.95	0.18				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		5.71	0.24					
43105110	**Good identification**							**43105110**
Eubacterium saburreum		99.00	0.52	IND 99				
Actinomyces meyeri		1.00	0.10					
43105112	**Good identification**							**43105112**
Eubacterium saburreum		100.0	0.39	IND 99				
43105120	**Low discrimination**							**43105120**
					GRA			
Eubacterium saburreum		49.84	0.40	IND 99	85			
Actinomyces meyeri		31.44	0.29		99			
Prevotella melaninogenica		12.28	0.14		1			
43105130	**Good identification**							**43105130**
Eubacterium saburreum		96.78	0.39	IND 99				
Bifidobacterium longum subsp. longum		3.22	0.10					
43105520	**Low discrimination**							**43105520**
Bifidobacterium longum subsp. longum		89.66	0.40	MNS 87				
Actinomyces meyeri		8.93	0.19	RAF 97				
Lactobacillus fermentum		1.41	0.16	FRU 90				
43105522	**Acceptable identification**							**43105522**
					GRA	CAT		
Bifidobacterium longum subsp. longum		94.98	0.40	MNS 87	99	1		
Capnocytophaga ochracea		3.53	0.07	RAF 97	1	90		
Lactobacillus fermentum		1.49	0.16	FRU 90	99	1		
43105530	**Good identification**							**43105530**
Bifidobacterium longum subsp. longum		100.0	0.44	MNS 87				
43105532	**Good identification**							**43105532**
Bifidobacterium longum subsp. longum		100.0	0.44	MNS 87				
43110000	**Low discrimination**							**43110000**
					COC	GRA		
Megasphaera elsdenii		40.89	0.66	ESL 1	99	1		
Clostridium botulinum biovar A		28.33	0.59		1	99		
Clostridium sporogenes		17.55	0.53		1	99		
43110002	**Low discrimination**							**43110002**
					COC	GRA	SPO	
Clostridium sporogenes		48.90	0.45	SUC 1	1	99	99	
Megasphaera elsdenii		37.99	0.50		99	1	1	
Fusobacterium mortiferum		5.53	0.44		1	1	1	
43110011	**Good identification**							**43110011**
Clostridium novyi biovar A		100.0	0.37	SUC 1				
43110040	**Low discrimination**							**43110040**
					COC	GRA		
Megasphaera elsdenii		66.50	0.63	ESL 1	99	1		
Clostridium botulinum biovar A		20.64	0.50		1	99		
Clostridium glycolicum		10.81	0.44		1	99		
43110042	**Acceptable identification**							**43110042**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Megasphaera elsdenii		93.12	0.47	ESL 1	99	1		
Clostridium botulinum biovar A		2.63	0.16	BGL 10	1	99		
Clostridium sporogenes		1.82	0.11		1	99		
43110100	**Low discrimination**							**43110100**
					COC	GRA		
Actinomyces meyeri		49.71	0.58	ESL 1	1	99		
Fusobacterium mortiferum		46.31	0.67		1	1		
Megasphaera elsdenii		1.38	0.32		99	1		
43110101	**Acceptable identification**							**43110101**
					GRA	SPO		
Clostridium chauvoei		93.68	0.47	MNS 90	99	99		
Actinomyces meyeri		3.27	0.24	GAL 99	99	1		
Fusobacterium mortiferum		3.05	0.33		1	1		
43110102	**Low discrimination**							**43110102**
					GRA	CAT		
Fusobacterium mortiferum		66.77	0.50	MLT 11	1	5		
Actinomyces meyeri		19.95	0.32	BGL 10	99	1		
Capnocytophaga ochracea		3.16	0.14		1	90		
43110120	**Low discrimination**							**43110120**
					GRA			
Actinomyces meyeri		58.48	0.43	ESL 1	99			
Prevotella melaninogenica		31.50	0.30	NAG 12	1			
Fusobacterium mortiferum		4.04	0.33		1			
43110121	**Good identification**							**43110121**
					GRA	SPO		
Clostridium chauvoei		97.85	0.38	MNS 90	99	99		
Actinomyces meyeri		1.40	0.09	GAL 99	99	1		
Prevotella melaninogenica		0.75	-.04		1	1		
43110400	**Low discrimination**							**43110400**
					COC	SPO	GRA	
Actinomyces meyeri		47.49	0.37	ESL 1	1	1	99	
Megasphaera elsdenii		21.04	0.32		99	1	1	
Clostridium botulinum biovar A		14.58	0.25		1	99	99	
43110500	**Acceptable identification**							**43110500**
					GRA			
Actinomyces meyeri		94.83	0.48	ESL 1	99			
Fusobacterium mortiferum		3.57	0.33		1			
Lactobacillus fermentum		1.60	0.28		99			
43111000	**Low discrimination**							**43111000**
					GRA			
Fusobacterium mortiferum		32.68	0.63	MLT 11	1			
Eubacterium saburreum		24.46	0.60		85			
Actinomyces meyeri		20.46	0.51		99			
43111001	**Acceptable identification**							**43111001**
					SPO	GRA		
Clostridium chauvoei		93.36	0.47	MNS 90	99	99		
Fusobacterium mortiferum		1.95	0.29	LAC 99	1	1		
Eubacterium saburreum		1.46	0.25		1	85		
43111002	**Low discrimination**							**43111002**
					GRA			
Lactobacillus fermentum		58.31	0.59	ESL 1	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium saburreum		18.06	0.47		85			
Fusobacterium mortiferum		15.19	0.47		1			
43111010	**Very good identification**							**43111010**
					GRA			
Eubacterium saburreum		95.93	0.58	IND 99	85			
Fusobacterium mortiferum		1.58	0.29		1			
Actinomyces meyeri		0.99	0.16		99			
43111012	**Acceptable identification**							**43111012**
					GRA			
Eubacterium saburreum		93.57	0.45	IND 99	85			
Lactobacillus fermentum		3.73	0.24		99			
Fusobacterium mortiferum		0.97	0.13		1			
43111020	**Low discrimination**							**43111020**
					GRA			
Lactobacillus fermentum		22.80	0.50	ESL 1	99			
Eubacterium saburreum		21.19	0.47		85			
Prevotella melaninogenica		19.67	0.30		1			
43111021	**Good identification**							**43111021**
					SPO			
Clostridium chauvoei		95.37	0.38	MNS 90	99			
Lactobacillus fermentum		0.85	0.16	LAC 99	1			
Eubacterium saburreum		0.79	0.13		1			
43111022	**Low discrimination**							**43111022**
					GRA	CAT		
Lactobacillus fermentum		43.09	0.50	ESL 1	99	1		
Capnocytophaga ochracea		22.34	0.30		1	90		
Clostridium perfringens		15.44	0.31		99	1		
43111030	**Acceptable identification**							**43111030**
					GRA			
Eubacterium saburreum		94.37	0.45	IND 99	85			
Lactobacillus fermentum		1.25	0.16		99			
Prevotella melaninogenica		1.08	-.04		1			
43111100	**Low discrimination**							**43111100**
					GRA			
Actinomyces meyeri		31.22	0.61	ESL 1	99			
Fusobacterium mortiferum		29.09	0.70		1			
Eubacterium saburreum		21.77	0.66		85			
43111101	**Very good identification**							**43111101**
					GRA	SPO		
Clostridium chauvoei		99.84	0.81	MNS 90	99	99		
Actinomyces meyeri		0.05	0.27		99	1		
Fusobacterium mortiferum		0.05	0.36		1	1		
43111102	**Low discrimination**							**43111102**
					CAT	GRA		
Capnocytophaga ochracea		35.33	0.48	MNS 90	90	1		
Lactobacillus fermentum		30.33	0.62	RAF 90	1	99		
Eubacterium saburreum		13.44	0.53	NAG 90	1	85		
43111103	**Very good identification**							**43111103**
					GRA	SPO	CAT	
Clostridium chauvoei		99.72	0.68	MNS 90	99	99	1	
Capnocytophaga ochracea		0.10	0.14		1	1	90	
Lactobacillus fermentum		0.08	0.28		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43111110	**Very good identification**							**43111110**
					GRA			
Eubacterium saburreum		95.13	0.64	IND 99	85			
Actinomyces meyeri		1.68	0.27		99			
Fusobacterium mortiferum		1.57	0.36		1			
43111111	**Good identification**							**43111111**
					SPO			
Clostridium chauvoei		97.10	0.47	MNS 90	99			
Eubacterium saburreum		2.90	0.30	MLZ 1	1			
43111112	**Low discrimination**							**43111112**
					CAT	GRA		
Eubacterium saburreum		70.15	0.52	IND 99	1	85		
Capnocytophaga ochracea		25.05	0.32		90	1		
Lactobacillus fermentum		1.96	0.28		1	99		
43111120	**Low discrimination**							**43111120**
					GRA	CAT		
Prevotella melaninogenica		81.87	0.64	MNS 90	1	1		
Capnocytophaga ochracea		5.46	0.48	RAF 90	1	90		
Clostridium perfringens		5.35	0.51		99	1		
43111121	**Very good identification**							**43111121**
					GRA	SPO		
Clostridium chauvoei		99.06	0.73	MNS 90	99	99		
Prevotella melaninogenica		0.76	0.30		1	1		
Clostridium perfringens		0.05	0.17		99	84		
43111122	**Low discrimination**							**43111122**
					GRA	CAT		
Capnocytophaga ochracea		69.77	0.64	MNS 90	1	90		
Prevotella melaninogenica		22.12	0.52	RAF 90	1	1		
Clostridium perfringens		3.57	0.45		99	1		
43111123	**Very good identification**							**43111123**
					GRA	CAT	SPO	
Clostridium chauvoei		96.33	0.60	MNS 90	99	1	99	
Capnocytophaga ochracea		2.53	0.30		1	90	1	
Prevotella melaninogenica		0.80	0.18		1	1	1	
43111130	**Low discrimination**							**43111130**
					CAT			
Eubacterium saburreum		52.02	0.52	IND 99	1			
Prevotella melaninogenica		25.30	0.30		1			
Capnocytophaga ochracea		18.57	0.32		90			
43111131	**Good identification**							**43111131**
					SPO	GRA		
Clostridium chauvoei		97.16	0.38	MNS 90	99	99		
Eubacterium saburreum		1.54	0.17	MLZ 1	1	85		
Prevotella melaninogenica		0.75	-.04		1	1		
43111132	**Acceptable identification**							**43111132**
					CAT			
Capnocytophaga ochracea		91.04	0.48	MNS 90	90			
Eubacterium saburreum		5.00	0.39	RAF 90	1			
Prevotella melaninogenica		2.62	0.18	MLZ 10	1			
43111141	**Good identification**							**43111141**
Clostridium chauvoei		100.0	0.47	MNS 90				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43111160	**Low discrimination**						**43111160**
					GRA	CAT	
Clostridium perfringens		40.38	0.39	MNS 99	99	1	
Prevotella melaninogenica		32.76	0.30		1	1	
Capnocytophaga ochracea		24.05	0.32		1	90	
43111161	**Good identification**						**43111161**
					GRA	SPO	
Clostridium chauvoei		97.77	0.38	MNS 90	99	99	
Clostridium perfringens		0.93	0.04	SOR 1	99	84	
Prevotella melaninogenica		0.75	-.04		1	1	
43111162	**Low discrimination**						**43111162**
					GRA	CAT	
Capnocytophaga ochracea		89.20	0.48	MNS 90	1	90	
Clostridium perfringens		7.83	0.33	RAF 90	99	1	
Prevotella melaninogenica		2.57	0.18	SOR 10	1	1	
43111301	**Good identification**						**43111301**
Clostridium chauvoei		100.0	0.47	MNS 90			
43111321	**Good identification**						**43111321**
					GRA	SPO	
Clostridium chauvoei		99.24	0.38	MNS 90	99	99	
Prevotella melaninogenica		0.76	-.04	RHA 1	1	1	
43111400	**Low discrimination**						**43111400**
					GRA		
Lactobacillus fermentum		70.61	0.59	ESL 1	99		
Actinomyces meyeri		25.92	0.40		99		
Fusobacterium mortiferum		1.67	0.29		1		
43111402	**Very good identification**						**43111402**
					GRA	CAT	
Lactobacillus fermentum		97.99	0.59	ESL 1	99	1	
Actinomyces meyeri		1.11	0.15		99	1	
Capnocytophaga ochracea		0.33	0.01		1	90	
43111420	**Low discrimination**						**43111420**
Lactobacillus fermentum		83.00	0.50	ESL 1			
Actinomyces meyeri		12.46	0.26				
Bifidobacterium longum subsp. longum		1.55	0.14				
43111422	**Acceptable identification**						**43111422**
					GRA	CAT	
Lactobacillus fermentum		90.08	0.50	ESL 1	99	1	
Capnocytophaga ochracea		8.24	0.17		1	90	
Bifidobacterium longum subsp. longum		1.68	0.14		99	1	
43111500	**Low discrimination**						**43111500**
Lactobacillus fermentum		47.96	0.62	ESL 1			
Actinomyces meyeri		43.16	0.51				
Bifidobacterium longum subsp. longum		4.53	0.38				
43111501	**Good identification**						**43111501**
					SPO		
Clostridium chauvoei		97.31	0.47	MNS 90	99		
Lactobacillus fermentum		1.42	0.28	ARA 1	1		
Actinomyces meyeri		1.28	0.16		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43111502	**Low discrimination**							**43111502**
					GRA	CAT		
Lactobacillus fermentum		72.38	0.62	ESL 1	99	1		
Capnocytophaga ochracea		14.88	0.35		1	90		
Bifidobacterium longum subsp. longum		6.83	0.38		99	1		
43111510	**Low discrimination**							**43111510**
Bifidobacterium longum subsp. longum		79.10	0.42	MNS 87				
Eubacterium saburreum		10.21	0.30	RAF 97				
Lactobacillus fermentum		4.97	0.28					
43111512	**Low discrimination**							**43111512**
					GRA	CAT		
Bifidobacterium longum subsp. longum		63.93	0.42	MNS 87	99	1		
Bifidobacterium adolescentis		21.52	0.35	RAF 97	99	1		
Capnocytophaga ochracea		9.08	0.19		1	90		
43111520	**Low discrimination**							**43111520**
					GRA	CAT		
Bifidobacterium longum subsp. longum		30.56	0.48	MNS 87	99	1		
Lactobacillus fermentum		26.97	0.54	RAF 97	99	1		
Capnocytophaga ochracea		16.63	0.35		1	90		
43111521	**Acceptable identification**							**43111521**
					SPO			
Clostridium chauvoei		94.96	0.38	MNS 90	99			
Bifidobacterium longum subsp. longum		1.57	0.14	ARA 1	1			
Lactobacillus fermentum		1.38	0.20		1			
43111522	**Low discrimination**							**43111522**
					GRA	CAT		
Capnocytophaga ochracea		71.00	0.51	MNS 90	1	90		
Bifidobacterium longum subsp. longum		14.49	0.48	RAF 90	99	1		
Lactobacillus fermentum		12.79	0.54		99	1		
43111530	**Good identification**							**43111530**
					CAT	GRA		
Bifidobacterium longum subsp. longum		95.13	0.52	MNS 87	1	99		
Capnocytophaga ochracea		3.38	0.19	RAF 97	90	1		
Eubacterium saburreum		0.54	0.17		1	85		
43111532	**Low discrimination**							**43111532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		75.48	0.52	MNS 87	99	1		
Capnocytophaga ochracea		24.13	0.35	RAF 97	1	90		
Lactobacillus fermentum		0.40	0.20		99	1		
43111562	**Good identification**							**43111562**
					GRA	CAT		
Capnocytophaga ochracea		96.62	0.35	MNS 90	1	90		
Bifidobacterium longum subsp. longum		1.79	0.14	RAF 90	99	1		
Lactobacillus fermentum		1.58	0.20	SOR 10	99	1		
43112000	**Very good identification**							**43112000**
					COC	SPO	GRA	
Megasphaera elsdenii		96.22	0.66	ESL 1	99	1	1	
Eubacterium limosum		2.11	0.29		1	1	99	
Clostridium botulinum biovar A		0.67	0.25		1	99	99	
43112002	**Good identification**							**43112002**
					COC	SPO	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		98.16	0.50	ESL 1	99	1	1	
Clostridium sporogenes		1.28	0.11	BGL 10	1	99	99	
Propionibacterium propionicum		0.37	0.18		1	1	99	
43112040	**Very good identification**							**43112040**
					COC	GRA		
Megasphaera elsdenii		99.45	0.63	ESL 1	99	1		
Clostridium botulinum biovar A		0.31	0.16		1	99		
Clostridium glycolicum		0.16	0.10		1	99		
43112042	**Good identification**							**43112042**
					COC	GRA		
Megasphaera elsdenii		99.44	0.47	ESL 1	99	1		
Propionibacterium propionicum		0.56	0.18	BGL 10	1	99		
43112100	**Low discrimination**							**43112100**
					COC	GRA		
Propionibacterium propionicum		61.25	0.49	NIT 86	1	99		
Megasphaera elsdenii		22.44	0.32	ESL 1	99	1		
Actinomyces meyeri		8.19	0.24		1	99		
43112102	**Good identification**							**43112102**
					COC	GRA		
Propionibacterium propionicum		96.49	0.52	NIT 86	1	99		
Megasphaera elsdenii		2.62	0.15	ESL 1	99	1		
Fusobacterium mortiferum		0.89	0.16		1	1		
43112103	**Good identification**							**43112103**
					SPO			
Propionibacterium propionicum		96.07	0.36	NIT 86	1			
Clostridium chauvoei		3.93	-.01	ESL 1	99			
43112140	**Low discrimination**							**43112140**
					COC	GRA		
Propionibacterium propionicum		80.37	0.49	NIT 86	1	99		
Megasphaera elsdenii		19.63	0.29	ESL 1	99	1		
43112142	**Good identification**							**43112142**
					COC	GRA		
Propionibacterium propionicum		98.22	0.52	NIT 86	1	99		
Megasphaera elsdenii		1.78	0.12	ESL 1	99	1		
43112143	**Good identification**							**43112143**
Propionibacterium propionicum		100.0	0.36	NIT 86				
43113100	**Low discrimination**							**43113100**
					GRA			
Propionibacterium propionicum		59.72	0.49	NIT 86	99			
Actinomyces meyeri		11.97	0.27	ESL 1	99			
Fusobacterium mortiferum		11.16	0.36		1			
43113101	**Good identification**							**43113101**
					SPO			
Clostridium chauvoei		96.18	0.47	MNS 90	99			
Propionibacterium propionicum		2.79	0.33	MAN 1	1			
Actinomyces naeslundii		1.03	0.36		1			
43113102	**Low discrimination**							**43113102**
					GRA	CAT		
Propionibacterium propionicum		86.06	0.52	NIT 86	99	1		
Capnocytophaga ochracea		3.72	0.14	ESL 1	1	90		
Lactobacillus fermentum		3.20	0.28		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43113121	**Good identification**						**43113121**
					GRA	SPO	
Clostridium chauvoei		96.18	0.38	MNS 90	99	99	
Actinomyces naeslundii		3.08	0.36	MAN 1	99	1	
Prevotella melaninogenica		0.74	-.04		1	1	
43113140	**Good identification**						**43113140**
Propionibacterium propionicum		100.0	0.49	NIT 86			
43113142	**Good identification**						**43113142**
					GRA	CAT	
Propionibacterium propionicum		99.52	0.52	NIT 86	99	1	
Capnocytophaga ochracea		0.48	-.03	ESL 1	1	90	
43113143	**Acceptable identification**						**43113143**
					SPO		
Propionibacterium propionicum		92.24	0.36	NIT 86	1		
Actinomyces naeslundii		3.99	0.23	ESL 1	1		
Clostridium chauvoei		3.77	-.01	URE 10	99		
43114000	**Low discrimination**						**43114000**
					GRA	SPO	
Actinomyces meyeri		55.08	0.53	ESL 1	99	1	
Clostridium glycolicum		33.71	0.44		99	99	
Fusobacterium mortiferum		3.41	0.41		1	1	
43114020	**Acceptable identification**						**43114020**
					SPO		
Actinomyces meyeri		94.97	0.38	ESL 1	1		
Clostridium glycolicum		4.31	0.10	NAG 12	99		
Lactobacillus fermentum		0.73	0.12		1		
43114040	**Very good identification**						**43114040**
					COC	GRA	
Clostridium glycolicum		99.49	0.66	SUC 1	1	99	
Megasphaera elsdenii		0.33	0.29		99	1	
Clostridium botulinum biovar A		0.10	0.16		1	99	
43114100	**Very good identification**						**43114100**
					GRA		
Actinomyces meyeri		95.95	0.63	ESL 1	99		
Fusobacterium mortiferum		3.47	0.47		1		
Eubacterium saburreum		0.28	0.27		85		
43114102	**Low discrimination**						**43114102**
					GRA		
Actinomyces meyeri		82.77	0.37	ESL 1	99		
Fusobacterium mortiferum		10.74	0.31	BGL 3	1		
Lactobacillus fermentum		3.41	0.24		99		
43114120	**Good identification**						**43114120**
					CAT	GRA	
Actinomyces meyeri		98.33	0.48	ESL 1	1	99	
Capnocytophaga ochracea		0.46	0.04	NAG 12	90	1	
Eubacterium saburreum		0.37	0.14		1	85	
43114400	**Good identification**						**43114400**
					SPO		
Actinomyces meyeri		96.41	0.42	ESL 1	1		
Clostridium glycolicum		2.38	0.10		99		
Lactobacillus fermentum		1.20	0.20		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
43114500	**Good identification**						**43114500**
Actinomyces meyeri		99.11	0.53	ESL 1			
Lactobacillus fermentum		0.51	0.24				
Bifidobacterium longum subsp. longum		0.24	0.12				
43114520	**Acceptable identification**						**43114520**
Actinomyces meyeri		92.21	0.38	ESL 1			
Bifidobacterium longum subsp. longum		6.64	0.22	NAG 12			
Lactobacillus fermentum		1.15	0.16				
43115000	**Low discrimination**						**43115000**
Actinomyces meyeri		60.61	0.56	ESL 1			
Eubacterium saburreum		20.10	0.55				
Lactobacillus fermentum		12.49	0.55				
43115002	**Low discrimination**						**43115002**
Lactobacillus fermentum		67.47	0.55	ESL 1			
Eubacterium saburreum		19.16	0.42				
Actinomyces meyeri		10.13	0.30				
43115010	**Good identification**						**43115010**
Eubacterium saburreum		95.70	0.53	IND 99			
Actinomyces meyeri		3.56	0.21				
Lactobacillus fermentum		0.73	0.20				
43115012	**Good identification**						**43115012**
Eubacterium saburreum		95.83	0.41	IND 99			
Lactobacillus fermentum		4.17	0.20				
43115020	**Low discrimination**						**43115020**
Actinomyces meyeri		51.04	0.41	ESL 1			
Lactobacillus fermentum		25.71	0.46	NAG 12			
Eubacterium saburreum		21.90	0.42				
43115022	**Low discrimination**						**43115022**
					CAT	GRA	
Lactobacillus fermentum		67.46	0.46	ESL 1	1	99	
Capnocytophaga ochracea		14.88	0.20		90	1	
Eubacterium saburreum		10.14	0.29		1	85	
43115030	**Good identification**						**43115030**
Eubacterium saburreum		95.85	0.41	IND 99			
Actinomyces meyeri		2.76	0.07				
Lactobacillus fermentum		1.39	0.12				
43115040	**Good identification**						**43115040**
					SPO		
Clostridium glycolicum		98.23	0.50	SUC 1	99		
Actinomyces meyeri		1.15	0.21	GAL 10	1		
Eubacterium saburreum		0.38	0.21		1		
43115100	**Low discrimination**						**43115100**
Actinomyces meyeri		74.60	0.66	ESL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium saburreum		14.43	0.61					
Lactobacillus fermentum		6.27	0.58					
43115101	**Low discrimination**							**43115101**
					SPO			
Clostridium chauvoei		88.21	0.47	MNS 90	99			
Actinomyces meyeri		8.98	0.32	XYL 1	1			
Eubacterium saburreum		1.74	0.27		1			
43115102	**Low discrimination**							**43115102**
					CAT	GRA		
Lactobacillus fermentum		40.53	0.58	ESL 1	1	99		
Capnocytophaga ochracea		20.10	0.38		90	1		
Eubacterium saburreum		16.46	0.48		1	85		
43115110	**Good identification**							**43115110**
Eubacterium saburreum		92.67	0.60	IND 99				
Actinomyces meyeri		5.91	0.32					
Lactobacillus fermentum		0.50	0.24					
43115112	**Low discrimination**							**43115112**
					CAT	GRA		
Eubacterium saburreum		66.24	0.47	IND 99	1	85		
Bifidobacterium adolescentis		18.37	0.35		1	99		
Capnocytophaga ochracea		10.98	0.21		90	1		
43115120	**Low discrimination**							**43115120**
					CAT	GRA		
Actinomyces meyeri		49.15	0.51	ESL 1	1	99		
Capnocytophaga ochracea		15.02	0.38	NAG 12	90	1		
Eubacterium saburreum		12.31	0.48		1	85		
43115121	**Acceptable identification**							**43115121**
					GRA	SPO	CAT	
Clostridium chauvoei		92.51	0.38	MNS 90	99	99	1	
Actinomyces meyeri		3.85	0.17	XYL 1	99	1	1	
Capnocytophaga ochracea		1.18	0.04		1	1	90	
43115122	**Low discrimination**							**43115122**
					GRA	CAT		
Capnocytophaga ochracea		76.46	0.54	MNS 90	1	90		
Bacteroides fragilis		13.14	0.32	RAF 90	1	99		
Lactobacillus fermentum		5.71	0.50		99	1		
43115130	**Low discrimination**							**43115130**
					CAT	GRA		
Eubacterium saburreum		74.95	0.47	IND 99	1	85		
Capnocytophaga ochracea		12.43	0.21		90	1		
Bifidobacterium longum subsp. longum		7.48	0.26		1	99		
43115132	**Low discrimination**							**43115132**
					CAT	GRA		
Capnocytophaga ochracea		82.39	0.38	MNS 90	90	1		
Eubacterium saburreum		9.75	0.34	RAF 90	1	85		
Bifidobacterium longum subsp. longum		5.51	0.26	MLZ 10	1	99		
43115162	**Good identification**							**43115162**
					GRA	CAT		
Capnocytophaga ochracea		97.81	0.38	MNS 90	1	90		
Bacteroides fragilis		1.53	-.03	RAF 90	1	99		
Lactobacillus fermentum		0.66	0.16	SOR 10	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43115300	**Very good identification**						**43115300**
Eubacterium contortum		99.14	0.66	SAL 99			
Actinomyces meyeri		0.64	0.32				
Eubacterium saburreum		0.12	0.27				
43115302	**Good identification**						**43115302**
					GRA	CAT	
Eubacterium contortum		99.24	0.53	SAL 99	99	1	
Lactobacillus fermentum		0.31	0.24		99	1	
Capnocytophaga ochracea		0.15	0.04		1	90	
43115320	**Good identification**						**43115320**
					GRA		
Eubacterium contortum		95.76	0.53	SAL 99	99		
Bacteroides vulgatus		3.31	0.21		1		
Actinomyces meyeri		0.48	0.17		99		
43115322	**Low discrimination**						**43115322**
					GRA	CAT	
Eubacterium contortum		86.58	0.40	SAL 99	99	1	
Capnocytophaga ochracea		6.73	0.20		1	90	
Parabacteroides distasonis		2.54	0.25		1	76	
43115400	**Low discrimination**						**43115400**
Actinomyces meyeri		54.22	0.45	ESL 1			
Lactobacillus fermentum		44.70	0.55				
Eubacterium saburreum		0.73	0.21				
43115402	**Acceptable identification**						**43115402**
Lactobacillus fermentum		94.90	0.55	ESL 1			
Actinomyces meyeri		3.56	0.19				
Bifidobacterium adolescentis		0.79	0.14				
43115420	**Low discrimination**						**43115420**
Lactobacillus fermentum		62.83	0.46	ESL 1			
Actinomyces meyeri		31.18	0.30				
Bifidobacterium longum subsp. longum		5.99	0.22				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43115422	**Low discrimination**						**43115422**
					GRA	CAT	
Lactobacillus fermentum		86.98	0.46	ESL 1	99	1	
Bifidobacterium longum subsp. longum		8.30	0.22		99	1	
Capnocytophaga ochracea		3.39	0.07		1	90	
43115500	**Low discrimination**						**43115500**
Actinomyces meyeri		63.86	0.56	ESL 1			
Lactobacillus fermentum		21.47	0.58				
Bifidobacterium longum subsp. longum		10.35	0.46				
43115502	**Low discrimination**						**43115502**
Lactobacillus fermentum		45.37	0.58	ESL 1			
Bifidobacterium adolescentis		22.89	0.48				
Bifidobacterium longum subsp. longum		21.88	0.46				
43115510	**Acceptable identification**						**43115510**
Bifidobacterium longum subsp. longum		92.62	0.50	MNS 87			
Actinomyces meyeri		3.39	0.21	RAF 97			
Eubacterium saburreum		2.15	0.26				
43115512	**Low discrimination**						**43115512**
Bifidobacterium longum subsp. longum		56.21	0.50	MNS 87			
Bifidobacterium adolescentis		42.21	0.49	RAF 97			
Lactobacillus fermentum		0.69	0.24				
43115520	**Low discrimination**						**43115520**
Bifidobacterium longum subsp. longum		65.23	0.56	MNS 87			
Actinomyces meyeri		13.72	0.41	RAF 97			
Lactobacillus fermentum		11.27	0.50				
43115522	**Low discrimination**						**43115522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		62.56	0.56	MNS 87	99	1	
Capnocytophaga ochracea		25.54	0.41	RAF 97	1	90	
Lactobacillus fermentum		10.81	0.50		99	1	
43115530	**Very good identification**						**43115530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.38	0.60	MNS 87	99	1	
Capnocytophaga ochracea		0.29	0.08	RAF 97	1	90	
Actinomyces meyeri		0.12	0.07		99	1	
43115532	**Very good identification**						**43115532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.13	0.60	MNS 87	99	1	
Capnocytophaga ochracea		2.59	0.25	RAF 97	1	90	
Bifidobacterium adolescentis		0.18	0.15		99	1	
43115700	**Very good identification**						**43115700**
					GRA		
Eubacterium contortum		99.61	0.64	SAL 99	99		
Actinomyces meyeri		0.20	0.21		99		
Bacteroides vulgatus		0.10	0.06		1		
43115702	**Good identification**						**43115702**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		99.26	0.52	SAL 99		
Lactobacillus fermentum		0.37	0.24			
Bifidobacterium adolescentis		0.19	0.14			
43115720	**Low discrimination**					**43115720**
					GRA	
Eubacterium contortum		63.64	0.52	SAL 99	99	
Bacteroides vulgatus		35.72	0.40		1	
Bifidobacterium longum subsp. longum		0.46	0.22		99	
43115722	**Low discrimination**					**43115722**
					GRA	
Eubacterium contortum		89.36	0.39	SAL 99	99	
Bifidobacterium longum subsp. longum		3.67	0.22		99	
Bacteroides vulgatus		2.87	0.06		1	
43117502	**Low discrimination**					**43117502**
Bifidobacterium adolescentis		88.64	0.46	MNS 87		
Actinomyces israelii		8.14	0.35	RAF 92		
Lactobacillus fermentum		2.17	0.24			
43117512	**Good identification**					**43117512**
Bifidobacterium adolescentis		97.31	0.48	MNS 87		
Bifidobacterium longum subsp. longum		1.60	0.16	RAF 92		
Actinomyces israelii		1.09	0.21			
43120000	**Low discrimination**					**43120000**
Clostridium botulinum biovar A		76.91	0.69	SUC 7		
Clostridium sporogenes		22.97	0.58			
Clostridium bifermentans		0.05	0.14			
43120002	**Acceptable identification**					**43120002**
Clostridium sporogenes		90.79	0.50	SUC 1		
Clostridium botulinum biovar A		9.21	0.35			
43120040	**Very good identification**					**43120040**
Clostridium botulinum biovar A		98.12	0.60	SUC 7		
Clostridium sporogenes		0.99	0.24			
Clostridium botulinum biovar B		0.79	0.21			
43121000	**Low discrimination**					**43121000**
					SPO	
Eubacterium saburreum		49.86	0.48	IND 99	1	
Clostridium botulinum biovar A		37.61	0.35		99	
Clostridium sporogenes		11.23	0.24		99	
43121002	**Low discrimination**					**43121002**
					SPO	
Eubacterium saburreum		68.09	0.36	IND 99	1	
Clostridium sporogenes		28.97	0.15		99	
Clostridium botulinum biovar A		2.94	0.01		99	
43121010	**Good identification**					**43121010**
					SPO	
Eubacterium saburreum		99.08	0.47	IND 99	1	
Clostridium botulinum biovar A		0.92	0.01		99	
43121020	**Low discrimination**					**43121020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Eubacterium saburreum		89.90	0.36	IND 99	1		
Clostridium perfringens		6.22	0.07		84		
Clostridium botulinum biovar A		3.88	0.01		99		
43121100	**Low discrimination**						**43121100**
					COC	SPO	
Eubacterium saburreum		89.51	0.55	IND 99	1	1	
Clostridium chauvoei		5.30	0.24		1	99	
Atopobium parvulum		2.63	0.19		99	1	
43121101	**Very good identification**						**43121101**
					SPO		
Clostridium chauvoei		99.83	0.58	MNS 90	99		
Eubacterium saburreum		0.17	0.21	TRE 10	1		
43121102	**Acceptable identification**						**43121102**
					COC	SPO	
Eubacterium saburreum		91.86	0.42	IND 99	1	1	
Clostridium chauvoei		5.44	0.11		1	99	
Atopobium parvulum		2.70	0.06		99	1	
43121103	**Good identification**						**43121103**
Clostridium chauvoei		100.0	0.45	MNS 90			
43121110	**Good identification**						**43121110**
Eubacterium saburreum		100.0	0.53	IND 99			
43121112	**Good identification**						**43121112**
Eubacterium saburreum		100.0	0.40	IND 99			
43121120	**Low discrimination**						**43121120**
					GRA		
Eubacterium saburreum		64.20	0.42	IND 99	85		
Clostridium perfringens		13.96	0.21		99		
Prevotella melaninogenica		12.77	0.14		1		
43121121	**Good identification**						**43121121**
Clostridium chauvoei		100.0	0.50	MNS 90			
43121123	**Good identification**						**43121123**
Clostridium chauvoei		100.0	0.37	MNS 90			
43121130	**Good identification**						**43121130**
Eubacterium saburreum		100.0	0.40	IND 99			
43125000	**Good identification**						**43125000**
					SPO		
Eubacterium saburreum		98.61	0.44	IND 99	1		
Clostridium botulinum biovar A		1.40	0.01		99		
43125010	**Good identification**						**43125010**
Eubacterium saburreum		100.0	0.42	IND 99			
43125100	**Low discrimination**						**43125100**
					COC		
Eubacterium saburreum		89.79	0.50	IND 99	1		
Atopobium parvulum		9.10	0.23		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		1.11	0.10		1		
43125102	**Acceptable identification**						**43125102**
					COC		
Eubacterium saburreum		90.80	0.37	IND 99	1		
Atopobium parvulum		9.20	0.10		99		
43125110	**Good identification**						**43125110**
Eubacterium saburreum		100.0	0.49	IND 99			
43125112	**Good identification**						**43125112**
Eubacterium saburreum		100.0	0.36	IND 99			
43125120	**Acceptable identification**						**43125120**
					COC		
Eubacterium saburreum		90.80	0.37	IND 99	1		
Atopobium parvulum		9.20	0.10		99		
43125130	**Good identification**						**43125130**
Eubacterium saburreum		100.0	0.36	IND 99			
43130000	**Low discrimination**						**43130000**
					GRA	SPO	
Clostridium botulinum biovar A		46.34	0.49	SUC 7	99	99	
Clostridium sporogenes		38.26	0.45	FRU 6	99	99	
Fusobacterium mortiferum		11.64	0.51		1	1	
43130002	**Low discrimination**						**43130002**
					GRA	SPO	
Clostridium sporogenes		85.14	0.37	SUC 1	99	99	
Fusobacterium mortiferum		8.63	0.35		1	1	
Clostridium botulinum biovar A		3.12	0.14		99	99	
43130040	**Low discrimination**						**43130040**
					COC	GRA	
Clostridium botulinum biovar A		58.99	0.40	SUC 7	1	99	
Clostridium botulinum biovar B		29.93	0.32	FRU 6	1	99	
Megasphaera elsdenii		7.68	0.29		99	1	
43130100	**Low discrimination**						**43130100**
					GRA		
Fusobacterium mortiferum		89.30	0.58	MLT 11	1		
Actinomyces meyeri		3.24	0.24		99		
Eubacterium saburreum		2.26	0.29		85		
43130102	**Low discrimination**						**43130102**
					SPO	GRA	
Fusobacterium mortiferum		86.79	0.41	MLT 11	1	1	
Clostridium sporogenes		3.71	0.03	BGL 10	99	99	
Eubacterium saburreum		3.49	0.16		1	85	
43131000	**Low discrimination**						**43131000**
					GRA		
Eubacterium saburreum		57.74	0.57	IND 99	85		
Fusobacterium mortiferum		34.56	0.54		1		
Lactobacillus jensenii		1.71	0.22		99		
43131002	**Low discrimination**						**43131002**
					GRA		
Eubacterium saburreum		60.46	0.44	IND 99	85		
Fusobacterium mortiferum		22.78	0.38		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		6.75	0.19		99		
43131010	**Good identification**						**43131010**
					GRA		
Eubacterium saburreum		97.85	0.55	IND 99	85		
Lactobacillus jensenii		1.18	0.14		99		
Fusobacterium mortiferum		0.72	0.20		1		
43131012	**Good identification**						**43131012**
Eubacterium saburreum		95.65	0.42	IND 99			
Lactobacillus jensenii		4.35	0.11				
43131020	**Low discrimination**						**43131020**
Clostridium perfringens		66.02	0.41	MNS 99			
Eubacterium saburreum		28.92	0.44	LAC 88			
Lactobacillus jensenii		1.21	0.12				
43131022	**Low discrimination**						**43131022**
Clostridium perfringens		82.09	0.35	MNS 99			
Eubacterium saburreum		13.49	0.31	LAC 88			
Lactobacillus jensenii		2.13	0.09				
43131030	**Good identification**						**43131030**
Eubacterium saburreum		95.67	0.42	IND 99			
Clostridium perfringens		2.70	0.07				
Lactobacillus jensenii		1.63	0.04				
43131100	**Low discrimination**						**43131100**
					COC	GRA	
Eubacterium saburreum		37.22	0.63	IND 99	1	85	
Atopobium parvulum		36.09	0.53		99	99	
Fusobacterium mortiferum		22.28	0.61		1	1	
43131101	**Very good identification**						**43131101**
					COC	SPO	
Clostridium chauvoei		99.34	0.64	MNS 90	1	99	
Eubacterium saburreum		0.22	0.29	TRE 10	1	1	
Atopobium parvulum		0.21	0.19		99	1	
43131102	**Low discrimination**						**43131102**
					COC	GRA	
Eubacterium saburreum		38.23	0.50	IND 99	1	85	
Atopobium parvulum		37.07	0.40		99	99	
Fusobacterium mortiferum		14.41	0.44		1	1	
43131103	**Good identification**						**43131103**
					SPO		
Clostridium chauvoei		99.02	0.51	MNS 90	99		
Actinomyces naeslundii		0.54	0.36	TRE 10	1		
Eubacterium saburreum		0.22	0.16		1		
43131110	**Very good identification**						**43131110**
					COC	GRA	
Eubacterium saburreum		98.10	0.61	IND 99	1	85	
Atopobium parvulum		1.17	0.19		99	99	
Fusobacterium mortiferum		0.73	0.27		1	1	
43131112	**Acceptable identification**						**43131112**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		94.57	0.48	IND 99	85		
Bifidobacterium adolescentis		2.11	0.18		99		
Leptotrichia buccalis		1.68	-.03		1		
43131120	**Low discrimination**						**43131120**
					COC		
Clostridium perfringens		71.72	0.56	MNS 99	1		
Eubacterium saburreum		10.00	0.50		1		
Atopobium parvulum		9.69	0.40		99		
43131121	**Very good identification**						**43131121**
					SPO		
Clostridium chauvoei		98.59	0.56	MNS 90	99		
Clostridium perfringens		0.83	0.21	TRE 10	84		
Actinomyces naeslundii		0.29	0.36		1		
43131122	**Low discrimination**						**43131122**
					CAT	GRA	
Clostridium perfringens		78.79	0.50	MNS 99	1	99	
Capnocytophaga ochracea		8.37	0.30		90	1	
Eubacterium saburreum		4.12	0.37		1	85	
43131123	**Good identification**						**43131123**
					SPO		
Clostridium chauvoei		96.25	0.43	MNS 90	99		
Clostridium perfringens		2.16	0.16	TRE 10	84		
Actinomyces naeslundii		1.59	0.36		1		
43131130	**Low discrimination**						**43131130**
					COC		
Eubacterium saburreum		89.28	0.48	IND 99	1		
Clostridium perfringens		7.91	0.21		1		
Atopobium parvulum		1.07	0.06		99		
43131132	**Low discrimination**						**43131132**
					CAT	GRA	
Eubacterium saburreum		64.51	0.36	IND 99	1	85	
Capnocytophaga ochracea		17.80	0.14		90	1	
Clostridium perfringens		15.24	0.16		1	99	
43131160	**Good identification**						**43131160**
					COC		
Clostridium perfringens		97.83	0.43	MNS 99	1		
Eubacterium saburreum		0.72	0.16		1		
Atopobium parvulum		0.70	0.06		99		
43131162	**Acceptable identification**						**43131162**
					GRA	CAT	
Clostridium perfringens		94.17	0.38	MNS 99	99	1	
Capnocytophaga ochracea		5.83	0.14		1	90	
43131520	**Low discrimination**						**43131520**
Bifidobacterium longum subsp. longum		72.54	0.38	MNS 87			
Clostridium perfringens		16.44	0.21	RAF 97			
Lactobacillus fermentum		2.59	0.20				
43131522	**Low discrimination**						**43131522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.61	0.38	MNS 87	99	1	
Capnocytophaga ochracea		14.77	0.17	RAF 97	1	90	
Clostridium perfringens		7.96	0.16		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43131530	**Good identification**						**43131530**
Bifidobacterium longum subsp. longum		98.50	0.42	MNS 87			
Eubacterium saburreum		1.50	0.14	RAF 97			
43131532	**Good identification**						**43131532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.73	0.42	MNS 87	99	1	
Capnocytophaga ochracea		1.28	0.01	RAF 97	1	90	
43132100	**Good identification**						**43132100**
					GRA		
Propionibacterium propionicum		95.65	0.48	NIT 86	99		
Fusobacterium mortiferum		4.35	0.24	ESL 1	1		
43132102	**Good identification**						**43132102**
Propionibacterium propionicum		100.0	0.51	NIT 86			
43132140	**Good identification**						**43132140**
Propionibacterium propionicum		100.0	0.48	NIT 86			
43132142	**Good identification**						**43132142**
Propionibacterium propionicum		100.0	0.51	NIT 86			
43133100	**Low discrimination**						**43133100**
					COC		
Propionibacterium propionicum		71.58	0.48	NIT 86	1		
Eubacterium saburreum		8.15	0.29	ESL 1	1		
Atopobium parvulum		7.91	0.19		99		
43133102	**Low discrimination**						**43133102**
Propionibacterium propionicum		88.84	0.51	NIT 86			
Actinomyces naeslundii		3.82	0.38	ESL 1			
Actinomyces israelii		3.55	0.28				
43133123	**Low discrimination**						**43133123**
					SPO		
Actinomyces naeslundii		76.02	0.41	ESL 12	1		
Clostridium chauvoei		23.99	0.09	MNS 88	99		
43133140	**Good identification**						**43133140**
Propionibacterium propionicum		97.38	0.48	NIT 86			
Actinomyces israelii		1.52	0.18	ESL 1			
Actinomyces naeslundii		1.11	0.25				
43133142	**Good identification**						**43133142**
Propionibacterium propionicum		97.74	0.51	NIT 86			
Actinomyces israelii		1.52	0.21	ESL 1			
Actinomyces naeslundii		0.74	0.25				
43134100	**Low discrimination**						**43134100**
					COC	GRA	
Fusobacterium mortiferum		36.73	0.38	MLT 11	1	1	
Actinomyces meyeri		34.38	0.29	XYL 7	1	99	
Atopobium parvulum		22.24	0.23		99	99	
43135000	**Low discrimination**						**43135000**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		81.61	0.52	IND 99	1	85	
Fusobacterium mortiferum		6.83	0.35		1	1	
Atopobium parvulum		6.43	0.23		99	99	
43135002	**Low discrimination**						**43135002**
					COC		
Eubacterium saburreum		79.93	0.39	IND 99	1		
Atopobium parvulum		6.30	0.10		99		
Actinomyces israelii		5.30	0.23		1		
43135010	**Good identification**						**43135010**
Eubacterium saburreum		100.0	0.50	IND 99			
43135012	**Good identification**						**43135012**
Eubacterium saburreum		100.0	0.38	IND 99			
43135020	**Low discrimination**						**43135020**
					COC		
Eubacterium saburreum		85.13	0.39	IND 99	1		
Atopobium parvulum		6.71	0.10		99		
Clostridium perfringens		3.65	0.07		1		
43135030	**Good identification**						**43135030**
Eubacterium saburreum		100.0	0.38	IND 99			
43135100	**Low discrimination**						**43135100**
					COC	GRA	
Atopobium parvulum		73.89	0.57	MNS 99	99	99	
Eubacterium saburreum		22.10	0.58		1	85	
Fusobacterium mortiferum		1.85	0.42		1	1	
43135102	**Low discrimination**						**43135102**
					COC		
Atopobium parvulum		70.17	0.45	MNS 99	99		
Eubacterium saburreum		20.98	0.45		1		
Bifidobacterium adolescentis		3.90	0.31		1		
43135110	**Very good identification**						**43135110**
					COC		
Eubacterium saburreum		95.66	0.57	IND 99	1		
Atopobium parvulum		3.95	0.23		99		
Actinomyces israelii		0.23	0.16		1		
43135112	**Low discrimination**						**43135112**
					COC		
Eubacterium saburreum		74.15	0.44	IND 99	1		
Bifidobacterium adolescentis		20.56	0.32		1		
Atopobium parvulum		3.06	0.10		99		
43135120	**Low discrimination**						**43135120**
					COC		
Atopobium parvulum		72.71	0.45	MNS 99	99		
Eubacterium saburreum		21.74	0.45		1		
Clostridium perfringens		2.93	0.21		1		
43135130	**Acceptable identification**						**43135130**
					COC		
Eubacterium saburreum		92.70	0.44	IND 99	1		
Atopobium parvulum		3.83	0.10		99		
Bifidobacterium longum subsp. longum		3.47	0.16		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43135322	**Acceptable identification**						**43135322**
					GRA		
Parabacteroides distasonis		92.98	0.36	MNS 98	1		
Eubacterium contortum		7.02	0.06	RAF 99	99		
43135500	**Low discrimination**						**43135500**
					COC		
Bifidobacterium longum subsp. longum		48.57	0.36	MNS 87	1		
Atopobium parvulum		20.88	0.23	RAF 97	99		
Actinomyces meyeri		12.10	0.21		1		
43135502	**Low discrimination**						**43135502**
Bifidobacterium adolescentis		66.73	0.45	MNS 87			
Bifidobacterium longum subsp. longum		23.92	0.36	RAF 92			
Actinomyces israelii		5.00	0.32				
43135510	**Acceptable identification**						**43135510**
Bifidobacterium longum subsp. longum		91.39	0.39	MNS 87			
Eubacterium saburreum		5.65	0.23	RAF 97			
Bifidobacterium adolescentis		1.85	0.12				
43135512	**Low discrimination**						**43135512**
Bifidobacterium adolescentis		65.72	0.46	MNS 87			
Bifidobacterium longum subsp. longum		32.82	0.39	RAF 92			
Actinomyces israelii		0.60	0.18				
43135520	**Acceptable identification**						**43135520**
					COC	GRA	
Bifidobacterium longum subsp. longum		94.82	0.46	MNS 87	1	99	
Atopobium parvulum		1.80	0.10	RAF 97	99	99	
Bacteroides vulgatus		1.37	0.00		1	1	
43135522	**Acceptable identification**						**43135522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		94.04	0.46	MNS 87	99	1	
Parabacteroides distasonis		3.09	0.25	RAF 97	1	76	
Capnocytophaga ochracea		1.55	0.07		1	90	
43135530	**Good identification**						**43135530**
Bifidobacterium longum subsp. longum		99.73	0.50	MNS 87			
Eubacterium saburreum		0.27	0.10	RAF 97			
43135532	**Good identification**						**43135532**
					GRA		
Bifidobacterium longum subsp. longum		98.11	0.50	MNS 87	99		
Parabacteroides distasonis		1.02	0.20	RAF 97	1		
Bifidobacterium adolescentis		0.50	0.12		99		
43137002	**Good identification**						**43137002**
Actinomyces israelii		100.0	0.36	SAL 92			
43137100	**Low discrimination**						**43137100**
					COC		
Actinomyces israelii		62.46	0.43	SAL 92	1		
Atopobium parvulum		27.85	0.23	CEL 93	99		
Eubacterium saburreum		8.33	0.24		1		
43137102	**Low discrimination**						**43137102**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC			
Actinomyces israelii		75.42	0.46	SAL 92	1			
Bifidobacterium adolescentis		17.79	0.29	CEL 93	1			
Atopobium parvulum		3.96	0.10		99			
43137140	**Acceptable identification**							**43137140**
Actinomyces israelii		94.66	0.36	SAL 92				
Propionibacterium propionicum		5.34	0.14	CEL 93				
43137142	**Low discrimination**							**43137142**
Actinomyces israelii		88.39	0.39	SAL 92				
Bifidobacterium adolescentis		6.62	0.14	CEL 93				
Propionibacterium propionicum		4.99	0.17					
43137500	**Good identification**							**43137500**
Actinomyces israelii		97.16	0.42	SAL 92				
Bifidobacterium adolescentis		2.84	0.09	CEL 93				
43137502	**Low discrimination**							**43137502**
Bifidobacterium adolescentis		65.86	0.43	MNS 87				
Actinomyces israelii		34.14	0.45	RAF 92				
43137512	**Acceptable identification**							**43137512**
Bifidobacterium adolescentis		93.50	0.45	MNS 87				
Actinomyces israelii		5.93	0.31	RAF 92				
Bifidobacterium longum subsp. longum		0.58	0.05					
43137542	**Low discrimination**							**43137542**
Actinomyces israelii		61.99	0.38	SAL 92				
Bifidobacterium adolescentis		38.01	0.28	CEL 93				
43140000	**Low discrimination**							**43140000**
Clostridium botulinum biovar A		79.53	0.45	SUC 7				
Clostridium sporogenes		17.82	0.32	CEL 1				
Clostridium bifermentans		1.28	0.14					
43140040	**Good identification**							**43140040**
					COC	GRA		
Clostridium botulinum biovar A		96.94	0.36	SUC 7	1	99		
Clostridium novyi biovar A		1.28	0.16	CEL 1	1	99		
Megasphaera elsdenii		1.05	0.07		99	1		
43141101	**Good identification**							**43141101**
Clostridium chauvoei		100.0	0.40	MNS 90				
43150000	**Low discrimination**							**43150000**
					COC	GRA	SPO	
Fusobacterium mortiferum		68.27	0.52	MLT 11	1	1	1	
Megasphaera elsdenii		14.21	0.32		99	1	1	
Clostridium botulinum biovar A		9.84	0.25		1	99	99	
43150002	**Low discrimination**							**43150002**
					COC	GRA	SPO	
Fusobacterium mortiferum		67.76	0.36	MLT 11	1	1	1	
Clostridium sporogenes		18.15	0.11	BGL 10	1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Megasphaera elsdenii		14.10	0.15		99	1	1	
43150100	**Very good identification**							**43150100**
					GRA			
Fusobacterium mortiferum		96.14	0.59	MLT 11	1			
Actinomyces meyeri		3.13	0.24		99			
Collinsella aerofaciens		0.73	-.03		99			
43150102	**Acceptable identification**							**43150102**
					GRA	CAT	SPO	
Fusobacterium mortiferum		93.60	0.42	MLT 11	1	5	1	
Capnocytophaga ochracea		3.32	0.04	BGL 10	1	90	1	
Clostridium barati		3.08	0.02		99	1	99	
43151000	**Low discrimination**							**43151000**
					GRA			
Fusobacterium mortiferum		68.23	0.55	MLT 11	1			
Lactobacillus jensenii		19.11	0.36		99			
Lactobacillus catenaformis		7.31	0.29		99			
43151100	**Low discrimination**							**43151100**
					GRA			
Fusobacterium mortiferum		57.15	0.62	MLT 11	1			
Collinsella aerofaciens		28.63	0.32		99			
Lactobacillus catenaformis		3.93	0.32		99			
43151101	**Good identification**							**43151101**
					GRA	SPO		
Clostridium chauvoei		96.93	0.47	MNS 90	99	99		
Fusobacterium mortiferum		1.58	0.27	CEL 1	1	1		
Collinsella aerofaciens		0.79	-.03		99	1		
43151102	**Low discrimination**							**43151102**
					GRA	CAT		
Capnocytophaga ochracea		41.25	0.38	MNS 90	1	90		
Fusobacterium mortiferum		17.59	0.45	RAF 90	1	5		
Lactobacillus catenaformis		10.90	0.32	NAG 90	99	1		
43151120	**Low discrimination**							**43151120**
					GRA	CAT	SPO	
Prevotella melaninogenica		68.93	0.48	MNS 90	1	1	1	
Clostridium paraputrificum		11.14	0.32	RAF 90	99	1	99	
Capnocytophaga ochracea		10.35	0.38	CEL 10	1	90	1	
43151121	**Low discrimination**							**43151121**
					GRA	SPO		
Clostridium chauvoei		88.26	0.38	MNS 90	99	99		
Prevotella melaninogenica		7.48	0.14	CEL 1	1	1		
Actinomyces naeslundii		1.93	0.33		99	1		
43151122	**Low discrimination**							**43151122**
					CAT			
Capnocytophaga ochracea		65.38	0.54	MNS 90	90			
Prevotella oralis		16.31	0.43	RAF 90	1			
Prevotella melaninogenica		9.21	0.36		1			
43151132	**Good identification**							**43151132**
					CAT			
Capnocytophaga ochracea		95.15	0.38	MNS 90	90			
Prevotella oralis		2.16	0.09	RAF 90	1			
Prevotella melaninogenica		1.22	0.02	MLZ 10	1			
43151162	**Low discrimination**							**43151162**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Capnocytophaga ochracea		87.50	0.38	MNS 90	1	1		
Clostridium barati		4.06	0.14	RAF 90	99	99		
Clostridium paraputrificum		3.04	0.06	SOR 10	99	99		
43151322	**Low discrimination**							**43151322**
					GRA	CAT	SPO	
Prevotella oralis		86.15	0.37	MNS 99	1	1	1	
Capnocytophaga ochracea		8.14	0.20	RAF 92	1	90	1	
Clostridium barati		4.15	0.14		99	1	99	
43151502	**Low discrimination**							**43151502**
					GRA	CAT		
Bifidobacterium adolescentis		58.31	0.39	MNS 87	99	1		
Capnocytophaga ochracea		31.84	0.25	RAF 92	1	90		
Lactobacillus fermentum		6.26	0.28	XYL 87	99	1		
43151512	**Acceptable identification**							**43151512**
					GRA	CAT		
Bifidobacterium adolescentis		94.01	0.41	MNS 87	99	1		
Capnocytophaga ochracea		4.67	0.08	RAF 92	1	90		
Bifidobacterium longum subsp. longum		1.33	0.07	XYL 87	99	1		
43151522	**Good identification**							**43151522**
					GRA	CAT		
Capnocytophaga ochracea		95.77	0.41	MNS 90	1	90		
Prevotella oralis		1.37	0.09	RAF 90	1	1		
Bifidobacterium longum subsp. longum		0.79	0.14		99	1		
43153101	**Low discrimination**							**43153101**
					SPO			
Actinomyces naeslundii		58.34	0.38	ESL 12	1			
Clostridium chauvoei		41.66	0.12	MNS 88	99			
43153102	**Low discrimination**							**43153102**
Actinomyces israelii		50.34	0.37	SAL 92				
Bifidobacterium adolescentis		18.41	0.24	XYL 92				
Actinomyces naeslundii		10.47	0.35					
43153103	**Low discrimination**							**43153103**
					SPO			
Actinomyces naeslundii		88.81	0.38	ESL 12	1			
Clostridium chauvoei		11.19	-.01	MNS 88	99			
43153121	**Low discrimination**							**43153121**
					SPO			
Actinomyces naeslundii		80.77	0.38	ESL 12	1			
Clostridium chauvoei		19.23	0.04	MNS 88	99			
43153123	**Good identification**							**43153123**
Actinomyces naeslundii		100.0	0.38	ESL 12				
43153502	**Low discrimination**							**43153502**
Bifidobacterium adolescentis		74.95	0.38	MNS 87				
Actinomyces israelii		25.05	0.36	RAF 92				
43153512	**Good identification**							**43153512**
Bifidobacterium adolescentis		96.07	0.39	MNS 87				
Actinomyces israelii		3.93	0.22	RAF 92				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43154100	**Low discrimination**						**43154100**
					GRA		
Fusobacterium mortiferum		54.38	0.39	MLT 11	1		
Actinomyces meyeri		45.62	0.29	XYL 7	99		
43155000	**Low discrimination**						**43155000**
					GRA		
Fusobacterium mortiferum		47.96	0.36	MLT 11	1		
Actinomyces meyeri		23.47	0.21	XYL 7	99		
Actinomyces israelii		14.15	0.29		99		
43155100	**Low discrimination**						**43155100**
					COC	GRA	
Fusobacterium mortiferum		27.90	0.42	MLT 11	1	1	
Atopobium parvulum		23.55	0.30	XYL 7	99	99	
Actinomyces meyeri		23.40	0.32		1	99	
43155102	**Low discrimination**						**43155102**
					GRA	CAT	
Bifidobacterium adolescentis		34.94	0.39	MNS 87	99	1	
Capnocytophaga ochracea		27.03	0.27	RAF 92	1	90	
Actinomyces israelii		23.70	0.43	ARA 87	99	1	
43155112	**Low discrimination**						**43155112**
					GRA	CAT	
Bifidobacterium adolescentis		85.44	0.41	MNS 87	99	1	
Actinomyces israelii		7.09	0.29	RAF 92	99	1	
Capnocytophaga ochracea		6.01	0.11	ARA 87	1	90	
43155122	**Low discrimination**						**43155122**
					CAT		
Capnocytophaga ochracea		69.28	0.44	MNS 90	90		
Bacteroides fragilis		20.41	0.25	RAF 90	99		
Prevotella oralis		8.55	0.28		1		
43155142	**Low discrimination**						**43155142**
					GRA	CAT	
Actinomyces israelii		55.73	0.36	SAL 92	99	1	
Bifidobacterium adolescentis		26.11	0.24		99	1	
Capnocytophaga ochracea		18.16	0.11		1	90	
43155500	**Low discrimination**						**43155500**
Actinomyces israelii		50.43	0.38	SAL 92			
Actinomyces meyeri		25.56	0.21				
Bifidobacterium adolescentis		9.21	0.19				
43155502	**Low discrimination**						**43155502**
					GRA	CAT	
Bifidobacterium adolescentis		89.50	0.53	MNS 87	99	1	
Actinomyces israelii		7.42	0.41	RAF 92	99	1	
Capnocytophaga ochracea		1.83	0.15		1	90	
43155512	**Good identification**						**43155512**
Bifidobacterium adolescentis		98.20	0.55	MNS 87			
Actinomyces israelii		1.00	0.27	RAF 92			
Bifidobacterium longum subsp. longum		0.62	0.16				
43155542	**Low discrimination**						**43155542**
					GRA	CAT	
Bifidobacterium adolescentis		78.17	0.38	MNS 87	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		20.40	0.34	RAF 92	99	1		
Capnocytophaga ochracea		1.43	-.02	SOR 11	1	90		
43155552	**Good identification**							**43155552**
Bifidobacterium adolescentis		96.91	0.39	MNS 87				
Actinomyces israelii		3.09	0.20	RAF 92				
43157000	**Good identification**							**43157000**
Actinomyces israelii		100.0	0.42	SAL 92				
43157002	**Good identification**							**43157002**
Actinomyces israelii		100.0	0.45	SAL 92				
43157040	**Good identification**							**43157040**
Actinomyces israelii		100.0	0.35	SAL 92				
43157042	**Good identification**							**43157042**
Actinomyces israelii		100.0	0.38	SAL 92				
43157100	**Good identification**							**43157100**
Actinomyces israelii		100.0	0.53	SAL 92				
43157102	**Low discrimination**							**43157102**
					SPO			
Actinomyces israelii		82.06	0.56	SAL 92	1			
Bifidobacterium adolescentis		17.47	0.38		1			
Clostridium tertium		0.48	0.03		99			
43157110	**Good identification**							**43157110**
					SPO			
Actinomyces israelii		96.64	0.38	SAL 92	1			
Clostridium tertium		3.36	-.01	MLZ 13	99			
43157112	**Low discrimination**							**43157112**
					SPO			
Bifidobacterium adolescentis		61.95	0.39	MNS 87	1			
Actinomyces israelii		35.58	0.41	RAF 92	1			
Clostridium tertium		2.47	0.07	ARA 87	99			
43157122	**Low discrimination**							**43157122**
					GRA	SPO	CAT	
Actinomyces israelii		60.76	0.35	SAL 92	99	1	1	
Capnocytophaga ochracea		17.42	0.10	NAG 6	1	1	90	
Clostridium tertium		16.69	0.11		99	99	1	
43157140	**Good identification**							**43157140**
Actinomyces israelii		100.0	0.46	SAL 92				
43157142	**Acceptable identification**							**43157142**
Actinomyces israelii		93.66	0.49	SAL 92				
Bifidobacterium adolescentis		6.34	0.22					
43157302	**Good identification**							**43157302**
Actinomyces israelii		100.0	0.36	SAL 92				
43157400	**Good identification**							**43157400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.41	SAL 92			
43157402	**Acceptable identification**						**43157402**
Actinomyces israelii		93.43	0.44	SAL 92			
Bifidobacterium adolescentis		6.57	0.18				
43157442	**Good identification**						**43157442**
Actinomyces israelii		100.0	0.37	SAL 92			
43157500	**Good identification**						**43157500**
Actinomyces israelii		97.43	0.51	SAL 92			
Bifidobacterium adolescentis		2.57	0.18				
43157502	**Low discrimination**						**43157502**
Bifidobacterium adolescentis		63.52	0.52	MNS 87			
Actinomyces israelii		36.48	0.54	RAF 92			
43157510	**Low discrimination**						**43157510**
Actinomyces israelii		82.25	0.37	SAL 92			
Bifidobacterium adolescentis		17.75	0.19	MLZ 13			
43157512	**Acceptable identification**						**43157512**
Bifidobacterium adolescentis		93.44	0.53	MNS 87			
Actinomyces israelii		6.56	0.40	RAF 92			
43157540	**Good identification**						**43157540**
Actinomyces israelii		100.0	0.44	SAL 92			
43157542	**Low discrimination**						**43157542**
Actinomyces israelii		64.38	0.47	SAL 92			
Bifidobacterium adolescentis		35.62	0.36				
43157552	**Low discrimination**						**43157552**
Bifidobacterium adolescentis		81.90	0.38	MNS 87			
Actinomyces israelii		18.10	0.33	RAF 92			
43170000	**Low discrimination**						**43170000**
					GRA	SPO	
Fusobacterium mortiferum		79.18	0.43	MLT 11	1	1	
Clostridium botulinum biovar A		9.55	0.14		99	99	
Clostridium sporogenes		7.89	0.11		99	99	
43170100	**Low discrimination**						**43170100**
					COC	GRA	
Fusobacterium mortiferum		85.34	0.50	MLT 11	1	1	
Atopobium parvulum		14.66	0.25		99	99	
43171000	**Low discrimination**						**43171000**
					COC	GRA	
Lactobacillus jensenii		67.44	0.44	MNS 99	1	99	
Fusobacterium mortiferum		23.97	0.46	SAL 95	1	1	
Atopobium parvulum		6.41	0.25		99	99	
43171002	**Low discrimination**						**43171002**
					COC	GRA	
Lactobacillus jensenii		87.83	0.41	MNS 99	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		5.20	0.30	SAL 95	1	1	
Atopobium parvulum		2.21	0.12		99	99	
43171010	**Acceptable identification**						**43171010**
					GRA		
Lactobacillus jensenii		94.79	0.36	MNS 99	99		
Eubacterium saburreum		4.19	0.21	SAL 95	85		
Fusobacterium mortiferum		1.02	0.12		1		
43171100	**Good identification**						**43171100**
					COC	GRA	
Atopobium parvulum		90.77	0.59	MNS 99	99	99	
Fusobacterium mortiferum		8.01	0.53		1	1	
Eubacterium saburreum		0.41	0.29		1	85	
43171102	**Low discrimination**						**43171102**
					COC	GRA	
Atopobium parvulum		82.30	0.46	MNS 99	99	99	
Lactobacillus acidophilus		6.03	0.25		1	99	
Fusobacterium mortiferum		4.57	0.36		1	1	
43171120	**Low discrimination**						**43171120**
					COC	SPO	
Atopobium parvulum		71.37	0.46	MNS 99	99	1	
Clostridium perfringens		9.43	0.32		1	84	
Clostridium paraputrificum		8.90	0.22		1	99	
43171123	**Low discrimination**						**43171123**
					SPO		
Actinomyces naeslundii		68.40	0.38	ESL 12	1		
Clostridium chauvoei		31.60	0.09	MNS 88	99		
43171502	**Low discrimination**						**43171502**
					COC		
Bifidobacterium adolescentis		72.47	0.36	MNS 87	1		
Actinomyces israelii		19.74	0.33	RAF 92	1		
Atopobium parvulum		7.80	0.12	XYL 87	99		
43171512	**Acceptable identification**						**43171512**
					GRA		
Bifidobacterium adolescentis		91.81	0.38	MNS 87	99		
Leptotrichia buccalis		5.13	-.03	RAF 92	1		
Actinomyces israelii		3.06	0.19	XYL 87	99		
43173001	**Good identification**						**43173001**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43173002	**Low discrimination**						**43173002**
Actinomyces israelii		80.48	0.37	SAL 92			
Actinomyces naeslundii		12.14	0.32	XYL 92			
Lactobacillus jensenii		7.38	0.07				
43173003	**Good identification**						**43173003**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43173021	**Good identification**						**43173021**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43173023	**Good identification**						**43173023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	ESL 12		
43173100	**Low discrimination**					**43173100**
					COC	
Actinomyces israelii		56.63	0.44	SAL 92	1	
Atopobium parvulum		27.46	0.25	XYL 92	99	
Actinomyces naeslundii		9.12	0.40		1	
43173101	**Good identification**					**43173101**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43173102	**Low discrimination**					**43173102**
					SPO	
Actinomyces israelii		72.13	0.47	SAL 92	1	
Clostridium tertium		8.40	0.17	XYL 92	99	
Actinomyces naeslundii		7.74	0.40		1	
43173103	**Good identification**					**43173103**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43173121	**Good identification**					**43173121**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43173123	**Good identification**					**43173123**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43173140	**Low discrimination**					**43173140**
Actinomyces israelii		89.13	0.37	SAL 92		
Actinomyces naeslundii		6.51	0.27	XYL 92		
Propionibacterium propionicum		4.36	0.14			
43173142	**Acceptable identification**					**43173142**
Actinomyces israelii		91.11	0.40	SAL 92		
Propionibacterium propionicum		4.45	0.17	XYL 92		
Actinomyces naeslundii		4.44	0.27			
43173402	**Good identification**					**43173402**
Actinomyces israelii		100.0	0.35	SAL 92		
43173500	**Good identification**					**43173500**
Actinomyces israelii		100.0	0.43	SAL 92		
43173502	**Low discrimination**					**43173502**
Actinomyces israelii		65.35	0.46	SAL 92		
Bifidobacterium adolescentis		34.65	0.35	XYL 92		
43173512	**Low discrimination**					**43173512**
Bifidobacterium adolescentis		81.26	0.36	MNS 87		
Actinomyces israelii		18.74	0.32	RAF 92		
43173540	**Good identification**					**43173540**
Actinomyces israelii		100.0	0.36	SAL 92		
43173542	**Low discrimination**					**43173542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		85.58	0.39	SAL 92			
Bifidobacterium adolescentis		14.42	0.19	XYL 92			
43175002	**Low discrimination**						**43175002**
					COC		
Actinomyces israelii		79.75	0.42	SAL 92	1		
Atopobium parvulum		16.65	0.17		99		
Lactobacillus jensenii		3.60	0.07		1		
43175042	**Good identification**						**43175042**
Actinomyces israelii		100.0	0.35	SAL 92			
43175100	**Very good identification**						**43175100**
					COC	GRA	
Atopobium parvulum		97.29	0.64	MNS 99	99	99	
Actinomyces israelii		2.22	0.49		1	99	
Fusobacterium mortiferum		0.35	0.33		1	1	
43175102	**Low discrimination**						**43175102**
					COC		
Atopobium parvulum		80.16	0.51	MNS 99	99		
Actinomyces israelii		15.51	0.52		1		
Bifidobacterium adolescentis		4.05	0.36		1		
43175112	**Low discrimination**						**43175112**
					COC		
Bifidobacterium adolescentis		58.67	0.38	MNS 87	1		
Actinomyces israelii		27.46	0.38	RAF 92	1		
Atopobium parvulum		9.59	0.17	ARA 87	99		
43175120	**Good identification**						**43175120**
					COC		
Atopobium parvulum		99.06	0.51	MNS 99	99		
Actinomyces israelii		0.82	0.29		1		
Eubacterium saburreum		0.13	0.11		1		
43175122	**Low discrimination**						**43175122**
					COC	GRA	
Atopobium parvulum		85.47	0.38	MNS 99	99	99	
Actinomyces israelii		5.98	0.32		1	99	
Parabacteroides distasonis		3.19	0.24		1	1	
43175142	**Low discrimination**						**43175142**
					COC		
Actinomyces israelii		82.15	0.45	SAL 92	1		
Atopobium parvulum		11.03	0.17		99		
Bifidobacterium adolescentis		6.82	0.21		1		
43175400	**Good identification**						**43175400**
Actinomyces israelii		100.0	0.38	SAL 92			
43175402	**Acceptable identification**						**43175402**
Actinomyces israelii		92.05	0.41	SAL 92			
Bifidobacterium adolescentis		7.95	0.16				
43175500	**Low discrimination**						**43175500**
					COC	SPO	
Actinomyces israelii		59.71	0.48	SAL 92	1	1	
Atopobium parvulum		32.38	0.30		99	1	
Clostridium butyricum		5.98	0.10		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43175502	**Low discrimination**						**43175502**
					COC		
Bifidobacterium adolescentis		66.09	0.50	MNS 87	1		
Actinomyces israelii		30.93	0.51	RAF 92	1		
Atopobium parvulum		1.97	0.17		99		
43175512	**Acceptable identification**						**43175512**
Bifidobacterium adolescentis		93.32	0.52	MNS 87			
Actinomyces israelii		5.34	0.37	RAF 92			
Clostridium clostridioforme		1.12	0.16				
43175540	**Good identification**						**43175540**
Actinomyces israelii		100.0	0.41	SAL 92			
43175542	**Low discrimination**						**43175542**
Actinomyces israelii		59.56	0.44	SAL 92			
Bifidobacterium adolescentis		40.45	0.35				
43175552	**Low discrimination**						**43175552**
Bifidobacterium adolescentis		84.74	0.36	MNS 87			
Actinomyces israelii		15.26	0.30	RAF 92			
43177000	**Good identification**						**43177000**
Actinomyces israelii		100.0	0.52	SAL 92			
43177002	**Good identification**						**43177002**
Actinomyces israelii		100.0	0.55	SAL 92			
43177010	**Good identification**						**43177010**
Actinomyces israelii		100.0	0.38	SAL 92			
43177012	**Good identification**						**43177012**
Actinomyces israelii		100.0	0.41	SAL 92			
43177040	**Good identification**						**43177040**
Actinomyces israelii		100.0	0.45	SAL 92			
43177042	**Good identification**						**43177042**
Actinomyces israelii		100.0	0.48	SAL 92			
43177100	**Good identification**						**43177100**
					COC	SPO	
Actinomyces israelii		92.14	0.62	SAL 92	1	1	
Atopobium parvulum		7.22	0.30		99	1	
Clostridium tertium		0.64	0.11		1	99	
43177102	**Good identification**						**43177102**
					SPO		
Actinomyces israelii		94.26	0.65	SAL 92	1		
Bifidobacterium adolescentis		3.56	0.35		1		
Clostridium tertium		1.32	0.19		99		
43177110	**Acceptable identification**						**43177110**
					SPO		
Actinomyces israelii		92.32	0.48	SAL 92	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		7.68	0.15	MLZ 13	99		
43177112	**Low discrimination**						**43177112**
					SPO		
Actinomyces israelii		67.80	0.51	SAL 92	1		
Bifidobacterium adolescentis		20.92	0.36	MLZ 13	1		
Clostridium tertium		11.28	0.24		99		
43177120	**Low discrimination**						**43177120**
					COC	SPO	
Actinomyces israelii		64.73	0.42	SAL 92	1	1	
Clostridium tertium		21.26	0.19	NAG 6	1	99	
Atopobium parvulum		14.01	0.17		99	1	
43177122	**Low discrimination**						**43177122**
					COC	SPO	
Actinomyces israelii		59.44	0.45	SAL 92	1	1	
Clostridium tertium		39.05	0.27	NAG 6	1	99	
Atopobium parvulum		1.51	0.04		99	1	
43177140	**Good identification**						**43177140**
Actinomyces israelii		100.0	0.55	SAL 92			
43177142	**Very good identification**						**43177142**
Actinomyces israelii		98.82	0.58	SAL 92			
Bifidobacterium adolescentis		1.19	0.19				
43177150	**Good identification**						**43177150**
Actinomyces israelii		100.0	0.41	SAL 92			
43177152	**Acceptable identification**						**43177152**
Actinomyces israelii		91.07	0.44	SAL 92			
Bifidobacterium adolescentis		8.93	0.21	MLZ 13			
43177162	**Good identification**						**43177162**
Actinomyces israelii		100.0	0.38	SAL 92			
43177202	**Good identification**						**43177202**
Actinomyces israelii		100.0	0.36	SAL 92			
43177300	**Good identification**						**43177300**
Actinomyces israelii		100.0	0.43	SAL 92			
43177302	**Good identification**						**43177302**
Actinomyces israelii		100.0	0.46	SAL 92			
43177340	**Good identification**						**43177340**
Actinomyces israelii		100.0	0.36	SAL 92			
43177342	**Good identification**						**43177342**
Actinomyces israelii		100.0	0.39	SAL 92			
43177400	**Good identification**						**43177400**
Actinomyces israelii		100.0	0.51	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43177402	**Good identification**					**43177402**
Actinomyces israelii		98.77	0.54	SAL 92		
Bifidobacterium adolescentis		1.23	0.15			
43177410	**Good identification**					**43177410**
Actinomyces israelii		100.0	0.36	SAL 92		
43177412	**Acceptable identification**					**43177412**
Actinomyces israelii		90.75	0.39	SAL 92		
Bifidobacterium adolescentis		9.25	0.16	MLZ 13		
43177440	**Good identification**					**43177440**
Actinomyces israelii		100.0	0.44	SAL 92		
43177442	**Good identification**					**43177442**
Actinomyces israelii		100.0	0.47	SAL 92		
43177500	**Very good identification**					**43177500**
					SPO	
Actinomyces israelii		99.34	0.61	SAL 92	1	
Bifidobacterium adolescentis		0.46	0.15		1	
Clostridium butyricum		0.20	-.07		99	
43177502	**Low discrimination**					**43177502**
Actinomyces israelii		76.42	0.64	SAL 92		
Bifidobacterium adolescentis		23.58	0.49			
43177510	**Good identification**					**43177510**
Actinomyces israelii		96.32	0.47	SAL 92		
Bifidobacterium adolescentis		3.68	0.16	MLZ 13		
43177512	**Low discrimination**					**43177512**
Bifidobacterium adolescentis		70.25	0.50	MNS 87		
Actinomyces israelii		27.84	0.50	RAF 92		
Bifidobacterium dentium		1.91	0.06			
43177520	**Good identification**					**43177520**
Actinomyces israelii		100.0	0.40	SAL 92		
43177522	**Good identification**					**43177522**
Actinomyces israelii		95.35	0.43	SAL 92		
Bifidobacterium adolescentis		4.66	0.15	NAG 6		
43177540	**Good identification**					**43177540**
Actinomyces israelii		100.0	0.54	SAL 92		
43177542	**Good identification**					**43177542**
Actinomyces israelii		91.07	0.57	SAL 92		
Bifidobacterium adolescentis		8.93	0.33			
43177550	**Good identification**					**43177550**
					COC	
Actinomyces israelii		95.22	0.40	SAL 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		4.78	-.03	MLZ 13	99		
43177552	**Low discrimination**						**43177552**
Actinomyces israelii		55.49	0.43	SAL 92			
Bifidobacterium adolescentis		44.51	0.35	MLZ 13			
43177562	**Good identification**						**43177562**
Actinomyces israelii		100.0	0.36	SAL 92			
43177700	**Good identification**						**43177700**
Actinomyces israelii		100.0	0.42	SAL 92			
43177702	**Good identification**						**43177702**
Actinomyces israelii		96.02	0.45	SAL 92			
Bifidobacterium adolescentis		3.98	0.15	RHA 7			
43177742	**Good identification**						**43177742**
Actinomyces israelii		100.0	0.38	SAL 92			
43200000	**Good identification**						**43200000**
					GRA	SPO	
Eubacterium tortuosum		93.52	0.66	GAL 99	99	1	
Clostridium botulinum biovar A		4.36	0.47		99	99	
Fusobacterium mortiferum		0.75	0.47		1	1	
43200002	**Good identification**						**43200002**
					SPO		
Eubacterium tortuosum		96.42	0.53	GAL 99	1		
Atopobium minutum		1.81	0.35		1		
Clostridium sporogenes		0.84	0.19		99		
43200020	**Good identification**						**43200020**
					SPO		
Eubacterium tortuosum		98.40	0.53	GAL 99	1		
Atopobium minutum		0.92	0.30		1		
Clostridium novyi biovar A		0.27	0.25		99		
43200022	**Good identification**						**43200022**
					GRA		
Eubacterium tortuosum		95.87	0.40	GAL 99	99		
Atopobium minutum		3.40	0.27		99		
Pseudoflavonifractor capillosus		0.47	-.03		1		
43200040	**Low discrimination**						**43200040**
					SPO		
Clostridium botulinum biovar A		54.37	0.38	SUC 7	99		
Eubacterium tortuosum		39.46	0.32	SAL 1	1		
Clostridium novyi biovar A		5.06	0.32		99		
43200100	**Low discrimination**						**43200100**
					GRA	SPO	
Fusobacterium mortiferum		63.94	0.53	FRU 99	1	1	
Eubacterium tortuosum		34.45	0.32		99	1	
Clostridium botulinum biovar A		1.61	0.13		99	99	
43200102	**Low discrimination**						**43200102**
					GRA		
Fusobacterium mortiferum		53.89	0.37	FRU 99	1		
Eubacterium tortuosum		46.11	0.19	BGL 10	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
43201000	**Excellent identification**						**43201000**
					GRA		
Eubacterium tortuosum		99.99	1.00		99		
Fusobacterium mortiferum		0.01	0.50		1		
Atopobium minutum		0.00	0.30		99		
43201001	**Very good identification**						**43201001**
					GRA	SPO	
Eubacterium tortuosum		99.97	0.66	URE 1	99	1	
Clostridium novyi biovar A		0.02	0.19		99	99	
Fusobacterium mortiferum		0.01	0.16		1	1	
43201002	**Very good identification**						**43201002**
					GRA		
Eubacterium tortuosum		99.99	0.87		99		
Fusobacterium mortiferum		0.01	0.33		1		
Atopobium minutum		0.01	0.27		99		
43201003	**Good identification**						**43201003**
Eubacterium tortuosum		100.0	0.53	URE 1			
43201010	**Very good identification**						**43201010**
					GRA	SPO	
Eubacterium tortuosum		99.97	0.66	MLZ 1	99	1	
Clostridium novyi biovar A		0.02	0.19		99	99	
Fusobacterium mortiferum		0.01	0.16		1	1	
43201012	**Good identification**						**43201012**
Eubacterium tortuosum		100.0	0.53	MLZ 1			
43201020	**Very good identification**						**43201020**
					GRA		
Eubacterium tortuosum		100.0	0.87		99		
Atopobium minutum		0.00	0.21		99		
Fusobacterium mortiferum		0.00	0.16		1		
43201021	**Good identification**						**43201021**
Eubacterium tortuosum		100.0	0.53	URE 1			
43201022	**Very good identification**						**43201022**
Eubacterium tortuosum		99.99	0.74				
Atopobium minutum		0.01	0.18				
43201023	**Good identification**						**43201023**
Eubacterium tortuosum		100.0	0.40	URE 1			
43201030	**Good identification**						**43201030**
Eubacterium tortuosum		100.0	0.53	MLZ 1			
43201032	**Good identification**						**43201032**
Eubacterium tortuosum		100.0	0.40	MLZ 1			
43201040	**Very good identification**						**43201040**
					GRA	SPO	
Eubacterium tortuosum		99.96	0.66	SOR 1	99	1	
Clostridium novyi biovar A		0.01	0.15		99	99	
Fusobacterium mortiferum		0.01	0.16		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43201042	**Good identification**					**43201042**
Eubacterium tortuosum		100.0	0.53	SOR 1		
43201060	**Good identification**					**43201060**
Eubacterium tortuosum		100.0	0.53	SOR 1		
43201062	**Good identification**					**43201062**
Eubacterium tortuosum		100.0	0.40	SOR 1		
43201100	**Very good identification**					**43201100**
					GRA	
Eubacterium tortuosum		97.25	0.66	LAC 1	99	
Fusobacterium mortiferum		2.74	0.56		1	
Eubacterium saburreum		0.01	0.13		85	
43201102	**Good identification**					**43201102**
					GRA	
Eubacterium tortuosum		98.26	0.53	LAC 1	99	
Fusobacterium mortiferum		1.74	0.40		1	
43201120	**Good identification**					**43201120**
					GRA	
Eubacterium tortuosum		99.84	0.53	LAC 1	99	
Fusobacterium mortiferum		0.16	0.22		1	
43201122	**Good identification**					**43201122**
Eubacterium tortuosum		100.0	0.40	LAC 1		
43201200	**Very good identification**					**43201200**
					GRA	
Eubacterium tortuosum		99.99	0.66	RHA 1	99	
Fusobacterium mortiferum		0.01	0.16		1	
43201202	**Good identification**					**43201202**
Eubacterium tortuosum		100.0	0.53	RHA 1		
43201220	**Good identification**					**43201220**
Eubacterium tortuosum		100.0	0.53	RHA 1		
43201222	**Good identification**					**43201222**
Eubacterium tortuosum		100.0	0.40	RHA 1		
43201400	**Very good identification**					**43201400**
					GRA	
Eubacterium tortuosum		99.99	0.66	ARA 1	99	
Fusobacterium mortiferum		0.01	0.16		1	
43201402	**Good identification**					**43201402**
Eubacterium tortuosum		100.0	0.53	ARA 1		
43201420	**Good identification**					**43201420**
Eubacterium tortuosum		100.0	0.53	ARA 1		
43201422	**Good identification**					**43201422**
Eubacterium tortuosum		100.0	0.40	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43203000	**Very good identification**						**43203000**
					GRA	SPO	
Eubacterium tortuosum		99.98	0.66	MAN 1	99	1	
Fusobacterium mortiferum		0.01	0.16		1	1	
Clostridium innocuum		0.01	0.00		99	99	
43203002	**Good identification**						**43203002**
					SPO		
Eubacterium tortuosum		99.95	0.53	MAN 1	1		
Clostridium innocuum		0.05	0.00		99		
43203020	**Good identification**						**43203020**
Eubacterium tortuosum		100.0	0.53	MAN 1			
43203022	**Good identification**						**43203022**
Eubacterium tortuosum		100.0	0.40	MAN 1			
43205000	**Very good identification**						**43205000**
					GRA		
Eubacterium tortuosum		99.91	0.66	XYL 1	99		
Fusobacterium mortiferum		0.09	0.30		1		
43205002	**Good identification**						**43205002**
					GRA		
Eubacterium tortuosum		99.94	0.53	XYL 1	99		
Fusobacterium mortiferum		0.06	0.14		1		
43205020	**Good identification**						**43205020**
Eubacterium tortuosum		100.0	0.53	XYL 1			
43205022	**Good identification**						**43205022**
Eubacterium tortuosum		100.0	0.40	XYL 1			
43210000	**Very good identification**						**43210000**
					GRA		
Fusobacterium mortiferum		96.86	0.81		1		
Eubacterium tortuosum		1.23	0.32		99		
Atopobium minutum		0.56	0.37		99		
43210001	**Good identification**						**43210001**
					GRA	SPO	
Fusobacterium mortiferum		96.19	0.47	URE 1	1	1	
Clostridium novyi biovar A		2.59	0.19		99	99	
Eubacterium tortuosum		1.22	-.03		99	1	
43210002	**Good identification**						**43210002**
					GRA		
Fusobacterium mortiferum		94.27	0.65	BGL 10	1		
Atopobium minutum		3.26	0.34		99		
Eubacterium tortuosum		1.90	0.19		99		
43210010	**Low discrimination**						**43210010**
					GRA	SPO	
Clostridium difficile		63.31	0.35	SUC 1	99	99	
Fusobacterium mortiferum		35.30	0.47	MAN 90	1	1	
Clostridium novyi biovar A		0.95	0.19		99	99	
43210020	**Low discrimination**						**43210020**
					GRA		
Fusobacterium mortiferum		70.81	0.47	NAG 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium tortuosum		15.71	0.19		99			
Atopobium minutum		13.48	0.29		99			
43210040	**Low discrimination**							**43210040**
					COC	GRA	SPO	
Fusobacterium mortiferum		71.50	0.47	SOR 1	1	1	1	
Megasphaera elsdenii		13.89	0.26		99	1	1	
Clostridium botulinum biovar A		7.91	0.17		1	99	99	
43210100	**Very good identification**							**43210100**
					GRA			
Fusobacterium mortiferum		99.93	0.87		1			
Actinomyces meyeri		0.06	0.23		99			
Eubacterium tortuosum		0.01	-.03		99			
43210101	**Good identification**							**43210101**
Fusobacterium mortiferum		100.0	0.53	URE 1				
43210102	**Very good identification**							**43210102**
Fusobacterium mortiferum		100.0	0.71	BGL 10				
43210103	**Good identification**							**43210103**
Fusobacterium mortiferum		100.0	0.37	URE 1				
43210110	**Good identification**							**43210110**
					GRA	SPO		
Fusobacterium mortiferum		99.23	0.53	MLZ 1	1	1		
Clostridium difficile		0.77	0.01		99	99		
43210112	**Good identification**							**43210112**
Fusobacterium mortiferum		100.0	0.37	MLZ 1				
43210120	**Good identification**							**43210120**
					GRA			
Fusobacterium mortiferum		99.18	0.53	NAG 1	1			
Actinomyces meyeri		0.82	0.08		99			
43210122	**Good identification**							**43210122**
					GRA	SPO		
Fusobacterium mortiferum		95.93	0.37	NAG 1	1	1		
Clostridium barati		4.08	-.02	BGL 10	99	99		
43210140	**Good identification**							**43210140**
Fusobacterium mortiferum		100.0	0.53	SOR 1				
43210142	**Good identification**							**43210142**
Fusobacterium mortiferum		100.0	0.37	SOR 1				
43210200	**Good identification**							**43210200**
					GRA			
Fusobacterium mortiferum		98.75	0.47	RHA 1	1			
Eubacterium tortuosum		1.25	-.03		99			
43210300	**Good identification**							**43210300**
Fusobacterium mortiferum		100.0	0.53	RHA 1				
43210302	**Good identification**							**43210302**
Fusobacterium mortiferum		100.0	0.37	RHA 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43210400	**Good identification**						**43210400**
					GRA		
Fusobacterium mortiferum		97.89	0.47	ARA 1	1		
Eubacterium tortuosum		1.24	-.03		99		
Actinomyces meyeri		0.87	0.02		99		
43210500	**Good identification**						**43210500**
					GRA		
Fusobacterium mortiferum		98.51	0.53	ARA 1	1		
Actinomyces meyeri		1.49	0.13		99		
43210502	**Good identification**						**43210502**
Fusobacterium mortiferum		100.0	0.37	ARA 1			
43211000	**Low discrimination**						**43211000**
					GRA		
Fusobacterium mortiferum		54.33	0.84		1		
Eubacterium tortuosum		45.54	0.66		99		
Atopobium minutum		0.07	0.29		99		
43211001	**Low discrimination**						**43211001**
					GRA		
Fusobacterium mortiferum		54.40	0.50	URE 1	1		
Eubacterium tortuosum		45.60	0.32		99		
43211002	**Low discrimination**						**43211002**
					GRA		
Eubacterium tortuosum		56.76	0.53	FRU 1	99		
Fusobacterium mortiferum		42.64	0.68		1		
Atopobium minutum		0.33	0.26		99		
43211010	**Low discrimination**						**43211010**
					GRA	SPO	
Fusobacterium mortiferum		53.47	0.50	MLZ 1	1	1	
Eubacterium tortuosum		44.81	0.32		99	1	
Clostridium difficile		0.65	0.01		99	99	
43211020	**Acceptable identification**						**43211020**
					GRA		
Eubacterium tortuosum		93.18	0.53	FRU 1	99		
Fusobacterium mortiferum		6.36	0.50		1		
Atopobium minutum		0.27	0.21		99		
43211022	**Acceptable identification**						**43211022**
					GRA	SPO	
Eubacterium tortuosum		93.08	0.40	FRU 1	99	1	
Fusobacterium mortiferum		4.00	0.33		1	1	
Clostridium barati		1.68	0.12		99	99	
43211040	**Low discrimination**						**43211040**
					GRA		
Fusobacterium mortiferum		54.40	0.50	SOR 1	1		
Eubacterium tortuosum		45.60	0.32		99		
43211100	**Very good identification**						**43211100**
					GRA		
Fusobacterium mortiferum		99.53	0.90		1		
Eubacterium tortuosum		0.36	0.32		99		
Actinomyces meyeri		0.06	0.26		99		
43211101	**Good identification**						**43211101**
					GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		98.35	0.56	URE 1	1	1	
Clostridium chauvoei		1.30	0.12		99	99	
Eubacterium tortuosum		0.36	-.03		99	1	
43211102	**Very good identification**						**43211102**
					GRA	SPO	
Fusobacterium mortiferum		98.94	0.74	BGL 10	1	1	
Eubacterium tortuosum		0.57	0.19		99	1	
Clostridium barati		0.18	0.12		99	99	
43211103	**Good identification**						**43211103**
					GRA	SPO	
Fusobacterium mortiferum		97.95	0.40	URE 1	1	1	
Clostridium chauvoei		2.05	-.01	BGL 10	99	99	
43211110	**Good identification**						**43211110**
					GRA		
Fusobacterium mortiferum		98.68	0.56	MLZ 1	1		
Eubacterium saburreum		0.96	0.20		85		
Eubacterium tortuosum		0.36	-.03		99		
43211112	**Good identification**						**43211112**
Fusobacterium mortiferum		100.0	0.40	MLZ 1			
43211120	**Low discrimination**						**43211120**
					GRA	SPO	
Fusobacterium mortiferum		89.85	0.56	NAG 1	1	1	
Eubacterium tortuosum		5.70	0.19		99	1	
Clostridium barati		1.79	0.12		99	99	
43211122	**Low discrimination**						**43211122**
					GRA	SPO	
Fusobacterium mortiferum		50.81	0.40	NAG 1	1	1	
Clostridium barati		27.35	0.20	BGL 10	99	99	
Bifidobacterium longum subsp. infantis		9.71	0.10		99	1	
43211140	**Good identification**						**43211140**
					GRA		
Fusobacterium mortiferum		99.64	0.56	SOR 1	1		
Eubacterium tortuosum		0.36	-.03		99		
43211142	**Good identification**						**43211142**
Fusobacterium mortiferum		100.0	0.40	SOR 1			
43211200	**Low discrimination**						**43211200**
					GRA		
Fusobacterium mortiferum		54.40	0.50	RHA 1	1		
Eubacterium tortuosum		45.60	0.32		99		
43211300	**Low discrimination**						**43211300**
					GRA		
Fusobacterium mortiferum		82.57	0.56	RHA 1	1		
Eubacterium contortum		17.14	0.32		99		
Eubacterium tortuosum		0.30	-.03		99		
43211302	**Low discrimination**						**43211302**
					GRA		
Fusobacterium mortiferum		75.21	0.40	RHA 1	1		
Eubacterium contortum		24.79	0.19	BGL 10	99		
43211400	**Low discrimination**						**43211400**
					GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		53.88	0.50	ARA 1	1		
Eubacterium tortuosum		45.16	0.32		99		
Lactobacillus fermentum		0.49	0.16		99		
43211500	**Good identification**						**43211500**
					GRA		
Fusobacterium mortiferum		97.55	0.56	ARA 1	1		
Actinomyces meyeri		1.48	0.16		99		
Lactobacillus fermentum		0.62	0.20		99		
43211502	**Acceptable identification**						**43211502**
					GRA		
Fusobacterium mortiferum		90.92	0.40	ARA 1	1		
Lactobacillus fermentum		5.18	0.20	BGL 10	99		
Bifidobacterium adolescentis		3.90	0.12		99		
43212000	**Low discrimination**						**43212000**
					GRA	SPO	
Clostridium difficile		59.26	0.40	SUC 1	99	99	
Fusobacterium mortiferum		17.93	0.47		1	1	
Eubacterium limosum		17.00	0.29		99	1	
43212010	**Good identification**						**43212010**
					GRA	SPO	
Clostridium difficile		99.88	0.52	SUC 1	99	99	
Fusobacterium mortiferum		0.06	0.13		1	1	
Eubacterium limosum		0.06	-.06		99	1	
43212100	**Good identification**						**43212100**
					GRA	SPO	
Fusobacterium mortiferum		98.19	0.53	MAN 1	1	1	
Clostridium difficile		1.41	0.06		99	99	
Eubacterium limosum		0.40	-.06		99	1	
43212102	**Good identification**						**43212102**
Fusobacterium mortiferum		100.0	0.37	MAN 1			
43213000	**Low discrimination**						**43213000**
					GRA	SPO	
Fusobacterium mortiferum		50.15	0.50	MAN 1	1	1	
Eubacterium tortuosum		42.03	0.32		99	1	
Clostridium innocuum		6.39	0.22		99	99	
43213100	**Good identification**						**43213100**
					GRA		
Fusobacterium mortiferum		99.64	0.56	MAN 1	1		
Eubacterium tortuosum		0.36	-.03		99		
43213102	**Good identification**						**43213102**
Fusobacterium mortiferum		100.0	0.40	MAN 1			
43214000	**Very good identification**						**43214000**
					GRA	SPO	
Fusobacterium mortiferum		98.10	0.62	XYL 7	1	1	
Actinomyces meyeri		0.90	0.18		99	1	
Clostridium glycolicum		0.58	0.09		99	99	
43214002	**Good identification**						**43214002**
Fusobacterium mortiferum		100.0	0.45	XYL 7			
43214100	**Very good identification**						**43214100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		98.45	0.68	XYL 7	1	
Actinomyces meyeri		1.55	0.28		99	
43214102	**Good identification**					**43214102**
					GRA	
Fusobacterium mortiferum		99.56	0.52	XYL 7	1	
Actinomyces meyeri		0.44	0.02	BGL 10	99	
43215000	**Low discrimination**					**43215000**
					GRA	
Fusobacterium mortiferum		89.03	0.65	XYL 7	1	
Eubacterium tortuosum		10.01	0.32		99	
Actinomyces meyeri		0.82	0.21		99	
43215002	**Low discrimination**					**43215002**
					GRA	
Fusobacterium mortiferum		84.38	0.48	XYL 7	1	
Eubacterium tortuosum		15.07	0.19	BGL 10	99	
Lactobacillus fermentum		0.54	0.12		99	
43215100	**Very good identification**					**43215100**
					GRA	
Fusobacterium mortiferum		95.65	0.71	XYL 7	1	
Eubacterium contortum		2.66	0.32		99	
Actinomyces meyeri		1.51	0.31		99	
43215101	**Good identification**					**43215101**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
43215102	**Acceptable identification**					**43215102**
					GRA	
Fusobacterium mortiferum		93.40	0.55	XYL 7	1	
Eubacterium contortum		4.13	0.19	BGL 10	99	
Bifidobacterium longum subsp. infantis		1.10	0.04		99	
43215110	**Acceptable identification**					**43215110**
Fusobacterium mortiferum		93.48	0.37	XYL 7		
Eubacterium saburreum		6.52	0.15	MLZ 1		
43215120	**Low discrimination**					**43215120**
					GRA	
Fusobacterium mortiferum		58.83	0.37	XYL 7	1	
Eubacterium contortum		28.63	0.19	NAG 1	99	
Actinomyces meyeri		12.54	0.16		99	
43215140	**Good identification**					**43215140**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
43215300	**Very good identification**					**43215300**
					GRA	
Eubacterium contortum		99.64	0.66	MLT 99	99	
Fusobacterium mortiferum		0.37	0.37		1	
43215302	**Good identification**					**43215302**
					GRA	
Eubacterium contortum		99.77	0.53	MLT 99	99	
Fusobacterium mortiferum		0.23	0.20		1	
43215320	**Good identification**					**43215320**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Eubacterium contortum		100.0	0.53	MLT 99			
43215322	**Good identification**						**43215322**
					GRA		
Eubacterium contortum		99.11	0.40	MLT 99	99		
Parabacteroides distasonis		0.89	0.16		1		
43215700	**Very good identification**						**43215700**
Eubacterium contortum		100.0	0.64	MLT 99			
43215702	**Good identification**						**43215702**
Eubacterium contortum		100.0	0.52	MLT 99			
43215720	**Good identification**						**43215720**
					GRA		
Eubacterium contortum		99.86	0.52	MLT 99	99		
Bacteroides vulgatus		0.14	-.05		1		
43215722	**Good identification**						**43215722**
					GRA		
Eubacterium contortum		99.33	0.39	MLT 99	99		
Parabacteroides distasonis		0.67	0.13		1		
43217100	**Acceptable identification**						**43217100**
					GRA		
Fusobacterium mortiferum		95.00	0.37	MAN 1	1		
Actinomyces israelii		5.00	0.17	XYL 7	99		
43220000	**Low discrimination**						**43220000**
					GRA	SPO	
Clostridium botulinum biovar A		45.36	0.36	SUC 7	99	99	
Eubacterium tortuosum		39.33	0.32	SAL 1	99	1	
Fusobacterium mortiferum		9.34	0.38		1	1	
43220100	**Good identification**						**43220100**
					GRA	SPO	
Fusobacterium mortiferum		96.22	0.44	FRU 99	1	1	
Clostridium botulinum biovar A		2.02	0.02		99	99	
Eubacterium tortuosum		1.75	-.03		99	1	
43221000	**Very good identification**						**43221000**
					GRA	SPO	
Eubacterium tortuosum		99.63	0.66	TRE 1	99	1	
Fusobacterium mortiferum		0.36	0.41		1	1	
Clostridium botulinum biovar A		0.01	0.02		99	99	
43221002	**Good identification**						**43221002**
					GRA		
Eubacterium tortuosum		99.77	0.53	TRE 1	99		
Fusobacterium mortiferum		0.23	0.24		1		
43221020	**Good identification**						**43221020**
Eubacterium tortuosum		100.0	0.53	TRE 1			
43221022	**Good identification**						**43221022**
Eubacterium tortuosum		100.0	0.40	TRE 1			
43230000	**Very good identification**						**43230000**
					GRA	SPO	
Fusobacterium mortiferum		99.44	0.72		1	1	
Clostridium botulinum biovar A		0.31	0.16		99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Clostridium sporogenes	0.11	0.06		99	99	
43230001	**Good identification**						**43230001**
	Fusobacterium mortiferum	100.0	0.38	URE 1			
43230002	**Good identification**						**43230002**
					GRA	SPO	
	Fusobacterium mortiferum	99.66	0.56	BGL 10	1	1	
	Clostridium sporogenes	0.34	-.02		99	99	
43230010	**Low discrimination**						**43230010**
					GRA	SPO	
	Fusobacterium mortiferum	68.87	0.38	MLZ 1	1	1	
	Clostridium difficile	31.13	0.16		99	99	
43230020	**Good identification**						**43230020**
	Fusobacterium mortiferum	100.0	0.38	NAG 1			
43230040	**Acceptable identification**						**43230040**
					GRA	SPO	
	Fusobacterium mortiferum	91.53	0.38	SOR 1	1	1	
	Clostridium botulinum biovar A	8.47	0.07		99	99	
43230100	**Very good identification**						**43230100**
	Fusobacterium mortiferum	100.0	0.78				
43230101	**Good identification**						**43230101**
	Fusobacterium mortiferum	100.0	0.44	URE 1			
43230102	**Very good identification**						**43230102**
	Fusobacterium mortiferum	100.0	0.62	BGL 10			
43230110	**Good identification**						**43230110**
	Fusobacterium mortiferum	100.0	0.44	MLZ 1			
43230120	**Good identification**						**43230120**
	Fusobacterium mortiferum	100.0	0.44	NAG 1			
43230140	**Good identification**						**43230140**
	Fusobacterium mortiferum	100.0	0.44	SOR 1			
43230200	**Good identification**						**43230200**
	Fusobacterium mortiferum	100.0	0.38	RHA 1			
43230300	**Good identification**						**43230300**
	Fusobacterium mortiferum	100.0	0.44	RHA 1			
43230400	**Good identification**						**43230400**
	Fusobacterium mortiferum	100.0	0.38	ARA 1			
43230500	**Good identification**						**43230500**
	Fusobacterium mortiferum	100.0	0.44	ARA 1			
43231000	**Very good identification**						**43231000**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		96.92	0.75		1		
Eubacterium tortuosum		2.75	0.32		99		
Lactobacillus jensenii		0.31	0.22		99		
43231001	**Good identification**						**43231001**
					GRA		
Fusobacterium mortiferum		97.24	0.41	URE 1	1		
Eubacterium tortuosum		2.76	-.03		99		
43231002	**Good identification**						**43231002**
					GRA		
Fusobacterium mortiferum		94.00	0.59	BGL 10	1		
Eubacterium tortuosum		4.23	0.19		99		
Lactobacillus jensenii		1.77	0.19		99		
43231010	**Low discrimination**						**43231010**
					GRA		
Fusobacterium mortiferum		86.67	0.41	MLZ 1	1		
Lactobacillus jensenii		8.99	0.14		99		
Eubacterium tortuosum		2.46	-.03		99		
43231020	**Low discrimination**						**43231020**
					GRA		
Fusobacterium mortiferum		63.58	0.41	NAG 1	1		
Eubacterium tortuosum		31.48	0.19		99		
Lactobacillus jensenii		4.94	0.12		99		
43231040	**Good identification**						**43231040**
					GRA		
Fusobacterium mortiferum		97.24	0.41	SOR 1	1		
Eubacterium tortuosum		2.76	-.03		99		
43231100	**Very good identification**						**43231100**
					COC	GRA	
Fusobacterium mortiferum		99.82	0.81		1	1	
Atopobium parvulum		0.14	0.21		99	99	
Eubacterium saburreum		0.03	0.18		1	85	
43231101	**Good identification**						**43231101**
					GRA		
Fusobacterium mortiferum		98.96	0.47	URE 1	1		
Actinomyces naeslundii		1.04	0.25		99		
43231102	**Very good identification**						**43231102**
					COC	GRA	
Fusobacterium mortiferum		99.60	0.65	BGL 10	1	1	
Atopobium parvulum		0.22	0.08		99	99	
Lactobacillus acidophilus		0.18	0.05		1	99	
43231110	**Good identification**						**43231110**
Fusobacterium mortiferum		97.87	0.47	MLZ 1			
Eubacterium saburreum		2.13	0.17				
43231120	**Acceptable identification**						**43231120**
					GRA	SPO	
Fusobacterium mortiferum		92.20	0.47	NAG 1	1	1	
Clostridium perfringens		3.10	0.11		99	84	
Clostridium septicum		2.48	0.03		99	99	
43231140	**Good identification**						**43231140**
Fusobacterium mortiferum		100.0	0.47	SOR 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43231200	**Good identification**						**43231200**
					GRA		
Fusobacterium mortiferum		97.24	0.41	RHA 1	1		
Eubacterium tortuosum		2.76	-.03		99		
43231300	**Good identification**						**43231300**
Fusobacterium mortiferum		100.0	0.47	RHA 1			
43231400	**Good identification**						**43231400**
					GRA		
Fusobacterium mortiferum		97.24	0.41	ARA 1	1		
Eubacterium tortuosum		2.76	-.03		99		
43231500	**Good identification**						**43231500**
Fusobacterium mortiferum		100.0	0.47	ARA 1			
43232000	**Low discrimination**						**43232000**
					GRA	SPO	
Fusobacterium mortiferum		53.62	0.38	MAN 1	1	1	
Clostridium difficile		44.66	0.21		99	99	
Eubacterium limosum		1.72	-.06		99	1	
43232100	**Good identification**						**43232100**
Fusobacterium mortiferum		100.0	0.44	MAN 1			
43233000	**Low discrimination**						**43233000**
					GRA	SPO	
Fusobacterium mortiferum		84.88	0.41	MAN 1	1	1	
Clostridium innocuum		12.71	0.14		99	99	
Eubacterium tortuosum		2.41	-.03		99	1	
43233100	**Good identification**						**43233100**
					GRA		
Fusobacterium mortiferum		98.65	0.47	MAN 1	1		
Actinomyces naeslundii		1.35	0.27		99		
43234000	**Good identification**						**43234000**
Fusobacterium mortiferum		100.0	0.53	XYL 7			
43234002	**Good identification**						**43234002**
Fusobacterium mortiferum		100.0	0.36	XYL 7			
43234100	**Very good identification**						**43234100**
Fusobacterium mortiferum		100.0	0.59	XYL 7			
43234102	**Good identification**						**43234102**
Fusobacterium mortiferum		100.0	0.43	XYL 7			
43235000	**Good identification**						**43235000**
					GRA		
Fusobacterium mortiferum		99.62	0.56	XYL 7	1		
Eubacterium tortuosum		0.38	-.03		99		
43235002	**Good identification**						**43235002**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
43235100	**Very good identification**						**43235100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Fusobacterium mortiferum		96.42	0.62	XYL 7	1	1	
Atopobium parvulum		3.29	0.25		99	99	
Eubacterium saburreum		0.19	0.14		1	85	
43235102	**Acceptable identification**						**43235102**
					COC	GRA	
Fusobacterium mortiferum		92.34	0.46	XYL 7	1	1	
Atopobium parvulum		5.00	0.13	BGL 10	99	99	
Actinomyces israelii		1.47	0.17		1	99	
43240000	**Good identification**						**43240000**
					GRA	SPO	
Eubacterium tortuosum		97.24	0.50	CEL 10	99	1	
Fusobacterium mortiferum		2.34	0.39	GAL 99	1	1	
Clostridium botulinum biovar A		0.41	0.13		99	99	
43240002	**Good identification**						**43240002**
					GRA		
Eubacterium tortuosum		98.51	0.37	CEL 10	99		
Fusobacterium mortiferum		1.50	0.22	GAL 99	1		
43240020	**Good identification**						**43240020**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43240100	**Low discrimination**						**43240100**
					GRA		
Fusobacterium mortiferum		84.78	0.45	FRU 99	1		
Eubacterium tortuosum		15.23	0.15		99		
43241000	**Very good identification**						**43241000**
					GRA		
Eubacterium tortuosum		99.96	0.84	CEL 10	99		
Fusobacterium mortiferum		0.04	0.42		1		
Lactobacillus catenaformis		0.00	0.12		99		
43241001	**Good identification**						**43241001**
Eubacterium tortuosum		100.0	0.50	CEL 10			
43241002	**Very good identification**						**43241002**
					GRA		
Eubacterium tortuosum		99.95	0.71	CEL 10	99		
Fusobacterium mortiferum		0.02	0.25		1		
Lactobacillus catenaformis		0.02	0.12		99		
43241003	**Good identification**						**43241003**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241010	**Good identification**						**43241010**
Eubacterium tortuosum		100.0	0.50	CEL 10			
43241012	**Good identification**						**43241012**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241020	**Very good identification**						**43241020**
					SPO		
Eubacterium tortuosum		99.99	0.71	CEL 10	1		
Clostridium barati		0.01	0.04		99		
Lactobacillus catenaformis		0.00	0.02		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
43241021	**Good identification**						**43241021**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241022	**Very good identification**						**43241022**
					SPO		
Eubacterium tortuosum		99.82	0.58	CEL 10	1		
Clostridium barati		0.16	0.12		99		
Lactobacillus catenaformis		0.02	0.02		1		
43241030	**Good identification**						**43241030**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241040	**Good identification**						**43241040**
Eubacterium tortuosum		100.0	0.50	CEL 10			
43241042	**Good identification**						**43241042**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241060	**Good identification**						**43241060**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241100	**Acceptable identification**						**43241100**
					GRA		
Eubacterium tortuosum		91.51	0.50	CEL 10	99		
Fusobacterium mortiferum		7.72	0.48	LAC 1	1		
Lactobacillus catenaformis		0.37	0.15		99		
43241102	**Acceptable identification**						**43241102**
					GRA	SPO	
Eubacterium tortuosum		90.52	0.37	CEL 10	99	1	
Fusobacterium mortiferum		4.81	0.32	LAC 1	1	1	
Clostridium barati		2.59	0.12		99	99	
43241120	**Good identification**						**43241120**
					SPO		
Eubacterium tortuosum		95.45	0.37	CEL 10	1		
Clostridium barati		2.73	0.12	LAC 1	99		
Clostridium paraputrificum		0.82	-.03		99		
43241200	**Good identification**						**43241200**
Eubacterium tortuosum		100.0	0.50	CEL 10			
43241202	**Good identification**						**43241202**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241220	**Good identification**						**43241220**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241400	**Good identification**						**43241400**
Eubacterium tortuosum		100.0	0.50	CEL 10			
43241402	**Good identification**						**43241402**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43241420	**Good identification**						**43241420**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium tortuosum		100.0	0.37	CEL 10			
43243000	**Acceptable identification**						**43243000**
					SPO		
Eubacterium tortuosum		93.29	0.50	CEL 10	1		
Clostridium innocuum		6.71	0.34	MAN 1	99		
43243002	**Low discrimination**						**43243002**
					SPO		
Eubacterium tortuosum		71.04	0.37	CEL 10	1		
Clostridium innocuum		28.96	0.34	MAN 1	99		
43243020	**Good identification**						**43243020**
					SPO		
Eubacterium tortuosum		99.59	0.37	CEL 10	1		
Clostridium innocuum		0.41	0.00	MAN 1	99		
43245000	**Good identification**						**43245000**
					GRA		
Eubacterium tortuosum		99.73	0.50	CEL 10	99		
Fusobacterium mortiferum		0.27	0.22	XYL 1	1		
43245002	**Good identification**						**43245002**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43245020	**Good identification**						**43245020**
Eubacterium tortuosum		100.0	0.37	CEL 10			
43250000	**Very good identification**						**43250000**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.73		1	1	
Eubacterium tortuosum		0.42	0.15		99	1	
Clostridium barati		0.12	0.08		99	99	
43250001	**Good identification**						**43250001**
Fusobacterium mortiferum		100.0	0.39	URE 1			
43250002	**Good identification**						**43250002**
					GRA	SPO	
Fusobacterium mortiferum		94.99	0.56	BGL 10	1	1	
Clostridium barati		3.14	0.16		99	99	
Lactobacillus catenaformis		0.97	0.12		99	1	
43250010	**Low discrimination**						**43250010**
					GRA	SPO	
Fusobacterium mortiferum		77.93	0.39	MLZ 1	1	1	
Clostridium difficile		22.07	0.13		99	99	
43250020	**Low discrimination**						**43250020**
					GRA	SPO	
Fusobacterium mortiferum		68.23	0.39	NAG 1	1	1	
Clostridium barati		24.80	0.16		99	99	
Eubacterium tortuosum		5.05	0.03		99	1	
43250040	**Good identification**						**43250040**
Fusobacterium mortiferum		100.0	0.39	SOR 1			
43250100	**Very good identification**						**43250100**
					GRA	SPO	
Fusobacterium mortiferum		99.72	0.79		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		0.16	0.16		99	99	
Lactobacillus catenaformis		0.07	0.15		99	1	
43250101	**Good identification**						**43250101**
Fusobacterium mortiferum		100.0	0.45	URE 1			
43250102	**Very good identification**						**43250102**
					GRA	SPO	
Fusobacterium mortiferum		95.32	0.63	BGL 10	1	1	
Clostridium barati		4.05	0.24		99	99	
Lactobacillus catenaformis		0.63	0.15		99	1	
43250110	**Good identification**						**43250110**
Fusobacterium mortiferum		100.0	0.45	MLZ 1			
43250120	**Low discrimination**						**43250120**
					GRA	SPO	
Fusobacterium mortiferum		66.13	0.45	NAG 1	1	1	
Clostridium barati		30.91	0.24		99	99	
Clostridium paraputrificum		1.77	-.03		99	99	
43250140	**Good identification**						**43250140**
					GRA	SPO	
Fusobacterium mortiferum		98.66	0.45	SOR 1	1	1	
Clostridium barati		1.34	-.02		99	99	
43250200	**Good identification**						**43250200**
Fusobacterium mortiferum		100.0	0.39	RHA 1			
43250300	**Good identification**						**43250300**
					GRA	SPO	
Fusobacterium mortiferum		98.66	0.45	RHA 1	1	1	
Clostridium barati		1.34	-.02		99	99	
43250400	**Good identification**						**43250400**
Fusobacterium mortiferum		100.0	0.39	ARA 1			
43250500	**Good identification**						**43250500**
Fusobacterium mortiferum		100.0	0.45	ARA 1			
43251000	**Low discrimination**						**43251000**
					GRA		
Fusobacterium mortiferum		71.82	0.76		1		
Eubacterium tortuosum		20.07	0.50		99		
Lactobacillus catenaformis		5.39	0.47		99		
43251001	**Low discrimination**						**43251001**
					GRA		
Fusobacterium mortiferum		72.88	0.42	URE 1	1		
Eubacterium tortuosum		20.36	0.15		99		
Lactobacillus catenaformis		5.47	0.12		99		
43251002	**Low discrimination**						**43251002**
					GRA		
Fusobacterium mortiferum		37.20	0.59	BGL 10	1		
Lactobacillus catenaformis		25.10	0.47		99		
Eubacterium tortuosum		16.50	0.37		99		
43251010	**Low discrimination**						**43251010**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		51.05	0.42	MLZ 1	1		
Lactobacillus jensenii		30.05	0.28		99		
Eubacterium tortuosum		14.26	0.15		99		
43251020	**Low discrimination**						**43251020**
					SPO		
Eubacterium tortuosum		38.06	0.37	FRU 1	1		
Clostridium barati		35.90	0.38	CEL 10	99		
Lactobacillus catenaformis		14.47	0.36		1		
43251022	**Low discrimination**						**43251022**
					SPO		
Clostridium barati		80.97	0.46	MNS 99	99		
Lactobacillus catenaformis		10.88	0.36		1		
Eubacterium tortuosum		5.05	0.24		1		
43251040	**Low discrimination**						**43251040**
					GRA	SPO	
Fusobacterium mortiferum		66.42	0.42	SOR 1	1	1	
Eubacterium tortuosum		18.56	0.15		99	1	
Clostridium barati		8.86	0.11		99	99	
43251100	**Good identification**						**43251100**
					GRA		
Fusobacterium mortiferum		90.79	0.82		1		
Lactobacillus catenaformis		4.38	0.50		99		
Collinsella aerofaciens		2.90	0.32		99		
43251101	**Acceptable identification**						**43251101**
					GRA		
Fusobacterium mortiferum		90.91	0.48	URE 1	1		
Lactobacillus catenaformis		4.38	0.15		99		
Collinsella aerofaciens		2.90	-.03		99		
43251102	**Low discrimination**						**43251102**
					GRA	SPO	
Fusobacterium mortiferum		48.69	0.66	BGL 10	1	1	
Clostridium barati		26.20	0.46		99	99	
Lactobacillus catenaformis		21.13	0.50		99	1	
43251110	**Acceptable identification**						**43251110**
					GRA		
Fusobacterium mortiferum		90.91	0.48	MLZ 1	1		
Lactobacillus catenaformis		4.38	0.15		99		
Collinsella aerofaciens		2.90	-.03		99		
43251120	**Low discrimination**						**43251120**
					SPO		
Clostridium barati		57.60	0.46	MNS 99	99		
Clostridium paraputrificum		17.20	0.32		99		
Lactobacillus catenaformis		11.61	0.39		1		
43251122	**Low discrimination**						**43251122**
					SPO		
Clostridium barati		86.82	0.54	MNS 99	99		
Lactobacillus catenaformis		5.83	0.39		1		
Clostridium paraputrificum		3.70	0.25		99		
43251140	**Low discrimination**						**43251140**
					GRA	SPO	
Fusobacterium mortiferum		79.89	0.48	SOR 1	1	1	
Clostridium barati		13.71	0.20		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		3.85	0.15		99	1	
43251162	**Acceptable identification**						**43251162**
					SPO		
Clostridium barati		91.56	0.36	MNS 99	99		
Lactobacillus acidophilus		3.82	0.15	SOR 8	1		
Clostridium paraputrificum		3.38	0.06		99		
43251200	**Low discrimination**						**43251200**
					GRA	SPO	
Fusobacterium mortiferum		66.42	0.42	RHA 1	1	1	
Eubacterium tortuosum		18.56	0.15		99	1	
Clostridium barati		8.86	0.11		99	99	
43251300	**Low discrimination**						**43251300**
					GRA	SPO	
Fusobacterium mortiferum		79.89	0.48	RHA 1	1	1	
Clostridium barati		13.71	0.20		99	99	
Lactobacillus catenaformis		3.85	0.15		99	1	
43251322	**Acceptable identification**						**43251322**
					GRA	SPO	
Clostridium barati		94.65	0.36	MNS 99	99	99	
Lactobacillus acidophilus		2.40	0.11	RHA 8	99	1	
Prevotella oralis		1.74	0.07		1	1	
43251400	**Low discrimination**						**43251400**
					GRA		
Fusobacterium mortiferum		69.46	0.42	ARA 1	1		
Eubacterium tortuosum		19.41	0.15		99		
Lactobacillus catenaformis		5.21	0.12		99		
43251500	**Acceptable identification**						**43251500**
					GRA		
Fusobacterium mortiferum		90.91	0.48	ARA 1	1		
Lactobacillus catenaformis		4.38	0.15		99		
Collinsella aerofaciens		2.90	-.03		99		
43252100	**Good identification**						**43252100**
					GRA	SPO	
Fusobacterium mortiferum		97.42	0.45	MAN 1	1	1	
Clostridium barati		1.32	-.02		99	99	
Clostridium innocuum		1.26	0.00		99	99	
43253000	**Good identification**						**43253000**
					GRA	SPO	
Clostridium innocuum		96.17	0.56	MNS 99	99	99	
Fusobacterium mortiferum		2.54	0.42		1	1	
Eubacterium tortuosum		0.71	0.15		99	1	
43253002	**Good identification**						**43253002**
					GRA	SPO	
Clostridium innocuum		98.35	0.56	MNS 99	99	99	
Clostridium barati		1.04	0.20		99	99	
Fusobacterium mortiferum		0.29	0.25		1	1	
43253100	**Low discrimination**						**43253100**
					GRA	SPO	
Fusobacterium mortiferum		69.57	0.48	MAN 1	1	1	
Clostridium barati		11.94	0.20		99	99	
Clostridium innocuum		11.39	0.22		99	99	
43253122	**Acceptable identification**						**43253122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium barati		93.81	0.36	MNS 99	99		
Lactobacillus acidophilus		1.67	0.08	MAN 8	1		
Clostridium tertium		1.57	0.09		99		
43253400	**Good identification**						**43253400**
Clostridium innocuum		100.0	0.43	MNS 99			
43253402	**Good identification**						**43253402**
Clostridium innocuum		100.0	0.43	MNS 99			
43254000	**Good identification**						**43254000**
Fusobacterium mortiferum		100.0	0.53	XYL 7			
43254002	**Good identification**						**43254002**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
43254100	**Very good identification**						**43254100**
Fusobacterium mortiferum		100.0	0.60	XYL 7			
43254102	**Good identification**						**43254102**
Fusobacterium mortiferum		100.0	0.43	XYL 7			
43255000	**Good identification**						**43255000**
					GRA		
Fusobacterium mortiferum		94.80	0.56	XYL 7	1		
Eubacterium tortuosum		3.55	0.15		99		
Lactobacillus catenaformis		0.95	0.12		99		
43255002	**Low discrimination**						**43255002**
					GRA		
Fusobacterium mortiferum		79.36	0.40	XYL 7	1		
Lactobacillus catenaformis		7.19	0.12	BGL 10	99		
Eubacterium tortuosum		4.73	0.03		99		
43255100	**Very good identification**						**43255100**
					GRA		
Fusobacterium mortiferum		98.31	0.63	XYL 7	1		
Lactobacillus catenaformis		0.64	0.15		99		
Collinsella aerofaciens		0.42	-.03		99		
43255102	**Low discrimination**						**43255102**
					GRA	SPO	
Fusobacterium mortiferum		82.13	0.46	XYL 7	1	1	
Clostridium barati		5.93	0.12	BGL 10	99	99	
Lactobacillus catenaformis		4.78	0.15		99	1	
43257100	**Low discrimination**						**43257100**
					GRA	SPO	
Actinomyces israelii		65.31	0.37	MLT 99	99	1	
Fusobacterium mortiferum		31.13	0.29		1	1	
Clostridium innocuum		3.56	0.00		99	99	
43257102	**Low discrimination**						**43257102**
					SPO		
Actinomyces israelii		81.51	0.40	MLT 99	1		
Bifidobacterium adolescentis		8.55	0.16		1		
Clostridium tertium		4.10	0.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43257500	**Good identification**						**43257500**
Actinomyces israelii		100.0	0.35	MLT 99			
43257502	**Low discrimination**						**43257502**
Actinomyces israelii		53.83	0.38	MLT 99			
Bifidobacterium adolescentis		46.18	0.31				
43260100	**Good identification**						**43260100**
Fusobacterium mortiferum		100.0	0.36	FRU 99			
43261000	**Good identification**						**43261000**
					GRA		
Eubacterium tortuosum		98.74	0.50	TRE 1	99		
Fusobacterium mortiferum		1.07	0.33	CEL 10	1		
Lactobacillus jensenii		0.19	0.10		99		
43261002	**Good identification**						**43261002**
					GRA		
Eubacterium tortuosum		98.61	0.37	TRE 1	99		
Lactobacillus jensenii		0.72	0.07	CEL 10	99		
Fusobacterium mortiferum		0.67	0.16		1		
43261020	**Good identification**						**43261020**
Eubacterium tortuosum		100.0	0.37	TRE 1			
43261100	**Low discrimination**						**43261100**
					GRA		
Fusobacterium mortiferum		71.39	0.39	FRU 99	1		
Eubacterium tortuosum		28.61	0.15		99		
43270000	**Very good identification**						**43270000**
					GRA		
Fusobacterium mortiferum		99.73	0.64		1		
Lactobacillus jensenii		0.27	0.10		99		
43270002	**Good identification**						**43270002**
					GRA	SPO	
Fusobacterium mortiferum		97.48	0.47	BGL 10	1	1	
Lactobacillus jensenii		1.59	0.07		99	1	
Clostridium barati		0.94	-.02		99	99	
43270100	**Very good identification**						**43270100**
					GRA	SPO	
Fusobacterium mortiferum		99.95	0.70		1	1	
Clostridium barati		0.05	-.02		99	99	
43270101	**Good identification**						**43270101**
Fusobacterium mortiferum		100.0	0.36	URE 1			
43270102	**Good identification**						**43270102**
					GRA	SPO	
Fusobacterium mortiferum		98.00	0.54	BGL 10	1	1	
Clostridium barati		1.21	0.06		99	99	
Lactobacillus acidophilus		0.79	0.05		99	1	
43270110	**Good identification**						**43270110**
Fusobacterium mortiferum		100.0	0.36	MLZ 1			
43270120	**Low discrimination**						**43270120**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Fusobacterium mortiferum		88.02	0.36	NAG 1	1	1		
Clostridium barati		11.98	0.06		99	99		
43270140	**Good identification**							**43270140**
Fusobacterium mortiferum		100.0	0.36	SOR 1				
43270300	**Good identification**							**43270300**
Fusobacterium mortiferum		100.0	0.36	RHA 1				
43270500	**Good identification**							**43270500**
Fusobacterium mortiferum		100.0	0.36	ARA 1				
43271000	**Low discrimination**							**43271000**
					GRA			
Fusobacterium mortiferum		83.32	0.67		1			
Lactobacillus jensenii		14.93	0.44		99			
Eubacterium tortuosum		0.79	0.15		99			
43271002	**Low discrimination**							**43271002**
					GRA	SPO		
Lactobacillus jensenii		45.10	0.41	MNS 99	99	1		
Fusobacterium mortiferum		41.97	0.50	MLT 95	1	1		
Clostridium barati		5.11	0.20		99	99		
43271010	**Low discrimination**							**43271010**
					GRA			
Lactobacillus jensenii		85.53	0.36	MNS 99	99			
Fusobacterium mortiferum		14.47	0.33	MLT 95	1			
43271100	**Very good identification**							**43271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.03	0.73		1	1	1	
Atopobium parvulum		0.95	0.27		99	99	1	
Clostridium barati		0.57	0.20		1	99	99	
43271101	**Good identification**							**43271101**
					GRA			
Fusobacterium mortiferum		95.98	0.39	URE 1	1			
Actinomyces naeslundii		4.02	0.27		99			
43271102	**Low discrimination**							**43271102**
					GRA	SPO		
Fusobacterium mortiferum		57.60	0.57	BGL 10	1	1		
Lactobacillus acidophilus		30.57	0.39		99	1		
Clostridium barati		9.02	0.28		99	99		
43271110	**Acceptable identification**							**43271110**
					GRA			
Fusobacterium mortiferum		90.38	0.39	MLZ 1	1			
Leptotrichia buccalis		7.31	-.03		1			
Lactobacillus jensenii		2.31	0.02		99			
43271122	**Low discrimination**							**43271122**
					SPO			
Lactobacillus acidophilus		48.65	0.39	MNS 99	1			
Clostridium barati		43.09	0.36	MLT 94	99			
Clostridium paraputrificum		5.62	0.16		99			
43271140	**Good identification**							**43271140**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		95.24	0.39	SOR 1	1	1	
Clostridium barati		4.76	0.01		99	99	
43271300	**Good identification**						**43271300**
					GRA	SPO	
Fusobacterium mortiferum		95.24	0.39	RHA 1	1	1	
Clostridium barati		4.76	0.01		99	99	
43271500	**Good identification**						**43271500**
					GRA		
Fusobacterium mortiferum		97.56	0.39	ARA 1	1		
Actinomyces israelii		2.44	0.14		99		
43272100	**Good identification**						**43272100**
Fusobacterium mortiferum		100.0	0.36	MAN 1			
43273000	**Good identification**						**43273000**
					GRA	SPO	
Clostridium innocuum		97.20	0.48	MNS 99	99	99	
Fusobacterium mortiferum		2.19	0.33		1	1	
Lactobacillus jensenii		0.39	0.10		99	1	
43273002	**Good identification**						**43273002**
					SPO		
Clostridium innocuum		98.76	0.48	MNS 99	99		
Actinomyces israelii		0.34	0.21		1		
Lactobacillus jensenii		0.27	0.07		1		
43273100	**Low discrimination**						**43273100**
					GRA	SPO	
Fusobacterium mortiferum		64.25	0.39	MAN 1	1	1	
Clostridium innocuum		12.37	0.14		99	99	
Actinomyces israelii		11.11	0.28		99	1	
43273400	**Good identification**						**43273400**
					SPO		
Clostridium innocuum		98.96	0.35	MNS 99	99		
Actinomyces israelii		1.04	0.16		1		
43273402	**Good identification**						**43273402**
					SPO		
Clostridium innocuum		98.45	0.35	MNS 99	99		
Actinomyces israelii		1.55	0.19		1		
43274000	**Good identification**						**43274000**
Fusobacterium mortiferum		100.0	0.44	XYL 7			
43274100	**Good identification**						**43274100**
Fusobacterium mortiferum		100.0	0.51	XYL 7			
43275000	**Good identification**						**43275000**
					GRA		
Fusobacterium mortiferum		95.10	0.48	XYL 7	1		
Actinomyces israelii		2.61	0.23		99		
Lactobacillus jensenii		2.29	0.10		99		
43275100	**Low discrimination**						**43275100**
					COC	GRA	
Fusobacterium mortiferum		77.56	0.54	XYL 7	1	1	
Atopobium parvulum		18.50	0.32		99	99	
Actinomyces israelii		3.65	0.33		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43275102	**Low discrimination**						**43275102**
					COC	GRA	
Fusobacterium mortiferum		46.13	0.37	XYL 7	1	1	
Actinomyces israelii		29.33	0.36	BGL 10	1	99	
Atopobium parvulum		17.48	0.19		99	99	
43277100	**Acceptable identification**						**43277100**
					GRA	SPO	
Actinomyces israelii		91.07	0.46	MLT 99	99	1	
Clostridium tertium		5.48	0.11		99	99	
Fusobacterium mortiferum		3.45	0.20		1	1	
43277102	**Low discrimination**						**43277102**
					SPO		
Actinomyces israelii		87.80	0.49	MLT 99	1		
Clostridium tertium		10.57	0.19		99		
Bifidobacterium adolescentis		1.63	0.13		1		
43277140	**Good identification**						**43277140**
Actinomyces israelii		100.0	0.39	MLT 99			
43277142	**Good identification**						**43277142**
Actinomyces israelii		100.0	0.42	MLT 99			
43277402	**Low discrimination**						**43277402**
					SPO		
Actinomyces israelii		77.51	0.38	MLT 99	1		
Clostridium innocuum		22.49	0.13		99		
43277500	**Good identification**						**43277500**
					SPO		
Actinomyces israelii		98.34	0.45	MLT 99	1		
Clostridium butyricum		1.66	-.07		99		
43277502	**Low discrimination**						**43277502**
Actinomyces israelii		86.80	0.48	MLT 99			
Bifidobacterium adolescentis		13.20	0.28				
43277540	**Good identification**						**43277540**
Actinomyces israelii		100.0	0.38	MLT 99			
43277542	**Good identification**						**43277542**
Actinomyces israelii		95.39	0.41	MLT 99			
Bifidobacterium adolescentis		4.61	0.12				
43300000	**Low discrimination**						**43300000**
					SPO		
Clostridium botulinum biovar A		57.06	0.45	SUC 7	99		
Eubacterium tortuosum		15.12	0.32	SAL 1	1		
Clostridium sporogenes		12.78	0.32		99		
43300040	**Low discrimination**						**43300040**
Clostridium botulinum biovar A		85.78	0.36	SUC 7			
Clostridium novyi biovar A		9.76	0.32	SAL 1			
Clostridium bifermentans		2.12	0.08				
43300100	**Low discrimination**						**43300100**
					GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		76.57	0.37	MLT 11	1	1		
Clostridium botulinum biovar A		12.73	0.11	FRU 99	99	99		
Eubacterium tortuosum		3.37	-.03		99	1		
43301000	**Very good identification**							**43301000**
					GRA	SPO		
Eubacterium tortuosum		99.74	0.66	MLT 1	99	1		
Fusobacterium mortiferum		0.15	0.34		1	1		
Clostridium botulinum biovar A		0.04	0.11		99	99		
43301002	**Good identification**							**43301002**
					GRA			
Eubacterium tortuosum		99.91	0.53	MLT 1	99			
Fusobacterium mortiferum		0.09	0.18		1			
43301020	**Good identification**							**43301020**
Eubacterium tortuosum		100.0	0.53	MLT 1				
43301022	**Good identification**							**43301022**
Eubacterium tortuosum		100.0	0.40	MLT 1				
43301101	**Good identification**							**43301101**
					SPO			
Clostridium chauvoei		99.36	0.40	MNS 90	99			
Eubacterium tortuosum		0.65	-.03	SAL 1	1			
43305300	**Very good identification**							**43305300**
Eubacterium contortum		100.0	0.66	FRU 99				
43305302	**Good identification**							**43305302**
Eubacterium contortum		100.0	0.53	FRU 99				
43305320	**Good identification**							**43305320**
Eubacterium contortum		100.0	0.53	FRU 99				
43305322	**Good identification**							**43305322**
Eubacterium contortum		100.0	0.40	FRU 99				
43305700	**Very good identification**							**43305700**
Eubacterium contortum		100.0	0.64	FRU 99				
43305702	**Good identification**							**43305702**
Eubacterium contortum		100.0	0.52	FRU 99				
43305720	**Good identification**							**43305720**
					GRA			
Eubacterium contortum		99.86	0.52	FRU 99	99			
Bacteroides vulgatus		0.14	-.05		1			
43305722	**Good identification**							**43305722**
Eubacterium contortum		100.0	0.39	FRU 99				
43310000	**Good identification**							**43310000**
					COC	GRA	SPO	
Fusobacterium mortiferum		91.02	0.65	MLT 11	1	1	1	
Megasphaera elsdenii		3.25	0.32		99	1	1	
Clostridium botulinum biovar A		2.25	0.25		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43310002	**Low discrimination**						**43310002**
					GRA	SPO	
Fusobacterium mortiferum		87.13	0.49	MLT 11	1	1	
Actinomyces odontolyticus		5.75	0.23	BGL 10	99	1	
Clostridium sporogenes		4.01	0.11		99	99	
43310100	**Very good identification**						**43310100**
					GRA		
Fusobacterium mortiferum		99.16	0.72	MLT 11	1		
Actinomyces meyeri		0.55	0.24		99		
Actinomyces odontolyticus		0.16	0.18		99		
43310101	**Low discrimination**						**43310101**
					GRA	SPO	
Fusobacterium mortiferum		86.22	0.37	MLT 11	1	1	
Clostridium chauvoei		13.78	0.12	URE 1	99	99	
43310102	**Good identification**						**43310102**
					GRA		
Fusobacterium mortiferum		97.30	0.55	MLT 11	1		
Bifidobacterium longum subsp. infantis		1.28	0.06	BGL 10	99		
Actinomyces odontolyticus		0.87	0.15		99		
43310110	**Good identification**						**43310110**
Fusobacterium mortiferum		100.0	0.37	MLT 11			
43310120	**Low discrimination**						**43310120**
					GRA	SPO	
Fusobacterium mortiferum		75.98	0.37	MLT 11	1	1	
Actinomyces meyeri		5.73	0.09	NAG 1	99	1	
Clostridium paraputrificum		5.49	-.03		99	99	
43310140	**Good identification**						**43310140**
Fusobacterium mortiferum		100.0	0.37	MLT 11			
43310500	**Low discrimination**						**43310500**
					GRA		
Fusobacterium mortiferum		87.86	0.37	MLT 11	1		
Actinomyces meyeri		12.14	0.13	ARA 1	99		
43311000	**Low discrimination**						**43311000**
					GRA		
Fusobacterium mortiferum		86.59	0.68	MLT 11	1		
Eubacterium tortuosum		5.93	0.32		99		
Lactobacillus jensenii		4.17	0.36		99		
43311002	**Low discrimination**						**43311002**
					GRA		
Fusobacterium mortiferum		58.52	0.52	MLT 11	1		
Lactobacillus jensenii		16.89	0.33	BGL 10	99		
Lactobacillus catenaformis		9.68	0.29		99		
43311100	**Low discrimination**						**43311100**
					GRA		
Fusobacterium mortiferum		87.12	0.75	MLT 11	1		
Collinsella aerofaciens		7.49	0.32		99		
Eubacterium contortum		1.48	0.32		99		
43311101	**Low discrimination**						**43311101**
					GRA	SPO	
Clostridium chauvoei		89.09	0.47	MNS 90	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		8.45	0.40	SAL 1	1	1	
Actinomyces naeslundii		1.74	0.40		99	1	
43311102	**Low discrimination**						**43311102**
					GRA		
Fusobacterium mortiferum		42.91	0.58	MLT 11	1		
Bifidobacterium longum subsp. infantis		37.32	0.40	BGL 10	99		
Lactobacillus catenaformis		4.56	0.32		99		
43311110	**Low discrimination**						**43311110**
					GRA		
Fusobacterium mortiferum		71.35	0.40	MLT 11	1		
Eubacterium saburreum		22.51	0.30	MLZ 1	85		
Collinsella aerofaciens		6.14	-.03		99		
43311121	**Acceptable identification**						**43311121**
					SPO		
Clostridium chauvoei		91.55	0.38	MNS 90	99		
Actinomyces naeslundii		5.35	0.40	SAL 1	1		
Clostridium paraputrificum		1.25	-.03		99		
43311122	**Low discrimination**						**43311122**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		53.90	0.44	MNS 99	99	1	
Prevotella oralis		15.80	0.38	RAF 95	1	1	
Clostridium barati		7.49	0.32		99	99	
43311140	**Low discrimination**						**43311140**
					GRA		
Fusobacterium mortiferum		84.92	0.40	MLT 11	1		
Collinsella aerofaciens		7.31	-.03	SOR 1	99		
Clostridium perfringens		2.17	0.03		99		
43311300	**Very good identification**						**43311300**
					GRA		
Eubacterium contortum		99.26	0.66	XYL 99	99		
Fusobacterium mortiferum		0.60	0.40		1		
Actinomyces odontolyticus		0.08	0.20		99		
43311302	**Good identification**						**43311302**
					GRA		
Eubacterium contortum		98.41	0.53	XYL 99	99		
Prevotella oralis		0.38	0.10		1		
Fusobacterium mortiferum		0.37	0.24		1		
43311320	**Good identification**						**43311320**
					GRA	SPO	
Eubacterium contortum		98.91	0.53	XYL 99	99	1	
Prevotella oralis		0.38	0.10		1	1	
Clostridium barati		0.23	0.05		99	99	
43311322	**Low discrimination**						**43311322**
					GRA	SPO	
Eubacterium contortum		66.95	0.40	XYL 99	99	1	
Prevotella oralis		27.64	0.32		1	1	
Clostridium barati		2.66	0.14		99	99	
43311502	**Low discrimination**						**43311502**
Bifidobacterium adolescentis		84.59	0.47	MNS 87			
Eubacterium contortum		5.38	0.17	RAF 92			
Lactobacillus fermentum		3.41	0.28	XYL 87			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43311512	**Good identification**					**43311512**
Bifidobacterium adolescentis		99.47	0.48	MNS 87		
Bifidobacterium longum subsp. longum		0.53	0.07	RAF 92		
43311700	**Very good identification**					**43311700**
Eubacterium contortum		100.0	0.64	XYL 99		
43311702	**Good identification**					**43311702**
Eubacterium contortum		99.84	0.52	XYL 99		
Bifidobacterium adolescentis		0.16	0.12			
43311720	**Good identification**					**43311720**
					GRA	
Eubacterium contortum		99.71	0.52	XYL 99	99	
Bacteroides vulgatus		0.29	0.01		1	
43311722	**Good identification**					**43311722**
					GRA	
Eubacterium contortum		99.49	0.39	XYL 99	99	
Prevotella oralis		0.51	-.02		1	
43312100	**Good identification**					**43312100**
					GRA	
Fusobacterium mortiferum		95.99	0.37	MLT 11	1	
Propionibacterium propionicum		4.01	0.15	MAN 1	99	
43313001	**Good identification**					**43313001**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43313003	**Good identification**					**43313003**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43313021	**Good identification**					**43313021**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43313023	**Good identification**					**43313023**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43313100	**Low discrimination**					**43313100**
					GRA	
Fusobacterium mortiferum		59.57	0.40	MLT 11	1	
Actinomyces israelii		16.44	0.33	MAN 1	99	
Actinomyces naeslundii		15.85	0.42		99	
43313101	**Low discrimination**					**43313101**
					SPO	
Actinomyces naeslundii		78.93	0.45	ESL 12	1	
Clostridium chauvoei		21.07	0.12	MNS 88	99	
43313103	**Good identification**					**43313103**
					SPO	
Actinomyces naeslundii		95.50	0.45	ESL 12	1	
Clostridium chauvoei		4.50	-.01	MNS 88	99	
43313120	**Low discrimination**					**43313120**
					SPO	
Actinomyces naeslundii		55.80	0.42	ESL 12	1	
Clostridium barati		14.36	0.05	MNS 88	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium paraputrificum		10.10	-.03	RAF 92	99	
43313121	**Acceptable identification**					**43313121**
					SPO	
Actinomyces naeslundii		91.83	0.45	ESL 12	1	
Clostridium chauvoei		8.17	0.04	MNS 88	99	
43313122	**Low discrimination**					**43313122**
					SPO	
Actinomyces naeslundii		30.01	0.42	ESL 12	1	
Clostridium barati		23.17	0.14	MNS 88	99	
Bifidobacterium longum subsp. infantis		19.38	0.10	RAF 92	1	
43313123	**Good identification**					**43313123**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43313502	**Acceptable identification**					**43313502**
Bifidobacterium adolescentis		90.21	0.45	MNS 87		
Actinomyces israelii		9.79	0.35	RAF 92		
43313512	**Good identification**					**43313512**
Bifidobacterium adolescentis		98.69	0.47	MNS 87		
Actinomyces israelii		1.31	0.21	RAF 92		
43314000	**Low discrimination**					**43314000**
					GRA	
Fusobacterium mortiferum		82.39	0.46	MLT 11	1	
Actinomyces meyeri		6.92	0.18	XYL 7	99	
Actinomyces odontolyticus		6.45	0.22		99	
43314100	**Low discrimination**					**43314100**
					GRA	
Fusobacterium mortiferum		67.00	0.52	MLT 11	1	
Eubacterium contortum		22.65	0.32	XYL 7	99	
Actinomyces meyeri		9.65	0.29		99	
43314102	**Low discrimination**					**43314102**
					GRA	
Fusobacterium mortiferum		54.10	0.36	MLT 11	1	
Eubacterium contortum		29.04	0.19	XYL 7	99	
Bifidobacterium longum subsp. infantis		7.75	0.04	BGL 10	99	
43314300	**Very good identification**					**43314300**
					GRA	
Eubacterium contortum		99.91	0.66	GAL 99	99	
Actinomyces odontolyticus		0.06	0.18		99	
Fusobacterium mortiferum		0.03	0.18		1	
43314302	**Good identification**					**43314302**
Eubacterium contortum		99.79	0.53	GAL 99		
Actinomyces odontolyticus		0.21	0.14			
43314320	**Good identification**					**43314320**
Eubacterium contortum		100.0	0.53	GAL 99		
43314322	**Good identification**					**43314322**
Eubacterium contortum		100.0	0.40	GAL 99		
43314700	**Very good identification**					**43314700**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		100.0	0.64	GAL 99		
43314702	**Good identification**					**43314702**
Eubacterium contortum		100.0	0.52	GAL 99		
43314720	**Good identification**					**43314720**
					GRA	
Eubacterium contortum		99.86	0.52	GAL 99	99	
Bacteroides vulgatus		0.14	-.05		1	
43314722	**Good identification**					**43314722**
Eubacterium contortum		100.0	0.39	GAL 99		
43315000	**Low discrimination**					**43315000**
					GRA	
Fusobacterium mortiferum		57.17	0.49	MLT 11	1	
Eubacterium contortum		30.06	0.32	XYL 7	99	
Actinomyces meyeri		4.80	0.21		99	
43315100	**Good identification**					**43315100**
					GRA	
Eubacterium contortum		93.83	0.66	RHA 99	99	
Fusobacterium mortiferum		4.21	0.55		1	
Actinomyces meyeri		0.61	0.32		99	
43315102	**Low discrimination**					**43315102**
Eubacterium contortum		68.20	0.53	RHA 99		
Bifidobacterium longum subsp. infantis		18.20	0.38			
Bifidobacterium adolescentis		8.87	0.47			
43315112	**Low discrimination**					**43315112**
Bifidobacterium adolescentis		89.78	0.48	MNS 87		
Eubacterium contortum		5.71	0.19	RAF 92		
Actinomyces israelii		2.42	0.27	ARA 87		
43315120	**Acceptable identification**					**43315120**
Eubacterium contortum		93.52	0.53	RHA 99		
Bifidobacterium longum subsp. infantis		4.39	0.25			
Actinomyces meyeri		0.47	0.17			
43315122	**Low discrimination**					**43315122**
					GRA	
Bifidobacterium longum subsp. infantis		68.07	0.43	MNS 99	99	
Eubacterium contortum		25.33	0.40	RAF 95	99	
Prevotella oralis		3.02	0.23		1	
43315200	**Very good identification**					**43315200**
					GRA	
Eubacterium contortum		99.85	0.66	LAC 99	99	
Actinomyces odontolyticus		0.13	0.23		99	
Fusobacterium mortiferum		0.02	0.15		1	
43315202	**Good identification**					**43315202**
Eubacterium contortum		99.55	0.53	LAC 99		
Actinomyces odontolyticus		0.45	0.20			
43315220	**Good identification**					**43315220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		99.88	0.53	LAC 99		
Actinomyces odontolyticus		0.12	0.10			
43315222	**Good identification**					**43315222**
Eubacterium contortum		99.59	0.40	LAC 99		
Actinomyces odontolyticus		0.41	0.06			
43315300	**Excellent identification**					**43315300**
					GRA	
Eubacterium contortum		100.0	1.00		99	
Fusobacterium mortiferum		0.00	0.21		1	
Actinomyces israelii		0.00	0.19		99	
43315301	**Very good identification**					**43315301**
Eubacterium contortum		100.0	0.66	URE 1		
43315302	**Very good identification**					**43315302**
Eubacterium contortum		99.99	0.87			
Actinomyces israelii		0.00	0.22			
Bifidobacterium adolescentis		0.00	0.12			
43315303	**Good identification**					**43315303**
Eubacterium contortum		100.0	0.53	URE 1		
43315310	**Very good identification**					**43315310**
Eubacterium contortum		100.0	0.66	MLZ 1		
43315312	**Good identification**					**43315312**
Eubacterium contortum		99.84	0.53	MLZ 1		
Bifidobacterium adolescentis		0.16	0.14			
43315320	**Very good identification**					**43315320**
					GRA	
Eubacterium contortum		99.99	0.87		99	
Bacteroides vulgatus		0.01	0.11		1	
43315321	**Good identification**					**43315321**
Eubacterium contortum		100.0	0.53	URE 1		
43315322	**Very good identification**					**43315322**
					GRA	
Eubacterium contortum		99.89	0.74		99	
Prevotella oralis		0.05	0.16		1	
Parabacteroides distasonis		0.03	0.26		1	
43315323	**Good identification**					**43315323**
Eubacterium contortum		100.0	0.40	URE 1		
43315330	**Good identification**					**43315330**
Eubacterium contortum		100.0	0.53	MLZ 1		
43315332	**Good identification**					**43315332**
					GRA	
Eubacterium contortum		98.32	0.40	MLZ 1	99	
Parabacteroides distasonis		1.68	0.21		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43315340	**Very good identification**					**43315340**
Eubacterium contortum		100.0	0.66	SOR 1		
43315342	**Good identification**					**43315342**
Eubacterium contortum		99.92	0.53	SOR 1		
Actinomyces israelii		0.08	0.15			
43315360	**Good identification**					**43315360**
Eubacterium contortum		100.0	0.53	SOR 1		
43315362	**Good identification**					**43315362**
Eubacterium contortum		100.0	0.40	SOR 1		
43315500	**Very good identification**					**43315500**
Eubacterium contortum		99.13	0.64	RHA 99		
Actinomyces israelii		0.33	0.37			
Actinomyces meyeri		0.20	0.21			
43315502	**Low discrimination**					**43315502**
Bifidobacterium adolescentis		50.63	0.61	MNS 87		
Eubacterium contortum		47.59	0.52	RAF 92		
Actinomyces israelii		1.36	0.40			
43315512	**Very good identification**					**43315512**
Bifidobacterium adolescentis		98.59	0.62	MNS 87		
Eubacterium contortum		0.77	0.17	RAF 92		
Actinomyces israelii		0.32	0.26			
43315520	**Good identification**					**43315520**
					GRA	
Eubacterium contortum		97.39	0.52	RHA 99	99	
Bacteroides vulgatus		1.52	0.13		1	
Bifidobacterium longum subsp. longum		0.71	0.22		99	
43315522	**Low discrimination**					**43315522**
Eubacterium contortum		84.94	0.39	RHA 99		
Bifidobacterium adolescentis		5.17	0.27			
Bifidobacterium longum subsp. longum		3.49	0.22			
43315542	**Low discrimination**					**43315542**
Bifidobacterium adolescentis		86.09	0.45	MNS 87		
Actinomyces israelii		7.29	0.33	RAF 92		
Eubacterium contortum		6.61	0.17	SOR 11		
43315552	**Good identification**					**43315552**
Bifidobacterium adolescentis		98.98	0.47	MNS 87		
Actinomyces israelii		1.03	0.19	RAF 92		
43315600	**Very good identification**					**43315600**
Eubacterium contortum		100.0	0.64	LAC 99		
43315602	**Good identification**					**43315602**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		100.0	0.52	LAC 99		
43315620	**Good identification**					**43315620**
					GRA	
Eubacterium contortum		99.86	0.52	LAC 99	99	
Bacteroides vulgatus		0.14	-.05		1	
43315622	**Good identification**					**43315622**
Eubacterium contortum		100.0	0.39	LAC 99		
43315700	**Excellent identification**					**43315700**
					GRA	
Eubacterium contortum		100.0	0.99		99	
Actinomyces israelii		0.00	0.18		99	
Bacteroides vulgatus		0.00	-.05		1	
43315701	**Very good identification**					**43315701**
Eubacterium contortum		100.0	0.64	URE 1		
43315702	**Very good identification**					**43315702**
Eubacterium contortum		99.99	0.86			
Bifidobacterium adolescentis		0.01	0.27			
Actinomyces israelii		0.00	0.21			
43315703	**Good identification**					**43315703**
Eubacterium contortum		100.0	0.52	URE 1		
43315710	**Very good identification**					**43315710**
Eubacterium contortum		100.0	0.64	MLZ 1		
43315712	**Good identification**					**43315712**
Eubacterium contortum		97.88	0.52	MLZ 1		
Bifidobacterium adolescentis		1.29	0.28			
Clostridium clostridioforme		0.83	0.22			
43315720	**Very good identification**					**43315720**
					GRA	
Eubacterium contortum		99.86	0.86		99	
Bacteroides vulgatus		0.14	0.30		1	
43315721	**Good identification**					**43315721**
					GRA	
Eubacterium contortum		99.86	0.52	URE 1	99	
Bacteroides vulgatus		0.14	-.05		1	
43315722	**Very good identification**					**43315722**
					GRA	
Eubacterium contortum		99.96	0.73		99	
Parabacteroides distasonis		0.02	0.22		1	
Clostridium clostridioforme		0.01	0.10		50	
43315723	**Good identification**					**43315723**
Eubacterium contortum		100.0	0.39	URE 1		
43315730	**Good identification**					**43315730**
					GRA	
Eubacterium contortum		99.86	0.52	MLZ 1	99	
Bacteroides vulgatus		0.14	-.05		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43315732	**Acceptable identification**					**43315732**
					GRA	
Eubacterium contortum		94.24	0.39	MLZ 1	99	
Clostridium clostridioforme		4.55	0.22		50	
Parabacteroides distasonis		1.21	0.17		1	
43315740	**Very good identification**					**43315740**
Eubacterium contortum		100.0	0.64	SOR 1		
43315742	**Good identification**					**43315742**
Eubacterium contortum		99.78	0.52	SOR 1		
Bifidobacterium adolescentis		0.13	0.11			
Actinomyces israelii		0.08	0.14			
43315760	**Good identification**					**43315760**
					GRA	
Eubacterium contortum		99.86	0.52	SOR 1	99	
Bacteroides vulgatus		0.14	-.05		1	
43315762	**Good identification**					**43315762**
Eubacterium contortum		100.0	0.39	SOR 1		
43317000	**Good identification**					**43317000**
					GRA	
Actinomyces israelii		96.09	0.41	CEL 93	99	
Fusobacterium mortiferum		3.91	0.15		1	
43317002	**Good identification**					**43317002**
Actinomyces israelii		97.42	0.44	CEL 93		
Bifidobacterium adolescentis		2.58	0.11			
43317042	**Good identification**					**43317042**
Actinomyces israelii		100.0	0.37	CEL 93		
43317100	**Low discrimination**					**43317100**
					GRA	
Actinomyces israelii		63.99	0.51	CEL 93	99	
Eubacterium contortum		33.86	0.32		99	
Fusobacterium mortiferum		1.52	0.21		1	
43317102	**Low discrimination**					**43317102**
Actinomyces israelii		57.43	0.54	CEL 93		
Bifidobacterium adolescentis		37.66	0.45			
Eubacterium contortum		3.57	0.19			
43317110	**Low discrimination**					**43317110**
					SPO	
Actinomyces israelii		89.17	0.37	CEL 93	1	
Bifidobacterium adolescentis		7.25	0.12	MLZ 13	1	
Clostridium tertium		3.58	-.01		99	
43317112	**Low discrimination**					**43317112**
					SPO	
Bifidobacterium adolescentis		83.24	0.47	MNS 87	1	
Actinomyces israelii		15.52	0.40	RAF 92	1	
Clostridium tertium		1.25	0.07	ARA 87	99	
43317140	**Good identification**					**43317140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.44	CEL 93		
43317142	**Low discrimination**					**43317142**
Actinomyces israelii		82.75	0.47	CEL 93		
Bifidobacterium adolescentis		17.25	0.30			
43317300	**Very good identification**					**43317300**
Eubacterium contortum		99.86	0.66	MAN 1		
Actinomyces israelii		0.14	0.32			
43317302	**Good identification**					**43317302**
Eubacterium contortum		98.69	0.53	MAN 1		
Actinomyces israelii		1.21	0.35			
Bifidobacterium adolescentis		0.11	0.11			
43317320	**Good identification**					**43317320**
Eubacterium contortum		100.0	0.53	MAN 1		
43317322	**Good identification**					**43317322**
Eubacterium contortum		99.56	0.40	MAN 1		
Actinomyces israelii		0.44	0.15			
43317400	**Good identification**					**43317400**
Actinomyces israelii		100.0	0.40	CEL 93		
43317402	**Low discrimination**					**43317402**
Actinomyces israelii		82.19	0.43	CEL 93		
Bifidobacterium adolescentis		17.81	0.25			
43317442	**Acceptable identification**					**43317442**
Actinomyces israelii		93.56	0.36	CEL 93		
Bifidobacterium adolescentis		6.45	0.09			
43317500	**Low discrimination**					**43317500**
Actinomyces israelii		62.10	0.50	CEL 93		
Eubacterium contortum		32.85	0.30			
Bifidobacterium adolescentis		5.05	0.25			
43317502	**Low discrimination**					**43317502**
Bifidobacterium adolescentis		83.47	0.59	MNS 87		
Actinomyces israelii		15.56	0.53	RAF 92		
Eubacterium contortum		0.97	0.17			
43317510	**Low discrimination**					**43317510**
Actinomyces israelii		60.07	0.36	CEL 93		
Bifidobacterium adolescentis		39.93	0.27	MLZ 13		
43317512	**Very good identification**					**43317512**
Bifidobacterium adolescentis		97.77	0.61	MNS 87		
Actinomyces israelii		2.23	0.39	RAF 92		
43317540	**Good identification**					**43317540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		97.48	0.43	CEL 93			
Bifidobacterium adolescentis		2.52	0.09				
43317542	**Low discrimination**						**43317542**
Bifidobacterium adolescentis		63.03	0.44	MNS 87			
Actinomyces israelii		36.97	0.46	RAF 92			
43317552	**Acceptable identification**						**43317552**
Bifidobacterium adolescentis		93.31	0.45	MNS 87			
Actinomyces israelii		6.69	0.32	RAF 92			
43317700	**Very good identification**						**43317700**
Eubacterium contortum		99.86	0.64	MAN 1			
Actinomyces israelii		0.14	0.31				
43317702	**Good identification**						**43317702**
Eubacterium contortum		97.94	0.52	MAN 1			
Actinomyces israelii		1.20	0.34				
Bifidobacterium adolescentis		0.86	0.25				
43317720	**Good identification**						**43317720**
					GRA		
Eubacterium contortum		99.86	0.52	MAN 1	99		
Bacteroides vulgatus		0.14	-.05		1		
43317722	**Good identification**						**43317722**
Eubacterium contortum		99.56	0.39	MAN 1			
Actinomyces israelii		0.44	0.13				
43330000	**Very good identification**						**43330000**
					GRA	SPO	
Fusobacterium mortiferum		95.68	0.56	MLT 11	1	1	
Clostridium botulinum biovar A		1.98	0.14		99	99	
Clostridium sporogenes		1.64	0.11		99	99	
43330002	**Acceptable identification**						**43330002**
					GRA	SPO	
Fusobacterium mortiferum		91.30	0.40	MLT 11	1	1	
Clostridium sporogenes		4.69	0.03	BGL 10	99	99	
Lactobacillus jensenii		4.01	0.07		99	1	
43330100	**Very good identification**						**43330100**
					COC	GRA	
Fusobacterium mortiferum		98.35	0.63	MLT 11	1	1	
Atopobium parvulum		1.65	0.21		99	99	
43330102	**Good identification**						**43330102**
					COC	GRA	
Fusobacterium mortiferum		97.41	0.46	MLT 11	1	1	
Atopobium parvulum		2.59	0.08	BGL 10	99	99	
43331000	**Low discrimination**						**43331000**
					COC	GRA	
Fusobacterium mortiferum		65.20	0.59	MLT 11	1	1	
Lactobacillus jensenii		31.50	0.44		1	99	
Atopobium parvulum		1.70	0.21		99	99	
43331001	**Low discrimination**						**43331001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		42.48	0.37	ESL 12	99		
Fusobacterium mortiferum		38.78	0.25	MNS 88	1		
Lactobacillus jensenii		18.74	0.10	RAF 92	99		
43331002	**Low discrimination**						**43331002**
					GRA		
Lactobacillus jensenii		70.16	0.41	MNS 99	99		
Fusobacterium mortiferum		24.21	0.43	CEL 95	1		
Actinomyces naeslundii		1.61	0.34		99		
43331003	**Low discrimination**						**43331003**
Actinomyces naeslundii		77.28	0.37	ESL 12			
Lactobacillus jensenii		22.72	0.07	MNS 88			
43331010	**Acceptable identification**						**43331010**
					GRA		
Lactobacillus jensenii		90.34	0.36	MNS 99	99		
Fusobacterium mortiferum		5.67	0.25	CEL 95	1		
Eubacterium saburreum		3.99	0.21		85		
43331021	**Good identification**						**43331021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43331023	**Good identification**						**43331023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43331100	**Low discrimination**						**43331100**
					COC	GRA	
Atopobium parvulum		51.59	0.55	MNS 99	99	99	
Fusobacterium mortiferum		46.64	0.66		1	1	
Actinomyces naeslundii		0.42	0.42		1	99	
43331102	**Low discrimination**						**43331102**
					COC	GRA	
Atopobium parvulum		51.74	0.42	MNS 99	99	99	
Fusobacterium mortiferum		29.45	0.49		1	1	
Lactobacillus acidophilus		6.67	0.25		1	99	
43331103	**Low discrimination**						**43331103**
					COC	SPO	
Actinomyces naeslundii		57.22	0.45	ESL 12	1	1	
Clostridium chauvoei		29.66	0.17	MNS 88	1	99	
Atopobium parvulum		8.36	0.08	RAF 92	99	1	
43331120	**Low discrimination**						**43331120**
					COC	SPO	
Clostridium perfringens		47.84	0.45	MNS 99	1	84	
Atopobium parvulum		34.28	0.42		99	1	
Clostridium paraputrificum		7.53	0.22		1	99	
43331121	**Low discrimination**						**43331121**
					SPO		
Actinomyces naeslundii		42.94	0.45	ESL 12	1		
Clostridium chauvoei		42.04	0.22	MNS 88	99		
Clostridium perfringens		8.75	0.11	RAF 92	84		
43331122	**Low discrimination**						**43331122**
					COC		
Clostridium perfringens		52.46	0.40	MNS 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Atopobium parvulum		14.10	0.29		99	
Lactobacillus acidophilus		10.30	0.25		1	
43331123	**Low discrimination**					**43331123**
					SPO	
Actinomyces naeslundii		78.82	0.45	ESL 12	1	
Clostridium chauvoei		13.62	0.09	MNS 88	99	
Clostridium perfringens		7.56	0.06	RAF 92	84	
43331502	**Low discrimination**					**43331502**
					COC	
Bifidobacterium adolescentis		88.94	0.44	MNS 87	1	
Actinomyces israelii		7.86	0.32	RAF 92	1	
Atopobium parvulum		2.04	0.08	XYL 87	99	
43331512	**Good identification**					**43331512**
					GRA	
Bifidobacterium adolescentis		96.31	0.45	MNS 87	99	
Leptotrichia buccalis		2.02	-.03	RAF 92	1	
Actinomyces israelii		1.04	0.18	XYL 87	99	
43333001	**Good identification**					**43333001**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43333002	**Low discrimination**					**43333002**
Actinomyces israelii		63.61	0.36	CEL 93		
Actinomyces naeslundii		29.65	0.39	XYL 92		
Lactobacillus jensenii		6.74	0.07			
43333003	**Good identification**					**43333003**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43333020	**Good identification**					**43333020**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43333021	**Good identification**					**43333021**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43333022	**Low discrimination**					**43333022**
Actinomyces naeslundii		87.95	0.39	ESL 12		
Actinomyces israelii		12.05	0.15	MNS 88		
43333023	**Good identification**					**43333023**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43333100	**Low discrimination**					**43333100**
					COC	
Actinomyces israelii		45.61	0.43	CEL 93	1	
Actinomyces naeslundii		22.69	0.47	XYL 92	1	
Atopobium parvulum		14.52	0.21		99	
43333101	**Good identification**					**43333101**
Actinomyces naeslundii		100.0	0.50	ESL 12		
43333102	**Low discrimination**					**43333102**
Actinomyces israelii		56.45	0.46	CEL 93		

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii	18.72	0.47	XYL 92		
Bifidobacterium adolescentis	11.27	0.28			
43333103 Good identification					**43333103**
Actinomyces naeslundii	100.0	0.50	ESL 12		
43333120 Low discrimination					**43333120**
				SPO	
Actinomyces naeslundii	49.00	0.47	ESL 12	1	
Clostridium tertium	19.87	0.17	MNS 88	99	
Clostridium ramosum	11.59	0.11	RAF 92	92	
43333121 Good identification					**43333121**
Actinomyces naeslundii	100.0	0.50	ESL 12		
43333123 Good identification					**43333123**
Actinomyces naeslundii	100.0	0.50	ESL 12		
43333140 Low discrimination					**43333140**
Actinomyces israelii	77.99	0.36	CEL 93		
Actinomyces naeslundii	17.60	0.34	XYL 92		
Propionibacterium propionicum	4.40	0.14			
43333141 Good identification					**43333141**
Actinomyces naeslundii	100.0	0.37	ESL 12		
43333142 Low discrimination					**43333142**
Actinomyces israelii	78.72	0.39	CEL 93		
Actinomyces naeslundii	11.85	0.34	XYL 92		
Bifidobacterium adolescentis	5.00	0.12			
43333143 Good identification					**43333143**
Actinomyces naeslundii	100.0	0.37	ESL 12		
43333161 Good identification					**43333161**
Actinomyces naeslundii	100.0	0.37	ESL 12		
43333163 Good identification					**43333163**
Actinomyces naeslundii	100.0	0.37	ESL 12		
43333500 Good identification					**43333500**
Actinomyces israelii	97.59	0.42	CEL 93		
Bifidobacterium adolescentis	2.42	0.08	XYL 92		
43333502 Low discrimination					**43333502**
Bifidobacterium adolescentis	62.03	0.42	MNS 87		
Actinomyces israelii	37.98	0.45	RAF 92		
43333512 Acceptable identification					**43333512**
Bifidobacterium adolescentis	93.04	0.44	MNS 87		
Actinomyces israelii	6.96	0.31	RAF 92		
43333542 Low discrimination					**43333542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		65.83	0.38	CEL 93			
Bifidobacterium adolescentis		34.17	0.26	XYL 92			
43334000	**Good identification**						**43334000**
Fusobacterium mortiferum		100.0	0.37	MLT 11			
43334100	**Low discrimination**						**43334100**
					COC	GRA	
Fusobacterium mortiferum		70.03	0.43	MLT 11	1	1	
Atopobium parvulum		28.96	0.25	XYL 7	99	99	
Actinomyces israelii		1.01	0.14		1	99	
43335000	**Low discrimination**						**43335000**
					COC	GRA	
Fusobacterium mortiferum		43.06	0.40	MLT 11	1	1	
Atopobium parvulum		27.70	0.25	XYL 7	99	99	
Actinomyces israelii		23.78	0.38		1	99	
43335002	**Low discrimination**						**43335002**
					COC	GRA	
Actinomyces israelii		73.75	0.41	CEL 93	1	99	
Atopobium parvulum		10.11	0.13		99	99	
Fusobacterium mortiferum		9.89	0.24		1	1	
43335100	**Good identification**						**43335100**
					COC	GRA	
Atopobium parvulum		90.74	0.60	MNS 99	99	99	
Fusobacterium mortiferum		3.33	0.46		1	1	
Actinomyces israelii		3.15	0.48		1	99	
43335102	**Low discrimination**						**43335102**
					COC		
Atopobium parvulum		62.80	0.47	MNS 99	99		
Actinomyces israelii		18.51	0.51		1		
Bifidobacterium adolescentis		14.90	0.44		1		
43335112	**Low discrimination**						**43335112**
					COC		
Bifidobacterium adolescentis		82.36	0.45	MNS 87	1		
Actinomyces israelii		12.51	0.37	RAF 92	1		
Atopobium parvulum		2.87	0.13	ARA 87	99		
43335120	**Good identification**						**43335120**
					COC		
Atopobium parvulum		95.04	0.47	MNS 99	99		
Eubacterium contortum		2.63	0.19		1		
Actinomyces israelii		1.19	0.28		1		
43335140	**Low discrimination**						**43335140**
					COC	GRA	
Actinomyces israelii		56.30	0.41	CEL 93	1	99	
Atopobium parvulum		42.16	0.25		99	99	
Fusobacterium mortiferum		1.55	0.12		1	1	
43335142	**Low discrimination**						**43335142**
					COC		
Actinomyces israelii		74.41	0.44	CEL 93	1		
Bifidobacterium adolescentis		19.03	0.28		1		
Atopobium parvulum		6.56	0.13		99		
43335300	**Very good identification**						**43335300**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		99.52	0.66	TRE 1	1	
Atopobium parvulum		0.37	0.25		99	
Actinomyces israelii		0.10	0.29		1	
43335302	**Good identification**					**43335302**
					COC	
Eubacterium contortum		98.68	0.53	TRE 1	1	
Actinomyces israelii		0.80	0.32		1	
Atopobium parvulum		0.36	0.13		99	
43335320	**Good identification**					**43335320**
					COC	
Eubacterium contortum		99.63	0.53	TRE 1	1	
Atopobium parvulum		0.37	0.13		99	
43335322	**Low discrimination**					**43335322**
					GRA	
Eubacterium contortum		86.75	0.40	TRE 1	99	
Parabacteroides distasonis		12.58	0.37		1	
Clostridium clostridioforme		0.35	0.05		50	
43335400	**Good identification**					**43335400**
Actinomyces israelii		100.0	0.37	CEL 93		
43335402	**Low discrimination**					**43335402**
Actinomyces israelii		78.99	0.40	CEL 93		
Bifidobacterium adolescentis		21.01	0.23			
43335500	**Low discrimination**					**43335500**
					COC	
Actinomyces israelii		41.99	0.47	CEL 93	1	
Eubacterium contortum		33.47	0.30		1	
Atopobium parvulum		14.95	0.25		99	
43335502	**Low discrimination**					**43335502**
Bifidobacterium adolescentis		84.77	0.58	MNS 87		
Actinomyces israelii		12.88	0.50	RAF 92		
Eubacterium contortum		1.21	0.17			
43335512	**Very good identification**					**43335512**
Bifidobacterium adolescentis		97.15	0.59	MNS 87		
Actinomyces israelii		1.80	0.36	RAF 92		
Clostridium clostridioforme		0.96	0.22			
43335540	**Good identification**					**43335540**
Actinomyces israelii		96.93	0.40	CEL 93		
Bifidobacterium adolescentis		3.07	0.08			
43335542	**Low discrimination**					**43335542**
Bifidobacterium adolescentis		67.66	0.42	MNS 87		
Actinomyces israelii		32.34	0.43	RAF 92		
43335552	**Acceptable identification**					**43335552**
Bifidobacterium adolescentis		94.48	0.44	MNS 87		
Actinomyces israelii		5.52	0.29	RAF 92		
43335700	**Very good identification**					**43335700**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium contortum		99.91	0.64	TRE 1			
Actinomyces israelii		0.10	0.28				
43335702	**Good identification**						**43335702**
Eubacterium contortum		97.18	0.52	TRE 1			
Clostridium clostridioforme		1.33	0.25				
Actinomyces israelii		0.79	0.31				
43335712	**Low discrimination**						**43335712**
Clostridium clostridioforme		75.70	0.36	MNS 99			
Eubacterium contortum		12.23	0.17	RAF 94			
Bifidobacterium adolescentis		10.60	0.25				
43335720	**Good identification**						**43335720**
					GRA		
Eubacterium contortum		99.42	0.52	TRE 1	99		
Bacteroides vulgatus		0.58	0.06		1		
43335722	**Low discrimination**						**43335722**
					GRA		
Eubacterium contortum		84.29	0.39	TRE 1	99		
Parabacteroides distasonis		9.15	0.33		1		
Clostridium clostridioforme		6.55	0.25		50		
43335732	**Low discrimination**						**43335732**
					GRA		
Clostridium clostridioforme		83.78	0.36	MNS 99	50		
Parabacteroides distasonis		13.83	0.29	RAF 94	1		
Eubacterium contortum		2.39	0.04		99		
43337000	**Good identification**						**43337000**
Actinomyces israelii		100.0	0.51	CEL 93			
43337002	**Good identification**						**43337002**
Actinomyces israelii		99.53	0.54	CEL 93			
Bifidobacterium adolescentis		0.47	0.08				
43337010	**Good identification**						**43337010**
Actinomyces israelii		100.0	0.37	CEL 93			
43337012	**Good identification**						**43337012**
Actinomyces israelii		96.30	0.40	CEL 93			
Bifidobacterium adolescentis		3.70	0.09	MLZ 13			
43337040	**Good identification**						**43337040**
Actinomyces israelii		100.0	0.44	CEL 93			
43337042	**Good identification**						**43337042**
Actinomyces israelii		100.0	0.47	CEL 93			
43337100	**Good identification**						**43337100**
					COC	SPO	
Actinomyces israelii		94.06	0.61	CEL 93	1	1	
Atopobium parvulum		4.83	0.25		99	1	
Clostridium tertium		0.76	0.11		1	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
43337102	**Low discrimination**						**43337102**
					SPO		
Actinomyces israelii		87.84	0.64	CEL 93	1		
Bifidobacterium adolescentis		10.21	0.42		1		
Clostridium tertium		1.42	0.19		99		
43337110	**Acceptable identification**						**43337110**
					SPO		
Actinomyces israelii		90.05	0.47	CEL 93	1		
Clostridium tertium		8.65	0.15	MLZ 13	99		
Bifidobacterium adolescentis		1.30	0.09		1		
43337112	**Low discrimination**						**43337112**
					SPO		
Actinomyces israelii		46.67	0.50	CEL 93	1		
Bifidobacterium adolescentis		44.36	0.44	MLZ 13	1		
Clostridium tertium		8.97	0.24		99		
43337120	**Low discrimination**						**43337120**
					COC	SPO	
Actinomyces israelii		65.72	0.41	CEL 93	1	1	
Clostridium tertium		24.94	0.19	NAG 6	1	99	
Atopobium parvulum		9.34	0.13		99	1	
43337122	**Low discrimination**						**43337122**
					SPO		
Actinomyces israelii		55.74	0.44	CEL 93	1		
Clostridium tertium		42.31	0.27	NAG 6	99		
Bifidobacterium adolescentis		1.03	0.08		1		
43337140	**Good identification**						**43337140**
Actinomyces israelii		100.0	0.54	CEL 93			
43337142	**Very good identification**						**43337142**
Actinomyces israelii		96.44	0.57	CEL 93			
Bifidobacterium adolescentis		3.56	0.26				
43337150	**Good identification**						**43337150**
Actinomyces israelii		100.0	0.40	CEL 93			
43337152	**Low discrimination**						**43337152**
Actinomyces israelii		76.80	0.43	CEL 93			
Bifidobacterium adolescentis		23.20	0.28	MLZ 13			
43337162	**Good identification**						**43337162**
Actinomyces israelii		100.0	0.37	CEL 93			
43337302	**Low discrimination**						**43337302**
Actinomyces israelii		80.95	0.45	CEL 93			
Eubacterium contortum		17.79	0.19	RHA 7			
Bifidobacterium adolescentis		1.26	0.08				
43337340	**Good identification**						**43337340**
Actinomyces israelii		100.0	0.35	CEL 93			
43337342	**Good identification**						**43337342**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.38	CEL 93		
43337400	**Good identification**					**43337400**
Actinomyces israelii		100.0	0.50	CEL 93		
43337402	**Good identification**					**43337402**
Actinomyces israelii		96.30	0.53	CEL 93		
Bifidobacterium adolescentis		3.70	0.22			
43337410	**Good identification**					**43337410**
Actinomyces israelii		100.0	0.35	CEL 93		
43337412	**Low discrimination**					**43337412**
Actinomyces israelii		76.10	0.38	CEL 93		
Bifidobacterium adolescentis		23.90	0.23	MLZ 13		
43337440	**Good identification**					**43337440**
Actinomyces israelii		100.0	0.42	CEL 93		
43337442	**Good identification**					**43337442**
Actinomyces israelii		100.0	0.45	CEL 93		
43337500	**Very good identification**					**43337500**
					SPO	
Actinomyces israelii		98.36	0.60	CEL 93	1	
Bifidobacterium adolescentis		1.42	0.22		1	
Clostridium butyricum		0.22	-.07		99	
43337502	**Low discrimination**					**43337502**
Actinomyces israelii		51.27	0.63	CEL 93		
Bifidobacterium adolescentis		48.73	0.56			
43337510	**Low discrimination**					**43337510**
Actinomyces israelii		89.46	0.46	CEL 93		
Bifidobacterium adolescentis		10.54	0.23	MLZ 13		
43337512	**Low discrimination**					**43337512**
Bifidobacterium adolescentis		87.81	0.58	MNS 87		
Actinomyces israelii		11.29	0.49	RAF 92		
Bifidobacterium dentium		0.90	0.06			
43337520	**Good identification**					**43337520**
Actinomyces israelii		100.0	0.39	CEL 93		
43337522	**Low discrimination**					**43337522**
Actinomyces israelii		86.93	0.42	CEL 93		
Bifidobacterium adolescentis		13.08	0.22	NAG 6		
43337540	**Good identification**					**43337540**
Actinomyces israelii		100.0	0.53	CEL 93		
43337542	**Low discrimination**					**43337542**
Actinomyces israelii		76.80	0.56	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		23.20	0.41			
43337550	**Acceptable identification**					**43337550**
					COC	
Actinomyces israelii		91.29	0.39	CEL 93	1	
Blautia producta		5.29	-.03	MLZ 13	99	
Bifidobacterium adolescentis		3.42	0.08		1	
43337552	**Low discrimination**					**43337552**
Bifidobacterium adolescentis		71.19	0.42	MNS 87		
Actinomyces israelii		28.81	0.42	RAF 92		
43337562	**Good identification**					**43337562**
Actinomyces israelii		100.0	0.35	CEL 93		
43337702	**Low discrimination**					**43337702**
Actinomyces israelii		74.22	0.44	CEL 93		
Eubacterium contortum		16.31	0.17	RHA 7		
Bifidobacterium adolescentis		9.47	0.22			
43337742	**Good identification**					**43337742**
Actinomyces israelii		100.0	0.37	CEL 93		
43341000	**Good identification**					**43341000**
Eubacterium tortuosum		95.55	0.50	MLT 1		
Lactobacillus catenaformis		2.33	0.29	CEL 10		
Lactobacillus jensenii		1.17	0.24			
43341002	**Low discrimination**					**43341002**
					SPO	
Eubacterium tortuosum		81.00	0.37	MLT 1	1	
Lactobacillus catenaformis		11.20	0.29	CEL 10	1	
Clostridium barati		3.77	0.15		99	
43341020	**Acceptable identification**					**43341020**
					SPO	
Eubacterium tortuosum		90.34	0.37	MLT 1	1	
Clostridium barati		4.20	0.15	CEL 10	99	
Lactobacillus catenaformis		3.12	0.18		1	
43347100	**Good identification**					**43347100**
Actinomyces israelii		100.0	0.37	FRU 99		
43347102	**Low discrimination**					**43347102**
					SPO	
Actinomyces israelii		86.57	0.40	FRU 99	1	
Bifidobacterium adolescentis		9.08	0.16		1	
Clostridium tertium		4.35	0.03		99	
43347500	**Good identification**					**43347500**
Actinomyces israelii		100.0	0.35	FRU 99		
43347502	**Low discrimination**					**43347502**
Actinomyces israelii		53.83	0.38	FRU 99		
Bifidobacterium adolescentis		46.18	0.31			
43350000	**Low discrimination**					**43350000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		84.62	0.57	MLT 11	1	1	
Lactobacillus catenaformis		7.00	0.29		99	1	
Clostridium barati		4.09	0.19		99	99	
43350022	**Acceptable identification**						**43350022**
					SPO		
Clostridium barati		93.18	0.36	MNS 99	99		
Lactobacillus catenaformis		4.43	0.18	GAL 95	1		
Lactobacillus jensenii		1.48	0.10		1		
43350100	**Low discrimination**						**43350100**
					GRA	SPO	
Fusobacterium mortiferum		66.46	0.63	MLT 11	1	1	
Collinsella aerofaciens		25.73	0.32		99	1	
Clostridium barati		4.13	0.28		99	99	
43350102	**Low discrimination**						**43350102**
					GRA	SPO	
Clostridium barati		46.06	0.36	MNS 99	99	99	
Fusobacterium mortiferum		27.45	0.47	GAL 95	1	1	
Lactobacillus catenaformis		13.14	0.32		99	1	
43350122	**Low discrimination**						**43350122**
					SPO		
Clostridium barati		80.98	0.44	MNS 99	99		
Clostridium paraputrificum		13.44	0.25	GAL 95	99		
Lactobacillus catenaformis		1.93	0.21		1		
43351000	**Low discrimination**						**43351000**
					GRA		
Lactobacillus catenaformis		52.20	0.63	MNS 99	99		
Lactobacillus jensenii		26.21	0.58		99		
Fusobacterium mortiferum		9.56	0.60		1		
43351002	**Low discrimination**						**43351002**
					SPO		
Lactobacillus catenaformis		57.66	0.63	MNS 99	1		
Clostridium barati		19.40	0.50		99		
Lactobacillus jensenii		19.30	0.55		1		
43351010	**Acceptable identification**						**43351010**
					GRA		
Lactobacillus jensenii		92.02	0.50	MNS 99	99		
Lactobacillus catenaformis		5.55	0.29		99		
Fusobacterium mortiferum		1.02	0.26		1		
43351012	**Low discrimination**						**43351012**
					SPO		
Lactobacillus jensenii		88.11	0.47	MNS 99	1		
Lactobacillus catenaformis		7.98	0.29		1		
Clostridium barati		2.68	0.15		99		
43351020	**Low discrimination**						**43351020**
					SPO		
Clostridium barati		42.55	0.50	MNS 99	99		
Lactobacillus catenaformis		31.62	0.53		1		
Lactobacillus jensenii		15.87	0.47		1		
43351022	**Low discrimination**						**43351022**
					SPO		
Clostridium barati		71.52	0.58	MNS 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		17.71	0.53		1	
Lactobacillus jensenii		5.93	0.44		1	
43351030	**Low discrimination**					**43351030**
					SPO	
Lactobacillus jensenii		86.47	0.39	MNS 99	1	
Clostridium barati		7.02	0.15		99	
Lactobacillus catenaformis		5.22	0.18		1	
43351032	**Low discrimination**					**43351032**
					SPO	
Lactobacillus jensenii		67.30	0.36	MNS 99	1	
Clostridium barati		24.60	0.24		99	
Lactobacillus catenaformis		6.09	0.18		1	
43351062	**Acceptable identification**					**43351062**
					SPO	
Clostridium barati		90.95	0.40	MNS 99	99	
Lactobacillus acidophilus		3.17	0.17	SOR 8	1	
Lactobacillus catenaformis		2.62	0.18		1	
43351100	**Low discrimination**					**43351100**
					GRA	
Collinsella aerofaciens		82.25	0.66	MNS 99	99	
Lactobacillus catenaformis		11.30	0.66		99	
Fusobacterium mortiferum		3.22	0.66		1	
43351102	**Low discrimination**					**43351102**
					SPO	
Lactobacillus catenaformis		36.15	0.66	MNS 99	1	
Collinsella aerofaciens		29.23	0.50		1	
Clostridium barati		24.33	0.58		99	
43351120	**Low discrimination**					**43351120**
					SPO	
Clostridium paraputrificum		69.77	0.66	MNS 99	99	
Collinsella aerofaciens		13.85	0.50		1	
Clostridium barati		11.52	0.58		99	
43351122	**Low discrimination**					**43351122**
					SPO	
Clostridium barati		45.03	0.66	MNS 99	99	
Clostridium paraputrificum		38.95	0.60		99	
Lactobacillus catenaformis		5.58	0.56		1	
43351132	**Low discrimination**					**43351132**
					SPO	
Bifidobacterium breve		28.12	0.44	MNS 99	1	
Clostridium barati		27.79	0.32	RAF 95	99	
Clostridium paraputrificum		24.03	0.25		99	
43351142	**Low discrimination**					**43351142**
					SPO	
Clostridium barati		49.44	0.40	MNS 99	99	
Lactobacillus acidophilus		19.84	0.35	SOR 8	1	
Bifidobacterium breve		9.81	0.40		1	
43351152	**Low discrimination**					**43351152**
					SPO	
Bifidobacterium breve		77.05	0.40	MNS 99	1	
Bifidobacterium adolescentis		10.49	0.24	RAF 95	1	
Clostridium barati		3.92	0.05		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43351160	**Low discrimination**						**43351160**
					SPO		
Clostridium paraputrificum		80.97	0.47	MNS 99	99		
Clostridium barati		15.45	0.40	SOR 7	99		
Collinsella aerofaciens		2.16	0.15		1		
43351162	**Low discrimination**						**43351162**
					SPO		
Clostridium barati		49.18	0.48	MNS 99	99		
Clostridium paraputrificum		36.82	0.40	SOR 8	99		
Lactobacillus acidophilus		6.58	0.35		1		
43351172	**Low discrimination**						**43351172**
					SPO		
Bifidobacterium breve		86.08	0.44	MNS 99	1		
Clostridium barati		7.40	0.14	RAF 95	99		
Clostridium paraputrificum		5.54	0.06		99		
43351222	**Low discrimination**						**43351222**
					GRA	SPO	
Clostridium barati		88.18	0.40	MNS 99	99	99	
Prevotella oralis		6.29	0.21	RHA 8	1	1	
Lactobacillus catenaformis		2.54	0.18		99	1	
43351302	**Low discrimination**						**43351302**
					SPO		
Clostridium barati		59.58	0.40	MNS 99	99		
Lactobacillus acidophilus		14.56	0.31	RHA 8	1		
Lactobacillus catenaformis		10.29	0.32		1		
43351320	**Low discrimination**						**43351320**
					SPO		
Clostridium barati		49.62	0.40	MNS 99	99		
Clostridium paraputrificum		34.90	0.32	RHA 8	99		
Collinsella aerofaciens		6.93	0.15		1		
43351322	**Low discrimination**						**43351322**
					GRA	SPO	
Clostridium barati		61.98	0.48	MNS 99	99	99	
Prevotella oralis		23.09	0.41	RHA 8	1	1	
Clostridium paraputrificum		6.23	0.25		99	99	
43351502	**Low discrimination**						**43351502**
Bifidobacterium adolescentis		67.09	0.52	MNS 87			
Lactobacillus catenaformis		9.68	0.32	RAF 92			
Collinsella aerofaciens		7.83	0.15	XYL 87			
43351512	**Good identification**						**43351512**
					GRA		
Bifidobacterium adolescentis		98.18	0.54	MNS 87	99		
Actinomyces israelii		1.18	0.27	RAF 92	99		
Leptotrichia buccalis		0.65	-.03	XYL 87	1		
43351542	**Low discrimination**						**43351542**
					SPO		
Bifidobacterium adolescentis		71.23	0.37	MNS 87	1		
Actinomyces israelii		21.95	0.34	RAF 92	1		
Clostridium barati		4.87	0.05	XYL 87	99		
43351552	**Good identification**						**43351552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		96.37	0.38	MNS 87		
Actinomyces israelii		3.63	0.20	RAF 92		
43353000	**Low discrimination**					**43353000**
					SPO	
Actinomyces israelii		25.21	0.42	XYL 92	1	
Lactobacillus catenaformis		21.46	0.29		1	
Clostridium barati		20.71	0.23		99	
43353001	**Good identification**					**43353001**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43353003	**Low discrimination**					**43353003**
					SPO	
Actinomyces naeslundii		89.06	0.40	ESL 12	1	
Clostridium barati		10.94	-.03	MNS 88	99	
43353021	**Low discrimination**					**43353021**
					SPO	
Actinomyces naeslundii		89.06	0.40	ESL 12	1	
Clostridium barati		10.94	-.03	MNS 88	99	
43353022	**Acceptable identification**					**43353022**
					SPO	
Clostridium barati		90.42	0.40	MNS 99	99	
Lactobacillus catenaformis		2.60	0.18	MAN 8	1	
Actinomyces naeslundii		1.65	0.37		1	
43353023	**Low discrimination**					**43353023**
					SPO	
Actinomyces naeslundii		73.08	0.40	ESL 12	1	
Clostridium barati		26.92	0.05	MNS 88	99	
43353040	**Low discrimination**					**43353040**
					SPO	
Actinomyces israelii		80.32	0.35	XYL 92	1	
Clostridium barati		14.76	0.05		99	
Actinomyces naeslundii		4.93	0.24		1	
43353042	**Low discrimination**					**43353042**
					SPO	
Actinomyces israelii		71.01	0.38	XYL 92	1	
Clostridium barati		26.09	0.13		99	
Actinomyces naeslundii		2.90	0.24		1	
43353101	**Low discrimination**					**43353101**
Actinomyces naeslundii		78.38	0.47	ESL 12		
Collinsella aerofaciens		12.75	-.03	MNS 88		
Actinomyces israelii		5.49	0.18	RAF 92		
43353102	**Low discrimination**					**43353102**
					SPO	
Clostridium barati		32.85	0.40	MNS 99	99	
Actinomyces israelii		26.66	0.56	MAN 8	1	
Clostridium tertium		11.17	0.35		99	
43353103	**Low discrimination**					**43353103**
					SPO	
Actinomyces naeslundii		78.22	0.47	ESL 12	1	
Clostridium barati		10.12	0.05	MNS 88	99	
Actinomyces israelii		8.22	0.21	RAF 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43353112	**Low discrimination**					**43353112**
					SPO	
Clostridium tertium		53.71	0.39	MNS 99	99	
Bifidobacterium breve		17.63	0.40	TRE 90	1	
Bifidobacterium adolescentis		15.89	0.38		1	
43353120	**Low discrimination**					**43353120**
Clostridium barati		40.63	0.40	MNS 99		
Clostridium paraputrificum		28.58	0.32	MAN 8		
Clostridium tertium		13.81	0.35			
43353121	**Low discrimination**					**43353121**
					SPO	
Actinomyces naeslundii		79.09	0.47	ESL 12	1	
Clostridium barati		10.24	0.05	MNS 88	99	
Clostridium paraputrificum		7.20	-.03	RAF 92	99	
43353122	**Low discrimination**					**43353122**
Clostridium barati		54.65	0.48	MNS 99		
Clostridium tertium		18.58	0.43	MAN 8		
Clostridium ramosum		8.30	0.35			
43353123	**Low discrimination**					**43353123**
					SPO	
Actinomyces naeslundii		60.10	0.47	ESL 12	1	
Clostridium barati		23.34	0.14	MNS 88	99	
Clostridium tertium		7.93	0.09	RAF 92	99	
43353130	**Acceptable identification**					**43353130**
					SPO	
Clostridium tertium		90.97	0.39	MNS 99	99	
Bifidobacterium breve		5.25	0.27	TRE 90	1	
Clostridium barati		1.52	0.05		99	
43353132	**Low discrimination**					**43353132**
					SPO	
Clostridium tertium		81.88	0.47	MNS 99	99	
Bifidobacterium breve		15.93	0.44	TRE 90	1	
Clostridium barati		1.37	0.14		99	
43353140	**Low discrimination**					**43353140**
					SPO	
Actinomyces israelii		74.91	0.46	XYL 92	1	
Clostridium barati		10.32	0.13		99	
Bifidobacterium breve		6.99	0.23		1	
43353142	**Low discrimination**					**43353142**
					SPO	
Actinomyces israelii		48.35	0.49	XYL 92	1	
Bifidobacterium breve		30.39	0.40		1	
Clostridium barati		13.32	0.21		99	
43353152	**Low discrimination**					**43353152**
Bifidobacterium breve		72.47	0.40	MNS 99		
Actinomyces israelii		17.23	0.34	RAF 95		
Bifidobacterium adolescentis		8.07	0.22			
43353162	**Low discrimination**					**43353162**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		50.27	0.44	MNS 99	1	
Clostridium barati		37.19	0.30	RAF 95	99	
Clostridium paraputrificum		3.23	0.06		99	
43353172	**Acceptable identification**					**43353172**
					SPO	
Bifidobacterium breve		94.30	0.44	MNS 99	1	
Clostridium tertium		4.90	0.13	RAF 95	99	
Actinomyces israelii		0.81	0.14		1	
43353322	**Low discrimination**					**43353322**
					SPO	
Clostridium ramosum		66.39	0.37	MNS 99	92	
Clostridium barati		28.67	0.30	TRE 94	99	
Lactobacillus acidophilus		1.41	0.09		1	
43353400	**Acceptable identification**					**43353400**
					SPO	
Actinomyces israelii		90.59	0.41	XYL 92	1	
Clostridium innocuum		9.41	0.09		99	
43353402	**Low discrimination**					**43353402**
					SPO	
Actinomyces israelii		87.02	0.44	XYL 92	1	
Clostridium innocuum		6.03	0.09		99	
Bifidobacterium adolescentis		5.18	0.16		1	
43353442	**Good identification**					**43353442**
Actinomyces israelii		100.0	0.37	XYL 92		
43353500	**Acceptable identification**					**43353500**
Actinomyces israelii		94.46	0.51	XYL 92		
Collinsella aerofaciens		2.71	-.03			
Bifidobacterium adolescentis		2.11	0.16			
43353502	**Low discrimination**					**43353502**
					SPO	
Bifidobacterium adolescentis		59.10	0.51	MNS 87	1	
Actinomyces israelii		40.09	0.54	RAF 92	1	
Clostridium barati		0.61	0.05	XYL 87	99	
43353510	**Low discrimination**					**43353510**
Actinomyces israelii		84.55	0.37	XYL 92		
Bifidobacterium adolescentis		15.45	0.18	MLZ 13		
43353512	**Acceptable identification**					**43353512**
					SPO	
Bifidobacterium adolescentis		91.91	0.52	MNS 87	1	
Actinomyces israelii		7.62	0.40	RAF 92	1	
Clostridium tertium		0.47	0.05	XYL 87	99	
43353540	**Good identification**					**43353540**
Actinomyces israelii		100.0	0.44	XYL 92		
43353542	**Low discrimination**					**43353542**
Actinomyces israelii		68.10	0.47	XYL 92		
Bifidobacterium adolescentis		31.90	0.35			
43353552	**Low discrimination**					**43353552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		79.31	0.37	MNS 87		
Actinomyces israelii		20.70	0.33	RAF 92		
43354000	**Good identification**					**43354000**
					GRA	
Fusobacterium mortiferum		97.42	0.38	MLT 11	1	
Actinomyces israelii		2.58	0.13	XYL 7	99	
43354100	**Acceptable identification**					**43354100**
					GRA	
Fusobacterium mortiferum		91.13	0.44	MLT 11	1	
Collinsella aerofaciens		4.73	-.03	XYL 7	99	
Actinomyces israelii		4.14	0.24		99	
43355000	**Low discrimination**					**43355000**
					GRA	
Actinomyces israelii		43.57	0.47		99	
Fusobacterium mortiferum		24.93	0.41		1	
Lactobacillus catenaformis		18.27	0.29		99	
43355002	**Low discrimination**					**43355002**
					SPO	
Actinomyces israelii		64.69	0.50		1	
Lactobacillus catenaformis		18.09	0.29		1	
Clostridium barati		6.08	0.15		99	
43355012	**Low discrimination**					**43355012**
Actinomyces israelii		69.02	0.36	MLZ 13		
Bifidobacterium adolescentis		16.57	0.19			
Lactobacillus jensenii		14.41	0.12			
43355040	**Good identification**					**43355040**
Actinomyces israelii		100.0	0.40			
43355042	**Good identification**					**43355042**
					SPO	
Actinomyces israelii		97.94	0.43		1	
Clostridium barati		2.06	-.03		99	
43355102	**Low discrimination**					**43355102**
Actinomyces israelii		47.45	0.61			
Bifidobacterium adolescentis		34.47	0.52			
Lactobacillus catenaformis		4.98	0.32			
43355110	**Low discrimination**					**43355110**
Actinomyces israelii		82.84	0.44	MLZ 13		
Bifidobacterium adolescentis		7.46	0.19			
Collinsella aerofaciens		6.41	-.03			
43355112	**Low discrimination**					**43355112**
					GRA	
Bifidobacterium adolescentis		84.70	0.54	MNS 87	99	
Actinomyces israelii		14.26	0.47	RAF 92	99	
Leptotrichia buccalis		0.56	-.03	ARA 87	1	
43355140	**Good identification**					**43355140**
					GRA	
Actinomyces israelii		95.58	0.51		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Collinsella aerofaciens		2.84	-.03		99		
Fusobacterium mortiferum		0.83	0.13		1		
43355142	**Low discrimination**						**43355142**
					SPO		
Actinomyces israelii		79.81	0.54		1		
Bifidobacterium adolescentis		18.43	0.37		1		
Clostridium barati		1.26	0.05		99		
43355150	**Good identification**						**43355150**
Actinomyces israelii		100.0	0.37	MLZ 13			
43355152	**Low discrimination**						**43355152**
Bifidobacterium adolescentis		65.38	0.38	MNS 87			
Actinomyces israelii		34.62	0.40	RAF 92			
43355300	**Very good identification**						**43355300**
Eubacterium contortum		99.58	0.66	CEL 1			
Actinomyces israelii		0.34	0.39				
Collinsella aerofaciens		0.05	-.03				
43355302	**Good identification**						**43355302**
Eubacterium contortum		96.58	0.53	CEL 1			
Actinomyces israelii		2.77	0.42				
Bifidobacterium adolescentis		0.27	0.18				
43355320	**Good identification**						**43355320**
					GRA	SPO	
Eubacterium contortum		99.32	0.53	CEL 1	99	1	
Clostridium barati		0.23	0.05		99	99	
Prevotella oralis		0.17	0.04		1	1	
43355322	**Low discrimination**						**43355322**
					GRA	SPO	
Eubacterium contortum		79.56	0.40	CEL 1	99	1	
Prevotella oralis		14.39	0.26		1	1	
Clostridium barati		3.16	0.14		99	99	
43355400	**Good identification**						**43355400**
					SPO		
Actinomyces israelii		96.66	0.46		1		
Clostridium butyricum		3.34	0.00		99		
43355402	**Low discrimination**						**43355402**
					SPO		
Actinomyces israelii		80.06	0.49		1		
Bifidobacterium adolescentis		19.22	0.32		1		
Clostridium butyricum		0.72	-.07		99		
43355440	**Good identification**						**43355440**
Actinomyces israelii		100.0	0.39				
43355442	**Acceptable identification**						**43355442**
Actinomyces israelii		92.91	0.42				
Bifidobacterium adolescentis		7.09	0.17				
43355500	**Low discrimination**						**43355500**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		64.89	0.56		1	
Eubacterium contortum		14.64	0.30		1	
Clostridium butyricum		13.44	0.24		99	
43355502	**Low discrimination**					**43355502**
					SPO	
Bifidobacterium adolescentis		84.59	0.66	MNS 87	1	
Actinomyces israelii		14.24	0.59	RAF 92	1	
Clostridium butyricum		0.76	0.17		99	
43355510	**Low discrimination**					**43355510**
					SPO	
Actinomyces israelii		55.27	0.42	MLZ 13	1	
Bifidobacterium adolescentis		40.70	0.34		1	
Clostridium butyricum		4.03	0.02		99	
43355512	**Very good identification**					**43355512**
Bifidobacterium adolescentis		97.81	0.68	MNS 87		
Actinomyces israelii		2.01	0.45	RAF 92		
Clostridium clostridioforme		0.14	0.16			
43355520	**Low discrimination**					**43355520**
					SPO	
Actinomyces israelii		48.06	0.36	NAG 6	1	
Eubacterium contortum		29.99	0.17		1	
Clostridium butyricum		13.56	0.05		99	
43355522	**Low discrimination**					**43355522**
					GRA	
Actinomyces israelii		38.50	0.39	NAG 6	99	
Bifidobacterium adolescentis		36.20	0.32		99	
Bacteroides uniformis		13.90	0.09		1	
43355540	**Acceptable identification**					**43355540**
					SPO	
Actinomyces israelii		93.03	0.49		1	
Clostridium butyricum		4.31	0.05		99	
Bifidobacterium adolescentis		2.66	0.17		1	
43355542	**Low discrimination**					**43355542**
					SPO	
Bifidobacterium adolescentis		65.11	0.51	MNS 87	1	
Actinomyces israelii		34.48	0.52	RAF 92	1	
Clostridium butyricum		0.41	-.02	SOR 11	99	
43355550	**Low discrimination**					**43355550**
Actinomyces israelii		81.04	0.35	MLZ 13		
Bifidobacterium adolescentis		18.96	0.18			
43355552	**Acceptable identification**					**43355552**
Bifidobacterium adolescentis		93.92	0.52	MNS 87		
Actinomyces israelii		6.08	0.38	RAF 92		
43355700	**Very good identification**					**43355700**
					SPO	
Eubacterium contortum		99.64	0.64	CEL 1	1	
Actinomyces israelii		0.34	0.37		1	
Clostridium butyricum		0.03	-.02		99	
43355702	**Acceptable identification**					**43355702**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		94.54	0.52	CEL 1		
Actinomyces israelii		2.71	0.40			
Bifidobacterium adolescentis		2.16	0.32			
43355720	**Good identification**					**43355720**
					GRA	
Eubacterium contortum		99.15	0.52	CEL 1	99	
Bacteroides vulgatus		0.73	0.08		1	
Actinomyces israelii		0.12	0.17		99	
43355722	**Low discrimination**					**43355722**
					GRA	
Eubacterium contortum		88.71	0.39	CEL 1	99	
Bacteroides uniformis		6.09	0.11		1	
Clostridium clostridioforme		3.15	0.19		50	
43356100	**Good identification**					**43356100**
Actinomyces israelii		100.0	0.37	GAL 99		
43356102	**Low discrimination**					**43356102**
					SPO	
Actinomyces israelii		86.57	0.40	GAL 99	1	
Bifidobacterium adolescentis		9.08	0.16		1	
Clostridium tertium		4.35	0.03		99	
43356500	**Good identification**					**43356500**
Actinomyces israelii		100.0	0.35	GAL 99		
43356502	**Low discrimination**					**43356502**
Actinomyces israelii		53.83	0.38	GAL 99		
Bifidobacterium adolescentis		46.18	0.31			
43357000	**Very good identification**					**43357000**
					SPO	
Actinomyces israelii		99.78	0.60		1	
Clostridium innocuum		0.22	0.00		99	
43357002	**Very good identification**					**43357002**
					SPO	
Actinomyces israelii		99.09	0.63		1	
Bifidobacterium adolescentis		0.42	0.16		1	
Clostridium tertium		0.20	0.03		99	
43357010	**Good identification**					**43357010**
					SPO	
Actinomyces israelii		98.81	0.46	MLZ 13	1	
Clostridium tertium		1.20	-.01		99	
43357012	**Acceptable identification**					**43357012**
					SPO	
Actinomyces israelii		94.44	0.49	MLZ 13	1	
Bifidobacterium adolescentis		3.27	0.18		1	
Clostridium tertium		2.28	0.07		99	
43357020	**Acceptable identification**					**43357020**
					SPO	
Actinomyces israelii		92.45	0.40	NAG 6	1	
Clostridium tertium		4.42	0.03		99	
Clostridium barati		3.14	-.03		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43357022	**Low discrimination**					**43357022**
					SPO	
Actinomyces israelii		85.96	0.43	NAG 6	1	
Clostridium tertium		8.21	0.11		99	
Clostridium barati		5.83	0.05		99	
43357040	**Good identification**					**43357040**
Actinomyces israelii		100.0	0.53			
43357042	**Very good identification**					**43357042**
Actinomyces israelii		100.0	0.56			
43357050	**Good identification**					**43357050**
Actinomyces israelii		100.0	0.39	MLZ 13		
43357052	**Good identification**					**43357052**
Actinomyces israelii		100.0	0.42	MLZ 13		
43357062	**Good identification**					**43357062**
Actinomyces israelii		100.0	0.36	NAG 6		
43357100	**Very good identification**					**43357100**
					SPO	
Actinomyces israelii		97.10	0.71		1	
Clostridium tertium		2.44	0.29		99	
Collinsella aerofaciens		0.20	-.03		1	
43357101	**Good identification**					**43357101**
Actinomyces israelii		100.0	0.37	URE 1		
43357102	**Low discrimination**					**43357102**
					SPO	
Actinomyces israelii		86.49	0.74		1	
Bifidobacterium adolescentis		9.07	0.51		1	
Clostridium tertium		4.35	0.37		99	
43357103	**Low discrimination**					**43357103**
					SPO	
Actinomyces israelii		86.57	0.40	URE 1	1	
Bifidobacterium adolescentis		9.08	0.16		1	
Clostridium tertium		4.35	0.03		99	
43357110	**Low discrimination**					**43357110**
					SPO	
Actinomyces israelii		76.21	0.57	MLZ 13	1	
Clostridium tertium		22.80	0.33		99	
Bifidobacterium adolescentis		0.99	0.18		1	
43357112	**Low discrimination**					**43357112**
					SPO	
Actinomyces israelii		40.71	0.60	MLZ 13	1	
Bifidobacterium adolescentis		34.93	0.52		1	
Clostridium tertium		24.36	0.41		99	
43357120	**Low discrimination**					**43357120**
					SPO	
Clostridium tertium		53.12	0.37	MNS 99	99	
Actinomyces israelii		44.94	0.50	TRE 90	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		1.14	0.05		99	
43357122	**Low discrimination**					**43357122**
					SPO	
Clostridium tertium		68.41	0.45	MNS 99	99	
Actinomyces israelii		28.94	0.53	TRE 90	1	
Clostridium barati		1.47	0.14		99	
43357130	**Acceptable identification**					**43357130**
					SPO	
Clostridium tertium		93.36	0.41	MNS 99	99	
Actinomyces israelii		6.64	0.36	TRE 90	1	
43357132	**Good identification**					**43357132**
					SPO	
Clostridium tertium		95.99	0.50	MNS 99	99	
Actinomyces israelii		3.41	0.39	TRE 90	1	
Bifidobacterium adolescentis		0.46	0.18		1	
43357140	**Very good identification**					**43357140**
Actinomyces israelii		100.0	0.64			
43357142	**Very good identification**					**43357142**
					SPO	
Actinomyces israelii		96.65	0.67		1	
Bifidobacterium adolescentis		3.22	0.35		1	
Clostridium tertium		0.13	0.03		99	
43357150	**Good identification**					**43357150**
					SPO	
Actinomyces israelii		99.23	0.50	MLZ 13	1	
Clostridium tertium		0.77	-.01		99	
43357152	**Low discrimination**					**43357152**
					SPO	
Actinomyces israelii		77.63	0.53	MLZ 13	1	
Bifidobacterium adolescentis		21.17	0.37		1	
Clostridium tertium		1.21	0.07		99	
43357160	**Good identification**					**43357160**
					SPO	
Actinomyces israelii		97.02	0.43	NAG 6	1	
Clostridium tertium		2.98	0.03		99	
43357162	**Acceptable identification**					**43357162**
					SPO	
Actinomyces israelii		92.87	0.46	NAG 6	1	
Clostridium tertium		5.70	0.11		99	
Bifidobacterium breve		1.43	0.10		1	
43357200	**Good identification**					**43357200**
Actinomyces israelii		100.0	0.41	RHA 7		
43357202	**Good identification**					**43357202**
Actinomyces israelii		100.0	0.44	RHA 7		
43357242	**Good identification**					**43357242**
Actinomyces israelii		100.0	0.37	RHA 7		
43357300	**Low discrimination**					**43357300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		65.40	0.51	RHA 7		
Eubacterium contortum		34.60	0.32			
43357302	**Acceptable identification**					**43357302**
Actinomyces israelii		92.33	0.54	RHA 7		
Eubacterium contortum		5.75	0.19			
Bifidobacterium adolescentis		1.30	0.16			
43357310	**Good identification**					**43357310**
					SPO	
Actinomyces israelii		96.14	0.37	RHA 7	1	
Clostridium tertium		3.86	-.01	MLZ 13	99	
43357312	**Low discrimination**					**43357312**
					SPO	
Actinomyces israelii		83.65	0.40	RHA 7	1	
Bifidobacterium adolescentis		9.63	0.18	MLZ 13	1	
Clostridium tertium		6.72	0.07		99	
43357340	**Good identification**					**43357340**
Actinomyces israelii		100.0	0.44	RHA 7		
43357342	**Good identification**					**43357342**
Actinomyces israelii		100.0	0.47	RHA 7		
43357400	**Very good identification**					**43357400**
Actinomyces israelii		100.0	0.59			
43357402	**Very good identification**					**43357402**
Actinomyces israelii		96.65	0.62			
Bifidobacterium adolescentis		3.35	0.31			
43357410	**Good identification**					**43357410**
Actinomyces israelii		100.0	0.45	MLZ 13		
43357412	**Low discrimination**					**43357412**
Actinomyces israelii		77.91	0.48	MLZ 13		
Bifidobacterium adolescentis		22.09	0.32			
43357420	**Good identification**					**43357420**
Actinomyces israelii		100.0	0.38	NAG 6		
43357422	**Good identification**					**43357422**
Actinomyces israelii		100.0	0.41	NAG 6		
43357440	**Good identification**					**43357440**
Actinomyces israelii		100.0	0.52			
43357442	**Good identification**					**43357442**
Actinomyces israelii		98.91	0.55			
Bifidobacterium adolescentis		1.09	0.15			
43357450	**Good identification**					**43357450**
Actinomyces israelii		100.0	0.38	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43357452	**Acceptable identification**					**43357452**
Actinomyces israelii		91.74	0.41	MLZ 13		
Bifidobacterium adolescentis		8.27	0.17			
43357500	**Very good identification**					**43357500**
					SPO	
Actinomyces israelii		98.32	0.69		1	
Bifidobacterium adolescentis		1.28	0.31		1	
Clostridium butyricum		0.40	0.07		99	
43357501	**Good identification**					**43357501**
Actinomyces israelii		100.0	0.35	URE 1		
43357502	**Low discrimination**					**43357502**
					SPO	
Actinomyces israelii		53.78	0.72		1	
Bifidobacterium adolescentis		46.13	0.65		1	
Clostridium butyricum		0.06	0.00		99	
43357503	**Low discrimination**					**43357503**
Actinomyces israelii		53.83	0.38	URE 1		
Bifidobacterium adolescentis		46.18	0.31			
43357510	**Acceptable identification**					**43357510**
					SPO	
Actinomyces israelii		90.09	0.55	MLZ 13	1	
Bifidobacterium adolescentis		9.58	0.32		1	
Clostridium tertium		0.33	-.01		99	
43357512	**Low discrimination**					**43357512**
Bifidobacterium adolescentis		87.20	0.66	MNS 87		
Actinomyces israelii		12.43	0.58	RAF 92		
Bifidobacterium dentium		0.28	0.06			
43357520	**Good identification**					**43357520**
					SPO	
Actinomyces israelii		98.56	0.49	NAG 6	1	
Clostridium tertium		1.44	0.03		99	
43357522	**Low discrimination**					**43357522**
					SPO	
Actinomyces israelii		85.84	0.52	NAG 6	1	
Bifidobacterium adolescentis		11.65	0.31		1	
Clostridium tertium		2.51	0.11		99	
43357540	**Very good identification**					**43357540**
Actinomyces israelii		99.59	0.62			
Bifidobacterium adolescentis		0.41	0.15			
43357542	**Low discrimination**					**43357542**
Actinomyces israelii		78.58	0.65			
Bifidobacterium adolescentis		21.42	0.49			
43357550	**Good identification**					**43357550**
					COC	
Actinomyces israelii		95.22	0.48	MLZ 13	1	
Bifidobacterium adolescentis		3.22	0.17		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Blautia producta		1.56	-.03		99	
43357552	**Low discrimination**					**43357552**
Bifidobacterium adolescentis		69.04	0.51	MNS 87		
Actinomyces israelii		30.96	0.51	RAF 92		
43357560	**Good identification**					**43357560**
Actinomyces israelii		100.0	0.42	NAG 6		
43357562	**Good identification**					**43357562**
Actinomyces israelii		95.86	0.45	NAG 6		
Bifidobacterium adolescentis		4.14	0.15			
43357600	**Good identification**					**43357600**
Actinomyces israelii		100.0	0.40	RHA 7		
43357602	**Good identification**					**43357602**
Actinomyces israelii		100.0	0.43	RHA 7		
43357642	**Good identification**					**43357642**
Actinomyces israelii		100.0	0.36	RHA 7		
43357700	**Low discrimination**					**43357700**
Actinomyces israelii		65.40	0.50	RHA 7		
Eubacterium contortum		34.60	0.30			
43357702	**Low discrimination**					**43357702**
Actinomyces israelii		84.94	0.53	RHA 7		
Bifidobacterium adolescentis		9.78	0.31			
Eubacterium contortum		5.29	0.17			
43357710	**Good identification**					**43357710**
Actinomyces israelii		100.0	0.36	RHA 7		
43357712	**Low discrimination**					**43357712**
Actinomyces israelii		51.50	0.39	RHA 7		
Bifidobacterium adolescentis		48.50	0.32	MLZ 13		
43357740	**Good identification**					**43357740**
Actinomyces israelii		100.0	0.43	RHA 7		
43357742	**Good identification**					**43357742**
Actinomyces israelii		96.47	0.46	RHA 7		
Bifidobacterium adolescentis		3.53	0.15			
43365102	**Low discrimination**					**43365102**
					COC	
Actinomyces israelii		57.99	0.36	FRU 99	1	
Atopobium parvulum		34.55	0.19		99	
Bifidobacterium adolescentis		7.47	0.15		1	
43367000	**Good identification**					**43367000**
Actinomyces israelii		100.0	0.36	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43367002	**Good identification**							**43367002**
Actinomyces israelii		100.0	0.39	FRU 99				
43367100	**Acceptable identification**							**43367100**
					SPO			
Actinomyces israelii		94.33	0.46	FRU 99	1			
Clostridium tertium		5.68	0.11		99			
43367102	**Low discrimination**							**43367102**
					SPO			
Actinomyces israelii		87.80	0.49	FRU 99	1			
Clostridium tertium		10.57	0.19		99			
Bifidobacterium adolescentis		1.63	0.13		1			
43367140	**Good identification**							**43367140**
Actinomyces israelii		100.0	0.39	FRU 99				
43367142	**Good identification**							**43367142**
Actinomyces israelii		100.0	0.42	FRU 99				
43367402	**Good identification**							**43367402**
Actinomyces israelii		100.0	0.38	FRU 99				
43367500	**Good identification**							**43367500**
					SPO			
Actinomyces israelii		98.34	0.45	FRU 99	1			
Clostridium butyricum		1.66	-.07		99			
43367502	**Low discrimination**							**43367502**
Actinomyces israelii		86.80	0.48	FRU 99				
Bifidobacterium adolescentis		13.20	0.28					
43367540	**Good identification**							**43367540**
Actinomyces israelii		100.0	0.38	FRU 99				
43367542	**Good identification**							**43367542**
Actinomyces israelii		95.39	0.41	FRU 99				
Bifidobacterium adolescentis		4.61	0.12					
43370000	**Low discrimination**							**43370000**
					GRA	SPO		
Fusobacterium mortiferum		69.87	0.48	MLT 11	1	1		
Lactobacillus jensenii		29.15	0.32		99	1		
Clostridium barati		0.98	0.01		99	99		
43370100	**Low discrimination**							**43370100**
					COC	GRA	SPO	
Fusobacterium mortiferum		86.79	0.54	MLT 11	1	1	1	
Atopobium parvulum		10.18	0.27		99	99	1	
Clostridium barati		1.57	0.10		1	99	99	
43371000	**Good identification**							**43371000**
					GRA			
Lactobacillus jensenii		94.01	0.66	MNS 99	99			
Fusobacterium mortiferum		3.41	0.51		1			
Lactobacillus catenaformis		0.63	0.29		99			
43371002	**Low discrimination**							**43371002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Lactobacillus jensenii		87.43	0.63	MNS 99	1		
Lactobacillus acidophilus		7.22	0.41		1		
Clostridium barati		2.55	0.31		99		
43371010	**Very good identification**						**43371010**
					GRA		
Lactobacillus jensenii		99.49	0.58	MNS 99	99		
Actinomyces israelii		0.23	0.25		99		
Leptotrichia buccalis		0.17	-.03		1		
43371012	**Low discrimination**						**43371012**
					GRA		
Lactobacillus jensenii		79.48	0.55	MNS 99	99		
Leptotrichia buccalis		19.84	0.32		1		
Actinomyces israelii		0.42	0.28		99		
43371020	**Low discrimination**						**43371020**
					SPO		
Lactobacillus jensenii		87.31	0.56	MNS 99	1		
Clostridium barati		6.78	0.31		99		
Clostridium paraputrificum		3.81	0.22		99		
43371022	**Low discrimination**						**43371022**
					SPO		
Lactobacillus jensenii		57.54	0.53	MNS 99	1		
Clostridium barati		20.12	0.40		99		
Lactobacillus acidophilus		19.01	0.41		1		
43371030	**Good identification**						**43371030**
					SPO		
Lactobacillus jensenii		99.77	0.47	MNS 99	1		
Clostridium barati		0.24	-.03		99		
43371032	**Low discrimination**						**43371032**
					GRA	SPO	
Lactobacillus jensenii		88.38	0.44	MNS 99	99	1	
Leptotrichia buccalis		9.80	0.15		1	1	
Clostridium barati		0.94	0.05		99	99	
43371100	**Low discrimination**						**43371100**
					COC	GRA	
Atopobium parvulum		73.08	0.61	MNS 99	99	99	
Fusobacterium mortiferum		9.44	0.57		1	1	
Collinsella aerofaciens		8.15	0.32		1	99	
43371102	**Low discrimination**						**43371102**
					COC	SPO	
Lactobacillus acidophilus		66.37	0.60	MNS 99	1	1	
Atopobium parvulum		12.14	0.48		99	1	
Clostridium barati		6.11	0.40		1	99	
43371112	**Very good identification**						**43371112**
					GRA		
Leptotrichia buccalis		99.37	0.66	MNS 99	1		
Bifidobacterium adolescentis		0.15	0.36		99		
Bifidobacterium breve		0.14	0.37		99		
43371120	**Low discrimination**						**43371120**
					COC	SPO	
Clostridium paraputrificum		81.10	0.56	MNS 99	1	99	
Atopobium parvulum		8.71	0.48		99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		4.38	0.40		1	99	
43371122	**Low discrimination**						**43371122**
					SPO		
Lactobacillus acidophilus		44.88	0.60	MNS 99	1		
Clostridium paraputrificum		32.76	0.50		99		
Clostridium barati		12.39	0.48		99		
43371132	**Acceptable identification**						**43371132**
					GRA	SPO	
Leptotrichia buccalis		94.12	0.50	MNS 99	1	1	
Bifidobacterium breve		2.11	0.41	NAG 10	99	1	
Clostridium tertium		1.48	0.29		99	99	
43371140	**Low discrimination**						**43371140**
					COC	SPO	
Actinomyces israelii		38.83	0.42	XYL 92	1	1	
Atopobium parvulum		31.62	0.27		99	1	
Clostridium barati		8.06	0.13		1	99	
43371142	**Low discrimination**						**43371142**
Lactobacillus acidophilus		73.34	0.44	MNS 99			
Actinomyces israelii		11.44	0.45	SOR 11			
Bifidobacterium breve		7.22	0.37				
43371160	**Low discrimination**						**43371160**
					SPO		
Clostridium paraputrificum		88.60	0.37	MNS 99	99		
Clostridium barati		5.53	0.21	SOR 7	99		
Bifidobacterium breve		1.48	0.24		1		
43371162	**Low discrimination**						**43371162**
					SPO		
Lactobacillus acidophilus		54.17	0.44	MNS 99	1		
Clostridium paraputrificum		24.08	0.31	SOR 11	99		
Clostridium barati		10.52	0.30		99		
43371172	**Low discrimination**						**43371172**
					GRA		
Bifidobacterium breve		65.22	0.41	MNS 99	99		
Leptotrichia buccalis		29.34	0.15	RAF 95	1		
Lactobacillus acidophilus		3.77	0.10		99		
43371302	**Low discrimination**						**43371302**
					SPO		
Lactobacillus acidophilus		82.15	0.40	MNS 99	1		
Clostridium barati		8.73	0.21	RHA 7	99		
Actinomyces israelii		4.07	0.33		1		
43371322	**Low discrimination**						**43371322**
					SPO		
Clostridium ramosum		57.38	0.45	MNS 99	92		
Lactobacillus acidophilus		28.11	0.40		1		
Clostridium barati		8.96	0.30		99		
43371500	**Low discrimination**						**43371500**
					COC	SPO	
Actinomyces israelii		54.77	0.48	XYL 92	1	1	
Atopobium parvulum		21.20	0.27		99	1	
Clostridium butyricum		16.80	0.18		1	99	
43371502	**Low discrimination**						**43371502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		56.05	0.49	MNS 87		
Actinomyces israelii		30.98	0.51	RAF 92		
Lactobacillus acidophilus		7.72	0.25	XYL 87		
43371512	**Low discrimination**					**43371512**
					GRA	
Bifidobacterium adolescentis		48.68	0.51	MNS 87	99	
Leptotrichia buccalis		47.53	0.32	RAF 92	1	
Actinomyces israelii		3.29	0.37	XYL 87	99	
43371540	**Acceptable identification**					**43371540**
					SPO	
Actinomyces israelii		93.58	0.41	XYL 92	1	
Clostridium butyricum		6.42	0.00		99	
43371542	**Low discrimination**					**43371542**
Actinomyces israelii		59.81	0.44	XYL 92		
Bifidobacterium adolescentis		34.39	0.34			
Lactobacillus acidophilus		4.74	0.10			
43371552	**Low discrimination**					**43371552**
					GRA	
Bifidobacterium adolescentis		77.38	0.35	MNS 87	99	
Actinomyces israelii		16.45	0.30	RAF 92	99	
Leptotrichia buccalis		6.18	-.03	XYL 87	1	
43373000	**Low discrimination**					**43373000**
Actinomyces israelii		66.35	0.52	XYL 92		
Lactobacillus jensenii		22.61	0.32			
Actinomyces naeslundii		4.63	0.41			
43373001	**Acceptable identification**					**43373001**
Actinomyces naeslundii		91.20	0.44	ESL 12		
Actinomyces israelii		8.80	0.18	MNS 88		
43373002	**Low discrimination**					**43373002**
Actinomyces israelii		72.91	0.55	XYL 92		
Lactobacillus jensenii		11.04	0.29			
Lactobacillus acidophilus		4.75	0.20			
43373003	**Low discrimination**					**43373003**
Actinomyces naeslundii		87.35	0.44	ESL 12		
Actinomyces israelii		12.65	0.21	MNS 88		
43373010	**Low discrimination**					**43373010**
					SPO	
Actinomyces israelii		49.98	0.38	XYL 92	1	
Lactobacillus jensenii		37.99	0.24	MLZ 13	1	
Clostridium tertium		12.04	0.13		99	
43373012	**Low discrimination**					**43373012**
					SPO	
Actinomyces israelii		52.53	0.41	XYL 92	1	
Clostridium tertium		25.30	0.21	MLZ 13	99	
Lactobacillus jensenii		17.74	0.21		1	
43373021	**Good identification**					**43373021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	ESL 12			
43373023	**Good identification**						**43373023**
Actinomyces naeslundii		100.0	0.44	ESL 12			
43373040	**Good identification**						**43373040**
Actinomyces israelii		96.93	0.45	XYL 92			
Actinomyces naeslundii		3.07	0.29				
43373042	**Good identification**						**43373042**
Actinomyces israelii		95.99	0.48	XYL 92			
Actinomyces naeslundii		2.03	0.29				
Lactobacillus acidophilus		1.99	0.04				
43373100	**Low discrimination**						**43373100**
					COC	SPO	
Actinomyces israelii		60.52	0.62	XYL 92	1	1	
Clostridium tertium		30.32	0.43		1	99	
Atopobium parvulum		3.38	0.27		99	1	
43373101	**Low discrimination**						**43373101**
					SPO		
Actinomyces naeslundii		83.08	0.52	ESL 12	1		
Actinomyces israelii		11.27	0.28	MNS 88	1		
Clostridium tertium		5.65	0.09	RAF 92	99		
43373102	**Low discrimination**						**43373102**
					SPO		
Clostridium tertium		42.47	0.51	MNS 99	99		
Actinomyces israelii		42.38	0.65		1		
Lactobacillus acidophilus		7.94	0.38		1		
43373103	**Low discrimination**						**43373103**
					SPO		
Actinomyces naeslundii		69.18	0.52	ESL 12	1		
Clostridium tertium		14.11	0.17	MNS 88	99		
Actinomyces israelii		14.08	0.31	RAF 92	1		
43373110	**Low discrimination**						**43373110**
					SPO		
Clostridium tertium		84.52	0.47	MNS 99	99		
Actinomyces israelii		14.18	0.48		1		
Bifidobacterium breve		0.61	0.20		1		
43373112	**Low discrimination**						**43373112**
					GRA	SPO	
Clostridium tertium		76.50	0.55	MNS 99	99	99	
Leptotrichia buccalis		13.39	0.32		1	1	
Actinomyces israelii		6.42	0.51		99	1	
43373120	**Low discrimination**						**43373120**
Clostridium tertium		56.62	0.51	MNS 99			
Clostridium ramosum		33.04	0.45				
Clostridium paraputrificum		3.46	0.22				
43373121	**Low discrimination**						**43373121**
					SPO		
Actinomyces naeslundii		75.59	0.52	ESL 12	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		15.42	0.17	MNS 88	99		
Clostridium ramosum		9.00	0.11	RAF 92	92		
43373122	**Low discrimination**						**43373122**
					SPO		
Clostridium tertium		51.66	0.59	MNS 99	99		
Clostridium ramosum		40.19	0.56		92		
Lactobacillus acidophilus		3.22	0.38		1		
43373123	**Low discrimination**						**43373123**
					SPO		
Actinomyces naeslundii		47.04	0.52	ESL 12	1		
Clostridium tertium		28.78	0.25	MNS 88	99		
Clostridium ramosum		22.39	0.21	RAF 92	92		
43373130	**Good identification**						**43373130**
					SPO		
Clostridium tertium		98.80	0.55	MNS 99	99		
Bifidobacterium breve		0.42	0.24		1		
Actinomyces israelii		0.35	0.28		1		
43373132	**Very good identification**						**43373132**
					GRA	SPO	
Clostridium tertium		97.30	0.63	MNS 99	99	99	
Bifidobacterium breve		1.40	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
43373140	**Good identification**						**43373140**
Actinomyces israelii		95.05	0.55	XYL 92			
Actinomyces naeslundii		2.14	0.36				
Bifidobacterium breve		1.57	0.20				
43373141	**Low discrimination**						**43373141**
Actinomyces naeslundii		76.98	0.39	ESL 12			
Actinomyces israelii		23.02	0.21	MNS 88			
43373142	**Low discrimination**						**43373142**
Actinomyces israelii		81.40	0.58	XYL 92			
Bifidobacterium breve		9.07	0.37				
Lactobacillus acidophilus		4.85	0.22				
43373143	**Low discrimination**						**43373143**
Actinomyces naeslundii		69.04	0.39	ESL 12			
Actinomyces israelii		30.96	0.24	MNS 88			
43373150	**Low discrimination**						**43373150**
					SPO		
Actinomyces israelii		79.03	0.41	XYL 92	1		
Clostridium tertium		12.24	0.13	MLZ 13	99		
Bifidobacterium breve		8.74	0.20		1		
43373152	**Low discrimination**						**43373152**
					SPO		
Actinomyces israelii		45.61	0.44	XYL 92	1		
Bifidobacterium breve		34.00	0.37	MLZ 13	1		
Clostridium tertium		14.13	0.21		99		
43373161	**Good identification**						**43373161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	ESL 12		
43373162	**Low discrimination**					**43373162**
					SPO	
Bifidobacterium breve		39.65	0.41	MNS 99	1	
Clostridium tertium		15.63	0.25	RAF 95	99	
Actinomyces israelii		12.78	0.38		1	
43373163	**Good identification**					**43373163**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43373172	**Low discrimination**					**43373172**
					SPO	
Bifidobacterium breve		57.17	0.41	MNS 99	1	
Clostridium tertium		40.08	0.29	RAF 95	99	
Actinomyces israelii		2.75	0.24		1	
43373202	**Acceptable identification**					**43373202**
Actinomyces israelii		93.88	0.36	XYL 92		
Lactobacillus acidophilus		6.12	0.00	RHA 7		
43373300	**Low discrimination**					**43373300**
					SPO	
Actinomyces israelii		81.75	0.43	XYL 92	1	
Clostridium ramosum		12.76	0.13	RHA 7	92	
Clostridium tertium		5.50	0.09		99	
43373302	**Low discrimination**					**43373302**
					SPO	
Actinomyces israelii		56.47	0.46	XYL 92	1	
Clostridium ramosum		23.50	0.24	RHA 7	92	
Lactobacillus acidophilus		10.58	0.18		1	
43373320	**Good identification**					**43373320**
					SPO	
Clostridium ramosum		98.00	0.47	MNS 99	92	
Clostridium tertium		1.28	0.17		99	
Actinomyces israelii		0.41	0.23		1	
43373322	**Very good identification**					**43373322**
					SPO	
Clostridium ramosum		98.20	0.58	MNS 99	92	
Clostridium tertium		0.96	0.25		99	
Lactobacillus acidophilus		0.45	0.18		1	
43373340	**Good identification**					**43373340**
Actinomyces israelii		100.0	0.36	XYL 92		
43373342	**Acceptable identification**					**43373342**
Actinomyces israelii		94.38	0.39	XYL 92		
Lactobacillus acidophilus		5.62	0.03	RHA 7		
43373400	**Good identification**					**43373400**
					SPO	
Actinomyces israelii		99.04	0.51	XYL 92	1	
Clostridium innocuum		0.96	0.01		99	
43373402	**Good identification**					**43373402**
					SPO	
Actinomyces israelii		98.33	0.54	XYL 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		1.04	0.13		1	
Clostridium innocuum		0.64	0.01		99	
43373410	**Good identification**					**43373410**
Actinomyces israelii		100.0	0.36	XYL 92		
43373412	**Acceptable identification**					**43373412**
Actinomyces israelii		92.05	0.39	XYL 92		
Bifidobacterium adolescentis		7.95	0.15	MLZ 13		
43373440	**Good identification**					**43373440**
Actinomyces israelii		100.0	0.44	XYL 92		
43373442	**Good identification**					**43373442**
Actinomyces israelii		100.0	0.47	XYL 92		
43373500	**Very good identification**					**43373500**
					SPO	
Actinomyces israelii		98.41	0.61	XYL 92	1	
Clostridium tertium		0.61	0.09		99	
Clostridium butyricum		0.59	0.02		99	
43373502	**Low discrimination**					**43373502**
					SPO	
Actinomyces israelii		78.28	0.64	XYL 92	1	
Bifidobacterium adolescentis		20.45	0.48		1	
Clostridium tertium		0.97	0.17		99	
43373510	**Acceptable identification**					**43373510**
					SPO	
Actinomyces israelii		90.42	0.47	XYL 92	1	
Clostridium tertium		6.65	0.13	MLZ 13	99	
Bifidobacterium adolescentis		2.93	0.15		1	
43373512	**Low discrimination**					**43373512**
					SPO	
Bifidobacterium adolescentis		63.25	0.49	MNS 87	1	
Actinomyces israelii		29.60	0.50	RAF 92	1	
Clostridium tertium		4.36	0.21	XYL 87	99	
43373520	**Low discrimination**					**43373520**
					SPO	
Actinomyces israelii		68.48	0.40	XYL 92	1	
Clostridium tertium		19.91	0.17	NAG 6	99	
Clostridium ramosum		11.62	0.11		92	
43373522	**Low discrimination**					**43373522**
					SPO	
Actinomyces israelii		47.36	0.43	XYL 92	1	
Clostridium tertium		27.54	0.25	NAG 6	99	
Clostridium ramosum		21.43	0.21		92	
43373540	**Good identification**					**43373540**
Actinomyces israelii		100.0	0.54	XYL 92		
43373542	**Good identification**					**43373542**
Actinomyces israelii		92.33	0.57	XYL 92		
Bifidobacterium adolescentis		7.67	0.32			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
43373550	**Good identification**						**43373550**
					COC		
Actinomyces israelii		95.22	0.40	XYL 92	1		
Blautia producta		4.78	-.03	MLZ 13	99		
43373552	**Low discrimination**						**43373552**
Actinomyces israelii		59.56	0.43	XYL 92			
Bifidobacterium adolescentis		40.45	0.34	MLZ 13			
43373562	**Good identification**						**43373562**
Actinomyces israelii		100.0	0.36	XYL 92			
43373700	**Good identification**						**43373700**
Actinomyces israelii		100.0	0.42	XYL 92			
43373702	**Good identification**						**43373702**
Actinomyces israelii		96.61	0.45	XYL 92			
Bifidobacterium adolescentis		3.39	0.13	RHA 7			
43373742	**Good identification**						**43373742**
Actinomyces israelii		100.0	0.38	XYL 92			
43374102	**Low discrimination**						**43374102**
					COC	GRA	
Actinomyces israelii		53.94	0.36	GAL 99	1	99	
Atopobium parvulum		32.13	0.19		99	99	
Fusobacterium mortiferum		6.99	0.19		1	1	
43374712	**Good identification**						**43374712**
Clostridium clostridioforme		100.0	0.37	MNS 99			
43374732	**Good identification**						**43374732**
Clostridium clostridioforme		100.0	0.37	MNS 99			
43375000	**Low discrimination**						**43375000**
					COC		
Actinomyces israelii		69.88	0.57		1		
Atopobium parvulum		14.30	0.32		99		
Lactobacillus jensenii		11.73	0.32		1		
43375002	**Low discrimination**						**43375002**
					COC		
Actinomyces israelii		89.37	0.60		1		
Lactobacillus jensenii		6.67	0.29		1		
Atopobium parvulum		2.15	0.19		99		
43375010	**Low discrimination**						**43375010**
Actinomyces israelii		72.75	0.43	MLZ 13			
Lactobacillus jensenii		27.25	0.24				
43375012	**Low discrimination**						**43375012**
Actinomyces israelii		79.97	0.46	MLZ 13			
Lactobacillus jensenii		13.31	0.21				
Bifidobacterium adolescentis		3.40	0.16				
43375020	**Low discrimination**						**43375020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces israelii		43.97	0.37	NAG 6	1		
Lactobacillus jensenii		28.91	0.21		1		
Atopobium parvulum		24.87	0.19		99		
43375022	**Low discrimination**						**43375022**
					SPO		
Actinomyces israelii		64.20	0.40	NAG 6	1		
Lactobacillus jensenii		18.76	0.18		1		
Clostridium barati		6.56	0.05		99		
43375040	**Good identification**						**43375040**
Actinomyces israelii		100.0	0.50				
43375042	**Good identification**						**43375042**
Actinomyces israelii		100.0	0.53				
43375050	**Good identification**						**43375050**
Actinomyces israelii		100.0	0.36	MLZ 13			
43375052	**Good identification**						**43375052**
Actinomyces israelii		100.0	0.39	MLZ 13			
43375100	**Low discrimination**						**43375100**
					COC	SPO	
Atopobium parvulum		81.97	0.66	MNS 99	99	1	
Actinomyces israelii		16.19	0.68		1	1	
Clostridium butyricum		1.27	0.28		1	99	
43375102	**Low discrimination**						**43375102**
					COC		
Actinomyces israelii		56.35	0.71		1		
Atopobium parvulum		33.57	0.53		99		
Bifidobacterium adolescentis		7.25	0.49		1		
43375103	**Low discrimination**						**43375103**
					COC		
Actinomyces israelii		57.99	0.36	URE 1	1		
Atopobium parvulum		34.55	0.19		99		
Bifidobacterium adolescentis		7.47	0.15		1		
43375110	**Low discrimination**						**43375110**
					COC	SPO	
Actinomyces israelii		69.33	0.54	MLZ 13	1	1	
Atopobium parvulum		23.73	0.32		99	1	
Clostridium tertium		2.84	0.15		1	99	
43375112	**Low discrimination**						**43375112**
					GRA		
Bifidobacterium adolescentis		32.73	0.51	MNS 87	99		
Leptotrichia buccalis		31.96	0.32	RAF 92	1		
Actinomyces israelii		31.08	0.57	ARA 87	99		
43375120	**Low discrimination**						**43375120**
					COC	SPO	
Atopobium parvulum		87.25	0.53	MNS 99	99	1	
Actinomyces israelii		6.23	0.47		1	1	
Clostridium paraputrificum		4.42	0.22		1	99	
43375122	**Low discrimination**						**43375122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Atopobium parvulum		44.39	0.40	MNS 99	99	1	
Actinomyces israelii		26.95	0.50		1	1	
Clostridium tertium		8.73	0.27		1	99	
43375140	**Low discrimination**						**43375140**
					COC	SPO	
Actinomyces israelii		87.03	0.61		1	1	
Atopobium parvulum		11.45	0.32		99	1	
Clostridium butyricum		1.53	0.10		1	99	
43375142	**Good identification**						**43375142**
					COC		
Actinomyces israelii		93.76	0.64		1		
Bifidobacterium adolescentis		3.84	0.34		1		
Atopobium parvulum		1.45	0.19		99		
43375150	**Good identification**						**43375150**
Actinomyces israelii		100.0	0.46	MLZ 13			
43375152	**Low discrimination**						**43375152**
					GRA		
Actinomyces israelii		73.46	0.49	MLZ 13	99		
Bifidobacterium adolescentis		24.58	0.35		99		
Leptotrichia buccalis		1.96	-.03		1		
43375160	**Low discrimination**						**43375160**
					COC	SPO	
Actinomyces israelii		66.63	0.40	NAG 6	1	1	
Atopobium parvulum		24.23	0.19		99	1	
Clostridium paraputrificum		9.15	0.03		1	99	
43375162	**Low discrimination**						**43375162**
					COC		
Actinomyces israelii		85.45	0.43	NAG 6	1		
Lactobacillus acidophilus		7.54	0.10		1		
Atopobium parvulum		3.66	0.06		99		
43375200	**Good identification**						**43375200**
Actinomyces israelii		100.0	0.38	RHA 7			
43375202	**Good identification**						**43375202**
Actinomyces israelii		97.41	0.41	RHA 7			
Clostridium clostridioforme		2.59	0.05				
43375300	**Low discrimination**						**43375300**
					COC		
Actinomyces israelii		39.86	0.48	RHA 7	1		
Eubacterium contortum		31.77	0.32		1		
Atopobium parvulum		27.09	0.32		99		
43375302	**Low discrimination**						**43375302**
					COC		
Actinomyces israelii		80.69	0.51	RHA 7	1		
Eubacterium contortum		7.57	0.19		1		
Atopobium parvulum		6.45	0.19		99		
43375312	**Low discrimination**						**43375312**
Actinomyces israelii		51.07	0.37	RHA 7			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		34.67	0.25	MLZ 13			
Bifidobacterium adolescentis		7.22	0.16				
43375340	**Good identification**						**43375340**
Actinomyces israelii		100.0	0.41	RHA 7			
43375342	**Good identification**						**43375342**
Actinomyces israelii		100.0	0.44	RHA 7			
43375400	**Low discrimination**						**43375400**
					SPO		
Actinomyces israelii		87.43	0.56		1		
Clostridium butyricum		12.57	0.20		99		
43375402	**Good identification**						**43375402**
					SPO		
Actinomyces israelii		92.16	0.59		1		
Bifidobacterium adolescentis		3.92	0.29		1		
Clostridium butyricum		3.44	0.13		99		
43375410	**Good identification**						**43375410**
					SPO		
Actinomyces israelii		95.18	0.42	MLZ 13	1		
Clostridium butyricum		4.82	-.02		99		
43375412	**Low discrimination**						**43375412**
Actinomyces israelii		66.34	0.45	MLZ 13			
Bifidobacterium adolescentis		23.09	0.31				
Clostridium clostridioforme		10.58	0.22				
43375420	**Low discrimination**						**43375420**
					SPO		
Actinomyces israelii		83.62	0.35	NAG 6	1		
Clostridium butyricum		16.38	0.02		99		
43375422	**Low discrimination**						**43375422**
					SPO		
Actinomyces israelii		88.29	0.38	NAG 6	1		
Clostridium clostridioforme		7.23	0.10		66		
Clostridium butyricum		4.48	-.05		99		
43375440	**Good identification**						**43375440**
					SPO		
Actinomyces israelii		96.89	0.49		1		
Clostridium butyricum		3.11	0.02		99		
43375442	**Good identification**						**43375442**
					SPO		
Actinomyces israelii		97.86	0.52		1		
Bifidobacterium adolescentis		1.32	0.14		1		
Clostridium butyricum		0.82	-.05		99		
43375452	**Acceptable identification**						**43375452**
Actinomyces israelii		90.04	0.38	MLZ 13			
Bifidobacterium adolescentis		9.96	0.15				
43375500	**Low discrimination**						**43375500**
					COC	SPO	
Actinomyces israelii		51.52	0.66		1	1	
Clostridium butyricum		44.44	0.44		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Atopobium parvulum		3.22	0.32		99 1	
43375502	**Low discrimination**					**43375502**
					SPO	
Bifidobacterium adolescentis		46.01	0.63	MNS 87	1	
Actinomyces israelii		43.70	0.69	RAF 92	1	
Clostridium butyricum		9.77	0.37		99	
43375510	**Low discrimination**					**43375510**
					SPO	
Actinomyces israelii		69.72	0.52	MLZ 13	1	
Clostridium butyricum		21.18	0.22		99	
Bifidobacterium adolescentis		9.10	0.31		1	
43375512	**Low discrimination**					**43375512**
Bifidobacterium adolescentis		87.25	0.65	MNS 87		
Actinomyces israelii		10.13	0.55	RAF 92		
Clostridium clostridioforme		1.35	0.31			
43375522	**Low discrimination**					**43375522**
					SPO	
Actinomyces israelii		62.59	0.49	NAG 6	1	
Clostridium butyricum		19.07	0.19		99	
Bifidobacterium adolescentis		10.43	0.29		1	
43375540	**Low discrimination**					**43375540**
					SPO	
Actinomyces israelii		83.48	0.59		1	
Clostridium butyricum		16.10	0.26		99	
Bifidobacterium adolescentis		0.42	0.14		1	
43375542	**Low discrimination**					**43375542**
					SPO	
Actinomyces israelii		72.22	0.62		1	
Bifidobacterium adolescentis		24.17	0.48		1	
Clostridium butyricum		3.61	0.19		99	
43375550	**Low discrimination**					**43375550**
					SPO	
Actinomyces israelii		88.17	0.45	MLZ 13	1	
Clostridium butyricum		5.99	0.04		99	
Bifidobacterium adolescentis		3.66	0.15		1	
43375552	**Low discrimination**					**43375552**
					SPO	
Bifidobacterium adolescentis		72.90	0.49	MNS 87	1	
Actinomyces israelii		26.63	0.48	RAF 92	1	
Clostridium butyricum		0.47	-.03	SOR 11	99	
43375560	**Low discrimination**					**43375560**
					SPO	
Actinomyces israelii		79.19	0.39	NAG 6	1	
Clostridium butyricum		20.81	0.08		99	
43375562	**Low discrimination**					**43375562**
					SPO	
Actinomyces israelii		89.20	0.42	NAG 6	1	
Clostridium butyricum		6.08	0.01		99	
Bifidobacterium adolescentis		4.72	0.14		1	
43375600	**Acceptable identification**					**43375600**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		94.42	0.37	RHA 7	1	
Clostridium butyricum		5.58	-.06		99	
43375602	**Low discrimination**					**43375602**
Actinomyces israelii		66.25	0.40	RHA 7		
Clostridium clostridioforme		33.75	0.25			
43375612	**Acceptable identification**					**43375612**
Clostridium clostridioforme		93.95	0.36	MNS 99		
Actinomyces israelii		6.05	0.25	RAF 94		
43375632	**Good identification**					**43375632**
Clostridium clostridioforme		100.0	0.36	MNS 99		
43375700	**Low discrimination**					**43375700**
					SPO	
Actinomyces israelii		46.48	0.47	RHA 7	1	
Eubacterium contortum		37.04	0.30		1	
Clostridium butyricum		16.47	0.18		99	
43375702	**Low discrimination**					**43375702**
Actinomyces israelii		57.04	0.50	RHA 7		
Clostridium clostridioforme		24.32	0.34			
Bifidobacterium adolescentis		8.06	0.29			
43375712	**Low discrimination**					**43375712**
Clostridium clostridioforme		85.78	0.45	MNS 99		
Bifidobacterium adolescentis		7.63	0.31	RAF 94		
Actinomyces israelii		6.60	0.36			
43375732	**Good identification**					**43375732**
Clostridium clostridioforme		95.74	0.45	MNS 99		
Parabacteroides distasonis		2.54	0.24	RAF 94		
Bacteroides ovatus		1.25	0.03			
43375740	**Acceptable identification**					**43375740**
					SPO	
Actinomyces israelii		92.66	0.40	RHA 7	1	
Clostridium butyricum		7.34	0.00		99	
43375742	**Acceptable identification**					**43375742**
					SPO	
Actinomyces israelii		93.86	0.43	RHA 7	1	
Bifidobacterium adolescentis		4.22	0.14		1	
Clostridium butyricum		1.93	-.07		99	
43376000	**Good identification**					**43376000**
Actinomyces israelii		100.0	0.36	GAL 99		
43376002	**Good identification**					**43376002**
Actinomyces israelii		100.0	0.39	GAL 99		
43376100	**Acceptable identification**					**43376100**
					SPO	
Actinomyces israelii		94.33	0.46	GAL 99	1	
Clostridium tertium		5.68	0.11		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43376102	**Low discrimination**					**43376102**
					SPO	
Actinomyces israelii		87.80	0.49	GAL 99	1	
Clostridium tertium		10.57	0.19		99	
Bifidobacterium adolescentis		1.63	0.13		1	
43376140	**Good identification**					**43376140**
Actinomyces israelii		100.0	0.39	GAL 99		
43376142	**Good identification**					**43376142**
Actinomyces israelii		100.0	0.42	GAL 99		
43376402	**Good identification**					**43376402**
Actinomyces israelii		100.0	0.38	GAL 99		
43376500	**Good identification**					**43376500**
					SPO	
Actinomyces israelii		98.34	0.45	GAL 99	1	
Clostridium butyricum		1.66	-.07		99	
43376502	**Low discrimination**					**43376502**
Actinomyces israelii		86.80	0.48	GAL 99		
Bifidobacterium adolescentis		13.20	0.28			
43376540	**Good identification**					**43376540**
Actinomyces israelii		100.0	0.38	GAL 99		
43376542	**Good identification**					**43376542**
Actinomyces israelii		95.39	0.41	GAL 99		
Bifidobacterium adolescentis		4.61	0.12			
43377000	**Very good identification**					**43377000**
					SPO	
Actinomyces israelii		99.76	0.70		1	
Clostridium tertium		0.24	0.11		99	
43377001	**Good identification**					**43377001**
Actinomyces israelii		100.0	0.36	URE 1		
43377002	**Very good identification**					**43377002**
					SPO	
Actinomyces israelii		99.44	0.73		1	
Clostridium tertium		0.48	0.19		99	
Bifidobacterium adolescentis		0.08	0.13		1	
43377003	**Good identification**					**43377003**
Actinomyces israelii		100.0	0.39	URE 1		
43377010	**Very good identification**					**43377010**
					SPO	
Actinomyces israelii		97.19	0.56	MLZ 13	1	
Clostridium tertium		2.81	0.15		99	
43377012	**Good identification**					**43377012**
					SPO	
Actinomyces israelii		93.99	0.59	MLZ 13	1	
Clostridium tertium		5.44	0.24		99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		0.58	0.15		1		
43377020	**Low discrimination**						**43377020**
					SPO		
Actinomyces israelii		89.75	0.50	NAG 6	1		
Clostridium tertium		10.25	0.19		99		
43377022	**Low discrimination**						**43377022**
					SPO		
Actinomyces israelii		81.40	0.53	NAG 6	1		
Clostridium tertium		18.60	0.27		99		
43377040	**Very good identification**						**43377040**
Actinomyces israelii		100.0	0.63				
43377042	**Very good identification**						**43377042**
Actinomyces israelii		100.0	0.66				
43377050	**Good identification**						**43377050**
Actinomyces israelii		100.0	0.49	MLZ 13			
43377052	**Good identification**						**43377052**
Actinomyces israelii		100.0	0.52	MLZ 13			
43377060	**Good identification**						**43377060**
Actinomyces israelii		100.0	0.43	NAG 6			
43377062	**Good identification**						**43377062**
Actinomyces israelii		100.0	0.46	NAG 6			
43377100	**Good identification**						**43377100**
					COC	SPO	
Actinomyces israelii		93.37	0.81		1	1	
Clostridium tertium		5.62	0.45		1	99	
Atopobium parvulum		0.84	0.32		99	1	
43377101	**Acceptable identification**						**43377101**
					SPO		
Actinomyces israelii		94.33	0.46	URE 1	1		
Clostridium tertium		5.68	0.11		99		
43377102	**Low discrimination**						**43377102**
					SPO		
Actinomyces israelii		87.68	0.84		1		
Clostridium tertium		10.55	0.53		99		
Bifidobacterium adolescentis		1.63	0.48		1		
43377103	**Low discrimination**						**43377103**
					SPO		
Actinomyces israelii		87.80	0.49	URE 1	1		
Clostridium tertium		10.57	0.19		99		
Bifidobacterium adolescentis		1.63	0.13		1		
43377110	**Low discrimination**						**43377110**
					SPO		
Actinomyces israelii		58.20	0.66	MLZ 13	1		
Clostridium tertium		41.66	0.50		99		
Bifidobacterium adolescentis		0.13	0.15		1		
43377112	**Low discrimination**						**43377112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium tertium		55.27	0.58	MNS 99	99		
Actinomyces israelii		38.60	0.69		1		
Bifidobacterium adolescentis		5.87	0.49		1		
43377120	**Low discrimination**						**43377120**
					COC	SPO	
Clostridium tertium		73.13	0.53	MNS 99	1	99	
Actinomyces israelii		25.86	0.60		1	1	
Atopobium parvulum		0.65	0.19		99	1	
43377122	**Low discrimination**						**43377122**
					SPO		
Clostridium tertium		84.46	0.62	MNS 99	99		
Actinomyces israelii		14.93	0.63		1		
Clostridium ramosum		0.48	0.21		92		
43377130	**Very good identification**						**43377130**
					SPO		
Clostridium tertium		97.06	0.58	MNS 99	99		
Actinomyces israelii		2.89	0.46		1		
Bifidobacterium dentium		0.06	-.07		1		
43377132	**Very good identification**						**43377132**
					SPO		
Clostridium tertium		98.31	0.66	MNS 99	99		
Actinomyces israelii		1.46	0.49		1		
Bifidobacterium dentium		0.20	0.10		1		
43377140	**Very good identification**						**43377140**
					SPO		
Actinomyces israelii		99.81	0.74		1		
Clostridium tertium		0.16	0.11		99		
Clostridium butyricum		0.03	-.07		99		
43377141	**Good identification**						**43377141**
Actinomyces israelii		100.0	0.39	URE 1			
43377142	**Very good identification**						**43377142**
					SPO		
Actinomyces israelii		99.11	0.77		1		
Bifidobacterium adolescentis		0.59	0.32		1		
Clostridium tertium		0.31	0.19		99		
43377143	**Good identification**						**43377143**
Actinomyces israelii		100.0	0.42	URE 1			
43377150	**Very good identification**						**43377150**
					COC	SPO	
Actinomyces israelii		97.83	0.59	MLZ 13	1	1	
Clostridium tertium		1.82	0.15		1	99	
Blautia producta		0.35	-.03		99	1	
43377152	**Good identification**						**43377152**
					SPO		
Actinomyces israelii		92.12	0.62	MLZ 13	1		
Bifidobacterium adolescentis		4.45	0.34		1		
Clostridium tertium		3.43	0.24		99		
43377160	**Acceptable identification**						**43377160**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		93.16	0.53	NAG 6	1	
Clostridium tertium		6.84	0.19		99	
43377162	**Low discrimination**					**43377162**
					SPO	
Actinomyces israelii		87.19	0.56	NAG 6	1	
Clostridium tertium		12.81	0.27		99	
43377170	**Low discrimination**					**43377170**
					SPO	
Actinomyces israelii		53.37	0.39	MLZ 13	1	
Clostridium tertium		46.64	0.24	NAG 6	99	
43377200	**Good identification**					**43377200**
Actinomyces israelii		100.0	0.51	RHA 7		
43377202	**Good identification**					**43377202**
Actinomyces israelii		100.0	0.54	RHA 7		
43377210	**Good identification**					**43377210**
Actinomyces israelii		100.0	0.37	RHA 7		
43377212	**Good identification**					**43377212**
Actinomyces israelii		100.0	0.40	RHA 7		
43377240	**Good identification**					**43377240**
Actinomyces israelii		100.0	0.44	RHA 7		
43377242	**Good identification**					**43377242**
Actinomyces israelii		100.0	0.47	RHA 7		
43377300	**Very good identification**					**43377300**
					SPO	
Actinomyces israelii		99.20	0.61	RHA 7	1	
Clostridium tertium		0.80	0.11		99	
43377302	**Very good identification**					**43377302**
					SPO	
Actinomyces israelii		98.17	0.64	RHA 7	1	
Clostridium tertium		1.59	0.19		99	
Bifidobacterium adolescentis		0.25	0.13		1	
43377310	**Acceptable identification**					**43377310**
					SPO	
Actinomyces israelii		91.24	0.47	RHA 7	1	
Clostridium tertium		8.77	0.15	MLZ 13	99	
43377312	**Low discrimination**					**43377312**
					SPO	
Actinomyces israelii		82.47	0.50	RHA 7	1	
Clostridium tertium		15.85	0.24	MLZ 13	99	
Bifidobacterium adolescentis		1.68	0.15		1	
43377320	**Low discrimination**					**43377320**
					SPO	
Actinomyces israelii		62.81	0.41	RHA 7	1	
Clostridium tertium		23.84	0.19	NAG 6	99	
Clostridium ramosum		13.35	0.13		92	
43377322	**Low discrimination**					**43377322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		43.00	0.44	RHA 7	1	
Clostridium tertium		32.63	0.27	NAG 6	99	
Clostridium ramosum		24.37	0.24		92	
43377340	**Good identification**					**43377340**
Actinomyces israelii		100.0	0.54	RHA 7		
43377342	**Very good identification**					**43377342**
Actinomyces israelii		100.0	0.57	RHA 7		
43377350	**Good identification**					**43377350**
					COC	
Actinomyces israelii		95.47	0.40	RHA 7	1	
Blautia producta		4.53	-.03	MLZ 13	99	
43377352	**Good identification**					**43377352**
Actinomyces israelii		100.0	0.43	RHA 7		
43377362	**Good identification**					**43377362**
Actinomyces israelii		100.0	0.37	RHA 7		
43377400	**Very good identification**					**43377400**
					SPO	
Actinomyces israelii		99.72	0.69		1	
Clostridium butyricum		0.28	0.04		99	
43377402	**Very good identification**					**43377402**
					SPO	
Actinomyces israelii		99.32	0.72		1	
Bifidobacterium adolescentis		0.61	0.28		1	
Clostridium butyricum		0.07	-.03		99	
43377403	**Good identification**					**43377403**
Actinomyces israelii		100.0	0.38	URE 1		
43377410	**Good identification**					**43377410**
Actinomyces israelii		100.0	0.55	MLZ 13		
43377412	**Good identification**					**43377412**
Actinomyces israelii		93.10	0.58	MLZ 13		
Bifidobacterium adolescentis		4.68	0.29			
Bifidobacterium dentium		2.23	0.06			
43377420	**Good identification**					**43377420**
Actinomyces israelii		100.0	0.48	NAG 6		
43377422	**Good identification**					**43377422**
Actinomyces israelii		100.0	0.51	NAG 6		
43377432	**Low discrimination**					**43377432**
Actinomyces israelii		60.02	0.37	MLZ 13		
Bifidobacterium dentium		39.98	0.10	NAG 6		
43377440	**Very good identification**					**43377440**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.62			
43377442	**Very good identification**					**43377442**
Actinomyces israelii		99.81	0.65			
Bifidobacterium adolescentis		0.20	0.12			
43377450	**Good identification**					**43377450**
					COC	
Actinomyces israelii		98.29	0.48	MLZ 13	1	
Blautia producta		1.72	-.03		99	
43377452	**Good identification**					**43377452**
Actinomyces israelii		98.43	0.51	MLZ 13		
Bifidobacterium adolescentis		1.57	0.14			
43377460	**Good identification**					**43377460**
Actinomyces israelii		100.0	0.41	NAG 6		
43377462	**Good identification**					**43377462**
Actinomyces israelii		100.0	0.44	NAG 6		
43377500	**Very good identification**					**43377500**
					SPO	
Actinomyces israelii		98.04	0.79		1	
Clostridium butyricum		1.66	0.28		99	
Bifidobacterium adolescentis		0.23	0.28		1	
43377501	**Good identification**					**43377501**
					SPO	
Actinomyces israelii		98.34	0.45	URE 1	1	
Clostridium butyricum		1.66	-.07		99	
43377502	**Low discrimination**					**43377502**
					SPO	
Actinomyces israelii		86.30	0.82		1	
Bifidobacterium adolescentis		13.12	0.62		1	
Clostridium butyricum		0.38	0.21		99	
43377503	**Low discrimination**					**43377503**
Actinomyces israelii		86.80	0.48	URE 1		
Bifidobacterium adolescentis		13.20	0.28			
43377510	**Low discrimination**					**43377510**
Actinomyces israelii		89.04	0.65	MLZ 13		
Bifidobacterium dentium		7.82	0.23			
Bifidobacterium adolescentis		1.68	0.29			
43377512	**Low discrimination**					**43377512**
Bifidobacterium adolescentis		43.56	0.63	MNS 87		
Actinomyces israelii		35.03	0.68	RAF 92		
Bifidobacterium dentium		20.74	0.40			
43377520	**Good identification**					**43377520**
					SPO	
Actinomyces israelii		94.19	0.59	NAG 6	1	
Clostridium tertium		3.29	0.19		99	
Clostridium butyricum		2.17	0.10		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43377522	**Low discrimination**						**43377522**
					SPO		
Actinomyces israelii		88.91	0.62	NAG 6	1		
Clostridium tertium		6.21	0.27		99		
Bifidobacterium dentium		2.21	0.10		1		
43377532	**Low discrimination**						**43377532**
					SPO		
Bifidobacterium dentium		89.05	0.44	MNS 99	1		
Actinomyces israelii		5.40	0.47	RAF 95	1		
Clostridium tertium		4.49	0.32		99		
43377540	**Very good identification**						**43377540**
					COC	SPO	
Actinomyces israelii		97.46	0.72		1	1	
Blautia producta		2.10	0.24		99	1	
Clostridium butyricum		0.37	0.10		1	99	
43377541	**Good identification**						**43377541**
Actinomyces israelii		100.0	0.38	URE 1			
43377542	**Very good identification**						**43377542**
					COC		
Actinomyces israelii		95.16	0.75		1		
Bifidobacterium adolescentis		4.60	0.46		1		
Blautia producta		0.15	0.07		99		
43377543	**Good identification**						**43377543**
Actinomyces israelii		95.39	0.41	URE 1			
Bifidobacterium adolescentis		4.61	0.12				
43377550	**Low discrimination**						**43377550**
					COC		
Actinomyces israelii		69.54	0.58	MLZ 13	1		
Blautia producta		30.04	0.32		99		
Bifidobacterium adolescentis		0.42	0.14		1		
43377552	**Low discrimination**						**43377552**
					COC		
Actinomyces israelii		69.32	0.61	MLZ 13	1		
Bifidobacterium adolescentis		27.40	0.48		1		
Blautia producta		2.22	0.15		99		
43377560	**Good identification**						**43377560**
					COC		
Actinomyces israelii		96.39	0.52	NAG 6	1		
Blautia producta		3.61	0.07		99		
43377562	**Good identification**						**43377562**
Actinomyces israelii		99.24	0.55	NAG 6			
Bifidobacterium adolescentis		0.76	0.12				
43377570	**Low discrimination**						**43377570**
					COC		
Actinomyces israelii		55.08	0.37	MLZ 13	1		
Blautia producta		41.42	0.15	NAG 6	99		
Bifidobacterium dentium		3.50	-.07		1		
43377572	**Low discrimination**						**43377572**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		64.66	0.40	MLZ 13		
Bifidobacterium dentium		27.69	0.10	NAG 6		
Bifidobacterium adolescentis		4.04	0.14			
43377600	**Good identification**					**43377600**
Actinomyces israelii		100.0	0.50	RHA 7		
43377602	**Good identification**					**43377602**
Actinomyces israelii		100.0	0.53	RHA 7		
43377610	**Good identification**					**43377610**
Actinomyces israelii		100.0	0.35	RHA 7		
43377612	**Good identification**					**43377612**
Actinomyces israelii		100.0	0.38	RHA 7		
43377640	**Good identification**					**43377640**
Actinomyces israelii		100.0	0.42	RHA 7		
43377642	**Good identification**					**43377642**
Actinomyces israelii		100.0	0.45	RHA 7		
43377700	**Very good identification**					**43377700**
					SPO	
Actinomyces israelii		99.31	0.60	RHA 7	1	
Clostridium butyricum		0.69	0.02		99	
43377702	**Very good identification**					**43377702**
					SPO	
Actinomyces israelii		97.83	0.63	RHA 7	1	
Bifidobacterium adolescentis		2.00	0.28		1	
Clostridium butyricum		0.18	-.05		99	
43377710	**Good identification**					**43377710**
					COC	
Actinomyces israelii		97.80	0.46	RHA 7	1	
Blautia producta		2.21	-.03	MLZ 13	99	
43377712	**Low discrimination**					**43377712**
Actinomyces israelii		78.77	0.49	RHA 7		
Bifidobacterium adolescentis		13.15	0.29	MLZ 13		
Bifidobacterium dentium		6.26	0.06			
43377720	**Good identification**					**43377720**
Actinomyces israelii		100.0	0.39	RHA 7		
43377722	**Good identification**					**43377722**
					GRA	
Actinomyces israelii		96.62	0.42	RHA 7	99	
Bacteroides ovatus		3.38	-.02	NAG 6	1	
43377740	**Low discrimination**					**43377740**
					COC	
Actinomyces israelii		77.77	0.53	RHA 7	1	
Blautia producta		22.23	0.24		99	
43377742	**Good identification**					**43377742**
					COC	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		97.31	0.56	RHA 7	1			
Blautia producta		2.06	0.07		99			
Bifidobacterium adolescentis		0.63	0.12		1			
43377752	**Low discrimination**							**43377752**
					COC			
Actinomyces israelii		67.65	0.42	RHA 7	1			
Blautia producta		28.76	0.15	MLZ 13	99			
Bifidobacterium adolescentis		3.59	0.14		1			
43377762	**Good identification**							**43377762**
Actinomyces israelii		100.0	0.35	RHA 7				
43400000	**Low discrimination**							**43400000**
					GRA	SPO		
Clostridium botulinum biovar A		68.23	0.47	SUC 7	99	99		
Fusobacterium mortiferum		11.76	0.47	RAF 1	1	1		
Atopobium minutum		7.28	0.38		99	1		
43400040	**Good identification**							**43400040**
					COC	GRA		
Clostridium botulinum biovar A		97.80	0.38	SUC 7	1	99		
Clostridium novyi biovar A		1.06	0.16	RAF 1	1	99		
Megasphaera elsdenii		0.58	0.04		99	1		
43400100	**Good identification**							**43400100**
					COC	GRA	SPO	
Fusobacterium mortiferum		96.99	0.53	FRU 99	1	1	1	
Clostridium botulinum biovar A		2.44	0.13		1	99	99	
Blautia hansenii		0.58	-.03		99	99	1	
43400102	**Good identification**							**43400102**
Fusobacterium mortiferum		100.0	0.37	FRU 99				
43401000	**Very good identification**							**43401000**
					GRA			
Eubacterium tortuosum		95.14	0.58	SAL 99	99			
Fusobacterium mortiferum		3.44	0.50		1			
Eubacterium saburreum		0.73	0.37		85			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43401002	**Good identification**						**43401002**
					GRA		
Eubacterium tortuosum		95.04	0.45	SAL 99	99		
Fusobacterium mortiferum		2.16	0.33		1		
Atopobium minutum		1.79	0.27		99		
43401020	**Good identification**						**43401020**
Eubacterium tortuosum		97.97	0.45	SAL 99			
Atopobium minutum		0.92	0.21				
Eubacterium saburreum		0.75	0.24				
43401100	**Low discrimination**						**43401100**
					COC	GRA	
Fusobacterium mortiferum		57.47	0.56	FRU 99	1	1	
Blautia hansenii		22.52	0.32		99	99	
Eubacterium saburreum		12.17	0.43		1	85	
43401102	**Low discrimination**						**43401102**
					COC	GRA	
Fusobacterium mortiferum		49.72	0.40	FRU 99	1	1	
Blautia hansenii		19.48	0.15	BGL 10	99	99	
Eubacterium saburreum		16.72	0.30		1	85	
43401110	**Acceptable identification**						**43401110**
					COC	GRA	
Eubacterium saburreum		92.49	0.41	IND 99	1	85	
Fusobacterium mortiferum		5.39	0.22		1	1	
Blautia hansenii		2.11	-.03		99	99	
43405100	**Low discrimination**						**43405100**
					COC	GRA	
Eubacterium saburreum		42.12	0.38	IND 99	1	85	
Fusobacterium mortiferum		27.81	0.37		1	1	
Blautia hansenii		25.55	0.19		99	99	
43405110	**Good identification**						**43405110**
Eubacterium saburreum		100.0	0.37	IND 99			
43405530	**Good identification**						**43405530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43405532	**Good identification**						**43405532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43410000	**Very good identification**						**43410000**
					GRA	SPO	
Fusobacterium mortiferum		98.34	0.81		1	1	
Atopobium minutum		0.57	0.37		99	1	
Clostridium botulinum biovar A		0.37	0.26		99	99	
43410001	**Good identification**						**43410001**
Fusobacterium mortiferum		100.0	0.47	URE 1			
43410002	**Very good identification**						**43410002**
					GRA	SPO	
Fusobacterium mortiferum		95.77	0.65	BGL 10	1	1	
Atopobium minutum		3.31	0.34		99	1	
Clostridium sporogenes		0.29	0.06		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43410010	**Acceptable identification**							**43410010**
					GRA	SPO	CAT	
Fusobacterium mortiferum		93.14	0.47	MLZ 1	1	1	5	
Clostridium difficile		5.98	0.10		99	99	1	
Propionibacterium granulosum		0.88	0.09		99	1	90	
43410020	**Low discrimination**							**43410020**
					GRA	CAT		
Fusobacterium mortiferum		82.81	0.47	NAG 1	1	5		
Atopobium minutum		15.76	0.29		99	1		
Propionibacterium granulosum		1.43	0.13		99	90		
43410040	**Low discrimination**							**43410040**
					COC	GRA	SPO	
Fusobacterium mortiferum		74.15	0.47	SOR 1	1	1	1	
Megasphaera elsdenii		14.41	0.26		99	1	1	
Clostridium botulinum biovar A		8.20	0.17		1	99	99	
43410100	**Very good identification**							**43410100**
					GRA			
Fusobacterium mortiferum		99.91	0.87		1			
Actinomyces meyeri		0.06	0.23		99			
Lactobacillus fermentum		0.02	0.24		99			
43410101	**Good identification**							**43410101**
Fusobacterium mortiferum		100.0	0.53	URE 1				
43410102	**Very good identification**							**43410102**
					GRA			
Fusobacterium mortiferum		99.86	0.71	BGL 10	1			
Lactobacillus fermentum		0.14	0.24		99			
43410103	**Good identification**							**43410103**
Fusobacterium mortiferum		100.0	0.37	URE 1				
43410110	**Good identification**							**43410110**
Fusobacterium mortiferum		99.22	0.53	MLZ 1				
Eubacterium saburreum		0.78	0.15					
43410112	**Good identification**							**43410112**
Fusobacterium mortiferum		100.0	0.37	MLZ 1				
43410120	**Acceptable identification**							**43410120**
					GRA			
Fusobacterium mortiferum		94.46	0.53	NAG 1	1			
Prevotella melaninogenica		4.27	0.12		1			
Actinomyces meyeri		0.78	0.08		99			
43410122	**Low discrimination**							**43410122**
					CAT			
Fusobacterium mortiferum		73.09	0.37	NAG 1	5			
Capnocytophaga ochracea		17.85	0.12	BGL 10	90			
Prevotella melaninogenica		5.66	0.00		1			
43410140	**Good identification**							**43410140**
Fusobacterium mortiferum		100.0	0.53	SOR 1				
43410142	**Good identification**							**43410142**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.37	SOR 1			
43410200	**Good identification**						**43410200**
Fusobacterium mortiferum		100.0	0.47	RHA 1			
43410300	**Good identification**						**43410300**
Fusobacterium mortiferum		100.0	0.53	RHA 1			
43410302	**Good identification**						**43410302**
Fusobacterium mortiferum		100.0	0.37	RHA 1			
43410400	**Good identification**						**43410400**
					GRA		
Fusobacterium mortiferum		96.99	0.47	ARA 1	1		
Lactobacillus fermentum		2.15	0.20		99		
Actinomyces meyeri		0.86	0.02		99		
43410500	**Good identification**						**43410500**
					GRA		
Fusobacterium mortiferum		97.03	0.53	ARA 1	1		
Lactobacillus fermentum		1.50	0.24		99		
Actinomyces meyeri		1.47	0.13		99		
43410502	**Low discrimination**						**43410502**
					GRA		
Fusobacterium mortiferum		87.77	0.37	ARA 1	1		
Lactobacillus fermentum		12.24	0.24	BGL 10	99		
43411000	**Very good identification**						**43411000**
					GRA		
Fusobacterium mortiferum		97.45	0.84		1		
Lactobacillus fermentum		1.44	0.54		99		
Eubacterium saburreum		0.63	0.45		85		
43411001	**Good identification**						**43411001**
					GRA		
Fusobacterium mortiferum		97.93	0.50	URE 1	1		
Lactobacillus fermentum		1.45	0.20		99		
Eubacterium saburreum		0.63	0.11		85		
43411002	**Low discrimination**						**43411002**
					GRA		
Fusobacterium mortiferum		86.48	0.68	BGL 10	1		
Lactobacillus fermentum		11.49	0.54		99		
Eubacterium saburreum		0.88	0.32		85		
43411010	**Low discrimination**						**43411010**
					GRA		
Fusobacterium mortiferum		64.90	0.50	MLZ 1	1		
Eubacterium saburreum		33.73	0.43		85		
Lactobacillus fermentum		0.96	0.20		99		
43411020	**Low discrimination**						**43411020**
					GRA		
Fusobacterium mortiferum		56.41	0.50	NAG 1	1		
Lactobacillus fermentum		27.49	0.46		99		
Eubacterium saburreum		6.32	0.32		85		
43411022	**Low discrimination**						**43411022**
					GRA	CAT	
Lactobacillus fermentum		67.99	0.46	ESL 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		15.50	0.33		1	5	
Capnocytophaga ochracea		5.89	0.12		1	90	
43411040	**Good identification**						**43411040**
					GRA		
Fusobacterium mortiferum		97.93	0.50	SOR 1	1		
Lactobacillus fermentum		1.45	0.20		99		
Eubacterium saburreum		0.63	0.11		85		
43411100	**Very good identification**						**43411100**
					GRA		
Fusobacterium mortiferum		98.24	0.90		1		
Lactobacillus fermentum		1.01	0.58		99		
Eubacterium saburreum		0.63	0.51		85		
43411101	**Good identification**						**43411101**
					GRA	SPO	
Fusobacterium mortiferum		96.60	0.56	URE 1	1	1	
Clostridium chauvoei		1.27	0.12		99	99	
Lactobacillus fermentum		1.00	0.24		99	1	
43411102	**Low discrimination**						**43411102**
					GRA	CAT	
Fusobacterium mortiferum		89.11	0.74	BGL 10	1	5	
Lactobacillus fermentum		8.28	0.58		99	1	
Capnocytophaga ochracea		1.61	0.30		1	90	
43411103	**Low discrimination**						**43411103**
					GRA		
Fusobacterium mortiferum		86.12	0.40	URE 1	1		
Lactobacillus fermentum		8.00	0.24	BGL 10	99		
Actinomyces naeslundii		4.08	0.29		99		
43411110	**Low discrimination**						**43411110**
					GRA		
Fusobacterium mortiferum		65.25	0.56	MLZ 1	1		
Eubacterium saburreum		33.91	0.49		85		
Lactobacillus fermentum		0.67	0.24		99		
43411112	**Low discrimination**						**43411112**
					CAT		
Fusobacterium mortiferum		47.23	0.40	MLZ 1	5		
Eubacterium saburreum		38.98	0.37	BGL 10	1		
Capnocytophaga ochracea		9.40	0.14		90		
43411120	**Low discrimination**						**43411120**
					GRA		
Prevotella melaninogenica		64.10	0.47	MNS 90	1		
Fusobacterium mortiferum		21.50	0.56	MLT 99	1		
Lactobacillus fermentum		7.33	0.50		99		
43411122	**Low discrimination**						**43411122**
					GRA	CAT	
Capnocytophaga ochracea		61.58	0.47	MNS 90	1	90	
Prevotella melaninogenica		19.53	0.34	MLT 99	1	1	
Lactobacillus fermentum		11.71	0.50		99	1	
43411130	**Low discrimination**						**43411130**
					CAT		
Eubacterium saburreum		56.80	0.37	IND 99	1		
Prevotella melaninogenica		18.65	0.12		1		
Capnocytophaga ochracea		13.70	0.14		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43411140	**Good identification**						**43411140**
					GRA		
Fusobacterium mortiferum		98.12	0.56	SOR 1	1		
Lactobacillus fermentum		1.01	0.24		99		
Eubacterium saburreum		0.63	0.17		85		
43411142	**Low discrimination**						**43411142**
					GRA	CAT	
Fusobacterium mortiferum		77.40	0.40	SOR 1	1	5	
Capnocytophaga ochracea		15.41	0.14	BGL 10	1	90	
Lactobacillus fermentum		7.19	0.24		99	1	
43411200	**Good identification**						**43411200**
					GRA		
Fusobacterium mortiferum		97.93	0.50	RHA 1	1		
Lactobacillus fermentum		1.45	0.20		99		
Eubacterium saburreum		0.63	0.11		85		
43411300	**Good identification**						**43411300**
					GRA		
Fusobacterium mortiferum		98.35	0.56	RHA 1	1		
Lactobacillus fermentum		1.02	0.24		99		
Eubacterium saburreum		0.63	0.17		85		
43411302	**Acceptable identification**						**43411302**
					GRA		
Fusobacterium mortiferum		91.50	0.40	RHA 1	1		
Lactobacillus fermentum		8.50	0.24	BGL 10	99		
43411400	**Low discrimination**						**43411400**
					GRA		
Lactobacillus fermentum		59.02	0.54	ESL 1	99		
Fusobacterium mortiferum		40.37	0.50		1		
Actinomyces meyeri		0.36	0.05		99		
43411402	**Acceptable identification**						**43411402**
					GRA		
Lactobacillus fermentum		92.94	0.54	ESL 1	99		
Fusobacterium mortiferum		7.06	0.33		1		
43411420	**Good identification**						**43411420**
					GRA		
Lactobacillus fermentum		96.90	0.46	ESL 1	99		
Fusobacterium mortiferum		2.01	0.16		1		
Bifidobacterium longum subsp. longum		1.09	0.05		99		
43411422	**Good identification**						**43411422**
					GRA	CAT	
Lactobacillus fermentum		97.42	0.46	ESL 1	99	1	
Capnocytophaga ochracea		1.49	-.01		1	90	
Bifidobacterium longum subsp. longum		1.09	0.05		99	1	
43411500	**Low discrimination**						**43411500**
					GRA		
Lactobacillus fermentum		48.55	0.58	ESL 1	99		
Fusobacterium mortiferum		47.49	0.56		1		
Bifidobacterium longum subsp. longum		2.75	0.29		99		
43411502	**Low discrimination**						**43411502**
					GRA		
Lactobacillus fermentum		82.70	0.58	ESL 1	99		
Fusobacterium mortiferum		8.99	0.40		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		4.69	0.29		99		
43411520	**Low discrimination**						**43411520**
					GRA	CAT	
Lactobacillus fermentum		52.59	0.50	ESL 1	99	1	
Bifidobacterium longum subsp. longum		35.78	0.40		99	1	
Capnocytophaga ochracea		5.42	0.17		1	90	
43411522	**Low discrimination**						**43411522**
					GRA	CAT	
Lactobacillus fermentum		37.60	0.50	ESL 1	99	1	
Capnocytophaga ochracea		34.88	0.34		1	90	
Bifidobacterium longum subsp. longum		25.58	0.40		99	1	
43411530	**Good identification**						**43411530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.17	0.44	MNS 87	99	1	
Capnocytophaga ochracea		0.97	0.01	MLT 99	1	90	
Lactobacillus fermentum		0.86	0.15		99	1	
43411532	**Low discrimination**						**43411532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.77	0.44	MNS 87	99	1	
Capnocytophaga ochracea		7.99	0.17	MLT 99	1	90	
Parabacteroides distasonis		1.46	0.17		1	76	
43412000	**Low discrimination**						**43412000**
					COC	GRA	
Fusobacterium mortiferum		41.32	0.47	MAN 1	1	1	
Eubacterium limosum		39.18	0.29		1	99	
Megasphaera elsdenii		12.04	0.29		99	1	
43412100	**Good identification**						**43412100**
					GRA		
Fusobacterium mortiferum		99.25	0.53	MAN 1	1		
Eubacterium limosum		0.41	-.06		99		
Propionibacterium propionicum		0.34	0.12		99		
43412102	**Good identification**						**43412102**
					GRA		
Fusobacterium mortiferum		95.56	0.37	MAN 1	1		
Propionibacterium propionicum		4.44	0.15	BGL 10	99		
43413000	**Acceptable identification**						**43413000**
					GRA	CAT	
Fusobacterium mortiferum		93.65	0.50	MAN 1	1	5	
Propionibacterium granulosum		3.25	0.21		99	90	
Lactobacillus fermentum		1.38	0.20		99	1	
43413100	**Good identification**						**43413100**
					GRA		
Fusobacterium mortiferum		97.48	0.56	MAN 1	1		
Lactobacillus fermentum		1.01	0.24		99		
Actinomyces naeslundii		0.66	0.31		99		
43413102	**Low discrimination**						**43413102**
					GRA		
Fusobacterium mortiferum		84.37	0.40	MAN 1	1		
Lactobacillus fermentum		7.84	0.24	BGL 10	99		
Actinomyces naeslundii		5.17	0.31		99		
43414000	**Very good identification**						**43414000**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		98.35	0.62	XYL 7	1	1	
Actinomyces meyeri		0.91	0.18		99	1	
Clostridium glycolicum		0.58	0.09		99	99	
43414002	**Good identification**						**43414002**
					GRA		
Fusobacterium mortiferum		98.45	0.45	XYL 7	1		
Lactobacillus fermentum		1.55	0.16	BGL 10	99		
43414100	**Very good identification**						**43414100**
					GRA		
Fusobacterium mortiferum		98.26	0.68	XYL 7	1		
Actinomyces meyeri		1.55	0.28		99		
Lactobacillus fermentum		0.12	0.20		99		
43414102	**Good identification**						**43414102**
					GRA		
Fusobacterium mortiferum		98.49	0.52	XYL 7	1		
Lactobacillus fermentum		1.08	0.20	BGL 10	99		
Actinomyces meyeri		0.43	0.02		99		
43415000	**Low discrimination**						**43415000**
					GRA		
Fusobacterium mortiferum		85.45	0.65	XYL 7	1		
Lactobacillus fermentum		9.85	0.50		99		
Eubacterium saburreum		3.92	0.40		85		
43415002	**Low discrimination**						**43415002**
					GRA		
Lactobacillus fermentum		49.15	0.50	ESL 1	99		
Fusobacterium mortiferum		47.40	0.48		1		
Eubacterium saburreum		3.46	0.27		85		
43415010	**Low discrimination**						**43415010**
					GRA		
Eubacterium saburreum		76.93	0.39	IND 99	85		
Fusobacterium mortiferum		20.69	0.30		1		
Lactobacillus fermentum		2.38	0.16		99		
43415020	**Low discrimination**						**43415020**
					GRA		
Lactobacillus fermentum		66.38	0.42	ESL 1	99		
Fusobacterium mortiferum		17.46	0.30		1		
Eubacterium saburreum		14.00	0.27		85		
43415022	**Low discrimination**						**43415022**
					CAT	GRA	
Lactobacillus fermentum		85.25	0.42	ESL 1	1	99	
Bacteroides fragilis		5.94	-.03		99	1	
Eubacterium saburreum		3.17	0.14		1	85	
43415100	**Low discrimination**						**43415100**
					GRA		
Fusobacterium mortiferum		87.49	0.71	XYL 7	1		
Lactobacillus fermentum		7.05	0.54		99		
Eubacterium saburreum		4.02	0.46		85		
43415101	**Low discrimination**						**43415101**
					GRA		
Fusobacterium mortiferum		88.77	0.37	XYL 7	1		
Lactobacillus fermentum		7.15	0.20	URE 1	99		
Eubacterium saburreum		4.08	0.12		85		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43415102	**Low discrimination**						**43415102**
					GRA		
Fusobacterium mortiferum		53.06	0.55	XYL 7	1		
Lactobacillus fermentum		38.48	0.54	BGL 10	99		
Eubacterium saburreum		3.87	0.33		85		
43415110	**Low discrimination**						**43415110**
					GRA		
Eubacterium saburreum		76.84	0.45	IND 99	85		
Fusobacterium mortiferum		20.67	0.37		1		
Lactobacillus fermentum		1.67	0.20		99		
43415120	**Low discrimination**						**43415120**
					GRA		
Lactobacillus fermentum		41.03	0.46	ESL 1	99		
Fusobacterium mortiferum		15.43	0.37		1		
Eubacterium saburreum		12.37	0.33		85		
43415132	**Low discrimination**						**43415132**
					GRA	CAT	
Parabacteroides distasonis		54.39	0.37	MNS 98	1	76	
Capnocytophaga ochracea		28.73	0.20		1	90	
Bifidobacterium longum subsp. longum		6.91	0.18		99	1	
43415140	**Low discrimination**						**43415140**
					GRA		
Fusobacterium mortiferum		88.77	0.37	XYL 7	1		
Lactobacillus fermentum		7.15	0.20	SOR 1	99		
Eubacterium saburreum		4.08	0.12		85		
43415322	**Good identification**						**43415322**
Parabacteroides distasonis		96.94	0.50	MNS 98			
Bacteroides fragilis		1.58	-.03				
Capnocytophaga ochracea		0.84	0.02				
43415332	**Good identification**						**43415332**
Parabacteroides distasonis		100.0	0.45	MNS 98			
43415400	**Acceptable identification**						**43415400**
					GRA		
Lactobacillus fermentum		90.28	0.50	ESL 1	99		
Fusobacterium mortiferum		7.92	0.30		1		
Actinomyces meyeri		1.80	0.10		99		
43415402	**Good identification**						**43415402**
					GRA		
Lactobacillus fermentum		99.04	0.50	ESL 1	99		
Fusobacterium mortiferum		0.97	0.14		1		
43415420	**Acceptable identification**						**43415420**
Lactobacillus fermentum		94.58	0.42	ESL 1			
Bifidobacterium longum subsp. longum		5.42	0.14				
43415422	**Acceptable identification**						**43415422**
Lactobacillus fermentum		94.58	0.42	ESL 1			
Bifidobacterium longum subsp. longum		5.42	0.14				
43415500	**Low discrimination**						**43415500**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		68.04	0.54	ESL 1	99		
Bifidobacterium longum subsp. longum		19.70	0.37		99		
Fusobacterium mortiferum		8.53	0.37		1		
43415502	**Low discrimination**						**43415502**
Lactobacillus fermentum		70.13	0.54	ESL 1			
Bifidobacterium longum subsp. longum		20.31	0.37				
Bifidobacterium adolescentis		7.56	0.32				
43415510	**Good identification**						**43415510**
Bifidobacterium longum subsp. longum		97.08	0.41	MNS 87			
Lactobacillus fermentum		1.99	0.20	MLT 99			
Eubacterium saburreum		0.93	0.11				
43415512	**Low discrimination**						**43415512**
Bifidobacterium longum subsp. longum		77.67	0.41	MNS 87			
Bifidobacterium adolescentis		20.74	0.33	MLT 99			
Lactobacillus fermentum		1.59	0.20				
43415520	**Low discrimination**						**43415520**
					GRA		
Bifidobacterium longum subsp. longum		71.79	0.48	MNS 87	99		
Lactobacillus fermentum		20.66	0.46	MLT 99	99		
Bacteroides vulgatus		6.24	0.15		1		
43415522	**Low discrimination**						**43415522**
					GRA		
Bifidobacterium longum subsp. longum		63.78	0.48	MNS 87	99		
Lactobacillus fermentum		18.35	0.46	MLT 99	99		
Parabacteroides distasonis		9.83	0.38		1		
43415530	**Good identification**						**43415530**
Bifidobacterium longum subsp. longum		100.0	0.52	MNS 87			
43415532	**Acceptable identification**						**43415532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		94.68	0.52	MNS 87	99	1	
Parabacteroides distasonis		4.62	0.33	MLT 99	1	76	
Capnocytophaga ochracea		0.70	0.07		1	90	
43415722	**Good identification**						**43415722**
					GRA		
Parabacteroides distasonis		95.44	0.46	MNS 98	1		
Bifidobacterium longum subsp. longum		2.08	0.14		99		
Bacteroides vulgatus		1.63	-.03		1		
43415732	**Acceptable identification**						**43415732**
					GRA		
Parabacteroides distasonis		93.54	0.42	MNS 98	1		
Bifidobacterium longum subsp. longum		6.46	0.18		99		
43417100	**Low discrimination**						**43417100**
					GRA		
Fusobacterium mortiferum		88.77	0.37	MAN 1	1		
Lactobacillus fermentum		7.15	0.20	XYL 7	99		
Eubacterium saburreum		4.08	0.12		85		
43420000	**Low discrimination**						**43420000**
					GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		74.76	0.36	SUC 7	99	99		
Fusobacterium mortiferum		15.40	0.38	RAF 1	1	1		
Clostridium sporogenes		9.84	0.19		99	99		
43420100	**Good identification**							**43420100**
					GRA	SPO		
Fusobacterium mortiferum		97.94	0.44	FRU 99	1	1		
Clostridium botulinum biovar A		2.06	0.02		99	99		
43421000	**Low discrimination**							**43421000**
					GRA			
Fusobacterium mortiferum		40.98	0.41	FRU 99	1			
Eubacterium tortuosum		38.32	0.24		99			
Eubacterium saburreum		19.36	0.34		85			
43421100	**Low discrimination**							**43421100**
					COC	GRA		
Fusobacterium mortiferum		67.30	0.47	FRU 99	1	1		
Eubacterium saburreum		31.80	0.40		1	85		
Blautia hansenii		0.89	-.03		99	99		
43421110	**Good identification**							**43421110**
Eubacterium saburreum		97.45	0.38	IND 99				
Fusobacterium mortiferum		2.55	0.13					
43425100	**Low discrimination**							**43425100**
Eubacterium saburreum		77.17	0.35	IND 99				
Fusobacterium mortiferum		22.83	0.28					
43430000	**Very good identification**							**43430000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		98.89	0.72		1	1	5	
Propionibacterium granulosum		0.69	0.32		99	1	90	
Clostridium botulinum biovar A		0.31	0.16		99	99	1	
43430001	**Good identification**							**43430001**
Fusobacterium mortiferum		100.0	0.38	URE 1				
43430002	**Good identification**							**43430002**
					GRA	SPO	CAT	
Fusobacterium mortiferum		98.12	0.56	BGL 10	1	1	5	
Propionibacterium granulosum		1.54	0.21		99	1	90	
Clostridium sporogenes		0.34	-.02		99	99	1	
43430010	**Acceptable identification**							**43430010**
					GRA	CAT		
Fusobacterium mortiferum		91.37	0.38	MLZ 1	1	5		
Propionibacterium granulosum		8.63	0.17		99	90		
43430020	**Low discrimination**							**43430020**
					GRA	CAT		
Fusobacterium mortiferum		85.24	0.38	NAG 1	1	5		
Propionibacterium granulosum		14.76	0.21		99	90		
43430040	**Acceptable identification**							**43430040**
					GRA	SPO		
Fusobacterium mortiferum		91.53	0.38	SOR 1	1	1		
Clostridium botulinum biovar A		8.47	0.07		99	99		
43430100	**Very good identification**							**43430100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		99.98	0.78				
Eubacterium saburreum		0.02	0.14				
43430101	**Good identification**						**43430101**
Fusobacterium mortiferum		100.0	0.44	URE 1			
43430102	**Very good identification**						**43430102**
Fusobacterium mortiferum		100.0	0.62	BGL 10			
43430110	**Good identification**						**43430110**
Fusobacterium mortiferum		98.27	0.44	MLZ 1			
Eubacterium saburreum		1.73	0.12				
43430120	**Good identification**						**43430120**
Fusobacterium mortiferum		100.0	0.44	NAG 1			
43430140	**Good identification**						**43430140**
Fusobacterium mortiferum		100.0	0.44	SOR 1			
43430200	**Good identification**						**43430200**
Fusobacterium mortiferum		100.0	0.38	RHA 1			
43430300	**Good identification**						**43430300**
Fusobacterium mortiferum		100.0	0.44	RHA 1			
43430400	**Good identification**						**43430400**
Fusobacterium mortiferum		100.0	0.38	ARA 1			
43430500	**Good identification**						**43430500**
Fusobacterium mortiferum		100.0	0.44	ARA 1			
43431000	**Very good identification**						**43431000**
					CAT	GRA	
Fusobacterium mortiferum		98.07	0.75		5	1	
Eubacterium saburreum		1.40	0.42		1	85	
Propionibacterium granulosum		0.46	0.32		90	99	
43431001	**Good identification**						**43431001**
					GRA		
Fusobacterium mortiferum		97.27	0.41	URE 1	1		
Actinomyces naeslundii		2.73	0.26		99		
43431002	**Very good identification**						**43431002**
					CAT	GRA	
Fusobacterium mortiferum		96.21	0.59	BGL 10	5	1	
Eubacterium saburreum		2.19	0.29		1	85	
Propionibacterium granulosum		1.01	0.21		90	99	
43431010	**Low discrimination**						**43431010**
					CAT	GRA	
Eubacterium saburreum		52.18	0.40	IND 99	1	85	
Fusobacterium mortiferum		44.99	0.41		5	1	
Propionibacterium granulosum		2.83	0.17		90	99	
43431020	**Low discrimination**						**43431020**
					CAT	GRA	
Fusobacterium mortiferum		72.24	0.41	NAG 1	5	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		18.07	0.29		1	85	
Propionibacterium granulosum		8.34	0.21		90	99	
43431040	**Good identification**						**43431040**
Fusobacterium mortiferum		100.0	0.41	SOR 1			
43431100	**Very good identification**						**43431100**
					GRA		
Fusobacterium mortiferum		98.53	0.81		1		
Eubacterium saburreum		1.41	0.48		85		
Lactobacillus fermentum		0.03	0.24		99		
43431101	**Good identification**						**43431101**
					GRA		
Fusobacterium mortiferum		95.37	0.47	URE 1	1		
Actinomyces naeslundii		3.26	0.34		99		
Eubacterium saburreum		1.37	0.14		85		
43431102	**Very good identification**						**43431102**
					GRA		
Fusobacterium mortiferum		97.28	0.65	BGL 10	1		
Eubacterium saburreum		2.21	0.35		85		
Lactobacillus fermentum		0.31	0.24		99		
43431110	**Low discrimination**						**43431110**
Eubacterium saburreum		53.70	0.46	IND 99			
Fusobacterium mortiferum		46.30	0.47				
43431120	**Low discrimination**						**43431120**
Fusobacterium mortiferum		70.87	0.47	NAG 1			
Eubacterium saburreum		17.73	0.35				
Prevotella melaninogenica		7.15	0.12				
43431122	**Low discrimination**						**43431122**
					CAT		
Parabacteroides distasonis		50.54	0.37	MNS 98	76		
Fusobacterium mortiferum		18.95	0.31	XYL 90	5		
Capnocytophaga ochracea		10.33	0.12		90		
43431140	**Good identification**						**43431140**
Fusobacterium mortiferum		98.59	0.47	SOR 1			
Eubacterium saburreum		1.41	0.14				
43431200	**Good identification**						**43431200**
Fusobacterium mortiferum		100.0	0.41	RHA 1			
43431300	**Good identification**						**43431300**
Fusobacterium mortiferum		98.59	0.47	RHA 1			
Eubacterium saburreum		1.41	0.14				
43431322	**Good identification**						**43431322**
Parabacteroides distasonis		100.0	0.45	MNS 98			
43431332	**Good identification**						**43431332**
Parabacteroides distasonis		100.0	0.40	MNS 98			
43431400	**Good identification**						**43431400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		95.29	0.41	ARA 1	1		
Lactobacillus fermentum		4.71	0.20		99		
43431500	**Acceptable identification**						**43431500**
					GRA		
Fusobacterium mortiferum		91.13	0.47	ARA 1	1		
Bifidobacterium longum subsp. longum		4.42	0.19		99		
Lactobacillus fermentum		3.15	0.24		99		
43431722	**Good identification**						**43431722**
Parabacteroides distasonis		100.0	0.41	MNS 98			
43431732	**Good identification**						**43431732**
Parabacteroides distasonis		100.0	0.37	MNS 98			
43432000	**Low discrimination**						**43432000**
					GRA	CAT	
Fusobacterium mortiferum		63.12	0.38	MAN 1	1	5	
Propionibacterium granulosum		32.97	0.30		99	90	
Eubacterium limosum		2.02	-.06		99	1	
43432100	**Good identification**						**43432100**
Fusobacterium mortiferum		100.0	0.44	MAN 1			
43433000	**Low discrimination**						**43433000**
					GRA	CAT	
Fusobacterium mortiferum		72.23	0.41	MAN 1	1	5	
Propionibacterium granulosum		25.15	0.30		99	90	
Actinomyces naeslundii		2.62	0.28		99	25	
43433100	**Acceptable identification**						**43433100**
					GRA		
Fusobacterium mortiferum		94.47	0.47	MAN 1	1		
Actinomyces naeslundii		4.18	0.35		99		
Eubacterium saburreum		1.35	0.14		85		
43433101	**Low discrimination**						**43433101**
					GRA		
Actinomyces naeslundii		86.79	0.38	ESL 12	99		
Fusobacterium mortiferum		13.21	0.13	MNS 88	1		
43433103	**Good identification**						**43433103**
Actinomyces naeslundii		100.0	0.38	ESL 12			
43433120	**Low discrimination**						**43433120**
					GRA		
Actinomyces naeslundii		81.42	0.35	ESL 12	99		
Fusobacterium mortiferum		18.58	0.13	MNS 88	1		
43433121	**Good identification**						**43433121**
Actinomyces naeslundii		100.0	0.38	ESL 12			
43433122	**Good identification**						**43433122**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43433123	**Good identification**						**43433123**
Actinomyces naeslundii		100.0	0.38	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43434000	**Good identification**					**43434000**
Fusobacterium mortiferum		100.0	0.53	XYL 7		
43434002	**Good identification**					**43434002**
Fusobacterium mortiferum		100.0	0.36	XYL 7		
43434100	**Very good identification**					**43434100**
Fusobacterium mortiferum		99.85	0.59	XYL 7		
Eubacterium saburreum		0.16	0.09			
43434102	**Good identification**					**43434102**
Fusobacterium mortiferum		100.0	0.43	XYL 7		
43435000	**Acceptable identification**					**43435000**
					GRA	
Fusobacterium mortiferum		90.39	0.56	XYL 7	1	
Eubacterium saburreum		9.26	0.37		85	
Lactobacillus fermentum		0.35	0.16		99	
43435002	**Low discrimination**					**43435002**
					GRA	
Fusobacterium mortiferum		83.49	0.39	XYL 7	1	
Eubacterium saburreum		13.58	0.24	BGL 10	85	
Lactobacillus fermentum		2.93	0.16		99	
43435010	**Low discrimination**					**43435010**
Eubacterium saburreum		89.24	0.36	IND 99		
Fusobacterium mortiferum		10.76	0.22			
43435100	**Good identification**					**43435100**
					GRA	
Fusobacterium mortiferum		90.48	0.62	XYL 7	1	
Eubacterium saburreum		9.27	0.43		85	
Lactobacillus fermentum		0.25	0.20		99	
43435102	**Low discrimination**					**43435102**
Fusobacterium mortiferum		80.27	0.46	XYL 7		
Eubacterium saburreum		13.06	0.30	BGL 10		
Parabacteroides distasonis		2.61	0.19			
43435110	**Low discrimination**					**43435110**
Eubacterium saburreum		89.25	0.42	IND 99		
Fusobacterium mortiferum		10.76	0.28			
43435122	**Good identification**					**43435122**
					CAT	
Parabacteroides distasonis		97.23	0.53	MNS 98	76	
Bacteroides fragilis		1.04	-.03		99	
Eubacterium saburreum		0.87	0.18		1	
43435132	**Good identification**					**43435132**
					GRA	
Parabacteroides distasonis		97.99	0.48	MNS 98	1	
Eubacterium saburreum		1.33	0.16		85	
Bifidobacterium longum subsp. longum		0.68	0.07		99	
43435322	**Very good identification**					**43435322**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis	100.0	0.61	MNS 98			
43435332 Very good identification						**43435332**
Parabacteroides distasonis	100.0	0.57	MNS 98			
43435520 Good identification						**43435520**
				GRA		
Bifidobacterium longum subsp. longum	95.89	0.37	MNS 87	99		
Parabacteroides distasonis	2.72	0.15	MLT 99	1		
Bacteroides vulgatus	1.39	-.08		1		
43435522 Low discrimination						**43435522**
				GRA		
Parabacteroides distasonis	73.75	0.49	MNS 98	1		
Bifidobacterium longum subsp. longum	26.25	0.37		99		
43435530 Good identification						**43435530**
				GRA		
Bifidobacterium longum subsp. longum	99.11	0.41	MNS 87	99		
Parabacteroides distasonis	0.89	0.11	MLT 99	1		
43435532 Low discrimination						**43435532**
				GRA		
Bifidobacterium longum subsp. longum	52.96	0.41	MNS 87	99		
Parabacteroides distasonis	47.04	0.45	MLT 99	1		
43435722 Very good identification						**43435722**
Parabacteroides distasonis	99.20	0.57	MNS 98			
Bacteroides thetaiotaomicron	0.81	0.06				
43435732 Good identification						**43435732**
Parabacteroides distasonis	98.55	0.53	MNS 98			
Bacteroides thetaiotaomicron	0.58	-.01				
Clostridium clostridioforme	0.50	0.08				
43440000 Low discrimination						**43440000**
				GRA	SPO	
Fusobacterium mortiferum	76.02	0.39	FRU 99	1	1	
Clostridium botulinum biovar A	13.36	0.13		99	99	
Eubacterium tortuosum	10.62	0.07		99	1	
43440100 Good identification						**43440100**
Fusobacterium mortiferum	100.0	0.45	FRU 99			
43441000 Acceptable identification						**43441000**
				GRA		
Eubacterium tortuosum	90.22	0.41	SAL 99	99		
Fusobacterium mortiferum	9.79	0.42	CEL 10	1		
43441100 Good identification						**43441100**
				COC	GRA	
Fusobacterium mortiferum	95.08	0.48	FRU 99	1	1	
Eubacterium tortuosum	3.80	0.07		1	99	
Blautia hansenii	1.13	-.03		99	99	
43450000 Very good identification						**43450000**
Fusobacterium mortiferum	100.0	0.73				
43450001 Good identification						**43450001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.39	URE 1		
43450002	**Very good identification**					**43450002**
Fusobacterium mortiferum		100.0	0.56	BGL 10		
43450010	**Good identification**					**43450010**
Fusobacterium mortiferum		100.0	0.39	MLZ 1		
43450020	**Good identification**					**43450020**
Fusobacterium mortiferum		100.0	0.39	NAG 1		
43450040	**Good identification**					**43450040**
Fusobacterium mortiferum		100.0	0.39	SOR 1		
43450100	**Very good identification**					**43450100**
Fusobacterium mortiferum		100.0	0.79			
43450101	**Good identification**					**43450101**
Fusobacterium mortiferum		100.0	0.45	URE 1		
43450102	**Very good identification**					**43450102**
Fusobacterium mortiferum		100.0	0.63	BGL 10		
43450110	**Good identification**					**43450110**
Fusobacterium mortiferum		100.0	0.45	MLZ 1		
43450120	**Good identification**					**43450120**
Fusobacterium mortiferum		98.52	0.45	NAG 1		
Prevotella melaninogenica		1.48	-.04			
43450140	**Good identification**					**43450140**
Fusobacterium mortiferum		100.0	0.45	SOR 1		
43450200	**Good identification**					**43450200**
Fusobacterium mortiferum		100.0	0.39	RHA 1		
43450300	**Good identification**					**43450300**
Fusobacterium mortiferum		100.0	0.45	RHA 1		
43450400	**Good identification**					**43450400**
Fusobacterium mortiferum		100.0	0.39	ARA 1		
43450500	**Good identification**					**43450500**
Fusobacterium mortiferum		100.0	0.45	ARA 1		
43451000	**Very good identification**					**43451000**
					GRA	
Fusobacterium mortiferum		99.84	0.76		1	
Eubacterium tortuosum		0.09	0.07		99	
Lactobacillus fermentum		0.05	0.20		99	
43451001	**Good identification**					**43451001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		98.31	0.42	URE 1	1		
Actinomyces naeslundii		1.69	0.23		99		
43451002	**Very good identification**						**43451002**
					GRA		
Fusobacterium mortiferum		99.45	0.59	BGL 10	1		
Lactobacillus fermentum		0.40	0.20		99		
Eubacterium tortuosum		0.15	-.06		99		
43451010	**Good identification**						**43451010**
Fusobacterium mortiferum		100.0	0.42	MLZ 1			
43451020	**Good identification**						**43451020**
					GRA		
Fusobacterium mortiferum		96.17	0.42	NAG 1	1		
Prevotella melaninogenica		2.25	-.04		1		
Eubacterium tortuosum		1.58	-.06		99		
43451040	**Good identification**						**43451040**
Fusobacterium mortiferum		100.0	0.42	SOR 1			
43451100	**Very good identification**						**43451100**
					GRA		
Fusobacterium mortiferum		99.91	0.82		1		
Lactobacillus fermentum		0.03	0.24		99		
Eubacterium saburreum		0.02	0.17		85		
43451101	**Good identification**						**43451101**
					GRA		
Fusobacterium mortiferum		97.95	0.48	URE 1	1		
Actinomyces naeslundii		2.05	0.31		99		
43451102	**Very good identification**						**43451102**
					GRA	CAT	
Fusobacterium mortiferum		97.99	0.66	BGL 10	1	5	
Capnocytophaga ochracea		1.33	0.20		1	90	
Lactobacillus fermentum		0.28	0.24		99	1	
43451110	**Good identification**						**43451110**
Fusobacterium mortiferum		98.45	0.48	MLZ 1			
Eubacterium saburreum		1.55	0.15				
43451120	**Low discrimination**						**43451120**
					CAT		
Fusobacterium mortiferum		45.66	0.48	NAG 1	5		
Prevotella melaninogenica		45.39	0.30		1		
Capnocytophaga ochracea		6.82	0.20		90		
43451122	**Low discrimination**						**43451122**
					CAT		
Capnocytophaga ochracea		62.81	0.36	MNS 90	90		
Prevotella oralis		20.02	0.27	MLT 99	1		
Prevotella melaninogenica		8.85	0.18		1		
43451140	**Good identification**						**43451140**
Fusobacterium mortiferum		100.0	0.48	SOR 1			
43451200	**Good identification**						**43451200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.42	RHA 1		
43451300	**Good identification**					**43451300**
Fusobacterium mortiferum		100.0	0.48	RHA 1		
43451400	**Good identification**					**43451400**
					GRA	
Fusobacterium mortiferum		95.76	0.42	ARA 1	1	
Lactobacillus fermentum		4.24	0.20		99	
43451500	**Good identification**					**43451500**
					GRA	
Fusobacterium mortiferum		97.00	0.48	ARA 1	1	
Lactobacillus fermentum		3.01	0.24		99	
43452000	**Good identification**					**43452000**
					GRA	
Fusobacterium mortiferum		97.21	0.39	MAN 1	1	
Eubacterium limosum		2.79	-.06		99	
43452100	**Good identification**					**43452100**
Fusobacterium mortiferum		100.0	0.45	MAN 1		
43453000	**Good identification**					**43453000**
					GRA	
Fusobacterium mortiferum		97.83	0.42	MAN 1	1	
Actinomyces naeslundii		2.17	0.25		99	
43453100	**Good identification**					**43453100**
					GRA	
Fusobacterium mortiferum		97.36	0.48	MAN 1	1	
Actinomyces naeslundii		2.64	0.33		99	
43453101	**Low discrimination**					**43453101**
					GRA	
Actinomyces naeslundii		80.08	0.36	ESL 12	99	
Fusobacterium mortiferum		19.92	0.14	MNS 88	1	
43453103	**Good identification**					**43453103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43453121	**Good identification**					**43453121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43453123	**Good identification**					**43453123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43454000	**Good identification**					**43454000**
Fusobacterium mortiferum		100.0	0.53	XYL 7		
43454002	**Good identification**					**43454002**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
43454100	**Very good identification**					**43454100**
Fusobacterium mortiferum		100.0	0.60	XYL 7		
43454102	**Good identification**					**43454102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.43	XYL 7			
43455000	**Very good identification**						**43455000**
					GRA		
Fusobacterium mortiferum		99.65	0.56	XYL 7	1		
Lactobacillus fermentum		0.35	0.16		99		
43455002	**Good identification**						**43455002**
					GRA		
Fusobacterium mortiferum		96.95	0.40	XYL 7	1		
Lactobacillus fermentum		3.05	0.16	BGL 10	99		
43455100	**Very good identification**						**43455100**
					GRA		
Fusobacterium mortiferum		99.50	0.63	XYL 7	1		
Lactobacillus fermentum		0.24	0.20		99		
Eubacterium saburreum		0.14	0.12		85		
43455102	**Low discrimination**						**43455102**
					GRA	CAT	
Fusobacterium mortiferum		85.38	0.46	XYL 7	1	5	
Bifidobacterium adolescentis		6.36	0.23	BGL 10	99	1	
Capnocytophaga ochracea		3.85	0.10		1	90	
43455322	**Good identification**						**43455322**
					CAT		
Parabacteroides distasonis		96.18	0.45	MNS 98	76		
Prevotella oralis		2.57	0.06		1		
Bacteroides fragilis		1.25	-.09		99		
43455332	**Good identification**						**43455332**
Parabacteroides distasonis		100.0	0.41	MNS 98			
43455502	**Acceptable identification**						**43455502**
Bifidobacterium adolescentis		91.62	0.37	MNS 87			
Lactobacillus fermentum		4.04	0.20	MLT 99			
Actinomyces israelii		2.49	0.17				
43455512	**Good identification**						**43455512**
Bifidobacterium adolescentis		98.25	0.39	MNS 87			
Bifidobacterium longum subsp. longum		1.75	0.07	MLT 99			
43455722	**Good identification**						**43455722**
Parabacteroides distasonis		95.17	0.42	MNS 98			
Bacteroides uniformis		4.83	0.01				
43455732	**Good identification**						**43455732**
Parabacteroides distasonis		100.0	0.37	MNS 98			
43457502	**Low discrimination**						**43457502**
Bifidobacterium adolescentis		84.18	0.36	MNS 87			
Actinomyces israelii		15.82	0.30	MLT 99			
43457512	**Good identification**						**43457512**
Bifidobacterium adolescentis		97.75	0.37	MNS 87			
Actinomyces israelii		2.25	0.16	MLT 99			
43460100	**Good identification**						**43460100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.36	FRU 99		
43461100	**Good identification**					**43461100**
Fusobacterium mortiferum		100.0	0.39	FRU 99		
43470000	**Very good identification**					**43470000**
Fusobacterium mortiferum		100.0	0.64			
43470002	**Good identification**					**43470002**
Fusobacterium mortiferum		100.0	0.47	BGL 10		
43470100	**Very good identification**					**43470100**
Fusobacterium mortiferum		100.0	0.70			
43470101	**Good identification**					**43470101**
Fusobacterium mortiferum		100.0	0.36	URE 1		
43470102	**Good identification**					**43470102**
Fusobacterium mortiferum		100.0	0.54	BGL 10		
43470110	**Good identification**					**43470110**
Fusobacterium mortiferum		100.0	0.36	MLZ 1		
43470120	**Good identification**					**43470120**
Fusobacterium mortiferum		100.0	0.36	NAG 1		
43470140	**Good identification**					**43470140**
Fusobacterium mortiferum		100.0	0.36	SOR 1		
43470300	**Good identification**					**43470300**
Fusobacterium mortiferum		100.0	0.36	RHA 1		
43470500	**Good identification**					**43470500**
Fusobacterium mortiferum		100.0	0.36	ARA 1		
43471000	**Very good identification**					**43471000**
					GRA	
Fusobacterium mortiferum		99.93	0.67		1	
Actinomyces naeslundii		0.08	0.25		99	
43471002	**Good identification**					**43471002**
					GRA	
Fusobacterium mortiferum		99.33	0.50	BGL 10	1	
Actinomyces naeslundii		0.67	0.25		99	
43471100	**Very good identification**					**43471100**
					GRA	
Fusobacterium mortiferum		99.87	0.73		1	
Actinomyces naeslundii		0.09	0.33		99	
Eubacterium saburreum		0.04	0.14		85	
43471101	**Low discrimination**					**43471101**
					GRA	
Fusobacterium mortiferum		88.03	0.39	URE 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		11.97	0.36		99	
43471102	**Very good identification**					**43471102**
					GRA	
Fusobacterium mortiferum		99.18	0.57	BGL 10	1	
Actinomyces naeslundii		0.82	0.33		99	
43471103	**Low discrimination**					**43471103**
					GRA	
Actinomyces naeslundii		55.03	0.36	ESL 12	99	
Fusobacterium mortiferum		44.97	0.23	MNS 88	1	
43471110	**Good identification**					**43471110**
Fusobacterium mortiferum		96.61	0.39	MLZ 1		
Eubacterium saburreum		3.40	0.12			
43471120	**Low discrimination**					**43471120**
					GRA	
Fusobacterium mortiferum		88.95	0.39	NAG 1	1	
Actinomyces naeslundii		8.06	0.33		99	
Prevotella melaninogenica		2.99	-.04		1	
43471121	**Good identification**					**43471121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43471123	**Good identification**					**43471123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43471140	**Good identification**					**43471140**
Fusobacterium mortiferum		100.0	0.39	SOR 1		
43471300	**Good identification**					**43471300**
Fusobacterium mortiferum		100.0	0.39	RHA 1		
43471322	**Good identification**					**43471322**
Parabacteroides distasonis		97.34	0.40	MNS 98		
Prevotella oralis		2.66	0.01	XYL 90		
43471332	**Good identification**					**43471332**
Parabacteroides distasonis		100.0	0.36	MNS 98		
43471500	**Good identification**					**43471500**
Fusobacterium mortiferum		100.0	0.39	ARA 1		
43471722	**Good identification**					**43471722**
Parabacteroides distasonis		100.0	0.37	MNS 98		
43472100	**Good identification**					**43472100**
Fusobacterium mortiferum		100.0	0.36	MAN 1		
43473100	**Low discrimination**					**43473100**
					GRA	
Fusobacterium mortiferum		81.13	0.39	MAN 1	1	
Actinomyces naeslundii		14.28	0.38		99	
Actinomyces israelii		4.59	0.20		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43473101	**Good identification**					**43473101**
Actinomyces naeslundii		100.0	0.41	ESL 12		
43473102	**Low discrimination**					**43473102**
					GRA	
Actinomyces naeslundii		47.31	0.38	ESL 12	99	
Fusobacterium mortiferum		29.87	0.23	MNS 88	1	
Actinomyces israelii		22.82	0.23	MLT 94	99	
43473103	**Good identification**					**43473103**
Actinomyces naeslundii		100.0	0.41	ESL 12		
43473120	**Good identification**					**43473120**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43473121	**Good identification**					**43473121**
Actinomyces naeslundii		100.0	0.41	ESL 12		
43473122	**Good identification**					**43473122**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43473123	**Good identification**					**43473123**
Actinomyces naeslundii		100.0	0.41	ESL 12		
43474000	**Good identification**					**43474000**
Fusobacterium mortiferum		100.0	0.44	XYL 7		
43474100	**Good identification**					**43474100**
Fusobacterium mortiferum		100.0	0.51	XYL 7		
43475000	**Good identification**					**43475000**
					GRA	
Fusobacterium mortiferum		99.11	0.48	XYL 7	1	
Actinomyces israelii		0.89	0.15		99	
43475100	**Good identification**					**43475100**
					GRA	
Fusobacterium mortiferum		98.18	0.54	XYL 7	1	
Actinomyces israelii		1.51	0.25		99	
Eubacterium saburreum		0.31	0.09		85	
43475102	**Low discrimination**					**43475102**
					GRA	
Fusobacterium mortiferum		70.09	0.37	XYL 7	1	
Actinomyces israelii		14.58	0.28	BGL 10	99	
Bifidobacterium adolescentis		11.64	0.20		99	
43475122	**Good identification**					**43475122**
Parabacteroides distasonis		99.15	0.48	MNS 98		
Bacteroides fragilis		0.85	-.09			
43475132	**Good identification**					**43475132**
Parabacteroides distasonis		100.0	0.44	MNS 98		
43475322	**Very good identification**					**43475322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.57	MNS 98			
43475332	**Good identification**						**43475332**
Parabacteroides distasonis		100.0	0.52	MNS 98			
43475512	**Good identification**						**43475512**
Bifidobacterium adolescentis		100.0	0.36	MNS 87			
43475522	**Good identification**						**43475522**
Parabacteroides distasonis		100.0	0.45	MNS 98			
43475532	**Good identification**						**43475532**
					GRA		
Parabacteroides distasonis		97.93	0.40	MNS 98	1		
Bifidobacterium longum subsp. longum		2.07	0.07		99		
43475722	**Good identification**						**43475722**
Parabacteroides distasonis		96.55	0.53	MNS 98			
Bacteroides thetaiotaomicron		3.09	0.11				
Clostridium clostridioforme		0.36	0.06				
43475732	**Acceptable identification**						**43475732**
Parabacteroides distasonis		94.82	0.48	MNS 98			
Clostridium clostridioforme		2.99	0.17				
Bacteroides thetaiotaomicron		2.19	0.04				
43477100	**Low discrimination**						**43477100**
					GRA		
Actinomyces israelii		89.63	0.38	MLT 99	99		
Fusobacterium mortiferum		10.37	0.20	SAL 92	1		
43477102	**Low discrimination**						**43477102**
Actinomyces israelii		89.66	0.41	MLT 99			
Bifidobacterium adolescentis		10.34	0.19	SAL 92			
43477500	**Good identification**						**43477500**
Actinomyces israelii		100.0	0.37	MLT 99			
43477502	**Low discrimination**						**43477502**
Actinomyces israelii		51.46	0.40	MLT 99			
Bifidobacterium adolescentis		48.54	0.33	SAL 92			
43500000	**Low discrimination**						**43500000**
					GRA	SPO	
Clostridium botulinum biovar A		77.70	0.45	SUC 7	99	99	
Clostridium sporogenes		17.41	0.32	RAF 1	99	99	
Fusobacterium mortiferum		2.02	0.31		1	1	
43500040	**Good identification**						**43500040**
					COC	GRA	
Clostridium botulinum biovar A		96.94	0.36	SUC 7	1	99	
Clostridium novyi biovar A		1.28	0.16	RAF 1	1	99	
Megasphaera elsdenii		1.05	0.07		99	1	
43501010	**Good identification**						**43501010**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium saburreum		98.69	0.45	IND 99	1			
Blautia hansenii		1.32	-.03		99			
43501100	**Very good identification**							**43501100**
					COC	GRA		
Blautia hansenii		96.58	0.66	SUC 1	99	99		
Eubacterium saburreum		2.11	0.53		1	85		
Prevotella melaninogenica		0.83	0.30		1	1		
43501101	**Low discrimination**							**43501101**
					COC	SPO		
Clostridium chauvoei		57.79	0.40	MNS 90	1	99		
Blautia hansenii		40.53	0.32	RAF 1	99	1		
Eubacterium saburreum		0.89	0.19		1	1		
43501102	**Acceptable identification**							**43501102**
					COC	CAT		
Blautia hansenii		90.21	0.50	SUC 1	99	1		
Capnocytophaga ochracea		4.17	0.30	BGL 10	1	90		
Eubacterium saburreum		3.13	0.40		1	1		
43501110	**Low discrimination**							**43501110**
					COC			
Eubacterium saburreum		62.91	0.52	IND 99	1			
Blautia hansenii		35.57	0.32		99			
Bifidobacterium longum subsp. longum		1.00	0.17		1			
43501112	**Low discrimination**							**43501112**
					COC	CAT		
Eubacterium saburreum		61.48	0.39	IND 99	1	1		
Blautia hansenii		21.89	0.15		99	1		
Capnocytophaga ochracea		11.12	0.14		1	90		
43501120	**Low discrimination**							**43501120**
					COC	CAT	GRA	
Prevotella melaninogenica		87.46	0.64	MNS 90	1	1	1	
Blautia hansenii		11.49	0.50	FRU 90	99	1	99	
Capnocytophaga ochracea		0.53	0.30		1	90	1	
43501122	**Low discrimination**							**43501122**
					COC	CAT	GRA	
Prevotella melaninogenica		72.34	0.52	MNS 90	1	1	1	
Capnocytophaga ochracea		20.74	0.47	FRU 90	1	90	1	
Blautia hansenii		5.55	0.33		99	1	99	
43501500	**Acceptable identification**							**43501500**
					COC			
Blautia hansenii		90.91	0.53	SUC 1	99			
Bifidobacterium longum subsp. longum		8.47	0.47		1			
Lactobacillus fermentum		0.41	0.32		1			
43501502	**Low discrimination**							**43501502**
					COC	GRA	CAT	
Blautia hansenii		51.61	0.37	SUC 1	99	99	1	
Bifidobacterium longum subsp. longum		43.25	0.47	BGL 10	1	99	1	
Capnocytophaga ochracea		2.38	0.17		1	1	90	
43501510	**Acceptable identification**							**43501510**
					COC			
Bifidobacterium longum subsp. longum		93.45	0.51	MNS 87	1			
Blautia hansenii		5.95	0.19	FRU 90	99			
Eubacterium saburreum		0.60	0.17		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43501512	**Good identification**							**43501512**
					COC			
Bifidobacterium longum subsp. longum		97.89	0.51	MNS 87	1			
Bifidobacterium adolescentis		1.07	0.19	FRU 90	1			
Blautia hansenii		0.69	0.03		99			
43501520	**Low discrimination**							**43501520**
					COC	GRA		
Bifidobacterium longum subsp. longum		69.15	0.57	MNS 87	1	99		
Blautia hansenii		20.63	0.37	FRU 90	99	99		
Prevotella melaninogenica		8.99	0.30		1	1		
43501522	**Low discrimination**							**43501522**
					COC	GRA	CAT	
Bifidobacterium longum subsp. longum		84.32	0.57	MNS 87	1	99	1	
Capnocytophaga ochracea		10.45	0.34	FRU 90	1	1	90	
Blautia hansenii		2.80	0.20		99	99	1	
43501530	**Very good identification**							**43501530**
					COC	GRA	CAT	
Bifidobacterium longum subsp. longum		99.66	0.61	MNS 87	1	99	1	
Blautia hansenii		0.18	0.03	FRU 90	99	99	1	
Capnocytophaga ochracea		0.09	0.01		1	1	90	
43501532	**Very good identification**							**43501532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		99.20	0.61	MNS 87	99	1		
Capnocytophaga ochracea		0.80	0.17	FRU 90	1	90		
43504530	**Good identification**							**43504530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87				
43504532	**Good identification**							**43504532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87				
43505000	**Low discrimination**							**43505000**
					COC			
Eubacterium saburreum		69.87	0.42	IND 99	1			
Blautia hansenii		24.72	0.19		99			
Lactobacillus fermentum		3.99	0.24		1			
43505010	**Good identification**							**43505010**
Eubacterium saburreum		100.0	0.41	IND 99				
43505100	**Acceptable identification**							**43505100**
					COC			
Blautia hansenii		92.96	0.53	SUC 1	99			
Eubacterium saburreum		6.19	0.49		1			
Bifidobacterium longum subsp. longum		0.26	0.21		1			
43505102	**Low discrimination**							**43505102**
					COC	CAT		
Blautia hansenii		80.36	0.37	SUC 1	99	1		
Eubacterium saburreum		8.50	0.36	BGL 10	1	1		
Capnocytophaga ochracea		5.26	0.20		1	90		
43505110	**Low discrimination**							**43505110**
					COC			
Eubacterium saburreum		78.52	0.47	IND 99	1			
Blautia hansenii		14.55	0.19		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		6.92	0.25		1			
43505120	**Low discrimination**							**43505120**
					COC	GRA		
Blautia hansenii		56.15	0.37	SUC 1	99	99		
Prevotella melaninogenica		24.46	0.30	NAG 10	1	1		
Eubacterium saburreum		5.94	0.36		1	85		
43505130	**Low discrimination**							**43505130**
					COC			
Bifidobacterium longum subsp. longum		61.49	0.35	MNS 87	1			
Eubacterium saburreum		30.77	0.34	FRU 90	1			
Blautia hansenii		3.59	0.03	ARA 99	99			
43505132	**Low discrimination**							**43505132**
					CAT	GRA		
Bifidobacterium longum subsp. longum		64.22	0.35	MNS 87	1	99		
Capnocytophaga ochracea		24.29	0.20	FRU 90	90	1		
Eubacterium saburreum		5.67	0.21	ARA 99	1	85		
43505430	**Good identification**							**43505430**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87				
43505432	**Good identification**							**43505432**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87				
43505500	**Low discrimination**							**43505500**
					COC			
Bifidobacterium longum subsp. longum		60.84	0.55	MNS 87	1			
Blautia hansenii		38.43	0.40	FRU 90	99			
Lactobacillus fermentum		0.58	0.28		1			
43505502	**Low discrimination**							**43505502**
					COC			
Bifidobacterium longum subsp. longum		89.44	0.55	MNS 87	1			
Blautia hansenii		6.28	0.24	FRU 90	99			
Bifidobacterium adolescentis		3.03	0.32		1			
43505510	**Very good identification**							**43505510**
					COC			
Bifidobacterium longum subsp. longum		99.51	0.59	MNS 87	1			
Blautia hansenii		0.37	0.06	FRU 90	99			
Eubacterium saburreum		0.12	0.13		1			
43505512	**Very good identification**							**43505512**
Bifidobacterium longum subsp. longum		97.63	0.59	MNS 87				
Bifidobacterium adolescentis		2.37	0.33	FRU 90				
43505520	**Very good identification**							**43505520**
					COC	GRA		
Bifidobacterium longum subsp. longum		97.29	0.66	MNS 87	1	99		
Blautia hansenii		1.71	0.24	FRU 90	99	99		
Bacteroides vulgatus		0.77	0.15		1	1		
43505522	**Very good identification**							**43505522**
					COC	GRA	CAT	
Bifidobacterium longum subsp. longum		98.55	0.66	MNS 87	1	99	1	
Capnocytophaga ochracea		1.02	0.23	FRU 90	1	1	90	
Blautia hansenii		0.19	0.07		99	99	1	
43505530	**Very good identification**							**43505530**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		100.0	0.70	MNS 87			
43505531	**Good identification**						**43505531**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43505532	**Very good identification**						**43505532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.93	0.70	MNS 87	99	1	
Capnocytophaga ochracea		0.07	0.07	FRU 90	1	90	
43505533	**Good identification**						**43505533**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43505570	**Good identification**						**43505570**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43505572	**Good identification**						**43505572**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43505730	**Good identification**						**43505730**
					GRA		
Bifidobacterium longum subsp. longum		95.99	0.35	MNS 87	99		
Bacteroides vulgatus		4.01	-.03	FRU 90	1		
43505732	**Good identification**						**43505732**
					GRA		
Bifidobacterium longum subsp. longum		95.50	0.35	MNS 87	99		
Parabacteroides distasonis		4.50	0.17	FRU 90	1		
43507530	**Good identification**						**43507530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43507532	**Good identification**						**43507532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87			
43510000	**Low discrimination**						**43510000**
					COC		
Fusobacterium mortiferum		84.58	0.65	MLT 11	1		
Prevotella intermedia		7.79	0.36		1		
Megasphaera elsdenii		3.02	0.32		99		
43510002	**Low discrimination**						**43510002**
					GRA	SPO	
Fusobacterium mortiferum		86.85	0.49	MLT 11	1	1	
Lactobacillus fermentum		4.25	0.28	BGL 10	99	1	
Clostridium sporogenes		4.00	0.11		99	99	
43510100	**Very good identification**						**43510100**
					GRA		
Fusobacterium mortiferum		98.00	0.72	MLT 11	1		
Actinomyces meyeri		0.55	0.24		99		
Lactobacillus fermentum		0.37	0.32		99		
43510101	**Low discrimination**						**43510101**
					GRA	SPO	
Fusobacterium mortiferum		83.23	0.37	MLT 11	1	1	
Clostridium chauvoei		13.30	0.12	URE 1	99	99	
Actinomyces naeslundii		3.47	0.25		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43510102	**Low discrimination**						**43510102**
					GRA	CAT	
Fusobacterium mortiferum		77.92	0.55	MLT 11	1	5	
Capnocytophaga ochracea		17.11	0.30	BGL 10	1	90	
Lactobacillus fermentum		2.66	0.32		99	1	
43510110	**Low discrimination**						**43510110**
					CAT		
Fusobacterium mortiferum		77.84	0.37	MLT 11	5		
Eubacterium saburreum		19.84	0.26	MLZ 1	1		
Capnocytophaga ochracea		2.32	-.03		90		
43510120	**Acceptable identification**						**43510120**
					CAT		
Prevotella melaninogenica		90.55	0.47	MNS 90	1		
Capnocytophaga ochracea		6.04	0.30	GAL 99	90		
Fusobacterium mortiferum		2.50	0.37		5		
43510122	**Low discrimination**						**43510122**
					CAT		
Capnocytophaga ochracea		73.53	0.47	MNS 90	90		
Prevotella melaninogenica		23.31	0.34	GAL 99	1		
Prevotella oralis		1.65	0.18		1		
43510140	**Acceptable identification**						**43510140**
					GRA	CAT	
Fusobacterium mortiferum		94.55	0.37	MLT 11	1	5	
Capnocytophaga ochracea		2.82	-.03	SOR 1	1	90	
Propionibacterium propionicum		2.63	0.12		99	1	
43510300	**Good identification**						**43510300**
Fusobacterium mortiferum		100.0	0.37	MLT 11			
43510500	**Low discrimination**						**43510500**
					GRA		
Fusobacterium mortiferum		44.15	0.37	MLT 11	1		
Bifidobacterium longum subsp. longum		31.05	0.29	ARA 1	99		
Lactobacillus fermentum		16.60	0.32		99		
43510520	**Low discrimination**						**43510520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		75.42	0.40	MNS 87	99	1	
Capnocytophaga ochracea		11.43	0.17	GAL 99	1	90	
Prevotella melaninogenica		9.80	0.12		1	1	
43510530	**Good identification**						**43510530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.02	0.44	MNS 87	99	1	
Capnocytophaga ochracea		0.98	0.01	GAL 99	1	90	
43510532	**Acceptable identification**						**43510532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.83	0.44	MNS 87	99	1	
Capnocytophaga ochracea		8.17	0.17	GAL 99	1	90	
43511000	**Low discrimination**						**43511000**
					GRA		
Fusobacterium mortiferum		60.72	0.68	MLT 11	1		
Lactobacillus fermentum		21.77	0.62		99		
Eubacterium saburreum		12.61	0.55		85		
43511002	**Low discrimination**						**43511002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Lactobacillus fermentum		64.81	0.62	ESL 1	99	1	
Fusobacterium mortiferum		20.09	0.52		1	5	
Capnocytophaga ochracea		6.86	0.30		1	90	
43511003	**Low discrimination**						**43511003**
					GRA		
Actinomyces naeslundii		51.57	0.41	ESL 12	99		
Lactobacillus fermentum		33.81	0.28	MNS 88	99		
Fusobacterium mortiferum		10.48	0.18		1		
43511010	**Acceptable identification**						**43511010**
					GRA		
Eubacterium saburreum		91.98	0.53	IND 99	85		
Fusobacterium mortiferum		5.47	0.34		1		
Lactobacillus fermentum		1.96	0.28		99		
43511012	**Low discrimination**						**43511012**
					CAT	GRA	
Eubacterium saburreum		76.99	0.41	IND 99	1	85	
Capnocytophaga ochracea		10.83	0.14		90	1	
Lactobacillus fermentum		9.30	0.28		1	99	
43511020	**Low discrimination**						**43511020**
					GRA	CAT	
Prevotella melaninogenica		71.87	0.47	MNS 90	1	1	
Lactobacillus fermentum		15.10	0.54	LAC 99	99	1	
Capnocytophaga ochracea		4.80	0.30		1	90	
43511022	**Low discrimination**						**43511022**
					GRA	CAT	
Capnocytophaga ochracea		53.62	0.47	MNS 90	1	90	
Lactobacillus fermentum		18.76	0.54	LAC 99	99	1	
Prevotella melaninogenica		17.00	0.34		1	1	
43511023	**Low discrimination**						**43511023**
					GRA	CAT	
Actinomyces naeslundii		46.94	0.41	ESL 12	99	25	
Capnocytophaga ochracea		29.33	0.12	MNS 88	1	90	
Lactobacillus fermentum		10.26	0.20		99	1	
43511030	**Low discrimination**						**43511030**
					CAT		
Eubacterium saburreum		71.54	0.41	IND 99	1		
Prevotella melaninogenica		13.71	0.12		1		
Capnocytophaga ochracea		10.06	0.14		90		
43511100	**Low discrimination**						**43511100**
					GRA		
Fusobacterium mortiferum		51.73	0.75	MLT 11	1		
Lactobacillus fermentum		12.97	0.66		99		
Prevotella melaninogenica		12.61	0.47		1		
43511101	**Low discrimination**						**43511101**
					GRA	SPO	
Clostridium chauvoei		80.35	0.47	MNS 90	99	99	
Fusobacterium mortiferum		7.62	0.40	RAF 1	1	1	
Actinomyces naeslundii		5.08	0.49		99	1	
43511102	**Low discrimination**						**43511102**
					GRA	CAT	
Capnocytophaga ochracea		76.04	0.64	MNS 90	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus fermentum		11.84	0.66	NAG 90	99	1		
Fusobacterium mortiferum		5.25	0.58		1	5		
43511110	**Low discrimination**							**43511110**
					CAT			
Eubacterium saburreum		79.58	0.60	IND 99	1			
Capnocytophaga ochracea		9.31	0.32		90			
Fusobacterium mortiferum		4.73	0.40		5			
43511112	**Low discrimination**							**43511112**
					CAT	GRA		
Capnocytophaga ochracea		75.88	0.48	MNS 90	90	1		
Eubacterium saburreum		12.71	0.47	MLZ 10	1	85		
Bifidobacterium adolescentis		6.06	0.39	NAG 90	1	99		
43511120	**Good identification**							**43511120**
					GRA	CAT		
Prevotella melaninogenica		92.96	0.81	MNS 90	1	1		
Capnocytophaga ochracea		6.21	0.64		1	90		
Lactobacillus fermentum		0.32	0.58		99	1		
43511121	**Low discrimination**							**43511121**
					GRA	CAT	SPO	
Prevotella melaninogenica		80.17	0.47	MNS 90	1	1	1	
Clostridium chauvoei		11.69	0.38	URE 1	99	1	99	
Capnocytophaga ochracea		5.35	0.30		1	90	1	
43511122	**Low discrimination**							**43511122**
					CAT			
Capnocytophaga ochracea		73.75	0.81	MNS 90	90			
Prevotella melaninogenica		23.38	0.68		1			
Prevotella oralis		1.66	0.52		1			
43511123	**Low discrimination**							**43511123**
					GRA	CAT		
Capnocytophaga ochracea		69.32	0.47	MNS 90	1	90		
Prevotella melaninogenica		21.98	0.34	URE 1	1	1		
Actinomyces naeslundii		3.19	0.49		99	25		
43511130	**Low discrimination**							**43511130**
					GRA	CAT		
Prevotella melaninogenica		50.12	0.47	MNS 90	1	1		
Capnocytophaga ochracea		36.79	0.48	MLZ 1	1	90		
Bifidobacterium longum subsp. longum		6.63	0.44		99	1		
43511132	**Good identification**							**43511132**
					GRA	CAT		
Capnocytophaga ochracea		94.57	0.64	MNS 90	1	90		
Prevotella melaninogenica		2.73	0.34	MLZ 10	1	1		
Bifidobacterium longum subsp. longum		1.90	0.44		99	1		
43511141	**Low discrimination**							**43511141**
					GRA	SPO		
Actinomyces naeslundii		48.21	0.36	ESL 12	99	1		
Clostridium chauvoei		43.65	0.12	MNS 88	99	99		
Capnocytophaga ochracea		8.14	-.03		1	1		
43511142	**Good identification**							**43511142**
					GRA	CAT		
Capnocytophaga ochracea		96.13	0.48	MNS 90	1	90		
Lactobacillus fermentum		1.36	0.32	NAG 90	99	1		
Bifidobacterium adolescentis		0.78	0.22	SOR 10	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43511160	**Low discrimination**						**43511160**
					GRA	CAT	
Prevotella melaninogenica		56.52	0.47	MNS 90	1	1	
Capnocytophaga ochracea		41.50	0.48	SOR 1	1	90	
Clostridium perfringens		1.48	0.26		99	1	
43511162	**Very good identification**						**43511162**
					CAT		
Capnocytophaga ochracea		96.67	0.64	MNS 90	90		
Prevotella melaninogenica		2.79	0.34	SOR 10	1		
Prevotella oralis		0.20	0.18		1		
43511172	**Good identification**						**43511172**
					GRA	CAT	
Capnocytophaga ochracea		99.56	0.48	MNS 90	1	90	
Prevotella melaninogenica		0.26	0.00	MLZ 10	1	1	
Bifidobacterium longum subsp. longum		0.18	0.09	SOR 10	99	1	
43511300	**Low discrimination**						**43511300**
					GRA		
Fusobacterium mortiferum		35.02	0.40	MLT 11	1		
Eubacterium contortum		31.67	0.27	RHA 1	99		
Lactobacillus fermentum		8.78	0.32		99		
43511320	**Low discrimination**						**43511320**
					CAT		
Prevotella melaninogenica		87.84	0.47	MNS 90	1		
Capnocytophaga ochracea		5.86	0.30	RHA 1	90		
Prevotella oralis		2.65	0.24		1		
43511322	**Low discrimination**						**43511322**
					CAT		
Capnocytophaga ochracea		41.63	0.47	MNS 90	90		
Prevotella oralis		39.70	0.46	RHA 1	1		
Prevotella melaninogenica		13.20	0.34		1		
43511400	**Good identification**						**43511400**
					GRA		
Lactobacillus fermentum		93.38	0.62	ESL 1	99		
Bifidobacterium longum subsp. longum		2.88	0.29		99		
Fusobacterium mortiferum		2.63	0.34		1		
43511402	**Good identification**						**43511402**
					GRA	CAT	
Lactobacillus fermentum		94.54	0.62	ESL 1	99	1	
Bifidobacterium longum subsp. longum		2.92	0.29		99	1	
Capnocytophaga ochracea		1.77	0.17		1	90	
43511420	**Low discrimination**						**43511420**
					GRA	CAT	
Lactobacillus fermentum		67.84	0.54	ESL 1	99	1	
Bifidobacterium longum subsp. longum		25.10	0.40		99	1	
Capnocytophaga ochracea		3.80	0.17		1	90	
43511422	**Low discrimination**						**43511422**
					GRA	CAT	
Lactobacillus fermentum		52.97	0.54	ESL 1	99	1	
Capnocytophaga ochracea		26.73	0.34		1	90	
Bifidobacterium longum subsp. longum		19.60	0.40		99	1	
43511430	**Good identification**						**43511430**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		97.47	0.44	MNS 87	99	1	
Lactobacillus fermentum		1.56	0.20	LAC 99	99	1	
Capnocytophaga ochracea		0.96	0.01		1	90	
43511432	**Acceptable identification**						**43511432**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.50	0.44	MNS 87	99	1	
Capnocytophaga ochracea		8.05	0.17	LAC 99	1	90	
Lactobacillus fermentum		1.45	0.20		99	1	
43511500	**Low discrimination**						**43511500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		60.61	0.63	MNS 87	99	1	
Lactobacillus fermentum		32.40	0.66		99	1	
Capnocytophaga ochracea		4.08	0.35		1	90	
43511502	**Low discrimination**						**43511502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		43.22	0.63	MNS 87	99	1	
Capnocytophaga ochracea		26.19	0.51		1	90	
Lactobacillus fermentum		23.10	0.66		99	1	
43511510	**Very good identification**						**43511510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.92	0.67	MNS 87	99	1	
Capnocytophaga ochracea		0.44	0.19		1	90	
Lactobacillus fermentum		0.31	0.32		99	1	
43511512	**Low discrimination**						**43511512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		86.00	0.67	MNS 87	99	1	
Bifidobacterium adolescentis		10.30	0.53		99	1	
Capnocytophaga ochracea		3.40	0.35		1	90	
43511520	**Low discrimination**						**43511520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		75.07	0.74	MNS 87	99	1	
Capnocytophaga ochracea		11.37	0.51		1	90	
Prevotella melaninogenica		9.75	0.47		1	1	
43511521	**Low discrimination**						**43511521**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.35	0.40	MNS 87	99	1	
Capnocytophaga ochracea		11.27	0.17	URE 1	1	90	
Prevotella melaninogenica		9.66	0.12		1	1	
43511522	**Low discrimination**						**43511522**
					GRA	CAT	
Capnocytophaga ochracea		55.94	0.68	MNS 90	1	90	
Bifidobacterium longum subsp. longum		41.03	0.74		99	1	
Lactobacillus fermentum		1.83	0.58		99	1	
43511530	**Very good identification**						**43511530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.91	0.78	MNS 87	99	1	
Capnocytophaga ochracea		0.98	0.35		1	90	
Prevotella melaninogenica		0.08	0.12		1	1	
43511531	**Good identification**						**43511531**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.02	0.44	MNS 87	99	1	
Capnocytophaga ochracea		0.98	0.01	URE 1	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43511532	**Good identification**						**43511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.72	0.78	MNS 87	99	1	
Capnocytophaga ochracea		8.16	0.51		1	90	
Parabacteroides distasonis		0.05	0.27		1	76	
43511533	**Acceptable identification**						**43511533**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.83	0.44	MNS 87	99	1	
Capnocytophaga ochracea		8.17	0.17	URE 1	1	90	
43511542	**Low discrimination**						**43511542**
					GRA	CAT	
Capnocytophaga ochracea		65.09	0.35	MNS 90	1	90	
Bifidobacterium adolescentis		19.93	0.36	NAG 90	99	1	
Bifidobacterium longum subsp. longum		9.76	0.29	SOR 10	99	1	
43511552	**Low discrimination**						**43511552**
					GRA	CAT	
Bifidobacterium adolescentis		50.52	0.38	MNS 87	99	1	
Bifidobacterium longum subsp. longum		34.48	0.33	XYL 87	99	1	
Capnocytophaga ochracea		15.00	0.19	SOR 11	1	90	
43511560	**Low discrimination**						**43511560**
					GRA	CAT	
Capnocytophaga ochracea		58.66	0.35	MNS 90	1	90	
Bifidobacterium longum subsp. longum		35.20	0.40	SOR 10	99	1	
Prevotella melaninogenica		4.57	0.12	BGL 90	1	1	
43511562	**Acceptable identification**						**43511562**
					GRA	CAT	
Capnocytophaga ochracea		93.35	0.51	MNS 90	1	90	
Bifidobacterium longum subsp. longum		6.22	0.40	SOR 10	99	1	
Lactobacillus fermentum		0.28	0.24		99	1	
43511570	**Acceptable identification**						**43511570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.19	0.44	MNS 87	99	1	
Capnocytophaga ochracea		9.81	0.19	SOR 1	1	90	
43511572	**Low discrimination**						**43511572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		50.54	0.44	MNS 87	99	1	
Capnocytophaga ochracea		49.47	0.35	SOR 1	1	90	
43511720	**Low discrimination**						**43511720**
					GRA	CAT	
Bacteroides vulgatus		77.25	0.37	MNS 99	1	3	
Bifidobacterium longum subsp. longum		16.30	0.40	XYL 98	99	1	
Capnocytophaga ochracea		2.47	0.17		1	90	
43511730	**Good identification**						**43511730**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.34	0.44	MNS 87	99	1	
Bacteroides vulgatus		2.71	0.03	RHA 1	1	3	
Capnocytophaga ochracea		0.95	0.01		1	90	
43511732	**Low discrimination**						**43511732**
					GRA	CAT	
Bifidobacterium longum subsp. longum		78.41	0.44	MNS 87	99	1	
Parabacteroides distasonis		13.55	0.35	RHA 1	1	76	
Capnocytophaga ochracea		6.98	0.17		1	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
43512100	**Low discrimination**						**43512100**
					GRA		
Propionibacterium propionicum		72.35	0.46	NIT 86	99		
Fusobacterium mortiferum		26.23	0.37	ESL 1	1		
Actinomyces naeslundii		1.42	0.27		99		
43512102	**Good identification**						**43512102**
					GRA		
Propionibacterium propionicum		96.16	0.49	NIT 86	99		
Fusobacterium mortiferum		2.58	0.21	ESL 1	1		
Actinomyces naeslundii		1.25	0.27		99		
43512140	**Good identification**						**43512140**
Propionibacterium propionicum		100.0	0.46	NIT 86			
43512142	**Good identification**						**43512142**
Propionibacterium propionicum		100.0	0.49	NIT 86			
43513001	**Good identification**						**43513001**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43513002	**Low discrimination**						**43513002**
					GRA		
Actinomyces naeslundii		53.56	0.43	ESL 12	99		
Lactobacillus fermentum		27.13	0.28	MNS 88	99		
Fusobacterium mortiferum		8.41	0.18		1		
43513003	**Good identification**						**43513003**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43513020	**Low discrimination**						**43513020**
					GRA		
Actinomyces naeslundii		50.70	0.43	ESL 12	99		
Prevotella melaninogenica		40.74	0.12	MNS 88	1		
Lactobacillus fermentum		8.56	0.20		99		
43513021	**Good identification**						**43513021**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43513022	**Low discrimination**						**43513022**
					GRA	CAT	
Actinomyces naeslundii		53.38	0.43	ESL 12	99	25	
Capnocytophaga ochracea		25.77	0.12	MNS 88	1	90	
Lactobacillus fermentum		9.02	0.20		99	1	
43513023	**Good identification**						**43513023**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43513100	**Low discrimination**						**43513100**
					GRA		
Propionibacterium propionicum		39.08	0.46	NIT 86	99		
Fusobacterium mortiferum		21.26	0.40	ESL 1	1		
Actinomyces naeslundii		18.34	0.51		99		
43513101	**Low discrimination**						**43513101**
					SPO		
Actinomyces naeslundii		80.64	0.54	ESL 12	1		
Propionibacterium propionicum		12.73	0.30	MNS 88	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		6.64	0.12		99		
43513102	**Low discrimination**						**43513102**
					GRA	CAT	
Propionibacterium propionicum		40.70	0.49	NIT 86	99	1	
Capnocytophaga ochracea		23.76	0.30	ESL 1	1	90	
Bifidobacterium adolescentis		13.97	0.36		99	1	
43513103	**Low discrimination**						**43513103**
					SPO		
Actinomyces naeslundii		79.92	0.54	ESL 12	1		
Propionibacterium propionicum		18.92	0.33	MNS 88	1		
Clostridium chauvoei		1.16	-.01		99		
43513112	**Low discrimination**						**43513112**
					GRA	CAT	
Bifidobacterium adolescentis		79.28	0.38	MNS 87	99	1	
Capnocytophaga ochracea		12.26	0.14	XYL 87	1	90	
Actinomyces naeslundii		3.11	0.29	ARA 87	99	25	
43513120	**Acceptable identification**						**43513120**
					GRA	CAT	
Prevotella melaninogenica		90.06	0.47	MNS 90	1	1	
Capnocytophaga ochracea		6.01	0.30	MAN 1	1	90	
Actinomyces naeslundii		3.22	0.51		99	25	
43513121	**Low discrimination**						**43513121**
					GRA	SPO	
Actinomyces naeslundii		82.26	0.54	ESL 12	99	1	
Prevotella melaninogenica		15.49	0.12	MNS 88	1	1	
Clostridium chauvoei		2.26	0.04		99	99	
43513122	**Low discrimination**						**43513122**
					GRA	CAT	
Capnocytophaga ochracea		70.66	0.47	MNS 90	1	90	
Prevotella melaninogenica		22.41	0.34	MAN 1	1	1	
Actinomyces naeslundii		4.21	0.51		99	25	
43513123	**Low discrimination**						**43513123**
					GRA	CAT	
Actinomyces naeslundii		87.04	0.54	ESL 12	99	25	
Capnocytophaga ochracea		9.84	0.12	MNS 88	1	90	
Prevotella melaninogenica		3.12	0.00		1	1	
43513140	**Low discrimination**						**43513140**
Propionibacterium propionicum		89.87	0.46	NIT 86			
Actinomyces naeslundii		7.44	0.38	ESL 1			
Actinomyces israelii		1.72	0.18				
43513141	**Low discrimination**						**43513141**
Actinomyces naeslundii		52.79	0.41	ESL 12			
Propionibacterium propionicum		47.21	0.30	MNS 88			
43513142	**Low discrimination**						**43513142**
					GRA	CAT	
Propionibacterium propionicum		84.63	0.49	NIT 86	99	1	
Capnocytophaga ochracea		5.49	0.14	ESL 1	1	90	
Actinomyces naeslundii		4.67	0.38		99	25	
43513161	**Good identification**						**43513161**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	ESL 12			
43513163	**Good identification**						**43513163**
Actinomyces naeslundii		100.0	0.41	ESL 12			
43513502	**Low discrimination**						**43513502**
					GRA	CAT	
Bifidobacterium adolescentis		84.72	0.50	MNS 87	99	1	
Bifidobacterium longum subsp. longum		6.27	0.29	XYL 87	99	1	
Capnocytophaga ochracea		3.80	0.17		1	90	
43513512	**Acceptable identification**						**43513512**
					GRA	CAT	
Bifidobacterium adolescentis		90.32	0.52	MNS 87	99	1	
Bifidobacterium longum subsp. longum		9.31	0.33	XYL 87	99	1	
Capnocytophaga ochracea		0.37	0.01		1	90	
43513520	**Low discrimination**						**43513520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		75.42	0.40	MNS 87	99	1	
Capnocytophaga ochracea		11.43	0.17	MAN 1	1	90	
Prevotella melaninogenica		9.80	0.12		1	1	
43513530	**Good identification**						**43513530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.02	0.44	MNS 87	99	1	
Capnocytophaga ochracea		0.98	0.01	MAN 1	1	90	
43513532	**Low discrimination**						**43513532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.81	0.44	MNS 87	99	1	
Capnocytophaga ochracea		7.99	0.17	MAN 1	1	90	
Bifidobacterium adolescentis		2.20	0.18		99	1	
43513552	**Good identification**						**43513552**
Bifidobacterium adolescentis		100.0	0.36	MNS 87			
43514000	**Low discrimination**						**43514000**
					GRA	SPO	
Fusobacterium mortiferum		82.46	0.46	MLT 11	1	1	
Actinomyces meyeri		6.93	0.18	XYL 7	99	1	
Clostridium glycolicum		4.24	0.10		99	99	
43514100	**Low discrimination**						**43514100**
					GRA		
Fusobacterium mortiferum		82.98	0.52	MLT 11	1		
Actinomyces meyeri		11.95	0.29	XYL 7	99		
Lactobacillus fermentum		2.46	0.28		99		
43514102	**Low discrimination**						**43514102**
					GRA	CAT	
Fusobacterium mortiferum		42.75	0.36	MLT 11	1	5	
Capnocytophaga ochracea		31.18	0.20	XYL 7	1	90	
Lactobacillus fermentum		11.40	0.28	BGL 10	99	1	
43514500	**Low discrimination**						**43514500**
Bifidobacterium longum subsp. longum		78.90	0.37	MNS 87			
Actinomyces meyeri		10.03	0.18	GAL 99			
Lactobacillus fermentum		8.26	0.28				
43514502	**Low discrimination**						**43514502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		65.48	0.37	MNS 87			
Bifidobacterium adolescentis		24.36	0.32	GAL 99			
Lactobacillus fermentum		6.85	0.28				
43514510	**Good identification**						**43514510**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87			
43514512	**Low discrimination**						**43514512**
Bifidobacterium longum subsp. longum		78.41	0.41	MNS 87			
Bifidobacterium adolescentis		20.94	0.33	GAL 99			
Clostridium clostridioforme		0.65	0.04				
43514520	**Low discrimination**						**43514520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.88	0.48	MNS 87	99	1	
Bacteroides vulgatus		7.81	0.15	GAL 99	1	3	
Capnocytophaga ochracea		1.14	0.07		1	90	
43514522	**Low discrimination**						**43514522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.71	0.48	MNS 87	99	1	
Capnocytophaga ochracea		9.97	0.23	GAL 99	1	90	
Bacteroides fragilis		1.08	-.03		1	99	
43514530	**Good identification**						**43514530**
Bifidobacterium longum subsp. longum		100.0	0.52	MNS 87			
43514532	**Good identification**						**43514532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.06	0.52	MNS 87	99	1	
Capnocytophaga ochracea		0.74	0.07	GAL 99	1	90	
Clostridium clostridioforme		0.21	0.04		50	33	
43515000	**Low discrimination**						**43515000**
					GRA		
Lactobacillus fermentum		51.77	0.58	ESL 1	99		
Eubacterium saburreum		27.49	0.50		85		
Fusobacterium mortiferum		18.51	0.49		1		
43515002	**Low discrimination**						**43515002**
					CAT	GRA	
Lactobacillus fermentum		83.15	0.58	ESL 1	1	99	
Eubacterium saburreum		7.79	0.37		1	85	
Capnocytophaga ochracea		3.75	0.20		90	1	
43515010	**Good identification**						**43515010**
					GRA		
Eubacterium saburreum		96.36	0.49	IND 99	85		
Lactobacillus fermentum		2.24	0.24		99		
Fusobacterium mortiferum		0.80	0.15		1		
43515012	**Low discrimination**						**43515012**
Eubacterium saburreum		72.89	0.36	IND 99			
Bifidobacterium adolescentis		10.17	0.19				
Lactobacillus fermentum		9.60	0.24				
43515020	**Low discrimination**						**43515020**
					CAT	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		59.74	0.50	ESL 1	1	99	
Eubacterium saburreum		16.80	0.37		1	85	
Capnocytophaga ochracea		8.08	0.20		90	1	
43515030	**Low discrimination**						**43515030**
					CAT	GRA	
Eubacterium saburreum		79.97	0.36	IND 99	1	85	
Bifidobacterium longum subsp. longum		11.30	0.18		1	99	
Capnocytophaga ochracea		5.22	0.04		90	1	
43515100	**Low discrimination**						**43515100**
					GRA		
Lactobacillus fermentum		34.28	0.62	ESL 1	99		
Eubacterium saburreum		26.04	0.57		85		
Fusobacterium mortiferum		17.53	0.55		1		
43515102	**Low discrimination**						**43515102**
					GRA	CAT	
Capnocytophaga ochracea		52.76	0.54	MNS 90	1	90	
Lactobacillus fermentum		19.29	0.62	NAG 90	99	1	
Bifidobacterium adolescentis		10.25	0.52		99	1	
43515110	**Low discrimination**						**43515110**
					CAT	GRA	
Eubacterium saburreum		73.70	0.55	IND 99	1	85	
Bifidobacterium longum subsp. longum		19.49	0.41		1	99	
Capnocytophaga ochracea		4.00	0.21		90	1	
43515112	**Low discrimination**						**43515112**
					GRA	CAT	
Bifidobacterium adolescentis		52.08	0.53	MNS 87	99	1	
Capnocytophaga ochracea		24.37	0.38	ARA 87	1	90	
Bifidobacterium longum subsp. longum		13.18	0.41		99	1	
43515120	**Low discrimination**						**43515120**
					CAT		
Capnocytophaga ochracea		34.14	0.54	MNS 90	90		
Bacteroides fragilis		30.57	0.44	BGL 90	99		
Prevotella melaninogenica		20.66	0.47		1		
43515122	**Low discrimination**						**43515122**
					GRA	CAT	
Bacteroides fragilis		63.73	0.66	MNS 99	1	99	
Capnocytophaga ochracea		33.71	0.70		1	90	
Bifidobacterium longum subsp. longum		0.53	0.48		99	1	
43515130	**Low discrimination**						**43515130**
					CAT	GRA	
Bifidobacterium longum subsp. longum		58.69	0.52	MNS 87	1	99	
Capnocytophaga ochracea		27.13	0.38	ARA 99	90	1	
Eubacterium saburreum		9.79	0.42		1	85	
43515132	**Low discrimination**						**43515132**
					GRA	CAT	
Capnocytophaga ochracea		66.95	0.54	MNS 90	1	90	
Bifidobacterium longum subsp. longum		16.09	0.52	MLZ 10	99	1	
Bacteroides fragilis		11.51	0.32		1	99	
43515142	**Low discrimination**						**43515142**
					GRA	CAT	
Capnocytophaga ochracea		77.31	0.38	MNS 90	1	90	
Bifidobacterium adolescentis		16.71	0.36	NAG 90	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		2.57	0.28	SOR 10	99	1	
43515152	**Low discrimination**						**43515152**
					GRA	CAT	
Bifidobacterium adolescentis		69.38	0.38	MNS 87	99	1	
Capnocytophaga ochracea		29.19	0.21	ARA 87	1	90	
Bifidobacterium longum subsp. longum		1.44	0.07	SOR 11	99	1	
43515160	**Low discrimination**						**43515160**
					CAT		
Capnocytophaga ochracea		85.24	0.38	MNS 90	90		
Bacteroides fragilis		6.94	0.10	SOR 10	99		
Prevotella melaninogenica		4.69	0.12	BGL 90	1		
43515162	**Low discrimination**						**43515162**
					GRA	CAT	
Capnocytophaga ochracea		84.84	0.54	MNS 90	1	90	
Bacteroides fragilis		14.58	0.32	SOR 10	1	99	
Bifidobacterium longum subsp. longum		0.12	0.14		99	1	
43515172	**Good identification**						**43515172**
					GRA	CAT	
Capnocytophaga ochracea		95.82	0.38	MNS 90	1	90	
Bifidobacterium longum subsp. longum		2.09	0.18	MLZ 10	99	1	
Bacteroides fragilis		1.50	-.03	SOR 10	1	99	
43515300	**Very good identification**						**43515300**
					GRA		
Eubacterium contortum		99.21	0.61	SAL 99	99		
Bacteroides vulgatus		0.37	0.13		1		
Lactobacillus fermentum		0.16	0.28		99		
43515302	**Acceptable identification**						**43515302**
					GRA	CAT	
Eubacterium contortum		94.16	0.48	SAL 99	99	1	
Capnocytophaga ochracea		2.40	0.20		1	90	
Parabacteroides distasonis		0.91	0.25		1	76	
43515320	**Low discrimination**						**43515320**
					GRA	CAT	
Bacteroides vulgatus		65.63	0.47	MNS 99	1	3	
Eubacterium contortum		31.62	0.48	ARA 93	99	1	
Capnocytophaga ochracea		0.81	0.20		1	90	
43515322	**Low discrimination**						**43515322**
Parabacteroides distasonis		49.30	0.59	MNS 98			
Bacteroides fragilis		22.44	0.32				
Capnocytophaga ochracea		11.87	0.36				
43515332	**Acceptable identification**						**43515332**
					GRA	CAT	
Parabacteroides distasonis		92.85	0.55	MNS 98	1	76	
Capnocytophaga ochracea		4.61	0.20		1	90	
Bifidobacterium longum subsp. longum		1.11	0.18		99	1	
43515400	**Low discrimination**						**43515400**
Lactobacillus fermentum		84.88	0.58	ESL 1			
Bifidobacterium longum subsp. longum		13.37	0.37				
Actinomyces meyeri		0.64	0.11				
43515402	**Low discrimination**						**43515402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		81.28	0.58	ESL 1			
Bifidobacterium longum subsp. longum		12.80	0.37				
Bifidobacterium adolescentis		4.76	0.32				
43515410	**Acceptable identification**						**43515410**
Bifidobacterium longum subsp. longum		94.87	0.41	MNS 87			
Lactobacillus fermentum		3.57	0.24	LAC 99			
Eubacterium saburreum		1.55	0.15				
43515412	**Low discrimination**						**43515412**
Bifidobacterium longum subsp. longum		76.64	0.41	MNS 87			
Bifidobacterium adolescentis		20.47	0.33	LAC 99			
Lactobacillus fermentum		2.89	0.24				
43515420	**Low discrimination**						**43515420**
					GRA		
Bifidobacterium longum subsp. longum		61.40	0.48	MNS 87	99		
Lactobacillus fermentum		32.49	0.50	LAC 99	99		
Bacteroides vulgatus		5.34	0.15		1		
43515422	**Low discrimination**						**43515422**
					GRA	CAT	
Bifidobacterium longum subsp. longum		60.22	0.48	MNS 87	99	1	
Lactobacillus fermentum		31.86	0.50	LAC 99	99	1	
Capnocytophaga ochracea		6.84	0.23		1	90	
43515430	**Good identification**						**43515430**
Bifidobacterium longum subsp. longum		99.69	0.52	MNS 87			
Lactobacillus fermentum		0.31	0.16	LAC 99			
43515432	**Good identification**						**43515432**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.96	0.52	MNS 87	99	1	
Capnocytophaga ochracea		0.73	0.07	LAC 99	1	90	
Lactobacillus fermentum		0.31	0.16		99	1	
43515500	**Low discrimination**						**43515500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.66	0.72	MNS 87	99	1	
Lactobacillus fermentum		9.28	0.62		99	1	
Capnocytophaga ochracea		0.50	0.25		1	90	
43515501	**Acceptable identification**						**43515501**
Bifidobacterium longum subsp. longum		90.53	0.37	MNS 87			
Lactobacillus fermentum		9.47	0.28	URE 1			
43515502	**Low discrimination**						**43515502**
Bifidobacterium longum subsp. longum		65.33	0.72	MNS 87			
Bifidobacterium adolescentis		24.31	0.66				
Lactobacillus fermentum		6.84	0.62				
43515503	**Low discrimination**						**43515503**
Bifidobacterium longum subsp. longum		65.48	0.37	MNS 87			
Bifidobacterium adolescentis		24.36	0.32	URE 1			
Lactobacillus fermentum		6.85	0.28				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43515510	**Very good identification**						**43515510**
Bifidobacterium longum subsp. longum		99.59	0.76	MNS 87			
Bifidobacterium adolescentis		0.27	0.33				
Lactobacillus fermentum		0.06	0.28				
43515511	**Good identification**						**43515511**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87			
43515512	**Low discrimination**						**43515512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		78.65	0.76	MNS 87	99	1	
Bifidobacterium adolescentis		21.01	0.67		99	1	
Capnocytophaga ochracea		0.26	0.25		1	90	
43515513	**Low discrimination**						**43515513**
Bifidobacterium longum subsp. longum		78.92	0.41	MNS 87			
Bifidobacterium adolescentis		21.08	0.33	URE 1			
43515520	**Good identification**						**43515520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.11	0.82	MNS 87	99	1	
Bacteroides vulgatus		7.83	0.50		1	3	
Capnocytophaga ochracea		1.14	0.41		1	90	
43515521	**Acceptable identification**						**43515521**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.23	0.48	MNS 87	99	1	
Bacteroides vulgatus		7.84	0.15	URE 1	1	3	
Capnocytophaga ochracea		1.14	0.07		1	90	
43515522	**Low discrimination**						**43515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.55	0.82	MNS 87	99	1	
Capnocytophaga ochracea		9.95	0.58		1	90	
Bacteroides fragilis		1.08	0.32		1	99	
43515523	**Low discrimination**						**43515523**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.71	0.48	MNS 87	99	1	
Capnocytophaga ochracea		9.97	0.23	URE 1	1	90	
Bacteroides fragilis		1.08	-.03		1	99	
43515530	**Very good identification**						**43515530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.86	0.86	MNS 87	99	1	
Capnocytophaga ochracea		0.08	0.25		1	90	
Bacteroides vulgatus		0.05	0.15		1	3	
43515531	**Good identification**						**43515531**
Bifidobacterium longum subsp. longum		100.0	0.52	MNS 87			
43515532	**Very good identification**						**43515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.01	0.86	MNS 87	99	1	
Capnocytophaga ochracea		0.73	0.41		1	90	
Parabacteroides distasonis		0.17	0.43		1	76	
43515533	**Good identification**						**43515533**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		99.26	0.52	MNS 87	99	1	
Capnocytophaga ochracea		0.74	0.07	URE 1	1	90	
43515540	**Low discrimination**						**43515540**
					GRA	CAT	
Bifidobacterium longum subsp. longum		79.76	0.37	MNS 87	99	1	
Lactobacillus fermentum		8.35	0.28	SOR 1	99	1	
Capnocytophaga ochracea		4.92	0.08		1	90	
43515542	**Low discrimination**						**43515542**
					GRA	CAT	
Bifidobacterium adolescentis		72.55	0.50	MNS 87	99	1	
Bifidobacterium longum subsp. longum		15.94	0.37	SOR 11	99	1	
Capnocytophaga ochracea		8.85	0.25		1	90	
43515550	**Good identification**						**43515550**
Bifidobacterium longum subsp. longum		96.80	0.41	MNS 87			
Bifidobacterium adolescentis		3.20	0.18	SOR 1			
43515552	**Low discrimination**						**43515552**
					GRA	CAT	
Bifidobacterium adolescentis		75.93	0.52	MNS 87	99	1	
Bifidobacterium longum subsp. longum		23.23	0.41	SOR 11	99	1	
Capnocytophaga ochracea		0.84	0.08		1	90	
43515560	**Low discrimination**						**43515560**
					GRA	CAT	
Bifidobacterium longum subsp. longum		81.01	0.48	MNS 87	99	1	
Capnocytophaga ochracea		11.25	0.25	SOR 1	1	90	
Bacteroides vulgatus		7.04	0.15		1	3	
43515562	**Low discrimination**						**43515562**
					GRA	CAT	
Capnocytophaga ochracea		54.63	0.41	MNS 90	1	90	
Bifidobacterium longum subsp. longum		43.71	0.48	SOR 10	99	1	
Bacteroides fragilis		0.54	-.03		1	99	
43515570	**Good identification**						**43515570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.10	0.52	MNS 87	99	1	
Capnocytophaga ochracea		0.90	0.08	SOR 1	1	90	
43515572	**Acceptable identification**						**43515572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.76	0.52	MNS 87	99	1	
Capnocytophaga ochracea		7.48	0.25	SOR 1	1	90	
Bifidobacterium adolescentis		0.76	0.18		99	1	
43515700	**Good identification**						**43515700**
					GRA		
Eubacterium contortum		92.48	0.60	SAL 99	99		
Bacteroides vulgatus		5.56	0.32		1		
Bifidobacterium longum subsp. longum		1.78	0.37		99		
43515702	**Low discrimination**						**43515702**
Eubacterium contortum		84.44	0.47	SAL 99			
Bifidobacterium longum subsp. longum		9.19	0.37				
Bifidobacterium adolescentis		3.42	0.32				
43515710	**Low discrimination**						**43515710**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		75.34	0.41	MNS 87	99		
Eubacterium contortum		23.27	0.26	RHA 1	99		
Bacteroides vulgatus		1.40	-.03		1		
43515712	**Low discrimination**						**43515712**
Bifidobacterium longum subsp. longum		65.57	0.41	MNS 87			
Bifidobacterium adolescentis		17.51	0.33	RHA 1			
Clostridium clostridioforme		11.97	0.27				
43515720	**Very good identification**						**43515720**
					GRA		
Bacteroides vulgatus		95.87	0.66	MNS 99	1		
Eubacterium contortum		2.84	0.47		99		
Bifidobacterium longum subsp. longum		1.24	0.48		99		
43515722	**Low discrimination**						**43515722**
					GRA		
Parabacteroides distasonis		41.00	0.56	MNS 98	1		
Bifidobacterium longum subsp. longum		25.01	0.48		99		
Bacteroides vulgatus		19.56	0.32		1		
43515730	**Low discrimination**						**43515730**
					GRA		
Bifidobacterium longum subsp. longum		67.65	0.52	MNS 87	99		
Bacteroides vulgatus		31.07	0.32	RHA 1	1		
Eubacterium contortum		0.92	0.13		99		
43515732	**Low discrimination**						**43515732**
					GRA		
Bifidobacterium longum subsp. longum		63.16	0.52	MNS 87	99		
Parabacteroides distasonis		32.75	0.51	RHA 1	1		
Clostridium clostridioforme		2.88	0.27		50		
43517002	**Low discrimination**						**43517002**
Actinomyces israelii		71.63	0.36	SAL 92			
Lactobacillus fermentum		16.60	0.24	CEL 93			
Bifidobacterium adolescentis		11.77	0.16				
43517100	**Low discrimination**						**43517100**
Actinomyces israelii		69.23	0.43	SAL 92			
Lactobacillus fermentum		9.82	0.28	CEL 93			
Eubacterium saburreum		7.45	0.23				
43517102	**Low discrimination**						**43517102**
					GRA	CAT	
Bifidobacterium adolescentis		74.02	0.50	MNS 87	99	1	
Actinomyces israelii		18.20	0.46	ARA 87	99	1	
Capnocytophaga ochracea		4.70	0.20		1	90	
43517112	**Good identification**						**43517112**
					GRA	CAT	
Bifidobacterium adolescentis		96.25	0.52	MNS 87	99	1	
Actinomyces israelii		2.89	0.32	ARA 87	99	1	
Capnocytophaga ochracea		0.56	0.04		1	90	
43517140	**Good identification**						**43517140**
Actinomyces israelii		95.60	0.36	SAL 92			
Propionibacterium propionicum		4.40	0.12	CEL 93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43517152	**Acceptable identification**						**43517152**
Bifidobacterium adolescentis		91.36	0.36	MNS 87			
Actinomyces israelii		8.64	0.25	ARA 87			
43517500	**Low discrimination**						**43517500**
Bifidobacterium longum subsp. longum		49.68	0.37	MNS 87			
Actinomyces israelii		30.00	0.42	MAN 1			
Bifidobacterium adolescentis		15.12	0.30				
43517502	**Good identification**						**43517502**
Bifidobacterium adolescentis		93.60	0.64	MNS 87			
Bifidobacterium longum subsp. longum		3.11	0.37				
Actinomyces israelii		2.81	0.45				
43517510	**Low discrimination**						**43517510**
Bifidobacterium longum subsp. longum		78.65	0.41	MNS 87			
Bifidobacterium adolescentis		17.19	0.32	MAN 1			
Actinomyces israelii		4.17	0.28				
43517512	**Very good identification**						**43517512**
Bifidobacterium adolescentis		95.25	0.66	MNS 87			
Bifidobacterium longum subsp. longum		4.40	0.41				
Actinomyces israelii		0.35	0.31				
43517520	**Low discrimination**						**43517520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.46	0.48	MNS 87	99	1	
Bacteroides vulgatus		7.77	0.15	MAN 1	1	3	
Capnocytophaga ochracea		1.13	0.07		1	90	
43517522	**Low discrimination**						**43517522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		81.25	0.48	MNS 87	99	1	
Capnocytophaga ochracea		9.23	0.23	MAN 1	1	90	
Bifidobacterium adolescentis		6.18	0.30		99	1	
43517530	**Good identification**						**43517530**
Bifidobacterium longum subsp. longum		100.0	0.52	MNS 87			
43517532	**Acceptable identification**						**43517532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		94.16	0.52	MNS 87	99	1	
Bifidobacterium adolescentis		5.14	0.32	MAN 1	99	1	
Capnocytophaga ochracea		0.70	0.07		1	90	
43517542	**Acceptable identification**						**43517542**
Bifidobacterium adolescentis		91.36	0.49	MNS 87			
Actinomyces israelii		8.64	0.38	SOR 11			
43517552	**Good identification**						**43517552**
Bifidobacterium adolescentis		98.49	0.50	MNS 87			
Actinomyces israelii		1.14	0.24	SOR 11			
Bifidobacterium longum subsp. longum		0.37	0.07				
43521000	**Acceptable identification**						**43521000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Eubacterium saburreum		91.25	0.44	IND 99	1	85	
Fusobacterium mortiferum		5.97	0.25		1	1	
Blautia hansenii		1.49	-.03		99	99	
43521010	**Good identification**						**43521010**
Eubacterium saburreum		100.0	0.42	IND 99			
43521100	**Low discrimination**						**43521100**
					COC	GRA	
Eubacterium saburreum		56.33	0.50	IND 99	1	85	
Blautia hansenii		39.09	0.32		99	99	
Fusobacterium mortiferum		3.68	0.32		1	1	
43521102	**Low discrimination**						**43521102**
					COC		
Eubacterium saburreum		65.17	0.37	IND 99	1		
Blautia hansenii		28.48	0.15		99		
Actinomyces naeslundii		3.67	0.29		1		
43521110	**Good identification**						**43521110**
					COC		
Eubacterium saburreum		98.57	0.49	IND 99	1		
Blautia hansenii		0.85	-.03		99		
Bifidobacterium longum subsp. longum		0.59	0.07		1		
43521112	**Good identification**						**43521112**
Eubacterium saburreum		96.75	0.36	IND 99			
Bifidobacterium longum subsp. longum		3.25	0.07				
43521130	**Low discrimination**						**43521130**
					GRA		
Eubacterium saburreum		85.07	0.36	IND 99	85		
Bifidobacterium longum subsp. longum		11.44	0.17		99		
Prevotella melaninogenica		3.49	-.04		1		
43521500	**Low discrimination**						**43521500**
					COC		
Bifidobacterium longum subsp. longum		68.04	0.37	MNS 87	1		
Blautia hansenii		29.52	0.19	FRU 90	99		
Eubacterium saburreum		2.44	0.16		1		
43521502	**Acceptable identification**						**43521502**
					COC		
Bifidobacterium longum subsp. longum		91.86	0.37	MNS 87	1		
Blautia hansenii		4.43	0.03	FRU 90	99		
Bifidobacterium adolescentis		3.72	0.15		1		
43521510	**Good identification**						**43521510**
Bifidobacterium longum subsp. longum		98.31	0.41	MNS 87			
Eubacterium saburreum		1.69	0.14	FRU 90			
43521512	**Good identification**						**43521512**
Bifidobacterium longum subsp. longum		97.18	0.41	MNS 87			
Bifidobacterium adolescentis		2.82	0.16	FRU 90			
43521520	**Good identification**						**43521520**
					COC	GRA	
Bifidobacterium longum subsp. longum		98.30	0.47	MNS 87	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		1.19	0.03	FRU 90	99	99	
Prevotella melaninogenica		0.52	-.04		1	1	
43521522	**Good identification**						**43521522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.50	0.47	MNS 87	99	1	
Capnocytophaga ochracea		0.50	-.01	FRU 90	1	90	
43521530	**Good identification**						**43521530**
Bifidobacterium longum subsp. longum		100.0	0.51	MNS 87			
43521532	**Good identification**						**43521532**
Bifidobacterium longum subsp. longum		100.0	0.51	MNS 87			
43523101	**Good identification**						**43523101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43523103	**Good identification**						**43523103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43523121	**Good identification**						**43523121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43523123	**Good identification**						**43523123**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43525000	**Good identification**						**43525000**
Eubacterium saburreum		100.0	0.39	IND 99			
43525010	**Good identification**						**43525010**
Eubacterium saburreum		100.0	0.38	IND 99			
43525100	**Low discrimination**						**43525100**
					COC		
Eubacterium saburreum		79.84	0.46	IND 99	1		
Blautia hansenii		18.16	0.19		99		
Bifidobacterium longum subsp. longum		1.27	0.11		1		
43525110	**Good identification**						**43525110**
Eubacterium saburreum		96.80	0.44	IND 99			
Bifidobacterium longum subsp. longum		3.20	0.15				
43525322	**Good identification**						**43525322**
Parabacteroides distasonis		100.0	0.36	MNS 98			
43525500	**Good identification**						**43525500**
					COC		
Bifidobacterium longum subsp. longum		96.90	0.45	MNS 87	1		
Blautia hansenii		2.47	0.06	FRU 90	99		
Eubacterium saburreum		0.62	0.11		1		
43525502	**Acceptable identification**						**43525502**
Bifidobacterium longum subsp. longum		91.73	0.45	MNS 87			
Bifidobacterium adolescentis		8.27	0.29	FRU 90			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43525510	**Good identification**							**43525510**
Bifidobacterium longum subsp. longum		99.69	0.49	MNS 87				
Eubacterium saburreum		0.31	0.10	FRU 90				
43525512	**Acceptable identification**							**43525512**
Bifidobacterium longum subsp. longum		93.92	0.49	MNS 87				
Bifidobacterium adolescentis		6.08	0.30	FRU 90				
43525520	**Good identification**							**43525520**
					GRA			
Bifidobacterium longum subsp. longum		99.87	0.55	MNS 87	99			
Bacteroides vulgatus		0.13	-.08	FRU 90	1			
43525522	**Good identification**							**43525522**
					GRA			
Bifidobacterium longum subsp. longum		99.09	0.55	MNS 87	99			
Parabacteroides distasonis		0.91	0.25	FRU 90	1			
43525530	**Very good identification**							**43525530**
Bifidobacterium longum subsp. longum		100.0	0.59	MNS 87				
43525532	**Very good identification**							**43525532**
					GRA			
Bifidobacterium longum subsp. longum		99.71	0.59	MNS 87	99			
Parabacteroides distasonis		0.29	0.20	FRU 90	1			
43530000	**Good identification**							**43530000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		93.16	0.56	MLT 11	1	1	5	
Propionibacterium granulosum		2.37	0.26		99	1	90	
Clostridium botulinum biovar A		1.93	0.14		99	99	1	
43530002	**Acceptable identification**							**43530002**
					GRA	SPO	CAT	
Fusobacterium mortiferum		90.21	0.40	MLT 11	1	1	5	
Propionibacterium granulosum		5.16	0.15	BGL 10	99	1	90	
Clostridium sporogenes		4.63	0.03		99	99	1	
43530100	**Very good identification**							**43530100**
					GRA			
Fusobacterium mortiferum		99.12	0.63	MLT 11	1			
Eubacterium saburreum		0.70	0.24		85			
Actinomyces naeslundii		0.18	0.27		99			
43530102	**Good identification**							**43530102**
					GRA			
Fusobacterium mortiferum		96.34	0.46	MLT 11	1			
Actinomyces naeslundii		1.58	0.27	BGL 10	99			
Eubacterium saburreum		1.08	0.11		85			
43531000	**Low discrimination**							**43531000**
					GRA			
Fusobacterium mortiferum		65.77	0.59	MLT 11	1			
Eubacterium saburreum		30.48	0.52		85			
Actinomyces naeslundii		1.57	0.43		99			
43531001	**Low discrimination**							**43531001**
					GRA			
Actinomyces naeslundii		70.83	0.46	ESL 12	99			
Fusobacterium mortiferum		19.94	0.25	MNS 88	1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		9.24	0.18		85		
43531002	**Low discrimination**						**43531002**
					GRA		
Fusobacterium mortiferum		47.29	0.43	MLT 11	1		
Eubacterium saburreum		34.81	0.39	BGL 10	85		
Actinomyces naeslundii		10.18	0.43		99		
43531003	**Good identification**						**43531003**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43531010	**Good identification**						**43531010**
					CAT	GRA	
Eubacterium saburreum		96.83	0.50	IND 99	1	85	
Fusobacterium mortiferum		2.58	0.25		5	1	
Propionibacterium granulosum		0.59	0.11		90	99	
43531012	**Good identification**						**43531012**
Eubacterium saburreum		100.0	0.38	IND 99			
43531020	**Low discrimination**						**43531020**
Eubacterium saburreum		39.07	0.39	IND 99			
Clostridium perfringens		31.54	0.28				
Actinomyces naeslundii		11.43	0.43				
43531021	**Good identification**						**43531021**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43531023	**Good identification**						**43531023**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43531030	**Good identification**						**43531030**
Eubacterium saburreum		100.0	0.38	IND 99			
43531100	**Low discrimination**						**43531100**
					GRA		
Fusobacterium mortiferum		65.05	0.66	MLT 11	1		
Eubacterium saburreum		30.15	0.58		85		
Actinomyces naeslundii		1.90	0.51		99		
43531101	**Low discrimination**						**43531101**
					GRA	SPO	
Actinomyces naeslundii		44.58	0.54	ESL 12	99	1	
Clostridium chauvoei		40.37	0.30	MNS 88	99	99	
Fusobacterium mortiferum		10.29	0.32		1	1	
43531102	**Low discrimination**						**43531102**
					CAT		
Fusobacterium mortiferum		33.66	0.49	MLT 11	5		
Eubacterium saburreum		24.78	0.45	BGL 10	1		
Capnocytophaga ochracea		16.50	0.30		90		
43531103	**Low discrimination**						**43531103**
					GRA	SPO	
Actinomyces naeslundii		83.04	0.54	ESL 12	99	1	
Clostridium chauvoei		13.27	0.17	MNS 88	99	99	
Fusobacterium mortiferum		2.13	0.15		1	1	
43531110	**Very good identification**						**43531110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Eubacterium saburreum		95.21	0.57	IND 99	85		
Fusobacterium mortiferum		2.54	0.32		1		
Bifidobacterium longum subsp. longum		1.70	0.23		99		
43531112	**Low discrimination**						**43531112**
Eubacterium saburreum		58.19	0.44	IND 99			
Bifidobacterium adolescentis		27.73	0.36				
Bifidobacterium longum subsp. longum		5.87	0.23				
43531120	**Low discrimination**						**43531120**
					GRA		
Prevotella melaninogenica		66.68	0.47	MNS 90	1		
Clostridium perfringens		16.96	0.43	TRE 1	99		
Eubacterium saburreum		6.68	0.45		85		
43531121	**Low discrimination**						**43531121**
					GRA	SPO	
Actinomyces naeslundii		64.23	0.54	ESL 12	99	1	
Clostridium chauvoei		19.39	0.22	MNS 88	99	99	
Prevotella melaninogenica		12.09	0.12		1	1	
43531122	**Low discrimination**						**43531122**
					GRA	CAT	
Capnocytophaga ochracea		50.76	0.47	MNS 90	1	90	
Prevotella melaninogenica		16.10	0.34	TRE 1	1	1	
Clostridium perfringens		10.12	0.38		99	1	
43531123	**Low discrimination**						**43531123**
					GRA	SPO	
Actinomyces naeslundii		79.33	0.54	ESL 12	99	1	
Capnocytophaga ochracea		8.97	0.12	MNS 88	1	1	
Clostridium chauvoei		4.23	0.09		99	99	
43531130	**Low discrimination**						**43531130**
					GRA		
Eubacterium saburreum		59.83	0.44	IND 99	85		
Bifidobacterium longum subsp. longum		24.14	0.33		99		
Prevotella melaninogenica		7.37	0.12		1		
43531141	**Good identification**						**43531141**
Actinomyces naeslundii		100.0	0.41	ESL 12			
43531143	**Good identification**						**43531143**
Actinomyces naeslundii		100.0	0.41	ESL 12			
43531161	**Good identification**						**43531161**
Actinomyces naeslundii		100.0	0.41	ESL 12			
43531163	**Good identification**						**43531163**
Actinomyces naeslundii		100.0	0.41	ESL 12			
43531322	**Low discrimination**						**43531322**
					CAT		
Parabacteroides distasonis		87.34	0.54	MNS 98	76		
Prevotella oralis		10.11	0.25	XYL 90	1		
Capnocytophaga ochracea		1.68	0.12		90		
43531332	**Good identification**						**43531332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		99.51	0.50	MNS 98			
Clostridium clostridioforme		0.49	0.04	XYL 90			
43531500	**Acceptable identification**						**43531500**
					GRA		
Bifidobacterium longum subsp. longum		93.40	0.53	MNS 87	99		
Fusobacterium mortiferum		2.40	0.32		1		
Lactobacillus fermentum		2.02	0.32		99		
43531502	**Low discrimination**						**43531502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		66.57	0.53	MNS 87	99	1	
Bifidobacterium adolescentis		29.61	0.49		99	1	
Capnocytophaga ochracea		1.63	0.17		1	90	
43531510	**Very good identification**						**43531510**
Bifidobacterium longum subsp. longum		99.11	0.57	MNS 87			
Eubacterium saburreum		0.57	0.23				
Bifidobacterium adolescentis		0.32	0.16				
43531512	**Low discrimination**						**43531512**
Bifidobacterium longum subsp. longum		75.52	0.57	MNS 87			
Bifidobacterium adolescentis		24.11	0.50				
Clostridium clostridioforme		0.17	0.10				
43531520	**Very good identification**						**43531520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.43	0.63	MNS 87	99	1	
Capnocytophaga ochracea		0.60	0.17		1	90	
Prevotella melaninogenica		0.52	0.12		1	1	
43531522	**Good identification**						**43531522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.44	0.63	MNS 87	99	1	
Capnocytophaga ochracea		5.04	0.34		1	90	
Parabacteroides distasonis		3.07	0.42		1	76	
43531530	**Very good identification**						**43531530**
					CAT	GRA	
Bifidobacterium longum subsp. longum		99.94	0.67	MNS 87	1	99	
Capnocytophaga ochracea		0.04	0.01		90	1	
Eubacterium saburreum		0.03	0.10		1	85	
43531532	**Very good identification**						**43531532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.47	0.67	MNS 87	99	1	
Parabacteroides distasonis		1.04	0.38		1	76	
Capnocytophaga ochracea		0.35	0.17		1	90	
43531722	**Low discrimination**						**43531722**
					GRA		
Parabacteroides distasonis		88.16	0.51	MNS 98	1		
Bifidobacterium longum subsp. longum		8.85	0.29	XYL 90	99		
Clostridium clostridioforme		1.31	0.14		50		
43531732	**Low discrimination**						**43531732**
					GRA		
Parabacteroides distasonis		68.81	0.46	MNS 98	1		
Bifidobacterium longum subsp. longum		21.85	0.33	XYL 90	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		8.67	0.25		50		
43532100	**Low discrimination**						**43532100**
					GRA		
Propionibacterium propionicum		84.83	0.45	NIT 86	99		
Fusobacterium mortiferum		11.23	0.28	ESL 1	1		
Actinomyces naeslundii		3.94	0.32		99		
43532102	**Good identification**						**43532102**
					GRA		
Propionibacterium propionicum		96.09	0.48	NIT 86	99		
Actinomyces naeslundii		2.97	0.32	ESL 1	99		
Fusobacterium mortiferum		0.94	0.12		1		
43532121	**Good identification**						**43532121**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43532123	**Good identification**						**43532123**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43532140	**Good identification**						**43532140**
Propionibacterium propionicum		100.0	0.45	NIT 86			
43532142	**Good identification**						**43532142**
Propionibacterium propionicum		100.0	0.48	NIT 86			
43533000	**Low discrimination**						**43533000**
					GRA	CAT	
Actinomyces naeslundii		57.54	0.48	ESL 12	99	25	
Propionibacterium granulosum		15.84	0.24	MNS 88	99	90	
Fusobacterium mortiferum		12.52	0.25		1	5	
43533001	**Good identification**						**43533001**
Actinomyces naeslundii		100.0	0.51	ESL 12			
43533002	**Low discrimination**						**43533002**
					CAT		
Actinomyces naeslundii		75.53	0.48	ESL 12	25		
Actinomyces israelii		16.35	0.28	MNS 88	1		
Propionibacterium granulosum		5.20	0.13		90		
43533003	**Good identification**						**43533003**
Actinomyces naeslundii		100.0	0.51	ESL 12			
43533020	**Acceptable identification**						**43533020**
Actinomyces naeslundii		93.56	0.48	ESL 12			
Propionibacterium granulosum		6.44	0.13	MNS 88			
43533021	**Good identification**						**43533021**
Actinomyces naeslundii		100.0	0.51	ESL 12			
43533022	**Good identification**						**43533022**
Actinomyces naeslundii		100.0	0.48	ESL 12			
43533023	**Good identification**						**43533023**
Actinomyces naeslundii		100.0	0.51	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43533040	**Low discrimination**					**43533040**
Actinomyces naeslundii		75.88	0.35	ESL 12		
Actinomyces israelii		24.12	0.18	MNS 88		
43533041	**Good identification**					**43533041**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43533042	**Low discrimination**					**43533042**
Actinomyces naeslundii		58.95	0.35	ESL 12		
Actinomyces israelii		28.11	0.21	MNS 88		
Propionibacterium propionicum		12.94	0.14			
43533043	**Good identification**					**43533043**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43533060	**Good identification**					**43533060**
Actinomyces naeslundii		100.0	0.35	ESL 12		
43533061	**Good identification**					**43533061**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43533062	**Good identification**					**43533062**
Actinomyces naeslundii		100.0	0.35	ESL 12		
43533063	**Good identification**					**43533063**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43533100	**Low discrimination**					**43533100**
Actinomyces naeslundii		42.34	0.56	ESL 12		
Propionibacterium propionicum		38.02	0.45	MNS 88		
Actinomyces israelii		8.59	0.35			
43533101	**Good identification**					**43533101**
Actinomyces naeslundii		93.76	0.59	ESL 12		
Propionibacterium propionicum		6.24	0.29	MNS 88		
43533102	**Low discrimination**					**43533102**
Propionibacterium propionicum		43.99	0.48	NIT 86		
Actinomyces naeslundii		32.66	0.56	ESL 1		
Bifidobacterium adolescentis		12.30	0.33			
43533103	**Good identification**					**43533103**
Actinomyces naeslundii		90.93	0.59	ESL 12		
Propionibacterium propionicum		9.07	0.32	MNS 88		
43533111	**Good identification**					**43533111**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43533113	**Good identification**					**43533113**
Actinomyces naeslundii		100.0	0.37	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43533120	**Low discrimination**						**43533120**
					GRA		
Actinomyces naeslundii		82.64	0.56	ESL 12	99		
Prevotella melaninogenica		12.02	0.12	MNS 88	1		
Clostridium perfringens		3.06	0.09		99		
43533121	**Very good identification**						**43533121**
Actinomyces naeslundii		100.0	0.59	ESL 12			
43533122	**Low discrimination**						**43533122**
					GRA	CAT	
Actinomyces naeslundii		83.81	0.56	ESL 12	99	25	
Capnocytophaga ochracea		7.32	0.12	MNS 88	1	90	
Prevotella melaninogenica		2.32	0.00		1	1	
43533123	**Very good identification**						**43533123**
Actinomyces naeslundii		100.0	0.59	ESL 12			
43533131	**Good identification**						**43533131**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43533133	**Good identification**						**43533133**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43533140	**Low discrimination**						**43533140**
Propionibacterium propionicum		77.86	0.45	NIT 86			
Actinomyces naeslundii		15.30	0.43	ESL 1			
Actinomyces israelii		6.84	0.28				
43533141	**Low discrimination**						**43533141**
Actinomyces naeslundii		72.63	0.46	ESL 12			
Propionibacterium propionicum		27.38	0.29	MNS 88			
43533142	**Low discrimination**						**43533142**
Propionibacterium propionicum		79.78	0.48	NIT 86			
Actinomyces naeslundii		10.45	0.43	ESL 1			
Actinomyces israelii		7.01	0.31				
43533143	**Low discrimination**						**43533143**
Actinomyces naeslundii		63.88	0.46	ESL 12			
Propionibacterium propionicum		36.12	0.32	MNS 88			
43533160	**Good identification**						**43533160**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43533161	**Good identification**						**43533161**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43533162	**Acceptable identification**						**43533162**
Actinomyces naeslundii		92.84	0.43	ESL 12			
Propionibacterium propionicum		7.16	0.14	MNS 88			
43533163	**Good identification**						**43533163**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.46	ESL 12		
43533502	**Low discrimination**					**43533502**
	Bifidobacterium adolescentis	88.35	0.47	MNS 87		
	Actinomyces israelii	8.72	0.36	XYL 87		
	Bifidobacterium longum subsp. longum	2.45	0.19			
43533512	**Good identification**					**43533512**
	Bifidobacterium adolescentis	95.17	0.49	MNS 87		
	Bifidobacterium longum subsp. longum	3.68	0.23	XYL 87		
	Actinomyces israelii	1.15	0.22			
43534000	**Good identification**					**43534000**
	Fusobacterium mortiferum	95.22	0.37	MLT 11		
	Eubacterium saburreum	4.78	0.13	XYL 7		
43534100	**Good identification**					**43534100**
	Fusobacterium mortiferum	95.22	0.43	MLT 11		
	Eubacterium saburreum	4.78	0.20	XYL 7		
43534322	**Good identification**					**43534322**
	Parabacteroides distasonis	100.0	0.36	MNS 98		
43534520	**Good identification**					**43534520**
					GRA	
	Bifidobacterium longum subsp. longum	98.57	0.37	MNS 87	99	
	Bacteroides vulgatus	1.43	-.08	GAL 99	1	
43534522	**Low discrimination**					**43534522**
					GRA	
	Bifidobacterium longum subsp. longum	89.02	0.37	MNS 87	99	
	Parabacteroides distasonis	8.96	0.25	GAL 99	1	
	Clostridium clostridioforme	2.02	0.08		50	
43534530	**Good identification**					**43534530**
	Bifidobacterium longum subsp. longum	100.0	0.41	MNS 87		
43534532	**Acceptable identification**					**43534532**
					GRA	
	Bifidobacterium longum subsp. longum	91.52	0.41	MNS 87	99	
	Clostridium clostridioforme	5.57	0.19	GAL 99	50	
	Parabacteroides distasonis	2.91	0.20		1	
43535000	**Low discrimination**					**43535000**
					GRA	
	Eubacterium saburreum	72.22	0.47	IND 99	85	
	Fusobacterium mortiferum	21.78	0.40		1	
	Actinomyces israelii	3.94	0.30		99	
43535010	**Good identification**					**43535010**
	Eubacterium saburreum	99.02	0.46	IND 99		
	Actinomyces israelii	0.99	0.16			
43535100	**Low discrimination**					**43535100**
					GRA	
	Eubacterium saburreum	65.00	0.54	IND 99	85	
	Fusobacterium mortiferum	19.61	0.46		1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		6.07	0.40		99		
43535102	**Low discrimination**						**43535102**
Bifidobacterium adolescentis		57.17	0.49	MNS 87			
Eubacterium saburreum		14.42	0.41	ARA 87			
Actinomyces israelii		11.45	0.43				
43535110	**Low discrimination**						**43535110**
Eubacterium saburreum		88.46	0.52	IND 99			
Bifidobacterium longum subsp. longum		8.77	0.31				
Actinomyces israelii		1.51	0.26				
43535112	**Low discrimination**						**43535112**
Bifidobacterium adolescentis		75.79	0.50	MNS 87			
Eubacterium saburreum		12.80	0.39	ARA 87			
Bifidobacterium longum subsp. longum		7.19	0.31				
43535120	**Low discrimination**						**43535120**
					CAT	GRA	
Bifidobacterium longum subsp. longum		32.28	0.37	MNS 87	1	99	
Eubacterium saburreum		29.88	0.41	ARA 99	1	85	
Capnocytophaga ochracea		9.24	0.20		90	1	
43535122	**Low discrimination**						**43535122**
Parabacteroides distasonis		64.40	0.62	MNS 98			
Bacteroides fragilis		19.30	0.32				
Capnocytophaga ochracea		10.21	0.36				
43535130	**Low discrimination**						**43535130**
					GRA		
Bifidobacterium longum subsp. longum		65.99	0.41	MNS 87	99		
Eubacterium saburreum		29.35	0.39	ARA 99	85		
Parabacteroides distasonis		3.43	0.24		1		
43535132	**Low discrimination**						**43535132**
					GRA	CAT	
Parabacteroides distasonis		79.64	0.58	MNS 98	1	76	
Bifidobacterium longum subsp. longum		15.50	0.41		99	1	
Capnocytophaga ochracea		2.61	0.20		1	90	
43535222	**Good identification**						**43535222**
Parabacteroides distasonis		100.0	0.36	MNS 98			
43535302	**Low discrimination**						**43535302**
					GRA		
Parabacteroides distasonis		67.44	0.36	MNS 98	1		
Eubacterium contortum		15.54	0.14	NAG 99	99		
Actinomyces israelii		7.58	0.24		99		
43535322	**Very good identification**						**43535322**
					CAT		
Parabacteroides distasonis		99.58	0.71	MNS 98	76		
Prevotella oralis		0.16	0.10		1		
Bacteroides fragilis		0.10	-.03		99		
43535323	**Good identification**						**43535323**
Parabacteroides distasonis		100.0	0.36	MNS 98			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43535332	**Very good identification**					**43535332**
					GRA	
Parabacteroides distasonis		99.38	0.66	MNS 98	1	
Clostridium clostridioforme		0.55	0.22		50	
Bifidobacterium longum subsp. longum		0.07	0.07		99	
43535362	**Good identification**					**43535362**
Parabacteroides distasonis		100.0	0.36	MNS 98		
43535420	**Good identification**					**43535420**
					GRA	
Bifidobacterium longum subsp. longum		96.54	0.37	MNS 87	99	
Lactobacillus fermentum		2.06	0.16	LAC 99	99	
Bacteroides vulgatus		1.40	-.08		1	
43535422	**Low discrimination**					**43535422**
					GRA	
Bifidobacterium longum subsp. longum		87.54	0.37	MNS 87	99	
Parabacteroides distasonis		8.81	0.25	LAC 99	1	
Lactobacillus fermentum		1.87	0.16		99	
43535430	**Good identification**					**43535430**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87		
43535432	**Acceptable identification**					**43535432**
					GRA	
Bifidobacterium longum subsp. longum		92.05	0.41	MNS 87	99	
Clostridium clostridioforme		5.02	0.18	LAC 99	50	
Parabacteroides distasonis		2.93	0.20		1	
43535500	**Very good identification**					**43535500**
Bifidobacterium longum subsp. longum		96.63	0.61	MNS 87		
Actinomyces israelii		1.57	0.39			
Bifidobacterium adolescentis		0.97	0.29			
43535502	**Low discrimination**					**43535502**
Bifidobacterium longum subsp. longum		49.27	0.61	MNS 87		
Bifidobacterium adolescentis		48.89	0.63			
Actinomyces israelii		1.20	0.42			
43535510	**Very good identification**					**43535510**
Bifidobacterium longum subsp. longum		99.05	0.65	MNS 87		
Bifidobacterium adolescentis		0.71	0.30			
Actinomyces israelii		0.14	0.24			
43535512	**Low discrimination**					**43535512**
Bifidobacterium longum subsp. longum		58.04	0.65	MNS 87		
Bifidobacterium adolescentis		41.34	0.64			
Clostridium clostridioforme		0.43	0.27			
43535520	**Very good identification**					**43535520**
					GRA	
Bifidobacterium longum subsp. longum		98.37	0.72	MNS 87	99	
Bacteroides vulgatus		1.42	0.26		1	
Parabacteroides distasonis		0.10	0.25		1	
43535521	**Good identification**					**43535521**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium longum subsp. longum		98.57	0.37	MNS 87	99		
Bacteroides vulgatus		1.43	-.08	URE 1	1		
43535522	**Good identification**						**43535522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.04	0.72	MNS 87	99	1	
Parabacteroides distasonis		9.06	0.59		1	76	
Capnocytophaga ochracea		0.41	0.23		1	90	
43535523	**Acceptable identification**						**43535523**
					GRA		
Bifidobacterium longum subsp. longum		90.86	0.37	MNS 87	99		
Parabacteroides distasonis		9.14	0.25	URE 1	1		
43535530	**Very good identification**						**43535530**
					GRA		
Bifidobacterium longum subsp. longum		99.96	0.76	MNS 87	99		
Parabacteroides distasonis		0.03	0.20		1		
Bacteroides vulgatus		0.01	-.08		1		
43535531	**Good identification**						**43535531**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87			
43535532	**Very good identification**						**43535532**
					GRA		
Bifidobacterium longum subsp. longum		96.53	0.76	MNS 87	99		
Parabacteroides distasonis		3.07	0.54		1		
Clostridium clostridioforme		0.18	0.27		50		
43535533	**Good identification**						**43535533**
					GRA		
Bifidobacterium longum subsp. longum		96.92	0.41	MNS 87	99		
Parabacteroides distasonis		3.08	0.20	URE 1	1		
43535542	**Low discrimination**						**43535542**
Bifidobacterium adolescentis		86.25	0.47	MNS 87			
Bifidobacterium longum subsp. longum		7.10	0.27	SOR 11			
Actinomyces israelii		6.65	0.35				
43535552	**Low discrimination**						**43535552**
Bifidobacterium adolescentis		88.96	0.49	MNS 87			
Bifidobacterium longum subsp. longum		10.21	0.31	SOR 11			
Actinomyces israelii		0.84	0.20				
43535560	**Good identification**						**43535560**
					GRA		
Bifidobacterium longum subsp. longum		98.57	0.37	MNS 87	99		
Bacteroides vulgatus		1.43	-.08	SOR 1	1		
43535562	**Low discrimination**						**43535562**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.56	0.37	MNS 87	99	1	
Parabacteroides distasonis		8.41	0.25	SOR 1	1	76	
Capnocytophaga ochracea		4.22	0.07		1	90	
43535570	**Good identification**						**43535570**
Bifidobacterium longum subsp. longum		100.0	0.41	MNS 87			
43535572	**Acceptable identification**						**43535572**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium longum subsp. longum		94.89	0.41	MNS 87	99		
Parabacteroides distasonis		3.02	0.20	SOR 1	1		
Bifidobacterium adolescentis		2.09	0.15		99		
43535712	**Low discrimination**						**43535712**
Clostridium clostridioforme		71.88	0.42	MNS 99			
Bifidobacterium longum subsp. longum		13.43	0.31	SAL 88			
Bifidobacterium adolescentis		9.57	0.30				
43535720	**Acceptable identification**						**43535720**
					GRA		
Bacteroides vulgatus		90.25	0.42	MNS 99	1		
Bifidobacterium longum subsp. longum		6.99	0.37	TRE 4	99		
Parabacteroides distasonis		2.11	0.33		1		
43535722	**Good identification**						**43535722**
					GRA	CAT	
Parabacteroides distasonis		91.66	0.67	MNS 98	1	76	
Bacteroides thetaiotaomicron		3.29	0.26		1	89	
Bifidobacterium longum subsp. longum		3.07	0.37		99	1	
43535730	**Low discrimination**						**43535730**
					GRA		
Bifidobacterium longum subsp. longum		84.35	0.41	MNS 87	99		
Parabacteroides distasonis		8.05	0.28	RHA 1	1		
Bacteroides vulgatus		6.46	0.08		1		
43535732	**Low discrimination**						**43535732**
					GRA		
Parabacteroides distasonis		78.53	0.62	MNS 98	1		
Clostridium clostridioforme		11.12	0.42		50		
Bifidobacterium longum subsp. longum		8.31	0.41		99		
43537000	**Good identification**						**43537000**
Actinomyces israelii		96.83	0.43	SAL 92			
Eubacterium saburreum		3.17	0.13	CEL 93			
43537002	**Good identification**						**43537002**
Actinomyces israelii		97.17	0.46	SAL 92			
Bifidobacterium adolescentis		2.83	0.13	CEL 93			
43537040	**Good identification**						**43537040**
Actinomyces israelii		100.0	0.36	SAL 92			
43537042	**Good identification**						**43537042**
Actinomyces israelii		100.0	0.39	SAL 92			
43537100	**Good identification**						**43537100**
Actinomyces israelii		96.55	0.53	SAL 92			
Eubacterium saburreum		1.84	0.20	CEL 93			
Bifidobacterium adolescentis		1.06	0.13				
43537102	**Low discrimination**						**43537102**
Actinomyces israelii		57.98	0.56	SAL 92			
Bifidobacterium adolescentis		41.80	0.47	CEL 93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		0.23	0.14				
43537110	**Low discrimination**						**43537110**
Actinomyces israelii		83.76	0.39	SAL 92			
Eubacterium saburreum		8.75	0.18	CEL 93			
Bifidobacterium adolescentis		7.48	0.15	MLZ 13			
43537112	**Low discrimination**						**43537112**
Bifidobacterium adolescentis		85.50	0.49	MNS 87			
Actinomyces israelii		14.50	0.42	ARA 87			
43537122	**Low discrimination**						**43537122**
					GRA	CAT	
Actinomyces israelii		49.64	0.36	SAL 92	99	1	
Parabacteroides distasonis		30.65	0.28	CEL 93	1	76	
Bacteroides fragilis		9.18	-.03	NAG 6	1	99	
43537140	**Good identification**						**43537140**
Actinomyces israelii		100.0	0.46	SAL 92			
43537142	**Low discrimination**						**43537142**
Actinomyces israelii		80.70	0.49	SAL 92			
Bifidobacterium adolescentis		18.49	0.32	CEL 93			
Propionibacterium propionicum		0.81	0.14				
43537302	**Acceptable identification**						**43537302**
Actinomyces israelii		91.18	0.37	SAL 92			
Bifidobacterium adolescentis		8.82	0.13	CEL 93			
43537322	**Good identification**						**43537322**
					GRA		
Parabacteroides distasonis		96.10	0.36	MNS 98	1		
Actinomyces israelii		3.91	0.16	MAN 1	99		
43537400	**Good identification**						**43537400**
Actinomyces israelii		100.0	0.41	SAL 92			
43537402	**Low discrimination**						**43537402**
Actinomyces israelii		80.76	0.44	SAL 92			
Bifidobacterium adolescentis		19.24	0.27	CEL 93			
43537442	**Acceptable identification**						**43537442**
Actinomyces israelii		92.96	0.37	SAL 92			
Bifidobacterium adolescentis		7.04	0.12	CEL 93			
43537500	**Low discrimination**						**43537500**
Actinomyces israelii		83.38	0.52	SAL 92			
Bifidobacterium longum subsp. longum		9.18	0.27	CEL 93			
Bifidobacterium adolescentis		7.45	0.27				
43537502	**Low discrimination**						**43537502**
Bifidobacterium adolescentis		84.60	0.61	MNS 87			
Actinomyces israelii		14.35	0.55				
Bifidobacterium longum subsp. longum		1.05	0.27				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
43537512	**Very good identification**							**43537512**
Bifidobacterium adolescentis		96.24	0.63	MNS 87				
Actinomyces israelii		2.00	0.40					
Bifidobacterium longum subsp. longum		1.67	0.31					
43537520	**Low discrimination**							**43537520**
					GRA			
Bifidobacterium longum subsp. longum		86.25	0.37	MNS 87	99			
Actinomyces israelii		12.51	0.31	MAN 1	99			
Bacteroides vulgatus		1.25	-.08		1			
43537522	**Low discrimination**							**43537522**
Bifidobacterium longum subsp. longum		65.74	0.37	MNS 87				
Actinomyces israelii		14.30	0.34	MAN 1				
Bifidobacterium adolescentis		13.34	0.27					
43537530	**Good identification**							**43537530**
Bifidobacterium longum subsp. longum		98.74	0.41	MNS 87				
Actinomyces israelii		1.26	0.17	MAN 1				
43537532	**Low discrimination**							**43537532**
					GRA			
Bifidobacterium longum subsp. longum		81.88	0.41	MNS 87	99			
Bifidobacterium adolescentis		11.93	0.29	MAN 1	99			
Parabacteroides distasonis		2.61	0.20		1			
43537540	**Good identification**							**43537540**
Actinomyces israelii		97.24	0.44	SAL 92				
Bifidobacterium adolescentis		2.76	0.12	CEL 93				
43537542	**Low discrimination**							**43537542**
Bifidobacterium adolescentis		65.21	0.46	MNS 87				
Actinomyces israelii		34.79	0.47	SOR 11				
43537552	**Acceptable identification**							**43537552**
Bifidobacterium adolescentis		93.88	0.47	MNS 87				
Actinomyces israelii		6.12	0.33	SOR 11				
43537702	**Low discrimination**							**43537702**
Actinomyces israelii		55.82	0.35	SAL 92				
Bifidobacterium adolescentis		44.18	0.27	CEL 93				
43541120	**Good identification**							**43541120**
					COC	CAT	GRA	
Prevotella melaninogenica		97.21	0.48	MNS 90	1	1	1	
Capnocytophaga ochracea		1.33	0.20	FRU 90	1	90	1	
Blautia hansenii		1.16	0.15	CEL 10	99	1	99	
43541122	**Low discrimination**							**43541122**
					CAT			
Prevotella melaninogenica		53.52	0.36	MNS 90	1			
Capnocytophaga ochracea		34.52	0.36	FRU 90	90			
Prevotella oralis		11.01	0.27	CEL 10	1			
43545502	**Low discrimination**							**43545502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		85.64	0.37	MNS 87		
Bifidobacterium longum subsp. longum		12.04	0.21	FRU 99		
Actinomyces israelii		2.32	0.17			
43545512	**Low discrimination**					**43545512**
Bifidobacterium adolescentis		83.63	0.39	MNS 87		
Bifidobacterium longum subsp. longum		16.37	0.25	FRU 99		
43545530	**Good identification**					**43545530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87		
43545532	**Good identification**					**43545532**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87		
43547502	**Low discrimination**					**43547502**
Bifidobacterium adolescentis		84.18	0.36	MNS 87		
Actinomyces israelii		15.82	0.30	FRU 99		
43547512	**Good identification**					**43547512**
Bifidobacterium adolescentis		97.75	0.37	MNS 87		
Actinomyces israelii		2.25	0.16	FRU 99		
43550000	**Very good identification**					**43550000**
Fusobacterium mortiferum		99.72	0.57	MLT 11		
Prevotella intermedia		0.28	0.02			
43550002	**Good identification**					**43550002**
Fusobacterium mortiferum		100.0	0.41	MLT 11		
43550100	**Very good identification**					**43550100**
					CAT	
Fusobacterium mortiferum		99.56	0.63	MLT 11	5	
Capnocytophaga ochracea		0.20	0.04		90	
Prevotella melaninogenica		0.12	-.04		1	
43550102	**Low discrimination**					**43550102**
					CAT	
Fusobacterium mortiferum		82.15	0.47	MLT 11	5	
Capnocytophaga ochracea		13.53	0.20	BGL 10	90	
Prevotella oralis		2.04	0.06		1	
43550122	**Low discrimination**					**43550122**
					CAT	
Capnocytophaga ochracea		67.50	0.36	MNS 90	90	
Prevotella oralis		21.52	0.27	GAL 99	1	
Prevotella melaninogenica		9.51	0.18		1	
43551000	**Very good identification**					**43551000**
					GRA	
Fusobacterium mortiferum		95.66	0.60	MLT 11	1	
Actinomyces naeslundii		1.40	0.40		99	
Lactobacillus fermentum		1.04	0.28		99	
43551001	**Low discrimination**					**43551001**
					GRA	
Actinomyces naeslundii		68.49	0.43	ESL 12	99	
Fusobacterium mortiferum		31.51	0.26	MNS 88	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43551002	**Low discrimination**						**43551002**
					CAT		
Fusobacterium mortiferum		55.67	0.44	MLT 11	5		
Capnocytophaga ochracea		14.26	0.20	BGL 10	90		
Prevotella oralis		13.61	0.19		1		
43551003	**Good identification**						**43551003**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43551021	**Acceptable identification**						**43551021**
					GRA		
Actinomyces naeslundii		91.98	0.43	ESL 12	99		
Prevotella melaninogenica		8.02	-.04	MNS 88	1		
43551022	**Low discrimination**						**43551022**
					CAT		
Prevotella oralis		62.27	0.41	MNS 99	1		
Capnocytophaga ochracea		30.91	0.36	LAC 94	90		
Prevotella melaninogenica		4.36	0.18		1		
43551023	**Low discrimination**						**43551023**
					GRA	CAT	
Actinomyces naeslundii		73.79	0.43	ESL 12	99	25	
Prevotella oralis		17.52	0.07	MNS 88	1	1	
Capnocytophaga ochracea		8.70	0.02		1	90	
43551100	**Low discrimination**						**43551100**
					CAT		
Fusobacterium mortiferum		78.85	0.66	MLT 11	5		
Capnocytophaga ochracea		10.58	0.38		90		
Prevotella melaninogenica		6.41	0.30		1		
43551101	**Low discrimination**						**43551101**
					GRA	SPO	
Actinomyces naeslundii		63.34	0.51	ESL 12	99	1	
Fusobacterium mortiferum		23.88	0.32	MNS 88	1	1	
Clostridium chauvoei		7.64	0.12		99	99	
43551102	**Low discrimination**						**43551102**
					CAT		
Capnocytophaga ochracea		73.74	0.54	MNS 90	90		
Prevotella oralis		11.13	0.40	NAG 90	1		
Fusobacterium mortiferum		6.78	0.50		5		
43551103	**Low discrimination**						**43551103**
					GRA	CAT	
Actinomyces naeslundii		61.90	0.51	ESL 12	99	25	
Capnocytophaga ochracea		28.19	0.20	MNS 88	1	90	
Prevotella oralis		4.26	0.06		1	1	
43551112	**Low discrimination**						**43551112**
					GRA	CAT	
Capnocytophaga ochracea		55.92	0.38	MNS 90	1	90	
Bifidobacterium adolescentis		37.95	0.45	MLZ 10	99	1	
Bifidobacterium breve		3.26	0.27	NAG 90	99	1	
43551120	**Low discrimination**						**43551120**
					CAT		
Prevotella melaninogenica		84.78	0.64	MNS 90	1		
Capnocytophaga ochracea		12.73	0.54	CEL 10	90		
Prevotella oralis		1.92	0.40		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43551122	**Low discrimination**						**43551122**
					CAT		
Capnocytophaga ochracea		67.89	0.70	MNS 90	90		
Prevotella oralis		21.64	0.62		1		
Prevotella melaninogenica		9.57	0.52		1		
43551123	**Low discrimination**						**43551123**
					GRA	CAT	
Capnocytophaga ochracea		58.42	0.36	MNS 90	1	90	
Prevotella oralis		18.62	0.27	URE 1	1	1	
Actinomyces naeslundii		14.25	0.51		99	25	
43551130	**Low discrimination**						**43551130**
					CAT		
Capnocytophaga ochracea		60.59	0.38	MNS 90	90		
Prevotella melaninogenica		36.68	0.30	MLZ 10	1		
Prevotella oralis		0.83	0.06	BGL 90	1		
43551132	**Acceptable identification**						**43551132**
					CAT		
Capnocytophaga ochracea		94.30	0.54	MNS 90	90		
Prevotella oralis		2.73	0.27	MLZ 10	1		
Prevotella melaninogenica		1.21	0.18		1		
43551141	**Good identification**						**43551141**
Actinomyces naeslundii		100.0	0.38	ESL 12			
43551142	**Low discrimination**						**43551142**
					GRA	CAT	
Capnocytophaga ochracea		83.31	0.38	MNS 90	1	90	
Bifidobacterium adolescentis		5.72	0.28	NAG 90	99	1	
Bifidobacterium breve		4.85	0.27	SOR 10	99	1	
43551143	**Low discrimination**						**43551143**
					GRA		
Actinomyces naeslundii		77.72	0.38	ESL 12	99		
Capnocytophaga ochracea		22.28	0.04	MNS 88	1		
43551160	**Low discrimination**						**43551160**
					GRA	CAT	
Capnocytophaga ochracea		60.07	0.38	MNS 90	1	90	
Prevotella melaninogenica		36.36	0.30	SOR 10	1	1	
Actinomyces naeslundii		1.41	0.35	BGL 90	99	25	
43551161	**Low discrimination**						**43551161**
					GRA	CAT	
Actinomyces naeslundii		68.48	0.38	ESL 12	99	25	
Capnocytophaga ochracea		19.64	0.04	MNS 88	1	90	
Prevotella melaninogenica		11.89	-.04		1	1	
43551162	**Acceptable identification**						**43551162**
					CAT		
Capnocytophaga ochracea		94.64	0.54	MNS 90	90		
Prevotella oralis		2.74	0.27	SOR 10	1		
Prevotella melaninogenica		1.21	0.18		1		
43551172	**Acceptable identification**						**43551172**
					GRA	CAT	
Capnocytophaga ochracea		90.61	0.38	MNS 90	1	90	
Bifidobacterium breve		9.39	0.32	MLZ 10	99	1	
43551322	**Acceptable identification**						**43551322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Prevotella oralis		90.60	0.55	MNS 99	1		
Capnocytophaga ochracea		6.70	0.36		90		
Parabacteroides distasonis		1.67	0.38		76		
43551502	**Low discrimination**						**43551502**
					GRA	CAT	
Bifidobacterium adolescentis		66.98	0.57	MNS 87	99	1	
Capnocytophaga ochracea		28.63	0.41	XYL 87	1	90	
Bifidobacterium longum subsp. longum		1.91	0.29		99	1	
43551512	**Good identification**						**43551512**
					GRA	CAT	
Bifidobacterium adolescentis		92.47	0.59	MNS 87	99	1	
Bifidobacterium longum subsp. longum		3.67	0.33	XYL 87	99	1	
Capnocytophaga ochracea		3.59	0.25		1	90	
43551520	**Low discrimination**						**43551520**
					GRA	CAT	
Capnocytophaga ochracea		59.68	0.41	MNS 90	1	90	
Prevotella melaninogenica		22.75	0.30	BGL 90	1	1	
Bifidobacterium longum subsp. longum		15.91	0.40		99	1	
43551522	**Good identification**						**43551522**
					GRA	CAT	
Capnocytophaga ochracea		94.05	0.58	MNS 90	1	90	
Bifidobacterium longum subsp. longum		2.79	0.40		99	1	
Prevotella oralis		1.72	0.27		1	1	
43551530	**Low discrimination**						**43551530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		79.80	0.44	MNS 87	99	1	
Capnocytophaga ochracea		19.53	0.25	CEL 1	1	90	
Prevotella melaninogenica		0.68	-.04		1	1	
43551532	**Low discrimination**						**43551532**
					GRA	CAT	
Capnocytophaga ochracea		66.67	0.41	MNS 90	1	90	
Bifidobacterium longum subsp. longum		30.27	0.44	MLZ 10	99	1	
Bifidobacterium adolescentis		1.93	0.25		99	1	
43551542	**Low discrimination**						**43551542**
					GRA	CAT	
Bifidobacterium adolescentis		69.74	0.42	MNS 87	99	1	
Capnocytophaga ochracea		26.80	0.25	XYL 87	1	90	
Actinomyces israelii		3.46	0.26	SOR 11	99	1	
43551552	**Good identification**						**43551552**
					GRA	CAT	
Bifidobacterium adolescentis		96.62	0.43	MNS 87	99	1	
Capnocytophaga ochracea		3.38	0.08	XYL 87	1	90	
43551562	**Good identification**						**43551562**
					GRA	CAT	
Capnocytophaga ochracea		99.44	0.41	MNS 90	1	90	
Bifidobacterium adolescentis		0.29	0.08	SOR 10	99	1	
Bifidobacterium longum subsp. longum		0.27	0.05		99	1	
43553000	**Low discrimination**						**43553000**
					GRA		
Actinomyces naeslundii		47.59	0.45	ESL 12	99		
Actinomyces israelii		35.50	0.34	MNS 88	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		16.91	0.26		1		
43553001	**Good identification**						**43553001**
Actinomyces naeslundii		100.0	0.48	ESL 12			
43553002	**Low discrimination**						**43553002**
Actinomyces israelii		51.58	0.37	SAL 92			
Actinomyces naeslundii		46.09	0.45	XYL 92			
Bifidobacterium adolescentis		2.33	0.08				
43553003	**Good identification**						**43553003**
Actinomyces naeslundii		100.0	0.48	ESL 12			
43553020	**Low discrimination**						**43553020**
					GRA		
Actinomyces naeslundii		89.69	0.45	ESL 12	99		
Prevotella melaninogenica		6.04	-.04	MNS 88	1		
Actinomyces israelii		4.27	0.13		99		
43553021	**Good identification**						**43553021**
Actinomyces naeslundii		100.0	0.48	ESL 12			
43553022	**Low discrimination**						**43553022**
					GRA	CAT	
Actinomyces naeslundii		74.30	0.45	ESL 12	99	25	
Prevotella oralis		13.63	0.07	MNS 88	1	1	
Capnocytophaga ochracea		6.77	0.02		1	90	
43553023	**Good identification**						**43553023**
Actinomyces naeslundii		100.0	0.48	ESL 12			
43553041	**Good identification**						**43553041**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43553043	**Good identification**						**43553043**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43553061	**Good identification**						**43553061**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43553063	**Good identification**						**43553063**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43553100	**Low discrimination**						**43553100**
					GRA		
Actinomyces israelii		42.64	0.44	SAL 92	99		
Actinomyces naeslundii		40.67	0.53	XYL 92	99		
Fusobacterium mortiferum		11.85	0.32		1		
43553101	**Good identification**						**43553101**
Actinomyces naeslundii		100.0	0.56	ESL 12			
43553102	**Low discrimination**						**43553102**
Bifidobacterium adolescentis		35.04	0.42	MNS 87			
Actinomyces israelii		31.34	0.47	XYL 87			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		19.93	0.53	ARA 87			
43553103	**Good identification**						**43553103**
Actinomyces naeslundii		98.83	0.56	ESL 12			
Bifidobacterium adolescentis		1.17	0.08	MNS 88			
43553112	**Low discrimination**						**43553112**
Bifidobacterium adolescentis		79.56	0.43	MNS 87			
Actinomyces israelii		8.70	0.33	XYL 87			
Bifidobacterium breve		8.35	0.27	ARA 87			
43553121	**Good identification**						**43553121**
					GRA		
Actinomyces naeslundii		98.45	0.56	ESL 12	99		
Prevotella melaninogenica		1.55	-.04	MNS 88	1		
43553122	**Low discrimination**						**43553122**
					GRA	CAT	
Capnocytophaga ochracea		50.68	0.36	MNS 90	1	90	
Prevotella oralis		16.15	0.27	MAN 1	1	1	
Actinomyces naeslundii		16.00	0.53		99	25	
43553123	**Good identification**						**43553123**
					GRA		
Actinomyces naeslundii		97.91	0.56	ESL 12	99		
Capnocytophaga ochracea		2.09	0.02	MNS 88	1		
43553140	**Low discrimination**						**43553140**
Actinomyces israelii		65.48	0.37	SAL 92			
Actinomyces naeslundii		28.34	0.40	XYL 92			
Bifidobacterium breve		3.58	0.10				
43553141	**Good identification**						**43553141**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43553142	**Low discrimination**						**43553142**
Actinomyces israelii		47.25	0.40	SAL 92			
Bifidobacterium breve		17.43	0.27	XYL 92			
Bifidobacterium adolescentis		16.79	0.26				
43553143	**Good identification**						**43553143**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43553160	**Low discrimination**						**43553160**
					GRA	CAT	
Actinomyces naeslundii		57.87	0.40	ESL 12	99	25	
Bifidobacterium breve		13.01	0.15	MNS 88	99	1	
Capnocytophaga ochracea		12.82	0.04		1	90	
43553161	**Good identification**						**43553161**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43553163	**Good identification**						**43553163**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43553402	**Low discrimination**						**43553402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		73.02	0.35	SAL 92			
Bifidobacterium adolescentis		26.98	0.22	XYL 92			
43553500	**Low discrimination**						**43553500**
Actinomyces israelii		87.83	0.43	SAL 92			
Bifidobacterium adolescentis		12.17	0.22	XYL 92			
43553502	**Low discrimination**						**43553502**
					GRA	CAT	
Bifidobacterium adolescentis		89.72	0.56	MNS 87	99	1	
Actinomyces israelii		9.81	0.46	XYL 87	99	1	
Capnocytophaga ochracea		0.47	0.07		1	90	
43553512	**Very good identification**						**43553512**
Bifidobacterium adolescentis		98.68	0.57	MNS 87			
Actinomyces israelii		1.32	0.32	XYL 87			
43553540	**Good identification**						**43553540**
Actinomyces israelii		100.0	0.36	SAL 92			
43553542	**Low discrimination**						**43553542**
Bifidobacterium adolescentis		74.40	0.40	MNS 87			
Actinomyces israelii		25.60	0.39	XYL 87			
43553552	**Good identification**						**43553552**
Bifidobacterium adolescentis		95.96	0.42	MNS 87			
Actinomyces israelii		4.04	0.25	XYL 87			
43554000	**Good identification**						**43554000**
Fusobacterium mortiferum		100.0	0.38	MLT 11			
43554100	**Good identification**						**43554100**
					GRA		
Fusobacterium mortiferum		98.54	0.44	MLT 11	1		
Actinomyces israelii		1.46	0.15	XYL 7	99		
43554502	**Good identification**						**43554502**
Bifidobacterium adolescentis		97.36	0.37	MNS 87			
Actinomyces israelii		2.64	0.17	GAL 99			
43554512	**Acceptable identification**						**43554512**
Bifidobacterium adolescentis		93.75	0.39	MNS 87			
Clostridium clostridioforme		4.59	0.13	GAL 99			
Bifidobacterium longum subsp. longum		1.67	0.07				
43555000	**Low discrimination**						**43555000**
					GRA		
Fusobacterium mortiferum		58.77	0.41	MLT 11	1		
Actinomyces israelii		33.60	0.39	XYL 7	99		
Lactobacillus fermentum		4.98	0.24		99		
43555002	**Low discrimination**						**43555002**
					GRA		
Actinomyces israelii		62.30	0.42	SAL 92	99		
Bifidobacterium adolescentis		11.34	0.23		99		
Fusobacterium mortiferum		8.07	0.25		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43555042	**Acceptable identification**						**43555042**
Actinomyces israelii		94.53	0.35	SAL 92			
Bifidobacterium adolescentis		5.47	0.08				
43555100	**Low discrimination**						**43555100**
					GRA	CAT	
Fusobacterium mortiferum		37.08	0.47	MLT 11	1	5	
Actinomyces israelii		36.34	0.49	XYL 7	99	1	
Capnocytophaga ochracea		16.53	0.27		1	90	
43555102	**Low discrimination**						**43555102**
					GRA	CAT	
Bifidobacterium adolescentis		48.09	0.57	MNS 87	99	1	
Capnocytophaga ochracea		29.12	0.44	ARA 87	1	90	
Actinomyces israelii		10.67	0.52		99	1	
43555110	**Low discrimination**						**43555110**
					GRA	CAT	
Actinomyces israelii		43.79	0.35	SAL 92	99	1	
Bifidobacterium adolescentis		24.46	0.25	MLZ 13	99	1	
Capnocytophaga ochracea		14.81	0.11		1	90	
43555112	**Good identification**						**43555112**
					GRA	CAT	
Bifidobacterium adolescentis		92.18	0.59	MNS 87	99	1	
Capnocytophaga ochracea		5.07	0.27	ARA 87	1	90	
Actinomyces israelii		2.50	0.38		99	1	
43555120	**Low discrimination**						**43555120**
					CAT		
Bacteroides fragilis		52.37	0.38	MNS 99	99		
Capnocytophaga ochracea		34.11	0.44	BGL 95	90		
Prevotella melaninogenica		9.18	0.30		1		
43555122	**Low discrimination**						**43555122**
					CAT		
Bacteroides fragilis		72.98	0.60	MNS 99	99		
Capnocytophaga ochracea		22.52	0.60		90		
Prevotella oralis		3.55	0.46		1		
43555132	**Low discrimination**						**43555132**
Capnocytophaga ochracea		67.18	0.44	MNS 90			
Bacteroides fragilis		19.79	0.25	MLZ 10			
Parabacteroides distasonis		9.81	0.42				
43555140	**Low discrimination**						**43555140**
					GRA	CAT	
Actinomyces israelii		84.88	0.42	SAL 92	99	1	
Capnocytophaga ochracea		11.03	0.11		1	90	
Fusobacterium mortiferum		2.25	0.13		1	5	
43555142	**Low discrimination**						**43555142**
					GRA	CAT	
Bifidobacterium adolescentis		44.30	0.42	MNS 87	99	1	
Actinomyces israelii		30.93	0.45	ARA 87	99	1	
Capnocytophaga ochracea		24.12	0.27	SOR 11	1	90	
43555152	**Low discrimination**						**43555152**
					GRA	CAT	
Bifidobacterium adolescentis		88.12	0.43	MNS 87	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		7.52	0.31	ARA 87	99	1	
Capnocytophaga ochracea		4.36	0.11	SOR 11	1	90	
43555162	**Low discrimination**						**43555162**
					CAT		
Capnocytophaga ochracea		75.59	0.44	MNS 90	90		
Bacteroides fragilis		22.27	0.25	SOR 10	99		
Prevotella oralis		1.08	0.12		1		
43555322	**Low discrimination**						**43555322**
					CAT		
Parabacteroides distasonis		44.72	0.55	MNS 98	76		
Prevotella oralis		33.42	0.40		1		
Bacteroides fragilis		16.20	0.25		99		
43555332	**Acceptable identification**						**43555332**
Parabacteroides distasonis		93.87	0.50	MNS 98			
Capnocytophaga ochracea		2.17	0.10				
Clostridium clostridioforme		2.01	0.16				
43555400	**Low discrimination**						**43555400**
Actinomyces israelii		84.66	0.38	SAL 92			
Lactobacillus fermentum		15.34	0.24				
43555402	**Low discrimination**						**43555402**
Bifidobacterium adolescentis		57.06	0.37	MNS 87			
Actinomyces israelii		38.31	0.41	LAC 99			
Lactobacillus fermentum		4.63	0.24				
43555412	**Low discrimination**						**43555412**
Bifidobacterium adolescentis		87.43	0.39	MNS 87			
Actinomyces israelii		7.18	0.27	LAC 99			
Clostridium clostridioforme		3.83	0.12				
43555500	**Low discrimination**						**43555500**
Actinomyces israelii		38.70	0.48	SAL 92			
Bifidobacterium longum subsp. longum		27.34	0.37				
Bifidobacterium adolescentis		21.62	0.37				
43555502	**Good identification**						**43555502**
					GRA	CAT	
Bifidobacterium adolescentis		94.45	0.72	MNS 87	99	1	
Actinomyces israelii		2.56	0.51		99	1	
Capnocytophaga ochracea		1.51	0.31		1	90	
43555503	**Good identification**						**43555503**
Bifidobacterium adolescentis		97.36	0.37	MNS 87			
Actinomyces israelii		2.64	0.17	URE 1			
43555510	**Low discrimination**						**43555510**
Bifidobacterium longum subsp. longum		58.79	0.41	MNS 87			
Bifidobacterium adolescentis		33.37	0.39	CEL 1			
Actinomyces israelii		7.31	0.34				
43555512	**Very good identification**						**43555512**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		97.65	0.73	MNS 87			
Bifidobacterium longum subsp. longum		1.74	0.41				
Actinomyces israelii		0.32	0.37				
43555513	**Good identification**						**43555513**
Bifidobacterium adolescentis		98.25	0.39	MNS 87			
Bifidobacterium longum subsp. longum		1.75	0.07	URE 1			
43555520	**Low discrimination**						**43555520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		54.57	0.48	MNS 87	99	1	
Bacteroides vulgatus		24.71	0.28	CEL 1	1	3	
Capnocytophaga ochracea		17.05	0.31		1	90	
43555522	**Low discrimination**						**43555522**
					GRA	CAT	
Capnocytophaga ochracea		56.12	0.47	MNS 90	1	90	
Bifidobacterium longum subsp. longum		19.95	0.48		99	1	
Bacteroides fragilis		10.41	0.25		1	99	
43555530	**Good identification**						**43555530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.56	0.52	MNS 87	99	1	
Capnocytophaga ochracea		1.99	0.15	CEL 1	1	90	
Bacteroides vulgatus		0.26	-.07		1	3	
43555532	**Low discrimination**						**43555532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		69.19	0.52	MNS 87	99	1	
Capnocytophaga ochracea		12.70	0.31	CEL 1	1	90	
Bifidobacterium adolescentis		9.82	0.39		99	1	
43555540	**Low discrimination**						**43555540**
					GRA	CAT	
Actinomyces israelii		81.12	0.41	SAL 92	99	1	
Bifidobacterium adolescentis		14.40	0.22		99	1	
Capnocytophaga ochracea		2.27	-.02		1	90	
43555542	**Acceptable identification**						**43555542**
					GRA	CAT	
Bifidobacterium adolescentis		90.93	0.56	MNS 87	99	1	
Actinomyces israelii		7.76	0.44	SOR 11	99	1	
Capnocytophaga ochracea		1.31	0.15		1	90	
43555552	**Very good identification**						**43555552**
Bifidobacterium adolescentis		98.70	0.57	MNS 87			
Actinomyces israelii		1.03	0.30	SOR 11			
Bifidobacterium longum subsp. longum		0.14	0.07				
43555702	**Low discrimination**						**43555702**
Bifidobacterium adolescentis		60.48	0.37	MNS 87			
Clostridium clostridioforme		17.47	0.25	RHA 1			
Actinomyces israelii		12.23	0.32				
43555712	**Low discrimination**						**43555712**
Clostridium clostridioforme		50.33	0.36	MNS 99			
Bifidobacterium adolescentis		46.76	0.39	SAL 88			
Actinomyces israelii		1.16	0.18	TRE 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43555720	**Good identification**						**43555720**
					GRA		
Bacteroides vulgatus		98.97	0.44	MNS 99	1		
Eubacterium contortum		0.56	0.13	CEL 5	99		
Bifidobacterium longum subsp. longum		0.25	0.14		99		
43555722	**Low discrimination**						**43555722**
Parabacteroides distasonis		70.56	0.51	MNS 98			
Bacteroides uniformis		11.61	0.19				
Clostridium clostridioforme		4.56	0.25				
43555732	**Low discrimination**						**43555732**
					GRA		
Parabacteroides distasonis		60.57	0.46	MNS 98	1		
Clostridium clostridioforme		33.13	0.36		50		
Bifidobacterium longum subsp. longum		2.19	0.18		99		
43556502	**Low discrimination**						**43556502**
Bifidobacterium adolescentis		84.18	0.36	MNS 87			
Actinomyces israelii		15.82	0.30	GAL 99			
43556512	**Good identification**						**43556512**
Bifidobacterium adolescentis		97.75	0.37	MNS 87			
Actinomyces israelii		2.25	0.16	GAL 99			
43557000	**Good identification**						**43557000**
Actinomyces israelii		100.0	0.52	SAL 92			
43557002	**Good identification**						**43557002**
Actinomyces israelii		97.44	0.55	SAL 92			
Bifidobacterium adolescentis		2.56	0.22				
43557010	**Good identification**						**43557010**
Actinomyces israelii		100.0	0.38	SAL 92			
43557012	**Low discrimination**						**43557012**
Actinomyces israelii		82.30	0.41	SAL 92			
Bifidobacterium adolescentis		17.70	0.23	MLZ 13			
43557040	**Good identification**						**43557040**
Actinomyces israelii		100.0	0.45	SAL 92			
43557042	**Good identification**						**43557042**
Actinomyces israelii		100.0	0.48	SAL 92			
43557100	**Very good identification**						**43557100**
					GRA		
Actinomyces israelii		98.85	0.62	SAL 92	99		
Bifidobacterium adolescentis		0.98	0.22		99		
Fusobacterium mortiferum		0.18	0.13		1		
43557102	**Low discrimination**						**43557102**
					GRA	CAT	
Actinomyces israelii		60.35	0.65	SAL 92	99	1	
Bifidobacterium adolescentis		39.27	0.56		99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		0.29	0.10		1	90	
43557110	**Acceptable identification**						**43557110**
Actinomyces israelii		92.54	0.48	SAL 92			
Bifidobacterium adolescentis		7.46	0.23	MLZ 13			
43557112	**Low discrimination**						**43557112**
Bifidobacterium adolescentis		84.18	0.57	MNS 87			
Actinomyces israelii		15.82	0.51	ARA 87			
43557120	**Low discrimination**						**43557120**
					GRA	CAT	
Actinomyces israelii		75.73	0.42	SAL 92	99	1	
Bacteroides fragilis		13.29	0.04	NAG 6	1	99	
Capnocytophaga ochracea		8.66	0.10		1	90	
43557140	**Good identification**						**43557140**
Actinomyces israelii		100.0	0.55	SAL 92			
43557142	**Low discrimination**						**43557142**
Actinomyces israelii		82.86	0.58	SAL 92			
Bifidobacterium adolescentis		17.14	0.40				
43557150	**Good identification**						**43557150**
Actinomyces israelii		97.50	0.41	SAL 92			
Bifidobacterium adolescentis		2.50	0.08	MLZ 13			
43557152	**Low discrimination**						**43557152**
Bifidobacterium adolescentis		62.85	0.42	MNS 87			
Actinomyces israelii		37.16	0.44	ARA 87			
43557162	**Low discrimination**						**43557162**
					GRA	CAT	
Actinomyces israelii		79.77	0.38	SAL 92	99	1	
Capnocytophaga ochracea		15.63	0.10	NAG 6	1	90	
Bacteroides fragilis		4.61	-.09		1	99	
43557202	**Good identification**						**43557202**
Actinomyces israelii		100.0	0.36	SAL 92			
43557300	**Good identification**						**43557300**
Actinomyces israelii		100.0	0.43	SAL 92			
43557302	**Acceptable identification**						**43557302**
Actinomyces israelii		91.97	0.46	SAL 92			
Bifidobacterium adolescentis		8.03	0.22	RHA 7			
43557340	**Good identification**						**43557340**
Actinomyces israelii		100.0	0.36	SAL 92			
43557342	**Good identification**						**43557342**
Actinomyces israelii		100.0	0.39	SAL 92			
43557400	**Good identification**						**43557400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.51	SAL 92		
43557402	**Low discrimination**					**43557402**
Actinomyces israelii		82.30	0.54	SAL 92		
Bifidobacterium adolescentis		17.70	0.36			
43557410	**Good identification**					**43557410**
Actinomyces israelii		100.0	0.36	SAL 92		
43557412	**Low discrimination**					**43557412**
Bifidobacterium adolescentis		63.75	0.37	MNS 87		
Actinomyces israelii		36.25	0.39	LAC 99		
43557440	**Good identification**					**43557440**
Actinomyces israelii		100.0	0.44	SAL 92		
43557442	**Acceptable identification**					**43557442**
Actinomyces israelii		93.60	0.47	SAL 92		
Bifidobacterium adolescentis		6.40	0.20			
43557500	**Good identification**					**43557500**
					SPO	
Actinomyces israelii		92.33	0.61	SAL 92	1	
Bifidobacterium adolescentis		7.45	0.36		1	
Clostridium butyricum		0.22	-.05		99	
43557502	**Low discrimination**					**43557502**
Bifidobacterium adolescentis		84.18	0.70	MNS 87		
Actinomyces israelii		15.82	0.64			
43557503	**Low discrimination**					**43557503**
Bifidobacterium adolescentis		84.18	0.36	MNS 87		
Actinomyces israelii		15.82	0.30	URE 1		
43557510	**Low discrimination**					**43557510**
Actinomyces israelii		59.74	0.47	SAL 92		
Bifidobacterium adolescentis		39.40	0.37	MLZ 13		
Bifidobacterium longum subsp. longum		0.86	0.07			
43557512	**Very good identification**					**43557512**
Bifidobacterium adolescentis		97.70	0.72	MNS 87		
Actinomyces israelii		2.24	0.50			
Bifidobacterium dentium		0.03	-.07			
43557513	**Good identification**					**43557513**
Bifidobacterium adolescentis		97.75	0.37	MNS 87		
Actinomyces israelii		2.25	0.16	URE 1		
43557520	**Low discrimination**					**43557520**
					GRA	
Actinomyces israelii		89.72	0.40	SAL 92	99	
Bifidobacterium longum subsp. longum		7.08	0.14	NAG 6	99	
Bacteroides vulgatus		3.20	-.07		1	
43557522	**Low discrimination**					**43557522**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces israelii		48.31	0.43	SAL 92	99	1	
Bifidobacterium adolescentis		40.68	0.36	NAG 6	99	1	
Capnocytophaga ochracea		7.15	0.13		1	90	
43557532	**Low discrimination**						**43557532**
Bifidobacterium adolescentis		77.79	0.37	MNS 87			
Actinomyces israelii		11.29	0.29	NAG 1			
Bifidobacterium longum subsp. longum		6.77	0.18				
43557540	**Good identification**						**43557540**
Actinomyces israelii		97.50	0.54	SAL 92			
Bifidobacterium adolescentis		2.50	0.20				
43557542	**Low discrimination**						**43557542**
Bifidobacterium adolescentis		62.85	0.54	MNS 87			
Actinomyces israelii		37.16	0.57	SOR 11			
43557550	**Low discrimination**						**43557550**
					COC		
Actinomyces israelii		79.38	0.40	SAL 92	1		
Bifidobacterium adolescentis		16.64	0.22	MLZ 13	1		
Blautia producta		3.98	-.03		99		
43557552	**Acceptable identification**						**43557552**
Bifidobacterium adolescentis		93.26	0.56	MNS 87			
Actinomyces israelii		6.74	0.43	SOR 11			
43557562	**Low discrimination**						**43557562**
Actinomyces israelii		78.89	0.36	SAL 92			
Bifidobacterium adolescentis		21.12	0.20	NAG 6			
43557700	**Good identification**						**43557700**
Actinomyces israelii		100.0	0.42	SAL 92			
43557702	**Low discrimination**						**43557702**
Actinomyces israelii		58.34	0.45	SAL 92			
Bifidobacterium adolescentis		41.67	0.36	RHA 7			
43557712	**Low discrimination**						**43557712**
Bifidobacterium adolescentis		85.38	0.37	MNS 87			
Actinomyces israelii		14.62	0.31	RHA 1			
43557742	**Low discrimination**						**43557742**
Actinomyces israelii		81.50	0.38	SAL 92			
Bifidobacterium adolescentis		18.50	0.20	RHA 7			
43563100	**Low discrimination**						**43563100**
Actinomyces naeslundii		71.94	0.36	ESL 12			
Actinomyces israelii		28.06	0.20	MNS 88			
43563101	**Good identification**						**43563101**
Actinomyces naeslundii		100.0	0.39	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43563102	**Low discrimination**					**43563102**
Actinomyces naeslundii		63.09	0.36	ESL 12		
Actinomyces israelii		36.91	0.23	MNS 88		
43563103	**Good identification**					**43563103**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43563120	**Good identification**					**43563120**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43563121	**Good identification**					**43563121**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43563122	**Good identification**					**43563122**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43563123	**Good identification**					**43563123**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43565512	**Acceptable identification**					**43565512**
Bifidobacterium adolescentis		93.16	0.36	MNS 87		
Bifidobacterium longum subsp. longum		6.84	0.15	FRU 99		
43567100	**Good identification**					**43567100**
Actinomyces israelii		100.0	0.38	SAL 92		
43567102	**Low discrimination**					**43567102**
Actinomyces israelii		89.66	0.41	SAL 92		
Bifidobacterium adolescentis		10.34	0.19	FRU 99		
43567500	**Good identification**					**43567500**
Actinomyces israelii		100.0	0.37	SAL 92		
43567502	**Low discrimination**					**43567502**
Actinomyces israelii		51.46	0.40	SAL 92		
Bifidobacterium adolescentis		48.54	0.33	FRU 99		
43570000	**Good identification**					**43570000**
Fusobacterium mortiferum		100.0	0.48	MLT 11		
43570100	**Good identification**					**43570100**
					GRA	
Fusobacterium mortiferum		99.28	0.54	MLT 11	1	
Actinomyces naeslundii		0.72	0.29		99	
43570102	**Acceptable identification**					**43570102**
					GRA	
Fusobacterium mortiferum		93.87	0.38	MLT 11	1	
Actinomyces naeslundii		6.13	0.29	BGL 10	99	
43571000	**Low discrimination**					**43571000**
					GRA	
Fusobacterium mortiferum		85.40	0.51	MLT 11	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		8.13	0.45		99	
Actinomyces israelii		4.02	0.31		99	
43571001	**Acceptable identification**					**43571001**
					GRA	
Actinomyces naeslundii		93.39	0.48	ESL 12	99	
Fusobacterium mortiferum		6.61	0.17	MNS 88	1	
43571003	**Good identification**					**43571003**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43571020	**Low discrimination**					**43571020**
					GRA	
Actinomyces naeslundii		75.68	0.45	ESL 12	99	
Fusobacterium mortiferum		8.03	0.17	MNS 88	1	
Clostridium perfringens		6.57	0.05		99	
43571021	**Good identification**					**43571021**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43571023	**Good identification**					**43571023**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43571041	**Good identification**					**43571041**
Actinomyces naeslundii		100.0	0.35	ESL 12		
43571043	**Good identification**					**43571043**
Actinomyces naeslundii		100.0	0.35	ESL 12		
43571061	**Good identification**					**43571061**
Actinomyces naeslundii		100.0	0.35	ESL 12		
43571063	**Good identification**					**43571063**
Actinomyces naeslundii		100.0	0.35	ESL 12		
43571100	**Low discrimination**					**43571100**
					GRA	
Fusobacterium mortiferum		78.17	0.57	MLT 11	1	
Actinomyces naeslundii		9.07	0.53		99	
Actinomyces israelii		6.32	0.41		99	
43571101	**Acceptable identification**					**43571101**
					GRA	
Actinomyces naeslundii		94.52	0.56	ESL 12	99	
Fusobacterium mortiferum		5.49	0.23	MNS 88	1	
43571102	**Low discrimination**					**43571102**
Bifidobacterium adolescentis		25.93	0.40	MNS 87		
Actinomyces israelii		18.89	0.44	XYL 87		
Actinomyces naeslundii		18.10	0.53	ARA 87		
43571103	**Good identification**					**43571103**
Actinomyces naeslundii		100.0	0.56	ESL 12		
43571121	**Good identification**					**43571121**
					GRA	
Actinomyces naeslundii		98.45	0.56	ESL 12	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		1.55	-.04	MNS 88	1		
43571122	**Low discrimination**						**43571122**
					CAT		
Prevotella oralis		48.93	0.41	MNS 99	1		
Capnocytophaga ochracea		24.29	0.36	TRE 6	90		
Parabacteroides distasonis		9.17	0.41		76		
43571123	**Acceptable identification**						**43571123**
					GRA	CAT	
Actinomyces naeslundii		93.96	0.56	ESL 12	99	25	
Prevotella oralis		4.04	0.07	MNS 88	1	1	
Capnocytophaga ochracea		2.00	0.02		1	90	
43571132	**Low discrimination**						**43571132**
					GRA	CAT	
Parabacteroides distasonis		40.07	0.37	MNS 98	1	76	
Capnocytophaga ochracea		21.90	0.20	XYL 90	1	90	
Bifidobacterium breve		16.64	0.29		99	1	
43571141	**Good identification**						**43571141**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43571142	**Low discrimination**						**43571142**
Actinomyces israelii		42.78	0.37	SAL 92			
Bifidobacterium adolescentis		18.66	0.25	XYL 92			
Actinomyces naeslundii		18.59	0.40				
43571143	**Good identification**						**43571143**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43571160	**Low discrimination**						**43571160**
					GRA	CAT	
Actinomyces naeslundii		54.21	0.40	ESL 12	99	25	
Clostridium perfringens		13.08	0.07	MNS 88	99	1	
Capnocytophaga ochracea		12.01	0.04		1	90	
43571161	**Good identification**						**43571161**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43571163	**Good identification**						**43571163**
Actinomyces naeslundii		100.0	0.43	ESL 12			
43571322	**Low discrimination**						**43571322**
					SPO		
Parabacteroides distasonis		53.91	0.50	MNS 98	1		
Prevotella oralis		41.10	0.35	XYL 90	1		
Clostridium ramosum		4.51	0.19		92		
43571332	**Good identification**						**43571332**
Parabacteroides distasonis		95.70	0.45	MNS 98			
Clostridium clostridioforme		2.93	0.13	XYL 90			
Prevotella oralis		1.37	0.01				
43571500	**Low discrimination**						**43571500**
					SPO		
Actinomyces israelii		59.42	0.40	SAL 92	1		
Clostridium butyricum		10.69	0.06	XYL 92	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		10.69	0.19		1		
43571502	**Low discrimination**						**43571502**
Bifidobacterium adolescentis		89.82	0.54	MNS 87			
Actinomyces israelii		8.00	0.43	XYL 87			
Bifidobacterium longum subsp. longum		0.96	0.19				
43571512	**Good identification**						**43571512**
Bifidobacterium adolescentis		96.32	0.56	MNS 87			
Bifidobacterium longum subsp. longum		1.43	0.23	XYL 87			
Actinomyces israelii		1.05	0.29				
43571522	**Low discrimination**						**43571522**
					GRA	CAT	
Parabacteroides distasonis		37.67	0.38	MNS 98	1	76	
Capnocytophaga ochracea		28.74	0.23	XYL 90	1	90	
Bifidobacterium longum subsp. longum		21.07	0.29		99	1	
43571542	**Low discrimination**						**43571542**
Bifidobacterium adolescentis		78.11	0.39	MNS 87			
Actinomyces israelii		21.89	0.36	XYL 87			
43571552	**Good identification**						**43571552**
Bifidobacterium adolescentis		96.69	0.40	MNS 87			
Actinomyces israelii		3.31	0.22	XYL 87			
43571722	**Low discrimination**						**43571722**
Parabacteroides distasonis		86.33	0.46	MNS 98			
Clostridium clostridioforme		8.00	0.23	XYL 90			
Bacteroides thetaiotaomicron		4.58	0.08				
43571732	**Low discrimination**						**43571732**
Parabacteroides distasonis		54.88	0.41	MNS 98			
Clostridium clostridioforme		43.02	0.34	XYL 90			
Bacteroides thetaiotaomicron		2.10	0.01				
43572101	**Good identification**						**43572101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43572103	**Good identification**						**43572103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43572121	**Good identification**						**43572121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43572123	**Good identification**						**43572123**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43573000	**Low discrimination**						**43573000**
					GRA		
Actinomyces israelii		57.82	0.44	SAL 92	99		
Actinomyces naeslundii		39.99	0.50	XYL 92	99		
Fusobacterium mortiferum		2.19	0.17		1		
43573001	**Good identification**						**43573001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	ESL 12		
43573002	**Low discrimination**					**43573002**
Actinomyces israelii		68.44	0.47	SAL 92		
Actinomyces naeslundii		31.56	0.50	XYL 92		
43573003	**Good identification**					**43573003**
Actinomyces naeslundii		100.0	0.53	ESL 12		
43573020	**Acceptable identification**					**43573020**
Actinomyces naeslundii		91.55	0.50	ESL 12		
Actinomyces israelii		8.45	0.23	MNS 88		
43573021	**Good identification**					**43573021**
Actinomyces naeslundii		100.0	0.53	ESL 12		
43573022	**Low discrimination**					**43573022**
Actinomyces naeslundii		87.84	0.50	ESL 12		
Actinomyces israelii		12.16	0.26	MNS 88		
43573023	**Good identification**					**43573023**
Actinomyces naeslundii		100.0	0.53	ESL 12		
43573040	**Low discrimination**					**43573040**
Actinomyces israelii		76.11	0.37	SAL 92		
Actinomyces naeslundii		23.89	0.37	XYL 92		
43573041	**Good identification**					**43573041**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43573042	**Low discrimination**					**43573042**
Actinomyces israelii		82.70	0.40	SAL 92		
Actinomyces naeslundii		17.30	0.37	XYL 92		
43573043	**Good identification**					**43573043**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43573060	**Low discrimination**					**43573060**
Actinomyces naeslundii		83.10	0.37	ESL 12		
Actinomyces israelii		16.90	0.16	MNS 88		
43573061	**Good identification**					**43573061**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43573062	**Low discrimination**					**43573062**
Actinomyces naeslundii		76.62	0.37	ESL 12		
Actinomyces israelii		23.38	0.19	MNS 88		
43573063	**Good identification**					**43573063**
Actinomyces naeslundii		100.0	0.40	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43573100	**Low discrimination**					**43573100**
					GRA	
Actinomyces israelii		66.04	0.54	SAL 92	99	
Actinomyces naeslundii		32.50	0.58	XYL 92	99	
Fusobacterium mortiferum		1.46	0.23		1	
43573101	**Very good identification**					**43573101**
Actinomyces naeslundii		98.65	0.61	ESL 12		
Actinomyces israelii		1.35	0.20	MNS 88		
43573102	**Low discrimination**					**43573102**
Actinomyces israelii		64.31	0.57	SAL 92		
Actinomyces naeslundii		21.10	0.58	XYL 92		
Bifidobacterium adolescentis		12.74	0.39			
43573103	**Very good identification**					**43573103**
Actinomyces naeslundii		97.99	0.61	ESL 12		
Actinomyces israelii		2.01	0.23	MNS 88		
43573110	**Low discrimination**					**43573110**
Actinomyces israelii		85.23	0.40	SAL 92		
Actinomyces naeslundii		14.77	0.36	XYL 92		
43573111	**Good identification**					**43573111**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43573112	**Low discrimination**					**43573112**
Bifidobacterium adolescentis		55.36	0.40	MNS 87		
Actinomyces israelii		34.16	0.43	XYL 87		
Bifidobacterium breve		5.81	0.24	ARA 87		
43573113	**Good identification**					**43573113**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43573120	**Low discrimination**					**43573120**
					SPO	
Actinomyces naeslundii		75.72	0.58	ESL 12	1	
Clostridium ramosum		12.50	0.19	MNS 88	92	
Actinomyces israelii		9.82	0.34		1	
43573121	**Very good identification**					**43573121**
Actinomyces naeslundii		100.0	0.61	ESL 12		
43573122	**Low discrimination**					**43573122**
					SPO	
Actinomyces naeslundii		48.96	0.58	ESL 12	1	
Clostridium ramosum		32.34	0.30	MNS 88	92	
Actinomyces israelii		9.53	0.37		1	
43573123	**Very good identification**					**43573123**
Actinomyces naeslundii		100.0	0.61	ESL 12		
43573130	**Low discrimination**					**43573130**
Actinomyces naeslundii		61.46	0.36	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		22.63	0.19	MNS 88		
Bifidobacterium breve		15.91	0.12	MLZ 5		
43573131	**Good identification**					**43573131**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43573133	**Good identification**					**43573133**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43573140	**Low discrimination**					**43573140**
Actinomyces israelii		81.75	0.47	SAL 92		
Actinomyces naeslundii		18.26	0.45	XYL 92		
43573141	**Good identification**					**43573141**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43573142	**Low discrimination**					**43573142**
Actinomyces israelii		77.77	0.50	SAL 92		
Actinomyces naeslundii		11.58	0.45	XYL 92		
Bifidobacterium breve		5.08	0.24			
43573143	**Good identification**					**43573143**
Actinomyces naeslundii		95.67	0.48	ESL 12		
Actinomyces israelii		4.33	0.16	MNS 88		
43573152	**Low discrimination**					**43573152**
Actinomyces israelii		49.88	0.36	SAL 92		
Bifidobacterium adolescentis		25.69	0.25	XYL 92		
Bifidobacterium breve		21.82	0.24	MLZ 13		
43573160	**Low discrimination**					**43573160**
Actinomyces naeslundii		73.37	0.45	ESL 12		
Actinomyces israelii		20.97	0.27	MNS 88		
Bifidobacterium breve		5.66	0.12			
43573161	**Good identification**					**43573161**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43573162	**Low discrimination**					**43573162**
Actinomyces naeslundii		45.27	0.45	ESL 12		
Bifidobacterium breve		35.33	0.29	MNS 88		
Actinomyces israelii		19.41	0.30			
43573163	**Good identification**					**43573163**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43573302	**Acceptable identification**					**43573302**
					SPO	
Actinomyces israelii		92.37	0.38	SAL 92	1	
Actinomyces naeslundii		4.07	0.24	XYL 92	1	
Clostridium ramosum		3.56	-.03	RHA 7	92	
43573400	**Good identification**					**43573400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.42	SAL 92		
43573402	**Acceptable identification**					**43573402**
Actinomyces israelii		93.85	0.45	SAL 92		
Bifidobacterium adolescentis		6.15	0.19	XYL 92		
43573440	**Good identification**					**43573440**
Actinomyces israelii		100.0	0.35	SAL 92		
43573442	**Good identification**					**43573442**
Actinomyces israelii		100.0	0.38	SAL 92		
43573500	**Good identification**					**43573500**
Actinomyces israelii		97.03	0.53	SAL 92		
Bifidobacterium adolescentis		2.38	0.19	XYL 92		
Actinomyces naeslundii		0.59	0.24			
43573502	**Low discrimination**					**43573502**
Bifidobacterium adolescentis		61.75	0.53	MNS 87		
Actinomyces israelii		38.10	0.56	XYL 87		
Actinomyces naeslundii		0.15	0.24			
43573510	**Low discrimination**					**43573510**
Actinomyces israelii		83.27	0.38	SAL 92		
Bifidobacterium adolescentis		16.73	0.20	XYL 92		
43573512	**Acceptable identification**					**43573512**
Bifidobacterium adolescentis		92.72	0.54	MNS 87		
Actinomyces israelii		7.00	0.41	XYL 87		
Bifidobacterium dentium		0.28	-.07			
43573522	**Low discrimination**					**43573522**
Actinomyces israelii		75.76	0.35	SAL 92		
Bifidobacterium adolescentis		19.43	0.19	XYL 92		
Actinomyces naeslundii		4.81	0.24	NAG 6		
43573540	**Good identification**					**43573540**
Actinomyces israelii		100.0	0.46	SAL 92		
43573542	**Low discrimination**					**43573542**
Actinomyces israelii		66.00	0.49	SAL 92		
Bifidobacterium adolescentis		34.00	0.37	XYL 92		
43573552	**Low discrimination**					**43573552**
Bifidobacterium adolescentis		80.82	0.39	MNS 87		
Actinomyces israelii		19.18	0.34	XYL 87		
43573702	**Low discrimination**					**43573702**
Actinomyces israelii		82.14	0.36	SAL 92		
Bifidobacterium adolescentis		17.86	0.19	XYL 92		
43574100	**Low discrimination**					**43574100**

GRA

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		84.25	0.35	MLT 11	1		
Actinomyces israelii		15.76	0.25	XYL 7	99		
43574512	**Low discrimination**						**43574512**
Bifidobacterium adolescentis		65.01	0.36	MNS 87			
Clostridium clostridioforme		34.99	0.28	GAL 99			
43574712	**Good identification**						**43574712**
Clostridium clostridioforme		100.0	0.43	MNS 99			
43574732	**Good identification**						**43574732**
Clostridium clostridioforme		96.67	0.43	MNS 99			
Parabacteroides distasonis		3.33	0.24	SAL 88			
43575000	**Low discrimination**						**43575000**
					GRA		
Actinomyces israelii		86.74	0.49	SAL 92	99		
Fusobacterium mortiferum		12.05	0.32		1		
Eubacterium saburreum		1.21	0.13		85		
43575002	**Good identification**						**43575002**
					GRA		
Actinomyces israelii		95.92	0.52	SAL 92	99		
Bifidobacterium adolescentis		3.10	0.20		99		
Fusobacterium mortiferum		0.99	0.16		1		
43575012	**Low discrimination**						**43575012**
Actinomyces israelii		75.86	0.38	SAL 92			
Bifidobacterium adolescentis		20.02	0.22	MLZ 13			
Clostridium clostridioforme		4.12	0.07				
43575040	**Good identification**						**43575040**
Actinomyces israelii		100.0	0.42	SAL 92			
43575042	**Good identification**						**43575042**
Actinomyces israelii		100.0	0.45	SAL 92			
43575100	**Low discrimination**						**43575100**
					COC	GRA	
Actinomyces israelii		79.04	0.59	SAL 92	1	99	
Atopobium parvulum		9.32	0.30		99	99	
Fusobacterium mortiferum		6.40	0.38		1	1	
43575102	**Low discrimination**						**43575102**
					COC		
Actinomyces israelii		53.94	0.62	SAL 92	1		
Bifidobacterium adolescentis		43.08	0.54		1		
Atopobium parvulum		0.75	0.17		99		
43575110	**Low discrimination**						**43575110**
Actinomyces israelii		86.23	0.45	SAL 92			
Bifidobacterium adolescentis		8.53	0.22	MLZ 13			
Eubacterium saburreum		3.84	0.18				
43575112	**Low discrimination**						**43575112**
Bifidobacterium adolescentis		85.71	0.56	MNS 87			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		13.12	0.48	ARA 87				
Clostridium clostridioforme		0.60	0.16					
43575120	**Low discrimination**							**43575120**
					CAT	COC	GRA	
Actinomyces israelii		46.18	0.39	SAL 92	1	1	99	
Atopobium parvulum		15.06	0.17	NAG 6	1	99	99	
Bacteroides fragilis		12.21	0.04		99	1	1	
43575122	**Low discrimination**							**43575122**
Parabacteroides distasonis		68.03	0.58	MNS 98				
Bacteroides fragilis		16.23	0.25					
Capnocytophaga ochracea		5.01	0.26					
43575132	**Acceptable identification**							**43575132**
					GRA			
Parabacteroides distasonis		92.80	0.53	MNS 98	1			
Bifidobacterium adolescentis		1.90	0.22		99			
Actinomyces israelii		1.83	0.28		99			
43575140	**Good identification**							**43575140**
					SPO			
Actinomyces israelii		98.98	0.52	SAL 92	1			
Clostridium butyricum		1.02	-.03		99			
43575142	**Low discrimination**							**43575142**
Actinomyces israelii		79.76	0.55	SAL 92				
Bifidobacterium adolescentis		20.24	0.39					
43575150	**Good identification**							**43575150**
Actinomyces israelii		100.0	0.38	SAL 92				
43575152	**Low discrimination**							**43575152**
Bifidobacterium adolescentis		67.49	0.40	MNS 87				
Actinomyces israelii		32.51	0.41	ARA 87				
43575300	**Good identification**							**43575300**
Actinomyces israelii		100.0	0.40	SAL 92				
43575302	**Low discrimination**							**43575302**
					GRA			
Actinomyces israelii		61.24	0.43	SAL 92	99			
Parabacteroides distasonis		22.09	0.32	RHA 7	1			
Clostridium clostridioforme		8.91	0.19		50			
43575322	**Very good identification**							**43575322**
Parabacteroides distasonis		98.12	0.66	MNS 98				
Prevotella oralis		1.03	0.19					
Clostridium clostridioforme		0.40	0.19					
43575332	**Very good identification**							**43575332**
Parabacteroides distasonis		96.54	0.61	MNS 98				
Clostridium clostridioforme		3.33	0.31					
Bacteroides ovatus		0.13	-.04					
43575342	**Good identification**							**43575342**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	SAL 92		
43575400	**Acceptable identification**					**43575400**
					SPO	
Actinomyces israelii		92.22	0.48	SAL 92	1	
Clostridium butyricum		7.78	0.08		99	
43575402	**Low discrimination**					**43575402**
Actinomyces israelii		75.76	0.51	SAL 92		
Bifidobacterium adolescentis		20.00	0.34			
Clostridium clostridioforme		2.59	0.16			
43575412	**Low discrimination**					**43575412**
Bifidobacterium adolescentis		51.40	0.36	MNS 87		
Clostridium clostridioforme		24.79	0.27	LAC 99		
Actinomyces israelii		23.81	0.36			
43575440	**Good identification**					**43575440**
Actinomyces israelii		100.0	0.41	SAL 92		
43575442	**Acceptable identification**					**43575442**
Actinomyces israelii		92.26	0.44	SAL 92		
Bifidobacterium adolescentis		7.74	0.19			
43575500	**Low discrimination**					**43575500**
					SPO	
Actinomyces israelii		60.54	0.58	SAL 92	1	
Clostridium butyricum		30.63	0.32		99	
Bifidobacterium adolescentis		5.99	0.34		1	
43575502	**Low discrimination**					**43575502**
					SPO	
Bifidobacterium adolescentis		84.50	0.69	MNS 87	1	
Actinomyces israelii		12.94	0.61		1	
Clostridium butyricum		1.70	0.25		99	
43575510	**Low discrimination**					**43575510**
Actinomyces israelii		39.64	0.44	SAL 92		
Bifidobacterium adolescentis		32.09	0.36	MLZ 13		
Bifidobacterium longum subsp. longum		21.20	0.31			
43575512	**Very good identification**					**43575512**
Bifidobacterium adolescentis		95.84	0.70	MNS 87		
Actinomyces israelii		1.79	0.47			
Clostridium clostridioforme		1.56	0.36			
43575513	**Good identification**					**43575513**
Bifidobacterium adolescentis		100.0	0.36	MNS 87		
43575520	**Low discrimination**					**43575520**
					SPO	
Bifidobacterium longum subsp. longum		58.69	0.37	MNS 87	1	
Actinomyces israelii		19.95	0.37	CEL 1	1	
Clostridium butyricum		13.75	0.14		99	
43575522	**Low discrimination**					**43575522**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		66.41	0.54	MNS 98	1	
Bifidobacterium longum subsp. longum		12.38	0.37		99	
Bifidobacterium adolescentis		6.53	0.34		99	
43575530	**Good identification**					**43575530**
					GRA	
Bifidobacterium longum subsp. longum		95.51	0.41	MNS 87	99	
Actinomyces israelii		2.85	0.23	CEL 1	99	
Parabacteroides distasonis		1.64	0.15		1	
43575532	**Low discrimination**					**43575532**
					GRA	
Parabacteroides distasonis		44.47	0.50	MNS 98	1	
Bifidobacterium longum subsp. longum		26.22	0.41		99	
Clostridium clostridioforme		16.02	0.36		50	
43575540	**Low discrimination**					**43575540**
					SPO	
Actinomyces israelii		87.37	0.51	SAL 92	1	
Clostridium butyricum		9.89	0.14		99	
Bifidobacterium adolescentis		2.75	0.19		1	
43575542	**Low discrimination**					**43575542**
					SPO	
Bifidobacterium adolescentis		66.85	0.53	MNS 87	1	
Actinomyces israelii		32.20	0.54	SOR 11	1	
Clostridium butyricum		0.95	0.07		99	
43575550	**Low discrimination**					**43575550**
					COC	
Actinomyces israelii		75.03	0.37	SAL 92	1	
Bifidobacterium adolescentis		19.30	0.20	MLZ 13	1	
Blautia producta		5.67	-.03		99	
43575552	**Acceptable identification**					**43575552**
Bifidobacterium adolescentis		94.44	0.54	MNS 87		
Actinomyces israelii		5.56	0.40	SOR 11		
43575612	**Good identification**					**43575612**
Clostridium clostridioforme		99.02	0.42	MNS 99		
Actinomyces israelii		0.98	0.17	SAL 88		
43575632	**Good identification**					**43575632**
Clostridium clostridioforme		95.57	0.42	MNS 99		
Parabacteroides distasonis		3.67	0.24	SAL 88		
Bacteroides thetaiotaomicron		0.76	-.04			
43575700	**Low discrimination**					**43575700**
					SPO	
Actinomyces israelii		79.99	0.39	SAL 92	1	
Clostridium butyricum		16.63	0.06	RHA 7	99	
Clostridium clostridioforme		3.38	0.06		66	
43575702	**Low discrimination**					**43575702**
Clostridium clostridioforme		55.43	0.40	MNS 99		
Actinomyces israelii		19.91	0.42	SAL 88		
Bifidobacterium adolescentis		17.45	0.34			
43575712	**Acceptable identification**					**43575712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		90.28	0.51	MNS 99		
Bifidobacterium adolescentis		7.63	0.36	SAL 88		
Actinomyces israelii		1.06	0.27			
43575722	**Low discrimination**					**43575722**
Parabacteroides distasonis		76.54	0.62	MNS 98		
Bacteroides thetaiotaomicron		10.83	0.32			
Clostridium clostridioforme		7.97	0.40			
43575732	**Low discrimination**					**43575732**
Parabacteroides distasonis		48.61	0.58	MNS 98		
Clostridium clostridioforme		42.80	0.51			
Bacteroides thetaiotaomicron		4.97	0.25			
43576100	**Good identification**					**43576100**
Actinomyces israelii		100.0	0.38	SAL 92		
43576102	**Low discrimination**					**43576102**
Actinomyces israelii		89.66	0.41	SAL 92		
Bifidobacterium adolescentis		10.34	0.19	GAL 99		
43576500	**Good identification**					**43576500**
Actinomyces israelii		100.0	0.37	SAL 92		
43576502	**Low discrimination**					**43576502**
Actinomyces israelii		51.46	0.40	SAL 92		
Bifidobacterium adolescentis		48.54	0.33	GAL 99		
43577000	**Very good identification**					**43577000**
Actinomyces israelii		100.0	0.62	SAL 92		
43577002	**Very good identification**					**43577002**
Actinomyces israelii		99.54	0.65	SAL 92		
Bifidobacterium adolescentis		0.46	0.19			
43577010	**Good identification**					**43577010**
Actinomyces israelii		100.0	0.48	SAL 92		
43577012	**Good identification**					**43577012**
Actinomyces israelii		96.33	0.51	SAL 92		
Bifidobacterium adolescentis		3.67	0.20	MLZ 13		
43577020	**Good identification**					**43577020**
Actinomyces israelii		100.0	0.41	SAL 92		
43577022	**Good identification**					**43577022**
Actinomyces israelii		100.0	0.44	SAL 92		
43577040	**Good identification**					**43577040**
Actinomyces israelii		100.0	0.55	SAL 92		
43577042	**Very good identification**					**43577042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.58	SAL 92		
43577050	**Good identification**					**43577050**
Actinomyces israelii		100.0	0.41	SAL 92		
43577052	**Good identification**					**43577052**
Actinomyces israelii		100.0	0.44	SAL 92		
43577062	**Good identification**					**43577062**
Actinomyces israelii		100.0	0.37	SAL 92		
43577100	**Very good identification**					**43577100**
					SPO	
Actinomyces israelii		99.69	0.72	SAL 92	1	
Bifidobacterium adolescentis		0.17	0.19		1	
Clostridium butyricum		0.09	-.01		99	
43577101	**Acceptable identification**					**43577101**
Actinomyces israelii		93.97	0.38	SAL 92		
Actinomyces naeslundii		6.03	0.27	URE 1		
43577102	**Low discrimination**					**43577102**
Actinomyces israelii		89.64	0.75	SAL 92		
Bifidobacterium adolescentis		10.34	0.53			
Actinomyces naeslundii		0.03	0.24			
43577103	**Low discrimination**					**43577103**
Actinomyces israelii		86.35	0.41	SAL 92		
Bifidobacterium adolescentis		9.96	0.19	URE 1		
Actinomyces naeslundii		3.70	0.27			
43577110	**Very good identification**					**43577110**
Actinomyces israelii		98.59	0.58	SAL 92		
Bifidobacterium adolescentis		1.41	0.20	MLZ 13		
43577112	**Low discrimination**					**43577112**
Actinomyces israelii		51.39	0.61	SAL 92		
Bifidobacterium adolescentis		48.47	0.54	MLZ 13		
Bifidobacterium dentium		0.15	-.07			
43577120	**Good identification**					**43577120**
Actinomyces israelii		99.33	0.52	SAL 92		
Actinomyces naeslundii		0.67	0.24	NAG 6		
43577122	**Acceptable identification**					**43577122**
					GRA	
Actinomyces israelii		94.90	0.55	SAL 92	99	
Parabacteroides distasonis		2.38	0.24	NAG 6	1	
Bifidobacterium adolescentis		1.73	0.19		99	
43577130	**Good identification**					**43577130**
Actinomyces israelii		100.0	0.38	SAL 92		
43577132	**Low discrimination**					**43577132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces israelii		75.79	0.41	SAL 92	99	
Bifidobacterium adolescentis		11.31	0.20	MLZ 13	99	
Parabacteroides distasonis		6.84	0.19	NAG 6	1	
43577140	**Very good identification**					**43577140**
Actinomyces israelii		100.0	0.65	SAL 92		
43577142	**Very good identification**					**43577142**
Actinomyces israelii		96.47	0.68	SAL 92		
Bifidobacterium adolescentis		3.54	0.37			
43577150	**Good identification**					**43577150**
					COC	
Actinomyces israelii		98.92	0.51	SAL 92	1	
Blautia producta		1.08	-.03	MLZ 13	99	
43577152	**Low discrimination**					**43577152**
Actinomyces israelii		76.94	0.54	SAL 92		
Bifidobacterium adolescentis		23.06	0.39	MLZ 13		
43577160	**Good identification**					**43577160**
Actinomyces israelii		100.0	0.45	SAL 92		
43577162	**Good identification**					**43577162**
Actinomyces israelii		100.0	0.48	SAL 92		
43577200	**Good identification**					**43577200**
Actinomyces israelii		100.0	0.43	SAL 92		
43577202	**Good identification**					**43577202**
Actinomyces israelii		100.0	0.46	SAL 92		
43577240	**Good identification**					**43577240**
Actinomyces israelii		100.0	0.36	SAL 92		
43577242	**Good identification**					**43577242**
Actinomyces israelii		100.0	0.39	SAL 92		
43577300	**Good identification**					**43577300**
Actinomyces israelii		100.0	0.53	SAL 92		
43577302	**Good identification**					**43577302**
Actinomyces israelii		98.48	0.56	SAL 92		
Bifidobacterium adolescentis		1.52	0.19	RHA 7		
43577310	**Good identification**					**43577310**
Actinomyces israelii		100.0	0.39	SAL 92		
43577312	**Low discrimination**					**43577312**
Actinomyces israelii		88.76	0.42	SAL 92		
Bifidobacterium adolescentis		11.24	0.20	RHA 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43577322	**Low discrimination**						**43577322**
					SPO	GRA	
Actinomyces israelii		48.78	0.36	SAL 92	1	99	
Parabacteroides distasonis		48.66	0.32	RHA 7	1	1	
Clostridium ramosum		2.56	-.03	NAG 6	92	50	
43577340	**Good identification**						**43577340**
Actinomyces israelii		100.0	0.46	SAL 92			
43577342	**Good identification**						**43577342**
Actinomyces israelii		100.0	0.49	SAL 92			
43577400	**Very good identification**						**43577400**
Actinomyces israelii		100.0	0.60	SAL 92			
43577402	**Very good identification**						**43577402**
Actinomyces israelii		96.33	0.63	SAL 92			
Bifidobacterium adolescentis		3.67	0.33				
43577410	**Good identification**						**43577410**
Actinomyces israelii		100.0	0.46	SAL 92			
43577412	**Low discrimination**						**43577412**
Actinomyces israelii		75.43	0.49	SAL 92			
Bifidobacterium adolescentis		23.51	0.34	MLZ 13			
Bifidobacterium dentium		1.06	-.07				
43577420	**Good identification**						**43577420**
Actinomyces israelii		100.0	0.40	SAL 92			
43577422	**Good identification**						**43577422**
Actinomyces israelii		100.0	0.43	SAL 92			
43577440	**Good identification**						**43577440**
Actinomyces israelii		100.0	0.53	SAL 92			
43577442	**Very good identification**						**43577442**
Actinomyces israelii		98.80	0.56	SAL 92			
Bifidobacterium adolescentis		1.20	0.17				
43577450	**Acceptable identification**						**43577450**
					COC		
Actinomyces israelii		94.94	0.39	SAL 92	1		
Blautia producta		5.07	-.03	MLZ 13	99		
43577452	**Acceptable identification**						**43577452**
Actinomyces israelii		90.99	0.42	SAL 92			
Bifidobacterium adolescentis		9.01	0.19	MLZ 13			
43577462	**Good identification**						**43577462**
Actinomyces israelii		100.0	0.36	SAL 92			
43577500	**Very good identification**						**43577500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		97.64	0.71	SAL 92	1	
Bifidobacterium adolescentis		1.40	0.33		1	
Clostridium butyricum		0.97	0.15		99	
43577501	**Good identification**					**43577501**
Actinomyces israelii		100.0	0.37	SAL 92		
43577502	**Low discrimination**					**43577502**
					SPO	
Actinomyces israelii		51.38	0.74	SAL 92	1	
Bifidobacterium adolescentis		48.46	0.67		1	
Clostridium butyricum		0.13	0.08		99	
43577503	**Low discrimination**					**43577503**
Actinomyces israelii		51.46	0.40	SAL 92		
Bifidobacterium adolescentis		48.54	0.33	URE 1		
43577510	**Low discrimination**					**43577510**
Actinomyces israelii		84.96	0.57	SAL 92		
Bifidobacterium adolescentis		9.93	0.34	MLZ 13		
Bifidobacterium dentium		4.38	0.10			
43577512	**Low discrimination**					**43577512**
Bifidobacterium adolescentis		85.13	0.69	MNS 87		
Actinomyces israelii		11.04	0.60			
Bifidobacterium dentium		3.83	0.27			
43577520	**Good identification**					**43577520**
					SPO	
Actinomyces israelii		98.67	0.50	SAL 92	1	
Clostridium butyricum		1.33	-.03	NAG 6	99	
43577522	**Low discrimination**					**43577522**
					GRA	
Actinomyces israelii		84.15	0.53	SAL 92	99	
Bifidobacterium adolescentis		12.56	0.33	NAG 6	99	
Parabacteroides distasonis		1.58	0.20		1	
43577540	**Good identification**					**43577540**
					COC	
Actinomyces israelii		93.24	0.64	SAL 92	1	
Blautia producta		6.13	0.24		99	
Bifidobacterium adolescentis		0.42	0.17		1	
43577542	**Low discrimination**					**43577542**
					COC	
Actinomyces israelii		76.65	0.67	SAL 92	1	
Bifidobacterium adolescentis		22.98	0.51		1	
Blautia producta		0.37	0.07		99	
43577552	**Low discrimination**					**43577552**
					COC	
Bifidobacterium adolescentis		68.90	0.53	MNS 87	1	
Actinomyces israelii		28.10	0.53	SOR 11	1	
Blautia producta		2.75	0.15		99	
43577560	**Low discrimination**					**43577560**
					COC	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		89.73	0.43	SAL 92	1		
Blautia producta		10.27	0.07	NAG 6	99		
43577562	**Good identification**						**43577562**
Actinomyces israelii		95.47	0.46	SAL 92			
Bifidobacterium adolescentis		4.53	0.17	NAG 6			
43577600	**Good identification**						**43577600**
Actinomyces israelii		100.0	0.41	SAL 92			
43577602	**Good identification**						**43577602**
Actinomyces israelii		100.0	0.44	SAL 92			
43577642	**Good identification**						**43577642**
Actinomyces israelii		100.0	0.37	SAL 92			
43577700	**Good identification**						**43577700**
Actinomyces israelii		100.0	0.52	SAL 92			
43577702	**Low discrimination**						**43577702**
Actinomyces israelii		88.38	0.55	SAL 92			
Bifidobacterium adolescentis		11.19	0.33	RHA 7			
Clostridium clostridioforme		0.44	0.06				
43577710	**Acceptable identification**						**43577710**
					COC		
Actinomyces israelii		93.55	0.37	SAL 92	1		
Blautia producta		6.45	-.03	RHA 7	99		
43577740	**Low discrimination**						**43577740**
					COC		
Actinomyces israelii		53.36	0.44	SAL 92	1		
Blautia producta		46.64	0.24	RHA 7	99		
43577742	**Acceptable identification**						**43577742**
					COC		
Actinomyces israelii		90.50	0.47	SAL 92	1		
Blautia producta		5.86	0.07	RHA 7	99		
Bifidobacterium adolescentis		3.64	0.17		1		
43600000	**Very good identification**						**43600000**
					GRA	SPO	
Eubacterium tortuosum		95.40	0.58	GAL 99	99	1	
Fusobacterium mortiferum		4.46	0.52		1	1	
Clostridium botulinum biovar A		0.14	0.13		99	99	
43600002	**Good identification**						**43600002**
					GRA		
Eubacterium tortuosum		97.14	0.45	GAL 99	99		
Fusobacterium mortiferum		2.86	0.35		1		
43600020	**Good identification**						**43600020**
					GRA		
Eubacterium tortuosum		99.73	0.45	GAL 99	99		
Fusobacterium mortiferum		0.27	0.17		1		
43600100	**Good identification**						**43600100**
					GRA		
Fusobacterium mortiferum		91.53	0.58	FRU 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		8.47	0.24		99	
43600102	**Low discrimination**					**43600102**
					GRA	
Fusobacterium mortiferum		87.19	0.42	FRU 99	1	
Eubacterium tortuosum		12.81	0.11	BGL 10	99	
43601000	**Excellent identification**					**43601000**
					GRA	
Eubacterium tortuosum		99.93	0.92		99	
Fusobacterium mortiferum		0.07	0.55		1	
43601001	**Very good identification**					**43601001**
					GRA	
Eubacterium tortuosum		99.93	0.58	URE 1	99	
Fusobacterium mortiferum		0.07	0.20		1	
43601002	**Very good identification**					**43601002**
					GRA	
Eubacterium tortuosum		99.96	0.79		99	
Fusobacterium mortiferum		0.05	0.38		1	
43601003	**Good identification**					**43601003**
Eubacterium tortuosum		100.0	0.45	URE 1		
43601010	**Very good identification**					**43601010**
					GRA	
Eubacterium tortuosum		99.93	0.58	MLZ 1	99	
Fusobacterium mortiferum		0.07	0.20		1	
43601012	**Good identification**					**43601012**
Eubacterium tortuosum		100.0	0.45	MLZ 1		
43601020	**Very good identification**					**43601020**
					GRA	
Eubacterium tortuosum		100.0	0.79		99	
Fusobacterium mortiferum		0.00	0.20		1	
43601021	**Good identification**					**43601021**
Eubacterium tortuosum		100.0	0.45	URE 1		
43601022	**Very good identification**					**43601022**
Eubacterium tortuosum		100.0	0.66			
43601030	**Good identification**					**43601030**
Eubacterium tortuosum		100.0	0.45	MLZ 1		
43601040	**Very good identification**					**43601040**
					GRA	
Eubacterium tortuosum		99.93	0.58	SOR 1	99	
Fusobacterium mortiferum		0.07	0.20		1	
43601042	**Good identification**					**43601042**
Eubacterium tortuosum		100.0	0.45	SOR 1		
43601060	**Good identification**					**43601060**
Eubacterium tortuosum		100.0	0.45	SOR 1		
43601100	**Low discrimination**					**43601100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Eubacterium tortuosum		85.90	0.58	LAC 1	1	99	
Fusobacterium mortiferum		14.07	0.61		1	1	
Blautia hansenii		0.03	-.03		99	99	
43601102	**Acceptable identification**						**43601102**
					GRA		
Eubacterium tortuosum		90.56	0.45	LAC 1	99		
Fusobacterium mortiferum		9.34	0.45		1		
Bifidobacterium longum subsp. infantis		0.10	-.07		99		
43601120	**Good identification**						**43601120**
					GRA		
Eubacterium tortuosum		98.92	0.45	LAC 1	99		
Fusobacterium mortiferum		0.93	0.27		1		
Prevotella melaninogenica		0.16	-.04		1		
43601200	**Very good identification**						**43601200**
					GRA		
Eubacterium tortuosum		99.93	0.58	RHA 1	99		
Fusobacterium mortiferum		0.07	0.20		1		
43601202	**Good identification**						**43601202**
Eubacterium tortuosum		100.0	0.45	RHA 1			
43601220	**Good identification**						**43601220**
Eubacterium tortuosum		100.0	0.45	RHA 1			
43601400	**Very good identification**						**43601400**
					GRA		
Eubacterium tortuosum		99.93	0.58	ARA 1	99		
Fusobacterium mortiferum		0.07	0.20		1		
43601402	**Good identification**						**43601402**
Eubacterium tortuosum		100.0	0.45	ARA 1			
43601420	**Good identification**						**43601420**
Eubacterium tortuosum		100.0	0.45	ARA 1			
43603000	**Very good identification**						**43603000**
					GRA		
Eubacterium tortuosum		99.93	0.58	MAN 1	99		
Fusobacterium mortiferum		0.07	0.20		1		
43603002	**Good identification**						**43603002**
Eubacterium tortuosum		100.0	0.45	MAN 1			
43603020	**Good identification**						**43603020**
Eubacterium tortuosum		100.0	0.45	MAN 1			
43604100	**Good identification**						**43604100**
Fusobacterium mortiferum		100.0	0.39	FRU 99			
43605000	**Very good identification**						**43605000**
					GRA		
Eubacterium tortuosum		99.47	0.58	XYL 1	99		
Fusobacterium mortiferum		0.53	0.35		1		
43605002	**Good identification**						**43605002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Eubacterium tortuosum		99.67	0.45	XYL 1	99		
Fusobacterium mortiferum		0.33	0.19		1		
43605020	**Good identification**						**43605020**
Eubacterium tortuosum		100.0	0.45	XYL 1			
43605100	**Low discrimination**						**43605100**
					GRA		
Fusobacterium mortiferum		54.96	0.42	FRU 99	1		
Eubacterium tortuosum		45.04	0.24	XYL 7	99		
43610000	**Very good identification**						**43610000**
					GRA		
Fusobacterium mortiferum		99.78	0.86		1		
Eubacterium tortuosum		0.22	0.24		99		
43610001	**Good identification**						**43610001**
Fusobacterium mortiferum		100.0	0.52	URE 1			
43610002	**Very good identification**						**43610002**
					GRA		
Fusobacterium mortiferum		99.66	0.69	BGL 10	1		
Eubacterium tortuosum		0.35	0.11		99		
43610003	**Good identification**						**43610003**
Fusobacterium mortiferum		100.0	0.35	URE 1			
43610010	**Good identification**						**43610010**
					GRA	SPO	
Fusobacterium mortiferum		99.08	0.52	MLZ 1	1	1	
Clostridium difficile		0.93	0.01		99	99	
43610012	**Good identification**						**43610012**
Fusobacterium mortiferum		100.0	0.35	MLZ 1			
43610020	**Good identification**						**43610020**
					GRA		
Fusobacterium mortiferum		96.33	0.52	NAG 1	1		
Eubacterium tortuosum		3.67	0.11		99		
43610022	**Acceptable identification**						**43610022**
					GRA		
Fusobacterium mortiferum		94.29	0.35	NAG 1	1		
Eubacterium tortuosum		5.71	-.02	BGL 10	99		
43610040	**Good identification**						**43610040**
Fusobacterium mortiferum		100.0	0.52	SOR 1			
43610042	**Good identification**						**43610042**
Fusobacterium mortiferum		100.0	0.35	SOR 1			
43610100	**Excellent identification**						**43610100**
Fusobacterium mortiferum		100.0	0.92				
43610101	**Very good identification**						**43610101**
Fusobacterium mortiferum		100.0	0.58	URE 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
43610102	**Very good identification**					**43610102**
					GRA	
Fusobacterium mortiferum		99.98	0.76	BGL 10	1	
Bifidobacterium longum subsp. infantis		0.02	-.07		99	
43610103	**Good identification**					**43610103**
Fusobacterium mortiferum		100.0	0.42	URE 1		
43610110	**Very good identification**					**43610110**
Fusobacterium mortiferum		100.0	0.58	MLZ 1		
43610112	**Good identification**					**43610112**
Fusobacterium mortiferum		100.0	0.42	MLZ 1		
43610120	**Very good identification**					**43610120**
Fusobacterium mortiferum		100.0	0.58	NAG 1		
43610122	**Good identification**					**43610122**
					GRA	
Fusobacterium mortiferum		97.24	0.42	NAG 1	1	
Bifidobacterium longum subsp. infantis		2.76	-.03	BGL 10	99	
43610140	**Very good identification**					**43610140**
Fusobacterium mortiferum		100.0	0.58	SOR 1		
43610142	**Good identification**					**43610142**
Fusobacterium mortiferum		100.0	0.42	SOR 1		
43610200	**Good identification**					**43610200**
Fusobacterium mortiferum		100.0	0.52	RHA 1		
43610202	**Good identification**					**43610202**
Fusobacterium mortiferum		100.0	0.35	RHA 1		
43610300	**Very good identification**					**43610300**
Fusobacterium mortiferum		100.0	0.58	RHA 1		
43610302	**Good identification**					**43610302**
Fusobacterium mortiferum		100.0	0.42	RHA 1		
43610400	**Good identification**					**43610400**
Fusobacterium mortiferum		100.0	0.52	ARA 1		
43610402	**Good identification**					**43610402**
Fusobacterium mortiferum		100.0	0.35	ARA 1		
43610500	**Very good identification**					**43610500**
Fusobacterium mortiferum		100.0	0.58	ARA 1		
43610502	**Good identification**					**43610502**
Fusobacterium mortiferum		100.0	0.42	ARA 1		
43611000	**Low discrimination**					**43611000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		87.41	0.89		1	
Eubacterium tortuosum		12.58	0.58		99	
Lactobacillus fermentum		0.01	0.20		99	
43611001	**Low discrimination**					**43611001**
					GRA	
Fusobacterium mortiferum		86.82	0.55	URE 1	1	
Eubacterium tortuosum		12.50	0.24		99	
Actinomyces naeslundii		0.68	0.30		99	
43611002	**Low discrimination**					**43611002**
					GRA	
Fusobacterium mortiferum		81.30	0.72	BGL 10	1	
Eubacterium tortuosum		18.58	0.45		99	
Lactobacillus fermentum		0.06	0.20		99	
43611003	**Low discrimination**					**43611003**
					GRA	
Fusobacterium mortiferum		76.96	0.38	URE 1	1	
Eubacterium tortuosum		17.59	0.11	BGL 10	99	
Actinomyces naeslundii		5.46	0.30		99	
43611010	**Low discrimination**					**43611010**
					GRA	
Fusobacterium mortiferum		87.42	0.55	MLZ 1	1	
Eubacterium tortuosum		12.58	0.24		99	
43611012	**Low discrimination**					**43611012**
					GRA	
Fusobacterium mortiferum		81.40	0.38	MLZ 1	1	
Eubacterium tortuosum		18.61	0.11	BGL 10	99	
43611020	**Low discrimination**					**43611020**
					GRA	
Eubacterium tortuosum		71.44	0.45	FRU 1	99	
Fusobacterium mortiferum		28.41	0.55		1	
Actinomyces naeslundii		0.15	0.27		99	
43611040	**Low discrimination**					**43611040**
					GRA	
Fusobacterium mortiferum		87.42	0.55	SOR 1	1	
Eubacterium tortuosum		12.58	0.24		99	
43611042	**Low discrimination**					**43611042**
					GRA	
Fusobacterium mortiferum		81.40	0.38	SOR 1	1	
Eubacterium tortuosum		18.61	0.11	BGL 10	99	
43611100	**Excellent identification**					**43611100**
					GRA	
Fusobacterium mortiferum		99.91	0.95		1	
Eubacterium tortuosum		0.06	0.24		99	
Actinomyces naeslundii		0.01	0.35		99	
43611101	**Very good identification**					**43611101**
					GRA	
Fusobacterium mortiferum		99.05	0.61	URE 1	1	
Actinomyces naeslundii		0.95	0.38		99	
43611102	**Very good identification**					**43611102**
					GRA	
Fusobacterium mortiferum		98.69	0.79	BGL 10	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. infantis		1.05	0.27		99	
Eubacterium tortuosum		0.10	0.11		99	
43611103	**Acceptable identification**					**43611103**
					GRA	
Fusobacterium mortiferum		91.15	0.45	URE 1	1	
Actinomyces naeslundii		7.88	0.38	BGL 10	99	
Bifidobacterium longum subsp. infantis		0.97	-.07		99	
43611110	**Very good identification**					**43611110**
Fusobacterium mortiferum		99.73	0.61	MLZ 1		
Eubacterium saburreum		0.27	0.15			
43611112	**Acceptable identification**					**43611112**
					GRA	
Fusobacterium mortiferum		94.61	0.45	MLZ 1	1	
Bifidobacterium adolescentis		4.39	0.18	BGL 10	99	
Bifidobacterium longum subsp. infantis		1.01	-.07		99	
43611120	**Good identification**					**43611120**
					GRA	
Fusobacterium mortiferum		94.60	0.61	NAG 1	1	
Bifidobacterium longum subsp. infantis		1.94	0.15		99	
Prevotella melaninogenica		1.47	0.12		1	
43611123	**Low discrimination**					**43611123**
Actinomyces naeslundii		82.07	0.38	ESL 12		
Bifidobacterium longum subsp. infantis		17.93	-.03	MNS 88		
43611140	**Very good identification**					**43611140**
Fusobacterium mortiferum		100.0	0.61	SOR 1		
43611142	**Good identification**					**43611142**
					GRA	
Fusobacterium mortiferum		98.95	0.45	SOR 1	1	
Bifidobacterium longum subsp. infantis		1.05	-.07	BGL 10	99	
43611200	**Low discrimination**					**43611200**
					GRA	
Fusobacterium mortiferum		87.42	0.55	RHA 1	1	
Eubacterium tortuosum		12.58	0.24		99	
43611202	**Low discrimination**					**43611202**
					GRA	
Fusobacterium mortiferum		81.40	0.38	RHA 1	1	
Eubacterium tortuosum		18.61	0.11	BGL 10	99	
43611300	**Good identification**					**43611300**
					GRA	
Fusobacterium mortiferum		94.56	0.61	RHA 1	1	
Eubacterium contortum		5.44	0.27		99	
43611302	**Acceptable identification**					**43611302**
					GRA	
Fusobacterium mortiferum		90.74	0.45	RHA 1	1	
Eubacterium contortum		8.30	0.14	BGL 10	99	
Bifidobacterium longum subsp. infantis		0.96	-.07		99	
43611400	**Low discrimination**					**43611400**
					GRA	
Fusobacterium mortiferum		86.84	0.55	ARA 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium tortuosum		12.50	0.24		99		
Lactobacillus fermentum		0.66	0.20		99		
43611402	**Low discrimination**						**43611402**
					GRA		
Fusobacterium mortiferum		77.10	0.38	ARA 1	1		
Eubacterium tortuosum		17.62	0.11	BGL 10	99		
Lactobacillus fermentum		5.28	0.20		99		
43611500	**Very good identification**						**43611500**
					GRA		
Fusobacterium mortiferum		99.47	0.61	ARA 1	1		
Lactobacillus fermentum		0.53	0.24		99		
43611502	**Low discrimination**						**43611502**
					GRA		
Fusobacterium mortiferum		76.19	0.45	ARA 1	1		
Bifidobacterium adolescentis		19.35	0.31	BGL 10	99		
Lactobacillus fermentum		3.65	0.24		99		
43612000	**Good identification**						**43612000**
					GRA	SPO	
Fusobacterium mortiferum		97.83	0.52	MAN 1	1	1	
Clostridium difficile		1.68	0.06		99	99	
Eubacterium limosum		0.48	-.06		99	1	
43612002	**Good identification**						**43612002**
Fusobacterium mortiferum		100.0	0.35	MAN 1			
43612100	**Very good identification**						**43612100**
Fusobacterium mortiferum		100.0	0.58	MAN 1			
43612102	**Good identification**						**43612102**
Fusobacterium mortiferum		100.0	0.42	MAN 1			
43613000	**Low discrimination**						**43613000**
					GRA		
Fusobacterium mortiferum		86.65	0.55	MAN 1	1		
Eubacterium tortuosum		12.47	0.24		99		
Actinomyces naeslundii		0.88	0.32		99		
43613001	**Low discrimination**						**43613001**
					GRA		
Actinomyces naeslundii		60.22	0.35	ESL 12	99		
Fusobacterium mortiferum		39.78	0.20	MNS 88	1		
43613002	**Low discrimination**						**43613002**
					GRA		
Fusobacterium mortiferum		75.74	0.38	MAN 1	1		
Eubacterium tortuosum		17.31	0.11	BGL 10	99		
Actinomyces naeslundii		6.95	0.32		99		
43613003	**Good identification**						**43613003**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43613021	**Good identification**						**43613021**
Actinomyces naeslundii		100.0	0.35	ESL 12			
43613023	**Good identification**						**43613023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	ESL 12		
43613100	**Very good identification**					**43613100**
					GRA	
Fusobacterium mortiferum		98.77	0.61	MAN 1	1	
Actinomyces naeslundii		1.23	0.40		99	
43613101	**Low discrimination**					**43613101**
					GRA	
Actinomyces naeslundii		64.87	0.43	ESL 12	99	
Fusobacterium mortiferum		35.13	0.27	MNS 88	1	
43613102	**Low discrimination**					**43613102**
					GRA	
Fusobacterium mortiferum		86.69	0.45	MAN 1	1	
Actinomyces naeslundii		9.70	0.40	BGL 10	99	
Bifidobacterium adolescentis		2.69	0.15		99	
43613103	**Good identification**					**43613103**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43613120	**Low discrimination**					**43613120**
					GRA	
Actinomyces naeslundii		55.18	0.40	ESL 12	99	
Fusobacterium mortiferum		44.82	0.27	MNS 88	1	
43613121	**Good identification**					**43613121**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43613122	**Low discrimination**					**43613122**
Actinomyces naeslundii		85.56	0.40	ESL 12		
Bifidobacterium longum subsp. infantis		14.44	-.03	MNS 88		
43613123	**Good identification**					**43613123**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43614000	**Very good identification**					**43614000**
Fusobacterium mortiferum		100.0	0.67	XYL 7		
43614002	**Good identification**					**43614002**
Fusobacterium mortiferum		100.0	0.50	XYL 7		
43614100	**Very good identification**					**43614100**
Fusobacterium mortiferum		100.0	0.73	XYL 7		
43614101	**Good identification**					**43614101**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
43614102	**Very good identification**					**43614102**
Fusobacterium mortiferum		100.0	0.57	XYL 7		
43614110	**Good identification**					**43614110**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
43614120	**Good identification**					**43614120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.39	XYL 7		
43614140	**Good identification**					**43614140**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
43614500	**Good identification**					**43614500**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
43615000	**Very good identification**					**43615000**
					GRA	
Fusobacterium mortiferum		98.05	0.70	XYL 7	1	
Eubacterium tortuosum		1.89	0.24		99	
Lactobacillus fermentum		0.06	0.16		99	
43615001	**Good identification**					**43615001**
Fusobacterium mortiferum		100.0	0.35	XYL 7		
43615002	**Good identification**					**43615002**
					GRA	
Fusobacterium mortiferum		96.52	0.53	XYL 7	1	
Eubacterium tortuosum		2.96	0.11	BGL 10	99	
Lactobacillus fermentum		0.52	0.16		99	
43615010	**Good identification**					**43615010**
Fusobacterium mortiferum		100.0	0.35	XYL 7		
43615020	**Low discrimination**					**43615020**
					GRA	
Fusobacterium mortiferum		74.77	0.35	XYL 7	1	
Eubacterium tortuosum		25.23	0.11	NAG 1	99	
43615040	**Good identification**					**43615040**
Fusobacterium mortiferum		100.0	0.35	XYL 7		
43615100	**Very good identification**					**43615100**
					GRA	
Fusobacterium mortiferum		99.03	0.76	XYL 7	1	
Eubacterium contortum		0.77	0.27		99	
Bifidobacterium longum subsp. infantis		0.13	0.09		99	
43615101	**Good identification**					**43615101**
Fusobacterium mortiferum		100.0	0.42	XYL 7		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43615102	**Low discrimination**					**43615102**
					GRA	
Fusobacterium mortiferum		85.61	0.60	XYL 7	1	
Bifidobacterium longum subsp. infantis		9.89	0.26	BGL 10	99	
Bifidobacterium adolescentis		2.92	0.31		99	
43615110	**Good identification**					**43615110**
Fusobacterium mortiferum		98.10	0.42	XYL 7		
Eubacterium saburreum		1.90	0.11	MLZ 1		
43615120	**Low discrimination**					**43615120**
					GRA	
Fusobacterium mortiferum		73.62	0.42	XYL 7	1	
Bifidobacterium longum subsp. infantis		16.45	0.13	NAG 1	99	
Eubacterium contortum		9.94	0.14		99	
43615132	**Good identification**					**43615132**
					GRA	
Parabacteroides distasonis		96.84	0.38	MNS 98	1	
Bifidobacterium longum subsp. infantis		3.16	-.04		99	
43615140	**Good identification**					**43615140**
Fusobacterium mortiferum		100.0	0.42	XYL 7		
43615300	**Very good identification**					**43615300**
					GRA	
Eubacterium contortum		98.70	0.61	MLT 99	99	
Fusobacterium mortiferum		1.30	0.42		1	
43615302	**Good identification**					**43615302**
					GRA	
Eubacterium contortum		98.89	0.48	MLT 99	99	
Fusobacterium mortiferum		0.82	0.25		1	
Parabacteroides distasonis		0.29	0.16		1	
43615320	**Good identification**					**43615320**
					GRA	
Eubacterium contortum		99.19	0.48	MLT 99	99	
Bacteroides vulgatus		0.52	0.02		1	
Parabacteroides distasonis		0.29	0.16		1	
43615322	**Low discrimination**					**43615322**
					GRA	
Parabacteroides distasonis		61.78	0.50	MNS 98	1	
Eubacterium contortum		37.44	0.35		99	
Prevotella oralis		0.41	0.00		1	
43615332	**Good identification**					**43615332**
					GRA	
Parabacteroides distasonis		98.88	0.46	MNS 98	1	
Eubacterium contortum		1.12	0.01		99	
43615400	**Acceptable identification**					**43615400**
					GRA	
Fusobacterium mortiferum		94.40	0.35	XYL 7	1	
Lactobacillus fermentum		5.60	0.16	ARA 1	99	
43615500	**Low discrimination**					**43615500**
					GRA	
Fusobacterium mortiferum		59.08	0.42	XYL 7	1	
Eubacterium contortum		36.97	0.26	ARA 1	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		2.45	0.20		99	
43615502	**Low discrimination**					**43615502**
					GRA	
Bifidobacterium adolescentis		89.80	0.45	MNS 87	99	
Fusobacterium mortiferum		3.98	0.25	MLT 99	1	
Eubacterium contortum		3.95	0.13		99	
43615512	**Good identification**					**43615512**
Bifidobacterium adolescentis		99.34	0.46	MNS 87		
Bifidobacterium longum subsp. longum		0.66	0.07	MLT 99		
43615522	**Low discrimination**					**43615522**
					GRA	
Parabacteroides distasonis		87.29	0.39	MNS 98	1	
Bifidobacterium longum subsp. longum		5.28	0.14		99	
Bifidobacterium adolescentis		2.78	0.11		99	
43615700	**Very good identification**					**43615700**
Eubacterium contortum		100.0	0.60	MLT 99		
43615702	**Good identification**					**43615702**
					GRA	
Eubacterium contortum		99.55	0.47	MLT 99	99	
Bifidobacterium adolescentis		0.23	0.11		99	
Parabacteroides distasonis		0.22	0.13		1	
43615720	**Acceptable identification**					**43615720**
					GRA	
Eubacterium contortum		91.97	0.47	MLT 99	99	
Bacteroides vulgatus		7.83	0.21		1	
Parabacteroides distasonis		0.20	0.13		1	
43615722	**Low discrimination**					**43615722**
					GRA	
Parabacteroides distasonis		55.28	0.47	MNS 98	1	
Eubacterium contortum		44.72	0.34		99	
43615732	**Good identification**					**43615732**
					GRA	
Parabacteroides distasonis		96.48	0.42	MNS 98	1	
Clostridium clostridioforme		2.06	0.08		50	
Eubacterium contortum		1.46	0.00		99	
43616100	**Good identification**					**43616100**
Fusobacterium mortiferum		100.0	0.39	MAN 1		
43617000	**Acceptable identification**					**43617000**
					GRA	
Fusobacterium mortiferum		94.38	0.35	MAN 1	1	
Actinomyces israelii		5.62	0.17	XYL 7	99	
43617100	**Acceptable identification**					**43617100**
					GRA	
Fusobacterium mortiferum		90.74	0.42	MAN 1	1	
Actinomyces israelii		9.26	0.27	XYL 7	99	
43617502	**Acceptable identification**					**43617502**
Bifidobacterium adolescentis		94.25	0.43	MNS 87		
Actinomyces israelii		5.75	0.29	MLT 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
43617512	**Good identification**						**43617512**
Bifidobacterium adolescentis		99.26	0.45	MNS 87			
Actinomyces israelii		0.74	0.15	MLT 99			
43620000	**Low discrimination**						**43620000**
					GRA	SPO	
Fusobacterium mortiferum		57.21	0.43	FRU 99	1	1	
Eubacterium tortuosum		41.35	0.24		99	1	
Clostridium botulinum biovar A		1.45	0.02		99	99	
43620100	**Good identification**						**43620100**
Fusobacterium mortiferum		100.0	0.49	FRU 99			
43621000	**Very good identification**						**43621000**
					GRA		
Eubacterium tortuosum		97.95	0.58	TRE 1	99		
Fusobacterium mortiferum		2.05	0.46		1		
43621002	**Good identification**						**43621002**
					GRA		
Eubacterium tortuosum		98.70	0.45	TRE 1	99		
Fusobacterium mortiferum		1.30	0.29		1		
43621020	**Good identification**						**43621020**
					GRA		
Eubacterium tortuosum		99.88	0.45	TRE 1	99		
Fusobacterium mortiferum		0.12	0.12		1		
43621100	**Low discrimination**						**43621100**
					GRA		
Fusobacterium mortiferum		82.89	0.52	FRU 99	1		
Eubacterium tortuosum		17.12	0.24		99		
43621102	**Low discrimination**						**43621102**
					GRA		
Fusobacterium mortiferum		75.30	0.36	FRU 99	1		
Eubacterium tortuosum		24.70	0.11	BGL 10	99		
43630000	**Very good identification**						**43630000**
Fusobacterium mortiferum		100.0	0.77				
43630001	**Good identification**						**43630001**
Fusobacterium mortiferum		100.0	0.43	URE 1			
43630002	**Very good identification**						**43630002**
Fusobacterium mortiferum		100.0	0.60	BGL 10			
43630010	**Good identification**						**43630010**
Fusobacterium mortiferum		100.0	0.43	MLZ 1			
43630020	**Good identification**						**43630020**
Fusobacterium mortiferum		100.0	0.43	NAG 1			
43630040	**Good identification**						**43630040**
Fusobacterium mortiferum		100.0	0.43	SOR 1			
43630100	**Very good identification**						**43630100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.83			
43630101	**Good identification**					**43630101**
Fusobacterium mortiferum		100.0	0.49	URE 1		
43630102	**Very good identification**					**43630102**
Fusobacterium mortiferum		100.0	0.67	BGL 10		
43630110	**Good identification**					**43630110**
Fusobacterium mortiferum		100.0	0.49	MLZ 1		
43630120	**Good identification**					**43630120**
Fusobacterium mortiferum		100.0	0.49	NAG 1		
43630140	**Good identification**					**43630140**
Fusobacterium mortiferum		100.0	0.49	SOR 1		
43630200	**Good identification**					**43630200**
Fusobacterium mortiferum		100.0	0.43	RHA 1		
43630300	**Good identification**					**43630300**
Fusobacterium mortiferum		100.0	0.49	RHA 1		
43630400	**Good identification**					**43630400**
Fusobacterium mortiferum		100.0	0.43	ARA 1		
43630500	**Good identification**					**43630500**
Fusobacterium mortiferum		100.0	0.49	ARA 1		
43631000	**Very good identification**					**43631000**
					GRA	
Fusobacterium mortiferum		99.48	0.80		1	
Eubacterium tortuosum		0.48	0.24		99	
Actinomyces naeslundii		0.03	0.32		99	
43631001	**Good identification**					**43631001**
					GRA	
Fusobacterium mortiferum		95.13	0.46	URE 1	1	
Actinomyces naeslundii		4.87	0.35		99	
43631002	**Very good identification**					**43631002**
					GRA	
Fusobacterium mortiferum		98.93	0.64	BGL 10	1	
Eubacterium tortuosum		0.77	0.11		99	
Actinomyces naeslundii		0.31	0.32		99	
43631010	**Good identification**					**43631010**
Fusobacterium mortiferum		100.0	0.46	MLZ 1		
43631020	**Low discrimination**					**43631020**
					GRA	
Fusobacterium mortiferum		89.35	0.46	NAG 1	1	
Eubacterium tortuosum		7.60	0.11		99	
Actinomyces naeslundii		3.05	0.32		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43631021	**Low discrimination**					**43631021**
					GRA	
Actinomyces naeslundii		83.52	0.35	ESL 12	99	
Fusobacterium mortiferum		16.48	0.12	MNS 88	1	
43631023	**Good identification**					**43631023**
Actinomyces naeslundii		100.0	0.35	ESL 12		
43631040	**Good identification**					**43631040**
Fusobacterium mortiferum		100.0	0.46	SOR 1		
43631100	**Very good identification**					**43631100**
					GRA	
Fusobacterium mortiferum		99.95	0.86		1	
Actinomyces naeslundii		0.04	0.40		99	
Eubacterium saburreum		0.01	0.14		85	
43631101	**Acceptable identification**					**43631101**
					GRA	
Fusobacterium mortiferum		94.12	0.52	URE 1	1	
Actinomyces naeslundii		5.88	0.43		99	
43631102	**Very good identification**					**43631102**
					GRA	
Fusobacterium mortiferum		99.50	0.70	BGL 10	1	
Actinomyces naeslundii		0.38	0.40		99	
Bifidobacterium adolescentis		0.09	0.13		99	
43631103	**Low discrimination**					**43631103**
					GRA	
Fusobacterium mortiferum		64.02	0.36	URE 1	1	
Actinomyces naeslundii		35.98	0.43	BGL 10	99	
43631110	**Good identification**					**43631110**
Fusobacterium mortiferum		99.40	0.52	MLZ 1		
Eubacterium saburreum		0.60	0.12			
43631112	**Low discrimination**					**43631112**
					GRA	
Fusobacterium mortiferum		87.32	0.36	MLZ 1	1	
Bifidobacterium adolescentis		9.04	0.15	BGL 10	99	
Leptotrichia buccalis		3.64	-.11		1	
43631120	**Good identification**					**43631120**
					GRA	
Fusobacterium mortiferum		96.00	0.52	NAG 1	1	
Actinomyces naeslundii		4.00	0.40		99	
43631121	**Low discrimination**					**43631121**
					GRA	
Actinomyces naeslundii		86.08	0.43	ESL 12	99	
Fusobacterium mortiferum		13.92	0.18	MNS 88	1	
43631122	**Low discrimination**					**43631122**
					GRA	
Parabacteroides distasonis		49.67	0.37	MNS 98	1	
Fusobacterium mortiferum		33.36	0.36	XYL 90	1	
Actinomyces naeslundii		12.50	0.40		99	
43631123	**Good identification**					**43631123**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ESL 12		
43631140	**Good identification**					**43631140**
					GRA	
Fusobacterium mortiferum		99.27	0.52	SOR 1	1	
Actinomyces naeslundii		0.73	0.27		99	
43631142	**Acceptable identification**					**43631142**
					GRA	
Fusobacterium mortiferum		93.80	0.36	SOR 1	1	
Actinomyces naeslundii		6.20	0.27	BGL 10	99	
43631200	**Good identification**					**43631200**
Fusobacterium mortiferum		100.0	0.46	RHA 1		
43631300	**Good identification**					**43631300**
Fusobacterium mortiferum		100.0	0.52	RHA 1		
43631302	**Good identification**					**43631302**
Fusobacterium mortiferum		95.68	0.36	RHA 1		
Parabacteroides distasonis		4.32	0.11	BGL 10		
43631322	**Good identification**					**43631322**
Parabacteroides distasonis		100.0	0.45	MNS 98		
43631332	**Good identification**					**43631332**
Parabacteroides distasonis		100.0	0.41	MNS 98		
43631400	**Good identification**					**43631400**
Fusobacterium mortiferum		100.0	0.46	ARA 1		
43631500	**Good identification**					**43631500**
Fusobacterium mortiferum		100.0	0.52	ARA 1		
43631502	**Low discrimination**					**43631502**
					GRA	
Fusobacterium mortiferum		63.82	0.36	ARA 1	1	
Bifidobacterium adolescentis		36.18	0.28	BGL 10	99	
43631722	**Good identification**					**43631722**
Parabacteroides distasonis		100.0	0.42	MNS 98		
43631732	**Good identification**					**43631732**
Parabacteroides distasonis		97.04	0.37	MNS 98		
Clostridium clostridioforme		2.97	0.06	XYL 90		
43632000	**Good identification**					**43632000**
Fusobacterium mortiferum		100.0	0.43	MAN 1		
43632100	**Good identification**					**43632100**
Fusobacterium mortiferum		100.0	0.49	MAN 1		
43633000	**Acceptable identification**					**43633000**
					GRA	
Fusobacterium mortiferum		93.79	0.46	MAN 1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		6.21	0.37		99	
43633001	**Acceptable identification**					**43633001**
					GRA	
Actinomyces naeslundii		90.77	0.40	ESL 12	99	
Fusobacterium mortiferum		9.23	0.12	MNS 88	1	
43633003	**Good identification**					**43633003**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43633020	**Low discrimination**					**43633020**
					GRA	
Actinomyces naeslundii		86.77	0.37	ESL 12	99	
Fusobacterium mortiferum		13.23	0.12	MNS 88	1	
43633021	**Good identification**					**43633021**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43633022	**Good identification**					**43633022**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43633023	**Good identification**					**43633023**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43633100	**Acceptable identification**					**43633100**
					GRA	
Fusobacterium mortiferum		91.81	0.52	MAN 1	1	
Actinomyces naeslundii		7.42	0.45		99	
Actinomyces israelii		0.77	0.19		99	
43633101	**Acceptable identification**					**43633101**
					GRA	
Actinomyces naeslundii		92.31	0.48	ESL 12	99	
Fusobacterium mortiferum		7.69	0.18	MNS 88	1	
43633102	**Low discrimination**					**43633102**
					GRA	
Fusobacterium mortiferum		52.35	0.36	MAN 1	1	
Actinomyces naeslundii		38.08	0.45	BGL 10	99	
Actinomyces israelii		5.94	0.22		99	
43633103	**Good identification**					**43633103**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43633120	**Low discrimination**					**43633120**
					GRA	
Actinomyces naeslundii		88.89	0.45	ESL 12	99	
Fusobacterium mortiferum		11.11	0.18	MNS 88	1	
43633121	**Good identification**					**43633121**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43633122	**Good identification**					**43633122**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43633123	**Good identification**					**43633123**
Actinomyces naeslundii		100.0	0.48	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43634000	**Very good identification**					**43634000**
Fusobacterium mortiferum		100.0	0.58	XYL 7		
43634002	**Good identification**					**43634002**
Fusobacterium mortiferum		100.0	0.41	XYL 7		
43634100	**Very good identification**					**43634100**
Fusobacterium mortiferum		100.0	0.64	XYL 7		
43634102	**Good identification**					**43634102**
Fusobacterium mortiferum		100.0	0.48	XYL 7		
43635000	**Very good identification**					**43635000**
					GRA	
Fusobacterium mortiferum		99.87	0.61	XYL 7	1	
Actinomyces israelii		0.13	0.14		99	
43635002	**Good identification**					**43635002**
					GRA	
Fusobacterium mortiferum		98.23	0.44	XYL 7	1	
Actinomyces israelii		1.77	0.17	BGL 10	99	
43635100	**Very good identification**					**43635100**
					GRA	
Fusobacterium mortiferum		99.72	0.67	XYL 7	1	
Actinomyces israelii		0.23	0.24		99	
Eubacterium saburreum		0.05	0.09		85	
43635102	**Low discrimination**					**43635102**
					GRA	
Fusobacterium mortiferum		88.88	0.51	XYL 7	1	
Bifidobacterium adolescentis		6.76	0.28	BGL 10	99	
Actinomyces israelii		2.75	0.27		99	
43635122	**Good identification**					**43635122**
					GRA	
Parabacteroides distasonis		99.06	0.54	MNS 98	1	
Fusobacterium mortiferum		0.56	0.16		1	
Bifidobacterium longum subsp. infantis		0.38	-.04		99	
43635132	**Good identification**					**43635132**
Parabacteroides distasonis		100.0	0.49	MNS 98		
43635322	**Very good identification**					**43635322**
					GRA	
Parabacteroides distasonis		99.87	0.62	MNS 98	1	
Eubacterium contortum		0.13	0.01		99	
43635332	**Very good identification**					**43635332**
Parabacteroides distasonis		100.0	0.57	MNS 98		
43635502	**Acceptable identification**					**43635502**
					GRA	
Bifidobacterium adolescentis		91.63	0.42	MNS 87	99	
Actinomyces israelii		4.55	0.26	MLT 99	99	
Parabacteroides distasonis		2.00	0.16		1	
43635512	**Good identification**					**43635512**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		97.75	0.43	MNS 87	99	
Clostridium clostridioforme		1.31	0.08	MLT 99	50	
Parabacteroides distasonis		0.94	0.11		1	
43635522	**Good identification**					**43635522**
					GRA	
Parabacteroides distasonis		99.54	0.50	MNS 98	1	
Bifidobacterium adolescentis		0.47	0.07		99	
43635532	**Good identification**					**43635532**
					GRA	
Parabacteroides distasonis		96.61	0.45	MNS 98	1	
Clostridium clostridioforme		1.36	0.08		50	
Bifidobacterium adolescentis		1.02	0.09		99	
43635722	**Very good identification**					**43635722**
					GRA	
Parabacteroides distasonis		99.42	0.58	MNS 98	1	
Clostridium clostridioforme		0.40	0.11		50	
Eubacterium contortum		0.18	0.00		99	
43635732	**Good identification**					**43635732**
Parabacteroides distasonis		96.68	0.53	MNS 98		
Clostridium clostridioforme		3.32	0.23			
43637100	**Low discrimination**					**43637100**
					GRA	
Actinomyces israelii		56.23	0.37	MLT 99	99	
Fusobacterium mortiferum		43.77	0.33	CEL 93	1	
43637102	**Low discrimination**					**43637102**
					GRA	
Actinomyces israelii		70.78	0.40	MLT 99	99	
Bifidobacterium adolescentis		25.14	0.26	CEL 93	99	
Fusobacterium mortiferum		4.08	0.16		1	
43637500	**Good identification**					**43637500**
Actinomyces israelii		100.0	0.35	MLT 99		
43637502	**Low discrimination**					**43637502**
Bifidobacterium adolescentis		74.40	0.40	MNS 87		
Actinomyces israelii		25.60	0.39	MLT 99		
43637512	**Acceptable identification**					**43637512**
Bifidobacterium adolescentis		94.43	0.42	MNS 87		
Actinomyces israelii		3.97	0.24	MLT 99		
Bifidobacterium dentium		1.59	-.07			
43640000	**Low discrimination**					**43640000**
					GRA	
Eubacterium tortuosum		87.69	0.41	CEL 10	99	
Fusobacterium mortiferum		12.31	0.43	GAL 99	1	
43640100	**Good identification**					**43640100**
					GRA	
Fusobacterium mortiferum		97.01	0.50	FRU 99	1	
Eubacterium tortuosum		2.99	0.07		99	
43641000	**Very good identification**					**43641000**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium tortuosum		99.79	0.76	CEL 10	99		
Fusobacterium mortiferum		0.21	0.46		1		
43641001	**Good identification**						**43641001**
					GRA		
Eubacterium tortuosum		99.79	0.41	CEL 10	99		
Fusobacterium mortiferum		0.21	0.12	URE 1	1		
43641002	**Very good identification**						**43641002**
					GRA		
Eubacterium tortuosum		99.87	0.63	CEL 10	99		
Fusobacterium mortiferum		0.13	0.30		1		
43641010	**Good identification**						**43641010**
					GRA		
Eubacterium tortuosum		99.79	0.41	CEL 10	99		
Fusobacterium mortiferum		0.21	0.12	MLZ 1	1		
43641020	**Very good identification**						**43641020**
					GRA		
Eubacterium tortuosum		99.99	0.63	CEL 10	99		
Fusobacterium mortiferum		0.01	0.12		1		
43641022	**Good identification**						**43641022**
Eubacterium tortuosum		100.0	0.50	CEL 10			
43641040	**Good identification**						**43641040**
					GRA		
Eubacterium tortuosum		99.79	0.41	CEL 10	99		
Fusobacterium mortiferum		0.21	0.12	SOR 1	1		
43641100	**Low discrimination**						**43641100**
					GRA		
Eubacterium tortuosum		67.06	0.41	CEL 10	99		
Fusobacterium mortiferum		32.94	0.53	LAC 1	1		
43641200	**Good identification**						**43641200**
					GRA		
Eubacterium tortuosum		99.79	0.41	CEL 10	99		
Fusobacterium mortiferum		0.21	0.12	RHA 1	1		
43641400	**Good identification**						**43641400**
					GRA		
Eubacterium tortuosum		99.79	0.41	CEL 10	99		
Fusobacterium mortiferum		0.21	0.12	ARA 1	1		
43643000	**Good identification**						**43643000**
					GRA	SPO	
Eubacterium tortuosum		99.57	0.41	CEL 10	99	1	
Clostridium innocuum		0.22	0.00	MAN 1	99	99	
Fusobacterium mortiferum		0.21	0.12		1	1	
43645000	**Good identification**						**43645000**
					GRA		
Eubacterium tortuosum		98.44	0.41	CEL 10	99		
Fusobacterium mortiferum		1.56	0.27	XYL 1	1		
43650000	**Very good identification**						**43650000**
					GRA		
Fusobacterium mortiferum		99.93	0.78		1		
Eubacterium tortuosum		0.07	0.07		99		
43650001	**Good identification**						**43650001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.43	URE 1		
43650002	**Very good identification**					**43650002**
					GRA	
Fusobacterium mortiferum		99.89	0.61	BGL 10	1	
Eubacterium tortuosum		0.12	-.06		99	
43650010	**Good identification**					**43650010**
Fusobacterium mortiferum		100.0	0.43	MLZ 1		
43650020	**Good identification**					**43650020**
					GRA	
Fusobacterium mortiferum		98.75	0.43	NAG 1	1	
Eubacterium tortuosum		1.25	-.06		99	
43650040	**Good identification**					**43650040**
Fusobacterium mortiferum		100.0	0.43	SOR 1		
43650100	**Very good identification**					**43650100**
Fusobacterium mortiferum		100.0	0.84			
43650101	**Good identification**					**43650101**
Fusobacterium mortiferum		100.0	0.50	URE 1		
43650102	**Very good identification**					**43650102**
Fusobacterium mortiferum		100.0	0.68	BGL 10		
43650110	**Good identification**					**43650110**
Fusobacterium mortiferum		100.0	0.50	MLZ 1		
43650120	**Good identification**					**43650120**
Fusobacterium mortiferum		100.0	0.50	NAG 1		
43650140	**Good identification**					**43650140**
Fusobacterium mortiferum		100.0	0.50	SOR 1		
43650200	**Good identification**					**43650200**
Fusobacterium mortiferum		100.0	0.43	RHA 1		
43650300	**Good identification**					**43650300**
Fusobacterium mortiferum		100.0	0.50	RHA 1		
43650400	**Good identification**					**43650400**
Fusobacterium mortiferum		100.0	0.43	ARA 1		
43650500	**Good identification**					**43650500**
Fusobacterium mortiferum		100.0	0.50	ARA 1		
43651000	**Very good identification**					**43651000**
					GRA	
Fusobacterium mortiferum		95.36	0.81		1	
Eubacterium tortuosum		4.58	0.41		99	
Lactobacillus catenaformis		0.04	0.12		99	
43651001	**Acceptable identification**					**43651001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		92.65	0.46	URE 1	1		
Eubacterium tortuosum		4.45	0.07		99		
Actinomyces naeslundii		2.90	0.32		99		
43651002	**Good identification**						**43651002**
					GRA		
Fusobacterium mortiferum		92.19	0.64	BGL 10	1		
Eubacterium tortuosum		7.02	0.28		99		
Lactobacillus catenaformis		0.32	0.12		99		
43651010	**Good identification**						**43651010**
					GRA		
Fusobacterium mortiferum		95.42	0.46	MLZ 1	1		
Eubacterium tortuosum		4.58	0.07		99		
43651020	**Low discrimination**						**43651020**
					GRA	SPO	
Fusobacterium mortiferum		52.84	0.46	NAG 1	1	1	
Eubacterium tortuosum		44.28	0.28		99	1	
Clostridium barati		1.27	0.04		99	99	
43651040	**Good identification**						**43651040**
					GRA		
Fusobacterium mortiferum		95.42	0.46	SOR 1	1		
Eubacterium tortuosum		4.58	0.07		99		
43651100	**Very good identification**						**43651100**
					GRA		
Fusobacterium mortiferum		99.89	0.87		1		
Actinomyces naeslundii		0.03	0.37		99		
Lactobacillus catenaformis		0.03	0.15		99		
43651101	**Good identification**						**43651101**
					GRA		
Fusobacterium mortiferum		96.32	0.53	URE 1	1		
Actinomyces naeslundii		3.68	0.40		99		
43651102	**Very good identification**						**43651102**
					GRA		
Fusobacterium mortiferum		97.84	0.71	BGL 10	1		
Lactobacillus acidophilus		0.46	0.18		99		
Bifidobacterium breve		0.44	0.27		99		
43651103	**Low discrimination**						**43651103**
					GRA		
Fusobacterium mortiferum		74.41	0.36	URE 1	1		
Actinomyces naeslundii		25.59	0.40	BGL 10	99		
43651110	**Good identification**						**43651110**
					GRA		
Fusobacterium mortiferum		99.52	0.53	MLZ 1	1		
Bifidobacterium breve		0.49	0.10		99		
43651112	**Low discrimination**						**43651112**
					GRA		
Fusobacterium mortiferum		56.29	0.36	MLZ 1	1		
Bifidobacterium breve		24.97	0.27	BGL 10	99		
Bifidobacterium adolescentis		16.64	0.23		99		
43651120	**Low discrimination**						**43651120**
					GRA	SPO	
Fusobacterium mortiferum		85.53	0.53	NAG 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		5.86	0.13		99	99	
Clostridium barati		2.64	0.12		99	99	
43651121	**Low discrimination**						**43651121**
					GRA		
Actinomyces naeslundii		79.09	0.40	ESL 12	99		
Fusobacterium mortiferum		20.91	0.19	MNS 88	1		
43651123	**Acceptable identification**						**43651123**
					GRA		
Actinomyces naeslundii		90.61	0.40	ESL 12	99		
Prevotella oralis		9.39	-.03	MNS 88	1		
43651140	**Good identification**						**43651140**
					GRA		
Fusobacterium mortiferum		99.07	0.53	SOR 1	1		
Bifidobacterium breve		0.48	0.10		99		
Actinomyces naeslundii		0.45	0.24		99		
43651142	**Low discrimination**						**43651142**
					GRA		
Fusobacterium mortiferum		64.90	0.36	SOR 1	1		
Bifidobacterium breve		28.79	0.27	BGL 10	99		
Lactobacillus acidophilus		3.69	0.02		99		
43651200	**Good identification**						**43651200**
					GRA		
Fusobacterium mortiferum		95.42	0.46	RHA 1	1		
Eubacterium tortuosum		4.58	0.07		99		
43651300	**Good identification**						**43651300**
Fusobacterium mortiferum		100.0	0.53	RHA 1			
43651302	**Low discrimination**						**43651302**
					GRA		
Fusobacterium mortiferum		89.75	0.36	RHA 1	1		
Prevotella oralis		7.15	0.04	BGL 10	1		
Lactobacillus acidophilus		3.11	-.02		99		
43651400	**Good identification**						**43651400**
					GRA		
Fusobacterium mortiferum		95.42	0.46	ARA 1	1		
Eubacterium tortuosum		4.58	0.07		99		
43651500	**Good identification**						**43651500**
					GRA		
Fusobacterium mortiferum		99.62	0.53	ARA 1	1		
Actinomyces israelii		0.38	0.14		99		
43651502	**Low discrimination**						**43651502**
					GRA		
Bifidobacterium adolescentis		60.62	0.36	MNS 87	99		
Fusobacterium mortiferum		37.44	0.36	MLT 99	1		
Actinomyces israelii		1.94	0.17	XYL 87	99		
43651512	**Good identification**						**43651512**
Bifidobacterium adolescentis		100.0	0.38	MNS 87			
43652000	**Good identification**						**43652000**
					GRA	SPO	
Fusobacterium mortiferum		98.47	0.43	MAN 1	1	1	
Clostridium innocuum		1.53	0.00		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43652100	**Good identification**						**43652100**
Fusobacterium mortiferum		100.0	0.50	MAN 1			
43653000	**Low discrimination**						**43653000**
					GRA	SPO	
Fusobacterium mortiferum		76.88	0.46	MAN 1	1	1	
Clostridium innocuum		15.13	0.22		99	99	
Eubacterium tortuosum		3.69	0.07		99	1	
43653001	**Low discrimination**						**43653001**
					GRA		
Actinomyces naeslundii		85.76	0.37	ESL 12	99		
Fusobacterium mortiferum		14.25	0.12	MNS 88	1		
43653003	**Good identification**						**43653003**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43653021	**Good identification**						**43653021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43653023	**Good identification**						**43653023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43653100	**Acceptable identification**						**43653100**
					GRA		
Fusobacterium mortiferum		92.51	0.53	MAN 1	1		
Actinomyces naeslundii		4.58	0.42		99		
Actinomyces israelii		2.46	0.28		99		
43653101	**Low discrimination**						**43653101**
					GRA		
Actinomyces naeslundii		88.01	0.45	ESL 12	99		
Fusobacterium mortiferum		11.99	0.19	MNS 88	1		
43653102	**Low discrimination**						**43653102**
					GRA		
Fusobacterium mortiferum		40.89	0.36	MAN 1	1		
Actinomyces naeslundii		18.20	0.42	BGL 10	99		
Bifidobacterium breve		18.14	0.27		99		
43653103	**Good identification**						**43653103**
Actinomyces naeslundii		100.0	0.45	ESL 12			
43653120	**Low discrimination**						**43653120**
					GRA		
Actinomyces naeslundii		68.03	0.42	ESL 12	99		
Fusobacterium mortiferum		13.90	0.19	MNS 88	1		
Bifidobacterium breve		11.92	0.15	MLT 94	99		
43653121	**Good identification**						**43653121**
Actinomyces naeslundii		100.0	0.45	ESL 12			
43653123	**Good identification**						**43653123**
Actinomyces naeslundii		100.0	0.45	ESL 12			
43653512	**Good identification**						**43653512**
Bifidobacterium adolescentis		97.36	0.36	MNS 87			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		2.64	0.16	MLT 99			
43654000	**Very good identification**						**43654000**
Fusobacterium mortiferum		100.0	0.58	XYL 7			
43654002	**Good identification**						**43654002**
Fusobacterium mortiferum		100.0	0.42	XYL 7			
43654100	**Very good identification**						**43654100**
Fusobacterium mortiferum		100.0	0.65	XYL 7			
43654102	**Good identification**						**43654102**
Fusobacterium mortiferum		100.0	0.48	XYL 7			
43655000	**Very good identification**						**43655000**
					GRA		
Fusobacterium mortiferum		98.95	0.61	XYL 7	1		
Eubacterium tortuosum		0.64	0.07		99		
Actinomyces israelii		0.42	0.23		99		
43655002	**Acceptable identification**						**43655002**
					GRA		
Fusobacterium mortiferum		93.69	0.45	XYL 7	1		
Actinomyces israelii		5.35	0.26	BGL 10	99		
Eubacterium tortuosum		0.96	-.06		99		
43655100	**Very good identification**						**43655100**
					GRA	SPO	
Fusobacterium mortiferum		99.21	0.68	XYL 7	1	1	
Actinomyces israelii		0.72	0.33		99	1	
Clostridium butyricum		0.07	-.05		99	99	
43655102	**Low discrimination**						**43655102**
					GRA		
Fusobacterium mortiferum		76.04	0.51	XYL 7	1		
Bifidobacterium adolescentis		16.52	0.36	BGL 10	99		
Actinomyces israelii		7.44	0.36		99		
43655112	**Acceptable identification**						**43655112**
					GRA		
Bifidobacterium adolescentis		91.48	0.38	MNS 87	99		
Actinomyces israelii		5.04	0.22	MLT 99	99		
Fusobacterium mortiferum		3.48	0.17	ARA 87	1		
43655122	**Low discrimination**						**43655122**
Parabacteroides distasonis		65.61	0.38	MNS 98			
Prevotella oralis		20.17	0.16				
Bacteroides uniformis		4.09	-.02				
43655322	**Acceptable identification**						**43655322**
Parabacteroides distasonis		92.25	0.46	MNS 98			
Prevotella oralis		4.05	0.10				
Bacteroides uniformis		2.65	0.01				
43655332	**Good identification**						**43655332**
Parabacteroides distasonis		100.0	0.41	MNS 98			
43655502	**Acceptable identification**						**43655502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium adolescentis		92.86	0.50	MNS 87	1		
Actinomyces israelii		5.11	0.35	MLT 99	1		
Clostridium butyricum		1.39	0.04		99		
43655512	**Good identification**						**43655512**
Bifidobacterium adolescentis		99.33	0.52	MNS 87			
Actinomyces israelii		0.67	0.21	MLT 99			
43655552	**Good identification**						**43655552**
Bifidobacterium adolescentis		97.93	0.36	MNS 87			
Actinomyces israelii		2.07	0.14	MLT 99			
43655722	**Low discrimination**						**43655722**
Parabacteroides distasonis		63.73	0.42	MNS 98			
Bacteroides uniformis		34.31	0.19				
Clostridium clostridioforme		1.00	0.06				
43655732	**Low discrimination**						**43655732**
Parabacteroides distasonis		88.29	0.38	MNS 98			
Clostridium clostridioforme		11.71	0.17				
43657000	**Low discrimination**						**43657000**
					GRA	SPO	
Actinomyces israelii		67.59	0.36	MLT 99	99	1	
Fusobacterium mortiferum		28.50	0.27		1	1	
Clostridium innocuum		3.92	0.00		99	99	
43657002	**Acceptable identification**						**43657002**
					GRA	SPO	
Actinomyces israelii		93.47	0.39	MLT 99	99	1	
Clostridium innocuum		3.61	0.00		99	99	
Fusobacterium mortiferum		2.92	0.11		1	1	
43657100	**Low discrimination**						**43657100**
					GRA		
Actinomyces israelii		80.26	0.46	MLT 99	99		
Fusobacterium mortiferum		19.74	0.34		1		
43657102	**Low discrimination**						**43657102**
					GRA		
Actinomyces israelii		74.69	0.49	MLT 99	99		
Bifidobacterium adolescentis		23.95	0.35		99		
Fusobacterium mortiferum		1.36	0.17		1		
43657112	**Low discrimination**						**43657112**
Bifidobacterium adolescentis		72.40	0.36	MNS 87			
Actinomyces israelii		27.61	0.35	MLT 99			
43657140	**Good identification**						**43657140**
Actinomyces israelii		100.0	0.39	MLT 99			
43657142	**Acceptable identification**						**43657142**
Actinomyces israelii		90.75	0.42	MLT 99			
Bifidobacterium adolescentis		9.25	0.19				
43657402	**Acceptable identification**						**43657402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		90.42	0.38	MLT 99		
Bifidobacterium adolescentis		9.58	0.15			
43657500	**Acceptable identification**					**43657500**
					SPO	
Actinomyces israelii		94.32	0.45	MLT 99	1	
Bifidobacterium adolescentis		3.75	0.15		1	
Clostridium butyricum		1.93	-.05		99	
43657502	**Low discrimination**					**43657502**
Bifidobacterium adolescentis		72.40	0.49	MNS 87		
Actinomyces israelii		27.61	0.48	MLT 99		
43657512	**Good identification**					**43657512**
Bifidobacterium adolescentis		95.07	0.50	MNS 87		
Actinomyces israelii		4.43	0.34	MLT 99		
Bifidobacterium dentium		0.50	-.07			
43657540	**Good identification**					**43657540**
Actinomyces israelii		100.0	0.38	MLT 99		
43657542	**Low discrimination**					**43657542**
Actinomyces israelii		54.54	0.41	MLT 99		
Bifidobacterium adolescentis		45.46	0.33			
43660100	**Good identification**					**43660100**
Fusobacterium mortiferum		100.0	0.41	FRU 99		
43661000	**Acceptable identification**					**43661000**
					GRA	
Eubacterium tortuosum		94.08	0.41	TRE 1	99	
Fusobacterium mortiferum		5.92	0.38	CEL 10	1	
43661100	**Acceptable identification**					**43661100**
					GRA	
Fusobacterium mortiferum		93.56	0.44	FRU 99	1	
Eubacterium tortuosum		6.44	0.07		99	
43670000	**Very good identification**					**43670000**
Fusobacterium mortiferum		100.0	0.69			
43670002	**Good identification**					**43670002**
Fusobacterium mortiferum		100.0	0.52	BGL 10		
43670100	**Very good identification**					**43670100**
Fusobacterium mortiferum		100.0	0.75			
43670101	**Good identification**					**43670101**
Fusobacterium mortiferum		100.0	0.41	URE 1		
43670102	**Very good identification**					**43670102**
Fusobacterium mortiferum		100.0	0.59	BGL 10		
43670110	**Good identification**					**43670110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.41	MLZ 1		
43670120	**Good identification**					**43670120**
Fusobacterium mortiferum		100.0	0.41	NAG 1		
43670140	**Good identification**					**43670140**
Fusobacterium mortiferum		100.0	0.41	SOR 1		
43670300	**Good identification**					**43670300**
Fusobacterium mortiferum		100.0	0.41	RHA 1		
43670500	**Good identification**					**43670500**
Fusobacterium mortiferum		100.0	0.41	ARA 1		
43671000	**Very good identification**					**43671000**
					GRA	
Fusobacterium mortiferum		99.57	0.72		1	
Eubacterium tortuosum		0.16	0.07		99	
Actinomyces naeslundii		0.14	0.34		99	
43671001	**Low discrimination**					**43671001**
					GRA	
Fusobacterium mortiferum		83.09	0.38	URE 1	1	
Actinomyces naeslundii		16.91	0.37		99	
43671002	**Good identification**					**43671002**
					GRA	
Fusobacterium mortiferum		96.55	0.55	BGL 10	1	
Actinomyces naeslundii		1.19	0.34		99	
Lactobacillus acidophilus		1.02	0.09		99	
43671003	**Low discrimination**					**43671003**
					GRA	
Actinomyces naeslundii		64.69	0.37	ESL 12	99	
Fusobacterium mortiferum		35.31	0.21	MNS 88	1	
43671010	**Good identification**					**43671010**
					GRA	
Fusobacterium mortiferum		97.02	0.38	MLZ 1	1	
Lactobacillus jensenii		2.98	0.02		99	
43671020	**Low discrimination**					**43671020**
					GRA	
Fusobacterium mortiferum		85.91	0.38	NAG 1	1	
Actinomyces naeslundii		11.66	0.34		99	
Eubacterium tortuosum		2.44	-.06		99	
43671021	**Good identification**					**43671021**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43671023	**Good identification**					**43671023**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43671040	**Good identification**					**43671040**
Fusobacterium mortiferum		100.0	0.38	SOR 1		
43671100	**Very good identification**					**43671100**
					GRA	
Fusobacterium mortiferum		99.77	0.78		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.17	0.42		99		
Actinomyces israelii		0.06	0.25		99		
43671101	**Low discrimination**						**43671101**
					GRA		
Fusobacterium mortiferum		80.11	0.44	URE 1	1		
Actinomyces naeslundii		19.89	0.45		99		
43671102	**Good identification**						**43671102**
					GRA		
Fusobacterium mortiferum		91.58	0.62	BGL 10	1		
Lactobacillus acidophilus		4.77	0.27		99		
Actinomyces naeslundii		1.38	0.42		99		
43671103	**Low discrimination**						**43671103**
					GRA		
Actinomyces naeslundii		69.09	0.45	ESL 12	99		
Fusobacterium mortiferum		30.91	0.27	MNS 88	1		
43671110	**Good identification**						**43671110**
Fusobacterium mortiferum		98.63	0.44	MLZ 1			
Leptotrichia buccalis		1.37	-.11				
43671120	**Low discrimination**						**43671120**
					GRA	SPO	
Fusobacterium mortiferum		77.69	0.44	NAG 1	1	1	
Actinomyces naeslundii		12.86	0.42		99	1	
Clostridium paraputrificum		4.74	0.03		99	99	
43671121	**Good identification**						**43671121**
Actinomyces naeslundii		100.0	0.45	ESL 12			
43671123	**Good identification**						**43671123**
Actinomyces naeslundii		100.0	0.45	ESL 12			
43671140	**Good identification**						**43671140**
					GRA		
Fusobacterium mortiferum		95.04	0.44	SOR 1	1		
Actinomyces naeslundii		2.78	0.29		99		
Actinomyces israelii		2.19	0.18		99		
43671200	**Good identification**						**43671200**
Fusobacterium mortiferum		100.0	0.38	RHA 1			
43671300	**Good identification**						**43671300**
Fusobacterium mortiferum		100.0	0.44	RHA 1			
43671322	**Low discrimination**						**43671322**
					SPO	GRA	
Parabacteroides distasonis		72.88	0.41	MNS 98	1	1	
Clostridium ramosum		19.94	0.19	XYL 90	92	50	
Lactobacillus acidophilus		3.92	0.08		1	99	
43671332	**Good identification**						**43671332**
Parabacteroides distasonis		100.0	0.36	MNS 98			
43671400	**Good identification**						**43671400**
					GRA		
Fusobacterium mortiferum		97.25	0.38	ARA 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		2.75	0.13		99		
43671500	**Low discrimination**						**43671500**
					GRA	SPO	
Fusobacterium mortiferum		89.02	0.44	ARA 1	1	1	
Clostridium butyricum		6.67	0.06		99	99	
Actinomyces israelii		4.31	0.24		99	1	
43671722	**Good identification**						**43671722**
Parabacteroides distasonis		100.0	0.37	MNS 98			
43672100	**Good identification**						**43672100**
					GRA		
Fusobacterium mortiferum		98.03	0.41	MAN 1	1		
Actinomyces naeslundii		1.97	0.23		99		
43673000	**Low discrimination**						**43673000**
					GRA	SPO	
Fusobacterium mortiferum		59.16	0.38	MAN 1	1	1	
Actinomyces naeslundii		15.59	0.39		99	1	
Clostridium innocuum		13.69	0.14		99	99	
43673001	**Good identification**						**43673001**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43673003	**Good identification**						**43673003**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43673020	**Good identification**						**43673020**
Actinomyces naeslundii		100.0	0.39	ESL 12			
43673021	**Good identification**						**43673021**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43673022	**Good identification**						**43673022**
Actinomyces naeslundii		100.0	0.39	ESL 12			
43673023	**Good identification**						**43673023**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43673100	**Low discrimination**						**43673100**
					GRA		
Fusobacterium mortiferum		60.37	0.44	MAN 1	1		
Actinomyces israelii		20.23	0.38		99		
Actinomyces naeslundii		19.40	0.47		99		
43673101	**Good identification**						**43673101**
Actinomyces naeslundii		100.0	0.50	ESL 12			
43673102	**Low discrimination**						**43673102**
					GRA		
Actinomyces israelii		44.71	0.41	MLT 99	99		
Actinomyces naeslundii		28.58	0.47	XYL 92	99		
Fusobacterium mortiferum		9.88	0.27		1		
43673103	**Good identification**						**43673103**
Actinomyces naeslundii		100.0	0.50	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43673120	**Low discrimination**						**43673120**
					SPO		
Actinomyces naeslundii		53.87	0.47	ESL 12	1		
Clostridium ramosum		39.30	0.19	MNS 88	92		
Actinomyces israelii		3.59	0.18	MLT 94	1		
43673121	**Good identification**						**43673121**
Actinomyces naeslundii		100.0	0.50	ESL 12			
43673123	**Good identification**						**43673123**
Actinomyces naeslundii		100.0	0.50	ESL 12			
43673141	**Good identification**						**43673141**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43673143	**Good identification**						**43673143**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43673161	**Good identification**						**43673161**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43673163	**Good identification**						**43673163**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43673500	**Good identification**						**43673500**
Actinomyces israelii		100.0	0.37	MLT 99			
43673502	**Low discrimination**						**43673502**
Actinomyces israelii		55.60	0.40	MLT 99			
Bifidobacterium adolescentis		44.40	0.32	XYL 92			
43674000	**Good identification**						**43674000**
Fusobacterium mortiferum		100.0	0.49	XYL 7			
43674100	**Good identification**						**43674100**
Fusobacterium mortiferum		100.0	0.56	XYL 7			
43674102	**Good identification**						**43674102**
Fusobacterium mortiferum		100.0	0.39	XYL 7			
43675000	**Acceptable identification**						**43675000**
					GRA		
Fusobacterium mortiferum		94.94	0.52	XYL 7	1		
Actinomyces israelii		5.06	0.33		99		
43675002	**Low discrimination**						**43675002**
					GRA		
Fusobacterium mortiferum		58.18	0.36	XYL 7	1		
Actinomyces israelii		41.82	0.36	BGL 10	99		
43675100	**Low discrimination**						**43675100**
					GRA	SPO	
Fusobacterium mortiferum		88.70	0.59	XYL 7	1	1	
Actinomyces israelii		8.10	0.43		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		3.21	0.15		99	99	
43675102	**Low discrimination**						**43675102**
					GRA		
Actinomyces israelii		42.89	0.46	MLT 99	99		
Fusobacterium mortiferum		34.81	0.42		1		
Bifidobacterium adolescentis		16.88	0.33		99		
43675122	**Good identification**						**43675122**
					GRA		
Parabacteroides distasonis		97.36	0.49	MNS 98	1		
Actinomyces israelii		2.64	0.26		99		
43675132	**Good identification**						**43675132**
Parabacteroides distasonis		100.0	0.44	MNS 98			
43675140	**Low discrimination**						**43675140**
					GRA	SPO	
Actinomyces israelii		72.83	0.36	MLT 99	99	1	
Fusobacterium mortiferum		20.72	0.25		1	1	
Clostridium butyricum		6.45	-.03		99	99	
43675142	**Low discrimination**						**43675142**
Actinomyces israelii		88.88	0.39	MLT 99			
Bifidobacterium adolescentis		11.12	0.18				
43675322	**Very good identification**						**43675322**
Parabacteroides distasonis		100.0	0.57	MNS 98			
43675332	**Good identification**						**43675332**
Parabacteroides distasonis		99.17	0.53	MNS 98			
Clostridium clostridioforme		0.83	0.11				
43675502	**Low discrimination**						**43675502**
					SPO		
Bifidobacterium adolescentis		59.78	0.47	MNS 87	1		
Clostridium butyricum		20.98	0.25	MLT 99	99		
Actinomyces israelii		18.57	0.45		1		
43675512	**Acceptable identification**						**43675512**
Bifidobacterium adolescentis		92.53	0.49	MNS 87			
Actinomyces israelii		3.52	0.31	MLT 99			
Clostridium clostridioforme		1.95	0.17				
43675522	**Low discrimination**						**43675522**
					GRA	SPO	
Parabacteroides distasonis		89.25	0.45	MNS 98	1	1	
Clostridium butyricum		4.96	0.07		99	99	
Actinomyces israelii		3.23	0.24		99	1	
43675532	**Low discrimination**						**43675532**
					GRA		
Parabacteroides distasonis		84.97	0.41	MNS 98	1		
Clostridium clostridioforme		7.42	0.17		50		
Bifidobacterium dentium		4.06	-.03		99		
43675722	**Good identification**						**43675722**
Parabacteroides distasonis		95.73	0.53	MNS 98			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		2.42	0.20				
Bacteroides ovatus		1.36	0.06				
43675732	**Low discrimination**						**43675732**
Parabacteroides distasonis		81.23	0.49	MNS 98			
Clostridium clostridioforme		17.34	0.32				
Bacteroides ovatus		1.43	0.03				
43677000	**Good identification**						**43677000**
					GRA		
Actinomyces israelii		96.76	0.46	MLT 99	99		
Fusobacterium mortiferum		3.24	0.18		1		
43677002	**Good identification**						**43677002**
Actinomyces israelii		100.0	0.49	MLT 99			
43677040	**Good identification**						**43677040**
Actinomyces israelii		100.0	0.39	MLT 99			
43677042	**Good identification**						**43677042**
Actinomyces israelii		100.0	0.42	MLT 99			
43677100	**Very good identification**						**43677100**
					GRA	SPO	
Actinomyces israelii		97.34	0.56	MLT 99	99	1	
Fusobacterium mortiferum		1.90	0.25		1	1	
Clostridium butyricum		0.76	-.01		99	99	
43677102	**Good identification**						**43677102**
Actinomyces israelii		94.62	0.59	MLT 99			
Bifidobacterium adolescentis		5.38	0.32				
43677110	**Good identification**						**43677110**
Actinomyces israelii		100.0	0.42	MLT 99			
43677112	**Low discrimination**						**43677112**
Actinomyces israelii		67.14	0.45	MLT 99			
Bifidobacterium adolescentis		31.20	0.33	MLZ 13			
Bifidobacterium dentium		1.66	-.07				
43677120	**Good identification**						**43677120**
Actinomyces israelii		100.0	0.36	MLT 99			
43677122	**Acceptable identification**						**43677122**
					GRA		
Actinomyces israelii		93.83	0.39	MLT 99	99		
Parabacteroides distasonis		6.18	0.15	NAG 6	1		
43677140	**Good identification**						**43677140**
Actinomyces israelii		100.0	0.49	MLT 99			
43677142	**Good identification**						**43677142**
Actinomyces israelii		98.23	0.52	MLT 99			
Bifidobacterium adolescentis		1.77	0.16				
43677150	**Acceptable identification**						**43677150**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Actinomyces israelii		91.41	0.35	MLT 99	1	
Blautia producta		8.59	-.03	MLZ 13	99	
43677152	**Low discrimination**					**43677152**
Actinomyces israelii		87.13	0.38	MLT 99		
Bifidobacterium adolescentis		12.87	0.18	MLZ 13		
43677300	**Good identification**					**43677300**
Actinomyces israelii		100.0	0.37	MLT 99		
43677302	**Good identification**					**43677302**
Actinomyces israelii		100.0	0.40	MLT 99		
43677400	**Good identification**					**43677400**
Actinomyces israelii		100.0	0.44	MLT 99		
43677402	**Good identification**					**43677402**
Actinomyces israelii		98.16	0.47	MLT 99		
Bifidobacterium adolescentis		1.84	0.12			
43677440	**Good identification**					**43677440**
Actinomyces israelii		100.0	0.37	MLT 99		
43677442	**Good identification**					**43677442**
Actinomyces israelii		100.0	0.40	MLT 99		
43677500	**Acceptable identification**					**43677500**
					SPO	
Actinomyces israelii		91.54	0.55	MLT 99	1	
Clostridium butyricum		7.82	0.15		99	
Bifidobacterium adolescentis		0.65	0.12		1	
43677502	**Low discrimination**					**43677502**
					SPO	
Actinomyces israelii		67.05	0.58	MLT 99	1	
Bifidobacterium adolescentis		31.16	0.46		1	
Clostridium butyricum		1.49	0.08		99	
43677510	**Low discrimination**					**43677510**
Actinomyces israelii		63.46	0.41	MLT 99		
Bifidobacterium dentium		28.15	0.10	MLZ 13		
Bifidobacterium adolescentis		3.66	0.13			
43677512	**Low discrimination**					**43677512**
Bifidobacterium adolescentis		48.79	0.47	MNS 87		
Bifidobacterium dentium		38.37	0.27	MLT 99		
Actinomyces israelii		12.84	0.44			
43677522	**Low discrimination**					**43677522**
Actinomyces israelii		80.09	0.37	MLT 99		
Bifidobacterium dentium		10.07	-.03	NAG 6		
Bifidobacterium adolescentis		5.89	0.12			
43677540	**Low discrimination**					**43677540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		63.08	0.48	MLT 99	1	1	
Blautia producta		35.72	0.24		99	1	
Clostridium butyricum		1.21	-.03		1	99	
43677542	**Low discrimination**						**43677542**
					COC		
Actinomyces israelii		84.06	0.51	MLT 99	1		
Bifidobacterium adolescentis		12.42	0.30		1		
Blautia producta		3.53	0.07		99		
43677700	**Good identification**						**43677700**
Actinomyces israelii		100.0	0.35	MLT 99			
43677702	**Acceptable identification**						**43677702**
Actinomyces israelii		94.13	0.39	MLT 99			
Bifidobacterium adolescentis		5.87	0.12	RHA 7			
43700100	**Good identification**						**43700100**
					COC	GRA	
Fusobacterium mortiferum		97.59	0.42	MLT 11	1	1	
Blautia hansenii		2.42	-.03	FRU 99	99	99	
43701000	**Very good identification**						**43701000**
					GRA		
Eubacterium tortuosum		99.05	0.58	MLT 1	99		
Fusobacterium mortiferum		0.86	0.39		1		
Actinomyces naeslundii		0.03	0.26		99		
43701002	**Good identification**						**43701002**
					GRA		
Eubacterium tortuosum		99.18	0.45	MLT 1	99		
Fusobacterium mortiferum		0.54	0.23		1		
Actinomyces naeslundii		0.17	0.26		99		
43701020	**Good identification**						**43701020**
					GRA		
Eubacterium tortuosum		99.67	0.45	MLT 1	99		
Actinomyces naeslundii		0.17	0.26		99		
Prevotella melaninogenica		0.16	-.04		1		
43701101	**Low discrimination**						**43701101**
					COC	SPO	
Actinomyces naeslundii		56.12	0.37	ESL 12	1	1	
Clostridium chauvoei		20.60	0.06	MNS 88	1	99	
Blautia hansenii		14.44	-.03	FRU 95	99	1	
43701103	**Low discrimination**						**43701103**
Actinomyces naeslundii		87.03	0.37	ESL 12			
Bifidobacterium longum subsp. infantis		12.97	-.07	MNS 88			
43701121	**Low discrimination**						**43701121**
					GRA		
Actinomyces naeslundii		82.13	0.37	ESL 12	99		
Prevotella melaninogenica		17.87	-.04	MNS 88	1		
43701123	**Low discrimination**						**43701123**
Actinomyces naeslundii		79.05	0.37	ESL 12			
Bifidobacterium longum subsp. infantis		20.95	-.03	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43703100	**Low discrimination**						**43703100**
					COC	GRA	
Actinomyces naeslundii		75.72	0.38	ESL 12	1	99	
Blautia hansenii		15.06	-.03	MNS 88	99	99	
Fusobacterium mortiferum		9.22	0.11	FRU 95	1	1	
43703101	**Good identification**						**43703101**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43703102	**Low discrimination**						**43703102**
Actinomyces naeslundii		68.89	0.38	ESL 12			
Bifidobacterium adolescentis		23.18	0.15	MNS 88			
Bifidobacterium longum subsp. infantis		7.93	-.07	FRU 95			
43703103	**Good identification**						**43703103**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43703120	**Low discrimination**						**43703120**
					GRA		
Actinomyces naeslundii		85.61	0.38	ESL 12	99		
Prevotella melaninogenica		14.40	-.04	MNS 88	1		
43703121	**Good identification**						**43703121**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43703122	**Low discrimination**						**43703122**
Actinomyces naeslundii		83.01	0.38	ESL 12			
Bifidobacterium longum subsp. infantis		17.00	-.03	MNS 88			
43703123	**Good identification**						**43703123**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43705300	**Very good identification**						**43705300**
Eubacterium contortum		100.0	0.61	FRU 99			
43705302	**Good identification**						**43705302**
Eubacterium contortum		100.0	0.48	FRU 99			
43705320	**Good identification**						**43705320**
					GRA		
Eubacterium contortum		99.48	0.48	FRU 99	99		
Bacteroides vulgatus		0.52	0.02		1		
43705322	**Acceptable identification**						**43705322**
					GRA		
Eubacterium contortum		92.61	0.35	FRU 99	99		
Parabacteroides distasonis		5.47	0.26		1		
Prevotella oralis		1.02	0.00		1		
43705502	**Acceptable identification**						**43705502**
Bifidobacterium adolescentis		90.46	0.45	MNS 87			
Bifidobacterium longum subsp. longum		4.77	0.21	FRU 99			
Eubacterium contortum		3.98	0.13				
43705512	**Acceptable identification**						**43705512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		93.16	0.46	MNS 87		
Bifidobacterium longum subsp. longum		6.84	0.25	FRU 99		
43705530	**Good identification**					**43705530**
Bifidobacterium longum subsp. longum		100.0	0.35	MNS 87		
43705532	**Good identification**					**43705532**
Bifidobacterium longum subsp. longum		96.67	0.35	MNS 87		
Bifidobacterium adolescentis		3.33	0.12	SAL 1		
43705700	**Very good identification**					**43705700**
Eubacterium contortum		100.0	0.60	FRU 99		
43705702	**Good identification**					**43705702**
Eubacterium contortum		99.77	0.47	FRU 99		
Bifidobacterium adolescentis		0.23	0.11			
43705720	**Acceptable identification**					**43705720**
					GRA	
Eubacterium contortum		92.16	0.47	FRU 99	99	
Bacteroides vulgatus		7.84	0.21		1	
43707502	**Acceptable identification**					**43707502**
Bifidobacterium adolescentis		94.25	0.43	MNS 87		
Actinomyces israelii		5.75	0.29	FRU 99		
43707512	**Good identification**					**43707512**
Bifidobacterium adolescentis		99.26	0.45	MNS 87		
Actinomyces israelii		0.74	0.15	FRU 99		
43710000	**Very good identification**					**43710000**
					GRA	
Fusobacterium mortiferum		99.91	0.70	MLT 11	1	
Prevotella intermedia		0.05	0.02		1	
Actinomyces naeslundii		0.04	0.24		99	
43710001	**Acceptable identification**					**43710001**
					GRA	
Fusobacterium mortiferum		94.13	0.36	MLT 11	1	
Actinomyces naeslundii		5.87	0.27	URE 1	99	
43710002	**Good identification**					**43710002**
					GRA	
Fusobacterium mortiferum		99.32	0.54	MLT 11	1	
Actinomyces naeslundii		0.38	0.24	BGL 10	99	
Bifidobacterium longum subsp. infantis		0.30	-.07		99	
43710010	**Good identification**					**43710010**
Fusobacterium mortiferum		100.0	0.36	MLT 11		
43710020	**Good identification**					**43710020**
					GRA	
Fusobacterium mortiferum		96.01	0.36	MLT 11	1	
Actinomyces naeslundii		3.99	0.24	NAG 1	99	
43710040	**Good identification**					**43710040**
Fusobacterium mortiferum		100.0	0.36	MLT 11		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43710100	**Very good identification**					**43710100**
					GRA	
Fusobacterium mortiferum		99.81	0.77	MLT 11	1	
Bifidobacterium longum subsp. infantis		0.14	0.10		99	
Actinomyces naeslundii		0.05	0.32		99	
43710101	**Acceptable identification**					**43710101**
					GRA	
Fusobacterium mortiferum		92.93	0.42	MLT 11	1	
Actinomyces naeslundii		7.07	0.35	URE 1	99	
43710102	**Low discrimination**					**43710102**
					GRA	
Fusobacterium mortiferum		87.59	0.60	MLT 11	1	
Bifidobacterium longum subsp. infantis		11.30	0.27	BGL 10	99	
Bifidobacterium adolescentis		0.41	0.16		99	
43710110	**Good identification**					**43710110**
Fusobacterium mortiferum		100.0	0.42	MLT 11		
43710120	**Low discrimination**					**43710120**
					GRA	
Fusobacterium mortiferum		66.14	0.42	MLT 11	1	
Bifidobacterium longum subsp. infantis		16.50	0.15	NAG 1	99	
Prevotella melaninogenica		12.45	0.12		1	
43710140	**Good identification**					**43710140**
Fusobacterium mortiferum		100.0	0.42	MLT 11		
43710200	**Good identification**					**43710200**
					GRA	
Fusobacterium mortiferum		96.47	0.36	MLT 11	1	
Actinomyces odontolyticus		3.53	0.06	RHA 1	99	
43710300	**Low discrimination**					**43710300**
					GRA	
Fusobacterium mortiferum		58.87	0.42	MLT 11	1	
Eubacterium contortum		41.13	0.27	RHA 1	99	
43710400	**Good identification**					**43710400**
Fusobacterium mortiferum		100.0	0.36	MLT 11		
43710500	**Good identification**					**43710500**
Fusobacterium mortiferum		100.0	0.42	MLT 11		
43711000	**Very good identification**					**43711000**
					GRA	
Fusobacterium mortiferum		97.58	0.73	MLT 11	1	
Eubacterium tortuosum		1.15	0.24		99	
Actinomyces naeslundii		0.66	0.48		99	
43711001	**Low discrimination**					**43711001**
					GRA	
Fusobacterium mortiferum		50.04	0.39	MLT 11	1	
Actinomyces naeslundii		49.96	0.51	URE 1	99	
43711002	**Low discrimination**					**43711002**
					GRA	
Fusobacterium mortiferum		75.05	0.57	MLT 11	1	
Bifidobacterium longum subsp. infantis		15.06	0.27	BGL 10	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		4.54	0.48		99	
43711003	**Low discrimination**					**43711003**
					GRA	
Actinomyces naeslundii		88.22	0.51	ESL 12	99	
Fusobacterium mortiferum		9.82	0.23	MNS 88	1	
Bifidobacterium longum subsp. infantis		1.97	-.07		99	
43711010	**Low discrimination**					**43711010**
					GRA	
Fusobacterium mortiferum		89.08	0.39	MLT 11	1	
Eubacterium saburreum		7.80	0.19	MLZ 1	85	
Actinomyces naeslundii		3.12	0.26		99	
43711020	**Low discrimination**					**43711020**
					GRA	
Fusobacterium mortiferum		33.55	0.39	MLT 11	1	
Actinomyces naeslundii		22.33	0.48	NAG 1	99	
Bifidobacterium longum subsp. infantis		13.02	0.15		99	
43711021	**Good identification**					**43711021**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43711022	**Low discrimination**					**43711022**
					GRA	
Prevotella oralis		46.34	0.36	MNS 99	1	
Bifidobacterium longum subsp. infantis		41.33	0.32	LAC 94	99	
Actinomyces naeslundii		7.01	0.48		99	
43711023	**Acceptable identification**					**43711023**
					GRA	
Actinomyces naeslundii		92.24	0.51	ESL 12	99	
Prevotella oralis		4.10	0.02	MNS 88	1	
Bifidobacterium longum subsp. infantis		3.66	-.03		99	
43711040	**Low discrimination**					**43711040**
					GRA	
Fusobacterium mortiferum		89.49	0.39	MLT 11	1	
Actinomyces naeslundii		10.51	0.35	SOR 1	99	
43711041	**Good identification**					**43711041**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43711043	**Good identification**					**43711043**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43711061	**Good identification**					**43711061**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43711063	**Good identification**					**43711063**
Actinomyces naeslundii		100.0	0.38	ESL 12		
43711100	**Low discrimination**					**43711100**
					GRA	
Fusobacterium mortiferum		89.76	0.80	MLT 11	1	
Bifidobacterium longum subsp. infantis		8.40	0.44		99	
Actinomyces naeslundii		0.74	0.55		99	
43711101	**Low discrimination**					**43711101**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		51.47	0.58	ESL 12	99		
Fusobacterium mortiferum		42.26	0.45	MNS 88	1		
Bifidobacterium longum subsp. infantis		3.95	0.10		99		
43711102	**Low discrimination**						**43711102**
					GRA		
Bifidobacterium longum subsp. infantis		83.78	0.62	MNS 99	99		
Fusobacterium mortiferum		9.84	0.63		1		
Bifidobacterium adolescentis		3.02	0.51		99		
43711103	**Low discrimination**						**43711103**
					GRA		
Actinomyces naeslundii		52.17	0.58	ESL 12	99		
Bifidobacterium longum subsp. infantis		40.52	0.27	MNS 88	99		
Fusobacterium mortiferum		4.76	0.29		1		
43711110	**Low discrimination**						**43711110**
					GRA		
Fusobacterium mortiferum		76.75	0.45	MLT 11	1		
Bifidobacterium longum subsp. infantis		7.18	0.10	MLZ 1	99		
Eubacterium saburreum		6.72	0.26		85		
43711111	**Low discrimination**						**43711111**
					GRA		
Actinomyces naeslundii		86.39	0.37	ESL 12	99		
Fusobacterium mortiferum		13.61	0.11	MNS 88	1		
43711112	**Low discrimination**						**43711112**
Bifidobacterium adolescentis		74.83	0.52	MNS 87			
Bifidobacterium longum subsp. infantis		17.15	0.27	XYL 87			
Bifidobacterium breve		2.41	0.27	ARA 87			
43711113	**Low discrimination**						**43711113**
Actinomyces naeslundii		55.57	0.37	ESL 12			
Bifidobacterium adolescentis		36.15	0.18	MNS 88			
Bifidobacterium longum subsp. infantis		8.28	-.07	MLZ 5			
43711120	**Low discrimination**						**43711120**
					GRA		
Bifidobacterium longum subsp. infantis		48.59	0.49	MNS 99	99		
Prevotella melaninogenica		36.68	0.47	BGL 91	1		
Prevotella oralis		4.59	0.35		1		
43711121	**Low discrimination**						**43711121**
					GRA		
Actinomyces naeslundii		78.38	0.58	ESL 12	99		
Bifidobacterium longum subsp. infantis		10.70	0.15	MNS 88	99		
Prevotella melaninogenica		8.08	0.12		1		
43711122	**Low discrimination**						**43711122**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		80.13	0.66	MNS 99	99	1	
Prevotella oralis		14.22	0.56		1	1	
Capnocytophaga ochracea		3.59	0.47		1	90	
43711131	**Good identification**						**43711131**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43711133	**Low discrimination**						**43711133**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		79.05	0.37	ESL 12			
Bifidobacterium longum subsp. infantis		20.95	-.03	MNS 88			
43711140	**Low discrimination**						**43711140**
					GRA		
Fusobacterium mortiferum		77.11	0.45	MLT 11	1		
Actinomyces naeslundii		11.05	0.43	SOR 1	99		
Bifidobacterium longum subsp. infantis		7.21	0.10		99		
43711141	**Good identification**						**43711141**
					GRA		
Actinomyces naeslundii		95.51	0.46	ESL 12	99		
Fusobacterium mortiferum		4.49	0.11	MNS 88	1		
43711143	**Good identification**						**43711143**
Actinomyces naeslundii		95.75	0.46	ESL 12			
Bifidobacterium longum subsp. infantis		4.26	-.07	MNS 88			
43711152	**Low discrimination**						**43711152**
					GRA	CAT	
Bifidobacterium adolescentis		76.96	0.37	MNS 87	99	1	
Bifidobacterium breve		20.05	0.27	XYL 87	99	1	
Capnocytophaga ochracea		1.55	-.03	ARA 87	1	90	
43711161	**Good identification**						**43711161**
Actinomyces naeslundii		100.0	0.46	ESL 12			
43711163	**Acceptable identification**						**43711163**
Actinomyces naeslundii		92.68	0.46	ESL 12			
Bifidobacterium longum subsp. infantis		7.32	-.03	MNS 88			
43711300	**Very good identification**						**43711300**
					GRA		
Eubacterium contortum		97.61	0.61	XYL 99	99		
Fusobacterium mortiferum		2.12	0.45		1		
Bifidobacterium longum subsp. infantis		0.20	0.10		99		
43711302	**Low discrimination**						**43711302**
					GRA		
Eubacterium contortum		82.15	0.48	XYL 99	99		
Bifidobacterium longum subsp. infantis		9.55	0.27		99		
Prevotella oralis		6.73	0.28		1		
43711320	**Low discrimination**						**43711320**
					GRA		
Eubacterium contortum		88.53	0.48	XYL 99	99		
Prevotella oralis		7.25	0.28		1		
Bifidobacterium longum subsp. infantis		1.81	0.15		99		
43711322	**Low discrimination**						**43711322**
					GRA		
Prevotella oralis		75.26	0.50	MNS 99	1		
Bifidobacterium longum subsp. infantis		10.00	0.32		99		
Eubacterium contortum		8.54	0.35		99		
43711332	**Low discrimination**						**43711332**
					GRA		
Parabacteroides distasonis		74.75	0.39	MNS 98	1		
Prevotella oralis		19.02	0.16	XYL 90	1		
Bifidobacterium longum subsp. infantis		2.53	-.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43711400	**Low discrimination**						**43711400**
					GRA		
Fusobacterium mortiferum		81.32	0.39	MLT 11	1		
Lactobacillus fermentum		15.02	0.28	ARA 1	99		
Actinomyces israelii		3.66	0.18		99		
43711500	**Low discrimination**						**43711500**
					GRA		
Fusobacterium mortiferum		46.15	0.45	MLT 11	1		
Eubacterium contortum		17.59	0.26	ARA 1	99		
Bifidobacterium longum subsp. longum		11.15	0.29		99		
43711502	**Good identification**						**43711502**
Bifidobacterium adolescentis		92.73	0.65	MNS 87			
Bifidobacterium longum subsp. infantis		3.88	0.27	XYL 87			
Bifidobacterium longum subsp. longum		0.99	0.29				
43711512	**Very good identification**						**43711512**
Bifidobacterium adolescentis		98.30	0.66	MNS 87			
Bifidobacterium longum subsp. longum		1.46	0.33	XYL 87			
Clostridium clostridioforme		0.08	0.10				
43711520	**Low discrimination**						**43711520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		57.94	0.40	MNS 87	99	1	
Bifidobacterium longum subsp. infantis		9.97	0.15	SAL 1	99	1	
Capnocytophaga ochracea		8.78	0.17		1	90	
43711530	**Good identification**						**43711530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.02	0.44	MNS 87	99	1	
Capnocytophaga ochracea		0.98	0.01	SAL 1	1	90	
43711532	**Low discrimination**						**43711532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.36	0.44	MNS 87	99	1	
Bifidobacterium adolescentis		12.61	0.32	SAL 1	99	1	
Capnocytophaga ochracea		6.62	0.17		1	90	
43711542	**Good identification**						**43711542**
					GRA	CAT	
Bifidobacterium adolescentis		97.53	0.49	MNS 87	99	1	
Actinomyces israelii		1.57	0.25	XYL 87	99	1	
Capnocytophaga ochracea		0.57	0.01	SOR 11	1	90	
43711552	**Good identification**						**43711552**
Bifidobacterium adolescentis		100.0	0.51	MNS 87			
43711700	**Very good identification**						**43711700**
					GRA		
Eubacterium contortum		99.94	0.60	XYL 99	99		
Bacteroides vulgatus		0.03	-.08		1		
Fusobacterium mortiferum		0.03	0.11		1		
43711702	**Good identification**						**43711702**
Eubacterium contortum		96.05	0.47	XYL 99			
Bifidobacterium adolescentis		3.29	0.31				
Clostridium clostridioforme		0.52	0.14				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43711720	**Low discrimination**					**43711720**
					GRA	
Eubacterium contortum		85.22	0.47	XYL 99	99	
Bacteroides vulgatus		14.66	0.27		1	
Bifidobacterium longum subsp. longum		0.13	0.05		99	
43711732	**Low discrimination**					**43711732**
					GRA	
Parabacteroides distasonis		61.63	0.35	MNS 98	1	
Clostridium clostridioforme		32.67	0.25	XYL 90	50	
Bifidobacterium longum subsp. longum		3.33	0.09		99	
43712000	**Acceptable identification**					**43712000**
					GRA	
Fusobacterium mortiferum		92.53	0.36	MLT 11	1	
Actinomyces naeslundii		7.47	0.29	MAN 1	99	
43712100	**Low discrimination**					**43712100**
					GRA	
Fusobacterium mortiferum		89.86	0.42	MLT 11	1	
Actinomyces naeslundii		8.85	0.37	MAN 1	99	
Propionibacterium propionicum		1.29	0.12		99	
43712101	**Good identification**					**43712101**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43712103	**Good identification**					**43712103**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43712120	**Good identification**					**43712120**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43712121	**Good identification**					**43712121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43712122	**Low discrimination**					**43712122**
Actinomyces naeslundii		79.45	0.37	ESL 12		
Bifidobacterium longum subsp. infantis		20.55	-.03	MNS 88		
43712123	**Good identification**					**43712123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43713000	**Low discrimination**					**43713000**
					GRA	
Actinomyces naeslundii		49.62	0.53	ESL 12	99	
Fusobacterium mortiferum		38.40	0.39	MNS 88	1	
Actinomyces israelii		11.98	0.33		99	
43713001	**Good identification**					**43713001**
Actinomyces naeslundii		100.0	0.56	ESL 12		
43713002	**Low discrimination**					**43713002**
					GRA	
Actinomyces naeslundii		65.97	0.53	ESL 12	99	
Actinomyces israelii		23.89	0.36	MNS 88	99	
Fusobacterium mortiferum		5.67	0.23		1	
43713003	**Good identification**					**43713003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.56	ESL 12		
43713020	**Good identification**					**43713020**
Actinomyces naeslundii		100.0	0.53	ESL 12		
43713021	**Good identification**					**43713021**
Actinomyces naeslundii		100.0	0.56	ESL 12		
43713022	**Acceptable identification**					**43713022**
					GRA	
Actinomyces naeslundii		91.90	0.53	ESL 12	99	
Prevotella oralis		3.16	0.02	MNS 88	1	
Bifidobacterium longum subsp. infantis		2.82	-.03		99	
43713023	**Good identification**					**43713023**
Actinomyces naeslundii		100.0	0.56	ESL 12		
43713040	**Low discrimination**					**43713040**
Actinomyces naeslundii		65.27	0.40	ESL 12		
Actinomyces israelii		34.73	0.26	MNS 88		
43713041	**Good identification**					**43713041**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43713042	**Low discrimination**					**43713042**
Actinomyces naeslundii		55.61	0.40	ESL 12		
Actinomyces israelii		44.39	0.29	MNS 88		
43713043	**Good identification**					**43713043**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43713060	**Good identification**					**43713060**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43713061	**Good identification**					**43713061**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43713062	**Good identification**					**43713062**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43713063	**Good identification**					**43713063**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43713100	**Low discrimination**					**43713100**
					GRA	
Actinomyces naeslundii		48.42	0.60	ESL 12	99	
Fusobacterium mortiferum		30.72	0.45	MNS 88	1	
Actinomyces israelii		16.43	0.43		99	
43713101	**Very good identification**					**43713101**
					GRA	
Actinomyces naeslundii		99.58	0.63	ESL 12	99	
Fusobacterium mortiferum		0.43	0.11	MNS 88	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43713102	**Low discrimination**					**43713102**
Bifidobacterium adolescentis		43.44	0.49	MNS 87		
Actinomyces naeslundii		24.78	0.60	XYL 87		
Bifidobacterium longum subsp. infantis		14.87	0.27	ARA 87		
43713103	**Very good identification**					**43713103**
Actinomyces naeslundii		98.44	0.63	ESL 12		
Bifidobacterium adolescentis		1.16	0.15	MNS 88		
Bifidobacterium longum subsp. infantis		0.40	-.07			
43713110	**Low discrimination**					**43713110**
Actinomyces naeslundii		37.67	0.38	ESL 12		
Actinomyces israelii		36.29	0.29	MNS 88		
Bifidobacterium adolescentis		15.49	0.16	MLZ 5		
43713111	**Good identification**					**43713111**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43713112	**Acceptable identification**					**43713112**
Bifidobacterium adolescentis		90.72	0.51	MNS 87		
Bifidobacterium breve		3.57	0.27	XYL 87		
Actinomyces israelii		3.22	0.32	ARA 87		
43713113	**Low discrimination**					**43713113**
Actinomyces naeslundii		78.48	0.42	ESL 12		
Bifidobacterium adolescentis		21.52	0.16	MNS 88		
43713120	**Low discrimination**					**43713120**
					GRA	
Actinomyces naeslundii		80.28	0.60	ESL 12	99	
Bifidobacterium longum subsp. infantis		8.47	0.15	MNS 88	99	
Prevotella melaninogenica		6.39	0.12		1	
43713121	**Very good identification**					**43713121**
Actinomyces naeslundii		100.0	0.63	ESL 12		
43713123	**Very good identification**					**43713123**
Actinomyces naeslundii		99.29	0.63	ESL 12		
Bifidobacterium longum subsp. infantis		0.71	-.03	MNS 88		
43713130	**Low discrimination**					**43713130**
Actinomyces naeslundii		78.02	0.38	ESL 12		
Bifidobacterium breve		21.98	0.15	MNS 88		
43713131	**Good identification**					**43713131**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43713133	**Good identification**					**43713133**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43713140	**Low discrimination**					**43713140**
Actinomyces naeslundii		53.60	0.47	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		40.08	0.36	MNS 88		
Bifidobacterium breve		2.53	0.10			
43713141	**Good identification**					**43713141**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43713143	**Good identification**					**43713143**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43713152	**Low discrimination**					**43713152**
Bifidobacterium adolescentis		68.25	0.35	MNS 87		
Bifidobacterium breve		21.73	0.27	XYL 87		
Actinomyces israelii		7.62	0.25	ARA 87		
43713160	**Low discrimination**					**43713160**
Actinomyces naeslundii		88.36	0.47	ESL 12		
Bifidobacterium breve		7.42	0.15	MNS 88		
Actinomyces israelii		4.22	0.16			
43713161	**Good identification**					**43713161**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43713162	**Low discrimination**					**43713162**
Actinomyces naeslundii		50.45	0.47	ESL 12		
Bifidobacterium breve		42.86	0.32	MNS 88		
Actinomyces israelii		3.61	0.19			
43713163	**Good identification**					**43713163**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43713500	**Low discrimination**					**43713500**
Actinomyces israelii		67.28	0.42	CEL 93		
Bifidobacterium adolescentis		28.72	0.29	XYL 92		
Actinomyces naeslundii		2.45	0.26			
43713502	**Very good identification**					**43713502**
Bifidobacterium adolescentis		96.44	0.63	MNS 87		
Actinomyces israelii		3.42	0.45	XYL 87		
Actinomyces naeslundii		0.08	0.26			
43713512	**Very good identification**					**43713512**
Bifidobacterium adolescentis		99.57	0.65	MNS 87		
Actinomyces israelii		0.43	0.31	XYL 87		
43713542	**Low discrimination**					**43713542**
Bifidobacterium adolescentis		89.95	0.48	MNS 87		
Actinomyces israelii		10.05	0.38	XYL 87		
43713552	**Good identification**					**43713552**
Bifidobacterium adolescentis		98.65	0.49	MNS 87		
Actinomyces israelii		1.35	0.24	XYL 87		
43714000	**Good identification**					**43714000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.51	MLT 11		
43714100	**Low discrimination**					**43714100**
					GRA	
Fusobacterium mortiferum		89.98	0.57	MLT 11	1	
Eubacterium contortum		8.44	0.27	XYL 7	99	
Bifidobacterium longum subsp. infantis		1.39	0.09		99	
43714300	**Very good identification**					**43714300**
					GRA	
Eubacterium contortum		99.89	0.61	GAL 99	99	
Fusobacterium mortiferum		0.11	0.23		1	
43714302	**Good identification**					**43714302**
Eubacterium contortum		100.0	0.48	GAL 99		
43714320	**Good identification**					**43714320**
					GRA	
Eubacterium contortum		99.48	0.48	GAL 99	99	
Bacteroides vulgatus		0.52	0.02		1	
43714322	**Acceptable identification**					**43714322**
					GRA	
Eubacterium contortum		92.61	0.35	GAL 99	99	
Parabacteroides distasonis		5.47	0.26		1	
Prevotella oralis		1.02	0.00		1	
43714502	**Acceptable identification**					**43714502**
Bifidobacterium adolescentis		94.02	0.45	MNS 87		
Eubacterium contortum		4.14	0.13	GAL 99		
Clostridium clostridioforme		1.01	0.08			
43714512	**Good identification**					**43714512**
Bifidobacterium adolescentis		95.52	0.46	MNS 87		
Clostridium clostridioforme		3.84	0.19	GAL 99		
Bifidobacterium longum subsp. longum		0.64	0.07			
43714700	**Very good identification**					**43714700**
Eubacterium contortum		100.0	0.60	GAL 99		
43714702	**Good identification**					**43714702**
Eubacterium contortum		98.00	0.47	GAL 99		
Clostridium clostridioforme		1.78	0.23			
Bifidobacterium adolescentis		0.23	0.11			
43714720	**Acceptable identification**					**43714720**
					GRA	
Eubacterium contortum		92.16	0.47	GAL 99	99	
Bacteroides vulgatus		7.84	0.21		1	
43715000	**Low discrimination**					**43715000**
					GRA	
Fusobacterium mortiferum		78.30	0.54	MLT 11	1	
Eubacterium contortum		11.42	0.27	XYL 7	99	
Actinomyces israelii		6.65	0.38		99	
43715100	**Low discrimination**					**43715100**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium contortum		73.62	0.61	RHA 99	99		
Bifidobacterium longum subsp. infantis		12.10	0.43		99		
Fusobacterium mortiferum		11.90	0.60		1		
43715102	**Low discrimination**						**43715102**
Bifidobacterium longum subsp. infantis		69.42	0.60	MNS 99			
Bifidobacterium adolescentis		20.48	0.65				
Eubacterium contortum		7.37	0.48				
43715112	**Very good identification**						**43715112**
Bifidobacterium adolescentis		95.96	0.66	MNS 87			
Bifidobacterium longum subsp. infantis		2.69	0.26	ARA 87			
Actinomyces israelii		0.85	0.37				
43715120	**Low discrimination**						**43715120**
					GRA		
Bifidobacterium longum subsp. infantis		58.50	0.47	MNS 99	99		
Eubacterium contortum		35.34	0.48	BGL 91	99		
Bacteroides vulgatus		2.04	0.20		1		
43715122	**Good identification**						**43715122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		90.87	0.64	MNS 99	1	99	
Prevotella oralis		2.44	0.41		1	1	
Bacteroides fragilis		2.35	0.32		99	1	
43715132	**Low discrimination**						**43715132**
					GRA		
Parabacteroides distasonis		42.63	0.47	MNS 98	1		
Bifidobacterium longum subsp. infantis		38.90	0.30		99		
Bifidobacterium adolescentis		7.89	0.32		99		
43715142	**Low discrimination**						**43715142**
Bifidobacterium adolescentis		64.88	0.49	MNS 87			
Bifidobacterium longum subsp. infantis		17.97	0.26	ARA 87			
Actinomyces israelii		14.70	0.44	SOR 11			
43715152	**Good identification**						**43715152**
Bifidobacterium adolescentis		97.30	0.51	MNS 87			
Actinomyces israelii		2.70	0.30	ARA 87			
43715200	**Very good identification**						**43715200**
					GRA		
Eubacterium contortum		99.89	0.61	LAC 99	99		
Fusobacterium mortiferum		0.07	0.20		1		
Actinomyces israelii		0.04	0.19		99		
43715202	**Good identification**						**43715202**
Eubacterium contortum		99.63	0.48	LAC 99			
Actinomyces israelii		0.38	0.22				
43715220	**Good identification**						**43715220**
					GRA		
Eubacterium contortum		99.48	0.48	LAC 99	99		
Bacteroides vulgatus		0.52	0.02		1		
43715222	**Low discrimination**						**43715222**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		87.85	0.35	LAC 99	99	
Prevotella oralis		6.11	0.14		1	
Parabacteroides distasonis		5.19	0.26		1	
43715300	**Excellent identification**					**43715300**
					GRA	
Eubacterium contortum		99.99	0.95		99	
Actinomyces israelii		0.00	0.29		99	
Fusobacterium mortiferum		0.00	0.26		1	
43715301	**Very good identification**					**43715301**
Eubacterium contortum		100.0	0.61	URE 1		
43715302	**Very good identification**					**43715302**
Eubacterium contortum		99.84	0.83			
Bifidobacterium longum subsp. infantis		0.10	0.26			
Bifidobacterium adolescentis		0.03	0.31			
43715303	**Good identification**					**43715303**
Eubacterium contortum		100.0	0.48	URE 1		
43715310	**Very good identification**					**43715310**
Eubacterium contortum		99.97	0.61	MLZ 1		
Actinomyces israelii		0.03	0.15			
43715312	**Acceptable identification**					**43715312**
Eubacterium contortum		94.78	0.48	MLZ 1		
Bifidobacterium adolescentis		3.25	0.32			
Clostridium clostridioforme		1.23	0.22			
43715320	**Very good identification**					**43715320**
					GRA	
Eubacterium contortum		99.44	0.83		99	
Bacteroides vulgatus		0.52	0.36		1	
Bifidobacterium longum subsp. infantis		0.02	0.13		99	
43715321	**Good identification**					**43715321**
					GRA	
Eubacterium contortum		99.48	0.48	URE 1	99	
Bacteroides vulgatus		0.52	0.02		1	
43715322	**Good identification**					**43715322**
					GRA	
Eubacterium contortum		92.54	0.70		99	
Parabacteroides distasonis		5.47	0.60		1	
Prevotella oralis		1.02	0.35		1	
43715323	**Acceptable identification**					**43715323**
					GRA	
Eubacterium contortum		92.61	0.35	URE 1	99	
Parabacteroides distasonis		5.47	0.26		1	
Prevotella oralis		1.02	0.00		1	
43715330	**Good identification**					**43715330**
					GRA	
Eubacterium contortum		98.93	0.48	MLZ 1	99	
Parabacteroides distasonis		0.56	0.21		1	
Bacteroides vulgatus		0.52	0.02		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43715332	**Low discrimination**					**43715332**
					GRA	
Parabacteroides distasonis		74.22	0.55	MNS 98	1	
Eubacterium contortum		23.56	0.35		99	
Clostridium clostridioforme		1.73	0.22		50	
43715340	**Very good identification**					**43715340**
Eubacterium contortum		99.93	0.61	SOR 1		
Actinomyces israelii		0.07	0.22			
43715342	**Good identification**					**43715342**
Eubacterium contortum		99.08	0.48	SOR 1		
Actinomyces israelii		0.58	0.25			
Bifidobacterium adolescentis		0.34	0.15			
43715360	**Good identification**					**43715360**
					GRA	
Eubacterium contortum		99.48	0.48	SOR 1	99	
Bacteroides vulgatus		0.52	0.02		1	
43715362	**Acceptable identification**					**43715362**
					GRA	
Eubacterium contortum		92.61	0.35	SOR 1	99	
Parabacteroides distasonis		5.47	0.26		1	
Prevotella oralis		1.02	0.00		1	
43715402	**Low discrimination**					**43715402**
Bifidobacterium adolescentis		76.80	0.45	MNS 87		
Actinomyces israelii		16.74	0.40	LAC 99		
Eubacterium contortum		3.38	0.13			
43715412	**Acceptable identification**					**43715412**
Bifidobacterium adolescentis		93.52	0.46	MNS 87		
Clostridium clostridioforme		3.37	0.18	LAC 99		
Actinomyces israelii		2.49	0.25			
43715500	**Good identification**					**43715500**
Eubacterium contortum		91.39	0.60	RHA 99		
Bifidobacterium adolescentis		3.70	0.45			
Actinomyces israelii		2.15	0.47			
43715502	**Good identification**					**43715502**
Bifidobacterium adolescentis		93.97	0.79	MNS 87		
Eubacterium contortum		4.14	0.47			
Actinomyces israelii		0.83	0.50			
43715503	**Acceptable identification**					**43715503**
Bifidobacterium adolescentis		94.98	0.45	MNS 87		
Eubacterium contortum		4.18	0.13	URE 1		
Actinomyces israelii		0.84	0.16			
43715510	**Low discrimination**					**43715510**
Bifidobacterium adolescentis		51.66	0.46	MNS 87		
Bifidobacterium longum subsp. longum		34.13	0.41	BGL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		10.54	0.26			
43715512	**Very good identification**					**43715512**
Bifidobacterium adolescentis		99.08	0.80	MNS 87		
Bifidobacterium longum subsp. longum		0.66	0.41			
Clostridium clostridioforme		0.12	0.27			
43715513	**Good identification**					**43715513**
Bifidobacterium adolescentis		99.34	0.46	MNS 87		
Bifidobacterium longum subsp. longum		0.66	0.07	URE 1		
43715520	**Low discrimination**					**43715520**
					GRA	
Eubacterium contortum		41.38	0.47	RHA 99	99	
Bacteroides vulgatus		38.75	0.39		1	
Bifidobacterium longum subsp. longum		18.02	0.48		99	
43715522	**Low discrimination**					**43715522**
					GRA	
Bifidobacterium longum subsp. longum		30.99	0.48	MNS 87	99	
Parabacteroides distasonis		18.35	0.48	SAL 1	1	
Bifidobacterium adolescentis		16.34	0.45		99	
43715530	**Good identification**					**43715530**
					GRA	
Bifidobacterium longum subsp. longum		97.07	0.52	MNS 87	99	
Eubacterium contortum		1.32	0.13	SAL 1	99	
Bacteroides vulgatus		1.24	0.05		1	
43715532	**Low discrimination**					**43715532**
					GRA	
Bifidobacterium longum subsp. longum		61.48	0.52	MNS 87	99	
Bifidobacterium adolescentis		23.27	0.46	SAL 1	99	
Parabacteroides distasonis		11.51	0.43		1	
43715542	**Very good identification**					**43715542**
Bifidobacterium adolescentis		96.97	0.63	MNS 87		
Actinomyces israelii		2.69	0.43	SOR 11		
Eubacterium contortum		0.35	0.13			
43715552	**Very good identification**					**43715552**
Bifidobacterium adolescentis		99.61	0.65	MNS 87		
Actinomyces israelii		0.34	0.29	SOR 11		
Bifidobacterium longum subsp. longum		0.05	0.07			
43715600	**Very good identification**					**43715600**
Eubacterium contortum		99.96	0.60	LAC 99		
Actinomyces israelii		0.04	0.17			
43715602	**Good identification**					**43715602**
Eubacterium contortum		97.81	0.47	LAC 99		
Clostridium clostridioforme		1.59	0.22			
Actinomyces israelii		0.37	0.20			
43715620	**Acceptable identification**					**43715620**
					GRA	
Eubacterium contortum		92.16	0.47	LAC 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		7.84	0.21		1	
43715700	**Excellent identification**					**43715700**
					GRA	
Eubacterium contortum		99.98	0.94		99	
Bacteroides vulgatus		0.02	0.21		1	
Actinomyces israelii		0.00	0.28		99	
43715701	**Very good identification**					**43715701**
Eubacterium contortum		100.0	0.60	URE 1		
43715702	**Very good identification**					**43715702**
Eubacterium contortum		99.69	0.81			
Bifidobacterium adolescentis		0.23	0.45			
Clostridium clostridioforme		0.06	0.31			
43715703	**Good identification**					**43715703**
Eubacterium contortum		99.77	0.47	URE 1		
Bifidobacterium adolescentis		0.23	0.11			
43715710	**Very good identification**					**43715710**
Eubacterium contortum		99.85	0.60	MLZ 1		
Bifidobacterium adolescentis		0.05	0.12			
Clostridium clostridioforme		0.04	0.08			
43715712	**Low discrimination**					**43715712**
Eubacterium contortum		65.07	0.47	MLZ 1		
Bifidobacterium adolescentis		18.25	0.46			
Clostridium clostridioforme		16.14	0.42			
43715720	**Good identification**					**43715720**
					GRA	
Eubacterium contortum		92.15	0.81		99	
Bacteroides vulgatus		7.84	0.56		1	
Parabacteroides distasonis		0.01	0.22		1	
43715721	**Acceptable identification**					**43715721**
					GRA	
Eubacterium contortum		92.16	0.47	URE 1	99	
Bacteroides vulgatus		7.84	0.21		1	
43715722	**Good identification**					**43715722**
					GRA	
Eubacterium contortum		94.93	0.68		99	
Parabacteroides distasonis		4.20	0.56		1	
Bacteroides vulgatus		0.46	0.21		1	
43715730	**Acceptable identification**					**43715730**
					GRA	
Eubacterium contortum		90.97	0.47	MLZ 1	99	
Bacteroides vulgatus		7.74	0.21		1	
Bifidobacterium longum subsp. longum		0.68	0.18		99	
43715732	**Low discrimination**					**43715732**
					GRA	
Parabacteroides distasonis		48.92	0.52	MNS 98	1	
Clostridium clostridioforme		29.14	0.42		50	
Eubacterium contortum		20.73	0.34		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43715740	**Very good identification**					**43715740**
Eubacterium contortum		99.93	0.60	SOR 1		
Actinomyces israelii		0.07	0.21			
43715742	**Good identification**					**43715742**
Eubacterium contortum		96.69	0.47	SOR 1		
Bifidobacterium adolescentis		2.74	0.29			
Actinomyces israelii		0.57	0.24			
43715760	**Acceptable identification**					**43715760**
					GRA	
Eubacterium contortum		92.16	0.47	SOR 1	99	
Bacteroides vulgatus		7.84	0.21		1	
43716502	**Acceptable identification**					**43716502**
Bifidobacterium adolescentis		94.25	0.43	MNS 87		
Actinomyces israelii		5.75	0.29	GAL 99		
43716512	**Good identification**					**43716512**
Bifidobacterium adolescentis		99.26	0.45	MNS 87		
Actinomyces israelii		0.74	0.15	GAL 99		
43717000	**Good identification**					**43717000**
					GRA	
Actinomyces israelii		97.95	0.51	CEL 93	99	
Fusobacterium mortiferum		2.06	0.20		1	
43717002	**Acceptable identification**					**43717002**
Actinomyces israelii		92.51	0.54	CEL 93		
Bifidobacterium adolescentis		7.49	0.29			
43717010	**Good identification**					**43717010**
Actinomyces israelii		100.0	0.37	CEL 93		
43717012	**Low discrimination**					**43717012**
Actinomyces israelii		60.15	0.40	CEL 93		
Bifidobacterium adolescentis		39.85	0.31	MLZ 13		
43717040	**Good identification**					**43717040**
Actinomyces israelii		100.0	0.44	CEL 93		
43717042	**Good identification**					**43717042**
Actinomyces israelii		97.49	0.47	CEL 93		
Bifidobacterium adolescentis		2.51	0.13			
43717100	**Low discrimination**					**43717100**
Actinomyces israelii		88.23	0.61	CEL 93		
Eubacterium contortum		6.68	0.27			
Bifidobacterium adolescentis		2.68	0.29			
43717102	**Low discrimination**					**43717102**
Bifidobacterium adolescentis		64.68	0.63	MNS 87		
Actinomyces israelii		32.27	0.64	ARA 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. infantis		2.71	0.26			
43717110	**Low discrimination**					**43717110**
Actinomyces israelii		80.10	0.47	CEL 93		
Bifidobacterium adolescentis		19.90	0.31	MLZ 13		
43717112	**Good identification**					**43717112**
Bifidobacterium adolescentis		94.25	0.65	MNS 87		
Actinomyces israelii		5.75	0.50	ARA 87		
43717120	**Low discrimination**					**43717120**
Actinomyces israelii		62.64	0.41	CEL 93		
Bifidobacterium longum subsp. infantis		21.71	0.13	NAG 6		
Eubacterium contortum		13.11	0.14			
43717140	**Good identification**					**43717140**
Actinomyces israelii		99.04	0.54	CEL 93		
Bifidobacterium adolescentis		0.96	0.13			
43717142	**Low discrimination**					**43717142**
Actinomyces israelii		61.09	0.57	CEL 93		
Bifidobacterium adolescentis		38.92	0.48			
43717150	**Acceptable identification**					**43717150**
Actinomyces israelii		92.68	0.40	CEL 93		
Bifidobacterium adolescentis		7.32	0.15	MLZ 13		
43717152	**Low discrimination**					**43717152**
Bifidobacterium adolescentis		83.90	0.49	MNS 87		
Actinomyces israelii		16.10	0.43	ARA 87		
43717162	**Low discrimination**					**43717162**
Actinomyces israelii		86.10	0.37	CEL 93		
Bifidobacterium adolescentis		8.68	0.13	NAG 6		
Bifidobacterium longum subsp. infantis		5.23	-.04			
43717300	**Very good identification**					**43717300**
Eubacterium contortum		99.01	0.61	MAN 1		
Actinomyces israelii		0.99	0.42			
43717302	**Acceptable identification**					**43717302**
Eubacterium contortum		90.23	0.48	MAN 1		
Actinomyces israelii		7.70	0.45			
Bifidobacterium adolescentis		2.07	0.29			
43717320	**Good identification**					**43717320**
					GRA	
Eubacterium contortum		99.12	0.48	MAN 1	99	
Bacteroides vulgatus		0.52	0.02		1	
Actinomyces israelii		0.36	0.22		99	
43717322	**Acceptable identification**					**43717322**
					GRA	
Eubacterium contortum		90.04	0.35	MAN 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		5.32	0.26		1	
Actinomyces israelii		2.78	0.25		99	
43717342	**Low discrimination**					**43717342**
Actinomyces israelii		71.95	0.38	CEL 93		
Eubacterium contortum		21.90	0.14	RHA 7		
Bifidobacterium adolescentis		6.15	0.13			
43717400	**Good identification**					**43717400**
Actinomyces israelii		99.01	0.50	CEL 93		
Bifidobacterium adolescentis		0.99	0.09			
43717402	**Low discrimination**					**43717402**
Actinomyces israelii		60.15	0.53	CEL 93		
Bifidobacterium adolescentis		39.85	0.43			
43717410	**Acceptable identification**					**43717410**
Actinomyces israelii		92.41	0.35	CEL 93		
Bifidobacterium adolescentis		7.59	0.11	MLZ 13		
43717412	**Low discrimination**					**43717412**
Bifidobacterium adolescentis		84.42	0.45	MNS 87		
Actinomyces israelii		15.58	0.38	LAC 99		
43717440	**Good identification**					**43717440**
Actinomyces israelii		100.0	0.42	CEL 93		
43717442	**Low discrimination**					**43717442**
Actinomyces israelii		82.61	0.45	CEL 93		
Bifidobacterium adolescentis		17.39	0.28			
43717500	**Low discrimination**					**43717500**
Actinomyces israelii		75.36	0.60	CEL 93		
Bifidobacterium adolescentis		18.72	0.43			
Eubacterium contortum		5.71	0.26			
43717502	**Good identification**					**43717502**
Bifidobacterium adolescentis		94.20	0.77	MNS 87		
Actinomyces israelii		5.75	0.63			
Eubacterium contortum		0.05	0.13			
43717503	**Acceptable identification**					**43717503**
Bifidobacterium adolescentis		94.25	0.43	MNS 87		
Actinomyces israelii		5.75	0.29	URE 1		
43717510	**Low discrimination**					**43717510**
Bifidobacterium adolescentis		66.65	0.45	MNS 87		
Actinomyces israelii		32.80	0.46	BGL 99		
Bifidobacterium longum subsp. longum		0.54	0.07			
43717512	**Very good identification**					**43717512**
Bifidobacterium adolescentis		99.24	0.79	MNS 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.74	0.49			
Bifidobacterium longum subsp. longum		0.01	0.07			
43717513	**Good identification**					**43717513**
Bifidobacterium adolescentis		99.26	0.45	MNS 87		
Actinomyces israelii		0.74	0.15	URE 1		
43717520	**Low discrimination**					**43717520**
					GRA	
Actinomyces israelii		65.11	0.39	CEL 93	99	
Eubacterium contortum		13.63	0.13	NAG 6	99	
Bacteroides vulgatus		12.76	0.05		1	
43717522	**Low discrimination**					**43717522**
Bifidobacterium adolescentis		69.29	0.43	MNS 87		
Actinomyces israelii		26.70	0.42	NAG 1		
Bifidobacterium longum subsp. longum		1.62	0.14			
43717532	**Acceptable identification**					**43717532**
Bifidobacterium adolescentis		90.93	0.45	MNS 87		
Actinomyces israelii		4.28	0.28	NAG 1		
Bifidobacterium longum subsp. longum		2.97	0.18			
43717540	**Acceptable identification**					**43717540**
Actinomyces israelii		92.68	0.53	CEL 93		
Bifidobacterium adolescentis		7.32	0.28			
43717542	**Low discrimination**					**43717542**
Bifidobacterium adolescentis		83.90	0.62	MNS 87		
Actinomyces israelii		16.10	0.56	SOR 11		
43717550	**Low discrimination**					**43717550**
					COC	
Actinomyces israelii		58.69	0.39	CEL 93	1	
Bifidobacterium adolescentis		37.90	0.29	MLZ 13	1	
Blautia producta		3.40	-.03		99	
43717552	**Very good identification**					**43717552**
Bifidobacterium adolescentis		97.71	0.63	MNS 87		
Actinomyces israelii		2.29	0.42	SOR 11		
43717562	**Low discrimination**					**43717562**
Actinomyces israelii		54.81	0.35	CEL 93		
Bifidobacterium adolescentis		45.19	0.28	NAG 6		
43717700	**Very good identification**					**43717700**
Eubacterium contortum		98.97	0.60	MAN 1		
Actinomyces israelii		0.99	0.41			
Bifidobacterium adolescentis		0.03	0.09			
43717702	**Low discrimination**					**43717702**
Eubacterium contortum		78.55	0.47	MAN 1		
Bifidobacterium adolescentis		14.75	0.43			
Actinomyces israelii		6.70	0.44			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43717712	**Acceptable identification**					**43717712**
Bifidobacterium adolescentis		90.05	0.45	MNS 87		
Actinomyces israelii		5.00	0.29	RHA 1		
Eubacterium contortum		3.96	0.13			
43717720	**Acceptable identification**					**43717720**
					GRA	
Eubacterium contortum		91.85	0.47	MAN 1	99	
Bacteroides vulgatus		7.82	0.21		1	
Actinomyces israelii		0.33	0.20		99	
43717742	**Low discrimination**					**43717742**
Actinomyces israelii		49.91	0.37	CEL 93		
Bifidobacterium adolescentis		34.90	0.28	RHA 7		
Eubacterium contortum		15.19	0.13			
43721100	**Low discrimination**					**43721100**
					GRA	
Fusobacterium mortiferum		70.79	0.36	MLT 11	1	
Actinomyces naeslundii		19.66	0.38	FRU 99	99	
Eubacterium saburreum		5.63	0.16		85	
43721101	**Good identification**					**43721101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43721102	**Low discrimination**					**43721102**
					GRA	
Actinomyces naeslundii		55.88	0.38	ESL 12	99	
Fusobacterium mortiferum		22.36	0.20	MNS 88	1	
Bifidobacterium adolescentis		15.32	0.13	FRU 95	99	
43721103	**Good identification**					**43721103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43721120	**Low discrimination**					**43721120**
					GRA	
Actinomyces naeslundii		85.61	0.38	ESL 12	99	
Prevotella melaninogenica		14.40	-.04	MNS 88	1	
43721121	**Good identification**					**43721121**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43721122	**Low discrimination**					**43721122**
					GRA	
Actinomyces naeslundii		62.23	0.38	ESL 12	99	
Prevotella oralis		14.29	0.02	MNS 88	1	
Bifidobacterium longum subsp. infantis		12.74	-.03	FRU 95	99	
43721123	**Good identification**					**43721123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43723000	**Good identification**					**43723000**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43723001	**Good identification**					**43723001**
Actinomyces naeslundii		100.0	0.39	ESL 12		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
43723002	**Good identification**					**43723002**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43723003	**Good identification**					**43723003**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43723020	**Good identification**					**43723020**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43723021	**Good identification**					**43723021**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43723022	**Good identification**					**43723022**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43723023	**Good identification**					**43723023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43723100	**Low discrimination**					**43723100**
Actinomyces naeslundii		88.80	0.43	ESL 12		
Actinomyces israelii		11.21	0.19	MNS 88		
43723101	**Good identification**					**43723101**
Actinomyces naeslundii		100.0	0.46	ESL 12		
43723102	**Low discrimination**					**43723102**
Actinomyces naeslundii		76.64	0.43	ESL 12		
Actinomyces israelii		14.51	0.22	MNS 88		
Bifidobacterium adolescentis		8.86	0.12	FRU 95		
43723103	**Good identification**					**43723103**
Actinomyces naeslundii		100.0	0.46	ESL 12		
43723120	**Good identification**					**43723120**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43723121	**Good identification**					**43723121**
Actinomyces naeslundii		100.0	0.46	ESL 12		
43723122	**Good identification**					**43723122**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43723123	**Good identification**					**43723123**
Actinomyces naeslundii		100.0	0.46	ESL 12		
43725322	**Good identification**					**43725322**
					GRA	
Parabacteroides distasonis		96.38	0.37	MNS 98	1	
Eubacterium contortum		3.62	0.01	FRU 99	99	
43725502	**Acceptable identification**					**43725502**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		93.51	0.42	MNS 87		
Actinomyces israelii		4.65	0.26	FRU 99		
Bifidobacterium longum subsp. longum		1.85	0.11			
43725512	**Good identification**					**43725512**
Bifidobacterium adolescentis		96.07	0.43	MNS 87		
Bifidobacterium longum subsp. longum		2.64	0.15	FRU 99		
Clostridium clostridioforme		1.29	0.08			
43727100	**Good identification**					**43727100**
Actinomyces israelii		100.0	0.37	FRU 99		
43727102	**Low discrimination**					**43727102**
Actinomyces israelii		73.79	0.40	FRU 99		
Bifidobacterium adolescentis		26.21	0.26	CEL 93		
43727500	**Good identification**					**43727500**
Actinomyces israelii		100.0	0.35	FRU 99		
43727502	**Low discrimination**					**43727502**
Bifidobacterium adolescentis		74.40	0.40	MNS 87		
Actinomyces israelii		25.60	0.39	FRU 99		
43727512	**Acceptable identification**					**43727512**
Bifidobacterium adolescentis		94.43	0.42	MNS 87		
Actinomyces israelii		3.97	0.24	FRU 99		
Bifidobacterium dentium		1.59	-.07			
43730000	**Very good identification**					**43730000**
					GRA	
Fusobacterium mortiferum		99.73	0.61	MLT 11	1	
Actinomyces naeslundii		0.27	0.29		99	
43730002	**Good identification**					**43730002**
					GRA	
Fusobacterium mortiferum		97.60	0.45	MLT 11	1	
Actinomyces naeslundii		2.40	0.29	BGL 10	99	
43730100	**Very good identification**					**43730100**
					GRA	
Fusobacterium mortiferum		99.67	0.68	MLT 11	1	
Actinomyces naeslundii		0.33	0.37		99	
43730102	**Good identification**					**43730102**
					GRA	
Fusobacterium mortiferum		95.72	0.51	MLT 11	1	
Actinomyces naeslundii		2.87	0.37	BGL 10	99	
Bifidobacterium adolescentis		0.99	0.13		99	
43730103	**Low discrimination**					**43730103**
					GRA	
Actinomyces naeslundii		81.66	0.40	ESL 12	99	
Fusobacterium mortiferum		18.34	0.17	MNS 88	1	
43730121	**Good identification**					**43730121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43730122	**Low discrimination**					**43730122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		47.54	0.37	ESL 12	99	
Fusobacterium mortiferum		16.02	0.17	MNS 88	1	
Prevotella oralis		13.79	0.02	GAL 96	1	
43730123	**Good identification**					**43730123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
43731000	**Good identification**					**43731000**
					GRA	
Fusobacterium mortiferum		94.73	0.64	MLT 11	1	
Actinomyces naeslundii		4.14	0.53		99	
Actinomyces israelii		0.66	0.30		99	
43731001	**Low discrimination**					**43731001**
					GRA	
Actinomyces naeslundii		86.65	0.56	ESL 12	99	
Fusobacterium mortiferum		13.35	0.30	MNS 88	1	
43731002	**Low discrimination**					**43731002**
					GRA	
Fusobacterium mortiferum		65.54	0.48	MLT 11	1	
Actinomyces naeslundii		25.77	0.53	BGL 10	99	
Actinomyces israelii		6.20	0.33		99	
43731003	**Good identification**					**43731003**
					GRA	
Actinomyces naeslundii		98.32	0.56	ESL 12	99	
Fusobacterium mortiferum		1.68	0.14	MNS 88	1	
43731020	**Low discrimination**					**43731020**
					GRA	
Actinomyces naeslundii		70.14	0.53	ESL 12	99	
Fusobacterium mortiferum		16.21	0.30	MNS 88	1	
Clostridium perfringens		13.65	0.18		99	
43731021	**Good identification**					**43731021**
Actinomyces naeslundii		100.0	0.56	ESL 12		
43731022	**Low discrimination**					**43731022**
					GRA	
Actinomyces naeslundii		71.89	0.53	ESL 12	99	
Prevotella oralis		15.62	0.16	MNS 88	1	
Clostridium perfringens		6.58	0.13		99	
43731023	**Good identification**					**43731023**
Actinomyces naeslundii		100.0	0.56	ESL 12		
43731041	**Good identification**					**43731041**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43731042	**Low discrimination**					**43731042**
					GRA	
Actinomyces naeslundii		59.69	0.40	ESL 12	99	
Actinomyces israelii		31.63	0.26	MNS 88	99	
Fusobacterium mortiferum		8.69	0.14		1	
43731043	**Good identification**					**43731043**
Actinomyces naeslundii		100.0	0.43	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43731060	**Low discrimination**					**43731060**
Actinomyces naeslundii		82.64	0.40	ESL 12		
Clostridium perfringens		17.36	0.06	MNS 88		
43731061	**Good identification**					**43731061**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43731062	**Acceptable identification**					**43731062**
Actinomyces naeslundii		91.00	0.40	ESL 12		
Clostridium perfringens		9.00	0.00	MNS 88		
43731063	**Good identification**					**43731063**
Actinomyces naeslundii		100.0	0.43	ESL 12		
43731100	**Good identification**					**43731100**
					GRA	
Fusobacterium mortiferum		92.74	0.71	MLT 11	1	
Actinomyces naeslundii		4.94	0.60		99	
Actinomyces israelii		1.11	0.40		99	
43731101	**Low discrimination**					**43731101**
					GRA	
Actinomyces naeslundii		88.78	0.63	ESL 12	99	
Fusobacterium mortiferum		11.22	0.36	MNS 88	1	
43731102	**Low discrimination**					**43731102**
					GRA	
Fusobacterium mortiferum		36.78	0.54	MLT 11	1	
Bifidobacterium adolescentis		25.21	0.48	BGL 10	99	
Actinomyces naeslundii		17.65	0.60		99	
43731103	**Very good identification**					**43731103**
					GRA	
Actinomyces naeslundii		97.31	0.63	ESL 12	99	
Fusobacterium mortiferum		1.37	0.20	MNS 88	1	
Bifidobacterium adolescentis		0.94	0.13		99	
43731110	**Low discrimination**					**43731110**
					GRA	
Fusobacterium mortiferum		56.14	0.36	MLT 11	1	
Actinomyces naeslundii		15.59	0.38	MLZ 1	99	
Eubacterium saburreum		10.97	0.23		85	
43731111	**Good identification**					**43731111**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43731112	**Low discrimination**					**43731112**
					GRA	
Bifidobacterium adolescentis		65.93	0.49	MNS 87	99	
Leptotrichia buccalis		26.54	0.24	XYL 87	1	
Bifidobacterium breve		2.12	0.24	ARA 87	99	
43731113	**Low discrimination**					**43731113**
					GRA	
Actinomyces naeslundii		76.14	0.42	ESL 12	99	
Bifidobacterium adolescentis		17.01	0.15	MNS 88	99	
Leptotrichia buccalis		6.85	-.11	MLZ 5	1	
43731120	**Low discrimination**					**43731120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		48.74	0.60	ESL 12	99	
Clostridium perfringens		24.44	0.33	MNS 88	99	
Fusobacterium mortiferum		9.24	0.36		1	
43731121	**Very good identification**					**43731121**
Actinomyces naeslundii		100.0	0.63	ESL 12		
43731122	**Low discrimination**					**43731122**
					GRA	
Prevotella oralis		25.12	0.36	MNS 99	1	
Bifidobacterium longum subsp. infantis		22.40	0.32	TRE 6	99	
Actinomyces naeslundii		21.00	0.60		99	
43731123	**Very good identification**					**43731123**
					GRA	
Actinomyces naeslundii		97.92	0.63	ESL 12	99	
Prevotella oralis		0.79	0.02	MNS 88	1	
Bifidobacterium longum subsp. infantis		0.70	-.03		99	
43731130	**Low discrimination**					**43731130**
Actinomyces naeslundii		76.22	0.38	ESL 12		
Bifidobacterium breve		14.32	0.12	MNS 88		
Eubacterium saburreum		9.46	0.10	MLZ 5		
43731131	**Good identification**					**43731131**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43731132	**Low discrimination**					**43731132**
					GRA	
Parabacteroides distasonis		63.55	0.42	MNS 98	1	
Bifidobacterium breve		13.12	0.29	XYL 90	99	
Leptotrichia buccalis		10.25	0.07		1	
43731133	**Good identification**					**43731133**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43731140	**Low discrimination**					**43731140**
					GRA	
Fusobacterium mortiferum		41.78	0.36	MLT 11	1	
Actinomyces naeslundii		38.91	0.47	SOR 1	99	
Actinomyces israelii		19.31	0.33		99	
43731141	**Good identification**					**43731141**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43731143	**Good identification**					**43731143**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43731160	**Low discrimination**					**43731160**
Actinomyces naeslundii		62.61	0.47	ESL 12		
Clostridium perfringens		33.89	0.21	MNS 88		
Bifidobacterium breve		3.51	0.12			
43731161	**Good identification**					**43731161**
Actinomyces naeslundii		100.0	0.51	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43731162	**Low discrimination**					**43731162**
Actinomyces naeslundii		48.79	0.47	ESL 12		
Bifidobacterium breve		27.63	0.29	MNS 88		
Clostridium perfringens		12.43	0.15			
43731163	**Good identification**					**43731163**
Actinomyces naeslundii		100.0	0.51	ESL 12		
43731322	**Low discrimination**					**43731322**
					SPO	
Parabacteroides distasonis		80.21	0.55	MNS 98	1	
Prevotella oralis		15.24	0.30	XYL 90	1	
Clostridium ramosum		3.33	0.19		92	
43731332	**Good identification**					**43731332**
Parabacteroides distasonis		96.77	0.50	MNS 98		
Clostridium clostridioforme		3.23	0.19	XYL 90		
43731500	**Low discrimination**					**43731500**
					GRA	
Fusobacterium mortiferum		34.02	0.36	MLT 11	1	
Actinomyces israelii		33.08	0.39	ARA 1	99	
Bifidobacterium adolescentis		17.33	0.28		99	
43731502	**Very good identification**					**43731502**
Bifidobacterium adolescentis		96.08	0.62	MNS 87		
Actinomyces israelii		2.78	0.42	XYL 87		
Bifidobacterium longum subsp. longum		0.39	0.19			
43731512	**Very good identification**					**43731512**
Bifidobacterium adolescentis		98.18	0.63	MNS 87		
Clostridium clostridioforme		0.87	0.25	XYL 87		
Bifidobacterium longum subsp. longum		0.55	0.23			
43731522	**Low discrimination**					**43731522**
					GRA	
Parabacteroides distasonis		62.44	0.43	MNS 98	1	
Bifidobacterium longum subsp. longum		17.36	0.29	XYL 90	99	
Bifidobacterium adolescentis		10.94	0.28		99	
43731532	**Low discrimination**					**43731532**
					GRA	
Parabacteroides distasonis		38.08	0.38	MNS 98	1	
Bifidobacterium longum subsp. longum		33.48	0.33	XYL 90	99	
Bifidobacterium adolescentis		15.15	0.29		99	
43731542	**Acceptable identification**					**43731542**
Bifidobacterium adolescentis		91.66	0.46	MNS 87		
Actinomyces israelii		8.34	0.35	XYL 87		
43731552	**Good identification**					**43731552**
Bifidobacterium adolescentis		98.90	0.48	MNS 87		
Actinomyces israelii		1.10	0.20	XYL 87		
43731712	**Low discrimination**					**43731712**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		85.68	0.40	MNS 99	50	
Bifidobacterium adolescentis		13.31	0.29	XYL 91	99	
Parabacteroides distasonis		1.01	0.12		1	
43731722	**Acceptable identification**					**43731722**
					GRA	
Parabacteroides distasonis		90.46	0.51	MNS 98	1	
Clostridium clostridioforme		9.12	0.28	XYL 90	50	
Eubacterium contortum		0.41	0.00		99	
43731732	**Low discrimination**					**43731732**
Parabacteroides distasonis		53.96	0.47	MNS 98		
Clostridium clostridioforme		46.04	0.40	XYL 90		
43732001	**Good identification**					**43732001**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43732003	**Good identification**					**43732003**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43732021	**Good identification**					**43732021**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43732023	**Good identification**					**43732023**
Actinomyces naeslundii		100.0	0.37	ESL 12		
43732101	**Good identification**					**43732101**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43732102	**Low discrimination**					**43732102**
					GRA	
Actinomyces naeslundii		60.58	0.42	ESL 12	99	
Actinomyces israelii		14.49	0.22	MNS 88	99	
Fusobacterium mortiferum		10.51	0.17	GAL 96	1	
43732103	**Good identification**					**43732103**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43732120	**Good identification**					**43732120**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43732121	**Good identification**					**43732121**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43732122	**Good identification**					**43732122**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43732123	**Good identification**					**43732123**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43733000	**Low discrimination**					**43733000**
					GRA	
Actinomyces naeslundii		63.01	0.58	ESL 12	99	
Actinomyces israelii		29.48	0.43	MNS 88	99	
Fusobacterium mortiferum		7.50	0.30		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43733001	**Very good identification**					**43733001**
Actinomyces naeslundii		100.0	0.61	ESL 12		
43733002	**Low discrimination**					**43733002**
Actinomyces naeslundii		57.72	0.58	ESL 12		
Actinomyces israelii		40.51	0.46	MNS 88		
Bifidobacterium adolescentis		1.00	0.12			
43733003	**Very good identification**					**43733003**
Actinomyces naeslundii		100.0	0.61	ESL 12		
43733011	**Good identification**					**43733011**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43733013	**Good identification**					**43733013**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43733020	**Very good identification**					**43733020**
Actinomyces naeslundii		97.10	0.58	ESL 12		
Actinomyces israelii		2.90	0.22	MNS 88		
43733021	**Very good identification**					**43733021**
Actinomyces naeslundii		100.0	0.61	ESL 12		
43733022	**Very good identification**					**43733022**
Actinomyces naeslundii		95.71	0.58	ESL 12		
Actinomyces israelii		4.29	0.25	MNS 88		
43733023	**Very good identification**					**43733023**
Actinomyces naeslundii		100.0	0.61	ESL 12		
43733030	**Good identification**					**43733030**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43733031	**Good identification**					**43733031**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43733032	**Good identification**					**43733032**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43733033	**Good identification**					**43733033**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43733040	**Low discrimination**					**43733040**
Actinomyces israelii		50.77	0.36	CEL 93		
Actinomyces naeslundii		49.23	0.45	XYL 92		
43733041	**Good identification**					**43733041**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43733042	**Low discrimination**					**43733042**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		60.73	0.39	CEL 93		
Actinomyces naeslundii		39.27	0.45	XYL 92		
43733043	**Good identification**					**43733043**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43733060	**Acceptable identification**					**43733060**
Actinomyces naeslundii		93.82	0.45	ESL 12		
Actinomyces israelii		6.18	0.15	MNS 88		
43733061	**Good identification**					**43733061**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43733062	**Acceptable identification**					**43733062**
Actinomyces naeslundii		91.02	0.45	ESL 12		
Actinomyces israelii		8.99	0.18	MNS 88		
43733063	**Good identification**					**43733063**
Actinomyces naeslundii		100.0	0.48	ESL 12		
43733100	**Low discrimination**					**43733100**
					GRA	
Actinomyces naeslundii		56.78	0.65	ESL 12	99	
Actinomyces israelii		37.34	0.53	MNS 88	99	
Fusobacterium mortiferum		5.54	0.36		1	
43733101	**Very good identification**					**43733101**
Actinomyces naeslundii		99.56	0.68	ESL 12		
Actinomyces israelii		0.44	0.19	MNS 88		
43733102	**Low discrimination**					**43733102**
Actinomyces naeslundii		37.95	0.65	ESL 12		
Actinomyces israelii		37.43	0.56	MNS 88		
Bifidobacterium adolescentis		22.85	0.46			
43733103	**Very good identification**					**43733103**
Actinomyces naeslundii		98.94	0.68	ESL 12		
Actinomyces israelii		0.66	0.22	MNS 88		
Bifidobacterium adolescentis		0.40	0.12			
43733110	**Low discrimination**					**43733110**
Actinomyces israelii		62.06	0.39	CEL 93		
Actinomyces naeslundii		33.24	0.43	XYL 92		
Bifidobacterium adolescentis		4.70	0.13	MLZ 13		
43733111	**Good identification**					**43733111**
Actinomyces naeslundii		100.0	0.46	ESL 12		
43733112	**Low discrimination**					**43733112**
Bifidobacterium adolescentis		75.98	0.48	MNS 87		
Actinomyces israelii		15.22	0.42	XYL 87		
Actinomyces naeslundii		5.43	0.43	ARA 87		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
43733113	**Acceptable identification**					**43733113**
Actinomyces naeslundii		91.39	0.46	ESL 12		
Bifidobacterium adolescentis		8.61	0.13	MNS 88		
43733120	**Good identification**					**43733120**
					SPO	
Actinomyces naeslundii		90.19	0.65	ESL 12	1	
Clostridium ramosum		5.57	0.19	MNS 88	92	
Actinomyces israelii		3.79	0.33		1	
43733121	**Very good identification**					**43733121**
Actinomyces naeslundii		100.0	0.68	ESL 12		
43733122	**Low discrimination**					**43733122**
					SPO	
Actinomyces naeslundii		72.26	0.65	ESL 12	1	
Clostridium ramosum		17.84	0.30	MNS 88	92	
Actinomyces israelii		4.55	0.36		1	
43733123	**Very good identification**					**43733123**
Actinomyces naeslundii		100.0	0.68	ESL 12		
43733130	**Low discrimination**					**43733130**
Actinomyces naeslundii		82.24	0.43	ESL 12		
Actinomyces israelii		9.80	0.18	MNS 88		
Bifidobacterium breve		7.96	0.12	MLZ 5		
43733131	**Good identification**					**43733131**
Actinomyces naeslundii		100.0	0.46	ESL 12		
43733132	**Low discrimination**					**43733132**
Actinomyces naeslundii		43.51	0.43	ESL 12		
Bifidobacterium breve		42.57	0.29	MNS 88		
Actinomyces israelii		7.78	0.21	MLZ 5		
43733133	**Good identification**					**43733133**
Actinomyces naeslundii		100.0	0.46	ESL 12		
43733140	**Low discrimination**					**43733140**
Actinomyces israelii		59.17	0.46	CEL 93		
Actinomyces naeslundii		40.83	0.52	XYL 92		
43733141	**Good identification**					**43733141**
Actinomyces naeslundii		100.0	0.55	ESL 12		
43733142	**Low discrimination**					**43733142**
Actinomyces israelii		57.49	0.49	CEL 93		
Actinomyces naeslundii		26.45	0.52	XYL 92		
Bifidobacterium adolescentis		11.15	0.31			
43733143	**Good identification**					**43733143**
Actinomyces naeslundii		98.56	0.55	ESL 12		
Actinomyces israelii		1.44	0.15	MNS 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
43733160	**Low discrimination**					**43733160**
Actinomyces naeslundii		89.18	0.52	ESL 12		
Actinomyces israelii		8.25	0.26	MNS 88		
Bifidobacterium breve		2.57	0.12			
43733161	**Good identification**					**43733161**
Actinomyces naeslundii		100.0	0.55	ESL 12		
43733162	**Low discrimination**					**43733162**
Actinomyces naeslundii		69.90	0.52	ESL 12		
Bifidobacterium breve		20.40	0.29	MNS 88		
Actinomyces israelii		9.70	0.29			
43733163	**Good identification**					**43733163**
Actinomyces naeslundii		100.0	0.55	ESL 12		
43733302	**Low discrimination**					**43733302**
Actinomyces israelii		79.21	0.37	CEL 93		
Actinomyces naeslundii		10.78	0.31	XYL 92		
Bifidobacterium adolescentis		6.49	0.12	RHA 7		
43733400	**Good identification**					**43733400**
Actinomyces israelii		97.43	0.41	CEL 93		
Actinomyces naeslundii		2.57	0.23	XYL 92		
43733402	**Low discrimination**					**43733402**
Actinomyces israelii		82.01	0.44	CEL 93		
Bifidobacterium adolescentis		16.55	0.26	XYL 92		
Actinomyces naeslundii		1.44	0.23			
43733442	**Acceptable identification**					**43733442**
Actinomyces israelii		93.98	0.37	CEL 93		
Bifidobacterium adolescentis		6.03	0.10	XYL 92		
43733500	**Acceptable identification**					**43733500**
Actinomyces israelii		91.37	0.52	CEL 93		
Bifidobacterium adolescentis		6.91	0.26	XYL 92		
Actinomyces naeslundii		1.72	0.31			
43733502	**Low discrimination**					**43733502**
Bifidobacterium adolescentis		83.14	0.60	MNS 87		
Actinomyces israelii		16.65	0.55	XYL 87		
Actinomyces naeslundii		0.21	0.31			
43733510	**Low discrimination**					**43733510**
Actinomyces israelii		61.78	0.37	CEL 93		
Bifidobacterium adolescentis		38.23	0.28	XYL 92		
43733512	**Very good identification**					**43733512**
Bifidobacterium adolescentis		97.50	0.62	MNS 87		
Actinomyces israelii		2.39	0.40	XYL 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium dentium		0.11	-.07			
43733540	**Good identification**					**43733540**
Actinomyces israelii		97.65	0.44	CEL 93		
Bifidobacterium adolescentis		2.35	0.10	XYL 92		
43733542	**Low discrimination**					**43733542**
Bifidobacterium adolescentis		61.34	0.45	MNS 87		
Actinomyces israelii		38.66	0.47	XYL 87		
43733552	**Acceptable identification**					**43733552**
Bifidobacterium adolescentis		92.85	0.46	MNS 87		
Actinomyces israelii		7.15	0.33	XYL 87		
43733702	**Low discrimination**					**43733702**
Actinomyces israelii		59.88	0.35	CEL 93		
Bifidobacterium adolescentis		40.12	0.26	XYL 92		
43734000	**Good identification**					**43734000**
					GRA	
Fusobacterium mortiferum		98.40	0.42	MLT 11	1	
Actinomyces israelii		1.60	0.14	XYL 7	99	
43734100	**Good identification**					**43734100**
					GRA	
Fusobacterium mortiferum		97.30	0.48	MLT 11	1	
Actinomyces israelii		2.70	0.24	XYL 7	99	
43734322	**Low discrimination**					**43734322**
					GRA	
Parabacteroides distasonis		84.47	0.37	MNS 98	1	
Clostridium clostridioforme		12.36	0.17	GAL 99	50	
Eubacterium contortum		3.17	0.01		99	
43734502	**Low discrimination**					**43734502**
Bifidobacterium adolescentis		85.60	0.42	MNS 87		
Clostridium clostridioforme		10.15	0.23	GAL 99		
Actinomyces israelii		4.25	0.26			
43734512	**Low discrimination**					**43734512**
Bifidobacterium adolescentis		69.35	0.43	MNS 87		
Clostridium clostridioforme		30.66	0.34	GAL 99		
43734612	**Good identification**					**43734612**
Clostridium clostridioforme		100.0	0.40	MNS 99		
43734632	**Good identification**					**43734632**
Clostridium clostridioforme		100.0	0.40	MNS 99		
43734702	**Acceptable identification**					**43734702**
Clostridium clostridioforme		91.95	0.37	MNS 99		
Eubacterium contortum		6.98	0.13			
Bifidobacterium adolescentis		1.07	0.07			
43734712	**Good identification**					**43734712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		99.69	0.49	MNS 99		
Bifidobacterium adolescentis		0.31	0.09			
43734722	**Low discrimination**					**43734722**
					GRA	
Clostridium clostridioforme		78.08	0.37	MNS 99	50	
Parabacteroides distasonis		20.88	0.33		1	
Eubacterium contortum		1.05	0.00		99	
43734732	**Good identification**					**43734732**
Clostridium clostridioforme		96.94	0.49	MNS 99		
Parabacteroides distasonis		3.06	0.29			
43735000	**Low discrimination**					**43735000**
					GRA	
Actinomyces israelii		51.27	0.48	CEL 93	99	
Fusobacterium mortiferum		47.91	0.45		1	
Eubacterium saburreum		0.83	0.13		85	
43735002	**Low discrimination**					**43735002**
					GRA	
Actinomyces israelii		85.57	0.51	CEL 93	99	
Bifidobacterium adolescentis		8.51	0.28		99	
Fusobacterium mortiferum		5.92	0.29		1	
43735012	**Low discrimination**					**43735012**
Actinomyces israelii		51.27	0.37	CEL 93		
Bifidobacterium adolescentis		41.68	0.29	MLZ 13		
Clostridium clostridioforme		7.05	0.13			
43735040	**Good identification**					**43735040**
					GRA	
Actinomyces israelii		97.63	0.41	CEL 93	99	
Fusobacterium mortiferum		2.37	0.11		1	
43735042	**Good identification**					**43735042**
Actinomyces israelii		96.94	0.44	CEL 93		
Bifidobacterium adolescentis		3.06	0.12			
43735100	**Low discrimination**					**43735100**
					GRA	
Actinomyces israelii		53.79	0.58	CEL 93	99	
Fusobacterium mortiferum		29.32	0.51		1	
Eubacterium contortum		6.14	0.27		99	
43735102	**Low discrimination**					**43735102**
Bifidobacterium adolescentis		66.63	0.62	MNS 87		
Actinomyces israelii		27.08	0.61	ARA 87		
Bifidobacterium longum subsp. infantis		3.42	0.26			
43735110	**Low discrimination**					**43735110**
Actinomyces israelii		70.82	0.44	CEL 93		
Bifidobacterium adolescentis		21.60	0.29	MLZ 13		
Eubacterium saburreum		3.65	0.18			
43735112	**Good identification**					**43735112**
Bifidobacterium adolescentis		94.36	0.63	MNS 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		4.69	0.47	ARA 87		
Clostridium clostridioforme		0.54	0.22			
43735120	**Low discrimination**					**43735120**
					GRA	
Actinomyces israelii		34.63	0.38	CEL 93	99	
Bifidobacterium longum subsp. infantis		18.08	0.13	NAG 6	99	
Parabacteroides distasonis		17.12	0.29		1	
43735122	**Low discrimination**					**43735122**
					GRA	
Parabacteroides distasonis		84.90	0.63	MNS 98	1	
Bifidobacterium longum subsp. infantis		9.16	0.30		99	
Actinomyces israelii		2.60	0.41		99	
43735132	**Good identification**					**43735132**
					GRA	
Parabacteroides distasonis		94.81	0.58	MNS 98	1	
Bifidobacterium adolescentis		2.57	0.29		99	
Clostridium clostridioforme		1.46	0.22		50	
43735140	**Good identification**					**43735140**
					GRA	
Actinomyces israelii		96.35	0.51	CEL 93	99	
Fusobacterium mortiferum		1.36	0.17		1	
Bifidobacterium adolescentis		1.14	0.12		99	
43735142	**Low discrimination**					**43735142**
Actinomyces israelii		56.12	0.54	CEL 93		
Bifidobacterium adolescentis		43.88	0.46			
43735150	**Acceptable identification**					**43735150**
Actinomyces israelii		91.17	0.37	CEL 93		
Bifidobacterium adolescentis		8.84	0.13	MLZ 13		
43735152	**Low discrimination**					**43735152**
Bifidobacterium adolescentis		86.48	0.48	MNS 87		
Actinomyces israelii		13.52	0.40	ARA 87		
43735222	**Low discrimination**					**43735222**
					GRA	
Parabacteroides distasonis		85.57	0.37	MNS 98	1	
Clostridium clostridioforme		11.22	0.16	LAC 99	50	
Eubacterium contortum		3.21	0.01		99	
43735300	**Very good identification**					**43735300**
					GRA	
Eubacterium contortum		99.29	0.61	TRE 1	99	
Actinomyces israelii		0.66	0.39		99	
Fusobacterium mortiferum		0.05	0.17		1	
43735302	**Low discrimination**					**43735302**
					GRA	
Eubacterium contortum		87.43	0.48	TRE 1	99	
Actinomyces israelii		4.95	0.42		99	
Parabacteroides distasonis		4.15	0.37		1	
43735312	**Low discrimination**					**43735312**
					GRA	
Clostridium clostridioforme		58.63	0.36	MNS 99	50	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		15.79	0.32	ARA 94	1	
Bifidobacterium adolescentis		14.11	0.29		99	
43735320	**Acceptable identification**					**43735320**
					GRA	
Eubacterium contortum		93.32	0.48	TRE 1	99	
Parabacteroides distasonis		4.43	0.37		1	
Bacteroides vulgatus		2.02	0.13		1	
43735322	**Very good identification**					**43735322**
					GRA	
Parabacteroides distasonis		95.63	0.71	MNS 98	1	
Eubacterium contortum		3.59	0.35		99	
Clostridium clostridioforme		0.42	0.25		50	
43735323	**Good identification**					**43735323**
					GRA	
Parabacteroides distasonis		96.38	0.37	MNS 98	1	
Eubacterium contortum		3.62	0.01	URE 1	99	
43735332	**Very good identification**					**43735332**
					GRA	
Parabacteroides distasonis		96.32	0.67	MNS 98	1	
Clostridium clostridioforme		3.61	0.36		50	
Eubacterium contortum		0.07	0.01		99	
43735342	**Low discrimination**					**43735342**
Actinomyces israelii		63.96	0.35	CEL 93		
Eubacterium contortum		29.33	0.14	RHA 7		
Bifidobacterium adolescentis		6.71	0.12			
43735362	**Acceptable identification**					**43735362**
					GRA	
Parabacteroides distasonis		93.71	0.37	MNS 98	1	
Eubacterium contortum		3.52	0.01	SOR 1	99	
Actinomyces israelii		2.78	0.15		99	
43735400	**Low discrimination**					**43735400**
					SPO	
Actinomyces israelii		89.18	0.46	CEL 93	1	
Clostridium butyricum		8.69	0.08		99	
Bifidobacterium adolescentis		1.10	0.07		1	
43735402	**Low discrimination**					**43735402**
Actinomyces israelii		51.95	0.49	CEL 93		
Bifidobacterium adolescentis		42.24	0.42			
Clostridium clostridioforme		4.49	0.22			
43735412	**Low discrimination**					**43735412**
Bifidobacterium adolescentis		64.66	0.43	MNS 87		
Clostridium clostridioforme		25.62	0.33	LAC 99		
Actinomyces israelii		9.72	0.35			
43735440	**Good identification**					**43735440**
Actinomyces israelii		100.0	0.39	CEL 93		
43735442	**Low discrimination**					**43735442**
Actinomyces israelii		79.47	0.42	CEL 93		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Bifidobacterium adolescentis	20.54	0.26			
43735500	**Low discrimination**					**43735500**
					SPO	
	Actinomyces israelii	48.44	0.57	CEL 93	1	
	Clostridium butyricum	28.32	0.32		99	
	Bifidobacterium adolescentis	14.77	0.42		1	
43735502	**Good identification**					**43735502**
					SPO	
	Bifidobacterium adolescentis	93.98	0.76	MNS 87	1	
	Actinomyces israelii	4.67	0.60		1	
	Clostridium butyricum	0.71	0.25		99	
43735503	**Good identification**					**43735503**
	Bifidobacterium adolescentis	95.27	0.42	MNS 87		
	Actinomyces israelii	4.73	0.26	URE 1		
43735510	**Low discrimination**					**43735510**
	Bifidobacterium adolescentis	57.32	0.43	MNS 87		
	Actinomyces israelii	22.98	0.43	BGL 99		
	Bifidobacterium longum subsp. longum	14.20	0.31			
43735512	**Very good identification**					**43735512**
	Bifidobacterium adolescentis	97.77	0.77	MNS 87		
	Clostridium clostridioforme	1.31	0.42			
	Actinomyces israelii	0.59	0.46			
43735513	**Good identification**					**43735513**
	Bifidobacterium adolescentis	98.68	0.43	MNS 87		
	Clostridium clostridioforme	1.32	0.08	URE 1		
43735520	**Low discrimination**					**43735520**
					GRA	
	Bifidobacterium longum subsp. longum	47.55	0.37	MNS 87	99	
	Bacteroides vulgatus	17.04	0.16	SAL 1	1	
	Actinomyces israelii	13.99	0.36		99	
43735522	**Low discrimination**					**43735522**
					GRA	
	Parabacteroides distasonis	75.57	0.59	MNS 98	1	
	Bifidobacterium adolescentis	9.85	0.42		99	
	Bifidobacterium longum subsp. longum	7.00	0.37		99	
43735530	**Acceptable identification**					**43735530**
					GRA	
	Bifidobacterium longum subsp. longum	92.25	0.41	MNS 87	99	
	Parabacteroides distasonis	3.18	0.21	SAL 1	1	
	Actinomyces israelii	2.38	0.22		99	
43735532	**Low discrimination**					**43735532**
					GRA	
	Parabacteroides distasonis	50.01	0.55	MNS 98	1	
	Clostridium clostridioforme	19.61	0.42		50	
	Bifidobacterium adolescentis	14.79	0.43		99	
43735540	**Low discrimination**					**43735540**
					SPO	
	Actinomyces israelii	81.46	0.50	CEL 93	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		10.65	0.14		99	
Bifidobacterium adolescentis		7.89	0.26		1	
43735542	**Low discrimination**					**43735542**
					SPO	
Bifidobacterium adolescentis		86.08	0.60	MNS 87	1	
Actinomyces israelii		13.46	0.53	SOR 11	1	
Clostridium butyricum		0.46	0.07		99	
43735550	**Low discrimination**					**43735550**
					COC	
Actinomyces israelii		53.19	0.36	CEL 93	1	
Bifidobacterium adolescentis		42.16	0.28	MLZ 13	1	
Blautia producta		4.65	-.03		99	
43735552	**Very good identification**					**43735552**
Bifidobacterium adolescentis		98.02	0.62	MNS 87		
Actinomyces israelii		1.87	0.39	SOR 11		
Clostridium clostridioforme		0.11	0.08			
43735602	**Low discrimination**					**43735602**
Clostridium clostridioforme		82.21	0.37	MNS 99		
Actinomyces israelii		9.76	0.30			
Eubacterium contortum		6.96	0.13			
43735612	**Good identification**					**43735612**
Clostridium clostridioforme		99.27	0.48	MNS 99		
Actinomyces israelii		0.39	0.16			
Bifidobacterium adolescentis		0.35	0.09			
43735622	**Low discrimination**					**43735622**
					GRA	
Clostridium clostridioforme		76.14	0.37	MNS 99	50	
Parabacteroides distasonis		22.72	0.33		1	
Eubacterium contortum		1.14	0.00		99	
43735632	**Good identification**					**43735632**
Clostridium clostridioforme		96.59	0.48	MNS 99		
Parabacteroides distasonis		3.41	0.29			
43735700	**Very good identification**					**43735700**
					SPO	
Eubacterium contortum		99.02	0.60	TRE 1	1	
Actinomyces israelii		0.66	0.38		1	
Clostridium butyricum		0.16	0.06		99	
43735702	**Low discrimination**					**43735702**
Eubacterium contortum		60.68	0.47	TRE 1		
Clostridium clostridioforme		24.23	0.46			
Bifidobacterium adolescentis		9.28	0.42			
43735712	**Low discrimination**					**43735712**
					GRA	
Clostridium clostridioforme		89.00	0.57	MNS 99	50	
Bifidobacterium adolescentis		9.15	0.43		99	
Parabacteroides distasonis		0.94	0.29		1	
43735720	**Low discrimination**					**43735720**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Eubacterium contortum		71.78	0.47	TRE 1	99	
Bacteroides vulgatus		25.20	0.32		1	
Parabacteroides distasonis		2.56	0.33		1	
43735722	**Low discrimination**					**43735722**
					GRA	
Parabacteroides distasonis		85.19	0.67	MNS 98	1	
Clostridium clostridioforme		9.65	0.46		50	
Eubacterium contortum		4.27	0.34		99	
43735732	**Low discrimination**					**43735732**
Parabacteroides distasonis		50.80	0.63	MNS 98		
Clostridium clostridioforme		48.70	0.57			
Bacteroides thetaiotaomicron		0.21	0.06			
43736100	**Acceptable identification**					**43736100**
					GRA	
Actinomyces israelii		93.98	0.37	CEL 93	99	
Fusobacterium mortiferum		6.03	0.14	GAL 99	1	
43736102	**Low discrimination**					**43736102**
Actinomyces israelii		73.79	0.40	CEL 93		
Bifidobacterium adolescentis		26.21	0.26	GAL 99		
43736500	**Good identification**					**43736500**
Actinomyces israelii		100.0	0.35	CEL 93		
43736502	**Low discrimination**					**43736502**
Bifidobacterium adolescentis		74.40	0.40	MNS 87		
Actinomyces israelii		25.60	0.39	GAL 99		
43736512	**Acceptable identification**					**43736512**
Bifidobacterium adolescentis		94.43	0.42	MNS 87		
Actinomyces israelii		3.97	0.24	GAL 99		
Bifidobacterium dentium		1.59	-.07			
43737000	**Very good identification**					**43737000**
					GRA	
Actinomyces israelii		99.65	0.61	CEL 93	99	
Actinomyces naeslundii		0.19	0.23		99	
Fusobacterium mortiferum		0.17	0.11		1	
43737002	**Very good identification**					**43737002**
Actinomyces israelii		98.46	0.64	CEL 93		
Bifidobacterium adolescentis		1.41	0.26			
Actinomyces naeslundii		0.12	0.23			
43737010	**Good identification**					**43737010**
Actinomyces israelii		100.0	0.47	CEL 93		
43737012	**Low discrimination**					**43737012**
Actinomyces israelii		89.49	0.50	CEL 93		
Bifidobacterium adolescentis		10.51	0.28	MLZ 13		
43737020	**Good identification**					**43737020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		97.14	0.40	CEL 93		
Actinomyces naeslundii		2.86	0.23	NAG 6		
43737022	**Good identification**					**43737022**
Actinomyces israelii		98.08	0.43	CEL 93		
Actinomyces naeslundii		1.92	0.23	NAG 6		
43737040	**Good identification**					**43737040**
Actinomyces israelii		100.0	0.54	CEL 93		
43737042	**Very good identification**					**43737042**
Actinomyces israelii		99.55	0.57	CEL 93		
Bifidobacterium adolescentis		0.45	0.10			
43737050	**Good identification**					**43737050**
Actinomyces israelii		100.0	0.40	CEL 93		
43737052	**Good identification**					**43737052**
Actinomyces israelii		96.40	0.43	CEL 93		
Bifidobacterium adolescentis		3.60	0.12	MLZ 13		
43737062	**Good identification**					**43737062**
Actinomyces israelii		100.0	0.36	CEL 93		
43737100	**Very good identification**					**43737100**
Actinomyces israelii		99.13	0.71	CEL 93		
Bifidobacterium adolescentis		0.53	0.26			
Actinomyces naeslundii		0.13	0.31			
43737101	**Low discrimination**					**43737101**
Actinomyces israelii		83.45	0.37	CEL 93		
Actinomyces naeslundii		16.55	0.34	URE 1		
43737102	**Low discrimination**					**43737102**
Actinomyces israelii		73.74	0.74	CEL 93		
Bifidobacterium adolescentis		26.20	0.60			
Actinomyces naeslundii		0.07	0.31			
43737103	**Low discrimination**					**43737103**
Actinomyces israelii		67.23	0.40	CEL 93		
Bifidobacterium adolescentis		23.88	0.26	URE 1		
Actinomyces naeslundii		8.89	0.34			
43737110	**Very good identification**					**43737110**
Actinomyces israelii		95.78	0.57	CEL 93		
Bifidobacterium adolescentis		4.22	0.28	MLZ 13		
43737112	**Low discrimination**					**43737112**
Bifidobacterium adolescentis		74.33	0.62	MNS 87		
Actinomyces israelii		25.58	0.60	ARA 87		
Bifidobacterium dentium		0.09	-.07			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
43737120	**Good identification**					**43737120**
Actinomyces israelii		97.95	0.51	CEL 93		
Actinomyces naeslundii		2.05	0.31	NAG 6		
43737122	**Low discrimination**					**43737122**
					GRA	
Actinomyces israelii		88.14	0.54	CEL 93	99	
Parabacteroides distasonis		5.13	0.29	NAG 6	1	
Bifidobacterium adolescentis		4.96	0.26		99	
43737130	**Good identification**					**43737130**
Actinomyces israelii		100.0	0.37	CEL 93		
43737132	**Low discrimination**					**43737132**
					GRA	
Actinomyces israelii		56.76	0.40	CEL 93	99	
Bifidobacterium adolescentis		26.10	0.28	MLZ 13	99	
Parabacteroides distasonis		11.90	0.24	NAG 6	1	
43737140	**Very good identification**					**43737140**
Actinomyces israelii		99.83	0.64	CEL 93		
Bifidobacterium adolescentis		0.17	0.10			
43737142	**Low discrimination**					**43737142**
Actinomyces israelii		89.86	0.67	CEL 93		
Bifidobacterium adolescentis		10.15	0.45			
43737150	**Good identification**					**43737150**
					COC	
Actinomyces israelii		97.41	0.50	CEL 93	1	
Bifidobacterium adolescentis		1.36	0.12	MLZ 13	1	
Blautia producta		1.23	-.03		99	
43737152	**Low discrimination**					**43737152**
Actinomyces israelii		51.99	0.53	CEL 93		
Bifidobacterium adolescentis		48.01	0.46	MLZ 13		
43737160	**Good identification**					**43737160**
Actinomyces israelii		100.0	0.44	CEL 93		
43737162	**Good identification**					**43737162**
Actinomyces israelii		98.25	0.47	CEL 93		
Bifidobacterium adolescentis		1.76	0.10	NAG 6		
43737200	**Good identification**					**43737200**
Actinomyces israelii		100.0	0.42	CEL 93		
43737202	**Good identification**					**43737202**
Actinomyces israelii		100.0	0.45	CEL 93		
43737242	**Good identification**					**43737242**
Actinomyces israelii		100.0	0.38	CEL 93		
43737300	**Low discrimination**					**43737300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		78.90	0.52	CEL 93		
Eubacterium contortum		21.10	0.27	RHA 7		
43737302	**Acceptable identification**					**43737302**
Actinomyces israelii		92.67	0.55	CEL 93		
Bifidobacterium adolescentis		4.42	0.26	RHA 7		
Eubacterium contortum		2.92	0.14			
43737310	**Good identification**					**43737310**
Actinomyces israelii		100.0	0.38	CEL 93		
43737312	**Low discrimination**					**43737312**
Actinomyces israelii		71.95	0.41	CEL 93		
Bifidobacterium adolescentis		28.05	0.28	RHA 7		
43737322	**Low discrimination**					**43737322**
					GRA	
Parabacteroides distasonis		66.89	0.37	MNS 98	1	
Actinomyces israelii		28.85	0.34	MAN 1	99	
Eubacterium contortum		2.51	0.01		99	
43737340	**Good identification**					**43737340**
Actinomyces israelii		100.0	0.45	CEL 93		
43737342	**Good identification**					**43737342**
Actinomyces israelii		98.51	0.48	CEL 93		
Bifidobacterium adolescentis		1.49	0.10	RHA 7		
43737400	**Very good identification**					**43737400**
Actinomyces israelii		100.0	0.59	CEL 93		
43737402	**Low discrimination**					**43737402**
Actinomyces israelii		89.49	0.62	CEL 93		
Bifidobacterium adolescentis		10.51	0.40			
43737410	**Good identification**					**43737410**
Actinomyces israelii		98.57	0.45	CEL 93		
Bifidobacterium adolescentis		1.43	0.07	MLZ 13		
43737412	**Low discrimination**					**43737412**
Actinomyces israelii		50.60	0.48	CEL 93		
Bifidobacterium adolescentis		48.59	0.42	MLZ 13		
Bifidobacterium dentium		0.82	-.07			
43737420	**Good identification**					**43737420**
Actinomyces israelii		100.0	0.39	CEL 93		
43737422	**Good identification**					**43737422**
Actinomyces israelii		100.0	0.42	CEL 93		
43737440	**Good identification**					**43737440**
Actinomyces israelii		100.0	0.52	CEL 93		
43737442	**Good identification**					**43737442**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		96.40	0.55	CEL 93		
Bifidobacterium adolescentis		3.60	0.25			
43737450	**Acceptable identification**					**43737450**
					COC	
Actinomyces israelii		94.19	0.38	CEL 93	1	
Blautia producta		5.81	-.03	MLZ 13	99	
43737452	**Low discrimination**					**43737452**
Actinomyces israelii		76.62	0.41	CEL 93		
Bifidobacterium adolescentis		23.38	0.26	MLZ 13		
43737500	**Good identification**					**43737500**
					SPO	
Actinomyces israelii		94.74	0.70	CEL 93	1	
Bifidobacterium adolescentis		4.17	0.40		1	
Clostridium butyricum		1.09	0.15		99	
43737501	**Good identification**					**43737501**
Actinomyces israelii		100.0	0.35	CEL 93		
43737502	**Low discrimination**					**43737502**
					SPO	
Bifidobacterium adolescentis		74.33	0.74	MNS 87	1	
Actinomyces israelii		25.58	0.73		1	
Clostridium butyricum		0.08	0.08		99	
43737503	**Low discrimination**					**43737503**
Bifidobacterium adolescentis		74.40	0.40	MNS 87		
Actinomyces israelii		25.60	0.39	URE 1		
43737510	**Low discrimination**					**43737510**
Actinomyces israelii		69.95	0.56	CEL 93		
Bifidobacterium adolescentis		25.19	0.42	MLZ 13		
Bifidobacterium dentium		4.16	0.10			
43737512	**Good identification**					**43737512**
Bifidobacterium adolescentis		94.42	0.76	MNS 87		
Actinomyces israelii		3.97	0.59			
Bifidobacterium dentium		1.59	0.27			
43737513	**Acceptable identification**					**43737513**
Bifidobacterium adolescentis		94.43	0.42	MNS 87		
Actinomyces israelii		3.97	0.24	URE 1		
Bifidobacterium dentium		1.59	-.07			
43737520	**Good identification**					**43737520**
					SPO	
Actinomyces israelii		98.46	0.49	CEL 93	1	
Clostridium butyricum		1.54	-.03	NAG 6	99	
43737522	**Low discrimination**					**43737522**
					GRA	
Actinomyces israelii		65.78	0.52	CEL 93	99	
Bifidobacterium adolescentis		30.25	0.40	NAG 6	99	
Parabacteroides distasonis		2.87	0.25		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43737540	**Good identification**					**43737540**
					COC	
Actinomyces israelii		91.53	0.63	CEL 93	1	
Blautia producta		6.96	0.24		99	
Bifidobacterium adolescentis		1.28	0.25		1	
43737542	**Low discrimination**					**43737542**
					COC	
Actinomyces israelii		51.84	0.66	CEL 93	1	
Bifidobacterium adolescentis		47.87	0.59		1	
Blautia producta		0.29	0.07		99	
43737552	**Low discrimination**					**43737552**
					COC	
Bifidobacterium adolescentis		87.05	0.60	MNS 87	1	
Actinomyces israelii		11.53	0.52	SOR 11	1	
Blautia producta		1.30	0.15		99	
43737560	**Low discrimination**					**43737560**
					COC	
Actinomyces israelii		88.32	0.42	CEL 93	1	
Blautia producta		11.68	0.07	NAG 6	99	
43737562	**Low discrimination**					**43737562**
Actinomyces israelii		87.25	0.45	CEL 93		
Bifidobacterium adolescentis		12.75	0.25	NAG 6		
43737600	**Good identification**					**43737600**
Actinomyces israelii		100.0	0.40	CEL 93		
43737602	**Good identification**					**43737602**
Actinomyces israelii		100.0	0.43	CEL 93		
43737642	**Good identification**					**43737642**
Actinomyces israelii		100.0	0.36	CEL 93		
43737700	**Low discrimination**					**43737700**
Actinomyces israelii		78.90	0.50	CEL 93		
Eubacterium contortum		21.10	0.26	RHA 7		
43737702	**Low discrimination**					**43737702**
Actinomyces israelii		69.74	0.53	CEL 93		
Bifidobacterium adolescentis		27.19	0.40	RHA 7		
Eubacterium contortum		2.19	0.13			
43737710	**Low discrimination**					**43737710**
					COC	
Actinomyces israelii		88.66	0.36	CEL 93	1	
Blautia producta		7.06	-.03	RHA 7	99	
Bifidobacterium adolescentis		4.29	0.07	MLZ 13	1	
43737712	**Low discrimination**					**43737712**
Bifidobacterium adolescentis		68.94	0.42	MNS 87		
Actinomyces israelii		21.62	0.39	RHA 1		
Clostridium clostridioforme		8.28	0.23			
43737742	**Low discrimination**					**43737742**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
				COC	
Actinomyces israelii	83.42	0.46	CEL 93	1	
Bifidobacterium adolescentis	10.34	0.25	RHA 7	1	
Blautia producta	6.24	0.07		99	
43741000 Good identification					**43741000**
				GRA	
Eubacterium tortuosum	97.12	0.41	MLT 1	99	
Fusobacterium mortiferum	2.53	0.31	CEL 10	1	
Actinomyces naeslundii	0.35	0.28		99	
43741100 Low discrimination					**43741100**
				GRA	
Fusobacterium mortiferum	62.65	0.37	MLT 11	1	
Actinomyces naeslundii	10.65	0.36	FRU 99	99	
Eubacterium tortuosum	10.42	0.07		99	
43741101 Good identification					**43741101**
Actinomyces naeslundii	100.0	0.39	ESL 12		
43741103 Good identification					**43741103**
Actinomyces naeslundii	100.0	0.39	ESL 12		
43741121 Good identification					**43741121**
Actinomyces naeslundii	100.0	0.39	ESL 12		
43741123 Low discrimination					**43741123**
				GRA	
Actinomyces naeslundii	88.83	0.39	ESL 12	99	
Prevotella oralis	11.17	-.03	MNS 88	1	
43741502 Good identification					**43741502**
Bifidobacterium adolescentis	96.90	0.36	MNS 87		
Actinomyces israelii	3.10	0.17	FRU 99		
43741512 Good identification					**43741512**
Bifidobacterium adolescentis	100.0	0.38	MNS 87		
43743001 Good identification					**43743001**
Actinomyces naeslundii	100.0	0.36	ESL 12		
43743003 Good identification					**43743003**
Actinomyces naeslundii	100.0	0.36	ESL 12		
43743021 Good identification					**43743021**
Actinomyces naeslundii	100.0	0.36	ESL 12		
43743023 Good identification					**43743023**
Actinomyces naeslundii	100.0	0.36	ESL 12		
43743100 Low discrimination					**43743100**
Actinomyces naeslundii	56.42	0.41	ESL 12		
Actinomyces israelii	36.83	0.28	MNS 88		
Bifidobacterium breve	6.75	0.10	FRU 95		
43743101 Good identification					**43743101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	ESL 12		
43743103	**Good identification**					**43743103**
Actinomyces naeslundii		100.0	0.44	ESL 12		
43743120	**Low discrimination**					**43743120**
					SPO	
Actinomyces naeslundii		75.64	0.41	ESL 12	1	
Bifidobacterium breve		16.07	0.15	MNS 88	1	
Clostridium ramosum		8.29	-.01	FRU 95	92	
43743121	**Good identification**					**43743121**
Actinomyces naeslundii		100.0	0.44	ESL 12		
43743123	**Good identification**					**43743123**
Actinomyces naeslundii		100.0	0.44	ESL 12		
43743512	**Good identification**					**43743512**
Bifidobacterium adolescentis		97.36	0.36	MNS 87		
Actinomyces israelii		2.64	0.16	FRU 99		
43745102	**Low discrimination**					**43745102**
Bifidobacterium adolescentis		68.95	0.36	MNS 87		
Actinomyces israelii		31.05	0.36	FRU 99		
43745112	**Acceptable identification**					**43745112**
Bifidobacterium adolescentis		94.78	0.38	MNS 87		
Actinomyces israelii		5.22	0.22	FRU 99		
43745502	**Acceptable identification**					**43745502**
					SPO	
Bifidobacterium adolescentis		93.46	0.50	MNS 87	1	
Actinomyces israelii		5.15	0.35	FRU 99	1	
Clostridium butyricum		1.40	0.04		99	
43745512	**Good identification**					**43745512**
Bifidobacterium adolescentis		99.33	0.52	MNS 87		
Actinomyces israelii		0.67	0.21	FRU 99		
43745552	**Good identification**					**43745552**
Bifidobacterium adolescentis		97.93	0.36	MNS 87		
Actinomyces israelii		2.07	0.14	FRU 99		
43747000	**Good identification**					**43747000**
Actinomyces israelii		100.0	0.36	FRU 99		
43747002	**Good identification**					**43747002**
Actinomyces israelii		100.0	0.39	FRU 99		
43747100	**Good identification**					**43747100**
Actinomyces israelii		100.0	0.46	FRU 99		
43747102	**Low discrimination**					**43747102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		75.72	0.49	FRU 99		
Bifidobacterium adolescentis		24.28	0.35			
43747112	**Low discrimination**					**43747112**
Bifidobacterium adolescentis		72.40	0.36	MNS 87		
Actinomyces israelii		27.61	0.35	FRU 99		
43747140	**Good identification**					**43747140**
Actinomyces israelii		100.0	0.39	FRU 99		
43747142	**Acceptable identification**					**43747142**
Actinomyces israelii		90.75	0.42	FRU 99		
Bifidobacterium adolescentis		9.25	0.19			
43747402	**Acceptable identification**					**43747402**
Actinomyces israelii		90.42	0.38	FRU 99		
Bifidobacterium adolescentis		9.58	0.15			
43747500	**Acceptable identification**					**43747500**
					SPO	
Actinomyces israelii		94.32	0.45	FRU 99	1	
Bifidobacterium adolescentis		3.75	0.15		1	
Clostridium butyricum		1.93	-.05		99	
43747502	**Low discrimination**					**43747502**
Bifidobacterium adolescentis		72.40	0.49	MNS 87		
Actinomyces israelii		27.61	0.48	FRU 99		
43747512	**Good identification**					**43747512**
Bifidobacterium adolescentis		95.07	0.50	MNS 87		
Actinomyces israelii		4.43	0.34	FRU 99		
Bifidobacterium dentium		0.50	-.07			
43747540	**Good identification**					**43747540**
Actinomyces israelii		100.0	0.38	FRU 99		
43747542	**Low discrimination**					**43747542**
Actinomyces israelii		54.54	0.41	FRU 99		
Bifidobacterium adolescentis		45.46	0.33			
43750000	**Very good identification**					**43750000**
					GRA	
Fusobacterium mortiferum		99.83	0.62	MLT 11	1	
Actinomyces naeslundii		0.17	0.26		99	
43750002	**Good identification**					**43750002**
					GRA	
Fusobacterium mortiferum		98.52	0.46	MLT 11	1	
Actinomyces naeslundii		1.48	0.26	BGL 10	99	
43750100	**Very good identification**					**43750100**
					GRA	
Fusobacterium mortiferum		99.48	0.68	MLT 11	1	
Actinomyces naeslundii		0.20	0.34		99	
Collinsella aerofaciens		0.20	-.03		99	
43750102	**Low discrimination**					**43750102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		86.62	0.52	MLT 11	1		
Bifidobacterium breve		4.71	0.27	BGL 10	99		
Bifidobacterium adolescentis		2.57	0.22		99		
43750103	**Low discrimination**						**43750103**
					GRA		
Actinomyces naeslundii		73.15	0.37	ESL 12	99		
Fusobacterium mortiferum		26.85	0.18	MNS 88	1		
43750121	**Good identification**						**43750121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
43750123	**Low discrimination**						**43750123**
					GRA		
Actinomyces naeslundii		86.29	0.37	ESL 12	99		
Prevotella oralis		13.71	-.03	MNS 88	1		
43750502	**Acceptable identification**						**43750502**
					GRA		
Bifidobacterium adolescentis		92.34	0.36	MNS 87	99		
Fusobacterium mortiferum		4.70	0.18	GAL 99	1		
Actinomyces israelii		2.96	0.17	XYL 87	99		
43750512	**Good identification**						**43750512**
Bifidobacterium adolescentis		95.98	0.38	MNS 87			
Clostridium clostridioforme		4.02	0.11	GAL 99			
43751000	**Good identification**						**43751000**
					GRA		
Fusobacterium mortiferum		90.57	0.65	MLT 11	1		
Lactobacillus catenaformis		2.57	0.29		99		
Actinomyces naeslundii		2.42	0.50		99		
43751001	**Low discrimination**						**43751001**
					GRA		
Actinomyces naeslundii		79.88	0.53	ESL 12	99		
Fusobacterium mortiferum		20.12	0.31	MNS 88	1		
43751002	**Low discrimination**						**43751002**
					GRA		
Fusobacterium mortiferum		40.76	0.49	MLT 11	1		
Actinomyces israelii		12.20	0.42	BGL 10	99		
Lactobacillus catenaformis		10.42	0.29		99		
43751003	**Good identification**						**43751003**
					GRA		
Actinomyces naeslundii		97.28	0.53	ESL 12	99		
Fusobacterium mortiferum		2.72	0.15	MNS 88	1		
43751020	**Low discrimination**						**43751020**
					GRA	SPO	
Actinomyces naeslundii		26.80	0.50	ESL 12	99	1	
Prevotella oralis		24.93	0.24	MNS 88	1	1	
Clostridium paraputrificum		13.10	0.13		99	99	
43751021	**Good identification**						**43751021**
Actinomyces naeslundii		100.0	0.53	ESL 12			
43751022	**Low discrimination**						**43751022**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		81.64	0.45	MNS 99	1	1	
Clostridium barati		4.95	0.24	LAC 94	99	99	
Actinomyces naeslundii		4.62	0.50		99	1	
43751023	**Low discrimination**						**43751023**
					GRA		
Actinomyces naeslundii		89.36	0.53	ESL 12	99		
Prevotella oralis		10.64	0.11	MNS 88	1		
43751041	**Good identification**						**43751041**
Actinomyces naeslundii		100.0	0.40	ESL 12			
43751042	**Low discrimination**						**43751042**
Actinomyces israelii		42.20	0.35	XYL 92			
Bifidobacterium breve		30.67	0.27				
Actinomyces naeslundii		15.40	0.37				
43751043	**Good identification**						**43751043**
Actinomyces naeslundii		100.0	0.40	ESL 12			
43751060	**Low discrimination**						**43751060**
					SPO		
Actinomyces naeslundii		57.64	0.37	ESL 12	1		
Bifidobacterium breve		20.18	0.15	MNS 88	1		
Clostridium paraputrificum		12.02	-.07		99		
43751061	**Good identification**						**43751061**
Actinomyces naeslundii		100.0	0.40	ESL 12			
43751063	**Good identification**						**43751063**
Actinomyces naeslundii		100.0	0.40	ESL 12			
43751100	**Low discrimination**						**43751100**
					GRA		
Fusobacterium mortiferum		77.65	0.71	MLT 11	1		
Collinsella aerofaciens		10.32	0.32		99		
Bifidobacterium breve		3.06	0.44		99		
43751101	**Low discrimination**						**43751101**
					GRA		
Actinomyces naeslundii		80.01	0.61	ESL 12	99		
Fusobacterium mortiferum		16.52	0.37	MNS 88	1		
Collinsella aerofaciens		2.20	-.03		99		
43751102	**Low discrimination**						**43751102**
					GRA		
Bifidobacterium breve		35.20	0.62	MNS 99	99		
Bifidobacterium adolescentis		19.20	0.56		99		
Prevotella oralis		14.74	0.44		1		
43751103	**Low discrimination**						**43751103**
Actinomyces naeslundii		82.25	0.61	ESL 12			
Bifidobacterium breve		6.77	0.27	MNS 88			
Bifidobacterium adolescentis		3.69	0.22				
43751110	**Low discrimination**						**43751110**
					GRA		
Bifidobacterium breve		64.34	0.44	MNS 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		16.48	0.37	BGL 91	1		
Actinomyces israelii		9.26	0.35		99		
43751111	**Low discrimination**						**43751111**
Actinomyces naeslundii		86.60	0.39	ESL 12			
Bifidobacterium breve		13.40	0.10	MNS 88			
43751112	**Low discrimination**						**43751112**
					GRA		
Bifidobacterium breve		55.94	0.62	MNS 99	99		
Bifidobacterium adolescentis		37.29	0.58		99		
Leptotrichia buccalis		4.71	0.24		1		
43751120	**Low discrimination**						**43751120**
					GRA	SPO	
Clostridium paraputrificum		59.68	0.47	MNS 99	99	99	
Prevotella oralis		17.97	0.44	RAF 7	1	1	
Bifidobacterium breve		7.55	0.49		99	1	
43751121	**Low discrimination**						**43751121**
					GRA	SPO	
Actinomyces naeslundii		85.32	0.61	ESL 12	99	1	
Clostridium paraputrificum		9.77	0.13	MNS 88	99	99	
Prevotella oralis		2.94	0.10		1	1	
43751122	**Low discrimination**						**43751122**
					GRA	SPO	
Prevotella oralis		71.88	0.66	MNS 99	1	1	
Bifidobacterium breve		16.06	0.66		99	1	
Clostridium paraputrificum		5.38	0.40		99	99	
43751123	**Low discrimination**						**43751123**
					GRA		
Actinomyces naeslundii		52.80	0.61	ESL 12	99		
Prevotella oralis		34.59	0.32	MNS 88	1		
Bifidobacterium breve		7.73	0.32		99		
43751130	**Low discrimination**						**43751130**
					SPO		
Bifidobacterium breve		86.99	0.49	MNS 99	1		
Clostridium paraputrificum		6.95	0.13	BGL 91	99		
Actinomyces naeslundii		2.13	0.36		1		
43751131	**Low discrimination**						**43751131**
Actinomyces naeslundii		78.43	0.39	ESL 12			
Bifidobacterium breve		21.57	0.15	MNS 88			
43751132	**Good identification**						**43751132**
					GRA	CAT	
Bifidobacterium breve		92.96	0.66	MNS 99	99	1	
Prevotella oralis		4.20	0.32		1	1	
Capnocytophaga ochracea		0.82	0.20		1	90	
43751140	**Low discrimination**						**43751140**
					GRA		
Bifidobacterium breve		54.63	0.44	MNS 99	99		
Actinomyces israelii		20.47	0.42	BGL 91	99		
Fusobacterium mortiferum		13.99	0.37		1		
43751141	**Good identification**						**43751141**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		95.59	0.48	ESL 12			
Bifidobacterium breve		4.41	0.10	MNS 88			
43751142	**Low discrimination**						**43751142**
Bifidobacterium breve		85.48	0.62	MNS 99			
Bifidobacterium adolescentis		5.76	0.41				
Actinomyces israelii		4.75	0.45				
43751143	**Low discrimination**						**43751143**
Actinomyces naeslundii		68.19	0.48	ESL 12			
Bifidobacterium breve		31.81	0.27	MNS 88			
43751150	**Acceptable identification**						**43751150**
Bifidobacterium breve		93.28	0.44	MNS 99			
Actinomyces israelii		5.22	0.28	BGL 91			
Bifidobacterium adolescentis		0.79	0.08				
43751152	**Good identification**						**43751152**
Bifidobacterium breve		91.55	0.62	MNS 99			
Bifidobacterium adolescentis		7.54	0.42				
Actinomyces israelii		0.76	0.31				
43751160	**Low discrimination**						**43751160**
					SPO		
Bifidobacterium breve		57.27	0.49	MNS 99	1		
Clostridium paraputrificum		34.09	0.27	BGL 91	99		
Actinomyces naeslundii		4.70	0.45		1		
43751161	**Low discrimination**						**43751161**
					SPO		
Actinomyces naeslundii		88.43	0.48	ESL 12	1		
Bifidobacterium breve		7.25	0.15	MNS 88	1		
Clostridium paraputrificum		4.32	-.07		99		
43751162	**Good identification**						**43751162**
					GRA	SPO	
Bifidobacterium breve		90.41	0.66	MNS 99	99	1	
Prevotella oralis		4.09	0.32		1	1	
Clostridium paraputrificum		2.28	0.21		99	99	
43751163	**Low discrimination**						**43751163**
					GRA		
Actinomyces naeslundii		53.56	0.48	ESL 12	99		
Bifidobacterium breve		44.43	0.32	MNS 88	99		
Prevotella oralis		2.01	-.03		1		
43751170	**Good identification**						**43751170**
					SPO		
Bifidobacterium breve		98.98	0.49	MNS 99	1		
Clostridium paraputrificum		0.60	-.07	BGL 91	99		
Actinomyces naeslundii		0.43	0.23		1		
43751172	**Very good identification**						**43751172**
					GRA	CAT	
Bifidobacterium breve		99.77	0.66	MNS 99	99	1	
Capnocytophaga ochracea		0.10	0.04		1	90	
Bifidobacterium adolescentis		0.05	0.08		99	1	
43751222	**Good identification**						**43751222**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Prevotella oralis		97.93	0.39	MNS 99	1	1	
Clostridium barati		1.21	0.05	LAC 94	99	99	
Lactobacillus acidophilus		0.48	0.01		99	1	
43751302	**Low discrimination**						**43751302**
					GRA		
Prevotella oralis		77.25	0.38	MNS 99	1		
Lactobacillus acidophilus		5.32	0.19	NAG 95	99		
Actinomyces israelii		4.63	0.33		99		
43751320	**Low discrimination**						**43751320**
					SPO	GRA	
Prevotella oralis		78.95	0.38	MNS 99	1	1	
Clostridium ramosum		9.33	0.22	BGL 95	92	50	
Clostridium paraputrificum		6.18	0.13		99	99	
43751322	**Very good identification**						**43751322**
					SPO		
Prevotella oralis		95.20	0.60	MNS 99	1		
Clostridium ramosum		2.37	0.33		92		
Parabacteroides distasonis		1.07	0.39		1		
43751400	**Low discrimination**						**43751400**
					GRA	SPO	
Actinomyces israelii		61.77	0.38	XYL 92	99	1	
Fusobacterium mortiferum		34.40	0.31		1	1	
Clostridium butyricum		3.83	-.04		99	99	
43751402	**Low discrimination**						**43751402**
					GRA		
Bifidobacterium adolescentis		54.77	0.36	MNS 87	99		
Actinomyces israelii		43.44	0.41	XYL 87	99		
Fusobacterium mortiferum		1.79	0.15	LAC 99	1		
43751412	**Low discrimination**						**43751412**
Bifidobacterium adolescentis		88.15	0.38	MNS 87			
Actinomyces israelii		8.55	0.27	XYL 87			
Clostridium clostridioforme		3.31	0.10	LAC 99			
43751500	**Low discrimination**						**43751500**
					SPO		
Actinomyces israelii		44.71	0.48	XYL 92	1		
Bifidobacterium adolescentis		21.15	0.36		1		
Clostridium butyricum		16.64	0.20		99		
43751502	**Very good identification**						**43751502**
					SPO		
Bifidobacterium adolescentis		95.99	0.70	MNS 87	1		
Actinomyces israelii		3.08	0.51	XYL 87	1		
Clostridium butyricum		0.30	0.13		99		
43751503	**Acceptable identification**						**43751503**
Bifidobacterium adolescentis		93.84	0.36	MNS 87			
Actinomyces naeslundii		3.15	0.26	XYL 87			
Actinomyces israelii		3.01	0.17	URE 1			
43751510	**Low discrimination**						**43751510**
					SPO		
Bifidobacterium adolescentis		76.07	0.38	MNS 87	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		19.67	0.34	XYL 87	1	
Clostridium butyricum		2.58	-.02	BGL 99	99	
43751512	**Very good identification**					**43751512**
Bifidobacterium adolescentis		99.24	0.72	MNS 87		
Actinomyces israelii		0.39	0.37	XYL 87		
Bifidobacterium breve		0.23	0.27			
43751513	**Good identification**					**43751513**
Bifidobacterium adolescentis		100.0	0.38	MNS 87		
43751532	**Low discrimination**					**43751532**
Bifidobacterium adolescentis		57.14	0.38	MNS 87		
Bifidobacterium breve		22.77	0.32	XYL 87		
Clostridium clostridioforme		7.18	0.19	NAG 1		
43751540	**Low discrimination**					**43751540**
					SPO	
Actinomyces israelii		78.96	0.41	XYL 92	1	
Bifidobacterium adolescentis		11.87	0.21		1	
Clostridium butyricum		6.57	0.01		99	
43751542	**Low discrimination**					**43751542**
Bifidobacterium adolescentis		88.86	0.55	MNS 87		
Actinomyces israelii		8.96	0.44	XYL 87		
Bifidobacterium breve		1.99	0.27	SOR 11		
43751552	**Very good identification**					**43751552**
Bifidobacterium adolescentis		97.03	0.56	MNS 87		
Bifidobacterium breve		1.78	0.27	XYL 87		
Actinomyces israelii		1.20	0.30	SOR 11		
43751702	**Low discrimination**					**43751702**
Bifidobacterium adolescentis		59.74	0.36	MNS 87		
Clostridium clostridioforme		14.78	0.23	XYL 87		
Actinomyces israelii		14.26	0.32	RHA 1		
43751712	**Low discrimination**					**43751712**
Bifidobacterium adolescentis		51.25	0.38	MNS 87		
Clostridium clostridioforme		47.25	0.34	XYL 87		
Actinomyces israelii		1.50	0.18	RHA 1		
43752101	**Good identification**					**43752101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43752103	**Good identification**					**43752103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43752120	**Low discrimination**					**43752120**
					SPO	
Actinomyces naeslundii		71.08	0.39	ESL 12	1	
Bifidobacterium breve		19.08	0.15	MNS 88	1	
Clostridium ramosum		9.84	-.01	GAL 96	92	
43752121	**Good identification**					**43752121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	ESL 12		
43752123	**Good identification**					**43752123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
43752512	**Good identification**					**43752512**
Bifidobacterium adolescentis		97.36	0.36	MNS 87		
Actinomyces israelii		2.64	0.16	GAL 99		
43753000	**Low discrimination**					**43753000**
					GRA	
Actinomyces israelii		66.62	0.52	XYL 92	99	
Actinomyces naeslundii		27.53	0.55		99	
Fusobacterium mortiferum		5.36	0.31		1	
43753001	**Very good identification**					**43753001**
Actinomyces naeslundii		98.40	0.58	ESL 12		
Actinomyces israelii		1.60	0.18	MNS 88		
43753002	**Low discrimination**					**43753002**
Actinomyces israelii		73.44	0.55	XYL 92		
Actinomyces naeslundii		20.23	0.55			
Bifidobacterium breve		3.66	0.27			
43753003	**Very good identification**					**43753003**
Actinomyces naeslundii		97.61	0.58	ESL 12		
Actinomyces israelii		2.39	0.21	MNS 88		
43753010	**Low discrimination**					**43753010**
Actinomyces israelii		83.68	0.38	XYL 92		
Actinomyces naeslundii		12.18	0.33	MLZ 13		
Bifidobacterium breve		4.15	0.10			
43753011	**Good identification**					**43753011**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43753012	**Low discrimination**					**43753012**
Actinomyces israelii		62.00	0.41	XYL 92		
Bifidobacterium breve		20.70	0.27	MLZ 13		
Bifidobacterium adolescentis		11.29	0.22			
43753013	**Good identification**					**43753013**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43753020	**Low discrimination**					**43753020**
Actinomyces naeslundii		82.05	0.55	ESL 12		
Actinomyces israelii		12.67	0.32	MNS 88		
Bifidobacterium breve		2.61	0.15			
43753021	**Very good identification**					**43753021**
Actinomyces naeslundii		100.0	0.58	ESL 12		
43753022	**Low discrimination**					**43753022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		56.41	0.55	ESL 12		
Bifidobacterium breve		18.16	0.32	MNS 88		
Actinomyces israelii		13.07	0.35			
43753023	**Very good identification**					**43753023**
Actinomyces naeslundii		100.0	0.58	ESL 12		
43753031	**Good identification**					**43753031**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43753033	**Good identification**					**43753033**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43753040	**Low discrimination**					**43753040**
Actinomyces israelii		82.88	0.45	XYL 92		
Actinomyces naeslundii		15.54	0.42			
Bifidobacterium breve		1.58	0.10			
43753041	**Good identification**					**43753041**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43753042	**Low discrimination**					**43753042**
Actinomyces israelii		79.79	0.48	XYL 92		
Bifidobacterium breve		10.24	0.27			
Actinomyces naeslundii		9.98	0.42			
43753043	**Acceptable identification**					**43753043**
Actinomyces naeslundii		94.89	0.45	ESL 12		
Actinomyces israelii		5.11	0.14	MNS 88		
43753060	**Low discrimination**					**43753060**
Actinomyces naeslundii		65.76	0.42	ESL 12		
Actinomyces israelii		22.38	0.25	MNS 88		
Bifidobacterium breve		11.86	0.15			
43753061	**Good identification**					**43753061**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43753063	**Good identification**					**43753063**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43753100	**Low discrimination**					**43753100**
Actinomyces israelii		64.85	0.62	XYL 92		
Actinomyces naeslundii		19.07	0.62			
Bifidobacterium breve		11.88	0.44			
43753101	**Very good identification**					**43753101**
Actinomyces naeslundii		97.36	0.65	ESL 12		
Actinomyces israelii		2.23	0.28	MNS 88		
Bifidobacterium breve		0.41	0.10			
43753102	**Low discrimination**					**43753102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		41.01	0.62	MNS 99		
Actinomyces israelii		33.21	0.65			
Bifidobacterium adolescentis		18.30	0.55			
43753103	**Good identification**					**43753103**
Actinomyces naeslundii		91.27	0.65	ESL 12		
Bifidobacterium breve		3.87	0.27	MNS 88		
Actinomyces israelii		3.14	0.31			
43753110	**Low discrimination**					**43753110**
Bifidobacterium breve		51.12	0.44	MNS 99		
Actinomyces israelii		41.71	0.48	BGL 91		
Actinomyces naeslundii		4.32	0.41			
43753111	**Low discrimination**					**43753111**
Actinomyces naeslundii		87.36	0.44	ESL 12		
Bifidobacterium breve		6.96	0.10	MNS 88		
Actinomyces israelii		5.68	0.14	MLZ 5		
43753112	**Low discrimination**					**43753112**
Bifidobacterium breve		59.62	0.62	MNS 99		
Bifidobacterium adolescentis		32.52	0.56			
Actinomyces israelii		7.22	0.51			
43753113	**Low discrimination**					**43753113**
Actinomyces naeslundii		42.68	0.44	ESL 12		
Bifidobacterium breve		34.40	0.27	MNS 88		
Bifidobacterium adolescentis		18.76	0.22	MLZ 5		
43753120	**Low discrimination**					**43753120**
					SPO	
Bifidobacterium breve		36.43	0.49	MNS 99	1	
Actinomyces naeslundii		32.89	0.62	BGL 91	1	
Clostridium ramosum		18.79	0.33		92	
43753121	**Very good identification**					**43753121**
					SPO	
Actinomyces naeslundii		98.88	0.65	ESL 12	1	
Bifidobacterium breve		0.74	0.15	MNS 88	1	
Clostridium ramosum		0.38	-.01		92	
43753122	**Low discrimination**					**43753122**
					SPO	
Bifidobacterium breve		72.15	0.66	MNS 99	1	
Clostridium ramosum		14.72	0.43		92	
Actinomyces naeslundii		6.44	0.62		1	
43753123	**Good identification**					**43753123**
					SPO	
Actinomyces naeslundii		91.39	0.65	ESL 12	1	
Bifidobacterium breve		6.89	0.32	MNS 88	1	
Clostridium ramosum		1.41	0.09		92	
43753130	**Acceptable identification**					**43753130**
Bifidobacterium breve		91.64	0.49	MNS 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		4.36	0.41	BGL 91		
Actinomyces israelii		2.68	0.28			
43753131	**Low discrimination**					**43753131**
Actinomyces naeslundii		87.59	0.44	ESL 12		
Bifidobacterium breve		12.41	0.15	MNS 88		
43753132	**Very good identification**					**43753132**
Bifidobacterium breve		98.29	0.66	MNS 99		
Actinomyces naeslundii		0.46	0.41			
Actinomyces israelii		0.43	0.31			
43753140	**Low discrimination**					**43753140**
Actinomyces israelii		62.33	0.55	XYL 92		
Bifidobacterium breve		29.36	0.44			
Actinomyces naeslundii		8.32	0.50			
43753141	**Acceptable identification**					**43753141**
Actinomyces naeslundii		93.09	0.53	ESL 12		
Actinomyces israelii		4.70	0.21	MNS 88		
Bifidobacterium breve		2.21	0.10			
43753142	**Low discrimination**					**43753142**
Bifidobacterium breve		71.47	0.62	MNS 99		
Actinomyces israelii		22.51	0.58			
Bifidobacterium adolescentis		3.94	0.39			
43753143	**Low discrimination**					**43753143**
Actinomyces naeslundii		75.99	0.53	ESL 12		
Bifidobacterium breve		18.26	0.27	MNS 88		
Actinomyces israelii		5.75	0.24			
43753150	**Low discrimination**					**43753150**
Bifidobacterium breve		75.07	0.44	MNS 99		
Actinomyces israelii		23.82	0.41	BGL 91		
Actinomyces naeslundii		1.12	0.28			
43753152	**Low discrimination**					**43753152**
Bifidobacterium breve		89.61	0.62	MNS 99		
Bifidobacterium adolescentis		6.04	0.41			
Actinomyces israelii		4.22	0.44			
43753160	**Low discrimination**					**43753160**
Bifidobacterium breve		80.21	0.49	MNS 99		
Actinomyces naeslundii		12.78	0.50	BGL 91		
Actinomyces israelii		6.11	0.35			
43753161	**Good identification**					**43753161**
Actinomyces naeslundii		95.95	0.53	ESL 12		
Bifidobacterium breve		4.05	0.15	MNS 88		
43753162	**Very good identification**					**43753162**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		97.07	0.66	MNS 99			
Actinomyces naeslundii		1.53	0.50				
Actinomyces israelii		1.10	0.38				
43753163	**Low discrimination**						**43753163**
Actinomyces naeslundii		70.06	0.53	ESL 12			
Bifidobacterium breve		29.94	0.32	MNS 88			
43753170	**Good identification**						**43753170**
Bifidobacterium breve		98.06	0.49	MNS 99			
Actinomyces israelii		1.12	0.21	BGL 91			
Actinomyces naeslundii		0.82	0.28				
43753172	**Very good identification**						**43753172**
Bifidobacterium breve		99.75	0.66	MNS 99			
Actinomyces israelii		0.17	0.24				
Actinomyces naeslundii		0.08	0.28				
43753202	**Good identification**						**43753202**
Actinomyces israelii		100.0	0.36	XYL 92			
43753300	**Acceptable identification**						**43753300**
					SPO		
Actinomyces israelii		91.41	0.43	XYL 92	1		
Actinomyces naeslundii		3.61	0.28	RHA 7	1		
Clostridium ramosum		2.73	0.01		92		
43753302	**Low discrimination**						**43753302**
					SPO		
Actinomyces israelii		72.71	0.46	XYL 92	1		
Bifidobacterium breve		12.05	0.27	RHA 7	1		
Clostridium ramosum		5.79	0.11		92		
43753320	**Acceptable identification**						**43753320**
					SPO	GRA	
Clostridium ramosum		93.92	0.35	MNS 99	92	50	
Actinomyces israelii		2.03	0.23	TRE 94	1	99	
Prevotella oralis		1.42	0.04		1	1	
43753322	**Low discrimination**						**43753322**
					SPO	GRA	
Clostridium ramosum		89.25	0.45	MNS 99	92	50	
Prevotella oralis		6.39	0.25	TRE 94	1	1	
Bifidobacterium breve		3.33	0.32		1	99	
43753340	**Acceptable identification**						**43753340**
Actinomyces israelii		94.05	0.36	XYL 92			
Bifidobacterium breve		5.95	0.10	RHA 7			
43753342	**Low discrimination**						**43753342**
Actinomyces israelii		70.12	0.39	XYL 92			
Bifidobacterium breve		29.88	0.27	RHA 7			
43753400	**Good identification**						**43753400**
Actinomyces israelii		100.0	0.51	XYL 92			
43753402	**Low discrimination**						**43753402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		84.60	0.54	XYL 92		
Bifidobacterium adolescentis		15.40	0.35			
43753410	**Good identification**					**43753410**
Actinomyces israelii		100.0	0.36	XYL 92		
43753412	**Low discrimination**					**43753412**
Bifidobacterium adolescentis		59.83	0.36	MNS 87		
Actinomyces israelii		40.17	0.39	XYL 87		
43753440	**Good identification**					**43753440**
Actinomyces israelii		100.0	0.44	XYL 92		
43753442	**Acceptable identification**					**43753442**
Actinomyces israelii		94.53	0.47	XYL 92		
Bifidobacterium adolescentis		5.47	0.19			
43753500	**Good identification**					**43753500**
					SPO	
Actinomyces israelii		92.47	0.61	XYL 92	1	
Bifidobacterium adolescentis		6.31	0.35		1	
Clostridium butyricum		0.68	0.03		99	
43753502	**Low discrimination**					**43753502**
Bifidobacterium adolescentis		81.55	0.69	MNS 87		
Actinomyces israelii		18.10	0.64	XYL 87		
Bifidobacterium breve		0.28	0.27			
43753510	**Low discrimination**					**43753510**
Actinomyces israelii		63.55	0.47	XYL 92		
Bifidobacterium adolescentis		35.49	0.36	MLZ 13		
Bifidobacterium breve		0.96	0.10			
43753512	**Very good identification**					**43753512**
Bifidobacterium adolescentis		97.06	0.70	MNS 87		
Actinomyces israelii		2.63	0.50	XYL 87		
Bifidobacterium breve		0.27	0.27			
43753513	**Good identification**					**43753513**
Bifidobacterium adolescentis		97.36	0.36	MNS 87		
Actinomyces israelii		2.64	0.16	XYL 87		
43753520	**Low discrimination**					**43753520**
Actinomyces israelii		86.78	0.40	XYL 92		
Bifidobacterium breve		5.47	0.15	NAG 6		
Actinomyces naeslundii		4.94	0.28			
43753522	**Low discrimination**					**43753522**
Actinomyces israelii		43.83	0.43	XYL 92		
Bifidobacterium adolescentis		31.26	0.35	NAG 6		
Bifidobacterium breve		18.61	0.32			
43753532	**Low discrimination**					**43753532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		58.09	0.36	MNS 87			
Bifidobacterium breve		28.29	0.32	XYL 87			
Actinomyces israelii		9.96	0.29	NAG 1			
43753540	**Good identification**						**43753540**
Actinomyces israelii		97.32	0.54	XYL 92			
Bifidobacterium adolescentis		2.11	0.19				
Bifidobacterium breve		0.57	0.10				
43753542	**Low discrimination**						**43753542**
Bifidobacterium adolescentis		57.95	0.53	MNS 87			
Actinomyces israelii		40.46	0.57	XYL 87			
Bifidobacterium breve		1.59	0.27	SOR 11			
43753550	**Low discrimination**						**43753550**
					COC		
Actinomyces israelii		78.95	0.40	XYL 92	1		
Bifidobacterium adolescentis		14.01	0.21	MLZ 13	1		
Blautia producta		3.96	-.03		99		
43753552	**Acceptable identification**						**43753552**
Bifidobacterium adolescentis		90.27	0.55	MNS 87			
Actinomyces israelii		7.71	0.43	XYL 87			
Bifidobacterium breve		2.02	0.27	SOR 11			
43753700	**Good identification**						**43753700**
Actinomyces israelii		100.0	0.42	XYL 92			
43753702	**Low discrimination**						**43753702**
Actinomyces israelii		62.32	0.45	XYL 92			
Bifidobacterium adolescentis		37.69	0.35	RHA 7			
43753712	**Low discrimination**						**43753712**
Bifidobacterium adolescentis		83.18	0.36	MNS 87			
Actinomyces israelii		16.82	0.31	XYL 87			
43753742	**Low discrimination**						**43753742**
Actinomyces israelii		83.88	0.38	XYL 92			
Bifidobacterium adolescentis		16.12	0.19	RHA 7			
43754000	**Good identification**						**43754000**
					GRA		
Fusobacterium mortiferum		95.12	0.43	MLT 11	1		
Actinomyces israelii		4.88	0.23	XYL 7	99		
43754100	**Acceptable identification**						**43754100**
					GRA	SPO	
Fusobacterium mortiferum		91.21	0.49	MLT 11	1	1	
Actinomyces israelii		8.02	0.33	XYL 7	99	1	
Clostridium butyricum		0.76	-.05		99	99	
43754102	**Low discrimination**						**43754102**
					GRA		
Bifidobacterium adolescentis		54.66	0.36	MNS 87	99		
Actinomyces israelii		24.62	0.36	GAL 99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		20.73	0.33	ARA 87	1		
43754112	**Acceptable identification**						**43754112**
Bifidobacterium adolescentis		92.41	0.38	MNS 87			
Actinomyces israelii		5.09	0.22	GAL 99			
Clostridium clostridioforme		2.50	0.08	ARA 87			
43754502	**Acceptable identification**						**43754502**
Bifidobacterium adolescentis		92.00	0.50	MNS 87			
Actinomyces israelii		5.07	0.35	GAL 99			
Clostridium clostridioforme		1.56	0.17				
43754512	**Acceptable identification**						**43754512**
Bifidobacterium adolescentis		93.45	0.52	MNS 87			
Clostridium clostridioforme		5.92	0.28	GAL 99			
Actinomyces israelii		0.63	0.21				
43754552	**Good identification**						**43754552**
Bifidobacterium adolescentis		97.93	0.36	MNS 87			
Actinomyces israelii		2.07	0.14	GAL 99			
43754712	**Good identification**						**43754712**
Clostridium clostridioforme		97.87	0.43	MNS 99			
Bifidobacterium adolescentis		2.13	0.18	TRE 88			
43754732	**Good identification**						**43754732**
Clostridium clostridioforme		99.19	0.43	MNS 99			
Parabacteroides distasonis		0.81	0.13	TRE 88			
43755000	**Low discrimination**						**43755000**
					GRA	SPO	
Actinomyces israelii		76.27	0.57		99	1	
Fusobacterium mortiferum		22.52	0.46		1	1	
Clostridium butyricum		1.21	0.05		99	99	
43755002	**Low discrimination**						**43755002**
					GRA		
Actinomyces israelii		88.99	0.60		99		
Bifidobacterium adolescentis		7.98	0.36		99		
Fusobacterium mortiferum		1.95	0.30		1		
43755010	**Good identification**						**43755010**
					GRA		
Actinomyces israelii		98.04	0.43	MLZ 13	99		
Fusobacterium mortiferum		1.96	0.12		1		
43755012	**Low discrimination**						**43755012**
Actinomyces israelii		57.09	0.46	MLZ 13			
Bifidobacterium adolescentis		41.90	0.38				
Clostridium clostridioforme		1.02	0.07				
43755020	**Low discrimination**						**43755020**
					GRA		
Actinomyces israelii		80.93	0.37	NAG 6	99		
Prevotella oralis		15.29	0.08		1		
Fusobacterium mortiferum		3.78	0.12		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43755040	**Good identification**						**43755040**
					GRA		
Actinomyces israelii		99.24	0.50		99		
Fusobacterium mortiferum		0.76	0.12		1		
43755042	**Good identification**						**43755042**
Actinomyces israelii		97.23	0.53				
Bifidobacterium adolescentis		2.77	0.21				
43755050	**Good identification**						**43755050**
Actinomyces israelii		100.0	0.36	MLZ 13			
43755052	**Low discrimination**						**43755052**
Actinomyces israelii		81.09	0.39	MLZ 13			
Bifidobacterium adolescentis		18.91	0.22				
43755100	**Low discrimination**						**43755100**
					GRA	SPO	
Actinomyces israelii		74.31	0.68		99	1	
Fusobacterium mortiferum		12.80	0.52		1	1	
Clostridium butyricum		7.07	0.29		99	99	
43755102	**Low discrimination**						**43755102**
Bifidobacterium adolescentis		66.46	0.70	MNS 87			
Actinomyces israelii		29.93	0.71	ARA 87			
Bifidobacterium longum subsp. infantis		1.07	0.26				
43755103	**Low discrimination**						**43755103**
Bifidobacterium adolescentis		67.39	0.36	MNS 87			
Actinomyces israelii		30.35	0.36	ARA 87			
Actinomyces naeslundii		2.27	0.26	URE 1			
43755110	**Low discrimination**						**43755110**
					SPO		
Actinomyces israelii		75.39	0.54	MLZ 13	1		
Bifidobacterium adolescentis		20.75	0.38		1		
Clostridium butyricum		2.53	0.07		99		
43755112	**Good identification**						**43755112**
Bifidobacterium adolescentis		94.45	0.72	MNS 87			
Actinomyces israelii		5.20	0.57	ARA 87			
Bifidobacterium breve		0.21	0.27				
43755113	**Acceptable identification**						**43755113**
Bifidobacterium adolescentis		94.78	0.38	MNS 87			
Actinomyces israelii		5.22	0.22	ARA 87			
43755120	**Low discrimination**						**43755120**
					GRA	SPO	
Actinomyces israelii		38.86	0.47	NAG 6	99	1	
Prevotella oralis		28.76	0.28		1	1	
Clostridium paraputrificum		7.80	0.13		99	99	
43755122	**Low discrimination**						**43755122**
					CAT		
Prevotella oralis		63.42	0.50	MNS 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides fragilis		7.41	0.25	XYL 11	99	
Parabacteroides distasonis		7.39	0.47		76	
43755132	**Low discrimination**					**43755132**
					GRA	
Parabacteroides distasonis		38.53	0.42	MNS 98	1	
Bifidobacterium adolescentis		28.14	0.38		99	
Bifidobacterium breve		11.21	0.32		99	
43755140	**Very good identification**					**43755140**
					SPO	
Actinomyces israelii		96.27	0.61		1	
Clostridium butyricum		2.05	0.11		99	
Bifidobacterium adolescentis		1.03	0.21		1	
43755142	**Low discrimination**					**43755142**
Actinomyces israelii		57.90	0.64			
Bifidobacterium adolescentis		40.86	0.55			
Bifidobacterium breve		0.92	0.27			
43755150	**Acceptable identification**					**43755150**
Actinomyces israelii		90.65	0.46	MLZ 13		
Bifidobacterium adolescentis		7.93	0.22			
Bifidobacterium breve		1.42	0.10			
43755152	**Low discrimination**					**43755152**
Bifidobacterium adolescentis		83.92	0.56	MNS 87		
Actinomyces israelii		14.54	0.49	ARA 87		
Bifidobacterium breve		1.54	0.27	SOR 11		
43755160	**Low discrimination**					**43755160**
					SPO	
Actinomyces israelii		88.25	0.40	NAG 6	1	
Bifidobacterium breve		5.76	0.15		1	
Clostridium paraputrificum		3.43	-.07		99	
43755162	**Low discrimination**					**43755162**
					GRA	
Actinomyces israelii		50.65	0.43	NAG 6	99	
Bifidobacterium breve		22.29	0.32		99	
Prevotella oralis		12.33	0.16		1	
43755202	**Low discrimination**					**43755202**
Actinomyces israelii		76.65	0.41	RHA 7		
Eubacterium contortum		15.48	0.14			
Clostridium clostridioforme		4.36	0.10			
43755300	**Very good identification**					**43755300**
					SPO	
Eubacterium contortum		97.53	0.61	CEL 1	1	
Actinomyces israelii		2.30	0.48		1	
Clostridium butyricum		0.09	0.03		99	
43755302	**Low discrimination**					**43755302**
Eubacterium contortum		76.18	0.48	CEL 1		
Actinomyces israelii		15.24	0.51			
Bifidobacterium adolescentis		4.54	0.36			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43755312	**Low discrimination**					**43755312**
Bifidobacterium adolescentis		45.76	0.38	MNS 87		
Clostridium clostridioforme		27.23	0.31	RHA 1		
Actinomyces israelii		18.77	0.37	ARA 87		
43755320	**Acceptable identification**					**43755320**
					GRA	
Eubacterium contortum		92.82	0.48	CEL 1	99	
Prevotella oralis		3.33	0.22		1	
Bacteroides vulgatus		2.54	0.14		1	
43755322	**Low discrimination**					**43755322**
					GRA	
Prevotella oralis		45.60	0.44	MNS 99	1	
Parabacteroides distasonis		37.19	0.55	XYL 11	1	
Eubacterium contortum		11.81	0.35		99	
43755332	**Low discrimination**					**43755332**
Parabacteroides distasonis		84.43	0.51	MNS 98		
Clostridium clostridioforme		12.23	0.31			
Prevotella oralis		1.94	0.10			
43755342	**Low discrimination**					**43755342**
Actinomyces israelii		81.66	0.44	RHA 7		
Eubacterium contortum		10.60	0.14			
Bifidobacterium adolescentis		7.73	0.21			
43755400	**Low discrimination**					**43755400**
					SPO	
Actinomyces israelii		84.10	0.56		1	
Clostridium butyricum		14.66	0.22		99	
Bifidobacterium adolescentis		0.94	0.16		1	
43755402	**Low discrimination**					**43755402**
					SPO	
Actinomyces israelii		55.86	0.59		1	
Bifidobacterium adolescentis		40.99	0.50		1	
Clostridium butyricum		2.53	0.15		99	
43755410	**Low discrimination**					**43755410**
					SPO	
Actinomyces israelii		86.78	0.42	MLZ 13	1	
Bifidobacterium adolescentis		7.89	0.18		1	
Clostridium butyricum		5.33	0.00		99	
43755412	**Low discrimination**					**43755412**
Bifidobacterium adolescentis		81.57	0.52	MNS 87		
Actinomyces israelii		13.59	0.45	LAC 99		
Clostridium clostridioforme		4.63	0.27			
43755420	**Low discrimination**					**43755420**
					SPO	
Actinomyces israelii		80.81	0.35	NAG 6	1	
Clostridium butyricum		19.19	0.04		99	
43755422	**Low discrimination**					**43755422**
					GRA	
Actinomyces israelii		62.26	0.38	NAG 6	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		15.77	0.06		1	
Clostridium clostridioforme		10.90	0.16		50	
43755440	**Good identification**					**43755440**
					SPO	
Actinomyces israelii		96.25	0.49		1	
Clostridium butyricum		3.75	0.04		99	
43755442	**Low discrimination**					**43755442**
					SPO	
Actinomyces israelii		80.43	0.52		1	
Bifidobacterium adolescentis		18.76	0.35		1	
Clostridium butyricum		0.81	-.03		99	
43755452	**Low discrimination**					**43755452**
Bifidobacterium adolescentis		65.61	0.36	MNS 87		
Actinomyces israelii		34.39	0.38	LAC 99		
43755500	**Low discrimination**					**43755500**
					SPO	
Clostridium butyricum		44.40	0.45	MNS 99	99	
Actinomyces israelii		42.45	0.66	TRE 94	1	
Bifidobacterium adolescentis		11.68	0.50		1	
43755502	**Good identification**					**43755502**
					SPO	
Bifidobacterium adolescentis		93.39	0.84	MNS 87	1	
Actinomyces israelii		5.14	0.69		1	
Clostridium butyricum		1.39	0.38		99	
43755503	**Acceptable identification**					**43755503**
					SPO	
Bifidobacterium adolescentis		93.46	0.50	MNS 87	1	
Actinomyces israelii		5.15	0.35	URE 1	1	
Clostridium butyricum		1.40	0.04		99	
43755510	**Low discrimination**					**43755510**
					SPO	
Bifidobacterium adolescentis		62.07	0.52	MNS 87	1	
Actinomyces israelii		27.58	0.52	BGL 99	1	
Clostridium butyricum		10.16	0.24		99	
43755512	**Very good identification**					**43755512**
Bifidobacterium adolescentis		99.07	0.86	MNS 87		
Actinomyces israelii		0.67	0.55			
Clostridium clostridioforme		0.19	0.36			
43755513	**Good identification**					**43755513**
Bifidobacterium adolescentis		99.33	0.52	MNS 87		
Actinomyces israelii		0.67	0.21	URE 1		
43755532	**Low discrimination**					**43755532**
					GRA	
Bifidobacterium adolescentis		71.58	0.52	MNS 87	99	
Clostridium clostridioforme		13.60	0.36	NAG 1	50	
Parabacteroides distasonis		8.98	0.39		1	
43755540	**Low discrimination**					**43755540**
					SPO	
Actinomyces israelii		75.68	0.59		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		17.70	0.27		99	
Bifidobacterium adolescentis		6.62	0.35		1	
43755542	**Low discrimination**					**43755542**
					SPO	
Bifidobacterium adolescentis		84.48	0.69	MNS 87	1	
Actinomyces israelii		14.63	0.62	SOR 11	1	
Clostridium butyricum		0.89	0.20		99	
43755550	**Low discrimination**					**43755550**
					SPO	
Actinomyces israelii		54.87	0.45	MLZ 13	1	
Bifidobacterium adolescentis		39.25	0.36		1	
Clostridium butyricum		4.52	0.05		99	
43755552	**Very good identification**					**43755552**
					SPO	
Bifidobacterium adolescentis		97.88	0.70	MNS 87	1	
Actinomyces israelii		2.07	0.48	SOR 11	1	
Clostridium butyricum		0.04	-.02		99	
43755553	**Good identification**					**43755553**
Bifidobacterium adolescentis		97.93	0.36	MNS 87		
Actinomyces israelii		2.07	0.14	SOR 11		
43755560	**Low discrimination**					**43755560**
					SPO	
Actinomyces israelii		75.84	0.39	NAG 6	1	
Clostridium butyricum		24.16	0.09		99	
43755562	**Low discrimination**					**43755562**
					SPO	
Actinomyces israelii		48.15	0.42	NAG 6	1	
Bifidobacterium adolescentis		43.99	0.35		1	
Clostridium butyricum		3.98	0.02		99	
43755572	**Low discrimination**					**43755572**
Bifidobacterium adolescentis		88.20	0.36	MNS 87		
Actinomyces israelii		11.80	0.28	NAG 1		
43755612	**Acceptable identification**					**43755612**
Clostridium clostridioforme		94.84	0.42	MNS 99		
Actinomyces israelii		2.86	0.25	TRE 88		
Bifidobacterium adolescentis		2.30	0.18			
43755632	**Good identification**					**43755632**
Clostridium clostridioforme		99.10	0.42	MNS 99		
Parabacteroides distasonis		0.91	0.13	TRE 88		
43755700	**Very good identification**					**43755700**
					SPO	
Eubacterium contortum		96.56	0.60	CEL 1	1	
Actinomyces israelii		2.27	0.47		1	
Clostridium butyricum		0.98	0.20		99	
43755702	**Low discrimination**					**43755702**
Eubacterium contortum		52.52	0.47	CEL 1		
Bifidobacterium adolescentis		25.62	0.50			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		10.51	0.50			
43755712	**Low discrimination**					**43755712**
Clostridium clostridioforme		56.54	0.51	MNS 99		
Bifidobacterium adolescentis		40.59	0.52	TRE 88		
Actinomyces israelii		2.04	0.36			
43755720	**Low discrimination**					**43755720**
					GRA	
Eubacterium contortum		68.01	0.47	CEL 1	99	
Bacteroides vulgatus		30.16	0.34		1	
Actinomyces israelii		0.58	0.26		99	
43755722	**Low discrimination**					**43755722**
Bacteroides uniformis		45.09	0.37	IND 98		
Parabacteroides distasonis		25.83	0.52	MNS 99		
Clostridium clostridioforme		11.30	0.40			
43755732	**Low discrimination**					**43755732**
Clostridium clostridioforme		74.06	0.51	MNS 99		
Parabacteroides distasonis		20.00	0.47	TRE 88		
Bacteroides ovatus		4.42	0.20			
43755742	**Low discrimination**					**43755742**
Actinomyces israelii		51.20	0.43	RHA 7		
Bifidobacterium adolescentis		39.66	0.35			
Eubacterium contortum		6.65	0.13			
43755752	**Low discrimination**					**43755752**
Bifidobacterium adolescentis		78.64	0.36	MNS 87		
Actinomyces israelii		12.41	0.29	RHA 1		
Clostridium clostridioforme		8.95	0.17	SOR 11		
43756000	**Good identification**					**43756000**
Actinomyces israelii		100.0	0.36	GAL 99		
43756002	**Good identification**					**43756002**
Actinomyces israelii		100.0	0.39	GAL 99		
43756100	**Good identification**					**43756100**
					GRA	
Actinomyces israelii		98.01	0.46	GAL 99	99	
Fusobacterium mortiferum		1.99	0.15		1	
43756102	**Low discrimination**					**43756102**
Actinomyces israelii		75.72	0.49	GAL 99		
Bifidobacterium adolescentis		24.28	0.35			
43756112	**Low discrimination**					**43756112**
Bifidobacterium adolescentis		72.40	0.36	MNS 87		
Actinomyces israelii		27.61	0.35	GAL 99		
43756140	**Good identification**					**43756140**
Actinomyces israelii		100.0	0.39	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43756142	**Acceptable identification**					**43756142**
Actinomyces israelii		90.75	0.42	GAL 99		
Bifidobacterium adolescentis		9.25	0.19			
43756402	**Acceptable identification**					**43756402**
Actinomyces israelii		90.42	0.38	GAL 99		
Bifidobacterium adolescentis		9.58	0.15			
43756500	**Acceptable identification**					**43756500**
					SPO	
Actinomyces israelii		94.32	0.45	GAL 99	1	
Bifidobacterium adolescentis		3.75	0.15		1	
Clostridium butyricum		1.93	-.05		99	
43756502	**Low discrimination**					**43756502**
Bifidobacterium adolescentis		72.40	0.49	MNS 87		
Actinomyces israelii		27.61	0.48	GAL 99		
43756512	**Good identification**					**43756512**
Bifidobacterium adolescentis		95.07	0.50	MNS 87		
Actinomyces israelii		4.43	0.34	GAL 99		
Bifidobacterium dentium		0.50	-.07			
43756540	**Good identification**					**43756540**
Actinomyces israelii		100.0	0.38	GAL 99		
43756542	**Low discrimination**					**43756542**
Actinomyces israelii		54.54	0.41	GAL 99		
Bifidobacterium adolescentis		45.46	0.33			
43757000	**Very good identification**					**43757000**
					GRA	
Actinomyces israelii		99.95	0.70		99	
Fusobacterium mortiferum		0.05	0.12		1	
43757001	**Acceptable identification**					**43757001**
Actinomyces israelii		94.89	0.36	URE 1		
Actinomyces naeslundii		5.12	0.24			
43757002	**Very good identification**					**43757002**
Actinomyces israelii		98.72	0.73			
Bifidobacterium adolescentis		1.28	0.35			
43757003	**Good identification**					**43757003**
Actinomyces israelii		96.53	0.39	URE 1		
Actinomyces naeslundii		3.47	0.24			
43757010	**Very good identification**					**43757010**
Actinomyces israelii		100.0	0.56	MLZ 13		
43757012	**Good identification**					**43757012**
Actinomyces israelii		90.42	0.59	MLZ 13		
Bifidobacterium adolescentis		9.58	0.36			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43757020	**Good identification**					**43757020**
Actinomyces israelii		100.0	0.50	NAG 6		
43757022	**Good identification**					**43757022**
Actinomyces israelii		100.0	0.53	NAG 6		
43757030	**Good identification**					**43757030**
Actinomyces israelii		100.0	0.36	MLZ 13		
43757032	**Good identification**					**43757032**
Actinomyces israelii		100.0	0.39	MLZ 13		
43757040	**Very good identification**					**43757040**
Actinomyces israelii		100.0	0.63			
43757042	**Very good identification**					**43757042**
Actinomyces israelii		99.59	0.66			
Bifidobacterium adolescentis		0.41	0.19			
43757050	**Good identification**					**43757050**
Actinomyces israelii		100.0	0.49	MLZ 13		
43757052	**Good identification**					**43757052**
Actinomyces israelii		96.74	0.52	MLZ 13		
Bifidobacterium adolescentis		3.26	0.21			
43757060	**Good identification**					**43757060**
Actinomyces israelii		100.0	0.43	NAG 6		
43757062	**Good identification**					**43757062**
Actinomyces israelii		100.0	0.46	NAG 6		
43757100	**Very good identification**					**43757100**
					SPO	
Actinomyces israelii		99.26	0.81		1	
Bifidobacterium adolescentis		0.48	0.35		1	
Clostridium butyricum		0.19	0.13		99	
43757101	**Good identification**					**43757101**
Actinomyces israelii		96.31	0.46	URE 1		
Actinomyces naeslundii		3.69	0.31			
43757102	**Low discrimination**					**43757102**
Actinomyces israelii		75.62	0.84			
Bifidobacterium adolescentis		24.24	0.69			
Bifidobacterium breve		0.08	0.27			
43757103	**Low discrimination**					**43757103**
Actinomyces israelii		74.28	0.49	URE 1		
Bifidobacterium adolescentis		23.82	0.35			
Actinomyces naeslundii		1.90	0.31			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43757110	**Very good identification**					**43757110**
Actinomyces israelii		96.01	0.66	MLZ 13		
Bifidobacterium adolescentis		3.82	0.36			
Bifidobacterium breve		0.10	0.10			
43757112	**Low discrimination**					**43757112**
Bifidobacterium adolescentis		72.20	0.70	MNS 87		
Actinomyces israelii		27.53	0.69	ARA 87		
Bifidobacterium breve		0.20	0.27			
43757113	**Low discrimination**					**43757113**
Bifidobacterium adolescentis		72.40	0.36	MNS 87		
Actinomyces israelii		27.61	0.35	ARA 87		
43757120	**Very good identification**					**43757120**
Actinomyces israelii		98.37	0.60	NAG 6		
Bifidobacterium breve		0.44	0.15			
Actinomyces naeslundii		0.40	0.28			
43757122	**Low discrimination**					**43757122**
Actinomyces israelii		89.40	0.63	NAG 6		
Bifidobacterium adolescentis		4.54	0.35			
Bifidobacterium breve		2.70	0.32			
43757130	**Acceptable identification**					**43757130**
					SPO	
Actinomyces israelii		93.66	0.46	MLZ 13	1	
Clostridium tertium		3.54	0.07	NAG 6	99	
Bifidobacterium breve		2.81	0.15		1	
43757132	**Low discrimination**					**43757132**
Actinomyces israelii		58.18	0.49	MLZ 13		
Bifidobacterium adolescentis		24.15	0.36	NAG 6		
Bifidobacterium breve		11.76	0.32			
43757140	**Very good identification**					**43757140**
Actinomyces israelii		99.76	0.74			
Bifidobacterium adolescentis		0.15	0.19			
Bifidobacterium breve		0.04	0.10			
43757141	**Good identification**					**43757141**
Actinomyces israelii		100.0	0.39	URE 1		
43757142	**Good identification**					**43757142**
Actinomyces israelii		90.52	0.77			
Bifidobacterium adolescentis		9.22	0.53			
Bifidobacterium breve		0.25	0.27			
43757143	**Acceptable identification**					**43757143**
Actinomyces israelii		90.75	0.42	URE 1		
Bifidobacterium adolescentis		9.25	0.19			
43757150	**Very good identification**					**43757150**

Code Taxon		ID score	T index	tests against	Differentiation tests COC	Code
Actinomyces israelii		98.14	0.59	MLZ 13	1	
Bifidobacterium adolescentis		1.24	0.21		1	
Blautia producta		0.35	-.03		99	
43757152	**Low discrimination**					**43757152**
Actinomyces israelii		53.99	0.62	MLZ 13		
Bifidobacterium adolescentis		45.00	0.55			
Bifidobacterium breve		1.01	0.27			
43757160	**Good identification**					**43757160**
Actinomyces israelii		98.86	0.53	NAG 6		
Bifidobacterium breve		1.14	0.15			
43757162	**Acceptable identification**					**43757162**
Actinomyces israelii		91.43	0.56	NAG 6		
Bifidobacterium breve		7.10	0.32			
Bifidobacterium adolescentis		1.47	0.19			
43757170	**Acceptable identification**					**43757170**
Actinomyces israelii		92.84	0.39	MLZ 13		
Bifidobacterium breve		7.16	0.15	NAG 6		
43757172	**Low discrimination**					**43757172**
Actinomyces israelii		60.55	0.42	MLZ 13		
Bifidobacterium breve		31.47	0.32	NAG 6		
Bifidobacterium adolescentis		7.99	0.21			
43757200	**Good identification**					**43757200**
Actinomyces israelii		100.0	0.51	RHA 7		
43757202	**Good identification**					**43757202**
Actinomyces israelii		100.0	0.54	RHA 7		
43757210	**Good identification**					**43757210**
Actinomyces israelii		100.0	0.37	RHA 7		
43757212	**Good identification**					**43757212**
Actinomyces israelii		100.0	0.40	RHA 7		
43757240	**Good identification**					**43757240**
Actinomyces israelii		100.0	0.44	RHA 7		
43757242	**Good identification**					**43757242**
Actinomyces israelii		100.0	0.47	RHA 7		
43757300	**Good identification**					**43757300**
Actinomyces israelii		92.96	0.61	RHA 7		
Eubacterium contortum		7.04	0.27			
43757302	**Very good identification**					**43757302**
Actinomyces israelii		95.06	0.64	RHA 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		4.09	0.35				
Eubacterium contortum		0.85	0.14				
43757310	**Good identification**						**43757310**
Actinomyces israelii		100.0	0.47	RHA 7			
43757312	**Low discrimination**						**43757312**
Actinomyces israelii		73.97	0.50	RHA 7			
Bifidobacterium adolescentis		26.03	0.36	MLZ 13			
43757320	**Low discrimination**						**43757320**
					SPO		
Actinomyces israelii		80.00	0.41	RHA 7	1		
Eubacterium contortum		16.75	0.14	NAG 6	1		
Clostridium ramosum		3.26	0.01		92		
43757322	**Low discrimination**						**43757322**
					SPO	GRA	
Actinomyces israelii		76.58	0.44	RHA 7	1	99	
Clostridium ramosum		8.31	0.11	NAG 6	92	50	
Prevotella oralis		7.29	0.10		1	1	
43757340	**Good identification**						**43757340**
Actinomyces israelii		100.0	0.54	RHA 7			
43757342	**Very good identification**						**43757342**
Actinomyces israelii		98.65	0.57	RHA 7			
Bifidobacterium adolescentis		1.35	0.19				
43757350	**Good identification**						**43757350**
					COC		
Actinomyces israelii		95.47	0.40	RHA 7	1		
Blautia producta		4.53	-.03	MLZ 13	99		
43757352	**Low discrimination**						**43757352**
Actinomyces israelii		89.94	0.43	RHA 7			
Bifidobacterium adolescentis		10.06	0.21	MLZ 13			
43757362	**Good identification**						**43757362**
Actinomyces israelii		100.0	0.37	RHA 7			
43757400	**Very good identification**						**43757400**
					SPO		
Actinomyces israelii		99.50	0.69		1		
Clostridium butyricum		0.34	0.05		99		
Bifidobacterium adolescentis		0.16	0.15		1		
43757402	**Good identification**						**43757402**
					SPO		
Actinomyces israelii		90.35	0.72		1		
Bifidobacterium adolescentis		9.57	0.49		1		
Clostridium butyricum		0.08	-.02		99		
43757403	**Acceptable identification**						**43757403**
Actinomyces israelii		90.42	0.38	URE 1			
Bifidobacterium adolescentis		9.58	0.15				
43757410	**Good identification**						**43757410**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		98.70	0.55	MLZ 13		
Bifidobacterium adolescentis		1.30	0.16			
43757412	**Low discrimination**					**43757412**
Actinomyces israelii		53.44	0.58	MLZ 13		
Bifidobacterium adolescentis		46.32	0.50			
Bifidobacterium dentium		0.25	-.07			
43757420	**Good identification**					**43757420**
Actinomyces israelii		100.0	0.48	NAG 6		
43757422	**Good identification**					**43757422**
Actinomyces israelii		98.35	0.51	NAG 6		
Bifidobacterium adolescentis		1.65	0.15			
43757432	**Low discrimination**					**43757432**
Actinomyces israelii		79.05	0.37	MLZ 13		
Bifidobacterium adolescentis		10.84	0.16	NAG 6		
Bifidobacterium dentium		10.11	-.03			
43757440	**Very good identification**					**43757440**
Actinomyces israelii		100.0	0.62			
43757442	**Very good identification**					**43757442**
Actinomyces israelii		96.74	0.65			
Bifidobacterium adolescentis		3.26	0.33			
43757450	**Good identification**					**43757450**
					COC	
Actinomyces israelii		98.29	0.48	MLZ 13	1	
Blautia producta		1.72	-.03		99	
43757452	**Low discrimination**					**43757452**
Actinomyces israelii		78.40	0.51	MLZ 13		
Bifidobacterium adolescentis		21.60	0.35			
43757460	**Good identification**					**43757460**
Actinomyces israelii		100.0	0.41	NAG 6		
43757462	**Good identification**					**43757462**
Actinomyces israelii		100.0	0.44	NAG 6		
43757500	**Good identification**					**43757500**
					SPO	
Actinomyces israelii		94.32	0.79		1	
Bifidobacterium adolescentis		3.75	0.49		1	
Clostridium butyricum		1.93	0.29		99	
43757501	**Acceptable identification**					**43757501**
					SPO	
Actinomyces israelii		94.32	0.45	URE 1	1	
Bifidobacterium adolescentis		3.75	0.15		1	
Clostridium butyricum		1.93	-.05		99	
43757502	**Low discrimination**					**43757502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Bifidobacterium adolescentis		72.29	0.83	MNS 87	1	
Actinomyces israelii		27.56	0.82		1	
Clostridium butyricum		0.15	0.22		99	
43757503	**Low discrimination**					**43757503**
Bifidobacterium adolescentis		72.40	0.49	MNS 87		
Actinomyces israelii		27.61	0.48	URE 1		
43757510	**Low discrimination**					**43757510**
Actinomyces israelii		74.03	0.65	MLZ 13		
Bifidobacterium adolescentis		24.06	0.50			
Bifidobacterium dentium		1.25	0.10			
43757512	**Very good identification**					**43757512**
Bifidobacterium adolescentis		95.06	0.84	MNS 87		
Actinomyces israelii		4.43	0.68			
Bifidobacterium dentium		0.50	0.27			
43757513	**Good identification**					**43757513**
Bifidobacterium adolescentis		95.07	0.50	MNS 87		
Actinomyces israelii		4.43	0.34	URE 1		
Bifidobacterium dentium		0.50	-.07			
43757520	**Very good identification**					**43757520**
					SPO	
Actinomyces israelii		96.69	0.59	NAG 6	1	
Clostridium butyricum		2.70	0.11		99	
Bifidobacterium adolescentis		0.61	0.15		1	
43757522	**Low discrimination**					**43757522**
					SPO	
Actinomyces israelii		69.81	0.62	NAG 6	1	
Bifidobacterium adolescentis		28.97	0.49		1	
Clostridium butyricum		0.51	0.04		99	
43757530	**Low discrimination**					**43757530**
Actinomyces israelii		65.75	0.44	MLZ 13		
Bifidobacterium dentium		30.87	0.15	NAG 6		
Bifidobacterium adolescentis		3.38	0.16			
43757532	**Low discrimination**					**43757532**
Bifidobacterium adolescentis		44.91	0.50	MNS 87		
Bifidobacterium dentium		41.86	0.32	NAG 1		
Actinomyces israelii		13.23	0.47			
43757540	**Very good identification**					**43757540**
					COC	
Actinomyces israelii		96.27	0.72		1	
Blautia producta		2.07	0.24		99	
Bifidobacterium adolescentis		1.22	0.33		1	
43757541	**Good identification**					**43757541**
Actinomyces israelii		100.0	0.38	URE 1		
43757542	**Low discrimination**					**43757542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces israelii		54.46	0.75		1		
Bifidobacterium adolescentis		45.39	0.67		1		
Blautia producta		0.09	0.07		99		
43757543	**Low discrimination**						**43757543**
Actinomyces israelii		54.54	0.41	URE 1			
Bifidobacterium adolescentis		45.46	0.33				
43757550	**Low discrimination**						**43757550**
					COC		
Actinomyces israelii		65.13	0.58	MLZ 13	1		
Blautia producta		28.14	0.32		99		
Bifidobacterium adolescentis		6.73	0.35		1		
43757552	**Low discrimination**						**43757552**
					COC		
Bifidobacterium adolescentis		86.82	0.69	MNS 87	1		
Actinomyces israelii		12.74	0.61	SOR 11	1		
Blautia producta		0.41	0.15		99		
43757560	**Good identification**						**43757560**
					COC	SPO	
Actinomyces israelii		95.81	0.52	NAG 6	1	1	
Blautia producta		3.59	0.07		99	1	
Clostridium butyricum		0.60	-.07		1	99	
43757562	**Low discrimination**						**43757562**
Actinomyces israelii		88.35	0.55	NAG 6			
Bifidobacterium adolescentis		11.65	0.33				
43757570	**Low discrimination**						**43757570**
					COC		
Actinomyces israelii		57.08	0.37	MLZ 13	1		
Blautia producta		42.92	0.15	NAG 6	99		
43757600	**Good identification**						**43757600**
Actinomyces israelii		100.0	0.50	RHA 7			
43757602	**Good identification**						**43757602**
Actinomyces israelii		98.60	0.53	RHA 7			
Bifidobacterium adolescentis		1.40	0.15				
43757610	**Good identification**						**43757610**
Actinomyces israelii		100.0	0.35	RHA 7			
43757612	**Low discrimination**						**43757612**
Actinomyces israelii		85.07	0.38	RHA 7			
Bifidobacterium adolescentis		9.89	0.16	MLZ 13			
Clostridium clostridioforme		5.03	0.08				
43757640	**Good identification**						**43757640**
Actinomyces israelii		100.0	0.42	RHA 7			
43757642	**Good identification**						**43757642**
Actinomyces israelii		100.0	0.45	RHA 7			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43757700	**Good identification**					**43757700**
					SPO	
Actinomyces israelii		91.79	0.60	RHA 7	1	
Eubacterium contortum		6.95	0.26		1	
Clostridium butyricum		0.77	0.03		99	
43757702	**Low discrimination**					**43757702**
Actinomyces israelii		73.28	0.63	RHA 7		
Bifidobacterium adolescentis		25.79	0.49			
Eubacterium contortum		0.65	0.13			
43757710	**Acceptable identification**					**43757710**
					COC	
Actinomyces israelii		93.80	0.46	RHA 7	1	
Bifidobacterium adolescentis		4.09	0.16	MLZ 13	1	
Blautia producta		2.11	-.03		99	
43757712	**Low discrimination**					**43757712**
Bifidobacterium adolescentis		73.00	0.50	MNS 87		
Actinomyces israelii		25.36	0.49	RHA 1		
Clostridium clostridioforme		1.26	0.17			
43757720	**Low discrimination**					**43757720**
					GRA	
Actinomyces israelii		76.79	0.39	RHA 7	99	
Eubacterium contortum		16.08	0.13	NAG 6	99	
Bacteroides vulgatus		7.13	-.01		1	
43757722	**Low discrimination**					**43757722**
					GRA	
Actinomyces israelii		62.19	0.42	RHA 7	99	
Bacteroides ovatus		21.30	0.15	NAG 6	1	
Bacteroides uniformis		6.31	0.03		1	
43757740	**Low discrimination**					**43757740**
					COC	
Actinomyces israelii		77.77	0.53	RHA 7	1	
Blautia producta		22.23	0.24		99	
43757742	**Low discrimination**					**43757742**
					COC	
Actinomyces israelii		88.26	0.56	RHA 7	1	
Bifidobacterium adolescentis		9.87	0.33		1	
Blautia producta		1.87	0.07		99	
43757752	**Low discrimination**					**43757752**
					COC	
Actinomyces israelii		42.74	0.42	RHA 7	1	
Bifidobacterium adolescentis		39.10	0.35	MLZ 13	1	
Blautia producta		18.17	0.15		99	
43757762	**Good identification**					**43757762**
Actinomyces israelii		100.0	0.35	RHA 7		
43761001	**Good identification**					**43761001**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43761003	**Good identification**					**43761003**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.36	ESL 12		
43761021	**Good identification**					**43761021**
	Actinomyces naeslundii	100.0	0.36	ESL 12		
43761023	**Good identification**					**43761023**
	Actinomyces naeslundii	100.0	0.36	ESL 12		
43761100	**Low discrimination**					**43761100**
					GRA	
	Actinomyces naeslundii	42.76	0.41	ESL 12	99	
	Fusobacterium mortiferum	38.72	0.28	MNS 88	1	
	Actinomyces israelii	18.53	0.25	FRU 95	99	
43761101	**Good identification**					**43761101**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
43761102	**Low discrimination**					**43761102**
	Actinomyces naeslundii	26.58	0.41	ESL 12		
	Bifidobacterium breve	21.42	0.24	MNS 88		
	Lactobacillus acidophilus	17.65	0.13	FRU 95		
43761103	**Good identification**					**43761103**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
43761120	**Low discrimination**					**43761120**
					SPO	
	Actinomyces naeslundii	52.87	0.41	ESL 12	1	
	Clostridium paraputrificum	23.62	0.03	MNS 88	99	
	Clostridium ramosum	16.02	0.06	FRU 95	92	
43761121	**Good identification**					**43761121**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
43761123	**Good identification**					**43761123**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
43763000	**Low discrimination**					**43763000**
	Actinomyces naeslundii	52.63	0.38	ESL 12		
	Actinomyces israelii	47.37	0.28	MNS 88		
43763001	**Good identification**					**43763001**
	Actinomyces naeslundii	100.0	0.41	ESL 12		
43763003	**Good identification**					**43763003**
	Actinomyces naeslundii	100.0	0.41	ESL 12		
43763020	**Good identification**					**43763020**
	Actinomyces naeslundii	100.0	0.38	ESL 12		
43763021	**Good identification**					**43763021**
	Actinomyces naeslundii	100.0	0.41	ESL 12		
43763022	**Good identification**					**43763022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	ESL 12		
43763023	**Good identification**					**43763023**
Actinomyces naeslundii		100.0	0.41	ESL 12		
43763100	**Low discrimination**					**43763100**
Actinomyces israelii		55.85	0.38	FRU 99		
Actinomyces naeslundii		44.15	0.46	XYL 92		
43763101	**Good identification**					**43763101**
Actinomyces naeslundii		100.0	0.49	ESL 12		
43763102	**Low discrimination**					**43763102**
Actinomyces israelii		54.25	0.41	FRU 99		
Actinomyces naeslundii		28.59	0.46	XYL 92		
Bifidobacterium breve		11.87	0.24			
43763103	**Good identification**					**43763103**
Actinomyces naeslundii		100.0	0.49	ESL 12		
43763120	**Low discrimination**					**43763120**
					SPO	
Actinomyces naeslundii		49.06	0.46	ESL 12	1	
Clostridium ramosum		43.40	0.19	MNS 88	92	
Actinomyces israelii		3.96	0.18	FRU 95	1	
43763121	**Good identification**					**43763121**
Actinomyces naeslundii		100.0	0.49	ESL 12		
43763123	**Good identification**					**43763123**
Actinomyces naeslundii		100.0	0.49	ESL 12		
43763141	**Good identification**					**43763141**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43763143	**Good identification**					**43763143**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43763161	**Good identification**					**43763161**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43763163	**Good identification**					**43763163**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43763500	**Good identification**					**43763500**
Actinomyces israelii		100.0	0.37	FRU 99		
43763502	**Low discrimination**					**43763502**
Actinomyces israelii		55.60	0.40	FRU 99		
Bifidobacterium adolescentis		44.40	0.32	XYL 92		
43765002	**Good identification**					**43765002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	FRU 99		
43765100	**Low discrimination**					**43765100**
					SPO	
Actinomyces israelii		71.63	0.43	FRU 99	1	
Clostridium butyricum		28.37	0.15		99	
43765102	**Low discrimination**					**43765102**
					SPO	
Actinomyces israelii		66.84	0.46	FRU 99	1	
Bifidobacterium adolescentis		26.30	0.33		1	
Clostridium butyricum		6.86	0.08		99	
43765140	**Acceptable identification**					**43765140**
					SPO	
Actinomyces israelii		91.86	0.36	FRU 99	1	
Clostridium butyricum		8.14	-.03		99	
43765142	**Low discrimination**					**43765142**
Actinomyces israelii		88.88	0.39	FRU 99		
Bifidobacterium adolescentis		11.12	0.18			
43765502	**Low discrimination**					**43765502**
					SPO	
Bifidobacterium adolescentis		59.98	0.47	MNS 87	1	
Clostridium butyricum		21.05	0.25	FRU 99	99	
Actinomyces israelii		18.64	0.45		1	
43765512	**Acceptable identification**					**43765512**
Bifidobacterium adolescentis		92.53	0.49	MNS 87		
Actinomyces israelii		3.52	0.31	FRU 99		
Clostridium clostridioforme		1.95	0.17			
43767000	**Good identification**					**43767000**
Actinomyces israelii		100.0	0.46	FRU 99		
43767002	**Good identification**					**43767002**
Actinomyces israelii		100.0	0.49	FRU 99		
43767040	**Good identification**					**43767040**
Actinomyces israelii		100.0	0.39	FRU 99		
43767042	**Good identification**					**43767042**
Actinomyces israelii		100.0	0.42	FRU 99		
43767100	**Very good identification**					**43767100**
					SPO	
Actinomyces israelii		99.23	0.56	FRU 99	1	
Clostridium butyricum		0.77	-.01		99	
43767102	**Good identification**					**43767102**
Actinomyces israelii		94.62	0.59	FRU 99		
Bifidobacterium adolescentis		5.38	0.32			
43767110	**Good identification**					**43767110**
Actinomyces israelii		100.0	0.42	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43767112	**Low discrimination**					**43767112**
Actinomyces israelii		67.14	0.45	FRU 99		
Bifidobacterium adolescentis		31.20	0.33	MLZ 13		
Bifidobacterium dentium		1.66	-.07			
43767120	**Good identification**					**43767120**
Actinomyces israelii		100.0	0.36	FRU 99		
43767122	**Good identification**					**43767122**
Actinomyces israelii		100.0	0.39	FRU 99		
43767140	**Good identification**					**43767140**
Actinomyces israelii		100.0	0.49	FRU 99		
43767142	**Good identification**					**43767142**
Actinomyces israelii		98.23	0.52	FRU 99		
Bifidobacterium adolescentis		1.77	0.16			
43767150	**Acceptable identification**					**43767150**
					COC	
Actinomyces israelii		91.41	0.35	FRU 99	1	
Blautia producta		8.59	-.03	MLZ 13	99	
43767152	**Low discrimination**					**43767152**
Actinomyces israelii		87.13	0.38	FRU 99		
Bifidobacterium adolescentis		12.87	0.18	MLZ 13		
43767300	**Good identification**					**43767300**
Actinomyces israelii		100.0	0.37	FRU 99		
43767302	**Good identification**					**43767302**
Actinomyces israelii		100.0	0.40	FRU 99		
43767400	**Good identification**					**43767400**
Actinomyces israelii		100.0	0.44	FRU 99		
43767402	**Good identification**					**43767402**
Actinomyces israelii		98.16	0.47	FRU 99		
Bifidobacterium adolescentis		1.84	0.12			
43767440	**Good identification**					**43767440**
Actinomyces israelii		100.0	0.37	FRU 99		
43767442	**Good identification**					**43767442**
Actinomyces israelii		100.0	0.40	FRU 99		
43767500	**Acceptable identification**					**43767500**
					SPO	
Actinomyces israelii		91.54	0.55	FRU 99	1	
Clostridium butyricum		7.82	0.15		99	
Bifidobacterium adolescentis		0.65	0.12		1	
43767502	**Low discrimination**					**43767502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		67.05	0.58	FRU 99	1		
Bifidobacterium adolescentis		31.16	0.46		1		
Clostridium butyricum		1.49	0.08		99		
43767510	**Low discrimination**						**43767510**
Actinomyces israelii		63.46	0.41	FRU 99			
Bifidobacterium dentium		28.15	0.10	MLZ 13			
Bifidobacterium adolescentis		3.66	0.13				
43767512	**Low discrimination**						**43767512**
Bifidobacterium adolescentis		48.79	0.47	MNS 87			
Bifidobacterium dentium		38.37	0.27	FRU 99			
Actinomyces israelii		12.84	0.44				
43767522	**Low discrimination**						**43767522**
Actinomyces israelii		83.39	0.37	FRU 99			
Bifidobacterium dentium		10.48	-.03	NAG 6			
Bifidobacterium adolescentis		6.13	0.12				
43767540	**Low discrimination**						**43767540**
					COC	SPO	
Actinomyces israelii		63.08	0.48	FRU 99	1	1	
Blautia producta		35.72	0.24		99	1	
Clostridium butyricum		1.21	-.03		1	99	
43767542	**Low discrimination**						**43767542**
					COC		
Actinomyces israelii		84.06	0.51	FRU 99	1		
Bifidobacterium adolescentis		12.42	0.30		1		
Blautia producta		3.53	0.07		99		
43767700	**Good identification**						**43767700**
Actinomyces israelii		100.0	0.35	FRU 99			
43767702	**Acceptable identification**						**43767702**
Actinomyces israelii		94.13	0.39	FRU 99			
Bifidobacterium adolescentis		5.87	0.12	RHA 7			
43770000	**Good identification**						**43770000**
					GRA		
Fusobacterium mortiferum		98.51	0.53	MLT 11	1		
Actinomyces naeslundii		1.07	0.31		99		
Actinomyces israelii		0.42	0.15		99		
43770002	**Low discrimination**						**43770002**
					GRA		
Fusobacterium mortiferum		86.59	0.37	MLT 11	1		
Actinomyces naeslundii		8.46	0.31	BGL 10	99		
Actinomyces israelii		4.95	0.18		99		
43770100	**Very good identification**						**43770100**
					GRA		
Fusobacterium mortiferum		97.99	0.59	MLT 11	1		
Actinomyces naeslundii		1.30	0.39		99		
Actinomyces israelii		0.71	0.25		99		
43770101	**Low discrimination**						**43770101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		66.30	0.42	ESL 12	99		
Fusobacterium mortiferum		33.70	0.25	MNS 88	1		
43770102	**Low discrimination**						**43770102**
					GRA		
Fusobacterium mortiferum		65.79	0.43	MLT 11	1		
Bifidobacterium breve		7.99	0.24	BGL 10	99		
Actinomyces naeslundii		7.84	0.39		99		
43770103	**Good identification**						**43770103**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43770120	**Low discrimination**						**43770120**
					GRA	SPO	
Actinomyces naeslundii		34.62	0.39	ESL 12	99	1	
Fusobacterium mortiferum		26.40	0.25	MNS 88	1	1	
Clostridium paraputrificum		19.54	0.03	GAL 96	99	99	
43770121	**Good identification**						**43770121**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43770123	**Good identification**						**43770123**
Actinomyces naeslundii		100.0	0.42	ESL 12			
43770712	**Good identification**						**43770712**
Clostridium clostridioforme		100.0	0.41	MNS 99			
43770732	**Good identification**						**43770732**
Clostridium clostridioforme		100.0	0.41	MNS 99			
43771000	**Low discrimination**						**43771000**
					GRA		
Fusobacterium mortiferum		62.39	0.56	MLT 11	1		
Actinomyces israelii		17.42	0.49		99		
Actinomyces naeslundii		10.84	0.55		99		
43771001	**Very good identification**						**43771001**
					GRA		
Actinomyces naeslundii		95.28	0.58	ESL 12	99		
Fusobacterium mortiferum		3.69	0.22	MNS 88	1		
Actinomyces israelii		1.03	0.15		99		
43771002	**Low discrimination**						**43771002**
Actinomyces israelii		42.14	0.52	XYL 92			
Actinomyces naeslundii		17.49	0.55				
Lactobacillus acidophilus		14.97	0.29				
43771003	**Very good identification**						**43771003**
Actinomyces naeslundii		98.40	0.58	ESL 12			
Actinomyces israelii		1.60	0.18	MNS 88			
43771011	**Good identification**						**43771011**
Actinomyces naeslundii		100.0	0.36	ESL 12			
43771013	**Low discrimination**						**43771013**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		83.82	0.36	ESL 12	99	
Leptotrichia buccalis		16.18	-.11	MNS 88	1	
43771020	**Low discrimination**					**43771020**
Actinomyces naeslundii		64.64	0.55	ESL 12		
Lactobacillus jensenii		13.33	0.21	MNS 88		
Actinomyces israelii		6.63	0.29			
43771021	**Very good identification**					**43771021**
Actinomyces naeslundii		100.0	0.58	ESL 12		
43771022	**Low discrimination**					**43771022**
					GRA	
Actinomyces naeslundii		31.23	0.55	ESL 12	99	
Lactobacillus acidophilus		26.75	0.29	MNS 88	99	
Prevotella oralis		18.15	0.25		1	
43771023	**Very good identification**					**43771023**
Actinomyces naeslundii		100.0	0.58	ESL 12		
43771031	**Good identification**					**43771031**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43771033	**Good identification**					**43771033**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43771040	**Low discrimination**					**43771040**
					GRA	
Actinomyces israelii		72.70	0.42	XYL 92	99	
Actinomyces naeslundii		20.54	0.42		99	
Fusobacterium mortiferum		6.77	0.22		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43771041	**Good identification**					**43771041**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43771042	**Low discrimination**					**43771042**
Actinomyces israelii		69.93	0.45	XYL 92		
Actinomyces naeslundii		13.17	0.42			
Bifidobacterium breve		9.01	0.24			
43771043	**Good identification**					**43771043**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43771060	**Low discrimination**					**43771060**
Actinomyces naeslundii		74.28	0.42	ESL 12		
Actinomyces israelii		16.79	0.22	MNS 88		
Bifidobacterium breve		8.93	0.12			
43771061	**Good identification**					**43771061**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43771063	**Good identification**					**43771063**
Actinomyces naeslundii		100.0	0.45	ESL 12		
43771100	**Low discrimination**					**43771100**
					GRA	
Fusobacterium mortiferum		52.20	0.62	MLT 11	1	
Actinomyces israelii		24.98	0.59		99	
Actinomyces naeslundii		11.07	0.62		99	
43771101	**Very good identification**					**43771101**
					GRA	
Actinomyces naeslundii		95.51	0.65	ESL 12	99	
Fusobacterium mortiferum		3.03	0.28	MNS 88	1	
Actinomyces israelii		1.45	0.25		99	
43771102	**Low discrimination**					**43771102**
Bifidobacterium breve		26.56	0.59	MNS 99		
Lactobacillus acidophilus		21.88	0.47			
Actinomyces israelii		21.42	0.62			
43771103	**Good identification**					**43771103**
Actinomyces naeslundii		91.17	0.65	ESL 12		
Bifidobacterium breve		2.58	0.24	MNS 88		
Lactobacillus acidophilus		2.12	0.13			
43771111	**Acceptable identification**					**43771111**
					GRA	
Actinomyces naeslundii		93.65	0.44	ESL 12	99	
Leptotrichia buccalis		6.35	-.11	MNS 88	1	
43771112	**Low discrimination**					**43771112**
					GRA	
Leptotrichia buccalis		87.32	0.58	MNS 99	1	
Bifidobacterium breve		6.99	0.59		99	
Bifidobacterium adolescentis		4.66	0.55		99	
43771120	**Low discrimination**					**43771120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium paraputrificum		37.86	0.37	MNS 99	99	
Clostridium ramosum		25.68	0.41	RAF 7	92	
Actinomyces naeslundii		16.26	0.62		1	
43771121	**Very good identification**					**43771121**
					SPO	
Actinomyces naeslundii		96.97	0.65	ESL 12	1	
Clostridium paraputrificum		1.52	0.03	MNS 88	99	
Clostridium ramosum		1.03	0.06		92	
43771122	**Low discrimination**					**43771122**
					SPO	
Bifidobacterium breve		32.18	0.63	MNS 99	1	
Clostridium ramosum		27.23	0.51		92	
Lactobacillus acidophilus		14.91	0.47		1	
43771123	**Low discrimination**					**43771123**
					SPO	
Actinomyces naeslundii		87.38	0.65	ESL 12	1	
Bifidobacterium breve		4.39	0.29	MNS 88	1	
Clostridium ramosum		3.72	0.17		92	
43771130	**Low discrimination**					**43771130**
					GRA	
Bifidobacterium breve		81.06	0.46	MNS 99	99	
Leptotrichia buccalis		6.47	0.07	BGL 91	1	
Actinomyces naeslundii		5.78	0.41		99	
43771131	**Acceptable identification**					**43771131**
Actinomyces naeslundii		91.37	0.44	ESL 12		
Bifidobacterium breve		8.63	0.12	MNS 88		
43771132	**Low discrimination**					**43771132**
					GRA	
Bifidobacterium breve		54.23	0.63	MNS 99	99	
Leptotrichia buccalis		42.37	0.41		1	
Parabacteroides distasonis		1.43	0.37		1	
43771133	**Low discrimination**					**43771133**
					GRA	
Actinomyces naeslundii		37.02	0.44	ESL 12	99	
Bifidobacterium breve		35.36	0.29	MNS 88	99	
Leptotrichia buccalis		27.62	0.07	MLZ 5	1	
43771140	**Low discrimination**					**43771140**
Actinomyces israelii		56.83	0.52	XYL 92		
Bifidobacterium breve		26.88	0.41			
Actinomyces naeslundii		11.42	0.50			
43771141	**Good identification**					**43771141**
Actinomyces naeslundii		96.76	0.53	ESL 12		
Actinomyces israelii		3.24	0.18	MNS 88		
43771142	**Low discrimination**					**43771142**
Bifidobacterium breve		65.36	0.59	MNS 99		
Actinomyces israelii		20.50	0.55			
Lactobacillus acidophilus		6.66	0.32			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43771143	**Low discrimination**						**43771143**
Actinomyces naeslundii		82.61	0.53	ESL 12			
Bifidobacterium breve		13.24	0.24	MNS 88			
Actinomyces israelii		4.15	0.21				
43771150	**Low discrimination**						**43771150**
Bifidobacterium breve		74.01	0.41	MNS 99			
Actinomyces israelii		23.38	0.38	BGL 91			
Actinomyces naeslundii		1.66	0.28				
43771152	**Low discrimination**						**43771152**
					GRA		
Bifidobacterium breve		79.51	0.59	MNS 99	99		
Leptotrichia buccalis		10.04	0.24		1		
Bifidobacterium adolescentis		6.55	0.39		99		
43771160	**Low discrimination**						**43771160**
					SPO		
Bifidobacterium breve		63.54	0.46	MNS 99	1		
Actinomyces naeslundii		15.19	0.50	BGL 91	1		
Clostridium paraputrificum		15.08	0.18		99		
43771161	**Good identification**						**43771161**
Actinomyces naeslundii		97.26	0.53	ESL 12			
Bifidobacterium breve		2.74	0.12	MNS 88			
43771162	**Low discrimination**						**43771162**
Bifidobacterium breve		89.68	0.63	MNS 99			
Lactobacillus acidophilus		5.14	0.32				
Actinomyces naeslundii		2.12	0.50				
43771163	**Low discrimination**						**43771163**
Actinomyces naeslundii		77.83	0.53	ESL 12			
Bifidobacterium breve		22.17	0.29	MNS 88			
43771170	**Good identification**						**43771170**
Bifidobacterium breve		97.66	0.46	MNS 99			
Actinomyces naeslundii		1.23	0.28	BGL 91			
Actinomyces israelii		1.11	0.18				
43771172	**Very good identification**						**43771172**
					GRA		
Bifidobacterium breve		98.93	0.63	MNS 99	99		
Leptotrichia buccalis		0.78	0.07		1		
Actinomyces israelii		0.17	0.21		99		
43771300	**Low discrimination**						**43771300**
					SPO	GRA	
Actinomyces israelii		65.70	0.40	XYL 92	1	99	
Fusobacterium mortiferum		18.42	0.28	RHA 7	1	1	
Clostridium ramosum		8.17	0.09		92	50	
43771320	**Acceptable identification**						**43771320**
					SPO	GRA	
Clostridium ramosum		93.94	0.43	MNS 99	92	50	
Prevotella oralis		3.24	0.17		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		1.06	0.03		99	99	
43771322	**Low discrimination**						**43771322**
					SPO		
Clostridium ramosum		74.50	0.53	MNS 99	92		
Prevotella oralis		12.20	0.39		1		
Parabacteroides distasonis		9.75	0.50		1		
43771332	**Low discrimination**						**43771332**
					SPO		
Parabacteroides distasonis		62.42	0.46	MNS 98	1		
Clostridium clostridioforme		12.96	0.28	XYL 90	66		
Clostridium ramosum		8.95	0.19		92		
43771342	**Low discrimination**						**43771342**
Actinomyces israelii		57.06	0.36	XYL 92			
Bifidobacterium breve		24.41	0.24	RHA 7			
Lactobacillus acidophilus		18.53	0.12				
43771400	**Low discrimination**						**43771400**
					GRA	SPO	
Actinomyces israelii		76.78	0.48	XYL 92	99	1	
Clostridium butyricum		19.83	0.16		99	99	
Fusobacterium mortiferum		3.40	0.22		1	1	
43771402	**Low discrimination**						**43771402**
					SPO		
Actinomyces israelii		76.04	0.51	XYL 92	1		
Bifidobacterium adolescentis		16.99	0.33		1		
Clostridium butyricum		5.09	0.09		99		
43771440	**Acceptable identification**						**43771440**
					SPO		
Actinomyces israelii		94.54	0.41	XYL 92	1		
Clostridium butyricum		5.46	-.02		99		
43771442	**Acceptable identification**						**43771442**
Actinomyces israelii		93.37	0.44	XYL 92			
Bifidobacterium adolescentis		6.63	0.18				
43771500	**Low discrimination**						**43771500**
					SPO		
Clostridium butyricum		58.15	0.40	MNS 99	99		
Actinomyces israelii		37.53	0.58	XYL 97	1		
Bifidobacterium adolescentis		3.15	0.33		1		
43771502	**Low discrimination**						**43771502**
					SPO		
Bifidobacterium adolescentis		79.14	0.67	MNS 87	1		
Actinomyces israelii		14.31	0.61	XYL 87	1		
Clostridium butyricum		5.75	0.33		99		
43771510	**Low discrimination**						**43771510**
					SPO		
Actinomyces israelii		44.43	0.44	XYL 92	1		
Bifidobacterium adolescentis		30.46	0.35	MLZ 13	1		
Clostridium butyricum		24.25	0.18		99		
43771512	**Good identification**						**43771512**
					GRA		
Bifidobacterium adolescentis		93.44	0.69	MNS 87	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Leptotrichia buccalis		2.65	0.24	XYL 87	1	
Actinomyces israelii		2.07	0.47		99	
43771522	**Low discrimination**					**43771522**
					GRA	
Parabacteroides distasonis		23.91	0.38	MNS 98	1	
Actinomyces israelii		18.89	0.40	XYL 90	99	
Bifidobacterium adolescentis		16.53	0.33		99	
43771540	**Low discrimination**					**43771540**
					SPO	
Actinomyces israelii		72.83	0.51	XYL 92	1	
Clostridium butyricum		25.23	0.22		99	
Bifidobacterium adolescentis		1.94	0.18		1	
43771542	**Low discrimination**					**43771542**
					SPO	
Bifidobacterium adolescentis		60.89	0.52	MNS 87	1	
Actinomyces israelii		34.64	0.54	XYL 87	1	
Clostridium butyricum		3.11	0.15	SOR 11	99	
43771550	**Low discrimination**					**43771550**
					SPO	
Actinomyces israelii		70.65	0.37	XYL 92	1	
Bifidobacterium adolescentis		15.39	0.19	MLZ 13	1	
Clostridium butyricum		8.62	0.00		99	
43771552	**Acceptable identification**					**43771552**
Bifidobacterium adolescentis		91.54	0.53	MNS 87		
Actinomyces israelii		6.37	0.40	XYL 87		
Bifidobacterium breve		1.68	0.24	SOR 11		
43771612	**Good identification**					**43771612**
Clostridium clostridioforme		98.66	0.40	MNS 99		
Actinomyces israelii		1.34	0.17	XYL 91		
43771632	**Good identification**					**43771632**
Clostridium clostridioforme		100.0	0.40	MNS 99		
43771700	**Low discrimination**					**43771700**
					SPO	
Actinomyces israelii		61.10	0.39	XYL 92	1	
Clostridium butyricum		38.90	0.14	RHA 7	99	
43771702	**Low discrimination**					**43771702**
Clostridium clostridioforme		51.01	0.37	MNS 99		
Actinomyces israelii		25.26	0.42	XYL 91		
Bifidobacterium adolescentis		18.75	0.33			
43771712	**Low discrimination**					**43771712**
Clostridium clostridioforme		89.47	0.49	MNS 99		
Bifidobacterium adolescentis		8.82	0.35	XYL 91		
Actinomyces israelii		1.45	0.27			
43771722	**Low discrimination**					**43771722**
					SPO	
Parabacteroides distasonis		54.40	0.47	MNS 98	1	
Clostridium clostridioforme		34.12	0.37	XYL 90	66	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		6.85	0.19		92	
43771732	**Low discrimination**					**43771732**
Clostridium clostridioforme		83.55	0.49	MNS 99		
Parabacteroides distasonis		15.75	0.42	XYL 91		
Bacteroides ovatus		0.70	0.03			
43772001	**Good identification**					**43772001**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43772003	**Good identification**					**43772003**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43772020	**Good identification**					**43772020**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43772021	**Good identification**					**43772021**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43772022	**Good identification**					**43772022**
Actinomyces naeslundii		100.0	0.36	ESL 12		
43772023	**Good identification**					**43772023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
43772100	**Low discrimination**					**43772100**
					GRA	
Actinomyces israelii		53.43	0.38	GAL 99	99	
Actinomyces naeslundii		33.44	0.44	XYL 92	99	
Fusobacterium mortiferum		13.13	0.25		1	
43772101	**Good identification**					**43772101**
Actinomyces naeslundii		100.0	0.47	ESL 12		
43772102	**Low discrimination**					**43772102**
Actinomyces israelii		57.68	0.41	GAL 99		
Actinomyces naeslundii		24.07	0.44	XYL 92		
Bifidobacterium breve		12.62	0.24			
43772103	**Good identification**					**43772103**
Actinomyces naeslundii		100.0	0.47	ESL 12		
43772121	**Good identification**					**43772121**
Actinomyces naeslundii		100.0	0.47	ESL 12		
43772123	**Good identification**					**43772123**
Actinomyces naeslundii		100.0	0.47	ESL 12		
43772500	**Good identification**					**43772500**
Actinomyces israelii		100.0	0.37	GAL 99		
43772502	**Low discrimination**					**43772502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		55.60	0.40	GAL 99		
Bifidobacterium adolescentis		44.40	0.32	XYL 92		
43773000	**Low discrimination**					**43773000**
					GRA	
Actinomyces israelii		81.99	0.62	XYL 92	99	
Actinomyces naeslundii		17.48	0.60		99	
Fusobacterium mortiferum		0.52	0.22		1	
43773001	**Very good identification**					**43773001**
Actinomyces naeslundii		96.94	0.63	ESL 12		
Actinomyces israelii		3.06	0.28	MNS 88		
43773002	**Low discrimination**					**43773002**
Actinomyces israelii		86.34	0.65	XYL 92		
Actinomyces naeslundii		12.27	0.60			
Bifidobacterium breve		0.76	0.24			
43773003	**Very good identification**					**43773003**
Actinomyces naeslundii		95.48	0.63	ESL 12		
Actinomyces israelii		4.52	0.31	MNS 88		
43773010	**Acceptable identification**					**43773010**
Actinomyces israelii		93.01	0.48	XYL 92		
Actinomyces naeslundii		6.99	0.38	MLZ 13		
43773011	**Acceptable identification**					**43773011**
Actinomyces naeslundii		91.77	0.41	ESL 12		
Actinomyces israelii		8.23	0.14	MNS 88		
43773012	**Low discrimination**					**43773012**
Actinomyces israelii		87.03	0.51	XYL 92		
Bifidobacterium breve		5.15	0.24	MLZ 13		
Actinomyces naeslundii		4.36	0.38			
43773013	**Low discrimination**					**43773013**
Actinomyces naeslundii		88.15	0.41	ESL 12		
Actinomyces israelii		11.85	0.17	MNS 88		
43773020	**Low discrimination**					**43773020**
					SPO	
Actinomyces naeslundii		69.31	0.60	ESL 12	1	
Actinomyces israelii		20.75	0.41	MNS 88	1	
Clostridium ramosum		9.19	0.19		92	
43773021	**Very good identification**					**43773021**
Actinomyces naeslundii		100.0	0.63	ESL 12		
43773022	**Low discrimination**					**43773022**
					SPO	
Actinomyces naeslundii		47.33	0.60	ESL 12	1	
Clostridium ramosum		25.09	0.30	MNS 88	92	
Actinomyces israelii		21.25	0.44		1	
43773023	**Very good identification**					**43773023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.63	ESL 12		
43773030	**Low discrimination**					**43773030**
Actinomyces naeslundii		48.60	0.38	ESL 12		
Actinomyces israelii		41.30	0.27	MNS 88		
Bifidobacterium breve		10.10	0.12	MLZ 5		
43773031	**Good identification**					**43773031**
Actinomyces naeslundii		100.0	0.41	ESL 12		
43773033	**Good identification**					**43773033**
Actinomyces naeslundii		100.0	0.41	ESL 12		
43773040	**Acceptable identification**					**43773040**
Actinomyces israelii		91.18	0.55	XYL 92		
Actinomyces naeslundii		8.82	0.47			
43773041	**Acceptable identification**					**43773041**
Actinomyces naeslundii		93.49	0.50	ESL 12		
Actinomyces israelii		6.51	0.21	MNS 88		
43773042	**Good identification**					**43773042**
Actinomyces israelii		91.98	0.58	XYL 92		
Actinomyces naeslundii		5.93	0.47			
Bifidobacterium breve		2.09	0.24			
43773043	**Acceptable identification**					**43773043**
Actinomyces naeslundii		90.55	0.50	ESL 12		
Actinomyces israelii		9.45	0.24	MNS 88		
43773050	**Good identification**					**43773050**
Actinomyces israelii		96.70	0.41	XYL 92		
Actinomyces naeslundii		3.30	0.25	MLZ 13		
43773052	**Low discrimination**					**43773052**
Actinomyces israelii		85.12	0.44	XYL 92		
Bifidobacterium breve		12.95	0.24	MLZ 13		
Actinomyces naeslundii		1.93	0.25			
43773060	**Low discrimination**					**43773060**
Actinomyces naeslundii		58.09	0.47	ESL 12		
Actinomyces israelii		38.32	0.34	MNS 88		
Bifidobacterium breve		3.60	0.12			
43773061	**Good identification**					**43773061**
Actinomyces naeslundii		100.0	0.50	ESL 12		
43773062	**Low discrimination**					**43773062**
Actinomyces naeslundii		38.23	0.47	ESL 12		
Actinomyces israelii		37.83	0.37	MNS 88		
Bifidobacterium breve		23.95	0.29			
43773063	**Good identification**					**43773063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	ESL 12		
43773100	**Low discrimination**					**43773100**
Actinomyces israelii		83.30	0.72	XYL 92		
Actinomyces naeslundii		12.64	0.67			
Bifidobacterium breve		2.70	0.41			
43773101	**Very good identification**					**43773101**
Actinomyces naeslundii		95.75	0.70	ESL 12		
Actinomyces israelii		4.25	0.38	MNS 88		
43773102	**Low discrimination**					**43773102**
Actinomyces israelii		68.97	0.75	XYL 92		
Bifidobacterium breve		15.09	0.59			
Actinomyces naeslundii		6.98	0.67			
43773103	**Good identification**					**43773103**
Actinomyces naeslundii		91.94	0.70	ESL 12		
Actinomyces israelii		6.12	0.41	MNS 88		
Bifidobacterium breve		1.34	0.24			
43773110	**Low discrimination**					**43773110**
Actinomyces israelii		76.83	0.58	XYL 92		
Bifidobacterium breve		16.69	0.41	MLZ 13		
Actinomyces naeslundii		4.11	0.46			
43773111	**Low discrimination**					**43773111**
Actinomyces naeslundii		88.81	0.49	ESL 12		
Actinomyces israelii		11.19	0.24	MNS 88		
43773112	**Low discrimination**					**43773112**
Bifidobacterium breve		41.62	0.59	MNS 99		
Actinomyces israelii		28.42	0.61			
Bifidobacterium adolescentis		22.70	0.53			
43773113	**Low discrimination**					**43773113**
Actinomyces naeslundii		60.55	0.49	ESL 12		
Bifidobacterium breve		16.76	0.24	MNS 88		
Actinomyces israelii		11.44	0.27	MLZ 5		
43773120	**Low discrimination**					**43773120**
					SPO	
Clostridium ramosum		71.48	0.53	MNS 99	92	
Actinomyces naeslundii		15.51	0.67		1	
Actinomyces israelii		6.52	0.52		1	
43773121	**Very good identification**					**43773121**
					SPO	
Actinomyces naeslundii		96.48	0.70	ESL 12	1	
Clostridium ramosum		3.00	0.19	MNS 88	92	
Actinomyces israelii		0.27	0.18		1	
43773122	**Low discrimination**					**43773122**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		76.45	0.64	MNS 99	92	
Bifidobacterium breve		15.94	0.63		1	
Actinomyces naeslundii		4.15	0.67		1	
43773123	**Low discrimination**					**43773123**
					SPO	
Actinomyces naeslundii		86.63	0.70	ESL 12	1	
Clostridium ramosum		10.76	0.30	MNS 88	92	
Bifidobacterium breve		2.24	0.29		1	
43773130	**Low discrimination**					**43773130**
Bifidobacterium breve		64.50	0.46	MNS 99		
Actinomyces israelii		10.66	0.38	BGL 91		
Actinomyces naeslundii		8.93	0.46			
43773131	**Good identification**					**43773131**
Actinomyces naeslundii		95.36	0.49	ESL 12		
Bifidobacterium breve		4.64	0.12	MNS 88		
43773132	**Low discrimination**					**43773132**
					SPO	
Bifidobacterium breve		87.99	0.63	MNS 99	1	
Clostridium ramosum		4.26	0.30		92	
Clostridium tertium		3.25	0.29		99	
43773133	**Low discrimination**					**43773133**
Actinomyces naeslundii		67.02	0.49	ESL 12		
Bifidobacterium breve		32.98	0.29	MNS 88		
43773140	**Low discrimination**					**43773140**
Actinomyces israelii		86.78	0.65	XYL 92		
Bifidobacterium breve		7.24	0.41			
Actinomyces naeslundii		5.98	0.55			
43773141	**Good identification**					**43773141**
Actinomyces naeslundii		91.09	0.58	ESL 12		
Actinomyces israelii		8.91	0.31	MNS 88		
43773142	**Low discrimination**					**43773142**
Actinomyces israelii		60.88	0.68	XYL 92		
Bifidobacterium breve		34.25	0.59			
Actinomyces naeslundii		2.79	0.55			
43773143	**Low discrimination**					**43773143**
Actinomyces naeslundii		81.35	0.58	ESL 12		
Actinomyces israelii		11.94	0.34	MNS 88		
Bifidobacterium breve		6.72	0.24			
43773150	**Low discrimination**					**43773150**
Actinomyces israelii		62.75	0.51	XYL 92		
Bifidobacterium breve		35.05	0.41	MLZ 13		
Actinomyces naeslundii		1.52	0.33			
43773151	**Low discrimination**					**43773151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		78.26	0.36	ESL 12		
Actinomyces israelii		21.74	0.17	MNS 88		
43773152	**Low discrimination**					**43773152**
Bifidobacterium breve		74.71	0.59	MNS 99		
Actinomyces israelii		19.84	0.54			
Bifidobacterium adolescentis		5.04	0.38			
43773160	**Low discrimination**					**43773160**
Bifidobacterium breve		49.59	0.46	MNS 99		
Actinomyces naeslundii		23.01	0.55	BGL 91		
Actinomyces israelii		21.33	0.45			
43773161	**Very good identification**					**43773161**
Actinomyces naeslundii		98.57	0.58	ESL 12		
Bifidobacterium breve		1.43	0.12	MNS 88		
43773162	**Low discrimination**					**43773162**
					SPO	
Bifidobacterium breve		86.12	0.63	MNS 99	1	
Actinomyces israelii		5.50	0.48		1	
Clostridium ramosum		4.17	0.30		92	
43773163	**Low discrimination**					**43773163**
Actinomyces naeslundii		86.50	0.58	ESL 12		
Bifidobacterium breve		12.69	0.29	MNS 88		
Actinomyces israelii		0.81	0.14			
43773170	**Acceptable identification**					**43773170**
Bifidobacterium breve		91.85	0.46	MNS 99		
Actinomyces israelii		5.90	0.31	BGL 91		
Actinomyces naeslundii		2.24	0.33			
43773171	**Low discrimination**					**43773171**
Actinomyces naeslundii		78.39	0.36	ESL 12		
Bifidobacterium breve		21.62	0.12	MNS 88		
43773172	**Very good identification**					**43773172**
Bifidobacterium breve		98.82	0.63	MNS 99		
Actinomyces israelii		0.94	0.34			
Actinomyces naeslundii		0.24	0.33			
43773200	**Good identification**					**43773200**
Actinomyces israelii		97.22	0.43	XYL 92		
Actinomyces naeslundii		2.78	0.25	RHA 7		
43773202	**Good identification**					**43773202**
					SPO	
Actinomyces israelii		96.85	0.46	XYL 92	1	
Actinomyces naeslundii		1.85	0.25	RHA 7	1	
Clostridium ramosum		1.30	-.03		92	
43773240	**Good identification**					**43773240**
Actinomyces israelii		100.0	0.36	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
43773242	**Good identification**					**43773242**
Actinomyces israelii		100.0	0.39	XYL 92		
43773300	**Low discrimination**					**43773300**
					SPO	
Actinomyces israelii		87.35	0.53	XYL 92	1	
Clostridium ramosum		10.87	0.21	RHA 7	92	
Actinomyces naeslundii		1.78	0.33		1	
43773301	**Low discrimination**					**43773301**
Actinomyces naeslundii		75.15	0.36	ESL 12		
Actinomyces israelii		24.85	0.19	MNS 88		
43773302	**Low discrimination**					**43773302**
					SPO	
Actinomyces israelii		71.57	0.56	XYL 92	1	
Clostridium ramosum		23.75	0.32	RHA 7	92	
Bifidobacterium breve		2.10	0.24		1	
43773303	**Low discrimination**					**43773303**
					SPO	
Actinomyces naeslundii		60.21	0.36	ESL 12	1	
Actinomyces israelii		29.88	0.22	MNS 88	1	
Clostridium ramosum		9.91	-.03	RHA 1	92	
43773310	**Good identification**					**43773310**
Actinomyces israelii		100.0	0.39	XYL 92		
43773312	**Low discrimination**					**43773312**
Actinomyces israelii		74.02	0.42	XYL 92		
Bifidobacterium breve		14.55	0.24	RHA 7		
Bifidobacterium adolescentis		7.93	0.19	MLZ 13		
43773320	**Good identification**					**43773320**
					SPO	
Clostridium ramosum		99.26	0.56	MNS 99	92	
Actinomyces israelii		0.51	0.33		1	
Actinomyces naeslundii		0.16	0.33		1	
43773322	**Very good identification**					**43773322**
					SPO	
Clostridium ramosum		99.53	0.66	MNS 99	92	
Actinomyces israelii		0.19	0.36		1	
Bifidobacterium breve		0.16	0.29		1	
43773340	**Good identification**					**43773340**
Actinomyces israelii		100.0	0.46	XYL 92		
43773342	**Acceptable identification**					**43773342**
					SPO	
Actinomyces israelii		92.24	0.49	XYL 92	1	
Bifidobacterium breve		6.97	0.24	RHA 7	1	
Clostridium ramosum		0.80	-.03		92	
43773400	**Very good identification**					**43773400**
					SPO	
Actinomyces israelii		99.24	0.60	XYL 92	1	
Clostridium butyricum		0.50	0.00		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.26	0.25		1	
43773402	**Very good identification**					**43773402**
Actinomyces israelii		96.59	0.63	XYL 92		
Bifidobacterium adolescentis		3.12	0.32			
Actinomyces naeslundii		0.17	0.25			
43773410	**Good identification**					**43773410**
Actinomyces israelii		100.0	0.46	XYL 92		
43773412	**Low discrimination**					**43773412**
Actinomyces israelii		78.25	0.49	XYL 92		
Bifidobacterium adolescentis		20.65	0.33	MLZ 13		
Bifidobacterium dentium		1.10	-.07			
43773420	**Good identification**					**43773420**
Actinomyces israelii		96.04	0.40	XYL 92		
Actinomyces naeslundii		3.96	0.25	NAG 6		
43773422	**Good identification**					**43773422**
Actinomyces israelii		97.32	0.43	XYL 92		
Actinomyces naeslundii		2.68	0.25	NAG 6		
43773440	**Good identification**					**43773440**
Actinomyces israelii		100.0	0.53	XYL 92		
43773442	**Very good identification**					**43773442**
Actinomyces israelii		98.99	0.56	XYL 92		
Bifidobacterium adolescentis		1.02	0.16			
43773450	**Acceptable identification**					**43773450**
					COC	
Actinomyces israelii		94.94	0.39	XYL 92	1	
Blautia producta		5.07	-.03	MLZ 13	99	
43773452	**Acceptable identification**					**43773452**
Actinomyces israelii		92.26	0.42	XYL 92		
Bifidobacterium adolescentis		7.74	0.18	MLZ 13		
43773462	**Good identification**					**43773462**
Actinomyces israelii		100.0	0.36	XYL 92		
43773500	**Very good identification**					**43773500**
					SPO	
Actinomyces israelii		95.75	0.71	XYL 92	1	
Clostridium butyricum		2.91	0.24		99	
Bifidobacterium adolescentis		1.16	0.32		1	
43773501	**Low discrimination**					**43773501**
Actinomyces israelii		78.24	0.37	XYL 92		
Actinomyces naeslundii		21.76	0.36	URE 1		
43773502	**Low discrimination**					**43773502**
					SPO	
Actinomyces israelii		55.22	0.74	XYL 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		44.10	0.66		1		
Clostridium butyricum		0.44	0.17		99		
43773503	**Low discrimination**						**43773503**
Actinomyces israelii		50.40	0.40	XYL 92			
Bifidobacterium adolescentis		40.25	0.32	URE 1			
Actinomyces naeslundii		9.35	0.36				
43773510	**Low discrimination**						**43773510**
Actinomyces israelii		85.74	0.57	XYL 92			
Bifidobacterium adolescentis		8.49	0.33	MLZ 13			
Bifidobacterium dentium		4.42	0.10				
43773512	**Low discrimination**						**43773512**
Bifidobacterium adolescentis		82.66	0.67	MNS 87			
Actinomyces israelii		12.65	0.60	XYL 87			
Bifidobacterium dentium		4.40	0.27				
43773520	**Low discrimination**						**43773520**
					SPO		
Actinomyces israelii		82.16	0.50	XYL 92	1		
Clostridium ramosum		11.11	0.19	NAG 6	92		
Clostridium butyricum		3.40	0.06		99		
43773521	**Low discrimination**						**43773521**
Actinomyces naeslundii		81.34	0.36	ESL 12			
Actinomyces israelii		18.66	0.16	MNS 88			
43773522	**Low discrimination**						**43773522**
					SPO		
Actinomyces israelii		62.22	0.53	XYL 92	1		
Clostridium ramosum		22.45	0.30	NAG 6	92		
Bifidobacterium adolescentis		7.86	0.32		1		
43773523	**Low discrimination**						**43773523**
Actinomyces naeslundii		74.39	0.36	ESL 12			
Actinomyces israelii		25.61	0.19	MNS 88			
43773540	**Good identification**						**43773540**
					COC	SPO	
Actinomyces israelii		92.90	0.64	XYL 92	1	1	
Blautia producta		6.11	0.24		99	1	
Clostridium butyricum		0.63	0.06		1	99	
43773542	**Low discrimination**						**43773542**
Actinomyces israelii		78.90	0.67	XYL 92			
Bifidobacterium adolescentis		20.03	0.50				
Bifidobacterium breve		0.55	0.24				
43773552	**Low discrimination**						**43773552**
					COC		
Bifidobacterium adolescentis		64.29	0.52	MNS 87	1		
Actinomyces israelii		30.97	0.53	XYL 87	1		
Blautia producta		3.03	0.15	SOR 11	99		
43773560	**Low discrimination**						**43773560**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		87.47	0.43	XYL 92	1		
Blautia producta		10.02	0.07	NAG 6	99		
Bifidobacterium breve		2.51	0.12		1		
43773562	**Low discrimination**						**43773562**
Actinomyces israelii		81.06	0.46	XYL 92			
Bifidobacterium breve		15.68	0.29	NAG 6			
Bifidobacterium adolescentis		3.26	0.16				
43773600	**Good identification**						**43773600**
Actinomyces israelii		100.0	0.41	XYL 92			
43773602	**Good identification**						**43773602**
Actinomyces israelii		100.0	0.44	XYL 92			
43773642	**Good identification**						**43773642**
Actinomyces israelii		100.0	0.37	XYL 92			
43773700	**Good identification**						**43773700**
					SPO		
Actinomyces israelii		98.77	0.52	XYL 92	1		
Clostridium butyricum		1.23	-.02	RHA 7	99		
43773702	**Low discrimination**						**43773702**
					SPO		
Actinomyces israelii		89.99	0.55	XYL 92	1		
Bifidobacterium adolescentis		9.64	0.32	RHA 7	1		
Clostridium ramosum		0.37	-.03		92		
43773710	**Acceptable identification**						**43773710**
					COC		
Actinomyces israelii		93.55	0.37	XYL 92	1		
Blautia producta		6.45	-.03	RHA 7	99		
43773712	**Low discrimination**						**43773712**
Actinomyces israelii		49.19	0.40	XYL 92			
Bifidobacterium adolescentis		43.12	0.33	RHA 7			
Clostridium clostridioforme		5.40	0.15	MLZ 13			
43773740	**Low discrimination**						**43773740**
					COC		
Actinomyces israelii		53.36	0.44	XYL 92	1		
Blautia producta		46.64	0.24	RHA 7	99		
43773742	**Acceptable identification**						**43773742**
					COC		
Actinomyces israelii		91.01	0.47	XYL 92	1		
Blautia producta		5.89	0.07	RHA 7	99		
Bifidobacterium adolescentis		3.10	0.16		1		
43774002	**Low discrimination**						**43774002**
					GRA		
Actinomyces israelii		89.72	0.36	GAL 99	99		
Fusobacterium mortiferum		10.28	0.18		1		
43774100	**Low discrimination**						**43774100**
					GRA	SPO	
Actinomyces israelii		43.50	0.43	GAL 99	99	1	
Fusobacterium mortiferum		39.27	0.40		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		17.23	0.15		99	99	
43774102	**Low discrimination**						**43774102**
					SPO		
Actinomyces israelii		62.72	0.46	GAL 99	1		
Bifidobacterium adolescentis		24.68	0.33		1		
Clostridium butyricum		6.44	0.08		99		
43774140	**Acceptable identification**						**43774140**
					SPO		
Actinomyces israelii		91.86	0.36	GAL 99	1		
Clostridium butyricum		8.14	-.03		99		
43774142	**Low discrimination**						**43774142**
Actinomyces israelii		88.88	0.39	GAL 99			
Bifidobacterium adolescentis		11.12	0.18				
43774312	**Good identification**						**43774312**
Clostridium clostridioforme		100.0	0.37	MNS 99			
43774332	**Low discrimination**						**43774332**
Clostridium clostridioforme		88.50	0.37	MNS 99			
Parabacteroides distasonis		11.50	0.28	ARA 94			
43774502	**Low discrimination**						**43774502**
					SPO		
Bifidobacterium adolescentis		54.10	0.47	MNS 87	1		
Clostridium butyricum		18.98	0.25	GAL 99	99		
Actinomyces israelii		16.81	0.45		1		
43774512	**Low discrimination**						**43774512**
Bifidobacterium adolescentis		56.95	0.49	MNS 87			
Clostridium clostridioforme		39.66	0.43	GAL 99			
Actinomyces israelii		2.16	0.31				
43774532	**Good identification**						**43774532**
Clostridium clostridioforme		95.83	0.43	MNS 99			
Bifidobacterium dentium		1.59	-.03	RHA 88			
Bifidobacterium adolescentis		1.39	0.15				
43774602	**Good identification**						**43774602**
Clostridium clostridioforme		98.59	0.37	MNS 99			
Actinomyces israelii		1.41	0.15				
43774612	**Good identification**						**43774612**
Clostridium clostridioforme		100.0	0.49	MNS 99			
43774622	**Good identification**						**43774622**
Clostridium clostridioforme		100.0	0.37	MNS 99			
43774632	**Good identification**						**43774632**
Clostridium clostridioforme		100.0	0.49	MNS 99			
43774702	**Good identification**						**43774702**
					SPO		
Clostridium clostridioforme		96.87	0.46	MNS 99	66		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.65	0.26		1		
Clostridium butyricum		0.77	-.01		99		
43774712	**Very good identification**						**43774712**
Clostridium clostridioforme		99.80	0.58	MNS 99			
Bifidobacterium adolescentis		0.20	0.15				
43774722	**Acceptable identification**						**43774722**
Clostridium clostridioforme		94.33	0.46	MNS 99			
Parabacteroides distasonis		4.06	0.29				
Bacteroides ovatus		1.61	0.06				
43774732	**Very good identification**						**43774732**
Clostridium clostridioforme		99.25	0.58	MNS 99			
Parabacteroides distasonis		0.51	0.24				
Bacteroides ovatus		0.25	0.03				
43775000	**Good identification**						**43775000**
					GRA	SPO	
Actinomyces israelii		91.79	0.67		99	1	
Clostridium butyricum		6.06	0.26		99	99	
Fusobacterium mortiferum		2.15	0.37		1	1	
43775002	**Very good identification**						**43775002**
					SPO		
Actinomyces israelii		96.54	0.70		1		
Clostridium butyricum		1.65	0.19		99		
Bifidobacterium adolescentis		1.54	0.33		1		
43775003	**Acceptable identification**						**43775003**
Actinomyces israelii		94.86	0.36	URE 1			
Actinomyces naeslundii		5.14	0.24				
43775010	**Good identification**						**43775010**
					SPO		
Actinomyces israelii		97.73	0.53	MLZ 13	1		
Clostridium butyricum		2.27	0.04		99		
43775012	**Low discrimination**						**43775012**
Actinomyces israelii		85.15	0.56	MLZ 13			
Bifidobacterium adolescentis		11.07	0.35				
Clostridium clostridioforme		2.95	0.22				
43775020	**Acceptable identification**						**43775020**
					SPO		
Actinomyces israelii		91.75	0.47	NAG 6	1		
Clostridium butyricum		8.25	0.08		99		
43775022	**Low discrimination**						**43775022**
					GRA		
Actinomyces israelii		88.09	0.50	NAG 6	99		
Parabacteroides distasonis		4.71	0.24		1		
Prevotella oralis		3.58	0.09		1		
43775032	**Low discrimination**						**43775032**
					GRA		
Actinomyces israelii		57.61	0.36	MLZ 13	99		
Clostridium clostridioforme		31.30	0.22	NAG 6	50		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		11.09	0.20		1		
43775040	**Very good identification**						**43775040**
					SPO		
Actinomyces israelii		98.55	0.60		1		
Clostridium butyricum		1.45	0.08		99		
43775042	**Very good identification**						**43775042**
					SPO		
Actinomyces israelii		99.12	0.63		1		
Bifidobacterium adolescentis		0.50	0.18		1		
Clostridium butyricum		0.38	0.01		99		
43775050	**Good identification**						**43775050**
Actinomyces israelii		100.0	0.46	MLZ 13			
43775052	**Good identification**						**43775052**
Actinomyces israelii		96.03	0.49	MLZ 13			
Bifidobacterium adolescentis		3.97	0.19				
43775060	**Good identification**						**43775060**
Actinomyces israelii		100.0	0.40	NAG 6			
43775062	**Good identification**						**43775062**
Actinomyces israelii		100.0	0.43	NAG 6			
43775100	**Low discrimination**						**43775100**
					GRA	SPO	
Actinomyces israelii		69.95	0.77		99	1	
Clostridium butyricum		27.70	0.50		99	99	
Fusobacterium mortiferum		0.96	0.43		1	1	
43775101	**Low discrimination**						**43775101**
					SPO		
Actinomyces israelii		68.79	0.43	URE 1	1		
Clostridium butyricum		27.24	0.15		99		
Actinomyces naeslundii		3.97	0.31		1		
43775102	**Low discrimination**						**43775102**
					SPO		
Actinomyces israelii		66.51	0.80		1		
Bifidobacterium adolescentis		26.17	0.67		1		
Clostridium butyricum		6.83	0.43		99		
43775103	**Low discrimination**						**43775103**
					SPO		
Actinomyces israelii		65.16	0.46	URE 1	1		
Bifidobacterium adolescentis		25.64	0.33		1		
Clostridium butyricum		6.69	0.08		99		
43775110	**Low discrimination**						**43775110**
					SPO		
Actinomyces israelii		84.06	0.63	MLZ 13	1		
Clostridium butyricum		11.73	0.28		99		
Bifidobacterium adolescentis		4.10	0.35		1		
43775112	**Low discrimination**						**43775112**
					GRA		
Bifidobacterium adolescentis		73.37	0.69	MNS 87	99		
Actinomyces israelii		22.80	0.66	ARA 87	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		2.08	0.24		1		
43775120	**Low discrimination**						**43775120**
					COC	SPO	
Actinomyces israelii		60.82	0.57	NAG 6	1	1	
Clostridium butyricum		32.81	0.31		1	99	
Atopobium parvulum		2.29	0.19		99	1	
43775122	**Low discrimination**						**43775122**
					GRA		
Parabacteroides distasonis		47.46	0.58	MNS 98	1		
Actinomyces israelii		35.88	0.60		99		
Prevotella oralis		5.71	0.30		1		
43775130	**Low discrimination**						**43775130**
					GRA	SPO	
Actinomyces israelii		77.38	0.43	MLZ 13	99	1	
Clostridium butyricum		14.70	0.10	NAG 6	99	99	
Parabacteroides distasonis		5.59	0.20		1	1	
43775132	**Low discrimination**						**43775132**
					GRA		
Parabacteroides distasonis		66.38	0.54	MNS 98	1		
Actinomyces israelii		13.93	0.46		99		
Bifidobacterium adolescentis		7.09	0.35		99		
43775140	**Good identification**						**43775140**
					SPO		
Actinomyces israelii		91.71	0.70		1		
Clostridium butyricum		8.12	0.31		99		
Bifidobacterium adolescentis		0.17	0.18		1		
43775141	**Acceptable identification**						**43775141**
					SPO		
Actinomyces israelii		91.86	0.36	URE 1	1		
Clostridium butyricum		8.14	-.03		99		
43775142	**Low discrimination**						**43775142**
					SPO		
Actinomyces israelii		86.89	0.73		1		
Bifidobacterium adolescentis		10.87	0.52		1		
Clostridium butyricum		2.00	0.24		99		
43775143	**Low discrimination**						**43775143**
Actinomyces israelii		88.88	0.39	URE 1			
Bifidobacterium adolescentis		11.12	0.18				
43775150	**Very good identification**						**43775150**
					SPO		
Actinomyces israelii		95.05	0.56	MLZ 13	1		
Clostridium butyricum		2.97	0.10		99		
Bifidobacterium adolescentis		1.47	0.19		1		
43775152	**Low discrimination**						**43775152**
Bifidobacterium adolescentis		49.85	0.53	MNS 87			
Actinomyces israelii		48.73	0.59	ARA 87			
Bifidobacterium breve		0.91	0.24	SOR 11			
43775160	**Low discrimination**						**43775160**
					SPO		
Actinomyces israelii		88.32	0.50	NAG 6	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		10.66	0.13		99	
Bifidobacterium breve		1.02	0.12		1	
43775162	**Low discrimination**					**43775162**
					GRA	
Actinomyces israelii		85.95	0.53	NAG 6	99	
Bifidobacterium breve		6.70	0.29		99	
Parabacteroides distasonis		2.95	0.24		1	
43775170	**Acceptable identification**					**43775170**
Actinomyces israelii		92.81	0.36	MLZ 13		
Bifidobacterium breve		7.19	0.12	NAG 6		
43775172	**Low discrimination**					**43775172**
Actinomyces israelii		55.32	0.39	MLZ 13		
Bifidobacterium breve		28.88	0.29	NAG 6		
Bifidobacterium adolescentis		8.96	0.19			
43775200	**Good identification**					**43775200**
					SPO	
Actinomyces israelii		97.36	0.48	RHA 7	1	
Clostridium butyricum		2.64	0.00		99	
43775202	**Low discrimination**					**43775202**
					SPO	
Actinomyces israelii		89.46	0.51	RHA 7	1	
Clostridium clostridioforme		9.91	0.25		66	
Clostridium butyricum		0.63	-.07		99	
43775212	**Low discrimination**					**43775212**
Clostridium clostridioforme		77.16	0.36	MNS 99		
Actinomyces israelii		22.84	0.37	ARA 94		
43775232	**Low discrimination**					**43775232**
					GRA	
Clostridium clostridioforme		85.92	0.36	MNS 99	50	
Parabacteroides distasonis		12.46	0.28	ARA 94	1	
Actinomyces israelii		1.62	0.16		99	
43775240	**Good identification**					**43775240**
Actinomyces israelii		100.0	0.41	RHA 7		
43775242	**Good identification**					**43775242**
Actinomyces israelii		100.0	0.44	RHA 7		
43775300	**Low discrimination**					**43775300**
					SPO	
Actinomyces israelii		78.32	0.58	RHA 7	1	
Clostridium butyricum		12.75	0.24		99	
Eubacterium contortum		8.93	0.27		1	
43775302	**Low discrimination**					**43775302**
Actinomyces israelii		80.76	0.61	RHA 7		
Clostridium clostridioforme		7.49	0.34			
Bifidobacterium adolescentis		4.27	0.33			
43775310	**Acceptable identification**					**43775310**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		90.90	0.44	RHA 7	1	
Clostridium butyricum		5.21	0.02	MLZ 13	99	
Clostridium clostridioforme		3.89	0.11		66	
43775312	**Low discrimination**					**43775312**
Clostridium clostridioforme		64.13	0.45	MNS 99		
Actinomyces israelii		22.68	0.47	ARA 94		
Bifidobacterium adolescentis		9.80	0.35			
43775320	**Low discrimination**					**43775320**
					GRA	
Actinomyces israelii		38.81	0.38	RHA 7	99	
Parabacteroides distasonis		31.00	0.32	NAG 6	1	
Eubacterium contortum		12.24	0.14		99	
43775322	**Good identification**					**43775322**
					GRA	
Parabacteroides distasonis		92.46	0.67	MNS 98	1	
Clostridium clostridioforme		2.55	0.34		50	
Actinomyces israelii		1.75	0.41		99	
43775332	**Low discrimination**					**43775332**
Parabacteroides distasonis		80.14	0.62	MNS 98		
Clostridium clostridioforme		18.69	0.45			
Bacteroides ovatus		0.75	0.11			
43775340	**Good identification**					**43775340**
					SPO	
Actinomyces israelii		96.49	0.51	RHA 7	1	
Clostridium butyricum		3.51	0.06		99	
43775342	**Good identification**					**43775342**
					SPO	
Actinomyces israelii		97.45	0.54	RHA 7	1	
Bifidobacterium adolescentis		1.64	0.18		1	
Clostridium butyricum		0.92	-.02		99	
43775350	**Acceptable identification**					**43775350**
					COC	
Actinomyces israelii		93.33	0.37	RHA 7	1	
Blautia producta		6.67	-.03	MLZ 13	99	
43775352	**Low discrimination**					**43775352**
Actinomyces israelii		82.60	0.40	RHA 7		
Bifidobacterium adolescentis		11.34	0.19	MLZ 13		
Clostridium clostridioforme		6.07	0.11			
43775400	**Low discrimination**					**43775400**
					SPO	
Actinomyces israelii		57.87	0.66		1	
Clostridium butyricum		42.02	0.42		99	
Bifidobacterium adolescentis		0.11	0.13		1	
43775402	**Low discrimination**					**43775402**
					SPO	
Actinomyces israelii		74.62	0.69		1	
Clostridium butyricum		14.05	0.35		99	
Bifidobacterium adolescentis		9.71	0.47		1	
43775410	**Low discrimination**					**43775410**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		78.01	0.52	MLZ 13	1		
Clostridium butyricum		19.95	0.20		99		
Bifidobacterium adolescentis		1.26	0.15		1		
43775412	**Low discrimination**						**43775412**
Bifidobacterium adolescentis		37.98	0.49	MNS 87			
Actinomyces israelii		35.70	0.55	LAC 99			
Clostridium clostridioforme		23.70	0.42				
43775420	**Low discrimination**						**43775420**
					SPO		
Actinomyces israelii		50.28	0.45	NAG 6	1		
Clostridium butyricum		49.73	0.24		99		
43775422	**Low discrimination**						**43775422**
					SPO		
Actinomyces israelii		60.31	0.48	NAG 6	1		
Clostridium clostridioforme		20.57	0.31		66		
Clostridium butyricum		15.46	0.17		99		
43775432	**Low discrimination**						**43775432**
Clostridium clostridioforme		86.70	0.42	MNS 99			
Actinomyces israelii		8.34	0.34	RHA 88			
Bifidobacterium dentium		1.61	-.03				
43775440	**Low discrimination**						**43775440**
					SPO		
Actinomyces israelii		86.03	0.59		1		
Clostridium butyricum		13.97	0.24		99		
43775442	**Good identification**						**43775442**
					SPO		
Actinomyces israelii		92.30	0.62		1		
Clostridium butyricum		3.89	0.17		99		
Bifidobacterium adolescentis		3.82	0.32		1		
43775450	**Acceptable identification**						**43775450**
					COC	SPO	
Actinomyces israelii		92.30	0.45	MLZ 13	1	1	
Clostridium butyricum		5.28	0.02		1	99	
Blautia producta		2.43	-.03		99	1	
43775452	**Low discrimination**						**43775452**
Actinomyces israelii		72.99	0.48	MLZ 13			
Bifidobacterium adolescentis		24.68	0.33				
Clostridium clostridioforme		1.26	0.08				
43775460	**Low discrimination**						**43775460**
					SPO		
Actinomyces israelii		81.89	0.38	NAG 6	1		
Clostridium butyricum		18.11	0.06		99		
43775462	**Acceptable identification**						**43775462**
					SPO		
Actinomyces israelii		94.58	0.41	NAG 6	1		
Clostridium butyricum		5.42	-.01		99		
43775500	**Low discrimination**						**43775500**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		80.54	0.66	MNS 99	99		
Actinomyces israelii		18.49	0.76		1		
Bifidobacterium adolescentis		0.90	0.47		1		
43775502	**Low discrimination**						**43775502**
					SPO		
Bifidobacterium adolescentis		59.97	0.81	MNS 87	1		
Clostridium butyricum		21.04	0.59		99		
Actinomyces israelii		18.63	0.79		1		
43775503	**Low discrimination**						**43775503**
					SPO		
Bifidobacterium adolescentis		59.98	0.47	MNS 87	1		
Clostridium butyricum		21.05	0.25	URE 1	99		
Actinomyces israelii		18.64	0.45		1		
43775510	**Low discrimination**						**43775510**
					SPO		
Clostridium butyricum		51.54	0.44	MNS 99	99		
Actinomyces israelii		33.59	0.62	MLZ 5	1		
Bifidobacterium adolescentis		13.40	0.49		1		
43775512	**Good identification**						**43775512**
Bifidobacterium adolescentis		92.52	0.83	MNS 87			
Actinomyces israelii		3.51	0.65				
Clostridium clostridioforme		1.95	0.51				
43775513	**Acceptable identification**						**43775513**
Bifidobacterium adolescentis		92.53	0.49	MNS 87			
Actinomyces israelii		3.52	0.31	URE 1			
Clostridium clostridioforme		1.95	0.17				
43775520	**Low discrimination**						**43775520**
					GRA	SPO	
Clostridium butyricum		85.03	0.48	MNS 99	99	99	
Actinomyces israelii		14.33	0.56	NAG 8	99	1	
Parabacteroides distasonis		0.22	0.21		1	1	
43775522	**Low discrimination**						**43775522**
					GRA	SPO	
Clostridium butyricum		34.91	0.41	MNS 99	99	99	
Actinomyces israelii		22.69	0.59	NAG 8	99	1	
Parabacteroides distasonis		22.48	0.55		1	1	
43775532	**Low discrimination**						**43775532**
Clostridium clostridioforme		37.77	0.51	MNS 99			
Bifidobacterium dentium		20.68	0.32	RHA 88			
Bifidobacterium adolescentis		18.08	0.49				
43775540	**Low discrimination**						**43775540**
					COC	SPO	
Actinomyces israelii		49.25	0.69		1	1	
Clostridium butyricum		47.97	0.48		1	99	
Blautia producta		1.60	0.24		99	1	
43775542	**Low discrimination**						**43775542**
					SPO		
Bifidobacterium adolescentis		44.89	0.66	MNS 87	1		
Actinomyces israelii		43.88	0.72	SOR 11	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		11.08	0.41		99		
43775550	**Low discrimination**						**43775550**
					COC	SPO	
Actinomyces israelii		46.74	0.55	MLZ 13	1	1	
Blautia producta		30.42	0.32		99	1	
Clostridium butyricum		16.04	0.26		1	99	
43775552	**Low discrimination**						**43775552**
					SPO		
Bifidobacterium adolescentis		87.86	0.67	MNS 87	1		
Actinomyces israelii		10.50	0.58	SOR 11	1		
Clostridium butyricum		0.93	0.19		99		
43775562	**Low discrimination**						**43775562**
					SPO		
Actinomyces israelii		64.70	0.52	NAG 6	1		
Clostridium butyricum		22.26	0.23		99		
Bifidobacterium adolescentis		10.47	0.32		1		
43775600	**Low discrimination**						**43775600**
					SPO		
Actinomyces israelii		75.16	0.46	RHA 7	1		
Clostridium butyricum		22.42	0.16		99		
Clostridium clostridioforme		2.42	0.11		66		
43775602	**Low discrimination**						**43775602**
					SPO		
Clostridium clostridioforme		65.96	0.46	MNS 99	66		
Actinomyces israelii		31.09	0.49		1		
Clostridium butyricum		2.41	0.09		99		
43775612	**Very good identification**						**43775612**
Clostridium clostridioforme		98.26	0.57	MNS 99			
Actinomyces israelii		1.52	0.35				
Bifidobacterium adolescentis		0.22	0.15				
43775622	**Acceptable identification**						**43775622**
					GRA		
Clostridium clostridioforme		91.15	0.46	MNS 99	50		
Parabacteroides distasonis		4.38	0.29		1		
Actinomyces israelii		2.74	0.29		99		
43775632	**Very good identification**						**43775632**
Clostridium clostridioforme		99.06	0.57	MNS 99			
Parabacteroides distasonis		0.56	0.24				
Bacteroides ovatus		0.28	0.03				
43775640	**Acceptable identification**						**43775640**
					SPO		
Actinomyces israelii		93.75	0.39	RHA 7	1		
Clostridium butyricum		6.25	-.02		99		
43775642	**Acceptable identification**						**43775642**
Actinomyces israelii		94.78	0.42	RHA 7			
Clostridium clostridioforme		5.22	0.11				
43775700	**Low discrimination**						**43775700**
					SPO		
Clostridium butyricum		59.30	0.40	MNS 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		33.13	0.57	RHA 3	1		
Eubacterium contortum		3.78	0.26		1		
43775702	**Low discrimination**						**43775702**
					SPO		
Clostridium clostridioforme		47.65	0.55	MNS 99	66		
Actinomyces israelii		26.83	0.60		1		
Clostridium butyricum		12.45	0.33		99		
43775710	**Low discrimination**						**43775710**
					SPO		
Actinomyces israelii		36.79	0.43	RHA 7	1		
Clostridium clostridioforme		30.17	0.32	MLZ 13	66		
Clostridium butyricum		23.19	0.18		99		
43775712	**Good identification**						**43775712**
Clostridium clostridioforme		91.75	0.66	MNS 99			
Bifidobacterium adolescentis		5.99	0.49				
Actinomyces israelii		1.70	0.46				
43775722	**Low discrimination**						**43775722**
Parabacteroides distasonis		46.04	0.63	MNS 98			
Clostridium clostridioforme		32.44	0.55				
Bacteroides ovatus		18.25	0.40				
43775732	**Low discrimination**						**43775732**
Clostridium clostridioforme		79.63	0.66	MNS 99			
Parabacteroides distasonis		13.36	0.58				
Bacteroides ovatus		6.56	0.37				
43775740	**Low discrimination**						**43775740**
					COC	SPO	
Actinomyces israelii		49.05	0.50	RHA 7	1	1	
Blautia producta		21.12	0.24		99	1	
Clostridium butyricum		19.63	0.22		1	99	
43775742	**Low discrimination**						**43775742**
					SPO		
Actinomyces israelii		74.94	0.53	RHA 7	1		
Bifidobacterium adolescentis		10.29	0.32		1		
Clostridium butyricum		7.78	0.15		99		
43776000	**Good identification**						**43776000**
Actinomyces israelii		100.0	0.46	GAL 99			
43776002	**Good identification**						**43776002**
Actinomyces israelii		100.0	0.49	GAL 99			
43776040	**Good identification**						**43776040**
Actinomyces israelii		100.0	0.39	GAL 99			
43776042	**Good identification**						**43776042**
Actinomyces israelii		100.0	0.42	GAL 99			
43776100	**Very good identification**						**43776100**
					SPO		
Actinomyces israelii		99.23	0.56	GAL 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		0.77	-.01		99	
43776102	**Good identification**					**43776102**
Actinomyces israelii		94.62	0.59	GAL 99		
Bifidobacterium adolescentis		5.38	0.32			
43776110	**Good identification**					**43776110**
Actinomyces israelii		100.0	0.42	GAL 99		
43776112	**Low discrimination**					**43776112**
Actinomyces israelii		67.14	0.45	GAL 99		
Bifidobacterium adolescentis		31.20	0.33	MLZ 13		
Bifidobacterium dentium		1.66	-.07			
43776120	**Good identification**					**43776120**
Actinomyces israelii		100.0	0.36	GAL 99		
43776122	**Good identification**					**43776122**
Actinomyces israelii		100.0	0.39	GAL 99		
43776140	**Good identification**					**43776140**
Actinomyces israelii		100.0	0.49	GAL 99		
43776142	**Good identification**					**43776142**
Actinomyces israelii		98.23	0.52	GAL 99		
Bifidobacterium adolescentis		1.77	0.16			
43776150	**Acceptable identification**					**43776150**
					COC	
Actinomyces israelii		91.41	0.35	GAL 99	1	
Blautia producta		8.59	-.03	MLZ 13	99	
43776152	**Low discrimination**					**43776152**
Actinomyces israelii		87.13	0.38	GAL 99		
Bifidobacterium adolescentis		12.87	0.18	MLZ 13		
43776300	**Good identification**					**43776300**
Actinomyces israelii		100.0	0.37	GAL 99		
43776302	**Good identification**					**43776302**
Actinomyces israelii		100.0	0.40	GAL 99		
43776400	**Good identification**					**43776400**
Actinomyces israelii		100.0	0.44	GAL 99		
43776402	**Good identification**					**43776402**
Actinomyces israelii		98.16	0.47	GAL 99		
Bifidobacterium adolescentis		1.84	0.12			
43776440	**Good identification**					**43776440**
Actinomyces israelii		100.0	0.37	GAL 99		
43776442	**Good identification**					**43776442**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	GAL 99			
43776500	**Acceptable identification**						**43776500**
					SPO		
Actinomyces israelii		91.54	0.55	GAL 99	1		
Clostridium butyricum		7.82	0.15		99		
Bifidobacterium adolescentis		0.65	0.12		1		
43776502	**Low discrimination**						**43776502**
					SPO		
Actinomyces israelii		67.05	0.58	GAL 99	1		
Bifidobacterium adolescentis		31.16	0.46		1		
Clostridium butyricum		1.49	0.08		99		
43776510	**Low discrimination**						**43776510**
Actinomyces israelii		63.46	0.41	GAL 99			
Bifidobacterium dentium		28.15	0.10	MLZ 13			
Bifidobacterium adolescentis		3.66	0.13				
43776512	**Low discrimination**						**43776512**
Bifidobacterium adolescentis		48.59	0.47	MNS 87			
Bifidobacterium dentium		38.21	0.27	GAL 99			
Actinomyces israelii		12.78	0.44				
43776522	**Low discrimination**						**43776522**
Actinomyces israelii		83.39	0.37	GAL 99			
Bifidobacterium dentium		10.48	-.03	NAG 6			
Bifidobacterium adolescentis		6.13	0.12				
43776540	**Low discrimination**						**43776540**
					COC	SPO	
Actinomyces israelii		63.08	0.48	GAL 99	1	1	
Blautia producta		35.72	0.24		99	1	
Clostridium butyricum		1.21	-.03		1	99	
43776542	**Low discrimination**						**43776542**
					COC		
Actinomyces israelii		84.06	0.51	GAL 99	1		
Bifidobacterium adolescentis		12.42	0.30		1		
Blautia producta		3.53	0.07		99		
43776700	**Good identification**						**43776700**
Actinomyces israelii		100.0	0.35	GAL 99			
43776702	**Low discrimination**						**43776702**
Actinomyces israelii		85.70	0.39	GAL 99			
Clostridium clostridioforme		8.95	0.12	RHA 7			
Bifidobacterium adolescentis		5.35	0.12				
43777000	**Very good identification**						**43777000**
					SPO		
Actinomyces israelii		99.85	0.80		1		
Clostridium butyricum		0.13	0.10		99		
Actinomyces naeslundii		0.02	0.25		1		
43777001	**Good identification**						**43777001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		97.29	0.46	URE 1		
Actinomyces naeslundii		2.71	0.28			
43777002	**Very good identification**					**43777002**
					SPO	
Actinomyces israelii		99.73	0.83		1	
Bifidobacterium adolescentis		0.23	0.32		1	
Clostridium butyricum		0.03	0.03		99	
43777003	**Good identification**					**43777003**
Actinomyces israelii		98.18	0.49	URE 1		
Actinomyces naeslundii		1.82	0.28			
43777010	**Very good identification**					**43777010**
Actinomyces israelii		100.0	0.66	MLZ 13		
43777012	**Very good identification**					**43777012**
Actinomyces israelii		98.06	0.69	MLZ 13		
Bifidobacterium adolescentis		1.84	0.33			
Bifidobacterium dentium		0.10	-.07			
43777020	**Very good identification**					**43777020**
Actinomyces israelii		99.71	0.60	NAG 6		
Actinomyces naeslundii		0.29	0.25			
43777022	**Very good identification**					**43777022**
Actinomyces israelii		99.81	0.63	NAG 6		
Actinomyces naeslundii		0.20	0.25			
43777030	**Good identification**					**43777030**
Actinomyces israelii		100.0	0.45	MLZ 13		
43777032	**Good identification**					**43777032**
Actinomyces israelii		97.30	0.49	MLZ 13		
Bifidobacterium dentium		2.71	-.03	NAG 6		
43777040	**Very good identification**					**43777040**
Actinomyces israelii		100.0	0.73			
43777041	**Good identification**					**43777041**
Actinomyces israelii		100.0	0.39	URE 1		
43777042	**Very good identification**					**43777042**
Actinomyces israelii		99.93	0.76			
Bifidobacterium adolescentis		0.07	0.16			
43777043	**Good identification**					**43777043**
Actinomyces israelii		100.0	0.42	URE 1		
43777050	**Very good identification**					**43777050**
					COC	
Actinomyces israelii		99.62	0.59	MLZ 13	1	
Blautia producta		0.38	-.03		99	
43777052	**Very good identification**					**43777052**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		99.41	0.62	MLZ 13		
Bifidobacterium adolescentis		0.59	0.18			
43777060	**Good identification**					**43777060**
Actinomyces israelii		100.0	0.53	NAG 6		
43777062	**Good identification**					**43777062**
Actinomyces israelii		100.0	0.56	NAG 6		
43777070	**Good identification**					**43777070**
Actinomyces israelii		100.0	0.38	MLZ 13		
43777072	**Good identification**					**43777072**
Actinomyces israelii		100.0	0.41	MLZ 13		
43777100	**Very good identification**					**43777100**
					SPO	
Actinomyces israelii		99.12	0.90		1	
Clostridium butyricum		0.77	0.33		99	
Bifidobacterium adolescentis		0.09	0.32		1	
43777101	**Very good identification**					**43777101**
					SPO	
Actinomyces israelii		97.32	0.56	URE 1	1	
Actinomyces naeslundii		1.93	0.36		1	
Clostridium butyricum		0.76	-.01		99	
43777102	**Good identification**					**43777102**
					SPO	
Actinomyces israelii		94.39	0.93		1	
Bifidobacterium adolescentis		5.36	0.66		1	
Clostridium butyricum		0.19	0.26		99	
43777103	**Good identification**					**43777103**
Actinomyces israelii		93.46	0.59	URE 1		
Bifidobacterium adolescentis		5.31	0.32			
Actinomyces naeslundii		1.23	0.36			
43777110	**Very good identification**					**43777110**
Actinomyces israelii		98.45	0.76	MLZ 13		
Bifidobacterium adolescentis		0.69	0.33			
Bifidobacterium dentium		0.36	0.10			
43777111	**Good identification**					**43777111**
Actinomyces israelii		100.0	0.42	MLZ 13		
43777112	**Low discrimination**					**43777112**
Actinomyces israelii		66.87	0.79	MLZ 13		
Bifidobacterium adolescentis		31.08	0.67			
Bifidobacterium dentium		1.65	0.27			
43777113	**Low discrimination**					**43777113**
Actinomyces israelii		67.14	0.45	MLZ 13		
Bifidobacterium adolescentis		31.20	0.33	URE 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		1.66	-.07				
43777120	**Very good identification**						**43777120**
					SPO		
Actinomyces israelii		97.02	0.70	NAG 6	1		
Clostridium butyricum		1.03	0.15		99		
Clostridium ramosum		0.93	0.19		92		
43777121	**Low discrimination**						**43777121**
Actinomyces israelii		76.33	0.36	NAG 6			
Actinomyces naeslundii		23.67	0.36	URE 1			
43777122	**Good identification**						**43777122**
					SPO		
Actinomyces israelii		94.10	0.73	NAG 6	1		
Clostridium ramosum		2.42	0.30		92		
Clostridium tertium		1.43	0.27		99		
43777123	**Low discrimination**						**43777123**
Actinomyces israelii		82.87	0.39	NAG 6			
Actinomyces naeslundii		17.13	0.36	URE 1			
43777130	**Low discrimination**						**43777130**
					SPO		
Actinomyces israelii		83.23	0.56	MLZ 13	1		
Bifidobacterium dentium		8.50	0.15	NAG 6	1		
Clostridium tertium		7.52	0.24		99		
43777132	**Low discrimination**						**43777132**
					SPO		
Actinomyces israelii		50.33	0.59	MLZ 13	1		
Bifidobacterium dentium		34.64	0.32	NAG 6	1		
Clostridium tertium		9.09	0.32		99		
43777140	**Very good identification**						**43777140**
					COC	SPO	
Actinomyces israelii		99.34	0.83		1	1	
Blautia producta		0.47	0.24		99	1	
Clostridium butyricum		0.17	0.15		1	99	
43777141	**Good identification**						**43777141**
Actinomyces israelii		99.11	0.49	URE 1			
Actinomyces naeslundii		0.89	0.23				
43777142	**Very good identification**						**43777142**
Actinomyces israelii		98.10	0.86				
Bifidobacterium adolescentis		1.77	0.50				
Bifidobacterium breve		0.05	0.24				
43777143	**Good identification**						**43777143**
Actinomyces israelii		97.65	0.52	URE 1			
Bifidobacterium adolescentis		1.76	0.16				
Actinomyces naeslundii		0.59	0.23				
43777150	**Good identification**						**43777150**
					COC		
Actinomyces israelii		91.17	0.69	MLZ 13	1		
Blautia producta		8.57	0.32		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		0.20	0.18		1		
43777151	**Acceptable identification**						**43777151**
					COC		
Actinomyces israelii		91.41	0.35	MLZ 13	1		
Blautia producta		8.59	-.03	URE 1	99		
43777152	**Low discrimination**						**43777152**
					COC		
Actinomyces israelii		86.31	0.72	MLZ 13	1		
Bifidobacterium adolescentis		12.75	0.52		1		
Blautia producta		0.60	0.15		99		
43777153	**Low discrimination**						**43777153**
Actinomyces israelii		87.13	0.38	MLZ 13			
Bifidobacterium adolescentis		12.87	0.18	URE 1			
43777160	**Very good identification**						**43777160**
					COC	SPO	
Actinomyces israelii		98.76	0.63	NAG 6	1	1	
Blautia producta		0.81	0.07		99	1	
Clostridium butyricum		0.23	-.03		1	99	
43777162	**Very good identification**						**43777162**
Actinomyces israelii		98.37	0.66	NAG 6			
Bifidobacterium breve		1.35	0.29				
Bifidobacterium adolescentis		0.28	0.16				
43777170	**Low discrimination**						**43777170**
					COC		
Actinomyces israelii		84.95	0.49	MLZ 13	1		
Blautia producta		13.89	0.15	NAG 6	99		
Bifidobacterium breve		1.16	0.12		1		
43777172	**Low discrimination**						**43777172**
Actinomyces israelii		87.30	0.52	MLZ 13			
Bifidobacterium breve		8.04	0.29	NAG 6			
Bifidobacterium adolescentis		2.04	0.18				
43777200	**Very good identification**						**43777200**
Actinomyces israelii		100.0	0.61	RHA 7			
43777202	**Very good identification**						**43777202**
Actinomyces israelii		100.0	0.64	RHA 7			
43777210	**Good identification**						**43777210**
Actinomyces israelii		100.0	0.47	RHA 7			
43777212	**Good identification**						**43777212**
Actinomyces israelii		100.0	0.50	RHA 7			
43777220	**Good identification**						**43777220**
Actinomyces israelii		100.0	0.40	RHA 7			
43777222	**Good identification**						**43777222**
					SPO		
Actinomyces israelii		98.21	0.43	RHA 7	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		1.79	-.03	NAG 6	92		
43777240	**Good identification**						**43777240**
Actinomyces israelii		100.0	0.54	RHA 7			
43777242	**Very good identification**						**43777242**
Actinomyces israelii		100.0	0.57	RHA 7			
43777250	**Good identification**						**43777250**
					COC		
Actinomyces israelii		95.20	0.40	RHA 7	1		
Blautia producta		4.80	-.03	MLZ 13	99		
43777252	**Good identification**						**43777252**
Actinomyces israelii		100.0	0.43	RHA 7			
43777262	**Good identification**						**43777262**
Actinomyces israelii		100.0	0.36	RHA 7			
43777300	**Very good identification**						**43777300**
					SPO		
Actinomyces israelii		99.68	0.71	RHA 7	1		
Clostridium butyricum		0.32	0.07		99		
43777301	**Good identification**						**43777301**
Actinomyces israelii		100.0	0.37	RHA 7			
43777302	**Very good identification**						**43777302**
					SPO		
Actinomyces israelii		99.13	0.74	RHA 7	1		
Bifidobacterium adolescentis		0.76	0.32		1		
Clostridium butyricum		0.08	0.00		99		
43777303	**Good identification**						**43777303**
Actinomyces israelii		100.0	0.40	RHA 7			
43777310	**Very good identification**						**43777310**
					COC		
Actinomyces israelii		99.51	0.57	RHA 7	1		
Blautia producta		0.49	-.03	MLZ 13	99		
43777312	**Good identification**						**43777312**
Actinomyces israelii		93.40	0.60	RHA 7			
Bifidobacterium adolescentis		5.83	0.33	MLZ 13			
Clostridium clostridioforme		0.47	0.11				
43777320	**Low discrimination**						**43777320**
					SPO		
Actinomyces israelii		85.51	0.51	RHA 7	1		
Clostridium ramosum		14.49	0.21	NAG 6	92		
43777322	**Low discrimination**						**43777322**
					SPO	GRA	
Actinomyces israelii		63.72	0.54	RHA 7	1	99	
Clostridium ramosum		28.80	0.32	NAG 6	92	50	
Parabacteroides distasonis		5.99	0.32		1	1	
43777330	**Good identification**						**43777330**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.37	RHA 7		
43777332	**Low discrimination**					**43777332**
					GRA	
Actinomyces israelii		60.81	0.40	RHA 7	99	
Parabacteroides distasonis		20.59	0.28	MLZ 13	1	
Bacteroides ovatus		6.32	0.03	NAG 6	1	
43777340	**Good identification**					**43777340**
					COC	
Actinomyces israelii		94.15	0.64	RHA 7	1	
Blautia producta		5.85	0.24		99	
43777342	**Very good identification**					**43777342**
					COC	
Actinomyces israelii		99.30	0.67	RHA 7	1	
Blautia producta		0.46	0.07		99	
Bifidobacterium adolescentis		0.24	0.16		1	
43777352	**Low discrimination**					**43777352**
					COC	
Actinomyces israelii		89.91	0.53	RHA 7	1	
Blautia producta		8.31	0.15	MLZ 13	99	
Bifidobacterium adolescentis		1.78	0.18		1	
43777360	**Acceptable identification**					**43777360**
					COC	
Actinomyces israelii		90.23	0.44	RHA 7	1	
Blautia producta		9.77	0.07	NAG 6	99	
43777362	**Good identification**					**43777362**
					SPO	
Actinomyces israelii		98.84	0.47	RHA 7	1	
Clostridium ramosum		1.16	-.03	NAG 6	92	
43777400	**Very good identification**					**43777400**
					SPO	
Actinomyces israelii		98.57	0.79		1	
Clostridium butyricum		1.40	0.26		99	
Bifidobacterium adolescentis		0.03	0.12		1	
43777401	**Good identification**					**43777401**
Actinomyces israelii		100.0	0.44	URE 1		
43777402	**Very good identification**					**43777402**
					SPO	
Actinomyces israelii		97.79	0.82		1	
Bifidobacterium adolescentis		1.84	0.46		1	
Clostridium butyricum		0.36	0.19		99	
43777403	**Good identification**					**43777403**
Actinomyces israelii		98.16	0.47	URE 1		
Bifidobacterium adolescentis		1.84	0.12			
43777410	**Very good identification**					**43777410**
					SPO	
Actinomyces israelii		97.37	0.64	MLZ 13	1	
Bifidobacterium dentium		1.75	0.10		1	
Clostridium butyricum		0.49	0.04		99	
43777412	**Low discrimination**					**43777412**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		78.32	0.67	MLZ 13			
Bifidobacterium adolescentis		12.03	0.47				
Bifidobacterium dentium		9.46	0.27				
43777420	**Very good identification**						**43777420**
					SPO		
Actinomyces israelii		98.10	0.58	NAG 6	1		
Clostridium butyricum		1.90	0.08		99		
43777422	**Very good identification**						**43777422**
					SPO		
Actinomyces israelii		98.71	0.61	NAG 6	1		
Bifidobacterium dentium		0.50	-.03		1		
Clostridium butyricum		0.50	0.01		99		
43777430	**Low discrimination**						**43777430**
Actinomyces israelii		66.71	0.44	MLZ 13			
Bifidobacterium dentium		33.30	0.15	NAG 6			
43777440	**Very good identification**						**43777440**
					COC	SPO	
Actinomyces israelii		97.46	0.72		1	1	
Blautia producta		2.23	0.24		99	1	
Clostridium butyricum		0.31	0.08		1	99	
43777441	**Good identification**						**43777441**
Actinomyces israelii		100.0	0.37	URE 1			
43777442	**Very good identification**						**43777442**
					COC		
Actinomyces israelii		99.16	0.75		1		
Bifidobacterium adolescentis		0.59	0.30		1		
Blautia producta		0.17	0.07		99		
43777443	**Good identification**						**43777443**
Actinomyces israelii		100.0	0.40	URE 1			
43777450	**Low discrimination**						**43777450**
					COC		
Actinomyces israelii		68.52	0.57	MLZ 13	1		
Blautia producta		31.48	0.32		99		
43777452	**Good identification**						**43777452**
					COC		
Actinomyces israelii		92.08	0.60	MLZ 13	1		
Bifidobacterium adolescentis		4.50	0.32		1		
Blautia producta		3.13	0.15		99		
43777460	**Good identification**						**43777460**
					COC		
Actinomyces israelii		96.17	0.51	NAG 6	1		
Blautia producta		3.83	0.07		99		
43777462	**Good identification**						**43777462**
Actinomyces israelii		100.0	0.54	NAG 6			
43777470	**Low discrimination**						**43777470**
					COC		
Actinomyces israelii		55.57	0.37	MLZ 13	1		
Blautia producta		44.43	0.15	NAG 6	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
43777472	**Low discrimination**						**43777472**
					COC		
Actinomyces israelii		87.21	0.40	MLZ 13	1		
Bifidobacterium dentium		7.62	-.03	NAG 6	1		
Blautia producta		5.17	-.01		99		
43777500	**Good identification**						**43777500**
					SPO		
Actinomyces israelii		91.18	0.89		1		
Clostridium butyricum		7.79	0.50		99		
Bifidobacterium adolescentis		0.64	0.46		1		
43777501	**Acceptable identification**						**43777501**
					GRA	SPO	
Actinomyces israelii		90.31	0.55	URE 1	99	1	
Clostridium butyricum		7.71	0.15		99	99	
Mitsuokella multacida		1.34	0.11		1	1	
43777502	**Low discrimination**						**43777502**
					SPO		
Actinomyces israelii		67.04	0.92		1		
Bifidobacterium adolescentis		31.16	0.80		1		
Clostridium butyricum		1.49	0.43		99		
43777503	**Low discrimination**						**43777503**
					SPO		
Actinomyces israelii		67.05	0.58	URE 1	1		
Bifidobacterium adolescentis		31.16	0.46		1		
Clostridium butyricum		1.49	0.08		99		
43777510	**Low discrimination**						**43777510**
Actinomyces israelii		63.34	0.75	MLZ 13			
Bifidobacterium dentium		28.09	0.44				
Bifidobacterium adolescentis		3.65	0.47				
43777511	**Low discrimination**						**43777511**
Actinomyces israelii		62.15	0.41	MLZ 13			
Bifidobacterium dentium		27.57	0.10	URE 1			
Bifidobacterium adolescentis		3.58	0.13				
43777512	**Low discrimination**						**43777512**
Bifidobacterium adolescentis		48.71	0.81	MNS 87			
Bifidobacterium dentium		38.31	0.62				
Actinomyces israelii		12.82	0.78				
43777513	**Low discrimination**						**43777513**
Bifidobacterium adolescentis		48.79	0.47	MNS 87			
Bifidobacterium dentium		38.37	0.27	URE 1			
Actinomyces israelii		12.84	0.44				
43777520	**Low discrimination**						**43777520**
					SPO		
Actinomyces israelii		87.32	0.68	NAG 6	1		
Clostridium butyricum		10.16	0.31		99		
Bifidobacterium dentium		1.63	0.15		1		
43777522	**Low discrimination**						**43777522**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		80.95	0.71	NAG 6			
Bifidobacterium dentium		10.17	0.32				
Bifidobacterium adolescentis		5.95	0.46				
43777523	**Low discrimination**						**43777523**
Actinomyces israelii		83.39	0.37	NAG 6			
Bifidobacterium dentium		10.48	-.03	URE 1			
Bifidobacterium adolescentis		6.13	0.12				
43777530	**Acceptable identification**						**43777530**
					COC		
Bifidobacterium dentium		91.55	0.49	MNS 99	1		
Actinomyces israelii		7.41	0.54	BGL 91	1		
Blautia producta		0.58	0.15		99		
43777532	**Very good identification**						**43777532**
Bifidobacterium dentium		98.05	0.66	MNS 99			
Actinomyces israelii		1.18	0.57				
Bifidobacterium adolescentis		0.71	0.47				
43777540	**Low discrimination**						**43777540**
					COC	SPO	
Actinomyces israelii		62.85	0.82		1	1	
Blautia producta		35.59	0.58		99	1	
Clostridium butyricum		1.20	0.31		1	99	
43777541	**Low discrimination**						**43777541**
					COC	GRA	
Actinomyces israelii		61.60	0.48	URE 1	1	99	
Blautia producta		34.88	0.24		99	99	
Mitsuokella multacida		2.35	0.11		1	1	
43777542	**Low discrimination**						**43777542**
					COC		
Actinomyces israelii		83.68	0.85		1		
Bifidobacterium adolescentis		12.36	0.64		1		
Blautia producta		3.51	0.41		99		
43777543	**Low discrimination**						**43777543**
					COC		
Actinomyces israelii		84.06	0.51	URE 1	1		
Bifidobacterium adolescentis		12.42	0.30		1		
Blautia producta		3.53	0.07		99		
43777550	**Good identification**						**43777550**
					COC		
Blautia producta		91.59	0.66	MNS 99	99		
Actinomyces israelii		8.06	0.68		1		
Bifidobacterium adolescentis		0.15	0.32		1		
43777552	**Low discrimination**						**43777552**
					COC		
Bifidobacterium adolescentis		38.60	0.66	MNS 87	1		
Actinomyces israelii		31.95	0.71	SOR 11	1		
Blautia producta		26.90	0.50		99		
43777560	**Low discrimination**						**43777560**
					COC	SPO	
Actinomyces israelii		49.38	0.61	NAG 6	1	1	
Blautia producta		48.67	0.41		99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		1.29	0.13		1	99	
43777562	**Good identification**						**43777562**
					COC		
Actinomyces israelii		90.38	0.64	NAG 6	1		
Blautia producta		6.60	0.25		99		
Bifidobacterium adolescentis		2.11	0.30		1		
43777570	**Acceptable identification**						**43777570**
					COC		
Blautia producta		93.61	0.50	MNS 99	99		
Actinomyces israelii		4.73	0.47	NAG 10	1		
Bifidobacterium dentium		1.52	0.15		1		
43777600	**Very good identification**						**43777600**
					SPO		
Actinomyces israelii		99.42	0.59	RHA 7	1		
Clostridium butyricum		0.58	0.00		99		
43777602	**Very good identification**						**43777602**
Actinomyces israelii		99.22	0.62	RHA 7			
Clostridium clostridioforme		0.38	0.11				
Bifidobacterium adolescentis		0.25	0.12				
43777610	**Good identification**						**43777610**
					COC		
Actinomyces israelii		97.66	0.45	RHA 7	1		
Blautia producta		2.34	-.03	MLZ 13	99		
43777612	**Low discrimination**						**43777612**
Actinomyces israelii		86.80	0.48	RHA 7			
Clostridium clostridioforme		10.01	0.23	MLZ 13			
Bifidobacterium adolescentis		1.79	0.13				
43777620	**Good identification**						**43777620**
Actinomyces israelii		100.0	0.39	RHA 7			
43777622	**Acceptable identification**						**43777622**
					GRA		
Actinomyces israelii		91.20	0.42	RHA 7	99		
Clostridium clostridioforme		5.40	0.11	NAG 6	50		
Bacteroides ovatus		3.39	-.02		1		
43777640	**Low discrimination**						**43777640**
					COC		
Actinomyces israelii		76.69	0.52	RHA 7	1		
Blautia producta		23.31	0.24		99		
43777642	**Good identification**						**43777642**
					COC		
Actinomyces israelii		97.80	0.55	RHA 7	1		
Blautia producta		2.20	0.07		99		
43777652	**Low discrimination**						**43777652**
					COC		
Actinomyces israelii		68.87	0.41	RHA 7	1		
Blautia producta		31.13	0.15	MLZ 13	99		
43777700	**Good identification**						**43777700**
					GRA	SPO	
Actinomyces israelii		91.00	0.70	RHA 7	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		3.19	0.24		99	99	
Mitsuokella multacida		3.03	0.32		1	1	
43777701	**Low discrimination**						**43777701**
					GRA		
Actinomyces israelii		73.21	0.35	RHA 7	99		
Mitsuokella multacida		26.79	0.15	URE 1	1		
43777702	**Good identification**						**43777702**
					SPO		
Actinomyces israelii		92.55	0.73	RHA 7	1		
Bifidobacterium adolescentis		5.77	0.46		1		
Clostridium butyricum		0.84	0.17		99		
43777703	**Acceptable identification**						**43777703**
Actinomyces israelii		94.13	0.39	RHA 7			
Bifidobacterium adolescentis		5.87	0.12	URE 1			
43777710	**Low discrimination**						**43777710**
					COC	GRA	
Actinomyces israelii		57.24	0.56	RHA 7	1	99	
Blautia producta		33.96	0.32	MLZ 13	99	99	
Mitsuokella multacida		4.25	0.24		1	1	
43777712	**Low discrimination**						**43777712**
Actinomyces israelii		48.47	0.59	RHA 7			
Bifidobacterium adolescentis		24.73	0.47	MLZ 13			
Bifidobacterium dentium		19.45	0.27				
43777720	**Low discrimination**						**43777720**
					COC	GRA	
Actinomyces israelii		80.42	0.49	RHA 7	1	99	
Mitsuokella multacida		10.48	0.21	NAG 6	1	1	
Blautia producta		4.14	0.07		99	99	
43777722	**Low discrimination**						**43777722**
					GRA		
Actinomyces israelii		47.73	0.52	RHA 7	99		
Bacteroides ovatus		43.95	0.32	NAG 6	1		
Parabacteroides distasonis		3.36	0.29		1		
43777740	**Low discrimination**						**43777740**
					COC	GRA	
Blautia producta		87.31	0.58	MNS 99	99	99	
Actinomyces israelii		11.61	0.63		1	99	
Mitsuokella multacida		0.99	0.32		1	1	
43777742	**Low discrimination**						**43777742**
					COC		
Actinomyces israelii		63.07	0.66	RHA 7	1		
Blautia producta		35.15	0.41		99		
Bifidobacterium adolescentis		1.25	0.30		1		
43777750	**Very good identification**						**43777750**
					COC	GRA	
Blautia producta		99.22	0.66	MNS 99	99	99	
Actinomyces israelii		0.66	0.49		1	99	
Mitsuokella multacida		0.13	0.24		1	1	
43777752	**Acceptable identification**						**43777752**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		90.41	0.50	MNS 99	99		
Actinomyces israelii		8.08	0.52	BGL 10	1		
Bifidobacterium adolescentis		1.31	0.32		1		
43777760	**Acceptable identification**						**43777760**
					COC	GRA	
Blautia producta		90.75	0.41	MNS 99	99	99	
Actinomyces israelii		6.93	0.42	NAG 10	1	99	
Mitsuokella multacida		2.32	0.21		1	1	
43777762	**Low discrimination**						**43777762**
					COC	GRA	
Actinomyces israelii		47.77	0.45	RHA 7	1	99	
Blautia producta		46.34	0.25	NAG 6	99	99	
Bacteroides ovatus		4.71	0.08		1	1	
43777770	**Good identification**						**43777770**
					COC	GRA	
Blautia producta		99.34	0.50	MNS 99	99	99	
Actinomyces israelii		0.38	0.28	NAG 10	1	99	
Mitsuokella multacida		0.28	0.13		1	1	
44000000	**Low discrimination**						**44000000**
					COC	SPO	
Atopobium minutum		48.28	0.80		1	1	
Clostridium botulinum biovar A		15.98	0.64		1	99	
Parvimonas micra		10.15	0.50		99	1	
44000001	**Very good identification**						**44000001**
					COC	GRA	
Anaerococcus prevotii		95.77	0.66	ESL 1	99	99	
Campylobacter ureolyticus		1.59	0.32		1	1	
Atopobium minutum		1.09	0.46		1	99	
44000002	**Low discrimination**						**44000002**
					COC		
Atopobium minutum		87.82	0.77		1		
Anaerococcus prevotii		3.25	0.39		99		
Parvimonas micra		3.08	0.33		99		
44000003	**Good identification**						**44000003**
					COC		
Anaerococcus prevotii		96.41	0.56	ESL 1	99		
Atopobium minutum		2.93	0.43		1		
Peptococcus niger		0.38	0.10		99		
44000010	**Low discrimination**						**44000010**
					COC	SPO	
Atopobium minutum		46.65	0.46	MLZ 1	1	1	
Clostridium botulinum biovar A		15.44	0.30		1	99	
Parvimonas micra		9.81	0.15		99	1	
44000012	**Low discrimination**						**44000012**
					COC		
Atopobium minutum		88.84	0.43	MLZ 1	1		
Anaerococcus prevotii		3.28	0.05		99		
Parvimonas micra		3.11	-.01		99		
44000020	**Good identification**						**44000020**
					GRA	SPO	
Atopobium minutum		90.39	0.72		99	1	
Clostridium bifermentans		3.06	0.39		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		2.62	0.32		1	1	
44000022	**Low discrimination**						**44000022**
					COC	GRA	
Pseudoflavonifractor capillosus		81.01	0.66	MNS 1	1	1	
Atopobium minutum		18.84	0.69		1	99	
Peptococcus niger		0.08	0.11		99	99	
44000030	**Acceptable identification**						**44000030**
					GRA	SPO	
Atopobium minutum		92.64	0.38	MLZ 1	99	1	
Clostridium bifermentans		3.14	0.05		99	99	
Pseudoflavonifractor capillosus		2.68	-.03		1	1	
44000040	**Low discrimination**						**44000040**
					SPO		
Clostridium botulinum biovar A		81.00	0.56	GLU 90	99		
Atopobium minutum		8.28	0.46	MNS 7	1		
Clostridium bifermentans		6.04	0.36		99		
44000042	**Low discrimination**						**44000042**
					COC	SPO	
Atopobium minutum		77.82	0.43	SOR 1	1	1	
Clostridium botulinum biovar A		11.54	0.21		1	99	
Anaerococcus prevotii		2.88	0.05		99	1	
44000060	**Low discrimination**						**44000060**
					SPO		
Atopobium minutum		67.48	0.38	SOR 1	1		
Clostridium botulinum biovar A		20.02	0.21		99		
Clostridium bifermentans		9.43	0.15		99		
44000100	**Low discrimination**						**44000100**
					COC	SPO	
Atopobium minutum		48.42	0.46	LAC 1	1	1	
Clostridium botulinum biovar A		16.03	0.30		1	99	
Parvimonas micra		10.18	0.15		99	1	
44000102	**Low discrimination**						**44000102**
					COC		
Atopobium minutum		88.84	0.43	LAC 1	1		
Anaerococcus prevotii		3.28	0.05		99		
Parvimonas micra		3.11	-.01		99		
44000120	**Low discrimination**						**44000120**
					GRA	SPO	
Atopobium minutum		88.22	0.38	LAC 1	99	1	
Prevotella bivia		4.78	-.03		1	1	
Clostridium bifermentans		2.99	0.05		99	99	
44000200	**Low discrimination**						**44000200**
					COC	SPO	
Atopobium minutum		48.42	0.46	RHA 1	1	1	
Clostridium botulinum biovar A		16.03	0.30		1	99	
Parvimonas micra		10.18	0.15		99	1	
44000202	**Low discrimination**						**44000202**
					COC		
Atopobium minutum		88.84	0.43	RHA 1	1		
Anaerococcus prevotii		3.28	0.05		99		
Parvimonas micra		3.11	-.01		99		
44000220	**Acceptable identification**						**44000220**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Atopobium minutum		92.64	0.38	RHA 1	99	1	
Clostridium bifermentans		3.14	0.05		99	99	
Pseudoflavonifractor capillosus		2.68	-.03		1	1	
44000400	**Low discrimination**						**44000400**
					COC	SPO	
Atopobium minutum		48.32	0.46	ARA 1	1	1	
Clostridium botulinum biovar A		15.99	0.30		1	99	
Parvimonas micra		10.16	0.15		99	1	
44000402	**Low discrimination**						**44000402**
					COC		
Atopobium minutum		88.84	0.43	ARA 1	1		
Anaerococcus prevotii		3.28	0.05		99		
Parvimonas micra		3.11	-.01		99		
44000420	**Acceptable identification**						**44000420**
					GRA	SPO	
Atopobium minutum		92.64	0.38	ARA 1	99	1	
Clostridium bifermentans		3.14	0.05		99	99	
Pseudoflavonifractor capillosus		2.68	-.03		1	1	
44001000	**Very good identification**						**44001000**
					COC	SPO	
Atopobium minutum		96.78	0.72		1	1	
Clostridium botulinum biovar A		0.97	0.30		1	99	
Parvimonas micra		0.62	0.15		99	1	
44001002	**Very good identification**						**44001002**
					COC	GRA	
Atopobium minutum		99.19	0.69		1	99	
Pseudoflavonifractor capillosus		0.48	0.15		1	1	
Anaerococcus prevotii		0.11	0.05		99	99	
44001003	**Low discrimination**						**44001003**
					COC		
Atopobium minutum		50.05	0.35	URE 1	1		
Anaerococcus prevotii		49.95	0.21		99		
44001010	**Good identification**						**44001010**
Atopobium minutum		100.0	0.38	MLZ 1			
44001012	**Good identification**						**44001012**
Atopobium minutum		100.0	0.35	MLZ 1			
44001020	**Very good identification**						**44001020**
					GRA	SPO	
Atopobium minutum		98.89	0.64		99	1	
Pseudoflavonifractor capillosus		0.96	0.15		1	1	
Clostridium bifermentans		0.10	0.05		99	99	
44001022	**Low discrimination**						**44001022**
					GRA		
Pseudoflavonifractor capillosus		58.90	0.50	MNS 1	1		
Atopobium minutum		41.10	0.61	GAL 10	99		
44001040	**Low discrimination**						**44001040**
					SPO		
Atopobium minutum		75.83	0.38	SOR 1	1		
Clostridium botulinum biovar A		22.49	0.21		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		1.68	0.01		99		
44001042	**Good identification**						**44001042**
Atopobium minutum		100.0	0.35	SOR 1			
44001100	**Good identification**						**44001100**
Atopobium minutum		100.0	0.38	LAC 1			
44001102	**Good identification**						**44001102**
Atopobium minutum		100.0	0.35	LAC 1			
44001200	**Good identification**						**44001200**
Atopobium minutum		100.0	0.38	RHA 1			
44001202	**Good identification**						**44001202**
Atopobium minutum		100.0	0.35	RHA 1			
44001400	**Good identification**						**44001400**
Atopobium minutum		100.0	0.38	ARA 1			
44001402	**Good identification**						**44001402**
Atopobium minutum		100.0	0.35	ARA 1			
44002000	**Low discrimination**						**44002000**
					SPO		
Atopobium minutum		34.32	0.46	MAN 1	1		
Eubacterium limosum		25.07	0.32		1		
Clostridium botulinum biovar A		11.36	0.30		99		
44002002	**Low discrimination**						**44002002**
					COC		
Atopobium minutum		87.98	0.43	MAN 1	1		
Anaerococcus prevotii		3.25	0.05		99		
Parvimonas micra		3.08	-.01		99		
44002020	**Acceptable identification**						**44002020**
					GRA	SPO	
Atopobium minutum		90.78	0.38	MAN 1	99	1	
Clostridium bifermentans		3.08	0.05		99	99	
Pseudoflavonifractor capillosus		2.63	-.03		1	1	
44003000	**Good identification**						**44003000**
Atopobium minutum		97.83	0.38	MAN 1			
Eubacterium limosum		2.17	-.03				
44003002	**Good identification**						**44003002**
Atopobium minutum		100.0	0.35	MAN 1			
44004000	**Low discrimination**						**44004000**
					COC	SPO	
Atopobium minutum		48.22	0.46	XYL 1	1	1	
Clostridium botulinum biovar A		15.96	0.30		1	99	
Parvimonas micra		10.14	0.15		99	1	
44004002	**Low discrimination**						**44004002**
					COC		
Atopobium minutum		88.84	0.43	XYL 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Anaerococcus prevotii		3.28	0.05		99		
Parvimonas micra		3.11	-.01		99		
44004020	**Acceptable identification**						**44004020**
					GRA	SPO	
Atopobium minutum		92.64	0.38	XYL 1	99	1	
Clostridium bifermentans		3.14	0.05		99	99	
Pseudoflavonifractor capillosus		2.68	-.03		1	1	
44005000	**Good identification**						**44005000**
Atopobium minutum		100.0	0.38	XYL 1			
44005002	**Good identification**						**44005002**
Atopobium minutum		100.0	0.35	XYL 1			
44010000	**Low discrimination**						**44010000**
					GRA		
Atopobium minutum		76.67	0.80		99		
Eubacterium limosum		9.88	0.52		99		
Fusobacterium varium		8.50	0.50		1		
44010001	**Low discrimination**						**44010001**
					COC		
Atopobium minutum		43.88	0.46	URE 1	1		
Anaerococcus prevotii		42.17	0.32		99		
Eubacterium limosum		5.65	0.18		1		
44010002	**Very good identification**						**44010002**
					SPO		
Atopobium minutum		98.80	0.77		1		
Clostridium sporogenes		0.49	0.27		99		
Eubacterium limosum		0.19	0.18		1		
44010003	**Low discrimination**						**44010003**
					COC		
Atopobium minutum		73.51	0.43	URE 1	1		
Anaerococcus prevotii		26.49	0.21		99		
44010010	**Low discrimination**						**44010010**
					SPO		
Clostridium difficile		84.26	0.50	GLU 99	99		
Atopobium minutum		12.27	0.46	MAN 90	1		
Eubacterium limosum		1.58	0.18		1		
44010012	**Acceptable identification**						**44010012**
					SPO		
Atopobium minutum		90.58	0.43	MLZ 1	1		
Clostridium difficile		9.43	0.15		99		
44010020	**Very good identification**						**44010020**
					GRA		
Atopobium minutum		98.43	0.72		99		
Eubacterium limosum		0.38	0.18		99		
Pseudoflavonifractor capillosus		0.34	0.15		1		
44010021	**Good identification**						**44010021**
					COC		
Atopobium minutum		97.17	0.37	URE 1	1		
Anaerococcus prevotii		2.83	-.03		99		
44010022	**Low discrimination**						**44010022**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Atopobium minutum		65.89	0.69		99	
Pseudoflavonifractor capillosus		34.11	0.50		1	
44010030	**Low discrimination**					**44010030**
					SPO	
Atopobium minutum		82.77	0.37	MLZ 1	1	
Clostridium difficile		17.23	0.15		99	
44010040	**Low discrimination**					**44010040**
					SPO	
Atopobium minutum		48.40	0.46	SOR 1	1	
Clostridium botulinum biovar A		32.76	0.35		99	
Eubacterium limosum		6.23	0.18		1	
44010042	**Good identification**					**44010042**
					SPO	
Atopobium minutum		98.99	0.43	SOR 1	1	
Clostridium botulinum biovar A		1.02	0.01		99	
44010060	**Good identification**					**44010060**
					SPO	
Atopobium minutum		97.99	0.37	SOR 1	1	
Clostridium botulinum biovar A		2.01	0.01		99	
44010100	**Low discrimination**					**44010100**
					GRA	
Atopobium minutum		65.54	0.46	LAC 1	99	
Fusobacterium mortiferum		15.02	0.40		1	
Eubacterium limosum		8.44	0.18		99	
44010102	**Good identification**					**44010102**
					GRA	
Atopobium minutum		96.32	0.43	LAC 1	99	
Fusobacterium mortiferum		3.68	0.24		1	
44010120	**Good identification**					**44010120**
Atopobium minutum		100.0	0.37	LAC 1		
44010200	**Low discrimination**					**44010200**
					GRA	
Atopobium minutum		77.12	0.46	RHA 1	99	
Eubacterium limosum		9.93	0.18		99	
Fusobacterium varium		8.55	0.15		1	
44010202	**Good identification**					**44010202**
Atopobium minutum		100.0	0.43	RHA 1		
44010220	**Good identification**					**44010220**
Atopobium minutum		100.0	0.37	RHA 1		
44010400	**Low discrimination**					**44010400**
					GRA	
Atopobium minutum		76.77	0.46	ARA 1	99	
Eubacterium limosum		9.89	0.18		99	
Fusobacterium varium		8.51	0.15		1	
44010402	**Good identification**					**44010402**
Atopobium minutum		100.0	0.43	ARA 1		
44010420	**Good identification**					**44010420**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		100.0	0.37	ARA 1			
44011000	**Very good identification**						**44011000**
					GRA		
Atopobium minutum		98.57	0.72		99		
Fusobacterium mortiferum		0.44	0.37		1		
Eubacterium limosum		0.39	0.18		99		
44011001	**Good identification**						**44011001**
					COC		
Atopobium minutum		97.17	0.37	URE 1	1		
Anaerococcus prevotii		2.83	-.03		99		
44011002	**Very good identification**						**44011002**
					GRA		
Atopobium minutum		99.87	0.69		99		
Fusobacterium mortiferum		0.07	0.20		1		
Pseudoflavonifractor capillosus		0.06	-.01		1		
44011010	**Low discrimination**						**44011010**
					SPO		
Atopobium minutum		82.77	0.37	MLZ 1	1		
Clostridium difficile		17.23	0.15		99		
44011020	**Very good identification**						**44011020**
					GRA	CAT	
Atopobium minutum		99.84	0.63		99	1	
Pseudoflavonifractor capillosus		0.12	-.01		1	1	
Propionibacterium acnes		0.05	-.01		99	95	
44011022	**Low discrimination**						**44011022**
					GRA		
Atopobium minutum		85.29	0.60		99		
Pseudoflavonifractor capillosus		14.71	0.33		1		
44011040	**Good identification**						**44011040**
					SPO		
Atopobium minutum		97.99	0.37	SOR 1	1		
Clostridium botulinum biovar A		2.01	0.01		99		
44011100	**Low discrimination**						**44011100**
					GRA		
Fusobacterium mortiferum		50.76	0.43	GLU 99	1		
Atopobium minutum		49.24	0.37		99		
44011200	**Good identification**						**44011200**
Atopobium minutum		100.0	0.37	RHA 1			
44011400	**Good identification**						**44011400**
Atopobium minutum		100.0	0.37	ARA 1			
44012000	**Low discrimination**						**44012000**
					SPO		
Eubacterium limosum		85.01	0.66	GLU 99	1		
Clostridium difficile		13.74	0.54		99		
Atopobium minutum		1.09	0.46		1		
44012010	**Very good identification**						**44012010**
					SPO		
Clostridium difficile		98.72	0.66	GLU 99	99		
Eubacterium limosum		1.26	0.32		1		
Atopobium minutum		0.02	0.11		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
44012050	**Good identification**					**44012050**
					SPO	
Clostridium difficile		99.76	0.44	GLU 99	99	
Eubacterium limosum		0.25	-.03	SOR 5	1	
44014000	**Low discrimination**					**44014000**
					GRA	
Atopobium minutum		57.46	0.46	XYL 1	99	
Fusobacterium varium		26.27	0.26		1	
Eubacterium limosum		7.40	0.18		99	
44014002	**Good identification**					**44014002**
					GRA	
Atopobium minutum		99.31	0.43	XYL 1	99	
Fusobacterium varium		0.69	-.08		1	
44014020	**Good identification**					**44014020**
					GRA	
Atopobium minutum		98.63	0.37	XYL 1	99	
Fusobacterium varium		1.37	-.08		1	
44015000	**Good identification**					**44015000**
					GRA	
Atopobium minutum		95.50	0.37	XYL 1	99	
Fusobacterium mortiferum		3.18	0.18		1	
Fusobacterium varium		1.32	-.08		1	
44016010	**Good identification**					**44016010**
					SPO	
Clostridium difficile		99.76	0.44	GLU 99	99	
Eubacterium limosum		0.25	-.03	XYL 5	1	
44020000	**Low discrimination**					**44020000**
					SPO	
Clostridium botulinum biovar A		70.70	0.54	GLU 90	99	
Clostridium sporogenes		14.78	0.40	MNS 7	99	
Atopobium minutum		8.63	0.46		1	
44020002	**Low discrimination**					**44020002**
					SPO	
Atopobium minutum		48.08	0.43	TRE 1	1	
Clostridium sporogenes		41.16	0.32		99	
Clostridium botulinum biovar A		5.97	0.20		99	
44020020	**Low discrimination**					**44020020**
					SPO	
Atopobium minutum		72.49	0.38	TRE 1	1	
Clostridium botulinum biovar A		18.00	0.20		99	
Clostridium sporogenes		3.76	0.06		99	
44020040	**Good identification**					**44020040**
					SPO	
Clostridium botulinum biovar A		98.60	0.45	GLU 90	99	
Clostridium sporogenes		0.70	0.06	MNS 7	99	
Atopobium minutum		0.41	0.12		1	
44021000	**Low discrimination**					**44021000**
					SPO	
Atopobium minutum		76.92	0.38	TRE 1	1	
Clostridium botulinum biovar A		19.09	0.20		99	
Clostridium sporogenes		3.99	0.06		99	
44021002	**Good identification**					**44021002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Atopobium minutum		97.47	0.35	TRE 1	1		
Clostridium sporogenes		2.53	-.02		99		
44030000	**Low discrimination**						**44030000**
					SPO		
Atopobium minutum		43.75	0.46	TRE 1	1		
Clostridium botulinum biovar A		24.79	0.34		99		
Clostridium sporogenes		14.32	0.27		99		
44030002	**Low discrimination**						**44030002**
					SPO		
Atopobium minutum		85.31	0.43	TRE 1	1		
Clostridium sporogenes		13.96	0.19		99		
Clostridium botulinum biovar A		0.73	-.01		99		
44030020	**Good identification**						**44030020**
					SPO		
Atopobium minutum		98.31	0.37	TRE 1	1		
Clostridium botulinum biovar A		1.69	-.01		99		
44031000	**Low discrimination**						**44031000**
					GRA	CAT	
Atopobium minutum		85.50	0.37	TRE 1	99	1	
Fusobacterium mortiferum		11.29	0.28		1	5	
Propionibacterium granulosum		1.75	0.11		99	90	
44032010	**Good identification**						**44032010**
					SPO		
Clostridium difficile		99.83	0.47	GLU 99	99		
Eubacterium limosum		0.17	-.03	TRE 7	1		
44040000	**Low discrimination**						**44040000**
					COC	SPO	
Atopobium minutum		48.42	0.46	CEL 1	1	1	
Clostridium botulinum biovar A		16.03	0.30		1	99	
Parvimonas micra		10.18	0.15		99	1	
44040002	**Low discrimination**						**44040002**
					COC		
Atopobium minutum		88.84	0.43	CEL 1	1		
Anaerococcus prevotii		3.28	0.05		99		
Parvimonas micra		3.11	-.01		99		
44040020	**Acceptable identification**						**44040020**
					GRA	SPO	
Atopobium minutum		92.64	0.38	CEL 1	99	1	
Clostridium bifermentans		3.14	0.05		99	99	
Pseudoflavonifractor capillosus		2.68	-.03		1	1	
44041000	**Good identification**						**44041000**
Atopobium minutum		100.0	0.38	CEL 1			
44041002	**Good identification**						**44041002**
Atopobium minutum		100.0	0.35	CEL 1			
44050000	**Low discrimination**						**44050000**
					GRA		
Atopobium minutum		71.98	0.46	CEL 1	99		
Eubacterium limosum		9.27	0.18		99		
Fusobacterium varium		7.98	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
44050002	**Good identification**					**44050002**
Atopobium minutum		100.0	0.43	CEL 1		
44050020	**Good identification**					**44050020**
Atopobium minutum		100.0	0.37	CEL 1		
44051000	**Low discrimination**					**44051000**
					GRA	
Atopobium minutum		87.16	0.37	CEL 1	99	
Fusobacterium mortiferum		12.84	0.29		1	
44052010	**Good identification**					**44052010**
					SPO	
Clostridium difficile		99.76	0.44	GLU 99	99	
Eubacterium limosum		0.25	-.03	CEL 5	1	
44071120	**Good identification**					**44071120**
Clostridium septicum		100.0	0.35	GLU 99		
44100000	**Low discrimination**					**44100000**
Clostridium bifermentans		50.14	0.66	IND 90		
Clostridium botulinum biovar A		32.90	0.63	GLU 92		
Clostridium sporogenes		11.71	0.53			
44100002	**Low discrimination**					**44100002**
					SPO	
Clostridium sporogenes		68.79	0.45	MNS 1	99	
Atopobium minutum		14.42	0.43		1	
Clostridium bifermentans		8.93	0.31		99	
44100020	**Low discrimination**					**44100020**
					SPO	
Clostridium bifermentans		76.88	0.45	IND 90	99	
Atopobium minutum		9.83	0.38	GLU 92	1	
Clostridium botulinum biovar A		7.98	0.29	NAG 6	99	
44100040	**Low discrimination**					**44100040**
Clostridium botulinum biovar A		81.24	0.54	GLU 90		
Clostridium bifermentans		17.27	0.42	MNS 7		
Clostridium sporogenes		0.98	0.19			
44101002	**Low discrimination**					**44101002**
					SPO	
Atopobium minutum		85.96	0.35	MLT 1	1	
Clostridium sporogenes		12.42	0.11		99	
Clostridium bifermentans		1.61	-.03		99	
44110000	**Low discrimination**					**44110000**
Clostridium bifermentans		25.70	0.44	IND 90		
Clostridium botulinum biovar A		20.46	0.43	GLU 92		
Clostridium sporogenes		20.12	0.40	FRU 5		
44110002	**Low discrimination**					**44110002**
					SPO	
Atopobium minutum		50.06	0.43	MLT 1	1	
Clostridium sporogenes		45.65	0.32		99	
Clostridium bifermentans		1.77	0.10		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
44110020	**Low discrimination**						**44110020**
					COC	SPO	
Atopobium minutum		58.81	0.37	MLT 1	1	1	
Clostridium bifermentans		26.23	0.23		1	99	
Gemella morbillorum		5.78	0.15		99	1	
44111000	**Low discrimination**						**44111000**
					COC	GRA	
Prevotella intermedia		47.95	0.35	ESL 1	1	1	
Atopobium minutum		29.89	0.37	GLU 90	1	99	
Gemella morbillorum		14.23	0.27		99	99	
44111120	**Low discrimination**						**44111120**
					GRA	SPO	
Clostridium septicum		64.80	0.39	GLU 99	99	99	
Prevotella melaninogenica		20.72	0.30		1	1	
Clostridium perfringens		13.26	0.34		99	84	
44120000	**Low discrimination**						**44120000**
Clostridium botulinum biovar A		64.88	0.53	GLU 90			
Clostridium sporogenes		30.78	0.45	MNS 7			
Clostridium bifermentans		4.00	0.31				
44120002	**Acceptable identification**						**44120002**
					SPO		
Clostridium sporogenes		93.10	0.37	MNS 1	99		
Clostridium botulinum biovar A		5.95	0.19		99		
Atopobium minutum		0.59	0.09		1		
44120040	**Good identification**						**44120040**
Clostridium botulinum biovar A		96.83	0.44	GLU 90			
Clostridium sporogenes		1.55	0.11	MNS 7			
Clostridium bifermentans		0.83	0.08				
44131120	**Low discrimination**						**44131120**
					COC	SPO	
Clostridium septicum		88.11	0.47	GLU 99	1	99	
Clostridium perfringens		11.16	0.38		1	84	
Atopobium parvulum		0.64	0.16		99	1	
44151120	**Good identification**						**44151120**
					GRA	SPO	
Clostridium septicum		96.97	0.47	GLU 99	99	99	
Clostridium paraputrificum		1.51	0.14		99	99	
Prevotella melaninogenica		1.09	0.14		1	1	
44171100	**Low discrimination**						**44171100**
					COC	SPO	
Atopobium parvulum		74.89	0.36	GLU 99	99	1	
Clostridium septicum		25.11	0.21	SUC 96	1	99	
44171120	**Good identification**						**44171120**
					COC	SPO	
Clostridium septicum		99.17	0.56	GLU 99	1	99	
Atopobium parvulum		0.53	0.23		99	1	
Clostridium perfringens		0.17	0.15		1	84	
44175100	**Good identification**						**44175100**
Atopobium parvulum		100.0	0.40	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
44200000	**Low discrimination**						**44200000**
					SPO		
Clostridium bifermentans		50.75	0.43	IND 90	99		
Atopobium minutum		26.08	0.46	GLU 92	1		
Clostridium botulinum biovar A		8.63	0.30	SAL 10	99		
44200002	**Low discrimination**						**44200002**
					COC		
Atopobium minutum		86.57	0.43	SAL 1	1		
Anaerococcus prevotii		3.20	0.05		99		
Parvimonas micra		3.03	-.01		99		
44200020	**Low discrimination**						**44200020**
					GRA	SPO	
Atopobium minutum		70.52	0.38	SAL 1	99	1	
Clostridium bifermentans		26.27	0.22		99	99	
Pseudoflavonifractor capillosus		2.04	-.03		1	1	
44201000	**Acceptable identification**						**44201000**
					SPO		
Atopobium minutum		92.72	0.38	SAL 1	1		
Clostridium bifermentans		5.47	0.09		99		
Eubacterium tortuosum		1.81	-.03		1		
44201002	**Good identification**						**44201002**
Atopobium minutum		100.0	0.35	SAL 1			
44210000	**Low discrimination**						**44210000**
					GRA	SPO	
Atopobium minutum		50.74	0.46	SAL 1	99	1	
Clostridium difficile		20.13	0.28		99	99	
Fusobacterium mortiferum		9.67	0.39		1	1	
44210002	**Good identification**						**44210002**
					GRA		
Atopobium minutum		96.92	0.43	SAL 1	99		
Fusobacterium mortiferum		3.08	0.22		1		
44210010	**Good identification**						**44210010**
					SPO		
Clostridium difficile		99.48	0.40	GLU 99	99		
Atopobium minutum		0.52	0.11	MAN 90	1		
44210020	**Good identification**						**44210020**
					SPO		
Atopobium minutum		97.92	0.37	SAL 1	1		
Clostridium bifermentans		2.08	0.01		99		
44210100	**Good identification**						**44210100**
					GRA		
Fusobacterium mortiferum		97.78	0.45	GLU 99	1		
Atopobium minutum		2.22	0.11		99		
44211000	**Low discrimination**						**44211000**
					GRA		
Atopobium minutum		53.83	0.37	SAL 1	99		
Fusobacterium mortiferum		46.17	0.42		1		
44211100	**Good identification**						**44211100**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.48	GLU 99	1	1	
Clostridium septicum		0.77	-.06		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
44212000	**Low discrimination**						**44212000**
					SPO		
Clostridium difficile		81.67	0.45	GLU 99	99		
Eubacterium limosum		18.10	0.32		1		
Atopobium minutum		0.23	0.11		1		
44212010	**Very good identification**						**44212010**
					SPO		
Clostridium difficile		99.95	0.56	GLU 99	99		
Eubacterium limosum		0.05	-.03		1		
44230100	**Good identification**						**44230100**
Fusobacterium mortiferum		100.0	0.36	GLU 99			
44231100	**Acceptable identification**						**44231100**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	GLU 99	1	1	
Clostridium septicum		7.27	0.03		99	99	
44231120	**Good identification**						**44231120**
Clostridium septicum		100.0	0.37	GLU 99			
44232010	**Good identification**						**44232010**
Clostridium difficile		100.0	0.37	GLU 99			
44250100	**Good identification**						**44250100**
Fusobacterium mortiferum		100.0	0.37	GLU 99			
44251100	**Low discrimination**						**44251100**
					GRA		
Fusobacterium mortiferum		73.20	0.40	GLU 99	1		
Lactobacillus catenaformis		10.30	0.15		99		
Collinsella aerofaciens		6.82	-.03		99		
44251120	**Acceptable identification**						**44251120**
Clostridium septicum		91.77	0.37	GLU 99			
Clostridium paraputrificum		5.61	0.14	MLT 94			
Clostridium barati		2.18	0.12				
44253000	**Good identification**						**44253000**
Clostridium innocuum		100.0	0.49	GLU 99			
44253002	**Good identification**						**44253002**
Clostridium innocuum		100.0	0.49	GLU 99			
44253400	**Good identification**						**44253400**
Clostridium innocuum		100.0	0.36	GLU 99			
44253402	**Good identification**						**44253402**
Clostridium innocuum		100.0	0.36	GLU 99			
44271120	**Good identification**						**44271120**
Clostridium septicum		99.46	0.45	GLU 99			
Clostridium paraputrificum		0.54	0.04	MLT 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
44273000	**Good identification**						**44273000**
Clostridium innocuum		100.0	0.41	GLU 99			
44273002	**Good identification**						**44273002**
Clostridium innocuum		100.0	0.41	GLU 99			
44300000	**Acceptable identification**						**44300000**
Clostridium bifermentans		91.85	0.49	IND 90			
Clostridium botulinum biovar A		5.48	0.29	GLU 92			
Clostridium sporogenes		1.95	0.19	SAL 10			
44311120	**Good identification**						**44311120**
Clostridium septicum		97.41	0.49	GLU 99			
Clostridium perfringens		1.25	0.23				
Clostridium paraputrificum		1.20	0.14				
44331120	**Very good identification**						**44331120**
					COC	SPO	
Clostridium septicum		98.87	0.57	GLU 99	1	99	
Clostridium perfringens		0.78	0.28		1	84	
Atopobium parvulum		0.24	0.18		99	1	
44335100	**Good identification**						**44335100**
Atopobium parvulum		100.0	0.36	GLU 99			
44351000	**Low discrimination**						**44351000**
					SPO		
Lactobacillus jensenii		81.12	0.41	GLU 99	1		
Lactobacillus catenaformis		14.69	0.29	SUC 90	1		
Clostridium barati		1.65	0.07		99		
44351002	**Low discrimination**						**44351002**
					SPO		
Lactobacillus jensenii		73.01	0.38	GLU 99	1		
Lactobacillus catenaformis		19.83	0.29	SUC 90	1		
Clostridium barati		6.67	0.15		99		
44351120	**Low discrimination**						**44351120**
					SPO		
Clostridium septicum		71.11	0.58	GLU 99	99		
Clostridium paraputrificum		27.48	0.48		99		
Collinsella aerofaciens		0.63	0.15		1		
44351122	**Low discrimination**						**44351122**
Clostridium paraputrificum		72.39	0.41	GLU 99			
Clostridium barati		9.72	0.32	SUC 92			
Clostridium septicum		8.92	0.29				
44371000	**Good identification**						**44371000**
					COC	SPO	
Lactobacillus jensenii		99.50	0.50	GLU 99	1	1	
Clostridium septicum		0.25	-.03	SUC 90	1	99	
Atopobium parvulum		0.25	0.04		99	1	
44371002	**Good identification**						**44371002**
					SPO		
Lactobacillus jensenii		98.99	0.47	GLU 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		0.74	0.07	SUC 90	1		
Clostridium barati		0.26	-.03		99		
44371010	**Good identification**						**44371010**
Lactobacillus jensenii		100.0	0.41	GLU 99			
44371012	**Good identification**						**44371012**
					GRA		
Lactobacillus jensenii		97.78	0.38	GLU 99	99		
Leptotrichia buccalis		2.22	-.03	SUC 90	1		
44371020	**Low discrimination**						**44371020**
					SPO		
Lactobacillus jensenii		49.09	0.39	GLU 99	1		
Clostridium septicum		48.89	0.32	SUC 90	99		
Clostridium paraputrificum		1.67	0.04		99		
44371022	**Low discrimination**						**44371022**
					SPO		
Lactobacillus jensenii		89.75	0.36	GLU 99	1		
Clostridium barati		2.85	0.05	SUC 90	99		
Clostridium septicum		2.74	0.03		99		
44371120	**Very good identification**						**44371120**
					COC	SPO	
Clostridium septicum		96.48	0.66	GLU 99	1	99	
Clostridium paraputrificum		3.30	0.38		1	99	
Atopobium parvulum		0.17	0.25		99	1	
44371122	**Low discrimination**						**44371122**
					SPO		
Clostridium septicum		52.30	0.37	GLU 99	99		
Clostridium paraputrificum		37.61	0.32	BGL 2	99		
Lactobacillus acidophilus		5.99	0.25		1		
44375100	**Good identification**						**44375100**
					COC	SPO	
Atopobium parvulum		99.08	0.42	GLU 99	99	1	
Clostridium septicum		0.55	-.03	SUC 96	1	99	
Clostridium butyricum		0.37	-.07		1	99	
44400000	**Low discrimination**						**44400000**
					COC	SPO	
Anaerococcus prevotii		62.32	0.41	ESL 1	99	1	
Atopobium minutum		19.16	0.46	URE 90	1	1	
Clostridium botulinum biovar A		6.34	0.30		1	99	
44400001	**Very good identification**						**44400001**
					COC	GRA	
Anaerococcus prevotii		99.90	0.58	ESL 1	99	99	
Campylobacter ureolyticus		0.05	-.03		1	1	
Atopobium minutum		0.03	0.12		1	99	
44400003	**Good identification**						**44400003**
					COC		
Anaerococcus prevotii		99.91	0.47	ESL 1	99		
Atopobium minutum		0.09	0.09		1		
44400020	**Low discrimination**						**44400020**
					COC	SPO	
Atopobium minutum		84.89	0.38	RAF 1	1	1	
Anaerococcus prevotii		8.37	0.07		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		2.88	0.05		1	99	
44401000	**Acceptable identification**						**44401000**
					COC		
Atopobium minutum		91.03	0.38	RAF 1	1		
Anaerococcus prevotii		8.97	0.07		99		
44401002	**Good identification**						**44401002**
					COC		
Atopobium minutum		96.44	0.35	RAF 1	1		
Anaerococcus prevotii		3.57	-.03		99		
44410000	**Low discrimination**						**44410000**
					GRA		
Atopobium minutum		65.32	0.46	RAF 1	99		
Fusobacterium mortiferum		12.45	0.39		1		
Eubacterium limosum		8.41	0.18		99		
44410002	**Good identification**						**44410002**
					COC	GRA	
Atopobium minutum		95.68	0.43	RAF 1	1	99	
Fusobacterium mortiferum		3.04	0.22		1	1	
Anaerococcus prevotii		1.28	-.03		99	99	
44410020	**Good identification**						**44410020**
Atopobium minutum		100.0	0.37	RAF 1			
44410100	**Good identification**						**44410100**
					GRA		
Fusobacterium mortiferum		97.78	0.45	GLU 99	1		
Atopobium minutum		2.22	0.11		99		
44411000	**Low discrimination**						**44411000**
					GRA		
Atopobium minutum		53.13	0.37	RAF 1	99		
Fusobacterium mortiferum		45.57	0.42		1		
Prevotella intermedia		1.29	0.04		1		
44411100	**Good identification**						**44411100**
Fusobacterium mortiferum		100.0	0.48	GLU 99			
44430100	**Good identification**						**44430100**
Fusobacterium mortiferum		100.0	0.36	GLU 99			
44431100	**Good identification**						**44431100**
Fusobacterium mortiferum		100.0	0.39	GLU 99			
44435122	**Good identification**						**44435122**
Parabacteroides distasonis		100.0	0.37	GLU 99			
44435322	**Good identification**						**44435322**
Parabacteroides distasonis		100.0	0.46	GLU 99			
44435332	**Good identification**						**44435332**
Parabacteroides distasonis		100.0	0.41	GLU 99			
44435722	**Good identification**						**44435722**
Parabacteroides distasonis		100.0	0.42	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
44435732	**Good identification**						**44435732**
Parabacteroides distasonis		100.0	0.37	GLU 99			
44450100	**Good identification**						**44450100**
Fusobacterium mortiferum		100.0	0.37	GLU 99			
44451100	**Good identification**						**44451100**
Fusobacterium mortiferum		100.0	0.40	GLU 99			
44475322	**Good identification**						**44475322**
Parabacteroides distasonis		100.0	0.41	GLU 99			
44475332	**Good identification**						**44475332**
Parabacteroides distasonis		100.0	0.36	GLU 99			
44475722	**Good identification**						**44475722**
Parabacteroides distasonis		100.0	0.37	GLU 99			
44510000	**Low discrimination**						**44510000**
					GRA	SPO	
Prevotella intermedia		88.54	0.38	ESL 1	1	1	
Fusobacterium mortiferum		2.60	0.23	GLU 90	1	1	
Clostridium bifermentans		2.60	0.10		99	99	
44511000	**Good identification**						**44511000**
Prevotella intermedia		95.78	0.38	ESL 1			
Fusobacterium mortiferum		4.22	0.26	GLU 90			
44511120	**Good identification**						**44511120**
					GRA	CAT	
Prevotella melaninogenica		97.16	0.47	GLU 90	1	1	
Clostridium perfringens		1.32	0.21	SUC 99	99	1	
Capnocytophaga ochracea		0.59	0.12		1	90	
44515322	**Low discrimination**						**44515322**
					CAT		
Parabacteroides distasonis		89.53	0.44	GLU 99	76		
Prevotella oralis		7.14	0.12		1		
Bacteroides fragilis		3.33	-.03		99		
44515332	**Good identification**						**44515332**
Parabacteroides distasonis		100.0	0.39	GLU 99			
44515720	**Good identification**						**44515720**
Bacteroides vulgatus		100.0	0.42	GLU 99			
44515722	**Low discrimination**						**44515722**
Parabacteroides distasonis		86.16	0.40	GLU 99			
Bacteroides vulgatus		13.84	0.08				
44515732	**Acceptable identification**						**44515732**
Parabacteroides distasonis		91.16	0.35	GLU 99			
Clostridium clostridioforme		8.84	0.12				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
44531322	**Acceptable identification**					**44531322**
Parabacteroides distasonis		92.48	0.39	GLU 99		
Prevotella oralis		7.53	0.07	XYL 90		
44531722	**Good identification**					**44531722**
Parabacteroides distasonis		100.0	0.35	GLU 99		
44535122	**Good identification**					**44535122**
					CAT	
Parabacteroides distasonis		96.87	0.47	GLU 99	76	
Bacteroides fragilis		2.37	-.03		99	
Prevotella oralis		0.76	-.02		1	
44535132	**Good identification**					**44535132**
Parabacteroides distasonis		100.0	0.42	GLU 99		
44535322	**Good identification**					**44535322**
Parabacteroides distasonis		100.0	0.55	GLU 99		
44535332	**Good identification**					**44535332**
Parabacteroides distasonis		99.39	0.50	GLU 99		
Clostridium clostridioforme		0.61	0.07			
44535522	**Acceptable identification**					**44535522**
					GRA	
Parabacteroides distasonis		94.80	0.43	GLU 99	1	
Bifidobacterium longum subsp. longum		5.20	0.17		99	
44535532	**Low discrimination**					**44535532**
					GRA	
Parabacteroides distasonis		80.82	0.39	GLU 99	1	
Bifidobacterium longum subsp. longum		14.02	0.21		99	
Clostridium clostridioforme		5.16	0.12		50	
44535722	**Good identification**					**44535722**
Parabacteroides distasonis		96.47	0.51	GLU 99		
Clostridium clostridioforme		1.78	0.16			
Bacteroides thetaiotaomicron		1.75	0.06			
44535732	**Low discrimination**					**44535732**
Parabacteroides distasonis		85.53	0.47	GLU 99		
Clostridium clostridioforme		13.35	0.27			
Bacteroides thetaiotaomicron		1.12	-.02			
44551122	**Acceptable identification**					**44551122**
					CAT	
Prevotella oralis		91.64	0.43	GLU 99	1	
Prevotella melaninogenica		4.71	0.18	SUC 92	1	
Capnocytophaga ochracea		3.04	0.18		90	
44551322	**Good identification**					**44551322**
Prevotella oralis		97.45	0.37	GLU 99		
Parabacteroides distasonis		2.55	0.23	SUC 92		
44555322	**Low discrimination**					**44555322**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		64.32	0.39	GLU 99	76	
Prevotella oralis		33.78	0.22		1	
Bacteroides fragilis		1.90	-.09		99	
44555722	**Acceptable identification**					**44555722**
Parabacteroides distasonis		93.35	0.35	GLU 99		
Clostridium clostridioforme		6.65	0.10			
44575122	**Acceptable identification**					**44575122**
					CAT	
Parabacteroides distasonis		93.37	0.42	GLU 99	76	
Prevotella oralis		4.81	0.07		1	
Bacteroides fragilis		1.82	-.09		99	
44575132	**Good identification**					**44575132**
Parabacteroides distasonis		100.0	0.38	GLU 99		
44575322	**Good identification**					**44575322**
Parabacteroides distasonis		98.83	0.50	GLU 99		
Prevotella oralis		0.73	0.01			
Clostridium clostridioforme		0.44	0.05			
44575332	**Good identification**					**44575332**
Parabacteroides distasonis		96.34	0.46	GLU 99		
Clostridium clostridioforme		3.66	0.16			
44575522	**Good identification**					**44575522**
Parabacteroides distasonis		95.52	0.39	GLU 99		
Clostridium clostridioforme		4.48	0.10			
44575712	**Good identification**					**44575712**
Clostridium clostridioforme		100.0	0.36	GLU 88		
44575722	**Low discrimination**					**44575722**
Parabacteroides distasonis		84.30	0.47	GLU 99		
Clostridium clostridioforme		9.67	0.25			
Bacteroides thetaiotaomicron		6.03	0.11			
44575732	**Low discrimination**					**44575732**
Parabacteroides distasonis		49.46	0.42	GLU 99		
Clostridium clostridioforme		47.99	0.36			
Bacteroides thetaiotaomicron		2.55	0.04			
44610000	**Good identification**					**44610000**
					GRA	
Fusobacterium mortiferum		97.34	0.44	GLU 99	1	
Atopobium minutum		2.66	0.11		99	
44610100	**Good identification**					**44610100**
Fusobacterium mortiferum		100.0	0.50	GLU 99		
44611000	**Good identification**					**44611000**
Fusobacterium mortiferum		100.0	0.47	GLU 99		
44611100	**Good identification**					**44611100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.53	GLU 99		
44611102	**Good identification**					**44611102**
					GRA	
Fusobacterium mortiferum		96.99	0.37	GLU 99	1	
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99	
44630100	**Good identification**					**44630100**
Fusobacterium mortiferum		100.0	0.41	GLU 99		
44631000	**Good identification**					**44631000**
Fusobacterium mortiferum		100.0	0.38	GLU 99		
44631100	**Good identification**					**44631100**
Fusobacterium mortiferum		100.0	0.44	GLU 99		
44635122	**Good identification**					**44635122**
Parabacteroides distasonis		100.0	0.38	GLU 99		
44635322	**Good identification**					**44635322**
Parabacteroides distasonis		100.0	0.46	GLU 99		
44635332	**Good identification**					**44635332**
Parabacteroides distasonis		100.0	0.42	GLU 99		
44635722	**Good identification**					**44635722**
Parabacteroides distasonis		100.0	0.43	GLU 99		
44635732	**Good identification**					**44635732**
Parabacteroides distasonis		96.36	0.38	GLU 99		
Clostridium clostridioforme		3.65	0.08			
44650000	**Good identification**					**44650000**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
44650100	**Good identification**					**44650100**
Fusobacterium mortiferum		100.0	0.42	GLU 99		
44651000	**Good identification**					**44651000**
Fusobacterium mortiferum		100.0	0.39	GLU 99		
44651100	**Good identification**					**44651100**
Fusobacterium mortiferum		100.0	0.45	GLU 99		
44671100	**Good identification**					**44671100**
Fusobacterium mortiferum		100.0	0.36	GLU 99		
44675322	**Good identification**					**44675322**
Parabacteroides distasonis		100.0	0.42	GLU 99		
44675332	**Good identification**					**44675332**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		100.0	0.37	GLU 99		
44675722	**Good identification**					**44675722**
Parabacteroides distasonis		97.29	0.38	GLU 99		
Clostridium clostridioforme		2.71	0.06			
44711100	**Low discrimination**					**44711100**
					GRA	
Fusobacterium mortiferum		78.55	0.37	GLU 99	1	
Bifidobacterium longum subsp. infantis		21.45	0.10	MLT 11	99	
44711122	**Low discrimination**					**44711122**
					GRA	
Prevotella oralis		59.29	0.38	GLU 99	1	
Bifidobacterium longum subsp. infantis		38.81	0.32	SUC 92	99	
Parabacteroides distasonis		0.89	0.20		1	
44715322	**Low discrimination**					**44715322**
					GRA	
Parabacteroides distasonis		86.69	0.44	GLU 99	1	
Prevotella oralis		11.35	0.16		1	
Bifidobacterium longum subsp. infantis		1.16	-.04		99	
44715332	**Good identification**					**44715332**
Parabacteroides distasonis		97.50	0.40	GLU 99		
Clostridium clostridioforme		2.50	0.07			
44715722	**Low discrimination**					**44715722**
Parabacteroides distasonis		89.72	0.41	GLU 99		
Clostridium clostridioforme		6.96	0.16			
Bacteroides vulgatus		3.33	-.02			
44715732	**Low discrimination**					**44715732**
Parabacteroides distasonis		60.38	0.36	GLU 99		
Clostridium clostridioforme		39.63	0.27			
44731322	**Low discrimination**					**44731322**
Parabacteroides distasonis		88.22	0.39	GLU 99		
Prevotella oralis		11.78	0.12	XYL 90		
44731722	**Low discrimination**					**44731722**
Parabacteroides distasonis		90.00	0.36	GLU 99		
Clostridium clostridioforme		10.00	0.14	XYL 90		
44735122	**Good identification**					**44735122**
					GRA	
Parabacteroides distasonis		97.88	0.47	GLU 99	1	
Prevotella oralis		1.26	0.02		1	
Bifidobacterium longum subsp. infantis		0.86	-.04		99	
44735132	**Good identification**					**44735132**
Parabacteroides distasonis		98.34	0.43	GLU 99		
Clostridium clostridioforme		1.66	0.07			
44735322	**Good identification**					**44735322**
Parabacteroides distasonis		99.51	0.56	GLU 99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.49	0.10				
44735332	**Good identification**						**44735332**
Parabacteroides distasonis		96.03	0.51	GLU 99			
Clostridium clostridioforme		3.97	0.22				
44735522	**Good identification**						**44735522**
Parabacteroides distasonis		95.14	0.44	GLU 99			
Clostridium clostridioforme		4.86	0.16				
44735532	**Low discrimination**						**44735532**
Parabacteroides distasonis		69.82	0.39	GLU 99			
Clostridium clostridioforme		30.18	0.27				
44735712	**Good identification**						**44735712**
Clostridium clostridioforme		99.05	0.42	GLU 88			
Parabacteroides distasonis		0.95	0.13	SUC 99			
44735722	**Low discrimination**						**44735722**
Parabacteroides distasonis		88.90	0.52	GLU 99			
Clostridium clostridioforme		11.10	0.31				
44735732	**Low discrimination**						**44735732**
Clostridium clostridioforme		51.37	0.42	GLU 88			
Parabacteroides distasonis		48.63	0.47	SUC 99			
44751122	**Acceptable identification**						**44751122**
					GRA	SPO	
Prevotella oralis		90.09	0.48	GLU 99	1	1	
Clostridium paraputrificum		6.75	0.22	SUC 92	99	99	
Bifidobacterium breve		2.34	0.32		99	1	
44751322	**Good identification**						**44751322**
					SPO		
Prevotella oralis		98.15	0.41	GLU 99	1		
Parabacteroides distasonis		1.56	0.23	SUC 92	1		
Clostridium ramosum		0.28	-.02		92		
44755322	**Low discrimination**						**44755322**
Parabacteroides distasonis		53.17	0.40	GLU 99			
Prevotella oralis		45.82	0.26				
Clostridium clostridioforme		1.00	0.05				
44755332	**Low discrimination**						**44755332**
Parabacteroides distasonis		86.23	0.35	GLU 99			
Clostridium clostridioforme		13.77	0.16				
44755712	**Good identification**						**44755712**
Clostridium clostridioforme		100.0	0.36	GLU 88			
44755722	**Low discrimination**						**44755722**
Parabacteroides distasonis		61.54	0.36	GLU 99			
Clostridium clostridioforme		29.68	0.25				
Bacteroides uniformis		8.77	0.03				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
44755732	**Low discrimination**						**44755732**
Clostridium clostridioforme		80.32	0.36	GLU 88			
Parabacteroides distasonis		19.68	0.31	SUC 99			
44774712	**Good identification**						**44774712**
Clostridium clostridioforme		100.0	0.43	GLU 88			
44774732	**Good identification**						**44774732**
Clostridium clostridioforme		100.0	0.43	GLU 88			
44775122	**Acceptable identification**						**44775122**
Parabacteroides distasonis		91.16	0.43	GLU 99			
Prevotella oralis		7.71	0.12				
Clostridium clostridioforme		1.13	0.05				
44775132	**Acceptable identification**						**44775132**
Parabacteroides distasonis		90.49	0.38	GLU 99			
Clostridium clostridioforme		9.51	0.16				
44775322	**Good identification**						**44775322**
Parabacteroides distasonis		95.93	0.51	GLU 99			
Clostridium clostridioforme		2.91	0.19				
Prevotella oralis		1.16	0.05				
44775332	**Low discrimination**						**44775332**
Parabacteroides distasonis		79.56	0.46	GLU 99			
Clostridium clostridioforme		20.44	0.31				
44775512	**Low discrimination**						**44775512**
					SPO		
Clostridium clostridioforme		76.16	0.36	GLU 88	66		
Bifidobacterium adolescentis		18.07	0.29	SUC 99	1		
Clostridium butyricum		4.04	0.03	RHA 88	99		
44775522	**Low discrimination**						**44775522**
					GRA	SPO	
Parabacteroides distasonis		69.23	0.39	GLU 99	1	1	
Clostridium clostridioforme		21.99	0.25		50	66	
Clostridium butyricum		8.78	0.07		99	99	
44775532	**Low discrimination**						**44775532**
					GRA		
Clostridium clostridioforme		70.79	0.36	GLU 88	50		
Parabacteroides distasonis		26.34	0.35	SUC 99	1		
Bifidobacterium dentium		2.87	-.03	RHA 88	99		
44775612	**Good identification**						**44775612**
Clostridium clostridioforme		100.0	0.42	GLU 88			
44775632	**Good identification**						**44775632**
Clostridium clostridioforme		100.0	0.42	GLU 88			
44775702	**Good identification**						**44775702**
					GRA	SPO	
Clostridium clostridioforme		96.87	0.40	GLU 88	50	66	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		1.88	-.01	SUC 99	99	99	
Parabacteroides distasonis		1.26	0.13		1	1	
44775712	**Good identification**						**44775712**
Clostridium clostridioforme		100.0	0.51	GLU 88			
44775722	**Low discrimination**						**44775722**
Parabacteroides distasonis		54.93	0.47	GLU 99			
Clostridium clostridioforme		42.65	0.40				
Bacteroides ovatus		1.78	0.06				
44775732	**Low discrimination**						**44775732**
Clostridium clostridioforme		86.33	0.51	GLU 88			
Parabacteroides distasonis		13.14	0.43	SUC 99			
Bacteroides ovatus		0.53	0.03				
44777102	**Acceptable identification**						**44777102**
Actinomyces israelii		92.86	0.37	GLU 99			
Bifidobacterium adolescentis		7.14	0.12	SUC 98			
44777502	**Low discrimination**						**44777502**
					SPO		
Actinomyces israelii		46.25	0.35	GLU 99	1		
Bifidobacterium adolescentis		29.08	0.26	SUC 98	1		
Clostridium butyricum		20.50	0.08		99		
45000000	**Low discrimination**						**45000000**
					SPO		
Clostridium botulinum biovar A		41.02	0.81	MNS 7	99		
Clostridium bifermentans		28.01	0.77		99		
Atopobium minutum		23.44	0.84		1		
45000001	**Low discrimination**						**45000001**
					SPO		
Clostridium botulinum biovar A		31.05	0.47	MNS 7	99		
Clostridium bifermentans		21.20	0.43	URE 1	99		
Atopobium minutum		17.75	0.50		1		
45000002	**Low discrimination**						**45000002**
					SPO		
Atopobium minutum		88.04	0.81		1		
Clostridium sporogenes		7.60	0.53		99		
Clostridium botulinum biovar A		2.33	0.47		99		
45000003	**Low discrimination**						**45000003**
					COC	SPO	
Atopobium minutum		64.58	0.47	URE 1	1	1	
Clostridium novyi biovar A		14.34	0.40		1	99	
Anaerococcus prevotii		12.62	0.21		99	1	
45000010	**Low discrimination**						**45000010**
Clostridium difficile		42.28	0.50	FRU 99			
Clostridium botulinum biovar A		19.87	0.47	MAN 90			
Clostridium bifermentans		13.57	0.43				
45000011	**Acceptable identification**						**45000011**
Clostridium novyi biovar A		91.03	0.53	MNS 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		4.08	0.15				
Clostridium botulinum biovar A		1.92	0.13				
45000012	**Low discrimination**						**45000012**
					SPO		
Atopobium minutum		70.95	0.47	MLZ 1	1		
Clostridium novyi biovar A		15.76	0.40		99		
Clostridium sporogenes		6.13	0.19		99		
45000013	**Good identification**						**45000013**
					SPO		
Clostridium novyi biovar A		95.65	0.40	MNS 1	99		
Atopobium minutum		4.35	0.13		1		
45000020	**Low discrimination**						**45000020**
					SPO		
Atopobium minutum		75.82	0.76		1		
Clostridium bifermentans		17.34	0.57		99		
Clostridium botulinum biovar A		4.02	0.47		99		
45000021	**Low discrimination**						**45000021**
					SPO		
Atopobium minutum		51.33	0.42	URE 1	1		
Clostridium novyi biovar A		32.30	0.43		99		
Clostridium bifermentans		11.74	0.23		99		
45000022	**Very good identification**						**45000022**
					GRA	SPO	
Atopobium minutum		96.67	0.73		99	1	
Pseudoflavonifractor capillosus		2.47	0.32		1	1	
Clostridium bifermentans		0.34	0.23		99	99	
45000023	**Low discrimination**						**45000023**
					GRA	SPO	
Atopobium minutum		83.89	0.39	URE 1	99	1	
Clostridium novyi biovar A		13.97	0.30		99	99	
Pseudoflavonifractor capillosus		2.14	-.03		1	1	
45000030	**Low discrimination**						**45000030**
					SPO		
Atopobium minutum		48.89	0.42	MLZ 1	1		
Clostridium novyi biovar A		30.77	0.43		99		
Clostridium bifermentans		11.18	0.23		99		
45000031	**Good identification**						**45000031**
					SPO		
Clostridium novyi biovar A		98.42	0.43	MNS 1	99		
Atopobium minutum		1.58	0.08		1		
45000032	**Low discrimination**						**45000032**
					GRA	SPO	
Atopobium minutum		83.89	0.39	MLZ 1	99	1	
Clostridium novyi biovar A		13.97	0.30		99	99	
Pseudoflavonifractor capillosus		2.14	-.03		1	1	
45000040	**Low discrimination**						**45000040**
					SPO		
Clostridium botulinum biovar A		88.12	0.72	MNS 7	99		
Clostridium bifermentans		8.39	0.54		99		
Atopobium minutum		1.70	0.50		1		
45000041	**Low discrimination**						**45000041**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		48.76	0.50	MNS 1			
Clostridium botulinum biovar A		45.56	0.38				
Clostridium bifermentans		4.34	0.20				
45000042	**Low discrimination**						**45000042**
					SPO		
Atopobium minutum		47.78	0.47	SOR 1	1		
Clostridium botulinum biovar A		37.45	0.38		99		
Clostridium novyi biovar A		7.07	0.37		99		
45000043	**Low discrimination**						**45000043**
					SPO		
Clostridium novyi biovar A		89.15	0.37	MNS 1	99		
Atopobium minutum		6.08	0.13		1		
Clostridium botulinum biovar A		4.77	0.04		99		
45000050	**Low discrimination**						**45000050**
Clostridium novyi biovar A		41.81	0.50	MNS 1			
Clostridium botulinum biovar A		39.06	0.38				
Clostridium difficile		14.65	0.28				
45000051	**Good identification**						**45000051**
Clostridium novyi biovar A		99.07	0.50	MNS 1			
Clostridium botulinum biovar A		0.94	0.04				
45000052	**Low discrimination**						**45000052**
					SPO		
Clostridium novyi biovar A		89.15	0.37	MNS 1	99		
Atopobium minutum		6.08	0.13		1		
Clostridium botulinum biovar A		4.77	0.04		99		
45000053	**Good identification**						**45000053**
Clostridium novyi biovar A		100.0	0.37	MNS 1			
45000060	**Low discrimination**						**45000060**
					SPO		
Clostridium botulinum biovar A		39.67	0.38	MNS 7	99		
Atopobium minutum		25.30	0.42	NAG 1	1		
Clostridium bifermentans		23.87	0.33		99		
45000061	**Acceptable identification**						**45000061**
					SPO		
Clostridium novyi biovar A		92.21	0.40	MNS 1	99		
Clostridium botulinum biovar A		3.48	0.04		99		
Atopobium minutum		2.22	0.08		1		
45000062	**Low discrimination**						**45000062**
					GRA	SPO	
Atopobium minutum		85.14	0.39	SOR 1	99	1	
Clostridium novyi biovar A		9.45	0.27		99	99	
Pseudoflavonifractor capillosus		2.17	-.03		1	1	
45000070	**Acceptable identification**						**45000070**
					SPO		
Clostridium novyi biovar A		92.21	0.40	MNS 1	99		
Clostridium botulinum biovar A		3.48	0.04		99		
Atopobium minutum		2.22	0.08		1		
45000071	**Good identification**						**45000071**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		100.0	0.40	MNS 1		
45000100	**Low discrimination**					**45000100**
					SPO	
Clostridium botulinum biovar A		39.81	0.47	MNS 7	99	
Clostridium bifermentans		27.18	0.43	LAC 1	99	
Atopobium minutum		22.75	0.50		1	
45000102	**Low discrimination**					**45000102**
					SPO	
Atopobium minutum		86.83	0.47	LAC 1	1	
Clostridium sporogenes		7.50	0.19		99	
Clostridium botulinum biovar A		2.30	0.13		99	
45000122	**Acceptable identification**					**45000122**
					GRA	
Atopobium minutum		93.18	0.39	LAC 1	99	
Prevotella bivia		4.45	-.03		1	
Pseudoflavonifractor capillosus		2.38	-.03		1	
45000140	**Low discrimination**					**45000140**
					SPO	
Clostridium botulinum biovar A		88.86	0.38	MNS 7	99	
Clostridium bifermentans		8.46	0.20	LAC 1	99	
Atopobium minutum		1.72	0.16		1	
45000200	**Low discrimination**					**45000200**
					SPO	
Clostridium botulinum biovar A		41.07	0.47	MNS 7	99	
Clostridium bifermentans		28.04	0.43	RHA 1	99	
Atopobium minutum		23.47	0.50		1	
45000202	**Low discrimination**					**45000202**
					SPO	
Atopobium minutum		88.42	0.47	RHA 1	1	
Clostridium sporogenes		7.63	0.19		99	
Clostridium botulinum biovar A		2.34	0.13		99	
45000220	**Low discrimination**					**45000220**
					SPO	
Atopobium minutum		76.73	0.42	RHA 1	1	
Clostridium bifermentans		17.55	0.23		99	
Clostridium botulinum biovar A		4.07	0.13		99	
45000222	**Good identification**					**45000222**
					GRA	
Atopobium minutum		97.51	0.39	RHA 1	99	
Pseudoflavonifractor capillosus		2.49	-.03		1	
45000240	**Low discrimination**					**45000240**
					SPO	
Clostridium botulinum biovar A		88.86	0.38	MNS 7	99	
Clostridium bifermentans		8.46	0.20	RHA 1	99	
Atopobium minutum		1.72	0.16		1	
45000400	**Low discrimination**					**45000400**
					SPO	
Clostridium botulinum biovar A		38.96	0.47	MNS 7	99	
Clostridium bifermentans		26.60	0.43	ARA 1	99	
Atopobium minutum		22.27	0.50		1	
45000402	**Low discrimination**					**45000402**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		88.12	0.47	ARA 1	1		
Clostridium sporogenes		7.61	0.19		99		
Clostridium botulinum biovar A		2.34	0.13		99		
45000420	**Low discrimination**						**45000420**
					COC	SPO	
Atopobium minutum		71.91	0.42	ARA 1	1	1	
Clostridium bifermentans		16.45	0.23		1	99	
Gemella morbillorum		6.08	0.19		99	1	
45000422	**Good identification**						**45000422**
					GRA		
Atopobium minutum		97.51	0.39	ARA 1	99		
Pseudoflavonifractor capillosus		2.49	-.03		1		
45000440	**Low discrimination**						**45000440**
					SPO		
Clostridium botulinum biovar A		88.47	0.38	MNS 7	99		
Clostridium bifermentans		8.43	0.20	ARA 1	99		
Atopobium minutum		1.71	0.16		1		
45001000	**Low discrimination**						**45001000**
					COC	SPO	
Atopobium minutum		87.68	0.76		1	1	
Clostridium botulinum biovar A		4.65	0.47		1	99	
Gemella morbillorum		3.27	0.47		99	1	
45001001	**Low discrimination**						**45001001**
					SPO		
Atopobium minutum		70.33	0.42	URE 1	1		
Clostridium novyi biovar A		19.67	0.37		99		
Clostridium botulinum biovar A		3.73	0.13		99		
45001002	**Very good identification**						**45001002**
					SPO		
Atopobium minutum		99.44	0.73		1		
Clostridium sporogenes		0.26	0.19		99		
Clostridium botulinum biovar A		0.08	0.13		99		
45001003	**Acceptable identification**						**45001003**
					SPO		
Atopobium minutum		93.11	0.39	URE 1	1		
Clostridium novyi biovar A		6.89	0.24		99		
45001010	**Low discrimination**						**45001010**
					SPO		
Atopobium minutum		65.83	0.42	MLZ 1	1		
Clostridium novyi biovar A		18.41	0.37		99		
Clostridium difficile		7.43	0.15		99		
45001011	**Good identification**						**45001011**
					SPO		
Clostridium novyi biovar A		96.51	0.37	MNS 1	99		
Atopobium minutum		3.49	0.08	GAL 10	1		
45001012	**Acceptable identification**						**45001012**
					SPO		
Atopobium minutum		93.11	0.39	MLZ 1	1		
Clostridium novyi biovar A		6.89	0.24		99		
45001020	**Very good identification**						**45001020**
					COC	SPO	
Atopobium minutum		97.65	0.68		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Gemella morbillorum		1.21	0.31		99	1	
Clostridium bifermentans		0.68	0.23		1	99	
45001022	**Very good identification**						**45001022**
					GRA	SPO	
Atopobium minutum		99.10	0.65		99	1	
Pseudoflavonifractor capillosus		0.84	0.15		1	1	
Clostridium novyi biovar A		0.06	0.13		99	99	
45001040	**Low discrimination**						**45001040**
					SPO		
Clostridium botulinum biovar A		52.45	0.38	MNS 7	99		
Atopobium minutum		33.45	0.42	GAL 1	1		
Clostridium novyi biovar A		6.24	0.34		99		
45001042	**Acceptable identification**						**45001042**
					SPO		
Atopobium minutum		93.19	0.39	SOR 1	1		
Clostridium novyi biovar A		4.60	0.21		99		
Clostridium botulinum biovar A		2.21	0.04		99		
45001100	**Low discrimination**						**45001100**
					GRA	SPO	
Atopobium minutum		58.53	0.42	LAC 1	99	1	
Fusobacterium mortiferum		32.71	0.43		1	1	
Clostridium botulinum biovar A		3.10	0.13		99	99	
45001102	**Acceptable identification**						**45001102**
					GRA		
Atopobium minutum		91.48	0.39	LAC 1	99		
Fusobacterium mortiferum		8.52	0.27		1		
45001200	**Low discrimination**						**45001200**
					COC	SPO	
Atopobium minutum		88.77	0.42	RHA 1	1	1	
Clostridium botulinum biovar A		4.71	0.13		1	99	
Gemella morbillorum		3.31	0.13		99	1	
45001202	**Good identification**						**45001202**
Atopobium minutum		100.0	0.39	RHA 1			
45001400	**Low discrimination**						**45001400**
					COC	SPO	
Atopobium minutum		66.24	0.42	ARA 1	1	1	
Gemella morbillorum		27.15	0.31		99	1	
Clostridium botulinum biovar A		3.51	0.13		1	99	
45001402	**Good identification**						**45001402**
Atopobium minutum		100.0	0.39	ARA 1			
45001520	**Very good identification**						**45001520**
Odoribacter splanchnicus		100.0	0.61	IND 99			
45001560	**Good identification**						**45001560**
Odoribacter splanchnicus		100.0	0.45	IND 99			
45002000	**Low discrimination**						**45002000**
					SPO		
Eubacterium limosum		79.35	0.66	FRU 99	1		
Clostridium difficile		12.82	0.54		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		3.27	0.47		99		
45002002	**Low discrimination**						**45002002**
					SPO		
Atopobium minutum		53.24	0.47	MAN 1	1		
Eubacterium limosum		34.26	0.32		1		
Clostridium difficile		5.54	0.20		99		
45002010	**Very good identification**						**45002010**
					SPO		
Clostridium difficile		98.60	0.66	FRU 99	99		
Eubacterium limosum		1.26	0.32		1		
Clostridium botulinum biovar A		0.05	0.13		99		
45002022	**Good identification**						**45002022**
					GRA		
Atopobium minutum		95.69	0.39	MAN 1	99		
Pseudoflavonifractor capillosus		2.44	-.03		1		
Eubacterium limosum		1.87	-.03		99		
45002040	**Low discrimination**						**45002040**
					SPO		
Clostridium botulinum biovar A		37.92	0.38	MNS 7	99		
Eubacterium limosum		31.12	0.32	MAN 1	1		
Clostridium difficile		26.21	0.32		99		
45002050	**Good identification**						**45002050**
Clostridium difficile		99.14	0.44	FRU 99			
Clostridium novyi biovar A		0.32	0.16	SOR 5			
Clostridium botulinum biovar A		0.30	0.04				
45003002	**Good identification**						**45003002**
Atopobium minutum		98.09	0.39	MAN 1			
Eubacterium limosum		1.91	-.03				
45004000	**Low discrimination**						**45004000**
					SPO		
Clostridium botulinum biovar A		37.38	0.47	MNS 7	99		
Clostridium bifermentans		25.52	0.43	XYL 1	99		
Atopobium minutum		21.37	0.50		1		
45004002	**Low discrimination**						**45004002**
					SPO		
Atopobium minutum		88.12	0.47	XYL 1	1		
Clostridium sporogenes		7.61	0.19		99		
Clostridium botulinum biovar A		2.34	0.13		99		
45004020	**Low discrimination**						**45004020**
					COC	SPO	
Atopobium minutum		72.46	0.42	XYL 1	1	1	
Clostridium bifermentans		16.57	0.23		1	99	
Gemella morbillorum		6.13	0.19		99	1	
45004022	**Good identification**						**45004022**
					GRA		
Atopobium minutum		97.51	0.39	XYL 1	99		
Pseudoflavonifractor capillosus		2.49	-.03		1		
45004040	**Low discrimination**						**45004040**
Clostridium glycolicum		82.35	0.49	MNS 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		15.62	0.38	FRU 90			
Clostridium bifermentans		1.49	0.20				
45005000	**Low discrimination**						**45005000**
					COC	SPO	
Atopobium minutum		63.88	0.42	XYL 1	1	1	
Gemella morbillorum		26.18	0.31		99	1	
Clostridium botulinum biovar A		3.39	0.13		1	99	
45005002	**Good identification**						**45005002**
Atopobium minutum		100.0	0.39	XYL 1			
45006010	**Good identification**						**45006010**
					SPO		
Clostridium difficile		99.76	0.44	FRU 99	99		
Eubacterium limosum		0.25	-.03	XYL 5	1		
45010000	**Low discrimination**						**45010000**
					SPO		
Eubacterium limosum		75.87	0.87	MAN 86	1		
Atopobium minutum		10.13	0.84		1		
Clostridium difficile		8.37	0.72		99		
45010001	**Low discrimination**						**45010001**
					SPO		
Eubacterium limosum		75.00	0.52	MAN 86	1		
Atopobium minutum		10.01	0.50	URE 1	1		
Clostridium difficile		8.27	0.38		99		
45010002	**Low discrimination**						**45010002**
					SPO		
Atopobium minutum		86.14	0.81		1		
Eubacterium limosum		9.78	0.52		1		
Clostridium sporogenes		1.42	0.40		99		
45010003	**Low discrimination**						**45010003**
					SPO		
Atopobium minutum		84.83	0.47	URE 1	1		
Eubacterium limosum		9.63	0.18		1		
Clostridium novyi biovar A		2.27	0.24		99		
45010010	**Very good identification**						**45010010**
					SPO		
Clostridium difficile		97.76	0.84	MAN 90	99		
Eubacterium limosum		1.83	0.52		1		
Atopobium minutum		0.25	0.50		1		
45010011	**Good identification**						**45010011**
					SPO		
Clostridium difficile		95.58	0.50	MAN 90	99		
Clostridium novyi biovar A		2.39	0.37	URE 1	99		
Eubacterium limosum		1.79	0.18		1		
45010012	**Low discrimination**						**45010012**
					SPO		
Clostridium difficile		83.76	0.50	MAN 90	99		
Atopobium minutum		13.84	0.47	BGL 1	1		
Eubacterium limosum		1.57	0.18		1		
45010020	**Low discrimination**						**45010020**
					SPO		
Atopobium minutum		75.88	0.76		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium limosum		17.22	0.52		1	
Clostridium difficile		1.90	0.38		99	
45010021	**Low discrimination**					**45010021**
					SPO	
Atopobium minutum		72.72	0.41	URE 1	1	
Eubacterium limosum		16.51	0.18		1	
Clostridium novyi biovar A		5.51	0.26		99	
45010022	**Very good identification**					**45010022**
					GRA	
Atopobium minutum		99.06	0.73		99	
Eubacterium limosum		0.34	0.18		99	
Pseudoflavonifractor capillosus		0.30	0.15		1	
45010023	**Good identification**					**45010023**
					SPO	
Atopobium minutum		98.03	0.38	URE 1	1	
Clostridium novyi biovar A		1.97	0.13		99	
45010030	**Low discrimination**					**45010030**
					SPO	
Clostridium difficile		89.87	0.50	MAN 90	99	
Atopobium minutum		7.43	0.41	NAG 1	1	
Eubacterium limosum		1.69	0.18		1	
45010032	**Low discrimination**					**45010032**
					SPO	
Atopobium minutum		83.10	0.38	MLZ 1	1	
Clostridium difficile		15.24	0.15		99	
Clostridium novyi biovar A		1.67	0.13		99	
45010040	**Low discrimination**					**45010040**
					SPO	
Eubacterium limosum		40.06	0.52	MAN 86	1	
Clostridium difficile		23.02	0.50	SOR 1	99	
Clostridium botulinum biovar A		19.13	0.51		99	
45010042	**Low discrimination**					**45010042**
					SPO	
Atopobium minutum		74.92	0.47	SOR 1	1	
Eubacterium limosum		8.50	0.18		1	
Clostridium difficile		4.89	0.16		99	
45010050	**Very good identification**					**45010050**
					SPO	
Clostridium difficile		99.00	0.62	MAN 90	99	
Eubacterium limosum		0.36	0.18	SOR 5	1	
Clostridium novyi biovar A		0.32	0.34		99	
45010060	**Low discrimination**					**45010060**
					CAT	
Atopobium minutum		51.57	0.41	SOR 1	1	
Eubacterium limosum		11.70	0.18		1	
Propionibacterium acnes		10.36	0.22		95	
45010062	**Good identification**					**45010062**
					SPO	
Atopobium minutum		97.40	0.38	SOR 1	1	
Clostridium novyi biovar A		1.30	0.10		99	
Pseudoramibacter alactolyticus		1.30	-.08		1	
45010100	**Low discrimination**					**45010100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		57.58	0.74		1		
Eubacterium limosum		32.38	0.52		99		
Atopobium minutum		4.32	0.50		99		
45010101	**Low discrimination**						**45010101**
					GRA		
Fusobacterium mortiferum		58.85	0.40	URE 1	1		
Eubacterium limosum		33.09	0.18		99		
Atopobium minutum		4.42	0.15		99		
45010102	**Low discrimination**						**45010102**
					GRA		
Fusobacterium mortiferum		66.02	0.58	BGL 10	1		
Atopobium minutum		29.74	0.47		99		
Eubacterium limosum		3.38	0.18		99		
45010110	**Acceptable identification**						**45010110**
					GRA	SPO	
Clostridium difficile		94.82	0.50	MAN 90	99	99	
Fusobacterium mortiferum		3.16	0.40	LAC 1	1	1	
Eubacterium limosum		1.78	0.18		99	1	
45010120	**Low discrimination**						**45010120**
					GRA		
Atopobium minutum		42.91	0.41	LAC 1	99		
Fusobacterium mortiferum		17.32	0.40		1		
Prevotella bivia		16.26	0.15		1		
45010122	**Acceptable identification**						**45010122**
					GRA		
Atopobium minutum		93.70	0.38	LAC 1	99		
Fusobacterium mortiferum		6.30	0.24		1		
45010140	**Low discrimination**						**45010140**
					GRA	SPO	
Fusobacterium mortiferum		42.44	0.40	SOR 1	1	1	
Eubacterium limosum		23.86	0.18		99	1	
Clostridium difficile		13.71	0.16		99	99	
45010200	**Low discrimination**						**45010200**
					SPO		
Eubacterium limosum		75.87	0.52	MAN 86	1		
Atopobium minutum		10.13	0.50	RHA 1	1		
Clostridium difficile		8.37	0.38		99		
45010202	**Low discrimination**						**45010202**
					SPO		
Atopobium minutum		86.80	0.47	RHA 1	1		
Eubacterium limosum		9.85	0.18		1		
Clostridium sporogenes		1.43	0.06		99		
45010210	**Good identification**						**45010210**
					SPO		
Clostridium difficile		97.92	0.50	MAN 90	99		
Eubacterium limosum		1.84	0.18	RHA 1	1		
Atopobium minutum		0.25	0.15		1		
45010220	**Low discrimination**						**45010220**
					SPO		
Atopobium minutum		76.96	0.41	RHA 1	1		
Eubacterium limosum		17.47	0.18		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium difficile		1.93	0.04		99	
45010222	**Good identification**					**45010222**
Atopobium minutum		100.0	0.38	RHA 1		
45010300	**Low discrimination**					**45010300**
					GRA	
Fusobacterium mortiferum		58.85	0.40	RHA 1	1	
Eubacterium limosum		33.09	0.18		99	
Atopobium minutum		4.42	0.15		99	
45010400	**Low discrimination**					**45010400**
					SPO	
Eubacterium limosum		73.63	0.52	MAN 86	1	
Atopobium minutum		9.83	0.50	ARA 1	1	
Clostridium difficile		8.12	0.38		99	
45010402	**Low discrimination**					**45010402**
					SPO	
Atopobium minutum		86.42	0.47	ARA 1	1	
Eubacterium limosum		9.81	0.18		1	
Clostridium sporogenes		1.43	0.06		99	
45010410	**Good identification**					**45010410**
					SPO	
Clostridium difficile		97.84	0.50	MAN 90	99	
Eubacterium limosum		1.84	0.18	ARA 1	1	
Atopobium minutum		0.25	0.15		1	
45010420	**Low discrimination**					**45010420**
					COC	
Atopobium minutum		70.30	0.41	ARA 1	1	
Eubacterium limosum		15.96	0.18		1	
Gemella morbillorum		7.87	0.20		99	
45010422	**Good identification**					**45010422**
Atopobium minutum		100.0	0.38	ARA 1		
45010500	**Low discrimination**					**45010500**
					GRA	
Fusobacterium mortiferum		56.27	0.40	ARA 1	1	
Eubacterium limosum		31.64	0.18		99	
Atopobium minutum		4.22	0.15		99	
45011000	**Low discrimination**					**45011000**
					GRA	
Atopobium minutum		63.24	0.76		99	
Fusobacterium mortiferum		16.41	0.71		1	
Eubacterium limosum		14.35	0.52		99	
45011001	**Low discrimination**					**45011001**
					GRA	
Atopobium minutum		62.72	0.41	URE 1	99	
Fusobacterium mortiferum		16.28	0.37		1	
Eubacterium limosum		14.23	0.18		99	
45011002	**Very good identification**					**45011002**
					GRA	
Atopobium minutum		95.14	0.73		99	
Fusobacterium mortiferum		4.12	0.55		1	
Eubacterium limosum		0.33	0.18		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45011003	**Good identification**						**45011003**
					GRA		
Atopobium minutum		95.85	0.38	URE 1	99		
Fusobacterium mortiferum		4.15	0.20		1		
45011010	**Low discrimination**						**45011010**
					GRA	SPO	
Clostridium difficile		88.11	0.50	GAL 1	99	99	
Atopobium minutum		7.28	0.41	MAN 90	99	1	
Fusobacterium mortiferum		1.89	0.37		1	1	
45011012	**Low discrimination**						**45011012**
					GRA	SPO	
Atopobium minutum		80.20	0.38	MLZ 1	99	1	
Clostridium difficile		14.70	0.15		99	99	
Fusobacterium mortiferum		3.47	0.20		1	1	
45011020	**Good identification**						**45011020**
					COC	CAT	
Atopobium minutum		93.03	0.67		1	1	
Propionibacterium acnes		2.58	0.34		1	95	
Gemella morbillorum		1.53	0.32		99	1	
45011022	**Very good identification**						**45011022**
					GRA		
Atopobium minutum		99.31	0.64		99		
Clostridium perfringens		0.21	0.13		99		
Fusobacterium mortiferum		0.13	0.20		1		
45011040	**Low discrimination**						**45011040**
					GRA	SPO	
Atopobium minutum		43.92	0.41	SOR 1	99	1	
Clostridium glycolicum		20.70	0.27		99	99	
Fusobacterium mortiferum		11.40	0.37		1	1	
45011042	**Good identification**						**45011042**
					GRA		
Atopobium minutum		95.85	0.38	SOR 1	99		
Fusobacterium mortiferum		4.15	0.20		1		
45011100	**Very good identification**						**45011100**
					GRA	SPO	
Fusobacterium mortiferum		97.23	0.77		1	1	
Atopobium minutum		1.62	0.41		99	1	
Clostridium septicum		0.37	0.18		99	99	
45011101	**Low discrimination**						**45011101**
					GRA	SPO	
Fusobacterium mortiferum		59.23	0.43	URE 1	1	1	
Clostridium chauvoei		39.78	0.29		99	99	
Atopobium minutum		0.99	0.07		99	1	
45011102	**Low discrimination**						**45011102**
					GRA		
Fusobacterium mortiferum		89.47	0.61	BGL 10	1		
Atopobium minutum		8.96	0.38		99		
Lactobacillus fermentum		1.44	0.32		99		
45011110	**Low discrimination**						**45011110**
					GRA	SPO	
Fusobacterium mortiferum		81.15	0.43	MLZ 1	1	1	
Clostridium difficile		16.38	0.15		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		1.35	0.07		99	1	
45011120	**Low discrimination**						**45011120**
					GRA	SPO	
Clostridium septicum		84.92	0.53	MLT 94	99	99	
Odoribacter splanchnicus		7.97	0.32		1	1	
Clostridium perfringens		2.75	0.34		99	84	
45011140	**Good identification**						**45011140**
					GRA		
Fusobacterium mortiferum		98.36	0.43	SOR 1	1		
Atopobium minutum		1.64	0.07		99		
45011200	**Low discrimination**						**45011200**
					GRA		
Atopobium minutum		63.57	0.41	RHA 1	99		
Fusobacterium mortiferum		16.50	0.37		1		
Eubacterium limosum		14.43	0.18		99		
45011202	**Good identification**						**45011202**
					GRA		
Atopobium minutum		95.85	0.38	RHA 1	99		
Fusobacterium mortiferum		4.15	0.20		1		
45011300	**Good identification**						**45011300**
					GRA		
Fusobacterium mortiferum		98.36	0.43	RHA 1	1		
Atopobium minutum		1.64	0.07		99		
45011400	**Low discrimination**						**45011400**
					COC	GRA	
Atopobium minutum		46.70	0.41	ARA 1	1	99	
Gemella morbillorum		25.35	0.32		99	99	
Fusobacterium mortiferum		12.12	0.37		1	1	
45011402	**Low discrimination**						**45011402**
					GRA		
Atopobium minutum		87.59	0.38	ARA 1	99		
Lactobacillus fermentum		8.62	0.28		99		
Fusobacterium mortiferum		3.79	0.20		1		
45011520	**Very good identification**						**45011520**
					GRA	SPO	
Odoribacter splanchnicus		99.87	0.66	IND 99	1	1	
Clostridium septicum		0.11	0.18		99	99	
Lactobacillus fermentum		0.02	0.24		99	1	
45011560	**Good identification**						**45011560**
Odoribacter splanchnicus		100.0	0.50	IND 99			
45012000	**Low discrimination**						**45012000**
					SPO		
Eubacterium limosum		86.06	1.00		1		
Clostridium difficile		13.91	0.88		99		
Atopobium minutum		0.02	0.50		1		
45012001	**Low discrimination**						**45012001**
					SPO		
Eubacterium limosum		86.07	0.66	URE 1	1		
Clostridium difficile		13.91	0.54		99		
Atopobium minutum		0.02	0.15		1		
45012002	**Low discrimination**						**45012002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Eubacterium limosum		84.89	0.66	BGL 1	1		
Clostridium difficile		13.72	0.54		99		
Atopobium minutum		1.23	0.47		1		
45012010	**Very good identification**						**45012010**
					SPO	CAT	
Clostridium difficile		98.74	1.00		99	1	
Eubacterium limosum		1.26	0.66		1	1	
Propionibacterium granulosum		0.00	0.20		1	90	
45012011	**Very good identification**						**45012011**
					SPO		
Clostridium difficile		98.74	0.66	URE 1	99		
Eubacterium limosum		1.26	0.32		1		
45012012	**Very good identification**						**45012012**
					SPO		
Clostridium difficile		98.71	0.66	BGL 1	99		
Eubacterium limosum		1.26	0.32		1		
Atopobium minutum		0.02	0.12		1		
45012020	**Low discrimination**						**45012020**
					SPO		
Eubacterium limosum		85.13	0.66	NAG 1	1		
Clostridium difficile		13.76	0.54		99		
Atopobium minutum		0.62	0.41		1		
45012030	**Very good identification**						**45012030**
					SPO	CAT	
Clostridium difficile		98.72	0.66	NAG 1	99	1	
Eubacterium limosum		1.26	0.32		1	1	
Propionibacterium granulosum		0.01	0.09		1	90	
45012040	**Low discrimination**						**45012040**
					SPO		
Eubacterium limosum		54.17	0.66	SOR 1	1		
Clostridium difficile		45.61	0.66		99		
Pseudoramibacter alactolyticus		0.10	0.17		1		
45012050	**Very good identification**						**45012050**
					SPO		
Clostridium difficile		99.76	0.78	SOR 5	99		
Eubacterium limosum		0.25	0.32		1		
45012051	**Good identification**						**45012051**
					SPO		
Clostridium difficile		99.76	0.44	SOR 5	99		
Eubacterium limosum		0.25	-.03	URE 1	1		
45012052	**Good identification**						**45012052**
					SPO		
Clostridium difficile		99.76	0.44	SOR 5	99		
Eubacterium limosum		0.25	-.03	BGL 1	1		
45012070	**Good identification**						**45012070**
					SPO		
Clostridium difficile		99.76	0.44	NAG 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		
45012100	**Low discrimination**						**45012100**
					GRA	SPO	
Eubacterium limosum		85.85	0.66	LAC 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		13.87	0.54		99	99	
Fusobacterium mortiferum		0.25	0.40		1	1	
45012110	**Very good identification**						**45012110**
					SPO		
Clostridium difficile		98.74	0.66	LAC 1	99		
Eubacterium limosum		1.26	0.32		1		
45012150	**Good identification**						**45012150**
					SPO		
Clostridium difficile		99.76	0.44	LAC 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		
45012200	**Low discrimination**						**45012200**
					SPO		
Eubacterium limosum		86.06	0.66	RHA 1	1		
Clostridium difficile		13.91	0.54		99		
Atopobium minutum		0.02	0.15		1		
45012210	**Very good identification**						**45012210**
					SPO		
Clostridium difficile		98.74	0.66	RHA 1	99		
Eubacterium limosum		1.26	0.32		1		
45012250	**Good identification**						**45012250**
					SPO		
Clostridium difficile		99.76	0.44	RHA 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		
45012400	**Low discrimination**						**45012400**
					SPO		
Eubacterium limosum		86.06	0.66	ARA 1	1		
Clostridium difficile		13.91	0.54		99		
Atopobium minutum		0.02	0.15		1		
45012410	**Very good identification**						**45012410**
					SPO		
Clostridium difficile		98.74	0.66	ARA 1	99		
Eubacterium limosum		1.26	0.32		1		
45012450	**Good identification**						**45012450**
					SPO		
Clostridium difficile		99.76	0.44	ARA 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		
45013000	**Low discrimination**						**45013000**
					SPO		
Eubacterium limosum		85.19	0.66	GAL 1	1		
Clostridium difficile		13.77	0.54		99		
Atopobium minutum		0.62	0.41		1		
45013010	**Very good identification**						**45013010**
					SPO	CAT	
Clostridium difficile		98.69	0.66	GAL 1	99	1	
Eubacterium limosum		1.26	0.32		1	1	
Propionibacterium granulosum		0.04	0.20		1	90	
45013050	**Good identification**						**45013050**
					SPO		
Clostridium difficile		99.76	0.44	GAL 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		
45014000	**Low discrimination**						**45014000**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		46.47	0.52	MAN 86	1		
Clostridium difficile		26.71	0.50	XYL 1	99		
Clostridium glycolicum		8.77	0.43		99		
45014002	**Low discrimination**						**45014002**
					GRA		
Atopobium minutum		76.51	0.47	XYL 1	99		
Eubacterium limosum		8.68	0.18		99		
Fusobacterium mortiferum		5.48	0.32		1		
45014010	**Very good identification**						**45014010**
					SPO		
Clostridium difficile		99.45	0.62	MAN 90	99		
Eubacterium limosum		0.36	0.18	XYL 5	1		
Clostridium glycolicum		0.07	0.09		99		
45014020	**Low discrimination**						**45014020**
					SPO		
Atopobium minutum		62.90	0.41	XYL 1	1		
Eubacterium limosum		14.28	0.18		1		
Clostridium difficile		8.21	0.16		99		
45014022	**Good identification**						**45014022**
Atopobium minutum		100.0	0.38	XYL 1			
45014040	**Very good identification**						**45014040**
					SPO		
Clostridium glycolicum		98.65	0.65	MNS 1	99		
Clostridium difficile		0.83	0.28		99		
Eubacterium limosum		0.28	0.18		1		
45014050	**Low discrimination**						**45014050**
Clostridium difficile		80.30	0.40	MAN 90			
Clostridium glycolicum		19.70	0.31	XYL 5			
45014100	**Low discrimination**						**45014100**
					GRA	SPO	
Fusobacterium mortiferum		86.04	0.55	XYL 7	1	1	
Eubacterium limosum		6.49	0.18		99	1	
Clostridium difficile		3.73	0.16		99	99	
45014102	**Acceptable identification**						**45014102**
					GRA		
Fusobacterium mortiferum		94.30	0.39	XYL 7	1		
Atopobium minutum		5.70	0.12	BGL 10	99		
45015000	**Low discrimination**						**45015000**
					COC	GRA	
Fusobacterium mortiferum		44.24	0.52	XYL 7	1	1	
Atopobium minutum		22.88	0.41		1	99	
Gemella morbillorum		12.42	0.32		99	99	
45015002	**Low discrimination**						**45015002**
					GRA		
Atopobium minutum		72.46	0.38	XYL 1	99		
Fusobacterium mortiferum		23.35	0.35		1		
Lactobacillus fermentum		4.19	0.24		99		
45015040	**Good identification**						**45015040**
					GRA	SPO	
Clostridium glycolicum		99.67	0.49	MNS 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.22	0.18	GAL 10	1	1	
Atopobium minutum		0.11	0.07		99	1	
45015100	**Very good identification**						**45015100**
					GRA		
Fusobacterium mortiferum		97.59	0.58	XYL 7	1		
Lactobacillus fermentum		1.36	0.28		99		
Actinomyces meyeri		0.66	0.12		99		
45015102	**Low discrimination**						**45015102**
					GRA		
Fusobacterium mortiferum		88.85	0.42	XYL 7	1		
Lactobacillus fermentum		11.15	0.28	BGL 10	99		
45016000	**Low discrimination**						**45016000**
					SPO		
Eubacterium limosum		54.26	0.66	XYL 1	1		
Clostridium difficile		45.69	0.66		99		
Clostridium glycolicum		0.02	0.09		99		
45016010	**Very good identification**						**45016010**
					SPO		
Clostridium difficile		99.76	0.78	XYL 5	99		
Eubacterium limosum		0.25	0.32		1		
45016011	**Good identification**						**45016011**
					SPO		
Clostridium difficile		99.76	0.44	XYL 5	99		
Eubacterium limosum		0.25	-.03	URE 1	1		
45016012	**Good identification**						**45016012**
					SPO		
Clostridium difficile		99.76	0.44	XYL 5	99		
Eubacterium limosum		0.25	-.03	BGL 1	1		
45016030	**Good identification**						**45016030**
					SPO		
Clostridium difficile		99.76	0.44	XYL 5	99		
Eubacterium limosum		0.25	-.03	NAG 1	1		
45016040	**Low discrimination**						**45016040**
					SPO		
Clostridium difficile		73.47	0.44	XYL 5	99		
Eubacterium limosum		16.75	0.32	SOR 5	1		
Clostridium glycolicum		9.78	0.31		99		
45016050	**Very good identification**						**45016050**
					SPO		
Clostridium difficile		99.93	0.56	XYL 5	99		
Eubacterium limosum		0.05	-.03	SOR 5	1		
Clostridium glycolicum		0.03	-.03		99		
45016110	**Good identification**						**45016110**
					SPO		
Clostridium difficile		99.76	0.44	XYL 5	99		
Eubacterium limosum		0.25	-.03	LAC 1	1		
45016210	**Good identification**						**45016210**
					SPO		
Clostridium difficile		99.76	0.44	XYL 5	99		
Eubacterium limosum		0.25	-.03	RHA 1	1		
45016410	**Good identification**						**45016410**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		99.76	0.44	XYL 5	99		
Eubacterium limosum		0.25	-.03	ARA 1	1		
45017010	**Good identification**						**45017010**
					SPO		
Clostridium difficile		99.76	0.44	GAL 1	99		
Eubacterium limosum		0.25	-.03	XYL 5	1		
45020000	**Low discrimination**						**45020000**
Clostridium botulinum biovar A		84.23	0.71	MNS 7			
Clostridium sporogenes		11.08	0.53				
Clostridium bifermentans		2.32	0.43				
45020001	**Low discrimination**						**45020001**
Clostridium botulinum biovar A		82.39	0.36	MNS 7			
Clostridium sporogenes		10.84	0.19	URE 1			
Clostridium bifermentans		2.27	0.09				
45020002	**Low discrimination**						**45020002**
					SPO		
Clostridium sporogenes		62.99	0.45	MNS 1	99		
Atopobium minutum		22.11	0.47		1		
Clostridium botulinum biovar A		14.51	0.36		99		
45020010	**Low discrimination**						**45020010**
Clostridium botulinum biovar A		54.28	0.36	MNS 7			
Clostridium difficile		34.78	0.30	MLZ 1			
Clostridium sporogenes		7.14	0.19				
45020020	**Low discrimination**						**45020020**
					SPO		
Clostridium botulinum biovar A		46.98	0.36	MNS 7	99		
Atopobium minutum		35.80	0.42	NAG 1	1		
Clostridium bifermentans		8.19	0.23		99		
45020022	**Low discrimination**						**45020022**
					GRA	SPO	
Atopobium minutum		88.36	0.39	TRE 1	99	1	
Clostridium sporogenes		7.63	0.11		99	99	
Pseudoflavonifractor capillosus		2.25	-.03		1	1	
45020040	**Very good identification**						**45020040**
Clostridium botulinum biovar A		98.96	0.62	MNS 7			
Clostridium sporogenes		0.44	0.19				
Clostridium bifermentans		0.38	0.20				
45020100	**Low discrimination**						**45020100**
					GRA	SPO	
Clostridium botulinum biovar A		78.92	0.36	MNS 7	99	99	
Clostridium sporogenes		10.39	0.19	LAC 1	99	99	
Fusobacterium mortiferum		6.69	0.31		1	1	
45020200	**Low discrimination**						**45020200**
Clostridium botulinum biovar A		84.58	0.36	MNS 7			
Clostridium sporogenes		11.13	0.19	RHA 1			
Clostridium bifermentans		2.33	0.09				
45020400	**Low discrimination**						**45020400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		84.17	0.36	MNS 7			
Clostridium sporogenes		11.08	0.19	ARA 1			
Clostridium bifermentans		2.32	0.09				
45021000	**Low discrimination**						**45021000**
					SPO		
Clostridium botulinum biovar A		49.69	0.36	MNS 7	99		
Atopobium minutum		37.87	0.42	GAL 1	1		
Clostridium sporogenes		6.54	0.19		99		
45021002	**Low discrimination**						**45021002**
					SPO		
Atopobium minutum		89.43	0.39	TRE 1	1		
Clostridium sporogenes		7.72	0.11		99		
Clostridium botulinum biovar A		1.78	0.02		99		
45022010	**Good identification**						**45022010**
					SPO		
Clostridium difficile		99.65	0.47	FRU 99	99		
Clostridium botulinum biovar A		0.18	0.02	TRE 7	99		
Eubacterium limosum		0.17	-.03		1		
45024000	**Low discrimination**						**45024000**
Clostridium botulinum biovar A		83.68	0.36	MNS 7			
Clostridium sporogenes		11.01	0.19	XYL 1			
Clostridium bifermentans		2.31	0.09				
45030000	**Low discrimination**						**45030000**
					SPO		
Eubacterium limosum		35.38	0.52	TRE 1	1		
Clostridium difficile		29.08	0.53	MAN 86	99		
Clostridium botulinum biovar A		14.14	0.50		99		
45030002	**Low discrimination**						**45030002**
					GRA	SPO	
Atopobium minutum		44.86	0.47	TRE 1	99	1	
Clostridium sporogenes		24.43	0.32		99	99	
Fusobacterium mortiferum		12.75	0.43		1	1	
45030010	**Very good identification**						**45030010**
					SPO	CAT	
Clostridium difficile		99.35	0.64	TRE 7	99	1	
Eubacterium limosum		0.25	0.18	MAN 90	1	1	
Propionibacterium granulosum		0.18	0.30		1	90	
45030020	**Low discrimination**						**45030020**
					CAT		
Atopobium minutum		50.10	0.41	TRE 1	1		
Propionibacterium granulosum		14.87	0.35		90		
Eubacterium limosum		11.37	0.18		1		
45030022	**Low discrimination**						**45030022**
					SPO	CAT	
Atopobium minutum		88.67	0.38	TRE 1	1	1	
Propionibacterium granulosum		9.87	0.24		1	90	
Clostridium sporogenes		1.46	-.02		99	1	
45030040	**Low discrimination**						**45030040**
					SPO		
Clostridium botulinum biovar A		64.13	0.41	MNS 7	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		23.23	0.31	FRU 6	99			
Eubacterium limosum		5.42	0.18		1			
45030050	**Good identification**							**45030050**
Clostridium difficile		98.18	0.43	TRE 7				
Clostridium botulinum biovar B		1.26	0.10	MAN 90				
Clostridium botulinum biovar A		0.56	0.07	SOR 5				
45030100	**Very good identification**							**45030100**
					GRA	SPO		
Fusobacterium mortiferum		95.37	0.65		1	1		
Eubacterium limosum		1.81	0.18		99	1		
Clostridium difficile		1.49	0.19		99	99		
45030102	**Good identification**							**45030102**
					GRA	SPO		
Fusobacterium mortiferum		97.70	0.49	BGL 10	1	1		
Atopobium minutum		1.49	0.12		99	1		
Clostridium sporogenes		0.81	-.02		99	99		
45031000	**Low discrimination**							**45031000**
					GRA	CAT		
Fusobacterium mortiferum		72.90	0.62		1	5		
Propionibacterium granulosum		11.28	0.45		99	90		
Atopobium minutum		9.50	0.41		99	1		
45031002	**Low discrimination**							**45031002**
					GRA	CAT		
Fusobacterium mortiferum		45.36	0.46	BGL 10	1	5		
Atopobium minutum		35.46	0.38		99	1		
Propionibacterium granulosum		15.79	0.35		99	90		
45031100	**Good identification**							**45031100**
					COC	GRA	SPO	
Fusobacterium mortiferum		93.81	0.68		1	1	1	
Clostridium septicum		3.57	0.27		1	99	99	
Atopobium parvulum		2.29	0.29		99	99	1	
45031102	**Acceptable identification**							**45031102**
					COC	GRA		
Fusobacterium mortiferum		94.74	0.52	BGL 10	1	1		
Atopobium parvulum		3.66	0.16		99	99		
Clostridium perfringens		0.94	0.07		1	99		
45031120	**Very good identification**							**45031120**
					GRA	SPO		
Clostridium septicum		97.64	0.61	MLT 94	99	99		
Clostridium perfringens		1.96	0.38		99	84		
Fusobacterium mortiferum		0.26	0.34		1	1		
45032000	**Low discrimination**							**45032000**
					SPO	CAT		
Clostridium difficile		54.43	0.69	TRE 7	99	1		
Eubacterium limosum		45.20	0.66		1	1		
Propionibacterium granulosum		0.29	0.43		1	90		
45032010	**Very good identification**							**45032010**
					SPO	CAT		
Clostridium difficile		99.81	0.81	TRE 7	99	1		
Eubacterium limosum		0.17	0.32		1	1		
Propionibacterium granulosum		0.02	0.28		1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45032011	**Good identification**							**45032011**
					SPO			
Clostridium difficile		99.83	0.47	TRE 7	99			
Eubacterium limosum		0.17	-.03	URE 1	1			
45032012	**Good identification**							**45032012**
					SPO	CAT		
Clostridium difficile		99.46	0.47	TRE 7	99	1		
Propionibacterium granulosum		0.37	0.18	BGL 1	1	90		
Eubacterium limosum		0.17	-.03		1	1		
45032030	**Good identification**							**45032030**
					SPO	CAT		
Clostridium difficile		99.46	0.47	TRE 7	99	1		
Propionibacterium granulosum		0.37	0.18	NAG 1	1	90		
Eubacterium limosum		0.17	-.03		1	1		
45032040	**Low discrimination**							**45032040**
					SPO			
Clostridium difficile		85.95	0.47	TRE 7	99			
Eubacterium limosum		13.70	0.32	SOR 5	1			
Clostridium botulinum biovar A		0.27	0.07		99			
45032050	**Very good identification**							**45032050**
					SPO			
Clostridium difficile		99.97	0.59	TRE 7	99			
Eubacterium limosum		0.03	-.03	SOR 5	1			
45032110	**Good identification**							**45032110**
					SPO			
Clostridium difficile		99.83	0.47	TRE 7	99			
Eubacterium limosum		0.17	-.03	LAC 1	1			
45032210	**Good identification**							**45032210**
					SPO			
Clostridium difficile		99.83	0.47	TRE 7	99			
Eubacterium limosum		0.17	-.03	RHA 1	1			
45032410	**Good identification**							**45032410**
					SPO			
Clostridium difficile		99.83	0.47	TRE 7	99			
Eubacterium limosum		0.17	-.03	ARA 1	1			
45033010	**Good identification**							**45033010**
					SPO	CAT		
Clostridium difficile		98.37	0.47	TRE 7	99	1		
Propionibacterium granulosum		1.47	0.28	GAL 1	1	90		
Eubacterium limosum		0.17	-.03		1	1		
45034010	**Good identification**							**45034010**
Clostridium difficile		100.0	0.43	TRE 7				
45034100	**Good identification**							**45034100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.04	0.46	XYL 7	1	1	1	
Clostridium difficile		1.07	-.03		1	99	99	
Atopobium parvulum		0.89	0.00		99	99	1	
45035000	**Acceptable identification**							**45035000**
					GRA	CAT		
Fusobacterium mortiferum		93.51	0.43	XYL 7	1	5		
Propionibacterium granulosum		1.94	0.11		99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium minutum		1.64	0.07		99	1		
45035100	**Low discrimination**							**45035100**
					COC	GRA		
Fusobacterium mortiferum		62.47	0.49	XYL 7	1	1		
Atopobium parvulum		37.53	0.34		99	99		
45035122	**Low discrimination**							**45035122**
					COC	GRA	SPO	
Parabacteroides distasonis		89.64	0.37	RAF 99	1	1	1	
Atopobium parvulum		6.68	0.08		99	99	1	
Clostridium septicum		3.68	-.02		1	99	99	
45035322	**Good identification**							**45035322**
Parabacteroides distasonis		100.0	0.46	RAF 99				
45035332	**Good identification**							**45035332**
Parabacteroides distasonis		100.0	0.41	RAF 99				
45035722	**Good identification**							**45035722**
Parabacteroides distasonis		100.0	0.42	RAF 99				
45035732	**Good identification**							**45035732**
Parabacteroides distasonis		100.0	0.37	RAF 99				
45036000	**Low discrimination**							**45036000**
					SPO	CAT		
Clostridium difficile		86.18	0.47	TRE 7	99	1		
Eubacterium limosum		13.74	0.32	XYL 5	1	1		
Propionibacterium granulosum		0.09	0.09		1	90		
45036010	**Very good identification**							**45036010**
					SPO			
Clostridium difficile		99.97	0.59	TRE 7	99			
Eubacterium limosum		0.03	-.03	XYL 5	1			
45036050	**Good identification**							**45036050**
Clostridium difficile		100.0	0.37	TRE 7				
45040000	**Low discrimination**							**45040000**
					SPO			
Clostridium botulinum biovar A		40.59	0.47	MNS 7	99			
Clostridium bifermentans		27.72	0.43	CEL 1	99			
Atopobium minutum		23.20	0.50		1			
45040002	**Low discrimination**							**45040002**
					SPO			
Atopobium minutum		88.42	0.47	CEL 1	1			
Clostridium sporogenes		7.63	0.19		99			
Clostridium botulinum biovar A		2.34	0.13		99			
45040020	**Low discrimination**							**45040020**
					SPO			
Atopobium minutum		76.73	0.42	CEL 1	1			
Clostridium bifermentans		17.55	0.23		99			
Clostridium botulinum biovar A		4.07	0.13		99			
45040022	**Good identification**							**45040022**
					GRA			
Atopobium minutum		97.51	0.39	CEL 1	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Pseudoflavonifractor capillosus		2.49	-.03		1		
45040040	**Low discrimination**						**45040040**
					SPO		
Clostridium botulinum biovar A		88.86	0.38	MNS 7	99		
Clostridium bifermentans		8.46	0.20	CEL 1	99		
Atopobium minutum		1.72	0.16		1		
45041000	**Low discrimination**						**45041000**
					GRA	SPO	
Atopobium minutum		82.90	0.42	CEL 1	99	1	
Fusobacterium mortiferum		6.62	0.29		1	1	
Clostridium botulinum biovar A		4.40	0.13		99	99	
45041002	**Good identification**						**45041002**
					GRA		
Atopobium minutum		98.69	0.39	CEL 1	99		
Fusobacterium mortiferum		1.31	0.12		1		
45042010	**Good identification**						**45042010**
					SPO		
Clostridium difficile		99.76	0.44	FRU 99	99		
Eubacterium limosum		0.25	-.03	CEL 5	1		
45050000	**Low discrimination**						**45050000**
					GRA	SPO	
Eubacterium limosum		49.31	0.52	CEL 1	99	1	
Clostridium difficile		28.34	0.50	MAN 86	99	99	
Fusobacterium mortiferum		12.53	0.60		1	1	
45050002	**Low discrimination**						**45050002**
					GRA		
Atopobium minutum		66.12	0.47	CEL 1	99		
Fusobacterium mortiferum		20.97	0.43		1		
Eubacterium limosum		7.50	0.18		99		
45050010	**Very good identification**						**45050010**
					GRA	SPO	
Clostridium difficile		99.50	0.62	CEL 5	99	99	
Eubacterium limosum		0.36	0.18	MAN 90	99	1	
Fusobacterium mortiferum		0.09	0.26		1	1	
45050020	**Low discrimination**						**45050020**
					SPO		
Atopobium minutum		68.38	0.41	CEL 1	1		
Eubacterium limosum		15.52	0.18		1		
Clostridium difficile		8.92	0.16		99		
45050022	**Good identification**						**45050022**
Atopobium minutum		100.0	0.38	CEL 1			
45050050	**Good identification**						**45050050**
Clostridium difficile		100.0	0.40	CEL 5			
45050100	**Very good identification**						**45050100**
					GRA	SPO	
Fusobacterium mortiferum		97.17	0.66		1	1	
Eubacterium limosum		1.66	0.18		99	1	
Clostridium difficile		0.95	0.16		99	99	
45050102	**Good identification**						**45050102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		98.65	0.50	BGL 10	1		
Atopobium minutum		1.35	0.12		99		
45051000	**Low discrimination**						**45051000**
					GRA		
Fusobacterium mortiferum		80.93	0.63		1		
Atopobium minutum		9.45	0.41		99		
Lactobacillus jensenii		4.74	0.32		99		
45051002	**Low discrimination**						**45051002**
					GRA		
Fusobacterium mortiferum		45.65	0.46	BGL 10	1		
Atopobium minutum		31.98	0.38		99		
Lactobacillus jensenii		16.02	0.29		99		
45051100	**Very good identification**						**45051100**
					GRA	SPO	
Fusobacterium mortiferum		95.90	0.69		1	1	
Clostridium septicum		3.45	0.27		99	99	
Lactobacillus catenaformis		0.27	0.15		99	1	
45051102	**Acceptable identification**						**45051102**
					GRA	SPO	
Fusobacterium mortiferum		93.38	0.53	BGL 10	1	1	
Clostridium barati		2.88	0.12		99	99	
Lactobacillus catenaformis		2.32	0.15		99	1	
45051120	**Very good identification**						**45051120**
					GRA	SPO	
Clostridium septicum		99.24	0.61	MLT 94	99	99	
Fusobacterium mortiferum		0.28	0.35		1	1	
Clostridium paraputrificum		0.25	0.14		99	99	
45052000	**Low discrimination**						**45052000**
					SPO		
Eubacterium limosum		54.19	0.66	CEL 1	1		
Clostridium difficile		45.63	0.66		99		
Clostridium innocuum		0.15	0.27		99		
45052010	**Very good identification**						**45052010**
					SPO		
Clostridium difficile		99.75	0.78	CEL 5	99		
Eubacterium limosum		0.25	0.32		1		
Clostridium innocuum		0.00	0.04		99		
45052011	**Good identification**						**45052011**
					SPO		
Clostridium difficile		99.76	0.44	CEL 5	99		
Eubacterium limosum		0.25	-.03	URE 1	1		
45052012	**Good identification**						**45052012**
					SPO		
Clostridium difficile		99.48	0.44	CEL 5	99		
Clostridium innocuum		0.27	0.04	BGL 1	99		
Eubacterium limosum		0.24	-.03		1		
45052030	**Good identification**						**45052030**
					SPO		
Clostridium difficile		99.76	0.44	CEL 5	99		
Eubacterium limosum		0.25	-.03	NAG 1	1		
45052040	**Low discrimination**						**45052040**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium difficile		81.44	0.44	CEL 5	99	
Eubacterium limosum		18.56	0.32	SOR 5	1	
45052050	**Very good identification**					**45052050**
					SPO	
Clostridium difficile		99.95	0.56	CEL 5	99	
Eubacterium limosum		0.05	-.03	SOR 5	1	
45052110	**Good identification**					**45052110**
					SPO	
Clostridium difficile		99.76	0.44	CEL 5	99	
Eubacterium limosum		0.25	-.03	LAC 1	1	
45052210	**Good identification**					**45052210**
					SPO	
Clostridium difficile		99.76	0.44	CEL 5	99	
Eubacterium limosum		0.25	-.03	RHA 1	1	
45052410	**Good identification**					**45052410**
					SPO	
Clostridium difficile		99.76	0.44	CEL 5	99	
Eubacterium limosum		0.25	-.03	ARA 1	1	
45053000	**Low discrimination**					**45053000**
					SPO	
Clostridium innocuum		73.01	0.49	SAL 99	99	
Eubacterium limosum		14.12	0.32		1	
Clostridium difficile		11.89	0.32		99	
45053002	**Good identification**					**45053002**
					SPO	
Clostridium innocuum		99.51	0.49	SAL 99	99	
Eubacterium limosum		0.19	-.03		1	
Clostridium difficile		0.16	-.02		99	
45053010	**Acceptable identification**					**45053010**
					SPO	
Clostridium difficile		94.80	0.44	CEL 5	99	
Clostridium innocuum		4.97	0.25	GAL 1	99	
Eubacterium limosum		0.23	-.03		1	
45053400	**Good identification**					**45053400**
					SPO	
Clostridium innocuum		98.00	0.36	SAL 99	99	
Eubacterium limosum		1.09	-.03		1	
Clostridium difficile		0.91	-.02		99	
45053402	**Good identification**					**45053402**
Clostridium innocuum		100.0	0.36	SAL 99		
45054010	**Good identification**					**45054010**
Clostridium difficile		100.0	0.40	CEL 5		
45054100	**Good identification**					**45054100**
Fusobacterium mortiferum		100.0	0.47	XYL 7		
45055000	**Good identification**					**45055000**
					GRA	
Fusobacterium mortiferum		98.46	0.44	XYL 7	1	
Atopobium minutum		1.54	0.07		99	
45055100	**Good identification**					**45055100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Fusobacterium mortiferum		98.75	0.50	XYL 7	1	1		
Atopobium parvulum		1.25	0.06		99	99		
45056000	**Low discrimination**							**45056000**
					SPO			
Clostridium difficile		81.22	0.44	CEL 5	99			
Eubacterium limosum		18.51	0.32	XYL 5	1			
Clostridium innocuum		0.27	0.05		99			
45056010	**Very good identification**							**45056010**
					SPO			
Clostridium difficile		99.95	0.56	CEL 5	99			
Eubacterium limosum		0.05	-.03	XYL 5	1			
45060000	**Low discrimination**							**45060000**
Clostridium botulinum biovar A		83.24	0.36	MNS 7				
Clostridium sporogenes		10.95	0.19	CEL 1				
Clostridium bifermentans		2.30	0.09					
45061120	**Good identification**							**45061120**
Clostridium septicum		100.0	0.35	MLT 94				
45070000	**Low discrimination**							**45070000**
					GRA	SPO		
Fusobacterium mortiferum		55.27	0.51		1	1		
Clostridium difficile		31.50	0.31		99	99		
Eubacterium limosum		7.36	0.18		99	1		
45070010	**Good identification**							**45070010**
					GRA	SPO		
Clostridium difficile		99.64	0.43	TRE 7	99	99		
Fusobacterium mortiferum		0.36	0.17	CEL 5	1	1		
45070100	**Very good identification**							**45070100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.96	0.57		1	1	1	
Clostridium septicum		0.54	0.01		1	99	99	
Atopobium parvulum		0.26	0.01		99	99	1	
45070102	**Good identification**							**45070102**
Fusobacterium mortiferum		100.0	0.41	BGL 10				
45070120	**Good identification**							**45070120**
					GRA	SPO		
Clostridium septicum		98.17	0.35	MLT 94	99	99		
Fusobacterium mortiferum		1.83	0.23	GAL 99	1	1		
45071000	**Low discrimination**							**45071000**
					GRA	SPO		
Fusobacterium mortiferum		61.63	0.54		1	1		
Lactobacillus jensenii		36.21	0.40		99	1		
Clostridium innocuum		0.62	0.07		99	99		
45071002	**Low discrimination**							**45071002**
					GRA	SPO		
Lactobacillus jensenii		76.39	0.37	SUC 90	99	1		
Fusobacterium mortiferum		21.67	0.38	MLT 95	1	1		
Clostridium innocuum		1.95	0.07	SAL 95	99	99		
45071020	**Low discrimination**							**45071020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Clostridium septicum		84.30	0.35	MLT 94	99	99		
Lactobacillus jensenii		14.69	0.30	LAC 99	99	1		
Fusobacterium mortiferum		1.01	0.20		1	1		
45071100	**Low discrimination**							**45071100**
					COC	GRA	SPO	
Fusobacterium mortiferum		65.18	0.60		1	1	1	
Clostridium septicum		23.54	0.35		1	99	99	
Atopobium parvulum		11.11	0.36		99	99	1	
45071102	**Low discrimination**							**45071102**
					COC	GRA	SPO	
Fusobacterium mortiferum		72.32	0.44	BGL 10	1	1	1	
Atopobium parvulum		19.58	0.23		99	99	1	
Clostridium septicum		4.80	0.06		1	99	99	
45071120	**Very good identification**							**45071120**
					COC	GRA	SPO	
Clostridium septicum		99.84	0.69	MLT 94	1	99	99	
Atopobium parvulum		0.08	0.23		99	99	1	
Fusobacterium mortiferum		0.03	0.26		1	1	1	
45071121	**Good identification**							**45071121**
Clostridium septicum		100.0	0.35	MLT 94				
45071122	**Good identification**							**45071122**
					COC	SPO		
Clostridium septicum		96.46	0.40	MLT 94	1	99		
Atopobium parvulum		0.70	0.10	BGL 2	99	1		
Clostridium perfringens		0.59	0.09		1	84		
45071130	**Good identification**							**45071130**
Clostridium septicum		100.0	0.35	MLT 94				
45071160	**Good identification**							**45071160**
Clostridium septicum		99.51	0.35	MLT 94				
Clostridium perfringens		0.50	0.02	SOR 1				
45071320	**Good identification**							**45071320**
Clostridium septicum		100.0	0.35	MLT 94				
45071520	**Good identification**							**45071520**
					GRA	SPO		
Clostridium septicum		99.02	0.35	MLT 94	99	99		
Odoribacter splanchnicus		0.98	-.03	ARA 1	1	1		
45072000	**Low discrimination**							**45072000**
					SPO			
Clostridium difficile		84.92	0.47	TRE 7	99			
Eubacterium limosum		13.53	0.32	CEL 5	1			
Clostridium innocuum		1.29	0.19		99			
45072010	**Very good identification**							**45072010**
					SPO			
Clostridium difficile		99.97	0.59	TRE 7	99			
Eubacterium limosum		0.03	-.03	CEL 5	1			
45072050	**Good identification**							**45072050**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		100.0	0.37	TRE 7				
45073000	**Acceptable identification**							**45073000**
					GRA	SPO		
Clostridium innocuum		94.32	0.41	SAL 99	99	99		
Clostridium difficile		3.29	0.13		99	99		
Fusobacterium mortiferum		0.96	0.20		1	1		
45073002	**Good identification**							**45073002**
					SPO			
Clostridium innocuum		99.60	0.41	SAL 99	99			
Lactobacillus jensenii		0.40	0.03		1			
45073120	**Good identification**							**45073120**
Clostridium septicum		100.0	0.35	MLT 94				
45074100	**Acceptable identification**							**45074100**
					COC	GRA		
Fusobacterium mortiferum		94.01	0.38	XYL 7	1	1		
Atopobium parvulum		5.99	0.06		99	99		
45075100	**Low discrimination**							**45075100**
					COC	GRA	SPO	
Atopobium parvulum		80.05	0.40	SUC 96	99	99	1	
Fusobacterium mortiferum		19.03	0.41	MLT 99	1	1	1	
Clostridium septicum		0.92	0.01		1	99	99	
45075120	**Low discrimination**							**45075120**
					COC	SPO		
Clostridium septicum		86.61	0.35	MLT 94	1	99		
Atopobium parvulum		13.39	0.27	XYL 1	99	1		
45075322	**Good identification**							**45075322**
Parabacteroides distasonis		100.0	0.41	RAF 99				
45075332	**Good identification**							**45075332**
Parabacteroides distasonis		100.0	0.36	RAF 99				
45075722	**Good identification**							**45075722**
Parabacteroides distasonis		100.0	0.37	RAF 99				
45076010	**Good identification**							**45076010**
Clostridium difficile		100.0	0.37	TRE 7				
45100000	**Low discrimination**							**45100000**
					GRA	SPO		
Clostridium bifermentans		53.47	0.84	IND 90	99	99		
Clostridium botulinum biovar A		27.46	0.79		99	99		
Prevotella disiens		9.80	0.66		1	1		
45100001	**Low discrimination**							**45100001**
Clostridium bifermentans		45.34	0.50	IND 90				
Clostridium botulinum biovar A		23.29	0.45	URE 1				
Clostridium novyi biovar A		13.65	0.53					
45100002	**Low discrimination**							**45100002**
Clostridium sporogenes		65.04	0.58	MNS 1				
Clostridium bifermentans		17.13	0.50					

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		8.80	0.45				
45100003	**Low discrimination**						**45100003**
Clostridium novyi biovar A		47.85	0.40	MNS 1			
Clostridium sporogenes		34.54	0.24				
Clostridium bifermentans		9.10	0.15				
45100010	**Low discrimination**						**45100010**
Clostridium bifermentans		45.85	0.50	IND 90			
Clostridium botulinum biovar A		23.54	0.45	MLZ 1			
Clostridium novyi biovar A		13.80	0.53				
45100011	**Acceptable identification**						**45100011**
Clostridium novyi biovar A		94.27	0.53	MNS 1			
Clostridium bifermentans		3.16	0.15				
Clostridium botulinum biovar A		1.63	0.11				
45100012	**Low discrimination**						**45100012**
Clostridium novyi biovar A		47.85	0.40	MNS 1			
Clostridium sporogenes		34.54	0.24				
Clostridium bifermentans		9.10	0.15				
45100013	**Good identification**						**45100013**
Clostridium novyi biovar A		100.0	0.40	MNS 1			
45100020	**Low discrimination**						**45100020**
					COC	SPO	
Clostridium bifermentans		76.77	0.63	IND 90	1	99	
Gemella morbillorum		6.27	0.48	NAG 6	99	1	
Clostridium botulinum biovar A		6.24	0.45		1	99	
45100021	**Low discrimination**						**45100021**
					COC	SPO	
Clostridium novyi biovar A		47.78	0.43	MNS 1	1	99	
Clostridium bifermentans		40.53	0.29		1	99	
Gemella morbillorum		3.31	0.14		99	1	
45100022	**Low discrimination**						**45100022**
					SPO		
Atopobium minutum		37.38	0.39	MLT 1	1		
Clostridium bifermentans		29.93	0.29		99		
Clostridium sporogenes		17.98	0.24		99		
45100030	**Low discrimination**						**45100030**
					COC	SPO	
Clostridium novyi biovar A		47.78	0.43	MNS 1	1	99	
Clostridium bifermentans		40.53	0.29		1	99	
Gemella morbillorum		3.31	0.14		99	1	
45100031	**Good identification**						**45100031**
Clostridium novyi biovar A		100.0	0.43	MNS 1			
45100040	**Low discrimination**						**45100040**
Clostridium botulinum biovar A		75.80	0.70	MNS 7			
Clostridium bifermentans		20.59	0.60				
Clostridium novyi biovar A		1.00	0.50				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45100041	**Low discrimination**						**45100041**
Clostridium novyi biovar A		50.00	0.50	MNS 1			
Clostridium botulinum biovar A		38.22	0.36				
Clostridium bifermentans		10.38	0.26				
45100042	**Low discrimination**						**45100042**
Clostridium botulinum biovar A		55.94	0.36	MNS 7			
Clostridium bifermentans		15.19	0.26	BGL 1			
Clostridium sporogenes		13.98	0.24				
45100043	**Good identification**						**45100043**
Clostridium novyi biovar A		95.81	0.37	MNS 1			
Clostridium botulinum biovar A		4.19	0.02				
45100050	**Low discrimination**						**45100050**
Clostridium novyi biovar A		44.18	0.50	MNS 1			
Clostridium botulinum biovar A		33.77	0.36				
Clostridium botulinum biovar B		12.22	0.26				
45100051	**Good identification**						**45100051**
Clostridium novyi biovar A		99.23	0.50	MNS 1			
Clostridium botulinum biovar A		0.77	0.02				
45100052	**Good identification**						**45100052**
Clostridium novyi biovar A		95.81	0.37	MNS 1			
Clostridium botulinum biovar A		4.19	0.02				
45100053	**Good identification**						**45100053**
Clostridium novyi biovar A		100.0	0.37	MNS 1			
45100060	**Low discrimination**						**45100060**
Clostridium bifermentans		49.16	0.40	IND 90			
Clostridium botulinum biovar A		28.64	0.36	NAG 6			
Clostridium botulinum biovar B		10.36	0.26	SOR 4			
45100061	**Acceptable identification**						**45100061**
Clostridium novyi biovar A		92.26	0.40	MNS 1			
Clostridium bifermentans		4.89	0.05				
Clostridium botulinum biovar A		2.85	0.02				
45100070	**Low discrimination**						**45100070**
Clostridium novyi biovar A		61.39	0.40	MNS 1			
Clostridium botulinum biovar B		33.46	0.21				
Clostridium bifermentans		3.26	0.05				
45100071	**Good identification**						**45100071**
Clostridium novyi biovar A		100.0	0.40	MNS 1			
45100100	**Low discrimination**						**45100100**
					GRA	SPO	
Clostridium bifermentans		44.38	0.50	IND 90	99	99	
Clostridium botulinum biovar A		22.79	0.45	LAC 1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella bivia		16.71	0.32		1	1		
45100120	**Very good identification**							**45100120**
					COC	GRA	SPO	
Prevotella bivia		99.78	0.66	ESL 1	1	1	1	
Clostridium bifermentans		0.17	0.29		1	99	99	
Gemella morbillorum		0.01	0.14		99	99	1	
45100140	**Low discrimination**							**45100140**
					GRA	SPO		
Clostridium botulinum biovar A		74.81	0.36	MNS 7	99	99		
Clostridium bifermentans		20.32	0.26	LAC 1	99	99		
Prevotella bivia		1.86	-.03		1	1		
45100200	**Low discrimination**							**45100200**
					GRA	SPO		
Clostridium bifermentans		53.48	0.50	IND 90	99	99		
Clostridium botulinum biovar A		27.47	0.45	RHA 1	99	99		
Prevotella disiens		9.80	0.32		1	1		
45100240	**Low discrimination**							**45100240**
Clostridium botulinum biovar A		76.23	0.36	MNS 7				
Clostridium bifermentans		20.71	0.26	RHA 1				
Clostridium novyi biovar A		1.01	0.16					
45100400	**Low discrimination**							**45100400**
					COC	SPO		
Clostridium bifermentans		42.76	0.50	IND 90	1	99		
Gemella morbillorum		22.06	0.48	ARA 1	99	1		
Clostridium botulinum biovar A		21.96	0.45		1	99		
45100440	**Low discrimination**							**45100440**
					COC	SPO		
Clostridium botulinum biovar A		74.30	0.36	MNS 7	1	99		
Clostridium bifermentans		20.18	0.26	ARA 1	1	99		
Gemella morbillorum		2.52	0.14		99	1		
45101000	**Low discrimination**							**45101000**
					COC	SPO		
Gemella morbillorum		55.76	0.60	ESL 1	99	1		
Clostridium bifermentans		22.30	0.50		1	99		
Clostridium botulinum biovar A		11.45	0.45		1	99		
45101001	**Low discrimination**							**45101001**
					COC	SPO		
Clostridium novyi biovar A		32.24	0.37	MNS 1	1	99		
Gemella morbillorum		24.34	0.26	GAL 10	99	1		
Clostridium chauvoei		18.54	0.22		1	99		
45101002	**Low discrimination**							**45101002**
					COC	SPO		
Atopobium minutum		47.92	0.39	MLT 1	1	1		
Clostridium sporogenes		23.05	0.24		1	99		
Gemella morbillorum		15.18	0.26		99	1		
45101010	**Low discrimination**							**45101010**
					COC	SPO		
Clostridium novyi biovar A		42.77	0.37	MNS 1	1	99		
Gemella morbillorum		32.29	0.26	GAL 10	99	1		
Clostridium bifermentans		12.92	0.15		1	99		
45101011	**Good identification**							**45101011**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium novyi biovar A		100.0	0.37	MNS 1				
45101020	**Low discrimination**							**45101020**
					COC	GRA	SPO	
Gemella morbillorum		57.79	0.43	ESL 1	99	99	1	
Prevotella bivia		13.82	0.19	NAG 10	1	1	1	
Clostridium bifermentans		13.28	0.29		1	99	99	
45101040	**Low discrimination**							**45101040**
					COC	SPO		
Clostridium botulinum biovar A		60.34	0.36	MNS 7	1	99		
Clostridium bifermentans		16.39	0.26	GAL 1	1	99		
Gemella morbillorum		9.93	0.26		99	1		
45101101	**Very good identification**							**45101101**
					COC	SPO		
Clostridium chauvoei		99.92	0.57	SUC 99	1	99		
Blautia hansenii		0.08	-.03		99	1		
45101103	**Good identification**							**45101103**
Clostridium chauvoei		100.0	0.44	SUC 99				
45101120	**Acceptable identification**							**45101120**
					GRA	SPO		
Prevotella bivia		91.32	0.53	ESL 1	1	1		
Clostridium septicum		5.45	0.39		99	99		
Prevotella melaninogenica		1.74	0.30		1	1		
45101121	**Low discrimination**							**45101121**
					GRA	SPO		
Clostridium chauvoei		89.86	0.48	SUC 99	99	99		
Prevotella bivia		9.40	0.19		1	1		
Clostridium septicum		0.56	0.05		99	99		
45101123	**Good identification**							**45101123**
Clostridium chauvoei		100.0	0.36	SUC 99				
45101400	**Acceptable identification**							**45101400**
					COC	SPO		
Gemella morbillorum		93.44	0.43	ESL 1	99	1		
Clostridium bifermentans		3.40	0.15	ARA 10	1	99		
Clostridium botulinum biovar A		1.75	0.11		1	99		
45101722	**Good identification**							**45101722**
Bacteroides eggerthii		100.0	0.36	IND 99				
45102000	**Low discrimination**							**45102000**
					GRA	SPO		
Clostridium bifermentans		48.78	0.50	IND 90	99	99		
Clostridium botulinum biovar A		25.05	0.45	MAN 1	99	99		
Prevotella disiens		8.94	0.32		1	1		
45102040	**Low discrimination**							**45102040**
Clostridium botulinum biovar A		74.75	0.36	MNS 7				
Clostridium bifermentans		20.30	0.26	MAN 1				
Clostridium novyi biovar A		0.99	0.16					
45104000	**Low discrimination**							**45104000**
					COC	SPO		

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Clostridium bifermentans		41.75	0.50	IND 90	1	99		
Gemella morbillorum		21.54	0.48	XYL 1	99	1		
Clostridium botulinum biovar A		21.44	0.45		1	99		
45104040	**Low discrimination**							**45104040**
Clostridium glycolicum		83.84	0.50	MNS 1				
Clostridium botulinum biovar A		12.01	0.36	FRU 90				
Clostridium bifermentans		3.26	0.26					
45105000	**Acceptable identification**							**45105000**
					COC	SPO		
Gemella morbillorum		91.51	0.43	ESL 1	99	1		
Clostridium bifermentans		3.33	0.15	XYL 10	1	99		
Clostridium glycolicum		2.06	0.11		1	99		
45105722	**Good identification**							**45105722**
Bacteroides eggerthii		100.0	0.56	IND 99				
45110000	**Low discrimination**							**45110000**
					COC	GRA	SPO	
Prevotella disiens		46.84	0.66	ESL 1	1	1	1	
Gemella morbillorum		14.65	0.66		99	99	1	
Clostridium bifermentans		13.45	0.62		1	99	99	
45110002	**Low discrimination**							**45110002**
					GRA	SPO		
Clostridium sporogenes		51.76	0.45	MNS 1	99	99		
Atopobium minutum		17.06	0.47		99	1		
Prevotella disiens		14.16	0.32		1	1		
45110010	**Low discrimination**							**45110010**
					COC	GRA	SPO	
Clostridium difficile		75.87	0.50	MLT 1	1	99	99	
Prevotella disiens		10.41	0.32	MAN 90	1	1	1	
Gemella morbillorum		3.26	0.32		99	99	1	
45110011	**Low discrimination**							**45110011**
					GRA	SPO		
Clostridium novyi biovar A		68.55	0.37	MNS 1	99	99		
Clostridium difficile		27.65	0.15	FRU 10	99	99		
Prevotella disiens		3.79	-.03		1	1		
45110020	**Low discrimination**							**45110020**
					COC	GRA	SPO	
Gemella morbillorum		44.05	0.50	ESL 1	99	99	1	
Clostridium bifermentans		23.24	0.41	NAG 10	1	99	99	
Prevotella disiens		12.80	0.32		1	1	1	
45110022	**Low discrimination**							**45110022**
					COC	SPO		
Atopobium minutum		73.74	0.38	MLT 1	1	1		
Clostridium sporogenes		6.78	0.11		1	99		
Gemella morbillorum		6.38	0.15		99	1		
45110040	**Low discrimination**							**45110040**
Clostridium botulinum biovar A		38.72	0.50	MNS 7				
Clostridium glycolicum		20.27	0.44	FRU 6				
Clostridium botulinum biovar B		18.01	0.41					
45110050	**Low discrimination**							**45110050**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		66.56	0.36	ESL 1				
Clostridium difficile		21.07	0.28	SUC 90				
Clostridium novyi biovar A		6.68	0.34					
45110060	**Low discrimination**							**45110060**
Clostridium botulinum biovar B		82.91	0.36	ESL 1				
Clostridium bifermentans		5.17	0.18	SUC 90				
Clostridium botulinum biovar A		3.66	0.16					
45110100	**Low discrimination**							**45110100**
					COC	GRA		
Fusobacterium mortiferum		55.54	0.59	MLT 11	1	1		
Prevotella disiens		18.67	0.32		1	1		
Gemella morbillorum		5.84	0.32		99	99		
45110102	**Low discrimination**							**45110102**
					GRA	SPO		
Fusobacterium mortiferum		83.45	0.42	MLT 11	1	1		
Clostridium sporogenes		9.32	0.11	BGL 10	99	99		
Atopobium minutum		3.07	0.12		99	1		
45110120	**Low discrimination**							**45110120**
					GRA	SPO		
Prevotella bivia		89.18	0.50	ESL 1	1	1		
Clostridium septicum		8.45	0.39	FRU 10	99	99		
Clostridium perfringens		1.73	0.34		99	84		
45110400	**Low discrimination**							**45110400**
					COC	GRA	SPO	
Gemella morbillorum		65.51	0.50	ESL 1	99	99	1	
Prevotella disiens		19.04	0.32	ARA 10	1	1	1	
Clostridium bifermentans		5.47	0.28		1	99	99	
45111000	**Low discrimination**							**45111000**
					COC	GRA		
Gemella morbillorum		62.21	0.61	ESL 1	99	99		
Prevotella intermedia		19.05	0.52		1	1		
Fusobacterium mortiferum		7.14	0.55		1	1		
45111002	**Low discrimination**							**45111002**
					GRA			
Atopobium minutum		30.09	0.38	MLT 1	99			
Fusobacterium mortiferum		15.92	0.39		1			
Lactobacillus jensenii		15.08	0.29		99			
45111020	**Low discrimination**							**45111020**
					COC	SPO		
Clostridium perfringens		56.42	0.53	SUC 91	1	84		
Clostridium septicum		20.42	0.39	LAC 88	1	99		
Gemella morbillorum		17.67	0.45		99	1		
45111022	**Low discrimination**							**45111022**
					GRA			
Clostridium perfringens		79.10	0.48	SUC 91	99			
Prevotella oralis		11.18	0.30	LAC 88	1			
Atopobium minutum		3.80	0.30		99			
45111060	**Good identification**							**45111060**
					COC	SPO		
Clostridium perfringens		95.54	0.41	SUC 91	1	84		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		1.83	0.05	LAC 88	1	99	
Gemella morbillorum		1.59	0.11		99	1	
45111062	**Good identification**						**45111062**
Clostridium perfringens		100.0	0.35	SUC 91			
45111100	**Low discrimination**						**45111100**
					GRA	SPO	
Fusobacterium mortiferum		48.96	0.62	MLT 11	1	1	
Clostridium septicum		23.50	0.39		99	99	
Clostridium perfringens		14.72	0.42		99	84	
45111101	**Very good identification**						**45111101**
					GRA	SPO	
Clostridium chauvoei		99.67	0.63	SUC 99	99	99	
Fusobacterium mortiferum		0.19	0.28		1	1	
Clostridium septicum		0.09	0.05		99	99	
45111102	**Low discrimination**						**45111102**
					GRA	CAT	
Clostridium perfringens		28.36	0.36	SUC 91	99	1	
Fusobacterium mortiferum		22.27	0.45	NAG 97	1	5	
Capnocytophaga ochracea		18.47	0.30		1	90	
45111103	**Good identification**						**45111103**
					GRA	SPO	
Clostridium chauvoei		99.74	0.50	SUC 99	99	99	
Clostridium perfringens		0.15	0.02		99	84	
Fusobacterium mortiferum		0.12	0.11		1	1	
45111120	**Low discrimination**						**45111120**
					GRA	SPO	
Clostridium septicum		79.51	0.73		99	99	
Clostridium perfringens		16.27	0.68		99	84	
Prevotella melaninogenica		2.31	0.47		1	1	
45111121	**Low discrimination**						**45111121**
Clostridium chauvoei		75.09	0.55	SUC 99			
Clostridium septicum		19.89	0.39				
Clostridium perfringens		4.07	0.34				
45111122	**Low discrimination**						**45111122**
					GRA	SPO	
Clostridium perfringens		56.64	0.62	SUC 91	99	84	
Prevotella oralis		17.11	0.50		1	1	
Clostridium septicum		12.00	0.44		99	99	
45111123	**Low discrimination**						**45111123**
					GRA	SPO	
Clostridium chauvoei		79.78	0.42	SUC 99	99	99	
Clostridium perfringens		11.54	0.28		99	84	
Prevotella oralis		3.48	0.16		1	1	
45111130	**Low discrimination**						**45111130**
					GRA	SPO	
Clostridium septicum		78.50	0.39	MLZ 1	99	99	
Clostridium perfringens		16.07	0.34		99	84	
Prevotella melaninogenica		2.28	0.12		1	1	
45111160	**Low discrimination**						**45111160**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		78.25	0.56	SUC 91	99	84	
Clostridium septicum		20.28	0.39		99	99	
Prevotella melaninogenica		0.59	0.12		1	1	
45111162	**Low discrimination**						**45111162**
					GRA	CAT	
Clostridium perfringens		88.02	0.50	SUC 91	99	1	
Capnocytophaga ochracea		9.31	0.30		1	90	
Prevotella oralis		1.41	0.16		1	1	
45111320	**Low discrimination**						**45111320**
					GRA	SPO	
Clostridium septicum		75.32	0.39	RHA 1	99	99	
Clostridium perfringens		15.42	0.34		99	84	
Prevotella oralis		4.89	0.22		1	1	
45111322	**Low discrimination**						**45111322**
					GRA		
Prevotella oralis		88.25	0.44	SUC 92	1		
Clostridium perfringens		6.89	0.28	RAF 92	99		
Parabacteroides distasonis		1.77	0.27		1		
45111400	**Acceptable identification**						**45111400**
					COC	GRA	
Gemella morbillorum		90.01	0.45	ESL 1	99	99	
Lactobacillus fermentum		5.76	0.36	ARA 10	1	99	
Prevotella intermedia		2.51	0.17		1	1	
45111402	**Low discrimination**						**45111402**
					COC		
Lactobacillus fermentum		86.37	0.36	ESL 1	1		
Gemella morbillorum		13.63	0.11	SUC 87	99		
45111500	**Low discrimination**						**45111500**
					GRA		
Lactobacillus fermentum		57.22	0.40	ESL 1	99		
Fusobacterium mortiferum		13.33	0.28	SUC 87	1		
Bifidobacterium longum subsp. longum		7.33	0.18		99		
45111502	**Low discrimination**						**45111502**
					GRA	CAT	
Lactobacillus fermentum		61.64	0.40	ESL 1	99	1	
Capnocytophaga ochracea		23.13	0.17	SUC 87	1	90	
Bifidobacterium longum subsp. longum		7.90	0.18		99	1	
45111520	**Low discrimination**						**45111520**
					GRA	SPO	
Clostridium septicum		50.75	0.39	ARA 1	99	99	
Odoribacter splanchnicus		30.08	0.32		1	1	
Clostridium perfringens		10.39	0.34		99	84	
45112000	**Low discrimination**						**45112000**
					GRA	SPO	
Eubacterium limosum		84.15	0.66	MLT 1	99	1	
Clostridium difficile		13.60	0.54		99	99	
Prevotella disiens		1.01	0.32		1	1	
45112010	**Very good identification**						**45112010**
					SPO	CAT	
Clostridium difficile		98.71	0.66	MLT 1	99	1	
Eubacterium limosum		1.26	0.32		1	1	
Propionibacterium granulosum		0.02	0.14		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45112050	**Good identification**							**45112050**
					SPO			
Clostridium difficile		99.40	0.44	MLT 1	99			
Clostridium botulinum biovar B		0.35	0.02	SOR 5	99			
Eubacterium limosum		0.24	-.03		1			
45113120	**Low discrimination**							**45113120**
					GRA	SPO		
Clostridium septicum		79.84	0.39	MAN 1	99	99		
Clostridium perfringens		16.34	0.34		99	84		
Prevotella melaninogenica		2.32	0.12		1	1		
45114000	**Low discrimination**							**45114000**
					COC	GRA	SPO	
Gemella morbillorum		42.04	0.50	ESL 1	99	99	1	
Clostridium glycolicum		33.83	0.44	XYL 10	1	99	99	
Prevotella disiens		12.22	0.32		1	1	1	
45114040	**Very good identification**							**45114040**
					COC	SPO		
Clostridium glycolicum		99.74	0.66	MNS 1	1	99		
Clostridium botulinum biovar A		0.10	0.16		1	99		
Gemella morbillorum		0.07	0.15		99	1		
45114100	**Low discrimination**							**45114100**
					COC	GRA	SPO	
Fusobacterium mortiferum		70.99	0.40	MLT 11	1	1	1	
Gemella morbillorum		11.02	0.15	XYL 7	99	99	1	
Clostridium glycolicum		8.87	0.10		1	99	99	
45115000	**Low discrimination**							**45115000**
					COC	GRA	SPO	
Gemella morbillorum		75.39	0.45	ESL 1	99	99	1	
Clostridium glycolicum		12.52	0.28	XYL 10	1	99	99	
Fusobacterium mortiferum		5.86	0.36		1	1	1	
45115040	**Good identification**							**45115040**
					COC	SPO		
Clostridium glycolicum		99.68	0.50	MNS 1	1	99		
Gemella morbillorum		0.32	0.11	GAL 10	99	1		
45115102	**Low discrimination**							**45115102**
					COC	GRA	CAT	
Lactobacillus fermentum		26.07	0.36	ESL 1	1	99	1	
Capnocytophaga ochracea		23.59	0.20	SUC 87	1	1	90	
Atopobium parvulum		19.75	0.21		99	99	1	
45115120	**Low discrimination**							**45115120**
					GRA	SPO		
Clostridium septicum		67.77	0.39	XYL 1	99	99		
Clostridium perfringens		13.87	0.34		99	84		
Bacteroides vulgatus		3.76	0.15		1	1		
45115322	**Low discrimination**							**45115322**
					CAT			
Parabacteroides distasonis		56.03	0.44	RAF 99	76			
Prevotella oralis		38.47	0.28		1			
Bacteroides fragilis		2.08	-.03		99			
45115332	**Good identification**							**45115332**
Parabacteroides distasonis		100.0	0.39	RAF 99				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45115500	**Low discrimination**						**45115500**
					GRA		
Lactobacillus fermentum		53.10	0.36	ESL 1	99		
Bifidobacterium longum subsp. longum		34.75	0.26	SUC 87	99		
Bacteroides vulgatus		7.45	0.00		1		
45115502	**Low discrimination**						**45115502**
Lactobacillus fermentum		40.02	0.36	ESL 1			
Bifidobacterium adolescentis		27.40	0.28	SUC 87			
Bifidobacterium longum subsp. longum		26.19	0.26				
45115522	**Low discrimination**						**45115522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		46.69	0.36	SUC 99	99	1	
Capnocytophaga ochracea		25.63	0.23	RAF 97	1	90	
Parabacteroides distasonis		15.26	0.32		1	76	
45115530	**Good identification**						**45115530**
					GRA		
Bifidobacterium longum subsp. longum		96.95	0.40	SUC 99	99		
Bacteroides vulgatus		3.05	0.00	RAF 97	1		
45115532	**Low discrimination**						**45115532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.78	0.40	SUC 99	99	1	
Parabacteroides distasonis		9.07	0.27	RAF 97	1	76	
Capnocytophaga ochracea		3.15	0.07		1	90	
45115720	**Good identification**						**45115720**
					GRA		
Bacteroides vulgatus		99.71	0.51	SUC 96	1		
Eubacterium contortum		0.29	0.14	RAF 97	99		
45115732	**Acceptable identification**						**45115732**
					GRA		
Parabacteroides distasonis		92.77	0.35	RAF 99	1		
Clostridium clostridioforme		4.21	0.07		50		
Bifidobacterium longum subsp. longum		3.02	0.06		99		
45116010	**Good identification**						**45116010**
					SPO		
Clostridium difficile		99.76	0.44	MLT 1	99		
Eubacterium limosum		0.25	-.03	XYL 5	1		
45120000	**Low discrimination**						**45120000**
Clostridium botulinum biovar A		71.60	0.69	MNS 7			
Clostridium sporogenes		21.39	0.58				
Clostridium bifermentans		5.63	0.50				
45120002	**Low discrimination**						**45120002**
Clostridium sporogenes		89.86	0.50	MNS 1			
Clostridium botulinum biovar A		9.12	0.35				
Clostridium bifermentans		0.72	0.15				
45120040	**Low discrimination**						**45120040**
Clostridium botulinum biovar A		90.00	0.60	MNS 7			
Clostridium botulinum biovar B		8.02	0.39				
Clostridium bifermentans		0.99	0.26				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
45121101	**Good identification**							**45121101**
Clostridium chauvoei		100.0	0.40	SUC 99				
45121120	**Low discrimination**							**45121120**
					GRA	SPO		
Clostridium septicum		83.93	0.47	FRU 99	99	99		
Clostridium perfringens		10.63	0.38		99	84		
Prevotella bivia		4.74	0.19		1	1		
45130000	**Low discrimination**							**45130000**
					GRA	SPO		
Clostridium botulinum biovar A		41.04	0.49	MNS 7	99	99		
Clostridium sporogenes		33.89	0.45	FRU 6	99	99		
Prevotella disiens		9.27	0.32		1	1		
45130002	**Low discrimination**							**45130002**
					SPO	CAT		
Clostridium sporogenes		86.16	0.37	MNS 1	99	1		
Propionibacterium granulosum		5.68	0.29		1	90		
Clostridium botulinum biovar A		3.16	0.14		99	1		
45130040	**Low discrimination**							**45130040**
Clostridium botulinum biovar B		83.91	0.50	ESL 1				
Clostridium botulinum biovar A		15.04	0.40	SUC 90				
Clostridium sporogenes		0.42	0.11					
45130050	**Good identification**							**45130050**
Clostridium botulinum biovar B		98.85	0.44	ESL 1				
Clostridium difficile		0.79	0.08	SUC 90				
Clostridium botulinum biovar A		0.36	0.05					
45130060	**Good identification**							**45130060**
Clostridium botulinum biovar B		99.06	0.44	ESL 1				
Clostridium perfringens		0.58	0.11	SUC 90				
Clostridium botulinum biovar A		0.36	0.05					
45130070	**Good identification**							**45130070**
Clostridium botulinum biovar B		100.0	0.39	ESL 1				
45130100	**Low discrimination**							**45130100**
					COC	GRA	SPO	
Fusobacterium mortiferum		66.66	0.50	MLT 11	1	1	1	
Atopobium parvulum		19.70	0.29		99	99	1	
Clostridium septicum		4.87	0.13		1	99	99	
45130120	**Low discrimination**							**45130120**
					GRA	SPO		
Clostridium septicum		85.41	0.47	GAL 99	99	99		
Clostridium perfringens		10.82	0.38		99	84		
Prevotella bivia		3.03	0.15		1	1		
45131000	**Low discrimination**							**45131000**
					GRA			
Lactobacillus jensenii		32.45	0.40	SUC 90	99			
Fusobacterium mortiferum		20.48	0.46	SAL 95	1			
Clostridium perfringens		12.18	0.32	CEL 95	99			
45131002	**Low discrimination**							**45131002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Lactobacillus jensenii		62.49	0.37	SUC 90	1		
Clostridium perfringens		16.56	0.26	SAL 95	1		
Propionibacterium granulosum		8.31	0.29	CEL 95	90		
45131020	**Low discrimination**						**45131020**
					SPO		
Clostridium perfringens		61.77	0.58	SUC 91	84		
Clostridium septicum		36.11	0.47	LAC 88	99		
Lactobacillus jensenii		1.27	0.30		1		
45131022	**Acceptable identification**						**45131022**
					SPO		
Clostridium perfringens		93.91	0.52	SUC 91	84		
Lactobacillus jensenii		2.74	0.27	LAC 88	1		
Clostridium septicum		2.38	0.18		99		
45131060	**Good identification**						**45131060**
Clostridium perfringens		95.37	0.45	SUC 91			
Clostridium septicum		2.96	0.13	LAC 88			
Clostridium botulinum biovar B		1.67	0.10				
45131062	**Good identification**						**45131062**
Clostridium perfringens		100.0	0.40	SUC 91			
45131100	**Low discrimination**						**45131100**
					COC	SPO	
Atopobium parvulum		71.62	0.63	SUC 96	99	1	
Clostridium septicum		17.70	0.47		1	99	
Clostridium perfringens		6.87	0.47		1	84	
45131101	**Low discrimination**						**45131101**
					COC	SPO	
Clostridium chauvoei		88.04	0.47	SUC 99	1	99	
Atopobium parvulum		8.58	0.29	TRE 10	99	1	
Clostridium septicum		2.12	0.13		1	99	
45131102	**Low discrimination**						**45131102**
					COC	GRA	
Atopobium parvulum		75.57	0.51	SUC 96	99	99	
Clostridium perfringens		19.32	0.41		1	99	
Fusobacterium mortiferum		2.44	0.36		1	1	
45131120	**Low discrimination**						**45131120**
					COC	SPO	
Clostridium septicum		88.17	0.81		1	99	
Clostridium perfringens		11.17	0.73		1	84	
Atopobium parvulum		0.64	0.51		99	1	
45131121	**Low discrimination**						**45131121**
Clostridium septicum		78.51	0.47	URE 1			
Clostridium chauvoei		10.98	0.38				
Clostridium perfringens		9.95	0.38				
45131122	**Low discrimination**						**45131122**
					COC	SPO	
Clostridium perfringens		72.58	0.67	SUC 91	1	84	
Clostridium septicum		24.85	0.52		1	99	
Atopobium parvulum		1.55	0.38		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45131130	**Low discrimination**						**45131130**
					COC	SPO	
Clostridium septicum		88.19	0.47	MLZ 1	1	99	
Clostridium perfringens		11.17	0.38		1	84	
Atopobium parvulum		0.64	0.16		99	1	
45131160	**Low discrimination**						**45131160**
					COC	SPO	
Clostridium perfringens		70.34	0.60	SUC 91	1	84	
Clostridium septicum		29.44	0.47		1	99	
Atopobium parvulum		0.21	0.16		99	1	
45131162	**Good identification**						**45131162**
					COC	SPO	
Clostridium perfringens		98.11	0.55	SUC 91	1	84	
Clostridium septicum		1.78	0.18		1	99	
Atopobium parvulum		0.11	0.03		99	1	
45131320	**Low discrimination**						**45131320**
					COC	SPO	
Clostridium septicum		88.08	0.47	RHA 1	1	99	
Clostridium perfringens		11.16	0.38		1	84	
Atopobium parvulum		0.64	0.16		99	1	
45131520	**Low discrimination**						**45131520**
					COC	SPO	
Clostridium septicum		87.74	0.47	ARA 1	1	99	
Clostridium perfringens		11.12	0.38		1	84	
Atopobium parvulum		0.63	0.16		99	1	
45131722	**Good identification**						**45131722**
Parabacteroides distasonis		100.0	0.35	RAF 99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45132010	**Good identification**							**45132010**
					SPO	CAT		
Clostridium difficile		99.17	0.47	MLT 1	99	1		
Propionibacterium granulosum		0.66	0.22	TRE 7	1	90		
Eubacterium limosum		0.17	-.03		1	1		
45133000	**Low discrimination**							**45133000**
					SPO	CAT		
Propionibacterium granulosum		87.82	0.37	ESL 1	1	90		
Clostridium difficile		3.70	0.01	SUC 93	99	1		
Lactobacillus jensenii		3.32	0.06		1	1		
45133120	**Low discrimination**							**45133120**
					COC	SPO		
Clostridium septicum		88.19	0.47	MAN 1	1	99		
Clostridium perfringens		11.17	0.38		1	84		
Atopobium parvulum		0.64	0.16		99	1		
45135100	**Very good identification**							**45135100**
					COC	GRA	SPO	
Atopobium parvulum		99.59	0.68	SUC 96	99	99	1	
Fusobacterium mortiferum		0.21	0.34		1	1	1	
Clostridium septicum		0.13	0.13		1	99	99	
45135102	**Good identification**							**45135102**
					COC	GRA		
Atopobium parvulum		99.56	0.55	SUC 96	99	99		
Clostridium perfringens		0.14	0.07		1	99		
Fusobacterium mortiferum		0.13	0.17		1	1		
45135120	**Low discrimination**							**45135120**
					COC	SPO		
Atopobium parvulum		54.03	0.55	SUC 96	99	1		
Clostridium septicum		40.73	0.47		1	99		
Clostridium perfringens		5.16	0.38		1	84		
45135122	**Low discrimination**							**45135122**
					COC	GRA		
Atopobium parvulum		53.40	0.42	SUC 96	99	99		
Parabacteroides distasonis		25.66	0.47		1	1		
Clostridium perfringens		13.60	0.33		1	99		
45135132	**Good identification**							**45135132**
					COC	GRA		
Parabacteroides distasonis		96.24	0.42	RAF 99	1	1		
Atopobium parvulum		3.76	0.08		99	99		
45135322	**Good identification**							**45135322**
					COC	GRA		
Parabacteroides distasonis		98.36	0.55	RAF 99	1	1		
Prevotella oralis		0.95	0.08		1	1		
Atopobium parvulum		0.69	0.08		99	99		
45135332	**Good identification**							**45135332**
Parabacteroides distasonis		100.0	0.50	RAF 99				
45135522	**Low discrimination**							**45135522**
					COC	GRA		
Parabacteroides distasonis		83.18	0.43	RAF 99	1	1		
Bifidobacterium longum subsp. longum		13.97	0.26		1	99		
Atopobium parvulum		2.85	0.08		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45135532	**Low discrimination**						**45135532**
					GRA		
Parabacteroides distasonis		64.06	0.39	RAF 99	1		
Bifidobacterium longum subsp. longum		34.02	0.30		99		
Clostridium clostridioforme		1.91	0.07		50		
45135722	**Good identification**						**45135722**
Parabacteroides distasonis		98.13	0.51	RAF 99			
Clostridium clostridioforme		0.85	0.10				
Bacteroides thetaiotaomicron		0.58	-.03				
45135732	**Acceptable identification**						**45135732**
Parabacteroides distasonis		93.19	0.47	RAF 99			
Clostridium clostridioforme		6.81	0.22				
45140000	**Low discrimination**						**45140000**
					GRA	SPO	
Clostridium bifermentans		53.48	0.50	IND 90	99	99	
Clostridium botulinum biovar A		27.47	0.45	CEL 1	99	99	
Prevotella disiens		9.80	0.32		1	1	
45140040	**Low discrimination**						**45140040**
Clostridium botulinum biovar A		76.23	0.36	MNS 7			
Clostridium bifermentans		20.71	0.26	CEL 1			
Clostridium novyi biovar A		1.01	0.16				
45141120	**Acceptable identification**						**45141120**
					GRA	SPO	
Clostridium septicum		92.19	0.47	FRU 99	99	99	
Prevotella bivia		4.93	0.19		1	1	
Clostridium paraputrificum		1.44	0.14		99	99	
45141722	**Good identification**						**45141722**
Bacteroides eggerthii		100.0	0.47	IND 99			
45145722	**Very good identification**						**45145722**
Bacteroides eggerthii		100.0	0.66	IND 99			
45150100	**Acceptable identification**						**45150100**
					GRA	SPO	
Fusobacterium mortiferum		90.28	0.51	MLT 11	1	1	
Clostridium septicum		6.24	0.13		99	99	
Collinsella aerofaciens		2.00	-.03		99	1	
45150120	**Acceptable identification**						**45150120**
					GRA	SPO	
Clostridium septicum		94.54	0.47	GAL 99	99	99	
Prevotella bivia		3.18	0.15		1	1	
Clostridium paraputrificum		1.48	0.14		99	99	
45151000	**Low discrimination**						**45151000**
					GRA		
Lactobacillus jensenii		82.70	0.54	SUC 90	99		
Fusobacterium mortiferum		9.20	0.47	SAL 95	1		
Lactobacillus catenaformis		2.87	0.29		99		
45151002	**Acceptable identification**						**45151002**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		90.20	0.51	SUC 90	99		
Lactobacillus catenaformis		4.70	0.29	SAL 95	99		
Prevotella oralis		1.73	0.17		1		
45151010	**Good identification**						**45151010**
					GRA		
Lactobacillus jensenii		99.66	0.45	SUC 90	99		
Fusobacterium mortiferum		0.34	0.13	SAL 95	1		
45151012	**Good identification**						**45151012**
Lactobacillus jensenii		100.0	0.42	SUC 90			
45151020	**Low discrimination**						**45151020**
					SPO		
Clostridium septicum		77.15	0.47	LAC 99	99		
Lactobacillus jensenii		16.31	0.43		1		
Clostridium perfringens		2.81	0.30		84		
45151022	**Low discrimination**						**45151022**
					GRA	SPO	
Prevotella oralis		47.86	0.39	SUC 92	1	1	
Lactobacillus jensenii		32.77	0.40	RAF 92	99	1	
Clostridium barati		6.90	0.24	LAC 94	99	99	
45151030	**Low discrimination**						**45151030**
					SPO		
Lactobacillus jensenii		87.46	0.35	SUC 90	1		
Clostridium septicum		12.54	0.13	SAL 95	99		
45151100	**Low discrimination**						**45151100**
					GRA	SPO	
Clostridium septicum		57.86	0.47	NAG 99	99	99	
Collinsella aerofaciens		18.56	0.32		99	1	
Fusobacterium mortiferum		12.69	0.54		1	1	
45151102	**Low discrimination**						**45151102**
					GRA		
Prevotella oralis		44.70	0.38	SUC 92	1		
Lactobacillus catenaformis		11.61	0.32	RAF 92	99		
Collinsella aerofaciens		9.39	0.15	NAG 95	99		
45151120	**Very good identification**						**45151120**
Clostridium septicum		97.83	0.82				
Clostridium paraputrificum		1.53	0.48				
Clostridium perfringens		0.26	0.44				
45151121	**Good identification**						**45151121**
Clostridium septicum		96.80	0.47	URE 1			
Clostridium paraputrificum		1.51	0.14				
Clostridium chauvoei		1.17	0.21				
45151122	**Low discrimination**						**45151122**
					GRA	SPO	
Prevotella oralis		51.11	0.59	SUC 92	1	1	
Clostridium septicum		32.03	0.53	RAF 92	99	99	
Clostridium paraputrificum		10.50	0.41		99	99	
45151130	**Good identification**						**45151130**
Clostridium septicum		97.71	0.47	MLZ 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium paraputrificum		1.53	0.14					
Clostridium perfringens		0.26	0.10					
45151160	**Low discrimination**							**45151160**
Clostridium septicum		85.17	0.47	SOR 1				
Clostridium paraputrificum		9.91	0.29					
Clostridium perfringens		4.33	0.32					
45151320	**Acceptable identification**							**45151320**
					GRA	SPO		
Clostridium septicum		91.36	0.47	RHA 1	99	99		
Prevotella oralis		6.64	0.31		1	1		
Clostridium paraputrificum		1.43	0.14		99	99		
45151322	**Good identification**							**45151322**
					GRA	SPO		
Prevotella oralis		97.07	0.53	SUC 92	1	1		
Clostridium septicum		1.43	0.18	RAF 92	99	99		
Clostridium barati		0.54	0.14		99	99		
45151520	**Good identification**							**45151520**
Clostridium septicum		97.50	0.47	ARA 1				
Clostridium paraputrificum		1.52	0.14					
Clostridium perfringens		0.26	0.10					
45152010	**Good identification**							**45152010**
					SPO			
Clostridium difficile		99.76	0.44	MLT 1	99			
Eubacterium limosum		0.25	-.03	CEL 5	1			
45153120	**Good identification**							**45153120**
Clostridium septicum		97.61	0.47	MAN 1				
Clostridium paraputrificum		1.52	0.14					
Clostridium perfringens		0.26	0.10					
45155100	**Low discrimination**							**45155100**
					COC	GRA	SPO	
Atopobium parvulum		83.91	0.40	SUC 96	99	99	1	
Fusobacterium mortiferum		8.26	0.34	TRE 99	1	1	1	
Clostridium septicum		5.05	0.13		1	99	99	
45155120	**Acceptable identification**							**45155120**
					COC	GRA	SPO	
Clostridium septicum		92.55	0.47	XYL 1	1	99	99	
Atopobium parvulum		2.74	0.27		99	99	1	
Prevotella oralis		1.94	0.22		1	1	1	
45155122	**Low discrimination**							**45155122**
					CAT			
Prevotella oralis		67.51	0.44	SUC 92	1			
Bacteroides fragilis		18.73	0.25	RAF 92	99			
Capnocytophaga ochracea		5.78	0.26	XYL 11	90			
45155322	**Low discrimination**							**45155322**
					CAT			
Prevotella oralis		80.41	0.38	SUC 92	1			
Parabacteroides distasonis		17.79	0.39	RAF 92	76			
Bacteroides eggerthii		1.27	-.03	XYL 11	93			
45161100	**Low discrimination**							**45161100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Atopobium parvulum		74.89	0.36	SUC 96	99	1		
Clostridium septicum		25.11	0.21	FRU 99	1	99		
45161120	**Good identification**							**45161120**
					COC	SPO		
Clostridium septicum		99.17	0.56	FRU 99	1	99		
Atopobium parvulum		0.53	0.23		99	1		
Clostridium perfringens		0.17	0.15		1	84		
45165100	**Good identification**							**45165100**
Atopobium parvulum		100.0	0.40	SUC 96				
45170100	**Low discrimination**							**45170100**
					COC	GRA	SPO	
Atopobium parvulum		54.98	0.36	SUC 96	99	99	1	
Fusobacterium mortiferum		26.58	0.42	GAL 99	1	1	1	
Clostridium septicum		18.44	0.21		1	99	99	
45170120	**Good identification**							**45170120**
					COC	SPO		
Clostridium septicum		99.17	0.56	GAL 99	1	99		
Atopobium parvulum		0.53	0.23		99	1		
Clostridium perfringens		0.17	0.15		1	84		
45171000	**Good identification**							**45171000**
					COC	SPO		
Lactobacillus jensenii		94.39	0.62	SUC 90	1	1		
Atopobium parvulum		3.36	0.36	SAL 95	99	1		
Clostridium septicum		1.13	0.21		1	99		
45171002	**Very good identification**							**45171002**
					COC	GRA		
Lactobacillus jensenii		98.60	0.59	SUC 90	1	99		
Atopobium parvulum		0.93	0.23	SAL 95	99	99		
Fusobacterium mortiferum		0.18	0.22		1	1		
45171010	**Good identification**							**45171010**
					COC			
Lactobacillus jensenii		99.89	0.54	SUC 90	1			
Atopobium parvulum		0.11	0.01	SAL 95	99			
45171012	**Good identification**							**45171012**
					GRA			
Lactobacillus jensenii		99.57	0.51	SUC 90	99			
Leptotrichia buccalis		0.43	-.03	SAL 95	1			
45171020	**Low discrimination**							**45171020**
					SPO			
Clostridium septicum		80.57	0.56	LAC 99	99			
Lactobacillus jensenii		17.03	0.52		1			
Clostridium perfringens		1.81	0.34		84			
45171022	**Low discrimination**							**45171022**
					SPO			
Lactobacillus jensenii		78.51	0.49	SUC 90	1			
Clostridium septicum		11.37	0.27	SAL 95	99			
Clostridium perfringens		5.90	0.29		84			
45171030	**Low discrimination**							**45171030**
					SPO			
Lactobacillus jensenii		87.46	0.43	SUC 90	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium septicum		12.54	0.21	SAL 95	99			
45171032	**Good identification**							**45171032**
Lactobacillus jensenii		100.0	0.40	SUC 90				
45171100	**Low discrimination**							**45171100**
					COC	GRA	SPO	
Atopobium parvulum		74.18	0.70	SUC 96	99	99	1	
Clostridium septicum		24.87	0.56		1	99	99	
Fusobacterium mortiferum		0.54	0.45		1	1	1	
45171101	**Low discrimination**							**45171101**
					COC	SPO		
Atopobium parvulum		72.48	0.36	SUC 96	99	1		
Clostridium septicum		24.30	0.21	URE 1	1	99		
Clostridium chauvoei		3.22	0.12		1	99		
45171102	**Good identification**							**45171102**
					COC	SPO		
Atopobium parvulum		91.94	0.57	SUC 96	99	1		
Clostridium septicum		3.57	0.27		1	99		
Lactobacillus acidophilus		1.63	0.25		1	1		
45171110	**Low discrimination**							**45171110**
					COC	SPO		
Atopobium parvulum		68.63	0.36	SUC 96	99	1		
Clostridium septicum		23.02	0.21	MLZ 1	1	99		
Lactobacillus jensenii		6.49	0.19		1	1		
45171120	**Very good identification**							**45171120**
					COC	SPO		
Clostridium septicum		99.16	0.90		1	99		
Atopobium parvulum		0.53	0.57		99	1		
Clostridium perfringens		0.17	0.49		1	84		
45171121	**Good identification**							**45171121**
					COC	SPO		
Clostridium septicum		99.13	0.56	URE 1	1	99		
Atopobium parvulum		0.53	0.23		99	1		
Clostridium perfringens		0.17	0.15		1	84		
45171122	**Low discrimination**							**45171122**
					COC	SPO		
Clostridium septicum		86.52	0.61	BGL 2	1	99		
Atopobium parvulum		3.98	0.44		99	1		
Clostridium perfringens		3.33	0.43		1	84		
45171130	**Good identification**							**45171130**
					COC	SPO		
Clostridium septicum		99.10	0.56	MLZ 1	1	99		
Atopobium parvulum		0.53	0.23		99	1		
Clostridium perfringens		0.17	0.15		1	84		
45171140	**Low discrimination**							**45171140**
					COC	SPO		
Atopobium parvulum		73.12	0.36	SUC 96	99	1		
Clostridium septicum		24.52	0.21	SOR 1	1	99		
Clostridium perfringens		2.36	0.11		1	84		
45171160	**Good identification**							**45171160**
Clostridium septicum		95.51	0.56	SOR 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		3.00	0.37				
Clostridium paraputrificum		0.99	0.19				
45171300	**Low discrimination**						**45171300**
					COC	SPO	
Atopobium parvulum		74.89	0.36	SUC 96	99	1	
Clostridium septicum		25.11	0.21	RHA 1	1	99	
45171320	**Good identification**						**45171320**
					COC	SPO	
Clostridium septicum		98.98	0.56	RHA 1	1	99	
Atopobium parvulum		0.53	0.23		99	1	
Clostridium perfringens		0.17	0.15		1	84	
45171500	**Low discrimination**						**45171500**
					COC	SPO	
Atopobium parvulum		74.89	0.36	SUC 96	99	1	
Clostridium septicum		25.11	0.21	ARA 1	1	99	
45171520	**Good identification**						**45171520**
					COC	SPO	
Clostridium septicum		99.17	0.56	ARA 1	1	99	
Atopobium parvulum		0.53	0.23		99	1	
Clostridium perfringens		0.17	0.15		1	84	
45173100	**Low discrimination**						**45173100**
					COC	SPO	
Atopobium parvulum		71.97	0.36	SUC 96	99	1	
Clostridium septicum		24.14	0.21	MAN 1	1	99	
Clostridium tertium		2.09	0.09		1	99	
45173120	**Good identification**						**45173120**
					COC	SPO	
Clostridium septicum		98.76	0.56	MAN 1	1	99	
Atopobium parvulum		0.53	0.23		99	1	
Clostridium tertium		0.26	0.17		1	99	
45174100	**Good identification**						**45174100**
					COC	GRA	
Atopobium parvulum		98.08	0.40	SUC 96	99	99	
Fusobacterium mortiferum		1.92	0.22	GAL 99	1	1	
45175000	**Low discrimination**						**45175000**
					COC	GRA	
Atopobium parvulum		85.32	0.40	SUC 96	99	99	
Lactobacillus jensenii		13.03	0.28	LAC 99	1	99	
Fusobacterium mortiferum		1.08	0.19		1	1	
45175100	**Very good identification**						**45175100**
					COC	SPO	
Atopobium parvulum		99.76	0.74	SUC 96	99	1	
Clostridium septicum		0.18	0.21		1	99	
Actinomyces israelii		0.03	0.27		1	1	
45175101	**Good identification**						**45175101**
Atopobium parvulum		100.0	0.40	SUC 96			
45175102	**Very good identification**						**45175102**
					COC		
Atopobium parvulum		99.64	0.61	SUC 96	99		
Actinomyces israelii		0.23	0.30		1		
Bifidobacterium adolescentis		0.08	0.16		1		

Code		ID	T	tests	Differentiation tests			Code
Taxon		score	index	against				
45175110	**Good identification**							**45175110**
Atopobium parvulum		100.0	0.40	SUC 96				
45175120	**Low discrimination**							**45175120**
					COC	SPO		
Clostridium septicum		50.49	0.56	XYL 1	1	99		
Atopobium parvulum		49.34	0.61		99	1		
Clostridium perfringens		0.08	0.15		1	84		
45175122	**Low discrimination**							**45175122**
					COC	GRA	SPO	
Atopobium parvulum		77.61	0.49	SUC 96	99	99	1	
Clostridium septicum		9.18	0.27		1	99	99	
Parabacteroides distasonis		8.61	0.42		1	1	1	
45175132	**Low discrimination**							**45175132**
					COC	GRA		
Parabacteroides distasonis		85.53	0.38	RAF 99	1	1		
Atopobium parvulum		14.47	0.14		99	99		
45175140	**Good identification**							**45175140**
					COC			
Atopobium parvulum		98.95	0.40	SUC 96	99			
Actinomyces israelii		1.05	0.20	SOR 1	1			
45175300	**Good identification**							**45175300**
Atopobium parvulum		100.0	0.40	SUC 96				
45175322	**Acceptable identification**							**45175322**
					COC	GRA		
Parabacteroides distasonis		91.43	0.50	RAF 99	1	1		
Prevotella oralis		5.79	0.17		1	1		
Atopobium parvulum		2.78	0.14		99	99		
45175332	**Good identification**							**45175332**
Parabacteroides distasonis		98.25	0.46	RAF 99				
Clostridium clostridioforme		1.75	0.10					
45175500	**Acceptable identification**							**45175500**
					COC	SPO		
Atopobium parvulum		93.71	0.40	SUC 96	99	1		
Clostridium butyricum		4.20	0.10	ARA 1	1	99		
Actinomyces israelii		2.09	0.26		1	1		
45175522	**Low discrimination**							**45175522**
					COC	GRA		
Parabacteroides distasonis		85.43	0.39	RAF 99	1	1		
Atopobium parvulum		12.70	0.14		99	99		
Clostridium clostridioforme		1.88	0.05		1	50		
45175722	**Acceptable identification**							**45175722**
Parabacteroides distasonis		90.69	0.47	RAF 99				
Clostridium clostridioforme		4.87	0.19					
Bacteroides eggerthii		2.33	-.03					
45175732	**Low discrimination**							**45175732**
Parabacteroides distasonis		67.98	0.42	RAF 99				
Clostridium clostridioforme		30.88	0.31					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		1.15	-.04				
45177100	**Low discrimination**						**45177100**
					COC	SPO	
Atopobium parvulum		85.01	0.40	SUC 96	99	1	
Actinomyces israelii		13.15	0.40	MAN 1	1	1	
Clostridium tertium		1.84	0.11		1	99	
45177102	**Low discrimination**						**45177102**
					COC	SPO	
Actinomyces israelii		47.82	0.43	SUC 98	1	1	
Atopobium parvulum		36.37	0.27	SAL 92	99	1	
Clostridium tertium		13.38	0.19		1	99	
45177142	**Good identification**						**45177142**
Actinomyces israelii		100.0	0.36	SUC 98			
45177500	**Low discrimination**						**45177500**
					COC	SPO	
Actinomyces israelii		89.36	0.39	SUC 98	1	1	
Atopobium parvulum		7.13	0.06	SAL 92	99	1	
Clostridium butyricum		3.51	-.07		1	99	
45177502	**Low discrimination**						**45177502**
Actinomyces israelii		70.55	0.42	SUC 98			
Bifidobacterium adolescentis		29.45	0.29	SAL 92			
45200000	**Low discrimination**						**45200000**
					SPO		
Clostridium bifermentans		79.23	0.61	IND 90	99		
Clostridium botulinum biovar A		10.55	0.47	SAL 10	99		
Atopobium minutum		6.03	0.50		1		
45200002	**Low discrimination**						**45200002**
					SPO		
Atopobium minutum		74.16	0.47	SAL 1	1		
Clostridium bifermentans		14.77	0.27		99		
Clostridium sporogenes		6.40	0.19		99		
45200010	**Low discrimination**						**45200010**
Clostridium difficile		81.81	0.40	FRU 99			
Clostridium bifermentans		10.34	0.27	MAN 90			
Clostridium novyi biovar A		5.68	0.35				
45200020	**Low discrimination**						**45200020**
					SPO		
Clostridium bifermentans		69.02	0.41	IND 90	99		
Atopobium minutum		27.43	0.42	SAL 10	1		
Clostridium novyi biovar A		1.50	0.25	NAG 6	99		
45200022	**Acceptable identification**						**45200022**
					GRA	SPO	
Atopobium minutum		92.75	0.39	SAL 1	99	1	
Clostridium bifermentans		3.54	0.06		99	99	
Pseudoflavonifractor capillosus		2.37	-.03		1	1	
45200040	**Low discrimination**						**45200040**
Clostridium bifermentans		48.02	0.37	IND 90			
Clostridium botulinum biovar A		45.84	0.38	SAL 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		4.27	0.32	SOR 4			
45200100	**Low discrimination**						**45200100**
					GRA	SPO	
Fusobacterium mortiferum		60.04	0.45	FRU 99	1	1	
Clostridium bifermentans		33.05	0.27		99	99	
Clostridium botulinum biovar A		4.40	0.13		99	99	
45201002	**Low discrimination**						**45201002**
					GRA		
Atopobium minutum		72.54	0.39	SAL 1	99		
Eubacterium tortuosum		21.84	0.19		99		
Fusobacterium mortiferum		5.62	0.25		1		
45201100	**Good identification**						**45201100**
					GRA		
Fusobacterium mortiferum		97.30	0.48	FRU 99	1		
Eubacterium tortuosum		1.03	-.03		99		
Atopobium minutum		0.91	0.08		99		
45202000	**Low discrimination**						**45202000**
					SPO		
Clostridium difficile		76.47	0.45	FRU 99	99		
Eubacterium limosum		16.95	0.32		1		
Clostridium bifermentans		5.24	0.27		99		
45202010	**Very good identification**						**45202010**
					SPO		
Clostridium difficile		99.95	0.56	FRU 99	99		
Eubacterium limosum		0.05	-.03		1		
45210000	**Low discrimination**						**45210000**
					GRA	SPO	
Clostridium difficile		52.64	0.63	MAN 90	99	99	
Fusobacterium mortiferum		25.29	0.73		1	1	
Eubacterium limosum		17.09	0.52		99	1	
45210002	**Low discrimination**						**45210002**
					GRA	SPO	
Fusobacterium mortiferum		55.34	0.57	BGL 10	1	1	
Atopobium minutum		29.96	0.47		99	1	
Clostridium difficile		10.47	0.28		99	99	
45210010	**Very good identification**						**45210010**
					GRA	SPO	
Clostridium difficile		99.81	0.74	MAN 90	99	99	
Fusobacterium mortiferum		0.10	0.39		1	1	
Eubacterium limosum		0.07	0.18		99	1	
45210011	**Good identification**						**45210011**
Clostridium difficile		99.23	0.40	MAN 90			
Clostridium novyi biovar A		0.77	0.19	URE 1			
45210012	**Good identification**						**45210012**
					GRA	SPO	
Clostridium difficile		98.34	0.40	MAN 90	99	99	
Fusobacterium mortiferum		1.08	0.22	BGL 1	1	1	
Atopobium minutum		0.58	0.12		99	1	
45210020	**Low discrimination**						**45210020**
					GRA	SPO	
Atopobium minutum		40.84	0.41	SAL 1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		28.55	0.28		99	99	
Fusobacterium mortiferum		13.71	0.39		1	1	
45210022	**Acceptable identification**						**45210022**
					GRA		
Atopobium minutum		94.70	0.38	SAL 1	99		
Fusobacterium mortiferum		5.30	0.22		1		
45210030	**Good identification**						**45210030**
					SPO		
Clostridium difficile		99.71	0.40	MAN 90	99		
Atopobium minutum		0.30	0.07	NAG 1	1		
45210040	**Low discrimination**						**45210040**
					GRA	SPO	
Clostridium difficile		80.98	0.41	MAN 90	99	99	
Fusobacterium mortiferum		7.47	0.39	SOR 5	1	1	
Eubacterium limosum		5.05	0.18		99	1	
45210050	**Good identification**						**45210050**
Clostridium difficile		99.90	0.52	MAN 90			
Clostridium novyi biovar A		0.10	0.15	SOR 5			
45210100	**Very good identification**						**45210100**
					GRA	SPO	
Fusobacterium mortiferum		98.74	0.79		1	1	
Clostridium difficile		0.89	0.28		99	99	
Eubacterium limosum		0.29	0.18		99	1	
45210101	**Good identification**						**45210101**
Fusobacterium mortiferum		100.0	0.45	URE 1			
45210102	**Very good identification**						**45210102**
					GRA		
Fusobacterium mortiferum		99.77	0.63	BGL 10	1		
Atopobium minutum		0.23	0.12		99		
45210110	**Low discrimination**						**45210110**
					GRA	SPO	
Clostridium difficile		81.33	0.40	MAN 90	99	99	
Fusobacterium mortiferum		18.67	0.45	LAC 1	1	1	
45210140	**Good identification**						**45210140**
					GRA	SPO	
Fusobacterium mortiferum		95.52	0.45	SOR 1	1	1	
Clostridium difficile		4.49	0.06		99	99	
45210210	**Good identification**						**45210210**
Clostridium difficile		100.0	0.40	MAN 90			
45210300	**Good identification**						**45210300**
Fusobacterium mortiferum		100.0	0.45	RHA 1			
45210410	**Good identification**						**45210410**
Clostridium difficile		100.0	0.40	MAN 90			
45210500	**Good identification**						**45210500**
Fusobacterium mortiferum		100.0	0.45	ARA 1			
45211000	**Good identification**						**45211000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		94.99	0.76		1	1	
Atopobium minutum		1.91	0.41		99	1	
Clostridium difficile		1.33	0.28		99	99	
45211001	**Good identification**						**45211001**
					GRA		
Fusobacterium mortiferum		98.03	0.42	URE 1	1		
Atopobium minutum		1.97	0.07		99		
45211002	**Low discrimination**						**45211002**
					GRA		
Fusobacterium mortiferum		83.02	0.60	BGL 10	1		
Atopobium minutum		9.99	0.38		99		
Lactobacillus jensenii		5.01	0.29		99		
45211010	**Low discrimination**						**45211010**
					GRA	SPO	
Clostridium difficile		83.37	0.40	GAL 1	99	99	
Fusobacterium mortiferum		12.31	0.42	MAN 90	1	1	
Lactobacillus jensenii		4.08	0.24		99	1	
45211040	**Acceptable identification**						**45211040**
					GRA	SPO	
Fusobacterium mortiferum		91.48	0.42	SOR 1	1	1	
Clostridium difficile		6.68	0.06		99	99	
Atopobium minutum		1.84	0.07		99	1	
45211100	**Very good identification**						**45211100**
					GRA	SPO	
Fusobacterium mortiferum		99.12	0.82		1	1	
Clostridium septicum		0.77	0.29		99	99	
Lactobacillus catenaformis		0.05	0.15		99	1	
45211101	**Good identification**						**45211101**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.48	URE 1	1	1	
Clostridium septicum		0.77	-.06		99	99	
45211102	**Very good identification**						**45211102**
					GRA	SPO	
Fusobacterium mortiferum		98.53	0.66	BGL 10	1	1	
Clostridium barati		0.52	0.12		99	99	
Lactobacillus catenaformis		0.42	0.15		99	1	
45211110	**Good identification**						**45211110**
					GRA	SPO	
Fusobacterium mortiferum		96.42	0.48	MLZ 1	1	1	
Clostridium difficile		2.83	0.06		99	99	
Clostridium septicum		0.75	-.06		99	99	
45211120	**Very good identification**						**45211120**
					GRA	SPO	
Clostridium septicum		98.20	0.63	MLT 94	99	99	
Fusobacterium mortiferum		1.28	0.48		1	1	
Clostridium perfringens		0.20	0.23		99	84	
45211140	**Good identification**						**45211140**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.48	SOR 1	1	1	
Clostridium septicum		0.77	-.06		99	99	
45211200	**Good identification**						**45211200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		98.03	0.42	RHA 1	1		
Atopobium minutum		1.97	0.07		99		
45211300	**Good identification**						**45211300**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.48	RHA 1	1	1	
Clostridium septicum		0.77	-.06		99	99	
45211400	**Good identification**						**45211400**
					GRA	SPO	
Fusobacterium mortiferum		95.64	0.42	ARA 1	1	1	
Clostridium innocuum		2.44	0.02		99	99	
Atopobium minutum		1.92	0.07		99	1	
45211500	**Good identification**						**45211500**
					GRA	SPO	
Fusobacterium mortiferum		97.47	0.48	ARA 1	1	1	
Odoribacter splanchnicus		1.77	-.03		1	1	
Clostridium septicum		0.76	-.06		99	99	
45212000	**Low discrimination**						**45212000**
					SPO		
Clostridium difficile		81.78	0.79		99		
Eubacterium limosum		18.12	0.66		1		
Clostridium innocuum		0.05	0.27		99		
45212001	**Low discrimination**						**45212001**
					SPO		
Clostridium difficile		81.86	0.45	URE 1	99		
Eubacterium limosum		18.14	0.32		1		
45212002	**Low discrimination**						**45212002**
					SPO		
Clostridium difficile		77.46	0.45	BGL 1	99		
Eubacterium limosum		17.17	0.32		1		
Clostridium innocuum		4.67	0.27		99		
45212010	**Very good identification**						**45212010**
					SPO		
Clostridium difficile		99.95	0.91		99		
Eubacterium limosum		0.05	0.32		1		
Clostridium innocuum		0.00	0.04		99		
45212011	**Very good identification**						**45212011**
					SPO		
Clostridium difficile		99.95	0.56	URE 1	99		
Eubacterium limosum		0.05	-.03		1		
45212012	**Very good identification**						**45212012**
					SPO		
Clostridium difficile		99.90	0.56	BGL 1	99		
Clostridium innocuum		0.05	0.04		99		
Eubacterium limosum		0.05	-.03		1		
45212020	**Low discrimination**						**45212020**
					SPO		
Clostridium difficile		81.75	0.45	NAG 1	99		
Eubacterium limosum		18.12	0.32		1		
Atopobium minutum		0.13	0.07		1		
45212030	**Very good identification**						**45212030**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		99.95	0.56	NAG 1	99		
Eubacterium limosum		0.05	-.03		1		
45212040	**Very good identification**						**45212040**
					SPO		
Clostridium difficile		95.92	0.57	SOR 5	99		
Eubacterium limosum		4.08	0.32		1		
45212050	**Very good identification**						**45212050**
					SPO		
Clostridium difficile		99.99	0.69	SOR 5	99		
Eubacterium limosum		0.01	-.03		1		
45212100	**Low discrimination**						**45212100**
					GRA	SPO	
Clostridium difficile		74.29	0.45	LAC 1	99	99	
Eubacterium limosum		16.46	0.32		99	1	
Fusobacterium mortiferum		9.25	0.45		1	1	
45212110	**Very good identification**						**45212110**
					GRA	SPO	
Clostridium difficile		99.93	0.56	LAC 1	99	99	
Eubacterium limosum		0.05	-.03		99	1	
Fusobacterium mortiferum		0.03	0.11		1	1	
45212200	**Low discrimination**						**45212200**
					SPO		
Clostridium difficile		81.86	0.45	RHA 1	99		
Eubacterium limosum		18.14	0.32		1		
45212210	**Very good identification**						**45212210**
					SPO		
Clostridium difficile		99.95	0.56	RHA 1	99		
Eubacterium limosum		0.05	-.03		1		
45212400	**Low discrimination**						**45212400**
					SPO		
Clostridium difficile		81.15	0.45	ARA 1	99		
Eubacterium limosum		17.99	0.32		1		
Clostridium innocuum		0.86	0.14		99		
45212410	**Very good identification**						**45212410**
					SPO		
Clostridium difficile		99.95	0.56	ARA 1	99		
Eubacterium limosum		0.05	-.03		1		
45213000	**Low discrimination**						**45213000**
					SPO		
Clostridium innocuum		46.78	0.49	CEL 99	99		
Clostridium difficile		40.83	0.45		99		
Eubacterium limosum		9.05	0.32		1		
45213002	**Good identification**						**45213002**
					GRA	SPO	
Clostridium innocuum		98.18	0.49	CEL 99	99	99	
Clostridium difficile		0.87	0.11		99	99	
Fusobacterium mortiferum		0.76	0.25		1	1	
45213010	**Very good identification**						**45213010**
					SPO		
Clostridium difficile		98.99	0.56	GAL 1	99		
Clostridium innocuum		0.97	0.25		99		
Eubacterium limosum		0.05	-.03		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45213100	**Low discrimination**							**45213100**
					GRA	SPO		
Fusobacterium mortiferum		88.04	0.48	MAN 1	1	1		
Clostridium innocuum		5.45	0.15		99	99		
Clostridium difficile		4.76	0.11		99	99		
45213400	**Acceptable identification**							**45213400**
					SPO			
Clostridium innocuum		94.25	0.36	CEL 99	99			
Clostridium difficile		4.71	0.11		99			
Eubacterium limosum		1.04	-.03		1			
45213402	**Good identification**							**45213402**
Clostridium innocuum		100.0	0.36	CEL 99				
45214000	**Low discrimination**							**45214000**
					GRA	SPO		
Clostridium difficile		55.99	0.41	MAN 90	99	99		
Fusobacterium mortiferum		38.47	0.54	XYL 5	1	1		
Eubacterium limosum		3.49	0.18		99	1		
45214002	**Low discrimination**							**45214002**
					GRA	SPO		
Fusobacterium mortiferum		82.99	0.37	XYL 7	1	1		
Clostridium difficile		10.98	0.06	BGL 10	99	99		
Atopobium minutum		6.03	0.12		99	1		
45214010	**Good identification**							**45214010**
					GRA	SPO		
Clostridium difficile		99.86	0.52	MAN 90	99	99		
Fusobacterium mortiferum		0.14	0.20	XYL 5	1	1		
45214100	**Very good identification**							**45214100**
					GRA	SPO		
Fusobacterium mortiferum		99.37	0.60	XYL 7	1	1		
Clostridium difficile		0.63	0.06		99	99		
45214102	**Good identification**							**45214102**
Fusobacterium mortiferum		100.0	0.44	XYL 7				
45215000	**Very good identification**							**45215000**
					GRA	SPO		
Fusobacterium mortiferum		98.77	0.57	XYL 7	1	1		
Clostridium difficile		0.97	0.06		99	99		
Atopobium minutum		0.27	0.07		99	1		
45215002	**Good identification**							**45215002**
Fusobacterium mortiferum		100.0	0.40	XYL 7				
45215100	**Very good identification**							**45215100**
					COC	GRA	SPO	
Fusobacterium mortiferum		99.77	0.63	XYL 7	1	1	1	
Atopobium parvulum		0.12	0.02		99	99	1	
Clostridium septicum		0.10	-.06		1	99	99	
45215102	**Good identification**							**45215102**
					GRA			
Fusobacterium mortiferum		96.67	0.47	XYL 7	1			
Bifidobacterium longum subsp. infantis		3.33	0.04	BGL 10	99			
45216000	**Very good identification**							**45216000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium difficile		95.79	0.57	XYL 5	99	99	
Eubacterium limosum		4.07	0.32		99	1	
Fusobacterium mortiferum		0.07	0.20		1	1	
45216010	**Very good identification**						**45216010**
					SPO		
Clostridium difficile		99.99	0.69	XYL 5	99		
Eubacterium limosum		0.01	-.03		1		
45216040	**Good identification**						**45216040**
					SPO		
Clostridium difficile		98.72	0.35	XYL 5	99		
Eubacterium limosum		0.81	-.03	SOR 5	1		
Clostridium glycolicum		0.47	-.03		99		
45216050	**Good identification**						**45216050**
Clostridium difficile		100.0	0.47	XYL 5			
45220000	**Low discrimination**						**45220000**
Clostridium botulinum biovar A		63.87	0.36	MNS 7			
Clostridium bifermentans		19.38	0.27	SAL 1			
Clostridium sporogenes		8.40	0.19				
45220100	**Acceptable identification**						**45220100**
					GRA	SPO	
Fusobacterium mortiferum		94.22	0.36	FRU 99	1	1	
Clostridium botulinum biovar A		5.78	0.02		99	99	
45221100	**Acceptable identification**						**45221100**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	FRU 99	1	1	
Clostridium septicum		7.27	0.03		99	99	
45221120	**Good identification**						**45221120**
Clostridium septicum		100.0	0.37	MLT 94			
45222010	**Good identification**						**45222010**
Clostridium difficile		100.0	0.37	FRU 99			
45230000	**Low discrimination**						**45230000**
					GRA	SPO	
Fusobacterium mortiferum		63.78	0.64		1	1	
Clostridium difficile		33.45	0.43		99	99	
Eubacterium limosum		1.46	0.18		99	1	
45230002	**Acceptable identification**						**45230002**
					GRA	SPO	
Fusobacterium mortiferum		92.15	0.48	BGL 10	1	1	
Clostridium difficile		4.39	0.09		99	99	
Atopobium minutum		1.69	0.12		99	1	
45230010	**Good identification**						**45230010**
					GRA	SPO	
Clostridium difficile		99.61	0.55	TRE 7	99	99	
Fusobacterium mortiferum		0.39	0.30	MAN 90	1	1	
45230100	**Very good identification**						**45230100**
					GRA	SPO	
Fusobacterium mortiferum		99.66	0.70		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		0.23	0.09		99	99		
Clostridium septicum		0.12	0.03		99	99		
45230101	**Good identification**							**45230101**
Fusobacterium mortiferum		100.0	0.36	URE 1				
45230102	**Good identification**							**45230102**
Fusobacterium mortiferum		100.0	0.54	BGL 10				
45230120	**Acceptable identification**							**45230120**
					GRA	SPO		
Clostridium septicum		92.09	0.37	MLT 94	99	99		
Fusobacterium mortiferum		7.91	0.36	GAL 99	1	1		
45230140	**Good identification**							**45230140**
Fusobacterium mortiferum		100.0	0.36	SOR 1				
45230300	**Good identification**							**45230300**
Fusobacterium mortiferum		100.0	0.36	RHA 1				
45230500	**Good identification**							**45230500**
Fusobacterium mortiferum		100.0	0.36	ARA 1				
45231000	**Good identification**							**45231000**
					GRA	SPO		
Fusobacterium mortiferum		90.13	0.67		1	1		
Lactobacillus jensenii		9.10	0.40		99	1		
Clostridium difficile		0.32	0.09		99	99		
45231002	**Low discrimination**							**45231002**
					GRA	SPO		
Fusobacterium mortiferum		61.70	0.51	BGL 10	1	1		
Lactobacillus jensenii		37.35	0.37		99	1		
Clostridium innocuum		0.95	0.07		99	99		
45231020	**Low discrimination**							**45231020**
					GRA	SPO		
Clostridium septicum		82.63	0.37	MLT 94	99	99		
Lactobacillus jensenii		11.40	0.30	LAC 99	99	1		
Fusobacterium mortiferum		4.57	0.33		1	1		
45231100	**Good identification**							**45231100**
					COC	GRA	SPO	
Fusobacterium mortiferum		91.26	0.73		1	1	1	
Clostridium septicum		7.15	0.37		1	99	99	
Atopobium parvulum		1.52	0.31		99	99	1	
45231101	**Acceptable identification**							**45231101**
					GRA	SPO		
Fusobacterium mortiferum		92.73	0.39	URE 1	1	1		
Clostridium septicum		7.27	0.03		99	99		
45231102	**Very good identification**							**45231102**
					COC	GRA	SPO	
Fusobacterium mortiferum		95.36	0.57	BGL 10	1	1	1	
Atopobium parvulum		2.52	0.18		99	99	1	
Clostridium septicum		1.37	0.08		1	99	99	
45231110	**Acceptable identification**							**45231110**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		92.73	0.39	MLZ 1	1	1	
Clostridium septicum		7.27	0.03		99	99	
45231120	**Very good identification**						**45231120**
					GRA	SPO	
Clostridium septicum		99.69	0.71	MLT 94	99	99	
Fusobacterium mortiferum		0.13	0.39		1	1	
Clostridium perfringens		0.13	0.28		99	84	
45231121	**Good identification**						**45231121**
Clostridium septicum		100.0	0.37	MLT 94			
45231122	**Acceptable identification**						**45231122**
					GRA	SPO	
Clostridium septicum		94.40	0.42	MLT 94	99	99	
Clostridium perfringens		2.73	0.22	BGL 2	99	84	
Parabacteroides distasonis		0.70	0.22		1	1	
45231130	**Good identification**						**45231130**
Clostridium septicum		100.0	0.37	MLT 94			
45231140	**Acceptable identification**						**45231140**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	SOR 1	1	1	
Clostridium septicum		7.27	0.03		99	99	
45231160	**Good identification**						**45231160**
Clostridium septicum		97.69	0.37	MLT 94			
Clostridium perfringens		2.31	0.16	SOR 1			
45231300	**Acceptable identification**						**45231300**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	RHA 1	1	1	
Clostridium septicum		7.27	0.03		99	99	
45231320	**Good identification**						**45231320**
Clostridium septicum		100.0	0.37	MLT 94			
45231500	**Acceptable identification**						**45231500**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	ARA 1	1	1	
Clostridium septicum		7.27	0.03		99	99	
45231520	**Good identification**						**45231520**
					GRA	SPO	
Clostridium septicum		99.22	0.37	MLT 94	99	99	
Odoribacter splanchnicus		0.78	-.03	ARA 1	1	1	
45232000	**Very good identification**						**45232000**
					SPO		
Clostridium difficile		96.63	0.60	TRE 7	99		
Eubacterium limosum		2.87	0.32		1		
Clostridium innocuum		0.28	0.19		99		
45232010	**Very good identification**						**45232010**
					SPO		
Clostridium difficile		99.99	0.71	TRE 7	99		
Eubacterium limosum		0.01	-.03		1		
45232011	**Good identification**						**45232011**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		100.0	0.37	TRE 7			
45232012	**Good identification**						**45232012**
Clostridium difficile		100.0	0.37	TRE 7			
45232030	**Good identification**						**45232030**
Clostridium difficile		100.0	0.37	TRE 7			
45232040	**Good identification**						**45232040**
					SPO		
Clostridium difficile		99.43	0.38	TRE 7	99		
Eubacterium limosum		0.57	-.03	SOR 5	1		
45232050	**Good identification**						**45232050**
Clostridium difficile		100.0	0.50	TRE 7			
45232110	**Good identification**						**45232110**
Clostridium difficile		100.0	0.37	TRE 7			
45232210	**Good identification**						**45232210**
Clostridium difficile		100.0	0.37	TRE 7			
45232410	**Good identification**						**45232410**
Clostridium difficile		100.0	0.37	TRE 7			
45233000	**Low discrimination**						**45233000**
					GRA	SPO	
Clostridium innocuum		79.27	0.41	CEL 99	99	99	
Clostridium difficile		14.82	0.25		99	99	
Fusobacterium mortiferum		4.71	0.33		1	1	
45233002	**Good identification**						**45233002**
					GRA	SPO	
Clostridium innocuum		98.95	0.41	CEL 99	99	99	
Fusobacterium mortiferum		0.65	0.16		1	1	
Lactobacillus jensenii		0.40	0.03		99	1	
45233010	**Good identification**						**45233010**
Clostridium difficile		95.64	0.37	TRE 7			
Clostridium innocuum		4.37	0.18	GAL 1			
45233100	**Low discrimination**						**45233100**
					GRA	SPO	
Fusobacterium mortiferum		86.87	0.39	MAN 1	1	1	
Clostridium septicum		6.81	0.03		99	99	
Clostridium innocuum		6.33	0.07		99	99	
45233120	**Good identification**						**45233120**
Clostridium septicum		100.0	0.37	MLT 94			
45234000	**Low discrimination**						**45234000**
					GRA	SPO	
Fusobacterium mortiferum		73.17	0.45	XYL 7	1	1	
Clostridium difficile		26.83	0.21		99	99	
45234100	**Good identification**						**45234100**
					COC	GRA	
Fusobacterium mortiferum		99.38	0.51	XYL 7	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		0.62	0.02		99	99		
45235000	**Good identification**							**45235000**
					COC	GRA		
Fusobacterium mortiferum		97.73	0.48	XYL 7	1	1		
Lactobacillus jensenii		1.32	0.06		1	99		
Atopobium parvulum		0.95	0.02		99	99		
45235100	**Low discrimination**							**45235100**
					COC	GRA	SPO	
Fusobacterium mortiferum		70.38	0.54	XYL 7	1	1	1	
Atopobium parvulum		28.88	0.36		99	99	1	
Clostridium septicum		0.74	0.03		1	99	99	
45235102	**Low discrimination**							**45235102**
					COC	GRA		
Fusobacterium mortiferum		60.55	0.38	XYL 7	1	1		
Atopobium parvulum		39.46	0.23	BGL 10	99	99		
45235120	**Acceptable identification**							**45235120**
					COC	GRA	SPO	
Clostridium septicum		92.66	0.37	MLT 94	1	99	99	
Atopobium parvulum		6.45	0.23	XYL 1	99	99	1	
Fusobacterium mortiferum		0.90	0.20		1	1	1	
45235122	**Low discrimination**							**45235122**
					COC	GRA	SPO	
Parabacteroides distasonis		80.46	0.38	RAF 99	1	1	1	
Clostridium septicum		12.20	0.08		1	99	99	
Atopobium parvulum		7.34	0.10		99	99	1	
45235322	**Good identification**							**45235322**
Parabacteroides distasonis		100.0	0.46	RAF 99				
45235332	**Good identification**							**45235332**
Parabacteroides distasonis		100.0	0.42	RAF 99				
45235722	**Good identification**							**45235722**
Parabacteroides distasonis		100.0	0.43	RAF 99				
45235732	**Good identification**							**45235732**
Parabacteroides distasonis		100.0	0.38	RAF 99				
45236000	**Good identification**							**45236000**
					SPO			
Clostridium difficile		99.43	0.38	TRE 7	99			
Eubacterium limosum		0.57	-.03	XYL 5	1			
45236010	**Good identification**							**45236010**
Clostridium difficile		100.0	0.50	TRE 7				
45240100	**Good identification**							**45240100**
Fusobacterium mortiferum		100.0	0.37	FRU 99				
45241100	**Low discrimination**							**45241100**
					GRA			
Fusobacterium mortiferum		73.20	0.40	FRU 99	1			
Lactobacillus catenaformis		10.30	0.15		99			
Collinsella aerofaciens		6.82	-.03		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45241120	**Acceptable identification**						**45241120**
Clostridium septicum		91.77	0.37	MLT 94			
Clostridium paraputrificum		5.61	0.14	FRU 99			
Clostridium barati		2.18	0.12				
45242000	**Low discrimination**						**45242000**
					SPO		
Clostridium innocuum		85.13	0.39	FRU 95	99		
Clostridium difficile		14.26	0.23	GAL 95	99		
Eubacterium limosum		0.61	-.03		1		
45242002	**Good identification**						**45242002**
Clostridium innocuum		100.0	0.39	FRU 95			
45243000	**Very good identification**						**45243000**
Clostridium innocuum		100.0	0.61	FRU 95			
45243002	**Very good identification**						**45243002**
Clostridium innocuum		100.0	0.61	FRU 95			
45243010	**Good identification**						**45243010**
Clostridium innocuum		98.97	0.38	FRU 95			
Clostridium difficile		1.03	0.00	MLZ 4			
45243012	**Good identification**						**45243012**
Clostridium innocuum		100.0	0.38	FRU 95			
45243400	**Good identification**						**45243400**
Clostridium innocuum		100.0	0.48	FRU 95			
45243402	**Good identification**						**45243402**
Clostridium innocuum		100.0	0.48	FRU 95			
45247000	**Good identification**						**45247000**
Clostridium innocuum		100.0	0.39	FRU 95			
45247002	**Good identification**						**45247002**
Clostridium innocuum		100.0	0.39	FRU 95			
45250000	**Low discrimination**						**45250000**
					GRA	SPO	
Fusobacterium mortiferum		71.24	0.65		1	1	
Clostridium difficile		23.41	0.41		99	99	
Clostridium innocuum		2.44	0.27		99	99	
45250002	**Low discrimination**						**45250002**
					GRA	SPO	
Fusobacterium mortiferum		65.24	0.48	BGL 10	1	1	
Clostridium innocuum		20.10	0.27		99	99	
Clostridium barati		6.29	0.16		99	99	
45250010	**Good identification**						**45250010**
					GRA	SPO	
Clostridium difficile		99.11	0.52	CEL 5	99	99	
Fusobacterium mortiferum		0.62	0.31	MAN 90	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		0.18	0.11		99	1	
45250100	**Very good identification**						**45250100**
					GRA	SPO	
Fusobacterium mortiferum		98.94	0.71		1	1	
Clostridium barati		0.46	0.16		99	99	
Lactobacillus catenaformis		0.21	0.15		99	1	
45250101	**Good identification**						**45250101**
Fusobacterium mortiferum		100.0	0.37	URE 1			
45250102	**Low discrimination**						**45250102**
					GRA	SPO	
Fusobacterium mortiferum		87.47	0.55	BGL 10	1	1	
Clostridium barati		10.85	0.24		99	99	
Lactobacillus catenaformis		1.68	0.15		99	1	
45250110	**Low discrimination**						**45250110**
					GRA	SPO	
Fusobacterium mortiferum		59.25	0.37	MLZ 1	1	1	
Clostridium difficile		40.75	0.18		99	99	
45250120	**Low discrimination**						**45250120**
					GRA	SPO	
Clostridium septicum		78.08	0.37	MLT 94	99	99	
Clostridium barati		9.68	0.24	GAL 99	99	99	
Fusobacterium mortiferum		7.09	0.37		1	1	
45250140	**Good identification**						**45250140**
					GRA	SPO	
Fusobacterium mortiferum		96.20	0.37	SOR 1	1	1	
Clostridium barati		3.80	-.02		99	99	
45250300	**Good identification**						**45250300**
					GRA	SPO	
Fusobacterium mortiferum		96.20	0.37	RHA 1	1	1	
Clostridium barati		3.80	-.02		99	99	
45250500	**Good identification**						**45250500**
Fusobacterium mortiferum		100.0	0.37	ARA 1			
45251000	**Low discrimination**						**45251000**
					GRA	SPO	
Fusobacterium mortiferum		43.91	0.68		1	1	
Lactobacillus jensenii		25.15	0.54		99	1	
Clostridium innocuum		19.04	0.49		99	99	
45251002	**Low discrimination**						**45251002**
					SPO		
Clostridium innocuum		33.76	0.49	MAN 99	99		
Lactobacillus jensenii		29.73	0.51		1		
Lactobacillus catenaformis		17.05	0.47		1		
45251010	**Low discrimination**						**45251010**
					SPO		
Lactobacillus jensenii		82.25	0.45	SUC 90	1		
Clostridium innocuum		7.79	0.25	MLT 95	99		
Clostridium difficile		4.66	0.18		99		
45251012	**Low discrimination**						**45251012**
					SPO		
Lactobacillus jensenii		84.82	0.42	SUC 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		12.04	0.25	MLT 95	99		
Lactobacillus catenaformis		1.47	0.12		1		
45251020	**Low discrimination**						**45251020**
					SPO		
Clostridium septicum		32.53	0.37	MLT 94	99		
Lactobacillus jensenii		26.93	0.43	LAC 99	1		
Clostridium barati		25.54	0.38		99		
45251022	**Low discrimination**						**45251022**
					SPO		
Clostridium barati		70.63	0.46	SUC 99	99		
Lactobacillus jensenii		16.55	0.40		1		
Lactobacillus catenaformis		9.49	0.36		1		
45251030	**Acceptable identification**						**45251030**
					SPO		
Lactobacillus jensenii		92.86	0.35	SUC 90	1		
Clostridium septicum		3.40	0.03	MLT 95	99		
Clostridium barati		2.67	0.04		99		
45251100	**Low discrimination**						**45251100**
					GRA		
Fusobacterium mortiferum		72.71	0.74		1		
Lactobacillus catenaformis		10.23	0.50		99		
Collinsella aerofaciens		6.77	0.32		99		
45251101	**Low discrimination**						**45251101**
					GRA		
Fusobacterium mortiferum		73.20	0.40	URE 1	1		
Lactobacillus catenaformis		10.30	0.15		99		
Collinsella aerofaciens		6.82	-.03		99		
45251102	**Low discrimination**						**45251102**
					GRA	SPO	
Clostridium barati		37.95	0.46	SUC 99	99	99	
Lactobacillus catenaformis		30.60	0.50		99	1	
Fusobacterium mortiferum		24.16	0.58		1	1	
45251110	**Low discrimination**						**45251110**
					GRA		
Fusobacterium mortiferum		69.06	0.40	MLZ 1	1		
Lactobacillus catenaformis		9.72	0.15		99		
Collinsella aerofaciens		6.43	-.03		99		
45251120	**Good identification**						**45251120**
Clostridium septicum		91.51	0.71	MLT 94			
Clostridium paraputrificum		5.60	0.48				
Clostridium barati		2.18	0.46				
45251121	**Acceptable identification**						**45251121**
Clostridium septicum		91.77	0.37	MLT 94			
Clostridium paraputrificum		5.61	0.14	URE 1			
Clostridium barati		2.18	0.12				
45251122	**Low discrimination**						**45251122**
Clostridium barati		55.95	0.54	SUC 99			
Clostridium paraputrificum		20.55	0.41				
Clostridium septicum		16.00	0.42				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45251130	**Acceptable identification**						**45251130**
Clostridium septicum		91.53	0.37	MLT 94			
Clostridium paraputrificum		5.60	0.14	MLZ 1			
Clostridium barati		2.18	0.12				
45251140	**Low discrimination**						**45251140**
					GRA	SPO	
Fusobacterium mortiferum		54.27	0.40	SOR 1	1	1	
Clostridium barati		27.18	0.20		99	99	
Lactobacillus catenaformis		7.64	0.15		99	1	
45251160	**Low discrimination**						**45251160**
Clostridium septicum		60.05	0.37	MLT 94			
Clostridium paraputrificum		27.37	0.29	SOR 1			
Clostridium barati		12.30	0.28				
45251162	**Low discrimination**						**45251162**
					SPO		
Clostridium barati		71.11	0.36	SUC 99	99		
Clostridium paraputrificum		22.60	0.22	SOR 8	99		
Lactobacillus acidophilus		2.96	0.15		1		
45251300	**Low discrimination**						**45251300**
					GRA	SPO	
Fusobacterium mortiferum		55.28	0.40	RHA 1	1	1	
Clostridium barati		27.69	0.20		99	99	
Lactobacillus catenaformis		7.78	0.15		99	1	
45251320	**Low discrimination**						**45251320**
Clostridium septicum		78.38	0.37	MLT 94			
Clostridium barati		16.05	0.28	RHA 1			
Clostridium paraputrificum		4.79	0.14				
45251322	**Low discrimination**						**45251322**
					GRA	SPO	
Clostridium barati		78.41	0.36	SUC 99	99	99	
Prevotella oralis		12.40	0.23	RHA 8	1	1	
Clostridium paraputrificum		3.35	0.07		99	99	
45251400	**Low discrimination**						**45251400**
					GRA	SPO	
Clostridium innocuum		80.87	0.36	MAN 99	99	99	
Fusobacterium mortiferum		10.68	0.34		1	1	
Lactobacillus jensenii		6.11	0.19		99	1	
45251402	**Low discrimination**						**45251402**
					SPO		
Clostridium innocuum		89.94	0.36	MAN 99	99		
Lactobacillus jensenii		4.53	0.16		1		
Lactobacillus catenaformis		2.60	0.12		1		
45251500	**Low discrimination**						**45251500**
					GRA		
Fusobacterium mortiferum		71.48	0.40	ARA 1	1		
Lactobacillus catenaformis		10.06	0.15		99		
Collinsella aerofaciens		6.66	-.03		99		
45251520	**Acceptable identification**						**45251520**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		91.15	0.37	MLT 94			
Clostridium paraputrificum		5.57	0.14	ARA 1			
Clostridium barati		2.17	0.12				
45252000	**Low discrimination**						**45252000**
					SPO		
Clostridium innocuum		52.28	0.61	GAL 95	99		
Clostridium difficile		45.63	0.57		99		
Eubacterium limosum		1.94	0.32		1		
45252002	**Very good identification**						**45252002**
					SPO		
Clostridium innocuum		99.03	0.61	GAL 95	99		
Clostridium difficile		0.87	0.23		99		
Eubacterium limosum		0.04	-.03		1		
45252010	**Very good identification**						**45252010**
					SPO		
Clostridium difficile		99.02	0.69	CEL 5	99		
Clostridium innocuum		0.97	0.38		99		
Eubacterium limosum		0.01	-.03		1		
45252040	**Low discrimination**						**45252040**
					SPO		
Clostridium difficile		81.43	0.35	CEL 5	99		
Clostridium innocuum		17.91	0.27	SOR 5	99		
Eubacterium limosum		0.67	-.03		1		
45252050	**Good identification**						**45252050**
Clostridium difficile		99.81	0.47	CEL 5			
Clostridium innocuum		0.19	0.04	SOR 5			
45252400	**Good identification**						**45252400**
					SPO		
Clostridium innocuum		95.05	0.48	GAL 95	99		
Clostridium difficile		4.75	0.23		99		
Eubacterium limosum		0.20	-.03		1		
45252402	**Good identification**						**45252402**
Clostridium innocuum		100.0	0.48	GAL 95			
45253000	**Very good identification**						**45253000**
					GRA	SPO	
Clostridium innocuum		99.90	0.83		99	99	
Clostridium difficile		0.05	0.23		99	99	
Fusobacterium mortiferum		0.02	0.34		1	1	
45253001	**Good identification**						**45253001**
Clostridium innocuum		100.0	0.49	URE 1			
45253002	**Very good identification**						**45253002**
					SPO		
Clostridium innocuum		99.96	0.83		99		
Clostridium barati		0.03	0.20		99		
Lactobacillus jensenii		0.01	0.16		1		
45253003	**Good identification**						**45253003**
Clostridium innocuum		100.0	0.49	URE 1			
45253010	**Good identification**						**45253010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium innocuum		94.75	0.60	MLZ 4	99		
Clostridium difficile		5.15	0.35		99		
Lactobacillus jensenii		0.10	0.11		1		
45253012	**Very good identification**						**45253012**
					SPO		
Clostridium innocuum		99.87	0.60	MLZ 4	99		
Lactobacillus jensenii		0.07	0.08		1		
Clostridium difficile		0.06	0.00		99		
45253020	**Good identification**						**45253020**
Clostridium innocuum		96.54	0.49	NAG 1			
Clostridium barati		2.63	0.20				
Clostridium septicum		0.39	0.03				
45253022	**Acceptable identification**						**45253022**
					SPO		
Clostridium innocuum		92.15	0.49	NAG 1	99		
Clostridium barati		7.53	0.28		99		
Lactobacillus jensenii		0.21	0.06		1		
45253040	**Good identification**						**45253040**
Clostridium innocuum		99.76	0.49	SOR 1			
Clostridium difficile		0.24	0.01				
45253042	**Good identification**						**45253042**
Clostridium innocuum		99.76	0.49	SOR 1			
Clostridium barati		0.24	0.01				
45253100	**Acceptable identification**						**45253100**
					GRA	SPO	
Clostridium innocuum		91.05	0.49	LAC 1	99	99	
Fusobacterium mortiferum		4.95	0.40		1	1	
Clostridium barati		2.48	0.20		99	99	
45253102	**Acceptable identification**						**45253102**
					SPO		
Clostridium innocuum		90.82	0.49	LAC 1	99		
Clostridium barati		7.42	0.28		99		
Lactobacillus catenaformis		0.70	0.15		1		
45253120	**Low discrimination**						**45253120**
Clostridium septicum		76.95	0.37	MLT 94			
Clostridium barati		15.76	0.28	MAN 1			
Clostridium paraputrificum		4.71	0.14				
45253122	**Low discrimination**						**45253122**
Clostridium barati		85.27	0.36	SUC 99			
Clostridium paraputrificum		3.64	0.07	MAN 8			
Clostridium innocuum		3.52	0.15				
45253200	**Good identification**						**45253200**
Clostridium innocuum		100.0	0.49	RHA 1			
45253202	**Good identification**						**45253202**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		99.76	0.49	RHA 1			
Clostridium barati		0.24	0.01				
45253400	**Very good identification**						**45253400**
Clostridium innocuum		100.0	0.70				
45253401	**Good identification**						**45253401**
Clostridium innocuum		100.0	0.36	URE 1			
45253402	**Very good identification**						**45253402**
Clostridium innocuum		100.0	0.70				
45253403	**Good identification**						**45253403**
Clostridium innocuum		100.0	0.36	URE 1			
45253410	**Good identification**						**45253410**
Clostridium innocuum		99.69	0.47	MLZ 4			
Clostridium difficile		0.31	0.00				
45253412	**Good identification**						**45253412**
Clostridium innocuum		100.0	0.47	MLZ 4			
45253420	**Good identification**						**45253420**
Clostridium innocuum		100.0	0.36	NAG 1			
45253422	**Good identification**						**45253422**
Clostridium innocuum		100.0	0.36	NAG 1			
45253440	**Good identification**						**45253440**
Clostridium innocuum		100.0	0.36	SOR 1			
45253442	**Good identification**						**45253442**
Clostridium innocuum		100.0	0.36	SOR 1			
45253500	**Good identification**						**45253500**
Clostridium innocuum		100.0	0.36	LAC 1			
45253502	**Good identification**						**45253502**
Clostridium innocuum		100.0	0.36	LAC 1			
45253600	**Good identification**						**45253600**
Clostridium innocuum		100.0	0.36	RHA 1			
45253602	**Good identification**						**45253602**
Clostridium innocuum		100.0	0.36	RHA 1			
45254000	**Low discrimination**						**45254000**
					GRA	SPO	
Fusobacterium mortiferum		79.76	0.46	XYL 7	1	1	
Clostridium difficile		18.33	0.19		99	99	
Clostridium innocuum		1.91	0.05		99	99	
45254100	**Good identification**						**45254100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.52	XYL 7			
45254102	**Good identification**						**45254102**
Fusobacterium mortiferum		100.0	0.35	XYL 7			
45255000	**Low discrimination**						**45255000**
					GRA	SPO	
Fusobacterium mortiferum		70.95	0.49	XYL 7	1	1	
Clostridium innocuum		21.51	0.27		99	99	
Lactobacillus jensenii		5.45	0.19		99	1	
45255100	**Acceptable identification**						**45255100**
					GRA		
Fusobacterium mortiferum		94.54	0.55	XYL 7	1		
Lactobacillus catenaformis		1.79	0.15		99		
Collinsella aerofaciens		1.18	-.03		99		
45255102	**Low discrimination**						**45255102**
					GRA	SPO	
Fusobacterium mortiferum		72.42	0.38	XYL 7	1	1	
Clostridium barati		15.27	0.12	BGL 10	99	99	
Lactobacillus catenaformis		12.31	0.15		99	1	
45255120	**Acceptable identification**						**45255120**
Clostridium septicum		90.91	0.37	MLT 94			
Clostridium paraputrificum		5.56	0.14	XYL 1			
Clostridium barati		2.16	0.12				
45256000	**Low discrimination**						**45256000**
					SPO		
Clostridium innocuum		53.07	0.39	GAL 95	99		
Clostridium difficile		46.32	0.35	XYL 5	99		
Eubacterium limosum		0.38	-.03		1		
45256002	**Good identification**						**45256002**
Clostridium innocuum		99.13	0.39	GAL 95			
Clostridium difficile		0.87	0.01	XYL 5			
45256010	**Good identification**						**45256010**
Clostridium difficile		99.03	0.47	CEL 5			
Clostridium innocuum		0.97	0.16	XYL 5			
45257000	**Very good identification**						**45257000**
					GRA	SPO	
Clostridium innocuum		99.92	0.61	XYL 5	99	99	
Clostridium difficile		0.05	0.01		99	99	
Fusobacterium mortiferum		0.03	0.14		1	1	
45257002	**Very good identification**						**45257002**
Clostridium innocuum		100.0	0.61	XYL 5			
45257010	**Acceptable identification**						**45257010**
Clostridium innocuum		94.84	0.38	XYL 5			
Clostridium difficile		5.16	0.13	MLZ 4			
45257012	**Good identification**						**45257012**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium innocuum		100.0	0.38	XYL 5				
45257400	**Good identification**							**45257400**
Clostridium innocuum		100.0	0.48	XYL 5				
45257402	**Good identification**							**45257402**
Clostridium innocuum		100.0	0.48	XYL 5				
45261120	**Good identification**							**45261120**
Clostridium septicum		99.46	0.45	MLT 94				
Clostridium paraputrificum		0.54	0.04	FRU 99				
45263000	**Good identification**							**45263000**
Clostridium innocuum		100.0	0.54	FRU 95				
45263002	**Good identification**							**45263002**
Clostridium innocuum		100.0	0.54	FRU 95				
45263400	**Good identification**							**45263400**
Clostridium innocuum		100.0	0.41	FRU 95				
45263402	**Good identification**							**45263402**
Clostridium innocuum		100.0	0.41	FRU 95				
45270000	**Low discrimination**							**45270000**
					GRA	SPO		
Fusobacterium mortiferum		82.63	0.56		1	1		
Lactobacillus jensenii		7.20	0.28		99	1		
Clostridium difficile		6.84	0.21		99	99		
45270002	**Low discrimination**							**45270002**
					GRA	SPO		
Fusobacterium mortiferum		52.27	0.39	BGL 10	1	1		
Lactobacillus jensenii		27.33	0.25		99	1		
Clostridium innocuum		18.94	0.19		99	99		
45270100	**Very good identification**							**45270100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.58	0.62		1	1	1	
Clostridium septicum		1.11	0.11		1	99	99	
Atopobium parvulum		0.17	0.04		99	99	1	
45270102	**Acceptable identification**							**45270102**
					GRA	SPO		
Fusobacterium mortiferum		94.38	0.46	BGL 10	1	1		
Clostridium barati		3.41	0.06		99	99		
Lactobacillus acidophilus		2.22	0.05		99	1		
45270120	**Good identification**							**45270120**
					GRA	SPO		
Clostridium septicum		98.23	0.45	MLT 94	99	99		
Fusobacterium mortiferum		0.89	0.28	GAL 99	1	1		
Clostridium paraputrificum		0.53	0.04		99	99		
45271000	**Low discrimination**							**45271000**
					GRA	SPO		
Lactobacillus jensenii		78.76	0.62	SUC 90	99	1		
Fusobacterium mortiferum		13.69	0.59	MLT 95	1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium innocuum		6.98	0.41		99	99		
45271002	**Low discrimination**							**45271002**
					GRA	SPO		
Lactobacillus jensenii		84.51	0.59	SUC 90	99	1		
Clostridium innocuum		11.24	0.41	MLT 95	99	99		
Fusobacterium mortiferum		2.45	0.42		1	1		
45271010	**Good identification**							**45271010**
					GRA	SPO		
Lactobacillus jensenii		98.25	0.54	SUC 90	99	1		
Clostridium innocuum		1.09	0.18	MLT 95	99	99		
Fusobacterium mortiferum		0.52	0.25		1	1		
45271012	**Good identification**							**45271012**
					GRA	SPO		
Lactobacillus jensenii		97.95	0.51	SUC 90	99	1		
Clostridium innocuum		1.63	0.18	MLT 95	99	99		
Leptotrichia buccalis		0.43	-.03		1	1		
45271020	**Low discrimination**							**45271020**
					SPO			
Clostridium septicum		53.59	0.45	MLT 94	99			
Lactobacillus jensenii		44.37	0.52	LAC 99	1			
Clostridium barati		1.22	0.20		99			
45271022	**Low discrimination**							**45271022**
					SPO			
Lactobacillus jensenii		82.74	0.49	SUC 90	1			
Clostridium barati		10.23	0.28	MLT 95	99			
Clostridium septicum		3.06	0.17		99			
45271030	**Good identification**							**45271030**
					SPO			
Lactobacillus jensenii		96.47	0.43	SUC 90	1			
Clostridium septicum		3.53	0.11	MLT 95	99			
45271032	**Good identification**							**45271032**
Lactobacillus jensenii		100.0	0.40	SUC 90				
45271100	**Low discrimination**							**45271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		52.13	0.65		1	1	1	
Clostridium septicum		38.80	0.45		1	99	99	
Atopobium parvulum		6.07	0.38		99	99	1	
45271102	**Low discrimination**							**45271102**
					GRA	SPO		
Lactobacillus acidophilus		43.31	0.39	SUC 99	99	1		
Fusobacterium mortiferum		27.96	0.49	MLT 94	1	1		
Clostridium barati		12.79	0.28		99	99		
45271120	**Very good identification**							**45271120**
Clostridium septicum		99.34	0.80	MLT 94				
Clostridium paraputrificum		0.54	0.38					
Clostridium barati		0.07	0.28					
45271121	**Good identification**							**45271121**
Clostridium septicum		99.46	0.45	MLT 94				
Clostridium paraputrificum		0.54	0.04	URE 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45271122	**Low discrimination**						**45271122**
					SPO		
Clostridium septicum		74.51	0.51	MLT 94	99		
Lactobacillus acidophilus		8.53	0.39	BGL 2	1		
Clostridium paraputrificum		8.48	0.32		99		
45271130	**Good identification**						**45271130**
					SPO		
Clostridium septicum		99.19	0.45	MLT 94	99		
Clostridium paraputrificum		0.54	0.04	MLZ 1	99		
Lactobacillus jensenii		0.28	0.09		1		
45271160	**Good identification**						**45271160**
Clostridium septicum		95.57	0.45	MLT 94			
Clostridium paraputrificum		3.86	0.19	SOR 1			
Clostridium barati		0.57	0.10				
45271320	**Good identification**						**45271320**
Clostridium septicum		98.74	0.45	MLT 94			
Clostridium barati		0.59	0.10	RHA 1			
Clostridium paraputrificum		0.54	0.04				
45271520	**Good identification**						**45271520**
Clostridium septicum		99.46	0.45	MLT 94			
Clostridium paraputrificum		0.54	0.04	ARA 1			
45272000	**Low discrimination**						**45272000**
					GRA	SPO	
Clostridium innocuum		83.99	0.54	GAL 95	99	99	
Clostridium difficile		15.71	0.38		99	99	
Fusobacterium mortiferum		0.21	0.22		1	1	
45272002	**Good identification**						**45272002**
Clostridium innocuum		99.81	0.54	GAL 95			
Clostridium difficile		0.19	0.04				
45272010	**Good identification**						**45272010**
Clostridium difficile		95.64	0.50	TRE 7			
Clostridium innocuum		4.37	0.30	CEL 5			
45272400	**Good identification**						**45272400**
Clostridium innocuum		98.94	0.41	GAL 95			
Clostridium difficile		1.06	0.04				
45272402	**Good identification**						**45272402**
Clostridium innocuum		100.0	0.41	GAL 95			
45273000	**Very good identification**						**45273000**
					GRA	SPO	
Clostridium innocuum		99.86	0.75		99	99	
Lactobacillus jensenii		0.12	0.28		99	1	
Fusobacterium mortiferum		0.02	0.25		1	1	
45273001	**Good identification**						**45273001**
Clostridium innocuum		100.0	0.41	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45273002	**Very good identification**						**45273002**
					SPO		
Clostridium innocuum		99.91	0.75		99		
Lactobacillus jensenii		0.08	0.25		1		
Clostridium barati		0.01	0.01		99		
45273003	**Good identification**						**45273003**
Clostridium innocuum		100.0	0.41	URE 1			
45273010	**Good identification**						**45273010**
					SPO		
Clostridium innocuum		97.96	0.52	MLZ 4	99		
Clostridium difficile		1.14	0.15		99		
Lactobacillus jensenii		0.90	0.19		1		
45273012	**Good identification**						**45273012**
					SPO		
Clostridium innocuum		99.39	0.52	MLZ 4	99		
Lactobacillus jensenii		0.61	0.16		1		
45273020	**Acceptable identification**						**45273020**
					SPO		
Clostridium innocuum		93.49	0.41	NAG 1	99		
Clostridium septicum		3.22	0.11		99		
Lactobacillus jensenii		2.66	0.17		1		
45273022	**Good identification**						**45273022**
					SPO		
Clostridium innocuum		95.90	0.41	NAG 1	99		
Clostridium barati		1.94	0.10		99		
Lactobacillus jensenii		1.82	0.14		1		
45273040	**Good identification**						**45273040**
Clostridium innocuum		100.0	0.41	SOR 1			
45273042	**Good identification**						**45273042**
Clostridium innocuum		100.0	0.41	SOR 1			
45273100	**Acceptable identification**						**45273100**
					GRA	SPO	
Clostridium innocuum		90.55	0.41	LAC 1	99	99	
Fusobacterium mortiferum		4.18	0.31		1	1	
Clostridium septicum		3.11	0.11		99	99	
45273102	**Acceptable identification**						**45273102**
					SPO		
Clostridium innocuum		90.76	0.41	LAC 1	99		
Lactobacillus acidophilus		3.76	0.17		1		
Clostridium tertium		3.18	0.17		99		
45273120	**Good identification**						**45273120**
Clostridium septicum		97.03	0.45	MLT 94			
Clostridium tertium		1.00	0.17	MAN 1			
Clostridium ramosum		0.58	0.11				
45273200	**Good identification**						**45273200**
Clostridium innocuum		100.0	0.41	RHA 1			
45273202	**Good identification**						**45273202**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium innocuum		100.0	0.41	RHA 1				
45273400	**Very good identification**							**45273400**
Clostridium innocuum		100.0	0.63					
45273402	**Very good identification**							**45273402**
Clostridium innocuum		100.0	0.63					
45273410	**Good identification**							**45273410**
Clostridium innocuum		100.0	0.39	MLZ 4				
45273412	**Good identification**							**45273412**
Clostridium innocuum		100.0	0.39	MLZ 4				
45274000	**Acceptable identification**							**45274000**
					GRA	SPO		
Fusobacterium mortiferum		94.53	0.37	XYL 7	1	1		
Clostridium difficile		5.47	-.01		99	99		
45274100	**Good identification**							**45274100**
					COC	GRA		
Fusobacterium mortiferum		95.83	0.43	XYL 7	1	1		
Atopobium parvulum		4.17	0.08		99	99		
45275000	**Low discrimination**							**45275000**
					GRA	SPO		
Fusobacterium mortiferum		45.53	0.40	XYL 7	1	1		
Lactobacillus jensenii		35.15	0.28		99	1		
Clostridium innocuum		16.24	0.19		99	99		
45275100	**Low discrimination**							**45275100**
					COC	GRA	SPO	
Atopobium parvulum		72.11	0.42	SUC 96	99	99	1	
Fusobacterium mortiferum		25.11	0.46	MLT 99	1	1	1	
Clostridium septicum		2.51	0.11		1	99	99	
45275120	**Acceptable identification**							**45275120**
					COC	SPO		
Clostridium septicum		94.55	0.45	MLT 94	1	99		
Atopobium parvulum		4.85	0.29	XYL 1	99	1		
Clostridium paraputrificum		0.51	0.04		1	99		
45275322	**Good identification**							**45275322**
Parabacteroides distasonis		100.0	0.42	RAF 99				
45275332	**Good identification**							**45275332**
Parabacteroides distasonis		100.0	0.37	RAF 99				
45275722	**Good identification**							**45275722**
Parabacteroides distasonis		100.0	0.38	RAF 99				
45277000	**Good identification**							**45277000**
					SPO			
Clostridium innocuum		99.94	0.54	XYL 5	99			
Actinomyces israelii		0.06	0.14		1			
45277002	**Good identification**							**45277002**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		99.90	0.54	XYL 5	99		
Actinomyces israelii		0.10	0.17		1		
45277400	**Good identification**						**45277400**
Clostridium innocuum		100.0	0.41	XYL 5			
45277402	**Good identification**						**45277402**
					SPO		
Clostridium innocuum		99.56	0.41	XYL 5	99		
Actinomyces israelii		0.45	0.15		1		
45300000	**Good identification**						**45300000**
					GRA	SPO	
Clostridium bifermentans		92.46	0.67	IND 90	99	99	
Clostridium botulinum biovar A		4.32	0.45	SAL 10	99	99	
Prevotella disiens		1.54	0.32		1	1	
45300020	**Good identification**						**45300020**
					COC	SPO	
Clostridium bifermentans		96.57	0.47	IND 90	1	99	
Clostridium novyi biovar A		0.90	0.25	SAL 10	1	99	
Gemella morbillorum		0.72	0.14	NAG 6	99	1	
45300040	**Low discrimination**						**45300040**
Clostridium bifermentans		72.37	0.44	IND 90			
Clostridium botulinum biovar A		24.22	0.36	SAL 10			
Clostridium novyi biovar A		2.76	0.32	SOR 4			
45301120	**Acceptable identification**						**45301120**
					GRA	SPO	
Clostridium septicum		93.55	0.49	FRU 99	99	99	
Prevotella bivia		3.96	0.19		1	1	
Clostridium perfringens		1.20	0.23		99	84	
45310000	**Low discrimination**						**45310000**
					GRA	SPO	
Clostridium bifermentans		39.80	0.45	IND 90	99	99	
Fusobacterium mortiferum		31.18	0.57	SAL 10	1	1	
Prevotella disiens		12.60	0.32	FRU 5	1	1	
45310002	**Low discrimination**						**45310002**
					GRA	SPO	
Fusobacterium mortiferum		67.95	0.41	MLT 11	1	1	
Lactobacillus jensenii		9.54	0.16	BGL 10	99	1	
Clostridium sporogenes		9.12	0.11		99	99	
45310010	**Good identification**						**45310010**
					GRA	SPO	
Clostridium difficile		95.26	0.40	MLT 1	99	99	
Clostridium bifermentans		1.48	0.11	MAN 90	99	99	
Fusobacterium mortiferum		1.16	0.23		1	1	
45310100	**Very good identification**						**45310100**
					GRA	SPO	
Fusobacterium mortiferum		97.46	0.64	MLT 11	1	1	
Clostridium septicum		1.46	0.15		99	99	
Clostridium bifermentans		0.54	0.11		99	99	
45310102	**Acceptable identification**						**45310102**
					GRA	SPO	
Fusobacterium mortiferum		93.98	0.47	MLT 11	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis		3.62	0.06	BGL 10	99	1	
Clostridium barati		1.55	0.02		99	99	
45310120	**Acceptable identification**						**45310120**
					GRA	SPO	
Clostridium septicum		94.29	0.49	GAL 99	99	99	
Prevotella bivia		2.51	0.15		1	1	
Clostridium perfringens		1.21	0.23		99	84	
45311000	**Low discrimination**						**45311000**
					GRA		
Lactobacillus jensenii		56.83	0.54	SUC 90	99		
Fusobacterium mortiferum		36.79	0.60	CEL 95	1		
Lactobacillus catenaformis		1.98	0.29		99		
45311002	**Low discrimination**						**45311002**
					GRA		
Lactobacillus jensenii		83.29	0.51	SUC 90	99		
Fusobacterium mortiferum		8.99	0.44	CEL 95	1		
Lactobacillus catenaformis		4.34	0.29		99		
45311010	**Good identification**						**45311010**
					GRA	SPO	
Lactobacillus jensenii		97.04	0.45	SUC 90	99	1	
Fusobacterium mortiferum		1.90	0.26	CEL 95	1	1	
Clostridium difficile		1.05	0.06		99	99	
45311012	**Good identification**						**45311012**
Lactobacillus jensenii		100.0	0.42	SUC 90			
45311020	**Low discrimination**						**45311020**
					SPO		
Clostridium septicum		72.63	0.49	LAC 99	99		
Clostridium perfringens		12.54	0.43		84		
Lactobacillus jensenii		12.15	0.43		1		
45311022	**Low discrimination**						**45311022**
					GRA		
Lactobacillus jensenii		33.28	0.40	SUC 90	99		
Prevotella oralis		24.37	0.34	CEL 95	1		
Clostridium perfringens		24.24	0.37		99		
45311030	**Low discrimination**						**45311030**
					SPO		
Lactobacillus jensenii		82.48	0.35	SUC 90	1		
Clostridium septicum		14.94	0.15	CEL 95	99		
Clostridium perfringens		2.58	0.09		84		
45311100	**Low discrimination**						**45311100**
					GRA	SPO	
Fusobacterium mortiferum		41.47	0.67	MLT 11	1	1	
Clostridium septicum		41.01	0.49		99	99	
Collinsella aerofaciens		10.41	0.32		99	1	
45311102	**Low discrimination**						**45311102**
					GRA		
Bifidobacterium longum subsp. infantis		46.37	0.40	SUC 99	99		
Fusobacterium mortiferum		18.27	0.50	RAF 95	1		
Prevotella oralis		10.95	0.32		1		
45311120	**Very good identification**						**45311120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		97.35	0.83				
Clostridium perfringens		1.25	0.58				
Clostridium paraputrificum		1.20	0.48				
45311121	**Good identification**						**45311121**
Clostridium septicum		96.58	0.49	URE 1			
Clostridium perfringens		1.24	0.23				
Clostridium paraputrificum		1.19	0.14				
45311122	**Low discrimination**						**45311122**
					GRA	SPO	
Clostridium septicum		40.18	0.54	BGL 2	99	99	
Prevotella oralis		25.45	0.54		1	1	
Clostridium perfringens		11.85	0.52		99	84	
45311130	**Good identification**						**45311130**
Clostridium septicum		97.43	0.49	MLZ 1			
Clostridium perfringens		1.25	0.23				
Clostridium paraputrificum		1.20	0.14				
45311160	**Low discrimination**						**45311160**
Clostridium septicum		74.85	0.49	SOR 1			
Clostridium perfringens		18.05	0.45				
Clostridium paraputrificum		6.90	0.29				
45311162	**Low discrimination**						**45311162**
Clostridium perfringens		57.01	0.40	SUC 91			
Clostridium paraputrificum		19.83	0.22				
Clostridium septicum		10.25	0.20				
45311320	**Acceptable identification**						**45311320**
					GRA	SPO	
Clostridium septicum		94.11	0.49	RHA 1	99	99	
Prevotella oralis		2.72	0.26		1	1	
Clostridium perfringens		1.20	0.23		99	84	
45311322	**Acceptable identification**						**45311322**
					GRA	SPO	
Prevotella oralis		91.54	0.48	SUC 92	1	1	
Clostridium septicum		3.41	0.20	RAF 92	99	99	
Parabacteroides distasonis		1.12	0.28		1	1	
45311520	**Good identification**						**45311520**
Clostridium septicum		97.25	0.49	ARA 1			
Clostridium perfringens		1.24	0.23				
Clostridium paraputrificum		1.20	0.14				
45312000	**Low discrimination**						**45312000**
					SPO		
Clostridium difficile		80.69	0.45	MLT 1	99		
Eubacterium limosum		17.88	0.32		1		
Clostridium bifermentans		0.68	0.11		99		
45312010	**Very good identification**						**45312010**
					SPO		
Clostridium difficile		99.95	0.56	MLT 1	99		
Eubacterium limosum		0.05	-.03		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45313120	**Good identification**							**45313120**
Clostridium septicum		97.17	0.49	MAN 1				
Clostridium perfringens		1.24	0.23					
Clostridium paraputrificum		1.20	0.14					
45314000	**Low discrimination**							**45314000**
					COC	GRA	SPO	
Fusobacterium mortiferum		59.48	0.38	MLT 11	1	1	1	
Gemella morbillorum		11.10	0.15	XYL 7	99	99	1	
Clostridium bifermentans		10.19	0.11		1	99	99	
45314100	**Good identification**							**45314100**
					COC	GRA		
Fusobacterium mortiferum		98.52	0.44	MLT 11	1	1		
Atopobium parvulum		1.48	0.02	XYL 7	99	99		
45315000	**Low discrimination**							**45315000**
					COC	GRA		
Fusobacterium mortiferum		77.07	0.41	MLT 11	1	1		
Lactobacillus jensenii		15.97	0.19	XYL 7	1	99		
Gemella morbillorum		5.16	0.11		99	99		
45315100	**Low discrimination**							**45315100**
					COC	GRA		
Fusobacterium mortiferum		34.36	0.47	MLT 11	1	1		
Atopobium parvulum		34.07	0.36	XYL 7	99	99		
Eubacterium contortum		15.07	0.29		1	99		
45315102	**Low discrimination**							**45315102**
					COC	GRA		
Bifidobacterium longum subsp. infantis		83.86	0.38	SUC 99	1	99		
Atopobium parvulum		4.79	0.23	RAF 95	99	99		
Fusobacterium mortiferum		3.04	0.31		1	1		
45315120	**Acceptable identification**							**45315120**
					COC	SPO		
Clostridium septicum		91.08	0.49	XYL 1	1	99		
Bifidobacterium longum subsp. infantis		3.74	0.25		1	1		
Atopobium parvulum		1.21	0.23		99	1		
45315122	**Low discrimination**							**45315122**
					GRA	SPO		
Bifidobacterium longum subsp. infantis		65.64	0.43	SUC 99	99	1		
Prevotella oralis		25.02	0.39	RAF 95	1	1		
Clostridium septicum		3.23	0.20		99	99		
45315300	**Very good identification**							**45315300**
					COC	GRA		
Eubacterium contortum		99.95	0.63	SUC 99	1	99		
Fusobacterium mortiferum		0.02	0.13		1	1		
Atopobium parvulum		0.02	0.02		99	99		
45315302	**Good identification**							**45315302**
					GRA			
Eubacterium contortum		98.89	0.50	SUC 99	99			
Prevotella oralis		0.61	0.11		1			
Bifidobacterium longum subsp. infantis		0.40	0.04		99			
45315320	**Acceptable identification**							**45315320**
					GRA	SPO		
Eubacterium contortum		92.77	0.50	SUC 99	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		4.96	0.21		1	1	
Clostridium septicum		1.61	0.15		99	99	
45315322	**Low discrimination**						**45315322**
					GRA		
Eubacterium contortum		43.78	0.37	SUC 99	99		
Prevotella oralis		28.84	0.32		1		
Parabacteroides distasonis		25.60	0.44		1		
45315332	**Acceptable identification**						**45315332**
					GRA		
Parabacteroides distasonis		94.95	0.40	RAF 99	1		
Eubacterium contortum		3.05	0.03		99		
Prevotella oralis		2.01	-.02		1		
45315502	**Low discrimination**						**45315502**
Bifidobacterium adolescentis		86.26	0.41	SUC 99			
Eubacterium contortum		8.08	0.14	RAF 92			
Bifidobacterium longum subsp. infantis		3.95	0.04				
45315512	**Good identification**						**45315512**
Bifidobacterium adolescentis		98.70	0.42	SUC 99			
Clostridium clostridioforme		1.31	0.07	RAF 92			
45315700	**Very good identification**						**45315700**
					GRA		
Eubacterium contortum		99.85	0.61	SUC 99	99		
Bacteroides vulgatus		0.16	0.06		1		
45315702	**Good identification**						**45315702**
Eubacterium contortum		99.72	0.48	SUC 99			
Clostridium clostridioforme		0.28	0.10				
45315720	**Low discrimination**						**45315720**
					GRA		
Eubacterium contortum		53.53	0.48	SUC 99	99		
Bacteroides vulgatus		46.47	0.40		1		
45315722	**Low discrimination**						**45315722**
					GRA		
Eubacterium contortum		66.15	0.36	SUC 99	99		
Parabacteroides distasonis		28.97	0.41		1		
Bacteroides vulgatus		3.29	0.06		1		
45315732	**Low discrimination**						**45315732**
					GRA		
Parabacteroides distasonis		74.07	0.36	RAF 99	1		
Clostridium clostridioforme		22.76	0.22		50		
Eubacterium contortum		3.17	0.01		99		
45317502	**Low discrimination**						**45317502**
Bifidobacterium adolescentis		87.89	0.39	SUC 99			
Actinomyces israelii		12.11	0.31	RAF 92			
45317512	**Good identification**						**45317512**
Bifidobacterium adolescentis		98.34	0.41	SUC 99			
Actinomyces israelii		1.66	0.17	RAF 92			
45321120	**Very good identification**						**45321120**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Clostridium septicum		98.87	0.57	FRU 99	1	99		
Clostridium perfringens		0.78	0.28		1	84		
Atopobium parvulum		0.24	0.18		99	1		
45325100	**Good identification**							**45325100**
Atopobium parvulum		100.0	0.36	SUC 96				
45330000	**Low discrimination**							**45330000**
					GRA	SPO		
Fusobacterium mortiferum		69.20	0.48	MLT 11	1	1		
Lactobacillus jensenii		16.26	0.28		99	1		
Clostridium botulinum biovar A		4.19	0.14		99	99		
45330100	**Low discrimination**							**45330100**
					COC	GRA	SPO	
Fusobacterium mortiferum		73.75	0.55	MLT 11	1	1	1	
Atopobium parvulum		14.88	0.31		99	99	1	
Clostridium septicum		11.10	0.23		1	99	99	
45330102	**Low discrimination**							**45330102**
					COC	GRA	SPO	
Fusobacterium mortiferum		74.18	0.38	MLT 11	1	1	1	
Atopobium parvulum		23.78	0.18	BGL 10	99	99	1	
Clostridium septicum		2.05	-.06		1	99	99	
45330120	**Very good identification**							**45330120**
					COC	SPO		
Clostridium septicum		98.81	0.57	GAL 99	1	99		
Clostridium perfringens		0.78	0.28		1	84		
Atopobium parvulum		0.24	0.18		99	1		
45331000	**Good identification**							**45331000**
					COC	GRA		
Lactobacillus jensenii		90.52	0.62	SUC 90	1	99		
Fusobacterium mortiferum		5.84	0.51	CEL 95	1	1		
Atopobium parvulum		1.83	0.31		99	99		
45331002	**Very good identification**							**45331002**
					COC	GRA		
Lactobacillus jensenii		97.85	0.59	SUC 90	1	99		
Fusobacterium mortiferum		1.05	0.35	CEL 95	1	1		
Atopobium parvulum		0.52	0.18		99	99		
45331010	**Good identification**							**45331010**
					GRA			
Lactobacillus jensenii		99.81	0.54	SUC 90	99			
Fusobacterium mortiferum		0.20	0.17	CEL 95	1			
45331012	**Good identification**							**45331012**
					GRA			
Lactobacillus jensenii		99.57	0.51	SUC 90	99			
Leptotrichia buccalis		0.43	-.03	CEL 95	1			
45331020	**Low discrimination**							**45331020**
					SPO			
Clostridium septicum		78.22	0.57	LAC 99	99			
Lactobacillus jensenii		13.09	0.52		1			
Clostridium perfringens		8.36	0.47		84			
45331022	**Low discrimination**							**45331022**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus jensenii		59.91	0.49	SUC 90	1			
Clostridium perfringens		27.01	0.42	CEL 95	84			
Clostridium septicum		10.96	0.28		99			
45331030	**Low discrimination**							**45331030**
					SPO			
Lactobacillus jensenii		83.30	0.43	SUC 90	1			
Clostridium septicum		15.09	0.23	CEL 95	99			
Clostridium perfringens		1.61	0.13		84			
45331032	**Good identification**							**45331032**
					SPO			
Lactobacillus jensenii		98.12	0.40	SUC 90	1			
Clostridium perfringens		1.34	0.08	CEL 95	84			
Clostridium septicum		0.54	-.06		99			
45331060	**Low discrimination**							**45331060**
					SPO			
Clostridium perfringens		63.33	0.35	SUC 91	84			
Clostridium septicum		31.42	0.23	LAC 88	99			
Lactobacillus jensenii		5.26	0.17		1			
45331100	**Low discrimination**							**45331100**
					COC	GRA	SPO	
Atopobium parvulum		54.18	0.66	SUC 96	99	99	1	
Clostridium septicum		40.40	0.57		1	99	99	
Fusobacterium mortiferum		4.07	0.58		1	1	1	
45331102	**Low discrimination**							**45331102**
					COC	SPO		
Atopobium parvulum		79.90	0.53	SUC 96	99	1		
Clostridium septicum		6.89	0.28		1	99		
Clostridium perfringens		3.85	0.31		1	84		
45331120	**Very good identification**							**45331120**
					COC	SPO		
Clostridium septicum		98.87	0.91		1	99		
Clostridium perfringens		0.78	0.62		1	84		
Atopobium parvulum		0.24	0.53		99	1		
45331121	**Very good identification**							**45331121**
					COC	SPO		
Clostridium septicum		98.81	0.57	URE 1	1	99		
Clostridium perfringens		0.78	0.28		1	84		
Atopobium parvulum		0.24	0.18		99	1		
45331122	**Low discrimination**							**45331122**
Clostridium septicum		80.13	0.63	BGL 2				
Clostridium perfringens		14.63	0.57					
Clostridium paraputrificum		1.84	0.32					
45331130	**Very good identification**							**45331130**
					COC	SPO		
Clostridium septicum		98.82	0.57	MLZ 1	1	99		
Clostridium perfringens		0.78	0.28		1	84		
Atopobium parvulum		0.24	0.18		99	1		
45331160	**Low discrimination**							**45331160**
Clostridium septicum		86.22	0.57	SOR 1				
Clostridium perfringens		12.87	0.50					

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium paraputrificum		0.70	0.19					
45331162	**Low discrimination**							**45331162**
Clostridium perfringens		73.36	0.44	SUC 91				
Clostridium septicum		21.31	0.28					
Clostridium paraputrificum		3.65	0.12					
45331320	**Very good identification**							**45331320**
					COC	SPO		
Clostridium septicum		98.79	0.57	RHA 1	1	99		
Clostridium perfringens		0.78	0.28		1	84		
Atopobium parvulum		0.24	0.18		99	1		
45331520	**Very good identification**							**45331520**
					COC	SPO		
Clostridium septicum		98.87	0.57	ARA 1	1	99		
Clostridium perfringens		0.78	0.28		1	84		
Atopobium parvulum		0.24	0.18		99	1		
45331722	**Acceptable identification**							**45331722**
					GRA	SPO		
Parabacteroides distasonis		92.15	0.36	RAF 99	1	1		
Clostridium clostridioforme		4.80	0.08	XYL 90	50	66		
Clostridium septicum		3.05	-.06		99	99		
45332010	**Good identification**							**45332010**
Clostridium difficile		100.0	0.37	MLT 1				
45333120	**Very good identification**							**45333120**
					COC	SPO		
Clostridium septicum		98.51	0.57	MAN 1	1	99		
Clostridium perfringens		0.78	0.28		1	84		
Atopobium parvulum		0.24	0.18		99	1		
45334100	**Low discrimination**							**45334100**
					COC	GRA		
Atopobium parvulum		83.28	0.36	SUC 96	99	99		
Fusobacterium mortiferum		16.72	0.35	GAL 99	1	1		
45335000	**Low discrimination**							**45335000**
					COC	GRA		
Atopobium parvulum		71.01	0.36	SUC 96	99	99		
Lactobacillus jensenii		19.08	0.28	LAC 99	1	99		
Fusobacterium mortiferum		9.17	0.32		1	1		
45335100	**Very good identification**							**45335100**
					COC	GRA	SPO	
Atopobium parvulum		99.24	0.70	SUC 96	99	99	1	
Clostridium septicum		0.40	0.23		1	99	99	
Fusobacterium mortiferum		0.30	0.38		1	1	1	
45335101	**Good identification**							**45335101**
Atopobium parvulum		100.0	0.36	SUC 96				
45335102	**Very good identification**							**45335102**
					COC			
Atopobium parvulum		98.78	0.57	SUC 96	99			
Bifidobacterium adolescentis		0.38	0.23		1			
Actinomyces israelii		0.35	0.29		1			
45335110	**Good identification**							**45335110**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		100.0	0.36	SUC 96				
45335120	**Low discrimination**							**45335120**
					COC	SPO		
Clostridium septicum		69.02	0.57	XYL 1	1	99		
Atopobium parvulum		30.34	0.57		99	1		
Clostridium perfringens		0.55	0.28		1	84		
45335122	**Low discrimination**							**45335122**
					COC	GRA	SPO	
Atopobium parvulum		56.46	0.44	SUC 96	99	99	1	
Parabacteroides distasonis		22.17	0.47		1	1	1	
Clostridium septicum		14.85	0.28		1	99	99	
45335132	**Acceptable identification**							**45335132**
					COC	GRA	SPO	
Parabacteroides distasonis		94.31	0.43	RAF 99	1	1	1	
Atopobium parvulum		4.51	0.10		99	99	1	
Clostridium septicum		1.19	-.06		1	99	99	
45335140	**Good identification**							**45335140**
					COC			
Atopobium parvulum		98.41	0.36	SUC 96	99			
Actinomyces israelii		1.59	0.19	SOR 1	1			
45335300	**Low discrimination**							**45335300**
					COC			
Atopobium parvulum		69.34	0.36	SUC 96	99			
Eubacterium contortum		30.66	0.29	RHA 1	1			
45335322	**Good identification**							**45335322**
					COC	GRA		
Parabacteroides distasonis		96.83	0.56	RAF 99	1	1		
Prevotella oralis		1.53	0.12		1	1		
Atopobium parvulum		0.83	0.10		99	99		
45335332	**Good identification**							**45335332**
Parabacteroides distasonis		98.10	0.51	RAF 99				
Clostridium clostridioforme		1.90	0.16					
45335500	**Low discrimination**							**45335500**
					COC	SPO		
Atopobium parvulum		89.86	0.36	SUC 96	99	1		
Clostridium butyricum		7.08	0.10	ARA 1	1	99		
Actinomyces israelii		3.06	0.25		1	1		
45335502	**Low discrimination**							**45335502**
					COC			
Bifidobacterium adolescentis		61.60	0.38	SUC 99	1			
Atopobium parvulum		23.91	0.23	RAF 92	99			
Actinomyces israelii		6.92	0.28		1			
45335512	**Low discrimination**							**45335512**
Bifidobacterium adolescentis		86.27	0.39	SUC 99				
Clostridium clostridioforme		12.55	0.22	RAF 92				
Actinomyces israelii		1.18	0.14					
45335522	**Acceptable identification**							**45335522**
					COC	GRA		
Parabacteroides distasonis		92.86	0.44	RAF 99	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		3.90	0.10		99	99	
Clostridium clostridioforme		2.22	0.10		1	50	
45335532	**Low discrimination**						**45335532**
Parabacteroides distasonis		83.17	0.39	RAF 99			
Clostridium clostridioforme		16.83	0.22				
45335712	**Good identification**						**45335712**
Clostridium clostridioforme		98.00	0.36	SUC 99			
Parabacteroides distasonis		2.00	0.13	RAF 94			
45335722	**Acceptable identification**						**45335722**
					GRA		
Parabacteroides distasonis		94.03	0.52	RAF 99	1		
Clostridium clostridioforme		5.50	0.25		50		
Eubacterium contortum		0.48	0.01		99		
45335732	**Low discrimination**						**45335732**
Parabacteroides distasonis		66.91	0.47	RAF 99			
Clostridium clostridioforme		33.09	0.36				
45337100	**Low discrimination**						**45337100**
					COC	SPO	
Atopobium parvulum		78.51	0.36	SUC 96	99	1	
Actinomyces israelii		18.50	0.39	MAN 1	1	1	
Clostridium tertium		2.99	0.11		1	99	
45337102	**Low discrimination**						**45337102**
					COC	SPO	
Actinomyces israelii		50.52	0.42	SUC 98	1	1	
Atopobium parvulum		25.22	0.23	CEL 93	99	1	
Clostridium tertium		16.33	0.19		1	99	
45337142	**Good identification**						**45337142**
Actinomyces israelii		100.0	0.35	SUC 98			
45337500	**Acceptable identification**						**45337500**
					COC	SPO	
Actinomyces israelii		91.09	0.38	SUC 98	1	1	
Atopobium parvulum		4.77	0.02	CEL 93	99	1	
Clostridium butyricum		4.14	-.07		1	99	
45337502	**Low discrimination**						**45337502**
Bifidobacterium adolescentis		56.26	0.36	SUC 99			
Actinomyces israelii		43.75	0.41	RAF 92			
45337512	**Low discrimination**						**45337512**
Bifidobacterium adolescentis		80.24	0.38	SUC 99			
Bifidobacterium dentium		12.13	0.06	RAF 92			
Actinomyces israelii		7.63	0.26				
45341000	**Low discrimination**						**45341000**
					SPO		
Lactobacillus jensenii		80.76	0.41	SUC 90	1		
Lactobacillus catenaformis		14.62	0.29	FRU 99	1		
Clostridium barati		1.64	0.07		99		
45341002	**Low discrimination**						**45341002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Lactobacillus jensenii		73.01	0.38	SUC 90	1	
Lactobacillus catenaformis		19.83	0.29	FRU 99	1	
Clostridium barati		6.67	0.15		99	
45341120	**Low discrimination**					**45341120**
					SPO	
Clostridium septicum		71.11	0.58	FRU 99	99	
Clostridium paraputrificum		27.48	0.48		99	
Collinsella aerofaciens		0.63	0.15		1	
45341122	**Low discrimination**					**45341122**
Clostridium paraputrificum		72.39	0.41	SUC 92		
Clostridium barati		9.72	0.32	FRU 99		
Clostridium septicum		8.92	0.29			
45350000	**Low discrimination**					**45350000**
					GRA	
Lactobacillus jensenii		47.29	0.41	SUC 90	99	
Fusobacterium mortiferum		35.45	0.49	GAL 99	1	
Lactobacillus catenaformis		8.56	0.29		99	
45350002	**Low discrimination**					**45350002**
					SPO	
Lactobacillus jensenii		53.21	0.38	SUC 90	1	
Clostridium barati		25.33	0.28	GAL 99	99	
Lactobacillus catenaformis		14.45	0.29		1	
45350022	**Low discrimination**					**45350022**
					SPO	
Clostridium barati		76.93	0.36	SUC 99	99	
Lactobacillus jensenii		13.47	0.28	GAL 95	1	
Lactobacillus catenaformis		3.66	0.18		1	
45350102	**Low discrimination**					**45350102**
					SPO	
Clostridium barati		54.07	0.36	SUC 99	99	
Lactobacillus catenaformis		15.42	0.32	GAL 95	1	
Collinsella aerofaciens		12.47	0.15		1	
45350120	**Low discrimination**					**45350120**
Clostridium septicum		69.53	0.58	GAL 99		
Clostridium paraputrificum		26.87	0.48			
Clostridium barati		2.69	0.36			
45350122	**Low discrimination**					**45350122**
Clostridium paraputrificum		51.36	0.41	SUC 92		
Clostridium barati		35.94	0.44	GAL 99		
Clostridium septicum		6.33	0.29			
45351000	**Low discrimination**					**45351000**
					SPO	
Lactobacillus jensenii		80.86	0.76	SUC 90	1	
Lactobacillus catenaformis		14.64	0.63		1	
Clostridium barati		1.64	0.41		99	
45351001	**Low discrimination**					**45351001**
					SPO	
Lactobacillus jensenii		81.12	0.41	SUC 90	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		14.69	0.29	URE 1	1	
Clostridium barati		1.65	0.07		99	
45351002	**Low discrimination**					**45351002**
					SPO	
Lactobacillus jensenii		72.57	0.73	SUC 90	1	
Lactobacillus catenaformis		19.71	0.63		1	
Clostridium barati		6.63	0.50		99	
45351003	**Low discrimination**					**45351003**
					SPO	
Lactobacillus jensenii		73.01	0.38	SUC 90	1	
Lactobacillus catenaformis		19.83	0.29	URE 1	1	
Clostridium barati		6.67	0.15		99	
45351010	**Very good identification**					**45351010**
					SPO	
Lactobacillus jensenii		99.30	0.67	SUC 90	1	
Lactobacillus catenaformis		0.55	0.29		1	
Clostridium barati		0.06	0.07		99	
45351012	**Very good identification**					**45351012**
					SPO	
Lactobacillus jensenii		98.83	0.64	SUC 90	1	
Lactobacillus catenaformis		0.81	0.29		1	
Clostridium barati		0.27	0.15		99	
45351020	**Low discrimination**					**45351020**
					SPO	
Lactobacillus jensenii		35.43	0.65	SUC 90	1	
Clostridium septicum		35.29	0.58		99	
Clostridium paraputrificum		13.64	0.48		99	
45351022	**Low discrimination**					**45351022**
					SPO	
Clostridium barati		38.98	0.58	SUC 99	99	
Lactobacillus jensenii		35.55	0.62		1	
Lactobacillus catenaformis		9.66	0.53		1	
45351030	**Good identification**					**45351030**
					SPO	
Lactobacillus jensenii		94.81	0.57	SUC 90	1	
Clostridium septicum		2.86	0.24		99	
Clostridium paraputrificum		1.11	0.14		99	
45351032	**Acceptable identification**					**45351032**
					SPO	
Lactobacillus jensenii		94.71	0.54	SUC 90	1	
Clostridium barati		3.15	0.24		99	
Lactobacillus catenaformis		0.78	0.18		1	
45351040	**Low discrimination**					**45351040**
					SPO	
Lactobacillus jensenii		71.71	0.41	SUC 90	1	
Lactobacillus catenaformis		12.99	0.29	SOR 1	1	
Clostridium barati		12.54	0.23		99	
45351042	**Low discrimination**					**45351042**
					SPO	
Lactobacillus jensenii		46.72	0.38	SUC 90	1	
Clostridium barati		36.75	0.31	SOR 1	99	
Lactobacillus catenaformis		12.69	0.29		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
45351062	**Low discrimination**					**45351062**
					SPO	
Clostridium barati		71.98	0.40	SUC 99	99	
Clostridium paraputrificum		14.06	0.22	SOR 8	99	
Lactobacillus jensenii		7.62	0.28		1	
45351100	**Low discrimination**					**45351100**
					SPO	
Collinsella aerofaciens		73.20	0.66	SUC 99	1	
Lactobacillus catenaformis		10.06	0.66		1	
Clostridium septicum		9.22	0.58		99	
45351102	**Low discrimination**					**45351102**
					SPO	
Lactobacillus catenaformis		33.71	0.66	SUC 99	1	
Collinsella aerofaciens		27.26	0.50		1	
Clostridium barati		22.69	0.58		99	
45351120	**Low discrimination**					**45351120**
					SPO	
Clostridium septicum		71.09	0.92		99	
Clostridium paraputrificum		27.47	0.82		99	
Collinsella aerofaciens		0.63	0.50		1	
45351121	**Low discrimination**					**45351121**
					SPO	
Clostridium septicum		71.10	0.58	URE 1	99	
Clostridium paraputrificum		27.48	0.48		99	
Collinsella aerofaciens		0.63	0.15		1	
45351122	**Low discrimination**					**45351122**
Clostridium paraputrificum		72.31	0.76	SUC 92		
Clostridium barati		9.71	0.66			
Clostridium septicum		8.91	0.63			
45351123	**Low discrimination**					**45351123**
Clostridium paraputrificum		72.33	0.41	SUC 92		
Clostridium barati		9.71	0.32	URE 1		
Clostridium septicum		8.91	0.29			
45351130	**Low discrimination**					**45351130**
					SPO	
Clostridium septicum		70.81	0.58	MLZ 1	99	
Clostridium paraputrificum		27.37	0.48		99	
Collinsella aerofaciens		0.63	0.15		1	
45351132	**Low discrimination**					**45351132**
					SPO	
Clostridium paraputrificum		63.35	0.41	SUC 92	99	
Bifidobacterium breve		8.61	0.44	MLZ 1	1	
Clostridium barati		8.51	0.32		99	
45351142	**Low discrimination**					**45351142**
					SPO	
Clostridium barati		46.96	0.40	SUC 99	99	
Lactobacillus acidophilus		18.84	0.35	SOR 8	1	
Bifidobacterium breve		9.32	0.40		1	
45351152	**Low discrimination**					**45351152**
					GRA SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		89.30	0.40	SUC 99	99	1	
Clostridium barati		4.55	0.05	RAF 95	99	99	
Leptotrichia buccalis		4.33	-.03		1	1	
45351160	**Low discrimination**						**45351160**
Clostridium paraputrificum		72.69	0.63	SUC 92			
Clostridium septicum		25.24	0.58	SOR 7			
Clostridium barati		1.61	0.40				
45351162	**Low discrimination**						**45351162**
					SPO		
Clostridium paraputrificum		81.52	0.56	SUC 92	99		
Clostridium barati		12.65	0.48	SOR 7	99		
Lactobacillus acidophilus		1.69	0.35		1		
45351172	**Low discrimination**						**45351172**
					SPO		
Bifidobacterium breve		60.22	0.44	SUC 99	1		
Clostridium paraputrificum		33.36	0.22	RAF 95	99		
Clostridium barati		5.18	0.14		99		
45351200	**Low discrimination**						**45351200**
					SPO		
Lactobacillus jensenii		72.09	0.41	SUC 90	1		
Lactobacillus catenaformis		13.05	0.29	RHA 1	1		
Clostridium barati		12.60	0.23		99		
45351202	**Low discrimination**						**45351202**
					SPO		
Lactobacillus jensenii		45.77	0.38	SUC 90	1		
Clostridium barati		36.01	0.31	RHA 1	99		
Lactobacillus catenaformis		12.43	0.29		1		
45351222	**Low discrimination**						**45351222**
					GRA	SPO	
Clostridium barati		55.57	0.40	SUC 99	99	99	
Prevotella oralis		34.13	0.37	RHA 8	1	1	
Lactobacillus jensenii		5.89	0.28		99	1	
45351302	**Low discrimination**						**45351302**
					GRA	SPO	
Clostridium barati		47.69	0.40	SUC 99	99	99	
Prevotella oralis		24.15	0.36	RHA 8	1	1	
Lactobacillus acidophilus		11.65	0.31		99	1	
45351320	**Low discrimination**						**45351320**
Clostridium septicum		66.85	0.58	RHA 1			
Clostridium paraputrificum		25.83	0.48				
Clostridium barati		4.27	0.40				
45351322	**Low discrimination**						**45351322**
					GRA	SPO	
Prevotella oralis		60.26	0.57	SUC 92	1	1	
Clostridium barati		18.79	0.48	RAF 92	99	99	
Clostridium paraputrificum		16.25	0.41		99	99	
45351400	**Low discrimination**						**45351400**
					SPO		
Lactobacillus jensenii		80.66	0.41	SUC 90	1		
Lactobacillus catenaformis		14.61	0.29	ARA 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		1.64	0.07		99	
45351402	**Low discrimination**					**45351402**
					SPO	
Lactobacillus jensenii		72.45	0.38	SUC 90	1	
Lactobacillus catenaformis		19.68	0.29	ARA 1	1	
Clostridium barati		6.62	0.15		99	
45351520	**Low discrimination**					**45351520**
					SPO	
Clostridium septicum		71.11	0.58	ARA 1	99	
Clostridium paraputrificum		27.48	0.48		99	
Collinsella aerofaciens		0.63	0.15		1	
45351522	**Low discrimination**					**45351522**
Clostridium paraputrificum		72.31	0.41	SUC 92		
Clostridium barati		9.71	0.32	ARA 1		
Clostridium septicum		8.91	0.29			
45353000	**Low discrimination**					**45353000**
					SPO	
Clostridium innocuum		73.84	0.49	MLT 1	99	
Lactobacillus jensenii		18.72	0.41		1	
Lactobacillus catenaformis		3.39	0.29		1	
45353002	**Low discrimination**					**45353002**
					SPO	
Clostridium innocuum		73.58	0.49	MLT 1	99	
Lactobacillus jensenii		12.43	0.38		1	
Clostridium barati		9.78	0.31		99	
45353022	**Low discrimination**					**45353022**
					SPO	
Clostridium barati		80.82	0.40	SUC 99	99	
Lactobacillus jensenii		8.56	0.28	MAN 8	1	
Lactobacillus catenaformis		2.33	0.18		1	
45353102	**Low discrimination**					**45353102**
					SPO	
Clostridium barati		46.49	0.40	SUC 99	99	
Clostridium tertium		15.80	0.35	MAN 8	99	
Bifidobacterium breve		9.22	0.40		1	
45353112	**Low discrimination**					**45353112**
					SPO	
Clostridium tertium		71.89	0.39	SUC 99	99	
Bifidobacterium breve		23.60	0.40	TRE 90	1	
Bifidobacterium adolescentis		1.44	0.18		1	
45353120	**Low discrimination**					**45353120**
Clostridium septicum		66.93	0.58	MAN 1		
Clostridium paraputrificum		25.87	0.48			
Clostridium barati		4.27	0.40			
45353122	**Low discrimination**					**45353122**
Clostridium barati		36.57	0.48	SUC 99		
Clostridium paraputrificum		31.64	0.41	MAN 8		
Clostridium tertium		12.43	0.43			
45353130	**Low discrimination**					**45353130**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		69.56	0.39	SUC 99			
Clostridium septicum		18.22	0.24	TRE 90			
Clostridium paraputrificum		7.04	0.14				
45353132	**Low discrimination**						**45353132**
					SPO		
Clostridium tertium		81.30	0.47	SUC 99	99		
Bifidobacterium breve		15.82	0.44	TRE 90	1		
Clostridium barati		1.36	0.14		99		
45353142	**Low discrimination**						**45353142**
					SPO		
Bifidobacterium breve		60.91	0.40	SUC 99	1		
Clostridium barati		26.70	0.21	RAF 95	99		
Lactobacillus acidophilus		6.49	0.13		1		
45353152	**Good identification**						**45353152**
					SPO		
Bifidobacterium breve		97.02	0.40	SUC 99	1		
Clostridium tertium		2.99	0.05	RAF 95	99		
45353162	**Low discrimination**						**45353162**
					SPO		
Bifidobacterium breve		41.58	0.44	SUC 99	1		
Clostridium barati		30.76	0.30	RAF 95	99		
Clostridium paraputrificum		23.03	0.22		99		
45353172	**Good identification**						**45353172**
					SPO		
Bifidobacterium breve		95.06	0.44	SUC 99	1		
Clostridium tertium		4.94	0.13	RAF 95	99		
45353322	**Low discrimination**						**45353322**
					SPO	GRA	
Clostridium ramosum		59.32	0.37	SUC 99	92	50	
Clostridium barati		25.62	0.30	TRE 94	99	99	
Prevotella oralis		9.54	0.23		1	1	
45353400	**Good identification**						**45353400**
					SPO		
Clostridium innocuum		97.37	0.36	MLT 1	99		
Lactobacillus jensenii		1.41	0.07		1		
Actinomyces israelii		1.22	0.18		1		
45353402	**Good identification**						**45353402**
					SPO		
Clostridium innocuum		96.52	0.36	MLT 1	99		
Actinomyces israelii		1.81	0.21		1		
Lactobacillus jensenii		0.93	0.04		1		
45354100	**Low discrimination**						**45354100**
					COC	GRA	
Fusobacterium mortiferum		79.57	0.36	MLT 11	1	1	
Collinsella aerofaciens		12.07	-.03	XYL 7	1	99	
Atopobium parvulum		8.37	0.08		99	99	
45355000	**Low discrimination**						**45355000**
					GRA		
Lactobacillus jensenii		74.55	0.41	SUC 90	99		
Lactobacillus catenaformis		13.50	0.29	XYL 1	99		
Fusobacterium mortiferum		6.31	0.33		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45355002	**Low discrimination**						**45355002**
					SPO		
Lactobacillus jensenii		68.76	0.38	SUC 90	1		
Lactobacillus catenaformis		18.68	0.29	XYL 1	1		
Clostridium barati		6.28	0.15		99		
45355120	**Low discrimination**						**45355120**
					COC	SPO	
Clostridium septicum		70.17	0.58	XYL 1	1	99	
Clostridium paraputrificum		27.12	0.48		1	99	
Atopobium parvulum		0.69	0.29		99	1	
45355122	**Low discrimination**						**45355122**
					GRA	SPO	
Prevotella oralis		44.34	0.48	SUC 92	1	1	
Clostridium paraputrificum		41.47	0.41	RAF 92	99	99	
Clostridium barati		5.57	0.32	XYL 11	99	99	
45355322	**Low discrimination**						**45355322**
					GRA	SPO	
Prevotella oralis		83.86	0.42	SUC 92	1	1	
Parabacteroides distasonis		11.30	0.40	RAF 92	1	1	
Clostridium barati		2.14	0.14	XYL 11	99	99	
45355332	**Low discrimination**						**45355332**
Parabacteroides distasonis		82.38	0.35	RAF 99			
Prevotella oralis		11.46	0.08				
Clostridium clostridioforme		6.16	0.10				
45355502	**Low discrimination**						**45355502**
					SPO		
Bifidobacterium adolescentis		79.13	0.46	SUC 99	1		
Clostridium butyricum		10.58	0.17	RAF 92	99		
Actinomyces israelii		9.85	0.37		1		
45355512	**Good identification**						**45355512**
Bifidobacterium adolescentis		95.99	0.48	SUC 99			
Clostridium clostridioforme		2.00	0.16	RAF 92			
Actinomyces israelii		1.46	0.23				
45355722	**Low discrimination**						**45355722**
Parabacteroides distasonis		51.68	0.36	RAF 99			
Bacteroides uniformis		22.55	0.11				
Clostridium clostridioforme		11.67	0.19				
45357100	**Low discrimination**						**45357100**
					SPO		
Actinomyces israelii		63.16	0.48	SUC 98	1		
Clostridium tertium		31.79	0.29		99		
Collinsella aerofaciens		2.58	-.03		1		
45357102	**Low discrimination**						**45357102**
					SPO		
Clostridium tertium		46.39	0.37	SUC 99	99		
Actinomyces israelii		46.08	0.51	TRE 90	1		
Bifidobacterium adolescentis		6.54	0.31		1		
45357112	**Low discrimination**						**45357112**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		84.72	0.41	SUC 99	99	
Bifidobacterium adolescentis		8.21	0.32	TRE 90	1	
Actinomyces israelii		7.07	0.37		1	
45357120	**Low discrimination**					**45357120**
Clostridium tertium		65.30	0.37	SUC 99		
Clostridium septicum		22.02	0.24	TRE 90		
Clostridium paraputrificum		8.51	0.14			
45357122	**Acceptable identification**					**45357122**
					SPO	
Clostridium tertium		91.80	0.45	SUC 99	99	
Clostridium barati		1.98	0.14	TRE 90	99	
Actinomyces israelii		1.94	0.31		1	
45357130	**Good identification**					**45357130**
					SPO	
Clostridium tertium		99.65	0.41	SUC 99	99	
Actinomyces israelii		0.35	0.14	TRE 90	1	
45357132	**Good identification**					**45357132**
					SPO	
Clostridium tertium		99.68	0.50	SUC 99	99	
Actinomyces israelii		0.18	0.17	TRE 90	1	
Bifidobacterium breve		0.14	0.10		1	
45357140	**Good identification**					**45357140**
Actinomyces israelii		100.0	0.41	SUC 98		
45357142	**Acceptable identification**					**45357142**
					SPO	
Actinomyces israelii		93.35	0.44	SUC 98	1	
Bifidobacterium adolescentis		4.21	0.15		1	
Clostridium tertium		2.44	0.03		99	
45357400	**Low discrimination**					**45357400**
					SPO	
Actinomyces israelii		73.19	0.37	SUC 98	1	
Clostridium innocuum		26.81	0.14		99	
45357402	**Low discrimination**					**45357402**
					SPO	
Actinomyces israelii		77.45	0.40	SUC 98	1	
Clostridium innocuum		18.92	0.14		99	
Bifidobacterium adolescentis		3.63	0.11		1	
45357500	**Acceptable identification**					**45357500**
					SPO	
Actinomyces israelii		91.00	0.47	SUC 98	1	
Clostridium butyricum		7.40	0.07		99	
Bifidobacterium adolescentis		1.60	0.11		1	
45357502	**Low discrimination**					**45357502**
					SPO	
Bifidobacterium adolescentis		52.90	0.45	SUC 99	1	
Actinomyces israelii		45.58	0.50	RAF 92	1	
Clostridium butyricum		0.96	0.00		99	
45357512	**Low discrimination**					**45357512**
Bifidobacterium adolescentis		85.59	0.46	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		9.02	0.36	RAF 92			
Bifidobacterium dentium		4.06	0.06				
45357540	**Good identification**						**45357540**
Actinomyces israelii		100.0	0.40	SUC 98			
45357542	**Low discrimination**						**45357542**
Actinomyces israelii		73.05	0.43	SUC 98			
Bifidobacterium adolescentis		26.95	0.29				
45361000	**Good identification**						**45361000**
					COC	SPO	
Lactobacillus jensenii		99.50	0.50	SUC 90	1	1	
Clostridium septicum		0.25	-.03	FRU 99	1	99	
Atopobium parvulum		0.25	0.04		99	1	
45361002	**Good identification**						**45361002**
					SPO		
Lactobacillus jensenii		98.99	0.47	SUC 90	1		
Lactobacillus acidophilus		0.74	0.07	FRU 99	1		
Clostridium barati		0.26	-.03		99		
45361010	**Good identification**						**45361010**
Lactobacillus jensenii		100.0	0.41	SUC 90			
45361012	**Good identification**						**45361012**
					GRA		
Lactobacillus jensenii		97.78	0.38	SUC 90	99		
Leptotrichia buccalis		2.22	-.03	FRU 99	1		
45361020	**Low discrimination**						**45361020**
					SPO		
Lactobacillus jensenii		49.09	0.39	SUC 90	1		
Clostridium septicum		48.89	0.32	FRU 99	99		
Clostridium paraputrificum		1.67	0.04		99		
45361022	**Low discrimination**						**45361022**
					SPO		
Lactobacillus jensenii		89.75	0.36	SUC 90	1		
Clostridium barati		2.85	0.05	FRU 99	99		
Clostridium septicum		2.74	0.03		99		
45361120	**Very good identification**						**45361120**
					COC	SPO	
Clostridium septicum		96.48	0.66	FRU 99	1	99	
Clostridium paraputrificum		3.30	0.38		1	99	
Atopobium parvulum		0.17	0.25		99	1	
45361122	**Low discrimination**						**45361122**
					SPO		
Clostridium septicum		52.30	0.37	FRU 99	99		
Clostridium paraputrificum		37.61	0.32	BGL 2	99		
Lactobacillus acidophilus		5.99	0.25		1		
45365100	**Good identification**						**45365100**
					COC	SPO	
Atopobium parvulum		99.08	0.42	SUC 96	99	1	
Clostridium septicum		0.55	-.03	FRU 99	1	99	
Clostridium butyricum		0.37	-.07		1	99	
45370000	**Acceptable identification**						**45370000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Lactobacillus jensenii		92.36	0.50	SUC 90	99	1	
Fusobacterium mortiferum		6.89	0.40	GAL 99	1	1	
Clostridium barati		0.28	0.01		99	99	
45370002	**Good identification**						**45370002**
					GRA	SPO	
Lactobacillus jensenii		96.74	0.47	SUC 90	99	1	
Clostridium barati		1.34	0.10	GAL 99	99	99	
Fusobacterium mortiferum		1.20	0.24		1	1	
45370010	**Good identification**						**45370010**
					SPO		
Lactobacillus jensenii		99.27	0.41	SUC 90	1		
Clostridium difficile		0.73	-.01	GAL 99	99		
45370012	**Good identification**						**45370012**
					GRA		
Lactobacillus jensenii		97.78	0.38	SUC 90	99		
Leptotrichia buccalis		2.22	-.03	GAL 99	1		
45370020	**Low discrimination**						**45370020**
					SPO		
Lactobacillus jensenii		48.38	0.39	SUC 90	1		
Clostridium septicum		48.19	0.32	GAL 99	99		
Clostridium barati		1.78	0.10		99		
45370022	**Low discrimination**						**45370022**
					SPO		
Lactobacillus jensenii		80.12	0.36	SUC 90	1		
Clostridium barati		13.27	0.18	GAL 99	99		
Clostridium septicum		2.44	0.03		99		
45370120	**Very good identification**						**45370120**
					COC	SPO	
Clostridium septicum		96.39	0.66	GAL 99	1	99	
Clostridium paraputrificum		3.30	0.38		1	99	
Atopobium parvulum		0.17	0.25		99	1	
45370122	**Low discrimination**						**45370122**
Clostridium septicum		48.90	0.37	GAL 99			
Clostridium paraputrificum		35.16	0.32	BGL 2			
Clostridium barati		8.05	0.26				
45371000	**Very good identification**						**45371000**
					COC	SPO	
Lactobacillus jensenii		99.25	0.84	SUC 90	1	1	
Clostridium septicum		0.25	0.32		1	99	
Atopobium parvulum		0.25	0.38		99	1	
45371001	**Good identification**						**45371001**
					COC	SPO	
Lactobacillus jensenii		99.50	0.50	SUC 90	1	1	
Clostridium septicum		0.25	-.03	URE 1	1	99	
Atopobium parvulum		0.25	0.04		99	1	
45371002	**Very good identification**						**45371002**
					SPO		
Lactobacillus jensenii		98.79	0.81	SUC 90	1		
Lactobacillus acidophilus		0.74	0.41		1		
Clostridium barati		0.26	0.31		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45371003	**Good identification**						**45371003**
					SPO		
Lactobacillus jensenii		98.99	0.47	SUC 90	1		
Lactobacillus acidophilus		0.74	0.07	URE 1	1		
Clostridium barati		0.26	-.03		99		
45371010	**Very good identification**						**45371010**
					COC	GRA	
Lactobacillus jensenii		99.97	0.76	SUC 90	1	99	
Leptotrichia buccalis		0.02	-.03		1	1	
Atopobium parvulum		0.01	0.04		99	99	
45371011	**Good identification**						**45371011**
Lactobacillus jensenii		100.0	0.41	SUC 90			
45371012	**Very good identification**						**45371012**
					GRA		
Lactobacillus jensenii		97.75	0.73	SUC 90	99		
Leptotrichia buccalis		2.22	0.32		1		
Lactobacillus acidophilus		0.02	0.07		99		
45371013	**Good identification**						**45371013**
					GRA		
Lactobacillus jensenii		97.78	0.38	SUC 90	99		
Leptotrichia buccalis		2.22	-.03	URE 1	1		
45371020	**Low discrimination**						**45371020**
					SPO		
Lactobacillus jensenii		48.99	0.73	SUC 90	1		
Clostridium septicum		48.79	0.66		99		
Clostridium paraputrificum		1.67	0.38		99		
45371021	**Low discrimination**						**45371021**
					SPO		
Lactobacillus jensenii		49.09	0.39	SUC 90	1		
Clostridium septicum		48.89	0.32	URE 1	99		
Clostridium paraputrificum		1.67	0.04		99		
45371022	**Low discrimination**						**45371022**
					SPO		
Lactobacillus jensenii		89.29	0.70	SUC 90	1		
Clostridium barati		2.84	0.40		99		
Clostridium septicum		2.72	0.37		99		
45371023	**Low discrimination**						**45371023**
					SPO		
Lactobacillus jensenii		89.75	0.36	SUC 90	1		
Clostridium barati		2.85	0.05	URE 1	99		
Clostridium septicum		2.74	0.03		99		
45371030	**Very good identification**						**45371030**
					SPO		
Lactobacillus jensenii		96.95	0.65	SUC 90	1		
Clostridium septicum		2.93	0.32		99		
Clostridium paraputrificum		0.10	0.04		99		
45371032	**Very good identification**						**45371032**
					GRA		
Lactobacillus jensenii		98.66	0.62	SUC 90	99		
Leptotrichia buccalis		1.00	0.15		1		
Lactobacillus acidophilus		0.09	0.07		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45371040	**Good identification**						**45371040**
					COC	SPO	
Lactobacillus jensenii		99.00	0.50	SUC 90	1	1	
Clostridium barati		0.50	0.05	SOR 1	1	99	
Atopobium parvulum		0.25	0.04		99	1	
45371042	**Low discrimination**						**45371042**
					SPO		
Lactobacillus jensenii		89.71	0.47	SUC 90	1		
Lactobacillus acidophilus		8.24	0.26	SOR 1	1		
Clostridium barati		2.05	0.13		99		
45371050	**Good identification**						**45371050**
Lactobacillus jensenii		100.0	0.41	SUC 90			
45371052	**Good identification**						**45371052**
					GRA		
Lactobacillus jensenii		97.78	0.38	SUC 90	99		
Leptotrichia buccalis		2.22	-.03	SOR 1	1		
45371060	**Low discrimination**						**45371060**
					SPO		
Lactobacillus jensenii		42.78	0.39	SUC 90	1		
Clostridium septicum		42.61	0.32	SOR 1	99		
Clostridium paraputrificum		10.87	0.19		99		
45371062	**Low discrimination**						**45371062**
					SPO		
Lactobacillus jensenii		53.95	0.36	SUC 90	1		
Lactobacillus acidophilus		19.83	0.26	SOR 1	1		
Clostridium barati		14.76	0.21		99		
45371100	**Low discrimination**						**45371100**
					COC	SPO	
Clostridium septicum		47.27	0.66	NAG 99	1	99	
Atopobium parvulum		46.72	0.72		99	1	
Lactobacillus jensenii		1.92	0.50		1	1	
45371102	**Low discrimination**						**45371102**
					COC		
Lactobacillus acidophilus		44.17	0.60	SUC 99	1		
Atopobium parvulum		33.31	0.59		99		
Lactobacillus jensenii		5.17	0.47		1		
45371112	**Very good identification**						**45371112**
					GRA		
Leptotrichia buccalis		99.11	0.66	SUC 99	1		
Lactobacillus jensenii		0.45	0.38		99		
Bifidobacterium breve		0.14	0.37		99		
45371120	**Very good identification**						**45371120**
					COC	SPO	
Clostridium septicum		96.47	1.00		1	99	
Clostridium paraputrificum		3.30	0.72		1	99	
Atopobium parvulum		0.17	0.59		99	1	
45371121	**Very good identification**						**45371121**
					COC	SPO	
Clostridium septicum		96.47	0.66	URE 1	1	99	
Clostridium paraputrificum		3.30	0.38		1	99	
Atopobium parvulum		0.17	0.25		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45371122	**Low discrimination**						**45371122**
					SPO		
Clostridium septicum		52.08	0.71	BGL 2	99		
Clostridium paraputrificum		37.46	0.66		99		
Lactobacillus acidophilus		5.96	0.60		1		
45371123	**Low discrimination**						**45371123**
					SPO		
Clostridium septicum		52.13	0.37	URE 1	99		
Clostridium paraputrificum		37.49	0.32	BGL 2	99		
Lactobacillus acidophilus		5.97	0.25		1		
45371130	**Very good identification**						**45371130**
					SPO		
Clostridium septicum		95.81	0.66	MLZ 1	99		
Clostridium paraputrificum		3.28	0.38		99		
Lactobacillus jensenii		0.32	0.31		1		
45371132	**Low discrimination**						**45371132**
					GRA	SPO	
Leptotrichia buccalis		81.66	0.50	SUC 99	1	1	
Clostridium septicum		7.48	0.37	NAG 10	99	99	
Clostridium paraputrificum		5.38	0.32		99	99	
45371142	**Low discrimination**						**45371142**
					SPO		
Lactobacillus acidophilus		77.21	0.44	SUC 99	1		
Bifidobacterium breve		7.60	0.37	SOR 11	1		
Clostridium barati		5.00	0.21		99		
45371160	**Low discrimination**						**45371160**
Clostridium septicum		79.26	0.66	SOR 1			
Clostridium paraputrificum		20.23	0.53				
Clostridium perfringens		0.16	0.26				
45371162	**Low discrimination**						**45371162**
					SPO		
Clostridium paraputrificum		64.04	0.47	SUC 92	99		
Lactobacillus acidophilus		16.73	0.44	SOR 7	1		
Clostridium septicum		11.95	0.37		99		
45371172	**Low discrimination**						**45371172**
					GRA	SPO	
Bifidobacterium breve		56.51	0.41	SUC 99	99	1	
Leptotrichia buccalis		25.42	0.15	RAF 95	1	1	
Clostridium paraputrificum		12.48	0.12		99	99	
45371200	**Good identification**						**45371200**
					COC	SPO	
Lactobacillus jensenii		99.00	0.50	SUC 90	1	1	
Clostridium barati		0.50	0.05	RHA 1	1	99	
Atopobium parvulum		0.25	0.04		99	1	
45371202	**Acceptable identification**						**45371202**
					SPO		
Lactobacillus jensenii		92.70	0.47	SUC 90	1		
Lactobacillus acidophilus		5.19	0.22	RHA 1	1		
Clostridium barati		2.11	0.13		99		
45371210	**Good identification**						**45371210**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		100.0	0.41	SUC 90			
45371212	**Good identification**						**45371212**
					GRA		
Lactobacillus jensenii		97.78	0.38	SUC 90	99		
Leptotrichia buccalis		2.22	-.03	RHA 1	1		
45371220	**Low discrimination**						**45371220**
					SPO		
Lactobacillus jensenii		47.63	0.39	SUC 90	1		
Clostridium septicum		47.43	0.32	RHA 1	99		
Clostridium barati		2.90	0.13		99		
45371222	**Low discrimination**						**45371222**
					SPO		
Lactobacillus jensenii		57.93	0.36	SUC 90	1		
Clostridium barati		15.85	0.21	RHA 1	99		
Lactobacillus acidophilus		12.97	0.22		1		
45371302	**Low discrimination**						**45371302**
					COC	SPO	
Lactobacillus acidophilus		76.02	0.40	SUC 99	1	1	
Clostridium barati		8.08	0.21	RHA 7	1	99	
Atopobium parvulum		7.69	0.25		99	1	
45371320	**Very good identification**						**45371320**
Clostridium septicum		95.44	0.66	RHA 1			
Clostridium paraputrificum		3.27	0.38				
Clostridium ramosum		0.85	0.35				
45371322	**Low discrimination**						**45371322**
					SPO		
Clostridium ramosum		32.68	0.45	SUC 99	92		
Clostridium septicum		18.77	0.37		99		
Lactobacillus acidophilus		16.01	0.40		1		
45371400	**Good identification**						**45371400**
					COC	SPO	
Lactobacillus jensenii		99.12	0.50	SUC 90	1	1	
Clostridium septicum		0.25	-.03	ARA 1	1	99	
Atopobium parvulum		0.25	0.04		99	1	
45371402	**Good identification**						**45371402**
Lactobacillus jensenii		98.57	0.47	SUC 90			
Lactobacillus acidophilus		0.74	0.07	ARA 1			
Actinomyces israelii		0.43	0.18				
45371410	**Good identification**						**45371410**
Lactobacillus jensenii		100.0	0.41	SUC 90			
45371412	**Good identification**						**45371412**
					GRA		
Lactobacillus jensenii		97.11	0.38	SUC 90	99		
Leptotrichia buccalis		2.20	-.03	ARA 1	1		
Clostridium clostridioforme		0.68	0.04		50		
45371420	**Low discrimination**						**45371420**
					SPO		
Lactobacillus jensenii		49.09	0.39	SUC 90	1		
Clostridium septicum		48.89	0.32	ARA 1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		1.67	0.04		99		
45371422	**Low discrimination**						**45371422**
					SPO		
Lactobacillus jensenii		89.75	0.36	SUC 90	1		
Clostridium barati		2.85	0.05	ARA 1	99		
Clostridium septicum		2.74	0.03		99		
45371520	**Very good identification**						**45371520**
					COC	SPO	
Clostridium septicum		96.46	0.66	ARA 1	1	99	
Clostridium paraputrificum		3.30	0.38		1	99	
Atopobium parvulum		0.17	0.25		99	1	
45371522	**Low discrimination**						**45371522**
					SPO		
Clostridium septicum		51.75	0.37	ARA 1	99		
Clostridium paraputrificum		37.21	0.32	BGL 2	99		
Lactobacillus acidophilus		5.92	0.25		1		
45373000	**Low discrimination**						**45373000**
					SPO		
Lactobacillus jensenii		67.08	0.50	SUC 90	1		
Clostridium innocuum		30.99	0.41	MAN 1	99		
Actinomyces israelii		0.89	0.30		1		
45373002	**Low discrimination**						**45373002**
					SPO		
Lactobacillus jensenii		55.27	0.47	SUC 90	1		
Clostridium innocuum		38.31	0.41	MAN 1	99		
Lactobacillus acidophilus		2.16	0.20		1		
45373010	**Acceptable identification**						**45373010**
					SPO		
Lactobacillus jensenii		91.53	0.41	SUC 90	1		
Clostridium innocuum		5.29	0.18	MAN 1	99		
Clostridium tertium		2.64	0.13		99		
45373012	**Low discrimination**						**45373012**
					SPO		
Lactobacillus jensenii		79.85	0.38	SUC 90	1		
Clostridium tertium		10.35	0.21	MAN 1	99		
Clostridium innocuum		6.92	0.18		99		
45373020	**Low discrimination**						**45373020**
					SPO		
Lactobacillus jensenii		45.20	0.39	SUC 90	1		
Clostridium septicum		45.02	0.32	MAN 1	99		
Clostridium tertium		2.93	0.17		99		
45373022	**Low discrimination**						**45373022**
					SPO		
Lactobacillus jensenii		49.28	0.36	SUC 90	1		
Clostridium tertium		14.37	0.25	MAN 1	99		
Clostridium barati		13.48	0.21		99		
45373100	**Low discrimination**						**45373100**
					COC	SPO	
Clostridium tertium		46.33	0.43	SUC 99	1	99	
Clostridium septicum		21.57	0.32		1	99	
Atopobium parvulum		21.32	0.38		99	1	
45373102	**Low discrimination**						**45373102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium tertium		72.49	0.51	SUC 99	99		
Lactobacillus acidophilus		13.56	0.38		1		
Actinomyces israelii		3.62	0.43		1		
45373110	**Good identification**						**45373110**
					SPO		
Clostridium tertium		97.12	0.47	SUC 99	99		
Actinomyces israelii		0.81	0.26		1		
Bifidobacterium breve		0.70	0.20		1		
45373112	**Low discrimination**						**45373112**
					GRA	SPO	
Clostridium tertium		82.86	0.55	SUC 99	99	99	
Leptotrichia buccalis		14.50	0.32		1	1	
Bifidobacterium breve		2.02	0.37		99	1	
45373120	**Low discrimination**						**45373120**
Clostridium septicum		87.56	0.66	MAN 1			
Clostridium tertium		5.70	0.51				
Clostridium ramosum		3.33	0.45				
45373122	**Low discrimination**						**45373122**
Clostridium tertium		48.09	0.59	SUC 99			
Clostridium ramosum		37.41	0.56				
Clostridium septicum		5.03	0.37				
45373130	**Acceptable identification**						**45373130**
					SPO		
Clostridium tertium		91.08	0.55	SUC 99	99		
Clostridium septicum		7.95	0.32		99		
Bifidobacterium breve		0.39	0.24		1		
45373132	**Very good identification**						**45373132**
					GRA	SPO	
Clostridium tertium		97.40	0.63	SUC 99	99	99	
Bifidobacterium breve		1.40	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
45373142	**Low discrimination**						**45373142**
Bifidobacterium breve		43.98	0.37	SUC 99			
Lactobacillus acidophilus		23.51	0.22	RAF 95			
Actinomyces israelii		19.72	0.36				
45373152	**Low discrimination**						**45373152**
					GRA	SPO	
Bifidobacterium breve		64.30	0.37	SUC 99	99	1	
Clostridium tertium		26.71	0.21	RAF 95	99	99	
Leptotrichia buccalis		4.68	-.03		1	1	
45373162	**Low discrimination**						**45373162**
					SPO		
Bifidobacterium breve		42.08	0.41	SUC 99	1		
Clostridium tertium		16.59	0.25	RAF 95	99		
Clostridium ramosum		12.91	0.21		92		
45373172	**Low discrimination**						**45373172**
					SPO		
Bifidobacterium breve		58.79	0.41	SUC 99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium tertium		41.21	0.29	RAF 95	99			
45373320	**Low discrimination**							**45373320**
Clostridium ramosum		81.71	0.47	SUC 99				
Clostridium septicum		16.40	0.32					
Clostridium tertium		1.07	0.17					
45373322	**Very good identification**							**45373322**
					SPO			
Clostridium ramosum		98.12	0.58	SUC 99	92			
Clostridium tertium		0.96	0.25		99			
Lactobacillus acidophilus		0.45	0.18		1			
45373500	**Low discrimination**							**45373500**
					SPO			
Actinomyces israelii		73.60	0.39	SUC 98	1			
Clostridium tertium		9.11	0.09	XYL 92	99			
Clostridium butyricum		8.86	0.02		99			
45373502	**Low discrimination**							**45373502**
					SPO			
Actinomyces israelii		59.58	0.42	SUC 98	1			
Bifidobacterium adolescentis		21.06	0.28	XYL 92	1			
Clostridium tertium		14.75	0.17		99			
45374100	**Good identification**							**45374100**
					COC	GRA	SPO	
Atopobium parvulum		96.34	0.42	SUC 96	99	99	1	
Fusobacterium mortiferum		2.76	0.27	GAL 99	1	1	1	
Clostridium septicum		0.53	-.03		1	99	99	
45374712	**Good identification**							**45374712**
Clostridium clostridioforme		100.0	0.37	SUC 99				
45374732	**Good identification**							**45374732**
Clostridium clostridioforme		100.0	0.37	SUC 99				
45375000	**Low discrimination**							**45375000**
					COC			
Lactobacillus jensenii		66.58	0.50	SUC 90	1			
Atopobium parvulum		30.42	0.42	XYL 1	99			
Actinomyces israelii		1.80	0.35		1			
45375002	**Low discrimination**							**45375002**
					COC			
Lactobacillus jensenii		83.60	0.47	SUC 90	1			
Atopobium parvulum		10.11	0.29	XYL 1	99			
Actinomyces israelii		5.09	0.38		1			
45375010	**Good identification**							**45375010**
					COC			
Lactobacillus jensenii		97.47	0.41	SUC 90	1			
Atopobium parvulum		1.35	0.08	XYL 1	99			
Actinomyces israelii		1.18	0.21		1			
45375012	**Good identification**							**45375012**
					GRA			
Lactobacillus jensenii		95.24	0.38	SUC 90	99			
Actinomyces israelii		2.60	0.24	XYL 1	99			
Leptotrichia buccalis		2.16	-.03		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45375020	**Low discrimination**						**45375020**
					COC	SPO	
Lactobacillus jensenii		42.26	0.39	SUC 90	1	1	
Clostridium septicum		42.08	0.32	XYL 1	1	99	
Atopobium parvulum		13.63	0.29		99	1	
45375022	**Low discrimination**						**45375022**
					COC	GRA	
Lactobacillus jensenii		79.95	0.36	SUC 90	1	99	
Atopobium parvulum		6.83	0.16	XYL 1	99	99	
Prevotella oralis		2.84	0.07		1	1	
45375100	**Very good identification**						**45375100**
					COC	SPO	
Atopobium parvulum		98.73	0.76	SUC 96	99	1	
Clostridium septicum		0.54	0.32		1	99	
Clostridium butyricum		0.37	0.28		1	99	
45375101	**Good identification**						**45375101**
					COC	SPO	
Atopobium parvulum		99.08	0.42	SUC 96	99	1	
Clostridium septicum		0.55	-.03	URE 1	1	99	
Clostridium butyricum		0.37	-.07		1	99	
45375102	**Very good identification**						**45375102**
					COC	SPO	
Atopobium parvulum		95.58	0.64	SUC 96	99	1	
Actinomyces israelii		1.94	0.48		1	1	
Clostridium butyricum		0.79	0.21		1	99	
45375110	**Acceptable identification**						**45375110**
					COC	SPO	
Atopobium parvulum		90.26	0.42	SUC 96	99	1	
Actinomyces israelii		3.19	0.31	MLZ 1	1	1	
Clostridium tertium		2.62	0.15		1	99	
45375120	**Low discrimination**						**45375120**
					COC	SPO	
Clostridium septicum		73.50	0.66	XYL 1	1	99	
Atopobium parvulum		23.80	0.64		99	1	
Clostridium paraputrificum		2.52	0.38		1	99	
45375122	**Low discrimination**						**45375122**
					COC	SPO	
Atopobium parvulum		52.98	0.51	SUC 96	99	1	
Clostridium septicum		18.92	0.37		1	99	
Clostridium paraputrificum		13.60	0.32		1	99	
45375140	**Low discrimination**						**45375140**
					COC	SPO	
Atopobium parvulum		88.50	0.42	SUC 96	99	1	
Actinomyces israelii		8.15	0.38	SOR 1	1	1	
Clostridium butyricum		2.86	0.10		1	99	
45375300	**Good identification**						**45375300**
					COC	SPO	
Atopobium parvulum		96.64	0.42	SUC 96	99	1	
Actinomyces israelii		1.72	0.26	RHA 1	1	1	
Clostridium butyricum		1.11	0.02		1	99	
45375322	**Low discrimination**						**45375322**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		82.47	0.51	RAF 99	1	1		
Prevotella oralis		8.57	0.21		1	1		
Atopobium parvulum		3.06	0.16		99	99		
45375332	**Low discrimination**							**45375332**
Parabacteroides distasonis		89.26	0.46	RAF 99				
Clostridium clostridioforme		10.74	0.25					
45375402	**Low discrimination**							**45375402**
					SPO			
Actinomyces israelii		47.37	0.37	SUC 98	1			
Clostridium butyricum		35.34	0.13		99			
Lactobacillus jensenii		9.61	0.12		1			
45375500	**Low discrimination**							**45375500**
					COC	SPO		
Clostridium butyricum		73.54	0.44	SUC 99	1	99		
Atopobium parvulum		21.99	0.42	RAF 95	99	1		
Actinomyces israelii		4.26	0.44		1	1		
45375502	**Low discrimination**							**45375502**
					SPO			
Clostridium butyricum		58.93	0.37	SUC 99	99			
Bifidobacterium adolescentis		18.76	0.43	RAF 95	1			
Actinomyces israelii		13.17	0.47		1			
45375512	**Low discrimination**							**45375512**
					SPO			
Bifidobacterium adolescentis		65.35	0.45	SUC 99	1			
Clostridium clostridioforme		14.98	0.31	RAF 92	66			
Clostridium butyricum		8.84	0.15		99			
45375522	**Low discrimination**							**45375522**
					COC	GRA	SPO	
Parabacteroides distasonis		44.73	0.39	RAF 99	1	1	1	
Clostridium butyricum		29.55	0.19		1	99	99	
Atopobium parvulum		8.14	0.16		99	99	1	
45375612	**Good identification**							**45375612**
Clostridium clostridioforme		100.0	0.36	SUC 99				
45375632	**Good identification**							**45375632**
Clostridium clostridioforme		100.0	0.36	SUC 99				
45375712	**Good identification**							**45375712**
Clostridium clostridioforme		99.02	0.45	SUC 99				
Bifidobacterium adolescentis		0.60	0.11	RAF 94				
Actinomyces israelii		0.38	0.14					
45375722	**Low discrimination**							**45375722**
Parabacteroides distasonis		71.30	0.47	RAF 99				
Clostridium clostridioforme		25.91	0.34					
Bacteroides ovatus		2.31	0.06					
45375732	**Low discrimination**							**45375732**
Clostridium clostridioforme		74.72	0.45	SUC 99				
Parabacteroides distasonis		24.30	0.43	RAF 94				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		0.97	0.03				
45377000	**Low discrimination**						**45377000**
					SPO		
Actinomyces israelii		76.73	0.48	SUC 98	1		
Clostridium innocuum		12.17	0.19		99		
Lactobacillus jensenii		5.06	0.15		1		
45377002	**Low discrimination**						**45377002**
					SPO		
Actinomyces israelii		81.15	0.51	SUC 98	1		
Clostridium innocuum		8.58	0.19		99		
Clostridium tertium		7.90	0.19		99		
45377040	**Good identification**						**45377040**
Actinomyces israelii		100.0	0.41	SUC 98			
45377042	**Good identification**						**45377042**
Actinomyces israelii		100.0	0.44	SUC 98			
45377100	**Low discrimination**						**45377100**
					COC	SPO	
Clostridium tertium		40.35	0.45	SUC 99	1	99	
Actinomyces israelii		33.51	0.58		1	1	
Atopobium parvulum		24.97	0.42		99	1	
45377102	**Low discrimination**						**45377102**
					COC	SPO	
Clostridium tertium		68.18	0.53	SUC 99	1	99	
Actinomyces israelii		28.31	0.61		1	1	
Atopobium parvulum		2.48	0.29		99	1	
45377110	**Acceptable identification**						**45377110**
					COC	SPO	
Clostridium tertium		93.17	0.50	SUC 99	1	99	
Actinomyces israelii		6.50	0.44		1	1	
Atopobium parvulum		0.33	0.08		99	1	
45377112	**Very good identification**						**45377112**
					SPO		
Clostridium tertium		95.53	0.58	SUC 99	99		
Actinomyces israelii		3.33	0.47		1		
Bifidobacterium adolescentis		0.69	0.29		1		
45377120	**Low discrimination**						**45377120**
					COC	SPO	
Clostridium tertium		85.08	0.53	SUC 99	1	99	
Clostridium septicum		9.56	0.32		1	99	
Atopobium parvulum		3.10	0.29		99	1	
45377122	**Very good identification**						**45377122**
					SPO		
Clostridium tertium		98.19	0.62	SUC 99	99		
Actinomyces israelii		0.87	0.41		1		
Clostridium ramosum		0.56	0.21		92		
45377130	**Very good identification**						**45377130**
					SPO		
Clostridium tertium		99.73	0.58	SUC 99	99		
Actinomyces israelii		0.15	0.24		1		
Clostridium septicum		0.06	-.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45377132	**Very good identification**						**45377132**
					SPO		
Clostridium tertium		99.72	0.66	SUC 99	99		
Bifidobacterium dentium		0.20	0.10		1		
Actinomyces israelii		0.07	0.27		1		
45377140	**Acceptable identification**						**45377140**
					COC	SPO	
Actinomyces israelii		94.56	0.51	SUC 98	1	1	
Clostridium tertium		2.96	0.11		1	99	
Atopobium parvulum		1.83	0.08		99	1	
45377142	**Acceptable identification**						**45377142**
					SPO		
Actinomyces israelii		93.41	0.54	SUC 98	1		
Clostridium tertium		5.84	0.19		99		
Bifidobacterium adolescentis		0.75	0.12		1		
45377150	**Low discrimination**						**45377150**
					COC	SPO	
Actinomyces israelii		69.27	0.37	SUC 98	1	1	
Clostridium tertium		25.78	0.15	MLZ 13	1	99	
Blautia producta		4.95	-.03		99	1	
45377152	**Low discrimination**						**45377152**
					SPO		
Actinomyces israelii		55.26	0.40	SUC 98	1		
Clostridium tertium		41.13	0.24	MLZ 13	99		
Bifidobacterium adolescentis		3.61	0.14		1		
45377300	**Low discrimination**						**45377300**
					COC	SPO	
Actinomyces israelii		79.27	0.39	SUC 98	1	1	
Clostridium tertium		12.81	0.11	RHA 7	1	99	
Atopobium parvulum		7.93	0.08		99	1	
45377302	**Low discrimination**						**45377302**
					SPO		
Actinomyces israelii		75.57	0.42	SUC 98	1		
Clostridium tertium		24.43	0.19	RHA 7	99		
45377342	**Good identification**						**45377342**
Actinomyces israelii		100.0	0.35	SUC 98			
45377400	**Acceptable identification**						**45377400**
					SPO		
Actinomyces israelii		91.69	0.46	SUC 98	1		
Clostridium butyricum		5.17	0.04		99		
Clostridium innocuum		3.14	0.06		99		
45377402	**Good identification**						**45377402**
					SPO		
Actinomyces israelii		95.63	0.49	SUC 98	1		
Clostridium innocuum		2.18	0.06		99		
Clostridium butyricum		1.40	-.03		99		
45377412	**Low discrimination**						**45377412**
Actinomyces israelii		64.66	0.35	SUC 98			
Bifidobacterium dentium		30.95	0.06	MLZ 13			
Bifidobacterium adolescentis		4.40	0.09				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45377440	**Good identification**						**45377440**
Actinomyces israelii		100.0	0.39	SUC 98			
45377442	**Good identification**						**45377442**
Actinomyces israelii		100.0	0.42	SUC 98			
45377500	**Low discrimination**						**45377500**
					SPO		
Actinomyces israelii		73.23	0.57	SUC 98	1		
Clostridium butyricum		24.79	0.28		99		
Clostridium tertium		1.09	0.11		99		
45377502	**Low discrimination**						**45377502**
					SPO		
Actinomyces israelii		74.57	0.60	SUC 98	1		
Bifidobacterium adolescentis		15.34	0.42		1		
Clostridium butyricum		6.54	0.21		99		
45377512	**Low discrimination**						**45377512**
Bifidobacterium dentium		79.44	0.40	SUC 99			
Bifidobacterium adolescentis		11.28	0.43	RAF 95			
Actinomyces israelii		6.71	0.46				
45377520	**Low discrimination**						**45377520**
					SPO		
Actinomyces israelii		43.25	0.36	SUC 98	1		
Clostridium tertium		30.22	0.19	NAG 6	99		
Clostridium butyricum		19.94	0.10		99		
45377532	**Acceptable identification**						**45377532**
					SPO		
Bifidobacterium dentium		94.86	0.44	SUC 99	1		
Clostridium tertium		4.78	0.32	RAF 95	99		
Actinomyces israelii		0.29	0.25		1		
45377540	**Low discrimination**						**45377540**
					COC	SPO	
Actinomyces israelii		66.39	0.50	SUC 98	1	1	
Blautia producta		28.59	0.24		99	1	
Clostridium butyricum		5.02	0.10		1	99	
45377542	**Low discrimination**						**45377542**
					COC		
Actinomyces israelii		89.53	0.53	SUC 98	1		
Bifidobacterium adolescentis		5.85	0.26		1		
Blautia producta		2.86	0.07		99		
45377552	**Low discrimination**						**45377552**
					COC		
Actinomyces israelii		40.28	0.39	SUC 98	1		
Blautia producta		25.80	0.15	MLZ 13	99		
Bifidobacterium adolescentis		21.54	0.28		1		
45377700	**Low discrimination**						**45377700**
					SPO		
Actinomyces israelii		87.79	0.38	SUC 98	1		
Clostridium butyricum		12.21	0.02	RHA 7	99		
45377702	**Acceptable identification**						**45377702**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		94.02	0.41	SUC 98	1		
Clostridium butyricum		3.39	-.05	RHA 7	99		
Bifidobacterium adolescentis		2.60	0.08		1		
45400000	**Low discrimination**						**45400000**
					SPO		
Clostridium botulinum biovar A		39.92	0.47	MNS 7	99		
Clostridium bifermentans		27.26	0.43	RAF 1	99		
Atopobium minutum		22.82	0.50		1		
45400002	**Low discrimination**						**45400002**
					SPO		
Atopobium minutum		86.55	0.47	RAF 1	1		
Clostridium sporogenes		7.47	0.19		99		
Clostridium botulinum biovar A		2.29	0.13		99		
45400020	**Low discrimination**						**45400020**
					SPO		
Atopobium minutum		76.73	0.42	RAF 1	1		
Clostridium bifermentans		17.55	0.23		99		
Clostridium botulinum biovar A		4.07	0.13		99		
45400022	**Good identification**						**45400022**
					GRA		
Atopobium minutum		97.51	0.39	RAF 1	99		
Pseudoflavonifractor capillosus		2.49	-.03		1		
45400040	**Low discrimination**						**45400040**
					SPO		
Clostridium botulinum biovar A		88.86	0.38	MNS 7	99		
Clostridium bifermentans		8.46	0.20	RAF 1	99		
Atopobium minutum		1.72	0.16		1		
45400100	**Low discrimination**						**45400100**
					GRA	SPO	
Fusobacterium mortiferum		84.56	0.45	FRU 99	1	1	
Clostridium botulinum biovar A		6.20	0.13		99	99	
Clostridium bifermentans		4.23	0.09		99	99	
45401000	**Low discrimination**						**45401000**
					GRA	SPO	
Atopobium minutum		62.34	0.42	RAF 1	99	1	
Fusobacterium mortiferum		28.99	0.42		1	1	
Clostridium botulinum biovar A		3.31	0.13		99	99	
45401002	**Acceptable identification**						**45401002**
					GRA		
Atopobium minutum		92.81	0.39	RAF 1	99		
Fusobacterium mortiferum		7.19	0.25		1		
45410000	**Low discrimination**						**45410000**
					GRA		
Fusobacterium mortiferum		52.52	0.73		1		
Eubacterium limosum		35.49	0.52		99		
Atopobium minutum		4.74	0.50		99		
45410001	**Low discrimination**						**45410001**
					GRA		
Fusobacterium mortiferum		54.33	0.39	URE 1	1		
Eubacterium limosum		36.72	0.18		99		
Atopobium minutum		4.90	0.15		99		
45410002	**Low discrimination**						**45410002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		60.51	0.57	BGL 10	1		
Atopobium minutum		32.76	0.47		99		
Eubacterium limosum		3.72	0.18		99		
45410010	**Acceptable identification**						**45410010**
					GRA	SPO	
Clostridium difficile		94.82	0.50	RAF 1	99	99	
Fusobacterium mortiferum		2.63	0.39	MAN 90	1	1	
Eubacterium limosum		1.78	0.18		99	1	
45410020	**Low discrimination**						**45410020**
					GRA		
Atopobium minutum		56.78	0.41	RAF 1	99		
Fusobacterium mortiferum		19.07	0.39		1		
Eubacterium limosum		12.89	0.18		99		
45410022	**Acceptable identification**						**45410022**
					GRA	CAT	
Atopobium minutum		90.65	0.38	RAF 1	99	1	
Fusobacterium mortiferum		5.07	0.22		1	5	
Propionibacterium granulosum		4.28	0.18		99	90	
45410040	**Low discrimination**						**45410040**
					GRA	SPO	
Fusobacterium mortiferum		38.02	0.39	SOR 1	1	1	
Eubacterium limosum		25.70	0.18		99	1	
Clostridium difficile		14.77	0.16		99	99	
45410100	**Very good identification**						**45410100**
					GRA		
Fusobacterium mortiferum		99.64	0.79		1		
Eubacterium limosum		0.29	0.18		99		
Atopobium minutum		0.04	0.15		99		
45410101	**Good identification**						**45410101**
Fusobacterium mortiferum		100.0	0.45	URE 1			
45410102	**Very good identification**						**45410102**
					GRA		
Fusobacterium mortiferum		99.77	0.63	BGL 10	1		
Atopobium minutum		0.23	0.12		99		
45410110	**Low discrimination**						**45410110**
					GRA	SPO	
Fusobacterium mortiferum		86.51	0.45	MLZ 1	1	1	
Clostridium difficile		13.49	0.15		99	99	
45410120	**Good identification**						**45410120**
					GRA		
Fusobacterium mortiferum		97.57	0.45	NAG 1	1		
Atopobium minutum		1.26	0.07		99		
Prevotella melaninogenica		1.17	-.06		1		
45410140	**Good identification**						**45410140**
Fusobacterium mortiferum		100.0	0.45	SOR 1			
45410200	**Low discrimination**						**45410200**
					GRA		
Fusobacterium mortiferum		52.99	0.39	RHA 1	1		
Eubacterium limosum		35.82	0.18		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium minutum		4.78	0.15		99			
45410300	**Good identification**							**45410300**
Fusobacterium mortiferum		100.0	0.45	RHA 1				
45410400	**Low discrimination**							**45410400**
					GRA			
Fusobacterium mortiferum		53.45	0.39	ARA 1	1			
Eubacterium limosum		36.12	0.18		99			
Atopobium minutum		4.82	0.15		99			
45410500	**Good identification**							**45410500**
Fusobacterium mortiferum		100.0	0.45	ARA 1				
45411000	**Very good identification**							**45411000**
					GRA	CAT		
Fusobacterium mortiferum		96.07	0.76		1	5		
Atopobium minutum		1.93	0.41		99	1		
Propionibacterium granulosum		0.97	0.38		99	90		
45411001	**Good identification**							**45411001**
					GRA			
Fusobacterium mortiferum		98.03	0.42	URE 1	1			
Atopobium minutum		1.97	0.07		99			
45411002	**Low discrimination**							**45411002**
					GRA	CAT		
Fusobacterium mortiferum		86.03	0.60	BGL 10	1	5		
Atopobium minutum		10.35	0.38		99	1		
Propionibacterium granulosum		1.96	0.28		99	90		
45411010	**Low discrimination**							**45411010**
					GRA	SPO	CAT	
Fusobacterium mortiferum		71.48	0.42	MLZ 1	1	1	5	
Clostridium difficile		17.34	0.15		99	99	1	
Propionibacterium granulosum		9.75	0.24		99	1	90	
45411020	**Low discrimination**							**45411020**
					GRA	CAT		
Fusobacterium mortiferum		48.61	0.42	NAG 1	1	5		
Atopobium minutum		32.17	0.33		99	1		
Propionibacterium granulosum		12.15	0.28		99	90		
45411040	**Good identification**							**45411040**
					GRA			
Fusobacterium mortiferum		98.03	0.42	SOR 1	1			
Atopobium minutum		1.97	0.07		99			
45411100	**Very good identification**							**45411100**
					GRA			
Fusobacterium mortiferum		99.82	0.82		1			
Lactobacillus fermentum		0.15	0.35		99			
Atopobium minutum		0.01	0.07		99			
45411101	**Good identification**							**45411101**
Fusobacterium mortiferum		100.0	0.48	URE 1				
45411102	**Very good identification**							**45411102**
					GRA	CAT		
Fusobacterium mortiferum		97.62	0.66	BGL 10	1	5		
Lactobacillus fermentum		1.32	0.35		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		0.47	0.12		1	90	
45411110	**Good identification**						**45411110**
Fusobacterium mortiferum		100.0	0.48	MLZ 1			
45411120	**Low discrimination**						**45411120**
					GRA	SPO	
Fusobacterium mortiferum		43.80	0.48	NAG 1	1	1	
Prevotella melaninogenica		34.66	0.29		1	1	
Clostridium septicum		8.46	0.18		99	99	
45411132	**Low discrimination**						**45411132**
					CAT		
Parabacteroides distasonis		85.58	0.39	XYL 90	76		
Capnocytophaga ochracea		12.08	0.12		90		
Prevotella oralis		2.34	0.00		1		
45411140	**Good identification**						**45411140**
Fusobacterium mortiferum		100.0	0.48	SOR 1			
45411200	**Good identification**						**45411200**
					GRA	CAT	
Fusobacterium mortiferum		95.15	0.42	RHA 1	1	5	
Propionibacterium granulosum		2.94	0.13		99	90	
Atopobium minutum		1.91	0.07		99	1	
45411300	**Good identification**						**45411300**
Fusobacterium mortiferum		100.0	0.48	RHA 1			
45411322	**Low discrimination**						**45411322**
Parabacteroides distasonis		82.78	0.52	XYL 90			
Prevotella oralis		17.22	0.28				
45411332	**Good identification**						**45411332**
Parabacteroides distasonis		100.0	0.48	XYL 90			
45411400	**Low discrimination**						**45411400**
					GRA		
Fusobacterium mortiferum		81.13	0.42	ARA 1	1		
Lactobacillus fermentum		17.25	0.32		99		
Atopobium minutum		1.63	0.07		99		
45411500	**Low discrimination**						**45411500**
					GRA		
Fusobacterium mortiferum		84.88	0.48	ARA 1	1		
Lactobacillus fermentum		12.62	0.35		99		
Odoribacter splanchnicus		1.54	-.03		1		
45411502	**Low discrimination**						**45411502**
					GRA		
Lactobacillus fermentum		52.99	0.35	ESL 1	99		
Fusobacterium mortiferum		39.61	0.32	SUC 87	1		
Bifidobacterium longum subsp. longum		4.08	0.09		99		
45411522	**Low discrimination**						**45411522**
					GRA	CAT	
Parabacteroides distasonis		67.24	0.40	XYL 90	1	76	
Capnocytophaga ochracea		13.24	0.16		1	90	
Lactobacillus fermentum		7.82	0.27		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45411532	**Low discrimination**						**45411532**
					GRA	CAT	
Parabacteroides distasonis		72.45	0.36	XYL 90	1	76	
Bifidobacterium longum subsp. longum		24.60	0.24		99	1	
Capnocytophaga ochracea		2.94	-.01		1	90	
45411722	**Good identification**						**45411722**
Parabacteroides distasonis		100.0	0.48	XYL 90			
45411732	**Good identification**						**45411732**
Parabacteroides distasonis		100.0	0.44	XYL 90			
45412000	**Low discrimination**						**45412000**
					SPO	CAT	
Eubacterium limosum		85.68	0.66	RAF 1	1	1	
Clostridium difficile		13.85	0.54		99	1	
Propionibacterium granulosum		0.24	0.36		1	90	
45412010	**Very good identification**						**45412010**
					SPO	CAT	
Clostridium difficile		98.69	0.66	RAF 1	99	1	
Eubacterium limosum		1.26	0.32		1	1	
Propionibacterium granulosum		0.05	0.22		1	90	
45412050	**Good identification**						**45412050**
					SPO		
Clostridium difficile		99.76	0.44	RAF 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		
45413100	**Good identification**						**45413100**
					GRA		
Fusobacterium mortiferum		98.82	0.48	MAN 1	1		
Eubacterium limosum		1.18	-.03		99		
45414000	**Low discrimination**						**45414000**
					GRA	SPO	
Fusobacterium mortiferum		84.25	0.54	XYL 7	1	1	
Eubacterium limosum		7.64	0.18		99	1	
Clostridium difficile		4.39	0.16		99	99	
45414002	**Acceptable identification**						**45414002**
					GRA		
Fusobacterium mortiferum		93.23	0.37	XYL 7	1		
Atopobium minutum		6.77	0.12	BGL 10	99		
45414100	**Very good identification**						**45414100**
Fusobacterium mortiferum		100.0	0.60	XYL 7			
45414102	**Good identification**						**45414102**
Fusobacterium mortiferum		100.0	0.44	XYL 7			
45415000	**Very good identification**						**45415000**
					GRA		
Fusobacterium mortiferum		98.09	0.57	XYL 7	1		
Lactobacillus fermentum		1.64	0.28		99		
Atopobium minutum		0.26	0.07		99		
45415002	**Low discrimination**						**45415002**
					GRA		
Fusobacterium mortiferum		86.90	0.40	XYL 7	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		13.10	0.28	BGL 10	99	
45415100	**Very good identification**					**45415100**
					GRA	
Fusobacterium mortiferum		98.84	0.63	XYL 7	1	
Lactobacillus fermentum		1.16	0.31		99	
45415102	**Low discrimination**					**45415102**
					GRA	
Fusobacterium mortiferum		81.44	0.47	XYL 7	1	
Lactobacillus fermentum		8.59	0.31	BGL 10	99	
Parabacteroides distasonis		6.22	0.26		1	
45415122	**Low discrimination**					**45415122**
					CAT	
Parabacteroides distasonis		69.37	0.60		76	
Bacteroides fragilis		27.41	0.32		99	
Prevotella oralis		1.39	0.19		1	
45415132	**Good identification**					**45415132**
Parabacteroides distasonis		98.88	0.56			
Bacteroides fragilis		0.73	-.03			
Capnocytophaga ochracea		0.39	0.02			
45415322	**Very good identification**					**45415322**
					CAT	
Parabacteroides distasonis		99.58	0.68		76	
Prevotella oralis		0.28	0.12		1	
Bacteroides fragilis		0.13	-.03		99	
45415332	**Very good identification**					**45415332**
Parabacteroides distasonis		100.0	0.64			
45415522	**Good identification**					**45415522**
					GRA	
Parabacteroides distasonis		94.58	0.57		1	
Bifidobacterium longum subsp. longum		3.39	0.28		99	
Lactobacillus fermentum		0.72	0.23		99	
45415532	**Low discrimination**					**45415532**
					GRA	
Parabacteroides distasonis		89.83	0.52		1	
Bifidobacterium longum subsp. longum		10.17	0.32		99	
45415720	**Good identification**					**45415720**
Bacteroides vulgatus		98.26	0.42	SUC 96		
Parabacteroides distasonis		1.74	0.31	MLT 99		
45415722	**Very good identification**					**45415722**
Parabacteroides distasonis		99.43	0.65			
Bacteroides vulgatus		0.57	0.08			
45415732	**Very good identification**					**45415732**
Parabacteroides distasonis		100.0	0.60			
45416010	**Good identification**					**45416010**
					SPO	
Clostridium difficile		99.76	0.44	RAF 1	99	
Eubacterium limosum		0.25	-.03	XYL 5	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45420000	**Low discrimination**						**45420000**
					GRA	SPO	
Clostridium botulinum biovar A		79.14	0.36	MNS 7	99	99	
Clostridium sporogenes		10.41	0.19	RAF 1	99	99	
Fusobacterium mortiferum		5.59	0.30		1	1	
45420100	**Acceptable identification**						**45420100**
					GRA	SPO	
Fusobacterium mortiferum		94.22	0.36	FRU 99	1	1	
Clostridium botulinum biovar A		5.78	0.02		99	99	
45421100	**Good identification**						**45421100**
					COC	GRA	
Fusobacterium mortiferum		96.28	0.39	FRU 99	1	1	
Blautia hansenii		3.73	-.03		99	99	
45425122	**Good identification**						**45425122**
Parabacteroides distasonis		100.0	0.37	FRU 99			
45425322	**Good identification**						**45425322**
Parabacteroides distasonis		100.0	0.46	FRU 99			
45425332	**Good identification**						**45425332**
Parabacteroides distasonis		100.0	0.41	FRU 99			
45425722	**Good identification**						**45425722**
Parabacteroides distasonis		100.0	0.42	FRU 99			
45425732	**Good identification**						**45425732**
Parabacteroides distasonis		100.0	0.37	FRU 99			
45430000	**Low discrimination**						**45430000**
					GRA	CAT	
Fusobacterium mortiferum		82.69	0.64		1	5	
Propionibacterium granulosum		12.58	0.47		99	90	
Eubacterium limosum		1.89	0.18		99	1	
45430002	**Low discrimination**						**45430002**
					GRA	CAT	
Fusobacterium mortiferum		72.96	0.48	BGL 10	1	5	
Propionibacterium granulosum		24.97	0.36		99	90	
Atopobium minutum		1.34	0.12		99	1	
45430020	**Low discrimination**						**45430020**
					GRA	CAT	
Propionibacterium granulosum		77.38	0.36	ESL 1	99	90	
Fusobacterium mortiferum		20.55	0.30	SUC 93	1	5	
Atopobium minutum		2.07	0.07		99	1	
45430100	**Very good identification**						**45430100**
					GRA	CAT	
Fusobacterium mortiferum		99.93	0.70		1	5	
Propionibacterium granulosum		0.07	0.13		99	90	
45430101	**Good identification**						**45430101**
Fusobacterium mortiferum		100.0	0.36	URE 1			
45430102	**Good identification**						**45430102**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.54	BGL 10			
45430110	**Good identification**						**45430110**
					GRA	SPO	
Fusobacterium mortiferum		96.22	0.36	MLZ 1	1	1	
Clostridium difficile		3.78	-.04		99	99	
45430120	**Good identification**						**45430120**
Fusobacterium mortiferum		100.0	0.36	NAG 1			
45430140	**Good identification**						**45430140**
Fusobacterium mortiferum		100.0	0.36	SOR 1			
45430300	**Good identification**						**45430300**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
45430500	**Good identification**						**45430500**
Fusobacterium mortiferum		100.0	0.36	ARA 1			
45431000	**Good identification**						**45431000**
					GRA	CAT	
Fusobacterium mortiferum		90.74	0.67		1	5	
Propionibacterium granulosum		9.20	0.47		99	90	
Atopobium minutum		0.06	0.07		99	1	
45431002	**Low discrimination**						**45431002**
					GRA	CAT	
Fusobacterium mortiferum		81.42	0.51	BGL 10	1	5	
Propionibacterium granulosum		18.58	0.36		99	90	
45431020	**Low discrimination**						**45431020**
					GRA	CAT	
Propionibacterium granulosum		67.06	0.36	ESL 1	99	90	
Fusobacterium mortiferum		26.71	0.33	SUC 93	1	5	
Clostridium perfringens		6.23	0.11		99	1	
45431100	**Very good identification**						**45431100**
					GRA	CAT	
Fusobacterium mortiferum		99.93	0.73		1	5	
Propionibacterium granulosum		0.04	0.13		99	90	
Clostridium perfringens		0.02	0.00		99	1	
45431101	**Good identification**						**45431101**
Fusobacterium mortiferum		100.0	0.39	URE 1			
45431102	**Very good identification**						**45431102**
Fusobacterium mortiferum		99.04	0.57	BGL 10			
Parabacteroides distasonis		0.96	0.21				
45431110	**Good identification**						**45431110**
Fusobacterium mortiferum		100.0	0.39	MLZ 1			
45431122	**Acceptable identification**						**45431122**
					GRA		
Parabacteroides distasonis		93.71	0.55	XYL 90	1		
Clostridium perfringens		3.05	0.20		99		
Prevotella oralis		1.91	0.14		1		
45431132	**Good identification**						**45431132**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		100.0	0.51	XYL 90				
45431140	**Good identification**							**45431140**
Fusobacterium mortiferum		100.0	0.39	SOR 1				
45431300	**Good identification**							**45431300**
Fusobacterium mortiferum		100.0	0.39	RHA 1				
45431322	**Very good identification**							**45431322**
Parabacteroides distasonis		99.71	0.63	XYL 90				
Prevotella oralis		0.29	0.07					
45431332	**Very good identification**							**45431332**
Parabacteroides distasonis		100.0	0.59	XYL 90				
45431500	**Good identification**							**45431500**
Fusobacterium mortiferum		100.0	0.39	ARA 1				
45431522	**Good identification**							**45431522**
					GRA			
Parabacteroides distasonis		99.41	0.52	XYL 90	1			
Bifidobacterium longum subsp. longum		0.59	0.09		99			
45431532	**Good identification**							**45431532**
					GRA			
Parabacteroides distasonis		98.17	0.47	XYL 90	1			
Bifidobacterium longum subsp. longum		1.83	0.13		99			
45431722	**Very good identification**							**45431722**
Parabacteroides distasonis		100.0	0.60	XYL 90				
45431732	**Good identification**							**45431732**
Parabacteroides distasonis		100.0	0.55	XYL 90				
45432010	**Good identification**							**45432010**
					SPO	CAT		
Clostridium difficile		97.97	0.47	RAF 1	99	1		
Propionibacterium granulosum		1.86	0.30	TRE 7	1	90		
Eubacterium limosum		0.17	-.03		1	1		
45432100	**Low discrimination**							**45432100**
					GRA	SPO		
Fusobacterium mortiferum		84.62	0.36	MAN 1	1	1		
Clostridium difficile		6.13	0.01		99	99		
Eubacterium limosum		5.09	-.03		99	1		
45433000	**Low discrimination**							**45433000**
					GRA	SPO	CAT	
Propionibacterium granulosum		86.26	0.45	ESL 1	99	1	90	
Fusobacterium mortiferum		11.39	0.33	SUC 93	1	1	5	
Clostridium difficile		1.28	0.01		99	99	1	
45433100	**Good identification**							**45433100**
					GRA	CAT		
Fusobacterium mortiferum		96.83	0.39	MAN 1	1	5		
Propionibacterium granulosum		3.17	0.10		99	90		
45434000	**Good identification**							**45434000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Fusobacterium mortiferum		96.75	0.45	XYL 7	1	1	5	
Propionibacterium granulosum		1.98	0.13		99	1	90	
Clostridium difficile		1.27	-.03		99	99	1	
45434100	**Good identification**							**45434100**
Fusobacterium mortiferum		100.0	0.51	XYL 7				
45434122	**Good identification**							**45434122**
Parabacteroides distasonis		100.0	0.37	GAL 99				
45434322	**Good identification**							**45434322**
Parabacteroides distasonis		100.0	0.46	GAL 99				
45434332	**Good identification**							**45434332**
Parabacteroides distasonis		100.0	0.41	GAL 99				
45434722	**Good identification**							**45434722**
Parabacteroides distasonis		100.0	0.42	GAL 99				
45434732	**Good identification**							**45434732**
Parabacteroides distasonis		100.0	0.37	GAL 99				
45435000	**Good identification**							**45435000**
					GRA	CAT		
Fusobacterium mortiferum		98.66	0.48	XYL 7	1	5		
Propionibacterium granulosum		1.34	0.13		99	90		
45435022	**Good identification**							**45435022**
Parabacteroides distasonis		100.0	0.37	LAC 99				
45435100	**Good identification**							**45435100**
					COC	GRA		
Fusobacterium mortiferum		99.69	0.54	XYL 7	1	1		
Atopobium parvulum		0.31	0.00		99	99		
45435102	**Low discrimination**							**45435102**
Parabacteroides distasonis		53.79	0.37	NAG 99				
Fusobacterium mortiferum		46.21	0.38					
45435120	**Acceptable identification**							**45435120**
Parabacteroides distasonis		92.76	0.37	BGL 99				
Fusobacterium mortiferum		7.25	0.20					
45435122	**Very good identification**							**45435122**
					CAT			
Parabacteroides distasonis		99.89	0.72		76			
Bacteroides fragilis		0.09	-.03		99			
Prevotella oralis		0.03	-.02		1			
45435123	**Good identification**							**45435123**
Parabacteroides distasonis		100.0	0.37	URE 1				
45435132	**Very good identification**							**45435132**
Parabacteroides distasonis		100.0	0.67					

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
45435162	**Good identification**					**45435162**
Parabacteroides distasonis		100.0	0.37	SOR 1		
45435222	**Good identification**					**45435222**
Parabacteroides distasonis		100.0	0.46	LAC 99		
45435232	**Good identification**					**45435232**
Parabacteroides distasonis		100.0	0.41	LAC 99		
45435302	**Good identification**					**45435302**
Parabacteroides distasonis		100.0	0.46	NAG 99		
45435312	**Good identification**					**45435312**
Parabacteroides distasonis		100.0	0.41	NAG 99		
45435320	**Good identification**					**45435320**
Parabacteroides distasonis		97.68	0.46	BGL 99		
Bacteroides vulgatus		2.32	-.01			
45435322	**Very good identification**					**45435322**
Parabacteroides distasonis		100.0	0.80			
45435323	**Good identification**					**45435323**
Parabacteroides distasonis		100.0	0.46	URE 1		
45435330	**Good identification**					**45435330**
Parabacteroides distasonis		100.0	0.41	BGL 99		
45435332	**Very good identification**					**45435332**
Parabacteroides distasonis		100.0	0.75			
45435333	**Good identification**					**45435333**
Parabacteroides distasonis		100.0	0.41	URE 1		
45435362	**Good identification**					**45435362**
Parabacteroides distasonis		100.0	0.46	SOR 1		
45435372	**Good identification**					**45435372**
Parabacteroides distasonis		100.0	0.41	SOR 1		
45435522	**Very good identification**					**45435522**
					GRA	
Parabacteroides distasonis		99.80	0.68		1	
Bifidobacterium longum subsp. longum		0.20	0.17		99	
45435532	**Very good identification**					**45435532**
					GRA	
Parabacteroides distasonis		99.38	0.63		1	
Bifidobacterium longum subsp. longum		0.62	0.21		99	
45435622	**Good identification**					**45435622**
Parabacteroides distasonis		100.0	0.42	LAC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
45435632	**Good identification**					**45435632**
Parabacteroides distasonis		100.0	0.37	LAC 99		
45435702	**Good identification**					**45435702**
Parabacteroides distasonis		100.0	0.42	NAG 99		
45435712	**Good identification**					**45435712**
Parabacteroides distasonis		96.06	0.37	NAG 99		
Clostridium clostridioforme		3.94	0.08			
45435720	**Low discrimination**					**45435720**
Parabacteroides distasonis		65.98	0.42	BGL 99		
Bacteroides vulgatus		34.02	0.19			
45435722	**Very good identification**					**45435722**
Parabacteroides distasonis		99.87	0.76			
Bacteroides thetaiotaomicron		0.13	0.11			
45435723	**Good identification**					**45435723**
Parabacteroides distasonis		100.0	0.42	URE 1		
45435730	**Good identification**					**45435730**
Parabacteroides distasonis		100.0	0.37	BGL 99		
45435732	**Very good identification**					**45435732**
Parabacteroides distasonis		99.86	0.71			
Bacteroides thetaiotaomicron		0.10	0.04			
Clostridium clostridioforme		0.04	0.08			
45435733	**Good identification**					**45435733**
Parabacteroides distasonis		100.0	0.37	URE 1		
45435762	**Good identification**					**45435762**
Parabacteroides distasonis		100.0	0.42	SOR 1		
45435772	**Good identification**					**45435772**
Parabacteroides distasonis		100.0	0.37	SOR 1		
45437122	**Good identification**					**45437122**
Parabacteroides distasonis		100.0	0.37	MAN 1		
45437322	**Good identification**					**45437322**
Parabacteroides distasonis		100.0	0.46	MAN 1		
45437332	**Good identification**					**45437332**
Parabacteroides distasonis		100.0	0.41	MAN 1		
45437722	**Good identification**					**45437722**
Parabacteroides distasonis		100.0	0.42	MAN 1		
45437732	**Good identification**					**45437732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.37	MAN 1			
45440100	**Good identification**						**45440100**
Fusobacterium mortiferum		100.0	0.37	FRU 99			
45441100	**Good identification**						**45441100**
					COC	GRA	
Fusobacterium mortiferum		96.65	0.40	FRU 99	1	1	
Blautia hansenii		3.35	-.03		99	99	
45450000	**Very good identification**						**45450000**
					GRA	SPO	
Fusobacterium mortiferum		96.62	0.65		1	1	
Eubacterium limosum		1.98	0.18		99	1	
Clostridium difficile		1.14	0.16		99	99	
45450002	**Good identification**						**45450002**
					GRA		
Fusobacterium mortiferum		98.39	0.48	BGL 10	1		
Atopobium minutum		1.61	0.12		99		
45450100	**Very good identification**						**45450100**
Fusobacterium mortiferum		100.0	0.71				
45450101	**Good identification**						**45450101**
Fusobacterium mortiferum		100.0	0.37	URE 1			
45450102	**Good identification**						**45450102**
Fusobacterium mortiferum		100.0	0.55	BGL 10			
45450110	**Good identification**						**45450110**
Fusobacterium mortiferum		100.0	0.37	MLZ 1			
45450120	**Good identification**						**45450120**
Fusobacterium mortiferum		100.0	0.37	NAG 1			
45450140	**Good identification**						**45450140**
Fusobacterium mortiferum		100.0	0.37	SOR 1			
45450300	**Good identification**						**45450300**
Fusobacterium mortiferum		100.0	0.37	RHA 1			
45450500	**Good identification**						**45450500**
Fusobacterium mortiferum		100.0	0.37	ARA 1			
45451000	**Very good identification**						**45451000**
					GRA		
Fusobacterium mortiferum		99.94	0.68		1		
Atopobium minutum		0.06	0.07		99		
45451002	**Good identification**						**45451002**
Fusobacterium mortiferum		99.24	0.51	BGL 10			
Prevotella oralis		0.76	0.01				
45451100	**Very good identification**						**45451100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		99.97	0.74			
Prevotella oralis		0.03	0.00			
45451101	**Good identification**					**45451101**
Fusobacterium mortiferum		100.0	0.40	URE 1		
45451102	**Good identification**					**45451102**
					CAT	
Fusobacterium mortiferum		94.78	0.58	BGL 10	5	
Prevotella oralis		4.88	0.22		1	
Capnocytophaga ochracea		0.34	0.02		90	
45451110	**Good identification**					**45451110**
Fusobacterium mortiferum		100.0	0.40	MLZ 1		
45451122	**Low discrimination**					**45451122**
					CAT	
Prevotella oralis		86.29	0.43	SUC 92	1	
Parabacteroides distasonis		9.00	0.39	MLT 99	76	
Capnocytophaga ochracea		2.86	0.18		90	
45451140	**Good identification**					**45451140**
Fusobacterium mortiferum		100.0	0.40	SOR 1		
45451300	**Good identification**					**45451300**
Fusobacterium mortiferum		100.0	0.40	RHA 1		
45451322	**Low discrimination**					**45451322**
Prevotella oralis		57.80	0.37	SUC 92		
Parabacteroides distasonis		42.20	0.48	MLT 99		
45451332	**Good identification**					**45451332**
Parabacteroides distasonis		97.50	0.43	XYL 90		
Prevotella oralis		2.51	0.03			
45451500	**Good identification**					**45451500**
Fusobacterium mortiferum		100.0	0.40	ARA 1		
45451522	**Low discrimination**					**45451522**
					CAT	
Parabacteroides distasonis		80.04	0.36	XYL 90	76	
Prevotella oralis		12.65	0.09		1	
Capnocytophaga ochracea		7.32	0.06		90	
45451722	**Good identification**					**45451722**
Parabacteroides distasonis		97.79	0.44	XYL 90		
Prevotella oralis		2.21	0.03			
45451732	**Good identification**					**45451732**
Parabacteroides distasonis		100.0	0.39	XYL 90		
45452010	**Good identification**					**45452010**
					SPO	
Clostridium difficile		99.76	0.44	RAF 1	99	
Eubacterium limosum		0.25	-.03	CEL 5	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45452100	**Acceptable identification**						**45452100**
					GRA	SPO	
Fusobacterium mortiferum		90.97	0.37	MAN 1	1	1	
Eubacterium limosum		4.91	-.03		99	1	
Clostridium difficile		4.13	-.02		99	99	
45453100	**Good identification**						**45453100**
Fusobacterium mortiferum		100.0	0.40	MAN 1			
45454000	**Good identification**						**45454000**
Fusobacterium mortiferum		100.0	0.46	XYL 7			
45454100	**Good identification**						**45454100**
Fusobacterium mortiferum		100.0	0.52	XYL 7			
45454102	**Good identification**						**45454102**
Fusobacterium mortiferum		100.0	0.35	XYL 7			
45455000	**Good identification**						**45455000**
Fusobacterium mortiferum		100.0	0.49	XYL 7			
45455100	**Good identification**						**45455100**
Fusobacterium mortiferum		100.0	0.55	XYL 7			
45455102	**Low discrimination**						**45455102**
Fusobacterium mortiferum		80.22	0.38	XYL 7			
Parabacteroides distasonis		9.89	0.22	BGL 10			
Prevotella oralis		6.78	0.06				
45455122	**Low discrimination**						**45455122**
					CAT		
Parabacteroides distasonis		68.73	0.56		76		
Bacteroides fragilis		21.61	0.25		99		
Prevotella oralis		9.05	0.28		1		
45455132	**Good identification**						**45455132**
Parabacteroides distasonis		99.41	0.51				
Bacteroides fragilis		0.59	-.09				
45455322	**Very good identification**						**45455322**
					CAT		
Parabacteroides distasonis		98.05	0.64		76		
Prevotella oralis		1.84	0.22		1		
Bacteroides fragilis		0.10	-.09		99		
45455332	**Very good identification**						**45455332**
Parabacteroides distasonis		100.0	0.59				
45455522	**Good identification**						**45455522**
					CAT		
Parabacteroides distasonis		98.60	0.52		76		
Bacteroides uniformis		0.89	-.02		5		
Bacteroides fragilis		0.51	-.09		99		
45455532	**Good identification**						**45455532**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		100.0	0.47					
45455722	**Very good identification**							**45455722**
					CAT			
Parabacteroides distasonis		99.41	0.60		76			
Bacteroides uniformis		0.41	0.01		5			
Bacteroides thetaiotaomicron		0.18	-.03		89			
45455732	**Good identification**							**45455732**
Parabacteroides distasonis		100.0	0.56					
45465322	**Good identification**							**45465322**
Parabacteroides distasonis		100.0	0.41	FRU 99				
45465332	**Good identification**							**45465332**
Parabacteroides distasonis		100.0	0.36	FRU 99				
45465722	**Good identification**							**45465722**
Parabacteroides distasonis		100.0	0.37	FRU 99				
45470000	**Good identification**							**45470000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		99.25	0.56		1	1	5	
Propionibacterium granulosum		0.46	0.13		99	1	90	
Clostridium difficile		0.29	-.03		99	99	1	
45470002	**Good identification**							**45470002**
Fusobacterium mortiferum		100.0	0.39	BGL 10				
45470100	**Very good identification**							**45470100**
Fusobacterium mortiferum		100.0	0.62					
45470102	**Good identification**							**45470102**
Fusobacterium mortiferum		100.0	0.46	BGL 10				
45471000	**Very good identification**							**45471000**
					GRA	CAT		
Fusobacterium mortiferum		99.39	0.59		1	5		
Propionibacterium granulosum		0.31	0.13		99	90		
Lactobacillus jensenii		0.30	0.06		99	1		
45471002	**Good identification**							**45471002**
					GRA			
Fusobacterium mortiferum		98.20	0.42	BGL 10	1			
Lactobacillus jensenii		1.80	0.03		99			
45471100	**Very good identification**							**45471100**
					COC	GRA	SPO	
Fusobacterium mortiferum		99.72	0.65		1	1	1	
Clostridium septicum		0.19	0.01		1	99	99	
Atopobium parvulum		0.09	0.01		99	99	1	
45471102	**Good identification**							**45471102**
Fusobacterium mortiferum		97.40	0.49	BGL 10				
Parabacteroides distasonis		1.53	0.16					
Prevotella oralis		1.07	0.01					
45471120	**Acceptable identification**							**45471120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium septicum		93.01	0.35	MLT 94	99	99	
Fusobacterium mortiferum		5.05	0.31	RAF 1	1	1	
Parabacteroides distasonis		0.87	0.16		1	1	
45471122	**Low discrimination**						**45471122**
					GRA	SPO	
Parabacteroides distasonis		86.00	0.51	XYL 90	1	1	
Prevotella oralis		11.55	0.23		1	1	
Clostridium septicum		1.89	0.06		99	99	
45471132	**Good identification**						**45471132**
Parabacteroides distasonis		100.0	0.46	XYL 90			
45471322	**Very good identification**						**45471322**
Parabacteroides distasonis		98.12	0.59	XYL 90			
Prevotella oralis		1.88	0.17				
45471332	**Good identification**						**45471332**
Parabacteroides distasonis		100.0	0.54	XYL 90			
45471522	**Good identification**						**45471522**
Parabacteroides distasonis		100.0	0.47	XYL 90			
45471532	**Good identification**						**45471532**
Parabacteroides distasonis		100.0	0.42	XYL 90			
45471722	**Good identification**						**45471722**
Parabacteroides distasonis		100.0	0.55	XYL 90			
45471732	**Good identification**						**45471732**
Parabacteroides distasonis		100.0	0.51	XYL 90			
45474000	**Good identification**						**45474000**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
45474100	**Good identification**						**45474100**
Fusobacterium mortiferum		100.0	0.43	XYL 7			
45474322	**Good identification**						**45474322**
Parabacteroides distasonis		100.0	0.41	GAL 99			
45474332	**Good identification**						**45474332**
Parabacteroides distasonis		100.0	0.36	GAL 99			
45474722	**Good identification**						**45474722**
Parabacteroides distasonis		100.0	0.37	GAL 99			
45475000	**Good identification**						**45475000**
Fusobacterium mortiferum		100.0	0.40	XYL 7			
45475100	**Good identification**						**45475100**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		97.86	0.46	XYL 7	1	1	
Atopobium parvulum		2.14	0.06		99	99	
45475122	**Very good identification**						**45475122**
					CAT		
Parabacteroides distasonis		99.75	0.67		76		
Prevotella oralis		0.18	0.07		1		
Bacteroides fragilis		0.07	-.09		99		
45475132	**Very good identification**						**45475132**
Parabacteroides distasonis		100.0	0.62				
45475222	**Good identification**						**45475222**
Parabacteroides distasonis		100.0	0.41	LAC 99			
45475232	**Good identification**						**45475232**
Parabacteroides distasonis		100.0	0.36	LAC 99			
45475302	**Good identification**						**45475302**
Parabacteroides distasonis		100.0	0.41	NAG 99			
45475312	**Good identification**						**45475312**
Parabacteroides distasonis		100.0	0.36	NAG 99			
45475320	**Good identification**						**45475320**
Parabacteroides distasonis		100.0	0.41	BGL 99			
45475322	**Very good identification**						**45475322**
Parabacteroides distasonis		99.97	0.75				
Prevotella oralis		0.03	0.01				
45475323	**Good identification**						**45475323**
Parabacteroides distasonis		100.0	0.41	URE 1			
45475330	**Good identification**						**45475330**
Parabacteroides distasonis		100.0	0.36	BGL 99			
45475332	**Very good identification**						**45475332**
Parabacteroides distasonis		100.0	0.71				
45475333	**Good identification**						**45475333**
Parabacteroides distasonis		100.0	0.36	URE 1			
45475362	**Good identification**						**45475362**
Parabacteroides distasonis		100.0	0.41	SOR 1			
45475372	**Good identification**						**45475372**
Parabacteroides distasonis		100.0	0.36	SOR 1			
45475522	**Very good identification**						**45475522**
Parabacteroides distasonis		99.88	0.63				
Bacteroides thetaiotaomicron		0.12	-.03				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45475532	**Very good identification**						**45475532**
Parabacteroides distasonis		100.0	0.59				
45475622	**Good identification**						**45475622**
Parabacteroides distasonis		100.0	0.37	LAC 99			
45475702	**Good identification**						**45475702**
Parabacteroides distasonis		97.08	0.37	NAG 99			
Clostridium clostridioforme		2.93	0.06				
45475720	**Good identification**						**45475720**
Parabacteroides distasonis		95.20	0.37	BGL 99			
Bacteroides vulgatus		4.80	-.03				
45475722	**Very good identification**						**45475722**
Parabacteroides distasonis		99.44	0.71				
Bacteroides thetaiotaomicron		0.53	0.17				
Clostridium clostridioforme		0.03	0.06				
45475723	**Good identification**						**45475723**
Parabacteroides distasonis		100.0	0.37	URE 1			
45475732	**Very good identification**						**45475732**
Parabacteroides distasonis		99.37	0.67				
Bacteroides thetaiotaomicron		0.38	0.09				
Clostridium clostridioforme		0.26	0.17				
45475762	**Good identification**						**45475762**
Parabacteroides distasonis		100.0	0.37	SOR 1			
45477322	**Good identification**						**45477322**
Parabacteroides distasonis		100.0	0.41	MAN 1			
45477332	**Good identification**						**45477332**
Parabacteroides distasonis		100.0	0.36	MAN 1			
45477722	**Good identification**						**45477722**
Parabacteroides distasonis		100.0	0.37	MAN 1			
45500000	**Low discrimination**						**45500000**
					GRA	SPO	
Clostridium bifermentans		52.91	0.50	IND 90	99	99	
Clostridium botulinum biovar A		27.17	0.45	RAF 1	99	99	
Prevotella disiens		9.70	0.32		1	1	
45500040	**Low discrimination**						**45500040**
Clostridium botulinum biovar A		76.23	0.36	MNS 7			
Clostridium bifermentans		20.71	0.26	RAF 1			
Clostridium novyi biovar A		1.01	0.16				
45501100	**Very good identification**						**45501100**
					COC	GRA	
Blautia hansenii		99.81	0.66	MNS 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		0.11	0.33		1	1		
Prevotella melaninogenica		0.08	0.12		1	1		
45501102	**Good identification**							**45501102**
					COC	CAT	GRA	
Blautia hansenii		98.87	0.50	MNS 1	99	1	99	
Prevotella oralis		0.42	0.12	BGL 10	1	1	1	
Capnocytophaga ochracea		0.42	0.12		1	90	1	
45501120	**Low discrimination**							**45501120**
					COC	GRA		
Blautia hansenii		57.05	0.50	MNS 1	99	99		
Prevotella melaninogenica		39.47	0.47	NAG 10	1	1		
Prevotella bivia		2.32	0.19		1	1		
45501500	**Good identification**							**45501500**
					COC			
Blautia hansenii		99.37	0.53	MNS 1	99			
Bifidobacterium longum subsp. longum		0.63	0.27		1			
45501502	**Acceptable identification**							**45501502**
					COC	GRA	CAT	
Blautia hansenii		94.26	0.37	MNS 1	99	99	1	
Bifidobacterium longum subsp. longum		5.34	0.27	BGL 10	1	99	1	
Capnocytophaga ochracea		0.40	-.01		1	1	90	
45501520	**Low discrimination**							**45501520**
					COC	GRA		
Blautia hansenii		78.03	0.37	MNS 1	99	99		
Bifidobacterium longum subsp. longum		17.68	0.37	NAG 10	1	99		
Prevotella melaninogenica		3.09	0.12		1	1		
45501522	**Low discrimination**							**45501522**
					COC	GRA	CAT	
Bifidobacterium longum subsp. longum		57.44	0.37	SUC 99	1	99	1	
Blautia hansenii		28.16	0.20	FRU 90	99	99	1	
Capnocytophaga ochracea		9.58	0.16		1	1	90	
45501530	**Good identification**							**45501530**
					COC			
Bifidobacterium longum subsp. longum		97.45	0.41	SUC 99	1			
Blautia hansenii		2.55	0.03	FRU 90	99			
45501532	**Good identification**							**45501532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		98.00	0.41	SUC 99	99	1		
Capnocytophaga ochracea		1.07	-.01	FRU 90	1	90		
Parabacteroides distasonis		0.94	0.11		1	76		
45505100	**Good identification**							**45505100**
					COC	GRA		
Blautia hansenii		99.95	0.53	MNS 1	99	99		
Fusobacterium mortiferum		0.05	0.13		1	1		
45505102	**Good identification**							**45505102**
					COC	GRA		
Blautia hansenii		98.31	0.37	MNS 1	99	99		
Bacteroides fragilis		1.11	-.03	BGL 10	1	1		
Capnocytophaga ochracea		0.59	0.02		1	1		
45505120	**Low discrimination**							**45505120**
					CAT	COC	GRA	
Blautia hansenii		88.06	0.37	MNS 1	1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides fragilis		5.16	0.10	NAG 10	99	1	1	
Prevotella melaninogenica		3.49	0.12		1	1	1	
45505322	**Low discrimination**							**45505322**
					CAT			
Parabacteroides distasonis		89.53	0.44	FRU 99	76			
Prevotella oralis		7.14	0.12		1			
Bacteroides fragilis		3.33	-.03		99			
45505332	**Good identification**							**45505332**
Parabacteroides distasonis		100.0	0.39	FRU 99				
45505500	**Acceptable identification**							**45505500**
					COC	GRA		
Blautia hansenii		90.16	0.40	MNS 1	99	99		
Bifidobacterium longum subsp. longum		9.65	0.35		1	99		
Bacteroides vulgatus		0.19	-.08		1	1		
45505510	**Acceptable identification**							**45505510**
					COC			
Bifidobacterium longum subsp. longum		94.75	0.39	SUC 99	1			
Blautia hansenii		5.25	0.06	FRU 90	99			
45505512	**Good identification**							**45505512**
Bifidobacterium longum subsp. longum		97.63	0.39	SUC 99				
Bifidobacterium adolescentis		2.37	0.13	FRU 90				
45505520	**Low discrimination**							**45505520**
					COC	GRA		
Bifidobacterium longum subsp. longum		57.42	0.46	SUC 99	1	99		
Bacteroides vulgatus		27.68	0.26	FRU 90	1	1		
Blautia hansenii		14.91	0.24		99	99		
45505522	**Low discrimination**							**45505522**
					COC	GRA		
Bifidobacterium longum subsp. longum		86.57	0.46	SUC 99	1	99		
Parabacteroides distasonis		7.87	0.32	FRU 90	1	1		
Blautia hansenii		2.50	0.07		99	99		
45505530	**Good identification**							**45505530**
					GRA			
Bifidobacterium longum subsp. longum		99.72	0.50	SUC 99	99			
Bacteroides vulgatus		0.29	-.08	FRU 90	1			
45505532	**Good identification**							**45505532**
					GRA			
Bifidobacterium longum subsp. longum		97.20	0.50	SUC 99	99			
Parabacteroides distasonis		2.80	0.27	FRU 90	1			
45505720	**Good identification**							**45505720**
					GRA			
Bacteroides vulgatus		99.77	0.42	SUC 96	1			
Bifidobacterium longum subsp. longum		0.23	0.11	FRU 99	99			
45505732	**Low discrimination**							**45505732**
					GRA			
Parabacteroides distasonis		89.52	0.35	FRU 99	1			
Bifidobacterium longum subsp. longum		10.48	0.15		99			
45510000	**Low discrimination**							**45510000**
Prevotella intermedia		62.12	0.55	ESL 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		20.07	0.57				
Prevotella disiens		8.11	0.32				
45510002	**Low discrimination**						**45510002**
					GRA	SPO	
Fusobacterium mortiferum		61.71	0.41	MLT 11	1	1	
Prevotella intermedia		17.36	0.20	BGL 10	1	1	
Clostridium sporogenes		8.29	0.11		99	99	
45510100	**Very good identification**						**45510100**
Fusobacterium mortiferum		98.28	0.64	MLT 11			
Prevotella intermedia		1.32	0.20				
Prevotella disiens		0.17	-.03				
45510102	**Low discrimination**						**45510102**
					CAT		
Fusobacterium mortiferum		88.94	0.47	MLT 11	5		
Prevotella oralis		5.23	0.12	BGL 10	1		
Capnocytophaga ochracea		5.18	0.12		90		
45511000	**Low discrimination**						**45511000**
					GRA		
Prevotella intermedia		64.08	0.55	ESL 1	1		
Fusobacterium mortiferum		31.06	0.60		1		
Lactobacillus fermentum		1.62	0.40		99		
45511002	**Low discrimination**						**45511002**
					GRA		
Fusobacterium mortiferum		40.67	0.44	MLT 11	1		
Prevotella oralis		23.49	0.26	BGL 10	1		
Lactobacillus fermentum		19.08	0.40		99		
45511020	**Low discrimination**						**45511020**
					GRA		
Clostridium perfringens		50.84	0.41	SUC 91	99		
Prevotella melaninogenica		25.27	0.29	LAC 88	1		
Prevotella oralis		10.75	0.26		1		
45511022	**Low discrimination**						**45511022**
					GRA	CAT	
Prevotella oralis		80.59	0.48	SUC 92	1	1	
Clostridium perfringens		9.44	0.35	LAC 94	99	1	
Capnocytophaga ochracea		5.99	0.29		1	90	
45511100	**Low discrimination**						**45511100**
					COC	GRA	
Fusobacterium mortiferum		76.08	0.67	MLT 11	1	1	
Blautia hansenii		7.11	0.32		99	99	
Prevotella melaninogenica		4.92	0.29		1	1	
45511102	**Low discrimination**						**45511102**
					CAT		
Prevotella oralis		39.00	0.46	SUC 92	1		
Capnocytophaga ochracea		38.68	0.47	NAG 95	90		
Fusobacterium mortiferum		10.06	0.50		5		
45511120	**Low discrimination**						**45511120**
					GRA		
Prevotella melaninogenica		75.01	0.63	SUC 99	1		
Clostridium perfringens		11.18	0.56		99		
Prevotella oralis		5.05	0.46		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45511122	**Low discrimination**						**45511122**
					CAT		
Prevotella oralis		57.55	0.68	SUC 92	1		
Capnocytophaga ochracea		27.04	0.63		90		
Prevotella melaninogenica		8.57	0.51		1		
45511132	**Low discrimination**						**45511132**
					CAT		
Capnocytophaga ochracea		66.29	0.47	SUC 99	90		
Parabacteroides distasonis		16.82	0.49	MLZ 10	76		
Prevotella oralis		12.83	0.34		1		
45511160	**Low discrimination**						**45511160**
					GRA	CAT	
Clostridium perfringens		60.43	0.43	SUC 91	99	1	
Prevotella melaninogenica		21.51	0.29		1	1	
Capnocytophaga ochracea		15.79	0.30		1	90	
45511162	**Low discrimination**						**45511162**
					GRA	CAT	
Capnocytophaga ochracea		69.72	0.47	SUC 99	1	90	
Clostridium perfringens		13.95	0.38	SOR 10	99	1	
Prevotella oralis		13.49	0.34		1	1	
45511222	**Good identification**						**45511222**
					GRA		
Prevotella oralis		97.09	0.41	SUC 92	1		
Parabacteroides distasonis		2.65	0.27	LAC 94	1		
Clostridium perfringens		0.27	0.01		99		
45511302	**Acceptable identification**						**45511302**
					CAT		
Prevotella oralis		93.86	0.40	SUC 92	1		
Parabacteroides distasonis		3.10	0.27	NAG 95	76		
Capnocytophaga ochracea		2.19	0.12		90		
45511320	**Low discrimination**						**45511320**
Prevotella oralis		52.27	0.40	SUC 92			
Bacteroides vulgatus		23.04	0.28	BGL 95			
Prevotella melaninogenica		18.31	0.29				
45511322	**Low discrimination**						**45511322**
					CAT		
Prevotella oralis		84.09	0.62	SUC 92	1		
Parabacteroides distasonis		14.48	0.62		76		
Capnocytophaga ochracea		0.93	0.29		90		
45511332	**Low discrimination**						**45511332**
					CAT		
Parabacteroides distasonis		89.11	0.57	XYL 90	76		
Prevotella oralis		9.71	0.28		1		
Capnocytophaga ochracea		1.18	0.12		90		
45511400	**Low discrimination**						**45511400**
					GRA		
Lactobacillus fermentum		57.73	0.40	ESL 1	99		
Prevotella intermedia		23.08	0.20	SUC 87	1		
Fusobacterium mortiferum		11.19	0.26		1		
45511402	**Acceptable identification**						**45511402**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		92.94	0.40	ESL 1	99	1	
Bifidobacterium longum subsp. longum		3.89	0.09	SUC 87	99	1	
Capnocytophaga ochracea		3.17	-.01		1	90	
45511500	**Low discrimination**						**45511500**
					COC		
Bifidobacterium longum subsp. longum		55.80	0.43	SUC 99	1		
Lactobacillus fermentum		21.98	0.44		1		
Blautia hansenii		9.95	0.19		99		
45511502	**Low discrimination**						**45511502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		40.42	0.43	SUC 99	99	1	
Capnocytophaga ochracea		32.94	0.34		1	90	
Lactobacillus fermentum		15.92	0.44		99	1	
45511510	**Good identification**						**45511510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.41	0.47	SUC 99	99	1	
Capnocytophaga ochracea		0.59	0.01		1	90	
45511512	**Low discrimination**						**45511512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		84.95	0.47	SUC 99	99	1	
Bifidobacterium adolescentis		10.17	0.33		99	1	
Capnocytophaga ochracea		4.52	0.17		1	90	
45511520	**Low discrimination**						**45511520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		55.78	0.54	SUC 99	99	1	
Bacteroides vulgatus		18.11	0.31		1	3	
Capnocytophaga ochracea		11.37	0.34		1	90	
45511522	**Low discrimination**						**45511522**
					GRA	CAT	
Capnocytophaga ochracea		52.69	0.50	SUC 99	1	90	
Bifidobacterium longum subsp. longum		28.73	0.54		99	1	
Parabacteroides distasonis		9.58	0.50		1	76	
45511530	**Very good identification**						**45511530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.30	0.58	SUC 99	99	1	
Capnocytophaga ochracea		1.31	0.17		1	90	
Bacteroides vulgatus		0.19	-.03		1	3	
45511532	**Low discrimination**						**45511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		81.54	0.58	SUC 99	99	1	
Capnocytophaga ochracea		9.76	0.34		1	90	
Parabacteroides distasonis		8.60	0.45		1	76	
45511720	**Good identification**						**45511720**
					GRA		
Bacteroides vulgatus		99.02	0.47	SUC 96	1		
Parabacteroides distasonis		0.34	0.24	XYL 98	1		
Bifidobacterium longum subsp. longum		0.34	0.20		99		
45511722	**Low discrimination**						**45511722**
Parabacteroides distasonis		86.67	0.58	XYL 90			
Prevotella oralis		8.30	0.28				
Bacteroides vulgatus		2.56	0.13				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45511732	**Good identification**						**45511732**
					GRA		
Parabacteroides distasonis		95.96	0.53	XYL 90	1		
Bifidobacterium longum subsp. longum		3.06	0.24		99		
Clostridium clostridioforme		0.61	0.10		50		
45514000	**Low discrimination**						**45514000**
					COC	GRA	
Fusobacterium mortiferum		55.37	0.38	MLT 11	1	1	
Prevotella intermedia		22.99	0.20	XYL 7	1	1	
Gemella morbillorum		10.33	0.15		99	99	
45514100	**Good identification**						**45514100**
Fusobacterium mortiferum		100.0	0.44	MLT 11			
45514322	**Low discrimination**						**45514322**
					CAT		
Parabacteroides distasonis		89.53	0.44	GAL 99	76		
Prevotella oralis		7.14	0.12		1		
Bacteroides fragilis		3.33	-.03		99		
45514332	**Good identification**						**45514332**
Parabacteroides distasonis		100.0	0.39	GAL 99			
45514720	**Good identification**						**45514720**
Bacteroides vulgatus		100.0	0.42	SUC 96			
45514722	**Low discrimination**						**45514722**
Parabacteroides distasonis		84.13	0.40	GAL 99			
Bacteroides vulgatus		13.51	0.08				
Clostridium clostridioforme		2.36	0.08				
45514732	**Low discrimination**						**45514732**
Parabacteroides distasonis		80.84	0.35	GAL 99			
Clostridium clostridioforme		19.16	0.19				
45515000	**Low discrimination**						**45515000**
					GRA		
Fusobacterium mortiferum		57.13	0.41	MLT 11	1		
Lactobacillus fermentum		23.23	0.36	XYL 7	99		
Prevotella intermedia		15.82	0.20		1		
45515002	**Low discrimination**						**45515002**
					GRA		
Lactobacillus fermentum		56.49	0.36	ESL 1	99		
Fusobacterium mortiferum		15.44	0.25	SUC 87	1		
Prevotella oralis		14.64	0.10		1		
45515100	**Low discrimination**						**45515100**
					COC	GRA	
Fusobacterium mortiferum		49.46	0.47	MLT 11	1	1	
Lactobacillus fermentum		14.06	0.40	XYL 7	1	99	
Blautia hansenii		10.84	0.19		99	99	
45515120	**Low discrimination**						**45515120**
					CAT		
Bacteroides fragilis		59.14	0.44	SUC 99	99		
Bacteroides vulgatus		24.90	0.41	BGL 95	3		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Capnocytophaga ochracea		6.00	0.36		90	
45515122	**Low discrimination**					**45515122**
					CAT	
Bacteroides fragilis		83.66	0.66	SUC 99	99	
Parabacteroides distasonis		7.58	0.70		76	
Prevotella oralis		4.23	0.53		1	
45515132	**Low discrimination**					**45515132**
Parabacteroides distasonis		74.22	0.65			
Bacteroides fragilis		15.36	0.32			
Capnocytophaga ochracea		8.13	0.36			
45515222	**Low discrimination**					**45515222**
					CAT	
Parabacteroides distasonis		64.89	0.44	LAC 99	76	
Prevotella oralis		32.70	0.26		1	
Bacteroides fragilis		2.41	-.03		99	
45515232	**Good identification**					**45515232**
Parabacteroides distasonis		100.0	0.39	LAC 99		
45515302	**Low discrimination**					**45515302**
					GRA	
Parabacteroides distasonis		65.38	0.44	NAG 99	1	
Prevotella oralis		27.17	0.24		1	
Eubacterium contortum		3.73	0.11		99	
45515312	**Good identification**					**45515312**
Parabacteroides distasonis		97.29	0.39	NAG 99		
Clostridium clostridioforme		2.71	0.07			
45515320	**Very good identification**					**45515320**
Bacteroides vulgatus		97.72	0.57	SUC 96		
Parabacteroides distasonis		1.35	0.44	ARA 93		
Prevotella oralis		0.56	0.24			
45515322	**Low discrimination**					**45515322**
					CAT	
Parabacteroides distasonis		88.78	0.78		76	
Prevotella oralis		7.08	0.46		1	
Bacteroides fragilis		3.30	0.32		99	
45515323	**Low discrimination**					**45515323**
					CAT	
Parabacteroides distasonis		89.53	0.44	URE 1	76	
Prevotella oralis		7.14	0.12		1	
Bacteroides fragilis		3.33	-.03		99	
45515332	**Very good identification**					**45515332**
					CAT	
Parabacteroides distasonis		99.72	0.73		76	
Prevotella oralis		0.15	0.12		1	
Bacteroides fragilis		0.07	-.03		99	
45515333	**Good identification**					**45515333**
Parabacteroides distasonis		100.0	0.39	URE 1		
45515362	**Low discrimination**					**45515362**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Parabacteroides distasonis		87.99	0.44	SOR 1	76		
Prevotella oralis		7.02	0.12		1		
Bacteroides fragilis		3.27	-.03		99		
45515372	**Good identification**						**45515372**
Parabacteroides distasonis		100.0	0.39	SOR 1			
45515400	**Low discrimination**						**45515400**
					GRA		
Lactobacillus fermentum		79.39	0.36	ESL 1	99		
Bifidobacterium longum subsp. longum		16.97	0.17	SUC 87	99		
Bacteroides vulgatus		3.64	-.08		1		
45515402	**Low discrimination**						**45515402**
Lactobacillus fermentum		77.32	0.36	ESL 1			
Bifidobacterium longum subsp. longum		16.53	0.17	SUC 87			
Bifidobacterium adolescentis		6.15	0.12				
45515500	**Low discrimination**						**45515500**
					GRA		
Bifidobacterium longum subsp. longum		76.00	0.52	SUC 99	99		
Bacteroides vulgatus		16.28	0.26		1		
Lactobacillus fermentum		5.86	0.40		99		
45515502	**Low discrimination**						**45515502**
Bifidobacterium longum subsp. longum		63.81	0.52	SUC 99			
Bifidobacterium adolescentis		23.75	0.46				
Lactobacillus fermentum		4.92	0.40				
45515510	**Good identification**						**45515510**
					GRA		
Bifidobacterium longum subsp. longum		99.61	0.56	SUC 99	99		
Bifidobacterium adolescentis		0.27	0.13		99		
Bacteroides vulgatus		0.13	-.08		1		
45515512	**Low discrimination**						**45515512**
					GRA		
Bifidobacterium longum subsp. longum		77.64	0.56	SUC 99	99		
Bifidobacterium adolescentis		20.74	0.47		99		
Parabacteroides distasonis		0.99	0.27		1		
45515520	**Low discrimination**						**45515520**
					GRA	CAT	
Bacteroides vulgatus		83.55	0.60	SUC 96	1	3	
Bifidobacterium longum subsp. longum		15.76	0.62	RHA 90	99	1	
Capnocytophaga ochracea		0.27	0.23		1	90	
45515522	**Low discrimination**						**45515522**
					GRA	CAT	
Parabacteroides distasonis		41.58	0.66		1	76	
Bifidobacterium longum subsp. longum		41.56	0.62		99	1	
Bacteroides fragilis		7.56	0.32		1	99	
45515530	**Very good identification**						**45515530**
					GRA		
Bifidobacterium longum subsp. longum		96.55	0.66	SUC 99	99		
Bacteroides vulgatus		3.04	0.26		1		
Parabacteroides distasonis		0.31	0.27		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45515532	**Low discrimination**						**45515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		75.22	0.66	SUC 99	99	1	
Parabacteroides distasonis		23.80	0.61		1	76	
Capnocytophaga ochracea		0.75	0.23		1	90	
45515620	**Good identification**						**45515620**
Bacteroides vulgatus		100.0	0.42	SUC 96			
45515622	**Low discrimination**						**45515622**
Parabacteroides distasonis		84.34	0.40	LAC 99			
Bacteroides vulgatus		13.54	0.08				
Clostridium clostridioforme		2.12	0.07				
45515632	**Low discrimination**						**45515632**
Parabacteroides distasonis		82.48	0.35	LAC 99			
Clostridium clostridioforme		17.52	0.18				
45515700	**Good identification**						**45515700**
					GRA		
Bacteroides vulgatus		96.86	0.42	SUC 96	1		
Eubacterium contortum		2.63	0.23	NAG 99	99		
Bifidobacterium longum subsp. longum		0.51	0.17		99		
45515702	**Low discrimination**						**45515702**
Parabacteroides distasonis		69.68	0.40	NAG 99			
Bacteroides vulgatus		11.19	0.08				
Clostridium clostridioforme		5.86	0.16				
45515712	**Low discrimination**						**45515712**
					GRA		
Parabacteroides distasonis		48.91	0.35	NAG 99	1		
Clostridium clostridioforme		34.77	0.27		50		
Bifidobacterium longum subsp. longum		12.88	0.21		99		
45515720	**Very good identification**						**45515720**
					GRA		
Bacteroides vulgatus		99.91	0.76	SUC 96	1		
Parabacteroides distasonis		0.06	0.40		1		
Bifidobacterium longum subsp. longum		0.02	0.28		99		
45515721	**Good identification**						**45515721**
Bacteroides vulgatus		100.0	0.42	SUC 96			
45515722	**Low discrimination**						**45515722**
					GRA		
Parabacteroides distasonis		85.52	0.74		1		
Bacteroides vulgatus		13.73	0.42		1		
Bifidobacterium longum subsp. longum		0.29	0.28		99		
45515723	**Low discrimination**						**45515723**
Parabacteroides distasonis		86.16	0.40	URE 1			
Bacteroides vulgatus		13.84	0.08				
45515730	**Acceptable identification**						**45515730**
					GRA		
Bacteroides vulgatus		93.50	0.42	SUC 96	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		3.34	0.32	MLZ 1	99	
Parabacteroides distasonis		3.17	0.35		1	
45515732	**Very good identification**					**45515732**
					GRA	
Parabacteroides distasonis		97.82	0.70		1	
Bifidobacterium longum subsp. longum		1.04	0.32		99	
Clostridium clostridioforme		0.70	0.27		50	
45515733	**Good identification**					**45515733**
Parabacteroides distasonis		100.0	0.35	URE 1		
45515760	**Good identification**					**45515760**
Bacteroides vulgatus		100.0	0.42	SUC 96		
45515762	**Low discrimination**					**45515762**
Parabacteroides distasonis		86.16	0.40	SOR 1		
Bacteroides vulgatus		13.84	0.08			
45515772	**Good identification**					**45515772**
Parabacteroides distasonis		100.0	0.35	SOR 1		
45517322	**Low discrimination**					**45517322**
					CAT	
Parabacteroides distasonis		89.53	0.44	MAN 1	76	
Prevotella oralis		7.14	0.12		1	
Bacteroides fragilis		3.33	-.03		99	
45517332	**Good identification**					**45517332**
Parabacteroides distasonis		100.0	0.39	MAN 1		
45517502	**Acceptable identification**					**45517502**
Bifidobacterium adolescentis		94.75	0.44	SUC 99		
Bifidobacterium longum subsp. longum		3.14	0.17			
Actinomyces israelii		2.11	0.22			
45517512	**Good identification**					**45517512**
Bifidobacterium adolescentis		95.58	0.46	SUC 99		
Bifidobacterium longum subsp. longum		4.42	0.21			
45517720	**Good identification**					**45517720**
Bacteroides vulgatus		100.0	0.42	SUC 96		
45517722	**Low discrimination**					**45517722**
Parabacteroides distasonis		86.16	0.40	MAN 1		
Bacteroides vulgatus		13.84	0.08			
45517732	**Good identification**					**45517732**
Parabacteroides distasonis		100.0	0.35	MAN 1		
45521322	**Acceptable identification**					**45521322**
Parabacteroides distasonis		92.48	0.39	FRU 99		
Prevotella oralis		7.53	0.07	XYL 90		
45521722	**Good identification**					**45521722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.35	FRU 99			
45525122	**Good identification**						**45525122**
					CAT		
Parabacteroides distasonis		96.87	0.47	FRU 99	76		
Bacteroides fragilis		2.37	-.03		99		
Prevotella oralis		0.76	-.02		1		
45525132	**Good identification**						**45525132**
Parabacteroides distasonis		100.0	0.42	FRU 99			
45525322	**Good identification**						**45525322**
Parabacteroides distasonis		100.0	0.55	FRU 99			
45525332	**Good identification**						**45525332**
Parabacteroides distasonis		100.0	0.50	FRU 99			
45525520	**Acceptable identification**						**45525520**
					GRA		
Bifidobacterium longum subsp. longum		92.56	0.35	SUC 99	99		
Bacteroides vulgatus		7.44	0.02	FRU 90	1		
45525522	**Low discrimination**						**45525522**
					GRA		
Parabacteroides distasonis		62.37	0.43	FRU 99	1		
Bifidobacterium longum subsp. longum		37.63	0.35		99		
45525530	**Good identification**						**45525530**
Bifidobacterium longum subsp. longum		100.0	0.39	SUC 99			
45525532	**Low discrimination**						**45525532**
					GRA		
Bifidobacterium longum subsp. longum		65.61	0.39	SUC 99	99		
Parabacteroides distasonis		34.39	0.39	FRU 90	1		
45525722	**Good identification**						**45525722**
Parabacteroides distasonis		99.41	0.51	FRU 99			
Bacteroides thetaiotaomicron		0.59	-.03				
45525732	**Good identification**						**45525732**
Parabacteroides distasonis		98.86	0.47	FRU 99			
Clostridium clostridioforme		1.14	0.08				
45530000	**Low discrimination**						**45530000**
					GRA	CAT	
Fusobacterium mortiferum		56.13	0.48	MLT 11	1	5	
Propionibacterium granulosum		31.04	0.41		99	90	
Prevotella intermedia		5.87	0.20		1	1	
45530100	**Good identification**						**45530100**
					GRA		
Fusobacterium mortiferum		99.72	0.55	MLT 11	1		
Clostridium perfringens		0.28	0.00		99		
45530102	**Good identification**						**45530102**
Fusobacterium mortiferum		100.0	0.38	MLT 11			
45530322	**Acceptable identification**						**45530322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		92.48	0.39	GAL 99			
Prevotella oralis		7.53	0.07	XYL 90			
45530722	**Good identification**						**45530722**
Parabacteroides distasonis		96.14	0.35	GAL 99			
Clostridium clostridioforme		3.86	0.05	XYL 90			
45531000	**Low discrimination**						**45531000**
					GRA	CAT	
Fusobacterium mortiferum		66.22	0.51	MLT 11	1	5	
Propionibacterium granulosum		24.41	0.41		99	90	
Prevotella intermedia		4.62	0.20		1	1	
45531002	**Low discrimination**						**45531002**
					GRA	CAT	
Fusobacterium mortiferum		43.55	0.35	MLT 11	1	5	
Propionibacterium granulosum		36.13	0.30	BGL 10	99	90	
Clostridium perfringens		10.77	0.14		99	1	
45531020	**Acceptable identification**						**45531020**
					CAT	SPO	
Clostridium perfringens		91.08	0.45	SUC 91	1	84	
Propionibacterium granulosum		4.45	0.30	LAC 88	90	1	
Clostridium septicum		2.82	0.13		1	99	
45531022	**Low discrimination**						**45531022**
					GRA		
Clostridium perfringens		71.09	0.40	SUC 91	99		
Prevotella oralis		20.86	0.27	LAC 88	1		
Parabacteroides distasonis		5.79	0.30		1		
45531100	**Low discrimination**						**45531100**
					COC	GRA	
Fusobacterium mortiferum		75.19	0.58	MLT 11	1	1	
Clostridium perfringens		13.79	0.34		1	99	
Atopobium parvulum		7.63	0.29		99	99	
45531102	**Low discrimination**						**45531102**
					GRA		
Fusobacterium mortiferum		30.34	0.41	MLT 11	1		
Prevotella oralis		25.14	0.26	BGL 10	1		
Clostridium perfringens		23.58	0.29		99		
45531120	**Low discrimination**						**45531120**
					GRA	SPO	
Clostridium perfringens		67.36	0.60	SUC 91	99	84	
Clostridium septicum		28.20	0.47		99	99	
Prevotella melaninogenica		2.46	0.29		1	1	
45531122	**Low discrimination**						**45531122**
					GRA		
Parabacteroides distasonis		39.09	0.65	XYL 90	1		
Clostridium perfringens		35.53	0.55		99		
Prevotella oralis		22.27	0.48		1		
45531132	**Very good identification**						**45531132**
					GRA		
Parabacteroides distasonis		96.18	0.60	XYL 90	1		
Clostridium perfringens		1.64	0.20		99		
Prevotella oralis		1.03	0.14		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45531160	**Good identification**						**45531160**
					GRA	SPO	
Clostridium perfringens		97.64	0.48	SUC 91	99	84	
Clostridium septicum		2.17	0.13		99	99	
Prevotella melaninogenica		0.19	-.06		1	1	
45531162	**Low discrimination**						**45531162**
					GRA		
Clostridium perfringens		89.39	0.42	SUC 91	99		
Parabacteroides distasonis		5.22	0.30		1		
Prevotella oralis		2.97	0.14		1		
45531222	**Low discrimination**						**45531222**
					GRA		
Parabacteroides distasonis		64.29	0.39	XYL 90	1		
Prevotella oralis		33.06	0.21	LAC 99	1		
Clostridium perfringens		2.66	0.06		99		
45531302	**Low discrimination**						**45531302**
Parabacteroides distasonis		70.23	0.39	XYL 90			
Prevotella oralis		29.78	0.20	NAG 99			
45531320	**Low discrimination**						**45531320**
					GRA		
Parabacteroides distasonis		40.28	0.39	XYL 90	1		
Clostridium perfringens		25.93	0.26	BGL 99	99		
Prevotella oralis		17.08	0.20		1		
45531322	**Good identification**						**45531322**
					GRA		
Parabacteroides distasonis		92.21	0.73	XYL 90	1		
Prevotella oralis		7.50	0.41		1		
Clostridium perfringens		0.28	0.20		99		
45531323	**Acceptable identification**						**45531323**
Parabacteroides distasonis		92.48	0.39	XYL 90			
Prevotella oralis		7.53	0.07	URE 1			
45531332	**Very good identification**						**45531332**
Parabacteroides distasonis		99.81	0.68	XYL 90			
Prevotella oralis		0.15	0.07				
Clostridium clostridioforme		0.04	0.04				
45531362	**Low discrimination**						**45531362**
					GRA		
Parabacteroides distasonis		87.79	0.39	XYL 90	1		
Prevotella oralis		7.14	0.07	SOR 1	1		
Clostridium perfringens		5.07	0.08		99		
45531510	**Good identification**						**45531510**
Bifidobacterium longum subsp. longum		100.0	0.37	SUC 99			
45531512	**Low discrimination**						**45531512**
					GRA		
Bifidobacterium longum subsp. longum		69.96	0.37	SUC 99	99		
Bifidobacterium adolescentis		22.34	0.30		99		
Parabacteroides distasonis		5.43	0.22		1		
45531520	**Low discrimination**						**45531520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium longum subsp. longum		71.17	0.43	SUC 99	1		
Clostridium perfringens		13.77	0.26		84		
Clostridium septicum		5.76	0.13		99		
45531522	**Low discrimination**						**45531522**
					GRA		
Parabacteroides distasonis		83.26	0.61	XYL 90	1		
Bifidobacterium longum subsp. longum		13.70	0.43		99		
Clostridium perfringens		1.25	0.20		99		
45531530	**Good identification**						**45531530**
					GRA		
Bifidobacterium longum subsp. longum		98.10	0.47	SUC 99	99		
Parabacteroides distasonis		1.90	0.22		1		
45531532	**Low discrimination**						**45531532**
					GRA		
Parabacteroides distasonis		65.48	0.56	XYL 90	1		
Bifidobacterium longum subsp. longum		34.08	0.47		99		
Clostridium clostridioforme		0.28	0.10		50		
45531622	**Good identification**						**45531622**
Parabacteroides distasonis		96.53	0.35	XYL 90			
Clostridium clostridioforme		3.47	0.05	LAC 99			
45531702	**Low discrimination**						**45531702**
Parabacteroides distasonis		89.25	0.35	XYL 90			
Clostridium clostridioforme		10.75	0.14	NAG 99			
45531722	**Very good identification**						**45531722**
					CAT		
Parabacteroides distasonis		99.47	0.69	XYL 90	76		
Bacteroides thetaiotaomicron		0.22	0.08		89		
Prevotella oralis		0.13	0.07		1		
45531723	**Good identification**						**45531723**
Parabacteroides distasonis		100.0	0.35	XYL 90			
45531732	**Very good identification**						**45531732**
					GRA		
Parabacteroides distasonis		98.65	0.65	XYL 90	1		
Clostridium clostridioforme		1.02	0.25		50		
Bifidobacterium longum subsp. longum		0.17	0.13		99		
45531762	**Good identification**						**45531762**
Parabacteroides distasonis		100.0	0.35	XYL 90			
45532000	**Acceptable identification**						**45532000**
					SPO	CAT	
Propionibacterium granulosum		92.18	0.39	ESL 1	1	90	
Clostridium difficile		3.05	0.01	SUC 93	99	1	
Eubacterium limosum		2.54	-.03		1	1	
45533000	**Acceptable identification**						**45533000**
					GRA	CAT	
Propionibacterium granulosum		93.25	0.39	ESL 1	99	90	
Fusobacterium mortiferum		3.39	0.17	SUC 93	1	5	
Actinomyces naeslundii		3.37	0.29		99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
45533100	**Low discrimination**					**45533100**
					GRA	
Actinomyces naeslundii		50.61	0.37	ESL 12	99	
Fusobacterium mortiferum		41.73	0.24	SUC 99	1	
Clostridium perfringens		7.66	0.00		99	
45533101	**Good identification**					**45533101**
Actinomyces naeslundii		100.0	0.40	ESL 12		
45533102	**Low discrimination**					**45533102**
Actinomyces naeslundii		64.56	0.37	ESL 12		
Bifidobacterium adolescentis		22.20	0.13	SUC 99		
Actinomyces israelii		13.24	0.16			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45533103	**Good identification**						**45533103**
Actinomyces naeslundii		100.0	0.40	ESL 12			
45533121	**Good identification**						**45533121**
Actinomyces naeslundii		100.0	0.40	ESL 12			
45533123	**Good identification**						**45533123**
Actinomyces naeslundii		100.0	0.40	ESL 12			
45533322	**Acceptable identification**						**45533322**
					SPO		
Parabacteroides distasonis		90.73	0.39	MAN 1	1		
Prevotella oralis		7.38	0.07	XYL 90	1		
Clostridium ramosum		1.89	-.03		92		
45533722	**Good identification**						**45533722**
Parabacteroides distasonis		100.0	0.35	MAN 1			
45534100	**Good identification**						**45534100**
					COC	GRA	
Fusobacterium mortiferum		96.34	0.35	MLT 11	1	1	
Atopobium parvulum		3.66	0.00	XYL 7	99	99	
45534122	**Good identification**						**45534122**
					CAT		
Parabacteroides distasonis		96.87	0.47	GAL 99	76		
Bacteroides fragilis		2.37	-.03		99		
Prevotella oralis		0.76	-.02		1		
45534132	**Good identification**						**45534132**
Parabacteroides distasonis		100.0	0.42	GAL 99			
45534322	**Good identification**						**45534322**
Parabacteroides distasonis		100.0	0.55	GAL 99			
45534332	**Good identification**						**45534332**
Parabacteroides distasonis		98.53	0.50	GAL 99			
Clostridium clostridioforme		1.47	0.13				
45534522	**Acceptable identification**						**45534522**
					GRA		
Parabacteroides distasonis		93.17	0.43	GAL 99	1		
Bifidobacterium longum subsp. longum		5.11	0.17		99		
Clostridium clostridioforme		1.72	0.08		50		
45534532	**Low discrimination**						**45534532**
					GRA		
Parabacteroides distasonis		75.22	0.39	GAL 99	1		
Bifidobacterium longum subsp. longum		13.05	0.21		99		
Clostridium clostridioforme		11.74	0.19		50		
45534722	**Good identification**						**45534722**
Parabacteroides distasonis		95.15	0.51	GAL 99			
Clostridium clostridioforme		4.29	0.23				
Bacteroides thetaiotaomicron		0.56	-.03				
45534732	**Low discrimination**						**45534732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		72.38	0.47	GAL 99			
Clostridium clostridioforme		27.62	0.34				
45535022	**Acceptable identification**						**45535022**
					CAT		
Parabacteroides distasonis		91.94	0.47	LAC 99	76		
Prevotella oralis		4.55	0.12		1		
Bacteroides fragilis		2.25	-.03		99		
45535032	**Good identification**						**45535032**
Parabacteroides distasonis		100.0	0.42	LAC 99			
45535102	**Low discrimination**						**45535102**
					COC	GRA	
Parabacteroides distasonis		72.61	0.47	NAG 99	1	1	
Bifidobacterium adolescentis		8.96	0.29		1	99	
Atopobium parvulum		8.65	0.21		99	99	
45535112	**Low discrimination**						**45535112**
					GRA		
Parabacteroides distasonis		75.48	0.42	NAG 99	1		
Bifidobacterium adolescentis		21.14	0.30		99		
Bifidobacterium longum subsp. longum		2.01	0.11		99		
45535120	**Low discrimination**						**45535120**
					GRA		
Parabacteroides distasonis		53.55	0.47	BGL 99	1		
Bacteroides vulgatus		11.85	0.17		1		
Clostridium perfringens		11.49	0.26		99		
45535122	**Very good identification**						**45535122**
					CAT		
Parabacteroides distasonis		96.61	0.81		76		
Bacteroides fragilis		2.36	0.32		99		
Prevotella oralis		0.76	0.32		1		
45535123	**Good identification**						**45535123**
					CAT		
Parabacteroides distasonis		96.87	0.47	URE 1	76		
Bacteroides fragilis		2.37	-.03		99		
Prevotella oralis		0.76	-.02		1		
45535130	**Acceptable identification**						**45535130**
					GRA		
Parabacteroides distasonis		90.39	0.42	BGL 99	1		
Bifidobacterium longum subsp. longum		9.61	0.21		99		
45535132	**Very good identification**						**45535132**
					GRA	CAT	
Parabacteroides distasonis		99.79	0.76		1	76	
Bifidobacterium longum subsp. longum		0.11	0.21		99	1	
Bacteroides fragilis		0.05	-.03		1	99	
45535133	**Good identification**						**45535133**
Parabacteroides distasonis		100.0	0.42	URE 1			
45535162	**Acceptable identification**						**45535162**
					GRA	CAT	
Parabacteroides distasonis		93.94	0.47	SOR 1	1	76	
Bacteroides fragilis		2.30	-.03		1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		1.81	0.08		99	1	
45535172	**Good identification**						**45535172**
Parabacteroides distasonis		100.0	0.42	SOR 1			
45535222	**Good identification**						**45535222**
Parabacteroides distasonis		99.30	0.55	LAC 99			
Prevotella oralis		0.70	0.05				
45535232	**Good identification**						**45535232**
Parabacteroides distasonis		98.68	0.50	LAC 99			
Clostridium clostridioforme		1.32	0.13				
45535302	**Good identification**						**45535302**
Parabacteroides distasonis		98.90	0.55	NAG 99			
Prevotella oralis		0.58	0.04				
Clostridium clostridioforme		0.52	0.10				
45535312	**Good identification**						**45535312**
Parabacteroides distasonis		95.71	0.50	NAG 99			
Clostridium clostridioforme		4.29	0.22				
45535320	**Low discrimination**						**45535320**
Parabacteroides distasonis		59.88	0.55	BGL 99			
Bacteroides vulgatus		39.77	0.34				
Prevotella oralis		0.35	0.04				
45535322	**Very good identification**						**45535322**
Parabacteroides distasonis		99.86	0.89				
Prevotella oralis		0.11	0.26				
Bacteroides vulgatus		0.01	-.01				
45535323	**Good identification**						**45535323**
Parabacteroides distasonis		100.0	0.55	URE 1			
45535330	**Good identification**						**45535330**
Parabacteroides distasonis		98.77	0.50	BGL 99			
Bacteroides vulgatus		1.23	-.01				
45535332	**Very good identification**						**45535332**
Parabacteroides distasonis		99.96	0.85				
Clostridium clostridioforme		0.05	0.22				
45535333	**Good identification**						**45535333**
Parabacteroides distasonis		100.0	0.50	URE 1			
45535362	**Good identification**						**45535362**
Parabacteroides distasonis		100.0	0.55	SOR 1			
45535372	**Good identification**						**45535372**
Parabacteroides distasonis		100.0	0.50	SOR 1			
45535422	**Acceptable identification**						**45535422**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		93.34	0.43	LAC 99	1		
Bifidobacterium longum subsp. longum		5.12	0.17		99		
Clostridium clostridioforme		1.54	0.07		50		
45535432	**Low discrimination**						**45535432**
					GRA		
Parabacteroides distasonis		76.14	0.39	LAC 99	1		
Bifidobacterium longum subsp. longum		13.21	0.21		99		
Clostridium clostridioforme		10.65	0.18		50		
45535500	**Acceptable identification**						**45535500**
					GRA	SPO	
Bifidobacterium longum subsp. longum		92.41	0.41	SUC 99	99	1	
Bacteroides vulgatus		3.30	0.02		1	1	
Clostridium butyricum		1.50	-.03		99	99	
45535502	**Low discrimination**						**45535502**
					GRA		
Bifidobacterium longum subsp. longum		35.87	0.41	SUC 99	99		
Bifidobacterium adolescentis		35.60	0.43		99		
Parabacteroides distasonis		26.42	0.43		1		
45535510	**Good identification**						**45535510**
Bifidobacterium longum subsp. longum		99.29	0.45	SUC 99			
Bifidobacterium adolescentis		0.71	0.10				
45535512	**Low discrimination**						**45535512**
					GRA		
Bifidobacterium longum subsp. longum		48.68	0.45	SUC 99	99		
Bifidobacterium adolescentis		34.67	0.44		99		
Parabacteroides distasonis		11.34	0.39		1		
45535520	**Low discrimination**						**45535520**
					GRA		
Bifidobacterium longum subsp. longum		48.36	0.52	SUC 99	99		
Bacteroides vulgatus		42.73	0.36		1		
Parabacteroides distasonis		8.91	0.43		1		
45535522	**Good identification**						**45535522**
					GRA	CAT	
Parabacteroides distasonis		94.50	0.77		1	76	
Bifidobacterium longum subsp. longum		5.18	0.52		99	1	
Bacteroides thetaiotaomicron		0.13	0.12		1	89	
45535523	**Acceptable identification**						**45535523**
					GRA		
Parabacteroides distasonis		94.80	0.43	URE 1	1		
Bifidobacterium longum subsp. longum		5.20	0.17		99		
45535530	**Acceptable identification**						**45535530**
					GRA		
Bifidobacterium longum subsp. longum		94.03	0.56	SUC 99	99		
Parabacteroides distasonis		5.48	0.39		1		
Bacteroides vulgatus		0.49	0.02		1		
45535532	**Low discrimination**						**45535532**
					GRA		
Parabacteroides distasonis		84.79	0.73		1		
Bifidobacterium longum subsp. longum		14.71	0.56		99		
Clostridium clostridioforme		0.40	0.27		50		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
45535533	**Low discrimination**					**45535533**
					GRA	
Parabacteroides distasonis		85.22	0.39	URE 1	1	
Bifidobacterium longum subsp. longum		14.78	0.21		99	
45535562	**Acceptable identification**					**45535562**
					GRA	
Parabacteroides distasonis		94.80	0.43	SOR 1	1	
Bifidobacterium longum subsp. longum		5.20	0.17		99	
45535572	**Low discrimination**					**45535572**
					GRA	
Parabacteroides distasonis		85.22	0.39	SOR 1	1	
Bifidobacterium longum subsp. longum		14.78	0.21		99	
45535622	**Good identification**					**45535622**
Parabacteroides distasonis		95.02	0.51	LAC 99		
Clostridium clostridioforme		3.84	0.22			
Bacteroides thetaiotaomicron		1.14	0.02			
45535632	**Low discrimination**					**45535632**
Parabacteroides distasonis		74.05	0.47	LAC 99		
Clostridium clostridioforme		25.32	0.33			
Bacteroides thetaiotaomicron		0.64	-.05			
45535702	**Low discrimination**					**45535702**
Parabacteroides distasonis		86.93	0.51	NAG 99		
Clostridium clostridioforme		11.76	0.31			
Bacteroides thetaiotaomicron		0.51	-.03			
45535712	**Low discrimination**					**45535712**
					GRA	
Clostridium clostridioforme		52.76	0.42	SUC 99	50	
Parabacteroides distasonis		46.10	0.47	SAL 88	1	
Bifidobacterium longum subsp. longum		0.67	0.11		99	
45535720	**Acceptable identification**					**45535720**
					GRA	
Bacteroides vulgatus		93.36	0.53	SUC 96	1	
Parabacteroides distasonis		6.49	0.51	TRE 4	1	
Bifidobacterium longum subsp. longum		0.12	0.17		99	
45535722	**Very good identification**					**45535722**
					CAT	
Parabacteroides distasonis		99.11	0.86		76	
Bacteroides thetaiotaomicron		0.59	0.31		89	
Bacteroides vulgatus		0.15	0.19		3	
45535723	**Good identification**					**45535723**
Parabacteroides distasonis		99.41	0.51	URE 1		
Bacteroides thetaiotaomicron		0.59	-.03			
45535730	**Low discrimination**					**45535730**
					GRA	
Parabacteroides distasonis		74.66	0.47	BGL 99	1	
Bacteroides vulgatus		20.16	0.19		1	
Bifidobacterium longum subsp. longum		4.32	0.21		99	
45535732	**Very good identification**					**45535732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		98.39	0.81			
Clostridium clostridioforme		1.14	0.42			
Bacteroides thetaiotaomicron		0.42	0.24			
45535733	**Good identification**					**45535733**
Parabacteroides distasonis		98.86	0.47	URE 1		
Clostridium clostridioforme		1.14	0.08			
45535762	**Good identification**					**45535762**
Parabacteroides distasonis		99.41	0.51	SOR 1		
Bacteroides thetaiotaomicron		0.59	-.03			
45535772	**Good identification**					**45535772**
Parabacteroides distasonis		98.86	0.47	SOR 1		
Clostridium clostridioforme		1.14	0.08			
45537122	**Good identification**					**45537122**
					CAT	
Parabacteroides distasonis		96.26	0.47	MAN 1	76	
Bacteroides fragilis		2.35	-.03		99	
Prevotella oralis		0.75	-.02		1	
45537132	**Good identification**					**45537132**
Parabacteroides distasonis		100.0	0.42	MAN 1		
45537322	**Good identification**					**45537322**
Parabacteroides distasonis		100.0	0.55	MAN 1		
45537332	**Good identification**					**45537332**
Parabacteroides distasonis		100.0	0.50	MAN 1		
45537502	**Low discrimination**					**45537502**
Bifidobacterium adolescentis		87.89	0.41	SUC 99		
Actinomyces israelii		11.02	0.32			
Bifidobacterium longum subsp. longum		1.09	0.07			
45537512	**Good identification**					**45537512**
Bifidobacterium adolescentis		95.52	0.43	SUC 99		
Bifidobacterium longum subsp. longum		1.66	0.11			
Actinomyces israelii		1.46	0.18			
45537522	**Acceptable identification**					**45537522**
					GRA	
Parabacteroides distasonis		93.81	0.43	MAN 1	1	
Bifidobacterium longum subsp. longum		5.15	0.17		99	
Bifidobacterium adolescentis		1.05	0.07		99	
45537532	**Low discrimination**					**45537532**
					GRA	
Parabacteroides distasonis		79.22	0.39	MAN 1	1	
Bifidobacterium longum subsp. longum		13.74	0.21		99	
Bifidobacterium dentium		5.04	-.03		99	
45537722	**Good identification**					**45537722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		99.41	0.51	MAN 1		
Bacteroides thetaiotaomicron		0.59	-.03			
45537732	**Good identification**					**45537732**
Parabacteroides distasonis		98.86	0.47	MAN 1		
Clostridium clostridioforme		1.14	0.08			
45541122	**Acceptable identification**					**45541122**
					CAT	
Prevotella oralis		91.31	0.43	SUC 92	1	
Prevotella melaninogenica		4.69	0.18	FRU 99	1	
Capnocytophaga ochracea		3.03	0.18		90	
45541322	**Good identification**					**45541322**
Prevotella oralis		97.45	0.37	SUC 92		
Parabacteroides distasonis		2.55	0.23	FRU 99		
45545322	**Low discrimination**					**45545322**
					CAT	
Parabacteroides distasonis		61.49	0.39	FRU 99	76	
Prevotella oralis		32.29	0.22		1	
Bacteroides eggerthii		4.40	-.03		93	
45550000	**Acceptable identification**					**45550000**
Fusobacterium mortiferum		90.41	0.49	MLT 11		
Prevotella intermedia		8.48	0.20			
Prevotella disiens		1.11	-.03			
45550100	**Good identification**					**45550100**
Fusobacterium mortiferum		99.64	0.56	MLT 11		
Prevotella oralis		0.36	0.00			
45550102	**Low discrimination**					**45550102**
					CAT	
Fusobacterium mortiferum		59.93	0.39	MLT 11	5	
Prevotella oralis		37.45	0.22	BGL 10	1	
Capnocytophaga ochracea		2.62	0.02		90	
45550122	**Good identification**					**45550122**
					CAT	
Prevotella oralis		95.74	0.43	SUC 92	1	
Capnocytophaga ochracea		3.17	0.18	GAL 99	90	
Prevotella melaninogenica		0.45	0.00		1	
45550322	**Good identification**					**45550322**
Prevotella oralis		97.45	0.37	SUC 92		
Parabacteroides distasonis		2.55	0.23	GAL 99		
45551000	**Low discrimination**					**45551000**
					GRA	
Fusobacterium mortiferum		87.32	0.52	MLT 11	1	
Prevotella intermedia		5.46	0.20		1	
Lactobacillus jensenii		4.09	0.19		99	
45551002	**Low discrimination**					**45551002**
					GRA	
Prevotella oralis		82.00	0.35	SUC 92	1	
Fusobacterium mortiferum		13.35	0.36	LAC 94	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		3.75	0.16	NAG 95	99		
45551020	**Low discrimination**						**45551020**
					GRA	SPO	
Prevotella oralis		79.19	0.35	SUC 92	1	1	
Clostridium septicum		6.42	0.13	LAC 94	99	99	
Prevotella melaninogenica		5.84	0.12	BGL 95	1	1	
45551022	**Very good identification**						**45551022**
					GRA	CAT	
Prevotella oralis		99.11	0.57	SUC 92	1	1	
Capnocytophaga ochracea		0.52	0.18	LAC 94	1	90	
Clostridium perfringens		0.14	0.12		99	1	
45551100	**Low discrimination**						**45551100**
					CAT		
Fusobacterium mortiferum		74.23	0.59	MLT 11	5		
Prevotella oralis		17.90	0.34		1		
Capnocytophaga ochracea		2.64	0.20		90		
45551102	**Acceptable identification**						**45551102**
					CAT		
Prevotella oralis		90.34	0.56	SUC 92	1		
Capnocytophaga ochracea		6.32	0.36	NAG 95	90		
Fusobacterium mortiferum		2.19	0.42		5		
45551120	**Low discrimination**						**45551120**
					GRA	SPO	
Prevotella oralis		46.70	0.56	SUC 92	1	1	
Clostridium septicum		23.93	0.47	BGL 95	99	99	
Prevotella melaninogenica		21.75	0.47		1	1	
45551122	**Very good identification**						**45551122**
					CAT		
Prevotella oralis		95.45	0.78	SUC 92	1		
Capnocytophaga ochracea		3.16	0.53		90		
Prevotella melaninogenica		0.45	0.34		1		
45551123	**Good identification**						**45551123**
					GRA	CAT	
Prevotella oralis		95.14	0.43	SUC 92	1	1	
Capnocytophaga ochracea		3.15	0.18	URE 1	1	90	
Actinomyces naeslundii		0.63	0.32		99	25	
45551132	**Low discrimination**						**45551132**
					CAT		
Prevotella oralis		61.83	0.43	SUC 92	1		
Capnocytophaga ochracea		22.53	0.36	MLZ 1	90		
Parabacteroides distasonis		12.31	0.44		76		
45551162	**Low discrimination**						**45551162**
					GRA	CAT	
Prevotella oralis		69.25	0.43	SUC 92	1	1	
Capnocytophaga ochracea		25.23	0.36	SOR 1	1	90	
Bifidobacterium breve		3.20	0.32		99	1	
45551222	**Good identification**						**45551222**
Prevotella oralis		99.59	0.51	SUC 92			
Parabacteroides distasonis		0.41	0.23	LAC 94			
45551302	**Good identification**						**45551302**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		99.34	0.49	SUC 92	1		
Parabacteroides distasonis		0.50	0.23	NAG 95	76		
Capnocytophaga ochracea		0.16	0.02		90		
45551320	**Good identification**						**45551320**
					GRA	SPO	
Prevotella oralis		96.37	0.49	SUC 92	1	1	
Clostridium septicum		1.16	0.13	BGL 95	99	99	
Prevotella melaninogenica		1.06	0.12		1	1	
45551322	**Very good identification**						**45551322**
					CAT		
Prevotella oralis		97.36	0.71	SUC 92	1		
Parabacteroides distasonis		2.55	0.57		76		
Capnocytophaga ochracea		0.08	0.18		90		
45551323	**Good identification**						**45551323**
Prevotella oralis		97.45	0.37	SUC 92			
Parabacteroides distasonis		2.55	0.23	URE 1			
45551332	**Low discrimination**						**45551332**
					CAT		
Parabacteroides distasonis		58.02	0.52	XYL 90	76		
Prevotella oralis		41.62	0.37		1		
Capnocytophaga ochracea		0.36	0.02		90		
45551362	**Good identification**						**45551362**
					CAT		
Prevotella oralis		96.64	0.37	SUC 92	1		
Parabacteroides distasonis		2.53	0.23	SOR 1	76		
Capnocytophaga ochracea		0.83	0.02		90		
45551502	**Low discrimination**						**45551502**
					GRA	CAT	
Bifidobacterium adolescentis		47.75	0.37	SUC 99	99	1	
Capnocytophaga ochracea		27.45	0.23	XYL 87	1	90	
Prevotella oralis		22.47	0.22		1	1	
45551512	**Acceptable identification**						**45551512**
					GRA	CAT	
Bifidobacterium adolescentis		90.38	0.39	SUC 99	99	1	
Capnocytophaga ochracea		4.72	0.07	XYL 87	1	90	
Bifidobacterium longum subsp. longum		3.59	0.13		99	1	
45551522	**Low discrimination**						**45551522**
					CAT		
Prevotella oralis		54.77	0.43	SUC 92	1		
Capnocytophaga ochracea		31.70	0.40	ARA 1	90		
Parabacteroides distasonis		12.42	0.45		76		
45551532	**Low discrimination**						**45551532**
					GRA	CAT	
Parabacteroides distasonis		55.44	0.41	XYL 90	1	76	
Capnocytophaga ochracea		29.21	0.23		1	90	
Bifidobacterium longum subsp. longum		9.86	0.24		99	1	
45551722	**Low discrimination**						**45551722**
					CAT		
Parabacteroides distasonis		60.63	0.53	XYL 90	76		
Prevotella oralis		38.21	0.37		1		
Capnocytophaga ochracea		0.52	0.06		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45551732	**Good identification**						**45551732**
Parabacteroides distasonis		95.10	0.49	XYL 90			
Clostridium clostridioforme		3.78	0.19				
Prevotella oralis		1.12	0.03				
45553101	**Good identification**						**45553101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
45553103	**Good identification**						**45553103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
45553121	**Good identification**						**45553121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
45553122	**Acceptable identification**						**45553122**
					GRA	CAT	
Prevotella oralis		90.21	0.43	SUC 92	1	1	
Bifidobacterium breve		4.16	0.32	MAN 1	99	1	
Capnocytophaga ochracea		2.99	0.18		1	90	
45553123	**Low discrimination**						**45553123**
					GRA		
Actinomyces naeslundii		55.86	0.37	ESL 12	99		
Prevotella oralis		44.14	0.09	SUC 99	1		
45553322	**Acceptable identification**						**45553322**
					SPO		
Prevotella oralis		94.76	0.37	SUC 92	1		
Clostridium ramosum		2.76	0.11	MAN 1	92		
Parabacteroides distasonis		2.48	0.23		1		
45553502	**Acceptable identification**						**45553502**
Bifidobacterium adolescentis		92.52	0.36	SUC 99			
Actinomyces israelii		7.48	0.23	XYL 87			
45553512	**Good identification**						**45553512**
Bifidobacterium adolescentis		100.0	0.37	SUC 99			
45554100	**Good identification**						**45554100**
Fusobacterium mortiferum		100.0	0.36	MLT 11			
45554322	**Low discrimination**						**45554322**
					CAT		
Parabacteroides distasonis		64.32	0.39	GAL 99	76		
Prevotella oralis		33.78	0.22		1		
Bacteroides fragilis		1.90	-.09		99		
45554722	**Low discrimination**						**45554722**
Parabacteroides distasonis		85.17	0.35	GAL 99			
Clostridium clostridioforme		14.83	0.17				
45555022	**Low discrimination**						**45555022**
					CAT		
Prevotella oralis		69.85	0.42	SUC 92	1		
Bacteroides fragilis		26.40	0.25	XYL 11	99		
Parabacteroides distasonis		3.01	0.31	LAC 94	76		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45555100	**Low discrimination**						**45555100**
					CAT		
Fusobacterium mortiferum		51.99	0.39	MLT 11	5		
Prevotella oralis		20.59	0.18	XYL 7	1		
Bacteroides fragilis		9.44	0.04		99		
45555102	**Low discrimination**						**45555102**
					CAT	GRA	
Prevotella oralis		53.50	0.40	SUC 92	1	1	
Bacteroides fragilis		24.53	0.25	XYL 11	99	1	
Bifidobacterium adolescentis		9.29	0.37	NAG 95	1	99	
45555112	**Low discrimination**						**45555112**
					GRA	CAT	
Bifidobacterium adolescentis		77.17	0.39	SUC 99	99	1	
Parabacteroides distasonis		10.22	0.26	ARA 87	1	76	
Capnocytophaga ochracea		5.71	0.10		1	90	
45555120	**Low discrimination**						**45555120**
					CAT		
Bacteroides fragilis		63.02	0.38	SUC 99	99		
Prevotella oralis		26.38	0.40	BGL 95	1		
Capnocytophaga ochracea		3.73	0.26		90		
45555122	**Low discrimination**						**45555122**
					CAT		
Bacteroides fragilis		64.06	0.60	SUC 99	99		
Prevotella oralis		26.82	0.62		1		
Parabacteroides distasonis		7.30	0.65		76		
45555132	**Low discrimination**						**45555132**
					CAT		
Parabacteroides distasonis		77.85	0.61		76		
Bacteroides fragilis		12.82	0.25		99		
Prevotella oralis		5.37	0.28		1		
45555222	**Low discrimination**						**45555222**
					CAT		
Prevotella oralis		76.32	0.35	SUC 92	1		
Parabacteroides distasonis		23.00	0.39	XYL 11	76		
Bacteroides fragilis		0.68	-.09	LAC 94	99		
45555320	**Low discrimination**						**45555320**
Bacteroides vulgatus		64.59	0.35	SUC 96			
Prevotella oralis		24.90	0.34	CEL 5			
Parabacteroides distasonis		9.10	0.39	ARA 93			
45555322	**Low discrimination**						**45555322**
					CAT		
Parabacteroides distasonis		64.22	0.73		76		
Prevotella oralis		33.72	0.56		1		
Bacteroides fragilis		1.90	0.25		99		
45555323	**Low discrimination**						**45555323**
					CAT		
Parabacteroides distasonis		64.32	0.39	URE 1	76		
Prevotella oralis		33.78	0.22		1		
Bacteroides fragilis		1.90	-.09		99		
45555332	**Very good identification**						**45555332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		98.80	0.69				
Prevotella oralis		0.97	0.22				
Clostridium clostridioforme		0.17	0.16				
45555362	**Low discrimination**						**45555362**
					CAT		
Parabacteroides distasonis		64.32	0.39	SOR 1	76		
Prevotella oralis		33.78	0.22		1		
Bacteroides fragilis		1.90	-.09		99		
45555502	**Low discrimination**						**45555502**
					GRA	CAT	
Bifidobacterium adolescentis		89.85	0.52	SUC 99	99	1	
Parabacteroides distasonis		2.47	0.27		1	76	
Capnocytophaga ochracea		1.93	0.13		1	90	
45555512	**Acceptable identification**						**45555512**
Bifidobacterium adolescentis		94.85	0.53	SUC 99			
Clostridium clostridioforme		2.08	0.21				
Bifidobacterium longum subsp. longum		1.69	0.21				
45555520	**Acceptable identification**						**45555520**
					GRA		
Bacteroides vulgatus		91.43	0.38	SUC 96	1		
Bifidobacterium longum subsp. longum		3.31	0.28	CEL 5	99		
Parabacteroides distasonis		1.78	0.27	RHA 90	1		
45555522	**Low discrimination**						**45555522**
Parabacteroides distasonis		74.79	0.61				
Bacteroides fragilis		10.82	0.25				
Capnocytophaga ochracea		5.30	0.29				
45555532	**Good identification**						**45555532**
					GRA		
Parabacteroides distasonis		90.64	0.57		1		
Bifidobacterium longum subsp. longum		5.37	0.32		99		
Clostridium clostridioforme		1.66	0.21		50		
45555542	**Acceptable identification**						**45555542**
Bifidobacterium adolescentis		94.07	0.36	SUC 99			
Actinomyces israelii		5.94	0.22	SOR 11			
45555552	**Good identification**						**45555552**
Bifidobacterium adolescentis		100.0	0.37	SUC 99			
45555622	**Low discrimination**						**45555622**
Parabacteroides distasonis		82.59	0.35	LAC 99			
Clostridium clostridioforme		12.89	0.16				
Prevotella oralis		4.52	0.01				
45555702	**Low discrimination**						**45555702**
					GRA		
Parabacteroides distasonis		59.17	0.35	NAG 99	1		
Clostridium clostridioforme		30.92	0.25		50		
Bifidobacterium adolescentis		7.24	0.17		99		
45555712	**Low discrimination**						**45555712**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		77.58	0.36	SUC 99	50	
Parabacteroides distasonis		17.55	0.31	SAL 88	1	
Bifidobacterium adolescentis		4.87	0.19	TRE 88	99	
45555720	**Good identification**					**45555720**
Bacteroides vulgatus		99.33	0.55	SUC 96		
Parabacteroides distasonis		0.65	0.35	CEL 5		
Prevotella oralis		0.03	0.00			
45555722	**Good identification**					**45555722**
Parabacteroides distasonis		94.80	0.70			
Bacteroides vulgatus		1.49	0.20			
Bacteroides uniformis		1.27	0.19			
45555723	**Good identification**					**45555723**
Parabacteroides distasonis		100.0	0.35	URE 1		
45555732	**Good identification**					**45555732**
Parabacteroides distasonis		94.96	0.65			
Clostridium clostridioforme		4.24	0.36			
Bacteroides thetaiotaomicron		0.55	0.11			
45555762	**Good identification**					**45555762**
Parabacteroides distasonis		100.0	0.35	SOR 1		
45557100	**Good identification**					**45557100**
Actinomyces israelii		100.0	0.40	SUC 98		
45557102	**Low discrimination**					**45557102**
					GRA	
Actinomyces israelii		50.72	0.43	SUC 98	99	
Bifidobacterium adolescentis		44.65	0.36	SAL 92	99	
Prevotella oralis		3.17	0.06		1	
45557112	**Low discrimination**					**45557112**
Bifidobacterium adolescentis		87.81	0.37	SUC 99		
Actinomyces israelii		12.20	0.29	ARA 87		
45557142	**Low discrimination**					**45557142**
Actinomyces israelii		78.14	0.36	SUC 98		
Bifidobacterium adolescentis		21.86	0.20	SAL 92		
45557322	**Low discrimination**					**45557322**
					CAT	
Parabacteroides distasonis		64.32	0.39	MAN 1	76	
Prevotella oralis		33.78	0.22		1	
Bacteroides fragilis		1.90	-.09		99	
45557500	**Low discrimination**					**45557500**
					SPO	
Actinomyces israelii		86.45	0.39	SUC 98	1	
Bifidobacterium adolescentis		9.43	0.16	SAL 92	1	
Clostridium butyricum		4.12	-.05		99	
45557502	**Low discrimination**					**45557502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		87.81	0.50	SUC 99			
Actinomyces israelii		12.20	0.42				
45557512	**Good identification**						**45557512**
Bifidobacterium adolescentis		97.90	0.52	SUC 99			
Actinomyces israelii		1.66	0.27				
Bifidobacterium dentium		0.44	-.07				
45557552	**Acceptable identification**						**45557552**
Bifidobacterium adolescentis		94.93	0.36	SUC 99			
Actinomyces israelii		5.07	0.20	SOR 11			
45557722	**Acceptable identification**						**45557722**
Parabacteroides distasonis		93.37	0.35	MAN 1			
Bacteroides ovatus		6.64	0.00				
45565122	**Acceptable identification**						**45565122**
					CAT		
Parabacteroides distasonis		93.37	0.42	FRU 99	76		
Prevotella oralis		4.81	0.07		1		
Bacteroides fragilis		1.82	-.09		99		
45565132	**Good identification**						**45565132**
Parabacteroides distasonis		100.0	0.38	FRU 99			
45565322	**Good identification**						**45565322**
Parabacteroides distasonis		99.27	0.50	FRU 99			
Prevotella oralis		0.73	0.01				
45565332	**Good identification**						**45565332**
Parabacteroides distasonis		100.0	0.46	FRU 99			
45565522	**Good identification**						**45565522**
Parabacteroides distasonis		100.0	0.39	FRU 99			
45565722	**Acceptable identification**						**45565722**
Parabacteroides distasonis		94.56	0.47	FRU 99			
Bacteroides eggerthii		2.43	-.03				
Bacteroides thetaiotaomicron		2.21	0.03				
45565732	**Acceptable identification**						**45565732**
Parabacteroides distasonis		91.85	0.42	FRU 99			
Clostridium clostridioforme		6.60	0.17				
Bacteroides thetaiotaomicron		1.55	-.04				
45570000	**Good identification**						**45570000**
Fusobacterium mortiferum		100.0	0.40	MLT 11			
45570100	**Good identification**						**45570100**
					COC	GRA	
Fusobacterium mortiferum		98.94	0.47	MLT 11	1	1	
Atopobium parvulum		1.07	0.01		99	99	
45571000	**Low discrimination**						**45571000**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		66.34	0.43	MLT 11	1			
Lactobacillus jensenii		31.18	0.28		99			
Actinomyces naeslundii		1.37	0.26		99			
45571022	**Low discrimination**							**45571022**
					GRA			
Prevotella oralis		89.45	0.37	SUC 92	1			
Parabacteroides distasonis		3.77	0.26	TRE 6	1			
Clostridium perfringens		3.58	0.16	LAC 94	99			
45571100	**Low discrimination**							**45571100**
					COC	GRA	SPO	
Fusobacterium mortiferum		48.54	0.50	MLT 11	1	1	1	
Atopobium parvulum		34.49	0.36		99	99	1	
Clostridium septicum		11.57	0.21		1	99	99	
45571101	**Low discrimination**							**45571101**
					COC	GRA		
Actinomyces naeslundii		68.55	0.37	ESL 12	1	99		
Fusobacterium mortiferum		18.39	0.15	SUC 99	1	1		
Atopobium parvulum		13.07	0.01		99	99		
45571102	**Low discrimination**							**45571102**
					COC	GRA		
Prevotella oralis		66.96	0.35	SUC 92	1	1		
Atopobium parvulum		8.57	0.23	TRE 6	99	99		
Fusobacterium mortiferum		7.60	0.33	NAG 95	1	1		
45571103	**Low discrimination**							**45571103**
					GRA			
Actinomyces naeslundii		79.19	0.37	ESL 12	99			
Prevotella oralis		20.81	0.01	SUC 99	1			
45571120	**Acceptable identification**							**45571120**
					GRA	SPO		
Clostridium septicum		91.17	0.56	RAF 1	99	99		
Prevotella oralis		3.79	0.35		1	1		
Clostridium perfringens		2.87	0.37		99	84		
45571122	**Low discrimination**							**45571122**
					GRA	SPO		
Prevotella oralis		74.79	0.57	SUC 92	1	1		
Parabacteroides distasonis		19.94	0.60	TRE 6	1	1		
Clostridium septicum		1.94	0.27		99	99		
45571132	**Low discrimination**							**45571132**
					GRA			
Parabacteroides distasonis		88.12	0.55	XYL 90	1			
Prevotella oralis		6.20	0.23		1			
Bifidobacterium breve		2.99	0.29		99			
45571322	**Low discrimination**							**45571322**
					SPO			
Parabacteroides distasonis		64.83	0.68	XYL 90	1			
Prevotella oralis		34.73	0.51		1			
Clostridium ramosum		0.44	0.19		92			
45571332	**Very good identification**							**45571332**
Parabacteroides distasonis		98.76	0.64	XYL 90				
Prevotella oralis		0.99	0.17					
Clostridium clostridioforme		0.25	0.13					

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
45571512	**Low discrimination**					**45571512**
					GRA	
Bifidobacterium adolescentis		80.30	0.36	SUC 99	99	
Clostridium clostridioforme		12.82	0.19	XYL 87	50	
Parabacteroides distasonis		4.95	0.18		1	
45571522	**Good identification**					**45571522**
					CAT	
Parabacteroides distasonis		92.97	0.56	XYL 90	76	
Prevotella oralis		5.75	0.23		1	
Capnocytophaga ochracea		0.53	0.06		90	
45571532	**Good identification**					**45571532**
					GRA	
Parabacteroides distasonis		96.53	0.52	XYL 90	1	
Clostridium clostridioforme		2.53	0.19		50	
Bifidobacterium longum subsp. longum		0.94	0.13		99	
45571722	**Very good identification**					**45571722**
					CAT	
Parabacteroides distasonis		97.55	0.65	XYL 90	76	
Prevotella oralis		0.86	0.17		1	
Bacteroides thetaiotaomicron		0.85	0.13		89	
45571732	**Good identification**					**45571732**
Parabacteroides distasonis		93.43	0.60	XYL 90		
Clostridium clostridioforme		5.98	0.34			
Bacteroides thetaiotaomicron		0.59	0.06			
45573101	**Good identification**					**45573101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
45573103	**Good identification**					**45573103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
45573121	**Good identification**					**45573121**
Actinomyces naeslundii		100.0	0.42	ESL 12		
45573123	**Good identification**					**45573123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
45574122	**Acceptable identification**					**45574122**
					CAT	
Parabacteroides distasonis		93.37	0.42	GAL 99	76	
Prevotella oralis		4.81	0.07		1	
Bacteroides fragilis		1.82	-.09		99	
45574132	**Good identification**					**45574132**
Parabacteroides distasonis		96.34	0.38	GAL 99		
Clostridium clostridioforme		3.66	0.08			
45574322	**Good identification**					**45574322**
Parabacteroides distasonis		98.20	0.50	GAL 99		
Clostridium clostridioforme		1.08	0.11			
Prevotella oralis		0.72	0.01			
45574332	**Acceptable identification**					**45574332**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		91.51	0.46	GAL 99				
Clostridium clostridioforme		8.49	0.22					
45574522	**Low discrimination**							**45574522**
Parabacteroides distasonis		89.71	0.39	GAL 99				
Clostridium clostridioforme		10.29	0.17					
45574712	**Good identification**							**45574712**
Clostridium clostridioforme		100.0	0.43	SUC 99				
45574722	**Low discrimination**							**45574722**
Parabacteroides distasonis		76.70	0.47	GAL 99				
Clostridium clostridioforme		21.51	0.32					
Bacteroides thetaiotaomicron		1.79	0.03					
45574732	**Low discrimination**							**45574732**
Clostridium clostridioforme		69.99	0.43	SUC 99				
Parabacteroides distasonis		29.51	0.42	SAL 88				
Bacteroides thetaiotaomicron		0.50	-.04					
45575022	**Low discrimination**							**45575022**
					CAT			
Parabacteroides distasonis		74.35	0.42	LAC 99	76			
Prevotella oralis		24.20	0.21		1			
Bacteroides fragilis		1.45	-.09		99			
45575032	**Good identification**							**45575032**
Parabacteroides distasonis		96.71	0.38	LAC 99				
Clostridium clostridioforme		3.29	0.07					
45575100	**Low discrimination**							**45575100**
					COC	SPO		
Atopobium parvulum		79.42	0.40	SUC 96	99	1		
Actinomyces israelii		8.16	0.37	RAF 1	1	1		
Clostridium butyricum		7.51	0.15		1	99		
45575102	**Low discrimination**							**45575102**
					COC	GRA		
Parabacteroides distasonis		34.71	0.42	NAG 99	1	1		
Atopobium parvulum		17.92	0.27		99	99		
Bifidobacterium adolescentis		16.90	0.34		1	99		
45575112	**Low discrimination**							**45575112**
					GRA			
Bifidobacterium adolescentis		46.62	0.36	SUC 99	99			
Parabacteroides distasonis		42.18	0.38	ARA 87	1			
Actinomyces israelii		5.28	0.26		99			
45575120	**Low discrimination**							**45575120**
					COC	GRA	SPO	
Parabacteroides distasonis		40.07	0.42	BGL 99	1	1	1	
Clostridium septicum		21.17	0.21		1	99	99	
Atopobium parvulum		20.69	0.27		99	99	1	
45575122	**Good identification**							**45575122**
					CAT			
Parabacteroides distasonis		93.21	0.76		76			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		4.80	0.42		1	
Bacteroides fragilis		1.82	0.25		99	
45575123	**Acceptable identification**					**45575123**
					CAT	
Parabacteroides distasonis		93.37	0.42	URE 1	76	
Prevotella oralis		4.81	0.07		1	
Bacteroides fragilis		1.82	-.09		99	
45575130	**Good identification**					**45575130**
Parabacteroides distasonis		100.0	0.38	BGL 99		
45575132	**Very good identification**					**45575132**
Parabacteroides distasonis		99.75	0.72			
Clostridium clostridioforme		0.12	0.16			
Prevotella oralis		0.10	0.07			
45575133	**Good identification**					**45575133**
Parabacteroides distasonis		100.0	0.38	URE 1		
45575162	**Acceptable identification**					**45575162**
					CAT	
Parabacteroides distasonis		93.37	0.42	SOR 1	76	
Prevotella oralis		4.81	0.07		1	
Bacteroides fragilis		1.82	-.09		99	
45575172	**Good identification**					**45575172**
Parabacteroides distasonis		100.0	0.38	SOR 1		
45575222	**Acceptable identification**					**45575222**
Parabacteroides distasonis		94.67	0.50	LAC 99		
Prevotella oralis		4.40	0.15			
Clostridium clostridioforme		0.93	0.10			
45575232	**Acceptable identification**					**45575232**
Parabacteroides distasonis		92.32	0.46	LAC 99		
Clostridium clostridioforme		7.68	0.22			
45575302	**Acceptable identification**					**45575302**
Parabacteroides distasonis		92.37	0.50	NAG 99		
Prevotella oralis		3.54	0.13			
Clostridium clostridioforme		3.04	0.19			
45575312	**Low discrimination**					**45575312**
Parabacteroides distasonis		78.22	0.46	NAG 99		
Clostridium clostridioforme		21.78	0.31			
45575320	**Acceptable identification**					**45575320**
Parabacteroides distasonis		90.64	0.50	BGL 99		
Bacteroides vulgatus		5.88	0.12			
Prevotella oralis		3.48	0.13			
45575322	**Very good identification**					**45575322**
Parabacteroides distasonis		99.21	0.85			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		0.73	0.35				
Clostridium clostridioforme		0.03	0.19				
45575323	**Good identification**						**45575323**
Parabacteroides distasonis		99.27	0.50	URE 1			
Prevotella oralis		0.73	0.01				
45575330	**Good identification**						**45575330**
Parabacteroides distasonis		100.0	0.46	BGL 99			
45575332	**Very good identification**						**45575332**
Parabacteroides distasonis		99.68	0.80				
Clostridium clostridioforme		0.28	0.31				
Prevotella oralis		0.01	0.01				
45575333	**Good identification**						**45575333**
Parabacteroides distasonis		100.0	0.46	URE 1			
45575362	**Good identification**						**45575362**
Parabacteroides distasonis		99.27	0.50	SOR 1			
Prevotella oralis		0.73	0.01				
45575372	**Good identification**						**45575372**
Parabacteroides distasonis		100.0	0.46	SOR 1			
45575422	**Acceptable identification**						**45575422**
Parabacteroides distasonis		90.68	0.39	LAC 99			
Clostridium clostridioforme		9.32	0.16				
45575502	**Low discrimination**						**45575502**
					GRA	SPO	
Bifidobacterium adolescentis		59.96	0.48	SUC 99	99	1	
Clostridium butyricum		17.81	0.25		99	99	
Parabacteroides distasonis		11.28	0.39		1	1	
45575512	**Low discrimination**						**45575512**
					GRA		
Bifidobacterium adolescentis		73.37	0.50	SUC 99	99		
Clostridium clostridioforme		17.70	0.36		50		
Parabacteroides distasonis		6.08	0.34		1		
45575520	**Low discrimination**						**45575520**
					GRA	SPO	
Parabacteroides distasonis		51.29	0.39	BGL 99	1	1	
Bacteroides vulgatus		24.06	0.15		1	1	
Clostridium butyricum		18.11	0.14		99	99	
45575522	**Very good identification**						**45575522**
Parabacteroides distasonis		98.62	0.73				
Bacteroides thetaiotaomicron		0.52	0.18				
Clostridium clostridioforme		0.34	0.25				
45575523	**Good identification**						**45575523**
Parabacteroides distasonis		100.0	0.39	URE 1			
45575532	**Very good identification**						**45575532**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		96.17	0.68			
Clostridium clostridioforme		2.83	0.36			
Bacteroides thetaiotaomicron		0.37	0.11			
45575562	**Good identification**					**45575562**
Parabacteroides distasonis		100.0	0.39	SOR 1		
45575612	**Good identification**					**45575612**
Clostridium clostridioforme		100.0	0.42	SUC 99		
45575622	**Low discrimination**					**45575622**
Parabacteroides distasonis		77.02	0.47	LAC 99		
Clostridium clostridioforme		19.35	0.31			
Bacteroides thetaiotaomicron		3.63	0.08			
45575632	**Low discrimination**					**45575632**
Clostridium clostridioforme		67.27	0.42	SUC 99		
Parabacteroides distasonis		31.65	0.42	SAL 88		
Bacteroides thetaiotaomicron		1.08	0.01			
45575702	**Low discrimination**					**45575702**
Parabacteroides distasonis		52.25	0.47	NAG 99		
Clostridium clostridioforme		43.95	0.40			
Bacteroides thetaiotaomicron		1.22	0.03			
45575712	**Low discrimination**					**45575712**
					GRA	
Clostridium clostridioforme		87.06	0.51	SUC 99	50	
Parabacteroides distasonis		12.23	0.42	SAL 88	1	
Bifidobacterium adolescentis		0.50	0.16		99	
45575722	**Very good identification**					**45575722**
Parabacteroides distasonis		96.44	0.81			
Bacteroides thetaiotaomicron		2.25	0.37			
Clostridium clostridioforme		0.82	0.40			
45575723	**Good identification**					**45575723**
Parabacteroides distasonis		96.91	0.47	URE 1		
Bacteroides thetaiotaomicron		2.26	0.03			
Clostridium clostridioforme		0.82	0.06			
45575730	**Low discrimination**					**45575730**
Parabacteroides distasonis		89.68	0.42	BGL 99		
Clostridium clostridioforme		6.45	0.17			
Bacteroides vulgatus		2.37	-.03			
45575732	**Good identification**					**45575732**
Parabacteroides distasonis		91.35	0.76			
Clostridium clostridioforme		6.57	0.51			
Bacteroides thetaiotaomicron		1.54	0.30			
45575733	**Acceptable identification**					**45575733**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		91.85	0.42	URE 1			
Clostridium clostridioforme		6.60	0.17				
Bacteroides thetaiotaomicron		1.55	-.04				
45575762	**Good identification**						**45575762**
Parabacteroides distasonis		95.10	0.47	SOR 1			
Bacteroides thetaiotaomicron		2.22	0.03				
Bacteroides ovatus		1.88	0.02				
45575772	**Low discrimination**						**45575772**
Parabacteroides distasonis		89.84	0.42	SOR 1			
Clostridium clostridioforme		6.46	0.17				
Bacteroides ovatus		2.19	-.02				
45577000	**Good identification**						**45577000**
Actinomyces israelii		100.0	0.40	SUC 98			
45577002	**Good identification**						**45577002**
Actinomyces israelii		100.0	0.43	SUC 98			
45577042	**Good identification**						**45577042**
Actinomyces israelii		100.0	0.36	SUC 98			
45577100	**Good identification**						**45577100**
					COC	SPO	
Actinomyces israelii		96.58	0.50	SUC 98	1	1	
Clostridium butyricum		1.74	-.01	SAL 92	1	99	
Atopobium parvulum		1.68	0.06		99	1	
45577102	**Low discrimination**						**45577102**
Actinomyces israelii		86.51	0.53	SUC 98			
Bifidobacterium adolescentis		13.50	0.33	SAL 92			
45577110	**Good identification**						**45577110**
Actinomyces israelii		100.0	0.36	SUC 98			
45577122	**Low discrimination**						**45577122**
					GRA		
Parabacteroides distasonis		81.04	0.42	MAN 1	1		
Actinomyces israelii		13.21	0.33		99		
Prevotella oralis		4.17	0.07		1		
45577132	**Low discrimination**						**45577132**
					GRA		
Parabacteroides distasonis		89.48	0.38	MAN 1	1		
Bifidobacterium dentium		6.48	-.03		99		
Actinomyces israelii		4.05	0.18		99		
45577140	**Good identification**						**45577140**
Actinomyces israelii		100.0	0.43	SUC 98			
45577142	**Good identification**						**45577142**
Actinomyces israelii		95.28	0.46	SUC 98			
Bifidobacterium adolescentis		4.72	0.17	SAL 92			
45577322	**Good identification**						**45577322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Parabacteroides distasonis		98.45	0.50	MAN 1	1		
Prevotella oralis		0.72	0.01		1		
Clostridium ramosum		0.42	-.03		92		
45577332	**Good identification**						**45577332**
Parabacteroides distasonis		100.0	0.46	MAN 1			
45577400	**Good identification**						**45577400**
Actinomyces israelii		100.0	0.38	SUC 98			
45577402	**Good identification**						**45577402**
Actinomyces israelii		95.10	0.41	SUC 98			
Bifidobacterium adolescentis		4.90	0.13	SAL 92			
45577500	**Low discrimination**						**45577500**
					SPO		
Actinomyces israelii		82.11	0.48	SUC 98	1		
Clostridium butyricum		16.31	0.15	SAL 92	99		
Bifidobacterium adolescentis		1.59	0.13		1		
45577502	**Low discrimination**						**45577502**
					SPO		
Bifidobacterium adolescentis		54.58	0.47	SUC 99	1		
Actinomyces israelii		42.77	0.51		1		
Clostridium butyricum		2.20	0.08		99		
45577512	**Low discrimination**						**45577512**
Bifidobacterium adolescentis		56.76	0.48	SUC 99			
Bifidobacterium dentium		37.80	0.27				
Actinomyces israelii		5.44	0.37				
45577522	**Low discrimination**						**45577522**
					GRA		
Parabacteroides distasonis		74.07	0.39	MAN 1	1		
Actinomyces israelii		16.12	0.31		99		
Bifidobacterium dentium		4.71	-.03		99		
45577542	**Low discrimination**						**45577542**
					COC		
Actinomyces israelii		66.53	0.44	SUC 98	1		
Bifidobacterium adolescentis		26.98	0.31	SAL 92	1		
Blautia producta		6.49	0.07		99		
45577722	**Low discrimination**						**45577722**
Parabacteroides distasonis		84.06	0.47	MAN 1			
Bacteroides ovatus		13.26	0.17				
Bacteroides thetaiotaomicron		1.96	0.03				
45577732	**Low discrimination**						**45577732**
Parabacteroides distasonis		75.56	0.42	MAN 1			
Bacteroides ovatus		14.76	0.14				
Clostridium clostridioforme		5.43	0.17				
45600000	**Low discrimination**						**45600000**
					GRA	SPO	
Fusobacterium mortiferum		55.55	0.44	FRU 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		36.76	0.27		99	99	
Clostridium botulinum biovar A		4.89	0.13		99	99	
45600100	**Good identification**						**45600100**
Fusobacterium mortiferum		100.0	0.50	FRU 99			
45601000	**Low discrimination**						**45601000**
					GRA		
Fusobacterium mortiferum		69.87	0.47	FRU 99	1		
Eubacterium tortuosum		29.35	0.24		99		
Atopobium minutum		0.78	0.08		99		
45601100	**Good identification**						**45601100**
					COC	GRA	
Fusobacterium mortiferum		99.41	0.53	FRU 99	1	1	
Blautia hansenii		0.59	-.03		99	99	
45601102	**Good identification**						**45601102**
					GRA		
Fusobacterium mortiferum		96.99	0.37	FRU 99	1		
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99		
45610000	**Very good identification**						**45610000**
					GRA	SPO	
Fusobacterium mortiferum		98.51	0.78		1	1	
Clostridium difficile		1.07	0.28		99	99	
Eubacterium limosum		0.35	0.18		99	1	
45610001	**Good identification**						**45610001**
Fusobacterium mortiferum		100.0	0.44	URE 1			
45610002	**Very good identification**						**45610002**
					GRA		
Fusobacterium mortiferum		99.72	0.62	BGL 10	1		
Atopobium minutum		0.28	0.12		99		
45610010	**Low discrimination**						**45610010**
					GRA	SPO	
Clostridium difficile		83.96	0.40	RAF 1	99	99	
Fusobacterium mortiferum		16.04	0.44	MAN 90	1	1	
45610020	**Good identification**						**45610020**
					GRA		
Fusobacterium mortiferum		98.47	0.44	NAG 1	1		
Atopobium minutum		1.53	0.07		99		
45610040	**Acceptable identification**						**45610040**
					GRA	SPO	
Fusobacterium mortiferum		94.66	0.44	SOR 1	1	1	
Clostridium difficile		5.34	0.06		99	99	
45610100	**Very good identification**						**45610100**
Fusobacterium mortiferum		100.0	0.84				
45610101	**Good identification**						**45610101**
Fusobacterium mortiferum		100.0	0.50	URE 1			
45610102	**Very good identification**						**45610102**
					GRA		
Fusobacterium mortiferum		99.95	0.68	BGL 10	1		
Bifidobacterium longum subsp. infantis		0.05	-.07		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45610110	**Good identification**						**45610110**
					GRA	SPO	
Fusobacterium mortiferum		97.78	0.50	MLZ 1	1	1	
Clostridium difficile		2.22	0.06		99	99	
45610120	**Good identification**						**45610120**
					GRA	SPO	
Fusobacterium mortiferum		99.40	0.50	NAG 1	1	1	
Clostridium septicum		0.60	-.06		99	99	
45610140	**Good identification**						**45610140**
Fusobacterium mortiferum		100.0	0.50	SOR 1			
45610200	**Good identification**						**45610200**
Fusobacterium mortiferum		100.0	0.44	RHA 1			
45610300	**Good identification**						**45610300**
Fusobacterium mortiferum		100.0	0.50	RHA 1			
45610400	**Good identification**						**45610400**
Fusobacterium mortiferum		100.0	0.44	ARA 1			
45610500	**Good identification**						**45610500**
Fusobacterium mortiferum		100.0	0.50	ARA 1			
45611000	**Very good identification**						**45611000**
					GRA		
Fusobacterium mortiferum		99.99	0.81		1		
Atopobium minutum		0.01	0.07		99		
45611001	**Good identification**						**45611001**
Fusobacterium mortiferum		100.0	0.47	URE 1			
45611002	**Very good identification**						**45611002**
					GRA		
Fusobacterium mortiferum		99.93	0.65	BGL 10	1		
Bifidobacterium longum subsp. infantis		0.07	-.07		99		
45611010	**Good identification**						**45611010**
					GRA	SPO	
Fusobacterium mortiferum		96.60	0.47	MLZ 1	1	1	
Clostridium difficile		3.41	0.06		99	99	
45611020	**Good identification**						**45611020**
					GRA	SPO	
Fusobacterium mortiferum		99.07	0.47	NAG 1	1	1	
Clostridium septicum		0.93	-.06		99	99	
45611040	**Good identification**						**45611040**
Fusobacterium mortiferum		100.0	0.47	SOR 1			
45611100	**Very good identification**						**45611100**
					GRA	SPO	
Fusobacterium mortiferum		99.96	0.87		1	1	
Bifidobacterium longum subsp. infantis		0.03	0.10		99	1	
Clostridium septicum		0.00	-.06		99	99	
45611101	**Good identification**						**45611101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.53	URE 1			
45611102	**Very good identification**						**45611102**
					GRA		
Fusobacterium mortiferum		96.54	0.71	BGL 10	1		
Bifidobacterium longum subsp. infantis		3.00	0.27		99		
Prevotella oralis		0.43	0.16		1		
45611103	**Good identification**						**45611103**
					GRA		
Fusobacterium mortiferum		96.99	0.37	URE 1	1		
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99		
45611110	**Good identification**						**45611110**
Fusobacterium mortiferum		100.0	0.53	MLZ 1			
45611112	**Good identification**						**45611112**
					GRA		
Fusobacterium mortiferum		96.99	0.37	MLZ 1	1		
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99		
45611120	**Low discrimination**						**45611120**
					GRA	SPO	
Fusobacterium mortiferum		65.12	0.53	NAG 1	1	1	
Clostridium septicum		25.91	0.29		99	99	
Bifidobacterium longum subsp. infantis		3.91	0.15		99	1	
45611122	**Low discrimination**						**45611122**
					GRA		
Prevotella oralis		45.26	0.38	SUC 92	1		
Bifidobacterium longum subsp. infantis		29.63	0.32	MLT 99	99		
Parabacteroides distasonis		18.94	0.45		1		
45611132	**Acceptable identification**						**45611132**
					GRA		
Parabacteroides distasonis		93.10	0.40	XYL 90	1		
Prevotella oralis		4.17	0.04		1		
Bifidobacterium longum subsp. infantis		2.73	-.03		99		
45611140	**Good identification**						**45611140**
Fusobacterium mortiferum		100.0	0.53	SOR 1			
45611142	**Good identification**						**45611142**
					GRA		
Fusobacterium mortiferum		96.99	0.37	SOR 1	1		
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99		
45611200	**Good identification**						**45611200**
Fusobacterium mortiferum		100.0	0.47	RHA 1			
45611300	**Good identification**						**45611300**
Fusobacterium mortiferum		100.0	0.53	RHA 1			
45611302	**Low discrimination**						**45611302**
Fusobacterium mortiferum		75.48	0.37	RHA 1			
Prevotella oralis		14.20	0.10	BGL 10			
Parabacteroides distasonis		7.99	0.19				
45611322	**Low discrimination**						**45611322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		74.26	0.53	XYL 90	1		
Prevotella oralis		25.35	0.32		1		
Bifidobacterium longum subsp. infantis		0.39	-.03		99		
45611332	**Good identification**						**45611332**
Parabacteroides distasonis		99.36	0.48	XYL 90			
Prevotella oralis		0.64	-.02				
45611400	**Good identification**						**45611400**
Fusobacterium mortiferum		100.0	0.47	ARA 1			
45611500	**Good identification**						**45611500**
Fusobacterium mortiferum		100.0	0.53	ARA 1			
45611502	**Acceptable identification**						**45611502**
					GRA		
Fusobacterium mortiferum		92.50	0.37	ARA 1	1		
Bifidobacterium adolescentis		4.64	0.11	BGL 10	99		
Bifidobacterium longum subsp. infantis		2.87	-.07		99		
45611522	**Acceptable identification**						**45611522**
					GRA		
Parabacteroides distasonis		93.88	0.41	XYL 90	1		
Prevotella oralis		3.70	0.04		1		
Bifidobacterium longum subsp. infantis		2.42	-.03		99		
45611532	**Good identification**						**45611532**
Parabacteroides distasonis		100.0	0.36	XYL 90			
45611722	**Good identification**						**45611722**
Parabacteroides distasonis		99.44	0.49	XYL 90			
Prevotella oralis		0.56	-.02				
45611732	**Good identification**						**45611732**
Parabacteroides distasonis		100.0	0.44	XYL 90			
45612000	**Low discrimination**						**45612000**
					GRA	SPO	
Clostridium difficile		75.46	0.45	RAF 1	99	99	
Eubacterium limosum		16.72	0.32		99	1	
Fusobacterium mortiferum		7.82	0.44		1	1	
45612010	**Very good identification**						**45612010**
					SPO		
Clostridium difficile		99.95	0.56	RAF 1	99		
Eubacterium limosum		0.05	-.03		1		
45612100	**Good identification**						**45612100**
					GRA	SPO	
Fusobacterium mortiferum		95.14	0.50	MAN 1	1	1	
Clostridium difficile		3.98	0.11		99	99	
Eubacterium limosum		0.88	-.03		99	1	
45613000	**Low discrimination**						**45613000**
					GRA	SPO	
Fusobacterium mortiferum		86.67	0.47	MAN 1	1	1	
Clostridium innocuum		6.45	0.15		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		5.63	0.11		99	99	
45613100	**Good identification**						**45613100**
Fusobacterium mortiferum		100.0	0.53	MAN 1			
45613102	**Good identification**						**45613102**
					GRA		
Fusobacterium mortiferum		96.99	0.37	MAN 1	1		
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99		
45614000	**Very good identification**						**45614000**
					GRA	SPO	
Fusobacterium mortiferum		99.25	0.59	XYL 7	1	1	
Clostridium difficile		0.75	0.06		99	99	
45614002	**Good identification**						**45614002**
Fusobacterium mortiferum		100.0	0.42	XYL 7			
45614100	**Very good identification**						**45614100**
Fusobacterium mortiferum		100.0	0.65	XYL 7			
45614102	**Good identification**						**45614102**
Fusobacterium mortiferum		100.0	0.49	XYL 7			
45615000	**Very good identification**						**45615000**
Fusobacterium mortiferum		100.0	0.62	XYL 7			
45615002	**Good identification**						**45615002**
Fusobacterium mortiferum		100.0	0.45	XYL 7			
45615100	**Very good identification**						**45615100**
					GRA		
Fusobacterium mortiferum		99.63	0.68	XYL 7	1		
Bifidobacterium longum subsp. infantis		0.37	0.09		99		
45615102	**Low discrimination**						**45615102**
					GRA		
Fusobacterium mortiferum		71.75	0.52	XYL 7	1		
Bifidobacterium longum subsp. infantis		24.20	0.26	BGL 10	99		
Parabacteroides distasonis		3.06	0.27		1		
45615122	**Low discrimination**						**45615122**
					GRA		
Parabacteroides distasonis		84.67	0.61		1		
Bifidobacterium longum subsp. infantis		12.04	0.30		99		
Prevotella oralis		2.78	0.23		1		
45615132	**Very good identification**						**45615132**
					GRA		
Parabacteroides distasonis		99.73	0.56		1		
Bifidobacterium longum subsp. infantis		0.27	-.04		99		
45615322	**Very good identification**						**45615322**
					GRA		
Parabacteroides distasonis		99.45	0.69		1		
Prevotella oralis		0.47	0.16		1		
Bifidobacterium longum subsp. infantis		0.05	-.04		99		
45615332	**Very good identification**						**45615332**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		100.0	0.64					
45615522	**Very good identification**							**45615522**
					GRA			
Parabacteroides distasonis		99.77	0.57		1			
Bifidobacterium longum subsp. infantis		0.23	-.04		99			
45615532	**Good identification**							**45615532**
Parabacteroides distasonis		100.0	0.53					
45615722	**Very good identification**							**45615722**
Parabacteroides distasonis		99.87	0.65					
Bacteroides vulgatus		0.13	-.02					
45615732	**Very good identification**							**45615732**
Parabacteroides distasonis		99.83	0.61					
Clostridium clostridioforme		0.17	0.08					
45620100	**Good identification**							**45620100**
Fusobacterium mortiferum		100.0	0.41	FRU 99				
45621000	**Good identification**							**45621000**
Fusobacterium mortiferum		100.0	0.38	FRU 99				
45621100	**Good identification**							**45621100**
Fusobacterium mortiferum		100.0	0.44	FRU 99				
45625122	**Good identification**							**45625122**
Parabacteroides distasonis		100.0	0.38	FRU 99				
45625322	**Good identification**							**45625322**
Parabacteroides distasonis		100.0	0.46	FRU 99				
45625332	**Good identification**							**45625332**
Parabacteroides distasonis		100.0	0.42	FRU 99				
45625722	**Good identification**							**45625722**
Parabacteroides distasonis		100.0	0.43	FRU 99				
45625732	**Good identification**							**45625732**
Parabacteroides distasonis		100.0	0.38	FRU 99				
45630000	**Very good identification**							**45630000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		99.65	0.69		1	1	5	
Clostridium difficile		0.27	0.09		99	99	1	
Propionibacterium granulosum		0.08	0.13		99	1	90	
45630002	**Good identification**							**45630002**
Fusobacterium mortiferum		100.0	0.53	BGL 10				
45630100	**Very good identification**							**45630100**
Fusobacterium mortiferum		100.0	0.75					

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
45630101	**Good identification**						**45630101**
Fusobacterium mortiferum		100.0	0.41	URE 1			
45630102	**Very good identification**						**45630102**
Fusobacterium mortiferum		100.0	0.59	BGL 10			
45630110	**Good identification**						**45630110**
Fusobacterium mortiferum		100.0	0.41	MLZ 1			
45630120	**Acceptable identification**						**45630120**
					GRA	SPO	
Fusobacterium mortiferum		94.29	0.41	NAG 1	1	1	
Clostridium septicum		5.71	0.03		99	99	
45630140	**Good identification**						**45630140**
Fusobacterium mortiferum		100.0	0.41	SOR 1			
45630300	**Good identification**						**45630300**
Fusobacterium mortiferum		100.0	0.41	RHA 1			
45630500	**Good identification**						**45630500**
Fusobacterium mortiferum		100.0	0.41	ARA 1			
45631000	**Very good identification**						**45631000**
					GRA	CAT	
Fusobacterium mortiferum		99.90	0.72		1	5	
Propionibacterium granulosum		0.05	0.13		99	90	
Lactobacillus jensenii		0.05	0.06		99	1	
45631001	**Good identification**						**45631001**
Fusobacterium mortiferum		100.0	0.38	URE 1			
45631002	**Good identification**						**45631002**
					GRA		
Fusobacterium mortiferum		99.69	0.56	BGL 10	1		
Lactobacillus jensenii		0.31	0.03		99		
45631010	**Good identification**						**45631010**
Fusobacterium mortiferum		100.0	0.38	MLZ 1			
45631020	**Low discrimination**						**45631020**
					GRA	SPO	
Fusobacterium mortiferum		88.95	0.38	NAG 1	1	1	
Clostridium septicum		8.38	0.03		99	99	
Clostridium perfringens		2.67	0.01		99	84	
45631040	**Good identification**						**45631040**
Fusobacterium mortiferum		100.0	0.38	SOR 1			
45631100	**Very good identification**						**45631100**
					GRA	SPO	
Fusobacterium mortiferum		99.96	0.78		1	1	
Clostridium septicum		0.04	0.03		99	99	
45631101	**Good identification**						**45631101**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		98.67	0.44	URE 1	1		
Actinomyces naeslundii		1.33	0.24		99		
45631102	**Very good identification**						**45631102**
					GRA		
Fusobacterium mortiferum		99.36	0.62	BGL 10	1		
Parabacteroides distasonis		0.54	0.22		1		
Bifidobacterium longum subsp. infantis		0.10	-.07		99		
45631110	**Good identification**						**45631110**
Fusobacterium mortiferum		100.0	0.44	MLZ 1			
45631120	**Low discrimination**						**45631120**
					GRA	SPO	
Clostridium septicum		77.59	0.37	MLT 94	99	99	
Fusobacterium mortiferum		19.42	0.44	RAF 1	1	1	
Clostridium perfringens		1.83	0.16		99	84	
45631122	**Acceptable identification**						**45631122**
Parabacteroides distasonis		92.85	0.56	XYL 90			
Prevotella oralis		3.11	0.18				
Fusobacterium mortiferum		1.74	0.28				
45631132	**Good identification**						**45631132**
Parabacteroides distasonis		100.0	0.51	XYL 90			
45631140	**Good identification**						**45631140**
Fusobacterium mortiferum		100.0	0.44	SOR 1			
45631200	**Good identification**						**45631200**
Fusobacterium mortiferum		100.0	0.38	RHA 1			
45631300	**Good identification**						**45631300**
Fusobacterium mortiferum		100.0	0.44	RHA 1			
45631322	**Very good identification**						**45631322**
Parabacteroides distasonis		99.52	0.64	XYL 90			
Prevotella oralis		0.48	0.12				
45631332	**Very good identification**						**45631332**
Parabacteroides distasonis		100.0	0.59	XYL 90			
45631400	**Good identification**						**45631400**
Fusobacterium mortiferum		100.0	0.38	ARA 1			
45631500	**Good identification**						**45631500**
Fusobacterium mortiferum		100.0	0.44	ARA 1			
45631522	**Good identification**						**45631522**
Parabacteroides distasonis		100.0	0.52	XYL 90			
45631532	**Good identification**						**45631532**
Parabacteroides distasonis		100.0	0.48	XYL 90			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45631722	**Very good identification**							**45631722**
Parabacteroides distasonis		100.0	0.60	XYL 90				
45631732	**Good identification**							**45631732**
Parabacteroides distasonis		99.75	0.56	XYL 90				
Clostridium clostridioforme		0.25	0.06					
45632010	**Good identification**							**45632010**
Clostridium difficile		100.0	0.37	RAF 1				
45632100	**Good identification**							**45632100**
Fusobacterium mortiferum		100.0	0.41	MAN 1				
45633000	**Low discrimination**							**45633000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		88.73	0.38	MAN 1	1	1	5	
Clostridium innocuum		7.77	0.07		99	99	1	
Propionibacterium granulosum		3.50	0.10		99	1	90	
45633100	**Good identification**							**45633100**
					GRA			
Fusobacterium mortiferum		98.28	0.44	MAN 1	1			
Actinomyces naeslundii		1.72	0.26		99			
45634000	**Good identification**							**45634000**
Fusobacterium mortiferum		100.0	0.50	XYL 7				
45634100	**Good identification**							**45634100**
Fusobacterium mortiferum		100.0	0.56	XYL 7				
45634102	**Good identification**							**45634102**
Fusobacterium mortiferum		100.0	0.40	XYL 7				
45634122	**Good identification**							**45634122**
Parabacteroides distasonis		100.0	0.38	GAL 99				
45634322	**Good identification**							**45634322**
Parabacteroides distasonis		100.0	0.46	GAL 99				
45634332	**Good identification**							**45634332**
Parabacteroides distasonis		100.0	0.42	GAL 99				
45634722	**Good identification**							**45634722**
Parabacteroides distasonis		100.0	0.43	GAL 99				
45634732	**Acceptable identification**							**45634732**
Parabacteroides distasonis		91.53	0.38	GAL 99				
Clostridium clostridioforme		8.47	0.15					
45635000	**Good identification**							**45635000**
Fusobacterium mortiferum		100.0	0.53	XYL 7				
45635002	**Good identification**							**45635002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.36	XYL 7			
45635022	**Good identification**						**45635022**
Parabacteroides distasonis		100.0	0.38	LAC 99			
45635100	**Very good identification**						**45635100**
					COC	GRA	
Fusobacterium mortiferum		99.79	0.59	XYL 7	1	1	
Atopobium parvulum		0.21	0.02		99	99	
45635102	**Low discrimination**						**45635102**
					GRA		
Fusobacterium mortiferum		60.08	0.43	XYL 7	1		
Parabacteroides distasonis		39.02	0.38	BGL 10	1		
Bifidobacterium adolescentis		0.90	0.07		99		
45635120	**Low discrimination**						**45635120**
					GRA	SPO	
Parabacteroides distasonis		82.30	0.38	BGL 99	1	1	
Fusobacterium mortiferum		11.52	0.25		1	1	
Clostridium septicum		6.18	0.03		99	99	
45635122	**Very good identification**						**45635122**
					GRA		
Parabacteroides distasonis		99.92	0.72		1		
Prevotella oralis		0.05	0.02		1		
Bifidobacterium longum subsp. infantis		0.03	-.04		99		
45635123	**Good identification**						**45635123**
Parabacteroides distasonis		100.0	0.38	URE 1			
45635132	**Very good identification**						**45635132**
Parabacteroides distasonis		100.0	0.68				
45635162	**Good identification**						**45635162**
Parabacteroides distasonis		100.0	0.38	SOR 1			
45635222	**Good identification**						**45635222**
Parabacteroides distasonis		100.0	0.46	LAC 99			
45635232	**Good identification**						**45635232**
Parabacteroides distasonis		100.0	0.42	LAC 99			
45635302	**Good identification**						**45635302**
Parabacteroides distasonis		100.0	0.46	NAG 99			
45635312	**Good identification**						**45635312**
Parabacteroides distasonis		100.0	0.42	NAG 99			
45635320	**Good identification**						**45635320**
Parabacteroides distasonis		100.0	0.46	BGL 99			
45635322	**Very good identification**						**45635322**
Parabacteroides distasonis		100.0	0.80				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
45635323	**Good identification**					**45635323**
Parabacteroides distasonis		100.0	0.46	URE 1		
45635330	**Good identification**					**45635330**
Parabacteroides distasonis		100.0	0.42	BGL 99		
45635332	**Very good identification**					**45635332**
Parabacteroides distasonis		100.0	0.76			
45635333	**Good identification**					**45635333**
Parabacteroides distasonis		100.0	0.42	URE 1		
45635362	**Good identification**					**45635362**
Parabacteroides distasonis		100.0	0.46	SOR 1		
45635372	**Good identification**					**45635372**
Parabacteroides distasonis		100.0	0.42	SOR 1		
45635522	**Very good identification**					**45635522**
Parabacteroides distasonis		100.0	0.69			
45635532	**Very good identification**					**45635532**
Parabacteroides distasonis		99.89	0.64			
Clostridium clostridioforme		0.12	0.08			
45635622	**Good identification**					**45635622**
Parabacteroides distasonis		100.0	0.43	LAC 99		
45635632	**Acceptable identification**					**45635632**
Parabacteroides distasonis		92.35	0.38	LAC 99		
Clostridium clostridioforme		7.65	0.14			
45635702	**Good identification**					**45635702**
Parabacteroides distasonis		96.82	0.43	NAG 99		
Clostridium clostridioforme		3.18	0.11			
45635712	**Low discrimination**					**45635712**
Parabacteroides distasonis		78.28	0.38	NAG 99		
Clostridium clostridioforme		21.72	0.23			
45635720	**Low discrimination**					**45635720**
Parabacteroides distasonis		89.37	0.43	BGL 99		
Bacteroides vulgatus		10.63	0.08			
45635722	**Very good identification**					**45635722**
Parabacteroides distasonis		99.95	0.77			
Clostridium clostridioforme		0.03	0.11			
Bacteroides thetaiotaomicron		0.02	-.02			
45635723	**Good identification**					**45635723**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.43	URE 1			
45635730	**Good identification**						**45635730**
Parabacteroides distasonis		100.0	0.38	BGL 99			
45635732	**Very good identification**						**45635732**
Parabacteroides distasonis		99.72	0.72				
Clostridium clostridioforme		0.28	0.23				
45635733	**Good identification**						**45635733**
Parabacteroides distasonis		100.0	0.38	URE 1			
45635762	**Good identification**						**45635762**
Parabacteroides distasonis		100.0	0.43	SOR 1			
45635772	**Good identification**						**45635772**
Parabacteroides distasonis		100.0	0.38	SOR 1			
45637122	**Good identification**						**45637122**
Parabacteroides distasonis		100.0	0.38	MAN 1			
45637322	**Good identification**						**45637322**
Parabacteroides distasonis		100.0	0.46	MAN 1			
45637332	**Good identification**						**45637332**
Parabacteroides distasonis		100.0	0.42	MAN 1			
45637722	**Good identification**						**45637722**
Parabacteroides distasonis		100.0	0.43	MAN 1			
45637732	**Good identification**						**45637732**
Parabacteroides distasonis		100.0	0.38	MAN 1			
45640000	**Good identification**						**45640000**
Fusobacterium mortiferum		100.0	0.36	FRU 99			
45640100	**Good identification**						**45640100**
Fusobacterium mortiferum		100.0	0.42	FRU 99			
45641000	**Low discrimination**						**45641000**
					GRA		
Fusobacterium mortiferum		87.72	0.39	FRU 99	1		
Eubacterium tortuosum		12.28	0.07		99		
45641100	**Good identification**						**45641100**
Fusobacterium mortiferum		100.0	0.45	FRU 99			
45650000	**Very good identification**						**45650000**
					GRA	SPO	
Fusobacterium mortiferum		99.83	0.70		1	1	
Clostridium difficile		0.17	0.06		99	99	
45650001	**Good identification**						**45650001**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.36	URE 1			
45650002	**Good identification**						**45650002**
Fusobacterium mortiferum		100.0	0.53	BGL 10			
45650010	**Low discrimination**						**45650010**
					GRA	SPO	
Fusobacterium mortiferum		54.75	0.36	MLZ 1	1	1	
Clostridium difficile		45.25	0.18		99	99	
45650020	**Good identification**						**45650020**
Fusobacterium mortiferum		100.0	0.36	NAG 1			
45650040	**Good identification**						**45650040**
Fusobacterium mortiferum		100.0	0.36	SOR 1			
45650100	**Very good identification**						**45650100**
Fusobacterium mortiferum		100.0	0.76				
45650101	**Good identification**						**45650101**
Fusobacterium mortiferum		100.0	0.42	URE 1			
45650102	**Very good identification**						**45650102**
Fusobacterium mortiferum		100.0	0.60	BGL 10			
45650110	**Good identification**						**45650110**
Fusobacterium mortiferum		100.0	0.42	MLZ 1			
45650120	**Acceptable identification**						**45650120**
					GRA	SPO	
Fusobacterium mortiferum		92.31	0.42	NAG 1	1	1	
Clostridium septicum		5.29	0.03		99	99	
Clostridium paraputrificum		2.41	-.06		99	99	
45650140	**Good identification**						**45650140**
Fusobacterium mortiferum		100.0	0.42	SOR 1			
45650200	**Good identification**						**45650200**
Fusobacterium mortiferum		100.0	0.36	RHA 1			
45650300	**Good identification**						**45650300**
Fusobacterium mortiferum		100.0	0.42	RHA 1			
45650400	**Good identification**						**45650400**
Fusobacterium mortiferum		100.0	0.36	ARA 1			
45650500	**Good identification**						**45650500**
Fusobacterium mortiferum		100.0	0.42	ARA 1			
45651000	**Very good identification**						**45651000**
					GRA	SPO	
Fusobacterium mortiferum		99.37	0.73		1	1	
Lactobacillus jensenii		0.30	0.19		99	1	
Clostridium innocuum		0.22	0.15		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45651001	**Good identification**						**45651001**
Fusobacterium mortiferum		100.0	0.39	URE 1			
45651002	**Good identification**						**45651002**
					GRA	SPO	
Fusobacterium mortiferum		93.94	0.56	BGL 10	1	1	
Clostridium innocuum		1.91	0.15		99	99	
Lactobacillus jensenii		1.68	0.16		99	1	
45651010	**Acceptable identification**						**45651010**
					GRA		
Fusobacterium mortiferum		91.05	0.39	MLZ 1	1		
Lactobacillus jensenii		8.95	0.11		99		
45651020	**Low discrimination**						**45651020**
					GRA	SPO	
Fusobacterium mortiferum		72.53	0.39	NAG 1	1	1	
Clostridium septicum		6.46	0.03		99	99	
Prevotella oralis		5.60	0.05		1	1	
45651040	**Good identification**						**45651040**
Fusobacterium mortiferum		100.0	0.39	SOR 1			
45651100	**Very good identification**						**45651100**
					GRA		
Fusobacterium mortiferum		99.75	0.79		1		
Lactobacillus catenaformis		0.07	0.15		99		
Collinsella aerofaciens		0.05	-.03		99		
45651101	**Good identification**						**45651101**
Fusobacterium mortiferum		100.0	0.45	URE 1			
45651102	**Good identification**						**45651102**
					GRA		
Fusobacterium mortiferum		91.79	0.63	BGL 10	1		
Prevotella oralis		4.33	0.26		1		
Lactobacillus acidophilus		1.25	0.18		99		
45651110	**Good identification**						**45651110**
					GRA		
Fusobacterium mortiferum		98.60	0.45	MLZ 1	1		
Bifidobacterium breve		1.40	0.10		99		
45651120	**Low discrimination**						**45651120**
					GRA	SPO	
Clostridium septicum		52.83	0.37	MLT 94	99	99	
Clostridium paraputrificum		24.08	0.29	RAF 1	99	99	
Fusobacterium mortiferum		13.98	0.45		1	1	
45651122	**Low discrimination**						**45651122**
					GRA	SPO	
Prevotella oralis		81.32	0.48	SUC 92	1	1	
Clostridium paraputrificum		6.09	0.22	MLT 99	99	99	
Parabacteroides distasonis		5.17	0.40		1	1	
45651132	**Low discrimination**						**45651132**
					GRA		
Parabacteroides distasonis		48.70	0.35	XYL 90	1		
Bifidobacterium breve		36.92	0.32		99		
Prevotella oralis		14.37	0.14		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45651140	**Good identification**						**45651140**
					GRA		
Fusobacterium mortiferum		98.60	0.45	SOR 1	1		
Bifidobacterium breve		1.40	0.10		99		
45651200	**Good identification**						**45651200**
Fusobacterium mortiferum		100.0	0.39	RHA 1			
45651300	**Good identification**						**45651300**
Fusobacterium mortiferum		98.84	0.45	RHA 1			
Prevotella oralis		1.16	-.02				
45651322	**Low discrimination**						**45651322**
					GRA	SPO	
Prevotella oralis		68.68	0.41	SUC 92	1	1	
Parabacteroides distasonis		30.56	0.48	MLT 99	1	1	
Clostridium barati		0.38	0.02		99	99	
45651332	**Good identification**						**45651332**
Parabacteroides distasonis		95.96	0.44	XYL 90			
Prevotella oralis		4.05	0.07				
45651400	**Good identification**						**45651400**
					GRA	SPO	
Fusobacterium mortiferum		96.21	0.39	ARA 1	1	1	
Clostridium innocuum		3.79	0.02		99	99	
45651500	**Good identification**						**45651500**
Fusobacterium mortiferum		100.0	0.45	ARA 1			
45651522	**Low discrimination**						**45651522**
Parabacteroides distasonis		79.41	0.36	XYL 90			
Prevotella oralis		20.59	0.14				
45651722	**Good identification**						**45651722**
Parabacteroides distasonis		96.43	0.44	XYL 90			
Prevotella oralis		3.57	0.07				
45651732	**Good identification**						**45651732**
Parabacteroides distasonis		100.0	0.40	XYL 90			
45652100	**Good identification**						**45652100**
Fusobacterium mortiferum		100.0	0.42	MAN 1			
45653000	**Good identification**						**45653000**
					GRA	SPO	
Clostridium innocuum		95.67	0.49	RAF 1	99	99	
Fusobacterium mortiferum		4.33	0.39		1	1	
45653002	**Good identification**						**45653002**
					GRA	SPO	
Clostridium innocuum		99.50	0.49	RAF 1	99	99	
Fusobacterium mortiferum		0.50	0.22		1	1	
45653100	**Low discrimination**						**45653100**
					GRA	SPO	
Fusobacterium mortiferum		89.23	0.45	MAN 1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		8.54	0.15		99	99	
Bifidobacterium breve		1.27	0.10		99	1	
45653400	**Good identification**						**45653400**
Clostridium innocuum		100.0	0.36	RAF 1			
45653402	**Good identification**						**45653402**
Clostridium innocuum		100.0	0.36	RAF 1			
45654000	**Good identification**						**45654000**
Fusobacterium mortiferum		100.0	0.50	XYL 7			
45654100	**Very good identification**						**45654100**
Fusobacterium mortiferum		100.0	0.57	XYL 7			
45654102	**Good identification**						**45654102**
Fusobacterium mortiferum		100.0	0.40	XYL 7			
45655000	**Good identification**						**45655000**
Fusobacterium mortiferum		100.0	0.53	XYL 7			
45655002	**Good identification**						**45655002**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
45655100	**Very good identification**						**45655100**
					GRA	SPO	
Fusobacterium mortiferum		99.80	0.60	XYL 7	1	1	
Clostridium butyricum		0.20	-.05		99	99	
45655102	**Low discrimination**						**45655102**
Fusobacterium mortiferum		83.10	0.43	XYL 7			
Prevotella oralis		6.44	0.10	BGL 10			
Parabacteroides distasonis		5.72	0.22				
45655122	**Low discrimination**						**45655122**
Parabacteroides distasonis		81.52	0.56				
Prevotella oralis		17.61	0.32				
Bacteroides uniformis		0.42	-.02				
45655132	**Good identification**						**45655132**
Parabacteroides distasonis		99.60	0.52				
Prevotella oralis		0.40	-.02				
45655322	**Very good identification**						**45655322**
Parabacteroides distasonis		96.79	0.64				
Prevotella oralis		2.99	0.26				
Bacteroides uniformis		0.23	0.01				
45655332	**Very good identification**						**45655332**
Parabacteroides distasonis		100.0	0.60				
45655522	**Acceptable identification**						**45655522**
Parabacteroides distasonis		90.99	0.53				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides uniformis		8.69	0.16					
Prevotella oralis		0.32	-.02					
45655532	**Good identification**							**45655532**
Parabacteroides distasonis		100.0	0.48					
45655722	**Very good identification**							**45655722**
Parabacteroides distasonis		95.67	0.61					
Bacteroides uniformis		4.20	0.19					
Clostridium clostridioforme		0.12	0.06					
45655732	**Very good identification**							**45655732**
Parabacteroides distasonis		98.93	0.56					
Clostridium clostridioforme		1.07	0.17					
45661100	**Good identification**							**45661100**
Fusobacterium mortiferum		100.0	0.36	FRU 99				
45665322	**Good identification**							**45665322**
Parabacteroides distasonis		100.0	0.42	FRU 99				
45665332	**Good identification**							**45665332**
Parabacteroides distasonis		100.0	0.37	FRU 99				
45665722	**Good identification**							**45665722**
Parabacteroides distasonis		100.0	0.38	FRU 99				
45670000	**Very good identification**							**45670000**
Fusobacterium mortiferum		100.0	0.61					
45670002	**Good identification**							**45670002**
Fusobacterium mortiferum		100.0	0.44	BGL 10				
45670100	**Very good identification**							**45670100**
Fusobacterium mortiferum		100.0	0.67					
45670102	**Good identification**							**45670102**
Fusobacterium mortiferum		100.0	0.51	BGL 10				
45671000	**Very good identification**							**45671000**
					GRA	SPO		
Fusobacterium mortiferum		96.85	0.64		1	1		
Lactobacillus jensenii		2.90	0.28		99	1		
Clostridium innocuum		0.26	0.07		99	99		
45671002	**Low discrimination**							**45671002**
					GRA			
Fusobacterium mortiferum		81.02	0.47	BGL 10	1			
Lactobacillus jensenii		14.55	0.25		99			
Lactobacillus acidophilus		2.50	0.09		99			
45671100	**Very good identification**							**45671100**
					COC	GRA	SPO	
Fusobacterium mortiferum		99.52	0.70		1	1	1	
Clostridium septicum		0.39	0.11		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		0.06	0.04		99	99	1	
45671101	**Acceptable identification**							**45671101**
					GRA			
Fusobacterium mortiferum		94.91	0.36	URE 1	1			
Actinomyces naeslundii		5.10	0.26		99			
45671102	**Low discrimination**							**45671102**
					GRA			
Fusobacterium mortiferum		82.85	0.54	BGL 10	1			
Lactobacillus acidophilus		12.59	0.27		99			
Bifidobacterium breve		2.42	0.24		99			
45671110	**Good identification**							**45671110**
Fusobacterium mortiferum		96.10	0.36	MLZ 1				
Leptotrichia buccalis		3.90	-.11					
45671120	**Acceptable identification**							**45671120**
					GRA	SPO		
Clostridium septicum		92.78	0.45	MLT 94	99	99		
Clostridium paraputrificum		3.75	0.19	RAF 1	99	99		
Fusobacterium mortiferum		2.44	0.36		1	1		
45671122	**Low discrimination**							**45671122**
					GRA			
Parabacteroides distasonis		59.56	0.51	XYL 90	1			
Prevotella oralis		13.13	0.27		1			
Lactobacillus acidophilus		10.42	0.27		99			
45671132	**Low discrimination**							**45671132**
					GRA			
Parabacteroides distasonis		83.50	0.47	XYL 90	1			
Bifidobacterium breve		9.26	0.29		99			
Leptotrichia buccalis		7.24	0.07		1			
45671140	**Good identification**							**45671140**
Fusobacterium mortiferum		100.0	0.36	SOR 1				
45671300	**Good identification**							**45671300**
Fusobacterium mortiferum		100.0	0.36	RHA 1				
45671322	**Good identification**							**45671322**
					SPO			
Parabacteroides distasonis		94.50	0.59	XYL 90	1			
Prevotella oralis		2.98	0.21		1			
Clostridium ramosum		2.11	0.19		92			
45671332	**Good identification**							**45671332**
Parabacteroides distasonis		100.0	0.55	XYL 90				
45671500	**Low discrimination**							**45671500**
					GRA	SPO		
Fusobacterium mortiferum		82.04	0.36	ARA 1	1	1		
Clostridium butyricum		17.96	0.06		99	99		
45671522	**Good identification**							**45671522**
Parabacteroides distasonis		100.0	0.48	XYL 90				
45671532	**Good identification**							**45671532**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		100.0	0.43	XYL 90				
45671722	**Good identification**							**45671722**
Parabacteroides distasonis		100.0	0.56	XYL 90				
45671732	**Good identification**							**45671732**
Parabacteroides distasonis		98.47	0.51	XYL 90				
Clostridium clostridioforme		1.53	0.15					
45673000	**Good identification**							**45673000**
					GRA	SPO		
Clostridium innocuum		96.30	0.41	RAF 1	99	99		
Fusobacterium mortiferum		3.70	0.30		1	1		
45673002	**Good identification**							**45673002**
					GRA	SPO		
Clostridium innocuum		99.58	0.41	RAF 1	99	99		
Fusobacterium mortiferum		0.43	0.13		1	1		
45673100	**Low discrimination**							**45673100**
					GRA	SPO		
Fusobacterium mortiferum		81.24	0.36	MAN 1	1	1		
Clostridium innocuum		9.15	0.07		99	99		
Actinomyces naeslundii		5.64	0.28		99	1		
45674000	**Good identification**							**45674000**
Fusobacterium mortiferum		100.0	0.41	XYL 7				
45674100	**Good identification**							**45674100**
Fusobacterium mortiferum		100.0	0.48	XYL 7				
45674322	**Good identification**							**45674322**
Parabacteroides distasonis		100.0	0.42	GAL 99				
45674332	**Good identification**							**45674332**
Parabacteroides distasonis		100.0	0.37	GAL 99				
45674722	**Acceptable identification**							**45674722**
Parabacteroides distasonis		93.64	0.38	GAL 99				
Clostridium clostridioforme		6.37	0.12					
45675000	**Good identification**							**45675000**
Fusobacterium mortiferum		100.0	0.44	XYL 7				
45675100	**Low discrimination**							**45675100**
					COC	GRA	SPO	
Fusobacterium mortiferum		88.20	0.51	XYL 7	1	1	1	
Clostridium butyricum		9.31	0.15		1	99	99	
Atopobium parvulum		1.32	0.08		99	99	1	
45675122	**Very good identification**							**45675122**
Parabacteroides distasonis		99.70	0.68					
Prevotella oralis		0.30	0.12					
45675132	**Very good identification**							**45675132**
Parabacteroides distasonis		100.0	0.63					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45675222	**Good identification**						**45675222**
Parabacteroides distasonis		100.0	0.42	LAC 99			
45675232	**Good identification**						**45675232**
Parabacteroides distasonis		100.0	0.37	LAC 99			
45675302	**Good identification**						**45675302**
Parabacteroides distasonis		100.0	0.42	NAG 99			
45675312	**Acceptable identification**						**45675312**
Parabacteroides distasonis		93.68	0.37	NAG 99			
Clostridium clostridioforme		6.32	0.11				
45675320	**Good identification**						**45675320**
Parabacteroides distasonis		100.0	0.42	BGL 99			
45675322	**Very good identification**						**45675322**
Parabacteroides distasonis		99.96	0.76				
Prevotella oralis		0.04	0.05				
45675323	**Good identification**						**45675323**
Parabacteroides distasonis		100.0	0.42	URE 1			
45675330	**Good identification**						**45675330**
Parabacteroides distasonis		100.0	0.37	BGL 99			
45675332	**Very good identification**						**45675332**
Parabacteroides distasonis		99.93	0.71				
Clostridium clostridioforme		0.07	0.11				
45675333	**Good identification**						**45675333**
Parabacteroides distasonis		100.0	0.37	URE 1			
45675362	**Good identification**						**45675362**
Parabacteroides distasonis		100.0	0.42	SOR 1			
45675372	**Good identification**						**45675372**
Parabacteroides distasonis		100.0	0.37	SOR 1			
45675522	**Very good identification**						**45675522**
					GRA	SPO	
Parabacteroides distasonis		99.47	0.64		1	1	
Clostridium butyricum		0.45	0.07		99	99	
Clostridium clostridioforme		0.08	0.06		50	66	
45675532	**Very good identification**						**45675532**
					GRA		
Parabacteroides distasonis		98.91	0.59		1		
Clostridium clostridioforme		0.71	0.17		50		
Bifidobacterium dentium		0.39	-.03		99		
45675622	**Acceptable identification**						**45675622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		94.26	0.38	LAC 99			
Clostridium clostridioforme		5.74	0.11				
45675702	**Low discrimination**						**45675702**
					GRA	SPO	
Parabacteroides distasonis		79.54	0.38	NAG 99	1	1	
Clostridium clostridioforme		16.22	0.20		50	66	
Clostridium butyricum		4.24	-.01		99	99	
45675720	**Good identification**						**45675720**
Parabacteroides distasonis		100.0	0.38	BGL 99			
45675722	**Very good identification**						**45675722**
Parabacteroides distasonis		99.59	0.72				
Clostridium clostridioforme		0.21	0.20				
Bacteroides ovatus		0.12	0.06				
45675723	**Good identification**						**45675723**
Parabacteroides distasonis		100.0	0.38	URE 1			
45675732	**Very good identification**						**45675732**
Parabacteroides distasonis		98.09	0.67				
Clostridium clostridioforme		1.71	0.32				
Bacteroides ovatus		0.14	0.03				
45675762	**Good identification**						**45675762**
Parabacteroides distasonis		100.0	0.38	SOR 1			
45677102	**Acceptable identification**						**45677102**
Actinomyces israelii		92.86	0.37	SUC 98			
Bifidobacterium adolescentis		7.14	0.12	MLT 99			
45677322	**Good identification**						**45677322**
					SPO		
Parabacteroides distasonis		98.61	0.42	MAN 1	1		
Clostridium ramosum		1.39	-.03		92		
45677332	**Good identification**						**45677332**
Parabacteroides distasonis		100.0	0.37	MAN 1			
45677502	**Low discrimination**						**45677502**
					SPO		
Actinomyces israelii		46.25	0.35	SUC 98	1		
Bifidobacterium adolescentis		29.08	0.26	MLT 99	1		
Clostridium butyricum		20.50	0.08		99		
45677722	**Good identification**						**45677722**
Parabacteroides distasonis		96.32	0.38	MAN 1			
Bacteroides ovatus		3.68	-.02				
45701122	**Low discrimination**						**45701122**
					GRA		
Prevotella oralis		59.01	0.38	SUC 92	1		
Bifidobacterium longum subsp. infantis		38.64	0.32	FRU 99	99		
Parabacteroides distasonis		0.89	0.20		1		
45705322	**Low discrimination**						**45705322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		86.69	0.44	FRU 99	1		
Prevotella oralis		11.35	0.16		1		
Bifidobacterium longum subsp. infantis		1.16	-.04		99		
45705332	**Good identification**						**45705332**
Parabacteroides distasonis		100.0	0.40	FRU 99			
45705722	**Good identification**						**45705722**
Parabacteroides distasonis		96.43	0.41	FRU 99			
Bacteroides vulgatus		3.57	-.02				
45705732	**Good identification**						**45705732**
Parabacteroides distasonis		95.36	0.36	FRU 99			
Clostridium clostridioforme		4.64	0.08				
45710000	**Very good identification**						**45710000**
					GRA	SPO	
Fusobacterium mortiferum		97.58	0.62	MLT 11	1	1	
Prevotella intermedia		1.57	0.20		1	1	
Clostridium bifermentans		0.65	0.11		99	99	
45710002	**Good identification**						**45710002**
					GRA		
Fusobacterium mortiferum		99.12	0.46	MLT 11	1		
Bifidobacterium longum subsp. infantis		0.88	-.07	BGL 10	99		
45710100	**Very good identification**						**45710100**
					GRA		
Fusobacterium mortiferum		99.59	0.69	MLT 11	1		
Bifidobacterium longum subsp. infantis		0.41	0.10		99		
45710102	**Low discrimination**						**45710102**
					GRA		
Fusobacterium mortiferum		69.91	0.52	MLT 11	1		
Bifidobacterium longum subsp. infantis		26.32	0.27	BGL 10	99		
Prevotella oralis		3.76	0.16		1		
45710122	**Low discrimination**						**45710122**
					GRA		
Prevotella oralis		59.27	0.38	SUC 92	1		
Bifidobacterium longum subsp. infantis		38.80	0.32	GAL 99	99		
Parabacteroides distasonis		0.89	0.20		1		
45711000	**Very good identification**						**45711000**
					GRA		
Fusobacterium mortiferum		97.00	0.65	MLT 11	1		
Prevotella intermedia		1.04	0.20		1		
Lactobacillus jensenii		0.78	0.19		99		
45711002	**Low discrimination**						**45711002**
					GRA		
Fusobacterium mortiferum		45.96	0.49	MLT 11	1		
Bifidobacterium longum subsp. infantis		26.92	0.27	BGL 10	99		
Prevotella oralis		24.31	0.30		1		
45711022	**Low discrimination**						**45711022**
					GRA		
Prevotella oralis		88.81	0.52	SUC 92	1		
Bifidobacterium longum subsp. infantis		9.20	0.32	LAC 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		1.46	0.25		99		
45711100	**Low discrimination**						**45711100**
					GRA		
Fusobacterium mortiferum		76.48	0.72	MLT 11	1		
Bifidobacterium longum subsp. infantis		20.88	0.44		99		
Prevotella oralis		1.59	0.29		1		
45711101	**Low discrimination**						**45711101**
					GRA		
Fusobacterium mortiferum		65.09	0.37	MLT 11	1		
Bifidobacterium longum subsp. infantis		17.77	0.10	URE 1	99		
Actinomyces naeslundii		17.14	0.39		99		
45711102	**Low discrimination**						**45711102**
					GRA		
Bifidobacterium longum subsp. infantis		84.05	0.62	SUC 99	99		
Prevotella oralis		12.01	0.51		1		
Fusobacterium mortiferum		3.38	0.55		1		
45711110	**Low discrimination**						**45711110**
					GRA		
Fusobacterium mortiferum		76.25	0.37	MLT 11	1		
Bifidobacterium longum subsp. infantis		20.82	0.10	MLZ 1	99		
Bifidobacterium breve		2.93	0.10		99		
45711120	**Low discrimination**						**45711120**
					GRA	SPO	
Clostridium septicum		31.65	0.49	RAF 1	99	99	
Bifidobacterium longum subsp. infantis		30.16	0.49		99	1	
Prevotella oralis		24.52	0.51		1	1	
45711122	**Low discrimination**						**45711122**
					GRA		
Prevotella oralis		59.29	0.72	SUC 92	1		
Bifidobacterium longum subsp. infantis		38.82	0.66		99		
Parabacteroides distasonis		0.89	0.54		1		
45711123	**Low discrimination**						**45711123**
					GRA		
Prevotella oralis		58.39	0.38	SUC 92	1		
Bifidobacterium longum subsp. infantis		38.23	0.32	URE 1	99		
Actinomyces naeslundii		2.05	0.39		99		
45711132	**Low discrimination**						**45711132**
					GRA		
Prevotella oralis		38.72	0.38	SUC 92	1		
Parabacteroides distasonis		30.94	0.49	MLZ 1	1		
Bifidobacterium longum subsp. infantis		25.35	0.32		99		
45711140	**Low discrimination**						**45711140**
					GRA		
Fusobacterium mortiferum		70.72	0.37	MLT 11	1		
Bifidobacterium longum subsp. infantis		19.31	0.10	SOR 1	99		
Clostridium perfringens		5.07	0.07		99		
45711162	**Low discrimination**						**45711162**
					GRA		
Prevotella oralis		51.01	0.38	SUC 92	1		
Bifidobacterium longum subsp. infantis		33.39	0.32	SOR 1	99		
Clostridium perfringens		7.42	0.27		99		
45711222	**Good identification**						**45711222**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Prevotella oralis		98.13	0.46	SUC 92	1		
Parabacteroides distasonis		1.63	0.28	LAC 94	1		
Bifidobacterium longum subsp. infantis		0.24	-.03		99		
45711300	**Low discrimination**						**45711300**
					GRA		
Fusobacterium mortiferum		32.67	0.37	MLT 11	1		
Eubacterium contortum		29.62	0.24	RHA 1	99		
Prevotella oralis		28.79	0.22		1		
45711302	**Low discrimination**						**45711302**
					GRA		
Prevotella oralis		83.25	0.44	SUC 92	1		
Bifidobacterium longum subsp. infantis		13.73	0.27	NAG 95	99		
Parabacteroides distasonis		1.68	0.28		1		
45711320	**Low discrimination**						**45711320**
					GRA	SPO	
Prevotella oralis		85.97	0.44	SUC 92	1	1	
Bacteroides vulgatus		5.33	0.18	BGL 95	1	1	
Clostridium septicum		2.62	0.15		99	99	
45711322	**Low discrimination**						**45711322**
					GRA		
Prevotella oralis		89.23	0.66	SUC 92	1		
Parabacteroides distasonis		9.36	0.62		1		
Bifidobacterium longum subsp. infantis		1.38	0.32		99		
45711332	**Low discrimination**						**45711332**
					GRA		
Parabacteroides distasonis		84.51	0.58	XYL 90	1		
Prevotella oralis		15.11	0.32		1		
Bifidobacterium longum subsp. infantis		0.23	-.03		99		
45711500	**Low discrimination**						**45711500**
					GRA		
Fusobacterium mortiferum		73.22	0.37	MLT 11	1		
Bifidobacterium longum subsp. infantis		19.99	0.10	ARA 1	99		
Bifidobacterium longum subsp. longum		3.49	0.09		99		
45711502	**Low discrimination**						**45711502**
					GRA		
Bifidobacterium adolescentis		56.59	0.45	SUC 99	99		
Bifidobacterium longum subsp. infantis		35.03	0.27	XYL 87	99		
Prevotella oralis		5.01	0.16		1		
45711512	**Good identification**						**45711512**
Bifidobacterium adolescentis		96.27	0.46	SUC 99			
Bifidobacterium longum subsp. longum		1.43	0.13	XYL 87			
Clostridium clostridioforme		1.15	0.10				
45711522	**Low discrimination**						**45711522**
					GRA		
Prevotella oralis		37.74	0.38	SUC 92	1		
Parabacteroides distasonis		34.32	0.50	ARA 1	1		
Bifidobacterium longum subsp. infantis		24.71	0.32		99		
45711532	**Low discrimination**						**45711532**
					GRA		
Parabacteroides distasonis		85.78	0.46	XYL 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		7.58	0.24		99	
Prevotella oralis		1.77	0.04		1	
45711720	**Good identification**					**45711720**
Bacteroides vulgatus		96.46	0.37	SUC 96		
Parabacteroides distasonis		1.45	0.24	XYL 98		
Prevotella oralis		1.18	0.10			
45711722	**Low discrimination**					**45711722**
Parabacteroides distasonis		85.37	0.58	XYL 90		
Prevotella oralis		13.41	0.32			
Bacteroides vulgatus		0.58	0.03			
45711732	**Good identification**					**45711732**
Parabacteroides distasonis		95.58	0.54	XYL 90		
Clostridium clostridioforme		4.14	0.25			
Prevotella oralis		0.28	-.02			
45713001	**Good identification**					**45713001**
Actinomyces naeslundii		100.0	0.36	ESL 12		
45713003	**Good identification**					**45713003**
Actinomyces naeslundii		100.0	0.36	ESL 12		
45713021	**Good identification**					**45713021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
45713023	**Good identification**					**45713023**
Actinomyces naeslundii		100.0	0.36	ESL 12		
45713100	**Low discrimination**					**45713100**
					GRA	
Fusobacterium mortiferum		57.79	0.37	MLT 11	1	
Actinomyces naeslundii		19.70	0.41	MAN 1	99	
Bifidobacterium longum subsp. infantis		15.78	0.10		99	
45713101	**Good identification**					**45713101**
Actinomyces naeslundii		100.0	0.44	ESL 12		
45713103	**Acceptable identification**					**45713103**
Actinomyces naeslundii		94.83	0.44	ESL 12		
Bifidobacterium longum subsp. infantis		5.17	-.07	SUC 99		
45713121	**Good identification**					**45713121**
Actinomyces naeslundii		100.0	0.44	ESL 12		
45713122	**Low discrimination**					**45713122**
					GRA	
Prevotella oralis		54.54	0.38	SUC 92	1	
Bifidobacterium longum subsp. infantis		35.70	0.32	MAN 1	99	
Bifidobacterium breve		5.02	0.32		99	
45713123	**Low discrimination**					**45713123**
					GRA	
Actinomyces naeslundii		80.31	0.44	ESL 12	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		11.90	0.04	SUC 99	1	
Bifidobacterium longum subsp. infantis		7.79	-.03		99	
45713502	**Good identification**					**45713502**
Bifidobacterium adolescentis		96.72	0.43	SUC 99		
Actinomyces israelii		2.54	0.22	XYL 87		
Bifidobacterium longum subsp. infantis		0.74	-.07			
45713512	**Good identification**					**45713512**
Bifidobacterium adolescentis		100.0	0.45	SUC 99		
45714000	**Good identification**					**45714000**
Fusobacterium mortiferum		100.0	0.43	MLT 11		
45714100	**Good identification**					**45714100**
					GRA	
Fusobacterium mortiferum		95.70	0.49	MLT 11	1	
Bifidobacterium longum subsp. infantis		4.30	0.09	XYL 7	99	
45714322	**Low discrimination**					**45714322**
					GRA	
Parabacteroides distasonis		86.69	0.44	GAL 99	1	
Prevotella oralis		11.35	0.16		1	
Bifidobacterium longum subsp. infantis		1.16	-.04		99	
45714332	**Acceptable identification**					**45714332**
Parabacteroides distasonis		94.09	0.40	GAL 99		
Clostridium clostridioforme		5.91	0.13			
45714722	**Low discrimination**					**45714722**
Parabacteroides distasonis		81.52	0.41	GAL 99		
Clostridium clostridioforme		15.46	0.23			
Bacteroides vulgatus		3.02	-.02			
45715000	**Acceptable identification**					**45715000**
					GRA	
Fusobacterium mortiferum		92.39	0.46	MLT 11	1	
Bifidobacterium longum subsp. infantis		6.46	0.09	XYL 7	99	
Actinomyces israelii		1.15	0.16		99	
45715022	**Low discrimination**					**45715022**
					GRA	
Prevotella oralis		53.30	0.36	SUC 92	1	
Bifidobacterium longum subsp. infantis		36.55	0.30	XYL 11	99	
Parabacteroides distasonis		9.21	0.36	LAC 94	1	
45715100	**Low discrimination**					**45715100**
					GRA	
Bifidobacterium longum subsp. infantis		71.19	0.43	SUC 99	99	
Fusobacterium mortiferum		23.98	0.52	BGL 91	1	
Eubacterium contortum		2.92	0.24		99	
45715102	**Very good identification**					**45715102**
					GRA	
Bifidobacterium longum subsp. infantis		95.04	0.60	SUC 99	99	
Prevotella oralis		2.05	0.35		1	
Bifidobacterium adolescentis		1.90	0.45		99	
45715112	**Low discrimination**					**45715112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium adolescentis		65.04	0.46	SUC 99	99	
Bifidobacterium longum subsp. infantis		26.95	0.26	ARA 87	99	
Parabacteroides distasonis		6.50	0.32		1	
45715120	**Low discrimination**					**45715120**
					GRA	
Bifidobacterium longum subsp. infantis		75.11	0.47	SUC 99	99	
Bacteroides vulgatus		10.79	0.31	BGL 91	1	
Prevotella oralis		9.22	0.35		1	
45715122	**Low discrimination**					**45715122**
					GRA	
Bifidobacterium longum subsp. infantis		66.22	0.64	SUC 99	99	
Parabacteroides distasonis		16.68	0.70		1	
Prevotella oralis		15.28	0.57		1	
45715132	**Good identification**					**45715132**
					GRA	
Parabacteroides distasonis		91.21	0.66		1	
Bifidobacterium longum subsp. infantis		6.79	0.30		99	
Prevotella oralis		1.57	0.23		1	
45715222	**Low discrimination**					**45715222**
					GRA	
Parabacteroides distasonis		54.06	0.44	LAC 99	1	
Prevotella oralis		44.71	0.30		1	
Bifidobacterium longum subsp. infantis		0.72	-.04		99	
45715232	**Acceptable identification**					**45715232**
Parabacteroides distasonis		94.68	0.40	LAC 99		
Clostridium clostridioforme		5.32	0.13			
45715300	**Very good identification**					**45715300**
					GRA	
Eubacterium contortum		98.95	0.58	SUC 99	99	
Bacteroides vulgatus		0.57	0.13		1	
Bifidobacterium longum subsp. infantis		0.25	0.09		99	
45715302	**Low discrimination**					**45715302**
					GRA	
Eubacterium contortum		67.93	0.45	SUC 99	99	
Parabacteroides distasonis		13.02	0.44		1	
Bifidobacterium longum subsp. infantis		9.69	0.26		99	
45715312	**Low discrimination**					**45715312**
					GRA	
Parabacteroides distasonis		75.76	0.40	NAG 99	1	
Clostridium clostridioforme		14.27	0.22		50	
Eubacterium contortum		7.42	0.11		99	
45715320	**Low discrimination**					**45715320**
					GRA	
Bacteroides vulgatus		70.44	0.47	SUC 96	1	
Eubacterium contortum		21.94	0.45	ARA 93	99	
Parabacteroides distasonis		4.20	0.44		1	
45715322	**Low discrimination**					**45715322**
					GRA	
Parabacteroides distasonis		86.52	0.78		1	
Prevotella oralis		11.32	0.51		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. infantis		1.16	0.30		99	
45715323	**Low discrimination**					**45715323**
					GRA	
Parabacteroides distasonis		86.69	0.44	URE 1	1	
Prevotella oralis		11.35	0.16		1	
Bifidobacterium longum subsp. infantis		1.16	-.04		99	
45715330	**Low discrimination**					**45715330**
					GRA	
Parabacteroides distasonis		70.81	0.40	BGL 99	1	
Bacteroides vulgatus		22.26	0.13		1	
Eubacterium contortum		6.93	0.11		99	
45715332	**Very good identification**					**45715332**
Parabacteroides distasonis		99.52	0.74			
Prevotella oralis		0.24	0.16			
Clostridium clostridioforme		0.19	0.22			
45715333	**Good identification**					**45715333**
Parabacteroides distasonis		100.0	0.40	URE 1		
45715362	**Low discrimination**					**45715362**
					GRA	
Parabacteroides distasonis		86.69	0.44	SOR 1	1	
Prevotella oralis		11.35	0.16		1	
Bifidobacterium longum subsp. infantis		1.16	-.04		99	
45715372	**Good identification**					**45715372**
Parabacteroides distasonis		100.0	0.40	SOR 1		
45715502	**Low discrimination**					**45715502**
					GRA	
Bifidobacterium adolescentis		89.02	0.59	SUC 99	99	
Bifidobacterium longum subsp. infantis		6.74	0.26		99	
Parabacteroides distasonis		1.85	0.33		1	
45715512	**Very good identification**					**45715512**
					GRA	
Bifidobacterium adolescentis		96.65	0.60	SUC 99	99	
Clostridium clostridioforme		1.74	0.27		50	
Parabacteroides distasonis		0.88	0.28		1	
45715520	**Good identification**					**45715520**
					GRA	
Bacteroides vulgatus		97.66	0.50	SUC 96	1	
Parabacteroides distasonis		0.81	0.33	RHA 90	1	
Bifidobacterium longum subsp. longum		0.74	0.28		99	
45715522	**Low discrimination**					**45715522**
					GRA	
Parabacteroides distasonis		89.74	0.67		1	
Bifidobacterium longum subsp. infantis		5.87	0.30		99	
Prevotella oralis		1.36	0.23		1	
45715532	**Good identification**					**45715532**
					GRA	
Parabacteroides distasonis		94.19	0.62		1	
Bifidobacterium longum subsp. longum		2.78	0.32		99	
Clostridium clostridioforme		1.87	0.27		50	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
45715542	**Good identification**					**45715542**
Bifidobacterium adolescentis		97.99	0.43	SUC 99		
Actinomyces israelii		2.01	0.21	SOR 11		
45715552	**Good identification**					**45715552**
Bifidobacterium adolescentis		100.0	0.45	SUC 99		
45715622	**Low discrimination**					**45715622**
Parabacteroides distasonis		82.85	0.41	LAC 99		
Clostridium clostridioforme		14.08	0.22			
Bacteroides vulgatus		3.07	-.02			
45715700	**Good identification**					**45715700**
					GRA	
Eubacterium contortum		91.50	0.57	SUC 99	99	
Bacteroides vulgatus		8.50	0.32		1	
45715702	**Low discrimination**					**45715702**
					GRA	
Eubacterium contortum		79.76	0.44	SUC 99	99	
Parabacteroides distasonis		11.45	0.41		1	
Clostridium clostridioforme		6.51	0.31		50	
45715712	**Low discrimination**					**45715712**
					GRA	
Clostridium clostridioforme		76.26	0.42	SUC 99	50	
Parabacteroides distasonis		15.84	0.36	TRE 88	1	
Bifidobacterium adolescentis		5.83	0.26		99	
45715720	**Very good identification**					**45715720**
					GRA	
Bacteroides vulgatus		97.85	0.66	SUC 96	1	
Eubacterium contortum		1.88	0.44		99	
Parabacteroides distasonis		0.27	0.41		1	
45715722	**Good identification**					**45715722**
					GRA	
Parabacteroides distasonis		94.46	0.75		1	
Bacteroides vulgatus		3.50	0.32		1	
Eubacterium contortum		1.17	0.31		99	
45715723	**Good identification**					**45715723**
Parabacteroides distasonis		96.43	0.41	URE 1		
Bacteroides vulgatus		3.57	-.02			
45715732	**Very good identification**					**45715732**
Parabacteroides distasonis		95.30	0.70			
Clostridium clostridioforme		4.63	0.42			
Bacteroides vulgatus		0.07	-.02			
45715733	**Good identification**					**45715733**
Parabacteroides distasonis		95.36	0.36	URE 1		
Clostridium clostridioforme		4.64	0.08			
45715762	**Good identification**					**45715762**
Parabacteroides distasonis		96.43	0.41	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		3.57	-.02			
45715772	**Good identification**					**45715772**
Parabacteroides distasonis		95.36	0.36	SOR 1		
Clostridium clostridioforme		4.64	0.08			
45717100	**Low discrimination**					**45717100**
					GRA	
Actinomyces israelii		72.78	0.39	SUC 98	99	
Bifidobacterium longum subsp. infantis		18.13	0.09	CEL 93	99	
Fusobacterium mortiferum		6.11	0.18		1	
45717102	**Low discrimination**					**45717102**
Bifidobacterium adolescentis		49.97	0.43	SUC 99		
Bifidobacterium longum subsp. infantis		30.93	0.26	ARA 87		
Actinomyces israelii		18.43	0.42			
45717112	**Good identification**					**45717112**
Bifidobacterium adolescentis		95.69	0.45	SUC 99		
Actinomyces israelii		4.31	0.28	ARA 87		
45717142	**Low discrimination**					**45717142**
Actinomyces israelii		53.71	0.35	SUC 98		
Bifidobacterium adolescentis		46.29	0.28	CEL 93		
45717322	**Low discrimination**					**45717322**
					GRA	
Parabacteroides distasonis		86.69	0.44	MAN 1	1	
Prevotella oralis		11.35	0.16		1	
Bifidobacterium longum subsp. infantis		1.16	-.04		99	
45717332	**Good identification**					**45717332**
Parabacteroides distasonis		100.0	0.40	MAN 1		
45717500	**Low discrimination**					**45717500**
					SPO	
Actinomyces israelii		71.88	0.38	SUC 98	1	
Bifidobacterium adolescentis		24.16	0.23	CEL 93	1	
Clostridium butyricum		3.96	-.05		99	
45717502	**Very good identification**					**45717502**
Bifidobacterium adolescentis		95.69	0.57	SUC 99		
Actinomyces israelii		4.31	0.41			
45717512	**Very good identification**					**45717512**
Bifidobacterium adolescentis		99.29	0.59	SUC 99		
Actinomyces israelii		0.55	0.26			
Bifidobacterium dentium		0.17	-.07			
45717542	**Low discrimination**					**45717542**
Bifidobacterium adolescentis		87.58	0.42	SUC 99		
Actinomyces israelii		12.42	0.34	SOR 11		
45717552	**Good identification**					**45717552**
Bifidobacterium adolescentis		98.30	0.43	SUC 99		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		1.70	0.19	SOR 11		
45717722	**Good identification**					**45717722**
Parabacteroides distasonis		96.43	0.41	MAN 1		
Bacteroides vulgatus		3.57	-.02			
45717732	**Good identification**					**45717732**
Parabacteroides distasonis		95.36	0.36	MAN 1		
Clostridium clostridioforme		4.64	0.08			
45721322	**Low discrimination**					**45721322**
Parabacteroides distasonis		88.22	0.39	FRU 99		
Prevotella oralis		11.78	0.12	XYL 90		
45721722	**Good identification**					**45721722**
Parabacteroides distasonis		100.0	0.36	FRU 99		
45725122	**Good identification**					**45725122**
					GRA	
Parabacteroides distasonis		97.88	0.47	FRU 99	1	
Prevotella oralis		1.26	0.02		1	
Bifidobacterium longum subsp. infantis		0.86	-.04		99	
45725132	**Good identification**					**45725132**
Parabacteroides distasonis		100.0	0.43	FRU 99		
45725322	**Good identification**					**45725322**
Parabacteroides distasonis		100.0	0.56	FRU 99		
45725332	**Good identification**					**45725332**
Parabacteroides distasonis		100.0	0.51	FRU 99		
45725522	**Good identification**					**45725522**
Parabacteroides distasonis		100.0	0.44	FRU 99		
45725532	**Good identification**					**45725532**
Parabacteroides distasonis		96.90	0.39	FRU 99		
Clostridium clostridioforme		3.10	0.08			
45725722	**Good identification**					**45725722**
Parabacteroides distasonis		99.08	0.52	FRU 99		
Clostridium clostridioforme		0.92	0.11			
45725732	**Acceptable identification**					**45725732**
Parabacteroides distasonis		92.74	0.47	FRU 99		
Clostridium clostridioforme		7.26	0.23			
45730000	**Good identification**					**45730000**
Fusobacterium mortiferum		100.0	0.53	MLT 11		
45730002	**Good identification**					**45730002**
Fusobacterium mortiferum		100.0	0.37	MLT 11		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45730100	**Very good identification**							**45730100**
Fusobacterium mortiferum		100.0	0.60	MLT 11				
45730102	**Good identification**							**45730102**
					GRA			
Fusobacterium mortiferum		98.74	0.43	MLT 11	1			
Bifidobacterium longum subsp. infantis		1.26	-.07	BGL 10	99			
45730322	**Low discrimination**							**45730322**
Parabacteroides distasonis		88.22	0.39	GAL 99				
Prevotella oralis		11.78	0.12	XYL 90				
45730722	**Low discrimination**							**45730722**
Parabacteroides distasonis		78.63	0.36	GAL 99				
Clostridium clostridioforme		21.37	0.20	XYL 90				
45731000	**Good identification**							**45731000**
					GRA			
Fusobacterium mortiferum		91.10	0.56	MLT 11	1			
Lactobacillus jensenii		7.35	0.28		99			
Actinomyces naeslundii		0.86	0.33		99			
45731001	**Low discrimination**							**45731001**
					GRA			
Actinomyces naeslundii		58.39	0.36	ESL 12	99			
Fusobacterium mortiferum		41.61	0.22	SUC 99	1			
45731002	**Low discrimination**							**45731002**
					GRA			
Fusobacterium mortiferum		57.67	0.40	MLT 11	1			
Lactobacillus jensenii		27.93	0.25	BGL 10	99			
Prevotella oralis		6.52	0.10		1			
45731003	**Good identification**							**45731003**
Actinomyces naeslundii		100.0	0.36	ESL 12				
45731020	**Low discrimination**							**45731020**
					SPO			
Clostridium perfringens		58.44	0.35	SUC 91	84			
Clostridium septicum		28.99	0.23	LAC 88	99			
Lactobacillus jensenii		4.85	0.17		1			
45731021	**Low discrimination**							**45731021**
Actinomyces naeslundii		85.24	0.36	ESL 12				
Clostridium perfringens		14.76	0.01	SUC 99				
45731023	**Low discrimination**							**45731023**
					GRA			
Actinomyces naeslundii		85.47	0.36	ESL 12	99			
Prevotella oralis		14.53	-.03	SUC 99	1			
45731100	**Low discrimination**							**45731100**
					COC	GRA	SPO	
Fusobacterium mortiferum		85.36	0.63	MLT 11	1	1	1	
Atopobium parvulum		5.92	0.31		99	99	1	
Clostridium septicum		4.41	0.23		1	99	99	
45731101	**Low discrimination**							**45731101**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		63.12	0.44	ESL 12	99		
Fusobacterium mortiferum		36.88	0.28	SUC 99	1		
45731102	**Low discrimination**						**45731102**
					GRA		
Fusobacterium mortiferum		29.49	0.46	MLT 11	1		
Bifidobacterium longum subsp. infantis		24.78	0.27	BGL 10	99		
Prevotella oralis		22.38	0.30		1		
45731103	**Low discrimination**						**45731103**
					GRA		
Actinomyces naeslundii		89.33	0.44	ESL 12	99		
Fusobacterium mortiferum		5.80	0.12	SUC 99	1		
Bifidobacterium longum subsp. infantis		4.87	-.07		99		
45731120	**Low discrimination**						**45731120**
					GRA	SPO	
Clostridium septicum		84.15	0.57	RAF 1	99	99	
Clostridium perfringens		12.57	0.50		99	84	
Prevotella oralis		1.39	0.30		1	1	
45731122	**Low discrimination**						**45731122**
					GRA		
Parabacteroides distasonis		42.69	0.65	XYL 90	1		
Prevotella oralis		39.91	0.52		1		
Clostridium perfringens		8.96	0.44		99		
45731132	**Good identification**						**45731132**
					GRA		
Parabacteroides distasonis		94.74	0.61	XYL 90	1		
Prevotella oralis		1.66	0.18		1		
Bifidobacterium breve		1.60	0.29		99		
45731160	**Low discrimination**						**45731160**
Clostridium perfringens		71.36	0.38	SUC 91			
Clostridium septicum		25.34	0.23				
Clostridium paraputrificum		1.54	-.01				
45731222	**Low discrimination**						**45731222**
Parabacteroides distasonis		54.24	0.39	XYL 90			
Prevotella oralis		45.77	0.25	LAC 99			
45731302	**Low discrimination**						**45731302**
Parabacteroides distasonis		56.58	0.39	XYL 90			
Prevotella oralis		39.37	0.24	NAG 99			
Clostridium clostridioforme		1.81	0.08				
45731320	**Low discrimination**						**45731320**
					GRA	SPO	
Parabacteroides distasonis		33.88	0.39	XYL 90	1	1	
Clostridium septicum		33.71	0.23	BGL 99	99	99	
Prevotella oralis		23.58	0.24		1	1	
45731322	**Low discrimination**						**45731322**
					SPO		
Parabacteroides distasonis		87.84	0.73	XYL 90	1		
Prevotella oralis		11.73	0.46		1		
Clostridium ramosum		0.30	0.19		92		
45731323	**Low discrimination**						**45731323**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		88.22	0.39	XYL 90		
Prevotella oralis		11.78	0.12	URE 1		
45731332	**Very good identification**					**45731332**
Parabacteroides distasonis		99.48	0.69	XYL 90		
Clostridium clostridioforme		0.27	0.19			
Prevotella oralis		0.25	0.12			
45731362	**Low discrimination**					**45731362**
Parabacteroides distasonis		88.22	0.39	XYL 90		
Prevotella oralis		11.78	0.12	SOR 1		
45731502	**Low discrimination**					**45731502**
					GRA	
Bifidobacterium adolescentis		82.94	0.42	SUC 99	99	
Parabacteroides distasonis		8.75	0.27	XYL 87	1	
Clostridium clostridioforme		2.92	0.14		50	
45731512	**Low discrimination**					**45731512**
					GRA	
Bifidobacterium adolescentis		84.27	0.43	SUC 99	99	
Clostridium clostridioforme		11.05	0.25	XYL 87	50	
Parabacteroides distasonis		3.92	0.23		1	
45731522	**Very good identification**					**45731522**
					GRA	
Parabacteroides distasonis		97.34	0.62	XYL 90	1	
Prevotella oralis		1.50	0.18		1	
Clostridium perfringens		0.34	0.10		99	
45731532	**Very good identification**					**45731532**
					GRA	
Parabacteroides distasonis		96.57	0.57	XYL 90	1	
Clostridium clostridioforme		2.75	0.25		50	
Bifidobacterium longum subsp. longum		0.47	0.13		99	
45731622	**Low discrimination**					**45731622**
Parabacteroides distasonis		80.42	0.36	XYL 90		
Clostridium clostridioforme		19.58	0.19	LAC 99		
45731702	**Low discrimination**					**45731702**
					GRA	
Parabacteroides distasonis		54.14	0.36	XYL 90	1	
Clostridium clostridioforme		44.13	0.28	NAG 99	50	
Bifidobacterium adolescentis		1.73	0.07		99	
45731712	**Low discrimination**					**45731712**
					GRA	
Clostridium clostridioforme		86.54	0.40	SUC 99	50	
Parabacteroides distasonis		12.55	0.31	XYL 91	1	
Bifidobacterium adolescentis		0.91	0.09		99	
45731720	**Low discrimination**					**45731720**
Parabacteroides distasonis		62.10	0.36	XYL 90		
Bacteroides vulgatus		37.90	0.14	BGL 99		
45731722	**Very good identification**					**45731722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		98.93	0.70	XYL 90		
Clostridium clostridioforme		0.82	0.28			
Prevotella oralis		0.22	0.12			
45731723	**Good identification**					**45731723**
Parabacteroides distasonis		100.0	0.36	XYL 90		
45731732	**Good identification**					**45731732**
Parabacteroides distasonis		93.49	0.65	XYL 90		
Clostridium clostridioforme		6.51	0.40			
45731762	**Good identification**					**45731762**
Parabacteroides distasonis		100.0	0.36	XYL 90		
45733000	**Low discrimination**					**45733000**
					GRA	
Actinomyces naeslundii		53.58	0.38	ESL 12	99	
Fusobacterium mortiferum		29.51	0.22	SUC 99	1	
Actinomyces israelii		16.91	0.20		99	
45733001	**Good identification**					**45733001**
Actinomyces naeslundii		100.0	0.41	ESL 12		
45733002	**Low discrimination**					**45733002**
Actinomyces naeslundii		67.87	0.38	ESL 12		
Actinomyces israelii		32.13	0.23	SUC 99		
45733003	**Good identification**					**45733003**
Actinomyces naeslundii		100.0	0.41	ESL 12		
45733020	**Low discrimination**					**45733020**
Actinomyces naeslundii		88.20	0.38	ESL 12		
Clostridium perfringens		11.81	0.01	SUC 99		
45733021	**Good identification**					**45733021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
45733022	**Low discrimination**					**45733022**
					GRA	
Actinomyces naeslundii		88.39	0.38	ESL 12	99	
Prevotella oralis		11.61	-.03	SUC 99	1	
45733023	**Good identification**					**45733023**
Actinomyces naeslundii		100.0	0.41	ESL 12		
45733100	**Low discrimination**					**45733100**
					GRA	
Actinomyces naeslundii		52.77	0.46	ESL 12	99	
Fusobacterium mortiferum		23.82	0.28	SUC 99	1	
Actinomyces israelii		23.41	0.31		99	
45733101	**Good identification**					**45733101**
Actinomyces naeslundii		100.0	0.49	ESL 12		
45733102	**Low discrimination**					**45733102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		37.06	0.46	ESL 12			
Actinomyces israelii		24.66	0.34	SUC 99			
Bifidobacterium adolescentis		20.37	0.26				
45733103	**Good identification**						**45733103**
Actinomyces naeslundii		100.0	0.49	ESL 12			
45733121	**Good identification**						**45733121**
Actinomyces naeslundii		100.0	0.49	ESL 12			
45733123	**Good identification**						**45733123**
Actinomyces naeslundii		100.0	0.49	ESL 12			
45733141	**Good identification**						**45733141**
Actinomyces naeslundii		100.0	0.36	ESL 12			
45733143	**Good identification**						**45733143**
Actinomyces naeslundii		100.0	0.36	ESL 12			
45733161	**Good identification**						**45733161**
Actinomyces naeslundii		100.0	0.36	ESL 12			
45733163	**Good identification**						**45733163**
Actinomyces naeslundii		100.0	0.36	ESL 12			
45733502	**Low discrimination**						**45733502**
Bifidobacterium adolescentis		87.11	0.40	SUC 99			
Actinomyces israelii		12.89	0.32	XYL 87			
45733512	**Good identification**						**45733512**
Bifidobacterium adolescentis		96.62	0.42	SUC 99			
Actinomyces israelii		1.75	0.18	XYL 87			
Bifidobacterium dentium		1.63	-.07				
45733722	**Good identification**						**45733722**
					SPO		
Parabacteroides distasonis		96.96	0.36	MAN 1	1		
Clostridium ramosum		3.04	-.03	XYL 90	92		
45734100	**Good identification**						**45734100**
					COC	GRA	
Fusobacterium mortiferum		97.48	0.40	MLT 11	1	1	
Atopobium parvulum		2.53	0.02	XYL 7	99	99	
45734122	**Good identification**						**45734122**
					GRA		
Parabacteroides distasonis		97.88	0.47	GAL 99	1		
Prevotella oralis		1.26	0.02		1		
Bifidobacterium longum subsp. infantis		0.86	-.04		99		
45734132	**Good identification**						**45734132**
Parabacteroides distasonis		96.03	0.43	GAL 99			
Clostridium clostridioforme		3.97	0.13				
45734322	**Good identification**						**45734322**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		98.82	0.56	GAL 99				
Clostridium clostridioforme		1.18	0.17					
45734332	**Acceptable identification**							**45734332**
Parabacteroides distasonis		90.82	0.51	GAL 99				
Clostridium clostridioforme		9.18	0.28					
45734522	**Low discrimination**							**45734522**
Parabacteroides distasonis		88.90	0.44	GAL 99				
Clostridium clostridioforme		11.10	0.23					
45734612	**Good identification**							**45734612**
Clostridium clostridioforme		100.0	0.40	SUC 99				
45734632	**Good identification**							**45734632**
Clostridium clostridioforme		98.71	0.40	SUC 99				
Parabacteroides distasonis		1.29	0.13					
45734702	**Good identification**							**45734702**
Clostridium clostridioforme		96.80	0.37	SUC 99				
Parabacteroides distasonis		3.20	0.18					
45734712	**Good identification**							**45734712**
Clostridium clostridioforme		99.61	0.49	SUC 99				
Parabacteroides distasonis		0.39	0.13					
45734722	**Low discrimination**							**45734722**
Parabacteroides distasonis		76.61	0.52	GAL 99				
Clostridium clostridioforme		23.39	0.37					
45734732	**Low discrimination**							**45734732**
Clostridium clostridioforme		72.09	0.49	SUC 99				
Parabacteroides distasonis		27.91	0.47					
45735000	**Low discrimination**							**45735000**
					COC	GRA		
Fusobacterium mortiferum		83.59	0.37	MLT 11	1	1		
Actinomyces israelii		13.05	0.26	XYL 7	1	99		
Atopobium parvulum		3.37	0.02		99	99		
45735022	**Acceptable identification**							**45735022**
					GRA			
Parabacteroides distasonis		91.75	0.47	LAC 99	1			
Prevotella oralis		7.44	0.16		1			
Bifidobacterium longum subsp. infantis		0.81	-.04		99			
45735032	**Good identification**							**45735032**
Parabacteroides distasonis		96.43	0.43	LAC 99				
Clostridium clostridioforme		3.57	0.13					
45735100	**Low discrimination**							**45735100**
					COC	GRA	SPO	
Atopobium parvulum		50.34	0.36	SUC 96	99	99	1	
Fusobacterium mortiferum		29.44	0.43	RAF 1	1	1	1	
Clostridium butyricum		8.38	0.15		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
45735102	**Low discrimination**							**45735102**
					GRA			
Parabacteroides distasonis		37.54	0.47	NAG 99	1			
Bifidobacterium adolescentis		24.23	0.42		99			
Bifidobacterium longum subsp. infantis		18.40	0.26		99			
45735112	**Low discrimination**							**45735112**
					GRA			
Bifidobacterium adolescentis		55.16	0.43	SUC 99	99			
Parabacteroides distasonis		37.65	0.43	ARA 87	1			
Clostridium clostridioforme		4.67	0.22		50			
45735120	**Low discrimination**							**45735120**
					COC	GRA	SPO	
Parabacteroides distasonis		57.10	0.47	BGL 99	1	1	1	
Clostridium septicum		18.94	0.23		1	99	99	
Atopobium parvulum		8.32	0.23		99	99	1	
45735122	**Very good identification**							**45735122**
					GRA			
Parabacteroides distasonis		97.77	0.82		1			
Prevotella oralis		1.26	0.36		1			
Bifidobacterium longum subsp. infantis		0.86	0.30		99			
45735123	**Good identification**							**45735123**
					GRA			
Parabacteroides distasonis		97.88	0.47	URE 1	1			
Prevotella oralis		1.26	0.02		1			
Bifidobacterium longum subsp. infantis		0.86	-.04		99			
45735130	**Good identification**							**45735130**
Parabacteroides distasonis		100.0	0.43	BGL 99				
45735132	**Very good identification**							**45735132**
Parabacteroides distasonis		99.82	0.77					
Clostridium clostridioforme		0.13	0.22					
Prevotella oralis		0.02	0.02					
45735133	**Good identification**							**45735133**
Parabacteroides distasonis		100.0	0.43	URE 1				
45735162	**Good identification**							**45735162**
					GRA			
Parabacteroides distasonis		97.88	0.47	SOR 1	1			
Prevotella oralis		1.26	0.02		1			
Bifidobacterium longum subsp. infantis		0.86	-.04		99			
45735172	**Good identification**							**45735172**
Parabacteroides distasonis		100.0	0.43	SOR 1				
45735222	**Good identification**							**45735222**
Parabacteroides distasonis		97.82	0.56	LAC 99				
Prevotella oralis		1.13	0.10					
Clostridium clostridioforme		1.05	0.16					
45735232	**Acceptable identification**							**45735232**
Parabacteroides distasonis		91.70	0.51	LAC 99				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		8.30	0.27				
45735302	**Acceptable identification**						**45735302**
					GRA		
Parabacteroides distasonis		93.99	0.56	NAG 99	1		
Clostridium clostridioforme		3.37	0.25		50		
Eubacterium contortum		1.09	0.11		99		
45735312	**Low discrimination**						**45735312**
					GRA		
Parabacteroides distasonis		76.45	0.51	NAG 99	1		
Clostridium clostridioforme		23.17	0.36		50		
Bifidobacterium adolescentis		0.38	0.09		99		
45735320	**Low discrimination**						**45735320**
					GRA		
Parabacteroides distasonis		85.15	0.56	BGL 99	1		
Bacteroides vulgatus		13.05	0.23		1		
Eubacterium contortum		0.99	0.11		99		
45735322	**Very good identification**						**45735322**
Parabacteroides distasonis		99.78	0.90				
Prevotella oralis		0.18	0.30				
Clostridium clostridioforme		0.04	0.25				
45735323	**Good identification**						**45735323**
Parabacteroides distasonis		100.0	0.56	URE 1			
45735330	**Good identification**						**45735330**
Parabacteroides distasonis		100.0	0.51	BGL 99			
45735332	**Very good identification**						**45735332**
Parabacteroides distasonis		99.70	0.85				
Clostridium clostridioforme		0.31	0.36				
45735333	**Good identification**						**45735333**
Parabacteroides distasonis		100.0	0.51	URE 1			
45735362	**Good identification**						**45735362**
Parabacteroides distasonis		100.0	0.56	SOR 1			
45735372	**Good identification**						**45735372**
Parabacteroides distasonis		100.0	0.51	SOR 1			
45735422	**Low discrimination**						**45735422**
Parabacteroides distasonis		89.94	0.44	LAC 99			
Clostridium clostridioforme		10.06	0.22				
45735432	**Low discrimination**						**45735432**
Parabacteroides distasonis		51.37	0.39	LAC 99			
Clostridium clostridioforme		48.63	0.33				
45735502	**Low discrimination**						**45735502**
					GRA	SPO	
Bifidobacterium adolescentis		74.35	0.56	SUC 99	99	1	
Parabacteroides distasonis		10.55	0.44		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		8.28	0.25		99	99	
45735512	**Low discrimination**						**45735512**
					GRA		
Bifidobacterium adolescentis		78.46	0.57	SUC 99	99		
Clostridium clostridioforme		15.55	0.42		50		
Parabacteroides distasonis		4.91	0.39		1		
45735522	**Very good identification**						**45735522**
					GRA		
Parabacteroides distasonis		99.35	0.78		1		
Clostridium clostridioforme		0.38	0.31		50		
Bifidobacterium adolescentis		0.07	0.22		99		
45735523	**Good identification**						**45735523**
Parabacteroides distasonis		100.0	0.44	URE 1			
45735530	**Low discrimination**						**45735530**
					GRA		
Parabacteroides distasonis		82.45	0.39	BGL 99	1		
Bifidobacterium longum subsp. longum		13.20	0.21		99		
Clostridium clostridioforme		2.64	0.08		50		
45735532	**Very good identification**						**45735532**
					GRA		
Parabacteroides distasonis		96.54	0.73		1		
Clostridium clostridioforme		3.09	0.42		50		
Bifidobacterium adolescentis		0.16	0.23		99		
45735533	**Good identification**						**45735533**
Parabacteroides distasonis		96.90	0.39	URE 1			
Clostridium clostridioforme		3.10	0.08				
45735542	**Low discrimination**						**45735542**
					SPO		
Bifidobacterium adolescentis		82.95	0.40	SUC 99	1		
Actinomyces israelii		9.59	0.30	SOR 11	1		
Clostridium butyricum		6.50	0.07		99		
45735552	**Good identification**						**45735552**
Bifidobacterium adolescentis		97.06	0.42	SUC 99			
Clostridium clostridioforme		1.57	0.08	SOR 11			
Actinomyces israelii		1.37	0.16				
45735562	**Good identification**						**45735562**
Parabacteroides distasonis		100.0	0.44	SOR 1			
45735572	**Good identification**						**45735572**
					GRA		
Parabacteroides distasonis		95.06	0.39	SOR 1	1		
Clostridium clostridioforme		3.04	0.08		50		
Bifidobacterium adolescentis		1.90	0.08		99		
45735602	**Good identification**						**45735602**
Clostridium clostridioforme		96.44	0.37	SUC 99			
Parabacteroides distasonis		3.56	0.18				
45735612	**Good identification**						**45735612**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		99.57	0.48	SUC 99		
Parabacteroides distasonis		0.44	0.13			
45735622	**Low discrimination**					**45735622**
Parabacteroides distasonis		78.52	0.52	LAC 99		
Clostridium clostridioforme		21.48	0.37			
45735632	**Low discrimination**					**45735632**
Clostridium clostridioforme		69.83	0.48	SUC 99		
Parabacteroides distasonis		30.17	0.47			
45735702	**Low discrimination**					**45735702**
					GRA	
Parabacteroides distasonis		50.77	0.52	NAG 99	1	
Clostridium clostridioforme		46.50	0.46		50	
Bifidobacterium adolescentis		1.21	0.22		99	
45735712	**Low discrimination**					**45735712**
					GRA	
Clostridium clostridioforme		88.03	0.57	SUC 99	50	
Parabacteroides distasonis		11.36	0.47		1	
Bifidobacterium adolescentis		0.61	0.23		99	
45735720	**Low discrimination**					**45735720**
					GRA	
Bacteroides vulgatus		76.43	0.42	SUC 96	1	
Parabacteroides distasonis		23.01	0.52	TRE 4	1	
Eubacterium contortum		0.36	0.10		99	
45735722	**Very good identification**					**45735722**
Parabacteroides distasonis		98.94	0.86			
Clostridium clostridioforme		0.92	0.46			
Bacteroides thetaiotaomicron		0.10	0.18			
45735723	**Good identification**					**45735723**
Parabacteroides distasonis		99.08	0.52	URE 1		
Clostridium clostridioforme		0.92	0.11			
45735730	**Low discrimination**					**45735730**
Parabacteroides distasonis		87.68	0.47	BGL 99		
Clostridium clostridioforme		6.86	0.23			
Bacteroides vulgatus		5.46	0.08			
45735732	**Good identification**					**45735732**
Parabacteroides distasonis		92.66	0.82			
Clostridium clostridioforme		7.25	0.57			
Bacteroides thetaiotaomicron		0.06	0.11			
45735733	**Acceptable identification**					**45735733**
Parabacteroides distasonis		92.74	0.47	URE 1		
Clostridium clostridioforme		7.26	0.23			
45735762	**Good identification**					**45735762**
Parabacteroides distasonis		99.08	0.52	SOR 1		
Clostridium clostridioforme		0.92	0.11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45735772	**Acceptable identification**						**45735772**
Parabacteroides distasonis		92.74	0.47	SOR 1			
Clostridium clostridioforme		7.26	0.23				
45737000	**Good identification**						**45737000**
Actinomyces israelii		100.0	0.39	SUC 98			
45737002	**Good identification**						**45737002**
Actinomyces israelii		100.0	0.42	SUC 98			
45737100	**Good identification**						**45737100**
					COC	SPO	
Actinomyces israelii		96.88	0.49	SUC 98	1	1	
Clostridium butyricum		2.02	-.01	CEL 93	1	99	
Atopobium parvulum		1.10	0.02		99	1	
45737102	**Low discrimination**						**45737102**
					GRA		
Actinomyces israelii		67.12	0.52	SUC 98	99		
Bifidobacterium adolescentis		32.26	0.40	CEL 93	99		
Parabacteroides distasonis		0.62	0.13		1		
45737112	**Low discrimination**						**45737112**
Bifidobacterium adolescentis		78.66	0.42	SUC 99			
Actinomyces israelii		20.01	0.38	ARA 87			
Bifidobacterium dentium		1.33	-.07				
45737122	**Acceptable identification**						**45737122**
					GRA		
Parabacteroides distasonis		91.60	0.47	MAN 1	1		
Actinomyces israelii		6.42	0.31		99		
Prevotella oralis		1.18	0.02		1		
45737132	**Acceptable identification**						**45737132**
					GRA		
Parabacteroides distasonis		93.67	0.43	MAN 1	1		
Bifidobacterium dentium		3.37	-.03		99		
Actinomyces israelii		1.82	0.17		99		
45737140	**Good identification**						**45737140**
Actinomyces israelii		100.0	0.42	SUC 98			
45737142	**Low discrimination**						**45737142**
Actinomyces israelii		86.75	0.45	SUC 98			
Bifidobacterium adolescentis		13.25	0.25	CEL 93			
45737322	**Good identification**						**45737322**
					SPO		
Parabacteroides distasonis		99.79	0.56	MAN 1	1		
Clostridium ramosum		0.21	-.03		92		
45737332	**Good identification**						**45737332**
Parabacteroides distasonis		100.0	0.51	MAN 1			
45737400	**Good identification**						**45737400**
Actinomyces israelii		100.0	0.37	SUC 98			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45737402	**Low discrimination**						**45737402**
Actinomyces israelii		86.29	0.40	SUC 98			
Bifidobacterium adolescentis		13.71	0.20	CEL 93			
45737500	**Low discrimination**						**45737500**
					SPO		
Actinomyces israelii		77.58	0.47	SUC 98	1		
Clostridium butyricum		17.80	0.15	CEL 93	99		
Bifidobacterium adolescentis		4.62	0.20		1		
45737502	**Low discrimination**						**45737502**
					SPO		
Bifidobacterium adolescentis		78.47	0.54	SUC 99	1		
Actinomyces israelii		19.96	0.50		1		
Clostridium butyricum		1.19	0.08		99		
45737512	**Low discrimination**						**45737512**
Bifidobacterium adolescentis		77.93	0.56	SUC 99			
Bifidobacterium dentium		19.46	0.27				
Actinomyces israelii		2.42	0.36				
45737522	**Low discrimination**						**45737522**
					GRA		
Parabacteroides distasonis		84.50	0.44	MAN 1	1		
Actinomyces israelii		7.91	0.30		99		
Bifidobacterium adolescentis		4.92	0.20		99		
45737542	**Low discrimination**						**45737542**
					COC		
Bifidobacterium adolescentis		52.89	0.39	SUC 99	1		
Actinomyces israelii		42.34	0.43	SOR 11	1		
Blautia producta		4.77	0.07		99		
45737552	**Low discrimination**						**45737552**
					COC		
Bifidobacterium adolescentis		74.65	0.40	SUC 99	1		
Blautia producta		16.52	0.15	SOR 11	99		
Actinomyces israelii		7.31	0.29		1		
45737722	**Good identification**						**45737722**
Parabacteroides distasonis		98.52	0.52	MAN 1			
Clostridium clostridioforme		0.91	0.11				
Bacteroides ovatus		0.57	-.02				
45737732	**Acceptable identification**						**45737732**
					GRA		
Parabacteroides distasonis		90.50	0.47	MAN 1	1		
Clostridium clostridioforme		7.08	0.23		50		
Bifidobacterium dentium		1.77	-.03		99		
45741122	**Acceptable identification**						**45741122**
					GRA	SPO	
Prevotella oralis		90.09	0.48	SUC 92	1	1	
Clostridium paraputrificum		6.75	0.22	FRU 99	99	99	
Bifidobacterium breve		2.34	0.32		99	1	
45741322	**Good identification**						**45741322**
					SPO		
Prevotella oralis		98.15	0.41	SUC 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		1.56	0.23	FRU 99	1		
Clostridium ramosum		0.28	-.02		92		
45745322	**Low discrimination**						**45745322**
Parabacteroides distasonis		53.71	0.40	FRU 99			
Prevotella oralis		46.29	0.26				
45745332	**Good identification**						**45745332**
Parabacteroides distasonis		100.0	0.35	FRU 99			
45745722	**Low discrimination**						**45745722**
					CAT		
Parabacteroides distasonis		77.75	0.36	FRU 99	76		
Bacteroides uniformis		11.08	0.03		5		
Bacteroides eggerthii		8.39	-.03		93		
45750000	**Good identification**						**45750000**
					GRA		
Fusobacterium mortiferum		99.31	0.54	MLT 11	1		
Lactobacillus jensenii		0.69	0.07		99		
45750002	**Low discrimination**						**45750002**
					GRA		
Fusobacterium mortiferum		88.74	0.38	MLT 11	1		
Prevotella oralis		7.57	0.05	BGL 10	1		
Lactobacillus jensenii		3.70	0.04		99		
45750100	**Very good identification**						**45750100**
					GRA		
Fusobacterium mortiferum		98.71	0.60	MLT 11	1		
Collinsella aerofaciens		0.58	-.03		99		
Prevotella oralis		0.33	0.04		1		
45750102	**Low discrimination**						**45750102**
					GRA		
Fusobacterium mortiferum		55.74	0.44	MLT 11	1		
Prevotella oralis		31.90	0.26	BGL 10	1		
Bifidobacterium breve		8.85	0.27		99		
45750122	**Low discrimination**						**45750122**
					GRA	SPO	
Prevotella oralis		89.63	0.48	SUC 92	1	1	
Clostridium paraputrificum		6.71	0.22	GAL 99	99	99	
Bifidobacterium breve		2.33	0.32		99	1	
45750322	**Good identification**						**45750322**
					SPO		
Prevotella oralis		98.15	0.41	SUC 92	1		
Parabacteroides distasonis		1.56	0.23	GAL 99	1		
Clostridium ramosum		0.28	-.02		92		
45751000	**Low discrimination**						**45751000**
					GRA		
Fusobacterium mortiferum		62.21	0.57	MLT 11	1		
Lactobacillus jensenii		28.50	0.41		99		
Lactobacillus catenaformis		5.16	0.29		99		
45751002	**Low discrimination**						**45751002**
					GRA		
Prevotella oralis		50.77	0.40	SUC 92	1		
Lactobacillus jensenii		24.80	0.38	LAC 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		9.02	0.41	NAG 95	1		
45751020	**Low discrimination**						**45751020**
					GRA	SPO	
Prevotella oralis		49.75	0.40	SUC 92	1	1	
Clostridium paraputrificum		26.14	0.29	LAC 94	99	99	
Lactobacillus jensenii		9.12	0.31	BGL 95	99	1	
45751022	**Very good identification**						**45751022**
					GRA	SPO	
Prevotella oralis		96.53	0.61	SUC 92	1	1	
Clostridium paraputrificum		1.14	0.22	LAC 94	99	99	
Clostridium barati		0.68	0.24		99	99	
45751100	**Low discrimination**						**45751100**
					GRA		
Fusobacterium mortiferum		49.78	0.63	MLT 11	1		
Collinsella aerofaciens		19.32	0.32		99		
Prevotella oralis		11.00	0.38		1		
45751101	**Low discrimination**						**45751101**
					GRA		
Actinomyces naeslundii		35.96	0.41	ESL 12	99		
Fusobacterium mortiferum		34.34	0.29	SUC 99	1		
Collinsella aerofaciens		13.33	-.03		99		
45751102	**Low discrimination**						**45751102**
					GRA		
Prevotella oralis		69.88	0.60	SUC 92	1		
Bifidobacterium breve		19.38	0.62	NAG 95	99		
Lactobacillus acidophilus		3.18	0.38		99		
45751110	**Low discrimination**						**45751110**
					GRA		
Bifidobacterium breve		86.05	0.44	SUC 99	99		
Fusobacterium mortiferum		7.55	0.29	BGL 91	1		
Collinsella aerofaciens		2.93	-.03		99		
45751112	**Low discrimination**						**45751112**
					GRA		
Bifidobacterium breve		85.40	0.62	SUC 99	99		
Leptotrichia buccalis		7.19	0.24		1		
Bifidobacterium adolescentis		3.85	0.38		99		
45751120	**Low discrimination**						**45751120**
					GRA	SPO	
Clostridium paraputrificum		59.68	0.63	SUC 92	99	99	
Clostridium septicum		20.72	0.58	RAF 7	99	99	
Prevotella oralis		17.97	0.60		1	1	
45751122	**Low discrimination**						**45751122**
					GRA	SPO	
Prevotella oralis		89.96	0.82	SUC 92	1	1	
Clostridium paraputrificum		6.74	0.56		99	99	
Bifidobacterium breve		2.34	0.66		99	1	
45751123	**Low discrimination**						**45751123**
					GRA	SPO	
Prevotella oralis		89.04	0.48	SUC 92	1	1	
Clostridium paraputrificum		6.67	0.22	URE 1	99	99	
Bifidobacterium breve		2.31	0.32		99	1	
45751130	**Low discrimination**						**45751130**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium breve		46.76	0.49	SUC 99	1		
Clostridium paraputrificum		32.16	0.29	BGL 91	99		
Clostridium septicum		11.17	0.24		99		
45751132	**Low discrimination**						**45751132**
					GRA		
Bifidobacterium breve		67.83	0.66	SUC 99	99		
Prevotella oralis		26.40	0.48		1		
Parabacteroides distasonis		3.20	0.45		1		
45751140	**Low discrimination**						**45751140**
					GRA	SPO	
Bifidobacterium breve		78.63	0.44	SUC 99	99	1	
Clostridium paraputrificum		7.24	0.09	BGL 91	99	99	
Fusobacterium mortiferum		6.90	0.29		1	1	
45751142	**Good identification**						**45751142**
					GRA		
Bifidobacterium breve		93.16	0.62	SUC 99	99		
Prevotella oralis		3.39	0.26		1		
Lactobacillus acidophilus		1.89	0.23		99		
45751150	**Good identification**						**45751150**
Bifidobacterium breve		100.0	0.44	SUC 99			
45751152	**Very good identification**						**45751152**
					GRA		
Bifidobacterium breve		99.36	0.62	SUC 99	99		
Bifidobacterium adolescentis		0.55	0.22		99		
Leptotrichia buccalis		0.08	-.11		1		
45751160	**Low discrimination**						**45751160**
					SPO		
Clostridium paraputrificum		77.58	0.43	SUC 92	99		
Bifidobacterium breve		15.14	0.49	RAF 7	1		
Clostridium septicum		3.62	0.24	SOR 7	99		
45751162	**Low discrimination**						**45751162**
					GRA	SPO	
Bifidobacterium breve		61.48	0.66	SUC 99	99	1	
Prevotella oralis		23.93	0.48		1	1	
Clostridium paraputrificum		13.36	0.37		99	99	
45751170	**Good identification**						**45751170**
					SPO		
Bifidobacterium breve		95.08	0.49	SUC 99	1		
Clostridium paraputrificum		4.92	0.09	BGL 91	99		
45751172	**Very good identification**						**45751172**
					GRA	SPO	
Bifidobacterium breve		99.34	0.66	SUC 99	99	1	
Prevotella oralis		0.39	0.14		1	1	
Clostridium paraputrificum		0.22	0.03		99	99	
45751222	**Good identification**						**45751222**
					GRA	SPO	
Prevotella oralis		99.50	0.55	SUC 92	1	1	
Parabacteroides distasonis		0.25	0.23	LAC 94	1	1	
Clostridium barati		0.14	0.05		99	99	
45751302	**Good identification**						**45751302**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Prevotella oralis		97.95	0.54	SUC 92	1		
Lactobacillus acidophilus		0.78	0.19	NAG 95	99		
Bifidobacterium breve		0.64	0.27		99		
45751320	**Low discrimination**						**45751320**
					GRA	SPO	
Prevotella oralis		88.81	0.54	SUC 92	1	1	
Clostridium paraputrificum		6.95	0.29	BGL 95	99	99	
Clostridium septicum		2.41	0.24		99	99	
45751322	**Very good identification**						**45751322**
					SPO		
Prevotella oralis		97.85	0.76	SUC 92	1		
Parabacteroides distasonis		1.56	0.58		1		
Clostridium ramosum		0.28	0.33		92		
45751323	**Good identification**						**45751323**
					SPO		
Prevotella oralis		98.15	0.41	SUC 92	1		
Parabacteroides distasonis		1.56	0.23	URE 1	1		
Clostridium ramosum		0.28	-.02		92		
45751332	**Low discrimination**						**45751332**
					GRA		
Prevotella oralis		52.03	0.41	SUC 92	1		
Parabacteroides distasonis		44.20	0.53	MLZ 1	1		
Bifidobacterium breve		3.15	0.32		99		
45751362	**Acceptable identification**						**45751362**
					GRA		
Prevotella oralis		91.10	0.41	SUC 92	1		
Bifidobacterium breve		5.52	0.32	SOR 1	99		
Parabacteroides distasonis		1.45	0.23		1		
45751502	**Low discrimination**						**45751502**
					GRA	SPO	
Bifidobacterium adolescentis		78.10	0.50	SUC 99	99	1	
Prevotella oralis		11.53	0.26	XYL 87	1	1	
Clostridium butyricum		3.57	0.13		99	99	
45751512	**Acceptable identification**						**45751512**
Bifidobacterium adolescentis		94.52	0.52	SUC 99			
Bifidobacterium breve		3.17	0.27	XYL 87			
Clostridium clostridioforme		1.78	0.19				
45751522	**Low discrimination**						**45751522**
					GRA	SPO	
Prevotella oralis		79.38	0.48	SUC 92	1	1	
Parabacteroides distasonis		10.97	0.46	ARA 1	1	1	
Clostridium paraputrificum		5.95	0.22		99	99	
45751532	**Low discrimination**						**45751532**
					GRA		
Parabacteroides distasonis		60.46	0.41	XYL 90	1		
Bifidobacterium breve		21.09	0.32		99		
Prevotella oralis		8.21	0.14		1		
45751552	**Low discrimination**						**45751552**
Bifidobacterium adolescentis		78.68	0.36	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		21.32	0.27	XYL 87		
45751722	**Low discrimination**					**45751722**
Prevotella oralis		49.67	0.41	SUC 92		
Parabacteroides distasonis		48.03	0.54	ARA 1		
Clostridium clostridioforme		1.53	0.23			
45751732	**Low discrimination**					**45751732**
Parabacteroides distasonis		77.61	0.49	XYL 90		
Clostridium clostridioforme		20.88	0.34			
Prevotella oralis		1.51	0.07			
45753001	**Good identification**					**45753001**
Actinomyces naeslundii		100.0	0.38	ESL 12		
45753003	**Good identification**					**45753003**
Actinomyces naeslundii		100.0	0.38	ESL 12		
45753020	**Low discrimination**					**45753020**
					GRA	
Actinomyces naeslundii		36.68	0.35	ESL 12	99	
Prevotella oralis		20.64	0.05	SUC 99	1	
Bifidobacterium breve		15.77	0.15		99	
45753021	**Good identification**					**45753021**
Actinomyces naeslundii		100.0	0.38	ESL 12		
45753023	**Good identification**					**45753023**
Actinomyces naeslundii		100.0	0.38	ESL 12		
45753100	**Low discrimination**					**45753100**
Bifidobacterium breve		63.96	0.44	SUC 99		
Actinomyces israelii		17.45	0.40	BGL 91		
Actinomyces naeslundii		7.61	0.43			
45753101	**Acceptable identification**					**45753101**
Actinomyces naeslundii		94.64	0.46	ESL 12		
Bifidobacterium breve		5.36	0.10	SUC 99		
45753102	**Low discrimination**					**45753102**
					GRA	
Bifidobacterium breve		88.00	0.62	SUC 99	99	
Actinomyces israelii		3.56	0.43		99	
Prevotella oralis		3.21	0.26		1	
45753103	**Low discrimination**					**45753103**
Actinomyces naeslundii		63.59	0.46	ESL 12		
Bifidobacterium breve		36.41	0.27	SUC 99		
45753110	**Good identification**					**45753110**
Bifidobacterium breve		96.08	0.44	SUC 99		
Actinomyces israelii		3.92	0.26	BGL 91		
45753112	**Very good identification**					**45753112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		95.66	0.62	SUC 99			
Bifidobacterium adolescentis		3.53	0.36				
Actinomyces israelii		0.58	0.29				
45753120	**Low discrimination**						**45753120**
					SPO		
Bifidobacterium breve		36.41	0.49	SUC 99	1		
Clostridium paraputrificum		25.04	0.29	BGL 91	99		
Clostridium ramosum		18.78	0.33		92		
45753121	**Low discrimination**						**45753121**
					SPO		
Actinomyces naeslundii		81.84	0.46	ESL 12	1		
Bifidobacterium breve		8.24	0.15	SUC 99	1		
Clostridium paraputrificum		5.67	-.06		99		
45753122	**Low discrimination**						**45753122**
					SPO	GRA	
Bifidobacterium breve		60.85	0.66	SUC 99	1	99	
Prevotella oralis		23.68	0.48		1	1	
Clostridium ramosum		12.42	0.43		92	50	
45753130	**Good identification**						**45753130**
					SPO		
Bifidobacterium breve		97.91	0.49	SUC 99	1		
Clostridium tertium		0.90	0.05	BGL 91	99		
Clostridium paraputrificum		0.68	-.06		99		
45753132	**Very good identification**						**45753132**
					GRA	SPO	
Bifidobacterium breve		99.09	0.66	SUC 99	99	1	
Prevotella oralis		0.39	0.14		1	1	
Clostridium tertium		0.27	0.13		99	99	
45753140	**Low discrimination**						**45753140**
Bifidobacterium breve		88.73	0.44	SUC 99			
Actinomyces israelii		9.41	0.33	BGL 91			
Actinomyces naeslundii		1.86	0.30				
45753142	**Very good identification**						**45753142**
Bifidobacterium breve		97.79	0.62	SUC 99			
Actinomyces israelii		1.54	0.36				
Bifidobacterium adolescentis		0.37	0.19				
45753150	**Good identification**						**45753150**
Bifidobacterium breve		98.44	0.44	SUC 99			
Actinomyces israelii		1.56	0.19	BGL 91			
45753152	**Very good identification**						**45753152**
Bifidobacterium breve		99.31	0.62	SUC 99			
Bifidobacterium adolescentis		0.45	0.21				
Actinomyces israelii		0.23	0.22				
45753160	**Acceptable identification**						**45753160**
					SPO		
Bifidobacterium breve		93.56	0.49	SUC 99	1		
Clostridium paraputrificum		4.84	0.09	BGL 91	99		
Actinomyces naeslundii		1.10	0.30		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45753162	**Very good identification**						**45753162**
					GRA	SPO	
Bifidobacterium breve		98.96	0.66	SUC 99	99	1	
Prevotella oralis		0.39	0.14		1	1	
Clostridium paraputrificum		0.22	0.03		99	99	
45753170	**Good identification**						**45753170**
Bifidobacterium breve		100.0	0.49	SUC 99			
45753172	**Very good identification**						**45753172**
Bifidobacterium breve		100.0	0.66	SUC 99			
45753320	**Low discrimination**						**45753320**
					SPO	GRA	
Clostridium ramosum		86.60	0.35	SUC 99	92	50	
Prevotella oralis		11.24	0.20	TRE 94	1	1	
Bifidobacterium breve		1.28	0.15		1	99	
45753322	**Low discrimination**						**45753322**
					SPO	GRA	
Clostridium ramosum		60.10	0.45	SUC 99	92	50	
Prevotella oralis		37.07	0.41	TRE 94	1	1	
Bifidobacterium breve		2.24	0.32		1	99	
45753500	**Low discrimination**						**45753500**
					SPO		
Actinomyces israelii		77.90	0.39	SUC 98	1		
Clostridium butyricum		11.37	0.03	XYL 92	99		
Bifidobacterium adolescentis		7.20	0.15		1		
45753502	**Low discrimination**						**45753502**
Bifidobacterium adolescentis		81.95	0.49	SUC 99			
Actinomyces israelii		13.44	0.42	XYL 87			
Bifidobacterium breve		4.10	0.27				
45753512	**Acceptable identification**						**45753512**
Bifidobacterium adolescentis		93.78	0.50	SUC 99			
Bifidobacterium breve		3.84	0.27	XYL 87			
Actinomyces israelii		1.88	0.27				
45754100	**Good identification**						**45754100**
					GRA	SPO	
Fusobacterium mortiferum		97.62	0.41	MLT 11	1	1	
Clostridium butyricum		2.38	-.05	XYL 7	99	99	
45754322	**Low discrimination**						**45754322**
Parabacteroides distasonis		52.41	0.40	GAL 99			
Prevotella oralis		45.17	0.26				
Clostridium clostridioforme		2.42	0.11				
45754332	**Low discrimination**						**45754332**
Parabacteroides distasonis		71.93	0.35	GAL 99			
Clostridium clostridioforme		28.07	0.22				
45754712	**Good identification**						**45754712**
Clostridium clostridioforme		100.0	0.43	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45754732	**Acceptable identification**						**45754732**
Clostridium clostridioforme		90.89	0.43	SUC 99			
Parabacteroides distasonis		9.11	0.31	TRE 88			
45755000	**Low discrimination**						**45755000**
					GRA	SPO	
Fusobacterium mortiferum		56.62	0.38	MLT 11	1	1	
Actinomyces israelii		27.97	0.35	XYL 7	99	1	
Clostridium butyricum		8.87	0.05		99	99	
45755022	**Good identification**						**45755022**
					GRA		
Prevotella oralis		96.83	0.46	SUC 92	1		
Parabacteroides distasonis		2.54	0.32	XYL 11	1		
Actinomyces israelii		0.24	0.18	LAC 94	99		
45755102	**Low discrimination**						**45755102**
					GRA		
Prevotella oralis		49.03	0.44	SUC 92	1		
Bifidobacterium adolescentis		27.15	0.50	XYL 11	99		
Actinomyces israelii		9.04	0.48	NAG 95	99		
45755112	**Low discrimination**						**45755112**
Bifidobacterium adolescentis		87.99	0.52	SUC 99			
Actinomyces israelii		3.58	0.34	ARA 87			
Bifidobacterium breve		2.95	0.27				
45755120	**Low discrimination**						**45755120**
					GRA	SPO	
Prevotella oralis		67.40	0.44	SUC 92	1	1	
Clostridium paraputrificum		18.29	0.29	XYL 11	99	99	
Clostridium septicum		6.35	0.24	BGL 95	99	99	
45755122	**Low discrimination**						**45755122**
					CAT		
Prevotella oralis		83.00	0.66	SUC 92	1		
Parabacteroides distasonis		13.76	0.66	XYL 11	76		
Bacteroides fragilis		1.13	0.25		99		
45755132	**Low discrimination**						**45755132**
					GRA		
Parabacteroides distasonis		86.98	0.61		1		
Prevotella oralis		9.84	0.32		1		
Bifidobacterium breve		2.07	0.32		99		
45755140	**Low discrimination**						**45755140**
					SPO		
Actinomyces israelii		67.91	0.38	SUC 98	1		
Clostridium butyricum		28.90	0.11		99		
Bifidobacterium breve		3.19	0.10		1		
45755142	**Low discrimination**						**45755142**
Actinomyces israelii		39.65	0.41	SUC 98			
Bifidobacterium adolescentis		37.85	0.35				
Bifidobacterium breve		12.54	0.27				
45755152	**Low discrimination**						**45755152**
Bifidobacterium adolescentis		71.48	0.36	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		19.37	0.27	ARA 87		
Actinomyces israelii		9.15	0.27	SOR 11		
45755222	**Low discrimination**					**45755222**
Prevotella oralis		83.95	0.40	SUC 92		
Parabacteroides distasonis		15.42	0.40	XYL 11		
Clostridium clostridioforme		0.64	0.10	LAC 94		
45755232	**Low discrimination**					**45755232**
Parabacteroides distasonis		68.88	0.35	LAC 99		
Clostridium clostridioforme		24.09	0.22			
Prevotella oralis		7.04	0.05			
45755302	**Low discrimination**					**45755302**
					GRA	
Prevotella oralis		75.34	0.38	SUC 92	1	
Parabacteroides distasonis		16.78	0.40	XYL 11	1	
Actinomyces israelii		2.44	0.29	NAG 95	99	
45755320	**Low discrimination**					**45755320**
Prevotella oralis		61.89	0.38	SUC 92		
Bacteroides vulgatus		22.58	0.25	XYL 11		
Parabacteroides distasonis		13.78	0.40	BGL 95		
45755322	**Low discrimination**					**45755322**
Parabacteroides distasonis		53.42	0.74			
Prevotella oralis		46.04	0.60			
Bacteroides uniformis		0.41	0.19			
45755323	**Low discrimination**					**45755323**
Parabacteroides distasonis		53.71	0.40	URE 1		
Prevotella oralis		46.29	0.26			
45755330	**Acceptable identification**					**45755330**
Parabacteroides distasonis		92.23	0.35	BGL 99		
Prevotella oralis		7.77	0.04			
45755332	**Very good identification**					**45755332**
Parabacteroides distasonis		97.25	0.69			
Prevotella oralis		1.57	0.26			
Clostridium clostridioforme		1.15	0.31			
45755333	**Good identification**					**45755333**
Parabacteroides distasonis		100.0	0.35	URE 1		
45755362	**Low discrimination**					**45755362**
Parabacteroides distasonis		53.71	0.40	SOR 1		
Prevotella oralis		46.29	0.26			
45755372	**Good identification**					**45755372**
Parabacteroides distasonis		100.0	0.35	SOR 1		
45755402	**Low discrimination**					**45755402**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		32.04	0.37	SUC 98	1	
Bifidobacterium adolescentis		31.81	0.30		1	
Clostridium butyricum		28.98	0.15		99	
45755500	**Acceptable identification**					**45755500**
					SPO	
Clostridium butyricum		93.75	0.45	SUC 99	99	
Actinomyces israelii		4.48	0.44	TRE 94	1	
Bifidobacterium adolescentis		1.67	0.30		1	
45755502	**Low discrimination**					**45755502**
					SPO	
Bifidobacterium adolescentis		78.20	0.64	SUC 99	1	
Clostridium butyricum		17.27	0.38		99	
Actinomyces israelii		3.18	0.47		1	
45755512	**Very good identification**					**45755512**
					SPO	
Bifidobacterium adolescentis		95.62	0.66	SUC 99	1	
Clostridium clostridioforme		2.71	0.36		66	
Clostridium butyricum		0.91	0.17		99	
45755522	**Low discrimination**					**45755522**
Parabacteroides distasonis		67.07	0.62			
Bacteroides uniformis		20.77	0.35			
Prevotella oralis		6.67	0.32			
45755532	**Low discrimination**					**45755532**
					GRA	
Parabacteroides distasonis		84.26	0.57		1	
Clostridium clostridioforme		10.42	0.36		50	
Bifidobacterium adolescentis		3.71	0.32		99	
45755542	**Low discrimination**					**45755542**
					SPO	
Bifidobacterium adolescentis		77.92	0.49	SUC 99	1	
Clostridium butyricum		12.11	0.20	SOR 11	99	
Actinomyces israelii		9.98	0.40		1	
45755552	**Good identification**					**45755552**
					SPO	
Bifidobacterium adolescentis		97.81	0.50	SUC 99	1	
Actinomyces israelii		1.53	0.26	SOR 11	1	
Clostridium butyricum		0.66	-.02		99	
45755612	**Good identification**					**45755612**
Clostridium clostridioforme		100.0	0.42	SUC 99		
45755632	**Low discrimination**					**45755632**
Clostridium clostridioforme		89.94	0.42	SUC 99		
Parabacteroides distasonis		10.06	0.31	TRE 88		
45755702	**Low discrimination**					**45755702**
					GRA	
Clostridium clostridioforme		57.51	0.40	SUC 99	50	
Parabacteroides distasonis		16.26	0.36	TRE 88	1	
Bifidobacterium adolescentis		10.40	0.30		99	
45755712	**Acceptable identification**					**45755712**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		92.27	0.51	SUC 99	50	
Bifidobacterium adolescentis		4.48	0.32	TRE 88	99	
Parabacteroides distasonis		3.08	0.31		1	
45755720	**Good identification**					**45755720**
Bacteroides vulgatus		96.00	0.44	SUC 96		
Parabacteroides distasonis		2.70	0.36	CEL 5		
Bacteroides uniformis		0.39	0.03			
45755722	**Low discrimination**					**45755722**
Parabacteroides distasonis		82.48	0.70			
Bacteroides uniformis		11.76	0.37			
Clostridium clostridioforme		2.95	0.40			
45755723	**Low discrimination**					**45755723**
Parabacteroides distasonis		84.87	0.36	URE 1		
Bacteroides uniformis		12.10	0.03			
Clostridium clostridioforme		3.03	0.06			
45755732	**Low discrimination**					**45755732**
Parabacteroides distasonis		75.53	0.66			
Clostridium clostridioforme		22.83	0.51			
Bacteroides ovatus		1.36	0.20			
45755762	**Low discrimination**					**45755762**
Parabacteroides distasonis		80.75	0.36	SOR 1		
Bacteroides uniformis		11.51	0.03			
Bacteroides ovatus		4.86	-.01			
45757000	**Good identification**					**45757000**
Actinomyces israelii		100.0	0.48	SUC 98		
45757002	**Good identification**					**45757002**
Actinomyces israelii		98.28	0.51	SUC 98		
Bifidobacterium adolescentis		1.72	0.15			
45757012	**Low discrimination**					**45757012**
Actinomyces israelii		87.46	0.37	SUC 98		
Bifidobacterium adolescentis		12.54	0.16	MLZ 13		
45757040	**Good identification**					**45757040**
Actinomyces israelii		100.0	0.41	SUC 98		
45757042	**Good identification**					**45757042**
Actinomyces israelii		100.0	0.44	SUC 98		
45757100	**Very good identification**					**45757100**
					SPO	
Actinomyces israelii		95.50	0.58	SUC 98	1	
Clostridium butyricum		3.56	0.13		99	
Bifidobacterium adolescentis		0.63	0.15		1	
45757102	**Low discrimination**					**45757102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		67.68	0.61	SUC 98		
Bifidobacterium adolescentis		29.36	0.49			
Bifidobacterium breve		1.47	0.27			
45757110	**Acceptable identification**					**45757110**
Actinomyces israelii		91.63	0.44	SUC 98		
Bifidobacterium adolescentis		4.93	0.16	MLZ 13		
Bifidobacterium breve		1.98	0.10			
45757112	**Low discrimination**					**45757112**
Bifidobacterium adolescentis		74.79	0.50	SUC 99		
Actinomyces israelii		21.08	0.47	ARA 87		
Bifidobacterium breve		3.06	0.27			
45757120	**Low discrimination**					**45757120**
					GRA	
Actinomyces israelii		64.97	0.38	SUC 98	99	
Prevotella oralis		14.77	0.10	NAG 6	1	
Bifidobacterium breve		5.83	0.15		99	
45757140	**Good identification**					**45757140**
					SPO	
Actinomyces israelii		98.37	0.51	SUC 98	1	
Clostridium butyricum		0.82	-.05		99	
Bifidobacterium breve		0.82	0.10		1	
45757142	**Low discrimination**					**45757142**
Actinomyces israelii		83.78	0.54	SUC 98		
Bifidobacterium adolescentis		11.55	0.33			
Bifidobacterium breve		4.68	0.27			
45757150	**Low discrimination**					**45757150**
					COC	
Actinomyces israelii		88.74	0.37	SUC 98	1	
Blautia producta		6.34	-.03	MLZ 13	99	
Bifidobacterium breve		4.92	0.10		1	
45757300	**Good identification**					**45757300**
Actinomyces israelii		100.0	0.39	SUC 98		
45757302	**Low discrimination**					**45757302**
					GRA	
Actinomyces israelii		89.82	0.42	SUC 98	99	
Bifidobacterium adolescentis		5.23	0.15	RHA 7	99	
Prevotella oralis		4.95	0.04		1	
45757322	**Low discrimination**					**45757322**
					SPO	
Parabacteroides distasonis		49.30	0.40	MAN 1	1	
Prevotella oralis		42.48	0.26		1	
Clostridium ramosum		5.63	0.11		92	
45757332	**Good identification**					**45757332**
Parabacteroides distasonis		100.0	0.35	MAN 1		
45757342	**Good identification**					**45757342**
Actinomyces israelii		100.0	0.35	SUC 98		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45757400	**Acceptable identification**						**45757400**
					SPO		
Actinomyces israelii		93.60	0.46	SUC 98	1		
Clostridium butyricum		6.40	0.05		99		
45757402	**Low discrimination**						**45757402**
					SPO		
Actinomyces israelii		86.13	0.49	SUC 98	1		
Bifidobacterium adolescentis		12.35	0.29		1		
Clostridium butyricum		1.53	-.02		99		
45757440	**Good identification**						**45757440**
Actinomyces israelii		100.0	0.39	SUC 98			
45757442	**Good identification**						**45757442**
Actinomyces israelii		95.64	0.42	SUC 98			
Bifidobacterium adolescentis		4.36	0.13				
45757500	**Low discrimination**						**45757500**
					SPO		
Actinomyces israelii		68.30	0.57	SUC 98	1		
Clostridium butyricum		28.03	0.29		99		
Bifidobacterium adolescentis		3.67	0.29		1		
45757502	**Low discrimination**						**45757502**
					SPO		
Bifidobacterium adolescentis		76.17	0.63	SUC 99	1		
Actinomyces israelii		21.47	0.60		1		
Clostridium butyricum		2.29	0.22		99		
45757510	**Low discrimination**						**45757510**
Actinomyces israelii		51.14	0.43	SUC 98			
Bifidobacterium adolescentis		22.49	0.30	MLZ 13			
Bifidobacterium dentium		17.25	0.10				
45757512	**Low discrimination**						**45757512**
Bifidobacterium adolescentis		89.76	0.64	SUC 99			
Bifidobacterium dentium		7.03	0.27				
Actinomyces israelii		3.09	0.46				
45757520	**Low discrimination**						**45757520**
					GRA	SPO	
Actinomyces israelii		61.60	0.36	SUC 98	99	1	
Clostridium butyricum		34.45	0.11	NAG 6	99	99	
Bacteroides vulgatus		3.95	-.06		1	1	
45757522	**Low discrimination**						**45757522**
					GRA		
Actinomyces israelii		43.37	0.39	SUC 98	99		
Bifidobacterium adolescentis		24.35	0.29	NAG 6	99		
Parabacteroides distasonis		15.51	0.28		1		
45757540	**Low discrimination**						**45757540**
					COC	SPO	
Actinomyces israelii		64.96	0.50	SUC 98	1	1	
Blautia producta		27.97	0.24		99	1	
Clostridium butyricum		5.96	0.11		1	99	
45757542	**Low discrimination**						**45757542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Bifidobacterium adolescentis		51.65	0.47	SUC 99	1		
Actinomyces israelii		45.80	0.53	SOR 11	1		
Blautia producta		1.46	0.07		99		
45757552	**Low discrimination**						**45757552**
					COC		
Bifidobacterium adolescentis		84.44	0.49	SUC 99	1		
Actinomyces israelii		9.16	0.39	SOR 11	1		
Blautia producta		5.86	0.15		99		
45757700	**Low discrimination**						**45757700**
					SPO		
Actinomyces israelii		85.57	0.38	SUC 98	1		
Clostridium butyricum		14.43	0.03	RHA 7	99		
45757702	**Low discrimination**						**45757702**
					SPO		
Actinomyces israelii		64.42	0.41	SUC 98	1		
Bifidobacterium adolescentis		30.67	0.29	RHA 7	1		
Clostridium butyricum		2.82	-.04		99		
45757722	**Low discrimination**						**45757722**
Parabacteroides distasonis		57.82	0.36	MAN 1			
Bacteroides ovatus		27.82	0.15				
Bacteroides uniformis		8.24	0.03				
45765122	**Acceptable identification**						**45765122**
Parabacteroides distasonis		92.21	0.43	FRU 99			
Prevotella oralis		7.79	0.12				
45765132	**Good identification**						**45765132**
Parabacteroides distasonis		100.0	0.38	FRU 99			
45765322	**Good identification**						**45765322**
Parabacteroides distasonis		98.81	0.51	FRU 99			
Prevotella oralis		1.19	0.05				
45765332	**Good identification**						**45765332**
Parabacteroides distasonis		98.13	0.46	FRU 99			
Clostridium clostridioforme		1.87	0.11				
45765522	**Low discrimination**						**45765522**
					GRA	SPO	
Parabacteroides distasonis		86.93	0.39	FRU 99	1	1	
Clostridium butyricum		11.02	0.07		99	99	
Clostridium clostridioforme		2.05	0.06		50	66	
45765722	**Acceptable identification**						**45765722**
Parabacteroides distasonis		91.75	0.47	FRU 99			
Clostridium clostridioforme		5.28	0.20				
Bacteroides ovatus		2.97	0.06				
45765732	**Low discrimination**						**45765732**
Parabacteroides distasonis		65.50	0.43	FRU 99			
Clostridium clostridioforme		31.87	0.32				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides ovatus		2.62	0.03					
45767102	**Acceptable identification**							**45767102**
Actinomyces israelii		92.86	0.37	SUC 98				
Bifidobacterium adolescentis		7.14	0.12	FRU 99				
45767502	**Low discrimination**							**45767502**
					SPO			
Actinomyces israelii		46.25	0.35	SUC 98	1			
Bifidobacterium adolescentis		29.08	0.26	FRU 99	1			
Clostridium butyricum		20.50	0.08		99			
45770000	**Acceptable identification**							**45770000**
					GRA			
Fusobacterium mortiferum		93.48	0.45	MLT 11	1			
Lactobacillus jensenii		6.52	0.15		99			
45770100	**Good identification**							**45770100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.54	0.51	MLT 11	1	1	1	
Clostridium septicum		0.73	-.03		1	99	99	
Atopobium parvulum		0.72	0.04		99	99	1	
45770102	**Low discrimination**							**45770102**
					GRA			
Fusobacterium mortiferum		55.15	0.35	MLT 11	1			
Bifidobacterium breve		19.54	0.24	BGL 10	99			
Lactobacillus acidophilus		16.10	0.13		99			
45770712	**Good identification**							**45770712**
Clostridium clostridioforme		100.0	0.41	SUC 99				
45770732	**Acceptable identification**							**45770732**
Clostridium clostridioforme		93.46	0.41	SUC 99				
Parabacteroides distasonis		6.54	0.26	XYL 91				
45771000	**Low discrimination**							**45771000**
					GRA			
Lactobacillus jensenii		80.32	0.50	SUC 90	99			
Fusobacterium mortiferum		17.46	0.48	RAF 1	1			
Actinomyces israelii		0.71	0.27		99			
45771001	**Low discrimination**							**45771001**
					GRA			
Actinomyces naeslundii		49.91	0.38	ESL 12	99			
Lactobacillus jensenii		41.15	0.15	SUC 99	99			
Fusobacterium mortiferum		8.94	0.14		1			
45771002	**Low discrimination**							**45771002**
					GRA			
Lactobacillus jensenii		78.06	0.47	SUC 90	99			
Lactobacillus acidophilus		11.05	0.29	RAF 1	99			
Prevotella oralis		3.40	0.19		1			
45771003	**Low discrimination**							**45771003**
Actinomyces naeslundii		64.53	0.38	ESL 12				
Lactobacillus jensenii		35.47	0.12	SUC 99				
45771010	**Good identification**							**45771010**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus jensenii		98.85	0.41	SUC 90	99			
Fusobacterium mortiferum		0.65	0.14	RAF 1	1			
Leptotrichia buccalis		0.50	-.11		1			
45771012	**Low discrimination**							**45771012**
					GRA			
Lactobacillus jensenii		54.81	0.38	SUC 90	99			
Leptotrichia buccalis		41.03	0.24	RAF 1	1			
Bifidobacterium breve		3.28	0.24		99			
45771020	**Low discrimination**							**45771020**
					SPO			
Lactobacillus jensenii		40.48	0.39	SUC 90	1			
Clostridium septicum		40.31	0.32	RAF 1	99			
Clostridium paraputrificum		10.29	0.19		99			
45771021	**Low discrimination**							**45771021**
					SPO			
Actinomyces naeslundii		70.85	0.38	ESL 12	1			
Lactobacillus jensenii		14.61	0.05	SUC 99	1			
Clostridium septicum		14.55	-.03		99			
45771022	**Low discrimination**							**45771022**
					GRA			
Prevotella oralis		60.00	0.41	SUC 92	1			
Lactobacillus jensenii		18.13	0.36	TRE 6	99			
Lactobacillus acidophilus		10.27	0.29	LAC 94	99			
45771023	**Low discrimination**							**45771023**
					GRA			
Actinomyces naeslundii		62.80	0.38	ESL 12	99			
Prevotella oralis		28.57	0.07	SUC 99	1			
Lactobacillus jensenii		8.63	0.02		99			
45771100	**Low discrimination**							**45771100**
					COC	GRA	SPO	
Fusobacterium mortiferum		35.81	0.54	MLT 11	1	1	1	
Clostridium septicum		17.58	0.32		1	99	99	
Atopobium parvulum		17.38	0.38		99	99	1	
45771101	**Low discrimination**							**45771101**
					GRA	SPO		
Actinomyces naeslundii		77.50	0.46	ESL 12	99	1		
Fusobacterium mortiferum		11.39	0.20	SUC 99	1	1		
Clostridium septicum		5.59	-.03		99	99		
45771102	**Low discrimination**							**45771102**
					GRA			
Bifidobacterium breve		39.12	0.59	SUC 99	99			
Lactobacillus acidophilus		32.24	0.47		99			
Prevotella oralis		13.51	0.40		1			
45771103	**Low discrimination**							**45771103**
Actinomyces naeslundii		53.30	0.46	ESL 12				
Bifidobacterium breve		20.35	0.24	SUC 99				
Lactobacillus acidophilus		16.76	0.13					
45771112	**Good identification**							**45771112**
					GRA			
Leptotrichia buccalis		92.08	0.58	SUC 99	1			
Bifidobacterium breve		7.37	0.59		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		0.33	0.35		99		
45771120	**Low discrimination**						**45771120**
Clostridium septicum		76.35	0.66	RAF 1			
Clostridium paraputrificum		19.48	0.53				
Clostridium ramosum		1.54	0.41				
45771122	**Low discrimination**						**45771122**
					GRA	SPO	
Prevotella oralis		46.08	0.61	SUC 92	1	1	
Clostridium paraputrificum		14.37	0.47	TRE 6	99	99	
Bifidobacterium breve		12.49	0.63		99	1	
45771132	**Low discrimination**						**45771132**
					GRA		
Bifidobacterium breve		46.07	0.63	SUC 99	99		
Leptotrichia buccalis		35.99	0.41		1		
Parabacteroides distasonis		14.87	0.56		1		
45771140	**Low discrimination**						**45771140**
					SPO		
Bifidobacterium breve		75.87	0.41	SUC 99	1		
Actinomyces israelii		8.02	0.30	BGL 91	1		
Clostridium butyricum		5.05	0.06		99		
45771142	**Low discrimination**						**45771142**
Bifidobacterium breve		88.18	0.59	SUC 99			
Lactobacillus acidophilus		8.98	0.32				
Actinomyces israelii		1.38	0.33				
45771150	**Good identification**						**45771150**
					GRA		
Bifidobacterium breve		97.21	0.41	SUC 99	99		
Actinomyces israelii		1.53	0.16	BGL 91	99		
Leptotrichia buccalis		1.25	-.11		1		
45771152	**Low discrimination**						**45771152**
					GRA		
Bifidobacterium breve		88.17	0.59	SUC 99	99		
Leptotrichia buccalis		11.13	0.24		1		
Bifidobacterium adolescentis		0.49	0.19		99		
45771162	**Low discrimination**						**45771162**
					SPO		
Bifidobacterium breve		83.23	0.63	SUC 99	1		
Clostridium paraputrificum		7.21	0.27		99		
Lactobacillus acidophilus		4.77	0.32		1		
45771170	**Good identification**						**45771170**
					SPO		
Bifidobacterium breve		96.95	0.46	SUC 99	1		
Clostridium paraputrificum		2.00	-.01	BGL 91	99		
Clostridium septicum		1.05	-.03		99		
45771172	**Very good identification**						**45771172**
					GRA		
Bifidobacterium breve		98.81	0.63	SUC 99	99		
Leptotrichia buccalis		0.78	0.07		1		
Parabacteroides distasonis		0.32	0.22		1		
45771320	**Low discrimination**						**45771320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO	GRA	
Clostridium ramosum		54.32	0.43	SUC 99	92	50	
Clostridium septicum		20.59	0.32		99	99	
Prevotella oralis		16.12	0.33		1	1	
45771322	**Low discrimination**						**45771322**
					SPO		
Parabacteroides distasonis		39.41	0.69	XYL 90	1		
Prevotella oralis		34.65	0.55		1		
Clostridium ramosum		24.58	0.53		92		
45771332	**Good identification**						**45771332**
Parabacteroides distasonis		94.01	0.64	XYL 90			
Clostridium clostridioforme		1.59	0.28				
Prevotella oralis		1.55	0.21				
45771500	**Good identification**						**45771500**
					GRA	SPO	
Clostridium butyricum		95.50	0.40	SUC 99	99	99	
Actinomyces israelii		3.08	0.36	XYL 97	99	1	
Fusobacterium mortiferum		0.55	0.20		1	1	
45771512	**Low discrimination**						**45771512**
					GRA		
Bifidobacterium adolescentis		57.48	0.49	SUC 99	99		
Leptotrichia buccalis		24.07	0.24	XYL 87	1		
Clostridium clostridioforme		11.88	0.34		50		
45771522	**Low discrimination**						**45771522**
					GRA	SPO	
Parabacteroides distasonis		79.01	0.57	XYL 90	1	1	
Prevotella oralis		8.01	0.27		1	1	
Clostridium butyricum		2.79	0.15		99	99	
45771532	**Low discrimination**						**45771532**
					GRA		
Parabacteroides distasonis		77.51	0.52	XYL 90	1		
Clostridium clostridioforme		13.73	0.34		50		
Bifidobacterium breve		3.96	0.29		99		
45771612	**Good identification**						**45771612**
Clostridium clostridioforme		100.0	0.40	SUC 99			
45771632	**Acceptable identification**						**45771632**
Clostridium clostridioforme		92.76	0.40	SUC 99			
Parabacteroides distasonis		7.24	0.26	XYL 91			
45771702	**Low discrimination**						**45771702**
					GRA	SPO	
Clostridium clostridioforme		74.43	0.37	SUC 99	50	66	
Parabacteroides distasonis		14.69	0.31	XYL 91	1	1	
Clostridium butyricum		6.08	0.07		99	99	
45771712	**Good identification**						**45771712**
					GRA		
Clostridium clostridioforme		96.83	0.49	SUC 99	50		
Parabacteroides distasonis		2.26	0.26	XYL 91	1		
Bifidobacterium adolescentis		0.65	0.15		99		
45771722	**Good identification**						**45771722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		92.56	0.65	XYL 90		
Clostridium clostridioforme		4.74	0.37			
Prevotella oralis		1.34	0.21			
45771732	**Low discrimination**					**45771732**
Parabacteroides distasonis		69.55	0.61	XYL 90		
Clostridium clostridioforme		30.12	0.49			
Bacteroides ovatus		0.25	0.03			
45773000	**Low discrimination**					**45773000**
Actinomyces israelii		60.45	0.40	SUC 98		
Actinomyces naeslundii		19.11	0.40	XYL 92		
Lactobacillus jensenii		12.18	0.15			
45773001	**Good identification**					**45773001**
Actinomyces naeslundii		100.0	0.43	ESL 12		
45773002	**Low discrimination**					**45773002**
Actinomyces israelii		62.29	0.43	SUC 98		
Actinomyces naeslundii		13.13	0.40	XYL 92		
Bifidobacterium breve		11.02	0.24			
45773003	**Good identification**					**45773003**
Actinomyces naeslundii		100.0	0.43	ESL 12		
45773021	**Good identification**					**45773021**
Actinomyces naeslundii		100.0	0.43	ESL 12		
45773023	**Good identification**					**45773023**
Actinomyces naeslundii		100.0	0.43	ESL 12		
45773042	**Low discrimination**					**45773042**
Actinomyces israelii		64.49	0.36	SUC 98		
Bifidobacterium breve		29.34	0.24	XYL 92		
Actinomyces naeslundii		6.17	0.27			
45773100	**Low discrimination**					**45773100**
Actinomyces israelii		46.54	0.50	SUC 98		
Bifidobacterium breve		30.24	0.41	XYL 92		
Actinomyces naeslundii		10.47	0.48			
45773101	**Good identification**					**45773101**
Actinomyces naeslundii		97.09	0.51	ESL 12		
Actinomyces israelii		2.91	0.16	SUC 99		
45773102	**Low discrimination**					**45773102**
					SPO	
Bifidobacterium breve		69.32	0.59	SUC 99	1	
Actinomyces israelii		15.83	0.53		1	
Clostridium ramosum		5.97	0.30		92	
45773103	**Low discrimination**					**45773103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		80.54	0.51	ESL 12			
Bifidobacterium breve		15.84	0.24	SUC 99			
Actinomyces israelii		3.62	0.19				
45773110	**Low discrimination**						**45773110**
					SPO		
Bifidobacterium breve		74.92	0.41	SUC 99	1		
Actinomyces israelii		17.23	0.36	BGL 91	1		
Clostridium tertium		5.52	0.13		99		
45773112	**Low discrimination**						**45773112**
					GRA		
Bifidobacterium breve		81.77	0.59	SUC 99	99		
Leptotrichia buccalis		10.33	0.24		1		
Bifidobacterium adolescentis		3.02	0.33		99		
45773120	**Low discrimination**						**45773120**
					SPO		
Clostridium ramosum		81.85	0.53	SUC 99	92		
Clostridium septicum		7.26	0.32		99		
Bifidobacterium breve		6.75	0.46		1		
45773121	**Low discrimination**						**45773121**
					SPO		
Actinomyces naeslundii		67.08	0.51	ESL 12	1		
Clostridium ramosum		28.11	0.19	SUC 99	92		
Clostridium septicum		2.49	-.03		99		
45773122	**Low discrimination**						**45773122**
					SPO	GRA	
Clostridium ramosum		80.93	0.64	SUC 99	92	50	
Bifidobacterium breve		16.88	0.63		1	99	
Prevotella oralis		0.63	0.27		1	1	
45773130	**Low discrimination**						**45773130**
					SPO		
Bifidobacterium breve		78.36	0.46	SUC 99	1		
Clostridium tertium		9.74	0.21	BGL 91	99		
Clostridium ramosum		9.59	0.19		92		
45773132	**Good identification**						**45773132**
					SPO		
Bifidobacterium breve		90.85	0.63	SUC 99	1		
Clostridium ramosum		4.40	0.30		92		
Clostridium tertium		3.35	0.29		99		
45773140	**Low discrimination**						**45773140**
Bifidobacterium breve		60.25	0.41	SUC 99			
Actinomyces israelii		36.07	0.43	BGL 91			
Actinomyces naeslundii		3.68	0.35				
45773141	**Good identification**						**45773141**
Actinomyces naeslundii		100.0	0.38	ESL 12			
45773142	**Good identification**						**45773142**
Bifidobacterium breve		90.59	0.59	SUC 99			
Actinomyces israelii		8.05	0.46				
Actinomyces naeslundii		0.55	0.35				
45773150	**Acceptable identification**						**45773150**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Bifidobacterium breve		90.17	0.41	SUC 99	1	
Actinomyces israelii		8.07	0.29	BGL 91	1	
Blautia producta		1.76	-.03		99	
45773152	**Very good identification**					**45773152**
Bifidobacterium breve		98.13	0.59	SUC 99		
Actinomyces israelii		1.30	0.32			
Bifidobacterium adolescentis		0.45	0.18			
45773160	**Low discrimination**					**45773160**
					SPO	
Bifidobacterium breve		82.66	0.46	SUC 99	1	
Clostridium ramosum		10.12	0.19	BGL 91	92	
Actinomyces naeslundii		2.84	0.35		1	
45773161	**Low discrimination**					**45773161**
Actinomyces naeslundii		83.62	0.38	ESL 12		
Bifidobacterium breve		16.38	0.12	SUC 99		
45773162	**Good identification**					**45773162**
					SPO	
Bifidobacterium breve		94.44	0.63	SUC 99	1	
Clostridium ramosum		4.57	0.30		92	
Actinomyces naeslundii		0.32	0.35		1	
45773170	**Good identification**					**45773170**
Bifidobacterium breve		100.0	0.46	SUC 99		
45773172	**Very good identification**					**45773172**
Bifidobacterium breve		100.0	0.63	SUC 99		
45773320	**Good identification**					**45773320**
					SPO	
Clostridium ramosum		99.82	0.56	SUC 99	92	
Clostridium septicum		0.07	-.03		99	
Bifidobacterium breve		0.06	0.12		1	
45773322	**Very good identification**					**45773322**
					SPO	
Clostridium ramosum		99.27	0.66	SUC 99	92	
Parabacteroides distasonis		0.28	0.35		1	
Prevotella oralis		0.25	0.21		1	
45773400	**Acceptable identification**					**45773400**
					SPO	
Actinomyces israelii		90.80	0.38	SUC 98	1	
Clostridium butyricum		9.20	0.00	XYL 92	99	
45773402	**Acceptable identification**					**45773402**
					SPO	
Actinomyces israelii		93.47	0.41	SUC 98	1	
Bifidobacterium adolescentis		4.08	0.12	XYL 92	1	
Clostridium butyricum		2.46	-.07		99	
45773500	**Low discrimination**					**45773500**
					SPO	
Actinomyces israelii		61.56	0.48	SUC 98	1	
Clostridium butyricum		37.43	0.24	XYL 92	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		1.01	0.12		1		
45773502	**Low discrimination**						**45773502**
					SPO		
Bifidobacterium adolescentis		46.92	0.46	SUC 99	1		
Actinomyces israelii		43.43	0.51	XYL 87	1		
Clostridium butyricum		6.85	0.17		99		
45773512	**Low discrimination**						**45773512**
Bifidobacterium adolescentis		51.23	0.47	SUC 99			
Bifidobacterium dentium		40.29	0.27	XYL 87			
Actinomyces israelii		5.80	0.37				
45773542	**Low discrimination**						**45773542**
Actinomyces israelii		61.91	0.44	SUC 98			
Bifidobacterium adolescentis		21.26	0.30	XYL 92			
Bifidobacterium breve		8.61	0.24				
45774122	**Low discrimination**						**45774122**
Parabacteroides distasonis		89.70	0.43	GAL 99			
Prevotella oralis		7.58	0.12				
Clostridium clostridioforme		2.72	0.11				
45774132	**Low discrimination**						**45774132**
Parabacteroides distasonis		79.56	0.38	GAL 99			
Clostridium clostridioforme		20.45	0.22				
45774312	**Good identification**						**45774312**
Clostridium clostridioforme		98.42	0.37	SUC 99			
Parabacteroides distasonis		1.58	0.12	ARA 94			
45774322	**Acceptable identification**						**45774322**
Parabacteroides distasonis		92.05	0.51	GAL 99			
Clostridium clostridioforme		6.84	0.26				
Prevotella oralis		1.11	0.05				
45774332	**Low discrimination**						**45774332**
Parabacteroides distasonis		61.42	0.46	GAL 99			
Clostridium clostridioforme		38.58	0.37				
45774512	**Low discrimination**						**45774512**
					SPO		
Clostridium clostridioforme		88.65	0.43	SUC 99	66		
Bifidobacterium adolescentis		8.60	0.29	RHA 88	1		
Clostridium butyricum		1.92	0.03		99		
45774522	**Low discrimination**						**45774522**
					GRA	SPO	
Parabacteroides distasonis		52.54	0.39	GAL 99	1	1	
Clostridium clostridioforme		40.80	0.32		50	66	
Clostridium butyricum		6.66	0.07		99	99	
45774532	**Low discrimination**						**45774532**
					GRA		
Clostridium clostridioforme		85.56	0.43	SUC 99	50		
Parabacteroides distasonis		13.03	0.35	RHA 88	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		1.42	-.03		99		
45774602	**Good identification**						**45774602**
Clostridium clostridioforme		100.0	0.37	SUC 99			
45774612	**Good identification**						**45774612**
Clostridium clostridioforme		100.0	0.49	SUC 99			
45774622	**Good identification**						**45774622**
Clostridium clostridioforme		98.25	0.37	SUC 99			
Parabacteroides distasonis		1.75	0.13				
45774632	**Good identification**						**45774632**
Clostridium clostridioforme		100.0	0.49	SUC 99			
45774702	**Good identification**						**45774702**
					GRA	SPO	
Clostridium clostridioforme		98.69	0.46	SUC 99	50	66	
Clostridium butyricum		0.78	-.01		99	99	
Parabacteroides distasonis		0.53	0.13		1	1	
45774712	**Very good identification**						**45774712**
Clostridium clostridioforme		100.0	0.58	SUC 99			
45774722	**Low discrimination**						**45774722**
Clostridium clostridioforme		64.77	0.46	SUC 99			
Parabacteroides distasonis		34.13	0.47				
Bacteroides ovatus		1.10	0.06				
45774732	**Good identification**						**45774732**
Clostridium clostridioforme		93.92	0.58	SUC 99			
Parabacteroides distasonis		5.85	0.43				
Bacteroides ovatus		0.23	0.03				
45775002	**Low discrimination**						**45775002**
					SPO		
Actinomyces israelii		68.20	0.48	SUC 98	1		
Clostridium butyricum		23.36	0.19		99		
Lactobacillus jensenii		3.01	0.12		1		
45775022	**Low discrimination**						**45775022**
					GRA		
Parabacteroides distasonis		59.30	0.43	LAC 99	1		
Prevotella oralis		31.67	0.25		1		
Actinomyces israelii		4.53	0.27		99		
45775032	**Low discrimination**						**45775032**
					GRA		
Parabacteroides distasonis		79.91	0.38	LAC 99	1		
Clostridium clostridioforme		18.40	0.22		50		
Actinomyces israelii		1.69	0.13		99		
45775040	**Low discrimination**						**45775040**
					SPO		
Actinomyces israelii		77.20	0.38	SUC 98	1		
Clostridium butyricum		22.80	0.08		99		
45775042	**Acceptable identification**						**45775042**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		92.89	0.41	SUC 98	1		
Clostridium butyricum		7.11	0.01		99		
45775100	**Low discrimination**						**45775100**
					COC	SPO	
Clostridium butyricum		77.98	0.50	SUC 99	1	99	
Atopobium parvulum		11.04	0.42	ARA 90	99	1	
Actinomyces israelii		9.84	0.55		1	1	
45775102	**Low discrimination**						**45775102**
					SPO		
Clostridium butyricum		50.25	0.43	SUC 99	99		
Actinomyces israelii		24.46	0.58	ARA 90	1		
Bifidobacterium adolescentis		13.02	0.47		1		
45775112	**Low discrimination**						**45775112**
					GRA		
Bifidobacterium adolescentis		46.28	0.49	SUC 99	99		
Leptotrichia buccalis		19.38	0.24	ARA 87	1		
Actinomyces israelii		10.63	0.44		99		
45775122	**Good identification**						**45775122**
					GRA	SPO	
Parabacteroides distasonis		90.43	0.77		1	1	
Prevotella oralis		7.64	0.46		1	1	
Clostridium butyricum		0.78	0.24		99	99	
45775123	**Acceptable identification**						**45775123**
Parabacteroides distasonis		92.21	0.43	URE 1			
Prevotella oralis		7.79	0.12				
45775130	**Low discrimination**						**45775130**
					GRA	SPO	
Parabacteroides distasonis		74.52	0.38	BGL 99	1	1	
Clostridium butyricum		16.01	0.10		99	99	
Actinomyces israelii		4.21	0.21		99	1	
45775132	**Very good identification**						**45775132**
					GRA		
Parabacteroides distasonis		98.18	0.72		1		
Clostridium clostridioforme		0.77	0.31		50		
Bifidobacterium breve		0.34	0.29		99		
45775133	**Good identification**						**45775133**
Parabacteroides distasonis		100.0	0.38	URE 1			
45775142	**Low discrimination**						**45775142**
					SPO		
Actinomyces israelii		59.36	0.51	SUC 98	1		
Clostridium butyricum		27.27	0.24		99		
Bifidobacterium adolescentis		10.04	0.32		1		
45775162	**Low discrimination**						**45775162**
					GRA		
Parabacteroides distasonis		67.61	0.43	SOR 1	1		
Bifidobacterium breve		12.53	0.29		99		
Actinomyces israelii		8.03	0.31		99		
45775172	**Low discrimination**						**45775172**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		72.61	0.38	SOR 1	1		
Bifidobacterium breve		25.00	0.29		99		
Actinomyces israelii		2.39	0.17		99		
45775212	**Good identification**						**45775212**
					GRA		
Clostridium clostridioforme		96.83	0.36	SUC 99	50		
Parabacteroides distasonis		1.74	0.12	ARA 94	1		
Actinomyces israelii		1.43	0.15		99		
45775222	**Low discrimination**						**45775222**
Parabacteroides distasonis		87.50	0.51	LAC 99			
Prevotella oralis		6.68	0.19				
Clostridium clostridioforme		5.82	0.25				
45775232	**Low discrimination**						**45775232**
Parabacteroides distasonis		63.98	0.46	LAC 99			
Clostridium clostridioforme		36.02	0.36				
45775302	**Low discrimination**						**45775302**
					GRA		
Parabacteroides distasonis		65.98	0.51	NAG 99	1		
Clostridium clostridioforme		14.70	0.34		50		
Actinomyces israelii		7.92	0.39		99		
45775312	**Low discrimination**						**45775312**
					GRA		
Clostridium clostridioforme		63.77	0.45	SUC 99	50		
Parabacteroides distasonis		33.84	0.46	ARA 94	1		
Actinomyces israelii		1.13	0.25		99		
45775320	**Low discrimination**						**45775320**
					GRA	SPO	
Parabacteroides distasonis		88.53	0.51	BGL 99	1	1	
Prevotella oralis		5.57	0.18		1	1	
Clostridium butyricum		2.01	0.06		99	99	
45775322	**Very good identification**						**45775322**
Parabacteroides distasonis		98.44	0.85				
Prevotella oralis		1.19	0.40				
Clostridium clostridioforme		0.22	0.34				
45775323	**Good identification**						**45775323**
Parabacteroides distasonis		98.81	0.51	URE 1			
Prevotella oralis		1.19	0.05				
45775330	**Good identification**						**45775330**
Parabacteroides distasonis		98.13	0.46	BGL 99			
Clostridium clostridioforme		1.87	0.11				
45775332	**Very good identification**						**45775332**
Parabacteroides distasonis		98.04	0.81				
Clostridium clostridioforme		1.87	0.45				
Bacteroides ovatus		0.07	0.11				
45775333	**Good identification**						**45775333**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		98.13	0.46	URE 1			
Clostridium clostridioforme		1.87	0.11				
45775362	**Good identification**						**45775362**
Parabacteroides distasonis		98.81	0.51	SOR 1			
Prevotella oralis		1.19	0.05				
45775372	**Good identification**						**45775372**
Parabacteroides distasonis		98.13	0.46	SOR 1			
Clostridium clostridioforme		1.87	0.11				
45775400	**Acceptable identification**						**45775400**
					SPO		
Clostridium butyricum		93.56	0.42	SUC 99	99		
Actinomyces israelii		6.44	0.43	LAC 96	1		
45775402	**Low discrimination**						**45775402**
					SPO		
Clostridium butyricum		70.04	0.35	SUC 99	99		
Actinomyces israelii		18.59	0.46	LAC 96	1		
Clostridium clostridioforme		8.10	0.31		66		
45775412	**Low discrimination**						**45775412**
					SPO		
Clostridium clostridioforme		77.29	0.42	SUC 99	66		
Bifidobacterium adolescentis		8.37	0.29	RHA 88	1		
Clostridium butyricum		7.72	0.13		99		
45775422	**Low discrimination**						**45775422**
					GRA	SPO	
Parabacteroides distasonis		43.09	0.39	LAC 99	1	1	
Clostridium clostridioforme		29.98	0.31		50	66	
Clostridium butyricum		22.54	0.17		99	99	
45775432	**Low discrimination**						**45775432**
					GRA		
Clostridium clostridioforme		83.54	0.42	SUC 99	50		
Parabacteroides distasonis		14.19	0.35	RHA 88	1		
Bifidobacterium dentium		1.55	-.03		99		
45775442	**Low discrimination**						**45775442**
					SPO		
Actinomyces israelii		52.68	0.39	SUC 98	1		
Clostridium butyricum		44.38	0.17		99		
Bifidobacterium adolescentis		2.95	0.12		1		
45775500	**Very good identification**						**45775500**
					SPO		
Clostridium butyricum		98.70	0.66	SUC 99	99		
Actinomyces israelii		1.13	0.54		1		
Bifidobacterium adolescentis		0.08	0.27		1		
45775502	**Low discrimination**						**45775502**
					SPO		
Clostridium butyricum		79.34	0.59	SUC 99	99		
Bifidobacterium adolescentis		15.29	0.61		1		
Actinomyces israelii		3.51	0.57		1		
45775510	**Acceptable identification**						**45775510**
					SPO		
Clostridium butyricum		92.71	0.44	SUC 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		3.02	0.40	MLZ 5	1		
Bifidobacterium adolescentis		1.63	0.29		1		
45775512	**Low discrimination**						**45775512**
					SPO		
Bifidobacterium adolescentis		59.65	0.63	SUC 99	1		
Clostridium clostridioforme		18.62	0.51		66		
Clostridium butyricum		13.33	0.37		99		
45775520	**Good identification**						**45775520**
					GRA	SPO	
Clostridium butyricum		95.54	0.48	SUC 99	99	99	
Parabacteroides distasonis		2.96	0.39	NAG 8	1	1	
Actinomyces israelii		0.80	0.33		99	1	
45775522	**Low discrimination**						**45775522**
					GRA	SPO	
Parabacteroides distasonis		85.77	0.73		1	1	
Clostridium butyricum		10.87	0.41		99	99	
Clostridium clostridioforme		2.02	0.40		50	66	
45775523	**Low discrimination**						**45775523**
					GRA	SPO	
Parabacteroides distasonis		86.93	0.39	URE 1	1	1	
Clostridium butyricum		11.02	0.07		99	99	
Clostridium clostridioforme		2.05	0.06		50	66	
45775532	**Low discrimination**						**45775532**
					GRA		
Parabacteroides distasonis		74.91	0.69		1		
Clostridium clostridioforme		14.91	0.51		50		
Bifidobacterium dentium		8.16	0.32		99		
45775540	**Acceptable identification**						**45775540**
					COC	SPO	
Clostridium butyricum		91.38	0.48	SUC 99	1	99	
Actinomyces israelii		4.69	0.47	SOR 8	1	1	
Blautia producta		3.04	0.24		99	1	
45775542	**Low discrimination**						**45775542**
					SPO		
Clostridium butyricum		67.32	0.41	SUC 99	99		
Bifidobacterium adolescentis		18.43	0.46	SOR 8	1		
Actinomyces israelii		13.32	0.50		1		
45775552	**Low discrimination**						**45775552**
					SPO		
Bifidobacterium adolescentis		73.30	0.47	SUC 99	1		
Clostridium butyricum		11.53	0.19	SOR 11	99		
Actinomyces israelii		6.48	0.36		1		
45775602	**Acceptable identification**						**45775602**
					SPO		
Clostridium clostridioforme		93.81	0.46	SUC 99	66		
Clostridium butyricum		3.42	0.09		99		
Actinomyces israelii		2.21	0.27		1		
45775612	**Very good identification**						**45775612**
Clostridium clostridioforme		100.0	0.57	SUC 99			
45775622	**Low discrimination**						**45775622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		62.05	0.46	SUC 99			
Parabacteroides distasonis		36.49	0.47				
Bacteroides ovatus		1.18	0.06				
45775632	**Good identification**						**45775632**
Clostridium clostridioforme		93.26	0.57	SUC 99			
Parabacteroides distasonis		6.48	0.43				
Bacteroides ovatus		0.26	0.03				
45775700	**Acceptable identification**						**45775700**
					GRA	SPO	
Clostridium butyricum		91.68	0.40	SUC 99	99	99	
Mitsuokella multacida		4.14	0.25	RHA 3	1	1	
Actinomyces israelii		2.56	0.34		99	1	
45775702	**Low discrimination**						**45775702**
					GRA	SPO	
Clostridium clostridioforme		67.09	0.55	SUC 99	50	66	
Clostridium butyricum		17.53	0.33		99	99	
Parabacteroides distasonis		11.78	0.47		1	1	
45775712	**Very good identification**						**45775712**
					GRA		
Clostridium clostridioforme		97.03	0.66	SUC 99	50		
Parabacteroides distasonis		2.01	0.43		1		
Bifidobacterium adolescentis		0.43	0.29		99		
45775720	**Low discrimination**						**45775720**
					GRA	SPO	
Parabacteroides distasonis		55.64	0.47	BGL 99	1	1	
Clostridium butyricum		18.51	0.22		99	99	
Bacteroides vulgatus		18.07	0.21		1	1	
45775722	**Good identification**						**45775722**
Parabacteroides distasonis		91.27	0.82				
Clostridium clostridioforme		5.25	0.55				
Bacteroides ovatus		2.95	0.40				
45775723	**Acceptable identification**						**45775723**
Parabacteroides distasonis		91.75	0.47	URE 1			
Clostridium clostridioforme		5.28	0.20				
Bacteroides ovatus		2.97	0.06				
45775730	**Low discrimination**						**45775730**
Parabacteroides distasonis		63.06	0.43	BGL 99			
Clostridium clostridioforme		30.68	0.32				
Bacteroides ovatus		2.53	0.03				
45775732	**Low discrimination**						**45775732**
Parabacteroides distasonis		65.37	0.77				
Clostridium clostridioforme		31.81	0.66				
Bacteroides ovatus		2.62	0.37				
45775733	**Low discrimination**						**45775733**
Parabacteroides distasonis		65.50	0.43	URE 1			
Clostridium clostridioforme		31.87	0.32				

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		2.62	0.03			
45775762	**Low discrimination**					**45775762**
Parabacteroides distasonis		83.18	0.47	SOR 1		
Bacteroides ovatus		11.10	0.16			
Clostridium clostridioforme		4.78	0.20			
45775772	**Low discrimination**					**45775772**
Parabacteroides distasonis		59.65	0.43	SOR 1		
Clostridium clostridioforme		29.02	0.32			
Bacteroides ovatus		9.86	0.13			
45776102	**Acceptable identification**					**45776102**
Actinomyces israelii		92.86	0.37	SUC 98		
Bifidobacterium adolescentis		7.14	0.12	GAL 99		
45776502	**Low discrimination**					**45776502**
					SPO	
Actinomyces israelii		46.25	0.35	SUC 98	1	
Bifidobacterium adolescentis		29.08	0.26	GAL 99	1	
Clostridium butyricum		20.50	0.08		99	
45777000	**Very good identification**					**45777000**
					SPO	
Actinomyces israelii		97.48	0.58	SUC 98	1	
Clostridium butyricum		2.53	0.10		99	
45777002	**Very good identification**					**45777002**
					SPO	
Actinomyces israelii		99.03	0.61	SUC 98	1	
Clostridium butyricum		0.67	0.03		99	
Bifidobacterium adolescentis		0.31	0.12		1	
45777010	**Good identification**					**45777010**
Actinomyces israelii		100.0	0.44	SUC 98		
45777012	**Good identification**					**45777012**
Actinomyces israelii		95.66	0.47	SUC 98		
Bifidobacterium adolescentis		2.43	0.13	MLZ 13		
Bifidobacterium dentium		1.91	-.07			
45777020	**Good identification**					**45777020**
Actinomyces israelii		100.0	0.37	SUC 98		
45777022	**Good identification**					**45777022**
Actinomyces israelii		100.0	0.40	SUC 98		
45777040	**Good identification**					**45777040**
Actinomyces israelii		100.0	0.51	SUC 98		
45777042	**Good identification**					**45777042**
Actinomyces israelii		100.0	0.54	SUC 98		
45777050	**Acceptable identification**					**45777050**
					COC	
Actinomyces israelii		92.94	0.37	SUC 98	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		7.06	-.03	MLZ 13	99		
45777052	**Good identification**						**45777052**
Actinomyces israelii		100.0	0.40	SUC 98			
45777100	**Low discrimination**						**45777100**
					SPO		
Actinomyces israelii		86.07	0.68	SUC 98	1		
Clostridium butyricum		13.38	0.33		99		
Clostridium tertium		0.28	0.11		99		
45777102	**Low discrimination**						**45777102**
					SPO		
Actinomyces israelii		88.64	0.71	SUC 98	1		
Bifidobacterium adolescentis		6.81	0.46		1		
Clostridium butyricum		3.57	0.26		99		
45777103	**Acceptable identification**						**45777103**
Actinomyces israelii		92.86	0.37	SUC 98			
Bifidobacterium adolescentis		7.14	0.12	URE 1			
45777110	**Low discrimination**						**45777110**
					SPO		
Actinomyces israelii		84.50	0.54	SUC 98	1		
Bifidobacterium dentium		6.20	0.10	MLZ 13	1		
Clostridium butyricum		4.63	0.11		99		
45777112	**Low discrimination**						**45777112**
Actinomyces israelii		44.58	0.57	SUC 98			
Bifidobacterium adolescentis		28.03	0.47	MLZ 13			
Bifidobacterium dentium		22.04	0.27				
45777120	**Low discrimination**						**45777120**
					SPO		
Actinomyces israelii		62.92	0.48	SUC 98	1		
Clostridium butyricum		13.32	0.15	NAG 6	99		
Clostridium ramosum		12.12	0.19		92		
45777122	**Low discrimination**						**45777122**
					SPO	GRA	
Actinomyces israelii		37.89	0.51	SUC 98	1	99	
Parabacteroides distasonis		21.90	0.43	NAG 6	1	1	
Clostridium ramosum		19.46	0.30		92	50	
45777140	**Low discrimination**						**45777140**
					COC	SPO	
Actinomyces israelii		88.62	0.61	SUC 98	1	1	
Blautia producta		8.30	0.24		99	1	
Clostridium butyricum		3.08	0.15		1	99	
45777142	**Very good identification**						**45777142**
Actinomyces israelii		95.21	0.64	SUC 98			
Bifidobacterium adolescentis		2.33	0.30				
Bifidobacterium breve		0.94	0.24				
45777152	**Low discrimination**						**45777152**
					COC		
Actinomyces israelii		70.51	0.50	SUC 98	1		
Bifidobacterium adolescentis		14.09	0.32	MLZ 13	1		

Code / Taxon		ID score	T index	tests against		Differentiation tests		Code
Blautia producta		9.82	0.15			99		
45777160	**Low discrimination**							**45777160**
						COC	SPO	
Actinomyces israelii		79.92	0.41	SUC	98	1	1	
Blautia producta		13.03	0.07	NAG	6	99	1	
Clostridium butyricum		3.78	-.03			1	99	
45777162	**Low discrimination**							**45777162**
Actinomyces israelii		78.40	0.44	SUC	98			
Bifidobacterium breve		21.60	0.29	NAG	6			
45777200	**Good identification**							**45777200**
Actinomyces israelii		100.0	0.39	SUC	98			
45777202	**Good identification**							**45777202**
Actinomyces israelii		100.0	0.42	SUC	98			
45777300	**Acceptable identification**							**45777300**
						SPO		
Actinomyces israelii		94.00	0.49	SUC	98	1		
Clostridium butyricum		6.00	0.07	RHA	7	99		
45777302	**Good identification**							**45777302**
						GRA	SPO	
Actinomyces israelii		95.46	0.52	SUC	98	99	1	
Clostridium butyricum		1.58	0.00	RHA	7	99	99	
Parabacteroides distasonis		1.42	0.17			1	1	
45777312	**Low discrimination**							**45777312**
Actinomyces israelii		76.62	0.38	SUC	98			
Clostridium clostridioforme		7.73	0.11	RHA	7			
Bifidobacterium adolescentis		6.47	0.13	MLZ	13			
45777322	**Low discrimination**							**45777322**
						SPO	GRA	
Parabacteroides distasonis		68.10	0.51	MAN	1	1	1	
Clostridium ramosum		26.74	0.32			92	50	
Actinomyces israelii		2.96	0.31			1	99	
45777332	**Acceptable identification**							**45777332**
						GRA		
Parabacteroides distasonis		92.10	0.46	MAN	1	1		
Bacteroides ovatus		2.31	0.03			1		
Bifidobacterium dentium		2.05	-.03			99		
45777342	**Acceptable identification**							**45777342**
						COC		
Actinomyces israelii		91.56	0.45	SUC	98	1		
Blautia producta		8.44	0.07	RHA	7	99		
45777400	**Low discrimination**							**45777400**
						SPO		
Actinomyces israelii		77.83	0.56	SUC	98	1		
Clostridium butyricum		22.17	0.26			99		
45777402	**Good identification**							**45777402**
						SPO		
Actinomyces israelii		90.67	0.59	SUC	98	1		
Clostridium butyricum		6.70	0.19			99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		2.30	0.26		1		
45777410	**Low discrimination**						**45777410**
					SPO		
Actinomyces israelii		66.88	0.42	SUC 98	1		
Bifidobacterium dentium		23.99	0.10	MLZ 13	1		
Clostridium butyricum		6.71	0.04		99		
45777420	**Low discrimination**						**45777420**
					SPO		
Actinomyces israelii		72.04	0.36	SUC 98	1		
Clostridium butyricum		27.96	0.08	NAG 6	99		
45777422	**Low discrimination**						**45777422**
					SPO		
Actinomyces israelii		83.18	0.39	SUC 98	1		
Bifidobacterium dentium		8.45	-.03	NAG 6	1		
Clostridium butyricum		8.37	0.01		99		
45777440	**Low discrimination**						**45777440**
					COC	SPO	
Actinomyces israelii		65.72	0.49	SUC 98	1	1	
Blautia producta		30.09	0.24		99	1	
Clostridium butyricum		4.19	0.08		1	99	
45777442	**Acceptable identification**						**45777442**
					COC	SPO	
Actinomyces israelii		94.47	0.52	SUC 98	1	1	
Blautia producta		3.20	0.07		99	1	
Clostridium butyricum		1.56	0.01		1	99	
45777452	**Low discrimination**						**45777452**
					COC		
Actinomyces israelii		55.25	0.38	SUC 98	1		
Blautia producta		37.63	0.15	MLZ 13	99		
Bifidobacterium adolescentis		3.65	0.12		1		
45777500	**Low discrimination**						**45777500**
					COC	SPO	
Clostridium butyricum		60.97	0.50	SUC 99	1	99	
Actinomyces israelii		35.66	0.67	MAN 10	1	1	
Blautia producta		1.59	0.24		99	1	
45777502	**Low discrimination**						**45777502**
					SPO		
Actinomyces israelii		46.12	0.70	SUC 98	1		
Bifidobacterium adolescentis		29.00	0.60		1		
Clostridium butyricum		20.44	0.43		99		
45777503	**Low discrimination**						**45777503**
					SPO		
Actinomyces israelii		46.25	0.35	SUC 98	1		
Bifidobacterium adolescentis		29.08	0.26	URE 1	1		
Clostridium butyricum		20.50	0.08		99		
45777510	**Low discrimination**						**45777510**
					COC		
Bifidobacterium dentium		77.12	0.44	SUC 99	1		
Actinomyces israelii		8.69	0.52	BGL 91	1		
Blautia producta		7.77	0.32		99		
45777512	**Good identification**						**45777512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		90.35	0.62	SUC 99			
Bifidobacterium adolescentis		7.77	0.61				
Actinomyces israelii		1.51	0.55				
45777530	**Good identification**						**45777530**
					COC		
Bifidobacterium dentium		98.56	0.49	SUC 99	1		
Blautia producta		0.62	0.15	BGL 91	99		
Actinomyces israelii		0.40	0.32		1		
45777532	**Very good identification**						**45777532**
					GRA		
Bifidobacterium dentium		99.74	0.66	SUC 99	99		
Parabacteroides distasonis		0.09	0.35		1		
Actinomyces israelii		0.06	0.35		99		
45777540	**Low discrimination**						**45777540**
					COC	SPO	
Blautia producta		88.63	0.58	SUC 99	99	1	
Actinomyces israelii		7.82	0.60		1	1	
Clostridium butyricum		2.99	0.31		1	99	
45777542	**Low discrimination**						**45777542**
					COC		
Actinomyces israelii		46.60	0.63	SUC 98	1		
Blautia producta		39.12	0.41		99		
Bifidobacterium adolescentis		9.31	0.44		1		
45777550	**Very good identification**						**45777550**
					COC		
Blautia producta		99.33	0.66	SUC 99	99		
Actinomyces israelii		0.44	0.45		1		
Bifidobacterium dentium		0.10	0.10		1		
45777552	**Low discrimination**						**45777552**
					COC		
Blautia producta		79.92	0.50	SUC 99	99		
Bifidobacterium adolescentis		7.75	0.46	BGL 10	1		
Bifidobacterium dentium		7.37	0.27		1		
45777560	**Acceptable identification**						**45777560**
					COC	SPO	
Blautia producta		91.67	0.41	SUC 99	99	1	
Actinomyces israelii		4.65	0.39	NAG 10	1	1	
Clostridium butyricum		2.42	0.13		1	99	
45777570	**Good identification**						**45777570**
					COC		
Blautia producta		98.01	0.50	SUC 99	99		
Bifidobacterium dentium		1.59	0.15	NAG 10	1		
Actinomyces israelii		0.25	0.25		1		
45777600	**Low discrimination**						**45777600**
					SPO		
Actinomyces israelii		89.52	0.37	SUC 98	1		
Clostridium butyricum		10.48	0.00	RHA 7	99		
45777602	**Acceptable identification**						**45777602**
					SPO		
Actinomyces israelii		90.41	0.40	SUC 98	1		
Clostridium clostridioforme		6.84	0.11	RHA 7	66		
Clostridium butyricum		2.74	-.07		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
45777700	**Low discrimination**						**45777700**
					GRA	SPO	
Actinomyces israelii		33.79	0.47	SUC 98	99	1	
Clostridium butyricum		23.73	0.24	RHA 7	99	99	
Mitsuokella multacida		22.49	0.32		1	1	
45777702	**Low discrimination**						**45777702**
					SPO		
Actinomyces israelii		68.58	0.50	SUC 98	1		
Clostridium butyricum		12.49	0.17	RHA 7	99		
Bifidobacterium adolescentis		5.79	0.26		1		
45777740	**Very good identification**						**45777740**
					COC	GRA	
Blautia producta		98.13	0.58	SUC 99	99	99	
Mitsuokella multacida		1.12	0.32		1	1	
Actinomyces israelii		0.65	0.40		1	99	
45777742	**Acceptable identification**						**45777742**
					COC	GRA	
Blautia producta		90.33	0.41	SUC 99	99	99	
Actinomyces israelii		8.10	0.43	BGL 10	1	99	
Mitsuokella multacida		1.03	0.15		1	1	
45777750	**Very good identification**						**45777750**
					COC	GRA	
Blautia producta		99.84	0.66	SUC 99	99	99	
Mitsuokella multacida		0.13	0.24		1	1	
Actinomyces israelii		0.03	0.26		1	99	
45777752	**Good identification**						**45777752**
					COC	GRA	
Blautia producta		99.24	0.50	SUC 99	99	99	
Actinomyces israelii		0.44	0.29	BGL 10	1	99	
Mitsuokella multacida		0.13	0.07		1	1	
45777760	**Good identification**						**45777760**
					COC	GRA	
Blautia producta		97.15	0.41	SUC 99	99	99	
Mitsuokella multacida		2.49	0.21	NAG 10	1	1	
Actinomyces israelii		0.37	0.20		1	99	
45777770	**Good identification**						**45777770**
					COC	GRA	
Blautia producta		99.72	0.50	SUC 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
46000000	**Low discrimination**						**46000000**
					COC	SPO	
Atopobium minutum		61.62	0.58	SUC 5	1	1	
Clostridium botulinum biovar A		29.17	0.45		1	99	
Parvimonas micra		2.49	0.15		99	1	
46000002	**Good identification**						**46000002**
					COC	SPO	
Atopobium minutum		96.97	0.55	SUC 5	1	1	
Clostridium botulinum biovar A		0.70	0.11		1	99	
Anaerococcus prevotii		0.69	0.05		99	1	
46000020	**Good identification**						**46000020**
					SPO		
Atopobium minutum		96.89	0.50	SUC 5	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		1.39	0.11		99		
Clostridium bifermentans		0.63	0.05		99		
46000022	**Low discrimination**						**46000022**
					GRA		
Atopobium minutum		54.79	0.47	SUC 5	99		
Pseudoflavonifractor capillosus		45.21	0.32		1		
46000040	**Acceptable identification**						**46000040**
					SPO		
Clostridium botulinum biovar A		92.47	0.36	GLU 90	99		
Atopobium minutum		6.61	0.24	SUC 7	1		
Clostridium bifermentans		0.93	0.01	MNS 7	99		
46001000	**Good identification**						**46001000**
					COC	SPO	
Atopobium minutum		96.64	0.50	SUC 5	1	1	
Clostridium botulinum biovar A		1.39	0.11		1	99	
Gemella morbillorum		1.18	0.13		99	1	
46001002	**Good identification**						**46001002**
Atopobium minutum		100.0	0.47	SUC 5			
46001020	**Good identification**						**46001020**
Atopobium minutum		100.0	0.42	SUC 5			
46001022	**Low discrimination**						**46001022**
					GRA		
Atopobium minutum		78.43	0.39	SUC 5	99		
Pseudoflavonifractor capillosus		21.57	0.15		1		
46010000	**Low discrimination**						**46010000**
					GRA	SPO	
Atopobium minutum		82.79	0.58	SUC 5	99	1	
Fusobacterium mortiferum		8.84	0.47		1	1	
Clostridium botulinum biovar A		2.71	0.25		99	99	
46010002	**Good identification**						**46010002**
					GRA	CAT	
Atopobium minutum		98.04	0.55	SUC 5	99	1	
Fusobacterium mortiferum		1.75	0.30		1	5	
Propionibacterium granulosum		0.21	0.11		99	90	
46010020	**Good identification**						**46010020**
					COC	CAT	
Atopobium minutum		98.92	0.50	SUC 5	1	1	
Propionibacterium granulosum		0.43	0.11		1	90	
Gemella morbillorum		0.33	0.03		99	1	
46010022	**Acceptable identification**						**46010022**
					GRA		
Atopobium minutum		90.96	0.47	SUC 5	99		
Pseudoflavonifractor capillosus		9.04	0.15		1		
46010100	**Good identification**						**46010100**
					GRA		
Fusobacterium mortiferum		95.68	0.53	GLU 99	1		
Atopobium minutum		3.88	0.24		99		
Actinomyces meyeri		0.44	0.04		99		
46010102	**Low discrimination**						**46010102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		80.44	0.37	GLU 99	1		
Atopobium minutum		19.57	0.21	BGL 10	99		
46011000	**Low discrimination**						**46011000**
					GRA	CAT	
Atopobium minutum		66.05	0.50	SUC 5	99	1	
Fusobacterium mortiferum		31.74	0.50		1	5	
Propionibacterium granulosum		1.15	0.22		99	90	
46011002	**Acceptable identification**						**46011002**
					GRA	CAT	
Atopobium minutum		92.03	0.47	SUC 5	99	1	
Fusobacterium mortiferum		7.37	0.33		1	5	
Propionibacterium granulosum		0.60	0.11		99	90	
46011020	**Good identification**						**46011020**
					GRA	CAT	
Atopobium minutum		95.35	0.42	SUC 5	99	1	
Fusobacterium mortiferum		1.39	0.16		1	5	
Propionibacterium granulosum		1.24	0.11		99	90	
46011022	**Good identification**						**46011022**
					GRA		
Atopobium minutum		96.80	0.39	SUC 5	99		
Pseudoflavonifractor capillosus		3.21	-.01		1		
46011100	**Good identification**						**46011100**
					GRA		
Fusobacterium mortiferum		98.05	0.56	GLU 99	1		
Atopobium minutum		0.88	0.16		99		
Actinomyces meyeri		0.45	0.07		99		
46011102	**Acceptable identification**						**46011102**
					GRA		
Fusobacterium mortiferum		94.87	0.40	GLU 99	1		
Atopobium minutum		5.13	0.13	BGL 10	99		
46015100	**Low discrimination**						**46015100**
					GRA		
Fusobacterium mortiferum		89.47	0.37	GLU 99	1		
Actinomyces meyeri		10.53	0.12	XYL 7	99		
46030000	**Low discrimination**						**46030000**
					GRA	CAT	
Fusobacterium mortiferum		47.23	0.38	GLU 99	1	5	
Propionibacterium granulosum		25.69	0.30		99	90	
Atopobium minutum		14.96	0.24		99	1	
46030100	**Good identification**						**46030100**
Fusobacterium mortiferum		100.0	0.44	GLU 99			
46031000	**Low discrimination**						**46031000**
					GRA	CAT	
Fusobacterium mortiferum		69.78	0.41	GLU 99	1	5	
Propionibacterium granulosum		25.31	0.30		99	90	
Atopobium minutum		4.91	0.16		99	1	
46031100	**Acceptable identification**						**46031100**
					COC	GRA	
Fusobacterium mortiferum		90.65	0.47	GLU 99	1	1	
Atopobium parvulum		9.35	0.19		99	99	
46050000	**Low discrimination**						**46050000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		77.90	0.39	GLU 99	1		
Atopobium minutum		22.11	0.24		99		
46050100	**Good identification**						**46050100**
Fusobacterium mortiferum		100.0	0.45	GLU 99			
46051000	**Low discrimination**						**46051000**
					GRA		
Fusobacterium mortiferum		81.24	0.42	GLU 99	1		
Lactobacillus jensenii		7.55	0.14		99		
Lactobacillus catenaformis		6.09	0.12		99		
46051100	**Acceptable identification**						**46051100**
					GRA		
Fusobacterium mortiferum		90.91	0.48	GLU 99	1		
Lactobacillus catenaformis		4.38	0.15		99		
Collinsella aerofaciens		2.90	-.03		99		
46070100	**Good identification**						**46070100**
Fusobacterium mortiferum		100.0	0.36	GLU 99			
46071100	**Low discrimination**						**46071100**
					COC	GRA	
Fusobacterium mortiferum		58.07	0.39	GLU 99	1	1	
Atopobium parvulum		41.93	0.25		99	99	
46100000	**Low discrimination**						**46100000**
					COC	SPO	
Clostridium botulinum biovar A		71.36	0.44	GLU 90	1	99	
Clostridium bifermentans		14.59	0.31	SUC 7	1	99	
Gemella morbillorum		7.87	0.30	MNS 7	99	1	
46101101	**Very good identification**						**46101101**
Clostridium chauvoei		100.0	0.57	GLU 99			
46101103	**Good identification**						**46101103**
Clostridium chauvoei		100.0	0.44	GLU 99			
46101120	**Good identification**						**46101120**
					GRA	SPO	
Prevotella melaninogenica		98.95	0.48	GLU 90	1	1	
Clostridium perfringens		0.59	0.17	RAF 90	99	84	
Clostridium chauvoei		0.46	0.14	FRU 90	99	99	
46101121	**Good identification**						**46101121**
					GRA	SPO	
Clostridium chauvoei		97.85	0.48	GLU 99	99	99	
Prevotella melaninogenica		2.15	0.14		1	1	
46101122	**Good identification**						**46101122**
					GRA	CAT	
Prevotella melaninogenica		95.23	0.36	GLU 90	1	1	
Capnocytophaga ochracea		2.48	0.12	RAF 90	1	90	
Clostridium perfringens		1.40	0.11	FRU 90	99	1	
46101123	**Good identification**						**46101123**
					GRA	SPO	
Clostridium chauvoei		97.68	0.36	GLU 99	99	99	
Prevotella melaninogenica		2.32	0.02		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
46110100	**Acceptable identification**							**46110100**
					GRA			
Fusobacterium mortiferum		92.40	0.37	GLU 99	1			
Actinomyces meyeri		3.85	0.05	MLT 11	99			
Prevotella melaninogenica		3.75	-.04		1			
46111101	**Very good identification**							**46111101**
					GRA	SPO		
Clostridium chauvoei		99.97	0.63	GLU 99	99	99		
Prevotella melaninogenica		0.03	-.04		1	1		
46111103	**Good identification**							**46111103**
Clostridium chauvoei		100.0	0.50	GLU 99				
46111120	**Good identification**							**46111120**
					GRA	CAT		
Prevotella melaninogenica		93.12	0.64	GLU 90	1	1		
Clostridium perfringens		6.09	0.51	RAF 90	99	1		
Capnocytophaga ochracea		0.57	0.30		1	90		
46111121	**Acceptable identification**							**46111121**
					GRA	SPO		
Clostridium chauvoei		91.72	0.55	GLU 99	99	99		
Prevotella melaninogenica		7.77	0.30		1	1		
Clostridium perfringens		0.51	0.17		99	84		
46111122	**Low discrimination**							**46111122**
					GRA	CAT		
Prevotella melaninogenica		66.21	0.52	GLU 90	1	1		
Capnocytophaga ochracea		18.98	0.47	RAF 90	1	90		
Clostridium perfringens		10.69	0.45		99	1		
46111123	**Low discrimination**							**46111123**
					GRA	CAT	SPO	
Clostridium chauvoei		87.91	0.42	GLU 99	99	1	99	
Prevotella melaninogenica		8.04	0.18		1	1	1	
Capnocytophaga ochracea		2.30	0.12		1	90	1	
46111160	**Low discrimination**							**46111160**
					GRA	CAT		
Clostridium perfringens		53.61	0.39	GLU 99	99	1		
Prevotella melaninogenica		43.49	0.30		1	1		
Capnocytophaga ochracea		2.90	0.14		1	90		
46115522	**Low discrimination**							**46115522**
					GRA	CAT		
Bifidobacterium longum subsp. longum		62.17	0.36	GLU 99	99	1		
Capnocytophaga ochracea		34.13	0.23	RAF 97	1	90		
Bacteroides fragilis		3.69	-.03		1	99		
46115530	**Good identification**							**46115530**
Bifidobacterium longum subsp. longum		100.0	0.40	GLU 99				
46115532	**Acceptable identification**							**46115532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		94.19	0.40	GLU 99	99	1		
Capnocytophaga ochracea		3.37	0.07	RAF 97	1	90		
Clostridium clostridioforme		2.44	0.11		50	33		
46115720	**Good identification**							**46115720**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		98.81	0.40	GLU 99	1		
Eubacterium contortum		1.19	0.14	RAF 97	99		
46121101	**Good identification**						**46121101**
Clostridium chauvoei		100.0	0.40	GLU 99			
46131020	**Good identification**						**46131020**
					COC		
Clostridium perfringens		95.78	0.41	GLU 99	1		
Lactobacillus jensenii		1.76	0.12	LAC 88	1		
Atopobium parvulum		0.96	0.06		99		
46131022	**Good identification**						**46131022**
Clostridium perfringens		97.47	0.35	GLU 99			
Lactobacillus jensenii		2.53	0.09	LAC 88			
46131100	**Acceptable identification**						**46131100**
					COC	GRA	
Atopobium parvulum		94.29	0.53	GLU 99	99	99	
Clostridium perfringens		3.81	0.30		1	99	
Fusobacterium mortiferum		1.14	0.32		1	1	
46131101	**Good identification**						**46131101**
					COC	SPO	
Clostridium chauvoei		97.69	0.47	GLU 99	1	99	
Atopobium parvulum		2.31	0.19	TRE 10	99	1	
46131102	**Low discrimination**						**46131102**
					COC	GRA	
Atopobium parvulum		89.32	0.40	GLU 99	99	99	
Clostridium perfringens		9.62	0.24		1	99	
Fusobacterium mortiferum		0.68	0.15		1	1	
46131120	**Low discrimination**						**46131120**
					COC	GRA	
Clostridium perfringens		81.48	0.56	GLU 99	1	99	
Atopobium parvulum		11.01	0.40		99	99	
Prevotella melaninogenica		6.78	0.30		1	1	
46131121	**Low discrimination**						**46131121**
					COC	SPO	
Clostridium chauvoei		89.87	0.38	GLU 99	1	99	
Clostridium perfringens		8.32	0.21	TRE 10	1	84	
Atopobium parvulum		1.12	0.06		99	1	
46131122	**Acceptable identification**						**46131122**
					COC	GRA	
Clostridium perfringens		90.43	0.50	GLU 99	1	99	
Atopobium parvulum		4.58	0.27		99	99	
Prevotella melaninogenica		3.05	0.18		1	1	
46131160	**Good identification**						**46131160**
					COC	GRA	
Clostridium perfringens		98.86	0.43	GLU 99	1	99	
Atopobium parvulum		0.71	0.06		99	99	
Prevotella melaninogenica		0.44	-.04		1	1	
46131162	**Good identification**						**46131162**
Clostridium perfringens		100.0	0.38	GLU 99			
46135100	**Very good identification**						**46135100**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				COC	GRA	
Atopobium parvulum	99.88	0.57	GLU 99	99	99	
Eubacterium saburreum	0.08	0.14		1	85	
Fusobacterium mortiferum	0.05	0.12		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46135102	**Good identification**						**46135102**
					COC		
Atopobium parvulum		99.63	0.45	GLU 99	99		
Bifidobacterium adolescentis		0.37	0.11		1		
46135120	**Good identification**						**46135120**
					COC	GRA	
Atopobium parvulum		95.82	0.45	GLU 99	99	99	
Clostridium perfringens		3.86	0.21		1	99	
Prevotella melaninogenica		0.32	-.04		1	1	
46135712	**Good identification**						**46135712**
Clostridium clostridioforme		100.0	0.41	GLU 88			
46135732	**Good identification**						**46135732**
Clostridium clostridioforme		96.06	0.41	GLU 88			
Parabacteroides distasonis		3.94	0.23	SAL 88			
46151000	**Low discrimination**						**46151000**
					GRA		
Lactobacillus jensenii		65.05	0.36	GLU 99	99		
Lactobacillus catenaformis		24.87	0.29	SAL 95	99		
Fusobacterium mortiferum		4.55	0.26		1		
46151120	**Low discrimination**						**46151120**
					GRA	SPO	
Prevotella melaninogenica		76.99	0.48	GLU 90	1	1	
Clostridium paraputrificum		12.45	0.32	RAF 90	99	99	
Collinsella aerofaciens		2.47	0.15	CEL 10	99	1	
46151122	**Low discrimination**						**46151122**
					CAT		
Prevotella oralis		40.42	0.43	GLU 99	1		
Prevotella melaninogenica		22.83	0.36	RAF 92	1		
Capnocytophaga ochracea		14.73	0.36		90		
46151322	**Acceptable identification**						**46151322**
					GRA	SPO	
Prevotella oralis		93.03	0.37	GLU 99	1	1	
Clostridium barati		4.48	0.14	RAF 92	99	99	
Prevotella melaninogenica		1.24	0.02		1	1	
46155712	**Good identification**						**46155712**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46155732	**Good identification**						**46155732**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46171000	**Acceptable identification**						**46171000**
					COC	GRA	
Lactobacillus jensenii		90.75	0.44	GLU 99	1	99	
Atopobium parvulum		8.62	0.25	SAL 95	99	99	
Fusobacterium mortiferum		0.63	0.17		1	1	
46171002	**Good identification**						**46171002**
					COC		
Lactobacillus jensenii		95.55	0.41	GLU 99	1		
Atopobium parvulum		2.40	0.12	SAL 95	99		
Lactobacillus acidophilus		1.52	0.07		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
46171010	**Good identification**						**46171010**
Lactobacillus jensenii		100.0	0.36	GLU 99			
46171100	**Very good identification**						**46171100**
					COC	GRA	
Atopobium parvulum		99.46	0.59	GLU 99	99	99	
Fusobacterium mortiferum		0.17	0.23		1	1	
Collinsella aerofaciens		0.15	-.03		1	99	
46171102	**Acceptable identification**						**46171102**
					COC	SPO	
Atopobium parvulum		91.27	0.46	GLU 99	99	1	
Lactobacillus acidophilus		6.68	0.25		1	1	
Clostridium barati		0.62	0.05		1	99	
46171120	**Low discrimination**						**46171120**
					COC	SPO	
Atopobium parvulum		72.60	0.46	GLU 99	99	1	
Clostridium perfringens		9.59	0.32		1	84	
Clostridium paraputrificum		9.06	0.22		1	99	
46174712	**Good identification**						**46174712**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46174732	**Good identification**						**46174732**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46175100	**Very good identification**						**46175100**
					COC	SPO	
Atopobium parvulum		99.92	0.64	GLU 99	99	1	
Actinomyces israelii		0.06	0.22		1	1	
Clostridium butyricum		0.02	-.07		1	99	
46175102	**Good identification**						**46175102**
					COC		
Atopobium parvulum		99.19	0.51	GLU 99	99		
Actinomyces israelii		0.48	0.25		1		
Bifidobacterium adolescentis		0.34	0.16		1		
46175120	**Good identification**						**46175120**
Atopobium parvulum		100.0	0.51	GLU 99			
46175122	**Good identification**						**46175122**
					COC	GRA	
Atopobium parvulum		95.15	0.38	GLU 99	99	99	
Prevotella oralis		2.24	0.07		1	1	
Parabacteroides distasonis		1.76	0.18		1	1	
46175512	**Low discrimination**						**46175512**
Clostridium clostridioforme		70.51	0.35	GLU 88			
Bifidobacterium adolescentis		29.49	0.32	SAL 88			
46175532	**Good identification**						**46175532**
Clostridium clostridioforme		98.42	0.35	GLU 88			
Parabacteroides distasonis		1.59	0.10	SAL 88			
46175612	**Good identification**						**46175612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.41	GLU 88			
46175632	**Good identification**						**46175632**
Clostridium clostridioforme		100.0	0.41	GLU 88			
46175702	**Good identification**						**46175702**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46175712	**Good identification**						**46175712**
Clostridium clostridioforme		100.0	0.50	GLU 88			
46175722	**Acceptable identification**						**46175722**
Clostridium clostridioforme		90.53	0.39	GLU 88			
Parabacteroides distasonis		5.05	0.23	SAL 88			
Bacteroides thetaiotaomicron		4.42	0.06	RAF 94			
46175732	**Good identification**						**46175732**
Clostridium clostridioforme		98.94	0.50	GLU 88			
Parabacteroides distasonis		0.65	0.18	SAL 88			
Bacteroides thetaiotaomicron		0.41	-.01	RAF 94			
46177102	**Low discrimination**						**46177102**
					COC	SPO	
Actinomyces israelii		49.00	0.38	GLU 99	1	1	
Clostridium tertium		27.70	0.19	SAL 92	1	99	
Atopobium parvulum		18.25	0.17		99	1	
46177502	**Low discrimination**						**46177502**
Actinomyces israelii		54.25	0.36	GLU 99			
Bifidobacterium adolescentis		45.75	0.29	SAL 92			
46210000	**Good identification**						**46210000**
					GRA		
Fusobacterium mortiferum		95.35	0.52	GLU 99	1		
Atopobium minutum		4.65	0.24		99		
46210002	**Low discrimination**						**46210002**
					GRA		
Fusobacterium mortiferum		77.38	0.35	GLU 99	1		
Atopobium minutum		22.62	0.21	BGL 10	99		
46210100	**Very good identification**						**46210100**
Fusobacterium mortiferum		100.0	0.58	GLU 99			
46210102	**Good identification**						**46210102**
Fusobacterium mortiferum		100.0	0.42	GLU 99			
46211000	**Good identification**						**46211000**
					GRA		
Fusobacterium mortiferum		95.78	0.55	GLU 99	1		
Lactobacillus jensenii		1.53	0.14		99		
Lactobacillus catenaformis		1.23	0.12		99		
46211002	**Low discrimination**						**46211002**
					GRA		
Fusobacterium mortiferum		74.16	0.38	GLU 99	1		
Lactobacillus catenaformis		8.60	0.12	BGL 10	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		7.10	0.11		99		
46211100	**Very good identification**						**46211100**
					GRA		
Fusobacterium mortiferum		98.31	0.61	GLU 99	1		
Lactobacillus catenaformis		0.81	0.15		99		
Collinsella aerofaciens		0.54	-.03		99		
46211102	**Low discrimination**						**46211102**
					GRA	SPO	
Fusobacterium mortiferum		81.81	0.45	GLU 99	1	1	
Clostridium barati		7.56	0.12	BGL 10	99	99	
Lactobacillus catenaformis		6.10	0.15		99	1	
46214100	**Good identification**						**46214100**
Fusobacterium mortiferum		100.0	0.39	GLU 99			
46215000	**Good identification**						**46215000**
Fusobacterium mortiferum		100.0	0.35	GLU 99			
46215100	**Good identification**						**46215100**
Fusobacterium mortiferum		100.0	0.42	GLU 99			
46230000	**Good identification**						**46230000**
Fusobacterium mortiferum		100.0	0.43	GLU 99			
46230100	**Good identification**						**46230100**
Fusobacterium mortiferum		100.0	0.49	GLU 99			
46231000	**Low discrimination**						**46231000**
					GRA		
Fusobacterium mortiferum		86.19	0.46	GLU 99	1		
Lactobacillus jensenii		13.81	0.22		99		
46231100	**Acceptable identification**						**46231100**
					COC	GRA	
Fusobacterium mortiferum		93.42	0.52	GLU 99	1	1	
Atopobium parvulum		6.58	0.21		99	99	
46231102	**Low discrimination**						**46231102**
					COC	GRA	
Fusobacterium mortiferum		83.13	0.36	GLU 99	1	1	
Atopobium parvulum		9.30	0.08	BGL 10	99	99	
Lactobacillus acidophilus		7.58	0.05		1	99	
46250000	**Low discrimination**						**46250000**
					GRA	SPO	
Fusobacterium mortiferum		86.98	0.43	GLU 99	1	1	
Clostridium barati		5.43	0.08		99	99	
Lactobacillus catenaformis		5.04	0.12		99	1	
46250100	**Low discrimination**						**46250100**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.50	GLU 99	1	1	
Clostridium barati		7.03	0.16		99	99	
Lactobacillus catenaformis		3.26	0.15		99	1	
46251000	**Low discrimination**						**46251000**
					GRA		
Lactobacillus catenaformis		56.58	0.47	GLU 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		14.79	0.46	MLT 90	1		
Lactobacillus jensenii		13.46	0.36		99		
46251002	**Low discrimination**						**46251002**
					SPO		
Lactobacillus catenaformis		53.38	0.47	GLU 99	1		
Clostridium barati		33.10	0.38	MLT 90	99		
Lactobacillus jensenii		8.46	0.33		1		
46251020	**Low discrimination**						**46251020**
					SPO		
Clostridium barati		66.13	0.38	GLU 99	99		
Lactobacillus catenaformis		26.66	0.36		1		
Lactobacillus jensenii		6.34	0.26		1		
46251022	**Low discrimination**						**46251022**
					SPO		
Clostridium barati		85.87	0.46	GLU 99	99		
Lactobacillus catenaformis		11.54	0.36		1		
Lactobacillus jensenii		1.83	0.23		1		
46251100	**Low discrimination**						**46251100**
					SPO		
Lactobacillus catenaformis		40.21	0.50	GLU 99	1		
Collinsella aerofaciens		26.60	0.32	MLT 90	1		
Clostridium barati		16.62	0.38		99		
46251102	**Low discrimination**						**46251102**
					SPO		
Clostridium barati		50.22	0.46	GLU 99	99		
Lactobacillus catenaformis		40.49	0.50		1		
Lactobacillus acidophilus		4.42	0.30		1		
46251120	**Low discrimination**						**46251120**
					SPO		
Clostridium barati		63.82	0.46	GLU 99	99		
Clostridium paraputrificum		19.06	0.32		99		
Lactobacillus catenaformis		12.86	0.39		1		
46251122	**Low discrimination**						**46251122**
					SPO		
Clostridium barati		87.29	0.54	GLU 99	99		
Lactobacillus catenaformis		5.87	0.39		1		
Clostridium paraputrificum		3.72	0.25		99		
46251162	**Acceptable identification**						**46251162**
					SPO		
Clostridium barati		91.56	0.36	GLU 99	99		
Lactobacillus acidophilus		3.82	0.15	SOR 8	1		
Clostridium paraputrificum		3.38	0.06		99		
46251322	**Acceptable identification**						**46251322**
					GRA	SPO	
Clostridium barati		94.65	0.36	GLU 99	99	99	
Lactobacillus acidophilus		2.40	0.11	RHA 8	99	1	
Prevotella oralis		1.74	0.07		1	1	
46253000	**Good identification**						**46253000**
					SPO		
Clostridium innocuum		99.35	0.56	GLU 99	99		
Clostridium barati		0.35	0.11		99		
Lactobacillus catenaformis		0.20	0.12		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
46253002	**Good identification**							**46253002**
					SPO			
Clostridium innocuum		98.76	0.56	GLU 99	99			
Clostridium barati		1.05	0.20		99			
Lactobacillus catenaformis		0.20	0.12		1			
46253122	**Acceptable identification**							**46253122**
					SPO			
Clostridium barati		94.29	0.36	GLU 99	99			
Lactobacillus acidophilus		1.67	0.08	MAN 8	1			
Clostridium tertium		1.58	0.09		99			
46253400	**Good identification**							**46253400**
Clostridium innocuum		100.0	0.43	GLU 99				
46253402	**Good identification**							**46253402**
Clostridium innocuum		100.0	0.43	GLU 99				
46270100	**Good identification**							**46270100**
					GRA	SPO		
Fusobacterium mortiferum		97.72	0.41	GLU 99	1	1		
Clostridium barati		2.28	-.02		99	99		
46271000	**Low discrimination**							**46271000**
					GRA	SPO		
Lactobacillus jensenii		85.17	0.44	GLU 99	99	1		
Fusobacterium mortiferum		9.32	0.38	MLT 95	1	1		
Clostridium innocuum		2.16	0.14		99	99		
46271002	**Low discrimination**							**46271002**
					SPO			
Lactobacillus jensenii		77.46	0.41	GLU 99	1			
Clostridium barati		8.78	0.20	MLT 95	99			
Lactobacillus acidophilus		7.76	0.21		1			
46271010	**Good identification**							**46271010**
Lactobacillus jensenii		100.0	0.36	GLU 99				
46271100	**Low discrimination**							**46271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		50.08	0.44	GLU 99	1	1	1	
Atopobium parvulum		24.69	0.27		99	99	1	
Clostridium barati		14.82	0.20		1	99	99	
46271102	**Low discrimination**							**46271102**
					GRA	SPO		
Lactobacillus acidophilus		70.85	0.39	GLU 99	99	1		
Clostridium barati		20.92	0.28	MLT 94	99	99		
Fusobacterium mortiferum		2.62	0.27		1	1		
46271122	**Low discrimination**							**46271122**
					SPO			
Lactobacillus acidophilus		49.22	0.39	GLU 99	1			
Clostridium barati		43.60	0.36	MLT 94	99			
Clostridium paraputrificum		5.69	0.16		99			
46273000	**Good identification**							**46273000**
					SPO			
Clostridium innocuum		99.60	0.48	GLU 99	99			
Lactobacillus jensenii		0.40	0.10		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46273002	**Good identification**						**46273002**
					SPO		
Clostridium innocuum		99.33	0.48	GLU 99	99		
Lactobacillus jensenii		0.27	0.07		1		
Clostridium barati		0.26	0.01		99		
46273400	**Good identification**						**46273400**
Clostridium innocuum		100.0	0.35	GLU 99			
46273402	**Good identification**						**46273402**
Clostridium innocuum		100.0	0.35	GLU 99			
46310000	**Good identification**						**46310000**
					GRA		
Fusobacterium mortiferum		96.42	0.36	GLU 99	1		
Lactobacillus jensenii		3.58	0.02	MLT 11	99		
46310100	**Acceptable identification**						**46310100**
					GRA		
Fusobacterium mortiferum		93.77	0.42	GLU 99	1		
Collinsella aerofaciens		6.23	-.03	MLT 11	99		
46311000	**Low discrimination**						**46311000**
					GRA		
Lactobacillus jensenii		53.34	0.36	GLU 99	99		
Fusobacterium mortiferum		21.74	0.39	CEL 95	1		
Lactobacillus catenaformis		20.39	0.29		99		
46311102	**Low discrimination**						**46311102**
Bifidobacterium longum subsp. infantis		71.65	0.40	GLU 99			
Lactobacillus catenaformis		8.76	0.32	RAF 95			
Collinsella aerofaciens		7.09	0.15				
46311122	**Low discrimination**						**46311122**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		57.19	0.44	GLU 99	99	1	
Prevotella oralis		16.76	0.38	RAF 95	1	1	
Clostridium barati		7.94	0.32		99	99	
46315102	**Acceptable identification**						**46315102**
Bifidobacterium longum subsp. infantis		92.69	0.38	GLU 99			
Bifidobacterium adolescentis		3.05	0.27	RAF 95			
Eubacterium contortum		2.34	0.16				
46315122	**Acceptable identification**						**46315122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		94.13	0.43	GLU 99	1	99	
Prevotella oralis		4.17	0.23	RAF 95	1	1	
Bacteroides fragilis		0.47	-.03		99	1	
46315300	**Very good identification**						**46315300**
Eubacterium contortum		100.0	0.63	GLU 99			
46315302	**Good identification**						**46315302**
Eubacterium contortum		99.40	0.50	GLU 99			
Bifidobacterium longum subsp. infantis		0.40	0.04				
Clostridium clostridioforme		0.20	0.09				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46315320	**Good identification**					**46315320**
					GRA	
Eubacterium contortum		98.72	0.50	GLU 99	99	
Bacteroides vulgatus		1.28	0.11		1	
46315322	**Low discrimination**					**46315322**
					GRA	
Eubacterium contortum		86.80	0.37	GLU 99	99	
Prevotella oralis		6.64	0.16		1	
Bifidobacterium longum subsp. infantis		3.54	0.08		99	
46315502	**Low discrimination**					**46315502**
Bifidobacterium adolescentis		84.22	0.41	GLU 99		
Eubacterium contortum		7.89	0.14	RAF 92		
Clostridium clostridioforme		4.03	0.15			
46315512	**Low discrimination**					**46315512**
Bifidobacterium adolescentis		84.85	0.42	GLU 99		
Clostridium clostridioforme		15.15	0.26	RAF 92		
46315700	**Very good identification**					**46315700**
					GRA	
Eubacterium contortum		99.96	0.61	GLU 99	99	
Bacteroides vulgatus		0.04	-.05		1	
46315702	**Good identification**					**46315702**
Eubacterium contortum		96.35	0.48	GLU 99		
Clostridium clostridioforme		3.65	0.30			
46315712	**Acceptable identification**					**46315712**
Clostridium clostridioforme		93.79	0.41	GLU 88		
Eubacterium contortum		5.49	0.14	RAF 94		
Bifidobacterium adolescentis		0.72	0.08	TRE 88		
46315720	**Low discrimination**					**46315720**
					GRA	
Eubacterium contortum		82.61	0.48	GLU 99	99	
Bacteroides vulgatus		17.39	0.30		1	
46315722	**Low discrimination**					**46315722**
					GRA	
Eubacterium contortum		80.36	0.36	GLU 99	99	
Clostridium clostridioforme		17.25	0.30		50	
Parabacteroides distasonis		1.42	0.17		1	
46315732	**Good identification**					**46315732**
					GRA	
Clostridium clostridioforme		98.03	0.41	GLU 88	50	
Eubacterium contortum		1.01	0.01	RAF 94	99	
Parabacteroides distasonis		0.96	0.12	TRE 88	1	
46317502	**Acceptable identification**					**46317502**
Bifidobacterium adolescentis		93.62	0.39	GLU 99		
Actinomyces israelii		6.39	0.25	RAF 92		
46317512	**Good identification**					**46317512**
Bifidobacterium adolescentis		100.0	0.41	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46331000	**Acceptable identification**						**46331000**
					COC	GRA	
Lactobacillus jensenii		90.90	0.44	GLU 99	1	99	
Atopobium parvulum		4.91	0.21	CEL 95	99	99	
Fusobacterium mortiferum		3.69	0.30		1	1	
46331002	**Good identification**						**46331002**
					COC		
Lactobacillus jensenii		95.92	0.41	GLU 99	1		
Lactobacillus acidophilus		1.52	0.07	CEL 95	1		
Atopobium parvulum		1.37	0.08		99		
46331010	**Good identification**						**46331010**
Lactobacillus jensenii		100.0	0.36	GLU 99			
46331100	**Good identification**						**46331100**
					COC	GRA	
Atopobium parvulum		96.98	0.55	GLU 99	99	99	
Fusobacterium mortiferum		1.72	0.36		1	1	
Clostridium perfringens		0.74	0.19		1	99	
46331102	**Low discrimination**						**46331102**
					COC		
Atopobium parvulum		82.29	0.42	GLU 99	99		
Lactobacillus acidophilus		10.61	0.25		1		
Clostridium perfringens		1.67	0.14		1		
46331120	**Low discrimination**						**46331120**
					COC	SPO	
Clostridium perfringens		49.58	0.45	GLU 99	1	84	
Atopobium parvulum		35.53	0.42		99	1	
Clostridium paraputrificum		7.80	0.22		1	99	
46331122	**Low discrimination**						**46331122**
					COC		
Clostridium perfringens		54.79	0.40	GLU 99	1		
Atopobium parvulum		14.72	0.29		99		
Lactobacillus acidophilus		10.76	0.25		1		
46331712	**Good identification**						**46331712**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46331732	**Good identification**						**46331732**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46334612	**Good identification**						**46334612**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46334632	**Good identification**						**46334632**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46334702	**Good identification**						**46334702**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46334712	**Good identification**						**46334712**
Clostridium clostridioforme		100.0	0.48	GLU 88			
46334722	**Good identification**						**46334722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.36	GLU 88			
46334732	**Good identification**						**46334732**
Clostridium clostridioforme		100.0	0.48	GLU 88			
46335100	**Very good identification**						**46335100**
					COC	GRA	
Atopobium parvulum		99.81	0.60	GLU 99	99	99	
Actinomyces israelii		0.09	0.21		1	99	
Fusobacterium mortiferum		0.07	0.17		1	1	
46335102	**Good identification**						**46335102**
					COC		
Atopobium parvulum		96.68	0.47	GLU 99	99		
Bifidobacterium adolescentis		1.55	0.23		1		
Bifidobacterium longum subsp. infantis		0.71	0.04		1		
46335120	**Good identification**						**46335120**
					COC		
Atopobium parvulum		99.25	0.47	GLU 99	99		
Clostridium perfringens		0.75	0.11		1		
46335312	**Good identification**						**46335312**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46335332	**Low discrimination**						**46335332**
Clostridium clostridioforme		86.60	0.35	GLU 88			
Parabacteroides distasonis		13.40	0.27	RAF 94			
46335502	**Low discrimination**						**46335502**
					COC		
Bifidobacterium adolescentis		57.34	0.38	GLU 99	1		
Clostridium clostridioforme		30.21	0.30	RAF 92	1		
Atopobium parvulum		5.40	0.13		99		
46335512	**Low discrimination**						**46335512**
Clostridium clostridioforme		66.26	0.41	GLU 88			
Bifidobacterium adolescentis		33.74	0.39	RAF 94			
46335532	**Good identification**						**46335532**
Clostridium clostridioforme		98.54	0.41	GLU 88			
Parabacteroides distasonis		1.46	0.15	RAF 94			
46335602	**Good identification**						**46335602**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46335612	**Good identification**						**46335612**
Clostridium clostridioforme		100.0	0.47	GLU 88			
46335622	**Good identification**						**46335622**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46335632	**Good identification**						**46335632**
Clostridium clostridioforme		100.0	0.47	GLU 88			
46335702	**Good identification**						**46335702**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		96.49	0.45	GLU 88		
Eubacterium contortum		3.51	0.14	RAF 94		
46335712	**Good identification**					**46335712**
Clostridium clostridioforme		100.0	0.56	GLU 88		
46335722	**Acceptable identification**					**46335722**
					GRA	
Clostridium clostridioforme		94.55	0.45	GLU 88	50	
Parabacteroides distasonis		4.84	0.28	RAF 94	1	
Eubacterium contortum		0.61	0.01		99	
46335732	**Good identification**					**46335732**
Clostridium clostridioforme		99.40	0.56	GLU 88		
Parabacteroides distasonis		0.60	0.23	RAF 94		
46337102	**Low discrimination**					**46337102**
					SPO	
Actinomyces israelii		45.14	0.37	GLU 99	1	
Clostridium tertium		29.48	0.19	CEL 93	99	
Bifidobacterium adolescentis		14.34	0.22		1	
46337502	**Low discrimination**					**46337502**
Bifidobacterium adolescentis		72.21	0.36	GLU 99		
Actinomyces israelii		27.79	0.35	RAF 92		
46337512	**Low discrimination**					**46337512**
Bifidobacterium adolescentis		81.80	0.38	GLU 99		
Bifidobacterium dentium		12.37	0.06	RAF 92		
Actinomyces israelii		3.85	0.21			
46350022	**Acceptable identification**					**46350022**
					SPO	
Clostridium barati		93.18	0.36	GLU 99	99	
Lactobacillus catenaformis		4.43	0.18	GAL 95	1	
Lactobacillus jensenii		1.48	0.10		1	
46350102	**Low discrimination**					**46350102**
					SPO	
Clostridium barati		63.02	0.36	GLU 99	99	
Lactobacillus catenaformis		17.98	0.32	GAL 95	1	
Collinsella aerofaciens		14.54	0.15		1	
46350122	**Low discrimination**					**46350122**
					SPO	
Clostridium barati		81.11	0.44	GLU 99	99	
Clostridium paraputrificum		13.47	0.25	GAL 95	99	
Lactobacillus catenaformis		1.93	0.21		1	
46351000	**Low discrimination**					**46351000**
					SPO	
Lactobacillus catenaformis		57.84	0.63	GLU 99	1	
Lactobacillus jensenii		29.04	0.58		1	
Clostridium barati		6.49	0.41		99	
46351002	**Low discrimination**					**46351002**
					SPO	
Lactobacillus catenaformis		58.49	0.63	GLU 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		19.68	0.50		99	
Lactobacillus jensenii		19.58	0.55		1	
46351010	**Acceptable identification**					**46351010**
					SPO	
Lactobacillus jensenii		93.13	0.50	GLU 99	1	
Lactobacillus catenaformis		5.62	0.29		1	
Clostridium barati		0.63	0.07		99	
46351012	**Low discrimination**					**46351012**
					SPO	
Lactobacillus jensenii		88.43	0.47	GLU 99	1	
Lactobacillus catenaformis		8.01	0.29		1	
Clostridium barati		2.69	0.15		99	
46351020	**Low discrimination**					**46351020**
					SPO	
Clostridium barati		42.74	0.50	GLU 99	99	
Lactobacillus catenaformis		31.76	0.53		1	
Lactobacillus jensenii		15.94	0.47		1	
46351022	**Low discrimination**					**46351022**
					SPO	
Clostridium barati		71.56	0.58	GLU 99	99	
Lactobacillus catenaformis		17.72	0.53		1	
Lactobacillus jensenii		5.93	0.44		1	
46351030	**Low discrimination**					**46351030**
					SPO	
Lactobacillus jensenii		86.47	0.39	GLU 99	1	
Clostridium barati		7.02	0.15		99	
Lactobacillus catenaformis		5.22	0.18		1	
46351032	**Low discrimination**					**46351032**
					SPO	
Lactobacillus jensenii		67.30	0.36	GLU 99	1	
Clostridium barati		24.60	0.24		99	
Lactobacillus catenaformis		6.09	0.18		1	
46351062	**Acceptable identification**					**46351062**
					SPO	
Clostridium barati		90.95	0.40	GLU 99	99	
Lactobacillus acidophilus		3.17	0.17	SOR 8	1	
Lactobacillus catenaformis		2.62	0.18		1	
46351100	**Low discrimination**					**46351100**
					SPO	
Collinsella aerofaciens		85.00	0.66	GLU 99	1	
Lactobacillus catenaformis		11.68	0.66		1	
Clostridium barati		2.62	0.50		99	
46351102	**Low discrimination**					**46351102**
					SPO	
Lactobacillus catenaformis		36.83	0.66	GLU 99	1	
Collinsella aerofaciens		29.78	0.50		1	
Clostridium barati		24.78	0.58		99	
46351120	**Low discrimination**					**46351120**
					SPO	
Clostridium paraputrificum		69.83	0.66	GLU 99	99	
Collinsella aerofaciens		13.86	0.50		1	
Clostridium barati		11.53	0.58		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46351122	**Low discrimination**						**46351122**
					SPO		
Clostridium barati		45.09	0.66	GLU 99	99		
Clostridium paraputrificum		39.00	0.60		99		
Lactobacillus catenaformis		5.58	0.56		1		
46351132	**Low discrimination**						**46351132**
					SPO		
Bifidobacterium breve		28.26	0.44	GLU 99	1		
Clostridium barati		27.93	0.32	RAF 95	99		
Clostridium paraputrificum		24.16	0.25		99		
46351142	**Low discrimination**						**46351142**
					SPO		
Clostridium barati		51.69	0.40	GLU 99	99		
Lactobacillus acidophilus		20.74	0.35	SOR 8	1		
Bifidobacterium breve		10.25	0.40		1		
46351152	**Low discrimination**						**46351152**
					GRA	SPO	
Bifidobacterium breve		89.30	0.40	GLU 99	99	1	
Clostridium barati		4.55	0.05	RAF 95	99	99	
Leptotrichia buccalis		4.33	-.03		1	1	
46351160	**Low discrimination**						**46351160**
					SPO		
Clostridium paraputrificum		81.03	0.47	GLU 99	99		
Clostridium barati		15.46	0.40	SOR 7	99		
Collinsella aerofaciens		2.16	0.15		1		
46351162	**Low discrimination**						**46351162**
					SPO		
Clostridium barati		49.30	0.48	GLU 99	99		
Clostridium paraputrificum		36.91	0.40	SOR 8	99		
Lactobacillus acidophilus		6.59	0.35		1		
46351172	**Low discrimination**						**46351172**
					SPO		
Bifidobacterium breve		86.08	0.44	GLU 99	1		
Clostridium barati		7.40	0.14	RAF 95	99		
Clostridium paraputrificum		5.54	0.06		99		
46351222	**Low discrimination**						**46351222**
					GRA	SPO	
Clostridium barati		88.18	0.40	GLU 99	99	99	
Prevotella oralis		6.29	0.21	RHA 8	1	1	
Lactobacillus catenaformis		2.54	0.18		99	1	
46351302	**Low discrimination**						**46351302**
					SPO		
Clostridium barati		61.79	0.40	GLU 99	99		
Lactobacillus acidophilus		15.10	0.31	RHA 8	1		
Lactobacillus catenaformis		10.67	0.32		1		
46351320	**Low discrimination**						**46351320**
					SPO		
Clostridium barati		50.72	0.40	GLU 99	99		
Clostridium paraputrificum		35.68	0.32	RHA 8	99		
Collinsella aerofaciens		7.08	0.15		1		
46351322	**Low discrimination**						**46351322**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		62.08	0.48	GLU 99	99	99	
Prevotella oralis		23.13	0.41	RHA 8	1	1	
Clostridium paraputrificum		6.24	0.25		99	99	
46353022	**Acceptable identification**						**46353022**
					SPO		
Clostridium barati		93.04	0.40	GLU 99	99		
Lactobacillus catenaformis		2.68	0.18	MAN 8	1		
Lactobacillus acidophilus		1.38	0.11		1		
46353102	**Low discrimination**						**46353102**
					SPO		
Clostridium barati		48.48	0.40	GLU 99	99		
Clostridium tertium		16.48	0.35	MAN 8	99		
Bifidobacterium breve		9.62	0.40		1		
46353112	**Low discrimination**						**46353112**
					SPO		
Clostridium tertium		72.16	0.39	GLU 99	99		
Bifidobacterium breve		23.69	0.40	TRE 90	1		
Bifidobacterium adolescentis		1.44	0.18		1		
46353120	**Low discrimination**						**46353120**
Clostridium barati		42.04	0.40	GLU 99			
Clostridium paraputrificum		29.57	0.32	MAN 8			
Clostridium tertium		14.29	0.35				
46353122	**Low discrimination**						**46353122**
Clostridium barati		55.67	0.48	GLU 99			
Clostridium tertium		18.92	0.43	MAN 8			
Clostridium ramosum		8.46	0.35				
46353130	**Acceptable identification**						**46353130**
					SPO		
Clostridium tertium		92.06	0.39	GLU 99	99		
Bifidobacterium breve		5.31	0.27	TRE 90	1		
Clostridium barati		1.54	0.05		99		
46353132	**Low discrimination**						**46353132**
					SPO		
Clostridium tertium		82.27	0.47	GLU 99	99		
Bifidobacterium breve		16.01	0.44	TRE 90	1		
Clostridium barati		1.38	0.14		99		
46353142	**Low discrimination**						**46353142**
					SPO		
Bifidobacterium breve		62.44	0.40	GLU 99	1		
Clostridium barati		27.37	0.21	RAF 95	99		
Lactobacillus acidophilus		6.65	0.13		1		
46353152	**Good identification**						**46353152**
					SPO		
Bifidobacterium breve		97.02	0.40	GLU 99	1		
Clostridium tertium		2.99	0.05	RAF 95	99		
46353162	**Low discrimination**						**46353162**
					SPO		
Bifidobacterium breve		52.46	0.44	GLU 99	1		
Clostridium barati		38.81	0.30	RAF 95	99		
Clostridium paraputrificum		3.38	0.06		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46353172	**Good identification**					**46353172**
					SPO	
Bifidobacterium breve		95.06	0.44	GLU 99	1	
Clostridium tertium		4.94	0.13	RAF 95	99	
46353322	**Low discrimination**					**46353322**
					SPO	
Clostridium ramosum		66.68	0.37	GLU 99	92	
Clostridium barati		28.79	0.30	TRE 94	99	
Lactobacillus acidophilus		1.42	0.09		1	
46354712	**Good identification**					**46354712**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46354732	**Good identification**					**46354732**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46355502	**Low discrimination**					**46355502**
					SPO	
Bifidobacterium adolescentis		78.70	0.46	GLU 99	1	
Clostridium butyricum		10.52	0.17	RAF 92	99	
Clostridium clostridioforme		5.94	0.24		66	
46355512	**Low discrimination**					**46355512**
Bifidobacterium adolescentis		77.25	0.48	GLU 99		
Clostridium clostridioforme		21.73	0.35	RAF 92		
Actinomyces israelii		0.58	0.18			
46355532	**Good identification**					**46355532**
Clostridium clostridioforme		96.53	0.35	GLU 88		
Bifidobacterium adolescentis		3.47	0.14	RAF 94		
46355612	**Good identification**					**46355612**
Clostridium clostridioforme		100.0	0.41	GLU 88		
46355632	**Good identification**					**46355632**
Clostridium clostridioforme		100.0	0.41	GLU 88		
46355702	**Acceptable identification**					**46355702**
Clostridium clostridioforme		91.08	0.39	GLU 88		
Eubacterium contortum		7.25	0.14	RAF 94		
Bifidobacterium adolescentis		1.66	0.12	TRE 88		
46355712	**Good identification**					**46355712**
Clostridium clostridioforme		99.51	0.50	GLU 88		
Bifidobacterium adolescentis		0.49	0.14	RAF 94		
46355722	**Low discrimination**					**46355722**
					GRA	
Clostridium clostridioforme		85.43	0.39	GLU 88	50	
Bacteroides uniformis		12.23	0.11	RAF 94	1	
Eubacterium contortum		1.20	0.01	TRE 88	99	
46355732	**Good identification**					**46355732**
Clostridium clostridioforme		100.0	0.50	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46357002	**Low discrimination**					**46357002**
					SPO	
Actinomyces israelii		83.34	0.36	GLU 99	1	
Clostridium tertium		6.85	0.03		99	
Clostridium innocuum		4.95	0.00		99	
46357102	**Low discrimination**					**46357102**
					SPO	
Clostridium tertium		60.45	0.37	GLU 99	99	
Actinomyces israelii		29.72	0.46	TRE 90	1	
Bifidobacterium adolescentis		8.52	0.31		1	
46357112	**Low discrimination**					**46357112**
					SPO	
Clostridium tertium		87.86	0.41	GLU 99	99	
Bifidobacterium adolescentis		8.51	0.32	TRE 90	1	
Actinomyces israelii		3.63	0.32		1	
46357120	**Acceptable identification**					**46357120**
					SPO	
Clostridium tertium		94.56	0.37	GLU 99	99	
Clostridium barati		2.04	0.05	TRE 90	99	
Actinomyces israelii		1.98	0.23		1	
46357122	**Good identification**					**46357122**
					SPO	
Clostridium tertium		95.95	0.45	GLU 99	99	
Clostridium barati		2.07	0.14	TRE 90	99	
Actinomyces israelii		1.00	0.26		1	
46357130	**Good identification**					**46357130**
Clostridium tertium		100.0	0.41	GLU 99		
46357132	**Good identification**					**46357132**
					SPO	
Clostridium tertium		99.86	0.50	GLU 99	99	
Bifidobacterium breve		0.14	0.10	TRE 90	1	
46357140	**Good identification**					**46357140**
Actinomyces israelii		100.0	0.36	GLU 99		
46357142	**Low discrimination**					**46357142**
					SPO	
Actinomyces israelii		87.42	0.39	GLU 99	1	
Bifidobacterium adolescentis		7.97	0.15		1	
Clostridium tertium		4.62	0.03		99	
46357500	**Low discrimination**					**46357500**
					SPO	
Actinomyces israelii		83.35	0.42	GLU 99	1	
Clostridium butyricum		13.69	0.07		99	
Bifidobacterium adolescentis		2.96	0.11		1	
46357502	**Low discrimination**					**46357502**
					SPO	
Bifidobacterium adolescentis		68.71	0.45	GLU 99	1	
Actinomyces israelii		29.30	0.45	RAF 92	1	
Clostridium butyricum		1.25	0.00		99	
46357512	**Low discrimination**					**46357512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		89.68	0.46	GLU 99			
Actinomyces israelii		4.68	0.31	RAF 92			
Bifidobacterium dentium		4.25	0.06				
46357542	**Low discrimination**						**46357542**
Actinomyces israelii		57.30	0.38	GLU 99			
Bifidobacterium adolescentis		42.70	0.29				
46370712	**Good identification**						**46370712**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46370732	**Good identification**						**46370732**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46371000	**Very good identification**						**46371000**
					COC		
Lactobacillus jensenii		97.87	0.66	GLU 99	1		
Lactobacillus catenaformis		0.66	0.29		1		
Atopobium parvulum		0.65	0.27		99		
46371002	**Low discrimination**						**46371002**
					SPO		
Lactobacillus jensenii		88.91	0.63	GLU 99	1		
Lactobacillus acidophilus		7.35	0.41		1		
Clostridium barati		2.59	0.31		99		
46371010	**Very good identification**						**46371010**
					GRA		
Lactobacillus jensenii		99.83	0.58	GLU 99	99		
Leptotrichia buccalis		0.17	-.03		1		
46371012	**Low discrimination**						**46371012**
					GRA		
Lactobacillus jensenii		79.81	0.55	GLU 99	99		
Leptotrichia buccalis		19.92	0.32		1		
Lactobacillus acidophilus		0.20	0.07		99		
46371020	**Low discrimination**						**46371020**
					SPO		
Lactobacillus jensenii		87.85	0.56	GLU 99	1		
Clostridium barati		6.83	0.31		99		
Clostridium paraputrificum		3.83	0.22		99		
46371022	**Low discrimination**						**46371022**
					SPO		
Lactobacillus jensenii		57.85	0.53	GLU 99	1		
Clostridium barati		20.23	0.40		99		
Lactobacillus acidophilus		19.12	0.41		1		
46371030	**Good identification**						**46371030**
					SPO		
Lactobacillus jensenii		99.77	0.47	GLU 99	1		
Clostridium barati		0.24	-.03		99		
46371032	**Low discrimination**						**46371032**
					GRA	SPO	
Lactobacillus jensenii		88.38	0.44	GLU 99	99	1	
Leptotrichia buccalis		9.80	0.15		1	1	
Clostridium barati		0.94	0.05		99	99	
46371100	**Low discrimination**						**46371100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Atopobium parvulum		82.89	0.61	GLU 99	99	1	
Collinsella aerofaciens		9.25	0.32		1	1	
Clostridium barati		2.45	0.31		1	99	
46371102	**Low discrimination**						**46371102**
					COC	SPO	
Lactobacillus acidophilus		69.97	0.60	GLU 99	1	1	
Atopobium parvulum		12.79	0.48		99	1	
Clostridium barati		6.44	0.40		1	99	
46371112	**Very good identification**						**46371112**
					GRA		
Leptotrichia buccalis		99.59	0.66	GLU 99	1		
Bifidobacterium breve		0.14	0.37		99		
Lactobacillus acidophilus		0.12	0.25		99		
46371120	**Low discrimination**						**46371120**
					COC	SPO	
Clostridium paraputrificum		81.40	0.56	GLU 99	1	99	
Atopobium parvulum		8.74	0.48		99	1	
Clostridium barati		4.40	0.40		1	99	
46371122	**Low discrimination**						**46371122**
					SPO		
Lactobacillus acidophilus		45.03	0.60	GLU 99	1		
Clostridium paraputrificum		32.87	0.50		99		
Clostridium barati		12.43	0.48		99		
46371132	**Acceptable identification**						**46371132**
					GRA	SPO	
Leptotrichia buccalis		94.12	0.50	GLU 99	1	1	
Bifidobacterium breve		2.11	0.41	NAG 10	99	1	
Clostridium tertium		1.48	0.29		99	99	
46371142	**Low discrimination**						**46371142**
					SPO		
Lactobacillus acidophilus		83.78	0.44	GLU 99	1		
Bifidobacterium breve		8.25	0.37	SOR 11	1		
Clostridium barati		5.42	0.21		99		
46371160	**Low discrimination**						**46371160**
					SPO		
Clostridium paraputrificum		89.63	0.37	GLU 99	99		
Clostridium barati		5.59	0.21	SOR 7	99		
Bifidobacterium breve		1.50	0.24		1		
46371162	**Low discrimination**						**46371162**
					SPO		
Lactobacillus acidophilus		54.67	0.44	GLU 99	1		
Clostridium paraputrificum		24.30	0.31	SOR 11	99		
Clostridium barati		10.62	0.30		99		
46371172	**Low discrimination**						**46371172**
					GRA		
Bifidobacterium breve		65.22	0.41	GLU 99	99		
Leptotrichia buccalis		29.34	0.15	RAF 95	1		
Lactobacillus acidophilus		3.77	0.10		99		
46371302	**Low discrimination**						**46371302**
					COC	SPO	
Lactobacillus acidophilus		84.41	0.40	GLU 99	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		8.97	0.21	RHA 7	1	99	
Atopobium parvulum		2.07	0.14		99	1	
46371322	**Low discrimination**						**46371322**
					SPO		
Clostridium ramosum		57.17	0.45	GLU 99	92		
Lactobacillus acidophilus		28.01	0.40		1		
Clostridium barati		8.93	0.30		99		
46371612	**Good identification**						**46371612**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46371632	**Good identification**						**46371632**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46371702	**Good identification**						**46371702**
Clostridium clostridioforme		96.33	0.36	GLU 88			
Lactobacillus acidophilus		3.67	0.06	RAF 94			
46371712	**Good identification**						**46371712**
Clostridium clostridioforme		99.04	0.48	GLU 88			
Leptotrichia buccalis		0.96	-.03	RAF 94			
46371722	**Low discrimination**						**46371722**
					SPO		
Clostridium clostridioforme		89.63	0.36	GLU 88	66		
Clostridium ramosum		6.96	0.11	RAF 94	92		
Lactobacillus acidophilus		3.41	0.06	XYL 91	1		
46371732	**Good identification**						**46371732**
Clostridium clostridioforme		100.0	0.48	GLU 88			
46373100	**Low discrimination**						**46373100**
					COC	SPO	
Clostridium tertium		80.30	0.43	GLU 99	1	99	
Atopobium parvulum		8.96	0.27		99	1	
Actinomyces israelii		3.96	0.35		1	1	
46373102	**Low discrimination**						**46373102**
					SPO		
Clostridium tertium		75.69	0.51	GLU 99	99		
Lactobacillus acidophilus		14.15	0.38		1		
Bifidobacterium breve		3.27	0.37		1		
46373110	**Good identification**						**46373110**
					GRA	SPO	
Clostridium tertium		98.36	0.47	GLU 99	99	99	
Bifidobacterium breve		0.71	0.20		99	1	
Leptotrichia buccalis		0.52	-.03		1	1	
46373112	**Low discrimination**						**46373112**
					GRA	SPO	
Clostridium tertium		83.06	0.55	GLU 99	99	99	
Leptotrichia buccalis		14.54	0.32		1	1	
Bifidobacterium breve		2.02	0.37		99	1	
46373120	**Low discrimination**						**46373120**
Clostridium tertium		59.03	0.51	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		34.44	0.45				
Clostridium paraputrificum		3.61	0.22				
46373122	**Low discrimination**						**46373122**
					SPO		
Clostridium tertium		52.50	0.59	GLU 99	99		
Clostridium ramosum		40.84	0.56		92		
Lactobacillus acidophilus		3.27	0.38		1		
46373130	**Good identification**						**46373130**
					SPO		
Clostridium tertium		99.25	0.55	GLU 99	99		
Bifidobacterium breve		0.42	0.24		1		
Clostridium ramosum		0.33	0.11		92		
46373132	**Very good identification**						**46373132**
					GRA	SPO	
Clostridium tertium		97.50	0.63	GLU 99	99	99	
Bifidobacterium breve		1.41	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
46373142	**Low discrimination**						**46373142**
					SPO		
Bifidobacterium breve		48.84	0.37	GLU 99	1		
Lactobacillus acidophilus		26.11	0.22	RAF 95	1		
Clostridium tertium		11.41	0.17		99		
46373152	**Low discrimination**						**46373152**
					GRA	SPO	
Bifidobacterium breve		65.74	0.37	GLU 99	99	1	
Clostridium tertium		27.31	0.21	RAF 95	99	99	
Leptotrichia buccalis		4.78	-.03		1	1	
46373162	**Low discrimination**						**46373162**
					SPO		
Bifidobacterium breve		47.08	0.41	GLU 99	1		
Clostridium tertium		18.56	0.25	RAF 95	99		
Clostridium ramosum		14.44	0.21		92		
46373172	**Low discrimination**						**46373172**
					SPO		
Bifidobacterium breve		58.79	0.41	GLU 99	1		
Clostridium tertium		41.21	0.29	RAF 95	99		
46373320	**Good identification**						**46373320**
Clostridium ramosum		98.40	0.47	GLU 99			
Clostridium tertium		1.29	0.17				
Clostridium barati		0.31	0.03				
46373322	**Very good identification**						**46373322**
					SPO		
Clostridium ramosum		98.35	0.58	GLU 99	92		
Clostridium tertium		0.96	0.25		99		
Lactobacillus acidophilus		0.45	0.18		1		
46373502	**Low discrimination**						**46373502**
					SPO		
Actinomyces israelii		42.18	0.36	GLU 99	1		
Bifidobacterium adolescentis		30.12	0.28	XYL 92	1		
Clostridium tertium		21.10	0.17		99		
46374312	**Good identification**						**46374312**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.36	GLU 88			
46374332	**Good identification**						**46374332**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46374512	**Good identification**						**46374512**
Clostridium clostridioforme		99.03	0.42	GLU 88			
Bifidobacterium adolescentis		0.97	0.11	RAF 94			
46374532	**Good identification**						**46374532**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46374602	**Good identification**						**46374602**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46374612	**Good identification**						**46374612**
Clostridium clostridioforme		100.0	0.48	GLU 88			
46374622	**Good identification**						**46374622**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46374632	**Good identification**						**46374632**
Clostridium clostridioforme		100.0	0.48	GLU 88			
46374702	**Good identification**						**46374702**
Clostridium clostridioforme		100.0	0.45	GLU 88			
46374712	**Very good identification**						**46374712**
Clostridium clostridioforme		100.0	0.57	GLU 88			
46374722	**Good identification**						**46374722**
Clostridium clostridioforme		100.0	0.45	GLU 88			
46374732	**Very good identification**						**46374732**
Clostridium clostridioforme		100.0	0.57	GLU 88			
46375100	**Very good identification**						**46375100**
					COC	SPO	
Atopobium parvulum		97.87	0.66	GLU 99	99	1	
Clostridium butyricum		1.52	0.28		1	99	
Actinomyces israelii		0.48	0.40		1	1	
46375102	**Low discrimination**						**46375102**
					COC	SPO	
Atopobium parvulum		87.56	0.53	GLU 99	99	1	
Actinomyces israelii		3.64	0.43		1	1	
Clostridium butyricum		2.99	0.21		1	99	
46375120	**Acceptable identification**						**46375120**
					COC	SPO	
Atopobium parvulum		92.89	0.53	GLU 99	99	1	
Clostridium paraputrificum		4.71	0.22		1	99	
Clostridium tertium		1.07	0.19		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46375122	**Low discrimination**						**46375122**
					COC	SPO	
Atopobium parvulum		60.16	0.40	GLU 99	99	1	
Clostridium tertium		11.83	0.27		1	99	
Lactobacillus acidophilus		10.14	0.25		1	1	
46375142	**Low discrimination**						**46375142**
					COC		
Actinomyces israelii		46.53	0.36	GLU 99	1		
Atopobium parvulum		29.11	0.19		99		
Lactobacillus acidophilus		10.60	0.10		1		
46375212	**Good identification**						**46375212**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46375232	**Good identification**						**46375232**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46375312	**Good identification**						**46375312**
Clostridium clostridioforme		98.52	0.44	GLU 88			
Leptotrichia buccalis		1.48	-.03	RAF 94			
46375332	**Good identification**						**46375332**
Clostridium clostridioforme		97.57	0.44	GLU 88			
Parabacteroides distasonis		2.43	0.23	RAF 94			
46375412	**Good identification**						**46375412**
Clostridium clostridioforme		97.81	0.41	GLU 88			
Actinomyces israelii		1.12	0.17	RAF 94			
Bifidobacterium adolescentis		1.07	0.11	RHA 88			
46375432	**Good identification**						**46375432**
Clostridium clostridioforme		100.0	0.41	GLU 88			
46375500	**Acceptable identification**						**46375500**
					COC	SPO	
Clostridium butyricum		90.63	0.44	GLU 99	1	99	
Atopobium parvulum		6.57	0.32	RAF 95	99	1	
Actinomyces israelii		2.60	0.39		1	1	
46375502	**Low discrimination**						**46375502**
					SPO		
Clostridium butyricum		57.94	0.37	GLU 99	99		
Bifidobacterium adolescentis		18.44	0.43	RAF 95	1		
Clostridium clostridioforme		15.31	0.39		66		
46375512	**Low discrimination**						**46375512**
					SPO		
Clostridium clostridioforme		71.10	0.50	GLU 88	66		
Bifidobacterium adolescentis		22.98	0.45	RAF 94	1		
Clostridium butyricum		3.11	0.15	RHA 88	99		
46375522	**Low discrimination**						**46375522**
					SPO		
Clostridium clostridioforme		70.92	0.39	GLU 88	66		
Clostridium butyricum		23.34	0.19	RAF 94	99		
Actinomyces israelii		1.90	0.21	RHA 88	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46375532	**Good identification**					**46375532**
					SPO	
Clostridium clostridioforme		95.89	0.50	GLU 88	66	
Bifidobacterium dentium		3.21	0.10	RAF 94	1	
Clostridium butyricum		0.37	-.03	RHA 88	99	
46375602	**Good identification**					**46375602**
Clostridium clostridioforme		100.0	0.45	GLU 88		
46375612	**Good identification**					**46375612**
Clostridium clostridioforme		100.0	0.56	GLU 88		
46375622	**Good identification**					**46375622**
Clostridium clostridioforme		100.0	0.45	GLU 88		
46375632	**Good identification**					**46375632**
Clostridium clostridioforme		100.0	0.56	GLU 88		
46375702	**Good identification**					**46375702**
					SPO	
Clostridium clostridioforme		97.86	0.54	GLU 88	66	
Clostridium butyricum		1.56	0.11	RAF 94	99	
Actinomyces israelii		0.42	0.22		1	
46375712	**Very good identification**					**46375712**
Clostridium clostridioforme		99.96	0.65	GLU 88		
Bifidobacterium adolescentis		0.04	0.11	RAF 94		
46375722	**Good identification**					**46375722**
Clostridium clostridioforme		98.40	0.54	GLU 88		
Parabacteroides distasonis		0.81	0.23	RAF 94		
Bacteroides ovatus		0.65	0.06			
46375732	**Very good identification**					**46375732**
Clostridium clostridioforme		99.81	0.65	GLU 88		
Parabacteroides distasonis		0.10	0.19	RAF 94		
Bacteroides ovatus		0.10	0.03			
46377000	**Acceptable identification**					**46377000**
					SPO	
Actinomyces israelii		91.05	0.43	GLU 99	1	
Clostridium tertium		8.95	0.11		99	
46377002	**Low discrimination**					**46377002**
					SPO	
Actinomyces israelii		83.57	0.46	GLU 99	1	
Clostridium tertium		16.43	0.19		99	
46377040	**Good identification**					**46377040**
Actinomyces israelii		100.0	0.36	GLU 99		
46377042	**Good identification**					**46377042**
Actinomyces israelii		100.0	0.39	GLU 99		
46377100	**Low discrimination**					**46377100**
					COC SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		63.03	0.45	GLU 99	1	99	
Actinomyces israelii		25.91	0.53		1	1	
Atopobium parvulum		9.46	0.32		99	1	
46377102	**Low discrimination**						**46377102**
					SPO		
Clostridium tertium		81.34	0.53	GLU 99	99		
Actinomyces israelii		16.72	0.56		1		
Bifidobacterium adolescentis		0.85	0.28		1		
46377110	**Good identification**						**46377110**
					SPO		
Clostridium tertium		96.66	0.50	GLU 99	99		
Actinomyces israelii		3.34	0.39		1		
46377112	**Very good identification**						**46377112**
					SPO		
Clostridium tertium		97.17	0.58	GLU 99	99		
Actinomyces israelii		1.68	0.42		1		
Bifidobacterium adolescentis		0.70	0.29		1		
46377120	**Good identification**						**46377120**
					COC	SPO	
Clostridium tertium		97.79	0.53	GLU 99	1	99	
Atopobium parvulum		0.86	0.19		99	1	
Actinomyces israelii		0.86	0.33		1	1	
46377122	**Very good identification**						**46377122**
					SPO		
Clostridium tertium		98.91	0.62	GLU 99	99		
Clostridium ramosum		0.56	0.21		92		
Actinomyces israelii		0.43	0.36		1		
46377130	**Very good identification**						**46377130**
					SPO		
Clostridium tertium		99.87	0.58	GLU 99	99		
Actinomyces israelii		0.07	0.18		1		
Bifidobacterium dentium		0.06	-.07		1		
46377132	**Very good identification**						**46377132**
					SPO		
Clostridium tertium		99.76	0.66	GLU 99	99		
Bifidobacterium dentium		0.20	0.10		1		
Actinomyces israelii		0.04	0.21		1		
46377140	**Acceptable identification**						**46377140**
					SPO		
Actinomyces israelii		92.84	0.46	GLU 99	1		
Clostridium tertium		5.87	0.11		99		
Clostridium butyricum		1.29	-.07		99		
46377142	**Low discrimination**						**46377142**
					SPO		
Actinomyces israelii		87.53	0.49	GLU 99	1		
Clostridium tertium		11.06	0.19		99		
Bifidobacterium adolescentis		1.41	0.12		1		
46377302	**Low discrimination**						**46377302**
					SPO		
Actinomyces israelii		60.50	0.37	GLU 99	1		
Clostridium tertium		39.50	0.19	RHA 7	99		
46377400	**Low discrimination**						**46377400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		89.77	0.41	GLU 99	1		
Clostridium butyricum		10.23	0.04		99		
46377402	**Good identification**						**46377402**
					SPO		
Actinomyces israelii		95.57	0.44	GLU 99	1		
Clostridium butyricum		2.82	-.03		99		
Bifidobacterium adolescentis		1.61	0.08		1		
46377442	**Good identification**						**46377442**
Actinomyces israelii		100.0	0.37	GLU 99			
46377500	**Low discrimination**						**46377500**
					SPO		
Actinomyces israelii		58.13	0.52	GLU 99	1		
Clostridium butyricum		39.76	0.28		99		
Clostridium tertium		1.75	0.11		99		
46377502	**Low discrimination**						**46377502**
					SPO		
Actinomyces israelii		59.06	0.55	GLU 99	1		
Bifidobacterium adolescentis		24.54	0.42		1		
Clostridium butyricum		10.47	0.21		99		
46377512	**Low discrimination**						**46377512**
Bifidobacterium dentium		81.86	0.40	GLU 99			
Bifidobacterium adolescentis		11.62	0.43	RAF 95			
Actinomyces israelii		3.42	0.40				
46377532	**Acceptable identification**						**46377532**
					SPO		
Bifidobacterium dentium		94.72	0.44	GLU 99	1		
Clostridium tertium		4.77	0.32	RAF 95	99		
Clostridium clostridioforme		0.29	0.16		66		
46377540	**Low discrimination**						**46377540**
					COC	SPO	
Actinomyces israelii		49.43	0.45	GLU 99	1	1	
Blautia producta		43.01	0.24		99	1	
Clostridium butyricum		7.56	0.10		1	99	
46377542	**Low discrimination**						**46377542**
					COC		
Actinomyces israelii		80.90	0.48	GLU 99	1		
Bifidobacterium adolescentis		10.68	0.26		1		
Blautia producta		5.21	0.07		99		
46377702	**Low discrimination**						**46377702**
					SPO		
Actinomyces israelii		64.79	0.35	GLU 99	1		
Clostridium clostridioforme		26.88	0.19	RHA 7	66		
Clostridium butyricum		4.72	-.05		99		
46410000	**Acceptable identification**						**46410000**
					GRA	CAT	
Fusobacterium mortiferum		92.23	0.52	GLU 99	1	5	
Atopobium minutum		4.49	0.24		99	1	
Propionibacterium granulosum		3.28	0.24		99	90	
46410002	**Low discrimination**						**46410002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Fusobacterium mortiferum		72.87	0.35	GLU 99	1	5	
Atopobium minutum		21.31	0.21	BGL 10	99	1	
Propionibacterium granulosum		5.82	0.13		99	90	
46410100	**Very good identification**						**46410100**
Fusobacterium mortiferum		100.0	0.58	GLU 99			
46410102	**Good identification**						**46410102**
Fusobacterium mortiferum		100.0	0.42	GLU 99			
46411000	**Good identification**						**46411000**
					GRA	CAT	
Fusobacterium mortiferum		96.67	0.55	GLU 99	1	5	
Propionibacterium granulosum		2.29	0.24		99	90	
Atopobium minutum		1.05	0.16		99	1	
46411002	**Low discrimination**						**46411002**
					GRA	CAT	
Fusobacterium mortiferum		89.43	0.38	GLU 99	1	5	
Atopobium minutum		5.81	0.13	BGL 10	99	1	
Propionibacterium granulosum		4.76	0.13		99	90	
46411100	**Very good identification**						**46411100**
					GRA		
Fusobacterium mortiferum		98.30	0.61	GLU 99	1		
Prevotella melaninogenica		1.53	0.12		1		
Lactobacillus fermentum		0.17	0.15		99		
46411102	**Low discrimination**						**46411102**
					CAT		
Fusobacterium mortiferum		87.95	0.45	GLU 99	5		
Capnocytophaga ochracea		7.38	0.12	BGL 10	90		
Prevotella melaninogenica		2.34	0.00		1		
46411120	**Good identification**						**46411120**
					CAT		
Prevotella melaninogenica		98.62	0.47	GLU 90	1		
Fusobacterium mortiferum		0.65	0.27	MLT 99	5		
Capnocytophaga ochracea		0.60	0.12		90		
46414100	**Good identification**						**46414100**
Fusobacterium mortiferum		100.0	0.39	GLU 99			
46415000	**Good identification**						**46415000**
Fusobacterium mortiferum		100.0	0.35	GLU 99			
46415100	**Good identification**						**46415100**
Fusobacterium mortiferum		100.0	0.42	GLU 99			
46415322	**Good identification**						**46415322**
Parabacteroides distasonis		96.81	0.45	GLU 99			
Bacteroides fragilis		3.19	-.03				
46415332	**Good identification**						**46415332**
Parabacteroides distasonis		100.0	0.40	GLU 99			
46415722	**Good identification**						**46415722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Parabacteroides distasonis		95.08	0.41	GLU 99	76		
Bacteroides vulgatus		3.28	-.03		3		
Bacteroides thetaiotaomicron		1.64	-.05		89		
46415732	**Acceptable identification**						**46415732**
Parabacteroides distasonis		92.09	0.36	GLU 99			
Clostridium clostridioforme		7.91	0.12				
46430000	**Low discrimination**						**46430000**
					GRA	CAT	
Fusobacterium mortiferum		73.71	0.43	GLU 99	1	5	
Propionibacterium granulosum		26.29	0.32		99	90	
46430100	**Good identification**						**46430100**
Fusobacterium mortiferum		100.0	0.49	GLU 99			
46431000	**Low discrimination**						**46431000**
					GRA	CAT	
Fusobacterium mortiferum		80.79	0.46	GLU 99	1	5	
Propionibacterium granulosum		19.21	0.32		99	90	
46431100	**Good identification**						**46431100**
Fusobacterium mortiferum		100.0	0.52	GLU 99			
46431102	**Good identification**						**46431102**
Fusobacterium mortiferum		100.0	0.36	GLU 99			
46431322	**Good identification**						**46431322**
Parabacteroides distasonis		100.0	0.40	GLU 99			
46431722	**Good identification**						**46431722**
Parabacteroides distasonis		100.0	0.36	GLU 99			
46435122	**Good identification**						**46435122**
Parabacteroides distasonis		97.88	0.48	GLU 99			
Bacteroides fragilis		2.12	-.03				
46435132	**Good identification**						**46435132**
Parabacteroides distasonis		100.0	0.43	GLU 99			
46435322	**Good identification**						**46435322**
Parabacteroides distasonis		100.0	0.56	GLU 99			
46435332	**Good identification**						**46435332**
Parabacteroides distasonis		99.46	0.51	GLU 99			
Clostridium clostridioforme		0.54	0.07				
46435522	**Acceptable identification**						**46435522**
					GRA	CAT	
Parabacteroides distasonis		94.34	0.44	GLU 99	1	76	
Bifidobacterium longum subsp. longum		4.59	0.17		99	1	
Bacteroides thetaiotaomicron		1.07	-.05		1	89	
46435532	**Low discrimination**						**46435532**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		82.62	0.39	GLU 99	1	
Bifidobacterium longum subsp. longum		12.70	0.21		99	
Clostridium clostridioforme		4.68	0.12		50	
46435722	**Acceptable identification**					**46435722**
Parabacteroides distasonis		93.76	0.52	GLU 99		
Bacteroides thetaiotaomicron		4.71	0.14			
Clostridium clostridioforme		1.53	0.16			
46435732	**Low discrimination**					**46435732**
Parabacteroides distasonis		85.14	0.48	GLU 99		
Clostridium clostridioforme		11.78	0.27			
Bacteroides thetaiotaomicron		3.09	0.07			
46450000	**Good identification**					**46450000**
Fusobacterium mortiferum		100.0	0.43	GLU 99		
46450100	**Good identification**					**46450100**
Fusobacterium mortiferum		100.0	0.50	GLU 99		
46451000	**Good identification**					**46451000**
Fusobacterium mortiferum		100.0	0.46	GLU 99		
46451100	**Good identification**					**46451100**
Fusobacterium mortiferum		99.49	0.53	GLU 99		
Prevotella melaninogenica		0.52	-.04			
46451102	**Low discrimination**					**46451102**
					CAT	
Fusobacterium mortiferum		85.66	0.36	GLU 99	5	
Prevotella oralis		8.95	0.06	BGL 10	1	
Capnocytophaga ochracea		5.39	0.02		90	
46455322	**Acceptable identification**					**46455322**
					CAT	
Parabacteroides distasonis		92.57	0.40	GLU 99	76	
Prevotella oralis		5.01	0.06		1	
Bacteroides fragilis		2.43	-.09		99	
46455332	**Good identification**					**46455332**
Parabacteroides distasonis		100.0	0.35	GLU 99		
46455722	**Low discrimination**					**46455722**
					CAT	
Parabacteroides distasonis		81.07	0.36	GLU 99	76	
Bacteroides uniformis		8.31	0.01		5	
Bacteroides thetaiotaomicron		5.51	0.00		89	
46470100	**Good identification**					**46470100**
Fusobacterium mortiferum		100.0	0.41	GLU 99		
46471000	**Good identification**					**46471000**
Fusobacterium mortiferum		100.0	0.38	GLU 99		
46471100	**Good identification**					**46471100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.44	GLU 99			
46475122	**Good identification**						**46475122**
Parabacteroides distasonis		98.30	0.43	GLU 99			
Bacteroides fragilis		1.70	-.09				
46475132	**Good identification**						**46475132**
Parabacteroides distasonis		100.0	0.39	GLU 99			
46475322	**Good identification**						**46475322**
Parabacteroides distasonis		99.60	0.51	GLU 99			
Clostridium clostridioforme		0.40	0.05				
46475332	**Good identification**						**46475332**
Parabacteroides distasonis		96.75	0.47	GLU 99			
Clostridium clostridioforme		3.26	0.16				
46475522	**Acceptable identification**						**46475522**
Parabacteroides distasonis		92.05	0.39	GLU 99			
Bacteroides thetaiotaomicron		4.12	0.00				
Clostridium clostridioforme		3.83	0.10				
46475712	**Good identification**						**46475712**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46475722	**Low discrimination**						**46475722**
Parabacteroides distasonis		76.93	0.48	GLU 99			
Bacteroides thetaiotaomicron		15.25	0.20				
Clostridium clostridioforme		7.82	0.25				
46475732	**Low discrimination**						**46475732**
Parabacteroides distasonis		49.92	0.43	GLU 99			
Clostridium clostridioforme		42.94	0.36				
Bacteroides thetaiotaomicron		7.15	0.13				
46501120	**Very good identification**						**46501120**
					GRA	CAT	
Prevotella melaninogenica		99.93	0.64	GLU 90	1	1	
Capnocytophaga ochracea		0.06	0.12	FRU 90	1	90	
Clostridium perfringens		0.01	0.04		99	1	
46501122	**Good identification**						**46501122**
					CAT		
Prevotella melaninogenica		96.66	0.52	GLU 90	1		
Capnocytophaga ochracea		2.52	0.29	FRU 90	90		
Prevotella oralis		0.62	0.18		1		
46501522	**Low discrimination**						**46501522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		65.24	0.37	GLU 99	99	1	
Prevotella melaninogenica		23.89	0.18	FRU 90	1	1	
Capnocytophaga ochracea		10.88	0.16		1	90	
46501530	**Good identification**						**46501530**
					GRA		
Bifidobacterium longum subsp. longum		98.87	0.41	GLU 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		1.13	-.04	FRU 90	1		
46501532	**Good identification**						**46501532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.92	0.41	GLU 99	99	1	
Capnocytophaga ochracea		1.08	-.01	FRU 90	1	90	
46505500	**Good identification**						**46505500**
Bifidobacterium longum subsp. longum		100.0	0.35	GLU 99			
46505502	**Good identification**						**46505502**
Bifidobacterium longum subsp. longum		96.73	0.35	GLU 99			
Bifidobacterium adolescentis		3.27	0.12	FRU 90			
46505510	**Good identification**						**46505510**
Bifidobacterium longum subsp. longum		100.0	0.39	GLU 99			
46505512	**Good identification**						**46505512**
Bifidobacterium longum subsp. longum		97.63	0.39	GLU 99			
Bifidobacterium adolescentis		2.37	0.13	FRU 90			
46505520	**Low discrimination**						**46505520**
					GRA		
Bifidobacterium longum subsp. longum		89.02	0.46	GLU 99	99		
Bacteroides vulgatus		10.40	0.15	FRU 90	1		
Prevotella melaninogenica		0.58	-.04		1		
46505522	**Good identification**						**46505522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.05	0.46	GLU 99	99	1	
Bacteroides fragilis		1.61	-.03	FRU 90	1	99	
Capnocytophaga ochracea		1.35	0.06		1	90	
46505530	**Good identification**						**46505530**
Bifidobacterium longum subsp. longum		100.0	0.50	GLU 99			
46505532	**Good identification**						**46505532**
Bifidobacterium longum subsp. longum		100.0	0.50	GLU 99			
46510100	**Low discrimination**						**46510100**
Fusobacterium mortiferum		84.15	0.42	GLU 99			
Prevotella melaninogenica		15.85	0.12	MLT 11			
46510120	**Good identification**						**46510120**
					GRA	CAT	
Prevotella melaninogenica		99.26	0.47	GLU 90	1	1	
Capnocytophaga ochracea		0.60	0.12	GAL 99	1	90	
Clostridium perfringens		0.14	0.04		99	1	
46511020	**Good identification**						**46511020**
					GRA	CAT	
Prevotella melaninogenica		97.16	0.47	GLU 90	1	1	
Clostridium perfringens		1.82	0.24	LAC 99	99	1	
Capnocytophaga ochracea		0.59	0.12		1	90	
46511100	**Low discrimination**						**46511100**
					CAT		
Prevotella melaninogenica		85.40	0.47	GLU 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		6.87	0.45	NAG 99	5			
Capnocytophaga ochracea		5.70	0.30		90			
46511102	**Low discrimination**							**46511102**
					GRA	CAT		
Capnocytophaga ochracea		62.52	0.47	GLU 99	1	90		
Prevotella melaninogenica		19.82	0.34	NAG 90	1	1		
Bifidobacterium longum subsp. infantis		7.92	0.27		99	1		
46511120	**Very good identification**							**46511120**
					GRA	CAT		
Prevotella melaninogenica		99.15	0.81	GLU 90	1	1		
Capnocytophaga ochracea		0.60	0.47		1	90		
Clostridium perfringens		0.14	0.39		99	1		
46511121	**Good identification**							**46511121**
					GRA	CAT	SPO	
Prevotella melaninogenica		97.78	0.47	GLU 90	1	1	1	
Clostridium chauvoei		1.30	0.21	URE 1	99	1	99	
Capnocytophaga ochracea		0.59	0.12		1	90	1	
46511122	**Low discrimination**							**46511122**
					CAT			
Prevotella melaninogenica		72.24	0.68	GLU 90	1			
Capnocytophaga ochracea		20.71	0.63		90			
Prevotella oralis		5.12	0.52		1			
46511130	**Acceptable identification**							**46511130**
					GRA	CAT		
Prevotella melaninogenica		92.84	0.47	GLU 90	1	1		
Capnocytophaga ochracea		6.20	0.30	MLZ 1	1	90		
Bifidobacterium longum subsp. longum		0.83	0.24		99	1		
46511132	**Low discrimination**							**46511132**
					CAT			
Capnocytophaga ochracea		72.90	0.47	GLU 99	90			
Prevotella melaninogenica		23.12	0.34	MLZ 10	1			
Prevotella oralis		1.64	0.18		1			
46511160	**Acceptable identification**							**46511160**
					GRA	CAT		
Prevotella melaninogenica		91.51	0.47	GLU 90	1	1		
Capnocytophaga ochracea		6.11	0.30	SOR 1	1	90		
Clostridium perfringens		2.39	0.26		99	1		
46511162	**Low discrimination**							**46511162**
					CAT			
Capnocytophaga ochracea		73.15	0.47	GLU 99	90			
Prevotella melaninogenica		23.20	0.34	SOR 10	1			
Prevotella oralis		1.64	0.18		1			
46511320	**Acceptable identification**							**46511320**
Prevotella melaninogenica		93.87	0.47	GLU 90				
Prevotella oralis		2.83	0.24	RHA 1				
Bacteroides vulgatus		2.60	0.18					
46511322	**Low discrimination**							**46511322**
					CAT			
Prevotella oralis		66.89	0.46	GLU 99	1			
Prevotella melaninogenica		22.24	0.34		1			
Capnocytophaga ochracea		6.38	0.29		90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46511500	**Low discrimination**						**46511500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.26	0.43	GLU 99	99	1	
Capnocytophaga ochracea		7.54	0.17		1	90	
Prevotella melaninogenica		6.47	0.12		1	1	
46511502	**Low discrimination**						**46511502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		49.28	0.43	GLU 99	99	1	
Capnocytophaga ochracea		40.17	0.34		1	90	
Bifidobacterium adolescentis		8.22	0.32		99	1	
46511510	**Good identification**						**46511510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.41	0.47	GLU 99	99	1	
Capnocytophaga ochracea		0.59	0.01		1	90	
46511512	**Low discrimination**						**46511512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		84.19	0.47	GLU 99	99	1	
Bifidobacterium adolescentis		10.08	0.33		99	1	
Capnocytophaga ochracea		4.48	0.17		1	90	
46511520	**Low discrimination**						**46511520**
					GRA	CAT	
Prevotella melaninogenica		59.89	0.47	GLU 90	1	1	
Bifidobacterium longum subsp. longum		31.16	0.54	ARA 1	99	1	
Capnocytophaga ochracea		6.35	0.34		1	90	
46511522	**Low discrimination**						**46511522**
					GRA	CAT	
Capnocytophaga ochracea		56.43	0.50	GLU 99	1	90	
Bifidobacterium longum subsp. longum		30.77	0.54		99	1	
Prevotella melaninogenica		11.27	0.34		1	1	
46511530	**Very good identification**						**46511530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.59	0.58	GLU 99	99	1	
Capnocytophaga ochracea		1.30	0.17		1	90	
Prevotella melaninogenica		1.11	0.12		1	1	
46511532	**Low discrimination**						**46511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.51	0.58	GLU 99	99	1	
Capnocytophaga ochracea		10.59	0.34		1	90	
Parabacteroides distasonis		0.38	0.21		1	76	
46511720	**Good identification**						**46511720**
					GRA		
Bacteroides vulgatus		95.73	0.37	GLU 99	1		
Prevotella melaninogenica		2.62	0.12	XYL 98	1		
Bifidobacterium longum subsp. longum		1.37	0.20		99		
46513120	**Good identification**						**46513120**
					GRA	CAT	
Prevotella melaninogenica		99.00	0.47	GLU 90	1	1	
Capnocytophaga ochracea		0.60	0.12	MAN 1	1	90	
Actinomyces naeslundii		0.26	0.32		99	25	
46514712	**Good identification**						**46514712**
Clostridium clostridioforme		100.0	0.38	GLU 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46514732	**Good identification**						**46514732**
Clostridium clostridioforme		98.75	0.38	GLU 88			
Parabacteroides distasonis		1.25	0.12	SAL 88			
46515120	**Low discrimination**						**46515120**
					CAT		
Bacteroides fragilis		52.49	0.44	GLU 99	99		
Prevotella melaninogenica		35.48	0.47	BGL 95	1		
Bacteroides vulgatus		5.36	0.30		3		
46515122	**Good identification**						**46515122**
					CAT		
Bacteroides fragilis		93.55	0.66	GLU 99	99		
Capnocytophaga ochracea		4.50	0.53		90		
Prevotella melaninogenica		0.63	0.34		1		
46515320	**Good identification**						**46515320**
					CAT		
Bacteroides vulgatus		97.29	0.47	GLU 99	3		
Bacteroides fragilis		1.07	0.10	ARA 93	99		
Prevotella melaninogenica		0.72	0.12		1		
46515322	**Low discrimination**						**46515322**
					CAT		
Parabacteroides distasonis		44.23	0.54	GLU 99	76		
Bacteroides fragilis		40.67	0.32		99		
Prevotella oralis		10.14	0.30		1		
46515332	**Low discrimination**						**46515332**
Parabacteroides distasonis		89.27	0.49	GLU 99			
Clostridium clostridioforme		8.38	0.26				
Bacteroides fragilis		1.54	-.03				
46515500	**Acceptable identification**						**46515500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		93.59	0.52	GLU 99	99	1	
Bacteroides vulgatus		4.86	0.15		1	3	
Capnocytophaga ochracea		0.71	0.07		1	90	
46515502	**Low discrimination**						**46515502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		67.97	0.52	GLU 99	99	1	
Bifidobacterium adolescentis		25.29	0.46		99	1	
Capnocytophaga ochracea		4.62	0.23		1	90	
46515510	**Good identification**						**46515510**
Bifidobacterium longum subsp. longum		99.73	0.56	GLU 99			
Bifidobacterium adolescentis		0.27	0.13				
46515512	**Low discrimination**						**46515512**
Bifidobacterium longum subsp. longum		75.66	0.56	GLU 99			
Bifidobacterium adolescentis		20.21	0.47				
Clostridium clostridioforme		3.80	0.32				
46515520	**Low discrimination**						**46515520**
					GRA	CAT	
Bacteroides vulgatus		55.44	0.50	GLU 99	1	3	
Bifidobacterium longum subsp. longum		43.13	0.62	RHA 90	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		0.73	0.23		1	90	
46515522	**Low discrimination**						**46515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		71.56	0.62	GLU 99	99	1	
Bacteroides fragilis		13.02	0.32		1	99	
Capnocytophaga ochracea		10.94	0.40		1	90	
46515530	**Very good identification**						**46515530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.13	0.66	GLU 99	99	1	
Bacteroides vulgatus		0.76	0.15		1	3	
Capnocytophaga ochracea		0.11	0.07		1	90	
46515532	**Very good identification**						**46515532**
					GRA		
Bifidobacterium longum subsp. longum		96.43	0.66	GLU 99	99		
Parabacteroides distasonis		1.23	0.38		1		
Clostridium clostridioforme		1.21	0.32		50		
46515612	**Good identification**						**46515612**
Clostridium clostridioforme		100.0	0.38	GLU 88			
46515632	**Good identification**						**46515632**
Clostridium clostridioforme		98.61	0.38	GLU 88			
Parabacteroides distasonis		1.39	0.12	SAL 88			
46515702	**Low discrimination**						**46515702**
Clostridium clostridioforme		80.78	0.35	GLU 88			
Bifidobacterium longum subsp. longum		5.93	0.17	SAL 88			
Eubacterium contortum		5.43	0.10	TRE 88			
46515712	**Good identification**						**46515712**
Clostridium clostridioforme		96.25	0.47	GLU 88			
Bifidobacterium longum subsp. longum		2.64	0.21	SAL 88			
Bifidobacterium adolescentis		0.71	0.13	TRE 88			
46515720	**Very good identification**						**46515720**
					GRA		
Bacteroides vulgatus		99.88	0.66	GLU 99	1		
Bifidobacterium longum subsp. longum		0.09	0.28		99		
Eubacterium contortum		0.02	0.10		99		
46515722	**Low discrimination**						**46515722**
Parabacteroides distasonis		41.08	0.50	GLU 99			
Bacteroides vulgatus		39.58	0.32				
Clostridium clostridioforme		11.65	0.35				
46515732	**Low discrimination**						**46515732**
					GRA		
Clostridium clostridioforme		64.23	0.47	GLU 88	50		
Parabacteroides distasonis		26.77	0.46	SAL 88	1		
Bifidobacterium longum subsp. longum		7.05	0.32	TRE 88	99		
46517502	**Good identification**						**46517502**
Bifidobacterium adolescentis		95.77	0.44	GLU 99			
Bifidobacterium longum subsp. longum		3.18	0.17				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.05	0.17				
46517512	**Good identification**						**46517512**
Bifidobacterium adolescentis		95.58	0.46	GLU 99			
Bifidobacterium longum subsp. longum		4.42	0.21				
46525520	**Good identification**						**46525520**
					GRA		
Bifidobacterium longum subsp. longum		98.09	0.35	GLU 99	99		
Bacteroides vulgatus		1.91	-.08	FRU 90	1		
46525522	**Acceptable identification**						**46525522**
					GRA		
Bifidobacterium longum subsp. longum		93.72	0.35	GLU 99	99		
Parabacteroides distasonis		6.28	0.19	FRU 90	1		
46525530	**Good identification**						**46525530**
Bifidobacterium longum subsp. longum		100.0	0.39	GLU 99			
46525532	**Acceptable identification**						**46525532**
					GRA		
Bifidobacterium longum subsp. longum		94.82	0.39	GLU 99	99		
Clostridium clostridioforme		3.17	0.12	FRU 90	50		
Parabacteroides distasonis		2.01	0.15		1		
46530712	**Good identification**						**46530712**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46530732	**Good identification**						**46530732**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46531100	**Low discrimination**						**46531100**
					COC	GRA	
Fusobacterium mortiferum		42.60	0.36	GLU 99	1	1	
Atopobium parvulum		18.31	0.19	MLT 11	99	99	
Prevotella melaninogenica		17.90	0.12		1	1	
46531120	**Low discrimination**						**46531120**
					GRA	CAT	
Prevotella melaninogenica		78.70	0.47	GLU 90	1	1	
Clostridium perfringens		20.02	0.43	TRE 1	99	1	
Capnocytophaga ochracea		0.48	0.12		1	90	
46531132	**Low discrimination**						**46531132**
					GRA	CAT	
Parabacteroides distasonis		66.96	0.36	GLU 99	1	76	
Capnocytophaga ochracea		14.48	0.12	XYL 90	1	90	
Bifidobacterium longum subsp. longum		5.34	0.13		99	1	
46531322	**Low discrimination**						**46531322**
Parabacteroides distasonis		78.75	0.49	GLU 99			
Prevotella oralis		18.42	0.25	XYL 90			
Clostridium clostridioforme		1.25	0.13				
46531332	**Low discrimination**						**46531332**
Parabacteroides distasonis		88.14	0.44	GLU 99			
Clostridium clostridioforme		11.86	0.24	XYL 90			
46531510	**Good identification**						**46531510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		100.0	0.37	GLU 99			
46531512	**Low discrimination**						**46531512**
Bifidobacterium longum subsp. longum		56.94	0.37	GLU 99			
Clostridium clostridioforme		24.88	0.29				
Bifidobacterium adolescentis		18.18	0.30				
46531520	**Low discrimination**						**46531520**
					GRA		
Bifidobacterium longum subsp. longum		89.38	0.43	GLU 99	99		
Prevotella melaninogenica		6.94	0.12		1		
Clostridium perfringens		1.77	0.09		99		
46531522	**Low discrimination**						**46531522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		71.88	0.43	GLU 99	99	1	
Parabacteroides distasonis		17.65	0.37		1	76	
Capnocytophaga ochracea		5.33	0.16		1	90	
46531530	**Good identification**						**46531530**
Bifidobacterium longum subsp. longum		100.0	0.47	GLU 99			
46531532	**Low discrimination**						**46531532**
					GRA		
Bifidobacterium longum subsp. longum		83.91	0.47	GLU 99	99		
Clostridium clostridioforme		9.17	0.29		50		
Parabacteroides distasonis		6.52	0.33		1		
46531612	**Good identification**						**46531612**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46531632	**Good identification**						**46531632**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46531712	**Good identification**						**46531712**
Clostridium clostridioforme		100.0	0.44	GLU 88			
46531722	**Low discrimination**						**46531722**
Parabacteroides distasonis		66.52	0.45	GLU 99			
Clostridium clostridioforme		27.03	0.33	XYL 90			
Bacteroides thetaiotaomicron		5.54	0.11				
46531732	**Low discrimination**						**46531732**
Clostridium clostridioforme		75.71	0.44	GLU 88			
Parabacteroides distasonis		22.02	0.41	SAL 88			
Bacteroides thetaiotaomicron		1.32	0.04	XYL 91			
46533100	**Good identification**						**46533100**
Actinomyces naeslundii		100.0	0.37	ESL 12			
46533101	**Good identification**						**46533101**
Actinomyces naeslundii		100.0	0.40	ESL 12			
46533102	**Low discrimination**						**46533102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		74.41	0.37	ESL 12		
Bifidobacterium adolescentis		25.59	0.13	GLU 99		
46533103	**Good identification**					**46533103**
Actinomyces naeslundii		100.0	0.40	ESL 12		
46533121	**Good identification**					**46533121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
46533122	**Low discrimination**					**46533122**
					GRA	
Actinomyces naeslundii		56.28	0.37	ESL 12	99	
Prevotella melaninogenica		21.05	0.00	GLU 99	1	
Clostridium perfringens		13.23	0.03		99	
46533123	**Good identification**					**46533123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
46534512	**Good identification**					**46534512**
Clostridium clostridioforme		96.60	0.38	GLU 88		
Bifidobacterium longum subsp. longum		1.99	0.11	SAL 88		
Bifidobacterium adolescentis		1.42	0.10	RHA 88		
46534532	**Acceptable identification**					**46534532**
					GRA	
Clostridium clostridioforme		90.79	0.38	GLU 88	50	
Bifidobacterium longum subsp. longum		7.47	0.21	SAL 88	99	
Parabacteroides distasonis		1.74	0.15	RHA 88	1	
46534612	**Good identification**					**46534612**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46534632	**Good identification**					**46534632**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46534702	**Good identification**					**46534702**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46534712	**Good identification**					**46534712**
Clostridium clostridioforme		100.0	0.53	GLU 88		
46534722	**Acceptable identification**					**46534722**
Clostridium clostridioforme		92.50	0.42	GLU 88		
Parabacteroides distasonis		6.14	0.27	SAL 88		
Bacteroides thetaiotaomicron		1.36	0.00			
46534732	**Good identification**					**46534732**
Clostridium clostridioforme		99.22	0.53	GLU 88		
Parabacteroides distasonis		0.78	0.23	SAL 88		
46535122	**Low discrimination**					**46535122**
Parabacteroides distasonis		59.46	0.57	GLU 99		
Bacteroides fragilis		36.00	0.32			
Capnocytophaga ochracea		1.73	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46535132	**Acceptable identification**					**46535132**
					GRA	
Parabacteroides distasonis		90.44	0.53	GLU 99	1	
Clostridium clostridioforme		5.59	0.26		50	
Bifidobacterium longum subsp. longum		2.40	0.21		99	
46535312	**Acceptable identification**					**46535312**
Clostridium clostridioforme		93.74	0.41	GLU 88		
Parabacteroides distasonis		6.27	0.27	SAL 88		
46535322	**Very good identification**					**46535322**
Parabacteroides distasonis		97.61	0.65	GLU 99		
Clostridium clostridioforme		1.74	0.30			
Prevotella oralis		0.31	0.10			
46535332	**Low discrimination**					**46535332**
Parabacteroides distasonis		86.87	0.61	GLU 99		
Clostridium clostridioforme		13.13	0.41			
46535412	**Good identification**					**46535412**
Clostridium clostridioforme		96.22	0.38	GLU 88		
Bifidobacterium longum subsp. longum		2.21	0.11	SAL 88		
Bifidobacterium adolescentis		1.57	0.10	RHA 88		
46535432	**Low discrimination**					**46535432**
					GRA	
Clostridium clostridioforme		89.83	0.38	GLU 88	50	
Bifidobacterium longum subsp. longum		8.25	0.21	SAL 88	99	
Parabacteroides distasonis		1.92	0.15	RHA 88	1	
46535500	**Good identification**					**46535500**
					SPO	
Bifidobacterium longum subsp. longum		96.62	0.41	GLU 99	1	
Clostridium butyricum		1.57	-.03		99	
Bifidobacterium adolescentis		0.97	0.09		1	
46535502	**Low discrimination**					**46535502**
Bifidobacterium longum subsp. longum		38.74	0.41	GLU 99		
Bifidobacterium adolescentis		38.44	0.43			
Clostridium clostridioforme		21.32	0.35			
46535510	**Good identification**					**46535510**
Bifidobacterium longum subsp. longum		97.84	0.45	GLU 99		
Clostridium clostridioforme		1.46	0.12			
Bifidobacterium adolescentis		0.70	0.10			
46535512	**Low discrimination**					**46535512**
Clostridium clostridioforme		46.10	0.47	GLU 88		
Bifidobacterium longum subsp. longum		31.31	0.45	SAL 88		
Bifidobacterium adolescentis		22.30	0.44	RHA 88		
46535520	**Low discrimination**					**46535520**
					GRA	
Bifidobacterium longum subsp. longum		81.86	0.52	GLU 99	99	
Bacteroides vulgatus		17.54	0.26		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		0.61	0.19		1	
46535522	**Low discrimination**					**46535522**
					GRA	
Bifidobacterium longum subsp. longum		51.83	0.52	GLU 99	99	
Parabacteroides distasonis		38.18	0.53		1	
Clostridium clostridioforme		7.13	0.35		50	
46535530	**Good identification**					**46535530**
					GRA	
Bifidobacterium longum subsp. longum		99.27	0.56	GLU 99	99	
Clostridium clostridioforme		0.37	0.12		50	
Parabacteroides distasonis		0.23	0.15		1	
46535532	**Low discrimination**					**46535532**
					GRA	
Bifidobacterium longum subsp. longum		62.06	0.56	GLU 99	99	
Clostridium clostridioforme		22.85	0.47		50	
Parabacteroides distasonis		14.46	0.49		1	
46535602	**Good identification**					**46535602**
Clostridium clostridioforme		100.0	0.41	GLU 88		
46535612	**Good identification**					**46535612**
Clostridium clostridioforme		100.0	0.52	GLU 88		
46535622	**Acceptable identification**					**46535622**
Clostridium clostridioforme		90.31	0.41	GLU 88		
Parabacteroides distasonis		6.69	0.27	SAL 88		
Bacteroides thetaiotaomicron		3.00	0.06			
46535632	**Good identification**					**46535632**
Clostridium clostridioforme		98.85	0.52	GLU 88		
Parabacteroides distasonis		0.87	0.23	SAL 88		
Bacteroides thetaiotaomicron		0.28	-.02			
46535702	**Good identification**					**46535702**
Clostridium clostridioforme		96.90	0.50	GLU 88		
Parabacteroides distasonis		2.14	0.27	SAL 88		
Bacteroides thetaiotaomicron		0.48	0.00			
46535712	**Very good identification**					**46535712**
					GRA	
Clostridium clostridioforme		99.58	0.61	GLU 88	50	
Parabacteroides distasonis		0.26	0.23	SAL 88	1	
Bifidobacterium longum subsp. longum		0.09	0.11		99	
46535720	**Good identification**					**46535720**
Bacteroides vulgatus		97.59	0.42	GLU 99		
Parabacteroides distasonis		1.13	0.27	TRE 4		
Clostridium clostridioforme		0.52	0.16			
46535722	**Low discrimination**					**46535722**
Parabacteroides distasonis		59.11	0.62	GLU 99		
Clostridium clostridioforme		26.99	0.50			
Bacteroides thetaiotaomicron		13.12	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46535732	**Low discrimination**					**46535732**
Clostridium clostridioforme		76.68	0.61	GLU 88		
Parabacteroides distasonis		19.85	0.57	SAL 88		
Bacteroides thetaiotaomicron		3.18	0.27			
46537502	**Acceptable identification**					**46537502**
Bifidobacterium adolescentis		93.07	0.41	GLU 99		
Actinomyces israelii		5.77	0.27			
Bifidobacterium longum subsp. longum		1.16	0.07			
46537512	**Acceptable identification**					**46537512**
Bifidobacterium adolescentis		94.59	0.43	GLU 99		
Clostridium clostridioforme		2.41	0.12			
Bifidobacterium longum subsp. longum		1.64	0.11			
46541120	**Good identification**					**46541120**
					CAT	
Prevotella melaninogenica		99.67	0.48	GLU 90	1	
Prevotella oralis		0.21	0.06	FRU 90	1	
Capnocytophaga ochracea		0.12	0.02	CEL 10	90	
46541122	**Low discrimination**					**46541122**
					CAT	
Prevotella melaninogenica		78.77	0.36	GLU 90	1	
Prevotella oralis		16.20	0.27	FRU 90	1	
Capnocytophaga ochracea		4.62	0.18	CEL 10	90	
46551022	**Low discrimination**					**46551022**
					CAT	
Prevotella oralis		89.35	0.41	GLU 99	1	
Prevotella melaninogenica		6.25	0.18	LAC 94	1	
Capnocytophaga ochracea		4.03	0.18		90	
46551102	**Low discrimination**					**46551102**
					CAT	
Prevotella oralis		55.82	0.40	GLU 99	1	
Capnocytophaga ochracea		33.60	0.36	NAG 95	90	
Prevotella melaninogenica		4.74	0.18		1	
46551120	**Very good identification**					**46551120**
					CAT	
Prevotella melaninogenica		96.24	0.64	GLU 90	1	
Prevotella oralis		2.18	0.40	CEL 10	1	
Capnocytophaga ochracea		1.31	0.36		90	
46551122	**Low discrimination**					**46551122**
					CAT	
Prevotella oralis		56.77	0.62	GLU 99	1	
Prevotella melaninogenica		25.10	0.52		1	
Capnocytophaga ochracea		16.19	0.53		90	
46551132	**Low discrimination**					**46551132**
					CAT	
Capnocytophaga ochracea		61.86	0.36	GLU 99	90	
Prevotella oralis		19.72	0.27	MLZ 10	1	
Prevotella melaninogenica		8.72	0.18		1	
46551162	**Low discrimination**					**46551162**
					CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		62.72	0.36	GLU 99	90		
Prevotella oralis		19.99	0.27	SOR 10	1		
Prevotella melaninogenica		8.84	0.18		1		
46551322	**Good identification**						**46551322**
Prevotella oralis		97.29	0.55	GLU 99			
Prevotella melaninogenica		1.01	0.18				
Parabacteroides distasonis		0.89	0.33				
46551502	**Low discrimination**						**46551502**
					GRA	CAT	
Bifidobacterium adolescentis		58.46	0.37	GLU 99	99	1	
Capnocytophaga ochracea		33.61	0.23	XYL 87	1	90	
Prevotella oralis		3.20	0.06		1	1	
46551512	**Low discrimination**						**46551512**
					GRA	CAT	
Bifidobacterium adolescentis		77.65	0.39	GLU 99	99	1	
Clostridium clostridioforme		15.22	0.24	XYL 87	50	33	
Capnocytophaga ochracea		4.06	0.07		1	90	
46551522	**Low discrimination**						**46551522**
					CAT		
Capnocytophaga ochracea		74.48	0.40	GLU 99	90		
Prevotella oralis		14.95	0.27		1		
Prevotella melaninogenica		6.61	0.18		1		
46551712	**Good identification**						**46551712**
Clostridium clostridioforme		100.0	0.38	GLU 88			
46551732	**Acceptable identification**						**46551732**
Clostridium clostridioforme		93.00	0.38	GLU 88			
Parabacteroides distasonis		7.00	0.25	SAL 88			
46553101	**Good identification**						**46553101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
46553103	**Good identification**						**46553103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
46553121	**Low discrimination**						**46553121**
					GRA		
Actinomyces naeslundii		82.44	0.37	ESL 12	99		
Prevotella melaninogenica		17.56	-.04	GLU 99	1		
46553123	**Good identification**						**46553123**
Actinomyces naeslundii		100.0	0.37	ESL 12			
46553502	**Good identification**						**46553502**
Bifidobacterium adolescentis		96.15	0.36	GLU 99			
Actinomyces israelii		3.85	0.18	XYL 87			
46553512	**Good identification**						**46553512**
Bifidobacterium adolescentis		100.0	0.37	GLU 99			
46554612	**Good identification**						**46554612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.38	GLU 88			
46554632	**Good identification**						**46554632**
Clostridium clostridioforme		100.0	0.38	GLU 88			
46554702	**Good identification**						**46554702**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46554712	**Good identification**						**46554712**
Clostridium clostridioforme		100.0	0.47	GLU 88			
46554722	**Good identification**						**46554722**
Clostridium clostridioforme		98.31	0.36	GLU 88			
Parabacteroides distasonis		1.69	0.12	SAL 88			
46554732	**Good identification**						**46554732**
Clostridium clostridioforme		100.0	0.47	GLU 88			
46555112	**Low discrimination**						**46555112**
					GRA	CAT	
Bifidobacterium adolescentis		81.81	0.39	GLU 99	99	1	
Clostridium clostridioforme		10.35	0.20	ARA 87	50	33	
Capnocytophaga ochracea		6.06	0.10		1	90	
46555120	**Low discrimination**						**46555120**
					CAT		
Bacteroides fragilis		77.18	0.38	GLU 99	99		
Prevotella melaninogenica		13.53	0.30	BGL 95	1		
Capnocytophaga ochracea		4.57	0.26		90		
46555122	**Good identification**						**46555122**
					CAT		
Bacteroides fragilis		92.30	0.60	GLU 99	99		
Prevotella oralis		4.49	0.46		1		
Capnocytophaga ochracea		2.59	0.42		90		
46555312	**Good identification**						**46555312**
Clostridium clostridioforme		98.30	0.35	GLU 88			
Parabacteroides distasonis		1.70	0.11	SAL 88			
46555322	**Low discrimination**						**46555322**
					CAT		
Prevotella oralis		45.09	0.40	GLU 99	1		
Parabacteroides distasonis		29.86	0.49	XYL 11	76		
Bacteroides fragilis		21.85	0.25		99		
46555332	**Low discrimination**						**46555332**
Parabacteroides distasonis		61.51	0.45	GLU 99			
Clostridium clostridioforme		35.90	0.35				
Prevotella oralis		1.74	0.06				
46555502	**Low discrimination**						**46555502**
					GRA	CAT	
Bifidobacterium adolescentis		87.70	0.52	GLU 99	99	1	
Clostridium clostridioforme		6.97	0.30		50	33	
Capnocytophaga ochracea		1.88	0.13		1	90	
46555512	**Low discrimination**						**46555512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		76.11	0.53	GLU 99		
Clostridium clostridioforme		22.54	0.41			
Bifidobacterium longum subsp. longum		1.36	0.21			
46555532	**Low discrimination**					**46555532**
					GRA	
Clostridium clostridioforme		65.93	0.41	GLU 88	50	
Bifidobacterium longum subsp. longum		15.85	0.32	SAL 88	99	
Parabacteroides distasonis		10.80	0.33	TRE 88	1	
46555542	**Good identification**					**46555542**
Bifidobacterium adolescentis		96.97	0.36	GLU 99		
Actinomyces israelii		3.03	0.16	SOR 11		
46555552	**Good identification**					**46555552**
Bifidobacterium adolescentis		97.64	0.37	GLU 99		
Clostridium clostridioforme		2.36	0.07	SOR 11		
46555602	**Good identification**					**46555602**
Clostridium clostridioforme		100.0	0.35	GLU 88		
46555612	**Good identification**					**46555612**
Clostridium clostridioforme		100.0	0.47	GLU 88		
46555622	**Good identification**					**46555622**
Clostridium clostridioforme		98.12	0.35	GLU 88		
Parabacteroides distasonis		1.88	0.12	SAL 88		
46555632	**Good identification**					**46555632**
Clostridium clostridioforme		100.0	0.47	GLU 88		
46555702	**Good identification**					**46555702**
					GRA	
Clostridium clostridioforme		97.75	0.44	GLU 88	50	
Bifidobacterium adolescentis		1.70	0.17	SAL 88	99	
Parabacteroides distasonis		0.56	0.12	TRE 88	1	
46555712	**Good identification**					**46555712**
Clostridium clostridioforme		99.54	0.56	GLU 88		
Bifidobacterium adolescentis		0.46	0.19	SAL 88		
46555720	**Good identification**					**46555720**
Bacteroides vulgatus		99.70	0.44	GLU 99		
Clostridium clostridioforme		0.19	0.10	CEL 5		
Parabacteroides distasonis		0.11	0.12			
46555722	**Low discrimination**					**46555722**
Clostridium clostridioforme		49.25	0.44	GLU 88		
Parabacteroides distasonis		27.93	0.46	SAL 88		
Bacteroides uniformis		9.28	0.19	TRE 88		
46555732	**Acceptable identification**					**46555732**
Clostridium clostridioforme		92.08	0.56	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		6.17	0.41	SAL 88			
Bacteroides thetaiotaomicron		1.34	0.14	TRE 88			
46557102	**Low discrimination**						**46557102**
					CAT	GRA	
Bifidobacterium adolescentis		62.70	0.36	GLU 99	1	99	
Actinomyces israelii		35.26	0.38	ARA 87	1	99	
Bacteroides fragilis		2.04	-.09		99	1	
46557112	**Acceptable identification**						**46557112**
Bifidobacterium adolescentis		93.57	0.37	GLU 99			
Actinomyces israelii		6.43	0.24	ARA 87			
46557502	**Acceptable identification**						**46557502**
Bifidobacterium adolescentis		93.57	0.50	GLU 99			
Actinomyces israelii		6.43	0.36				
46557512	**Good identification**						**46557512**
Bifidobacterium adolescentis		98.37	0.52	GLU 99			
Actinomyces israelii		0.83	0.22				
Bifidobacterium dentium		0.44	-.07				
46557552	**Good identification**						**46557552**
Bifidobacterium adolescentis		97.42	0.36	GLU 99			
Actinomyces israelii		2.58	0.15	SOR 11			
46565712	**Good identification**						**46565712**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46565732	**Acceptable identification**						**46565732**
Clostridium clostridioforme		93.64	0.36	GLU 88			
Parabacteroides distasonis		3.90	0.18	SAL 88			
Bacteroides thetaiotaomicron		2.47	-.01	FRU 99			
46570612	**Good identification**						**46570612**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46570632	**Good identification**						**46570632**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46570712	**Good identification**						**46570712**
Clostridium clostridioforme		100.0	0.45	GLU 88			
46570732	**Good identification**						**46570732**
Clostridium clostridioforme		100.0	0.45	GLU 88			
46571101	**Good identification**						**46571101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
46571103	**Good identification**						**46571103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
46571121	**Low discrimination**						**46571121**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		82.44	0.37	ESL 12	99	
Prevotella melaninogenica		17.56	-.04	GLU 99	1	
46571122	**Low discrimination**					**46571122**
Prevotella oralis		74.29	0.41	GLU 99		
Parabacteroides distasonis		6.89	0.36	TRE 6		
Prevotella melaninogenica		5.20	0.18			
46571123	**Low discrimination**					**46571123**
					GRA	
Actinomyces naeslundii		63.29	0.37	ESL 12	99	
Prevotella oralis		36.71	0.07	GLU 99	1	
46571332	**Low discrimination**					**46571332**
Parabacteroides distasonis		53.61	0.40	GLU 99		
Clostridium clostridioforme		44.84	0.33	XYL 90		
Prevotella oralis		1.55	0.01			
46571512	**Low discrimination**					**46571512**
					GRA	
Clostridium clostridioforme		67.80	0.38	GLU 88	50	
Bifidobacterium adolescentis		31.45	0.36	SAL 88	99	
Leptotrichia buccalis		0.75	-.11	XYL 91	1	
46571532	**Low discrimination**					**46571532**
					GRA	
Clostridium clostridioforme		87.57	0.38	GLU 88	50	
Parabacteroides distasonis		10.01	0.28	SAL 88	1	
Bifidobacterium longum subsp. longum		2.42	0.13	XYL 91	99	
46571612	**Good identification**					**46571612**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46571632	**Good identification**					**46571632**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46571702	**Good identification**					**46571702**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46571712	**Good identification**					**46571712**
Clostridium clostridioforme		100.0	0.53	GLU 88		
46571722	**Low discrimination**					**46571722**
Clostridium clostridioforme		65.11	0.42	GLU 88		
Parabacteroides distasonis		25.77	0.41	SAL 88		
Bacteroides thetaiotaomicron		8.46	0.16	XYL 91		
46571732	**Acceptable identification**					**46571732**
Clostridium clostridioforme		94.53	0.53	GLU 88		
Parabacteroides distasonis		4.42	0.36	SAL 88		
Bacteroides thetaiotaomicron		1.05	0.09	XYL 91		
46573100	**Low discrimination**					**46573100**
Actinomyces naeslundii		59.58	0.39	ESL 12		
Actinomyces israelii		40.42	0.27	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46573101	**Good identification**					**46573101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
46573103	**Good identification**					**46573103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
46573121	**Good identification**					**46573121**
Actinomyces naeslundii		100.0	0.42	ESL 12		
46573123	**Good identification**					**46573123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
46574312	**Good identification**					**46574312**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46574332	**Good identification**					**46574332**
Clostridium clostridioforme		96.88	0.42	GLU 88		
Parabacteroides distasonis		3.13	0.22	SAL 88		
46574412	**Good identification**					**46574412**
Clostridium clostridioforme		100.0	0.38	GLU 88		
46574432	**Good identification**					**46574432**
Clostridium clostridioforme		100.0	0.38	GLU 88		
46574502	**Good identification**					**46574502**
Clostridium clostridioforme		96.65	0.36	GLU 88		
Bifidobacterium adolescentis		3.35	0.14	SAL 88		
46574512	**Good identification**					**46574512**
Clostridium clostridioforme		99.08	0.47	GLU 88		
Bifidobacterium adolescentis		0.92	0.16	SAL 88		
46574522	**Good identification**					**46574522**
Clostridium clostridioforme		97.46	0.36	GLU 88		
Parabacteroides distasonis		2.54	0.15	SAL 88		
46574532	**Good identification**					**46574532**
Clostridium clostridioforme		99.69	0.47	GLU 88		
Parabacteroides distasonis		0.31	0.10	SAL 88		
46574602	**Good identification**					**46574602**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46574612	**Good identification**					**46574612**
Clostridium clostridioforme		100.0	0.53	GLU 88		
46574622	**Good identification**					**46574622**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46574632	**Good identification**					**46574632**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.53	GLU 88		
46574702	**Good identification**					**46574702**
Clostridium clostridioforme		100.0	0.51	GLU 88		
46574712	**Very good identification**					**46574712**
Clostridium clostridioforme		100.0	0.62	GLU 88		
46574722	**Good identification**					**46574722**
Clostridium clostridioforme		98.04	0.51	GLU 88		
Parabacteroides distasonis		1.05	0.23	SAL 88		
Bacteroides thetaiotaomicron		0.92	0.06			
46574732	**Very good identification**					**46574732**
Clostridium clostridioforme		99.79	0.62	GLU 88		
Parabacteroides distasonis		0.13	0.18	SAL 88		
Bacteroides thetaiotaomicron		0.08	-.01			
46575112	**Low discrimination**					**46575112**
Clostridium clostridioforme		55.48	0.35	GLU 88		
Bifidobacterium adolescentis		39.88	0.36	SAL 88		
Actinomyces israelii		2.23	0.21	RHA 88		
46575122	**Low discrimination**					**46575122**
					CAT	
Parabacteroides distasonis		58.89	0.53	GLU 99	76	
Bacteroides fragilis		28.38	0.25		99	
Prevotella oralis		8.72	0.26		1	
46575132	**Low discrimination**					**46575132**
Parabacteroides distasonis		71.77	0.48	GLU 99		
Clostridium clostridioforme		27.59	0.35			
Bacteroides fragilis		0.65	-.09			
46575212	**Good identification**					**46575212**
Clostridium clostridioforme		100.0	0.41	GLU 88		
46575232	**Good identification**					**46575232**
Clostridium clostridioforme		96.53	0.41	GLU 88		
Parabacteroides distasonis		3.47	0.22	SAL 88		
46575302	**Low discrimination**					**46575302**
					GRA	
Clostridium clostridioforme		89.80	0.39	GLU 88	50	
Parabacteroides distasonis		8.17	0.27	SAL 88	1	
Actinomyces israelii		1.13	0.16	ARA 94	99	
46575312	**Good identification**					**46575312**
Clostridium clostridioforme		98.94	0.50	GLU 88		
Parabacteroides distasonis		1.06	0.22	SAL 88		
46575322	**Low discrimination**					**46575322**
Parabacteroides distasonis		87.60	0.61	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		9.73	0.39			
Prevotella oralis		1.85	0.19			
46575332	**Low discrimination**					**46575332**
Parabacteroides distasonis		51.38	0.56	GLU 99		
Clostridium clostridioforme		48.28	0.50			
Bacteroides thetaiotaomicron		0.20	-.01			
46575402	**Low discrimination**					**46575402**
					SPO	
Clostridium clostridioforme		87.74	0.35	GLU 88	66	
Actinomyces israelii		4.71	0.23	SAL 88	1	
Clostridium butyricum		4.16	0.01	RHA 88	99	
46575412	**Good identification**					**46575412**
Clostridium clostridioforme		98.97	0.47	GLU 88		
Bifidobacterium adolescentis		1.03	0.16	SAL 88		
46575422	**Good identification**					**46575422**
Clostridium clostridioforme		97.17	0.35	GLU 88		
Parabacteroides distasonis		2.83	0.15	SAL 88		
46575432	**Good identification**					**46575432**
Clostridium clostridioforme		99.66	0.47	GLU 88		
Parabacteroides distasonis		0.34	0.10	SAL 88		
46575502	**Low discrimination**					**46575502**
					SPO	
Bifidobacterium adolescentis		44.66	0.48	GLU 99	1	
Clostridium clostridioforme		39.02	0.44		66	
Clostridium butyricum		13.27	0.25		99	
46575512	**Low discrimination**					**46575512**
					SPO	
Clostridium clostridioforme		75.88	0.56	GLU 88	66	
Bifidobacterium adolescentis		23.30	0.50	SAL 88	1	
Clostridium butyricum		0.30	0.03	RHA 88	99	
46575522	**Low discrimination**					**46575522**
Clostridium clostridioforme		47.23	0.44	GLU 88		
Parabacteroides distasonis		40.67	0.49	SAL 88		
Bacteroides thetaiotaomicron		8.04	0.21	RHA 88		
46575532	**Low discrimination**					**46575532**
Clostridium clostridioforme		88.26	0.56	GLU 88		
Parabacteroides distasonis		8.98	0.44	SAL 88		
Bacteroides thetaiotaomicron		1.28	0.14	RHA 88		
46575602	**Good identification**					**46575602**
Clostridium clostridioforme		100.0	0.50	GLU 88		
46575612	**Very good identification**					**46575612**
Clostridium clostridioforme		100.0	0.61	GLU 88		
46575622	**Good identification**					**46575622**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		96.81	0.50	GLU 88		
Bacteroides thetaiotaomicron		2.04	0.11	SAL 88		
Parabacteroides distasonis		1.15	0.23			
46575632	**Very good identification**					**46575632**
Clostridium clostridioforme		99.68	0.61	GLU 88		
Bacteroides thetaiotaomicron		0.18	0.04	SAL 88		
Parabacteroides distasonis		0.14	0.18			
46575702	**Very good identification**					**46575702**
Clostridium clostridioforme		98.98	0.59	GLU 88		
Parabacteroides distasonis		0.35	0.23	SAL 88		
Bacteroides thetaiotaomicron		0.31	0.06			
46575710	**Good identification**					**46575710**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46575712	**Very good identification**					**46575712**
					GRA	
Clostridium clostridioforme		99.89	0.70	GLU 88	50	
Parabacteroides distasonis		0.04	0.18	SAL 88	1	
Bifidobacterium adolescentis		0.04	0.16		99	
46575713	**Good identification**					**46575713**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46575722	**Low discrimination**					**46575722**
Clostridium clostridioforme		58.74	0.59	GLU 88		
Parabacteroides distasonis		20.69	0.57	SAL 88		
Bacteroides thetaiotaomicron		18.12	0.40			
46575730	**Acceptable identification**					**46575730**
Clostridium clostridioforme		93.64	0.36	GLU 88		
Parabacteroides distasonis		3.90	0.18	SAL 88		
Bacteroides thetaiotaomicron		2.47	-.01	BGL 99		
46575732	**Good identification**					**46575732**
Clostridium clostridioforme		93.10	0.70	GLU 88		
Parabacteroides distasonis		3.88	0.52	SAL 88		
Bacteroides thetaiotaomicron		2.45	0.33			
46575733	**Acceptable identification**					**46575733**
Clostridium clostridioforme		93.64	0.36	GLU 88		
Parabacteroides distasonis		3.90	0.18	SAL 88		
Bacteroides thetaiotaomicron		2.47	-.01	URE 1		
46575752	**Good identification**					**46575752**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46575772	**Acceptable identification**					**46575772**
Clostridium clostridioforme		91.48	0.36	GLU 88		
Parabacteroides distasonis		3.81	0.18	SAL 88		
Bacteroides thetaiotaomicron		2.41	-.01	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46577002	**Good identification**					**46577002**
Actinomyces israelii		100.0	0.37	GLU 99		
46577100	**Good identification**					**46577100**
					SPO	
Actinomyces israelii		96.48	0.45	GLU 99	1	
Clostridium butyricum		3.52	-.01	SAL 92	99	
46577102	**Low discrimination**					**46577102**
Actinomyces israelii		76.04	0.48	GLU 99		
Bifidobacterium adolescentis		23.97	0.33	SAL 92		
46577140	**Good identification**					**46577140**
Actinomyces israelii		100.0	0.38	GLU 99		
46577142	**Acceptable identification**					**46577142**
Actinomyces israelii		90.90	0.41	GLU 99		
Bifidobacterium adolescentis		9.11	0.17	SAL 92		
46577402	**Acceptable identification**					**46577402**
Actinomyces israelii		90.57	0.36	GLU 99		
Bifidobacterium adolescentis		9.43	0.13	SAL 92		
46577500	**Low discrimination**					**46577500**
					SPO	
Actinomyces israelii		69.43	0.43	GLU 99	1	
Clostridium butyricum		27.86	0.15	SAL 92	99	
Bifidobacterium adolescentis		2.71	0.13		1	
46577502	**Low discrimination**					**46577502**
					SPO	
Bifidobacterium adolescentis		69.10	0.47	GLU 99	1	
Actinomyces israelii		26.80	0.46		1	
Clostridium butyricum		2.79	0.08		99	
46577512	**Low discrimination**					**46577512**
Bifidobacterium adolescentis		57.03	0.48	GLU 99		
Bifidobacterium dentium		37.98	0.27			
Actinomyces israelii		2.70	0.32			
46577542	**Low discrimination**					**46577542**
					COC	
Actinomyces israelii		49.59	0.39	GLU 99	1	
Bifidobacterium adolescentis		40.64	0.31	SAL 92	1	
Blautia producta		9.77	0.07		99	
46577712	**Acceptable identification**					**46577712**
Clostridium clostridioforme		94.60	0.36	GLU 88		
Bifidobacterium adolescentis		3.24	0.14	SAL 88		
Bifidobacterium dentium		2.16	-.07	MAN 1		
46577732	**Low discrimination**					**46577732**
Clostridium clostridioforme		76.35	0.36	GLU 88		
Bacteroides ovatus		15.37	0.14	SAL 88		
Parabacteroides distasonis		3.18	0.18	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46610000	**Very good identification**					**46610000**
Fusobacterium mortiferum		100.0	0.57	GLU 99		
46610002	**Good identification**					**46610002**
Fusobacterium mortiferum		100.0	0.40	GLU 99		
46610100	**Very good identification**					**46610100**
Fusobacterium mortiferum		100.0	0.63	GLU 99		
46610102	**Good identification**					**46610102**
					GRA	
Fusobacterium mortiferum		99.19	0.47	GLU 99	1	
Bifidobacterium longum subsp. infantis		0.81	-.07	BGL 10	99	
46611000	**Very good identification**					**46611000**
Fusobacterium mortiferum		100.0	0.60	GLU 99		
46611002	**Good identification**					**46611002**
					GRA	
Fusobacterium mortiferum		98.74	0.43	GLU 99	1	
Bifidobacterium longum subsp. infantis		1.26	-.07	BGL 10	99	
46611100	**Very good identification**					**46611100**
					GRA	
Fusobacterium mortiferum		99.41	0.66	GLU 99	1	
Bifidobacterium longum subsp. infantis		0.59	0.10		99	
46611102	**Low discrimination**					**46611102**
					GRA	
Fusobacterium mortiferum		64.48	0.50	GLU 99	1	
Bifidobacterium longum subsp. infantis		34.95	0.27	BGL 10	99	
Prevotella oralis		0.58	0.00		1	
46614000	**Good identification**					**46614000**
Fusobacterium mortiferum		100.0	0.37	GLU 99		
46614100	**Good identification**					**46614100**
Fusobacterium mortiferum		100.0	0.44	GLU 99		
46615000	**Good identification**					**46615000**
Fusobacterium mortiferum		100.0	0.40	GLU 99		
46615100	**Acceptable identification**					**46615100**
					GRA	
Fusobacterium mortiferum		93.92	0.47	GLU 99	1	
Bifidobacterium longum subsp. infantis		6.08	0.09	XYL 7	99	
46615322	**Good identification**					**46615322**
					GRA	
Parabacteroides distasonis		96.75	0.45	GLU 99	1	
Prevotella oralis		1.30	0.00		1	
Bifidobacterium longum subsp. infantis		1.15	-.04		99	
46615332	**Good identification**					**46615332**
Parabacteroides distasonis		97.78	0.41	GLU 99		
Clostridium clostridioforme		2.23	0.07			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46615722	**Acceptable identification**					**46615722**
Parabacteroides distasonis		93.57	0.42	GLU 99		
Clostridium clostridioforme		6.43	0.16			
46615732	**Low discrimination**					**46615732**
Parabacteroides distasonis		63.22	0.37	GLU 99		
Clostridium clostridioforme		36.78	0.27			
46630000	**Good identification**					**46630000**
Fusobacterium mortiferum		100.0	0.48	GLU 99		
46630100	**Good identification**					**46630100**
Fusobacterium mortiferum		100.0	0.54	GLU 99		
46630102	**Good identification**					**46630102**
Fusobacterium mortiferum		100.0	0.38	GLU 99		
46631000	**Good identification**					**46631000**
Fusobacterium mortiferum		100.0	0.51	GLU 99		
46631100	**Very good identification**					**46631100**
Fusobacterium mortiferum		100.0	0.57	GLU 99		
46631102	**Good identification**					**46631102**
					GRA	
Fusobacterium mortiferum		98.20	0.41	GLU 99	1	
Bifidobacterium longum subsp. infantis		1.80	-.07	BGL 10	99	
46631322	**Good identification**					**46631322**
Parabacteroides distasonis		100.0	0.40	GLU 99		
46631332	**Good identification**					**46631332**
Parabacteroides distasonis		96.84	0.36	GLU 99		
Clostridium clostridioforme		3.16	0.04	XYL 90		
46631722	**Acceptable identification**					**46631722**
Parabacteroides distasonis		91.03	0.37	GLU 99		
Clostridium clostridioforme		8.97	0.14	XYL 90		
46635100	**Good identification**					**46635100**
Fusobacterium mortiferum		100.0	0.38	GLU 99		
46635122	**Good identification**					**46635122**
					GRA	
Parabacteroides distasonis		99.23	0.48	GLU 99	1	
Bifidobacterium longum subsp. infantis		0.77	-.04		99	
46635132	**Good identification**					**46635132**
Parabacteroides distasonis		98.52	0.44	GLU 99		
Clostridium clostridioforme		1.48	0.07			
46635322	**Very good identification**					**46635322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		99.57	0.56	GLU 99			
Clostridium clostridioforme		0.43	0.10				
46635332	**Good identification**						**46635332**
Parabacteroides distasonis		96.47	0.52	GLU 99			
Clostridium clostridioforme		3.53	0.22				
46635522	**Good identification**						**46635522**
Parabacteroides distasonis		95.67	0.45	GLU 99			
Clostridium clostridioforme		4.33	0.16				
46635532	**Low discrimination**						**46635532**
Parabacteroides distasonis		72.30	0.40	GLU 99			
Clostridium clostridioforme		27.70	0.27				
46635712	**Good identification**						**46635712**
Clostridium clostridioforme		98.93	0.42	GLU 88			
Parabacteroides distasonis		1.07	0.14	MLT 99			
46635722	**Low discrimination**						**46635722**
Parabacteroides distasonis		89.38	0.53	GLU 99			
Clostridium clostridioforme		9.89	0.31				
Bacteroides thetaiotaomicron		0.73	0.01				
46635732	**Low discrimination**						**46635732**
Parabacteroides distasonis		51.48	0.48	GLU 99			
Clostridium clostridioforme		48.21	0.42				
Bacteroides thetaiotaomicron		0.30	-.06				
46650000	**Good identification**						**46650000**
Fusobacterium mortiferum		100.0	0.48	GLU 99			
46650100	**Good identification**						**46650100**
Fusobacterium mortiferum		100.0	0.55	GLU 99			
46650102	**Good identification**						**46650102**
Fusobacterium mortiferum		100.0	0.38	GLU 99			
46651000	**Good identification**						**46651000**
					GRA		
Fusobacterium mortiferum		97.60	0.51	GLU 99	1		
Lactobacillus catenaformis		1.94	0.12		99		
Lactobacillus jensenii		0.46	0.02		99		
46651002	**Low discrimination**						**46651002**
					GRA	SPO	
Fusobacterium mortiferum		74.72	0.35	GLU 99	1	1	
Lactobacillus catenaformis		13.39	0.12	BGL 10	99	1	
Clostridium barati		8.30	0.04		99	99	
46651100	**Very good identification**						**46651100**
					GRA		
Fusobacterium mortiferum		97.18	0.58	GLU 99	1		
Lactobacillus catenaformis		1.24	0.15		99		
Collinsella aerofaciens		0.82	-.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46651102	**Low discrimination**						**46651102**
					GRA		
Fusobacterium mortiferum		54.48	0.41	GLU 99	1		
Lactobacillus acidophilus		12.91	0.18	BGL 10	99		
Bifidobacterium breve		12.45	0.27		99		
46654100	**Good identification**						**46654100**
Fusobacterium mortiferum		100.0	0.35	GLU 99			
46655100	**Good identification**						**46655100**
					GRA	SPO	
Fusobacterium mortiferum		96.60	0.38	GLU 99	1	1	
Clostridium butyricum		3.40	-.05	XYL 7	99	99	
46655322	**Low discrimination**						**46655322**
Parabacteroides distasonis		85.95	0.41	GLU 99			
Prevotella oralis		7.63	0.10				
Bacteroides uniformis		4.98	0.01				
46655332	**Low discrimination**						**46655332**
Parabacteroides distasonis		87.60	0.36	GLU 99			
Clostridium clostridioforme		12.40	0.16				
46655712	**Good identification**						**46655712**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46655732	**Low discrimination**						**46655732**
Clostridium clostridioforme		78.34	0.36	GLU 88			
Parabacteroides distasonis		21.66	0.32	MLT 99			
46670000	**Good identification**						**46670000**
Fusobacterium mortiferum		100.0	0.39	GLU 99			
46670100	**Good identification**						**46670100**
Fusobacterium mortiferum		100.0	0.46	GLU 99			
46671000	**Good identification**						**46671000**
					GRA		
Fusobacterium mortiferum		95.46	0.42	GLU 99	1		
Lactobacillus jensenii		4.54	0.10		99		
46671100	**Good identification**						**46671100**
					GRA		
Fusobacterium mortiferum		99.37	0.49	GLU 99	1		
Actinomyces naeslundii		0.63	0.23		99		
46671322	**Low discrimination**						**46671322**
					SPO	GRA	
Parabacteroides distasonis		57.08	0.36	GLU 99	1	1	
Clostridium ramosum		31.56	0.19	XYL 90	92	50	
Lactobacillus acidophilus		6.20	0.08		1	99	
46674712	**Good identification**						**46674712**
Clostridium clostridioforme		100.0	0.43	GLU 88			
46674732	**Good identification**						**46674732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		99.48	0.43	GLU 88			
Parabacteroides distasonis		0.52	0.09	MLT 99			
46675122	**Good identification**						**46675122**
Parabacteroides distasonis		98.91	0.44	GLU 99			
Clostridium clostridioforme		1.09	0.05				
46675132	**Acceptable identification**						**46675132**
Parabacteroides distasonis		91.48	0.39	GLU 99			
Clostridium clostridioforme		8.52	0.16				
46675322	**Good identification**						**46675322**
Parabacteroides distasonis		97.38	0.52	GLU 99			
Clostridium clostridioforme		2.62	0.19				
46675332	**Low discrimination**						**46675332**
Parabacteroides distasonis		81.45	0.47	GLU 99			
Clostridium clostridioforme		18.55	0.31				
46675512	**Low discrimination**						**46675512**
					SPO		
Clostridium clostridioforme		76.16	0.36	GLU 88	66		
Bifidobacterium adolescentis		18.07	0.29	MLT 99	1		
Clostridium butyricum		4.04	0.03	RHA 88	99		
46675522	**Low discrimination**						**46675522**
					GRA	SPO	
Parabacteroides distasonis		71.74	0.40	GLU 99	1	1	
Clostridium clostridioforme		20.20	0.25		50	66	
Clostridium butyricum		8.06	0.07		99	99	
46675532	**Low discrimination**						**46675532**
					GRA		
Clostridium clostridioforme		68.48	0.36	GLU 88	50		
Parabacteroides distasonis		28.75	0.35	MLT 99	1		
Bifidobacterium dentium		2.78	-.03	RHA 88	99		
46675612	**Good identification**						**46675612**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46675632	**Good identification**						**46675632**
Clostridium clostridioforme		99.42	0.42	GLU 88			
Parabacteroides distasonis		0.58	0.09	MLT 99			
46675702	**Good identification**						**46675702**
					GRA	SPO	
Clostridium clostridioforme		96.71	0.40	GLU 88	50	66	
Clostridium butyricum		1.87	-.01	MLT 99	99	99	
Parabacteroides distasonis		1.42	0.14		1	1	
46675712	**Good identification**						**46675712**
Clostridium clostridioforme		99.83	0.51	GLU 88			
Parabacteroides distasonis		0.17	0.09	MLT 99			
46675722	**Low discrimination**						**46675722**
Parabacteroides distasonis		57.17	0.48	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		39.34	0.40			
Bacteroides thetaiotaomicron		1.85	0.07			
46675732	**Low discrimination**					**46675732**
Clostridium clostridioforme		84.61	0.51	GLU 88		
Parabacteroides distasonis		14.53	0.44	MLT 99		
Bacteroides ovatus		0.52	0.03			
46710000	**Good identification**					**46710000**
Fusobacterium mortiferum		100.0	0.41	GLU 99		
46710100	**Acceptable identification**					**46710100**
					GRA	
Fusobacterium mortiferum		93.26	0.47	GLU 99	1	
Bifidobacterium longum subsp. infantis		6.74	0.10	MLT 11	99	
46710712	**Good identification**					**46710712**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46710732	**Good identification**					**46710732**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46711000	**Low discrimination**					**46711000**
					GRA	
Fusobacterium mortiferum		86.91	0.44	GLU 99	1	
Bifidobacterium longum subsp. infantis		9.77	0.10	MLT 11	99	
Actinomyces naeslundii		2.21	0.28		99	
46711022	**Low discrimination**					**46711022**
					GRA	
Prevotella oralis		51.81	0.36	GLU 99	1	
Bifidobacterium longum subsp. infantis		46.21	0.32	LAC 94	99	
Clostridium perfringens		0.75	0.08		99	
46711100	**Low discrimination**					**46711100**
					GRA	
Bifidobacterium longum subsp. infantis		80.25	0.44	GLU 99	99	
Fusobacterium mortiferum		16.82	0.50	BGL 91	1	
Prevotella melaninogenica		1.09	0.12		1	
46711102	**Very good identification**					**46711102**
					GRA	
Bifidobacterium longum subsp. infantis		97.59	0.62	GLU 99	99	
Prevotella oralis		1.62	0.35		1	
Bifidobacterium adolescentis		0.24	0.31		99	
46711120	**Low discrimination**					**46711120**
					GRA	
Bifidobacterium longum subsp. infantis		52.69	0.49	GLU 99	99	
Prevotella melaninogenica		39.78	0.47	BGL 91	1	
Prevotella oralis		4.98	0.35		1	
46711122	**Low discrimination**					**46711122**
					GRA	
Bifidobacterium longum subsp. infantis		83.30	0.66	GLU 99	99	
Prevotella oralis		14.78	0.56		1	
Prevotella melaninogenica		1.19	0.34		1	
46711322	**Low discrimination**					**46711322**
					GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		85.08	0.50	GLU 99	1	
Bifidobacterium longum subsp. infantis		11.30	0.32		99	
Parabacteroides distasonis		3.11	0.38		1	
46711502	**Low discrimination**					**46711502**
Bifidobacterium adolescentis		59.11	0.45	GLU 99		
Bifidobacterium longum subsp. infantis		36.59	0.27	XYL 87		
Clostridium clostridioforme		2.55	0.18			
46711512	**Low discrimination**					**46711512**
Bifidobacterium adolescentis		84.68	0.46	GLU 99		
Clostridium clostridioforme		13.63	0.29	XYL 87		
Bifidobacterium longum subsp. longum		1.26	0.13			
46711612	**Good identification**					**46711612**
Clostridium clostridioforme		100.0	0.35	GLU 88		
46711632	**Good identification**					**46711632**
Clostridium clostridioforme		100.0	0.35	GLU 88		
46711712	**Good identification**					**46711712**
Clostridium clostridioforme		99.15	0.44	GLU 88		
Bifidobacterium adolescentis		0.85	0.12	TRE 88		
46711732	**Acceptable identification**					**46711732**
Clostridium clostridioforme		93.53	0.44	GLU 88		
Parabacteroides distasonis		6.47	0.30	TRE 88		
46713001	**Good identification**					**46713001**
Actinomyces naeslundii		100.0	0.36	ESL 12		
46713003	**Good identification**					**46713003**
Actinomyces naeslundii		100.0	0.36	ESL 12		
46713021	**Good identification**					**46713021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
46713023	**Good identification**					**46713023**
Actinomyces naeslundii		100.0	0.36	ESL 12		
46713100	**Low discrimination**					**46713100**
					GRA	
Actinomyces naeslundii		45.56	0.41	ESL 12	99	
Bifidobacterium longum subsp. infantis		36.50	0.10	GLU 99	99	
Fusobacterium mortiferum		7.65	0.16		1	
46713101	**Good identification**					**46713101**
Actinomyces naeslundii		100.0	0.44	ESL 12		
46713103	**Acceptable identification**					**46713103**
Actinomyces naeslundii		94.83	0.44	ESL 12		
Bifidobacterium longum subsp. infantis		5.17	-.07	GLU 99		
46713121	**Good identification**					**46713121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	ESL 12		
46713123	**Acceptable identification**					**46713123**
Actinomyces naeslundii		91.16	0.44	ESL 12		
Bifidobacterium longum subsp. infantis		8.84	-.03	GLU 99		
46713502	**Good identification**					**46713502**
Bifidobacterium adolescentis		97.98	0.43	GLU 99		
Actinomyces israelii		1.27	0.17	XYL 87		
Bifidobacterium longum subsp. infantis		0.75	-.07			
46713512	**Good identification**					**46713512**
Bifidobacterium adolescentis		100.0	0.45	GLU 99		
46714512	**Low discrimination**					**46714512**
Clostridium clostridioforme		88.92	0.38	GLU 88		
Bifidobacterium adolescentis		11.08	0.26	TRE 88		
46714532	**Good identification**					**46714532**
Clostridium clostridioforme		100.0	0.38	GLU 88		
46714612	**Good identification**					**46714612**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46714632	**Good identification**					**46714632**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46714702	**Good identification**					**46714702**
Clostridium clostridioforme		97.33	0.42	GLU 88		
Eubacterium contortum		2.68	0.10	TRE 88		
46714712	**Good identification**					**46714712**
Clostridium clostridioforme		100.0	0.53	GLU 88		
46714722	**Good identification**					**46714722**
Clostridium clostridioforme		98.45	0.42	GLU 88		
Parabacteroides distasonis		1.55	0.17	TRE 88		
46714732	**Good identification**					**46714732**
Clostridium clostridioforme		99.81	0.53	GLU 88		
Parabacteroides distasonis		0.19	0.12	TRE 88		
46715100	**Acceptable identification**					**46715100**
					GRA	
Bifidobacterium longum subsp. infantis		93.55	0.43	GLU 99	99	
Eubacterium contortum		3.83	0.24	BGL 91	99	
Fusobacterium mortiferum		1.80	0.31		1	
46715102	**Very good identification**					**46715102**
					GRA	
Bifidobacterium longum subsp. infantis		97.53	0.60	GLU 99	99	
Bifidobacterium adolescentis		1.95	0.45		99	
Prevotella oralis		0.25	0.19		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46715112	**Low discrimination**						**46715112**
Bifidobacterium adolescentis		65.87	0.46	GLU 99			
Bifidobacterium longum subsp. infantis		27.29	0.26	ARA 87			
Clostridium clostridioforme		6.84	0.26				
46715120	**Acceptable identification**						**46715120**
					GRA		
Bifidobacterium longum subsp. infantis		92.70	0.47	GLU 99	99		
Bacteroides vulgatus		3.23	0.20	BGL 91	1		
Prevotella oralis		1.32	0.19		1		
46715122	**Good identification**						**46715122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		93.91	0.64	GLU 99	1	99	
Prevotella oralis		2.52	0.41		1	1	
Bacteroides fragilis		2.43	0.32		99	1	
46715300	**Very good identification**						**46715300**
					GRA		
Eubacterium contortum		99.61	0.58	GLU 99	99		
Bifidobacterium longum subsp. infantis		0.25	0.09		99		
Bacteroides vulgatus		0.14	0.02		1		
46715302	**Low discrimination**						**46715302**
Eubacterium contortum		81.57	0.45	GLU 99			
Bifidobacterium longum subsp. infantis		11.64	0.26				
Clostridium clostridioforme		4.70	0.30				
46715312	**Acceptable identification**						**46715312**
					GRA		
Clostridium clostridioforme		93.66	0.41	GLU 88	50		
Eubacterium contortum		3.61	0.11	TRE 88	99		
Parabacteroides distasonis		1.49	0.16	ARA 94	1		
46715320	**Low discrimination**						**46715320**
					GRA		
Eubacterium contortum		54.75	0.45	GLU 99	99		
Bacteroides vulgatus		42.62	0.36		1		
Bifidobacterium longum subsp. infantis		1.37	0.13		99		
46715322	**Low discrimination**						**46715322**
					GRA		
Parabacteroides distasonis		49.23	0.55	GLU 99	1		
Prevotella oralis		18.52	0.35		1		
Bifidobacterium longum subsp. infantis		16.29	0.30		99		
46715332	**Low discrimination**						**46715332**
Parabacteroides distasonis		60.49	0.50	GLU 99			
Clostridium clostridioforme		38.44	0.41				
Prevotella oralis		0.43	0.00				
46715412	**Low discrimination**						**46715412**
Clostridium clostridioforme		87.79	0.38	GLU 88			
Bifidobacterium adolescentis		12.21	0.26	TRE 88			
46715432	**Good identification**						**46715432**
Clostridium clostridioforme		100.0	0.38	GLU 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46715502	**Low discrimination**					**46715502**
Bifidobacterium adolescentis		86.35	0.59	GLU 99		
Bifidobacterium longum subsp. infantis		6.54	0.26			
Clostridium clostridioforme		5.63	0.35			
46715512	**Low discrimination**					**46715512**
Bifidobacterium adolescentis		80.01	0.60	GLU 99		
Clostridium clostridioforme		19.46	0.47			
Bifidobacterium longum subsp. longum		0.53	0.21			
46715520	**Acceptable identification**					**46715520**
					GRA	
Bacteroides vulgatus		94.15	0.39	GLU 99	1	
Bifidobacterium longum subsp. longum		2.96	0.28	RHA 90	99	
Bifidobacterium longum subsp. infantis		2.06	0.13		99	
46715532	**Low discrimination**					**46715532**
					GRA	
Clostridium clostridioforme		76.55	0.47	GLU 88	50	
Parabacteroides distasonis		11.52	0.38	TRE 88	1	
Bifidobacterium longum subsp. longum		8.40	0.32	RHA 88	99	
46715542	**Good identification**					**46715542**
Bifidobacterium adolescentis		99.00	0.43	GLU 99		
Actinomyces israelii		1.00	0.15	SOR 11		
46715552	**Good identification**					**46715552**
Bifidobacterium adolescentis		98.05	0.45	GLU 99		
Clostridium clostridioforme		1.95	0.12	SOR 11		
46715602	**Good identification**					**46715602**
Clostridium clostridioforme		97.02	0.41	GLU 88		
Eubacterium contortum		2.98	0.10	TRE 88		
46715612	**Good identification**					**46715612**
Clostridium clostridioforme		100.0	0.52	GLU 88		
46715622	**Good identification**					**46715622**
Clostridium clostridioforme		98.27	0.41	GLU 88		
Parabacteroides distasonis		1.73	0.17	TRE 88		
46715632	**Good identification**					**46715632**
Clostridium clostridioforme		99.79	0.52	GLU 88		
Parabacteroides distasonis		0.21	0.12	TRE 88		
46715700	**Very good identification**					**46715700**
					GRA	
Eubacterium contortum		97.61	0.57	GLU 99	99	
Bacteroides vulgatus		2.20	0.21		1	
Clostridium clostridioforme		0.19	0.16		50	
46715702	**Low discrimination**					**46715702**
Clostridium clostridioforme		51.72	0.50	GLU 88		
Eubacterium contortum		46.92	0.44	TRE 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		1.09	0.25			
46715712	**Very good identification**					**46715712**
Clostridium clostridioforme		99.18	0.61	GLU 88		
Bifidobacterium adolescentis		0.56	0.26	TRE 88		
Eubacterium contortum		0.20	0.10			
46715720	**Acceptable identification**					**46715720**
					GRA	
Bacteroides vulgatus		92.55	0.56	GLU 99	1	
Eubacterium contortum		7.32	0.44		99	
Clostridium clostridioforme		0.08	0.16		50	
46715722	**Low discrimination**					**46715722**
					GRA	
Clostridium clostridioforme		54.97	0.50	GLU 88	50	
Parabacteroides distasonis		28.63	0.51	TRE 88	1	
Eubacterium contortum		8.80	0.31		99	
46715732	**Good identification**					**46715732**
Clostridium clostridioforme		94.15	0.61	GLU 88		
Parabacteroides distasonis		5.80	0.46	TRE 88		
Bacteroides thetaiotaomicron		0.05	-.05			
46717102	**Low discrimination**					**46717102**
Bifidobacterium adolescentis		55.51	0.43	GLU 99		
Bifidobacterium longum subsp. infantis		34.36	0.26	ARA 87		
Actinomyces israelii		10.13	0.37			
46717112	**Good identification**					**46717112**
Bifidobacterium adolescentis		97.82	0.45	GLU 99		
Actinomyces israelii		2.18	0.23	ARA 87		
46717502	**Very good identification**					**46717502**
Bifidobacterium adolescentis		97.82	0.57	GLU 99		
Actinomyces israelii		2.18	0.35			
46717512	**Very good identification**					**46717512**
Bifidobacterium adolescentis		99.26	0.59	GLU 99		
Clostridium clostridioforme		0.30	0.12			
Actinomyces israelii		0.27	0.21			
46717542	**Acceptable identification**					**46717542**
Bifidobacterium adolescentis		93.44	0.42	GLU 99		
Actinomyces israelii		6.56	0.28	SOR 11		
46717552	**Good identification**					**46717552**
Bifidobacterium adolescentis		99.15	0.43	GLU 99		
Actinomyces israelii		0.85	0.14	SOR 11		
46725712	**Good identification**					**46725712**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46725732	**Good identification**					**46725732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		96.32	0.42	GLU 88			
Parabacteroides distasonis		3.68	0.23	FRU 99			
46730100	**Good identification**						**46730100**
Fusobacterium mortiferum		100.0	0.38	GLU 99			
46730512	**Good identification**						**46730512**
Clostridium clostridioforme		98.32	0.36	GLU 88			
Bifidobacterium adolescentis		1.68	0.09	XYL 91			
46730532	**Good identification**						**46730532**
Clostridium clostridioforme		100.0	0.36	GLU 88			
46730612	**Good identification**						**46730612**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46730632	**Good identification**						**46730632**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46730702	**Good identification**						**46730702**
Clostridium clostridioforme		100.0	0.40	GLU 88			
46730712	**Good identification**						**46730712**
Clostridium clostridioforme		100.0	0.51	GLU 88			
46730722	**Good identification**						**46730722**
Clostridium clostridioforme		98.91	0.40	GLU 88			
Parabacteroides distasonis		1.09	0.12	XYL 91			
46730732	**Good identification**						**46730732**
Clostridium clostridioforme		100.0	0.51	GLU 88			
46731000	**Low discrimination**						**46731000**
					GRA		
Fusobacterium mortiferum		77.32	0.35	GLU 99	1		
Actinomyces naeslundii		12.77	0.33	MLT 11	99		
Lactobacillus jensenii		9.91	0.10		99		
46731001	**Good identification**						**46731001**
Actinomyces naeslundii		100.0	0.36	ESL 12			
46731003	**Good identification**						**46731003**
Actinomyces naeslundii		100.0	0.36	ESL 12			
46731021	**Good identification**						**46731021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
46731023	**Good identification**						**46731023**
Actinomyces naeslundii		100.0	0.36	ESL 12			
46731100	**Low discrimination**						**46731100**
					COC	GRA	
Fusobacterium mortiferum		58.88	0.41	GLU 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		17.28	0.21	MLT 11	99	99	
Actinomyces naeslundii		11.86	0.41		1	99	
46731101	**Good identification**						**46731101**
Actinomyces naeslundii		100.0	0.44	ESL 12			
46731103	**Acceptable identification**						**46731103**
Actinomyces naeslundii		94.83	0.44	ESL 12			
Bifidobacterium longum subsp. infantis		5.17	-.07	GLU 99			
46731121	**Good identification**						**46731121**
Actinomyces naeslundii		100.0	0.44	ESL 12			
46731122	**Low discrimination**						**46731122**
					GRA		
Prevotella oralis		36.28	0.36	GLU 99	1		
Bifidobacterium longum subsp. infantis		32.36	0.32	TRE 6	99		
Parabacteroides distasonis		13.50	0.41		1		
46731123	**Low discrimination**						**46731123**
					GRA		
Actinomyces naeslundii		82.94	0.44	ESL 12	99		
Prevotella oralis		9.02	0.02	GLU 99	1		
Bifidobacterium longum subsp. infantis		8.04	-.03		99		
46731132	**Low discrimination**						**46731132**
					GRA		
Parabacteroides distasonis		45.23	0.37	GLU 99	1		
Bifidobacterium breve		18.86	0.29	XYL 90	99		
Clostridium clostridioforme		16.85	0.24		50		
46731312	**Good identification**						**46731312**
Clostridium clostridioforme		97.85	0.39	GLU 88			
Parabacteroides distasonis		1.09	0.11	XYL 91			
Leptotrichia buccalis		1.06	-.11	ARA 94			
46731322	**Low discrimination**						**46731322**
Parabacteroides distasonis		63.15	0.50	GLU 99			
Prevotella oralis		24.24	0.30	XYL 90			
Clostridium clostridioforme		6.80	0.27				
46731332	**Low discrimination**						**46731332**
Parabacteroides distasonis		52.34	0.45	GLU 99			
Clostridium clostridioforme		47.66	0.39	XYL 90			
46731412	**Good identification**						**46731412**
Clostridium clostridioforme		98.13	0.35	GLU 88			
Bifidobacterium adolescentis		1.87	0.09	XYL 91			
46731432	**Good identification**						**46731432**
Clostridium clostridioforme		100.0	0.35	GLU 88			
46731502	**Low discrimination**						**46731502**
					SPO		
Bifidobacterium adolescentis		65.87	0.42	GLU 99	1		
Clostridium clostridioforme		31.29	0.33	XYL 87	66		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		1.52	-.01		99	
46731512	**Low discrimination**					**46731512**
					GRA	
Clostridium clostridioforme		63.70	0.44	GLU 88	50	
Bifidobacterium adolescentis		35.98	0.43	XYL 91	99	
Leptotrichia buccalis		0.32	-.11	RHA 88	1	
46731532	**Low discrimination**					**46731532**
					GRA	
Clostridium clostridioforme		89.02	0.44	GLU 88	50	
Parabacteroides distasonis		9.35	0.33	XYL 91	1	
Bifidobacterium longum subsp. longum		1.12	0.13	RHA 88	99	
46731602	**Good identification**					**46731602**
Clostridium clostridioforme		100.0	0.39	GLU 88		
46731612	**Good identification**					**46731612**
Clostridium clostridioforme		100.0	0.50	GLU 88		
46731622	**Good identification**					**46731622**
Clostridium clostridioforme		98.79	0.39	GLU 88		
Parabacteroides distasonis		1.21	0.12	XYL 91		
46731632	**Good identification**					**46731632**
Clostridium clostridioforme		100.0	0.50	GLU 88		
46731702	**Good identification**					**46731702**
					GRA	
Clostridium clostridioforme		99.35	0.48	GLU 88	50	
Parabacteroides distasonis		0.37	0.12	XYL 91	1	
Bifidobacterium adolescentis		0.29	0.07		99	
46731712	**Very good identification**					**46731712**
Clostridium clostridioforme		99.92	0.59	GLU 88		
Bifidobacterium adolescentis		0.08	0.09	XYL 91		
46731722	**Low discrimination**					**46731722**
Clostridium clostridioforme		73.08	0.48	GLU 88		
Parabacteroides distasonis		26.56	0.46	XYL 91		
Bacteroides thetaiotaomicron		0.36	-.02			
46731732	**Very good identification**					**46731732**
Clostridium clostridioforme		95.88	0.59	GLU 88		
Parabacteroides distasonis		4.12	0.41	XYL 91		
46733000	**Low discrimination**					**46733000**
Actinomyces naeslundii		86.49	0.38	ESL 12		
Actinomyces israelii		13.51	0.15	GLU 99		
46733001	**Good identification**					**46733001**
Actinomyces naeslundii		100.0	0.41	ESL 12		
46733002	**Low discrimination**					**46733002**
Actinomyces naeslundii		81.02	0.38	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		18.98	0.18	GLU 99		
46733003	**Good identification**					**46733003**
Actinomyces naeslundii		100.0	0.41	ESL 12		
46733020	**Good identification**					**46733020**
Actinomyces naeslundii		100.0	0.38	ESL 12		
46733021	**Good identification**					**46733021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
46733022	**Good identification**					**46733022**
Actinomyces naeslundii		100.0	0.38	ESL 12		
46733023	**Good identification**					**46733023**
Actinomyces naeslundii		100.0	0.41	ESL 12		
46733100	**Low discrimination**					**46733100**
Actinomyces naeslundii		82.00	0.46	ESL 12		
Actinomyces israelii		18.00	0.25	GLU 99		
46733101	**Good identification**					**46733101**
Actinomyces naeslundii		100.0	0.49	ESL 12		
46733102	**Low discrimination**					**46733102**
Actinomyces naeslundii		43.25	0.46	ESL 12		
Bifidobacterium adolescentis		23.77	0.26	GLU 99		
Bifidobacterium breve		16.91	0.24			
46733103	**Good identification**					**46733103**
Actinomyces naeslundii		100.0	0.49	ESL 12		
46733120	**Low discrimination**					**46733120**
					SPO	
Actinomyces naeslundii		52.58	0.46	ESL 12	1	
Clostridium ramosum		43.81	0.19	GLU 99	92	
Bifidobacterium breve		3.62	0.12		1	
46733121	**Good identification**					**46733121**
Actinomyces naeslundii		100.0	0.49	ESL 12		
46733123	**Good identification**					**46733123**
Actinomyces naeslundii		100.0	0.49	ESL 12		
46733141	**Good identification**					**46733141**
Actinomyces naeslundii		100.0	0.36	ESL 12		
46733143	**Good identification**					**46733143**
Actinomyces naeslundii		100.0	0.36	ESL 12		
46733161	**Good identification**					**46733161**
Actinomyces naeslundii		100.0	0.36	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46733163	**Good identification**					**46733163**
Actinomyces naeslundii		100.0	0.36	ESL 12		
46733502	**Acceptable identification**					**46733502**
Bifidobacterium adolescentis		93.17	0.40	GLU 99		
Actinomyces israelii		6.83	0.27	XYL 87		
46733512	**Good identification**					**46733512**
Bifidobacterium adolescentis		96.27	0.42	GLU 99		
Clostridium clostridioforme		2.11	0.10	XYL 87		
Bifidobacterium dentium		1.63	-.07			
46734212	**Good identification**					**46734212**
Clostridium clostridioforme		100.0	0.39	GLU 88		
46734232	**Good identification**					**46734232**
Clostridium clostridioforme		100.0	0.39	GLU 88		
46734302	**Good identification**					**46734302**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46734312	**Good identification**					**46734312**
Clostridium clostridioforme		100.0	0.48	GLU 88		
46734322	**Low discrimination**					**46734322**
Clostridium clostridioforme		79.96	0.36	GLU 88		
Parabacteroides distasonis		20.04	0.32	ARA 94		
46734332	**Good identification**					**46734332**
Clostridium clostridioforme		97.12	0.48	GLU 88		
Parabacteroides distasonis		2.88	0.27	ARA 94		
46734412	**Good identification**					**46734412**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46734432	**Good identification**					**46734432**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46734502	**Good identification**					**46734502**
Clostridium clostridioforme		95.95	0.42	GLU 88		
Bifidobacterium adolescentis		4.05	0.22	RHA 88		
46734512	**Good identification**					**46734512**
Clostridium clostridioforme		98.88	0.53	GLU 88		
Bifidobacterium adolescentis		1.12	0.23	RHA 88		
46734522	**Good identification**					**46734522**
Clostridium clostridioforme		97.66	0.42	GLU 88		
Parabacteroides distasonis		2.34	0.20	RHA 88		
46734532	**Good identification**					**46734532**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		99.72	0.53	GLU 88		
Parabacteroides distasonis		0.28	0.15	RHA 88		
46734602	**Good identification**					**46734602**
Clostridium clostridioforme		100.0	0.48	GLU 88		
46734612	**Very good identification**					**46734612**
Clostridium clostridioforme		100.0	0.59	GLU 88		
46734622	**Good identification**					**46734622**
Clostridium clostridioforme		100.0	0.48	GLU 88		
46734632	**Very good identification**					**46734632**
Clostridium clostridioforme		100.0	0.59	GLU 88		
46734702	**Very good identification**					**46734702**
Clostridium clostridioforme		100.0	0.57	GLU 88		
46734712	**Very good identification**					**46734712**
Clostridium clostridioforme		100.0	0.68	GLU 88		
46734722	**Very good identification**					**46734722**
Clostridium clostridioforme		99.03	0.57	GLU 88		
Parabacteroides distasonis		0.97	0.28			
46734732	**Very good identification**					**46734732**
Clostridium clostridioforme		99.88	0.68	GLU 88		
Parabacteroides distasonis		0.12	0.23			
46735102	**Low discrimination**					**46735102**
Bifidobacterium adolescentis		41.20	0.42	GLU 99		
Bifidobacterium longum subsp. infantis		31.30	0.26	ARA 87		
Clostridium clostridioforme		12.63	0.30			
46735112	**Low discrimination**					**46735112**
					GRA	
Clostridium clostridioforme		52.00	0.41	GLU 88	50	
Bifidobacterium adolescentis		45.51	0.43	RHA 88	99	
Parabacteroides distasonis		1.26	0.19	ARA 94	1	
46735122	**Low discrimination**					**46735122**
					GRA	
Parabacteroides distasonis		76.07	0.58	GLU 99	1	
Bifidobacterium longum subsp. infantis		16.57	0.30		99	
Clostridium clostridioforme		3.76	0.30		50	
46735132	**Low discrimination**					**46735132**
					GRA	
Parabacteroides distasonis		70.11	0.53	GLU 99	1	
Clostridium clostridioforme		29.34	0.41		50	
Bifidobacterium longum subsp. infantis		0.29	-.04		99	
46735202	**Good identification**					**46735202**
Clostridium clostridioforme		100.0	0.36	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46735212	**Good identification**					**46735212**
Clostridium clostridioforme		100.0	0.47	GLU 88		
46735222	**Low discrimination**					**46735222**
Clostridium clostridioforme		78.15	0.36	GLU 88		
Parabacteroides distasonis		21.86	0.32	ARA 94		
46735232	**Good identification**					**46735232**
Clostridium clostridioforme		96.80	0.47	GLU 88		
Parabacteroides distasonis		3.20	0.27	ARA 94		
46735302	**Low discrimination**					**46735302**
					GRA	
Clostridium clostridioforme		89.53	0.45	GLU 88	50	
Parabacteroides distasonis		7.48	0.32	ARA 94	1	
Eubacterium contortum		2.14	0.11		99	
46735312	**Good identification**					**46735312**
					GRA	
Clostridium clostridioforme		98.90	0.56	GLU 88	50	
Parabacteroides distasonis		0.98	0.27	ARA 94	1	
Bifidobacterium adolescentis		0.12	0.09		99	
46735322	**Low discrimination**					**46735322**
Parabacteroides distasonis		88.70	0.66	GLU 99		
Clostridium clostridioforme		10.73	0.45			
Prevotella oralis		0.47	0.14			
46735332	**Low discrimination**					**46735332**
Clostridium clostridioforme		50.57	0.56	GLU 88		
Parabacteroides distasonis		49.43	0.61	ARA 94		
46735402	**Acceptable identification**					**46735402**
					SPO	
Clostridium clostridioforme		91.75	0.41	GLU 88	66	
Bifidobacterium adolescentis		4.32	0.22	RHA 88	1	
Clostridium butyricum		1.99	0.01		99	
46735412	**Good identification**					**46735412**
Clostridium clostridioforme		98.75	0.52	GLU 88		
Bifidobacterium adolescentis		1.25	0.23	RHA 88		
46735422	**Good identification**					**46735422**
Clostridium clostridioforme		97.40	0.41	GLU 88		
Parabacteroides distasonis		2.61	0.20	RHA 88		
46735432	**Good identification**					**46735432**
Clostridium clostridioforme		99.69	0.52	GLU 88		
Parabacteroides distasonis		0.32	0.15	RHA 88		
46735502	**Low discrimination**					**46735502**
					SPO	
Bifidobacterium adolescentis		53.91	0.56	GLU 99	1	
Clostridium clostridioforme		38.70	0.50		66	
Clostridium butyricum		6.01	0.25		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46735512	**Low discrimination**					**46735512**
					SPO	
Clostridium clostridioforme		72.51	0.61	GLU 88	66	
Bifidobacterium adolescentis		27.11	0.57	RHA 88	1	
Clostridium butyricum		0.13	0.03		99	
46735522	**Low discrimination**					**46735522**
					GRA	
Clostridium clostridioforme		54.41	0.50	GLU 88	50	
Parabacteroides distasonis		43.04	0.54	RHA 88	1	
Bifidobacterium adolescentis		0.77	0.22		99	
46735532	**Good identification**					**46735532**
					GRA	
Clostridium clostridioforme		90.67	0.61	GLU 88	50	
Parabacteroides distasonis		8.48	0.49	RHA 88	1	
Bifidobacterium adolescentis		0.34	0.23		99	
46735542	**Low discrimination**					**46735542**
					SPO	
Bifidobacterium adolescentis		83.74	0.40	GLU 99	1	
Clostridium butyricum		6.56	0.07	SOR 11	99	
Clostridium clostridioforme		4.91	0.16		66	
46735552	**Low discrimination**					**46735552**
Bifidobacterium adolescentis		82.06	0.42	GLU 99		
Clostridium clostridioforme		17.94	0.27	SOR 11		
46735602	**Good identification**					**46735602**
Clostridium clostridioforme		100.0	0.56	GLU 88		
46735612	**Very good identification**					**46735612**
Clostridium clostridioforme		100.0	0.67	GLU 88		
46735622	**Good identification**					**46735622**
Clostridium clostridioforme		98.92	0.56	GLU 88		
Parabacteroides distasonis		1.08	0.28			
46735632	**Very good identification**					**46735632**
Clostridium clostridioforme		99.87	0.67	GLU 88		
Parabacteroides distasonis		0.13	0.23			
46735702	**Very good identification**					**46735702**
					GRA	
Clostridium clostridioforme		99.30	0.65	GLU 88	50	
Parabacteroides distasonis		0.33	0.28		1	
Bifidobacterium adolescentis		0.19	0.22		99	
46735710	**Good identification**					**46735710**
Clostridium clostridioforme		100.0	0.42	GLU 88		
46735712	**Very good identification**					**46735712**
					GRA	
Clostridium clostridioforme		99.91	0.76	GLU 88	50	
Bifidobacterium adolescentis		0.05	0.23		99	
Parabacteroides distasonis		0.04	0.23		1	
46735713	**Good identification**					**46735713**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.42	GLU 88		
46735722	**Low discrimination**					**46735722**
Clostridium clostridioforme		74.78	0.65	GLU 88		
Parabacteroides distasonis		24.19	0.62			
Bacteroides thetaiotaomicron		0.88	0.22			
46735730	**Good identification**					**46735730**
Clostridium clostridioforme		95.01	0.42	GLU 88		
Parabacteroides distasonis		3.63	0.23	BGL 99		
Bacteroides vulgatus		1.36	-.02			
46735732	**Very good identification**					**46735732**
Clostridium clostridioforme		96.21	0.76	GLU 88		
Parabacteroides distasonis		3.68	0.58			
Bacteroides thetaiotaomicron		0.10	0.15			
46735733	**Good identification**					**46735733**
Clostridium clostridioforme		96.32	0.42	GLU 88		
Parabacteroides distasonis		3.68	0.23	URE 1		
46735752	**Good identification**					**46735752**
Clostridium clostridioforme		99.37	0.42	GLU 88		
Bifidobacterium adolescentis		0.63	0.08	SOR 1		
46735772	**Good identification**					**46735772**
Clostridium clostridioforme		96.32	0.42	GLU 88		
Parabacteroides distasonis		3.68	0.23	SOR 1		
46737002	**Good identification**					**46737002**
Actinomyces israelii		100.0	0.36	GLU 99		
46737100	**Good identification**					**46737100**
					SPO	
Actinomyces israelii		95.96	0.44	GLU 99	1	
Clostridium butyricum		4.04	-.01	CEL 93	99	
46737102	**Low discrimination**					**46737102**
Actinomyces israelii		50.74	0.47	GLU 99		
Bifidobacterium adolescentis		49.27	0.40	CEL 93		
46737112	**Low discrimination**					**46737112**
Bifidobacterium adolescentis		86.44	0.42	GLU 99		
Actinomyces israelii		10.88	0.33	ARA 87		
Bifidobacterium dentium		1.46	-.07			
46737140	**Good identification**					**46737140**
Actinomyces israelii		100.0	0.37	GLU 99		
46737142	**Low discrimination**					**46737142**
Actinomyces israelii		76.42	0.40	GLU 99		
Bifidobacterium adolescentis		23.58	0.25	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46737500	**Low discrimination**					**46737500**
					SPO	
Actinomyces israelii		63.14	0.42	GLU 99	1	
Clostridium butyricum		29.27	0.15	CEL 93	99	
Bifidobacterium adolescentis		7.60	0.20		1	
46737502	**Low discrimination**					**46737502**
					SPO	
Bifidobacterium adolescentis		86.73	0.54	GLU 99	1	
Actinomyces israelii		10.92	0.45		1	
Clostridium butyricum		1.31	0.08		99	
46737512	**Low discrimination**					**46737512**
Bifidobacterium adolescentis		77.03	0.56	GLU 99		
Bifidobacterium dentium		19.24	0.27			
Clostridium clostridioforme		2.54	0.27			
46737542	**Low discrimination**					**46737542**
					COC	
Bifidobacterium adolescentis		67.28	0.39	GLU 99	1	
Actinomyces israelii		26.65	0.38	SOR 11	1	
Blautia producta		6.07	0.07		99	
46737552	**Low discrimination**					**46737552**
					COC	
Bifidobacterium adolescentis		77.51	0.40	GLU 99	1	
Blautia producta		17.16	0.15	SOR 11	99	
Actinomyces israelii		3.75	0.24		1	
46737712	**Good identification**					**46737712**
Clostridium clostridioforme		95.05	0.42	GLU 88		
Bifidobacterium adolescentis		3.96	0.22	MAN 1		
Bifidobacterium dentium		0.99	-.07			
46737732	**Acceptable identification**					**46737732**
					GRA	
Clostridium clostridioforme		94.02	0.42	GLU 88	50	
Parabacteroides distasonis		3.60	0.23	MAN 1	1	
Bifidobacterium dentium		1.74	-.03		99	
46745712	**Good identification**					**46745712**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46745732	**Good identification**					**46745732**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46750100	**Low discrimination**					**46750100**
					GRA	
Fusobacterium mortiferum		88.25	0.39	GLU 99	1	
Collinsella aerofaciens		9.07	-.03	MLT 11	99	
Bifidobacterium breve		2.69	0.10		99	
46750612	**Good identification**					**46750612**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46750632	**Good identification**					**46750632**
Clostridium clostridioforme		100.0	0.36	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46750712	**Good identification**						**46750712**
Clostridium clostridioforme		100.0	0.45	GLU 88			
46750732	**Good identification**						**46750732**
Clostridium clostridioforme		100.0	0.45	GLU 88			
46751022	**Low discrimination**						**46751022**
					GRA	SPO	
Prevotella oralis		86.00	0.45	GLU 99	1	1	
Clostridium barati		5.22	0.24	LAC 94	99	99	
Bifidobacterium breve		3.04	0.32		99	1	
46751101	**Low discrimination**						**46751101**
Actinomyces naeslundii		67.55	0.41	ESL 12			
Collinsella aerofaciens		25.03	-.03	GLU 99			
Bifidobacterium breve		7.43	0.10				
46751102	**Low discrimination**						**46751102**
					GRA		
Bifidobacterium breve		54.62	0.62	GLU 99	99		
Prevotella oralis		22.87	0.44		1		
Lactobacillus acidophilus		8.96	0.38		99		
46751110	**Good identification**						**46751110**
					GRA		
Bifidobacterium breve		95.91	0.44	GLU 99	99		
Collinsella aerofaciens		3.27	-.03	BGL 91	99		
Leptotrichia buccalis		0.83	-.11		1		
46751112	**Low discrimination**						**46751112**
					GRA		
Bifidobacterium breve		87.66	0.62	GLU 99	99		
Leptotrichia buccalis		7.38	0.24		1		
Bifidobacterium adolescentis		3.95	0.38		99		
46751120	**Low discrimination**						**46751120**
					GRA	SPO	
Clostridium paraputrificum		62.96	0.47	GLU 99	99	99	
Prevotella oralis		18.96	0.44	RAF 7	1	1	
Bifidobacterium breve		7.96	0.49		99	1	
46751122	**Low discrimination**						**46751122**
					GRA	SPO	
Prevotella oralis		73.39	0.66	GLU 99	1	1	
Bifidobacterium breve		16.40	0.66		99	1	
Clostridium paraputrificum		5.50	0.40		99	99	
46751130	**Acceptable identification**						**46751130**
					GRA	SPO	
Bifidobacterium breve		90.09	0.49	GLU 99	99	1	
Clostridium paraputrificum		7.20	0.13	BGL 91	99	99	
Prevotella oralis		2.17	0.10		1	1	
46751132	**Good identification**						**46751132**
					GRA		
Bifidobacterium breve		94.21	0.66	GLU 99	99		
Prevotella oralis		4.26	0.32		1		
Leptotrichia buccalis		0.50	0.07		1		
46751140	**Acceptable identification**						**46751140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		93.12	0.44	GLU 99			
Collinsella aerofaciens		3.17	-.03	BGL 91			
Actinomyces naeslundii		1.01	0.25				
46751142	**Very good identification**						**46751142**
Bifidobacterium breve		96.63	0.62	GLU 99			
Lactobacillus acidophilus		1.96	0.23				
Bifidobacterium adolescentis		0.44	0.21				
46751150	**Good identification**						**46751150**
Bifidobacterium breve		100.0	0.44	GLU 99			
46751152	**Very good identification**						**46751152**
					GRA		
Bifidobacterium breve		99.36	0.62	GLU 99	99		
Bifidobacterium adolescentis		0.55	0.22		99		
Leptotrichia buccalis		0.08	-.11		1		
46751160	**Low discrimination**						**46751160**
					GRA	SPO	
Bifidobacterium breve		60.73	0.49	GLU 99	99	1	
Clostridium paraputrificum		36.15	0.27	BGL 91	99	99	
Prevotella oralis		1.46	0.10		1	1	
46751162	**Good identification**						**46751162**
					GRA	SPO	
Bifidobacterium breve		91.87	0.66	GLU 99	99	1	
Prevotella oralis		4.15	0.32		1	1	
Clostridium paraputrificum		2.32	0.21		99	99	
46751170	**Good identification**						**46751170**
					SPO		
Bifidobacterium breve		99.40	0.49	GLU 99	1		
Clostridium paraputrificum		0.60	-.07	BGL 91	99		
46751172	**Very good identification**						**46751172**
					GRA		
Bifidobacterium breve		99.95	0.66	GLU 99	99		
Prevotella oralis		0.05	-.03		1		
46751222	**Good identification**						**46751222**
					GRA	SPO	
Prevotella oralis		96.90	0.39	GLU 99	1	1	
Clostridium barati		1.19	0.05	LAC 94	99	99	
Clostridium clostridioforme		1.05	0.13		50	66	
46751302	**Low discrimination**						**46751302**
					GRA		
Prevotella oralis		83.85	0.38	GLU 99	1		
Lactobacillus acidophilus		5.77	0.19	NAG 95	99		
Bifidobacterium breve		4.72	0.27		99		
46751320	**Low discrimination**						**46751320**
					SPO	GRA	
Prevotella oralis		81.49	0.38	GLU 99	1	1	
Clostridium ramosum		9.63	0.22	BGL 95	92	50	
Clostridium paraputrificum		6.38	0.13		99	99	
46751322	**Very good identification**						**46751322**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		95.59	0.60	GLU 99	1	
Clostridium ramosum		2.38	0.33		92	
Parabacteroides distasonis		0.53	0.34		1	
46751502	**Low discrimination**					**46751502**
					SPO	
Bifidobacterium adolescentis		83.81	0.50	GLU 99	1	
Clostridium clostridioforme		5.70	0.27	XYL 87	66	
Clostridium butyricum		3.83	0.13		99	
46751512	**Low discrimination**					**46751512**
Bifidobacterium adolescentis		77.52	0.52	GLU 99		
Clostridium clostridioforme		19.66	0.38	XYL 87		
Bifidobacterium breve		2.60	0.27			
46751532	**Low discrimination**					**46751532**
Clostridium clostridioforme		76.19	0.38	GLU 88		
Bifidobacterium breve		17.89	0.32	TRE 88		
Bifidobacterium adolescentis		3.04	0.18	XYL 91		
46751552	**Low discrimination**					**46751552**
Bifidobacterium adolescentis		77.42	0.36	GLU 99		
Bifidobacterium breve		20.98	0.27	XYL 87		
Clostridium clostridioforme		1.61	0.04	SOR 11		
46751612	**Good identification**					**46751612**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46751632	**Good identification**					**46751632**
Clostridium clostridioforme		100.0	0.44	GLU 88		
46751702	**Good identification**					**46751702**
					GRA	
Clostridium clostridioforme		96.62	0.42	GLU 88	50	
Bifidobacterium adolescentis		1.96	0.16	TRE 88	99	
Prevotella oralis		1.42	0.04	XYL 91	1	
46751712	**Good identification**					**46751712**
Clostridium clostridioforme		99.46	0.53	GLU 88		
Bifidobacterium adolescentis		0.54	0.18	TRE 88		
46751722	**Low discrimination**					**46751722**
Clostridium clostridioforme		70.87	0.42	GLU 88		
Prevotella oralis		19.83	0.25	TRE 88		
Parabacteroides distasonis		6.67	0.30	XYL 91		
46751732	**Good identification**					**46751732**
Clostridium clostridioforme		98.90	0.53	GLU 88		
Parabacteroides distasonis		1.10	0.25	TRE 88		
46753000	**Low discrimination**					**46753000**
Actinomyces naeslundii		48.79	0.35	ESL 12		
Actinomyces israelii		39.42	0.25	GLU 99		
Bifidobacterium breve		11.80	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46753001	**Good identification**					**46753001**
Actinomyces naeslundii		100.0	0.38	ESL 12		
46753003	**Good identification**					**46753003**
Actinomyces naeslundii		100.0	0.38	ESL 12		
46753020	**Low discrimination**					**46753020**
					SPO	
Actinomyces naeslundii		53.54	0.35	ESL 12	1	
Bifidobacterium breve		23.01	0.15	GLU 99	1	
Clostridium ramosum		11.87	-.01		92	
46753021	**Good identification**					**46753021**
Actinomyces naeslundii		100.0	0.38	ESL 12		
46753023	**Good identification**					**46753023**
Actinomyces naeslundii		100.0	0.38	ESL 12		
46753100	**Low discrimination**					**46753100**
Bifidobacterium breve		76.55	0.44	GLU 99		
Actinomyces israelii		10.34	0.35	BGL 91		
Actinomyces naeslundii		9.10	0.43			
46753101	**Acceptable identification**					**46753101**
Actinomyces naeslundii		94.64	0.46	ESL 12		
Bifidobacterium breve		5.36	0.10	GLU 99		
46753102	**Good identification**					**46753102**
Bifidobacterium breve		92.35	0.62	GLU 99		
Bifidobacterium adolescentis		2.79	0.35			
Actinomyces israelii		1.85	0.38			
46753103	**Low discrimination**					**46753103**
Actinomyces naeslundii		63.59	0.46	ESL 12		
Bifidobacterium breve		36.41	0.27	GLU 99		
46753110	**Good identification**					**46753110**
Bifidobacterium breve		98.02	0.44	GLU 99		
Actinomyces israelii		1.98	0.21	BGL 91		
46753112	**Very good identification**					**46753112**
Bifidobacterium breve		95.94	0.62	GLU 99		
Bifidobacterium adolescentis		3.54	0.36			
Actinomyces israelii		0.29	0.24			
46753120	**Low discrimination**					**46753120**
					SPO	
Bifidobacterium breve		58.21	0.49	GLU 99	1	
Clostridium ramosum		30.02	0.33	BGL 91	92	
Clostridium paraputrificum		4.65	0.13		99	
46753121	**Low discrimination**					**46753121**
					SPO	
Actinomyces naeslundii		86.76	0.46	ESL 12	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		8.73	0.15	GLU 99	1		
Clostridium ramosum		4.51	-.01		92		
46753122	**Low discrimination**						**46753122**
					SPO	GRA	
Bifidobacterium breve		78.64	0.66	GLU 99	1	99	
Clostridium ramosum		16.05	0.43		92	50	
Prevotella oralis		3.56	0.32		1	1	
46753130	**Good identification**						**46753130**
					SPO		
Bifidobacterium breve		98.58	0.49	GLU 99	1		
Clostridium tertium		0.91	0.05	BGL 91	99		
Clostridium ramosum		0.51	-.01		92		
46753132	**Very good identification**						**46753132**
					SPO		
Bifidobacterium breve		99.48	0.66	GLU 99	1		
Clostridium tertium		0.27	0.13		99		
Clostridium ramosum		0.21	0.09		92		
46753140	**Acceptable identification**						**46753140**
Bifidobacterium breve		93.15	0.44	GLU 99			
Actinomyces israelii		4.89	0.28	BGL 91			
Actinomyces naeslundii		1.96	0.30				
46753142	**Very good identification**						**46753142**
Bifidobacterium breve		98.56	0.62	GLU 99			
Actinomyces israelii		0.77	0.31				
Bifidobacterium adolescentis		0.37	0.19				
46753150	**Good identification**						**46753150**
Bifidobacterium breve		99.22	0.44	GLU 99			
Actinomyces israelii		0.78	0.14	BGL 91			
46753152	**Very good identification**						**46753152**
Bifidobacterium breve		99.43	0.62	GLU 99			
Bifidobacterium adolescentis		0.45	0.21				
Actinomyces israelii		0.12	0.17				
46753160	**Good identification**						**46753160**
					SPO		
Bifidobacterium breve		97.75	0.49	GLU 99	1		
Actinomyces naeslundii		1.15	0.30	BGL 91	1		
Clostridium paraputrificum		0.59	-.07		99		
46753162	**Very good identification**						**46753162**
					SPO		
Bifidobacterium breve		99.57	0.66	GLU 99	1		
Clostridium ramosum		0.21	0.09		92		
Actinomyces naeslundii		0.12	0.30		1		
46753170	**Good identification**						**46753170**
Bifidobacterium breve		100.0	0.49	GLU 99			
46753172	**Very good identification**						**46753172**
Bifidobacterium breve		100.0	0.66	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46753320	**Good identification**						**46753320**
					SPO	GRA	
Clostridium ramosum		97.10	0.35	GLU 99	92	50	
Prevotella oralis		1.47	0.04	TRE 94	1	1	
Bifidobacterium breve		1.43	0.15		1	99	
46753322	**Acceptable identification**						**46753322**
					SPO	GRA	
Clostridium ramosum		90.17	0.45	GLU 99	92	50	
Prevotella oralis		6.46	0.25	TRE 94	1	1	
Bifidobacterium breve		3.37	0.32		1	99	
46753502	**Low discrimination**						**46753502**
Bifidobacterium adolescentis		87.92	0.49	GLU 99			
Actinomyces israelii		7.14	0.36	XYL 87			
Bifidobacterium breve		4.40	0.27				
46753512	**Acceptable identification**						**46753512**
Bifidobacterium adolescentis		94.40	0.50	GLU 99			
Bifidobacterium breve		3.87	0.27	XYL 87			
Actinomyces israelii		0.94	0.22				
46754312	**Good identification**						**46754312**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46754332	**Good identification**						**46754332**
Clostridium clostridioforme		99.24	0.42	GLU 88			
Parabacteroides distasonis		0.76	0.11	TRE 88			
46754412	**Good identification**						**46754412**
Clostridium clostridioforme		100.0	0.38	GLU 88			
46754432	**Good identification**						**46754432**
Clostridium clostridioforme		100.0	0.38	GLU 88			
46754502	**Low discrimination**						**46754502**
					SPO		
Clostridium clostridioforme		73.54	0.36	GLU 88	66		
Bifidobacterium adolescentis		21.68	0.30	TRE 88	1		
Clostridium butyricum		4.79	0.04	RHA 88	99		
46754512	**Acceptable identification**						**46754512**
Clostridium clostridioforme		92.67	0.47	GLU 88			
Bifidobacterium adolescentis		7.33	0.32	TRE 88			
46754522	**Good identification**						**46754522**
Clostridium clostridioforme		95.46	0.36	GLU 88			
Bacteroides uniformis		4.54	0.00	TRE 88			
46754532	**Good identification**						**46754532**
Clostridium clostridioforme		100.0	0.47	GLU 88			
46754602	**Good identification**						**46754602**
Clostridium clostridioforme		100.0	0.42	GLU 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46754612	**Good identification**						**46754612**
Clostridium clostridioforme		100.0	0.53	GLU 88			
46754622	**Good identification**						**46754622**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46754632	**Good identification**						**46754632**
Clostridium clostridioforme		100.0	0.53	GLU 88			
46754702	**Good identification**						**46754702**
Clostridium clostridioforme		100.0	0.51	GLU 88			
46754712	**Very good identification**						**46754712**
Clostridium clostridioforme		100.0	0.62	GLU 88			
46754722	**Good identification**						**46754722**
Clostridium clostridioforme		98.86	0.51	GLU 88			
Bacteroides uniformis		0.89	0.03	TRE 88			
Parabacteroides distasonis		0.25	0.12				
46754732	**Very good identification**						**46754732**
Clostridium clostridioforme		100.0	0.62	GLU 88			
46755102	**Low discrimination**						**46755102**
					GRA		
Bifidobacterium adolescentis		53.38	0.50	GLU 99	99		
Bifidobacterium longum subsp. infantis		12.72	0.26	ARA 87	99		
Prevotella oralis		11.20	0.28		1		
46755112	**Low discrimination**						**46755112**
Bifidobacterium adolescentis		81.49	0.52	GLU 99			
Clostridium clostridioforme		13.34	0.35	ARA 87			
Bifidobacterium breve		2.73	0.27				
46755122	**Low discrimination**						**46755122**
					CAT	GRA	
Prevotella oralis		73.83	0.50	GLU 99	1	1	
Bacteroides fragilis		8.63	0.25	XYL 11	99	1	
Bifidobacterium longum subsp. infantis		7.85	0.30		1	99	
46755132	**Low discrimination**						**46755132**
					GRA		
Clostridium clostridioforme		43.66	0.35	GLU 88	50		
Parabacteroides distasonis		27.01	0.37	TRE 88	1		
Bifidobacterium breve		15.88	0.32	RHA 88	99		
46755152	**Low discrimination**						**46755152**
Bifidobacterium adolescentis		74.94	0.36	GLU 99			
Bifidobacterium breve		20.31	0.27	ARA 87			
Actinomyces israelii		4.75	0.22	SOR 11			
46755212	**Good identification**						**46755212**
Clostridium clostridioforme		100.0	0.41	GLU 88			
46755232	**Good identification**						**46755232**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		99.15	0.41	GLU 88			
Parabacteroides distasonis		0.85	0.11	TRE 88			
46755302	**Low discrimination**						**46755302**
					GRA		
Clostridium clostridioforme		69.50	0.39	GLU 88	50		
Prevotella oralis		19.41	0.22	TRE 88	1		
Eubacterium contortum		3.64	0.11	ARA 94	99		
46755312	**Good identification**						**46755312**
					GRA		
Clostridium clostridioforme		98.92	0.50	GLU 88	50		
Bifidobacterium adolescentis		0.83	0.18	TRE 88	99		
Parabacteroides distasonis		0.25	0.11	ARA 94	1		
46755322	**Low discrimination**						**46755322**
Prevotella oralis		59.53	0.44	GLU 99			
Parabacteroides distasonis		24.03	0.50	XYL 11			
Clostridium clostridioforme		11.22	0.39				
46755332	**Low discrimination**						**46755332**
Clostridium clostridioforme		78.93	0.50	GLU 88			
Parabacteroides distasonis		19.98	0.45	TRE 88			
Prevotella oralis		0.93	0.10	ARA 94			
46755402	**Low discrimination**						**46755402**
					SPO		
Clostridium clostridioforme		55.78	0.35	GLU 88	66		
Bifidobacterium adolescentis		18.35	0.30	TRE 88	1		
Clostridium butyricum		16.72	0.15	RHA 88	99		
46755412	**Acceptable identification**						**46755412**
Clostridium clostridioforme		91.15	0.47	GLU 88			
Bifidobacterium adolescentis		8.05	0.32	TRE 88			
Actinomyces israelii		0.49	0.17	RHA 88			
46755422	**Low discrimination**						**46755422**
					GRA	SPO	
Clostridium clostridioforme		88.24	0.35	GLU 88	50	66	
Bacteroides uniformis		9.46	0.06	TRE 88	1	1	
Clostridium butyricum		2.30	-.03	RHA 88	99	99	
46755432	**Good identification**						**46755432**
Clostridium clostridioforme		100.0	0.47	GLU 88			
46755500	**Good identification**						**46755500**
					SPO		
Clostridium butyricum		95.85	0.45	GLU 99	99		
Actinomyces israelii		2.27	0.39	TRE 94	1		
Bifidobacterium adolescentis		1.71	0.30		1		
46755502	**Low discrimination**						**46755502**
					SPO		
Bifidobacterium adolescentis		74.33	0.64	GLU 99	1		
Clostridium butyricum		16.42	0.38		99		
Clostridium clostridioforme		7.64	0.44		66		
46755512	**Low discrimination**						**46755512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Bifidobacterium adolescentis		71.65	0.66	GLU 99	1	
Clostridium clostridioforme		27.45	0.56		66	
Clostridium butyricum		0.68	0.17		99	
46755532	**Acceptable identification**					**46755532**
					GRA	
Clostridium clostridioforme		94.26	0.56	GLU 88	50	
Bifidobacterium adolescentis		2.49	0.32	TRE 88	99	
Parabacteroides distasonis		2.28	0.34	RHA 88	1	
46755542	**Low discrimination**					**46755542**
					SPO	
Bifidobacterium adolescentis		81.49	0.49	GLU 99	1	
Clostridium butyricum		12.66	0.20	SOR 11	99	
Actinomyces israelii		5.16	0.35		1	
46755552	**Good identification**					**46755552**
Bifidobacterium adolescentis		95.63	0.50	GLU 99		
Clostridium clostridioforme		2.99	0.21	SOR 11		
Actinomyces israelii		0.74	0.21			
46755602	**Good identification**					**46755602**
Clostridium clostridioforme		100.0	0.50	GLU 88		
46755612	**Very good identification**					**46755612**
Clostridium clostridioforme		100.0	0.61	GLU 88		
46755622	**Good identification**					**46755622**
Clostridium clostridioforme		97.75	0.50	GLU 88		
Bacteroides uniformis		1.97	0.08	TRE 88		
Parabacteroides distasonis		0.28	0.12			
46755632	**Very good identification**					**46755632**
Clostridium clostridioforme		100.0	0.61	GLU 88		
46755702	**Very good identification**					**46755702**
					SPO	
Clostridium clostridioforme		96.99	0.59	GLU 88	66	
Bifidobacterium adolescentis		1.30	0.30	TRE 88	1	
Clostridium butyricum		0.88	0.13		99	
46755710	**Good identification**					**46755710**
Clostridium clostridioforme		97.40	0.36	GLU 88		
Clostridium butyricum		2.60	-.02	TRE 88		
46755712	**Very good identification**					**46755712**
Clostridium clostridioforme		99.64	0.70	GLU 88		
Bifidobacterium adolescentis		0.36	0.32	TRE 88		
46755713	**Good identification**					**46755713**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46755722	**Low discrimination**					**46755722**
Clostridium clostridioforme		70.44	0.59	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		20.82	0.37	TRE 88		
Parabacteroides distasonis		5.90	0.46			
46755730	**Good identification**					**46755730**
Clostridium clostridioforme		96.19	0.36	GLU 88		
Bacteroides vulgatus		3.81	-.01	TRE 88		
46755732	**Very good identification**					**46755732**
Clostridium clostridioforme		98.49	0.70	GLU 88		
Parabacteroides distasonis		0.98	0.42	TRE 88		
Bacteroides ovatus		0.44	0.20			
46755733	**Good identification**					**46755733**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46755752	**Good identification**					**46755752**
Clostridium clostridioforme		95.79	0.36	GLU 88		
Bifidobacterium adolescentis		4.21	0.16	TRE 88		
46755772	**Good identification**					**46755772**
Clostridium clostridioforme		98.21	0.36	GLU 88		
Bacteroides ovatus		1.79	-.04	TRE 88		
46757000	**Good identification**					**46757000**
Actinomyces israelii		100.0	0.43	GLU 99		
46757002	**Good identification**					**46757002**
Actinomyces israelii		96.58	0.46	GLU 99		
Bifidobacterium adolescentis		3.42	0.15			
46757040	**Good identification**					**46757040**
Actinomyces israelii		100.0	0.36	GLU 99		
46757042	**Good identification**					**46757042**
Actinomyces israelii		100.0	0.39	GLU 99		
46757100	**Acceptable identification**					**46757100**
					SPO	
Actinomyces israelii		91.31	0.53	GLU 99	1	
Clostridium butyricum		6.88	0.13		99	
Bifidobacterium adolescentis		1.21	0.15		1	
46757102	**Low discrimination**					**46757102**
Actinomyces israelii		51.41	0.56	GLU 99		
Bifidobacterium adolescentis		45.05	0.49			
Bifidobacterium breve		2.25	0.27			
46757110	**Low discrimination**					**46757110**
Actinomyces israelii		84.41	0.39	GLU 99		
Bifidobacterium adolescentis		9.17	0.16	MLZ 13		
Bifidobacterium breve		3.68	0.10			
46757112	**Low discrimination**					**46757112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		83.70	0.50	GLU 99		
Actinomyces israelii		11.68	0.42	ARA 87		
Bifidobacterium breve		3.43	0.27			
46757140	**Good identification**					**46757140**
					SPO	
Actinomyces israelii		96.75	0.46	GLU 99	1	
Clostridium butyricum		1.63	-.05		99	
Bifidobacterium breve		1.62	0.10		1	
46757142	**Low discrimination**					**46757142**
Actinomyces israelii		71.88	0.49	GLU 99		
Bifidobacterium adolescentis		20.02	0.33			
Bifidobacterium breve		8.11	0.27			
46757302	**Low discrimination**					**46757302**
Actinomyces israelii		85.92	0.37	GLU 99		
Bifidobacterium adolescentis		10.11	0.15	RHA 7		
Clostridium clostridioforme		3.98	0.05			
46757400	**Low discrimination**					**46757400**
					SPO	
Actinomyces israelii		87.86	0.41	GLU 99	1	
Clostridium butyricum		12.14	0.05		99	
46757402	**Low discrimination**					**46757402**
					SPO	
Actinomyces israelii		75.44	0.44	GLU 99	1	
Bifidobacterium adolescentis		21.85	0.29		1	
Clostridium butyricum		2.70	-.02		99	
46757442	**Acceptable identification**					**46757442**
Actinomyces israelii		91.57	0.37	GLU 99		
Bifidobacterium adolescentis		8.43	0.13			
46757500	**Low discrimination**					**46757500**
					SPO	
Actinomyces israelii		51.60	0.52	GLU 99	1	
Clostridium butyricum		42.80	0.29		99	
Bifidobacterium adolescentis		5.61	0.29		1	
46757502	**Low discrimination**					**46757502**
					SPO	
Bifidobacterium adolescentis		85.34	0.63	GLU 99	1	
Actinomyces israelii		11.91	0.55		1	
Clostridium butyricum		2.56	0.22		99	
46757510	**Low discrimination**					**46757510**
Actinomyces israelii		34.12	0.37	GLU 99		
Bifidobacterium adolescentis		30.31	0.30	MLZ 13		
Bifidobacterium dentium		23.26	0.10			
46757512	**Good identification**					**46757512**
Bifidobacterium adolescentis		90.79	0.64	GLU 99		
Bifidobacterium dentium		7.11	0.27			
Actinomyces israelii		1.55	0.40			
46757540	**Low discrimination**					**46757540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		47.85	0.45	GLU 99	1	1	
Blautia producta		41.63	0.24		99	1	
Clostridium butyricum		8.87	0.11		1	99	
46757542	**Low discrimination**						**46757542**
					COC		
Bifidobacterium adolescentis		67.19	0.47	GLU 99	1		
Actinomyces israelii		29.49	0.48	SOR 11	1		
Blautia producta		1.90	0.07		99		
46757552	**Low discrimination**						**46757552**
					COC		
Bifidobacterium adolescentis		88.53	0.49	GLU 99	1		
Blautia producta		6.15	0.15	SOR 11	99		
Actinomyces israelii		4.75	0.33		1		
46757702	**Low discrimination**						**46757702**
Actinomyces israelii		34.05	0.35	GLU 99			
Bifidobacterium adolescentis		32.76	0.29	RHA 7			
Clostridium clostridioforme		30.18	0.25				
46757712	**Low discrimination**						**46757712**
Clostridium clostridioforme		74.02	0.36	GLU 88			
Bifidobacterium adolescentis		21.55	0.30	TRE 88			
Actinomyces israelii		2.74	0.21	MAN 1			
46757732	**Low discrimination**						**46757732**
					GRA		
Clostridium clostridioforme		84.28	0.36	GLU 88	50		
Bacteroides ovatus		12.30	0.12	TRE 88	1		
Bifidobacterium dentium		3.42	-.03	MAN 1	99		
46764712	**Good identification**						**46764712**
Clostridium clostridioforme		100.0	0.43	GLU 88			
46764732	**Good identification**						**46764732**
Clostridium clostridioforme		100.0	0.43	GLU 88			
46765512	**Low discrimination**						**46765512**
					SPO		
Clostridium clostridioforme		76.16	0.36	GLU 88	66		
Bifidobacterium adolescentis		18.07	0.29	FRU 99	1		
Clostridium butyricum		4.04	0.03	RHA 88	99		
46765532	**Acceptable identification**						**46765532**
					GRA		
Clostridium clostridioforme		94.73	0.36	GLU 88	50		
Bifidobacterium dentium		3.84	-.03	FRU 99	99		
Parabacteroides distasonis		1.42	0.11	RHA 88	1		
46765612	**Good identification**						**46765612**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46765632	**Good identification**						**46765632**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46765702	**Good identification**						**46765702**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		98.10	0.40	GLU 88		
Clostridium butyricum		1.90	-.01	FRU 99		
46765712	**Good identification**					**46765712**
Clostridium clostridioforme		100.0	0.51	GLU 88		
46765722	**Acceptable identification**					**46765722**
Clostridium clostridioforme		91.43	0.40	GLU 88		
Parabacteroides distasonis		4.76	0.23	FRU 99		
Bacteroides ovatus		3.81	0.06			
46765732	**Good identification**					**46765732**
Clostridium clostridioforme		98.79	0.51	GLU 88		
Parabacteroides distasonis		0.61	0.19	FRU 99		
Bacteroides ovatus		0.60	0.03			
46770312	**Good identification**					**46770312**
Clostridium clostridioforme		99.04	0.39	GLU 88		
Leptotrichia buccalis		0.96	-.11	XYL 91		
46770332	**Good identification**					**46770332**
Clostridium clostridioforme		100.0	0.39	GLU 88		
46770412	**Good identification**					**46770412**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46770432	**Good identification**					**46770432**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46770512	**Good identification**					**46770512**
					GRA	
Clostridium clostridioforme		98.48	0.45	GLU 88	50	
Bifidobacterium adolescentis		1.07	0.15	XYL 91	99	
Leptotrichia buccalis		0.45	-.11	RHA 88	1	
46770532	**Good identification**					**46770532**
Clostridium clostridioforme		100.0	0.45	GLU 88		
46770602	**Good identification**					**46770602**
Clostridium clostridioforme		100.0	0.40	GLU 88		
46770612	**Good identification**					**46770612**
Clostridium clostridioforme		100.0	0.51	GLU 88		
46770622	**Good identification**					**46770622**
Clostridium clostridioforme		100.0	0.40	GLU 88		
46770632	**Good identification**					**46770632**
Clostridium clostridioforme		100.0	0.51	GLU 88		
46770702	**Good identification**					**46770702**
Clostridium clostridioforme		100.0	0.49	GLU 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46770712	**Very good identification**						**46770712**
Clostridium clostridioforme		100.0	0.60	GLU 88			
46770722	**Good identification**						**46770722**
Clostridium clostridioforme		100.0	0.49	GLU 88			
46770732	**Very good identification**						**46770732**
Clostridium clostridioforme		100.0	0.60	GLU 88			
46771001	**Good identification**						**46771001**
Actinomyces naeslundii		100.0	0.38	ESL 12			
46771003	**Good identification**						**46771003**
Actinomyces naeslundii		100.0	0.38	ESL 12			
46771021	**Good identification**						**46771021**
Actinomyces naeslundii		100.0	0.38	ESL 12			
46771023	**Good identification**						**46771023**
Actinomyces naeslundii		100.0	0.38	ESL 12			
46771100	**Low discrimination**						**46771100**
					COC	SPO	
Bifidobacterium breve		32.93	0.41	GLU 99	1	1	
Clostridium butyricum		25.21	0.24	BGL 91	1	99	
Atopobium parvulum		15.07	0.27		99	1	
46771101	**Good identification**						**46771101**
Actinomyces naeslundii		100.0	0.46	ESL 12			
46771102	**Low discrimination**						**46771102**
					GRA		
Bifidobacterium breve		46.75	0.59	GLU 99	99		
Lactobacillus acidophilus		38.52	0.47		99		
Leptotrichia buccalis		5.90	0.24		1		
46771103	**Low discrimination**						**46771103**
Actinomyces naeslundii		57.32	0.46	ESL 12			
Bifidobacterium breve		21.88	0.24	GLU 99			
Lactobacillus acidophilus		18.03	0.13				
46771112	**Good identification**						**46771112**
					GRA		
Leptotrichia buccalis		91.83	0.58	GLU 99	1		
Bifidobacterium breve		7.35	0.59		99		
Clostridium clostridioforme		0.39	0.33		50		
46771120	**Low discrimination**						**46771120**
					SPO		
Clostridium paraputrificum		46.27	0.37	GLU 99	99		
Clostridium ramosum		31.38	0.41	RAF 7	92		
Bifidobacterium breve		14.67	0.46		1		
46771121	**Low discrimination**						**46771121**
					SPO		
Actinomyces naeslundii		70.31	0.46	ESL 12	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		14.88	0.03	GLU 99	99		
Clostridium ramosum		10.09	0.06		92		
46771122	**Low discrimination**						**46771122**
					SPO		
Bifidobacterium breve		34.10	0.63	GLU 99	1		
Clostridium ramosum		28.86	0.51		92		
Lactobacillus acidophilus		15.81	0.47		1		
46771123	**Low discrimination**						**46771123**
					SPO		
Actinomyces naeslundii		33.91	0.46	ESL 12	1		
Bifidobacterium breve		23.01	0.29	GLU 99	1		
Clostridium ramosum		19.47	0.17		92		
46771130	**Low discrimination**						**46771130**
					GRA	SPO	
Bifidobacterium breve		88.24	0.46	GLU 99	99	1	
Leptotrichia buccalis		7.04	0.07	BGL 91	1	1	
Clostridium paraputrificum		2.81	0.03		99	99	
46771132	**Low discrimination**						**46771132**
					GRA		
Bifidobacterium breve		54.25	0.63	GLU 99	99		
Leptotrichia buccalis		42.38	0.41		1		
Clostridium clostridioforme		1.64	0.33		50		
46771140	**Low discrimination**						**46771140**
					SPO		
Bifidobacterium breve		86.93	0.41	GLU 99	1		
Clostridium butyricum		5.79	0.06	BGL 91	99		
Actinomyces israelii		4.55	0.25		1		
46771142	**Low discrimination**						**46771142**
Bifidobacterium breve		89.20	0.59	GLU 99			
Lactobacillus acidophilus		9.08	0.32				
Actinomyces israelii		0.69	0.28				
46771150	**Good identification**						**46771150**
					GRA		
Bifidobacterium breve		98.73	0.41	GLU 99	99		
Leptotrichia buccalis		1.27	-.11	BGL 91	1		
46771152	**Low discrimination**						**46771152**
					GRA		
Bifidobacterium breve		88.26	0.59	GLU 99	99		
Leptotrichia buccalis		11.15	0.24		1		
Bifidobacterium adolescentis		0.49	0.19		99		
46771160	**Low discrimination**						**46771160**
					SPO		
Bifidobacterium breve		78.33	0.46	GLU 99	1		
Clostridium paraputrificum		18.59	0.18	BGL 91	99		
Clostridium ramosum		1.69	0.06		92		
46771162	**Good identification**						**46771162**
					SPO		
Bifidobacterium breve		92.42	0.63	GLU 99	1		
Lactobacillus acidophilus		5.29	0.32		1		
Clostridium paraputrificum		0.93	0.11		99		
46771170	**Good identification**						**46771170**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		100.0	0.46	GLU 99			
46771172	**Very good identification**						**46771172**
					GRA		
Bifidobacterium breve		99.22	0.63	GLU 99	99		
Leptotrichia buccalis		0.78	0.07		1		
46771212	**Good identification**						**46771212**
Clostridium clostridioforme		98.93	0.39	GLU 88			
Leptotrichia buccalis		1.07	-.11	XYL 91			
46771232	**Good identification**						**46771232**
Clostridium clostridioforme		100.0	0.39	GLU 88			
46771302	**Low discrimination**						**46771302**
					SPO		
Clostridium clostridioforme		43.23	0.36	GLU 88	66		
Lactobacillus acidophilus		31.01	0.28	XYL 91	1		
Clostridium ramosum		10.07	0.19	ARA 94	92		
46771312	**Low discrimination**						**46771312**
					GRA		
Clostridium clostridioforme		74.28	0.48	GLU 88	50		
Leptotrichia buccalis		23.82	0.24	XYL 91	1		
Bifidobacterium breve		1.91	0.24	ARA 94	99		
46771320	**Good identification**						**46771320**
					SPO	GRA	
Clostridium ramosum		95.07	0.43	GLU 99	92	50	
Prevotella oralis		3.28	0.17		1	1	
Clostridium paraputrificum		1.07	0.03		99	99	
46771322	**Low discrimination**						**46771322**
					SPO		
Clostridium ramosum		76.15	0.53	GLU 99	92		
Prevotella oralis		12.47	0.39		1		
Parabacteroides distasonis		4.93	0.45		1		
46771332	**Low discrimination**						**46771332**
					SPO		
Clostridium clostridioforme		75.91	0.48	GLU 88	66		
Parabacteroides distasonis		13.41	0.40	XYL 91	1		
Clostridium ramosum		3.88	0.19	ARA 94	92		
46771412	**Good identification**						**46771412**
					GRA		
Clostridium clostridioforme		98.31	0.44	GLU 88	50		
Bifidobacterium adolescentis		1.19	0.15	XYL 91	99		
Leptotrichia buccalis		0.50	-.11	RHA 88	1		
46771432	**Good identification**						**46771432**
Clostridium clostridioforme		100.0	0.44	GLU 88			
46771500	**Good identification**						**46771500**
					SPO		
Clostridium butyricum		97.81	0.40	GLU 99	99		
Actinomyces israelii		1.56	0.30	XYL 97	1		
Bifidobacterium adolescentis		0.36	0.13		1		
46771512	**Low discrimination**						**46771512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Clostridium clostridioforme		64.84	0.53	GLU 88	50		
Bifidobacterium adolescentis		23.24	0.49	XYL 91	99		
Leptotrichia buccalis		9.73	0.24	RHA 88	1		
46771522	**Low discrimination**						**46771522**
					GRA	SPO	
Clostridium clostridioforme		64.08	0.42	GLU 88	50	66	
Parabacteroides distasonis		9.16	0.33	XYL 91	1	1	
Clostridium butyricum		8.00	0.15	RHA 88	99	99	
46771532	**Acceptable identification**						**46771532**
					GRA		
Clostridium clostridioforme		93.97	0.53	GLU 88	50		
Bifidobacterium breve		2.01	0.29	XYL 91	99		
Parabacteroides distasonis		1.59	0.29	RHA 88	1		
46771602	**Good identification**						**46771602**
Clostridium clostridioforme		100.0	0.48	GLU 88			
46771612	**Very good identification**						**46771612**
Clostridium clostridioforme		100.0	0.59	GLU 88			
46771622	**Good identification**						**46771622**
Clostridium clostridioforme		100.0	0.48	GLU 88			
46771632	**Very good identification**						**46771632**
Clostridium clostridioforme		100.0	0.59	GLU 88			
46771702	**Very good identification**						**46771702**
					SPO		
Clostridium clostridioforme		99.13	0.57	GLU 88	66		
Clostridium butyricum		0.60	0.07	XYL 91	99		
Bifidobacterium adolescentis		0.18	0.13		1		
46771712	**Very good identification**						**46771712**
					GRA		
Clostridium clostridioforme		99.93	0.68	GLU 88	50		
Bifidobacterium adolescentis		0.05	0.15	XYL 91	99		
Leptotrichia buccalis		0.02	-.11		1		
46771722	**Good identification**						**46771722**
					SPO		
Clostridium clostridioforme		92.32	0.57	GLU 88	66		
Parabacteroides distasonis		5.40	0.41	XYL 91	1		
Clostridium ramosum		1.37	0.19		92		
46771732	**Very good identification**						**46771732**
Clostridium clostridioforme		99.23	0.68	GLU 88			
Parabacteroides distasonis		0.69	0.37	XYL 91			
Bacteroides ovatus		0.06	0.03				
46773001	**Good identification**						**46773001**
Actinomyces naeslundii		100.0	0.43	ESL 12			
46773002	**Low discrimination**						**46773002**
Actinomyces israelii		52.17	0.37	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		22.22	0.40	XYL 92		
Bifidobacterium breve		18.65	0.24			
46773003	**Good identification**					**46773003**
Actinomyces naeslundii		100.0	0.43	ESL 12		
46773021	**Good identification**					**46773021**
Actinomyces naeslundii		100.0	0.43	ESL 12		
46773023	**Good identification**					**46773023**
Actinomyces naeslundii		100.0	0.43	ESL 12		
46773100	**Low discrimination**					**46773100**
Bifidobacterium breve		40.78	0.41	GLU 99		
Actinomyces israelii		31.07	0.45	BGL 91		
Actinomyces naeslundii		14.12	0.48			
46773101	**Good identification**					**46773101**
Actinomyces naeslundii		100.0	0.51	ESL 12		
46773102	**Low discrimination**					**46773102**
					SPO	
Bifidobacterium breve		75.54	0.59	GLU 99	1	
Actinomyces israelii		8.54	0.48		1	
Clostridium ramosum		6.51	0.30		92	
46773103	**Low discrimination**					**46773103**
Actinomyces naeslundii		82.04	0.51	ESL 12		
Bifidobacterium breve		16.13	0.24	GLU 99		
Actinomyces israelii		1.82	0.14			
46773110	**Low discrimination**					**46773110**
					SPO	
Bifidobacterium breve		82.06	0.41	GLU 99	1	
Actinomyces israelii		9.34	0.31	BGL 91	1	
Clostridium tertium		6.05	0.13		99	
46773112	**Low discrimination**					**46773112**
					GRA	
Bifidobacterium breve		82.94	0.59	GLU 99	99	
Leptotrichia buccalis		10.47	0.24		1	
Bifidobacterium adolescentis		3.06	0.33		99	
46773120	**Acceptable identification**					**46773120**
					SPO	
Clostridium ramosum		90.15	0.53	GLU 99	92	
Bifidobacterium breve		7.44	0.46		1	
Actinomyces naeslundii		1.45	0.48		1	
46773121	**Low discrimination**					**46773121**
					SPO	
Actinomyces naeslundii		68.79	0.51	ESL 12	1	
Clostridium ramosum		28.83	0.19	GLU 99	92	
Bifidobacterium breve		2.38	0.12		1	
46773122	**Low discrimination**					**46773122**
					SPO	
Clostridium ramosum		81.70	0.64	GLU 99	92	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		17.04	0.63		1	
Lactobacillus acidophilus		0.42	0.25		1	
46773130	**Low discrimination**					**46773130**
					SPO	
Bifidobacterium breve		79.55	0.46	GLU 99	1	
Clostridium tertium		9.89	0.21	BGL 91	99	
Clostridium ramosum		9.74	0.19		92	
46773132	**Good identification**					**46773132**
					SPO	
Bifidobacterium breve		91.17	0.63	GLU 99	1	
Clostridium ramosum		4.42	0.30		92	
Clostridium tertium		3.36	0.29		99	
46773140	**Low discrimination**					**46773140**
Bifidobacterium breve		73.67	0.41	GLU 99		
Actinomyces israelii		21.83	0.38	BGL 91		
Actinomyces naeslundii		4.50	0.35			
46773141	**Good identification**					**46773141**
Actinomyces naeslundii		100.0	0.38	ESL 12		
46773142	**Good identification**					**46773142**
Bifidobacterium breve		94.42	0.59	GLU 99		
Actinomyces israelii		4.15	0.41			
Actinomyces naeslundii		0.57	0.35			
46773150	**Acceptable identification**					**46773150**
					COC	
Bifidobacterium breve		94.00	0.41	GLU 99	1	
Actinomyces israelii		4.16	0.24	BGL 91	1	
Blautia producta		1.84	-.03		99	
46773152	**Very good identification**					**46773152**
Bifidobacterium breve		98.78	0.59	GLU 99		
Actinomyces israelii		0.65	0.27			
Bifidobacterium adolescentis		0.45	0.18			
46773160	**Low discrimination**					**46773160**
					SPO	
Bifidobacterium breve		85.66	0.46	GLU 99	1	
Clostridium ramosum		10.49	0.19	BGL 91	92	
Actinomyces naeslundii		2.94	0.35		1	
46773161	**Low discrimination**					**46773161**
Actinomyces naeslundii		83.62	0.38	ESL 12		
Bifidobacterium breve		16.38	0.12	GLU 99		
46773162	**Good identification**					**46773162**
					SPO	
Bifidobacterium breve		94.66	0.63	GLU 99	1	
Clostridium ramosum		4.59	0.30		92	
Actinomyces naeslundii		0.32	0.35		1	
46773170	**Good identification**					**46773170**
Bifidobacterium breve		100.0	0.46	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46773172	**Very good identification**					**46773172**
Bifidobacterium breve		100.0	0.63	GLU 99		
46773320	**Good identification**					**46773320**
					SPO	
Clostridium ramosum		99.94	0.56	GLU 99	92	
Bifidobacterium breve		0.06	0.12		1	
46773322	**Very good identification**					**46773322**
					SPO	
Clostridium ramosum		99.77	0.66	GLU 99	92	
Bifidobacterium breve		0.16	0.29		1	
Lactobacillus acidophilus		0.03	0.06		1	
46773402	**Low discrimination**					**46773402**
					SPO	
Actinomyces israelii		87.62	0.36	GLU 99	1	
Bifidobacterium adolescentis		7.73	0.12	XYL 92	1	
Clostridium butyricum		4.65	-.07		99	
46773502	**Low discrimination**					**46773502**
					SPO	
Bifidobacterium adolescentis		59.77	0.46	GLU 99	1	
Actinomyces israelii		27.38	0.46	XYL 87	1	
Clostridium butyricum		8.72	0.17		99	
46773512	**Low discrimination**					**46773512**
Bifidobacterium adolescentis		51.83	0.47	GLU 99		
Bifidobacterium dentium		40.77	0.27	XYL 87		
Actinomyces israelii		2.90	0.32			
46773542	**Low discrimination**					**46773542**
Actinomyces israelii		44.59	0.39	GLU 99		
Bifidobacterium adolescentis		30.93	0.30	XYL 92		
Bifidobacterium breve		12.52	0.24			
46774112	**Good identification**					**46774112**
					GRA	
Clostridium clostridioforme		97.67	0.42	GLU 88	50	
Bifidobacterium adolescentis		1.64	0.15	RHA 88	99	
Leptotrichia buccalis		0.69	-.11	ARA 94	1	
46774132	**Good identification**					**46774132**
Clostridium clostridioforme		98.85	0.42	GLU 88		
Parabacteroides distasonis		1.15	0.14	RHA 88		
46774202	**Good identification**					**46774202**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46774212	**Good identification**					**46774212**
Clostridium clostridioforme		100.0	0.48	GLU 88		
46774222	**Good identification**					**46774222**
Clostridium clostridioforme		100.0	0.36	GLU 88		
46774232	**Good identification**					**46774232**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.48	GLU 88			
46774302	**Good identification**						**46774302**
Clostridium clostridioforme		100.0	0.45	GLU 88			
46774312	**Very good identification**						**46774312**
Clostridium clostridioforme		100.0	0.57	GLU 88			
46774322	**Good identification**						**46774322**
Clostridium clostridioforme		96.13	0.45	GLU 88			
Parabacteroides distasonis		3.88	0.27	ARA 94			
46774332	**Very good identification**						**46774332**
Clostridium clostridioforme		99.53	0.57	GLU 88			
Parabacteroides distasonis		0.47	0.23	ARA 94			
46774402	**Good identification**						**46774402**
Clostridium clostridioforme		98.10	0.42	GLU 88			
Clostridium butyricum		1.90	0.01	RHA 88			
46774412	**Good identification**						**46774412**
Clostridium clostridioforme		100.0	0.53	GLU 88			
46774422	**Good identification**						**46774422**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46774432	**Good identification**						**46774432**
Clostridium clostridioforme		100.0	0.53	GLU 88			
46774502	**Low discrimination**						**46774502**
					SPO		
Clostridium clostridioforme		85.55	0.51	GLU 88	66		
Clostridium butyricum		11.90	0.25	RHA 88	99		
Bifidobacterium adolescentis		2.29	0.27		1		
46774512	**Very good identification**						**46774512**
					SPO		
Clostridium clostridioforme		99.06	0.62	GLU 88	66		
Bifidobacterium adolescentis		0.71	0.29	RHA 88	1		
Clostridium butyricum		0.16	0.03		99		
46774522	**Good identification**						**46774522**
					GRA	SPO	
Clostridium clostridioforme		98.43	0.51	GLU 88	50	66	
Clostridium butyricum		1.19	0.07	RHA 88	99	99	
Parabacteroides distasonis		0.38	0.15		1	1	
46774532	**Very good identification**						**46774532**
					GRA		
Clostridium clostridioforme		99.83	0.62	GLU 88	50		
Bifidobacterium dentium		0.12	-.03	RHA 88	99		
Parabacteroides distasonis		0.05	0.11		1		
46774602	**Very good identification**						**46774602**
Clostridium clostridioforme		100.0	0.57	GLU 88			
46774612	**Very good identification**						**46774612**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.68	GLU 88		
46774622	**Very good identification**					**46774622**
Clostridium clostridioforme		100.0	0.57	GLU 88		
46774632	**Very good identification**					**46774632**
Clostridium clostridioforme		100.0	0.68	GLU 88		
46774702	**Very good identification**					**46774702**
Clostridium clostridioforme		99.94	0.66	GLU 88		
Clostridium butyricum		0.06	-.01			
46774710	**Good identification**					**46774710**
Clostridium clostridioforme		100.0	0.43	GLU 88		
46774712	**Very good identification**					**46774712**
Clostridium clostridioforme		100.0	0.77	GLU 88		
46774713	**Good identification**					**46774713**
Clostridium clostridioforme		100.0	0.43	GLU 88		
46774722	**Very good identification**					**46774722**
Clostridium clostridioforme		99.72	0.66	GLU 88		
Parabacteroides distasonis		0.16	0.23			
Bacteroides ovatus		0.13	0.06			
46774730	**Good identification**					**46774730**
Clostridium clostridioforme		100.0	0.43	GLU 88		
46774732	**Very good identification**					**46774732**
Clostridium clostridioforme		99.96	0.77	GLU 88		
Parabacteroides distasonis		0.02	0.19			
Bacteroides ovatus		0.02	0.03			
46774733	**Good identification**					**46774733**
Clostridium clostridioforme		100.0	0.43	GLU 88		
46774752	**Good identification**					**46774752**
Clostridium clostridioforme		100.0	0.43	GLU 88		
46774772	**Good identification**					**46774772**
Clostridium clostridioforme		100.0	0.43	GLU 88		
46775002	**Low discrimination**					**46775002**
					SPO	
Actinomyces israelii		42.44	0.43	GLU 99	1	
Clostridium butyricum		29.37	0.19		99	
Clostridium clostridioforme		26.35	0.30		66	
46775012	**Acceptable identification**					**46775012**
Clostridium clostridioforme		91.55	0.41	GLU 88		
Actinomyces israelii		4.84	0.29	RHA 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		1.72	0.15	ARA 94			
46775032	**Good identification**						**46775032**
Clostridium clostridioforme		98.72	0.41	GLU 88			
Parabacteroides distasonis		1.28	0.14	RHA 88			
46775042	**Low discrimination**						**46775042**
					SPO		
Actinomyces israelii		86.60	0.36	GLU 99	1		
Clostridium butyricum		13.40	0.01		99		
46775100	**Acceptable identification**						**46775100**
					COC	SPO	
Clostridium butyricum		91.00	0.50	GLU 99	1	99	
Actinomyces israelii		5.68	0.50	ARA 90	1	1	
Atopobium parvulum		3.12	0.32		99	1	
46775102	**Low discrimination**						**46775102**
					SPO		
Clostridium butyricum		59.86	0.43	GLU 99	99		
Bifidobacterium adolescentis		15.51	0.47	ARA 90	1		
Actinomyces israelii		14.42	0.53		1		
46775112	**Low discrimination**						**46775112**
					GRA		
Clostridium clostridioforme		50.77	0.50	GLU 88	50		
Bifidobacterium adolescentis		28.20	0.49	RHA 88	99		
Leptotrichia buccalis		11.81	0.24	ARA 94	1		
46775122	**Low discrimination**						**46775122**
Parabacteroides distasonis		51.63	0.53	GLU 99			
Clostridium clostridioforme		15.88	0.39				
Prevotella oralis		12.55	0.30				
46775132	**Low discrimination**						**46775132**
					GRA		
Clostridium clostridioforme		68.15	0.50	GLU 88	50		
Parabacteroides distasonis		26.19	0.49	RHA 88	1		
Bifidobacterium breve		2.25	0.29	ARA 94	99		
46775142	**Low discrimination**						**46775142**
					SPO		
Actinomyces israelii		41.72	0.46	GLU 99	1		
Clostridium butyricum		38.73	0.24		99		
Bifidobacterium adolescentis		14.26	0.32		1		
46775202	**Good identification**						**46775202**
					SPO		
Clostridium clostridioforme		97.92	0.45	GLU 88	66		
Actinomyces israelii		1.62	0.23	ARA 94	1		
Clostridium butyricum		0.46	-.07		99		
46775212	**Good identification**						**46775212**
Clostridium clostridioforme		100.0	0.56	GLU 88			
46775222	**Acceptable identification**						**46775222**
Clostridium clostridioforme		94.80	0.45	GLU 88			
Parabacteroides distasonis		4.26	0.27	ARA 94			
Prevotella oralis		0.94	0.03				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
46775232	**Good identification**					**46775232**
Clostridium clostridioforme		99.47	0.56	GLU 88		
Parabacteroides distasonis		0.53	0.23	ARA 94		
46775302	**Acceptable identification**					**46775302**
					SPO	
Clostridium clostridioforme		93.27	0.54	GLU 88	66	
Clostridium butyricum		3.14	0.17	ARA 94	99	
Actinomyces israelii		1.84	0.34		1	
46775312	**Very good identification**					**46775312**
					GRA	
Clostridium clostridioforme		99.63	0.65	GLU 88	50	
Parabacteroides distasonis		0.16	0.23	ARA 94	1	
Bifidobacterium adolescentis		0.08	0.15		99	
46775322	**Low discrimination**					**46775322**
Parabacteroides distasonis		54.88	0.61	GLU 99		
Clostridium clostridioforme		41.26	0.54			
Prevotella oralis		1.91	0.24			
46775332	**Low discrimination**					**46775332**
Clostridium clostridioforme		86.19	0.65	GLU 88		
Parabacteroides distasonis		13.55	0.57	ARA 94		
Bacteroides ovatus		0.26	0.11			
46775400	**Good identification**					**46775400**
					SPO	
Clostridium butyricum		96.14	0.42	GLU 99	99	
Actinomyces israelii		3.28	0.38	LAC 96	1	
Clostridium clostridioforme		0.59	0.16		66	
46775402	**Low discrimination**					**46775402**
					SPO	
Clostridium clostridioforme		57.00	0.50	GLU 88	66	
Clostridium butyricum		36.50	0.35	RHA 88	99	
Actinomyces israelii		4.80	0.41		1	
46775412	**Very good identification**					**46775412**
					SPO	
Clostridium clostridioforme		98.14	0.61	GLU 88	66	
Bifidobacterium adolescentis		0.79	0.29	RHA 88	1	
Clostridium butyricum		0.73	0.13		99	
46775422	**Acceptable identification**					**46775422**
					SPO	
Clostridium clostridioforme		93.87	0.50	GLU 88	66	
Clostridium butyricum		5.23	0.17	RHA 88	99	
Actinomyces israelii		0.50	0.21		1	
46775432	**Very good identification**					**46775432**
					SPO	
Clostridium clostridioforme		99.75	0.61	GLU 88	66	
Bifidobacterium dentium		0.14	-.03	RHA 88	1	
Clostridium butyricum		0.06	-.05		99	
46775500	**Very good identification**					**46775500**
					SPO	
Clostridium butyricum		99.20	0.66	GLU 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.56	0.48		1		
Clostridium clostridioforme		0.09	0.25		66		
46775502	**Low discrimination**						**46775502**
					SPO		
Clostridium butyricum		69.78	0.59	GLU 99	99		
Clostridium clostridioforme		15.20	0.59		66		
Bifidobacterium adolescentis		13.44	0.61		1		
46775510	**Low discrimination**						**46775510**
					SPO		
Clostridium butyricum		88.44	0.44	GLU 99	99		
Clostridium clostridioforme		6.55	0.36	MLZ 5	66		
Bifidobacterium adolescentis		1.55	0.29		1		
46775512	**Low discrimination**						**46775512**
					SPO		
Clostridium clostridioforme		75.95	0.70	GLU 88	66		
Bifidobacterium adolescentis		18.02	0.63	RHA 88	1		
Clostridium butyricum		4.03	0.37		99		
46775513	**Low discrimination**						**46775513**
					SPO		
Clostridium clostridioforme		76.16	0.36	GLU 88	66		
Bifidobacterium adolescentis		18.07	0.29	RHA 88	1		
Clostridium butyricum		4.04	0.03	URE 1	99		
46775520	**Good identification**						**46775520**
					SPO		
Clostridium butyricum		98.21	0.48	GLU 99	99		
Clostridium clostridioforme		0.97	0.25	NAG 8	66		
Actinomyces israelii		0.41	0.28		1		
46775522	**Low discrimination**						**46775522**
					GRA	SPO	
Clostridium clostridioforme		63.86	0.59	GLU 88	50	66	
Clostridium butyricum		25.49	0.41	RHA 88	99	99	
Parabacteroides distasonis		8.12	0.49		1	1	
46775532	**Good identification**						**46775532**
					GRA		
Clostridium clostridioforme		93.85	0.70	GLU 88	50		
Bifidobacterium dentium		3.81	0.32	RHA 88	99		
Parabacteroides distasonis		1.41	0.45		1		
46775533	**Acceptable identification**						**46775533**
					GRA		
Clostridium clostridioforme		94.73	0.36	GLU 88	50		
Bifidobacterium dentium		3.84	-.03	RHA 88	99		
Parabacteroides distasonis		1.42	0.11	URE 1	1		
46775540	**Acceptable identification**						**46775540**
					COC	SPO	
Clostridium butyricum		93.60	0.48	GLU 99	1	99	
Blautia producta		3.12	0.24	SOR 8	99	1	
Actinomyces israelii		2.38	0.41		1	1	
46775542	**Low discrimination**						**46775542**
					SPO		
Clostridium butyricum		70.97	0.41	GLU 99	99		
Bifidobacterium adolescentis		19.43	0.46	SOR 8	1		
Actinomyces israelii		6.95	0.44		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46775552	**Low discrimination**						**46775552**
					SPO		
Bifidobacterium adolescentis		61.03	0.47	GLU 99	1		
Clostridium clostridioforme		21.02	0.36	SOR 11	66		
Clostridium butyricum		9.60	0.19		99		
46775572	**Low discrimination**						**46775572**
					SPO		
Clostridium clostridioforme		86.80	0.36	GLU 88	66		
Bifidobacterium dentium		3.52	-.03	RHA 88	1		
Clostridium butyricum		3.45	0.01	SOR 1	99		
46775602	**Very good identification**						**46775602**
					SPO		
Clostridium clostridioforme		99.65	0.65	GLU 88	66		
Clostridium butyricum		0.27	0.09		99		
Actinomyces israelii		0.09	0.22		1		
46775610	**Good identification**						**46775610**
Clostridium clostridioforme		99.21	0.42	GLU 88			
Clostridium butyricum		0.79	-.06	BGL 99			
46775612	**Very good identification**						**46775612**
Clostridium clostridioforme		100.0	0.76	GLU 88			
46775613	**Good identification**						**46775613**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46775622	**Very good identification**						**46775622**
Clostridium clostridioforme		99.63	0.65	GLU 88			
Parabacteroides distasonis		0.18	0.23				
Bacteroides ovatus		0.14	0.06				
46775630	**Good identification**						**46775630**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46775632	**Very good identification**						**46775632**
Clostridium clostridioforme		99.96	0.76	GLU 88			
Parabacteroides distasonis		0.02	0.19				
Bacteroides ovatus		0.02	0.03				
46775633	**Good identification**						**46775633**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46775652	**Good identification**						**46775652**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46775672	**Good identification**						**46775672**
Clostridium clostridioforme		100.0	0.42	GLU 88			
46775700	**Low discrimination**						**46775700**
					GRA	SPO	
Clostridium butyricum		79.25	0.40	GLU 99	99	99	
Clostridium clostridioforme		16.08	0.40	RHA 3	50	66	
Mitsuokella multacida		3.58	0.25		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46775702	**Very good identification**						**46775702**
					SPO		
Clostridium clostridioforme		97.77	0.74	GLU 88	66		
Clostridium butyricum		1.89	0.33		99		
Bifidobacterium adolescentis		0.12	0.27		1		
46775703	**Good identification**						**46775703**
Clostridium clostridioforme		98.10	0.40	GLU 88			
Clostridium butyricum		1.90	-.01	URE 1			
46775710	**Acceptable identification**						**46775710**
					GRA	SPO	
Clostridium clostridioforme		92.72	0.51	GLU 88	50	66	
Clostridium butyricum		5.28	0.18	BGL 99	99	99	
Mitsuokella multacida		1.51	0.17		1	1	
46775712	**Very good identification**						**46775712**
					SPO		
Clostridium clostridioforme		99.92	0.85	GLU 88	66		
Bifidobacterium adolescentis		0.03	0.29		1		
Clostridium butyricum		0.02	0.11		99		
46775713	**Good identification**						**46775713**
Clostridium clostridioforme		100.0	0.51	GLU 88			
46775720	**Low discrimination**						**46775720**
					GRA	SPO	
Clostridium clostridioforme		59.55	0.40	GLU 88	50	66	
Clostridium butyricum		25.52	0.22	BGL 99	99	99	
Bacteroides vulgatus		6.04	0.10		1	1	
46775722	**Good identification**						**46775722**
Clostridium clostridioforme		90.64	0.74	GLU 88			
Parabacteroides distasonis		4.72	0.58				
Bacteroides ovatus		3.78	0.40				
46775723	**Acceptable identification**						**46775723**
Clostridium clostridioforme		91.43	0.40	GLU 88			
Parabacteroides distasonis		4.76	0.23	URE 1			
Bacteroides ovatus		3.81	0.06				
46775730	**Good identification**						**46775730**
Clostridium clostridioforme		97.92	0.51	GLU 88			
Parabacteroides distasonis		0.60	0.19	BGL 99			
Bacteroides ovatus		0.60	0.03				
46775732	**Very good identification**						**46775732**
Clostridium clostridioforme		98.72	0.85	GLU 88			
Parabacteroides distasonis		0.61	0.53				
Bacteroides ovatus		0.60	0.37				
46775733	**Good identification**						**46775733**
Clostridium clostridioforme		98.79	0.51	GLU 88			
Parabacteroides distasonis		0.61	0.19	URE 1			
Bacteroides ovatus		0.60	0.03				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46775742	**Low discrimination**						**46775742**
					COC	SPO	
Clostridium clostridioforme		77.09	0.40	GLU 88	1	66	
Clostridium butyricum		12.85	0.15	SOR 1	1	99	
Blautia producta		3.95	0.07		99	1	
46775752	**Good identification**						**46775752**
					COC		
Clostridium clostridioforme		96.02	0.51	GLU 88	1		
Blautia producta		3.24	0.15	SOR 1	99		
Bifidobacterium adolescentis		0.38	0.13		1		
46775762	**Low discrimination**						**46775762**
Clostridium clostridioforme		80.75	0.40	GLU 88			
Bacteroides ovatus		13.88	0.16	SOR 1			
Parabacteroides distasonis		4.20	0.23				
46775772	**Good identification**						**46775772**
Clostridium clostridioforme		96.61	0.51	GLU 88			
Bacteroides ovatus		2.43	0.13	SOR 1			
Parabacteroides distasonis		0.59	0.19				
46776712	**Good identification**						**46776712**
Clostridium clostridioforme		99.08	0.43	GLU 88			
Bifidobacterium dentium		0.93	-.07	MAN 1			
46776732	**Good identification**						**46776732**
					GRA		
Clostridium clostridioforme		97.78	0.43	GLU 88	50		
Bifidobacterium dentium		1.62	-.03	MAN 1	99		
Bacteroides ovatus		0.60	-.05		1		
46777000	**Good identification**						**46777000**
					SPO		
Actinomyces israelii		95.03	0.53	GLU 99	1		
Clostridium butyricum		4.97	0.10		99		
46777002	**Good identification**						**46777002**
					SPO		
Actinomyces israelii		98.05	0.56	GLU 99	1		
Clostridium butyricum		1.33	0.03		99		
Bifidobacterium adolescentis		0.62	0.12		1		
46777010	**Good identification**						**46777010**
Actinomyces israelii		100.0	0.38	GLU 99			
46777012	**Low discrimination**						**46777012**
Actinomyces israelii		88.86	0.41	GLU 99			
Bifidobacterium adolescentis		4.56	0.13	MLZ 13			
Bifidobacterium dentium		3.59	-.07				
46777022	**Good identification**						**46777022**
Actinomyces israelii		100.0	0.35	GLU 99			
46777040	**Good identification**						**46777040**
Actinomyces israelii		100.0	0.46	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
46777042	**Good identification**						**46777042**
Actinomyces israelii		100.0	0.49	GLU 99			
46777100	**Low discrimination**						**46777100**
					SPO		
Actinomyces israelii		75.59	0.63	GLU 99	1		
Clostridium butyricum		23.74	0.33		99		
Clostridium tertium		0.49	0.11		99		
46777102	**Low discrimination**						**46777102**
					SPO		
Actinomyces israelii		79.37	0.66	GLU 99	1		
Bifidobacterium adolescentis		12.33	0.46		1		
Clostridium butyricum		6.46	0.26		99		
46777110	**Low discrimination**						**46777110**
					SPO		
Actinomyces israelii		72.96	0.49	GLU 99	1		
Bifidobacterium dentium		10.81	0.10	MLZ 13	1		
Clostridium butyricum		8.07	0.11		99		
46777112	**Low discrimination**						**46777112**
Bifidobacterium adolescentis		35.88	0.47	GLU 99			
Actinomyces israelii		28.25	0.52	ARA 87			
Bifidobacterium dentium		28.22	0.27				
46777120	**Low discrimination**						**46777120**
					SPO		
Actinomyces israelii		46.38	0.42	GLU 99	1		
Clostridium butyricum		19.84	0.15	NAG 6	99		
Clostridium ramosum		18.05	0.19		92		
46777140	**Low discrimination**						**46777140**
					COC	SPO	
Actinomyces israelii		79.40	0.56	GLU 99	1	1	
Blautia producta		15.03	0.24		99	1	
Clostridium butyricum		5.57	0.15		1	99	
46777142	**Good identification**						**46777142**
Actinomyces israelii		90.78	0.59	GLU 99			
Bifidobacterium adolescentis		4.48	0.30				
Bifidobacterium breve		1.81	0.24				
46777152	**Low discrimination**						**46777152**
					COC		
Actinomyces israelii		54.21	0.45	GLU 99	1		
Bifidobacterium adolescentis		21.89	0.32	MLZ 13	1		
Blautia producta		15.26	0.15		99		
46777160	**Low discrimination**						**46777160**
					COC	SPO	
Actinomyces israelii		66.33	0.35	GLU 99	1	1	
Blautia producta		21.85	0.07	NAG 6	99	1	
Clostridium butyricum		6.34	-.03		1	99	
46777162	**Low discrimination**						**46777162**
Actinomyces israelii		64.24	0.38	GLU 99			
Bifidobacterium breve		35.76	0.29	NAG 6			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
46777202	**Acceptable identification**						**46777202**
Actinomyces israelii		90.27	0.36	GLU 99			
Clostridium clostridioforme		9.73	0.10	RHA 7			
46777300	**Low discrimination**						**46777300**
					SPO		
Actinomyces israelii		88.57	0.44	GLU 99	1		
Clostridium butyricum		11.43	0.07	RHA 7	99		
46777302	**Low discrimination**						**46777302**
					SPO		
Actinomyces israelii		86.48	0.47	GLU 99	1		
Clostridium clostridioforme		7.80	0.19	RHA 7	66		
Clostridium butyricum		2.89	0.00		99		
46777342	**Low discrimination**						**46777342**
					COC		
Actinomyces israelii		84.30	0.40	GLU 99	1		
Blautia producta		15.70	0.07	RHA 7	99		
46777400	**Low discrimination**						**46777400**
					SPO		
Actinomyces israelii		63.47	0.51	GLU 99	1		
Clostridium butyricum		36.53	0.26		99		
46777402	**Low discrimination**						**46777402**
					SPO		
Actinomyces israelii		81.36	0.54	GLU 99	1		
Clostridium butyricum		12.14	0.19		99		
Bifidobacterium adolescentis		4.18	0.26		1		
46777410	**Low discrimination**						**46777410**
					SPO		
Actinomyces israelii		49.99	0.37	GLU 99	1		
Bifidobacterium dentium		36.23	0.10	MLZ 13	1		
Clostridium butyricum		10.14	0.04		99		
46777440	**Low discrimination**						**46777440**
					COC	SPO	
Actinomyces israelii		48.69	0.44	GLU 99	1	1	
Blautia producta		45.05	0.24		99	1	
Clostridium butyricum		6.27	0.08		1	99	
46777442	**Low discrimination**						**46777442**
					COC	SPO	
Actinomyces israelii		89.43	0.47	GLU 99	1	1	
Blautia producta		6.13	0.07		99	1	
Clostridium butyricum		2.98	0.01		1	99	
46777500	**Low discrimination**						**46777500**
					COC	SPO	
Clostridium butyricum		74.36	0.50	GLU 99	1	99	
Actinomyces israelii		21.53	0.61	MAN 10	1	1	
Blautia producta		1.94	0.24		99	1	
46777502	**Low discrimination**						**46777502**
					SPO		
Bifidobacterium adolescentis		37.62	0.60	GLU 99	1		
Actinomyces israelii		29.61	0.64		1		
Clostridium butyricum		26.52	0.43		99		
46777510	**Low discrimination**						**46777510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Bifidobacterium dentium		80.66	0.44	GLU 99	1	1	
Blautia producta		8.12	0.32	BGL 91	99	1	
Clostridium butyricum		5.47	0.28		1	99	
46777512	**Good identification**						**46777512**
Bifidobacterium dentium		90.70	0.62	GLU 99			
Bifidobacterium adolescentis		7.80	0.61				
Actinomyces israelii		0.75	0.50				
46777530	**Good identification**						**46777530**
					COC	SPO	
Bifidobacterium dentium		98.76	0.49	GLU 99	1	1	
Blautia producta		0.62	0.15	BGL 91	99	1	
Clostridium butyricum		0.33	0.10		1	99	
46777532	**Very good identification**						**46777532**
Bifidobacterium dentium		99.63	0.66	GLU 99			
Clostridium clostridioforme		0.25	0.36				
Bifidobacterium adolescentis		0.05	0.27				
46777540	**Good identification**						**46777540**
					COC	SPO	
Blautia producta		92.27	0.58	GLU 99	99	1	
Actinomyces israelii		4.03	0.54		1	1	
Clostridium butyricum		3.11	0.31		1	99	
46777542	**Low discrimination**						**46777542**
					COC		
Blautia producta		51.16	0.41	GLU 99	99		
Actinomyces israelii		30.16	0.57	BGL 10	1		
Bifidobacterium adolescentis		12.18	0.44		1		
46777550	**Very good identification**						**46777550**
					COC		
Blautia producta		99.55	0.66	GLU 99	99		
Actinomyces israelii		0.22	0.40		1		
Bifidobacterium dentium		0.10	0.10		1		
46777552	**Low discrimination**						**46777552**
					COC		
Blautia producta		81.88	0.50	GLU 99	99		
Bifidobacterium adolescentis		7.94	0.46	BGL 10	1		
Bifidobacterium dentium		7.55	0.27		1		
46777560	**Acceptable identification**						**46777560**
					COC	SPO	
Blautia producta		93.87	0.41	GLU 99	99	1	
Clostridium butyricum		2.48	0.13	NAG 10	1	99	
Actinomyces israelii		2.36	0.34		1	1	
46777570	**Good identification**						**46777570**
					COC	GRA	
Blautia producta		98.14	0.50	GLU 99	99	99	
Bifidobacterium dentium		1.59	0.15	NAG 10	1	99	
Mitsuokella multacida		0.15	0.09		1	1	
46777612	**Good identification**						**46777612**
Clostridium clostridioforme		97.44	0.42	GLU 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.55	0.21	MAN 1			
Bifidobacterium dentium		1.02	-.07				
46777632	**Good identification**						**46777632**
					GRA		
Clostridium clostridioforme		97.53	0.42	GLU 88	50		
Bifidobacterium dentium		1.81	-.03	MAN 1	99		
Bacteroides ovatus		0.66	-.05		1		
46777702	**Low discrimination**						**46777702**
					SPO		
Clostridium clostridioforme		49.33	0.40	GLU 88	66		
Actinomyces israelii		28.55	0.45	MAN 1	1		
Clostridium butyricum		10.50	0.17		99		
46777712	**Low discrimination**						**46777712**
					COC		
Clostridium clostridioforme		71.64	0.51	GLU 88	1		
Bifidobacterium dentium		22.06	0.27	MAN 1	1		
Blautia producta		2.42	0.15		99		
46777732	**Low discrimination**						**46777732**
					GRA		
Clostridium clostridioforme		56.86	0.51	GLU 88	50		
Bifidobacterium dentium		31.13	0.32	MAN 1	99		
Bacteroides ovatus		11.44	0.29		1		
46777740	**Very good identification**						**46777740**
					COC	GRA	
Blautia producta		98.45	0.58	GLU 99	99	99	
Mitsuokella multacida		1.12	0.32		1	1	
Actinomyces israelii		0.32	0.35		1	99	
46777742	**Acceptable identification**						**46777742**
					COC	GRA	
Blautia producta		94.00	0.41	GLU 99	99	99	
Actinomyces israelii		4.17	0.38	BGL 10	1	99	
Mitsuokella multacida		1.07	0.15		1	1	
46777750	**Very good identification**						**46777750**
					COC	GRA	
Blautia producta		99.86	0.66	GLU 99	99	99	
Mitsuokella multacida		0.13	0.24		1	1	
Actinomyces israelii		0.02	0.21		1	99	
46777752	**Good identification**						**46777752**
					COC		
Blautia producta		99.17	0.50	GLU 99	99		
Clostridium clostridioforme		0.30	0.17	BGL 10	1		
Actinomyces israelii		0.22	0.24		1		
46777760	**Good identification**						**46777760**
					COC	GRA	
Blautia producta		97.33	0.41	GLU 99	99	99	
Mitsuokella multacida		2.49	0.21	NAG 10	1	1	
Actinomyces israelii		0.18	0.15		1	99	
46777770	**Good identification**						**46777770**
					COC	GRA	
Blautia producta		99.72	0.50	GLU 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
47000000	**Low discrimination**						**47000000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium botulinum biovar A		58.34	0.62	SUC 7	1	99	
Atopobium minutum		23.32	0.62	MNS 7	1	1	
Gemella morbillorum		10.22	0.52		99	1	
47000002	**Good identification**						**47000002**
					SPO		
Atopobium minutum		92.99	0.59	SUC 5	1		
Clostridium botulinum biovar A		3.53	0.27		99		
Clostridium sporogenes		1.54	0.19		99		
47000020	**Low discrimination**						**47000020**
					COC	SPO	
Atopobium minutum		77.54	0.54	SUC 5	1	1	
Gemella morbillorum		11.33	0.35		99	1	
Clostridium botulinum biovar A		5.88	0.27		1	99	
47000022	**Good identification**						**47000022**
					COC	GRA	
Atopobium minutum		99.29	0.51	SUC 5	1	99	
Pseudoflavonifractor capillosus		0.49	-.03		1	1	
Gemella morbillorum		0.22	0.01		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47000040	**Good identification**						**47000040**
					SPO		
Clostridium botulinum biovar A		95.92	0.53	SUC 7	99		
Atopobium minutum		1.30	0.28	MNS 7	1		
Clostridium bifermentans		1.23	0.20		99		
47000100	**Low discrimination**						**47000100**
					GRA	SPO	
Fusobacterium mortiferum		76.23	0.53	FRU 99	1	1	
Clostridium botulinum biovar A		14.26	0.27		99	99	
Atopobium minutum		5.70	0.28		99	1	
47000102	**Low discrimination**						**47000102**
					GRA		
Fusobacterium mortiferum		69.03	0.37	FRU 99	1		
Atopobium minutum		30.97	0.25	BGL 10	99		
47000400	**Low discrimination**						**47000400**
					COC	SPO	
Gemella morbillorum		55.99	0.35	ESL 1	99	1	
Clostridium botulinum biovar A		29.06	0.27	ARA 10	1	99	
Atopobium minutum		11.61	0.28		1	1	
47001000	**Low discrimination**						**47001000**
					COC	GRA	
Atopobium minutum		43.39	0.54	SUC 5	1	99	
Gemella morbillorum		30.72	0.47		99	99	
Fusobacterium mortiferum		11.30	0.50		1	1	
47001002	**Low discrimination**						**47001002**
					GRA		
Atopobium minutum		89.11	0.51	SUC 5	99		
Eubacterium tortuosum		5.15	0.19		99		
Fusobacterium mortiferum		3.87	0.33		1		
47001020	**Low discrimination**						**47001020**
					COC		
Atopobium minutum		71.86	0.46	SUC 5	1		
Gemella morbillorum		16.96	0.31		99		
Eubacterium tortuosum		8.31	0.19		1		
47001022	**Good identification**						**47001022**
Atopobium minutum		97.03	0.43	SUC 5			
Eubacterium tortuosum		2.97	0.06				
47001100	**Low discrimination**						**47001100**
					SPO	GRA	
Fusobacterium mortiferum		84.20	0.56	FRU 99	1	1	
Eubacterium saburreum		8.28	0.37		1	85	
Clostridium chauvoei		4.28	0.22		99	99	
47001101	**Very good identification**						**47001101**
					GRA	SPO	
Clostridium chauvoei		99.80	0.57	MLT 99	99	99	
Fusobacterium mortiferum		0.20	0.22		1	1	
47001102	**Low discrimination**						**47001102**
					GRA		
Fusobacterium mortiferum		74.82	0.40	FRU 99	1		
Eubacterium saburreum		11.68	0.24	BGL 10	85		
Atopobium minutum		7.46	0.17		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47001103	**Good identification**							**47001103**
Clostridium chauvoei		100.0	0.44	MLT 99				
47001110	**Low discrimination**							**47001110**
Eubacterium saburreum		88.84	0.36	IND 99				
Fusobacterium mortiferum		11.16	0.22					
47001121	**Good identification**							**47001121**
					GRA	SPO		
Clostridium chauvoei		99.80	0.48	MLT 99	99	99		
Prevotella melaninogenica		0.20	-.04		1	1		
47001123	**Good identification**							**47001123**
Clostridium chauvoei		100.0	0.36	MLT 99				
47004000	**Low discrimination**							**47004000**
					COC	SPO		
Gemella morbillorum		53.67	0.35	ESL 1	99	1		
Clostridium botulinum biovar A		27.85	0.27	XYL 10	1	99		
Atopobium minutum		11.13	0.28		1	1		
47005100	**Low discrimination**							**47005100**
					GRA			
Fusobacterium mortiferum		56.01	0.37	FRU 99	1			
Eubacterium saburreum		39.39	0.33	XYL 7	85			
Actinomyces meyeri		4.61	0.09		99			
47010000	**Low discrimination**							**47010000**
					GRA			
Fusobacterium mortiferum		62.19	0.81		1			
Eubacterium limosum		14.40	0.52		99			
Atopobium minutum		10.02	0.62		99			
47010001	**Low discrimination**							**47010001**
					GRA			
Fusobacterium mortiferum		62.75	0.47	URE 1	1			
Eubacterium limosum		14.53	0.18		99			
Atopobium minutum		10.11	0.28		99			
47010002	**Low discrimination**							**47010002**
					GRA	CAT		
Fusobacterium mortiferum		46.98	0.65	BGL 10	1	5		
Atopobium minutum		45.40	0.59		99	1		
Propionibacterium granulosum		5.73	0.46		99	90		
47010010	**Low discrimination**							**47010010**
					GRA	SPO	CAT	
Clostridium difficile		84.61	0.50	SUC 1	99	99	1	
Fusobacterium mortiferum		6.86	0.47	MAN 90	1	1	5	
Propionibacterium granulosum		5.01	0.41		99	1	90	
47010020	**Low discrimination**							**47010020**
					COC	CAT		
Atopobium minutum		58.31	0.54	SUC 5	1	1		
Propionibacterium granulosum		14.71	0.46		1	90		
Gemella morbillorum		11.28	0.37		99	1		
47010022	**Low discrimination**							**47010022**
					GRA	CAT		
Atopobium minutum		88.58	0.51	SUC 5	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		8.38	0.35		99	90	
Fusobacterium mortiferum		2.78	0.30		1	5	
47010040	**Low discrimination**						**47010040**
					GRA	SPO	
Fusobacterium mortiferum		32.02	0.47	SOR 1	1	1	
Clostridium botulinum biovar A		26.39	0.32		99	99	
Clostridium botulinum biovar B		12.13	0.24		99	99	
47010100	**Very good identification**						**47010100**
					GRA		
Fusobacterium mortiferum		99.68	0.87		1		
Eubacterium limosum		0.10	0.18		99		
Atopobium minutum		0.07	0.28		99		
47010101	**Low discrimination**						**47010101**
					GRA	SPO	
Fusobacterium mortiferum		84.90	0.53	URE 1	1	1	
Clostridium chauvoei		15.10	0.29		99	99	
47010102	**Very good identification**						**47010102**
					GRA	CAT	
Fusobacterium mortiferum		99.53	0.71	BGL 10	1	5	
Atopobium minutum		0.42	0.25		99	1	
Propionibacterium granulosum		0.05	0.11		99	90	
47010103	**Low discrimination**						**47010103**
					GRA	SPO	
Fusobacterium mortiferum		77.98	0.37	URE 1	1	1	
Clostridium chauvoei		22.02	0.16	BGL 10	99	99	
47010110	**Acceptable identification**						**47010110**
					SPO	GRA	
Fusobacterium mortiferum		94.60	0.53	MLZ 1	1	1	
Clostridium difficile		5.06	0.15		99	99	
Eubacterium saburreum		0.35	0.10		1	85	
47010112	**Good identification**						**47010112**
Fusobacterium mortiferum		100.0	0.37	MLZ 1			
47010120	**Low discrimination**						**47010120**
					GRA		
Fusobacterium mortiferum		89.35	0.53	NAG 1	1		
Prevotella melaninogenica		4.04	0.12		1		
Clostridium perfringens		2.90	0.17		99		
47010122	**Low discrimination**						**47010122**
					GRA	CAT	
Fusobacterium mortiferum		59.27	0.37	NAG 1	1	5	
Capnocytophaga ochracea		14.48	0.12	BGL 10	1	90	
Atopobium minutum		8.18	0.17		99	1	
47010140	**Good identification**						**47010140**
					GRA	SPO	
Fusobacterium mortiferum		99.65	0.53	SOR 1	1	1	
Clostridium botulinum biovar A		0.36	-.02		99	99	
47010142	**Good identification**						**47010142**
Fusobacterium mortiferum		100.0	0.37	SOR 1			
47010200	**Low discrimination**						**47010200**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		58.64	0.47	RHA 1	1	5		
Eubacterium limosum		13.58	0.18		99	1		
Propionibacterium granulosum		9.73	0.30		99	90		
47010300	**Good identification**							**47010300**
Fusobacterium mortiferum		100.0	0.53	RHA 1				
47010302	**Good identification**							**47010302**
Fusobacterium mortiferum		100.0	0.37	RHA 1				
47010400	**Low discrimination**							**47010400**
					COC	GRA		
Gemella morbillorum		40.53	0.37	ESL 1	99	99		
Fusobacterium mortiferum		39.41	0.47	ARA 10	1	1		
Eubacterium limosum		9.13	0.18		1	99		
47010500	**Good identification**							**47010500**
					COC	GRA		
Fusobacterium mortiferum		98.08	0.53	ARA 1	1	1		
Actinomyces meyeri		1.49	0.13		1	99		
Gemella morbillorum		0.44	0.03		99	99		
47010502	**Good identification**							**47010502**
Fusobacterium mortiferum		100.0	0.37	ARA 1				
47011000	**Low discrimination**							**47011000**
					GRA	CAT		
Fusobacterium mortiferum		89.59	0.84		1	5		
Propionibacterium granulosum		3.24	0.56		99	90		
Atopobium minutum		3.21	0.54		99	1		
47011001	**Low discrimination**							**47011001**
					GRA	SPO	CAT	
Fusobacterium mortiferum		72.36	0.50	URE 1	1	1	5	
Clostridium chauvoei		20.01	0.29		99	99	1	
Propionibacterium granulosum		2.61	0.22		99	1	90	
47011002	**Low discrimination**							**47011002**
					GRA	CAT		
Fusobacterium mortiferum		74.33	0.68	BGL 10	1	5		
Atopobium minutum		15.96	0.51		99	1		
Propionibacterium granulosum		6.04	0.46		99	90		
47011010	**Low discrimination**							**47011010**
					CAT	GRA		
Fusobacterium mortiferum		51.68	0.50	MLZ 1	5	1		
Propionibacterium granulosum		25.19	0.41		90	99		
Eubacterium saburreum		12.47	0.37		1	85		
47011020	**Low discrimination**							**47011020**
					GRA	CAT		
Atopobium minutum		26.74	0.46	SUC 5	99	1		
Fusobacterium mortiferum		22.64	0.50		1	5		
Propionibacterium granulosum		20.24	0.46		99	90		
47011022	**Low discrimination**							**47011022**
					CAT			
Atopobium minutum		45.29	0.43	SUC 5	1			
Clostridium perfringens		18.47	0.31		1			
Propionibacterium granulosum		12.85	0.35		90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47011040	**Acceptable identification**						**47011040**
					GRA	CAT	
Fusobacterium mortiferum		90.01	0.50	SOR 1	1	5	
Propionibacterium granulosum		3.25	0.22		99	90	
Atopobium minutum		3.22	0.20		99	1	
47011100	**Very good identification**						**47011100**
					GRA		
Fusobacterium mortiferum		99.09	0.90		1		
Eubacterium saburreum		0.30	0.45		85		
Lactobacillus fermentum		0.21	0.46		99		
47011101	**Good identification**						**47011101**
					GRA	SPO	
Clostridium chauvoei		92.10	0.63	MLT 99	99	99	
Fusobacterium mortiferum		7.85	0.56		1	1	
Actinomyces naeslundii		0.03	0.25		99	1	
47011102	**Good identification**						**47011102**
					GRA	CAT	
Fusobacterium mortiferum		94.80	0.74	BGL 10	1	5	
Lactobacillus fermentum		1.80	0.46		99	1	
Capnocytophaga ochracea		1.72	0.30		1	90	
47011103	**Acceptable identification**						**47011103**
					GRA	SPO	
Clostridium chauvoei		94.76	0.50	MLT 99	99	99	
Fusobacterium mortiferum		5.09	0.40		1	1	
Actinomyces naeslundii		0.15	0.25		99	1	
47011110	**Low discrimination**						**47011110**
					CAT		
Fusobacterium mortiferum		80.40	0.56	MLZ 1	5		
Eubacterium saburreum		19.40	0.44		1		
Capnocytophaga ochracea		0.20	-.03		90		
47011112	**Low discrimination**						**47011112**
					CAT		
Fusobacterium mortiferum		63.20	0.40	MLZ 1	5		
Eubacterium saburreum		24.22	0.31	BGL 10	1		
Capnocytophaga ochracea		12.58	0.14		90		
47011120	**Low discrimination**						**47011120**
					GRA		
Prevotella melaninogenica		44.98	0.47	MLT 99	1		
Clostridium perfringens		32.34	0.51	RAF 90	99		
Fusobacterium mortiferum		15.08	0.56		1		
47011121	**Good identification**						**47011121**
					GRA	SPO	
Clostridium chauvoei		98.36	0.55	MLT 99	99	99	
Prevotella melaninogenica		0.76	0.12		1	1	
Clostridium perfringens		0.55	0.17		99	84	
47011122	**Low discrimination**						**47011122**
					GRA	CAT	
Capnocytophaga ochracea		44.55	0.47	MLT 99	1	90	
Clostridium perfringens		25.10	0.45	RAF 90	99	1	
Prevotella melaninogenica		14.13	0.34		1	1	
47011123	**Acceptable identification**						**47011123**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		94.41	0.42	MLT 99	99	1	
Capnocytophaga ochracea		2.48	0.12		1	90	
Clostridium perfringens		1.39	0.11		99	1	
47011140	**Good identification**						**47011140**
					GRA		
Fusobacterium mortiferum		98.23	0.56	SOR 1	1		
Clostridium perfringens		1.24	0.13		99		
Eubacterium saburreum		0.29	0.11		85		
47011142	**Low discrimination**						**47011142**
					GRA	CAT	
Fusobacterium mortiferum		79.84	0.40	SOR 1	1	5	
Capnocytophaga ochracea		15.89	0.14	BGL 10	1	90	
Clostridium perfringens		4.27	0.07		99	1	
47011160	**Low discrimination**						**47011160**
					GRA	CAT	
Clostridium perfringens		86.76	0.39	MLT 99	99	1	
Prevotella melaninogenica		6.40	0.12		1	1	
Capnocytophaga ochracea		4.70	0.14		1	90	
47011200	**Low discrimination**						**47011200**
					GRA	CAT	
Fusobacterium mortiferum		84.75	0.50	RHA 1	1	5	
Propionibacterium granulosum		9.37	0.30		99	90	
Atopobium minutum		3.03	0.20		99	1	
47011300	**Good identification**						**47011300**
Fusobacterium mortiferum		99.70	0.56	RHA 1			
Eubacterium saburreum		0.30	0.11				
47011302	**Acceptable identification**						**47011302**
Fusobacterium mortiferum		93.43	0.40	RHA 1			
Prevotella oralis		6.57	0.06	BGL 10			
47011400	**Low discrimination**						**47011400**
					COC	GRA	
Fusobacterium mortiferum		57.19	0.50	ARA 1	1	1	
Gemella morbillorum		21.11	0.32		99	99	
Lactobacillus fermentum		17.08	0.42		1	99	
47011402	**Low discrimination**						**47011402**
					GRA		
Lactobacillus fermentum		67.47	0.42	ESL 1	99		
Fusobacterium mortiferum		25.10	0.33		1		
Atopobium minutum		5.39	0.17		99		
47011500	**Low discrimination**						**47011500**
					GRA		
Fusobacterium mortiferum		80.06	0.56	ARA 1	1		
Lactobacillus fermentum		16.72	0.46		99		
Actinomyces meyeri		1.21	0.16		99		
47011502	**Low discrimination**						**47011502**
					GRA	CAT	
Lactobacillus fermentum		56.04	0.46	ESL 1	99	1	
Fusobacterium mortiferum		29.81	0.40		1	5	
Capnocytophaga ochracea		9.42	0.17		1	90	
47012000	**Low discrimination**						**47012000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	CAT		
Eubacterium limosum		83.22	0.66	SUC 1	1	1		
Clostridium difficile		13.45	0.54		99	1		
Propionibacterium granulosum		2.39	0.54		1	90		
47012010	**Very good identification**							**47012010**
					SPO	CAT		
Clostridium difficile		98.25	0.66	SUC 1	99	1		
Eubacterium limosum		1.26	0.32		1	1		
Propionibacterium granulosum		0.49	0.39		1	90		
47012050	**Good identification**							**47012050**
					SPO			
Clostridium difficile		99.76	0.44	SUC 1	99			
Eubacterium limosum		0.25	-.03	SOR 5	1			
47012100	**Low discrimination**							**47012100**
					GRA	SPO		
Fusobacterium mortiferum		57.84	0.53	MAN 1	1	1		
Eubacterium limosum		35.25	0.32		99	1		
Clostridium difficile		5.70	0.20		99	99		
47012102	**Low discrimination**							**47012102**
					GRA			
Fusobacterium mortiferum		87.62	0.37	MAN 1	1			
Eubacterium limosum		4.86	-.03	BGL 10	99			
Propionibacterium propionicum		4.07	0.15		99			
47013000	**Low discrimination**							**47013000**
					GRA	CAT		
Propionibacterium granulosum		53.99	0.54	ESL 1	99	90		
Fusobacterium mortiferum		20.02	0.50		1	5		
Eubacterium limosum		18.98	0.32		99	1		
47013002	**Low discrimination**							**47013002**
					GRA	SPO	CAT	
Propionibacterium granulosum		71.26	0.44	ESL 1	99	1	90	
Clostridium innocuum		13.46	0.22		99	99	1	
Fusobacterium mortiferum		11.75	0.33		1	1	5	
47013020	**Acceptable identification**							**47013020**
					GRA	CAT		
Propionibacterium granulosum		94.60	0.44	ESL 1	99	90		
Atopobium minutum		1.67	0.11		99	1		
Fusobacterium mortiferum		1.42	0.16		1	5		
47013100	**Good identification**							**47013100**
					GRA	CAT		
Fusobacterium mortiferum		97.52	0.56	MAN 1	1	5		
Propionibacterium granulosum		1.14	0.20		99	90		
Actinomyces naeslundii		0.42	0.27		99	25		
47013102	**Acceptable identification**							**47013102**
					GRA			
Fusobacterium mortiferum		91.21	0.40	MAN 1	1			
Actinomyces naeslundii		3.56	0.27	BGL 10	99			
Propionibacterium propionicum		2.83	0.15		99			
47014000	**Low discrimination**							**47014000**
					COC	GRA		
Fusobacterium mortiferum		80.75	0.62	XYL 7	1	1		
Gemella morbillorum		11.14	0.37		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium limosum		2.51	0.18		1	99	
47014002	**Low discrimination**						**47014002**
					GRA	CAT	
Fusobacterium mortiferum		86.31	0.45	XYL 7	1	5	
Atopobium minutum		11.19	0.25	BGL 10	99	1	
Propionibacterium granulosum		1.41	0.11		99	90	
47014100	**Very good identification**						**47014100**
					COC	GRA	
Fusobacterium mortiferum		98.39	0.68	XYL 7	1	1	
Actinomyces meyeri		1.55	0.28		1	99	
Gemella morbillorum		0.06	0.03		99	99	
47014102	**Good identification**						**47014102**
					GRA		
Fusobacterium mortiferum		99.56	0.52	XYL 7	1		
Actinomyces meyeri		0.44	0.02	BGL 10	99		
47015000	**Low discrimination**						**47015000**
					COC	GRA	
Fusobacterium mortiferum		89.83	0.65	XYL 7	1	1	
Gemella morbillorum		4.45	0.32		99	99	
Lactobacillus fermentum		2.12	0.38		1	99	
47015002	**Low discrimination**						**47015002**
					GRA		
Fusobacterium mortiferum		77.80	0.48	XYL 7	1		
Lactobacillus fermentum		16.48	0.38	BGL 10	99		
Eubacterium saburreum		2.63	0.21		85		
47015100	**Good identification**						**47015100**
					GRA		
Fusobacterium mortiferum		94.16	0.71	XYL 7	1		
Eubacterium saburreum		2.01	0.41		85		
Lactobacillus fermentum		1.55	0.42		99		
47015102	**Low discrimination**						**47015102**
					GRA	CAT	
Fusobacterium mortiferum		77.66	0.55	XYL 7	1	5	
Lactobacillus fermentum		11.51	0.42	BGL 10	99	1	
Capnocytophaga ochracea		4.67	0.20		1	90	
47015110	**Low discrimination**						**47015110**
Eubacterium saburreum		63.32	0.39	IND 99			
Fusobacterium mortiferum		36.68	0.37				
47015120	**Low discrimination**						**47015120**
					CAT		
Fusobacterium mortiferum		22.79	0.37	XYL 7	5		
Capnocytophaga ochracea		15.06	0.20	NAG 1	90		
Bacteroides fragilis		13.49	0.10		99		
47015140	**Good identification**						**47015140**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
47015322	**Low discrimination**						**47015322**
					CAT		
Parabacteroides distasonis		87.90	0.45	RAF 99	76		
Prevotella oralis		6.21	0.12		1		
Bacteroides fragilis		2.89	-.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47015332	**Good identification**						**47015332**
Parabacteroides distasonis		100.0	0.40	RAF 99			
47015400	**Low discrimination**						**47015400**
					COC	GRA	
Lactobacillus fermentum		56.80	0.38	ESL 1	1	99	
Fusobacterium mortiferum		24.37	0.30		1	1	
Gemella morbillorum		13.28	0.16		99	99	
47015402	**Good identification**						**47015402**
					GRA		
Lactobacillus fermentum		95.45	0.38	ESL 1	99		
Fusobacterium mortiferum		4.55	0.14		1		
47015500	**Low discrimination**						**47015500**
					GRA		
Lactobacillus fermentum		46.58	0.42	ESL 1	99		
Fusobacterium mortiferum		28.58	0.37		1		
Bifidobacterium longum subsp. longum		13.67	0.26		99		
47015502	**Low discrimination**						**47015502**
Lactobacillus fermentum		57.47	0.42	ESL 1			
Bifidobacterium adolescentis		17.64	0.28				
Bifidobacterium longum subsp. longum		16.86	0.26				
47015522	**Low discrimination**						**47015522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		44.23	0.36	MLT 99	99	1	
Capnocytophaga ochracea		24.28	0.23	RAF 97	1	90	
Parabacteroides distasonis		16.30	0.33		1	76	
47015530	**Good identification**						**47015530**
Bifidobacterium longum subsp. longum		100.0	0.40	MLT 99			
47015532	**Low discrimination**						**47015532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		86.78	0.40	MLT 99	99	1	
Parabacteroides distasonis		10.12	0.28	RAF 97	1	76	
Capnocytophaga ochracea		3.11	0.07		1	90	
47015720	**Good identification**						**47015720**
					GRA		
Bacteroides vulgatus		98.81	0.40	MLT 99	1		
Eubacterium contortum		1.19	0.14	RAF 97	99		
47015722	**Low discrimination**						**47015722**
					GRA		
Parabacteroides distasonis		88.67	0.41	RAF 99	1		
Bacteroides vulgatus		9.37	0.06		1		
Eubacterium contortum		1.96	0.01		99		
47015732	**Acceptable identification**						**47015732**
					GRA		
Parabacteroides distasonis		93.54	0.36	RAF 99	1		
Clostridium clostridioforme		3.76	0.07		50		
Bifidobacterium longum subsp. longum		2.70	0.06		99		
47016010	**Good identification**						**47016010**
					SPO		
Clostridium difficile		99.76	0.44	SUC 1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium limosum		0.25	-.03	XYL 5	1			
47017100	**Good identification**							**47017100**
Fusobacterium mortiferum		100.0	0.37	MAN 1				
47020000	**Acceptable identification**							**47020000**
					GRA	SPO		
Clostridium botulinum biovar A		91.97	0.51	SUC 7	99	99		
Fusobacterium mortiferum		2.54	0.38	MNS 7	1	1		
Clostridium sporogenes		1.62	0.19		99	99		
47020040	**Acceptable identification**							**47020040**
Clostridium botulinum biovar A		91.91	0.42	SUC 7				
Clostridium botulinum biovar B		8.09	0.21	MNS 7				
47020100	**Low discrimination**							**47020100**
					GRA	SPO		
Fusobacterium mortiferum		86.46	0.44	FRU 99	1	1		
Clostridium botulinum biovar A		13.54	0.17		99	99		
47021000	**Low discrimination**							**47021000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		48.18	0.41	FRU 99	1	1	5	
Propionibacterium granulosum		17.47	0.30		99	1	90	
Clostridium botulinum biovar A		11.74	0.17		99	99	1	
47021100	**Low discrimination**							**47021100**
					COC	GRA		
Fusobacterium mortiferum		74.55	0.47	FRU 99	1	1		
Eubacterium saburreum		16.36	0.34		1	85		
Atopobium parvulum		7.69	0.19		99	99		
47021101	**Good identification**							**47021101**
					GRA	SPO		
Clostridium chauvoei		99.46	0.40	MLT 99	99	99		
Fusobacterium mortiferum		0.54	0.13	TRE 10	1	1		
47030000	**Low discrimination**							**47030000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		62.83	0.72		1	1	5	
Propionibacterium granulosum		34.18	0.64		99	1	90	
Clostridium botulinum biovar A		1.47	0.31		99	99	1	
47030001	**Low discrimination**							**47030001**
					GRA	CAT		
Fusobacterium mortiferum		64.77	0.38	URE 1	1	5		
Propionibacterium granulosum		35.23	0.30		99	90		
47030002	**Low discrimination**							**47030002**
					GRA	CAT		
Propionibacterium granulosum		54.16	0.54	ESL 1	99	90		
Fusobacterium mortiferum		44.25	0.56		1	5		
Atopobium minutum		1.45	0.25		99	1		
47030010	**Low discrimination**							**47030010**
					GRA	SPO	CAT	
Propionibacterium granulosum		64.12	0.49	ESL 1	99	1	90	
Clostridium difficile		27.15	0.30	MLZ 12	99	99	1	
Fusobacterium mortiferum		8.73	0.38		1	1	5	
47030012	**Acceptable identification**							**47030012**
					GRA	SPO	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		92.80	0.39	ESL 1	99	1	90	
Fusobacterium mortiferum		5.62	0.21	MLZ 12	1	1	5	
Clostridium difficile		1.59	-.04		99	99	1	
47030020	**Acceptable identification**							**47030020**
					GRA	CAT		
Propionibacterium granulosum		91.33	0.54	ESL 1	99	90		
Fusobacterium mortiferum		6.78	0.38		1	5		
Atopobium minutum		1.22	0.20		99	1		
47030022	**Acceptable identification**							**47030022**
					GRA	CAT		
Propionibacterium granulosum		92.80	0.43	ESL 1	99	90		
Atopobium minutum		3.30	0.17		99	1		
Fusobacterium mortiferum		3.06	0.21		1	5		
47030030	**Good identification**							**47030030**
					SPO	CAT		
Propionibacterium granulosum		98.32	0.39	ESL 1	1	90		
Clostridium difficile		1.68	-.04	MLZ 12	99	1		
47030100	**Very good identification**							**47030100**
					COC	GRA	CAT	
Fusobacterium mortiferum		99.61	0.78		1	1	5	
Propionibacterium granulosum		0.24	0.30		1	99	90	
Atopobium parvulum		0.16	0.19		99	99	1	
47030101	**Acceptable identification**							**47030101**
					GRA	SPO		
Fusobacterium mortiferum		93.80	0.44	URE 1	1	1		
Clostridium chauvoei		6.20	0.12		99	99		
47030102	**Very good identification**							**47030102**
					COC	GRA	CAT	
Fusobacterium mortiferum		99.23	0.62	BGL 10	1	1	5	
Propionibacterium granulosum		0.53	0.20		1	99	90	
Atopobium parvulum		0.25	0.06		99	99	1	
47030110	**Good identification**							**47030110**
					GRA	SPO	CAT	
Fusobacterium mortiferum		95.67	0.44	MLZ 1	1	1	5	
Propionibacterium granulosum		3.04	0.15		99	1	90	
Clostridium difficile		1.29	-.04		99	99	1	
47030120	**Low discrimination**							**47030120**
					GRA	CAT		
Fusobacterium mortiferum		77.67	0.44	NAG 1	1	5		
Clostridium perfringens		15.69	0.21		99	1		
Propionibacterium granulosum		4.53	0.20		99	90		
47030140	**Good identification**							**47030140**
					GRA	SPO		
Fusobacterium mortiferum		98.38	0.44	SOR 1	1	1		
Clostridium botulinum biovar B		1.62	-.03		99	99		
47030200	**Low discrimination**							**47030200**
					GRA	CAT		
Propionibacterium granulosum		62.49	0.38	ESL 1	99	90		
Fusobacterium mortiferum		37.51	0.38	RHA 3	1	5		
47030300	**Good identification**							**47030300**
Fusobacterium mortiferum		100.0	0.44	RHA 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47030400	**Low discrimination**							**47030400**
					COC	GRA	CAT	
Fusobacterium mortiferum		63.34	0.38	ARA 1	1	1	5	
Propionibacterium granulosum		34.46	0.30		1	99	90	
Gemella morbillorum		2.20	0.03		99	99	1	
47030500	**Good identification**							**47030500**
Fusobacterium mortiferum		100.0	0.44	ARA 1				
47031000	**Low discrimination**							**47031000**
					GRA	CAT		
Fusobacterium mortiferum		71.85	0.75		1	5		
Propionibacterium granulosum		26.06	0.64		99	90		
Lactobacillus jensenii		1.18	0.34		99	1		
47031001	**Low discrimination**							**47031001**
					GRA	SPO	CAT	
Fusobacterium mortiferum		67.41	0.41	URE 1	1	1	5	
Propionibacterium granulosum		24.45	0.30		99	1	90	
Clostridium chauvoei		6.94	0.12		99	99	1	
47031002	**Low discrimination**							**47031002**
					GRA	CAT		
Fusobacterium mortiferum		51.51	0.59	BGL 10	1	5		
Propionibacterium granulosum		42.03	0.54		99	90		
Lactobacillus jensenii		5.06	0.31		99	1		
47031010	**Low discrimination**							**47031010**
					GRA	CAT		
Propionibacterium granulosum		69.99	0.49	ESL 1	99	90		
Fusobacterium mortiferum		14.29	0.41	MLZ 12	1	5		
Lactobacillus jensenii		7.72	0.26		99	1		
47031012	**Low discrimination**							**47031012**
					GRA	CAT		
Propionibacterium granulosum		68.37	0.39	ESL 1	99	90		
Lactobacillus jensenii		20.12	0.23	MLZ 12	99	1		
Fusobacterium mortiferum		6.21	0.24		1	5		
47031020	**Low discrimination**							**47031020**
					GRA	CAT		
Propionibacterium granulosum		60.56	0.54	ESL 1	99	90		
Clostridium perfringens		28.62	0.41		99	1		
Fusobacterium mortiferum		6.75	0.41		1	5		
47031022	**Low discrimination**							**47031022**
					CAT			
Propionibacterium granulosum		48.07	0.43	ESL 1	90			
Clostridium perfringens		42.76	0.35		1			
Lactobacillus jensenii		5.79	0.21		1			
47031030	**Low discrimination**							**47031030**
					CAT			
Propionibacterium granulosum		81.77	0.39	ESL 1	90			
Lactobacillus jensenii		9.02	0.16	MLZ 12	1			
Eubacterium saburreum		6.35	0.22		1			
47031040	**Low discrimination**							**47031040**
					GRA	SPO	CAT	
Fusobacterium mortiferum		70.76	0.41	SOR 1	1	1	5	
Propionibacterium granulosum		25.66	0.30		99	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar B		1.81	-.03		99	99	1	
47031100	**Low discrimination**							**47031100**
					COC	GRA		
Fusobacterium mortiferum		89.60	0.81		1	1		
Atopobium parvulum		9.24	0.53		99	99		
Eubacterium saburreum		0.60	0.42		1	85		
47031101	**Low discrimination**							**47031101**
					COC	GRA	SPO	
Clostridium chauvoei		79.51	0.47	MLT 99	1	99	99	
Fusobacterium mortiferum		18.22	0.47	TRE 10	1	1	1	
Atopobium parvulum		1.88	0.19		99	99	1	
47031102	**Low discrimination**							**47031102**
					COC	GRA		
Fusobacterium mortiferum		83.38	0.65	BGL 10	1	1		
Atopobium parvulum		13.66	0.40		99	99		
Clostridium perfringens		1.47	0.24		1	99		
47031110	**Low discrimination**							**47031110**
					COC	GRA		
Fusobacterium mortiferum		60.14	0.47	MLZ 1	1	1		
Eubacterium saburreum		32.39	0.41		1	85		
Atopobium parvulum		6.20	0.19		99	99		
47031120	**Low discrimination**							**47031120**
					COC	GRA		
Clostridium perfringens		78.04	0.56	MLT 99	1	99		
Atopobium parvulum		10.55	0.40		99	99		
Fusobacterium mortiferum		5.85	0.47		1	1		
47031121	**Low discrimination**							**47031121**
					SPO			
Clostridium chauvoei		88.73	0.38	MLT 99	99			
Clostridium perfringens		8.21	0.21	TRE 10	84			
Actinomyces naeslundii		1.33	0.30		1			
47031122	**Low discrimination**							**47031122**
					COC	GRA		
Clostridium perfringens		88.38	0.50	MLT 99	1	99		
Atopobium parvulum		4.48	0.27		99	99		
Parabacteroides distasonis		2.07	0.31		1	1		
47031140	**Low discrimination**							**47031140**
					COC	GRA		
Fusobacterium mortiferum		84.63	0.47	SOR 1	1	1		
Atopobium parvulum		8.73	0.19		99	99		
Clostridium perfringens		6.65	0.17		1	99		
47031160	**Good identification**							**47031160**
					COC	GRA		
Clostridium perfringens		98.90	0.43	MLT 99	1	99		
Atopobium parvulum		0.71	0.06		99	99		
Fusobacterium mortiferum		0.39	0.13		1	1		
47031162	**Good identification**							**47031162**
Clostridium perfringens		100.0	0.38	MLT 99				
47031200	**Low discrimination**							**47031200**
					GRA	CAT		
Propionibacterium granulosum		52.62	0.38	ESL 1	99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		47.38	0.41	RHA 3	1	5		
47031300	**Acceptable identification**							**47031300**
					COC	GRA		
Fusobacterium mortiferum		90.65	0.47	RHA 1	1	1		
Atopobium parvulum		9.35	0.19		99	99		
47031322	**Low discrimination**							**47031322**
					GRA			
Parabacteroides distasonis		82.23	0.40	RAF 99	1			
Clostridium perfringens		11.84	0.16	XYL 90	99			
Prevotella oralis		5.93	0.07		1			
47031400	**Low discrimination**							**47031400**
					GRA	CAT		
Fusobacterium mortiferum		73.39	0.41	ARA 1	1	5		
Propionibacterium granulosum		26.62	0.30		99	90		
47031500	**Low discrimination**							**47031500**
					COC	GRA		
Fusobacterium mortiferum		89.83	0.47	ARA 1	1	1		
Atopobium parvulum		9.27	0.19		99	99		
Bifidobacterium longum subsp. longum		0.90	0.07		1	99		
47031722	**Good identification**							**47031722**
Parabacteroides distasonis		100.0	0.36	RAF 99				
47032000	**Low discrimination**							**47032000**
					SPO	CAT		
Propionibacterium granulosum		77.95	0.62	ESL 1	1	90		
Clostridium difficile		11.00	0.35		99	1		
Eubacterium limosum		9.14	0.32		1	1		
47032002	**Good identification**							**47032002**
					GRA	SPO	CAT	
Propionibacterium granulosum		97.91	0.52	ESL 1	99	1	90	
Fusobacterium mortiferum		1.07	0.21		1	1	5	
Clostridium difficile		0.56	0.01		99	99	1	
47032010	**Low discrimination**							**47032010**
					SPO	CAT		
Clostridium difficile		83.36	0.47	SUC 1	99	1		
Propionibacterium granulosum		16.50	0.47	TRE 7	1	90		
Eubacterium limosum		0.14	-.03		1	1		
47032012	**Low discrimination**							**47032012**
					SPO	CAT		
Propionibacterium granulosum		83.05	0.37	ESL 1	1	90		
Clostridium difficile		16.95	0.13	MLZ 12	99	1		
47032020	**Good identification**							**47032020**
					SPO	CAT		
Propionibacterium granulosum		98.97	0.52	ESL 1	1	90		
Clostridium difficile		0.56	0.01		99	1		
Eubacterium limosum		0.47	-.03		1	1		
47032022	**Good identification**							**47032022**
Propionibacterium granulosum		100.0	0.41	ESL 1				
47032030	**Low discrimination**							**47032030**
					SPO	CAT		
Propionibacterium granulosum		83.05	0.37	ESL 1	1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium difficile		16.95	0.13	MLZ 12	99	1		
47032100	**Low discrimination**							**47032100**
					GRA	SPO	CAT	
Fusobacterium mortiferum		81.88	0.44	MAN 1	1	1	5	
Propionibacterium granulosum		14.40	0.28		99	1	90	
Clostridium difficile		2.03	0.01		99	99	1	
47032200	**Acceptable identification**							**47032200**
					SPO	CAT		
Propionibacterium granulosum		92.22	0.36	ESL 1	1	90		
Clostridium difficile		4.25	0.01	RHA 3	99	1		
Eubacterium limosum		3.53	-.03		1	1		
47033000	**Very good identification**							**47033000**
					GRA	SPO	CAT	
Propionibacterium granulosum		95.61	0.62	ESL 1	99	1	90	
Fusobacterium mortiferum		3.53	0.41		1	1	5	
Clostridium innocuum		0.53	0.14		99	99	1	
47033002	**Good identification**							**47033002**
					GRA	SPO	CAT	
Propionibacterium granulosum		95.98	0.52	ESL 1	99	1	90	
Clostridium innocuum		2.12	0.14		99	99	1	
Fusobacterium mortiferum		1.58	0.24		1	1	5	
47033010	**Good identification**							**47033010**
					SPO	CAT		
Propionibacterium granulosum		95.14	0.47	ESL 1	1	90		
Clostridium difficile		4.86	0.13	MLZ 12	99	1		
47033012	**Good identification**							**47033012**
Propionibacterium granulosum		100.0	0.37	ESL 1				
47033020	**Good identification**							**47033020**
					CAT			
Propionibacterium granulosum		99.04	0.52	ESL 1	90			
Clostridium perfringens		0.63	0.07		1			
Actinomyces naeslundii		0.34	0.24		25			
47033022	**Good identification**							**47033022**
					CAT			
Propionibacterium granulosum		97.51	0.41	ESL 1	90			
Actinomyces naeslundii		1.33	0.24		25			
Clostridium perfringens		1.16	0.01		1			
47033030	**Good identification**							**47033030**
Propionibacterium granulosum		100.0	0.37	ESL 1				
47033100	**Low discrimination**							**47033100**
					COC	GRA	CAT	
Fusobacterium mortiferum		80.09	0.47	MAN 1	1	1	5	
Propionibacterium granulosum		9.39	0.28		1	99	90	
Atopobium parvulum		8.26	0.19		99	99	1	
47033103	**Low discrimination**							**47033103**
					SPO			
Actinomyces naeslundii		84.47	0.35	ESL 12	1			
Clostridium chauvoei		15.53	-.01	MLT 94	99			
47033121	**Low discrimination**							**47033121**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		74.23	0.35	ESL 12	1			
Clostridium chauvoei		25.77	0.04	MLT 94	99			
47033123	**Good identification**							**47033123**
Actinomyces naeslundii		100.0	0.35	ESL 12				
47033200	**Good identification**							**47033200**
Propionibacterium granulosum		100.0	0.36	ESL 1				
47034000	**Acceptable identification**							**47034000**
					COC	GRA	CAT	
Fusobacterium mortiferum		92.41	0.53	XYL 7	1	1	5	
Propionibacterium granulosum		6.75	0.30		1	99	90	
Gemella morbillorum		0.43	0.03		99	99	1	
47034002	**Low discrimination**							**47034002**
					GRA	CAT		
Fusobacterium mortiferum		85.89	0.36	XYL 7	1	5		
Propionibacterium granulosum		14.11	0.20	BGL 10	99	90		
47034100	**Very good identification**							**47034100**
					COC	GRA		
Fusobacterium mortiferum		96.29	0.59	XYL 7	1	1		
Atopobium parvulum		3.71	0.23		99	99		
47034102	**Acceptable identification**							**47034102**
					COC	GRA		
Fusobacterium mortiferum		94.23	0.43	XYL 7	1	1		
Atopobium parvulum		5.77	0.10	BGL 10	99	99		
47035000	**Low discrimination**							**47035000**
					COC	GRA	CAT	
Fusobacterium mortiferum		86.49	0.56	XYL 7	1	1	5	
Atopobium parvulum		5.19	0.23		99	99	1	
Propionibacterium granulosum		4.21	0.30		1	99	90	
47035002	**Low discrimination**							**47035002**
					COC	GRA	CAT	
Fusobacterium mortiferum		78.11	0.39	XYL 7	1	1	5	
Propionibacterium granulosum		8.55	0.20	BGL 10	1	99	90	
Atopobium parvulum		7.44	0.10		99	99	1	
47035100	**Low discrimination**							**47035100**
					COC	GRA		
Atopobium parvulum		70.84	0.57	MLT 99	99	99		
Fusobacterium mortiferum		27.84	0.62		1	1		
Eubacterium saburreum		1.32	0.38		1	85		
47035102	**Low discrimination**							**47035102**
					COC	GRA		
Atopobium parvulum		78.50	0.45	MLT 99	99	99		
Fusobacterium mortiferum		19.42	0.46		1	1		
Eubacterium saburreum		1.47	0.25		1	85		
47035110	**Low discrimination**							**47035110**
					COC	GRA		
Eubacterium saburreum		52.08	0.36	IND 99	1	85		
Atopobium parvulum		34.40	0.23		99	99		
Fusobacterium mortiferum		13.52	0.28		1	1		
47035120	**Acceptable identification**							**47035120**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		92.13	0.45	MLT 99	99	99	
Clostridium perfringens		3.71	0.21		1	99	
Fusobacterium mortiferum		2.07	0.28		1	1	
47035122	**Low discrimination**						**47035122**
					COC	GRA	
Parabacteroides distasonis		64.52	0.48	RAF 99	1	1	
Atopobium parvulum		28.85	0.32		99	99	
Clostridium perfringens		3.10	0.16		1	99	
47035132	**Good identification**						**47035132**
Parabacteroides distasonis		98.75	0.43	RAF 99			
Eubacterium saburreum		1.25	0.10				
47035322	**Good identification**						**47035322**
Parabacteroides distasonis		100.0	0.56	RAF 99			
47035332	**Good identification**						**47035332**
Parabacteroides distasonis		100.0	0.51	RAF 99			
47035522	**Low discrimination**						**47035522**
					GRA		
Parabacteroides distasonis		87.04	0.44	RAF 99	1		
Bifidobacterium longum subsp. longum		12.96	0.26		99		
47035532	**Low discrimination**						**47035532**
					GRA		
Parabacteroides distasonis		66.79	0.39	RAF 99	1		
Bifidobacterium longum subsp. longum		31.44	0.30		99		
Clostridium clostridioforme		1.77	0.07		50		
47035722	**Good identification**						**47035722**
Parabacteroides distasonis		97.65	0.52	RAF 99			
Bacteroides thetaiotaomicron		1.60	0.06				
Clostridium clostridioforme		0.75	0.10				
47035732	**Acceptable identification**						**47035732**
Parabacteroides distasonis		92.89	0.48	RAF 99			
Clostridium clostridioforme		6.02	0.22				
Bacteroides thetaiotaomicron		1.10	-.01				
47040100	**Good identification**						**47040100**
Fusobacterium mortiferum		100.0	0.45	FRU 99			
47041000	**Low discrimination**						**47041000**
					GRA		
Fusobacterium mortiferum		60.02	0.42	FRU 99	1		
Eubacterium tortuosum		16.77	0.15		99		
Atopobium minutum		6.98	0.20		99		
47041100	**Acceptable identification**						**47041100**
					GRA		
Fusobacterium mortiferum		90.91	0.48	FRU 99	1		
Lactobacillus catenaformis		4.38	0.15		99		
Collinsella aerofaciens		2.90	-.03		99		
47050000	**Very good identification**						**47050000**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		97.53	0.73		1		
Eubacterium limosum		0.68	0.18		99		
Atopobium minutum		0.48	0.28		99		
47050001	**Good identification**						**47050001**
Fusobacterium mortiferum		100.0	0.39	URE 1			
47050002	**Good identification**						**47050002**
					GRA	SPO	
Fusobacterium mortiferum		91.96	0.56	BGL 10	1	1	
Clostridium barati		3.04	0.16		99	99	
Atopobium minutum		2.69	0.25		99	1	
47050020	**Low discrimination**						**47050020**
					GRA	SPO	
Fusobacterium mortiferum		60.27	0.39	NAG 1	1	1	
Clostridium barati		21.91	0.16		99	99	
Atopobium minutum		9.71	0.20		99	1	
47050040	**Good identification**						**47050040**
					GRA	SPO	
Fusobacterium mortiferum		97.56	0.39	SOR 1	1	1	
Clostridium botulinum biovar A		2.44	-.02		99	99	
47050100	**Very good identification**						**47050100**
					GRA	SPO	
Fusobacterium mortiferum		99.72	0.79		1	1	
Clostridium barati		0.16	0.16		99	99	
Lactobacillus catenaformis		0.07	0.15		99	1	
47050101	**Good identification**						**47050101**
Fusobacterium mortiferum		100.0	0.45	URE 1			
47050102	**Very good identification**						**47050102**
					GRA	SPO	
Fusobacterium mortiferum		95.32	0.63	BGL 10	1	1	
Clostridium barati		4.05	0.24		99	99	
Lactobacillus catenaformis		0.63	0.15		99	1	
47050110	**Good identification**						**47050110**
Fusobacterium mortiferum		100.0	0.45	MLZ 1			
47050120	**Low discrimination**						**47050120**
					GRA	SPO	
Fusobacterium mortiferum		65.48	0.45	NAG 1	1	1	
Clostridium barati		30.60	0.24		99	99	
Clostridium paraputrificum		1.75	-.03		99	99	
47050140	**Good identification**						**47050140**
					GRA	SPO	
Fusobacterium mortiferum		98.66	0.45	SOR 1	1	1	
Clostridium barati		1.34	-.02		99	99	
47050200	**Good identification**						**47050200**
Fusobacterium mortiferum		100.0	0.39	RHA 1			
47050300	**Good identification**						**47050300**
					GRA	SPO	
Fusobacterium mortiferum		98.66	0.45	RHA 1	1	1	
Clostridium barati		1.34	-.02		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47050400	**Good identification**						**47050400**
					COC	GRA	
Fusobacterium mortiferum		96.98	0.39	ARA 1	1	1	
Gemella morbillorum		3.02	0.03		99	99	
47050500	**Good identification**						**47050500**
Fusobacterium mortiferum		100.0	0.45	ARA 1			
47051000	**Low discrimination**						**47051000**
					GRA		
Fusobacterium mortiferum		83.95	0.76		1		
Lactobacillus jensenii		7.80	0.48		99		
Lactobacillus catenaformis		6.30	0.47		99		
47051001	**Low discrimination**						**47051001**
					GRA		
Fusobacterium mortiferum		85.62	0.42	URE 1	1		
Lactobacillus jensenii		7.96	0.14		99		
Lactobacillus catenaformis		6.42	0.12		99		
47051002	**Low discrimination**						**47051002**
					GRA		
Fusobacterium mortiferum		36.94	0.59	BGL 10	1		
Lactobacillus catenaformis		24.93	0.47		99		
Lactobacillus jensenii		20.60	0.45		99		
47051010	**Low discrimination**						**47051010**
					GRA		
Lactobacillus jensenii		73.76	0.40	MLT 95	99		
Fusobacterium mortiferum		24.05	0.42	SAL 95	1		
Lactobacillus catenaformis		1.80	0.12		99		
47051012	**Low discrimination**						**47051012**
					GRA		
Lactobacillus jensenii		89.16	0.37	MLT 95	99		
Fusobacterium mortiferum		4.85	0.25	SAL 95	1		
Lactobacillus catenaformis		3.27	0.12		99		
47051020	**Low discrimination**						**47051020**
					SPO		
Clostridium barati		46.03	0.38	SAL 99	99		
Lactobacillus jensenii		22.99	0.38		1		
Lactobacillus catenaformis		18.56	0.36		1		
47051022	**Low discrimination**						**47051022**
					SPO		
Clostridium barati		76.39	0.46	SAL 99	99		
Lactobacillus catenaformis		10.27	0.36		1		
Lactobacillus jensenii		8.48	0.35		1		
47051040	**Low discrimination**						**47051040**
					GRA	SPO	
Fusobacterium mortiferum		76.84	0.42	SOR 1	1	1	
Clostridium barati		10.26	0.11		99	99	
Lactobacillus jensenii		7.14	0.14		99	1	
47051100	**Good identification**						**47051100**
					GRA		
Fusobacterium mortiferum		90.63	0.82		1		
Lactobacillus catenaformis		4.37	0.50		99		
Collinsella aerofaciens		2.89	0.32		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47051101	**Low discrimination**							**47051101**
					GRA	SPO		
Fusobacterium mortiferum		68.08	0.48	URE 1	1	1		
Clostridium chauvoei		24.21	0.29		99	99		
Lactobacillus catenaformis		3.28	0.15		99	1		
47051102	**Low discrimination**							**47051102**
					GRA	SPO		
Fusobacterium mortiferum		47.85	0.66	BGL 10	1	1		
Clostridium barati		25.75	0.46		99	99		
Lactobacillus catenaformis		20.76	0.50		99	1		
47051110	**Low discrimination**							**47051110**
					GRA			
Fusobacterium mortiferum		89.24	0.48	MLZ 1	1			
Lactobacillus catenaformis		4.30	0.15		99			
Collinsella aerofaciens		2.85	-.03		99			
47051120	**Low discrimination**							**47051120**
					SPO			
Clostridium barati		47.28	0.46	SAL 99	99			
Clostridium paraputrificum		14.12	0.32		99			
Lactobacillus catenaformis		9.53	0.39		1			
47051122	**Low discrimination**							**47051122**
					GRA	CAT	SPO	
Clostridium barati		68.88	0.54	SAL 99	99	1	99	
Prevotella oralis		14.30	0.43		1	1	1	
Capnocytophaga ochracea		5.21	0.36		1	90	1	
47051140	**Low discrimination**							**47051140**
					GRA	SPO		
Fusobacterium mortiferum		79.89	0.48	SOR 1	1	1		
Clostridium barati		13.71	0.20		99	99		
Lactobacillus catenaformis		3.85	0.15		99	1		
47051162	**Low discrimination**							**47051162**
					GRA	SPO	CAT	
Clostridium barati		81.40	0.36	SAL 99	99	99	1	
Capnocytophaga ochracea		7.87	0.20	SOR 8	1	1	90	
Lactobacillus acidophilus		3.39	0.15		99	1	1	
47051200	**Low discrimination**							**47051200**
					GRA	SPO		
Fusobacterium mortiferum		76.84	0.42	RHA 1	1	1		
Clostridium barati		10.26	0.11		99	99		
Lactobacillus jensenii		7.14	0.14		99	1		
47051300	**Low discrimination**							**47051300**
					GRA	SPO		
Fusobacterium mortiferum		79.89	0.48	RHA 1	1	1		
Clostridium barati		13.71	0.20		99	99		
Lactobacillus catenaformis		3.85	0.15		99	1		
47051322	**Low discrimination**							**47051322**
					GRA	SPO		
Prevotella oralis		48.72	0.37	MLT 99	1	1		
Clostridium barati		47.61	0.36	RAF 92	99	99		
Parabacteroides distasonis		1.44	0.24		1	1		
47051400	**Low discrimination**							**47051400**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		80.95	0.42	ARA 1	1		
Lactobacillus jensenii		7.52	0.14		99		
Lactobacillus catenaformis		6.07	0.12		99		
47051500	**Acceptable identification**						**47051500**
					GRA		
Fusobacterium mortiferum		90.91	0.48	ARA 1	1		
Lactobacillus catenaformis		4.38	0.15		99		
Collinsella aerofaciens		2.90	-.03		99		
47052010	**Good identification**						**47052010**
					SPO		
Clostridium difficile		99.05	0.44	SUC 1	99		
Clostridium innocuum		0.70	0.11	CEL 5	99		
Eubacterium limosum		0.24	-.03		1		
47052100	**Acceptable identification**						**47052100**
					GRA	SPO	
Fusobacterium mortiferum		94.30	0.45	MAN 1	1	1	
Eubacterium limosum		1.74	-.03		99	1	
Clostridium difficile		1.47	-.02		99	99	
47053000	**Good identification**						**47053000**
					GRA	SPO	
Clostridium innocuum		96.34	0.56	SAL 99	99	99	
Fusobacterium mortiferum		2.55	0.42		1	1	
Clostridium barati		0.34	0.11		99	99	
47053002	**Good identification**						**47053002**
					GRA	SPO	
Clostridium innocuum		98.26	0.56	SAL 99	99	99	
Clostridium barati		1.04	0.20		99	99	
Fusobacterium mortiferum		0.29	0.25		1	1	
47053100	**Low discrimination**						**47053100**
					GRA	SPO	
Fusobacterium mortiferum		69.80	0.48	MAN 1	1	1	
Clostridium barati		11.98	0.20		99	99	
Clostridium innocuum		11.43	0.22		99	99	
47053122	**Acceptable identification**						**47053122**
					GRA	SPO	
Clostridium barati		90.53	0.36	SAL 99	99	99	
Prevotella oralis		2.18	0.09	MAN 8	1	1	
Lactobacillus acidophilus		1.61	0.08		99	1	
47053400	**Good identification**						**47053400**
Clostridium innocuum		100.0	0.43	SAL 99			
47053402	**Good identification**						**47053402**
Clostridium innocuum		100.0	0.43	SAL 99			
47054000	**Good identification**						**47054000**
					COC	GRA	
Fusobacterium mortiferum		99.58	0.53	XYL 7	1	1	
Gemella morbillorum		0.42	0.03		99	99	
47054002	**Good identification**						**47054002**
Fusobacterium mortiferum		100.0	0.37	XYL 7			
47054100	**Very good identification**						**47054100**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		100.0	0.60	XYL 7				
47054102	**Good identification**							**47054102**
Fusobacterium mortiferum		100.0	0.43	XYL 7				
47055000	**Very good identification**							**47055000**
					GRA			
Fusobacterium mortiferum		97.54	0.56	XYL 7	1			
Lactobacillus jensenii		1.22	0.14		99			
Lactobacillus catenaformis		0.98	0.12		99			
47055002	**Low discrimination**							**47055002**
					GRA			
Fusobacterium mortiferum		80.27	0.40	XYL 7	1			
Lactobacillus catenaformis		7.27	0.12	BGL 10	99			
Lactobacillus jensenii		6.01	0.11		99			
47055100	**Good identification**							**47055100**
					COC	GRA		
Fusobacterium mortiferum		93.71	0.63	XYL 7	1	1		
Atopobium parvulum		5.04	0.30		99	99		
Lactobacillus catenaformis		0.61	0.15		1	99		
47055102	**Low discrimination**							**47055102**
					COC	GRA	SPO	
Fusobacterium mortiferum		73.68	0.46	XYL 7	1	1	1	
Atopobium parvulum		6.29	0.17	BGL 10	99	99	1	
Clostridium barati		5.32	0.12		1	99	99	
47055322	**Low discrimination**							**47055322**
					GRA	SPO		
Parabacteroides distasonis		64.70	0.40	RAF 99	1	1		
Prevotella oralis		30.12	0.22		1	1		
Clostridium barati		2.41	0.02		99	99		
47055332	**Good identification**							**47055332**
Parabacteroides distasonis		100.0	0.35	RAF 99				
47055722	**Low discrimination**							**47055722**
Parabacteroides distasonis		74.45	0.36	RAF 99				
Bacteroides uniformis		23.36	0.09					
Clostridium clostridioforme		2.20	0.05					
47060100	**Good identification**							**47060100**
Fusobacterium mortiferum		100.0	0.36	FRU 99				
47061100	**Low discrimination**							**47061100**
					COC	GRA		
Fusobacterium mortiferum		58.07	0.39	FRU 99	1	1		
Atopobium parvulum		41.93	0.25		99	99		
47070000	**Very good identification**							**47070000**
					GRA	CAT		
Fusobacterium mortiferum		96.93	0.64		1	5		
Propionibacterium granulosum		1.60	0.30		99	90		
Lactobacillus jensenii		1.37	0.22		99	1		
47070002	**Low discrimination**							**47070002**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		88.37	0.47	BGL 10	1	5		
Lactobacillus jensenii		7.50	0.19		99	1		
Propionibacterium granulosum		3.28	0.20		99	90		
47070100	**Very good identification**							**47070100**
					COC	GRA	SPO	
Fusobacterium mortiferum		98.87	0.70		1	1	1	
Atopobium parvulum		1.08	0.25		99	99	1	
Clostridium barati		0.05	-.02		1	99	99	
47070101	**Good identification**							**47070101**
Fusobacterium mortiferum		100.0	0.36	URE 1				
47070102	**Good identification**							**47070102**
					COC	GRA	SPO	
Fusobacterium mortiferum		96.36	0.54	BGL 10	1	1	1	
Atopobium parvulum		1.67	0.12		99	99	1	
Clostridium barati		1.19	0.06		1	99	99	
47070110	**Good identification**							**47070110**
Fusobacterium mortiferum		100.0	0.36	MLZ 1				
47070120	**Low discrimination**							**47070120**
					COC	GRA	SPO	
Fusobacterium mortiferum		70.28	0.36	NAG 1	1	1	1	
Atopobium parvulum		13.43	0.12		99	99	1	
Clostridium barati		9.56	0.06		1	99	99	
47070140	**Good identification**							**47070140**
Fusobacterium mortiferum		100.0	0.36	SOR 1				
47070300	**Good identification**							**47070300**
Fusobacterium mortiferum		100.0	0.36	RHA 1				
47070500	**Good identification**							**47070500**
Fusobacterium mortiferum		100.0	0.36	ARA 1				
47071000	**Low discrimination**							**47071000**
					COC	GRA		
Fusobacterium mortiferum		50.69	0.67		1	1		
Lactobacillus jensenii		47.31	0.56		1	99		
Atopobium parvulum		0.86	0.25		99	99		
47071002	**Low discrimination**							**47071002**
					GRA	SPO		
Lactobacillus jensenii		80.63	0.53	MLT 95	99	1		
Fusobacterium mortiferum		14.40	0.50	SAL 95	1	1		
Clostridium barati		1.76	0.20		99	99		
47071010	**Good identification**							**47071010**
					GRA	CAT		
Lactobacillus jensenii		96.41	0.48	MLT 95	99	1		
Fusobacterium mortiferum		3.13	0.33	SAL 95	1	5		
Propionibacterium granulosum		0.46	0.15		99	90		
47071012	**Good identification**							**47071012**
					GRA			
Lactobacillus jensenii		98.56	0.45	MLT 95	99			
Leptotrichia buccalis		0.91	-.03	SAL 95	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		0.53	0.16		1			
47071020	**Low discrimination**							**47071020**
					GRA	SPO		
Lactobacillus jensenii		86.35	0.46	MLT 95	99	1		
Clostridium barati		5.01	0.20	SAL 95	99	99		
Fusobacterium mortiferum		3.74	0.33		1	1		
47071022	**Low discrimination**							**47071022**
					SPO			
Lactobacillus jensenii		72.59	0.43	MLT 95	1			
Clostridium barati		18.96	0.28	SAL 95	99			
Lactobacillus acidophilus		5.58	0.21		1			
47071030	**Good identification**							**47071030**
Lactobacillus jensenii		100.0	0.38	MLT 95				
47071100	**Low discrimination**							**47071100**
					COC	GRA	SPO	
Fusobacterium mortiferum		57.58	0.73		1	1	1	
Atopobium parvulum		41.57	0.59		99	99	1	
Clostridium barati		0.33	0.20		1	99	99	
47071101	**Low discrimination**							**47071101**
					COC	GRA	SPO	
Fusobacterium mortiferum		51.52	0.39	URE 1	1	1	1	
Atopobium parvulum		37.20	0.25		99	99	1	
Clostridium chauvoei		6.82	0.12		1	99	99	
47071102	**Low discrimination**							**47071102**
					COC	GRA		
Atopobium parvulum		39.66	0.46	MLT 99	99	99		
Fusobacterium mortiferum		34.59	0.57		1	1		
Lactobacillus acidophilus		18.35	0.39		1	99		
47071110	**Low discrimination**							**47071110**
					COC	GRA		
Fusobacterium mortiferum		51.65	0.39	MLZ 1	1	1		
Atopobium parvulum		37.29	0.25		99	99		
Lactobacillus jensenii		6.89	0.14		1	99		
47071120	**Low discrimination**							**47071120**
					COC	SPO		
Atopobium parvulum		49.96	0.46	MLT 99	99	1		
Clostridium septicum		25.00	0.35		1	99		
Clostridium barati		6.83	0.28		1	99		
47071122	**Low discrimination**							**47071122**
					COC	SPO		
Lactobacillus acidophilus		35.87	0.39	MLT 94	1	1		
Clostridium barati		31.77	0.36	SAL 99	1	99		
Atopobium parvulum		13.68	0.33		99	1		
47071140	**Low discrimination**							**47071140**
					COC	GRA	SPO	
Fusobacterium mortiferum		56.44	0.39	SOR 1	1	1	1	
Atopobium parvulum		40.75	0.25		99	99	1	
Clostridium barati		2.82	0.01		1	99	99	
47071300	**Low discrimination**							**47071300**
					COC	GRA	SPO	
Fusobacterium mortiferum		56.44	0.39	RHA 1	1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		40.75	0.25		99	99	1	
Clostridium barati		2.82	0.01		1	99	99	
47071500	**Low discrimination**							**47071500**
					COC	GRA		
Fusobacterium mortiferum		58.07	0.39	ARA 1	1	1		
Atopobium parvulum		41.93	0.25		99	99		
47072100	**Good identification**							**47072100**
Fusobacterium mortiferum		100.0	0.36	MAN 1				
47073000	**Acceptable identification**							**47073000**
					GRA	SPO		
Clostridium innocuum		93.99	0.48	SAL 99	99	99		
Fusobacterium mortiferum		2.11	0.33		1	1		
Lactobacillus jensenii		1.97	0.22		99	1		
47073002	**Good identification**							**47073002**
					SPO	CAT		
Clostridium innocuum		97.35	0.48	SAL 99	99	1		
Lactobacillus jensenii		1.36	0.19		1	1		
Propionibacterium granulosum		0.45	0.18		1	90		
47073100	**Low discrimination**							**47073100**
					COC	GRA	SPO	
Fusobacterium mortiferum		45.45	0.39	MAN 1	1	1	1	
Atopobium parvulum		32.82	0.25		99	99	1	
Clostridium innocuum		8.75	0.14		1	99	99	
47073101	**Good identification**							**47073101**
Actinomyces naeslundii		100.0	0.37	ESL 12				
47073103	**Good identification**							**47073103**
Actinomyces naeslundii		100.0	0.37	ESL 12				
47073121	**Good identification**							**47073121**
Actinomyces naeslundii		100.0	0.37	ESL 12				
47073123	**Good identification**							**47073123**
Actinomyces naeslundii		100.0	0.37	ESL 12				
47073400	**Good identification**							**47073400**
Clostridium innocuum		100.0	0.35	SAL 99				
47073402	**Good identification**							**47073402**
Clostridium innocuum		100.0	0.35	SAL 99				
47074000	**Good identification**							**47074000**
Fusobacterium mortiferum		100.0	0.44	XYL 7				
47074100	**Low discrimination**							**47074100**
					COC	GRA		
Fusobacterium mortiferum		78.75	0.51	XYL 7	1	1		
Atopobium parvulum		21.25	0.30		99	99		
47075000	**Low discrimination**							**47075000**
					COC	GRA		
Fusobacterium mortiferum		64.72	0.48	XYL 7	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		27.17	0.30		99	99		
Lactobacillus jensenii		8.11	0.22		1	99		
47075100	**Good identification**							**47075100**
					COC	GRA		
Atopobium parvulum		94.62	0.64	MLT 99	99	99		
Fusobacterium mortiferum		5.31	0.54		1	1		
Actinomyces israelii		0.05	0.22		1	99		
47075102	**Good identification**							**47075102**
					COC	GRA		
Atopobium parvulum		95.60	0.51	MLT 99	99	99		
Fusobacterium mortiferum		3.38	0.37		1	1		
Actinomyces israelii		0.46	0.25		1	99		
47075120	**Good identification**							**47075120**
					COC	GRA	SPO	
Atopobium parvulum		99.41	0.51	MLT 99	99	99	1	
Fusobacterium mortiferum		0.32	0.20		1	1	1	
Clostridium septicum		0.27	0.01		1	99	99	
47075122	**Low discrimination**							**47075122**
					COC	GRA		
Atopobium parvulum		63.48	0.38	MLT 99	99	99		
Parabacteroides distasonis		32.75	0.43		1	1		
Prevotella oralis		1.50	0.07		1	1		
47075132	**Good identification**							**47075132**
					COC	GRA		
Parabacteroides distasonis		96.49	0.39	RAF 99	1	1		
Atopobium parvulum		3.51	0.04		99	99		
47075322	**Good identification**							**47075322**
					COC	GRA		
Parabacteroides distasonis		98.71	0.51	RAF 99	1	1		
Atopobium parvulum		0.64	0.04		99	99		
Prevotella oralis		0.64	0.01		1	1		
47075332	**Good identification**							**47075332**
Parabacteroides distasonis		98.45	0.47	RAF 99				
Clostridium clostridioforme		1.55	0.10					
47075522	**Good identification**							**47075522**
					COC	GRA		
Parabacteroides distasonis		95.11	0.39	RAF 99	1	1		
Atopobium parvulum		3.04	0.04		99	99		
Clostridium clostridioforme		1.85	0.05		1	50		
47075722	**Low discrimination**							**47075722**
Parabacteroides distasonis		89.90	0.48	RAF 99				
Bacteroides thetaiotaomicron		5.82	0.11					
Clostridium clostridioforme		4.28	0.19					
47075732	**Low discrimination**							**47075732**
Parabacteroides distasonis		69.00	0.43	RAF 99				
Clostridium clostridioforme		27.78	0.31					
Bacteroides thetaiotaomicron		3.23	0.04					
47077102	**Low discrimination**							**47077102**
					COC	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		49.00	0.38	MLT 99	1	1	
Clostridium tertium		27.70	0.19	SAL 92	1	99	
Atopobium parvulum		18.25	0.17		99	1	
47077502	**Low discrimination**						**47077502**
Actinomyces israelii		54.25	0.36	MLT 99			
Bifidobacterium adolescentis		45.75	0.29	SAL 92			
47100000	**Low discrimination**						**47100000**
					COC	SPO	
Gemella morbillorum		47.24	0.64	ESL 1	99	1	
Clostridium botulinum biovar A		38.94	0.60		1	99	
Clostridium bifermentans		10.18	0.50		1	99	
47100020	**Low discrimination**						**47100020**
					COC	SPO	
Gemella morbillorum		80.74	0.48	ESL 1	99	1	
Clostridium bifermentans		9.99	0.29	NAG 10	1	99	
Clostridium botulinum biovar A		6.05	0.26		1	99	
47100040	**Low discrimination**						**47100040**
					COC	SPO	
Clostridium botulinum biovar A		50.74	0.51	SUC 7	1	99	
Clostridium botulinum biovar B		45.03	0.47	MNS 7	1	99	
Gemella morbillorum		2.08	0.30		99	1	
47100050	**Good identification**						**47100050**
Clostridium botulinum biovar B		97.36	0.42	ESL 1			
Clostridium botulinum biovar A		2.25	0.17				
Clostridium novyi biovar A		0.40	0.16				
47100060	**Good identification**						**47100060**
					COC	SPO	
Clostridium botulinum biovar B		96.26	0.42	ESL 1	1	99	
Clostridium botulinum biovar A		2.23	0.17		1	99	
Gemella morbillorum		1.00	0.14		99	1	
47100070	**Good identification**						**47100070**
Clostridium botulinum biovar B		100.0	0.37	ESL 1			
47100101	**Very good identification**						**47100101**
Clostridium chauvoei		100.0	0.57	GAL 99			
47100103	**Good identification**						**47100103**
Clostridium chauvoei		100.0	0.44	GAL 99			
47100121	**Good identification**						**47100121**
					GRA	SPO	
Clostridium chauvoei		99.21	0.48	GAL 99	99	99	
Prevotella bivia		0.59	-.03		1	1	
Prevotella melaninogenica		0.20	-.04		1	1	
47100123	**Good identification**						**47100123**
Clostridium chauvoei		100.0	0.36	GAL 99			
47100400	**Acceptable identification**						**47100400**
					COC	SPO	
Gemella morbillorum		90.88	0.48	ESL 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium botulinum biovar A		6.81	0.26	ARA 10	1	99		
Clostridium bifermentans		1.78	0.15		1	99		
47101000	**Good identification**							**47101000**
					COC	SPO		
Gemella morbillorum		93.25	0.60	ESL 1	99	1		
Eubacterium saburreum		2.38	0.41		1	1		
Clostridium botulinum biovar A		1.44	0.26		1	99		
47101001	**Very good identification**							**47101001**
					COC	SPO		
Clostridium chauvoei		98.69	0.57	LAC 99	1	99		
Gemella morbillorum		1.31	0.26		99	1		
47101003	**Good identification**							**47101003**
Clostridium chauvoei		100.0	0.44	LAC 99				
47101010	**Low discrimination**							**47101010**
					COC			
Eubacterium saburreum		65.98	0.40	IND 99	1			
Gemella morbillorum		31.91	0.26		99			
Lactobacillus jensenii		2.12	0.06		1			
47101020	**Low discrimination**							**47101020**
					COC	GRA		
Gemella morbillorum		51.92	0.43	ESL 1	99	99		
Prevotella melaninogenica		23.45	0.30	NAG 10	1	1		
Clostridium perfringens		20.70	0.36		1	99		
47101021	**Good identification**							**47101021**
					COC	GRA	SPO	
Clostridium chauvoei		99.19	0.48	LAC 99	1	99	99	
Gemella morbillorum		0.44	0.09		99	99	1	
Prevotella melaninogenica		0.20	-.04		1	1	1	
47101023	**Good identification**							**47101023**
Clostridium chauvoei		100.0	0.36	LAC 99				
47101100	**Low discrimination**							**47101100**
					SPO	GRA		
Clostridium chauvoei		82.61	0.57	URE 99	99	99		
Eubacterium saburreum		6.46	0.47		1	85		
Prevotella melaninogenica		5.44	0.30		1	1		
47101101	**Very good identification**							**47101101**
					SPO			
Clostridium chauvoei		100.0	0.91		99			
Actinomyces naeslundii		0.00	0.24		1			
Eubacterium saburreum		0.00	0.13		1			
47101102	**Low discrimination**							**47101102**
					SPO	CAT		
Clostridium chauvoei		67.96	0.44	URE 99	99	1		
Capnocytophaga ochracea		15.24	0.30		1	90		
Eubacterium saburreum		5.31	0.35		1	1		
47101103	**Very good identification**							**47101103**
					SPO			
Clostridium chauvoei		100.0	0.78		99			
Actinomyces naeslundii		0.00	0.24		1			
47101110	**Low discrimination**							**47101110**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO	GRA		
Eubacterium saburreum		84.53	0.46	IND 99	1	85		
Clostridium chauvoei		13.35	0.22		99	99		
Prevotella melaninogenica		0.88	-.04		1	1		
47101111	**Very good identification**							**47101111**
					SPO			
Clostridium chauvoei		99.94	0.57	MLZ 1	99			
Eubacterium saburreum		0.07	0.12		1			
47101113	**Good identification**							**47101113**
Clostridium chauvoei		100.0	0.44	MLZ 1				
47101120	**Low discrimination**							**47101120**
					GRA	SPO		
Prevotella melaninogenica		88.37	0.64	RAF 90	1	1		
Clostridium perfringens		5.78	0.51	FRU 90	99	84		
Clostridium chauvoei		4.52	0.48		99	99		
47101121	**Very good identification**							**47101121**
					GRA	SPO		
Clostridium chauvoei		99.79	0.83		99	99		
Prevotella melaninogenica		0.20	0.30		1	1		
Clostridium perfringens		0.01	0.17		99	84		
47101122	**Low discrimination**							**47101122**
					GRA	CAT		
Prevotella melaninogenica		64.12	0.52	RAF 90	1	1		
Capnocytophaga ochracea		18.38	0.47	FRU 90	1	90		
Clostridium perfringens		10.36	0.45		99	1		
47101123	**Very good identification**							**47101123**
					GRA	CAT	SPO	
Clostridium chauvoei		99.67	0.70		99	1	99	
Prevotella melaninogenica		0.22	0.18		1	1	1	
Capnocytophaga ochracea		0.06	0.12		1	90	1	
47101131	**Good identification**							**47101131**
					GRA	SPO		
Clostridium chauvoei		99.80	0.48	MLZ 1	99	99		
Prevotella melaninogenica		0.20	-.04		1	1		
47101133	**Good identification**							**47101133**
Clostridium chauvoei		100.0	0.36	MLZ 1				
47101141	**Very good identification**							**47101141**
Clostridium chauvoei		100.0	0.57	SOR 1				
47101143	**Good identification**							**47101143**
Clostridium chauvoei		100.0	0.44	SOR 1				
47101160	**Low discrimination**							**47101160**
					GRA	CAT		
Clostridium perfringens		52.44	0.39	FRU 99	99	1		
Prevotella melaninogenica		42.54	0.30		1	1		
Capnocytophaga ochracea		2.84	0.14		1	90		
47101161	**Good identification**							**47101161**
					GRA	SPO		
Clostridium chauvoei		99.56	0.48	SOR 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		0.25	0.04		99	84		
Prevotella melaninogenica		0.20	-.04		1	1		
47101163	**Good identification**							**47101163**
Clostridium chauvoei		100.0	0.36	SOR 1				
47101301	**Very good identification**							**47101301**
Clostridium chauvoei		100.0	0.57	RHA 1				
47101303	**Good identification**							**47101303**
Clostridium chauvoei		100.0	0.44	RHA 1				
47101321	**Good identification**							**47101321**
					GRA	SPO		
Clostridium chauvoei		99.80	0.48	RHA 1	99	99		
Prevotella melaninogenica		0.20	-.04		1	1		
47101323	**Good identification**							**47101323**
Clostridium chauvoei		100.0	0.36	RHA 1				
47101400	**Good identification**							**47101400**
					COC			
Gemella morbillorum		99.47	0.43	ESL 1	99			
Lactobacillus fermentum		0.53	0.16	ARA 10	1			
47101500	**Low discrimination**							**47101500**
					COC	SPO		
Bifidobacterium longum subsp. longum		63.53	0.35	RAF 97	1	1		
Clostridium chauvoei		24.46	0.22	FRU 90	1	99		
Gemella morbillorum		3.57	0.09		99	1		
47101501	**Very good identification**							**47101501**
Clostridium chauvoei		100.0	0.57	ARA 1				
47101502	**Low discrimination**							**47101502**
					GRA	SPO	CAT	
Bifidobacterium longum subsp. longum		70.19	0.35	RAF 97	99	1	1	
Capnocytophaga ochracea		18.68	0.17	FRU 90	1	1	90	
Clostridium chauvoei		4.77	0.10		99	99	1	
47101503	**Good identification**							**47101503**
Clostridium chauvoei		100.0	0.44	ARA 1				
47101510	**Good identification**							**47101510**
Bifidobacterium longum subsp. longum		98.57	0.39	RAF 97				
Eubacterium saburreum		1.43	0.12	FRU 90				
47101512	**Good identification**							**47101512**
					GRA	CAT		
Bifidobacterium longum subsp. longum		95.42	0.39	RAF 97	99	1		
Bifidobacterium adolescentis		2.92	0.15	FRU 90	99	1		
Capnocytophaga ochracea		1.66	0.01		1	90		
47101520	**Low discrimination**							**47101520**
					GRA	CAT		
Bifidobacterium longum subsp. longum		56.15	0.46	RAF 97	99	1		
Prevotella melaninogenica		35.25	0.30	FRU 90	1	1		
Capnocytophaga ochracea		3.74	0.17		1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47101521	**Good identification**							**47101521**
					GRA	SPO		
Clostridium chauvoei		99.49	0.48	ARA 1	99	99		
Bifidobacterium longum subsp. longum		0.32	0.12		99	1		
Prevotella melaninogenica		0.20	-.04		1	1		
47101522	**Low discrimination**							**47101522**
					GRA	CAT		
Bifidobacterium longum subsp. longum		57.14	0.46	RAF 97	99	1		
Capnocytophaga ochracea		34.22	0.34	FRU 90	1	90		
Prevotella melaninogenica		6.83	0.18		1	1		
47101523	**Good identification**							**47101523**
					GRA	SPO	CAT	
Clostridium chauvoei		97.20	0.36	ARA 1	99	99	1	
Bifidobacterium longum subsp. longum		1.75	0.12		99	1	1	
Capnocytophaga ochracea		1.05	-.01		1	1	90	
47101530	**Good identification**							**47101530**
					GRA	CAT		
Bifidobacterium longum subsp. longum		99.20	0.50	RAF 97	99	1		
Capnocytophaga ochracea		0.43	0.01	FRU 90	1	90		
Prevotella melaninogenica		0.37	-.04		1	1		
47101532	**Good identification**							**47101532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		96.24	0.50	RAF 97	99	1		
Capnocytophaga ochracea		3.76	0.17	FRU 90	1	90		
47103101	**Very good identification**							**47103101**
					SPO			
Clostridium chauvoei		99.89	0.57	MAN 1	99			
Actinomyces naeslundii		0.11	0.29		1			
47103103	**Good identification**							**47103103**
					SPO			
Clostridium chauvoei		99.40	0.44	MAN 1	99			
Actinomyces naeslundii		0.60	0.29		1			
47103121	**Good identification**							**47103121**
					GRA	SPO		
Clostridium chauvoei		99.48	0.48	MAN 1	99	99		
Actinomyces naeslundii		0.32	0.29		99	1		
Prevotella melaninogenica		0.20	-.04		1	1		
47103123	**Good identification**							**47103123**
					SPO			
Clostridium chauvoei		98.22	0.36	MAN 1	99			
Actinomyces naeslundii		1.78	0.29		1			
47104000	**Acceptable identification**							**47104000**
					COC	SPO		
Gemella morbillorum		90.72	0.48	ESL 1	99	1		
Clostridium botulinum biovar A		6.80	0.26	XYL 10	1	99		
Clostridium bifermentans		1.78	0.15		1	99		
47105000	**Low discrimination**							**47105000**
					COC	GRA		
Gemella morbillorum		88.32	0.43	ESL 1	99	99		
Eubacterium saburreum		10.93	0.37	XYL 10	1	85		
Fusobacterium mortiferum		0.48	0.15		1	1		
47105010	**Acceptable identification**							**47105010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Eubacterium saburreum		90.93	0.35	IND 99	1		
Gemella morbillorum		9.07	0.09		99		
47105100	**Low discrimination**						**47105100**
					COC	SPO	
Eubacterium saburreum		58.81	0.43	IND 99	1	1	
Atopobium parvulum		15.89	0.23		99	1	
Clostridium chauvoei		14.11	0.22		1	99	
47105101	**Very good identification**						**47105101**
Clostridium chauvoei		100.0	0.57	XYL 1			
47105103	**Good identification**						**47105103**
Clostridium chauvoei		100.0	0.44	XYL 1			
47105110	**Good identification**						**47105110**
Eubacterium saburreum		96.22	0.41	IND 99			
Bifidobacterium longum subsp. longum		3.78	0.13				
47105121	**Good identification**						**47105121**
					GRA	SPO	
Clostridium chauvoei		99.80	0.48	XYL 1	99	99	
Prevotella melaninogenica		0.20	-.04		1	1	
47105123	**Good identification**						**47105123**
					GRA	SPO	
Clostridium chauvoei		95.77	0.36	XYL 1	99	99	
Bacteroides fragilis		2.77	-.03		1	1	
Capnocytophaga ochracea		1.46	0.02		1	1	
47105500	**Good identification**						**47105500**
Bifidobacterium longum subsp. longum		99.07	0.44	RAF 97			
Lactobacillus fermentum		0.93	0.16	FRU 90			
47105502	**Low discrimination**						**47105502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.76	0.44	RAF 97	99	1	
Bifidobacterium adolescentis		8.44	0.28	FRU 90	99	1	
Capnocytophaga ochracea		1.97	0.07		1	90	
47105510	**Good identification**						**47105510**
Bifidobacterium longum subsp. longum		100.0	0.48	RAF 97			
47105512	**Acceptable identification**						**47105512**
Bifidobacterium longum subsp. longum		93.61	0.48	RAF 97			
Bifidobacterium adolescentis		6.39	0.29	FRU 90			
47105520	**Low discrimination**						**47105520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.93	0.54	RAF 97	99	1	
Bacteroides vulgatus		10.39	0.24	FRU 90	1	3	
Capnocytophaga ochracea		0.49	0.07		1	90	
47105522	**Acceptable identification**						**47105522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		94.76	0.54	RAF 97	99	1	
Capnocytophaga ochracea		4.73	0.23	FRU 90	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides fragilis		0.51	-.03		1	99		
47105530	**Very good identification**							**47105530**
Bifidobacterium longum subsp. longum		100.0	0.58	RAF 97				
47105532	**Very good identification**							**47105532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		99.68	0.58	RAF 97	99	1		
Capnocytophaga ochracea		0.33	0.07	FRU 90	1	90		
47105720	**Good identification**							**47105720**
					GRA			
Bacteroides vulgatus		97.89	0.40	RAF 97	1			
Eubacterium contortum		1.17	0.14	FRU 99	99			
Bifidobacterium longum subsp. longum		0.94	0.20		99			
47110000	**Low discrimination**							**47110000**
					COC	GRA		
Gemella morbillorum		67.79	0.66	ESL 1	99	99		
Fusobacterium mortiferum		15.82	0.65		1	1		
Prevotella intermedia		5.51	0.46		1	1		
47110002	**Low discrimination**							**47110002**
					COC	GRA	CAT	
Fusobacterium mortiferum		47.23	0.49	MLT 11	1	1	5	
Propionibacterium granulosum		20.93	0.40	BGL 10	1	99	90	
Gemella morbillorum		18.40	0.32		99	99	1	
47110020	**Low discrimination**							**47110020**
					COC	CAT		
Gemella morbillorum		72.41	0.50	ESL 1	99	1		
Clostridium perfringens		12.72	0.36	NAG 10	1	1		
Propionibacterium granulosum		7.49	0.40		1	90		
47110040	**Very good identification**							**47110040**
					COC	SPO		
Clostridium botulinum biovar B		95.49	0.58	ESL 1	1	99		
Clostridium botulinum biovar A		1.72	0.31		1	99		
Gemella morbillorum		1.35	0.32		99	1		
47110050	**Good identification**							**47110050**
Clostridium botulinum biovar B		100.0	0.52	ESL 1				
47110060	**Good identification**							**47110060**
					COC	SPO		
Clostridium botulinum biovar B		98.64	0.52	ESL 1	1	99		
Clostridium perfringens		1.04	0.24		1	84		
Gemella morbillorum		0.31	0.15		99	1		
47110070	**Good identification**							**47110070**
Clostridium botulinum biovar B		100.0	0.47	ESL 1				
47110100	**Good identification**							**47110100**
					COC	GRA	SPO	
Fusobacterium mortiferum		93.94	0.72	MLT 11	1	1	1	
Gemella morbillorum		1.74	0.32		99	99	1	
Clostridium chauvoei		1.37	0.29		1	99	99	
47110101	**Very good identification**							**47110101**
					GRA	SPO		
Clostridium chauvoei		99.30	0.63	GAL 99	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.70	0.37		1	1	
47110102	**Low discrimination**						**47110102**
					GRA	CAT	
Fusobacterium mortiferum		73.87	0.55	MLT 11	1	5	
Capnocytophaga ochracea		16.22	0.30	BGL 10	1	90	
Clostridium perfringens		2.54	0.20		99	1	
47110103	**Good identification**						**47110103**
					GRA	SPO	
Clostridium chauvoei		99.35	0.50	GAL 99	99	99	
Fusobacterium mortiferum		0.44	0.21		1	1	
Actinomyces viscosus		0.21	0.03		99	1	
47110110	**Low discrimination**						**47110110**
					CAT		
Fusobacterium mortiferum		87.10	0.37	MLT 11	5		
Eubacterium saburreum		10.31	0.20	MLZ 1	1		
Capnocytophaga ochracea		2.60	-.03		90		
47110120	**Low discrimination**						**47110120**
					GRA	CAT	
Prevotella melaninogenica		53.00	0.47	RAF 90	1	1	
Clostridium perfringens		38.12	0.51	GAL 99	99	1	
Capnocytophaga ochracea		3.54	0.30		1	90	
47110121	**Good identification**						**47110121**
					GRA	SPO	
Clostridium chauvoei		98.69	0.55	GAL 99	99	99	
Prevotella melaninogenica		0.76	0.12		1	1	
Clostridium perfringens		0.55	0.17		99	84	
47110122	**Low discrimination**						**47110122**
					GRA	CAT	
Capnocytophaga ochracea		47.40	0.47	RAF 90	1	90	
Clostridium perfringens		26.70	0.45	GAL 99	99	1	
Prevotella melaninogenica		15.03	0.34		1	1	
47110123	**Acceptable identification**						**47110123**
					GRA	CAT	
Clostridium chauvoei		94.84	0.42	GAL 99	99	1	
Capnocytophaga ochracea		2.49	0.12		1	90	
Clostridium perfringens		1.40	0.11		99	1	
47110160	**Low discrimination**						**47110160**
					GRA	SPO	
Clostridium perfringens		79.47	0.39	GAL 99	99	84	
Clostridium botulinum biovar B		10.37	0.18		99	99	
Prevotella melaninogenica		5.86	0.12		1	1	
47110300	**Good identification**						**47110300**
Fusobacterium mortiferum		100.0	0.37	MLT 11			
47110400	**Good identification**						**47110400**
					COC	GRA	
Gemella morbillorum		95.82	0.50	ESL 1	99	99	
Fusobacterium mortiferum		2.03	0.31	ARA 10	1	1	
Prevotella intermedia		0.71	0.12		1	1	
47110500	**Low discrimination**						**47110500**
					COC	GRA	
Fusobacterium mortiferum		62.03	0.37	MLT 11	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		12.66	0.15	ARA 1	99	99		
Bifidobacterium longum subsp. longum		9.03	0.18		1	99		
47111000	**Low discrimination**							**47111000**
					COC	GRA		
Gemella morbillorum		43.15	0.61	ESL 1	99	99		
Fusobacterium mortiferum		28.05	0.68		1	1		
Lactobacillus jensenii		7.03	0.48		1	99		
47111001	**Very good identification**							**47111001**
					COC	GRA	SPO	
Clostridium chauvoei		98.30	0.63	LAC 99	1	99	99	
Gemella morbillorum		0.68	0.27		99	99	1	
Fusobacterium mortiferum		0.44	0.34		1	1	1	
47111002	**Low discrimination**							**47111002**
					GRA			
Lactobacillus jensenii		27.94	0.45	SAL 95	99			
Fusobacterium mortiferum		18.58	0.52	CEL 95	1			
Clostridium perfringens		13.44	0.39		99			
47111003	**Good identification**							**47111003**
					SPO			
Clostridium chauvoei		98.04	0.50	LAC 99	99			
Actinomyces naeslundii		0.87	0.38		1			
Lactobacillus jensenii		0.42	0.11		1			
47111010	**Low discrimination**							**47111010**
					CAT			
Lactobacillus jensenii		39.55	0.40	SAL 95	1			
Eubacterium saburreum		37.34	0.48	CEL 95	1			
Propionibacterium granulosum		8.47	0.35		90			
47111012	**Low discrimination**							**47111012**
					CAT			
Lactobacillus jensenii		68.78	0.37	SAL 95	1			
Eubacterium saburreum		17.19	0.35	CEL 95	1			
Propionibacterium granulosum		5.53	0.25		90			
47111020	**Low discrimination**							**47111020**
					COC	GRA		
Clostridium perfringens		84.99	0.70	LAC 88	1	99		
Prevotella melaninogenica		8.76	0.47		1	1		
Gemella morbillorum		2.63	0.45		99	99		
47111021	**Acceptable identification**							**47111021**
					GRA	SPO		
Clostridium chauvoei		91.72	0.55	LAC 99	99	99		
Clostridium perfringens		6.86	0.36		99	84		
Prevotella melaninogenica		0.71	0.12		1	1		
47111022	**Low discrimination**							**47111022**
					GRA	CAT		
Clostridium perfringens		72.57	0.65	LAC 88	99	1		
Prevotella oralis		11.67	0.48		1	1		
Capnocytophaga ochracea		9.54	0.47		1	90		
47111023	**Low discrimination**							**47111023**
					GRA	SPO		
Clostridium chauvoei		77.36	0.42	LAC 99	99	99		
Clostridium perfringens		15.42	0.31		99	84		
Prevotella oralis		2.48	0.14		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47111030	**Low discrimination**						**47111030**
Clostridium perfringens		52.25	0.36	LAC 88			
Lactobacillus jensenii		19.57	0.30	MLZ 1			
Eubacterium saburreum		13.04	0.35				
47111060	**Very good identification**						**47111060**
					GRA	SPO	
Clostridium perfringens		97.86	0.58	LAC 88	99	84	
Clostridium botulinum biovar B		0.95	0.18		99	99	
Prevotella melaninogenica		0.54	0.12		1	1	
47111062	**Acceptable identification**						**47111062**
					GRA	CAT	
Clostridium perfringens		91.68	0.52	LAC 88	99	1	
Capnocytophaga ochracea		7.03	0.30		1	90	
Prevotella oralis		0.78	0.14		1	1	
47111100	**Low discrimination**						**47111100**
					GRA	SPO	
Fusobacterium mortiferum		28.96	0.75	MLT 11	1	1	
Clostridium chauvoei		27.77	0.63		99	99	
Clostridium perfringens		15.54	0.59		99	84	
47111101	**Excellent identification**						**47111101**
					GRA	SPO	
Clostridium chauvoei		99.97	0.97		99	99	
Fusobacterium mortiferum		0.01	0.40		1	1	
Clostridium perfringens		0.01	0.25		99	84	
47111102	**Low discrimination**						**47111102**
					GRA	CAT	
Capnocytophaga ochracea		62.15	0.64	RAF 90	1	90	
Clostridium perfringens		9.75	0.54	NAG 90	99	1	
Clostridium chauvoei		6.53	0.50		99	1	
47111103	**Very good identification**						**47111103**
					GRA	SPO	
Clostridium chauvoei		99.83	0.84		99	99	
Capnocytophaga ochracea		0.10	0.30		1	1	
Actinomyces naeslundii		0.03	0.46		99	1	
47111110	**Low discrimination**						**47111110**
					CAT		
Eubacterium saburreum		59.59	0.54	IND 99	1		
Capnocytophaga ochracea		15.02	0.32		90		
Fusobacterium mortiferum		7.63	0.40		5		
47111111	**Very good identification**						**47111111**
					SPO		
Clostridium chauvoei		99.87	0.63	MLZ 1	99		
Eubacterium saburreum		0.08	0.20		1		
Actinomyces naeslundii		0.02	0.24		1		
47111112	**Low discrimination**						**47111112**
					CAT	GRA	
Capnocytophaga ochracea		84.00	0.48	RAF 90	90	1	
Eubacterium saburreum		6.54	0.41	MLZ 10	1	85	
Bifidobacterium adolescentis		3.90	0.35	NAG 90	1	99	
47111113	**Good identification**						**47111113**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium chauvoei		98.81	0.50	MLZ 1	99	99		
Capnocytophaga ochracea		1.06	0.14		1	1		
Actinomyces naeslundii		0.13	0.24		99	1		
47111120	**Low discrimination**							**47111120**
					GRA	CAT		
Prevotella melaninogenica		55.05	0.81	RAF 90	1	1		
Clostridium perfringens		39.59	0.85		99	1		
Capnocytophaga ochracea		3.67	0.64		1	90		
47111121	**Very good identification**							**47111121**
					GRA	SPO		
Clostridium chauvoei		98.62	0.89		99	99		
Prevotella melaninogenica		0.76	0.47		1	1		
Clostridium perfringens		0.55	0.51		99	84		
47111122	**Low discrimination**							**47111122**
					GRA	CAT		
Capnocytophaga ochracea		47.60	0.81	RAF 90	1	90		
Clostridium perfringens		26.82	0.80		99	1		
Prevotella melaninogenica		15.09	0.68		1	1		
47111123	**Good identification**							**47111123**
					GRA	CAT		
Clostridium chauvoei		94.73	0.76		99	1		
Capnocytophaga ochracea		2.48	0.47		1	90		
Clostridium perfringens		1.40	0.45		99	1		
47111130	**Low discrimination**							**47111130**
					GRA	CAT		
Prevotella melaninogenica		38.99	0.47	RAF 90	1	1		
Capnocytophaga ochracea		28.63	0.48	MLZ 1	1	90		
Clostridium perfringens		28.04	0.51		99	1		
47111131	**Good identification**							**47111131**
					GRA	CAT	SPO	
Clostridium chauvoei		98.08	0.55	MLZ 1	99	1	99	
Prevotella melaninogenica		0.76	0.12		1	1	1	
Capnocytophaga ochracea		0.56	0.14		1	90	1	
47111132	**Good identification**							**47111132**
					GRA	CAT		
Capnocytophaga ochracea		90.37	0.64	RAF 90	1	90		
Clostridium perfringens		4.63	0.45	MLZ 10	99	1		
Prevotella melaninogenica		2.61	0.34		1	1		
47111133	**Low discrimination**							**47111133**
					GRA	CAT		
Clostridium chauvoei		75.73	0.42	MLZ 1	99	1		
Capnocytophaga ochracea		21.84	0.30		1	90		
Clostridium perfringens		1.12	0.11		99	1		
47111140	**Low discrimination**							**47111140**
					GRA	CAT		
Clostridium perfringens		68.08	0.47	NAG 97	99	1		
Capnocytophaga ochracea		13.24	0.32		1	90		
Fusobacterium mortiferum		6.73	0.40		1	5		
47111141	**Very good identification**							**47111141**
					SPO			
Clostridium chauvoei		99.79	0.63	SOR 1	99			
Clostridium perfringens		0.11	0.13		84			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.08	0.33		1		
47111142	**Low discrimination**						**47111142**
					GRA	CAT	
Capnocytophaga ochracea		75.70	0.48	RAF 90	1	90	
Clostridium perfringens		20.35	0.41	NAG 90	99	1	
Clostridium chauvoei		0.72	0.16	SOR 10	99	1	
47111143	**Good identification**						**47111143**
					GRA	SPO	
Clostridium chauvoei		98.23	0.50	SOR 1	99	99	
Capnocytophaga ochracea		1.05	0.14		1	1	
Actinomyces naeslundii		0.44	0.33		99	1	
47111160	**Low discrimination**						**47111160**
					GRA	CAT	
Clostridium perfringens		88.40	0.73		99	1	
Prevotella melaninogenica		6.52	0.47		1	1	
Capnocytophaga ochracea		4.79	0.48		1	90	
47111161	**Low discrimination**						**47111161**
					GRA	SPO	
Clostridium chauvoei		89.27	0.55	SOR 1	99	99	
Clostridium perfringens		9.32	0.39		99	84	
Prevotella melaninogenica		0.69	0.12		1	1	
47111162	**Low discrimination**						**47111162**
					GRA	CAT	
Capnocytophaga ochracea		49.58	0.64	RAF 90	1	90	
Clostridium perfringens		47.88	0.67	SOR 10	99	1	
Prevotella melaninogenica		1.43	0.34		1	1	
47111163	**Low discrimination**						**47111163**
					GRA	CAT	
Clostridium chauvoei		62.75	0.42	SOR 1	99	1	
Capnocytophaga ochracea		18.09	0.30		1	90	
Clostridium perfringens		17.47	0.33		99	1	
47111170	**Low discrimination**						**47111170**
					GRA	CAT	
Clostridium perfringens		59.91	0.39	MLZ 1	99	1	
Capnocytophaga ochracea		35.68	0.32		1	90	
Prevotella melaninogenica		4.42	0.12		1	1	
47111172	**Acceptable identification**						**47111172**
					GRA	CAT	
Capnocytophaga ochracea		91.46	0.48	RAF 90	1	90	
Clostridium perfringens		8.03	0.33	MLZ 10	99	1	
Prevotella melaninogenica		0.24	0.00	SOR 10	1	1	
47111220	**Low discrimination**						**47111220**
					GRA		
Clostridium perfringens		74.03	0.36	LAC 88	99		
Prevotella oralis		12.51	0.20	RHA 1	1		
Prevotella melaninogenica		7.63	0.12		1		
47111222	**Low discrimination**						**47111222**
					GRA	CAT	
Prevotella oralis		84.68	0.41	RAF 92	1	1	
Clostridium perfringens		12.41	0.31	LAC 94	99	1	
Capnocytophaga ochracea		1.63	0.12		1	90	
47111301	**Very good identification**						**47111301**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium chauvoei		100.0	0.63	RHA 1				
47111302	**Low discrimination**							**47111302**
					GRA	CAT		
Prevotella oralis		70.29	0.40	RAF 92	1	1		
Capnocytophaga ochracea		18.07	0.30	NAG 95	1	90		
Clostridium perfringens		2.83	0.20		99	1		
47111303	**Good identification**							**47111303**
					GRA	SPO		
Clostridium chauvoei		99.62	0.50	RHA 1	99	99		
Prevotella oralis		0.38	0.06		1	1		
47111320	**Low discrimination**							**47111320**
					GRA			
Prevotella melaninogenica		46.26	0.47	RAF 90	1			
Clostridium perfringens		33.26	0.51	RHA 1	99			
Prevotella oralis		12.01	0.40		1			
47111321	**Good identification**							**47111321**
					GRA	SPO		
Clostridium chauvoei		98.44	0.55	RHA 1	99	99		
Prevotella melaninogenica		0.76	0.12		1	1		
Clostridium perfringens		0.55	0.17		99	84		
47111322	**Low discrimination**							**47111322**
					GRA	CAT		
Prevotella oralis		80.59	0.62	RAF 92	1	1		
Capnocytophaga ochracea		9.82	0.47		1	90		
Clostridium perfringens		5.53	0.45		99	1		
47111323	**Low discrimination**							**47111323**
					GRA	CAT	SPO	
Clostridium chauvoei		79.08	0.42	RHA 1	99	1	99	
Prevotella oralis		17.02	0.28		1	1	1	
Capnocytophaga ochracea		2.07	0.12		1	90	1	
47111360	**Low discrimination**							**47111360**
					GRA	CAT		
Clostridium perfringens		86.71	0.39	RHA 1	99	1		
Prevotella melaninogenica		6.39	0.12		1	1		
Capnocytophaga ochracea		4.70	0.14		1	90		
47111400	**Low discrimination**							**47111400**
					COC	GRA		
Gemella morbillorum		64.02	0.45	ESL 1	99	99		
Lactobacillus fermentum		27.43	0.50	ARA 10	1	99		
Fusobacterium mortiferum		3.78	0.34		1	1		
47111402	**Low discrimination**							**47111402**
					GRA	CAT		
Lactobacillus fermentum		82.62	0.50	ESL 1	99	1		
Capnocytophaga ochracea		7.56	0.17		1	90		
Bifidobacterium longum subsp. longum		2.58	0.18		99	1		
47111420	**Low discrimination**							**47111420**
					COC			
Clostridium perfringens		45.67	0.36	LAC 88	1			
Lactobacillus fermentum		19.99	0.42	ARA 1	1			
Gemella morbillorum		15.55	0.29		99			
47111500	**Low discrimination**							**47111500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Bifidobacterium longum subsp. longum		45.19	0.52	RAF 97	99	1		
Lactobacillus fermentum		23.84	0.54		99	1		
Capnocytophaga ochracea		14.70	0.35		1	90		
47111501	**Very good identification**							**47111501**
					SPO			
Clostridium chauvoei		99.81	0.63	ARA 1	99			
Bifidobacterium longum subsp. longum		0.10	0.18		1			
Lactobacillus fermentum		0.05	0.20		1			
47111502	**Low discrimination**							**47111502**
					GRA	CAT		
Capnocytophaga ochracea		58.39	0.51	RAF 90	1	90		
Bifidobacterium longum subsp. longum		19.94	0.52	NAG 90	99	1		
Lactobacillus fermentum		10.52	0.54		99	1		
47111503	**Good identification**							**47111503**
					GRA	SPO	CAT	
Clostridium chauvoei		97.23	0.50	ARA 1	99	99	1	
Capnocytophaga ochracea		1.65	0.17		1	1	90	
Bifidobacterium longum subsp. longum		0.56	0.18		99	1	1	
47111510	**Good identification**							**47111510**
					CAT	GRA		
Bifidobacterium longum subsp. longum		96.85	0.56	RAF 97	1	99		
Capnocytophaga ochracea		2.06	0.19		90	1		
Eubacterium saburreum		0.47	0.20		1	85		
47111512	**Low discrimination**							**47111512**
					GRA	CAT		
Bifidobacterium longum subsp. longum		65.23	0.56	RAF 97	99	1		
Bifidobacterium adolescentis		21.96	0.49		99	1		
Capnocytophaga ochracea		12.46	0.35		1	90		
47111520	**Low discrimination**							**47111520**
					GRA	CAT		
Bifidobacterium longum subsp. longum		33.87	0.62	RAF 97	99	1		
Capnocytophaga ochracea		24.79	0.51		1	90		
Prevotella melaninogenica		21.26	0.47		1	1		
47111521	**Good identification**							**47111521**
					GRA	SPO	CAT	
Clostridium chauvoei		96.58	0.55	ARA 1	99	99	1	
Bifidobacterium longum subsp. longum		1.19	0.28		99	1	1	
Capnocytophaga ochracea		0.87	0.17		1	1	90	
47111522	**Low discrimination**							**47111522**
					GRA	CAT		
Capnocytophaga ochracea		81.82	0.68	RAF 90	1	90		
Bifidobacterium longum subsp. longum		12.42	0.62		99	1		
Clostridium perfringens		2.64	0.45		99	1		
47111523	**Low discrimination**							**47111523**
					GRA	SPO	CAT	
Clostridium chauvoei		64.28	0.42	ARA 1	99	99	1	
Capnocytophaga ochracea		29.44	0.34		1	1	90	
Bifidobacterium longum subsp. longum		4.47	0.28		99	1	1	
47111530	**Good identification**							**47111530**
					GRA	CAT		
Bifidobacterium longum subsp. longum		94.82	0.66	RAF 97	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Capnocytophaga ochracea		4.53	0.35		1	90		
Prevotella melaninogenica		0.35	0.12		1	1		
47111532	**Low discrimination**							**47111532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		69.68	0.66	RAF 97	99	1		
Capnocytophaga ochracea		29.96	0.51		1	90		
Parabacteroides distasonis		0.10	0.21		1	76		
47111542	**Low discrimination**							**47111542**
					GRA	CAT		
Capnocytophaga ochracea		80.58	0.35	RAF 90	1	90		
Bifidobacterium adolescentis		14.36	0.32	NAG 90	99	1		
Bifidobacterium longum subsp. longum		2.50	0.18	SOR 10	99	1		
47111560	**Low discrimination**							**47111560**
					GRA	CAT		
Clostridium perfringens		46.59	0.39	ARA 1	99	1		
Capnocytophaga ochracea		44.07	0.35		1	90		
Bifidobacterium longum subsp. longum		5.47	0.28		99	1		
47111562	**Acceptable identification**							**47111562**
					GRA	CAT		
Capnocytophaga ochracea		93.31	0.51	RAF 90	1	90		
Clostridium perfringens		5.16	0.33	SOR 10	99	1		
Bifidobacterium longum subsp. longum		1.29	0.28		99	1		
47111572	**Low discrimination**							**47111572**
					GRA	CAT		
Capnocytophaga ochracea		82.55	0.35	RAF 90	1	90		
Bifidobacterium longum subsp. longum		17.46	0.32	MLZ 10	99	1		
47111720	**Good identification**							**47111720**
					GRA	CAT		
Bacteroides vulgatus		95.39	0.45	RAF 97	1	3		
Bifidobacterium longum subsp. longum		1.36	0.28	XYL 98	99	1		
Capnocytophaga ochracea		1.00	0.17		1	90		
47112000	**Low discrimination**							**47112000**
					COC	CAT		
Propionibacterium granulosum		39.08	0.48	ESL 1	1	90		
Eubacterium limosum		30.58	0.32		1	1		
Gemella morbillorum		11.39	0.32		99	1		
47112002	**Low discrimination**							**47112002**
					COC	CAT	GRA	
Propionibacterium granulosum		83.66	0.38	ESL 1	1	90	99	
Megasphaera elsdenii		8.94	0.15		99	1	1	
Eubacterium limosum		2.65	-.03		1	1	99	
47112020	**Low discrimination**							**47112020**
					COC	CAT		
Propionibacterium granulosum		84.47	0.38	ESL 1	1	90		
Gemella morbillorum		10.94	0.15		99	1		
Eubacterium limosum		2.67	-.03		1	1		
47112100	**Low discrimination**							**47112100**
					GRA	CAT		
Propionibacterium propionicum		70.61	0.46	NIT 86	99	1		
Fusobacterium mortiferum		25.60	0.37	ESL 1	1	5		
Propionibacterium granulosum		1.63	0.14		99	90		
47112102	**Good identification**							**47112102**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA			
Propionibacterium propionicum		96.60	0.49	NIT 86	99			
Fusobacterium mortiferum		2.60	0.21	ESL 1	1			
Actinomyces naeslundii		0.80	0.24		99			
47112140	**Good identification**							**47112140**
Propionibacterium propionicum		100.0	0.46	NIT 86				
47112142	**Good identification**							**47112142**
Propionibacterium propionicum		100.0	0.49	NIT 86				
47113000	**Low discrimination**							**47113000**
					COC	GRA	CAT	
Propionibacterium granulosum		71.37	0.48	ESL 1	1	99	90	
Gemella morbillorum		11.20	0.27		99	99	1	
Fusobacterium mortiferum		7.28	0.34		1	1	5	
47113002	**Low discrimination**							**47113002**
					CAT			
Propionibacterium granulosum		69.96	0.38	ESL 1	90			
Actinomyces naeslundii		12.88	0.40		25			
Lactobacillus jensenii		4.77	0.11		1			
47113003	**Low discrimination**							**47113003**
					SPO			
Actinomyces naeslundii		63.15	0.43	ESL 12	1			
Clostridium chauvoei		36.85	0.16	RAF 92	99			
47113020	**Low discrimination**							**47113020**
					GRA	CAT		
Clostridium perfringens		59.50	0.36	MAN 1	99	1		
Propionibacterium granulosum		26.42	0.38	LAC 88	99	90		
Prevotella melaninogenica		6.13	0.12		1	1		
47113023	**Low discrimination**							**47113023**
					SPO			
Actinomyces naeslundii		83.72	0.43	ESL 12	1			
Clostridium chauvoei		16.28	0.08	RAF 92	99			
47113100	**Low discrimination**							**47113100**
					GRA	SPO		
Propionibacterium propionicum		31.09	0.46	NIT 86	99	1		
Fusobacterium mortiferum		16.91	0.40	ESL 1	1	1		
Clostridium chauvoei		16.22	0.29		99	99		
47113101	**Very good identification**							**47113101**
					SPO			
Clostridium chauvoei		98.93	0.63	MAN 1	99			
Actinomyces naeslundii		0.86	0.51		1			
Propionibacterium propionicum		0.21	0.30		1			
47113102	**Low discrimination**							**47113102**
					GRA	CAT		
Propionibacterium propionicum		41.99	0.49	NIT 86	99	1		
Capnocytophaga ochracea		24.52	0.30	ESL 1	1	90		
Bifidobacterium adolescentis		8.39	0.32		99	1		
47113103	**Acceptable identification**							**47113103**
					SPO			
Clostridium chauvoei		93.67	0.50	MAN 1	99			
Actinomyces naeslundii		4.62	0.51		1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		1.71	0.33		1		
47113120	**Low discrimination**						**47113120**
					GRA	CAT	
Prevotella melaninogenica		54.35	0.47	RAF 90	1	1	
Clostridium perfringens		39.08	0.51	MAN 1	99	1	
Capnocytophaga ochracea		3.63	0.30		1	90	
47113121	**Good identification**						**47113121**
					GRA	SPO	
Clostridium chauvoei		96.22	0.55	MAN 1	99	99	
Actinomyces naeslundii		2.51	0.51		99	1	
Prevotella melaninogenica		0.74	0.12		1	1	
47113122	**Low discrimination**						**47113122**
					GRA	CAT	
Capnocytophaga ochracea		46.33	0.47	RAF 90	1	90	
Clostridium perfringens		26.10	0.45	MAN 1	99	1	
Prevotella melaninogenica		14.69	0.34		1	1	
47113123	**Low discrimination**						**47113123**
					GRA	SPO	
Clostridium chauvoei		83.18	0.42	MAN 1	99	99	
Actinomyces naeslundii		12.30	0.51		99	1	
Capnocytophaga ochracea		2.18	0.12		1	1	
47113140	**Low discrimination**						**47113140**
Propionibacterium propionicum		88.37	0.46	NIT 86			
Clostridium perfringens		4.91	0.13	ESL 1			
Actinomyces naeslundii		4.67	0.35				
47113142	**Low discrimination**						**47113142**
					GRA	CAT	
Propionibacterium propionicum		86.61	0.49	NIT 86	99	1	
Capnocytophaga ochracea		5.62	0.14	ESL 1	1	90	
Actinomyces naeslundii		3.05	0.35		99	25	
47113160	**Low discrimination**						**47113160**
					GRA	CAT	
Clostridium perfringens		85.99	0.39	MAN 1	99	1	
Prevotella melaninogenica		6.34	0.12		1	1	
Capnocytophaga ochracea		4.66	0.14		1	90	
47113163	**Low discrimination**						**47113163**
					SPO		
Actinomyces naeslundii		72.08	0.38	ESL 12	1		
Clostridium chauvoei		27.92	0.08	RAF 92	99		
47113502	**Low discrimination**						**47113502**
					GRA	CAT	
Bifidobacterium adolescentis		87.38	0.46	RAF 92	99	1	
Capnocytophaga ochracea		6.74	0.17	XYL 87	1	90	
Bifidobacterium longum subsp. longum		2.30	0.18		99	1	
47113512	**Good identification**						**47113512**
					GRA	CAT	
Bifidobacterium adolescentis		95.82	0.48	RAF 92	99	1	
Bifidobacterium longum subsp. longum		3.51	0.22	XYL 87	99	1	
Capnocytophaga ochracea		0.67	0.01		1	90	
47114000	**Low discrimination**						**47114000**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		83.27	0.50	ESL 1	99	99		
Fusobacterium mortiferum		13.16	0.46	XYL 10	1	1		
Actinomyces meyeri		1.11	0.18		1	99		
47114100	**Low discrimination**							**47114100**
					COC	GRA		
Fusobacterium mortiferum		77.51	0.52	MLT 11	1	1		
Actinomyces meyeri		11.17	0.29	XYL 7	1	99		
Atopobium parvulum		7.22	0.23		99	99		
47114102	**Low discrimination**							**47114102**
					COC	CAT	GRA	
Fusobacterium mortiferum		43.05	0.36	MLT 11	1	5	1	
Capnocytophaga ochracea		31.40	0.20	XYL 7	1	90	1	
Atopobium parvulum		6.37	0.10	BGL 10	99	1	99	
47114522	**Low discrimination**							**47114522**
					GRA	CAT		
Bifidobacterium longum subsp. longum		61.27	0.36	RAF 97	99	1		
Capnocytophaga ochracea		33.64	0.23	GAL 99	1	90		
Bacteroides fragilis		3.64	-.03		1	99		
47114530	**Good identification**							**47114530**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97				
47114532	**Acceptable identification**							**47114532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		90.98	0.40	RAF 97	99	1		
Clostridium clostridioforme		5.76	0.18	GAL 99	50	33		
Capnocytophaga ochracea		3.26	0.07		1	90		
47114720	**Good identification**							**47114720**
					GRA			
Bacteroides vulgatus		98.81	0.40	RAF 97	1			
Eubacterium contortum		1.19	0.14	GAL 99	99			
47115000	**Low discrimination**							**47115000**
					COC	GRA		
Gemella morbillorum		46.45	0.45	ESL 1	99	99		
Fusobacterium mortiferum		20.45	0.49	XYL 10	1	1		
Eubacterium saburreum		14.11	0.45		1	85		
47115002	**Low discrimination**							**47115002**
					CAT	GRA		
Lactobacillus fermentum		50.89	0.46	ESL 1	1	99		
Capnocytophaga ochracea		11.23	0.20		90	1		
Eubacterium saburreum		10.84	0.32		1	85		
47115010	**Acceptable identification**							**47115010**
					COC			
Eubacterium saburreum		91.86	0.43	IND 99	1			
Gemella morbillorum		3.73	0.11		99			
Lactobacillus jensenii		1.83	0.06		1			
47115020	**Low discrimination**							**47115020**
					COC			
Clostridium perfringens		42.39	0.36	XYL 1	1			
Gemella morbillorum		14.44	0.29	LAC 88	99			
Lactobacillus fermentum		10.90	0.38		1			
47115100	**Low discrimination**							**47115100**
					COC	GRA		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		63.90	0.57	TRE 99	99	99		
Fusobacterium mortiferum		10.39	0.55		1	1		
Eubacterium saburreum		7.17	0.51		1	85		
47115101	**Very good identification**							**47115101**
					COC	GRA	SPO	
Clostridium chauvoei		99.31	0.63	XYL 1	1	99	99	
Atopobium parvulum		0.48	0.23		99	99	1	
Fusobacterium mortiferum		0.08	0.21		1	1	1	
47115102	**Low discrimination**							**47115102**
					COC	GRA	CAT	
Capnocytophaga ochracea		53.26	0.54	RAF 90	1	1	90	
Atopobium parvulum		10.81	0.45	NAG 90	99	99	1	
Bifidobacterium longum subsp. infantis		10.46	0.38		1	99	1	
47115103	**Good identification**							**47115103**
					COC	GRA	SPO	
Clostridium chauvoei		95.95	0.50	XYL 1	1	99	99	
Capnocytophaga ochracea		2.31	0.20		1	1	1	
Atopobium parvulum		0.47	0.10		99	99	1	
47115110	**Low discrimination**							**47115110**
					CAT	GRA		
Eubacterium saburreum		72.29	0.49	IND 99	1	85		
Bifidobacterium longum subsp. longum		8.52	0.30		1	99		
Capnocytophaga ochracea		8.46	0.21		90	1		
47115112	**Low discrimination**							**47115112**
					CAT	GRA		
Bifidobacterium adolescentis		47.96	0.49	RAF 92	1	99		
Capnocytophaga ochracea		38.56	0.38	ARA 87	90	1		
Eubacterium saburreum		6.46	0.37		1	85		
47115120	**Low discrimination**							**47115120**
					CAT			
Capnocytophaga ochracea		27.79	0.54	RAF 90	90			
Bacteroides fragilis		24.89	0.44	BGL 90	99			
Prevotella melaninogenica		16.82	0.47		1			
47115121	**Good identification**							**47115121**
					GRA	SPO		
Clostridium chauvoei		95.83	0.55	XYL 1	99	99		
Capnocytophaga ochracea		1.22	0.20		1	1		
Bacteroides fragilis		1.09	0.10		1	1		
47115122	**Low discrimination**							**47115122**
					CAT			
Bacteroides fragilis		61.41	0.66	RAF 99	99			
Capnocytophaga ochracea		32.48	0.70		90			
Prevotella oralis		3.11	0.53		1			
47115130	**Low discrimination**							**47115130**
					CAT	GRA		
Capnocytophaga ochracea		54.49	0.38	RAF 90	90	1		
Bifidobacterium longum subsp. longum		24.40	0.40	MLZ 10	1	99		
Eubacterium saburreum		9.13	0.37	BGL 90	1	85		
47115132	**Low discrimination**							**47115132**
					GRA	CAT		
Capnocytophaga ochracea		78.38	0.54	RAF 90	1	90		
Bacteroides fragilis		13.47	0.32	MLZ 10	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		3.90	0.40		99	1		
47115142	**Low discrimination**							**47115142**
					COC	GRA	CAT	
Capnocytophaga ochracea		82.57	0.38	RAF 90	1	1	90	
Bifidobacterium adolescentis		10.39	0.32	NAG 90	1	99	1	
Atopobium parvulum		1.52	0.10	SOR 10	99	99	1	
47115160	**Low discrimination**							**47115160**
					GRA	CAT		
Capnocytophaga ochracea		51.91	0.38	RAF 90	1	90		
Clostridium perfringens		38.73	0.39	SOR 10	99	1		
Bacteroides fragilis		4.23	0.10	BGL 90	1	99		
47115162	**Low discrimination**							**47115162**
					GRA	CAT		
Capnocytophaga ochracea		81.64	0.54	RAF 90	1	90		
Bacteroides fragilis		14.03	0.32	SOR 10	1	99		
Clostridium perfringens		3.19	0.33		99	1		
47115172	**Good identification**							**47115172**
					GRA	CAT		
Capnocytophaga ochracea		98.03	0.38	RAF 90	1	90		
Bacteroides fragilis		1.53	-.03	MLZ 10	1	99		
Bifidobacterium longum subsp. longum		0.44	0.06	SOR 10	99	1		
47115300	**Very good identification**							**47115300**
					COC	GRA		
Eubacterium contortum		98.47	0.63	SAL 99	1	99		
Bacteroides vulgatus		0.90	0.21		1	1		
Atopobium parvulum		0.41	0.23		99	99		
47115302	**Acceptable identification**							**47115302**
					GRA	CAT		
Eubacterium contortum		92.45	0.50	SAL 99	99	1		
Prevotella oralis		3.66	0.24		1	1		
Capnocytophaga ochracea		1.90	0.20		1	90		
47115320	**Low discrimination**							**47115320**
					GRA			
Bacteroides vulgatus		82.25	0.55	RAF 97	1			
Eubacterium contortum		16.02	0.50	ARA 93	99			
Prevotella oralis		0.63	0.24		1			
47115322	**Low discrimination**							**47115322**
					CAT			
Prevotella oralis		39.32	0.46	RAF 92	1			
Parabacteroides distasonis		19.92	0.54	XYL 11	76			
Bacteroides fragilis		18.31	0.32		99			
47115332	**Low discrimination**							**47115332**
Parabacteroides distasonis		82.49	0.49	RAF 99				
Capnocytophaga ochracea		8.28	0.20					
Clostridium clostridioforme		3.63	0.21					
47115400	**Low discrimination**							**47115400**
					COC			
Lactobacillus fermentum		59.48	0.46	ESL 1	1			
Gemella morbillorum		26.27	0.29		99			
Bifidobacterium longum subsp. longum		9.49	0.26		1			
47115402	**Low discrimination**							**47115402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus fermentum		71.48	0.46	ESL 1			
Bifidobacterium adolescentis		11.93	0.28				
Bifidobacterium longum subsp. longum		11.41	0.26				
47115422	**Low discrimination**						**47115422**
					GRA	CAT	
Bifidobacterium longum subsp. longum		46.06	0.36	RAF 97	99	1	
Capnocytophaga ochracea		25.29	0.23	LAC 99	1	90	
Lactobacillus fermentum		24.06	0.38		99	1	
47115430	**Good identification**						**47115430**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97			
47115432	**Acceptable identification**						**47115432**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.53	0.40	RAF 97	99	1	
Clostridium clostridioforme		5.19	0.17	LAC 99	50	33	
Capnocytophaga ochracea		3.28	0.07		1	90	
47115500	**Low discrimination**						**47115500**
					GRA		
Bifidobacterium longum subsp. longum		79.50	0.60	RAF 97	99		
Lactobacillus fermentum		8.21	0.50		99		
Bacteroides vulgatus		4.13	0.24		1		
47115502	**Low discrimination**						**47115502**
					GRA	CAT	
Bifidobacterium adolescentis		43.32	0.62	RAF 92	99	1	
Bifidobacterium longum subsp. longum		41.41	0.60		99	1	
Capnocytophaga ochracea		10.10	0.41		1	90	
47115510	**Very good identification**						**47115510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.93	0.64	RAF 97	99	1	
Bifidobacterium adolescentis		0.75	0.29		99	1	
Capnocytophaga ochracea		0.18	0.08		1	90	
47115512	**Low discrimination**						**47115512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		56.32	0.64	RAF 97	99	1	
Bifidobacterium adolescentis		42.29	0.63		99	1	
Capnocytophaga ochracea		0.90	0.25		1	90	
47115520	**Low discrimination**						**47115520**
					GRA	CAT	
Bacteroides vulgatus		54.37	0.58	RAF 97	1	3	
Bifidobacterium longum subsp. longum		42.30	0.70	RHA 90	99	1	
Capnocytophaga ochracea		2.58	0.41		1	90	
47115522	**Low discrimination**						**47115522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		60.53	0.70	RAF 97	99	1	
Capnocytophaga ochracea		33.23	0.58		1	90	
Bacteroides fragilis		3.60	0.32		1	99	
47115523	**Low discrimination**						**47115523**
					GRA	CAT	
Bifidobacterium longum subsp. longum		60.36	0.36	RAF 97	99	1	
Capnocytophaga ochracea		33.14	0.23	URE 1	1	90	
Bacteroides fragilis		3.59	-.03		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47115530	**Very good identification**						**47115530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.85	0.74	RAF 97	99	1	
Bacteroides vulgatus		0.75	0.24		1	3	
Capnocytophaga ochracea		0.39	0.25		1	90	
47115531	**Good identification**						**47115531**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97			
47115532	**Very good identification**						**47115532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		95.77	0.74	RAF 97	99	1	
Capnocytophaga ochracea		3.43	0.41		1	90	
Parabacteroides distasonis		0.40	0.38		1	76	
47115533	**Good identification**						**47115533**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.54	0.40	RAF 97	99	1	
Capnocytophaga ochracea		3.46	0.07	URE 1	1	90	
47115542	**Low discrimination**						**47115542**
					GRA	CAT	
Bifidobacterium adolescentis		76.27	0.46	RAF 92	99	1	
Capnocytophaga ochracea		15.99	0.25	SOR 11	1	90	
Bifidobacterium longum subsp. longum		5.96	0.26		99	1	
47115552	**Low discrimination**						**47115552**
					GRA	CAT	
Bifidobacterium adolescentis		88.66	0.48	RAF 92	99	1	
Bifidobacterium longum subsp. longum		9.65	0.30	SOR 11	99	1	
Capnocytophaga ochracea		1.69	0.08		1	90	
47115562	**Low discrimination**						**47115562**
					GRA	CAT	
Capnocytophaga ochracea		84.69	0.41	RAF 90	1	90	
Bifidobacterium longum subsp. longum		14.02	0.36	SOR 10	99	1	
Bacteroides fragilis		0.83	-.03		1	99	
47115570	**Good identification**						**47115570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		95.81	0.40	RAF 97	99	1	
Capnocytophaga ochracea		4.20	0.08	SOR 1	1	90	
47115572	**Low discrimination**						**47115572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		70.56	0.40	RAF 97	99	1	
Capnocytophaga ochracea		27.81	0.25	SOR 1	1	90	
Bifidobacterium adolescentis		1.64	0.14		99	1	
47115620	**Good identification**						**47115620**
					GRA		
Bacteroides vulgatus		98.81	0.40	RAF 97	1		
Eubacterium contortum		1.19	0.14	LAC 99	99		
47115700	**Low discrimination**						**47115700**
					GRA		
Eubacterium contortum		86.80	0.61	SAL 99	99		
Bacteroides vulgatus		12.90	0.40		1		
Bifidobacterium longum subsp. longum		0.28	0.26		99		
47115702	**Acceptable identification**						**47115702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium contortum		91.51	0.48	SAL 99			
Clostridium clostridioforme		3.47	0.30				
Bifidobacterium adolescentis		1.74	0.28				
47115712	**Low discrimination**						**47115712**
Clostridium clostridioforme		72.37	0.41	SAL 88			
Bifidobacterium longum subsp. longum		12.99	0.30	RAF 94			
Bifidobacterium adolescentis		9.76	0.29	TRE 88			
47115720	**Very good identification**						**47115720**
					GRA		
Bacteroides vulgatus		98.72	0.74	RAF 97	1		
Eubacterium contortum		1.18	0.48		99		
Bifidobacterium longum subsp. longum		0.09	0.36		99		
47115721	**Good identification**						**47115721**
					GRA		
Bacteroides vulgatus		98.81	0.40	RAF 97	1		
Eubacterium contortum		1.19	0.14	URE 1	99		
47115722	**Low discrimination**						**47115722**
					GRA		
Bacteroides vulgatus		57.02	0.40	RAF 97	1		
Parabacteroides distasonis		19.33	0.50	BGL 1	1		
Eubacterium contortum		11.95	0.36		99		
47115730	**Low discrimination**						**47115730**
					GRA		
Bacteroides vulgatus		85.98	0.40	RAF 97	1		
Bifidobacterium longum subsp. longum		12.66	0.40	MLZ 1	99		
Eubacterium contortum		1.03	0.14		99		
47115732	**Low discrimination**						**47115732**
					GRA		
Clostridium clostridioforme		36.91	0.41	SAL 88	50		
Parabacteroides distasonis		32.86	0.46	RAF 94	1		
Bifidobacterium longum subsp. longum		26.50	0.40	TRE 88	99		
47115760	**Good identification**						**47115760**
					GRA		
Bacteroides vulgatus		98.81	0.40	RAF 97	1		
Eubacterium contortum		1.19	0.14	SOR 1	99		
47117100	**Low discrimination**						**47117100**
					COC	GRA	
Actinomyces israelii		48.39	0.40	SAL 92	1	99	
Atopobium parvulum		33.15	0.23	CEL 93	99	99	
Fusobacterium mortiferum		5.39	0.21		1	1	
47117102	**Low discrimination**						**47117102**
					GRA	CAT	
Bifidobacterium adolescentis		67.99	0.46	RAF 92	99	1	
Actinomyces israelii		18.69	0.43	ARA 87	99	1	
Capnocytophaga ochracea		7.42	0.20		1	90	
47117112	**Acceptable identification**						**47117112**
					SPO		
Bifidobacterium adolescentis		94.68	0.48	RAF 92	1		
Actinomyces israelii		3.18	0.29	ARA 87	1		
Clostridium tertium		1.20	0.07		99		
47117500	**Low discrimination**						**47117500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		49.22	0.38	SAL 92			
Bifidobacterium longum subsp. longum		25.92	0.26	CEL 93			
Bifidobacterium adolescentis		22.19	0.26				
47117502	**Very good identification**						**47117502**
Bifidobacterium adolescentis		95.28	0.60	RAF 92			
Actinomyces israelii		3.20	0.41				
Bifidobacterium longum subsp. longum		1.12	0.26				
47117512	**Very good identification**						**47117512**
Bifidobacterium adolescentis		97.99	0.62	RAF 92			
Bifidobacterium longum subsp. longum		1.61	0.30				
Actinomyces israelii		0.40	0.27				
47117522	**Low discrimination**						**47117522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		53.54	0.36	RAF 97	99	1	
Capnocytophaga ochracea		29.39	0.23	MAN 1	1	90	
Bifidobacterium adolescentis		11.46	0.26		99	1	
47117530	**Good identification**						**47117530**
Bifidobacterium longum subsp. longum		100.0	0.40	RAF 97			
47117532	**Low discrimination**						**47117532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		84.07	0.40	RAF 97	99	1	
Bifidobacterium adolescentis		12.92	0.28	MAN 1	99	1	
Capnocytophaga ochracea		3.01	0.07		1	90	
47117542	**Acceptable identification**						**47117542**
Bifidobacterium adolescentis		90.44	0.45	RAF 92			
Actinomyces israelii		9.56	0.34	SOR 11			
47117552	**Good identification**						**47117552**
Bifidobacterium adolescentis		98.72	0.46	RAF 92			
Actinomyces israelii		1.28	0.20	SOR 11			
47117720	**Good identification**						**47117720**
					GRA		
Bacteroides vulgatus		98.81	0.40	RAF 97	1		
Eubacterium contortum		1.19	0.14	MAN 1	99		
47120000	**Low discrimination**						**47120000**
					COC	SPO	
Clostridium botulinum biovar A		87.25	0.50	SUC 7	1	99	
Gemella morbillorum		4.28	0.30	MNS 7	99	1	
Clostridium sporogenes		3.50	0.24		1	99	
47120040	**Acceptable identification**						**47120040**
Clostridium botulinum biovar B		91.42	0.56	ESL 1			
Clostridium botulinum biovar A		8.59	0.41				
47120050	**Good identification**						**47120050**
Clostridium botulinum biovar B		99.81	0.50	ESL 1			
Clostridium botulinum biovar A		0.19	0.07				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47120060	**Good identification**						**47120060**
Clostridium botulinum biovar B		99.81	0.50	ESL 1			
Clostridium botulinum biovar A		0.19	0.07				
47120070	**Good identification**						**47120070**
Clostridium botulinum biovar B		100.0	0.45	ESL 1			
47120101	**Good identification**						**47120101**
Clostridium chauvoei		100.0	0.40	TRE 10			
47121000	**Low discrimination**						**47121000**
					COC		
Eubacterium saburreum		33.61	0.38	IND 99	1		
Gemella morbillorum		19.95	0.26		99		
Lactobacillus jensenii		11.91	0.22		1		
47121001	**Good identification**						**47121001**
Clostridium chauvoei		100.0	0.40	TRE 10			
47121010	**Low discrimination**						**47121010**
					CAT		
Eubacterium saburreum		85.08	0.37	IND 99	1		
Lactobacillus jensenii		12.29	0.14		1		
Propionibacterium granulosum		2.63	0.09		90		
47121020	**Acceptable identification**						**47121020**
Clostridium perfringens		91.39	0.41	FRU 99			
Eubacterium saburreum		3.34	0.25	LAC 88			
Lactobacillus jensenii		1.68	0.12				
47121022	**Good identification**						**47121022**
Clostridium perfringens		96.19	0.35	FRU 99			
Lactobacillus jensenii		2.50	0.09	LAC 88			
Eubacterium saburreum		1.32	0.12				
47121100	**Low discrimination**						**47121100**
					COC	SPO	
Atopobium parvulum		75.53	0.53	FRU 99	99	1	
Clostridium chauvoei		13.84	0.40		1	99	
Eubacterium saburreum		6.49	0.44		1	1	
47121101	**Very good identification**						**47121101**
					COC	SPO	
Clostridium chauvoei		99.93	0.74	TRE 10	1	99	
Atopobium parvulum		0.06	0.19		99	1	
Actinomyces naeslundii		0.01	0.29		1	1	
47121102	**Low discrimination**						**47121102**
					COC	SPO	
Atopobium parvulum		71.89	0.40	FRU 99	99	1	
Clostridium chauvoei		13.17	0.27		1	99	
Clostridium perfringens		7.74	0.24		1	84	
47121103	**Very good identification**						**47121103**
					COC	SPO	
Clostridium chauvoei		99.89	0.62	TRE 10	1	99	
Atopobium parvulum		0.06	0.06		99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.06	0.29		1	1	
47121110	**Low discrimination**						**47121110**
					COC	SPO	
Eubacterium saburreum		85.47	0.43	IND 99	1	1	
Atopobium parvulum		12.28	0.19		99	1	
Clostridium chauvoei		2.25	0.06		1	99	
47121111	**Good identification**						**47121111**
Clostridium chauvoei		100.0	0.40	TRE 10			
47121120	**Low discrimination**						**47121120**
					COC	GRA	
Clostridium perfringens		77.79	0.56	FRU 99	1	99	
Atopobium parvulum		10.51	0.40		99	99	
Prevotella melaninogenica		6.47	0.30		1	1	
47121121	**Very good identification**						**47121121**
					SPO		
Clostridium chauvoei		99.71	0.66	TRE 10	99		
Clostridium perfringens		0.22	0.21		84		
Actinomyces naeslundii		0.03	0.29		1		
47121122	**Low discrimination**						**47121122**
					COC	GRA	
Clostridium perfringens		88.53	0.50	FRU 99	1	99	
Atopobium parvulum		4.49	0.27		99	99	
Prevotella melaninogenica		2.98	0.18		1	1	
47121123	**Good identification**						**47121123**
					SPO		
Clostridium chauvoei		99.26	0.53	TRE 10	99		
Clostridium perfringens		0.58	0.16		84		
Actinomyces naeslundii		0.16	0.29		1		
47121141	**Good identification**						**47121141**
Clostridium chauvoei		100.0	0.40	TRE 10			
47121160	**Good identification**						**47121160**
					COC	GRA	
Clostridium perfringens		98.86	0.43	FRU 99	1	99	
Atopobium parvulum		0.71	0.06		99	99	
Prevotella melaninogenica		0.44	-.04		1	1	
47121162	**Good identification**						**47121162**
Clostridium perfringens		100.0	0.38	FRU 99			
47121301	**Good identification**						**47121301**
Clostridium chauvoei		100.0	0.40	TRE 10			
47121501	**Good identification**						**47121501**
Clostridium chauvoei		100.0	0.40	TRE 10			
47121520	**Low discrimination**						**47121520**
					COC		
Bifidobacterium longum subsp. longum		72.92	0.35	RAF 97	1		
Clostridium perfringens		22.23	0.21	FRU 90	1		
Atopobium parvulum		3.01	0.06		99		
47121522	**Low discrimination**						**47121522**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Bifidobacterium longum subsp. longum		85.64	0.35	RAF 97	99	1	
Clostridium perfringens		12.29	0.16	FRU 90	99	1	
Capnocytophaga ochracea		2.07	-.01		1	90	
47121530	**Good identification**						**47121530**
Bifidobacterium longum subsp. longum		100.0	0.39	RAF 97			
47121532	**Good identification**						**47121532**
Bifidobacterium longum subsp. longum		100.0	0.39	RAF 97			
47123101	**Good identification**						**47123101**
					SPO		
Clostridium chauvoei		98.17	0.40	TRE 10	99		
Actinomyces naeslundii		1.83	0.34	MAN 1	1		
47125100	**Very good identification**						**47125100**
					COC	SPO	
Atopobium parvulum		97.43	0.57	FRU 99	99	1	
Eubacterium saburreum		2.43	0.40		1	1	
Clostridium chauvoei		0.10	0.06		1	99	
47125101	**Acceptable identification**						**47125101**
					COC	SPO	
Clostridium chauvoei		90.71	0.40	TRE 10	1	99	
Atopobium parvulum		9.29	0.23	XYL 1	99	1	
47125102	**Good identification**						**47125102**
					COC		
Atopobium parvulum		97.21	0.45	FRU 99	99		
Eubacterium saburreum		2.42	0.27		1		
Bifidobacterium adolescentis		0.37	0.11		1		
47125110	**Low discrimination**						**47125110**
					COC		
Eubacterium saburreum		66.87	0.38	IND 99	1		
Atopobium parvulum		33.13	0.23		99		
47125120	**Acceptable identification**						**47125120**
					COC		
Atopobium parvulum		93.24	0.45	FRU 99	99		
Clostridium perfringens		3.75	0.21		1		
Eubacterium saburreum		2.32	0.27		1		
47125510	**Good identification**						**47125510**
Bifidobacterium longum subsp. longum		100.0	0.37	RAF 97			
47125512	**Low discrimination**						**47125512**
Bifidobacterium longum subsp. longum		83.16	0.37	RAF 97			
Bifidobacterium adolescentis		15.14	0.26	FRU 90			
Clostridium clostridioforme		1.70	0.07				
47125520	**Good identification**						**47125520**
					COC	GRA	
Bifidobacterium longum subsp. longum		95.70	0.44	RAF 97	1	99	
Atopobium parvulum		2.44	0.10	FRU 90	99	99	
Bacteroides vulgatus		1.86	0.00		1	1	
47125522	**Good identification**						**47125522**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		97.86	0.44	RAF 97	99			
Parabacteroides distasonis		2.14	0.19	FRU 90	1			
47125530	**Good identification**							**47125530**
Bifidobacterium longum subsp. longum		100.0	0.48	RAF 97				
47125532	**Good identification**							**47125532**
					GRA			
Bifidobacterium longum subsp. longum		98.81	0.48	RAF 97	99			
Parabacteroides distasonis		0.68	0.15	FRU 90	1			
Clostridium clostridioforme		0.51	0.07		50			
47130000	**Low discrimination**							**47130000**
					GRA	SPO	CAT	
Propionibacterium granulosum		53.33	0.58	ESL 1	99	1	90	
Fusobacterium mortiferum		26.97	0.56		1	1	5	
Clostridium botulinum biovar B		8.38	0.32		99	99	1	
47130002	**Low discrimination**							**47130002**
					GRA	CAT		
Propionibacterium granulosum		75.76	0.48	ESL 1	99	90		
Fusobacterium mortiferum		17.03	0.40		1	5		
Lactobacillus jensenii		3.90	0.19		99	1		
47130010	**Low discrimination**							**47130010**
					SPO	CAT		
Propionibacterium granulosum		59.82	0.43	ESL 1	1	90		
Clostridium botulinum biovar B		33.96	0.26	MLZ 12	99	1		
Lactobacillus jensenii		2.82	0.14		1	1		
47130020	**Low discrimination**							**47130020**
					CAT	SPO		
Clostridium perfringens		43.03	0.41	GAL 99	1	84		
Propionibacterium granulosum		40.91	0.48	LAC 88	90	1		
Clostridium botulinum biovar B		12.67	0.26		1	99		
47130022	**Low discrimination**							**47130022**
					CAT			
Clostridium perfringens		65.32	0.35	GAL 99	1			
Propionibacterium granulosum		32.99	0.38	LAC 88	90			
Lactobacillus jensenii		1.70	0.09		1			
47130040	**Very good identification**							**47130040**
					SPO	CAT		
Clostridium botulinum biovar B		99.74	0.66	ESL 1	99	1		
Clostridium botulinum biovar A		0.15	0.20		99	1		
Propionibacterium granulosum		0.07	0.24		1	90		
47130050	**Very good identification**							**47130050**
					SPO	CAT		
Clostridium botulinum biovar B		99.98	0.61	ESL 1	99	1		
Propionibacterium granulosum		0.02	0.09		1	90		
47130060	**Very good identification**							**47130060**
					CAT	SPO		
Clostridium botulinum biovar B		99.32	0.61	ESL 1	1	99		
Clostridium perfringens		0.65	0.28		1	84		
Propionibacterium granulosum		0.03	0.14		90	1		
47130070	**Good identification**							**47130070**
Clostridium botulinum biovar B		100.0	0.55	ESL 1				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
47130100	**Low discrimination**							**47130100**
					COC	GRA		
Atopobium parvulum		53.91	0.53	GAL 99	99	99		
Fusobacterium mortiferum		43.07	0.63		1	1		
Clostridium perfringens		2.18	0.30		1	99		
47130101	**Good identification**							**47130101**
					COC	GRA	SPO	
Clostridium chauvoei		95.62	0.47	TRE 10	1	99	99	
Atopobium parvulum		2.26	0.19	GAL 99	99	99	1	
Fusobacterium mortiferum		1.81	0.28		1	1	1	
47130102	**Low discrimination**							**47130102**
					COC	GRA		
Atopobium parvulum		61.15	0.40	GAL 99	99	99		
Fusobacterium mortiferum		30.76	0.46		1	1		
Clostridium perfringens		6.59	0.24		1	99		
47130120	**Low discrimination**							**47130120**
					COC	SPO		
Clostridium perfringens		86.21	0.56	GAL 99	1	84		
Atopobium parvulum		11.65	0.40		99	1		
Clostridium septicum		0.68	0.13		1	99		
47130121	**Low discrimination**							**47130121**
					COC	SPO		
Clostridium chauvoei		89.70	0.38	TRE 10	1	99		
Clostridium perfringens		8.30	0.21	GAL 99	1	84		
Atopobium parvulum		1.12	0.06		99	1		
47130122	**Acceptable identification**							**47130122**
					COC	GRA		
Clostridium perfringens		92.75	0.50	GAL 99	1	99		
Atopobium parvulum		4.70	0.27		99	99		
Prevotella oralis		1.10	0.14		1	1		
47130160	**Low discrimination**							**47130160**
					COC	SPO		
Clostridium perfringens		82.11	0.43	GAL 99	1	84		
Clostridium botulinum biovar B		17.31	0.26		1	99		
Atopobium parvulum		0.59	0.06		99	1		
47130162	**Good identification**							**47130162**
Clostridium perfringens		100.0	0.38	GAL 99				
47131000	**Low discrimination**							**47131000**
					COC	CAT		
Lactobacillus jensenii		30.90	0.56	SAL 95	1	1		
Atopobium parvulum		23.90	0.53	CEL 95	99	1		
Propionibacterium granulosum		16.18	0.58		1	90		
47131001	**Low discrimination**							**47131001**
					SPO			
Clostridium chauvoei		88.87	0.47	TRE 10	99			
Lactobacillus jensenii		2.71	0.22	LAC 99	1			
Actinomyces naeslundii		2.44	0.43		1			
47131002	**Low discrimination**							**47131002**
					COC			
Lactobacillus jensenii		55.37	0.53	SAL 95	1			
Clostridium perfringens		16.48	0.43	CEL 95	1			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Atopobium parvulum		11.34	0.40		99	
47131010	**Low discrimination**					**47131010**
					CAT	
Lactobacillus jensenii		67.92	0.48	SAL 95	1	
Propionibacterium granulosum		14.55	0.43	CEL 95	90	
Eubacterium saburreum		14.25	0.45		1	
47131012	**Low discrimination**					**47131012**
					CAT	
Lactobacillus jensenii		86.18	0.45	SAL 95	1	
Propionibacterium granulosum		6.93	0.33	CEL 95	90	
Eubacterium saburreum		4.79	0.32		1	
47131020	**Very good identification**					**47131020**
					COC	
Clostridium perfringens		96.03	0.75	LAC 88	1	
Lactobacillus jensenii		1.76	0.46		1	
Atopobium parvulum		0.96	0.40		99	
47131021	**Low discrimination**					**47131021**
					SPO	
Clostridium perfringens		52.51	0.41	LAC 88	84	
Clostridium chauvoei		42.03	0.38	URE 1	99	
Actinomyces naeslundii		3.47	0.43		1	
47131022	**Very good identification**					**47131022**
					GRA	
Clostridium perfringens		95.85	0.69	LAC 88	99	
Lactobacillus jensenii		2.49	0.43		99	
Prevotella oralis		0.53	0.27		1	
47131023	**Low discrimination**					**47131023**
					SPO	
Clostridium perfringens		68.20	0.35	LAC 88	84	
Clostridium chauvoei		20.47	0.25	URE 1	99	
Actinomyces naeslundii		9.57	0.43		1	
47131030	**Low discrimination**					**47131030**
					CAT	
Clostridium perfringens		54.52	0.41	LAC 88	1	
Lactobacillus jensenii		32.99	0.38	MLZ 1	1	
Propionibacterium granulosum		7.07	0.33		90	
47131032	**Low discrimination**					**47131032**
					CAT	
Clostridium perfringens		51.03	0.35	LAC 88	1	
Lactobacillus jensenii		43.74	0.35	MLZ 1	1	
Propionibacterium granulosum		3.51	0.23		90	
47131060	**Very good identification**					**47131060**
					SPO	
Clostridium perfringens		98.23	0.63	LAC 88	84	
Clostridium botulinum biovar B		1.53	0.26		99	
Lactobacillus jensenii		0.10	0.12		1	
47131062	**Very good identification**					**47131062**
Clostridium perfringens		99.78	0.57	LAC 88		
Lactobacillus jensenii		0.14	0.09			
Actinomyces naeslundii		0.09	0.27			
47131100	**Good identification**					**47131100**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				COC	GRA	
Atopobium parvulum	94.32	0.87		99	99	
Clostridium perfringens	3.81	0.64		1	99	
Fusobacterium mortiferum	1.14	0.66		1	1	
47131101 Very good identification						**47131101**
				COC	SPO	
Clostridium chauvoei	97.49	0.81	TRE 10	1	99	
Atopobium parvulum	2.30	0.53		99	1	
Clostridium perfringens	0.09	0.30		1	84	
47131102 Low discrimination						**47131102**
				COC	GRA	
Atopobium parvulum	88.20	0.74		99	99	
Clostridium perfringens	9.50	0.58		1	99	
Fusobacterium mortiferum	0.67	0.49		1	1	
47131103 Very good identification						**47131103**
				COC	SPO	
Clostridium chauvoei	97.01	0.68	TRE 10	1	99	
Atopobium parvulum	2.29	0.40		99	1	
Actinomyces naeslundii	0.43	0.51		1	1	
47131110 Low discrimination						**47131110**
				COC		
Atopobium parvulum	78.17	0.53	MLZ 1	99		
Eubacterium saburreum	16.49	0.51		1		
Clostridium perfringens	3.16	0.30		1		
47131111 Good identification						**47131111**
				COC	SPO	
Clostridium chauvoei	96.83	0.47	TRE 10	1	99	
Atopobium parvulum	2.29	0.19	MLZ 1	99	1	
Eubacterium saburreum	0.48	0.17		1	1	
47131120 Low discrimination						**47131120**
				COC	SPO	
Clostridium perfringens	86.68	0.90		1	84	
Atopobium parvulum	11.72	0.74		99	1	
Clostridium septicum	0.68	0.47		1	99	
47131121 Good identification						**47131121**
				COC	SPO	
Clostridium chauvoei	90.18	0.73	TRE 10	1	99	
Clostridium perfringens	8.35	0.56		1	84	
Atopobium parvulum	1.13	0.40		99	1	
47131122 Good identification						**47131122**
				COC	GRA	
Clostridium perfringens	92.71	0.84		1	99	
Atopobium parvulum	4.70	0.61		99	99	
Prevotella oralis	1.10	0.48		1	1	
47131123 Low discrimination						**47131123**
				SPO		
Clostridium chauvoei	78.20	0.60	TRE 10	99		
Clostridium perfringens	19.30	0.50		84		
Actinomyces naeslundii	1.05	0.51		1		
47131130 Low discrimination						**47131130**
				COC		
Clostridium perfringens	83.85	0.56	MLZ 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		11.33	0.40		99			
Eubacterium saburreum		2.39	0.38		1			
47131131	**Low discrimination**							**47131131**
					COC	SPO		
Clostridium chauvoei		89.49	0.38	TRE 10	1	99		
Clostridium perfringens		8.28	0.21	MLZ 1	1	84		
Atopobium parvulum		1.12	0.06		99	1		
47131132	**Low discrimination**							**47131132**
					COC	GRA	CAT	
Clostridium perfringens		77.69	0.50	MLZ 1	1	99	1	
Capnocytophaga ochracea		8.25	0.30		1	1	90	
Atopobium parvulum		3.94	0.27		99	99	1	
47131140	**Low discrimination**							**47131140**
					COC	GRA		
Atopobium parvulum		55.98	0.53	SOR 1	99	99		
Clostridium perfringens		42.63	0.52		1	99		
Fusobacterium mortiferum		0.68	0.32		1	1		
47131141	**Acceptable identification**							**47131141**
					COC	SPO		
Clostridium chauvoei		94.75	0.47	TRE 10	1	99		
Atopobium parvulum		2.24	0.19	SOR 1	99	1		
Clostridium perfringens		1.71	0.17		1	84		
47131142	**Low discrimination**							**47131142**
					COC	GRA	CAT	
Clostridium perfringens		64.46	0.46	NAG 97	1	99	1	
Atopobium parvulum		31.74	0.40		99	99	1	
Capnocytophaga ochracea		1.30	0.14		1	1	90	
47131160	**Very good identification**							**47131160**
					COC	SPO		
Clostridium perfringens		99.14	0.77		1	84		
Atopobium parvulum		0.71	0.40		99	1		
Clostridium septicum		0.04	0.13		1	99		
47131161	**Low discrimination**							**47131161**
					SPO			
Clostridium perfringens		62.35	0.43	URE 1	84			
Clostridium chauvoei		35.72	0.38		99			
Actinomyces naeslundii		1.48	0.38		1			
47131162	**Very good identification**							**47131162**
					COC	GRA	CAT	
Clostridium perfringens		99.02	0.72		1	99	1	
Capnocytophaga ochracea		0.56	0.30		1	1	90	
Atopobium parvulum		0.27	0.27		99	99	1	
47131163	**Low discrimination**							**47131163**
					SPO			
Clostridium perfringens		79.03	0.38	URE 1	84			
Clostridium chauvoei		16.98	0.25		99			
Actinomyces naeslundii		3.99	0.38		1			
47131170	**Good identification**							**47131170**
					COC	GRA	CAT	
Clostridium perfringens		98.97	0.43	MLZ 1	1	99	1	
Atopobium parvulum		0.71	0.06		99	99	1	
Capnocytophaga ochracea		0.32	-.03		1	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47131172	**Acceptable identification**						**47131172**
					GRA	CAT	
Clostridium perfringens		94.17	0.38	MLZ 1	99	1	
Capnocytophaga ochracea		5.83	0.14		1	90	
47131220	**Acceptable identification**						**47131220**
					CAT		
Clostridium perfringens		94.02	0.41	LAC 88	1		
Propionibacterium granulosum		2.77	0.22	RHA 1	90		
Lactobacillus jensenii		1.72	0.12		1		
47131222	**Low discrimination**						**47131222**
					GRA		
Clostridium perfringens		77.25	0.35	LAC 88	99		
Prevotella oralis		18.11	0.21	RHA 1	1		
Lactobacillus jensenii		2.01	0.09		99		
47131300	**Acceptable identification**						**47131300**
					COC	GRA	
Atopobium parvulum		94.32	0.53	RHA 1	99	99	
Clostridium perfringens		3.81	0.30		1	99	
Fusobacterium mortiferum		1.14	0.32		1	1	
47131301	**Good identification**						**47131301**
					COC	SPO	
Clostridium chauvoei		97.69	0.47	TRE 10	1	99	
Atopobium parvulum		2.31	0.19	RHA 1	99	1	
47131302	**Low discrimination**						**47131302**
					COC	GRA	
Atopobium parvulum		80.68	0.40	RHA 1	99	99	
Clostridium perfringens		8.69	0.24		1	99	
Prevotella oralis		7.41	0.20		1	1	
47131320	**Low discrimination**						**47131320**
					COC	GRA	
Clostridium perfringens		85.56	0.56	RHA 1	1	99	
Atopobium parvulum		11.56	0.40		99	99	
Prevotella oralis		1.06	0.20		1	1	
47131321	**Acceptable identification**						**47131321**
					COC	SPO	
Clostridium chauvoei		90.49	0.38	TRE 10	1	99	
Clostridium perfringens		8.38	0.21	RHA 1	1	84	
Atopobium parvulum		1.13	0.06		99	1	
47131322	**Low discrimination**						**47131322**
					GRA		
Clostridium perfringens		54.76	0.50	RHA 1	99		
Prevotella oralis		27.43	0.41		1		
Parabacteroides distasonis		13.62	0.49		1		
47131332	**Low discrimination**						**47131332**
					GRA		
Parabacteroides distasonis		85.02	0.44	RAF 99	1		
Clostridium perfringens		6.41	0.16	XYL 90	99		
Clostridium clostridioforme		5.35	0.18		50		
47131360	**Good identification**						**47131360**
					COC		
Clostridium perfringens		99.29	0.43	RHA 1	1		
Atopobium parvulum		0.71	0.06		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47131362	**Good identification**						**47131362**
					GRA		
Clostridium perfringens		96.18	0.38	RHA 1	99		
Prevotella oralis		2.56	0.07		1		
Parabacteroides distasonis		1.27	0.15		1		
47131420	**Acceptable identification**						**47131420**
Clostridium perfringens		94.34	0.41	LAC 88			
Bifidobacterium longum subsp. longum		2.08	0.18	ARA 1			
Lactobacillus jensenii		1.73	0.12				
47131422	**Acceptable identification**						**47131422**
Clostridium perfringens		92.13	0.35	LAC 88			
Bifidobacterium longum subsp. longum		4.33	0.18	ARA 1			
Lactobacillus jensenii		2.39	0.09				
47131500	**Low discrimination**						**47131500**
					COC		
Atopobium parvulum		86.01	0.53	ARA 1	99		
Bifidobacterium longum subsp. longum		8.37	0.41		1		
Clostridium perfringens		3.47	0.30		1		
47131501	**Good identification**						**47131501**
					COC	SPO	
Clostridium chauvoei		97.47	0.47	TRE 10	1	99	
Atopobium parvulum		2.30	0.19	ARA 1	99	1	
Bifidobacterium longum subsp. longum		0.22	0.07		1	1	
47131502	**Low discrimination**						**47131502**
					COC		
Atopobium parvulum		40.60	0.40	ARA 1	99		
Bifidobacterium adolescentis		28.01	0.45		1		
Bifidobacterium longum subsp. longum		22.40	0.41		1		
47131510	**Acceptable identification**						**47131510**
					COC		
Bifidobacterium longum subsp. longum		92.35	0.45	RAF 97	1		
Atopobium parvulum		5.63	0.19		99		
Eubacterium saburreum		1.19	0.17		1		
47131512	**Low discrimination**						**47131512**
Bifidobacterium longum subsp. longum		50.03	0.45	RAF 97			
Bifidobacterium adolescentis		44.91	0.46				
Clostridium clostridioforme		3.34	0.24				
47131520	**Low discrimination**						**47131520**
					COC		
Clostridium perfringens		68.37	0.56	ARA 1	1		
Bifidobacterium longum subsp. longum		20.39	0.52		1		
Atopobium parvulum		9.24	0.40		99		
47131521	**Low discrimination**						**47131521**
					SPO		
Clostridium chauvoei		88.29	0.38	TRE 10	99		
Clostridium perfringens		8.17	0.21	ARA 1	84		
Bifidobacterium longum subsp. longum		2.44	0.18		1		
47131522	**Low discrimination**						**47131522**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium perfringens		51.96	0.50	ARA 1	99	1		
Bifidobacterium longum subsp. longum		32.92	0.52		99	1		
Capnocytophaga ochracea		8.76	0.34		1	90		
47131530	**Good identification**							**47131530**
					COC			
Bifidobacterium longum subsp. longum		97.61	0.56	RAF 97	1			
Clostridium perfringens		1.94	0.21		1			
Atopobium parvulum		0.26	0.06		99			
47131532	**Acceptable identification**							**47131532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		93.37	0.56	RAF 97	99	1		
Parabacteroides distasonis		2.37	0.33		1	76		
Capnocytophaga ochracea		1.62	0.17		1	90		
47131560	**Good identification**							**47131560**
					COC			
Clostridium perfringens		97.26	0.43	ARA 1	1			
Bifidobacterium longum subsp. longum		1.54	0.18		1			
Atopobium parvulum		0.70	0.06		99			
47131562	**Low discrimination**							**47131562**
					GRA	CAT		
Clostridium perfringens		88.34	0.38	ARA 1	99	1		
Capnocytophaga ochracea		8.69	0.17		1	90		
Bifidobacterium longum subsp. longum		2.97	0.18		99	1		
47131712	**Good identification**							**47131712**
Clostridium clostridioforme		96.23	0.39	SAL 88				
Bifidobacterium longum subsp. longum		1.99	0.11	RAF 94				
Bifidobacterium adolescentis		1.78	0.12	XYL 91				
47131722	**Low discrimination**							**47131722**
					GRA			
Parabacteroides distasonis		72.99	0.45	RAF 99	1			
Clostridium clostridioforme		13.89	0.27	XYL 90	50			
Clostridium perfringens		4.84	0.16		99			
47131732	**Low discrimination**							**47131732**
					GRA			
Clostridium clostridioforme		58.27	0.39	SAL 88	50			
Parabacteroides distasonis		36.21	0.41	RAF 94	1			
Bifidobacterium longum subsp. longum		4.81	0.22	XYL 91	99			
47132000	**Good identification**							**47132000**
					GRA	SPO	CAT	
Propionibacterium granulosum		98.55	0.56	ESL 1	99	1	90	
Fusobacterium mortiferum		0.67	0.22		1	1	5	
Clostridium difficile		0.31	0.01		99	99	1	
47132002	**Good identification**							**47132002**
					CAT			
Propionibacterium granulosum		99.28	0.46	ESL 1	90			
Propionibacterium propionicum		0.72	0.14		1			
47132010	**Low discrimination**							**47132010**
					SPO	CAT		
Propionibacterium granulosum		89.80	0.41	ESL 1	1	90		
Clostridium difficile		10.20	0.13	MLZ 12	99	1		
47132020	**Good identification**							**47132020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Propionibacterium granulosum		98.61	0.46	ESL 1	90		
Clostridium perfringens		1.39	0.07		1		
47132022	**Good identification**						**47132022**
					CAT		
Propionibacterium granulosum		97.42	0.35	ESL 1	90		
Clostridium perfringens		2.58	0.01		1		
47132100	**Low discrimination**						**47132100**
					COC	GRA	
Propionibacterium propionicum		70.81	0.45	NIT 86	1	99	
Atopobium parvulum		11.73	0.19	ESL 1	99	99	
Fusobacterium mortiferum		9.37	0.28		1	1	
47132102	**Acceptable identification**						**47132102**
					COC		
Propionibacterium propionicum		94.06	0.48	NIT 86	1		
Actinomyces naeslundii		1.86	0.29	ESL 1	1		
Atopobium parvulum		1.83	0.06		99		
47132140	**Good identification**						**47132140**
					SPO		
Propionibacterium propionicum		98.27	0.45	NIT 86	1		
Clostridium botulinum biovar B		1.73	-.03	ESL 1	99		
47132142	**Good identification**						**47132142**
Propionibacterium propionicum		100.0	0.48	NIT 86			
47133000	**Acceptable identification**						**47133000**
					CAT		
Propionibacterium granulosum		90.69	0.56	ESL 1	90		
Actinomyces naeslundii		2.70	0.45		25		
Lactobacillus jensenii		2.32	0.22		1		
47133001	**Low discrimination**						**47133001**
					SPO	CAT	
Actinomyces naeslundii		70.65	0.48	ESL 12	1	25	
Propionibacterium granulosum		15.98	0.22	RAF 92	1	90	
Clostridium chauvoei		13.37	0.12		99	1	
47133002	**Low discrimination**						**47133002**
					CAT		
Propionibacterium granulosum		79.68	0.46	ESL 1	90		
Actinomyces naeslundii		9.49	0.45		25		
Lactobacillus jensenii		5.43	0.19		1		
47133003	**Acceptable identification**						**47133003**
					SPO	CAT	
Actinomyces naeslundii		91.75	0.48	ESL 12	1	25	
Propionibacterium granulosum		5.19	0.12	RAF 92	1	90	
Clostridium chauvoei		3.07	-.01		99	1	
47133010	**Acceptable identification**						**47133010**
					CAT		
Propionibacterium granulosum		91.99	0.41	ESL 1	90		
Lactobacillus jensenii		5.75	0.14	MLZ 12	1		
Eubacterium saburreum		1.21	0.11		1		
47133020	**Low discrimination**						**47133020**
					CAT		
Clostridium perfringens		54.62	0.41	MAN 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		39.17	0.46	LAC 88	90		
Actinomyces naeslundii		4.67	0.45		25		
47133021	**Low discrimination**						**47133021**
					SPO		
Actinomyces naeslundii		83.44	0.48	ESL 12	1		
Clostridium perfringens		6.58	0.07	RAF 92	84		
Clostridium chauvoei		5.26	0.04		99		
47133022	**Low discrimination**						**47133022**
					CAT		
Clostridium perfringens		62.95	0.35	MAN 1	1		
Propionibacterium granulosum		23.99	0.35	LAC 88	90		
Actinomyces naeslundii		11.43	0.45		25		
47133023	**Good identification**						**47133023**
Actinomyces naeslundii		96.42	0.48	ESL 12			
Clostridium perfringens		3.58	0.01	RAF 92			
47133100	**Low discrimination**						**47133100**
					COC		
Atopobium parvulum		84.70	0.53	MAN 1	99		
Propionibacterium propionicum		5.16	0.45		1		
Actinomyces naeslundii		3.67	0.53		1		
47133101	**Low discrimination**						**47133101**
					COC	SPO	
Clostridium chauvoei		83.94	0.47	TRE 10	1	99	
Actinomyces naeslundii		12.75	0.56	MAN 1	1	1	
Atopobium parvulum		1.98	0.19		99	1	
47133102	**Low discrimination**						**47133102**
					COC		
Atopobium parvulum		47.44	0.40	MAN 1	99		
Propionibacterium propionicum		24.58	0.48		1		
Actinomyces naeslundii		11.64	0.53		1		
47133120	**Low discrimination**						**47133120**
					COC		
Clostridium perfringens		83.60	0.56	MAN 1	1		
Atopobium parvulum		11.30	0.40		99		
Actinomyces naeslundii		2.77	0.53		1		
47133121	**Low discrimination**						**47133121**
					SPO		
Clostridium chauvoei		64.07	0.38	TRE 10	99		
Actinomyces naeslundii		29.20	0.56	MAN 1	1		
Clostridium perfringens		5.93	0.21		84		
47133122	**Low discrimination**						**47133122**
					COC		
Clostridium perfringens		81.65	0.50	MAN 1	1		
Actinomyces naeslundii		5.75	0.53		1		
Atopobium parvulum		4.14	0.27		99		
47133123	**Low discrimination**						**47133123**
					SPO		
Actinomyces naeslundii		67.44	0.56	ESL 12	1		
Clostridium chauvoei		26.12	0.25	RAF 92	99		
Clostridium perfringens		6.45	0.16		84		
47133140	**Low discrimination**						**47133140**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC			
Propionibacterium propionicum		67.83	0.45	NIT 86	1			
Atopobium parvulum		11.24	0.19	ESL 1	99			
Clostridium perfringens		8.56	0.17		1			
47133141	**Low discrimination**							**47133141**
					SPO			
Actinomyces naeslundii		50.82	0.43	ESL 12	1			
Propionibacterium propionicum		30.04	0.29	RAF 92	1			
Clostridium chauvoei		19.15	0.12		99			
47133142	**Low discrimination**							**47133142**
Propionibacterium propionicum		81.98	0.48	NIT 86				
Actinomyces naeslundii		6.85	0.40	ESL 1				
Actinomyces israelii		4.68	0.28					
47133143	**Low discrimination**							**47133143**
					SPO			
Actinomyces naeslundii		51.20	0.43	ESL 12	1			
Propionibacterium propionicum		45.40	0.32	RAF 92	1			
Clostridium chauvoei		3.41	-.01		99			
47133160	**Good identification**							**47133160**
					COC			
Clostridium perfringens		96.35	0.43	MAN 1	1			
Actinomyces naeslundii		2.96	0.40		1			
Atopobium parvulum		0.69	0.06		99			
47133161	**Low discrimination**							**47133161**
					SPO			
Actinomyces naeslundii		74.36	0.43	ESL 12	1			
Clostridium perfringens		16.30	0.09	RAF 92	84			
Clostridium chauvoei		9.34	0.04		99			
47133162	**Acceptable identification**							**47133162**
Clostridium perfringens		93.18	0.38	MAN 1				
Actinomyces naeslundii		6.08	0.40					
Propionibacterium propionicum		0.74	0.14					
47133163	**Acceptable identification**							**47133163**
Actinomyces naeslundii		90.65	0.43	ESL 12				
Clostridium perfringens		9.35	0.03	RAF 92				
47133502	**Low discrimination**							**47133502**
					COC			
Bifidobacterium adolescentis		87.15	0.43	RAF 92	1			
Actinomyces israelii		9.62	0.33	XYL 87	1			
Atopobium parvulum		1.56	0.06		99			
47133512	**Good identification**							**47133512**
Bifidobacterium adolescentis		97.35	0.45	RAF 92				
Bifidobacterium longum subsp. longum		1.34	0.11	XYL 87				
Actinomyces israelii		1.31	0.19					
47134000	**Low discrimination**							**47134000**
					COC	GRA	CAT	
Fusobacterium mortiferum		44.47	0.37	MLT 11	1	1	5	
Atopobium parvulum		32.37	0.23	XYL 7	99	99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Propionibacterium granulosum	11.80	0.24		1	99	90	
47134100	**Very good identification**							**47134100**
					COC	GRA		
	Atopobium parvulum	96.79	0.57	GAL 99	99	99		
	Fusobacterium mortiferum	3.13	0.43		1	1		
	Eubacterium saburreum	0.07	0.14		1	85		
47134102	**Good identification**							**47134102**
					COC	GRA		
	Atopobium parvulum	97.64	0.45	GAL 99	99	99		
	Fusobacterium mortiferum	1.99	0.27		1	1		
	Bifidobacterium adolescentis	0.37	0.11		1	99		
47134120	**Good identification**							**47134120**
					COC			
	Atopobium parvulum	96.13	0.45	GAL 99	99			
	Clostridium perfringens	3.87	0.21		1			
47134612	**Good identification**							**47134612**
	Clostridium clostridioforme	100.0	0.39	SAL 88				
47134632	**Good identification**							**47134632**
	Clostridium clostridioforme	100.0	0.39	SAL 88				
47134702	**Good identification**							**47134702**
	Clostridium clostridioforme	100.0	0.36	SAL 88				
47134712	**Good identification**							**47134712**
	Clostridium clostridioforme	100.0	0.48	SAL 88				
47134722	**Low discrimination**							**47134722**
	Clostridium clostridioforme	87.58	0.36	SAL 88				
	Parabacteroides distasonis	12.42	0.27	RAF 94				
47134732	**Good identification**							**47134732**
	Clostridium clostridioforme	98.35	0.48	SAL 88				
	Parabacteroides distasonis	1.65	0.23	RAF 94				
47135000	**Good identification**							**47135000**
					COC	GRA		
	Atopobium parvulum	93.37	0.57	LAC 99	99	99		
	Eubacterium saburreum	2.99	0.42		1	85		
	Fusobacterium mortiferum	1.94	0.40		1	1		
47135002	**Low discrimination**							**47135002**
					COC			
	Atopobium parvulum	89.96	0.45	LAC 99	99			
	Eubacterium saburreum	2.88	0.29		1			
	Lactobacillus jensenii	2.39	0.19		1			
47135010	**Low discrimination**							**47135010**
					COC			
	Eubacterium saburreum	66.94	0.40	IND 99	1			
	Atopobium parvulum	25.79	0.23		99			
	Lactobacillus jensenii	5.99	0.14		1			
47135020	**Low discrimination**							**47135020**
					COC			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		62.87	0.45	LAC 99	99		
Clostridium perfringens		34.16	0.41		1		
Eubacterium saburreum		2.01	0.29		1		
47135022	**Low discrimination**						**47135022**
					COC	GRA	
Clostridium perfringens		52.29	0.35	XYL 1	1	99	
Atopobium parvulum		36.09	0.32	LAC 88	99	99	
Prevotella oralis		3.54	0.12		1	1	
47135100	**Excellent identification**						**47135100**
					COC	GRA	
Atopobium parvulum		99.84	0.92		99	99	
Eubacterium saburreum		0.08	0.48		1	85	
Fusobacterium mortiferum		0.05	0.46		1	1	
47135101	**Low discrimination**						**47135101**
					COC	SPO	
Atopobium parvulum		81.21	0.57	URE 1	99	1	
Clostridium chauvoei		18.69	0.47		1	99	
Eubacterium saburreum		0.06	0.14		1	1	
47135102	**Very good identification**						**47135102**
					COC		
Atopobium parvulum		99.26	0.79		99		
Bifidobacterium adolescentis		0.37	0.45		1		
Actinomyces israelii		0.08	0.40		1		
47135103	**Low discrimination**						**47135103**
					COC	SPO	
Atopobium parvulum		81.04	0.45	URE 1	99	1	
Clostridium chauvoei		18.66	0.34		1	99	
Bifidobacterium adolescentis		0.30	0.11		1	1	
47135110	**Good identification**						**47135110**
					COC		
Atopobium parvulum		93.75	0.57	MLZ 1	99		
Eubacterium saburreum		5.74	0.46		1		
Bifidobacterium longum subsp. longum		0.25	0.19		1		
47135112	**Low discrimination**						**47135112**
					COC		
Atopobium parvulum		63.06	0.45	MLZ 1	99		
Bifidobacterium adolescentis		28.63	0.46		1		
Eubacterium saburreum		3.86	0.34		1		
47135120	**Very good identification**						**47135120**
					COC		
Atopobium parvulum		95.79	0.79		99		
Clostridium perfringens		3.86	0.56		1		
Eubacterium saburreum		0.07	0.35		1		
47135121	**Low discrimination**						**47135121**
					COC	SPO	
Atopobium parvulum		67.79	0.45	URE 1	99	1	
Clostridium chauvoei		29.48	0.38		1	99	
Clostridium perfringens		2.73	0.21		1	84	
47135122	**Low discrimination**						**47135122**
					COC	GRA	
Atopobium parvulum		78.01	0.66		99	99	
Clostridium perfringens		8.37	0.50		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		6.25	0.57		1	1		
47135130	**Low discrimination**							**47135130**
					COC			
Atopobium parvulum		85.05	0.45	MLZ 1	99			
Bifidobacterium longum subsp. longum		5.21	0.30		1			
Eubacterium saburreum		5.20	0.34		1			
47135132	**Low discrimination**							**47135132**
					COC	GRA		
Parabacteroides distasonis		67.97	0.53	RAF 99	1	1		
Atopobium parvulum		15.92	0.32		99	99		
Bifidobacterium longum subsp. longum		5.53	0.30		1	99		
47135140	**Very good identification**							**47135140**
					COC			
Atopobium parvulum		99.09	0.57	SOR 1	99			
Clostridium perfringens		0.41	0.17		1			
Actinomyces israelii		0.38	0.30		1			
47135142	**Acceptable identification**							**47135142**
					COC			
Atopobium parvulum		91.34	0.45	SOR 1	99			
Bifidobacterium adolescentis		4.19	0.29		1			
Actinomyces israelii		2.96	0.33		1			
47135160	**Low discrimination**							**47135160**
					COC	GRA	CAT	
Atopobium parvulum		56.68	0.45	SOR 1	99	99	1	
Clostridium perfringens		43.01	0.43		1	99	1	
Capnocytophaga ochracea		0.31	0.04		1	1	90	
47135162	**Low discrimination**							**47135162**
					COC	GRA	CAT	
Clostridium perfringens		58.66	0.38	XYL 1	1	99	1	
Atopobium parvulum		28.99	0.32		99	99	1	
Capnocytophaga ochracea		8.18	0.20		1	1	90	
47135300	**Very good identification**							**47135300**
					COC	GRA		
Atopobium parvulum		97.34	0.57	RHA 1	99	99		
Eubacterium contortum		2.38	0.29		1	99		
Bacteroides vulgatus		0.09	-.03		1	1		
47135302	**Low discrimination**							**47135302**
					COC	GRA		
Atopobium parvulum		89.44	0.45	RHA 1	99	99		
Parabacteroides distasonis		3.79	0.31		1	1		
Clostridium clostridioforme		3.14	0.24		1	50		
47135312	**Low discrimination**							**47135312**
					COC	GRA		
Clostridium clostridioforme		80.99	0.35	SAL 88	1	50		
Parabacteroides distasonis		11.57	0.27	RAF 94	1	1		
Atopobium parvulum		5.12	0.10	ARA 94	99	99		
47135320	**Low discrimination**							**47135320**
					COC	GRA		
Atopobium parvulum		61.34	0.45	RHA 1	99	99		
Bacteroides vulgatus		31.72	0.31		1	1		
Parabacteroides distasonis		2.60	0.31		1	1		
47135322	**Good identification**							**47135322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Parabacteroides distasonis		91.91	0.65	RAF 99	1	1	
Atopobium parvulum		3.87	0.32		99	99	
Prevotella oralis		2.54	0.26		1	1	
47135332	**Good identification**						**47135332**
Parabacteroides distasonis		93.39	0.61	RAF 99			
Clostridium clostridioforme		6.61	0.35				
47135500	**Low discrimination**						**47135500**
					COC		
Atopobium parvulum		84.94	0.57	ARA 1	99		
Bifidobacterium longum subsp. longum		13.36	0.50		1		
Actinomyces israelii		0.68	0.35		1		
47135502	**Low discrimination**						**47135502**
					COC		
Bifidobacterium adolescentis		54.63	0.59	RAF 92	1		
Atopobium parvulum		21.98	0.45		99		
Bifidobacterium longum subsp. longum		19.58	0.50		1		
47135510	**Acceptable identification**						**47135510**
					COC		
Bifidobacterium longum subsp. longum		93.72	0.54	RAF 97	1		
Atopobium parvulum		3.54	0.23		99		
Bifidobacterium adolescentis		1.90	0.26		1		
47135512	**Low discrimination**						**47135512**
Bifidobacterium adolescentis		61.73	0.60	RAF 92			
Bifidobacterium longum subsp. longum		30.83	0.54				
Clostridium clostridioforme		6.94	0.41				
47135520	**Low discrimination**						**47135520**
					COC	GRA	
Bifidobacterium longum subsp. longum		66.15	0.60	RAF 97	1	99	
Atopobium parvulum		18.56	0.45		99	99	
Bacteroides vulgatus		14.17	0.34		1	1	
47135522	**Low discrimination**						**47135522**
					COC	GRA	
Bifidobacterium longum subsp. longum		73.74	0.60	RAF 97	1	99	
Parabacteroides distasonis		17.74	0.53		1	1	
Atopobium parvulum		3.65	0.32		99	99	
47135530	**Very good identification**						**47135530**
					COC	GRA	
Bifidobacterium longum subsp. longum		99.58	0.64	RAF 97	1	99	
Atopobium parvulum		0.17	0.10		99	99	
Bacteroides vulgatus		0.13	0.00		1	1	
47135532	**Low discrimination**						**47135532**
					GRA		
Bifidobacterium longum subsp. longum		87.74	0.64	RAF 97	99		
Parabacteroides distasonis		6.68	0.49		1		
Clostridium clostridioforme		4.94	0.41		50		
47135542	**Low discrimination**						**47135542**
					COC		
Bifidobacterium adolescentis		87.08	0.43	RAF 92	1		
Actinomyces israelii		7.50	0.31	SOR 11	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		2.86	0.10		99		
47135552	**Acceptable identification**						**47135552**
Bifidobacterium adolescentis		94.29	0.45	RAF 92			
Bifidobacterium longum subsp. longum		3.85	0.19	SOR 11			
Actinomyces israelii		0.99	0.17				
47135602	**Good identification**						**47135602**
Clostridium clostridioforme		100.0	0.36	SAL 88			
47135612	**Good identification**						**47135612**
Clostridium clostridioforme		100.0	0.47	SAL 88			
47135622	**Low discrimination**						**47135622**
Clostridium clostridioforme		84.65	0.36	SAL 88			
Parabacteroides distasonis		13.39	0.27	RAF 94			
Bacteroides thetaiotaomicron		1.96	-.03				
47135632	**Good identification**						**47135632**
Clostridium clostridioforme		98.16	0.47	SAL 88			
Parabacteroides distasonis		1.84	0.23	RAF 94			
47135702	**Low discrimination**						**47135702**
					GRA		
Clostridium clostridioforme		85.23	0.45	SAL 88	50		
Parabacteroides distasonis		4.03	0.27	RAF 94	1		
Bifidobacterium adolescentis		3.89	0.25		99		
47135712	**Good identification**						**47135712**
Clostridium clostridioforme		97.66	0.56	SAL 88			
Bifidobacterium adolescentis		1.20	0.26	RAF 94			
Bifidobacterium longum subsp. longum		0.60	0.19				
47135720	**Good identification**						**47135720**
					GRA		
Bacteroides vulgatus		98.60	0.51	RAF 97	1		
Bifidobacterium longum subsp. longum		0.52	0.26	TRE 4	99		
Parabacteroides distasonis		0.37	0.27		1		
47135722	**Low discrimination**						**47135722**
Parabacteroides distasonis		75.27	0.62	RAF 99			
Clostridium clostridioforme		16.09	0.45				
Bacteroides thetaiotaomicron		5.46	0.26				
47135732	**Low discrimination**						**47135732**
Clostridium clostridioforme		62.25	0.56	SAL 88			
Parabacteroides distasonis		34.43	0.57	RAF 94			
Bacteroides thetaiotaomicron		1.80	0.19				
47137000	**Low discrimination**						**47137000**
					COC	CAT	
Actinomyces israelii		51.85	0.39	SAL 92	1	1	
Atopobium parvulum		37.77	0.23	CEL 93	99	1	
Propionibacterium granulosum		10.39	0.22		1	90	
47137002	**Low discrimination**						**47137002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	CAT	
Actinomyces israelii		87.33	0.43	SAL 92	1	1	
Atopobium parvulum		7.48	0.10	CEL 93	99	1	
Propionibacterium granulosum		2.92	0.12		1	90	
47137042	**Good identification**						**47137042**
Actinomyces israelii		100.0	0.35	SAL 92			
47137100	**Good identification**						**47137100**
					COC	SPO	
Atopobium parvulum		94.40	0.57	MAN 1	99	1	
Actinomyces israelii		5.24	0.50		1	1	
Clostridium tertium		0.20	0.11		1	99	
47137102	**Low discrimination**						**47137102**
					COC		
Atopobium parvulum		55.13	0.45	MAN 1	99		
Actinomyces israelii		25.99	0.53		1		
Bifidobacterium adolescentis		16.75	0.43		1		
47137112	**Low discrimination**						**47137112**
					SPO		
Bifidobacterium adolescentis		72.05	0.45	RAF 92	1		
Actinomyces israelii		13.66	0.39	ARA 87	1		
Clostridium tertium		12.33	0.24		99		
47137120	**Acceptable identification**						**47137120**
					COC	SPO	
Atopobium parvulum		91.24	0.45	MAN 1	99	1	
Clostridium perfringens		3.67	0.21		1	84	
Clostridium tertium		3.26	0.19		1	99	
47137140	**Low discrimination**						**47137140**
					COC		
Actinomyces israelii		68.11	0.43	SAL 92	1		
Atopobium parvulum		31.89	0.23	CEL 93	99		
47137142	**Low discrimination**						**47137142**
					COC		
Actinomyces israelii		78.41	0.46	SAL 92	1		
Bifidobacterium adolescentis		16.06	0.28	CEL 93	1		
Atopobium parvulum		4.32	0.10		99		
47137400	**Good identification**						**47137400**
Actinomyces israelii		100.0	0.38	SAL 92			
47137402	**Low discrimination**						**47137402**
Actinomyces israelii		82.44	0.41	SAL 92			
Bifidobacterium adolescentis		17.56	0.23	CEL 93			
47137500	**Low discrimination**						**47137500**
					COC		
Actinomyces israelii		74.18	0.48	SAL 92	1		
Atopobium parvulum		16.51	0.23	CEL 93	99		
Bifidobacterium adolescentis		5.93	0.23		1		
47137502	**Low discrimination**						**47137502**
					COC		
Bifidobacterium adolescentis		83.40	0.57	RAF 92	1		
Actinomyces israelii		15.82	0.51		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		0.41	0.10		99		
47137512	**Very good identification**						**47137512**
Bifidobacterium adolescentis		96.21	0.59	RAF 92			
Actinomyces israelii		2.23	0.37				
Bifidobacterium dentium		0.83	0.06				
47137540	**Good identification**						**47137540**
Actinomyces israelii		97.52	0.41	SAL 92			
Bifidobacterium adolescentis		2.48	0.08	CEL 93			
47137542	**Low discrimination**						**47137542**
Bifidobacterium adolescentis		62.63	0.42	RAF 92			
Actinomyces israelii		37.37	0.44	SOR 11			
47137552	**Acceptable identification**						**47137552**
Bifidobacterium adolescentis		93.20	0.43	RAF 92			
Actinomyces israelii		6.80	0.30	SOR 11			
47141000	**Low discrimination**						**47141000**
					COC		
Lactobacillus jensenii		55.51	0.36	SAL 95	1		
Lactobacillus catenaformis		21.22	0.29	FRU 99	1		
Gemella morbillorum		14.67	0.26		99		
47141101	**Very good identification**						**47141101**
					SPO		
Clostridium chauvoei		99.72	0.57	CEL 1	99		
Collinsella aerofaciens		0.21	-.03		1		
Actinomyces naeslundii		0.07	0.26		1		
47141103	**Good identification**						**47141103**
					SPO		
Clostridium chauvoei		99.59	0.44	CEL 1	99		
Actinomyces naeslundii		0.41	0.26		1		
47141120	**Low discrimination**						**47141120**
					GRA	SPO	
Prevotella melaninogenica		76.68	0.48	RAF 90	1	1	
Clostridium paraputrificum		12.40	0.32	FRU 90	99	99	
Collinsella aerofaciens		2.46	0.15	CEL 10	99	1	
47141121	**Good identification**						**47141121**
					GRA	SPO	
Clostridium chauvoei		97.31	0.48	CEL 1	99	99	
Prevotella melaninogenica		2.14	0.14		1	1	
Clostridium paraputrificum		0.35	-.03		99	99	
47141122	**Low discrimination**						**47141122**
					CAT		
Prevotella oralis		40.34	0.43	RAF 92	1		
Prevotella melaninogenica		22.79	0.36	FRU 99	1		
Capnocytophaga ochracea		14.70	0.36		90		
47141123	**Low discrimination**						**47141123**
					GRA	SPO	
Clostridium chauvoei		89.93	0.36	CEL 1	99	99	
Prevotella oralis		3.77	0.09		1	1	
Prevotella melaninogenica		2.13	0.02		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47141322	**Acceptable identification**							**47141322**
					GRA	SPO		
Prevotella oralis		93.03	0.37	RAF 92	1	1		
Clostridium barati		4.48	0.14	FRU 99	99	99		
Prevotella melaninogenica		1.24	0.02		1	1		
47150000	**Low discrimination**							**47150000**
					COC	GRA		
Fusobacterium mortiferum		66.10	0.57	MLT 11	1	1		
Lactobacillus jensenii		14.31	0.36		1	99		
Gemella morbillorum		8.58	0.32		99	99		
47150022	**Low discrimination**							**47150022**
					GRA	SPO		
Clostridium barati		76.04	0.36	SAL 99	99	99		
Prevotella oralis		11.76	0.23	GAL 95	1	1		
Lactobacillus jensenii		6.30	0.23		99	1		
47150100	**Low discrimination**							**47150100**
					GRA	SPO		
Fusobacterium mortiferum		65.04	0.63	MLT 11	1	1		
Collinsella aerofaciens		25.18	0.32		99	1		
Clostridium barati		4.04	0.28		99	99		
47150102	**Low discrimination**							**47150102**
					GRA	SPO		
Clostridium barati		41.18	0.36	SAL 99	99	99		
Fusobacterium mortiferum		24.54	0.47	GAL 95	1	1		
Lactobacillus catenaformis		11.75	0.32		99	1		
47150122	**Low discrimination**							**47150122**
					GRA	CAT	SPO	
Clostridium barati		41.77	0.44	SAL 99	99	1	99	
Prevotella oralis		33.74	0.43	GAL 95	1	1	1	
Capnocytophaga ochracea		12.30	0.36		1	90	1	
47150322	**Low discrimination**							**47150322**
					GRA	CAT	SPO	
Prevotella oralis		79.08	0.37	RAF 92	1	1	1	
Clostridium barati		19.86	0.26	GAL 99	99	1	99	
Capnocytophaga ochracea		0.68	0.02		1	90	1	
47151000	**Low discrimination**							**47151000**
					GRA			
Lactobacillus jensenii		64.58	0.70	SAL 95	99			
Lactobacillus catenaformis		24.69	0.63		99			
Fusobacterium mortiferum		4.52	0.60		1			
47151001	**Low discrimination**							**47151001**
					SPO			
Lactobacillus jensenii		47.52	0.36	SAL 95	1			
Clostridium chauvoei		22.33	0.29	URE 1	99			
Lactobacillus catenaformis		18.17	0.29		1			
47151002	**Low discrimination**							**47151002**
					SPO			
Lactobacillus jensenii		54.64	0.67	SAL 95	1			
Lactobacillus catenaformis		31.33	0.63		1			
Clostridium barati		10.54	0.50		99			
47151010	**Very good identification**							**47151010**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		98.35	0.62	SAL 95	99		
Lactobacillus catenaformis		1.14	0.29		99		
Fusobacterium mortiferum		0.21	0.26		1		
47151012	**Very good identification**						**47151012**
					SPO		
Lactobacillus jensenii		97.39	0.59	SAL 95	1		
Lactobacillus catenaformis		1.69	0.29		1		
Clostridium barati		0.57	0.15		99		
47151020	**Low discrimination**						**47151020**
					SPO		
Lactobacillus jensenii		42.66	0.60	SAL 95	1		
Clostridium barati		21.95	0.50		99		
Lactobacillus catenaformis		16.31	0.53		1		
47151022	**Low discrimination**						**47151022**
					GRA	SPO	
Clostridium barati		37.71	0.58	SAL 99	99	99	
Prevotella oralis		30.39	0.57		1	1	
Lactobacillus jensenii		16.29	0.57		99	1	
47151030	**Good identification**						**47151030**
					SPO		
Lactobacillus jensenii		95.96	0.52	SAL 95	1		
Clostridium barati		1.50	0.15		99		
Lactobacillus catenaformis		1.11	0.18		1		
47151032	**Low discrimination**						**47151032**
					GRA	SPO	
Lactobacillus jensenii		84.05	0.49	SAL 95	99	1	
Clostridium barati		5.90	0.24		99	99	
Prevotella oralis		4.75	0.23		1	1	
47151040	**Low discrimination**						**47151040**
					SPO		
Lactobacillus jensenii		52.55	0.36	SAL 95	1		
Lactobacillus catenaformis		20.09	0.29	SOR 1	1		
Clostridium barati		19.39	0.23		99		
47151062	**Low discrimination**						**47151062**
					GRA	SPO	
Clostridium barati		70.15	0.40	SAL 99	99	99	
Clostridium perfringens		9.05	0.29	SOR 8	99	84	
Prevotella oralis		6.57	0.23		1	1	
47151100	**Low discrimination**						**47151100**
					COC		
Collinsella aerofaciens		76.88	0.66	SAL 99	1		
Lactobacillus catenaformis		10.57	0.66		1		
Atopobium parvulum		5.25	0.59		99		
47151101	**Low discrimination**						**47151101**
					SPO		
Clostridium chauvoei		89.10	0.63	CEL 1	99		
Collinsella aerofaciens		7.99	0.32		1		
Lactobacillus catenaformis		1.10	0.32		1		
47151102	**Low discrimination**						**47151102**
					SPO		
Lactobacillus catenaformis		27.34	0.66	SAL 99	1		
Collinsella aerofaciens		22.11	0.50		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium barati		18.40	0.58		99			
47151103	**Low discrimination**							**47151103**
					SPO			
Clostridium chauvoei		77.82	0.50	CEL 1	99			
Lactobacillus catenaformis		5.44	0.32		1			
Collinsella aerofaciens		4.40	0.15		1			
47151120	**Low discrimination**							**47151120**
					GRA	SPO		
Clostridium paraputrificum		41.40	0.66	SAL 99	99	99		
Prevotella melaninogenica		23.28	0.64		1	1		
Collinsella aerofaciens		8.22	0.50		99	1		
47151121	**Low discrimination**							**47151121**
					GRA	SPO		
Clostridium chauvoei		72.41	0.55	CEL 1	99	99		
Clostridium paraputrificum		10.91	0.32		99	99		
Prevotella melaninogenica		6.13	0.30		1	1		
47151122	**Low discrimination**							**47151122**
					GRA	CAT	SPO	
Prevotella oralis		50.89	0.78	RAF 92	1	1	1	
Capnocytophaga ochracea		18.55	0.70		1	90	1	
Clostridium barati		12.09	0.66		99	1	99	
47151123	**Low discrimination**							**47151123**
					GRA	CAT	SPO	
Prevotella oralis		38.77	0.43	RAF 92	1	1	1	
Clostridium chauvoei		21.78	0.42	URE 1	99	1	99	
Capnocytophaga ochracea		14.13	0.36		1	90	1	
47151132	**Low discrimination**							**47151132**
					GRA	CAT		
Capnocytophaga ochracea		67.38	0.54	RAF 90	1	90		
Prevotella oralis		16.81	0.43	MLZ 10	1	1		
Bifidobacterium breve		4.04	0.44		99	1		
47151142	**Low discrimination**							**47151142**
					GRA	SPO	CAT	
Clostridium barati		36.20	0.40	SAL 99	99	99	1	
Capnocytophaga ochracea		23.65	0.38	SOR 8	1	1	90	
Lactobacillus acidophilus		14.52	0.35		99	1	1	
47151152	**Low discrimination**							**47151152**
					GRA	CAT		
Bifidobacterium breve		60.73	0.40	SAL 99	99	1		
Capnocytophaga ochracea		22.23	0.21	RAF 95	1	90		
Bifidobacterium adolescentis		9.76	0.25		99	1		
47151160	**Low discrimination**							**47151160**
Clostridium paraputrificum		54.23	0.47	SAL 99				
Clostridium perfringens		20.81	0.49	SOR 7				
Clostridium barati		10.35	0.40					
47151162	**Low discrimination**							**47151162**
					GRA	SPO		
Capnocytophaga ochracea		40.72	0.54	RAF 90	1	1		
Clostridium barati		20.77	0.48	SOR 10	99	99		
Clostridium paraputrificum		15.55	0.40		99	99		
47151172	**Low discrimination**							**47151172**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Capnocytophaga ochracea		60.08	0.38	RAF 90	1	1	90	
Bifidobacterium breve		32.43	0.44	MLZ 10	99	1	1	
Clostridium barati		2.79	0.14	SOR 10	99	99	1	
47151200	**Low discrimination**							**47151200**
					SPO			
Lactobacillus jensenii		52.63	0.36	SAL 95	1			
Lactobacillus catenaformis		20.12	0.29	RHA 1	1			
Clostridium barati		19.42	0.23		99			
47151222	**Low discrimination**							**47151222**
					GRA	SPO		
Prevotella oralis		78.08	0.51	RAF 92	1	1		
Clostridium barati		19.65	0.40	LAC 94	99	99		
Lactobacillus jensenii		0.99	0.23		99	1		
47151302	**Low discrimination**							**47151302**
					GRA	SPO		
Prevotella oralis		66.64	0.49	RAF 92	1	1		
Clostridium barati		20.34	0.40	NAG 95	99	99		
Lactobacillus acidophilus		4.97	0.31		99	1		
47151320	**Low discrimination**							**47151320**
					GRA	SPO		
Prevotella oralis		56.28	0.49	RAF 92	1	1		
Clostridium barati		17.18	0.40	BGL 95	99	99		
Clostridium paraputrificum		12.08	0.32		99	99		
47151322	**Good identification**							**47151322**
					GRA	CAT	SPO	
Prevotella oralis		93.34	0.71	RAF 92	1	1	1	
Clostridium barati		4.50	0.48		99	1	99	
Capnocytophaga ochracea		0.80	0.36		1	90	1	
47151323	**Acceptable identification**							**47151323**
					GRA	SPO		
Prevotella oralis		93.03	0.37	RAF 92	1	1		
Clostridium barati		4.48	0.14	URE 1	99	99		
Clostridium chauvoei		1.23	0.08		99	99		
47151332	**Low discrimination**							**47151332**
					CAT			
Prevotella oralis		81.75	0.37	RAF 92	1			
Capnocytophaga ochracea		7.72	0.20	MLZ 1	90			
Parabacteroides distasonis		4.61	0.28		76			
47151362	**Low discrimination**							**47151362**
					GRA	CAT	SPO	
Prevotella oralis		61.94	0.37	RAF 92	1	1	1	
Clostridium barati		25.70	0.30	SOR 1	99	1	99	
Capnocytophaga ochracea		5.85	0.20		1	90	1	
47151400	**Low discrimination**							**47151400**
					GRA			
Lactobacillus jensenii		62.41	0.36	SAL 95	99			
Lactobacillus catenaformis		23.86	0.29	ARA 1	99			
Fusobacterium mortiferum		4.37	0.26		1			
47151502	**Low discrimination**							**47151502**
					GRA	CAT		
Bifidobacterium adolescentis		46.12	0.53	RAF 92	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Capnocytophaga ochracea		33.87	0.41	XYL 87	1	90		
Lactobacillus catenaformis		5.64	0.32		99	1		
47151512	**Acceptable identification**							**47151512**
					GRA	CAT		
Bifidobacterium adolescentis		90.97	0.55	RAF 92	99	1		
Capnocytophaga ochracea		6.07	0.25	XYL 87	1	90		
Bifidobacterium longum subsp. longum		1.28	0.22		99	1		
47151520	**Low discrimination**							**47151520**
					GRA	CAT	SPO	
Capnocytophaga ochracea		37.67	0.41	RAF 90	1	90	1	
Clostridium paraputrificum		25.53	0.32	BGL 90	99	1	99	
Prevotella melaninogenica		14.36	0.30		1	1	1	
47151522	**Low discrimination**							**47151522**
					GRA	CAT	SPO	
Capnocytophaga ochracea		79.35	0.58	RAF 90	1	90	1	
Prevotella oralis		12.46	0.43		1	1	1	
Clostridium barati		2.96	0.32		99	1	99	
47151532	**Low discrimination**							**47151532**
					GRA	CAT		
Capnocytophaga ochracea		86.24	0.41	RAF 90	1	90		
Bifidobacterium longum subsp. longum		8.11	0.32	MLZ 10	99	1		
Clostridium clostridioforme		1.53	0.18		50	33		
47151542	**Low discrimination**							**47151542**
					GRA	SPO	CAT	
Bifidobacterium adolescentis		55.65	0.38	RAF 92	99	1	1	
Capnocytophaga ochracea		36.75	0.25	XYL 87	1	1	90	
Clostridium barati		3.22	0.05	SOR 11	99	99	1	
47151552	**Acceptable identification**							**47151552**
					GRA	CAT		
Bifidobacterium adolescentis		94.34	0.39	RAF 92	99	1		
Capnocytophaga ochracea		5.66	0.08	XYL 87	1	90		
47151562	**Acceptable identification**							**47151562**
					GRA	SPO		
Capnocytophaga ochracea		92.43	0.41	RAF 90	1	1		
Clostridium barati		2.70	0.14	SOR 10	99	99		
Clostridium paraputrificum		2.02	0.06		99	99		
47151722	**Low discrimination**							**47151722**
					CAT			
Prevotella oralis		75.74	0.37	RAF 92	1			
Capnocytophaga ochracea		11.37	0.23	ARA 1	90			
Parabacteroides distasonis		4.86	0.29		76			
47153000	**Low discrimination**							**47153000**
					SPO			
Lactobacillus jensenii		43.07	0.36	SAL 95	1			
Lactobacillus catenaformis		16.46	0.29	MAN 1	1			
Clostridium barati		15.90	0.23		99			
47153001	**Acceptable identification**							**47153001**
Actinomyces naeslundii		94.91	0.45	ESL 12				
Lactobacillus jensenii		5.09	0.02	RAF 92				
47153003	**Acceptable identification**							**47153003**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		94.40	0.45	ESL 12	1		
Clostridium barati		5.61	-.03	RAF 92	99		
47153021	**Acceptable identification**						**47153021**
					SPO		
Actinomyces naeslundii		94.40	0.45	ESL 12	1		
Clostridium barati		5.61	-.03	RAF 92	99		
47153022	**Low discrimination**						**47153022**
					GRA	SPO	
Clostridium barati		79.25	0.40	SAL 99	99	99	
Prevotella oralis		7.42	0.23	MAN 8	1	1	
Lactobacillus jensenii		3.98	0.23		99	1	
47153023	**Low discrimination**						**47153023**
					SPO		
Actinomyces naeslundii		84.88	0.45	ESL 12	1		
Clostridium barati		15.12	0.05	RAF 92	99		
47153101	**Low discrimination**						**47153101**
					SPO		
Actinomyces naeslundii		50.61	0.53	ESL 12	1		
Clostridium chauvoei		44.36	0.29	RAF 92	99		
Collinsella aerofaciens		3.98	-.03		1		
47153102	**Low discrimination**						**47153102**
					SPO		
Clostridium barati		37.83	0.40	SAL 99	99		
Clostridium tertium		12.86	0.35	MAN 8	99		
Bifidobacterium breve		7.50	0.40		1		
47153103	**Low discrimination**						**47153103**
					SPO		
Actinomyces naeslundii		80.74	0.53	ESL 12	1		
Clostridium chauvoei		12.49	0.16	RAF 92	99		
Clostridium barati		5.05	0.05		99		
47153112	**Low discrimination**						**47153112**
					SPO		
Clostridium tertium		56.39	0.39	SAL 99	99		
Bifidobacterium adolescentis		19.70	0.39	TRE 90	1		
Bifidobacterium breve		18.51	0.40		1		
47153120	**Low discrimination**						**47153120**
					GRA	SPO	
Clostridium barati		31.42	0.40	SAL 99	99	99	
Clostridium paraputrificum		22.10	0.32	MAN 8	99	99	
Prevotella melaninogenica		12.42	0.30		1	1	
47153121	**Low discrimination**						**47153121**
					SPO		
Actinomyces naeslundii		69.22	0.53	ESL 12	1		
Clostridium chauvoei		20.22	0.21	RAF 92	99		
Clostridium barati		4.33	0.05		99		
47153122	**Low discrimination**						**47153122**
					GRA	SPO	
Clostridium barati		39.43	0.48	SAL 99	99	99	
Prevotella oralis		19.28	0.43	MAN 8	1	1	
Clostridium tertium		13.40	0.43		99	99	
47153123	**Low discrimination**						**47153123**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces naeslundii		66.77	0.53	ESL 12	99	1		
Clostridium barati		12.53	0.14	RAF 92	99	99		
Prevotella oralis		6.13	0.09		1	1		
47153130	**Low discrimination**							**47153130**
					SPO			
Clostridium tertium		89.84	0.39	SAL 99	99			
Bifidobacterium breve		5.19	0.27	TRE 90	1			
Clostridium barati		1.50	0.05		99			
47153132	**Low discrimination**							**47153132**
					GRA	SPO	CAT	
Clostridium tertium		79.39	0.47	SAL 99	99	99	1	
Bifidobacterium breve		15.45	0.44	TRE 90	99	1	1	
Capnocytophaga ochracea		2.60	0.20		1	1	90	
47153141	**Good identification**							**47153141**
Actinomyces naeslundii		100.0	0.40	ESL 12				
47153142	**Low discrimination**							**47153142**
					SPO			
Bifidobacterium breve		46.34	0.40	SAL 99	1			
Clostridium barati		20.31	0.21	RAF 95	99			
Actinomyces israelii		15.69	0.37		1			
47153143	**Good identification**							**47153143**
Actinomyces naeslundii		100.0	0.40	ESL 12				
47153152	**Low discrimination**							**47153152**
Bifidobacterium breve		82.45	0.40	SAL 99				
Bifidobacterium adolescentis		10.84	0.24	RAF 95				
Actinomyces israelii		4.17	0.23					
47153161	**Good identification**							**47153161**
Actinomyces naeslundii		100.0	0.40	ESL 12				
47153162	**Low discrimination**							**47153162**
					GRA	SPO	CAT	
Bifidobacterium breve		45.25	0.44	SAL 99	99	1	1	
Clostridium barati		33.48	0.30	RAF 95	99	99	1	
Capnocytophaga ochracea		7.62	0.20		1	1	90	
47153163	**Low discrimination**							**47153163**
Actinomyces naeslundii		88.89	0.40	ESL 12				
Bifidobacterium breve		11.11	0.10	RAF 92				
47153172	**Acceptable identification**							**47153172**
					GRA	SPO	CAT	
Bifidobacterium breve		93.40	0.44	SAL 99	99	1	1	
Clostridium tertium		4.85	0.13	RAF 95	99	99	1	
Capnocytophaga ochracea		1.75	0.04		1	1	90	
47153322	**Low discrimination**							**47153322**
					SPO	GRA		
Prevotella oralis		41.12	0.37	RAF 92	1	1		
Clostridium ramosum		39.51	0.37	MAN 1	92	50		
Clostridium barati		17.06	0.30		99	99		
47153500	**Low discrimination**							**47153500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		77.26	0.40	SAL 92			
Collinsella aerofaciens		10.41	-.03	XYL 92			
Bifidobacterium adolescentis		9.57	0.18				
47153502	**Low discrimination**						**47153502**
					GRA	CAT	
Bifidobacterium adolescentis		87.48	0.52	RAF 92	99	1	
Actinomyces israelii		10.70	0.43	XYL 87	99	1	
Capnocytophaga ochracea		0.79	0.07		1	90	
47153512	**Good identification**						**47153512**
					SPO		
Bifidobacterium adolescentis		98.11	0.53	RAF 92	1		
Actinomyces israelii		1.47	0.28	XYL 87	1		
Clostridium tertium		0.42	0.05		99		
47153542	**Low discrimination**						**47153542**
Bifidobacterium adolescentis		72.22	0.36	RAF 92			
Actinomyces israelii		27.79	0.36	XYL 87			
47153552	**Good identification**						**47153552**
Bifidobacterium adolescentis		95.51	0.38	RAF 92			
Actinomyces israelii		4.49	0.21	XYL 87			
47154000	**Low discrimination**						**47154000**
					COC	GRA	
Fusobacterium mortiferum		81.92	0.38	MLT 11	1	1	
Gemella morbillorum		15.71	0.15	XYL 7	99	99	
Lactobacillus jensenii		2.38	0.02		1	99	
47154100	**Low discrimination**						**47154100**
					COC	GRA	
Fusobacterium mortiferum		58.68	0.44	MLT 11	1	1	
Atopobium parvulum		38.27	0.30	XYL 7	99	99	
Collinsella aerofaciens		3.05	-.03		1	99	
47154712	**Good identification**						**47154712**
Clostridium clostridioforme		100.0	0.42	SAL 88			
47154732	**Good identification**						**47154732**
Clostridium clostridioforme		100.0	0.42	SAL 88			
47155000	**Low discrimination**						**47155000**
					COC	GRA	
Lactobacillus jensenii		35.99	0.36	SAL 95	1	99	
Atopobium parvulum		19.05	0.30	XYL 1	99	99	
Fusobacterium mortiferum		18.77	0.41		1	1	
47155022	**Low discrimination**						**47155022**
					CAT		
Prevotella oralis		59.34	0.42	RAF 92	1		
Bacteroides fragilis		22.43	0.25	XYL 11	99		
Capnocytophaga ochracea		6.92	0.26	LAC 94	90		
47155100	**Low discrimination**						**47155100**
					COC	GRA	
Atopobium parvulum		86.95	0.64	TRE 99	99	99	
Collinsella aerofaciens		6.92	0.32		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		2.02	0.47		1	1		
47155102	**Low discrimination**							**47155102**
					COC	GRA	CAT	
Atopobium parvulum		35.11	0.51	TRE 99	99	99	1	
Capnocytophaga ochracea		18.54	0.44		1	1	90	
Bifidobacterium adolescentis		17.82	0.53		1	99	1	
47155112	**Low discrimination**							**47155112**
					GRA	CAT		
Bifidobacterium adolescentis		85.72	0.55	RAF 92	99	1		
Capnocytophaga ochracea		8.11	0.27	ARA 87	1	90		
Actinomyces israelii		2.60	0.35		99	1		
47155120	**Low discrimination**							**47155120**
					COC	GRA		
Atopobium parvulum		30.56	0.51	TRE 99	99	99		
Bacteroides fragilis		24.77	0.38		1	1		
Capnocytophaga ochracea		16.14	0.44		1	1		
47155122	**Low discrimination**							**47155122**
					CAT			
Bacteroides fragilis		56.63	0.60	RAF 99	99			
Prevotella oralis		23.71	0.62		1			
Capnocytophaga ochracea		17.47	0.60		90			
47155132	**Low discrimination**							**47155132**
					CAT			
Capnocytophaga ochracea		64.85	0.44	RAF 90	90			
Bacteroides fragilis		19.11	0.25	MLZ 10	99			
Prevotella oralis		8.00	0.28		1			
47155142	**Low discrimination**							**47155142**
					GRA	CAT		
Bifidobacterium adolescentis		32.92	0.38	RAF 92	99	1		
Capnocytophaga ochracea		30.79	0.27	ARA 87	1	90		
Actinomyces israelii		25.70	0.42	SOR 11	99	1		
47155152	**Low discrimination**							**47155152**
					GRA	CAT		
Bifidobacterium adolescentis		84.71	0.39	RAF 92	99	1		
Actinomyces israelii		8.09	0.28	ARA 87	99	1		
Capnocytophaga ochracea		7.20	0.11	SOR 11	1	90		
47155162	**Low discrimination**							**47155162**
					CAT			
Capnocytophaga ochracea		67.80	0.44	RAF 90	90			
Bacteroides fragilis		19.98	0.25	SOR 10	99			
Prevotella oralis		8.36	0.28		1			
47155222	**Acceptable identification**							**47155222**
					GRA	SPO		
Prevotella oralis		94.94	0.35	RAF 92	1	1		
Clostridium barati		1.95	0.05	XYL 11	99	99		
Parabacteroides distasonis		1.16	0.15	LAC 94	1	1		
47155322	**Low discrimination**							**47155322**
					CAT			
Prevotella oralis		85.92	0.56	RAF 92	1			
Parabacteroides distasonis		6.61	0.49	XYL 11	76			
Bacteroides fragilis		4.84	0.25		99			
47155332	**Low discrimination**							**47155332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		63.40	0.45	RAF 99			
Clostridium clostridioforme		17.32	0.30				
Prevotella oralis		15.46	0.22				
47155500	**Low discrimination**						**47155500**
					COC	SPO	
Actinomyces israelii		25.97	0.45	SAL 92	1	1	
Clostridium butyricum		25.26	0.24		1	99	
Atopobium parvulum		21.64	0.30		99	1	
47155502	**Good identification**						**47155502**
					GRA	CAT	
Bifidobacterium adolescentis		92.66	0.68	RAF 92	99	1	
Actinomyces israelii		2.81	0.48		99	1	
Capnocytophaga ochracea		2.54	0.31		1	90	
47155512	**Very good identification**						**47155512**
Bifidobacterium adolescentis		97.19	0.69	RAF 92			
Clostridium clostridioforme		1.57	0.35				
Bifidobacterium longum subsp. longum		0.62	0.30				
47155520	**Low discrimination**						**47155520**
					GRA	CAT	
Bacteroides vulgatus		68.42	0.36	RAF 97	1	3	
Capnocytophaga ochracea		15.42	0.31	CEL 5	1	90	
Bifidobacterium longum subsp. longum		10.22	0.36	RHA 90	99	1	
47155522	**Low discrimination**						**47155522**
					CAT		
Capnocytophaga ochracea		63.22	0.47	RAF 90	90		
Bacteroides fragilis		11.73	0.25		99		
Bacteroides uniformis		7.24	0.25		5		
47155530	**Low discrimination**						**47155530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.85	0.40	RAF 97	99	1	
Capnocytophaga ochracea		8.66	0.15	CEL 1	1	90	
Bacteroides vulgatus		3.49	0.02		1	3	
47155532	**Low discrimination**						**47155532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		31.56	0.40	RAF 97	99	1	
Capnocytophaga ochracea		27.99	0.31	CEL 1	1	90	
Clostridium clostridioforme		20.07	0.35		50	33	
47155540	**Low discrimination**						**47155540**
					SPO		
Actinomyces israelii		70.46	0.38	SAL 92	1		
Clostridium butyricum		15.33	0.05		99		
Bifidobacterium adolescentis		11.18	0.18		1		
47155542	**Low discrimination**						**47155542**
					GRA	CAT	
Bifidobacterium adolescentis		88.85	0.52	RAF 92	99	1	
Actinomyces israelii		8.48	0.41	SOR 11	99	1	
Capnocytophaga ochracea		2.19	0.15		1	90	
47155552	**Good identification**						**47155552**
					GRA	CAT	
Bifidobacterium adolescentis		98.63	0.53	RAF 92	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.15	0.27	SOR 11	99	1	
Capnocytophaga ochracea		0.22	-.02		1	90	
47155612	**Good identification**						**47155612**
Clostridium clostridioforme		100.0	0.41	SAL 88			
47155632	**Good identification**						**47155632**
Clostridium clostridioforme		100.0	0.41	SAL 88			
47155702	**Low discrimination**						**47155702**
Clostridium clostridioforme		66.27	0.39	SAL 88			
Bifidobacterium adolescentis		21.13	0.33	RAF 94			
Eubacterium contortum		5.28	0.14	TRE 88			
47155712	**Acceptable identification**						**47155712**
Clostridium clostridioforme		91.92	0.50	SAL 88			
Bifidobacterium adolescentis		7.87	0.35	RAF 94			
Actinomyces israelii		0.22	0.14	TRE 88			
47155720	**Good identification**						**47155720**
					GRA		
Bacteroides vulgatus		99.63	0.52	RAF 97	1		
Eubacterium contortum		0.23	0.14	CEL 5	99		
Prevotella oralis		0.04	0.00		1		
47155732	**Low discrimination**						**47155732**
Clostridium clostridioforme		85.64	0.50	SAL 88			
Parabacteroides distasonis		12.26	0.41	RAF 94			
Bacteroides thetaiotaomicron		0.87	0.05	TRE 88			
47157000	**Good identification**						**47157000**
					SPO		
Actinomyces israelii		98.08	0.49	SAL 92	1		
Clostridium innocuum		1.02	0.00		99		
Lactobacillus jensenii		0.90	0.02		1		
47157002	**Good identification**						**47157002**
					SPO		
Actinomyces israelii		95.54	0.52	SAL 92	1		
Bifidobacterium adolescentis		2.25	0.18		1		
Clostridium tertium		0.91	0.03		99		
47157012	**Low discrimination**						**47157012**
					SPO		
Actinomyces israelii		76.58	0.38	SAL 92	1		
Bifidobacterium adolescentis		14.72	0.19	MLZ 13	1		
Clostridium tertium		8.70	0.07		99		
47157040	**Good identification**						**47157040**
Actinomyces israelii		100.0	0.42	SAL 92			
47157042	**Good identification**						**47157042**
Actinomyces israelii		100.0	0.45	SAL 92			
47157100	**Low discrimination**						**47157100**
					COC	SPO	
Actinomyces israelii		79.23	0.59	SAL 92	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		9.53	0.30		99	1		
Clostridium tertium		9.36	0.29		1	99		
47157102	**Low discrimination**							**47157102**
					SPO			
Actinomyces israelii		54.00	0.62	SAL 92	1			
Bifidobacterium adolescentis		31.42	0.52		1			
Clostridium tertium		12.76	0.37		99			
47157112	**Low discrimination**							**47157112**
					SPO			
Bifidobacterium adolescentis		55.53	0.53	RAF 92	1			
Clostridium tertium		32.80	0.41	ARA 87	99			
Actinomyces israelii		11.67	0.48		1			
47157120	**Low discrimination**							**47157120**
					COC	SPO		
Clostridium tertium		72.16	0.37	SAL 99	1	99		
Actinomyces israelii		13.00	0.39	TRE 90	1	1		
Atopobium parvulum		4.32	0.17		99	1		
47157122	**Low discrimination**							**47157122**
					CAT	GRA	SPO	
Clostridium tertium		60.15	0.45	SAL 99	1	99	99	
Bacteroides fragilis		18.50	0.25	TRE 90	99	1	1	
Prevotella oralis		7.75	0.28		1	1	1	
47157130	**Good identification**							**47157130**
					SPO			
Clostridium tertium		98.51	0.41	SAL 99	99			
Actinomyces israelii		1.49	0.25	TRE 90	1			
47157132	**Good identification**							**47157132**
					GRA	SPO	CAT	
Clostridium tertium		97.81	0.50	SAL 99	99	99	1	
Actinomyces israelii		0.74	0.28	TRE 90	99	1	1	
Capnocytophaga ochracea		0.58	0.10		1	1	90	
47157140	**Good identification**							**47157140**
Actinomyces israelii		100.0	0.52	SAL 92				
47157142	**Low discrimination**							**47157142**
					SPO			
Actinomyces israelii		83.96	0.55	SAL 92	1			
Bifidobacterium adolescentis		15.53	0.36		1			
Clostridium tertium		0.52	0.03		99			
47157150	**Good identification**							**47157150**
					SPO			
Actinomyces israelii		96.48	0.38	SAL 92	1			
Clostridium tertium		3.52	-.01	MLZ 13	99			
47157152	**Low discrimination**							**47157152**
					SPO			
Bifidobacterium adolescentis		58.50	0.38	RAF 92	1			
Actinomyces israelii		38.68	0.41	ARA 87	1			
Clostridium tertium		2.82	0.07	SOR 11	99			
47157300	**Good identification**							**47157300**
Actinomyces israelii		100.0	0.40	SAL 92				
47157302	**Low discrimination**							**47157302**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		87.94	0.43	SAL 92	1		
Bifidobacterium adolescentis		6.87	0.18	RHA 7	1		
Clostridium tertium		2.79	0.03		99		
47157342	**Good identification**						**47157342**
Actinomyces israelii		100.0	0.36	SAL 92			
47157400	**Good identification**						**47157400**
Actinomyces israelii		100.0	0.47	SAL 92			
47157402	**Low discrimination**						**47157402**
Actinomyces israelii		83.87	0.50	SAL 92			
Bifidobacterium adolescentis		16.13	0.32				
47157440	**Good identification**						**47157440**
Actinomyces israelii		100.0	0.40	SAL 92			
47157442	**Acceptable identification**						**47157442**
Actinomyces israelii		94.24	0.43	SAL 92			
Bifidobacterium adolescentis		5.76	0.16				
47157500	**Good identification**						**47157500**
					SPO		
Actinomyces israelii		91.64	0.58	SAL 92	1		
Bifidobacterium adolescentis		6.61	0.32		1		
Clostridium butyricum		1.75	0.07		99		
47157502	**Low discrimination**						**47157502**
					SPO		
Bifidobacterium adolescentis		82.52	0.66	RAF 92	1		
Actinomyces israelii		17.34	0.61		1		
Clostridium butyricum		0.09	0.00		99		
47157510	**Low discrimination**						**47157510**
					SPO		
Actinomyces israelii		62.22	0.44	SAL 92	1		
Bifidobacterium adolescentis		36.70	0.33	MLZ 13	1		
Clostridium tertium		1.08	-.01		99		
47157512	**Very good identification**						**47157512**
Bifidobacterium adolescentis		97.15	0.68	RAF 92			
Actinomyces israelii		2.50	0.47				
Bifidobacterium dentium		0.26	0.06				
47157520	**Low discrimination**						**47157520**
					GRA	SPO	
Actinomyces israelii		80.87	0.37	SAL 92	99	1	
Bacteroides vulgatus		13.59	0.02	NAG 6	1	1	
Clostridium tertium		5.54	0.03		99	99	
47157522	**Low discrimination**						**47157522**
					GRA	CAT	
Actinomyces israelii		46.30	0.40	SAL 92	99	1	
Bifidobacterium adolescentis		34.87	0.32	NAG 6	99	1	
Capnocytophaga ochracea		10.53	0.13		1	90	
47157540	**Good identification**						**47157540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		97.76	0.51	SAL 92			
Bifidobacterium adolescentis		2.24	0.16				
47157542	**Low discrimination**						**47157542**
Bifidobacterium adolescentis		60.20	0.50	RAF 92			
Actinomyces israelii		39.80	0.54	SOR 11			
47157550	**Low discrimination**						**47157550**
					COC		
Actinomyces israelii		79.08	0.37	SAL 92	1		
Bifidobacterium adolescentis		14.82	0.18	MLZ 13	1		
Blautia producta		6.10	-.03		99		
47157552	**Acceptable identification**						**47157552**
Bifidobacterium adolescentis		92.52	0.52	RAF 92			
Actinomyces israelii		7.48	0.40	SOR 11			
47157700	**Good identification**						**47157700**
Actinomyces israelii		100.0	0.38	SAL 92			
47157702	**Low discrimination**						**47157702**
Actinomyces israelii		60.12	0.41	SAL 92			
Bifidobacterium adolescentis		38.40	0.32	RHA 7			
Clostridium clostridioforme		1.49	0.05				
47161000	**Acceptable identification**						**47161000**
					COC	GRA	
Lactobacillus jensenii		90.75	0.44	SAL 95	1	99	
Atopobium parvulum		8.62	0.25	FRU 99	99	99	
Fusobacterium mortiferum		0.63	0.17		1	1	
47161002	**Good identification**						**47161002**
					COC		
Lactobacillus jensenii		95.55	0.41	SAL 95	1		
Atopobium parvulum		2.40	0.12	FRU 99	99		
Lactobacillus acidophilus		1.52	0.07		1		
47161010	**Good identification**						**47161010**
Lactobacillus jensenii		100.0	0.36	SAL 95			
47161100	**Very good identification**						**47161100**
					COC	GRA	
Atopobium parvulum		99.28	0.59	FRU 99	99	99	
Fusobacterium mortiferum		0.17	0.23		1	1	
Collinsella aerofaciens		0.15	-.03		1	99	
47161101	**Low discrimination**						**47161101**
					COC	SPO	
Clostridium chauvoei		87.61	0.40	TRE 10	1	99	
Atopobium parvulum		11.27	0.25	CEL 1	99	1	
Actinomyces naeslundii		1.12	0.31		1	1	
47161102	**Acceptable identification**						**47161102**
					COC	SPO	
Atopobium parvulum		90.96	0.46	FRU 99	99	1	
Lactobacillus acidophilus		6.66	0.25		1	1	
Clostridium barati		0.61	0.05		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47161120	**Low discrimination**						**47161120**
					COC	SPO	
Atopobium parvulum		72.40	0.46	FRU 99	99	1	
Clostridium perfringens		9.57	0.32		1	84	
Clostridium paraputrificum		9.03	0.22		1	99	
47163101	**Low discrimination**						**47163101**
					SPO		
Actinomyces naeslundii		71.00	0.36	ESL 12	1		
Clostridium chauvoei		29.00	0.06	RAF 92	99		
47163103	**Good identification**						**47163103**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47163121	**Good identification**						**47163121**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47163123	**Good identification**						**47163123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47165100	**Very good identification**						**47165100**
					COC	SPO	
Atopobium parvulum		99.92	0.64	FRU 99	99	1	
Actinomyces israelii		0.06	0.22		1	1	
Clostridium butyricum		0.02	-.07		1	99	
47165102	**Good identification**						**47165102**
					COC		
Atopobium parvulum		99.19	0.51	FRU 99	99		
Actinomyces israelii		0.48	0.25		1		
Bifidobacterium adolescentis		0.34	0.16		1		
47165120	**Good identification**						**47165120**
Atopobium parvulum		100.0	0.51	FRU 99			
47165122	**Good identification**						**47165122**
					COC	GRA	
Atopobium parvulum		95.15	0.38	FRU 99	99	99	
Prevotella oralis		2.24	0.07		1	1	
Parabacteroides distasonis		1.76	0.18		1	1	
47167102	**Low discrimination**						**47167102**
					COC	SPO	
Actinomyces israelii		49.00	0.38	SAL 92	1	1	
Clostridium tertium		27.70	0.19	FRU 99	1	99	
Atopobium parvulum		18.25	0.17		99	1	
47167502	**Low discrimination**						**47167502**
Actinomyces israelii		54.25	0.36	SAL 92			
Bifidobacterium adolescentis		45.75	0.29	FRU 99			
47170000	**Low discrimination**						**47170000**
					COC	GRA	
Lactobacillus jensenii		62.76	0.44	SAL 95	1	99	
Fusobacterium mortiferum		28.87	0.48	GAL 99	1	1	
Atopobium parvulum		5.96	0.25		99	99	
47170002	**Low discrimination**						**47170002**
					GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus jensenii		86.42	0.41	SAL 95	99	1		
Fusobacterium mortiferum		6.63	0.32	GAL 99	1	1		
Clostridium barati		2.52	0.10		99	99		
47170010	**Good identification**							**47170010**
					GRA	CAT		
Lactobacillus jensenii		97.54	0.36	SAL 95	99	1		
Fusobacterium mortiferum		1.36	0.14	GAL 99	1	5		
Propionibacterium granulosum		1.10	0.09		99	90		
47170100	**Low discrimination**							**47170100**
					COC	GRA	SPO	
Atopobium parvulum		89.29	0.59	GAL 99	99	99	1	
Fusobacterium mortiferum		10.19	0.54		1	1	1	
Clostridium barati		0.18	0.10		1	99	99	
47170102	**Low discrimination**							**47170102**
					COC	GRA		
Atopobium parvulum		83.41	0.46	GAL 99	99	99		
Lactobacillus acidophilus		6.11	0.25		1	99		
Fusobacterium mortiferum		5.99	0.38		1	1		
47170120	**Low discrimination**							**47170120**
					COC	SPO		
Atopobium parvulum		72.01	0.46	GAL 99	99	1		
Clostridium perfringens		9.51	0.32		1	84		
Clostridium paraputrificum		8.98	0.22		1	99		
47170712	**Good identification**							**47170712**
Clostridium clostridioforme		100.0	0.39	SAL 88				
47170732	**Good identification**							**47170732**
Clostridium clostridioforme		100.0	0.39	SAL 88				
47171000	**Good identification**							**47171000**
					COC	GRA		
Lactobacillus jensenii		90.36	0.78	SAL 95	1	99		
Atopobium parvulum		8.58	0.59		99	99		
Fusobacterium mortiferum		0.63	0.51		1	1		
47171001	**Low discrimination**							**47171001**
					COC			
Lactobacillus jensenii		84.59	0.44	SAL 95	1			
Atopobium parvulum		8.04	0.25	URE 1	99			
Actinomyces naeslundii		5.31	0.45		1			
47171002	**Very good identification**							**47171002**
					COC			
Lactobacillus jensenii		95.04	0.75	SAL 95	1			
Atopobium parvulum		2.39	0.46		99			
Lactobacillus acidophilus		1.51	0.41		1			
47171003	**Low discrimination**							**47171003**
					COC			
Lactobacillus jensenii		87.31	0.41	SAL 95	1			
Actinomyces naeslundii		8.22	0.45	URE 1	1			
Atopobium parvulum		2.20	0.12		99			
47171010	**Very good identification**							**47171010**
					COC	GRA		
Lactobacillus jensenii		99.63	0.70	SAL 95	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		0.29	0.25		99	99	
Leptotrichia buccalis		0.03	-.03		1	1	
47171011	**Good identification**						**47171011**
Lactobacillus jensenii		99.02	0.36	SAL 95			
Actinomyces naeslundii		0.98	0.23	URE 1			
47171012	**Very good identification**						**47171012**
					COC	GRA	
Lactobacillus jensenii		95.28	0.67	SAL 95	1	99	
Leptotrichia buccalis		4.56	0.32		1	1	
Atopobium parvulum		0.07	0.12		99	99	
47171020	**Low discrimination**						**47171020**
					COC		
Lactobacillus jensenii		81.91	0.68	SAL 95	1		
Clostridium perfringens		9.80	0.51		1		
Atopobium parvulum		5.49	0.46		99		
47171022	**Low discrimination**						**47171022**
					SPO		
Lactobacillus jensenii		77.91	0.65	SAL 95	1		
Clostridium perfringens		6.58	0.46		84		
Clostridium barati		5.23	0.40		99		
47171030	**Very good identification**						**47171030**
					COC		
Lactobacillus jensenii		99.39	0.60	SAL 95	1		
Clostridium perfringens		0.36	0.17		1		
Atopobium parvulum		0.20	0.12		99		
47171032	**Very good identification**						**47171032**
					GRA		
Lactobacillus jensenii		97.18	0.57	SAL 95	99		
Leptotrichia buccalis		2.07	0.15		1		
Clostridium perfringens		0.25	0.12		99		
47171040	**Low discrimination**						**47171040**
					COC		
Lactobacillus jensenii		86.95	0.44	SAL 95	1		
Atopobium parvulum		8.26	0.25	SOR 1	99		
Clostridium perfringens		1.52	0.13		1		
47171042	**Low discrimination**						**47171042**
					SPO		
Lactobacillus jensenii		76.38	0.41	SAL 95	1		
Lactobacillus acidophilus		14.82	0.26	SOR 1	1		
Clostridium barati		3.68	0.13		99		
47171050	**Good identification**						**47171050**
Lactobacillus jensenii		100.0	0.36	SAL 95			
47171060	**Low discrimination**						**47171060**
					SPO		
Clostridium perfringens		63.66	0.39	CEL 4	84		
Lactobacillus jensenii		28.23	0.34	LAC 88	1		
Clostridium barati		3.62	0.13		99		
47171100	**Excellent identification**						**47171100**
					COC	GRA	
Atopobium parvulum		99.36	0.93		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.17	0.57		1	1	
Collinsella aerofaciens		0.15	0.32		1	99	
47171101	**Low discrimination**						**47171101**
					COC	SPO	
Atopobium parvulum		82.81	0.59	URE 1	99	1	
Clostridium chauvoei		15.17	0.47		1	99	
Actinomyces naeslundii		1.57	0.53		1	1	
47171102	**Good identification**						**47171102**
					COC	SPO	
Atopobium parvulum		90.62	0.80		99	1	
Lactobacillus acidophilus		6.64	0.60		1	1	
Clostridium barati		0.61	0.40		1	99	
47171103	**Low discrimination**						**47171103**
					COC	SPO	
Atopobium parvulum		72.12	0.46	URE 1	99	1	
Clostridium chauvoei		13.21	0.34		1	99	
Actinomyces naeslundii		7.77	0.53		1	1	
47171110	**Low discrimination**						**47171110**
					COC	GRA	
Atopobium parvulum		86.32	0.59	MLZ 1	99	99	
Leptotrichia buccalis		9.67	0.32		1	1	
Lactobacillus jensenii		3.06	0.36		1	99	
47171112	**Very good identification**						**47171112**
					COC	GRA	
Leptotrichia buccalis		97.69	0.66	SAL 99	1	1	
Atopobium parvulum		1.55	0.46		99	99	
Lactobacillus jensenii		0.21	0.33		1	99	
47171120	**Low discrimination**						**47171120**
					COC	SPO	
Atopobium parvulum		73.75	0.80		99	1	
Clostridium perfringens		9.74	0.66		1	84	
Clostridium paraputrificum		9.20	0.56		1	99	
47171121	**Low discrimination**						**47171121**
					COC	SPO	
Atopobium parvulum		55.41	0.46	URE 1	99	1	
Clostridium chauvoei		19.18	0.38		1	99	
Clostridium perfringens		7.32	0.32		1	84	
47171122	**Low discrimination**						**47171122**
					COC	GRA	
Atopobium parvulum		37.71	0.68		99	99	
Lactobacillus acidophilus		15.65	0.60		1	99	
Prevotella oralis		13.35	0.57		1	1	
47171130	**Low discrimination**						**47171130**
					COC	SPO	
Atopobium parvulum		65.28	0.46	MLZ 1	99	1	
Clostridium perfringens		8.63	0.32		1	84	
Clostridium paraputrificum		8.14	0.22		1	99	
47171132	**Low discrimination**						**47171132**
					COC	GRA	
Leptotrichia buccalis		89.24	0.50	SAL 99	1	1	
Atopobium parvulum		2.26	0.33	NAG 10	99	99	
Bifidobacterium breve		2.00	0.41		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47171140	**Very good identification**							**47171140**
					COC			
Atopobium parvulum		96.92	0.59	SOR 1	99			
Clostridium perfringens		1.32	0.28		1			
Actinomyces israelii		0.34	0.31		1			
47171142	**Low discrimination**							**47171142**
					COC			
Atopobium parvulum		46.41	0.46	SOR 1	99			
Lactobacillus acidophilus		41.59	0.44		1			
Bifidobacterium breve		4.09	0.37		1			
47171143	**Low discrimination**							**47171143**
					COC			
Actinomyces naeslundii		47.50	0.40	ESL 12	1			
Atopobium parvulum		25.25	0.12	RAF 92	99			
Lactobacillus acidophilus		22.63	0.10		1			
47171160	**Low discrimination**							**47171160**
					COC	SPO		
Clostridium perfringens		54.07	0.54	CEL 4	1	84		
Atopobium parvulum		21.70	0.46		99	1		
Clostridium paraputrificum		20.17	0.37		1	99		
47171162	**Low discrimination**							**47171162**
					SPO			
Clostridium perfringens		37.83	0.48	CEL 4	84			
Lactobacillus acidophilus		28.91	0.44		1			
Clostridium paraputrificum		12.85	0.31		99			
47171163	**Low discrimination**							**47171163**
Actinomyces naeslundii		43.27	0.40	ESL 12				
Clostridium perfringens		26.96	0.14	RAF 92				
Lactobacillus acidophilus		20.61	0.10					
47171172	**Low discrimination**							**47171172**
					GRA			
Bifidobacterium breve		61.10	0.41	SAL 99	99			
Leptotrichia buccalis		27.48	0.15	RAF 95	1			
Clostridium perfringens		4.61	0.14		99			
47171200	**Low discrimination**							**47171200**
					COC	SPO		
Lactobacillus jensenii		89.87	0.44	SAL 95	1	1		
Atopobium parvulum		8.54	0.25	RHA 1	99	1		
Clostridium barati		0.96	0.05		1	99		
47171202	**Low discrimination**							**47171202**
					SPO			
Lactobacillus jensenii		81.94	0.41	SAL 95	1			
Lactobacillus acidophilus		9.68	0.22	RHA 1	1			
Clostridium barati		3.94	0.13		99			
47171210	**Good identification**							**47171210**
Lactobacillus jensenii		100.0	0.36	SAL 95				
47171300	**Very good identification**							**47171300**
					COC	GRA	SPO	
Atopobium parvulum		98.83	0.59	RHA 1	99	99	1	
Clostridium barati		0.34	0.13		1	99	99	

Code Taxon	ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum	0.17	0.23		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47171302	**Low discrimination**							**47171302**
					COC	GRA		
Atopobium parvulum		56.06	0.46	RHA 1	99	99		
Lactobacillus acidophilus		30.59	0.40		1	99		
Prevotella oralis		7.82	0.29		1	1		
47171320	**Low discrimination**							**47171320**
					COC	SPO	GRA	
Atopobium parvulum		54.88	0.46	RHA 1	99	1	99	
Clostridium ramosum		15.29	0.35		1	92	50	
Prevotella oralis		7.66	0.29		1	1	1	
47171322	**Low discrimination**							**47171322**
					SPO	GRA		
Prevotella oralis		53.12	0.51	RAF 92	1	1		
Clostridium ramosum		22.33	0.45	TRE 6	92	50		
Lactobacillus acidophilus		10.94	0.40		1	99		
47171332	**Low discrimination**							**47171332**
Parabacteroides distasonis		40.79	0.40	RAF 99				
Leptotrichia buccalis		26.74	0.15	XYL 90				
Clostridium clostridioforme		15.97	0.27					
47171400	**Low discrimination**							**47171400**
					COC			
Lactobacillus jensenii		89.01	0.44	SAL 95	1			
Atopobium parvulum		8.46	0.25	ARA 1	99			
Actinomyces israelii		1.54	0.26		1			
47171402	**Low discrimination**							**47171402**
					COC			
Lactobacillus jensenii		89.60	0.41	SAL 95	1			
Actinomyces israelii		3.49	0.29	ARA 1	1			
Atopobium parvulum		2.25	0.12		99			
47171410	**Good identification**							**47171410**
Lactobacillus jensenii		100.0	0.36	SAL 95				
47171500	**Very good identification**							**47171500**
					COC	SPO		
Atopobium parvulum		97.57	0.59	ARA 1	99	1		
Clostridium butyricum		1.04	0.18		1	99		
Actinomyces israelii		0.72	0.36		1	1		
47171502	**Low discrimination**							**47171502**
					COC			
Atopobium parvulum		54.38	0.46	ARA 1	99			
Bifidobacterium adolescentis		34.16	0.50		1			
Lactobacillus acidophilus		3.98	0.25		1			
47171512	**Low discrimination**							**47171512**
					GRA			
Bifidobacterium adolescentis		50.64	0.52	RAF 92	99			
Leptotrichia buccalis		41.87	0.32	XYL 87	1			
Clostridium clostridioforme		5.94	0.33		50			
47171520	**Low discrimination**							**47171520**
					COC	SPO		
Atopobium parvulum		72.71	0.46	ARA 1	99	1		
Clostridium perfringens		9.61	0.32		1	84		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		9.07	0.22		1	99	
47171552	**Acceptable identification**						**47171552**
					GRA		
Bifidobacterium adolescentis		90.42	0.36	RAF 92	99		
Leptotrichia buccalis		6.11	-.03	XYL 87	1		
Actinomyces israelii		3.47	0.18	SOR 11	99		
47171612	**Good identification**						**47171612**
Clostridium clostridioforme		100.0	0.39	SAL 88			
47171632	**Good identification**						**47171632**
Clostridium clostridioforme		100.0	0.39	SAL 88			
47171702	**Low discrimination**						**47171702**
					COC		
Clostridium clostridioforme		84.45	0.36	SAL 88	1		
Atopobium parvulum		5.89	0.12	RAF 94	99		
Bifidobacterium adolescentis		3.70	0.16	XYL 91	1		
47171712	**Good identification**						**47171712**
					GRA		
Clostridium clostridioforme		97.90	0.48	SAL 88	50		
Bifidobacterium adolescentis		1.15	0.18	RAF 94	99		
Leptotrichia buccalis		0.95	-.03	XYL 91	1		
47171722	**Low discrimination**						**47171722**
Clostridium clostridioforme		44.71	0.36	SAL 88			
Parabacteroides distasonis		37.80	0.41	RAF 94			
Prevotella oralis		8.26	0.17	XYL 91			
47171732	**Acceptable identification**						**47171732**
Clostridium clostridioforme		90.28	0.48	SAL 88			
Parabacteroides distasonis		9.02	0.36	RAF 94			
Bacteroides thetaiotaomicron		0.70	0.01	XYL 91			
47173000	**Low discrimination**						**47173000**
					COC		
Lactobacillus jensenii		72.92	0.44	SAL 95	1		
Actinomyces israelii		8.74	0.41	MAN 1	1		
Atopobium parvulum		6.93	0.25		99		
47173001	**Acceptable identification**						**47173001**
Actinomyces naeslundii		92.35	0.50	ESL 12			
Lactobacillus jensenii		7.65	0.10	RAF 92			
47173002	**Low discrimination**						**47173002**
Lactobacillus jensenii		60.80	0.41	SAL 95			
Actinomyces israelii		16.41	0.44	MAN 1			
Actinomyces naeslundii		7.41	0.47				
47173003	**Acceptable identification**						**47173003**
Actinomyces naeslundii		94.77	0.50	ESL 12			
Lactobacillus jensenii		5.24	0.07	RAF 92			
47173010	**Low discrimination**						**47173010**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		88.70	0.36	SAL 95	1		
Clostridium tertium		5.39	0.13	MAN 1	99		
Actinomyces israelii		4.77	0.26		1		
47173021	**Good identification**						**47173021**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47173023	**Good identification**						**47173023**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47173041	**Good identification**						**47173041**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47173042	**Low discrimination**						**47173042**
Actinomyces israelii		71.50	0.37	SAL 92			
Actinomyces naeslundii		14.66	0.34	XYL 92			
Lactobacillus acidophilus		6.96	0.04				
47173043	**Good identification**						**47173043**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47173061	**Good identification**						**47173061**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47173063	**Good identification**						**47173063**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47173100	**Low discrimination**						**47173100**
					COC	SPO	
Atopobium parvulum		82.75	0.59	MAN 1	99	1	
Clostridium tertium		9.93	0.43		1	99	
Actinomyces israelii		4.22	0.51		1	1	
47173101	**Low discrimination**						**47173101**
					COC	SPO	
Actinomyces naeslundii		72.95	0.58	ESL 12	1	1	
Atopobium parvulum		19.98	0.25	RAF 92	99	1	
Clostridium chauvoei		3.66	0.12		1	99	
47173102	**Low discrimination**						**47173102**
					COC	SPO	
Clostridium tertium		47.63	0.51	SAL 99	1	99	
Atopobium parvulum		23.34	0.46		99	1	
Actinomyces israelii		10.12	0.54		1	1	
47173103	**Low discrimination**						**47173103**
					COC	SPO	
Actinomyces naeslundii		83.67	0.58	ESL 12	1	1	
Clostridium tertium		8.25	0.17	RAF 92	1	99	
Atopobium parvulum		4.04	0.12		99	1	
47173110	**Acceptable identification**						**47173110**
					COC	SPO	
Clostridium tertium		90.64	0.47	SAL 99	1	99	
Atopobium parvulum		4.29	0.25		99	1	
Actinomyces israelii		3.24	0.37		1	1	
47173112	**Low discrimination**						**47173112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium tertium		80.07	0.55	SAL 99	99	99	
Leptotrichia buccalis		14.01	0.32		1	1	
Bifidobacterium adolescentis		2.07	0.36		99	1	
47173120	**Low discrimination**						**47173120**
					COC	SPO	
Clostridium tertium		42.44	0.51	SAL 99	1	99	
Clostridium ramosum		24.76	0.45		1	92	
Atopobium parvulum		20.80	0.46		99	1	
47173121	**Low discrimination**						**47173121**
					SPO		
Actinomyces naeslundii		81.89	0.58	ESL 12	1		
Clostridium tertium		8.07	0.17	RAF 92	99		
Clostridium ramosum		4.71	0.11		92		
47173122	**Low discrimination**						**47173122**
					SPO		
Clostridium tertium		50.50	0.59	SAL 99	99		
Clostridium ramosum		39.29	0.56		92		
Lactobacillus acidophilus		3.15	0.38		1		
47173123	**Low discrimination**						**47173123**
					SPO		
Actinomyces naeslundii		64.75	0.58	ESL 12	1		
Clostridium tertium		19.15	0.25	RAF 92	99		
Clostridium ramosum		14.90	0.21		92		
47173130	**Good identification**						**47173130**
					SPO		
Clostridium tertium		98.70	0.55	SAL 99	99		
Bifidobacterium breve		0.42	0.24		1		
Clostridium ramosum		0.33	0.11		92		
47173132	**Very good identification**						**47173132**
					GRA	SPO	
Clostridium tertium		97.40	0.63	SAL 99	99	99	
Bifidobacterium breve		1.40	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
47173140	**Low discrimination**						**47173140**
					COC		
Actinomyces israelii		53.57	0.44	SAL 92	1		
Atopobium parvulum		27.28	0.25	XYL 92	99		
Actinomyces naeslundii		11.72	0.42		1		
47173141	**Good identification**						**47173141**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47173142	**Low discrimination**						**47173142**
Actinomyces israelii		44.83	0.47	SAL 92			
Bifidobacterium breve		23.46	0.37	XYL 92			
Lactobacillus acidophilus		12.54	0.22				
47173143	**Good identification**						**47173143**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47173152	**Low discrimination**						**47173152**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		52.49	0.37	SAL 99	1		
Clostridium tertium		21.80	0.21	RAF 95	99		
Actinomyces israelii		14.99	0.33		1		
47173161	**Good identification**						**47173161**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47173162	**Low discrimination**						**47173162**
					SPO		
Bifidobacterium breve		41.24	0.41	SAL 99	1		
Clostridium tertium		16.26	0.25	RAF 95	99		
Clostridium ramosum		12.65	0.21		92		
47173163	**Good identification**						**47173163**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47173172	**Low discrimination**						**47173172**
					SPO		
Bifidobacterium breve		58.79	0.41	SAL 99	1		
Clostridium tertium		41.21	0.29	RAF 95	99		
47173320	**Good identification**						**47173320**
					COC	SPO	
Clostridium ramosum		97.79	0.47	SAL 99	1	92	
Clostridium tertium		1.28	0.17		1	99	
Atopobium parvulum		0.63	0.12		99	1	
47173322	**Very good identification**						**47173322**
					SPO		
Clostridium ramosum		97.88	0.58	SAL 99	92		
Clostridium tertium		0.96	0.25		99		
Lactobacillus acidophilus		0.45	0.18		1		
47173400	**Low discrimination**						**47173400**
					SPO		
Actinomyces israelii		87.07	0.39	SAL 92	1		
Lactobacillus jensenii		8.96	0.10	XYL 92	1		
Clostridium innocuum		3.97	0.01		99		
47173402	**Low discrimination**						**47173402**
Actinomyces israelii		88.13	0.42	SAL 92			
Bifidobacterium adolescentis		5.16	0.15	XYL 92			
Lactobacillus jensenii		4.03	0.07				
47173442	**Good identification**						**47173442**
Actinomyces israelii		100.0	0.35	SAL 92			
47173500	**Low discrimination**						**47173500**
					COC	SPO	
Actinomyces israelii		75.68	0.49	SAL 92	1	1	
Atopobium parvulum		18.32	0.25	XYL 92	99	1	
Clostridium tertium		2.20	0.09		1	99	
47173502	**Low discrimination**						**47173502**
					SPO		
Bifidobacterium adolescentis		56.74	0.49	RAF 92	1		
Actinomyces israelii		39.15	0.52	XYL 87	1		
Clostridium tertium		2.28	0.17		99		
47173510	**Low discrimination**						**47173510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		65.56	0.35	SAL 92	1		
Clostridium tertium		22.66	0.13	XYL 92	99		
Bifidobacterium adolescentis		11.77	0.16	MLZ 13	1		
47173512	**Low discrimination**						**47173512**
					SPO		
Bifidobacterium adolescentis		84.74	0.50	RAF 92	1		
Actinomyces israelii		7.15	0.38	XYL 87	1		
Clostridium tertium		4.94	0.21		99		
47173540	**Good identification**						**47173540**
Actinomyces israelii		100.0	0.42	SAL 92			
47173542	**Low discrimination**						**47173542**
Actinomyces israelii		68.47	0.45	SAL 92			
Bifidobacterium adolescentis		31.53	0.33	XYL 92			
47174100	**Very good identification**						**47174100**
					COC	GRA	
Atopobium parvulum		99.46	0.64	GAL 99	99	99	
Fusobacterium mortiferum		0.46	0.35		1	1	
Actinomyces israelii		0.06	0.22		1	99	
47174102	**Good identification**						**47174102**
					COC		
Atopobium parvulum		98.68	0.51	GAL 99	99		
Actinomyces israelii		0.47	0.25		1		
Bifidobacterium adolescentis		0.34	0.16		1		
47174120	**Good identification**						**47174120**
Atopobium parvulum		100.0	0.51	GAL 99			
47174122	**Acceptable identification**						**47174122**
					COC	GRA	
Atopobium parvulum		93.99	0.38	GAL 99	99	99	
Prevotella oralis		2.21	0.07		1	1	
Parabacteroides distasonis		1.74	0.18		1	1	
47174312	**Good identification**						**47174312**
Clostridium clostridioforme		100.0	0.36	SAL 88			
47174332	**Acceptable identification**						**47174332**
Clostridium clostridioforme		93.55	0.36	SAL 88			
Parabacteroides distasonis		6.45	0.22	RAF 94			
47174512	**Low discrimination**						**47174512**
Clostridium clostridioforme		85.39	0.42	SAL 88			
Bifidobacterium adolescentis		14.61	0.32	RAF 94			
47174532	**Good identification**						**47174532**
Clostridium clostridioforme		99.35	0.42	SAL 88			
Parabacteroides distasonis		0.66	0.10	RAF 94			
47174602	**Good identification**						**47174602**
Clostridium clostridioforme		100.0	0.36	SAL 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47174612	**Good identification**						**47174612**
Clostridium clostridioforme		100.0	0.48	SAL 88			
47174622	**Good identification**						**47174622**
Clostridium clostridioforme		100.0	0.36	SAL 88			
47174632	**Good identification**						**47174632**
Clostridium clostridioforme		100.0	0.48	SAL 88			
47174702	**Good identification**						**47174702**
Clostridium clostridioforme		100.0	0.45	SAL 88			
47174712	**Very good identification**						**47174712**
Clostridium clostridioforme		100.0	0.57	SAL 88			
47174722	**Good identification**						**47174722**
Clostridium clostridioforme		97.15	0.45	SAL 88			
Parabacteroides distasonis		2.22	0.23	RAF 94			
Bacteroides thetaiotaomicron		0.63	-.03				
47174732	**Very good identification**						**47174732**
Clostridium clostridioforme		99.73	0.57	SAL 88			
Parabacteroides distasonis		0.27	0.18	RAF 94			
47175000	**Good identification**						**47175000**
					COC		
Atopobium parvulum		93.02	0.64	LAC 99	99		
Lactobacillus jensenii		5.33	0.44		1		
Actinomyces israelii		1.30	0.46		1		
47175002	**Low discrimination**						**47175002**
					COC		
Atopobium parvulum		74.08	0.51	LAC 99	99		
Lactobacillus jensenii		16.02	0.41		1		
Actinomyces israelii		8.77	0.49		1		
47175010	**Low discrimination**						**47175010**
					COC		
Lactobacillus jensenii		61.04	0.36	SAL 95	1		
Atopobium parvulum		32.31	0.30	XYL 1	99		
Actinomyces israelii		6.66	0.32		1		
47175020	**Acceptable identification**						**47175020**
					COC		
Atopobium parvulum		91.04	0.51	LAC 99	99		
Lactobacillus jensenii		7.38	0.34		1		
Clostridium perfringens		0.88	0.17		1		
47175022	**Low discrimination**						**47175022**
					COC	GRA	
Atopobium parvulum		62.38	0.38	LAC 99	99	99	
Lactobacillus jensenii		19.12	0.31		1	99	
Prevotella oralis		9.29	0.21		1	1	
47175042	**Low discrimination**						**47175042**
					COC		
Actinomyces israelii		78.94	0.42	SAL 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		17.31	0.17		99		
Lactobacillus jensenii		3.75	0.07		1		
47175100	**Excellent identification**						**47175100**
					COC	SPO	
Atopobium parvulum		99.91	0.98		99	1	
Actinomyces israelii		0.06	0.56		1	1	
Clostridium butyricum		0.02	0.28		1	99	
47175101	**Very good identification**						**47175101**
					COC	SPO	
Atopobium parvulum		99.82	0.64	URE 1	99	1	
Clostridium chauvoei		0.10	0.12		1	99	
Actinomyces israelii		0.06	0.22		1	1	
47175102	**Very good identification**						**47175102**
					COC		
Atopobium parvulum		99.03	0.85		99		
Actinomyces israelii		0.47	0.59		1		
Bifidobacterium adolescentis		0.34	0.50		1		
47175103	**Good identification**						**47175103**
					COC		
Atopobium parvulum		99.09	0.51	URE 1	99		
Actinomyces israelii		0.47	0.25		1		
Bifidobacterium adolescentis		0.34	0.16		1		
47175110	**Very good identification**						**47175110**
					COC	SPO	
Atopobium parvulum		98.74	0.64	MLZ 1	99	1	
Actinomyces israelii		0.82	0.42		1	1	
Clostridium tertium		0.16	0.15		1	99	
47175112	**Low discrimination**						**47175112**
					COC	GRA	
Atopobium parvulum		52.94	0.51	MLZ 1	99	99	
Bifidobacterium adolescentis		21.89	0.52		1	99	
Leptotrichia buccalis		18.10	0.32		1	1	
47175120	**Very good identification**						**47175120**
					COC	SPO	
Atopobium parvulum		99.73	0.85		99	1	
Clostridium perfringens		0.07	0.32		1	84	
Clostridium paraputrificum		0.07	0.22		1	99	
47175121	**Good identification**						**47175121**
					COC	SPO	
Atopobium parvulum		99.81	0.51	URE 1	99	1	
Clostridium chauvoei		0.19	0.04		1	99	
47175122	**Good identification**						**47175122**
					COC	GRA	
Atopobium parvulum		93.85	0.72		99	99	
Prevotella oralis		2.21	0.42		1	1	
Parabacteroides distasonis		1.73	0.53		1	1	
47175123	**Good identification**						**47175123**
					COC	GRA	
Atopobium parvulum		95.15	0.38	URE 1	99	99	
Prevotella oralis		2.24	0.07		1	1	
Parabacteroides distasonis		1.76	0.18		1	1	
47175130	**Good identification**						**47175130**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Atopobium parvulum		96.90	0.51	MLZ 1	99	1		
Clostridium tertium		2.64	0.24		1	99		
Actinomyces israelii		0.29	0.22		1	1		
47175132	**Low discrimination**							**47175132**
					COC	GRA	SPO	
Atopobium parvulum		33.87	0.38	MLZ 1	99	99	1	
Parabacteroides distasonis		33.36	0.48		1	1	1	
Clostridium tertium		15.69	0.32		1	99	99	
47175140	**Very good identification**							**47175140**
					COC	SPO		
Atopobium parvulum		97.71	0.64	SOR 1	99	1		
Actinomyces israelii		2.12	0.49		1	1		
Clostridium butyricum		0.17	0.10		1	99		
47175142	**Low discrimination**							**47175142**
					COC			
Atopobium parvulum		80.98	0.51	SOR 1	99			
Actinomyces israelii		14.92	0.52		1			
Bifidobacterium adolescentis		3.39	0.35		1			
47175152	**Low discrimination**							**47175152**
					COC			
Bifidobacterium adolescentis		55.44	0.36	RAF 92	1			
Actinomyces israelii		29.86	0.38	ARA 87	1			
Atopobium parvulum		10.96	0.17	SOR 11	99			
47175160	**Good identification**							**47175160**
					COC			
Atopobium parvulum		97.42	0.51	SOR 1	99			
Clostridium perfringens		1.32	0.20		1			
Actinomyces israelii		0.76	0.29		1			
47175162	**Low discrimination**							**47175162**
					COC			
Atopobium parvulum		81.71	0.38	SOR 1	99			
Actinomyces israelii		5.45	0.32		1			
Clostridium perfringens		2.95	0.14		1			
47175212	**Good identification**							**47175212**
Clostridium clostridioforme		98.09	0.35	SAL 88				
Actinomyces israelii		1.91	0.15	RAF 94				
47175232	**Acceptable identification**							**47175232**
Clostridium clostridioforme		92.86	0.35	SAL 88				
Parabacteroides distasonis		7.14	0.22	RAF 94				
47175300	**Very good identification**							**47175300**
					COC	SPO		
Atopobium parvulum		99.52	0.64	RHA 1	99	1		
Actinomyces israelii		0.42	0.37		1	1		
Clostridium butyricum		0.06	0.02		1	99		
47175302	**Low discrimination**							**47175302**
					COC			
Atopobium parvulum		89.86	0.51	RHA 1	99			
Clostridium clostridioforme		4.52	0.33		1			
Actinomyces israelii		3.20	0.40		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47175312	**Low discrimination**						**47175312**
					COC		
Clostridium clostridioforme		89.01	0.44	SAL 88	1		
Atopobium parvulum		3.92	0.17	RAF 94	99		
Actinomyces israelii		2.07	0.26	ARA 94	1		
47175320	**Good identification**						**47175320**
					COC	GRA	
Atopobium parvulum		96.73	0.51	RHA 1	99	99	
Bacteroides vulgatus		1.13	0.09		1	1	
Parabacteroides distasonis		0.95	0.27		1	1	
47175322	**Low discrimination**						**47175322**
					COC	GRA	
Parabacteroides distasonis		69.75	0.61	RAF 99	1	1	
Atopobium parvulum		12.71	0.38		99	99	
Prevotella oralis		12.70	0.35		1	1	
47175332	**Low discrimination**						**47175332**
					COC	GRA	
Parabacteroides distasonis		68.99	0.56	RAF 99	1	1	
Clostridium clostridioforme		30.34	0.44		1	50	
Atopobium parvulum		0.24	0.04		99	99	
47175400	**Low discrimination**						**47175400**
					COC	SPO	
Actinomyces israelii		38.29	0.44	SAL 92	1	1	
Atopobium parvulum		33.92	0.30		99	1	
Clostridium butyricum		25.85	0.20		1	99	
47175402	**Low discrimination**						**47175402**
Actinomyces israelii		53.49	0.47	SAL 92			
Clostridium clostridioforme		17.74	0.30				
Bifidobacterium adolescentis		12.62	0.30				
47175412	**Low discrimination**						**47175412**
Clostridium clostridioforme		77.53	0.41	SAL 88			
Bifidobacterium adolescentis		14.81	0.32	RAF 94			
Actinomyces israelii		7.67	0.33	RHA 88			
47175432	**Good identification**						**47175432**
Clostridium clostridioforme		99.27	0.41	SAL 88			
Parabacteroides distasonis		0.73	0.10	RAF 94			
47175440	**Low discrimination**						**47175440**
					SPO		
Actinomyces israelii		86.88	0.37	SAL 92	1		
Clostridium butyricum		13.12	0.02		99		
47175442	**Low discrimination**						**47175442**
					SPO		
Actinomyces israelii		89.76	0.40	SAL 92	1		
Bifidobacterium adolescentis		6.73	0.15		1		
Clostridium butyricum		3.51	-.05		99		
47175500	**Low discrimination**						**47175500**
					COC	SPO	
Atopobium parvulum		80.96	0.64	ARA 1	99	1	
Clostridium butyricum		14.96	0.44		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		3.69	0.55		1	1		
47175502	**Low discrimination**							**47175502**
					COC	SPO		
Bifidobacterium adolescentis		54.26	0.64	RAF 92	1	1		
Atopobium parvulum		23.97	0.51		99	1		
Clostridium butyricum		9.76	0.37		1	99		
47175512	**Low discrimination**							**47175512**
Bifidobacterium adolescentis		82.19	0.66	RAF 92				
Clostridium clostridioforme		14.55	0.50					
Actinomyces israelii		1.72	0.44					
47175520	**Low discrimination**							**47175520**
					COC	GRA	SPO	
Atopobium parvulum		87.47	0.51	ARA 1	99	99	1	
Clostridium butyricum		7.96	0.26		1	99	99	
Bacteroides vulgatus		1.51	0.12		1	1	1	
47175522	**Low discrimination**							**47175522**
					COC	GRA		
Parabacteroides distasonis		32.38	0.49	RAF 99	1	1		
Atopobium parvulum		28.88	0.38		99	99		
Clostridium clostridioforme		17.60	0.39		1	50		
47175532	**Low discrimination**							**47175532**
					GRA			
Clostridium clostridioforme		71.86	0.50	SAL 88	50			
Parabacteroides distasonis		15.62	0.44	RAF 94	1			
Bifidobacterium adolescentis		4.10	0.32	RHA 88	99			
47175540	**Low discrimination**							**47175540**
					COC	SPO		
Actinomyces israelii		39.94	0.48	SAL 92	1	1		
Clostridium butyricum		36.18	0.26		1	99		
Atopobium parvulum		22.75	0.30		99	1		
47175542	**Low discrimination**							**47175542**
					SPO			
Bifidobacterium adolescentis		58.65	0.49	RAF 92	1			
Actinomyces israelii		31.59	0.51	SOR 11	1			
Clostridium butyricum		7.42	0.19		99			
47175552	**Acceptable identification**							**47175552**
Bifidobacterium adolescentis		92.10	0.50	RAF 92				
Actinomyces israelii		6.07	0.37	SOR 11				
Clostridium clostridioforme		1.33	0.16					
47175602	**Good identification**							**47175602**
Clostridium clostridioforme		97.00	0.45	SAL 88				
Actinomyces israelii		3.00	0.28	RAF 94				
47175612	**Good identification**							**47175612**
Clostridium clostridioforme		99.90	0.56	SAL 88				
Actinomyces israelii		0.10	0.14	RAF 94				
47175622	**Good identification**							**47175622**
Clostridium clostridioforme		96.14	0.45	SAL 88				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		2.45	0.23	RAF 94			
Bacteroides thetaiotaomicron		1.41	0.03				
47175632	**Good identification**						**47175632**
Clostridium clostridioforme		99.58	0.56	SAL 88			
Parabacteroides distasonis		0.30	0.18	RAF 94			
Bacteroides thetaiotaomicron		0.13	-.04				
47175702	**Acceptable identification**						**47175702**
Clostridium clostridioforme		90.54	0.54	SAL 88			
Actinomyces israelii		3.35	0.38	RAF 94			
Bifidobacterium adolescentis		2.63	0.30				
47175712	**Very good identification**						**47175712**
Clostridium clostridioforme		99.02	0.65	SAL 88			
Bifidobacterium adolescentis		0.77	0.32	RAF 94			
Actinomyces israelii		0.12	0.24				
47175722	**Low discrimination**						**47175722**
Clostridium clostridioforme		48.34	0.54	SAL 88			
Parabacteroides distasonis		36.38	0.57	RAF 94			
Bacteroides thetaiotaomicron		10.40	0.32				
47175732	**Low discrimination**						**47175732**
Clostridium clostridioforme		89.26	0.65	SAL 88			
Parabacteroides distasonis		7.94	0.52	RAF 94			
Bacteroides thetaiotaomicron		1.64	0.25				
47176102	**Low discrimination**						**47176102**
					COC	SPO	
Actinomyces israelii		49.00	0.38	SAL 92	1	1	
Clostridium tertium		27.70	0.19	GAL 99	1	99	
Atopobium parvulum		18.25	0.17		99	1	
47176502	**Low discrimination**						**47176502**
Actinomyces israelii		54.25	0.36	SAL 92			
Bifidobacterium adolescentis		45.75	0.29	GAL 99			
47177000	**Low discrimination**						**47177000**
					COC	SPO	
Actinomyces israelii		87.21	0.59	SAL 92	1	1	
Atopobium parvulum		11.16	0.30		99	1	
Clostridium tertium		1.00	0.11		1	99	
47177002	**Very good identification**						**47177002**
					COC	SPO	
Actinomyces israelii		95.67	0.62	SAL 92	1	1	
Clostridium tertium		2.19	0.19		1	99	
Atopobium parvulum		1.44	0.17		99	1	
47177010	**Low discrimination**						**47177010**
					SPO		
Actinomyces israelii		86.79	0.45	SAL 92	1		
Clostridium tertium		11.79	0.15	MLZ 13	99		
Lactobacillus jensenii		1.42	0.02		1		
47177012	**Low discrimination**						**47177012**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		76.58	0.48	SAL 92	1		
Clostridium tertium		20.81	0.24	MLZ 13	99		
Bifidobacterium adolescentis		2.61	0.16		1		
47177020	**Low discrimination**						**47177020**
					COC	SPO	
Actinomyces israelii		52.90	0.38	SAL 92	1	1	
Clostridium tertium		28.39	0.19	NAG 6	1	99	
Atopobium parvulum		18.71	0.17		99	1	
47177040	**Good identification**						**47177040**
Actinomyces israelii		100.0	0.52	SAL 92			
47177042	**Good identification**						**47177042**
Actinomyces israelii		100.0	0.55	SAL 92			
47177050	**Good identification**						**47177050**
Actinomyces israelii		100.0	0.38	SAL 92			
47177052	**Good identification**						**47177052**
Actinomyces israelii		100.0	0.41	SAL 92			
47177100	**Low discrimination**						**47177100**
					COC	SPO	
Atopobium parvulum		71.03	0.64	MAN 1	99	1	
Actinomyces israelii		22.43	0.69		1	1	
Clostridium tertium		6.34	0.45		1	99	
47177102	**Low discrimination**						**47177102**
					COC	SPO	
Actinomyces israelii		48.94	0.72	SAL 92	1	1	
Clostridium tertium		27.66	0.53		1	99	
Atopobium parvulum		18.23	0.51		99	1	
47177103	**Low discrimination**						**47177103**
					COC	SPO	
Actinomyces israelii		48.01	0.38	SAL 92	1	1	
Clostridium tertium		27.14	0.19	URE 1	1	99	
Atopobium parvulum		17.89	0.17		99	1	
47177110	**Low discrimination**						**47177110**
					COC	SPO	
Clostridium tertium		73.27	0.50	SAL 99	1	99	
Actinomyces israelii		21.79	0.55		1	1	
Atopobium parvulum		4.66	0.30		99	1	
47177112	**Low discrimination**						**47177112**
					SPO		
Clostridium tertium		77.97	0.58	SAL 99	99		
Actinomyces israelii		11.59	0.58		1		
Bifidobacterium adolescentis		9.78	0.50		1		
47177120	**Low discrimination**						**47177120**
					COC	SPO	
Clostridium tertium		57.49	0.53	SAL 99	1	99	
Atopobium parvulum		37.89	0.51		99	1	
Actinomyces israelii		4.33	0.49		1	1	
47177122	**Good identification**						**47177122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium tertium		92.15	0.62	SAL 99	1	99	
Atopobium parvulum		3.57	0.38		99	1	
Actinomyces israelii		3.47	0.52		1	1	
47177130	**Very good identification**						**47177130**
					COC	SPO	
Clostridium tertium		98.94	0.58	SAL 99	1	99	
Actinomyces israelii		0.63	0.34		1	1	
Atopobium parvulum		0.37	0.17		99	1	
47177132	**Very good identification**						**47177132**
					SPO		
Clostridium tertium		99.40	0.66	SAL 99	99		
Actinomyces israelii		0.31	0.37		1		
Bifidobacterium dentium		0.20	0.10		1		
47177140	**Good identification**						**47177140**
					COC	SPO	
Actinomyces israelii		91.64	0.62	SAL 92	1	1	
Atopobium parvulum		7.54	0.30		99	1	
Clostridium tertium		0.67	0.11		1	99	
47177142	**Good identification**						**47177142**
					SPO		
Actinomyces israelii		94.60	0.65	SAL 92	1		
Bifidobacterium adolescentis		3.10	0.33		1		
Clostridium tertium		1.39	0.19		99		
47177150	**Acceptable identification**						**47177150**
					COC	SPO	
Actinomyces israelii		90.57	0.48	SAL 92	1	1	
Clostridium tertium		7.91	0.15	MLZ 13	1	99	
Blautia producta		1.52	-.03		99	1	
47177152	**Low discrimination**						**47177152**
					SPO		
Actinomyces israelii		69.31	0.51	SAL 92	1		
Bifidobacterium adolescentis		18.58	0.35	MLZ 13	1		
Clostridium tertium		12.11	0.24		99		
47177160	**Low discrimination**						**47177160**
					COC	SPO	
Actinomyces israelii		63.60	0.42	SAL 92	1	1	
Clostridium tertium		21.94	0.19	NAG 6	1	99	
Atopobium parvulum		14.46	0.17		99	1	
47177162	**Low discrimination**						**47177162**
					COC	SPO	
Actinomyces israelii		58.25	0.45	SAL 92	1	1	
Clostridium tertium		40.19	0.27	NAG 6	1	99	
Atopobium parvulum		1.56	0.04		99	1	
47177200	**Good identification**						**47177200**
Actinomyces israelii		100.0	0.39	SAL 92			
47177202	**Good identification**						**47177202**
Actinomyces israelii		100.0	0.43	SAL 92			
47177242	**Good identification**						**47177242**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.35	SAL 92			
47177300	**Low discrimination**						**47177300**
					COC	SPO	
Actinomyces israelii		68.36	0.50	SAL 92	1	1	
Atopobium parvulum		29.05	0.30	RHA 7	99	1	
Clostridium tertium		2.59	0.11		1	99	
47177302	**Low discrimination**						**47177302**
					COC	SPO	
Actinomyces israelii		87.74	0.53	SAL 92	1	1	
Clostridium tertium		6.66	0.19	RHA 7	1	99	
Atopobium parvulum		4.39	0.17		99	1	
47177310	**Low discrimination**						**47177310**
					SPO		
Actinomyces israelii		68.91	0.36	SAL 92	1		
Clostridium tertium		31.09	0.15	RHA 7	99		
47177312	**Low discrimination**						**47177312**
					SPO		
Actinomyces israelii		47.79	0.39	SAL 92	1		
Clostridium tertium		43.13	0.24	RHA 7	99		
Bifidobacterium adolescentis		5.41	0.16	MLZ 13	1		
47177340	**Good identification**						**47177340**
Actinomyces israelii		100.0	0.43	SAL 92			
47177342	**Good identification**						**47177342**
Actinomyces israelii		100.0	0.46	SAL 92			
47177400	**Very good identification**						**47177400**
					SPO		
Actinomyces israelii		98.69	0.57	SAL 92	1		
Clostridium butyricum		1.31	0.04		99		
47177402	**Very good identification**						**47177402**
					SPO		
Actinomyces israelii		96.39	0.60	SAL 92	1		
Bifidobacterium adolescentis		3.28	0.29		1		
Clostridium butyricum		0.33	-.03		99		
47177410	**Good identification**						**47177410**
Actinomyces israelii		100.0	0.43	SAL 92			
47177412	**Low discrimination**						**47177412**
Actinomyces israelii		70.97	0.46	SAL 92			
Bifidobacterium adolescentis		19.78	0.30	MLZ 13			
Bifidobacterium dentium		7.97	0.06				
47177420	**Good identification**						**47177420**
Actinomyces israelii		100.0	0.37	SAL 92			
47177422	**Good identification**						**47177422**
Actinomyces israelii		100.0	0.40	SAL 92			
47177440	**Good identification**						**47177440**
Actinomyces israelii		100.0	0.50	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47177442	**Good identification**						**47177442**
Actinomyces israelii		98.93	0.53	SAL 92			
Bifidobacterium adolescentis		1.07	0.13				
47177450	**Acceptable identification**						**47177450**
					COC		
Actinomyces israelii		92.42	0.36	SAL 92	1		
Blautia producta		7.58	-.03	MLZ 13	99		
47177452	**Acceptable identification**						**47177452**
Actinomyces israelii		91.86	0.39	SAL 92			
Bifidobacterium adolescentis		8.14	0.15	MLZ 13			
47177500	**Low discrimination**						**47177500**
					COC	SPO	
Actinomyces israelii		88.12	0.68	SAL 92	1	1	
Clostridium butyricum		7.00	0.28		1	99	
Atopobium parvulum		3.45	0.30		99	1	
47177502	**Low discrimination**						**47177502**
					SPO		
Actinomyces israelii		53.19	0.71	SAL 92	1		
Bifidobacterium adolescentis		44.85	0.63		1		
Clostridium butyricum		1.10	0.21		99		
47177503	**Low discrimination**						**47177503**
Actinomyces israelii		54.25	0.36	SAL 92			
Bifidobacterium adolescentis		45.75	0.29	URE 1			
47177510	**Low discrimination**						**47177510**
Actinomyces israelii		62.72	0.53	SAL 92			
Bifidobacterium dentium		25.87	0.23	MLZ 13			
Bifidobacterium adolescentis		6.56	0.30				
47177512	**Low discrimination**						**47177512**
Bifidobacterium adolescentis		63.97	0.64	RAF 92			
Bifidobacterium dentium		25.78	0.40				
Actinomyces israelii		9.27	0.56				
47177520	**Low discrimination**						**47177520**
					SPO		
Actinomyces israelii		71.55	0.47	SAL 92	1		
Clostridium tertium		11.73	0.19	NAG 6	99		
Clostridium butyricum		7.74	0.10		99		
47177522	**Low discrimination**						**47177522**
					SPO		
Actinomyces israelii		60.61	0.50	SAL 92	1		
Clostridium tertium		19.88	0.27	NAG 6	99		
Bifidobacterium adolescentis		8.09	0.29		1		
47177532	**Acceptable identification**						**47177532**
					SPO		
Bifidobacterium dentium		92.50	0.44	SAL 99	1		
Clostridium tertium		4.66	0.32	RAF 95	99		
Bifidobacterium adolescentis		1.30	0.30		1		
47177540	**Low discrimination**						**47177540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		89.06	0.61	SAL 92	1	1	
Blautia producta		9.00	0.24		99	1	
Clostridium butyricum		1.58	0.10		1	99	
47177542	**Low discrimination**						**47177542**
					COC		
Actinomyces israelii		78.12	0.64	SAL 92	1		
Bifidobacterium adolescentis		20.94	0.47		1		
Blautia producta		0.59	0.07		99		
47177552	**Low discrimination**						**47177552**
					COC		
Bifidobacterium adolescentis		64.20	0.49	RAF 92	1		
Actinomyces israelii		29.28	0.49	SOR 11	1		
Blautia producta		4.40	0.15		99		
47177560	**Low discrimination**						**47177560**
					COC		
Actinomyces israelii		85.04	0.40	SAL 92	1		
Blautia producta		14.96	0.07	NAG 6	99		
47177562	**Good identification**						**47177562**
Actinomyces israelii		95.93	0.43	SAL 92			
Bifidobacterium adolescentis		4.07	0.13	NAG 6			
47177600	**Good identification**						**47177600**
Actinomyces israelii		100.0	0.38	SAL 92			
47177602	**Acceptable identification**						**47177602**
Actinomyces israelii		94.56	0.41	SAL 92			
Clostridium clostridioforme		5.44	0.10	RHA 7			
47177700	**Good identification**						**47177700**
					SPO		
Actinomyces israelii		96.84	0.48	SAL 92	1		
Clostridium butyricum		3.16	0.02	RHA 7	99		
47177702	**Low discrimination**						**47177702**
Actinomyces israelii		85.48	0.51	SAL 92			
Bifidobacterium adolescentis		9.68	0.29	RHA 7			
Clostridium clostridioforme		4.12	0.19				
47177742	**Low discrimination**						**47177742**
					COC		
Actinomyces israelii		88.07	0.44	SAL 92	1		
Blautia producta		8.76	0.07	RHA 7	99		
Bifidobacterium adolescentis		3.17	0.13		1		
47200000	**Low discrimination**						**47200000**
					GRA	SPO	
Fusobacterium mortiferum		44.87	0.52	FRU 99	1	1	
Eubacterium tortuosum		29.06	0.32		99	1	
Clostridium bifermentans		10.17	0.27		99	99	
47200100	**Very good identification**						**47200100**
					GRA		
Fusobacterium mortiferum		99.72	0.58	FRU 99	1		
Eubacterium tortuosum		0.28	-.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47200102	**Good identification**						**47200102**
Fusobacterium mortiferum		100.0	0.42	FRU 99			
47201000	**Very good identification**						**47201000**
					GRA		
Eubacterium tortuosum		97.57	0.66	MNS 1	99		
Fusobacterium mortiferum		2.28	0.55		1		
Atopobium minutum		0.05	0.20		99		
47201002	**Good identification**						**47201002**
					GRA		
Eubacterium tortuosum		97.93	0.53	MNS 1	99		
Fusobacterium mortiferum		1.44	0.38		1		
Atopobium minutum		0.17	0.17		99		
47201020	**Good identification**						**47201020**
					GRA	SPO	
Eubacterium tortuosum		99.58	0.53	MNS 1	99	1	
Fusobacterium mortiferum		0.13	0.20		1	1	
Clostridium barati		0.11	0.04		99	99	
47201022	**Good identification**						**47201022**
					SPO		
Eubacterium tortuosum		97.68	0.40	MNS 1	1		
Clostridium barati		1.76	0.12		99		
Atopobium minutum		0.33	0.09		1		
47201100	**Low discrimination**						**47201100**
					GRA		
Fusobacterium mortiferum		83.18	0.61	FRU 99	1		
Eubacterium tortuosum		15.39	0.32		99		
Lactobacillus catenaformis		0.69	0.15		99		
47201102	**Low discrimination**						**47201102**
					GRA	SPO	
Fusobacterium mortiferum		65.96	0.45	FRU 99	1	1	
Eubacterium tortuosum		19.38	0.19	BGL 10	99	1	
Clostridium barati		6.10	0.12		99	99	
47204100	**Good identification**						**47204100**
Fusobacterium mortiferum		100.0	0.39	FRU 99			
47205100	**Good identification**						**47205100**
					GRA		
Fusobacterium mortiferum		97.58	0.42	FRU 99	1		
Eubacterium tortuosum		2.42	-.03	XYL 7	99		
47210000	**Very good identification**						**47210000**
					GRA	SPO	
Fusobacterium mortiferum		99.25	0.86		1	1	
Clostridium difficile		0.37	0.28		99	99	
Eubacterium limosum		0.12	0.18		99	1	
47210001	**Good identification**						**47210001**
Fusobacterium mortiferum		100.0	0.52	URE 1			
47210002	**Very good identification**						**47210002**
					GRA	SPO	
Fusobacterium mortiferum		98.52	0.69	BGL 10	1	1	
Clostridium barati		0.56	0.16		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium minutum		0.50	0.25		99	1	
47210003	**Good identification**						**47210003**
Fusobacterium mortiferum		100.0	0.35	URE 1			
47210010	**Low discrimination**						**47210010**
					GRA	SPO	
Clostridium difficile		64.02	0.40	SUC 1	99	99	
Fusobacterium mortiferum		35.70	0.52	MAN 90	1	1	
Lactobacillus jensenii		0.29	0.06		99	1	
47210012	**Low discrimination**						**47210012**
					GRA	SPO	
Fusobacterium mortiferum		82.58	0.35	MLZ 1	1	1	
Clostridium difficile		13.46	0.06	BGL 10	99	99	
Lactobacillus jensenii		3.95	0.03		99	1	
47210020	**Low discrimination**						**47210020**
					GRA	SPO	
Fusobacterium mortiferum		89.83	0.52	NAG 1	1	1	
Clostridium barati		5.61	0.16		99	99	
Atopobium minutum		2.48	0.20		99	1	
47210040	**Good identification**						**47210040**
					GRA	SPO	
Fusobacterium mortiferum		97.69	0.52	SOR 1	1	1	
Clostridium difficile		1.89	0.06		99	99	
Clostridium botulinum biovar A		0.42	-.02		99	99	
47210042	**Good identification**						**47210042**
					GRA	SPO	
Fusobacterium mortiferum		95.34	0.35	SOR 1	1	1	
Clostridium barati		4.66	-.02	BGL 10	99	99	
47210100	**Excellent identification**						**47210100**
					GRA	SPO	
Fusobacterium mortiferum		99.95	0.92		1	1	
Clostridium barati		0.03	0.16		99	99	
Lactobacillus catenaformis		0.01	0.15		99	1	
47210101	**Very good identification**						**47210101**
Fusobacterium mortiferum		100.0	0.58	URE 1			
47210102	**Very good identification**						**47210102**
					GRA	SPO	
Fusobacterium mortiferum		99.08	0.76	BGL 10	1	1	
Clostridium barati		0.72	0.24		99	99	
Lactobacillus catenaformis		0.11	0.15		99	1	
47210103	**Good identification**						**47210103**
Fusobacterium mortiferum		100.0	0.42	URE 1			
47210110	**Very good identification**						**47210110**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.58	MLZ 1	1	1	
Clostridium difficile		0.77	0.06		99	99	
47210112	**Good identification**						**47210112**
Fusobacterium mortiferum		100.0	0.42	MLZ 1			
47210120	**Good identification**						**47210120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		91.39	0.58	NAG 1	1	1	
Clostridium barati		7.33	0.24		99	99	
Clostridium paraputrificum		0.42	-.03		99	99	
47210140	**Very good identification**						**47210140**
					GRA	SPO	
Fusobacterium mortiferum		99.77	0.58	SOR 1	1	1	
Clostridium barati		0.23	-.02		99	99	
47210142	**Acceptable identification**						**47210142**
					GRA	SPO	
Fusobacterium mortiferum		94.09	0.42	SOR 1	1	1	
Clostridium barati		5.91	0.06	BGL 10	99	99	
47210200	**Good identification**						**47210200**
Fusobacterium mortiferum		100.0	0.52	RHA 1			
47210202	**Good identification**						**47210202**
					GRA	SPO	
Fusobacterium mortiferum		95.34	0.35	RHA 1	1	1	
Clostridium barati		4.66	-.02	BGL 10	99	99	
47210300	**Very good identification**						**47210300**
					GRA	SPO	
Fusobacterium mortiferum		99.77	0.58	RHA 1	1	1	
Clostridium barati		0.23	-.02		99	99	
47210302	**Acceptable identification**						**47210302**
					GRA	SPO	
Fusobacterium mortiferum		94.09	0.42	RHA 1	1	1	
Clostridium barati		5.91	0.06	BGL 10	99	99	
47210400	**Good identification**						**47210400**
					COC	GRA	
Fusobacterium mortiferum		99.47	0.52	ARA 1	1	1	
Gemella morbillorum		0.53	0.03		99	99	
47210402	**Good identification**						**47210402**
Fusobacterium mortiferum		100.0	0.35	ARA 1			
47210500	**Very good identification**						**47210500**
Fusobacterium mortiferum		100.0	0.58	ARA 1			
47210502	**Good identification**						**47210502**
Fusobacterium mortiferum		100.0	0.42	ARA 1			
47211000	**Very good identification**						**47211000**
					GRA		
Fusobacterium mortiferum		96.41	0.89		1		
Lactobacillus jensenii		1.54	0.48		99		
Lactobacillus catenaformis		1.24	0.47		99		
47211001	**Good identification**						**47211001**
					GRA		
Fusobacterium mortiferum		96.32	0.55	URE 1	1		
Lactobacillus jensenii		1.54	0.14		99		
Lactobacillus catenaformis		1.24	0.12		99		
47211002	**Low discrimination**						**47211002**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		76.85	0.72	BGL 10	1		
Lactobacillus catenaformis		8.91	0.47		99		
Lactobacillus jensenii		7.36	0.45		99		
47211003	**Low discrimination**						**47211003**
					GRA		
Fusobacterium mortiferum		75.26	0.38	URE 1	1		
Lactobacillus catenaformis		8.72	0.12	BGL 10	99		
Lactobacillus jensenii		7.21	0.11		99		
47211010	**Low discrimination**						**47211010**
					GRA		
Fusobacterium mortiferum		64.16	0.55	MLZ 1	1		
Lactobacillus jensenii		33.79	0.40		99		
Lactobacillus catenaformis		0.83	0.12		99		
47211012	**Low discrimination**						**47211012**
					GRA		
Lactobacillus jensenii		72.27	0.37	MLT 95	99		
Fusobacterium mortiferum		22.87	0.38	CEL 95	1		
Lactobacillus catenaformis		2.65	0.12		99		
47211020	**Low discrimination**						**47211020**
					GRA	SPO	
Fusobacterium mortiferum		37.14	0.55	NAG 1	1	1	
Clostridium barati		29.36	0.38		99	99	
Lactobacillus jensenii		14.67	0.38		99	1	
47211022	**Low discrimination**						**47211022**
					SPO		
Clostridium barati		73.63	0.46	CEL 99	99		
Lactobacillus catenaformis		9.89	0.36		1		
Lactobacillus jensenii		8.17	0.35		1		
47211040	**Acceptable identification**						**47211040**
					GRA	SPO	
Fusobacterium mortiferum		94.69	0.55	SOR 1	1	1	
Clostridium barati		2.17	0.11		99	99	
Lactobacillus jensenii		1.51	0.14		99	1	
47211042	**Low discrimination**						**47211042**
					GRA	SPO	
Fusobacterium mortiferum		54.63	0.38	SOR 1	1	1	
Clostridium barati		33.81	0.20	BGL 10	99	99	
Lactobacillus catenaformis		6.33	0.12		99	1	
47211100	**Very good identification**						**47211100**
					GRA		
Fusobacterium mortiferum		98.21	0.95		1		
Lactobacillus catenaformis		0.81	0.50		99		
Collinsella aerofaciens		0.54	0.32		99		
47211101	**Good identification**						**47211101**
					GRA	SPO	
Fusobacterium mortiferum		92.22	0.61	URE 1	1	1	
Clostridium chauvoei		5.63	0.29		99	99	
Lactobacillus catenaformis		0.76	0.15		99	1	
47211102	**Low discrimination**						**47211102**
					GRA	SPO	
Fusobacterium mortiferum		80.69	0.79	BGL 10	1	1	
Clostridium barati		7.46	0.46		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		6.01	0.50		99	1	
47211103	**Low discrimination**						**47211103**
					GRA	SPO	
Fusobacterium mortiferum		72.76	0.45	URE 1	1	1	
Clostridium chauvoei		7.06	0.16	BGL 10	99	99	
Clostridium barati		6.72	0.12		99	99	
47211110	**Very good identification**						**47211110**
					GRA		
Fusobacterium mortiferum		97.97	0.61	MLZ 1	1		
Lactobacillus catenaformis		0.81	0.15		99		
Collinsella aerofaciens		0.54	-.03		99		
47211112	**Low discrimination**						**47211112**
					GRA	SPO	
Fusobacterium mortiferum		76.91	0.45	MLZ 1	1	1	
Clostridium barati		7.11	0.12	BGL 10	99	99	
Lactobacillus catenaformis		5.73	0.15		99	1	
47211120	**Low discrimination**						**47211120**
					GRA	SPO	
Clostridium barati		32.31	0.46	CEL 99	99	99	
Fusobacterium mortiferum		31.78	0.61		1	1	
Clostridium paraputrificum		9.65	0.32		99	99	
47211122	**Low discrimination**						**47211122**
					GRA	SPO	
Clostridium barati		59.24	0.54	CEL 99	99	99	
Bifidobacterium longum subsp. infantis		21.03	0.44		99	1	
Prevotella oralis		6.17	0.38		1	1	
47211140	**Very good identification**						**47211140**
					GRA	SPO	
Fusobacterium mortiferum		95.71	0.61	SOR 1	1	1	
Clostridium barati		2.82	0.20		99	99	
Lactobacillus catenaformis		0.79	0.15		99	1	
47211142	**Low discrimination**						**47211142**
					GRA	SPO	
Fusobacterium mortiferum		49.39	0.45	SOR 1	1	1	
Clostridium barati		39.29	0.28	BGL 10	99	99	
Lactobacillus acidophilus		4.91	0.15		99	1	
47211162	**Low discrimination**						**47211162**
					SPO		
Clostridium barati		80.72	0.36	CEL 99	99		
Clostridium perfringens		7.53	0.23	SOR 8	84		
Lactobacillus acidophilus		3.37	0.15		1		
47211200	**Acceptable identification**						**47211200**
					GRA	SPO	
Fusobacterium mortiferum		94.69	0.55	RHA 1	1	1	
Clostridium barati		2.17	0.11		99	99	
Lactobacillus jensenii		1.51	0.14		99	1	
47211202	**Low discrimination**						**47211202**
					GRA	SPO	
Fusobacterium mortiferum		54.63	0.38	RHA 1	1	1	
Clostridium barati		33.81	0.20	BGL 10	99	99	
Lactobacillus catenaformis		6.33	0.12		99	1	
47211300	**Low discrimination**						**47211300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		89.73	0.61	RHA 1	1	1	
Eubacterium contortum		6.40	0.29		99	1	
Clostridium barati		2.64	0.20		99	99	
47211302	**Low discrimination**						**47211302**
					GRA	SPO	
Fusobacterium mortiferum		46.22	0.45	RHA 1	1	1	
Clostridium barati		36.77	0.28	BGL 10	99	99	
Eubacterium contortum		5.23	0.16		99	1	
47211322	**Low discrimination**						**47211322**
					GRA	SPO	
Clostridium barati		60.01	0.36	CEL 99	99	99	
Prevotella oralis		30.79	0.32	RHA 8	1	1	
Parabacteroides distasonis		3.64	0.29		1	1	
47211400	**Good identification**						**47211400**
					GRA		
Fusobacterium mortiferum		95.72	0.55	ARA 1	1		
Lactobacillus jensenii		1.53	0.14		99		
Lactobacillus catenaformis		1.23	0.12		99		
47211402	**Low discrimination**						**47211402**
					GRA	SPO	
Fusobacterium mortiferum		72.06	0.38	ARA 1	1	1	
Lactobacillus catenaformis		8.35	0.12	BGL 10	99	1	
Clostridium innocuum		7.51	0.09		99	99	
47211500	**Very good identification**						**47211500**
					GRA		
Fusobacterium mortiferum		98.31	0.61	ARA 1	1		
Lactobacillus catenaformis		0.81	0.15		99		
Collinsella aerofaciens		0.54	-.03		99		
47211502	**Low discrimination**						**47211502**
					GRA	SPO	
Fusobacterium mortiferum		72.99	0.45	ARA 1	1	1	
Bifidobacterium adolescentis		10.79	0.27	BGL 10	99	1	
Clostridium barati		6.75	0.12		99	99	
47212000	**Low discrimination**						**47212000**
					GRA	SPO	
Clostridium difficile		59.33	0.45	SUC 1	99	99	
Fusobacterium mortiferum		17.95	0.52		1	1	
Eubacterium limosum		13.15	0.32		99	1	
47212010	**Very good identification**						**47212010**
					GRA	SPO	
Clostridium difficile		99.76	0.56	SUC 1	99	99	
Clostridium innocuum		0.13	0.11		99	99	
Fusobacterium mortiferum		0.06	0.17		1	1	
47212100	**Very good identification**						**47212100**
					GRA	SPO	
Fusobacterium mortiferum		97.85	0.58	MAN 1	1	1	
Clostridium difficile		1.40	0.11		99	99	
Eubacterium limosum		0.31	-.03		99	1	
47212102	**Acceptable identification**						**47212102**
					GRA	SPO	
Fusobacterium mortiferum		92.36	0.42	MAN 1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium barati		5.80	0.06	BGL 10	99	99		
Clostridium innocuum		1.85	0.00		99	99		
47213000	**Low discrimination**							**47213000**
					GRA	SPO		
Clostridium innocuum		85.47	0.56	CEL 99	99	99		
Fusobacterium mortiferum		13.16	0.55		1	1		
Clostridium barati		0.30	0.11		99	99		
47213002	**Good identification**							**47213002**
					GRA	SPO		
Clostridium innocuum		96.82	0.56	CEL 99	99	99		
Fusobacterium mortiferum		1.66	0.38		1	1		
Clostridium barati		1.03	0.20		99	99		
47213100	**Good identification**							**47213100**
					GRA	SPO		
Fusobacterium mortiferum		92.66	0.61	MAN 1	1	1		
Clostridium barati		2.73	0.20		99	99		
Clostridium innocuum		2.60	0.22		99	99		
47213101	**Low discrimination**							**47213101**
					GRA			
Actinomyces naeslundii		54.08	0.40	ESL 12	99			
Fusobacterium mortiferum		45.92	0.27	MLT 94	1			
47213102	**Low discrimination**							**47213102**
					GRA	SPO		
Fusobacterium mortiferum		43.04	0.45	MAN 1	1	1		
Clostridium barati		34.24	0.28	BGL 10	99	99		
Clostridium innocuum		10.89	0.22		99	99		
47213103	**Good identification**							**47213103**
Actinomyces naeslundii		100.0	0.40	ESL 12				
47213121	**Good identification**							**47213121**
Actinomyces naeslundii		100.0	0.40	ESL 12				
47213122	**Low discrimination**							**47213122**
					SPO			
Clostridium barati		87.05	0.36	CEL 99	99			
Bifidobacterium longum subsp. infantis		3.59	0.10	MAN 8	1			
Actinomyces naeslundii		2.60	0.37		1			
47213123	**Low discrimination**							**47213123**
					SPO			
Actinomyces naeslundii		81.62	0.40	ESL 12	1			
Clostridium barati		18.38	0.02	MLT 94	99			
47213400	**Good identification**							**47213400**
					GRA	SPO		
Clostridium innocuum		99.13	0.43	CEL 99	99	99		
Fusobacterium mortiferum		0.87	0.20		1	1		
47213402	**Good identification**							**47213402**
Clostridium innocuum		100.0	0.43	CEL 99				
47214000	**Very good identification**							**47214000**
					COC	GRA	SPO	
Fusobacterium mortiferum		99.67	0.67	XYL 7	1	1	1	
Clostridium difficile		0.26	0.06		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Gemella morbillorum		0.07	0.03		99	99	1	
47214002	**Good identification**							**47214002**
Fusobacterium mortiferum		100.0	0.50	XYL 7				
47214100	**Very good identification**							**47214100**
Fusobacterium mortiferum		100.0	0.73	XYL 7				
47214101	**Good identification**							**47214101**
Fusobacterium mortiferum		100.0	0.39	XYL 7				
47214102	**Very good identification**							**47214102**
					GRA			
Fusobacterium mortiferum		99.10	0.57	XYL 7	1			
Bifidobacterium longum subsp. infantis		0.90	0.04	BGL 10	99			
47214110	**Good identification**							**47214110**
Fusobacterium mortiferum		100.0	0.39	XYL 7				
47214120	**Good identification**							**47214120**
Fusobacterium mortiferum		100.0	0.39	XYL 7				
47214140	**Good identification**							**47214140**
Fusobacterium mortiferum		100.0	0.39	XYL 7				
47214500	**Good identification**							**47214500**
Fusobacterium mortiferum		100.0	0.39	XYL 7				
47215000	**Very good identification**							**47215000**
					GRA			
Fusobacterium mortiferum		99.51	0.70	XYL 7	1			
Lactobacillus jensenii		0.21	0.14		99			
Lactobacillus catenaformis		0.17	0.12		99			
47215001	**Good identification**							**47215001**
Fusobacterium mortiferum		100.0	0.35	XYL 7				
47215002	**Acceptable identification**							**47215002**
					GRA			
Fusobacterium mortiferum		94.66	0.53	XYL 7	1			
Lactobacillus catenaformis		1.47	0.12	BGL 10	99			
Bifidobacterium longum subsp. infantis		1.34	0.04		99			
47215010	**Acceptable identification**							**47215010**
					GRA			
Fusobacterium mortiferum		93.40	0.35	XYL 7	1			
Lactobacillus jensenii		6.60	0.06	MLZ 1	99			
47215020	**Low discrimination**							**47215020**
					GRA	SPO		
Fusobacterium mortiferum		83.20	0.35	XYL 7	1	1		
Clostridium barati		8.83	0.04	NAG 1	99	99		
Lactobacillus jensenii		4.41	0.04		99	1		
47215040	**Good identification**							**47215040**
Fusobacterium mortiferum		100.0	0.35	XYL 7				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47215100	**Very good identification**					**47215100**
					GRA	
Fusobacterium mortiferum		97.68	0.76	XYL 7	1	
Eubacterium contortum		0.93	0.29		99	
Bifidobacterium longum subsp. infantis		0.65	0.21		99	
47215101	**Good identification**					**47215101**
Fusobacterium mortiferum		100.0	0.42	XYL 7		
47215102	**Low discrimination**					**47215102**
					GRA	
Fusobacterium mortiferum		59.82	0.60	XYL 7	1	
Bifidobacterium longum subsp. infantis		36.01	0.38	BGL 10	99	
Bifidobacterium adolescentis		1.19	0.27		99	
47215110	**Good identification**					**47215110**
Fusobacterium mortiferum		100.0	0.42	XYL 7		
47215122	**Low discrimination**					**47215122**
					GRA	
Bifidobacterium longum subsp. infantis		86.22	0.43	MLT 99	99	
Parabacteroides distasonis		4.70	0.37	RAF 95	1	
Prevotella oralis		3.82	0.23		1	
47215140	**Good identification**					**47215140**
Fusobacterium mortiferum		100.0	0.42	XYL 7		
47215300	**Very good identification**					**47215300**
					GRA	
Eubacterium contortum		98.95	0.63	MLT 99	99	
Fusobacterium mortiferum		1.06	0.42		1	
47215302	**Good identification**					**47215302**
					GRA	
Eubacterium contortum		98.82	0.50	MLT 99	99	
Fusobacterium mortiferum		0.66	0.25		1	
Bifidobacterium longum subsp. infantis		0.40	0.04		99	
47215320	**Good identification**					**47215320**
					GRA	
Eubacterium contortum		98.61	0.50	MLT 99	99	
Bacteroides vulgatus		1.28	0.11		1	
Parabacteroides distasonis		0.12	0.11		1	
47215322	**Low discrimination**					**47215322**
					GRA	
Eubacterium contortum		55.90	0.37	MLT 99	99	
Parabacteroides distasonis		36.87	0.45		1	
Prevotella oralis		4.28	0.16		1	
47215332	**Good identification**					**47215332**
					GRA	
Parabacteroides distasonis		97.23	0.41	RAF 99	1	
Eubacterium contortum		2.77	0.03		99	
47215400	**Good identification**					**47215400**
Fusobacterium mortiferum		100.0	0.35	XYL 7		
47215500	**Low discrimination**					**47215500**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		56.35	0.42	XYL 7	1		
Eubacterium contortum		43.65	0.27	ARA 1	99		
47215502	**Low discrimination**						**47215502**
					GRA		
Bifidobacterium adolescentis		82.27	0.41	MLT 99	99		
Eubacterium contortum		7.70	0.14	RAF 92	99		
Fusobacterium mortiferum		6.26	0.25		1		
47215512	**Good identification**						**47215512**
Bifidobacterium adolescentis		98.70	0.42	MLT 99			
Clostridium clostridioforme		1.31	0.07	RAF 92			
47215700	**Very good identification**						**47215700**
					GRA		
Eubacterium contortum		99.96	0.61	MLT 99	99		
Bacteroides vulgatus		0.04	-.05		1		
47215702	**Good identification**						**47215702**
Eubacterium contortum		99.72	0.48	MLT 99			
Clostridium clostridioforme		0.28	0.10				
47215720	**Low discrimination**						**47215720**
					GRA		
Eubacterium contortum		82.61	0.48	MLT 99	99		
Bacteroides vulgatus		17.39	0.30		1		
47215722	**Low discrimination**						**47215722**
					GRA		
Eubacterium contortum		65.32	0.36	MLT 99	99		
Parabacteroides distasonis		32.28	0.42		1		
Clostridium clostridioforme		1.04	0.10		50		
47215732	**Low discrimination**						**47215732**
					GRA		
Parabacteroides distasonis		76.32	0.37	RAF 99	1		
Clostridium clostridioforme		20.78	0.22		50		
Eubacterium contortum		2.90	0.01		99		
47216100	**Good identification**						**47216100**
Fusobacterium mortiferum		100.0	0.39	MAN 1			
47217100	**Acceptable identification**						**47217100**
					GRA	SPO	
Fusobacterium mortiferum		92.08	0.42	MAN 1	1	1	
Actinomyces israelii		6.11	0.24	XYL 7	99	1	
Clostridium innocuum		1.81	0.00		99	99	
47217502	**Acceptable identification**						**47217502**
Bifidobacterium adolescentis		93.62	0.39	MLT 99			
Actinomyces israelii		6.39	0.25	RAF 92			
47217512	**Good identification**						**47217512**
Bifidobacterium adolescentis		100.0	0.41	MLT 99			
47220000	**Low discrimination**						**47220000**
					GRA	SPO	
Fusobacterium mortiferum		82.64	0.43	FRU 99	1	1	
Clostridium botulinum biovar A		15.55	0.17		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium tortuosum		1.81	-.03		99	1		
47220100	**Good identification**							**47220100**
Fusobacterium mortiferum		100.0	0.49	FRU 99				
47221100	**Acceptable identification**							**47221100**
					COC	GRA		
Fusobacterium mortiferum		92.88	0.52	FRU 99	1	1		
Atopobium parvulum		6.54	0.21		99	99		
Eubacterium tortuosum		0.58	-.03		1	99		
47221102	**Low discrimination**							**47221102**
					COC	GRA		
Fusobacterium mortiferum		83.13	0.36	FRU 99	1	1		
Atopobium parvulum		9.30	0.08	BGL 10	99	99		
Lactobacillus acidophilus		7.58	0.05		1	99		
47230000	**Very good identification**							**47230000**
					GRA	CAT		
Fusobacterium mortiferum		99.38	0.77		1	5		
Propionibacterium granulosum		0.28	0.30		99	90		
Lactobacillus jensenii		0.24	0.22		99	1		
47230001	**Good identification**							**47230001**
Fusobacterium mortiferum		100.0	0.43	URE 1				
47230002	**Very good identification**							**47230002**
					GRA	CAT		
Fusobacterium mortiferum		97.79	0.60	BGL 10	1	5		
Lactobacillus jensenii		1.43	0.19		99	1		
Propionibacterium granulosum		0.62	0.20		99	90		
47230010	**Low discrimination**							**47230010**
					GRA	SPO		
Fusobacterium mortiferum		63.68	0.43	MLZ 1	1	1		
Clostridium difficile		28.78	0.21		99	99		
Lactobacillus jensenii		5.10	0.14		99	1		
47230020	**Low discrimination**							**47230020**
					GRA	CAT		
Fusobacterium mortiferum		87.08	0.43	NAG 1	1	5		
Propionibacterium granulosum		6.10	0.20		99	90		
Lactobacillus jensenii		5.23	0.12		99	1		
47230040	**Good identification**							**47230040**
					GRA	SPO		
Fusobacterium mortiferum		98.06	0.43	SOR 1	1	1		
Clostridium botulinum biovar B		1.94	-.03		99	99		
47230100	**Very good identification**							**47230100**
					COC	GRA	SPO	
Fusobacterium mortiferum		99.89	0.83		1	1	1	
Atopobium parvulum		0.11	0.21		99	99	1	
Clostridium barati		0.01	-.02		1	99	99	
47230101	**Good identification**							**47230101**
Fusobacterium mortiferum		100.0	0.49	URE 1				
47230102	**Very good identification**							**47230102**
					COC	GRA	SPO	
Fusobacterium mortiferum		99.48	0.67	BGL 10	1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium barati		0.21	0.06		1	99	99	
Atopobium parvulum		0.17	0.08		99	99	1	
47230110	**Good identification**							**47230110**
Fusobacterium mortiferum		100.0	0.49	MLZ 1				
47230120	**Acceptable identification**							**47230120**
					GRA	SPO		
Fusobacterium mortiferum		91.85	0.49	NAG 1	1	1		
Clostridium perfringens		2.39	0.11		99	84		
Clostridium barati		2.15	0.06		99	99		
47230140	**Good identification**							**47230140**
Fusobacterium mortiferum		100.0	0.49	SOR 1				
47230200	**Good identification**							**47230200**
Fusobacterium mortiferum		100.0	0.43	RHA 1				
47230300	**Good identification**							**47230300**
Fusobacterium mortiferum		100.0	0.49	RHA 1				
47230400	**Good identification**							**47230400**
Fusobacterium mortiferum		100.0	0.43	ARA 1				
47230500	**Good identification**							**47230500**
Fusobacterium mortiferum		100.0	0.49	ARA 1				
47231000	**Low discrimination**							**47231000**
					GRA	CAT		
Fusobacterium mortiferum		85.74	0.80		1	5		
Lactobacillus jensenii		13.74	0.56		99	1		
Propionibacterium granulosum		0.16	0.30		99	90		
47231001	**Low discrimination**							**47231001**
					GRA			
Fusobacterium mortiferum		83.83	0.46	URE 1	1			
Lactobacillus jensenii		13.44	0.22		99			
Actinomyces naeslundii		2.74	0.32		99			
47231002	**Low discrimination**							**47231002**
					GRA	SPO		
Fusobacterium mortiferum		49.49	0.64	BGL 10	1	1		
Lactobacillus jensenii		47.58	0.53		99	1		
Clostridium barati		1.04	0.20		99	99		
47231010	**Low discrimination**							**47231010**
					GRA	CAT		
Lactobacillus jensenii		83.76	0.48	MLT 95	99	1		
Fusobacterium mortiferum		15.84	0.46	CEL 95	1	5		
Propionibacterium granulosum		0.40	0.15		99	90		
47231012	**Good identification**							**47231012**
					GRA			
Lactobacillus jensenii		96.09	0.45	MLT 95	99			
Fusobacterium mortiferum		3.03	0.29	CEL 95	1			
Leptotrichia buccalis		0.88	-.03		1			
47231020	**Low discrimination**							**47231020**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus jensenii		67.42	0.46	MLT 95	99			
Fusobacterium mortiferum		17.00	0.46	CEL 95	1			
Clostridium perfringens		9.29	0.30		99			
47231022	**Low discrimination**							**47231022**
					SPO			
Lactobacillus jensenii		66.76	0.43	MLT 95	1			
Clostridium barati		17.43	0.28	CEL 95	99			
Clostridium perfringens		6.49	0.25		84			
47231030	**Good identification**							**47231030**
					GRA			
Lactobacillus jensenii		99.24	0.38	MLT 95	99			
Fusobacterium mortiferum		0.76	0.12	CEL 95	1			
47231040	**Low discrimination**							**47231040**
					GRA			
Fusobacterium mortiferum		86.19	0.46	SOR 1	1			
Lactobacillus jensenii		13.81	0.22		99			
47231100	**Good identification**							**47231100**
					COC	GRA	SPO	
Fusobacterium mortiferum		93.14	0.86		1	1	1	
Atopobium parvulum		6.56	0.55		99	99	1	
Clostridium barati		0.09	0.20		1	99	99	
47231101	**Low discrimination**							**47231101**
					COC	GRA		
Fusobacterium mortiferum		88.26	0.52	URE 1	1	1		
Atopobium parvulum		6.22	0.21		99	99		
Actinomyces naeslundii		3.52	0.40		1	99		
47231102	**Low discrimination**							**47231102**
					COC	GRA		
Fusobacterium mortiferum		80.33	0.70	BGL 10	1	1		
Atopobium parvulum		8.99	0.42		99	99		
Lactobacillus acidophilus		7.32	0.39		1	99		
47231103	**Low discrimination**							**47231103**
					COC	GRA		
Fusobacterium mortiferum		62.60	0.36	URE 1	1	1		
Actinomyces naeslundii		22.43	0.40	BGL 10	1	99		
Atopobium parvulum		7.00	0.08		99	99		
47231110	**Acceptable identification**							**47231110**
					COC	GRA		
Fusobacterium mortiferum		90.32	0.52	MLZ 1	1	1		
Atopobium parvulum		6.36	0.21		99	99		
Lactobacillus jensenii		2.07	0.14		1	99		
47231120	**Low discrimination**							**47231120**
					COC	SPO		
Clostridium perfringens		28.87	0.45	MLT 99	1	84		
Clostridium septicum		23.02	0.37		1	99		
Atopobium parvulum		20.69	0.42		99	1		
47231121	**Low discrimination**							**47231121**
					SPO			
Actinomyces naeslundii		39.39	0.40	ESL 12	1			
Clostridium perfringens		17.16	0.11	MLT 94	84			
Clostridium septicum		13.68	0.03	RAF 92	99			
47231122	**Low discrimination**							**47231122**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Lactobacillus acidophilus		29.31	0.39	MLT 94	1			
Clostridium barati		25.96	0.36	CEL 99	99			
Clostridium perfringens		23.63	0.40		84			
47231123	**Low discrimination**							**47231123**
					SPO			
Actinomyces naeslundii		59.37	0.40	ESL 12	1			
Lactobacillus acidophilus		15.10	0.05	MLT 94	1			
Clostridium barati		13.37	0.02	RAF 92	99			
47231140	**Acceptable identification**							**47231140**
					COC	GRA		
Fusobacterium mortiferum		91.42	0.52	SOR 1	1	1		
Atopobium parvulum		6.44	0.21		99	99		
Clostridium perfringens		0.93	0.07		1	99		
47231200	**Low discrimination**							**47231200**
					GRA			
Fusobacterium mortiferum		86.19	0.46	RHA 1	1			
Lactobacillus jensenii		13.81	0.22		99			
47231300	**Acceptable identification**							**47231300**
					COC	GRA	SPO	
Fusobacterium mortiferum		92.68	0.52	RHA 1	1	1	1	
Atopobium parvulum		6.53	0.21		99	99	1	
Clostridium barati		0.80	0.01		1	99	99	
47231302	**Low discrimination**							**47231302**
					GRA	SPO		
Fusobacterium mortiferum		49.45	0.36	RHA 1	1	1		
Lactobacillus acidophilus		33.57	0.20	BGL 10	99	1		
Clostridium barati		11.45	0.10		99	99		
47231322	**Low discrimination**							**47231322**
					GRA	SPO		
Parabacteroides distasonis		53.41	0.40	RAF 99	1	1		
Clostridium barati		16.79	0.18	XYL 90	99	99		
Lactobacillus acidophilus		16.41	0.20		99	1		
47231332	**Good identification**							**47231332**
Parabacteroides distasonis		100.0	0.36	RAF 99				
47231400	**Low discrimination**							**47231400**
					GRA	SPO		
Fusobacterium mortiferum		85.19	0.46	ARA 1	1	1		
Lactobacillus jensenii		13.65	0.22		99	1		
Clostridium innocuum		1.16	0.01		99	99		
47231500	**Acceptable identification**							**47231500**
					COC	GRA		
Fusobacterium mortiferum		93.42	0.52	ARA 1	1	1		
Atopobium parvulum		6.58	0.21		99	99		
47231502	**Low discrimination**							**47231502**
					COC	GRA		
Fusobacterium mortiferum		65.24	0.36	ARA 1	1	1		
Bifidobacterium adolescentis		21.52	0.23	BGL 10	1	99		
Atopobium parvulum		7.30	0.08		99	99		
47231722	**Good identification**							**47231722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		95.59	0.37	RAF 99			
Clostridium clostridioforme		4.41	0.08	XYL 90			
47232000	**Low discrimination**						**47232000**
					GRA	SPO	
Fusobacterium mortiferum		37.42	0.43	MAN 1	1	1	
Clostridium difficile		31.18	0.25		99	99	
Clostridium innocuum		22.56	0.26		99	99	
47232010	**Good identification**						**47232010**
					SPO	CAT	
Clostridium difficile		98.69	0.37	SUC 1	99	1	
Propionibacterium granulosum		0.70	0.13	TRE 7	1	90	
Clostridium innocuum		0.61	0.03		99	1	
47232100	**Good identification**						**47232100**
Fusobacterium mortiferum		100.0	0.49	MAN 1			
47233000	**Low discrimination**						**47233000**
					GRA	SPO	
Clostridium innocuum		85.04	0.48	CEL 99	99	99	
Fusobacterium mortiferum		11.14	0.46		1	1	
Lactobacillus jensenii		1.78	0.22		99	1	
47233002	**Good identification**						**47233002**
					GRA	SPO	
Clostridium innocuum		95.91	0.48	CEL 99	99	99	
Fusobacterium mortiferum		1.40	0.29		1	1	
Lactobacillus jensenii		1.34	0.19		99	1	
47233021	**Good identification**						**47233021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47233023	**Good identification**						**47233023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47233100	**Low discrimination**						**47233100**
					COC	GRA	
Fusobacterium mortiferum		84.46	0.52	MAN 1	1	1	
Atopobium parvulum		5.95	0.21		99	99	
Actinomyces naeslundii		4.35	0.42		1	99	
47233101	**Low discrimination**						**47233101**
					GRA		
Actinomyces naeslundii		88.44	0.45	ESL 12	99		
Fusobacterium mortiferum		11.56	0.18	MLT 94	1		
47233102	**Low discrimination**						**47233102**
					GRA		
Fusobacterium mortiferum		32.30	0.36	MAN 1	1		
Lactobacillus acidophilus		15.34	0.17	BGL 10	99		
Actinomyces naeslundii		14.98	0.42		99		
47233103	**Good identification**						**47233103**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47233120	**Low discrimination**						**47233120**
					SPO		
Actinomyces naeslundii		25.55	0.42	ESL 12	1		
Clostridium tertium		22.14	0.17	MLT 94	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		12.92	0.11	RAF 92	92		
47233121	**Good identification**						**47233121**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47233123	**Good identification**						**47233123**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47233400	**Good identification**						**47233400**
					GRA	SPO	
Clostridium innocuum		99.26	0.35	CEL 99	99	99	
Fusobacterium mortiferum		0.74	0.12		1	1	
47233402	**Good identification**						**47233402**
Clostridium innocuum		100.0	0.35	CEL 99			
47234000	**Very good identification**						**47234000**
Fusobacterium mortiferum		100.0	0.58	XYL 7			
47234002	**Good identification**						**47234002**
Fusobacterium mortiferum		100.0	0.41	XYL 7			
47234100	**Very good identification**						**47234100**
					COC	GRA	
Fusobacterium mortiferum		97.44	0.64	XYL 7	1	1	
Atopobium parvulum		2.57	0.25		99	99	
47234102	**Good identification**						**47234102**
					COC	GRA	
Fusobacterium mortiferum		95.99	0.48	XYL 7	1	1	
Atopobium parvulum		4.01	0.13	BGL 10	99	99	
47235000	**Good identification**						**47235000**
					COC	GRA	
Fusobacterium mortiferum		94.12	0.61	XYL 7	1	1	
Atopobium parvulum		3.86	0.25		99	99	
Lactobacillus jensenii		2.02	0.22		1	99	
47235002	**Low discrimination**						**47235002**
					COC	GRA	
Fusobacterium mortiferum		82.93	0.44	XYL 7	1	1	
Lactobacillus jensenii		10.70	0.19	BGL 10	1	99	
Atopobium parvulum		5.40	0.13		99	99	
47235100	**Low discrimination**						**47235100**
					COC	GRA	
Atopobium parvulum		63.42	0.60	MLT 99	99	99	
Fusobacterium mortiferum		36.50	0.67		1	1	
Actinomyces israelii		0.05	0.21		1	99	
47235102	**Low discrimination**						**47235102**
					COC	GRA	
Atopobium parvulum		71.39	0.47	MLT 99	99	99	
Fusobacterium mortiferum		25.87	0.51		1	1	
Bifidobacterium adolescentis		1.15	0.23		1	99	
47235120	**Good identification**						**47235120**
					COC	GRA	
Atopobium parvulum		95.25	0.47	MLT 99	99	99	
Fusobacterium mortiferum		3.14	0.33		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.72	0.11		1	99	
47235122	**Low discrimination**						**47235122**
					COC	GRA	
Parabacteroides distasonis		60.69	0.48	RAF 99	1	1	
Atopobium parvulum		33.21	0.34		99	99	
Bifidobacterium longum subsp. infantis		2.47	0.08		1	99	
47235132	**Good identification**						**47235132**
					COC	GRA	
Parabacteroides distasonis		98.98	0.44	RAF 99	1	1	
Atopobium parvulum		1.02	0.00		99	99	
47235322	**Very good identification**						**47235322**
					GRA		
Parabacteroides distasonis		99.28	0.56	RAF 99	1		
Eubacterium contortum		0.33	0.03		99		
Clostridium clostridioforme		0.20	0.05		50		
47235332	**Good identification**						**47235332**
Parabacteroides distasonis		98.31	0.52	RAF 99			
Clostridium clostridioforme		1.69	0.16				
47235502	**Low discrimination**						**47235502**
					COC	SPO	
Bifidobacterium adolescentis		76.41	0.38	MLT 99	1	1	
Atopobium parvulum		7.19	0.13	RAF 92	99	1	
Clostridium butyricum		5.15	0.03		1	99	
47235512	**Low discrimination**						**47235512**
Bifidobacterium adolescentis		87.30	0.39	MLT 99			
Clostridium clostridioforme		12.70	0.22	RAF 92			
47235522	**Good identification**						**47235522**
					COC	GRA	
Parabacteroides distasonis		97.07	0.45	RAF 99	1	1	
Clostridium clostridioforme		2.06	0.10		1	50	
Atopobium parvulum		0.88	0.00		99	99	
47235532	**Low discrimination**						**47235532**
Parabacteroides distasonis		84.79	0.40	RAF 99			
Clostridium clostridioforme		15.21	0.22				
47235712	**Good identification**						**47235712**
Clostridium clostridioforme		97.75	0.36	MLT 99			
Parabacteroides distasonis		2.25	0.14	RAF 94			
47235722	**Acceptable identification**						**47235722**
					GRA		
Parabacteroides distasonis		94.67	0.53	RAF 99	1		
Clostridium clostridioforme		4.91	0.25		50		
Eubacterium contortum		0.43	0.01		99		
47235732	**Low discrimination**						**47235732**
Parabacteroides distasonis		69.53	0.48	RAF 99			
Clostridium clostridioforme		30.48	0.36				
47237102	**Low discrimination**						**47237102**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		43.41	0.37	MLT 99	1		
Clostridium tertium		28.35	0.19	CEL 93	99		
Bifidobacterium adolescentis		13.79	0.22		1		
47237502	**Low discrimination**						**47237502**
Bifidobacterium adolescentis		72.21	0.36	MLT 99			
Actinomyces israelii		27.79	0.35	RAF 92			
47237512	**Low discrimination**						**47237512**
Bifidobacterium adolescentis		83.46	0.38	MLT 99			
Bifidobacterium dentium		12.62	0.06	RAF 92			
Actinomyces israelii		3.93	0.21				
47240000	**Low discrimination**						**47240000**
					GRA	SPO	
Fusobacterium mortiferum		69.88	0.43	FRU 99	1	1	
Eubacterium tortuosum		15.09	0.15		99	1	
Clostridium innocuum		5.66	0.12		99	99	
47240100	**Low discrimination**						**47240100**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.50	FRU 99	1	1	
Clostridium barati		7.03	0.16		99	99	
Lactobacillus catenaformis		3.26	0.15		99	1	
47241000	**Low discrimination**						**47241000**
					SPO		
Eubacterium tortuosum		65.25	0.50	MNS 1	1		
Lactobacillus catenaformis		17.52	0.47	CEL 10	1		
Clostridium innocuum		4.70	0.34		99		
47241002	**Low discrimination**						**47241002**
					SPO		
Lactobacillus catenaformis		36.40	0.47	MLT 90	1		
Eubacterium tortuosum		23.93	0.37	FRU 99	1		
Clostridium barati		22.57	0.38		99		
47241020	**Low discrimination**						**47241020**
					SPO		
Eubacterium tortuosum		41.14	0.37	MNS 1	1		
Clostridium barati		38.81	0.38	CEL 10	99		
Lactobacillus catenaformis		15.64	0.36		1		
47241022	**Low discrimination**						**47241022**
					SPO		
Clostridium barati		81.41	0.46	FRU 99	99		
Lactobacillus catenaformis		10.94	0.36		1		
Eubacterium tortuosum		5.08	0.24		1		
47241100	**Low discrimination**						**47241100**
					SPO		
Lactobacillus catenaformis		39.78	0.50	MLT 90	1		
Collinsella aerofaciens		26.32	0.32	FRU 99	1		
Clostridium barati		16.45	0.38		99		
47241102	**Low discrimination**						**47241102**
					SPO		
Clostridium barati		50.10	0.46	FRU 99	99		
Lactobacillus catenaformis		40.39	0.50		1		
Lactobacillus acidophilus		4.41	0.30		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47241120	**Low discrimination**						**47241120**
					SPO		
Clostridium barati		63.67	0.46	FRU 99	99		
Clostridium paraputrificum		19.01	0.32		99		
Lactobacillus catenaformis		12.83	0.39		1		
47241122	**Low discrimination**						**47241122**
					SPO		
Clostridium barati		87.29	0.54	FRU 99	99		
Lactobacillus catenaformis		5.87	0.39		1		
Clostridium paraputrificum		3.72	0.25		99		
47241162	**Acceptable identification**						**47241162**
					SPO		
Clostridium barati		91.56	0.36	FRU 99	99		
Lactobacillus acidophilus		3.82	0.15	SOR 8	1		
Clostridium paraputrificum		3.38	0.06		99		
47241322	**Acceptable identification**						**47241322**
					GRA	SPO	
Clostridium barati		94.65	0.36	FRU 99	99	99	
Lactobacillus acidophilus		2.40	0.11	RHA 8	99	1	
Prevotella oralis		1.74	0.07		1	1	
47242000	**Good identification**						**47242000**
Clostridium innocuum		100.0	0.46	FRU 95			
47242002	**Good identification**						**47242002**
Clostridium innocuum		99.80	0.46	FRU 95			
Clostridium barati		0.20	-.02	GAL 95			
47243000	**Very good identification**						**47243000**
					SPO		
Clostridium innocuum		99.73	0.68	FRU 95	99		
Eubacterium tortuosum		0.14	0.15		1		
Clostridium barati		0.07	0.11		99		
47243002	**Very good identification**						**47243002**
					SPO		
Clostridium innocuum		99.73	0.68	FRU 95	99		
Clostridium barati		0.20	0.20		99		
Lactobacillus catenaformis		0.04	0.12		1		
47243010	**Good identification**						**47243010**
Clostridium innocuum		100.0	0.45	FRU 95			
47243012	**Good identification**						**47243012**
Clostridium innocuum		100.0	0.45	FRU 95			
47243122	**Acceptable identification**						**47243122**
					SPO		
Clostridium barati		93.80	0.36	FRU 99	99		
Lactobacillus acidophilus		1.67	0.08	MAN 8	1		
Clostridium tertium		1.57	0.09		99		
47243400	**Good identification**						**47243400**
Clostridium innocuum		100.0	0.55	FRU 95			
47243402	**Good identification**						**47243402**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.55	FRU 95			
47247000	**Good identification**						**47247000**
Clostridium innocuum		100.0	0.46	FRU 95			
47247002	**Good identification**						**47247002**
Clostridium innocuum		100.0	0.46	FRU 95			
47250000	**Low discrimination**						**47250000**
					GRA	SPO	
Fusobacterium mortiferum		86.89	0.78		1	1	
Clostridium barati		5.42	0.42		99	99	
Lactobacillus catenaformis		5.04	0.47		99	1	
47250001	**Low discrimination**						**47250001**
					GRA	SPO	
Fusobacterium mortiferum		86.98	0.43	URE 1	1	1	
Clostridium barati		5.43	0.08		99	99	
Lactobacillus catenaformis		5.04	0.12		99	1	
47250002	**Low discrimination**						**47250002**
					GRA	SPO	
Clostridium barati		49.04	0.50	GAL 95	99	99	
Fusobacterium mortiferum		29.10	0.61		1	1	
Lactobacillus catenaformis		15.18	0.47		99	1	
47250010	**Low discrimination**						**47250010**
					GRA	SPO	
Fusobacterium mortiferum		52.02	0.43	MLZ 1	1	1	
Lactobacillus jensenii		23.66	0.28		99	1	
Clostridium difficile		14.73	0.18		99	99	
47250020	**Low discrimination**						**47250020**
					GRA	SPO	
Clostridium barati		86.79	0.50	GAL 95	99	99	
Lactobacillus catenaformis		6.71	0.36		99	1	
Fusobacterium mortiferum		4.68	0.43		1	1	
47250022	**Very good identification**						**47250022**
					SPO		
Clostridium barati		96.72	0.58	GAL 95	99		
Lactobacillus catenaformis		2.49	0.36		1		
Lactobacillus jensenii		0.40	0.23		1		
47250040	**Low discrimination**						**47250040**
					GRA	SPO	
Fusobacterium mortiferum		61.55	0.43	SOR 1	1	1	
Clostridium barati		33.08	0.24		99	99	
Lactobacillus catenaformis		3.57	0.12		99	1	
47250062	**Good identification**						**47250062**
					SPO		
Clostridium barati		99.70	0.40	GAL 95	99		
Lactobacillus catenaformis		0.30	0.02	SOR 8	1		
47250100	**Low discrimination**						**47250100**
					GRA	SPO	
Fusobacterium mortiferum		87.53	0.84		1	1	
Clostridium barati		7.03	0.50		99	99	
Lactobacillus catenaformis		3.26	0.50		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47250101	**Low discrimination**						**47250101**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.50	URE 1	1	1	
Clostridium barati		7.03	0.16		99	99	
Lactobacillus catenaformis		3.26	0.15		99	1	
47250102	**Low discrimination**						**47250102**
					GRA	SPO	
Clostridium barati		60.79	0.58	GAL 95	99	99	
Fusobacterium mortiferum		28.06	0.68		1	1	
Lactobacillus catenaformis		9.41	0.50		99	1	
47250110	**Low discrimination**						**47250110**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.50	MLZ 1	1	1	
Clostridium barati		7.03	0.16		99	99	
Lactobacillus catenaformis		3.26	0.15		99	1	
47250120	**Low discrimination**						**47250120**
					GRA	SPO	
Clostridium barati		86.95	0.58	GAL 95	99	99	
Clostridium paraputrificum		4.98	0.32		99	99	
Fusobacterium mortiferum		3.65	0.50		1	1	
47250122	**Very good identification**						**47250122**
					SPO		
Clostridium barati		97.14	0.66	GAL 95	99		
Lactobacillus catenaformis		1.25	0.39		1		
Clostridium paraputrificum		0.80	0.25		99		
47250140	**Low discrimination**						**47250140**
					GRA	SPO	
Fusobacterium mortiferum		57.05	0.50	SOR 1	1	1	
Clostridium barati		39.42	0.32		99	99	
Lactobacillus catenaformis		2.13	0.15		99	1	
47250142	**Acceptable identification**						**47250142**
					GRA	SPO	
Clostridium barati		91.28	0.40	GAL 95	99	99	
Fusobacterium mortiferum		4.89	0.33	SOR 8	1	1	
Lactobacillus acidophilus		2.19	0.15		99	1	
47250160	**Acceptable identification**						**47250160**
					GRA	SPO	
Clostridium barati		94.43	0.40	GAL 95	99	99	
Clostridium paraputrificum		4.69	0.13	SOR 8	99	99	
Fusobacterium mortiferum		0.46	0.16		1	1	
47250162	**Good identification**						**47250162**
					SPO		
Clostridium barati		98.26	0.48	GAL 95	99		
Lactobacillus acidophilus		0.79	0.15	SOR 8	1		
Clostridium paraputrificum		0.70	0.06		99		
47250200	**Low discrimination**						**47250200**
					GRA	SPO	
Fusobacterium mortiferum		61.55	0.43	RHA 1	1	1	
Clostridium barati		33.08	0.24		99	99	
Lactobacillus catenaformis		3.57	0.12		99	1	
47250222	**Good identification**						**47250222**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		99.70	0.40	GAL 95	99		
Lactobacillus catenaformis		0.30	0.02	RHA 8	1		
47250300	**Low discrimination**						**47250300**
					GRA	SPO	
Fusobacterium mortiferum		57.05	0.50	RHA 1	1	1	
Clostridium barati		39.42	0.32		99	99	
Lactobacillus catenaformis		2.13	0.15		99	1	
47250302	**Acceptable identification**						**47250302**
					GRA	SPO	
Clostridium barati		92.06	0.40	GAL 95	99	99	
Fusobacterium mortiferum		4.94	0.33	RHA 8	1	1	
Lactobacillus catenaformis		1.66	0.15		99	1	
47250320	**Good identification**						**47250320**
					GRA	SPO	
Clostridium barati		98.42	0.40	GAL 95	99	99	
Clostridium paraputrificum		0.66	-.03	RHA 8	99	99	
Fusobacterium mortiferum		0.48	0.16		1	1	
47250322	**Good identification**						**47250322**
					GRA	SPO	
Clostridium barati		98.93	0.48	GAL 95	99	99	
Lactobacillus acidophilus		0.48	0.11	RHA 8	99	1	
Prevotella oralis		0.35	0.07		1	1	
47250400	**Low discrimination**						**47250400**
					GRA	SPO	
Fusobacterium mortiferum		71.15	0.43	ARA 1	1	1	
Clostridium innocuum		19.31	0.21		99	99	
Clostridium barati		4.44	0.08		99	99	
47250500	**Low discrimination**						**47250500**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.50	ARA 1	1	1	
Clostridium barati		7.03	0.16		99	99	
Lactobacillus catenaformis		3.26	0.15		99	1	
47251000	**Low discrimination**						**47251000**
					GRA		
Lactobacillus catenaformis		56.56	0.81	MLT 90	99		
Fusobacterium mortiferum		14.79	0.81		1		
Lactobacillus jensenii		13.45	0.70		99		
47251001	**Low discrimination**						**47251001**
					GRA		
Lactobacillus catenaformis		56.41	0.47	MLT 90	99		
Fusobacterium mortiferum		14.75	0.46	URE 1	1		
Lactobacillus jensenii		13.42	0.36		99		
47251002	**Low discrimination**						**47251002**
					SPO		
Lactobacillus catenaformis		53.34	0.81	MLT 90	1		
Clostridium barati		33.08	0.72		99		
Lactobacillus jensenii		8.46	0.67		1		
47251003	**Low discrimination**						**47251003**
					SPO		
Lactobacillus catenaformis		53.23	0.47	MLT 90	1		
Clostridium barati		33.01	0.38	URE 1	99		
Lactobacillus jensenii		8.44	0.33		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
47251010	**Low discrimination**						**47251010**
					GRA		
Lactobacillus jensenii		82.28	0.62	MLT 95	99		
Lactobacillus catenaformis		10.48	0.47		99		
Fusobacterium mortiferum		2.74	0.46		1		
47251012	**Low discrimination**						**47251012**
					SPO		
Lactobacillus jensenii		73.53	0.59	MLT 95	1		
Lactobacillus catenaformis		14.05	0.47		1		
Clostridium barati		8.72	0.38		99		
47251020	**Low discrimination**						**47251020**
					SPO		
Clostridium barati		65.98	0.72		99		
Lactobacillus catenaformis		26.60	0.70		1		
Lactobacillus jensenii		6.33	0.60		1		
47251021	**Low discrimination**						**47251021**
					SPO		
Clostridium barati		65.76	0.38	URE 1	99		
Lactobacillus catenaformis		26.51	0.36		1		
Lactobacillus jensenii		6.30	0.26		1		
47251022	**Low discrimination**						**47251022**
					SPO		
Clostridium barati		85.77	0.80		99		
Lactobacillus catenaformis		11.53	0.70		1		
Lactobacillus jensenii		1.83	0.57		1		
47251023	**Low discrimination**						**47251023**
					SPO		
Clostridium barati		85.66	0.46	URE 1	99		
Lactobacillus catenaformis		11.51	0.36		1		
Lactobacillus jensenii		1.83	0.23		1		
47251030	**Low discrimination**						**47251030**
					SPO		
Lactobacillus jensenii		69.02	0.52	MLT 95	1		
Clostridium barati		21.82	0.38		99		
Lactobacillus catenaformis		8.80	0.36		1		
47251032	**Low discrimination**						**47251032**
					SPO		
Clostridium barati		54.08	0.46	MLZ 1	99		
Lactobacillus jensenii		38.02	0.49		1		
Lactobacillus catenaformis		7.27	0.36		1		
47251040	**Low discrimination**						**47251040**
					GRA	SPO	
Clostridium barati		53.28	0.46	SOR 8	99	99	
Lactobacillus catenaformis		29.94	0.47		99	1	
Fusobacterium mortiferum		7.83	0.46		1	1	
47251042	**Low discrimination**						**47251042**
					SPO		
Clostridium barati		79.07	0.54	SOR 8	99		
Lactobacillus catenaformis		14.81	0.47		1		
Lactobacillus acidophilus		2.58	0.31		1		
47251060	**Acceptable identification**						**47251060**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		93.78	0.54	SOR 8	99		
Lactobacillus catenaformis		4.39	0.36		1		
Lactobacillus jensenii		1.04	0.26		1		
47251062	**Very good identification**						**47251062**
					SPO		
Clostridium barati		97.06	0.62	SOR 8	99		
Lactobacillus catenaformis		1.52	0.36		1		
Lactobacillus acidophilus		1.06	0.31		1		
47251100	**Low discrimination**						**47251100**
					SPO		
Lactobacillus catenaformis		40.17	0.84	MLT 90	1		
Collinsella aerofaciens		26.58	0.66		1		
Clostridium barati		16.61	0.72		99		
47251101	**Low discrimination**						**47251101**
					SPO		
Lactobacillus catenaformis		40.05	0.50	MLT 90	1		
Collinsella aerofaciens		26.49	0.32	URE 1	1		
Clostridium barati		16.56	0.38		99		
47251102	**Low discrimination**						**47251102**
					SPO		
Clostridium barati		50.15	0.80		99		
Lactobacillus catenaformis		40.43	0.84		1		
Lactobacillus acidophilus		4.41	0.64		1		
47251103	**Low discrimination**						**47251103**
					SPO		
Clostridium barati		50.02	0.46	URE 1	99		
Lactobacillus catenaformis		40.33	0.50		1		
Lactobacillus acidophilus		4.40	0.30		1		
47251110	**Low discrimination**						**47251110**
					SPO		
Lactobacillus catenaformis		39.14	0.50	MLT 90	1		
Collinsella aerofaciens		25.89	0.32	MLZ 1	1		
Clostridium barati		16.18	0.38		99		
47251112	**Low discrimination**						**47251112**
					GRA	SPO	
Clostridium barati		39.72	0.46	MLZ 1	99	99	
Lactobacillus catenaformis		32.02	0.50		99	1	
Leptotrichia buccalis		16.05	0.32		1	1	
47251120	**Low discrimination**						**47251120**
					SPO		
Clostridium barati		63.72	0.80		99		
Clostridium paraputrificum		19.03	0.66		99		
Lactobacillus catenaformis		12.84	0.73		1		
47251121	**Low discrimination**						**47251121**
					SPO		
Clostridium barati		63.49	0.46	URE 1	99		
Clostridium paraputrificum		18.96	0.32		99		
Lactobacillus catenaformis		12.80	0.39		1		
47251122	**Low discrimination**						**47251122**
					SPO		
Clostridium barati		87.26	0.88		99		
Lactobacillus catenaformis		5.86	0.73		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		3.72	0.60		99		
47251123	**Low discrimination**						**47251123**
					SPO		
Clostridium barati		87.09	0.54	URE 1	99		
Lactobacillus catenaformis		5.85	0.39		1		
Clostridium paraputrificum		3.72	0.25		99		
47251130	**Low discrimination**						**47251130**
					SPO		
Clostridium barati		62.69	0.46	MLZ 1	99		
Clostridium paraputrificum		18.72	0.32		99		
Lactobacillus catenaformis		12.64	0.39		1		
47251132	**Low discrimination**						**47251132**
					SPO		
Clostridium barati		82.28	0.54	MLZ 1	99		
Lactobacillus catenaformis		5.53	0.39		1		
Bifidobacterium breve		4.11	0.44		1		
47251140	**Low discrimination**						**47251140**
					SPO		
Clostridium barati		62.65	0.54	SOR 8	99		
Lactobacillus catenaformis		17.60	0.50		1		
Collinsella aerofaciens		11.65	0.32		1		
47251142	**Low discrimination**						**47251142**
					SPO		
Clostridium barati		80.60	0.62	SOR 8	99		
Lactobacillus acidophilus		10.08	0.49		1		
Lactobacillus catenaformis		7.55	0.50		1		
47251160	**Low discrimination**						**47251160**
					SPO		
Clostridium barati		77.35	0.62	SOR 8	99		
Clostridium paraputrificum		19.99	0.47		99		
Lactobacillus catenaformis		1.81	0.39		1		
47251162	**Good identification**						**47251162**
					SPO		
Clostridium barati		91.50	0.70	SOR 8	99		
Lactobacillus acidophilus		3.81	0.49		1		
Clostridium paraputrificum		3.38	0.40		99		
47251163	**Acceptable identification**						**47251163**
					SPO		
Clostridium barati		91.11	0.36	SOR 8	99		
Lactobacillus acidophilus		3.80	0.15	URE 1	1		
Clostridium paraputrificum		3.36	0.06		99		
47251172	**Low discrimination**						**47251172**
					SPO		
Clostridium barati		60.23	0.36	MLZ 1	99		
Bifidobacterium breve		34.57	0.44	SOR 8	1		
Lactobacillus acidophilus		2.51	0.15		1		
47251200	**Low discrimination**						**47251200**
					GRA	SPO	
Clostridium barati		53.28	0.46	RHA 8	99	99	
Lactobacillus catenaformis		29.94	0.47		99	1	
Fusobacterium mortiferum		7.83	0.46		1	1	
47251202	**Low discrimination**						**47251202**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium barati		79.84	0.54	RHA 8	99		
Lactobacillus catenaformis		14.95	0.47		1		
Lactobacillus jensenii		2.37	0.33		1		
47251220	**Acceptable identification**						**47251220**
					SPO		
Clostridium barati		94.33	0.54	RHA 8	99		
Lactobacillus catenaformis		4.42	0.36		1		
Lactobacillus jensenii		1.05	0.26		1		
47251222	**Very good identification**						**47251222**
					SPO		
Clostridium barati		97.24	0.62	RHA 8	99		
Lactobacillus catenaformis		1.52	0.36		1		
Lactobacillus acidophilus		0.64	0.27		1		
47251242	**Good identification**						**47251242**
					SPO		
Clostridium barati		95.24	0.36	RHA 8	99		
Lactobacillus acidophilus		2.69	0.11	SOR 8	1		
Lactobacillus catenaformis		2.07	0.12		1		
47251260	**Good identification**						**47251260**
					SPO		
Clostridium barati		99.46	0.36	RHA 8	99		
Lactobacillus catenaformis		0.54	0.02	SOR 8	1		
47251262	**Good identification**						**47251262**
					SPO		
Clostridium barati		98.89	0.44	RHA 8	99		
Lactobacillus acidophilus		0.93	0.11	SOR 8	1		
Lactobacillus catenaformis		0.18	0.02		1		
47251300	**Low discrimination**						**47251300**
					SPO		
Clostridium barati		63.08	0.54	RHA 8	99		
Lactobacillus catenaformis		17.72	0.50		1		
Collinsella aerofaciens		11.72	0.32		1		
47251302	**Low discrimination**						**47251302**
					SPO		
Clostridium barati		84.46	0.62	RHA 8	99		
Lactobacillus catenaformis		7.91	0.50		1		
Lactobacillus acidophilus		6.43	0.45		1		
47251320	**Good identification**						**47251320**
					SPO		
Clostridium barati		93.38	0.62	RHA 8	99		
Clostridium paraputrificum		3.24	0.32		99		
Lactobacillus catenaformis		2.19	0.39		1		
47251322	**Good identification**						**47251322**
					GRA	SPO	
Clostridium barati		94.44	0.70	RHA 8	99	99	
Lactobacillus acidophilus		2.40	0.45		99	1	
Prevotella oralis		1.74	0.41		1	1	
47251323	**Acceptable identification**						**47251323**
					GRA	SPO	
Clostridium barati		94.65	0.36	RHA 8	99	99	
Lactobacillus acidophilus		2.40	0.11	URE 1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		1.74	0.07		1	1	
47251332	**Acceptable identification**						**47251332**
					GRA	SPO	
Clostridium barati		92.60	0.36	RHA 8	99	99	
Lactobacillus acidophilus		2.35	0.11	MLZ 1	99	1	
Prevotella oralis		1.70	0.07		1	1	
47251340	**Acceptable identification**						**47251340**
					SPO		
Clostridium barati		93.68	0.36	RHA 8	99		
Lactobacillus catenaformis		3.06	0.15	SOR 8	1		
Collinsella aerofaciens		2.02	-.03		1		
47251342	**Low discrimination**						**47251342**
					SPO		
Clostridium barati		89.36	0.44	RHA 8	99		
Lactobacillus acidophilus		9.67	0.30	SOR 8	1		
Lactobacillus catenaformis		0.97	0.15		1		
47251360	**Good identification**						**47251360**
					SPO		
Clostridium barati		96.83	0.44	RHA 8	99		
Clostridium paraputrificum		2.91	0.13	SOR 8	99		
Lactobacillus catenaformis		0.26	0.05		1		
47251362	**Good identification**						**47251362**
					SPO		
Clostridium barati		95.78	0.52	RHA 8	99		
Lactobacillus acidophilus		3.46	0.30	SOR 8	1		
Clostridium paraputrificum		0.41	0.06		99		
47251400	**Low discrimination**						**47251400**
					GRA	SPO	
Lactobacillus catenaformis		38.24	0.47	MLT 90	99	1	
Clostridium innocuum		34.37	0.43	ARA 1	99	99	
Fusobacterium mortiferum		10.00	0.46		1	1	
47251402	**Low discrimination**						**47251402**
					SPO		
Lactobacillus catenaformis		36.75	0.47	MLT 90	1		
Clostridium innocuum		33.03	0.43	ARA 1	99		
Clostridium barati		22.79	0.38		99		
47251420	**Low discrimination**						**47251420**
					SPO		
Clostridium barati		65.49	0.38	ARA 1	99		
Lactobacillus catenaformis		26.40	0.36		1		
Lactobacillus jensenii		6.28	0.26		1		
47251422	**Low discrimination**						**47251422**
					SPO		
Clostridium barati		85.51	0.46	ARA 1	99		
Lactobacillus catenaformis		11.49	0.36		1		
Lactobacillus jensenii		1.82	0.23		1		
47251500	**Low discrimination**						**47251500**
					SPO		
Lactobacillus catenaformis		40.06	0.50	MLT 90	1		
Collinsella aerofaciens		26.50	0.32	ARA 1	1		
Clostridium barati		16.56	0.38		99		
47251502	**Low discrimination**						**47251502**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium barati		49.22	0.46	ARA 1	99		
Lactobacillus catenaformis		39.68	0.50		1		
Lactobacillus acidophilus		4.33	0.30		1		
47251520	**Low discrimination**						**47251520**
					SPO		
Clostridium barati		63.82	0.46	ARA 1	99		
Clostridium paraputrificum		19.06	0.32		99		
Lactobacillus catenaformis		12.86	0.39		1		
47251522	**Low discrimination**						**47251522**
					SPO		
Clostridium barati		87.25	0.54	ARA 1	99		
Lactobacillus catenaformis		5.86	0.39		1		
Clostridium paraputrificum		3.72	0.25		99		
47251562	**Acceptable identification**						**47251562**
					SPO		
Clostridium barati		91.56	0.36	ARA 1	99		
Lactobacillus acidophilus		3.82	0.15	SOR 8	1		
Clostridium paraputrificum		3.38	0.06		99		
47251722	**Acceptable identification**						**47251722**
					GRA	SPO	
Clostridium barati		92.37	0.36	RHA 8	99	99	
Lactobacillus acidophilus		2.35	0.11	ARA 1	99	1	
Parabacteroides distasonis		1.85	0.21		1	1	
47252000	**Very good identification**						**47252000**
					GRA	SPO	
Clostridium innocuum		98.61	0.68	GAL 95	99	99	
Fusobacterium mortiferum		0.65	0.43		1	1	
Clostridium barati		0.35	0.24		99	99	
47252002	**Very good identification**						**47252002**
					GRA	SPO	
Clostridium innocuum		98.84	0.68	GAL 95	99	99	
Clostridium barati		1.05	0.32		99	99	
Fusobacterium mortiferum		0.07	0.27		1	1	
47252010	**Low discrimination**						**47252010**
Clostridium innocuum		71.34	0.45	GAL 95			
Clostridium difficile		28.66	0.35	MLZ 4			
47252012	**Good identification**						**47252012**
Clostridium innocuum		99.34	0.45	GAL 95			
Clostridium difficile		0.40	0.00	MLZ 4			
Clostridium barati		0.26	-.02				
47252022	**Low discrimination**						**47252022**
					SPO		
Clostridium barati		75.70	0.40	GAL 95	99		
Clostridium innocuum		24.07	0.34	MAN 8	99		
Lactobacillus catenaformis		0.23	0.02		1		
47252100	**Low discrimination**						**47252100**
					GRA	SPO	
Fusobacterium mortiferum		41.46	0.50	MAN 1	1	1	
Clostridium barati		28.65	0.32		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		27.33	0.34		99	99	
47252102	**Low discrimination**						**47252102**
					GRA	SPO	
Clostridium barati		71.28	0.40	GAL 95	99	99	
Clostridium innocuum		22.67	0.34	MAN 8	99	99	
Fusobacterium mortiferum		3.82	0.33		1	1	
47252120	**Good identification**						**47252120**
					GRA	SPO	
Clostridium barati		97.80	0.40	GAL 95	99	99	
Clostridium paraputrificum		0.65	-.03	MAN 8	99	99	
Fusobacterium mortiferum		0.48	0.16		1	1	
47252122	**Good identification**						**47252122**
					SPO		
Clostridium barati		98.75	0.48	GAL 95	99		
Lactobacillus acidophilus		0.34	0.08	MAN 8	1		
Clostridium tertium		0.32	0.09		99		
47252400	**Good identification**						**47252400**
Clostridium innocuum		100.0	0.55	GAL 95			
47252402	**Good identification**						**47252402**
Clostridium innocuum		99.94	0.55	GAL 95			
Clostridium barati		0.06	-.02				
47253000	**Very good identification**						**47253000**
					SPO		
Clostridium innocuum		99.35	0.90		99		
Clostridium barati		0.35	0.46		99		
Lactobacillus catenaformis		0.20	0.47		1		
47253001	**Good identification**						**47253001**
					SPO		
Clostridium innocuum		99.16	0.56	URE 1	99		
Clostridium barati		0.35	0.11		99		
Actinomyces naeslundii		0.20	0.34		1		
47253002	**Very good identification**						**47253002**
					SPO		
Clostridium innocuum		98.71	0.90		99		
Clostridium barati		1.05	0.54		99		
Lactobacillus catenaformis		0.20	0.47		1		
47253003	**Good identification**						**47253003**
					SPO		
Clostridium innocuum		98.57	0.56	URE 1	99		
Clostridium barati		1.04	0.20		99		
Actinomyces naeslundii		0.20	0.34		1		
47253010	**Very good identification**						**47253010**
					SPO		
Clostridium innocuum		99.46	0.67	MLZ 4	99		
Lactobacillus jensenii		0.38	0.28		1		
Clostridium barati		0.09	0.11		99		
47253012	**Very good identification**						**47253012**
					SPO		
Clostridium innocuum		99.42	0.67	MLZ 4	99		
Clostridium barati		0.26	0.20		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		0.25	0.25		1	
47253020	**Low discrimination**					**47253020**
					SPO	
Clostridium barati		49.62	0.54	MAN 8	99	
Clostridium innocuum		47.34	0.56		99	
Lactobacillus catenaformis		2.32	0.36		1	
47253022	**Low discrimination**					**47253022**
					SPO	
Clostridium barati		74.51	0.62	MAN 8	99	
Clostridium innocuum		23.69	0.56		99	
Lactobacillus catenaformis		1.16	0.36		1	
47253040	**Good identification**					**47253040**
					SPO	
Clostridium innocuum		96.77	0.56	SOR 1	99	
Clostridium barati		2.94	0.27		99	
Lactobacillus catenaformis		0.19	0.12		1	
47253042	**Acceptable identification**					**47253042**
					SPO	
Clostridium innocuum		91.33	0.56	SOR 1	99	
Clostridium barati		8.33	0.36		99	
Lactobacillus catenaformis		0.18	0.12		1	
47253060	**Low discrimination**					**47253060**
					SPO	
Clostridium barati		89.59	0.36	MAN 8	99	
Clostridium innocuum		9.93	0.22	SOR 8	99	
Lactobacillus catenaformis		0.49	0.02		1	
47253062	**Good identification**					**47253062**
					SPO	
Clostridium barati		95.48	0.44	MAN 8	99	
Clostridium innocuum		3.53	0.22	SOR 8	99	
Lactobacillus acidophilus		0.63	0.09		1	
47253100	**Low discrimination**					**47253100**
					SPO	
Clostridium barati		39.01	0.54	MAN 8	99	
Clostridium innocuum		37.21	0.56		99	
Lactobacillus catenaformis		10.96	0.50		1	
47253102	**Low discrimination**					**47253102**
					SPO	
Clostridium barati		66.42	0.62	MAN 8	99	
Clostridium innocuum		21.12	0.56		99	
Lactobacillus catenaformis		6.22	0.50		1	
47253112	**Low discrimination**					**47253112**
Clostridium tertium		45.40	0.39	MLT 99		
Clostridium innocuum		20.19	0.32	TRE 90		
Clostridium barati		15.39	0.28			
47253120	**Good identification**					**47253120**
					SPO	
Clostridium barati		91.33	0.62	MAN 8	99	
Clostridium paraputrificum		3.17	0.32		99	
Lactobacillus catenaformis		2.14	0.39		1	
47253122	**Good identification**					**47253122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium barati		94.10	0.70	MAN 8	99	
Lactobacillus acidophilus		1.67	0.42		1	
Clostridium tertium		1.58	0.43		99	
47253123	**Low discrimination**					**47253123**
					SPO	
Clostridium barati		89.29	0.36	MAN 8	99	
Actinomyces naeslundii		5.31	0.42	URE 1	1	
Lactobacillus acidophilus		1.59	0.08		1	
47253130	**Low discrimination**					**47253130**
					SPO	
Clostridium tertium		70.38	0.39	MLT 99	99	
Clostridium barati		23.86	0.28	TRE 90	99	
Bifidobacterium breve		4.06	0.27		1	
47253132	**Low discrimination**					**47253132**
					SPO	
Clostridium tertium		64.60	0.47	MLT 99	99	
Clostridium barati		21.90	0.36	TRE 90	99	
Bifidobacterium breve		12.57	0.44		1	
47253140	**Low discrimination**					**47253140**
					SPO	
Clostridium barati		81.44	0.36	MAN 8	99	
Clostridium innocuum		9.03	0.22	SOR 8	99	
Bifidobacterium breve		2.72	0.23		1	
47253142	**Low discrimination**					**47253142**
					SPO	
Clostridium barati		80.33	0.44	MAN 8	99	
Bifidobacterium breve		9.04	0.40	SOR 8	1	
Lactobacillus acidophilus		6.08	0.27		1	
47253152	**Low discrimination**					**47253152**
					SPO	
Bifidobacterium breve		88.16	0.40	MLT 99	1	
Clostridium barati		7.92	0.10	RAF 95	99	
Clostridium tertium		2.71	0.05		99	
47253160	**Acceptable identification**					**47253160**
					SPO	
Clostridium barati		94.62	0.44	MAN 8	99	
Clostridium paraputrificum		2.84	0.13	SOR 8	99	
Bifidobacterium breve		1.87	0.27		1	
47253162	**Acceptable identification**					**47253162**
					SPO	
Clostridium barati		90.85	0.52	MAN 8	99	
Bifidobacterium breve		6.06	0.44	SOR 8	1	
Lactobacillus acidophilus		2.29	0.27		1	
47253172	**Low discrimination**					**47253172**
					SPO	
Bifidobacterium breve		83.10	0.44	MLT 99	1	
Clostridium barati		12.59	0.18	RAF 95	99	
Clostridium tertium		4.32	0.13		99	
47253200	**Good identification**					**47253200**
					SPO	
Clostridium innocuum		96.77	0.56	RHA 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		2.94	0.27		99	
Lactobacillus catenaformis		0.19	0.12		1	
47253202	**Acceptable identification**					**47253202**
					SPO	
Clostridium innocuum		91.39	0.56	RHA 1	99	
Clostridium barati		8.33	0.36		99	
Lactobacillus catenaformis		0.18	0.12		1	
47253220	**Low discrimination**					**47253220**
					SPO	
Clostridium barati		89.59	0.36	MAN 8	99	
Clostridium innocuum		9.93	0.22	RHA 8	99	
Lactobacillus catenaformis		0.49	0.02		1	
47253222	**Good identification**					**47253222**
					SPO	
Clostridium barati		95.58	0.44	MAN 8	99	
Clostridium innocuum		3.53	0.22	RHA 8	99	
Lactobacillus acidophilus		0.38	0.05		1	
47253300	**Low discrimination**					**47253300**
					SPO	
Clostridium barati		84.87	0.36	MAN 8	99	
Clostridium innocuum		9.40	0.22	RHA 8	99	
Lactobacillus catenaformis		2.77	0.15		1	
47253302	**Acceptable identification**					**47253302**
					SPO	
Clostridium barati		90.97	0.44	MAN 8	99	
Lactobacillus acidophilus		4.19	0.23	RHA 8	1	
Clostridium innocuum		3.36	0.22		99	
47253320	**Acceptable identification**					**47253320**
Clostridium barati		91.38	0.44	MAN 8		
Clostridium ramosum		7.83	0.27	RHA 8		
Clostridium paraputrificum		0.37	-.03			
47253322	**Low discrimination**					**47253322**
					SPO	
Clostridium barati		88.04	0.52	MAN 8	99	
Clostridium ramosum		10.06	0.37	RHA 8	92	
Lactobacillus acidophilus		1.35	0.23		1	
47253400	**Very good identification**					**47253400**
					SPO	
Clostridium innocuum		99.96	0.77		99	
Clostridium barati		0.02	0.11		99	
Lactobacillus catenaformis		0.01	0.12		1	
47253401	**Good identification**					**47253401**
Clostridium innocuum		100.0	0.43	URE 1		
47253402	**Very good identification**					**47253402**
					SPO	
Clostridium innocuum		99.92	0.77		99	
Clostridium barati		0.06	0.20		99	
Lactobacillus catenaformis		0.01	0.12		1	
47253403	**Good identification**					**47253403**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.43	URE 1			
47253410	**Good identification**						**47253410**
Clostridium innocuum		100.0	0.54	MLZ 4			
47253412	**Good identification**						**47253412**
Clostridium innocuum		100.0	0.54	MLZ 4			
47253420	**Acceptable identification**						**47253420**
					SPO		
Clostridium innocuum		94.09	0.43	NAG 1	99		
Clostridium barati		5.65	0.20		99		
Lactobacillus catenaformis		0.26	0.02		1		
47253422	**Low discrimination**						**47253422**
					SPO		
Clostridium innocuum		84.54	0.43	NAG 1	99		
Clostridium barati		15.22	0.28		99		
Lactobacillus catenaformis		0.24	0.02		1		
47253440	**Good identification**						**47253440**
Clostridium innocuum		100.0	0.43	SOR 1			
47253442	**Good identification**						**47253442**
Clostridium innocuum		99.48	0.43	SOR 1			
Clostridium barati		0.52	0.01				
47253500	**Acceptable identification**						**47253500**
					SPO		
Clostridium innocuum		90.54	0.43	LAC 1	99		
Clostridium barati		5.43	0.20		99		
Lactobacillus catenaformis		1.53	0.15		1		
47253502	**Low discrimination**						**47253502**
					SPO		
Clostridium innocuum		78.00	0.43	LAC 1	99		
Clostridium barati		14.04	0.28		99		
Bifidobacterium adolescentis		4.54	0.31		1		
47253522	**Acceptable identification**						**47253522**
					SPO		
Clostridium barati		92.66	0.36	MAN 8	99		
Clostridium innocuum		1.73	0.09	ARA 1	99		
Lactobacillus acidophilus		1.64	0.08		1		
47253600	**Good identification**						**47253600**
Clostridium innocuum		100.0	0.43	RHA 1			
47253602	**Good identification**						**47253602**
Clostridium innocuum		99.48	0.43	RHA 1			
Clostridium barati		0.52	0.01				
47254000	**Very good identification**						**47254000**
					GRA	SPO	
Fusobacterium mortiferum		97.20	0.58	XYL 7	1	1	
Clostridium innocuum		1.06	0.12		99	99	
Clostridium barati		0.81	0.08		99	99	
47254002	**Low discrimination**						**47254002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Fusobacterium mortiferum		71.74	0.42	XYL 7	1	1	
Clostridium barati		16.23	0.16	BGL 10	99	99	
Clostridium innocuum		7.01	0.12		99	99	
47254100	**Very good identification**						**47254100**
					GRA	SPO	
Fusobacterium mortiferum		98.13	0.65	XYL 7	1	1	
Clostridium barati		1.06	0.16		99	99	
Lactobacillus catenaformis		0.49	0.15		99	1	
47254102	**Low discrimination**						**47254102**
					GRA	SPO	
Fusobacterium mortiferum		74.86	0.48	XYL 7	1	1	
Clostridium barati		21.77	0.24	BGL 10	99	99	
Lactobacillus catenaformis		3.37	0.15		99	1	
47255000	**Low discrimination**						**47255000**
					GRA	SPO	
Fusobacterium mortiferum		53.00	0.61	XYL 7	1	1	
Lactobacillus catenaformis		27.20	0.47		99	1	
Clostridium innocuum		7.29	0.34		99	99	
47255002	**Low discrimination**						**47255002**
					SPO		
Lactobacillus catenaformis		43.75	0.47	MLT 90	1		
Clostridium barati		27.14	0.38	XYL 1	99		
Clostridium innocuum		11.73	0.34		99		
47255020	**Low discrimination**						**47255020**
					SPO		
Clostridium barati		64.76	0.38	XYL 1	99		
Lactobacillus catenaformis		26.11	0.36		1		
Lactobacillus jensenii		6.21	0.26		1		
47255022	**Low discrimination**						**47255022**
					SPO		
Clostridium barati		85.04	0.46	XYL 1	99		
Lactobacillus catenaformis		11.43	0.36		1		
Lactobacillus jensenii		1.81	0.23		1		
47255100	**Low discrimination**						**47255100**
					GRA		
Fusobacterium mortiferum		57.74	0.68	XYL 7	1		
Lactobacillus catenaformis		19.05	0.50		99		
Collinsella aerofaciens		12.60	0.32		99		
47255102	**Low discrimination**						**47255102**
					GRA	SPO	
Clostridium barati		43.24	0.46	XYL 1	99	99	
Lactobacillus catenaformis		34.86	0.50		99	1	
Fusobacterium mortiferum		11.74	0.51		1	1	
47255120	**Low discrimination**						**47255120**
					SPO		
Clostridium barati		61.98	0.46	XYL 1	99		
Clostridium paraputrificum		18.51	0.32		99		
Lactobacillus catenaformis		12.49	0.39		1		
47255122	**Low discrimination**						**47255122**
					GRA	SPO	
Clostridium barati		82.87	0.54	XYL 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus catenaformis		5.57	0.39		99	1	
Prevotella oralis		3.78	0.32		1	1	
47255162	**Acceptable identification**						**47255162**
					SPO		
Clostridium barati		91.12	0.36	XYL 1	99		
Lactobacillus acidophilus		3.80	0.15	SOR 8	1		
Clostridium paraputrificum		3.37	0.06		99		
47255322	**Low discrimination**						**47255322**
					GRA	SPO	
Clostridium barati		61.81	0.36	XYL 1	99	99	
Parabacteroides distasonis		18.19	0.41	RHA 8	1	1	
Prevotella oralis		13.90	0.26		1	1	
47255332	**Low discrimination**						**47255332**
					GRA	SPO	
Parabacteroides distasonis		88.50	0.36	RAF 99	1	1	
Clostridium clostridioforme		5.86	0.10		50	66	
Clostridium barati		5.64	0.02		99	99	
47255502	**Low discrimination**						**47255502**
					SPO		
Bifidobacterium adolescentis		77.39	0.46	MLT 99	1		
Clostridium butyricum		10.35	0.17	RAF 92	99		
Actinomyces israelii		4.77	0.32		1		
47255512	**Good identification**						**47255512**
Bifidobacterium adolescentis		96.70	0.48	MLT 99			
Clostridium clostridioforme		2.02	0.16	RAF 92			
Actinomyces israelii		0.73	0.18				
47256000	**Good identification**						**47256000**
					GRA	SPO	
Clostridium innocuum		98.73	0.46	GAL 95	99	99	
Fusobacterium mortiferum		0.93	0.24	XYL 5	1	1	
Clostridium difficile		0.34	0.01		99	99	
47256002	**Good identification**						**47256002**
Clostridium innocuum		99.80	0.46	GAL 95			
Clostridium barati		0.20	-.02	XYL 5			
47257000	**Very good identification**						**47257000**
					GRA	SPO	
Clostridium innocuum		99.70	0.68	XYL 5	99	99	
Actinomyces israelii		0.11	0.33		99	1	
Fusobacterium mortiferum		0.07	0.27		1	1	
47257002	**Very good identification**						**47257002**
					SPO		
Clostridium innocuum		99.57	0.68	XYL 5	99		
Clostridium barati		0.20	0.20		99		
Actinomyces israelii		0.17	0.36		1		
47257010	**Good identification**						**47257010**
					SPO		
Clostridium innocuum		99.39	0.45	XYL 5	99		
Actinomyces israelii		0.41	0.19	MLZ 4	1		
Clostridium tertium		0.20	-.01		99		
47257012	**Good identification**						**47257012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium innocuum		98.80	0.45	XYL 5	99	
Actinomyces israelii		0.61	0.22	MLZ 4	1	
Clostridium tertium		0.60	0.07		99	
47257102	**Low discrimination**					**47257102**
					SPO	
Clostridium tertium		34.84	0.37	MLT 99	99	
Clostridium innocuum		25.19	0.34	TRE 90	99	
Actinomyces israelii		17.13	0.46		1	
47257112	**Low discrimination**					**47257112**
					SPO	
Clostridium tertium		86.57	0.41	MLT 99	99	
Bifidobacterium adolescentis		8.39	0.32	TRE 90	1	
Actinomyces israelii		3.58	0.32		1	
47257120	**Low discrimination**					**47257120**
					SPO	
Clostridium tertium		67.12	0.37	MLT 99	99	
Clostridium barati		29.29	0.28	TRE 90	99	
Actinomyces israelii		1.40	0.23		1	
47257122	**Low discrimination**					**47257122**
					SPO	
Clostridium tertium		67.99	0.45	MLT 99	99	
Clostridium barati		29.67	0.36	TRE 90	99	
Actinomyces israelii		0.71	0.26		1	
47257130	**Good identification**					**47257130**
Clostridium tertium		100.0	0.41	MLT 99		
47257132	**Good identification**					**47257132**
					SPO	
Clostridium tertium		99.61	0.50	MLT 99	99	
Clostridium barati		0.25	0.02	TRE 90	99	
Bifidobacterium breve		0.14	0.10		1	
47257140	**Low discrimination**					**47257140**
					SPO	
Actinomyces israelii		86.47	0.36	MLT 99	1	
Clostridium barati		8.58	0.01		99	
Clostridium innocuum		4.95	0.00		99	
47257142	**Low discrimination**					**47257142**
					SPO	
Actinomyces israelii		72.43	0.39	MLT 99	1	
Clostridium barati		14.38	0.10		99	
Bifidobacterium adolescentis		6.60	0.15		1	
47257400	**Good identification**					**47257400**
					SPO	
Clostridium innocuum		99.47	0.55	XYL 5	99	
Actinomyces israelii		0.53	0.31		1	
47257402	**Good identification**					**47257402**
					SPO	
Clostridium innocuum		99.14	0.55	XYL 5	99	
Actinomyces israelii		0.79	0.34		1	
Bifidobacterium adolescentis		0.08	0.11		1	
47257500	**Low discrimination**					**47257500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Actinomyces israelii		59.68	0.42	MLT 99	1			
Clostridium innocuum		28.40	0.21		99			
Clostridium butyricum		9.80	0.07		99			
47257502	**Low discrimination**							**47257502**
					SPO			
Bifidobacterium adolescentis		62.87	0.45	MLT 99	1			
Actinomyces israelii		26.81	0.45	RAF 92	1			
Clostridium innocuum		8.50	0.21		99			
47257512	**Low discrimination**							**47257512**
Bifidobacterium adolescentis		89.68	0.46	MLT 99				
Actinomyces israelii		4.68	0.31	RAF 92				
Bifidobacterium dentium		4.25	0.06					
47257542	**Low discrimination**							**47257542**
Actinomyces israelii		57.30	0.38	MLT 99				
Bifidobacterium adolescentis		42.70	0.29					
47260100	**Good identification**							**47260100**
					GRA	SPO		
Fusobacterium mortiferum		97.72	0.41	FRU 99	1	1		
Clostridium barati		2.28	-.02		99	99		
47261000	**Low discrimination**							**47261000**
					GRA	SPO		
Lactobacillus jensenii		74.99	0.44	MLT 95	99	1		
Clostridium innocuum		9.90	0.26	FRU 99	99	99		
Fusobacterium mortiferum		8.21	0.38		1	1		
47261002	**Low discrimination**							**47261002**
					SPO			
Lactobacillus jensenii		68.26	0.41	MLT 95	1			
Clostridium innocuum		13.52	0.26	FRU 99	99			
Clostridium barati		7.74	0.20		99			
47261010	**Good identification**							**47261010**
					SPO			
Lactobacillus jensenii		98.38	0.36	MLT 95	1			
Clostridium innocuum		1.62	0.03	FRU 99	99			
47261100	**Low discrimination**							**47261100**
					COC	GRA	SPO	
Fusobacterium mortiferum		50.08	0.44	FRU 99	1	1	1	
Atopobium parvulum		24.69	0.27		99	99	1	
Clostridium barati		14.82	0.20		1	99	99	
47261102	**Low discrimination**							**47261102**
					GRA	SPO		
Lactobacillus acidophilus		70.85	0.39	MLT 94	99	1		
Clostridium barati		20.92	0.28	FRU 99	99	99		
Fusobacterium mortiferum		2.62	0.27		1	1		
47261122	**Low discrimination**							**47261122**
					SPO			
Lactobacillus acidophilus		49.22	0.39	MLT 94	1			
Clostridium barati		43.60	0.36	FRU 99	99			
Clostridium paraputrificum		5.69	0.16		99			
47262000	**Good identification**							**47262000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.39	FRU 95			
47262002	**Good identification**						**47262002**
Clostridium innocuum		100.0	0.39	FRU 95			
47263000	**Very good identification**						**47263000**
					SPO		
Clostridium innocuum		99.92	0.61	FRU 95	99		
Lactobacillus jensenii		0.08	0.10		1		
47263002	**Very good identification**						**47263002**
					SPO		
Clostridium innocuum		99.87	0.61	FRU 95	99		
Lactobacillus jensenii		0.05	0.07		1		
Clostridium barati		0.05	0.01		99		
47263010	**Good identification**						**47263010**
					SPO		
Clostridium innocuum		99.39	0.37	FRU 95	99		
Lactobacillus jensenii		0.62	0.02	MLZ 4	1		
47263012	**Good identification**						**47263012**
Clostridium innocuum		100.0	0.37	FRU 95			
47263400	**Good identification**						**47263400**
Clostridium innocuum		100.0	0.48	FRU 95			
47263402	**Good identification**						**47263402**
Clostridium innocuum		100.0	0.48	FRU 95			
47267000	**Good identification**						**47267000**
Clostridium innocuum		100.0	0.39	FRU 95			
47267002	**Good identification**						**47267002**
Clostridium innocuum		100.0	0.39	FRU 95			
47270000	**Low discrimination**						**47270000**
					GRA	SPO	
Fusobacterium mortiferum		84.98	0.69		1	1	
Lactobacillus jensenii		11.76	0.44		99	1	
Clostridium innocuum		1.55	0.26		99	99	
47270002	**Low discrimination**						**47270002**
					GRA	SPO	
Fusobacterium mortiferum		38.67	0.52	BGL 10	1	1	
Lactobacillus jensenii		32.11	0.41		99	1	
Clostridium barati		18.97	0.32		99	99	
47270010	**Low discrimination**						**47270010**
					GRA	SPO	
Lactobacillus jensenii		79.93	0.36	MLT 95	99	1	
Fusobacterium mortiferum		17.50	0.34	GAL 99	1	1	
Clostridium innocuum		1.32	0.03		99	99	
47270022	**Low discrimination**						**47270022**
					SPO		
Clostridium barati		82.82	0.40	TRE 8	99		
Lactobacillus jensenii		11.68	0.31	GAL 95	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Lactobacillus acidophilus		4.68	0.21		1			
47270100	**Very good identification**							**47270100**
					COC	GRA	SPO	
Fusobacterium mortiferum		96.71	0.75		1	1	1	
Clostridium barati		2.26	0.32		1	99	99	
Atopobium parvulum		0.72	0.27		99	99	1	
47270101	**Good identification**							**47270101**
					GRA	SPO		
Fusobacterium mortiferum		97.72	0.41	URE 1	1	1		
Clostridium barati		2.28	-.02		99	99		
47270102	**Low discrimination**							**47270102**
					GRA	SPO		
Fusobacterium mortiferum		48.23	0.59	BGL 10	1	1		
Clostridium barati		30.42	0.40		99	99		
Lactobacillus acidophilus		19.78	0.39		99	1		
47270110	**Acceptable identification**							**47270110**
					GRA	SPO		
Fusobacterium mortiferum		90.45	0.41	MLZ 1	1	1		
Leptotrichia buccalis		5.65	-.03		1	1		
Clostridium barati		2.11	-.02		99	99		
47270120	**Low discrimination**							**47270120**
					GRA	SPO		
Clostridium barati		72.43	0.40	TRE 8	99	99		
Clostridium paraputrificum		12.69	0.22	GAL 95	99	99		
Fusobacterium mortiferum		10.44	0.41		1	1		
47270122	**Low discrimination**							**47270122**
					SPO			
Clostridium barati		79.77	0.48	TRE 8	99			
Lactobacillus acidophilus		17.28	0.39	GAL 95	1			
Clostridium paraputrificum		2.00	0.16		99			
47270140	**Low discrimination**							**47270140**
					GRA	SPO		
Fusobacterium mortiferum		83.26	0.41	SOR 1	1	1		
Clostridium barati		16.75	0.14		99	99		
47270300	**Low discrimination**							**47270300**
					GRA	SPO		
Fusobacterium mortiferum		83.26	0.41	RHA 1	1	1		
Clostridium barati		16.75	0.14		99	99		
47270500	**Good identification**							**47270500**
					GRA	SPO		
Fusobacterium mortiferum		97.72	0.41	ARA 1	1	1		
Clostridium barati		2.28	-.02		99	99		
47271000	**Low discrimination**							**47271000**
					GRA	SPO		
Lactobacillus jensenii		85.01	0.78	MLT 95	99	1		
Fusobacterium mortiferum		9.31	0.72		1	1		
Clostridium innocuum		2.15	0.48		99	99		
47271001	**Low discrimination**							**47271001**
					GRA	SPO		
Lactobacillus jensenii		84.15	0.44	MLT 95	99	1		
Fusobacterium mortiferum		9.21	0.38	URE 1	1	1		
Clostridium innocuum		2.13	0.14		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47271002	**Low discrimination**						**47271002**
					SPO		
Lactobacillus jensenii		77.41	0.75	MLT 95	1		
Clostridium barati		8.78	0.54		99		
Lactobacillus acidophilus		7.76	0.55		1		
47271003	**Low discrimination**						**47271003**
					SPO		
Lactobacillus jensenii		76.20	0.41	MLT 95	1		
Clostridium barati		8.64	0.20	URE 1	99		
Lactobacillus acidophilus		7.63	0.21		1		
47271010	**Very good identification**						**47271010**
					GRA	SPO	
Lactobacillus jensenii		99.21	0.70	MLT 95	99	1	
Fusobacterium mortiferum		0.33	0.38		1	1	
Clostridium innocuum		0.31	0.25		99	99	
47271011	**Good identification**						**47271011**
Lactobacillus jensenii		100.0	0.36	MLT 95			
47271012	**Good identification**						**47271012**
					GRA	SPO	
Lactobacillus jensenii		94.31	0.67	MLT 95	99	1	
Leptotrichia buccalis		4.52	0.32		1	1	
Clostridium innocuum		0.45	0.25		99	99	
47271020	**Low discrimination**						**47271020**
					SPO		
Lactobacillus jensenii		74.85	0.68	MLT 95	1		
Clostridium barati		22.63	0.54		99		
Lactobacillus catenaformis		1.06	0.36		1		
47271022	**Low discrimination**						**47271022**
					SPO		
Clostridium barati		48.72	0.62	TRE 8	99		
Lactobacillus jensenii		35.81	0.65		1		
Lactobacillus acidophilus		14.35	0.55		1		
47271030	**Very good identification**						**47271030**
					SPO		
Lactobacillus jensenii		99.05	0.60	MLT 95	1		
Clostridium barati		0.91	0.20		99		
Lactobacillus catenaformis		0.04	0.02		1		
47271032	**Good identification**						**47271032**
					GRA	SPO	
Lactobacillus jensenii		93.00	0.57	MLT 95	99	1	
Clostridium barati		3.83	0.28		99	99	
Leptotrichia buccalis		1.98	0.15		1	1	
47271040	**Low discrimination**						**47271040**
					GRA	SPO	
Lactobacillus jensenii		72.77	0.44	MLT 95	99	1	
Clostridium barati		15.79	0.27	SOR 1	99	99	
Fusobacterium mortiferum		7.97	0.38		1	1	
47271042	**Low discrimination**						**47271042**
					SPO		
Lactobacillus acidophilus		37.38	0.40	MLT 94	1		
Lactobacillus jensenii		30.49	0.41	LAC 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium barati		29.76	0.36	SOR 11	99			
47271050	**Good identification**							**47271050**
Lactobacillus jensenii		100.0	0.36	MLT 95				
47271060	**Low discrimination**							**47271060**
					SPO			
Clostridium barati		70.11	0.36	TRE 8	99			
Lactobacillus jensenii		26.94	0.34	SOR 8	1			
Clostridium paraputrificum		1.68	0.03		99			
47271062	**Low discrimination**							**47271062**
					SPO			
Clostridium barati		66.26	0.44	TRE 8	99			
Lactobacillus acidophilus		27.74	0.40	SOR 8	1			
Lactobacillus jensenii		5.66	0.31		1			
47271100	**Low discrimination**							**47271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		49.73	0.78		1	1	1	
Atopobium parvulum		24.52	0.61		99	99	1	
Clostridium barati		14.72	0.54		1	99	99	
47271101	**Low discrimination**							**47271101**
					COC	GRA	SPO	
Fusobacterium mortiferum		46.40	0.44	URE 1	1	1	1	
Atopobium parvulum		22.87	0.27		99	99	1	
Clostridium barati		13.73	0.20		1	99	99	
47271102	**Low discrimination**							**47271102**
					GRA	SPO		
Lactobacillus acidophilus		70.51	0.73	MLT 94	99	1		
Clostridium barati		20.82	0.62		99	99		
Fusobacterium mortiferum		2.61	0.62		1	1		
47271103	**Low discrimination**							**47271103**
					SPO			
Lactobacillus acidophilus		68.30	0.39	MLT 94	1			
Clostridium barati		20.17	0.28	URE 1	99			
Actinomyces naeslundii		3.60	0.42		1			
47271112	**Very good identification**							**47271112**
					GRA			
Leptotrichia buccalis		98.58	0.66	MLT 99	1			
Lactobacillus acidophilus		0.73	0.39		99			
Lactobacillus jensenii		0.21	0.33		99			
47271120	**Low discrimination**							**47271120**
Clostridium barati		44.09	0.62	TRE 8				
Clostridium paraputrificum		40.25	0.56					
Clostridium septicum		6.52	0.45					
47271122	**Low discrimination**							**47271122**
					SPO			
Lactobacillus acidophilus		48.90	0.73	MLT 94	1			
Clostridium barati		43.31	0.70		99			
Clostridium paraputrificum		5.65	0.50		99			
47271123	**Low discrimination**							**47271123**
					SPO			
Lactobacillus acidophilus		47.98	0.39	MLT 94	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium barati		42.50	0.36	URE 1	99			
Clostridium paraputrificum		5.54	0.16		99			
47271132	**Low discrimination**							**47271132**
					GRA	SPO		
Leptotrichia buccalis		84.81	0.50	MLT 99	1	1		
Lactobacillus acidophilus		5.62	0.39	NAG 10	99	1		
Clostridium barati		4.98	0.36		99	99		
47271140	**Low discrimination**							**47271140**
					COC	GRA	SPO	
Clostridium barati		53.73	0.36	TRE 8	1	99	99	
Fusobacterium mortiferum		21.09	0.44	SOR 8	1	1	1	
Atopobium parvulum		10.40	0.27		99	99	1	
47271142	**Low discrimination**							**47271142**
					SPO			
Lactobacillus acidophilus		81.00	0.58	MLT 94	1			
Clostridium barati		16.83	0.44	SOR 11	99			
Bifidobacterium breve		1.26	0.37		1			
47271160	**Low discrimination**							**47271160**
					SPO			
Clostridium barati		52.83	0.44	TRE 8	99			
Clostridium paraputrificum		41.74	0.37	SOR 8	99			
Lactobacillus acidophilus		2.57	0.24		1			
47271162	**Low discrimination**							**47271162**
					SPO			
Lactobacillus acidophilus		57.98	0.58	MLT 94	1			
Clostridium barati		36.13	0.52	SOR 11	99			
Clostridium paraputrificum		4.08	0.31		99			
47271172	**Low discrimination**							**47271172**
					GRA			
Bifidobacterium breve		48.37	0.41	MLT 99	99			
Leptotrichia buccalis		21.76	0.15	RAF 95	1			
Lactobacillus acidophilus		17.64	0.24		99			
47271200	**Low discrimination**							**47271200**
					GRA	SPO		
Lactobacillus jensenii		72.95	0.44	MLT 95	99	1		
Clostridium barati		15.82	0.27	RHA 1	99	99		
Fusobacterium mortiferum		7.99	0.38		1	1		
47271202	**Low discrimination**							**47271202**
					SPO			
Lactobacillus jensenii		35.71	0.41	MLT 95	1			
Clostridium barati		34.86	0.36	RHA 1	99			
Lactobacillus acidophilus		26.66	0.36		1			
47271210	**Good identification**							**47271210**
Lactobacillus jensenii		100.0	0.36	MLT 95				
47271220	**Low discrimination**							**47271220**
					SPO			
Clostridium barati		71.20	0.36	TRE 8	99			
Lactobacillus jensenii		27.35	0.34	RHA 8	1			
Lactobacillus acidophilus		0.55	0.02		1			
47271222	**Low discrimination**							**47271222**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium barati		73.84	0.44	TRE 8	99			
Lactobacillus acidophilus		18.83	0.36	RHA 8	1			
Lactobacillus jensenii		6.30	0.31		1			
47271300	**Low discrimination**							**47271300**
					COC	GRA	SPO	
Clostridium barati		57.08	0.36	TRE 8	1	99	99	
Fusobacterium mortiferum		22.41	0.44	RHA 8	1	1	1	
Atopobium parvulum		11.05	0.27		99	99	1	
47271302	**Low discrimination**							**47271302**
					GRA	SPO		
Lactobacillus acidophilus		73.49	0.54	MLT 94	99	1		
Clostridium barati		25.07	0.44	RHA 7	99	99		
Fusobacterium mortiferum		0.36	0.27		1	1		
47271320	**Low discrimination**							**47271320**
Clostridium barati		71.13	0.44	TRE 8				
Clostridium ramosum		16.84	0.35	RHA 8				
Clostridium paraputrificum		7.54	0.22					
47271322	**Low discrimination**							**47271322**
					SPO			
Clostridium barati		42.73	0.52	TRE 8	99			
Lactobacillus acidophilus		41.76	0.54	RHA 8	1			
Clostridium ramosum		13.49	0.45		92			
47271342	**Low discrimination**							**47271342**
					SPO			
Lactobacillus acidophilus		80.65	0.39	MLT 94	1			
Clostridium barati		19.35	0.26	RHA 7	99			
47271362	**Low discrimination**							**47271362**
					SPO			
Lactobacillus acidophilus		56.96	0.39	MLT 94	1			
Clostridium barati		41.00	0.34	RHA 7	99			
Clostridium ramosum		1.50	0.11	SOR 11	92			
47271400	**Low discrimination**							**47271400**
					GRA	SPO		
Lactobacillus jensenii		62.67	0.44	MLT 95	99	1		
Clostridium innocuum		27.74	0.35	ARA 1	99	99		
Fusobacterium mortiferum		6.86	0.38		1	1		
47271402	**Low discrimination**							**47271402**
					SPO			
Lactobacillus jensenii		52.04	0.41	MLT 95	1			
Clostridium innocuum		34.55	0.35	ARA 1	99			
Clostridium barati		5.90	0.20		99			
47271410	**Acceptable identification**							**47271410**
					SPO			
Lactobacillus jensenii		94.76	0.36	MLT 95	1			
Clostridium innocuum		5.24	0.12	ARA 1	99			
47271500	**Low discrimination**							**47271500**
					COC	GRA	SPO	
Fusobacterium mortiferum		41.04	0.44	ARA 1	1	1	1	
Atopobium parvulum		20.23	0.27		99	99	1	
Clostridium butyricum		16.03	0.18		1	99	99	
47271502	**Low discrimination**							**47271502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Lactobacillus acidophilus		64.06	0.39	MLT 94	1		
Clostridium barati		18.91	0.28	ARA 1	99		
Bifidobacterium adolescentis		4.98	0.29		1		
47271522	**Low discrimination**						**47271522**
					SPO		
Lactobacillus acidophilus		48.47	0.39	MLT 94	1		
Clostridium barati		42.94	0.36	ARA 1	99		
Clostridium paraputrificum		5.60	0.16		99		
47272000	**Very good identification**						**47272000**
					GRA	SPO	
Clostridium innocuum		99.21	0.61	GAL 95	99	99	
Fusobacterium mortiferum		0.55	0.34		1	1	
Clostridium barati		0.09	0.06		99	99	
47272002	**Very good identification**						**47272002**
					GRA	SPO	
Clostridium innocuum		99.60	0.61	GAL 95	99	99	
Clostridium barati		0.26	0.14		99	99	
Fusobacterium mortiferum		0.06	0.18		1	1	
47272010	**Acceptable identification**						**47272010**
					SPO		
Clostridium innocuum		91.55	0.37	GAL 95	99		
Clostridium difficile		7.88	0.15	MLZ 4	99		
Lactobacillus jensenii		0.57	0.02		1		
47272012	**Good identification**						**47272012**
Clostridium innocuum		100.0	0.37	GAL 95			
47272100	**Low discrimination**						**47272100**
					GRA	SPO	
Fusobacterium mortiferum		48.94	0.41	MAN 1	1	1	
Clostridium innocuum		37.94	0.26		99	99	
Clostridium barati		9.84	0.14		99	99	
47272400	**Good identification**						**47272400**
Clostridium innocuum		100.0	0.48	GAL 95			
47272402	**Good identification**						**47272402**
Clostridium innocuum		100.0	0.48	GAL 95			
47273000	**Very good identification**						**47273000**
					SPO		
Clostridium innocuum		99.44	0.82		99		
Lactobacillus jensenii		0.40	0.44		1		
Clostridium barati		0.09	0.27		99		
47273001	**Good identification**						**47273001**
					SPO		
Clostridium innocuum		98.52	0.48	URE 1	99		
Actinomyces naeslundii		1.08	0.39		1		
Lactobacillus jensenii		0.40	0.10		1		
47273002	**Very good identification**						**47273002**
					SPO		
Clostridium innocuum		99.29	0.82		99		
Lactobacillus jensenii		0.27	0.41		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		0.26	0.36		99	
47273003	**Good identification**					**47273003**
					SPO	
Clostridium innocuum		98.26	0.48	URE 1	99	
Actinomyces naeslundii		1.08	0.39		1	
Lactobacillus jensenii		0.26	0.07		1	
47273010	**Very good identification**					**47273010**
					SPO	
Clostridium innocuum		96.70	0.59	MLZ 4	99	
Lactobacillus jensenii		3.12	0.36		1	
Clostridium tertium		0.19	0.13		99	
47273012	**Very good identification**					**47273012**
					SPO	
Clostridium innocuum		97.12	0.59	MLZ 4	99	
Lactobacillus jensenii		2.09	0.33		1	
Clostridium tertium		0.57	0.21		99	
47273020	**Low discrimination**					**47273020**
					SPO	
Clostridium innocuum		71.90	0.48	NAG 1	99	
Clostridium barati		18.65	0.36		99	
Lactobacillus jensenii		7.17	0.34		1	
47273021	**Low discrimination**					**47273021**
					SPO	
Actinomyces naeslundii		46.37	0.39	ESL 12	1	
Clostridium innocuum		42.59	0.14	MLT 94	99	
Clostridium barati		11.05	0.01	RAF 92	99	
47273022	**Low discrimination**					**47273022**
					SPO	
Clostridium innocuum		48.43	0.48	NAG 1	99	
Clostridium barati		37.69	0.44		99	
Lactobacillus acidophilus		6.72	0.33		1	
47273023	**Low discrimination**					**47273023**
					SPO	
Actinomyces naeslundii		36.22	0.39	ESL 12	1	
Clostridium innocuum		33.27	0.14	MLT 94	99	
Clostridium barati		25.89	0.10	RAF 92	99	
47273040	**Good identification**					**47273040**
					SPO	
Clostridium innocuum		98.52	0.48	SOR 1	99	
Clostridium barati		0.74	0.09		99	
Lactobacillus jensenii		0.40	0.10		1	
47273042	**Good identification**					**47273042**
					SPO	
Clostridium innocuum		95.52	0.48	SOR 1	99	
Clostridium barati		2.16	0.17		99	
Lactobacillus acidophilus		1.64	0.18		1	
47273100	**Low discrimination**					**47273100**
Clostridium innocuum		51.21	0.48	LAC 1		
Clostridium tertium		23.05	0.43			
Clostridium barati		13.29	0.36			
47273101	**Low discrimination**					**47273101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		64.45	0.47	ESL 12	1		
Clostridium innocuum		20.80	0.14	MLT 94	99		
Clostridium tertium		9.36	0.09	RAF 92	99		
47273102	**Low discrimination**						**47273102**
					SPO		
Lactobacillus acidophilus		32.52	0.51	MLT 94	1		
Clostridium tertium		27.52	0.51	MAN 5	99		
Clostridium innocuum		20.38	0.48		99		
47273103	**Low discrimination**						**47273103**
					SPO		
Actinomyces naeslundii		39.61	0.47	ESL 12	1		
Lactobacillus acidophilus		20.40	0.17	MLT 94	1		
Clostridium tertium		17.26	0.17	RAF 92	99		
47273110	**Acceptable identification**						**47273110**
					SPO		
Clostridium tertium		93.14	0.47	MLT 99	99		
Clostridium innocuum		4.85	0.25		99		
Bifidobacterium breve		0.67	0.20		1		
47273112	**Low discrimination**						**47273112**
					GRA	SPO	
Clostridium tertium		81.29	0.55	MLT 99	99	99	
Leptotrichia buccalis		14.23	0.32		1	1	
Bifidobacterium breve		1.98	0.37		99	1	
47273120	**Low discrimination**						**47273120**
Clostridium tertium		43.81	0.51	MLT 99			
Clostridium ramosum		25.56	0.45				
Clostridium barati		25.25	0.44				
47273121	**Low discrimination**						**47273121**
					SPO		
Actinomyces naeslundii		51.52	0.47	ESL 12	1		
Clostridium tertium		22.45	0.17	MLT 94	99		
Clostridium ramosum		13.10	0.11	RAF 92	92		
47273122	**Low discrimination**						**47273122**
Clostridium tertium		35.82	0.59	MLT 99			
Clostridium ramosum		27.87	0.56				
Clostridium barati		20.65	0.52				
47273130	**Good identification**						**47273130**
					SPO		
Clostridium tertium		98.93	0.55	MLT 99	99		
Bifidobacterium breve		0.42	0.24		1		
Clostridium ramosum		0.33	0.11		92		
47273132	**Very good identification**						**47273132**
					GRA	SPO	
Clostridium tertium		97.01	0.63	MLT 99	99	99	
Bifidobacterium breve		1.40	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
47273142	**Low discrimination**						**47273142**
					SPO		
Lactobacillus acidophilus		54.23	0.36	MLT 94	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		18.61	0.26	MAN 5	99		
Bifidobacterium breve		16.05	0.37	SOR 11	1		
47273152	**Low discrimination**						**47273152**
					GRA	SPO	
Bifidobacterium breve		64.29	0.37	MLT 99	99	1	
Clostridium tertium		26.71	0.21	RAF 95	99	99	
Leptotrichia buccalis		4.67	-.03		1	1	
47273172	**Low discrimination**						**47273172**
					SPO		
Bifidobacterium breve		57.47	0.41	MLT 99	1		
Clostridium tertium		40.29	0.29	RAF 95	99		
Clostridium barati		1.14	0.00		99		
47273200	**Good identification**						**47273200**
					SPO		
Clostridium innocuum		98.86	0.48	RHA 1	99		
Clostridium barati		0.74	0.09		99		
Lactobacillus jensenii		0.40	0.10		1		
47273202	**Good identification**						**47273202**
					SPO		
Clostridium innocuum		96.55	0.48	RHA 1	99		
Clostridium barati		2.18	0.17		99		
Lactobacillus acidophilus		1.01	0.14		1		
47273320	**Acceptable identification**						**47273320**
Clostridium ramosum		92.68	0.47	MLT 99			
Clostridium barati		6.01	0.26				
Clostridium tertium		1.21	0.17				
47273322	**Good identification**						**47273322**
					SPO		
Clostridium ramosum		91.98	0.58	MLT 99	92		
Clostridium barati		4.47	0.34		99		
Lactobacillus acidophilus		2.64	0.32		1		
47273400	**Very good identification**						**47273400**
					SPO		
Clostridium innocuum		99.95	0.70		99		
Actinomyces israelii		0.03	0.23		1		
Lactobacillus jensenii		0.02	0.10		1		
47273401	**Good identification**						**47273401**
Clostridium innocuum		100.0	0.35	URE 1			
47273402	**Very good identification**						**47273402**
					SPO		
Clostridium innocuum		99.92	0.70		99		
Actinomyces israelii		0.04	0.26		1		
Lactobacillus jensenii		0.02	0.07		1		
47273403	**Good identification**						**47273403**
Clostridium innocuum		100.0	0.35	URE 1			
47273410	**Good identification**						**47273410**
					SPO		
Clostridium innocuum		99.82	0.46	MLZ 4	99		
Lactobacillus jensenii		0.18	0.02		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47273412	**Good identification**							**47273412**
Clostridium innocuum		100.0	0.46	MLZ 4				
47273420	**Good identification**							**47273420**
Clostridium innocuum		98.54	0.35	NAG 1				
Clostridium barati		1.46	0.01					
47273422	**Good identification**							**47273422**
					SPO			
Clostridium innocuum		95.01	0.35	NAG 1	99			
Clostridium barati		4.23	0.10		99			
Lactobacillus acidophilus		0.75	-.01		1			
47273440	**Good identification**							**47273440**
					SPO			
Clostridium innocuum		99.01	0.35	SOR 1	99			
Actinomyces israelii		0.99	0.16		1			
47273442	**Good identification**							**47273442**
					SPO			
Clostridium innocuum		98.52	0.35	SOR 1	99			
Actinomyces israelii		1.48	0.19		1			
47273500	**Low discrimination**							**47273500**
					SPO			
Clostridium innocuum		85.57	0.35	LAC 1	99			
Actinomyces israelii		8.81	0.33		1			
Clostridium tertium		2.20	0.09		99			
47273502	**Low discrimination**							**47273502**
					SPO			
Clostridium innocuum		67.22	0.35	LAC 1	99			
Actinomyces israelii		10.39	0.36		1			
Bifidobacterium adolescentis		7.42	0.28		1			
47273600	**Good identification**							**47273600**
Clostridium innocuum		100.0	0.35	RHA 1				
47273602	**Good identification**							**47273602**
Clostridium innocuum		100.0	0.35	RHA 1				
47274000	**Good identification**							**47274000**
					GRA	SPO		
Fusobacterium mortiferum		96.96	0.49	XYL 7	1	1		
Lactobacillus jensenii		1.80	0.10		99	1		
Clostridium innocuum		1.24	0.04		99	99		
47274100	**Low discrimination**							**47274100**
					COC	GRA	SPO	
Fusobacterium mortiferum		84.01	0.56	XYL 7	1	1	1	
Atopobium parvulum		15.48	0.32		99	99	1	
Clostridium barati		0.26	-.02		1	99	99	
47274102	**Low discrimination**							**47274102**
					COC	GRA	SPO	
Fusobacterium mortiferum		69.81	0.39	XYL 7	1	1	1	
Atopobium parvulum		20.44	0.19	BGL 10	99	99	1	
Clostridium barati		5.91	0.06		1	99	99	
47274712	**Good identification**							**47274712**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium clostridioforme		100.0	0.37	MLT 99				
47274732	**Good identification**							**47274732**
Clostridium clostridioforme		98.90	0.37	MLT 99				
Parabacteroides distasonis		1.10	0.09	RAF 94				
47275000	**Low discrimination**							**47275000**
					COC	GRA		
Lactobacillus jensenii		44.17	0.44	MLT 95	1	99		
Fusobacterium mortiferum		36.03	0.52	XYL 1	1	1		
Atopobium parvulum		10.33	0.32		99	99		
47275002	**Low discrimination**							**47275002**
					GRA	SPO		
Lactobacillus jensenii		58.72	0.41	MLT 95	99	1		
Clostridium innocuum		11.63	0.26	XYL 1	99	99		
Fusobacterium mortiferum		7.98	0.36		1	1		
47275010	**Acceptable identification**							**47275010**
					GRA	SPO		
Lactobacillus jensenii		94.89	0.36	MLT 95	99	1		
Fusobacterium mortiferum		2.35	0.18	XYL 1	1	1		
Clostridium innocuum		1.57	0.03		99	99		
47275100	**Good identification**							**47275100**
					COC	GRA	SPO	
Atopobium parvulum		90.19	0.66	MLT 99	99	99	1	
Fusobacterium mortiferum		7.41	0.59		1	1	1	
Clostridium butyricum		1.40	0.28		1	99	99	
47275102	**Low discrimination**							**47275102**
					COC	SPO		
Atopobium parvulum		71.06	0.53	MLT 99	99	1		
Lactobacillus acidophilus		13.36	0.39		1	1		
Clostridium barati		3.94	0.28		1	99		
47275120	**Low discrimination**							**47275120**
					COC	SPO		
Atopobium parvulum		87.35	0.53	MLT 99	99	1		
Clostridium barati		4.85	0.28		1	99		
Clostridium paraputrificum		4.43	0.22		1	99		
47275122	**Low discrimination**							**47275122**
					COC	SPO		
Lactobacillus acidophilus		27.74	0.39	MLT 94	1	1		
Atopobium parvulum		26.04	0.40	XYL 1	99	1		
Clostridium barati		24.57	0.36		1	99		
47275322	**Low discrimination**							**47275322**
					GRA	SPO		
Parabacteroides distasonis		84.75	0.52	RAF 99	1	1		
Clostridium barati		5.50	0.18		99	99		
Lactobacillus acidophilus		5.37	0.20		99	1		
47275332	**Acceptable identification**							**47275332**
Parabacteroides distasonis		90.37	0.47	RAF 99				
Clostridium clostridioforme		9.64	0.25					
47275500	**Acceptable identification**							**47275500**
					COC	SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		90.22	0.44	MLT 99	1	99	
Atopobium parvulum		6.54	0.32	RAF 95	99	1	
Actinomyces israelii		2.59	0.39		1	1	
47275502	**Low discrimination**						**47275502**
					SPO		
Clostridium butyricum		67.23	0.37	MLT 99	99		
Bifidobacterium adolescentis		21.40	0.43	RAF 95	1		
Actinomyces israelii		7.44	0.42		1		
47275512	**Low discrimination**						**47275512**
					SPO		
Bifidobacterium adolescentis		67.25	0.45	MLT 99	1		
Clostridium clostridioforme		15.41	0.31	RAF 92	66		
Clostridium butyricum		9.10	0.15		99		
47275522	**Low discrimination**						**47275522**
					GRA	SPO	
Parabacteroides distasonis		52.52	0.40	RAF 99	1	1	
Clostridium butyricum		30.75	0.19		99	99	
Clostridium clostridioforme		6.92	0.19		50	66	
47275612	**Good identification**						**47275612**
Clostridium clostridioforme		100.0	0.36	MLT 99			
47275632	**Good identification**						**47275632**
Clostridium clostridioforme		98.77	0.36	MLT 99			
Parabacteroides distasonis		1.23	0.09	RAF 94			
47275712	**Good identification**						**47275712**
					GRA		
Clostridium clostridioforme		99.04	0.45	MLT 99	50		
Bifidobacterium adolescentis		0.60	0.11	RAF 94	99		
Parabacteroides distasonis		0.37	0.09		1		
47275722	**Low discrimination**						**47275722**
Parabacteroides distasonis		72.74	0.48	RAF 99			
Clostridium clostridioforme		23.43	0.34				
Bacteroides ovatus		2.09	0.06				
47275732	**Low discrimination**						**47275732**
Clostridium clostridioforme		72.47	0.45	MLT 99			
Parabacteroides distasonis		26.59	0.44	RAF 94			
Bacteroides ovatus		0.94	0.03				
47276000	**Good identification**						**47276000**
					GRA	SPO	
Clostridium innocuum		99.21	0.39	GAL 95	99	99	
Fusobacterium mortiferum		0.79	0.15	XYL 5	1	1	
47276002	**Good identification**						**47276002**
Clostridium innocuum		100.0	0.39	GAL 95			
47277000	**Very good identification**						**47277000**
					SPO		
Clostridium innocuum		98.54	0.61	XYL 5	99		
Actinomyces israelii		1.21	0.43		1		
Clostridium tertium		0.12	0.11		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47277002	**Very good identification**						**47277002**
					SPO		
Clostridium innocuum		97.72	0.61	XYL 5	99		
Actinomyces israelii		1.80	0.46		1		
Clostridium tertium		0.35	0.19		99		
47277010	**Acceptable identification**						**47277010**
					SPO		
Clostridium innocuum		90.78	0.37	XYL 5	99		
Clostridium tertium		4.67	0.15	MLZ 4	99		
Actinomyces israelii		3.99	0.29		1		
47277012	**Low discrimination**						**47277012**
					SPO		
Clostridium innocuum		81.96	0.37	XYL 5	99		
Clostridium tertium		12.64	0.24	MLZ 4	99		
Actinomyces israelii		5.40	0.32		1		
47277100	**Low discrimination**						**47277100**
					COC	SPO	
Clostridium tertium		59.25	0.45	MLT 99	1	99	
Actinomyces israelii		24.35	0.53		1	1	
Atopobium parvulum		8.89	0.32		99	1	
47277102	**Low discrimination**						**47277102**
					SPO		
Clostridium tertium		78.79	0.53	MLT 99	99		
Actinomyces israelii		16.19	0.56		1		
Clostridium innocuum		2.22	0.26		99		
47277110	**Good identification**						**47277110**
					SPO		
Clostridium tertium		96.48	0.50	MLT 99	99		
Actinomyces israelii		3.33	0.39		1		
Clostridium innocuum		0.19	0.03		99		
47277112	**Very good identification**						**47277112**
					SPO		
Clostridium tertium		97.11	0.58	MLT 99	99		
Actinomyces israelii		1.68	0.42		1		
Bifidobacterium adolescentis		0.70	0.29		1		
47277120	**Good identification**						**47277120**
					COC	SPO	
Clostridium tertium		97.39	0.53	MLT 99	1	99	
Atopobium parvulum		0.86	0.19		99	1	
Actinomyces israelii		0.85	0.33		1	1	
47277122	**Very good identification**						**47277122**
					SPO		
Clostridium tertium		98.24	0.62	MLT 99	99		
Clostridium ramosum		0.56	0.21		92		
Actinomyces israelii		0.43	0.36		1		
47277130	**Very good identification**						**47277130**
					SPO		
Clostridium tertium		99.87	0.58	MLT 99	99		
Actinomyces israelii		0.07	0.18		1		
Bifidobacterium dentium		0.06	-.07		1		
47277132	**Very good identification**						**47277132**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		99.76	0.66	MLT 99	99	
Bifidobacterium dentium		0.20	0.10		1	
Actinomyces israelii		0.04	0.21		1	
47277140	**Acceptable identification**					**47277140**
					SPO	
Actinomyces israelii		92.84	0.46	MLT 99	1	
Clostridium tertium		5.87	0.11		99	
Clostridium butyricum		1.29	-.07		99	
47277142	**Low discrimination**					**47277142**
					SPO	
Actinomyces israelii		86.51	0.49	MLT 99	1	
Clostridium tertium		10.93	0.19		99	
Bifidobacterium adolescentis		1.40	0.12		1	
47277302	**Low discrimination**					**47277302**
					SPO	
Actinomyces israelii		59.00	0.37	MLT 99	1	
Clostridium tertium		38.52	0.19	RHA 7	99	
Lactobacillus acidophilus		2.48	-.02		1	
47277400	**Acceptable identification**					**47277400**
					SPO	
Clostridium innocuum		94.05	0.48	XYL 5	99	
Actinomyces israelii		5.35	0.41		1	
Clostridium butyricum		0.61	0.04		99	
47277402	**Acceptable identification**					**47277402**
					SPO	
Clostridium innocuum		91.81	0.48	XYL 5	99	
Actinomyces israelii		7.83	0.44		1	
Clostridium butyricum		0.23	-.03		99	
47277442	**Low discrimination**					**47277442**
					SPO	
Actinomyces israelii		76.65	0.37	MLT 99	1	
Clostridium innocuum		23.35	0.13		99	
47277500	**Low discrimination**					**47277500**
					SPO	
Actinomyces israelii		56.67	0.52	MLT 99	1	
Clostridium butyricum		38.75	0.28		99	
Clostridium innocuum		2.52	0.13		99	
47277502	**Low discrimination**					**47277502**
					SPO	
Actinomyces israelii		58.18	0.55	MLT 99	1	
Bifidobacterium adolescentis		24.18	0.42		1	
Clostridium butyricum		10.32	0.21		99	
47277512	**Low discrimination**					**47277512**
Bifidobacterium dentium		82.22	0.40	MLT 99		
Bifidobacterium adolescentis		11.68	0.43	RAF 95		
Actinomyces israelii		3.44	0.40			
47277532	**Acceptable identification**					**47277532**
					SPO	
Bifidobacterium dentium		95.00	0.44	MLT 99	1	
Clostridium tertium		4.79	0.32	RAF 95	99	
Actinomyces israelii		0.14	0.20		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47277540	**Low discrimination**						**47277540**
					COC	SPO	
Actinomyces israelii		49.43	0.45	MLT 99	1	1	
Blautia producta		43.01	0.24		99	1	
Clostridium butyricum		7.56	0.10		1	99	
47277542	**Low discrimination**						**47277542**
					COC		
Actinomyces israelii		80.90	0.48	MLT 99	1		
Bifidobacterium adolescentis		10.68	0.26		1		
Blautia producta		5.21	0.07		99		
47277702	**Low discrimination**						**47277702**
					SPO		
Actinomyces israelii		88.60	0.35	MLT 99	1		
Clostridium butyricum		6.46	-.05	RHA 7	99		
Bifidobacterium adolescentis		4.94	0.08		1		
47300100	**Acceptable identification**						**47300100**
					GRA	SPO	
Fusobacterium mortiferum		92.17	0.42	MLT 11	1	1	
Collinsella aerofaciens		6.13	-.03	FRU 99	99	1	
Clostridium bifermentans		1.71	-.01		99	99	
47301101	**Very good identification**						**47301101**
					SPO		
Clostridium chauvoei		99.55	0.57	SAL 1	99		
Collinsella aerofaciens		0.21	-.03		1		
Actinomyces naeslundii		0.19	0.33		1		
47301102	**Low discrimination**						**47301102**
Bifidobacterium longum subsp. infantis		70.99	0.40	RAF 95			
Lactobacillus catenaformis		8.68	0.32	FRU 99			
Collinsella aerofaciens		7.02	0.15				
47301103	**Good identification**						**47301103**
					SPO		
Clostridium chauvoei		97.61	0.44	SAL 1	99		
Bifidobacterium longum subsp. infantis		1.31	0.06		1		
Actinomyces naeslundii		1.08	0.33		1		
47301121	**Good identification**						**47301121**
					SPO		
Clostridium chauvoei		98.56	0.48	SAL 1	99		
Actinomyces naeslundii		0.58	0.33		1		
Clostridium paraputrificum		0.35	-.03		99		
47301122	**Low discrimination**						**47301122**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		57.04	0.44	RAF 95	99	1	
Prevotella oralis		16.72	0.38	FRU 99	1	1	
Clostridium barati		7.92	0.32		99	99	
47301123	**Low discrimination**						**47301123**
					SPO		
Clostridium chauvoei		86.72	0.36	SAL 1	99		
Bifidobacterium longum subsp. infantis		6.22	0.10		1		
Actinomyces naeslundii		2.88	0.33		1		
47303100	**Low discrimination**						**47303100**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		44.43	0.35	ESL 12	1		
Collinsella aerofaciens		37.21	-.03	RAF 92	1		
Clostridium barati		9.88	-.03	FRU 95	99		
47303102	**Low discrimination**						**47303102**
					SPO		
Actinomyces naeslundii		31.83	0.35	ESL 12	1		
Bifidobacterium longum subsp. infantis		29.96	0.06	RAF 92	1		
Clostridium barati		21.22	0.05	FRU 95	99		
47303103	**Low discrimination**						**47303103**
					SPO		
Actinomyces naeslundii		67.99	0.38	ESL 12	1		
Clostridium chauvoei		32.01	0.10	RAF 92	99		
47303120	**Low discrimination**						**47303120**
					SPO		
Actinomyces naeslundii		32.76	0.35	ESL 12	1		
Clostridium barati		21.84	0.05	RAF 92	99		
Clostridium paraputrificum		15.37	-.03	FRU 95	99		
47303121	**Low discrimination**						**47303121**
					SPO		
Actinomyces naeslundii		52.93	0.38	ESL 12	1		
Clostridium chauvoei		47.07	0.14	RAF 92	99		
47303123	**Low discrimination**						**47303123**
					SPO		
Actinomyces naeslundii		86.43	0.38	ESL 12	1		
Clostridium chauvoei		13.57	0.01	RAF 92	99		
47305102	**Acceptable identification**						**47305102**
Bifidobacterium longum subsp. infantis		92.69	0.38	RAF 95			
Bifidobacterium adolescentis		3.05	0.27	FRU 99			
Eubacterium contortum		2.34	0.16				
47305122	**Acceptable identification**						**47305122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		94.13	0.43	RAF 95	1	99	
Prevotella oralis		4.17	0.23	FRU 99	1	1	
Bacteroides fragilis		0.47	-.03		99	1	
47305300	**Very good identification**						**47305300**
Eubacterium contortum		100.0	0.63	FRU 99			
47305302	**Good identification**						**47305302**
Eubacterium contortum		99.60	0.50	FRU 99			
Bifidobacterium longum subsp. infantis		0.40	0.04				
47305320	**Good identification**						**47305320**
					GRA		
Eubacterium contortum		98.72	0.50	FRU 99	99		
Bacteroides vulgatus		1.28	0.11		1		
47305322	**Low discrimination**						**47305322**
					GRA		
Eubacterium contortum		87.65	0.37	FRU 99	99		
Prevotella oralis		6.71	0.16		1		
Bifidobacterium longum subsp. infantis		3.57	0.08		99		
47305502	**Low discrimination**						**47305502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		86.34	0.41	RAF 92			
Eubacterium contortum		8.09	0.14	FRU 99			
Bifidobacterium longum subsp. infantis		3.96	0.04				
47305512	**Good identification**						**47305512**
Bifidobacterium adolescentis		96.22	0.42	RAF 92			
Bifidobacterium longum subsp. longum		2.51	0.13	FRU 99			
Clostridium clostridioforme		1.27	0.07				
47305700	**Very good identification**						**47305700**
					GRA		
Eubacterium contortum		99.96	0.61	FRU 99	99		
Bacteroides vulgatus		0.04	-.05		1		
47305702	**Good identification**						**47305702**
Eubacterium contortum		99.72	0.48	FRU 99			
Clostridium clostridioforme		0.28	0.10				
47305720	**Low discrimination**						**47305720**
					GRA		
Eubacterium contortum		82.61	0.48	FRU 99	99		
Bacteroides vulgatus		17.39	0.30		1		
47305722	**Good identification**						**47305722**
					GRA		
Eubacterium contortum		95.63	0.36	FRU 99	99		
Parabacteroides distasonis		1.69	0.17		1		
Clostridium clostridioforme		1.52	0.10		50		
47307502	**Acceptable identification**						**47307502**
Bifidobacterium adolescentis		93.62	0.39	RAF 92			
Actinomyces israelii		6.39	0.25	FRU 99			
47307512	**Good identification**						**47307512**
Bifidobacterium adolescentis		100.0	0.41	RAF 92			
47310000	**Good identification**						**47310000**
					COC	GRA	
Fusobacterium mortiferum		91.66	0.70	MLT 11	1	1	
Lactobacillus jensenii		3.41	0.36		1	99	
Gemella morbillorum		2.04	0.32		99	99	
47310001	**Acceptable identification**						**47310001**
					GRA		
Fusobacterium mortiferum		92.85	0.36	MLT 11	1		
Actinomyces naeslundii		3.70	0.24	URE 1	99		
Lactobacillus jensenii		3.45	0.02		99		
47310002	**Low discrimination**						**47310002**
					GRA	SPO	
Fusobacterium mortiferum		62.60	0.54	MLT 11	1	1	
Clostridium barati		14.03	0.28	BGL 10	99	99	
Lactobacillus jensenii		13.96	0.33		99	1	
47310022	**Low discrimination**						**47310022**
					GRA	SPO	
Clostridium barati		74.49	0.36	CEL 99	99	99	
Lactobacillus jensenii		6.18	0.23	GAL 95	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		5.78	0.18		1	1	
47310100	**Good identification**						**47310100**
					GRA	SPO	
Fusobacterium mortiferum		91.06	0.77	MLT 11	1	1	
Collinsella aerofaciens		6.05	0.32		99	1	
Clostridium barati		0.97	0.28		99	99	
47310101	**Low discrimination**						**47310101**
					GRA	SPO	
Fusobacterium mortiferum		53.88	0.42	MLT 11	1	1	
Clostridium chauvoei		39.93	0.29	URE 1	99	99	
Collinsella aerofaciens		3.58	-.03		99	1	
47310102	**Low discrimination**						**47310102**
					GRA	SPO	
Fusobacterium mortiferum		46.28	0.60	MLT 11	1	1	
Bifidobacterium longum subsp. infantis		31.11	0.40	BGL 10	99	1	
Clostridium barati		13.34	0.36		99	99	
47310110	**Acceptable identification**						**47310110**
					GRA		
Fusobacterium mortiferum		93.77	0.42	MLT 11	1		
Collinsella aerofaciens		6.23	-.03	MLZ 1	99		
47310122	**Low discrimination**						**47310122**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		43.16	0.44	RAF 95	99	1	
Clostridium barati		31.23	0.44	GAL 99	99	99	
Prevotella oralis		12.65	0.38		1	1	
47310140	**Low discrimination**						**47310140**
					GRA	SPO	
Fusobacterium mortiferum		84.88	0.42	MLT 11	1	1	
Clostridium barati		7.80	0.10	SOR 1	99	99	
Collinsella aerofaciens		5.64	-.03		99	1	
47310200	**Low discrimination**						**47310200**
					GRA	SPO	
Fusobacterium mortiferum		81.59	0.36	MLT 11	1	1	
Actinomyces odontolyticus		9.55	0.15	RHA 1	99	1	
Clostridium barati		5.83	0.01		99	99	
47310300	**Low discrimination**						**47310300**
					GRA	SPO	
Fusobacterium mortiferum		49.04	0.42	MLT 11	1	1	
Eubacterium contortum		42.42	0.29	RHA 1	99	1	
Clostridium barati		4.51	0.10		99	99	
47310400	**Low discrimination**						**47310400**
					COC	GRA	
Fusobacterium mortiferum		77.97	0.36	MLT 11	1	1	
Gemella morbillorum		19.13	0.15	ARA 1	99	99	
Lactobacillus jensenii		2.90	0.02		1	99	
47310500	**Acceptable identification**						**47310500**
					GRA		
Fusobacterium mortiferum		93.77	0.42	MLT 11	1		
Collinsella aerofaciens		6.23	-.03	ARA 1	99		
47311000	**Low discrimination**						**47311000**
					GRA		
Lactobacillus jensenii		52.70	0.70	CEL 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		21.48	0.73		1		
Lactobacillus catenaformis		20.15	0.63		99		
47311001	**Low discrimination**						**47311001**
					GRA	SPO	
Lactobacillus jensenii		38.40	0.36	CEL 95	99	1	
Clostridium chauvoei		18.04	0.29	URE 1	99	99	
Fusobacterium mortiferum		15.65	0.39		1	1	
47311002	**Low discrimination**						**47311002**
					SPO		
Lactobacillus jensenii		51.30	0.67	CEL 95	1		
Lactobacillus catenaformis		29.42	0.63		1		
Clostridium barati		9.90	0.50		99		
47311010	**Very good identification**						**47311010**
					GRA		
Lactobacillus jensenii		97.34	0.62	CEL 95	99		
Fusobacterium mortiferum		1.20	0.39		1		
Lactobacillus catenaformis		1.13	0.29		99		
47311012	**Very good identification**						**47311012**
					SPO		
Lactobacillus jensenii		97.01	0.59	CEL 95	1		
Lactobacillus catenaformis		1.69	0.29		1		
Clostridium barati		0.57	0.15		99		
47311020	**Low discrimination**						**47311020**
					SPO		
Clostridium perfringens		35.42	0.60	LAC 88	84		
Lactobacillus jensenii		30.55	0.60		1		
Clostridium barati		15.72	0.50		99		
47311022	**Low discrimination**						**47311022**
					GRA	SPO	
Clostridium barati		36.82	0.58	CEL 99	99	99	
Lactobacillus jensenii		15.90	0.57		99	1	
Prevotella oralis		14.88	0.52		1	1	
47311030	**Acceptable identification**						**47311030**
					SPO		
Lactobacillus jensenii		93.79	0.52	CEL 95	1		
Clostridium perfringens		3.30	0.26		84		
Clostridium barati		1.46	0.15		99		
47311032	**Low discrimination**						**47311032**
					GRA	SPO	
Lactobacillus jensenii		85.96	0.49	CEL 95	99	1	
Clostridium barati		6.03	0.24		99	99	
Prevotella oralis		2.44	0.18		1	1	
47311040	**Low discrimination**						**47311040**
					GRA		
Lactobacillus jensenii		41.66	0.36	CEL 95	99		
Fusobacterium mortiferum		16.98	0.39	SOR 1	1		
Lactobacillus catenaformis		15.92	0.29		99		
47311060	**Low discrimination**						**47311060**
					SPO		
Clostridium perfringens		76.71	0.48	LAC 88	84		
Clostridium barati		15.54	0.31		99		
Lactobacillus jensenii		3.51	0.26		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47311062	**Low discrimination**						**47311062**
					SPO		
Clostridium barati		50.10	0.40	CEL 99	99		
Clostridium perfringens		38.80	0.42	SOR 8	84		
Lactobacillus jensenii		2.51	0.23		1		
47311100	**Low discrimination**						**47311100**
					GRA		
Collinsella aerofaciens		63.42	0.66	CEL 99	99		
Fusobacterium mortiferum		14.46	0.80		1		
Lactobacillus catenaformis		8.71	0.66		99		
47311101	**Low discrimination**						**47311101**
					GRA	SPO	
Clostridium chauvoei		86.45	0.63	SAL 1	99	99	
Collinsella aerofaciens		7.75	0.32		99	1	
Fusobacterium mortiferum		1.77	0.45		1	1	
47311102	**Low discrimination**						**47311102**
Bifidobacterium longum subsp. infantis		71.33	0.74	RAF 95			
Lactobacillus catenaformis		8.72	0.66				
Collinsella aerofaciens		7.05	0.50				
47311103	**Low discrimination**						**47311103**
					SPO		
Clostridium chauvoei		53.00	0.50	SAL 1	99		
Bifidobacterium longum subsp. infantis		30.26	0.40		1		
Actinomyces naeslundii		4.77	0.55		1		
47311112	**Low discrimination**						**47311112**
					GRA		
Bifidobacterium longum subsp. infantis		36.07	0.40	RAF 95	99		
Leptotrichia buccalis		24.32	0.32	MLZ 1	1		
Bifidobacterium adolescentis		17.58	0.48		99		
47311120	**Low discrimination**						**47311120**
					SPO		
Clostridium paraputrificum		34.60	0.66	CEL 99	99		
Clostridium perfringens		31.48	0.75		84		
Bifidobacterium longum subsp. infantis		12.21	0.61		1		
47311121	**Low discrimination**						**47311121**
Clostridium chauvoei		67.49	0.55	SAL 1			
Clostridium paraputrificum		10.17	0.32				
Clostridium perfringens		9.25	0.41				
47311122	**Low discrimination**						**47311122**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		58.01	0.78	RAF 95	99	1	
Prevotella oralis		17.00	0.72		1	1	
Clostridium barati		8.06	0.66		99	99	
47311123	**Low discrimination**						**47311123**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		46.76	0.44	RAF 95	99	1	
Clostridium chauvoei		15.36	0.42	URE 1	99	99	
Prevotella oralis		13.71	0.38		1	1	
47311132	**Low discrimination**						**47311132**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis		49.55	0.44	RAF 95	99		
Prevotella oralis		14.53	0.38	MLZ 1	1		
Bifidobacterium breve		6.96	0.44		99		
47311142	**Low discrimination**						**47311142**
					SPO		
Bifidobacterium longum subsp. infantis		37.81	0.40	RAF 95	1		
Clostridium barati		26.79	0.40	SOR 1	99		
Lactobacillus acidophilus		10.75	0.35		1		
47311143	**Low discrimination**						**47311143**
					SPO		
Actinomyces naeslundii		42.37	0.42	ESL 12	1		
Clostridium chauvoei		26.96	0.16	RAF 92	99		
Bifidobacterium longum subsp. infantis		15.39	0.06		1		
47311152	**Low discrimination**						**47311152**
Bifidobacterium breve		61.16	0.40	RAF 95			
Bifidobacterium adolescentis		26.22	0.32	CEL 99			
Bifidobacterium longum subsp. infantis		4.40	0.06				
47311160	**Low discrimination**						**47311160**
Clostridium perfringens		63.62	0.63				
Clostridium paraputrificum		27.64	0.47				
Clostridium barati		5.27	0.40				
47311162	**Low discrimination**						**47311162**
					SPO		
Clostridium perfringens		37.23	0.57		84		
Clostridium barati		19.67	0.48		99		
Bifidobacterium longum subsp. infantis		16.45	0.44		1		
47311172	**Low discrimination**						**47311172**
					SPO		
Bifidobacterium breve		67.01	0.44	RAF 95	1		
Clostridium perfringens		10.90	0.23	CEL 99	84		
Clostridium barati		5.76	0.14		99		
47311200	**Low discrimination**						**47311200**
					GRA		
Lactobacillus jensenii		35.85	0.36	CEL 95	99		
Eubacterium contortum		19.67	0.29	RHA 1	99		
Fusobacterium mortiferum		14.62	0.39		1		
47311222	**Low discrimination**						**47311222**
					GRA	SPO	
Prevotella oralis		62.90	0.46	RAF 92	1	1	
Clostridium barati		31.58	0.40	LAC 94	99	99	
Lactobacillus jensenii		1.59	0.23		99	1	
47311300	**Low discrimination**						**47311300**
					SPO		
Eubacterium contortum		87.42	0.63	XYL 99	1		
Collinsella aerofaciens		6.72	0.32		1		
Clostridium barati		1.78	0.31		99		
47311302	**Low discrimination**						**47311302**
					GRA		
Eubacterium contortum		38.17	0.50	XYL 99	99		
Prevotella oralis		21.73	0.44		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis		18.67	0.40		99		
47311320	**Low discrimination**						**47311320**
					GRA	SPO	
Eubacterium contortum		38.46	0.50	XYL 99	99	1	
Prevotella oralis		21.90	0.44		1	1	
Clostridium barati		13.33	0.40		99	99	
47311322	**Low discrimination**						**47311322**
					GRA	SPO	
Prevotella oralis		81.26	0.66	RAF 92	1	1	
Clostridium barati		7.81	0.48		99	99	
Bifidobacterium longum subsp. infantis		6.53	0.44		99	1	
47311400	**Low discrimination**						**47311400**
					GRA		
Lactobacillus jensenii		51.60	0.36	CEL 95	99		
Fusobacterium mortiferum		21.03	0.39	ARA 1	1		
Lactobacillus catenaformis		19.72	0.29		99		
47311502	**Low discrimination**						**47311502**
Bifidobacterium adolescentis		64.40	0.61	RAF 92			
Bifidobacterium longum subsp. infantis		24.13	0.40	XYL 87			
Lactobacillus catenaformis		2.95	0.32				
47311512	**Very good identification**						**47311512**
Bifidobacterium adolescentis		97.95	0.62	RAF 92			
Clostridium clostridioforme		0.86	0.24	XYL 87			
Bifidobacterium longum subsp. longum		0.52	0.22				
47311522	**Low discrimination**						**47311522**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		52.16	0.44	RAF 95	99	1	
Prevotella oralis		15.29	0.38	ARA 1	1	1	
Capnocytophaga ochracea		7.84	0.34		1	90	
47311542	**Acceptable identification**						**47311542**
					SPO		
Bifidobacterium adolescentis		91.82	0.45	RAF 92	1		
Bifidobacterium longum subsp. infantis		2.81	0.06	XYL 87	1		
Clostridium barati		1.99	0.05	SOR 11	99		
47311552	**Good identification**						**47311552**
Bifidobacterium adolescentis		100.0	0.47	RAF 92			
47311700	**Very good identification**						**47311700**
					GRA		
Eubacterium contortum		99.78	0.61	XYL 99	99		
Collinsella aerofaciens		0.10	-.03		99		
Bacteroides vulgatus		0.08	0.01		1		
47311702	**Acceptable identification**						**47311702**
Eubacterium contortum		94.26	0.48	XYL 99			
Clostridium clostridioforme		2.59	0.27				
Bifidobacterium adolescentis		1.52	0.27				
47311712	**Low discrimination**						**47311712**
Clostridium clostridioforme		80.77	0.39	RAF 94			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium adolescentis		12.71	0.28	TRE 88				
Eubacterium contortum		6.52	0.14	XYL 91				
47311720	**Low discrimination**							**47311720**
					GRA			
Eubacterium contortum		69.33	0.48	XYL 99	99			
Bacteroides vulgatus		29.49	0.35		1			
Prevotella oralis		0.49	0.10		1			
47311722	**Low discrimination**							**47311722**
					GRA			
Eubacterium contortum		43.69	0.36	XYL 99	99			
Prevotella oralis		33.06	0.32		1			
Parabacteroides distasonis		8.51	0.35		1			
47311732	**Low discrimination**							**47311732**
					GRA			
Clostridium clostridioforme		85.27	0.39	RAF 94	50			
Parabacteroides distasonis		12.60	0.30	TRE 88	1			
Eubacterium contortum		1.21	0.01	XYL 91	99			
47312000	**Low discrimination**							**47312000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		62.17	0.36	MLT 11	1	1	5	
Clostridium difficile		16.80	0.11	MAN 1	99	99	1	
Propionibacterium granulosum		4.76	0.14		99	1	90	
47312100	**Low discrimination**							**47312100**
					GRA	SPO		
Fusobacterium mortiferum		80.94	0.42	MLT 11	1	1		
Clostridium barati		7.44	0.10	MAN 1	99	99		
Collinsella aerofaciens		5.38	-.03		99	1		
47312101	**Good identification**							**47312101**
Actinomyces naeslundii		100.0	0.36	ESL 12				
47312103	**Good identification**							**47312103**
Actinomyces naeslundii		100.0	0.36	ESL 12				
47312121	**Good identification**							**47312121**
Actinomyces naeslundii		100.0	0.36	ESL 12				
47312123	**Good identification**							**47312123**
Actinomyces naeslundii		100.0	0.36	ESL 12				
47313000	**Low discrimination**							**47313000**
					GRA			
Lactobacillus jensenii		35.00	0.36	CEL 95	99			
Fusobacterium mortiferum		14.27	0.39	MAN 1	1			
Lactobacillus catenaformis		13.38	0.29		99			
47313001	**Good identification**							**47313001**
Actinomyces naeslundii		98.04	0.52	ESL 12				
Lactobacillus jensenii		1.97	0.02	RAF 92				
47313003	**Good identification**							**47313003**
					SPO			
Actinomyces naeslundii		97.83	0.52	ESL 12	1			
Clostridium barati		2.17	-.03	RAF 92	99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47313021	**Good identification**					**47313021**
					SPO	
Actinomyces naeslundii		97.83	0.52	ESL 12	1	
Clostridium barati		2.17	-.03	RAF 92	99	
47313022	**Low discrimination**					**47313022**
					SPO	
Clostridium barati		75.20	0.40	CEL 99	99	
Actinomyces naeslundii		7.61	0.49	MAN 8	1	
Lactobacillus jensenii		3.77	0.23		1	
47313023	**Acceptable identification**					**47313023**
					SPO	
Actinomyces naeslundii		93.76	0.52	ESL 12	1	
Clostridium barati		6.24	0.05	RAF 92	99	
47313040	**Low discrimination**					**47313040**
					SPO	
Actinomyces naeslundii		44.35	0.36	ESL 12	1	
Actinomyces israelii		24.08	0.23	RAF 92	1	
Clostridium barati		24.01	0.05		99	
47313041	**Good identification**					**47313041**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47313043	**Good identification**					**47313043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47313061	**Good identification**					**47313061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47313063	**Good identification**					**47313063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47313101	**Low discrimination**					**47313101**
					SPO	
Actinomyces naeslundii		72.91	0.60	ESL 12	1	
Clostridium chauvoei		23.89	0.29	RAF 92	99	
Collinsella aerofaciens		2.14	-.03		1	
47313102	**Low discrimination**					**47313102**
					SPO	
Bifidobacterium longum subsp. infantis		31.85	0.40	RAF 95	1	
Clostridium barati		22.57	0.40	MAN 1	99	
Bifidobacterium adolescentis		10.39	0.45		1	
47313103	**Low discrimination**					**47313103**
					SPO	
Actinomyces naeslundii		88.26	0.60	ESL 12	1	
Clostridium chauvoei		5.10	0.16	RAF 92	99	
Bifidobacterium longum subsp. infantis		2.91	0.06		1	
47313111	**Good identification**					**47313111**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47313112	**Low discrimination**					**47313112**
					SPO	
Clostridium tertium		42.05	0.39	TRE 90	99	
Bifidobacterium adolescentis		39.17	0.47	CEL 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		13.80	0.40		1	
47313113	**Low discrimination**					**47313113**
					SPO	
Actinomyces naeslundii		65.84	0.38	ESL 12	1	
Clostridium tertium		17.68	0.05	RAF 92	99	
Bifidobacterium adolescentis		16.48	0.12	MLZ 5	1	
47313120	**Low discrimination**					**47313120**
Clostridium barati		27.08	0.40	CEL 99		
Clostridium paraputrificum		19.05	0.32	MAN 8		
Clostridium perfringens		17.33	0.41			
47313121	**Low discrimination**					**47313121**
					SPO	
Actinomyces naeslundii		85.32	0.60	ESL 12	1	
Clostridium chauvoei		9.32	0.21	RAF 92	99	
Clostridium barati		2.00	0.05		99	
47313122	**Low discrimination**					**47313122**
					SPO	
Clostridium barati		32.48	0.48	CEL 99	99	
Bifidobacterium longum subsp. infantis		27.17	0.44	MAN 8	1	
Clostridium tertium		11.04	0.43		99	
47313123	**Low discrimination**					**47313123**
					SPO	
Actinomyces naeslundii		81.82	0.60	ESL 12	1	
Clostridium barati		5.74	0.14	RAF 92	99	
Bifidobacterium longum subsp. infantis		4.80	0.10		1	
47313130	**Low discrimination**					**47313130**
					SPO	
Clostridium tertium		89.13	0.39	TRE 90	99	
Bifidobacterium breve		5.14	0.27	CEL 99	1	
Actinomyces naeslundii		2.24	0.35		1	
47313131	**Low discrimination**					**47313131**
					SPO	
Actinomyces naeslundii		78.83	0.38	ESL 12	1	
Clostridium tertium		21.17	0.05	RAF 92	99	
47313132	**Low discrimination**					**47313132**
					SPO	
Clostridium tertium		80.21	0.47	TRE 90	99	
Bifidobacterium breve		15.60	0.44	CEL 99	1	
Clostridium barati		1.34	0.14		99	
47313133	**Low discrimination**					**47313133**
					SPO	
Actinomyces naeslundii		50.96	0.38	ESL 12	1	
Clostridium tertium		41.06	0.13	RAF 92	99	
Bifidobacterium breve		7.99	0.10	MLZ 5	1	
47313140	**Low discrimination**					**47313140**
					SPO	
Actinomyces naeslundii		31.70	0.44	ESL 12	1	
Actinomyces israelii		24.19	0.33	RAF 92	1	
Clostridium barati		18.09	0.13		99	
47313141	**Good identification**					**47313141**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.47	ESL 12			
47313142	**Low discrimination**						**47313142**
					SPO		
Bifidobacterium breve		40.15	0.40	RAF 95	1		
Clostridium barati		17.60	0.21	CEL 99	99		
Actinomyces israelii		11.77	0.36		1		
47313143	**Good identification**						**47313143**
Actinomyces naeslundii		100.0	0.47	ESL 12			
47313152	**Low discrimination**						**47313152**
Bifidobacterium breve		70.16	0.40	RAF 95			
Bifidobacterium adolescentis		24.61	0.31	CEL 99			
Actinomyces israelii		3.07	0.22				
47313161	**Good identification**						**47313161**
Actinomyces naeslundii		100.0	0.47	ESL 12			
47313162	**Low discrimination**						**47313162**
					SPO		
Bifidobacterium breve		43.17	0.44	RAF 95	1		
Clostridium barati		31.94	0.30	CEL 99	99		
Clostridium perfringens		7.02	0.23		84		
47313163	**Good identification**						**47313163**
Actinomyces naeslundii		95.54	0.47	ESL 12			
Bifidobacterium breve		4.47	0.10	RAF 92			
47313172	**Good identification**						**47313172**
					SPO		
Bifidobacterium breve		95.06	0.44	RAF 95	1		
Clostridium tertium		4.94	0.13	CEL 99	99		
47313322	**Low discrimination**						**47313322**
					SPO	GRA	
Clostridium ramosum		48.47	0.37	TRE 94	92	50	
Prevotella oralis		25.29	0.32	CEL 99	1	1	
Clostridium barati		20.93	0.30		99	99	
47313500	**Low discrimination**						**47313500**
Actinomyces israelii		60.62	0.38	CEL 93			
Bifidobacterium adolescentis		23.14	0.25	XYL 92			
Collinsella aerofaciens		9.43	-.03				
47313502	**Very good identification**						**47313502**
Bifidobacterium adolescentis		95.27	0.59	RAF 92			
Actinomyces israelii		3.78	0.41	XYL 87			
Bifidobacterium longum subsp. infantis		0.44	0.06				
47313512	**Very good identification**						**47313512**
					SPO		
Bifidobacterium adolescentis		99.36	0.61	RAF 92	1		
Actinomyces israelii		0.48	0.27	XYL 87	1		
Clostridium tertium		0.16	0.05		99		
47313542	**Low discrimination**						**47313542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		88.90	0.44	RAF 92			
Actinomyces israelii		11.10	0.34	XYL 87			
47313552	**Good identification**						**47313552**
Bifidobacterium adolescentis		98.50	0.45	RAF 92			
Actinomyces israelii		1.50	0.20	XYL 87			
47314000	**Good identification**						**47314000**
					COC	GRA	
Fusobacterium mortiferum		96.35	0.51	MLT 11	1	1	
Gemella morbillorum		3.17	0.15	XYL 7	99	99	
Lactobacillus jensenii		0.48	0.02		1	99	
47314100	**Low discrimination**						**47314100**
					GRA		
Fusobacterium mortiferum		78.81	0.57	MLT 11	1		
Eubacterium contortum		9.15	0.29	XYL 7	99		
Bifidobacterium longum subsp. infantis		6.33	0.21		99		
47314102	**Low discrimination**						**47314102**
					GRA		
Bifidobacterium longum subsp. infantis		81.90	0.38	RAF 95	99		
Fusobacterium mortiferum		11.21	0.41	GAL 99	1		
Bifidobacterium adolescentis		2.70	0.27		99		
47314122	**Acceptable identification**						**47314122**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		93.49	0.43	RAF 95	99	1	
Prevotella oralis		4.14	0.23	GAL 99	1	1	
Clostridium barati		0.84	0.10		99	99	
47314300	**Very good identification**						**47314300**
					GRA		
Eubacterium contortum		99.91	0.63	GAL 99	99		
Fusobacterium mortiferum		0.09	0.23		1		
47314302	**Good identification**						**47314302**
Eubacterium contortum		99.12	0.50	GAL 99			
Clostridium clostridioforme		0.48	0.16				
Bifidobacterium longum subsp. infantis		0.40	0.04				
47314320	**Good identification**						**47314320**
					GRA		
Eubacterium contortum		98.72	0.50	GAL 99	99		
Bacteroides vulgatus		1.28	0.11		1		
47314322	**Low discrimination**						**47314322**
					GRA		
Eubacterium contortum		85.60	0.37	GAL 99	99		
Prevotella oralis		6.55	0.16		1		
Bifidobacterium longum subsp. infantis		3.49	0.08		99		
47314502	**Low discrimination**						**47314502**
Bifidobacterium adolescentis		79.58	0.41	RAF 92			
Clostridium clostridioforme		9.32	0.22	GAL 99			
Eubacterium contortum		7.45	0.14				
47314512	**Low discrimination**						**47314512**
Bifidobacterium adolescentis		69.62	0.42	RAF 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		30.38	0.33	GAL 99			
47314612	**Good identification**						**47314612**
Clostridium clostridioforme		100.0	0.39	RAF 94			
47314632	**Good identification**						**47314632**
Clostridium clostridioforme		100.0	0.39	RAF 94			
47314700	**Very good identification**						**47314700**
					GRA		
Eubacterium contortum		99.96	0.61	GAL 99	99		
Bacteroides vulgatus		0.04	-.05		1		
47314702	**Acceptable identification**						**47314702**
Eubacterium contortum		91.52	0.48	GAL 99			
Clostridium clostridioforme		8.48	0.36				
47314712	**Good identification**						**47314712**
Clostridium clostridioforme		97.36	0.48	RAF 94			
Eubacterium contortum		2.33	0.14	TRE 88			
Bifidobacterium adolescentis		0.31	0.08				
47314720	**Low discrimination**						**47314720**
					GRA		
Eubacterium contortum		82.61	0.48	GAL 99	99		
Bacteroides vulgatus		17.39	0.30		1		
47314722	**Low discrimination**						**47314722**
					GRA		
Eubacterium contortum		64.33	0.36	GAL 99	99		
Clostridium clostridioforme		33.76	0.36		50		
Parabacteroides distasonis		1.14	0.17		1		
47314732	**Good identification**						**47314732**
					GRA		
Clostridium clostridioforme		99.19	0.48	RAF 94	50		
Eubacterium contortum		0.42	0.01	TRE 88	99		
Parabacteroides distasonis		0.40	0.12		1		
47315000	**Low discrimination**						**47315000**
					GRA		
Fusobacterium mortiferum		50.99	0.54	MLT 11	1		
Lactobacillus jensenii		16.79	0.36	XYL 7	99		
Eubacterium contortum		9.21	0.29		99		
47315002	**Low discrimination**						**47315002**
Bifidobacterium longum subsp. infantis		63.95	0.38	RAF 95			
Lactobacillus jensenii		11.12	0.33	LAC 99			
Lactobacillus catenaformis		6.37	0.29				
47315022	**Low discrimination**						**47315022**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		70.74	0.43	RAF 95	99	1	
Prevotella oralis		19.80	0.36	LAC 99	1	1	
Clostridium barati		4.00	0.24		99	99	
47315100	**Low discrimination**						**47315100**
					COC		
Eubacterium contortum		40.31	0.63	RHA 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis		27.88	0.55		1		
Atopobium parvulum		22.10	0.60		99		
47315102	**Good identification**						**47315102**
Bifidobacterium longum subsp. infantis		92.09	0.72	RAF 95			
Bifidobacterium adolescentis		3.04	0.61				
Eubacterium contortum		2.32	0.50				
47315103	**Acceptable identification**						**47315103**
Bifidobacterium longum subsp. infantis		90.87	0.38	RAF 95			
Bifidobacterium adolescentis		2.99	0.27	URE 1			
Eubacterium contortum		2.29	0.16				
47315112	**Low discrimination**						**47315112**
Bifidobacterium adolescentis		77.97	0.62	RAF 92			
Bifidobacterium longum subsp. infantis		19.56	0.38	ARA 87			
Actinomyces israelii		0.77	0.34				
47315120	**Low discrimination**						**47315120**
					COC		
Bifidobacterium longum subsp. infantis		73.53	0.59	RAF 95	1		
Eubacterium contortum		10.56	0.50	BGL 91	1		
Atopobium parvulum		5.79	0.47		99		
47315122	**Good identification**						**47315122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		94.05	0.77	RAF 95	1	99	
Prevotella oralis		4.17	0.57		1	1	
Bacteroides fragilis		0.47	0.32		99	1	
47315123	**Acceptable identification**						**47315123**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		93.77	0.43	RAF 95	1	99	
Prevotella oralis		4.15	0.23	URE 1	1	1	
Bacteroides fragilis		0.47	-.03		99	1	
47315132	**Low discrimination**						**47315132**
					GRA		
Bifidobacterium longum subsp. infantis		80.45	0.43	RAF 95	99		
Parabacteroides distasonis		8.38	0.42	MLZ 1	1		
Prevotella oralis		3.56	0.23		1		
47315142	**Low discrimination**						**47315142**
Bifidobacterium longum subsp. infantis		63.83	0.38	RAF 95			
Bifidobacterium adolescentis		25.74	0.45	SOR 1			
Actinomyces israelii		6.52	0.41				
47315152	**Good identification**						**47315152**
Bifidobacterium adolescentis		95.10	0.47	RAF 92			
Actinomyces israelii		2.95	0.27	ARA 87			
Bifidobacterium longum subsp. infantis		1.95	0.04	SOR 11			
47315162	**Low discrimination**						**47315162**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		87.15	0.43	RAF 95	99	1	
Prevotella oralis		3.86	0.23	SOR 1	1	1	
Capnocytophaga ochracea		2.52	0.20		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47315200	**Very good identification**					**47315200**
					GRA	
Eubacterium contortum		99.88	0.63	LAC 99	99	
Fusobacterium mortiferum		0.06	0.20		1	
Actinomyces israelii		0.02	0.16		99	
47315202	**Good identification**					**47315202**
					GRA	
Eubacterium contortum		98.42	0.50	LAC 99	99	
Prevotella oralis		0.44	0.08		1	
Clostridium clostridioforme		0.43	0.15		50	
47315220	**Good identification**					**47315220**
					GRA	
Eubacterium contortum		98.17	0.50	LAC 99	99	
Bacteroides vulgatus		1.27	0.11		1	
Prevotella oralis		0.44	0.08		1	
47315222	**Low discrimination**					**47315222**
					GRA	
Eubacterium contortum		62.80	0.37	LAC 99	99	
Prevotella oralis		30.37	0.30		1	
Bifidobacterium longum subsp. infantis		2.56	0.08		99	
47315300	**Excellent identification**					**47315300**
					COC	
Eubacterium contortum		99.98	0.97		1	
Bifidobacterium longum subsp. infantis		0.01	0.21		1	
Atopobium parvulum		0.01	0.25		99	
47315301	**Very good identification**					**47315301**
Eubacterium contortum		100.0	0.63	URE 1		
47315302	**Very good identification**					**47315302**
					GRA	
Eubacterium contortum		99.48	0.84		99	
Bifidobacterium longum subsp. infantis		0.40	0.38		99	
Prevotella oralis		0.07	0.29		1	
47315303	**Good identification**					**47315303**
Eubacterium contortum		99.60	0.50	URE 1		
Bifidobacterium longum subsp. infantis		0.40	0.04			
47315310	**Very good identification**					**47315310**
Eubacterium contortum		100.0	0.63	MLZ 1		
47315312	**Acceptable identification**					**47315312**
Eubacterium contortum		91.77	0.50	MLZ 1		
Clostridium clostridioforme		6.07	0.35			
Bifidobacterium adolescentis		1.48	0.28			
47315320	**Very good identification**					**47315320**
					GRA	
Eubacterium contortum		98.56	0.84		99	
Bacteroides vulgatus		1.28	0.45		1	
Prevotella oralis		0.07	0.29		1	
47315321	**Good identification**					**47315321**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		98.72	0.50	URE 1	99	
Bacteroides vulgatus		1.28	0.11		1	
47315322	**Low discrimination**					**47315322**
					GRA	
Eubacterium contortum		87.44	0.71		99	
Prevotella oralis		6.69	0.51		1	
Bifidobacterium longum subsp. infantis		3.56	0.43		99	
47315323	**Low discrimination**					**47315323**
					GRA	
Eubacterium contortum		87.65	0.37	URE 1	99	
Prevotella oralis		6.71	0.16		1	
Bifidobacterium longum subsp. infantis		3.57	0.08		99	
47315330	**Good identification**					**47315330**
					GRA	
Eubacterium contortum		98.50	0.50	MLZ 1	99	
Bacteroides vulgatus		1.28	0.11		1	
Parabacteroides distasonis		0.22	0.16		1	
47315332	**Low discrimination**					**47315332**
					GRA	
Parabacteroides distasonis		45.77	0.50	RAF 99	1	
Eubacterium contortum		36.35	0.37		99	
Clostridium clostridioforme		13.62	0.35		50	
47315340	**Very good identification**					**47315340**
Eubacterium contortum		99.96	0.63	SOR 1		
Actinomyces israelii		0.04	0.19			
47315342	**Good identification**					**47315342**
Eubacterium contortum		99.13	0.50	SOR 1		
Bifidobacterium longum subsp. infantis		0.40	0.04			
Actinomyces israelii		0.31	0.22			
47315360	**Good identification**					**47315360**
					GRA	
Eubacterium contortum		98.72	0.50	SOR 1	99	
Bacteroides vulgatus		1.28	0.11		1	
47315362	**Low discrimination**					**47315362**
					GRA	
Eubacterium contortum		87.65	0.37	SOR 1	99	
Prevotella oralis		6.71	0.16		1	
Bifidobacterium longum subsp. infantis		3.57	0.08		99	
47315402	**Low discrimination**					**47315402**
Bifidobacterium adolescentis		66.20	0.41	RAF 92		
Actinomyces israelii		16.14	0.36	LAC 99		
Clostridium clostridioforme		6.95	0.21			
47315412	**Low discrimination**					**47315412**
Bifidobacterium adolescentis		70.38	0.42	RAF 92		
Clostridium clostridioforme		27.52	0.32	LAC 99		
Actinomyces israelii		2.10	0.22			
47315500	**Good identification**					**47315500**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		93.38	0.61	RHA 99	1	
Bifidobacterium adolescentis		1.78	0.41		1	
Clostridium butyricum		1.30	0.24		99	
47315502	**Low discrimination**					**47315502**
Bifidobacterium adolescentis		86.29	0.75	RAF 92		
Eubacterium contortum		8.08	0.48			
Bifidobacterium longum subsp. infantis		3.95	0.38			
47315503	**Low discrimination**					**47315503**
Bifidobacterium adolescentis		87.76	0.41	RAF 92		
Eubacterium contortum		8.22	0.14	URE 1		
Bifidobacterium longum subsp. infantis		4.02	0.04			
47315510	**Low discrimination**					**47315510**
Bifidobacterium adolescentis		55.76	0.42	RAF 92		
Eubacterium contortum		24.21	0.27	BGL 99		
Bifidobacterium longum subsp. longum		13.10	0.30			
47315512	**Very good identification**					**47315512**
Bifidobacterium adolescentis		98.22	0.76	RAF 92		
Clostridium clostridioforme		1.30	0.41			
Bifidobacterium longum subsp. longum		0.23	0.30			
47315513	**Good identification**					**47315513**
Bifidobacterium adolescentis		98.70	0.42	RAF 92		
Clostridium clostridioforme		1.31	0.07	URE 1		
47315520	**Low discrimination**					**47315520**
					GRA	
Bacteroides vulgatus		66.07	0.48	RAF 97	1	
Eubacterium contortum		28.54	0.48	RHA 90	99	
Bifidobacterium longum subsp. infantis		2.45	0.25		99	
47315522	**Low discrimination**					**47315522**
Bifidobacterium longum subsp. infantis		57.91	0.43	RAF 95		
Eubacterium contortum		11.75	0.36	ARA 1		
Bifidobacterium adolescentis		7.18	0.41			
47315530	**Low discrimination**					**47315530**
					GRA	
Bifidobacterium longum subsp. longum		77.22	0.40	SAL 1	99	
Bacteroides vulgatus		14.57	0.13	RAF 97	1	
Eubacterium contortum		6.29	0.14		99	
47315532	**Low discrimination**					**47315532**
Clostridium clostridioforme		34.60	0.41	RAF 94		
Bifidobacterium adolescentis		26.43	0.42	TRE 88		
Bifidobacterium longum subsp. longum		24.84	0.40	RHA 88		
47315542	**Very good identification**					**47315542**
Bifidobacterium adolescentis		95.75	0.59	RAF 92		
Actinomyces israelii		2.97	0.40	SOR 11		
Eubacterium contortum		0.73	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47315552	**Very good identification**					**47315552**
Bifidobacterium adolescentis		99.52	0.61	RAF 92		
Actinomyces israelii		0.38	0.26	SOR 11		
Clostridium clostridioforme		0.11	0.07			
47315600	**Very good identification**					**47315600**
					GRA	
Eubacterium contortum		99.94	0.61	LAC 99	99	
Bacteroides vulgatus		0.04	-.05		1	
Actinomyces israelii		0.02	0.14		99	
47315602	**Acceptable identification**					**47315602**
Eubacterium contortum		92.17	0.48	LAC 99		
Clostridium clostridioforme		7.65	0.36			
Actinomyces israelii		0.18	0.17			
47315612	**Good identification**					**47315612**
Clostridium clostridioforme		97.07	0.47	RAF 94		
Eubacterium contortum		2.59	0.14	TRE 88		
Bifidobacterium adolescentis		0.34	0.08			
47315620	**Low discrimination**					**47315620**
					GRA	
Eubacterium contortum		82.61	0.48	LAC 99	99	
Bacteroides vulgatus		17.39	0.30		1	
47315622	**Low discrimination**					**47315622**
					GRA	
Eubacterium contortum		66.66	0.36	LAC 99	99	
Clostridium clostridioforme		31.35	0.36		50	
Parabacteroides distasonis		1.18	0.17		1	
47315632	**Good identification**					**47315632**
					GRA	
Clostridium clostridioforme		99.09	0.47	RAF 94	50	
Eubacterium contortum		0.47	0.01	TRE 88	99	
Parabacteroides distasonis		0.44	0.12		1	
47315700	**Excellent identification**					**47315700**
					GRA	
Eubacterium contortum		99.96	0.96		99	
Bacteroides vulgatus		0.04	0.30		1	
Actinomyces israelii		0.00	0.24		99	
47315701	**Very good identification**					**47315701**
					GRA	
Eubacterium contortum		99.96	0.61	URE 1	99	
Bacteroides vulgatus		0.04	-.05		1	
47315702	**Very good identification**					**47315702**
Eubacterium contortum		99.59	0.83			
Clostridium clostridioforme		0.28	0.45			
Bifidobacterium adolescentis		0.11	0.41			
47315703	**Good identification**					**47315703**
Eubacterium contortum		99.72	0.48	URE 1		
Clostridium clostridioforme		0.28	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47315710	**Very good identification**					**47315710**
					GRA	
Eubacterium contortum		99.71	0.61	MLZ 1	99	
Clostridium clostridioforme		0.23	0.22		50	
Bacteroides vulgatus		0.04	-.05		1	
47315712	**Low discrimination**					**47315712**
Clostridium clostridioforme		52.73	0.56	RAF 94		
Eubacterium contortum		41.66	0.48	TRE 88		
Bifidobacterium adolescentis		5.49	0.42			
47315720	**Low discrimination**					**47315720**
					GRA	
Eubacterium contortum		82.61	0.83		99	
Bacteroides vulgatus		17.39	0.64		1	
Parabacteroides distasonis		0.00	0.17		1	
47315721	**Low discrimination**					**47315721**
					GRA	
Eubacterium contortum		82.61	0.48	URE 1	99	
Bacteroides vulgatus		17.39	0.30		1	
47315722	**Very good identification**					**47315722**
					GRA	
Eubacterium contortum		95.40	0.70		99	
Parabacteroides distasonis		1.69	0.51		1	
Clostridium clostridioforme		1.52	0.45		50	
47315723	**Good identification**					**47315723**
					GRA	
Eubacterium contortum		95.63	0.36	URE 1	99	
Parabacteroides distasonis		1.69	0.17		1	
Clostridium clostridioforme		1.52	0.10		50	
47315730	**Low discrimination**					**47315730**
					GRA	
Eubacterium contortum		81.55	0.48	MLZ 1	99	
Bacteroides vulgatus		17.17	0.30		1	
Clostridium clostridioforme		1.04	0.22		50	
47315732	**Low discrimination**					**47315732**
					GRA	
Clostridium clostridioforme		78.45	0.56	RAF 94	50	
Eubacterium contortum		10.94	0.36	TRE 88	99	
Parabacteroides distasonis		10.32	0.46		1	
47315740	**Very good identification**					**47315740**
					GRA	
Eubacterium contortum		99.93	0.61	SOR 1	99	
Bacteroides vulgatus		0.04	-.05		1	
Actinomyces israelii		0.04	0.17		99	
47315742	**Good identification**					**47315742**
Eubacterium contortum		98.12	0.48	SOR 1		
Bifidobacterium adolescentis		1.31	0.25			
Actinomyces israelii		0.30	0.20			
47315760	**Low discrimination**					**47315760**
					GRA	
Eubacterium contortum		82.61	0.48	SOR 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		17.39	0.30		1		
47315762	**Good identification**						**47315762**
					GRA		
Eubacterium contortum		95.63	0.36	SOR 1	99		
Parabacteroides distasonis		1.69	0.17		1		
Clostridium clostridioforme		1.52	0.10		50		
47316502	**Acceptable identification**						**47316502**
Bifidobacterium adolescentis		93.62	0.39	RAF 92			
Actinomyces israelii		6.39	0.25	GAL 99			
47316512	**Good identification**						**47316512**
Bifidobacterium adolescentis		100.0	0.41	RAF 92			
47317000	**Acceptable identification**						**47317000**
					GRA	SPO	
Actinomyces israelii		94.80	0.48	CEL 93	99	1	
Fusobacterium mortiferum		3.06	0.20		1	1	
Clostridium innocuum		1.13	0.00		99	99	
47317002	**Low discrimination**						**47317002**
Actinomyces israelii		88.78	0.51	CEL 93			
Bifidobacterium adolescentis		6.43	0.25				
Bifidobacterium longum subsp. infantis		2.41	0.04				
47317012	**Low discrimination**						**47317012**
					SPO		
Actinomyces israelii		58.02	0.37	CEL 93	1		
Bifidobacterium adolescentis		34.37	0.27	MLZ 13	1		
Clostridium tertium		7.61	0.07		99		
47317040	**Good identification**						**47317040**
Actinomyces israelii		100.0	0.41	CEL 93			
47317042	**Good identification**						**47317042**
Actinomyces israelii		97.75	0.44	CEL 93			
Bifidobacterium adolescentis		2.25	0.09				
47317100	**Low discrimination**						**47317100**
					SPO		
Actinomyces israelii		65.72	0.58	CEL 93	1		
Eubacterium contortum		9.47	0.29		1		
Clostridium tertium		8.97	0.29		99		
47317102	**Low discrimination**						**47317102**
Bifidobacterium adolescentis		47.50	0.59	RAF 92			
Actinomyces israelii		26.50	0.61	ARA 87			
Bifidobacterium longum subsp. infantis		17.80	0.38				
47317112	**Low discrimination**						**47317112**
					SPO		
Bifidobacterium adolescentis		77.35	0.61	RAF 92	1		
Clostridium tertium		17.13	0.41	ARA 87	99		
Actinomyces israelii		5.28	0.47		1		
47317120	**Low discrimination**						**47317120**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		56.81	0.37	TRE 90	99	
Bifidobacterium longum subsp. infantis		24.58	0.25	CEL 99	1	
Actinomyces israelii		8.86	0.38		1	
47317122	**Low discrimination**					**47317122**
					SPO	
Bifidobacterium longum subsp. infantis		54.49	0.43	RAF 95	1	
Clostridium tertium		37.37	0.45	MAN 1	99	
Actinomyces israelii		2.91	0.41		1	
47317130	**Good identification**					**47317130**
					SPO	
Clostridium tertium		98.71	0.41	TRE 90	99	
Actinomyces israelii		1.29	0.24	CEL 99	1	
47317132	**Good identification**					**47317132**
					SPO	
Clostridium tertium		96.95	0.50	TRE 90	99	
Bifidobacterium adolescentis		1.47	0.27	CEL 99	1	
Bifidobacterium longum subsp. infantis		0.80	0.08		1	
47317140	**Good identification**					**47317140**
Actinomyces israelii		99.14	0.51	CEL 93		
Bifidobacterium adolescentis		0.86	0.09			
47317142	**Low discrimination**					**47317142**
Actinomyces israelii		62.73	0.54	CEL 93		
Bifidobacterium adolescentis		35.73	0.44			
Bifidobacterium longum subsp. infantis		1.09	0.04			
47317150	**Low discrimination**					**47317150**
					SPO	
Actinomyces israelii		89.87	0.37	CEL 93	1	
Bifidobacterium adolescentis		6.34	0.11	MLZ 13	1	
Clostridium tertium		3.79	-.01		99	
47317152	**Low discrimination**					**47317152**
					SPO	
Bifidobacterium adolescentis		81.12	0.45	RAF 92	1	
Actinomyces israelii		17.41	0.40	ARA 87	1	
Clostridium tertium		1.47	0.07	SOR 11	99	
47317300	**Very good identification**					**47317300**
Eubacterium contortum		99.48	0.63	MAN 1		
Actinomyces israelii		0.53	0.39			
47317302	**Acceptable identification**					**47317302**
Eubacterium contortum		94.22	0.50	MAN 1		
Actinomyces israelii		4.23	0.42			
Bifidobacterium adolescentis		1.02	0.25			
47317320	**Good identification**					**47317320**
					GRA	
Eubacterium contortum		98.38	0.50	MAN 1	99	
Bacteroides vulgatus		1.28	0.11		1	
Actinomyces israelii		0.19	0.18		99	
47317322	**Low discrimination**					**47317322**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		83.54	0.37	MAN 1	99	
Prevotella oralis		6.39	0.16		1	
Bifidobacterium longum subsp. infantis		3.40	0.08		99	
47317400	**Good identification**					**47317400**
Actinomyces israelii		100.0	0.46	CEL 93		
47317402	**Low discrimination**					**47317402**
Actinomyces israelii		62.80	0.49	CEL 93		
Bifidobacterium adolescentis		37.20	0.39			
47317412	**Low discrimination**					**47317412**
Bifidobacterium adolescentis		82.89	0.41	RAF 92		
Actinomyces israelii		17.11	0.35	LAC 99		
47317440	**Good identification**					**47317440**
Actinomyces israelii		100.0	0.39	CEL 93		
47317442	**Low discrimination**					**47317442**
Actinomyces israelii		84.16	0.42	CEL 93		
Bifidobacterium adolescentis		15.84	0.24			
47317500	**Low discrimination**					**47317500**
Actinomyces israelii		72.03	0.57	CEL 93		
Bifidobacterium adolescentis		16.00	0.39			
Eubacterium contortum		10.38	0.27			
47317502	**Good identification**					**47317502**
Bifidobacterium adolescentis		93.41	0.73	RAF 92		
Actinomyces israelii		6.37	0.60			
Eubacterium contortum		0.11	0.14			
47317503	**Acceptable identification**					**47317503**
Bifidobacterium adolescentis		93.62	0.39	RAF 92		
Actinomyces israelii		6.39	0.25	URE 1		
47317510	**Low discrimination**					**47317510**
					SPO	
Bifidobacterium adolescentis		64.05	0.41	RAF 92	1	
Actinomyces israelii		35.25	0.42	BGL 99	1	
Clostridium tertium		0.71	-.01		99	
47317512	**Very good identification**					**47317512**
Bifidobacterium adolescentis		99.02	0.75	RAF 92		
Actinomyces israelii		0.83	0.45			
Bifidobacterium dentium		0.10	0.06			
47317513	**Good identification**					**47317513**
Bifidobacterium adolescentis		100.0	0.41	RAF 92		
47317520	**Low discrimination**					**47317520**
					GRA	
Actinomyces israelii		41.67	0.36	CEL 93	99	
Bacteroides vulgatus		38.43	0.13	NAG 6	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		16.60	0.14		99	
47317522	**Low discrimination**					**47317522**
Bifidobacterium adolescentis		61.77	0.39	RAF 92		
Actinomyces israelii		26.62	0.39	NAG 1		
Bifidobacterium longum subsp. infantis		6.15	0.08			
47317532	**Low discrimination**					**47317532**
					SPO	
Bifidobacterium adolescentis		73.59	0.41	RAF 92	1	
Bifidobacterium dentium		13.18	0.10	NAG 1	1	
Clostridium tertium		7.31	0.15		99	
47317540	**Acceptable identification**					**47317540**
Actinomyces israelii		93.41	0.50	CEL 93		
Bifidobacterium adolescentis		6.59	0.24			
47317542	**Low discrimination**					**47317542**
Bifidobacterium adolescentis		82.33	0.58	RAF 92		
Actinomyces israelii		17.67	0.53	SOR 11		
47317550	**Low discrimination**					**47317550**
					COC	
Actinomyces israelii		60.01	0.35	CEL 93	1	
Bifidobacterium adolescentis		34.65	0.25	MLZ 13	1	
Blautia producta		5.35	-.03		99	
47317552	**Very good identification**					**47317552**
Bifidobacterium adolescentis		97.44	0.59	RAF 92		
Actinomyces israelii		2.56	0.38	SOR 11		
47317700	**Very good identification**					**47317700**
					GRA	
Eubacterium contortum		99.44	0.61	MAN 1	99	
Actinomyces israelii		0.53	0.37		99	
Bacteroides vulgatus		0.04	-.05		1	
47317702	**Low discrimination**					**47317702**
Eubacterium contortum		88.03	0.48	MAN 1		
Bifidobacterium adolescentis		7.77	0.39			
Actinomyces israelii		3.95	0.40			
47317712	**Low discrimination**					**47317712**
Bifidobacterium adolescentis		78.47	0.41	RAF 92		
Clostridium clostridioforme		9.30	0.22	RHA 1		
Eubacterium contortum		7.35	0.14			
47317720	**Low discrimination**					**47317720**
					GRA	
Eubacterium contortum		82.48	0.48	MAN 1	99	
Bacteroides vulgatus		17.36	0.30		1	
Actinomyces israelii		0.16	0.17		99	
47317722	**Acceptable identification**					**47317722**
					GRA	
Eubacterium contortum		94.17	0.36	MAN 1	99	
Parabacteroides distasonis		1.67	0.17		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.53	0.20		99		
47321000	**Acceptable identification**						**47321000**
					COC	GRA	
Lactobacillus jensenii		90.03	0.44	FRU 99	1	99	
Atopobium parvulum		4.86	0.21	CEL 95	99	99	
Fusobacterium mortiferum		3.65	0.30		1	1	
47321002	**Good identification**						**47321002**
					COC		
Lactobacillus jensenii		95.12	0.41	FRU 99	1		
Lactobacillus acidophilus		1.51	0.07	CEL 95	1		
Atopobium parvulum		1.36	0.08		99		
47321010	**Good identification**						**47321010**
Lactobacillus jensenii		100.0	0.36	FRU 99			
47321100	**Good identification**						**47321100**
					COC	GRA	
Atopobium parvulum		96.55	0.55	FRU 99	99	99	
Fusobacterium mortiferum		1.71	0.36		1	1	
Clostridium perfringens		0.74	0.19		1	99	
47321101	**Acceptable identification**						**47321101**
					COC	SPO	
Clostridium chauvoei		90.32	0.40	SAL 1	1	99	
Atopobium parvulum		6.60	0.21	TRE 10	99	1	
Actinomyces naeslundii		3.08	0.38		1	1	
47321102	**Low discrimination**						**47321102**
					COC		
Atopobium parvulum		81.32	0.42	FRU 99	99		
Lactobacillus acidophilus		10.48	0.25		1		
Clostridium perfringens		1.65	0.14		1		
47321120	**Low discrimination**						**47321120**
					COC	SPO	
Clostridium perfringens		49.32	0.45	FRU 99	1	84	
Atopobium parvulum		35.35	0.42		99	1	
Clostridium paraputrificum		7.76	0.22		1	99	
47321122	**Low discrimination**						**47321122**
					COC		
Clostridium perfringens		54.14	0.40	FRU 99	1		
Atopobium parvulum		14.55	0.29		99		
Lactobacillus acidophilus		10.63	0.25		1		
47323001	**Good identification**						**47323001**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47323003	**Good identification**						**47323003**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47323021	**Good identification**						**47323021**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47323023	**Good identification**						**47323023**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47323101	**Low discrimination**						**47323101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		86.75	0.43	ESL 12	1		
Clostridium chauvoei		13.25	0.06	RAF 92	99		
47323103	**Good identification**						**47323103**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47323121	**Good identification**						**47323121**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47323123	**Good identification**						**47323123**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47325100	**Very good identification**						**47325100**
					COC	GRA	
Atopobium parvulum		99.81	0.60	FRU 99	99	99	
Actinomyces israelii		0.09	0.21		1	99	
Fusobacterium mortiferum		0.07	0.17		1	1	
47325102	**Good identification**						**47325102**
					COC		
Atopobium parvulum		97.02	0.47	FRU 99	99		
Bifidobacterium adolescentis		1.56	0.23		1		
Bifidobacterium longum subsp. infantis		0.72	0.04		1		
47325120	**Good identification**						**47325120**
					COC		
Atopobium parvulum		99.25	0.47	FRU 99	99		
Clostridium perfringens		0.75	0.11		1		
47325502	**Low discrimination**						**47325502**
					COC	SPO	
Bifidobacterium adolescentis		79.61	0.38	RAF 92	1	1	
Atopobium parvulum		7.49	0.13	FRU 99	99	1	
Clostridium butyricum		5.37	0.03		1	99	
47325512	**Low discrimination**						**47325512**
Bifidobacterium adolescentis		87.30	0.39	RAF 92			
Clostridium clostridioforme		12.70	0.22	FRU 99			
47325712	**Good identification**						**47325712**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47325732	**Acceptable identification**						**47325732**
Clostridium clostridioforme		92.45	0.36	RAF 94			
Parabacteroides distasonis		7.55	0.23	FRU 99			
47327102	**Low discrimination**						**47327102**
					SPO		
Actinomyces israelii		45.14	0.37	FRU 99	1		
Clostridium tertium		29.48	0.19	CEL 93	99		
Bifidobacterium adolescentis		14.34	0.22		1		
47327502	**Low discrimination**						**47327502**
Bifidobacterium adolescentis		72.21	0.36	RAF 92			
Actinomyces israelii		27.79	0.35	FRU 99			
47327512	**Low discrimination**						**47327512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		83.46	0.38	RAF 92			
Bifidobacterium dentium		12.62	0.06	FRU 99			
Actinomyces israelii		3.93	0.21				
47330000	**Low discrimination**						**47330000**
					COC	GRA	
Fusobacterium mortiferum		70.84	0.61	MLT 11	1	1	
Lactobacillus jensenii		26.45	0.44		1	99	
Atopobium parvulum		1.43	0.21		99	99	
47330002	**Low discrimination**						**47330002**
					GRA	SPO	
Lactobacillus jensenii		65.66	0.41	CEL 95	99	1	
Fusobacterium mortiferum		29.31	0.45	GAL 99	1	1	
Clostridium barati		1.91	0.10		99	99	
47330010	**Acceptable identification**						**47330010**
					GRA	CAT	
Lactobacillus jensenii		91.54	0.36	CEL 95	99	1	
Fusobacterium mortiferum		7.43	0.27	GAL 99	1	5	
Propionibacterium granulosum		1.03	0.09		99	90	
47330100	**Low discrimination**						**47330100**
					COC	GRA	
Fusobacterium mortiferum		53.47	0.68	MLT 11	1	1	
Atopobium parvulum		45.70	0.55		99	99	
Clostridium perfringens		0.35	0.19		1	99	
47330102	**Low discrimination**						**47330102**
					COC	GRA	
Atopobium parvulum		49.97	0.42	GAL 99	99	99	
Fusobacterium mortiferum		36.81	0.51		1	1	
Lactobacillus acidophilus		6.44	0.25		1	99	
47330103	**Low discrimination**						**47330103**
					COC	GRA	
Actinomyces naeslundii		50.39	0.36	ESL 12	1	99	
Atopobium parvulum		24.09	0.08	RAF 92	99	99	
Fusobacterium mortiferum		17.75	0.17	GAL 96	1	1	
47330120	**Low discrimination**						**47330120**
					COC	SPO	
Clostridium perfringens		47.50	0.45	GAL 99	1	84	
Atopobium parvulum		34.04	0.42		99	1	
Clostridium paraputrificum		7.47	0.22		1	99	
47330121	**Low discrimination**						**47330121**
					COC		
Actinomyces naeslundii		41.04	0.36	ESL 12	1		
Clostridium perfringens		27.38	0.11	RAF 92	1		
Atopobium parvulum		19.62	0.08	GAL 96	99		
47330122	**Low discrimination**						**47330122**
					COC	SPO	
Clostridium perfringens		48.06	0.40	GAL 99	1	84	
Clostridium barati		13.57	0.26		1	99	
Atopobium parvulum		12.91	0.29		99	1	
47330123	**Low discrimination**						**47330123**
Actinomyces naeslundii		76.10	0.36	ESL 12			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		23.90	0.06	RAF 92			
47330612	**Good identification**						**47330612**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47330632	**Good identification**						**47330632**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47330712	**Good identification**						**47330712**
Clostridium clostridioforme		100.0	0.45	RAF 94			
47330732	**Good identification**						**47330732**
Clostridium clostridioforme		100.0	0.45	RAF 94			
47331000	**Good identification**						**47331000**
					COC	GRA	
Lactobacillus jensenii		90.54	0.78	CEL 95	1	99	
Atopobium parvulum		4.89	0.55		99	99	
Fusobacterium mortiferum		3.68	0.64		1	1	
47331001	**Low discrimination**						**47331001**
					COC		
Lactobacillus jensenii		77.79	0.44	CEL 95	1		
Actinomyces naeslundii		13.07	0.52	URE 1	1		
Atopobium parvulum		4.20	0.21		99		
47331002	**Very good identification**						**47331002**
					COC		
Lactobacillus jensenii		95.12	0.75	CEL 95	1		
Lactobacillus acidophilus		1.51	0.41		1		
Atopobium parvulum		1.36	0.42		99		
47331003	**Low discrimination**						**47331003**
Lactobacillus jensenii		76.98	0.41	CEL 95			
Actinomyces naeslundii		19.39	0.52	URE 1			
Lactobacillus acidophilus		1.22	0.07				
47331010	**Very good identification**						**47331010**
					COC	GRA	
Lactobacillus jensenii		99.63	0.70	CEL 95	1	99	
Atopobium parvulum		0.16	0.21		99	99	
Fusobacterium mortiferum		0.12	0.30		1	1	
47331011	**Good identification**						**47331011**
Lactobacillus jensenii		97.42	0.36	CEL 95			
Actinomyces naeslundii		2.58	0.30	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47331012	**Very good identification**					**47331012**
					GRA	
Lactobacillus jensenii		95.22	0.67	CEL 95	99	
Leptotrichia buccalis		4.56	0.32		1	
Lactobacillus acidophilus		0.05	0.07		99	
47331020	**Low discrimination**					**47331020**
					COC	
Lactobacillus jensenii		55.72	0.68	CEL 95	1	
Clostridium perfringens		39.99	0.65		1	
Atopobium parvulum		2.12	0.42		99	
47331022	**Low discrimination**					**47331022**
					SPO	
Lactobacillus jensenii		59.38	0.65	CEL 95	1	
Clostridium perfringens		30.08	0.59		84	
Clostridium barati		3.98	0.40		99	
47331023	**Low discrimination**					**47331023**
Actinomyces naeslundii		37.82	0.52	ESL 12		
Lactobacillus jensenii		37.53	0.31	RAF 92		
Clostridium perfringens		19.01	0.25			
47331030	**Very good identification**					**47331030**
					COC	
Lactobacillus jensenii		97.65	0.60	CEL 95	1	
Clostridium perfringens		2.12	0.30		1	
Atopobium parvulum		0.11	0.08		99	
47331032	**Very good identification**					**47331032**
					GRA	
Lactobacillus jensenii		95.84	0.57	CEL 95	99	
Leptotrichia buccalis		2.04	0.15		1	
Clostridium perfringens		1.47	0.25		99	
47331040	**Low discrimination**					**47331040**
					COC	
Lactobacillus jensenii		80.57	0.44	CEL 95	1	
Clostridium perfringens		8.43	0.26	SOR 1	1	
Atopobium parvulum		4.35	0.21		99	
47331041	**Low discrimination**					**47331041**
Actinomyces naeslundii		74.58	0.39	ESL 12		
Lactobacillus jensenii		25.42	0.10	RAF 92		
47331042	**Low discrimination**					**47331042**
Lactobacillus jensenii		72.32	0.41	CEL 95		
Lactobacillus acidophilus		14.03	0.26	SOR 1		
Clostridium perfringens		5.34	0.21			
47331043	**Low discrimination**					**47331043**
Actinomyces naeslundii		81.49	0.39	ESL 12		
Lactobacillus jensenii		18.51	0.07	RAF 92		
47331050	**Good identification**					**47331050**
Lactobacillus jensenii		100.0	0.36	CEL 95		
47331060	**Acceptable identification**					**47331060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium perfringens		91.19	0.52	LAC 88	84		
Lactobacillus jensenii		6.74	0.34		1		
Clostridium barati		0.87	0.13		99		
47331062	**Low discrimination**						**47331062**
Clostridium perfringens		79.05	0.47	LAC 88			
Lactobacillus jensenii		8.28	0.31				
Lactobacillus acidophilus		6.42	0.26				
47331063	**Low discrimination**						**47331063**
Actinomyces naeslundii		64.83	0.39	ESL 12			
Clostridium perfringens		35.17	0.13	RAF 92			
47331100	**Very good identification**						**47331100**
					COC	GRA	
Atopobium parvulum		96.80	0.89		99	99	
Fusobacterium mortiferum		1.72	0.71		1	1	
Clostridium perfringens		0.74	0.54		1	99	
47331101	**Low discrimination**						**47331101**
					COC	SPO	
Atopobium parvulum		69.32	0.55	URE 1	99	1	
Clostridium chauvoei		22.36	0.47		1	99	
Actinomyces naeslundii		6.20	0.60		1	1	
47331102	**Low discrimination**						**47331102**
					COC		
Atopobium parvulum		81.55	0.76		99		
Lactobacillus acidophilus		10.51	0.60		1		
Clostridium perfringens		1.66	0.48		1		
47331103	**Low discrimination**						**47331103**
					COC	SPO	
Atopobium parvulum		48.99	0.42	URE 1	99	1	
Actinomyces naeslundii		24.84	0.60		1	1	
Clostridium chauvoei		15.80	0.34		1	99	
47331110	**Low discrimination**						**47331110**
					COC	GRA	
Atopobium parvulum		76.77	0.55	MLZ 1	99	99	
Leptotrichia buccalis		15.14	0.32		1	1	
Lactobacillus jensenii		4.79	0.36		1	99	
47331112	**Very good identification**						**47331112**
					COC	GRA	
Leptotrichia buccalis		98.02	0.66	CEL 99	1	1	
Atopobium parvulum		0.89	0.42		99	99	
Bifidobacterium adolescentis		0.48	0.45		1	99	
47331120	**Low discrimination**						**47331120**
					COC	SPO	
Clostridium perfringens		49.41	0.79		1	84	
Atopobium parvulum		35.41	0.76		99	1	
Clostridium paraputrificum		7.77	0.56		1	99	
47331121	**Low discrimination**						**47331121**
					COC	SPO	
Clostridium perfringens		35.57	0.45	URE 1	1	84	
Atopobium parvulum		25.49	0.42		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		15.53	0.38		1	99	
47331122	**Low discrimination**						**47331122**
					COC		
Clostridium perfringens		54.31	0.74		1		
Atopobium parvulum		14.59	0.63		99		
Lactobacillus acidophilus		10.66	0.60		1		
47331123	**Low discrimination**						**47331123**
					COC		
Clostridium perfringens		36.29	0.40	URE 1	1		
Actinomyces naeslundii		28.02	0.60		1		
Atopobium parvulum		9.75	0.29		99		
47331130	**Low discrimination**						**47331130**
					COC	SPO	
Clostridium perfringens		44.60	0.45	MLZ 1	1	84	
Atopobium parvulum		31.96	0.42		99	1	
Clostridium paraputrificum		7.02	0.22		1	99	
47331131	**Low discrimination**						**47331131**
					COC		
Actinomyces naeslundii		46.79	0.38	ESL 12	1		
Clostridium perfringens		24.71	0.11	RAF 92	1		
Atopobium parvulum		17.71	0.08	MLZ 5	99		
47331132	**Low discrimination**						**47331132**
					GRA		
Leptotrichia buccalis		86.64	0.50	CEL 99	1		
Clostridium perfringens		4.63	0.40	NAG 10	99		
Bifidobacterium breve		1.95	0.41		99		
47331140	**Low discrimination**						**47331140**
					COC	GRA	
Atopobium parvulum		83.60	0.55	SOR 1	99	99	
Clostridium perfringens		12.01	0.41		1	99	
Fusobacterium mortiferum		1.48	0.36		1	1	
47331141	**Low discrimination**						**47331141**
					COC	SPO	
Actinomyces naeslundii		51.60	0.47	ESL 12	1	1	
Atopobium parvulum		33.01	0.21	RAF 92	99	1	
Clostridium chauvoei		10.65	0.12		1	99	
47331142	**Low discrimination**						**47331142**
					COC		
Lactobacillus acidophilus		45.98	0.44	CEL 99	1		
Atopobium parvulum		29.15	0.42	SOR 11	99		
Clostridium perfringens		11.16	0.36		1		
47331143	**Low discrimination**						**47331143**
					COC		
Actinomyces naeslundii		72.96	0.47	ESL 12	1		
Lactobacillus acidophilus		12.99	0.10	RAF 92	1		
Atopobium parvulum		8.24	0.08		99		
47331160	**Low discrimination**						**47331160**
					COC	SPO	
Clostridium perfringens		89.70	0.67		1	84	
Clostridium paraputrificum		5.58	0.37		1	99	
Atopobium parvulum		3.41	0.42		99	1	
47331162	**Low discrimination**						**47331162**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium perfringens		79.46	0.62		84		
Lactobacillus acidophilus		10.12	0.44		1		
Clostridium paraputrificum		4.50	0.31		99		
47331172	**Low discrimination**						**47331172**
					GRA		
Bifidobacterium breve		50.34	0.41	RAF 95	99		
Clostridium perfringens		22.81	0.27	CEL 99	99		
Leptotrichia buccalis		22.65	0.15		1		
47331200	**Acceptable identification**						**47331200**
					COC	GRA	
Lactobacillus jensenii		90.03	0.44	CEL 95	1	99	
Atopobium parvulum		4.86	0.21	RHA 1	99	99	
Fusobacterium mortiferum		3.65	0.30		1	1	
47331202	**Low discrimination**						**47331202**
					SPO		
Lactobacillus jensenii		82.41	0.41	CEL 95	1		
Lactobacillus acidophilus		9.74	0.22	RHA 1	1		
Clostridium barati		3.97	0.13		99		
47331210	**Good identification**						**47331210**
Lactobacillus jensenii		100.0	0.36	CEL 95			
47331300	**Acceptable identification**						**47331300**
					COC	GRA	
Atopobium parvulum		92.98	0.55	RHA 1	99	99	
Eubacterium contortum		3.18	0.29		1	99	
Fusobacterium mortiferum		1.65	0.36		1	1	
47331302	**Low discrimination**						**47331302**
					COC	GRA	
Atopobium parvulum		42.27	0.42	RHA 1	99	99	
Lactobacillus acidophilus		40.60	0.40		1	99	
Prevotella oralis		5.20	0.24		1	1	
47331320	**Low discrimination**						**47331320**
					COC	SPO	
Clostridium perfringens		39.15	0.45	RHA 1	1	84	
Atopobium parvulum		28.05	0.42		99	1	
Clostridium ramosum		13.75	0.35		1	92	
47331322	**Low discrimination**						**47331322**
					SPO	GRA	
Prevotella oralis		32.13	0.46	RAF 92	1	1	
Clostridium ramosum		26.94	0.45	TRE 6	92	50	
Lactobacillus acidophilus		13.19	0.40		1	99	
47331332	**Low discrimination**						**47331332**
Parabacteroides distasonis		52.39	0.45	RAF 99			
Clostridium clostridioforme		22.33	0.33	XYL 90			
Leptotrichia buccalis		17.07	0.15				
47331400	**Low discrimination**						**47331400**
					COC	GRA	
Lactobacillus jensenii		89.35	0.44	CEL 95	1	99	
Atopobium parvulum		4.82	0.21	ARA 1	99	99	
Fusobacterium mortiferum		3.63	0.30		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47331402	**Low discrimination**							**47331402**
Lactobacillus jensenii		86.92	0.41	CEL 95				
Bifidobacterium adolescentis		3.66	0.23	ARA 1				
Actinomyces israelii		2.93	0.28					
47331410	**Good identification**							**47331410**
Lactobacillus jensenii		100.0	0.36	CEL 95				
47331500	**Acceptable identification**							**47331500**
					COC	GRA	SPO	
Atopobium parvulum		93.72	0.55	ARA 1	99	99	1	
Clostridium butyricum		1.75	0.18		1	99	99	
Fusobacterium mortiferum		1.66	0.36		1	1	1	
47331502	**Low discrimination**							**47331502**
					COC			
Bifidobacterium adolescentis		67.38	0.58	RAF 92	1			
Atopobium parvulum		22.85	0.42	XYL 87	99			
Lactobacillus acidophilus		2.95	0.25		1			
47331512	**Low discrimination**							**47331512**
					GRA			
Bifidobacterium adolescentis		70.67	0.59	RAF 92	99			
Leptotrichia buccalis		21.91	0.32	XYL 87	1			
Clostridium clostridioforme		6.80	0.39		50			
47331520	**Low discrimination**							**47331520**
					COC	SPO		
Clostridium perfringens		49.04	0.45	ARA 1	1	84		
Atopobium parvulum		35.14	0.42		99	1		
Clostridium paraputrificum		7.72	0.22		1	99		
47331522	**Low discrimination**							**47331522**
					COC	GRA		
Clostridium perfringens		44.37	0.40	ARA 1	1	99		
Atopobium parvulum		11.92	0.29		99	99		
Parabacteroides distasonis		9.77	0.38		1	1		
47331532	**Low discrimination**							**47331532**
Clostridium clostridioforme		55.91	0.39	RAF 94				
Leptotrichia buccalis		20.01	0.15	XYL 91				
Parabacteroides distasonis		12.54	0.33	RHA 88				
47331542	**Low discrimination**							**47331542**
Bifidobacterium adolescentis		83.78	0.42	RAF 92				
Actinomyces israelii		8.53	0.31	XYL 87				
Lactobacillus acidophilus		3.66	0.10	SOR 11				
47331552	**Good identification**							**47331552**
					GRA			
Bifidobacterium adolescentis		95.63	0.44	RAF 92	99			
Leptotrichia buccalis		2.42	-.03	XYL 87	1			
Actinomyces israelii		1.19	0.17	SOR 11	99			
47331612	**Good identification**							**47331612**
Clostridium clostridioforme		100.0	0.44	RAF 94				
47331632	**Good identification**							**47331632**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.44	RAF 94		
47331702	**Low discrimination**					**47331702**
Clostridium clostridioforme		86.24	0.42	RAF 94		
Bifidobacterium adolescentis		4.60	0.23	XYL 91		
Eubacterium contortum		4.32	0.14			
47331712	**Good identification**					**47331712**
					GRA	
Clostridium clostridioforme		98.16	0.53	RAF 94	50	
Bifidobacterium adolescentis		1.40	0.25	XYL 91	99	
Leptotrichia buccalis		0.44	-.03		1	
47331722	**Low discrimination**					**47331722**
Clostridium clostridioforme		52.84	0.42	RAF 94		
Parabacteroides distasonis		41.04	0.46	XYL 91		
Prevotella oralis		2.24	0.12			
47331732	**Acceptable identification**					**47331732**
Clostridium clostridioforme		91.59	0.53	RAF 94		
Parabacteroides distasonis		8.41	0.41	XYL 91		
47332101	**Good identification**					**47332101**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47332103	**Good identification**					**47332103**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47332121	**Good identification**					**47332121**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47332123	**Good identification**					**47332123**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47333000	**Low discrimination**					**47333000**
Lactobacillus jensenii		67.14	0.44	CEL 95		
Actinomyces naeslundii		14.59	0.54	MAN 1		
Actinomyces israelii		6.97	0.39			
47333001	**Very good identification**					**47333001**
Actinomyces naeslundii		97.00	0.57	ESL 12		
Lactobacillus jensenii		3.01	0.10	RAF 92		
47333002	**Low discrimination**					**47333002**
Lactobacillus jensenii		55.14	0.41	CEL 95		
Actinomyces naeslundii		17.98	0.54	MAN 1		
Actinomyces israelii		12.88	0.43			
47333003	**Very good identification**					**47333003**
Actinomyces naeslundii		97.98	0.57	ESL 12		
Lactobacillus jensenii		2.02	0.07	RAF 92		
47333010	**Low discrimination**					**47333010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Lactobacillus jensenii		87.59	0.36	CEL 95	1	
Clostridium tertium		5.33	0.13	MAN 1	99	
Actinomyces israelii		4.08	0.25		1	
47333011	**Low discrimination**					**47333011**
Actinomyces naeslundii		83.60	0.35	ESL 12		
Lactobacillus jensenii		16.40	0.02	RAF 92		
47333013	**Good identification**					**47333013**
Actinomyces naeslundii		100.0	0.35	ESL 12		
47333021	**Very good identification**					**47333021**
Actinomyces naeslundii		100.0	0.57	ESL 12		
47333022	**Low discrimination**					**47333022**
					SPO	
Actinomyces naeslundii		26.65	0.54	ESL 12	1	
Lactobacillus jensenii		20.44	0.31	RAF 92	1	
Clostridium tertium		12.58	0.25		99	
47333023	**Very good identification**					**47333023**
Actinomyces naeslundii		100.0	0.57	ESL 12		
47333031	**Good identification**					**47333031**
Actinomyces naeslundii		100.0	0.35	ESL 12		
47333033	**Good identification**					**47333033**
Actinomyces naeslundii		100.0	0.35	ESL 12		
47333041	**Good identification**					**47333041**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47333042	**Low discrimination**					**47333042**
Actinomyces israelii		53.85	0.35	CEL 93		
Actinomyces naeslundii		34.11	0.41	XYL 92		
Lactobacillus acidophilus		6.05	0.04			
47333043	**Good identification**					**47333043**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47333060	**Low discrimination**					**47333060**
Actinomyces naeslundii		52.88	0.41	ESL 12		
Clostridium perfringens		47.12	0.18	RAF 92		
47333061	**Good identification**					**47333061**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47333062	**Low discrimination**					**47333062**
					SPO	
Actinomyces naeslundii		52.20	0.41	ESL 12	1	
Clostridium perfringens		21.89	0.13	RAF 92	84	
Clostridium barati		11.39	0.03		99	
47333063	**Good identification**					**47333063**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	ESL 12			
47333100	**Low discrimination**						**47333100**
					COC	SPO	
Atopobium parvulum		69.02	0.55	MAN 1	99	1	
Clostridium tertium		14.58	0.43		1	99	
Actinomyces naeslundii		7.99	0.62		1	1	
47333101	**Good identification**						**47333101**
					COC	SPO	
Actinomyces naeslundii		91.43	0.65	ESL 12	1	1	
Atopobium parvulum		5.32	0.21	RAF 92	99	1	
Clostridium chauvoei		1.72	0.12		1	99	
47333102	**Low discrimination**						**47333102**
					COC	SPO	
Clostridium tertium		48.85	0.51	CEL 99	1	99	
Atopobium parvulum		13.60	0.42		99	1	
Lactobacillus acidophilus		9.14	0.38		1	1	
47333103	**Good identification**						**47333103**
					COC	SPO	
Actinomyces naeslundii		93.65	0.65	ESL 12	1	1	
Clostridium tertium		3.45	0.17	RAF 92	1	99	
Atopobium parvulum		0.96	0.08		99	1	
47333110	**Acceptable identification**						**47333110**
					COC	SPO	
Clostridium tertium		91.72	0.47	CEL 99	1	99	
Actinomyces israelii		2.84	0.36		1	1	
Atopobium parvulum		2.47	0.21		99	1	
47333111	**Low discrimination**						**47333111**
					SPO		
Actinomyces naeslundii		70.67	0.43	ESL 12	1		
Clostridium tertium		29.33	0.13	RAF 92	99		
47333112	**Low discrimination**						**47333112**
					GRA	SPO	
Clostridium tertium		77.41	0.55	CEL 99	99	99	
Leptotrichia buccalis		13.55	0.32		1	1	
Bifidobacterium adolescentis		5.34	0.44		99	1	
47333120	**Low discrimination**						**47333120**
Clostridium tertium		38.63	0.51	CEL 99			
Clostridium ramosum		22.54	0.45				
Clostridium perfringens		15.01	0.45				
47333121	**Good identification**						**47333121**
					SPO		
Actinomyces naeslundii		91.81	0.65	ESL 12	1		
Clostridium tertium		3.39	0.17	RAF 92	99		
Clostridium ramosum		1.98	0.11		92		
47333122	**Low discrimination**						**47333122**
					SPO		
Clostridium tertium		48.69	0.59	CEL 99	99		
Clostridium ramosum		37.88	0.56		92		
Lactobacillus acidophilus		3.04	0.38		1		
47333123	**Low discrimination**						**47333123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		82.63	0.65	ESL 12	1		
Clostridium tertium		9.14	0.25	RAF 92	99		
Clostridium ramosum		7.11	0.21		92		
47333130	**Good identification**						**47333130**
					SPO		
Clostridium tertium		98.29	0.55	CEL 99	99		
Actinomyces naeslundii		0.53	0.40		1		
Bifidobacterium breve		0.42	0.24		1		
47333132	**Very good identification**						**47333132**
					GRA	SPO	
Clostridium tertium		97.24	0.63	CEL 99	99	99	
Bifidobacterium breve		1.40	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
47333140	**Low discrimination**						**47333140**
					COC		
Actinomyces israelii		45.07	0.43	CEL 93	1		
Actinomyces naeslundii		30.47	0.49	XYL 92	1		
Atopobium parvulum		15.07	0.21		99		
47333141	**Good identification**						**47333141**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47333142	**Low discrimination**						**47333142**
Actinomyces israelii		35.75	0.46	CEL 93			
Bifidobacterium breve		21.61	0.37	XYL 92			
Actinomyces naeslundii		16.11	0.49				
47333143	**Good identification**						**47333143**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47333152	**Low discrimination**						**47333152**
					SPO		
Bifidobacterium breve		47.05	0.37	RAF 95	1		
Clostridium tertium		19.55	0.21	CEL 99	99		
Bifidobacterium adolescentis		16.50	0.28		1		
47333161	**Good identification**						**47333161**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47333162	**Low discrimination**						**47333162**
Bifidobacterium breve		32.72	0.41	RAF 95			
Clostridium perfringens		14.83	0.27	CEL 99			
Actinomyces naeslundii		13.73	0.49				
47333163	**Good identification**						**47333163**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47333172	**Low discrimination**						**47333172**
					SPO		
Bifidobacterium breve		58.04	0.41	RAF 95	1		
Clostridium tertium		40.68	0.29	CEL 99	99		
Actinomyces naeslundii		1.28	0.27		1		
47333320	**Good identification**						**47333320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		97.34	0.47	CEL 99			
Clostridium tertium		1.27	0.17				
Clostridium perfringens		0.49	0.11				
47333322	**Very good identification**						**47333322**
					SPO		
Clostridium ramosum		97.94	0.58	CEL 99	92		
Clostridium tertium		0.96	0.25		99		
Lactobacillus acidophilus		0.45	0.18		1		
47333400	**Low discrimination**						**47333400**
					SPO		
Actinomyces israelii		85.35	0.38	CEL 93	1		
Lactobacillus jensenii		10.15	0.10	XYL 92	1		
Clostridium innocuum		4.49	0.01		99		
47333402	**Low discrimination**						**47333402**
Actinomyces israelii		78.84	0.41	CEL 93			
Bifidobacterium adolescentis		14.22	0.22	XYL 92			
Lactobacillus jensenii		4.17	0.07				
47333500	**Low discrimination**						**47333500**
					COC		
Actinomyces israelii		76.27	0.48	CEL 93	1		
Atopobium parvulum		12.12	0.21	XYL 92	99		
Bifidobacterium adolescentis		5.16	0.22		1		
47333502	**Low discrimination**						**47333502**
					SPO		
Bifidobacterium adolescentis		79.96	0.56	RAF 92	1		
Actinomyces israelii		17.91	0.51	XYL 87	1		
Clostridium tertium		1.20	0.17		99		
47333512	**Good identification**						**47333512**
					SPO		
Bifidobacterium adolescentis		93.94	0.58	RAF 92	1		
Actinomyces israelii		2.57	0.37	XYL 87	1		
Clostridium tertium		2.06	0.21		99		
47333540	**Good identification**						**47333540**
Actinomyces israelii		100.0	0.41	CEL 93			
47333542	**Low discrimination**						**47333542**
Bifidobacterium adolescentis		58.66	0.41	RAF 92			
Actinomyces israelii		41.34	0.44	XYL 87			
47333552	**Acceptable identification**						**47333552**
Bifidobacterium adolescentis		92.07	0.42	RAF 92			
Actinomyces israelii		7.93	0.30	XYL 87			
47334000	**Low discrimination**						**47334000**
					COC	GRA	
Fusobacterium mortiferum		64.63	0.42	MLT 11	1	1	
Atopobium parvulum		32.13	0.25	XYL 7	99	99	
Lactobacillus jensenii		3.24	0.10		1	99	
47334100	**Very good identification**						**47334100**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		95.36	0.60	GAL 99	99	99	
Fusobacterium mortiferum		4.52	0.48		1	1	
Actinomyces israelii		0.08	0.21		1	99	
47334102	**Acceptable identification**						**47334102**
					COC	GRA	
Atopobium parvulum		93.51	0.47	GAL 99	99	99	
Fusobacterium mortiferum		2.79	0.32		1	1	
Bifidobacterium adolescentis		1.50	0.23		1	99	
47334120	**Good identification**						**47334120**
					COC	GRA	
Atopobium parvulum		98.98	0.47	GAL 99	99	99	
Clostridium perfringens		0.75	0.11		1	99	
Fusobacterium mortiferum		0.27	0.14		1	1	
47334312	**Good identification**						**47334312**
Clostridium clostridioforme		100.0	0.42	RAF 94			
47334332	**Acceptable identification**						**47334332**
Clostridium clostridioforme		94.05	0.42	RAF 94			
Parabacteroides distasonis		5.95	0.27	ARA 94			
47334412	**Good identification**						**47334412**
Clostridium clostridioforme		100.0	0.39	RAF 94			
47334432	**Good identification**						**47334432**
Clostridium clostridioforme		100.0	0.39	RAF 94			
47334502	**Low discrimination**						**47334502**
					COC		
Clostridium clostridioforme		51.41	0.36	RAF 94	1		
Bifidobacterium adolescentis		39.92	0.38	RHA 88	1		
Atopobium parvulum		3.76	0.13		99		
47334512	**Low discrimination**						**47334512**
Clostridium clostridioforme		82.76	0.48	RAF 94			
Bifidobacterium adolescentis		17.24	0.39	RHA 88			
47334522	**Acceptable identification**						**47334522**
					COC	GRA	
Clostridium clostridioforme		93.98	0.36	RAF 94	1	50	
Parabacteroides distasonis		4.81	0.20	RHA 88	1	1	
Atopobium parvulum		1.21	0.00		99	99	
47334532	**Good identification**						**47334532**
Clostridium clostridioforme		99.40	0.48	RAF 94			
Parabacteroides distasonis		0.60	0.15	RHA 88			
47334602	**Good identification**						**47334602**
Clostridium clostridioforme		100.0	0.42	RAF 94			
47334612	**Good identification**						**47334612**
Clostridium clostridioforme		100.0	0.53	RAF 94			
47334622	**Good identification**						**47334622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.42	RAF 94			
47334632	**Good identification**						**47334632**
Clostridium clostridioforme		100.0	0.53	RAF 94			
47334702	**Good identification**						**47334702**
Clostridium clostridioforme		98.54	0.51	RAF 94			
Eubacterium contortum		1.47	0.14				
47334712	**Very good identification**						**47334712**
Clostridium clostridioforme		100.0	0.62	RAF 94			
47334722	**Good identification**						**47334722**
					GRA		
Clostridium clostridioforme		97.70	0.51	RAF 94	50		
Parabacteroides distasonis		2.05	0.28		1		
Eubacterium contortum		0.26	0.01		99		
47334732	**Very good identification**						**47334732**
Clostridium clostridioforme		99.75	0.62	RAF 94			
Parabacteroides distasonis		0.25	0.23				
47335000	**Low discrimination**						**47335000**
					COC	GRA	
Atopobium parvulum		86.61	0.60	LAC 99	99	99	
Lactobacillus jensenii		8.73	0.44		1	99	
Fusobacterium mortiferum		2.64	0.45		1	1	
47335002	**Low discrimination**						**47335002**
					COC		
Atopobium parvulum		61.61	0.47	LAC 99	99		
Lactobacillus jensenii		23.46	0.41		1		
Actinomyces israelii		11.12	0.48		1		
47335010	**Low discrimination**						**47335010**
					COC		
Lactobacillus jensenii		71.21	0.36	CEL 95	1		
Atopobium parvulum		21.41	0.25	XYL 1	99		
Actinomyces israelii		6.72	0.31		1		
47335020	**Low discrimination**						**47335020**
					COC		
Atopobium parvulum		79.57	0.47	LAC 99	99		
Lactobacillus jensenii		11.36	0.34		1		
Clostridium perfringens		8.15	0.30		1		
47335042	**Low discrimination**						**47335042**
					COC		
Actinomyces israelii		81.50	0.41	CEL 93	1		
Atopobium parvulum		11.73	0.13		99		
Lactobacillus jensenii		4.47	0.07		1		
47335100	**Excellent identification**						**47335100**
					COC	GRA	
Atopobium parvulum		99.76	0.94		99	99	
Actinomyces israelii		0.09	0.55		1	99	
Fusobacterium mortiferum		0.07	0.51		1	1	
47335101	**Very good identification**						**47335101**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		99.58	0.60	URE 1	99	1		
Clostridium chauvoei		0.18	0.12		1	99		
Actinomyces israelii		0.09	0.21		1	1		
47335102	**Very good identification**							**47335102**
					COC			
Atopobium parvulum		96.71	0.81		99			
Bifidobacterium adolescentis		1.55	0.58		1			
Bifidobacterium longum subsp. infantis		0.71	0.38		1			
47335103	**Good identification**							**47335103**
					COC			
Atopobium parvulum		96.60	0.47	URE 1	99			
Bifidobacterium adolescentis		1.55	0.23		1			
Bifidobacterium longum subsp. infantis		0.71	0.04		1			
47335110	**Very good identification**							**47335110**
					COC			
Atopobium parvulum		97.71	0.60	MLZ 1	99			
Actinomyces israelii		1.24	0.41		1			
Bifidobacterium adolescentis		0.34	0.25		1			
47335112	**Low discrimination**							**47335112**
					COC	GRA		
Bifidobacterium adolescentis		50.58	0.59	RAF 92	1	99		
Atopobium parvulum		26.06	0.47	ARA 87	99	99		
Leptotrichia buccalis		15.68	0.32		1	1		
47335120	**Very good identification**							**47335120**
					COC			
Atopobium parvulum		98.76	0.81		99			
Clostridium perfringens		0.75	0.45		1			
Bifidobacterium longum subsp. infantis		0.13	0.25		1			
47335121	**Good identification**							**47335121**
					COC	SPO		
Atopobium parvulum		98.65	0.47	URE 1	99	1		
Clostridium perfringens		0.75	0.11		1	84		
Clostridium chauvoei		0.33	0.04		1	99		
47335122	**Low discrimination**							**47335122**
					COC	GRA		
Atopobium parvulum		83.32	0.68		99	99		
Bifidobacterium longum subsp. infantis		6.19	0.43		1	99		
Parabacteroides distasonis		5.45	0.58		1	1		
47335130	**Acceptable identification**							**47335130**
					COC	SPO		
Atopobium parvulum		93.78	0.47	MLZ 1	99	1		
Clostridium tertium		4.50	0.24		1	99		
Clostridium perfringens		0.71	0.11		1	84		
47335132	**Low discrimination**							**47335132**
					COC	GRA	SPO	
Parabacteroides distasonis		52.61	0.53	RAF 99	1	1	1	
Atopobium parvulum		15.08	0.34		99	99	1	
Clostridium tertium		12.30	0.32		1	99	99	
47335140	**Very good identification**							**47335140**
					COC	SPO		
Atopobium parvulum		96.34	0.60	SOR 1	99	1		
Actinomyces israelii		3.18	0.48		1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.30	0.10		1	99	
47335142	**Low discrimination**						**47335142**
					COC		
Atopobium parvulum		66.58	0.47	SOR 1	99		
Actinomyces israelii		18.69	0.51		1		
Bifidobacterium adolescentis		13.07	0.42		1		
47335152	**Low discrimination**						**47335152**
					COC		
Bifidobacterium adolescentis		80.50	0.44	RAF 92	1		
Actinomyces israelii		14.08	0.37	ARA 87	1		
Atopobium parvulum		3.39	0.13	SOR 11	99		
47335160	**Low discrimination**						**47335160**
					COC		
Atopobium parvulum		85.91	0.47	SOR 1	99		
Clostridium perfringens		12.30	0.33		1		
Actinomyces israelii		1.03	0.28		1		
47335212	**Good identification**						**47335212**
Clostridium clostridioforme		99.24	0.41	RAF 94			
Actinomyces israelii		0.76	0.14	ARA 94			
47335232	**Acceptable identification**						**47335232**
Clostridium clostridioforme		93.40	0.41	RAF 94			
Parabacteroides distasonis		6.60	0.27	ARA 94			
47335300	**Low discrimination**						**47335300**
					COC		
Eubacterium contortum		64.40	0.63	TRE 1	1		
Atopobium parvulum		35.30	0.60		99		
Actinomyces israelii		0.23	0.36		1		
47335302	**Low discrimination**						**47335302**
					COC		
Eubacterium contortum		57.98	0.50	TRE 1	1		
Atopobium parvulum		31.78	0.47		99		
Clostridium clostridioforme		6.17	0.39		1		
47335312	**Acceptable identification**						**47335312**
					GRA		
Clostridium clostridioforme		91.61	0.50	RAF 94	50		
Bifidobacterium adolescentis		2.03	0.25	ARA 94	99		
Parabacteroides distasonis		1.93	0.27		1		
47335320	**Low discrimination**						**47335320**
					COC	GRA	
Eubacterium contortum		61.23	0.50	TRE 1	1	99	
Atopobium parvulum		33.57	0.47		99	99	
Bacteroides vulgatus		3.27	0.21		1	1	
47335322	**Low discrimination**						**47335322**
					GRA		
Parabacteroides distasonis		79.50	0.66	RAF 99	1		
Eubacterium contortum		7.46	0.37		99		
Clostridium clostridioforme		4.50	0.39		50		
47335332	**Low discrimination**						**47335332**
					GRA		
Parabacteroides distasonis		67.50	0.61	RAF 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		32.32	0.50		50		
Eubacterium contortum		0.12	0.03		99		
47335400	**Low discrimination**						**47335400**
					COC	SPO	
Actinomyces israelii		41.02	0.43	CEL 93	1	1	
Clostridium butyricum		32.00	0.20		1	99	
Atopobium parvulum		23.85	0.25		99	1	
47335402	**Low discrimination**						**47335402**
Actinomyces israelii		34.93	0.46	CEL 93			
Clostridium clostridioforme		29.31	0.36				
Bifidobacterium adolescentis		25.40	0.38				
47335412	**Low discrimination**						**47335412**
Clostridium clostridioforme		78.64	0.47	RAF 94			
Bifidobacterium adolescentis		18.28	0.39	RHA 88			
Actinomyces israelii		3.07	0.32				
47335422	**Low discrimination**						**47335422**
					GRA		
Clostridium clostridioforme		85.58	0.36	RAF 94	50		
Actinomyces israelii		6.51	0.26	RHA 88	99		
Parabacteroides distasonis		4.89	0.20		1		
47335432	**Good identification**						**47335432**
Clostridium clostridioforme		99.33	0.47	RAF 94			
Parabacteroides distasonis		0.67	0.15	RHA 88			
47335440	**Low discrimination**						**47335440**
					SPO		
Actinomyces israelii		85.15	0.36	CEL 93	1		
Clostridium butyricum		14.85	0.02		99		
47335442	**Low discrimination**						**47335442**
					SPO		
Actinomyces israelii		78.35	0.39	CEL 93	1		
Bifidobacterium adolescentis		18.11	0.22		1		
Clostridium butyricum		3.54	-.05		99		
47335500	**Low discrimination**						**47335500**
					COC	SPO	
Atopobium parvulum		69.87	0.60	ARA 1	99	1	
Clostridium butyricum		22.72	0.44		1	99	
Actinomyces israelii		4.85	0.54		1	1	
47335502	**Low discrimination**						**47335502**
					COC	SPO	
Bifidobacterium adolescentis		79.39	0.72	RAF 92	1	1	
Atopobium parvulum		7.47	0.47		99	1	
Clostridium butyricum		5.35	0.37		1	99	
47335503	**Low discrimination**						**47335503**
					COC	SPO	
Bifidobacterium adolescentis		79.61	0.38	RAF 92	1	1	
Atopobium parvulum		7.49	0.13	URE 1	99	1	
Clostridium butyricum		5.37	0.03		1	99	
47335510	**Low discrimination**						**47335510**
					SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium adolescentis		35.00	0.39	RAF 92	1			
Clostridium butyricum		25.87	0.22	BGL 99	99			
Actinomyces israelii		15.69	0.39		1			
47335512	**Low discrimination**							**47335512**
Bifidobacterium adolescentis		86.29	0.73	RAF 92				
Clostridium clostridioforme		12.55	0.56					
Actinomyces israelii		0.59	0.42					
47335513	**Low discrimination**							**47335513**
Bifidobacterium adolescentis		87.30	0.39	RAF 92				
Clostridium clostridioforme		12.70	0.22	URE 1				
47335520	**Low discrimination**							**47335520**
					COC	GRA	SPO	
Atopobium parvulum		71.92	0.47	ARA 1	99	99	1	
Clostridium butyricum		11.52	0.26		1	99	99	
Bacteroides vulgatus		10.36	0.24		1	1	1	
47335522	**Low discrimination**							**47335522**
					COC	GRA		
Parabacteroides distasonis		45.17	0.54	RAF 99	1	1		
Clostridium clostridioforme		26.73	0.45		1	50		
Atopobium parvulum		11.38	0.34		99	99		
47335532	**Low discrimination**							**47335532**
					GRA			
Clostridium clostridioforme		76.07	0.56	RAF 94	50			
Parabacteroides distasonis		15.19	0.49	RHA 88	1			
Bifidobacterium adolescentis		5.28	0.39		99			
47335542	**Low discrimination**							**47335542**
					SPO			
Bifidobacterium adolescentis		81.09	0.56	RAF 92	1			
Actinomyces israelii		14.18	0.50	SOR 11	1			
Clostridium butyricum		3.85	0.19		99			
47335552	**Very good identification**							**47335552**
Bifidobacterium adolescentis		96.59	0.58	RAF 92				
Actinomyces israelii		2.07	0.35	SOR 11				
Clostridium clostridioforme		1.15	0.22					
47335602	**Good identification**							**47335602**
Clostridium clostridioforme		97.20	0.50	RAF 94				
Eubacterium contortum		1.61	0.14					
Actinomyces israelii		1.19	0.27					
47335612	**Very good identification**							**47335612**
Clostridium clostridioforme		100.0	0.62	RAF 94				
47335622	**Good identification**							**47335622**
					GRA			
Clostridium clostridioforme		97.44	0.50	RAF 94	50			
Parabacteroides distasonis		2.28	0.28		1			
Eubacterium contortum		0.29	0.01		99			
47335632	**Very good identification**							**47335632**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		99.72	0.62	RAF 94			
Parabacteroides distasonis		0.28	0.23				
47335700	**Very good identification**						**47335700**
					COC	SPO	
Eubacterium contortum		97.83	0.61	TRE 1	1	1	
Atopobium parvulum		0.66	0.25		99	1	
Clostridium butyricum		0.66	0.18		1	99	
47335702	**Low discrimination**						**47335702**
Clostridium clostridioforme		64.17	0.59	RAF 94			
Eubacterium contortum		31.50	0.48				
Bifidobacterium adolescentis		2.27	0.38				
47335710	**Low discrimination**						**47335710**
Clostridium clostridioforme		60.11	0.36	RAF 94			
Eubacterium contortum		36.70	0.27	BGL 99			
Actinomyces israelii		1.90	0.20				
47335712	**Very good identification**						**47335712**
Clostridium clostridioforme		98.83	0.71	RAF 94			
Bifidobacterium adolescentis		0.94	0.39				
Eubacterium contortum		0.11	0.14				
47335713	**Good identification**						**47335713**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47335720	**Low discrimination**						**47335720**
					GRA		
Eubacterium contortum		52.24	0.48	TRE 1	99		
Bacteroides vulgatus		45.35	0.40		1		
Clostridium clostridioforme		1.08	0.25		50		
47335722	**Low discrimination**						**47335722**
					GRA		
Clostridium clostridioforme		55.70	0.59	RAF 94	50		
Parabacteroides distasonis		38.50	0.62		1		
Eubacterium contortum		4.82	0.36		99		
47335730	**Low discrimination**						**47335730**
					GRA		
Clostridium clostridioforme		77.94	0.36	RAF 94	50		
Eubacterium contortum		8.40	0.14	BGL 99	99		
Bacteroides vulgatus		7.29	0.06		1		
47335732	**Good identification**						**47335732**
Clostridium clostridioforme		92.33	0.71	RAF 94			
Parabacteroides distasonis		7.54	0.58				
Bacteroides thetaiotaomicron		0.06	0.06				
47335733	**Acceptable identification**						**47335733**
Clostridium clostridioforme		92.45	0.36	RAF 94			
Parabacteroides distasonis		7.55	0.23	URE 1			
47335752	**Low discrimination**						**47335752**
Clostridium clostridioforme		88.16	0.36	RAF 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		10.22	0.24	SOR 1			
Actinomyces israelii		1.63	0.16				
47335772	**Acceptable identification**						**47335772**
Clostridium clostridioforme		92.45	0.36	RAF 94			
Parabacteroides distasonis		7.55	0.23	SOR 1			
47336102	**Low discrimination**						**47336102**
					SPO		
Actinomyces israelii		45.14	0.37	CEL 93	1		
Clostridium tertium		29.48	0.19	GAL 99	99		
Bifidobacterium adolescentis		14.34	0.22		1		
47336502	**Low discrimination**						**47336502**
Bifidobacterium adolescentis		72.21	0.36	RAF 92			
Actinomyces israelii		27.79	0.35	GAL 99			
47336512	**Low discrimination**						**47336512**
Bifidobacterium adolescentis		79.53	0.38	RAF 92			
Bifidobacterium dentium		12.02	0.06	GAL 99			
Clostridium clostridioforme		4.71	0.13				
47337000	**Good identification**						**47337000**
					COC	SPO	
Actinomyces israelii		90.24	0.58	CEL 93	1	1	
Atopobium parvulum		7.58	0.25		99	1	
Clostridium tertium		1.19	0.11		1	99	
47337002	**Good identification**						**47337002**
					SPO		
Actinomyces israelii		94.98	0.61	CEL 93	1		
Clostridium tertium		2.51	0.19		99		
Bifidobacterium adolescentis		1.22	0.22		1		
47337010	**Low discrimination**						**47337010**
					SPO		
Actinomyces israelii		85.05	0.44	CEL 93	1		
Clostridium tertium		13.35	0.15	MLZ 13	99		
Lactobacillus jensenii		1.61	0.02		1		
47337012	**Low discrimination**						**47337012**
					SPO		
Actinomyces israelii		70.48	0.47	CEL 93	1		
Clostridium tertium		22.12	0.24	MLZ 13	99		
Bifidobacterium adolescentis		7.40	0.23		1		
47337020	**Low discrimination**						**47337020**
					COC	SPO	
Actinomyces israelii		53.99	0.37	CEL 93	1	1	
Clostridium tertium		33.48	0.19	NAG 6	1	99	
Atopobium parvulum		12.53	0.13		99	1	
47337040	**Good identification**						**47337040**
Actinomyces israelii		100.0	0.51	CEL 93			
47337042	**Good identification**						**47337042**
Actinomyces israelii		100.0	0.54	CEL 93			
47337050	**Good identification**						**47337050**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.36	CEL 93			
47337052	**Good identification**						**47337052**
Actinomyces israelii		96.77	0.40	CEL 93			
Bifidobacterium adolescentis		3.23	0.08	MLZ 13			
47337100	**Low discrimination**						**47337100**
					COC	SPO	
Atopobium parvulum		60.75	0.60	MAN 1	99	1	
Actinomyces israelii		29.23	0.68		1	1	
Clostridium tertium		9.55	0.45		1	99	
47337102	**Low discrimination**						**47337102**
					SPO		
Actinomyces israelii		45.03	0.71	CEL 93	1		
Clostridium tertium		29.40	0.53		99		
Bifidobacterium adolescentis		14.30	0.56		1		
47337103	**Low discrimination**						**47337103**
					SPO		
Actinomyces israelii		42.65	0.37	CEL 93	1		
Clostridium tertium		27.85	0.19	URE 1	99		
Bifidobacterium adolescentis		13.55	0.22		1		
47337110	**Low discrimination**						**47337110**
					COC	SPO	
Clostridium tertium		76.70	0.50	CEL 99	1	99	
Actinomyces israelii		19.75	0.54		1	1	
Atopobium parvulum		2.77	0.25		99	1	
47337112	**Low discrimination**						**47337112**
					SPO		
Clostridium tertium		68.03	0.58	CEL 99	99		
Bifidobacterium adolescentis		22.75	0.58		1		
Actinomyces israelii		8.76	0.57		1		
47337120	**Low discrimination**						**47337120**
					COC	SPO	
Clostridium tertium		69.02	0.53	CEL 99	1	99	
Atopobium parvulum		25.84	0.47		99	1	
Actinomyces israelii		4.50	0.48		1	1	
47337122	**Good identification**						**47337122**
					COC	SPO	
Clostridium tertium		93.74	0.62	CEL 99	1	99	
Actinomyces israelii		3.05	0.51		1	1	
Atopobium parvulum		2.06	0.34		99	1	
47337130	**Very good identification**						**47337130**
					COC	SPO	
Clostridium tertium		99.19	0.58	CEL 99	1	99	
Actinomyces israelii		0.54	0.33		1	1	
Atopobium parvulum		0.21	0.13		99	1	
47337132	**Very good identification**						**47337132**
					SPO		
Clostridium tertium		99.36	0.66	CEL 99	99		
Actinomyces israelii		0.27	0.36		1		
Bifidobacterium dentium		0.20	0.10		1		
47337140	**Good identification**						**47337140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		93.96	0.61	CEL 93	1	1	
Atopobium parvulum		5.07	0.25		99	1	
Clostridium tertium		0.80	0.11		1	99	
47337142	**Low discrimination**						**47337142**
					SPO		
Actinomyces israelii		88.95	0.64	CEL 93	1		
Bifidobacterium adolescentis		8.98	0.41		1		
Clostridium tertium		1.51	0.19		99		
47337150	**Low discrimination**						**47337150**
					COC	SPO	
Actinomyces israelii		88.28	0.47	CEL 93	1	1	
Clostridium tertium		8.91	0.15	MLZ 13	1	99	
Blautia producta		1.71	-.03		99	1	
47337152	**Low discrimination**						**47337152**
					SPO		
Actinomyces israelii		49.32	0.50	CEL 93	1		
Bifidobacterium adolescentis		40.73	0.42	MLZ 13	1		
Clostridium tertium		9.95	0.24		99		
47337160	**Low discrimination**						**47337160**
					COC	SPO	
Actinomyces israelii		64.61	0.40	CEL 93	1	1	
Clostridium tertium		25.75	0.19	NAG 6	1	99	
Atopobium parvulum		9.64	0.13		99	1	
47337162	**Low discrimination**						**47337162**
					COC	SPO	
Actinomyces israelii		55.11	0.43	CEL 93	1	1	
Clostridium tertium		43.93	0.27	NAG 6	1	99	
Atopobium parvulum		0.97	0.00		99	1	
47337200	**Good identification**						**47337200**
Actinomyces israelii		100.0	0.38	CEL 93			
47337202	**Good identification**						**47337202**
Actinomyces israelii		100.0	0.41	CEL 93			
47337300	**Low discrimination**						**47337300**
					COC		
Actinomyces israelii		54.60	0.49	CEL 93	1		
Eubacterium contortum		27.78	0.29	RHA 7	1		
Atopobium parvulum		15.23	0.25		99		
47337302	**Low discrimination**						**47337302**
					SPO		
Actinomyces israelii		81.35	0.52	CEL 93	1		
Clostridium tertium		7.13	0.19	RHA 7	99		
Eubacterium contortum		4.87	0.16		1		
47337340	**Good identification**						**47337340**
Actinomyces israelii		100.0	0.42	CEL 93			
47337342	**Good identification**						**47337342**
Actinomyces israelii		100.0	0.45	CEL 93			
47337400	**Very good identification**						**47337400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		98.49	0.56	CEL 93	1	
Clostridium butyricum		1.51	0.04		99	
47337402	**Good identification**					**47337402**
					SPO	
Actinomyces israelii		90.18	0.59	CEL 93	1	
Bifidobacterium adolescentis		9.47	0.36		1	
Clostridium butyricum		0.36	-.03		99	
47337410	**Good identification**					**47337410**
Actinomyces israelii		100.0	0.42	CEL 93		
47337412	**Low discrimination**					**47337412**
Actinomyces israelii		49.16	0.45	CEL 93		
Bifidobacterium adolescentis		42.21	0.38	MLZ 13		
Bifidobacterium dentium		6.38	0.06			
47337420	**Good identification**					**47337420**
Actinomyces israelii		100.0	0.36	CEL 93		
47337422	**Good identification**					**47337422**
Actinomyces israelii		100.0	0.39	CEL 93		
47337440	**Good identification**					**47337440**
Actinomyces israelii		100.0	0.49	CEL 93		
47337442	**Good identification**					**47337442**
Actinomyces israelii		96.77	0.52	CEL 93		
Bifidobacterium adolescentis		3.23	0.21			
47337450	**Acceptable identification**					**47337450**
					COC	
Actinomyces israelii		91.35	0.35	CEL 93	1	
Blautia producta		8.65	-.03	MLZ 13	99	
47337452	**Low discrimination**					**47337452**
Actinomyces israelii		78.56	0.38	CEL 93		
Bifidobacterium adolescentis		21.44	0.22	MLZ 13		
47337500	**Low discrimination**					**47337500**
					SPO	
Actinomyces israelii		86.15	0.66	CEL 93	1	
Clostridium butyricum		7.91	0.28		99	
Bifidobacterium adolescentis		3.39	0.36		1	
47337502	**Low discrimination**					**47337502**
					SPO	
Bifidobacterium adolescentis		71.40	0.70	RAF 92	1	
Actinomyces israelii		27.48	0.69		1	
Clostridium butyricum		0.65	0.21		99	
47337503	**Low discrimination**					**47337503**
Bifidobacterium adolescentis		72.21	0.36	RAF 92		
Actinomyces israelii		27.79	0.35	URE 1		
47337510	**Low discrimination**					**47337510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		52.97	0.52	CEL 93			
Bifidobacterium dentium		25.24	0.23	MLZ 13			
Bifidobacterium adolescentis		17.05	0.38				
47337512	**Low discrimination**						**47337512**
Bifidobacterium adolescentis		82.99	0.72	RAF 92			
Bifidobacterium dentium		12.55	0.40				
Actinomyces israelii		3.91	0.55				
47337513	**Low discrimination**						**47337513**
Bifidobacterium adolescentis		83.46	0.38	RAF 92			
Bifidobacterium dentium		12.62	0.06	URE 1			
Actinomyces israelii		3.93	0.21				
47337520	**Low discrimination**						**47337520**
					SPO		
Actinomyces israelii		71.15	0.46	CEL 93	1		
Clostridium tertium		13.48	0.19	NAG 6	99		
Clostridium butyricum		8.90	0.10		99		
47337522	**Low discrimination**						**47337522**
					SPO		
Actinomyces israelii		49.54	0.49	CEL 93	1		
Bifidobacterium adolescentis		20.37	0.36	NAG 6	1		
Clostridium tertium		18.77	0.27		99		
47337532	**Acceptable identification**						**47337532**
					SPO		
Bifidobacterium dentium		90.32	0.44	RAF 95	1		
Clostridium tertium		4.55	0.32	CEL 99	99		
Bifidobacterium adolescentis		3.40	0.38		1		
47337540	**Low discrimination**						**47337540**
					COC	SPO	
Actinomyces israelii		86.97	0.59	CEL 93	1	1	
Blautia producta		10.16	0.24		99	1	
Clostridium butyricum		1.79	0.10		1	99	
47337542	**Low discrimination**						**47337542**
					COC		
Actinomyces israelii		54.36	0.62	CEL 93	1		
Bifidobacterium adolescentis		44.88	0.55		1		
Blautia producta		0.47	0.07		99		
47337552	**Low discrimination**						**47337552**
					COC		
Bifidobacterium adolescentis		84.31	0.56	RAF 92	1		
Actinomyces israelii		12.48	0.48	SOR 11	1		
Blautia producta		2.17	0.15		99		
47337560	**Low discrimination**						**47337560**
					COC		
Actinomyces israelii		83.11	0.39	CEL 93	1		
Blautia producta		16.89	0.07	NAG 6	99		
47337562	**Low discrimination**						**47337562**
Actinomyces israelii		88.44	0.42	CEL 93			
Bifidobacterium adolescentis		11.56	0.21	NAG 6			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47337600	**Good identification**					**47337600**
Actinomyces israelii		100.0	0.37	CEL 93		
47337602	**Low discrimination**					**47337602**
Actinomyces israelii		87.28	0.40	CEL 93		
Clostridium clostridioforme		12.72	0.16	RHA 7		
47337700	**Low discrimination**					**47337700**
					SPO	
Actinomyces israelii		64.66	0.47	CEL 93	1	
Eubacterium contortum		32.90	0.27	RHA 7	1	
Clostridium butyricum		2.44	0.02		99	
47337702	**Low discrimination**					**47337702**
Actinomyces israelii		64.92	0.50	CEL 93		
Bifidobacterium adolescentis		22.64	0.36	RHA 7		
Clostridium clostridioforme		7.92	0.25			
47337712	**Low discrimination**					**47337712**
Clostridium clostridioforme		46.47	0.36	RAF 94		
Bifidobacterium adolescentis		35.64	0.38	MAN 1		
Actinomyces israelii		12.50	0.36			
47337732	**Low discrimination**					**47337732**
					GRA	
Clostridium clostridioforme		75.79	0.36	RAF 94	50	
Bifidobacterium dentium		15.62	0.10	MAN 1	99	
Parabacteroides distasonis		6.19	0.23		1	
47337742	**Low discrimination**					**47337742**
					COC	
Actinomyces israelii		81.58	0.43	CEL 93	1	
Blautia producta		9.38	0.07	RHA 7	99	
Bifidobacterium adolescentis		9.04	0.21		1	
47340022	**Acceptable identification**					**47340022**
					SPO	
Clostridium barati		93.18	0.36	FRU 99	99	
Lactobacillus catenaformis		4.43	0.18	GAL 95	1	
Lactobacillus jensenii		1.48	0.10		1	
47340102	**Low discrimination**					**47340102**
					SPO	
Clostridium barati		63.02	0.36	FRU 99	99	
Lactobacillus catenaformis		17.98	0.32	GAL 95	1	
Collinsella aerofaciens		14.54	0.15		1	
47340122	**Low discrimination**					**47340122**
					SPO	
Clostridium barati		81.11	0.44	FRU 99	99	
Clostridium paraputrificum		13.47	0.25	GAL 95	99	
Lactobacillus catenaformis		1.93	0.21		1	
47341000	**Low discrimination**					**47341000**
					SPO	
Lactobacillus catenaformis		57.68	0.63	FRU 99	1	
Lactobacillus jensenii		28.96	0.58		1	
Clostridium barati		6.47	0.41		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47341002	**Low discrimination**					**47341002**
					SPO	
Lactobacillus catenaformis		58.45	0.63	FRU 99	1	
Clostridium barati		19.66	0.50		99	
Lactobacillus jensenii		19.56	0.55		1	
47341010	**Acceptable identification**					**47341010**
					SPO	
Lactobacillus jensenii		93.13	0.50	FRU 99	1	
Lactobacillus catenaformis		5.62	0.29		1	
Clostridium barati		0.63	0.07		99	
47341012	**Low discrimination**					**47341012**
					SPO	
Lactobacillus jensenii		88.43	0.47	FRU 99	1	
Lactobacillus catenaformis		8.01	0.29		1	
Clostridium barati		2.69	0.15		99	
47341020	**Low discrimination**					**47341020**
					SPO	
Clostridium barati		42.68	0.50	FRU 99	99	
Lactobacillus catenaformis		31.71	0.53		1	
Lactobacillus jensenii		15.92	0.47		1	
47341022	**Low discrimination**					**47341022**
					SPO	
Clostridium barati		71.54	0.58	FRU 99	99	
Lactobacillus catenaformis		17.72	0.53		1	
Lactobacillus jensenii		5.93	0.44		1	
47341030	**Low discrimination**					**47341030**
					SPO	
Lactobacillus jensenii		86.47	0.39	FRU 99	1	
Clostridium barati		7.02	0.15		99	
Lactobacillus catenaformis		5.22	0.18		1	
47341032	**Low discrimination**					**47341032**
					SPO	
Lactobacillus jensenii		67.30	0.36	FRU 99	1	
Clostridium barati		24.60	0.24		99	
Lactobacillus catenaformis		6.09	0.18		1	
47341062	**Acceptable identification**					**47341062**
					SPO	
Clostridium barati		90.95	0.40	FRU 99	99	
Lactobacillus acidophilus		3.17	0.17	SOR 8	1	
Lactobacillus catenaformis		2.62	0.18		1	
47341100	**Low discrimination**					**47341100**
					SPO	
Collinsella aerofaciens		84.99	0.66	FRU 99	1	
Lactobacillus catenaformis		11.68	0.66		1	
Clostridium barati		2.62	0.50		99	
47341102	**Low discrimination**					**47341102**
					SPO	
Lactobacillus catenaformis		36.82	0.66	FRU 99	1	
Collinsella aerofaciens		29.77	0.50		1	
Clostridium barati		24.78	0.58		99	
47341120	**Low discrimination**					**47341120**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		69.82	0.66	FRU 99	99		
Collinsella aerofaciens		13.86	0.50		1		
Clostridium barati		11.53	0.58		99		
47341122	**Low discrimination**						**47341122**
					SPO		
Clostridium barati		45.08	0.66	FRU 99	99		
Clostridium paraputrificum		39.00	0.60		99		
Lactobacillus catenaformis		5.58	0.56		1		
47341132	**Low discrimination**						**47341132**
					SPO		
Bifidobacterium breve		28.26	0.44	RAF 95	1		
Clostridium barati		27.93	0.32	FRU 99	99		
Clostridium paraputrificum		24.16	0.25		99		
47341142	**Low discrimination**						**47341142**
					SPO		
Clostridium barati		51.69	0.40	FRU 99	99		
Lactobacillus acidophilus		20.74	0.35	SOR 8	1		
Bifidobacterium breve		10.25	0.40		1		
47341152	**Low discrimination**						**47341152**
					GRA	SPO	
Bifidobacterium breve		89.30	0.40	RAF 95	99	1	
Clostridium barati		4.55	0.05	FRU 99	99	99	
Leptotrichia buccalis		4.33	-.03		1	1	
47341160	**Low discrimination**						**47341160**
					SPO		
Clostridium paraputrificum		81.03	0.47	FRU 99	99		
Clostridium barati		15.46	0.40	SOR 7	99		
Collinsella aerofaciens		2.16	0.15		1		
47341162	**Low discrimination**						**47341162**
					SPO		
Clostridium barati		49.30	0.48	FRU 99	99		
Clostridium paraputrificum		36.91	0.40	SOR 8	99		
Lactobacillus acidophilus		6.59	0.35		1		
47341172	**Low discrimination**						**47341172**
					SPO		
Bifidobacterium breve		86.08	0.44	RAF 95	1		
Clostridium barati		7.40	0.14	FRU 99	99		
Clostridium paraputrificum		5.54	0.06		99		
47341222	**Low discrimination**						**47341222**
					GRA	SPO	
Clostridium barati		88.18	0.40	FRU 99	99	99	
Prevotella oralis		6.29	0.21	RHA 8	1	1	
Lactobacillus catenaformis		2.54	0.18		99	1	
47341302	**Low discrimination**						**47341302**
					SPO		
Clostridium barati		61.79	0.40	FRU 99	99		
Lactobacillus acidophilus		15.10	0.31	RHA 8	1		
Lactobacillus catenaformis		10.67	0.32		1		
47341320	**Low discrimination**						**47341320**
					SPO		
Clostridium barati		50.72	0.40	FRU 99	99		
Clostridium paraputrificum		35.68	0.32	RHA 8	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Collinsella aerofaciens		7.08	0.15		1		
47341322	**Low discrimination**						**47341322**
					GRA	SPO	
Clostridium barati		62.08	0.48	FRU 99	99	99	
Prevotella oralis		23.13	0.41	RHA 8	1	1	
Clostridium paraputrificum		6.24	0.25		99	99	
47343022	**Acceptable identification**						**47343022**
					SPO		
Clostridium barati		92.15	0.40	FRU 99	99		
Lactobacillus catenaformis		2.65	0.18	MAN 8	1		
Lactobacillus acidophilus		1.37	0.11		1		
47343101	**Low discrimination**						**47343101**
					SPO		
Actinomyces naeslundii		65.23	0.40	ESL 12	1		
Collinsella aerofaciens		27.48	-.03	RAF 92	1		
Clostridium barati		7.29	-.03	FRU 95	99		
47343102	**Low discrimination**						**47343102**
					SPO		
Clostridium barati		48.02	0.40	FRU 99	99		
Clostridium tertium		16.32	0.35	MAN 8	99		
Bifidobacterium breve		9.53	0.40		1		
47343103	**Low discrimination**						**47343103**
					SPO		
Actinomyces naeslundii		69.00	0.40	ESL 12	1		
Clostridium barati		23.14	0.05	RAF 92	99		
Clostridium tertium		7.86	0.01	FRU 95	99		
47343112	**Low discrimination**						**47343112**
					SPO		
Clostridium tertium		72.16	0.39	FRU 99	99		
Bifidobacterium breve		23.69	0.40	TRE 90	1		
Bifidobacterium adolescentis		1.44	0.18		1		
47343120	**Low discrimination**						**47343120**
Clostridium barati		41.76	0.40	FRU 99			
Clostridium paraputrificum		29.37	0.32	MAN 8			
Clostridium tertium		14.19	0.35				
47343121	**Low discrimination**						**47343121**
					SPO		
Actinomyces naeslundii		59.34	0.40	ESL 12	1		
Clostridium barati		19.90	0.05	RAF 92	99		
Clostridium paraputrificum		14.00	-.03	FRU 95	99		
47343122	**Low discrimination**						**47343122**
Clostridium barati		55.50	0.48	FRU 99			
Clostridium tertium		18.86	0.43	MAN 8			
Clostridium ramosum		8.43	0.35				
47343130	**Acceptable identification**						**47343130**
					SPO		
Clostridium tertium		92.06	0.39	FRU 99	99		
Bifidobacterium breve		5.31	0.27	TRE 90	1		
Clostridium barati		1.54	0.05		99		
47343132	**Low discrimination**						**47343132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium tertium		82.27	0.47	FRU 99	99	
Bifidobacterium breve		16.01	0.44	TRE 90	1	
Clostridium barati		1.38	0.14		99	
47343142	**Low discrimination**					**47343142**
					SPO	
Bifidobacterium breve		61.75	0.40	RAF 95	1	
Clostridium barati		27.07	0.21	FRU 99	99	
Lactobacillus acidophilus		6.58	0.13		1	
47343152	**Good identification**					**47343152**
					SPO	
Bifidobacterium breve		97.02	0.40	RAF 95	1	
Clostridium tertium		2.99	0.05	FRU 99	99	
47343162	**Low discrimination**					**47343162**
					SPO	
Bifidobacterium breve		52.18	0.44	RAF 95	1	
Clostridium barati		38.60	0.30	FRU 99	99	
Clostridium paraputrificum		3.36	0.06		99	
47343172	**Good identification**					**47343172**
					SPO	
Bifidobacterium breve		95.06	0.44	RAF 95	1	
Clostridium tertium		4.94	0.13	FRU 99	99	
47343322	**Low discrimination**					**47343322**
					SPO	
Clostridium ramosum		66.68	0.37	FRU 99	92	
Clostridium barati		28.79	0.30	TRE 94	99	
Lactobacillus acidophilus		1.42	0.09		1	
47345502	**Low discrimination**					**47345502**
					SPO	
Bifidobacterium adolescentis		83.28	0.46	RAF 92	1	
Clostridium butyricum		11.13	0.17	FRU 99	99	
Actinomyces israelii		5.13	0.32		1	
47345512	**Good identification**					**47345512**
Bifidobacterium adolescentis		96.70	0.48	RAF 92		
Clostridium clostridioforme		2.02	0.16	FRU 99		
Actinomyces israelii		0.73	0.18			
47347002	**Low discrimination**					**47347002**
					SPO	
Actinomyces israelii		68.96	0.36	FRU 99	1	
Clostridium innocuum		21.35	0.12		99	
Clostridium tertium		5.67	0.03		99	
47347102	**Low discrimination**					**47347102**
					SPO	
Clostridium tertium		60.45	0.37	FRU 99	99	
Actinomyces israelii		29.72	0.46	TRE 90	1	
Bifidobacterium adolescentis		8.52	0.31		1	
47347112	**Low discrimination**					**47347112**
					SPO	
Clostridium tertium		87.86	0.41	FRU 99	99	
Bifidobacterium adolescentis		8.51	0.32	TRE 90	1	
Actinomyces israelii		3.63	0.32		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47347120	**Acceptable identification**					**47347120**
					SPO	
Clostridium tertium		94.56	0.37	FRU 99	99	
Clostridium barati		2.04	0.05	TRE 90	99	
Actinomyces israelii		1.98	0.23		1	
47347122	**Good identification**					**47347122**
					SPO	
Clostridium tertium		95.95	0.45	FRU 99	99	
Clostridium barati		2.07	0.14	TRE 90	99	
Actinomyces israelii		1.00	0.26		1	
47347130	**Good identification**					**47347130**
Clostridium tertium		100.0	0.41	FRU 99		
47347132	**Good identification**					**47347132**
					SPO	
Clostridium tertium		99.86	0.50	FRU 99	99	
Bifidobacterium breve		0.14	0.10	TRE 90	1	
47347140	**Good identification**					**47347140**
Actinomyces israelii		100.0	0.36	FRU 99		
47347142	**Low discrimination**					**47347142**
					SPO	
Actinomyces israelii		87.42	0.39	FRU 99	1	
Bifidobacterium adolescentis		7.97	0.15		1	
Clostridium tertium		4.62	0.03		99	
47347500	**Low discrimination**					**47347500**
					SPO	
Actinomyces israelii		83.35	0.42	FRU 99	1	
Clostridium butyricum		13.69	0.07		99	
Bifidobacterium adolescentis		2.96	0.11		1	
47347502	**Low discrimination**					**47347502**
					SPO	
Bifidobacterium adolescentis		68.71	0.45	RAF 92	1	
Actinomyces israelii		29.30	0.45	FRU 99	1	
Clostridium butyricum		1.25	0.00		99	
47347512	**Low discrimination**					**47347512**
Bifidobacterium adolescentis		89.68	0.46	RAF 92		
Actinomyces israelii		4.68	0.31	FRU 99		
Bifidobacterium dentium		4.25	0.06			
47347542	**Low discrimination**					**47347542**
Actinomyces israelii		57.30	0.38	FRU 99		
Bifidobacterium adolescentis		42.70	0.29			
47350000	**Low discrimination**					**47350000**
					SPO	
Lactobacillus catenaformis		41.07	0.63	GAL 99	1	
Clostridium barati		24.00	0.54		99	
Lactobacillus jensenii		20.62	0.58		1	
47350002	**Low discrimination**					**47350002**
					SPO	
Clostridium barati		55.60	0.62	GAL 95	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus catenaformis		31.72	0.63		1	
Lactobacillus jensenii		10.62	0.55		1	
47350010	**Low discrimination**					**47350010**
					SPO	
Lactobacillus jensenii		89.56	0.50	GAL 99	1	
Lactobacillus catenaformis		5.41	0.29		1	
Clostridium barati		3.16	0.19		99	
47350012	**Low discrimination**					**47350012**
					SPO	
Lactobacillus jensenii		79.28	0.47	GAL 99	1	
Clostridium barati		12.58	0.28		99	
Lactobacillus catenaformis		7.18	0.29		1	
47350020	**Low discrimination**					**47350020**
					SPO	
Clostridium barati		79.45	0.62	GAL 95	99	
Lactobacillus catenaformis		11.33	0.53		1	
Lactobacillus jensenii		5.69	0.47		1	
47350022	**Good identification**					**47350022**
					SPO	
Clostridium barati		92.90	0.70	GAL 95	99	
Lactobacillus catenaformis		4.42	0.53		1	
Lactobacillus jensenii		1.48	0.44		1	
47350023	**Acceptable identification**					**47350023**
					SPO	
Clostridium barati		92.57	0.36	GAL 95	99	
Lactobacillus catenaformis		4.40	0.18	URE 1	1	
Lactobacillus jensenii		1.47	0.10		1	
47350030	**Low discrimination**					**47350030**
					SPO	
Lactobacillus jensenii		66.73	0.39	GAL 99	1	
Clostridium barati		28.24	0.28		99	
Lactobacillus catenaformis		4.03	0.18		1	
47350032	**Low discrimination**					**47350032**
					SPO	
Clostridium barati		62.96	0.36	GAL 95	99	
Lactobacillus jensenii		33.06	0.36	MLZ 1	1	
Lactobacillus catenaformis		2.99	0.18		1	
47350040	**Low discrimination**					**47350040**
					SPO	
Clostridium barati		73.12	0.36	GAL 95	99	
Lactobacillus catenaformis		14.54	0.29	SOR 8	1	
Lactobacillus jensenii		7.30	0.24		1	
47350042	**Acceptable identification**					**47350042**
					SPO	
Clostridium barati		90.07	0.44	GAL 95	99	
Lactobacillus catenaformis		5.97	0.29	SOR 8	1	
Lactobacillus jensenii		2.00	0.21		1	
47350060	**Acceptable identification**					**47350060**
					SPO	
Clostridium barati		94.75	0.44	GAL 95	99	
Clostridium paraputrificum		2.89	0.13	SOR 8	99	
Lactobacillus catenaformis		1.57	0.18		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47350062	**Good identification**					**47350062**
					SPO	
Clostridium barati		98.13	0.52	GAL 95	99	
Lactobacillus acidophilus		0.66	0.17	SOR 8	1	
Lactobacillus catenaformis		0.54	0.18		1	
47350100	**Low discrimination**					**47350100**
					SPO	
Collinsella aerofaciens		73.73	0.66	GAL 99	1	
Clostridium barati		11.84	0.62		99	
Lactobacillus catenaformis		10.13	0.66		1	
47350102	**Low discrimination**					**47350102**
					SPO	
Clostridium barati		62.73	0.70	GAL 95	99	
Lactobacillus catenaformis		17.89	0.66		1	
Collinsella aerofaciens		14.47	0.50		1	
47350103	**Low discrimination**					**47350103**
					SPO	
Clostridium barati		62.23	0.36	GAL 95	99	
Lactobacillus catenaformis		17.75	0.32	URE 1	1	
Collinsella aerofaciens		14.35	0.15		1	
47350120	**Low discrimination**					**47350120**
					SPO	
Clostridium paraputrificum		47.00	0.66	GAL 99	99	
Clostridium barati		40.44	0.70		99	
Collinsella aerofaciens		9.33	0.50		1	
47350122	**Low discrimination**					**47350122**
					SPO	
Clostridium barati		81.05	0.78	GAL 95	99	
Clostridium paraputrificum		13.46	0.60		99	
Lactobacillus catenaformis		1.93	0.56		1	
47350123	**Low discrimination**					**47350123**
					SPO	
Clostridium barati		80.67	0.44	GAL 95	99	
Clostridium paraputrificum		13.39	0.25	URE 1	99	
Lactobacillus catenaformis		1.92	0.21		1	
47350132	**Low discrimination**					**47350132**
					SPO	
Clostridium barati		66.81	0.44	GAL 95	99	
Bifidobacterium breve		12.98	0.44	MLZ 1	1	
Clostridium paraputrificum		11.09	0.25		99	
47350140	**Low discrimination**					**47350140**
					SPO	
Clostridium barati		52.54	0.44	GAL 95	99	
Collinsella aerofaciens		38.00	0.32	SOR 8	1	
Lactobacillus catenaformis		5.22	0.32		1	
47350142	**Low discrimination**					**47350142**
					SPO	
Clostridium barati		84.70	0.52	GAL 95	99	
Lactobacillus acidophilus		6.52	0.35	SOR 8	1	
Bifidobacterium breve		3.22	0.40		1	
47350152	**Low discrimination**					**47350152**
					GRA SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		74.95	0.40	RAF 95	99	1	
Clostridium barati		19.89	0.18	GAL 99	99	99	
Leptotrichia buccalis		3.63	-.03		1	1	
47350160	**Low discrimination**						**47350160**
					SPO		
Clostridium paraputrificum		49.08	0.47	GAL 99	99		
Clostridium barati		48.80	0.52	SOR 7	99		
Collinsella aerofaciens		1.31	0.15		1		
47350162	**Low discrimination**						**47350162**
					SPO		
Clostridium barati		83.52	0.60	GAL 95	99		
Clostridium paraputrificum		12.00	0.40	SOR 8	99		
Lactobacillus acidophilus		2.14	0.35		1		
47350172	**Low discrimination**						**47350172**
					SPO		
Bifidobacterium breve		65.63	0.44	RAF 95	1		
Clostridium barati		29.39	0.26	GAL 99	99		
Clostridium paraputrificum		4.22	0.06		99		
47350200	**Low discrimination**						**47350200**
					SPO		
Clostridium barati		73.12	0.36	GAL 95	99		
Lactobacillus catenaformis		14.54	0.29	RHA 8	1		
Lactobacillus jensenii		7.30	0.24		1		
47350202	**Acceptable identification**						**47350202**
					SPO		
Clostridium barati		90.53	0.44	GAL 95	99		
Lactobacillus catenaformis		6.00	0.29	RHA 8	1		
Lactobacillus jensenii		2.01	0.21		1		
47350220	**Good identification**						**47350220**
					SPO		
Clostridium barati		96.98	0.44	GAL 95	99		
Lactobacillus catenaformis		1.61	0.18	RHA 8	1		
Lactobacillus jensenii		0.81	0.13		1		
47350222	**Good identification**						**47350222**
					GRA	SPO	
Clostridium barati		97.49	0.52	GAL 95	99	99	
Prevotella oralis		1.34	0.21	RHA 8	1	1	
Lactobacillus catenaformis		0.54	0.18		99	1	
47350300	**Low discrimination**						**47350300**
					SPO		
Clostridium barati		53.61	0.44	GAL 95	99		
Collinsella aerofaciens		38.78	0.32	RHA 8	1		
Lactobacillus catenaformis		5.33	0.32		1		
47350302	**Low discrimination**						**47350302**
					SPO		
Clostridium barati		89.21	0.52	GAL 95	99		
Lactobacillus acidophilus		4.18	0.31	RHA 8	1		
Lactobacillus catenaformis		2.96	0.32		1		
47350320	**Low discrimination**						**47350320**
					SPO		
Clostridium barati		84.28	0.52	GAL 95	99		
Clostridium paraputrificum		11.38	0.32	RHA 8	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Collinsella aerofaciens		2.26	0.15		1		
47350322	**Low discrimination**						**47350322**
					GRA	SPO	
Clostridium barati		89.48	0.60	GAL 95	99	99	
Prevotella oralis		6.40	0.41	RHA 8	1	1	
Clostridium paraputrificum		1.73	0.25		99	99	
47350362	**Good identification**						**47350362**
					SPO		
Clostridium barati		95.25	0.42	GAL 95	99		
Lactobacillus acidophilus		2.12	0.16	RHA 8	1		
Clostridium paraputrificum		1.59	0.06	SOR 8	99		
47350422	**Acceptable identification**						**47350422**
					SPO		
Clostridium barati		92.76	0.36	GAL 95	99		
Lactobacillus catenaformis		4.41	0.18	ARA 1	1		
Lactobacillus jensenii		1.48	0.10		1		
47350502	**Low discrimination**						**47350502**
					SPO		
Clostridium barati		57.42	0.36	GAL 95	99		
Lactobacillus catenaformis		16.38	0.32	ARA 1	1		
Collinsella aerofaciens		13.24	0.15		1		
47350522	**Low discrimination**						**47350522**
					SPO		
Clostridium barati		80.76	0.44	GAL 95	99		
Clostridium paraputrificum		13.41	0.25	ARA 1	99		
Lactobacillus catenaformis		1.92	0.21		1		
47350712	**Good identification**						**47350712**
Clostridium clostridioforme		100.0	0.39	RAF 94			
47350732	**Good identification**						**47350732**
Clostridium clostridioforme		100.0	0.39	RAF 94			
47351000	**Low discrimination**						**47351000**
					SPO		
Lactobacillus catenaformis		57.83	0.97		1		
Lactobacillus jensenii		29.03	0.92		1		
Clostridium barati		6.49	0.76		99		
47351001	**Low discrimination**						**47351001**
					SPO		
Lactobacillus catenaformis		57.54	0.63	URE 1	1		
Lactobacillus jensenii		28.89	0.58		1		
Clostridium barati		6.45	0.41		99		
47351002	**Low discrimination**						**47351002**
					SPO		
Lactobacillus catenaformis		58.48	0.97		1		
Clostridium barati		19.67	0.84		99		
Lactobacillus jensenii		19.57	0.89		1		
47351003	**Low discrimination**						**47351003**
					SPO		
Lactobacillus catenaformis		58.18	0.63	URE 1	1		
Clostridium barati		19.58	0.50		99		
Lactobacillus jensenii		19.47	0.55		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47351010	**Good identification**					**47351010**
					SPO	
Lactobacillus jensenii		93.09	0.84		1	
Lactobacillus catenaformis		5.62	0.63		1	
Clostridium barati		0.63	0.41		99	
47351011	**Acceptable identification**					**47351011**
					SPO	
Lactobacillus jensenii		92.88	0.50	URE 1	1	
Lactobacillus catenaformis		5.61	0.29		1	
Clostridium barati		0.63	0.07		99	
47351012	**Low discrimination**					**47351012**
					SPO	
Lactobacillus jensenii		88.19	0.81		1	
Lactobacillus catenaformis		7.98	0.63		1	
Clostridium barati		2.69	0.50		99	
47351013	**Low discrimination**					**47351013**
					SPO	
Lactobacillus jensenii		88.10	0.47	URE 1	1	
Lactobacillus catenaformis		7.98	0.29		1	
Clostridium barati		2.68	0.15		99	
47351020	**Low discrimination**					**47351020**
					SPO	
Clostridium barati		42.72	0.84		99	
Lactobacillus catenaformis		31.74	0.87		1	
Lactobacillus jensenii		15.94	0.82		1	
47351021	**Low discrimination**					**47351021**
					SPO	
Clostridium barati		42.25	0.50	URE 1	99	
Lactobacillus catenaformis		31.39	0.53		1	
Lactobacillus jensenii		15.76	0.47		1	
47351022	**Low discrimination**					**47351022**
					SPO	
Clostridium barati		71.54	0.92		99	
Lactobacillus catenaformis		17.72	0.87		1	
Lactobacillus jensenii		5.93	0.79		1	
47351023	**Low discrimination**					**47351023**
					SPO	
Clostridium barati		71.10	0.58	URE 1	99	
Lactobacillus catenaformis		17.61	0.53		1	
Lactobacillus jensenii		5.89	0.44		1	
47351030	**Low discrimination**					**47351030**
					SPO	
Lactobacillus jensenii		86.15	0.73		1	
Clostridium barati		7.00	0.50		99	
Lactobacillus catenaformis		5.20	0.53		1	
47351031	**Low discrimination**					**47351031**
					SPO	
Lactobacillus jensenii		85.62	0.39	URE 1	1	
Clostridium barati		6.96	0.15		99	
Lactobacillus catenaformis		5.17	0.18		1	
47351032	**Low discrimination**					**47351032**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		66.86	0.70		1	
Clostridium barati		24.44	0.58		99	
Lactobacillus catenaformis		6.05	0.53		1	
47351033	**Low discrimination**					**47351033**
					SPO	
Lactobacillus jensenii		66.54	0.36	URE 1	1	
Clostridium barati		24.32	0.24		99	
Lactobacillus catenaformis		6.02	0.18		1	
47351040	**Low discrimination**					**47351040**
					SPO	
Lactobacillus catenaformis		38.55	0.63	SOR 1	1	
Clostridium barati		37.22	0.57		99	
Lactobacillus jensenii		19.36	0.58		1	
47351042	**Low discrimination**					**47351042**
					SPO	
Clostridium barati		63.36	0.66	SOR 8	99	
Lactobacillus catenaformis		21.88	0.63		1	
Lactobacillus jensenii		7.32	0.55		1	
47351050	**Low discrimination**					**47351050**
					SPO	
Lactobacillus jensenii		87.92	0.50	SOR 1	1	
Lactobacillus catenaformis		5.31	0.29		1	
Clostridium barati		5.12	0.23		99	
47351052	**Low discrimination**					**47351052**
					SPO	
Lactobacillus jensenii		64.61	0.47	SOR 1	1	
Clostridium barati		16.94	0.31		99	
Bifidobacterium breve		10.08	0.40		1	
47351060	**Low discrimination**					**47351060**
					SPO	
Clostridium barati		77.07	0.66	SOR 8	99	
Clostridium paraputrificum		12.24	0.47		99	
Lactobacillus catenaformis		6.65	0.53		1	
47351062	**Good identification**					**47351062**
					SPO	
Clostridium barati		90.76	0.74	SOR 8	99	
Lactobacillus acidophilus		3.17	0.51		1	
Lactobacillus catenaformis		2.61	0.53		1	
47351063	**Low discrimination**					**47351063**
					SPO	
Clostridium barati		89.46	0.40	SOR 8	99	
Lactobacillus acidophilus		3.12	0.17	URE 1	1	
Lactobacillus catenaformis		2.57	0.18		1	
47351070	**Low discrimination**					**47351070**
					SPO	
Lactobacillus jensenii		51.45	0.39	SOR 1	1	
Clostridium barati		35.97	0.31		99	
Clostridium paraputrificum		5.71	0.13		99	
47351072	**Low discrimination**					**47351072**
					SPO	
Clostridium barati		56.92	0.40	MLZ 1	99	
Bifidobacterium breve		20.07	0.44	SOR 8	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		18.09	0.36		1	
47351100	**Low discrimination**					**47351100**
					SPO	
Collinsella aerofaciens		84.99	1.00		1	
Lactobacillus catenaformis		11.68	1.00		1	
Clostridium barati		2.62	0.84		99	
47351101	**Low discrimination**					**47351101**
					SPO	
Collinsella aerofaciens		84.74	0.66	URE 1	1	
Lactobacillus catenaformis		11.65	0.66		1	
Clostridium barati		2.61	0.50		99	
47351102	**Low discrimination**					**47351102**
					SPO	
Lactobacillus catenaformis		36.83	1.00		1	
Collinsella aerofaciens		29.78	0.84		1	
Clostridium barati		24.78	0.92		99	
47351103	**Low discrimination**					**47351103**
					SPO	
Lactobacillus catenaformis		36.58	0.66	URE 1	1	
Collinsella aerofaciens		29.58	0.50		1	
Clostridium barati		24.61	0.58		99	
47351110	**Low discrimination**					**47351110**
					SPO	
Collinsella aerofaciens		82.20	0.66	MLZ 1	1	
Lactobacillus catenaformis		11.30	0.66		1	
Clostridium barati		2.53	0.50		99	
47351112	**Low discrimination**					**47351112**
					GRA	
Leptotrichia buccalis		57.66	0.66	TRE 99	1	
Bifidobacterium breve		12.02	0.74		99	
Lactobacillus catenaformis		10.46	0.66		99	
47351120	**Low discrimination**					**47351120**
					SPO	
Clostridium paraputrificum		69.85	1.00		99	
Collinsella aerofaciens		13.86	0.84		1	
Clostridium barati		11.54	0.92		99	
47351121	**Low discrimination**					**47351121**
					SPO	
Clostridium paraputrificum		69.62	0.66	URE 1	99	
Collinsella aerofaciens		13.82	0.50		1	
Clostridium barati		11.50	0.58		99	
47351122	**Low discrimination**					**47351122**
					SPO	
Clostridium barati		45.09	1.00		99	
Clostridium paraputrificum		39.00	0.94		99	
Lactobacillus catenaformis		5.58	0.90		1	
47351123	**Low discrimination**					**47351123**
					SPO	
Clostridium barati		44.92	0.66	URE 1	99	
Clostridium paraputrificum		38.85	0.60		99	
Lactobacillus catenaformis		5.56	0.56		1	
47351130	**Low discrimination**					**47351130**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium paraputrificum		66.78	0.66	MLZ 1	99		
Collinsella aerofaciens		13.25	0.50		1		
Clostridium barati		11.03	0.58		99		
47351132	**Low discrimination**						**47351132**
					SPO		
Bifidobacterium breve		28.21	0.78	RAF 95	1		
Clostridium barati		27.88	0.66		99		
Clostridium paraputrificum		24.12	0.60		99		
47351133	**Low discrimination**						**47351133**
					SPO		
Bifidobacterium breve		27.92	0.44	RAF 95	1		
Clostridium barati		27.59	0.32	URE 1	99		
Clostridium paraputrificum		23.86	0.25		99		
47351140	**Low discrimination**						**47351140**
					SPO		
Collinsella aerofaciens		68.16	0.66	SOR 1	1		
Clostridium barati		18.09	0.66		99		
Lactobacillus catenaformis		9.37	0.66		1		
47351142	**Low discrimination**						**47351142**
					SPO		
Clostridium barati		51.54	0.74	SOR 8	99		
Lactobacillus acidophilus		20.68	0.69		1		
Bifidobacterium breve		10.22	0.74		1		
47351143	**Low discrimination**						**47351143**
					SPO		
Clostridium barati		50.34	0.40	SOR 8	99		
Lactobacillus acidophilus		20.20	0.35	URE 1	1		
Bifidobacterium breve		9.99	0.40		1		
47351150	**Low discrimination**						**47351150**
					SPO		
Bifidobacterium breve		51.06	0.57	RAF 95	1		
Collinsella aerofaciens		33.03	0.32	BGL 91	1		
Clostridium barati		8.76	0.31		99		
47351152	**Low discrimination**						**47351152**
					GRA	SPO	
Bifidobacterium breve		87.10	0.74	RAF 95	99	1	
Clostridium barati		4.44	0.40		99	99	
Leptotrichia buccalis		4.22	0.32		1	1	
47351153	**Low discrimination**						**47351153**
					GRA	SPO	
Bifidobacterium breve		88.21	0.40	RAF 95	99	1	
Clostridium barati		4.49	0.05	URE 1	99	99	
Leptotrichia buccalis		4.28	-.03		1	1	
47351160	**Low discrimination**						**47351160**
					SPO		
Clostridium paraputrificum		80.98	0.81	SOR 7	99		
Clostridium barati		15.45	0.74		99		
Collinsella aerofaciens		2.16	0.50		1		
47351161	**Low discrimination**						**47351161**
					SPO		
Clostridium paraputrificum		80.39	0.47	SOR 7	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		15.34	0.40	URE 1	99	
Collinsella aerofaciens		2.14	0.15		1	
47351162	**Low discrimination**					**47351162**
					SPO	
Clostridium barati		49.26	0.82	SOR 8	99	
Clostridium paraputrificum		36.89	0.75		99	
Lactobacillus acidophilus		6.59	0.69		1	
47351163	**Low discrimination**					**47351163**
					SPO	
Clostridium barati		48.88	0.48	SOR 8	99	
Clostridium paraputrificum		36.60	0.40	URE 1	99	
Lactobacillus acidophilus		6.54	0.35		1	
47351170	**Low discrimination**					**47351170**
					SPO	
Clostridium paraputrificum		52.96	0.47	MLZ 1	99	
Bifidobacterium breve		34.89	0.61	SOR 7	1	
Clostridium barati		10.10	0.40		99	
47351172	**Low discrimination**					**47351172**
					SPO	
Bifidobacterium breve		85.61	0.78	RAF 95	1	
Clostridium barati		7.36	0.48		99	
Clostridium paraputrificum		5.51	0.40		99	
47351173	**Low discrimination**					**47351173**
					SPO	
Bifidobacterium breve		85.50	0.44	RAF 95	1	
Clostridium barati		7.35	0.14	URE 1	99	
Clostridium paraputrificum		5.50	0.06		99	
47351200	**Low discrimination**					**47351200**
					SPO	
Lactobacillus catenaformis		38.67	0.63	RHA 1	1	
Clostridium barati		37.33	0.57		99	
Lactobacillus jensenii		19.42	0.58		1	
47351202	**Low discrimination**					**47351202**
					SPO	
Clostridium barati		64.91	0.66	RHA 8	99	
Lactobacillus catenaformis		22.41	0.63		1	
Lactobacillus jensenii		7.50	0.55		1	
47351210	**Low discrimination**					**47351210**
					SPO	
Lactobacillus jensenii		88.87	0.50	RHA 1	1	
Lactobacillus catenaformis		5.36	0.29		1	
Clostridium barati		5.18	0.23		99	
47351212	**Low discrimination**					**47351212**
					SPO	
Lactobacillus jensenii		71.69	0.47	RHA 1	1	
Clostridium barati		18.80	0.31		99	
Lactobacillus catenaformis		6.49	0.29		1	
47351220	**Low discrimination**					**47351220**
					SPO	
Clostridium barati		85.62	0.66	RHA 8	99	
Lactobacillus catenaformis		7.39	0.53		1	
Lactobacillus jensenii		3.71	0.47		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47351222	**Low discrimination**						**47351222**
					GRA	SPO	
Clostridium barati		88.08	0.74	RHA 8	99	99	
Prevotella oralis		6.28	0.55		1	1	
Lactobacillus catenaformis		2.53	0.53		99	1	
47351223	**Low discrimination**						**47351223**
					GRA	SPO	
Clostridium barati		88.18	0.40	RHA 8	99	99	
Prevotella oralis		6.29	0.21	URE 1	1	1	
Lactobacillus catenaformis		2.54	0.18		99	1	
47351230	**Low discrimination**						**47351230**
					SPO		
Lactobacillus jensenii		56.10	0.39	RHA 1	1		
Clostridium barati		39.23	0.31		99		
Lactobacillus catenaformis		3.39	0.18		1		
47351232	**Low discrimination**						**47351232**
					GRA	SPO	
Clostridium barati		68.28	0.40	RHA 8	99	99	
Lactobacillus jensenii		21.70	0.36	MLZ 1	99	1	
Prevotella oralis		4.87	0.21		1	1	
47351240	**Low discrimination**						**47351240**
					SPO		
Clostridium barati		83.75	0.39	RHA 8	99		
Lactobacillus catenaformis		10.08	0.29	SOR 8	1		
Lactobacillus jensenii		5.06	0.24		1		
47351242	**Low discrimination**						**47351242**
					SPO		
Clostridium barati		87.30	0.47	RHA 8	99		
Lactobacillus acidophilus		7.91	0.32	SOR 8	1		
Lactobacillus catenaformis		3.50	0.29		1		
47351260	**Good identification**						**47351260**
					SPO		
Clostridium barati		96.64	0.47	RHA 8	99		
Clostridium paraputrificum		1.78	0.13	SOR 8	99		
Lactobacillus catenaformis		0.97	0.18		1		
47351262	**Good identification**						**47351262**
					GRA	SPO	
Clostridium barati		95.60	0.56	RHA 8	99	99	
Lactobacillus acidophilus		2.89	0.32	SOR 8	99	1	
Prevotella oralis		0.79	0.21		1	1	
47351300	**Low discrimination**						**47351300**
					SPO		
Collinsella aerofaciens		70.58	0.66	RHA 1	1		
Clostridium barati		18.73	0.66		99		
Lactobacillus catenaformis		9.70	0.66		1		
47351302	**Low discrimination**						**47351302**
					SPO		
Clostridium barati		61.60	0.74	RHA 8	99		
Lactobacillus acidophilus		15.05	0.66		1		
Lactobacillus catenaformis		10.64	0.66		1		
47351303	**Low discrimination**						**47351303**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		61.67	0.40	RHA 8	99		
Lactobacillus acidophilus		15.07	0.31	URE 1	1		
Lactobacillus catenaformis		10.65	0.32		1		
47351312	**Low discrimination**						**47351312**
					GRA	SPO	
Clostridium barati		34.85	0.40	RHA 8	99	99	
Leptotrichia buccalis		33.17	0.32	MLZ 1	1	1	
Lactobacillus acidophilus		8.52	0.31		99	1	
47351320	**Low discrimination**						**47351320**
					SPO		
Clostridium barati		50.58	0.74	RHA 8	99		
Clostridium paraputrificum		35.58	0.66		99		
Collinsella aerofaciens		7.06	0.50		1		
47351321	**Low discrimination**						**47351321**
					SPO		
Clostridium barati		50.64	0.40	RHA 8	99		
Clostridium paraputrificum		35.62	0.32	URE 1	99		
Collinsella aerofaciens		7.07	0.15		1		
47351322	**Low discrimination**						**47351322**
					GRA	SPO	
Clostridium barati		62.06	0.82	RHA 8	99	99	
Prevotella oralis		23.12	0.76		1	1	
Clostridium paraputrificum		6.24	0.60		99	99	
47351323	**Low discrimination**						**47351323**
					GRA	SPO	
Clostridium barati		62.04	0.48	RHA 8	99	99	
Prevotella oralis		23.12	0.41	URE 1	1	1	
Clostridium paraputrificum		6.23	0.25		99	99	
47351330	**Low discrimination**						**47351330**
					SPO		
Clostridium barati		49.57	0.40	RHA 8	99		
Clostridium paraputrificum		34.87	0.32	MLZ 1	99		
Collinsella aerofaciens		6.92	0.15		1		
47351332	**Low discrimination**						**47351332**
					GRA	SPO	
Clostridium barati		55.45	0.48	RHA 8	99	99	
Prevotella oralis		20.66	0.41	MLZ 1	1	1	
Bifidobacterium breve		6.52	0.44		99	1	
47351340	**Low discrimination**						**47351340**
					SPO		
Clostridium barati		65.05	0.47	RHA 8	99		
Collinsella aerofaciens		28.48	0.32	SOR 8	1		
Lactobacillus catenaformis		3.91	0.32		1		
47351342	**Low discrimination**						**47351342**
					SPO		
Clostridium barati		70.56	0.56	RHA 8	99		
Lactobacillus acidophilus		24.51	0.50	SOR 8	1		
Bifidobacterium breve		1.63	0.40		1		
47351352	**Low discrimination**						**47351352**
					SPO		
Bifidobacterium breve		61.01	0.40	RAF 95	1		
Clostridium barati		26.75	0.21	RHA 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		9.29	0.16		1	
47351360	**Low discrimination**					**47351360**
					SPO	
Clostridium barati		60.68	0.56	RHA 8	99	
Clostridium paraputrificum		36.94	0.47	SOR 8	99	
Collinsella aerofaciens		0.98	0.15		1	
47351362	**Low discrimination**					**47351362**
					SPO	
Clostridium barati		78.98	0.64	RHA 8	99	
Lactobacillus acidophilus		9.14	0.50	SOR 8	1	
Clostridium paraputrificum		6.87	0.40		99	
47351372	**Low discrimination**					**47351372**
					SPO	
Bifidobacterium breve		52.04	0.44	RAF 95	1	
Clostridium barati		38.49	0.30	RHA 1	99	
Lactobacillus acidophilus		4.46	0.16		1	
47351400	**Low discrimination**					**47351400**
					SPO	
Lactobacillus catenaformis		57.60	0.63	ARA 1	1	
Lactobacillus jensenii		28.92	0.58		1	
Clostridium barati		6.46	0.41		99	
47351402	**Low discrimination**					**47351402**
					SPO	
Lactobacillus catenaformis		57.95	0.63	ARA 1	1	
Clostridium barati		19.50	0.50		99	
Lactobacillus jensenii		19.40	0.55		1	
47351410	**Acceptable identification**					**47351410**
					SPO	
Lactobacillus jensenii		92.81	0.50	ARA 1	1	
Lactobacillus catenaformis		5.60	0.29		1	
Clostridium barati		0.63	0.07		99	
47351412	**Low discrimination**					**47351412**
Lactobacillus jensenii		80.09	0.47	ARA 1		
Lactobacillus catenaformis		7.25	0.29			
Bifidobacterium adolescentis		6.23	0.34			
47351420	**Low discrimination**					**47351420**
					SPO	
Clostridium barati		42.72	0.50	ARA 1	99	
Lactobacillus catenaformis		31.74	0.53		1	
Lactobacillus jensenii		15.94	0.47		1	
47351422	**Low discrimination**					**47351422**
					SPO	
Clostridium barati		71.50	0.58	ARA 1	99	
Lactobacillus catenaformis		17.71	0.53		1	
Lactobacillus jensenii		5.93	0.44		1	
47351430	**Low discrimination**					**47351430**
					SPO	
Lactobacillus jensenii		86.47	0.39	ARA 1	1	
Clostridium barati		7.02	0.15		99	
Lactobacillus catenaformis		5.22	0.18		1	
47351432	**Low discrimination**					**47351432**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Lactobacillus jensenii		62.01	0.36	ARA 1	1	
Clostridium barati		22.67	0.24		99	
Clostridium clostridioforme		7.86	0.24		66	
47351462	**Acceptable identification**					**47351462**
					SPO	
Clostridium barati		90.95	0.40	ARA 1	99	
Lactobacillus acidophilus		3.17	0.17	SOR 8	1	
Lactobacillus catenaformis		2.62	0.18		1	
47351500	**Low discrimination**					**47351500**
					SPO	
Collinsella aerofaciens		84.51	0.66	ARA 1	1	
Lactobacillus catenaformis		11.61	0.66		1	
Clostridium barati		2.61	0.50		99	
47351502	**Low discrimination**					**47351502**
					SPO	
Lactobacillus catenaformis		31.11	0.66	ARA 1	1	
Collinsella aerofaciens		25.15	0.50		1	
Clostridium barati		20.93	0.58		99	
47351512	**Low discrimination**					**47351512**
					GRA	
Bifidobacterium adolescentis		84.30	0.68	RAF 92	99	
Leptotrichia buccalis		8.20	0.32	XYL 87	1	
Bifidobacterium breve		1.71	0.40		99	
47351520	**Low discrimination**					**47351520**
					SPO	
Clostridium paraputrificum		69.80	0.66	ARA 1	99	
Collinsella aerofaciens		13.85	0.50		1	
Clostridium barati		11.53	0.58		99	
47351522	**Low discrimination**					**47351522**
					SPO	
Clostridium barati		44.93	0.66	ARA 1	99	
Clostridium paraputrificum		38.86	0.60		99	
Lactobacillus catenaformis		5.56	0.56		1	
47351532	**Low discrimination**					**47351532**
					SPO	
Bifidobacterium breve		23.47	0.44	RAF 95	1	
Clostridium barati		23.20	0.32	ARA 1	99	
Clostridium paraputrificum		20.06	0.25		99	
47351542	**Low discrimination**					**47351542**
					SPO	
Clostridium barati		32.74	0.40	ARA 1	99	
Bifidobacterium adolescentis		32.39	0.51	SOR 8	1	
Lactobacillus acidophilus		13.14	0.35		1	
47351552	**Low discrimination**					**47351552**
Bifidobacterium adolescentis		83.51	0.52	RAF 92		
Bifidobacterium breve		13.70	0.40	XYL 87		
Actinomyces israelii		1.15	0.27	SOR 11		
47351560	**Low discrimination**					**47351560**
					SPO	
Clostridium paraputrificum		80.92	0.47	ARA 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		15.44	0.40	SOR 7	99		
Collinsella aerofaciens		2.16	0.15		1		
47351562	**Low discrimination**						**47351562**
					SPO		
Clostridium barati		49.08	0.48	ARA 1	99		
Clostridium paraputrificum		36.75	0.40	SOR 8	99		
Lactobacillus acidophilus		6.56	0.35		1		
47351572	**Low discrimination**						**47351572**
					SPO		
Bifidobacterium breve		83.58	0.44	RAF 95	1		
Clostridium barati		7.18	0.14	ARA 1	99		
Clostridium paraputrificum		5.38	0.06		99		
47351612	**Acceptable identification**						**47351612**
					SPO		
Clostridium clostridioforme		94.80	0.39	RAF 94	66		
Lactobacillus jensenii		4.12	0.12	TRE 88	1		
Clostridium barati		1.08	-.03	XYL 91	99		
47351622	**Low discrimination**						**47351622**
					GRA	SPO	
Clostridium barati		83.41	0.40	RHA 8	99	99	
Prevotella oralis		5.95	0.21	ARA 1	1	1	
Clostridium clostridioforme		5.41	0.27		50	66	
47351632	**Good identification**						**47351632**
					SPO		
Clostridium clostridioforme		95.69	0.39	RAF 94	66		
Clostridium barati		3.27	0.05	TRE 88	99		
Lactobacillus jensenii		1.04	0.02	XYL 91	1		
47351702	**Low discrimination**						**47351702**
					SPO		
Clostridium barati		50.79	0.40	RHA 8	99		
Lactobacillus acidophilus		12.41	0.31	ARA 1	1		
Clostridium clostridioforme		11.03	0.36		66		
47351712	**Low discrimination**						**47351712**
					SPO		
Clostridium clostridioforme		89.04	0.48	RAF 94	66		
Bifidobacterium adolescentis		8.89	0.34	TRE 88	1		
Clostridium barati		0.91	0.05	XYL 91	99		
47351720	**Low discrimination**						**47351720**
					SPO		
Clostridium barati		48.82	0.40	RHA 8	99		
Clostridium paraputrificum		34.34	0.32	ARA 1	99		
Collinsella aerofaciens		6.82	0.15		1		
47351722	**Low discrimination**						**47351722**
					GRA	SPO	
Clostridium barati		58.36	0.48	RHA 8	99	99	
Prevotella oralis		21.74	0.41	ARA 1	1	1	
Clostridium paraputrificum		5.86	0.25		99	99	
47351732	**Acceptable identification**						**47351732**
					GRA	SPO	
Clostridium clostridioforme		93.24	0.48	RAF 94	50	66	
Clostridium barati		2.86	0.14	TRE 88	99	99	
Parabacteroides distasonis		2.22	0.25	XYL 91	1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47352000	**Low discrimination**					**47352000**
					SPO	
Clostridium barati		50.78	0.36	GAL 95	99	
Clostridium innocuum		29.77	0.34	MAN 8	99	
Lactobacillus catenaformis		10.10	0.29		1	
47352002	**Low discrimination**					**47352002**
					SPO	
Clostridium barati		76.79	0.44	GAL 95	99	
Clostridium innocuum		15.00	0.34	MAN 8	99	
Lactobacillus catenaformis		5.09	0.29		1	
47352020	**Good identification**					**47352020**
					SPO	
Clostridium barati		96.51	0.44	GAL 95	99	
Lactobacillus catenaformis		1.60	0.18	MAN 8	1	
Lactobacillus jensenii		0.80	0.13		1	
47352022	**Good identification**					**47352022**
					SPO	
Clostridium barati		98.42	0.52	GAL 95	99	
Lactobacillus catenaformis		0.54	0.18	MAN 8	1	
Lactobacillus acidophilus		0.28	0.11		1	
47352100	**Low discrimination**					**47352100**
					SPO	
Clostridium barati		51.05	0.44	GAL 95	99	
Collinsella aerofaciens		36.93	0.32	MAN 8	1	
Lactobacillus catenaformis		5.07	0.32		1	
47352101	**Low discrimination**					**47352101**
					SPO	
Actinomyces naeslundii		44.10	0.39	ESL 12	1	
Clostridium barati		32.44	0.10	RAF 92	99	
Collinsella aerofaciens		23.46	-.03	GAL 96	1	
47352102	**Low discrimination**					**47352102**
					SPO	
Clostridium barati		82.76	0.52	GAL 95	99	
Clostridium tertium		5.40	0.35	MAN 8	99	
Bifidobacterium breve		3.15	0.40		1	
47352112	**Low discrimination**					**47352112**
					SPO	
Clostridium tertium		68.67	0.39	TRE 90	99	
Bifidobacterium breve		22.54	0.40	GAL 99	1	
Clostridium barati		5.98	0.18		99	
47352120	**Low discrimination**					**47352120**
Clostridium barati		78.93	0.52	GAL 95		
Clostridium paraputrificum		10.66	0.32	MAN 8		
Clostridium tertium		5.15	0.35			
47352122	**Low discrimination**					**47352122**
Clostridium barati		86.68	0.60	GAL 95		
Clostridium tertium		5.65	0.43	MAN 8		
Clostridium ramosum		2.53	0.35			
47352130	**Low discrimination**					**47352130**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		86.46	0.39	TRE 90	99	
Clostridium barati		7.53	0.18	GAL 99	99	
Bifidobacterium breve		4.99	0.27		1	
47352132	**Low discrimination**					**47352132**
					SPO	
Clostridium tertium		77.77	0.47	TRE 90	99	
Bifidobacterium breve		15.13	0.44	GAL 99	1	
Clostridium barati		6.77	0.26		99	
47352152	**Acceptable identification**					**47352152**
					SPO	
Bifidobacterium breve		94.89	0.40	RAF 95	1	
Clostridium tertium		2.92	0.05	GAL 99	99	
Clostridium barati		2.19	-.01		99	
47352162	**Low discrimination**					**47352162**
					SPO	
Clostridium barati		76.65	0.42	GAL 95	99	
Bifidobacterium breve		19.88	0.44	MAN 8	1	
Clostridium paraputrificum		1.28	0.06	SOR 8	99	
47352172	**Acceptable identification**					**47352172**
					SPO	
Bifidobacterium breve		91.67	0.44	RAF 95	1	
Clostridium tertium		4.76	0.13	GAL 99	99	
Clostridium barati		3.57	0.08		99	
47352322	**Low discrimination**					**47352322**
					SPO	
Clostridium barati		67.81	0.42	GAL 95	99	
Clostridium ramosum		30.14	0.37	MAN 8	92	
Lactobacillus acidophilus		0.64	0.09	RHA 8	1	
47353000	**Low discrimination**					**47353000**
					SPO	
Lactobacillus catenaformis		31.23	0.63	MAN 1	1	
Clostridium barati		30.15	0.57		99	
Clostridium innocuum		17.68	0.56		99	
47353001	**Low discrimination**					**47353001**
					SPO	
Actinomyces naeslundii		35.51	0.54	ESL 12	1	
Lactobacillus catenaformis		20.33	0.29	RAF 92	1	
Clostridium barati		19.62	0.23		99	
47353002	**Low discrimination**					**47353002**
					SPO	
Clostridium barati		57.43	0.66	MAN 8	99	
Lactobacillus catenaformis		19.83	0.63		1	
Clostridium innocuum		11.22	0.56		99	
47353010	**Low discrimination**					**47353010**
					SPO	
Lactobacillus jensenii		70.83	0.50	MAN 1	1	
Clostridium innocuum		9.98	0.32		99	
Clostridium tertium		7.48	0.31		99	
47353012	**Low discrimination**					**47353012**
					SPO	
Lactobacillus jensenii		43.70	0.47	MAN 1	1	
Clostridium tertium		20.77	0.39		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		11.46	0.31		99	
47353020	**Low discrimination**					**47353020**
					SPO	
Clostridium barati		84.93	0.66	MAN 8	99	
Lactobacillus catenaformis		7.33	0.53		1	
Lactobacillus jensenii		3.68	0.47		1	
47353022	**Good identification**					**47353022**
					SPO	
Clostridium barati		92.68	0.74	MAN 8	99	
Lactobacillus catenaformis		2.67	0.53		1	
Lactobacillus acidophilus		1.38	0.45		1	
47353023	**Low discrimination**					**47353023**
					SPO	
Clostridium barati		78.38	0.40	MAN 8	99	
Actinomyces naeslundii		15.76	0.54	URE 1	1	
Lactobacillus catenaformis		2.26	0.18		1	
47353030	**Low discrimination**					**47353030**
					SPO	
Clostridium tertium		40.34	0.39	TRE 90	99	
Lactobacillus jensenii		31.82	0.39	LAC 99	1	
Clostridium barati		22.25	0.31		99	
47353032	**Low discrimination**					**47353032**
					SPO	
Clostridium tertium		51.08	0.47	TRE 90	99	
Clostridium barati		28.18	0.40	LAC 99	99	
Bifidobacterium breve		9.94	0.44		1	
47353040	**Low discrimination**					**47353040**
					SPO	
Clostridium barati		69.35	0.39	MAN 8	99	
Actinomyces israelii		9.34	0.42	SOR 8	1	
Lactobacillus catenaformis		8.34	0.29		1	
47353041	**Low discrimination**					**47353041**
					SPO	
Actinomyces naeslundii		78.60	0.41	ESL 12	1	
Clostridium barati		21.40	0.05	RAF 92	99	
47353042	**Low discrimination**					**47353042**
					SPO	
Clostridium barati		77.79	0.47	MAN 8	99	
Bifidobacterium breve		5.38	0.40	SOR 8	1	
Actinomyces israelii		5.24	0.45		1	
47353043	**Low discrimination**					**47353043**
					SPO	
Actinomyces naeslundii		55.04	0.41	ESL 12	1	
Clostridium barati		44.96	0.13	RAF 92	99	
47353052	**Low discrimination**					**47353052**
					SPO	
Bifidobacterium breve		71.39	0.40	RAF 95	1	
Clostridium barati		10.43	0.13	LAC 99	99	
Actinomyces israelii		10.39	0.31		1	
47353060	**Acceptable identification**					**47353060**
					SPO	
Clostridium barati		94.52	0.47	MAN 8	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium paraputrificum		1.74	0.13	SOR 8	99	
Bifidobacterium breve		1.15	0.27		1	
47353061	**Low discrimination**					**47353061**
					SPO	
Actinomyces naeslundii		55.04	0.41	ESL 12	1	
Clostridium barati		44.96	0.13	RAF 92	99	
47353062	**Acceptable identification**					**47353062**
					SPO	
Clostridium barati		93.02	0.56	MAN 8	99	
Bifidobacterium breve		3.81	0.44	SOR 8	1	
Lactobacillus acidophilus		1.96	0.29		1	
47353072	**Low discrimination**					**47353072**
					SPO	
Bifidobacterium breve		76.48	0.44	RAF 95	1	
Clostridium barati		18.86	0.21	LAC 99	99	
Clostridium tertium		3.97	0.13		99	
47353100	**Low discrimination**					**47353100**
					SPO	
Collinsella aerofaciens		65.23	0.66	MAN 1	1	
Clostridium barati		17.31	0.66		99	
Lactobacillus catenaformis		8.96	0.66		1	
47353102	**Low discrimination**					**47353102**
					SPO	
Clostridium barati		48.41	0.74	MAN 8	99	
Clostridium tertium		16.45	0.69		99	
Bifidobacterium breve		9.60	0.74		1	
47353103	**Low discrimination**					**47353103**
					SPO	
Clostridium barati		38.01	0.40	MAN 8	99	
Actinomyces naeslundii		21.75	0.62	URE 1	1	
Clostridium tertium		12.92	0.35		99	
47353110	**Low discrimination**					**47353110**
					SPO	
Clostridium tertium		83.39	0.65	TRE 90	99	
Bifidobacterium breve		8.12	0.57		1	
Collinsella aerofaciens		5.25	0.32		1	
47353112	**Low discrimination**					**47353112**
					SPO	
Clostridium tertium		71.70	0.73	TRE 90	99	
Bifidobacterium breve		23.54	0.74		1	
Bifidobacterium adolescentis		1.43	0.52		1	
47353113	**Low discrimination**					**47353113**
					SPO	
Clostridium tertium		69.65	0.39	TRE 90	99	
Bifidobacterium breve		22.87	0.40	URE 1	1	
Actinomyces naeslundii		3.47	0.40		1	
47353120	**Low discrimination**					**47353120**
Clostridium barati		41.93	0.74	MAN 8		
Clostridium paraputrificum		29.49	0.66			
Clostridium tertium		14.25	0.69			
47353121	**Low discrimination**					**47353121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium barati		33.93	0.40	MAN 8	99	
Clostridium paraputrificum		23.87	0.32	URE 1	99	
Actinomyces naeslundii		19.42	0.62		1	
47353122	**Low discrimination**					**47353122**
Clostridium barati		55.65	0.82	MAN 8		
Clostridium tertium		18.91	0.77			
Clostridium ramosum		8.45	0.69			
47353123	**Low discrimination**					**47353123**
					SPO	
Clostridium barati		50.36	0.48	MAN 8	99	
Clostridium tertium		17.12	0.43	URE 1	99	
Actinomyces naeslundii		9.61	0.62		1	
47353130	**Good identification**					**47353130**
					SPO	
Clostridium tertium		91.59	0.73	TRE 90	99	
Bifidobacterium breve		5.29	0.61		1	
Clostridium barati		1.53	0.40		99	
47353131	**Low discrimination**					**47353131**
					SPO	
Clostridium tertium		88.02	0.39	TRE 90	99	
Bifidobacterium breve		5.08	0.27	URE 1	1	
Actinomyces naeslundii		4.39	0.40		1	
47353132	**Low discrimination**					**47353132**
					SPO	
Clostridium tertium		82.12	0.81	TRE 90	99	
Bifidobacterium breve		15.98	0.78		1	
Clostridium barati		1.37	0.48		99	
47353133	**Low discrimination**					**47353133**
					SPO	
Clostridium tertium		81.16	0.47	TRE 90	99	
Bifidobacterium breve		15.79	0.44	URE 1	1	
Clostridium barati		1.36	0.14		99	
47353140	**Low discrimination**					**47353140**
					SPO	
Clostridium barati		40.93	0.47	MAN 8	99	
Bifidobacterium breve		27.70	0.57	SOR 8	1	
Collinsella aerofaciens		17.92	0.32		1	
47353141	**Low discrimination**					**47353141**
					SPO	
Actinomyces naeslundii		60.30	0.49	ESL 12	1	
Clostridium barati		17.31	0.13	RAF 92	99	
Bifidobacterium breve		11.71	0.23		1	
47353142	**Low discrimination**					**47353142**
					SPO	
Bifidobacterium breve		61.41	0.74	RAF 95	1	
Clostridium barati		26.92	0.56		99	
Lactobacillus acidophilus		6.54	0.47		1	
47353143	**Low discrimination**					**47353143**
					SPO	
Bifidobacterium breve		47.38	0.40	RAF 95	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		24.13	0.49	URE 1	1	
Clostridium barati		20.77	0.21		99	
47353150	**Low discrimination**					**47353150**
					SPO	
Bifidobacterium breve		85.72	0.57	RAF 95	1	
Clostridium tertium		8.89	0.31	BGL 91	99	
Actinomyces israelii		3.40	0.38		1	
47353152	**Very good identification**					**47353152**
					SPO	
Bifidobacterium breve		95.21	0.74	RAF 95	1	
Clostridium tertium		2.93	0.39		99	
Bifidobacterium adolescentis		0.72	0.37		1	
47353153	**Acceptable identification**					**47353153**
					SPO	
Bifidobacterium breve		94.56	0.40	RAF 95	1	
Clostridium tertium		2.91	0.05	URE 1	99	
Actinomyces naeslundii		2.53	0.27		1	
47353160	**Low discrimination**					**47353160**
					SPO	
Clostridium barati		47.65	0.56	MAN 8	99	
Clostridium paraputrificum		29.01	0.47	SOR 8	99	
Bifidobacterium breve		19.11	0.61		1	
47353161	**Low discrimination**					**47353161**
					SPO	
Actinomyces naeslundii		36.18	0.49	ESL 12	1	
Clostridium barati		31.14	0.21	RAF 92	99	
Clostridium paraputrificum		18.96	0.13		99	
47353162	**Low discrimination**					**47353162**
					SPO	
Bifidobacterium breve		52.30	0.78	RAF 95	1	
Clostridium barati		38.69	0.64		99	
Clostridium paraputrificum		3.37	0.40		99	
47353163	**Low discrimination**					**47353163**
					SPO	
Bifidobacterium breve		45.61	0.44	RAF 95	1	
Clostridium barati		33.74	0.30	URE 1	99	
Actinomyces naeslundii		13.06	0.49		1	
47353170	**Low discrimination**					**47353170**
					SPO	
Bifidobacterium breve		82.10	0.61	RAF 95	1	
Clostridium tertium		14.37	0.39	BGL 91	99	
Clostridium barati		2.07	0.21		99	
47353172	**Good identification**					**47353172**
					SPO	
Bifidobacterium breve		94.24	0.78	RAF 95	1	
Clostridium tertium		4.89	0.47		99	
Clostridium barati		0.70	0.30		99	
47353173	**Acceptable identification**					**47353173**
					SPO	
Bifidobacterium breve		93.72	0.44	RAF 95	1	
Clostridium tertium		4.87	0.13	URE 1	99	
Actinomyces naeslundii		1.41	0.27		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47353200	**Low discrimination**					**47353200**
					SPO	
Clostridium barati		77.63	0.39	MAN 8	99	
Lactobacillus catenaformis		9.34	0.29	RHA 8	1	
Clostridium innocuum		5.29	0.22		99	
47353202	**Low discrimination**					**47353202**
					SPO	
Clostridium barati		88.30	0.47	MAN 8	99	
Lactobacillus catenaformis		3.54	0.29	RHA 8	1	
Lactobacillus acidophilus		3.41	0.25		1	
47353220	**Acceptable identification**					**47353220**
					SPO	
Clostridium barati		93.23	0.47	MAN 8	99	
Clostridium ramosum		4.91	0.27	RHA 8	92	
Lactobacillus catenaformis		0.94	0.18		1	
47353222	**Acceptable identification**					**47353222**
					SPO	
Clostridium barati		91.10	0.56	MAN 8	99	
Clostridium ramosum		6.39	0.37	RHA 8	92	
Lactobacillus acidophilus		1.17	0.25		1	
47353262	**Good identification**					**47353262**
					SPO	
Clostridium barati		96.98	0.37	MAN 8	99	
Lactobacillus acidophilus		1.77	0.10	RHA 8	1	
Clostridium ramosum		0.79	0.03	SOR 8	92	
47353300	**Low discrimination**					**47353300**
					SPO	
Clostridium barati		60.94	0.47	MAN 8	99	
Collinsella aerofaciens		26.68	0.32	RHA 8	1	
Lactobacillus catenaformis		3.67	0.32		1	
47353302	**Low discrimination**					**47353302**
					SPO	
Clostridium barati		74.46	0.56	MAN 8	99	
Lactobacillus acidophilus		11.01	0.44	RHA 8	1	
Clostridium ramosum		5.23	0.37		92	
47353312	**Low discrimination**					**47353312**
					SPO	
Clostridium tertium		63.43	0.39	TRE 90	99	
Bifidobacterium breve		20.82	0.40	RHA 1	1	
Clostridium barati		9.13	0.21		99	
47353320	**Low discrimination**					**47353320**
Clostridium ramosum		60.04	0.61	TRE 94		
Clostridium barati		34.57	0.56			
Clostridium paraputrificum		2.83	0.32			
47353322	**Low discrimination**					**47353322**
					SPO	
Clostridium ramosum		66.62	0.71	TRE 94	92	
Clostridium barati		28.77	0.64		99	
Lactobacillus acidophilus		1.42	0.44		1	
47353323	**Low discrimination**					**47353323**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		66.25	0.37	TRE 94	92	
Clostridium barati		28.61	0.30	URE 1	99	
Lactobacillus acidophilus		1.41	0.09		1	
47353330	**Low discrimination**					**47353330**
Clostridium tertium		68.34	0.39	TRE 90		
Clostridium ramosum		17.08	0.27	RHA 1		
Clostridium barati		9.84	0.21			
47353332	**Low discrimination**					**47353332**
					SPO	
Clostridium tertium		59.29	0.47	TRE 90	99	
Clostridium ramosum		19.76	0.37	RHA 1	92	
Bifidobacterium breve		11.53	0.44		1	
47353342	**Low discrimination**					**47353342**
					SPO	
Clostridium barati		63.87	0.37	MAN 8	99	
Bifidobacterium breve		16.93	0.40	RHA 8	1	
Lactobacillus acidophilus		13.43	0.28	SOR 8	1	
47353352	**Low discrimination**					**47353352**
					SPO	
Bifidobacterium breve		89.87	0.40	RAF 95	1	
Actinomyces israelii		3.94	0.22	RHA 1	1	
Clostridium barati		3.43	0.03		99	
47353360	**Low discrimination**					**47353360**
Clostridium barati		75.55	0.37	MAN 8		
Clostridium ramosum		15.24	0.27	RHA 8		
Clostridium paraputrificum		5.34	0.13	SOR 8		
47353362	**Low discrimination**					**47353362**
					SPO	
Clostridium barati		65.90	0.46	MAN 8	99	
Clostridium ramosum		17.73	0.37	RHA 8	92	
Bifidobacterium breve		10.35	0.44	SOR 8	1	
47353372	**Low discrimination**					**47353372**
					SPO	
Bifidobacterium breve		88.22	0.44	RAF 95	1	
Clostridium barati		5.68	0.11	RHA 1	99	
Clostridium tertium		4.58	0.13		99	
47353400	**Low discrimination**					**47353400**
					SPO	
Clostridium innocuum		66.72	0.43	MLT 1	99	
Actinomyces israelii		15.89	0.47		1	
Lactobacillus catenaformis		6.75	0.29		1	
47353402	**Low discrimination**					**47353402**
					SPO	
Clostridium innocuum		53.78	0.43	MLT 1	99	
Actinomyces israelii		19.21	0.50		1	
Clostridium barati		15.76	0.31		99	
47353422	**Low discrimination**					**47353422**
					SPO	
Clostridium barati		89.91	0.40	MAN 8	99	
Lactobacillus catenaformis		2.59	0.18	ARA 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		2.33	0.30		1	
47353440	**Low discrimination**					**47353440**
					SPO	
Actinomyces israelii		83.28	0.40	XYL 92	1	
Clostridium innocuum		9.09	0.09		99	
Clostridium barati		7.64	0.05		99	
47353442	**Low discrimination**					**47353442**
					SPO	
Actinomyces israelii		76.46	0.43	XYL 92	1	
Clostridium barati		14.03	0.13		99	
Clostridium innocuum		5.56	0.09		99	
47353502	**Low discrimination**					**47353502**
					SPO	
Bifidobacterium adolescentis		63.94	0.65	RAF 92	1	
Actinomyces israelii		15.87	0.61	XYL 87	1	
Clostridium barati		9.76	0.40		99	
47353512	**Low discrimination**					**47353512**
					SPO	
Bifidobacterium adolescentis		88.01	0.66	RAF 92	1	
Clostridium tertium		6.64	0.39	XYL 87	99	
Actinomyces israelii		2.67	0.47		1	
47353520	**Low discrimination**					**47353520**
Clostridium barati		40.76	0.40	MAN 8		
Clostridium paraputrificum		28.67	0.32	ARA 1		
Clostridium tertium		13.85	0.35			
47353522	**Low discrimination**					**47353522**
Clostridium barati		53.97	0.48	MAN 8		
Clostridium tertium		18.34	0.43	ARA 1		
Clostridium ramosum		8.20	0.35			
47353530	**Acceptable identification**					**47353530**
					SPO	
Clostridium tertium		90.03	0.39	TRE 90	99	
Bifidobacterium breve		5.20	0.27	ARA 1	1	
Actinomyces israelii		1.54	0.23		1	
47353532	**Low discrimination**					**47353532**
					SPO	
Clostridium tertium		77.29	0.47	TRE 90	99	
Bifidobacterium breve		15.04	0.44	ARA 1	1	
Bifidobacterium adolescentis		3.45	0.32		1	
47353540	**Low discrimination**					**47353540**
					SPO	
Actinomyces israelii		84.90	0.51	XYL 92	1	
Clostridium barati		5.84	0.13		99	
Bifidobacterium breve		3.95	0.23		1	
47353542	**Low discrimination**					**47353542**
Bifidobacterium adolescentis		46.22	0.49	RAF 92		
Actinomyces israelii		36.09	0.54	XYL 87		
Bifidobacterium breve		11.33	0.40	SOR 11		
47353550	**Low discrimination**					**47353550**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		64.63	0.37	XYL 92			
Bifidobacterium breve		20.14	0.23	MLZ 13			
Bifidobacterium adolescentis		10.26	0.17				
47353552	**Low discrimination**						**47353552**
Bifidobacterium adolescentis		76.80	0.51	RAF 92			
Bifidobacterium breve		15.40	0.40	XYL 87			
Actinomyces israelii		7.33	0.40	SOR 11			
47353562	**Low discrimination**						**47353562**
					SPO		
Bifidobacterium breve		48.93	0.44	RAF 95	1		
Clostridium barati		36.20	0.30	ARA 1	99		
Actinomyces israelii		5.60	0.33		1		
47353572	**Acceptable identification**						**47353572**
					SPO		
Bifidobacterium breve		91.13	0.44	RAF 95	1		
Clostridium tertium		4.73	0.13	ARA 1	99		
Bifidobacterium adolescentis		2.58	0.17		1		
47353700	**Low discrimination**						**47353700**
					SPO		
Actinomyces israelii		66.18	0.38	XYL 92	1		
Clostridium barati		23.52	0.13	RHA 7	99		
Collinsella aerofaciens		10.30	-.03		1		
47353702	**Low discrimination**						**47353702**
					SPO		
Actinomyces israelii		41.08	0.41	XYL 92	1		
Clostridium barati		29.19	0.21	RHA 7	99		
Bifidobacterium adolescentis		22.21	0.31		1		
47353722	**Low discrimination**						**47353722**
					SPO		
Clostridium ramosum		66.11	0.37	TRE 94	92		
Clostridium barati		28.55	0.30	ARA 1	99		
Lactobacillus acidophilus		1.41	0.09		1		
47354000	**Low discrimination**						**47354000**
					GRA	SPO	
Fusobacterium mortiferum		43.91	0.43	MLT 11	1	1	
Lactobacillus catenaformis		24.87	0.29	XYL 7	99	1	
Clostridium barati		14.53	0.19		99	99	
47354022	**Acceptable identification**						**47354022**
					GRA	SPO	
Clostridium barati		90.32	0.36	GAL 95	99	99	
Lactobacillus catenaformis		4.29	0.18	XYL 1	99	1	
Prevotella oralis		3.07	0.12		1	1	
47354102	**Low discrimination**						**47354102**
					SPO		
Clostridium barati		49.91	0.36	GAL 95	99		
Lactobacillus catenaformis		14.24	0.32	XYL 1	1		
Collinsella aerofaciens		11.51	0.15		1		
47354122	**Low discrimination**						**47354122**
					GRA	SPO	
Clostridium barati		70.19	0.44	GAL 95	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		12.46	0.32	XYL 1	1	1	
Clostridium paraputrificum		11.65	0.25		99	99	
47354312	**Good identification**						**47354312**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47354332	**Good identification**						**47354332**
Clostridium clostridioforme		98.39	0.36	RAF 94			
Parabacteroides distasonis		1.61	0.11	TRE 88			
47354502	**Low discrimination**						**47354502**
					SPO		
Bifidobacterium adolescentis		71.89	0.46	RAF 92	1		
Clostridium clostridioforme		13.26	0.31	GAL 99	66		
Clostridium butyricum		9.61	0.17		99		
47354512	**Low discrimination**						**47354512**
Bifidobacterium adolescentis		58.80	0.48	RAF 92			
Clostridium clostridioforme		40.42	0.42	GAL 99			
Actinomyces israelii		0.44	0.18				
47354532	**Good identification**						**47354532**
Clostridium clostridioforme		98.55	0.42	RAF 94			
Bifidobacterium adolescentis		1.45	0.14	TRE 88			
47354602	**Good identification**						**47354602**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47354612	**Good identification**						**47354612**
Clostridium clostridioforme		100.0	0.48	RAF 94			
47354622	**Good identification**						**47354622**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47354632	**Good identification**						**47354632**
Clostridium clostridioforme		100.0	0.48	RAF 94			
47354702	**Good identification**						**47354702**
Clostridium clostridioforme		96.15	0.45	RAF 94			
Eubacterium contortum		3.13	0.14	TRE 88			
Bifidobacterium adolescentis		0.72	0.12				
47354712	**Very good identification**						**47354712**
Clostridium clostridioforme		99.80	0.57	RAF 94			
Bifidobacterium adolescentis		0.20	0.14	TRE 88			
47354722	**Acceptable identification**						**47354722**
					GRA		
Clostridium clostridioforme		93.48	0.45	RAF 94	50		
Bacteroides uniformis		5.48	0.11	TRE 88	1		
Eubacterium contortum		0.54	0.01		99		
47354732	**Very good identification**						**47354732**
Clostridium clostridioforme		100.0	0.57	RAF 94			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47355000	**Low discrimination**						**47355000**
					SPO		
Lactobacillus catenaformis		54.41	0.63	XYL 1	1		
Lactobacillus jensenii		27.32	0.58		1		
Clostridium barati		6.10	0.41		99		
47355002	**Low discrimination**						**47355002**
					SPO		
Lactobacillus catenaformis		54.80	0.63	XYL 1	1		
Clostridium barati		18.44	0.50		99		
Lactobacillus jensenii		18.34	0.55		1		
47355010	**Low discrimination**						**47355010**
Lactobacillus jensenii		88.35	0.50	XYL 1			
Lactobacillus catenaformis		5.33	0.29				
Actinomyces israelii		4.65	0.40				
47355012	**Low discrimination**						**47355012**
Lactobacillus jensenii		73.85	0.47	XYL 1			
Actinomyces israelii		8.75	0.43				
Lactobacillus catenaformis		6.69	0.29				
47355020	**Low discrimination**						**47355020**
					SPO		
Clostridium barati		41.83	0.50	XYL 1	99		
Lactobacillus catenaformis		31.08	0.53		1		
Lactobacillus jensenii		15.61	0.47		1		
47355022	**Low discrimination**						**47355022**
					GRA	SPO	
Clostridium barati		63.66	0.58	XYL 1	99	99	
Lactobacillus catenaformis		15.77	0.53		99	1	
Prevotella oralis		11.27	0.46		1	1	
47355030	**Low discrimination**						**47355030**
					SPO		
Lactobacillus jensenii		84.20	0.39	XYL 1	1		
Clostridium barati		6.84	0.15		99		
Lactobacillus catenaformis		5.08	0.18		1		
47355032	**Low discrimination**						**47355032**
					SPO		
Lactobacillus jensenii		57.40	0.36	XYL 1	1		
Clostridium barati		20.98	0.24		99		
Lactobacillus catenaformis		5.20	0.18		1		
47355040	**Low discrimination**						**47355040**
					SPO		
Actinomyces israelii		45.97	0.47		1		
Lactobacillus catenaformis		20.25	0.29		1		
Clostridium barati		19.55	0.23		99		
47355042	**Low discrimination**						**47355042**
					SPO		
Actinomyces israelii		42.30	0.50		1		
Clostridium barati		35.97	0.31		99		
Lactobacillus catenaformis		12.42	0.29		1		
47355052	**Low discrimination**						**47355052**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		67.33	0.36	MLZ 13			
Lactobacillus jensenii		14.76	0.12				
Bifidobacterium adolescentis		14.04	0.18				
47355062	**Low discrimination**						**47355062**
					SPO		
Clostridium barati		87.33	0.40	XYL 1	99		
Lactobacillus acidophilus		3.05	0.17	SOR 8	1		
Lactobacillus catenaformis		2.51	0.18		1		
47355100	**Low discrimination**						**47355100**
					COC		
Collinsella aerofaciens		71.86	0.66	XYL 1	1		
Atopobium parvulum		12.08	0.66		99		
Lactobacillus catenaformis		9.88	0.66		1		
47355102	**Low discrimination**						**47355102**
					SPO		
Lactobacillus catenaformis		25.83	0.66	XYL 1	1		
Collinsella aerofaciens		20.88	0.50		1		
Clostridium barati		17.38	0.58		99		
47355112	**Low discrimination**						**47355112**
					GRA	SPO	
Bifidobacterium adolescentis		75.01	0.68	RAF 92	99	1	
Leptotrichia buccalis		7.30	0.32	ARA 87	1	1	
Clostridium tertium		6.40	0.41		99	99	
47355120	**Low discrimination**						**47355120**
					SPO		
Clostridium paraputrificum		65.69	0.66	XYL 1	99		
Collinsella aerofaciens		13.04	0.50		1		
Clostridium barati		10.85	0.58		99		
47355122	**Low discrimination**						**47355122**
					GRA	SPO	
Clostridium barati		30.98	0.66	XYL 1	99	99	
Prevotella oralis		28.66	0.66		1	1	
Clostridium paraputrificum		26.80	0.60		99	99	
47355130	**Low discrimination**						**47355130**
					SPO		
Clostridium tertium		42.45	0.41	TRE 90	99		
Clostridium paraputrificum		35.78	0.32	MAN 99	99		
Collinsella aerofaciens		7.10	0.15		1		
47355132	**Low discrimination**						**47355132**
					SPO		
Clostridium tertium		56.66	0.50	TRE 90	99		
Bifidobacterium breve		7.98	0.44	MAN 99	1		
Clostridium barati		7.89	0.32		99		
47355142	**Low discrimination**						**47355142**
					SPO		
Actinomyces israelii		33.85	0.60		1		
Clostridium barati		21.59	0.40		99		
Bifidobacterium adolescentis		21.36	0.51		1		
47355150	**Low discrimination**						**47355150**
Actinomyces israelii		72.23	0.43	MLZ 13			
Bifidobacterium breve		9.07	0.23				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Collinsella aerofaciens		5.87	-.03				
47355152	**Low discrimination**						**47355152**
Bifidobacterium adolescentis		72.08	0.52	RAF 92			
Actinomyces israelii		13.96	0.46	ARA 87			
Bifidobacterium breve		11.82	0.40	SOR 11			
47355160	**Low discrimination**						**47355160**
					SPO		
Clostridium paraputrificum		79.30	0.47	XYL 1	99		
Clostridium barati		15.13	0.40	SOR 7	99		
Collinsella aerofaciens		2.11	0.15		1		
47355162	**Low discrimination**						**47355162**
					SPO		
Clostridium barati		45.75	0.48	XYL 1	99		
Clostridium paraputrificum		34.25	0.40	SOR 8	99		
Lactobacillus acidophilus		6.12	0.35		1		
47355172	**Low discrimination**						**47355172**
					SPO		
Bifidobacterium breve		75.77	0.44	RAF 95	1		
Clostridium barati		6.51	0.14	XYL 1	99		
Clostridium tertium		5.43	0.15		99		
47355212	**Low discrimination**						**47355212**
Clostridium clostridioforme		87.38	0.35	RAF 94			
Lactobacillus jensenii		5.88	0.12	TRE 88			
Actinomyces israelii		5.19	0.24	ARA 94			
47355222	**Low discrimination**						**47355222**
					GRA	SPO	
Clostridium barati		49.96	0.40	XYL 1	99	99	
Prevotella oralis		43.61	0.40	RHA 8	1	1	
Clostridium clostridioforme		2.09	0.24		50	66	
47355232	**Low discrimination**						**47355232**
					GRA	SPO	
Clostridium clostridioforme		88.16	0.35	RAF 94	50	66	
Clostridium barati		4.67	0.05	TRE 88	99	99	
Prevotella oralis		4.08	0.05	ARA 94	1	1	
47355300	**Low discrimination**						**47355300**
					SPO		
Eubacterium contortum		87.71	0.63	CEL 1	1		
Collinsella aerofaciens		6.74	0.32		1		
Clostridium barati		1.79	0.31		99		
47355302	**Low discrimination**						**47355302**
					GRA	SPO	
Eubacterium contortum		49.94	0.50	CEL 1	99	1	
Clostridium barati		17.31	0.40		99	99	
Prevotella oralis		12.46	0.38		1	1	
47355312	**Low discrimination**						**47355312**
Clostridium clostridioforme		75.06	0.44	RAF 94			
Bifidobacterium adolescentis		11.62	0.34	TRE 88			
Actinomyces israelii		5.33	0.34	ARA 94			
47355320	**Low discrimination**						**47355320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Eubacterium contortum		49.89	0.50	CEL 1	99	1	
Clostridium barati		17.29	0.40		99	99	
Prevotella oralis		12.45	0.38		1	1	
47355322	**Low discrimination**						**47355322**
					GRA	SPO	
Prevotella oralis		71.53	0.60	RAF 92	1	1	
Clostridium barati		15.69	0.48	XYL 11	99	99	
Parabacteroides distasonis		3.35	0.50		1	1	
47355332	**Low discrimination**						**47355332**
Clostridium clostridioforme		53.26	0.44	RAF 94			
Parabacteroides distasonis		28.80	0.45	TRE 88			
Prevotella oralis		11.52	0.26	ARA 94			
47355342	**Low discrimination**						**47355342**
					SPO		
Actinomyces israelii		41.91	0.41	RHA 7	1		
Clostridium barati		30.88	0.21		99		
Lactobacillus acidophilus		10.73	0.16		1		
47355402	**Low discrimination**						**47355402**
					SPO		
Actinomyces israelii		42.32	0.56		1		
Bifidobacterium adolescentis		27.77	0.46		1		
Clostridium butyricum		15.31	0.27		99		
47355410	**Low discrimination**						**47355410**
					SPO		
Actinomyces israelii		53.84	0.39	MLZ 13	1		
Clostridium butyricum		26.47	0.12		99		
Lactobacillus jensenii		12.62	0.15		1		
47355412	**Low discrimination**						**47355412**
Bifidobacterium adolescentis		54.18	0.48	RAF 92			
Clostridium clostridioforme		33.38	0.41	LAC 99			
Actinomyces israelii		10.10	0.42				
47355432	**Good identification**						**47355432**
Clostridium clostridioforme		95.82	0.41	RAF 94			
Actinomyces israelii		1.85	0.21	TRE 88			
Bifidobacterium adolescentis		1.57	0.14	RHA 88			
47355440	**Low discrimination**						**47355440**
					SPO		
Actinomyces israelii		76.21	0.46		1		
Clostridium butyricum		23.79	0.16		99		
47355442	**Low discrimination**						**47355442**
					SPO		
Actinomyces israelii		76.93	0.49		1		
Bifidobacterium adolescentis		16.04	0.31		1		
Clostridium butyricum		6.22	0.09		99		
47355500	**Low discrimination**						**47355500**
					SPO		
Clostridium butyricum		80.10	0.58	RAF 95	99		
Actinomyces israelii		9.57	0.63	TRE 94	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Collinsella aerofaciens		5.46	0.32		1		
47355502	**Low discrimination**						**47355502**
					SPO		
Bifidobacterium adolescentis		82.52	0.80	RAF 92	1		
Clostridium butyricum		11.03	0.51		99		
Actinomyces israelii		5.08	0.66		1		
47355503	**Low discrimination**						**47355503**
					SPO		
Bifidobacterium adolescentis		83.28	0.46	RAF 92	1		
Clostridium butyricum		11.13	0.17	URE 1	99		
Actinomyces israelii		5.13	0.32		1		
47355510	**Low discrimination**						**47355510**
					SPO		
Clostridium butyricum		48.74	0.36	RAF 95	99		
Bifidobacterium adolescentis		33.26	0.48	TRE 94	1		
Actinomyces israelii		16.53	0.49	MLZ 5	1		
47355512	**Very good identification**						**47355512**
Bifidobacterium adolescentis		96.64	0.82	RAF 92			
Clostridium clostridioforme		2.01	0.50				
Actinomyces israelii		0.73	0.52				
47355513	**Good identification**						**47355513**
Bifidobacterium adolescentis		96.70	0.48	RAF 92			
Clostridium clostridioforme		2.02	0.16	URE 1			
Actinomyces israelii		0.73	0.18				
47355520	**Low discrimination**						**47355520**
					GRA	SPO	
Clostridium butyricum		50.77	0.40	RAF 95	99	99	
Clostridium paraputrificum		22.30	0.32	TRE 94	99	99	
Bacteroides vulgatus		9.12	0.26	NAG 8	1	1	
47355522	**Low discrimination**						**47355522**
					GRA	SPO	
Bacteroides uniformis		50.98	0.43	IND 98	1	1	
Clostridium butyricum		10.33	0.33	RAF 97	99	99	
Bifidobacterium adolescentis		8.97	0.46		99	1	
47355532	**Low discrimination**						**47355532**
					GRA		
Clostridium clostridioforme		59.74	0.50	RAF 94	50		
Bifidobacterium adolescentis		28.96	0.48	TRE 88	99		
Parabacteroides distasonis		3.09	0.34	RHA 88	1		
47355540	**Low discrimination**						**47355540**
					SPO		
Clostridium butyricum		63.06	0.40	RAF 95	99		
Actinomyces israelii		33.68	0.56	TRE 94	1		
Bifidobacterium adolescentis		2.63	0.31	SOR 8	1		
47355542	**Low discrimination**						**47355542**
					SPO		
Bifidobacterium adolescentis		77.50	0.65	RAF 92	1		
Actinomyces israelii		15.01	0.59	SOR 11	1		
Clostridium butyricum		7.29	0.33		99		
47355550	**Low discrimination**						**47355550**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		42.79	0.42	MLZ 13	1		
Clostridium butyricum		28.22	0.18		99		
Bifidobacterium adolescentis		27.37	0.32		1		
47355552	**Very good identification**						**47355552**
					SPO		
Bifidobacterium adolescentis		97.14	0.66	RAF 92	1		
Actinomyces israelii		2.30	0.45	SOR 11	1		
Clostridium butyricum		0.39	0.11		99		
47355562	**Low discrimination**						**47355562**
					SPO		
Actinomyces israelii		27.22	0.38	NAG 6	1		
Bifidobacterium adolescentis		22.23	0.31		1		
Clostridium butyricum		18.00	0.14		99		
47355602	**Low discrimination**						**47355602**
Clostridium clostridioforme		86.32	0.45	RAF 94			
Actinomyces israelii		8.17	0.36	TRE 88			
Eubacterium contortum		3.14	0.14				
47355612	**Good identification**						**47355612**
Clostridium clostridioforme		99.47	0.56	RAF 94			
Actinomyces israelii		0.31	0.22	TRE 88			
Bifidobacterium adolescentis		0.22	0.14				
47355622	**Low discrimination**						**47355622**
					GRA	SPO	
Clostridium clostridioforme		84.79	0.45	RAF 94	50	66	
Bacteroides uniformis		11.20	0.16	TRE 88	1	1	
Clostridium barati		1.31	0.05		99	99	
47355632	**Good identification**						**47355632**
Clostridium clostridioforme		100.0	0.56	RAF 94			
47355700	**Good identification**						**47355700**
					SPO		
Eubacterium contortum		94.32	0.61	CEL 1	1		
Clostridium butyricum		4.02	0.32		99		
Actinomyces israelii		1.17	0.44		1		
47355702	**Low discrimination**						**47355702**
Eubacterium contortum		41.92	0.48	CEL 1			
Clostridium clostridioforme		38.99	0.54				
Bifidobacterium adolescentis		9.61	0.46				
47355712	**Good identification**						**47355712**
Clostridium clostridioforme		93.16	0.65	RAF 94			
Bifidobacterium adolescentis		6.16	0.48	TRE 88			
Actinomyces israelii		0.35	0.33				
47355720	**Low discrimination**						**47355720**
					GRA	SPO	
Bacteroides vulgatus		50.73	0.42	RAF 97	1	1	
Eubacterium contortum		46.26	0.48	CEL 5	99	1	
Clostridium butyricum		0.97	0.14		99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47355722	**Low discrimination**					**47355722**
					GRA	
Bacteroides uniformis		55.59	0.45	IND 98	1	
Clostridium clostridioforme		28.76	0.54	RAF 97	50	
Eubacterium contortum		5.46	0.36		99	
47355732	**Very good identification**					**47355732**
Clostridium clostridioforme		96.53	0.65	RAF 94		
Parabacteroides distasonis		2.04	0.42	TRE 88		
Bacteroides ovatus		0.91	0.20			
47355742	**Low discrimination**					**47355742**
Actinomyces israelii		42.21	0.40	RHA 7		
Bifidobacterium adolescentis		29.24	0.31			
Eubacterium contortum		10.43	0.14			
47356002	**Low discrimination**					**47356002**
					SPO	
Actinomyces israelii		58.97	0.36	GAL 99	1	
Clostridium innocuum		18.26	0.12		99	
Clostridium barati		17.93	0.10		99	
47356102	**Low discrimination**					**47356102**
					SPO	
Clostridium tertium		57.31	0.37	TRE 90	99	
Actinomyces israelii		28.18	0.46	GAL 99	1	
Bifidobacterium adolescentis		8.08	0.31		1	
47356112	**Low discrimination**					**47356112**
					SPO	
Clostridium tertium		87.86	0.41	TRE 90	99	
Bifidobacterium adolescentis		8.51	0.32	GAL 99	1	
Actinomyces israelii		3.63	0.32		1	
47356120	**Low discrimination**					**47356120**
					SPO	
Clostridium tertium		87.09	0.37	TRE 90	99	
Clostridium barati		9.77	0.18	GAL 99	99	
Actinomyces israelii		1.82	0.23		1	
47356122	**Low discrimination**					**47356122**
					SPO	
Clostridium tertium		88.27	0.45	TRE 90	99	
Clostridium barati		9.90	0.26	GAL 99	99	
Actinomyces israelii		0.92	0.26		1	
47356130	**Good identification**					**47356130**
Clostridium tertium		100.0	0.41	TRE 90		
47356132	**Good identification**					**47356132**
					SPO	
Clostridium tertium		99.86	0.50	TRE 90	99	
Bifidobacterium breve		0.14	0.10	GAL 99	1	
47356140	**Good identification**					**47356140**
Actinomyces israelii		100.0	0.36	GAL 99		
47356142	**Low discrimination**					**47356142**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		83.69	0.39	GAL 99	1	
Bifidobacterium adolescentis		7.63	0.15		1	
Clostridium tertium		4.42	0.03		99	
47356500	**Low discrimination**					**47356500**
					SPO	
Actinomyces israelii		83.35	0.42	GAL 99	1	
Clostridium butyricum		13.69	0.07		99	
Bifidobacterium adolescentis		2.96	0.11		1	
47356502	**Low discrimination**					**47356502**
					SPO	
Bifidobacterium adolescentis		68.71	0.45	RAF 92	1	
Actinomyces israelii		29.30	0.45	GAL 99	1	
Clostridium butyricum		1.25	0.00		99	
47356512	**Low discrimination**					**47356512**
Bifidobacterium adolescentis		89.00	0.46	RAF 92		
Actinomyces israelii		4.64	0.31	GAL 99		
Bifidobacterium dentium		4.22	0.06			
47356542	**Low discrimination**					**47356542**
Actinomyces israelii		57.30	0.38	GAL 99		
Bifidobacterium adolescentis		42.70	0.29			
47357000	**Low discrimination**					**47357000**
					SPO	
Actinomyces israelii		82.55	0.67		1	
Clostridium innocuum		7.36	0.34		99	
Clostridium tertium		3.39	0.29		99	
47357002	**Low discrimination**					**47357002**
					SPO	
Actinomyces israelii		80.54	0.70		1	
Clostridium tertium		6.62	0.37		99	
Clostridium innocuum		4.79	0.34		99	
47357003	**Low discrimination**					**47357003**
					SPO	
Actinomyces israelii		83.34	0.36	URE 1	1	
Clostridium tertium		6.85	0.03		99	
Clostridium innocuum		4.95	0.00		99	
47357010	**Low discrimination**					**47357010**
					SPO	
Actinomyces israelii		64.62	0.53	MLZ 13	1	
Clostridium tertium		31.59	0.33		99	
Lactobacillus jensenii		2.19	0.15		1	
47357012	**Low discrimination**					**47357012**
					SPO	
Actinomyces israelii		47.44	0.56	MLZ 13	1	
Clostridium tertium		46.38	0.41		99	
Bifidobacterium adolescentis		4.50	0.32		1	
47357020	**Low discrimination**					**47357020**
					SPO	
Clostridium tertium		42.69	0.37	TRE 90	99	
Clostridium barati		30.32	0.31	LAC 99	99	
Actinomyces israelii		22.11	0.47		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47357022	**Low discrimination**					**47357022**
					SPO	
Clostridium tertium		49.21	0.45	TRE 90	99	
Clostridium barati		34.95	0.40	LAC 99	99	
Actinomyces israelii		12.74	0.50		1	
47357030	**Acceptable identification**					**47357030**
					SPO	
Clostridium tertium		94.94	0.41	TRE 90	99	
Actinomyces israelii		4.13	0.32	LAC 99	1	
Lactobacillus jensenii		0.55	0.05		1	
47357032	**Good identification**					**47357032**
					SPO	
Clostridium tertium		97.23	0.50	TRE 90	99	
Actinomyces israelii		2.12	0.35	LAC 99	1	
Clostridium barati		0.39	0.05		99	
47357040	**Very good identification**					**47357040**
					SPO	
Actinomyces israelii		99.12	0.60		1	
Clostridium barati		0.65	0.05		99	
Clostridium innocuum		0.23	0.00		99	
47357042	**Very good identification**					**47357042**
					SPO	
Actinomyces israelii		98.00	0.63		1	
Clostridium barati		1.28	0.13		99	
Bifidobacterium adolescentis		0.36	0.15		1	
47357050	**Good identification**					**47357050**
					SPO	
Actinomyces israelii		98.75	0.46	MLZ 13	1	
Clostridium tertium		1.25	-.01		99	
47357052	**Acceptable identification**					**47357052**
					SPO	
Actinomyces israelii		94.74	0.49	MLZ 13	1	
Bifidobacterium adolescentis		2.85	0.17		1	
Clostridium tertium		2.41	0.07		99	
47357060	**Low discrimination**					**47357060**
					SPO	
Actinomyces israelii		73.70	0.40	NAG 6	1	
Clostridium barati		22.60	0.13		99	
Clostridium tertium		3.70	0.03		99	
47357062	**Low discrimination**					**47357062**
					SPO	
Actinomyces israelii		57.51	0.43	NAG 6	1	
Clostridium barati		35.27	0.21		99	
Clostridium tertium		5.77	0.11		99	
47357100	**Low discrimination**					**47357100**
					SPO	
Clostridium tertium		46.90	0.63	TRE 90	99	
Actinomyces israelii		46.12	0.77		1	
Collinsella aerofaciens		3.80	0.32		1	
47357102	**Low discrimination**					**47357102**
					SPO	
Clostridium tertium		59.79	0.71	TRE 90	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		29.40	0.80		1	
Bifidobacterium adolescentis		8.43	0.65		1	
47357103	**Low discrimination**					**47357103**
					SPO	
Clostridium tertium		60.01	0.37	TRE 90	99	
Actinomyces israelii		29.50	0.46	URE 1	1	
Bifidobacterium adolescentis		8.46	0.31		1	
47357110	**Good identification**					**47357110**
					SPO	
Clostridium tertium		91.97	0.67	TRE 90	99	
Actinomyces israelii		7.60	0.63		1	
Bifidobacterium adolescentis		0.27	0.32		1	
47357112	**Low discrimination**					**47357112**
					SPO	
Clostridium tertium		87.62	0.76	TRE 90	99	
Bifidobacterium adolescentis		8.49	0.66		1	
Actinomyces israelii		3.62	0.66		1	
47357113	**Low discrimination**					**47357113**
					SPO	
Clostridium tertium		87.86	0.41	TRE 90	99	
Bifidobacterium adolescentis		8.51	0.32	URE 1	1	
Actinomyces israelii		3.63	0.32		1	
47357120	**Good identification**					**47357120**
					SPO	
Clostridium tertium		93.77	0.71	TRE 90	99	
Clostridium barati		2.02	0.40		99	
Actinomyces israelii		1.96	0.57		1	
47357121	**Acceptable identification**					**47357121**
					SPO	
Clostridium tertium		93.47	0.37	TRE 90	99	
Clostridium barati		2.01	0.05	URE 1	99	
Actinomyces israelii		1.96	0.23		1	
47357122	**Very good identification**					**47357122**
					SPO	
Clostridium tertium		95.72	0.79	TRE 90	99	
Clostridium barati		2.06	0.48		99	
Actinomyces israelii		1.00	0.60		1	
47357123	**Good identification**					**47357123**
					SPO	
Clostridium tertium		95.57	0.45	TRE 90	99	
Clostridium barati		2.06	0.14	URE 1	99	
Actinomyces israelii		1.00	0.26		1	
47357130	**Very good identification**					**47357130**
					SPO	
Clostridium tertium		99.76	0.76	TRE 90	99	
Actinomyces israelii		0.18	0.43		1	
Bifidobacterium breve		0.04	0.27		1	
47357131	**Good identification**					**47357131**
Clostridium tertium		100.0	0.41	TRE 90		
47357132	**Very good identification**					**47357132**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		99.70	0.84	TRE 90	99	
Bifidobacterium breve		0.14	0.44		1	
Actinomyces israelii		0.09	0.46		1	
47357133	**Good identification**					**47357133**
					SPO	
Clostridium tertium		99.86	0.50	TRE 90	99	
Bifidobacterium breve		0.14	0.10	URE 1	1	
47357140	**Very good identification**					**47357140**
					SPO	
Actinomyces israelii		96.02	0.70		1	
Clostridium tertium		2.54	0.29		99	
Clostridium barati		0.47	0.13		99	
47357141	**Good identification**					**47357141**
Actinomyces israelii		100.0	0.36	URE 1		
47357142	**Low discrimination**					**47357142**
					SPO	
Actinomyces israelii		84.80	0.73		1	
Bifidobacterium adolescentis		7.73	0.49		1	
Clostridium tertium		4.48	0.37		99	
47357143	**Low discrimination**					**47357143**
					SPO	
Actinomyces israelii		87.42	0.39	URE 1	1	
Bifidobacterium adolescentis		7.97	0.15		1	
Clostridium tertium		4.62	0.03		99	
47357150	**Low discrimination**					**47357150**
					SPO	
Actinomyces israelii		73.90	0.56	MLZ 13	1	
Clostridium tertium		23.22	0.33		99	
Bifidobacterium breve		1.64	0.23		1	
47357152	**Low discrimination**					**47357152**
					SPO	
Actinomyces israelii		39.63	0.59	MLZ 13	1	
Bifidobacterium adolescentis		29.54	0.51		1	
Clostridium tertium		24.91	0.41		99	
47357160	**Low discrimination**					**47357160**
					SPO	
Clostridium tertium		45.92	0.37	TRE 90	99	
Actinomyces israelii		36.99	0.50	SOR 1	1	
Clostridium barati		8.51	0.21		99	
47357162	**Low discrimination**					**47357162**
					SPO	
Clostridium tertium		53.16	0.45	TRE 90	99	
Actinomyces israelii		21.41	0.53	SOR 1	1	
Bifidobacterium breve		13.31	0.44		1	
47357170	**Acceptable identification**					**47357170**
					SPO	
Clostridium tertium		90.13	0.41	TRE 90	99	
Actinomyces israelii		6.10	0.36	SOR 1	1	
Bifidobacterium breve		3.77	0.27		1	
47357172	**Low discrimination**					**47357172**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium tertium		84.84	0.50	TRE 90	99	
Bifidobacterium breve		11.95	0.44	SOR 1	1	
Actinomyces israelii		2.87	0.39		1	
47357200	**Good identification**					**47357200**
					SPO	
Actinomyces israelii		95.63	0.48	RHA 7	1	
Clostridium barati		3.22	0.05		99	
Clostridium innocuum		1.14	0.00		99	
47357202	**Acceptable identification**					**47357202**
					SPO	
Actinomyces israelii		92.05	0.51	RHA 7	1	
Clostridium barati		6.21	0.13		99	
Clostridium tertium		1.02	0.03		99	
47357212	**Low discrimination**					**47357212**
					SPO	
Actinomyces israelii		88.40	0.37	RHA 7	1	
Clostridium tertium		11.60	0.07	MLZ 13	99	
47357240	**Good identification**					**47357240**
Actinomyces israelii		100.0	0.41	RHA 7		
47357242	**Good identification**					**47357242**
Actinomyces israelii		100.0	0.44	RHA 7		
47357300	**Low discrimination**					**47357300**
					SPO	
Actinomyces israelii		75.59	0.58	RHA 7	1	
Eubacterium contortum		10.89	0.29		1	
Clostridium tertium		10.32	0.29		99	
47357302	**Low discrimination**					**47357302**
					SPO	
Actinomyces israelii		71.64	0.61	RHA 7	1	
Clostridium tertium		19.56	0.37		99	
Clostridium barati		3.62	0.21		99	
47357312	**Low discrimination**					**47357312**
					SPO	
Clostridium tertium		70.79	0.41	TRE 90	99	
Actinomyces israelii		21.80	0.47	RHA 1	1	
Bifidobacterium adolescentis		6.86	0.32		1	
47357320	**Low discrimination**					**47357320**
Clostridium tertium		56.96	0.37	TRE 90		
Clostridium ramosum		18.33	0.27	RHA 1		
Clostridium barati		10.55	0.21			
47357322	**Low discrimination**					**47357322**
Clostridium tertium		55.09	0.45	TRE 90		
Clostridium ramosum		23.64	0.37	RHA 1		
Clostridium barati		10.21	0.30			
47357330	**Good identification**					**47357330**
					SPO	
Clostridium tertium		98.71	0.41	TRE 90	99	
Actinomyces israelii		1.29	0.24	RHA 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47357332	**Good identification**						**47357332**
					SPO		
Clostridium tertium		98.58	0.50	TRE 90	99		
Actinomyces israelii		0.65	0.27	RHA 1	1		
Clostridium clostridioforme		0.25	0.10		66		
47357340	**Good identification**						**47357340**
Actinomyces israelii		100.0	0.51	RHA 7			
47357342	**Good identification**						**47357342**
					SPO		
Actinomyces israelii		97.03	0.54	RHA 7	1		
Bifidobacterium adolescentis		1.19	0.15		1		
Clostridium barati		1.10	0.03		99		
47357350	**Low discrimination**						**47357350**
					COC	SPO	
Actinomyces israelii		89.68	0.37	RHA 7	1	1	
Blautia producta		6.54	-.03	MLZ 13	99	1	
Clostridium tertium		3.78	-.01		1	99	
47357352	**Low discrimination**						**47357352**
					SPO		
Actinomyces israelii		84.44	0.40	RHA 7	1		
Bifidobacterium adolescentis		8.44	0.17	MLZ 13	1		
Clostridium tertium		7.12	0.07		99		
47357400	**Very good identification**						**47357400**
					SPO		
Actinomyces israelii		95.42	0.66		1		
Clostridium butyricum		2.61	0.18		99		
Clostridium innocuum		1.83	0.21		99		
47357402	**Low discrimination**						**47357402**
					SPO		
Actinomyces israelii		89.57	0.69		1		
Bifidobacterium adolescentis		8.49	0.45		1		
Clostridium innocuum		1.15	0.21		99		
47357410	**Good identification**						**47357410**
					SPO		
Actinomyces israelii		97.33	0.51	MLZ 13	1		
Bifidobacterium adolescentis		1.14	0.12		1		
Clostridium butyricum		0.94	-.04		99		
47357412	**Low discrimination**						**47357412**
Actinomyces israelii		54.65	0.54	MLZ 13			
Bifidobacterium adolescentis		42.36	0.46				
Bifidobacterium dentium		2.01	0.06				
47357420	**Acceptable identification**						**47357420**
					SPO		
Actinomyces israelii		92.76	0.45	NAG 6	1		
Clostridium butyricum		3.46	-.01		99		
Clostridium tertium		2.21	0.03		99		
47357422	**Acceptable identification**						**47357422**
					SPO		
Actinomyces israelii		90.40	0.48	NAG 6	1		
Clostridium tertium		4.31	0.11		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		3.06	0.05		99	
47357440	**Very good identification**					**47357440**
					SPO	
Actinomyces israelii		99.39	0.59		1	
Clostridium butyricum		0.61	-.01		99	
47357442	**Very good identification**					**47357442**
					SPO	
Actinomyces israelii		96.93	0.62		1	
Bifidobacterium adolescentis		2.92	0.29		1	
Clostridium butyricum		0.15	-.08		99	
47357450	**Good identification**					**47357450**
					COC	
Actinomyces israelii		97.39	0.44	MLZ 13	1	
Blautia producta		2.61	-.03		99	
47357452	**Low discrimination**					**47357452**
Actinomyces israelii		80.24	0.47	MLZ 13		
Bifidobacterium adolescentis		19.76	0.31			
47357460	**Good identification**					**47357460**
Actinomyces israelii		100.0	0.38	NAG 6		
47357462	**Good identification**					**47357462**
Actinomyces israelii		100.0	0.41	NAG 6		
47357500	**Low discrimination**					**47357500**
					SPO	
Actinomyces israelii		82.40	0.76		1	
Clostridium butyricum		13.53	0.41		99	
Bifidobacterium adolescentis		2.93	0.45		1	
47357501	**Low discrimination**					**47357501**
					SPO	
Actinomyces israelii		83.35	0.42	URE 1	1	
Clostridium butyricum		13.69	0.07		99	
Bifidobacterium adolescentis		2.96	0.11		1	
47357502	**Low discrimination**					**47357502**
					SPO	
Bifidobacterium adolescentis		68.67	0.79	RAF 92	1	
Actinomyces israelii		29.29	0.79		1	
Clostridium butyricum		1.25	0.34		99	
47357503	**Low discrimination**					**47357503**
					SPO	
Bifidobacterium adolescentis		68.71	0.45	RAF 92	1	
Actinomyces israelii		29.30	0.45	URE 1	1	
Clostridium butyricum		1.25	0.00		99	
47357510	**Low discrimination**					**47357510**
					SPO	
Actinomyces israelii		61.15	0.62	MLZ 13	1	
Bifidobacterium adolescentis		17.77	0.46		1	
Clostridium tertium		9.13	0.33		99	
47357512	**Low discrimination**					**47357512**
Bifidobacterium adolescentis		89.59	0.80	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		4.67	0.65				
Bifidobacterium dentium		4.25	0.40				
47357513	**Low discrimination**						**47357513**
Bifidobacterium adolescentis		89.68	0.46	RAF 92			
Actinomyces israelii		4.68	0.31	URE 1			
Bifidobacterium dentium		4.25	0.06				
47357520	**Low discrimination**						**47357520**
					SPO		
Actinomyces israelii		54.04	0.55	NAG 6	1		
Clostridium tertium		31.89	0.37		99		
Clostridium butyricum		12.09	0.23		99		
47357522	**Low discrimination**						**47357522**
					SPO		
Clostridium tertium		43.49	0.45	TRE 90	99		
Actinomyces israelii		36.85	0.58	ARA 1	1		
Bifidobacterium adolescentis		13.68	0.45		1		
47357530	**Low discrimination**						**47357530**
					SPO		
Clostridium tertium		58.98	0.41	TRE 90	99		
Bifidobacterium dentium		31.57	0.27	ARA 1	1		
Actinomyces israelii		8.40	0.41		1		
47357532	**Low discrimination**						**47357532**
					SPO		
Bifidobacterium dentium		58.20	0.44	RAF 95	1		
Clostridium tertium		32.26	0.50	TRE 99	99		
Bifidobacterium adolescentis		6.97	0.46		1		
47357540	**Good identification**						**47357540**
					COC	SPO	
Actinomyces israelii		92.50	0.69		1	1	
Clostridium butyricum		3.40	0.23		1	99	
Blautia producta		3.06	0.24		99	1	
47357542	**Low discrimination**						**47357542**
					SPO		
Actinomyces israelii		56.89	0.72		1		
Bifidobacterium adolescentis		42.40	0.63		1		
Clostridium butyricum		0.54	0.16		99		
47357543	**Low discrimination**						**47357543**
Actinomyces israelii		57.30	0.38	URE 1			
Bifidobacterium adolescentis		42.70	0.29				
47357550	**Low discrimination**						**47357550**
					COC		
Actinomyces israelii		56.40	0.55	MLZ 13	1		
Blautia producta		37.44	0.32		99		
Bifidobacterium adolescentis		5.21	0.31		1		
47357552	**Low discrimination**						**47357552**
					COC		
Bifidobacterium adolescentis		84.90	0.65	RAF 92	1		
Actinomyces israelii		13.93	0.58	SOR 11	1		
Blautia producta		0.69	0.15		99		
47357560	**Low discrimination**						**47357560**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		89.06	0.48	NAG 6	1	1	
Blautia producta		5.13	0.07		99	1	
Clostridium butyricum		4.46	0.05		1	99	
47357562	**Low discrimination**						**47357562**
					SPO		
Actinomyces israelii		85.53	0.51	NAG 6	1		
Bifidobacterium adolescentis		10.09	0.29		1		
Clostridium tertium		2.62	0.11		99		
47357572	**Low discrimination**						**47357572**
Actinomyces israelii		32.01	0.37	MLZ 13			
Bifidobacterium adolescentis		30.88	0.31	NAG 6			
Bifidobacterium dentium		21.06	0.10				
47357600	**Good identification**						**47357600**
Actinomyces israelii		100.0	0.46	RHA 7			
47357602	**Good identification**						**47357602**
Actinomyces israelii		96.94	0.49	RHA 7			
Clostridium clostridioforme		1.83	0.10				
Bifidobacterium adolescentis		1.23	0.11				
47357612	**Low discrimination**						**47357612**
Actinomyces israelii		59.58	0.35	RHA 7			
Clostridium clostridioforme		34.22	0.22	MLZ 13			
Bifidobacterium adolescentis		6.20	0.12				
47357640	**Good identification**						**47357640**
Actinomyces israelii		100.0	0.39	RHA 7			
47357642	**Good identification**						**47357642**
Actinomyces israelii		100.0	0.42	RHA 7			
47357700	**Low discrimination**						**47357700**
					SPO		
Actinomyces israelii		82.22	0.57	RHA 7	1		
Eubacterium contortum		11.84	0.27		1		
Clostridium butyricum		5.55	0.16		99		
47357702	**Low discrimination**						**47357702**
					SPO		
Actinomyces israelii		73.09	0.60	RHA 7	1		
Bifidobacterium adolescentis		23.00	0.45		1		
Clostridium butyricum		1.28	0.09		99		
47357710	**Low discrimination**						**47357710**
					COC		
Actinomyces israelii		89.49	0.42	RHA 7	1		
Bifidobacterium adolescentis		3.49	0.12	MLZ 13	1		
Blautia producta		3.10	-.03		99		
47357712	**Low discrimination**						**47357712**
Bifidobacterium adolescentis		61.04	0.46	RAF 92			
Actinomyces israelii		23.72	0.45	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		11.40	0.31				
47357720	**Low discrimination**						**47357720**
					GRA		
Actinomyces israelii		49.85	0.36	RHA 7	99		
Bacteroides vulgatus		21.77	0.08	NAG 6	1		
Eubacterium contortum		19.85	0.14		99		
47357722	**Low discrimination**						**47357722**
					GRA		
Actinomyces israelii		38.20	0.39	RHA 7	99		
Bacteroides ovatus		20.10	0.15	NAG 6	1		
Bacteroides uniformis		18.24	0.11		1		
47357740	**Low discrimination**						**47357740**
					COC	SPO	
Actinomyces israelii		68.76	0.50	RHA 7	1	1	
Blautia producta		30.20	0.24		99	1	
Clostridium butyricum		1.04	-.03		1	99	
47357742	**Low discrimination**						**47357742**
					COC		
Actinomyces israelii		88.30	0.53	RHA 7	1		
Bifidobacterium adolescentis		8.83	0.29		1		
Blautia producta		2.87	0.07		99		
47357752	**Low discrimination**						**47357752**
					COC		
Actinomyces israelii		40.46	0.38	RHA 7	1		
Bifidobacterium adolescentis		33.10	0.31	MLZ 13	1		
Blautia producta		26.43	0.15		99		
47361000	**Very good identification**						**47361000**
					COC		
Lactobacillus jensenii		97.83	0.66	FRU 99	1		
Lactobacillus catenaformis		0.66	0.29		1		
Atopobium parvulum		0.65	0.27		99		
47361002	**Low discrimination**						**47361002**
					SPO		
Lactobacillus jensenii		88.87	0.63	FRU 99	1		
Lactobacillus acidophilus		7.34	0.41		1		
Clostridium barati		2.59	0.31		99		
47361010	**Very good identification**						**47361010**
					GRA		
Lactobacillus jensenii		99.83	0.58	FRU 99	99		
Leptotrichia buccalis		0.17	-.03		1		
47361012	**Low discrimination**						**47361012**
					GRA		
Lactobacillus jensenii		79.81	0.55	FRU 99	99		
Leptotrichia buccalis		19.92	0.32		1		
Lactobacillus acidophilus		0.20	0.07		99		
47361020	**Low discrimination**						**47361020**
					SPO		
Lactobacillus jensenii		87.72	0.56	FRU 99	1		
Clostridium barati		6.82	0.31		99		
Clostridium paraputrificum		3.82	0.22		99		
47361022	**Low discrimination**						**47361022**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		57.77	0.53	FRU 99	1		
Clostridium barati		20.20	0.40		99		
Lactobacillus acidophilus		19.09	0.41		1		
47361030	**Good identification**						**47361030**
					SPO		
Lactobacillus jensenii		99.77	0.47	FRU 99	1		
Clostridium barati		0.24	-.03		99		
47361032	**Low discrimination**						**47361032**
					GRA	SPO	
Lactobacillus jensenii		88.38	0.44	FRU 99	99	1	
Leptotrichia buccalis		9.80	0.15		1	1	
Clostridium barati		0.94	0.05		99	99	
47361100	**Low discrimination**						**47361100**
					COC	SPO	
Atopobium parvulum		82.79	0.61	FRU 99	99	1	
Collinsella aerofaciens		9.24	0.32		1	1	
Clostridium barati		2.45	0.31		1	99	
47361102	**Low discrimination**						**47361102**
					COC	SPO	
Lactobacillus acidophilus		69.90	0.60	FRU 99	1	1	
Atopobium parvulum		12.78	0.48		99	1	
Clostridium barati		6.43	0.40		1	99	
47361112	**Very good identification**						**47361112**
					GRA		
Leptotrichia buccalis		99.60	0.66	FRU 99	1		
Bifidobacterium breve		0.14	0.37		99		
Lactobacillus acidophilus		0.12	0.25		99		
47361120	**Low discrimination**						**47361120**
					COC	SPO	
Clostridium paraputrificum		81.34	0.56	FRU 99	1	99	
Atopobium parvulum		8.73	0.48		99	1	
Clostridium barati		4.40	0.40		1	99	
47361122	**Low discrimination**						**47361122**
					SPO		
Lactobacillus acidophilus		45.00	0.60	FRU 99	1		
Clostridium paraputrificum		32.85	0.50		99		
Clostridium barati		12.42	0.48		99		
47361132	**Acceptable identification**						**47361132**
					GRA	SPO	
Leptotrichia buccalis		94.19	0.50	FRU 99	1	1	
Bifidobacterium breve		2.12	0.41	NAG 10	99	1	
Clostridium tertium		1.48	0.29		99	99	
47361142	**Low discrimination**						**47361142**
					SPO		
Lactobacillus acidophilus		83.59	0.44	FRU 99	1		
Bifidobacterium breve		8.23	0.37	SOR 11	1		
Clostridium barati		5.41	0.21		99		
47361160	**Low discrimination**						**47361160**
					SPO		
Clostridium paraputrificum		89.42	0.37	FRU 99	99		
Clostridium barati		5.58	0.21	SOR 7	99		
Bifidobacterium breve		1.49	0.24		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47361162	**Low discrimination**						**47361162**
					SPO		
Lactobacillus acidophilus		54.59	0.44	FRU 99	1		
Clostridium paraputrificum		24.27	0.31	SOR 11	99		
Clostridium barati		10.60	0.30		99		
47361172	**Low discrimination**						**47361172**
					GRA		
Bifidobacterium breve		65.22	0.41	RAF 95	99		
Leptotrichia buccalis		29.34	0.15	FRU 99	1		
Lactobacillus acidophilus		3.77	0.10		99		
47361302	**Low discrimination**						**47361302**
					COC	SPO	
Lactobacillus acidophilus		85.63	0.40	FRU 99	1	1	
Clostridium barati		9.10	0.21	RHA 7	1	99	
Atopobium parvulum		2.10	0.14		99	1	
47361322	**Low discrimination**						**47361322**
					SPO		
Clostridium ramosum		57.45	0.45	FRU 99	92		
Lactobacillus acidophilus		28.14	0.40		1		
Clostridium barati		8.97	0.30		99		
47363001	**Good identification**						**47363001**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47363003	**Good identification**						**47363003**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47363021	**Good identification**						**47363021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47363023	**Good identification**						**47363023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47363100	**Low discrimination**						**47363100**
					COC	SPO	
Clostridium tertium		78.36	0.43	FRU 99	1	99	
Atopobium parvulum		8.74	0.27		99	1	
Actinomyces israelii		3.87	0.35		1	1	
47363101	**Low discrimination**						**47363101**
					SPO		
Actinomyces naeslundii		85.03	0.45	ESL 12	1		
Clostridium tertium		14.98	0.09	RAF 92	99		
47363102	**Low discrimination**						**47363102**
					SPO		
Clostridium tertium		75.10	0.51	FRU 99	99		
Lactobacillus acidophilus		14.05	0.38		1		
Bifidobacterium breve		3.25	0.37		1		
47363103	**Low discrimination**						**47363103**
					SPO		
Actinomyces naeslundii		61.45	0.45	ESL 12	1		
Clostridium tertium		32.47	0.17	RAF 92	99		
Lactobacillus acidophilus		6.07	0.04	FRU 95	1		
47363110	**Good identification**						**47363110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium tertium		98.36	0.47	FRU 99	99	99	
Bifidobacterium breve		0.71	0.20		99	1	
Leptotrichia buccalis		0.52	-.03		1	1	
47363112	**Low discrimination**						**47363112**
					GRA	SPO	
Clostridium tertium		83.06	0.55	FRU 99	99	99	
Leptotrichia buccalis		14.54	0.32		1	1	
Bifidobacterium breve		2.02	0.37		99	1	
47363120	**Low discrimination**						**47363120**
Clostridium tertium		58.67	0.51	FRU 99			
Clostridium ramosum		34.23	0.45				
Clostridium paraputrificum		3.59	0.22				
47363121	**Low discrimination**						**47363121**
					SPO		
Actinomyces naeslundii		54.45	0.45	ESL 12	1		
Clostridium tertium		28.77	0.17	RAF 92	99		
Clostridium ramosum		16.79	0.11	FRU 95	92		
47363122	**Low discrimination**						**47363122**
					SPO		
Clostridium tertium		52.40	0.59	FRU 99	99		
Clostridium ramosum		40.77	0.56		92		
Lactobacillus acidophilus		3.27	0.38		1		
47363130	**Good identification**						**47363130**
					SPO		
Clostridium tertium		99.25	0.55	FRU 99	99		
Bifidobacterium breve		0.42	0.24		1		
Clostridium ramosum		0.33	0.11		92		
47363132	**Very good identification**						**47363132**
					GRA	SPO	
Clostridium tertium		97.50	0.63	FRU 99	99	99	
Bifidobacterium breve		1.41	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
47363142	**Low discrimination**						**47363142**
					SPO		
Bifidobacterium breve		47.63	0.37	RAF 95	1		
Lactobacillus acidophilus		25.47	0.22	FRU 99	1		
Clostridium tertium		11.13	0.17		99		
47363152	**Low discrimination**						**47363152**
					GRA	SPO	
Bifidobacterium breve		65.74	0.37	RAF 95	99	1	
Clostridium tertium		27.31	0.21	FRU 99	99	99	
Leptotrichia buccalis		4.78	-.03		1	1	
47363162	**Low discrimination**						**47363162**
					SPO		
Bifidobacterium breve		46.44	0.41	RAF 95	1		
Clostridium tertium		18.31	0.25	FRU 99	99		
Clostridium ramosum		14.25	0.21		92		
47363172	**Low discrimination**						**47363172**
					SPO		
Bifidobacterium breve		58.79	0.41	RAF 95	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		41.21	0.29	FRU 99	99		
47363320	**Good identification**						**47363320**
Clostridium ramosum		98.40	0.47	FRU 99			
Clostridium tertium		1.29	0.17				
Clostridium barati		0.31	0.03				
47363322	**Very good identification**						**47363322**
					SPO		
Clostridium ramosum		98.35	0.58	FRU 99	92		
Clostridium tertium		0.96	0.25		99		
Lactobacillus acidophilus		0.45	0.18		1		
47363502	**Low discrimination**						**47363502**
					SPO		
Actinomyces israelii		42.18	0.36	FRU 99	1		
Bifidobacterium adolescentis		30.12	0.28	XYL 92	1		
Clostridium tertium		21.10	0.17		99		
47364712	**Good identification**						**47364712**
Clostridium clostridioforme		100.0	0.37	RAF 94			
47364732	**Good identification**						**47364732**
Clostridium clostridioforme		100.0	0.37	RAF 94			
47365100	**Very good identification**						**47365100**
					COC	SPO	
Atopobium parvulum		97.87	0.66	FRU 99	99	1	
Clostridium butyricum		1.52	0.28		1	99	
Actinomyces israelii		0.48	0.40		1	1	
47365102	**Low discrimination**						**47365102**
					COC	SPO	
Atopobium parvulum		87.96	0.53	FRU 99	99	1	
Actinomyces israelii		3.65	0.43		1	1	
Clostridium butyricum		3.00	0.21		1	99	
47365120	**Acceptable identification**						**47365120**
					COC	SPO	
Atopobium parvulum		92.89	0.53	FRU 99	99	1	
Clostridium paraputrificum		4.71	0.22		1	99	
Clostridium tertium		1.07	0.19		1	99	
47365122	**Low discrimination**						**47365122**
					COC	SPO	
Atopobium parvulum		61.25	0.40	FRU 99	99	1	
Clostridium tertium		12.04	0.27		1	99	
Lactobacillus acidophilus		10.32	0.25		1	1	
47365142	**Low discrimination**						**47365142**
					COC		
Actinomyces israelii		46.53	0.36	FRU 99	1		
Atopobium parvulum		29.11	0.19		99		
Lactobacillus acidophilus		10.60	0.10		1		
47365500	**Acceptable identification**						**47365500**
					COC	SPO	
Clostridium butyricum		90.71	0.44	RAF 95	1	99	
Atopobium parvulum		6.58	0.32	FRU 99	99	1	
Actinomyces israelii		2.60	0.39		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47365502	**Low discrimination**						**47365502**
					SPO		
Clostridium butyricum		67.51	0.37	RAF 95	99		
Bifidobacterium adolescentis		21.49	0.43	FRU 99	1		
Actinomyces israelii		7.47	0.42		1		
47365512	**Low discrimination**						**47365512**
					SPO		
Bifidobacterium adolescentis		67.25	0.45	RAF 92	1		
Clostridium clostridioforme		15.41	0.31	FRU 99	66		
Clostridium butyricum		9.10	0.15		99		
47365612	**Good identification**						**47365612**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47365632	**Good identification**						**47365632**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47365712	**Good identification**						**47365712**
Clostridium clostridioforme		99.40	0.45	RAF 94			
Bifidobacterium adolescentis		0.60	0.11	FRU 99			
47365732	**Good identification**						**47365732**
Clostridium clostridioforme		97.45	0.45	RAF 94			
Parabacteroides distasonis		1.28	0.19	FRU 99			
Bacteroides ovatus		1.27	0.03				
47367000	**Low discrimination**						**47367000**
					SPO		
Actinomyces israelii		87.59	0.43	FRU 99	1		
Clostridium tertium		8.61	0.11		99		
Clostridium innocuum		3.80	0.04		99		
47367002	**Low discrimination**						**47367002**
					SPO		
Actinomyces israelii		81.60	0.46	FRU 99	1		
Clostridium tertium		16.04	0.19		99		
Clostridium innocuum		2.36	0.04		99		
47367040	**Good identification**						**47367040**
Actinomyces israelii		100.0	0.36	FRU 99			
47367042	**Good identification**						**47367042**
Actinomyces israelii		100.0	0.39	FRU 99			
47367100	**Low discrimination**						**47367100**
					COC	SPO	
Clostridium tertium		63.03	0.45	FRU 99	1	99	
Actinomyces israelii		25.91	0.53		1	1	
Atopobium parvulum		9.46	0.32		99	1	
47367102	**Low discrimination**						**47367102**
					SPO		
Clostridium tertium		81.34	0.53	FRU 99	99		
Actinomyces israelii		16.72	0.56		1		
Bifidobacterium adolescentis		0.85	0.28		1		
47367110	**Good identification**						**47367110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium tertium		96.66	0.50	FRU 99	99		
Actinomyces israelii		3.34	0.39		1		
47367112	**Very good identification**						**47367112**
					SPO		
Clostridium tertium		97.17	0.58	FRU 99	99		
Actinomyces israelii		1.68	0.42		1		
Bifidobacterium adolescentis		0.70	0.29		1		
47367120	**Good identification**						**47367120**
					COC	SPO	
Clostridium tertium		97.79	0.53	FRU 99	1	99	
Atopobium parvulum		0.86	0.19		99	1	
Actinomyces israelii		0.86	0.33		1	1	
47367122	**Very good identification**						**47367122**
					SPO		
Clostridium tertium		98.91	0.62	FRU 99	99		
Clostridium ramosum		0.56	0.21		92		
Actinomyces israelii		0.43	0.36		1		
47367130	**Very good identification**						**47367130**
					SPO		
Clostridium tertium		99.87	0.58	FRU 99	99		
Actinomyces israelii		0.07	0.18		1		
Bifidobacterium dentium		0.06	-.07		1		
47367132	**Very good identification**						**47367132**
					SPO		
Clostridium tertium		99.76	0.66	FRU 99	99		
Bifidobacterium dentium		0.20	0.10		1		
Actinomyces israelii		0.04	0.21		1		
47367140	**Acceptable identification**						**47367140**
					SPO		
Actinomyces israelii		92.84	0.46	FRU 99	1		
Clostridium tertium		5.87	0.11		99		
Clostridium butyricum		1.29	-.07		99		
47367142	**Low discrimination**						**47367142**
					SPO		
Actinomyces israelii		87.53	0.49	FRU 99	1		
Clostridium tertium		11.06	0.19		99		
Bifidobacterium adolescentis		1.41	0.12		1		
47367302	**Low discrimination**						**47367302**
					SPO		
Actinomyces israelii		60.50	0.37	FRU 99	1		
Clostridium tertium		39.50	0.19	RHA 7	99		
47367400	**Low discrimination**						**47367400**
					SPO		
Actinomyces israelii		89.77	0.41	FRU 99	1		
Clostridium butyricum		10.23	0.04		99		
47367402	**Good identification**						**47367402**
					SPO		
Actinomyces israelii		95.57	0.44	FRU 99	1		
Clostridium butyricum		2.82	-.03		99		
Bifidobacterium adolescentis		1.61	0.08		1		
47367442	**Good identification**						**47367442**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.37	FRU 99			
47367500	**Low discrimination**						**47367500**
					SPO		
Actinomyces israelii		58.13	0.52	FRU 99	1		
Clostridium butyricum		39.76	0.28		99		
Clostridium tertium		1.75	0.11		99		
47367502	**Low discrimination**						**47367502**
					SPO		
Actinomyces israelii		59.21	0.55	FRU 99	1		
Bifidobacterium adolescentis		24.60	0.42		1		
Clostridium butyricum		10.50	0.21		99		
47367512	**Low discrimination**						**47367512**
Bifidobacterium dentium		82.22	0.40	RAF 95			
Bifidobacterium adolescentis		11.68	0.43	FRU 99			
Actinomyces israelii		3.44	0.40				
47367532	**Acceptable identification**						**47367532**
					SPO		
Bifidobacterium dentium		95.00	0.44	RAF 95	1		
Clostridium tertium		4.79	0.32	FRU 99	99		
Actinomyces israelii		0.14	0.20		1		
47367540	**Low discrimination**						**47367540**
					COC	SPO	
Actinomyces israelii		49.43	0.45	FRU 99	1	1	
Blautia producta		43.01	0.24		99	1	
Clostridium butyricum		7.56	0.10		1	99	
47367542	**Low discrimination**						**47367542**
					COC		
Actinomyces israelii		80.90	0.48	FRU 99	1		
Bifidobacterium adolescentis		10.68	0.26		1		
Blautia producta		5.21	0.07		99		
47367702	**Low discrimination**						**47367702**
					SPO		
Actinomyces israelii		88.60	0.35	FRU 99	1		
Clostridium butyricum		6.46	-.05	RHA 7	99		
Bifidobacterium adolescentis		4.94	0.08		1		
47370000	**Good identification**						**47370000**
					GRA	SPO	
Lactobacillus jensenii		91.27	0.66	GAL 99	99	1	
Fusobacterium mortiferum		4.29	0.53		1	1	
Clostridium barati		3.08	0.36		99	99	
47370002	**Low discrimination**						**47370002**
					SPO		
Lactobacillus jensenii		79.64	0.63	GAL 99	1		
Clostridium barati		12.09	0.44		99		
Lactobacillus acidophilus		6.58	0.41		1		
47370010	**Very good identification**						**47370010**
					GRA		
Lactobacillus jensenii		99.59	0.58	GAL 99	99		
Leptotrichia buccalis		0.17	-.03		1		
Fusobacterium mortiferum		0.14	0.19		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47370012	**Low discrimination**							**47370012**
					GRA	SPO		
Lactobacillus jensenii		79.47	0.55	GAL 99	99	1		
Leptotrichia buccalis		19.83	0.32		1	1		
Clostridium barati		0.37	0.10		99	99		
47370020	**Low discrimination**							**47370020**
					SPO			
Lactobacillus jensenii		68.09	0.56	GAL 99	1			
Clostridium barati		27.57	0.44		99			
Clostridium paraputrificum		2.97	0.22		99			
47370022	**Low discrimination**							**47370022**
					SPO			
Clostridium barati		56.88	0.52	TRE 8	99			
Lactobacillus jensenii		31.22	0.53	GAL 95	1			
Lactobacillus acidophilus		10.32	0.41		1			
47370030	**Good identification**							**47370030**
					SPO			
Lactobacillus jensenii		98.79	0.47	GAL 99	1			
Clostridium barati		1.21	0.10		99			
47370032	**Low discrimination**							**47370032**
					GRA	SPO		
Lactobacillus jensenii		84.55	0.44	GAL 99	99	1		
Leptotrichia buccalis		9.38	0.15		1	1		
Clostridium barati		4.67	0.18		99	99		
47370100	**Low discrimination**							**47370100**
					COC	GRA	SPO	
Atopobium parvulum		66.81	0.61	GAL 99	99	99	1	
Fusobacterium mortiferum		11.17	0.59		1	1	1	
Clostridium barati		10.31	0.44		1	99	99	
47370102	**Low discrimination**							**47370102**
					COC	SPO		
Lactobacillus acidophilus		54.44	0.60	GAL 99	1	1		
Clostridium barati		26.10	0.52		1	99		
Atopobium parvulum		9.95	0.48		99	1		
47370112	**Very good identification**							**47370112**
					GRA			
Leptotrichia buccalis		99.54	0.66	GAL 99	1			
Bifidobacterium breve		0.14	0.37		99			
Lactobacillus acidophilus		0.12	0.25		99			
47370120	**Low discrimination**							**47370120**
					COC	SPO		
Clostridium paraputrificum		68.61	0.56	GAL 99	1	99		
Clostridium barati		19.31	0.52		1	99		
Atopobium parvulum		7.37	0.48		99	1		
47370122	**Low discrimination**							**47370122**
					SPO			
Clostridium barati		42.50	0.60	TRE 8	99			
Lactobacillus acidophilus		29.55	0.60	GAL 95	1			
Clostridium paraputrificum		21.57	0.50		99			
47370132	**Acceptable identification**							**47370132**
					GRA	SPO		
Leptotrichia buccalis		92.95	0.50	GAL 99	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		2.09	0.41	NAG 10	99	1	
Clostridium tertium		1.46	0.29		99	99	
47370142	**Low discrimination**						**47370142**
					SPO		
Lactobacillus acidophilus		68.11	0.44	GAL 99	1		
Clostridium barati		22.97	0.34	SOR 11	99		
Bifidobacterium breve		6.71	0.37		1		
47370160	**Low discrimination**						**47370160**
					SPO		
Clostridium paraputrificum		72.44	0.37	GAL 99	99		
Clostridium barati		23.56	0.34	SOR 7	99		
Bifidobacterium breve		1.21	0.24		1		
47370162	**Low discrimination**						**47370162**
					SPO		
Clostridium barati		38.20	0.42	TRE 8	99		
Lactobacillus acidophilus		37.75	0.44	GAL 95	1		
Clostridium paraputrificum		16.78	0.31	SOR 8	99		
47370172	**Low discrimination**						**47370172**
					GRA	SPO	
Bifidobacterium breve		62.83	0.41	RAF 95	99	1	
Leptotrichia buccalis		28.26	0.15	GAL 99	1	1	
Clostridium barati		3.67	0.08		99	99	
47370302	**Low discrimination**						**47370302**
					SPO		
Lactobacillus acidophilus		60.36	0.40	GAL 99	1		
Clostridium barati		33.43	0.34	RHA 7	99		
Clostridium clostridioforme		2.50	0.23		66		
47370322	**Low discrimination**						**47370322**
					SPO		
Clostridium ramosum		41.34	0.45	GAL 99	92		
Clostridium barati		33.65	0.42		99		
Lactobacillus acidophilus		20.25	0.40		1		
47370532	**Low discrimination**						**47370532**
Clostridium clostridioforme		75.72	0.39	RAF 94			
Leptotrichia buccalis		24.28	0.15	XYL 91			
47370612	**Good identification**						**47370612**
Clostridium clostridioforme		100.0	0.45	RAF 94			
47370632	**Good identification**						**47370632**
Clostridium clostridioforme		100.0	0.45	RAF 94			
47370702	**Good identification**						**47370702**
					SPO		
Clostridium clostridioforme		97.64	0.43	RAF 94	66		
Lactobacillus acidophilus		1.52	0.06	XYL 91	1		
Clostridium barati		0.84	-.01		99		
47370712	**Good identification**						**47370712**
Clostridium clostridioforme		99.60	0.54	RAF 94			
Leptotrichia buccalis		0.40	-.03	XYL 91			
47370722	**Acceptable identification**						**47370722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		93.18	0.43	RAF 94		
Clostridium ramosum		2.96	0.11	XYL 91		
Clostridium barati		2.41	0.08			
47370732	**Good identification**					**47370732**
Clostridium clostridioforme		100.0	0.54	RAF 94		
47371000	**Very good identification**					**47371000**
					COC	
Lactobacillus jensenii		97.83	1.00		1	
Lactobacillus catenaformis		0.66	0.63		1	
Atopobium parvulum		0.65	0.61		99	
47371001	**Very good identification**					**47371001**
Lactobacillus jensenii		96.76	0.66	URE 1		
Actinomyces naeslundii		1.13	0.54			
Lactobacillus catenaformis		0.65	0.29			
47371002	**Low discrimination**					**47371002**
					SPO	
Lactobacillus jensenii		88.86	0.97		1	
Lactobacillus acidophilus		7.34	0.76		1	
Clostridium barati		2.59	0.66		99	
47371003	**Low discrimination**					**47371003**
					SPO	
Lactobacillus jensenii		87.55	0.63	URE 1	1	
Lactobacillus acidophilus		7.23	0.41		1	
Clostridium barati		2.55	0.31		99	
47371010	**Excellent identification**					**47371010**
					GRA	
Lactobacillus jensenii		99.75	0.92		99	
Leptotrichia buccalis		0.17	0.32		1	
Lactobacillus catenaformis		0.02	0.29		99	
47371011	**Very good identification**					**47371011**
					GRA	
Lactobacillus jensenii		99.65	0.58	URE 1	99	
Actinomyces naeslundii		0.18	0.32		99	
Leptotrichia buccalis		0.17	-.03		1	
47371012	**Low discrimination**					**47371012**
					GRA	
Lactobacillus jensenii		79.74	0.89		99	
Leptotrichia buccalis		19.90	0.66		1	
Lactobacillus acidophilus		0.20	0.41		99	
47371013	**Low discrimination**					**47371013**
					GRA	
Lactobacillus jensenii		79.64	0.55	URE 1	99	
Leptotrichia buccalis		19.87	0.32		1	
Actinomyces naeslundii		0.22	0.32		99	
47371020	**Low discrimination**					**47371020**
					SPO	
Lactobacillus jensenii		87.77	0.90		1	
Clostridium barati		6.82	0.66		99	
Clostridium paraputrificum		3.83	0.56		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47371021	**Low discrimination**						**47371021**
					SPO		
Lactobacillus jensenii		84.38	0.56	URE 1	1		
Clostridium barati		6.56	0.31		99		
Actinomyces naeslundii		3.96	0.54		1		
47371022	**Low discrimination**						**47371022**
					SPO		
Lactobacillus jensenii		57.78	0.87		1		
Clostridium barati		20.20	0.74		99		
Lactobacillus acidophilus		19.09	0.76		1		
47371023	**Low discrimination**						**47371023**
					SPO		
Lactobacillus jensenii		55.59	0.53	URE 1	1		
Clostridium barati		19.44	0.40		99		
Lactobacillus acidophilus		18.37	0.41		1		
47371030	**Very good identification**						**47371030**
					SPO		
Lactobacillus jensenii		99.45	0.82		1		
Clostridium barati		0.23	0.31		99		
Clostridium paraputrificum		0.13	0.22		99		
47371031	**Good identification**						**47371031**
					SPO		
Lactobacillus jensenii		99.03	0.47	URE 1	1		
Actinomyces naeslundii		0.73	0.32		1		
Clostridium barati		0.23	-.03		99		
47371032	**Low discrimination**						**47371032**
					GRA	SPO	
Lactobacillus jensenii		87.91	0.79		99	1	
Leptotrichia buccalis		9.75	0.50		1	1	
Clostridium barati		0.93	0.40		99	99	
47371033	**Low discrimination**						**47371033**
					GRA		
Lactobacillus jensenii		87.52	0.44	URE 1	99		
Leptotrichia buccalis		9.71	0.15		1		
Actinomyces naeslundii		0.97	0.32		99		
47371040	**Good identification**						**47371040**
					SPO		
Lactobacillus jensenii		92.11	0.66	SOR 1	1		
Clostridium barati		5.13	0.39		99		
Lactobacillus acidophilus		0.63	0.26		1		
47371042	**Low discrimination**						**47371042**
					SPO		
Lactobacillus acidophilus		43.95	0.60	LAC 92	1		
Lactobacillus jensenii		43.49	0.63	SOR 11	1		
Clostridium barati		10.91	0.47		99		
47371050	**Very good identification**						**47371050**
Lactobacillus jensenii		99.22	0.58	SOR 1			
Bifidobacterium breve		0.23	0.20				
Actinomyces israelii		0.22	0.25				
47371052	**Low discrimination**						**47371052**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		75.33	0.55	SOR 1	99		
Leptotrichia buccalis		18.80	0.32		1		
Bifidobacterium breve		2.61	0.37		99		
47371060	**Low discrimination**						**47371060**
					SPO		
Lactobacillus jensenii		48.17	0.56	SOR 1	1		
Clostridium barati		32.22	0.47		99		
Clostridium paraputrificum		15.64	0.37		99		
47371062	**Low discrimination**						**47371062**
					SPO		
Lactobacillus acidophilus		48.16	0.60	LAC 92	1		
Clostridium barati		35.85	0.56	SOR 11	99		
Lactobacillus jensenii		11.91	0.53		1		
47371070	**Good identification**						**47371070**
					SPO		
Lactobacillus jensenii		95.57	0.47	SOR 1	1		
Clostridium barati		1.94	0.13		99		
Bifidobacterium breve		1.55	0.24		1		
47371072	**Low discrimination**						**47371072**
Lactobacillus jensenii		63.39	0.44	SOR 1			
Bifidobacterium breve		15.63	0.41				
Lactobacillus acidophilus		7.77	0.26				
47371100	**Low discrimination**						**47371100**
					COC	SPO	
Atopobium parvulum		82.88	0.95		99	1	
Collinsella aerofaciens		9.25	0.66		1	1	
Clostridium barati		2.45	0.66		1	99	
47371101	**Low discrimination**						**47371101**
					COC		
Atopobium parvulum		79.47	0.61	URE 1	99		
Collinsella aerofaciens		8.87	0.32		1		
Actinomyces naeslundii		4.04	0.62		1		
47371102	**Low discrimination**						**47371102**
					COC	SPO	
Lactobacillus acidophilus		69.97	0.94		1	1	
Atopobium parvulum		12.79	0.83		99	1	
Clostridium barati		6.44	0.74		1	99	
47371103	**Low discrimination**						**47371103**
					COC	SPO	
Lactobacillus acidophilus		67.49	0.60	URE 1	1	1	
Atopobium parvulum		12.34	0.48		99	1	
Clostridium barati		6.21	0.40		1	99	
47371110	**Low discrimination**						**47371110**
					COC	GRA	
Leptotrichia buccalis		80.80	0.66	BGL 99	1	1	
Atopobium parvulum		9.66	0.61		99	99	
Lactobacillus jensenii		4.91	0.58		1	99	
47371112	**Excellent identification**						**47371112**
					GRA		
Leptotrichia buccalis		99.60	1.00		1		
Bifidobacterium breve		0.14	0.71		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Lactobacillus acidophilus	0.12	0.60		99		
47371113	**Very good identification**						**47371113**
					GRA		
	Leptotrichia buccalis	99.57	0.66	URE 1	1		
	Bifidobacterium breve	0.14	0.37		99		
	Lactobacillus acidophilus	0.12	0.25		99		
47371120	**Low discrimination**						**47371120**
					COC	SPO	
	Clostridium paraputrificum	81.39	0.90		1	99	
	Atopobium parvulum	8.74	0.83		99	1	
	Clostridium barati	4.40	0.74		1	99	
47371121	**Low discrimination**						**47371121**
					COC	SPO	
	Clostridium paraputrificum	79.40	0.56	URE 1	1	99	
	Atopobium parvulum	8.52	0.48		99	1	
	Clostridium barati	4.29	0.40		1	99	
47371122	**Low discrimination**						**47371122**
					SPO		
	Lactobacillus acidophilus	45.02	0.94		1		
	Clostridium paraputrificum	32.86	0.84		99		
	Clostridium barati	12.43	0.82		99		
47371123	**Low discrimination**						**47371123**
					SPO		
	Lactobacillus acidophilus	43.99	0.60	URE 1	1		
	Clostridium paraputrificum	32.11	0.50		99		
	Clostridium barati	12.14	0.48		99		
47371130	**Low discrimination**						**47371130**
					GRA	SPO	
	Clostridium paraputrificum	43.75	0.56	MLZ 1	99	99	
	Leptotrichia buccalis	24.74	0.50		1	1	
	Clostridium tertium	12.85	0.55		99	99	
47371132	**Good identification**						**47371132**
					GRA	SPO	
	Leptotrichia buccalis	94.15	0.84	NAG 10	1	1	
	Bifidobacterium breve	2.11	0.75		99	1	
	Clostridium tertium	1.48	0.63		99	99	
47371133	**Acceptable identification**						**47371133**
					GRA	SPO	
	Leptotrichia buccalis	93.93	0.50	NAG 10	1	1	
	Bifidobacterium breve	2.11	0.41	URE 1	99	1	
	Clostridium tertium	1.48	0.29		99	99	
47371140	**Low discrimination**						**47371140**
					COC	SPO	
	Atopobium parvulum	55.35	0.61	SOR 1	99	1	
	Clostridium barati	14.11	0.47		1	99	
	Clostridium paraputrificum	6.85	0.37		1	99	
47371142	**Low discrimination**						**47371142**
					SPO		
	Lactobacillus acidophilus	83.41	0.78	SOR 11	1		
	Bifidobacterium breve	8.21	0.71		1		
	Clostridium barati	5.40	0.56		99		
47371143	**Low discrimination**						**47371143**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		78.81	0.44	SOR 11			
Bifidobacterium breve		7.76	0.37	URE 1			
Actinomyces naeslundii		5.93	0.49				
47371150	**Low discrimination**						**47371150**
					COC	GRA	
Bifidobacterium breve		50.48	0.54	RAF 95	1	99	
Leptotrichia buccalis		37.11	0.32	BGL 91	1	1	
Atopobium parvulum		4.44	0.27		99	99	
47371152	**Low discrimination**						**47371152**
					GRA		
Leptotrichia buccalis		86.39	0.66	SOR 1	1		
Bifidobacterium breve		12.00	0.71		99		
Lactobacillus acidophilus		1.23	0.44		99		
47371160	**Low discrimination**						**47371160**
					SPO		
Clostridium paraputrificum		89.42	0.71	SOR 7	99		
Clostridium barati		5.58	0.56		99		
Bifidobacterium breve		1.49	0.58		1		
47371161	**Low discrimination**						**47371161**
					SPO		
Clostridium paraputrificum		84.16	0.37	SOR 7	99		
Actinomyces naeslundii		6.10	0.49	URE 1	1		
Clostridium barati		5.25	0.21		99		
47371162	**Low discrimination**						**47371162**
					SPO		
Lactobacillus acidophilus		54.54	0.78	SOR 11	1		
Clostridium paraputrificum		24.25	0.65		99		
Clostridium barati		10.59	0.64		99		
47371163	**Low discrimination**						**47371163**
					SPO		
Lactobacillus acidophilus		52.51	0.44	SOR 11	1		
Clostridium paraputrificum		23.34	0.31	URE 1	99		
Clostridium barati		10.20	0.30		99		
47371170	**Low discrimination**						**47371170**
					GRA	SPO	
Bifidobacterium breve		57.33	0.58	RAF 95	99	1	
Clostridium paraputrificum		34.70	0.37	BGL 91	99	99	
Leptotrichia buccalis		2.63	0.15		1	1	
47371172	**Low discrimination**						**47371172**
					GRA		
Bifidobacterium breve		64.39	0.75	RAF 95	99		
Leptotrichia buccalis		28.96	0.50		1		
Lactobacillus acidophilus		3.72	0.44		99		
47371173	**Low discrimination**						**47371173**
					GRA		
Bifidobacterium breve		64.27	0.41	RAF 95	99		
Leptotrichia buccalis		28.91	0.15	URE 1	1		
Lactobacillus acidophilus		3.71	0.10		99		
47371200	**Good identification**						**47371200**
					SPO		
Lactobacillus jensenii		92.95	0.66	RHA 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		5.18	0.39		99		
Lactobacillus catenaformis		0.62	0.29		1		
47371202	**Low discrimination**						**47371202**
					SPO		
Lactobacillus jensenii		52.98	0.63	RHA 1	1		
Lactobacillus acidophilus		32.61	0.56		1		
Clostridium barati		13.29	0.47		99		
47371210	**Very good identification**						**47371210**
					GRA	SPO	
Lactobacillus jensenii		99.66	0.58	RHA 1	99	1	
Clostridium barati		0.17	0.05		99	99	
Leptotrichia buccalis		0.17	-.03		1	1	
47371212	**Low discrimination**						**47371212**
					GRA		
Lactobacillus jensenii		76.16	0.55	RHA 1	99		
Leptotrichia buccalis		19.01	0.32		1		
Clostridium clostridioforme		2.76	0.33		50		
47371220	**Low discrimination**						**47371220**
					SPO		
Lactobacillus jensenii		54.17	0.56	RHA 1	1		
Clostridium barati		36.23	0.47		99		
Clostridium ramosum		5.27	0.35		92		
47371222	**Low discrimination**						**47371222**
					SPO		
Clostridium barati		41.43	0.56	TRE 8	99		
Lactobacillus acidophilus		33.89	0.56	RHA 8	1		
Lactobacillus jensenii		13.76	0.53		1		
47371230	**Good identification**						**47371230**
					SPO		
Lactobacillus jensenii		97.73	0.47	RHA 1	1		
Clostridium barati		1.98	0.13		99		
Clostridium ramosum		0.29	0.00		92		
47371232	**Low discrimination**						**47371232**
					GRA		
Lactobacillus jensenii		69.24	0.44	RHA 1	99		
Clostridium clostridioforme		10.05	0.33		50		
Leptotrichia buccalis		7.68	0.15		1		
47371242	**Low discrimination**						**47371242**
					SPO		
Lactobacillus acidophilus		69.90	0.41	LAC 92	1		
Clostridium barati		20.04	0.29	RHA 7	99		
Lactobacillus jensenii		9.28	0.29	SOR 11	1		
47371262	**Low discrimination**						**47371262**
					SPO		
Lactobacillus acidophilus		51.95	0.41	LAC 92	1		
Clostridium barati		44.68	0.37	RHA 7	99		
Lactobacillus jensenii		1.72	0.18	SOR 11	1		
47371300	**Low discrimination**						**47371300**
					COC	SPO	
Atopobium parvulum		64.93	0.61	RHA 1	99	1	
Clostridium barati		16.55	0.47		1	99	
Collinsella aerofaciens		7.25	0.32		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47371302	**Low discrimination**						**47371302**
					COC	SPO	
Lactobacillus acidophilus		84.82	0.75	RHA 7	1	1	
Clostridium barati		9.02	0.56		1	99	
Atopobium parvulum		2.08	0.48		99	1	
47371303	**Low discrimination**						**47371303**
					COC	SPO	
Lactobacillus acidophilus		85.12	0.40	RHA 7	1	1	
Clostridium barati		9.05	0.21	URE 1	1	99	
Atopobium parvulum		2.09	0.14		99	1	
47371312	**Very good identification**						**47371312**
					GRA		
Leptotrichia buccalis		98.27	0.66	RHA 1	1		
Lactobacillus acidophilus		0.85	0.40		99		
Clostridium clostridioforme		0.48	0.42		50		
47371320	**Low discrimination**						**47371320**
Clostridium ramosum		57.06	0.69				
Clostridium paraputrificum		25.55	0.56				
Clostridium barati		11.88	0.56				
47371322	**Low discrimination**						**47371322**
					SPO		
Clostridium ramosum		57.31	0.79		92		
Lactobacillus acidophilus		28.07	0.75		1		
Clostridium barati		8.95	0.64		99		
47371323	**Low discrimination**						**47371323**
					SPO		
Clostridium ramosum		57.33	0.45	URE 1	92		
Lactobacillus acidophilus		28.08	0.40		1		
Clostridium barati		8.96	0.30		99		
47371332	**Low discrimination**						**47371332**
					SPO	GRA	
Leptotrichia buccalis		72.60	0.50	RHA 1	1	1	
Clostridium ramosum		11.58	0.45	NAG 10	92	50	
Lactobacillus acidophilus		5.67	0.40		1	99	
47371342	**Good identification**						**47371342**
					SPO		
Lactobacillus acidophilus		91.12	0.59	RHA 7	1		
Clostridium barati		6.81	0.37	SOR 11	99		
Bifidobacterium breve		1.20	0.37		1		
47371360	**Low discrimination**						**47371360**
Clostridium paraputrificum		51.19	0.37	RHA 1			
Clostridium barati		27.50	0.37	SOR 7			
Clostridium ramosum		15.34	0.35				
47371362	**Low discrimination**						**47371362**
					SPO		
Lactobacillus acidophilus		67.45	0.59	RHA 7	1		
Clostridium barati		15.13	0.46	SOR 11	99		
Clostridium ramosum		11.25	0.45		92		
47371372	**Low discrimination**						**47371372**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		47.76	0.41	RAF 95	99		
Leptotrichia buccalis		21.49	0.15	RHA 1	1		
Lactobacillus acidophilus		20.54	0.25		99		
47371400	**Good identification**						**47371400**
					SPO		
Lactobacillus jensenii		94.86	0.66	ARA 1	1		
Clostridium butyricum		2.06	0.29		99		
Actinomyces israelii		0.99	0.44		1		
47371402	**Low discrimination**						**47371402**
					SPO		
Lactobacillus jensenii		85.37	0.63	ARA 1	1		
Lactobacillus acidophilus		7.05	0.41		1		
Clostridium barati		2.49	0.31		99		
47371410	**Very good identification**						**47371410**
					SPO		
Lactobacillus jensenii		98.98	0.58	ARA 1	1		
Actinomyces israelii		0.47	0.30		1		
Clostridium butyricum		0.34	0.07		99		
47371412	**Low discrimination**						**47371412**
					GRA		
Lactobacillus jensenii		73.51	0.55	ARA 1	99		
Leptotrichia buccalis		18.35	0.32		1		
Clostridium clostridioforme		5.70	0.39		50		
47371420	**Low discrimination**						**47371420**
					SPO		
Lactobacillus jensenii		87.07	0.56	ARA 1	1		
Clostridium barati		6.77	0.31		99		
Clostridium paraputrificum		3.79	0.22		99		
47371422	**Low discrimination**						**47371422**
					SPO		
Lactobacillus jensenii		56.76	0.53	ARA 1	1		
Clostridium barati		19.85	0.40		99		
Lactobacillus acidophilus		18.76	0.41		1		
47371430	**Good identification**						**47371430**
					SPO		
Lactobacillus jensenii		99.56	0.47	ARA 1	1		
Clostridium barati		0.23	-.03		99		
Clostridium clostridioforme		0.21	0.04		66		
47371432	**Low discrimination**						**47371432**
					GRA		
Lactobacillus jensenii		69.37	0.44	ARA 1	99		
Clostridium clostridioforme		21.50	0.39		50		
Leptotrichia buccalis		7.70	0.15		1		
47371500	**Low discrimination**						**47371500**
					COC	SPO	
Atopobium parvulum		48.41	0.61	ARA 1	99	1	
Clostridium butyricum		38.36	0.52		1	99	
Collinsella aerofaciens		5.40	0.32		1	1	
47371502	**Low discrimination**						**47371502**
					SPO		
Lactobacillus acidophilus		42.44	0.60	ARA 1	1		
Bifidobacterium adolescentis		20.84	0.63		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		13.55	0.45		99		
47371512	**Good identification**						**47371512**
					GRA		
Leptotrichia buccalis		91.95	0.66	ARA 1	1		
Bifidobacterium adolescentis		6.37	0.65		99		
Clostridium clostridioforme		0.97	0.48		50		
47371520	**Low discrimination**						**47371520**
					COC	SPO	
Clostridium paraputrificum		78.56	0.56	ARA 1	1	99	
Atopobium parvulum		8.43	0.48		99	1	
Clostridium barati		4.24	0.40		1	99	
47371522	**Low discrimination**						**47371522**
					SPO		
Lactobacillus acidophilus		43.75	0.60	ARA 1	1		
Clostridium paraputrificum		31.93	0.50		99		
Clostridium barati		12.08	0.48		99		
47371532	**Low discrimination**						**47371532**
					GRA		
Leptotrichia buccalis		85.27	0.50	ARA 1	1		
Clostridium clostridioforme		8.06	0.48	NAG 10	50		
Bifidobacterium breve		1.92	0.41		99		
47371542	**Low discrimination**						**47371542**
Lactobacillus acidophilus		45.09	0.44	ARA 1			
Bifidobacterium adolescentis		22.14	0.48	SOR 11			
Actinomyces israelii		14.09	0.51				
47371560	**Low discrimination**						**47371560**
Clostridium paraputrificum		84.96	0.37	ARA 1			
Clostridium barati		5.30	0.21	SOR 7			
Clostridium butyricum		4.12	0.16				
47371562	**Low discrimination**						**47371562**
					SPO		
Lactobacillus acidophilus		53.37	0.44	ARA 1	1		
Clostridium paraputrificum		23.73	0.31	SOR 11	99		
Clostridium barati		10.37	0.30		99		
47371572	**Low discrimination**						**47371572**
					GRA		
Bifidobacterium breve		61.45	0.41	RAF 95	99		
Leptotrichia buccalis		27.64	0.15	ARA 1	1		
Lactobacillus acidophilus		3.55	0.10		99		
47371602	**Low discrimination**						**47371602**
Clostridium clostridioforme		66.43	0.42	RAF 94			
Lactobacillus jensenii		16.14	0.29	XYL 91			
Lactobacillus acidophilus		9.93	0.22				
47371612	**Good identification**						**47371612**
					GRA		
Clostridium clostridioforme		97.69	0.53	RAF 94	50		
Lactobacillus jensenii		1.74	0.21	XYL 91	99		
Leptotrichia buccalis		0.43	-.03		1		
47371622	**Low discrimination**						**47371622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium clostridioforme		69.54	0.42	RAF 94	66		
Clostridium barati		12.71	0.21	XYL 91	99		
Lactobacillus acidophilus		10.40	0.22		1		
47371632	**Good identification**						**47371632**
Clostridium clostridioforme		99.56	0.53	RAF 94			
Lactobacillus jensenii		0.44	0.10	XYL 91			
47371702	**Low discrimination**						**47371702**
					SPO		
Clostridium clostridioforme		57.08	0.51	RAF 94	66		
Lactobacillus acidophilus		29.32	0.40	XYL 91	1		
Clostridium butyricum		3.85	0.19		99		
47371712	**Low discrimination**						**47371712**
					GRA		
Clostridium clostridioforme		87.42	0.62	RAF 94	50		
Leptotrichia buccalis		11.47	0.32	XYL 91	1		
Bifidobacterium adolescentis		0.79	0.31		99		
47371722	**Low discrimination**						**47371722**
					SPO		
Clostridium ramosum		35.32	0.45	ARA 1	92		
Clostridium clostridioforme		33.69	0.51		66		
Lactobacillus acidophilus		17.30	0.40		1		
47371732	**Very good identification**						**47371732**
Clostridium clostridioforme		96.70	0.62	RAF 94			
Parabacteroides distasonis		1.43	0.37	XYL 91			
Leptotrichia buccalis		1.41	0.15				
47372001	**Good identification**						**47372001**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47372003	**Good identification**						**47372003**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47372021	**Good identification**						**47372021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47372023	**Low discrimination**						**47372023**
					SPO		
Actinomyces naeslundii		78.04	0.36	ESL 12	1		
Clostridium barati		21.96	-.01	RAF 92	99		
47372100	**Low discrimination**						**47372100**
					COC	SPO	
Clostridium tertium		71.10	0.43	GAL 99	1	99	
Clostridium barati		10.53	0.26		1	99	
Atopobium parvulum		7.93	0.27		99	1	
47372101	**Low discrimination**						**47372101**
					SPO		
Actinomyces naeslundii		81.80	0.43	ESL 12	1		
Clostridium tertium		18.20	0.09	RAF 92	99		
47372102	**Low discrimination**						**47372102**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		69.03	0.51	GAL 99	99		
Lactobacillus acidophilus		12.91	0.38		1		
Clostridium barati		10.23	0.34		99		
47372103	**Low discrimination**						**47372103**
					SPO		
Actinomyces naeslundii		52.88	0.43	ESL 12	1		
Clostridium tertium		35.29	0.17	RAF 92	99		
Lactobacillus acidophilus		6.60	0.04	GAL 96	1		
47372110	**Good identification**						**47372110**
					GRA	SPO	
Clostridium tertium		98.36	0.47	GAL 99	99	99	
Bifidobacterium breve		0.71	0.20		99	1	
Leptotrichia buccalis		0.52	-.03		1	1	
47372112	**Low discrimination**						**47372112**
					GRA	SPO	
Clostridium tertium		83.00	0.55	GAL 99	99	99	
Leptotrichia buccalis		14.53	0.32		1	1	
Bifidobacterium breve		2.02	0.37		99	1	
47372120	**Low discrimination**						**47372120**
Clostridium tertium		54.90	0.51	GAL 99			
Clostridium ramosum		32.03	0.45				
Clostridium barati		8.13	0.34				
47372121	**Low discrimination**						**47372121**
					SPO		
Actinomyces naeslundii		46.39	0.43	ESL 12	1		
Clostridium tertium		30.96	0.17	RAF 92	99		
Clostridium ramosum		18.07	0.11	GAL 96	92		
47372122	**Low discrimination**						**47372122**
Clostridium tertium		49.33	0.59	GAL 99			
Clostridium ramosum		38.38	0.56				
Clostridium barati		7.31	0.42				
47372130	**Good identification**						**47372130**
					SPO		
Clostridium tertium		99.16	0.55	GAL 99	99		
Bifidobacterium breve		0.42	0.24		1		
Clostridium ramosum		0.33	0.11		92		
47372132	**Very good identification**						**47372132**
					GRA	SPO	
Clostridium tertium		97.42	0.63	GAL 99	99	99	
Bifidobacterium breve		1.40	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
47372142	**Low discrimination**						**47372142**
					SPO		
Bifidobacterium breve		42.93	0.37	RAF 95	1		
Lactobacillus acidophilus		22.95	0.22	GAL 99	1		
Clostridium barati		12.79	0.16		99		
47372152	**Low discrimination**						**47372152**
					GRA	SPO	
Bifidobacterium breve		65.74	0.37	RAF 95	99	1	
Clostridium tertium		27.31	0.21	GAL 99	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Leptotrichia buccalis		4.78	-.03		1 1	
47372162	**Low discrimination**					**47372162**
					SPO	
Bifidobacterium breve		39.16	0.41	RAF 95	1	
Clostridium barati		19.69	0.24	GAL 99	99	
Clostridium tertium		15.44	0.25		99	
47372172	**Low discrimination**					**47372172**
					SPO	
Bifidobacterium breve		58.79	0.41	RAF 95	1	
Clostridium tertium		41.21	0.29	GAL 99	99	
47372320	**Good identification**					**47372320**
Clostridium ramosum		97.11	0.47	GAL 99		
Clostridium barati		1.62	0.16			
Clostridium tertium		1.27	0.17			
47372322	**Very good identification**					**47372322**
Clostridium ramosum		97.39	0.58	GAL 99		
Clostridium barati		1.22	0.24			
Clostridium tertium		0.95	0.25			
47372502	**Low discrimination**					**47372502**
					SPO	
Actinomyces israelii		40.90	0.36	GAL 99	1	
Bifidobacterium adolescentis		29.21	0.28	XYL 92	1	
Clostridium tertium		20.46	0.17		99	
47373000	**Low discrimination**					**47373000**
					SPO	
Lactobacillus jensenii		73.82	0.66	MAN 1	1	
Clostridium innocuum		9.75	0.48		99	
Actinomyces israelii		5.36	0.59		1	
47373001	**Low discrimination**					**47373001**
					SPO	
Actinomyces naeslundii		63.05	0.59	ESL 12	1	
Lactobacillus jensenii		28.00	0.32	RAF 92	1	
Clostridium innocuum		3.70	0.14		99	
47373002	**Low discrimination**					**47373002**
					SPO	
Lactobacillus jensenii		42.31	0.63	MAN 1	1	
Lactobacillus acidophilus		18.21	0.54		1	
Clostridium tertium		11.31	0.51		99	
47373003	**Low discrimination**					**47373003**
Actinomyces naeslundii		59.39	0.59	ESL 12		
Lactobacillus jensenii		17.58	0.29	RAF 92		
Lactobacillus acidophilus		7.57	0.20			
47373010	**Low discrimination**					**47373010**
					SPO	
Lactobacillus jensenii		72.78	0.58	MAN 1	1	
Clostridium tertium		23.06	0.47		99	
Actinomyces israelii		2.37	0.45		1	
47373012	**Low discrimination**					**47373012**
					GRA SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		50.31	0.55	LAC 99	99	99	
Lactobacillus jensenii		35.28	0.55		99	1	
Leptotrichia buccalis		8.81	0.32		1	1	
47373020	**Low discrimination**						**47373020**
					SPO		
Lactobacillus jensenii		33.82	0.56	MAN 1	1		
Clostridium tertium		24.11	0.51		99		
Clostridium barati		22.62	0.47		99		
47373021	**Low discrimination**						**47373021**
					SPO		
Actinomyces naeslundii		76.30	0.59	ESL 12	1		
Lactobacillus jensenii		8.47	0.21	RAF 92	1		
Clostridium tertium		6.04	0.17		99		
47373022	**Low discrimination**						**47373022**
Clostridium tertium		27.42	0.59	LAC 99			
Clostridium barati		25.72	0.56				
Clostridium ramosum		21.33	0.56				
47373023	**Low discrimination**						**47373023**
					SPO		
Actinomyces naeslundii		54.16	0.59	ESL 12	1		
Clostridium tertium		12.86	0.25	RAF 92	99		
Clostridium barati		12.07	0.21		99		
47373030	**Low discrimination**						**47373030**
					SPO		
Clostridium tertium		78.09	0.55	LAC 99	99		
Lactobacillus jensenii		20.54	0.47		1		
Clostridium barati		0.42	0.13		99		
47373032	**Good identification**						**47373032**
					SPO		
Clostridium tertium		91.40	0.63	LAC 99	99		
Lactobacillus jensenii		5.34	0.44		1		
Bifidobacterium breve		1.32	0.41		1		
47373040	**Low discrimination**						**47373040**
					SPO		
Actinomyces israelii		57.72	0.52	XYL 92	1		
Lactobacillus jensenii		20.66	0.32		1		
Clostridium barati		9.91	0.21		99		
47373041	**Acceptable identification**						**47373041**
Actinomyces naeslundii		93.37	0.46	ESL 12			
Actinomyces israelii		6.63	0.18	RAF 92			
47373042	**Low discrimination**						**47373042**
					SPO		
Actinomyces israelii		37.60	0.55	XYL 92	1		
Lactobacillus acidophilus		31.50	0.38		1		
Clostridium barati		12.91	0.29		99		
47373043	**Low discrimination**						**47373043**
Actinomyces naeslundii		83.63	0.46	ESL 12			
Actinomyces israelii		8.91	0.21	RAF 92			
Lactobacillus acidophilus		7.46	0.04				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47373050	**Low discrimination**						**47373050**
					SPO		
Actinomyces israelii		44.80	0.38	XYL 92	1		
Lactobacillus jensenii		35.77	0.24	MLZ 13	1		
Clostridium tertium		11.33	0.13		99		
47373052	**Low discrimination**						**47373052**
					SPO		
Bifidobacterium breve		37.78	0.37	RAF 95	1		
Actinomyces israelii		31.02	0.41	LAC 99	1		
Clostridium tertium		15.69	0.21		99		
47373061	**Good identification**						**47373061**
Actinomyces naeslundii		100.0	0.46	ESL 12			
47373062	**Low discrimination**						**47373062**
					SPO		
Clostridium barati		39.73	0.37	TRE 8	99		
Lactobacillus acidophilus		32.30	0.38	MAN 8	1		
Bifidobacterium breve		12.48	0.41	SOR 8	1		
47373063	**Low discrimination**						**47373063**
					SPO		
Actinomyces naeslundii		83.40	0.46	ESL 12	1		
Clostridium barati		9.15	0.03	RAF 92	99		
Lactobacillus acidophilus		7.44	0.04		1		
47373072	**Low discrimination**						**47373072**
					SPO		
Bifidobacterium breve		54.65	0.41	RAF 95	1		
Clostridium tertium		38.31	0.29	LAC 99	99		
Lactobacillus jensenii		2.24	0.10		1		
47373100	**Low discrimination**						**47373100**
					COC	SPO	
Clostridium tertium		79.85	0.77		1	99	
Atopobium parvulum		8.91	0.61		99	1	
Actinomyces israelii		3.94	0.69		1	1	
47373101	**Low discrimination**						**47373101**
					COC	SPO	
Actinomyces naeslundii		46.81	0.67	ESL 12	1	1	
Clostridium tertium		42.96	0.43	RAF 92	1	99	
Atopobium parvulum		4.79	0.27		99	1	
47373102	**Low discrimination**						**47373102**
					SPO		
Clostridium tertium		75.58	0.85		99		
Lactobacillus acidophilus		14.13	0.72		1		
Bifidobacterium breve		3.27	0.71		1		
47373103	**Low discrimination**						**47373103**
					SPO		
Clostridium tertium		59.45	0.51	URE 1	99		
Actinomyces naeslundii		21.59	0.67		1		
Lactobacillus acidophilus		11.12	0.38		1		
47373110	**Very good identification**						**47373110**
					GRA	SPO	
Clostridium tertium		98.21	0.81		99	99	
Bifidobacterium breve		0.71	0.54		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		0.52	0.32		1	1	
47373111	**Good identification**						**47373111**
					SPO		
Clostridium tertium		95.34	0.47	URE 1	99		
Actinomyces naeslundii		3.08	0.45		1		
Bifidobacterium breve		0.69	0.20		1		
47373112	**Low discrimination**						**47373112**
					GRA	SPO	
Clostridium tertium		83.02	0.89		99	99	
Leptotrichia buccalis		14.53	0.66		1	1	
Bifidobacterium breve		2.02	0.71		99	1	
47373113	**Low discrimination**						**47373113**
					GRA	SPO	
Clostridium tertium		82.32	0.55	URE 1	99	99	
Leptotrichia buccalis		14.41	0.32		1	1	
Bifidobacterium breve		2.00	0.37		99	1	
47373120	**Low discrimination**						**47373120**
Clostridium tertium		58.99	0.85				
Clostridium ramosum		34.42	0.79				
Clostridium paraputrificum		3.61	0.56				
47373121	**Low discrimination**						**47373121**
					SPO		
Clostridium tertium		48.66	0.51	URE 1	99		
Clostridium ramosum		28.40	0.45		92		
Actinomyces naeslundii		17.68	0.67		1		
47373122	**Low discrimination**						**47373122**
					SPO		
Clostridium tertium		52.49	0.93		99		
Clostridium ramosum		40.83	0.90		92		
Lactobacillus acidophilus		3.27	0.72		1		
47373123	**Low discrimination**						**47373123**
					SPO		
Clostridium tertium		49.38	0.59	URE 1	99		
Clostridium ramosum		38.42	0.56		92		
Actinomyces naeslundii		5.98	0.67		1		
47373130	**Very good identification**						**47373130**
					SPO		
Clostridium tertium		99.15	0.89		99		
Bifidobacterium breve		0.42	0.58		1		
Clostridium ramosum		0.33	0.45		92		
47373131	**Good identification**						**47373131**
					SPO		
Clostridium tertium		98.20	0.55	URE 1	99		
Actinomyces naeslundii		1.06	0.45		1		
Bifidobacterium breve		0.42	0.24		1		
47373132	**Very good identification**						**47373132**
					GRA	SPO	
Clostridium tertium		97.47	0.98		99	99	
Bifidobacterium breve		1.41	0.75		99	1	
Leptotrichia buccalis		0.63	0.50		1	1	
47373133	**Very good identification**						**47373133**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium tertium		97.16	0.63	URE 1	99	99	
Bifidobacterium breve		1.40	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
47373140	**Low discrimination**						**47373140**
					SPO		
Actinomyces israelii		39.78	0.62	XYL 92	1		
Bifidobacterium breve		26.58	0.54		1		
Clostridium tertium		20.93	0.43		99		
47373141	**Low discrimination**						**47373141**
Actinomyces naeslundii		82.03	0.54	ESL 12			
Actinomyces israelii		8.19	0.28	RAF 92			
Bifidobacterium breve		5.47	0.20				
47373142	**Low discrimination**						**47373142**
					SPO		
Bifidobacterium breve		48.26	0.71	RAF 95	1		
Lactobacillus acidophilus		25.80	0.56		1		
Clostridium tertium		11.28	0.51		99		
47373143	**Low discrimination**						**47373143**
Actinomyces naeslundii		42.00	0.54	ESL 12			
Bifidobacterium breve		28.33	0.37	RAF 92			
Lactobacillus acidophilus		15.15	0.22				
47373150	**Low discrimination**						**47373150**
					SPO		
Clostridium tertium		53.03	0.47	SOR 1	99		
Bifidobacterium breve		37.88	0.54		1		
Actinomyces israelii		8.47	0.48		1		
47373152	**Low discrimination**						**47373152**
					GRA	SPO	
Bifidobacterium breve		65.13	0.71	RAF 95	99	1	
Clostridium tertium		27.06	0.55		99	99	
Leptotrichia buccalis		4.74	0.32		1	1	
47373153	**Low discrimination**						**47373153**
					SPO		
Bifidobacterium breve		62.53	0.37	RAF 95	1		
Clostridium tertium		25.97	0.21	URE 1	99		
Actinomyces naeslundii		4.88	0.32		1		
47373160	**Low discrimination**						**47373160**
					SPO		
Clostridium tertium		31.71	0.51	SOR 1	99		
Bifidobacterium breve		23.86	0.58		1		
Clostridium ramosum		18.50	0.45		92		
47373161	**Low discrimination**						**47373161**
					SPO		
Actinomyces naeslundii		67.64	0.54	ESL 12	1		
Clostridium tertium		10.66	0.17	RAF 92	99		
Bifidobacterium breve		8.02	0.24		1		
47373162	**Low discrimination**						**47373162**
					SPO		
Bifidobacterium breve		46.78	0.75	RAF 95	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		18.44	0.59		99		
Clostridium ramosum		14.35	0.56		92		
47373163	**Low discrimination**						**47373163**
					SPO		
Bifidobacterium breve		33.81	0.41	RAF 95	1		
Actinomyces naeslundii		28.20	0.54	URE 1	1		
Clostridium tertium		13.33	0.25		99		
47373170	**Low discrimination**						**47373170**
					SPO		
Clostridium tertium		69.68	0.55	SOR 1	99		
Bifidobacterium breve		29.49	0.58		1		
Actinomyces israelii		0.24	0.27		1		
47373172	**Low discrimination**						**47373172**
					GRA	SPO	
Bifidobacterium breve		58.33	0.75	RAF 95	99	1	
Clostridium tertium		40.89	0.63		99	99	
Leptotrichia buccalis		0.27	0.15		1	1	
47373173	**Low discrimination**						**47373173**
					SPO		
Bifidobacterium breve		57.31	0.41	RAF 95	1		
Clostridium tertium		40.17	0.29	URE 1	99		
Actinomyces naeslundii		2.52	0.32		1		
47373220	**Low discrimination**						**47373220**
					SPO		
Clostridium ramosum		87.65	0.47	LAC 99	92		
Clostridium barati		9.24	0.29		99		
Lactobacillus jensenii		1.61	0.21		1		
47373222	**Low discrimination**						**47373222**
					SPO		
Clostridium ramosum		88.30	0.58	LAC 99	92		
Clostridium barati		6.98	0.37		99		
Lactobacillus acidophilus		3.46	0.34		1		
47373242	**Low discrimination**						**47373242**
					SPO		
Actinomyces israelii		44.75	0.35	XYL 92	1		
Lactobacillus acidophilus		37.49	0.19	RHA 7	1		
Clostridium barati		17.76	0.11		99		
47373300	**Low discrimination**						**47373300**
					SPO		
Clostridium ramosum		56.29	0.47	NAG 99	92		
Clostridium tertium		24.26	0.43		99		
Actinomyces israelii		8.92	0.50		1		
47373302	**Low discrimination**						**47373302**
					SPO		
Clostridium ramosum		51.78	0.58	NAG 99	92		
Lactobacillus acidophilus		23.32	0.53		1		
Clostridium tertium		16.74	0.51		99		
47373310	**Acceptable identification**						**47373310**
					SPO		
Clostridium tertium		94.64	0.47	RHA 1	99		
Actinomyces israelii		2.93	0.36		1		
Clostridium ramosum		1.25	0.13		92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47373312	**Low discrimination**						**47373312**
					GRA	SPO	
Clostridium tertium		80.39	0.55	RHA 1	99	99	
Leptotrichia buccalis		14.07	0.32		1	1	
Bifidobacterium breve		1.96	0.37		99	1	
47373320	**Very good identification**						**47373320**
Clostridium ramosum		98.27	0.82				
Clostridium tertium		1.28	0.51				
Clostridium barati		0.31	0.37				
47373321	**Good identification**						**47373321**
					SPO		
Clostridium ramosum		97.94	0.47	URE 1	92		
Clostridium tertium		1.28	0.17		99		
Actinomyces naeslundii		0.47	0.33		1		
47373322	**Very good identification**						**47373322**
					SPO		
Clostridium ramosum		98.31	0.92		92		
Clostridium tertium		0.96	0.59		99		
Lactobacillus acidophilus		0.45	0.53		1		
47373323	**Very good identification**						**47373323**
					SPO		
Clostridium ramosum		98.24	0.58	URE 1	92		
Clostridium tertium		0.96	0.25		99		
Lactobacillus acidophilus		0.45	0.18		1		
47373330	**Low discrimination**						**47373330**
					SPO		
Clostridium tertium		69.38	0.55	RHA 1	99		
Clostridium ramosum		30.19	0.47		92		
Bifidobacterium breve		0.30	0.24		1		
47373332	**Low discrimination**						**47373332**
					SPO		
Clostridium tertium		62.28	0.63	RHA 1	99		
Clostridium ramosum		36.13	0.58		92		
Bifidobacterium breve		0.90	0.41		1		
47373340	**Low discrimination**						**47373340**
					SPO		
Actinomyces israelii		66.25	0.43	XYL 92	1		
Clostridium ramosum		10.86	0.13	RHA 7	92		
Clostridium barati		9.86	0.11		99		
47373342	**Low discrimination**						**47373342**
Lactobacillus acidophilus		49.26	0.37	MAN 5			
Actinomyces israelii		20.46	0.46	RHA 7			
Bifidobacterium breve		12.37	0.37	SOR 11			
47373352	**Low discrimination**						**47373352**
					SPO		
Bifidobacterium breve		56.32	0.37	RAF 95	1		
Clostridium tertium		23.40	0.21	RHA 1	99		
Actinomyces israelii		13.92	0.32		1		
47373360	**Acceptable identification**						**47373360**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		94.12	0.47	SOR 1		
Clostridium barati		2.59	0.19			
Clostridium tertium		1.23	0.17			
47373362	**Low discrimination**					**47373362**
					SPO	
Clostridium ramosum		89.84	0.58	SOR 1	92	
Lactobacillus acidophilus		5.00	0.37		1	
Bifidobacterium breve		2.23	0.41		1	
47373372	**Low discrimination**					**47373372**
					SPO	
Bifidobacterium breve		46.94	0.41	RAF 95	1	
Clostridium tertium		32.91	0.29	RHA 1	99	
Clostridium ramosum		19.09	0.24		92	
47373400	**Low discrimination**					**47373400**
					SPO	
Actinomyces israelii		61.63	0.57	XYL 92	1	
Clostridium innocuum		24.18	0.35		99	
Lactobacillus jensenii		10.48	0.32		1	
47373402	**Low discrimination**					**47373402**
					SPO	
Actinomyces israelii		69.05	0.60	XYL 92	1	
Clostridium innocuum		18.06	0.35		99	
Lactobacillus jensenii		5.22	0.29		1	
47373410	**Low discrimination**					**47373410**
					SPO	
Actinomyces israelii		62.15	0.43	XYL 92	1	
Lactobacillus jensenii		23.58	0.24	MLZ 13	1	
Clostridium tertium		7.47	0.13		99	
47373412	**Low discrimination**					**47373412**
					SPO	
Actinomyces israelii		53.04	0.46	XYL 92	1	
Clostridium tertium		12.75	0.21	MLZ 13	99	
Bifidobacterium adolescentis		12.52	0.29		1	
47373420	**Low discrimination**					**47373420**
					SPO	
Actinomyces israelii		33.54	0.37	XYL 92	1	
Lactobacillus jensenii		22.35	0.21	NAG 6	1	
Clostridium tertium		15.93	0.17		99	
47373422	**Low discrimination**					**47373422**
					SPO	
Actinomyces israelii		22.59	0.40	XYL 92	1	
Clostridium tertium		21.46	0.25	NAG 6	99	
Clostridium barati		20.13	0.21		99	
47373440	**Good identification**					**47373440**
					SPO	
Actinomyces israelii		98.11	0.50	XYL 92	1	
Clostridium innocuum		1.00	0.01		99	
Clostridium butyricum		0.89	-.06		99	
47373442	**Good identification**					**47373442**
Actinomyces israelii		97.44	0.53	XYL 92		
Lactobacillus acidophilus		1.01	0.04			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		0.89	0.12			
47373450	**Acceptable identification**					**47373450**
					COC	
Actinomyces israelii		92.42	0.36	XYL 92	1	
Blautia producta		7.58	-.03	MLZ 13	99	
47373452	**Acceptable identification**					**47373452**
Actinomyces israelii		93.02	0.39	XYL 92		
Bifidobacterium adolescentis		6.98	0.14	MLZ 13		
47373500	**Low discrimination**					**47373500**
					SPO	
Actinomyces israelii		64.34	0.68	XYL 92	1	
Clostridium tertium		16.09	0.43		99	
Clostridium butyricum		15.65	0.36		99	
47373502	**Low discrimination**					**47373502**
					SPO	
Actinomyces israelii		41.16	0.71	XYL 92	1	
Bifidobacterium adolescentis		29.39	0.62		1	
Clostridium tertium		20.59	0.51		99	
47373503	**Low discrimination**					**47373503**
					SPO	
Actinomyces israelii		39.17	0.36	XYL 92	1	
Bifidobacterium adolescentis		27.97	0.28	URE 1	1	
Clostridium tertium		19.60	0.17		99	
47373510	**Low discrimination**					**47373510**
					SPO	
Clostridium tertium		64.58	0.47	ARA 1	99	
Actinomyces israelii		21.70	0.53		1	
Bifidobacterium dentium		8.95	0.23		1	
47373512	**Low discrimination**					**47373512**
					SPO	
Clostridium tertium		35.43	0.55	ARA 1	99	
Bifidobacterium adolescentis		34.77	0.63		1	
Bifidobacterium dentium		16.55	0.40		1	
47373520	**Low discrimination**					**47373520**
					SPO	
Clostridium tertium		55.32	0.51	ARA 1	99	
Clostridium ramosum		32.28	0.45		92	
Actinomyces israelii		4.71	0.47		1	
47373522	**Low discrimination**					**47373522**
					SPO	
Clostridium tertium		51.01	0.59	ARA 1	99	
Clostridium ramosum		39.69	0.56		92	
Lactobacillus acidophilus		3.18	0.38		1	
47373530	**Acceptable identification**					**47373530**
					SPO	
Clostridium tertium		91.10	0.55	ARA 1	99	
Bifidobacterium dentium		7.48	0.27		1	
Actinomyces israelii		0.65	0.33		1	
47373532	**Low discrimination**					**47373532**
					SPO	
Clostridium tertium		76.34	0.63	ARA 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		21.13	0.44		1		
Bifidobacterium breve		1.10	0.41		1		
47373540	**Low discrimination**						**47373540**
					COC	SPO	
Actinomyces israelii		85.21	0.61	XYL 92	1	1	
Blautia producta		8.61	0.24		99	1	
Clostridium butyricum		4.63	0.18		1	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47373542	**Low discrimination**					**47373542**
Actinomyces israelii		74.07	0.64	XYL 92		
Bifidobacterium adolescentis		16.81	0.46			
Bifidobacterium breve		4.12	0.37			
47373552	**Low discrimination**					**47373552**
Bifidobacterium adolescentis		51.10	0.48	RAF 92		
Actinomyces israelii		27.53	0.49	XYL 87		
Bifidobacterium breve		10.25	0.37	SOR 11		
47373560	**Low discrimination**					**47373560**
					SPO	
Actinomyces israelii		45.93	0.40	XYL 92	1	
Clostridium tertium		14.03	0.17	NAG 6	99	
Bifidobacterium breve		10.55	0.24		1	
47373562	**Low discrimination**					**47373562**
					SPO	
Bifidobacterium breve		35.65	0.41	RAF 95	1	
Actinomyces israelii		23.02	0.43	ARA 1	1	
Clostridium tertium		14.06	0.25		99	
47373572	**Low discrimination**					**47373572**
					SPO	
Bifidobacterium breve		48.90	0.41	RAF 95	1	
Clostridium tertium		34.28	0.29	ARA 1	99	
Bifidobacterium dentium		9.49	0.10		1	
47373600	**Acceptable identification**					**47373600**
					SPO	
Actinomyces israelii		95.00	0.38	XYL 92	1	
Clostridium innocuum		5.00	0.01	RHA 7	99	
47373602	**Acceptable identification**					**47373602**
					SPO	
Actinomyces israelii		90.14	0.41	XYL 92	1	
Clostridium clostridioforme		3.77	0.08	RHA 7	66	
Clostridium innocuum		3.16	0.01		99	
47373700	**Low discrimination**					**47373700**
					SPO	
Actinomyces israelii		82.55	0.48	XYL 92	1	
Clostridium butyricum		8.25	0.10	RHA 7	99	
Clostridium ramosum		6.43	0.13		92	
47373702	**Low discrimination**					**47373702**
					SPO	
Actinomyces israelii		64.87	0.51	XYL 92	1	
Clostridium ramosum		13.47	0.24	RHA 7	92	
Bifidobacterium adolescentis		6.22	0.28		1	
47373720	**Good identification**					**47373720**
					SPO	
Clostridium ramosum		97.61	0.47	ARA 1	92	
Clostridium tertium		1.28	0.17		99	
Actinomyces israelii		0.81	0.28		1	
47373722	**Very good identification**					**47373722**
					SPO	
Clostridium ramosum		97.84	0.58	ARA 1	92	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium tertium		0.96	0.25		99			
Lactobacillus acidophilus		0.45	0.18		1			
47373742	**Low discrimination**							**47373742**
					COC			
Actinomyces israelii		86.23	0.44	XYL 92	1			
Blautia producta		8.58	0.07	RHA 7	99			
Bifidobacterium adolescentis		2.63	0.12		1			
47374100	**Very good identification**							**47374100**
					COC	GRA	SPO	
Atopobium parvulum		97.16	0.66	GAL 99	99	99	1	
Clostridium butyricum		1.50	0.28		1	99	99	
Fusobacterium mortiferum		0.66	0.40		1	1	1	
47374102	**Low discrimination**							**47374102**
					COC	SPO		
Atopobium parvulum		85.81	0.53	GAL 99	99	1		
Actinomyces israelii		3.56	0.43		1	1		
Clostridium butyricum		2.93	0.21		1	99		
47374120	**Acceptable identification**							**47374120**
					COC	SPO		
Atopobium parvulum		91.91	0.53	GAL 99	99	1		
Clostridium paraputrificum		4.66	0.22		1	99		
Clostridium barati		1.31	0.18		1	99		
47374122	**Low discrimination**							**47374122**
					COC	SPO		
Atopobium parvulum		52.62	0.40	GAL 99	99	1		
Clostridium barati		12.76	0.26		1	99		
Clostridium tertium		10.34	0.27		1	99		
47374142	**Low discrimination**							**47374142**
					COC			
Actinomyces israelii		44.93	0.36	GAL 99	1			
Atopobium parvulum		28.11	0.19		99			
Lactobacillus acidophilus		10.23	0.10		1			
47374212	**Good identification**							**47374212**
Clostridium clostridioforme		100.0	0.42	RAF 94				
47374232	**Good identification**							**47374232**
Clostridium clostridioforme		100.0	0.42	RAF 94				
47374302	**Low discrimination**							**47374302**
					COC			
Clostridium clostridioforme		83.94	0.40	RAF 94	1			
Atopobium parvulum		9.13	0.19	ARA 94	99			
Actinomyces israelii		2.83	0.24		1			
47374312	**Good identification**							**47374312**
Clostridium clostridioforme		99.39	0.51	RAF 94				
Leptotrichia buccalis		0.61	-.03	ARA 94				
47374322	**Low discrimination**							**47374322**
					SPO			
Clostridium clostridioforme		80.36	0.40	RAF 94	66			
Parabacteroides distasonis		6.92	0.27	ARA 94	1			
Clostridium ramosum		3.96	0.11		92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47374332	**Good identification**						**47374332**
Clostridium clostridioforme		98.99	0.51	RAF 94			
Parabacteroides distasonis		1.01	0.23	ARA 94			
47374402	**Low discrimination**						**47374402**
					SPO		
Clostridium clostridioforme		72.68	0.36	RAF 94	66		
Clostridium butyricum		15.70	0.13	RHA 88	99		
Actinomyces israelii		10.42	0.31		1		
47374412	**Good identification**						**47374412**
Clostridium clostridioforme		99.09	0.48	RAF 94			
Actinomyces israelii		0.47	0.17	RHA 88			
Bifidobacterium adolescentis		0.44	0.11				
47374422	**Good identification**						**47374422**
Clostridium clostridioforme		98.16	0.36	RAF 94			
Clostridium butyricum		1.84	-.05	RHA 88			
47374432	**Good identification**						**47374432**
Clostridium clostridioforme		100.0	0.48	RAF 94			
47374500	**Acceptable identification**						**47374500**
					COC	SPO	
Clostridium butyricum		90.50	0.44	RAF 95	1	99	
Atopobium parvulum		6.56	0.32	GAL 99	99	1	
Actinomyces israelii		2.60	0.39		1	1	
47374502	**Low discrimination**						**47374502**
					SPO		
Clostridium butyricum		47.45	0.37	RAF 95	99		
Clostridium clostridioforme		30.64	0.45	GAL 99	66		
Bifidobacterium adolescentis		15.10	0.43		1		
47374512	**Low discrimination**						**47374512**
					SPO		
Clostridium clostridioforme		85.74	0.57	RAF 94	66		
Bifidobacterium adolescentis		11.34	0.45	RHA 88	1		
Clostridium butyricum		1.53	0.15		99		
47374522	**Low discrimination**						**47374522**
					SPO		
Clostridium clostridioforme		85.63	0.45	RAF 94	66		
Clostridium butyricum		11.53	0.19	RHA 88	99		
Actinomyces israelii		0.94	0.21		1		
47374532	**Very good identification**						**47374532**
					SPO		
Clostridium clostridioforme		98.28	0.57	RAF 94	66		
Bifidobacterium dentium		1.35	0.10	RHA 88	1		
Clostridium butyricum		0.15	-.03		99		
47374602	**Good identification**						**47374602**
Clostridium clostridioforme		100.0	0.51	RAF 94			
47374612	**Very good identification**						**47374612**
Clostridium clostridioforme		100.0	0.62	RAF 94			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47374622	**Good identification**					**47374622**
Clostridium clostridioforme		100.0	0.51	RAF 94		
47374632	**Very good identification**					**47374632**
Clostridium clostridioforme		100.0	0.62	RAF 94		
47374702	**Very good identification**					**47374702**
					SPO	
Clostridium clostridioforme		99.11	0.60	RAF 94	66	
Clostridium butyricum		0.65	0.11		99	
Actinomyces israelii		0.17	0.22		1	
47374710	**Good identification**					**47374710**
Clostridium clostridioforme		98.12	0.37	RAF 94		
Clostridium butyricum		1.88	-.04	BGL 99		
47374712	**Very good identification**					**47374712**
Clostridium clostridioforme		99.98	0.71	RAF 94		
Bifidobacterium adolescentis		0.02	0.11			
47374713	**Good identification**					**47374713**
Clostridium clostridioforme		100.0	0.37	RAF 94		
47374722	**Very good identification**					**47374722**
Clostridium clostridioforme		99.34	0.60	RAF 94		
Parabacteroides distasonis		0.34	0.23			
Bacteroides ovatus		0.27	0.06			
47374730	**Good identification**					**47374730**
Clostridium clostridioforme		100.0	0.37	RAF 94		
47374732	**Very good identification**					**47374732**
Clostridium clostridioforme		99.92	0.71	RAF 94		
Parabacteroides distasonis		0.04	0.19			
Bacteroides ovatus		0.04	0.03			
47374733	**Good identification**					**47374733**
Clostridium clostridioforme		100.0	0.37	RAF 94		
47374752	**Good identification**					**47374752**
Clostridium clostridioforme		100.0	0.37	RAF 94		
47374772	**Good identification**					**47374772**
Clostridium clostridioforme		100.0	0.37	RAF 94		
47375000	**Low discrimination**					**47375000**
					COC	
Atopobium parvulum		49.45	0.66	LAC 99	99	
Lactobacillus jensenii		40.58	0.66		1	
Actinomyces israelii		5.98	0.64		1	
47375002	**Low discrimination**					**47375002**
					COC	
Lactobacillus jensenii		54.48	0.63	XYL 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		18.06	0.67		1		
Atopobium parvulum		17.57	0.53		99		
47375010	**Low discrimination**						**47375010**
					COC		
Lactobacillus jensenii		89.18	0.58	XYL 1	1		
Actinomyces israelii		5.89	0.50		1		
Atopobium parvulum		3.29	0.32		99		
47375012	**Low discrimination**						**47375012**
					GRA		
Lactobacillus jensenii		66.42	0.55	XYL 1	99		
Leptotrichia buccalis		16.58	0.32		1		
Actinomyces israelii		9.87	0.53		99		
47375020	**Low discrimination**						**47375020**
					COC	SPO	
Lactobacillus jensenii		48.31	0.56	XYL 1	1	1	
Atopobium parvulum		41.55	0.53		99	1	
Clostridium barati		3.75	0.31		1	99	
47375022	**Low discrimination**						**47375022**
					SPO		
Lactobacillus jensenii		46.30	0.53	XYL 1	1		
Clostridium barati		16.19	0.40		99		
Lactobacillus acidophilus		15.30	0.41		1		
47375030	**Acceptable identification**						**47375030**
					COC	SPO	
Lactobacillus jensenii		90.70	0.47	XYL 1	1	1	
Clostridium tertium		4.81	0.24		1	99	
Atopobium parvulum		2.36	0.19		99	1	
47375032	**Low discrimination**						**47375032**
					SPO		
Lactobacillus jensenii		61.70	0.44	XYL 1	1		
Clostridium tertium		14.73	0.32		99		
Clostridium clostridioforme		12.34	0.35		66		
47375040	**Low discrimination**						**47375040**
					COC		
Actinomyces israelii		65.82	0.57		1		
Atopobium parvulum		14.14	0.32		99		
Lactobacillus jensenii		11.61	0.32		1		
47375042	**Low discrimination**						**47375042**
Actinomyces israelii		80.80	0.60				
Lactobacillus acidophilus		6.40	0.26				
Lactobacillus jensenii		6.33	0.29				
47375050	**Low discrimination**						**47375050**
					SPO		
Actinomyces israelii		69.69	0.43	MLZ 13	1		
Lactobacillus jensenii		27.41	0.24		1		
Clostridium butyricum		2.90	-.02		99		
47375052	**Low discrimination**						**47375052**
					GRA		
Actinomyces israelii		79.65	0.46	MLZ 13	99		
Lactobacillus jensenii		13.93	0.21		99		
Leptotrichia buccalis		3.48	-.03		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47375060	**Low discrimination**						**47375060**
					COC		
Actinomyces israelii		31.93	0.37	NAG 6	1		
Lactobacillus jensenii		22.05	0.21		1		
Atopobium parvulum		18.97	0.19		99		
47375100	**Very good identification**						**47375100**
					COC	SPO	
Atopobium parvulum		97.82	1.00		99	1	
Clostridium butyricum		1.52	0.62		1	99	
Actinomyces israelii		0.48	0.74		1	1	
47375101	**Very good identification**						**47375101**
					COC	SPO	
Atopobium parvulum		97.84	0.66	URE 1	99	1	
Clostridium butyricum		1.52	0.28		1	99	
Actinomyces israelii		0.48	0.40		1	1	
47375102	**Low discrimination**						**47375102**
					COC	SPO	
Atopobium parvulum		87.76	0.87		99	1	
Actinomyces israelii		3.64	0.77		1	1	
Clostridium butyricum		3.00	0.55		1	99	
47375103	**Low discrimination**						**47375103**
					COC	SPO	
Atopobium parvulum		87.84	0.53	URE 1	99	1	
Actinomyces israelii		3.65	0.43		1	1	
Clostridium butyricum		3.00	0.21		1	99	
47375110	**Low discrimination**						**47375110**
					COC	SPO	
Atopobium parvulum		75.43	0.66	MLZ 1	99	1	
Clostridium tertium		9.03	0.50		1	99	
Clostridium butyricum		6.09	0.40		1	99	
47375112	**Low discrimination**						**47375112**
					GRA	SPO	
Leptotrichia buccalis		81.35	0.66	XYL 1	1	1	
Clostridium tertium		6.48	0.58		99	99	
Bifidobacterium adolescentis		5.63	0.65		99	1	
47375120	**Good identification**						**47375120**
					COC	SPO	
Atopobium parvulum		92.77	0.87		99	1	
Clostridium paraputrificum		4.70	0.56		1	99	
Clostridium tertium		1.07	0.53		1	99	
47375121	**Acceptable identification**						**47375121**
					COC	SPO	
Atopobium parvulum		92.76	0.53	URE 1	99	1	
Clostridium paraputrificum		4.70	0.22		1	99	
Clostridium tertium		1.07	0.19		1	99	
47375122	**Low discrimination**						**47375122**
					COC	SPO	
Atopobium parvulum		60.85	0.74		99	1	
Clostridium tertium		11.96	0.62		1	99	
Lactobacillus acidophilus		10.26	0.60		1	1	
47375123	**Low discrimination**						**47375123**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium parvulum		60.91	0.40	URE 1	99	1		
Clostridium tertium		11.97	0.27		1	99		
Lactobacillus acidophilus		10.27	0.25		1	1		
47375130	**Low discrimination**							**47375130**
					COC	SPO		
Clostridium tertium		63.60	0.58	MAN 99	1	99		
Atopobium parvulum		31.25	0.53		99	1		
Clostridium paraputrificum		1.58	0.22		1	99		
47375132	**Low discrimination**							**47375132**
					COC	GRA	SPO	
Clostridium tertium		63.24	0.66	MAN 99	1	99	99	
Leptotrichia buccalis		29.39	0.50		1	1	1	
Atopobium parvulum		1.83	0.40		99	99	1	
47375140	**Low discrimination**							**47375140**
					COC	SPO		
Atopobium parvulum		75.40	0.66	SOR 1	99	1		
Actinomyces israelii		14.18	0.67		1	1		
Clostridium butyricum		10.05	0.44		1	99		
47375142	**Low discrimination**							**47375142**
					COC			
Actinomyces israelii		45.54	0.70		1			
Atopobium parvulum		28.49	0.53		99			
Lactobacillus acidophilus		10.37	0.44		1			
47375143	**Low discrimination**							**47375143**
					COC			
Actinomyces israelii		46.53	0.36	URE 1	1			
Atopobium parvulum		29.11	0.19		99			
Lactobacillus acidophilus		10.60	0.10		1			
47375150	**Low discrimination**							**47375150**
					COC	SPO		
Actinomyces israelii		58.38	0.53	MLZ 13	1	1		
Atopobium parvulum		20.99	0.32		99	1		
Clostridium butyricum		14.58	0.22		1	99		
47375160	**Low discrimination**							**47375160**
					COC	SPO		
Atopobium parvulum		64.06	0.53	SOR 1	99	1		
Clostridium paraputrificum		24.19	0.37		1	99		
Actinomyces israelii		4.36	0.47		1	1		
47375162	**Low discrimination**							**47375162**
					COC	SPO		
Lactobacillus acidophilus		35.91	0.44	XYL 1	1	1		
Atopobium parvulum		17.41	0.40	SOR 11	99	1		
Clostridium paraputrificum		15.96	0.31		1	99		
47375172	**Low discrimination**							**47375172**
					GRA	SPO		
Bifidobacterium breve		34.79	0.41	RAF 95	99	1		
Clostridium tertium		33.68	0.32	XYL 1	99	99		
Leptotrichia buccalis		15.65	0.15		1	1		
47375202	**Low discrimination**							**47375202**
Clostridium clostridioforme		35.52	0.39	RAF 94				
Actinomyces israelii		33.02	0.48	ARA 94				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		13.37	0.29				
47375212	**Acceptable identification**						**47375212**
Clostridium clostridioforme		93.91	0.50	RAF 94			
Actinomyces israelii		2.86	0.34	ARA 94			
Lactobacillus jensenii		2.59	0.21				
47375222	**Low discrimination**						**47375222**
					SPO		
Clostridium clostridioforme		49.45	0.39	RAF 94	66		
Clostridium barati		14.01	0.21	ARA 94	99		
Lactobacillus acidophilus		11.46	0.22		1		
47375232	**Good identification**						**47375232**
					GRA		
Clostridium clostridioforme		98.02	0.50	RAF 94	50		
Parabacteroides distasonis		1.11	0.23	ARA 94	1		
Lactobacillus jensenii		0.68	0.10		99		
47375240	**Good identification**						**47375240**
					SPO		
Actinomyces israelii		95.37	0.38	RHA 7	1		
Clostridium butyricum		4.63	-.06		99		
47375242	**Acceptable identification**						**47375242**
Actinomyces israelii		90.32	0.41	RHA 7			
Lactobacillus acidophilus		7.16	0.07				
Clostridium clostridioforme		2.52	0.05				
47375300	**Good identification**						**47375300**
					COC	SPO	
Atopobium parvulum		91.29	0.66	RHA 1	99	1	
Clostridium butyricum		4.33	0.36		1	99	
Actinomyces israelii		3.32	0.55		1	1	
47375302	**Low discrimination**						**47375302**
					COC		
Atopobium parvulum		50.02	0.53	RHA 1	99		
Actinomyces israelii		15.47	0.58		1		
Clostridium clostridioforme		13.93	0.48		1		
47375312	**Low discrimination**						**47375312**
					GRA		
Clostridium clostridioforme		77.68	0.59	RAF 94	50		
Leptotrichia buccalis		15.79	0.32	ARA 94	1		
Actinomyces israelii		2.83	0.44		99		
47375320	**Low discrimination**						**47375320**
					COC	SPO	
Atopobium parvulum		78.60	0.53	RHA 1	99	1	
Clostridium ramosum		8.89	0.35		1	92	
Clostridium paraputrificum		3.98	0.22		1	99	
47375322	**Low discrimination**						**47375322**
					SPO		
Parabacteroides distasonis		34.01	0.61	RAF 99	1		
Clostridium ramosum		19.44	0.45		92		
Clostridium clostridioforme		11.97	0.48		66		
47375332	**Low discrimination**						**47375332**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		70.21	0.59	RAF 94	50	66	
Parabacteroides distasonis		23.58	0.57	ARA 94	1	1	
Clostridium tertium		3.41	0.32		99	99	
47375340	**Low discrimination**						**47375340**
					COC	SPO	
Actinomyces israelii		49.88	0.48	RHA 7	1	1	
Atopobium parvulum		35.60	0.32		99	1	
Clostridium butyricum		14.53	0.18		1	99	
47375342	**Low discrimination**						**47375342**
					COC		
Actinomyces israelii		69.32	0.51	RHA 7	1		
Lactobacillus acidophilus		15.79	0.25		1		
Atopobium parvulum		5.82	0.19		99		
47375352	**Low discrimination**						**47375352**
					GRA		
Actinomyces israelii		50.49	0.37	RHA 7	99		
Clostridium clostridioforme		36.00	0.25	MLZ 13	50		
Leptotrichia buccalis		7.32	-.03		1		
47375400	**Low discrimination**						**47375400**
					COC	SPO	
Clostridium butyricum		83.00	0.55	RAF 95	1	99	
Actinomyces israelii		14.28	0.62	LAC 96	1	1	
Atopobium parvulum		1.46	0.32		99	1	
47375402	**Low discrimination**						**47375402**
					SPO		
Clostridium butyricum		52.18	0.48	RAF 95	99		
Actinomyces israelii		34.62	0.66	LAC 96	1		
Clostridium clostridioforme		7.32	0.45		66		
47375412	**Low discrimination**						**47375412**
Clostridium clostridioforme		70.94	0.56	RAF 94			
Actinomyces israelii		11.00	0.51	RHA 88			
Bifidobacterium adolescentis		10.47	0.45				
47375420	**Low discrimination**						**47375420**
					SPO		
Clostridium butyricum		82.44	0.36	RAF 95	99		
Actinomyces israelii		10.41	0.42	LAC 96	1		
Lactobacillus jensenii		3.42	0.21	NAG 8	1		
47375422	**Low discrimination**						**47375422**
					SPO		
Clostridium clostridioforme		49.49	0.45	RAF 94	66		
Clostridium butyricum		30.68	0.29	RHA 88	99		
Actinomyces israelii		14.94	0.45		1		
47375432	**Good identification**						**47375432**
Clostridium clostridioforme		96.33	0.56	RAF 94			
Bifidobacterium dentium		1.47	0.10	RHA 88			
Actinomyces israelii		0.95	0.31				
47375440	**Low discrimination**						**47375440**
					SPO		
Clostridium butyricum		56.52	0.36	RAF 95	99		
Actinomyces israelii		43.48	0.55	LAC 96	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47375442	**Low discrimination**						**47375442**
					SPO		
Actinomyces israelii		72.49	0.58		1		
Clostridium butyricum		24.43	0.29		99		
Bifidobacterium adolescentis		2.68	0.28		1		
47375450	**Low discrimination**						**47375450**
					COC	SPO	
Actinomyces israelii		66.75	0.41	MLZ 13	1	1	
Clostridium butyricum		30.56	0.14		1	99	
Blautia producta		2.70	-.03		99	1	
47375452	**Low discrimination**						**47375452**
Actinomyces israelii		62.96	0.44	MLZ 13			
Bifidobacterium adolescentis		19.03	0.29				
Clostridium clostridioforme		10.54	0.22				
47375462	**Low discrimination**						**47375462**
					SPO		
Actinomyces israelii		64.72	0.38	NAG 6	1		
Clostridium butyricum		29.71	0.11		99		
Clostridium clostridioforme		5.57	0.10		66		
47375500	**Good identification**						**47375500**
					COC	SPO	
Clostridium butyricum		90.68	0.78	RAF 95	1	99	
Atopobium parvulum		6.57	0.66		99	1	
Actinomyces israelii		2.60	0.73		1	1	
47375501	**Acceptable identification**						**47375501**
					COC	SPO	
Clostridium butyricum		90.58	0.44	RAF 95	1	99	
Atopobium parvulum		6.57	0.32	URE 1	99	1	
Actinomyces israelii		2.60	0.39		1	1	
47375502	**Low discrimination**						**47375502**
					SPO		
Clostridium butyricum		67.42	0.71	RAF 95	99		
Bifidobacterium adolescentis		21.46	0.77		1		
Actinomyces israelii		7.46	0.76		1		
47375503	**Low discrimination**						**47375503**
					SPO		
Clostridium butyricum		67.51	0.37	RAF 95	99		
Bifidobacterium adolescentis		21.49	0.43	URE 1	1		
Actinomyces israelii		7.47	0.42		1		
47375510	**Low discrimination**						**47375510**
					SPO		
Clostridium butyricum		86.88	0.56	RAF 95	99		
Actinomyces israelii		7.07	0.59	MLZ 5	1		
Bifidobacterium adolescentis		2.52	0.45		1		
47375512	**Low discrimination**						**47375512**
					SPO		
Bifidobacterium adolescentis		67.13	0.79	RAF 92	1		
Clostridium clostridioforme		15.39	0.65		66		
Clostridium butyricum		9.08	0.49		99		
47375513	**Low discrimination**						**47375513**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		67.25	0.45	RAF 92	1		
Clostridium clostridioforme		15.41	0.31	URE 1	66		
Clostridium butyricum		9.10	0.15		99		
47375520	**Low discrimination**						**47375520**
					COC	SPO	
Clostridium butyricum		84.72	0.60	RAF 95	1	99	
Atopobium parvulum		12.46	0.53	NAG 8	99	1	
Actinomyces israelii		1.78	0.52		1	1	
47375522	**Low discrimination**						**47375522**
					SPO		
Clostridium butyricum		65.51	0.53	RAF 95	99		
Clostridium clostridioforme		14.74	0.54	NAG 8	66		
Actinomyces israelii		5.32	0.55		1		
47375530	**Low discrimination**						**47375530**
					SPO		
Clostridium butyricum		63.75	0.38	RAF 95	99		
Bifidobacterium dentium		21.57	0.27	MLZ 5	1		
Clostridium clostridioforme		4.88	0.31	NAG 8	66		
47375532	**Low discrimination**						**47375532**
					SPO		
Clostridium clostridioforme		61.36	0.65	RAF 94	66		
Bifidobacterium dentium		27.70	0.44	RHA 88	1		
Clostridium butyricum		3.15	0.31		99		
47375540	**Low discrimination**						**47375540**
					COC	SPO	
Clostridium butyricum		87.22	0.60	RAF 95	1	99	
Actinomyces israelii		11.19	0.66	SOR 8	1	1	
Atopobium parvulum		0.73	0.32		99	1	
47375542	**Low discrimination**						**47375542**
					SPO		
Clostridium butyricum		51.10	0.53	RAF 95	99		
Actinomyces israelii		25.28	0.69	SOR 8	1		
Bifidobacterium adolescentis		23.12	0.62		1		
47375550	**Low discrimination**						**47375550**
					COC	SPO	
Clostridium butyricum		56.18	0.38	RAF 95	1	99	
Actinomyces israelii		20.45	0.52	MLZ 5	1	1	
Blautia producta		20.45	0.32	SOR 8	99	1	
47375552	**Low discrimination**						**47375552**
					SPO		
Bifidobacterium adolescentis		79.11	0.63	RAF 92	1		
Actinomyces israelii		10.57	0.55	SOR 11	1		
Clostridium butyricum		7.53	0.31		99		
47375560	**Low discrimination**						**47375560**
					COC	SPO	
Clostridium butyricum		88.45	0.42	RAF 95	1	99	
Actinomyces israelii		8.33	0.45	NAG 8	1	1	
Atopobium parvulum		1.51	0.19	SOR 8	99	1	
47375602	**Good identification**						**47375602**
					SPO		
Clostridium clostridioforme		92.65	0.59	RAF 94	66		
Actinomyces israelii		4.50	0.46		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		2.79	0.22		99		
47375610	**Low discrimination**						**47375610**
					SPO		
Clostridium clostridioforme		83.79	0.36	RAF 94	66		
Actinomyces israelii		8.81	0.29	BGL 99	1		
Clostridium butyricum		7.41	0.07		99		
47375612	**Very good identification**						**47375612**
					SPO		
Clostridium clostridioforme		99.79	0.71	RAF 94	66		
Actinomyces israelii		0.16	0.32		1		
Clostridium butyricum		0.04	0.00		99		
47375613	**Good identification**						**47375613**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47375622	**Very good identification**						**47375622**
					GRA		
Clostridium clostridioforme		98.77	0.59	RAF 94	50		
Parabacteroides distasonis		0.37	0.23		1		
Actinomyces israelii		0.31	0.26		99		
47375630	**Good identification**						**47375630**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47375632	**Very good identification**						**47375632**
Clostridium clostridioforme		99.91	0.71	RAF 94			
Parabacteroides distasonis		0.04	0.19				
Bacteroides ovatus		0.04	0.03				
47375633	**Good identification**						**47375633**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47375640	**Low discrimination**						**47375640**
					SPO		
Actinomyces israelii		65.19	0.36	RHA 7	1		
Clostridium butyricum		34.81	0.11		99		
47375642	**Low discrimination**						**47375642**
					SPO		
Actinomyces israelii		59.75	0.39	RHA 7	1		
Clostridium clostridioforme		31.97	0.25		66		
Clostridium butyricum		8.27	0.04		99		
47375652	**Acceptable identification**						**47375652**
Clostridium clostridioforme		94.22	0.36	RAF 94			
Actinomyces israelii		5.78	0.25	SOR 1			
47375672	**Good identification**						**47375672**
Clostridium clostridioforme		100.0	0.36	RAF 94			
47375700	**Low discrimination**						**47375700**
					COC	SPO	
Clostridium butyricum		88.04	0.52	RAF 95	1	99	
Actinomyces israelii		6.14	0.54	RHA 3	1	1	
Atopobium parvulum		2.08	0.32		99	1	
47375702	**Low discrimination**						**47375702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium clostridioforme		76.76	0.68	RAF 94	66		
Clostridium butyricum		16.54	0.45		99		
Actinomyces israelii		4.45	0.57		1		
47375710	**Low discrimination**						**47375710**
					SPO		
Clostridium clostridioforme		55.35	0.45	RAF 94	66		
Clostridium butyricum		35.08	0.30	BGL 99	99		
Actinomyces israelii		6.95	0.39		1		
47375712	**Very good identification**						**47375712**
					SPO		
Clostridium clostridioforme		98.89	0.80	RAF 94	66		
Bifidobacterium adolescentis		0.59	0.45		1		
Clostridium butyricum		0.25	0.23		99		
47375713	**Good identification**						**47375713**
Clostridium clostridioforme		99.40	0.45	RAF 94			
Bifidobacterium adolescentis		0.60	0.11	URE 1			
47375722	**Low discrimination**						**47375722**
Clostridium clostridioforme		81.07	0.68	RAF 94			
Parabacteroides distasonis		9.01	0.58				
Bacteroides ovatus		7.22	0.40				
47375730	**Acceptable identification**						**47375730**
					GRA	SPO	
Clostridium clostridioforme		90.57	0.45	RAF 94	50	66	
Clostridium butyricum		4.99	0.12	BGL 99	99	99	
Parabacteroides distasonis		1.19	0.19		1	1	
47375732	**Very good identification**						**47375732**
Clostridium clostridioforme		97.31	0.80	RAF 94			
Parabacteroides distasonis		1.28	0.53				
Bacteroides ovatus		1.27	0.37				
47375733	**Good identification**						**47375733**
Clostridium clostridioforme		97.45	0.45	RAF 94			
Parabacteroides distasonis		1.28	0.19	URE 1			
Bacteroides ovatus		1.27	0.03				
47375742	**Low discrimination**						**47375742**
					SPO		
Actinomyces israelii		40.42	0.50	RHA 7	1		
Clostridium butyricum		33.58	0.27		99		
Clostridium clostridioforme		18.10	0.34		66		
47375752	**Low discrimination**						**47375752**
Clostridium clostridioforme		80.38	0.45	RAF 94			
Bifidobacterium adolescentis		5.91	0.29	SOR 1			
Actinomyces israelii		5.89	0.35				
47375772	**Acceptable identification**						**47375772**
Clostridium clostridioforme		92.63	0.45	RAF 94			
Bacteroides ovatus		4.98	0.13	SOR 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		1.22	0.19				
47376000	**Low discrimination**						**47376000**
					SPO		
Actinomyces israelii		87.59	0.43	GAL 99	1		
Clostridium tertium		8.61	0.11		99		
Clostridium innocuum		3.80	0.04		99		
47376002	**Low discrimination**						**47376002**
					SPO		
Actinomyces israelii		81.60	0.46	GAL 99	1		
Clostridium tertium		16.04	0.19		99		
Clostridium innocuum		2.36	0.04		99		
47376040	**Good identification**						**47376040**
Actinomyces israelii		100.0	0.36	GAL 99			
47376042	**Good identification**						**47376042**
Actinomyces israelii		100.0	0.39	GAL 99			
47376100	**Low discrimination**						**47376100**
					COC	SPO	
Clostridium tertium		63.03	0.45	GAL 99	1	99	
Actinomyces israelii		25.91	0.53		1	1	
Atopobium parvulum		9.46	0.32		99	1	
47376102	**Low discrimination**						**47376102**
					SPO		
Clostridium tertium		81.27	0.53	GAL 99	99		
Actinomyces israelii		16.70	0.56		1		
Bifidobacterium adolescentis		0.85	0.28		1		
47376110	**Good identification**						**47376110**
					SPO		
Clostridium tertium		96.66	0.50	GAL 99	99		
Actinomyces israelii		3.34	0.39		1		
47376112	**Very good identification**						**47376112**
					SPO		
Clostridium tertium		97.17	0.58	GAL 99	99		
Actinomyces israelii		1.68	0.42		1		
Bifidobacterium adolescentis		0.70	0.29		1		
47376120	**Good identification**						**47376120**
					COC	SPO	
Clostridium tertium		97.69	0.53	GAL 99	1	99	
Atopobium parvulum		0.86	0.19		99	1	
Actinomyces israelii		0.85	0.33		1	1	
47376122	**Very good identification**						**47376122**
					SPO		
Clostridium tertium		98.80	0.62	GAL 99	99		
Clostridium ramosum		0.56	0.21		92		
Actinomyces israelii		0.43	0.36		1		
47376130	**Very good identification**						**47376130**
					SPO		
Clostridium tertium		99.87	0.58	GAL 99	99		
Actinomyces israelii		0.07	0.18		1		
Bifidobacterium dentium		0.06	-.07		1		
47376132	**Very good identification**						**47376132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Clostridium tertium		99.76	0.66	GAL 99	99	
Bifidobacterium dentium		0.20	0.10		1	
Actinomyces israelii		0.04	0.21		1	
47376140	**Acceptable identification**					**47376140**
					SPO	
Actinomyces israelii		92.84	0.46	GAL 99	1	
Clostridium tertium		5.87	0.11		99	
Clostridium butyricum		1.29	-.07		99	
47376142	**Low discrimination**					**47376142**
					SPO	
Actinomyces israelii		87.53	0.49	GAL 99	1	
Clostridium tertium		11.06	0.19		99	
Bifidobacterium adolescentis		1.41	0.12		1	
47376302	**Low discrimination**					**47376302**
					SPO	
Actinomyces israelii		58.62	0.37	GAL 99	1	
Clostridium tertium		38.28	0.19	RHA 7	99	
Clostridium clostridioforme		3.11	0.06		66	
47376400	**Low discrimination**					**47376400**
					SPO	
Actinomyces israelii		89.77	0.41	GAL 99	1	
Clostridium butyricum		10.23	0.04		99	
47376402	**Good identification**					**47376402**
					SPO	
Actinomyces israelii		95.57	0.44	GAL 99	1	
Clostridium butyricum		2.82	-.03		99	
Bifidobacterium adolescentis		1.61	0.08		1	
47376442	**Good identification**					**47376442**
Actinomyces israelii		100.0	0.37	GAL 99		
47376500	**Low discrimination**					**47376500**
					SPO	
Actinomyces israelii		58.13	0.52	GAL 99	1	
Clostridium butyricum		39.76	0.28		99	
Clostridium tertium		1.75	0.11		99	
47376502	**Low discrimination**					**47376502**
					SPO	
Actinomyces israelii		58.84	0.55	GAL 99	1	
Bifidobacterium adolescentis		24.45	0.42		1	
Clostridium butyricum		10.43	0.21		99	
47376512	**Low discrimination**					**47376512**
Bifidobacterium dentium		81.33	0.40	RAF 95		
Bifidobacterium adolescentis		11.55	0.43	GAL 99		
Actinomyces israelii		3.40	0.40			
47376532	**Acceptable identification**					**47376532**
					SPO	
Bifidobacterium dentium		94.33	0.44	RAF 95	1	
Clostridium tertium		4.75	0.32	GAL 99	99	
Clostridium clostridioforme		0.70	0.22		66	
47376540	**Low discrimination**					**47376540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		49.43	0.45	GAL 99	1	1	
Blautia producta		43.01	0.24		99	1	
Clostridium butyricum		7.56	0.10		1	99	
47376542	**Low discrimination**						**47376542**
					COC		
Actinomyces israelii		80.90	0.48	GAL 99	1		
Bifidobacterium adolescentis		10.68	0.26		1		
Blautia producta		5.21	0.07		99		
47376712	**Low discrimination**						**47376712**
Clostridium clostridioforme		86.88	0.37	RAF 94			
Bifidobacterium dentium		9.03	0.06	MAN 1			
Actinomyces israelii		2.81	0.21				
47376732	**Low discrimination**						**47376732**
					GRA		
Clostridium clostridioforme		83.49	0.37	RAF 94	50		
Bifidobacterium dentium		15.42	0.10	MAN 1	99		
Bacteroides ovatus		1.09	-.05		1		
47377000	**Low discrimination**						**47377000**
					COC	SPO	
Actinomyces israelii		87.37	0.77		1	1	
Clostridium tertium		8.59	0.45		1	99	
Atopobium parvulum		1.29	0.32		99	1	
47377001	**Low discrimination**						**47377001**
					SPO		
Actinomyces israelii		88.85	0.43	URE 1	1		
Clostridium tertium		8.73	0.11		99		
Actinomyces naeslundii		2.42	0.25		1		
47377002	**Low discrimination**						**47377002**
					SPO		
Actinomyces israelii		82.12	0.80		1		
Clostridium tertium		16.15	0.53		99		
Clostridium innocuum		0.46	0.26		99		
47377003	**Low discrimination**						**47377003**
					SPO		
Actinomyces israelii		82.32	0.46	URE 1	1		
Clostridium tertium		16.18	0.19		99		
Actinomyces naeslundii		1.50	0.25		1		
47377010	**Low discrimination**						**47377010**
					SPO		
Clostridium tertium		53.10	0.50	LAC 99	99		
Actinomyces israelii		45.40	0.63		1		
Lactobacillus jensenii		1.23	0.24		1		
47377012	**Low discrimination**						**47377012**
					SPO		
Clostridium tertium		69.18	0.58	LAC 99	99		
Actinomyces israelii		29.57	0.66		1		
Bifidobacterium adolescentis		0.50	0.29		1		
47377020	**Low discrimination**						**47377020**
					SPO		
Clostridium tertium		80.02	0.53	LAC 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		17.32	0.56		1		
Lactobacillus jensenii		0.82	0.21		1		
47377022	**Low discrimination**						**47377022**
					SPO		
Clostridium tertium		88.66	0.62	LAC 99	99		
Actinomyces israelii		9.60	0.59		1		
Clostridium barati		0.61	0.21		99		
47377030	**Very good identification**						**47377030**
					SPO		
Clostridium tertium		97.97	0.58	LAC 99	99		
Actinomyces israelii		1.78	0.42		1		
Lactobacillus jensenii		0.19	0.13		1		
47377032	**Very good identification**						**47377032**
					SPO		
Clostridium tertium		98.86	0.66	LAC 99	99		
Actinomyces israelii		0.90	0.45		1		
Bifidobacterium dentium		0.20	0.10		1		
47377040	**Very good identification**						**47377040**
					SPO		
Actinomyces israelii		99.52	0.70		1		
Clostridium tertium		0.25	0.11		99		
Clostridium butyricum		0.23	0.04		99		
47377041	**Good identification**						**47377041**
Actinomyces israelii		100.0	0.36	URE 1			
47377042	**Very good identification**						**47377042**
					SPO		
Actinomyces israelii		99.30	0.73		1		
Clostridium tertium		0.51	0.19		99		
Lactobacillus acidophilus		0.07	0.04		1		
47377043	**Good identification**						**47377043**
Actinomyces israelii		100.0	0.39	URE 1			
47377050	**Good identification**						**47377050**
					COC	SPO	
Actinomyces israelii		96.51	0.56	MLZ 13	1	1	
Clostridium tertium		2.93	0.15		1	99	
Blautia producta		0.56	-.03		99	1	
47377052	**Good identification**						**47377052**
					SPO		
Actinomyces israelii		93.80	0.59	MLZ 13	1		
Clostridium tertium		5.70	0.24		99		
Bifidobacterium adolescentis		0.50	0.14		1		
47377060	**Low discrimination**						**47377060**
					SPO		
Actinomyces israelii		89.29	0.49	NAG 6	1		
Clostridium tertium		10.71	0.19		99		
47377062	**Low discrimination**						**47377062**
					SPO		
Actinomyces israelii		79.01	0.52	NAG 6	1		
Clostridium tertium		18.96	0.27		99		
Clostridium barati		1.12	0.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47377100	**Low discrimination**						**47377100**
					COC	SPO	
Clostridium tertium		62.99	0.79		1	99	
Actinomyces israelii		25.89	0.87		1	1	
Atopobium parvulum		9.45	0.66		99	1	
47377101	**Low discrimination**						**47377101**
					COC	SPO	
Clostridium tertium		62.71	0.45	URE 1	1	99	
Actinomyces israelii		25.78	0.53		1	1	
Atopobium parvulum		9.41	0.32		99	1	
47377102	**Low discrimination**						**47377102**
					SPO		
Clostridium tertium		81.28	0.88		99		
Actinomyces israelii		16.71	0.90		1		
Bifidobacterium adolescentis		0.85	0.62		1		
47377103	**Low discrimination**						**47377103**
					SPO		
Clostridium tertium		81.16	0.53	URE 1	99		
Actinomyces israelii		16.68	0.56		1		
Bifidobacterium adolescentis		0.85	0.28		1		
47377110	**Very good identification**						**47377110**
					SPO		
Clostridium tertium		96.39	0.84		99		
Actinomyces israelii		3.33	0.73		1		
Bifidobacterium dentium		0.10	0.23		1		
47377111	**Good identification**						**47377111**
					SPO		
Clostridium tertium		96.66	0.50	URE 1	99		
Actinomyces israelii		3.34	0.39		1		
47377112	**Very good identification**						**47377112**
					SPO		
Clostridium tertium		97.14	0.92		99		
Actinomyces israelii		1.68	0.76		1		
Bifidobacterium adolescentis		0.70	0.63		1		
47377113	**Very good identification**						**47377113**
					SPO		
Clostridium tertium		97.17	0.58	URE 1	99		
Actinomyces israelii		1.68	0.42		1		
Bifidobacterium adolescentis		0.70	0.29		1		
47377120	**Very good identification**						**47377120**
					COC	SPO	
Clostridium tertium		97.72	0.88		1	99	
Atopobium parvulum		0.86	0.53		99	1	
Actinomyces israelii		0.86	0.67		1	1	
47377121	**Good identification**						**47377121**
					COC	SPO	
Clostridium tertium		97.54	0.53	URE 1	1	99	
Atopobium parvulum		0.86	0.19		99	1	
Actinomyces israelii		0.85	0.33		1	1	
47377122	**Very good identification**						**47377122**
					SPO		
Clostridium tertium		98.84	0.96		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		0.56	0.56		92		
Actinomyces israelii		0.43	0.70		1		
47377123	**Very good identification**						**47377123**
					SPO		
Clostridium tertium		98.82	0.62	URE 1	99		
Clostridium ramosum		0.56	0.21		92		
Actinomyces israelii		0.43	0.36		1		
47377130	**Excellent identification**						**47377130**
					SPO		
Clostridium tertium		99.85	0.92		99		
Actinomyces israelii		0.07	0.53		1		
Bifidobacterium dentium		0.06	0.27		1		
47377131	**Very good identification**						**47377131**
					SPO		
Clostridium tertium		99.87	0.58	URE 1	99		
Actinomyces israelii		0.07	0.18		1		
Bifidobacterium dentium		0.06	-.07		1		
47377132	**Excellent identification**						**47377132**
					SPO		
Clostridium tertium		99.74	1.00		99		
Bifidobacterium dentium		0.20	0.44		1		
Actinomyces israelii		0.04	0.56		1		
47377133	**Very good identification**						**47377133**
					SPO		
Clostridium tertium		99.76	0.66	URE 1	99		
Bifidobacterium dentium		0.20	0.10		1		
Actinomyces israelii		0.04	0.21		1		
47377140	**Good identification**						**47377140**
					SPO		
Actinomyces israelii		91.36	0.80		1		
Clostridium tertium		5.77	0.45		99		
Clostridium butyricum		1.27	0.27		99		
47377141	**Acceptable identification**						**47377141**
					SPO		
Actinomyces israelii		92.84	0.46	URE 1	1		
Clostridium tertium		5.87	0.11		99		
Clostridium butyricum		1.29	-.07		99		
47377142	**Low discrimination**						**47377142**
					SPO		
Actinomyces israelii		86.65	0.83		1		
Clostridium tertium		10.95	0.53		99		
Bifidobacterium adolescentis		1.40	0.46		1		
47377143	**Low discrimination**						**47377143**
					SPO		
Actinomyces israelii		87.53	0.49	URE 1	1		
Clostridium tertium		11.06	0.19		99		
Bifidobacterium adolescentis		1.41	0.12		1		
47377150	**Low discrimination**						**47377150**
					COC	SPO	
Actinomyces israelii		52.44	0.66	MLZ 13	1	1	
Clostridium tertium		39.42	0.50		1	99	
Blautia producta		7.57	0.32		99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
47377152	**Low discrimination**						**47377152**
					SPO		
Clostridium tertium		56.08	0.58	SOR 1	99		
Actinomyces israelii		37.30	0.69		1		
Bifidobacterium adolescentis		4.93	0.48		1		
47377160	**Low discrimination**						**47377160**
					COC	SPO	
Clostridium tertium		72.94	0.53	SOR 1	1	99	
Actinomyces israelii		24.56	0.60		1	1	
Atopobium parvulum		0.64	0.19		99	1	
47377162	**Low discrimination**						**47377162**
					SPO		
Clostridium tertium		83.18	0.62	SOR 1	99		
Actinomyces israelii		14.00	0.63		1		
Bifidobacterium breve		1.54	0.41		1		
47377170	**Very good identification**						**47377170**
					COC	SPO	
Clostridium tertium		96.24	0.58	SOR 1	1	99	
Actinomyces israelii		2.72	0.46		1	1	
Blautia producta		0.68	0.15		99	1	
47377172	**Very good identification**						**47377172**
					SPO		
Clostridium tertium		97.36	0.66	SOR 1	99		
Actinomyces israelii		1.38	0.49		1		
Bifidobacterium breve		1.02	0.41		1		
47377200	**Very good identification**						**47377200**
					SPO		
Actinomyces israelii		98.29	0.58	RHA 7	1		
Clostridium tertium		1.30	0.11		99		
Clostridium butyricum		0.42	-.04		99		
47377202	**Very good identification**						**47377202**
					SPO		
Actinomyces israelii		97.03	0.61	RHA 7	1		
Clostridium tertium		2.56	0.19		99		
Lactobacillus acidophilus		0.23	0.00		1		
47377210	**Low discrimination**						**47377210**
					SPO		
Actinomyces israelii		86.43	0.44	RHA 7	1		
Clostridium tertium		13.57	0.15	MLZ 13	99		
47377212	**Low discrimination**						**47377212**
					SPO		
Actinomyces israelii		72.87	0.47	RHA 7	1		
Clostridium tertium		22.87	0.24	MLZ 13	99		
Clostridium clostridioforme		4.26	0.16		66		
47377220	**Low discrimination**						**47377220**
					SPO		
Actinomyces israelii		50.83	0.37	RHA 7	1		
Clostridium tertium		31.52	0.19	NAG 6	99		
Clostridium ramosum		17.65	0.13		92		
47377240	**Good identification**						**47377240**
Actinomyces israelii		100.0	0.51	RHA 7			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47377242	**Good identification**						**47377242**
Actinomyces israelii		100.0	0.54	RHA 7			
47377250	**Acceptable identification**						**47377250**
					COC		
Actinomyces israelii		92.81	0.36	RHA 7	1		
Blautia producta		7.19	-.03	MLZ 13	99		
47377252	**Good identification**						**47377252**
Actinomyces israelii		100.0	0.40	RHA 7			
47377300	**Low discrimination**						**47377300**
					COC	SPO	
Actinomyces israelii		71.10	0.68	RHA 7	1	1	
Clostridium tertium		23.21	0.45		1	99	
Atopobium parvulum		3.48	0.32		99	1	
47377302	**Low discrimination**						**47377302**
					SPO		
Actinomyces israelii		58.95	0.71	RHA 7	1		
Clostridium tertium		38.49	0.53		99		
Clostridium ramosum		0.87	0.24		92		
47377303	**Low discrimination**						**47377303**
					SPO		
Actinomyces israelii		60.50	0.37	RHA 7	1		
Clostridium tertium		39.50	0.19	URE 1	99		
47377310	**Low discrimination**						**47377310**
					SPO		
Clostridium tertium		79.26	0.50	RHA 1	99		
Actinomyces israelii		20.40	0.54		1		
Clostridium butyricum		0.18	-.02		99		
47377312	**Low discrimination**						**47377312**
					SPO		
Clostridium tertium		87.19	0.58	RHA 1	99		
Actinomyces israelii		11.22	0.57		1		
Bifidobacterium adolescentis		0.63	0.29		1		
47377320	**Low discrimination**						**47377320**
					SPO		
Clostridium tertium		61.04	0.53	RHA 1	99		
Clostridium ramosum		34.19	0.47		92		
Actinomyces israelii		3.98	0.48		1		
47377322	**Low discrimination**						**47377322**
					SPO		
Clostridium tertium		55.85	0.62	RHA 1	99		
Clostridium ramosum		41.71	0.58		92		
Actinomyces israelii		1.82	0.51		1		
47377330	**Very good identification**						**47377330**
					SPO		
Clostridium tertium		99.08	0.58	RHA 1	99		
Actinomyces israelii		0.54	0.33		1		
Clostridium ramosum		0.32	0.13		92		
47377332	**Very good identification**						**47377332**
					SPO		
Clostridium tertium		98.79	0.66	RHA 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		0.42	0.24		92		
Actinomyces israelii		0.27	0.36		1		
47377340	**Good identification**						**47377340**
					COC	SPO	
Actinomyces israelii		90.11	0.61	RHA 7	1	1	
Blautia producta		8.61	0.24		99	1	
Clostridium tertium		0.76	0.11		1	99	
47377342	**Very good identification**						**47377342**
					COC	SPO	
Actinomyces israelii		97.11	0.64	RHA 7	1	1	
Clostridium tertium		1.65	0.19		1	99	
Blautia producta		0.69	0.07		99	1	
47377352	**Low discrimination**						**47377352**
					COC	SPO	
Actinomyces israelii		73.44	0.50	RHA 7	1	1	
Clostridium tertium		14.82	0.24	MLZ 13	1	99	
Blautia producta		10.44	0.15		99	1	
47377360	**Low discrimination**						**47377360**
					SPO		
Actinomyces israelii		55.93	0.40	RHA 7	1		
Clostridium tertium		22.29	0.19	NAG 6	99		
Clostridium ramosum		12.49	0.13		92		
47377362	**Low discrimination**						**47377362**
					SPO		
Actinomyces israelii		41.23	0.43	RHA 7	1		
Clostridium tertium		32.86	0.27	NAG 6	99		
Clostridium ramosum		24.54	0.24		92		
47377400	**Low discrimination**						**47377400**
					SPO		
Actinomyces israelii		89.51	0.75		1		
Clostridium butyricum		10.20	0.38		99		
Clostridium innocuum		0.16	0.13		99		
47377401	**Low discrimination**						**47377401**
					SPO		
Actinomyces israelii		89.77	0.41	URE 1	1		
Clostridium butyricum		10.23	0.04		99		
47377402	**Very good identification**						**47377402**
					SPO		
Actinomyces israelii		95.08	0.78		1		
Clostridium butyricum		2.81	0.31		99		
Bifidobacterium adolescentis		1.60	0.42		1		
47377403	**Good identification**						**47377403**
					SPO		
Actinomyces israelii		95.57	0.44	URE 1	1		
Clostridium butyricum		2.82	-.03		99		
Bifidobacterium adolescentis		1.61	0.08		1		
47377410	**Low discrimination**						**47377410**
					SPO		
Actinomyces israelii		83.13	0.61	MLZ 13	1		
Bifidobacterium dentium		11.93	0.23		1		
Clostridium butyricum		3.34	0.16		99		
47377412	**Low discrimination**						**47377412**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		46.40	0.64	MLZ 13			
Bifidobacterium dentium		44.87	0.40				
Bifidobacterium adolescentis		6.37	0.43				
47377420	**Low discrimination**						**47377420**
					SPO		
Actinomyces israelii		82.08	0.55	NAG 6	1		
Clostridium butyricum		12.75	0.20		99		
Clostridium tertium		4.68	0.19		99		
47377422	**Low discrimination**						**47377422**
					SPO		
Actinomyces israelii		83.09	0.58	NAG 6	1		
Clostridium tertium		9.48	0.27		99		
Bifidobacterium dentium		3.38	0.10		1		
47377432	**Acceptable identification**						**47377432**
					SPO		
Bifidobacterium dentium		91.32	0.44	RAF 95	1		
Clostridium tertium		4.60	0.32	LAC 99	99		
Actinomyces israelii		3.39	0.44		1		
47377440	**Good identification**						**47377440**
					COC	SPO	
Actinomyces israelii		94.28	0.68		1	1	
Blautia producta		3.32	0.24		99	1	
Clostridium butyricum		2.40	0.20		1	99	
47377442	**Very good identification**						**47377442**
					SPO		
Actinomyces israelii		98.57	0.71		1		
Clostridium butyricum		0.65	0.13		99		
Bifidobacterium adolescentis		0.53	0.26		1		
47377443	**Good identification**						**47377443**
Actinomyces israelii		100.0	0.37	URE 1			
47377450	**Low discrimination**						**47377450**
					COC	SPO	
Actinomyces israelii		58.32	0.54	MLZ 13	1	1	
Blautia producta		41.16	0.32		99	1	
Clostridium butyricum		0.52	-.02		1	99	
47377452	**Low discrimination**						**47377452**
					COC		
Actinomyces israelii		89.20	0.57	MLZ 13	1		
Blautia producta		4.66	0.15		99		
Bifidobacterium adolescentis		3.89	0.28		1		
47377460	**Acceptable identification**						**47377460**
					COC	SPO	
Actinomyces israelii		91.25	0.48	NAG 6	1	1	
Blautia producta		5.58	0.07		99	1	
Clostridium butyricum		3.17	0.02		1	99	
47377462	**Good identification**						**47377462**
					SPO		
Actinomyces israelii		99.11	0.51	NAG 6	1		
Clostridium butyricum		0.89	-.05		99		
47377472	**Low discrimination**						**47377472**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COC	
Actinomyces israelii		55.85	0.37	MLZ 13	1	
Bifidobacterium dentium		39.07	0.10	NAG 6	1	
Blautia producta		5.08	-.01		99	
47377500	**Low discrimination**					**47377500**
					SPO	
Actinomyces israelii		57.62	0.86		1	
Clostridium butyricum		39.40	0.62		99	
Clostridium tertium		1.73	0.45		99	
47377501	**Low discrimination**					**47377501**
					SPO	
Actinomyces israelii		57.37	0.52	URE 1	1	
Clostridium butyricum		39.24	0.28		99	
Clostridium tertium		1.72	0.11		99	
47377502	**Low discrimination**					**47377502**
					SPO	
Actinomyces israelii		59.16	0.89		1	
Bifidobacterium adolescentis		24.58	0.76		1	
Clostridium butyricum		10.49	0.55		99	
47377503	**Low discrimination**					**47377503**
					SPO	
Actinomyces israelii		59.21	0.55	URE 1	1	
Bifidobacterium adolescentis		24.60	0.42		1	
Clostridium butyricum		10.50	0.21		99	
47377510	**Low discrimination**					**47377510**
					SPO	
Bifidobacterium dentium		67.33	0.57	RAF 95	1	
Actinomyces israelii		18.96	0.72	BGL 91	1	
Clostridium tertium		6.78	0.50		99	
47377512	**Low discrimination**					**47377512**
Bifidobacterium dentium		82.18	0.74	RAF 95		
Bifidobacterium adolescentis		11.67	0.77			
Actinomyces israelii		3.43	0.75			
47377513	**Low discrimination**					**47377513**
Bifidobacterium dentium		82.22	0.40	RAF 95		
Bifidobacterium adolescentis		11.68	0.43	URE 1		
Actinomyces israelii		3.44	0.40			
47377520	**Low discrimination**					**47377520**
					SPO	
Clostridium tertium		40.04	0.53	ARA 1	99	
Actinomyces israelii		28.36	0.65		1	
Clostridium butyricum		26.42	0.44		99	
47377522	**Low discrimination**					**47377522**
					SPO	
Clostridium tertium		54.64	0.62	ARA 1	99	
Bifidobacterium dentium		19.47	0.44		1	
Actinomyces israelii		19.35	0.68		1	
47377530	**Low discrimination**					**47377530**
					SPO	
Bifidobacterium dentium		84.41	0.61	RAF 95	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium tertium		14.33	0.58	BGL 91	99		
Actinomyces israelii		0.85	0.51		1		
47377532	**Good identification**						**47377532**
					SPO		
Bifidobacterium dentium		94.96	0.78	RAF 95	1		
Clostridium tertium		4.78	0.66		99		
Actinomyces israelii		0.14	0.54		1		
47377533	**Acceptable identification**						**47377533**
					SPO		
Bifidobacterium dentium		95.00	0.44	RAF 95	1		
Clostridium tertium		4.79	0.32	URE 1	99		
Actinomyces israelii		0.14	0.20		1		
47377540	**Low discrimination**						**47377540**
					COC	SPO	
Actinomyces israelii		49.23	0.79		1	1	
Blautia producta		42.84	0.58		99	1	
Clostridium butyricum		7.53	0.44		1	99	
47377541	**Low discrimination**						**47377541**
					COC	SPO	
Actinomyces israelii		48.04	0.45	URE 1	1	1	
Blautia producta		41.80	0.24		99	1	
Clostridium butyricum		7.35	0.10		1	99	
47377542	**Low discrimination**						**47377542**
					COC		
Actinomyces israelii		80.71	0.82		1		
Bifidobacterium adolescentis		10.66	0.60		1		
Blautia producta		5.20	0.41		99		
47377543	**Low discrimination**						**47377543**
					COC		
Actinomyces israelii		80.90	0.48	URE 1	1		
Bifidobacterium adolescentis		10.68	0.26		1		
Blautia producta		5.21	0.07		99		
47377550	**Good identification**						**47377550**
					COC		
Blautia producta		93.65	0.66	RAF 99	99		
Actinomyces israelii		5.36	0.65		1		
Bifidobacterium dentium		0.49	0.23		1		
47377552	**Low discrimination**						**47377552**
					COC		
Blautia producta		32.11	0.50	RAF 99	99		
Bifidobacterium adolescentis		26.81	0.62	BGL 10	1		
Actinomyces israelii		24.82	0.68		1		
47377560	**Low discrimination**						**47377560**
					COC	SPO	
Blautia producta		54.40	0.41	RAF 99	99	1	
Actinomyces israelii		35.92	0.58	NAG 10	1	1	
Clostridium butyricum		7.48	0.26		1	99	
47377562	**Low discrimination**						**47377562**
					COC	SPO	
Actinomyces israelii		77.73	0.61	NAG 6	1	1	
Blautia producta		8.72	0.25		99	1	
Clostridium tertium		5.70	0.27		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47377570	**Low discrimination**						**47377570**
					COC		
Blautia producta		88.06	0.50	RAF 99	99		
Bifidobacterium dentium		7.44	0.27	NAG 10	1		
Actinomyces israelii		2.90	0.44		1		
47377572	**Low discrimination**						**47377572**
					COC		
Bifidobacterium dentium		80.05	0.44	RAF 95	1		
Blautia producta		10.41	0.33	SOR 1	99		
Actinomyces israelii		4.62	0.47		1		
47377600	**Very good identification**						**47377600**
					SPO		
Actinomyces israelii		95.53	0.56	RHA 7	1		
Clostridium butyricum		4.47	0.12		99		
47377602	**Very good identification**						**47377602**
					SPO		
Actinomyces israelii		95.14	0.59	RHA 7	1		
Clostridium clostridioforme		3.49	0.25		66		
Clostridium butyricum		1.16	0.05		99		
47377610	**Good identification**						**47377610**
					COC		
Actinomyces israelii		96.45	0.42	RHA 7	1		
Blautia producta		3.55	-.03	MLZ 13	99		
47377612	**Low discrimination**						**47377612**
Clostridium clostridioforme		49.37	0.36	RAF 94			
Actinomyces israelii		44.10	0.45	MAN 1			
Bifidobacterium dentium		5.72	0.06				
47377620	**Acceptable identification**						**47377620**
					SPO		
Actinomyces israelii		94.00	0.36	RHA 7	1		
Clostridium butyricum		6.00	-.06	NAG 6	99		
47377622	**Low discrimination**						**47377622**
					GRA		
Actinomyces israelii		61.26	0.39	RHA 7	99		
Clostridium clostridioforme		35.24	0.25	NAG 6	50		
Bacteroides ovatus		3.50	-.02		1		
47377632	**Low discrimination**						**47377632**
Clostridium clostridioforme		78.27	0.36	RAF 94			
Bifidobacterium dentium		16.13	0.10	MAN 1			
Actinomyces israelii		4.46	0.25				
47377640	**Low discrimination**						**47377640**
					COC	SPO	
Actinomyces israelii		67.68	0.49	RHA 7	1	1	
Blautia producta		31.61	0.24		99	1	
Clostridium butyricum		0.71	-.06		1	99	
47377642	**Good identification**						**47377642**
					COC		
Actinomyces israelii		96.66	0.52	RHA 7	1		
Blautia producta		3.34	0.07		99		
47377652	**Low discrimination**						**47377652**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces israelii		59.01	0.38	RHA 7	1		
Blautia producta		40.99	0.15	MLZ 13	99		
47377700	**Low discrimination**						**47377700**
					GRA	SPO	
Actinomyces israelii		72.34	0.66	RHA 7	99	1	
Clostridium butyricum		20.33	0.36		99	99	
Mitsuokella multacida		3.70	0.32		1	1	
47377702	**Low discrimination**						**47377702**
					SPO		
Actinomyces israelii		84.72	0.69	RHA 7	1		
Clostridium butyricum		6.17	0.29		99		
Bifidobacterium adolescentis		4.73	0.42		1		
47377703	**Low discrimination**						**47377703**
					SPO		
Actinomyces israelii		88.60	0.35	RHA 7	1		
Clostridium butyricum		6.46	-.05	URE 1	99		
Bifidobacterium adolescentis		4.94	0.08		1		
47377710	**Low discrimination**						**47377710**
					COC		
Actinomyces israelii		37.45	0.52	RHA 7	1		
Blautia producta		34.14	0.32	MLZ 13	99		
Bifidobacterium dentium		17.85	0.23		1		
47377712	**Low discrimination**						**47377712**
Bifidobacterium dentium		55.32	0.40	RAF 95			
Actinomyces israelii		17.22	0.55	RHA 1			
Clostridium clostridioforme		16.14	0.45				
47377720	**Low discrimination**						**47377720**
					GRA	SPO	
Actinomyces israelii		50.02	0.46	RHA 7	99	1	
Clostridium butyricum		19.15	0.18	NAG 6	99	99	
Mitsuokella multacida		10.01	0.21		1	1	
47377732	**Low discrimination**						**47377732**
					GRA		
Bifidobacterium dentium		76.90	0.44	RAF 95	99		
Clostridium clostridioforme		12.62	0.45	RHA 1	50		
Bacteroides ovatus		5.42	0.29		1		
47377740	**Good identification**						**47377740**
					COC	GRA	
Blautia producta		90.64	0.58	RAF 99	99	99	
Actinomyces israelii		7.84	0.59		1	99	
Mitsuokella multacida		1.03	0.32		1	1	
47377742	**Low discrimination**						**47377742**
					COC		
Actinomyces israelii		52.63	0.62	RHA 7	1		
Blautia producta		45.07	0.41		99		
Bifidobacterium adolescentis		0.93	0.26		1		
47377750	**Very good identification**						**47377750**
					COC	GRA	
Blautia producta		99.44	0.66	RAF 99	99	99	
Actinomyces israelii		0.43	0.45		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		0.13	0.24		1	1	
47377752	**Acceptable identification**						**47377752**
					COC		
Blautia producta		93.10	0.50	RAF 99	99		
Actinomyces israelii		5.42	0.48	BGL 10	1		
Bifidobacterium adolescentis		0.79	0.28		1		
47377760	**Acceptable identification**						**47377760**
					COC	GRA	
Blautia producta		92.63	0.41	RAF 99	99	99	
Actinomyces israelii		4.60	0.39	NAG 10	1	99	
Mitsuokella multacida		2.37	0.21		1	1	
47377770	**Good identification**						**47377770**
					COC	GRA	
Blautia producta		99.39	0.50	RAF 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
Actinomyces israelii		0.25	0.25		1	99	
47400000	**Low discrimination**						**47400000**
					GRA	SPO	
Fusobacterium mortiferum		70.91	0.52	FRU 99	1	1	
Clostridium botulinum biovar A		15.95	0.27		99	99	
Atopobium minutum		6.37	0.28		99	1	
47400002	**Low discrimination**						**47400002**
					GRA	CAT	
Fusobacterium mortiferum		61.76	0.35	FRU 99	1	5	
Atopobium minutum		33.31	0.25	BGL 10	99	1	
Propionibacterium granulosum		4.94	0.13		99	90	
47400100	**Very good identification**						**47400100**
Fusobacterium mortiferum		100.0	0.58	FRU 99			
47400102	**Good identification**						**47400102**
Fusobacterium mortiferum		100.0	0.42	FRU 99			
47401000	**Low discrimination**						**47401000**
					GRA		
Fusobacterium mortiferum		81.37	0.55	FRU 99	1		
Eubacterium tortuosum		11.71	0.24		99		
Eubacterium saburreum		2.22	0.26		85		
47401002	**Low discrimination**						**47401002**
					GRA		
Fusobacterium mortiferum		69.21	0.38	FRU 99	1		
Eubacterium tortuosum		15.82	0.11	BGL 10	99		
Atopobium minutum		8.29	0.17		99		
47401100	**Very good identification**						**47401100**
Fusobacterium mortiferum		95.55	0.61	FRU 99			
Eubacterium saburreum		2.61	0.33				
Prevotella melaninogenica		1.48	0.12				
47401102	**Low discrimination**						**47401102**
					CAT		
Fusobacterium mortiferum		84.72	0.45	FRU 99	5		
Capnocytophaga ochracea		7.11	0.12	BGL 10	90		
Eubacterium saburreum		3.67	0.20		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47401120	**Good identification**						**47401120**
					CAT		
Prevotella melaninogenica		98.31	0.47	MLT 99	1		
Fusobacterium mortiferum		0.65	0.27	FRU 90	5		
Capnocytophaga ochracea		0.60	0.12		90		
47401520	**Low discrimination**						**47401520**
					GRA		
Bifidobacterium longum subsp. longum		82.47	0.37	MLT 99	99		
Prevotella melaninogenica		14.41	0.12	FRU 90	1		
Odoribacter splanchnicus		1.60	-.07		1		
47401522	**Low discrimination**						**47401522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		81.02	0.37	MLT 99	99	1	
Capnocytophaga ochracea		13.51	0.16	FRU 90	1	90	
Parabacteroides distasonis		2.77	0.16		1	76	
47401530	**Good identification**						**47401530**
Bifidobacterium longum subsp. longum		100.0	0.41	MLT 99			
47401532	**Good identification**						**47401532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.88	0.41	MLT 99	99	1	
Capnocytophaga ochracea		1.07	-.01	FRU 90	1	90	
Parabacteroides distasonis		1.06	0.12		1	76	
47404100	**Good identification**						**47404100**
Fusobacterium mortiferum		100.0	0.39	FRU 99			
47405000	**Low discrimination**						**47405000**
Fusobacterium mortiferum		83.68	0.35	FRU 99			
Eubacterium saburreum		16.32	0.22	XYL 7			
47405100	**Low discrimination**						**47405100**
Fusobacterium mortiferum		83.68	0.42	FRU 99			
Eubacterium saburreum		16.32	0.28	XYL 7			
47405322	**Good identification**						**47405322**
Parabacteroides distasonis		96.81	0.45	FRU 99			
Bacteroides fragilis		3.19	-.03				
47405332	**Good identification**						**47405332**
Parabacteroides distasonis		100.0	0.40	FRU 99			
47405500	**Good identification**						**47405500**
Bifidobacterium longum subsp. longum		100.0	0.35	MLT 99			
47405502	**Good identification**						**47405502**
Bifidobacterium longum subsp. longum		96.73	0.35	MLT 99			
Bifidobacterium adolescentis		3.27	0.12	FRU 90			
47405510	**Good identification**						**47405510**
Bifidobacterium longum subsp. longum		100.0	0.39	MLT 99			
47405512	**Good identification**						**47405512**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. longum		97.63	0.39	MLT 99				
Bifidobacterium adolescentis		2.37	0.13	FRU 90				
47405520	**Low discrimination**							**47405520**
					GRA			
Bifidobacterium longum subsp. longum		89.54	0.46	MLT 99	99			
Bacteroides vulgatus		10.46	0.15	FRU 90	1			
47405522	**Low discrimination**							**47405522**
					GRA	CAT		
Bifidobacterium longum subsp. longum		88.26	0.46	MLT 99	99	1		
Parabacteroides distasonis		9.06	0.33	FRU 90	1	76		
Bacteroides fragilis		1.46	-.03		1	99		
47405530	**Good identification**							**47405530**
Bifidobacterium longum subsp. longum		100.0	0.50	MLT 99				
47405532	**Good identification**							**47405532**
					GRA			
Bifidobacterium longum subsp. longum		96.86	0.50	MLT 99	99			
Parabacteroides distasonis		3.14	0.28	FRU 90	1			
47405722	**Acceptable identification**							**47405722**
					GRA			
Parabacteroides distasonis		93.69	0.41	FRU 99	1			
Bacteroides vulgatus		3.23	-.03		1			
Bifidobacterium longum subsp. longum		3.07	0.11		99			
47405732	**Acceptable identification**							**47405732**
					GRA			
Parabacteroides distasonis		90.60	0.36	FRU 99	1			
Bifidobacterium longum subsp. longum		9.40	0.15		99			
47410000	**Very good identification**							**47410000**
					GRA	CAT		
Fusobacterium mortiferum		96.28	0.86		1	5		
Propionibacterium granulosum		3.42	0.58		99	90		
Eubacterium limosum		0.12	0.18		99	1		
47410001	**Good identification**							**47410001**
					GRA	CAT		
Fusobacterium mortiferum		96.57	0.52	URE 1	1	5		
Propionibacterium granulosum		3.43	0.24		99	90		
47410002	**Good identification**							**47410002**
					GRA	CAT		
Fusobacterium mortiferum		92.17	0.69	BGL 10	1	5		
Propionibacterium granulosum		7.37	0.47		99	90		
Atopobium minutum		0.46	0.25		99	1		
47410003	**Acceptable identification**							**47410003**
					GRA	CAT		
Fusobacterium mortiferum		92.60	0.35	URE 1	1	5		
Propionibacterium granulosum		7.40	0.13	BGL 10	99	90		
47410010	**Low discrimination**							**47410010**
					GRA	SPO	CAT	
Fusobacterium mortiferum		64.78	0.52	MLZ 1	1	1	5	
Propionibacterium granulosum		31.06	0.43		99	1	90	
Clostridium difficile		4.16	0.15		99	99	1	
47410020	**Low discrimination**							**47410020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Fusobacterium mortiferum		52.17	0.52	NAG 1	1	5		
Propionibacterium granulosum		45.85	0.47		99	90		
Atopobium minutum		1.44	0.20		99	1		
47410022	**Low discrimination**							**47410022**
					GRA	CAT		
Propionibacterium granulosum		62.09	0.37	ESL 1	99	90		
Fusobacterium mortiferum		31.40	0.35		1	5		
Atopobium minutum		5.21	0.17		99	1		
47410040	**Good identification**							**47410040**
					GRA	SPO	CAT	
Fusobacterium mortiferum		96.17	0.52	SOR 1	1	1	5	
Propionibacterium granulosum		3.42	0.24		99	1	90	
Clostridium botulinum biovar A		0.41	-.02		99	99	1	
47410042	**Acceptable identification**							**47410042**
					GRA	CAT		
Fusobacterium mortiferum		92.60	0.35	SOR 1	1	5		
Propionibacterium granulosum		7.40	0.13	BGL 10	99	90		
47410100	**Excellent identification**							**47410100**
					GRA	CAT		
Fusobacterium mortiferum		99.98	0.92		1	5		
Propionibacterium granulosum		0.02	0.24		99	90		
Lactobacillus fermentum		0.00	0.15		99	1		
47410101	**Very good identification**							**47410101**
Fusobacterium mortiferum		100.0	0.58	URE 1				
47410102	**Very good identification**							**47410102**
					GRA	CAT		
Fusobacterium mortiferum		99.80	0.76	BGL 10	1	5		
Capnocytophaga ochracea		0.13	0.12		1	90		
Propionibacterium granulosum		0.04	0.13		99	90		
47410103	**Good identification**							**47410103**
Fusobacterium mortiferum		100.0	0.42	URE 1				
47410110	**Very good identification**							**47410110**
					GRA	CAT		
Fusobacterium mortiferum		99.79	0.58	MLZ 1	1	5		
Propionibacterium granulosum		0.21	0.09		99	90		
47410112	**Good identification**							**47410112**
Fusobacterium mortiferum		100.0	0.42	MLZ 1				
47410120	**Low discrimination**							**47410120**
					CAT			
Fusobacterium mortiferum		81.27	0.58	NAG 1	5			
Prevotella melaninogenica		17.02	0.29		1			
Capnocytophaga ochracea		1.14	0.12		90			
47410140	**Very good identification**							**47410140**
Fusobacterium mortiferum		100.0	0.58	SOR 1				
47410142	**Good identification**							**47410142**
Fusobacterium mortiferum		100.0	0.42	SOR 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47410200	**Acceptable identification**							**47410200**
					GRA	CAT		
Fusobacterium mortiferum		90.19	0.52	RHA 1	1	5		
Propionibacterium granulosum		9.81	0.32		99	90		
47410202	**Low discrimination**							**47410202**
					GRA	CAT		
Fusobacterium mortiferum		80.34	0.35	RHA 1	1	5		
Propionibacterium granulosum		19.66	0.22	BGL 10	99	90		
47410300	**Very good identification**							**47410300**
Fusobacterium mortiferum		100.0	0.58	RHA 1				
47410302	**Good identification**							**47410302**
Fusobacterium mortiferum		100.0	0.42	RHA 1				
47410400	**Good identification**							**47410400**
					COC	GRA	CAT	
Fusobacterium mortiferum		96.07	0.52	ARA 1	1	1	5	
Propionibacterium granulosum		3.41	0.24		1	99	90	
Gemella morbillorum		0.51	0.03		99	99	1	
47410402	**Acceptable identification**							**47410402**
					GRA	CAT		
Fusobacterium mortiferum		92.60	0.35	ARA 1	1	5		
Propionibacterium granulosum		7.40	0.13	BGL 10	99	90		
47410500	**Very good identification**							**47410500**
					GRA			
Fusobacterium mortiferum		99.43	0.58	ARA 1	1			
Bifidobacterium longum subsp. longum		0.30	0.09		99			
Lactobacillus fermentum		0.27	0.15		99			
47410502	**Acceptable identification**							**47410502**
					GRA			
Fusobacterium mortiferum		93.16	0.42	ARA 1	1			
Bifidobacterium longum subsp. longum		2.54	0.09	BGL 10	99			
Lactobacillus fermentum		2.23	0.15		99			
47411000	**Very good identification**							**47411000**
					GRA	CAT		
Fusobacterium mortiferum		97.30	0.89		1	5		
Propionibacterium granulosum		2.30	0.58		99	90		
Lactobacillus fermentum		0.25	0.46		99	1		
47411001	**Good identification**							**47411001**
					GRA	CAT		
Fusobacterium mortiferum		96.16	0.55	URE 1	1	5		
Propionibacterium granulosum		2.28	0.24		99	90		
Actinomyces naeslundii		1.57	0.36		99	25		
47411002	**Good identification**							**47411002**
					GRA	CAT		
Fusobacterium mortiferum		92.32	0.72	BGL 10	1	5		
Propionibacterium granulosum		4.92	0.47		99	90		
Lactobacillus fermentum		2.11	0.46		99	1		
47411003	**Low discrimination**							**47411003**
					GRA	CAT		
Fusobacterium mortiferum		83.34	0.38	URE 1	1	5		
Actinomyces naeslundii		12.22	0.36	BGL 10	99	25		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		4.44	0.13		99	90	
47411010	**Low discrimination**						**47411010**
					CAT	GRA	
Fusobacterium mortiferum		72.12	0.55	MLZ 1	5	1	
Propionibacterium granulosum		23.05	0.43		90	99	
Eubacterium saburreum		4.83	0.33		1	85	
47411012	**Low discrimination**						**47411012**
					CAT	GRA	
Fusobacterium mortiferum		54.78	0.38	MLZ 1	5	1	
Propionibacterium granulosum		39.40	0.33	BGL 10	90	99	
Eubacterium saburreum		5.82	0.20		1	85	
47411020	**Low discrimination**						**47411020**
					GRA	CAT	
Fusobacterium mortiferum		46.91	0.55	NAG 1	1	5	
Propionibacterium granulosum		27.49	0.47		99	90	
Prevotella melaninogenica		15.29	0.29		1	1	
47411021	**Low discrimination**						**47411021**
					GRA	CAT	
Actinomyces naeslundii		45.76	0.36	ESL 12	99	25	
Fusobacterium mortiferum		28.37	0.20	MLT 94	1	5	
Propionibacterium granulosum		16.62	0.13		99	90	
47411023	**Low discrimination**						**47411023**
					GRA		
Actinomyces naeslundii		84.06	0.36	ESL 12	99		
Prevotella oralis		15.94	-.03	MLT 94	1		
47411040	**Good identification**						**47411040**
					GRA	CAT	
Fusobacterium mortiferum		97.69	0.55	SOR 1	1	5	
Propionibacterium granulosum		2.31	0.24		99	90	
47411042	**Acceptable identification**						**47411042**
					GRA	CAT	
Fusobacterium mortiferum		94.94	0.38	SOR 1	1	5	
Propionibacterium granulosum		5.06	0.13	BGL 10	99	90	
47411100	**Excellent identification**						**47411100**
					GRA		
Fusobacterium mortiferum		99.45	0.95		1		
Lactobacillus fermentum		0.18	0.50		99		
Prevotella melaninogenica		0.14	0.29		1		
47411101	**Good identification**						**47411101**
					GRA	SPO	
Fusobacterium mortiferum		92.24	0.61	URE 1	1	1	
Clostridium chauvoei		5.63	0.29		99	99	
Actinomyces naeslundii		1.83	0.43		99	1	
47411102	**Low discrimination**						**47411102**
					GRA	CAT	
Fusobacterium mortiferum		88.71	0.79	BGL 10	1	5	
Capnocytophaga ochracea		7.44	0.47		1	90	
Lactobacillus fermentum		1.42	0.50		99	1	
47411103	**Low discrimination**						**47411103**
					GRA	SPO	
Fusobacterium mortiferum		72.13	0.45	URE 1	1	1	
Actinomyces naeslundii		12.90	0.43	BGL 10	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		7.00	0.16		99	99	
47411110	**Good identification**						**47411110**
					CAT		
Fusobacterium mortiferum		92.05	0.61	MLZ 1	5		
Eubacterium saburreum		6.16	0.39		1		
Capnocytophaga ochracea		1.05	0.14		90		
47411112	**Low discrimination**						**47411112**
					CAT		
Fusobacterium mortiferum		46.18	0.45	MLZ 1	5		
Capnocytophaga ochracea		42.62	0.30	BGL 10	90		
Eubacterium saburreum		4.91	0.26		1		
47411120	**Low discrimination**						**47411120**
					CAT		
Prevotella melaninogenica		85.25	0.63	MLT 99	1		
Fusobacterium mortiferum		6.17	0.61		5		
Capnocytophaga ochracea		5.69	0.47		90		
47411122	**Low discrimination**						**47411122**
					CAT		
Capnocytophaga ochracea		57.60	0.63	MLT 99	90		
Prevotella melaninogenica		18.26	0.51		1		
Prevotella oralis		14.24	0.52		1		
47411132	**Low discrimination**						**47411132**
					CAT		
Capnocytophaga ochracea		73.43	0.47	MLT 99	90		
Parabacteroides distasonis		21.02	0.50	MLZ 10	76		
Prevotella melaninogenica		2.12	0.16		1		
47411140	**Very good identification**						**47411140**
					GRA	CAT	
Fusobacterium mortiferum		98.22	0.61	SOR 1	1	5	
Capnocytophaga ochracea		1.12	0.14		1	90	
Actinomyces naeslundii		0.23	0.28		99	25	
47411142	**Low discrimination**						**47411142**
					GRA	CAT	
Fusobacterium mortiferum		50.75	0.45	SOR 1	1	5	
Capnocytophaga ochracea		46.83	0.30	BGL 10	1	90	
Actinomyces naeslundii		1.07	0.28		99	25	
47411162	**Acceptable identification**						**47411162**
					CAT		
Capnocytophaga ochracea		92.02	0.47	MLT 99	90		
Prevotella melaninogenica		2.65	0.16	SOR 10	1		
Prevotella oralis		2.07	0.18		1		
47411200	**Acceptable identification**						**47411200**
					GRA	CAT	
Fusobacterium mortiferum		93.24	0.55	RHA 1	1	5	
Propionibacterium granulosum		6.76	0.32		99	90	
47411202	**Low discrimination**						**47411202**
					GRA	CAT	
Fusobacterium mortiferum		81.62	0.38	RHA 1	1	5	
Propionibacterium granulosum		13.31	0.22	BGL 10	99	90	
Prevotella oralis		5.07	0.04		1	1	
47411300	**Very good identification**						**47411300**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		99.44	0.61	RHA 1	1		
Prevotella oralis		0.24	0.02		1		
Lactobacillus fermentum		0.18	0.15		99		
47411302	**Low discrimination**						**47411302**
Fusobacterium mortiferum		60.20	0.45	RHA 1			
Prevotella oralis		25.10	0.24	BGL 10			
Parabacteroides distasonis		8.06	0.28				
47411322	**Low discrimination**						**47411322**
					CAT		
Parabacteroides distasonis		59.62	0.62	XYL 90	76		
Prevotella oralis		35.65	0.46		1		
Capnocytophaga ochracea		3.40	0.29		90		
47411332	**Very good identification**						**47411332**
					CAT		
Parabacteroides distasonis		97.75	0.58	XYL 90	76		
Capnocytophaga ochracea		1.15	0.12		90		
Prevotella oralis		1.10	0.12		1		
47411400	**Low discrimination**						**47411400**
					GRA	CAT	
Fusobacterium mortiferum		78.16	0.55	ARA 1	1	5	
Lactobacillus fermentum		19.62	0.46		99	1	
Propionibacterium granulosum		1.85	0.24		99	90	
47411402	**Low discrimination**						**47411402**
					GRA	CAT	
Lactobacillus fermentum		66.66	0.46	ESL 1	99	1	
Fusobacterium mortiferum		29.50	0.38		1	5	
Propionibacterium granulosum		1.57	0.13		99	90	
47411420	**Low discrimination**						**47411420**
					GRA		
Lactobacillus fermentum		66.59	0.38	ESL 1	99		
Bifidobacterium longum subsp. longum		14.99	0.20		99		
Fusobacterium mortiferum		8.04	0.20		1		
47411422	**Low discrimination**						**47411422**
					GRA	CAT	
Lactobacillus fermentum		56.84	0.38	ESL 1	99	1	
Capnocytophaga ochracea		23.46	0.16		1	90	
Bifidobacterium longum subsp. longum		12.79	0.20		99	1	
47411500	**Low discrimination**						**47411500**
					GRA		
Fusobacterium mortiferum		71.69	0.61	ARA 1	1		
Bifidobacterium longum subsp. longum		14.32	0.43		99		
Lactobacillus fermentum		12.59	0.50		99		
47411502	**Low discrimination**						**47411502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		29.09	0.43	MLT 99	99	1	
Lactobacillus fermentum		25.56	0.50		99	1	
Capnocytophaga ochracea		23.71	0.34		1	90	
47411510	**Good identification**						**47411510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.08	0.47	MLT 99	99	1	
Fusobacterium mortiferum		2.85	0.27		1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		0.57	0.01		1	90	
47411512	**Low discrimination**						**47411512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		84.54	0.47	MLT 99	99	1	
Bifidobacterium adolescentis		10.12	0.33		99	1	
Capnocytophaga ochracea		4.50	0.17		1	90	
47411520	**Low discrimination**						**47411520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		64.37	0.54	MLT 99	99	1	
Capnocytophaga ochracea		13.12	0.34		1	90	
Prevotella melaninogenica		11.25	0.29		1	1	
47411522	**Low discrimination**						**47411522**
					GRA	CAT	
Capnocytophaga ochracea		54.73	0.50	MLT 99	1	90	
Bifidobacterium longum subsp. longum		29.85	0.54		99	1	
Parabacteroides distasonis		11.23	0.51		1	76	
47411530	**Very good identification**						**47411530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.47	0.58	MLT 99	99	1	
Capnocytophaga ochracea		1.31	0.17		1	90	
Parabacteroides distasonis		0.12	0.12		1	76	
47411532	**Low discrimination**						**47411532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		80.73	0.58	MLT 99	99	1	
Capnocytophaga ochracea		9.66	0.34		1	90	
Parabacteroides distasonis		9.61	0.46		1	76	
47411720	**Good identification**						**47411720**
					GRA		
Bacteroides vulgatus		96.55	0.37	MLT 99	1		
Parabacteroides distasonis		1.55	0.25	XYL 98	1		
Bifidobacterium longum subsp. longum		1.38	0.20		99		
47411722	**Very good identification**						**47411722**
					CAT		
Parabacteroides distasonis		95.82	0.59	XYL 90	76		
Capnocytophaga ochracea		1.57	0.16		90		
Prevotella oralis		0.94	0.12		1		
47411732	**Good identification**						**47411732**
					GRA		
Parabacteroides distasonis		96.40	0.54	XYL 90	1		
Bifidobacterium longum subsp. longum		2.73	0.24		99		
Clostridium clostridioforme		0.55	0.10		50		
47412000	**Low discrimination**						**47412000**
					GRA	CAT	
Propionibacterium granulosum		58.90	0.56	ESL 1	99	90	
Fusobacterium mortiferum		22.21	0.52		1	5	
Eubacterium limosum		16.27	0.32		99	1	
47412002	**Low discrimination**						**47412002**
					GRA	CAT	
Propionibacterium granulosum		84.84	0.45	ESL 1	99	90	
Fusobacterium mortiferum		14.22	0.35		1	5	
Eubacterium limosum		0.95	-.03		99	1	
47412020	**Good identification**						**47412020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Propionibacterium granulosum		97.43	0.45	ESL 1	99	90		
Fusobacterium mortiferum		1.48	0.17		1	5		
Eubacterium limosum		1.09	-.03		99	1		
47412022	**Good identification**							**47412022**
Propionibacterium granulosum		100.0	0.35	ESL 1				
47412100	**Very good identification**							**47412100**
					GRA	CAT		
Fusobacterium mortiferum		98.56	0.58	MAN 1	1	5		
Propionibacterium granulosum		1.13	0.21		99	90		
Eubacterium limosum		0.31	-.03		99	1		
47412102	**Good identification**							**47412102**
					GRA	CAT		
Fusobacterium mortiferum		95.99	0.42	MAN 1	1	5		
Propionibacterium granulosum		2.48	0.11	BGL 10	99	90		
Propionibacterium propionicum		1.53	0.12		99	1		
47413000	**Low discrimination**							**47413000**
					GRA	CAT		
Propionibacterium granulosum		63.28	0.56	ESL 1	99	90		
Fusobacterium mortiferum		35.79	0.55		1	5		
Actinomyces naeslundii		0.76	0.38		99	25		
47413001	**Low discrimination**							**47413001**
					GRA	CAT		
Actinomyces naeslundii		53.08	0.41	ESL 12	99	25		
Propionibacterium granulosum		29.97	0.21	MLT 94	99	90		
Fusobacterium mortiferum		16.95	0.20		1	5		
47413002	**Low discrimination**							**47413002**
					GRA	CAT		
Propionibacterium granulosum		76.98	0.45	ESL 1	99	90		
Fusobacterium mortiferum		19.35	0.38		1	5		
Actinomyces naeslundii		3.67	0.38		99	25		
47413003	**Low discrimination**							**47413003**
Actinomyces naeslundii		87.63	0.41	ESL 12				
Propionibacterium granulosum		12.37	0.11	MLT 94				
47413010	**Acceptable identification**							**47413010**
					GRA	SPO	CAT	
Propionibacterium granulosum		94.52	0.41	ESL 1	99	1	90	
Fusobacterium mortiferum		3.96	0.20	MLZ 12	1	1	5	
Clostridium difficile		1.53	-.03		99	99	1	
47413020	**Acceptable identification**							**47413020**
					GRA	CAT		
Propionibacterium granulosum		92.77	0.45	ESL 1	99	90		
Actinomyces naeslundii		4.43	0.38		99	25		
Fusobacterium mortiferum		2.12	0.20		1	5		
47413021	**Low discrimination**							**47413021**
Actinomyces naeslundii		87.63	0.41	ESL 12				
Propionibacterium granulosum		12.37	0.11	MLT 94				
47413022	**Low discrimination**							**47413022**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Propionibacterium granulosum		82.05	0.35	ESL 1	99	90		
Actinomyces naeslundii		15.66	0.38		99	25		
Prevotella oralis		2.29	-.03		1	1		
47413023	**Good identification**							**47413023**
Actinomyces naeslundii		100.0	0.41	ESL 12				
47413100	**Very good identification**							**47413100**
					GRA	CAT		
Fusobacterium mortiferum		96.47	0.61	MAN 1	1	5		
Actinomyces naeslundii		2.48	0.45		99	25		
Propionibacterium granulosum		0.74	0.21		99	90		
47413101	**Low discrimination**							**47413101**
					GRA			
Actinomyces naeslundii		79.25	0.48	ESL 12	99			
Fusobacterium mortiferum		20.75	0.27	MLT 94	1			
47413102	**Low discrimination**							**47413102**
					GRA	CAT		
Fusobacterium mortiferum		71.10	0.45	MAN 1	1	5		
Actinomyces naeslundii		16.46	0.45	BGL 10	99	25		
Capnocytophaga ochracea		5.96	0.12		1	90		
47413103	**Good identification**							**47413103**
Actinomyces naeslundii		100.0	0.48	ESL 12				
47413121	**Good identification**							**47413121**
					GRA			
Actinomyces naeslundii		96.47	0.48	ESL 12	99			
Prevotella melaninogenica		3.53	-.06	MLT 94	1			
47413123	**Good identification**							**47413123**
Actinomyces naeslundii		100.0	0.48	ESL 12				
47413141	**Good identification**							**47413141**
Actinomyces naeslundii		100.0	0.35	ESL 12				
47413143	**Good identification**							**47413143**
Actinomyces naeslundii		100.0	0.35	ESL 12				
47413161	**Good identification**							**47413161**
Actinomyces naeslundii		100.0	0.35	ESL 12				
47413163	**Good identification**							**47413163**
Actinomyces naeslundii		100.0	0.35	ESL 12				
47414000	**Very good identification**							**47414000**
					COC	GRA	CAT	
Fusobacterium mortiferum		99.46	0.67	XYL 7	1	1	5	
Propionibacterium granulosum		0.47	0.24		1	99	90	
Gemella morbillorum		0.07	0.03		99	99	1	
47414002	**Good identification**							**47414002**
					GRA	CAT		
Fusobacterium mortiferum		98.94	0.50	XYL 7	1	5		
Propionibacterium granulosum		1.06	0.13	BGL 10	99	90		
47414100	**Very good identification**							**47414100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.73	XYL 7		
47414101	**Good identification**					**47414101**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
47414102	**Very good identification**					**47414102**
					CAT	
Fusobacterium mortiferum		98.79	0.57	XYL 7	5	
Bacteroides fragilis		0.79	-.03	BGL 10	99	
Capnocytophaga ochracea		0.42	0.02		90	
47414110	**Good identification**					**47414110**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
47414120	**Low discrimination**					**47414120**
					CAT	
Fusobacterium mortiferum		63.34	0.39	XYL 7	5	
Bacteroides fragilis		28.98	0.10	NAG 1	99	
Bacteroides vulgatus		2.96	-.04		3	
47414140	**Good identification**					**47414140**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
47414300	**Good identification**					**47414300**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
47414322	**Good identification**					**47414322**
Parabacteroides distasonis		96.81	0.45	GAL 99		
Bacteroides fragilis		3.19	-.03			
47414332	**Good identification**					**47414332**
Parabacteroides distasonis		100.0	0.40	GAL 99		
47414500	**Low discrimination**					**47414500**
					GRA	
Fusobacterium mortiferum		89.23	0.39	XYL 7	1	
Bifidobacterium longum subsp. longum		10.77	0.17	ARA 1	99	
47414722	**Acceptable identification**					**47414722**
Parabacteroides distasonis		94.40	0.41	GAL 99		
Bacteroides vulgatus		3.26	-.03			
Clostridium clostridioforme		2.34	0.08			
47414732	**Low discrimination**					**47414732**
Parabacteroides distasonis		82.64	0.36	GAL 99		
Clostridium clostridioforme		17.36	0.19			
47415000	**Very good identification**					**47415000**
					GRA	
Fusobacterium mortiferum		97.19	0.70	XYL 7	1	
Lactobacillus fermentum		1.92	0.42		99	
Eubacterium saburreum		0.58	0.30		85	
47415001	**Good identification**					**47415001**
Fusobacterium mortiferum		100.0	0.35	XYL 7		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47415002	**Low discrimination**						**47415002**
					GRA	CAT	
Fusobacterium mortiferum		82.40	0.53	XYL 7	1	5	
Lactobacillus fermentum		14.67	0.42	BGL 10	99	1	
Bacteroides fragilis		1.02	-.03		1	99	
47415010	**Low discrimination**						**47415010**
					CAT	GRA	
Fusobacterium mortiferum		65.72	0.35	XYL 7	5	1	
Eubacterium saburreum		31.46	0.28	MLZ 1	1	85	
Propionibacterium granulosum		2.82	0.09		90	99	
47415020	**Low discrimination**						**47415020**
					GRA	CAT	
Fusobacterium mortiferum		36.56	0.35	XYL 7	1	5	
Bacteroides fragilis		26.02	0.10	NAG 1	1	99	
Lactobacillus fermentum		23.87	0.34		99	1	
47415040	**Good identification**						**47415040**
Fusobacterium mortiferum		100.0	0.35	XYL 7			
47415100	**Very good identification**						**47415100**
					GRA		
Fusobacterium mortiferum		97.20	0.76	XYL 7	1		
Lactobacillus fermentum		1.35	0.46		99		
Eubacterium saburreum		0.58	0.36		85		
47415101	**Good identification**						**47415101**
Fusobacterium mortiferum		100.0	0.42	XYL 7			
47415102	**Low discrimination**						**47415102**
					CAT		
Fusobacterium mortiferum		46.03	0.60	XYL 7	5		
Bacteroides fragilis		24.25	0.32	BGL 10	99		
Capnocytophaga ochracea		12.83	0.36		90		
47415110	**Low discrimination**						**47415110**
					GRA		
Fusobacterium mortiferum		60.47	0.42	XYL 7	1		
Eubacterium saburreum		28.96	0.35	MLZ 1	85		
Bifidobacterium longum subsp. longum		8.28	0.21		99		
47415120	**Low discrimination**						**47415120**
					CAT		
Bacteroides fragilis		73.47	0.44	MLT 99	99		
Bacteroides vulgatus		7.50	0.30	BGL 95	3		
Capnocytophaga ochracea		7.46	0.36		90		
47415122	**Low discrimination**						**47415122**
Bacteroides fragilis		86.03	0.66	MLT 99			
Parabacteroides distasonis		8.80	0.71				
Capnocytophaga ochracea		4.14	0.53				
47415132	**Low discrimination**						**47415132**
Parabacteroides distasonis		76.95	0.66				
Bacteroides fragilis		14.12	0.32				
Capnocytophaga ochracea		7.47	0.36				
47415140	**Good identification**						**47415140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Fusobacterium mortiferum		96.35	0.42	XYL 7	5	
Capnocytophaga ochracea		3.65	0.04	SOR 1	90	
47415200	**Good identification**					**47415200**
Fusobacterium mortiferum		100.0	0.35	XYL 7		
47415222	**Acceptable identification**					**47415222**
					CAT	
Parabacteroides distasonis		92.18	0.45	LAC 99	76	
Prevotella oralis		4.78	0.10		1	
Bacteroides fragilis		3.04	-.03		99	
47415232	**Good identification**					**47415232**
Parabacteroides distasonis		100.0	0.40	LAC 99		
47415300	**Low discrimination**					**47415300**
					GRA	
Fusobacterium mortiferum		55.31	0.42	XYL 7	1	
Eubacterium contortum		28.20	0.24	RHA 1	99	
Bacteroides vulgatus		15.50	0.13		1	
47415302	**Low discrimination**					**47415302**
					GRA	
Parabacteroides distasonis		82.90	0.45	NAG 99	1	
Fusobacterium mortiferum		5.18	0.25		1	
Eubacterium contortum		4.20	0.11		99	
47415312	**Good identification**					**47415312**
Parabacteroides distasonis		97.59	0.40	NAG 99		
Clostridium clostridioforme		2.41	0.07			
47415320	**Acceptable identification**					**47415320**
					CAT	
Bacteroides vulgatus		92.35	0.47	MLT 99	3	
Parabacteroides distasonis		5.92	0.45	ARA 93	76	
Bacteroides fragilis		1.02	0.10		99	
47415322	**Very good identification**					**47415322**
					CAT	
Parabacteroides distasonis		95.74	0.79		76	
Bacteroides fragilis		3.15	0.32		99	
Prevotella oralis		0.79	0.30		1	
47415323	**Good identification**					**47415323**
Parabacteroides distasonis		96.81	0.45	URE 1		
Bacteroides fragilis		3.19	-.03			
47415330	**Low discrimination**					**47415330**
Parabacteroides distasonis		77.36	0.40	BGL 99		
Bacteroides vulgatus		22.64	0.13			
47415332	**Very good identification**					**47415332**
Parabacteroides distasonis		99.88	0.74			
Bacteroides fragilis		0.06	-.03			
Capnocytophaga ochracea		0.03	0.02			
47415333	**Good identification**					**47415333**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.40	URE 1			
47415362	**Good identification**						**47415362**
Parabacteroides distasonis		95.21	0.45	SOR 1			
Bacteroides fragilis		3.14	-.03				
Capnocytophaga ochracea		1.66	0.02				
47415372	**Good identification**						**47415372**
Parabacteroides distasonis		100.0	0.40	SOR 1			
47415400	**Low discrimination**						**47415400**
					GRA		
Lactobacillus fermentum		62.25	0.42	ESL 1	99		
Fusobacterium mortiferum		31.78	0.35		1		
Bifidobacterium longum subsp. longum		5.97	0.17		99		
47415402	**Low discrimination**						**47415402**
					GRA		
Lactobacillus fermentum		84.16	0.42	ESL 1	99		
Bifidobacterium longum subsp. longum		8.06	0.17		99		
Fusobacterium mortiferum		4.77	0.19		1		
47415500	**Low discrimination**						**47415500**
					GRA		
Bifidobacterium longum subsp. longum		73.49	0.52	MLT 99	99		
Lactobacillus fermentum		12.64	0.46		99		
Fusobacterium mortiferum		9.23	0.42		1		
47415502	**Low discrimination**						**47415502**
Bifidobacterium longum subsp. longum		59.56	0.52	MLT 99			
Bifidobacterium adolescentis		22.16	0.46				
Lactobacillus fermentum		10.25	0.46				
47415510	**Good identification**						**47415510**
Bifidobacterium longum subsp. longum		99.73	0.56	MLT 99			
Bifidobacterium adolescentis		0.27	0.13				
47415512	**Low discrimination**						**47415512**
					GRA		
Bifidobacterium longum subsp. longum		77.54	0.56	MLT 99	99		
Bifidobacterium adolescentis		20.71	0.47		99		
Parabacteroides distasonis		1.12	0.28		1		
47415520	**Low discrimination**						**47415520**
					GRA	CAT	
Bacteroides vulgatus		54.97	0.50	MLT 99	1	3	
Bifidobacterium longum subsp. longum		42.77	0.62	RHA 90	99	1	
Capnocytophaga ochracea		0.73	0.23		1	90	
47415522	**Low discrimination**						**47415522**
					GRA	CAT	
Parabacteroides distasonis		45.22	0.67		1	76	
Bifidobacterium longum subsp. longum		40.06	0.62		99	1	
Bacteroides fragilis		7.29	0.32		1	99	
47415530	**Very good identification**						**47415530**
					GRA		
Bifidobacterium longum subsp. longum		98.78	0.66	MLT 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		0.75	0.15		1		
Parabacteroides distasonis		0.36	0.28		1		
47415532	**Low discrimination**						**47415532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		73.00	0.66	MLT 99	99	1	
Parabacteroides distasonis		26.06	0.62		1	76	
Capnocytophaga ochracea		0.73	0.23		1	90	
47415622	**Acceptable identification**						**47415622**
Parabacteroides distasonis		94.63	0.41	LAC 99			
Bacteroides vulgatus		3.27	-.03				
Clostridium clostridioforme		2.11	0.07				
47415632	**Low discrimination**						**47415632**
Parabacteroides distasonis		84.16	0.36	LAC 99			
Clostridium clostridioforme		15.84	0.18				
47415702	**Low discrimination**						**47415702**
					GRA		
Parabacteroides distasonis		78.26	0.41	NAG 99	1		
Clostridium clostridioforme		5.83	0.16		50		
Bifidobacterium longum subsp. longum		5.78	0.17		99		
47415712	**Low discrimination**						**47415712**
					GRA		
Parabacteroides distasonis		51.92	0.36	NAG 99	1		
Clostridium clostridioforme		32.72	0.27		50		
Bifidobacterium longum subsp. longum		12.12	0.21		99		
47415720	**Very good identification**						**47415720**
					GRA		
Bacteroides vulgatus		99.60	0.66	MLT 99	1		
Parabacteroides distasonis		0.29	0.41		1		
Bifidobacterium longum subsp. longum		0.09	0.28		99		
47415722	**Very good identification**						**47415722**
					CAT		
Parabacteroides distasonis		95.66	0.75		76		
Bacteroides vulgatus		3.30	0.32		3		
Bacteroides thetaiotaomicron		0.54	0.21		89		
47415723	**Good identification**						**47415723**
Parabacteroides distasonis		96.66	0.41	URE 1			
Bacteroides vulgatus		3.34	-.03				
47415732	**Very good identification**						**47415732**
					GRA		
Parabacteroides distasonis		97.99	0.71		1		
Bifidobacterium longum subsp. longum		0.92	0.32		99		
Clostridium clostridioforme		0.62	0.27		50		
47415733	**Good identification**						**47415733**
Parabacteroides distasonis		100.0	0.36	URE 1			
47415762	**Good identification**						**47415762**
Parabacteroides distasonis		96.66	0.41	SOR 1			
Bacteroides vulgatus		3.34	-.03				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47415772	**Good identification**							**47415772**
Parabacteroides distasonis		100.0	0.36	SOR 1				
47416100	**Good identification**							**47416100**
Fusobacterium mortiferum		100.0	0.39	MAN 1				
47417000	**Low discrimination**							**47417000**
					GRA	CAT		
Fusobacterium mortiferum		80.82	0.35	MAN 1	1	5		
Propionibacterium granulosum		19.18	0.21	XYL 7	99	90		
47417100	**Good identification**							**47417100**
					GRA			
Fusobacterium mortiferum		97.87	0.42	MAN 1	1			
Actinomyces israelii		2.13	0.16	XYL 7	99			
47417322	**Good identification**							**47417322**
Parabacteroides distasonis		96.81	0.45	MAN 1				
Bacteroides fragilis		3.19	-.03					
47417332	**Good identification**							**47417332**
Parabacteroides distasonis		100.0	0.40	MAN 1				
47417502	**Good identification**							**47417502**
Bifidobacterium adolescentis		95.77	0.44	MLT 99				
Bifidobacterium longum subsp. longum		3.18	0.17					
Actinomyces israelii		1.05	0.17					
47417512	**Good identification**							**47417512**
Bifidobacterium adolescentis		95.58	0.46	MLT 99				
Bifidobacterium longum subsp. longum		4.42	0.21					
47417722	**Good identification**							**47417722**
Parabacteroides distasonis		96.66	0.41	MAN 1				
Bacteroides vulgatus		3.34	-.03					
47417732	**Good identification**							**47417732**
Parabacteroides distasonis		100.0	0.36	MAN 1				
47420000	**Low discrimination**							**47420000**
					GRA	SPO	CAT	
Fusobacterium mortiferum		64.73	0.43	FRU 99	1	1	5	
Propionibacterium granulosum		23.09	0.32		99	1	90	
Clostridium botulinum biovar A		12.18	0.17		99	99	1	
47420100	**Good identification**							**47420100**
Fusobacterium mortiferum		100.0	0.49	FRU 99				
47421000	**Low discrimination**							**47421000**
					CAT	GRA		
Fusobacterium mortiferum		77.00	0.46	FRU 99	5	1		
Propionibacterium granulosum		18.31	0.32		90	99		
Eubacterium saburreum		4.69	0.23		1	85		
47421100	**Acceptable identification**							**47421100**
					GRA			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		93.86	0.52	FRU 99	1	
Eubacterium saburreum		5.71	0.29		85	
Actinomyces naeslundii		0.43	0.23		99	
47421102	**Low discrimination**					**47421102**
					GRA	
Fusobacterium mortiferum		87.92	0.36	FRU 99	1	
Eubacterium saburreum		8.50	0.17	BGL 10	85	
Actinomyces naeslundii		3.59	0.23		99	
47421322	**Good identification**					**47421322**
Parabacteroides distasonis		100.0	0.40	FRU 99		
47421722	**Good identification**					**47421722**
Parabacteroides distasonis		100.0	0.36	FRU 99		
47425122	**Good identification**					**47425122**
Parabacteroides distasonis		97.88	0.48	FRU 99		
Bacteroides fragilis		2.12	-.03			
47425132	**Good identification**					**47425132**
Parabacteroides distasonis		100.0	0.43	FRU 99		
47425322	**Good identification**					**47425322**
Parabacteroides distasonis		100.0	0.56	FRU 99		
47425332	**Good identification**					**47425332**
Parabacteroides distasonis		100.0	0.51	FRU 99		
47425520	**Good identification**					**47425520**
					GRA	
Bifidobacterium longum subsp. longum		96.31	0.35	MLT 99	99	
Bacteroides vulgatus		1.88	-.08	FRU 90	1	
Parabacteroides distasonis		1.82	0.10		1	
47425522	**Low discrimination**					**47425522**
					GRA	
Parabacteroides distasonis		65.16	0.44	FRU 99	1	
Bifidobacterium longum subsp. longum		34.85	0.35		99	
47425530	**Good identification**					**47425530**
Bifidobacterium longum subsp. longum		100.0	0.39	MLT 99		
47425532	**Low discrimination**					**47425532**
					GRA	
Bifidobacterium longum subsp. longum		62.84	0.39	MLT 99	99	
Parabacteroides distasonis		37.16	0.39	FRU 90	1	
47425722	**Good identification**					**47425722**
Parabacteroides distasonis		98.39	0.52	FRU 99		
Bacteroides thetaiotaomicron		1.61	0.06			
47425732	**Good identification**					**47425732**
Parabacteroides distasonis		97.84	0.48	FRU 99		
Bacteroides thetaiotaomicron		1.16	-.01			
Clostridium clostridioforme		1.00	0.08			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47430000	**Low discrimination**							**47430000**
					GRA	CAT		
Fusobacterium mortiferum		73.71	0.77		1	5		
Propionibacterium granulosum		26.29	0.66		99	90		
47430001	**Low discrimination**							**47430001**
					GRA	CAT		
Fusobacterium mortiferum		73.71	0.43	URE 1	1	5		
Propionibacterium granulosum		26.29	0.32		99	90		
47430002	**Low discrimination**							**47430002**
					GRA	CAT		
Fusobacterium mortiferum		55.48	0.60	BGL 10	1	5		
Propionibacterium granulosum		44.52	0.56		99	90		
47430010	**Low discrimination**							**47430010**
					GRA	SPO	CAT	
Propionibacterium granulosum		82.57	0.51	ESL 1	99	1	90	
Fusobacterium mortiferum		17.15	0.43	MLZ 12	1	1	5	
Clostridium difficile		0.28	-.04		99	99	1	
47430012	**Acceptable identification**							**47430012**
					GRA	CAT		
Propionibacterium granulosum		91.55	0.41	ESL 1	99	90		
Fusobacterium mortiferum		8.45	0.26	MLZ 12	1	5		
47430020	**Low discrimination**							**47430020**
					GRA	CAT		
Propionibacterium granulosum		89.83	0.56	ESL 1	99	90		
Fusobacterium mortiferum		10.18	0.43		1	5		
47430022	**Good identification**							**47430022**
					GRA	CAT		
Propionibacterium granulosum		95.21	0.45	ESL 1	99	90		
Fusobacterium mortiferum		4.79	0.26		1	5		
47430030	**Good identification**							**47430030**
Propionibacterium granulosum		100.0	0.41	ESL 1				
47430040	**Low discrimination**							**47430040**
					GRA	SPO	CAT	
Fusobacterium mortiferum		72.65	0.43	SOR 1	1	1	5	
Propionibacterium granulosum		25.91	0.32		99	1	90	
Clostridium botulinum biovar B		1.44	-.03		99	99	1	
47430100	**Very good identification**							**47430100**
					GRA	CAT		
Fusobacterium mortiferum		99.85	0.83		1	5		
Propionibacterium granulosum		0.15	0.32		99	90		
47430101	**Good identification**							**47430101**
					GRA			
Fusobacterium mortiferum		99.20	0.49	URE 1	1			
Actinomyces naeslundii		0.80	0.25		99			
47430102	**Very good identification**							**47430102**
					GRA	CAT		
Fusobacterium mortiferum		99.65	0.67	BGL 10	1	5		
Propionibacterium granulosum		0.35	0.21		99	90		
47430110	**Good identification**							**47430110**
					GRA	CAT		
Fusobacterium mortiferum		97.96	0.49	MLZ 1	1	5		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		2.04	0.17		99	90	
47430120	**Acceptable identification**						**47430120**
					GRA	CAT	
Fusobacterium mortiferum		93.89	0.49	NAG 1	1	5	
Propionibacterium granulosum		3.59	0.21		99	90	
Clostridium perfringens		1.86	0.09		99	1	
47430140	**Good identification**						**47430140**
Fusobacterium mortiferum		100.0	0.49	SOR 1			
47430200	**Low discrimination**						**47430200**
					GRA	CAT	
Propionibacterium granulosum		52.20	0.40	ESL 1	99	90	
Fusobacterium mortiferum		47.80	0.43	RHA 3	1	5	
47430300	**Good identification**						**47430300**
Fusobacterium mortiferum		100.0	0.49	RHA 1			
47430322	**Good identification**						**47430322**
Parabacteroides distasonis		100.0	0.40	GAL 99			
47430400	**Low discrimination**						**47430400**
					GRA	CAT	
Fusobacterium mortiferum		73.71	0.43	ARA 1	1	5	
Propionibacterium granulosum		26.29	0.32		99	90	
47430500	**Good identification**						**47430500**
Fusobacterium mortiferum		100.0	0.49	ARA 1			
47430722	**Good identification**						**47430722**
Parabacteroides distasonis		96.56	0.36	GAL 99			
Clostridium clostridioforme		3.44	0.05	XYL 90			
47431000	**Low discrimination**						**47431000**
					CAT	GRA	
Fusobacterium mortiferum		80.62	0.80		5	1	
Propionibacterium granulosum		19.17	0.66		90	99	
Eubacterium saburreum		0.15	0.31		1	85	
47431001	**Low discrimination**						**47431001**
					GRA	CAT	
Fusobacterium mortiferum		74.42	0.46	URE 1	1	5	
Propionibacterium granulosum		17.70	0.32		99	90	
Actinomyces naeslundii		7.88	0.41		99	25	
47431002	**Low discrimination**						**47431002**
					GRA	CAT	
Fusobacterium mortiferum		64.75	0.64	BGL 10	1	5	
Propionibacterium granulosum		34.64	0.56		99	90	
Actinomyces naeslundii		0.42	0.38		99	25	
47431010	**Low discrimination**						**47431010**
					CAT	GRA	
Propionibacterium granulosum		73.64	0.51	ESL 1	90	99	
Fusobacterium mortiferum		22.94	0.46	MLZ 12	5	1	
Eubacterium saburreum		3.43	0.30		1	85	
47431012	**Low discrimination**						**47431012**
					CAT	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		85.38	0.41	ESL 1	90	99	
Fusobacterium mortiferum		11.82	0.29	MLZ 12	5	1	
Eubacterium saburreum		2.80	0.17		1	85	
47431020	**Low discrimination**						**47431020**
					GRA	CAT	
Propionibacterium granulosum		79.37	0.56	ESL 1	99	90	
Fusobacterium mortiferum		13.49	0.46		1	5	
Clostridium perfringens		5.61	0.28		99	1	
47431021	**Low discrimination**						**47431021**
					GRA	CAT	
Actinomyces naeslundii		60.36	0.41	ESL 12	99	25	
Propionibacterium granulosum		33.89	0.21	MLT 94	99	90	
Fusobacterium mortiferum		5.76	0.12		1	5	
47431022	**Low discrimination**						**47431022**
					GRA	CAT	
Propionibacterium granulosum		70.38	0.45	ESL 1	99	90	
Clostridium perfringens		9.37	0.23		99	1	
Parabacteroides distasonis		8.44	0.31		1	76	
47431023	**Low discrimination**						**47431023**
Actinomyces naeslundii		87.69	0.41	ESL 12			
Propionibacterium granulosum		12.31	0.11	MLT 94			
47431030	**Good identification**						**47431030**
					CAT	GRA	
Propionibacterium granulosum		95.65	0.41	ESL 1	90	99	
Eubacterium saburreum		3.14	0.17	MLZ 12	1	85	
Fusobacterium mortiferum		1.20	0.12		5	1	
47431040	**Low discrimination**						**47431040**
					GRA	CAT	
Fusobacterium mortiferum		79.99	0.46	SOR 1	1	5	
Propionibacterium granulosum		19.02	0.32		99	90	
Actinomyces naeslundii		1.00	0.25		99	25	
47431100	**Very good identification**						**47431100**
					CAT	GRA	
Fusobacterium mortiferum		99.52	0.86		5	1	
Eubacterium saburreum		0.18	0.38		1	85	
Propionibacterium granulosum		0.10	0.32		90	99	
47431101	**Low discrimination**						**47431101**
					GRA	SPO	
Fusobacterium mortiferum		86.82	0.52	URE 1	1	1	
Actinomyces naeslundii		11.21	0.48		99	1	
Clostridium chauvoei		1.97	0.12		99	99	
47431102	**Very good identification**						**47431102**
					GRA		
Fusobacterium mortiferum		97.15	0.70	BGL 10	1		
Actinomyces naeslundii		0.76	0.45		99		
Parabacteroides distasonis		0.67	0.31		1		
47431103	**Low discrimination**						**47431103**
					GRA	SPO	
Actinomyces naeslundii		52.87	0.48	ESL 12	99	1	
Fusobacterium mortiferum		45.49	0.36	MLT 94	1	1	
Clostridium chauvoei		1.64	-.01		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47431110	**Low discrimination**						**47431110**
					CAT	GRA	
Fusobacterium mortiferum		85.63	0.52	MLZ 1	5	1	
Eubacterium saburreum		12.79	0.36		1	85	
Propionibacterium granulosum		1.19	0.17		90	99	
47431112	**Low discrimination**						**47431112**
Fusobacterium mortiferum		55.30	0.36	MLZ 1			
Parabacteroides distasonis		20.26	0.27	BGL 10			
Eubacterium saburreum		13.12	0.23				
47431120	**Low discrimination**						**47431120**
					GRA		
Clostridium perfringens		42.67	0.43	MLT 99	99		
Fusobacterium mortiferum		32.62	0.52		1		
Prevotella melaninogenica		15.25	0.29		1		
47431121	**Low discrimination**						**47431121**
					GRA		
Actinomyces naeslundii		78.39	0.48	ESL 12	99		
Clostridium perfringens		8.02	0.09	MLT 94	99		
Fusobacterium mortiferum		6.13	0.18		1		
47431122	**Low discrimination**						**47431122**
					GRA		
Parabacteroides distasonis		81.82	0.66	XYL 90	1		
Clostridium perfringens		6.73	0.38		99		
Prevotella oralis		4.80	0.32		1		
47431123	**Low discrimination**						**47431123**
					GRA		
Actinomyces naeslundii		59.97	0.48	ESL 12	99		
Parabacteroides distasonis		35.09	0.31	MLT 94	1		
Clostridium perfringens		2.89	0.03		99		
47431132	**Very good identification**						**47431132**
					GRA	CAT	
Parabacteroides distasonis		98.47	0.61	XYL 90	1	76	
Capnocytophaga ochracea		0.76	0.12		1	90	
Bifidobacterium longum subsp. longum		0.28	0.13		99	1	
47431140	**Good identification**						**47431140**
					GRA		
Fusobacterium mortiferum		97.76	0.52	SOR 1	1		
Actinomyces naeslundii		1.49	0.32		99		
Clostridium perfringens		0.75	0.05		99		
47431141	**Low discrimination**						**47431141**
					GRA		
Actinomyces naeslundii		69.29	0.35	ESL 12	99		
Fusobacterium mortiferum		30.71	0.18	MLT 94	1		
47431142	**Low discrimination**						**47431142**
					GRA		
Fusobacterium mortiferum		87.97	0.36	SOR 1	1		
Actinomyces naeslundii		12.03	0.32	BGL 10	99		
47431143	**Good identification**						**47431143**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47431161	**Good identification**						**47431161**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.35	ESL 12			
47431163	**Good identification**						**47431163**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47431200	**Low discrimination**						**47431200**
					GRA	CAT	
Fusobacterium mortiferum		57.87	0.46	RHA 1	1	5	
Propionibacterium granulosum		42.13	0.40		99	90	
47431222	**Low discrimination**						**47431222**
					GRA	CAT	
Parabacteroides distasonis		87.80	0.40	XYL 90	1	76	
Propionibacterium granulosum		7.55	0.19	LAC 99	99	90	
Prevotella oralis		4.65	0.05		1	1	
47431300	**Good identification**						**47431300**
Fusobacterium mortiferum		100.0	0.52	RHA 1			
47431302	**Low discrimination**						**47431302**
Parabacteroides distasonis		65.20	0.40	XYL 90			
Fusobacterium mortiferum		31.95	0.36	NAG 99			
Prevotella oralis		2.85	0.04				
47431320	**Low discrimination**						**47431320**
					GRA		
Parabacteroides distasonis		83.79	0.40	XYL 90	1		
Clostridium perfringens		4.88	0.09	BGL 99	99		
Fusobacterium mortiferum		3.73	0.18		1		
47431322	**Very good identification**						**47431322**
					GRA		
Parabacteroides distasonis		99.14	0.74	XYL 90	1		
Prevotella oralis		0.83	0.25		1		
Clostridium perfringens		0.03	0.03		99		
47431323	**Good identification**						**47431323**
Parabacteroides distasonis		100.0	0.40	XYL 90			
47431332	**Very good identification**						**47431332**
Parabacteroides distasonis		99.96	0.69	XYL 90			
Clostridium clostridioforme		0.04	0.04				
47431362	**Good identification**						**47431362**
Parabacteroides distasonis		100.0	0.40	XYL 90			
47431400	**Low discrimination**						**47431400**
					GRA	CAT	
Fusobacterium mortiferum		80.79	0.46	ARA 1	1	5	
Propionibacterium granulosum		19.21	0.32		99	90	
47431500	**Low discrimination**						**47431500**
					GRA		
Fusobacterium mortiferum		85.24	0.52	ARA 1	1		
Bifidobacterium longum subsp. longum		14.25	0.33		99		
Lactobacillus fermentum		0.51	0.15		99		
47431510	**Good identification**						**47431510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium longum subsp. longum		96.57	0.37	MLT 99	99		
Fusobacterium mortiferum		3.43	0.18		1		
47431512	**Low discrimination**						**47431512**
					GRA		
Bifidobacterium longum subsp. longum		69.48	0.37	MLT 99	99		
Bifidobacterium adolescentis		22.19	0.30		99		
Parabacteroides distasonis		6.09	0.23		1		
47431520	**Low discrimination**						**47431520**
					GRA		
Bifidobacterium longum subsp. longum		88.30	0.43	MLT 99	99		
Parabacteroides distasonis		6.12	0.28		1		
Clostridium perfringens		1.74	0.09		99		
47431522	**Low discrimination**						**47431522**
					GRA	CAT	
Parabacteroides distasonis		86.18	0.62	XYL 90	1	76	
Bifidobacterium longum subsp. longum		12.57	0.43		99	1	
Capnocytophaga ochracea		0.93	0.16		1	90	
47431530	**Good identification**						**47431530**
					GRA		
Bifidobacterium longum subsp. longum		97.86	0.47	MLT 99	99		
Parabacteroides distasonis		2.14	0.23		1		
47431532	**Low discrimination**						**47431532**
					GRA		
Parabacteroides distasonis		68.16	0.57	XYL 90	1		
Bifidobacterium longum subsp. longum		31.44	0.47		99		
Clostridium clostridioforme		0.25	0.10		50		
47431622	**Good identification**						**47431622**
Parabacteroides distasonis		96.91	0.36	XYL 90			
Clostridium clostridioforme		3.09	0.05	LAC 99			
47431702	**Acceptable identification**						**47431702**
Parabacteroides distasonis		90.36	0.36	XYL 90			
Clostridium clostridioforme		9.64	0.14	NAG 99			
47431720	**Low discrimination**						**47431720**
					GRA		
Parabacteroides distasonis		61.85	0.36	XYL 90	1		
Bacteroides vulgatus		35.14	0.13	BGL 99	1		
Bifidobacterium longum subsp. longum		3.01	0.09		99		
47431722	**Very good identification**						**47431722**
Parabacteroides distasonis		99.23	0.70	XYL 90			
Bacteroides thetaiotaomicron		0.61	0.16				
Clostridium clostridioforme		0.11	0.14				
47431723	**Good identification**						**47431723**
Parabacteroides distasonis		100.0	0.36	XYL 90			
47431732	**Very good identification**						**47431732**
Parabacteroides distasonis		98.51	0.65	XYL 90			
Clostridium clostridioforme		0.90	0.25				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides thetaiotaomicron		0.44	0.09					
47431762	**Good identification**							**47431762**
Parabacteroides distasonis		100.0	0.36	XYL 90				
47432000	**Very good identification**							**47432000**
					GRA	SPO	CAT	
Propionibacterium granulosum		96.19	0.64	ESL 1	99	1	90	
Fusobacterium mortiferum		3.61	0.43		1	1	5	
Clostridium difficile		0.11	0.01		99	99	1	
47432002	**Good identification**							**47432002**
					GRA	CAT		
Propionibacterium granulosum		98.36	0.53	ESL 1	99	90		
Fusobacterium mortiferum		1.64	0.26		1	5		
47432010	**Good identification**							**47432010**
					SPO	CAT		
Propionibacterium granulosum		96.15	0.49	ESL 1	1	90		
Clostridium difficile		3.86	0.13	MLZ 12	99	1		
47432012	**Good identification**							**47432012**
Propionibacterium granulosum		100.0	0.39	ESL 1				
47432020	**Good identification**							**47432020**
Propionibacterium granulosum		100.0	0.53	ESL 1				
47432022	**Good identification**							**47432022**
Propionibacterium granulosum		100.0	0.43	ESL 1				
47432030	**Good identification**							**47432030**
Propionibacterium granulosum		100.0	0.39	ESL 1				
47432100	**Low discrimination**							**47432100**
					GRA	CAT		
Fusobacterium mortiferum		88.83	0.49	MAN 1	1	5		
Propionibacterium granulosum		10.24	0.30		99	90		
Actinomyces naeslundii		0.93	0.27		99	25		
47432200	**Good identification**							**47432200**
Propionibacterium granulosum		100.0	0.38	ESL 1				
47433000	**Good identification**							**47433000**
					GRA	CAT		
Propionibacterium granulosum		93.98	0.64	ESL 1	99	90		
Fusobacterium mortiferum		5.29	0.46		1	5		
Actinomyces naeslundii		0.73	0.42		99	25		
47433001	**Low discrimination**							**47433001**
					GRA	CAT		
Actinomyces naeslundii		52.03	0.46	ESL 12	99	25		
Propionibacterium granulosum		45.41	0.30	MLT 94	99	90		
Fusobacterium mortiferum		2.56	0.12		1	5		
47433002	**Acceptable identification**							**47433002**
					GRA	CAT		
Propionibacterium granulosum		94.71	0.53	ESL 1	99	90		
Actinomyces naeslundii		2.92	0.42		99	25		
Fusobacterium mortiferum		2.37	0.29		1	5		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47433003	**Low discrimination**						**47433003**
Actinomyces naeslundii		82.09	0.46	ESL 12			
Propionibacterium granulosum		17.91	0.19	MLT 94			
47433010	**Good identification**						**47433010**
					GRA	CAT	
Propionibacterium granulosum		99.59	0.49	ESL 1	99	90	
Fusobacterium mortiferum		0.42	0.12	MLZ 12	1	5	
47433012	**Good identification**						**47433012**
Propionibacterium granulosum		100.0	0.39	ESL 1			
47433020	**Good identification**						**47433020**
					GRA	CAT	
Propionibacterium granulosum		96.79	0.53	ESL 1	99	90	
Actinomyces naeslundii		2.99	0.42		99	25	
Fusobacterium mortiferum		0.22	0.12		1	5	
47433021	**Low discrimination**						**47433021**
Actinomyces naeslundii		82.09	0.46	ESL 12			
Propionibacterium granulosum		17.91	0.19	MLT 94			
47433022	**Low discrimination**						**47433022**
Propionibacterium granulosum		89.01	0.43	ESL 1			
Actinomyces naeslundii		10.99	0.42				
47433023	**Acceptable identification**						**47433023**
Actinomyces naeslundii		94.83	0.46	ESL 12			
Propionibacterium granulosum		5.17	0.09	MLT 94			
47433030	**Good identification**						**47433030**
Propionibacterium granulosum		100.0	0.39	ESL 1			
47433100	**Low discrimination**						**47433100**
					GRA	CAT	
Fusobacterium mortiferum		80.39	0.52	MAN 1	1	5	
Actinomyces naeslundii		13.44	0.50		99	25	
Propionibacterium granulosum		6.18	0.30		99	90	
47433101	**Good identification**						**47433101**
					GRA		
Actinomyces naeslundii		96.13	0.53	ESL 12	99		
Fusobacterium mortiferum		3.87	0.18	MLT 94	1		
47433102	**Low discrimination**						**47433102**
					GRA	CAT	
Actinomyces naeslundii		54.52	0.50	ESL 12	99	25	
Fusobacterium mortiferum		36.24	0.36	MLT 94	1	5	
Propionibacterium granulosum		6.27	0.19		99	90	
47433103	**Good identification**						**47433103**
Actinomyces naeslundii		100.0	0.53	ESL 12			
47433120	**Low discrimination**						**47433120**
					CAT		
Actinomyces naeslundii		77.96	0.50	ESL 12	25		
Propionibacterium granulosum		8.97	0.19	MLT 94	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		6.16	0.09		1		
47433121	**Good identification**						**47433121**
Actinomyces naeslundii		100.0	0.53	ESL 12			
47433122	**Low discrimination**						**47433122**
					GRA		
Actinomyces naeslundii		64.74	0.50	ESL 12	99		
Parabacteroides distasonis		29.27	0.31	MLT 94	1		
Clostridium perfringens		2.41	0.03		99		
47433123	**Good identification**						**47433123**
Actinomyces naeslundii		100.0	0.53	ESL 12			
47433140	**Low discrimination**						**47433140**
					GRA		
Actinomyces naeslundii		74.49	0.37	ESL 12	99		
Fusobacterium mortiferum		25.51	0.18	MLT 94	1		
47433141	**Good identification**						**47433141**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47433142	**Good identification**						**47433142**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47433143	**Good identification**						**47433143**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47433160	**Good identification**						**47433160**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47433161	**Good identification**						**47433161**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47433162	**Good identification**						**47433162**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47433163	**Good identification**						**47433163**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47433200	**Good identification**						**47433200**
					GRA	CAT	
Propionibacterium granulosum		98.19	0.38	ESL 1	99	90	
Fusobacterium mortiferum		1.81	0.12	RHA 3	1	5	
47433322	**Good identification**						**47433322**
					SPO		
Parabacteroides distasonis		98.19	0.40	MAN 1	1		
Clostridium ramosum		1.81	-.03	XYL 90	92		
47433722	**Good identification**						**47433722**
Parabacteroides distasonis		100.0	0.36	MAN 1			
47434000	**Very good identification**						**47434000**
					GRA	CAT	
Fusobacterium mortiferum		95.43	0.58	XYL 7	1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		4.57	0.32		99	90	
47434002	**Acceptable identification**						**47434002**
					GRA	CAT	
Fusobacterium mortiferum		90.28	0.41	XYL 7	1	5	
Propionibacterium granulosum		9.72	0.21	BGL 10	99	90	
47434100	**Very good identification**						**47434100**
Fusobacterium mortiferum		100.0	0.64	XYL 7			
47434102	**Good identification**						**47434102**
Fusobacterium mortiferum		98.77	0.48	XYL 7			
Parabacteroides distasonis		1.23	0.14	BGL 10			
47434122	**Good identification**						**47434122**
					CAT		
Parabacteroides distasonis		97.10	0.48	GAL 99	76		
Bacteroides fragilis		2.11	-.03		99		
Fusobacterium mortiferum		0.80	0.13		5		
47434132	**Good identification**						**47434132**
Parabacteroides distasonis		100.0	0.43	GAL 99			
47434322	**Good identification**						**47434322**
Parabacteroides distasonis		100.0	0.56	GAL 99			
47434332	**Good identification**						**47434332**
Parabacteroides distasonis		98.69	0.51	GAL 99			
Clostridium clostridioforme		1.31	0.13				
47434522	**Acceptable identification**						**47434522**
					GRA		
Parabacteroides distasonis		93.90	0.44	GAL 99	1		
Bifidobacterium longum subsp. longum		4.57	0.17		99		
Clostridium clostridioforme		1.54	0.08		50		
47434532	**Low discrimination**						**47434532**
					GRA		
Parabacteroides distasonis		77.39	0.39	GAL 99	1		
Bifidobacterium longum subsp. longum		11.90	0.21		99		
Clostridium clostridioforme		10.71	0.19		50		
47434722	**Acceptable identification**						**47434722**
Parabacteroides distasonis		94.66	0.52	GAL 99			
Clostridium clostridioforme		3.78	0.23				
Bacteroides thetaiotaomicron		1.55	0.06				
47434732	**Low discrimination**						**47434732**
Parabacteroides distasonis		74.07	0.48	GAL 99			
Clostridium clostridioforme		25.05	0.34				
Bacteroides thetaiotaomicron		0.88	-.01				
47435000	**Very good identification**						**47435000**
					CAT	GRA	
Fusobacterium mortiferum		95.68	0.61	XYL 7	5	1	
Propionibacterium granulosum		3.05	0.32		90	99	
Eubacterium saburreum		1.26	0.27		1	85	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47435002	**Low discrimination**						**47435002**
					CAT	GRA	
Fusobacterium mortiferum		89.92	0.44	XYL 7	5	1	
Propionibacterium granulosum		6.46	0.21	BGL 10	90	99	
Eubacterium saburreum		1.88	0.14		1	85	
47435022	**Good identification**						**47435022**
					GRA		
Parabacteroides distasonis		96.99	0.48	LAC 99	1		
Bacteroides fragilis		2.10	-.03		1		
Propionibacterium granulosum		0.91	0.11		99		
47435032	**Good identification**						**47435032**
Parabacteroides distasonis		100.0	0.43	LAC 99			
47435100	**Very good identification**						**47435100**
					COC	GRA	
Fusobacterium mortiferum		97.27	0.67	XYL 7	1	1	
Atopobium parvulum		1.29	0.23		99	99	
Eubacterium saburreum		1.28	0.33		1	85	
47435102	**Low discrimination**						**47435102**
					GRA		
Fusobacterium mortiferum		50.17	0.51	XYL 7	1		
Parabacteroides distasonis		41.22	0.48	BGL 10	1		
Bifidobacterium adolescentis		4.51	0.29		99		
47435112	**Low discrimination**						**47435112**
					GRA		
Parabacteroides distasonis		74.10	0.43	NAG 99	1		
Bifidobacterium adolescentis		18.39	0.30		99		
Eubacterium saburreum		2.87	0.19		85		
47435120	**Low discrimination**						**47435120**
					CAT		
Parabacteroides distasonis		71.95	0.48	BGL 99	76		
Bacteroides fragilis		8.13	0.10		99		
Fusobacterium mortiferum		7.96	0.33		5		
47435122	**Very good identification**						**47435122**
Parabacteroides distasonis		97.64	0.82				
Bacteroides fragilis		2.12	0.32				
Capnocytophaga ochracea		0.10	0.18				
47435123	**Good identification**						**47435123**
Parabacteroides distasonis		97.88	0.48	URE 1			
Bacteroides fragilis		2.12	-.03				
47435130	**Low discrimination**						**47435130**
					GRA		
Parabacteroides distasonis		88.26	0.43	BGL 99	1		
Bifidobacterium longum subsp. longum		8.32	0.21		99		
Eubacterium saburreum		3.42	0.19		85		
47435132	**Very good identification**						**47435132**
					GRA	CAT	
Parabacteroides distasonis		99.83	0.77		1	76	
Bifidobacterium longum subsp. longum		0.10	0.21		99	1	
Bacteroides fragilis		0.04	-.03		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47435133	**Good identification**					**47435133**
Parabacteroides distasonis		100.0	0.43	URE 1		
47435162	**Good identification**					**47435162**
Parabacteroides distasonis		96.79	0.48	SOR 1		
Bacteroides fragilis		2.10	-.03			
Capnocytophaga ochracea		1.11	0.02			
47435172	**Good identification**					**47435172**
Parabacteroides distasonis		100.0	0.43	SOR 1		
47435222	**Good identification**					**47435222**
Parabacteroides distasonis		100.0	0.56	LAC 99		
47435232	**Good identification**					**47435232**
Parabacteroides distasonis		98.83	0.51	LAC 99		
Clostridium clostridioforme		1.17	0.13			
47435302	**Good identification**					**47435302**
Parabacteroides distasonis		99.13	0.56	NAG 99		
Clostridium clostridioforme		0.47	0.10			
Fusobacterium mortiferum		0.41	0.16			
47435312	**Good identification**					**47435312**
Parabacteroides distasonis		96.18	0.51	NAG 99		
Clostridium clostridioforme		3.82	0.22			
47435320	**Low discrimination**					**47435320**
Parabacteroides distasonis		87.51	0.56	BGL 99		
Bacteroides vulgatus		12.49	0.23			
47435322	**Very good identification**					**47435322**
					CAT	
Parabacteroides distasonis		99.97	0.90		76	
Prevotella oralis		0.01	0.10		1	
Bacteroides thetaiotaomicron		0.01	0.06		89	
47435323	**Good identification**					**47435323**
Parabacteroides distasonis		100.0	0.56	URE 1		
47435330	**Good identification**					**47435330**
Parabacteroides distasonis		100.0	0.51	BGL 99		
47435332	**Very good identification**					**47435332**
Parabacteroides distasonis		99.95	0.85			
Clostridium clostridioforme		0.04	0.22			
Bacteroides thetaiotaomicron		0.01	-.01			
47435333	**Good identification**					**47435333**
Parabacteroides distasonis		100.0	0.51	URE 1		
47435362	**Good identification**					**47435362**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.56	SOR 1			
47435372	**Good identification**						**47435372**
Parabacteroides distasonis		100.0	0.51	SOR 1			
47435422	**Acceptable identification**						**47435422**
					GRA		
Parabacteroides distasonis		94.05	0.44	LAC 99	1		
Bifidobacterium longum subsp. longum		4.57	0.17		99		
Clostridium clostridioforme		1.38	0.07		50		
47435432	**Low discrimination**						**47435432**
					GRA		
Parabacteroides distasonis		78.27	0.39	LAC 99	1		
Bifidobacterium longum subsp. longum		12.03	0.21		99		
Clostridium clostridioforme		9.70	0.18		50		
47435500	**Low discrimination**						**47435500**
					GRA	SPO	
Bifidobacterium longum subsp. longum		83.80	0.41	MLT 99	99	1	
Fusobacterium mortiferum		12.57	0.33		1	1	
Clostridium butyricum		1.36	-.03		99	99	
47435502	**Low discrimination**						**47435502**
					GRA		
Bifidobacterium longum subsp. longum		34.61	0.41	MLT 99	99		
Bifidobacterium adolescentis		34.34	0.43		99		
Parabacteroides distasonis		28.76	0.44		1		
47435510	**Good identification**						**47435510**
Bifidobacterium longum subsp. longum		99.29	0.45	MLT 99			
Bifidobacterium adolescentis		0.71	0.10				
47435512	**Low discrimination**						**47435512**
					GRA		
Bifidobacterium longum subsp. longum		47.98	0.45	MLT 99	99		
Bifidobacterium adolescentis		34.18	0.44		99		
Parabacteroides distasonis		12.61	0.39		1		
47435520	**Low discrimination**						**47435520**
					GRA		
Bifidobacterium longum subsp. longum		70.32	0.52	MLT 99	99		
Bacteroides vulgatus		15.07	0.26		1		
Parabacteroides distasonis		14.61	0.44		1		
47435522	**Good identification**						**47435522**
					GRA	CAT	
Parabacteroides distasonis		94.90	0.78		1	76	
Bifidobacterium longum subsp. longum		4.61	0.52		99	1	
Bacteroides thetaiotaomicron		0.35	0.21		1	89	
47435523	**Good identification**						**47435523**
					GRA		
Parabacteroides distasonis		95.36	0.44	URE 1	1		
Bifidobacterium longum subsp. longum		4.64	0.17		99		
47435530	**Acceptable identification**						**47435530**
					GRA		
Bifidobacterium longum subsp. longum		93.72	0.56	MLT 99	99		
Parabacteroides distasonis		6.16	0.39		1		
Bacteroides vulgatus		0.12	-.08		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47435532	**Low discrimination**					**47435532**
					GRA	
Parabacteroides distasonis		86.14	0.74		1	
Bifidobacterium longum subsp. longum		13.24	0.56		99	
Clostridium clostridioforme		0.36	0.27		50	
47435533	**Low discrimination**					**47435533**
					GRA	
Parabacteroides distasonis		86.67	0.39	URE 1	1	
Bifidobacterium longum subsp. longum		13.33	0.21		99	
47435562	**Good identification**					**47435562**
					GRA	
Parabacteroides distasonis		95.36	0.44	SOR 1	1	
Bifidobacterium longum subsp. longum		4.64	0.17		99	
47435572	**Low discrimination**					**47435572**
					GRA	
Parabacteroides distasonis		86.67	0.39	SOR 1	1	
Bifidobacterium longum subsp. longum		13.33	0.21		99	
47435622	**Acceptable identification**					**47435622**
Parabacteroides distasonis		93.55	0.52	LAC 99		
Clostridium clostridioforme		3.35	0.22			
Bacteroides thetaiotaomicron		3.10	0.11			
47435632	**Low discrimination**					**47435632**
Parabacteroides distasonis		75.36	0.48	LAC 99		
Clostridium clostridioforme		22.84	0.33			
Bacteroides thetaiotaomicron		1.80	0.04			
47435702	**Low discrimination**					**47435702**
Parabacteroides distasonis		87.38	0.52	NAG 99		
Clostridium clostridioforme		10.48	0.31			
Bacteroides thetaiotaomicron		1.43	0.06			
47435712	**Low discrimination**					**47435712**
					GRA	
Clostridium clostridioforme		49.53	0.42	MLT 99	50	
Parabacteroides distasonis		48.82	0.48	SAL 88	1	
Bifidobacterium longum subsp. longum		0.63	0.11		99	
47435720	**Low discrimination**					**47435720**
					CAT	
Bacteroides vulgatus		74.97	0.42	MLT 99	3	
Parabacteroides distasonis		24.24	0.52	TRE 4	76	
Bacteroides thetaiotaomicron		0.40	0.06		89	
47435722	**Very good identification**					**47435722**
Parabacteroides distasonis		98.22	0.86			
Bacteroides thetaiotaomicron		1.61	0.40			
Clostridium clostridioforme		0.12	0.31			
47435723	**Good identification**					**47435723**
Parabacteroides distasonis		98.39	0.52	URE 1		
Bacteroides thetaiotaomicron		1.61	0.06			
47435730	**Low discrimination**					**47435730**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Parabacteroides distasonis		88.39	0.48	BGL 99	1	
Bacteroides vulgatus		5.13	0.08		1	
Bifidobacterium longum subsp. longum		4.53	0.21		99	
47435732	**Very good identification**					**47435732**
Parabacteroides distasonis		97.79	0.82			
Bacteroides thetaiotaomicron		1.16	0.33			
Clostridium clostridioforme		1.00	0.42			
47435733	**Good identification**					**47435733**
Parabacteroides distasonis		97.84	0.48	URE 1		
Bacteroides thetaiotaomicron		1.16	-.01			
Clostridium clostridioforme		1.00	0.08			
47435762	**Good identification**					**47435762**
Parabacteroides distasonis		98.39	0.52	SOR 1		
Bacteroides thetaiotaomicron		1.61	0.06			
47435772	**Good identification**					**47435772**
Parabacteroides distasonis		97.84	0.48	SOR 1		
Bacteroides thetaiotaomicron		1.16	-.01			
Clostridium clostridioforme		1.00	0.08			
47437122	**Good identification**					**47437122**
Parabacteroides distasonis		97.88	0.48	MAN 1		
Bacteroides fragilis		2.12	-.03			
47437132	**Good identification**					**47437132**
Parabacteroides distasonis		100.0	0.43	MAN 1		
47437322	**Good identification**					**47437322**
Parabacteroides distasonis		100.0	0.56	MAN 1		
47437332	**Good identification**					**47437332**
Parabacteroides distasonis		100.0	0.51	MAN 1		
47437502	**Acceptable identification**					**47437502**
Bifidobacterium adolescentis		92.18	0.41	MLT 99		
Actinomyces israelii		5.72	0.27			
Bifidobacterium longum subsp. longum		1.15	0.07			
47437512	**Good identification**					**47437512**
Bifidobacterium adolescentis		96.93	0.43	MLT 99		
Bifidobacterium longum subsp. longum		1.68	0.11			
Bifidobacterium dentium		1.39	-.07			
47437522	**Acceptable identification**					**47437522**
					GRA	
Parabacteroides distasonis		94.47	0.44	MAN 1	1	
Bifidobacterium longum subsp. longum		4.59	0.17		99	
Bifidobacterium adolescentis		0.93	0.07		99	
47437532	**Low discrimination**					**47437532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		81.14	0.39	MAN 1	1		
Bifidobacterium longum subsp. longum		12.47	0.21		99		
Bifidobacterium dentium		4.57	-.03		99		
47437722	**Good identification**						**47437722**
Parabacteroides distasonis		98.39	0.52	MAN 1			
Bacteroides thetaiotaomicron		1.61	0.06				
47437732	**Good identification**						**47437732**
Parabacteroides distasonis		97.84	0.48	MAN 1			
Bacteroides thetaiotaomicron		1.16	-.01				
Clostridium clostridioforme		1.00	0.08				
47440000	**Good identification**						**47440000**
Fusobacterium mortiferum		100.0	0.43	FRU 99			
47440100	**Good identification**						**47440100**
Fusobacterium mortiferum		100.0	0.50	FRU 99			
47441000	**Good identification**						**47441000**
					GRA		
Fusobacterium mortiferum		95.42	0.46	FRU 99	1		
Eubacterium tortuosum		4.58	0.07		99		
47441100	**Good identification**						**47441100**
Fusobacterium mortiferum		99.49	0.53	FRU 99			
Prevotella melaninogenica		0.52	-.04				
47441102	**Low discrimination**						**47441102**
					CAT		
Fusobacterium mortiferum		85.66	0.36	FRU 99	5		
Prevotella oralis		8.95	0.06	BGL 10	1		
Capnocytophaga ochracea		5.39	0.02		90		
47445322	**Acceptable identification**						**47445322**
					CAT		
Parabacteroides distasonis		92.57	0.40	FRU 99	76		
Prevotella oralis		5.01	0.06		1		
Bacteroides fragilis		2.43	-.09		99		
47445332	**Good identification**						**47445332**
Parabacteroides distasonis		100.0	0.35	FRU 99			
47445722	**Acceptable identification**						**47445722**
Parabacteroides distasonis		90.71	0.36	FRU 99			
Bacteroides uniformis		9.29	0.01				
47450000	**Very good identification**						**47450000**
					GRA	CAT	
Fusobacterium mortiferum		99.89	0.78		1	5	
Propionibacterium granulosum		0.11	0.24		99	90	
47450001	**Good identification**						**47450001**
Fusobacterium mortiferum		100.0	0.43	URE 1			
47450002	**Very good identification**						**47450002**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Fusobacterium mortiferum		99.76	0.61	BGL 10	1	5	
Propionibacterium granulosum		0.24	0.13		99	90	
47450010	**Good identification**						**47450010**
					GRA	CAT	
Fusobacterium mortiferum		98.57	0.43	MLZ 1	1	5	
Propionibacterium granulosum		1.43	0.09		99	90	
47450020	**Good identification**						**47450020**
					GRA	CAT	
Fusobacterium mortiferum		97.41	0.43	NAG 1	1	5	
Propionibacterium granulosum		2.59	0.13		99	90	
47450040	**Good identification**						**47450040**
Fusobacterium mortiferum		100.0	0.43	SOR 1			
47450100	**Very good identification**						**47450100**
Fusobacterium mortiferum		100.0	0.84				
47450101	**Good identification**						**47450101**
Fusobacterium mortiferum		100.0	0.50	URE 1			
47450102	**Very good identification**						**47450102**
					CAT		
Fusobacterium mortiferum		99.75	0.68	BGL 10	5		
Prevotella oralis		0.16	0.06		1		
Capnocytophaga ochracea		0.10	0.02		90		
47450110	**Good identification**						**47450110**
Fusobacterium mortiferum		100.0	0.50	MLZ 1			
47450120	**Acceptable identification**						**47450120**
Fusobacterium mortiferum		91.10	0.50	NAG 1			
Prevotella melaninogenica		6.36	0.12				
Prevotella oralis		1.59	0.06				
47450140	**Good identification**						**47450140**
Fusobacterium mortiferum		100.0	0.50	SOR 1			
47450200	**Good identification**						**47450200**
Fusobacterium mortiferum		100.0	0.43	RHA 1			
47450300	**Good identification**						**47450300**
Fusobacterium mortiferum		100.0	0.50	RHA 1			
47450400	**Good identification**						**47450400**
Fusobacterium mortiferum		100.0	0.43	ARA 1			
47450500	**Good identification**						**47450500**
Fusobacterium mortiferum		100.0	0.50	ARA 1			
47451000	**Very good identification**						**47451000**
					GRA	CAT	
Fusobacterium mortiferum		99.79	0.81		1	5	
Propionibacterium granulosum		0.07	0.24		99	90	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		0.05	0.14		99	1	
47451001	**Acceptable identification**						**47451001**
					GRA		
Fusobacterium mortiferum		93.92	0.46	URE 1	1		
Actinomyces naeslundii		6.09	0.38		99		
47451002	**Very good identification**						**47451002**
					GRA		
Fusobacterium mortiferum		96.89	0.64	BGL 10	1		
Prevotella oralis		1.51	0.19		1		
Actinomyces naeslundii		0.38	0.35		99		
47451010	**Good identification**						**47451010**
					GRA	CAT	
Fusobacterium mortiferum		97.50	0.46	MLZ 1	1	5	
Lactobacillus jensenii		1.56	0.06		99	1	
Propionibacterium granulosum		0.94	0.09		99	90	
47451020	**Low discrimination**						**47451020**
Fusobacterium mortiferum		70.87	0.46	NAG 1			
Prevotella oralis		12.13	0.19				
Prevotella melaninogenica		7.70	0.12				
47451021	**Low discrimination**						**47451021**
					GRA		
Actinomyces naeslundii		86.51	0.38	ESL 12	99		
Fusobacterium mortiferum		13.49	0.12	MLT 94	1		
47451022	**Low discrimination**						**47451022**
					CAT		
Prevotella oralis		87.95	0.41	MLT 99	1		
Capnocytophaga ochracea		3.97	0.18	LAC 94	90		
Fusobacterium mortiferum		3.00	0.30		5		
47451023	**Low discrimination**						**47451023**
					GRA		
Actinomyces naeslundii		66.35	0.38	ESL 12	99		
Prevotella oralis		33.65	0.07	MLT 94	1		
47451040	**Good identification**						**47451040**
Fusobacterium mortiferum		100.0	0.46	SOR 1			
47451100	**Very good identification**						**47451100**
					CAT		
Fusobacterium mortiferum		99.70	0.87		5		
Capnocytophaga ochracea		0.08	0.20		90		
Prevotella oralis		0.06	0.18		1		
47451101	**Acceptable identification**						**47451101**
					GRA		
Fusobacterium mortiferum		92.68	0.53	URE 1	1		
Actinomyces naeslundii		7.32	0.46		99		
47451102	**Low discrimination**						**47451102**
					CAT		
Fusobacterium mortiferum		83.88	0.71	BGL 10	5		
Prevotella oralis		8.77	0.40		1		
Capnocytophaga ochracea		5.28	0.36		90		
47451103	**Low discrimination**						**47451103**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		53.23	0.36	URE 1	1		
Actinomyces naeslundii		37.86	0.46	BGL 10	99		
Prevotella oralis		5.56	0.06		1		
47451110	**Good identification**						**47451110**
					GRA	CAT	
Fusobacterium mortiferum		98.68	0.53	MLZ 1	1	5	
Capnocytophaga ochracea		0.84	0.04		1	90	
Bifidobacterium breve		0.48	0.10		99	1	
47451112	**Low discrimination**						**47451112**
					GRA	CAT	
Fusobacterium mortiferum		37.52	0.36	MLZ 1	1	5	
Capnocytophaga ochracea		25.96	0.20	BGL 10	1	90	
Bifidobacterium breve		16.64	0.27		99	1	
47451120	**Low discrimination**						**47451120**
Prevotella melaninogenica		60.00	0.47	MLT 99			
Prevotella oralis		14.97	0.40	CEL 10			
Fusobacterium mortiferum		13.02	0.53				
47451121	**Low discrimination**						**47451121**
					GRA		
Actinomyces naeslundii		51.23	0.46	ESL 12	99		
Prevotella melaninogenica		30.17	0.12	MLT 94	1		
Prevotella oralis		7.53	0.06		1		
47451122	**Low discrimination**						**47451122**
					CAT		
Prevotella oralis		71.03	0.62	MLT 99	1		
Capnocytophaga ochracea		20.26	0.53		90		
Prevotella melaninogenica		2.86	0.34		1		
47451132	**Low discrimination**						**47451132**
					CAT		
Capnocytophaga ochracea		48.01	0.36	MLT 99	90		
Parabacteroides distasonis		29.60	0.45	MLZ 10	76		
Prevotella oralis		15.30	0.27		1		
47451140	**Good identification**						**47451140**
					GRA	CAT	
Fusobacterium mortiferum		97.78	0.53	SOR 1	1	5	
Actinomyces naeslundii		0.91	0.30		99	25	
Capnocytophaga ochracea		0.84	0.04		1	90	
47451142	**Low discrimination**						**47451142**
					GRA	CAT	
Fusobacterium mortiferum		41.39	0.36	SOR 1	1	5	
Capnocytophaga ochracea		28.65	0.20	BGL 10	1	90	
Bifidobacterium breve		18.36	0.27		99	1	
47451162	**Low discrimination**						**47451162**
					GRA	CAT	
Capnocytophaga ochracea		66.13	0.36	MLT 99	1	90	
Prevotella oralis		21.08	0.27	SOR 10	1	1	
Bifidobacterium breve		8.37	0.32		99	1	
47451200	**Good identification**						**47451200**
Fusobacterium mortiferum		100.0	0.46	RHA 1			
47451300	**Good identification**						**47451300**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		97.47	0.53	RHA 1			
Prevotella oralis		2.53	0.12				
47451322	**Low discrimination**						**47451322**
					CAT		
Prevotella oralis		79.12	0.55	MLT 99	1		
Parabacteroides distasonis		20.09	0.58		76		
Capnocytophaga ochracea		0.53	0.18		90		
47451332	**Acceptable identification**						**47451332**
					CAT		
Parabacteroides distasonis		92.65	0.53	XYL 90	76		
Prevotella oralis		6.84	0.21		1		
Capnocytophaga ochracea		0.51	0.02		90		
47451400	**Good identification**						**47451400**
Fusobacterium mortiferum		100.0	0.46	ARA 1			
47451500	**Good identification**						**47451500**
					GRA	CAT	
Fusobacterium mortiferum		97.57	0.53	ARA 1	1	5	
Capnocytophaga ochracea		1.32	0.07		1	90	
Bifidobacterium longum subsp. longum		0.59	0.09		99	1	
47451502	**Low discrimination**						**47451502**
					GRA	CAT	
Bifidobacterium adolescentis		44.86	0.37	MLT 99	99	1	
Capnocytophaga ochracea		25.79	0.23	XYL 87	1	90	
Fusobacterium mortiferum		23.46	0.36		1	5	
47451512	**Acceptable identification**						**47451512**
					GRA	CAT	
Bifidobacterium adolescentis		90.38	0.39	MLT 99	99	1	
Capnocytophaga ochracea		4.72	0.07	XYL 87	1	90	
Bifidobacterium longum subsp. longum		3.59	0.13		99	1	
47451522	**Low discrimination**						**47451522**
					CAT		
Capnocytophaga ochracea		59.28	0.40	MLT 99	90		
Parabacteroides distasonis		26.20	0.46		76		
Prevotella oralis		11.90	0.27		1		
47451532	**Low discrimination**						**47451532**
					GRA	CAT	
Parabacteroides distasonis		61.01	0.41	XYL 90	1	76	
Capnocytophaga ochracea		28.49	0.23		1	90	
Bifidobacterium longum subsp. longum		9.62	0.24		99	1	
47451722	**Acceptable identification**						**47451722**
Parabacteroides distasonis		91.38	0.54	XYL 90			
Prevotella oralis		5.93	0.21				
Bacteroides uniformis		0.85	0.01				
47451732	**Good identification**						**47451732**
Parabacteroides distasonis		96.04	0.50	XYL 90			
Clostridium clostridioforme		3.38	0.19				
Bacteroides thetaiotaomicron		0.58	-.05				
47452000	**Low discrimination**						**47452000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Fusobacterium mortiferum		87.97	0.43	MAN 1	1	5	
Propionibacterium granulosum		7.07	0.21		99	90	
Eubacterium limosum		1.95	-.03		99	1	
47452100	**Good identification**						**47452100**
					GRA		
Fusobacterium mortiferum		99.37	0.50	MAN 1	1		
Actinomyces naeslundii		0.64	0.24		99		
47453000	**Low discrimination**						**47453000**
					GRA	SPO	
Fusobacterium mortiferum		74.95	0.46	MAN 1	1	1	
Clostridium innocuum		14.75	0.22		99	99	
Actinomyces naeslundii		6.29	0.40		99	1	
47453001	**Acceptable identification**						**47453001**
					GRA		
Actinomyces naeslundii		92.57	0.43	ESL 12	99		
Fusobacterium mortiferum		7.43	0.12	MLT 94	1		
47453003	**Good identification**						**47453003**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47453020	**Low discrimination**						**47453020**
					GRA	CAT	
Actinomyces naeslundii		78.11	0.40	ESL 12	99	25	
Propionibacterium granulosum		12.48	0.11	MLT 94	99	90	
Fusobacterium mortiferum		9.41	0.12		1	5	
47453021	**Good identification**						**47453021**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47453022	**Low discrimination**						**47453022**
					GRA		
Actinomyces naeslundii		71.85	0.40	ESL 12	99		
Prevotella oralis		28.16	0.07	MLT 94	1		
47453023	**Good identification**						**47453023**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47453100	**Low discrimination**						**47453100**
					GRA		
Fusobacterium mortiferum		89.86	0.53	MAN 1	1		
Actinomyces naeslundii		9.19	0.47		99		
Actinomyces israelii		0.51	0.17		99		
47453101	**Acceptable identification**						**47453101**
					GRA		
Actinomyces naeslundii		93.82	0.50	ESL 12	99		
Fusobacterium mortiferum		6.18	0.19	MLT 94	1		
47453102	**Low discrimination**						**47453102**
					GRA		
Fusobacterium mortiferum		35.19	0.36	MAN 1	1		
Actinomyces naeslundii		32.39	0.47	BGL 10	99		
Bifidobacterium breve		15.61	0.27		99		
47453103	**Good identification**						**47453103**
Actinomyces naeslundii		100.0	0.50	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47453120	**Low discrimination**					**47453120**
					GRA	
Actinomyces naeslundii		53.64	0.47	ESL 12	99	
Prevotella melaninogenica		24.41	0.12	MLT 94	1	
Prevotella oralis		6.09	0.06		1	
47453121	**Good identification**					**47453121**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47453123	**Good identification**					**47453123**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47453141	**Good identification**					**47453141**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47453143	**Good identification**					**47453143**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47453161	**Good identification**					**47453161**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47453163	**Good identification**					**47453163**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47453502	**Good identification**					**47453502**
Bifidobacterium adolescentis		96.15	0.36	MLT 99		
Actinomyces israelii		3.85	0.18	XYL 87		
47453512	**Good identification**					**47453512**
Bifidobacterium adolescentis		100.0	0.37	MLT 99		
47454000	**Very good identification**					**47454000**
Fusobacterium mortiferum		100.0	0.58	XYL 7		
47454002	**Good identification**					**47454002**
Fusobacterium mortiferum		100.0	0.42	XYL 7		
47454100	**Very good identification**					**47454100**
Fusobacterium mortiferum		100.0	0.65	XYL 7		
47454102	**Good identification**					**47454102**
					CAT	
Fusobacterium mortiferum		98.98	0.48	XYL 7	5	
Bacteroides fragilis		1.02	-.09	BGL 10	99	
47454322	**Acceptable identification**					**47454322**
					CAT	
Parabacteroides distasonis		92.57	0.40	GAL 99	76	
Prevotella oralis		5.01	0.06		1	
Bacteroides fragilis		2.43	-.09		99	
47454332	**Good identification**					**47454332**
Parabacteroides distasonis		95.14	0.35	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		4.86	0.08				
47454722	**Low discrimination**						**47454722**
Parabacteroides distasonis		79.57	0.36	GAL 99			
Clostridium clostridioforme		12.28	0.17				
Bacteroides uniformis		8.15	0.01				
47455000	**Very good identification**						**47455000**
Fusobacterium mortiferum		100.0	0.61	XYL 7			
47455002	**Acceptable identification**						**47455002**
					CAT		
Fusobacterium mortiferum		94.91	0.45	XYL 7	5		
Prevotella oralis		2.43	0.04	BGL 10	1		
Bacteroides fragilis		1.52	-.09		99		
47455100	**Very good identification**						**47455100**
					CAT		
Fusobacterium mortiferum		99.03	0.68	XYL 7	5		
Bacteroides fragilis		0.39	0.04		99		
Capnocytophaga ochracea		0.26	0.10		90		
47455102	**Low discrimination**						**47455102**
					GRA	CAT	
Fusobacterium mortiferum		41.28	0.51	XYL 7	1	5	
Bacteroides fragilis		27.96	0.25	BGL 10	1	99	
Bifidobacterium adolescentis		10.59	0.37		99	1	
47455112	**Low discrimination**						**47455112**
					GRA	CAT	
Bifidobacterium adolescentis		78.28	0.39	MLT 99	99	1	
Parabacteroides distasonis		11.70	0.27	ARA 87	1	76	
Capnocytophaga ochracea		5.80	0.10		1	90	
47455120	**Low discrimination**						**47455120**
					CAT		
Bacteroides fragilis		84.27	0.38	MLT 99	99		
Capnocytophaga ochracea		4.99	0.26	BGL 95	90		
Prevotella oralis		4.10	0.24		1		
47455122	**Low discrimination**						**47455122**
					CAT		
Bacteroides fragilis		82.84	0.60	MLT 99	99		
Parabacteroides distasonis		10.64	0.66		76		
Prevotella oralis		4.03	0.46		1		
47455132	**Low discrimination**						**47455132**
Parabacteroides distasonis		83.46	0.61				
Bacteroides fragilis		12.19	0.25				
Capnocytophaga ochracea		3.76	0.26				
47455222	**Low discrimination**						**47455222**
					CAT		
Parabacteroides distasonis		73.11	0.40	LAC 99	76		
Prevotella oralis		24.98	0.19		1		
Bacteroides fragilis		1.92	-.09		99		
47455232	**Good identification**						**47455232**
Parabacteroides distasonis		95.62	0.35	LAC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		4.38	0.07				
47455302	**Low discrimination**						**47455302**
Parabacteroides distasonis		73.27	0.40	NAG 99			
Prevotella oralis		20.64	0.18				
Fusobacterium mortiferum		2.84	0.17				
47455312	**Low discrimination**						**47455312**
Parabacteroides distasonis		86.71	0.35	NAG 99			
Clostridium clostridioforme		13.29	0.16				
47455322	**Good identification**						**47455322**
					CAT		
Parabacteroides distasonis		92.03	0.74		76		
Prevotella oralis		4.98	0.40		1		
Bacteroides fragilis		2.41	0.25		99		
47455323	**Acceptable identification**						**47455323**
					CAT		
Parabacteroides distasonis		92.57	0.40	URE 1	76		
Prevotella oralis		5.01	0.06		1		
Bacteroides fragilis		2.43	-.09		99		
47455330	**Good identification**						**47455330**
Parabacteroides distasonis		100.0	0.35	BGL 99			
47455332	**Very good identification**						**47455332**
Parabacteroides distasonis		99.70	0.70				
Clostridium clostridioforme		0.15	0.16				
Prevotella oralis		0.10	0.06				
47455333	**Good identification**						**47455333**
Parabacteroides distasonis		100.0	0.35	URE 1			
47455362	**Acceptable identification**						**47455362**
					CAT		
Parabacteroides distasonis		92.57	0.40	SOR 1	76		
Prevotella oralis		5.01	0.06		1		
Bacteroides fragilis		2.43	-.09		99		
47455372	**Good identification**						**47455372**
Parabacteroides distasonis		100.0	0.35	SOR 1			
47455502	**Acceptable identification**						**47455502**
					GRA	CAT	
Bifidobacterium adolescentis		90.05	0.52	MLT 99	99	1	
Parabacteroides distasonis		2.80	0.28		1	76	
Capnocytophaga ochracea		1.93	0.13		1	90	
47455512	**Acceptable identification**						**47455512**
Bifidobacterium adolescentis		94.93	0.53	MLT 99			
Clostridium clostridioforme		2.08	0.21				
Bifidobacterium longum subsp. longum		1.69	0.21				
47455522	**Low discrimination**						**47455522**
					CAT		
Parabacteroides distasonis		69.07	0.62		76		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		15.37	0.32		5	
Bacteroides fragilis		8.86	0.25		99	
47455532	**Good identification**					**47455532**
					GRA	
Parabacteroides distasonis		90.96	0.58		1	
Bifidobacterium longum subsp. longum		4.78	0.32		99	
Clostridium clostridioforme		1.47	0.21		50	
47455542	**Good identification**					**47455542**
Bifidobacterium adolescentis		96.97	0.36	MLT 99		
Actinomyces israelii		3.03	0.16	SOR 11		
47455552	**Good identification**					**47455552**
Bifidobacterium adolescentis		100.0	0.37	MLT 99		
47455622	**Low discrimination**					**47455622**
Parabacteroides distasonis		71.93	0.36	LAC 99		
Bacteroides uniformis		14.89	0.06			
Clostridium clostridioforme		9.95	0.16			
47455702	**Low discrimination**					**47455702**
					GRA	
Parabacteroides distasonis		59.74	0.36	NAG 99	1	
Clostridium clostridioforme		27.67	0.25		50	
Bifidobacterium adolescentis		6.48	0.17		99	
47455712	**Low discrimination**					**47455712**
					GRA	
Clostridium clostridioforme		75.87	0.36	MLT 99	50	
Parabacteroides distasonis		19.36	0.32	SAL 88	1	
Bifidobacterium adolescentis		4.77	0.19	TRE 88	99	
47455720	**Good identification**					**47455720**
Bacteroides vulgatus		96.77	0.44	MLT 99		
Parabacteroides distasonis		2.93	0.36	CEL 5		
Bacteroides uniformis		0.30	0.01			
47455722	**Low discrimination**					**47455722**
					CAT	
Parabacteroides distasonis		88.03	0.71		76	
Bacteroides uniformis		9.02	0.35		5	
Bacteroides thetaiotaomicron		1.95	0.26		89	
47455723	**Acceptable identification**					**47455723**
Parabacteroides distasonis		90.71	0.36	URE 1		
Bacteroides uniformis		9.29	0.01			
47455732	**Good identification**					**47455732**
Parabacteroides distasonis		94.35	0.66			
Clostridium clostridioforme		3.73	0.36			
Bacteroides thetaiotaomicron		1.51	0.19			
47455762	**Acceptable identification**					**47455762**
Parabacteroides distasonis		90.71	0.36	SOR 1		
Bacteroides uniformis		9.29	0.01			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47457102	**Low discrimination**					**47457102**
					GRA	
Bifidobacterium adolescentis		60.87	0.36	MLT 99	99	
Actinomyces israelii		34.22	0.38	ARA 87	99	
Fusobacterium mortiferum		2.93	0.17		1	
47457112	**Acceptable identification**					**47457112**
Bifidobacterium adolescentis		93.57	0.37	MLT 99		
Actinomyces israelii		6.43	0.24	ARA 87		
47457322	**Acceptable identification**					**47457322**
					CAT	
Parabacteroides distasonis		92.57	0.40	MAN 1	76	
Prevotella oralis		5.01	0.06		1	
Bacteroides fragilis		2.43	-.09		99	
47457332	**Good identification**					**47457332**
Parabacteroides distasonis		100.0	0.35	MAN 1		
47457502	**Acceptable identification**					**47457502**
Bifidobacterium adolescentis		93.57	0.50	MLT 99		
Actinomyces israelii		6.43	0.36			
47457512	**Good identification**					**47457512**
Bifidobacterium adolescentis		98.73	0.52	MLT 99		
Actinomyces israelii		0.83	0.22			
Bifidobacterium dentium		0.44	-.07			
47457552	**Good identification**					**47457552**
Bifidobacterium adolescentis		97.42	0.36	MLT 99		
Actinomyces israelii		2.58	0.15	SOR 11		
47457722	**Low discrimination**					**47457722**
Parabacteroides distasonis		85.80	0.36	MAN 1		
Bacteroides uniformis		8.79	0.01			
Bacteroides ovatus		5.41	0.00			
47460100	**Good identification**					**47460100**
Fusobacterium mortiferum		100.0	0.41	FRU 99		
47461000	**Good identification**					**47461000**
Fusobacterium mortiferum		100.0	0.38	FRU 99		
47461100	**Good identification**					**47461100**
					GRA	
Fusobacterium mortiferum		98.23	0.44	FRU 99	1	
Actinomyces naeslundii		1.77	0.26		99	
47465122	**Good identification**					**47465122**
Parabacteroides distasonis		98.30	0.43	FRU 99		
Bacteroides fragilis		1.70	-.09			
47465132	**Good identification**					**47465132**
Parabacteroides distasonis		100.0	0.39	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47465322	**Good identification**						**47465322**
Parabacteroides distasonis		100.0	0.51	FRU 99			
47465332	**Good identification**						**47465332**
Parabacteroides distasonis		100.0	0.47	FRU 99			
47465522	**Good identification**						**47465522**
Parabacteroides distasonis		100.0	0.39	FRU 99			
47465722	**Acceptable identification**						**47465722**
Parabacteroides distasonis		93.26	0.48	FRU 99			
Bacteroides thetaiotaomicron		6.04	0.11				
Clostridium clostridioforme		0.70	0.06				
47465732	**Acceptable identification**						**47465732**
Parabacteroides distasonis		90.05	0.43	FRU 99			
Clostridium clostridioforme		5.74	0.17				
Bacteroides thetaiotaomicron		4.21	0.04				
47470000	**Very good identification**						**47470000**
					GRA	CAT	
Fusobacterium mortiferum		98.93	0.69		1	5	
Propionibacterium granulosum		1.07	0.32		99	90	
47470002	**Good identification**						**47470002**
					GRA	CAT	
Fusobacterium mortiferum		97.63	0.52	BGL 10	1	5	
Propionibacterium granulosum		2.37	0.21		99	90	
47470100	**Very good identification**						**47470100**
					GRA		
Fusobacterium mortiferum		99.98	0.75		1		
Actinomyces naeslundii		0.02	0.24		99		
47470101	**Good identification**						**47470101**
					GRA		
Fusobacterium mortiferum		96.89	0.41	URE 1	1		
Actinomyces naeslundii		3.11	0.27		99		
47470102	**Very good identification**						**47470102**
					GRA		
Fusobacterium mortiferum		99.81	0.59	BGL 10	1		
Actinomyces naeslundii		0.19	0.24		99		
47470110	**Good identification**						**47470110**
Fusobacterium mortiferum		100.0	0.41	MLZ 1			
47470120	**Good identification**						**47470120**
					GRA		
Fusobacterium mortiferum		97.91	0.41	NAG 1	1		
Actinomyces naeslundii		2.10	0.24		99		
47470140	**Good identification**						**47470140**
Fusobacterium mortiferum		100.0	0.41	SOR 1			
47470300	**Good identification**						**47470300**
Fusobacterium mortiferum		100.0	0.41	RHA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47470500	**Good identification**						**47470500**
Fusobacterium mortiferum		100.0	0.41	ARA 1			
47471000	**Very good identification**						**47471000**
					GRA	CAT	
Fusobacterium mortiferum		98.53	0.72		1	5	
Propionibacterium granulosum		0.71	0.32		99	90	
Lactobacillus jensenii		0.48	0.22		99	1	
47471001	**Low discrimination**						**47471001**
					GRA		
Fusobacterium mortiferum		70.37	0.38	URE 1	1		
Actinomyces naeslundii		29.63	0.43		99		
47471002	**Acceptable identification**						**47471002**
					GRA		
Fusobacterium mortiferum		92.19	0.55	BGL 10	1		
Lactobacillus jensenii		2.69	0.19		99		
Actinomyces naeslundii		2.35	0.40		99		
47471003	**Low discrimination**						**47471003**
					GRA		
Actinomyces naeslundii		79.12	0.43	ESL 12	99		
Fusobacterium mortiferum		20.88	0.21	MLT 94	1		
47471010	**Low discrimination**						**47471010**
					GRA	CAT	
Fusobacterium mortiferum		79.52	0.38	MLZ 1	1	5	
Lactobacillus jensenii		12.75	0.14		99	1	
Propionibacterium granulosum		7.74	0.17		99	90	
47471020	**Low discrimination**						**47471020**
					GRA	CAT	
Fusobacterium mortiferum		59.96	0.38	NAG 1	1	5	
Actinomyces naeslundii		16.83	0.40		99	25	
Propionibacterium granulosum		10.69	0.21		99	90	
47471021	**Good identification**						**47471021**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47471023	**Good identification**						**47471023**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47471040	**Good identification**						**47471040**
					GRA		
Fusobacterium mortiferum		95.28	0.38	SOR 1	1		
Actinomyces naeslundii		4.72	0.27		99		
47471100	**Very good identification**						**47471100**
					COC	GRA	
Fusobacterium mortiferum		99.26	0.78		1	1	
Atopobium parvulum		0.37	0.25		99	99	
Actinomyces naeslundii		0.34	0.47		1	99	
47471101	**Low discrimination**						**47471101**
					GRA		
Fusobacterium mortiferum		66.07	0.44	URE 1	1		
Actinomyces naeslundii		33.93	0.50		99		
47471102	**Low discrimination**						**47471102**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		87.40	0.62	BGL 10	1	
Lactobacillus acidophilus		4.55	0.27		99	
Actinomyces naeslundii		2.72	0.47		99	
47471103	**Low discrimination**					**47471103**
					GRA	
Actinomyces naeslundii		82.21	0.50	ESL 12	99	
Fusobacterium mortiferum		17.79	0.27	MLT 94	1	
47471110	**Good identification**					**47471110**
					GRA	
Fusobacterium mortiferum		96.91	0.44	MLZ 1	1	
Actinomyces naeslundii		1.75	0.26		99	
Leptotrichia buccalis		1.35	-.11		1	
47471120	**Low discrimination**					**47471120**
					GRA	
Fusobacterium mortiferum		43.99	0.44	NAG 1	1	
Actinomyces naeslundii		15.06	0.47		99	
Prevotella oralis		10.81	0.19		1	
47471121	**Good identification**					**47471121**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47471122	**Low discrimination**					**47471122**
					GRA	
Parabacteroides distasonis		65.23	0.61	XYL 90	1	
Prevotella oralis		25.19	0.41		1	
Lactobacillus acidophilus		3.09	0.27		99	
47471123	**Low discrimination**					**47471123**
					GRA	
Actinomyces naeslundii		75.22	0.50	ESL 12	99	
Parabacteroides distasonis		17.88	0.27	MLT 94	1	
Prevotella oralis		6.90	0.07		1	
47471132	**Good identification**					**47471132**
					GRA	
Parabacteroides distasonis		93.71	0.56	XYL 90	1	
Bifidobacterium breve		2.82	0.29		99	
Leptotrichia buccalis		2.20	0.07		1	
47471140	**Acceptable identification**					**47471140**
					GRA	
Fusobacterium mortiferum		94.30	0.44	SOR 1	1	
Actinomyces naeslundii		5.70	0.35		99	
47471141	**Good identification**					**47471141**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47471143	**Good identification**					**47471143**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47471161	**Good identification**					**47471161**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47471163	**Good identification**					**47471163**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47471200	**Good identification**					**47471200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.38	RHA 1			
47471300	**Good identification**						**47471300**
Fusobacterium mortiferum		100.0	0.44	RHA 1			
47471322	**Good identification**						**47471322**
					SPO		
Parabacteroides distasonis		94.13	0.69	XYL 90	1		
Prevotella oralis		5.19	0.35		1		
Clostridium ramosum		0.57	0.19		92		
47471332	**Very good identification**						**47471332**
Parabacteroides distasonis		99.68	0.65	XYL 90			
Clostridium clostridioforme		0.22	0.13				
Prevotella oralis		0.10	0.01				
47471400	**Good identification**						**47471400**
Fusobacterium mortiferum		100.0	0.38	ARA 1			
47471500	**Acceptable identification**						**47471500**
					GRA	SPO	
Fusobacterium mortiferum		93.03	0.44	ARA 1	1	1	
Clostridium butyricum		6.98	0.06		99	99	
47471512	**Low discrimination**						**47471512**
					GRA		
Bifidobacterium adolescentis		79.80	0.36	MLT 99	99		
Clostridium clostridioforme		12.74	0.19	XYL 87	50		
Parabacteroides distasonis		5.55	0.19		1		
47471522	**Very good identification**						**47471522**
					CAT		
Parabacteroides distasonis		97.82	0.57	XYL 90	76		
Prevotella oralis		0.62	0.07		1		
Bacteroides thetaiotaomicron		0.54	0.03		89		
47471532	**Good identification**						**47471532**
					GRA		
Parabacteroides distasonis		96.54	0.53	XYL 90	1		
Clostridium clostridioforme		2.24	0.19		50		
Bifidobacterium longum subsp. longum		0.84	0.13		99		
47471722	**Very good identification**						**47471722**
Parabacteroides distasonis		96.91	0.65	XYL 90			
Bacteroides thetaiotaomicron		2.35	0.22				
Clostridium clostridioforme		0.65	0.23				
47471732	**Good identification**						**47471732**
Parabacteroides distasonis		93.09	0.61	XYL 90			
Clostridium clostridioforme		5.28	0.34				
Bacteroides thetaiotaomicron		1.63	0.15				
47472100	**Good identification**						**47472100**
					GRA		
Fusobacterium mortiferum		96.01	0.41	MAN 1	1		
Actinomyces naeslundii		3.99	0.29		99		
47473000	**Low discrimination**						**47473000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Fusobacterium mortiferum		42.44	0.38	MAN 1	1	5	
Actinomyces naeslundii		23.13	0.45		99	25	
Propionibacterium granulosum		22.84	0.30		99	90	
47473001	**Good identification**						**47473001**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47473002	**Low discrimination**						**47473002**
					SPO	CAT	
Actinomyces naeslundii		50.25	0.45	ESL 12	1	25	
Clostridium innocuum		21.34	0.14	MLT 94	99	1	
Propionibacterium granulosum		12.41	0.19		1	90	
47473003	**Good identification**						**47473003**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47473020	**Low discrimination**						**47473020**
Actinomyces naeslundii		80.20	0.45	ESL 12			
Propionibacterium granulosum		19.80	0.19	MLT 94			
47473021	**Good identification**						**47473021**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47473022	**Acceptable identification**						**47473022**
Actinomyces naeslundii		94.19	0.45	ESL 12			
Propionibacterium granulosum		5.81	0.09	MLT 94			
47473023	**Good identification**						**47473023**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47473100	**Low discrimination**						**47473100**
					GRA		
Fusobacterium mortiferum		57.60	0.44	MAN 1	1		
Actinomyces naeslundii		38.29	0.52		99		
Actinomyces israelii		4.11	0.27		99		
47473101	**Good identification**						**47473101**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47473102	**Low discrimination**						**47473102**
					GRA		
Actinomyces naeslundii		61.49	0.52	ESL 12	99		
Fusobacterium mortiferum		10.28	0.27	MLT 94	1		
Bifidobacterium breve		10.18	0.24		99		
47473103	**Good identification**						**47473103**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47473120	**Low discrimination**						**47473120**
					SPO		
Actinomyces naeslundii		72.37	0.52	ESL 12	1		
Clostridium ramosum		25.53	0.19	MLT 94	92		
Bifidobacterium breve		2.11	0.12		1		
47473121	**Good identification**						**47473121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	ESL 12		
47473123	**Good identification**					**47473123**
Actinomyces naeslundii		100.0	0.55	ESL 12		
47473140	**Low discrimination**					**47473140**
Actinomyces naeslundii		80.87	0.39	ESL 12		
Actinomyces israelii		19.13	0.20	MLT 94		
47473141	**Good identification**					**47473141**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47473142	**Low discrimination**					**47473142**
Actinomyces naeslundii		43.62	0.39	ESL 12		
Bifidobacterium breve		40.91	0.24	MLT 94		
Actinomyces israelii		15.48	0.23			
47473143	**Good identification**					**47473143**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47473160	**Low discrimination**					**47473160**
Actinomyces naeslundii		85.84	0.39	ESL 12		
Bifidobacterium breve		14.16	0.12	MLT 94		
47473161	**Good identification**					**47473161**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47473163	**Good identification**					**47473163**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47474000	**Good identification**					**47474000**
Fusobacterium mortiferum		100.0	0.49	XYL 7		
47474100	**Good identification**					**47474100**
Fusobacterium mortiferum		100.0	0.56	XYL 7		
47474102	**Good identification**					**47474102**
Fusobacterium mortiferum		100.0	0.39	XYL 7		
47474122	**Good identification**					**47474122**
Parabacteroides distasonis		98.30	0.43	GAL 99		
Bacteroides fragilis		1.70	-.09			
47474132	**Good identification**					**47474132**
Parabacteroides distasonis		96.74	0.39	GAL 99		
Clostridium clostridioforme		3.26	0.08			
47474322	**Good identification**					**47474322**
Parabacteroides distasonis		99.04	0.51	GAL 99		
Clostridium clostridioforme		0.96	0.11			
47474332	**Acceptable identification**					**47474332**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		92.40	0.47	GAL 99				
Clostridium clostridioforme		7.60	0.22					
47474522	**Acceptable identification**							**47474522**
Parabacteroides distasonis		90.77	0.39	GAL 99				
Clostridium clostridioforme		9.23	0.17					
47474712	**Good identification**							**47474712**
Clostridium clostridioforme		100.0	0.43	MLT 99				
47474722	**Low discrimination**							**47474722**
Parabacteroides distasonis		76.15	0.48	GAL 99				
Clostridium clostridioforme		18.92	0.32					
Bacteroides thetaiotaomicron		4.93	0.11					
47474732	**Low discrimination**							**47474732**
Clostridium clostridioforme		66.76	0.43	MLT 99				
Parabacteroides distasonis		31.75	0.43	SAL 88				
Bacteroides thetaiotaomicron		1.48	0.04					
47475000	**Good identification**							**47475000**
					GRA			
Fusobacterium mortiferum		98.88	0.52	XYL 7	1			
Actinomyces israelii		1.12	0.22		99			
47475002	**Low discrimination**							**47475002**
					GRA			
Fusobacterium mortiferum		86.73	0.36	XYL 7	1			
Actinomyces israelii		13.27	0.25	BGL 10	99			
47475022	**Good identification**							**47475022**
					CAT			
Parabacteroides distasonis		95.17	0.43	LAC 99	76			
Prevotella oralis		3.19	0.05		1			
Bacteroides fragilis		1.64	-.09		99			
47475032	**Good identification**							**47475032**
Parabacteroides distasonis		97.07	0.39	LAC 99				
Clostridium clostridioforme		2.93	0.07					
47475100	**Low discrimination**							**47475100**
					COC	GRA	SPO	
Fusobacterium mortiferum		87.09	0.59	XYL 7	1	1	1	
Atopobium parvulum		8.07	0.30		99	99	1	
Clostridium butyricum		3.15	0.15		1	99	99	
47475102	**Low discrimination**							**47475102**
					GRA			
Parabacteroides distasonis		37.80	0.43	NAG 99	1			
Fusobacterium mortiferum		28.48	0.42		1			
Bifidobacterium adolescentis		16.31	0.34		99			
47475112	**Low discrimination**							**47475112**
					GRA			
Parabacteroides distasonis		46.32	0.39	NAG 99	1			
Bifidobacterium adolescentis		45.37	0.36		99			
Clostridium clostridioforme		4.68	0.16		50			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47475120	**Low discrimination**						**47475120**
					COC	GRA	
Parabacteroides distasonis		75.86	0.43	BGL 99	1	1	
Atopobium parvulum		8.42	0.17		99	99	
Bacteroides fragilis		6.82	0.04		1	1	
47475122	**Very good identification**						**47475122**
					CAT		
Parabacteroides distasonis		97.71	0.77		76		
Bacteroides fragilis		1.69	0.25		99		
Prevotella oralis		0.52	0.26		1		
47475123	**Good identification**						**47475123**
Parabacteroides distasonis		98.30	0.43	URE 1			
Bacteroides fragilis		1.70	-.09				
47475130	**Good identification**						**47475130**
Parabacteroides distasonis		100.0	0.39	BGL 99			
47475132	**Very good identification**						**47475132**
Parabacteroides distasonis		99.87	0.73				
Clostridium clostridioforme		0.10	0.16				
Bacteroides fragilis		0.03	-.09				
47475133	**Good identification**						**47475133**
Parabacteroides distasonis		100.0	0.39	URE 1			
47475162	**Good identification**						**47475162**
Parabacteroides distasonis		98.30	0.43	SOR 1			
Bacteroides fragilis		1.70	-.09				
47475172	**Good identification**						**47475172**
Parabacteroides distasonis		100.0	0.39	SOR 1			
47475222	**Good identification**						**47475222**
Parabacteroides distasonis		98.67	0.51	LAC 99			
Clostridium clostridioforme		0.86	0.10				
Prevotella oralis		0.47	-.01				
47475232	**Acceptable identification**						**47475232**
Parabacteroides distasonis		93.14	0.47	LAC 99			
Clostridium clostridioforme		6.87	0.22				
47475302	**Good identification**						**47475302**
					GRA		
Parabacteroides distasonis		96.33	0.51	NAG 99	1		
Clostridium clostridioforme		2.81	0.19		50		
Actinomyces israelii		0.48	0.16		99		
47475312	**Low discrimination**						**47475312**
Parabacteroides distasonis		80.21	0.47	NAG 99			
Clostridium clostridioforme		19.79	0.31				
47475320	**Good identification**						**47475320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		98.24	0.51	BGL 99			
Bacteroides vulgatus		1.37	0.01				
Prevotella oralis		0.39	-.03				
47475322	**Very good identification**						**47475322**
					CAT		
Parabacteroides distasonis		99.84	0.85		76		
Prevotella oralis		0.08	0.19		1		
Bacteroides thetaiotaomicron		0.04	0.11		89		
47475323	**Good identification**						**47475323**
Parabacteroides distasonis		100.0	0.51	URE 1			
47475330	**Good identification**						**47475330**
Parabacteroides distasonis		100.0	0.47	BGL 99			
47475332	**Very good identification**						**47475332**
Parabacteroides distasonis		99.71	0.81				
Clostridium clostridioforme		0.25	0.31				
Bacteroides thetaiotaomicron		0.03	0.04				
47475333	**Good identification**						**47475333**
Parabacteroides distasonis		100.0	0.47	URE 1			
47475362	**Good identification**						**47475362**
Parabacteroides distasonis		100.0	0.51	SOR 1			
47475372	**Good identification**						**47475372**
Parabacteroides distasonis		100.0	0.47	SOR 1			
47475422	**Low discrimination**						**47475422**
Parabacteroides distasonis		89.24	0.39	LAC 99			
Clostridium clostridioforme		8.13	0.16				
Bacteroides thetaiotaomicron		2.64	-.03				
47475502	**Low discrimination**						**47475502**
					GRA	SPO	
Bifidobacterium adolescentis		61.17	0.48	MLT 99	99	1	
Clostridium butyricum		18.17	0.25		99	99	
Parabacteroides distasonis		12.98	0.39		1	1	
47475512	**Low discrimination**						**47475512**
					GRA		
Bifidobacterium adolescentis		73.18	0.50	MLT 99	99		
Clostridium clostridioforme		17.65	0.36		50		
Parabacteroides distasonis		6.84	0.35		1		
47475520	**Low discrimination**						**47475520**
					GRA	SPO	
Parabacteroides distasonis		66.48	0.39	BGL 99	1	1	
Clostridium butyricum		20.81	0.14		99	99	
Bacteroides vulgatus		6.70	0.04		1	1	
47475522	**Very good identification**						**47475522**
Parabacteroides distasonis		97.87	0.74				
Bacteroides thetaiotaomicron		1.43	0.26				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		0.30	0.25			
47475523	**Good identification**					**47475523**
Parabacteroides distasonis		100.0	0.39	URE 1		
47475532	**Very good identification**					**47475532**
Parabacteroides distasonis		95.93	0.69			
Clostridium clostridioforme		2.50	0.36			
Bacteroides thetaiotaomicron		1.01	0.19			
47475562	**Good identification**					**47475562**
Parabacteroides distasonis		100.0	0.39	SOR 1		
47475612	**Good identification**					**47475612**
Clostridium clostridioforme		100.0	0.42	MLT 99		
47475622	**Low discrimination**					**47475622**
Parabacteroides distasonis		73.89	0.48	LAC 99		
Clostridium clostridioforme		16.45	0.31			
Bacteroides thetaiotaomicron		9.66	0.17			
47475632	**Low discrimination**					**47475632**
Clostridium clostridioforme		63.26	0.42	MLT 99		
Parabacteroides distasonis		33.57	0.43	SAL 88		
Bacteroides thetaiotaomicron		3.17	0.09			
47475702	**Low discrimination**					**47475702**
Parabacteroides distasonis		54.14	0.48	NAG 99		
Clostridium clostridioforme		40.36	0.40			
Bacteroides thetaiotaomicron		3.50	0.11			
47475712	**Low discrimination**					**47475712**
Clostridium clostridioforme		85.35	0.51	MLT 99		
Parabacteroides distasonis		13.53	0.43	SAL 88		
Bacteroides thetaiotaomicron		0.63	0.04			
47475720	**Low discrimination**					**47475720**
					CAT	
Parabacteroides distasonis		72.75	0.48	BGL 99	76	
Bacteroides vulgatus		22.00	0.20		3	
Bacteroides thetaiotaomicron		4.71	0.11		89	
47475722	**Good identification**					**47475722**
Parabacteroides distasonis		92.88	0.82			
Bacteroides thetaiotaomicron		6.01	0.45			
Clostridium clostridioforme		0.70	0.40			
47475723	**Acceptable identification**					**47475723**
Parabacteroides distasonis		93.26	0.48	URE 1		
Bacteroides thetaiotaomicron		6.04	0.11			
Clostridium clostridioforme		0.70	0.06			
47475730	**Acceptable identification**					**47475730**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		90.05	0.43	BGL 99			
Clostridium clostridioforme		5.74	0.17				
Bacteroides thetaiotaomicron		4.21	0.04				
47475732	**Low discrimination**						**47475732**
Parabacteroides distasonis		89.63	0.77				
Clostridium clostridioforme		5.71	0.51				
Bacteroides thetaiotaomicron		4.19	0.38				
47475733	**Acceptable identification**						**47475733**
Parabacteroides distasonis		90.05	0.43	URE 1			
Clostridium clostridioforme		5.74	0.17				
Bacteroides thetaiotaomicron		4.21	0.04				
47475762	**Acceptable identification**						**47475762**
Parabacteroides distasonis		91.77	0.48	SOR 1			
Bacteroides thetaiotaomicron		5.94	0.11				
Bacteroides ovatus		1.60	0.02				
47475772	**Low discrimination**						**47475772**
Parabacteroides distasonis		88.33	0.43	SOR 1			
Clostridium clostridioforme		5.63	0.17				
Bacteroides thetaiotaomicron		4.13	0.04				
47477002	**Good identification**						**47477002**
Actinomyces israelii		100.0	0.37	MLT 99			
47477100	**Low discrimination**						**47477100**
					GRA	SPO	
Actinomyces israelii		88.64	0.45	MLT 99	99	1	
Fusobacterium mortiferum		8.13	0.25	SAL 92	1	1	
Clostridium butyricum		3.23	-.01		99	99	
47477102	**Low discrimination**						**47477102**
Actinomyces israelii		76.04	0.48	MLT 99			
Bifidobacterium adolescentis		23.97	0.33	SAL 92			
47477122	**Acceptable identification**						**47477122**
					GRA	CAT	
Parabacteroides distasonis		91.85	0.43	MAN 1	1	76	
Actinomyces israelii		6.57	0.27		99	1	
Bacteroides fragilis		1.59	-.09		1	99	
47477132	**Acceptable identification**						**47477132**
					GRA		
Parabacteroides distasonis		92.25	0.39	MAN 1	1		
Bifidobacterium dentium		5.92	-.03		99		
Actinomyces israelii		1.83	0.13		99		
47477140	**Good identification**						**47477140**
Actinomyces israelii		100.0	0.38	MLT 99			
47477142	**Acceptable identification**						**47477142**
Actinomyces israelii		90.90	0.41	MLT 99			
Bifidobacterium adolescentis		9.11	0.17	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47477322	**Good identification**						**47477322**
					SPO		
Parabacteroides distasonis		99.62	0.51	MAN 1	1		
Clostridium ramosum		0.38	-.03		92		
47477332	**Good identification**						**47477332**
Parabacteroides distasonis		100.0	0.47	MAN 1			
47477402	**Acceptable identification**						**47477402**
Actinomyces israelii		90.57	0.36	MLT 99			
Bifidobacterium adolescentis		9.43	0.13	SAL 92			
47477500	**Low discrimination**						**47477500**
					SPO		
Actinomyces israelii		69.43	0.43	MLT 99	1		
Clostridium butyricum		27.86	0.15	SAL 92	99		
Bifidobacterium adolescentis		2.71	0.13		1		
47477502	**Low discrimination**						**47477502**
					SPO		
Bifidobacterium adolescentis		69.62	0.47	MLT 99	1		
Actinomyces israelii		27.00	0.46		1		
Clostridium butyricum		2.81	0.08		99		
47477512	**Low discrimination**						**47477512**
Bifidobacterium adolescentis		58.36	0.48	MLT 99			
Bifidobacterium dentium		38.87	0.27				
Actinomyces israelii		2.77	0.32				
47477522	**Low discrimination**						**47477522**
					GRA		
Parabacteroides distasonis		82.45	0.39	MAN 1	1		
Actinomyces israelii		7.87	0.26		99		
Bifidobacterium dentium		4.65	-.03		99		
47477542	**Low discrimination**						**47477542**
					COC		
Actinomyces israelii		49.59	0.39	MLT 99	1		
Bifidobacterium adolescentis		40.64	0.31	SAL 92	1		
Blautia producta		9.77	0.07		99		
47477722	**Low discrimination**						**47477722**
Parabacteroides distasonis		82.50	0.48	MAN 1			
Bacteroides ovatus		11.54	0.17				
Bacteroides thetaiotaomicron		5.34	0.11				
47477732	**Low discrimination**						**47477732**
Parabacteroides distasonis		75.85	0.43	MAN 1			
Bacteroides ovatus		13.13	0.14				
Clostridium clostridioforme		4.83	0.17				
47500100	**Low discrimination**						**47500100**
					COC	GRA	
Fusobacterium mortiferum		82.44	0.42	MLT 11	1	1	
Prevotella melaninogenica		15.52	0.12	FRU 99	1	1	
Blautia hansenii		2.04	-.03		99	99	
47500120	**Good identification**						**47500120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Prevotella melaninogenica		98.93	0.47	FRU 90	1		
Capnocytophaga ochracea		0.60	0.12	GAL 99	90		
Prevotella bivia		0.33	-.03		1		
47500520	**Low discrimination**						**47500520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.80	0.37	FRU 90	99	1	
Prevotella melaninogenica		14.64	0.12	GAL 99	1	1	
Capnocytophaga ochracea		1.55	-.01		1	90	
47500522	**Low discrimination**						**47500522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.33	0.37	FRU 90	99	1	
Capnocytophaga ochracea		13.89	0.16	GAL 99	1	90	
Prevotella melaninogenica		2.77	0.00		1	1	
47500530	**Good identification**						**47500530**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90			
47500532	**Good identification**						**47500532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.92	0.41	FRU 90	99	1	
Capnocytophaga ochracea		1.08	-.01	GAL 99	1	90	
47501000	**Low discrimination**						**47501000**
					COC	GRA	
Fusobacterium mortiferum		30.66	0.39	MLT 11	1	1	
Eubacterium saburreum		27.06	0.37	FRU 99	1	85	
Gemella morbillorum		19.88	0.26		99	99	
47501010	**Good identification**						**47501010**
Eubacterium saburreum		100.0	0.35	IND 99			
47501020	**Good identification**						**47501020**
					GRA	CAT	
Prevotella melaninogenica		96.19	0.47	FRU 90	1	1	
Clostridium perfringens		1.80	0.24	LAC 99	99	1	
Capnocytophaga ochracea		0.58	0.12		1	90	
47501100	**Low discrimination**						**47501100**
					COC	GRA	
Prevotella melaninogenica		71.07	0.47	FRU 90	1	1	
Blautia hansenii		9.34	0.32	NAG 99	99	99	
Fusobacterium mortiferum		5.72	0.45		1	1	
47501101	**Very good identification**						**47501101**
					GRA	SPO	
Clostridium chauvoei		98.66	0.57	RAF 1	99	99	
Actinomyces naeslundii		0.62	0.42		99	1	
Prevotella melaninogenica		0.59	0.12		1	1	
47501102	**Low discrimination**						**47501102**
					GRA	CAT	
Capnocytophaga ochracea		59.35	0.47	FRU 99	1	90	
Prevotella melaninogenica		18.82	0.34	NAG 90	1	1	
Bifidobacterium longum subsp. infantis		7.52	0.27		99	1	
47501103	**Acceptable identification**						**47501103**
					GRA	SPO	
Clostridium chauvoei		93.87	0.44	RAF 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		3.37	0.42		99	1	
Capnocytophaga ochracea		1.91	0.12		1	1	
47501110	**Low discrimination**						**47501110**
					GRA		
Eubacterium saburreum		56.76	0.41	IND 99	85		
Bifidobacterium longum subsp. longum		24.04	0.31		99		
Prevotella melaninogenica		9.87	0.12		1		
47501120	**Very good identification**						**47501120**
					GRA	CAT	
Prevotella melaninogenica		99.06	0.81	FRU 90	1	1	
Capnocytophaga ochracea		0.60	0.47		1	90	
Clostridium perfringens		0.14	0.39		99	1	
47501121	**Low discrimination**						**47501121**
					GRA	SPO	
Prevotella melaninogenica		63.28	0.47	FRU 90	1	1	
Clostridium chauvoei		35.57	0.48	URE 1	99	99	
Actinomyces naeslundii		0.68	0.42		99	1	
47501122	**Low discrimination**						**47501122**
					CAT		
Prevotella melaninogenica		72.07	0.68	FRU 90	1		
Capnocytophaga ochracea		20.66	0.63		90		
Prevotella oralis		5.11	0.52		1		
47501130	**Low discrimination**						**47501130**
					GRA	CAT	
Prevotella melaninogenica		84.98	0.47	FRU 90	1	1	
Bifidobacterium longum subsp. longum		8.36	0.41	MLZ 1	99	1	
Capnocytophaga ochracea		5.67	0.30		1	90	
47501132	**Low discrimination**						**47501132**
					GRA	CAT	
Capnocytophaga ochracea		65.63	0.47	FRU 99	1	90	
Prevotella melaninogenica		20.81	0.34	MLZ 10	1	1	
Bifidobacterium longum subsp. longum		10.75	0.41		99	1	
47501160	**Acceptable identification**						**47501160**
					GRA	CAT	
Prevotella melaninogenica		91.40	0.47	FRU 90	1	1	
Capnocytophaga ochracea		6.10	0.30	SOR 1	1	90	
Clostridium perfringens		2.39	0.26		99	1	
47501162	**Low discrimination**						**47501162**
					CAT		
Capnocytophaga ochracea		73.04	0.47	FRU 99	90		
Prevotella melaninogenica		23.16	0.34	SOR 10	1		
Prevotella oralis		1.64	0.18		1		
47501320	**Acceptable identification**						**47501320**
Prevotella melaninogenica		93.87	0.47	FRU 90			
Prevotella oralis		2.83	0.24	RHA 1			
Bacteroides vulgatus		2.60	0.18				
47501322	**Low discrimination**						**47501322**
					CAT		
Prevotella oralis		66.89	0.46	FRU 99	1		
Prevotella melaninogenica		22.24	0.34		1		
Capnocytophaga ochracea		6.38	0.29		90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47501420	**Low discrimination**							**47501420**
					GRA	CAT		
Bifidobacterium longum subsp. longum		83.80	0.37	FRU 90	99	1		
Prevotella melaninogenica		14.64	0.12	LAC 99	1	1		
Capnocytophaga ochracea		1.55	-.01		1	90		
47501422	**Low discrimination**							**47501422**
					GRA	CAT		
Bifidobacterium longum subsp. longum		82.31	0.37	FRU 90	99	1		
Capnocytophaga ochracea		13.72	0.16	LAC 99	1	90		
Prevotella melaninogenica		2.74	0.00		1	1		
47501430	**Good identification**							**47501430**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90				
47501432	**Good identification**							**47501432**
					GRA	CAT		
Bifidobacterium longum subsp. longum		98.92	0.41	FRU 90	99	1		
Capnocytophaga ochracea		1.08	-.01	LAC 99	1	90		
47501500	**Very good identification**							**47501500**
					COC	GRA	CAT	
Bifidobacterium longum subsp. longum		96.67	0.61	FRU 90	1	99	1	
Blautia hansenii		1.57	0.19		99	99	1	
Capnocytophaga ochracea		0.80	0.17		1	1	90	
47501502	**Good identification**							**47501502**
					GRA	CAT		
Bifidobacterium longum subsp. longum		91.29	0.61	FRU 90	99	1		
Capnocytophaga ochracea		6.76	0.34		1	90		
Bifidobacterium adolescentis		1.38	0.32		99	1		
47501510	**Very good identification**							**47501510**
					GRA	CAT		
Bifidobacterium longum subsp. longum		99.95	0.65	FRU 90	99	1		
Capnocytophaga ochracea		0.05	0.01		1	90		
47501512	**Very good identification**							**47501512**
					GRA	CAT		
Bifidobacterium longum subsp. longum		98.45	0.65	FRU 90	99	1		
Bifidobacterium adolescentis		1.07	0.33		99	1		
Capnocytophaga ochracea		0.48	0.17		1	90		
47501520	**Low discrimination**							**47501520**
					GRA	CAT		
Bifidobacterium longum subsp. longum		83.24	0.72	FRU 90	99	1		
Prevotella melaninogenica		14.55	0.47		1	1		
Capnocytophaga ochracea		1.54	0.34		1	90		
47501521	**Low discrimination**							**47501521**
					GRA	SPO		
Bifidobacterium longum subsp. longum		77.43	0.37	FRU 90	99	1		
Prevotella melaninogenica		13.53	0.12	URE 1	1	1		
Clostridium chauvoei		7.61	0.14		99	99		
47501522	**Low discrimination**							**47501522**
					GRA	CAT		
Bifidobacterium longum subsp. longum		83.02	0.72	FRU 90	99	1		
Capnocytophaga ochracea		13.84	0.50		1	90		
Prevotella melaninogenica		2.76	0.34		1	1		
47501523	**Low discrimination**							**47501523**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Bifidobacterium longum subsp. longum		82.15	0.37	FRU 90	99	1	
Capnocytophaga ochracea		13.70	0.16	URE 1	1	90	
Prevotella melaninogenica		2.73	0.00		1	1	
47501530	**Very good identification**						**47501530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.78	0.76	FRU 90	99	1	
Capnocytophaga ochracea		0.12	0.17		1	90	
Prevotella melaninogenica		0.10	0.12		1	1	
47501531	**Good identification**						**47501531**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90			
47501532	**Very good identification**						**47501532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.87	0.76	FRU 90	99	1	
Capnocytophaga ochracea		1.08	0.34		1	90	
Parabacteroides distasonis		0.04	0.21		1	76	
47501533	**Good identification**						**47501533**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.92	0.41	FRU 90	99	1	
Capnocytophaga ochracea		1.08	-.01	URE 1	1	90	
47501560	**Low discrimination**						**47501560**
					GRA	CAT	
Bifidobacterium longum subsp. longum		72.54	0.37	FRU 90	99	1	
Capnocytophaga ochracea		14.78	0.17	SOR 1	1	90	
Prevotella melaninogenica		12.68	0.12		1	1	
47501570	**Good identification**						**47501570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.69	0.41	FRU 90	99	1	
Capnocytophaga ochracea		1.31	0.01	SOR 1	1	90	
47501572	**Low discrimination**						**47501572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.31	0.41	FRU 90	99	1	
Capnocytophaga ochracea		10.69	0.17	SOR 1	1	90	
47501720	**Low discrimination**						**47501720**
					GRA		
Bacteroides vulgatus		84.23	0.37	FRU 99	1		
Bifidobacterium longum subsp. longum		13.21	0.37	XYL 98	99		
Prevotella melaninogenica		2.31	0.12		1		
47501722	**Low discrimination**						**47501722**
					GRA	CAT	
Bifidobacterium longum subsp. longum		57.86	0.37	FRU 90	99	1	
Parabacteroides distasonis		21.05	0.34	RHA 1	1	76	
Capnocytophaga ochracea		9.65	0.16		1	90	
47501730	**Good identification**						**47501730**
					GRA		
Bifidobacterium longum subsp. longum		96.36	0.41	FRU 90	99		
Bacteroides vulgatus		3.64	0.03	RHA 1	1		
47501732	**Low discrimination**						**47501732**
					GRA		
Bifidobacterium longum subsp. longum		87.40	0.41	FRU 90	99		
Parabacteroides distasonis		10.06	0.29	RHA 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		1.59	0.10		50		
47503000	**Low discrimination**						**47503000**
Actinomyces naeslundii		68.42	0.36	ESL 12			
Propionibacterium granulosum		31.58	0.16	FRU 95			
47503001	**Good identification**						**47503001**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47503002	**Good identification**						**47503002**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47503003	**Good identification**						**47503003**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47503021	**Good identification**						**47503021**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47503022	**Low discrimination**						**47503022**
					GRA		
Actinomyces naeslundii		63.52	0.36	ESL 12	99		
Prevotella melaninogenica		25.20	0.00	FRU 95	1		
Prevotella oralis		11.29	-.03		1		
47503023	**Good identification**						**47503023**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47503100	**Low discrimination**						**47503100**
					COC	GRA	
Actinomyces naeslundii		53.00	0.44	ESL 12	1	99	
Prevotella melaninogenica		38.78	0.12	FRU 95	1	1	
Blautia hansenii		5.10	-.03		99	99	
47503101	**Low discrimination**						**47503101**
					SPO		
Actinomyces naeslundii		54.87	0.47	ESL 12	1		
Clostridium chauvoei		45.13	0.22	FRU 95	99		
47503102	**Low discrimination**						**47503102**
					GRA	CAT	
Actinomyces naeslundii		53.12	0.44	ESL 12	99	25	
Capnocytophaga ochracea		23.35	0.12	FRU 95	1	90	
Bifidobacterium adolescentis		10.21	0.16		99	1	
47503103	**Low discrimination**						**47503103**
					SPO		
Actinomyces naeslundii		87.32	0.47	ESL 12	1		
Clostridium chauvoei		12.68	0.10	FRU 95	99		
47503120	**Good identification**						**47503120**
					GRA	CAT	
Prevotella melaninogenica		97.92	0.47	FRU 90	1	1	
Actinomyces naeslundii		1.35	0.44	MAN 1	99	25	
Capnocytophaga ochracea		0.59	0.12		1	90	
47503121	**Low discrimination**						**47503121**
					GRA	SPO	
Actinomyces naeslundii		56.76	0.47	ESL 12	99	1	
Prevotella melaninogenica		27.68	0.12	FRU 95	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		15.56	0.14		99	99	
47503123	**Low discrimination**						**47503123**
					GRA	SPO	
Actinomyces naeslundii		87.62	0.47	ESL 12	99	1	
Prevotella melaninogenica		8.14	0.00	FRU 95	1	1	
Clostridium chauvoei		4.24	0.01		99	99	
47503520	**Low discrimination**						**47503520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.80	0.37	FRU 90	99	1	
Prevotella melaninogenica		14.64	0.12	MAN 1	1	1	
Capnocytophaga ochracea		1.55	-.01		1	90	
47503522	**Low discrimination**						**47503522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.33	0.37	FRU 90	99	1	
Capnocytophaga ochracea		13.89	0.16	MAN 1	1	90	
Prevotella melaninogenica		2.77	0.00		1	1	
47503530	**Good identification**						**47503530**
Bifidobacterium longum subsp. longum		100.0	0.41	FRU 90			
47503532	**Good identification**						**47503532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.92	0.41	FRU 90	99	1	
Capnocytophaga ochracea		1.08	-.01	MAN 1	1	90	
47504500	**Good identification**						**47504500**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90			
47504502	**Good identification**						**47504502**
Bifidobacterium longum subsp. longum		96.73	0.35	FRU 90			
Bifidobacterium adolescentis		3.27	0.12	GAL 99			
47504510	**Good identification**						**47504510**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47504512	**Good identification**						**47504512**
Bifidobacterium longum subsp. longum		96.58	0.39	FRU 90			
Bifidobacterium adolescentis		2.35	0.13	GAL 99			
Clostridium clostridioforme		1.08	0.04				
47504520	**Low discrimination**						**47504520**
					GRA		
Bifidobacterium longum subsp. longum		89.54	0.46	FRU 90	99		
Bacteroides vulgatus		10.46	0.15	GAL 99	1		
47504522	**Good identification**						**47504522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.05	0.46	FRU 90	99	1	
Bacteroides fragilis		1.61	-.03	GAL 99	1	99	
Capnocytophaga ochracea		1.35	0.06		1	90	
47504530	**Good identification**						**47504530**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
47504532	**Good identification**						**47504532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		99.72	0.50	FRU 90			
Clostridium clostridioforme		0.28	0.04	GAL 99			
47505100	**Low discrimination**						**47505100**
					COC		
Bifidobacterium longum subsp. longum		26.72	0.35	FRU 90	1		
Eubacterium saburreum		23.56	0.38	ARA 99	1		
Blautia hansenii		14.29	0.19		99		
47505110	**Low discrimination**						**47505110**
					CAT	GRA	
Bifidobacterium longum subsp. longum		69.02	0.39	FRU 90	1	99	
Eubacterium saburreum		29.25	0.37	ARA 99	1	85	
Capnocytophaga ochracea		1.73	0.04		90	1	
47505112	**Low discrimination**						**47505112**
					GRA	CAT	
Bifidobacterium longum subsp. longum		58.86	0.39	FRU 90	99	1	
Bifidobacterium adolescentis		21.14	0.33	ARA 99	99	1	
Capnocytophaga ochracea		13.31	0.20		1	90	
47505120	**Low discrimination**						**47505120**
					CAT	GRA	
Bacteroides fragilis		49.55	0.44	FRU 99	99	1	
Prevotella melaninogenica		33.50	0.47	BGL 95	1	1	
Bifidobacterium longum subsp. longum		5.81	0.46		1	99	
47505122	**Good identification**						**47505122**
					CAT		
Bacteroides fragilis		93.06	0.66	FRU 99	99		
Capnocytophaga ochracea		4.48	0.53		90		
Prevotella melaninogenica		0.63	0.34		1		
47505130	**Low discrimination**						**47505130**
					GRA	CAT	
Bifidobacterium longum subsp. longum		85.82	0.50	FRU 90	99	1	
Capnocytophaga ochracea		4.85	0.20	ARA 99	1	90	
Bacteroides fragilis		4.34	0.10		1	99	
47505132	**Low discrimination**						**47505132**
					GRA	CAT	
Bifidobacterium longum subsp. longum		37.35	0.50	FRU 90	99	1	
Bacteroides fragilis		35.92	0.32	ARA 99	1	99	
Capnocytophaga ochracea		19.00	0.36		1	90	
47505320	**Good identification**						**47505320**
					CAT		
Bacteroides vulgatus		97.17	0.47	FRU 99	3		
Bacteroides fragilis		1.07	0.10	ARA 93	99		
Prevotella melaninogenica		0.72	0.12		1		
47505322	**Low discrimination**						**47505322**
					CAT		
Parabacteroides distasonis		44.34	0.54	FRU 99	76		
Bacteroides fragilis		40.76	0.32		99		
Prevotella oralis		10.17	0.30		1		
47505332	**Good identification**						**47505332**
					GRA	CAT	
Parabacteroides distasonis		95.12	0.49	FRU 99	1	76	
Bifidobacterium longum subsp. longum		1.71	0.15		99	1	
Bacteroides fragilis		1.64	-.03		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47505400	**Good identification**						**47505400**
Bifidobacterium longum subsp. longum		97.21	0.35	FRU 90			
Lactobacillus fermentum		2.79	0.16	LAC 99			
47505402	**Acceptable identification**						**47505402**
Bifidobacterium longum subsp. longum		94.11	0.35	FRU 90			
Bifidobacterium adolescentis		3.18	0.12	LAC 99			
Lactobacillus fermentum		2.71	0.16				
47505410	**Good identification**						**47505410**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47505412	**Good identification**						**47505412**
Bifidobacterium longum subsp. longum		97.63	0.39	FRU 90			
Bifidobacterium adolescentis		2.37	0.13	LAC 99			
47505420	**Low discrimination**						**47505420**
					GRA		
Bifidobacterium longum subsp. longum		89.54	0.46	FRU 90	99		
Bacteroides vulgatus		10.46	0.15	LAC 99	1		
47505422	**Good identification**						**47505422**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.05	0.46	FRU 90	99	1	
Bacteroides fragilis		1.61	-.03	LAC 99	1	99	
Capnocytophaga ochracea		1.35	0.06		1	90	
47505430	**Good identification**						**47505430**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
47505432	**Good identification**						**47505432**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
47505500	**Very good identification**						**47505500**
					COC	GRA	
Bifidobacterium longum subsp. longum		99.29	0.69	FRU 90	1	99	
Bacteroides vulgatus		0.47	0.15		1	1	
Blautia hansenii		0.10	0.06		99	99	
47505501	**Good identification**						**47505501**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90			
47505502	**Very good identification**						**47505502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.05	0.69	FRU 90	99	1	
Bifidobacterium adolescentis		3.25	0.46		99	1	
Capnocytophaga ochracea		0.59	0.23		1	90	
47505503	**Good identification**						**47505503**
Bifidobacterium longum subsp. longum		96.73	0.35	FRU 90			
Bifidobacterium adolescentis		3.27	0.12	URE 1			
47505510	**Very good identification**						**47505510**
Bifidobacterium longum subsp. longum		99.98	0.73	FRU 90			
Bifidobacterium adolescentis		0.03	0.13				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47505511	**Good identification**						**47505511**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47505512	**Very good identification**						**47505512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.56	0.73	FRU 90	99	1	
Bifidobacterium adolescentis		2.37	0.47		99	1	
Capnocytophaga ochracea		0.04	0.07		1	90	
47505513	**Good identification**						**47505513**
Bifidobacterium longum subsp. longum		97.63	0.39	FRU 90			
Bifidobacterium adolescentis		2.37	0.13	URE 1			
47505520	**Low discrimination**						**47505520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.30	0.80	FRU 90	99	1	
Bacteroides vulgatus		10.43	0.50		1	3	
Capnocytophaga ochracea		0.14	0.23		1	90	
47505521	**Low discrimination**						**47505521**
					GRA		
Bifidobacterium longum subsp. longum		89.54	0.46	FRU 90	99		
Bacteroides vulgatus		10.46	0.15	URE 1	1		
47505522	**Very good identification**						**47505522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.56	0.80	FRU 90	99	1	
Bacteroides fragilis		1.60	0.32		1	99	
Capnocytophaga ochracea		1.34	0.40		1	90	
47505523	**Good identification**						**47505523**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.05	0.46	FRU 90	99	1	
Bacteroides fragilis		1.61	-.03	URE 1	1	99	
Capnocytophaga ochracea		1.35	0.06		1	90	
47505530	**Very good identification**						**47505530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.92	0.84	FRU 90	99	1	
Bacteroides vulgatus		0.07	0.15		1	3	
Capnocytophaga ochracea		0.01	0.07		1	90	
47505531	**Good identification**						**47505531**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
47505532	**Very good identification**						**47505532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.77	0.84	FRU 90	99	1	
Parabacteroides distasonis		0.12	0.38		1	76	
Capnocytophaga ochracea		0.09	0.23		1	90	
47505533	**Good identification**						**47505533**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
47505540	**Good identification**						**47505540**
Bifidobacterium longum subsp. longum		100.0	0.35	FRU 90			
47505542	**Low discrimination**						**47505542**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		67.48	0.35	FRU 90	99	1	
Bifidobacterium adolescentis		27.93	0.30	SOR 1	99	1	
Capnocytophaga ochracea		4.58	0.07		1	90	
47505550	**Good identification**						**47505550**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47505552	**Low discrimination**						**47505552**
Bifidobacterium longum subsp. longum		77.09	0.39	FRU 90			
Bifidobacterium adolescentis		22.91	0.32	SOR 1			
47505560	**Low discrimination**						**47505560**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.20	0.46	FRU 90	99	1	
Bacteroides vulgatus		10.31	0.15	SOR 1	1	3	
Capnocytophaga ochracea		1.50	0.07		1	90	
47505562	**Low discrimination**						**47505562**
					GRA	CAT	
Bifidobacterium longum subsp. longum		85.52	0.46	FRU 90	99	1	
Capnocytophaga ochracea		13.07	0.23	SOR 1	1	90	
Bacteroides fragilis		1.41	-.03		1	99	
47505570	**Good identification**						**47505570**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
47505572	**Good identification**						**47505572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.01	0.50	FRU 90	99	1	
Capnocytophaga ochracea		0.99	0.07	SOR 1	1	90	
47505702	**Low discrimination**						**47505702**
Bifidobacterium longum subsp. longum		77.20	0.35	FRU 90			
Clostridium clostridioforme		7.08	0.16	RHA 1			
Eubacterium contortum		6.42	0.10				
47505710	**Good identification**						**47505710**
					GRA		
Bifidobacterium longum subsp. longum		97.57	0.39	FRU 90	99		
Bacteroides vulgatus		2.44	-.03	RHA 1	1		
47505712	**Low discrimination**						**47505712**
Bifidobacterium longum subsp. longum		77.90	0.39	FRU 90			
Clostridium clostridioforme		19.12	0.27	RHA 1			
Bifidobacterium adolescentis		1.89	0.13				
47505720	**Very good identification**						**47505720**
					GRA		
Bacteroides vulgatus		99.02	0.66	FRU 99	1		
Bifidobacterium longum subsp. longum		0.95	0.46		99		
Eubacterium contortum		0.02	0.10		99		
47505722	**Low discrimination**						**47505722**
					GRA		
Parabacteroides distasonis		33.86	0.50	FRU 99	1		
Bacteroides vulgatus		32.62	0.32		1		
Bifidobacterium longum subsp. longum		31.02	0.46		99		
47505730	**Low discrimination**						**47505730**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium longum subsp. longum		61.69	0.50	FRU 90	99		
Bacteroides vulgatus		38.10	0.32	RHA 1	1		
Parabacteroides distasonis		0.22	0.12		1		
47505732	**Low discrimination**						**47505732**
					GRA		
Bifidobacterium longum subsp. longum		70.48	0.50	FRU 90	99		
Parabacteroides distasonis		24.33	0.46	RHA 1	1		
Clostridium clostridioforme		4.32	0.27		50		
47507500	**Good identification**						**47507500**
Bifidobacterium longum subsp. longum		95.44	0.35	FRU 90			
Bifidobacterium adolescentis		2.64	0.10	MAN 1			
Actinomyces israelii		1.92	0.14				
47507502	**Low discrimination**						**47507502**
Bifidobacterium adolescentis		72.68	0.44	FRU 99			
Bifidobacterium longum subsp. longum		26.52	0.35				
Actinomyces israelii		0.80	0.17				
47507510	**Good identification**						**47507510**
Bifidobacterium longum subsp. longum		98.05	0.39	FRU 90			
Bifidobacterium adolescentis		1.95	0.12	MAN 1			
47507512	**Low discrimination**						**47507512**
Bifidobacterium adolescentis		66.29	0.46	FRU 99			
Bifidobacterium longum subsp. longum		33.71	0.39				
47507520	**Low discrimination**						**47507520**
					GRA		
Bifidobacterium longum subsp. longum		89.54	0.46	FRU 90	99		
Bacteroides vulgatus		10.46	0.15	MAN 1	1		
47507522	**Good identification**						**47507522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.40	0.46	FRU 90	99	1	
Bacteroides fragilis		1.59	-.03	MAN 1	1	99	
Capnocytophaga ochracea		1.34	0.06		1	90	
47507530	**Good identification**						**47507530**
Bifidobacterium longum subsp. longum		100.0	0.50	FRU 90			
47507532	**Good identification**						**47507532**
Bifidobacterium longum subsp. longum		99.51	0.50	FRU 90			
Bifidobacterium adolescentis		0.49	0.12	MAN 1			
47510000	**Low discrimination**						**47510000**
					GRA	CAT	
Fusobacterium mortiferum		70.19	0.70	MLT 11	1	5	
Prevotella intermedia		18.88	0.49		1	1	
Propionibacterium granulosum		9.06	0.52		99	90	
47510001	**Low discrimination**						**47510001**
					GRA		
Fusobacterium mortiferum		65.48	0.36	MLT 11	1		
Prevotella intermedia		17.62	0.15	URE 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		8.45	0.32		99		
47510002	**Low discrimination**						**47510002**
					GRA	CAT	
Fusobacterium mortiferum		65.36	0.54	MLT 11	1	5	
Propionibacterium granulosum		18.99	0.41	BGL 10	99	90	
Actinomyces viscosus		10.07	0.27		99	76	
47510010	**Low discrimination**						**47510010**
					GRA	CAT	
Propionibacterium granulosum		57.87	0.37	ESL 1	99	90	
Fusobacterium mortiferum		33.20	0.36	MLZ 12	1	5	
Prevotella intermedia		8.93	0.15		1	1	
47510100	**Very good identification**						**47510100**
					CAT		
Fusobacterium mortiferum		96.46	0.77	MLT 11	5		
Prevotella melaninogenica		1.65	0.29		1		
Capnocytophaga ochracea		1.21	0.30		90		
47510101	**Low discrimination**						**47510101**
					GRA	SPO	
Fusobacterium mortiferum		52.20	0.42	MLT 11	1	1	
Clostridium chauvoei		38.69	0.29	URE 1	99	99	
Actinomyces naeslundii		8.22	0.40		99	1	
47510102	**Low discrimination**						**47510102**
					GRA	CAT	
Capnocytophaga ochracea		41.63	0.47	GAL 99	1	90	
Fusobacterium mortiferum		40.89	0.60	NAG 90	1	5	
Bifidobacterium longum subsp. infantis		5.27	0.27		99	1	
47510110	**Low discrimination**						**47510110**
					GRA	CAT	
Fusobacterium mortiferum		81.31	0.42	MLT 11	1	5	
Capnocytophaga ochracea		11.24	0.14	MLZ 1	1	90	
Bifidobacterium longum subsp. longum		3.39	0.13		99	1	
47510120	**Good identification**						**47510120**
					GRA	CAT	
Prevotella melaninogenica		90.86	0.63	GAL 99	1	1	
Capnocytophaga ochracea		6.07	0.47		1	90	
Clostridium perfringens		1.38	0.39		99	1	
47510122	**Low discrimination**						**47510122**
					CAT		
Capnocytophaga ochracea		60.27	0.63	GAL 99	90		
Prevotella melaninogenica		19.11	0.51		1		
Prevotella oralis		14.90	0.52		1		
47510132	**Acceptable identification**						**47510132**
					CAT		
Capnocytophaga ochracea		92.29	0.47	GAL 99	90		
Prevotella melaninogenica		2.66	0.16	MLZ 10	1		
Prevotella oralis		2.08	0.18		1		
47510140	**Low discrimination**						**47510140**
					GRA	CAT	
Fusobacterium mortiferum		84.11	0.42	MLT 11	1	5	
Capnocytophaga ochracea		11.63	0.14	SOR 1	1	90	
Actinomyces naeslundii		1.56	0.24		99	25	
47510162	**Acceptable identification**						**47510162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Capnocytophaga ochracea		92.55	0.47	GAL 99	90		
Prevotella melaninogenica		2.67	0.16	SOR 10	1		
Prevotella oralis		2.08	0.18		1		
47510200	**Low discrimination**						**47510200**
					GRA	CAT	
Fusobacterium mortiferum		60.09	0.36	MLT 11	1	5	
Propionibacterium granulosum		23.75	0.26	RHA 1	99	90	
Prevotella intermedia		16.16	0.15		1	1	
47510300	**Good identification**						**47510300**
Fusobacterium mortiferum		95.54	0.42	MLT 11			
Prevotella oralis		2.83	0.02	RHA 1			
Prevotella melaninogenica		1.64	-.06				
47510322	**Low discrimination**						**47510322**
					CAT		
Prevotella oralis		83.73	0.46	GAL 99	1		
Capnocytophaga ochracea		7.98	0.29		90		
Parabacteroides distasonis		5.02	0.38		76		
47510400	**Low discrimination**						**47510400**
					COC	GRA	
Fusobacterium mortiferum		55.78	0.36	MLT 11	1	1	
Prevotella intermedia		15.00	0.15	ARA 1	1	1	
Gemella morbillorum		13.68	0.15		99	99	
47510500	**Low discrimination**						**47510500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		65.08	0.43	GAL 99	99	1	
Fusobacterium mortiferum		26.84	0.42		1	5	
Capnocytophaga ochracea		5.89	0.17		1	90	
47510502	**Low discrimination**						**47510502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		48.22	0.43	GAL 99	99	1	
Capnocytophaga ochracea		39.30	0.34		1	90	
Bifidobacterium adolescentis		8.04	0.32		99	1	
47510510	**Good identification**						**47510510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.41	0.47	GAL 99	99	1	
Capnocytophaga ochracea		0.59	0.01		1	90	
47510512	**Low discrimination**						**47510512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		82.69	0.47	GAL 99	99	1	
Bifidobacterium adolescentis		9.90	0.33		99	1	
Capnocytophaga ochracea		4.40	0.17		1	90	
47510520	**Low discrimination**						**47510520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		68.40	0.54	GAL 99	99	1	
Capnocytophaga ochracea		13.94	0.34		1	90	
Prevotella melaninogenica		11.95	0.29		1	1	
47510522	**Low discrimination**						**47510522**
					GRA	CAT	
Capnocytophaga ochracea		62.80	0.50	GAL 99	1	90	
Bifidobacterium longum subsp. longum		34.24	0.54		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		1.14	0.16		1	1	
47510530	**Very good identification**						**47510530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.59	0.58	GAL 99	99	1	
Capnocytophaga ochracea		1.31	0.17		1	90	
Prevotella melaninogenica		0.10	-.06		1	1	
47510532	**Low discrimination**						**47510532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.26	0.58	GAL 99	99	1	
Capnocytophaga ochracea		10.56	0.34		1	90	
Clostridium clostridioforme		0.80	0.21		50	33	
47510712	**Good identification**						**47510712**
Clostridium clostridioforme		96.36	0.36	SAL 88			
Bifidobacterium longum subsp. longum		3.65	0.13	TRE 88			
47510720	**Good identification**						**47510720**
					GRA	CAT	
Bacteroides vulgatus		98.07	0.37	GAL 99	1	3	
Bifidobacterium longum subsp. longum		1.40	0.20	XYL 98	99	1	
Capnocytophaga ochracea		0.29	-.01		1	90	
47510732	**Low discrimination**						**47510732**
					GRA		
Clostridium clostridioforme		73.48	0.36	SAL 88	50		
Parabacteroides distasonis		14.07	0.29	TRE 88	1		
Bifidobacterium longum subsp. longum		11.12	0.24	XYL 91	99		
47511000	**Low discrimination**						**47511000**
					GRA	CAT	
Fusobacterium mortiferum		69.00	0.73	MLT 11	1	5	
Prevotella intermedia		12.37	0.49		1	1	
Propionibacterium granulosum		5.94	0.52		99	90	
47511001	**Low discrimination**						**47511001**
					GRA	SPO	
Actinomyces naeslundii		44.91	0.56	ESL 12	99	1	
Clostridium chauvoei		25.07	0.29		99	99	
Fusobacterium mortiferum		21.75	0.39		1	1	
47511002	**Low discrimination**						**47511002**
					CAT		
Capnocytophaga ochracea		31.08	0.47	LAC 99	90		
Prevotella oralis		23.01	0.44	NAG 90	1		
Fusobacterium mortiferum		19.63	0.57		5		
47511003	**Low discrimination**						**47511003**
					SPO		
Actinomyces naeslundii		68.83	0.56	ESL 12	1		
Actinomyces viscosus		7.69	0.16		1		
Clostridium chauvoei		6.78	0.16		99		
47511010	**Low discrimination**						**47511010**
					CAT	GRA	
Eubacterium saburreum		43.40	0.43	IND 99	1	85	
Propionibacterium granulosum		23.28	0.37		90	99	
Fusobacterium mortiferum		20.04	0.39		5	1	
47511020	**Low discrimination**						**47511020**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		73.87	0.63	LAC 99	1	1	
Clostridium perfringens		15.18	0.58		99	1	
Capnocytophaga ochracea		4.93	0.47		1	90	
47511022	**Low discrimination**						**47511022**
					CAT		
Prevotella oralis		49.64	0.66	LAC 94	1		
Capnocytophaga ochracea		31.77	0.63		90		
Prevotella melaninogenica		10.07	0.51		1		
47511032	**Low discrimination**						**47511032**
					CAT		
Capnocytophaga ochracea		81.31	0.47	LAC 99	90		
Prevotella oralis		11.55	0.32	MLZ 10	1		
Prevotella melaninogenica		2.34	0.16		1		
47511040	**Low discrimination**						**47511040**
					GRA		
Fusobacterium mortiferum		46.42	0.39	MLT 11	1		
Clostridium perfringens		14.66	0.20	SOR 1	99		
Actinomyces naeslundii		11.28	0.40		99		
47511041	**Good identification**						**47511041**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47511043	**Good identification**						**47511043**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47511060	**Low discrimination**						**47511060**
					GRA	CAT	
Clostridium perfringens		67.38	0.46	LAC 88	99	1	
Prevotella melaninogenica		17.39	0.29		1	1	
Capnocytophaga ochracea		12.77	0.30		1	90	
47511061	**Low discrimination**						**47511061**
					GRA		
Actinomyces naeslundii		73.74	0.43	ESL 12	99		
Clostridium perfringens		20.87	0.12		99		
Prevotella melaninogenica		5.39	-.06		1		
47511062	**Low discrimination**						**47511062**
					GRA	CAT	
Capnocytophaga ochracea		68.13	0.47	LAC 99	1	90	
Clostridium perfringens		18.80	0.40	SOR 10	99	1	
Prevotella oralis		9.68	0.32		1	1	
47511063	**Low discrimination**						**47511063**
					GRA	CAT	
Actinomyces naeslundii		59.36	0.43	ESL 12	99	25	
Capnocytophaga ochracea		28.66	0.12		1	90	
Clostridium perfringens		7.91	0.06		99	1	
47511100	**Low discrimination**						**47511100**
					CAT		
Prevotella melaninogenica		34.60	0.63	NAG 99	1		
Fusobacterium mortiferum		30.62	0.80		5		
Capnocytophaga ochracea		25.40	0.64		90		
47511101	**Low discrimination**						**47511101**
					GRA	SPO	
Clostridium chauvoei		89.47	0.63	RAF 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		4.61	0.64		99	1	
Prevotella melaninogenica		2.07	0.29		1	1	
47511102	**Low discrimination**						**47511102**
					GRA	CAT	
Capnocytophaga ochracea		76.15	0.81	NAG 90	1	90	
Bifidobacterium longum subsp. infantis		9.65	0.62		99	1	
Prevotella oralis		8.92	0.64		1	1	
47511103	**Low discrimination**						**47511103**
					GRA	SPO	
Clostridium chauvoei		40.91	0.50	RAF 1	99	99	
Capnocytophaga ochracea		35.39	0.47		1	1	
Actinomyces naeslundii		11.94	0.64		99	1	
47511110	**Low discrimination**						**47511110**
					CAT	GRA	
Capnocytophaga ochracea		55.10	0.48	MLZ 10	90	1	
Bifidobacterium longum subsp. longum		16.62	0.47	NAG 90	1	99	
Eubacterium saburreum		13.08	0.49	BGL 90	1	85	
47511112	**Low discrimination**						**47511112**
					GRA	CAT	
Capnocytophaga ochracea		88.58	0.64	MLZ 10	1	90	
Bifidobacterium adolescentis		5.26	0.53	NAG 90	99	1	
Bifidobacterium longum subsp. longum		2.97	0.47		99	1	
47511120	**Good identification**						**47511120**
					GRA	CAT	
Prevotella melaninogenica		91.38	0.97		1	1	
Capnocytophaga ochracea		6.10	0.81		1	90	
Clostridium perfringens		1.39	0.73		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47511121	**Low discrimination**							**47511121**
					GRA	CAT	SPO	
Prevotella melaninogenica		79.22	0.63	URE 1	1	1	1	
Clostridium chauvoei		11.55	0.55		99	1	99	
Capnocytophaga ochracea		5.29	0.47		1	90	1	
47511122	**Low discrimination**							**47511122**
					CAT			
Capnocytophaga ochracea		60.33	0.97		90			
Prevotella melaninogenica		19.13	0.85		1			
Prevotella oralis		14.92	0.86		1			
47511123	**Low discrimination**							**47511123**
					CAT			
Capnocytophaga ochracea		57.56	0.63	URE 1	90			
Prevotella melaninogenica		18.25	0.51		1			
Prevotella oralis		14.23	0.52		1			
47511130	**Low discrimination**							**47511130**
					GRA	CAT		
Prevotella melaninogenica		53.64	0.63	MLZ 1	1	1		
Capnocytophaga ochracea		39.38	0.64		1	90		
Bifidobacterium longum subsp. longum		5.28	0.58		99	1		
47511132	**Good identification**							**47511132**
					CAT			
Capnocytophaga ochracea		92.15	0.81	MLZ 10	90			
Prevotella melaninogenica		2.66	0.51		1			
Prevotella oralis		2.07	0.52		1			
47511133	**Acceptable identification**							**47511133**
					CAT			
Capnocytophaga ochracea		90.49	0.47	MLZ 10	90			
Prevotella melaninogenica		2.61	0.16	URE 1	1			
Prevotella oralis		2.03	0.18		1			
47511140	**Low discrimination**							**47511140**
					CAT			
Capnocytophaga ochracea		71.38	0.48	NAG 90	90			
Prevotella melaninogenica		8.84	0.29	SOR 10	1			
Fusobacterium mortiferum		7.82	0.45	BGL 90	5			
47511142	**Very good identification**							**47511142**
					GRA	CAT		
Capnocytophaga ochracea		95.74	0.64	NAG 90	1	90		
Bifidobacterium longum subsp. infantis		1.10	0.27	SOR 10	99	1		
Prevotella oralis		1.02	0.30		1	1		
47511152	**Acceptable identification**							**47511152**
					GRA	CAT		
Capnocytophaga ochracea		92.05	0.48	MLZ 10	1	90		
Bifidobacterium adolescentis		6.08	0.38	NAG 90	99	1		
Bifidobacterium breve		1.34	0.27	SOR 10	99	1		
47511160	**Low discrimination**							**47511160**
					GRA	CAT		
Prevotella melaninogenica		49.11	0.63	SOR 1	1	1		
Capnocytophaga ochracea		36.06	0.64		1	90		
Clostridium perfringens		14.09	0.60		99	1		
47511162	**Good identification**							**47511162**
					CAT			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		92.61	0.81	SOR 10	90		
Prevotella melaninogenica		2.67	0.51		1		
Prevotella oralis		2.08	0.52		1		
47511163	**Low discrimination**						**47511163**
					GRA	CAT	
Capnocytophaga ochracea		87.55	0.47	SOR 10	1	90	
Actinomyces naeslundii		5.21	0.51	URE 1	99	25	
Prevotella melaninogenica		2.52	0.16		1	1	
47511170	**Low discrimination**						**47511170**
					GRA	CAT	
Capnocytophaga ochracea		85.05	0.48	MLZ 10	1	90	
Prevotella melaninogenica		10.53	0.29	SOR 10	1	1	
Clostridium perfringens		3.02	0.26	BGL 90	99	1	
47511172	**Very good identification**						**47511172**
					GRA	CAT	
Capnocytophaga ochracea		98.76	0.64	MLZ 10	1	90	
Bifidobacterium breve		0.28	0.32	SOR 10	99	1	
Prevotella melaninogenica		0.26	0.16		1	1	
47511200	**Low discrimination**						**47511200**
					GRA	CAT	
Fusobacterium mortiferum		54.10	0.39	MLT 11	1	5	
Prevotella oralis		15.73	0.16	RHA 1	1	1	
Propionibacterium granulosum		14.26	0.26		99	90	
47511202	**Acceptable identification**						**47511202**
					CAT		
Prevotella oralis		92.68	0.38	LAC 94	1		
Capnocytophaga ochracea		2.95	0.12	NAG 95	90		
Fusobacterium mortiferum		1.86	0.23		5		
47511220	**Low discrimination**						**47511220**
Prevotella oralis		56.68	0.38	LAC 94			
Prevotella melaninogenica		27.04	0.29	BGL 95			
Bacteroides vulgatus		8.25	0.18				
47511222	**Very good identification**						**47511222**
					CAT		
Prevotella oralis		96.81	0.60	LAC 94	1		
Capnocytophaga ochracea		1.46	0.29		90		
Parabacteroides distasonis		0.92	0.38		76		
47511300	**Low discrimination**						**47511300**
Prevotella oralis		33.57	0.36	NAG 95			
Prevotella melaninogenica		19.43	0.29	BGL 95			
Fusobacterium mortiferum		17.19	0.45				
47511302	**Low discrimination**						**47511302**
					GRA	CAT	
Prevotella oralis		79.74	0.58	NAG 95	1	1	
Capnocytophaga ochracea		16.05	0.47		1	90	
Bifidobacterium longum subsp. infantis		2.03	0.27		99	1	
47511320	**Low discrimination**						**47511320**
Prevotella melaninogenica		57.90	0.63	RHA 1			
Prevotella oralis		19.20	0.58				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		17.66	0.52				
47511322	**Low discrimination**						**47511322**
					CAT		
Prevotella oralis		83.71	0.80		1		
Capnocytophaga ochracea		7.98	0.63		90		
Parabacteroides distasonis		5.01	0.72		76		
47511323	**Low discrimination**						**47511323**
					CAT		
Prevotella oralis		83.30	0.46	URE 1	1		
Capnocytophaga ochracea		7.94	0.29		90		
Parabacteroides distasonis		4.99	0.38		76		
47511332	**Low discrimination**						**47511332**
					CAT		
Parabacteroides distasonis		60.08	0.67	XYL 90	76		
Capnocytophaga ochracea		19.73	0.47		90		
Prevotella oralis		18.82	0.46		1		
47511362	**Low discrimination**						**47511362**
					CAT		
Capnocytophaga ochracea		48.38	0.47	RHA 1	90		
Prevotella oralis		46.14	0.46	SOR 10	1		
Parabacteroides distasonis		2.76	0.38		76		
47511400	**Low discrimination**						**47511400**
					GRA		
Lactobacillus fermentum		52.47	0.54	ESL 1	99		
Bifidobacterium longum subsp. longum		32.48	0.43		99		
Fusobacterium mortiferum		8.61	0.39		1		
47511402	**Low discrimination**						**47511402**
					GRA	CAT	
Lactobacillus fermentum		43.91	0.54	ESL 1	99	1	
Bifidobacterium longum subsp. longum		27.18	0.43		99	1	
Capnocytophaga ochracea		22.15	0.34		1	90	
47511410	**Good identification**						**47511410**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.47	0.47	LAC 99	99	1	
Lactobacillus fermentum		0.94	0.20		99	1	
Capnocytophaga ochracea		0.58	0.01		1	90	
47511412	**Low discrimination**						**47511412**
					GRA	CAT	
Bifidobacterium longum subsp. longum		82.30	0.47	LAC 99	99	1	
Bifidobacterium adolescentis		9.85	0.33		99	1	
Capnocytophaga ochracea		4.38	0.17		1	90	
47511420	**Low discrimination**						**47511420**
					GRA	CAT	
Bifidobacterium longum subsp. longum		61.11	0.54	LAC 99	99	1	
Capnocytophaga ochracea		12.45	0.34		1	90	
Prevotella melaninogenica		10.68	0.29		1	1	
47511422	**Low discrimination**						**47511422**
					GRA	CAT	
Capnocytophaga ochracea		57.18	0.50	LAC 99	1	90	
Bifidobacterium longum subsp. longum		31.18	0.54		99	1	
Prevotella oralis		5.11	0.32		1	1	
47511430	**Very good identification**						**47511430**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Bifidobacterium longum subsp. longum		98.59	0.58	LAC 99	99	1		
Capnocytophaga ochracea		1.31	0.17		1	90		
Prevotella melaninogenica		0.10	-.06		1	1		
47511432	**Low discrimination**							**47511432**
					GRA	CAT		
Bifidobacterium longum subsp. longum		88.26	0.58	LAC 99	99	1		
Capnocytophaga ochracea		10.56	0.34		1	90		
Clostridium clostridioforme		0.72	0.20		50	33		
47511500	**Low discrimination**							**47511500**
					GRA	CAT		
Bifidobacterium longum subsp. longum		87.96	0.78		99	1		
Capnocytophaga ochracea		7.97	0.51		1	90		
Lactobacillus fermentum		2.34	0.58		99	1		
47511501	**Low discrimination**							**47511501**
					GRA	SPO	CAT	
Bifidobacterium longum subsp. longum		68.88	0.43	URE 1	99	1	1	
Clostridium chauvoei		21.05	0.29		99	99	1	
Capnocytophaga ochracea		6.24	0.17		1	1	90	
47511502	**Low discrimination**							**47511502**
					GRA	CAT		
Bifidobacterium longum subsp. longum		49.42	0.78		99	1		
Capnocytophaga ochracea		40.28	0.68		1	90		
Bifidobacterium adolescentis		8.24	0.66		99	1		
47511503	**Low discrimination**							**47511503**
					GRA	CAT		
Bifidobacterium longum subsp. longum		47.98	0.43	URE 1	99	1		
Capnocytophaga ochracea		39.11	0.34		1	90		
Bifidobacterium adolescentis		8.00	0.32		99	1		
47511510	**Very good identification**							**47511510**
					GRA	CAT		
Bifidobacterium longum subsp. longum		99.26	0.82		99	1		
Capnocytophaga ochracea		0.59	0.35		1	90		
Bifidobacterium adolescentis		0.12	0.33		99	1		
47511511	**Good identification**							**47511511**
					GRA	CAT		
Bifidobacterium longum subsp. longum		99.41	0.47	URE 1	99	1		
Capnocytophaga ochracea		0.59	0.01		1	90		
47511512	**Low discrimination**							**47511512**
					GRA	CAT		
Bifidobacterium longum subsp. longum		85.15	0.82		99	1		
Bifidobacterium adolescentis		10.20	0.67		99	1		
Capnocytophaga ochracea		4.53	0.51		1	90		
47511513	**Low discrimination**							**47511513**
					GRA	CAT		
Bifidobacterium longum subsp. longum		85.26	0.47	URE 1	99	1		
Bifidobacterium adolescentis		10.21	0.33		99	1		
Capnocytophaga ochracea		4.53	0.17		1	90		
47511520	**Low discrimination**							**47511520**
					GRA	CAT		
Bifidobacterium longum subsp. longum		68.30	0.88		99	1		
Capnocytophaga ochracea		13.92	0.68		1	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		11.93	0.63		1	1	
47511521	**Low discrimination**						**47511521**
					GRA	CAT	
Bifidobacterium longum subsp. longum		67.05	0.54	URE 1	99	1	
Capnocytophaga ochracea		13.66	0.34		1	90	
Prevotella melaninogenica		11.72	0.29		1	1	
47511522	**Low discrimination**						**47511522**
					GRA	CAT	
Capnocytophaga ochracea		62.81	0.84		1	90	
Bifidobacterium longum subsp. longum		34.25	0.88		99	1	
Prevotella melaninogenica		1.14	0.51		1	1	
47511523	**Low discrimination**						**47511523**
					GRA	CAT	
Capnocytophaga ochracea		62.69	0.50	URE 1	1	90	
Bifidobacterium longum subsp. longum		34.18	0.54		99	1	
Prevotella melaninogenica		1.14	0.16		1	1	
47511530	**Very good identification**						**47511530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.54	0.92		99	1	
Capnocytophaga ochracea		1.31	0.51		1	90	
Prevotella melaninogenica		0.10	0.29		1	1	
47511531	**Very good identification**						**47511531**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.59	0.58	URE 1	99	1	
Capnocytophaga ochracea		1.31	0.17		1	90	
Prevotella melaninogenica		0.10	-.06		1	1	
47511532	**Low discrimination**						**47511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.90	0.92		99	1	
Capnocytophaga ochracea		10.64	0.68		1	90	
Parabacteroides distasonis		0.38	0.55		1	76	
47511533	**Low discrimination**						**47511533**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.97	0.58	URE 1	99	1	
Capnocytophaga ochracea		10.65	0.34		1	90	
Parabacteroides distasonis		0.38	0.21		1	76	
47511540	**Low discrimination**						**47511540**
					GRA	CAT	
Bifidobacterium longum subsp. longum		48.48	0.43	SOR 1	99	1	
Capnocytophaga ochracea		48.29	0.35		1	90	
Lactobacillus fermentum		1.29	0.24		99	1	
47511542	**Low discrimination**						**47511542**
					GRA	CAT	
Capnocytophaga ochracea		74.43	0.51	NAG 90	1	90	
Bifidobacterium adolescentis		16.95	0.50	SOR 10	99	1	
Bifidobacterium longum subsp. longum		8.30	0.43		99	1	
47511550	**Acceptable identification**						**47511550**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.61	0.47	SOR 1	99	1	
Capnocytophaga ochracea		6.02	0.19		1	90	
Bifidobacterium adolescentis		1.37	0.18		99	1	
47511552	**Low discrimination**						**47511552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Bifidobacterium adolescentis		48.03	0.52	XYL 87	99	1	
Bifidobacterium longum subsp. longum		32.79	0.47	SOR 11	99	1	
Capnocytophaga ochracea		19.18	0.35		1	90	
47511560	**Low discrimination**						**47511560**
					GRA	CAT	
Capnocytophaga ochracea		63.19	0.51	SOR 10	1	90	
Bifidobacterium longum subsp. longum		28.19	0.54	BGL 90	99	1	
Prevotella melaninogenica		4.93	0.29		1	1	
47511562	**Good identification**						**47511562**
					GRA	CAT	
Capnocytophaga ochracea		94.78	0.68	SOR 10	1	90	
Bifidobacterium longum subsp. longum		4.70	0.54		99	1	
Prevotella melaninogenica		0.16	0.16		1	1	
47511570	**Low discrimination**						**47511570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.16	0.58	SOR 1	99	1	
Capnocytophaga ochracea		12.75	0.35		1	90	
Prevotella melaninogenica		0.09	-.06		1	1	
47511572	**Low discrimination**						**47511572**
					GRA	CAT	
Capnocytophaga ochracea		56.63	0.51	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		43.02	0.58	SOR 10	99	1	
Parabacteroides distasonis		0.18	0.21		1	76	
47511612	**Good identification**						**47511612**
Clostridium clostridioforme		95.95	0.35	SAL 88			
Bifidobacterium longum subsp. longum		4.05	0.13	TRE 88			
47511620	**Good identification**						**47511620**
					GRA		
Bacteroides vulgatus		97.57	0.37	XYL 98	1		
Bifidobacterium longum subsp. longum		1.39	0.20	LAC 99	99		
Prevotella oralis		0.51	0.04		1		
47511632	**Low discrimination**						**47511632**
					GRA		
Clostridium clostridioforme		71.29	0.35	SAL 88	50		
Parabacteroides distasonis		15.24	0.29	TRE 88	1		
Bifidobacterium longum subsp. longum		12.04	0.24	XYL 91	99		
47511700	**Low discrimination**						**47511700**
					GRA		
Bacteroides vulgatus		68.55	0.37	XYL 98	1		
Bifidobacterium longum subsp. longum		24.20	0.43	NAG 99	99		
Eubacterium contortum		3.80	0.23		99		
47511702	**Low discrimination**						**47511702**
					GRA	CAT	
Bifidobacterium longum subsp. longum		36.19	0.43	RHA 1	99	1	
Capnocytophaga ochracea		29.49	0.34		1	90	
Clostridium clostridioforme		10.83	0.33		50	33	
47511710	**Good identification**						**47511710**
					GRA		
Bifidobacterium longum subsp. longum		97.01	0.47	RHA 1	99		
Bacteroides vulgatus		1.63	0.03		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.79	0.10		50		
47511712	**Low discrimination**						**47511712**
Bifidobacterium longum subsp. longum		49.38	0.47	RHA 1			
Clostridium clostridioforme		39.55	0.44				
Bifidobacterium adolescentis		5.91	0.33				
47511720	**Very good identification**						**47511720**
					GRA	CAT	
Bacteroides vulgatus		97.92	0.71	XYL 98	1	3	
Bifidobacterium longum subsp. longum		1.40	0.54		99	1	
Capnocytophaga ochracea		0.29	0.34		1	90	
47511721	**Good identification**						**47511721**
					GRA	CAT	
Bacteroides vulgatus		98.07	0.37	XYL 98	1	3	
Bifidobacterium longum subsp. longum		1.40	0.20	URE 1	99	1	
Capnocytophaga ochracea		0.29	-.01		1	90	
47511722	**Low discrimination**						**47511722**
					CAT		
Parabacteroides distasonis		45.47	0.68	XYL 90	76		
Capnocytophaga ochracea		20.84	0.50		90		
Prevotella oralis		12.51	0.46		1		
47511730	**Low discrimination**						**47511730**
					GRA	CAT	
Bifidobacterium longum subsp. longum		69.20	0.58	RHA 1	99	1	
Bacteroides vulgatus		28.78	0.37		1	3	
Capnocytophaga ochracea		0.92	0.17		1	90	
47511732	**Low discrimination**						**47511732**
					GRA		
Parabacteroides distasonis		48.59	0.64	XYL 90	1		
Bifidobacterium longum subsp. longum		38.39	0.58		99		
Clostridium clostridioforme		7.69	0.44		50		
47511760	**Good identification**						**47511760**
					GRA	CAT	
Bacteroides vulgatus		95.35	0.37	XYL 98	1	3	
Capnocytophaga ochracea		3.05	0.17	SOR 1	1	90	
Bifidobacterium longum subsp. longum		1.36	0.20		99	1	
47512000	**Low discrimination**						**47512000**
					GRA	CAT	
Propionibacterium granulosum		86.57	0.50	ESL 1	99	90	
Fusobacterium mortiferum		8.98	0.36		1	5	
Prevotella intermedia		2.42	0.15		1	1	
47512001	**Low discrimination**						**47512001**
Actinomyces naeslundii		72.01	0.37	ESL 12			
Propionibacterium granulosum		27.99	0.16	GAL 96			
47512002	**Low discrimination**						**47512002**
					GRA	CAT	
Propionibacterium granulosum		88.55	0.39	ESL 1	99	90	
Actinomyces naeslundii		6.14	0.34		99	25	
Fusobacterium mortiferum		4.08	0.20		1	5	
47512003	**Good identification**						**47512003**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	ESL 12			
47512020	**Acceptable identification**						**47512020**
					GRA	CAT	
Propionibacterium granulosum		92.09	0.39	ESL 1	99	90	
Actinomyces naeslundii		6.38	0.34		99	25	
Prevotella melaninogenica		1.53	-.06		1	1	
47512021	**Good identification**						**47512021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47512023	**Good identification**						**47512023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47512100	**Low discrimination**						**47512100**
					GRA		
Fusobacterium mortiferum		45.25	0.42	MLT 11	1		
Propionibacterium propionicum		42.86	0.43	MAN 1	99		
Actinomyces naeslundii		9.22	0.42		99		
47512101	**Low discrimination**						**47512101**
Actinomyces naeslundii		74.39	0.45	ESL 12			
Propionibacterium propionicum		25.61	0.27	GAL 96			
47512102	**Low discrimination**						**47512102**
					GRA	CAT	
Propionibacterium propionicum		74.29	0.46	NIT 86	99	1	
Actinomyces naeslundii		10.65	0.42	ESL 1	99	25	
Capnocytophaga ochracea		5.92	0.12		1	90	
47512103	**Low discrimination**						**47512103**
Actinomyces naeslundii		65.94	0.45	ESL 12			
Propionibacterium propionicum		34.06	0.30	GAL 96			
47512121	**Acceptable identification**						**47512121**
					GRA		
Actinomyces naeslundii		94.70	0.45	ESL 12	99		
Prevotella melaninogenica		5.30	-.06	GAL 96	1		
47512123	**Good identification**						**47512123**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47512140	**Good identification**						**47512140**
Propionibacterium propionicum		96.34	0.43	NIT 86			
Actinomyces naeslundii		3.66	0.29	ESL 1			
47512142	**Good identification**						**47512142**
Propionibacterium propionicum		97.53	0.46	NIT 86			
Actinomyces naeslundii		2.47	0.29	ESL 1			
47513000	**Low discrimination**						**47513000**
					GRA	CAT	
Propionibacterium granulosum		61.44	0.50	ESL 1	99	90	
Actinomyces naeslundii		25.55	0.58		99	25	
Fusobacterium mortiferum		9.56	0.39		1	5	
47513001	**Very good identification**						**47513001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		98.41	0.61	ESL 12			
Propionibacterium granulosum		1.59	0.16				
47513002	**Low discrimination**						**47513002**
					GRA		
Actinomyces naeslundii		53.67	0.58	ESL 12	99		
Propionibacterium granulosum		32.27	0.39		99		
Capnocytophaga ochracea		3.53	0.12		1		
47513003	**Very good identification**						**47513003**
Actinomyces naeslundii		100.0	0.61	ESL 12			
47513011	**Good identification**						**47513011**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47513013	**Good identification**						**47513013**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47513020	**Low discrimination**						**47513020**
					GRA	CAT	
Actinomyces naeslundii		34.43	0.58	ESL 12	99	25	
Prevotella melaninogenica		33.96	0.29		1	1	
Propionibacterium granulosum		20.70	0.39		99	90	
47513021	**Very good identification**						**47513021**
					GRA		
Actinomyces naeslundii		99.34	0.61	ESL 12	99		
Prevotella melaninogenica		0.66	-.06		1		
47513022	**Low discrimination**						**47513022**
					GRA	CAT	
Actinomyces naeslundii		33.36	0.58	ESL 12	99	25	
Prevotella oralis		30.88	0.32		1	1	
Capnocytophaga ochracea		19.77	0.29		1	90	
47513023	**Very good identification**						**47513023**
					GRA		
Actinomyces naeslundii		99.38	0.61	ESL 12	99		
Prevotella oralis		0.62	-.03		1		
47513031	**Good identification**						**47513031**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47513033	**Good identification**						**47513033**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47513040	**Low discrimination**						**47513040**
					CAT		
Actinomyces naeslundii		83.86	0.45	ESL 12	25		
Propionibacterium granulosum		11.55	0.16		90		
Actinomyces israelii		4.59	0.14		1		
47513041	**Good identification**						**47513041**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47513042	**Low discrimination**						**47513042**
Actinomyces naeslundii		88.54	0.45	ESL 12			
Actinomyces israelii		7.27	0.17				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		4.19	0.12				
47513043	**Good identification**						**47513043**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47513060	**Low discrimination**						**47513060**
					GRA		
Actinomyces naeslundii		78.42	0.45	ESL 12	99		
Clostridium perfringens		17.15	0.12		99		
Prevotella melaninogenica		4.43	-.06		1		
47513061	**Good identification**						**47513061**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47513062	**Low discrimination**						**47513062**
					GRA	CAT	
Actinomyces naeslundii		65.40	0.45	ESL 12	99	25	
Capnocytophaga ochracea		24.40	0.12		1	90	
Clostridium perfringens		6.73	0.06		99	1	
47513063	**Good identification**						**47513063**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47513100	**Low discrimination**						**47513100**
					GRA		
Actinomyces naeslundii		44.56	0.66	ESL 12	99		
Prevotella melaninogenica		15.44	0.29		1		
Fusobacterium mortiferum		13.67	0.45		1		
47513101	**Low discrimination**						**47513101**
					SPO		
Actinomyces naeslundii		89.30	0.69	ESL 12	1		
Clostridium chauvoei		9.02	0.29		99		
Propionibacterium propionicum		1.28	0.27		1		
47513102	**Low discrimination**						**47513102**
					GRA	CAT	
Capnocytophaga ochracea		42.65	0.47	MAN 1	1	90	
Bifidobacterium adolescentis		18.64	0.50	NAG 90	99	1	
Actinomyces naeslundii		18.62	0.66		99	25	
47513103	**Good identification**						**47513103**
					SPO		
Actinomyces naeslundii		94.03	0.69	ESL 12	1		
Propionibacterium propionicum		2.02	0.30		1		
Clostridium chauvoei		1.68	0.16		99		
47513110	**Low discrimination**						**47513110**
					GRA	CAT	
Actinomyces naeslundii		42.32	0.44	ESL 12	99	25	
Capnocytophaga ochracea		22.74	0.14	MLZ 5	1	90	
Bifidobacterium adolescentis		9.94	0.18		99	1	
47513111	**Good identification**						**47513111**
Actinomyces naeslundii		100.0	0.47	ESL 12			
47513112	**Low discrimination**						**47513112**
					GRA	CAT	
Bifidobacterium adolescentis		76.78	0.52	XYL 87	99	1	
Capnocytophaga ochracea		15.97	0.30	ARA 87	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		3.30	0.44		99	25	
47513113	**Low discrimination**						**47513113**
Actinomyces naeslundii		86.47	0.47	ESL 12			
Bifidobacterium adolescentis		13.54	0.18	MLZ 5			
47513120	**Low discrimination**						**47513120**
					GRA	CAT	
Prevotella melaninogenica		89.01	0.63	MAN 1	1	1	
Capnocytophaga ochracea		5.94	0.47		1	90	
Actinomyces naeslundii		2.59	0.66		99	25	
47513121	**Low discrimination**						**47513121**
					GRA	SPO	
Actinomyces naeslundii		77.90	0.69	ESL 12	99	1	
Prevotella melaninogenica		18.00	0.29		1	1	
Clostridium chauvoei		2.62	0.21		99	99	
47513122	**Low discrimination**						**47513122**
					CAT		
Capnocytophaga ochracea		58.46	0.63	MAN 1	90		
Prevotella melaninogenica		18.54	0.51		1		
Prevotella oralis		14.45	0.52		1		
47513123	**Low discrimination**						**47513123**
					GRA	CAT	
Actinomyces naeslundii		81.05	0.69	ESL 12	99	25	
Capnocytophaga ochracea		11.25	0.29		1	90	
Prevotella melaninogenica		3.57	0.16		1	1	
47513131	**Good identification**						**47513131**
					GRA		
Actinomyces naeslundii		95.75	0.47	ESL 12	99		
Prevotella melaninogenica		4.25	-.06	MLZ 5	1		
47513132	**Low discrimination**						**47513132**
					GRA	CAT	
Capnocytophaga ochracea		87.45	0.47	MAN 1	1	90	
Bifidobacterium breve		2.77	0.32	MLZ 10	99	1	
Prevotella melaninogenica		2.52	0.16		1	1	
47513133	**Low discrimination**						**47513133**
					GRA		
Actinomyces naeslundii		77.34	0.47	ESL 12	99		
Capnocytophaga ochracea		22.66	0.12	MLZ 5	1		
47513140	**Low discrimination**						**47513140**
					GRA	CAT	
Propionibacterium propionicum		45.50	0.43	NIT 86	99	1	
Actinomyces naeslundii		41.45	0.53	ESL 1	99	25	
Capnocytophaga ochracea		6.64	0.14		1	90	
47513141	**Acceptable identification**						**47513141**
Actinomyces naeslundii		92.48	0.56	ESL 12			
Propionibacterium propionicum		7.52	0.27				
47513142	**Low discrimination**						**47513142**
					GRA	CAT	
Propionibacterium propionicum		31.95	0.46	NIT 86	99	1	
Capnocytophaga ochracea		27.99	0.30	ESL 1	1	90	
Actinomyces naeslundii		19.41	0.53		99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47513143	**Low discrimination**						**47513143**
Actinomyces naeslundii		89.13	0.56	ESL 12			
Propionibacterium propionicum		10.87	0.30				
47513152	**Low discrimination**						**47513152**
					GRA	CAT	
Bifidobacterium adolescentis		64.20	0.36	XYL 87	99	1	
Bifidobacterium breve		17.31	0.27	ARA 87	99	1	
Capnocytophaga ochracea		12.00	0.14	SOR 11	1	90	
47513161	**Good identification**						**47513161**
					GRA		
Actinomyces naeslundii		98.70	0.56	ESL 12	99		
Prevotella melaninogenica		1.31	-.06		1		
47513162	**Low discrimination**						**47513162**
					GRA	CAT	
Capnocytophaga ochracea		84.12	0.47	MAN 1	1	90	
Actinomyces naeslundii		6.48	0.53	SOR 10	99	25	
Bifidobacterium breve		2.66	0.32		99	1	
47513163	**Acceptable identification**						**47513163**
					GRA		
Actinomyces naeslundii		91.96	0.56	ESL 12	99		
Capnocytophaga ochracea		8.04	0.12		1		
47513322	**Low discrimination**						**47513322**
					CAT		
Prevotella oralis		82.89	0.46	MAN 1	1		
Capnocytophaga ochracea		7.90	0.29		90		
Parabacteroides distasonis		4.96	0.38		76		
47513500	**Low discrimination**						**47513500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.35	0.43	MAN 1	99	1	
Bifidobacterium adolescentis		10.15	0.30		99	1	
Capnocytophaga ochracea		6.73	0.17		1	90	
47513502	**Low discrimination**						**47513502**
					GRA	CAT	
Bifidobacterium adolescentis		87.33	0.64	XYL 87	99	1	
Bifidobacterium longum subsp. longum		6.46	0.43		99	1	
Capnocytophaga ochracea		5.27	0.34		1	90	
47513510	**Acceptable identification**						**47513510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.18	0.47	MAN 1	99	1	
Bifidobacterium adolescentis		8.83	0.32		99	1	
Capnocytophaga ochracea		0.53	0.01		1	90	
47513512	**Good identification**						**47513512**
					GRA	CAT	
Bifidobacterium adolescentis		90.14	0.66	XYL 87	99	1	
Bifidobacterium longum subsp. longum		9.29	0.47		99	1	
Capnocytophaga ochracea		0.49	0.17		1	90	
47513520	**Low discrimination**						**47513520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		68.16	0.54	MAN 1	99	1	
Capnocytophaga ochracea		13.89	0.34		1	90	
Prevotella melaninogenica		11.91	0.29		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47513522	**Low discrimination**						**47513522**
					GRA	CAT	
Capnocytophaga ochracea		62.04	0.50	MAN 1	1	90	
Bifidobacterium longum subsp. longum		33.83	0.54		99	1	
Bifidobacterium adolescentis		1.15	0.30		99	1	
47513530	**Very good identification**						**47513530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.59	0.58	MAN 1	99	1	
Capnocytophaga ochracea		1.31	0.17		1	90	
Prevotella melaninogenica		0.10	-.06		1	1	
47513532	**Low discrimination**						**47513532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.08	0.58	MAN 1	99	1	
Capnocytophaga ochracea		10.42	0.34		1	90	
Bifidobacterium adolescentis		2.13	0.32		99	1	
47513542	**Acceptable identification**						**47513542**
					GRA	CAT	
Bifidobacterium adolescentis		92.26	0.49	XYL 87	99	1	
Capnocytophaga ochracea		5.00	0.17	SOR 11	1	90	
Actinomyces israelii		1.86	0.26		99	1	
47513552	**Good identification**						**47513552**
					GRA	CAT	
Bifidobacterium adolescentis		98.68	0.50	XYL 87	99	1	
Bifidobacterium longum subsp. longum		0.83	0.13	SOR 11	99	1	
Capnocytophaga ochracea		0.49	0.01		1	90	
47513720	**Good identification**						**47513720**
					GRA	CAT	
Bacteroides vulgatus		98.07	0.37	MAN 1	1	3	
Bifidobacterium longum subsp. longum		1.40	0.20	XYL 98	99	1	
Capnocytophaga ochracea		0.29	-.01		1	90	
47514000	**Acceptable identification**						**47514000**
					COC	GRA	
Fusobacterium mortiferum		91.44	0.51	MLT 11	1	1	
Prevotella intermedia		3.30	0.15	XYL 7	1	1	
Gemella morbillorum		3.01	0.15		99	99	
47514100	**Low discrimination**						**47514100**
					CAT		
Fusobacterium mortiferum		89.41	0.57	MLT 11	5		
Capnocytophaga ochracea		3.73	0.20	XYL 7	90		
Bacteroides fragilis		3.34	0.10		99		
47514120	**Low discrimination**						**47514120**
					CAT		
Bacteroides fragilis		77.32	0.44	GAL 99	99		
Bacteroides vulgatus		7.89	0.30	BGL 95	3		
Capnocytophaga ochracea		7.85	0.36		90		
47514122	**Good identification**						**47514122**
					CAT		
Bacteroides fragilis		94.09	0.66	GAL 99	99		
Capnocytophaga ochracea		4.52	0.53		90		
Prevotella oralis		0.55	0.37		1		
47514320	**Good identification**						**47514320**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		97.94	0.47	GAL 99	1	3	
Bacteroides fragilis		1.08	0.10	ARA 93	1	99	
Eubacterium contortum		0.32	0.11		99	1	
47514322	**Low discrimination**						**47514322**
					CAT		
Parabacteroides distasonis		44.04	0.54	GAL 99	76		
Bacteroides fragilis		40.49	0.32		99		
Prevotella oralis		10.10	0.30		1		
47514332	**Low discrimination**						**47514332**
Parabacteroides distasonis		79.63	0.49	GAL 99			
Clostridium clostridioforme		18.27	0.33				
Bacteroides fragilis		1.37	-.03				
47514500	**Acceptable identification**						**47514500**
					GRA		
Bifidobacterium longum subsp. longum		92.70	0.52	GAL 99	99		
Bacteroides vulgatus		4.81	0.15		1		
Fusobacterium mortiferum		0.96	0.23		1		
47514502	**Low discrimination**						**47514502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		66.74	0.52	GAL 99	99	1	
Bifidobacterium adolescentis		24.83	0.46		99	1	
Capnocytophaga ochracea		4.53	0.23		1	90	
47514510	**Good identification**						**47514510**
Bifidobacterium longum subsp. longum		99.61	0.56	GAL 99			
Bifidobacterium adolescentis		0.27	0.13				
Clostridium clostridioforme		0.12	0.04				
47514512	**Low discrimination**						**47514512**
Bifidobacterium longum subsp. longum		71.72	0.56	GAL 99			
Bifidobacterium adolescentis		19.16	0.47				
Clostridium clostridioforme		8.80	0.38				
47514520	**Low discrimination**						**47514520**
					GRA	CAT	
Bacteroides vulgatus		55.58	0.50	GAL 99	1	3	
Bifidobacterium longum subsp. longum		43.24	0.62	RHA 90	99	1	
Capnocytophaga ochracea		0.73	0.23		1	90	
47514522	**Low discrimination**						**47514522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		71.31	0.62	GAL 99	99	1	
Bacteroides fragilis		12.97	0.32		1	99	
Capnocytophaga ochracea		10.90	0.40		1	90	
47514530	**Very good identification**						**47514530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.10	0.66	GAL 99	99	1	
Bacteroides vulgatus		0.76	0.15		1	3	
Capnocytophaga ochracea		0.11	0.07		1	90	
47514532	**Good identification**						**47514532**
					GRA		
Bifidobacterium longum subsp. longum		94.77	0.66	GAL 99	99		
Clostridium clostridioforme		2.91	0.38		50		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		1.21	0.38		1	
47514612	**Good identification**					**47514612**
Clostridium clostridioforme		100.0	0.44	SAL 88		
47514632	**Good identification**					**47514632**
Clostridium clostridioforme		99.43	0.44	SAL 88		
Parabacteroides distasonis		0.57	0.12	TRE 88		
47514702	**Acceptable identification**					**47514702**
Clostridium clostridioforme		91.13	0.42	SAL 88		
Bifidobacterium longum subsp. longum		2.74	0.17	TRE 88		
Eubacterium contortum		2.51	0.10			
47514712	**Good identification**					**47514712**
Clostridium clostridioforme		98.43	0.53	SAL 88		
Bifidobacterium longum subsp. longum		1.11	0.21	TRE 88		
Bifidobacterium adolescentis		0.30	0.13			
47514720	**Very good identification**					**47514720**
					GRA	
Bacteroides vulgatus		99.88	0.66	GAL 99	1	
Bifidobacterium longum subsp. longum		0.09	0.28		99	
Eubacterium contortum		0.02	0.10		99	
47514722	**Low discrimination**					**47514722**
Parabacteroides distasonis		35.81	0.50	GAL 99		
Bacteroides vulgatus		34.50	0.32			
Clostridium clostridioforme		24.83	0.42			
47514732	**Low discrimination**					**47514732**
					GRA	
Clostridium clostridioforme		81.87	0.53	SAL 88	50	
Parabacteroides distasonis		13.96	0.46	TRE 88	1	
Bifidobacterium longum subsp. longum		3.68	0.32		99	
47515000	**Low discrimination**					**47515000**
					GRA	
Fusobacterium mortiferum		52.08	0.54	MLT 11	1	
Lactobacillus fermentum		25.01	0.50	XYL 7	99	
Eubacterium saburreum		9.96	0.40		85	
47515010	**Low discrimination**					**47515010**
					GRA	
Eubacterium saburreum		76.42	0.39	IND 99	85	
Bifidobacterium longum subsp. longum		12.75	0.21		99	
Fusobacterium mortiferum		4.93	0.20		1	
47515020	**Low discrimination**					**47515020**
					CAT	
Bacteroides fragilis		73.05	0.44	LAC 99	99	
Bacteroides vulgatus		7.46	0.30	BGL 95	3	
Capnocytophaga ochracea		7.42	0.36		90	
47515022	**Good identification**					**47515022**
					CAT	
Bacteroides fragilis		91.25	0.66	LAC 99	99	
Capnocytophaga ochracea		4.39	0.53		90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		3.39	0.50		1		
47515100	**Low discrimination**						**47515100**
					GRA	CAT	
Capnocytophaga ochracea		30.44	0.54	NAG 90	1	90	
Bacteroides fragilis		27.26	0.44	BGL 90	1	99	
Bifidobacterium longum subsp. infantis		11.24	0.43		99	1	
47515102	**Low discrimination**						**47515102**
					GRA	CAT	
Bacteroides fragilis		53.04	0.66	NAG 99	1	99	
Capnocytophaga ochracea		28.06	0.70		1	90	
Bifidobacterium longum subsp. infantis		11.63	0.60		99	1	
47515110	**Low discrimination**						**47515110**
					CAT	GRA	
Bifidobacterium longum subsp. longum		66.26	0.56	ARA 99	1	99	
Capnocytophaga ochracea		18.31	0.38		90	1	
Eubacterium saburreum		9.36	0.45		1	85	
47515112	**Low discrimination**						**47515112**
					GRA	CAT	
Bifidobacterium adolescentis		48.70	0.67	ARA 87	99	1	
Capnocytophaga ochracea		30.65	0.54		1	90	
Bifidobacterium longum subsp. longum		12.33	0.56		99	1	
47515120	**Low discrimination**						**47515120**
					CAT		
Bacteroides fragilis		77.44	0.78	BGL 95	99		
Bacteroides vulgatus		7.90	0.64		3		
Capnocytophaga ochracea		7.86	0.70		90		
47515121	**Low discrimination**						**47515121**
					CAT		
Bacteroides fragilis		76.87	0.44	URE 1	99		
Bacteroides vulgatus		7.85	0.30	BGL 95	3		
Capnocytophaga ochracea		7.80	0.36		90		
47515122	**Good identification**						**47515122**
					CAT		
Bacteroides fragilis		94.09	1.00		99		
Capnocytophaga ochracea		4.52	0.87		90		
Prevotella oralis		0.55	0.71		1		
47515123	**Good identification**						**47515123**
					CAT		
Bacteroides fragilis		94.08	0.66	URE 1	99		
Capnocytophaga ochracea		4.52	0.53		90		
Prevotella oralis		0.55	0.37		1		
47515130	**Low discrimination**						**47515130**
					GRA	CAT	
Bifidobacterium longum subsp. longum		43.40	0.66	ARA 99	99	1	
Capnocytophaga ochracea		26.98	0.54		1	90	
Bacteroides fragilis		24.16	0.44		1	99	
47515132	**Low discrimination**						**47515132**
Bacteroides fragilis		54.62	0.66	MLZ 1			
Capnocytophaga ochracea		28.89	0.70				
Parabacteroides distasonis		10.66	0.75				
47515140	**Low discrimination**						**47515140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		79.46	0.38	NAG 90	1	90	
Bacteroides fragilis		6.47	0.10	SOR 10	1	99	
Actinomyces israelii		3.13	0.30	BGL 90	99	1	
47515142	**Low discrimination**						**47515142**
					GRA	CAT	
Capnocytophaga ochracea		72.13	0.54	NAG 90	1	90	
Bacteroides fragilis		12.40	0.32	SOR 10	1	99	
Bifidobacterium adolescentis		11.59	0.50		99	1	
47515152	**Low discrimination**						**47515152**
					GRA	CAT	
Bifidobacterium adolescentis		62.49	0.52	ARA 87	99	1	
Capnocytophaga ochracea		35.35	0.38	SOR 11	1	90	
Bifidobacterium longum subsp. longum		1.29	0.21		99	1	
47515160	**Low discrimination**						**47515160**
					CAT		
Capnocytophaga ochracea		48.08	0.54	SOR 10	90		
Bacteroides fragilis		43.06	0.44	BGL 90	99		
Bacteroides vulgatus		4.40	0.30		3		
47515162	**Low discrimination**						**47515162**
					CAT		
Bacteroides fragilis		64.76	0.66	SOR 1	99		
Capnocytophaga ochracea		34.25	0.70		90		
Prevotella oralis		0.38	0.37		1		
47515170	**Low discrimination**						**47515170**
					GRA	CAT	
Capnocytophaga ochracea		80.58	0.38	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		11.79	0.32	SOR 10	99	1	
Bacteroides fragilis		6.56	0.10	BGL 90	1	99	
47515172	**Low discrimination**						**47515172**
Capnocytophaga ochracea		81.61	0.54	MLZ 10			
Bacteroides fragilis		14.03	0.32	SOR 10			
Parabacteroides distasonis		2.74	0.41				
47515220	**Good identification**						**47515220**
					CAT		
Bacteroides vulgatus		96.56	0.47	LAC 99	3		
Prevotella oralis		1.67	0.22	ARA 93	1		
Bacteroides fragilis		1.06	0.10		99		
47515222	**Low discrimination**						**47515222**
					CAT		
Prevotella oralis		41.55	0.44	XYL 11	1		
Parabacteroides distasonis		28.67	0.54	LAC 94	76		
Bacteroides fragilis		26.36	0.32		99		
47515232	**Low discrimination**						**47515232**
Parabacteroides distasonis		79.42	0.49	LAC 99			
Clostridium clostridioforme		16.33	0.32				
Prevotella oralis		2.16	0.10				
47515300	**Low discrimination**						**47515300**
					GRA		
Eubacterium contortum		63.29	0.58	SAL 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		34.77	0.47		1		
Prevotella oralis		0.50	0.21		1		
47515302	**Low discrimination**						**47515302**
					GRA		
Eubacterium contortum		25.99	0.45	SAL 99	99		
Prevotella oralis		21.97	0.43		1		
Parabacteroides distasonis		18.39	0.54		1		
47515312	**Low discrimination**						**47515312**
					GRA		
Parabacteroides distasonis		48.67	0.49	NAG 99	1		
Clostridium clostridioforme		33.50	0.41		50		
Bifidobacterium adolescentis		7.76	0.33		99		
47515320	**Very good identification**						**47515320**
					GRA	CAT	
Bacteroides vulgatus		97.92	0.81	ARA 93	1	3	
Bacteroides fragilis		1.08	0.44		1	99	
Eubacterium contortum		0.32	0.45		99	1	
47515321	**Good identification**						**47515321**
					GRA	CAT	
Bacteroides vulgatus		97.94	0.47	ARA 93	1	3	
Bacteroides fragilis		1.08	0.10	URE 1	1	99	
Eubacterium contortum		0.32	0.11		99	1	
47515322	**Low discrimination**						**47515322**
					CAT		
Parabacteroides distasonis		44.53	0.88		76		
Bacteroides fragilis		40.94	0.66		99		
Prevotella oralis		10.21	0.65		1		
47515323	**Low discrimination**						**47515323**
					CAT		
Parabacteroides distasonis		44.57	0.54	URE 1	76		
Bacteroides fragilis		40.98	0.32		99		
Prevotella oralis		10.22	0.30		1		
47515330	**Low discrimination**						**47515330**
					GRA		
Bacteroides vulgatus		85.26	0.47	ARA 93	1		
Parabacteroides distasonis		10.43	0.49	MLZ 1	1		
Bifidobacterium longum subsp. longum		1.68	0.32		99		
47515332	**Very good identification**						**47515332**
Parabacteroides distasonis		96.13	0.84				
Bacteroides fragilis		1.66	0.32				
Capnocytophaga ochracea		0.88	0.36				
47515333	**Good identification**						**47515333**
Parabacteroides distasonis		96.78	0.49	URE 1			
Bacteroides fragilis		1.67	-.03				
Capnocytophaga ochracea		0.88	0.02				
47515360	**Good identification**						**47515360**
					CAT		
Bacteroides vulgatus		96.88	0.47	ARA 93	3		
Capnocytophaga ochracea		1.19	0.20	SOR 1	90		
Bacteroides fragilis		1.07	0.10		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47515362	**Low discrimination**						**47515362**
Parabacteroides distasonis		37.24	0.54	SOR 1			
Bacteroides fragilis		34.23	0.32				
Capnocytophaga ochracea		18.11	0.36				
47515372	**Low discrimination**						**47515372**
Parabacteroides distasonis		88.93	0.49	SOR 1			
Capnocytophaga ochracea		8.92	0.20				
Bacteroides fragilis		1.53	-.03				
47515400	**Low discrimination**						**47515400**
					GRA		
Bifidobacterium longum subsp. longum		71.64	0.52	LAC 99	99		
Lactobacillus fermentum		22.66	0.50		99		
Bacteroides vulgatus		3.72	0.15		1		
47515402	**Low discrimination**						**47515402**
Bifidobacterium longum subsp. longum		54.97	0.52	LAC 99			
Bifidobacterium adolescentis		20.45	0.46				
Lactobacillus fermentum		17.38	0.50				
47515410	**Good identification**						**47515410**
Bifidobacterium longum subsp. longum		99.55	0.56	LAC 99			
Bifidobacterium adolescentis		0.27	0.13				
Lactobacillus fermentum		0.19	0.16				
47515412	**Low discrimination**						**47515412**
Bifidobacterium longum subsp. longum		72.22	0.56	LAC 99			
Bifidobacterium adolescentis		19.29	0.47				
Clostridium clostridioforme		7.94	0.38				
47515420	**Low discrimination**						**47515420**
					GRA		
Bacteroides vulgatus		54.96	0.50	LAC 99	1		
Bifidobacterium longum subsp. longum		42.76	0.62	RHA 90	99		
Lactobacillus fermentum		1.13	0.42		99		
47515422	**Low discrimination**						**47515422**
					GRA	CAT	
Bifidobacterium longum subsp. longum		69.78	0.62	LAC 99	99	1	
Bacteroides fragilis		12.69	0.32		1	99	
Capnocytophaga ochracea		10.66	0.40		1	90	
47515430	**Very good identification**						**47515430**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.13	0.66	LAC 99	99	1	
Bacteroides vulgatus		0.76	0.15		1	3	
Capnocytophaga ochracea		0.11	0.07		1	90	
47515432	**Very good identification**						**47515432**
					GRA		
Bifidobacterium longum subsp. longum		95.06	0.66	LAC 99	99		
Clostridium clostridioforme		2.61	0.38		50		
Parabacteroides distasonis		1.22	0.38		1		
47515500	**Good identification**						**47515500**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		93.42	0.86		99	1	
Bacteroides vulgatus		4.85	0.50		1	3	
Capnocytophaga ochracea		0.71	0.41		1	90	
47515501	**Acceptable identification**						**47515501**
					GRA	CAT	
Bifidobacterium longum subsp. longum		93.59	0.52	URE 1	99	1	
Bacteroides vulgatus		4.86	0.15		1	3	
Capnocytophaga ochracea		0.71	0.07		1	90	
47515502	**Low discrimination**						**47515502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		68.55	0.86		99	1	
Bifidobacterium adolescentis		25.51	0.80		99	1	
Capnocytophaga ochracea		4.66	0.58		1	90	
47515503	**Low discrimination**						**47515503**
					GRA	CAT	
Bifidobacterium longum subsp. longum		68.84	0.52	URE 1	99	1	
Bifidobacterium adolescentis		25.62	0.46		99	1	
Capnocytophaga ochracea		4.68	0.23		1	90	
47515510	**Very good identification**						**47515510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.64	0.90		99	1	
Bifidobacterium adolescentis		0.27	0.47		99	1	
Capnocytophaga ochracea		0.05	0.25		1	90	
47515511	**Good identification**						**47515511**
Bifidobacterium longum subsp. longum		99.73	0.56	URE 1			
Bifidobacterium adolescentis		0.27	0.13				
47515512	**Low discrimination**						**47515512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		78.38	0.90		99	1	
Bifidobacterium adolescentis		20.94	0.82		99	1	
Capnocytophaga ochracea		0.35	0.41		1	90	
47515513	**Low discrimination**						**47515513**
					GRA	CAT	
Bifidobacterium longum subsp. longum		78.42	0.56	URE 1	99	1	
Bifidobacterium adolescentis		20.95	0.47		99	1	
Capnocytophaga ochracea		0.35	0.07		1	90	
47515520	**Low discrimination**						**47515520**
					GRA	CAT	
Bacteroides vulgatus		55.56	0.84	RHA 90	1	3	
Bifidobacterium longum subsp. longum		43.22	0.96		99	1	
Capnocytophaga ochracea		0.73	0.58		1	90	
47515521	**Low discrimination**						**47515521**
					GRA	CAT	
Bacteroides vulgatus		55.58	0.50	RHA 90	1	3	
Bifidobacterium longum subsp. longum		43.24	0.62	URE 1	99	1	
Capnocytophaga ochracea		0.73	0.23		1	90	
47515522	**Low discrimination**						**47515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		71.83	0.96		99	1	
Bacteroides fragilis		13.07	0.66		1	99	
Capnocytophaga ochracea		10.98	0.74		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47515523	**Low discrimination**						**47515523**
					GRA	CAT	
Bifidobacterium longum subsp. longum		71.90	0.62	URE 1	99	1	
Bacteroides fragilis		13.08	0.32		1	99	
Capnocytophaga ochracea		10.99	0.40		1	90	
47515530	**Excellent identification**						**47515530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.11	1.00		99	1	
Bacteroides vulgatus		0.76	0.50		1	3	
Capnocytophaga ochracea		0.11	0.41		1	90	
47515531	**Very good identification**						**47515531**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.13	0.66	URE 1	99	1	
Bacteroides vulgatus		0.76	0.15		1	3	
Capnocytophaga ochracea		0.11	0.07		1	90	
47515532	**Very good identification**						**47515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.51	1.00		99	1	
Parabacteroides distasonis		1.25	0.72		1	76	
Capnocytophaga ochracea		0.97	0.58		1	90	
47515533	**Very good identification**						**47515533**
					GRA	CAT	
Bifidobacterium longum subsp. longum		97.52	0.66	URE 1	99	1	
Parabacteroides distasonis		1.25	0.38		1	76	
Capnocytophaga ochracea		0.97	0.23		1	90	
47515540	**Low discrimination**						**47515540**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.09	0.52	SOR 1	99	1	
Capnocytophaga ochracea		6.90	0.25		1	90	
Bacteroides vulgatus		4.32	0.15		1	3	
47515542	**Low discrimination**						**47515542**
					GRA	CAT	
Bifidobacterium adolescentis		71.87	0.65	SOR 11	99	1	
Bifidobacterium longum subsp. longum		15.78	0.52		99	1	
Capnocytophaga ochracea		11.79	0.41		1	90	
47515550	**Good identification**						**47515550**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.18	0.56	SOR 1	99	1	
Bifidobacterium adolescentis		3.18	0.32		99	1	
Capnocytophaga ochracea		0.52	0.08		1	90	
47515552	**Low discrimination**						**47515552**
					GRA	CAT	
Bifidobacterium adolescentis		75.61	0.66	SOR 11	99	1	
Bifidobacterium longum subsp. longum		23.13	0.56		99	1	
Capnocytophaga ochracea		1.13	0.25		1	90	
47515560	**Low discrimination**						**47515560**
					GRA	CAT	
Bacteroides vulgatus		51.78	0.50	RHA 90	1	3	
Bifidobacterium longum subsp. longum		40.29	0.62	SOR 1	99	1	
Capnocytophaga ochracea		7.52	0.41		1	90	
47515562	**Low discrimination**						**47515562**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		57.38	0.58	SOR 10	1	90	
Bifidobacterium longum subsp. longum		34.13	0.62		99	1	
Bacteroides fragilis		6.21	0.32		1	99	
47515570	**Very good identification**						**47515570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.06	0.66	SOR 1	99	1	
Capnocytophaga ochracea		1.20	0.25		1	90	
Bacteroides vulgatus		0.75	0.15		1	3	
47515572	**Low discrimination**						**47515572**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.28	0.66	SOR 1	99	1	
Capnocytophaga ochracea		9.68	0.41		1	90	
Parabacteroides distasonis		1.13	0.38		1	76	
47515602	**Low discrimination**						**47515602**
Clostridium clostridioforme		89.35	0.41	SAL 88			
Bifidobacterium longum subsp. longum		3.00	0.17	TRE 88			
Eubacterium contortum		2.74	0.10				
47515612	**Good identification**						**47515612**
Clostridium clostridioforme		98.25	0.52	SAL 88			
Bifidobacterium longum subsp. longum		1.23	0.21	TRE 88			
Bifidobacterium adolescentis		0.33	0.13				
47515620	**Very good identification**						**47515620**
					GRA		
Bacteroides vulgatus		99.88	0.66	LAC 99	1		
Bifidobacterium longum subsp. longum		0.09	0.28		99		
Eubacterium contortum		0.02	0.10		99		
47515622	**Low discrimination**						**47515622**
Parabacteroides distasonis		36.10	0.50	LAC 99			
Bacteroides vulgatus		34.78	0.32				
Clostridium clostridioforme		22.43	0.41				
47515632	**Low discrimination**						**47515632**
					GRA		
Clostridium clostridioforme		79.96	0.52	SAL 88	50		
Parabacteroides distasonis		15.21	0.46	TRE 88	1		
Bifidobacterium longum subsp. longum		4.01	0.32		99		
47515700	**Low discrimination**						**47515700**
					GRA		
Bacteroides vulgatus		88.15	0.66	NAG 99	1		
Eubacterium contortum		9.88	0.57		99		
Bifidobacterium longum subsp. longum		1.91	0.52		99		
47515702	**Low discrimination**						**47515702**
Clostridium clostridioforme		23.17	0.50	SAL 88			
Bifidobacterium longum subsp. longum		22.96	0.52	TRE 88			
Eubacterium contortum		21.02	0.44				
47515710	**Low discrimination**						**47515710**
					GRA		
Bifidobacterium longum subsp. longum		74.81	0.56	RHA 1	99		
Bacteroides vulgatus		20.54	0.32		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		2.30	0.23		99	
47515712	**Low discrimination**					**47515712**
Clostridium clostridioforme		65.28	0.61	SAL 88		
Bifidobacterium longum subsp. longum		24.18	0.56	TRE 88		
Bifidobacterium adolescentis		6.46	0.47			
47515720	**Excellent identification**					**47515720**
					GRA	
Bacteroides vulgatus		99.88	1.00		1	
Bifidobacterium longum subsp. longum		0.09	0.62		99	
Eubacterium contortum		0.02	0.44		99	
47515721	**Very good identification**					**47515721**
					GRA	
Bacteroides vulgatus		99.88	0.66	URE 1	1	
Bifidobacterium longum subsp. longum		0.09	0.28		99	
Eubacterium contortum		0.02	0.10		99	
47515722	**Low discrimination**					**47515722**
					GRA	
Parabacteroides distasonis		46.95	0.85		1	
Bacteroides vulgatus		45.24	0.66		1	
Bifidobacterium longum subsp. longum		3.91	0.62		99	
47515723	**Low discrimination**					**47515723**
					GRA	
Parabacteroides distasonis		47.16	0.50	URE 1	1	
Bacteroides vulgatus		45.44	0.32		1	
Bifidobacterium longum subsp. longum		3.93	0.28		99	
47515730	**Low discrimination**					**47515730**
					GRA	
Bacteroides vulgatus		86.63	0.66	MLZ 1	1	
Bifidobacterium longum subsp. longum		12.75	0.66		99	
Parabacteroides distasonis		0.49	0.46		1	
47515732	**Low discrimination**					**47515732**
					GRA	
Parabacteroides distasonis		67.54	0.80		1	
Bifidobacterium longum subsp. longum		17.79	0.66		99	
Clostridium clostridioforme		12.01	0.61		50	
47515733	**Low discrimination**					**47515733**
					GRA	
Parabacteroides distasonis		67.70	0.46	URE 1	1	
Bifidobacterium longum subsp. longum		17.83	0.32		99	
Clostridium clostridioforme		12.03	0.27		50	
47515760	**Very good identification**					**47515760**
					GRA	
Bacteroides vulgatus		99.87	0.66	SOR 1	1	
Bifidobacterium longum subsp. longum		0.09	0.28		99	
Eubacterium contortum		0.02	0.10		99	
47515762	**Low discrimination**					**47515762**
					CAT	
Parabacteroides distasonis		44.49	0.50	SOR 1	76	
Bacteroides vulgatus		42.87	0.32		3	
Capnocytophaga ochracea		6.23	0.23		90	
47515772	**Low discrimination**					**47515772**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		66.40	0.46	SOR 1	1		
Bifidobacterium longum subsp. longum		17.49	0.32		99		
Clostridium clostridioforme		11.80	0.27		50		
47516502	**Good identification**						**47516502**
Bifidobacterium adolescentis		95.77	0.44	GAL 99			
Bifidobacterium longum subsp. longum		3.18	0.17				
Actinomyces israelii		1.05	0.17				
47516512	**Good identification**						**47516512**
Bifidobacterium adolescentis		95.07	0.46	GAL 99			
Bifidobacterium longum subsp. longum		4.39	0.21				
Clostridium clostridioforme		0.54	0.04				
47517000	**Low discrimination**						**47517000**
					GRA	CAT	
Actinomyces israelii		78.98	0.39	SAL 92	99	1	
Fusobacterium mortiferum		7.78	0.20	CEL 93	1	5	
Propionibacterium granulosum		6.71	0.16		99	90	
47517002	**Low discrimination**						**47517002**
					CAT	GRA	
Actinomyces israelii		61.91	0.43	SAL 92	1	99	
Bifidobacterium adolescentis		27.81	0.30	CEL 93	1	99	
Bacteroides fragilis		4.49	-.03		99	1	
47517042	**Low discrimination**						**47517042**
Actinomyces israelii		87.50	0.35	SAL 92			
Bifidobacterium adolescentis		12.50	0.15	CEL 93			
47517100	**Low discrimination**						**47517100**
					GRA	CAT	
Actinomyces israelii		58.80	0.50	SAL 92	99	1	
Bifidobacterium adolescentis		9.91	0.30	CEL 93	99	1	
Capnocytophaga ochracea		9.31	0.20		1	90	
47517102	**Low discrimination**						**47517102**
					CAT	GRA	
Bifidobacterium adolescentis		72.29	0.64	ARA 87	1	99	
Bacteroides fragilis		11.68	0.32		99	1	
Actinomyces israelii		6.50	0.53		1	99	
47517112	**Very good identification**						**47517112**
					GRA	CAT	
Bifidobacterium adolescentis		97.73	0.66	ARA 87	99	1	
Actinomyces israelii		1.07	0.39		99	1	
Capnocytophaga ochracea		0.76	0.20		1	90	
47517120	**Low discrimination**						**47517120**
					CAT		
Bacteroides fragilis		77.10	0.44	MAN 1	99		
Bacteroides vulgatus		7.87	0.30	BGL 95	3		
Capnocytophaga ochracea		7.83	0.36		90		
47517122	**Good identification**						**47517122**
					CAT		
Bacteroides fragilis		94.00	0.66	MAN 1	99		
Capnocytophaga ochracea		4.52	0.53		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		0.55	0.37		1		
47517140	**Acceptable identification**						**47517140**
					GRA	CAT	
Actinomyces israelii		91.01	0.43	SAL 92	99	1	
Bifidobacterium adolescentis		4.87	0.15	CEL 93	99	1	
Capnocytophaga ochracea		4.12	0.04		1	90	
47517142	**Low discrimination**						**47517142**
					GRA	CAT	
Bifidobacterium adolescentis		72.65	0.49	ARA 87	99	1	
Actinomyces israelii		20.56	0.46	SOR 11	99	1	
Capnocytophaga ochracea		5.58	0.20		1	90	
47517152	**Good identification**						**47517152**
					GRA	CAT	
Bifidobacterium adolescentis		96.01	0.50	ARA 87	99	1	
Actinomyces israelii		3.32	0.32	SOR 11	99	1	
Capnocytophaga ochracea		0.67	0.04		1	90	
47517320	**Good identification**						**47517320**
					GRA	CAT	
Bacteroides vulgatus		97.94	0.47	MAN 1	1	3	
Bacteroides fragilis		1.08	0.10	ARA 93	1	99	
Eubacterium contortum		0.32	0.11		99	1	
47517322	**Low discrimination**						**47517322**
					CAT		
Parabacteroides distasonis		44.53	0.54	MAN 1	76		
Bacteroides fragilis		40.94	0.32		99		
Prevotella oralis		10.21	0.30		1		
47517332	**Good identification**						**47517332**
Parabacteroides distasonis		96.78	0.49	MAN 1			
Bacteroides fragilis		1.67	-.03				
Capnocytophaga ochracea		0.88	0.02				
47517400	**Low discrimination**						**47517400**
Actinomyces israelii		77.14	0.38	SAL 92			
Bifidobacterium longum subsp. longum		14.11	0.17	CEL 93			
Lactobacillus fermentum		4.46	0.16				
47517402	**Low discrimination**						**47517402**
Bifidobacterium adolescentis		76.00	0.44	LAC 99			
Actinomyces israelii		20.68	0.41				
Bifidobacterium longum subsp. longum		2.52	0.17				
47517412	**Acceptable identification**						**47517412**
Bifidobacterium adolescentis		92.64	0.46	LAC 99			
Bifidobacterium longum subsp. longum		4.28	0.21				
Actinomyces israelii		3.08	0.27				
47517500	**Low discrimination**						**47517500**
Bifidobacterium longum subsp. longum		62.78	0.52	MAN 1			
Bifidobacterium adolescentis		19.11	0.44				
Actinomyces israelii		13.87	0.48				
47517502	**Very good identification**						**47517502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		95.53	0.79				
Bifidobacterium longum subsp. longum		3.17	0.52				
Actinomyces israelii		1.05	0.51				
47517503	**Good identification**						**47517503**
Bifidobacterium adolescentis		95.77	0.44	URE 1			
Bifidobacterium longum subsp. longum		3.18	0.17				
Actinomyces israelii		1.05	0.17				
47517510	**Low discrimination**						**47517510**
Bifidobacterium longum subsp. longum		80.78	0.56	MAN 1			
Bifidobacterium adolescentis		17.65	0.46				
Actinomyces israelii		1.57	0.34				
47517512	**Very good identification**						**47517512**
Bifidobacterium adolescentis		95.42	0.80				
Bifidobacterium longum subsp. longum		4.41	0.56				
Actinomyces israelii		0.13	0.37				
47517513	**Good identification**						**47517513**
Bifidobacterium adolescentis		95.58	0.46	URE 1			
Bifidobacterium longum subsp. longum		4.42	0.21				
47517520	**Low discrimination**						**47517520**
					GRA	CAT	
Bacteroides vulgatus		55.48	0.50	MAN 1	1	3	
Bifidobacterium longum subsp. longum		43.16	0.62	RHA 90	99	1	
Capnocytophaga ochracea		0.73	0.23		1	90	
47517522	**Low discrimination**						**47517522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		67.97	0.62	MAN 1	99	1	
Bacteroides fragilis		12.36	0.32		1	99	
Capnocytophaga ochracea		10.39	0.40		1	90	
47517530	**Very good identification**						**47517530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.05	0.66	MAN 1	99	1	
Bacteroides vulgatus		0.76	0.15		1	3	
Capnocytophaga ochracea		0.11	0.07		1	90	
47517532	**Good identification**						**47517532**
					GRA		
Bifidobacterium longum subsp. longum		92.53	0.66	MAN 1	99		
Bifidobacterium adolescentis		5.06	0.46		99		
Parabacteroides distasonis		1.18	0.38		1		
47517540	**Low discrimination**						**47517540**
Actinomyces israelii		64.29	0.41	SAL 92			
Bifidobacterium adolescentis		28.16	0.29	CEL 93			
Bifidobacterium longum subsp. longum		7.56	0.17				
47517542	**Very good identification**						**47517542**
Bifidobacterium adolescentis		96.22	0.63	SOR 11			
Actinomyces israelii		3.33	0.44				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		0.26	0.17			
47517552	**Very good identification**					**47517552**
Bifidobacterium adolescentis		99.21	0.65	SOR 11		
Actinomyces israelii		0.42	0.30			
Bifidobacterium longum subsp. longum		0.38	0.21			
47517702	**Low discrimination**					**47517702**
Bifidobacterium adolescentis		82.61	0.44	RHA 1		
Actinomyces israelii		6.77	0.32			
Clostridium clostridioforme		2.77	0.16			
47517712	**Low discrimination**					**47517712**
Bifidobacterium adolescentis		84.17	0.46	RHA 1		
Clostridium clostridioforme		10.50	0.27			
Bifidobacterium longum subsp. longum		3.89	0.21			
47517720	**Very good identification**					**47517720**
					GRA	
Bacteroides vulgatus		99.88	0.66	MAN 1	1	
Bifidobacterium longum subsp. longum		0.09	0.28		99	
Eubacterium contortum		0.02	0.10		99	
47517722	**Low discrimination**					**47517722**
					GRA	
Parabacteroides distasonis		46.81	0.50	MAN 1	1	
Bacteroides vulgatus		45.10	0.32		1	
Bifidobacterium longum subsp. longum		3.90	0.28		99	
47517732	**Low discrimination**					**47517732**
					GRA	
Parabacteroides distasonis		66.52	0.46	MAN 1	1	
Bifidobacterium longum subsp. longum		17.52	0.32		99	
Clostridium clostridioforme		11.82	0.27		50	
47521001	**Low discrimination**					**47521001**
					SPO	
Actinomyces naeslundii		79.36	0.39	ESL 12	1	
Clostridium chauvoei		20.65	0.06	FRU 95	99	
47521002	**Low discrimination**					**47521002**
					CAT	
Actinomyces naeslundii		41.10	0.36	ESL 12	25	
Eubacterium saburreum		30.34	0.21	FRU 95	1	
Propionibacterium granulosum		18.86	0.15		90	
47521003	**Good identification**					**47521003**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47521021	**Good identification**					**47521021**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47521023	**Good identification**					**47521023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47521100	**Low discrimination**					**47521100**
					GRA	
Eubacterium saburreum		39.35	0.40	IND 99	85	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		19.98	0.36		1		
Actinomyces naeslundii		11.48	0.44		99		
47521101	**Acceptable identification**						**47521101**
					SPO		
Clostridium chauvoei		90.05	0.40	RAF 1	99		
Actinomyces naeslundii		9.95	0.47	TRE 10	1		
47521102	**Low discrimination**						**47521102**
					GRA	CAT	
Actinomyces naeslundii		27.96	0.44	ESL 12	99	25	
Eubacterium saburreum		16.92	0.27	FRU 95	85	1	
Capnocytophaga ochracea		12.29	0.12		1	90	
47521110	**Low discrimination**						**47521110**
Eubacterium saburreum		86.29	0.38	IND 99			
Bifidobacterium longum subsp. longum		13.71	0.21				
47521120	**Low discrimination**						**47521120**
					GRA		
Prevotella melaninogenica		76.72	0.47	FRU 90	1		
Clostridium perfringens		19.52	0.43	TRE 1	99		
Bifidobacterium longum subsp. longum		1.11	0.27		99		
47521123	**Low discrimination**						**47521123**
					GRA	SPO	
Actinomyces naeslundii		57.96	0.47	ESL 12	99	1	
Clostridium chauvoei		30.85	0.19	FRU 95	99	99	
Prevotella melaninogenica		5.39	0.00		1	1	
47521132	**Low discrimination**						**47521132**
					GRA	CAT	
Parabacteroides distasonis		43.88	0.36	FRU 99	1	76	
Bifidobacterium longum subsp. longum		38.49	0.31	XYL 90	99	1	
Capnocytophaga ochracea		9.49	0.12		1	90	
47521322	**Low discrimination**						**47521322**
Parabacteroides distasonis		79.75	0.49	FRU 99			
Prevotella oralis		18.66	0.25	XYL 90			
Prevotella melaninogenica		0.98	0.00				
47521332	**Good identification**						**47521332**
Parabacteroides distasonis		99.01	0.44	FRU 99			
Clostridium clostridioforme		0.99	0.04	XYL 90			
47521500	**Good identification**						**47521500**
Bifidobacterium longum subsp. longum		100.0	0.51	FRU 90			
47521502	**Good identification**						**47521502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		95.84	0.51	FRU 90	99	1	
Bifidobacterium adolescentis		3.88	0.29		99	1	
Capnocytophaga ochracea		0.29	-.01		1	90	
47521510	**Good identification**						**47521510**
Bifidobacterium longum subsp. longum		100.0	0.55	FRU 90			
47521512	**Good identification**						**47521512**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		96.90	0.55	FRU 90			
Bifidobacterium adolescentis		2.81	0.30				
Clostridium clostridioforme		0.29	0.10				
47521520	**Very good identification**						**47521520**
					GRA		
Bifidobacterium longum subsp. longum		98.93	0.61	FRU 90	99		
Prevotella melaninogenica		0.70	0.12		1		
Clostridium perfringens		0.18	0.09		99		
47521522	**Very good identification**						**47521522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.91	0.61	FRU 90	99	1	
Parabacteroides distasonis		2.16	0.37		1	76	
Capnocytophaga ochracea		0.65	0.16		1	90	
47521530	**Very good identification**						**47521530**
Bifidobacterium longum subsp. longum		100.0	0.65	FRU 90			
47521532	**Very good identification**						**47521532**
					GRA		
Bifidobacterium longum subsp. longum		99.18	0.65	FRU 90	99		
Parabacteroides distasonis		0.70	0.33		1		
Clostridium clostridioforme		0.07	0.10		50		
47521722	**Low discrimination**						**47521722**
					GRA		
Parabacteroides distasonis		82.77	0.45	FRU 99	1		
Bifidobacterium longum subsp. longum		12.49	0.27	XYL 90	99		
Clostridium clostridioforme		2.49	0.14		50		
47521732	**Low discrimination**						**47521732**
					GRA		
Parabacteroides distasonis		57.10	0.41	FRU 99	1		
Bifidobacterium longum subsp. longum		27.24	0.31	XYL 90	99		
Clostridium clostridioforme		14.54	0.25		50		
47523000	**Low discrimination**						**47523000**
Actinomyces naeslundii		58.37	0.41	ESL 12			
Propionibacterium granulosum		41.63	0.24	FRU 95			
47523001	**Good identification**						**47523001**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47523002	**Low discrimination**						**47523002**
Actinomyces naeslundii		84.87	0.41	ESL 12			
Propionibacterium granulosum		15.13	0.13	FRU 95			
47523003	**Good identification**						**47523003**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47523020	**Low discrimination**						**47523020**
Actinomyces naeslundii		84.87	0.41	ESL 12			
Propionibacterium granulosum		15.13	0.13	FRU 95			
47523021	**Good identification**						**47523021**
Actinomyces naeslundii		100.0	0.44	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47523022	**Good identification**					**47523022**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47523023	**Good identification**					**47523023**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47523100	**Good identification**					**47523100**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47523101	**Good identification**					**47523101**
					SPO	
Actinomyces naeslundii		95.50	0.52	ESL 12	1	
Clostridium chauvoei		4.50	0.06	FRU 95	99	
47523102	**Acceptable identification**					**47523102**
Actinomyces naeslundii		93.81	0.49	ESL 12		
Bifidobacterium adolescentis		6.19	0.13	FRU 95		
47523103	**Good identification**					**47523103**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47523120	**Low discrimination**					**47523120**
					GRA	
Actinomyces naeslundii		67.90	0.49	ESL 12	99	
Prevotella melaninogenica		25.59	0.12	FRU 95	1	
Clostridium perfringens		6.51	0.09		99	
47523121	**Good identification**					**47523121**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47523122	**Low discrimination**					**47523122**
					GRA	
Actinomyces naeslundii		87.03	0.49	ESL 12	99	
Prevotella melaninogenica		6.25	0.00	FRU 95	1	
Clostridium perfringens		3.93	0.03		99	
47523123	**Good identification**					**47523123**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47523140	**Good identification**					**47523140**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47523141	**Good identification**					**47523141**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47523142	**Good identification**					**47523142**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47523143	**Good identification**					**47523143**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47523160	**Good identification**					**47523160**
Actinomyces naeslundii		100.0	0.36	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47523161	**Good identification**						**47523161**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47523162	**Good identification**						**47523162**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47523163	**Good identification**						**47523163**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47524520	**Good identification**						**47524520**
					GRA		
Bifidobacterium longum subsp. longum		98.09	0.35	FRU 90	99		
Bacteroides vulgatus		1.91	-.08	GAL 99	1		
47524522	**Acceptable identification**						**47524522**
					GRA		
Bifidobacterium longum subsp. longum		91.11	0.35	FRU 90	99		
Parabacteroides distasonis		6.10	0.19	GAL 99	1		
Clostridium clostridioforme		2.79	0.08		50		
47524530	**Good identification**						**47524530**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47524532	**Acceptable identification**						**47524532**
					GRA		
Bifidobacterium longum subsp. longum		90.66	0.39	FRU 90	99		
Clostridium clostridioforme		7.42	0.19	GAL 99	50		
Parabacteroides distasonis		1.92	0.15		1		
47525100	**Low discrimination**						**47525100**
					COC		
Eubacterium saburreum		44.47	0.35	IND 99	1		
Atopobium parvulum		33.47	0.23		99		
Bifidobacterium longum subsp. longum		18.91	0.25		1		
47525120	**Low discrimination**						**47525120**
					CAT	GRA	
Bifidobacterium longum subsp. longum		47.44	0.35	FRU 90	1	99	
Bacteroides fragilis		16.35	0.10	ARA 99	99	1	
Prevotella melaninogenica		11.05	0.12		1	1	
47525122	**Low discrimination**						**47525122**
					GRA	CAT	
Parabacteroides distasonis		56.81	0.57	FRU 99	1	76	
Bacteroides fragilis		34.40	0.32		1	99	
Bifidobacterium longum subsp. longum		5.25	0.35		99	1	
47525130	**Acceptable identification**						**47525130**
					GRA		
Bifidobacterium longum subsp. longum		92.22	0.39	FRU 90	99		
Eubacterium saburreum		4.60	0.21	ARA 99	85		
Parabacteroides distasonis		3.19	0.18		1		
47525132	**Low discrimination**						**47525132**
					GRA	CAT	
Parabacteroides distasonis		76.09	0.53	FRU 99	1	76	
Bifidobacterium longum subsp. longum		22.24	0.39		99	1	
Bacteroides fragilis		0.86	-.03		1	99	
47525322	**Very good identification**						**47525322**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Parabacteroides distasonis		99.35	0.65	FRU 99	76	
Prevotella oralis		0.32	0.10		1	
Bacteroides fragilis		0.20	-.03		99	
47525332	**Very good identification**					**47525332**
Parabacteroides distasonis		98.89	0.61	FRU 99		
Clostridium clostridioforme		1.11	0.22			
47525420	**Good identification**					**47525420**
					GRA	
Bifidobacterium longum subsp. longum		98.09	0.35	FRU 90	99	
Bacteroides vulgatus		1.91	-.08	LAC 99	1	
47525422	**Acceptable identification**					**47525422**
					GRA	
Bifidobacterium longum subsp. longum		91.38	0.35	FRU 90	99	
Parabacteroides distasonis		6.12	0.19	LAC 99	1	
Clostridium clostridioforme		2.50	0.07		50	
47525430	**Good identification**					**47525430**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90		
47525432	**Acceptable identification**					**47525432**
					GRA	
Bifidobacterium longum subsp. longum		91.37	0.39	FRU 90	99	
Clostridium clostridioforme		6.70	0.18	LAC 99	50	
Parabacteroides distasonis		1.94	0.15		1	
47525500	**Very good identification**					**47525500**
					SPO	
Bifidobacterium longum subsp. longum		99.68	0.59	FRU 90	1	
Clostridium butyricum		0.15	-.03		99	
Bifidobacterium adolescentis		0.09	0.09		1	
47525502	**Good identification**					**47525502**
Bifidobacterium longum subsp. longum		91.12	0.59	FRU 90		
Bifidobacterium adolescentis		8.22	0.43			
Clostridium clostridioforme		0.34	0.16			
47525510	**Very good identification**					**47525510**
Bifidobacterium longum subsp. longum		99.94	0.63	FRU 90		
Bifidobacterium adolescentis		0.07	0.10			
47525512	**Good identification**					**47525512**
Bifidobacterium longum subsp. longum		92.98	0.63	FRU 90		
Bifidobacterium adolescentis		6.02	0.44			
Clostridium clostridioforme		0.92	0.27			
47525520	**Very good identification**					**47525520**
					GRA	
Bifidobacterium longum subsp. longum		98.03	0.69	FRU 90	99	
Bacteroides vulgatus		1.91	0.26		1	
Parabacteroides distasonis		0.07	0.19		1	
47525521	**Good identification**					**47525521**
					GRA	
Bifidobacterium longum subsp. longum		98.09	0.35	FRU 90	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		1.91	-.08	URE 1	1		
47525522	**Good identification**						**47525522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		93.40	0.69	FRU 90	99	1	
Parabacteroides distasonis		6.26	0.53		1	76	
Bacteroides thetaiotaomicron		0.10	0.07		1	89	
47525523	**Acceptable identification**						**47525523**
					GRA		
Bifidobacterium longum subsp. longum		93.72	0.35	FRU 90	99		
Parabacteroides distasonis		6.28	0.19	URE 1	1		
47525530	**Very good identification**						**47525530**
					GRA		
Bifidobacterium longum subsp. longum		99.97	0.73	FRU 90	99		
Parabacteroides distasonis		0.02	0.15		1		
Bacteroides vulgatus		0.01	-.08		1		
47525531	**Good identification**						**47525531**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47525532	**Very good identification**						**47525532**
					GRA		
Bifidobacterium longum subsp. longum		97.65	0.73	FRU 90	99		
Parabacteroides distasonis		2.07	0.49		1		
Clostridium clostridioforme		0.24	0.27		50		
47525533	**Good identification**						**47525533**
					GRA		
Bifidobacterium longum subsp. longum		97.93	0.39	FRU 90	99		
Parabacteroides distasonis		2.07	0.15	URE 1	1		
47525560	**Good identification**						**47525560**
					GRA		
Bifidobacterium longum subsp. longum		98.09	0.35	FRU 90	99		
Bacteroides vulgatus		1.91	-.08	SOR 1	1		
47525562	**Acceptable identification**						**47525562**
					GRA		
Bifidobacterium longum subsp. longum		93.72	0.35	FRU 90	99		
Parabacteroides distasonis		6.28	0.19	SOR 1	1		
47525570	**Good identification**						**47525570**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47525572	**Good identification**						**47525572**
					GRA		
Bifidobacterium longum subsp. longum		97.93	0.39	FRU 90	99		
Parabacteroides distasonis		2.07	0.15	SOR 1	1		
47525712	**Low discrimination**						**47525712**
					GRA		
Clostridium clostridioforme		84.52	0.42	SAL 88	50		
Bifidobacterium longum subsp. longum		11.74	0.29	FRU 99	99		
Parabacteroides distasonis		2.98	0.23		1		
47525720	**Acceptable identification**						**47525720**
					GRA		
Bacteroides vulgatus		93.53	0.42	FRU 99	1		
Bifidobacterium longum subsp. longum		5.39	0.35	TRE 4	99		
Parabacteroides distasonis		1.08	0.27		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47525722	**Low discrimination**						**47525722**
					GRA	CAT	
Parabacteroides distasonis		85.81	0.62	FRU 99	1	76	
Bacteroides thetaiotaomicron		6.22	0.26		1	89	
Bifidobacterium longum subsp. longum		4.32	0.35		99	1	
47525730	**Low discrimination**						**47525730**
					GRA		
Bifidobacterium longum subsp. longum		84.41	0.39	FRU 90	99		
Bacteroides vulgatus		8.69	0.08	RHA 1	1		
Parabacteroides distasonis		5.36	0.23		1		
47525732	**Low discrimination**						**47525732**
					GRA		
Parabacteroides distasonis		66.78	0.57	FRU 99	1		
Clostridium clostridioforme		19.11	0.42		50		
Bifidobacterium longum subsp. longum		10.62	0.39		99		
47527502	**Low discrimination**						**47527502**
Bifidobacterium adolescentis		83.41	0.41	FRU 99			
Bifidobacterium longum subsp. longum		11.42	0.25				
Actinomyces israelii		5.18	0.27				
47527512	**Low discrimination**						**47527512**
Bifidobacterium adolescentis		82.99	0.43	FRU 99			
Bifidobacterium longum subsp. longum		15.82	0.29				
Bifidobacterium dentium		1.19	-.07				
47527520	**Good identification**						**47527520**
					GRA		
Bifidobacterium longum subsp. longum		98.09	0.35	FRU 90	99		
Bacteroides vulgatus		1.91	-.08	MAN 1	1		
47527522	**Acceptable identification**						**47527522**
					GRA		
Bifidobacterium longum subsp. longum		92.13	0.35	FRU 90	99		
Parabacteroides distasonis		6.17	0.19	MAN 1	1		
Bifidobacterium adolescentis		1.70	0.07		99		
47527530	**Good identification**						**47527530**
Bifidobacterium longum subsp. longum		100.0	0.39	FRU 90			
47527532	**Acceptable identification**						**47527532**
					GRA		
Bifidobacterium longum subsp. longum		93.66	0.39	FRU 90	99		
Bifidobacterium dentium		3.12	-.03	MAN 1	99		
Parabacteroides distasonis		1.98	0.15		1		
47530000	**Low discrimination**						**47530000**
					GRA	CAT	
Propionibacterium granulosum		56.02	0.60	ESL 1	99	90	
Fusobacterium mortiferum		43.21	0.61		1	5	
Prevotella intermedia		0.39	0.15		1	1	
47530002	**Low discrimination**						**47530002**
					GRA	CAT	
Propionibacterium granulosum		72.81	0.50	ESL 1	99	90	
Fusobacterium mortiferum		24.96	0.45		1	5	
Actinomyces naeslundii		1.27	0.34		99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47530003	**Low discrimination**						**47530003**
Actinomyces naeslundii		59.23	0.37	ESL 12			
Propionibacterium granulosum		22.89	0.15	GAL 96			
Actinomyces viscosus		10.04	0.00				
47530010	**Acceptable identification**						**47530010**
					GRA	CAT	
Propionibacterium granulosum		94.60	0.45	ESL 1	99	90	
Fusobacterium mortiferum		5.40	0.27	MLZ 12	1	5	
47530020	**Low discrimination**						**47530020**
					GRA	CAT	
Propionibacterium granulosum		82.63	0.50	ESL 1	99	90	
Clostridium perfringens		13.01	0.28		99	1	
Fusobacterium mortiferum		2.58	0.27		1	5	
47530021	**Low discrimination**						**47530021**
Actinomyces naeslundii		72.13	0.37	ESL 12			
Propionibacterium granulosum		27.87	0.15	GAL 96			
47530022	**Low discrimination**						**47530022**
					GRA	CAT	
Propionibacterium granulosum		67.62	0.39	ESL 1	99	90	
Clostridium perfringens		20.04	0.23		99	1	
Prevotella oralis		6.69	0.11		1	1	
47530023	**Good identification**						**47530023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47530100	**Very good identification**						**47530100**
					COC	GRA	
Fusobacterium mortiferum		97.52	0.68	MLT 11	1	1	
Actinomyces naeslundii		0.67	0.42		1	99	
Atopobium parvulum		0.64	0.19		99	99	
47530101	**Low discrimination**						**47530101**
					GRA	SPO	
Actinomyces naeslundii		44.51	0.45	ESL 12	99	1	
Fusobacterium mortiferum		43.50	0.33	GAL 96	1	1	
Clostridium chauvoei		11.99	0.12		99	99	
47530102	**Low discrimination**						**47530102**
					GRA	CAT	
Fusobacterium mortiferum		82.18	0.51	MLT 11	1	5	
Actinomyces naeslundii		5.10	0.42	BGL 10	99	25	
Capnocytophaga ochracea		2.83	0.12		1	90	
47530103	**Low discrimination**						**47530103**
					GRA		
Actinomyces naeslundii		79.24	0.45	ESL 12	99		
Fusobacterium mortiferum		8.60	0.17	GAL 96	1		
Actinomyces viscosus		8.39	0.04		99		
47530120	**Low discrimination**						**47530120**
					GRA		
Clostridium perfringens		65.53	0.43	GAL 99	99		
Prevotella melaninogenica		23.42	0.29		1		
Fusobacterium mortiferum		4.13	0.33		1		
47530121	**Low discrimination**						**47530121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		76.77	0.45	ESL 12	1		
Clostridium perfringens		12.03	0.09	GAL 96	84		
Clostridium chauvoei		6.89	0.04		99		
47530122	**Low discrimination**						**47530122**
					GRA	CAT	
Clostridium perfringens		34.38	0.38	GAL 99	99	1	
Prevotella oralis		24.51	0.32		1	1	
Capnocytophaga ochracea		15.69	0.29		1	90	
47530123	**Low discrimination**						**47530123**
					GRA		
Actinomyces naeslundii		88.78	0.45	ESL 12	99		
Clostridium perfringens		6.55	0.03	GAL 96	99		
Prevotella oralis		4.67	-.03		1		
47530132	**Low discrimination**						**47530132**
Parabacteroides distasonis		66.48	0.36	GAL 99			
Capnocytophaga ochracea		14.38	0.12	XYL 90			
Clostridium clostridioforme		8.94	0.16				
47530322	**Low discrimination**						**47530322**
Parabacteroides distasonis		78.09	0.49	GAL 99			
Prevotella oralis		18.27	0.25	XYL 90			
Clostridium clostridioforme		3.04	0.19				
47530332	**Low discrimination**						**47530332**
Parabacteroides distasonis		75.26	0.44	GAL 99			
Clostridium clostridioforme		24.75	0.30	XYL 90			
47530510	**Good identification**						**47530510**
Bifidobacterium longum subsp. longum		100.0	0.37	GAL 99			
47530512	**Low discrimination**						**47530512**
Clostridium clostridioforme		44.74	0.36	SAL 88			
Bifidobacterium longum subsp. longum		41.89	0.37	XYL 91			
Bifidobacterium adolescentis		13.38	0.30	RHA 88			
47530520	**Good identification**						**47530520**
					GRA		
Bifidobacterium longum subsp. longum		95.41	0.43	GAL 99	99		
Clostridium perfringens		1.89	0.09		99		
Bacteroides vulgatus		1.25	-.03		1		
47530522	**Low discrimination**						**47530522**
					GRA		
Bifidobacterium longum subsp. longum		69.67	0.43	GAL 99	99		
Parabacteroides distasonis		17.11	0.37		1		
Clostridium clostridioforme		6.95	0.25		50		
47530530	**Good identification**						**47530530**
Bifidobacterium longum subsp. longum		100.0	0.47	GAL 99			
47530532	**Low discrimination**						**47530532**
					GRA		
Bifidobacterium longum subsp. longum		74.10	0.47	GAL 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		19.78	0.36		50		
Parabacteroides distasonis		5.75	0.33		1		
47530612	**Good identification**						**47530612**
Clostridium clostridioforme		100.0	0.42	SAL 88			
47530632	**Good identification**						**47530632**
Clostridium clostridioforme		100.0	0.42	SAL 88			
47530702	**Good identification**						**47530702**
Clostridium clostridioforme		98.99	0.40	SAL 88			
Parabacteroides distasonis		1.01	0.11	XYL 91			
47530712	**Good identification**						**47530712**
Clostridium clostridioforme		100.0	0.51	SAL 88			
47530722	**Low discrimination**						**47530722**
Parabacteroides distasonis		49.16	0.45	GAL 99			
Clostridium clostridioforme		48.83	0.40	XYL 90			
Bacteroides thetaiotaomicron		1.34	0.02				
47530732	**Low discrimination**						**47530732**
					GRA		
Clostridium clostridioforme		88.77	0.51	SAL 88	50		
Parabacteroides distasonis		10.56	0.41	XYL 91	1		
Bifidobacterium longum subsp. longum		0.46	0.13		99		
47531000	**Low discrimination**						**47531000**
					GRA	CAT	
Fusobacterium mortiferum		46.43	0.64	MLT 11	1	5	
Propionibacterium granulosum		40.13	0.60		99	90	
Clostridium perfringens		4.84	0.37		99	1	
47531001	**Low discrimination**						**47531001**
					GRA	CAT	
Actinomyces naeslundii		84.85	0.61	ESL 12	99	25	
Fusobacterium mortiferum		6.32	0.30		1	5	
Propionibacterium granulosum		5.46	0.26		99	90	
47531002	**Low discrimination**						**47531002**
					GRA	CAT	
Propionibacterium granulosum		40.74	0.50	ESL 1	99	90	
Fusobacterium mortiferum		20.95	0.48		1	5	
Actinomyces naeslundii		17.04	0.58		99	25	
47531003	**Very good identification**						**47531003**
					GRA	CAT	
Actinomyces naeslundii		97.09	0.61	ESL 12	99	25	
Propionibacterium granulosum		1.56	0.15		99	90	
Fusobacterium mortiferum		0.80	0.14		1	5	
47531010	**Low discrimination**						**47531010**
					CAT	GRA	
Propionibacterium granulosum		63.04	0.45	ESL 1	90	99	
Eubacterium saburreum		26.12	0.40	MLZ 12	1	85	
Fusobacterium mortiferum		5.40	0.30		5	1	
47531011	**Low discrimination**						**47531011**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		85.70	0.39	ESL 12			
Propionibacterium granulosum		14.30	0.11	MLZ 5			
47531013	**Good identification**						**47531013**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47531020	**Low discrimination**						**47531020**
					CAT		
Clostridium perfringens		88.07	0.63	LAC 88	1		
Propionibacterium granulosum		5.65	0.50		90		
Actinomyces naeslundii		2.36	0.58		25		
47531021	**Low discrimination**						**47531021**
					CAT		
Actinomyces naeslundii		77.86	0.61	ESL 12	25		
Clostridium perfringens		19.54	0.28		1		
Propionibacterium granulosum		1.25	0.15		90		
47531022	**Low discrimination**						**47531022**
					GRA		
Clostridium perfringens		66.30	0.57	LAC 88	99		
Prevotella oralis		22.12	0.45		1		
Actinomyces naeslundii		3.78	0.58		99		
47531023	**Low discrimination**						**47531023**
					GRA		
Actinomyces naeslundii		86.40	0.61	ESL 12	99		
Clostridium perfringens		10.20	0.23		99		
Prevotella oralis		3.40	0.11		1		
47531031	**Good identification**						**47531031**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47531032	**Low discrimination**						**47531032**
					GRA	CAT	
Parabacteroides distasonis		37.20	0.36	XYL 90	1	76	
Clostridium perfringens		21.64	0.23	LAC 99	99	1	
Propionibacterium granulosum		9.96	0.25		99	90	
47531033	**Good identification**						**47531033**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47531041	**Good identification**						**47531041**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47531042	**Low discrimination**						**47531042**
					CAT		
Actinomyces naeslundii		54.14	0.45	ESL 12	25		
Clostridium perfringens		31.69	0.19		1		
Propionibacterium granulosum		7.41	0.15		90		
47531043	**Good identification**						**47531043**
Actinomyces naeslundii		100.0	0.48	ESL 12			
47531060	**Good identification**						**47531060**
					CAT		
Clostridium perfringens		97.12	0.50	LAC 88	1		
Actinomyces naeslundii		2.42	0.45		25		
Propionibacterium granulosum		0.33	0.15		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47531061	**Low discrimination**						**47531061**
Actinomyces naeslundii		78.69	0.48	ESL 12			
Clostridium perfringens		21.31	0.16				
47531062	**Acceptable identification**						**47531062**
					GRA	CAT	
Clostridium perfringens		91.72	0.45	LAC 88	99	1	
Actinomyces naeslundii		4.85	0.45		99	25	
Capnocytophaga ochracea		1.81	0.12		1	90	
47531063	**Low discrimination**						**47531063**
Actinomyces naeslundii		88.70	0.48	ESL 12			
Clostridium perfringens		11.30	0.11				
47531100	**Low discrimination**						**47531100**
					COC	GRA	
Fusobacterium mortiferum		49.25	0.71	MLT 11	1	1	
Atopobium parvulum		21.17	0.53		99	99	
Clostridium perfringens		16.12	0.52		1	99	
47531101	**Low discrimination**						**47531101**
					GRA	SPO	
Clostridium chauvoei		49.95	0.47	RAF 1	99	99	
Actinomyces naeslundii		44.95	0.69	TRE 10	99	1	
Fusobacterium mortiferum		2.75	0.36		1	1	
47531102	**Low discrimination**						**47531102**
					GRA	CAT	
Capnocytophaga ochracea		23.34	0.47	TRE 1	1	90	
Prevotella oralis		17.27	0.44	NAG 90	1	1	
Clostridium perfringens		14.24	0.46		99	1	
47531103	**Low discrimination**						**47531103**
					GRA	SPO	
Actinomyces naeslundii		79.75	0.69	ESL 12	99	1	
Clostridium chauvoei		15.64	0.34		99	99	
Capnocytophaga ochracea		1.23	0.12		1	1	
47531110	**Low discrimination**						**47531110**
					GRA		
Eubacterium saburreum		48.09	0.46	IND 99	85		
Bifidobacterium longum subsp. longum		22.92	0.37		99		
Fusobacterium mortiferum		9.95	0.36		1		
47531111	**Low discrimination**						**47531111**
					SPO		
Actinomyces naeslundii		78.74	0.47	ESL 12	1		
Clostridium chauvoei		16.79	0.12	MLZ 5	99		
Eubacterium saburreum		4.46	0.12		1		
47531112	**Low discrimination**						**47531112**
					GRA	CAT	
Bifidobacterium adolescentis		46.01	0.50	XYL 87	99	1	
Leptotrichia buccalis		15.68	0.24	ARA 87	1	1	
Capnocytophaga ochracea		11.74	0.30		1	90	
47531113	**Low discrimination**						**47531113**
					GRA		
Actinomyces naeslundii		82.74	0.47	ESL 12	99		
Bifidobacterium adolescentis		10.55	0.16	MLZ 5	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		3.60	-.11		1		
47531120	**Low discrimination**						**47531120**
					GRA	CAT	
Clostridium perfringens		70.02	0.77		99	1	
Prevotella melaninogenica		25.02	0.63		1	1	
Capnocytophaga ochracea		1.67	0.47		1	90	
47531121	**Low discrimination**						**47531121**
					SPO		
Actinomyces naeslundii		43.81	0.69	ESL 12	1		
Clostridium perfringens		28.32	0.43		84		
Clostridium chauvoei		16.23	0.38		99		
47531122	**Low discrimination**						**47531122**
					GRA	CAT	
Clostridium perfringens		35.41	0.72		99	1	
Prevotella oralis		25.24	0.66		1	1	
Capnocytophaga ochracea		16.16	0.63		1	90	
47531123	**Low discrimination**						**47531123**
					GRA		
Actinomyces naeslundii		52.20	0.69	ESL 12	99		
Clostridium perfringens		15.88	0.38		99		
Prevotella oralis		11.32	0.32		1		
47531130	**Low discrimination**						**47531130**
					GRA		
Clostridium perfringens		36.16	0.43	MLZ 1	99		
Bifidobacterium longum subsp. longum		31.48	0.47		99		
Prevotella melaninogenica		12.92	0.29		1		
47531131	**Low discrimination**						**47531131**
Actinomyces naeslundii		74.21	0.47	ESL 12			
Clostridium perfringens		9.21	0.09	MLZ 5			
Bifidobacterium longum subsp. longum		8.02	0.13				
47531132	**Low discrimination**						**47531132**
					GRA	CAT	
Parabacteroides distasonis		71.58	0.70	XYL 90	1	76	
Capnocytophaga ochracea		15.48	0.47		1	90	
Bifidobacterium longum subsp. longum		5.71	0.47		99	1	
47531133	**Low discrimination**						**47531133**
					GRA		
Parabacteroides distasonis		47.44	0.36	XYL 90	1		
Actinomyces naeslundii		35.02	0.47	URE 1	99		
Capnocytophaga ochracea		10.26	0.12		1		
47531140	**Low discrimination**						**47531140**
					GRA		
Clostridium perfringens		61.52	0.39	NAG 97	99		
Actinomyces naeslundii		19.20	0.53		99		
Fusobacterium mortiferum		9.97	0.36		1		
47531141	**Acceptable identification**						**47531141**
					SPO		
Actinomyces naeslundii		92.15	0.56	ESL 12	1		
Clostridium chauvoei		5.86	0.12		99		
Clostridium perfringens		1.99	0.05		84		
47531143	**Good identification**						**47531143**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		98.89	0.56	ESL 12	1		
Clostridium chauvoei		1.11	-.01		99		
47531160	**Very good identification**						**47531160**
					GRA	CAT	
Clostridium perfringens		95.75	0.65		99	1	
Prevotella melaninogenica		1.82	0.29		1	1	
Capnocytophaga ochracea		1.33	0.30		1	90	
47531161	**Low discrimination**						**47531161**
					SPO		
Actinomyces naeslundii		57.73	0.56	ESL 12	1		
Clostridium perfringens		40.28	0.31		84		
Clostridium chauvoei		1.22	0.04		99		
47531162	**Low discrimination**						**47531162**
					GRA	CAT	
Clostridium perfringens		73.35	0.59		99	1	
Capnocytophaga ochracea		19.52	0.47		1	90	
Prevotella oralis		2.77	0.32		1	1	
47531163	**Low discrimination**						**47531163**
					GRA	CAT	
Actinomyces naeslundii		70.02	0.56	ESL 12	99	25	
Clostridium perfringens		22.99	0.25		99	1	
Capnocytophaga ochracea		6.12	0.12		1	90	
47531222	**Low discrimination**						**47531222**
					GRA		
Prevotella oralis		56.12	0.39	TRE 6	1		
Parabacteroides distasonis		37.96	0.49	LAC 94	1		
Clostridium perfringens		3.96	0.23		99		
47531232	**Low discrimination**						**47531232**
Parabacteroides distasonis		75.62	0.44	XYL 90			
Clostridium clostridioforme		22.28	0.30	LAC 99			
Prevotella oralis		2.10	0.05				
47531300	**Low discrimination**						**47531300**
					COC	GRA	
Fusobacterium mortiferum		40.06	0.36	MLT 11	1	1	
Atopobium parvulum		17.22	0.19	RHA 1	99	99	
Prevotella oralis		16.72	0.16		1	1	
47531302	**Low discrimination**						**47531302**
Prevotella oralis		49.50	0.38	TRE 6			
Parabacteroides distasonis		40.61	0.49	NAG 95			
Clostridium clostridioforme		4.74	0.27				
47531312	**Low discrimination**						**47531312**
					GRA		
Parabacteroides distasonis		48.75	0.44	XYL 90	1		
Clostridium clostridioforme		48.09	0.39	NAG 99	50		
Bifidobacterium adolescentis		1.53	0.16		99		
47531320	**Low discrimination**						**47531320**
					GRA		
Clostridium perfringens		30.95	0.43	RHA 1	99		
Prevotella oralis		23.18	0.38		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Parabacteroides distasonis	19.02	0.49		1	
47531322	**Low discrimination**					**47531322**
					GRA	
	Parabacteroides distasonis	80.05	0.83	XYL 90	1	
	Prevotella oralis	18.73	0.60		1	
	Clostridium perfringens	0.62	0.38		99	
47531323	**Low discrimination**					**47531323**
					GRA	
	Parabacteroides distasonis	78.92	0.49	XYL 90	1	
	Prevotella oralis	18.46	0.25	URE 1	1	
	Actinomyces naeslundii	2.01	0.35		99	
47531330	**Low discrimination**					**47531330**
					GRA	
	Parabacteroides distasonis	89.72	0.44	XYL 90	1	
	Clostridium perfringens	2.74	0.09	BGL 99	99	
	Bifidobacterium longum subsp. longum	2.38	0.13		99	
47531332	**Very good identification**					**47531332**
	Parabacteroides distasonis	98.46	0.79	XYL 90		
	Clostridium clostridioforme	0.98	0.39			
	Prevotella oralis	0.43	0.25			
47531333	**Good identification**					**47531333**
	Parabacteroides distasonis	99.01	0.44	XYL 90		
	Clostridium clostridioforme	0.99	0.04	URE 1		
47531362	**Low discrimination**					**47531362**
					GRA	
	Parabacteroides distasonis	70.49	0.49	XYL 90	1	
	Prevotella oralis	16.49	0.25	SOR 1	1	
	Clostridium perfringens	10.28	0.25		99	
47531372	**Good identification**					**47531372**
	Parabacteroides distasonis	99.01	0.44	XYL 90		
	Clostridium clostridioforme	0.99	0.04	SOR 1		
47531410	**Good identification**					**47531410**
					CAT	
	Bifidobacterium longum subsp. longum	97.85	0.37	LAC 99	1	
	Propionibacterium granulosum	2.15	0.11		90	
47531412	**Low discrimination**					**47531412**
	Bifidobacterium longum subsp. longum	43.93	0.37	LAC 99		
	Clostridium clostridioforme	42.04	0.35			
	Bifidobacterium adolescentis	14.03	0.30			
47531420	**Low discrimination**					**47531420**
					CAT	
	Bifidobacterium longum subsp. longum	75.79	0.43	LAC 99	1	
	Clostridium perfringens	20.21	0.28		1	
	Propionibacterium granulosum	1.30	0.15		90	
47531422	**Low discrimination**					**47531422**
					GRA	
	Bifidobacterium longum subsp. longum	63.15	0.43	LAC 99	99	
	Parabacteroides distasonis	15.51	0.37		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		7.93	0.23		99		
47531430	**Good identification**						**47531430**
Bifidobacterium longum subsp. longum		100.0	0.47	LAC 99			
47531432	**Low discrimination**						**47531432**
					GRA		
Bifidobacterium longum subsp. longum		75.66	0.47	LAC 99	99		
Clostridium clostridioforme		18.10	0.35		50		
Parabacteroides distasonis		5.88	0.33		1		
47531500	**Very good identification**						**47531500**
					GRA		
Bifidobacterium longum subsp. longum		97.33	0.67		99		
Fusobacterium mortiferum		0.73	0.36		1		
Bifidobacterium adolescentis		0.44	0.29		99		
47531502	**Low discrimination**						**47531502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		66.28	0.67		99	1	
Bifidobacterium adolescentis		29.48	0.63		99	1	
Capnocytophaga ochracea		2.18	0.34		1	90	
47531510	**Very good identification**						**47531510**
Bifidobacterium longum subsp. longum		99.60	0.71				
Bifidobacterium adolescentis		0.32	0.30				
Clostridium clostridioforme		0.03	0.10				
47531511	**Good identification**						**47531511**
Bifidobacterium longum subsp. longum		100.0	0.37	URE 1			
47531512	**Low discrimination**						**47531512**
Bifidobacterium longum subsp. longum		73.68	0.71				
Bifidobacterium adolescentis		23.52	0.64				
Clostridium clostridioforme		2.38	0.44				
47531513	**Low discrimination**						**47531513**
Bifidobacterium longum subsp. longum		73.98	0.37	URE 1			
Bifidobacterium adolescentis		23.62	0.30				
Clostridium clostridioforme		2.39	0.10				
47531520	**Very good identification**						**47531520**
					GRA		
Bifidobacterium longum subsp. longum		95.11	0.78		99		
Clostridium perfringens		1.88	0.43		99		
Bacteroides vulgatus		1.25	0.31		1		
47531521	**Acceptable identification**						**47531521**
Bifidobacterium longum subsp. longum		91.74	0.43	URE 1			
Actinomyces naeslundii		2.80	0.35				
Clostridium perfringens		1.81	0.09				
47531522	**Low discrimination**						**47531522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.44	0.78		99	1	
Parabacteroides distasonis		18.28	0.71		1	76	
Capnocytophaga ochracea		5.52	0.50		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47531523	**Low discrimination**						**47531523**
					GRA	CAT	
Bifidobacterium longum subsp. longum		73.20	0.43	URE 1	99	1	
Parabacteroides distasonis		17.97	0.37		1	76	
Capnocytophaga ochracea		5.42	0.16		1	90	
47531530	**Very good identification**						**47531530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.84	0.82		99	1	
Parabacteroides distasonis		0.08	0.33		1	76	
Capnocytophaga ochracea		0.05	0.17		1	90	
47531531	**Good identification**						**47531531**
Bifidobacterium longum subsp. longum		100.0	0.47	URE 1			
47531532	**Good identification**						**47531532**
					GRA		
Bifidobacterium longum subsp. longum		91.59	0.82		99		
Parabacteroides distasonis		7.11	0.67		1		
Clostridium clostridioforme		0.74	0.44		50		
47531533	**Acceptable identification**						**47531533**
					GRA		
Bifidobacterium longum subsp. longum		91.70	0.47	URE 1	99		
Parabacteroides distasonis		7.12	0.33		1		
Clostridium clostridioforme		0.74	0.10		50		
47531542	**Low discrimination**						**47531542**
					GRA	CAT	
Bifidobacterium adolescentis		78.95	0.47	XYL 87	99	1	
Bifidobacterium longum subsp. longum		14.51	0.33	SOR 11	99	1	
Capnocytophaga ochracea		5.25	0.17		1	90	
47531550	**Good identification**						**47531550**
Bifidobacterium longum subsp. longum		96.20	0.37	SOR 1			
Bifidobacterium adolescentis		3.80	0.15				
47531552	**Low discrimination**						**47531552**
Bifidobacterium adolescentis		78.72	0.49	XYL 87			
Bifidobacterium longum subsp. longum		20.15	0.37	SOR 11			
Clostridium clostridioforme		0.65	0.10				
47531560	**Low discrimination**						**47531560**
					GRA	CAT	
Bifidobacterium longum subsp. longum		67.42	0.43	SOR 1	99	1	
Clostridium perfringens		25.12	0.31		99	1	
Capnocytophaga ochracea		6.11	0.17		1	90	
47531562	**Low discrimination**						**47531562**
					GRA	CAT	
Bifidobacterium longum subsp. longum		44.32	0.43	SOR 1	99	1	
Capnocytophaga ochracea		36.12	0.34		1	90	
Parabacteroides distasonis		10.88	0.37		1	76	
47531570	**Good identification**						**47531570**
					GRA	CAT	
Bifidobacterium longum subsp. longum		99.41	0.47	SOR 1	99	1	
Capnocytophaga ochracea		0.59	0.01		1	90	
47531572	**Low discrimination**						**47531572**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Bifidobacterium longum subsp. longum		87.05	0.47	SOR 1	99	1	
Parabacteroides distasonis		6.76	0.33		1	76	
Capnocytophaga ochracea		4.63	0.17		1	90	
47531602	**Good identification**						**47531602**
Clostridium clostridioforme		98.88	0.39	SAL 88			
Parabacteroides distasonis		1.12	0.11	XYL 91			
47531612	**Good identification**						**47531612**
Clostridium clostridioforme		100.0	0.50	SAL 88			
47531622	**Low discrimination**						**47531622**
Parabacteroides distasonis		50.42	0.45	XYL 90			
Clostridium clostridioforme		44.89	0.39	LAC 99			
Bacteroides thetaiotaomicron		2.77	0.08				
47531632	**Low discrimination**						**47531632**
					GRA		
Clostridium clostridioforme		87.43	0.50	SAL 88	50		
Parabacteroides distasonis		11.61	0.41	XYL 91	1		
Bifidobacterium longum subsp. longum		0.50	0.13		99		
47531702	**Low discrimination**						**47531702**
					GRA		
Clostridium clostridioforme		65.80	0.48	SAL 88	50		
Parabacteroides distasonis		22.08	0.45	XYL 91	1		
Bifidobacterium longum subsp. longum		7.49	0.33		99		
47531710	**Low discrimination**						**47531710**
Bifidobacterium longum subsp. longum		80.82	0.37	RHA 1			
Clostridium clostridioforme		19.18	0.25				
47531712	**Good identification**						**47531712**
					GRA		
Clostridium clostridioforme		91.19	0.59	SAL 88	50		
Bifidobacterium longum subsp. longum		3.88	0.37	XYL 91	99		
Parabacteroides distasonis		3.62	0.41		1		
47531720	**Low discrimination**						**47531720**
					GRA		
Bacteroides vulgatus		86.47	0.47	TRE 4	1		
Bifidobacterium longum subsp. longum		7.40	0.43	XYL 98	99		
Parabacteroides distasonis		5.45	0.45		1		
47531722	**Good identification**						**47531722**
Parabacteroides distasonis		92.80	0.79	XYL 90			
Clostridium clostridioforme		2.79	0.48				
Bacteroides thetaiotaomicron		2.52	0.37				
47531723	**Acceptable identification**						**47531723**
Parabacteroides distasonis		93.37	0.45	XYL 90			
Clostridium clostridioforme		2.81	0.14	URE 1			
Bacteroides thetaiotaomicron		2.54	0.02				
47531730	**Low discrimination**						**47531730**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		73.20	0.47	RHA 1	99		
Parabacteroides distasonis		17.05	0.41		1		
Bacteroides vulgatus		5.07	0.13		1		
47531732	**Low discrimination**						**47531732**
					GRA		
Parabacteroides distasonis		75.88	0.75	XYL 90	1		
Clostridium clostridioforme		19.33	0.59		50		
Bifidobacterium longum subsp. longum		3.29	0.47		99		
47531733	**Low discrimination**						**47531733**
					GRA		
Parabacteroides distasonis		75.89	0.41	XYL 90	1		
Clostridium clostridioforme		19.33	0.25	URE 1	50		
Bifidobacterium longum subsp. longum		3.29	0.13		99		
47531762	**Acceptable identification**						**47531762**
Parabacteroides distasonis		92.41	0.45	XYL 90			
Clostridium clostridioforme		2.78	0.14	SOR 1			
Bacteroides thetaiotaomicron		2.51	0.02				
47531772	**Low discrimination**						**47531772**
					GRA		
Parabacteroides distasonis		75.89	0.41	XYL 90	1		
Clostridium clostridioforme		19.33	0.25	SOR 1	50		
Bifidobacterium longum subsp. longum		3.29	0.13		99		
47532000	**Very good identification**						**47532000**
					GRA	CAT	
Propionibacterium granulosum		97.89	0.58	ESL 1	99	90	
Actinomyces naeslundii		1.10	0.39		99	25	
Fusobacterium mortiferum		1.01	0.27		1	5	
47532001	**Low discrimination**						**47532001**
Actinomyces naeslundii		62.47	0.42	ESL 12			
Propionibacterium granulosum		37.53	0.24	GAL 96			
47532002	**Good identification**						**47532002**
					GRA	CAT	
Propionibacterium granulosum		95.29	0.48	ESL 1	99	90	
Actinomyces naeslundii		4.27	0.39		99	25	
Fusobacterium mortiferum		0.44	0.11		1	5	
47532003	**Low discrimination**						**47532003**
Actinomyces naeslundii		86.95	0.42	ESL 12			
Propionibacterium granulosum		13.05	0.13	GAL 96			
47532010	**Good identification**						**47532010**
Propionibacterium granulosum		100.0	0.43	ESL 1			
47532020	**Good identification**						**47532020**
Propionibacterium granulosum		95.71	0.48	ESL 1			
Actinomyces naeslundii		4.29	0.39				
47532021	**Low discrimination**						**47532021**
Actinomyces naeslundii		86.95	0.42	ESL 12			
Propionibacterium granulosum		13.05	0.13	GAL 96			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47532022	**Low discrimination**					**47532022**
Propionibacterium granulosum		84.79	0.37	ESL 1		
Actinomyces naeslundii		15.21	0.39			
47532023	**Good identification**					**47532023**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47532100	**Low discrimination**					**47532100**
					GRA	
Propionibacterium propionicum		48.61	0.42	NIT 86	99	
Actinomyces naeslundii		24.81	0.47	ESL 1	99	
Fusobacterium mortiferum		18.74	0.33		1	
47532101	**Low discrimination**					**47532101**
Actinomyces naeslundii		87.33	0.50	ESL 12		
Propionibacterium propionicum		12.67	0.26	GAL 96		
47532102	**Low discrimination**					**47532102**
					GRA	
Propionibacterium propionicum		70.22	0.45	NIT 86	99	
Actinomyces naeslundii		23.89	0.47	ESL 1	99	
Fusobacterium mortiferum		2.01	0.17		1	
47532103	**Low discrimination**					**47532103**
Actinomyces naeslundii		82.12	0.50	ESL 12		
Propionibacterium propionicum		17.88	0.29	GAL 96		
47532120	**Low discrimination**					**47532120**
					CAT	
Actinomyces naeslundii		80.42	0.47	ESL 12	25	
Clostridium perfringens		9.74	0.09	GAL 96	1	
Propionibacterium granulosum		6.36	0.13		90	
47532121	**Good identification**					**47532121**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47532122	**Acceptable identification**					**47532122**
					GRA	
Actinomyces naeslundii		91.11	0.47	ESL 12	99	
Clostridium perfringens		5.19	0.03	GAL 96	99	
Prevotella oralis		3.70	-.03		1	
47532123	**Good identification**					**47532123**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47532140	**Acceptable identification**					**47532140**
Propionibacterium propionicum		91.74	0.42	NIT 86		
Actinomyces naeslundii		8.26	0.34	ESL 1		
47532141	**Low discrimination**					**47532141**
Actinomyces naeslundii		54.87	0.37	ESL 12		
Propionibacterium propionicum		45.13	0.26	GAL 96		
47532142	**Acceptable identification**					**47532142**
Propionibacterium propionicum		94.34	0.45	NIT 86		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		5.67	0.34	ESL 1			
47532161	**Good identification**						**47532161**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47532163	**Good identification**						**47532163**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47533000	**Low discrimination**						**47533000**
					GRA	CAT	
Propionibacterium granulosum		76.98	0.58	ESL 1	99	90	
Actinomyces naeslundii		20.71	0.63		99	25	
Fusobacterium mortiferum		1.19	0.30		1	5	
47533001	**Very good identification**						**47533001**
Actinomyces naeslundii		97.56	0.66	ESL 12			
Propionibacterium granulosum		2.44	0.24				
47533002	**Low discrimination**						**47533002**
					CAT		
Actinomyces naeslundii		49.56	0.63	ESL 12	25		
Propionibacterium granulosum		46.05	0.48		90		
Actinomyces israelii		3.58	0.34		1		
47533003	**Very good identification**						**47533003**
Actinomyces naeslundii		99.38	0.66	ESL 12			
Propionibacterium granulosum		0.62	0.13				
47533010	**Low discrimination**						**47533010**
					CAT		
Propionibacterium granulosum		89.44	0.43	ESL 1	90		
Actinomyces naeslundii		9.29	0.41	MLZ 12	25		
Actinomyces israelii		1.27	0.17		1		
47533011	**Acceptable identification**						**47533011**
Actinomyces naeslundii		93.91	0.44	ESL 12			
Propionibacterium granulosum		6.09	0.09	MLZ 5			
47533013	**Good identification**						**47533013**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47533020	**Low discrimination**						**47533020**
					CAT		
Actinomyces naeslundii		46.99	0.63	ESL 12	25		
Propionibacterium granulosum		43.66	0.48		90		
Clostridium perfringens		9.11	0.28		1		
47533021	**Very good identification**						**47533021**
Actinomyces naeslundii		99.38	0.66	ESL 12			
Propionibacterium granulosum		0.62	0.13				
47533022	**Low discrimination**						**47533022**
					CAT		
Actinomyces naeslundii		73.61	0.63	ESL 12	25		
Propionibacterium granulosum		17.10	0.37		90		
Clostridium perfringens		6.72	0.23		1		
47533023	**Very good identification**						**47533023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.66	ESL 12		
47533031	**Good identification**					**47533031**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47533032	**Low discrimination**					**47533032**
Actinomyces naeslundii		62.43	0.41	ESL 12		
Propionibacterium granulosum		37.57	0.22	MLZ 5		
47533033	**Good identification**					**47533033**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47533040	**Low discrimination**					**47533040**
					CAT	
Actinomyces naeslundii		75.82	0.50	ESL 12	25	
Propionibacterium granulosum		16.13	0.24		90	
Actinomyces israelii		8.05	0.24		1	
47533041	**Good identification**					**47533041**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47533042	**Low discrimination**					**47533042**
					CAT	
Actinomyces naeslundii		82.48	0.50	ESL 12	25	
Actinomyces israelii		13.13	0.27		1	
Propionibacterium granulosum		4.39	0.13		90	
47533043	**Good identification**					**47533043**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47533060	**Low discrimination**					**47533060**
					CAT	
Actinomyces naeslundii		79.21	0.50	ESL 12	25	
Clostridium perfringens		16.58	0.16		1	
Propionibacterium granulosum		4.21	0.13		90	
47533061	**Good identification**					**47533061**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47533062	**Acceptable identification**					**47533062**
Actinomyces naeslundii		91.04	0.50	ESL 12		
Clostridium perfringens		8.97	0.11			
47533063	**Good identification**					**47533063**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47533100	**Low discrimination**					**47533100**
Actinomyces naeslundii		79.77	0.71	ESL 12		
Propionibacterium propionicum		6.51	0.42			
Actinomyces israelii		5.40	0.41			
47533101	**Very good identification**					**47533101**
					SPO	
Actinomyces naeslundii		98.83	0.74	ESL 12	1	
Propionibacterium propionicum		0.60	0.26		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		0.57	0.12		99	
47533102	**Low discrimination**					**47533102**
Actinomyces naeslundii		61.60	0.71	ESL 12		
Bifidobacterium adolescentis		21.18	0.47			
Propionibacterium propionicum		7.54	0.45			
47533103	**Very good identification**					**47533103**
Actinomyces naeslundii		98.78	0.74	ESL 12		
Propionibacterium propionicum		0.90	0.29			
Bifidobacterium adolescentis		0.23	0.13			
47533110	**Low discrimination**					**47533110**
Actinomyces naeslundii		74.03	0.49	ESL 12		
Actinomyces israelii		14.23	0.27	MLZ 5		
Bifidobacterium adolescentis		5.97	0.15			
47533111	**Good identification**					**47533111**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47533112	**Low discrimination**					**47533112**
Bifidobacterium adolescentis		83.41	0.49	XYL 87		
Actinomyces naeslundii		10.45	0.49	ARA 87		
Actinomyces israelii		3.01	0.30			
47533113	**Acceptable identification**					**47533113**
Actinomyces naeslundii		94.90	0.52	ESL 12		
Bifidobacterium adolescentis		5.10	0.15	MLZ 5		
47533120	**Low discrimination**					**47533120**
					GRA	
Actinomyces naeslundii		57.31	0.71	ESL 12	99	
Clostridium perfringens		28.63	0.43		99	
Prevotella melaninogenica		10.23	0.29		1	
47533121	**Very good identification**					**47533121**
					SPO	
Actinomyces naeslundii		99.36	0.74	ESL 12	1	
Clostridium perfringens		0.33	0.09		84	
Clostridium chauvoei		0.19	0.04		99	
47533122	**Low discrimination**					**47533122**
					GRA	
Actinomyces naeslundii		55.21	0.71	ESL 12	99	
Clostridium perfringens		12.98	0.38		99	
Prevotella oralis		9.25	0.32		1	
47533123	**Very good identification**					**47533123**
					GRA	
Actinomyces naeslundii		99.73	0.74	ESL 12	99	
Clostridium perfringens		0.16	0.03		99	
Prevotella oralis		0.11	-.03		1	
47533130	**Low discrimination**					**47533130**
Actinomyces naeslundii		79.34	0.49	ESL 12		
Clostridium perfringens		7.61	0.09	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		6.62	0.13			
47533131	**Good identification**					**47533131**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47533132	**Low discrimination**					**47533132**
					GRA	
Parabacteroides distasonis		35.04	0.36	MAN 1	1	
Actinomyces naeslundii		33.47	0.49	XYL 90	99	
Bifidobacterium breve		15.83	0.29		99	
47533133	**Good identification**					**47533133**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47533140	**Low discrimination**					**47533140**
Actinomyces naeslundii		61.41	0.58	ESL 12		
Propionibacterium propionicum		28.41	0.42			
Actinomyces israelii		9.16	0.34			
47533141	**Very good identification**					**47533141**
Actinomyces naeslundii		96.69	0.61	ESL 12		
Propionibacterium propionicum		3.31	0.26			
47533142	**Low discrimination**					**47533142**
Actinomyces naeslundii		44.69	0.58	ESL 12		
Propionibacterium propionicum		31.01	0.45			
Bifidobacterium adolescentis		10.76	0.32			
47533143	**Very good identification**					**47533143**
Actinomyces naeslundii		95.11	0.61	ESL 12		
Propionibacterium propionicum		4.89	0.29			
47533150	**Low discrimination**					**47533150**
Actinomyces naeslundii		70.25	0.36	ESL 12		
Actinomyces israelii		29.75	0.20	MLZ 5		
47533151	**Good identification**					**47533151**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47533153	**Good identification**					**47533153**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47533160	**Low discrimination**					**47533160**
Actinomyces naeslundii		63.58	0.58	ESL 12		
Clostridium perfringens		34.28	0.31			
Bifidobacterium breve		0.89	0.12			
47533161	**Very good identification**					**47533161**
Actinomyces naeslundii		100.0	0.61	ESL 12		
47533162	**Low discrimination**					**47533162**
Actinomyces naeslundii		67.29	0.58	ESL 12		
Clostridium perfringens		17.07	0.25			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		9.49	0.29				
47533163	**Very good identification**						**47533163**
Actinomyces naeslundii		100.0	0.61	ESL 12			
47533170	**Low discrimination**						**47533170**
Actinomyces naeslundii		79.05	0.36	ESL 12			
Bifidobacterium breve		20.95	0.12	MLZ 5			
47533171	**Good identification**						**47533171**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47533173	**Good identification**						**47533173**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47533300	**Low discrimination**						**47533300**
Actinomyces naeslundii		66.48	0.37	ESL 12			
Actinomyces israelii		33.52	0.22	RHA 1			
47533301	**Good identification**						**47533301**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47533302	**Low discrimination**						**47533302**
					GRA		
Actinomyces naeslundii		34.02	0.37	ESL 12	99		
Actinomyces israelii		25.74	0.25	RHA 1	99		
Prevotella oralis		12.73	0.04		1		
47533303	**Good identification**						**47533303**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47533321	**Good identification**						**47533321**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47533322	**Low discrimination**						**47533322**
					SPO		
Parabacteroides distasonis		55.87	0.49	MAN 1	1		
Clostridium ramosum		28.80	0.32	XYL 90	92		
Prevotella oralis		13.07	0.25		1		
47533323	**Low discrimination**						**47533323**
					SPO	GRA	
Actinomyces naeslundii		76.33	0.40	ESL 12	1	99	
Parabacteroides distasonis		15.62	0.15	RHA 1	1	1	
Clostridium ramosum		8.05	-.03		92	50	
47533332	**Good identification**						**47533332**
					SPO		
Parabacteroides distasonis		98.08	0.44	MAN 1	1		
Clostridium clostridioforme		0.98	0.04	XYL 90	66		
Clostridium ramosum		0.95	-.03		92		
47533501	**Good identification**						**47533501**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47533502	**Good identification**						**47533502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		93.53	0.61	XYL 87		
Actinomyces israelii		3.38	0.43			
Bifidobacterium longum subsp. longum		2.60	0.33			
47533510	**Low discrimination**					**47533510**
Bifidobacterium longum subsp. longum		74.78	0.37	MAN 1		
Bifidobacterium adolescentis		19.53	0.29			
Actinomyces israelii		5.69	0.26			
47533512	**Very good identification**					**47533512**
Bifidobacterium adolescentis		95.67	0.63	XYL 87		
Bifidobacterium longum subsp. longum		3.70	0.37			
Actinomyces israelii		0.42	0.29			
47533520	**Acceptable identification**					**47533520**
Bifidobacterium longum subsp. longum		90.78	0.43	MAN 1		
Actinomyces naeslundii		3.59	0.37			
Clostridium perfringens		1.79	0.09			
47533521	**Low discrimination**					**47533521**
Actinomyces naeslundii		85.45	0.40	ESL 12		
Bifidobacterium longum subsp. longum		14.55	0.09	ARA 1		
47533522	**Low discrimination**					**47533522**
					GRA	
Bifidobacterium longum subsp. longum		67.26	0.43	MAN 1	99	
Parabacteroides distasonis		16.51	0.37		1	
Bifidobacterium adolescentis		6.12	0.27		99	
47533523	**Low discrimination**					**47533523**
Actinomyces naeslundii		85.45	0.40	ESL 12		
Bifidobacterium longum subsp. longum		14.55	0.09	ARA 1		
47533530	**Good identification**					**47533530**
Bifidobacterium longum subsp. longum		100.0	0.47	MAN 1		
47533532	**Low discrimination**					**47533532**
					GRA	
Bifidobacterium longum subsp. longum		85.69	0.47	MAN 1	99	
Parabacteroides distasonis		6.65	0.33		1	
Bifidobacterium adolescentis		5.60	0.29		99	
47533542	**Low discrimination**					**47533542**
Bifidobacterium adolescentis		89.30	0.46	XYL 87		
Actinomyces israelii		10.14	0.36	SOR 11		
Actinomyces naeslundii		0.56	0.24			
47533552	**Good identification**					**47533552**
Bifidobacterium adolescentis		98.63	0.47	XYL 87		
Actinomyces israelii		1.37	0.22	SOR 11		
47533722	**Acceptable identification**					**47533722**
Parabacteroides distasonis		92.64	0.45	MAN 1		
Clostridium clostridioforme		2.79	0.14	XYL 90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		2.52	0.02				
47533732	**Low discrimination**						**47533732**
					GRA		
Parabacteroides distasonis		75.89	0.41	MAN 1	1		
Clostridium clostridioforme		19.33	0.25	XYL 90	50		
Bifidobacterium longum subsp. longum		3.29	0.13		99		
47534000	**Low discrimination**						**47534000**
					GRA	CAT	
Fusobacterium mortiferum		85.18	0.42	MLT 11	1	5	
Propionibacterium granulosum		14.82	0.26	XYL 7	99	90	
47534100	**Low discrimination**						**47534100**
					COC	GRA	
Fusobacterium mortiferum		85.08	0.48	MLT 11	1	1	
Atopobium parvulum		13.67	0.23	XYL 7	99	99	
Bifidobacterium longum subsp. longum		0.70	0.07		1	99	
47534122	**Low discrimination**						**47534122**
Parabacteroides distasonis		59.23	0.57	GAL 99			
Bacteroides fragilis		35.86	0.32				
Capnocytophaga ochracea		1.72	0.18				
47534132	**Low discrimination**						**47534132**
					GRA		
Parabacteroides distasonis		83.68	0.53	GAL 99	1		
Clostridium clostridioforme		12.65	0.33		50		
Bifidobacterium longum subsp. longum		2.22	0.21		99		
47534212	**Good identification**						**47534212**
Clostridium clostridioforme		100.0	0.39	SAL 88			
47534232	**Acceptable identification**						**47534232**
Clostridium clostridioforme		91.61	0.39	SAL 88			
Parabacteroides distasonis		8.39	0.27	ARA 94			
47534302	**Low discrimination**						**47534302**
Clostridium clostridioforme		81.21	0.36	SAL 88			
Parabacteroides distasonis		18.79	0.31	ARA 94			
47534312	**Good identification**						**47534312**
Clostridium clostridioforme		97.34	0.48	SAL 88			
Parabacteroides distasonis		2.66	0.27	ARA 94			
47534322	**Very good identification**						**47534322**
Parabacteroides distasonis		95.34	0.65	GAL 99			
Clostridium clostridioforme		4.16	0.36				
Prevotella oralis		0.31	0.10				
47534332	**Low discrimination**						**47534332**
Parabacteroides distasonis		73.02	0.61	GAL 99			
Clostridium clostridioforme		26.98	0.48				
47534412	**Good identification**						**47534412**
Clostridium clostridioforme		98.42	0.44	SAL 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		0.93	0.11	RHA 88		
Bifidobacterium adolescentis		0.66	0.10			
47534432	**Good identification**					**47534432**
					GRA	
Clostridium clostridioforme		95.57	0.44	SAL 88	50	
Bifidobacterium longum subsp. longum		3.59	0.21	RHA 88	99	
Parabacteroides distasonis		0.84	0.15		1	
47534500	**Acceptable identification**					**47534500**
					SPO	
Bifidobacterium longum subsp. longum		94.26	0.41	GAL 99	1	
Clostridium butyricum		1.53	-.03		99	
Clostridium clostridioforme		1.28	0.08		66	
47534502	**Low discrimination**					**47534502**
Clostridium clostridioforme		39.85	0.42	SAL 88		
Bifidobacterium longum subsp. longum		29.62	0.41	RHA 88		
Bifidobacterium adolescentis		29.39	0.43			
47534510	**Good identification**					**47534510**
Bifidobacterium longum subsp. longum		95.83	0.45	GAL 99		
Clostridium clostridioforme		3.48	0.19			
Bifidobacterium adolescentis		0.69	0.10			
47534512	**Low discrimination**					**47534512**
Clostridium clostridioforme		67.64	0.53	SAL 88		
Bifidobacterium longum subsp. longum		18.79	0.45	RHA 88		
Bifidobacterium adolescentis		13.39	0.44			
47534520	**Low discrimination**					**47534520**
					GRA	
Bifidobacterium longum subsp. longum		81.63	0.52	GAL 99	99	
Bacteroides vulgatus		17.49	0.26		1	
Parabacteroides distasonis		0.61	0.19		1	
47534522	**Low discrimination**					**47534522**
					GRA	
Bifidobacterium longum subsp. longum		47.55	0.52	GAL 99	99	
Parabacteroides distasonis		35.02	0.53		1	
Clostridium clostridioforme		15.99	0.42		50	
47534530	**Good identification**					**47534530**
					GRA	
Bifidobacterium longum subsp. longum		98.75	0.56	GAL 99	99	
Clostridium clostridioforme		0.90	0.19		50	
Parabacteroides distasonis		0.23	0.15		1	
47534532	**Low discrimination**					**47534532**
					GRA	
Bifidobacterium longum subsp. longum		46.79	0.56	GAL 99	99	
Clostridium clostridioforme		42.10	0.53		50	
Parabacteroides distasonis		10.90	0.49		1	
47534602	**Good identification**					**47534602**
Clostridium clostridioforme		100.0	0.48	SAL 88		
47534612	**Very good identification**					**47534612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.59	SAL 88			
47534622	**Good identification**						**47534622**
Clostridium clostridioforme		96.64	0.48	SAL 88			
Parabacteroides distasonis		2.93	0.27				
Bacteroides thetaiotaomicron		0.43	-.03				
47534632	**Very good identification**						**47534632**
Clostridium clostridioforme		99.64	0.59	SAL 88			
Parabacteroides distasonis		0.36	0.23				
47534702	**Very good identification**						**47534702**
					GRA		
Clostridium clostridioforme		98.90	0.57	SAL 88	50		
Parabacteroides distasonis		0.90	0.27		1		
Bifidobacterium adolescentis		0.10	0.09		99		
47534712	**Very good identification**						**47534712**
					GRA		
Clostridium clostridioforme		99.83	0.68	SAL 88	50		
Parabacteroides distasonis		0.11	0.23		1		
Bifidobacterium longum subsp. longum		0.04	0.11		99		
47534720	**Good identification**						**47534720**
Bacteroides vulgatus		97.11	0.42	TRE 4			
Clostridium clostridioforme		1.26	0.23	GAL 99			
Parabacteroides distasonis		1.12	0.27				
47534722	**Low discrimination**						**47534722**
Clostridium clostridioforme		50.69	0.57	SAL 88			
Parabacteroides distasonis		45.41	0.62				
Bacteroides thetaiotaomicron		3.29	0.26				
47534732	**Low discrimination**						**47534732**
Clostridium clostridioforme		89.85	0.68	SAL 88			
Parabacteroides distasonis		9.52	0.57				
Bacteroides thetaiotaomicron		0.50	0.19				
47535000	**Low discrimination**						**47535000**
					COC	GRA	
Fusobacterium mortiferum		48.17	0.45	MLT 11	1	1	
Eubacterium saburreum		20.57	0.37	XYL 7	1	85	
Atopobium parvulum		12.04	0.23		99	99	
47535002	**Low discrimination**						**47535002**
					GRA		
Actinomyces israelii		31.26	0.39	SAL 92	99		
Bifidobacterium adolescentis		17.24	0.29	CEL 93	99		
Fusobacterium mortiferum		10.16	0.29		1		
47535010	**Low discrimination**						**47535010**
Eubacterium saburreum		81.02	0.36	IND 99			
Actinomyces israelii		7.90	0.22				
Bifidobacterium longum subsp. longum		5.07	0.11				
47535022	**Low discrimination**						**47535022**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		54.17	0.57	LAC 99	76		
Bacteroides fragilis		32.80	0.32		99		
Prevotella oralis		7.70	0.30		1		
47535032	**Low discrimination**						**47535032**
					GRA		
Parabacteroides distasonis		84.55	0.53	LAC 99	1		
Clostridium clostridioforme		11.45	0.32		50		
Bifidobacterium longum subsp. longum		2.25	0.21		99		
47535100	**Low discrimination**						**47535100**
					COC	GRA	
Atopobium parvulum		78.50	0.57	RAF 1	99	99	
Fusobacterium mortiferum		7.40	0.51		1	1	
Bifidobacterium longum subsp. longum		4.03	0.41		1	99	
47535102	**Low discrimination**						**47535102**
					COC	GRA	
Bifidobacterium adolescentis		46.74	0.63	ARA 87	1	99	
Parabacteroides distasonis		15.31	0.57		1	1	
Atopobium parvulum		10.94	0.45		99	99	
47535110	**Low discrimination**						**47535110**
					COC		
Bifidobacterium longum subsp. longum		58.43	0.45	ARA 99	1		
Eubacterium saburreum		22.01	0.42		1		
Atopobium parvulum		6.75	0.23		99		
47535112	**Low discrimination**						**47535112**
					GRA		
Bifidobacterium adolescentis		74.99	0.64	ARA 87	99		
Parabacteroides distasonis		10.82	0.53		1		
Bifidobacterium longum subsp. longum		7.12	0.45		99		
47535120	**Low discrimination**						**47535120**
					CAT		
Bacteroides fragilis		38.11	0.44	TRE 1	99		
Bacteroides vulgatus		16.05	0.41	BGL 95	3		
Parabacteroides distasonis		12.08	0.57		76		
47535122	**Low discrimination**						**47535122**
Parabacteroides distasonis		59.76	0.91				
Bacteroides fragilis		36.18	0.66				
Capnocytophaga ochracea		1.74	0.53				
47535123	**Low discrimination**						**47535123**
Parabacteroides distasonis		59.56	0.57	URE 1			
Bacteroides fragilis		36.06	0.32				
Capnocytophaga ochracea		1.73	0.18				
47535130	**Low discrimination**						**47535130**
					GRA	CAT	
Bifidobacterium longum subsp. longum		68.22	0.56	ARA 99	99	1	
Parabacteroides distasonis		25.93	0.53		1	76	
Capnocytophaga ochracea		1.71	0.20		1	90	
47535132	**Very good identification**						**47535132**
					GRA	CAT	
Parabacteroides distasonis		95.24	0.87		1	76	
Bifidobacterium longum subsp. longum		2.53	0.56		99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides fragilis		1.08	0.32		1	99		
47535133	**Good identification**							**47535133**
					GRA	CAT		
Parabacteroides distasonis		95.37	0.53	URE 1	1	76		
Bifidobacterium longum subsp. longum		2.54	0.21		99	1		
Bacteroides fragilis		1.08	-.03		1	99		
47535140	**Low discrimination**							**47535140**
					COC	GRA	CAT	
Actinomyces israelii		48.05	0.40	SAL 92	1	99	1	
Atopobium parvulum		33.89	0.23	CEL 93	99	99	1	
Capnocytophaga ochracea		3.28	0.04		1	1	90	
47535142	**Low discrimination**							**47535142**
					GRA	CAT		
Bifidobacterium adolescentis		71.81	0.47	ARA 87	99	1		
Actinomyces israelii		16.56	0.43	SOR 11	99	1		
Capnocytophaga ochracea		6.77	0.20		1	90		
47535152	**Acceptable identification**							**47535152**
					GRA			
Bifidobacterium adolescentis		94.19	0.49	ARA 87	99			
Actinomyces israelii		2.66	0.29	SOR 11	99			
Parabacteroides distasonis		1.11	0.18		1			
47535162	**Low discrimination**							**47535162**
Parabacteroides distasonis		49.58	0.57	SOR 1				
Bacteroides fragilis		30.02	0.32					
Capnocytophaga ochracea		15.88	0.36					
47535172	**Low discrimination**							**47535172**
					GRA	CAT		
Parabacteroides distasonis		89.51	0.53	SOR 1	1	76		
Capnocytophaga ochracea		5.92	0.20		1	90		
Bifidobacterium longum subsp. longum		2.38	0.21		99	1		
47535202	**Low discrimination**							**47535202**
					GRA			
Clostridium clostridioforme		75.68	0.36	SAL 88	50			
Parabacteroides distasonis		19.54	0.31	ARA 94	1			
Actinomyces israelii		2.71	0.20		99			
47535212	**Good identification**							**47535212**
Clostridium clostridioforme		97.04	0.47	SAL 88				
Parabacteroides distasonis		2.96	0.27	ARA 94				
47535222	**Good identification**							**47535222**
Parabacteroides distasonis		94.13	0.65	LAC 99				
Clostridium clostridioforme		3.68	0.36					
Prevotella oralis		1.91	0.24					
47535232	**Low discrimination**							**47535232**
Parabacteroides distasonis		75.13	0.61	LAC 99				
Clostridium clostridioforme		24.87	0.47					
47535302	**Low discrimination**							**47535302**
Parabacteroides distasonis		85.25	0.65	NAG 99				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		11.17	0.45			
Prevotella oralis		1.43	0.22			
47535312	**Low discrimination**					**47535312**
					GRA	
Clostridium clostridioforme		51.90	0.56	SAL 88	50	
Parabacteroides distasonis		46.83	0.61	ARA 94	1	
Bifidobacterium adolescentis		1.09	0.30		99	
47535320	**Low discrimination**					**47535320**
Bacteroides vulgatus		79.29	0.57	TRE 4		
Parabacteroides distasonis		19.90	0.65	ARA 93		
Prevotella oralis		0.33	0.22			
47535322	**Excellent identification**					**47535322**
					CAT	
Parabacteroides distasonis		99.25	0.99		76	
Prevotella oralis		0.32	0.44		1	
Bacteroides fragilis		0.20	0.32		99	
47535323	**Very good identification**					**47535323**
					CAT	
Parabacteroides distasonis		99.35	0.65	URE 1	76	
Prevotella oralis		0.32	0.10		1	
Bacteroides fragilis		0.20	-.03		99	
47535330	**Good identification**					**47535330**
Parabacteroides distasonis		91.34	0.61	BGL 99		
Bacteroides vulgatus		6.83	0.23			
Clostridium clostridioforme		1.02	0.22			
47535332	**Very good identification**					**47535332**
Parabacteroides distasonis		98.84	0.95			
Clostridium clostridioforme		1.11	0.56			
Bacteroides thetaiotaomicron		0.03	0.19			
47535333	**Very good identification**					**47535333**
Parabacteroides distasonis		98.89	0.61	URE 1		
Clostridium clostridioforme		1.11	0.22			
47535362	**Very good identification**					**47535362**
					CAT	
Parabacteroides distasonis		99.24	0.65	SOR 1	76	
Prevotella oralis		0.32	0.10		1	
Bacteroides fragilis		0.20	-.03		99	
47535372	**Very good identification**					**47535372**
Parabacteroides distasonis		98.89	0.61	SOR 1		
Clostridium clostridioforme		1.11	0.22			
47535400	**Low discrimination**					**47535400**
					SPO	
Bifidobacterium longum subsp. longum		79.03	0.41	LAC 99	1	
Actinomyces israelii		11.59	0.35		1	
Clostridium butyricum		5.30	0.08		99	
47535402	**Low discrimination**					**47535402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		34.58	0.41	SAL 88		
Bifidobacterium longum subsp. longum		28.68	0.41	RHA 88		
Bifidobacterium adolescentis		28.46	0.43			
47535410	**Acceptable identification**					**47535410**
Bifidobacterium longum subsp. longum		95.00	0.45	LAC 99		
Clostridium clostridioforme		3.10	0.18			
Actinomyces israelii		1.22	0.21			
47535412	**Low discrimination**					**47535412**
Clostridium clostridioforme		64.94	0.52	SAL 88		
Bifidobacterium longum subsp. longum		20.14	0.45	RHA 88		
Bifidobacterium adolescentis		14.34	0.44			
47535420	**Low discrimination**					**47535420**
					GRA	
Bifidobacterium longum subsp. longum		81.50	0.52	LAC 99	99	
Bacteroides vulgatus		17.46	0.26		1	
Parabacteroides distasonis		0.61	0.19		1	
47535422	**Low discrimination**					**47535422**
					GRA	
Bifidobacterium longum subsp. longum		47.98	0.52	LAC 99	99	
Parabacteroides distasonis		35.35	0.53		1	
Clostridium clostridioforme		14.46	0.41		50	
47535430	**Good identification**					**47535430**
					GRA	
Bifidobacterium longum subsp. longum		98.84	0.56	LAC 99	99	
Clostridium clostridioforme		0.81	0.18		50	
Parabacteroides distasonis		0.23	0.15		1	
47535432	**Low discrimination**					**47535432**
					GRA	
Bifidobacterium longum subsp. longum		48.86	0.56	LAC 99	99	
Clostridium clostridioforme		39.40	0.52		50	
Parabacteroides distasonis		11.38	0.49		1	
47535500	**Very good identification**					**47535500**
					SPO	
Bifidobacterium longum subsp. longum		95.75	0.75		1	
Clostridium butyricum		1.56	0.32		99	
Bifidobacterium adolescentis		0.96	0.43		1	
47535501	**Good identification**					**47535501**
					SPO	
Bifidobacterium longum subsp. longum		96.62	0.41	URE 1	1	
Clostridium butyricum		1.57	-.03		99	
Bifidobacterium adolescentis		0.97	0.09		1	
47535502	**Low discrimination**					**47535502**
Bifidobacterium longum subsp. longum		48.03	0.75			
Bifidobacterium adolescentis		47.66	0.77			
Clostridium clostridioforme		1.96	0.50			
47535503	**Low discrimination**					**47535503**
Bifidobacterium longum subsp. longum		48.27	0.41	URE 1		
Bifidobacterium adolescentis		47.90	0.43			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		1.97	0.16				
47535510	**Very good identification**						**47535510**
Bifidobacterium longum subsp. longum		99.06	0.79				
Bifidobacterium adolescentis		0.71	0.44				
Clostridium clostridioforme		0.11	0.27				
47535511	**Good identification**						**47535511**
Bifidobacterium longum subsp. longum		99.29	0.45	URE 1			
Bifidobacterium adolescentis		0.71	0.10				
47535512	**Low discrimination**						**47535512**
Bifidobacterium longum subsp. longum		54.58	0.79				
Bifidobacterium adolescentis		38.88	0.79				
Clostridium clostridioforme		5.95	0.61				
47535513	**Low discrimination**						**47535513**
Bifidobacterium longum subsp. longum		54.62	0.45	URE 1			
Bifidobacterium adolescentis		38.91	0.44				
Clostridium clostridioforme		5.96	0.27				
47535520	**Low discrimination**						**47535520**
					GRA		
Bifidobacterium longum subsp. longum		81.73	0.86		99		
Bacteroides vulgatus		17.51	0.60		1		
Parabacteroides distasonis		0.61	0.53		1		
47535521	**Low discrimination**						**47535521**
					GRA		
Bifidobacterium longum subsp. longum		81.86	0.52	URE 1	99		
Bacteroides vulgatus		17.54	0.26		1		
Parabacteroides distasonis		0.61	0.19		1		
47535522	**Low discrimination**						**47535522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		56.25	0.86		99	1	
Parabacteroides distasonis		41.44	0.88		1	76	
Bacteroides thetaiotaomicron		0.68	0.41		1	89	
47535523	**Low discrimination**						**47535523**
					GRA	CAT	
Bifidobacterium longum subsp. longum		56.27	0.52	URE 1	99	1	
Parabacteroides distasonis		41.45	0.53		1	76	
Bacteroides thetaiotaomicron		0.68	0.07		1	89	
47535530	**Very good identification**						**47535530**
					GRA		
Bifidobacterium longum subsp. longum		99.60	0.90		99		
Parabacteroides distasonis		0.23	0.49		1		
Bacteroides vulgatus		0.13	0.26		1		
47535531	**Good identification**						**47535531**
					GRA		
Bifidobacterium longum subsp. longum		99.64	0.56	URE 1	99		
Parabacteroides distasonis		0.23	0.15		1		
Bacteroides vulgatus		0.13	-.08		1		
47535532	**Low discrimination**						**47535532**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		79.03	0.90		99		
Parabacteroides distasonis		18.41	0.83		1		
Clostridium clostridioforme		2.16	0.61		50		
47535533	**Low discrimination**						**47535533**
					GRA		
Bifidobacterium longum subsp. longum		79.07	0.56	URE 1	99		
Parabacteroides distasonis		18.42	0.49		1		
Clostridium clostridioforme		2.16	0.27		50		
47535540	**Low discrimination**						**47535540**
					SPO		
Bifidobacterium longum subsp. longum		66.69	0.41	SOR 1	1		
Actinomyces israelii		15.21	0.38		1		
Clostridium butyricum		9.34	0.14		99		
47535542	**Low discrimination**						**47535542**
Bifidobacterium adolescentis		89.02	0.62	SOR 11			
Bifidobacterium longum subsp. longum		7.33	0.41				
Actinomyces israelii		2.51	0.41				
47535550	**Low discrimination**						**47535550**
Bifidobacterium longum subsp. longum		89.58	0.45	SOR 1			
Bifidobacterium adolescentis		7.89	0.29				
Actinomyces israelii		1.79	0.24				
47535552	**Low discrimination**						**47535552**
Bifidobacterium adolescentis		88.36	0.63	SOR 11			
Bifidobacterium longum subsp. longum		10.14	0.45				
Clostridium clostridioforme		1.11	0.27				
47535560	**Low discrimination**						**47535560**
					GRA	CAT	
Bifidobacterium longum subsp. longum		80.91	0.52	SOR 1	99	1	
Bacteroides vulgatus		17.33	0.26		1	3	
Capnocytophaga ochracea		0.61	0.07		1	90	
47535562	**Low discrimination**						**47535562**
					GRA	CAT	
Bifidobacterium longum subsp. longum		53.41	0.52	SOR 1	99	1	
Parabacteroides distasonis		39.34	0.53		1	76	
Capnocytophaga ochracea		3.63	0.23		1	90	
47535570	**Good identification**						**47535570**
					GRA		
Bifidobacterium longum subsp. longum		99.64	0.56	SOR 1	99		
Parabacteroides distasonis		0.23	0.15		1		
Bacteroides vulgatus		0.13	-.08		1		
47535572	**Low discrimination**						**47535572**
					GRA		
Bifidobacterium longum subsp. longum		77.56	0.56	SOR 1	99		
Parabacteroides distasonis		18.07	0.49		1		
Clostridium clostridioforme		2.12	0.27		50		
47535602	**Good identification**						**47535602**
					GRA		
Clostridium clostridioforme		98.45	0.56	SAL 88	50		
Parabacteroides distasonis		0.99	0.27		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		0.18	0.19		99	
47535612	**Very good identification**					**47535612**
					GRA	
Clostridium clostridioforme		99.81	0.67	SAL 88	50	
Parabacteroides distasonis		0.12	0.23		1	
Bifidobacterium longum subsp. longum		0.04	0.11		99	
47535620	**Good identification**					**47535620**
Bacteroides vulgatus		97.08	0.42	TRE 4		
Parabacteroides distasonis		1.12	0.27	LAC 99		
Clostridium clostridioforme		1.12	0.22			
47535622	**Low discrimination**					**47535622**
Clostridium clostridioforme		46.30	0.56	SAL 88		
Parabacteroides distasonis		46.30	0.62			
Bacteroides thetaiotaomicron		6.78	0.31			
47535632	**Low discrimination**					**47535632**
Clostridium clostridioforme		88.31	0.67	SAL 88		
Parabacteroides distasonis		10.44	0.57			
Bacteroides thetaiotaomicron		1.10	0.24			
47535700	**Low discrimination**					**47535700**
					GRA	
Bacteroides vulgatus		81.92	0.42	TRE 4	1	
Bifidobacterium longum subsp. longum		10.62	0.41	NAG 99	99	
Clostridium clostridioforme		3.18	0.31		50	
47535702	**Low discrimination**					**47535702**
					GRA	
Clostridium clostridioforme		71.72	0.65	SAL 88	50	
Parabacteroides distasonis		21.42	0.62		1	
Bifidobacterium longum subsp. longum		2.42	0.41		99	
47535710	**Low discrimination**					**47535710**
					GRA	
Bifidobacterium longum subsp. longum		53.16	0.45	RHA 1	99	
Clostridium clostridioforme		42.52	0.42		50	
Bacteroides vulgatus		2.43	0.08		1	
47535712	**Good identification**					**47535712**
					GRA	
Clostridium clostridioforme		94.44	0.76	SAL 88	50	
Parabacteroides distasonis		3.33	0.57		1	
Bifidobacterium longum subsp. longum		1.19	0.45		99	
47535713	**Acceptable identification**					**47535713**
					GRA	
Clostridium clostridioforme		94.61	0.42	SAL 88	50	
Parabacteroides distasonis		3.34	0.23	URE 1	1	
Bifidobacterium longum subsp. longum		1.20	0.11		99	
47535720	**Very good identification**					**47535720**
					GRA	
Bacteroides vulgatus		98.22	0.76	TRE 4	1	
Parabacteroides distasonis		1.14	0.62		1	
Bifidobacterium longum subsp. longum		0.52	0.52		99	
47535721	**Good identification**					**47535721**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bacteroides vulgatus		98.35	0.42	TRE 4	1	
Parabacteroides distasonis		1.14	0.27	URE 1	1	
Bifidobacterium longum subsp. longum		0.52	0.17		99	
47535722	**Low discrimination**					**47535722**
Parabacteroides distasonis		89.25	0.96			
Bacteroides thetaiotaomicron		6.47	0.60			
Clostridium clostridioforme		3.02	0.65			
47535723	**Low discrimination**					**47535723**
Parabacteroides distasonis		89.31	0.62	URE 1		
Bacteroides thetaiotaomicron		6.47	0.26			
Clostridium clostridioforme		3.02	0.31			
47535730	**Low discrimination**					**47535730**
					GRA	
Bacteroides vulgatus		36.91	0.42	TRE 4	1	
Bifidobacterium longum subsp. longum		32.60	0.56	MLZ 1	99	
Parabacteroides distasonis		22.78	0.57		1	
47535732	**Low discrimination**					**47535732**
Parabacteroides distasonis		73.86	0.91			
Clostridium clostridioforme		21.13	0.76			
Bacteroides thetaiotaomicron		3.87	0.53			
47535733	**Low discrimination**					**47535733**
Parabacteroides distasonis		73.91	0.57	URE 1		
Clostridium clostridioforme		21.15	0.42			
Bacteroides thetaiotaomicron		3.87	0.19			
47535752	**Low discrimination**					**47535752**
					GRA	
Clostridium clostridioforme		86.36	0.42	SAL 88	50	
Bifidobacterium adolescentis		9.51	0.29	SOR 1	99	
Parabacteroides distasonis		3.05	0.23		1	
47535760	**Good identification**					**47535760**
					GRA	
Bacteroides vulgatus		98.35	0.42	TRE 4	1	
Parabacteroides distasonis		1.14	0.27	SOR 1	1	
Bifidobacterium longum subsp. longum		0.52	0.17		99	
47535762	**Low discrimination**					**47535762**
Parabacteroides distasonis		89.10	0.62	SOR 1		
Bacteroides thetaiotaomicron		6.46	0.26			
Clostridium clostridioforme		3.01	0.31			
47535772	**Low discrimination**					**47535772**
Parabacteroides distasonis		73.73	0.57	SOR 1		
Clostridium clostridioforme		21.10	0.42			
Bacteroides thetaiotaomicron		3.86	0.19			
47536502	**Acceptable identification**					**47536502**
Bifidobacterium adolescentis		91.64	0.41	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		5.69	0.27			
Clostridium clostridioforme		1.53	0.08			
47536512	**Acceptable identification**					**47536512**
Bifidobacterium adolescentis		91.41	0.43	GAL 99		
Clostridium clostridioforme		5.70	0.19			
Bifidobacterium longum subsp. longum		1.58	0.11			
47537000	**Acceptable identification**					**47537000**
					CAT	
Actinomyces israelii		91.42	0.49	SAL 92	1	
Propionibacterium granulosum		6.20	0.24	CEL 93	90	
Actinomyces naeslundii		1.67	0.29		25	
47537002	**Acceptable identification**					**47537002**
Actinomyces israelii		90.66	0.52	SAL 92		
Bifidobacterium adolescentis		7.22	0.27	CEL 93		
Actinomyces naeslundii		1.10	0.29			
47537010	**Acceptable identification**					**47537010**
					CAT	
Actinomyces israelii		94.18	0.35	SAL 92	1	
Propionibacterium granulosum		5.82	0.09	CEL 93	90	
47537012	**Low discrimination**					**47537012**
Actinomyces israelii		60.56	0.38	SAL 92		
Bifidobacterium adolescentis		39.44	0.29	CEL 93		
47537040	**Good identification**					**47537040**
Actinomyces israelii		100.0	0.42	SAL 92		
47537042	**Good identification**					**47537042**
Actinomyces israelii		97.53	0.45	SAL 92		
Bifidobacterium adolescentis		2.47	0.12	CEL 93		
47537100	**Good identification**					**47537100**
					COC	
Actinomyces israelii		90.49	0.60	SAL 92	1	
Atopobium parvulum		4.38	0.23	CEL 93	99	
Bifidobacterium adolescentis		2.70	0.27		1	
47537101	**Low discrimination**					**47537101**
Actinomyces naeslundii		65.83	0.40	ESL 12		
Actinomyces israelii		34.17	0.25	XYL 1		
47537102	**Low discrimination**					**47537102**
Bifidobacterium adolescentis		65.64	0.61	ARA 87		
Actinomyces israelii		33.31	0.63			
Actinomyces naeslundii		0.29	0.37			
47537110	**Low discrimination**					**47537110**
Actinomyces israelii		78.58	0.46	SAL 92		
Bifidobacterium adolescentis		19.19	0.29	CEL 93		
Bifidobacterium longum subsp. longum		2.23	0.11	MLZ 13		
47537112	**Good identification**					**47537112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium adolescentis		93.74	0.63	ARA 87	99		
Actinomyces israelii		5.82	0.49		99		
Parabacteroides distasonis		0.17	0.18		1		
47537120	**Low discrimination**						**47537120**
					GRA	CAT	
Actinomyces israelii		36.80	0.39	SAL 92	99	1	
Bacteroides fragilis		21.74	0.10	CEL 93	1	99	
Bacteroides vulgatus		9.15	0.07	NAG 6	1	3	
47537121	**Good identification**						**47537121**
Actinomyces naeslundii		100.0	0.40	ESL 12			
47537122	**Low discrimination**						**47537122**
					GRA	CAT	
Parabacteroides distasonis		55.94	0.57	MAN 1	1	76	
Bacteroides fragilis		33.87	0.32		1	99	
Actinomyces israelii		4.53	0.42		99	1	
47537123	**Low discrimination**						**47537123**
					GRA		
Actinomyces naeslundii		50.36	0.40	ESL 12	99		
Parabacteroides distasonis		30.92	0.23	XYL 1	1		
Bacteroides fragilis		18.72	-.03		1		
47537132	**Low discrimination**						**47537132**
					GRA		
Parabacteroides distasonis		87.87	0.53	MAN 1	1		
Bifidobacterium adolescentis		5.03	0.29		99		
Bifidobacterium longum subsp. longum		2.34	0.21		99		
47537140	**Good identification**						**47537140**
Actinomyces israelii		98.49	0.53	SAL 92			
Bifidobacterium adolescentis		0.94	0.12	CEL 93			
Actinomyces naeslundii		0.58	0.24				
47537142	**Low discrimination**						**47537142**
Actinomyces israelii		61.34	0.56	SAL 92			
Bifidobacterium adolescentis		38.42	0.46	CEL 93			
Actinomyces naeslundii		0.24	0.24				
47537150	**Low discrimination**						**47537150**
					COC		
Actinomyces israelii		87.96	0.38	SAL 92	1		
Bifidobacterium adolescentis		6.83	0.13	CEL 93	1		
Blautia producta		5.21	-.03	MLZ 13	99		
47537152	**Low discrimination**						**47537152**
Bifidobacterium adolescentis		83.67	0.47	ARA 87			
Actinomyces israelii		16.33	0.42	SOR 11			
47537162	**Low discrimination**						**47537162**
					GRA	CAT	
Actinomyces israelii		56.22	0.35	SAL 92	99	1	
Parabacteroides distasonis		18.05	0.23	CEL 93	1	76	
Bacteroides fragilis		10.93	-.03	NAG 6	1	99	
47537300	**Good identification**						**47537300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.40	SAL 92		
47537302	**Low discrimination**					**47537302**
					GRA	
Actinomyces israelii		61.59	0.43	SAL 92	99	
Parabacteroides distasonis		19.56	0.31	CEL 93	1	
Bifidobacterium adolescentis		16.29	0.27	RHA 7	99	
47537322	**Very good identification**					**47537322**
					GRA	
Parabacteroides distasonis		99.09	0.65	MAN 1	1	
Prevotella oralis		0.32	0.10		1	
Actinomyces israelii		0.20	0.23		99	
47537332	**Very good identification**					**47537332**
Parabacteroides distasonis		98.89	0.61	MAN 1		
Clostridium clostridioforme		1.11	0.22			
47537342	**Acceptable identification**					**47537342**
Actinomyces israelii		92.25	0.36	SAL 92		
Bifidobacterium adolescentis		7.75	0.12	CEL 93		
47537400	**Good identification**					**47537400**
Actinomyces israelii		97.85	0.48	SAL 92		
Bifidobacterium longum subsp. longum		1.19	0.07	CEL 93		
Bifidobacterium adolescentis		0.97	0.07			
47537402	**Low discrimination**					**47537402**
Actinomyces israelii		59.91	0.51	SAL 92		
Bifidobacterium adolescentis		39.02	0.41	CEL 93		
Clostridium clostridioforme		0.59	0.07			
47537412	**Low discrimination**					**47537412**
Bifidobacterium adolescentis		78.42	0.43	LAC 99		
Actinomyces israelii		14.72	0.37			
Clostridium clostridioforme		4.38	0.18			
47537440	**Good identification**					**47537440**
Actinomyces israelii		100.0	0.41	SAL 92		
47537442	**Low discrimination**					**47537442**
Actinomyces israelii		82.85	0.44	SAL 92		
Bifidobacterium adolescentis		17.15	0.26	CEL 93		
47537500	**Low discrimination**					**47537500**
Actinomyces israelii		62.44	0.58	SAL 92		
Bifidobacterium longum subsp. longum		18.78	0.41	CEL 93		
Bifidobacterium adolescentis		15.25	0.41			
47537502	**Good identification**					**47537502**
Bifidobacterium adolescentis		92.90	0.76			
Actinomyces israelii		5.76	0.61			
Bifidobacterium longum subsp. longum		1.16	0.41			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47537503	**Acceptable identification**					**47537503**
Bifidobacterium adolescentis		93.07	0.41	URE 1		
Actinomyces israelii		5.77	0.27			
Bifidobacterium longum subsp. longum		1.16	0.07			
47537510	**Low discrimination**					**47537510**
Bifidobacterium longum subsp. longum		50.77	0.45	MAN 1		
Bifidobacterium adolescentis		29.59	0.43			
Actinomyces israelii		14.81	0.44			
47537512	**Very good identification**					**47537512**
Bifidobacterium adolescentis		96.04	0.77			
Bifidobacterium longum subsp. longum		1.67	0.45			
Bifidobacterium dentium		1.37	0.27			
47537513	**Good identification**					**47537513**
Bifidobacterium adolescentis		96.93	0.43	URE 1		
Bifidobacterium longum subsp. longum		1.68	0.11			
Bifidobacterium dentium		1.39	-.07			
47537520	**Low discrimination**					**47537520**
					GRA	
Bifidobacterium longum subsp. longum		78.09	0.52	MAN 1	99	
Bacteroides vulgatus		16.73	0.26		1	
Actinomyces israelii		4.14	0.38		99	
47537522	**Low discrimination**					**47537522**
					GRA	
Bifidobacterium longum subsp. longum		48.41	0.52	MAN 1	99	
Parabacteroides distasonis		35.66	0.53		1	
Bifidobacterium adolescentis		9.83	0.41		99	
47537530	**Good identification**					**47537530**
Bifidobacterium longum subsp. longum		95.60	0.56	MAN 1		
Bifidobacterium dentium		3.47	0.15			
Actinomyces israelii		0.45	0.24			
47537532	**Low discrimination**					**47537532**
					GRA	
Bifidobacterium longum subsp. longum		56.11	0.56	MAN 1	99	
Bifidobacterium dentium		20.57	0.32		99	
Parabacteroides distasonis		13.07	0.49		1	
47537540	**Low discrimination**					**47537540**
					COC	
Actinomyces israelii		68.76	0.51	SAL 92	1	
Blautia producta		24.54	0.24	CEL 93	99	
Bifidobacterium adolescentis		5.34	0.26		1	
47537542	**Low discrimination**					**47537542**
					COC	
Bifidobacterium adolescentis		83.24	0.60	SOR 11	1	
Actinomyces israelii		16.25	0.54		1	
Blautia producta		0.43	0.07		99	
47537552	**Very good identification**					**47537552**
					COC	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		96.24	0.62	SOR 11	1	
Actinomyces israelii		2.30	0.40		1	
Blautia producta		1.22	0.15		99	
47537700	**Low discrimination**					**47537700**
					GRA	
Actinomyces israelii		72.23	0.39	SAL 92	99	
Bacteroides vulgatus		22.49	0.08	CEL 93	1	
Bifidobacterium longum subsp. longum		2.92	0.07	RHA 7	99	
47537702	**Low discrimination**					**47537702**
Bifidobacterium adolescentis		51.21	0.41	RHA 1		
Actinomyces israelii		23.67	0.42			
Clostridium clostridioforme		18.86	0.31			
47537712	**Low discrimination**					**47537712**
Clostridium clostridioforme		54.70	0.42	SAL 88		
Bifidobacterium adolescentis		39.86	0.43	MAN 1		
Actinomyces israelii		2.25	0.28			
47537720	**Good identification**					**47537720**
					GRA	
Bacteroides vulgatus		98.15	0.42	TRE 4	1	
Parabacteroides distasonis		1.14	0.27	MAN 1	1	
Bifidobacterium longum subsp. longum		0.51	0.17		99	
47537722	**Low discrimination**					**47537722**
Parabacteroides distasonis		87.37	0.62	MAN 1		
Bacteroides thetaiotaomicron		6.33	0.26			
Clostridium clostridioforme		2.96	0.31			
47537732	**Low discrimination**					**47537732**
Parabacteroides distasonis		72.12	0.57	MAN 1		
Clostridium clostridioforme		20.64	0.42			
Bacteroides thetaiotaomicron		3.78	0.19			
47541001	**Good identification**					**47541001**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47541003	**Good identification**					**47541003**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47541021	**Low discrimination**					**47541021**
					GRA	
Actinomyces naeslundii		81.57	0.36	ESL 12	99	
Prevotella melaninogenica		18.43	-.04	FRU 95	1	
47541022	**Low discrimination**					**47541022**
					CAT	
Prevotella oralis		88.49	0.41	FRU 99	1	
Prevotella melaninogenica		6.19	0.18	LAC 94	1	
Capnocytophaga ochracea		3.99	0.18		90	
47541023	**Low discrimination**					**47541023**
					GRA	
Actinomyces naeslundii		61.92	0.36	ESL 12	99	
Prevotella oralis		38.08	0.07	FRU 95	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47541102	**Low discrimination**						**47541102**
					CAT		
Prevotella oralis		54.88	0.40	FRU 99	1		
Capnocytophaga ochracea		33.04	0.36	NAG 95	90		
Prevotella melaninogenica		4.66	0.18		1		
47541103	**Low discrimination**						**47541103**
					GRA	SPO	
Actinomyces naeslundii		66.76	0.44	ESL 12	99	1	
Clostridium chauvoei		14.19	0.10	FRU 95	99	99	
Prevotella oralis		11.89	0.06		1	1	
47541120	**Very good identification**						**47541120**
					CAT		
Prevotella melaninogenica		96.17	0.64	FRU 90	1		
Prevotella oralis		2.18	0.40	CEL 10	1		
Capnocytophaga ochracea		1.31	0.36		90		
47541122	**Low discrimination**						**47541122**
					CAT		
Prevotella oralis		56.72	0.62	FRU 99	1		
Prevotella melaninogenica		25.08	0.52		1		
Capnocytophaga ochracea		16.18	0.53		90		
47541132	**Low discrimination**						**47541132**
					CAT		
Capnocytophaga ochracea		61.61	0.36	FRU 99	90		
Prevotella oralis		19.64	0.27	MLZ 10	1		
Prevotella melaninogenica		8.68	0.18		1		
47541162	**Low discrimination**						**47541162**
					CAT		
Capnocytophaga ochracea		62.28	0.36	FRU 99	90		
Prevotella oralis		19.86	0.27	SOR 10	1		
Prevotella melaninogenica		8.78	0.18		1		
47541322	**Good identification**						**47541322**
Prevotella oralis		97.34	0.55	FRU 99			
Prevotella melaninogenica		1.01	0.18				
Parabacteroides distasonis		0.89	0.33				
47541502	**Low discrimination**						**47541502**
					GRA	CAT	
Bifidobacterium adolescentis		51.47	0.37	FRU 99	99	1	
Capnocytophaga ochracea		29.59	0.23	XYL 87	1	90	
Bifidobacterium longum subsp. longum		16.13	0.27		99	1	
47541512	**Low discrimination**						**47541512**
					GRA	CAT	
Bifidobacterium adolescentis		66.51	0.39	FRU 99	99	1	
Bifidobacterium longum subsp. longum		29.05	0.31	XYL 87	99	1	
Capnocytophaga ochracea		3.48	0.07		1	90	
47541522	**Low discrimination**						**47541522**
					GRA	CAT	
Capnocytophaga ochracea		64.40	0.40	FRU 99	1	90	
Bifidobacterium longum subsp. longum		15.61	0.37		99	1	
Prevotella oralis		12.93	0.27		1	1	
47541530	**Good identification**						**47541530**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		96.03	0.41	FRU 90	99	1	
Capnocytophaga ochracea		2.87	0.07	CEL 1	1	90	
Prevotella melaninogenica		1.10	-.04		1	1	
47541532	**Low discrimination**						**47541532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		77.03	0.41	FRU 90	99	1	
Capnocytophaga ochracea		20.74	0.23	CEL 1	1	90	
Parabacteroides distasonis		1.59	0.17		1	76	
47543000	**Good identification**						**47543000**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47543001	**Good identification**						**47543001**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47543002	**Good identification**						**47543002**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47543003	**Good identification**						**47543003**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47543020	**Low discrimination**						**47543020**
					GRA		
Actinomyces naeslundii		85.14	0.38	ESL 12	99		
Prevotella melaninogenica		14.86	-.04	FRU 95	1		
47543021	**Good identification**						**47543021**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47543022	**Low discrimination**						**47543022**
					GRA		
Actinomyces naeslundii		67.78	0.38	ESL 12	99		
Prevotella oralis		32.22	0.07	FRU 95	1		
47543023	**Good identification**						**47543023**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47543100	**Low discrimination**						**47543100**
					GRA		
Actinomyces naeslundii		84.29	0.46	ESL 12	99		
Actinomyces israelii		5.66	0.17	FRU 95	99		
Prevotella melaninogenica		5.17	-.04		1		
47543101	**Good identification**						**47543101**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47543102	**Low discrimination**						**47543102**
Actinomyces naeslundii		45.17	0.46	ESL 12			
Bifidobacterium breve		26.40	0.27	FRU 95			
Bifidobacterium adolescentis		13.91	0.22				
47543103	**Good identification**						**47543103**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47543121	**Good identification**						**47543121**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		96.07	0.49	ESL 12	99		
Prevotella melaninogenica		3.93	-.04	FRU 95	1		
47543123	**Good identification**						**47543123**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47543141	**Good identification**						**47543141**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47543143	**Good identification**						**47543143**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47543161	**Good identification**						**47543161**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47543163	**Good identification**						**47543163**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47543502	**Good identification**						**47543502**
Bifidobacterium adolescentis		96.15	0.36	FRU 99			
Actinomyces israelii		3.85	0.18	XYL 87			
47543512	**Good identification**						**47543512**
Bifidobacterium adolescentis		100.0	0.37	FRU 99			
47545112	**Acceptable identification**						**47545112**
					GRA	CAT	
Bifidobacterium adolescentis		91.25	0.39	FRU 99	99	1	
Capnocytophaga ochracea		6.76	0.10	ARA 87	1	90	
Bacteroides fragilis		1.99	-.09		1	99	
47545120	**Low discrimination**						**47545120**
					CAT		
Bacteroides fragilis		77.02	0.38	FRU 99	99		
Prevotella melaninogenica		13.50	0.30	BGL 95	1		
Capnocytophaga ochracea		4.56	0.26		90		
47545122	**Good identification**						**47545122**
					CAT		
Bacteroides fragilis		92.30	0.60	FRU 99	99		
Prevotella oralis		4.49	0.46		1		
Capnocytophaga ochracea		2.59	0.42		90		
47545322	**Low discrimination**						**47545322**
					CAT		
Prevotella oralis		45.96	0.40	FRU 99	1		
Parabacteroides distasonis		30.44	0.49	XYL 11	76		
Bacteroides fragilis		22.28	0.25		99		
47545332	**Acceptable identification**						**47545332**
Parabacteroides distasonis		92.14	0.45	FRU 99			
Clostridium clostridioforme		3.98	0.16				
Prevotella oralis		2.61	0.06				
47545500	**Low discrimination**						**47545500**
					SPO		
Bifidobacterium longum subsp. longum		72.73	0.35	FRU 90	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		16.83	0.11	CEL 1	99		
Bifidobacterium adolescentis		5.23	0.17		1		
47545502	**Low discrimination**						**47545502**
					GRA	CAT	
Bifidobacterium adolescentis		83.72	0.52	FRU 99	99	1	
Bifidobacterium longum subsp. longum		11.76	0.35		99	1	
Capnocytophaga ochracea		1.80	0.13		1	90	
47545510	**Good identification**						**47545510**
Bifidobacterium longum subsp. longum		95.09	0.39	FRU 90			
Bifidobacterium adolescentis		4.91	0.19	CEL 1			
47545512	**Low discrimination**						**47545512**
Bifidobacterium adolescentis		82.12	0.53	FRU 99			
Bifidobacterium longum subsp. longum		16.08	0.39				
Clostridium clostridioforme		1.80	0.21				
47545520	**Low discrimination**						**47545520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		58.98	0.46	FRU 90	99	1	
Bacteroides vulgatus		35.91	0.28	CEL 1	1	3	
Capnocytophaga ochracea		2.25	0.13		1	90	
47545522	**Low discrimination**						**47545522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		40.79	0.46	FRU 90	99	1	
Bacteroides fragilis		28.62	0.25	CEL 1	1	99	
Capnocytophaga ochracea		14.03	0.29		1	90	
47545530	**Good identification**						**47545530**
					GRA		
Bifidobacterium longum subsp. longum		99.64	0.50	FRU 90	99		
Bacteroides vulgatus		0.36	-.07	CEL 1	1		
47545532	**Low discrimination**						**47545532**
					GRA		
Bifidobacterium longum subsp. longum		88.54	0.50	FRU 90	99		
Parabacteroides distasonis		5.49	0.33	CEL 1	1		
Clostridium clostridioforme		2.48	0.21		50		
47545542	**Good identification**						**47545542**
Bifidobacterium adolescentis		96.97	0.36	FRU 99			
Actinomyces israelii		3.03	0.16	SOR 11			
47545552	**Good identification**						**47545552**
Bifidobacterium adolescentis		100.0	0.37	FRU 99			
47545712	**Acceptable identification**						**47545712**
Clostridium clostridioforme		94.09	0.36	SAL 88			
Bifidobacterium adolescentis		5.91	0.19	FRU 99			
47545720	**Good identification**						**47545720**
					GRA		
Bacteroides vulgatus		99.71	0.44	FRU 99	1		
Bifidobacterium longum subsp. longum		0.18	0.11	CEL 5	99		
Parabacteroides distasonis		0.11	0.12		1		
47545722	**Low discrimination**						**47545722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		55.81	0.46	FRU 99			
Bacteroides uniformis		18.55	0.19				
Clostridium clostridioforme		7.29	0.25				
47545732	**Low discrimination**						**47545732**
Clostridium clostridioforme		48.13	0.36	SAL 88			
Parabacteroides distasonis		43.55	0.41	FRU 99			
Bacteroides thetaiotaomicron		3.08	0.05	TRE 88			
47547102	**Low discrimination**						**47547102**
					CAT	GRA	
Bifidobacterium adolescentis		62.70	0.36	FRU 99	1	99	
Actinomyces israelii		35.25	0.38	ARA 87	1	99	
Bacteroides fragilis		2.04	-.09		99	1	
47547112	**Acceptable identification**						**47547112**
Bifidobacterium adolescentis		93.57	0.37	FRU 99			
Actinomyces israelii		6.43	0.24	ARA 87			
47547502	**Acceptable identification**						**47547502**
Bifidobacterium adolescentis		93.57	0.50	FRU 99			
Actinomyces israelii		6.43	0.36				
47547512	**Good identification**						**47547512**
Bifidobacterium adolescentis		98.73	0.52	FRU 99			
Actinomyces israelii		0.83	0.22				
Bifidobacterium dentium		0.44	-.07				
47547552	**Good identification**						**47547552**
Bifidobacterium adolescentis		97.42	0.36	FRU 99			
Actinomyces israelii		2.58	0.15	SOR 11			
47550000	**Very good identification**						**47550000**
					GRA	CAT	
Fusobacterium mortiferum		98.26	0.62	MLT 11	1	5	
Prevotella intermedia		0.80	0.15		1	1	
Propionibacterium granulosum		0.38	0.18		99	90	
47550002	**Low discrimination**						**47550002**
					GRA		
Fusobacterium mortiferum		73.49	0.46	MLT 11	1		
Prevotella oralis		13.88	0.19	BGL 10	1		
Actinomyces viscosus		9.02	0.18		99		
47550022	**Acceptable identification**						**47550022**
					GRA	CAT	
Prevotella oralis		93.22	0.41	GAL 99	1	1	
Capnocytophaga ochracea		4.21	0.18	LAC 94	1	90	
Actinomyces naeslundii		0.81	0.32		99	25	
47550100	**Very good identification**						**47550100**
					CAT		
Fusobacterium mortiferum		97.15	0.68	MLT 11	5		
Capnocytophaga ochracea		0.92	0.20		90		
Prevotella oralis		0.72	0.18		1		
47550102	**Low discrimination**						**47550102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Prevotella oralis		38.55	0.40	GAL 99	1		
Fusobacterium mortiferum		30.39	0.52	NAG 95	5		
Capnocytophaga ochracea		23.21	0.36		90		
47550103	**Low discrimination**						**47550103**
					GRA		
Actinomyces naeslundii		47.94	0.42	ESL 12	99		
Actinomyces viscosus		26.27	0.14	GAL 96	99		
Prevotella oralis		10.79	0.06		1		
47550120	**Low discrimination**						**47550120**
					CAT		
Prevotella melaninogenica		68.70	0.47	CEL 10	1		
Prevotella oralis		17.14	0.40	GAL 99	1		
Capnocytophaga ochracea		10.32	0.36		90		
47550121	**Low discrimination**						**47550121**
					GRA		
Actinomyces naeslundii		44.19	0.42	ESL 12	99		
Prevotella melaninogenica		39.87	0.12	GAL 96	1		
Prevotella oralis		9.95	0.06		1		
47550122	**Low discrimination**						**47550122**
					CAT		
Prevotella oralis		73.42	0.62	GAL 99	1		
Capnocytophaga ochracea		20.94	0.53		90		
Prevotella melaninogenica		2.95	0.34		1		
47550132	**Low discrimination**						**47550132**
					GRA	CAT	
Capnocytophaga ochracea		66.67	0.36	GAL 99	1	90	
Prevotella oralis		21.25	0.27	MLZ 10	1	1	
Bifidobacterium breve		8.44	0.32		99	1	
47550162	**Low discrimination**						**47550162**
					GRA	CAT	
Capnocytophaga ochracea		67.79	0.36	GAL 99	1	90	
Prevotella oralis		21.61	0.27	SOR 10	1	1	
Bifidobacterium breve		8.58	0.32		99	1	
47550322	**Good identification**						**47550322**
					CAT		
Prevotella oralis		98.12	0.55	GAL 99	1		
Parabacteroides distasonis		0.89	0.33		76		
Capnocytophaga ochracea		0.66	0.18		90		
47550502	**Low discrimination**						**47550502**
					GRA	CAT	
Bifidobacterium adolescentis		54.66	0.37	GAL 99	99	1	
Capnocytophaga ochracea		31.42	0.23	XYL 87	1	90	
Clostridium clostridioforme		7.03	0.19		50	33	
47550512	**Low discrimination**						**47550512**
					GRA	CAT	
Bifidobacterium adolescentis		63.65	0.39	GAL 99	99	1	
Clostridium clostridioforme		30.49	0.30	XYL 87	50	33	
Capnocytophaga ochracea		3.33	0.07		1	90	
47550522	**Low discrimination**						**47550522**
					CAT		
Capnocytophaga ochracea		78.33	0.40	GAL 99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		15.72	0.27		1	
Clostridium clostridioforme		1.95	0.19		33	
47550612	**Good identification**					**47550612**
Clostridium clostridioforme		100.0	0.36	SAL 88		
47550632	**Good identification**					**47550632**
Clostridium clostridioforme		100.0	0.36	SAL 88		
47550712	**Good identification**					**47550712**
Clostridium clostridioforme		100.0	0.45	SAL 88		
47550732	**Good identification**					**47550732**
Clostridium clostridioforme		97.01	0.45	SAL 88		
Parabacteroides distasonis		2.99	0.25	TRE 88		
47551000	**Low discrimination**					**47551000**
					GRA	
Fusobacterium mortiferum		78.17	0.65	MLT 11	1	
Lactobacillus jensenii		5.81	0.36		99	
Prevotella oralis		5.70	0.32		1	
47551001	**Low discrimination**					**47551001**
					GRA	
Actinomyces naeslundii		87.74	0.58	ESL 12	99	
Fusobacterium mortiferum		10.68	0.31		1	
Lactobacillus jensenii		0.79	0.02		99	
47551002	**Low discrimination**					**47551002**
					CAT	
Prevotella oralis		76.16	0.53	LAC 94	1	
Capnocytophaga ochracea		7.26	0.36	NAG 95	90	
Fusobacterium mortiferum		6.11	0.49		5	
47551003	**Low discrimination**					**47551003**
					GRA	
Actinomyces naeslundii		81.92	0.58	ESL 12	99	
Prevotella oralis		13.82	0.19		1	
Actinomyces viscosus		1.84	0.06		99	
47551011	**Good identification**					**47551011**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47551013	**Good identification**					**47551013**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47551020	**Low discrimination**					**47551020**
					CAT	
Prevotella oralis		53.59	0.53	LAC 94	1	
Prevotella melaninogenica		34.00	0.47	BGL 95	1	
Capnocytophaga ochracea		5.11	0.36		90	
47551021	**Low discrimination**					**47551021**
					GRA	
Actinomyces naeslundii		76.93	0.58	ESL 12	99	
Prevotella oralis		12.97	0.19		1	
Prevotella melaninogenica		8.23	0.12		1	
47551022	**Good identification**					**47551022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Prevotella oralis		94.13	0.75	LAC 94	1		
Capnocytophaga ochracea		4.25	0.53		90		
Prevotella melaninogenica		0.60	0.34		1		
47551023	**Low discrimination**						**47551023**
					GRA	CAT	
Prevotella oralis		73.12	0.41	LAC 94	1	1	
Actinomyces naeslundii		22.82	0.58	URE 1	99	25	
Capnocytophaga ochracea		3.30	0.18		1	90	
47551031	**Good identification**						**47551031**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47551032	**Low discrimination**						**47551032**
					GRA	CAT	
Prevotella oralis		62.03	0.41	LAC 94	1	1	
Capnocytophaga ochracea		30.79	0.36	MLZ 1	1	90	
Bifidobacterium breve		3.90	0.32		99	1	
47551033	**Low discrimination**						**47551033**
					GRA	CAT	
Actinomyces naeslundii		52.07	0.36	ESL 12	99	25	
Prevotella oralis		32.03	0.07	MLZ 5	1	1	
Capnocytophaga ochracea		15.90	0.02		1	90	
47551041	**Good identification**						**47551041**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47551043	**Good identification**						**47551043**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47551061	**Good identification**						**47551061**
Actinomyces naeslundii		100.0	0.45	ESL 12			
47551062	**Low discrimination**						**47551062**
					GRA	CAT	
Prevotella oralis		60.80	0.41	LAC 94	1	1	
Capnocytophaga ochracea		30.18	0.36	SOR 1	1	90	
Bifidobacterium breve		3.82	0.32		99	1	
47551063	**Low discrimination**						**47551063**
					GRA	CAT	
Actinomyces naeslundii		78.46	0.45	ESL 12	99	25	
Prevotella oralis		14.39	0.07		1	1	
Capnocytophaga ochracea		7.14	0.02		1	90	
47551100	**Low discrimination**						**47551100**
					CAT		
Fusobacterium mortiferum		35.81	0.71	MLT 11	5		
Capnocytophaga ochracea		22.28	0.54		90		
Prevotella oralis		17.53	0.52		1		
47551101	**Low discrimination**						**47551101**
					GRA	SPO	
Actinomyces naeslundii		70.75	0.66	ESL 12	99	1	
Clostridium chauvoei		10.47	0.29		99	99	
Fusobacterium mortiferum		7.06	0.37		1	1	
47551102	**Low discrimination**						**47551102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Prevotella oralis		58.06	0.74	NAG 95	1	1	
Capnocytophaga ochracea		34.95	0.70		1	90	
Bifidobacterium breve		2.49	0.62		99	1	
47551103	**Low discrimination**						**47551103**
					GRA	CAT	
Actinomyces naeslundii		38.09	0.66	ESL 12	99	25	
Prevotella oralis		35.37	0.40		1	1	
Capnocytophaga ochracea		21.29	0.36		1	90	
47551110	**Low discrimination**						**47551110**
					GRA	CAT	
Capnocytophaga ochracea		49.90	0.38	MLZ 10	1	90	
Bifidobacterium breve		28.47	0.44	NAG 90	99	1	
Fusobacterium mortiferum		7.29	0.37	BGL 90	1	5	
47551111	**Low discrimination**						**47551111**
					GRA	CAT	
Actinomyces naeslundii		82.93	0.44	ESL 12	99	25	
Capnocytophaga ochracea		10.87	0.04	MLZ 5	1	90	
Bifidobacterium breve		6.20	0.10		99	1	
47551112	**Low discrimination**						**47551112**
					GRA	CAT	
Capnocytophaga ochracea		42.18	0.54	MLZ 10	1	90	
Bifidobacterium breve		27.04	0.62	NAG 90	99	1	
Bifidobacterium adolescentis		21.28	0.59		99	1	
47551120	**Low discrimination**						**47551120**
					CAT		
Prevotella melaninogenica		69.82	0.81	CEL 10	1		
Prevotella oralis		17.42	0.74		1		
Capnocytophaga ochracea		10.49	0.70		90		
47551121	**Low discrimination**						**47551121**
					GRA		
Prevotella melaninogenica		58.47	0.47	CEL 10	1		
Actinomyces naeslundii		15.71	0.66	URE 1	99		
Prevotella oralis		14.58	0.40		1		
47551122	**Low discrimination**						**47551122**
					CAT		
Prevotella oralis		73.54	0.96		1		
Capnocytophaga ochracea		20.97	0.87		90		
Prevotella melaninogenica		2.96	0.68		1		
47551123	**Low discrimination**						**47551123**
					GRA	CAT	
Prevotella oralis		70.61	0.62	URE 1	1	1	
Capnocytophaga ochracea		20.14	0.53		1	90	
Actinomyces naeslundii		4.00	0.66		99	25	
47551130	**Low discrimination**						**47551130**
					CAT		
Capnocytophaga ochracea		52.67	0.54	MLZ 10	90		
Prevotella melaninogenica		31.89	0.47	BGL 90	1		
Prevotella oralis		7.95	0.40		1		
47551132	**Low discrimination**						**47551132**
					GRA	CAT	
Capnocytophaga ochracea		67.02	0.70	MLZ 10	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		21.36	0.62		1	1	
Bifidobacterium breve		8.49	0.66		99	1	
47551133	**Low discrimination**						**47551133**
					GRA	CAT	
Capnocytophaga ochracea		63.19	0.36	MLZ 10	1	90	
Prevotella oralis		20.14	0.27	URE 1	1	1	
Bifidobacterium breve		8.00	0.32		99	1	
47551140	**Low discrimination**						**47551140**
					GRA	CAT	
Capnocytophaga ochracea		47.30	0.38	NAG 90	1	90	
Bifidobacterium breve		26.98	0.44	SOR 10	99	1	
Actinomyces naeslundii		8.15	0.50	BGL 90	99	25	
47551141	**Acceptable identification**						**47551141**
					GRA	CAT	
Actinomyces naeslundii		94.21	0.53	ESL 12	99	25	
Capnocytophaga ochracea		3.68	0.04		1	90	
Bifidobacterium breve		2.10	0.10		99	1	
47551142	**Low discrimination**						**47551142**
					GRA	CAT	
Capnocytophaga ochracea		52.92	0.54	NAG 90	1	90	
Bifidobacterium breve		33.92	0.62	SOR 10	99	1	
Prevotella oralis		7.99	0.40		1	1	
47551143	**Low discrimination**						**47551143**
					GRA	CAT	
Actinomyces naeslundii		60.66	0.53	ESL 12	99	25	
Capnocytophaga ochracea		21.35	0.20		1	90	
Bifidobacterium breve		13.68	0.27		99	1	
47551150	**Low discrimination**						**47551150**
					GRA	CAT	
Bifidobacterium breve		81.14	0.44	SAL 99	99	1	
Capnocytophaga ochracea		15.80	0.21	BGL 91	1	90	
Actinomyces naeslundii		1.29	0.28		99	25	
47551152	**Low discrimination**						**47551152**
					GRA	CAT	
Bifidobacterium breve		78.30	0.62	SAL 99	99	1	
Capnocytophaga ochracea		13.57	0.38		1	90	
Bifidobacterium adolescentis		7.62	0.43		99	1	
47551160	**Low discrimination**						**47551160**
					CAT		
Capnocytophaga ochracea		49.36	0.54	SOR 10	90		
Prevotella melaninogenica		29.88	0.47	BGL 90	1		
Prevotella oralis		7.45	0.40		1		
47551161	**Low discrimination**						**47551161**
					GRA	CAT	
Actinomyces naeslundii		58.68	0.53	ESL 12	99	25	
Capnocytophaga ochracea		20.65	0.20		1	90	
Prevotella melaninogenica		12.50	0.12		1	1	
47551162	**Low discrimination**						**47551162**
					GRA	CAT	
Capnocytophaga ochracea		67.67	0.70	SOR 10	1	90	
Prevotella oralis		21.57	0.62		1	1	
Bifidobacterium breve		8.57	0.66		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47551163	**Low discrimination**						**47551163**
					GRA	CAT	
Capnocytophaga ochracea		56.11	0.36	SOR 10	1	90	
Prevotella oralis		17.89	0.27	URE 1	1	1	
Actinomyces naeslundii		17.72	0.53		99	25	
47551170	**Low discrimination**						**47551170**
					GRA	CAT	
Bifidobacterium breve		48.20	0.49	SAL 99	99	1	
Capnocytophaga ochracea		47.53	0.38	BGL 91	1	90	
Prevotella melaninogenica		2.62	0.12		1	1	
47551172	**Low discrimination**						**47551172**
					GRA	CAT	
Bifidobacterium breve		52.41	0.66	SAL 99	99	1	
Capnocytophaga ochracea		46.00	0.54		1	90	
Prevotella oralis		1.33	0.27		1	1	
47551202	**Good identification**						**47551202**
					CAT		
Prevotella oralis		98.92	0.47	LAC 94	1		
Clostridium clostridioforme		0.36	0.13	NAG 95	33		
Capnocytophaga ochracea		0.22	0.02		90		
47551220	**Good identification**						**47551220**
					CAT		
Prevotella oralis		97.86	0.47	LAC 94	1		
Prevotella melaninogenica		1.46	0.12	BGL 95	1		
Capnocytophaga ochracea		0.22	0.02		90		
47551222	**Very good identification**						**47551222**
					CAT		
Prevotella oralis		99.67	0.69	LAC 94	1		
Parabacteroides distasonis		0.14	0.33		76		
Capnocytophaga ochracea		0.11	0.18		90		
47551300	**Low discrimination**						**47551300**
					CAT		
Prevotella oralis		89.69	0.46	NAG 95	1		
Fusobacterium mortiferum		4.32	0.37	BGL 95	5		
Capnocytophaga ochracea		2.69	0.20		90		
47551302	**Very good identification**						**47551302**
					CAT		
Prevotella oralis		97.94	0.68	NAG 95	1		
Capnocytophaga ochracea		1.39	0.36		90		
Parabacteroides distasonis		0.17	0.33		76		
47551320	**Low discrimination**						**47551320**
					CAT		
Prevotella oralis		88.55	0.68	BGL 95	1		
Prevotella melaninogenica		8.37	0.47		1		
Capnocytophaga ochracea		1.26	0.36		90		
47551322	**Very good identification**						**47551322**
					CAT		
Prevotella oralis		98.22	0.89		1		
Parabacteroides distasonis		0.89	0.67		76		
Capnocytophaga ochracea		0.66	0.53		90		
47551323	**Good identification**						**47551323**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		98.12	0.55	URE 1	1		
Parabacteroides distasonis		0.89	0.33		76		
Capnocytophaga ochracea		0.66	0.18		90		
47551332	**Low discrimination**						**47551332**
					CAT		
Prevotella oralis		62.94	0.55	MLZ 1	1		
Parabacteroides distasonis		30.52	0.63		76		
Capnocytophaga ochracea		4.65	0.36		90		
47551362	**Acceptable identification**						**47551362**
					GRA	CAT	
Prevotella oralis		91.31	0.55	SOR 1	1	1	
Capnocytophaga ochracea		6.75	0.36		1	90	
Bifidobacterium breve		0.86	0.32		99	1	
47551402	**Low discrimination**						**47551402**
					GRA	CAT	
Bifidobacterium adolescentis		43.85	0.37	XYL 87	99	1	
Capnocytophaga ochracea		25.21	0.23	LAC 99	1	90	
Prevotella oralis		15.14	0.19		1	1	
47551412	**Low discrimination**						**47551412**
					GRA	CAT	
Bifidobacterium adolescentis		64.99	0.39	XYL 87	99	1	
Clostridium clostridioforme		27.90	0.29	LAC 99	50	33	
Capnocytophaga ochracea		3.40	0.07		1	90	
47551422	**Low discrimination**						**47551422**
					CAT		
Prevotella oralis		54.16	0.41	LAC 94	1		
Capnocytophaga ochracea		42.70	0.40	ARA 1	90		
Clostridium clostridioforme		0.95	0.18		33		
47551500	**Low discrimination**						**47551500**
					GRA	CAT	
Capnocytophaga ochracea		50.43	0.41	NAG 90	1	90	
Bifidobacterium longum subsp. longum		22.50	0.43	BGL 90	99	1	
Bifidobacterium adolescentis		7.98	0.37		99	1	
47551502	**Low discrimination**						**47551502**
					GRA	CAT	
Bifidobacterium adolescentis		59.65	0.72	XYL 87	99	1	
Capnocytophaga ochracea		34.29	0.58		1	90	
Prevotella oralis		3.26	0.40		1	1	
47551503	**Low discrimination**						**47551503**
					GRA	CAT	
Bifidobacterium adolescentis		58.25	0.37	XYL 87	99	1	
Capnocytophaga ochracea		33.48	0.23	URE 1	1	90	
Actinomyces naeslundii		3.43	0.32		99	25	
47551510	**Low discrimination**						**47551510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		70.01	0.47	CEL 1	99	1	
Bifidobacterium adolescentis		17.81	0.39		99	1	
Capnocytophaga ochracea		10.24	0.25		1	90	
47551512	**Good identification**						**47551512**
					GRA	CAT	
Bifidobacterium adolescentis		90.08	0.73	XYL 87	99	1	
Capnocytophaga ochracea		4.71	0.41		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		3.58	0.47		99	1	
47551513	**Acceptable identification**						**47551513**
					GRA	CAT	
Bifidobacterium adolescentis		90.38	0.39	XYL 87	99	1	
Capnocytophaga ochracea		4.72	0.07	URE 1	1	90	
Bifidobacterium longum subsp. longum		3.59	0.13		99	1	
47551520	**Low discrimination**						**47551520**
					GRA	CAT	
Capnocytophaga ochracea		56.47	0.58	BGL 90	1	90	
Prevotella melaninogenica		21.52	0.47		1	1	
Bifidobacterium longum subsp. longum		11.20	0.54		99	1	
47551522	**Low discrimination**						**47551522**
					GRA	CAT	
Capnocytophaga ochracea		79.53	0.74		1	90	
Prevotella oralis		15.96	0.62		1	1	
Bifidobacterium longum subsp. longum		1.75	0.54		99	1	
47551523	**Low discrimination**						**47551523**
					GRA	CAT	
Capnocytophaga ochracea		79.17	0.40	URE 1	1	90	
Prevotella oralis		15.89	0.27		1	1	
Bifidobacterium longum subsp. longum		1.74	0.20		99	1	
47551530	**Low discrimination**						**47551530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.01	0.58	CEL 1	99	1	
Capnocytophaga ochracea		24.36	0.41		1	90	
Prevotella melaninogenica		0.84	0.12		1	1	
47551532	**Low discrimination**						**47551532**
					GRA	CAT	
Capnocytophaga ochracea		66.94	0.58	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		22.60	0.58		99	1	
Parabacteroides distasonis		5.13	0.51		1	76	
47551542	**Low discrimination**						**47551542**
					GRA	CAT	
Bifidobacterium adolescentis		64.00	0.56	XYL 87	99	1	
Capnocytophaga ochracea		33.07	0.41	SOR 11	1	90	
Bifidobacterium breve		1.21	0.27		99	1	
47551552	**Good identification**						**47551552**
					GRA	CAT	
Bifidobacterium adolescentis		93.53	0.57	XYL 87	99	1	
Capnocytophaga ochracea		4.39	0.25	SOR 11	1	90	
Bifidobacterium breve		1.45	0.27		99	1	
47551560	**Acceptable identification**						**47551560**
					GRA	CAT	
Capnocytophaga ochracea		92.36	0.41	SOR 10	1	90	
Prevotella melaninogenica		3.20	0.12	BGL 90	1	1	
Bifidobacterium longum subsp. longum		1.67	0.20		99	1	
47551562	**Very good identification**						**47551562**
					GRA	CAT	
Capnocytophaga ochracea		96.87	0.58	SOR 10	1	90	
Prevotella oralis		1.77	0.27		1	1	
Bifidobacterium breve		0.70	0.32		99	1	
47551572	**Low discrimination**						**47551572**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		88.53	0.41	MLZ 10	1	90	
Bifidobacterium breve		5.77	0.32	SOR 10	99	1	
Bifidobacterium longum subsp. longum		2.72	0.24		99	1	
47551612	**Good identification**						**47551612**
Clostridium clostridioforme		100.0	0.44	SAL 88			
47551632	**Good identification**						**47551632**
Clostridium clostridioforme		95.99	0.44	SAL 88			
Parabacteroides distasonis		3.30	0.25	TRE 88			
Prevotella oralis		0.71	0.01	XYL 91			
47551702	**Low discrimination**						**47551702**
					GRA		
Clostridium clostridioforme		40.17	0.42	SAL 88	50		
Prevotella oralis		32.95	0.33	TRE 88	1		
Bifidobacterium adolescentis		14.21	0.37	XYL 91	99		
47551712	**Low discrimination**						**47551712**
					GRA		
Clostridium clostridioforme		89.63	0.53	SAL 88	50		
Bifidobacterium adolescentis		8.51	0.39	TRE 88	99		
Parabacteroides distasonis		0.92	0.25	XYL 91	1		
47551720	**Acceptable identification**						**47551720**
					CAT		
Bacteroides vulgatus		92.25	0.49	CEL 5	3		
Prevotella oralis		5.14	0.33	XYL 98	1		
Capnocytophaga ochracea		1.27	0.23		90		
47551722	**Low discrimination**						**47551722**
					CAT		
Prevotella oralis		55.95	0.55	ARA 1	1		
Parabacteroides distasonis		30.88	0.64		76		
Capnocytophaga ochracea		6.57	0.40		90		
47551732	**Low discrimination**						**47551732**
Parabacteroides distasonis		47.42	0.59	XYL 90			
Clostridium clostridioforme		46.64	0.53				
Capnocytophaga ochracea		2.08	0.23				
47552001	**Good identification**						**47552001**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47552002	**Low discrimination**						**47552002**
					GRA		
Actinomyces naeslundii		85.67	0.36	ESL 12	99		
Fusobacterium mortiferum		14.33	0.12	GAL 96	1		
47552003	**Good identification**						**47552003**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47552020	**Good identification**						**47552020**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47552021	**Good identification**						**47552021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	ESL 12		
47552022	**Low discrimination**					**47552022**
					GRA	
Actinomyces naeslundii		62.49	0.36	ESL 12	99	
Prevotella oralis		37.51	0.07	GAL 96	1	
47552023	**Good identification**					**47552023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47552101	**Good identification**					**47552101**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47552102	**Low discrimination**					**47552102**
Actinomyces naeslundii		36.72	0.44	ESL 12		
Bifidobacterium breve		27.11	0.27	GAL 96		
Bifidobacterium adolescentis		14.29	0.22			
47552103	**Good identification**					**47552103**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47552120	**Low discrimination**					**47552120**
					GRA	
Actinomyces naeslundii		46.03	0.44	ESL 12	99	
Prevotella melaninogenica		32.09	0.12	GAL 96	1	
Prevotella oralis		8.01	0.06		1	
47552121	**Good identification**					**47552121**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47552123	**Good identification**					**47552123**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47552502	**Good identification**					**47552502**
Bifidobacterium adolescentis		96.15	0.36	GAL 99		
Actinomyces israelii		3.85	0.18	XYL 87		
47552512	**Good identification**					**47552512**
Bifidobacterium adolescentis		100.0	0.37	GAL 99		
47553000	**Low discrimination**					**47553000**
					GRA	
Actinomyces naeslundii		72.25	0.60	ESL 12	99	
Actinomyces israelii		18.00	0.41		99	
Fusobacterium mortiferum		6.80	0.31		1	
47553001	**Very good identification**					**47553001**
Actinomyces naeslundii		100.0	0.63	ESL 12		
47553002	**Low discrimination**					**47553002**
					GRA	
Actinomyces naeslundii		59.71	0.60	ESL 12	99	
Actinomyces israelii		22.31	0.44		99	
Prevotella oralis		7.78	0.19		1	
47553003	**Very good identification**					**47553003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.63	ESL 12		
47553010	**Low discrimination**					**47553010**
Actinomyces naeslundii		53.43	0.38	ESL 12		
Actinomyces israelii		37.78	0.26	MLZ 5		
Bifidobacterium breve		8.79	0.10			
47553011	**Good identification**					**47553011**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47553013	**Good identification**					**47553013**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47553020	**Low discrimination**					**47553020**
					GRA	
Actinomyces naeslundii		78.20	0.60	ESL 12	99	
Prevotella oralis		10.19	0.19		1	
Prevotella melaninogenica		6.47	0.12		1	
47553021	**Very good identification**					**47553021**
Actinomyces naeslundii		100.0	0.63	ESL 12		
47553022	**Low discrimination**					**47553022**
					GRA	
Prevotella oralis		64.24	0.41	MAN 1	1	
Actinomyces naeslundii		25.94	0.60	LAC 94	99	
Bifidobacterium breve		4.04	0.32		99	
47553023	**Very good identification**					**47553023**
					GRA	
Actinomyces naeslundii		98.36	0.63	ESL 12	99	
Prevotella oralis		1.64	0.07		1	
47553030	**Low discrimination**					**47553030**
Actinomyces naeslundii		77.37	0.38	ESL 12		
Bifidobacterium breve		22.63	0.15	MLZ 5		
47553031	**Good identification**					**47553031**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47553033	**Good identification**					**47553033**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47553040	**Low discrimination**					**47553040**
Actinomyces naeslundii		62.58	0.47	ESL 12		
Actinomyces israelii		34.35	0.34			
Bifidobacterium breve		3.07	0.10			
47553041	**Good identification**					**47553041**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47553042	**Low discrimination**					**47553042**
Actinomyces naeslundii		43.11	0.47	ESL 12		
Actinomyces israelii		35.50	0.37			
Bifidobacterium breve		21.39	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47553043	**Good identification**					**47553043**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47553060	**Acceptable identification**					**47553060**
Actinomyces naeslundii		91.98	0.47	ESL 12		
Bifidobacterium breve		8.02	0.15			
47553061	**Good identification**					**47553061**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47553062	**Low discrimination**					**47553062**
					GRA	
Actinomyces naeslundii		46.59	0.47	ESL 12	99	
Bifidobacterium breve		41.09	0.32		99	
Prevotella oralis		6.60	0.07		1	
47553063	**Good identification**					**47553063**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47553100	**Low discrimination**					**47553100**
Actinomyces naeslundii		53.33	0.68	ESL 12		
Actinomyces israelii		18.67	0.51			
Bifidobacterium breve		16.06	0.44			
47553101	**Very good identification**					**47553101**
Actinomyces naeslundii		99.56	0.71	ESL 12		
Actinomyces israelii		0.24	0.17			
Bifidobacterium breve		0.20	0.10			
47553102	**Low discrimination**					**47553102**
Bifidobacterium breve		41.18	0.62	SAL 99		
Bifidobacterium adolescentis		21.70	0.56			
Actinomyces naeslundii		13.52	0.68			
47553103	**Very good identification**					**47553103**
Actinomyces naeslundii		95.92	0.71	ESL 12		
Bifidobacterium breve		1.97	0.27			
Bifidobacterium adolescentis		1.04	0.22			
47553110	**Low discrimination**					**47553110**
Bifidobacterium breve		69.84	0.44	SAL 99		
Actinomyces naeslundii		12.20	0.46	BGL 91		
Actinomyces israelii		12.13	0.37			
47553111	**Good identification**					**47553111**
Actinomyces naeslundii		96.29	0.49	ESL 12		
Bifidobacterium breve		3.71	0.10	MLZ 5		
47553112	**Low discrimination**					**47553112**
Bifidobacterium breve		58.56	0.62	SAL 99		
Bifidobacterium adolescentis		37.72	0.57			
Actinomyces israelii		1.51	0.40			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47553113	**Low discrimination**						**47553113**
Actinomyces naeslundii		60.96	0.49	ESL 12			
Bifidobacterium breve		23.75	0.27	MLZ 5			
Bifidobacterium adolescentis		15.30	0.23				
47553120	**Low discrimination**						**47553120**
					GRA		
Prevotella melaninogenica		48.35	0.47	CEL 10	1		
Actinomyces naeslundii		16.81	0.68	MAN 1	99		
Prevotella oralis		12.06	0.40		1		
47553121	**Very good identification**						**47553121**
					GRA		
Actinomyces naeslundii		96.85	0.71	ESL 12	99		
Prevotella melaninogenica		1.88	0.12		1		
Prevotella oralis		0.47	0.06		1		
47553122	**Low discrimination**						**47553122**
					GRA	CAT	
Prevotella oralis		52.24	0.62	MAN 1	1	1	
Bifidobacterium breve		20.75	0.66		99	1	
Capnocytophaga ochracea		14.90	0.53		1	90	
47553123	**Low discrimination**						**47553123**
					GRA		
Actinomyces naeslundii		85.62	0.71	ESL 12	99		
Prevotella oralis		7.86	0.27		1		
Bifidobacterium breve		3.12	0.32		99		
47553130	**Low discrimination**						**47553130**
					GRA	CAT	
Bifidobacterium breve		78.35	0.49	SAL 99	99	1	
Actinomyces naeslundii		7.70	0.46	BGL 91	99	25	
Capnocytophaga ochracea		7.02	0.20		1	90	
47553131	**Acceptable identification**						**47553131**
Actinomyces naeslundii		93.59	0.49	ESL 12			
Bifidobacterium breve		6.41	0.15	MLZ 5			
47553132	**Low discrimination**						**47553132**
					GRA	CAT	
Bifidobacterium breve		88.67	0.66	SAL 99	99	1	
Capnocytophaga ochracea		7.07	0.36		1	90	
Prevotella oralis		2.26	0.27		1	1	
47553133	**Low discrimination**						**47553133**
					GRA	CAT	
Actinomyces naeslundii		57.21	0.49	ESL 12	99	25	
Bifidobacterium breve		39.63	0.32	MLZ 5	99	1	
Capnocytophaga ochracea		3.16	0.02		1	90	
47553140	**Low discrimination**						**47553140**
Bifidobacterium breve		48.65	0.44	SAL 99			
Actinomyces naeslundii		28.50	0.55	BGL 91			
Actinomyces israelii		21.99	0.44				
47553141	**Very good identification**						**47553141**
Actinomyces naeslundii		98.86	0.58	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		1.14	0.10				
47553142	**Low discrimination**						**47553142**
Bifidobacterium breve		82.64	0.62	SAL 99			
Actinomyces israelii		5.54	0.47				
Bifidobacterium adolescentis		5.38	0.40				
47553143	**Low discrimination**						**47553143**
Actinomyces naeslundii		89.59	0.58	ESL 12			
Bifidobacterium breve		10.41	0.27				
47553150	**Acceptable identification**						**47553150**
Bifidobacterium breve		90.37	0.44	SAL 99			
Actinomyces israelii		6.10	0.30	BGL 91			
Actinomyces naeslundii		2.79	0.33				
47553151	**Low discrimination**						**47553151**
Actinomyces naeslundii		82.08	0.36	ESL 12			
Bifidobacterium breve		17.93	0.10	MLZ 5			
47553152	**Good identification**						**47553152**
Bifidobacterium breve		91.37	0.62	SAL 99			
Bifidobacterium adolescentis		7.27	0.42				
Actinomyces israelii		0.92	0.33				
47553160	**Low discrimination**						**47553160**
					GRA	CAT	
Bifidobacterium breve		65.78	0.49	SAL 99	99	1	
Actinomyces naeslundii		21.68	0.55	BGL 91	99	25	
Capnocytophaga ochracea		5.90	0.20		1	90	
47553161	**Very good identification**						**47553161**
Actinomyces naeslundii		98.00	0.58	ESL 12			
Bifidobacterium breve		2.00	0.15				
47553162	**Low discrimination**						**47553162**
					GRA	CAT	
Bifidobacterium breve		87.37	0.66	SAL 99	99	1	
Capnocytophaga ochracea		6.97	0.36		1	90	
Actinomyces naeslundii		2.85	0.55		99	25	
47553163	**Low discrimination**						**47553163**
					GRA	CAT	
Actinomyces naeslundii		81.76	0.58	ESL 12	99	25	
Bifidobacterium breve		16.89	0.32		99	1	
Capnocytophaga ochracea		1.35	0.02		1	90	
47553170	**Good identification**						**47553170**
					GRA	CAT	
Bifidobacterium breve		97.34	0.49	SAL 99	99	1	
Actinomyces naeslundii		1.69	0.33	BGL 91	99	25	
Capnocytophaga ochracea		0.97	0.04		1	90	
47553171	**Low discrimination**						**47553171**
Actinomyces naeslundii		72.04	0.36	ESL 12			
Bifidobacterium breve		27.96	0.15	MLZ 5			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47553172	**Very good identification**						**47553172**
					GRA	CAT	
Bifidobacterium breve		98.91	0.66	SAL 99	99	1	
Capnocytophaga ochracea		0.88	0.20		1	90	
Actinomyces naeslundii		0.17	0.33		99	25	
47553301	**Good identification**						**47553301**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47553303	**Low discrimination**						**47553303**
					GRA		
Actinomyces naeslundii		82.99	0.37	ESL 12	99		
Prevotella oralis		17.01	-.01	RHA 1	1		
47553320	**Low discrimination**						**47553320**
					SPO		
Clostridium ramosum		50.38	0.35	SAL 99	92		
Prevotella oralis		42.31	0.33	TRE 94	1		
Prevotella melaninogenica		4.00	0.12		1		
47553321	**Low discrimination**						**47553321**
					SPO	GRA	
Actinomyces naeslundii		69.01	0.37	ESL 12	1	99	
Clostridium ramosum		16.84	0.01	RHA 1	92	50	
Prevotella oralis		14.14	-.01		1	1	
47553322	**Low discrimination**						**47553322**
					SPO	GRA	
Prevotella oralis		78.15	0.55	MAN 1	1	1	
Clostridium ramosum		19.59	0.45		92	50	
Bifidobacterium breve		0.73	0.32		1	99	
47553400	**Good identification**						**47553400**
Actinomyces israelii		95.28	0.39	SAL 92			
Actinomyces naeslundii		4.72	0.26	XYL 92			
47553402	**Low discrimination**						**47553402**
Bifidobacterium adolescentis		49.44	0.36	XYL 87			
Actinomyces israelii		48.95	0.42	LAC 99			
Actinomyces naeslundii		1.62	0.26				
47553412	**Low discrimination**						**47553412**
Bifidobacterium adolescentis		89.20	0.37	XYL 87			
Actinomyces israelii		10.80	0.28	LAC 99			
47553442	**Low discrimination**						**47553442**
Actinomyces israelii		75.70	0.35	SAL 92			
Bifidobacterium adolescentis		24.30	0.20	XYL 92			
47553500	**Low discrimination**						**47553500**
Actinomyces israelii		66.59	0.49	SAL 92			
Bifidobacterium adolescentis		25.22	0.36	XYL 92			
Actinomyces naeslundii		2.35	0.34				
47553501	**Low discrimination**						**47553501**
Actinomyces naeslundii		83.97	0.37	ESL 12			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		16.03	0.15	ARA 1			
47553502	**Very good identification**						**47553502**
					GRA	CAT	
Bifidobacterium adolescentis		95.03	0.70	XYL 87	99	1	
Actinomyces israelii		3.80	0.52		99	1	
Capnocytophaga ochracea		0.67	0.23		1	90	
47553503	**Low discrimination**						**47553503**
Bifidobacterium adolescentis		84.77	0.36	XYL 87			
Actinomyces naeslundii		11.84	0.37	URE 1			
Actinomyces israelii		3.39	0.18				
47553510	**Low discrimination**						**47553510**
Bifidobacterium adolescentis		71.72	0.37	XYL 87			
Actinomyces israelii		23.15	0.35	BGL 99			
Bifidobacterium longum subsp. longum		3.48	0.13				
47553512	**Very good identification**						**47553512**
Bifidobacterium adolescentis		99.12	0.72	XYL 87			
Actinomyces israelii		0.49	0.38				
Bifidobacterium breve		0.23	0.27				
47553513	**Good identification**						**47553513**
Bifidobacterium adolescentis		100.0	0.37	XYL 87			
47553521	**Good identification**						**47553521**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47553522	**Low discrimination**						**47553522**
					GRA	CAT	
Capnocytophaga ochracea		64.08	0.40	MAN 1	1	90	
Prevotella oralis		12.86	0.27		1	1	
Bifidobacterium adolescentis		10.13	0.36		99	1	
47553523	**Low discrimination**						**47553523**
					GRA		
Actinomyces naeslundii		68.62	0.37	ESL 12	99		
Capnocytophaga ochracea		31.38	0.06	ARA 1	1		
47553532	**Low discrimination**						**47553532**
					GRA	CAT	
Bifidobacterium adolescentis		42.97	0.37	XYL 87	99	1	
Capnocytophaga ochracea		24.70	0.23	NAG 1	1	90	
Bifidobacterium breve		17.72	0.32		99	1	
47553540	**Low discrimination**						**47553540**
Actinomyces israelii		87.13	0.42	SAL 92			
Bifidobacterium adolescentis		10.49	0.20	XYL 92			
Bifidobacterium breve		2.38	0.10				
47553542	**Low discrimination**						**47553542**
Bifidobacterium adolescentis		86.55	0.54	XYL 87			
Actinomyces israelii		10.89	0.45	SOR 11			
Bifidobacterium breve		2.01	0.27				
47553552	**Good identification**						**47553552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		96.68	0.56	XYL 87			
Bifidobacterium breve		1.83	0.27	SOR 11			
Actinomyces israelii		1.49	0.31				
47553702	**Low discrimination**						**47553702**
Bifidobacterium adolescentis		73.44	0.36	XYL 87			
Actinomyces israelii		21.89	0.33	RHA 1			
Clostridium clostridioforme		2.56	0.08				
47553712	**Low discrimination**						**47553712**
Bifidobacterium adolescentis		85.72	0.37	XYL 87			
Clostridium clostridioforme		11.15	0.19	RHA 1			
Actinomyces israelii		3.12	0.19				
47554000	**Good identification**						**47554000**
Fusobacterium mortiferum		100.0	0.43	MLT 11			
47554100	**Low discrimination**						**47554100**
					CAT		
Fusobacterium mortiferum		89.39	0.49	MLT 11	5		
Bacteroides fragilis		4.30	0.04	XYL 7	99		
Capnocytophaga ochracea		2.80	0.10		90		
47554112	**Low discrimination**						**47554112**
					GRA	CAT	
Bifidobacterium adolescentis		71.17	0.39	GAL 99	99	1	
Clostridium clostridioforme		22.00	0.27	ARA 87	50	33	
Capnocytophaga ochracea		5.27	0.10		1	90	
47554120	**Low discrimination**						**47554120**
					CAT		
Bacteroides fragilis		87.84	0.38	GAL 99	99		
Capnocytophaga ochracea		5.20	0.26	BGL 95	90		
Prevotella oralis		4.27	0.24		1		
47554122	**Good identification**						**47554122**
					CAT		
Bacteroides fragilis		92.42	0.60	GAL 99	99		
Prevotella oralis		4.49	0.46		1		
Capnocytophaga ochracea		2.59	0.42		90		
47554312	**Good identification**						**47554312**
Clostridium clostridioforme		99.30	0.42	SAL 88			
Parabacteroides distasonis		0.70	0.11	TRE 88			
47554322	**Low discrimination**						**47554322**
					CAT		
Prevotella oralis		43.78	0.40	GAL 99	1		
Parabacteroides distasonis		29.00	0.49	XYL 11	76		
Bacteroides fragilis		21.22	0.25		99		
47554332	**Low discrimination**						**47554332**
Clostridium clostridioforme		57.79	0.42	SAL 88			
Parabacteroides distasonis		40.51	0.45	TRE 88			
Prevotella oralis		1.15	0.06	ARA 94			
47554412	**Good identification**						**47554412**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		95.54	0.38	SAL 88			
Bifidobacterium adolescentis		4.46	0.19	TRE 88			
47554432	**Good identification**						**47554432**
Clostridium clostridioforme		100.0	0.38	SAL 88			
47554502	**Low discrimination**						**47554502**
					GRA	CAT	
Bifidobacterium adolescentis		79.68	0.52	GAL 99	99	1	
Clostridium clostridioforme		15.47	0.36		50	33	
Capnocytophaga ochracea		1.71	0.13		1	90	
47554512	**Low discrimination**						**47554512**
Bifidobacterium adolescentis		57.42	0.53	GAL 99			
Clostridium clostridioforme		41.56	0.47				
Bifidobacterium longum subsp. longum		1.02	0.21				
47554532	**Low discrimination**						**47554532**
					GRA		
Clostridium clostridioforme		82.77	0.47	SAL 88	50		
Bifidobacterium longum subsp. longum		8.14	0.32	TRE 88	99		
Parabacteroides distasonis		5.55	0.33	RHA 88	1		
47554542	**Good identification**						**47554542**
Bifidobacterium adolescentis		96.97	0.36	GAL 99			
Actinomyces israelii		3.03	0.16	SOR 11			
47554552	**Acceptable identification**						**47554552**
Bifidobacterium adolescentis		94.42	0.37	GAL 99			
Clostridium clostridioforme		5.59	0.13	SOR 11			
47554602	**Good identification**						**47554602**
Clostridium clostridioforme		100.0	0.42	SAL 88			
47554612	**Good identification**						**47554612**
Clostridium clostridioforme		100.0	0.53	SAL 88			
47554622	**Good identification**						**47554622**
Clostridium clostridioforme		99.22	0.42	SAL 88			
Parabacteroides distasonis		0.78	0.12	TRE 88			
47554632	**Good identification**						**47554632**
Clostridium clostridioforme		100.0	0.53	SAL 88			
47554702	**Good identification**						**47554702**
					GRA		
Clostridium clostridioforme		99.07	0.51	SAL 88	50		
Bifidobacterium adolescentis		0.70	0.17	TRE 88	99		
Parabacteroides distasonis		0.23	0.12		1		
47554712	**Very good identification**						**47554712**
Clostridium clostridioforme		99.81	0.62	SAL 88			
Bifidobacterium adolescentis		0.19	0.19	TRE 88			
47554720	**Good identification**						**47554720**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		99.43	0.44	CEL 5			
Clostridium clostridioforme		0.46	0.17	GAL 99			
Parabacteroides distasonis		0.11	0.12				
47554722	**Low discrimination**						**47554722**
Clostridium clostridioforme		72.75	0.51	SAL 88			
Parabacteroides distasonis		16.88	0.46	TRE 88			
Bacteroides uniformis		5.61	0.19				
47554732	**Very good identification**						**47554732**
Clostridium clostridioforme		96.98	0.62	SAL 88			
Parabacteroides distasonis		2.66	0.41	TRE 88			
Bacteroides thetaiotaomicron		0.19	0.05				
47555000	**Low discrimination**						**47555000**
					GRA		
Fusobacterium mortiferum		48.96	0.46	MLT 11	1		
Actinomyces israelii		35.30	0.46	XYL 7	99		
Prevotella oralis		5.86	0.16		1		
47555002	**Low discrimination**						**47555002**
					CAT	GRA	
Prevotella oralis		38.29	0.38	XYL 11	1	1	
Bacteroides fragilis		23.92	0.25	LAC 94	99	1	
Actinomyces israelii		18.21	0.49	NAG 95	1	99	
47555012	**Low discrimination**						**47555012**
Bifidobacterium adolescentis		60.48	0.39	LAC 99			
Clostridium clostridioforme		16.76	0.26	ARA 87			
Actinomyces israelii		14.86	0.35				
47555020	**Low discrimination**						**47555020**
					CAT		
Bacteroides fragilis		71.24	0.38	LAC 99	99		
Prevotella oralis		21.89	0.38	BGL 95	1		
Capnocytophaga ochracea		4.22	0.26		90		
47555022	**Low discrimination**						**47555022**
					CAT		
Bacteroides fragilis		74.55	0.60	LAC 99	99		
Prevotella oralis		22.91	0.60		1		
Capnocytophaga ochracea		2.09	0.42		90		
47555040	**Good identification**						**47555040**
					GRA		
Actinomyces israelii		96.52	0.39	SAL 92	99		
Fusobacterium mortiferum		3.48	0.12		1		
47555042	**Low discrimination**						**47555042**
					GRA	CAT	
Actinomyces israelii		73.38	0.42	SAL 92	99	1	
Bifidobacterium adolescentis		11.61	0.22		99	1	
Capnocytophaga ochracea		8.50	0.10		1	90	
47555100	**Low discrimination**						**47555100**
					GRA	CAT	
Bacteroides fragilis		33.48	0.38	NAG 99	1	99	
Capnocytophaga ochracea		21.81	0.44	BGL 95	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		13.05	0.56		99	1	
47555102	**Low discrimination**						**47555102**
					GRA	CAT	
Bacteroides fragilis		50.24	0.60	NAG 99	1	99	
Bifidobacterium adolescentis		19.04	0.72		99	1	
Capnocytophaga ochracea		15.50	0.60		1	90	
47555110	**Low discrimination**						**47555110**
					GRA	CAT	
Bifidobacterium adolescentis		34.42	0.39	ARA 87	99	1	
Capnocytophaga ochracea		28.03	0.27	BGL 99	1	90	
Actinomyces israelii		22.55	0.42		99	1	
47555112	**Low discrimination**						**47555112**
					GRA	CAT	
Bifidobacterium adolescentis		88.52	0.73	ARA 87	99	1	
Capnocytophaga ochracea		6.55	0.44		1	90	
Bacteroides fragilis		1.93	0.25		1	99	
47555113	**Acceptable identification**						**47555113**
					GRA	CAT	
Bifidobacterium adolescentis		91.25	0.39	ARA 87	99	1	
Capnocytophaga ochracea		6.76	0.10	URE 1	1	90	
Bacteroides fragilis		1.99	-.09		1	99	
47555120	**Low discrimination**						**47555120**
					CAT		
Bacteroides fragilis		87.83	0.72	BGL 95	99		
Capnocytophaga ochracea		5.20	0.60		90		
Prevotella oralis		4.27	0.58		1		
47555121	**Low discrimination**						**47555121**
					CAT		
Bacteroides fragilis		87.67	0.38	URE 1	99		
Capnocytophaga ochracea		5.19	0.26	BGL 95	90		
Prevotella oralis		4.26	0.24		1		
47555122	**Good identification**						**47555122**
					CAT		
Bacteroides fragilis		92.44	0.94		99		
Prevotella oralis		4.50	0.80		1		
Capnocytophaga ochracea		2.59	0.76		90		
47555123	**Good identification**						**47555123**
					CAT		
Bacteroides fragilis		92.43	0.60	URE 1	99		
Prevotella oralis		4.50	0.46		1		
Capnocytophaga ochracea		2.59	0.42		90		
47555130	**Low discrimination**						**47555130**
					CAT		
Bacteroides fragilis		54.81	0.38	MLZ 1	99		
Capnocytophaga ochracea		35.70	0.44	BGL 95	90		
Prevotella oralis		2.67	0.24		1		
47555132	**Low discrimination**						**47555132**
Bacteroides fragilis		61.88	0.60	MLZ 1			
Capnocytophaga ochracea		19.09	0.60				
Parabacteroides distasonis		15.18	0.71				
47555140	**Low discrimination**						**47555140**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO	CAT	
Actinomyces israelii		57.06	0.49	SAL 92	99	1	1	
Capnocytophaga ochracea		27.26	0.27		1	1	90	
Clostridium butyricum		5.70	0.11		99	99	1	
47555142	**Low discrimination**							**47555142**
					GRA	CAT		
Bifidobacterium adolescentis		43.79	0.56	ARA 87	99	1		
Capnocytophaga ochracea		32.06	0.44	SOR 11	1	90		
Actinomyces israelii		11.19	0.52		99	1		
47555152	**Low discrimination**							**47555152**
					GRA	CAT		
Bifidobacterium adolescentis		89.68	0.57	ARA 87	99	1		
Capnocytophaga ochracea		5.97	0.27	SOR 11	1	90		
Actinomyces israelii		2.80	0.38		99	1		
47555160	**Low discrimination**							**47555160**
					CAT			
Bacteroides fragilis		57.28	0.38	SOR 1	99			
Capnocytophaga ochracea		37.31	0.44	BGL 95	90			
Prevotella oralis		2.79	0.24		1			
47555162	**Low discrimination**							**47555162**
					CAT			
Bacteroides fragilis		73.26	0.60	SOR 1	99			
Capnocytophaga ochracea		22.60	0.60		90			
Prevotella oralis		3.56	0.46		1			
47555172	**Low discrimination**							**47555172**
Capnocytophaga ochracea		69.09	0.44	MLZ 10				
Bacteroides fragilis		20.36	0.25	SOR 10				
Parabacteroides distasonis		4.99	0.37					
47555212	**Good identification**							**47555212**
					GRA			
Clostridium clostridioforme		97.62	0.41	SAL 88	50			
Actinomyces israelii		0.89	0.15	TRE 88	99			
Parabacteroides distasonis		0.77	0.11	ARA 94	1			
47555222	**Low discrimination**							**47555222**
					CAT			
Prevotella oralis		83.11	0.53	XYL 11	1			
Parabacteroides distasonis		8.71	0.49	LAC 94	76			
Bacteroides fragilis		6.37	0.25		99			
47555232	**Low discrimination**							**47555232**
Clostridium clostridioforme		51.74	0.41	SAL 88				
Parabacteroides distasonis		40.47	0.45	TRE 88				
Prevotella oralis		7.24	0.19	ARA 94				
47555302	**Low discrimination**							**47555302**
					CAT			
Prevotella oralis		72.66	0.52	XYL 11	1			
Parabacteroides distasonis		9.24	0.49	NAG 95	76			
Bacteroides fragilis		6.76	0.25		99			
47555312	**Low discrimination**							**47555312**
					GRA			
Clostridium clostridioforme		69.44	0.50	SAL 88	50			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		16.22	0.45	TRE 88	1	
Bifidobacterium adolescentis		10.21	0.39	ARA 94	99	
47555320	**Low discrimination**					**47555320**
					CAT	
Bacteroides vulgatus		76.58	0.59	CEL 5	3	
Prevotella oralis		14.15	0.52	ARA 93	1	
Bacteroides fragilis		6.86	0.38		99	
47555322	**Low discrimination**					**47555322**
					CAT	
Prevotella oralis		45.87	0.74	XYL 11	1	
Parabacteroides distasonis		30.38	0.84		76	
Bacteroides fragilis		22.23	0.60		99	
47555323	**Low discrimination**					**47555323**
					CAT	
Prevotella oralis		45.96	0.40	XYL 11	1	
Parabacteroides distasonis		30.44	0.49	URE 1	76	
Bacteroides fragilis		22.28	0.25		99	
47555330	**Low discrimination**					**47555330**
Parabacteroides distasonis		47.44	0.45	BGL 99		
Bacteroides vulgatus		37.90	0.25			
Prevotella oralis		7.00	0.18			
47555332	**Good identification**					**47555332**
Parabacteroides distasonis		91.61	0.79			
Clostridium clostridioforme		3.96	0.50			
Prevotella oralis		2.60	0.40			
47555333	**Acceptable identification**					**47555333**
Parabacteroides distasonis		92.14	0.45	URE 1		
Clostridium clostridioforme		3.98	0.16			
Prevotella oralis		2.61	0.06			
47555362	**Low discrimination**					**47555362**
					CAT	
Prevotella oralis		43.26	0.40	XYL 11	1	
Parabacteroides distasonis		28.65	0.49	SOR 1	76	
Bacteroides fragilis		20.97	0.25		99	
47555372	**Low discrimination**					**47555372**
Parabacteroides distasonis		88.35	0.45	SOR 1		
Capnocytophaga ochracea		4.12	0.10			
Clostridium clostridioforme		3.82	0.16			
47555400	**Low discrimination**					**47555400**
					SPO	
Actinomyces israelii		49.47	0.44	SAL 92	1	
Clostridium butyricum		40.51	0.22		99	
Bifidobacterium longum subsp. longum		3.86	0.17		1	
47555402	**Low discrimination**					**47555402**
Bifidobacterium adolescentis		65.77	0.52	LAC 99		
Actinomyces israelii		16.16	0.47			
Clostridium clostridioforme		11.44	0.35			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47555412	**Low discrimination**						**47555412**
Bifidobacterium adolescentis		58.87	0.53	LAC 99			
Clostridium clostridioforme		38.18	0.47				
Actinomyces israelii		1.77	0.33				
47555432	**Low discrimination**						**47555432**
					GRA		
Clostridium clostridioforme		81.15	0.47	SAL 88	50		
Bifidobacterium longum subsp. longum		8.91	0.32	TRE 88	99		
Parabacteroides distasonis		6.07	0.33	RHA 88	1		
47555440	**Low discrimination**						**47555440**
					SPO		
Actinomyces israelii		84.52	0.37	SAL 92	1		
Clostridium butyricum		15.48	0.04		99		
47555442	**Low discrimination**						**47555442**
					SPO		
Bifidobacterium adolescentis		55.26	0.36	LAC 99	1		
Actinomyces israelii		42.71	0.40	SOR 11	1		
Clostridium butyricum		2.03	-.03		99		
47555452	**Low discrimination**						**47555452**
Bifidobacterium adolescentis		87.15	0.37	LAC 99			
Actinomyces israelii		8.24	0.26	SOR 11			
Clostridium clostridioforme		4.62	0.12				
47555500	**Low discrimination**						**47555500**
					SPO		
Clostridium butyricum		47.60	0.45	SAL 99	99		
Bifidobacterium longum subsp. longum		18.70	0.52	TRE 94	1		
Bifidobacterium adolescentis		14.79	0.52		1		
47555502	**Good identification**						**47555502**
					GRA	CAT	
Bifidobacterium adolescentis		93.48	0.86		99	1	
Capnocytophaga ochracea		2.01	0.47		1	90	
Bifidobacterium longum subsp. longum		1.19	0.52		99	1	
47555503	**Acceptable identification**						**47555503**
					GRA	CAT	
Bifidobacterium adolescentis		93.75	0.52	URE 1	99	1	
Capnocytophaga ochracea		2.01	0.13		1	90	
Bifidobacterium longum subsp. longum		1.20	0.17		99	1	
47555510	**Low discrimination**						**47555510**
					SPO		
Bifidobacterium longum subsp. longum		58.21	0.56	CEL 1	1		
Bifidobacterium adolescentis		33.04	0.53		1		
Clostridium butyricum		4.58	0.24		99		
47555512	**Very good identification**						**47555512**
Bifidobacterium adolescentis		95.79	0.87				
Clostridium clostridioforme		2.10	0.56				
Bifidobacterium longum subsp. longum		1.71	0.56				
47555513	**Good identification**						**47555513**
Bifidobacterium adolescentis		96.18	0.53	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		2.11	0.21				
Bifidobacterium longum subsp. longum		1.71	0.21				
47555520	**Low discrimination**						**47555520**
					GRA	CAT	
Bacteroides vulgatus		77.39	0.62	CEL 5	1	3	
Bifidobacterium longum subsp. longum		11.56	0.62	RHA 90	99	1	
Capnocytophaga ochracea		4.86	0.47		1	90	
47555522	**Low discrimination**						**47555522**
Bacteroides fragilis		45.22	0.60	ARA 1			
Capnocytophaga ochracea		22.16	0.64				
Parabacteroides distasonis		12.63	0.72				
47555530	**Good identification**						**47555530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.30	0.66	CEL 1	99	1	
Bacteroides vulgatus		3.67	0.28		1	3	
Capnocytophaga ochracea		2.53	0.31		1	90	
47555532	**Low discrimination**						**47555532**
					GRA		
Bifidobacterium longum subsp. longum		40.71	0.66	CEL 1	99		
Parabacteroides distasonis		27.74	0.67		1		
Clostridium clostridioforme		12.54	0.56		50		
47555542	**Good identification**						**47555542**
					GRA	CAT	
Bifidobacterium adolescentis		94.23	0.70	SOR 11	99	1	
Actinomyces israelii		2.94	0.51		99	1	
Capnocytophaga ochracea		1.82	0.31		1	90	
47555543	**Good identification**						**47555543**
Bifidobacterium adolescentis		96.97	0.36	SOR 11			
Actinomyces israelii		3.03	0.16	URE 1			
47555550	**Low discrimination**						**47555550**
Bifidobacterium adolescentis		64.85	0.37	SOR 11			
Actinomyces israelii		16.34	0.33	BGL 99			
Bifidobacterium longum subsp. longum		9.34	0.21				
47555552	**Very good identification**						**47555552**
Bifidobacterium adolescentis		99.09	0.72	SOR 11			
Actinomyces israelii		0.38	0.37				
Clostridium clostridioforme		0.18	0.21				
47555553	**Good identification**						**47555553**
Bifidobacterium adolescentis		100.0	0.37	SOR 11			
47555562	**Low discrimination**						**47555562**
					GRA	CAT	
Capnocytophaga ochracea		72.05	0.47	SOR 10	1	90	
Bacteroides fragilis		13.37	0.25		1	99	
Bifidobacterium adolescentis		4.19	0.36		99	1	
47555602	**Good identification**						**47555602**
Clostridium clostridioforme		97.36	0.50	SAL 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		1.41	0.28	TRE 88		
Bifidobacterium adolescentis		0.77	0.17			
47555612	**Very good identification**					**47555612**
Clostridium clostridioforme		99.74	0.61	SAL 88		
Bifidobacterium adolescentis		0.21	0.19	TRE 88		
Actinomyces israelii		0.05	0.14			
47555620	**Good identification**					**47555620**
Bacteroides vulgatus		99.39	0.44	CEL 5		
Clostridium clostridioforme		0.42	0.16	LAC 99		
Parabacteroides distasonis		0.11	0.12			
47555622	**Low discrimination**					**47555622**
Clostridium clostridioforme		63.86	0.50	SAL 88		
Parabacteroides distasonis		16.53	0.46	TRE 88		
Bacteroides uniformis		11.10	0.24			
47555632	**Very good identification**					**47555632**
Clostridium clostridioforme		96.43	0.61	SAL 88		
Parabacteroides distasonis		2.95	0.41	TRE 88		
Bacteroides thetaiotaomicron		0.42	0.11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47555700	**Acceptable identification**						**47555700**
					GRA	SPO	
Bacteroides vulgatus		91.55	0.44	CEL 5	1	1	
Clostridium butyricum		2.96	0.20	NAG 99	99	99	
Eubacterium contortum		1.97	0.23		99	1	
47555702	**Low discrimination**						**47555702**
					GRA		
Clostridium clostridioforme		71.22	0.59	SAL 88	50		
Bifidobacterium adolescentis		16.67	0.52	TRE 88	99		
Parabacteroides distasonis		5.51	0.46		1		
47555710	**Low discrimination**						**47555710**
					GRA		
Clostridium clostridioforme		71.64	0.36	SAL 88	50		
Bacteroides vulgatus		11.34	0.10	TRE 88	1		
Bifidobacterium longum subsp. longum		7.93	0.21	BGL 99	99		
47555712	**Good identification**						**47555712**
					GRA		
Clostridium clostridioforme		93.03	0.70	SAL 88	50		
Bifidobacterium adolescentis		5.84	0.53	TRE 88	99		
Parabacteroides distasonis		0.85	0.41		1		
47555713	**Acceptable identification**						**47555713**
Clostridium clostridioforme		94.09	0.36	SAL 88			
Bifidobacterium adolescentis		5.91	0.19	TRE 88			
47555720	**Very good identification**						**47555720**
Bacteroides vulgatus		99.77	0.78	CEL 5			
Parabacteroides distasonis		0.11	0.46				
Bacteroides uniformis		0.04	0.19				
47555721	**Good identification**						**47555721**
Bacteroides vulgatus		99.89	0.44	CEL 5			
Parabacteroides distasonis		0.11	0.12	URE 1			
47555722	**Low discrimination**						**47555722**
Parabacteroides distasonis		57.03	0.80				
Bacteroides uniformis		18.95	0.53				
Clostridium clostridioforme		7.45	0.59				
47555723	**Low discrimination**						**47555723**
Parabacteroides distasonis		57.29	0.46	URE 1			
Bacteroides uniformis		19.04	0.19				
Clostridium clostridioforme		7.48	0.25				
47555730	**Low discrimination**						**47555730**
Bacteroides vulgatus		86.38	0.44	CEL 5			
Clostridium clostridioforme		5.51	0.36	MLZ 1			
Parabacteroides distasonis		4.99	0.41				
47555732	**Low discrimination**						**47555732**
Clostridium clostridioforme		48.95	0.70	SAL 88			
Parabacteroides distasonis		44.30	0.75	TRE 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		3.14	0.40				
47555733	**Low discrimination**						**47555733**
Clostridium clostridioforme		49.29	0.36	SAL 88			
Parabacteroides distasonis		44.60	0.41	TRE 88			
Bacteroides thetaiotaomicron		3.16	0.05	URE 1			
47555742	**Low discrimination**						**47555742**
Bifidobacterium adolescentis		61.12	0.36	RHA 1			
Clostridium clostridioforme		21.34	0.25	SOR 11			
Actinomyces israelii		14.22	0.31				
47555752	**Low discrimination**						**47555752**
Clostridium clostridioforme		55.86	0.36	SAL 88			
Bifidobacterium adolescentis		42.92	0.37	TRE 88			
Actinomyces israelii		1.22	0.17	SOR 1			
47555760	**Good identification**						**47555760**
Bacteroides vulgatus		99.89	0.44	CEL 5			
Parabacteroides distasonis		0.11	0.12	SOR 1			
47555762	**Low discrimination**						**47555762**
Parabacteroides distasonis		50.58	0.46	SOR 1			
Bacteroides uniformis		16.81	0.19				
Bacteroides ovatus		11.12	0.19				
47555772	**Low discrimination**						**47555772**
Clostridium clostridioforme		45.14	0.36	SAL 88			
Parabacteroides distasonis		40.85	0.41	TRE 88			
Bacteroides ovatus		11.12	0.16	SOR 1			
47556102	**Low discrimination**						**47556102**
					CAT	GRA	
Bifidobacterium adolescentis		62.70	0.36	GAL 99	1	99	
Actinomyces israelii		35.25	0.38	ARA 87	1	99	
Bacteroides fragilis		2.04	-.09		99	1	
47556112	**Acceptable identification**						**47556112**
Bifidobacterium adolescentis		93.57	0.37	GAL 99			
Actinomyces israelii		6.43	0.24	ARA 87			
47556502	**Acceptable identification**						**47556502**
Bifidobacterium adolescentis		93.57	0.50	GAL 99			
Actinomyces israelii		6.43	0.36				
47556512	**Good identification**						**47556512**
Bifidobacterium adolescentis		97.86	0.52	GAL 99			
Clostridium clostridioforme		0.87	0.13				
Actinomyces israelii		0.82	0.22				
47556552	**Good identification**						**47556552**
Bifidobacterium adolescentis		97.42	0.36	GAL 99			
Actinomyces israelii		2.58	0.15	SOR 11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47557000	**Very good identification**						**47557000**
					GRA		
Actinomyces israelii		99.40	0.59	SAL 92	99		
Actinomyces naeslundii		0.35	0.26		99		
Fusobacterium mortiferum		0.25	0.12		1		
47557002	**Good identification**						**47557002**
					GRA		
Actinomyces israelii		92.57	0.62	SAL 92	99		
Bifidobacterium adolescentis		6.65	0.36		99		
Prevotella oralis		0.35	0.04		1		
47557010	**Good identification**						**47557010**
Actinomyces israelii		100.0	0.45	SAL 92			
47557012	**Low discrimination**						**47557012**
Actinomyces israelii		62.98	0.48	SAL 92			
Bifidobacterium adolescentis		37.02	0.37	MLZ 13			
47557020	**Low discrimination**						**47557020**
					CAT	GRA	
Actinomyces israelii		69.93	0.38	SAL 92	1	99	
Bacteroides fragilis		20.05	0.04	NAG 6	99	1	
Prevotella oralis		6.16	0.04		1	1	
47557040	**Good identification**						**47557040**
Actinomyces israelii		100.0	0.52	SAL 92			
47557042	**Good identification**						**47557042**
Actinomyces israelii		97.77	0.55	SAL 92			
Bifidobacterium adolescentis		2.23	0.20				
47557050	**Good identification**						**47557050**
Actinomyces israelii		100.0	0.38	SAL 92			
47557052	**Low discrimination**						**47557052**
Actinomyces israelii		84.26	0.41	SAL 92			
Bifidobacterium adolescentis		15.74	0.22	MLZ 13			
47557100	**Very good identification**						**47557100**
					SPO		
Actinomyces israelii		95.32	0.69	SAL 92	1		
Bifidobacterium adolescentis		2.57	0.36		1		
Clostridium butyricum		0.84	0.13		99		
47557102	**Low discrimination**						**47557102**
					CAT	GRA	
Bifidobacterium adolescentis		61.78	0.70	ARA 87	1	99	
Actinomyces israelii		34.74	0.72		1	99	
Bacteroides fragilis		2.01	0.25		99	1	
47557103	**Low discrimination**						**47557103**
Bifidobacterium adolescentis		57.65	0.36	ARA 87			
Actinomyces israelii		32.42	0.38	URE 1			
Actinomyces naeslundii		8.05	0.37				
47557110	**Low discrimination**						**47557110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		81.35	0.55	SAL 92			
Bifidobacterium adolescentis		17.93	0.37	MLZ 13			
Bifidobacterium breve		0.41	0.10				
47557112	**Good identification**						**47557112**
Bifidobacterium adolescentis		93.19	0.72	ARA 87			
Actinomyces israelii		6.41	0.58				
Bifidobacterium breve		0.22	0.27				
47557113	**Acceptable identification**						**47557113**
Bifidobacterium adolescentis		93.57	0.37	ARA 87			
Actinomyces israelii		6.43	0.24	URE 1			
47557120	**Low discrimination**						**47557120**
					GRA	CAT	
Bacteroides fragilis		77.46	0.38	MAN 1	1	99	
Actinomyces israelii		10.92	0.49	BGL 95	99	1	
Capnocytophaga ochracea		4.59	0.26		1	90	
47557122	**Good identification**						**47557122**
					CAT		
Bacteroides fragilis		91.03	0.60	MAN 1	99		
Prevotella oralis		4.43	0.46		1		
Capnocytophaga ochracea		2.55	0.42		90		
47557140	**Very good identification**						**47557140**
Actinomyces israelii		98.77	0.62	SAL 92			
Bifidobacterium adolescentis		0.85	0.20				
Bifidobacterium breve		0.19	0.10				
47557142	**Low discrimination**						**47557142**
Actinomyces israelii		63.09	0.65	SAL 92			
Bifidobacterium adolescentis		35.66	0.54				
Bifidobacterium breve		0.83	0.27				
47557150	**Acceptable identification**						**47557150**
					COC		
Actinomyces israelii		90.92	0.48	SAL 92	1		
Bifidobacterium adolescentis		6.37	0.22	MLZ 13	1		
Blautia producta		1.53	-.03		99		
47557152	**Low discrimination**						**47557152**
Bifidobacterium adolescentis		80.96	0.56	ARA 87			
Actinomyces israelii		17.51	0.51	SOR 11			
Bifidobacterium breve		1.54	0.27				
47557160	**Low discrimination**						**47557160**
					GRA	CAT	
Actinomyces israelii		73.61	0.42	SAL 92	99	1	
Bacteroides fragilis		13.57	0.04	NAG 6	1	99	
Capnocytophaga ochracea		8.84	0.10		1	90	
47557200	**Good identification**						**47557200**
Actinomyces israelii		100.0	0.39	SAL 92			
47557202	**Good identification**						**47557202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces israelii		97.91	0.43	SAL 92	99	
Prevotella oralis		2.09	-.03	RHA 7	1	
47557242	**Good identification**					**47557242**
Actinomyces israelii		100.0	0.35	SAL 92		
47557300	**Good identification**					**47557300**
Actinomyces israelii		100.0	0.50	SAL 92		
47557302	**Low discrimination**					**47557302**
					GRA	
Actinomyces israelii		74.29	0.53	SAL 92	99	
Bifidobacterium adolescentis		17.73	0.36	RHA 7	99	
Prevotella oralis		6.21	0.18		1	
47557310	**Good identification**					**47557310**
Actinomyces israelii		100.0	0.36	SAL 92		
47557312	**Low discrimination**					**47557312**
Bifidobacterium adolescentis		61.89	0.37	RHA 1		
Actinomyces israelii		31.70	0.39	ARA 87		
Clostridium clostridioforme		5.20	0.16			
47557322	**Low discrimination**					**47557322**
					CAT	
Prevotella oralis		44.28	0.40	MAN 1	1	
Parabacteroides distasonis		29.33	0.49	XYL 11	76	
Bacteroides fragilis		21.46	0.25		99	
47557332	**Low discrimination**					**47557332**
Parabacteroides distasonis		87.46	0.45	MAN 1		
Clostridium clostridioforme		3.78	0.16			
Bacteroides ovatus		3.61	0.05			
47557340	**Good identification**					**47557340**
Actinomyces israelii		100.0	0.43	SAL 92		
47557342	**Acceptable identification**					**47557342**
Actinomyces israelii		92.95	0.46	SAL 92		
Bifidobacterium adolescentis		7.05	0.20	RHA 7		
47557400	**Very good identification**					**47557400**
					SPO	
Actinomyces israelii		97.57	0.57	SAL 92	1	
Clostridium butyricum		1.57	0.05		99	
Bifidobacterium adolescentis		0.87	0.16		1	
47557402	**Low discrimination**					**47557402**
					SPO	
Actinomyces israelii		62.82	0.60	SAL 92	1	
Bifidobacterium adolescentis		36.92	0.50		1	
Clostridium butyricum		0.26	-.02		99	
47557410	**Acceptable identification**					**47557410**
Actinomyces israelii		93.21	0.43	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		6.79	0.17	MLZ 13			
47557412	**Low discrimination**						**47557412**
Bifidobacterium adolescentis		81.94	0.52	LAC 99			
Actinomyces israelii		17.04	0.46				
Clostridium clostridioforme		0.66	0.12				
47557420	**Acceptable identification**						**47557420**
					GRA		
Actinomyces israelii		94.49	0.37	SAL 92	99		
Bacteroides vulgatus		5.51	-.07	NAG 6	1		
47557422	**Low discrimination**						**47557422**
					CAT	GRA	
Actinomyces israelii		87.88	0.40	SAL 92	1	99	
Bifidobacterium adolescentis		8.18	0.16	NAG 6	1	99	
Bacteroides fragilis		3.94	-.09		99	1	
47557440	**Good identification**						**47557440**
Actinomyces israelii		100.0	0.50	SAL 92			
47557442	**Low discrimination**						**47557442**
Actinomyces israelii		84.26	0.53	SAL 92			
Bifidobacterium adolescentis		15.74	0.34				
47557450	**Acceptable identification**						**47557450**
					COC		
Actinomyces israelii		92.42	0.36	SAL 92	1		
Blautia producta		7.58	-.03	MLZ 13	99		
47557452	**Low discrimination**						**47557452**
Bifidobacterium adolescentis		60.44	0.36	LAC 99			
Actinomyces israelii		39.56	0.39	SOR 11			
47557500	**Low discrimination**						**47557500**
					SPO		
Actinomyces israelii		75.69	0.68	SAL 92	1		
Bifidobacterium adolescentis		16.68	0.50		1		
Clostridium butyricum		7.29	0.29		99		
47557502	**Good identification**						**47557502**
					SPO		
Bifidobacterium adolescentis		93.37	0.84		1		
Actinomyces israelii		6.42	0.71		1		
Clostridium butyricum		0.16	0.22		99		
47557503	**Acceptable identification**						**47557503**
Bifidobacterium adolescentis		93.57	0.50	URE 1			
Actinomyces israelii		6.43	0.36				
47557510	**Low discrimination**						**47557510**
Bifidobacterium adolescentis		60.84	0.52	BGL 99			
Actinomyces israelii		33.75	0.53				
Bifidobacterium dentium		2.67	0.10				
47557512	**Very good identification**						**47557512**
Bifidobacterium adolescentis		98.67	0.86				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.83	0.56				
Bifidobacterium dentium		0.44	0.27				
47557513	**Good identification**						**47557513**
Bifidobacterium adolescentis		98.73	0.52	URE 1			
Actinomyces israelii		0.83	0.22				
Bifidobacterium dentium		0.44	-.07				
47557522	**Low discrimination**						**47557522**
					CAT	GRA	
Bifidobacterium adolescentis		39.32	0.50	NAG 1	1	99	
Bacteroides fragilis		18.95	0.25		99	1	
Actinomyces israelii		17.08	0.50		1	99	
47557532	**Low discrimination**						**47557532**
Bifidobacterium adolescentis		48.51	0.52	NAG 1			
Bifidobacterium dentium		38.29	0.32				
Bifidobacterium longum subsp. longum		4.22	0.32				
47557540	**Low discrimination**						**47557540**
					COC		
Actinomyces israelii		83.85	0.61	SAL 92	1		
Blautia producta		8.47	0.24		99		
Bifidobacterium adolescentis		5.87	0.34		1		
47557542	**Low discrimination**						**47557542**
					COC		
Bifidobacterium adolescentis		82.03	0.69	SOR 11	1		
Actinomyces israelii		17.74	0.64		1		
Blautia producta		0.13	0.07		99		
47557552	**Very good identification**						**47557552**
					COC		
Bifidobacterium adolescentis		97.01	0.70	SOR 11	1		
Actinomyces israelii		2.57	0.49		1		
Blautia producta		0.39	0.15		99		
47557553	**Good identification**						**47557553**
Bifidobacterium adolescentis		97.42	0.36	SOR 11			
Actinomyces israelii		2.58	0.15	URE 1			
47557560	**Low discrimination**						**47557560**
					COC	GRA	
Actinomyces israelii		80.46	0.40	SAL 92	1	99	
Blautia producta		14.16	0.07	NAG 6	99	99	
Bacteroides vulgatus		3.02	-.07		1	1	
47557562	**Low discrimination**						**47557562**
					GRA	CAT	
Actinomyces israelii		52.20	0.43	SAL 92	99	1	
Bifidobacterium adolescentis		38.19	0.34	NAG 6	99	1	
Capnocytophaga ochracea		8.11	0.13		1	90	
47557572	**Low discrimination**						**47557572**
Bifidobacterium adolescentis		78.41	0.36	NAG 1			
Actinomyces israelii		13.10	0.29	SOR 11			
Bifidobacterium dentium		5.06	-.03				
47557600	**Good identification**						**47557600**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.38	SAL 92			
47557602	**Low discrimination**						**47557602**
Actinomyces israelii		83.20	0.41	SAL 92			
Clostridium clostridioforme		10.24	0.16	RHA 7			
Bifidobacterium adolescentis		6.56	0.16				
47557700	**Low discrimination**						**47557700**
					GRA	SPO	
Actinomyces israelii		84.71	0.48	SAL 92	99	1	
Bacteroides vulgatus		9.43	0.10	RHA 7	1	1	
Clostridium butyricum		3.35	0.03		99	99	
47557702	**Low discrimination**						**47557702**
Bifidobacterium adolescentis		63.23	0.50	RHA 1			
Actinomyces israelii		32.39	0.51				
Clostridium clostridioforme		3.34	0.25				
47557712	**Low discrimination**						**47557712**
Bifidobacterium adolescentis		78.89	0.52	RHA 1			
Clostridium clostridioforme		15.51	0.36				
Actinomyces israelii		4.94	0.37				
47557720	**Good identification**						**47557720**
					GRA		
Bacteroides vulgatus		99.13	0.44	CEL 5	1		
Actinomyces israelii		0.57	0.28	MAN 1	99		
Bacteroides ovatus		0.19	0.00		1		
47557742	**Low discrimination**						**47557742**
					COC		
Actinomyces israelii		58.14	0.44	SAL 92	1		
Bifidobacterium adolescentis		36.08	0.34	RHA 7	1		
Blautia producta		5.78	0.07		99		
47557752	**Low discrimination**						**47557752**
					COC		
Bifidobacterium adolescentis		62.87	0.36	RHA 1	1		
Blautia producta		24.74	0.15	SOR 11	99		
Actinomyces israelii		12.39	0.30		1		
47561000	**Low discrimination**						**47561000**
					GRA		
Actinomyces naeslundii		51.74	0.38	ESL 12	99		
Fusobacterium mortiferum		27.63	0.22	FRU 95	1		
Lactobacillus jensenii		20.63	0.10		99		
47561001	**Good identification**						**47561001**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47561002	**Low discrimination**						**47561002**
					GRA		
Actinomyces naeslundii		70.23	0.38	ESL 12	99		
Lactobacillus jensenii		18.67	0.07	FRU 95	99		
Prevotella oralis		11.10	-.01		1		
47561003	**Good identification**						**47561003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
47561020	**Low discrimination**					**47561020**
					GRA	
Actinomyces naeslundii		64.21	0.38	ESL 12	99	
Clostridium perfringens		14.43	0.05	FRU 95	99	
Prevotella melaninogenica		11.21	-.04		1	
47561021	**Good identification**					**47561021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47561023	**Good identification**					**47561023**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47561101	**Acceptable identification**					**47561101**
					SPO	
Actinomyces naeslundii		93.55	0.49	ESL 12	1	
Clostridium chauvoei		6.45	0.06	FRU 95	99	
47561102	**Low discrimination**					**47561102**
					GRA	
Actinomyces naeslundii		30.20	0.46	ESL 12	99	
Prevotella oralis		26.27	0.19	FRU 95	1	
Bifidobacterium breve		11.77	0.24		99	
47561103	**Good identification**					**47561103**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47561121	**Good identification**					**47561121**
					GRA	
Actinomyces naeslundii		96.07	0.49	ESL 12	99	
Prevotella melaninogenica		3.93	-.04	FRU 95	1	
47561122	**Low discrimination**					**47561122**
Prevotella oralis		71.88	0.41	FRU 99		
Parabacteroides distasonis		6.67	0.36	TRE 6		
Prevotella melaninogenica		5.03	0.18			
47561123	**Low discrimination**					**47561123**
					GRA	
Actinomyces naeslundii		89.98	0.49	ESL 12	99	
Prevotella oralis		10.02	0.07	FRU 95	1	
47561141	**Good identification**					**47561141**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47561143	**Good identification**					**47561143**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47561161	**Good identification**					**47561161**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47561163	**Good identification**					**47561163**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47561332	**Acceptable identification**					**47561332**
Parabacteroides distasonis		91.67	0.40	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		5.68	0.13	XYL 90		
Prevotella oralis		2.65	0.01			
47561512	**Low discrimination**					**47561512**
Bifidobacterium adolescentis		74.22	0.36	FRU 99		
Bifidobacterium longum subsp. longum		12.15	0.21	XYL 87		
Clostridium clostridioforme		11.85	0.19			
47561722	**Low discrimination**					**47561722**
Parabacteroides distasonis		75.77	0.41	FRU 99		
Clostridium clostridioforme		14.18	0.23	XYL 90		
Bacteroides thetaiotaomicron		8.13	0.08			
47563000	**Acceptable identification**					**47563000**
Actinomyces naeslundii		91.52	0.43	ESL 12		
Actinomyces israelii		8.48	0.16	FRU 95		
47563001	**Good identification**					**47563001**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47563002	**Low discrimination**					**47563002**
Actinomyces naeslundii		87.80	0.43	ESL 12		
Actinomyces israelii		12.20	0.19	FRU 95		
47563003	**Good identification**					**47563003**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47563020	**Good identification**					**47563020**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47563021	**Good identification**					**47563021**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47563022	**Good identification**					**47563022**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47563023	**Good identification**					**47563023**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47563100	**Low discrimination**					**47563100**
Actinomyces naeslundii		88.48	0.51	ESL 12		
Actinomyces israelii		11.52	0.27	FRU 95		
47563101	**Good identification**					**47563101**
Actinomyces naeslundii		100.0	0.54	ESL 12		
47563102	**Low discrimination**					**47563102**
Actinomyces naeslundii		66.59	0.51	ESL 12		
Bifidobacterium breve		13.37	0.24	FRU 95		
Actinomyces israelii		13.00	0.30			
47563103	**Good identification**					**47563103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.54	ESL 12		
47563120	**Low discrimination**					**47563120**
					SPO	
Actinomyces naeslundii		66.91	0.51	ESL 12	1	
Clostridium ramosum		28.62	0.19	FRU 95	92	
Bifidobacterium breve		2.36	0.12		1	
47563121	**Good identification**					**47563121**
Actinomyces naeslundii		100.0	0.54	ESL 12		
47563123	**Good identification**					**47563123**
Actinomyces naeslundii		100.0	0.54	ESL 12		
47563140	**Low discrimination**					**47563140**
Actinomyces naeslundii		77.71	0.38	ESL 12		
Actinomyces israelii		22.29	0.20	FRU 95		
47563141	**Good identification**					**47563141**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47563143	**Good identification**					**47563143**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47563160	**Low discrimination**					**47563160**
Actinomyces naeslundii		83.34	0.38	ESL 12		
Bifidobacterium breve		16.67	0.12	FRU 95		
47563161	**Good identification**					**47563161**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47563163	**Good identification**					**47563163**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47564712	**Good identification**					**47564712**
Clostridium clostridioforme		100.0	0.43	SAL 88		
47564732	**Good identification**					**47564732**
Clostridium clostridioforme		98.33	0.43	SAL 88		
Parabacteroides distasonis		1.68	0.18	FRU 99		
47565112	**Low discrimination**					**47565112**
Bifidobacterium adolescentis		82.00	0.36	FRU 99		
Clostridium clostridioforme		8.45	0.16	ARA 87		
Actinomyces israelii		4.59	0.21			
47565122	**Low discrimination**					**47565122**
					CAT	
Parabacteroides distasonis		60.39	0.53	FRU 99	76	
Bacteroides fragilis		29.10	0.25		99	
Prevotella oralis		8.94	0.26		1	
47565132	**Good identification**					**47565132**
Parabacteroides distasonis		96.38	0.48	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		2.74	0.16				
Bacteroides fragilis		0.87	-.09				
47565322	**Very good identification**						**47565322**
Parabacteroides distasonis		96.62	0.61	FRU 99			
Prevotella oralis		2.04	0.19				
Clostridium clostridioforme		0.80	0.19				
47565332	**Acceptable identification**						**47565332**
Parabacteroides distasonis		93.25	0.56	FRU 99			
Clostridium clostridioforme		6.49	0.31				
Bacteroides ovatus		0.26	-.04				
47565502	**Low discrimination**						**47565502**
					SPO		
Bifidobacterium adolescentis		67.65	0.48	FRU 99	1		
Clostridium butyricum		20.10	0.25		99		
Clostridium clostridioforme		4.38	0.25		66		
47565512	**Low discrimination**						**47565512**
Bifidobacterium adolescentis		74.45	0.50	FRU 99			
Clostridium clostridioforme		17.96	0.36				
Bifidobacterium longum subsp. longum		5.47	0.29				
47565520	**Low discrimination**						**47565520**
					GRA	SPO	
Bifidobacterium longum subsp. longum		68.84	0.35	FRU 90	99	1	
Clostridium butyricum		21.70	0.14	CEL 1	99	99	
Bacteroides vulgatus		6.99	0.04		1	1	
47565522	**Low discrimination**						**47565522**
					GRA		
Parabacteroides distasonis		66.44	0.49	FRU 99	1		
Bifidobacterium longum subsp. longum		18.61	0.35		99		
Clostridium clostridioforme		5.72	0.25		50		
47565530	**Good identification**						**47565530**
					GRA		
Bifidobacterium longum subsp. longum		98.87	0.39	FRU 90	99		
Parabacteroides distasonis		1.13	0.10	CEL 1	1		
47565532	**Low discrimination**						**47565532**
					GRA		
Parabacteroides distasonis		36.45	0.44	FRU 99	1		
Bifidobacterium longum subsp. longum		32.29	0.39		99		
Clostridium clostridioforme		26.53	0.36		50		
47565612	**Good identification**						**47565612**
Clostridium clostridioforme		100.0	0.42	SAL 88			
47565632	**Good identification**						**47565632**
Clostridium clostridioforme		97.38	0.42	SAL 88			
Parabacteroides distasonis		1.85	0.18	FRU 99			
Bacteroides thetaiotaomicron		0.77	-.04				
47565702	**Low discrimination**						**47565702**
					GRA		
Clostridium clostridioforme		89.99	0.40	SAL 88	50		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		4.32	0.23	FRU 99	1	
Bifidobacterium adolescentis		1.92	0.14		99	
47565712	**Good identification**					**47565712**
					GRA	
Clostridium clostridioforme		98.87	0.51	SAL 88	50	
Bifidobacterium adolescentis		0.57	0.16	FRU 99	99	
Parabacteroides distasonis		0.56	0.18		1	
47565722	**Low discrimination**					**47565722**
Parabacteroides distasonis		61.94	0.57	FRU 99		
Bacteroides thetaiotaomicron		17.71	0.32			
Clostridium clostridioforme		13.02	0.40			
47565732	**Low discrimination**					**47565732**
Clostridium clostridioforme		56.80	0.51	SAL 88		
Parabacteroides distasonis		31.93	0.52	FRU 99		
Bacteroides thetaiotaomicron		6.60	0.25			
47567002	**Good identification**					**47567002**
Actinomyces israelii		100.0	0.37	SAL 92		
47567100	**Good identification**					**47567100**
					SPO	
Actinomyces israelii		96.48	0.45	SAL 92	1	
Clostridium butyricum		3.52	-.01	FRU 99	99	
47567102	**Low discrimination**					**47567102**
Actinomyces israelii		76.04	0.48	SAL 92		
Bifidobacterium adolescentis		23.97	0.33	FRU 99		
47567140	**Good identification**					**47567140**
Actinomyces israelii		100.0	0.38	SAL 92		
47567142	**Acceptable identification**					**47567142**
Actinomyces israelii		90.90	0.41	SAL 92		
Bifidobacterium adolescentis		9.11	0.17	FRU 99		
47567402	**Acceptable identification**					**47567402**
Actinomyces israelii		90.57	0.36	SAL 92		
Bifidobacterium adolescentis		9.43	0.13	FRU 99		
47567500	**Low discrimination**					**47567500**
					SPO	
Actinomyces israelii		69.43	0.43	SAL 92	1	
Clostridium butyricum		27.86	0.15	FRU 99	99	
Bifidobacterium adolescentis		2.71	0.13		1	
47567502	**Low discrimination**					**47567502**
					SPO	
Bifidobacterium adolescentis		69.62	0.47	FRU 99	1	
Actinomyces israelii		27.00	0.46		1	
Clostridium butyricum		2.81	0.08		99	
47567512	**Low discrimination**					**47567512**
Bifidobacterium adolescentis		58.36	0.48	FRU 99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		38.87	0.27				
Actinomyces israelii		2.77	0.32				
47567542	**Low discrimination**						**47567542**
					COC		
Actinomyces israelii		49.59	0.39	SAL 92	1		
Bifidobacterium adolescentis		40.64	0.31	FRU 99	1		
Blautia producta		9.77	0.07		99		
47570000	**Acceptable identification**						**47570000**
					GRA	CAT	
Fusobacterium mortiferum		93.19	0.53	MLT 11	1	5	
Propionibacterium granulosum		3.66	0.26		99	90	
Actinomyces naeslundii		2.09	0.36		99	25	
47570001	**Low discrimination**						**47570001**
					GRA		
Actinomyces naeslundii		76.94	0.39	ESL 12	99		
Fusobacterium mortiferum		23.06	0.19	GAL 96	1		
47570002	**Low discrimination**						**47570002**
					GRA	CAT	
Fusobacterium mortiferum		68.37	0.37	MLT 11	1	5	
Actinomyces naeslundii		13.82	0.36	BGL 10	99	25	
Propionibacterium granulosum		6.04	0.15		99	90	
47570003	**Good identification**						**47570003**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47570020	**Low discrimination**						**47570020**
					GRA	CAT	
Actinomyces naeslundii		42.19	0.36	ESL 12	99	25	
Fusobacterium mortiferum		18.97	0.19	GAL 96	1	5	
Propionibacterium granulosum		18.45	0.15		99	90	
47570021	**Good identification**						**47570021**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47570023	**Good identification**						**47570023**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47570100	**Good identification**						**47570100**
					COC	GRA	
Fusobacterium mortiferum		92.93	0.59	MLT 11	1	1	
Atopobium parvulum		4.24	0.25		99	99	
Actinomyces naeslundii		2.55	0.44		1	99	
47570101	**Low discrimination**						**47570101**
					GRA		
Actinomyces naeslundii		80.27	0.47	ESL 12	99		
Fusobacterium mortiferum		19.73	0.25	GAL 96	1		
47570102	**Low discrimination**						**47570102**
					GRA		
Fusobacterium mortiferum		49.45	0.43	MLT 11	1		
Prevotella oralis		13.40	0.19	BGL 10	1		
Actinomyces naeslundii		12.19	0.44		99		
47570103	**Good identification**						**47570103**
Actinomyces naeslundii		100.0	0.47	ESL 12			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47570121	**Good identification**					**47570121**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47570122	**Low discrimination**					**47570122**
					GRA	
Prevotella oralis		75.52	0.41	TRE 6	1	
Parabacteroides distasonis		7.00	0.36	GAL 99	1	
Actinomyces naeslundii		3.62	0.44		99	
47570123	**Low discrimination**					**47570123**
					GRA	
Actinomyces naeslundii		87.67	0.47	ESL 12	99	
Prevotella oralis		12.33	0.07	GAL 96	1	
47570312	**Good identification**					**47570312**
Clostridium clostridioforme		99.04	0.39	SAL 88		
Leptotrichia buccalis		0.96	-.11	XYL 91		
47570332	**Low discrimination**					**47570332**
Clostridium clostridioforme		66.52	0.39	SAL 88		
Parabacteroides distasonis		32.54	0.40	XYL 91		
Prevotella oralis		0.94	0.01	ARA 94		
47570412	**Good identification**					**47570412**
Clostridium clostridioforme		100.0	0.36	SAL 88		
47570432	**Good identification**					**47570432**
Clostridium clostridioforme		100.0	0.36	SAL 88		
47570512	**Low discrimination**					**47570512**
					GRA	
Clostridium clostridioforme		83.73	0.45	SAL 88	50	
Bifidobacterium adolescentis		15.89	0.36	XYL 91	99	
Leptotrichia buccalis		0.38	-.11	RHA 88	1	
47570532	**Acceptable identification**					**47570532**
					GRA	
Clostridium clostridioforme		94.51	0.45	SAL 88	50	
Parabacteroides distasonis		4.42	0.28	XYL 91	1	
Bifidobacterium longum subsp. longum		1.07	0.13	RHA 88	99	
47570602	**Good identification**					**47570602**
Clostridium clostridioforme		100.0	0.40	SAL 88		
47570612	**Good identification**					**47570612**
Clostridium clostridioforme		100.0	0.51	SAL 88		
47570622	**Good identification**					**47570622**
Clostridium clostridioforme		100.0	0.40	SAL 88		
47570632	**Good identification**					**47570632**
Clostridium clostridioforme		100.0	0.51	SAL 88		
47570702	**Good identification**					**47570702**
Clostridium clostridioforme		100.0	0.49	SAL 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47570712	**Very good identification**					**47570712**
Clostridium clostridioforme		100.0	0.60	SAL 88		
47570722	**Low discrimination**					**47570722**
Clostridium clostridioforme		84.50	0.49	SAL 88		
Parabacteroides distasonis		13.68	0.41	XYL 91		
Bacteroides thetaiotaomicron		1.47	0.08			
47570732	**Very good identification**					**47570732**
Clostridium clostridioforme		97.98	0.60	SAL 88		
Parabacteroides distasonis		1.88	0.36	XYL 91		
Bacteroides thetaiotaomicron		0.15	0.01			
47571000	**Low discrimination**					**47571000**
					GRA	
Fusobacterium mortiferum		43.52	0.56	MLT 11	1	
Lactobacillus jensenii		32.49	0.44		99	
Actinomyces naeslundii		15.64	0.60		99	
47571001	**Very good identification**					**47571001**
					GRA	
Actinomyces naeslundii		96.83	0.63	ESL 12	99	
Fusobacterium mortiferum		1.81	0.22		1	
Lactobacillus jensenii		1.35	0.10		99	
47571002	**Low discrimination**					**47571002**
					GRA	
Lactobacillus jensenii		31.47	0.41	SAL 95	99	
Actinomyces naeslundii		22.73	0.60	RAF 1	99	
Prevotella oralis		18.72	0.33		1	
47571003	**Very good identification**					**47571003**
					GRA	
Actinomyces naeslundii		98.54	0.63	ESL 12	99	
Lactobacillus jensenii		0.92	0.07		99	
Prevotella oralis		0.55	-.01		1	
47571010	**Low discrimination**					**47571010**
					GRA	
Lactobacillus jensenii		84.94	0.36	SAL 95	99	
Actinomyces naeslundii		6.46	0.38	RAF 1	99	
Fusobacterium mortiferum		3.45	0.22		1	
47571011	**Acceptable identification**					**47571011**
Actinomyces naeslundii		91.86	0.41	ESL 12		
Lactobacillus jensenii		8.14	0.02	MLZ 5		
47571013	**Acceptable identification**					**47571013**
					GRA	
Actinomyces naeslundii		91.46	0.41	ESL 12	99	
Leptotrichia buccalis		8.54	-.11	MLZ 5	1	
47571020	**Low discrimination**					**47571020**
					GRA	
Clostridium perfringens		31.04	0.39	CEL 4	99	
Actinomyces naeslundii		26.50	0.60	LAC 88	99	
Prevotella oralis		21.83	0.33		1	
47571021	**Very good identification**					**47571021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		98.68	0.63	ESL 12	99		
Clostridium perfringens		0.78	0.05		99		
Prevotella oralis		0.55	-.01		1		
47571022	**Low discrimination**						**47571022**
					GRA		
Prevotella oralis		84.17	0.55	TRE 6	1		
Actinomyces naeslundii		5.38	0.60	LAC 94	99		
Clostridium perfringens		2.97	0.33		99		
47571023	**Good identification**						**47571023**
					GRA		
Actinomyces naeslundii		90.47	0.63	ESL 12	99		
Prevotella oralis		9.53	0.21		1		
47571031	**Good identification**						**47571031**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47571033	**Good identification**						**47571033**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47571040	**Low discrimination**						**47571040**
					GRA		
Actinomyces naeslundii		59.47	0.47	ESL 12	99		
Actinomyces israelii		21.67	0.31		99		
Fusobacterium mortiferum		9.47	0.22		1		
47571041	**Good identification**						**47571041**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47571042	**Low discrimination**						**47571042**
Actinomyces naeslundii		43.60	0.47	ESL 12			
Actinomyces israelii		23.83	0.34				
Bifidobacterium breve		14.42	0.24				
47571043	**Good identification**						**47571043**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47571061	**Good identification**						**47571061**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47571062	**Low discrimination**						**47571062**
					GRA		
Actinomyces naeslundii		29.08	0.47	ESL 12	99		
Prevotella oralis		26.05	0.21		1		
Clostridium perfringens		17.30	0.21		99		
47571063	**Good identification**						**47571063**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47571100	**Low discrimination**						**47571100**
					COC	GRA	
Atopobium parvulum		61.19	0.59	RAF 1	99	99	
Fusobacterium mortiferum		20.34	0.62		1	1	
Actinomyces naeslundii		8.92	0.68		1	99	
47571101	**Good identification**						**47571101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Actinomyces naeslundii		93.18	0.71	ESL 12	1	99	
Atopobium parvulum		4.31	0.25		99	99	
Fusobacterium mortiferum		1.43	0.28		1	1	
47571102	**Low discrimination**						**47571102**
					GRA		
Prevotella oralis		34.04	0.53	TRE 6	1		
Bifidobacterium breve		15.25	0.59	NAG 95	99		
Lactobacillus acidophilus		12.56	0.47		99		
47571103	**Good identification**						**47571103**
					GRA		
Actinomyces naeslundii		92.37	0.71	ESL 12	99		
Prevotella oralis		2.82	0.19		1		
Bifidobacterium breve		1.26	0.24		99		
47571111	**Good identification**						**47571111**
					GRA		
Actinomyces naeslundii		96.82	0.49	ESL 12	99		
Leptotrichia buccalis		3.18	-.11	MLZ 5	1		
47571112	**Low discrimination**						**47571112**
					GRA		
Leptotrichia buccalis		86.34	0.58	SAL 99	1		
Bifidobacterium breve		6.91	0.59		99		
Bifidobacterium adolescentis		5.44	0.56		99		
47571120	**Low discrimination**						**47571120**
					GRA		
Prevotella oralis		29.08	0.53	TRE 6	1		
Clostridium perfringens		19.36	0.54	BGL 95	99		
Prevotella melaninogenica		18.45	0.47		1		
47571121	**Good identification**						**47571121**
					GRA		
Actinomyces naeslundii		90.97	0.71	ESL 12	99		
Prevotella oralis		2.78	0.19		1		
Clostridium perfringens		1.85	0.20		99		
47571122	**Low discrimination**						**47571122**
					CAT		
Prevotella oralis		77.96	0.75	TRE 6	1		
Parabacteroides distasonis		7.23	0.70		76		
Capnocytophaga ochracea		3.52	0.53		90		
47571123	**Low discrimination**						**47571123**
					GRA		
Actinomyces naeslundii		57.64	0.71	ESL 12	99		
Prevotella oralis		33.43	0.41		1		
Parabacteroides distasonis		3.10	0.36		1		
47571130	**Low discrimination**						**47571130**
					GRA	CAT	
Bifidobacterium breve		52.90	0.46	SAL 99	99	1	
Actinomyces naeslundii		7.80	0.46	BGL 91	99	25	
Capnocytophaga ochracea		7.11	0.20		1	90	
47571131	**Good identification**						**47571131**
Actinomyces naeslundii		95.63	0.49	ESL 12			
Bifidobacterium breve		4.37	0.12	MLZ 5			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47571132	**Low discrimination**					**47571132**
					GRA	
Parabacteroides distasonis		34.98	0.66	XYL 90	1	
Bifidobacterium breve		29.35	0.63		99	
Leptotrichia buccalis		22.93	0.41		1	
47571133	**Low discrimination**					**47571133**
					GRA	
Actinomyces naeslundii		39.38	0.49	ESL 12	99	
Parabacteroides distasonis		21.67	0.32	MLZ 5	1	
Bifidobacterium breve		18.18	0.29		99	
47571140	**Low discrimination**					**47571140**
Bifidobacterium breve		33.53	0.41	SAL 99		
Actinomyces naeslundii		29.47	0.55	BGL 91		
Actinomyces israelii		15.09	0.41			
47571141	**Very good identification**					**47571141**
Actinomyces naeslundii		100.0	0.58	ESL 12		
47571142	**Low discrimination**					**47571142**
Bifidobacterium breve		71.60	0.59	SAL 99		
Lactobacillus acidophilus		7.29	0.32			
Actinomyces naeslundii		6.22	0.55			
47571143	**Good identification**					**47571143**
Actinomyces naeslundii		92.81	0.58	ESL 12		
Bifidobacterium breve		7.19	0.24			
47571150	**Low discrimination**					**47571150**
Bifidobacterium breve		88.78	0.41	SAL 99		
Actinomyces israelii		5.97	0.27	BGL 91		
Actinomyces naeslundii		4.11	0.33			
47571151	**Good identification**					**47571151**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47571152	**Low discrimination**					**47571152**
					GRA	
Bifidobacterium breve		80.60	0.59	SAL 99	99	
Leptotrichia buccalis		10.18	0.24		1	
Bifidobacterium adolescentis		7.84	0.40		99	
47571160	**Low discrimination**					**47571160**
Clostridium perfringens		42.85	0.41	CEL 4		
Bifidobacterium breve		26.61	0.46			
Actinomyces naeslundii		13.16	0.55			
47571161	**Very good identification**					**47571161**
Actinomyces naeslundii		96.57	0.58	ESL 12		
Clostridium perfringens		2.12	0.07			
Bifidobacterium breve		1.32	0.12			
47571162	**Low discrimination**					**47571162**
					GRA CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		62.98	0.63	SAL 99	99	1	
Prevotella oralis		15.18	0.41		1	1	
Capnocytophaga ochracea		7.54	0.36		1	90	
47571163	**Low discrimination**						**47571163**
					GRA		
Actinomyces naeslundii		83.49	0.58	ESL 12	99		
Bifidobacterium breve		11.50	0.29		99		
Prevotella oralis		2.77	0.07		1		
47571170	**Acceptable identification**						**47571170**
Bifidobacterium breve		94.59	0.46	SAL 99			
Actinomyces naeslundii		2.46	0.33	BGL 91			
Clostridium perfringens		1.54	0.07				
47571171	**Low discrimination**						**47571171**
Actinomyces naeslundii		79.44	0.36	ESL 12			
Bifidobacterium breve		20.56	0.12	MLZ 5			
47571172	**Very good identification**						**47571172**
					GRA	CAT	
Bifidobacterium breve		96.25	0.63	SAL 99	99	1	
Capnocytophaga ochracea		1.28	0.20		1	90	
Parabacteroides distasonis		1.16	0.32		1	76	
47571212	**Good identification**						**47571212**
Clostridium clostridioforme		98.93	0.39	SAL 88			
Leptotrichia buccalis		1.07	-.11	XYL 91			
47571222	**Low discrimination**						**47571222**
Prevotella oralis		87.07	0.49	TRE 6			
Parabacteroides distasonis		8.95	0.44	LAC 94			
Clostridium clostridioforme		1.94	0.27				
47571232	**Low discrimination**						**47571232**
Clostridium clostridioforme		60.77	0.39	SAL 88			
Parabacteroides distasonis		33.17	0.40	XYL 91			
Prevotella oralis		6.06	0.14	ARA 94			
47571301	**Good identification**						**47571301**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47571302	**Low discrimination**						**47571302**
Prevotella oralis		74.18	0.47	TRE 6			
Parabacteroides distasonis		9.24	0.44	NAG 95			
Clostridium clostridioforme		6.70	0.36				
47571312	**Low discrimination**						**47571312**
Clostridium clostridioforme		64.14	0.48	SAL 88			
Leptotrichia buccalis		20.57	0.24	XYL 91			
Parabacteroides distasonis		10.46	0.40	ARA 94			
47571320	**Low discrimination**						**47571320**
					SPO		
Prevotella oralis		58.60	0.47	TRE 6	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		30.53	0.43	BGL 95	92	
Parabacteroides distasonis		7.30	0.44		1	
47571322	**Low discrimination**					**47571322**
					SPO	
Prevotella oralis		56.48	0.69	TRE 6	1	
Parabacteroides distasonis		36.66	0.79		1	
Clostridium ramosum		6.19	0.53		92	
47571330	**Low discrimination**					**47571330**
					SPO	
Parabacteroides distasonis		75.84	0.40	XYL 90	1	
Prevotella oralis		11.42	0.13	BGL 99	1	
Clostridium ramosum		5.95	0.09		92	
47571332	**Good identification**					**47571332**
Parabacteroides distasonis		90.90	0.74	XYL 90		
Clostridium clostridioforme		5.63	0.48			
Prevotella oralis		2.63	0.35			
47571333	**Acceptable identification**					**47571333**
Parabacteroides distasonis		91.67	0.40	XYL 90		
Clostridium clostridioforme		5.68	0.13	URE 1		
Prevotella oralis		2.65	0.01			
47571372	**Low discrimination**					**47571372**
					GRA	
Parabacteroides distasonis		72.97	0.40	XYL 90	1	
Bifidobacterium breve		20.41	0.29	SOR 1	99	
Clostridium clostridioforme		4.52	0.13		50	
47571412	**Low discrimination**					**47571412**
Clostridium clostridioforme		80.79	0.44	SAL 88		
Bifidobacterium adolescentis		17.11	0.36	XYL 91		
Actinomyces israelii		1.69	0.25	RHA 88		
47571432	**Acceptable identification**					**47571432**
					GRA	
Clostridium clostridioforme		93.91	0.44	SAL 88	50	
Parabacteroides distasonis		4.90	0.28	XYL 91	1	
Bifidobacterium longum subsp. longum		1.18	0.13	RHA 88	99	
47571500	**Low discrimination**					**47571500**
					SPO	
Clostridium butyricum		73.88	0.40	SAL 99	99	
Actinomyces israelii		10.15	0.46	XYL 97	1	
Bifidobacterium longum subsp. longum		4.99	0.33		1	
47571501	**Low discrimination**					**47571501**
					SPO	
Actinomyces naeslundii		52.02	0.37	ESL 12	1	
Clostridium butyricum		47.98	0.06	ARA 1	99	
47571502	**Low discrimination**					**47571502**
					SPO	
Bifidobacterium adolescentis		85.13	0.69	XYL 87	1	
Clostridium butyricum		5.24	0.33		99	
Clostridium clostridioforme		3.65	0.42		66	
47571510	**Low discrimination**					**47571510**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Bifidobacterium longum subsp. longum		41.00	0.37	CEL 1	1			
Bifidobacterium adolescentis		27.82	0.36		1			
Clostridium butyricum		18.75	0.18		99			
47571512	**Low discrimination**							**47571512**
					GRA			
Bifidobacterium adolescentis		82.60	0.70	XYL 87	99			
Clostridium clostridioforme		13.19	0.53		50			
Leptotrichia buccalis		1.98	0.24		1			
47571513	**Low discrimination**							**47571513**
					GRA			
Bifidobacterium adolescentis		84.49	0.36	XYL 87	99			
Clostridium clostridioforme		13.49	0.19	URE 1	50			
Leptotrichia buccalis		2.03	-.11		1			
47571520	**Low discrimination**							**47571520**
					GRA	SPO	CAT	
Bifidobacterium longum subsp. longum		46.07	0.43	CEL 1	99	1	1	
Clostridium butyricum		14.82	0.22		99	99	1	
Capnocytophaga ochracea		9.39	0.23		1	1	90	
47571521	**Low discrimination**							**47571521**
Actinomyces naeslundii		80.04	0.37	ESL 12				
Bifidobacterium longum subsp. longum		19.96	0.09	ARA 1				
47571522	**Low discrimination**							**47571522**
					CAT			
Parabacteroides distasonis		64.39	0.67	XYL 90	76			
Prevotella oralis		11.44	0.41		1			
Capnocytophaga ochracea		9.02	0.40		90			
47571530	**Acceptable identification**							**47571530**
					GRA			
Bifidobacterium longum subsp. longum		91.09	0.47	CEL 1	99			
Parabacteroides distasonis		3.81	0.28		1			
Clostridium clostridioforme		2.47	0.19		50			
47571532	**Low discrimination**							**47571532**
					GRA			
Parabacteroides distasonis		49.71	0.62	XYL 90	1			
Clostridium clostridioforme		32.21	0.53		50			
Bifidobacterium longum subsp. longum		12.01	0.47		99			
47571542	**Low discrimination**							**47571542**
					SPO			
Bifidobacterium adolescentis		85.03	0.53	XYL 87	1			
Actinomyces israelii		8.72	0.42	SOR 11	1			
Clostridium butyricum		3.68	0.15		99			
47571552	**Good identification**							**47571552**
Bifidobacterium adolescentis		95.70	0.54	XYL 87				
Bifidobacterium breve		1.49	0.24	SOR 11				
Clostridium clostridioforme		1.25	0.19					
47571602	**Good identification**							**47571602**
Clostridium clostridioforme		99.35	0.48	SAL 88				
Actinomyces israelii		0.65	0.20	XYL 91				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47571612	**Very good identification**						**47571612**
Clostridium clostridioforme		100.0	0.59	SAL 88			
47571622	**Low discrimination**						**47571622**
Clostridium clostridioforme		80.08	0.48	SAL 88			
Parabacteroides distasonis		14.47	0.41	XYL 91			
Bacteroides thetaiotaomicron		3.14	0.13				
47571632	**Very good identification**						**47571632**
Clostridium clostridioforme		97.59	0.59	SAL 88			
Parabacteroides distasonis		2.08	0.36	XYL 91			
Bacteroides thetaiotaomicron		0.33	0.06				
47571702	**Low discrimination**						**47571702**
					GRA		
Clostridium clostridioforme		89.86	0.57	SAL 88	50		
Parabacteroides distasonis		4.85	0.41	XYL 91	1		
Bifidobacterium adolescentis		2.89	0.34		99		
47571712	**Very good identification**						**47571712**
					GRA		
Clostridium clostridioforme		98.43	0.68	SAL 88	50		
Bifidobacterium adolescentis		0.85	0.36	XYL 91	99		
Parabacteroides distasonis		0.63	0.36		1		
47571722	**Low discrimination**						**47571722**
Parabacteroides distasonis		74.95	0.75	XYL 90			
Clostridium clostridioforme		14.03	0.57				
Bacteroides thetaiotaomicron		8.04	0.42				
47571723	**Low discrimination**						**47571723**
Parabacteroides distasonis		75.77	0.41	XYL 90			
Clostridium clostridioforme		14.18	0.23	URE 1			
Bacteroides thetaiotaomicron		8.13	0.08				
47571732	**Low discrimination**						**47571732**
Clostridium clostridioforme		59.18	0.68	SAL 88			
Parabacteroides distasonis		37.38	0.70	XYL 91			
Bacteroides thetaiotaomicron		2.90	0.35				
47571762	**Low discrimination**						**47571762**
Parabacteroides distasonis		73.31	0.41	XYL 90			
Clostridium clostridioforme		13.72	0.23	SOR 1			
Bacteroides thetaiotaomicron		7.86	0.08				
47572000	**Low discrimination**						**47572000**
					GRA	CAT	
Actinomyces naeslundii		49.30	0.41	ESL 12	99	25	
Propionibacterium granulosum		33.51	0.24	GAL 96	99	90	
Fusobacterium mortiferum		11.42	0.19		1	5	
47572001	**Good identification**						**47572001**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47572002	**Low discrimination**						**47572002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Actinomyces naeslundii		74.33	0.41	ESL 12	25	
Actinomyces israelii		13.05	0.19	GAL 96	1	
Propionibacterium granulosum		12.63	0.13		90	
47572003	**Good identification**					**47572003**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47572020	**Low discrimination**					**47572020**
Actinomyces naeslundii		85.48	0.41	ESL 12		
Propionibacterium granulosum		14.52	0.13	GAL 96		
47572021	**Good identification**					**47572021**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47572022	**Good identification**					**47572022**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47572023	**Good identification**					**47572023**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47572100	**Low discrimination**					**47572100**
					GRA	
Actinomyces naeslundii		73.84	0.49	ESL 12	99	
Fusobacterium mortiferum		14.02	0.25	GAL 96	1	
Actinomyces israelii		12.14	0.27		99	
47572101	**Good identification**					**47572101**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47572102	**Low discrimination**					**47572102**
Actinomyces naeslundii		61.21	0.49	ESL 12		
Bifidobacterium breve		15.52	0.24	GAL 96		
Actinomyces israelii		15.10	0.30			
47572103	**Good identification**					**47572103**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47572120	**Low discrimination**					**47572120**
					SPO	
Actinomyces naeslundii		63.10	0.49	ESL 12	1	
Clostridium ramosum		34.09	0.19	GAL 96	92	
Bifidobacterium breve		2.81	0.12		1	
47572121	**Good identification**					**47572121**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47572123	**Good identification**					**47572123**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47572140	**Low discrimination**					**47572140**
Actinomyces naeslundii		73.40	0.36	ESL 12		
Actinomyces israelii		26.60	0.20	GAL 96		
47572141	**Good identification**					**47572141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	ESL 12		
47572143	**Good identification**					**47572143**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47572160	**Low discrimination**					**47572160**
Actinomyces naeslundii		79.83	0.36	ESL 12		
Bifidobacterium breve		20.17	0.12	GAL 96		
47572161	**Good identification**					**47572161**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47572163	**Good identification**					**47572163**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47573000	**Low discrimination**					**47573000**
					CAT	
Actinomyces naeslundii		65.09	0.65	ESL 12	25	
Actinomyces israelii		31.42	0.50		1	
Propionibacterium granulosum		1.84	0.24		90	
47573001	**Very good identification**					**47573001**
Actinomyces naeslundii		99.68	0.68	ESL 12		
Actinomyces israelii		0.32	0.16			
47573002	**Low discrimination**					**47573002**
Actinomyces naeslundii		55.56	0.65	ESL 12		
Actinomyces israelii		40.23	0.53			
Bifidobacterium breve		1.67	0.24			
47573003	**Very good identification**					**47573003**
Actinomyces naeslundii		99.52	0.68	ESL 12		
Actinomyces israelii		0.49	0.19			
47573010	**Low discrimination**					**47573010**
					CAT	
Actinomyces israelii		54.55	0.36	SAL 92	1	
Actinomyces naeslundii		39.81	0.43	XYL 92	25	
Propionibacterium granulosum		2.92	0.09	MLZ 13	90	
47573011	**Good identification**					**47573011**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47573012	**Low discrimination**					**47573012**
Actinomyces israelii		50.54	0.39	SAL 92		
Actinomyces naeslundii		24.59	0.43	XYL 92		
Bifidobacterium breve		14.05	0.24	MLZ 13		
47573013	**Good identification**					**47573013**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47573020	**Low discrimination**					**47573020**
					SPO	
Actinomyces naeslundii		89.48	0.65	ESL 12	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		5.73	0.19		92		
Actinomyces israelii		2.76	0.30		1		
47573021	**Very good identification**						**47573021**
Actinomyces naeslundii		100.0	0.68	ESL 12			
47573022	**Low discrimination**						**47573022**
					SPO	GRA	
Actinomyces naeslundii		69.03	0.65	ESL 12	1	99	
Clostridium ramosum		17.69	0.30		92	50	
Prevotella oralis		5.62	0.21		1	1	
47573023	**Very good identification**						**47573023**
Actinomyces naeslundii		100.0	0.68	ESL 12			
47573030	**Low discrimination**						**47573030**
Actinomyces naeslundii		84.18	0.43	ESL 12			
Bifidobacterium breve		8.46	0.12	MLZ 5			
Actinomyces israelii		7.36	0.16				
47573031	**Good identification**						**47573031**
Actinomyces naeslundii		100.0	0.46	ESL 12			
47573033	**Good identification**						**47573033**
Actinomyces naeslundii		100.0	0.46	ESL 12			
47573040	**Low discrimination**						**47573040**
Actinomyces israelii		51.54	0.43	SAL 92			
Actinomyces naeslundii		48.46	0.52	XYL 92			
47573041	**Good identification**						**47573041**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47573042	**Low discrimination**						**47573042**
Actinomyces israelii		57.69	0.46	SAL 92			
Actinomyces naeslundii		36.15	0.52	XYL 92			
Bifidobacterium breve		6.16	0.24				
47573043	**Good identification**						**47573043**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47573060	**Acceptable identification**						**47573060**
Actinomyces naeslundii		91.09	0.52	ESL 12			
Actinomyces israelii		6.18	0.23				
Bifidobacterium breve		2.73	0.12				
47573061	**Good identification**						**47573061**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47573062	**Low discrimination**						**47573062**
Actinomyces naeslundii		71.19	0.52	ESL 12			
Bifidobacterium breve		21.56	0.29				
Actinomyces israelii		7.25	0.26				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47573063	**Good identification**					**47573063**
Actinomyces naeslundii		100.0	0.55	ESL 12		
47573100	**Low discrimination**					**47573100**
Actinomyces naeslundii		53.42	0.73	ESL 12		
Actinomyces israelii		36.24	0.61			
Bifidobacterium breve		5.53	0.41			
47573101	**Very good identification**					**47573101**
Actinomyces naeslundii		99.55	0.76	ESL 12		
Actinomyces israelii		0.46	0.27			
47573102	**Low discrimination**					**47573102**
Bifidobacterium breve		27.51	0.59	SAL 99		
Actinomyces israelii		26.77	0.64			
Actinomyces naeslundii		26.31	0.73			
47573103	**Very good identification**					**47573103**
Actinomyces naeslundii		98.27	0.76	ESL 12		
Bifidobacterium breve		0.69	0.24			
Actinomyces israelii		0.67	0.30			
47573110	**Low discrimination**					**47573110**
Bifidobacterium breve		37.86	0.41	SAL 99		
Actinomyces israelii		37.11	0.47	BGL 91		
Actinomyces naeslundii		19.27	0.51			
47573111	**Good identification**					**47573111**
Actinomyces naeslundii		100.0	0.54	ESL 12		
47573112	**Low discrimination**					**47573112**
Bifidobacterium breve		50.25	0.59	SAL 99		
Bifidobacterium adolescentis		32.37	0.54			
Actinomyces israelii		7.31	0.50			
47573113	**Low discrimination**					**47573113**
Actinomyces naeslundii		79.60	0.54	ESL 12		
Bifidobacterium breve		10.65	0.24	MLZ 5		
Bifidobacterium adolescentis		6.86	0.20			
47573120	**Low discrimination**					**47573120**
					SPO	
Clostridium ramosum		63.01	0.53	SAL 99	92	
Actinomyces naeslundii		28.27	0.73		1	
Bifidobacterium breve		5.20	0.46		1	
47573121	**Very good identification**					**47573121**
					SPO	
Actinomyces naeslundii		98.40	0.76	ESL 12	1	
Clostridium ramosum		1.48	0.19		92	
Bifidobacterium breve		0.12	0.12		1	
47573122	**Low discrimination**					**47573122**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		71.59	0.64	SAL 99	92	
Bifidobacterium breve		14.93	0.63		1	
Actinomyces naeslundii		8.03	0.73		1	
47573123	**Good identification**					**47573123**
					SPO	
Actinomyces naeslundii		92.97	0.76	ESL 12	1	
Clostridium ramosum		5.58	0.30		92	
Bifidobacterium breve		1.16	0.29		1	
47573130	**Low discrimination**					**47573130**
					SPO	
Bifidobacterium breve		63.77	0.46	SAL 99	1	
Actinomyces naeslundii		18.25	0.51	BGL 91	1	
Clostridium tertium		7.93	0.21		99	
47573131	**Good identification**					**47573131**
Actinomyces naeslundii		97.70	0.54	ESL 12		
Bifidobacterium breve		2.30	0.12	MLZ 5		
47573132	**Low discrimination**					**47573132**
					SPO	
Bifidobacterium breve		87.10	0.63	SAL 99	1	
Clostridium ramosum		4.22	0.30		92	
Clostridium tertium		3.21	0.29		99	
47573133	**Low discrimination**					**47573133**
Actinomyces naeslundii		80.78	0.54	ESL 12		
Bifidobacterium breve		19.22	0.29	MLZ 5		
47573140	**Low discrimination**					**47573140**
Actinomyces israelii		48.52	0.54	SAL 92		
Actinomyces naeslundii		32.46	0.60	XYL 92		
Bifidobacterium breve		19.02	0.41			
47573141	**Very good identification**					**47573141**
Actinomyces naeslundii		99.00	0.63	ESL 12		
Actinomyces israelii		1.00	0.20			
47573142	**Low discrimination**					**47573142**
Bifidobacterium breve		61.82	0.59	SAL 99		
Actinomyces israelii		23.39	0.57			
Actinomyces naeslundii		10.43	0.60			
47573143	**Good identification**					**47573143**
Actinomyces naeslundii		94.79	0.63	ESL 12		
Bifidobacterium breve		3.78	0.24			
Actinomyces israelii		1.43	0.23			
47573150	**Low discrimination**					**47573150**
Bifidobacterium breve		67.09	0.41	SAL 99		
Actinomyces israelii		25.57	0.40	BGL 91		
Actinomyces naeslundii		6.03	0.38			
47573151	**Good identification**					**47573151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
47573152	**Low discrimination**					**47573152**
Bifidobacterium breve		87.24	0.59	SAL 99		
Bifidobacterium adolescentis		6.95	0.39			
Actinomyces israelii		4.93	0.43			
47573153	**Low discrimination**					**47573153**
Actinomyces naeslundii		56.87	0.41	ESL 12		
Bifidobacterium breve		43.13	0.24	MLZ 5		
47573160	**Low discrimination**					**47573160**
					SPO	
Bifidobacterium breve		45.66	0.46	SAL 99	1	
Actinomyces naeslundii		43.82	0.60	BGL 91	1	
Clostridium ramosum		5.59	0.19		92	
47573161	**Very good identification**					**47573161**
Actinomyces naeslundii		99.30	0.63	ESL 12		
Bifidobacterium breve		0.70	0.12			
47573162	**Low discrimination**					**47573162**
					SPO	
Bifidobacterium breve		85.88	0.63	SAL 99	1	
Actinomyces naeslundii		8.15	0.60		1	
Clostridium ramosum		4.16	0.30		92	
47573163	**Good identification**					**47573163**
Actinomyces naeslundii		93.38	0.63	ESL 12		
Bifidobacterium breve		6.63	0.29			
47573170	**Acceptable identification**					**47573170**
Bifidobacterium breve		93.97	0.46	SAL 99		
Actinomyces naeslundii		4.75	0.38	BGL 91		
Actinomyces israelii		1.29	0.19			
47573171	**Low discrimination**					**47573171**
Actinomyces naeslundii		88.24	0.41	ESL 12		
Bifidobacterium breve		11.76	0.12	MLZ 5		
47573172	**Very good identification**					**47573172**
Bifidobacterium breve		99.30	0.63	SAL 99		
Actinomyces naeslundii		0.50	0.38			
Actinomyces israelii		0.20	0.22			
47573300	**Low discrimination**					**47573300**
					SPO	
Actinomyces israelii		56.11	0.41	SAL 92	1	
Clostridium ramosum		32.79	0.21	XYL 92	92	
Actinomyces naeslundii		11.10	0.39	RHA 7	1	
47573301	**Good identification**					**47573301**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47573303	**Acceptable identification**					**47573303**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		92.63	0.42	ESL 12	1		
Clostridium ramosum		7.37	-.03	RHA 1	92		
47573320	**Good identification**						**47573320**
					SPO	GRA	
Clostridium ramosum		99.11	0.56	SAL 99	92	50	
Actinomyces naeslundii		0.34	0.39		1	99	
Prevotella oralis		0.34	0.13		1	1	
47573322	**Very good identification**						**47573322**
					SPO		
Clostridium ramosum		97.09	0.66	SAL 99	92		
Prevotella oralis		1.58	0.35		1		
Parabacteroides distasonis		1.03	0.44		1		
47573342	**Low discrimination**						**47573342**
Actinomyces israelii		68.73	0.37	SAL 92			
Bifidobacterium breve		24.38	0.24	XYL 92			
Actinomyces naeslundii		4.11	0.26	RHA 7			
47573400	**Good identification**						**47573400**
					SPO		
Actinomyces israelii		95.30	0.49	SAL 92	1		
Actinomyces naeslundii		2.44	0.31	XYL 92	1		
Clostridium butyricum		2.27	0.00		99		
47573402	**Low discrimination**						**47573402**
Actinomyces israelii		83.18	0.52	SAL 92			
Bifidobacterium adolescentis		14.89	0.33	XYL 92			
Actinomyces naeslundii		1.42	0.31				
47573440	**Good identification**						**47573440**
Actinomyces israelii		100.0	0.42	SAL 92			
47573442	**Acceptable identification**						**47573442**
Actinomyces israelii		94.62	0.45	SAL 92			
Bifidobacterium adolescentis		5.38	0.17	XYL 92			
47573500	**Low discrimination**						**47573500**
					SPO		
Actinomyces israelii		81.43	0.59	SAL 92	1		
Clostridium butyricum		11.62	0.24	XYL 92	99		
Bifidobacterium adolescentis		5.47	0.33		1		
47573501	**Low discrimination**						**47573501**
Actinomyces naeslundii		72.99	0.42	ESL 12			
Actinomyces israelii		27.01	0.25	ARA 1			
47573502	**Low discrimination**						**47573502**
					SPO		
Bifidobacterium adolescentis		80.60	0.67	XYL 87	1		
Actinomyces israelii		18.19	0.62		1		
Clostridium butyricum		0.67	0.17		99		
47573510	**Low discrimination**						**47573510**
Actinomyces israelii		53.60	0.45	SAL 92			
Bifidobacterium adolescentis		29.43	0.34	XYL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		12.97	0.10	MLZ 13			
47573512	**Good identification**						**47573512**
Bifidobacterium adolescentis		92.80	0.69	XYL 87			
Bifidobacterium dentium		4.18	0.27				
Actinomyces israelii		2.56	0.48				
47573520	**Low discrimination**						**47573520**
					SPO		
Actinomyces israelii		45.00	0.39	SAL 92	1		
Clostridium ramosum		28.59	0.19	XYL 92	92		
Actinomyces naeslundii		12.83	0.39	NAG 6	1		
47573521	**Good identification**						**47573521**
Actinomyces naeslundii		100.0	0.42	ESL 12			
47573523	**Good identification**						**47573523**
Actinomyces naeslundii		100.0	0.42	ESL 12			
47573540	**Low discrimination**						**47573540**
					COC	SPO	
Actinomyces israelii		72.97	0.52	SAL 92	1	1	
Blautia producta		22.55	0.24	XYL 92	99	1	
Clostridium butyricum		2.33	0.06		1	99	
47573542	**Low discrimination**						**47573542**
Bifidobacterium adolescentis		56.84	0.51	XYL 87			
Actinomyces israelii		40.37	0.55	SOR 11			
Bifidobacterium breve		1.32	0.24				
47573552	**Low discrimination**						**47573552**
					COC		
Bifidobacterium adolescentis		87.00	0.53	XYL 87	1		
Actinomyces israelii		7.56	0.41	SOR 11	1		
Blautia producta		3.47	0.15		99		
47573700	**Acceptable identification**						**47573700**
					SPO		
Actinomyces israelii		94.46	0.40	SAL 92	1		
Clostridium butyricum		5.54	-.02	XYL 92	99		
47573702	**Low discrimination**						**47573702**
Actinomyces israelii		54.29	0.43	SAL 92			
Bifidobacterium adolescentis		32.28	0.33	XYL 92			
Clostridium clostridioforme		12.39	0.23	RHA 7			
47573742	**Low discrimination**						**47573742**
					COC		
Actinomyces israelii		66.98	0.36	SAL 92	1		
Blautia producta		20.37	0.07	XYL 92	99		
Bifidobacterium adolescentis		12.66	0.17	RHA 7	1		
47574112	**Low discrimination**						**47574112**
Clostridium clostridioforme		75.28	0.42	SAL 88			
Bifidobacterium adolescentis		22.14	0.36	RHA 88			
Actinomyces israelii		1.24	0.21	ARA 94			
47574122	**Low discrimination**						**47574122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Parabacteroides distasonis		56.70	0.53	GAL 99	76	
Bacteroides fragilis		27.32	0.25		99	
Prevotella oralis		8.40	0.26		1	
47574132	**Low discrimination**					**47574132**
Parabacteroides distasonis		51.32	0.48	GAL 99		
Clostridium clostridioforme		48.22	0.42			
Bacteroides fragilis		0.46	-.09			
47574202	**Good identification**					**47574202**
Clostridium clostridioforme		100.0	0.36	SAL 88		
47574212	**Good identification**					**47574212**
Clostridium clostridioforme		100.0	0.48	SAL 88		
47574222	**Low discrimination**					**47574222**
Clostridium clostridioforme		87.63	0.36	SAL 88		
Parabacteroides distasonis		10.92	0.27	ARA 94		
Prevotella oralis		1.46	-.01			
47574232	**Good identification**					**47574232**
Clostridium clostridioforme		98.55	0.48	SAL 88		
Parabacteroides distasonis		1.45	0.22	ARA 94		
47574302	**Good identification**					**47574302**
					GRA	
Clostridium clostridioforme		95.56	0.45	SAL 88	50	
Parabacteroides distasonis		3.56	0.27	ARA 94	1	
Actinomyces israelii		0.49	0.16		99	
47574312	**Very good identification**					**47574312**
Clostridium clostridioforme		99.56	0.57	SAL 88		
Parabacteroides distasonis		0.44	0.22	ARA 94		
47574322	**Low discrimination**					**47574322**
Parabacteroides distasonis		77.03	0.61	GAL 99		
Clostridium clostridioforme		20.91	0.45			
Prevotella oralis		1.63	0.19			
47574332	**Low discrimination**					**47574332**
Clostridium clostridioforme		69.61	0.57	SAL 88		
Parabacteroides distasonis		30.31	0.56	ARA 94		
Bacteroides ovatus		0.08	-.04			
47574402	**Acceptable identification**					**47574402**
					SPO	
Clostridium clostridioforme		94.59	0.42	SAL 88	66	
Actinomyces israelii		2.08	0.23	RHA 88	1	
Clostridium butyricum		1.84	0.01		99	
47574412	**Good identification**					**47574412**
Clostridium clostridioforme		99.58	0.53	SAL 88		
Bifidobacterium adolescentis		0.42	0.16	RHA 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47574422	**Good identification**					**47574422**
Clostridium clostridioforme		98.82	0.42	SAL 88		
Parabacteroides distasonis		1.18	0.15	RHA 88		
47574432	**Good identification**					**47574432**
Clostridium clostridioforme		99.86	0.53	SAL 88		
Parabacteroides distasonis		0.14	0.10	RHA 88		
47574502	**Low discrimination**					**47574502**
					SPO	
Clostridium clostridioforme		61.00	0.51	SAL 88	66	
Bifidobacterium adolescentis		28.56	0.48	RHA 88	1	
Clostridium butyricum		8.49	0.25		99	
47574512	**Low discrimination**					**47574512**
					SPO	
Clostridium clostridioforme		88.49	0.62	SAL 88	66	
Bifidobacterium adolescentis		11.12	0.50	RHA 88	1	
Clostridium butyricum		0.14	0.03		99	
47574522	**Low discrimination**					**47574522**
Clostridium clostridioforme		70.91	0.51	SAL 88		
Parabacteroides distasonis		24.98	0.49	RHA 88		
Bacteroides thetaiotaomicron		1.61	0.13			
47574532	**Very good identification**					**47574532**
					GRA	
Clostridium clostridioforme		95.20	0.62	SAL 88	50	
Parabacteroides distasonis		3.96	0.44	RHA 88	1	
Bifidobacterium longum subsp. longum		0.32	0.21		99	
47574602	**Very good identification**					**47574602**
Clostridium clostridioforme		100.0	0.57	SAL 88		
47574612	**Very good identification**					**47574612**
Clostridium clostridioforme		100.0	0.68	SAL 88		
47574622	**Very good identification**					**47574622**
Clostridium clostridioforme		99.24	0.57	SAL 88		
Parabacteroides distasonis		0.48	0.23			
Bacteroides thetaiotaomicron		0.28	0.03			
47574632	**Very good identification**					**47574632**
Clostridium clostridioforme		99.92	0.68	SAL 88		
Parabacteroides distasonis		0.06	0.18			
Bacteroides thetaiotaomicron		0.02	-.04			
47574702	**Very good identification**					**47574702**
					GRA	
Clostridium clostridioforme		99.66	0.66	SAL 88	50	
Parabacteroides distasonis		0.15	0.23		1	
Bifidobacterium adolescentis		0.06	0.14		99	
47574710	**Good identification**					**47574710**
Clostridium clostridioforme		100.0	0.43	SAL 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47574712	**Very good identification**						**47574712**
					GRA		
Clostridium clostridioforme		99.97	0.77	SAL 88	50		
Parabacteroides distasonis		0.02	0.18		1		
Bifidobacterium adolescentis		0.02	0.16		99		
47574713	**Good identification**						**47574713**
Clostridium clostridioforme		100.0	0.43	SAL 88			
47574722	**Low discrimination**						**47574722**
Clostridium clostridioforme		83.17	0.66	SAL 88			
Parabacteroides distasonis		11.99	0.57				
Bacteroides thetaiotaomicron		3.43	0.32				
47574730	**Good identification**						**47574730**
Clostridium clostridioforme		98.33	0.43	SAL 88			
Parabacteroides distasonis		1.68	0.18	BGL 99			
47574732	**Very good identification**						**47574732**
Clostridium clostridioforme		97.75	0.77	SAL 88			
Parabacteroides distasonis		1.67	0.52				
Bacteroides thetaiotaomicron		0.34	0.25				
47574733	**Good identification**						**47574733**
Clostridium clostridioforme		98.33	0.43	SAL 88			
Parabacteroides distasonis		1.68	0.18	URE 1			
47574752	**Good identification**						**47574752**
Clostridium clostridioforme		100.0	0.43	SAL 88			
47574772	**Good identification**						**47574772**
Clostridium clostridioforme		97.34	0.43	SAL 88			
Parabacteroides distasonis		1.66	0.18	SOR 1			
Bacteroides ovatus		1.00	-.02				
47575000	**Low discrimination**						**47575000**
					COC	SPO	
Actinomyces israelii		61.38	0.56	SAL 92	1	1	
Clostridium butyricum		19.03	0.26		1	99	
Atopobium parvulum		11.83	0.30		99	1	
47575002	**Low discrimination**						**47575002**
					SPO		
Actinomyces israelii		75.39	0.59	SAL 92	1		
Bifidobacterium adolescentis		6.65	0.34		1		
Clostridium butyricum		6.06	0.19		99		
47575010	**Low discrimination**						**47575010**
					SPO		
Actinomyces israelii		85.68	0.42	SAL 92	1		
Clostridium butyricum		9.36	0.04	MLZ 13	99		
Clostridium clostridioforme		2.85	0.07		66		
47575012	**Low discrimination**						**47575012**
Clostridium clostridioforme		55.07	0.41	SAL 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		25.05	0.45	RHA 88			
Bifidobacterium adolescentis		18.07	0.36	ARA 94			
47575020	**Low discrimination**						**47575020**
					COC	GRA	
Actinomyces israelii		27.95	0.35	SAL 92	1	99	
Prevotella oralis		23.44	0.17	NAG 6	1	1	
Atopobium parvulum		14.89	0.17		99	99	
47575022	**Low discrimination**						**47575022**
					CAT		
Parabacteroides distasonis		37.90	0.53	LAC 99	76		
Prevotella oralis		35.46	0.39		1		
Bacteroides fragilis		18.26	0.25		99		
47575032	**Low discrimination**						**47575032**
					GRA		
Parabacteroides distasonis		52.83	0.48	LAC 99	1		
Clostridium clostridioforme		44.48	0.41		50		
Actinomyces israelii		1.29	0.24		99		
47575040	**Acceptable identification**						**47575040**
					SPO		
Actinomyces israelii		93.52	0.49	SAL 92	1		
Clostridium butyricum		6.48	0.08		99		
47575042	**Good identification**						**47575042**
					SPO		
Actinomyces israelii		95.60	0.52	SAL 92	1		
Bifidobacterium adolescentis		2.68	0.19		1		
Clostridium butyricum		1.72	0.01		99		
47575052	**Low discrimination**						**47575052**
Actinomyces israelii		77.74	0.38	SAL 92			
Bifidobacterium adolescentis		17.83	0.20	MLZ 13			
Clostridium clostridioforme		4.44	0.07				
47575100	**Low discrimination**						**47575100**
					COC	SPO	
Atopobium parvulum		61.47	0.64	RAF 1	99	1	
Clostridium butyricum		23.98	0.50		1	99	
Actinomyces israelii		12.89	0.66		1	1	
47575102	**Low discrimination**						**47575102**
					COC		
Bifidobacterium adolescentis		46.06	0.69	ARA 87	1		
Actinomyces israelii		21.10	0.69		1		
Atopobium parvulum		11.84	0.51		99		
47575110	**Low discrimination**						**47575110**
					COC	SPO	
Actinomyces israelii		43.39	0.52	SAL 92	1	1	
Clostridium butyricum		28.43	0.28	MLZ 13	1	99	
Atopobium parvulum		13.99	0.30		99	1	
47575112	**Low discrimination**						**47575112**
Bifidobacterium adolescentis		80.91	0.70	ARA 87			
Clostridium clostridioforme		8.34	0.50				
Actinomyces israelii		4.53	0.55				
47575113	**Low discrimination**						**47575113**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		82.00	0.36	ARA 87			
Clostridium clostridioforme		8.45	0.16	URE 1			
Actinomyces israelii		4.59	0.21				
47575120	**Low discrimination**						**47575120**
					COC	GRA	
Atopobium parvulum		35.87	0.51	RAF 1	99	99	
Bacteroides fragilis		29.08	0.38		1	1	
Parabacteroides distasonis		11.58	0.53		1	1	
47575122	**Low discrimination**						**47575122**
					CAT		
Parabacteroides distasonis		60.16	0.87		76		
Bacteroides fragilis		28.99	0.60		99		
Prevotella oralis		8.91	0.60		1		
47575123	**Low discrimination**						**47575123**
					CAT		
Parabacteroides distasonis		59.64	0.53	URE 1	76		
Bacteroides fragilis		28.74	0.25		99		
Prevotella oralis		8.83	0.26		1		
47575130	**Low discrimination**						**47575130**
					GRA	SPO	
Parabacteroides distasonis		73.49	0.48	BGL 99	1	1	
Actinomyces israelii		4.79	0.31		99	1	
Clostridium butyricum		4.28	0.10		99	99	
47575132	**Very good identification**						**47575132**
Parabacteroides distasonis		95.21	0.82				
Clostridium clostridioforme		2.71	0.50				
Bacteroides fragilis		0.86	0.25				
47575133	**Good identification**						**47575133**
Parabacteroides distasonis		96.38	0.48	URE 1			
Clostridium clostridioforme		2.74	0.16				
Bacteroides fragilis		0.87	-.09				
47575140	**Low discrimination**						**47575140**
					COC	SPO	
Actinomyces israelii		64.50	0.59	SAL 92	1	1	
Clostridium butyricum		26.83	0.31		1	99	
Atopobium parvulum		7.99	0.30		99	1	
47575142	**Low discrimination**						**47575142**
					SPO		
Actinomyces israelii		54.08	0.62	SAL 92	1		
Bifidobacterium adolescentis		37.52	0.53		1		
Clostridium butyricum		5.83	0.24		99		
47575150	**Low discrimination**						**47575150**
					SPO		
Actinomyces israelii		79.51	0.45	SAL 92	1		
Clostridium butyricum		11.65	0.10	MLZ 13	99		
Bifidobacterium adolescentis		6.84	0.20		1		
47575152	**Low discrimination**						**47575152**
Bifidobacterium adolescentis		82.52	0.54	ARA 87			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		14.54	0.48	SOR 11			
Bifidobacterium breve		1.28	0.24				
47575160	**Low discrimination**						**47575160**
					COC	SPO	
Actinomyces israelii		36.96	0.39	SAL 92	1	1	
Clostridium butyricum		20.95	0.13	NAG 6	1	99	
Atopobium parvulum		12.66	0.17		99	1	
47575162	**Low discrimination**						**47575162**
Parabacteroides distasonis		49.66	0.53	SOR 1			
Bacteroides fragilis		23.93	0.25				
Capnocytophaga ochracea		7.38	0.26				
47575172	**Low discrimination**						**47575172**
					GRA		
Parabacteroides distasonis		81.06	0.48	SOR 1	1		
Bifidobacterium breve		7.56	0.29		99		
Actinomyces israelii		3.08	0.27		99		
47575200	**Low discrimination**						**47575200**
					SPO		
Actinomyces israelii		81.05	0.36	SAL 92	1		
Clostridium butyricum		10.32	0.00	RHA 7	99		
Clostridium clostridioforme		8.63	0.10		66		
47575202	**Low discrimination**						**47575202**
					GRA		
Clostridium clostridioforme		82.14	0.45	SAL 88	50		
Actinomyces israelii		11.69	0.39	ARA 94	99		
Parabacteroides distasonis		3.41	0.27		1		
47575212	**Good identification**						**47575212**
					GRA		
Clostridium clostridioforme		99.05	0.56	SAL 88	50		
Parabacteroides distasonis		0.49	0.22	ARA 94	1		
Actinomyces israelii		0.46	0.25		99		
47575222	**Low discrimination**						**47575222**
Parabacteroides distasonis		72.12	0.61	LAC 99			
Clostridium clostridioforme		17.54	0.45				
Prevotella oralis		9.64	0.33				
47575232	**Low discrimination**						**47575232**
Clostridium clostridioforme		67.13	0.56	SAL 88			
Parabacteroides distasonis		32.62	0.56	ARA 94			
Bacteroides ovatus		0.09	-.04				
47575300	**Low discrimination**						**47575300**
					COC	SPO	
Actinomyces israelii		36.92	0.47	SAL 92	1	1	
Clostridium butyricum		28.22	0.24	RHA 7	1	99	
Atopobium parvulum		23.63	0.30		99	1	
47575302	**Low discrimination**						**47575302**
					GRA		
Parabacteroides distasonis		46.57	0.61	NAG 99	1		
Clostridium clostridioforme		37.93	0.54		50		
Actinomyces israelii		6.45	0.50		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47575312	**Low discrimination**					**47575312**
					GRA	
Clostridium clostridioforme		85.80	0.65	SAL 88	50	
Parabacteroides distasonis		12.45	0.56	ARA 94	1	
Bifidobacterium adolescentis		1.15	0.36		99	
47575320	**Low discrimination**					**47575320**
Parabacteroides distasonis		64.62	0.61	BGL 99		
Bacteroides vulgatus		25.17	0.35			
Prevotella oralis		7.12	0.32			
47575322	**Very good identification**					**47575322**
Parabacteroides distasonis		96.58	0.95			
Prevotella oralis		2.04	0.53			
Clostridium clostridioforme		0.79	0.54			
47575323	**Very good identification**					**47575323**
Parabacteroides distasonis		96.62	0.61	URE 1		
Prevotella oralis		2.04	0.19			
Clostridium clostridioforme		0.80	0.19			
47575330	**Acceptable identification**					**47575330**
Parabacteroides distasonis		92.44	0.56	BGL 99		
Clostridium clostridioforme		6.43	0.31			
Bacteroides vulgatus		0.68	0.01			
47575332	**Good identification**					**47575332**
Parabacteroides distasonis		93.10	0.90			
Clostridium clostridioforme		6.48	0.65			
Bacteroides ovatus		0.26	0.31			
47575333	**Acceptable identification**					**47575333**
Parabacteroides distasonis		93.25	0.56	URE 1		
Clostridium clostridioforme		6.49	0.31			
Bacteroides ovatus		0.26	-.04			
47575340	**Low discrimination**					**47575340**
					SPO	
Actinomyces israelii		85.40	0.40	SAL 92	1	
Clostridium butyricum		14.60	0.06	RHA 7	99	
47575342	**Low discrimination**					**47575342**
					GRA	
Actinomyces israelii		66.74	0.43	SAL 92	99	
Parabacteroides distasonis		12.52	0.27	RHA 7	1	
Clostridium clostridioforme		10.19	0.19		50	
47575362	**Very good identification**					**47575362**
Parabacteroides distasonis		95.65	0.61	SOR 1		
Prevotella oralis		2.02	0.19			
Bacteroides ovatus		0.89	0.10			
47575372	**Acceptable identification**					**47575372**
Parabacteroides distasonis		92.50	0.56	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		6.44	0.31				
Bacteroides ovatus		1.06	0.07				
47575400	**Low discrimination**						**47575400**
					SPO		
Clostridium butyricum		76.68	0.42	SAL 99	99		
Actinomyces israelii		22.49	0.54	LAC 96	1		
Clostridium clostridioforme		0.47	0.16		66		
47575402	**Low discrimination**						**47575402**
					SPO		
Clostridium clostridioforme		34.56	0.50	SAL 88	66		
Actinomyces israelii		25.03	0.57	RHA 88	1		
Clostridium butyricum		22.13	0.35		99		
47575412	**Low discrimination**						**47575412**
Clostridium clostridioforme		85.22	0.61	SAL 88			
Bifidobacterium adolescentis		11.95	0.50	RHA 88			
Actinomyces israelii		2.02	0.43				
47575422	**Low discrimination**						**47575422**
					GRA	SPO	
Clostridium clostridioforme		63.37	0.50	SAL 88	50	66	
Parabacteroides distasonis		24.91	0.49	RHA 88	1	1	
Clostridium butyricum		3.53	0.17		99	99	
47575432	**Good identification**						**47575432**
Clostridium clostridioforme		94.29	0.61	SAL 88			
Parabacteroides distasonis		4.38	0.44	RHA 88			
Bacteroides thetaiotaomicron		0.41	0.11				
47575440	**Low discrimination**						**47575440**
					SPO		
Actinomyces israelii		56.74	0.47	SAL 92	1		
Clostridium butyricum		43.26	0.24		99		
47575442	**Low discrimination**						**47575442**
					SPO		
Actinomyces israelii		68.36	0.50	SAL 92	1		
Bifidobacterium adolescentis		15.67	0.33		1		
Clostridium butyricum		13.52	0.17		99		
47575500	**Good identification**						**47575500**
					SPO		
Clostridium butyricum		93.18	0.66	SAL 99	99		
Actinomyces israelii		4.55	0.64		1		
Bifidobacterium adolescentis		1.23	0.48		1		
47575502	**Low discrimination**						**47575502**
					SPO		
Bifidobacterium adolescentis		69.84	0.83		1		
Clostridium butyricum		20.75	0.59		99		
Clostridium clostridioforme		4.52	0.59		66		
47575503	**Low discrimination**						**47575503**
					SPO		
Bifidobacterium adolescentis		69.92	0.48	URE 1	1		
Clostridium butyricum		20.77	0.25		99		
Clostridium clostridioforme		4.53	0.25		66		
47575510	**Low discrimination**						**47575510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium butyricum		57.25	0.44	SAL 99	99		
Bifidobacterium adolescentis		17.58	0.50	MLZ 5	1		
Bifidobacterium longum subsp. longum		11.61	0.45		1		
47575512	**Low discrimination**						**47575512**
					SPO		
Bifidobacterium adolescentis		78.33	0.84		1		
Clostridium clostridioforme		18.90	0.70		66		
Clostridium butyricum		1.00	0.37		99		
47575513	**Low discrimination**						**47575513**
					SPO		
Bifidobacterium adolescentis		78.35	0.50	URE 1	1		
Clostridium clostridioforme		18.90	0.36		66		
Clostridium butyricum		1.00	0.03		99		
47575520	**Low discrimination**						**47575520**
					GRA	SPO	
Clostridium butyricum		55.54	0.48	SAL 99	99	99	
Bacteroides vulgatus		17.88	0.38	NAG 8	1	1	
Bifidobacterium longum subsp. longum		16.02	0.52		99	1	
47575522	**Low discrimination**						**47575522**
Parabacteroides distasonis		79.13	0.83				
Clostridium clostridioforme		6.81	0.59				
Bacteroides thetaiotaomicron		5.11	0.47				
47575523	**Low discrimination**						**47575523**
Parabacteroides distasonis		79.97	0.49	URE 1			
Clostridium clostridioforme		6.88	0.25				
Bacteroides thetaiotaomicron		5.17	0.13				
47575530	**Low discrimination**						**47575530**
					GRA	SPO	
Bifidobacterium longum subsp. longum		71.85	0.56	CEL 1	99	1	
Parabacteroides distasonis		9.01	0.44		1	1	
Clostridium butyricum		7.70	0.26		99	99	
47575532	**Low discrimination**						**47575532**
					GRA		
Parabacteroides distasonis		51.45	0.78		1		
Clostridium clostridioforme		37.44	0.70		50		
Bifidobacterium longum subsp. longum		4.14	0.56		99		
47575533	**Low discrimination**						**47575533**
					GRA		
Parabacteroides distasonis		51.60	0.44	URE 1	1		
Clostridium clostridioforme		37.55	0.36		50		
Bifidobacterium longum subsp. longum		4.16	0.21		99		
47575540	**Low discrimination**						**47575540**
					COC	SPO	
Clostridium butyricum		78.13	0.48	SAL 99	1	99	
Actinomyces israelii		17.08	0.57	SOR 8	1	1	
Blautia producta		2.60	0.24		99	1	
47575542	**Low discrimination**						**47575542**
					SPO		
Bifidobacterium adolescentis		71.73	0.67	SOR 11	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		14.99	0.41		99		
Actinomyces israelii		12.64	0.60		1		
47575552	**Good identification**						**47575552**
Bifidobacterium adolescentis		94.66	0.69	SOR 11			
Actinomyces israelii		2.04	0.46				
Clostridium clostridioforme		1.87	0.36				
47575562	**Low discrimination**						**47575562**
					GRA	SPO	
Parabacteroides distasonis		51.37	0.49	SOR 1	1	1	
Clostridium butyricum		15.18	0.23		99	99	
Actinomyces israelii		9.40	0.40		99	1	
47575572	**Low discrimination**						**47575572**
					GRA		
Parabacteroides distasonis		40.78	0.44	SOR 1	1		
Clostridium clostridioforme		29.68	0.36		50		
Bifidobacterium adolescentis		15.20	0.34		99		
47575602	**Very good identification**						**47575602**
					SPO		
Clostridium clostridioforme		98.68	0.65	SAL 88	66		
Actinomyces israelii		0.73	0.38		1		
Clostridium butyricum		0.27	0.09		99		
47575610	**Good identification**						**47575610**
					SPO		
Clostridium clostridioforme		97.65	0.42	SAL 88	66		
Actinomyces israelii		1.57	0.21	BGL 99	1		
Clostridium butyricum		0.78	-.06		99		
47575612	**Very good identification**						**47575612**
					GRA		
Clostridium clostridioforme		99.93	0.76	SAL 88	50		
Actinomyces israelii		0.02	0.24		99		
Parabacteroides distasonis		0.02	0.18		1		
47575613	**Good identification**						**47575613**
Clostridium clostridioforme		100.0	0.42	SAL 88			
47575622	**Low discrimination**						**47575622**
Clostridium clostridioforme		78.52	0.65	SAL 88			
Parabacteroides distasonis		12.63	0.57				
Bacteroides thetaiotaomicron		7.30	0.37				
47575630	**Good identification**						**47575630**
Clostridium clostridioforme		97.38	0.42	SAL 88			
Parabacteroides distasonis		1.85	0.18	BGL 99			
Bacteroides thetaiotaomicron		0.77	-.04				
47575632	**Very good identification**						**47575632**
Clostridium clostridioforme		97.11	0.76	SAL 88			
Parabacteroides distasonis		1.85	0.52				
Bacteroides thetaiotaomicron		0.77	0.30				
47575633	**Good identification**						**47575633**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		97.38	0.42	SAL 88		
Parabacteroides distasonis		1.85	0.18	URE 1		
Bacteroides thetaiotaomicron		0.77	-.04			
47575652	**Good identification**					**47575652**
Clostridium clostridioforme		99.07	0.42	SAL 88		
Actinomyces israelii		0.93	0.17	SOR 1		
47575672	**Good identification**					**47575672**
Clostridium clostridioforme		96.30	0.42	SAL 88		
Parabacteroides distasonis		1.83	0.18	SOR 1		
Bacteroides ovatus		1.11	-.02			
47575700	**Low discrimination**					**47575700**
					SPO	
Clostridium butyricum		68.11	0.40	SAL 99	99	
Clostridium clostridioforme		13.82	0.40	RHA 3	66	
Actinomyces israelii		8.10	0.45		1	
47575702	**Low discrimination**					**47575702**
					GRA	
Clostridium clostridioforme		89.49	0.74	SAL 88	50	
Parabacteroides distasonis		4.30	0.57		1	
Bifidobacterium adolescentis		1.90	0.48		99	
47575703	**Low discrimination**					**47575703**
					GRA	
Clostridium clostridioforme		89.99	0.40	SAL 88	50	
Parabacteroides distasonis		4.32	0.23	URE 1	1	
Bifidobacterium adolescentis		1.92	0.14		99	
47575710	**Acceptable identification**					**47575710**
					SPO	
Clostridium clostridioforme		90.04	0.51	SAL 88	66	
Clostridium butyricum		5.13	0.18	BGL 99	99	
Actinomyces israelii		1.73	0.31		1	
47575712	**Very good identification**					**47575712**
					GRA	
Clostridium clostridioforme		98.62	0.85	SAL 88	50	
Bifidobacterium adolescentis		0.56	0.50		99	
Parabacteroides distasonis		0.56	0.52		1	
47575713	**Good identification**					**47575713**
					GRA	
Clostridium clostridioforme		98.87	0.51	SAL 88	50	
Bifidobacterium adolescentis		0.57	0.16	URE 1	99	
Parabacteroides distasonis		0.56	0.18		1	
47575720	**Low discrimination**					**47575720**
					CAT	
Bacteroides vulgatus		82.96	0.55	TRE 4	3	
Parabacteroides distasonis		9.83	0.57	CEL 5	76	
Bacteroides thetaiotaomicron		2.81	0.32		89	
47575722	**Low discrimination**					**47575722**
Parabacteroides distasonis		61.84	0.91			
Bacteroides thetaiotaomicron		17.69	0.66			
Clostridium clostridioforme		13.00	0.74			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47575723	**Low discrimination**					**47575723**
Parabacteroides distasonis		61.94	0.57	URE 1		
Bacteroides thetaiotaomicron		17.71	0.32			
Clostridium clostridioforme		13.02	0.40			
47575730	**Low discrimination**					**47575730**
Clostridium clostridioforme		53.37	0.51	SAL 88		
Parabacteroides distasonis		30.00	0.52	BGL 99		
Bacteroides thetaiotaomicron		6.20	0.25			
47575732	**Low discrimination**					**47575732**
Clostridium clostridioforme		56.79	0.85	SAL 88		
Parabacteroides distasonis		31.92	0.87			
Bacteroides thetaiotaomicron		6.59	0.59			
47575733	**Low discrimination**					**47575733**
Clostridium clostridioforme		56.80	0.51	SAL 88		
Parabacteroides distasonis		31.93	0.52	URE 1		
Bacteroides thetaiotaomicron		6.60	0.25			
47575742	**Low discrimination**					**47575742**
Clostridium clostridioforme		51.81	0.40	SAL 88		
Actinomyces israelii		17.72	0.41	SOR 1		
Bifidobacterium adolescentis		13.49	0.33			
47575752	**Low discrimination**					**47575752**
					COC	
Clostridium clostridioforme		88.68	0.51	SAL 88	1	
Bifidobacterium adolescentis		6.20	0.34	SOR 1	1	
Blautia producta		2.99	0.15		99	
47575762	**Low discrimination**					**47575762**
Parabacteroides distasonis		50.20	0.57	SOR 1		
Bacteroides ovatus		24.50	0.36			
Bacteroides thetaiotaomicron		14.36	0.32			
47575772	**Low discrimination**					**47575772**
Clostridium clostridioforme		49.47	0.51	SAL 88		
Parabacteroides distasonis		27.81	0.52	SOR 1		
Bacteroides ovatus		16.80	0.33			
47576002	**Good identification**					**47576002**
Actinomyces israelii		100.0	0.37	SAL 92		
47576100	**Good identification**					**47576100**
					SPO	
Actinomyces israelii		96.48	0.45	SAL 92	1	
Clostridium butyricum		3.52	-.01	GAL 99	99	
47576102	**Low discrimination**					**47576102**
Actinomyces israelii		76.04	0.48	SAL 92		
Bifidobacterium adolescentis		23.97	0.33	GAL 99		
47576140	**Good identification**					**47576140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.38	SAL 92		
47576142	**Acceptable identification**					**47576142**
Actinomyces israelii		90.90	0.41	SAL 92		
Bifidobacterium adolescentis		9.11	0.17	GAL 99		
47576402	**Low discrimination**					**47576402**
Actinomyces israelii		84.36	0.36	SAL 92		
Bifidobacterium adolescentis		8.79	0.13	GAL 99		
Clostridium clostridioforme		6.85	0.08			
47576500	**Low discrimination**					**47576500**
					SPO	
Actinomyces israelii		69.43	0.43	SAL 92	1	
Clostridium butyricum		27.86	0.15	GAL 99	99	
Bifidobacterium adolescentis		2.71	0.13		1	
47576502	**Low discrimination**					**47576502**
					SPO	
Bifidobacterium adolescentis		68.36	0.47	GAL 99	1	
Actinomyces israelii		26.52	0.46		1	
Clostridium butyricum		2.76	0.08		99	
47576512	**Low discrimination**					**47576512**
Bifidobacterium adolescentis		55.20	0.48	GAL 99		
Bifidobacterium dentium		36.76	0.27			
Clostridium clostridioforme		5.43	0.28			
47576542	**Low discrimination**					**47576542**
					COC	
Actinomyces israelii		49.59	0.39	SAL 92	1	
Bifidobacterium adolescentis		40.64	0.31	GAL 99	1	
Blautia producta		9.77	0.07		99	
47576712	**Good identification**					**47576712**
Clostridium clostridioforme		97.72	0.43	SAL 88		
Bifidobacterium adolescentis		1.37	0.14	MAN 1		
Bifidobacterium dentium		0.91	-.07			
47576732	**Low discrimination**					**47576732**
Clostridium clostridioforme		89.61	0.43	SAL 88		
Bacteroides ovatus		7.38	0.14	MAN 1		
Parabacteroides distasonis		1.53	0.18			
47577000	**Very good identification**					**47577000**
					SPO	
Actinomyces israelii		99.22	0.69	SAL 92	1	
Clostridium butyricum		0.60	0.10		99	
Actinomyces naeslundii		0.18	0.31		1	
47577002	**Very good identification**					**47577002**
					SPO	
Actinomyces israelii		98.47	0.72	SAL 92	1	
Bifidobacterium adolescentis		1.25	0.33		1	
Clostridium butyricum		0.16	0.03		99	
47577003	**Low discrimination**					**47577003**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		84.73	0.37	SAL 92			
Actinomyces naeslundii		15.27	0.34	URE 1			
47577010	**Good identification**						**47577010**
Actinomyces israelii		100.0	0.54	SAL 92			
47577012	**Low discrimination**						**47577012**
Actinomyces israelii		89.87	0.57	SAL 92			
Bifidobacterium adolescentis		9.36	0.34	MLZ 13			
Bifidobacterium dentium		0.42	-.07				
47577020	**Good identification**						**47577020**
Actinomyces israelii		97.23	0.48	SAL 92			
Actinomyces naeslundii		2.77	0.31	NAG 6			
47577022	**Acceptable identification**						**47577022**
					GRA		
Actinomyces israelii		93.64	0.51	SAL 92	99		
Parabacteroides distasonis		1.90	0.18	NAG 6	1		
Actinomyces naeslundii		1.78	0.31		99		
47577032	**Low discrimination**						**47577032**
					GRA		
Actinomyces israelii		79.06	0.37	SAL 92	99		
Bifidobacterium dentium		10.33	-.03	MLZ 13	99		
Parabacteroides distasonis		5.77	0.14	NAG 6	1		
47577040	**Very good identification**						**47577040**
Actinomyces israelii		100.0	0.62	SAL 92			
47577042	**Very good identification**						**47577042**
Actinomyces israelii		99.60	0.65	SAL 92			
Bifidobacterium adolescentis		0.40	0.17				
47577050	**Good identification**						**47577050**
					COC		
Actinomyces israelii		98.25	0.47	SAL 92	1		
Blautia producta		1.75	-.03	MLZ 13	99		
47577052	**Good identification**						**47577052**
Actinomyces israelii		96.80	0.50	SAL 92			
Bifidobacterium adolescentis		3.20	0.19	MLZ 13			
47577060	**Good identification**						**47577060**
Actinomyces israelii		100.0	0.41	SAL 92			
47577062	**Good identification**						**47577062**
Actinomyces israelii		100.0	0.44	SAL 92			
47577100	**Very good identification**						**47577100**
					COC	SPO	
Actinomyces israelii		95.08	0.79	SAL 92	1	1	
Clostridium butyricum		3.47	0.33		1	99	
Atopobium parvulum		0.81	0.30		99	1	
47577101	**Low discrimination**						**47577101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		81.38	0.45	SAL 92	1		
Actinomyces naeslundii		15.65	0.42	URE 1	1		
Clostridium butyricum		2.97	-.01		99		
47577102	**Low discrimination**						**47577102**
					SPO		
Actinomyces israelii		75.19	0.82	SAL 92	1		
Bifidobacterium adolescentis		23.70	0.67		1		
Clostridium butyricum		0.71	0.26		99		
47577103	**Low discrimination**						**47577103**
Actinomyces israelii		69.28	0.48	SAL 92			
Bifidobacterium adolescentis		21.84	0.33	URE 1			
Actinomyces naeslundii		8.88	0.42				
47577110	**Good identification**						**47577110**
Actinomyces israelii		92.60	0.65	SAL 92			
Bifidobacterium adolescentis		3.62	0.34	MLZ 13			
Bifidobacterium dentium		1.59	0.10				
47577112	**Low discrimination**						**47577112**
Bifidobacterium adolescentis		69.18	0.69	ARA 87			
Actinomyces israelii		26.83	0.68				
Bifidobacterium dentium		3.11	0.27				
47577120	**Low discrimination**						**47577120**
					SPO		
Actinomyces israelii		82.61	0.58	SAL 92	1		
Clostridium butyricum		4.10	0.15	NAG 6	99		
Clostridium ramosum		3.74	0.19		92		
47577121	**Low discrimination**						**47577121**
Actinomyces naeslundii		75.08	0.42	ESL 12			
Actinomyces israelii		24.92	0.24	XYL 1			
47577122	**Low discrimination**						**47577122**
					GRA	CAT	
Actinomyces israelii		46.91	0.61	SAL 92	99	1	
Parabacteroides distasonis		23.50	0.53	NAG 6	1	76	
Bacteroides fragilis		11.32	0.25		1	99	
47577123	**Low discrimination**						**47577123**
					GRA		
Actinomyces naeslundii		53.55	0.42	ESL 12	99		
Actinomyces israelii		26.66	0.27	XYL 1	99		
Parabacteroides distasonis		13.36	0.18		1		
47577130	**Low discrimination**						**47577130**
					SPO		
Actinomyces israelii		50.67	0.44	SAL 92	1		
Bifidobacterium dentium		24.30	0.15	MLZ 13	1		
Clostridium tertium		21.50	0.24	NAG 6	99		
47577140	**Very good identification**						**47577140**
					COC	SPO	
Actinomyces israelii		96.88	0.72	SAL 92	1	1	
Blautia producta		2.13	0.24		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.79	0.15		1	99	
47577141	**Acceptable identification**						**47577141**
Actinomyces israelii		91.98	0.38	SAL 92			
Actinomyces naeslundii		8.03	0.29	URE 1			
47577142	**Good identification**						**47577142**
Actinomyces israelii		90.37	0.75	SAL 92			
Bifidobacterium adolescentis		9.05	0.51				
Bifidobacterium breve		0.21	0.24				
47577143	**Low discrimination**						**47577143**
Actinomyces israelii		86.33	0.41	SAL 92			
Bifidobacterium adolescentis		8.65	0.17	URE 1			
Actinomyces naeslundii		5.02	0.29				
47577150	**Low discrimination**						**47577150**
					COC		
Actinomyces israelii		68.65	0.58	SAL 92	1		
Blautia producta		30.30	0.32	MLZ 13	99		
Bifidobacterium adolescentis		0.85	0.19		1		
47577152	**Low discrimination**						**47577152**
					COC		
Actinomyces israelii		53.46	0.61	SAL 92	1		
Bifidobacterium adolescentis		43.80	0.53	MLZ 13	1		
Blautia producta		1.75	0.15		99		
47577160	**Acceptable identification**						**47577160**
					COC	SPO	
Actinomyces israelii		93.62	0.51	SAL 92	1	1	
Blautia producta		3.58	0.07	NAG 6	99	1	
Clostridium butyricum		1.04	-.03		1	99	
47577162	**Acceptable identification**						**47577162**
Actinomyces israelii		90.42	0.54	SAL 92			
Bifidobacterium breve		5.85	0.29	NAG 6			
Bifidobacterium adolescentis		1.43	0.17				
47577170	**Low discrimination**						**47577170**
					COC		
Actinomyces israelii		54.57	0.37	SAL 92	1		
Blautia producta		41.93	0.15	MLZ 13	99		
Bifidobacterium breve		3.50	0.12	NAG 6	1		
47577172	**Low discrimination**						**47577172**
Actinomyces israelii		57.14	0.40	SAL 92			
Bifidobacterium breve		24.72	0.29	MLZ 13			
Bifidobacterium adolescentis		7.41	0.19	NAG 6			
47577200	**Good identification**						**47577200**
Actinomyces israelii		100.0	0.49	SAL 92			
47577202	**Good identification**						**47577202**
Actinomyces israelii		98.76	0.52	SAL 92			
Clostridium clostridioforme		1.24	0.10	RHA 7			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47577210	**Good identification**						**47577210**
Actinomyces israelii		100.0	0.35	SAL 92			
47577212	**Low discrimination**						**47577212**
Actinomyces israelii		72.37	0.38	SAL 92			
Clostridium clostridioforme		27.63	0.22	RHA 7			
47577240	**Good identification**						**47577240**
Actinomyces israelii		100.0	0.42	SAL 92			
47577242	**Good identification**						**47577242**
Actinomyces israelii		100.0	0.45	SAL 92			
47577300	**Very good identification**						**47577300**
					SPO		
Actinomyces israelii		98.52	0.60	SAL 92	1		
Clostridium butyricum		1.48	0.07	RHA 7	99		
47577302	**Good identification**						**47577302**
					GRA		
Actinomyces israelii		93.26	0.63	SAL 92	99		
Bifidobacterium adolescentis		3.95	0.33	RHA 7	99		
Parabacteroides distasonis		1.20	0.27		1		
47577310	**Good identification**						**47577310**
					COC		
Actinomyces israelii		97.75	0.46	SAL 92	1		
Blautia producta		2.25	-.03	RHA 7	99		
47577312	**Low discrimination**						**47577312**
Actinomyces israelii		57.89	0.49	SAL 92			
Bifidobacterium adolescentis		20.03	0.34	RHA 7			
Clostridium clostridioforme		18.50	0.31	MLZ 13			
47577320	**Low discrimination**						**47577320**
					SPO	GRA	
Actinomyces israelii		44.45	0.39	SAL 92	1	99	
Clostridium ramosum		35.39	0.21	RHA 7	92	50	
Parabacteroides distasonis		13.44	0.27	NAG 6	1	1	
47577322	**Low discrimination**						**47577322**
					SPO		
Parabacteroides distasonis		79.16	0.61	MAN 1	1		
Clostridium ramosum		8.42	0.32		92		
Bacteroides ovatus		5.86	0.25		1		
47577332	**Low discrimination**						**47577332**
Parabacteroides distasonis		84.52	0.56	MAN 1			
Bacteroides ovatus		7.74	0.22				
Clostridium clostridioforme		5.88	0.31				
47577340	**Low discrimination**						**47577340**
					COC		
Actinomyces israelii		77.40	0.53	SAL 92	1		
Blautia producta		22.60	0.24	RHA 7	99		
47577342	**Good identification**						**47577342**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		96.61	0.56	SAL 92	1		
Blautia producta		2.09	0.07	RHA 7	99		
Bifidobacterium adolescentis		1.30	0.17		1		
47577352	**Low discrimination**						**47577352**
					COC		
Actinomyces israelii		64.76	0.42	SAL 92	1		
Blautia producta		28.13	0.15	RHA 7	99		
Bifidobacterium adolescentis		7.12	0.19	MLZ 13	1		
47577362	**Low discrimination**						**47577362**
					GRA		
Actinomyces israelii		57.74	0.35	SAL 92	99		
Parabacteroides distasonis		29.94	0.27	RHA 7	1		
Bacteroides ovatus		9.14	0.02	NAG 6	1		
47577400	**Good identification**						**47577400**
					SPO		
Actinomyces israelii		93.59	0.67	SAL 92	1		
Clostridium butyricum		6.26	0.26		99		
Bifidobacterium adolescentis		0.15	0.13		1		
47577402	**Low discrimination**						**47577402**
					SPO		
Actinomyces israelii		88.90	0.70	SAL 92	1		
Bifidobacterium adolescentis		9.26	0.47		1		
Clostridium butyricum		1.54	0.19		99		
47577403	**Acceptable identification**						**47577403**
Actinomyces israelii		90.57	0.36	SAL 92			
Bifidobacterium adolescentis		9.43	0.13	URE 1			
47577410	**Low discrimination**						**47577410**
					SPO		
Actinomyces israelii		88.57	0.53	SAL 92	1		
Bifidobacterium dentium		7.46	0.10	MLZ 13	1		
Clostridium butyricum		2.09	0.04		99		
47577412	**Low discrimination**						**47577412**
Actinomyces israelii		39.99	0.56	SAL 92			
Bifidobacterium adolescentis		34.07	0.48	MLZ 13			
Bifidobacterium dentium		22.69	0.27				
47577420	**Acceptable identification**						**47577420**
					SPO		
Actinomyces israelii		91.65	0.47	SAL 92	1		
Clostridium butyricum		8.35	0.08	NAG 6	99		
47577422	**Low discrimination**						**47577422**
Actinomyces israelii		88.73	0.50	SAL 92			
Clostridium clostridioforme		3.42	0.16	NAG 6			
Bifidobacterium dentium		2.12	-.03				
47577440	**Low discrimination**						**47577440**
					COC	SPO	
Actinomyces israelii		89.09	0.60	SAL 92	1	1	
Blautia producta		9.58	0.24		99	1	
Clostridium butyricum		1.33	0.08		1	99	
47577442	**Very good identification**						**47577442**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces israelii		95.70	0.63	SAL 92	1		
Bifidobacterium adolescentis		3.17	0.31		1		
Blautia producta		0.76	0.07		99		
47577452	**Low discrimination**						**47577452**
					COC		
Actinomyces israelii		69.19	0.49	SAL 92	1		
Bifidobacterium adolescentis		18.74	0.33	MLZ 13	1		
Blautia producta		11.06	0.15		99		
47577460	**Low discrimination**						**47577460**
					COC		
Actinomyces israelii		84.24	0.40	SAL 92	1		
Blautia producta		15.76	0.07	NAG 6	99		
47577462	**Good identification**						**47577462**
Actinomyces israelii		100.0	0.43	SAL 92			
47577500	**Low discrimination**						**47577500**
					SPO		
Actinomyces israelii		68.47	0.77	SAL 92	1		
Clostridium butyricum		27.47	0.50		99		
Bifidobacterium adolescentis		2.68	0.47		1		
47577501	**Low discrimination**						**47577501**
					GRA	SPO	
Actinomyces israelii		66.23	0.43	SAL 92	99	1	
Clostridium butyricum		26.57	0.15	URE 1	99	99	
Mitsuokella multacida		4.61	0.11		1	1	
47577502	**Low discrimination**						**47577502**
					SPO		
Bifidobacterium adolescentis		69.54	0.81		1		
Actinomyces israelii		26.98	0.80		1		
Clostridium butyricum		2.81	0.43		99		
47577503	**Low discrimination**						**47577503**
					SPO		
Bifidobacterium adolescentis		69.62	0.47	URE 1	1		
Actinomyces israelii		27.00	0.46		1		
Clostridium butyricum		2.81	0.08		99		
47577510	**Low discrimination**						**47577510**
Bifidobacterium dentium		55.25	0.44	SAL 99			
Actinomyces israelii		26.52	0.63	BGL 91			
Bifidobacterium adolescentis		8.47	0.48				
47577512	**Low discrimination**						**47577512**
Bifidobacterium adolescentis		58.17	0.83				
Bifidobacterium dentium		38.74	0.62				
Actinomyces israelii		2.76	0.66				
47577513	**Low discrimination**						**47577513**
Bifidobacterium adolescentis		58.36	0.48	URE 1			
Bifidobacterium dentium		38.87	0.27				
Actinomyces israelii		2.77	0.32				
47577520	**Low discrimination**						**47577520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		58.04	0.57	SAL 92	1		
Clostridium butyricum		31.73	0.31	NAG 6	99		
Bifidobacterium dentium		5.08	0.15		1		
47577522	**Low discrimination**						**47577522**
Actinomyces israelii		35.52	0.60	SAL 92			
Bifidobacterium dentium		20.97	0.32	NAG 6			
Bifidobacterium adolescentis		14.49	0.47				
47577530	**Good identification**						**47577530**
					COC		
Bifidobacterium dentium		96.92	0.49	SAL 99	1		
Actinomyces israelii		1.67	0.43	BGL 91	1		
Blautia producta		0.61	0.15		99		
47577532	**Very good identification**						**47577532**
					GRA		
Bifidobacterium dentium		98.01	0.66	SAL 99	99		
Bifidobacterium adolescentis		0.84	0.48		99		
Parabacteroides distasonis		0.34	0.44		1		
47577540	**Low discrimination**						**47577540**
					COC	SPO	
Blautia producta		70.40	0.58	SAL 99	99	1	
Actinomyces israelii		26.47	0.70		1	1	
Clostridium butyricum		2.37	0.31		1	99	
47577542	**Low discrimination**						**47577542**
					COC		
Actinomyces israelii		48.99	0.73	SAL 92	1		
Bifidobacterium adolescentis		40.14	0.66		1		
Blautia producta		9.65	0.41		99		
47577543	**Low discrimination**						**47577543**
					COC		
Actinomyces israelii		49.59	0.39	SAL 92	1		
Bifidobacterium adolescentis		40.64	0.31	URE 1	1		
Blautia producta		9.77	0.07		99		
47577550	**Very good identification**						**47577550**
					COC		
Blautia producta		97.76	0.66	SAL 99	99		
Actinomyces israelii		1.83	0.56		1		
Bifidobacterium adolescentis		0.19	0.33		1		
47577552	**Low discrimination**						**47577552**
					COC		
Bifidobacterium adolescentis		55.70	0.67	SOR 11	1		
Blautia producta		32.87	0.50		99		
Actinomyces israelii		8.31	0.59		1		
47577560	**Low discrimination**						**47577560**
					COC	SPO	
Blautia producta		79.61	0.41	SAL 99	99	1	
Actinomyces israelii		17.19	0.50	NAG 10	1	1	
Clostridium butyricum		2.10	0.13		1	99	
47577562	**Low discrimination**						**47577562**
					COC		
Actinomyces israelii		64.34	0.53	SAL 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		22.07	0.25	NAG 6	99		
Bifidobacterium adolescentis		8.34	0.31		1		
47577570	**Good identification**						**47577570**
					COC		
Blautia producta		97.23	0.50	SAL 99	99		
Bifidobacterium dentium		1.58	0.15	NAG 10	1		
Actinomyces israelii		1.05	0.36		1		
47577600	**Good identification**						**47577600**
					SPO		
Actinomyces israelii		97.33	0.48	SAL 92	1		
Clostridium butyricum		2.67	0.00	RHA 7	99		
47577602	**Low discrimination**						**47577602**
Actinomyces israelii		79.31	0.51	SAL 92			
Clostridium clostridioforme		19.02	0.31	RHA 7			
Bifidobacterium adolescentis		1.11	0.13				
47577612	**Low discrimination**						**47577612**
Clostridium clostridioforme		86.00	0.42	SAL 88			
Actinomyces israelii		11.76	0.37	MAN 1			
Bifidobacterium adolescentis		1.35	0.14				
47577632	**Low discrimination**						**47577632**
Clostridium clostridioforme		87.26	0.42	SAL 88			
Bacteroides ovatus		8.02	0.14	MAN 1			
Parabacteroides distasonis		1.66	0.18				
47577642	**Acceptable identification**						**47577642**
					COC		
Actinomyces israelii		90.44	0.44	SAL 92	1		
Blautia producta		9.57	0.07	RHA 7	99		
47577700	**Low discrimination**						**47577700**
					GRA	SPO	
Actinomyces israelii		68.11	0.58	SAL 92	99	1	
Clostridium butyricum		11.23	0.24	RHA 7	99	99	
Mitsuokella multacida		10.64	0.32		1	1	
47577702	**Low discrimination**						**47577702**
Actinomyces israelii		60.09	0.61	SAL 92			
Bifidobacterium adolescentis		20.79	0.47	RHA 7			
Clostridium clostridioforme		12.06	0.40				
47577712	**Low discrimination**						**47577712**
Clostridium clostridioforme		49.82	0.51	SAL 88			
Bifidobacterium adolescentis		23.04	0.48	MAN 1			
Bifidobacterium dentium		15.34	0.27				
47577722	**Low discrimination**						**47577722**
Bacteroides ovatus		71.18	0.51	IND 97			
Parabacteroides distasonis		18.23	0.57	SAL 88			
Bacteroides thetaiotaomicron		5.21	0.32				
47577732	**Low discrimination**						**47577732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		54.76	0.48	IND 97			
Clostridium clostridioforme		20.16	0.51	SAL 88			
Parabacteroides distasonis		11.33	0.52				
47577740	**Very good identification**						**47577740**
					COC	GRA	
Blautia producta		96.09	0.58	SAL 99	99	99	
Actinomyces israelii		2.72	0.51		1	99	
Mitsuokella multacida		1.09	0.32		1	1	
47577742	**Low discrimination**						**47577742**
					COC		
Blautia producta		69.32	0.41	SAL 99	99		
Actinomyces israelii		26.48	0.54	BGL 10	1		
Bifidobacterium adolescentis		2.91	0.31		1		
47577750	**Very good identification**						**47577750**
					COC	GRA	
Blautia producta		99.73	0.66	SAL 99	99	99	
Actinomyces israelii		0.14	0.37		1	99	
Mitsuokella multacida		0.13	0.24		1	1	
47577752	**Good identification**						**47577752**
					COC		
Blautia producta		96.03	0.50	SAL 99	99		
Actinomyces israelii		1.83	0.40	BGL 10	1		
Bifidobacterium adolescentis		1.64	0.33		1		
47577760	**Good identification**						**47577760**
					COC	GRA	
Blautia producta		95.84	0.41	SAL 99	99	99	
Mitsuokella multacida		2.45	0.21	NAG 10	1	1	
Actinomyces israelii		1.56	0.31		1	99	
47577770	**Good identification**						**47577770**
					COC	GRA	
Blautia producta		99.64	0.50	SAL 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
Actinomyces israelii		0.08	0.17		1	99	
47600000	**Low discrimination**						**47600000**
					GRA		
Fusobacterium mortiferum		89.99	0.57	FRU 99	1		
Eubacterium tortuosum		10.01	0.24		99		
47600002	**Low discrimination**						**47600002**
					GRA		
Fusobacterium mortiferum		84.99	0.40	FRU 99	1		
Eubacterium tortuosum		15.01	0.11	BGL 10	99		
47600100	**Very good identification**						**47600100**
Fusobacterium mortiferum		100.0	0.63	FRU 99			
47600102	**Good identification**						**47600102**
					GRA		
Fusobacterium mortiferum		99.19	0.47	FRU 99	1		
Bifidobacterium longum subsp. infantis		0.81	-.07	BGL 10	99		
47601000	**Low discrimination**						**47601000**
					GRA		
Eubacterium tortuosum		88.01	0.58	MNS 1	99		
Fusobacterium mortiferum		11.99	0.60		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47601002	**Acceptable identification**					**47601002**
					GRA	
Eubacterium tortuosum		92.01	0.45	MNS 1	99	
Fusobacterium mortiferum		7.89	0.43		1	
Bifidobacterium longum subsp. infantis		0.10	-.07		99	
47601020	**Good identification**					**47601020**
					GRA	
Eubacterium tortuosum		99.23	0.45	MNS 1	99	
Fusobacterium mortiferum		0.77	0.25		1	
47601100	**Very good identification**					**47601100**
					GRA	
Fusobacterium mortiferum		96.25	0.66	FRU 99	1	
Eubacterium tortuosum		3.06	0.24		99	
Bifidobacterium longum subsp. infantis		0.57	0.10		99	
47601102	**Low discrimination**					**47601102**
					GRA	
Fusobacterium mortiferum		62.00	0.50	FRU 99	1	
Bifidobacterium longum subsp. infantis		33.60	0.27	BGL 10	99	
Eubacterium tortuosum		3.13	0.11		99	
47603101	**Good identification**					**47603101**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47603103	**Good identification**					**47603103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47603121	**Good identification**					**47603121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47603123	**Good identification**					**47603123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47604000	**Good identification**					**47604000**
Fusobacterium mortiferum		100.0	0.37	FRU 99		
47604100	**Good identification**					**47604100**
Fusobacterium mortiferum		100.0	0.44	FRU 99		
47605000	**Low discrimination**					**47605000**
					GRA	
Fusobacterium mortiferum		50.38	0.40	FRU 99	1	
Eubacterium tortuosum		49.62	0.24	XYL 7	99	
47605100	**Acceptable identification**					**47605100**
					GRA	
Fusobacterium mortiferum		93.92	0.47	FRU 99	1	
Bifidobacterium longum subsp. infantis		6.08	0.09	XYL 7	99	
47605322	**Good identification**					**47605322**
					GRA	
Parabacteroides distasonis		96.75	0.45	FRU 99	1	
Prevotella oralis		1.30	0.00		1	
Bifidobacterium longum subsp. infantis		1.15	-.04		99	
47605332	**Good identification**					**47605332**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		100.0	0.41	FRU 99				
47605722	**Good identification**							**47605722**
Parabacteroides distasonis		100.0	0.42	FRU 99				
47605732	**Good identification**							**47605732**
Parabacteroides distasonis		95.87	0.37	FRU 99				
Clostridium clostridioforme		4.13	0.08					
47610000	**Very good identification**							**47610000**
					GRA	CAT		
Fusobacterium mortiferum		99.98	0.91		1	5		
Propionibacterium granulosum		0.02	0.24		99	90		
47610001	**Very good identification**							**47610001**
Fusobacterium mortiferum		100.0	0.57	URE 1				
47610002	**Very good identification**							**47610002**
					GRA	CAT		
Fusobacterium mortiferum		99.94	0.74	BGL 10	1	5		
Propionibacterium granulosum		0.04	0.13		99	90		
Bifidobacterium longum subsp. infantis		0.02	-.07		99	1		
47610003	**Good identification**							**47610003**
Fusobacterium mortiferum		100.0	0.40	URE 1				
47610010	**Very good identification**							**47610010**
					GRA	SPO	CAT	
Fusobacterium mortiferum		98.83	0.57	MLZ 1	1	1	5	
Clostridium difficile		0.92	0.06		99	99	1	
Propionibacterium granulosum		0.25	0.09		99	1	90	
47610012	**Good identification**							**47610012**
Fusobacterium mortiferum		100.0	0.40	MLZ 1				
47610020	**Very good identification**							**47610020**
					GRA	CAT		
Fusobacterium mortiferum		99.55	0.57	NAG 1	1	5		
Propionibacterium granulosum		0.46	0.13		99	90		
47610022	**Acceptable identification**							**47610022**
					GRA			
Fusobacterium mortiferum		93.26	0.40	NAG 1	1			
Prevotella oralis		3.56	0.02	BGL 10	1			
Bifidobacterium longum subsp. infantis		3.18	-.03		99			
47610040	**Very good identification**							**47610040**
Fusobacterium mortiferum		100.0	0.57	SOR 1				
47610042	**Good identification**							**47610042**
Fusobacterium mortiferum		100.0	0.40	SOR 1				
47610100	**Excellent identification**							**47610100**
					GRA			
Fusobacterium mortiferum		99.99	0.97		1			
Bifidobacterium longum subsp. infantis		0.01	0.10		99			
Actinomyces naeslundii		0.00	0.26		99			
47610101	**Very good identification**							**47610101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		99.77	0.63	URE 1	1	
Actinomyces naeslundii		0.23	0.29		99	
47610102	**Very good identification**					**47610102**
					GRA	
Fusobacterium mortiferum		99.16	0.81	BGL 10	1	
Bifidobacterium longum subsp. infantis		0.81	0.27		99	
Actinomyces naeslundii		0.01	0.26		99	
47610103	**Good identification**					**47610103**
					GRA	
Fusobacterium mortiferum		97.22	0.47	URE 1	1	
Actinomyces naeslundii		1.99	0.29	BGL 10	99	
Bifidobacterium longum subsp. infantis		0.80	-.07		99	
47610110	**Very good identification**					**47610110**
Fusobacterium mortiferum		100.0	0.63	MLZ 1		
47610112	**Good identification**					**47610112**
					GRA	
Fusobacterium mortiferum		99.19	0.47	MLZ 1	1	
Bifidobacterium longum subsp. infantis		0.81	-.07	BGL 10	99	
47610120	**Very good identification**					**47610120**
					GRA	
Fusobacterium mortiferum		98.04	0.63	NAG 1	1	
Bifidobacterium longum subsp. infantis		1.56	0.15		99	
Actinomyces naeslundii		0.15	0.26		99	
47610140	**Very good identification**					**47610140**
Fusobacterium mortiferum		100.0	0.63	SOR 1		
47610142	**Good identification**					**47610142**
					GRA	
Fusobacterium mortiferum		99.19	0.47	SOR 1	1	
Bifidobacterium longum subsp. infantis		0.81	-.07	BGL 10	99	
47610200	**Very good identification**					**47610200**
Fusobacterium mortiferum		100.0	0.57	RHA 1		
47610202	**Good identification**					**47610202**
Fusobacterium mortiferum		100.0	0.40	RHA 1		
47610300	**Very good identification**					**47610300**
Fusobacterium mortiferum		100.0	0.63	RHA 1		
47610302	**Good identification**					**47610302**
					GRA	
Fusobacterium mortiferum		99.19	0.47	RHA 1	1	
Bifidobacterium longum subsp. infantis		0.81	-.07	BGL 10	99	
47610400	**Very good identification**					**47610400**
Fusobacterium mortiferum		100.0	0.57	ARA 1		
47610402	**Good identification**					**47610402**
Fusobacterium mortiferum		100.0	0.40	ARA 1		
47610500	**Very good identification**					**47610500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.63	ARA 1			
47610502	**Good identification**						**47610502**
					GRA		
Fusobacterium mortiferum		97.90	0.47	ARA 1	1		
Bifidobacterium adolescentis		1.30	0.11	BGL 10	99		
Bifidobacterium longum subsp. infantis		0.80	-.07		99		
47611000	**Excellent identification**						**47611000**
					GRA		
Fusobacterium mortiferum		99.86	0.94		1		
Eubacterium tortuosum		0.08	0.24		99		
Actinomyces naeslundii		0.02	0.42		99		
47611001	**Very good identification**						**47611001**
					GRA		
Fusobacterium mortiferum		97.11	0.60	URE 1	1		
Actinomyces naeslundii		2.89	0.45		99		
47611002	**Very good identification**						**47611002**
					GRA		
Fusobacterium mortiferum		98.13	0.77	BGL 10	1		
Bifidobacterium longum subsp. infantis		1.25	0.27		99		
Actinomyces naeslundii		0.18	0.42		99		
47611003	**Low discrimination**						**47611003**
					GRA		
Fusobacterium mortiferum		78.09	0.43	URE 1	1		
Actinomyces naeslundii		20.91	0.45	BGL 10	99		
Bifidobacterium longum subsp. infantis		1.00	-.07		99		
47611010	**Very good identification**						**47611010**
					GRA	CAT	
Fusobacterium mortiferum		99.56	0.60	MLZ 1	1	5	
Lactobacillus jensenii		0.27	0.06		99	1	
Propionibacterium granulosum		0.17	0.09		99	90	
47611012	**Good identification**						**47611012**
					GRA		
Fusobacterium mortiferum		97.16	0.43	MLZ 1	1		
Lactobacillus jensenii		1.60	0.03	BGL 10	99		
Bifidobacterium longum subsp. infantis		1.24	-.07		99		
47611020	**Good identification**						**47611020**
					GRA		
Fusobacterium mortiferum		91.44	0.60	NAG 1	1		
Bifidobacterium longum subsp. infantis		2.26	0.15		99		
Actinomyces naeslundii		1.81	0.42		99		
47611021	**Low discrimination**						**47611021**
					GRA		
Actinomyces naeslundii		74.66	0.45	ESL 12	99		
Fusobacterium mortiferum		25.34	0.25	MLT 94	1		
47611022	**Low discrimination**						**47611022**
					GRA		
Prevotella oralis		40.63	0.36	MLT 99	1		
Bifidobacterium longum subsp. infantis		36.23	0.32	LAC 94	99		
Fusobacterium mortiferum		16.12	0.43		1		
47611023	**Low discrimination**						**47611023**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		84.76	0.45	ESL 12	99	
Prevotella oralis		8.06	0.02	MLT 94	1	
Bifidobacterium longum subsp. infantis		7.19	-.03		99	
47611040	**Very good identification**					**47611040**
					GRA	
Fusobacterium mortiferum		99.65	0.60	SOR 1	1	
Actinomyces naeslundii		0.35	0.29		99	
47611042	**Good identification**					**47611042**
					GRA	
Fusobacterium mortiferum		95.76	0.43	SOR 1	1	
Actinomyces naeslundii		3.02	0.29	BGL 10	99	
Bifidobacterium longum subsp. infantis		1.22	-.07		99	
47611100	**Excellent identification**					**47611100**
					GRA	
Fusobacterium mortiferum		99.37	1.00		1	
Bifidobacterium longum subsp. infantis		0.59	0.44		99	
Actinomyces naeslundii		0.02	0.50		99	
47611101	**Very good identification**					**47611101**
					GRA	
Fusobacterium mortiferum		95.95	0.66	URE 1	1	
Actinomyces naeslundii		3.48	0.53		99	
Bifidobacterium longum subsp. infantis		0.57	0.10		99	
47611102	**Low discrimination**					**47611102**
					GRA	
Fusobacterium mortiferum		64.20	0.84	BGL 10	1	
Bifidobacterium longum subsp. infantis		34.79	0.62		99	
Prevotella oralis		0.58	0.35		1	
47611103	**Low discrimination**					**47611103**
					GRA	
Fusobacterium mortiferum		53.26	0.50	URE 1	1	
Bifidobacterium longum subsp. infantis		28.87	0.27	BGL 10	99	
Actinomyces naeslundii		17.40	0.53		99	
47611110	**Very good identification**					**47611110**
					GRA	
Fusobacterium mortiferum		99.20	0.66	MLZ 1	1	
Bifidobacterium longum subsp. infantis		0.59	0.10		99	
Actinomyces naeslundii		0.13	0.28		99	
47611112	**Low discrimination**					**47611112**
					GRA	
Fusobacterium mortiferum		55.00	0.50	MLZ 1	1	
Bifidobacterium longum subsp. infantis		29.81	0.27	BGL 10	99	
Bifidobacterium adolescentis		8.79	0.32		99	
47611120	**Low discrimination**					**47611120**
					GRA	
Bifidobacterium longum subsp. infantis		45.60	0.49	MLT 99	99	
Fusobacterium mortiferum		43.50	0.66	BGL 91	1	
Prevotella oralis		4.31	0.35		1	
47611121	**Low discrimination**					**47611121**
					GRA	
Actinomyces naeslundii		61.82	0.53	ESL 12	99	
Bifidobacterium longum subsp. infantis		18.03	0.15	MLT 94	99	
Fusobacterium mortiferum		17.20	0.32		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47611122	**Low discrimination**					**47611122**
					GRA	
Bifidobacterium longum subsp. infantis		81.49	0.66	MLT 99	99	
Prevotella oralis		14.46	0.56		1	
Parabacteroides distasonis		2.11	0.55		1	
47611132	**Low discrimination**					**47611132**
					GRA	
Parabacteroides distasonis		49.85	0.50	XYL 90	1	
Bifidobacterium longum subsp. infantis		36.20	0.32		99	
Prevotella oralis		6.42	0.22		1	
47611140	**Very good identification**					**47611140**
					GRA	
Fusobacterium mortiferum		98.91	0.66	SOR 1	1	
Bifidobacterium longum subsp. infantis		0.59	0.10		99	
Actinomyces naeslundii		0.42	0.37		99	
47611142	**Low discrimination**					**47611142**
					GRA	
Fusobacterium mortiferum		59.12	0.50	SOR 1	1	
Bifidobacterium longum subsp. infantis		32.04	0.27	BGL 10	99	
Bifidobacterium breve		4.50	0.27		99	
47611143	**Low discrimination**					**47611143**
					GRA	
Actinomyces naeslundii		78.73	0.40	ESL 12	99	
Fusobacterium mortiferum		13.79	0.15	MLT 94	1	
Bifidobacterium longum subsp. infantis		7.47	-.07		99	
47611161	**Good identification**					**47611161**
Actinomyces naeslundii		100.0	0.40	ESL 12		
47611163	**Low discrimination**					**47611163**
Actinomyces naeslundii		85.56	0.40	ESL 12		
Bifidobacterium longum subsp. infantis		14.44	-.03	MLT 94		
47611200	**Very good identification**					**47611200**
Fusobacterium mortiferum		100.0	0.60	RHA 1		
47611202	**Acceptable identification**					**47611202**
					GRA	
Fusobacterium mortiferum		93.49	0.43	RHA 1	1	
Prevotella oralis		5.32	0.08	BGL 10	1	
Bifidobacterium longum subsp. infantis		1.19	-.07		99	
47611300	**Very good identification**					**47611300**
					GRA	
Fusobacterium mortiferum		97.28	0.66	RHA 1	1	
Eubacterium contortum		1.92	0.24		99	
Bifidobacterium longum subsp. infantis		0.58	0.10		99	
47611302	**Low discrimination**					**47611302**
					GRA	
Fusobacterium mortiferum		49.12	0.50	RHA 1	1	
Bifidobacterium longum subsp. infantis		26.62	0.27	BGL 10	99	
Prevotella oralis		18.75	0.28		1	
47611322	**Low discrimination**					**47611322**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		47.24	0.63	XYL 90	1	
Prevotella oralis		46.35	0.50		1	
Bifidobacterium longum subsp. infantis		6.16	0.32		99	
47611332	**Very good identification**					**47611332**
					GRA	
Parabacteroides distasonis		97.81	0.58	XYL 90	1	
Prevotella oralis		1.80	0.16		1	
Bifidobacterium longum subsp. infantis		0.24	-.03		99	
47611400	**Very good identification**					**47611400**
Fusobacterium mortiferum		100.0	0.60	ARA 1		
47611402	**Good identification**					**47611402**
					GRA	
Fusobacterium mortiferum		96.77	0.43	ARA 1	1	
Bifidobacterium adolescentis		2.00	0.11	BGL 10	99	
Bifidobacterium longum subsp. infantis		1.24	-.07		99	
47611500	**Very good identification**					**47611500**
					GRA	
Fusobacterium mortiferum		99.12	0.66	ARA 1	1	
Bifidobacterium longum subsp. infantis		0.59	0.10		99	
Bifidobacterium adolescentis		0.10	0.11		99	
47611502	**Low discrimination**					**47611502**
					GRA	
Fusobacterium mortiferum		40.53	0.50	ARA 1	1	
Bifidobacterium adolescentis		35.49	0.45	BGL 10	99	
Bifidobacterium longum subsp. infantis		21.97	0.27		99	
47611512	**Good identification**					**47611512**
Bifidobacterium adolescentis		95.33	0.46	MLT 99		
Bifidobacterium longum subsp. longum		1.42	0.13	XYL 87		
Clostridium clostridioforme		1.14	0.10			
47611522	**Low discrimination**					**47611522**
					GRA	
Parabacteroides distasonis		54.53	0.51	XYL 90	1	
Bifidobacterium longum subsp. infantis		34.79	0.32		99	
Prevotella oralis		6.17	0.22		1	
47611532	**Low discrimination**					**47611532**
					GRA	
Parabacteroides distasonis		88.60	0.47	XYL 90	1	
Bifidobacterium longum subsp. longum		6.94	0.24		99	
Clostridium clostridioforme		1.39	0.10		50	
47611722	**Very good identification**					**47611722**
Parabacteroides distasonis		97.63	0.59	XYL 90		
Prevotella oralis		1.58	0.16			
Clostridium clostridioforme		0.44	0.14			
47611732	**Good identification**					**47611732**
Parabacteroides distasonis		96.31	0.55	XYL 90		
Clostridium clostridioforme		3.69	0.25			
47612000	**Very good identification**					**47612000**
					GRA SPO CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium mortiferum		96.16	0.57	MAN 1	1	1	5	
Clostridium difficile		1.65	0.11		99	99	1	
Propionibacterium granulosum		1.33	0.21		99	1	90	
47612002	**Acceptable identification**							**47612002**
					GRA	SPO	CAT	
Fusobacterium mortiferum		92.87	0.40	MAN 1	1	1	5	
Propionibacterium granulosum		2.88	0.11	BGL 10	99	1	90	
Clostridium innocuum		2.23	0.00		99	99	1	
47612100	**Very good identification**							**47612100**
					GRA			
Fusobacterium mortiferum		99.71	0.63	MAN 1	1			
Actinomyces naeslundii		0.29	0.31		99			
47612102	**Good identification**							**47612102**
					GRA			
Fusobacterium mortiferum		96.65	0.47	MAN 1	1			
Actinomyces naeslundii		2.55	0.31	BGL 10	99			
Bifidobacterium longum subsp. infantis		0.79	-.07		99			
47613000	**Good identification**							**47613000**
					GRA	SPO		
Fusobacterium mortiferum		92.46	0.60	MAN 1	1	1		
Actinomyces naeslundii		3.56	0.47		99	1		
Clostridium innocuum		3.12	0.22		99	99		
47613001	**Low discrimination**							**47613001**
					GRA			
Actinomyces naeslundii		85.12	0.50	ESL 12	99			
Fusobacterium mortiferum		14.88	0.25	MLT 94	1			
47613002	**Low discrimination**							**47613002**
					GRA	SPO		
Fusobacterium mortiferum		59.38	0.43	MAN 1	1	1		
Actinomyces naeslundii		20.58	0.47	BGL 10	99	1		
Clostridium innocuum		18.06	0.22		99	99		
47613003	**Good identification**							**47613003**
Actinomyces naeslundii		100.0	0.50	ESL 12				
47613020	**Low discrimination**							**47613020**
					GRA	CAT		
Actinomyces naeslundii		75.64	0.47	ESL 12	99	25		
Fusobacterium mortiferum		19.84	0.25	MLT 94	1	5		
Propionibacterium granulosum		4.52	0.11		99	90		
47613021	**Good identification**							**47613021**
Actinomyces naeslundii		100.0	0.50	ESL 12				
47613022	**Low discrimination**							**47613022**
					GRA			
Actinomyces naeslundii		87.80	0.47	ESL 12	99			
Prevotella oralis		6.45	0.02	MLT 94	1			
Bifidobacterium longum subsp. infantis		5.75	-.03		99			
47613023	**Good identification**							**47613023**
Actinomyces naeslundii		100.0	0.50	ESL 12				
47613041	**Good identification**							**47613041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ESL 12		
47613043	**Good identification**					**47613043**
Actinomyces naeslundii		100.0	0.37	ESL 12		
47613061	**Good identification**					**47613061**
Actinomyces naeslundii		100.0	0.37	ESL 12		
47613063	**Good identification**					**47613063**
Actinomyces naeslundii		100.0	0.37	ESL 12		
47613100	**Good identification**					**47613100**
					GRA	
Fusobacterium mortiferum		94.82	0.66	MAN 1	1	
Actinomyces naeslundii		4.45	0.55		99	
Bifidobacterium longum subsp. infantis		0.57	0.10		99	
47613101	**Low discrimination**					**47613101**
					GRA	
Actinomyces naeslundii		87.46	0.58	ESL 12	99	
Fusobacterium mortiferum		12.54	0.32	MLT 94	1	
47613102	**Low discrimination**					**47613102**
					GRA	
Fusobacterium mortiferum		46.12	0.50	MAN 1	1	
Bifidobacterium longum subsp. infantis		25.00	0.27	BGL 10	99	
Actinomyces naeslundii		19.50	0.55		99	
47613103	**Very good identification**					**47613103**
					GRA	
Actinomyces naeslundii		97.60	0.58	ESL 12	99	
Fusobacterium mortiferum		1.56	0.15	MLT 94	1	
Bifidobacterium longum subsp. infantis		0.84	-.07		99	
47613111	**Good identification**					**47613111**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47613113	**Good identification**					**47613113**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47613120	**Low discrimination**					**47613120**
					GRA	
Actinomyces naeslundii		65.57	0.55	ESL 12	99	
Bifidobacterium longum subsp. infantis		14.78	0.15	MLT 94	99	
Fusobacterium mortiferum		14.10	0.32		1	
47613121	**Very good identification**					**47613121**
Actinomyces naeslundii		100.0	0.58	ESL 12		
47613123	**Very good identification**					**47613123**
Actinomyces naeslundii		98.49	0.58	ESL 12		
Bifidobacterium longum subsp. infantis		1.51	-.03	MLT 94		
47613131	**Good identification**					**47613131**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47613133	**Good identification**					**47613133**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	ESL 12		
47613141	**Good identification**					**47613141**
Actinomyces naeslundii		100.0	0.45	ESL 12		
47613143	**Good identification**					**47613143**
Actinomyces naeslundii		100.0	0.45	ESL 12		
47613160	**Low discrimination**					**47613160**
Actinomyces naeslundii		84.78	0.42	ESL 12		
Bifidobacterium breve		15.22	0.15	MLT 94		
47613161	**Good identification**					**47613161**
Actinomyces naeslundii		100.0	0.45	ESL 12		
47613163	**Good identification**					**47613163**
Actinomyces naeslundii		100.0	0.45	ESL 12		
47613502	**Good identification**					**47613502**
					GRA	
Bifidobacterium adolescentis		96.64	0.43	MLT 99	99	
Fusobacterium mortiferum		1.36	0.15	XYL 87	1	
Actinomyces israelii		1.26	0.17		99	
47613512	**Good identification**					**47613512**
Bifidobacterium adolescentis		100.0	0.45	MLT 99		
47614000	**Very good identification**					**47614000**
Fusobacterium mortiferum		100.0	0.72	XYL 7		
47614001	**Good identification**					**47614001**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
47614002	**Good identification**					**47614002**
Fusobacterium mortiferum		100.0	0.55	XYL 7		
47614010	**Good identification**					**47614010**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
47614020	**Good identification**					**47614020**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
47614040	**Good identification**					**47614040**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
47614100	**Very good identification**					**47614100**
					GRA	
Fusobacterium mortiferum		99.90	0.78	XYL 7	1	
Bifidobacterium longum subsp. infantis		0.10	0.09		99	
47614101	**Good identification**					**47614101**
Fusobacterium mortiferum		100.0	0.44	XYL 7		
47614102	**Good identification**					**47614102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Fusobacterium mortiferum		91.66	0.61	XYL 7	1	
Bifidobacterium longum subsp. infantis		8.18	0.26	BGL 10	99	
Bifidobacterium adolescentis		0.16	0.11		99	
47614110	**Good identification**					**47614110**
Fusobacterium mortiferum		100.0	0.44	XYL 7		
47614120	**Low discrimination**					**47614120**
					GRA	
Fusobacterium mortiferum		85.28	0.44	XYL 7	1	
Bifidobacterium longum subsp. infantis		14.72	0.13	NAG 1	99	
47614140	**Good identification**					**47614140**
Fusobacterium mortiferum		100.0	0.44	XYL 7		
47614200	**Good identification**					**47614200**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
47614300	**Low discrimination**					**47614300**
					GRA	
Fusobacterium mortiferum		71.74	0.44	XYL 7	1	
Eubacterium contortum		28.26	0.24	RHA 1	99	
47614322	**Good identification**					**47614322**
					GRA	
Parabacteroides distasonis		96.75	0.45	GAL 99	1	
Prevotella oralis		1.30	0.00		1	
Bifidobacterium longum subsp. infantis		1.15	-.04		99	
47614332	**Acceptable identification**					**47614332**
Parabacteroides distasonis		94.73	0.41	GAL 99		
Clostridium clostridioforme		5.27	0.13			
47614400	**Good identification**					**47614400**
Fusobacterium mortiferum		100.0	0.37	XYL 7		
47614500	**Good identification**					**47614500**
Fusobacterium mortiferum		100.0	0.44	XYL 7		
47614722	**Low discrimination**					**47614722**
Parabacteroides distasonis		85.61	0.42	GAL 99		
Clostridium clostridioforme		14.39	0.23			
47615000	**Very good identification**					**47615000**
					GRA	
Fusobacterium mortiferum		99.85	0.75	XYL 7	1	
Bifidobacterium longum subsp. infantis		0.15	0.09		99	
47615001	**Good identification**					**47615001**
Fusobacterium mortiferum		100.0	0.40	XYL 7		
47615002	**Low discrimination**					**47615002**
					GRA	
Fusobacterium mortiferum		87.29	0.58	XYL 7	1	
Bifidobacterium longum subsp. infantis		12.12	0.26	BGL 10	99	
Bifidobacterium adolescentis		0.24	0.11		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47615010	**Good identification**					**47615010**
Fusobacterium mortiferum		100.0	0.40	XYL 7		
47615020	**Low discrimination**					**47615020**
					GRA	
Fusobacterium mortiferum		77.35	0.40	XYL 7	1	
Bifidobacterium longum subsp. infantis		20.78	0.13	NAG 1	99	
Prevotella oralis		1.87	-.02		1	
47615040	**Good identification**					**47615040**
Fusobacterium mortiferum		100.0	0.40	XYL 7		
47615100	**Good identification**					**47615100**
					GRA	
Fusobacterium mortiferum		93.64	0.81	XYL 7	1	
Bifidobacterium longum subsp. infantis		6.06	0.43		99	
Eubacterium contortum		0.25	0.24		99	
47615101	**Acceptable identification**					**47615101**
					GRA	
Fusobacterium mortiferum		93.92	0.47	XYL 7	1	
Bifidobacterium longum subsp. infantis		6.08	0.09	URE 1	99	
47615102	**Low discrimination**					**47615102**
					GRA	
Bifidobacterium longum subsp. infantis		83.39	0.60	MLT 99	99	
Fusobacterium mortiferum		14.15	0.64		1	
Bifidobacterium adolescentis		1.66	0.45		99	
47615110	**Acceptable identification**					**47615110**
					GRA	
Fusobacterium mortiferum		92.53	0.47	XYL 7	1	
Bifidobacterium longum subsp. infantis		5.99	0.09	MLZ 1	99	
Bifidobacterium adolescentis		1.48	0.12		99	
47615112	**Low discrimination**					**47615112**
					GRA	
Bifidobacterium adolescentis		62.30	0.46	MLT 99	99	
Bifidobacterium longum subsp. infantis		25.82	0.26	ARA 87	99	
Parabacteroides distasonis		7.02	0.32		1	
47615120	**Low discrimination**					**47615120**
					GRA	
Bifidobacterium longum subsp. infantis		84.32	0.47	MLT 99	99	
Fusobacterium mortiferum		7.40	0.47	BGL 91	1	
Bacteroides vulgatus		2.94	0.20		1	
47615122	**Low discrimination**					**47615122**
					GRA	
Bifidobacterium longum subsp. infantis		74.66	0.64	MLT 99	99	
Parabacteroides distasonis		21.21	0.71		1	
Prevotella oralis		2.00	0.41		1	
47615132	**Good identification**					**47615132**
					GRA	
Parabacteroides distasonis		93.28	0.67		1	
Bifidobacterium longum subsp. infantis		6.16	0.30		99	
Prevotella oralis		0.17	0.07		1	
47615140	**Acceptable identification**					**47615140**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		93.15	0.47	XYL 7	1	
Bifidobacterium longum subsp. infantis		6.03	0.09	SOR 1	99	
Actinomyces israelii		0.82	0.14		99	
47615200	**Low discrimination**					**47615200**
					GRA	
Fusobacterium mortiferum		62.00	0.40	XYL 7	1	
Eubacterium contortum		38.00	0.24	RHA 1	99	
47615222	**Acceptable identification**					**47615222**
					GRA	
Parabacteroides distasonis		90.48	0.45	LAC 99	1	
Prevotella oralis		7.70	0.14		1	
Bifidobacterium longum subsp. infantis		1.07	-.04		99	
47615232	**Good identification**					**47615232**
Parabacteroides distasonis		95.25	0.41	LAC 99		
Clostridium clostridioforme		4.75	0.13			
47615300	**Very good identification**					**47615300**
					GRA	
Eubacterium contortum		95.90	0.58	MLT 99	99	
Fusobacterium mortiferum		3.69	0.47		1	
Bifidobacterium longum subsp. infantis		0.24	0.09		99	
47615302	**Low discrimination**					**47615302**
					GRA	
Eubacterium contortum		71.16	0.45	MLT 99	99	
Parabacteroides distasonis		15.38	0.45		1	
Bifidobacterium longum subsp. infantis		10.15	0.26		99	
47615312	**Low discrimination**					**47615312**
					GRA	
Parabacteroides distasonis		77.91	0.41	NAG 99	1	
Clostridium clostridioforme		13.00	0.22		50	
Eubacterium contortum		6.76	0.11		99	
47615320	**Low discrimination**					**47615320**
					GRA	
Eubacterium contortum		49.09	0.45	MLT 99	99	
Bacteroides vulgatus		38.21	0.36		1	
Parabacteroides distasonis		10.61	0.45		1	
47615322	**Very good identification**					**47615322**
					GRA	
Parabacteroides distasonis		96.64	0.79		1	
Prevotella oralis		1.30	0.35		1	
Bifidobacterium longum subsp. infantis		1.15	0.30		99	
47615323	**Good identification**					**47615323**
					GRA	
Parabacteroides distasonis		96.75	0.45	URE 1	1	
Prevotella oralis		1.30	0.00		1	
Bifidobacterium longum subsp. infantis		1.15	-.04		99	
47615330	**Low discrimination**					**47615330**
					GRA	
Parabacteroides distasonis		86.63	0.41	BGL 99	1	
Eubacterium contortum		7.52	0.11		99	
Bacteroides vulgatus		5.85	0.02		1	
47615332	**Very good identification**					**47615332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		99.77	0.75				
Clostridium clostridioforme		0.17	0.22				
Prevotella oralis		0.03	0.00				
47615333	**Good identification**						**47615333**
Parabacteroides distasonis		100.0	0.41	URE 1			
47615362	**Good identification**						**47615362**
					GRA		
Parabacteroides distasonis		96.75	0.45	SOR 1	1		
Prevotella oralis		1.30	0.00		1		
Bifidobacterium longum subsp. infantis		1.15	-.04		99		
47615372	**Good identification**						**47615372**
Parabacteroides distasonis		100.0	0.41	SOR 1			
47615400	**Good identification**						**47615400**
Fusobacterium mortiferum		100.0	0.40	XYL 7			
47615500	**Low discrimination**						**47615500**
					GRA	SPO	
Fusobacterium mortiferum		63.06	0.47	XYL 7	1	1	
Eubacterium contortum		13.55	0.23	ARA 1	99	1	
Clostridium butyricum		6.65	0.11		99	99	
47615502	**Low discrimination**						**47615502**
					GRA		
Bifidobacterium adolescentis		88.19	0.59	MLT 99	99		
Bifidobacterium longum subsp. infantis		6.67	0.26		99		
Parabacteroides distasonis		2.07	0.33		1		
47615512	**Very good identification**						**47615512**
					GRA		
Bifidobacterium adolescentis		96.62	0.60	MLT 99	99		
Clostridium clostridioforme		1.74	0.27		50		
Parabacteroides distasonis		1.00	0.29		1		
47615520	**Acceptable identification**						**47615520**
					GRA		
Bacteroides vulgatus		90.70	0.39	MLT 99	1		
Parabacteroides distasonis		3.49	0.33	RHA 90	1		
Bifidobacterium longum subsp. longum		2.85	0.28		99		
47615522	**Good identification**						**47615522**
					GRA		
Parabacteroides distasonis		91.53	0.68		1		
Bifidobacterium longum subsp. infantis		5.31	0.30		99		
Bacteroides uniformis		1.18	0.16		1		
47615532	**Good identification**						**47615532**
					GRA		
Parabacteroides distasonis		94.81	0.63		1		
Bifidobacterium longum subsp. longum		2.48	0.32		99		
Clostridium clostridioforme		1.67	0.27		50		
47615542	**Good identification**						**47615542**
Bifidobacterium adolescentis		99.00	0.43	MLT 99			
Actinomyces israelii		1.00	0.15	SOR 11			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47615552	**Good identification**					**47615552**
Bifidobacterium adolescentis		100.0	0.45	MLT 99		
47615622	**Low discrimination**					**47615622**
Parabacteroides distasonis		86.91	0.42	LAC 99		
Clostridium clostridioforme		13.09	0.22			
47615700	**Very good identification**					**47615700**
					GRA	
Eubacterium contortum		97.75	0.57	MLT 99	99	
Bacteroides vulgatus		2.20	0.21		1	
Fusobacterium mortiferum		0.05	0.13		1	
47615702	**Low discrimination**					**47615702**
					GRA	
Eubacterium contortum		78.93	0.44	MLT 99	99	
Parabacteroides distasonis		12.78	0.42		1	
Clostridium clostridioforme		6.45	0.31		50	
47615712	**Low discrimination**					**47615712**
					GRA	
Clostridium clostridioforme		74.74	0.42	MLT 99	50	
Parabacteroides distasonis		17.52	0.37	TRE 88	1	
Bifidobacterium adolescentis		5.71	0.26		99	
47615720	**Acceptable identification**					**47615720**
					GRA	
Bacteroides vulgatus		91.58	0.56	MLT 99	1	
Eubacterium contortum		7.25	0.44		99	
Parabacteroides distasonis		1.17	0.42		1	
47615722	**Very good identification**					**47615722**
					GRA	
Parabacteroides distasonis		96.96	0.76		1	
Eubacterium contortum		1.07	0.31		99	
Bacteroides vulgatus		0.77	0.21		1	
47615723	**Good identification**					**47615723**
Parabacteroides distasonis		100.0	0.42	URE 1		
47615732	**Very good identification**					**47615732**
Parabacteroides distasonis		95.81	0.71			
Clostridium clostridioforme		4.13	0.42			
Bacteroides thetaiotaomicron		0.06	0.01			
47615733	**Good identification**					**47615733**
Parabacteroides distasonis		95.87	0.37	URE 1		
Clostridium clostridioforme		4.13	0.08			
47615762	**Good identification**					**47615762**
Parabacteroides distasonis		100.0	0.42	SOR 1		
47615772	**Good identification**					**47615772**
Parabacteroides distasonis		95.87	0.37	SOR 1		
Clostridium clostridioforme		4.13	0.08			
47616000	**Good identification**					**47616000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.37	MAN 1			
47616100	**Good identification**						**47616100**
Fusobacterium mortiferum		100.0	0.44	MAN 1			
47617000	**Acceptable identification**						**47617000**
					GRA	SPO	
Fusobacterium mortiferum		91.02	0.40	MAN 1	1	1	
Actinomyces israelii		6.83	0.23	XYL 7	99	1	
Clostridium innocuum		2.15	0.00		99	99	
47617100	**Low discrimination**						**47617100**
					GRA		
Fusobacterium mortiferum		83.05	0.47	MAN 1	1		
Actinomyces israelii		10.69	0.34	XYL 7	99		
Bifidobacterium longum subsp. infantis		5.38	0.09		99		
47617102	**Low discrimination**						**47617102**
Bifidobacterium adolescentis		52.45	0.43	MLT 99			
Bifidobacterium longum subsp. infantis		32.47	0.26	ARA 87			
Actinomyces israelii		9.57	0.37				
47617112	**Good identification**						**47617112**
Bifidobacterium adolescentis		97.82	0.45	MLT 99			
Actinomyces israelii		2.18	0.23	ARA 87			
47617322	**Good identification**						**47617322**
					GRA		
Parabacteroides distasonis		96.75	0.45	MAN 1	1		
Prevotella oralis		1.30	0.00		1		
Bifidobacterium longum subsp. infantis		1.15	-.04		99		
47617332	**Good identification**						**47617332**
Parabacteroides distasonis		100.0	0.41	MAN 1			
47617502	**Very good identification**						**47617502**
Bifidobacterium adolescentis		97.82	0.57	MLT 99			
Actinomyces israelii		2.18	0.35				
47617512	**Very good identification**						**47617512**
Bifidobacterium adolescentis		99.56	0.59	MLT 99			
Actinomyces israelii		0.27	0.21				
Bifidobacterium dentium		0.17	-.07				
47617542	**Acceptable identification**						**47617542**
Bifidobacterium adolescentis		93.44	0.42	MLT 99			
Actinomyces israelii		6.56	0.28	SOR 11			
47617552	**Good identification**						**47617552**
Bifidobacterium adolescentis		99.15	0.43	MLT 99			
Actinomyces israelii		0.85	0.14	SOR 11			
47617722	**Good identification**						**47617722**
Parabacteroides distasonis		100.0	0.42	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47617732	**Good identification**					**47617732**
Parabacteroides distasonis		95.87	0.37	MAN 1		
Clostridium clostridioforme		4.13	0.08			
47620000	**Good identification**					**47620000**
Fusobacterium mortiferum		100.0	0.48	FRU 99		
47620100	**Good identification**					**47620100**
Fusobacterium mortiferum		100.0	0.54	FRU 99		
47620102	**Good identification**					**47620102**
Fusobacterium mortiferum		100.0	0.38	FRU 99		
47621000	**Low discrimination**					**47621000**
					GRA	
Fusobacterium mortiferum		79.68	0.51	FRU 99	1	
Eubacterium tortuosum		19.78	0.24		99	
Actinomyces naeslundii		0.54	0.25		99	
47621100	**Very good identification**					**47621100**
					GRA	
Fusobacterium mortiferum		99.18	0.57	FRU 99	1	
Actinomyces naeslundii		0.82	0.33		99	
47621101	**Low discrimination**					**47621101**
					GRA	
Actinomyces naeslundii		55.14	0.36	ESL 12	99	
Fusobacterium mortiferum		44.86	0.23	MLT 94	1	
47621102	**Acceptable identification**					**47621102**
					GRA	
Fusobacterium mortiferum		91.51	0.41	FRU 99	1	
Actinomyces naeslundii		6.82	0.33	BGL 10	99	
Bifidobacterium longum subsp. infantis		1.68	-.07		99	
47621103	**Good identification**					**47621103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47621121	**Good identification**					**47621121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47621123	**Good identification**					**47621123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47621322	**Good identification**					**47621322**
Parabacteroides distasonis		100.0	0.40	FRU 99		
47621332	**Good identification**					**47621332**
Parabacteroides distasonis		100.0	0.36	FRU 99		
47621722	**Good identification**					**47621722**
Parabacteroides distasonis		100.0	0.37	FRU 99		
47623100	**Low discrimination**					**47623100**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		61.40	0.38	ESL 12	99	
Fusobacterium mortiferum		38.60	0.23	MLT 94	1	
47623101	**Good identification**					**47623101**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47623102	**Good identification**					**47623102**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47623103	**Good identification**					**47623103**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47623120	**Good identification**					**47623120**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47623121	**Good identification**					**47623121**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47623122	**Good identification**					**47623122**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47623123	**Good identification**					**47623123**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47625100	**Good identification**					**47625100**
Fusobacterium mortiferum		100.0	0.38	FRU 99		
47625122	**Good identification**					**47625122**
					GRA	
Parabacteroides distasonis		99.23	0.48	FRU 99	1	
Bifidobacterium longum subsp. infantis		0.77	-.04		99	
47625132	**Good identification**					**47625132**
Parabacteroides distasonis		100.0	0.44	FRU 99		
47625322	**Very good identification**					**47625322**
Parabacteroides distasonis		100.0	0.56	FRU 99		
47625332	**Good identification**					**47625332**
Parabacteroides distasonis		100.0	0.52	FRU 99		
47625522	**Good identification**					**47625522**
Parabacteroides distasonis		100.0	0.45	FRU 99		
47625532	**Good identification**					**47625532**
Parabacteroides distasonis		97.24	0.40	FRU 99		
Clostridium clostridioforme		2.76	0.08			
47625722	**Good identification**					**47625722**
Parabacteroides distasonis		99.19	0.53	FRU 99		
Clostridium clostridioforme		0.81	0.11			
47625732	**Acceptable identification**					**47625732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		93.51	0.48	FRU 99			
Clostridium clostridioforme		6.49	0.23				
47630000	**Very good identification**						**47630000**
					GRA	CAT	
Fusobacterium mortiferum		99.81	0.82		1	5	
Propionibacterium granulosum		0.19	0.32		99	90	
Actinomyces naeslundii		0.01	0.23		99	25	
47630001	**Good identification**						**47630001**
					GRA		
Fusobacterium mortiferum		98.81	0.48	URE 1	1		
Actinomyces naeslundii		1.20	0.26		99		
47630002	**Very good identification**						**47630002**
					GRA	CAT	
Fusobacterium mortiferum		99.51	0.65	BGL 10	1	5	
Propionibacterium granulosum		0.42	0.21		99	90	
Actinomyces naeslundii		0.07	0.23		99	25	
47630010	**Good identification**						**47630010**
					GRA	CAT	
Fusobacterium mortiferum		97.56	0.48	MLZ 1	1	5	
Propionibacterium granulosum		2.44	0.17		99	90	
47630020	**Acceptable identification**						**47630020**
					GRA	CAT	
Fusobacterium mortiferum		94.88	0.48	NAG 1	1	5	
Propionibacterium granulosum		4.36	0.21		99	90	
Actinomyces naeslundii		0.77	0.23		99	25	
47630040	**Good identification**						**47630040**
Fusobacterium mortiferum		100.0	0.48	SOR 1			
47630100	**Very good identification**						**47630100**
					GRA		
Fusobacterium mortiferum		99.99	0.88		1		
Actinomyces naeslundii		0.01	0.31		99		
47630101	**Good identification**						**47630101**
					GRA		
Fusobacterium mortiferum		98.55	0.54	URE 1	1		
Actinomyces naeslundii		1.45	0.34		99		
47630102	**Very good identification**						**47630102**
					GRA		
Fusobacterium mortiferum		99.88	0.72	BGL 10	1		
Actinomyces naeslundii		0.09	0.31		99		
Bifidobacterium longum subsp. infantis		0.03	-.07		99		
47630103	**Low discrimination**						**47630103**
					GRA		
Fusobacterium mortiferum		88.29	0.38	URE 1	1		
Actinomyces naeslundii		11.71	0.34	BGL 10	99		
47630110	**Good identification**						**47630110**
Fusobacterium mortiferum		100.0	0.54	MLZ 1			
47630112	**Good identification**						**47630112**
Fusobacterium mortiferum		96.88	0.38	MLZ 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		3.12	-.11	BGL 10			
47630120	**Good identification**						**47630120**
					GRA		
Fusobacterium mortiferum		99.03	0.54	NAG 1	1		
Actinomyces naeslundii		0.97	0.31		99		
47630122	**Low discrimination**						**47630122**
					GRA		
Fusobacterium mortiferum		56.77	0.38	NAG 1	1		
Parabacteroides distasonis		32.33	0.32	BGL 10	1		
Actinomyces naeslundii		5.02	0.31		99		
47630140	**Good identification**						**47630140**
Fusobacterium mortiferum		100.0	0.54	SOR 1			
47630142	**Good identification**						**47630142**
Fusobacterium mortiferum		100.0	0.38	SOR 1			
47630200	**Good identification**						**47630200**
Fusobacterium mortiferum		100.0	0.48	RHA 1			
47630300	**Good identification**						**47630300**
Fusobacterium mortiferum		100.0	0.54	RHA 1			
47630302	**Good identification**						**47630302**
Fusobacterium mortiferum		100.0	0.38	RHA 1			
47630322	**Good identification**						**47630322**
Parabacteroides distasonis		100.0	0.40	GAL 99			
47630332	**Acceptable identification**						**47630332**
Parabacteroides distasonis		92.62	0.36	GAL 99			
Clostridium clostridioforme		7.38	0.11	XYL 90			
47630400	**Good identification**						**47630400**
Fusobacterium mortiferum		100.0	0.48	ARA 1			
47630500	**Good identification**						**47630500**
Fusobacterium mortiferum		100.0	0.54	ARA 1			
47630502	**Acceptable identification**						**47630502**
					GRA		
Fusobacterium mortiferum		93.99	0.38	ARA 1	1		
Clostridium clostridioforme		3.23	0.05	BGL 10	50		
Bifidobacterium adolescentis		2.78	0.07		99		
47630722	**Low discrimination**						**47630722**
Parabacteroides distasonis		80.59	0.37	GAL 99			
Clostridium clostridioforme		19.41	0.20	XYL 90			
47631000	**Very good identification**						**47631000**
					GRA	CAT	
Fusobacterium mortiferum		99.66	0.85		1	5	
Actinomyces naeslundii		0.13	0.47		99	25	
Propionibacterium granulosum		0.12	0.32		99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47631001	**Low discrimination**						**47631001**
					GRA		
Fusobacterium mortiferum		83.80	0.51	URE 1	1		
Actinomyces naeslundii		16.21	0.50		99		
47631002	**Very good identification**						**47631002**
					GRA		
Fusobacterium mortiferum		97.87	0.68	BGL 10	1		
Actinomyces naeslundii		1.15	0.47		99		
Lactobacillus jensenii		0.49	0.19		99		
47631003	**Low discrimination**						**47631003**
					GRA		
Actinomyces naeslundii		63.51	0.50	ESL 12	99		
Fusobacterium mortiferum		36.49	0.34	MLT 94	1		
47631010	**Good identification**						**47631010**
					GRA	CAT	
Fusobacterium mortiferum		95.15	0.51	MLZ 1	1	5	
Lactobacillus jensenii		2.62	0.14		99	1	
Propionibacterium granulosum		1.59	0.17		99	90	
47631020	**Low discrimination**						**47631020**
					GRA		
Fusobacterium mortiferum		81.05	0.51	NAG 1	1		
Actinomyces naeslundii		10.45	0.47		99		
Clostridium perfringens		4.35	0.18		99		
47631021	**Good identification**						**47631021**
					GRA		
Actinomyces naeslundii		95.04	0.50	ESL 12	99		
Fusobacterium mortiferum		4.96	0.16	MLT 94	1		
47631022	**Low discrimination**						**47631022**
					GRA		
Actinomyces naeslundii		27.24	0.47	ESL 12	99		
Fusobacterium mortiferum		23.47	0.34	MLT 94	1		
Parabacteroides distasonis		20.79	0.32		1		
47631023	**Good identification**						**47631023**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47631040	**Good identification**						**47631040**
					GRA		
Fusobacterium mortiferum		97.78	0.51	SOR 1	1		
Actinomyces naeslundii		2.22	0.34		99		
47631041	**Low discrimination**						**47631041**
					GRA		
Actinomyces naeslundii		77.16	0.37	ESL 12	99		
Fusobacterium mortiferum		22.84	0.16	MLT 94	1		
47631043	**Good identification**						**47631043**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47631061	**Good identification**						**47631061**
Actinomyces naeslundii		100.0	0.37	ESL 12			
47631063	**Good identification**						**47631063**
Actinomyces naeslundii		100.0	0.37	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47631100	**Very good identification**						**47631100**
					COC	GRA	
Fusobacterium mortiferum		99.78	0.91		1	1	
Actinomyces naeslundii		0.16	0.55		1	99	
Atopobium parvulum		0.04	0.21		99	99	
47631101	**Low discrimination**						**47631101**
					GRA		
Fusobacterium mortiferum		80.91	0.57	URE 1	1		
Actinomyces naeslundii		19.09	0.58		99		
47631102	**Good identification**						**47631102**
					GRA		
Fusobacterium mortiferum		94.94	0.75	BGL 10	1		
Bifidobacterium longum subsp. infantis		1.74	0.27		99		
Actinomyces naeslundii		1.36	0.55		99		
47631103	**Low discrimination**						**47631103**
					GRA		
Actinomyces naeslundii		67.58	0.58	ESL 12	99		
Fusobacterium mortiferum		31.83	0.41	MLT 94	1		
Bifidobacterium longum subsp. infantis		0.58	-.07		99		
47631110	**Very good identification**						**47631110**
					GRA		
Fusobacterium mortiferum		98.95	0.57	MLZ 1	1		
Actinomyces naeslundii		0.82	0.33		99		
Leptotrichia buccalis		0.24	-.11		1		
47631111	**Low discrimination**						**47631111**
					GRA		
Actinomyces naeslundii		55.14	0.36	ESL 12	99		
Fusobacterium mortiferum		44.86	0.23	MLT 94	1		
47631113	**Low discrimination**						**47631113**
					GRA		
Actinomyces naeslundii		83.88	0.36	ESL 12	99		
Leptotrichia buccalis		16.12	-.11	MLT 94	1		
47631120	**Low discrimination**						**47631120**
					GRA		
Fusobacterium mortiferum		68.43	0.57	NAG 1	1		
Clostridium perfringens		11.53	0.33		99		
Actinomyces naeslundii		10.76	0.55		99		
47631121	**Very good identification**						**47631121**
					GRA		
Actinomyces naeslundii		95.89	0.58	ESL 12	99		
Fusobacterium mortiferum		4.11	0.23	MLT 94	1		
47631122	**Low discrimination**						**47631122**
					GRA		
Parabacteroides distasonis		76.57	0.66	XYL 90	1		
Prevotella oralis		7.37	0.36		1		
Bifidobacterium longum subsp. infantis		6.57	0.32		99		
47631123	**Low discrimination**						**47631123**
					GRA		
Actinomyces naeslundii		82.55	0.58	ESL 12	99		
Parabacteroides distasonis		14.76	0.32	MLT 94	1		
Prevotella oralis		1.42	0.02		1		
47631131	**Good identification**						**47631131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	ESL 12		
47631132	**Very good identification**					**47631132**
					GRA	
Parabacteroides distasonis		96.65	0.62	XYL 90	1	
Bifidobacterium breve		1.44	0.29		99	
Leptotrichia buccalis		1.13	0.07		1	
47631140	**Very good identification**					**47631140**
					GRA	
Fusobacterium mortiferum		97.30	0.57	SOR 1	1	
Actinomyces naeslundii		2.70	0.42		99	
47631141	**Low discrimination**					**47631141**
					GRA	
Actinomyces naeslundii		80.47	0.45	ESL 12	99	
Fusobacterium mortiferum		19.53	0.23	MLT 94	1	
47631142	**Low discrimination**					**47631142**
					GRA	
Fusobacterium mortiferum		63.15	0.41	SOR 1	1	
Actinomyces naeslundii		15.77	0.42	BGL 10	99	
Bifidobacterium breve		10.74	0.24		99	
47631143	**Good identification**					**47631143**
Actinomyces naeslundii		100.0	0.45	ESL 12		
47631161	**Good identification**					**47631161**
Actinomyces naeslundii		100.0	0.45	ESL 12		
47631163	**Good identification**					**47631163**
Actinomyces naeslundii		100.0	0.45	ESL 12		
47631200	**Good identification**					**47631200**
Fusobacterium mortiferum		100.0	0.51	RHA 1		
47631222	**Acceptable identification**					**47631222**
Parabacteroides distasonis		92.01	0.40	XYL 90		
Prevotella oralis		7.99	0.09	LAC 99		
47631232	**Acceptable identification**					**47631232**
Parabacteroides distasonis		93.33	0.36	XYL 90		
Clostridium clostridioforme		6.67	0.10	LAC 99		
47631300	**Very good identification**					**47631300**
Fusobacterium mortiferum		100.0	0.57	RHA 1		
47631302	**Low discrimination**					**47631302**
Parabacteroides distasonis		48.72	0.40	XYL 90		
Fusobacterium mortiferum		42.78	0.41	NAG 99		
Prevotella oralis		3.49	0.08			
47631312	**Low discrimination**					**47631312**
Parabacteroides distasonis		78.48	0.36	XYL 90		
Clostridium clostridioforme		18.77	0.19	NAG 99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		2.75	-.11				
47631320	**Low discrimination**						**47631320**
					SPO		
Parabacteroides distasonis		81.57	0.40	XYL 90	1		
Fusobacterium mortiferum		6.51	0.23	BGL 99	1		
Clostridium ramosum		6.07	0.09		92		
47631322	**Very good identification**						**47631322**
					SPO		
Parabacteroides distasonis		98.24	0.74	XYL 90	1		
Prevotella oralis		1.35	0.30		1		
Clostridium ramosum		0.30	0.19		92		
47631323	**Good identification**						**47631323**
					GRA		
Parabacteroides distasonis		98.15	0.40	XYL 90	1		
Actinomyces naeslundii		1.85	0.24	URE 1	99		
47631330	**Good identification**						**47631330**
Parabacteroides distasonis		100.0	0.36	XYL 90			
47631332	**Very good identification**						**47631332**
Parabacteroides distasonis		99.76	0.70	XYL 90			
Clostridium clostridioforme		0.24	0.19				
47631333	**Good identification**						**47631333**
Parabacteroides distasonis		100.0	0.36	XYL 90			
47631362	**Good identification**						**47631362**
Parabacteroides distasonis		100.0	0.40	XYL 90			
47631372	**Good identification**						**47631372**
Parabacteroides distasonis		100.0	0.36	XYL 90			
47631400	**Good identification**						**47631400**
Fusobacterium mortiferum		100.0	0.51	ARA 1			
47631500	**Very good identification**						**47631500**
					GRA	SPO	
Fusobacterium mortiferum		98.52	0.57	ARA 1	1	1	
Clostridium butyricum		1.27	0.06		99	99	
Bifidobacterium adolescentis		0.22	0.07		99	1	
47631502	**Low discrimination**						**47631502**
					GRA		
Bifidobacterium adolescentis		58.52	0.42	MLT 99	99		
Fusobacterium mortiferum		29.94	0.41	XYL 87	1		
Parabacteroides distasonis		6.97	0.28		1		
47631512	**Low discrimination**						**47631512**
					GRA		
Bifidobacterium adolescentis		83.85	0.43	MLT 99	99		
Clostridium clostridioforme		11.00	0.25	XYL 87	50		
Parabacteroides distasonis		4.40	0.24		1		
47631522	**Very good identification**						**47631522**
Parabacteroides distasonis		99.19	0.62	XYL 90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.30	0.14				
Prevotella oralis		0.16	0.02				
47631532	**Very good identification**						**47631532**
					GRA		
Parabacteroides distasonis		96.95	0.58	XYL 90	1		
Clostridium clostridioforme		2.45	0.25		50		
Bifidobacterium longum subsp. longum		0.42	0.13		99		
47631622	**Low discrimination**						**47631622**
Parabacteroides distasonis		82.25	0.37	XYL 90			
Clostridium clostridioforme		17.75	0.19	LAC 99			
47631702	**Low discrimination**						**47631702**
					GRA		
Parabacteroides distasonis		57.12	0.37	XYL 90	1		
Clostridium clostridioforme		41.27	0.28	NAG 99	50		
Bifidobacterium adolescentis		1.62	0.07		99		
47631712	**Low discrimination**						**47631712**
					GRA		
Clostridium clostridioforme		85.17	0.40	MLT 99	50		
Parabacteroides distasonis		13.93	0.32	XYL 91	1		
Bifidobacterium adolescentis		0.89	0.09		99		
47631720	**Low discrimination**						**47631720**
Parabacteroides distasonis		88.41	0.37	XYL 90			
Bacteroides vulgatus		11.59	0.03	BGL 99			
47631722	**Very good identification**						**47631722**
Parabacteroides distasonis		99.18	0.71	XYL 90			
Clostridium clostridioforme		0.72	0.28				
Bacteroides thetaiotaomicron		0.10	0.03				
47631723	**Good identification**						**47631723**
Parabacteroides distasonis		100.0	0.37	XYL 90			
47631732	**Good identification**						**47631732**
Parabacteroides distasonis		94.12	0.66	XYL 90			
Clostridium clostridioforme		5.81	0.40				
Bacteroides thetaiotaomicron		0.07	-.04				
47631762	**Good identification**						**47631762**
Parabacteroides distasonis		100.0	0.37	XYL 90			
47632000	**Low discrimination**						**47632000**
					GRA	CAT	
Fusobacterium mortiferum		86.64	0.48	MAN 1	1	5	
Propionibacterium granulosum		12.01	0.30		99	90	
Actinomyces naeslundii		1.36	0.28		99	25	
47632100	**Good identification**						**47632100**
					GRA		
Fusobacterium mortiferum		98.13	0.54	MAN 1	1		
Actinomyces naeslundii		1.87	0.36		99		
47632101	**Low discrimination**						**47632101**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		73.92	0.39	ESL 12	99		
Fusobacterium mortiferum		26.09	0.20	MLT 94	1		
47632102	**Low discrimination**						**47632102**
					GRA		
Fusobacterium mortiferum		85.35	0.38	MAN 1	1		
Actinomyces naeslundii		14.66	0.36	BGL 10	99		
47632103	**Good identification**						**47632103**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47632120	**Low discrimination**						**47632120**
					GRA		
Actinomyces naeslundii		65.39	0.36	ESL 12	99		
Fusobacterium mortiferum		34.61	0.20	MLT 94	1		
47632121	**Good identification**						**47632121**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47632122	**Good identification**						**47632122**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47632123	**Good identification**						**47632123**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47633000	**Low discrimination**						**47633000**
					GRA	CAT	
Fusobacterium mortiferum		72.01	0.51	MAN 1	1	5	
Actinomyces naeslundii		18.02	0.52		99	25	
Propionibacterium granulosum		6.66	0.30		99	90	
47633001	**Good identification**						**47633001**
					GRA		
Actinomyces naeslundii		97.38	0.55	ESL 12	99		
Fusobacterium mortiferum		2.62	0.16	MLT 94	1		
47633002	**Low discrimination**						**47633002**
					GRA	SPO	
Actinomyces naeslundii		57.73	0.52	ESL 12	99	1	
Fusobacterium mortiferum		25.63	0.34	MLT 94	1	1	
Clostridium innocuum		9.17	0.14		99	99	
47633003	**Good identification**						**47633003**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47633020	**Low discrimination**						**47633020**
					GRA	CAT	
Actinomyces naeslundii		88.29	0.52	ESL 12	99	25	
Propionibacterium granulosum		8.15	0.19	MLT 94	99	90	
Fusobacterium mortiferum		3.56	0.16		1	5	
47633021	**Good identification**						**47633021**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47633022	**Good identification**						**47633022**
Actinomyces naeslundii		97.75	0.52	ESL 12			
Propionibacterium granulosum		2.26	0.09	MLT 94			
47633023	**Good identification**						**47633023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	ESL 12		
47633040	**Low discrimination**					**47633040**
					GRA	
Actinomyces naeslundii		81.39	0.39	ESL 12	99	
Fusobacterium mortiferum		18.61	0.16	MLT 94	1	
47633041	**Good identification**					**47633041**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47633042	**Good identification**					**47633042**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47633043	**Good identification**					**47633043**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47633060	**Good identification**					**47633060**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47633061	**Good identification**					**47633061**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47633062	**Good identification**					**47633062**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47633063	**Good identification**					**47633063**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47633100	**Low discrimination**					**47633100**
					GRA	
Fusobacterium mortiferum		75.99	0.57	MAN 1	1	
Actinomyces naeslundii		23.20	0.60		99	
Actinomyces israelii		0.81	0.25		99	
47633101	**Very good identification**					**47633101**
					GRA	
Actinomyces naeslundii		97.84	0.63	ESL 12	99	
Fusobacterium mortiferum		2.16	0.23	MLT 94	1	
47633102	**Low discrimination**					**47633102**
					GRA	
Actinomyces naeslundii		62.95	0.60	ESL 12	99	
Fusobacterium mortiferum		22.91	0.41	MLT 94	1	
Bifidobacterium adolescentis		5.47	0.26		99	
47633103	**Very good identification**					**47633103**
Actinomyces naeslundii		100.0	0.63	ESL 12		
47633110	**Low discrimination**					**47633110**
					GRA	
Actinomyces naeslundii		61.40	0.38	ESL 12	99	
Fusobacterium mortiferum		38.60	0.23	MLT 94	1	
47633111	**Good identification**					**47633111**
Actinomyces naeslundii		100.0	0.41	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47633113	**Good identification**						**47633113**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47633120	**Low discrimination**						**47633120**
					SPO	GRA	
Actinomyces naeslundii		85.05	0.60	ESL 12	1	99	
Clostridium ramosum		11.22	0.19	MLT 94	92	50	
Fusobacterium mortiferum		2.81	0.23		1	1	
47633121	**Very good identification**						**47633121**
Actinomyces naeslundii		100.0	0.63	ESL 12			
47633122	**Low discrimination**						**47633122**
					SPO	GRA	
Actinomyces naeslundii		55.01	0.60	ESL 12	1	99	
Clostridium ramosum		29.02	0.30	MLT 94	92	50	
Parabacteroides distasonis		7.60	0.32		1	1	
47633123	**Very good identification**						**47633123**
Actinomyces naeslundii		100.0	0.63	ESL 12			
47633130	**Low discrimination**						**47633130**
Actinomyces naeslundii		82.87	0.38	ESL 12			
Bifidobacterium breve		17.13	0.12	MLT 94			
47633131	**Good identification**						**47633131**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47633133	**Good identification**						**47633133**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47633140	**Low discrimination**						**47633140**
					GRA		
Actinomyces naeslundii		79.11	0.47	ESL 12	99		
Fusobacterium mortiferum		14.83	0.23	MLT 94	1		
Actinomyces israelii		6.06	0.18		99		
47633141	**Good identification**						**47633141**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47633142	**Low discrimination**						**47633142**
Actinomyces naeslundii		65.51	0.47	ESL 12			
Bifidobacterium breve		22.97	0.24	MLT 94			
Actinomyces israelii		7.52	0.21				
47633143	**Good identification**						**47633143**
Actinomyces naeslundii		100.0	0.50	ESL 12			
47633160	**Acceptable identification**						**47633160**
Actinomyces naeslundii		94.19	0.47	ESL 12			
Bifidobacterium breve		5.81	0.12	MLT 94			
47633161	**Good identification**						**47633161**
Actinomyces naeslundii		100.0	0.50	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47633162	**Low discrimination**					**47633162**
Actinomyces naeslundii		61.60	0.47	ESL 12		
Bifidobacterium breve		38.40	0.29	MLT 94		
47633163	**Good identification**					**47633163**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47633332	**Good identification**					**47633332**
					SPO	
Parabacteroides distasonis		96.93	0.36	MAN 1	1	
Clostridium ramosum		3.07	-.03	XYL 90	92	
47633502	**Acceptable identification**					**47633502**
Bifidobacterium adolescentis		91.69	0.40	MLT 99		
Actinomyces israelii		6.72	0.27	XYL 87		
Actinomyces naeslundii		1.59	0.26			
47633512	**Good identification**					**47633512**
Bifidobacterium adolescentis		98.34	0.42	MLT 99		
Bifidobacterium dentium		1.66	-.07	XYL 87		
47633722	**Good identification**					**47633722**
					SPO	
Parabacteroides distasonis		97.30	0.37	MAN 1	1	
Clostridium ramosum		2.71	-.03	XYL 90	92	
47634000	**Very good identification**					**47634000**
Fusobacterium mortiferum		100.0	0.63	XYL 7		
47634002	**Good identification**					**47634002**
Fusobacterium mortiferum		100.0	0.46	XYL 7		
47634100	**Very good identification**					**47634100**
Fusobacterium mortiferum		100.0	0.69	XYL 7		
47634102	**Good identification**					**47634102**
					GRA	
Fusobacterium mortiferum		98.92	0.52	XYL 7	1	
Parabacteroides distasonis		0.69	0.14	BGL 10	1	
Bifidobacterium adolescentis		0.39	0.07		99	
47634122	**Good identification**					**47634122**
					GRA	
Parabacteroides distasonis		97.80	0.48	GAL 99	1	
Fusobacterium mortiferum		1.44	0.18		1	
Bifidobacterium longum subsp. infantis		0.76	-.04		99	
47634132	**Good identification**					**47634132**
Parabacteroides distasonis		96.47	0.44	GAL 99		
Clostridium clostridioforme		3.53	0.13			
47634322	**Very good identification**					**47634322**
Parabacteroides distasonis		98.95	0.56	GAL 99		
Clostridium clostridioforme		1.05	0.17			
47634332	**Acceptable identification**					**47634332**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		91.78	0.52	GAL 99		
Clostridium clostridioforme		8.22	0.28			
47634522	**Acceptable identification**					**47634522**
Parabacteroides distasonis		90.03	0.45	GAL 99		
Clostridium clostridioforme		9.97	0.23			
47634532	**Low discrimination**					**47634532**
Parabacteroides distasonis		51.64	0.40	GAL 99		
Clostridium clostridioforme		48.36	0.34			
47634612	**Good identification**					**47634612**
Clostridium clostridioforme		100.0	0.40	MLT 99		
47634632	**Good identification**					**47634632**
Clostridium clostridioforme		98.54	0.40	MLT 99		
Parabacteroides distasonis		1.46	0.14			
47634702	**Good identification**					**47634702**
Clostridium clostridioforme		96.40	0.37	MLT 99		
Parabacteroides distasonis		3.60	0.19			
47634712	**Good identification**					**47634712**
Clostridium clostridioforme		99.56	0.49	MLT 99		
Parabacteroides distasonis		0.44	0.14			
47634722	**Low discrimination**					**47634722**
Parabacteroides distasonis		78.71	0.53	GAL 99		
Clostridium clostridioforme		21.30	0.37			
47634732	**Low discrimination**					**47634732**
Clostridium clostridioforme		69.60	0.49	MLT 99		
Parabacteroides distasonis		30.40	0.48			
47635000	**Very good identification**					**47635000**
					GRA	
Fusobacterium mortiferum		99.83	0.66	XYL 7	1	
Actinomyces israelii		0.17	0.20		99	
47635002	**Good identification**					**47635002**
					GRA	
Fusobacterium mortiferum		96.18	0.49	XYL 7	1	
Actinomyces israelii		2.19	0.23	BGL 10	99	
Parabacteroides distasonis		1.04	0.14		1	
47635022	**Good identification**					**47635022**
Parabacteroides distasonis		97.50	0.48	LAC 99		
Fusobacterium mortiferum		0.92	0.15			
Prevotella oralis		0.81	0.00			
47635032	**Good identification**					**47635032**
Parabacteroides distasonis		96.82	0.44	LAC 99		
Clostridium clostridioforme		3.18	0.13			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47635100	**Very good identification**							**47635100**
					COC	GRA	SPO	
Fusobacterium mortiferum		97.93	0.72	XYL 7	1	1	1	
Atopobium parvulum		0.89	0.25		99	99	1	
Clostridium butyricum		0.61	0.15		1	99	99	
47635101	**Good identification**							**47635101**
					GRA			
Fusobacterium mortiferum		96.93	0.38	XYL 7	1			
Actinomyces naeslundii		3.07	0.24	URE 1	99			
47635102	**Low discrimination**							**47635102**
					GRA			
Fusobacterium mortiferum		49.90	0.56	XYL 7	1			
Parabacteroides distasonis		22.88	0.48	BGL 10	1			
Bifidobacterium adolescentis		13.09	0.42		99			
47635110	**Low discrimination**							**47635110**
					GRA			
Fusobacterium mortiferum		87.86	0.38	XYL 7	1			
Actinomyces israelii		3.76	0.17	MLZ 1	99			
Bifidobacterium adolescentis		3.13	0.09		99			
47635112	**Low discrimination**							**47635112**
					GRA			
Bifidobacterium adolescentis		52.26	0.43	MLT 99	99			
Parabacteroides distasonis		40.25	0.44	ARA 87	1			
Clostridium clostridioforme		4.42	0.22		50			
47635120	**Low discrimination**							**47635120**
					GRA			
Parabacteroides distasonis		74.94	0.48	BGL 99	1			
Fusobacterium mortiferum		14.86	0.38		1			
Bifidobacterium longum subsp. infantis		5.73	0.13		99			
47635122	**Very good identification**							**47635122**
					GRA			
Parabacteroides distasonis		99.03	0.82		1			
Bifidobacterium longum subsp. infantis		0.77	0.30		99			
Prevotella oralis		0.13	0.20		1			
47635123	**Good identification**							**47635123**
					GRA			
Parabacteroides distasonis		98.61	0.48	URE 1	1			
Bifidobacterium longum subsp. infantis		0.77	-.04		99			
Actinomyces naeslundii		0.62	0.24		99			
47635130	**Good identification**							**47635130**
Parabacteroides distasonis		100.0	0.44	BGL 99				
47635132	**Very good identification**							**47635132**
					GRA			
Parabacteroides distasonis		99.86	0.78		1			
Clostridium clostridioforme		0.11	0.22		50			
Bifidobacterium adolescentis		0.01	0.09		99			
47635133	**Good identification**							**47635133**
Parabacteroides distasonis		100.0	0.44	URE 1				
47635140	**Low discrimination**							**47635140**
					GRA	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		85.86	0.38	XYL 7	1	1	
Actinomyces israelii		9.55	0.24	SOR 1	99	1	
Clostridium butyricum		4.59	-.03		99	99	
47635162	**Good identification**						**47635162**
					GRA		
Parabacteroides distasonis		99.23	0.48	SOR 1	1		
Bifidobacterium longum subsp. infantis		0.77	-.04		99		
47635172	**Good identification**						**47635172**
Parabacteroides distasonis		100.0	0.44	SOR 1			
47635222	**Very good identification**						**47635222**
Parabacteroides distasonis		99.06	0.56	LAC 99			
Clostridium clostridioforme		0.94	0.16				
47635232	**Acceptable identification**						**47635232**
Parabacteroides distasonis		92.57	0.52	LAC 99			
Clostridium clostridioforme		7.43	0.27				
47635300	**Low discrimination**						**47635300**
					GRA		
Fusobacterium mortiferum		49.21	0.38	XYL 7	1		
Eubacterium contortum		43.27	0.24	RHA 1	99		
Parabacteroides distasonis		7.52	0.22		1		
47635302	**Good identification**						**47635302**
					GRA		
Parabacteroides distasonis		94.93	0.56	NAG 99	1		
Clostridium clostridioforme		3.02	0.25		50		
Eubacterium contortum		0.97	0.11		99		
47635312	**Low discrimination**						**47635312**
					GRA		
Parabacteroides distasonis		78.55	0.52	NAG 99	1		
Clostridium clostridioforme		21.10	0.36		50		
Bifidobacterium adolescentis		0.34	0.09		99		
47635320	**Very good identification**						**47635320**
					GRA		
Parabacteroides distasonis		95.86	0.56	BGL 99	1		
Bacteroides vulgatus		3.16	0.13		1		
Eubacterium contortum		0.98	0.11		99		
47635322	**Very good identification**						**47635322**
Parabacteroides distasonis		99.95	0.91				
Clostridium clostridioforme		0.03	0.25				
Prevotella oralis		0.02	0.14				
47635323	**Very good identification**						**47635323**
Parabacteroides distasonis		100.0	0.56	URE 1			
47635330	**Good identification**						**47635330**
Parabacteroides distasonis		100.0	0.52	BGL 99			
47635332	**Very good identification**						**47635332**
Parabacteroides distasonis		99.73	0.86				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.27	0.36				
47635333	**Good identification**						**47635333**
Parabacteroides distasonis		100.0	0.52	URE 1			
47635362	**Very good identification**						**47635362**
Parabacteroides distasonis		100.0	0.56	SOR 1			
47635372	**Good identification**						**47635372**
Parabacteroides distasonis		100.0	0.52	SOR 1			
47635422	**Acceptable identification**						**47635422**
Parabacteroides distasonis		90.98	0.45	LAC 99			
Clostridium clostridioforme		9.02	0.22				
47635432	**Low discrimination**						**47635432**
Parabacteroides distasonis		54.37	0.40	LAC 99			
Clostridium clostridioforme		45.63	0.33				
47635502	**Low discrimination**						**47635502**
					GRA	SPO	
Bifidobacterium adolescentis		74.05	0.56	MLT 99	99	1	
Parabacteroides distasonis		11.86	0.45		1	1	
Clostridium butyricum		8.25	0.25		99	99	
47635512	**Low discrimination**						**47635512**
					GRA		
Bifidobacterium adolescentis		78.11	0.57	MLT 99	99		
Clostridium clostridioforme		15.48	0.42		50		
Parabacteroides distasonis		5.51	0.40		1		
47635520	**Low discrimination**						**47635520**
					GRA	SPO	
Parabacteroides distasonis		69.52	0.45	BGL 99	1	1	
Bacteroides vulgatus		16.54	0.16		1	1	
Clostridium butyricum		10.82	0.14		99	99	
47635522	**Very good identification**						**47635522**
					GRA		
Parabacteroides distasonis		99.43	0.79		1		
Clostridium clostridioforme		0.33	0.31		50		
Bifidobacterium adolescentis		0.06	0.22		99		
47635523	**Good identification**						**47635523**
Parabacteroides distasonis		100.0	0.45	URE 1			
47635530	**Low discrimination**						**47635530**
					GRA		
Parabacteroides distasonis		85.45	0.40	BGL 99	1		
Bifidobacterium longum subsp. longum		12.13	0.21		99		
Clostridium clostridioforme		2.43	0.08		50		
47635532	**Very good identification**						**47635532**
					GRA		
Parabacteroides distasonis		96.88	0.74		1		
Clostridium clostridioforme		2.75	0.42		50		
Bifidobacterium adolescentis		0.14	0.23		99		
47635533	**Good identification**						**47635533**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		97.24	0.40	URE 1		
Clostridium clostridioforme		2.76	0.08			
47635542	**Low discrimination**					**47635542**
					SPO	
Bifidobacterium adolescentis		87.06	0.40	MLT 99	1	
Clostridium butyricum		6.82	0.07	SOR 11	99	
Actinomyces israelii		4.98	0.25		1	
47635552	**Good identification**					**47635552**
Bifidobacterium adolescentis		98.41	0.42	MLT 99		
Clostridium clostridioforme		1.59	0.08	SOR 11		
47635562	**Good identification**					**47635562**
Parabacteroides distasonis		100.0	0.45	SOR 1		
47635572	**Good identification**					**47635572**
					GRA	
Parabacteroides distasonis		95.60	0.40	SOR 1	1	
Clostridium clostridioforme		2.71	0.08		50	
Bifidobacterium adolescentis		1.69	0.08		99	
47635602	**Good identification**					**47635602**
Clostridium clostridioforme		96.00	0.37	MLT 99		
Parabacteroides distasonis		4.00	0.19			
47635612	**Good identification**					**47635612**
Clostridium clostridioforme		99.51	0.48	MLT 99		
Parabacteroides distasonis		0.49	0.14			
47635622	**Low discrimination**					**47635622**
Parabacteroides distasonis		80.14	0.53	LAC 99		
Clostridium clostridioforme		19.43	0.37			
Bacteroides thetaiotaomicron		0.43	-.02			
47635632	**Low discrimination**					**47635632**
Clostridium clostridioforme		67.23	0.48	MLT 99		
Parabacteroides distasonis		32.77	0.48			
47635702	**Low discrimination**					**47635702**
					GRA	
Parabacteroides distasonis		53.95	0.53	NAG 99	1	
Clostridium clostridioforme		43.79	0.46		50	
Bifidobacterium adolescentis		1.14	0.22		99	
47635712	**Low discrimination**					**47635712**
					GRA	
Clostridium clostridioforme		86.76	0.57	MLT 99	50	
Parabacteroides distasonis		12.63	0.48		1	
Bifidobacterium adolescentis		0.60	0.23		99	
47635720	**Low discrimination**					**47635720**
					GRA	
Parabacteroides distasonis		57.61	0.53	BGL 99	1	
Bacteroides vulgatus		41.13	0.32		1	
Eubacterium contortum		0.79	0.10		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47635722	**Very good identification**						**47635722**
Parabacteroides distasonis		98.90	0.87				
Clostridium clostridioforme		0.81	0.46				
Bacteroides thetaiotaomicron		0.26	0.27				
47635723	**Good identification**						**47635723**
Parabacteroides distasonis		99.19	0.53	URE 1			
Clostridium clostridioforme		0.81	0.11				
47635730	**Acceptable identification**						**47635730**
Parabacteroides distasonis		92.36	0.48	BGL 99			
Clostridium clostridioforme		6.41	0.23				
Bacteroides vulgatus		1.24	-.02				
47635732	**Good identification**						**47635732**
Parabacteroides distasonis		93.33	0.82				
Clostridium clostridioforme		6.47	0.57				
Bacteroides thetaiotaomicron		0.18	0.20				
47635733	**Acceptable identification**						**47635733**
Parabacteroides distasonis		93.51	0.48	URE 1			
Clostridium clostridioforme		6.49	0.23				
47635762	**Good identification**						**47635762**
Parabacteroides distasonis		99.19	0.53	SOR 1			
Clostridium clostridioforme		0.81	0.11				
47635772	**Acceptable identification**						**47635772**
Parabacteroides distasonis		93.51	0.48	SOR 1			
Clostridium clostridioforme		6.49	0.23				
47637002	**Acceptable identification**						**47637002**
					GRA		
Actinomyces israelii		92.73	0.36	MLT 99	99		
Fusobacterium mortiferum		7.27	0.15	CEL 93	1		
47637100	**Low discrimination**						**47637100**
					GRA	SPO	
Actinomyces israelii		59.36	0.44	MLT 99	99	1	
Fusobacterium mortiferum		36.63	0.38	CEL 93	1	1	
Clostridium butyricum		2.50	-.01		99	99	
47637102	**Low discrimination**						**47637102**
					GRA		
Actinomyces israelii		48.67	0.47	MLT 99	99		
Bifidobacterium adolescentis		47.26	0.40	CEL 93	99		
Fusobacterium mortiferum		2.23	0.21		1		
47637112	**Low discrimination**						**47637112**
Bifidobacterium adolescentis		86.78	0.42	MLT 99			
Actinomyces israelii		10.93	0.33	ARA 87			
Bifidobacterium dentium		1.47	-.07				
47637122	**Good identification**						**47637122**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		95.54	0.48	MAN 1	1	
Actinomyces israelii		2.94	0.26		99	
Actinomyces naeslundii		0.78	0.26		99	
47637132	**Good identification**					**47637132**
					GRA	
Parabacteroides distasonis		95.91	0.44	MAN 1	1	
Bifidobacterium dentium		3.06	-.03		99	
Bifidobacterium adolescentis		1.03	0.07		99	
47637140	**Good identification**					**47637140**
Actinomyces israelii		100.0	0.37	MLT 99		
47637142	**Low discrimination**					**47637142**
Actinomyces israelii		76.42	0.40	MLT 99		
Bifidobacterium adolescentis		23.58	0.25	CEL 93		
47637322	**Very good identification**					**47637322**
					SPO	
Parabacteroides distasonis		99.81	0.56	MAN 1	1	
Clostridium ramosum		0.19	-.03		92	
47637332	**Good identification**					**47637332**
Parabacteroides distasonis		100.0	0.52	MAN 1		
47637500	**Low discrimination**					**47637500**
					SPO	
Actinomyces israelii		63.14	0.42	MLT 99	1	
Clostridium butyricum		29.27	0.15	CEL 93	99	
Bifidobacterium adolescentis		7.60	0.20		1	
47637502	**Low discrimination**					**47637502**
					SPO	
Bifidobacterium adolescentis		87.25	0.54	MLT 99	1	
Actinomyces israelii		10.99	0.45		1	
Clostridium butyricum		1.32	0.08		99	
47637512	**Low discrimination**					**47637512**
Bifidobacterium adolescentis		78.89	0.56	MLT 99		
Bifidobacterium dentium		19.70	0.27			
Actinomyces israelii		1.21	0.31			
47637522	**Low discrimination**					**47637522**
					GRA	
Parabacteroides distasonis		89.23	0.45	MAN 1	1	
Bifidobacterium adolescentis		4.61	0.20		99	
Actinomyces israelii		3.66	0.25		99	
47637542	**Low discrimination**					**47637542**
					COC	
Bifidobacterium adolescentis		67.28	0.39	MLT 99	1	
Actinomyces israelii		26.65	0.38	SOR 11	1	
Blautia producta		6.07	0.07		99	
47637552	**Low discrimination**					**47637552**
					COC	
Bifidobacterium adolescentis		77.51	0.40	MLT 99	1	
Blautia producta		17.16	0.15	SOR 11	99	
Actinomyces israelii		3.75	0.24		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47637722	**Good identification**					**47637722**
Parabacteroides distasonis		98.68	0.53	MAN 1		
Clostridium clostridioforme		0.81	0.11			
Bacteroides ovatus		0.51	-.02			
47637732	**Acceptable identification**					**47637732**
					GRA	
Parabacteroides distasonis		91.49	0.48	MAN 1	1	
Clostridium clostridioforme		6.35	0.23		50	
Bifidobacterium dentium		1.59	-.03		99	
47640000	**Good identification**					**47640000**
					GRA	
Fusobacterium mortiferum		96.43	0.48	FRU 99	1	
Eubacterium tortuosum		3.58	0.07		99	
47640100	**Good identification**					**47640100**
Fusobacterium mortiferum		100.0	0.55	FRU 99		
47640102	**Good identification**					**47640102**
Fusobacterium mortiferum		100.0	0.38	FRU 99		
47641000	**Low discrimination**					**47641000**
					GRA	
Eubacterium tortuosum		70.37	0.41	MNS 1	99	
Fusobacterium mortiferum		28.76	0.51	CEL 10	1	
Lactobacillus catenaformis		0.57	0.12		99	
47641100	**Very good identification**					**47641100**
					GRA	
Fusobacterium mortiferum		95.72	0.58	FRU 99	1	
Lactobacillus catenaformis		1.23	0.15		99	
Eubacterium tortuosum		1.01	0.07		99	
47641102	**Low discrimination**					**47641102**
					GRA	
Fusobacterium mortiferum		52.69	0.41	FRU 99	1	
Lactobacillus acidophilus		12.48	0.18	BGL 10	99	
Bifidobacterium breve		12.04	0.27		99	
47643101	**Good identification**					**47643101**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47643103	**Good identification**					**47643103**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47643121	**Good identification**					**47643121**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47643123	**Good identification**					**47643123**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47644100	**Good identification**					**47644100**
Fusobacterium mortiferum		100.0	0.35	FRU 99		
47645100	**Good identification**					**47645100**
					GRA SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		96.60	0.38	FRU 99	1	1	
Clostridium butyricum		3.40	-.05	XYL 7	99	99	
47645322	**Low discrimination**						**47645322**
Parabacteroides distasonis		87.21	0.41	FRU 99			
Prevotella oralis		7.74	0.10				
Bacteroides uniformis		5.06	0.01				
47645332	**Good identification**						**47645332**
Parabacteroides distasonis		100.0	0.36	FRU 99			
47650000	**Very good identification**						**47650000**
					GRA	SPO	
Fusobacterium mortiferum		99.92	0.83		1	1	
Clostridium barati		0.03	0.08		99	99	
Lactobacillus catenaformis		0.03	0.12		99	1	
47650001	**Good identification**						**47650001**
					GRA		
Fusobacterium mortiferum		99.27	0.48	URE 1	1		
Actinomyces naeslundii		0.73	0.24		99		
47650002	**Very good identification**						**47650002**
					GRA	SPO	
Fusobacterium mortiferum		98.72	0.66	BGL 10	1	1	
Clostridium barati		0.87	0.16		99	99	
Lactobacillus catenaformis		0.27	0.12		99	1	
47650010	**Good identification**						**47650010**
Fusobacterium mortiferum		100.0	0.48	MLZ 1			
47650020	**Acceptable identification**						**47650020**
					GRA	SPO	
Fusobacterium mortiferum		90.59	0.48	NAG 1	1	1	
Clostridium barati		8.74	0.16		99	99	
Lactobacillus catenaformis		0.68	0.02		99	1	
47650040	**Good identification**						**47650040**
Fusobacterium mortiferum		100.0	0.48	SOR 1			
47650100	**Very good identification**						**47650100**
					GRA	SPO	
Fusobacterium mortiferum		99.92	0.89		1	1	
Clostridium barati		0.04	0.16		99	99	
Lactobacillus catenaformis		0.02	0.15		99	1	
47650101	**Good identification**						**47650101**
					GRA		
Fusobacterium mortiferum		99.11	0.55	URE 1	1		
Actinomyces naeslundii		0.89	0.31		99		
47650102	**Very good identification**						**47650102**
					GRA	SPO	
Fusobacterium mortiferum		97.82	0.73	BGL 10	1	1	
Clostridium barati		1.10	0.24		99	99	
Lactobacillus acidophilus		0.35	0.18		99	1	
47650103	**Acceptable identification**						**47650103**
					GRA		
Fusobacterium mortiferum		92.49	0.38	URE 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		7.51	0.31	BGL 10	99		
47650110	**Good identification**						**47650110**
					GRA		
Fusobacterium mortiferum		99.63	0.55	MLZ 1	1		
Bifidobacterium breve		0.38	0.10		99		
47650112	**Low discrimination**						**47650112**
					GRA		
Fusobacterium mortiferum		72.91	0.38	MLZ 1	1		
Bifidobacterium breve		24.99	0.27	BGL 10	99		
Leptotrichia buccalis		2.10	-.11		1		
47650120	**Low discrimination**						**47650120**
					GRA	SPO	
Fusobacterium mortiferum		82.61	0.55	NAG 1	1	1	
Clostridium barati		10.25	0.24		99	99	
Clostridium paraputrificum		4.38	0.13		99	99	
47650140	**Good identification**						**47650140**
					GRA	SPO	
Fusobacterium mortiferum		99.27	0.55	SOR 1	1	1	
Bifidobacterium breve		0.37	0.10		99	1	
Clostridium barati		0.36	-.02		99	99	
47650142	**Low discrimination**						**47650142**
					GRA	SPO	
Fusobacterium mortiferum		67.40	0.38	SOR 1	1	1	
Bifidobacterium breve		23.10	0.27	BGL 10	99	1	
Clostridium barati		6.54	0.06		99	99	
47650200	**Good identification**						**47650200**
Fusobacterium mortiferum		100.0	0.48	RHA 1			
47650300	**Good identification**						**47650300**
					GRA	SPO	
Fusobacterium mortiferum		99.64	0.55	RHA 1	1	1	
Clostridium barati		0.36	-.02		99	99	
47650302	**Low discrimination**						**47650302**
					GRA	SPO	
Fusobacterium mortiferum		84.36	0.38	RHA 1	1	1	
Clostridium barati		8.19	0.06	BGL 10	99	99	
Prevotella oralis		5.19	0.04		1	1	
47650400	**Good identification**						**47650400**
Fusobacterium mortiferum		100.0	0.48	ARA 1			
47650500	**Good identification**						**47650500**
Fusobacterium mortiferum		100.0	0.55	ARA 1			
47650502	**Acceptable identification**						**47650502**
					GRA		
Fusobacterium mortiferum		92.20	0.38	ARA 1	1		
Bifidobacterium adolescentis		7.80	0.16	BGL 10	99		
47651000	**Very good identification**						**47651000**
					GRA		
Fusobacterium mortiferum		96.97	0.86		1		
Lactobacillus catenaformis		1.93	0.47		99		
Lactobacillus jensenii		0.46	0.36		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47651001	**Low discrimination**						**47651001**
					GRA		
Fusobacterium mortiferum		87.49	0.51	URE 1	1		
Actinomyces naeslundii		10.35	0.47		99		
Lactobacillus catenaformis		1.74	0.12		99		
47651002	**Low discrimination**						**47651002**
					GRA	SPO	
Fusobacterium mortiferum		71.23	0.69	BGL 10	1	1	
Lactobacillus catenaformis		12.76	0.47		99	1	
Clostridium barati		7.91	0.38		99	99	
47651003	**Low discrimination**						**47651003**
					GRA		
Actinomyces naeslundii		44.32	0.47	ESL 12	99		
Fusobacterium mortiferum		41.61	0.35	MLT 94	1		
Lactobacillus catenaformis		7.45	0.12		99		
47651010	**Low discrimination**						**47651010**
					GRA		
Fusobacterium mortiferum		84.31	0.51	MLZ 1	1		
Lactobacillus jensenii		13.17	0.28		99		
Lactobacillus catenaformis		1.68	0.12		99		
47651012	**Low discrimination**						**47651012**
					GRA		
Fusobacterium mortiferum		34.58	0.35	MLZ 1	1		
Lactobacillus jensenii		32.41	0.25	BGL 10	99		
Bifidobacterium breve		18.44	0.27		99		
47651020	**Low discrimination**						**47651020**
					GRA	SPO	
Clostridium barati		38.55	0.38	RAF 1	99	99	
Fusobacterium mortiferum		31.55	0.51		1	1	
Lactobacillus catenaformis		15.54	0.36		99	1	
47651021	**Low discrimination**						**47651021**
					GRA	SPO	
Actinomyces naeslundii		81.19	0.47	ESL 12	99	1	
Clostridium barati		8.47	0.04	MLT 94	99	99	
Fusobacterium mortiferum		6.93	0.17		1	1	
47651022	**Low discrimination**						**47651022**
					GRA	SPO	
Clostridium barati		45.30	0.46	RAF 1	99	99	
Prevotella oralis		36.81	0.45		1	1	
Lactobacillus acidophilus		6.53	0.34		99	1	
47651023	**Low discrimination**						**47651023**
					GRA	SPO	
Actinomyces naeslundii		60.45	0.47	ESL 12	99	1	
Clostridium barati		18.91	0.12	MLT 94	99	99	
Prevotella oralis		15.37	0.11		1	1	
47651040	**Acceptable identification**						**47651040**
					GRA	SPO	
Fusobacterium mortiferum		92.61	0.51	SOR 1	1	1	
Clostridium barati		3.28	0.11		99	99	
Lactobacillus catenaformis		1.84	0.12		99	1	
47651042	**Low discrimination**						**47651042**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		29.57	0.35	SOR 1	1	1	
Clostridium barati		28.27	0.20	BGL 10	99	99	
Lactobacillus acidophilus		17.39	0.18		99	1	
47651100	**Very good identification**						**47651100**
					GRA		
Fusobacterium mortiferum		96.94	0.92		1		
Lactobacillus catenaformis		1.24	0.50		99		
Collinsella aerofaciens		0.82	0.32		99		
47651101	**Low discrimination**						**47651101**
					GRA		
Fusobacterium mortiferum		85.22	0.58	URE 1	1		
Actinomyces naeslundii		12.30	0.55		99		
Lactobacillus catenaformis		1.09	0.15		99		
47651102	**Low discrimination**						**47651102**
					GRA		
Fusobacterium mortiferum		53.63	0.76	BGL 10	1		
Lactobacillus acidophilus		12.71	0.52		99		
Bifidobacterium breve		12.26	0.62		99		
47651103	**Low discrimination**						**47651103**
					GRA		
Actinomyces naeslundii		41.45	0.55	ESL 12	99		
Fusobacterium mortiferum		31.90	0.41	MLT 94	1		
Lactobacillus acidophilus		7.56	0.18		99		
47651110	**Low discrimination**						**47651110**
					GRA		
Fusobacterium mortiferum		77.98	0.58	MLZ 1	1		
Bifidobacterium breve		19.39	0.44		99		
Lactobacillus catenaformis		1.00	0.15		99		
47651112	**Low discrimination**						**47651112**
					GRA		
Bifidobacterium breve		83.11	0.62	MLT 99	99		
Leptotrichia buccalis		7.00	0.24		1		
Bifidobacterium adolescentis		3.75	0.38		99		
47651120	**Low discrimination**						**47651120**
					GRA	SPO	
Clostridium paraputrificum		42.59	0.47	MLT 99	99	99	
Clostridium barati		19.14	0.46	RAF 7	99	99	
Prevotella oralis		12.83	0.44		1	1	
47651121	**Low discrimination**						**47651121**
					SPO		
Actinomyces naeslundii		64.46	0.55	ESL 12	1		
Clostridium paraputrificum		15.77	0.13	MLT 94	99		
Clostridium barati		7.09	0.12		99		
47651122	**Low discrimination**						**47651122**
					GRA	SPO	
Prevotella oralis		57.49	0.66	MLT 99	1	1	
Clostridium barati		13.55	0.54		99	99	
Bifidobacterium breve		12.84	0.66		99	1	
47651130	**Low discrimination**						**47651130**
					SPO		
Bifidobacterium breve		84.26	0.49	MLT 99	1		
Clostridium paraputrificum		6.73	0.13	BGL 91	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		3.03	0.12		99		
47651132	**Low discrimination**						**47651132**
					GRA		
Bifidobacterium breve		88.73	0.66	MLT 99	99		
Parabacteroides distasonis		4.73	0.46		1		
Prevotella oralis		4.01	0.32		1		
47651140	**Low discrimination**						**47651140**
					GRA	SPO	
Fusobacterium mortiferum		74.59	0.58	SOR 1	1	1	
Bifidobacterium breve		18.54	0.44		99	1	
Clostridium barati		3.40	0.20		99	99	
47651141	**Low discrimination**						**47651141**
					GRA		
Actinomyces naeslundii		66.88	0.42	ESL 12	99		
Fusobacterium mortiferum		26.52	0.24	MLT 94	1		
Bifidobacterium breve		6.60	0.10		99		
47651142	**Low discrimination**						**47651142**
					SPO		
Bifidobacterium breve		80.04	0.62	MLT 99	1		
Lactobacillus acidophilus		10.26	0.36		1		
Clostridium barati		4.35	0.28		99		
47651143	**Low discrimination**						**47651143**
Actinomyces naeslundii		47.07	0.42	ESL 12			
Bifidobacterium breve		46.92	0.27	MLT 94			
Lactobacillus acidophilus		6.01	0.02				
47651150	**Good identification**						**47651150**
					GRA		
Bifidobacterium breve		96.10	0.44	MLT 99	99		
Fusobacterium mortiferum		3.90	0.24	BGL 91	1		
47651152	**Very good identification**						**47651152**
Bifidobacterium breve		99.23	0.62	MLT 99			
Bifidobacterium adolescentis		0.55	0.22				
Lactobacillus acidophilus		0.13	0.02				
47651160	**Low discrimination**						**47651160**
					SPO		
Bifidobacterium breve		49.89	0.49	MLT 99	1		
Clostridium paraputrificum		29.70	0.27	BGL 91	99		
Clostridium barati		15.42	0.28		99		
47651161	**Low discrimination**						**47651161**
					SPO		
Actinomyces naeslundii		78.15	0.42	ESL 12	1		
Bifidobacterium breve		13.70	0.15	MLT 94	1		
Clostridium paraputrificum		8.16	-.07		99		
47651162	**Low discrimination**						**47651162**
					SPO		
Bifidobacterium breve		80.56	0.66	MLT 99	1		
Clostridium barati		7.39	0.36		99		
Lactobacillus acidophilus		5.81	0.36		1		
47651170	**Good identification**						**47651170**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		99.40	0.49	MLT 99	1		
Clostridium paraputrificum		0.60	-.07	BGL 91	99		
47651172	**Very good identification**						**47651172**
					SPO		
Bifidobacterium breve		99.74	0.66	MLT 99	1		
Clostridium barati		0.09	0.02		99		
Lactobacillus acidophilus		0.07	0.02		1		
47651200	**Acceptable identification**						**47651200**
					GRA	SPO	
Fusobacterium mortiferum		94.34	0.51	RHA 1	1	1	
Clostridium barati		3.34	0.11		99	99	
Lactobacillus catenaformis		1.88	0.12		99	1	
47651202	**Low discrimination**						**47651202**
					GRA	SPO	
Fusobacterium mortiferum		32.27	0.35	RHA 1	1	1	
Clostridium barati		30.86	0.20	BGL 10	99	99	
Prevotella oralis		19.52	0.17		1	1	
47651222	**Low discrimination**						**47651222**
					GRA	SPO	
Prevotella oralis		76.15	0.39	MLT 99	1	1	
Clostridium barati		19.01	0.28	LAC 94	99	99	
Lactobacillus acidophilus		2.37	0.15		99	1	
47651300	**Good identification**						**47651300**
					GRA	SPO	
Fusobacterium mortiferum		91.32	0.58	RHA 1	1	1	
Clostridium barati		4.16	0.20		99	99	
Prevotella oralis		2.17	0.16		1	1	
47651302	**Low discrimination**						**47651302**
					GRA	SPO	
Prevotella oralis		47.46	0.38	MLT 99	1	1	
Lactobacillus acidophilus		20.64	0.33	NAG 95	99	1	
Clostridium barati		14.37	0.28		99	99	
47651320	**Low discrimination**						**47651320**
					SPO	GRA	
Prevotella oralis		63.59	0.38	MLT 99	1	1	
Clostridium barati		19.25	0.28	BGL 95	99	99	
Clostridium ramosum		7.52	0.22		92	50	
47651322	**Low discrimination**						**47651322**
					GRA	SPO	
Prevotella oralis		79.43	0.60	MLT 99	1	1	
Parabacteroides distasonis		12.29	0.58		1	1	
Clostridium barati		3.80	0.36		99	99	
47651332	**Low discrimination**						**47651332**
					GRA		
Parabacteroides distasonis		82.97	0.54	XYL 90	1		
Prevotella oralis		10.06	0.25		1		
Bifidobacterium breve		5.24	0.32		99		
47651400	**Good identification**						**47651400**
					GRA	SPO	
Fusobacterium mortiferum		95.27	0.51	ARA 1	1	1	
Lactobacillus catenaformis		1.90	0.12		99	1	
Clostridium innocuum		1.70	0.09		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47651402	**Low discrimination**						**47651402**
					GRA	SPO	
Fusobacterium mortiferum		59.92	0.35	ARA 1	1	1	
Lactobacillus catenaformis		10.73	0.12	BGL 10	99	1	
Clostridium innocuum		9.65	0.09		99	99	
47651500	**Low discrimination**						**47651500**
					GRA	SPO	
Fusobacterium mortiferum		89.84	0.58	ARA 1	1	1	
Clostridium butyricum		6.55	0.20		99	99	
Lactobacillus catenaformis		1.15	0.15		99	1	
47651502	**Low discrimination**						**47651502**
					GRA	SPO	
Bifidobacterium adolescentis		71.65	0.50	MLT 99	99	1	
Fusobacterium mortiferum		12.84	0.41	XYL 87	1	1	
Clostridium butyricum		3.28	0.13		99	99	
47651512	**Acceptable identification**						**47651512**
Bifidobacterium adolescentis		94.78	0.52	MLT 99			
Bifidobacterium breve		3.18	0.27	XYL 87			
Clostridium clostridioforme		1.78	0.19				
47651522	**Low discrimination**						**47651522**
Parabacteroides distasonis		39.31	0.47	XYL 90			
Prevotella oralis		29.30	0.32				
Bacteroides uniformis		8.44	0.16				
47651532	**Low discrimination**						**47651532**
					GRA		
Parabacteroides distasonis		67.88	0.42	XYL 90	1		
Bifidobacterium breve		20.99	0.32		99		
Clostridium clostridioforme		6.62	0.19		50		
47651552	**Low discrimination**						**47651552**
Bifidobacterium adolescentis		78.68	0.36	MLT 99			
Bifidobacterium breve		21.32	0.27	XYL 87			
47651722	**Low discrimination**						**47651722**
Parabacteroides distasonis		80.40	0.55	XYL 90			
Prevotella oralis		8.56	0.25				
Bacteroides uniformis		7.95	0.19				
47651732	**Low discrimination**						**47651732**
Parabacteroides distasonis		80.74	0.50	XYL 90			
Clostridium clostridioforme		19.26	0.34				
47652000	**Low discrimination**						**47652000**
					GRA	SPO	
Fusobacterium mortiferum		55.50	0.48	MAN 1	1	1	
Clostridium innocuum		43.97	0.34		99	99	
Actinomyces naeslundii		0.53	0.25		99	1	
47652100	**Good identification**						**47652100**
					GRA		
Fusobacterium mortiferum		97.80	0.55	MAN 1	1		
Actinomyces naeslundii		1.14	0.33		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		0.37	0.10		99		
47652101	**Low discrimination**						**47652101**
					GRA		
Actinomyces naeslundii		63.42	0.36	ESL 12	99		
Fusobacterium mortiferum		36.58	0.21	MLT 94	1		
47652102	**Low discrimination**						**47652102**
					GRA		
Fusobacterium mortiferum		63.46	0.38	MAN 1	1		
Bifidobacterium breve		21.75	0.27	BGL 10	99		
Actinomyces naeslundii		6.67	0.33		99		
47652103	**Good identification**						**47652103**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47652121	**Good identification**						**47652121**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47652123	**Good identification**						**47652123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47653000	**Low discrimination**						**47653000**
					GRA	SPO	
Clostridium innocuum		89.01	0.56	RAF 1	99	99	
Fusobacterium mortiferum		8.87	0.51		1	1	
Actinomyces naeslundii		1.36	0.49		99	1	
47653001	**Low discrimination**						**47653001**
					GRA	SPO	
Actinomyces naeslundii		67.33	0.52	ESL 12	99	1	
Clostridium innocuum		29.71	0.22	MLT 94	99	99	
Fusobacterium mortiferum		2.96	0.17		1	1	
47653002	**Good identification**						**47653002**
					GRA	SPO	
Clostridium innocuum		95.19	0.56	RAF 1	99	99	
Actinomyces naeslundii		1.45	0.49		99	1	
Fusobacterium mortiferum		1.05	0.35		1	1	
47653003	**Low discrimination**						**47653003**
					SPO		
Actinomyces naeslundii		69.39	0.52	ESL 12	1		
Clostridium innocuum		30.62	0.22	MLT 94	99		
47653020	**Low discrimination**						**47653020**
					SPO		
Actinomyces naeslundii		39.11	0.49	ESL 12	1		
Clostridium barati		27.13	0.20	MLT 94	99		
Clostridium innocuum		25.88	0.22		99		
47653021	**Good identification**						**47653021**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47653023	**Good identification**						**47653023**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47653041	**Good identification**						**47653041**
Actinomyces naeslundii		100.0	0.39	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47653043	**Good identification**					**47653043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47653060	**Low discrimination**					**47653060**
					SPO	
Actinomyces naeslundii		57.90	0.36	ESL 12	1	
Bifidobacterium breve		22.31	0.15	MLT 94	1	
Clostridium barati		19.79	0.01		99	
47653061	**Good identification**					**47653061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47653063	**Good identification**					**47653063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47653100	**Low discrimination**					**47653100**
					GRA	
Fusobacterium mortiferum		63.23	0.58	MAN 1	1	
Bifidobacterium breve		15.72	0.44		99	
Actinomyces naeslundii		11.81	0.57		99	
47653101	**Very good identification**					**47653101**
					GRA	
Actinomyces naeslundii		95.69	0.60	ESL 12	99	
Fusobacterium mortiferum		3.45	0.24	MLT 94	1	
Bifidobacterium breve		0.86	0.10		99	
47653102	**Low discrimination**					**47653102**
Bifidobacterium breve		76.36	0.62	MLT 99		
Actinomyces naeslundii		5.68	0.57			
Lactobacillus acidophilus		4.17	0.30			
47653103	**Good identification**					**47653103**
Actinomyces naeslundii		91.69	0.60	ESL 12		
Bifidobacterium breve		8.31	0.27	MLT 94		
47653110	**Acceptable identification**					**47653110**
					GRA	
Bifidobacterium breve		90.28	0.44	MLT 99	99	
Fusobacterium mortiferum		3.67	0.24	BGL 91	1	
Actinomyces naeslundii		3.57	0.35		99	
47653111	**Low discrimination**					**47653111**
Actinomyces naeslundii		85.45	0.38	ESL 12		
Bifidobacterium breve		14.55	0.10	MLT 94		
47653112	**Very good identification**					**47653112**
Bifidobacterium breve		95.51	0.62	MLT 99		
Bifidobacterium adolescentis		3.52	0.36			
Actinomyces naeslundii		0.37	0.35			
47653120	**Low discrimination**					**47653120**
					SPO	
Bifidobacterium breve		41.81	0.49	MLT 99	1	
Clostridium ramosum		21.57	0.33	BGL 91	92	
Actinomyces naeslundii		17.67	0.57		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47653121	**Very good identification**					**47653121**
					SPO	
Actinomyces naeslundii		97.64	0.60	ESL 12	1	
Bifidobacterium breve		1.56	0.15	MLT 94	1	
Clostridium ramosum		0.80	-.01		92	
47653122	**Low discrimination**					**47653122**
					SPO	
Bifidobacterium breve		70.35	0.66	MLT 99	1	
Clostridium ramosum		14.35	0.43		92	
Clostridium barati		6.45	0.36		99	
47653123	**Low discrimination**					**47653123**
					SPO	
Actinomyces naeslundii		82.24	0.60	ESL 12	1	
Bifidobacterium breve		13.25	0.32	MLT 94	1	
Clostridium ramosum		2.70	0.09		92	
47653130	**Good identification**					**47653130**
					SPO	
Bifidobacterium breve		96.46	0.49	MLT 99	1	
Actinomyces naeslundii		2.15	0.35	BGL 91	1	
Clostridium tertium		0.89	0.05		99	
47653131	**Low discrimination**					**47653131**
Actinomyces naeslundii		76.77	0.38	ESL 12		
Bifidobacterium breve		23.24	0.15	MLT 94		
47653132	**Very good identification**					**47653132**
					SPO	
Bifidobacterium breve		99.12	0.66	MLT 99	1	
Clostridium tertium		0.27	0.13		99	
Actinomyces naeslundii		0.22	0.35		1	
47653140	**Low discrimination**					**47653140**
Bifidobacterium breve		80.54	0.44	MLT 99		
Actinomyces naeslundii		10.68	0.44	BGL 91		
Actinomyces israelii		4.23	0.28			
47653141	**Good identification**					**47653141**
Actinomyces naeslundii		95.17	0.47	ESL 12		
Bifidobacterium breve		4.83	0.10	MLT 94		
47653142	**Very good identification**					**47653142**
Bifidobacterium breve		96.52	0.62	MLT 99		
Actinomyces naeslundii		1.27	0.44			
Actinomyces israelii		0.75	0.31			
47653143	**Low discrimination**					**47653143**
Actinomyces naeslundii		66.07	0.47	ESL 12		
Bifidobacterium breve		33.93	0.27	MLT 94		
47653150	**Good identification**					**47653150**
Bifidobacterium breve		99.22	0.44	MLT 99		
Actinomyces israelii		0.78	0.14	BGL 91		
47653152	**Very good identification**					**47653152**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		99.43	0.62	MLT 99			
Bifidobacterium adolescentis		0.45	0.21				
Actinomyces israelii		0.12	0.17				
47653160	**Low discrimination**						**47653160**
					SPO		
Bifidobacterium breve		89.87	0.49	MLT 99	1		
Actinomyces naeslundii		6.70	0.44	BGL 91	1		
Clostridium barati		2.42	0.10		99		
47653161	**Acceptable identification**						**47653161**
Actinomyces naeslundii		91.72	0.47	ESL 12			
Bifidobacterium breve		8.28	0.15	MLT 94			
47653162	**Very good identification**						**47653162**
					SPO		
Bifidobacterium breve		97.88	0.66	MLT 99	1		
Clostridium barati		0.78	0.18		99		
Actinomyces naeslundii		0.72	0.44		1		
47653163	**Low discrimination**						**47653163**
Actinomyces naeslundii		52.28	0.47	ESL 12			
Bifidobacterium breve		47.72	0.32	MLT 94			
47653170	**Good identification**						**47653170**
Bifidobacterium breve		100.0	0.49	MLT 99			
47653172	**Very good identification**						**47653172**
Bifidobacterium breve		100.0	0.66	MLT 99			
47653320	**Acceptable identification**						**47653320**
					SPO	GRA	
Clostridium ramosum		92.99	0.35	MLT 99	92	50	
Clostridium barati		3.66	0.10	TRE 94	99	99	
Prevotella oralis		1.40	0.04		1	1	
47653322	**Low discrimination**						**47653322**
					SPO	GRA	
Clostridium ramosum		86.23	0.45	MLT 99	92	50	
Prevotella oralis		6.18	0.25	TRE 94	1	1	
Bifidobacterium breve		3.22	0.32		1	99	
47653400	**Good identification**						**47653400**
					GRA	SPO	
Clostridium innocuum		98.55	0.43	RAF 1	99	99	
Actinomyces israelii		0.89	0.23		99	1	
Fusobacterium mortiferum		0.56	0.17		1	1	
47653402	**Good identification**						**47653402**
					SPO		
Clostridium innocuum		98.01	0.43	RAF 1	99		
Actinomyces israelii		1.33	0.26		1		
Bifidobacterium adolescentis		0.66	0.15		1		
47653502	**Low discrimination**						**47653502**
Bifidobacterium adolescentis		86.49	0.49	MLT 99			
Actinomyces israelii		7.02	0.36	XYL 87			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Bifidobacterium breve	4.33	0.27				
47653512	**Acceptable identification**						**47653512**
	Bifidobacterium adolescentis	94.68	0.50	MLT 99			
	Bifidobacterium breve	3.88	0.27	XYL 87			
	Actinomyces israelii	0.94	0.22				
47654000	**Very good identification**						**47654000**
	Fusobacterium mortiferum	100.0	0.63	XYL 7			
47654002	**Good identification**						**47654002**
	Fusobacterium mortiferum	100.0	0.47	XYL 7			
47654100	**Very good identification**						**47654100**
					GRA	SPO	
	Fusobacterium mortiferum	99.95	0.70	XYL 7	1	1	
	Clostridium butyricum	0.05	-.05		99	99	
47654101	**Good identification**						**47654101**
	Fusobacterium mortiferum	100.0	0.35	XYL 7			
47654102	**Good identification**						**47654102**
					GRA		
	Fusobacterium mortiferum	98.88	0.53	XYL 7	1		
	Bifidobacterium adolescentis	1.12	0.16	BGL 10	99		
47654110	**Good identification**						**47654110**
	Fusobacterium mortiferum	100.0	0.35	XYL 7			
47654120	**Good identification**						**47654120**
	Fusobacterium mortiferum	100.0	0.35	XYL 7			
47654140	**Good identification**						**47654140**
	Fusobacterium mortiferum	100.0	0.35	XYL 7			
47654300	**Good identification**						**47654300**
	Fusobacterium mortiferum	100.0	0.35	XYL 7			
47654322	**Low discrimination**						**47654322**
	Parabacteroides distasonis	84.21	0.41	GAL 99			
	Prevotella oralis	7.47	0.10				
	Bacteroides uniformis	4.88	0.01				
47654332	**Low discrimination**						**47654332**
	Parabacteroides distasonis	74.30	0.36	GAL 99			
	Clostridium clostridioforme	25.70	0.22				
47654500	**Low discrimination**						**47654500**
					GRA	SPO	
	Fusobacterium mortiferum	67.82	0.35	XYL 7	1	1	
	Clostridium butyricum	32.18	0.11	ARA 1	99	99	
47654712	**Good identification**						**47654712**
	Clostridium clostridioforme	100.0	0.43	MLT 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47654732	**Low discrimination**						**47654732**
Clostridium clostridioforme		89.84	0.43	MLT 99			
Parabacteroides distasonis		10.16	0.32	TRE 88			
47655000	**Very good identification**						**47655000**
					GRA	SPO	
Fusobacterium mortiferum		98.74	0.66	XYL 7	1	1	
Actinomyces israelii		0.53	0.30		99	1	
Clostridium butyricum		0.34	0.05		99	99	
47655002	**Low discrimination**						**47655002**
					GRA		
Fusobacterium mortiferum		84.97	0.50	XYL 7	1		
Actinomyces israelii		6.12	0.33	BGL 10	99		
Lactobacillus catenaformis		2.04	0.12		99		
47655100	**Very good identification**						**47655100**
					GRA	SPO	
Fusobacterium mortiferum		95.07	0.73	XYL 7	1	1	
Clostridium butyricum		3.34	0.29		99	99	
Actinomyces israelii		0.87	0.40		99	1	
47655101	**Good identification**						**47655101**
					GRA	SPO	
Fusobacterium mortiferum		96.60	0.38	XYL 7	1	1	
Clostridium butyricum		3.40	-.05	URE 1	99	99	
47655102	**Low discrimination**						**47655102**
					GRA		
Fusobacterium mortiferum		40.18	0.56	XYL 7	1		
Bifidobacterium adolescentis		30.10	0.50	BGL 10	99		
Bifidobacterium longum subsp. infantis		7.17	0.26		99		
47655110	**Low discrimination**						**47655110**
					GRA	SPO	
Fusobacterium mortiferum		68.80	0.38	XYL 7	1	1	
Clostridium butyricum		12.60	0.07	MLZ 1	99	99	
Actinomyces israelii		9.30	0.26		99	1	
47655112	**Low discrimination**						**47655112**
					GRA		
Bifidobacterium adolescentis		89.67	0.52	MLT 99	99		
Bifidobacterium breve		3.00	0.27	ARA 87	99		
Parabacteroides distasonis		2.56	0.28		1		
47655122	**Low discrimination**						**47655122**
Parabacteroides distasonis		51.81	0.67				
Prevotella oralis		32.18	0.50				
Bacteroides uniformis		6.52	0.32				
47655132	**Very good identification**						**47655132**
					GRA		
Parabacteroides distasonis		95.65	0.62		1		
Bifidobacterium breve		2.01	0.32		99		
Prevotella oralis		1.11	0.16		1		
47655140	**Low discrimination**						**47655140**
					GRA	SPO	
Fusobacterium mortiferum		59.24	0.38	XYL 7	1	1	
Actinomyces israelii		20.85	0.33	SOR 1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		17.93	0.11		99	99	
47655152	**Low discrimination**						**47655152**
Bifidobacterium adolescentis		74.94	0.36	MLT 99			
Bifidobacterium breve		20.31	0.27	ARA 87			
Actinomyces israelii		4.75	0.22	SOR 11			
47655222	**Low discrimination**						**47655222**
Parabacteroides distasonis		58.33	0.41	LAC 99			
Prevotella oralis		32.70	0.24				
Bacteroides uniformis		6.83	0.06				
47655232	**Low discrimination**						**47655232**
Parabacteroides distasonis		76.34	0.36	LAC 99			
Clostridium clostridioforme		23.66	0.22				
47655300	**Low discrimination**						**47655300**
					GRA	SPO	
Fusobacterium mortiferum		49.93	0.38	XYL 7	1	1	
Eubacterium contortum		39.35	0.24	RHA 1	99	1	
Clostridium butyricum		5.37	0.03		99	99	
47655302	**Low discrimination**						**47655302**
Parabacteroides distasonis		51.09	0.41	NAG 99			
Prevotella oralis		23.62	0.22				
Clostridium clostridioforme		6.27	0.19				
47655320	**Low discrimination**						**47655320**
Parabacteroides distasonis		51.05	0.41	BGL 99			
Prevotella oralis		23.60	0.22				
Bacteroides vulgatus		17.97	0.14				
47655322	**Low discrimination**						**47655322**
Parabacteroides distasonis		87.00	0.75				
Prevotella oralis		7.72	0.44				
Bacteroides uniformis		5.04	0.35				
47655323	**Low discrimination**						**47655323**
Parabacteroides distasonis		87.21	0.41	URE 1			
Prevotella oralis		7.74	0.10				
Bacteroides uniformis		5.06	0.01				
47655330	**Good identification**						**47655330**
Parabacteroides distasonis		100.0	0.36	BGL 99			
47655332	**Very good identification**						**47655332**
Parabacteroides distasonis		98.66	0.70				
Clostridium clostridioforme		1.03	0.31				
Prevotella oralis		0.16	0.10				
47655333	**Good identification**						**47655333**
Parabacteroides distasonis		100.0	0.36	URE 1			
47655362	**Low discrimination**						**47655362**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		87.21	0.41	SOR 1		
Prevotella oralis		7.74	0.10			
Bacteroides uniformis		5.06	0.01			
47655372	**Good identification**					**47655372**
Parabacteroides distasonis		100.0	0.36	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47655500	**Acceptable identification**						**47655500**
					GRA	SPO	
Clostridium butyricum		93.17	0.45	MLT 99	99	99	
Fusobacterium mortiferum		2.98	0.38	TRE 94	1	1	
Actinomyces israelii		2.20	0.39		99	1	
47655502	**Low discrimination**						**47655502**
					SPO		
Bifidobacterium adolescentis		78.70	0.64	MLT 99	1		
Clostridium butyricum		17.38	0.38		99		
Actinomyces israelii		1.59	0.42		1		
47655512	**Very good identification**						**47655512**
					SPO		
Bifidobacterium adolescentis		95.82	0.66	MLT 99	1		
Clostridium clostridioforme		2.72	0.36		66		
Clostridium butyricum		0.91	0.17		99		
47655522	**Low discrimination**						**47655522**
					GRA	SPO	
Bacteroides uniformis		68.62	0.51	IND 98	1	1	
Parabacteroides distasonis		29.04	0.63	MLT 92	1	1	
Clostridium butyricum		0.95	0.20		99	99	
47655532	**Low discrimination**						**47655532**
Parabacteroides distasonis		83.12	0.58				
Clostridium clostridioforme		9.11	0.36				
Bacteroides uniformis		3.68	0.16				
47655542	**Low discrimination**						**47655542**
					SPO		
Bifidobacterium adolescentis		82.05	0.49	MLT 99	1		
Clostridium butyricum		12.75	0.20	SOR 11	99		
Actinomyces israelii		5.20	0.35		1		
47655552	**Good identification**						**47655552**
					SPO		
Bifidobacterium adolescentis		98.58	0.50	MLT 99	1		
Actinomyces israelii		0.76	0.21	SOR 11	1		
Clostridium butyricum		0.66	-.02		99		
47655612	**Good identification**						**47655612**
Clostridium clostridioforme		100.0	0.42	MLT 99			
47655632	**Low discrimination**						**47655632**
Clostridium clostridioforme		88.79	0.42	MLT 99			
Parabacteroides distasonis		11.21	0.32	TRE 88			
47655702	**Low discrimination**						**47655702**
Clostridium clostridioforme		49.18	0.40	MLT 99			
Bacteroides uniformis		17.06	0.19	TRE 88			
Parabacteroides distasonis		15.69	0.37				
47655712	**Acceptable identification**						**47655712**
					GRA		
Clostridium clostridioforme		92.06	0.51	MLT 99	50		
Bifidobacterium adolescentis		4.47	0.32	TRE 88	99		
Parabacteroides distasonis		3.47	0.32		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47655722	**Low discrimination**						**47655722**
Bacteroides uniformis		50.86	0.53	IND 98			
Parabacteroides distasonis		46.76	0.71	MLT 92			
Clostridium clostridioforme		1.48	0.40				
47655732	**Low discrimination**						**47655732**
Parabacteroides distasonis		76.52	0.67				
Clostridium clostridioforme		20.50	0.51				
Bacteroides uniformis		1.56	0.19				
47656100	**Acceptable identification**						**47656100**
					GRA		
Fusobacterium mortiferum		92.79	0.35	MAN 1	1		
Actinomyces israelii		7.21	0.19	XYL 7	99		
47657040	**Acceptable identification**						**47657040**
					SPO		
Actinomyces israelii		94.26	0.36	MLT 99	1		
Clostridium innocuum		5.74	0.00		99		
47657042	**Good identification**						**47657042**
					SPO		
Actinomyces israelii		96.10	0.39	MLT 99	1		
Clostridium innocuum		3.90	0.00		99		
47657100	**Low discrimination**						**47657100**
					GRA	SPO	
Actinomyces israelii		76.87	0.53	MLT 99	99	1	
Fusobacterium mortiferum		14.99	0.38		1	1	
Clostridium butyricum		5.80	0.13		99	99	
47657102	**Low discrimination**						**47657102**
Actinomyces israelii		50.84	0.56	MLT 99			
Bifidobacterium adolescentis		44.56	0.49				
Bifidobacterium breve		2.23	0.27				
47657110	**Low discrimination**						**47657110**
Actinomyces israelii		84.41	0.39	MLT 99			
Bifidobacterium adolescentis		9.17	0.16	MLZ 13			
Bifidobacterium breve		3.68	0.10				
47657112	**Low discrimination**						**47657112**
Bifidobacterium adolescentis		83.70	0.50	MLT 99			
Actinomyces israelii		11.68	0.42	ARA 87			
Bifidobacterium breve		3.43	0.27				
47657140	**Good identification**						**47657140**
					SPO		
Actinomyces israelii		96.75	0.46	MLT 99	1		
Clostridium butyricum		1.63	-.05		99		
Bifidobacterium breve		1.62	0.10		1		
47657142	**Low discrimination**						**47657142**
Actinomyces israelii		71.88	0.49	MLT 99			
Bifidobacterium adolescentis		20.02	0.33				
Bifidobacterium breve		8.11	0.27				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
47657302	**Low discrimination**						**47657302**
Actinomyces israelii		89.48	0.37	MLT 99			
Bifidobacterium adolescentis		10.52	0.15	RHA 7			
47657322	**Low discrimination**						**47657322**
					SPO		
Parabacteroides distasonis		78.68	0.41	MAN 1	1		
Clostridium ramosum		7.96	0.11		92		
Prevotella oralis		6.98	0.10		1		
47657332	**Good identification**						**47657332**
Parabacteroides distasonis		100.0	0.36	MAN 1			
47657400	**Low discrimination**						**47657400**
					SPO		
Actinomyces israelii		60.82	0.41	MLT 99	1		
Clostridium innocuum		30.77	0.21		99		
Clostridium butyricum		8.41	0.05		99		
47657402	**Low discrimination**						**47657402**
					SPO		
Actinomyces israelii		60.14	0.44	MLT 99	1		
Clostridium innocuum		20.28	0.21		99		
Bifidobacterium adolescentis		17.42	0.29		1		
47657442	**Acceptable identification**						**47657442**
Actinomyces israelii		91.57	0.37	MLT 99			
Bifidobacterium adolescentis		8.43	0.13				
47657500	**Low discrimination**						**47657500**
					SPO		
Actinomyces israelii		51.60	0.52	MLT 99	1		
Clostridium butyricum		42.80	0.29		99		
Bifidobacterium adolescentis		5.61	0.29		1		
47657502	**Low discrimination**						**47657502**
					SPO		
Bifidobacterium adolescentis		85.44	0.63	MLT 99	1		
Actinomyces israelii		11.92	0.55		1		
Clostridium butyricum		2.56	0.22		99		
47657510	**Low discrimination**						**47657510**
Actinomyces israelii		34.12	0.37	MLT 99			
Bifidobacterium adolescentis		30.31	0.30	MLZ 13			
Bifidobacterium dentium		23.26	0.10				
47657512	**Good identification**						**47657512**
Bifidobacterium adolescentis		91.18	0.64	MLT 99			
Bifidobacterium dentium		7.14	0.27				
Actinomyces israelii		1.56	0.40				
47657540	**Low discrimination**						**47657540**
					COC	SPO	
Actinomyces israelii		47.85	0.45	MLT 99	1	1	
Blautia producta		41.63	0.24		99	1	
Clostridium butyricum		8.87	0.11		1	99	
47657542	**Low discrimination**						**47657542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Bifidobacterium adolescentis		67.19	0.47	MLT 99	1		
Actinomyces israelii		29.49	0.48	SOR 11	1		
Blautia producta		1.90	0.07		99		
47657552	**Low discrimination**						**47657552**
					COC		
Bifidobacterium adolescentis		88.53	0.49	MLT 99	1		
Blautia producta		6.15	0.15	SOR 11	99		
Actinomyces israelii		4.75	0.33		1		
47657702	**Low discrimination**						**47657702**
					SPO		
Actinomyces israelii		47.26	0.35	MLT 99	1		
Bifidobacterium adolescentis		45.46	0.29	RHA 7	1		
Clostridium butyricum		4.18	-.04		99		
47660000	**Good identification**						**47660000**
Fusobacterium mortiferum		100.0	0.39	FRU 99			
47660100	**Good identification**						**47660100**
Fusobacterium mortiferum		100.0	0.46	FRU 99			
47661000	**Low discrimination**						**47661000**
					GRA		
Fusobacterium mortiferum		86.41	0.42	FRU 99	1		
Eubacterium tortuosum		7.15	0.07		99		
Lactobacillus jensenii		4.11	0.10		99		
47661100	**Good identification**						**47661100**
					GRA		
Fusobacterium mortiferum		96.81	0.49	FRU 99	1		
Actinomyces naeslundii		3.19	0.35		99		
47661101	**Low discrimination**						**47661101**
					GRA		
Actinomyces naeslundii		83.01	0.38	ESL 12	99		
Fusobacterium mortiferum		16.99	0.15	MLT 94	1		
47661103	**Good identification**						**47661103**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47661121	**Good identification**						**47661121**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47661123	**Good identification**						**47661123**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47661322	**Low discrimination**						**47661322**
					SPO	GRA	
Parabacteroides distasonis		57.08	0.36	FRU 99	1	1	
Clostridium ramosum		31.56	0.19	XYL 90	92	50	
Lactobacillus acidophilus		6.20	0.08		1	99	
47663001	**Good identification**						**47663001**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47663003	**Good identification**						**47663003**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.35	ESL 12			
47663021	**Good identification**						**47663021**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47663023	**Good identification**						**47663023**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47663100	**Low discrimination**						**47663100**
					GRA		
Actinomyces naeslundii		86.35	0.40	ESL 12	99		
Fusobacterium mortiferum		13.65	0.15	MLT 94	1		
47663101	**Good identification**						**47663101**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47663102	**Low discrimination**						**47663102**
Actinomyces naeslundii		44.84	0.40	ESL 12			
Bifidobacterium breve		39.77	0.24	MLT 94			
Lactobacillus acidophilus		10.90	0.05	FRU 95			
47663103	**Good identification**						**47663103**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47663121	**Good identification**						**47663121**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47663123	**Good identification**						**47663123**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47665122	**Good identification**						**47665122**
Parabacteroides distasonis		100.0	0.44	FRU 99			
47665132	**Good identification**						**47665132**
Parabacteroides distasonis		100.0	0.39	FRU 99			
47665322	**Good identification**						**47665322**
Parabacteroides distasonis		100.0	0.52	FRU 99			
47665332	**Good identification**						**47665332**
Parabacteroides distasonis		98.34	0.47	FRU 99			
Clostridium clostridioforme		1.66	0.11				
47665522	**Low discrimination**						**47665522**
					GRA	SPO	
Parabacteroides distasonis		88.24	0.40	FRU 99	1	1	
Clostridium butyricum		9.92	0.07		99	99	
Clostridium clostridioforme		1.84	0.06		50	66	
47665532	**Low discrimination**						**47665532**
					GRA		
Parabacteroides distasonis		78.55	0.35	FRU 99	1		
Clostridium clostridioforme		13.86	0.17		50		
Bifidobacterium dentium		7.59	-.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47665722	**Acceptable identification**						**47665722**
Parabacteroides distasonis		91.73	0.48	FRU 99			
Clostridium clostridioforme		4.68	0.20				
Bacteroides ovatus		2.63	0.06				
47665732	**Low discrimination**						**47665732**
Parabacteroides distasonis		68.18	0.44	FRU 99			
Clostridium clostridioforme		29.40	0.32				
Bacteroides ovatus		2.42	0.03				
47670000	**Very good identification**						**47670000**
					GRA		
Fusobacterium mortiferum		99.90	0.74		1		
Lactobacillus jensenii		0.07	0.10		99		
Actinomyces naeslundii		0.03	0.25		99		
47670001	**Good identification**						**47670001**
					GRA		
Fusobacterium mortiferum		95.41	0.39	URE 1	1		
Actinomyces naeslundii		4.59	0.28		99		
47670002	**Very good identification**						**47670002**
					GRA		
Fusobacterium mortiferum		98.24	0.57	BGL 10	1		
Lactobacillus acidophilus		0.80	0.09		99		
Lactobacillus jensenii		0.43	0.07		99		
47670010	**Good identification**						**47670010**
					GRA		
Fusobacterium mortiferum		97.68	0.39	MLZ 1	1		
Lactobacillus jensenii		2.32	0.02		99		
47670020	**Acceptable identification**						**47670020**
					GRA	SPO	
Fusobacterium mortiferum		94.33	0.39	NAG 1	1	1	
Actinomyces naeslundii		3.02	0.25		99	1	
Clostridium barati		2.65	-.02		99	99	
47670040	**Good identification**						**47670040**
Fusobacterium mortiferum		100.0	0.39	SOR 1			
47670100	**Very good identification**						**47670100**
					GRA	SPO	
Fusobacterium mortiferum		99.95	0.80		1	1	
Actinomyces naeslundii		0.04	0.33		99	1	
Clostridium barati		0.01	-.02		99	99	
47670101	**Acceptable identification**						**47670101**
					GRA		
Fusobacterium mortiferum		94.46	0.46	URE 1	1		
Actinomyces naeslundii		5.54	0.36		99		
47670102	**Good identification**						**47670102**
					GRA		
Fusobacterium mortiferum		94.72	0.64	BGL 10	1		
Lactobacillus acidophilus		3.81	0.27		99		
Bifidobacterium breve		0.73	0.24		99		
47670110	**Good identification**						**47670110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		98.94	0.46	MLZ 1			
Leptotrichia buccalis		1.06	-.11				
47670120	**Low discrimination**						**47670120**
					GRA	SPO	
Fusobacterium mortiferum		85.53	0.46	NAG 1	1	1	
Clostridium paraputrificum		4.03	0.03		99	99	
Actinomyces naeslundii		3.34	0.33		99	1	
47670121	**Low discrimination**						**47670121**
					GRA		
Actinomyces naeslundii		85.31	0.36	ESL 12	99		
Fusobacterium mortiferum		14.69	0.12	MLT 94	1		
47670123	**Good identification**						**47670123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47670140	**Good identification**						**47670140**
Fusobacterium mortiferum		100.0	0.46	SOR 1			
47670200	**Good identification**						**47670200**
Fusobacterium mortiferum		100.0	0.39	RHA 1			
47670300	**Good identification**						**47670300**
Fusobacterium mortiferum		100.0	0.46	RHA 1			
47670322	**Low discrimination**						**47670322**
					SPO	GRA	
Parabacteroides distasonis		54.31	0.36	GAL 99	1	1	
Clostridium ramosum		30.03	0.19	XYL 90	92	50	
Lactobacillus acidophilus		5.90	0.08		1	99	
47670400	**Good identification**						**47670400**
Fusobacterium mortiferum		100.0	0.39	ARA 1			
47670500	**Acceptable identification**						**47670500**
					GRA	SPO	
Fusobacterium mortiferum		94.52	0.46	ARA 1	1	1	
Clostridium butyricum		5.48	0.06		99	99	
47670712	**Good identification**						**47670712**
Clostridium clostridioforme		100.0	0.41	MLT 99			
47670732	**Acceptable identification**						**47670732**
Clostridium clostridioforme		92.68	0.41	MLT 99			
Parabacteroides distasonis		7.32	0.27	XYL 91			
47671000	**Good identification**						**47671000**
					GRA		
Fusobacterium mortiferum		94.59	0.77		1		
Lactobacillus jensenii		4.50	0.44		99		
Actinomyces naeslundii		0.49	0.49		99		
47671001	**Low discrimination**						**47671001**
					GRA		
Fusobacterium mortiferum		55.05	0.42	URE 1	1		
Actinomyces naeslundii		42.34	0.52		99		
Lactobacillus jensenii		2.62	0.10		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47671002	**Low discrimination**					**47671002**
					GRA	
Fusobacterium mortiferum		51.45	0.60	BGL 10	1	
Lactobacillus acidophilus		27.72	0.43		99	
Lactobacillus jensenii		14.67	0.41		99	
47671003	**Low discrimination**					**47671003**
					GRA	
Actinomyces naeslundii		79.14	0.52	ESL 12	99	
Fusobacterium mortiferum		11.43	0.26	MLT 94	1	
Lactobacillus acidophilus		6.16	0.09		99	
47671010	**Low discrimination**					**47671010**
					GRA	
Lactobacillus jensenii		60.05	0.36	MLT 95	99	
Fusobacterium mortiferum		38.28	0.42	RAF 1	1	
Actinomyces naeslundii		1.03	0.27		99	
47671021	**Good identification**					**47671021**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47671022	**Low discrimination**					**47671022**
					SPO	
Lactobacillus acidophilus		61.23	0.43	MLT 94	1	
Clostridium barati		11.02	0.28	LAC 92	99	
Lactobacillus jensenii		8.10	0.31		1	
47671023	**Acceptable identification**					**47671023**
Actinomyces naeslundii		92.78	0.52	ESL 12		
Lactobacillus acidophilus		7.22	0.09	MLT 94		
47671040	**Low discrimination**					**47671040**
					GRA	
Fusobacterium mortiferum		86.59	0.42	SOR 1	1	
Actinomyces naeslundii		7.83	0.36		99	
Lactobacillus jensenii		4.12	0.10		99	
47671041	**Good identification**					**47671041**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47671043	**Good identification**					**47671043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47671060	**Low discrimination**					**47671060**
					SPO	
Actinomyces naeslundii		62.55	0.36	ESL 12	1	
Clostridium barati		21.38	0.01	MLT 94	99	
Bifidobacterium breve		16.07	0.12		1	
47671061	**Good identification**					**47671061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47671063	**Good identification**					**47671063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47671100	**Very good identification**					**47671100**
					GRA	
Fusobacterium mortiferum		97.49	0.83		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.62	0.57		99		
Bifidobacterium breve		0.55	0.41		99		
47671101	**Low discrimination**						**47671101**
					GRA		
Fusobacterium mortiferum		51.60	0.49	URE 1	1		
Actinomyces naeslundii		48.40	0.60		99		
47671102	**Low discrimination**						**47671102**
					GRA		
Lactobacillus acidophilus		60.11	0.61	MLT 94	99		
Fusobacterium mortiferum		22.64	0.67		1		
Bifidobacterium breve		11.55	0.59		99		
47671103	**Low discrimination**						**47671103**
					GRA		
Actinomyces naeslundii		66.62	0.60	ESL 12	99		
Lactobacillus acidophilus		20.95	0.27	MLT 94	99		
Fusobacterium mortiferum		7.89	0.32		1		
47671110	**Low discrimination**						**47671110**
					GRA		
Fusobacterium mortiferum		42.80	0.49	MLZ 1	1		
Leptotrichia buccalis		30.32	0.24		1		
Bifidobacterium breve		23.75	0.41		99		
47671111	**Low discrimination**						**47671111**
					GRA		
Actinomyces naeslundii		74.10	0.38	ESL 12	99		
Fusobacterium mortiferum		15.16	0.15	MLT 94	1		
Leptotrichia buccalis		10.74	-.11	MLZ 5	1		
47671112	**Good identification**						**47671112**
					GRA		
Leptotrichia buccalis		91.65	0.58	MLT 99	1		
Bifidobacterium breve		7.33	0.59		99		
Lactobacillus acidophilus		0.39	0.27		99		
47671120	**Low discrimination**						**47671120**
					SPO	GRA	
Clostridium paraputrificum		34.34	0.37	MLT 99	99	99	
Clostridium ramosum		23.29	0.41	RAF 7	92	50	
Fusobacterium mortiferum		11.04	0.49		1	1	
47671121	**Good identification**						**47671121**
					SPO		
Actinomyces naeslundii		92.80	0.60	ESL 12	1		
Clostridium paraputrificum		3.11	0.03	MLT 94	99		
Clostridium ramosum		2.11	0.06		92		
47671122	**Low discrimination**						**47671122**
					SPO		
Lactobacillus acidophilus		46.94	0.61	MLT 94	1		
Bifidobacterium breve		16.03	0.63		1		
Clostridium ramosum		13.56	0.51		92		
47671123	**Low discrimination**						**47671123**
Actinomyces naeslundii		60.22	0.60	ESL 12			
Lactobacillus acidophilus		18.94	0.27	MLT 94			
Bifidobacterium breve		6.47	0.29				
47671130	**Low discrimination**						**47671130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bifidobacterium breve		82.37	0.46	MLT 99	99	
Leptotrichia buccalis		6.57	0.07	BGL 91	1	
Parabacteroides distasonis		3.06	0.23		1	
47671131	**Low discrimination**					**47671131**
Actinomyces naeslundii		83.21	0.38	ESL 12		
Bifidobacterium breve		16.79	0.12	MLT 94		
47671132	**Low discrimination**					**47671132**
					GRA	
Bifidobacterium breve		45.53	0.63	MLT 99	99	
Leptotrichia buccalis		35.58	0.41		1	
Parabacteroides distasonis		16.58	0.57		1	
47671140	**Low discrimination**					**47671140**
					GRA	
Fusobacterium mortiferum		56.15	0.49	SOR 1	1	
Bifidobacterium breve		31.16	0.41		99	
Actinomyces naeslundii		6.20	0.44		99	
47671141	**Acceptable identification**					**47671141**
					GRA	
Actinomyces naeslundii		94.25	0.47	ESL 12	99	
Fusobacterium mortiferum		5.75	0.15	MLT 94	1	
47671142	**Low discrimination**					**47671142**
					GRA	
Bifidobacterium breve		58.61	0.59	MLT 99	99	
Lactobacillus acidophilus		37.72	0.45		99	
Fusobacterium mortiferum		1.16	0.32		1	
47671143	**Low discrimination**					**47671143**
Actinomyces naeslundii		64.00	0.47	ESL 12		
Bifidobacterium breve		21.91	0.24	MLT 94		
Lactobacillus acidophilus		14.10	0.11			
47671150	**Good identification**					**47671150**
					GRA	
Bifidobacterium breve		96.98	0.41	MLT 99	99	
Fusobacterium mortiferum		1.77	0.15	BGL 91	1	
Leptotrichia buccalis		1.25	-.11		1	
47671152	**Low discrimination**					**47671152**
					GRA	
Bifidobacterium breve		87.76	0.59	MLT 99	99	
Leptotrichia buccalis		11.08	0.24		1	
Lactobacillus acidophilus		0.57	0.11		99	
47671160	**Low discrimination**					**47671160**
					SPO	
Bifidobacterium breve		68.57	0.46	MLT 99	1	
Clostridium paraputrificum		16.28	0.18	BGL 91	99	
Actinomyces naeslundii		7.67	0.44		1	
47671161	**Acceptable identification**					**47671161**
Actinomyces naeslundii		94.32	0.47	ESL 12		
Bifidobacterium breve		5.68	0.12	MLT 94		
47671162	**Low discrimination**					**47671162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium breve		70.68	0.63	MLT 99	1		
Lactobacillus acidophilus		25.59	0.45		1		
Clostridium barati		0.85	0.18		99		
47671163	**Low discrimination**						**47671163**
Actinomyces naeslundii		54.68	0.47	ESL 12			
Bifidobacterium breve		33.28	0.29	MLT 94			
Lactobacillus acidophilus		12.05	0.11				
47671170	**Good identification**						**47671170**
Bifidobacterium breve		100.0	0.46	MLT 99			
47671172	**Very good identification**						**47671172**
					GRA		
Bifidobacterium breve		98.50	0.63	MLT 99	99		
Leptotrichia buccalis		0.78	0.07		1		
Parabacteroides distasonis		0.36	0.23		1		
47671200	**Good identification**						**47671200**
					GRA		
Fusobacterium mortiferum		95.46	0.42	RHA 1	1		
Lactobacillus jensenii		4.54	0.10		99		
47671222	**Low discrimination**						**47671222**
					GRA		
Parabacteroides distasonis		30.24	0.36	XYL 90	1		
Lactobacillus acidophilus		28.28	0.24	LAC 99	99		
Prevotella oralis		17.30	0.19		1		
47671300	**Acceptable identification**						**47671300**
					SPO	GRA	
Fusobacterium mortiferum		92.32	0.49	RHA 1	1	1	
Clostridium ramosum		2.61	0.09		92	50	
Lactobacillus acidophilus		2.05	0.08		1	99	
47671302	**Low discrimination**						**47671302**
					SPO	GRA	
Lactobacillus acidophilus		76.40	0.42	MLT 94	1	99	
Parabacteroides distasonis		7.10	0.36	RHA 7	1	1	
Clostridium ramosum		3.93	0.19		92	50	
47671320	**Low discrimination**						**47671320**
					SPO		
Clostridium ramosum		86.93	0.43	MLT 99	92		
Parabacteroides distasonis		6.35	0.36		1		
Prevotella oralis		3.00	0.17		1		
47671322	**Low discrimination**						**47671322**
					SPO	GRA	
Parabacteroides distasonis		56.64	0.70	XYL 90	1	1	
Clostridium ramosum		31.31	0.53		92	50	
Lactobacillus acidophilus		6.15	0.42		1	99	
47671323	**Low discrimination**						**47671323**
					SPO	GRA	
Parabacteroides distasonis		55.61	0.36	XYL 90	1	1	
Clostridium ramosum		30.74	0.19	URE 1	92	50	
Lactobacillus acidophilus		6.04	0.08		1	99	
47671332	**Very good identification**						**47671332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Parabacteroides distasonis		95.64	0.65	XYL 90	1		
Clostridium clostridioforme		1.44	0.28		66		
Clostridium ramosum		0.99	0.19		92		
47671400	**Low discrimination**						**47671400**
					GRA	SPO	
Fusobacterium mortiferum		67.75	0.42	ARA 1	1	1	
Clostridium butyricum		25.19	0.16		99	99	
Lactobacillus jensenii		3.22	0.10		99	1	
47671500	**Low discrimination**						**47671500**
					GRA	SPO	
Clostridium butyricum		77.96	0.40	MLT 99	99	99	
Fusobacterium mortiferum		20.39	0.49	XYL 97	1	1	
Actinomyces israelii		1.24	0.30		99	1	
47671512	**Low discrimination**						**47671512**
					GRA		
Bifidobacterium adolescentis		57.70	0.49	MLT 99	99		
Leptotrichia buccalis		24.16	0.24	XYL 87	1		
Clostridium clostridioforme		11.92	0.34		50		
47671522	**Low discrimination**						**47671522**
					GRA	SPO	
Parabacteroides distasonis		84.16	0.58	XYL 90	1	1	
Lactobacillus acidophilus		6.01	0.27		99	1	
Clostridium butyricum		2.63	0.15		99	99	
47671532	**Low discrimination**						**47671532**
					GRA		
Parabacteroides distasonis		79.54	0.53	XYL 90	1		
Clostridium clostridioforme		12.49	0.34		50		
Bifidobacterium breve		3.60	0.29		99		
47671612	**Good identification**						**47671612**
Clostridium clostridioforme		100.0	0.40	MLT 99			
47671632	**Acceptable identification**						**47671632**
Clostridium clostridioforme		91.90	0.40	MLT 99			
Parabacteroides distasonis		8.10	0.27	XYL 91			
47671702	**Low discrimination**						**47671702**
					GRA	SPO	
Clostridium clostridioforme		72.42	0.37	MLT 99	50	66	
Parabacteroides distasonis		16.12	0.32	XYL 91	1	1	
Clostridium butyricum		5.92	0.07		99	99	
47671712	**Good identification**						**47671712**
					GRA		
Clostridium clostridioforme		96.55	0.49	MLT 99	50		
Parabacteroides distasonis		2.54	0.27	XYL 91	1		
Bifidobacterium adolescentis		0.64	0.15		99		
47671722	**Good identification**						**47671722**
					SPO		
Parabacteroides distasonis		93.96	0.66	XYL 90	1		
Clostridium clostridioforme		4.26	0.37		66		
Clostridium ramosum		0.86	0.19		92		
47671732	**Low discrimination**						**47671732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		71.94	0.61	XYL 90			
Clostridium clostridioforme		27.62	0.49				
Bacteroides ovatus		0.23	0.03				
47672000	**Low discrimination**						**47672000**
					GRA	SPO	
Fusobacterium mortiferum		50.15	0.39	MAN 1	1	1	
Clostridium innocuum		46.73	0.26		99	99	
Actinomyces naeslundii		3.12	0.30		99	1	
47672100	**Acceptable identification**						**47672100**
					GRA		
Fusobacterium mortiferum		92.95	0.46	MAN 1	1		
Actinomyces naeslundii		7.05	0.38		99		
47672101	**Acceptable identification**						**47672101**
					GRA		
Actinomyces naeslundii		91.85	0.41	ESL 12	99		
Fusobacterium mortiferum		8.15	0.12	MLT 94	1		
47672103	**Good identification**						**47672103**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47672121	**Good identification**						**47672121**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47672123	**Good identification**						**47672123**
Actinomyces naeslundii		100.0	0.41	ESL 12			
47673000	**Low discrimination**						**47673000**
					GRA	SPO	
Clostridium innocuum		83.76	0.48	RAF 1	99	99	
Fusobacterium mortiferum		7.10	0.42		1	1	
Actinomyces naeslundii		7.06	0.54		99	1	
47673001	**Good identification**						**47673001**
					SPO		
Actinomyces naeslundii		92.61	0.57	ESL 12	1		
Clostridium innocuum		7.40	0.14	MLT 94	99		
47673002	**Low discrimination**						**47673002**
					SPO		
Clostridium innocuum		85.62	0.48	RAF 1	99		
Actinomyces naeslundii		7.22	0.54		1		
Actinomyces israelii		2.68	0.37		1		
47673003	**Good identification**						**47673003**
					SPO		
Actinomyces naeslundii		92.61	0.57	ESL 12	1		
Clostridium innocuum		7.40	0.14	MLT 94	99		
47673011	**Good identification**						**47673011**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47673013	**Good identification**						**47673013**
Actinomyces naeslundii		100.0	0.35	ESL 12			
47673020	**Low discrimination**						**47673020**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		67.88	0.54	ESL 12	1	
Clostridium ramosum		19.22	0.19	MLT 94	92	
Clostridium innocuum		8.13	0.14		99	
47673021	**Very good identification**					**47673021**
Actinomyces naeslundii		100.0	0.57	ESL 12		
47673023	**Very good identification**					**47673023**
Actinomyces naeslundii		100.0	0.57	ESL 12		
47673031	**Good identification**					**47673031**
Actinomyces naeslundii		100.0	0.35	ESL 12		
47673033	**Good identification**					**47673033**
Actinomyces naeslundii		100.0	0.35	ESL 12		
47673040	**Low discrimination**					**47673040**
					SPO	
Actinomyces naeslundii		44.95	0.41	ESL 12	1	
Clostridium innocuum		30.51	0.14	MLT 94	99	
Actinomyces israelii		24.54	0.27		1	
47673041	**Good identification**					**47673041**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47673042	**Low discrimination**					**47673042**
Actinomyces naeslundii		28.82	0.41	ESL 12		
Actinomyces israelii		23.60	0.30	MLT 94		
Bifidobacterium breve		21.70	0.24			
47673043	**Good identification**					**47673043**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47673060	**Low discrimination**					**47673060**
Actinomyces naeslundii		88.31	0.41	ESL 12		
Bifidobacterium breve		11.69	0.12	MLT 94		
47673061	**Good identification**					**47673061**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47673063	**Good identification**					**47673063**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47673100	**Low discrimination**					**47673100**
					GRA	
Actinomyces naeslundii		35.06	0.62	ESL 12	99	
Fusobacterium mortiferum		28.88	0.49	MLT 94	1	
Bifidobacterium breve		16.03	0.41		99	
47673101	**Very good identification**					**47673101**
					GRA	
Actinomyces naeslundii		99.45	0.65	ESL 12	99	
Fusobacterium mortiferum		0.55	0.15	MLT 94	1	
47673102	**Low discrimination**					**47673102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		56.42	0.59	MLT 99		
Lactobacillus acidophilus		15.46	0.39			
Actinomyces naeslundii		12.21	0.62			
47673103	**Very good identification**					**47673103**
Actinomyces naeslundii		95.86	0.65	ESL 12		
Bifidobacterium breve		2.98	0.24	MLT 94		
Lactobacillus acidophilus		0.82	0.05			
47673110	**Low discrimination**					**47673110**
Bifidobacterium breve		74.97	0.41	MLT 99		
Actinomyces naeslundii		8.63	0.40	BGL 91		
Actinomyces israelii		8.54	0.31			
47673111	**Good identification**					**47673111**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47673112	**Low discrimination**					**47673112**
					GRA	
Bifidobacterium breve		82.10	0.59	MLT 99	99	
Leptotrichia buccalis		10.37	0.24		1	
Bifidobacterium adolescentis		3.03	0.33		99	
47673113	**Low discrimination**					**47673113**
					GRA	
Actinomyces naeslundii		60.03	0.43	ESL 12	99	
Bifidobacterium breve		35.49	0.24	MLT 94	99	
Leptotrichia buccalis		4.48	-.11	MLZ 5	1	
47673120	**Low discrimination**					**47673120**
					SPO	
Clostridium ramosum		83.32	0.53	MLT 99	92	
Actinomyces naeslundii		8.46	0.62		1	
Bifidobacterium breve		6.87	0.46		1	
47673121	**Good identification**					**47673121**
					SPO	
Actinomyces naeslundii		93.30	0.65	ESL 12	1	
Clostridium ramosum		6.19	0.19	MLT 94	92	
Bifidobacterium breve		0.51	0.12		1	
47673122	**Low discrimination**					**47673122**
					SPO	
Clostridium ramosum		78.35	0.64	MLT 99	92	
Bifidobacterium breve		16.34	0.63		1	
Lactobacillus acidophilus		2.52	0.39		1	
47673123	**Low discrimination**					**47673123**
					SPO	
Actinomyces naeslundii		75.24	0.65	ESL 12	1	
Clostridium ramosum		19.96	0.30	MLT 94	92	
Bifidobacterium breve		4.16	0.29		1	
47673130	**Low discrimination**					**47673130**
					SPO	
Bifidobacterium breve		76.25	0.46	MLT 99	1	
Clostridium tertium		9.48	0.21	BGL 91	99	
Clostridium ramosum		9.34	0.19		92	
47673131	**Acceptable identification**					**47673131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		90.58	0.43	ESL 12		
Bifidobacterium breve		9.42	0.12	MLT 94		
47673132	**Good identification**					**47673132**
					SPO	
Bifidobacterium breve		90.30	0.63	MLT 99	1	
Clostridium ramosum		4.37	0.30		92	
Clostridium tertium		3.33	0.29		99	
47673140	**Low discrimination**					**47673140**
Bifidobacterium breve		58.80	0.41	MLT 99		
Actinomyces naeslundii		22.70	0.49	BGL 91		
Actinomyces israelii		17.42	0.38			
47673141	**Good identification**					**47673141**
Actinomyces naeslundii		100.0	0.52	ESL 12		
47673142	**Low discrimination**					**47673142**
Bifidobacterium breve		89.31	0.59	MLT 99		
Actinomyces israelii		3.93	0.41			
Actinomyces naeslundii		3.41	0.49			
47673143	**Low discrimination**					**47673143**
Actinomyces naeslundii		85.01	0.52	ESL 12		
Bifidobacterium breve		14.99	0.24	MLT 94		
47673150	**Acceptable identification**					**47673150**
Bifidobacterium breve		92.24	0.41	MLT 99		
Actinomyces israelii		4.08	0.24	BGL 91		
Actinomyces naeslundii		1.88	0.27			
47673152	**Very good identification**					**47673152**
Bifidobacterium breve		98.58	0.59	MLT 99		
Actinomyces israelii		0.65	0.27			
Bifidobacterium adolescentis		0.45	0.18			
47673160	**Low discrimination**					**47673160**
					SPO	
Bifidobacterium breve		74.06	0.46	MLT 99	1	
Actinomyces naeslundii		16.09	0.49	BGL 91	1	
Clostridium ramosum		9.07	0.19		92	
47673161	**Good identification**					**47673161**
Actinomyces naeslundii		96.99	0.52	ESL 12		
Bifidobacterium breve		3.01	0.12	MLT 94		
47673162	**Good identification**					**47673162**
					SPO	
Bifidobacterium breve		91.61	0.63	MLT 99	1	
Clostridium ramosum		4.44	0.30		92	
Actinomyces naeslundii		1.97	0.49		1	
47673163	**Low discrimination**					**47673163**
Actinomyces naeslundii		76.13	0.52	ESL 12		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		23.87	0.29	MLT 94			
47673170	**Good identification**						**47673170**
Bifidobacterium breve		98.87	0.46	MLT 99			
Actinomyces naeslundii		1.13	0.27	BGL 91			
47673172	**Very good identification**						**47673172**
Bifidobacterium breve		99.89	0.63	MLT 99			
Actinomyces naeslundii		0.11	0.27				
47673320	**Good identification**						**47673320**
					SPO		
Clostridium ramosum		99.86	0.56	MLT 99	92		
Actinomyces naeslundii		0.08	0.28		1		
Bifidobacterium breve		0.06	0.12		1		
47673322	**Very good identification**						**47673322**
					SPO	GRA	
Clostridium ramosum		99.28	0.66	MLT 99	92	50	
Parabacteroides distasonis		0.32	0.36		1	1	
Lactobacillus acidophilus		0.18	0.20		1	99	
47673400	**Low discrimination**						**47673400**
					SPO		
Clostridium innocuum		89.55	0.35	RAF 1	99		
Actinomyces israelii		8.67	0.33		1		
Clostridium butyricum		1.78	0.00		99		
47673402	**Low discrimination**						**47673402**
					SPO		
Clostridium innocuum		85.78	0.35	RAF 1	99		
Actinomyces israelii		12.46	0.36		1		
Bifidobacterium adolescentis		1.10	0.12		1		
47673502	**Low discrimination**						**47673502**
					SPO		
Bifidobacterium adolescentis		58.95	0.46	MLT 99	1		
Actinomyces israelii		27.01	0.46	XYL 87	1		
Clostridium butyricum		8.60	0.17		99		
47673512	**Low discrimination**						**47673512**
Bifidobacterium adolescentis		52.77	0.47	MLT 99			
Bifidobacterium dentium		41.51	0.27	XYL 87			
Actinomyces israelii		2.96	0.32				
47673542	**Low discrimination**						**47673542**
Actinomyces israelii		44.59	0.39	MLT 99			
Bifidobacterium adolescentis		30.93	0.30	XYL 92			
Bifidobacterium breve		12.52	0.24				
47674000	**Good identification**						**47674000**
Fusobacterium mortiferum		100.0	0.54	XYL 7			
47674002	**Good identification**						**47674002**
Fusobacterium mortiferum		100.0	0.38	XYL 7			
47674100	**Very good identification**						**47674100**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		97.12	0.61	XYL 7	1	1	
Clostridium butyricum		2.71	0.15		99	99	
Actinomyces israelii		0.17	0.16		99	1	
47674102	**Low discrimination**						**47674102**
					GRA	SPO	
Fusobacterium mortiferum		84.19	0.44	XYL 7	1	1	
Clostridium butyricum		8.23	0.08	BGL 10	99	99	
Clostridium clostridioforme		2.52	0.11		50	66	
47674122	**Good identification**						**47674122**
Parabacteroides distasonis		97.38	0.44	GAL 99			
Clostridium clostridioforme		2.62	0.11				
47674132	**Low discrimination**						**47674132**
Parabacteroides distasonis		81.45	0.39	GAL 99			
Clostridium clostridioforme		18.55	0.22				
47674312	**Good identification**						**47674312**
Clostridium clostridioforme		98.22	0.37	MLT 99			
Parabacteroides distasonis		1.78	0.13	ARA 94			
47674322	**Acceptable identification**						**47674322**
Parabacteroides distasonis		93.83	0.52	GAL 99			
Clostridium clostridioforme		6.18	0.26				
47674332	**Low discrimination**						**47674332**
Parabacteroides distasonis		64.24	0.47	GAL 99			
Clostridium clostridioforme		35.77	0.37				
47674512	**Low discrimination**						**47674512**
					SPO		
Clostridium clostridioforme		88.65	0.43	MLT 99	66		
Bifidobacterium adolescentis		8.60	0.29	RHA 88	1		
Clostridium butyricum		1.92	0.03		99		
47674522	**Low discrimination**						**47674522**
					GRA	SPO	
Parabacteroides distasonis		55.54	0.40	GAL 99	1	1	
Clostridium clostridioforme		38.22	0.32		50	66	
Clostridium butyricum		6.24	0.07		99	99	
47674532	**Low discrimination**						**47674532**
					GRA		
Clostridium clostridioforme		84.15	0.43	MLT 99	50		
Parabacteroides distasonis		14.45	0.35	RHA 88	1		
Bifidobacterium dentium		1.40	-.03		99		
47674602	**Good identification**						**47674602**
Clostridium clostridioforme		100.0	0.37	MLT 99			
47674612	**Good identification**						**47674612**
Clostridium clostridioforme		100.0	0.49	MLT 99			
47674622	**Good identification**						**47674622**
Clostridium clostridioforme		98.03	0.37	MLT 99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		1.97	0.14				
47674632	**Good identification**						**47674632**
Clostridium clostridioforme		99.76	0.49	MLT 99			
Parabacteroides distasonis		0.24	0.09				
47674702	**Good identification**						**47674702**
					GRA	SPO	
Clostridium clostridioforme		98.63	0.46	MLT 99	50	66	
Clostridium butyricum		0.78	-.01		99	99	
Parabacteroides distasonis		0.59	0.14		1	1	
47674712	**Very good identification**						**47674712**
Clostridium clostridioforme		99.93	0.58	MLT 99			
Parabacteroides distasonis		0.07	0.09				
47674722	**Low discrimination**						**47674722**
Clostridium clostridioforme		61.81	0.46	MLT 99			
Parabacteroides distasonis		36.75	0.48				
Bacteroides ovatus		1.05	0.06				
47674732	**Good identification**						**47674732**
Clostridium clostridioforme		93.22	0.58	MLT 99			
Parabacteroides distasonis		6.55	0.44				
Bacteroides ovatus		0.23	0.03				
47675000	**Low discrimination**						**47675000**
					GRA	SPO	
Fusobacterium mortiferum		79.82	0.57	XYL 7	1	1	
Clostridium butyricum		14.31	0.26		99	99	
Actinomyces israelii		5.36	0.40		99	1	
47675002	**Low discrimination**						**47675002**
					GRA	SPO	
Fusobacterium mortiferum		36.45	0.41	XYL 7	1	1	
Actinomyces israelii		33.05	0.43	BGL 10	99	1	
Clostridium butyricum		22.87	0.19		99	99	
47675022	**Low discrimination**						**47675022**
					GRA		
Parabacteroides distasonis		84.32	0.44	LAC 99	1		
Prevotella oralis		4.64	0.09		1		
Lactobacillus acidophilus		3.53	0.09		99		
47675032	**Low discrimination**						**47675032**
Parabacteroides distasonis		83.05	0.39	LAC 99			
Clostridium clostridioforme		16.95	0.22				
47675042	**Low discrimination**						**47675042**
					SPO		
Actinomyces israelii		86.60	0.36	MLT 99	1		
Clostridium butyricum		13.40	0.01		99		
47675100	**Low discrimination**						**47675100**
					GRA	SPO	
Clostridium butyricum		60.93	0.50	MLT 99	99	99	
Fusobacterium mortiferum		33.05	0.64	ARA 90	1	1	
Actinomyces israelii		3.81	0.50		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47675102	**Low discrimination**						**47675102**
					SPO		
Clostridium butyricum		53.53	0.43	MLT 99	99		
Bifidobacterium adolescentis		13.87	0.47	ARA 90	1		
Actinomyces israelii		12.89	0.53		1		
47675112	**Low discrimination**						**47675112**
					GRA		
Bifidobacterium adolescentis		48.26	0.49	MLT 99	99		
Leptotrichia buccalis		20.21	0.24	ARA 87	1		
Parabacteroides distasonis		9.42	0.39		1		
47675122	**Very good identification**						**47675122**
					GRA	SPO	
Parabacteroides distasonis		97.26	0.78		1	1	
Prevotella oralis		0.85	0.30		1	1	
Clostridium butyricum		0.74	0.24		99	99	
47675123	**Good identification**						**47675123**
					GRA		
Parabacteroides distasonis		98.48	0.44	URE 1	1		
Actinomyces naeslundii		1.52	0.26		99		
47675130	**Low discrimination**						**47675130**
					GRA	SPO	
Parabacteroides distasonis		80.30	0.39	BGL 99	1	1	
Clostridium butyricum		15.29	0.10		99	99	
Bifidobacterium breve		2.42	0.12		99	1	
47675132	**Very good identification**						**47675132**
					GRA		
Parabacteroides distasonis		98.56	0.73		1		
Clostridium clostridioforme		0.68	0.31		50		
Bifidobacterium breve		0.30	0.29		99		
47675133	**Good identification**						**47675133**
Parabacteroides distasonis		100.0	0.39	URE 1			
47675142	**Low discrimination**						**47675142**
					SPO		
Actinomyces israelii		40.24	0.46	MLT 99	1		
Clostridium butyricum		37.35	0.24		99		
Bifidobacterium adolescentis		13.76	0.32		1		
47675162	**Low discrimination**						**47675162**
					GRA	SPO	
Parabacteroides distasonis		74.53	0.44	SOR 1	1	1	
Bifidobacterium breve		12.25	0.29		99	1	
Clostridium butyricum		4.91	0.06		99	99	
47675172	**Low discrimination**						**47675172**
					GRA		
Parabacteroides distasonis		76.62	0.39	SOR 1	1		
Bifidobacterium breve		23.38	0.29		99		
47675212	**Good identification**						**47675212**
Clostridium clostridioforme		98.02	0.36	MLT 99			
Parabacteroides distasonis		1.98	0.13	ARA 94			
47675222	**Acceptable identification**						**47675222**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		93.74	0.52	LAC 99			
Clostridium clostridioforme		5.53	0.25				
Prevotella oralis		0.74	0.03				
47675232	**Low discrimination**						**47675232**
Parabacteroides distasonis		66.71	0.47	LAC 99			
Clostridium clostridioforme		33.29	0.36				
47675302	**Low discrimination**						**47675302**
					GRA	SPO	
Parabacteroides distasonis		72.96	0.52	NAG 99	1	1	
Clostridium clostridioforme		14.41	0.34		50	66	
Clostridium butyricum		6.55	0.17		99	99	
47675312	**Low discrimination**						**47675312**
					GRA		
Clostridium clostridioforme		61.45	0.45	MLT 99	50		
Parabacteroides distasonis		36.79	0.47	ARA 94	1		
Bifidobacterium adolescentis		0.64	0.15		99		
47675320	**Good identification**						**47675320**
					SPO	GRA	
Parabacteroides distasonis		95.98	0.52	BGL 99	1	1	
Clostridium butyricum		1.93	0.06		99	99	
Clostridium ramosum		1.47	0.09		92	50	
47675322	**Very good identification**						**47675322**
Parabacteroides distasonis		99.52	0.86				
Clostridium clostridioforme		0.20	0.34				
Prevotella oralis		0.12	0.24				
47675323	**Good identification**						**47675323**
Parabacteroides distasonis		100.0	0.52	URE 1			
47675330	**Good identification**						**47675330**
Parabacteroides distasonis		98.34	0.47	BGL 99			
Clostridium clostridioforme		1.66	0.11				
47675332	**Very good identification**						**47675332**
Parabacteroides distasonis		98.28	0.81				
Clostridium clostridioforme		1.66	0.45				
Bacteroides ovatus		0.07	0.11				
47675333	**Good identification**						**47675333**
Parabacteroides distasonis		98.34	0.47	URE 1			
Clostridium clostridioforme		1.66	0.11				
47675362	**Good identification**						**47675362**
Parabacteroides distasonis		100.0	0.52	SOR 1			
47675372	**Good identification**						**47675372**
Parabacteroides distasonis		98.34	0.47	SOR 1			
Clostridium clostridioforme		1.66	0.11				
47675400	**Good identification**						**47675400**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		96.12	0.42	MLT 99	99	99	
Actinomyces israelii		3.27	0.38	LAC 96	99	1	
Fusobacterium mortiferum		0.60	0.23		1	1	
47675402	**Low discrimination**						**47675402**
					SPO		
Clostridium butyricum		77.29	0.35	MLT 99	99		
Actinomyces israelii		10.16	0.41	LAC 96	1		
Clostridium clostridioforme		8.94	0.31		66		
47675412	**Low discrimination**						**47675412**
					SPO		
Clostridium clostridioforme		79.63	0.42	MLT 99	66		
Bifidobacterium adolescentis		8.63	0.29	RHA 88	1		
Clostridium butyricum		7.95	0.13		99		
47675422	**Low discrimination**						**47675422**
					GRA	SPO	
Parabacteroides distasonis		47.06	0.40	LAC 99	1	1	
Clostridium clostridioforme		29.02	0.31		50	66	
Clostridium butyricum		21.81	0.17		99	99	
47675432	**Low discrimination**						**47675432**
					GRA		
Clostridium clostridioforme		82.04	0.42	MLT 99	50		
Parabacteroides distasonis		15.73	0.35	RHA 88	1		
Bifidobacterium dentium		1.52	-.03		99		
47675500	**Very good identification**						**47675500**
					SPO		
Clostridium butyricum		99.22	0.66	MLT 99	99		
Actinomyces israelii		0.56	0.48		1		
Bifidobacterium adolescentis		0.08	0.27		1		
47675502	**Low discrimination**						**47675502**
					SPO		
Clostridium butyricum		80.70	0.59	MLT 99	99		
Bifidobacterium adolescentis		15.55	0.61		1		
Actinomyces israelii		1.77	0.52		1		
47675510	**Acceptable identification**						**47675510**
					SPO		
Clostridium butyricum		94.15	0.44	MLT 99	99		
Bifidobacterium adolescentis		1.66	0.29	MLZ 5	1		
Bifidobacterium dentium		1.56	0.10		1		
47675512	**Low discrimination**						**47675512**
					SPO		
Bifidobacterium adolescentis		60.09	0.63	MLT 99	1		
Clostridium clostridioforme		18.76	0.51		66		
Clostridium butyricum		13.43	0.37		99		
47675520	**Good identification**						**47675520**
					GRA	SPO	
Clostridium butyricum		95.89	0.48	MLT 99	99	99	
Parabacteroides distasonis		3.35	0.40	NAG 8	1	1	
Actinomyces israelii		0.40	0.28		99	1	
47675522	**Low discrimination**						**47675522**
					GRA	SPO	
Parabacteroides distasonis		86.94	0.74		1	1	
Clostridium butyricum		9.77	0.41		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		1.81	0.40		50	66	
47675523	**Low discrimination**						**47675523**
					GRA	SPO	
Parabacteroides distasonis		88.24	0.40	URE 1	1	1	
Clostridium butyricum		9.92	0.07		99	99	
Clostridium clostridioforme		1.84	0.06		50	66	
47675532	**Low discrimination**						**47675532**
					GRA		
Parabacteroides distasonis		77.07	0.70		1		
Clostridium clostridioforme		13.60	0.51		50		
Bifidobacterium dentium		7.45	0.32		99		
47675533	**Low discrimination**						**47675533**
					GRA		
Parabacteroides distasonis		78.55	0.35	URE 1	1		
Clostridium clostridioforme		13.86	0.17		50		
Bifidobacterium dentium		7.59	-.03		99		
47675540	**Acceptable identification**						**47675540**
					COC	SPO	
Clostridium butyricum		93.60	0.48	MLT 99	1	99	
Blautia producta		3.12	0.24	SOR 8	99	1	
Actinomyces israelii		2.38	0.41		1	1	
47675542	**Low discrimination**						**47675542**
					SPO		
Clostridium butyricum		72.17	0.41	MLT 99	99		
Bifidobacterium adolescentis		19.76	0.46	SOR 8	1		
Actinomyces israelii		7.07	0.44		1		
47675552	**Low discrimination**						**47675552**
					COC	SPO	
Bifidobacterium adolescentis		75.78	0.47	MLT 99	1	1	
Clostridium butyricum		11.92	0.19	SOR 11	1	99	
Blautia producta		6.46	0.15		99	1	
47675562	**Low discrimination**						**47675562**
					GRA	SPO	
Parabacteroides distasonis		47.46	0.40	SOR 1	1	1	
Clostridium butyricum		45.91	0.23		99	99	
Actinomyces israelii		3.30	0.24		99	1	
47675572	**Low discrimination**						**47675572**
					GRA		
Parabacteroides distasonis		66.54	0.35	SOR 1	1		
Clostridium clostridioforme		11.74	0.17		50		
Bifidobacterium dentium		6.43	-.03		99		
47675602	**Acceptable identification**						**47675602**
					SPO		
Clostridium clostridioforme		94.80	0.46	MLT 99	66		
Clostridium butyricum		3.46	0.09		99		
Actinomyces israelii		1.11	0.22		1		
47675612	**Very good identification**						**47675612**
Clostridium clostridioforme		99.92	0.57	MLT 99			
Parabacteroides distasonis		0.08	0.09				
47675622	**Low discrimination**						**47675622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		58.94	0.46	MLT 99			
Parabacteroides distasonis		39.10	0.48				
Bacteroides ovatus		1.12	0.06				
47675632	**Good identification**						**47675632**
Clostridium clostridioforme		92.39	0.57	MLT 99			
Parabacteroides distasonis		7.24	0.44				
Bacteroides ovatus		0.26	0.03				
47675700	**Acceptable identification**						**47675700**
					GRA	SPO	
Clostridium butyricum		92.86	0.40	MLT 99	99	99	
Mitsuokella multacida		4.19	0.25	RHA 3	1	1	
Clostridium clostridioforme		1.40	0.20		50	66	
47675702	**Low discrimination**						**47675702**
					GRA	SPO	
Clostridium clostridioforme		66.62	0.55	MLT 99	50	66	
Clostridium butyricum		17.41	0.33		99	99	
Parabacteroides distasonis		13.20	0.48		1	1	
47675712	**Very good identification**						**47675712**
					GRA		
Clostridium clostridioforme		96.82	0.66	MLT 99	50		
Parabacteroides distasonis		2.27	0.44		1		
Bifidobacterium adolescentis		0.43	0.29		99		
47675720	**Low discrimination**						**47675720**
					GRA	SPO	
Parabacteroides distasonis		66.97	0.48	BGL 99	1	1	
Clostridium butyricum		19.75	0.22		99	99	
Bacteroides vulgatus		4.67	0.10		1	1	
47675722	**Good identification**						**47675722**
Parabacteroides distasonis		91.43	0.82				
Clostridium clostridioforme		4.66	0.55				
Bacteroides ovatus		2.62	0.40				
47675723	**Acceptable identification**						**47675723**
Parabacteroides distasonis		91.73	0.48	URE 1			
Clostridium clostridioforme		4.68	0.20				
Bacteroides ovatus		2.63	0.06				
47675730	**Low discrimination**						**47675730**
Parabacteroides distasonis		65.82	0.44	BGL 99			
Clostridium clostridioforme		28.39	0.32				
Bacteroides ovatus		2.34	0.03				
47675732	**Low discrimination**						**47675732**
Parabacteroides distasonis		67.80	0.78				
Clostridium clostridioforme		29.24	0.66				
Bacteroides ovatus		2.41	0.37				
47675733	**Low discrimination**						**47675733**
Parabacteroides distasonis		68.18	0.44	URE 1			
Clostridium clostridioforme		29.40	0.32				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		2.42	0.03				
47675762	**Low discrimination**						**47675762**
Parabacteroides distasonis		84.05	0.48	SOR 1			
Bacteroides ovatus		9.94	0.16				
Clostridium clostridioforme		4.29	0.20				
47675772	**Low discrimination**						**47675772**
Parabacteroides distasonis		62.52	0.44	SOR 1			
Clostridium clostridioforme		26.96	0.32				
Bacteroides ovatus		9.16	0.13				
47677000	**Low discrimination**						**47677000**
					SPO		
Actinomyces israelii		77.05	0.53	MLT 99	1		
Clostridium innocuum		16.88	0.26		99		
Clostridium butyricum		4.03	0.10		99		
47677002	**Low discrimination**						**47677002**
					SPO		
Actinomyces israelii		85.77	0.56	MLT 99	1		
Clostridium innocuum		12.53	0.26		99		
Clostridium butyricum		1.16	0.03		99		
47677010	**Acceptable identification**						**47677010**
					SPO		
Actinomyces israelii		94.24	0.38	MLT 99	1		
Clostridium innocuum		5.76	0.03	MLZ 13	99		
47677012	**Low discrimination**						**47677012**
					SPO		
Actinomyces israelii		88.31	0.41	MLT 99	1		
Bifidobacterium adolescentis		4.53	0.13	MLZ 13	1		
Clostridium innocuum		3.60	0.03		99		
47677022	**Acceptable identification**						**47677022**
					GRA		
Actinomyces israelii		94.95	0.35	MLT 99	99		
Parabacteroides distasonis		5.06	0.10	NAG 6	1		
47677040	**Good identification**						**47677040**
Actinomyces israelii		100.0	0.46	MLT 99			
47677042	**Good identification**						**47677042**
Actinomyces israelii		100.0	0.49	MLT 99			
47677100	**Low discrimination**						**47677100**
					GRA	SPO	
Actinomyces israelii		74.58	0.63	MLT 99	99	1	
Clostridium butyricum		23.42	0.33		99	99	
Fusobacterium mortiferum		1.16	0.30		1	1	
47677102	**Low discrimination**						**47677102**
					SPO		
Actinomyces israelii		79.06	0.66	MLT 99	1		
Bifidobacterium adolescentis		12.28	0.46		1		
Clostridium butyricum		6.44	0.26		99		
47677110	**Low discrimination**						**47677110**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		72.96	0.49	MLT 99	1		
Bifidobacterium dentium		10.81	0.10	MLZ 13	1		
Clostridium butyricum		8.07	0.11		99		
47677112	**Low discrimination**						**47677112**
Bifidobacterium adolescentis		36.17	0.47	MLT 99			
Actinomyces israelii		28.47	0.52	ARA 87			
Bifidobacterium dentium		28.45	0.27				
47677120	**Low discrimination**						**47677120**
					SPO		
Actinomyces israelii		45.14	0.42	MLT 99	1		
Clostridium butyricum		19.31	0.15	NAG 6	99		
Clostridium ramosum		17.57	0.19		92		
47677122	**Low discrimination**						**47677122**
					SPO	GRA	
Parabacteroides distasonis		29.79	0.44	MAN 1	1	1	
Clostridium ramosum		23.47	0.30		92	50	
Actinomyces israelii		22.61	0.45		1	99	
47677140	**Low discrimination**						**47677140**
					COC	SPO	
Actinomyces israelii		79.40	0.56	MLT 99	1	1	
Blautia producta		15.03	0.24		99	1	
Clostridium butyricum		5.57	0.15		1	99	
47677142	**Good identification**						**47677142**
Actinomyces israelii		90.78	0.59	MLT 99			
Bifidobacterium adolescentis		4.48	0.30				
Bifidobacterium breve		1.81	0.24				
47677152	**Low discrimination**						**47677152**
					COC		
Actinomyces israelii		54.21	0.45	MLT 99	1		
Bifidobacterium adolescentis		21.89	0.32	MLZ 13	1		
Blautia producta		15.26	0.15		99		
47677160	**Low discrimination**						**47677160**
					COC	SPO	
Actinomyces israelii		66.33	0.35	MLT 99	1	1	
Blautia producta		21.85	0.07	NAG 6	99	1	
Clostridium butyricum		6.34	-.03		1	99	
47677162	**Low discrimination**						**47677162**
					GRA		
Actinomyces israelii		62.86	0.38	MLT 99	99		
Bifidobacterium breve		34.99	0.29	NAG 6	99		
Parabacteroides distasonis		2.15	0.10		1		
47677202	**Good identification**						**47677202**
Actinomyces israelii		100.0	0.36	MLT 99			
47677300	**Low discrimination**						**47677300**
					SPO		
Actinomyces israelii		88.57	0.44	MLT 99	1		
Clostridium butyricum		11.43	0.07	RHA 7	99		
47677302	**Acceptable identification**						**47677302**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		90.92	0.47	MLT 99	99	1	
Parabacteroides distasonis		3.08	0.18	RHA 7	1	1	
Clostridium butyricum		3.04	0.00		99	99	
47677322	**Low discrimination**						**47677322**
					SPO	GRA	
Parabacteroides distasonis		72.20	0.52	MAN 1	1	1	
Clostridium ramosum		25.13	0.32		92	50	
Actinomyces israelii		1.38	0.26		1	99	
47677332	**Acceptable identification**						**47677332**
					GRA		
Parabacteroides distasonis		93.87	0.47	MAN 1	1		
Bacteroides ovatus		2.09	0.03		1		
Bifidobacterium dentium		1.85	-.03		99		
47677342	**Low discrimination**						**47677342**
					COC		
Actinomyces israelii		84.30	0.40	MLT 99	1		
Blautia producta		15.70	0.07	RHA 7	99		
47677400	**Low discrimination**						**47677400**
					SPO		
Actinomyces israelii		61.62	0.51	MLT 99	1		
Clostridium butyricum		35.47	0.26		99		
Clostridium innocuum		2.91	0.13		99		
47677402	**Low discrimination**						**47677402**
					SPO		
Actinomyces israelii		80.68	0.54	MLT 99	1		
Clostridium butyricum		12.04	0.19		99		
Bifidobacterium adolescentis		4.14	0.26		1		
47677410	**Low discrimination**						**47677410**
					SPO		
Actinomyces israelii		49.99	0.37	MLT 99	1		
Bifidobacterium dentium		36.23	0.10	MLZ 13	1		
Clostridium butyricum		10.14	0.04		99		
47677440	**Low discrimination**						**47677440**
					COC	SPO	
Actinomyces israelii		48.69	0.44	MLT 99	1	1	
Blautia producta		45.05	0.24		99	1	
Clostridium butyricum		6.27	0.08		1	99	
47677442	**Low discrimination**						**47677442**
					COC	SPO	
Actinomyces israelii		89.43	0.47	MLT 99	1	1	
Blautia producta		6.13	0.07		99	1	
Clostridium butyricum		2.98	0.01		1	99	
47677500	**Low discrimination**						**47677500**
					COC	SPO	
Clostridium butyricum		74.36	0.50	MLT 99	1	99	
Actinomyces israelii		21.53	0.61	MAN 10	1	1	
Blautia producta		1.94	0.24		99	1	
47677502	**Low discrimination**						**47677502**
					SPO		
Bifidobacterium adolescentis		37.81	0.60	MLT 99	1		
Actinomyces israelii		29.76	0.64		1		
Clostridium butyricum		26.65	0.43		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47677510	**Low discrimination**						**47677510**
					COC	SPO	
Bifidobacterium dentium		80.66	0.44	MLT 99	1	1	
Blautia producta		8.12	0.32	BGL 91	99	1	
Clostridium butyricum		5.47	0.28		1	99	
47677512	**Good identification**						**47677512**
Bifidobacterium dentium		91.05	0.62	MLT 99			
Bifidobacterium adolescentis		7.83	0.61				
Actinomyces israelii		0.75	0.50				
47677530	**Good identification**						**47677530**
					COC	SPO	
Bifidobacterium dentium		98.76	0.49	MLT 99	1	1	
Blautia producta		0.62	0.15	BGL 91	99	1	
Clostridium butyricum		0.33	0.10		1	99	
47677532	**Very good identification**						**47677532**
					GRA		
Bifidobacterium dentium		99.76	0.66	MLT 99	99		
Parabacteroides distasonis		0.11	0.35		1		
Bifidobacterium adolescentis		0.05	0.27		99		
47677540	**Good identification**						**47677540**
					COC	SPO	
Blautia producta		92.27	0.58	MLT 99	99	1	
Actinomyces israelii		4.03	0.54		1	1	
Clostridium butyricum		3.11	0.31		1	99	
47677542	**Low discrimination**						**47677542**
					COC		
Blautia producta		51.16	0.41	MLT 99	99		
Actinomyces israelii		30.16	0.57	BGL 10	1		
Bifidobacterium adolescentis		12.18	0.44		1		
47677550	**Very good identification**						**47677550**
					COC		
Blautia producta		99.55	0.66	MLT 99	99		
Actinomyces israelii		0.22	0.40		1		
Bifidobacterium dentium		0.10	0.10		1		
47677552	**Low discrimination**						**47677552**
					COC		
Blautia producta		81.88	0.50	MLT 99	99		
Bifidobacterium adolescentis		7.94	0.46	BGL 10	1		
Bifidobacterium dentium		7.55	0.27		1		
47677560	**Acceptable identification**						**47677560**
					COC	SPO	
Blautia producta		93.87	0.41	MLT 99	99	1	
Clostridium butyricum		2.48	0.13	NAG 10	1	99	
Actinomyces israelii		2.36	0.34		1	1	
47677570	**Good identification**						**47677570**
					COC	GRA	
Blautia producta		98.14	0.50	MLT 99	99	99	
Bifidobacterium dentium		1.59	0.15	NAG 10	1	99	
Mitsuokella multacida		0.15	0.09		1	1	
47677702	**Low discrimination**						**47677702**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		51.86	0.45	MLT 99	1		
Clostridium butyricum		19.08	0.17	RHA 7	99		
Bifidobacterium adolescentis		8.84	0.26		1		
47677722	**Low discrimination**						**47677722**
Parabacteroides distasonis		48.10	0.48	MAN 1			
Bacteroides ovatus		45.52	0.32				
Clostridium clostridioforme		2.45	0.20				
47677740	**Very good identification**						**47677740**
					COC	GRA	
Blautia producta		98.45	0.58	MLT 99	99	99	
Mitsuokella multacida		1.12	0.32		1	1	
Actinomyces israelii		0.32	0.35		1	99	
47677742	**Acceptable identification**						**47677742**
					COC	GRA	
Blautia producta		94.18	0.41	MLT 99	99	99	
Actinomyces israelii		4.18	0.38	BGL 10	1	99	
Mitsuokella multacida		1.07	0.15		1	1	
47677750	**Very good identification**						**47677750**
					COC	GRA	
Blautia producta		99.86	0.66	MLT 99	99	99	
Mitsuokella multacida		0.13	0.24		1	1	
Actinomyces israelii		0.02	0.21		1	99	
47677752	**Good identification**						**47677752**
					COC	GRA	
Blautia producta		99.46	0.50	MLT 99	99	99	
Actinomyces israelii		0.22	0.24	BGL 10	1	99	
Mitsuokella multacida		0.13	0.07		1	1	
47677760	**Good identification**						**47677760**
					COC	GRA	
Blautia producta		97.33	0.41	MLT 99	99	99	
Mitsuokella multacida		2.49	0.21	NAG 10	1	1	
Actinomyces israelii		0.18	0.15		1	99	
47677770	**Good identification**						**47677770**
					COC	GRA	
Blautia producta		99.72	0.50	MLT 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
47700000	**Good identification**						**47700000**
					GRA	SPO	
Fusobacterium mortiferum		97.82	0.41	MLT 11	1	1	
Clostridium bifermentans		2.18	-.01	FRU 99	99	99	
47700100	**Acceptable identification**						**47700100**
					GRA		
Fusobacterium mortiferum		92.39	0.47	MLT 11	1		
Bifidobacterium longum subsp. infantis		6.68	0.10	FRU 99	99		
Actinomyces naeslundii		0.93	0.25		99		
47701000	**Low discrimination**						**47701000**
					GRA		
Fusobacterium mortiferum		53.84	0.44	MLT 11	1		
Eubacterium tortuosum		32.30	0.24	FRU 99	99		
Actinomyces naeslundii		7.13	0.41		99		
47701001	**Good identification**						**47701001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.44	ESL 12		
47701003	**Acceptable identification**					**47701003**
Actinomyces naeslundii		94.53	0.44	ESL 12		
Bifidobacterium longum subsp. infantis		5.47	-.07	FRU 95		
47701021	**Good identification**					**47701021**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47701022	**Low discrimination**					**47701022**
					GRA	
Prevotella oralis		50.36	0.36	FRU 99	1	
Bifidobacterium longum subsp. infantis		44.92	0.32	LAC 94	99	
Actinomyces naeslundii		2.94	0.41		99	
47701023	**Low discrimination**					**47701023**
					GRA	
Actinomyces naeslundii		82.09	0.44	ESL 12	99	
Prevotella oralis		9.47	0.02	FRU 95	1	
Bifidobacterium longum subsp. infantis		8.44	-.03		99	
47701100	**Low discrimination**					**47701100**
					GRA	
Bifidobacterium longum subsp. infantis		78.42	0.44	FRU 99	99	
Fusobacterium mortiferum		16.44	0.50	BGL 91	1	
Actinomyces naeslundii		2.65	0.48		99	
47701101	**Low discrimination**					**47701101**
					SPO	
Actinomyces naeslundii		59.13	0.51	ESL 12	1	
Clostridium chauvoei		26.63	0.22	FRU 95	99	
Bifidobacterium longum subsp. infantis		11.77	0.10		1	
47701102	**Very good identification**					**47701102**
					GRA	
Bifidobacterium longum subsp. infantis		97.33	0.62	FRU 99	99	
Prevotella oralis		1.62	0.35		1	
Actinomyces naeslundii		0.33	0.48		99	
47701120	**Low discrimination**					**47701120**
					GRA	
Bifidobacterium longum subsp. infantis		52.27	0.49	FRU 99	99	
Prevotella melaninogenica		39.46	0.47	BGL 91	1	
Prevotella oralis		4.94	0.35		1	
47701121	**Low discrimination**					**47701121**
					GRA	
Actinomyces naeslundii		55.42	0.51	ESL 12	99	
Bifidobacterium longum subsp. infantis		19.61	0.15	FRU 95	99	
Prevotella melaninogenica		14.80	0.12		1	
47701122	**Low discrimination**					**47701122**
					GRA	
Bifidobacterium longum subsp. infantis		83.19	0.66	FRU 99	99	
Prevotella oralis		14.76	0.56		1	
Prevotella melaninogenica		1.18	0.34		1	
47701141	**Good identification**					**47701141**
Actinomyces naeslundii		100.0	0.38	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47701143	**Low discrimination**					**47701143**
Actinomyces naeslundii		89.67	0.38	ESL 12		
Bifidobacterium longum subsp. infantis		10.33	-.07	FRU 95		
47701161	**Good identification**					**47701161**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47701163	**Low discrimination**					**47701163**
Actinomyces naeslundii		83.01	0.38	ESL 12		
Bifidobacterium longum subsp. infantis		17.00	-.03	FRU 95		
47701322	**Low discrimination**					**47701322**
					GRA	
Prevotella oralis		85.26	0.50	FRU 99	1	
Bifidobacterium longum subsp. infantis		11.33	0.32		99	
Parabacteroides distasonis		3.11	0.38		1	
47701502	**Low discrimination**					**47701502**
Bifidobacterium adolescentis		56.96	0.45	FRU 99		
Bifidobacterium longum subsp. infantis		35.26	0.27	XYL 87		
Bifidobacterium longum subsp. longum		6.70	0.27			
47701512	**Low discrimination**					**47701512**
Bifidobacterium adolescentis		84.69	0.46	FRU 99		
Bifidobacterium longum subsp. longum		13.87	0.31	XYL 87		
Clostridium clostridioforme		1.01	0.10			
47701520	**Low discrimination**					**47701520**
					GRA	
Bifidobacterium longum subsp. longum		61.59	0.37	SAL 1	99	
Bifidobacterium longum subsp. infantis		14.26	0.15	FRU 90	99	
Bacteroides vulgatus		10.91	0.10		1	
47701530	**Good identification**					**47701530**
Bifidobacterium longum subsp. longum		100.0	0.41	SAL 1		
47701532	**Acceptable identification**					**47701532**
					GRA	
Bifidobacterium longum subsp. longum		90.92	0.41	SAL 1	99	
Parabacteroides distasonis		3.78	0.22	FRU 90	1	
Clostridium clostridioforme		1.66	0.10		50	
47703000	**Good identification**					**47703000**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47703001	**Good identification**					**47703001**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47703002	**Good identification**					**47703002**
Actinomyces naeslundii		95.72	0.46	ESL 12		
Bifidobacterium longum subsp. infantis		4.28	-.07	FRU 95		
47703003	**Good identification**					**47703003**
Actinomyces naeslundii		100.0	0.49	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47703020	**Good identification**					**47703020**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47703021	**Good identification**					**47703021**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47703022	**Low discrimination**					**47703022**
					GRA	
Actinomyces naeslundii		85.58	0.46	ESL 12	99	
Prevotella oralis		7.62	0.02	FRU 95	1	
Bifidobacterium longum subsp. infantis		6.80	-.03		99	
47703023	**Good identification**					**47703023**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47703041	**Good identification**					**47703041**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47703043	**Good identification**					**47703043**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47703061	**Good identification**					**47703061**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47703063	**Good identification**					**47703063**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47703100	**Low discrimination**					**47703100**
					GRA	
Actinomyces naeslundii		81.34	0.53	ESL 12	99	
Bifidobacterium longum subsp. infantis		12.51	0.10	FRU 95	99	
Fusobacterium mortiferum		2.62	0.16		1	
47703101	**Very good identification**					**47703101**
Actinomyces naeslundii		100.0	0.56	ESL 12		
47703103	**Very good identification**					**47703103**
Actinomyces naeslundii		98.96	0.56	ESL 12		
Bifidobacterium longum subsp. infantis		1.04	-.07	FRU 95		
47703120	**Low discrimination**					**47703120**
					GRA	
Actinomyces naeslundii		63.95	0.53	ESL 12	99	
Bifidobacterium longum subsp. infantis		17.48	0.15	FRU 95	99	
Prevotella melaninogenica		13.20	0.12		1	
47703121	**Very good identification**					**47703121**
Actinomyces naeslundii		100.0	0.56	ESL 12		
47703123	**Very good identification**					**47703123**
Actinomyces naeslundii		98.17	0.56	ESL 12		
Bifidobacterium longum subsp. infantis		1.83	-.03	FRU 95		
47703140	**Low discrimination**					**47703140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		89.09	0.40	ESL 12			
Bifidobacterium breve		10.91	0.10	FRU 95			
47703141	**Good identification**						**47703141**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47703143	**Good identification**						**47703143**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47703160	**Low discrimination**						**47703160**
Actinomyces naeslundii		82.12	0.40	ESL 12			
Bifidobacterium breve		17.88	0.15	FRU 95			
47703161	**Good identification**						**47703161**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47703163	**Good identification**						**47703163**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47703502	**Good identification**						**47703502**
Bifidobacterium adolescentis		97.98	0.43	FRU 99			
Actinomyces israelii		1.27	0.17	XYL 87			
Bifidobacterium longum subsp. infantis		0.75	-.07				
47703512	**Good identification**						**47703512**
Bifidobacterium adolescentis		100.0	0.45	FRU 99			
47705100	**Acceptable identification**						**47705100**
					GRA		
Bifidobacterium longum subsp. infantis		93.55	0.43	FRU 99	99		
Eubacterium contortum		3.83	0.24	BGL 91	99		
Fusobacterium mortiferum		1.80	0.31		1		
47705102	**Very good identification**						**47705102**
					GRA		
Bifidobacterium longum subsp. infantis		97.58	0.60	FRU 99	99		
Bifidobacterium adolescentis		1.95	0.45		99		
Prevotella oralis		0.25	0.19		1		
47705112	**Low discrimination**						**47705112**
Bifidobacterium adolescentis		70.32	0.46	FRU 99			
Bifidobacterium longum subsp. infantis		29.14	0.26	ARA 87			
Clostridium clostridioforme		0.54	0.07				
47705120	**Acceptable identification**						**47705120**
					GRA		
Bifidobacterium longum subsp. infantis		92.57	0.47	FRU 99	99		
Bacteroides vulgatus		3.22	0.20	BGL 91	1		
Prevotella oralis		1.32	0.19		1		
47705122	**Good identification**						**47705122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		93.93	0.64	FRU 99	1	99	
Prevotella oralis		2.52	0.41		1	1	
Bacteroides fragilis		2.43	0.32		99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47705300	**Very good identification**					**47705300**
					GRA	
Eubacterium contortum		99.61	0.58	FRU 99	99	
Bifidobacterium longum subsp. infantis		0.25	0.09		99	
Bacteroides vulgatus		0.14	0.02		1	
47705302	**Low discrimination**					**47705302**
					GRA	
Eubacterium contortum		85.27	0.45	FRU 99	99	
Bifidobacterium longum subsp. infantis		12.17	0.26		99	
Prevotella oralis		1.29	0.13		1	
47705320	**Low discrimination**					**47705320**
					GRA	
Eubacterium contortum		54.75	0.45	FRU 99	99	
Bacteroides vulgatus		42.62	0.36		1	
Bifidobacterium longum subsp. infantis		1.37	0.13		99	
47705322	**Low discrimination**					**47705322**
					GRA	
Parabacteroides distasonis		50.97	0.55	FRU 99	1	
Prevotella oralis		19.18	0.35		1	
Bifidobacterium longum subsp. infantis		16.87	0.30		99	
47705332	**Acceptable identification**					**47705332**
Parabacteroides distasonis		93.93	0.50	FRU 99		
Clostridium clostridioforme		4.42	0.22			
Prevotella oralis		0.66	0.00			
47705500	**Low discrimination**					**47705500**
					SPO	
Bifidobacterium longum subsp. longum		45.58	0.35	SAL 1	1	
Eubacterium contortum		21.49	0.23	FRU 90	1	
Clostridium butyricum		10.54	0.11		99	
47705502	**Low discrimination**					**47705502**
Bifidobacterium adolescentis		87.29	0.59	FRU 99		
Bifidobacterium longum subsp. infantis		6.61	0.26			
Bifidobacterium longum subsp. longum		4.60	0.35			
47705510	**Low discrimination**					**47705510**
Bifidobacterium longum subsp. longum		87.90	0.39	SAL 1		
Bifidobacterium adolescentis		12.10	0.26	FRU 90		
47705512	**Good identification**					**47705512**
Bifidobacterium adolescentis		91.62	0.60	FRU 99		
Bifidobacterium longum subsp. longum		6.73	0.39			
Clostridium clostridioforme		1.65	0.27			
47705520	**Low discrimination**					**47705520**
					GRA	
Bacteroides vulgatus		72.65	0.39	FRU 99	1	
Bifidobacterium longum subsp. longum		25.12	0.46	RHA 90	99	
Bifidobacterium longum subsp. infantis		1.59	0.13		99	
47705522	**Low discrimination**					**47705522**
					GRA	
Bifidobacterium longum subsp. longum		45.53	0.46	SAL 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. infantis		29.06	0.30	FRU 90	99	
Parabacteroides distasonis		17.94	0.43		1	
47705530	**Good identification**					**47705530**
					GRA	
Bifidobacterium longum subsp. longum		98.31	0.50	SAL 1	99	
Bacteroides vulgatus		1.69	0.05	FRU 90	1	
47705532	**Low discrimination**					**47705532**
					GRA	
Bifidobacterium longum subsp. longum		81.69	0.50	SAL 1	99	
Parabacteroides distasonis		10.18	0.38	FRU 90	1	
Clostridium clostridioforme		5.01	0.27		50	
47705542	**Good identification**					**47705542**
Bifidobacterium adolescentis		99.00	0.43	FRU 99		
Actinomyces israelii		1.00	0.15	SOR 11		
47705552	**Good identification**					**47705552**
Bifidobacterium adolescentis		100.0	0.45	FRU 99		
47705700	**Very good identification**					**47705700**
					GRA	
Eubacterium contortum		97.80	0.57	FRU 99	99	
Bacteroides vulgatus		2.20	0.21		1	
47705702	**Acceptable identification**					**47705702**
Eubacterium contortum		90.03	0.44	FRU 99		
Clostridium clostridioforme		7.35	0.31			
Bifidobacterium adolescentis		2.10	0.25			
47705712	**Low discrimination**					**47705712**
Clostridium clostridioforme		89.93	0.42	FRU 99		
Bifidobacterium adolescentis		6.88	0.26	TRE 88		
Eubacterium contortum		2.44	0.10			
47705720	**Acceptable identification**					**47705720**
					GRA	
Bacteroides vulgatus		92.60	0.56	FRU 99	1	
Eubacterium contortum		7.33	0.44		99	
Parabacteroides distasonis		0.04	0.17		1	
47705722	**Low discrimination**					**47705722**
					GRA	
Parabacteroides distasonis		58.30	0.51	FRU 99	1	
Eubacterium contortum		17.92	0.31		99	
Bacteroides vulgatus		12.96	0.21		1	
47705732	**Low discrimination**					**47705732**
					GRA	
Clostridium clostridioforme		53.95	0.42	FRU 99	50	
Parabacteroides distasonis		44.84	0.46	TRE 88	1	
Bifidobacterium longum subsp. longum		1.21	0.15		99	
47707102	**Low discrimination**					**47707102**
Bifidobacterium adolescentis		55.51	0.43	FRU 99		
Bifidobacterium longum subsp. infantis		34.36	0.26	ARA 87		
Actinomyces israelii		10.13	0.37			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47707112	**Good identification**					**47707112**
Bifidobacterium adolescentis		97.82	0.45	FRU 99		
Actinomyces israelii		2.18	0.23	ARA 87		
47707502	**Very good identification**					**47707502**
Bifidobacterium adolescentis		97.82	0.57	FRU 99		
Actinomyces israelii		2.18	0.35			
47707512	**Very good identification**					**47707512**
Bifidobacterium adolescentis		99.56	0.59	FRU 99		
Actinomyces israelii		0.27	0.21			
Bifidobacterium dentium		0.17	-.07			
47707542	**Acceptable identification**					**47707542**
Bifidobacterium adolescentis		93.44	0.42	FRU 99		
Actinomyces israelii		6.56	0.28	SOR 11		
47707552	**Good identification**					**47707552**
Bifidobacterium adolescentis		99.15	0.43	FRU 99		
Actinomyces israelii		0.85	0.14	SOR 11		
47710000	**Very good identification**					**47710000**
					GRA	
Fusobacterium mortiferum		99.45	0.75	MLT 11	1	
Bifidobacterium longum subsp. infantis		0.17	0.10		99	
Actinomyces naeslundii		0.16	0.39		99	
47710001	**Low discrimination**					**47710001**
					GRA	
Fusobacterium mortiferum		80.92	0.41	MLT 11	1	
Actinomyces naeslundii		19.08	0.42	URE 1	99	
47710002	**Low discrimination**					**47710002**
					GRA	
Fusobacterium mortiferum		82.87	0.59	MLT 11	1	
Bifidobacterium longum subsp. infantis		12.85	0.27	BGL 10	99	
Actinomyces viscosus		1.75	0.18		99	
47710003	**Low discrimination**					**47710003**
					GRA	
Actinomyces naeslundii		53.41	0.42	ESL 12	99	
Fusobacterium mortiferum		25.17	0.25	GAL 96	1	
Actinomyces viscosus		17.52	0.09		99	
47710010	**Good identification**					**47710010**
Fusobacterium mortiferum		100.0	0.41	MLT 11		
47710020	**Low discrimination**					**47710020**
					GRA	
Fusobacterium mortiferum		56.78	0.41	MLT 11	1	
Bifidobacterium longum subsp. infantis		17.03	0.15	NAG 1	99	
Prevotella oralis		10.16	0.14		1	
47710021	**Good identification**					**47710021**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47710022	**Low discrimination**					**47710022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Prevotella oralis		50.09	0.36	GAL 99	1		
Bifidobacterium longum subsp. infantis		44.67	0.32	LAC 94	99		
Actinomyces naeslundii		2.32	0.39		99		
47710023	**Low discrimination**						**47710023**
					GRA		
Actinomyces naeslundii		78.40	0.42	ESL 12	99		
Prevotella oralis		11.42	0.02	GAL 96	1		
Bifidobacterium longum subsp. infantis		10.18	-.03		99		
47710040	**Good identification**						**47710040**
					GRA		
Fusobacterium mortiferum		97.30	0.41	MLT 11	1		
Actinomyces naeslundii		2.70	0.26	SOR 1	99		
47710100	**Good identification**						**47710100**
					GRA		
Fusobacterium mortiferum		92.98	0.81	MLT 11	1		
Bifidobacterium longum subsp. infantis		6.72	0.44		99		
Actinomyces naeslundii		0.18	0.47		99		
47710101	**Low discrimination**						**47710101**
					GRA		
Fusobacterium mortiferum		73.54	0.47	MLT 11	1		
Actinomyces naeslundii		21.15	0.50	URE 1	99		
Bifidobacterium longum subsp. infantis		5.32	0.10		99		
47710102	**Low discrimination**						**47710102**
					GRA		
Bifidobacterium longum subsp. infantis		84.82	0.62	GAL 99	99		
Fusobacterium mortiferum		12.90	0.65		1		
Prevotella oralis		1.41	0.35		1		
47710110	**Acceptable identification**						**47710110**
					GRA		
Fusobacterium mortiferum		91.53	0.47	MLT 11	1		
Bifidobacterium longum subsp. infantis		6.62	0.10	MLZ 1	99		
Bifidobacterium breve		0.93	0.10		99		
47710120	**Low discrimination**						**47710120**
					GRA		
Bifidobacterium longum subsp. infantis		76.82	0.49	GAL 99	99		
Prevotella oralis		7.25	0.35	BGL 91	1		
Fusobacterium mortiferum		6.04	0.47		1		
47710121	**Low discrimination**						**47710121**
					GRA		
Actinomyces naeslundii		64.32	0.50	ESL 12	99		
Bifidobacterium longum subsp. infantis		28.74	0.15	GAL 96	99		
Prevotella oralis		2.71	0.00		1		
47710122	**Low discrimination**						**47710122**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		84.04	0.66	GAL 99	99	1	
Prevotella oralis		14.91	0.56		1	1	
Capnocytophaga ochracea		0.34	0.29		1	90	
47710140	**Low discrimination**						**47710140**
					GRA		
Fusobacterium mortiferum		89.58	0.47	MLT 11	1		
Bifidobacterium longum subsp. infantis		6.48	0.10	SOR 1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		3.03	0.34		99	
47710141	**Low discrimination**					**47710141**
					GRA	
Actinomyces naeslundii		83.40	0.37	ESL 12	99	
Fusobacterium mortiferum		16.60	0.13	GAL 96	1	
47710143	**Low discrimination**					**47710143**
Actinomyces naeslundii		87.30	0.37	ESL 12		
Bifidobacterium longum subsp. infantis		12.71	-.07	GAL 96		
47710161	**Good identification**					**47710161**
Actinomyces naeslundii		100.0	0.37	ESL 12		
47710163	**Low discrimination**					**47710163**
Actinomyces naeslundii		79.45	0.37	ESL 12		
Bifidobacterium longum subsp. infantis		20.55	-.03	GAL 96		
47710200	**Good identification**					**47710200**
Fusobacterium mortiferum		100.0	0.41	MLT 11		
47710300	**Low discrimination**					**47710300**
					GRA	
Fusobacterium mortiferum		74.67	0.47	MLT 11	1	
Eubacterium contortum		17.92	0.24	RHA 1	99	
Bifidobacterium longum subsp. infantis		5.40	0.10		99	
47710322	**Low discrimination**					**47710322**
					GRA	
Prevotella oralis		84.97	0.50	GAL 99	1	
Bifidobacterium longum subsp. infantis		11.29	0.32		99	
Parabacteroides distasonis		3.10	0.38		1	
47710400	**Good identification**					**47710400**
Fusobacterium mortiferum		100.0	0.41	MLT 11		
47710500	**Acceptable identification**					**47710500**
					GRA	
Fusobacterium mortiferum		91.17	0.47	MLT 11	1	
Bifidobacterium longum subsp. infantis		6.59	0.10	ARA 1	99	
Bifidobacterium longum subsp. longum		1.15	0.09		99	
47710502	**Low discrimination**					**47710502**
Bifidobacterium adolescentis		54.10	0.45	GAL 99		
Bifidobacterium longum subsp. infantis		33.49	0.27	XYL 87		
Clostridium clostridioforme		5.71	0.25			
47710512	**Low discrimination**					**47710512**
Bifidobacterium adolescentis		70.75	0.46	GAL 99		
Clostridium clostridioforme		27.84	0.36	XYL 87		
Bifidobacterium longum subsp. longum		1.05	0.13			
47710532	**Low discrimination**					**47710532**
					GRA	
Clostridium clostridioforme		77.68	0.36	TRE 88	50	
Bifidobacterium longum subsp. longum		11.76	0.24	XYL 91	99	
Parabacteroides distasonis		5.37	0.22	RHA 88	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47710612	**Good identification**					**47710612**
Clostridium clostridioforme		100.0	0.42	TRE 88		
47710632	**Good identification**					**47710632**
Clostridium clostridioforme		100.0	0.42	TRE 88		
47710702	**Acceptable identification**					**47710702**
Clostridium clostridioforme		94.43	0.40	TRE 88		
Eubacterium contortum		3.58	0.10	XYL 91		
Bifidobacterium adolescentis		1.23	0.11			
47710712	**Good identification**					**47710712**
Clostridium clostridioforme		99.65	0.51	TRE 88		
Bifidobacterium adolescentis		0.35	0.12	XYL 91		
47710722	**Low discrimination**					**47710722**
Clostridium clostridioforme		72.91	0.40	TRE 88		
Parabacteroides distasonis		17.45	0.35	XYL 91		
Prevotella oralis		7.88	0.16			
47710732	**Good identification**					**47710732**
Clostridium clostridioforme		97.25	0.51	TRE 88		
Parabacteroides distasonis		2.75	0.30	XYL 91		
47711000	**Low discrimination**					**47711000**
					GRA	
Fusobacterium mortiferum		85.63	0.78	MLT 11	1	
Bifidobacterium longum subsp. infantis		9.63	0.44		99	
Actinomyces naeslundii		2.18	0.62		99	
47711001	**Low discrimination**					**47711001**
					GRA	
Actinomyces naeslundii		77.02	0.66	ESL 12	99	
Fusobacterium mortiferum		20.42	0.44		1	
Bifidobacterium longum subsp. infantis		2.30	0.10		99	
47711002	**Low discrimination**					**47711002**
					GRA	
Bifidobacterium longum subsp. infantis		80.14	0.62	LAC 99	99	
Prevotella oralis		8.41	0.48		1	
Fusobacterium mortiferum		7.83	0.62		1	
47711003	**Low discrimination**					**47711003**
					GRA	
Actinomyces naeslundii		72.95	0.66	ESL 12	99	
Bifidobacterium longum subsp. infantis		21.99	0.27		99	
Prevotella oralis		2.31	0.14		1	
47711010	**Low discrimination**					**47711010**
					GRA	
Fusobacterium mortiferum		59.46	0.44	MLT 11	1	
Lactobacillus jensenii		25.05	0.28	MLZ 1	99	
Actinomyces naeslundii		7.87	0.41		99	
47711011	**Good identification**					**47711011**
Actinomyces naeslundii		100.0	0.44	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47711013	**Acceptable identification**					**47711013**
Actinomyces naeslundii		94.53	0.44	ESL 12		
Bifidobacterium longum subsp. infantis		5.47	-.07	MLZ 5		
47711020	**Low discrimination**					**47711020**
					GRA	
Bifidobacterium longum subsp. infantis		44.64	0.49	LAC 99	99	
Prevotella oralis		26.64	0.48	BGL 91	1	
Clostridium perfringens		15.58	0.48		99	
47711021	**Good identification**					**47711021**
					GRA	
Actinomyces naeslundii		90.36	0.66	ESL 12	99	
Bifidobacterium longum subsp. infantis		4.79	0.15		99	
Prevotella oralis		2.86	0.14		1	
47711022	**Low discrimination**					**47711022**
					GRA	
Prevotella oralis		51.81	0.70	LAC 94	1	
Bifidobacterium longum subsp. infantis		46.21	0.66		99	
Clostridium perfringens		0.75	0.42		99	
47711023	**Low discrimination**					**47711023**
					GRA	
Actinomyces naeslundii		46.61	0.66	ESL 12	99	
Prevotella oralis		28.00	0.36		1	
Bifidobacterium longum subsp. infantis		24.98	0.32		99	
47711031	**Good identification**					**47711031**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47711032	**Low discrimination**					**47711032**
					GRA	
Prevotella oralis		44.90	0.36	LAC 94	1	
Bifidobacterium longum subsp. infantis		40.04	0.32	MLZ 1	99	
Bifidobacterium breve		5.63	0.32		99	
47711033	**Low discrimination**					**47711033**
					GRA	
Actinomyces naeslundii		82.09	0.44	ESL 12	99	
Prevotella oralis		9.47	0.02	MLZ 5	1	
Bifidobacterium longum subsp. infantis		8.44	-.03		99	
47711040	**Low discrimination**					**47711040**
					GRA	
Fusobacterium mortiferum		60.52	0.44	MLT 11	1	
Actinomyces naeslundii		26.86	0.50	SOR 1	99	
Bifidobacterium longum subsp. infantis		6.80	0.10		99	
47711041	**Good identification**					**47711041**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47711043	**Good identification**					**47711043**
Actinomyces naeslundii		98.30	0.53	ESL 12		
Bifidobacterium longum subsp. infantis		1.70	-.07			
47711060	**Low discrimination**					**47711060**
Clostridium perfringens		61.15	0.35	LAC 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		20.63	0.50			
Bifidobacterium longum subsp. infantis		9.29	0.15			
47711061	**Good identification**					**47711061**
Actinomyces naeslundii		98.04	0.53	ESL 12		
Clostridium perfringens		1.96	0.01			
47711062	**Low discrimination**					**47711062**
					GRA	
Prevotella oralis		39.11	0.36	LAC 94	1	
Bifidobacterium longum subsp. infantis		34.88	0.32	SOR 1	99	
Clostridium perfringens		10.68	0.30		99	
47711063	**Acceptable identification**					**47711063**
					GRA	
Actinomyces naeslundii		93.89	0.53	ESL 12	99	
Prevotella oralis		3.23	0.02		1	
Bifidobacterium longum subsp. infantis		2.88	-.03		99	
47711100	**Low discrimination**					**47711100**
					GRA	
Bifidobacterium longum subsp. infantis		80.85	0.79	BGL 91	99	
Fusobacterium mortiferum		16.95	0.85		1	
Prevotella oralis		0.71	0.47		1	
47711101	**Low discrimination**					**47711101**
					GRA	
Bifidobacterium longum subsp. infantis		44.54	0.44	URE 1	99	
Actinomyces naeslundii		42.96	0.73	BGL 91	99	
Fusobacterium mortiferum		9.34	0.50		1	
47711102	**Very good identification**					**47711102**
					GRA	
Bifidobacterium longum subsp. infantis		97.58	0.96		99	
Prevotella oralis		1.62	0.69		1	
Bifidobacterium adolescentis		0.24	0.65		99	
47711103	**Low discrimination**					**47711103**
					GRA	
Bifidobacterium longum subsp. infantis		89.24	0.62	URE 1	99	
Actinomyces naeslundii		8.51	0.73		99	
Prevotella oralis		1.48	0.35		1	
47711110	**Low discrimination**					**47711110**
					GRA	
Bifidobacterium longum subsp. infantis		68.94	0.44	MLZ 1	99	
Fusobacterium mortiferum		14.45	0.50	BGL 91	1	
Bifidobacterium breve		9.69	0.44		99	
47711111	**Low discrimination**					**47711111**
					GRA	
Actinomyces naeslundii		78.82	0.51	ESL 12	99	
Bifidobacterium longum subsp. infantis		15.69	0.10	MLZ 5	99	
Fusobacterium mortiferum		3.29	0.16		1	
47711112	**Low discrimination**					**47711112**
Bifidobacterium longum subsp. infantis		67.48	0.62	MLZ 1		
Bifidobacterium adolescentis		19.91	0.66			
Bifidobacterium breve		9.48	0.62			
47711120	**Low discrimination**					**47711120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium longum subsp. infantis		82.46	0.83	BGL 91	99		
Prevotella oralis		7.79	0.69		1		
Prevotella melaninogenica		5.66	0.63		1		
47711121	**Low discrimination**						**47711121**
					GRA		
Bifidobacterium longum subsp. infantis		56.81	0.49	URE 1	99		
Actinomyces naeslundii		30.82	0.73	BGL 91	99		
Prevotella oralis		5.36	0.35		1		
47711122	**Low discrimination**						**47711122**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		84.20	1.00		99	1	
Prevotella oralis		14.94	0.91		1	1	
Capnocytophaga ochracea		0.34	0.63		1	90	
47711123	**Low discrimination**						**47711123**
					GRA		
Bifidobacterium longum subsp. infantis		80.58	0.66	URE 1	99		
Prevotella oralis		14.30	0.56		1		
Actinomyces naeslundii		4.32	0.73		99		
47711130	**Low discrimination**						**47711130**
					GRA		
Bifidobacterium longum subsp. infantis		70.24	0.49	MLZ 1	99		
Bifidobacterium breve		9.87	0.49	BGL 91	99		
Prevotella oralis		6.63	0.35		1		
47711131	**Low discrimination**						**47711131**
Actinomyces naeslundii		68.44	0.51	ESL 12			
Bifidobacterium longum subsp. infantis		24.21	0.15	MLZ 5			
Bifidobacterium breve		3.40	0.15				
47711132	**Low discrimination**						**47711132**
					GRA		
Bifidobacterium longum subsp. infantis		70.36	0.66	MLZ 1	99		
Prevotella oralis		12.48	0.56		1		
Bifidobacterium breve		9.89	0.66		99		
47711140	**Low discrimination**						**47711140**
					GRA		
Bifidobacterium longum subsp. infantis		65.33	0.44	SOR 1	99		
Fusobacterium mortiferum		13.69	0.50	BGL 91	1		
Bifidobacterium breve		9.18	0.44		99		
47711141	**Good identification**						**47711141**
					GRA		
Actinomyces naeslundii		92.58	0.60	ESL 12	99		
Bifidobacterium longum subsp. infantis		5.50	0.10		99		
Fusobacterium mortiferum		1.15	0.16		1		
47711142	**Low discrimination**						**47711142**
Bifidobacterium longum subsp. infantis		82.29	0.62	SOR 1			
Bifidobacterium breve		11.56	0.62				
Bifidobacterium adolescentis		2.46	0.49				
47711143	**Low discrimination**						**47711143**
Actinomyces naeslundii		58.40	0.60	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis		35.05	0.27				
Bifidobacterium breve		4.93	0.27				
47711150	**Low discrimination**						**47711150**
Bifidobacterium breve		85.17	0.44	CEL 99			
Bifidobacterium longum subsp. infantis		6.12	0.10	BGL 91			
Actinomyces naeslundii		3.62	0.35				
47711151	**Low discrimination**						**47711151**
Actinomyces naeslundii		86.32	0.38	ESL 12			
Bifidobacterium breve		13.68	0.10	MLZ 5			
47711152	**Low discrimination**						**47711152**
Bifidobacterium breve		74.29	0.62	CEL 99			
Bifidobacterium adolescentis		19.28	0.51				
Bifidobacterium longum subsp. infantis		5.34	0.27				
47711160	**Low discrimination**						**47711160**
Bifidobacterium longum subsp. infantis		50.18	0.49	SOR 1			
Clostridium perfringens		24.46	0.50	BGL 91			
Bifidobacterium breve		7.05	0.49				
47711161	**Low discrimination**						**47711161**
Actinomyces naeslundii		83.49	0.60	ESL 12			
Bifidobacterium longum subsp. infantis		8.81	0.15				
Clostridium perfringens		4.29	0.16				
47711162	**Low discrimination**						**47711162**
					GRA		
Bifidobacterium longum subsp. infantis		71.42	0.66	SOR 1	99		
Prevotella oralis		12.67	0.56		1		
Bifidobacterium breve		10.04	0.66		99		
47711170	**Low discrimination**						**47711170**
					GRA	CAT	
Bifidobacterium breve		84.15	0.49	CEL 99	99	1	
Bifidobacterium longum subsp. infantis		6.05	0.15	BGL 91	99	1	
Capnocytophaga ochracea		3.35	0.14		1	90	
47711171	**Low discrimination**						**47711171**
Actinomyces naeslundii		78.02	0.38	ESL 12			
Bifidobacterium breve		21.98	0.15	MLZ 5			
47711172	**Low discrimination**						**47711172**
					GRA	CAT	
Bifidobacterium breve		88.57	0.66	CEL 99	99	1	
Bifidobacterium longum subsp. infantis		6.37	0.32		99	1	
Capnocytophaga ochracea		3.14	0.30		1	90	
47711200	**Low discrimination**						**47711200**
					GRA		
Fusobacterium mortiferum		55.86	0.44	MLT 11	1		
Eubacterium contortum		20.85	0.24	RHA 1	99		
Prevotella oralis		14.88	0.20		1		
47711202	**Low discrimination**						**47711202**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		77.71	0.42	LAC 94	1	
Bifidobacterium longum subsp. infantis		17.46	0.27	NAG 95	99	
Fusobacterium mortiferum		1.71	0.28		1	
47711220	**Acceptable identification**					**47711220**
					GRA	
Prevotella oralis		91.13	0.42	LAC 94	1	
Bifidobacterium longum subsp. infantis		3.60	0.15	BGL 95	99	
Bacteroides vulgatus		1.87	0.07		1	
47711222	**Very good identification**					**47711222**
					GRA	
Prevotella oralis		97.17	0.64	LAC 94	1	
Bifidobacterium longum subsp. infantis		2.04	0.32		99	
Parabacteroides distasonis		0.56	0.38		1	
47711300	**Low discrimination**					**47711300**
					GRA	
Eubacterium contortum		67.31	0.58	XYL 99	99	
Bifidobacterium longum subsp. infantis		20.28	0.44		99	
Prevotella oralis		7.60	0.41		1	
47711302	**Low discrimination**					**47711302**
					GRA	
Bifidobacterium longum subsp. infantis		56.19	0.62	RHA 1	99	
Prevotella oralis		39.58	0.62		1	
Eubacterium contortum		3.26	0.45		99	
47711312	**Low discrimination**					**47711312**
					GRA	
Clostridium clostridioforme		30.61	0.39	TRE 88	50	
Bifidobacterium longum subsp. infantis		28.06	0.27	XYL 91	99	
Prevotella oralis		19.77	0.28	ARA 94	1	
47711320	**Low discrimination**					**47711320**
					GRA	
Prevotella oralis		66.58	0.62	BGL 95	1	
Bifidobacterium longum subsp. infantis		16.62	0.49		99	
Bacteroides vulgatus		8.61	0.42		1	
47711322	**Low discrimination**					**47711322**
					GRA	
Prevotella oralis		85.28	0.84		1	
Bifidobacterium longum subsp. infantis		11.33	0.66		99	
Parabacteroides distasonis		3.11	0.72		1	
47711323	**Low discrimination**					**47711323**
					GRA	
Prevotella oralis		84.88	0.50	URE 1	1	
Bifidobacterium longum subsp. infantis		11.28	0.32		99	
Parabacteroides distasonis		3.10	0.38		1	
47711332	**Low discrimination**					**47711332**
					GRA	
Parabacteroides distasonis		61.06	0.68	XYL 90	1	
Prevotella oralis		31.39	0.50		1	
Bifidobacterium longum subsp. infantis		4.17	0.32		99	
47711362	**Low discrimination**					**47711362**
					GRA	
Prevotella oralis		83.26	0.50	SOR 1	1	
Bifidobacterium longum subsp. infantis		11.06	0.32		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		3.04	0.38		1	
47711400	**Low discrimination**					**47711400**
					GRA	
Fusobacterium mortiferum		77.14	0.44	MLT 11	1	
Bifidobacterium longum subsp. infantis		8.67	0.10	ARA 1	99	
Actinomyces israelii		4.38	0.25		99	
47711402	**Low discrimination**					**47711402**
Bifidobacterium adolescentis		51.66	0.45	XYL 87		
Bifidobacterium longum subsp. infantis		31.98	0.27	LAC 99		
Clostridium clostridioforme		4.89	0.24			
47711412	**Low discrimination**					**47711412**
Bifidobacterium adolescentis		72.55	0.46	XYL 87		
Clostridium clostridioforme		25.58	0.35	LAC 99		
Bifidobacterium longum subsp. longum		1.08	0.13			
47711422	**Low discrimination**					**47711422**
					GRA	
Prevotella oralis		46.50	0.36	LAC 94	1	
Bifidobacterium longum subsp. infantis		41.47	0.32	ARA 1	99	
Clostridium clostridioforme		3.57	0.24		50	
47711432	**Low discrimination**					**47711432**
					GRA	
Clostridium clostridioforme		74.09	0.35	TRE 88	50	
Bifidobacterium longum subsp. longum		12.51	0.24	XYL 91	99	
Parabacteroides distasonis		5.72	0.22	RHA 88	1	
47711500	**Low discrimination**					**47711500**
					GRA	
Bifidobacterium longum subsp. infantis		58.94	0.44	ARA 1	99	
Fusobacterium mortiferum		12.36	0.50	BGL 91	1	
Bifidobacterium longum subsp. longum		10.29	0.43		99	
47711502	**Low discrimination**					**47711502**
Bifidobacterium adolescentis		60.29	0.79	XYL 87		
Bifidobacterium longum subsp. infantis		37.32	0.62			
Bifidobacterium longum subsp. longum		0.64	0.43			
47711503	**Low discrimination**					**47711503**
Bifidobacterium adolescentis		58.56	0.45	XYL 87		
Bifidobacterium longum subsp. infantis		36.25	0.27	URE 1		
Actinomyces naeslundii		3.46	0.39			
47711510	**Low discrimination**					**47711510**
Bifidobacterium longum subsp. longum		56.95	0.47	SAL 1		
Bifidobacterium adolescentis		38.64	0.46			
Bifidobacterium longum subsp. infantis		1.94	0.10			
47711512	**Very good identification**					**47711512**
Bifidobacterium adolescentis		96.70	0.80	XYL 87		
Bifidobacterium longum subsp. longum		1.44	0.47			
Clostridium clostridioforme		1.15	0.44			
47711513	**Good identification**					**47711513**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		96.91	0.46	XYL 87			
Bifidobacterium longum subsp. longum		1.44	0.13	URE 1			
Clostridium clostridioforme		1.16	0.10				
47711520	**Low discrimination**						**47711520**
					GRA		
Bifidobacterium longum subsp. infantis		40.70	0.49	ARA 1	99		
Bacteroides vulgatus		31.13	0.44	BGL 91	1		
Bifidobacterium longum subsp. longum		15.98	0.54		99		
47711522	**Low discrimination**						**47711522**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		73.41	0.66	ARA 1	99	1	
Prevotella oralis		13.03	0.56		1	1	
Capnocytophaga ochracea		5.23	0.50		1	90	
47711530	**Good identification**						**47711530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		94.98	0.58	SAL 1	99	1	
Bifidobacterium longum subsp. infantis		1.44	0.15		99	1	
Capnocytophaga ochracea		1.26	0.17		1	90	
47711532	**Low discrimination**						**47711532**
					GRA		
Bifidobacterium longum subsp. longum		46.56	0.58	SAL 1	99		
Parabacteroides distasonis		21.28	0.56		1		
Clostridium clostridioforme		9.32	0.44		50		
47711542	**Good identification**						**47711542**
					GRA	CAT	
Bifidobacterium adolescentis		93.08	0.63	XYL 87	99	1	
Bifidobacterium longum subsp. infantis		4.71	0.27	SOR 11	99	1	
Capnocytophaga ochracea		0.73	0.17		1	90	
47711552	**Very good identification**						**47711552**
Bifidobacterium adolescentis		99.02	0.65	XYL 87			
Bifidobacterium breve		0.58	0.27	SOR 11			
Bifidobacterium longum subsp. longum		0.12	0.13				
47711602	**Acceptable identification**						**47711602**
					GRA		
Clostridium clostridioforme		90.42	0.39	TRE 88	50		
Eubacterium contortum		3.82	0.10	XYL 91	99		
Prevotella oralis		3.63	0.08		1		
47711612	**Good identification**						**47711612**
Clostridium clostridioforme		99.61	0.50	TRE 88			
Bifidobacterium adolescentis		0.39	0.12	XYL 91			
47711622	**Low discrimination**						**47711622**
Clostridium clostridioforme		48.63	0.39	TRE 88			
Prevotella oralis		37.07	0.30	XYL 91			
Parabacteroides distasonis		12.99	0.35				
47711632	**Good identification**						**47711632**
Clostridium clostridioforme		96.94	0.50	TRE 88			
Parabacteroides distasonis		3.06	0.30	XYL 91			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47711700	**Good identification**						**47711700**
					GRA		
Eubacterium contortum		94.78	0.57	XYL 99	99		
Bacteroides vulgatus		4.31	0.27		1		
Bifidobacterium longum subsp. infantis		0.35	0.10		99		
47711702	**Low discrimination**						**47711702**
Eubacterium contortum		38.69	0.44	XYL 99			
Clostridium clostridioforme		30.93	0.48				
Bifidobacterium adolescentis		13.32	0.45				
47711712	**Low discrimination**						**47711712**
					GRA		
Clostridium clostridioforme		88.49	0.59	TRE 88	50		
Bifidobacterium adolescentis		10.22	0.46	XYL 91	99		
Parabacteroides distasonis		0.84	0.30		1		
47711720	**Very good identification**						**47711720**
					GRA		
Bacteroides vulgatus		95.23	0.61	XYL 98	1		
Eubacterium contortum		3.73	0.44		99		
Prevotella oralis		0.56	0.28		1		
47711722	**Low discrimination**						**47711722**
Parabacteroides distasonis		58.22	0.69	XYL 90			
Prevotella oralis		26.29	0.50				
Clostridium clostridioforme		7.37	0.48				
47711732	**Low discrimination**						**47711732**
Clostridium clostridioforme		51.20	0.59	TRE 88			
Parabacteroides distasonis		47.80	0.64	XYL 91			
Prevotella oralis		0.41	0.16				
47712000	**Low discrimination**						**47712000**
					GRA	CAT	
Fusobacterium mortiferum		73.79	0.41	MLT 11	1	5	
Actinomyces naeslundii		22.51	0.44	MAN 1	99	25	
Propionibacterium granulosum		3.70	0.16		99	90	
47712001	**Good identification**						**47712001**
Actinomyces naeslundii		100.0	0.47	ESL 12			
47712002	**Low discrimination**						**47712002**
					GRA		
Actinomyces naeslundii		70.39	0.44	ESL 12	99		
Fusobacterium mortiferum		25.64	0.25	GAL 96	1		
Bifidobacterium longum subsp. infantis		3.98	-.07		99		
47712003	**Good identification**						**47712003**
Actinomyces naeslundii		100.0	0.47	ESL 12			
47712020	**Good identification**						**47712020**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47712021	**Good identification**						**47712021**
Actinomyces naeslundii		100.0	0.47	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47712022	**Low discrimination**					**47712022**
					GRA	
Actinomyces naeslundii		82.45	0.44	ESL 12	99	
Prevotella oralis		9.28	0.02	GAL 96	1	
Bifidobacterium longum subsp. infantis		8.28	-.03		99	
47712023	**Good identification**					**47712023**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47712100	**Low discrimination**					**47712100**
					GRA	
Fusobacterium mortiferum		68.27	0.47	MLT 11	1	
Actinomyces naeslundii		25.41	0.52	MAN 1	99	
Bifidobacterium longum subsp. infantis		4.94	0.10		99	
47712101	**Good identification**					**47712101**
					GRA	
Actinomyces naeslundii		98.22	0.55	ESL 12	99	
Fusobacterium mortiferum		1.78	0.13	GAL 96	1	
47712103	**Good identification**					**47712103**
Actinomyces naeslundii		98.69	0.55	ESL 12		
Bifidobacterium longum subsp. infantis		1.31	-.07	GAL 96		
47712120	**Low discrimination**					**47712120**
Actinomyces naeslundii		66.56	0.52	ESL 12		
Bifidobacterium longum subsp. infantis		22.99	0.15	GAL 96		
Bifidobacterium breve		3.23	0.15			
47712121	**Good identification**					**47712121**
Actinomyces naeslundii		100.0	0.55	ESL 12		
47712123	**Good identification**					**47712123**
Actinomyces naeslundii		97.70	0.55	ESL 12		
Bifidobacterium longum subsp. infantis		2.30	-.03	GAL 96		
47712140	**Low discrimination**					**47712140**
					GRA	
Actinomyces naeslundii		76.42	0.39	ESL 12	99	
Bifidobacterium breve		11.82	0.10	GAL 96	99	
Fusobacterium mortiferum		11.75	0.13		1	
47712141	**Good identification**					**47712141**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47712143	**Good identification**					**47712143**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47712160	**Low discrimination**					**47712160**
Actinomyces naeslundii		78.43	0.39	ESL 12		
Bifidobacterium breve		21.57	0.15	GAL 96		
47712161	**Good identification**					**47712161**
Actinomyces naeslundii		100.0	0.42	ESL 12		
47712163	**Good identification**					**47712163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	ESL 12		
47712502	**Good identification**					**47712502**
Bifidobacterium adolescentis		97.98	0.43	GAL 99		
Actinomyces israelii		1.27	0.17	XYL 87		
Bifidobacterium longum subsp. infantis		0.75	-.07			
47712512	**Good identification**					**47712512**
Bifidobacterium adolescentis		100.0	0.45	GAL 99		
47713000	**Low discrimination**					**47713000**
					GRA	
Actinomyces naeslundii		75.69	0.67	ESL 12	99	
Fusobacterium mortiferum		15.51	0.44		1	
Actinomyces israelii		6.10	0.39		99	
47713001	**Very good identification**					**47713001**
Actinomyces naeslundii		100.0	0.70	ESL 12		
47713002	**Low discrimination**					**47713002**
Actinomyces naeslundii		67.34	0.67	ESL 12		
Bifidobacterium longum subsp. infantis		15.68	0.27			
Actinomyces israelii		8.14	0.43			
47713003	**Very good identification**					**47713003**
Actinomyces naeslundii		99.84	0.70	ESL 12		
Bifidobacterium longum subsp. infantis		0.16	-.07			
47713010	**Low discrimination**					**47713010**
Actinomyces naeslundii		77.50	0.46	ESL 12		
Actinomyces israelii		17.74	0.25	MLZ 5		
Bifidobacterium breve		4.77	0.10			
47713011	**Good identification**					**47713011**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47713013	**Good identification**					**47713013**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47713020	**Good identification**					**47713020**
					GRA	
Actinomyces naeslundii		90.97	0.67	ESL 12	99	
Bifidobacterium longum subsp. infantis		3.73	0.15		99	
Prevotella oralis		2.22	0.14		1	
47713021	**Very good identification**					**47713021**
Actinomyces naeslundii		100.0	0.70	ESL 12		
47713022	**Low discrimination**					**47713022**
					GRA	
Actinomyces naeslundii		50.66	0.67	ESL 12	99	
Prevotella oralis		23.52	0.36		1	
Bifidobacterium longum subsp. infantis		20.98	0.32		99	
47713023	**Very good identification**					**47713023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		99.41	0.70	ESL 12	99	
Prevotella oralis		0.31	0.02		1	
Bifidobacterium longum subsp. infantis		0.28	-.03		99	
47713030	**Acceptable identification**					**47713030**
Actinomyces naeslundii		90.14	0.46	ESL 12		
Bifidobacterium breve		9.86	0.15	MLZ 5		
47713031	**Good identification**					**47713031**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47713033	**Good identification**					**47713033**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47713040	**Low discrimination**					**47713040**
Actinomyces naeslundii		83.61	0.55	ESL 12		
Actinomyces israelii		14.85	0.32			
Bifidobacterium breve		1.53	0.10			
47713041	**Very good identification**					**47713041**
Actinomyces naeslundii		100.0	0.58	ESL 12		
47713042	**Low discrimination**					**47713042**
Actinomyces naeslundii		66.78	0.55	ESL 12		
Actinomyces israelii		17.79	0.35			
Bifidobacterium breve		12.39	0.27			
47713043	**Very good identification**					**47713043**
Actinomyces naeslundii		100.0	0.58	ESL 12		
47713051	**Good identification**					**47713051**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47713053	**Good identification**					**47713053**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47713060	**Good identification**					**47713060**
Actinomyces naeslundii		95.42	0.55	ESL 12		
Bifidobacterium breve		3.11	0.15			
Clostridium perfringens		1.47	0.01			
47713061	**Very good identification**					**47713061**
Actinomyces naeslundii		100.0	0.58	ESL 12		
47713062	**Low discrimination**					**47713062**
					GRA	
Actinomyces naeslundii		71.58	0.55	ESL 12	99	
Bifidobacterium breve		23.60	0.32		99	
Prevotella oralis		1.90	0.02		1	
47713063	**Very good identification**					**47713063**
Actinomyces naeslundii		100.0	0.58	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47713071	**Good identification**					**47713071**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47713073	**Good identification**					**47713073**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47713100	**Low discrimination**					**47713100**
					GRA	
Actinomyces naeslundii		44.92	0.75	ESL 12	99	
Bifidobacterium longum subsp. infantis		35.99	0.44		99	
Fusobacterium mortiferum		7.54	0.50		1	
47713101	**Very good identification**					**47713101**
					GRA	
Actinomyces naeslundii		99.20	0.78	ESL 12	99	
Bifidobacterium longum subsp. infantis		0.54	0.10		99	
Fusobacterium mortiferum		0.11	0.16		1	
47713102	**Low discrimination**					**47713102**
Bifidobacterium longum subsp. infantis		66.47	0.62	MAN 1		
Bifidobacterium adolescentis		13.13	0.63			
Bifidobacterium breve		9.34	0.62			
47713103	**Good identification**					**47713103**
Actinomyces naeslundii		93.02	0.78	ESL 12		
Bifidobacterium longum subsp. infantis		5.08	0.27			
Bifidobacterium adolescentis		1.00	0.29			
47713110	**Low discrimination**					**47713110**
Bifidobacterium breve		53.19	0.44	CEL 99		
Actinomyces naeslundii		24.86	0.53	BGL 91		
Bifidobacterium adolescentis		9.33	0.31			
47713111	**Very good identification**					**47713111**
Actinomyces naeslundii		98.58	0.56	ESL 12		
Bifidobacterium breve		1.42	0.10	MLZ 5		
47713112	**Low discrimination**					**47713112**
Bifidobacterium adolescentis		60.02	0.65	XYL 87		
Bifidobacterium breve		34.94	0.62	ARA 87		
Bifidobacterium longum subsp. infantis		2.51	0.27			
47713113	**Low discrimination**					**47713113**
Actinomyces naeslundii		71.11	0.56	ESL 12		
Bifidobacterium adolescentis		17.79	0.31	MLZ 5		
Bifidobacterium breve		10.36	0.27			
47713120	**Low discrimination**					**47713120**
Bifidobacterium longum subsp. infantis		46.96	0.49	MAN 1		
Actinomyces naeslundii		32.97	0.75	BGL 91		
Bifidobacterium breve		6.60	0.49			
47713121	**Very good identification**					**47713121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		98.70	0.78	ESL 12		
Bifidobacterium longum subsp. infantis		0.95	0.15			
Bifidobacterium breve		0.13	0.15			
47713122	**Low discrimination**					**47713122**
					GRA	
Bifidobacterium longum subsp. infantis		70.04	0.66	MAN 1	99	
Prevotella oralis		12.43	0.56		1	
Bifidobacterium breve		9.84	0.66		99	
47713123	**Low discrimination**					**47713123**
					GRA	
Actinomyces naeslundii		88.45	0.78	ESL 12	99	
Bifidobacterium longum subsp. infantis		8.58	0.32		99	
Prevotella oralis		1.52	0.22		1	
47713130	**Low discrimination**					**47713130**
Bifidobacterium breve		73.19	0.49	CEL 99		
Actinomyces naeslundii		19.24	0.53	BGL 91		
Bifidobacterium longum subsp. infantis		5.26	0.15			
47713131	**Very good identification**					**47713131**
Actinomyces naeslundii		97.50	0.56	ESL 12		
Bifidobacterium breve		2.50	0.15	MLZ 5		
47713132	**Low discrimination**					**47713132**
Bifidobacterium breve		88.28	0.66	CEL 99		
Bifidobacterium longum subsp. infantis		6.35	0.32			
Actinomyces naeslundii		2.30	0.53			
47713133	**Low discrimination**					**47713133**
Actinomyces naeslundii		78.27	0.56	ESL 12		
Bifidobacterium breve		20.27	0.32	MLZ 5		
Bifidobacterium longum subsp. infantis		1.46	-.03			
47713140	**Low discrimination**					**47713140**
Actinomyces naeslundii		51.16	0.62	ESL 12		
Bifidobacterium breve		32.65	0.44			
Actinomyces israelii		12.77	0.43			
47713141	**Very good identification**					**47713141**
Actinomyces naeslundii		99.57	0.65	ESL 12		
Bifidobacterium breve		0.43	0.10			
47713142	**Low discrimination**					**47713142**
Bifidobacterium breve		68.42	0.62	CEL 99		
Bifidobacterium adolescentis		11.88	0.48			
Actinomyces naeslundii		10.61	0.62			
47713143	**Good identification**					**47713143**
Actinomyces naeslundii		94.87	0.65	ESL 12		
Bifidobacterium breve		4.12	0.27			
Bifidobacterium adolescentis		0.72	0.13			
47713150	**Low discrimination**					**47713150**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		86.01	0.44	CEL 99		
Actinomyces naeslundii		7.10	0.40	BGL 91		
Actinomyces israelii		5.03	0.29			
47713151	**Acceptable identification**					**47713151**
Actinomyces naeslundii		92.45	0.43	ESL 12		
Bifidobacterium breve		7.55	0.10	MLZ 5		
47713152	**Low discrimination**					**47713152**
Bifidobacterium breve		81.31	0.62	CEL 99		
Bifidobacterium adolescentis		17.26	0.49			
Actinomyces israelii		0.71	0.32			
47713153	**Low discrimination**					**47713153**
Actinomyces naeslundii		49.98	0.43	ESL 12		
Bifidobacterium breve		41.26	0.27	MLZ 5		
Bifidobacterium adolescentis		8.76	0.15			
47713160	**Low discrimination**					**47713160**
Bifidobacterium breve		49.37	0.49	CEL 99		
Actinomyces naeslundii		43.52	0.62	BGL 91		
Bifidobacterium longum subsp. infantis		3.55	0.15			
47713161	**Very good identification**					**47713161**
Actinomyces naeslundii		99.24	0.65	ESL 12		
Bifidobacterium breve		0.76	0.15			
47713162	**Low discrimination**					**47713162**
Bifidobacterium breve		84.57	0.66	CEL 99		
Actinomyces naeslundii		7.37	0.62			
Bifidobacterium longum subsp. infantis		6.08	0.32			
47713163	**Good identification**					**47713163**
Actinomyces naeslundii		92.35	0.65	ESL 12		
Bifidobacterium breve		7.13	0.32			
Bifidobacterium longum subsp. infantis		0.51	-.03			
47713170	**Good identification**					**47713170**
Bifidobacterium breve		95.57	0.49	CEL 99		
Actinomyces naeslundii		4.43	0.40	BGL 91		
47713171	**Low discrimination**					**47713171**
Actinomyces naeslundii		87.33	0.43	ESL 12		
Bifidobacterium breve		12.67	0.15	MLZ 5		
47713172	**Very good identification**					**47713172**
Bifidobacterium breve		99.35	0.66	CEL 99		
Actinomyces naeslundii		0.46	0.40			
Bifidobacterium adolescentis		0.12	0.15			
47713201	**Good identification**					**47713201**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Actinomyces naeslundii	100.0	0.36	ESL 12			
47713203	**Good identification**						**47713203**
	Actinomyces naeslundii	100.0	0.36	ESL 12			
47713221	**Good identification**						**47713221**
	Actinomyces naeslundii	100.0	0.36	ESL 12			
47713223	**Good identification**						**47713223**
	Actinomyces naeslundii	100.0	0.36	ESL 12			
47713301	**Good identification**						**47713301**
	Actinomyces naeslundii	100.0	0.44	ESL 12			
47713303	**Acceptable identification**						**47713303**
	Actinomyces naeslundii	94.83	0.44	ESL 12			
	Bifidobacterium longum subsp. infantis	5.18	-.07	RHA 1			
47713320	**Low discrimination**						**47713320**
					SPO	GRA	
	Clostridium ramosum	58.44	0.35	TRE 94	92	50	
	Prevotella oralis	24.61	0.28	CEL 99	1	1	
	Bifidobacterium longum subsp. infantis	6.14	0.15		1	99	
47713321	**Acceptable identification**						**47713321**
					SPO		
	Actinomyces naeslundii	91.64	0.44	ESL 12	1		
	Clostridium ramosum	8.36	0.01	RHA 1	92		
47713322	**Low discrimination**						**47713322**
					SPO	GRA	
	Prevotella oralis	58.87	0.50	MAN 1	1	1	
	Clostridium ramosum	29.43	0.45		92	50	
	Bifidobacterium longum subsp. infantis	7.82	0.32		1	99	
47713323	**Low discrimination**						**47713323**
					SPO	GRA	
	Actinomyces naeslundii	45.62	0.44	ESL 12	1	99	
	Prevotella oralis	33.31	0.16	RHA 1	1	1	
	Clostridium ramosum	16.65	0.11		92	50	
47713400	**Low discrimination**						**47713400**
	Actinomyces israelii	83.31	0.38	CEL 93			
	Actinomyces naeslundii	12.76	0.33	XYL 92			
	Bifidobacterium adolescentis	3.93	0.09				
47713401	**Good identification**						**47713401**
	Actinomyces naeslundii	100.0	0.36	ESL 12			
47713402	**Low discrimination**						**47713402**
	Bifidobacterium adolescentis	73.43	0.43	XYL 87			
	Actinomyces israelii	23.60	0.41	LAC 99			
	Actinomyces naeslundii	2.41	0.33				
47713403	**Low discrimination**						**47713403**
	Actinomyces naeslundii	82.98	0.36	ESL 12			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		17.03	0.09	ARA 1		
47713412	**Good identification**					**47713412**
Bifidobacterium adolescentis		96.22	0.45	XYL 87		
Actinomyces israelii		3.78	0.27	LAC 99		
47713421	**Good identification**					**47713421**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47713423	**Good identification**					**47713423**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47713500	**Low discrimination**					**47713500**
Bifidobacterium adolescentis		47.66	0.43	XYL 87		
Actinomyces israelii		40.84	0.48	BGL 99		
Actinomyces naeslundii		4.45	0.41			
47713501	**Low discrimination**					**47713501**
Actinomyces naeslundii		88.19	0.44	ESL 12		
Bifidobacterium adolescentis		6.36	0.09	ARA 1		
Actinomyces israelii		5.45	0.14			
47713502	**Very good identification**					**47713502**
Bifidobacterium adolescentis		97.74	0.77	XYL 87		
Actinomyces israelii		1.27	0.51			
Bifidobacterium longum subsp. infantis		0.75	0.27			
47713503	**Low discrimination**					**47713503**
Bifidobacterium adolescentis		86.15	0.43	XYL 87		
Actinomyces naeslundii		12.07	0.44	URE 1		
Actinomyces israelii		1.12	0.17			
47713510	**Low discrimination**					**47713510**
Bifidobacterium adolescentis		88.37	0.45	XYL 87		
Actinomyces israelii		9.26	0.34	BGL 99		
Bifidobacterium longum subsp. longum		1.61	0.13			
47713512	**Very good identification**					**47713512**
Bifidobacterium adolescentis		99.70	0.79	XYL 87		
Actinomyces israelii		0.16	0.37			
Bifidobacterium breve		0.09	0.27			
47713513	**Good identification**					**47713513**
Bifidobacterium adolescentis		100.0	0.45	XYL 87		
47713521	**Good identification**					**47713521**
Actinomyces naeslundii		100.0	0.44	ESL 12		
47713523	**Low discrimination**					**47713523**
Actinomyces naeslundii		85.54	0.44	ESL 12		
Bifidobacterium longum subsp. infantis		8.30	-.03	ARA 1		
Bifidobacterium adolescentis		6.17	0.09			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47713532	**Low discrimination**					**47713532**
Bifidobacterium adolescentis		75.46	0.45	XYL 87		
Bifidobacterium breve		11.67	0.32	NAG 1		
Bifidobacterium longum subsp. longum		5.49	0.24			
47713540	**Low discrimination**					**47713540**
Actinomyces israelii		68.88	0.41	CEL 93		
Bifidobacterium adolescentis		25.54	0.28	XYL 92		
Actinomyces naeslundii		3.41	0.28			
47713542	**Very good identification**					**47713542**
Bifidobacterium adolescentis		95.10	0.62	XYL 87		
Actinomyces israelii		3.89	0.44	SOR 11		
Bifidobacterium breve		0.83	0.27			
47713552	**Very good identification**					**47713552**
Bifidobacterium adolescentis		98.80	0.63	XYL 87		
Bifidobacterium breve		0.70	0.27	SOR 11		
Actinomyces israelii		0.49	0.30			
47713702	**Low discrimination**					**47713702**
Bifidobacterium adolescentis		85.55	0.43	XYL 87		
Actinomyces israelii		8.28	0.32	RHA 1		
Eubacterium contortum		3.07	0.10			
47713712	**Low discrimination**					**47713712**
Bifidobacterium adolescentis		89.39	0.45	XYL 87		
Clostridium clostridioforme		9.55	0.25	RHA 1		
Actinomyces israelii		1.06	0.18			
47714000	**Good identification**					**47714000**
					GRA	
Fusobacterium mortiferum		98.18	0.56	MLT 11	1	
Bifidobacterium longum subsp. infantis		1.82	0.09	XYL 7	99	
47714100	**Low discrimination**					**47714100**
					GRA	
Fusobacterium mortiferum		54.80	0.62	MLT 11	1	
Bifidobacterium longum subsp. infantis		43.06	0.43	XYL 7	99	
Eubacterium contortum		1.77	0.24		99	
47714102	**Very good identification**					**47714102**
					GRA	
Bifidobacterium longum subsp. infantis		96.16	0.60	GAL 99	99	
Bifidobacterium adolescentis		1.92	0.45		99	
Fusobacterium mortiferum		1.35	0.46		1	
47714112	**Low discrimination**					**47714112**
Bifidobacterium adolescentis		59.73	0.46	GAL 99		
Bifidobacterium longum subsp. infantis		24.75	0.26	ARA 87		
Clostridium clostridioforme		15.17	0.33			
47714120	**Acceptable identification**					**47714120**
					GRA	
Bifidobacterium longum subsp. infantis		92.81	0.47	GAL 99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		3.23	0.20	BGL 91	1		
Prevotella oralis		1.32	0.19		1		
47714122	**Good identification**						**47714122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		93.88	0.64	GAL 99	1	99	
Prevotella oralis		2.52	0.41		1	1	
Bacteroides fragilis		2.43	0.32		99	1	
47714212	**Good identification**						**47714212**
Clostridium clostridioforme		100.0	0.39	TRE 88			
47714232	**Good identification**						**47714232**
Clostridium clostridioforme		97.87	0.39	TRE 88			
Parabacteroides distasonis		2.13	0.16	ARA 94			
47714300	**Very good identification**						**47714300**
					GRA		
Eubacterium contortum		99.30	0.58	GAL 99	99		
Fusobacterium mortiferum		0.32	0.28		1		
Bifidobacterium longum subsp. infantis		0.25	0.09		99		
47714302	**Low discrimination**						**47714302**
Eubacterium contortum		76.27	0.45	GAL 99			
Bifidobacterium longum subsp. infantis		10.88	0.26				
Clostridium clostridioforme		10.73	0.36				
47714312	**Good identification**						**47714312**
					GRA		
Clostridium clostridioforme		97.31	0.48	TRE 88	50		
Eubacterium contortum		1.53	0.11	ARA 94	99		
Parabacteroides distasonis		0.63	0.16		1		
47714320	**Low discrimination**						**47714320**
					GRA		
Eubacterium contortum		54.75	0.45	GAL 99	99		
Bacteroides vulgatus		42.62	0.36		1		
Bifidobacterium longum subsp. infantis		1.37	0.13		99		
47714322	**Low discrimination**						**47714322**
					GRA		
Parabacteroides distasonis		46.73	0.55	GAL 99	1		
Prevotella oralis		17.58	0.35		1		
Bifidobacterium longum subsp. infantis		15.46	0.30		99		
47714332	**Low discrimination**						**47714332**
Clostridium clostridioforme		60.42	0.48	TRE 88			
Parabacteroides distasonis		38.90	0.50	ARA 94			
Prevotella oralis		0.28	0.00				
47714412	**Acceptable identification**						**47714412**
Clostridium clostridioforme		94.62	0.44	TRE 88			
Bifidobacterium adolescentis		5.38	0.26	RHA 88			
47714432	**Good identification**						**47714432**
Clostridium clostridioforme		100.0	0.44	TRE 88			
47714502	**Low discrimination**						**47714502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		79.79	0.59	GAL 99		
Clostridium clostridioforme		12.73	0.42			
Bifidobacterium longum subsp. infantis		6.04	0.26			
47714512	**Low discrimination**					**47714512**
Bifidobacterium adolescentis		62.45	0.60	GAL 99		
Clostridium clostridioforme		37.13	0.53			
Bifidobacterium longum subsp. longum		0.42	0.21			
47714520	**Acceptable identification**					**47714520**
					GRA	
Bacteroides vulgatus		93.91	0.39	GAL 99	1	
Bifidobacterium longum subsp. longum		2.95	0.28	RHA 90	99	
Bifidobacterium longum subsp. infantis		2.05	0.13		99	
47714522	**Low discrimination**					**47714522**
					GRA	
Clostridium clostridioforme		37.44	0.42	TRE 88	50	
Bifidobacterium longum subsp. infantis		31.58	0.30	RHA 88	99	
Parabacteroides distasonis		19.50	0.43		1	
47714532	**Low discrimination**					**47714532**
					GRA	
Clostridium clostridioforme		88.86	0.53	TRE 88	50	
Parabacteroides distasonis		5.47	0.38	RHA 88	1	
Bifidobacterium longum subsp. longum		3.99	0.32		99	
47714542	**Good identification**					**47714542**
Bifidobacterium adolescentis		97.74	0.43	GAL 99		
Clostridium clostridioforme		1.27	0.08	SOR 11		
Actinomyces israelii		0.99	0.15			
47714552	**Good identification**					**47714552**
Bifidobacterium adolescentis		95.37	0.45	GAL 99		
Clostridium clostridioforme		4.63	0.19	SOR 11		
47714602	**Good identification**					**47714602**
Clostridium clostridioforme		98.76	0.48	TRE 88		
Eubacterium contortum		1.24	0.10			
47714612	**Very good identification**					**47714612**
Clostridium clostridioforme		100.0	0.59	TRE 88		
47714622	**Good identification**					**47714622**
Clostridium clostridioforme		99.29	0.48	TRE 88		
Parabacteroides distasonis		0.72	0.17			
47714632	**Very good identification**					**47714632**
Clostridium clostridioforme		99.92	0.59	TRE 88		
Parabacteroides distasonis		0.09	0.12			
47714700	**Very good identification**					**47714700**
					GRA	
Eubacterium contortum		97.34	0.57	GAL 99	99	
Bacteroides vulgatus		2.19	0.21		1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.47	0.23		50		
47714702	**Low discrimination**						**47714702**
Clostridium clostridioforme		72.37	0.57	TRE 88			
Eubacterium contortum		26.85	0.44				
Bifidobacterium adolescentis		0.63	0.25				
47714712	**Very good identification**						**47714712**
Clostridium clostridioforme		99.66	0.68	TRE 88			
Bifidobacterium adolescentis		0.23	0.26				
Eubacterium contortum		0.08	0.10				
47714720	**Acceptable identification**						**47714720**
					GRA		
Bacteroides vulgatus		92.44	0.56	GAL 99	1		
Eubacterium contortum		7.31	0.44		99		
Clostridium clostridioforme		0.20	0.23		50		
47714722	**Low discrimination**						**47714722**
					GRA		
Clostridium clostridioforme		75.00	0.57	TRE 88	50		
Parabacteroides distasonis		15.98	0.51		1		
Eubacterium contortum		4.91	0.31		99		
47714732	**Very good identification**						**47714732**
Clostridium clostridioforme		97.54	0.68	TRE 88			
Parabacteroides distasonis		2.46	0.46				
47715000	**Low discrimination**						**47715000**
					GRA		
Bifidobacterium longum subsp. infantis		50.73	0.43	LAC 99	99		
Fusobacterium mortiferum		41.51	0.59	BGL 91	1		
Actinomyces israelii		4.45	0.45		99		
47715002	**Good identification**						**47715002**
					GRA		
Bifidobacterium longum subsp. infantis		94.22	0.60	LAC 99	99		
Bifidobacterium adolescentis		1.88	0.45		99		
Prevotella oralis		1.49	0.33		1		
47715012	**Low discrimination**						**47715012**
Bifidobacterium adolescentis		57.86	0.46	LAC 99			
Bifidobacterium longum subsp. infantis		23.98	0.26	ARA 87			
Clostridium clostridioforme		13.17	0.32				
47715020	**Low discrimination**						**47715020**
					GRA		
Bifidobacterium longum subsp. infantis		86.22	0.47	LAC 99	99		
Prevotella oralis		7.77	0.33	BGL 91	1		
Bacteroides vulgatus		3.00	0.20		1		
47715022	**Low discrimination**						**47715022**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		82.75	0.64	LAC 99	1	99	
Prevotella oralis		14.02	0.55		1	1	
Bacteroides fragilis		2.14	0.32		99	1	
47715040	**Low discrimination**						**47715040**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		64.99	0.38	CEL 93	99	
Bifidobacterium longum subsp. infantis		19.25	0.09		99	
Fusobacterium mortiferum		15.75	0.25		1	
47715100	**Good identification**					**47715100**
					GRA	
Bifidobacterium longum subsp. infantis		93.38	0.77	BGL 91	99	
Eubacterium contortum		3.83	0.58		99	
Fusobacterium mortiferum		1.80	0.65		1	
47715101	**Acceptable identification**					**47715101**
					GRA	
Bifidobacterium longum subsp. infantis		92.51	0.43	URE 1	99	
Eubacterium contortum		3.79	0.24	BGL 91	99	
Fusobacterium mortiferum		1.78	0.31		1	
47715102	**Very good identification**					**47715102**
					GRA	
Bifidobacterium longum subsp. infantis		97.57	0.94		99	
Bifidobacterium adolescentis		1.95	0.79		99	
Prevotella oralis		0.25	0.53		1	
47715103	**Very good identification**					**47715103**
					GRA	
Bifidobacterium longum subsp. infantis		97.47	0.60	URE 1	99	
Bifidobacterium adolescentis		1.94	0.45		99	
Prevotella oralis		0.24	0.19		1	
47715110	**Low discrimination**					**47715110**
Bifidobacterium longum subsp. infantis		71.98	0.43	MLZ 1		
Bifidobacterium adolescentis		17.74	0.46	BGL 91		
Actinomyces israelii		3.77	0.41			
47715112	**Low discrimination**					**47715112**
Bifidobacterium adolescentis		69.76	0.80	ARA 87		
Bifidobacterium longum subsp. infantis		28.91	0.60			
Clostridium clostridioforme		0.54	0.41			
47715113	**Low discrimination**					**47715113**
Bifidobacterium adolescentis		70.32	0.46	ARA 87		
Bifidobacterium longum subsp. infantis		29.14	0.26	URE 1		
Clostridium clostridioforme		0.54	0.07			
47715120	**Good identification**					**47715120**
					GRA	
Bifidobacterium longum subsp. infantis		93.35	0.81	BGL 91	99	
Bacteroides vulgatus		3.25	0.54		1	
Prevotella oralis		1.33	0.53		1	
47715121	**Acceptable identification**					**47715121**
					GRA	
Bifidobacterium longum subsp. infantis		92.86	0.47	URE 1	99	
Bacteroides vulgatus		3.23	0.20	BGL 91	1	
Prevotella oralis		1.32	0.19		1	
47715122	**Good identification**					**47715122**
					CAT GRA	
Bifidobacterium longum subsp. infantis		93.94	0.99		1 99	
Prevotella oralis		2.52	0.75		1 1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides fragilis		2.43	0.66		99	1	
47715123	**Good identification**						**47715123**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		93.90	0.64	URE 1	1	99	
Prevotella oralis		2.52	0.41		1	1	
Bacteroides fragilis		2.43	0.32		99	1	
47715130	**Low discrimination**						**47715130**
					GRA		
Bifidobacterium longum subsp. infantis		85.51	0.47	MLZ 1	99		
Parabacteroides distasonis		4.74	0.42	BGL 91	1		
Bacteroides vulgatus		2.98	0.20		1		
47715132	**Low discrimination**						**47715132**
					GRA		
Bifidobacterium longum subsp. infantis		61.07	0.64	MLZ 1	99		
Parabacteroides distasonis		33.13	0.76		1		
Prevotella oralis		1.64	0.41		1		
47715140	**Low discrimination**						**47715140**
Bifidobacterium longum subsp. infantis		80.33	0.43	SOR 1			
Actinomyces israelii		10.96	0.48	BGL 91			
Eubacterium contortum		3.29	0.24				
47715142	**Low discrimination**						**47715142**
Bifidobacterium longum subsp. infantis		78.52	0.60	SOR 1			
Bifidobacterium adolescentis		19.16	0.63				
Actinomyces israelii		1.59	0.51				
47715152	**Very good identification**						**47715152**
Bifidobacterium adolescentis		95.11	0.65	ARA 87			
Bifidobacterium longum subsp. infantis		3.22	0.26	SOR 11			
Actinomyces israelii		0.96	0.37				
47715160	**Acceptable identification**						**47715160**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		90.95	0.47	SOR 1	99	1	
Bacteroides vulgatus		3.17	0.20	BGL 91	1	3	
Capnocytophaga ochracea		1.40	0.20		1	90	
47715162	**Good identification**						**47715162**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		92.52	0.64	SOR 1	1	99	
Prevotella oralis		2.48	0.41		1	1	
Bacteroides fragilis		2.40	0.32		99	1	
47715200	**Very good identification**						**47715200**
					GRA		
Eubacterium contortum		99.16	0.58	LAC 99	99		
Bifidobacterium longum subsp. infantis		0.25	0.09		99		
Fusobacterium mortiferum		0.20	0.25		1		
47715202	**Low discrimination**						**47715202**
Eubacterium contortum		71.99	0.45	LAC 99			
Bifidobacterium longum subsp. infantis		10.27	0.26				
Clostridium clostridioforme		9.08	0.36				
47715212	**Good identification**						**47715212**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Clostridium clostridioforme		96.67	0.47	TRE 88	50	
Eubacterium contortum		1.70	0.11	ARA 94	99	
Parabacteroides distasonis		0.70	0.16		1	
47715220	**Low discrimination**					**47715220**
					GRA	
Eubacterium contortum		52.44	0.45	LAC 99	99	
Bacteroides vulgatus		40.81	0.36		1	
Prevotella oralis		5.03	0.26		1	
47715222	**Low discrimination**					**47715222**
					GRA	
Prevotella oralis		57.68	0.48	XYL 11	1	
Parabacteroides distasonis		24.26	0.55	LAC 94	1	
Bifidobacterium longum subsp. infantis		8.03	0.30		99	
47715232	**Low discrimination**					**47715232**
Clostridium clostridioforme		56.88	0.47	TRE 88		
Parabacteroides distasonis		40.86	0.50	ARA 94		
Prevotella oralis		1.82	0.14			
47715300	**Excellent identification**					**47715300**
					GRA	
Eubacterium contortum		99.58	0.92		99	
Bifidobacterium longum subsp. infantis		0.25	0.43		99	
Bacteroides vulgatus		0.14	0.36		1	
47715301	**Very good identification**					**47715301**
					GRA	
Eubacterium contortum		99.61	0.58	URE 1	99	
Bifidobacterium longum subsp. infantis		0.25	0.09		99	
Bacteroides vulgatus		0.14	0.02		1	
47715302	**Low discrimination**					**47715302**
					GRA	
Eubacterium contortum		85.22	0.80		99	
Bifidobacterium longum subsp. infantis		12.16	0.60		99	
Prevotella oralis		1.29	0.47		1	
47715303	**Low discrimination**					**47715303**
					GRA	
Eubacterium contortum		85.27	0.45	URE 1	99	
Bifidobacterium longum subsp. infantis		12.17	0.26		99	
Prevotella oralis		1.29	0.13		1	
47715310	**Very good identification**					**47715310**
Eubacterium contortum		99.05	0.58	MLZ 1		
Clostridium clostridioforme		0.34	0.22			
Bifidobacterium longum subsp. infantis		0.25	0.09			
47715312	**Low discrimination**					**47715312**
					GRA	
Clostridium clostridioforme		50.00	0.56	TRE 88	50	
Eubacterium contortum		26.00	0.45	ARA 94	99	
Parabacteroides distasonis		10.73	0.50		1	
47715320	**Low discrimination**					**47715320**
					GRA	
Eubacterium contortum		54.74	0.80		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides vulgatus		42.61	0.71		1	
Bifidobacterium longum subsp. infantis		1.37	0.47		99	
47715321	**Low discrimination**					**47715321**
					GRA	
Eubacterium contortum		54.75	0.45	URE 1	99	
Bacteroides vulgatus		42.62	0.36		1	
Bifidobacterium longum subsp. infantis		1.37	0.13		99	
47715322	**Low discrimination**					**47715322**
					GRA	
Parabacteroides distasonis		50.93	0.89		1	
Prevotella oralis		19.17	0.69		1	
Bifidobacterium longum subsp. infantis		16.85	0.64		99	
47715323	**Low discrimination**					**47715323**
					GRA	
Parabacteroides distasonis		50.97	0.55	URE 1	1	
Prevotella oralis		19.18	0.35		1	
Bifidobacterium longum subsp. infantis		16.87	0.30		99	
47715330	**Low discrimination**					**47715330**
					GRA	
Eubacterium contortum		44.43	0.45	MLZ 1	99	
Bacteroides vulgatus		34.58	0.36		1	
Parabacteroides distasonis		18.34	0.50		1	
47715332	**Good identification**					**47715332**
Parabacteroides distasonis		93.88	0.84			
Clostridium clostridioforme		4.42	0.56			
Prevotella oralis		0.66	0.35			
47715333	**Acceptable identification**					**47715333**
Parabacteroides distasonis		93.93	0.50	URE 1		
Clostridium clostridioforme		4.42	0.22			
Prevotella oralis		0.66	0.00			
47715340	**Very good identification**					**47715340**
Eubacterium contortum		99.36	0.58	SOR 1		
Actinomyces israelii		0.25	0.29			
Bifidobacterium longum subsp. infantis		0.25	0.09			
47715342	**Low discrimination**					**47715342**
Eubacterium contortum		81.56	0.45	SOR 1		
Bifidobacterium longum subsp. infantis		11.63	0.26			
Bifidobacterium adolescentis		2.84	0.29			
47715360	**Low discrimination**					**47715360**
					GRA	
Eubacterium contortum		54.75	0.45	SOR 1	99	
Bacteroides vulgatus		42.62	0.36		1	
Bifidobacterium longum subsp. infantis		1.37	0.13		99	
47715362	**Low discrimination**					**47715362**
					GRA	
Parabacteroides distasonis		50.86	0.55	SOR 1	1	
Prevotella oralis		19.14	0.35		1	
Bifidobacterium longum subsp. infantis		16.83	0.30		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47715372	**Acceptable identification**					**47715372**
Parabacteroides distasonis		93.93	0.50	SOR 1		
Clostridium clostridioforme		4.42	0.22			
Prevotella oralis		0.66	0.00			
47715400	**Low discrimination**					**47715400**
					SPO	
Actinomyces israelii		32.04	0.43	CEL 93	1	
Clostridium butyricum		30.31	0.22		99	
Eubacterium contortum		14.98	0.23		1	
47715402	**Low discrimination**					**47715402**
Bifidobacterium adolescentis		75.20	0.59	LAC 99		
Clostridium clostridioforme		10.75	0.41			
Actinomyces israelii		6.00	0.46			
47715412	**Low discrimination**					**47715412**
Bifidobacterium adolescentis		64.52	0.60	LAC 99		
Clostridium clostridioforme		34.37	0.52			
Actinomyces israelii		0.63	0.32			
47715420	**Acceptable identification**					**47715420**
					GRA	
Bacteroides vulgatus		92.81	0.39	LAC 99	1	
Bifidobacterium longum subsp. longum		2.92	0.28	RHA 90	99	
Bifidobacterium longum subsp. infantis		2.03	0.13		99	
47715422	**Low discrimination**					**47715422**
					GRA	
Clostridium clostridioforme		32.56	0.41	TRE 88	50	
Bifidobacterium longum subsp. infantis		30.65	0.30	RHA 88	99	
Parabacteroides distasonis		18.93	0.43		1	
47715432	**Low discrimination**					**47715432**
					GRA	
Clostridium clostridioforme		87.73	0.52	TRE 88	50	
Parabacteroides distasonis		6.03	0.38	RHA 88	1	
Bifidobacterium longum subsp. longum		4.40	0.32		99	
47715440	**Low discrimination**					**47715440**
					SPO	
Actinomyces israelii		78.62	0.36	CEL 93	1	
Clostridium butyricum		16.63	0.04		99	
Bifidobacterium adolescentis		4.75	0.09		1	
47715442	**Low discrimination**					**47715442**
					SPO	
Bifidobacterium adolescentis		78.35	0.43	LAC 99	1	
Actinomyces israelii		19.66	0.39	SOR 11	1	
Clostridium butyricum		1.08	-.03		99	
47715452	**Acceptable identification**					**47715452**
Bifidobacterium adolescentis		93.09	0.45	LAC 99		
Clostridium clostridioforme		4.05	0.18	SOR 11		
Actinomyces israelii		2.86	0.25			
47715500	**Low discrimination**					**47715500**
					SPO	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		36.55	0.57	RHA 99	1	
Clostridium butyricum		17.93	0.45		99	
Bifidobacterium adolescentis		14.86	0.59		1	
47715502	**Good identification**					**47715502**
Bifidobacterium adolescentis		90.98	0.93			
Bifidobacterium longum subsp. infantis		6.89	0.60			
Bifidobacterium longum subsp. longum		0.44	0.52			
47715503	**Good identification**					**47715503**
Bifidobacterium adolescentis		91.10	0.59	URE 1		
Bifidobacterium longum subsp. infantis		6.90	0.26			
Bifidobacterium longum subsp. longum		0.44	0.17			
47715510	**Low discrimination**					**47715510**
					SPO	
Bifidobacterium adolescentis		55.92	0.60	BGL 99	1	
Bifidobacterium longum subsp. longum		36.95	0.56		1	
Clostridium butyricum		2.91	0.24		99	
47715512	**Very good identification**					**47715512**
Bifidobacterium adolescentis		97.43	0.94			
Clostridium clostridioforme		1.76	0.61			
Bifidobacterium longum subsp. longum		0.65	0.56			
47715513	**Very good identification**					**47715513**
Bifidobacterium adolescentis		97.59	0.60	URE 1		
Clostridium clostridioforme		1.76	0.27			
Bifidobacterium longum subsp. longum		0.65	0.21			
47715520	**Good identification**					**47715520**
					GRA	
Bacteroides vulgatus		93.87	0.73	RHA 90	1	
Bifidobacterium longum subsp. longum		2.95	0.62		99	
Bifidobacterium longum subsp. infantis		2.05	0.47		99	
47715521	**Acceptable identification**					**47715521**
					GRA	
Bacteroides vulgatus		94.15	0.39	RHA 90	1	
Bifidobacterium longum subsp. longum		2.96	0.28	URE 1	99	
Bifidobacterium longum subsp. infantis		2.06	0.13		99	
47715522	**Low discrimination**					**47715522**
					GRA	
Bifidobacterium longum subsp. infantis		49.31	0.64	ARA 1	99	
Parabacteroides distasonis		30.45	0.77		1	
Bifidobacterium longum subsp. longum		7.02	0.62		99	
47715530	**Low discrimination**					**47715530**
					GRA	
Bifidobacterium longum subsp. longum		81.82	0.66	SAL 1	99	
Bacteroides vulgatus		15.44	0.39		1	
Parabacteroides distasonis		1.13	0.38		1	
47715532	**Low discrimination**					**47715532**
					GRA	
Parabacteroides distasonis		39.34	0.72		1	
Bifidobacterium longum subsp. longum		28.69	0.66		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		19.37	0.61		50	
47715533	**Low discrimination**					**47715533**
					GRA	
Parabacteroides distasonis		39.56	0.38	URE 1	1	
Bifidobacterium longum subsp. longum		28.85	0.32		99	
Clostridium clostridioforme		19.47	0.27		50	
47715540	**Low discrimination**					**47715540**
					SPO	
Bifidobacterium adolescentis		34.86	0.43	SOR 11	1	
Clostridium butyricum		29.60	0.27	BGL 99	99	
Actinomyces israelii		23.33	0.47		1	
47715542	**Very good identification**					**47715542**
Bifidobacterium adolescentis		97.93	0.77	SOR 11		
Actinomyces israelii		0.99	0.50			
Bifidobacterium longum subsp. infantis		0.61	0.26			
47715543	**Good identification**					**47715543**
Bifidobacterium adolescentis		99.00	0.43	SOR 11		
Actinomyces israelii		1.00	0.15	URE 1		
47715550	**Low discrimination**					**47715550**
Bifidobacterium adolescentis		84.51	0.45	SOR 11		
Actinomyces israelii		6.91	0.32	BGL 99		
Bifidobacterium longum subsp. longum		4.56	0.21			
47715552	**Very good identification**					**47715552**
Bifidobacterium adolescentis		99.66	0.79	SOR 11		
Clostridium clostridioforme		0.15	0.27			
Actinomyces israelii		0.12	0.35			
47715553	**Good identification**					**47715553**
Bifidobacterium adolescentis		100.0	0.45	SOR 11		
47715560	**Acceptable identification**					**47715560**
					GRA	
Bacteroides vulgatus		91.58	0.39	RHA 90	1	
Bifidobacterium longum subsp. longum		2.88	0.28	SOR 1	99	
Bifidobacterium longum subsp. infantis		2.00	0.13		99	
47715572	**Low discrimination**					**47715572**
					GRA	
Bifidobacterium adolescentis		58.88	0.45	NAG 1	99	
Parabacteroides distasonis		17.44	0.38	SOR 11	1	
Bifidobacterium longum subsp. longum		12.72	0.32		99	
47715600	**Very good identification**					**47715600**
					GRA	
Eubacterium contortum		97.17	0.57	LAC 99	99	
Bacteroides vulgatus		2.19	0.21		1	
Clostridium clostridioforme		0.42	0.22		50	
47715602	**Low discrimination**					**47715602**
Clostridium clostridioforme		69.84	0.56	TRE 88		
Eubacterium contortum		28.92	0.44			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Bifidobacterium adolescentis	0.67	0.25			
47715612	**Very good identification**					**47715612**
	Clostridium clostridioforme	99.62	0.67	TRE 88		
	Bifidobacterium adolescentis	0.26	0.26			
	Eubacterium contortum	0.09	0.10			
47715620	**Acceptable identification**					**47715620**
					GRA	
	Bacteroides vulgatus	92.46	0.56	LAC 99	1	
	Eubacterium contortum	7.32	0.44		99	
	Clostridium clostridioforme	0.18	0.22		50	
47715622	**Low discrimination**					**47715622**
					GRA	
	Clostridium clostridioforme	72.18	0.56	TRE 88	50	
	Parabacteroides distasonis	17.16	0.51		1	
	Eubacterium contortum	5.28	0.31		99	
47715632	**Very good identification**					**47715632**
	Clostridium clostridioforme	97.27	0.67	TRE 88		
	Parabacteroides distasonis	2.73	0.46			
47715700	**Very good identification**					**47715700**
					GRA	
	Eubacterium contortum	97.75	0.91		99	
	Bacteroides vulgatus	2.20	0.56		1	
	Actinomyces israelii	0.01	0.34		99	
47715701	**Very good identification**					**47715701**
					GRA	
	Eubacterium contortum	97.80	0.57	URE 1	99	
	Bacteroides vulgatus	2.20	0.21		1	
47715702	**Low discrimination**					**47715702**
	Eubacterium contortum	89.68	0.78			
	Clostridium clostridioforme	7.32	0.65			
	Bifidobacterium adolescentis	2.09	0.59			
47715703	**Acceptable identification**					**47715703**
	Eubacterium contortum	90.03	0.44	URE 1		
	Clostridium clostridioforme	7.35	0.31			
	Bifidobacterium adolescentis	2.10	0.25			
47715710	**Good identification**					**47715710**
					GRA	
	Eubacterium contortum	91.01	0.57	MLZ 1	99	
	Clostridium clostridioforme	5.97	0.42		50	
	Bacteroides vulgatus	2.05	0.21		1	
47715712	**Low discrimination**					**47715712**
	Clostridium clostridioforme	89.87	0.76	TRE 88		
	Bifidobacterium adolescentis	6.87	0.60			
	Eubacterium contortum	2.44	0.44			
47715713	**Low discrimination**					**47715713**
	Clostridium clostridioforme	89.93	0.42	TRE 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		6.88	0.26	URE 1		
Eubacterium contortum		2.44	0.10			
47715720	**Good identification**					**47715720**
					GRA	
Bacteroides vulgatus		92.62	0.90		1	
Eubacterium contortum		7.33	0.78		99	
Parabacteroides distasonis		0.04	0.51		1	
47715721	**Acceptable identification**					**47715721**
					GRA	
Bacteroides vulgatus		92.63	0.56	URE 1	1	
Eubacterium contortum		7.33	0.44		99	
Parabacteroides distasonis		0.04	0.17		1	
47715722	**Low discrimination**					**47715722**
					GRA	
Parabacteroides distasonis		58.47	0.85		1	
Eubacterium contortum		17.97	0.65		99	
Bacteroides vulgatus		13.00	0.56		1	
47715723	**Low discrimination**					**47715723**
					GRA	
Parabacteroides distasonis		58.59	0.51	URE 1	1	
Eubacterium contortum		18.01	0.31		99	
Bacteroides vulgatus		13.03	0.21		1	
47715730	**Low discrimination**					**47715730**
					GRA	
Bacteroides vulgatus		87.80	0.56	MLZ 1	1	
Eubacterium contortum		6.95	0.44		99	
Clostridium clostridioforme		2.58	0.42		50	
47715732	**Low discrimination**					**47715732**
					GRA	
Clostridium clostridioforme		54.14	0.76	TRE 88	50	
Parabacteroides distasonis		44.99	0.81		1	
Eubacterium contortum		0.26	0.31		99	
47715733	**Low discrimination**					**47715733**
Clostridium clostridioforme		54.61	0.42	TRE 88		
Parabacteroides distasonis		45.39	0.46	URE 1		
47715740	**Very good identification**					**47715740**
					GRA	
Eubacterium contortum		97.38	0.57	SOR 1	99	
Bacteroides vulgatus		2.19	0.21		1	
Actinomyces israelii		0.25	0.27		99	
47715742	**Low discrimination**					**47715742**
Eubacterium contortum		71.60	0.44	SOR 1		
Bifidobacterium adolescentis		20.39	0.43			
Clostridium clostridioforme		5.85	0.31			
47715752	**Low discrimination**					**47715752**
Clostridium clostridioforme		50.51	0.42	TRE 88		
Bifidobacterium adolescentis		47.26	0.45	SOR 1		
Eubacterium contortum		1.37	0.10			
47715760	**Acceptable identification**					**47715760**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Bacteroides vulgatus		92.63	0.56	SOR 1	1	
Eubacterium contortum		7.33	0.44		99	
Parabacteroides distasonis		0.04	0.17		1	
47715762	**Low discrimination**					**47715762**
					GRA	
Parabacteroides distasonis		58.14	0.51	SOR 1	1	
Eubacterium contortum		17.87	0.31		99	
Bacteroides vulgatus		12.93	0.21		1	
47715772	**Low discrimination**					**47715772**
					GRA	
Clostridium clostridioforme		54.09	0.42	TRE 88	50	
Parabacteroides distasonis		44.95	0.46	SOR 1	1	
Bifidobacterium adolescentis		0.51	0.11		99	
47716102	**Low discrimination**					**47716102**
Bifidobacterium adolescentis		55.24	0.43	GAL 99		
Bifidobacterium longum subsp. infantis		34.20	0.26	ARA 87		
Actinomyces israelii		10.08	0.37			
47716112	**Good identification**					**47716112**
Bifidobacterium adolescentis		97.82	0.45	GAL 99		
Actinomyces israelii		2.18	0.23	ARA 87		
47716502	**Very good identification**					**47716502**
Bifidobacterium adolescentis		97.63	0.57	GAL 99		
Actinomyces israelii		2.18	0.35			
Clostridium clostridioforme		0.19	0.08			
47716512	**Very good identification**					**47716512**
Bifidobacterium adolescentis		98.84	0.59	GAL 99		
Clostridium clostridioforme		0.73	0.19			
Actinomyces israelii		0.27	0.21			
47716542	**Acceptable identification**					**47716542**
Bifidobacterium adolescentis		93.44	0.42	GAL 99		
Actinomyces israelii		6.56	0.28	SOR 11		
47716552	**Good identification**					**47716552**
Bifidobacterium adolescentis		99.15	0.43	GAL 99		
Actinomyces israelii		0.85	0.14	SOR 11		
47717000	**Very good identification**					**47717000**
					GRA	
Actinomyces israelii		95.13	0.58	CEL 93	99	
Bifidobacterium longum subsp. infantis		1.93	0.09		99	
Fusobacterium mortiferum		1.58	0.25		1	
47717001	**Low discrimination**					**47717001**
Actinomyces naeslundii		61.81	0.36	ESL 12		
Actinomyces israelii		38.20	0.23	XYL 1		
47717002	**Low discrimination**					**47717002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		73.11	0.61	CEL 93		
Bifidobacterium adolescentis		16.19	0.43			
Bifidobacterium longum subsp. infantis		10.02	0.26			
47717003	**Low discrimination**					**47717003**
Actinomyces naeslundii		46.90	0.36	ESL 12		
Actinomyces israelii		43.48	0.27	XYL 1		
Bifidobacterium adolescentis		9.62	0.09			
47717010	**Good identification**					**47717010**
Actinomyces israelii		97.33	0.44	CEL 93		
Bifidobacterium adolescentis		2.67	0.11	MLZ 13		
47717012	**Low discrimination**					**47717012**
Bifidobacterium adolescentis		64.42	0.45	LAC 99		
Actinomyces israelii		35.58	0.47	ARA 87		
47717020	**Low discrimination**					**47717020**
Actinomyces israelii		55.93	0.37	CEL 93		
Bifidobacterium longum subsp. infantis		31.67	0.13	NAG 6		
Actinomyces naeslundii		9.55	0.33			
47717021	**Good identification**					**47717021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47717023	**Low discrimination**					**47717023**
Actinomyces naeslundii		81.58	0.36	ESL 12		
Bifidobacterium longum subsp. infantis		18.42	-.04	XYL 1		
47717040	**Good identification**					**47717040**
Actinomyces israelii		100.0	0.51	CEL 93		
47717042	**Acceptable identification**					**47717042**
Actinomyces israelii		93.43	0.54	CEL 93		
Bifidobacterium adolescentis		6.57	0.28			
47717050	**Good identification**					**47717050**
Actinomyces israelii		100.0	0.36	CEL 93		
47717052	**Low discrimination**					**47717052**
Actinomyces israelii		63.47	0.40	CEL 93		
Bifidobacterium adolescentis		36.53	0.29	MLZ 13		
47717100	**Low discrimination**					**47717100**
Actinomyces israelii		61.13	0.68	CEL 93		
Bifidobacterium longum subsp. infantis		30.76	0.43			
Bifidobacterium adolescentis		5.08	0.43			
47717101	**Low discrimination**					**47717101**
Actinomyces naeslundii		42.06	0.44	ESL 12		
Actinomyces israelii		36.53	0.34	XYL 1		
Bifidobacterium longum subsp. infantis		18.38	0.09			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47717102	**Low discrimination**						**47717102**
Bifidobacterium adolescentis		55.35	0.77	ARA 87			
Bifidobacterium longum subsp. infantis		34.26	0.60				
Actinomyces israelii		10.10	0.71				
47717103	**Low discrimination**						**47717103**
Bifidobacterium adolescentis		51.50	0.43	ARA 87			
Bifidobacterium longum subsp. infantis		31.88	0.26	URE 1			
Actinomyces israelii		9.40	0.37				
47717110	**Low discrimination**						**47717110**
Actinomyces israelii		58.03	0.54	CEL 93			
Bifidobacterium adolescentis		39.41	0.45	MLZ 13			
Bifidobacterium longum subsp. infantis		1.97	0.09				
47717112	**Very good identification**						**47717112**
Bifidobacterium adolescentis		97.21	0.79	ARA 87			
Actinomyces israelii		2.17	0.57				
Bifidobacterium longum subsp. infantis		0.50	0.26				
47717113	**Good identification**						**47717113**
Bifidobacterium adolescentis		97.82	0.45	ARA 87			
Actinomyces israelii		2.18	0.23	URE 1			
47717120	**Low discrimination**						**47717120**
					GRA		
Bifidobacterium longum subsp. infantis		86.50	0.47	MAN 1	99		
Actinomyces israelii		6.17	0.48	BGL 91	99		
Bacteroides vulgatus		3.01	0.20		1		
47717121	**Low discrimination**						**47717121**
Actinomyces naeslundii		54.58	0.44	ESL 12			
Bifidobacterium longum subsp. infantis		42.40	0.13	XYL 1			
Actinomyces israelii		3.03	0.13				
47717122	**Good identification**						**47717122**
					CAT	GRA	
Bifidobacterium longum subsp. infantis		91.97	0.64	MAN 1	1	99	
Prevotella oralis		2.47	0.41		1	1	
Bacteroides fragilis		2.38	0.32		99	1	
47717132	**Low discrimination**						**47717132**
					GRA		
Bifidobacterium adolescentis		35.32	0.45	ARA 87	99		
Bifidobacterium longum subsp. infantis		31.80	0.30	NAG 1	99		
Parabacteroides distasonis		17.25	0.42		1		
47717140	**Very good identification**						**47717140**
Actinomyces israelii		95.47	0.61	CEL 93			
Bifidobacterium adolescentis		2.52	0.28				
Bifidobacterium longum subsp. infantis		1.25	0.09				
47717142	**Low discrimination**						**47717142**
Bifidobacterium adolescentis		61.17	0.62	ARA 87			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		35.13	0.64	SOR 11		
Bifidobacterium longum subsp. infantis		3.09	0.26			
47717150	**Low discrimination**					**47717150**
					COC	
Actinomyces israelii		79.99	0.47	CEL 93	1	
Bifidobacterium adolescentis		17.26	0.29	MLZ 13	1	
Blautia producta		1.55	-.03		99	
47717152	**Good identification**					**47717152**
Bifidobacterium adolescentis		92.82	0.63	ARA 87		
Actinomyces israelii		6.52	0.50	SOR 11		
Bifidobacterium breve		0.66	0.27			
47717160	**Low discrimination**					**47717160**
Actinomyces israelii		67.49	0.40	CEL 93		
Bifidobacterium longum subsp. infantis		24.57	0.13	NAG 6		
Bifidobacterium breve		4.22	0.15			
47717202	**Low discrimination**					**47717202**
Actinomyces israelii		86.31	0.41	CEL 93		
Eubacterium contortum		11.13	0.11	RHA 7		
Bifidobacterium adolescentis		2.56	0.09			
47717300	**Very good identification**					**47717300**
Eubacterium contortum		96.05	0.58	MAN 1		
Actinomyces israelii		3.54	0.49			
Bifidobacterium longum subsp. infantis		0.24	0.09			
47717302	**Low discrimination**					**47717302**
Eubacterium contortum		58.35	0.45	MAN 1		
Actinomyces israelii		18.29	0.52			
Bifidobacterium adolescentis		13.45	0.43			
47717312	**Low discrimination**					**47717312**
Bifidobacterium adolescentis		77.77	0.45	RHA 1		
Actinomyces israelii		12.93	0.38	ARA 87		
Clostridium clostridioforme		5.36	0.22			
47717320	**Low discrimination**					**47717320**
					GRA	
Eubacterium contortum		54.27	0.45	MAN 1	99	
Bacteroides vulgatus		42.24	0.36		1	
Bifidobacterium longum subsp. infantis		1.36	0.13		99	
47717322	**Low discrimination**					**47717322**
					GRA	
Parabacteroides distasonis		49.84	0.55	MAN 1	1	
Prevotella oralis		18.76	0.35		1	
Bifidobacterium longum subsp. infantis		16.49	0.30		99	
47717332	**Acceptable identification**					**47717332**
					GRA	
Parabacteroides distasonis		92.70	0.50	MAN 1	1	
Clostridium clostridioforme		4.36	0.22		50	
Actinomyces israelii		0.67	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47717340	**Low discrimination**					**47717340**
Actinomyces israelii		58.67	0.42	CEL 93		
Eubacterium contortum		41.33	0.24	RHA 7		
47717342	**Low discrimination**					**47717342**
Actinomyces israelii		75.96	0.45	CEL 93		
Bifidobacterium adolescentis		17.75	0.28	RHA 7		
Eubacterium contortum		6.30	0.11			
47717400	**Very good identification**					**47717400**
					SPO	
Actinomyces israelii		95.60	0.56	CEL 93	1	
Bifidobacterium adolescentis		2.62	0.23		1	
Clostridium butyricum		1.77	0.05		99	
47717402	**Low discrimination**					**47717402**
					SPO	
Bifidobacterium adolescentis		64.24	0.57	LAC 99	1	
Actinomyces israelii		35.48	0.59		1	
Clostridium butyricum		0.17	-.02		99	
47717410	**Low discrimination**					**47717410**
Actinomyces israelii		81.67	0.42	CEL 93		
Bifidobacterium adolescentis		18.33	0.25	MLZ 13		
47717412	**Good identification**					**47717412**
Bifidobacterium adolescentis		92.96	0.59	LAC 99		
Actinomyces israelii		6.28	0.45			
Clostridium clostridioforme		0.61	0.18			
47717420	**Low discrimination**					**47717420**
					GRA	
Actinomyces israelii		75.74	0.36	CEL 93	99	
Bacteroides vulgatus		24.26	0.05	NAG 6	1	
47717422	**Low discrimination**					**47717422**
Actinomyces israelii		72.27	0.39	CEL 93		
Bifidobacterium adolescentis		20.71	0.23	NAG 6		
Clostridium clostridioforme		3.62	0.07			
47717440	**Good identification**					**47717440**
Actinomyces israelii		99.14	0.49	CEL 93		
Bifidobacterium adolescentis		0.86	0.08			
47717442	**Low discrimination**					**47717442**
Actinomyces israelii		63.47	0.52	CEL 93		
Bifidobacterium adolescentis		36.53	0.42			
47717450	**Low discrimination**					**47717450**
					COC	
Actinomyces israelii		85.76	0.35	CEL 93	1	
Blautia producta		8.12	-.03	MLZ 13	99	
Bifidobacterium adolescentis		6.12	0.09		1	
47717452	**Low discrimination**					**47717452**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		82.48	0.43	LAC 99		
Actinomyces israelii		17.52	0.38	SOR 11		
47717500	**Low discrimination**					**47717500**
					SPO	
Actinomyces israelii		54.76	0.66	CEL 93	1	
Bifidobacterium adolescentis		37.18	0.57		1	
Clostridium butyricum		6.09	0.29		99	
47717502	**Very good identification**					**47717502**
Bifidobacterium adolescentis		97.65	0.91			
Actinomyces israelii		2.18	0.69			
Bifidobacterium longum subsp. infantis		0.09	0.26			
47717503	**Very good identification**					**47717503**
Bifidobacterium adolescentis		97.82	0.57	URE 1		
Actinomyces israelii		2.18	0.35			
47717510	**Low discrimination**					**47717510**
Bifidobacterium adolescentis		82.42	0.59	BGL 99		
Actinomyces israelii		14.84	0.52			
Bifidobacterium dentium		1.36	0.10			
47717512	**Excellent identification**					**47717512**
Bifidobacterium adolescentis		99.53	0.93			
Actinomyces israelii		0.27	0.55			
Bifidobacterium dentium		0.17	0.27			
47717513	**Very good identification**					**47717513**
Bifidobacterium adolescentis		99.56	0.59	URE 1		
Actinomyces israelii		0.27	0.21			
Bifidobacterium dentium		0.17	-.07			
47717520	**Low discrimination**					**47717520**
					GRA	
Bacteroides vulgatus		82.01	0.39	MAN 1	1	
Actinomyces israelii		10.35	0.46	RHA 90	99	
Bifidobacterium longum subsp. longum		2.58	0.28		99	
47717522	**Low discrimination**					**47717522**
Bifidobacterium adolescentis		68.05	0.57	NAG 1		
Bifidobacterium longum subsp. infantis		11.19	0.30			
Actinomyces israelii		9.60	0.49			
47717532	**Low discrimination**					**47717532**
					GRA	
Bifidobacterium adolescentis		70.72	0.59	NAG 1	99	
Bifidobacterium dentium		20.93	0.32		99	
Parabacteroides distasonis		3.16	0.38		1	
47717540	**Low discrimination**					**47717540**
					COC	
Actinomyces israelii		73.67	0.59	CEL 93	1	
Bifidobacterium adolescentis		15.90	0.42		1	
Blautia producta		8.60	0.24		99	
47717542	**Good identification**					**47717542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Bifidobacterium adolescentis		93.35	0.76	SOR 11	1		
Actinomyces israelii		6.55	0.62		1		
Blautia producta		0.06	0.07		99		
47717543	**Acceptable identification**						**47717543**
Bifidobacterium adolescentis		93.44	0.42	SOR 11			
Actinomyces israelii		6.56	0.28	URE 1			
47717552	**Very good identification**						**47717552**
					COC		
Bifidobacterium adolescentis		98.99	0.77	SOR 11	1		
Actinomyces israelii		0.85	0.48		1		
Blautia producta		0.15	0.15		99		
47717553	**Good identification**						**47717553**
Bifidobacterium adolescentis		99.15	0.43	SOR 11			
Actinomyces israelii		0.85	0.14	URE 1			
47717560	**Low discrimination**						**47717560**
					COC	GRA	
Actinomyces israelii		67.70	0.39	CEL 93	1	99	
Bacteroides vulgatus		13.94	0.05	NAG 6	1	1	
Blautia producta		13.76	0.07		99	99	
47717562	**Low discrimination**						**47717562**
Bifidobacterium adolescentis		68.63	0.42	NAG 1			
Actinomyces israelii		30.45	0.42	SOR 11			
Bifidobacterium longum subsp. infantis		0.92	-.04				
47717572	**Acceptable identification**						**47717572**
Bifidobacterium adolescentis		91.34	0.43	NAG 1			
Actinomyces israelii		4.95	0.28	SOR 11			
Bifidobacterium dentium		2.21	-.03				
47717602	**Low discrimination**						**47717602**
Actinomyces israelii		59.41	0.40	CEL 93			
Clostridium clostridioforme		18.50	0.22	RHA 7			
Bifidobacterium adolescentis		14.44	0.23				
47717700	**Good identification**						**47717700**
					GRA		
Eubacterium contortum		93.95	0.57	MAN 1	99		
Actinomyces israelii		3.46	0.47		99		
Bacteroides vulgatus		2.12	0.21		1		
47717702	**Low discrimination**						**47717702**
Bifidobacterium adolescentis		57.30	0.57	RHA 1			
Eubacterium contortum		30.40	0.44				
Actinomyces israelii		9.53	0.50				
47717712	**Low discrimination**						**47717712**
Bifidobacterium adolescentis		84.09	0.59	RHA 1			
Clostridium clostridioforme		13.58	0.42				
Actinomyces israelii		1.71	0.36				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47717720	**Acceptable identification**					**47717720**
					GRA	
Bacteroides vulgatus		92.54	0.56	MAN 1	1	
Eubacterium contortum		7.32	0.44		99	
Actinomyces israelii		0.10	0.27		99	
47717722	**Low discrimination**					**47717722**
					GRA	
Parabacteroides distasonis		54.36	0.51	MAN 1	1	
Eubacterium contortum		16.71	0.31		99	
Bacteroides vulgatus		12.09	0.21		1	
47717732	**Low discrimination**					**47717732**
Clostridium clostridioforme		50.33	0.42	TRE 88		
Parabacteroides distasonis		41.83	0.46	MAN 1		
Bacteroides ovatus		3.35	0.12			
47717742	**Low discrimination**					**47717742**
					COC	
Bifidobacterium adolescentis		61.48	0.42	RHA 1	1	
Actinomyces israelii		32.16	0.43	SOR 11	1	
Blautia producta		3.70	0.07		99	
47717752	**Low discrimination**					**47717752**
					COC	
Bifidobacterium adolescentis		81.65	0.43	RHA 1	1	
Blautia producta		12.05	0.15	SOR 11	99	
Actinomyces israelii		5.22	0.29		1	
47720100	**Acceptable identification**					**47720100**
					GRA	
Fusobacterium mortiferum		93.85	0.38	MLT 11	1	
Actinomyces naeslundii		6.15	0.30	FRU 99	99	
47721000	**Low discrimination**					**47721000**
					GRA	
Fusobacterium mortiferum		50.29	0.35	MLT 11	1	
Actinomyces naeslundii		43.26	0.46	FRU 99	99	
Lactobacillus jensenii		6.45	0.10		99	
47721001	**Good identification**					**47721001**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47721002	**Low discrimination**					**47721002**
					GRA	
Actinomyces naeslundii		78.54	0.46	ESL 12	99	
Fusobacterium mortiferum		10.15	0.19	FRU 95	1	
Lactobacillus jensenii		7.80	0.07		99	
47721003	**Good identification**					**47721003**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47721020	**Low discrimination**					**47721020**
Actinomyces naeslundii		66.48	0.46	ESL 12		
Clostridium perfringens		33.52	0.18	FRU 95		
47721021	**Good identification**					**47721021**
Actinomyces naeslundii		100.0	0.49	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47721022	**Low discrimination**						**47721022**
					GRA		
Actinomyces naeslundii		53.20	0.46	ESL 12	99		
Prevotella oralis		29.95	0.16	FRU 95	1		
Clostridium perfringens		12.62	0.13		99		
47721023	**Good identification**						**47721023**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47721041	**Good identification**						**47721041**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47721043	**Good identification**						**47721043**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47721061	**Good identification**						**47721061**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47721063	**Good identification**						**47721063**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47721100	**Low discrimination**						**47721100**
					COC	GRA	
Actinomyces naeslundii		41.21	0.53	ESL 12	1	99	
Fusobacterium mortiferum		39.27	0.41	FRU 95	1	1	
Atopobium parvulum		11.53	0.21		99	99	
47721101	**Very good identification**						**47721101**
					SPO		
Actinomyces naeslundii		97.49	0.56	ESL 12	1		
Clostridium chauvoei		2.51	0.06	FRU 95	99		
47721103	**Very good identification**						**47721103**
Actinomyces naeslundii		98.96	0.56	ESL 12			
Bifidobacterium longum subsp. infantis		1.04	-.07	FRU 95			
47721121	**Very good identification**						**47721121**
Actinomyces naeslundii		100.0	0.56	ESL 12			
47721122	**Low discrimination**						**47721122**
					GRA		
Prevotella oralis		33.32	0.36	FRU 99	1		
Bifidobacterium longum subsp. infantis		29.72	0.32	TRE 6	99		
Parabacteroides distasonis		12.40	0.41		1		
47721123	**Very good identification**						**47721123**
					GRA		
Actinomyces naeslundii		96.20	0.56	ESL 12	99		
Prevotella oralis		2.01	0.02	FRU 95	1		
Bifidobacterium longum subsp. infantis		1.79	-.03		99		
47721132	**Low discrimination**						**47721132**
					GRA		
Parabacteroides distasonis		51.27	0.37	FRU 99	1		
Bifidobacterium breve		21.38	0.29	XYL 90	99		
Leptotrichia buccalis		16.70	0.07		1		
47721140	**Good identification**						**47721140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	ESL 12		
47721141	**Good identification**					**47721141**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47721142	**Low discrimination**					**47721142**
Actinomyces naeslundii		47.85	0.40	ESL 12		
Bifidobacterium breve		39.50	0.24	FRU 95		
Bifidobacterium adolescentis		8.39	0.12			
47721143	**Good identification**					**47721143**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47721161	**Good identification**					**47721161**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47721163	**Good identification**					**47721163**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47721322	**Low discrimination**					**47721322**
					SPO	
Parabacteroides distasonis		67.39	0.50	FRU 99	1	
Prevotella oralis		25.87	0.30	XYL 90	1	
Clostridium ramosum		5.66	0.19		92	
47721332	**Acceptable identification**					**47721332**
Parabacteroides distasonis		93.68	0.45	FRU 99		
Clostridium clostridioforme		6.32	0.19	XYL 90		
47721502	**Low discrimination**					**47721502**
Bifidobacterium adolescentis		89.10	0.42	FRU 99		
Bifidobacterium longum subsp. longum		3.93	0.17	XYL 87		
Clostridium clostridioforme		3.14	0.14			
47721512	**Low discrimination**					**47721512**
Bifidobacterium adolescentis		83.23	0.43	FRU 99		
Clostridium clostridioforme		10.92	0.25	XYL 87		
Bifidobacterium longum subsp. longum		5.11	0.21			
47721522	**Low discrimination**					**47721522**
					GRA	
Parabacteroides distasonis		62.16	0.38	FRU 99	1	
Bifidobacterium longum subsp. longum		25.97	0.27	XYL 90	99	
Clostridium clostridioforme		5.18	0.14		50	
47721712	**Good identification**					**47721712**
Clostridium clostridioforme		98.96	0.40	FRU 99		
Bifidobacterium adolescentis		1.04	0.09	XYL 91		
47721722	**Low discrimination**					**47721722**
Parabacteroides distasonis		83.07	0.46	FRU 99		
Clostridium clostridioforme		16.93	0.28	XYL 90		
47721732	**Low discrimination**					**47721732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		63.29	0.40	FRU 99		
Parabacteroides distasonis		36.71	0.41	XYL 91		
47722101	**Good identification**					**47722101**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47722103	**Good identification**					**47722103**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47722121	**Good identification**					**47722121**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47722123	**Good identification**					**47722123**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47723000	**Good identification**					**47723000**
Actinomyces naeslundii		97.09	0.50	ESL 12		
Actinomyces israelii		2.91	0.15	FRU 95		
47723001	**Good identification**					**47723001**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47723002	**Good identification**					**47723002**
Actinomyces naeslundii		95.70	0.50	ESL 12		
Actinomyces israelii		4.30	0.18	FRU 95		
47723003	**Good identification**					**47723003**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47723020	**Good identification**					**47723020**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47723021	**Good identification**					**47723021**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47723022	**Good identification**					**47723022**
Actinomyces naeslundii		100.0	0.50	ESL 12		
47723023	**Good identification**					**47723023**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47723040	**Good identification**					**47723040**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47723041	**Good identification**					**47723041**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47723042	**Good identification**					**47723042**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47723043	**Good identification**					**47723043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
47723060	**Good identification**					**47723060**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47723061	**Good identification**					**47723061**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47723062	**Good identification**					**47723062**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47723063	**Good identification**					**47723063**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47723100	**Very good identification**					**47723100**
Actinomyces naeslundii		95.96	0.58	ESL 12		
Actinomyces israelii		4.04	0.25	FRU 95		
47723101	**Very good identification**					**47723101**
Actinomyces naeslundii		100.0	0.61	ESL 12		
47723102	**Low discrimination**					**47723102**
Actinomyces naeslundii		79.89	0.58	ESL 12		
Bifidobacterium adolescentis		8.43	0.26	FRU 95		
Bifidobacterium breve		5.99	0.24			
47723103	**Very good identification**					**47723103**
Actinomyces naeslundii		100.0	0.61	ESL 12		
47723110	**Good identification**					**47723110**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47723111	**Good identification**					**47723111**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47723113	**Good identification**					**47723113**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47723120	**Low discrimination**					**47723120**
					SPO	
Actinomyces naeslundii		85.24	0.58	ESL 12	1	
Clostridium ramosum		13.63	0.19	FRU 95	92	
Bifidobacterium breve		1.12	0.12		1	
47723121	**Very good identification**					**47723121**
Actinomyces naeslundii		100.0	0.61	ESL 12		
47723122	**Low discrimination**					**47723122**
					SPO	
Actinomyces naeslundii		55.44	0.58	ESL 12	1	
Clostridium ramosum		35.47	0.30	FRU 95	92	
Bifidobacterium breve		7.40	0.29		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47723123	**Very good identification**					**47723123**
Actinomyces naeslundii		100.0	0.61	ESL 12		
47723130	**Low discrimination**					**47723130**
Actinomyces naeslundii		79.96	0.36	ESL 12		
Bifidobacterium breve		20.04	0.12	FRU 95		
47723131	**Good identification**					**47723131**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47723133	**Good identification**					**47723133**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47723140	**Acceptable identification**					**47723140**
Actinomyces naeslundii		91.50	0.45	ESL 12		
Actinomyces israelii		8.50	0.18	FRU 95		
47723141	**Good identification**					**47723141**
Actinomyces naeslundii		100.0	0.48	ESL 12		
47723142	**Low discrimination**					**47723142**
Actinomyces naeslundii		61.04	0.45	ESL 12		
Bifidobacterium breve		25.95	0.24	FRU 95		
Actinomyces israelii		8.50	0.21			
47723143	**Good identification**					**47723143**
Actinomyces naeslundii		100.0	0.48	ESL 12		
47723160	**Acceptable identification**					**47723160**
Actinomyces naeslundii		93.05	0.45	ESL 12		
Bifidobacterium breve		6.95	0.12	FRU 95		
47723161	**Good identification**					**47723161**
Actinomyces naeslundii		100.0	0.48	ESL 12		
47723162	**Low discrimination**					**47723162**
Actinomyces naeslundii		56.95	0.45	ESL 12		
Bifidobacterium breve		43.05	0.29	FRU 95		
47723163	**Good identification**					**47723163**
Actinomyces naeslundii		100.0	0.48	ESL 12		
47723502	**Acceptable identification**					**47723502**
Bifidobacterium adolescentis		91.95	0.40	FRU 99		
Actinomyces israelii		6.74	0.27	XYL 87		
Actinomyces naeslundii		1.32	0.24			
47723512	**Good identification**					**47723512**
Bifidobacterium adolescentis		98.34	0.42	FRU 99		
Bifidobacterium dentium		1.66	-.07	XYL 87		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47724612	**Good identification**					**47724612**
Clostridium clostridioforme		100.0	0.40	FRU 99		
47724632	**Good identification**					**47724632**
Clostridium clostridioforme		100.0	0.40	FRU 99		
47724702	**Good identification**					**47724702**
Clostridium clostridioforme		100.0	0.37	FRU 99		
47724712	**Good identification**					**47724712**
Clostridium clostridioforme		100.0	0.49	FRU 99		
47724722	**Low discrimination**					**47724722**
Clostridium clostridioforme		88.31	0.37	FRU 99		
Parabacteroides distasonis		11.69	0.28			
47724732	**Good identification**					**47724732**
Clostridium clostridioforme		98.46	0.49	FRU 99		
Parabacteroides distasonis		1.54	0.23			
47725102	**Low discrimination**					**47725102**
Bifidobacterium adolescentis		46.66	0.42	FRU 99		
Bifidobacterium longum subsp. infantis		35.44	0.26	ARA 87		
Actinomyces israelii		6.94	0.34			
47725112	**Low discrimination**					**47725112**
					GRA	
Bifidobacterium adolescentis		87.77	0.43	FRU 99	99	
Clostridium clostridioforme		7.43	0.22	ARA 87	50	
Parabacteroides distasonis		2.42	0.19		1	
47725122	**Low discrimination**					**47725122**
					GRA	
Parabacteroides distasonis		78.81	0.58	FRU 99	1	
Bifidobacterium longum subsp. infantis		17.17	0.30		99	
Prevotella oralis		2.91	0.20		1	
47725132	**Good identification**					**47725132**
					GRA	
Parabacteroides distasonis		96.27	0.53	FRU 99	1	
Clostridium clostridioforme		2.98	0.22		50	
Bifidobacterium longum subsp. infantis		0.39	-.04		99	
47725312	**Low discrimination**					**47725312**
					GRA	
Clostridium clostridioforme		86.99	0.36	FRU 99	50	
Parabacteroides distasonis		11.60	0.27	ARA 94	1	
Bifidobacterium adolescentis		1.42	0.09		99	
47725322	**Very good identification**					**47725322**
Parabacteroides distasonis		98.48	0.66	FRU 99		
Clostridium clostridioforme		0.88	0.25			
Prevotella oralis		0.52	0.14			
47725332	**Good identification**					**47725332**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		92.96	0.61	FRU 99			
Clostridium clostridioforme		7.04	0.36				
47725502	**Low discrimination**						**47725502**
					SPO		
Bifidobacterium adolescentis		82.76	0.56	FRU 99	1		
Clostridium butyricum		9.22	0.25		99		
Clostridium clostridioforme		4.40	0.31		66		
47725512	**Low discrimination**						**47725512**
Bifidobacterium adolescentis		80.82	0.57	FRU 99			
Clostridium clostridioforme		16.02	0.42				
Bifidobacterium longum subsp. longum		2.23	0.29				
47725520	**Low discrimination**						**47725520**
					GRA	SPO	
Bifidobacterium longum subsp. longum		53.48	0.35	SAL 1	99	1	
Bacteroides vulgatus		25.78	0.16	FRU 90	1	1	
Clostridium butyricum		16.86	0.14		99	99	
47725522	**Low discrimination**						**47725522**
					GRA		
Parabacteroides distasonis		78.49	0.54	FRU 99	1		
Bifidobacterium longum subsp. longum		10.93	0.35		99		
Clostridium clostridioforme		7.35	0.31		50		
47725530	**Good identification**						**47725530**
					GRA		
Bifidobacterium longum subsp. longum		96.05	0.39	SAL 1	99		
Parabacteroides distasonis		2.20	0.15	FRU 90	1		
Clostridium clostridioforme		1.75	0.08		50		
47725532	**Low discrimination**						**47725532**
					GRA		
Parabacteroides distasonis		43.71	0.49	FRU 99	1		
Clostridium clostridioforme		34.64	0.42		50		
Bifidobacterium longum subsp. longum		19.25	0.39		99		
47725542	**Low discrimination**						**47725542**
					SPO		
Bifidobacterium adolescentis		88.06	0.40	FRU 99	1		
Clostridium butyricum		6.90	0.07	SOR 11	99		
Actinomyces israelii		5.04	0.25		1		
47725552	**Good identification**						**47725552**
Bifidobacterium adolescentis		98.41	0.42	FRU 99			
Clostridium clostridioforme		1.59	0.08	SOR 11			
47725602	**Good identification**						**47725602**
Clostridium clostridioforme		100.0	0.37	FRU 99			
47725612	**Good identification**						**47725612**
Clostridium clostridioforme		100.0	0.48	FRU 99			
47725622	**Low discrimination**						**47725622**
Clostridium clostridioforme		87.13	0.37	FRU 99			
Parabacteroides distasonis		12.87	0.28				
47725632	**Good identification**						**47725632**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		98.28	0.48	FRU 99		
Parabacteroides distasonis		1.72	0.23			
47725702	**Acceptable identification**					**47725702**
					GRA	
Clostridium clostridioforme		91.26	0.46	FRU 99	50	
Parabacteroides distasonis		4.03	0.28		1	
Bifidobacterium adolescentis		2.36	0.22		99	
47725712	**Very good identification**					**47725712**
					GRA	
Clostridium clostridioforme		98.80	0.57	FRU 99	50	
Bifidobacterium adolescentis		0.69	0.23		99	
Parabacteroides distasonis		0.52	0.23		1	
47725722	**Low discrimination**					**47725722**
Parabacteroides distasonis		80.18	0.62	FRU 99		
Clostridium clostridioforme		18.36	0.46			
Bacteroides thetaiotaomicron		0.95	0.13			
47725732	**Low discrimination**					**47725732**
Clostridium clostridioforme		65.64	0.57	FRU 99		
Parabacteroides distasonis		33.89	0.58			
Bacteroides thetaiotaomicron		0.29	0.06			
47727002	**Good identification**					**47727002**
Actinomyces israelii		100.0	0.36	FRU 99		
47727100	**Acceptable identification**					**47727100**
					SPO	
Actinomyces israelii		94.08	0.44	FRU 99	1	
Clostridium butyricum		3.96	-.01	CEL 93	99	
Actinomyces naeslundii		1.96	0.24		1	
47727102	**Low discrimination**					**47727102**
Actinomyces israelii		50.38	0.47	FRU 99		
Bifidobacterium adolescentis		48.92	0.40	CEL 93		
Actinomyces naeslundii		0.70	0.24			
47727112	**Low discrimination**					**47727112**
Bifidobacterium adolescentis		87.51	0.42	FRU 99		
Actinomyces israelii		11.02	0.33	ARA 87		
Bifidobacterium dentium		1.48	-.07			
47727140	**Good identification**					**47727140**
Actinomyces israelii		100.0	0.37	FRU 99		
47727142	**Low discrimination**					**47727142**
Actinomyces israelii		76.42	0.40	FRU 99		
Bifidobacterium adolescentis		23.58	0.25	CEL 93		
47727500	**Low discrimination**					**47727500**
					SPO	
Actinomyces israelii		63.14	0.42	FRU 99	1	
Clostridium butyricum		29.27	0.15	CEL 93	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		7.60	0.20		1		
47727502	**Low discrimination**						**47727502**
					SPO		
Bifidobacterium adolescentis		87.40	0.54	FRU 99	1		
Actinomyces israelii		11.01	0.45		1		
Clostridium butyricum		1.32	0.08		99		
47727512	**Low discrimination**						**47727512**
Bifidobacterium adolescentis		78.89	0.56	FRU 99			
Bifidobacterium dentium		19.70	0.27				
Actinomyces israelii		1.21	0.31				
47727542	**Low discrimination**						**47727542**
					COC		
Bifidobacterium adolescentis		67.28	0.39	FRU 99	1		
Actinomyces israelii		26.65	0.38	SOR 11	1		
Blautia producta		6.07	0.07		99		
47727552	**Low discrimination**						**47727552**
					COC		
Bifidobacterium adolescentis		77.51	0.40	FRU 99	1		
Blautia producta		17.16	0.15	SOR 11	99		
Actinomyces israelii		3.75	0.24		1		
47730000	**Very good identification**						**47730000**
					GRA	CAT	
Fusobacterium mortiferum		98.14	0.66	MLT 11	1	5	
Actinomyces naeslundii		1.01	0.44		99	25	
Propionibacterium granulosum		0.66	0.26		99	90	
47730001	**Low discrimination**						**47730001**
					GRA		
Actinomyces naeslundii		60.51	0.47	ESL 12	99		
Fusobacterium mortiferum		39.49	0.32	GAL 96	1		
47730002	**Low discrimination**						**47730002**
					GRA	CAT	
Fusobacterium mortiferum		87.52	0.50	MLT 11	1	5	
Actinomyces naeslundii		8.13	0.44	BGL 10	99	25	
Propionibacterium granulosum		1.33	0.15		99	90	
47730003	**Acceptable identification**						**47730003**
					GRA		
Actinomyces naeslundii		93.24	0.47	ESL 12	99		
Fusobacterium mortiferum		6.76	0.16	GAL 96	1		
47730020	**Low discrimination**						**47730020**
					GRA		
Actinomyces naeslundii		35.98	0.44	ESL 12	99		
Fusobacterium mortiferum		35.22	0.32	GAL 96	1		
Clostridium perfringens		22.92	0.18		99		
47730021	**Good identification**						**47730021**
Actinomyces naeslundii		100.0	0.47	ESL 12			
47730022	**Low discrimination**						**47730022**
					GRA		
Actinomyces naeslundii		42.49	0.44	ESL 12	99		
Prevotella oralis		30.22	0.16	GAL 96	1		
Clostridium perfringens		12.74	0.13		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47730023	**Good identification**						**47730023**
Actinomyces naeslundii		100.0	0.47	ESL 12			
47730100	**Very good identification**						**47730100**
					COC	GRA	
Fusobacterium mortiferum		98.03	0.73	MLT 11	1	1	
Actinomyces naeslundii		1.23	0.52		1	99	
Atopobium parvulum		0.44	0.21		99	99	
47730101	**Low discrimination**						**47730101**
					GRA		
Actinomyces naeslundii		65.14	0.55	ESL 12	99		
Fusobacterium mortiferum		34.86	0.38	GAL 96	1		
47730102	**Low discrimination**						**47730102**
					GRA		
Fusobacterium mortiferum		68.02	0.56	MLT 11	1		
Bifidobacterium longum subsp. infantis		15.13	0.27	BGL 10	99		
Actinomyces naeslundii		7.70	0.52		99		
47730103	**Acceptable identification**						**47730103**
					GRA		
Actinomyces naeslundii		93.22	0.55	ESL 12	99		
Fusobacterium mortiferum		5.54	0.22	GAL 96	1		
Bifidobacterium longum subsp. infantis		1.23	-.07		99		
47730110	**Acceptable identification**						**47730110**
					GRA		
Fusobacterium mortiferum		91.36	0.38	MLT 11	1		
Actinomyces naeslundii		5.99	0.30	MLZ 1	99		
Leptotrichia buccalis		2.65	-.11		1		
47730121	**Good identification**						**47730121**
Actinomyces naeslundii		100.0	0.55	ESL 12			
47730122	**Low discrimination**						**47730122**
					GRA		
Prevotella oralis		33.52	0.36	TRE 6	1		
Bifidobacterium longum subsp. infantis		29.89	0.32	GAL 99	99		
Parabacteroides distasonis		12.47	0.41		1		
47730123	**Good identification**						**47730123**
					GRA		
Actinomyces naeslundii		95.25	0.55	ESL 12	99		
Prevotella oralis		2.51	0.02	GAL 96	1		
Bifidobacterium longum subsp. infantis		2.24	-.03		99		
47730132	**Low discrimination**						**47730132**
					GRA		
Parabacteroides distasonis		35.51	0.37	GAL 99	1		
Clostridium clostridioforme		32.34	0.30	XYL 90	50		
Bifidobacterium breve		14.81	0.29		99		
47730140	**Low discrimination**						**47730140**
					GRA		
Fusobacterium mortiferum		81.98	0.38	MLT 11	1		
Actinomyces naeslundii		18.02	0.39	SOR 1	99		
47730141	**Good identification**						**47730141**
Actinomyces naeslundii		100.0	0.42	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47730143	**Good identification**						**47730143**
Actinomyces naeslundii		100.0	0.42	ESL 12			
47730161	**Good identification**						**47730161**
Actinomyces naeslundii		100.0	0.42	ESL 12			
47730163	**Good identification**						**47730163**
Actinomyces naeslundii		100.0	0.42	ESL 12			
47730212	**Good identification**						**47730212**
Clostridium clostridioforme		100.0	0.36	XYL 91			
47730232	**Good identification**						**47730232**
Clostridium clostridioforme		98.50	0.36	XYL 91			
Parabacteroides distasonis		1.50	0.11	ARA 94			
47730300	**Good identification**						**47730300**
Fusobacterium mortiferum		100.0	0.38	MLT 11			
47730312	**Good identification**						**47730312**
Clostridium clostridioforme		99.11	0.45	XYL 91			
Parabacteroides distasonis		0.45	0.11	ARA 94			
Leptotrichia buccalis		0.44	-.11				
47730322	**Low discrimination**						**47730322**
Parabacteroides distasonis		57.50	0.50	GAL 99			
Prevotella oralis		22.08	0.30	XYL 90			
Clostridium clostridioforme		15.13	0.34				
47730332	**Low discrimination**						**47730332**
Clostridium clostridioforme		69.00	0.45	XYL 91			
Parabacteroides distasonis		31.00	0.45	ARA 94			
47730412	**Good identification**						**47730412**
Clostridium clostridioforme		99.23	0.42	XYL 91			
Bifidobacterium adolescentis		0.78	0.09	RHA 88			
47730432	**Good identification**						**47730432**
Clostridium clostridioforme		100.0	0.42	XYL 91			
47730500	**Low discrimination**						**47730500**
					GRA	SPO	
Fusobacterium mortiferum		82.39	0.38	MLT 11	1	1	
Clostridium butyricum		12.87	0.06	ARA 1	99	99	
Clostridium clostridioforme		2.55	0.05		50	66	
47730502	**Low discrimination**						**47730502**
					GRA		
Clostridium clostridioforme		51.69	0.40	XYL 91	50		
Bifidobacterium adolescentis		44.52	0.42	RHA 88	99		
Fusobacterium mortiferum		1.88	0.22		1		
47730512	**Low discrimination**						**47730512**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		81.10	0.51	XYL 91	50	
Bifidobacterium adolescentis		18.74	0.43	RHA 88	99	
Leptotrichia buccalis		0.17	-.11		1	
47730522	**Low discrimination**					**47730522**
Clostridium clostridioforme		70.60	0.40	XYL 91		
Parabacteroides distasonis		25.66	0.38	RHA 88		
Prevotella oralis		1.14	0.02			
47730532	**Good identification**					**47730532**
					GRA	
Clostridium clostridioforme		95.20	0.51	XYL 91	50	
Parabacteroides distasonis		4.09	0.33	RHA 88	1	
Bifidobacterium longum subsp. longum		0.49	0.13		99	
47730602	**Good identification**					**47730602**
Clostridium clostridioforme		100.0	0.45	XYL 91		
47730612	**Very good identification**					**47730612**
Clostridium clostridioforme		100.0	0.57	XYL 91		
47730622	**Good identification**					**47730622**
Clostridium clostridioforme		99.50	0.45	XYL 91		
Parabacteroides distasonis		0.50	0.12			
47730632	**Very good identification**					**47730632**
Clostridium clostridioforme		100.0	0.57	XYL 91		
47730702	**Good identification**					**47730702**
					GRA	
Clostridium clostridioforme		99.73	0.54	XYL 91	50	
Parabacteroides distasonis		0.15	0.12		1	
Bifidobacterium adolescentis		0.12	0.07		99	
47730712	**Very good identification**					**47730712**
Clostridium clostridioforme		99.97	0.66	XYL 91		
Bifidobacterium adolescentis		0.03	0.09			
47730722	**Low discrimination**					**47730722**
Clostridium clostridioforme		87.06	0.54	XYL 91		
Parabacteroides distasonis		12.95	0.46			
47730732	**Very good identification**					**47730732**
Clostridium clostridioforme		98.27	0.66	XYL 91		
Parabacteroides distasonis		1.73	0.41			
47731000	**Low discrimination**					**47731000**
					GRA	
Fusobacterium mortiferum		75.01	0.69	MLT 11	1	
Actinomyces naeslundii		12.38	0.67		99	
Lactobacillus jensenii		9.62	0.44		99	
47731001	**Very good identification**					**47731001**
					GRA	
Actinomyces naeslundii		95.60	0.70	ESL 12	99	
Fusobacterium mortiferum		3.90	0.35		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		0.50	0.10		99	
47731002	**Low discrimination**					**47731002**
					GRA	
Actinomyces naeslundii		33.90	0.67	ESL 12	99	
Fusobacterium mortiferum		22.82	0.53		1	
Lactobacillus jensenii		17.55	0.41		99	
47731003	**Very good identification**					**47731003**
					GRA	
Actinomyces naeslundii		99.05	0.70	ESL 12	99	
Fusobacterium mortiferum		0.45	0.19		1	
Lactobacillus jensenii		0.35	0.07		99	
47731010	**Low discrimination**					**47731010**
					GRA	
Lactobacillus jensenii		66.86	0.36	RAF 1	99	
Fusobacterium mortiferum		15.81	0.35	CEL 95	1	
Actinomyces naeslundii		13.59	0.46		99	
47731011	**Good identification**					**47731011**
Actinomyces naeslundii		96.79	0.49	ESL 12		
Lactobacillus jensenii		3.21	0.02	MLZ 5		
47731013	**Good identification**					**47731013**
					GRA	
Actinomyces naeslundii		96.63	0.49	ESL 12	99	
Leptotrichia buccalis		3.37	-.11	MLZ 5	1	
47731020	**Low discrimination**					**47731020**
Clostridium perfringens		63.69	0.52	LAC 88		
Actinomyces naeslundii		24.24	0.67			
Lactobacillus jensenii		4.71	0.34			
47731021	**Very good identification**					**47731021**
Actinomyces naeslundii		98.26	0.70	ESL 12		
Clostridium perfringens		1.74	0.18			
47731022	**Low discrimination**					**47731022**
					GRA	
Prevotella oralis		47.66	0.50	TRE 6	1	
Clostridium perfringens		20.09	0.47	LAC 94	99	
Actinomyces naeslundii		16.25	0.67		99	
47731023	**Very good identification**					**47731023**
					GRA	
Actinomyces naeslundii		97.01	0.70	ESL 12	99	
Prevotella oralis		1.92	0.16		1	
Clostridium perfringens		0.81	0.13		99	
47731031	**Good identification**					**47731031**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47731032	**Low discrimination**					**47731032**
					GRA	
Parabacteroides distasonis		24.47	0.37	XYL 90	1	
Clostridium clostridioforme		19.97	0.30	LAC 99	50	
Actinomyces naeslundii		13.85	0.46		99	
47731033	**Good identification**					**47731033**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.49	ESL 12			
47731040	**Low discrimination**						**47731040**
					GRA		
Actinomyces naeslundii		62.62	0.55	ESL 12	99		
Fusobacterium mortiferum		21.71	0.35		1		
Actinomyces israelii		7.39	0.29		99		
47731041	**Very good identification**						**47731041**
Actinomyces naeslundii		100.0	0.58	ESL 12			
47731042	**Low discrimination**						**47731042**
Actinomyces naeslundii		64.19	0.55	ESL 12			
Actinomyces israelii		11.35	0.32				
Bifidobacterium breve		7.94	0.24				
47731043	**Very good identification**						**47731043**
Actinomyces naeslundii		100.0	0.58	ESL 12			
47731051	**Good identification**						**47731051**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47731053	**Good identification**						**47731053**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47731060	**Low discrimination**						**47731060**
Clostridium perfringens		73.51	0.40	LAC 88			
Actinomyces naeslundii		25.92	0.55				
Bifidobacterium breve		0.56	0.12				
47731061	**Very good identification**						**47731061**
Actinomyces naeslundii		98.13	0.58	ESL 12			
Clostridium perfringens		1.87	0.06				
47731063	**Very good identification**						**47731063**
Actinomyces naeslundii		99.11	0.58	ESL 12			
Clostridium perfringens		0.89	0.00				
47731071	**Good identification**						**47731071**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47731073	**Good identification**						**47731073**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47731100	**Low discrimination**						**47731100**
					COC	GRA	
Fusobacterium mortiferum		56.92	0.76	MLT 11	1	1	
Atopobium parvulum		16.71	0.55		99	99	
Actinomyces naeslundii		11.46	0.75		1	99	
47731101	**Good identification**						**47731101**
					COC	GRA	
Actinomyces naeslundii		94.95	0.78	ESL 12	1	99	
Fusobacterium mortiferum		3.18	0.41		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		0.93	0.21		99	99	
47731102	**Low discrimination**						**47731102**
Bifidobacterium longum subsp. infantis		58.35	0.62	TRE 1			
Bifidobacterium adolescentis		9.39	0.62				
Actinomyces naeslundii		7.20	0.75				
47731103	**Good identification**						**47731103**
Actinomyces naeslundii		92.04	0.78	ESL 12			
Bifidobacterium longum subsp. infantis		5.02	0.27				
Bifidobacterium adolescentis		0.81	0.28				
47731111	**Very good identification**						**47731111**
					GRA		
Actinomyces naeslundii		98.79	0.56	ESL 12	99		
Leptotrichia buccalis		1.21	-.11	MLZ 5	1		
47731112	**Low discrimination**						**47731112**
					GRA		
Leptotrichia buccalis		78.16	0.58	CEL 99	1		
Bifidobacterium adolescentis		13.13	0.63		99		
Bifidobacterium breve		6.25	0.59		99		
47731120	**Low discrimination**						**47731120**
Clostridium perfringens		59.25	0.67				
Bifidobacterium longum subsp. infantis		12.47	0.49				
Actinomyces naeslundii		8.75	0.75				
47731121	**Good identification**						**47731121**
Actinomyces naeslundii		93.46	0.78	ESL 12			
Clostridium perfringens		4.26	0.33				
Bifidobacterium longum subsp. infantis		0.90	0.15				
47731122	**Low discrimination**						**47731122**
					GRA		
Prevotella oralis		36.50	0.70	TRE 6	1		
Bifidobacterium longum subsp. infantis		32.56	0.66		99		
Parabacteroides distasonis		13.58	0.76		1		
47731123	**Low discrimination**						**47731123**
					GRA		
Actinomyces naeslundii		77.53	0.78	ESL 12	99		
Prevotella oralis		8.43	0.36		1		
Bifidobacterium longum subsp. infantis		7.52	0.32		99		
47731130	**Low discrimination**						**47731130**
Bifidobacterium breve		39.86	0.46	CEL 99			
Clostridium perfringens		20.43	0.33	BGL 91			
Actinomyces naeslundii		15.72	0.53				
47731131	**Very good identification**						**47731131**
Actinomyces naeslundii		98.32	0.56	ESL 12			
Bifidobacterium breve		1.68	0.12	MLZ 5			
47731132	**Low discrimination**						**47731132**
					GRA		
Parabacteroides distasonis		52.46	0.71	XYL 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		21.88	0.63		99	
Leptotrichia buccalis		17.09	0.41		1	
47731133	**Low discrimination**					**47731133**
					GRA	
Actinomyces naeslundii		56.42	0.56	ESL 12	99	
Parabacteroides distasonis		23.36	0.37	MLZ 5	1	
Bifidobacterium breve		9.74	0.29		99	
47731140	**Low discrimination**					**47731140**
					GRA	
Actinomyces naeslundii		44.09	0.62	ESL 12	99	
Bifidobacterium breve		18.75	0.41		99	
Fusobacterium mortiferum		12.53	0.41		1	
47731141	**Very good identification**					**47731141**
Actinomyces naeslundii		100.0	0.65	ESL 12		
47731142	**Low discrimination**					**47731142**
Bifidobacterium breve		56.73	0.59	CEL 99		
Actinomyces naeslundii		13.19	0.62			
Bifidobacterium adolescentis		12.04	0.46			
47731143	**Very good identification**					**47731143**
Actinomyces naeslundii		96.32	0.65	ESL 12		
Bifidobacterium breve		2.79	0.24			
Bifidobacterium adolescentis		0.59	0.12			
47731150	**Low discrimination**					**47731150**
Bifidobacterium breve		81.88	0.41	CEL 99		
Actinomyces naeslundii		10.13	0.40	BGL 91		
Actinomyces israelii		4.77	0.26			
47731151	**Good identification**					**47731151**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47731152	**Low discrimination**					**47731152**
					GRA	
Bifidobacterium breve		70.98	0.59	CEL 99	99	
Bifidobacterium adolescentis		18.42	0.48		99	
Leptotrichia buccalis		8.96	0.24		1	
47731153	**Low discrimination**					**47731153**
Actinomyces naeslundii		56.74	0.43	ESL 12		
Bifidobacterium breve		31.22	0.24	MLZ 5		
Bifidobacterium adolescentis		8.10	0.13			
47731160	**Low discrimination**					**47731160**
Clostridium perfringens		77.57	0.55			
Actinomyces naeslundii		10.62	0.62			
Bifidobacterium breve		8.03	0.46			
47731161	**Good identification**					**47731161**
Actinomyces naeslundii		94.85	0.65	ESL 12		
Clostridium perfringens		4.67	0.21			

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		0.48	0.12			
47731162	**Low discrimination**					**47731162**
Bifidobacterium breve		51.60	0.63	CEL 99		
Clostridium perfringens		23.20	0.49			
Actinomyces naeslundii		6.75	0.62			
47731163	**Good identification**					**47731163**
Actinomyces naeslundii		92.05	0.65	ESL 12		
Bifidobacterium breve		4.74	0.29			
Clostridium perfringens		2.13	0.15			
47731170	**Low discrimination**					**47731170**
Bifidobacterium breve		85.68	0.46	CEL 99		
Clostridium perfringens		8.36	0.21	BGL 91		
Actinomyces naeslundii		5.96	0.40			
47731171	**Acceptable identification**					**47731171**
Actinomyces naeslundii		91.18	0.43	ESL 12		
Bifidobacterium breve		8.82	0.12	MLZ 5		
47731172	**Very good identification**					**47731172**
					GRA	
Bifidobacterium breve		95.42	0.63	CEL 99	99	
Parabacteroides distasonis		2.31	0.37		1	
Leptotrichia buccalis		0.75	0.07		1	
47731173	**Low discrimination**					**47731173**
Actinomyces naeslundii		50.55	0.43	ESL 12		
Bifidobacterium breve		49.45	0.29	MLZ 5		
47731200	**Low discrimination**					**47731200**
					GRA	
Fusobacterium mortiferum		70.62	0.35	MLT 11	1	
Actinomyces naeslundii		11.66	0.33	RHA 1	99	
Lactobacillus jensenii		9.05	0.10		99	
47731201	**Good identification**					**47731201**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47731203	**Good identification**					**47731203**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47731212	**Good identification**					**47731212**
Clostridium clostridioforme		99.01	0.44	XYL 91		
Parabacteroides distasonis		0.50	0.11	ARA 94		
Leptotrichia buccalis		0.49	-.11			
47731221	**Good identification**					**47731221**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47731222	**Low discrimination**					**47731222**
Prevotella oralis		63.51	0.43	TRE 6		
Parabacteroides distasonis		26.18	0.50	LAC 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		6.17	0.33				
47731223	**Low discrimination**						**47731223**
					GRA		
Actinomyces naeslundii		45.80	0.36	ESL 12	99		
Prevotella oralis		38.38	0.09	RHA 1	1		
Parabacteroides distasonis		15.82	0.15		1		
47731232	**Low discrimination**						**47731232**
Clostridium clostridioforme		65.61	0.44	XYL 91			
Parabacteroides distasonis		32.89	0.45	ARA 94			
Prevotella oralis		1.50	0.09				
47731300	**Low discrimination**						**47731300**
					GRA		
Fusobacterium mortiferum		34.41	0.41	MLT 11	1		
Eubacterium contortum		18.43	0.24	RHA 1	99		
Prevotella oralis		13.15	0.20		1		
47731301	**Good identification**						**47731301**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47731302	**Low discrimination**						**47731302**
Prevotella oralis		40.99	0.42	TRE 6			
Parabacteroides distasonis		20.49	0.50	NAG 95			
Clostridium clostridioforme		16.17	0.42				
47731303	**Low discrimination**						**47731303**
					GRA		
Actinomyces naeslundii		66.02	0.44	ESL 12	99		
Prevotella oralis		16.03	0.08	RHA 1	1		
Parabacteroides distasonis		8.01	0.15		1		
47731312	**Low discrimination**						**47731312**
Clostridium clostridioforme		74.56	0.53	XYL 91			
Parabacteroides distasonis		11.17	0.45	ARA 94			
Leptotrichia buccalis		10.91	0.24				
47731320	**Low discrimination**						**47731320**
					SPO		
Clostridium ramosum		35.57	0.43	CEL 99	92		
Prevotella oralis		34.24	0.42		1		
Parabacteroides distasonis		17.12	0.50		1		
47731321	**Low discrimination**						**47731321**
					SPO	GRA	
Actinomyces naeslundii		61.86	0.44	ESL 12	1	99	
Clostridium ramosum		15.61	0.09	RHA 1	92	50	
Prevotella oralis		15.02	0.08		1	1	
47731322	**Low discrimination**						**47731322**
					SPO		
Parabacteroides distasonis		67.12	0.84	XYL 90	1		
Prevotella oralis		25.77	0.64		1		
Clostridium ramosum		5.64	0.53		92		
47731323	**Low discrimination**						**47731323**
					SPO		
Parabacteroides distasonis		63.81	0.50	XYL 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		24.50	0.30	URE 1	1	
Clostridium ramosum		5.36	0.19		92	
47731330	**Low discrimination**					**47731330**
					SPO	
Parabacteroides distasonis		86.38	0.45	XYL 90	1	
Clostridium clostridioforme		5.83	0.19	BGL 99	66	
Clostridium ramosum		3.37	0.09		92	
47731332	**Good identification**					**47731332**
Parabacteroides distasonis		92.69	0.79	XYL 90		
Clostridium clostridioforme		6.25	0.53			
Prevotella oralis		0.67	0.30			
47731333	**Acceptable identification**					**47731333**
Parabacteroides distasonis		93.68	0.45	XYL 90		
Clostridium clostridioforme		6.32	0.19	URE 1		
47731362	**Low discrimination**					**47731362**
					SPO	
Parabacteroides distasonis		61.19	0.50	XYL 90	1	
Prevotella oralis		23.49	0.30	SOR 1	1	
Clostridium ramosum		5.14	0.19		92	
47731372	**Low discrimination**					**47731372**
					GRA	
Parabacteroides distasonis		82.89	0.45	XYL 90	1	
Bifidobacterium breve		11.52	0.29	SOR 1	99	
Clostridium clostridioforme		5.59	0.19		50	
47731401	**Good identification**					**47731401**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47731402	**Low discrimination**					**47731402**
Clostridium clostridioforme		41.71	0.39	XYL 91		
Bifidobacterium adolescentis		40.08	0.42	RHA 88		
Actinomyces israelii		10.49	0.38			
47731403	**Low discrimination**					**47731403**
Actinomyces naeslundii		74.56	0.36	ESL 12		
Clostridium clostridioforme		12.97	0.05	ARA 1		
Bifidobacterium adolescentis		12.47	0.07			
47731412	**Low discrimination**					**47731412**
Clostridium clostridioforme		78.84	0.50	XYL 91		
Bifidobacterium adolescentis		20.33	0.43	RHA 88		
Actinomyces israelii		0.65	0.24			
47731421	**Good identification**					**47731421**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47731422	**Low discrimination**					**47731422**
Clostridium clostridioforme		60.14	0.39	XYL 91		
Parabacteroides distasonis		24.39	0.38	RHA 88		
Prevotella oralis		6.83	0.16			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47731423	**Low discrimination**					**47731423**
Actinomyces naeslundii		85.18	0.36	ESL 12		
Clostridium clostridioforme		14.82	0.05	ARA 1		
47731432	**Acceptable identification**					**47731432**
					GRA	
Clostridium clostridioforme		94.67	0.50	XYL 91	50	
Parabacteroides distasonis		4.54	0.33	RHA 88	1	
Bifidobacterium longum subsp. longum		0.55	0.13		99	
47731500	**Low discrimination**					**47731500**
					SPO	
Clostridium butyricum		65.46	0.40	CEL 99	99	
Bifidobacterium adolescentis		11.15	0.42	XYL 97	1	
Actinomyces israelii		7.79	0.45		1	
47731501	**Low discrimination**					**47731501**
					SPO	
Actinomyces naeslundii		71.24	0.44	ESL 12	1	
Clostridium butyricum		24.57	0.06	ARA 1	99	
Bifidobacterium adolescentis		4.19	0.07		1	
47731502	**Good identification**					**47731502**
					SPO	
Bifidobacterium adolescentis		91.62	0.76	XYL 87	1	
Clostridium clostridioforme		3.22	0.48		66	
Clostridium butyricum		2.11	0.33		99	
47731503	**Low discrimination**					**47731503**
Bifidobacterium adolescentis		79.99	0.42	XYL 87		
Actinomyces naeslundii		13.75	0.44	URE 1		
Clostridium clostridioforme		2.82	0.14			
47731510	**Low discrimination**					**47731510**
					SPO	
Bifidobacterium adolescentis		49.24	0.43	XYL 87	1	
Bifidobacterium longum subsp. longum		27.21	0.37	BGL 99	1	
Clostridium butyricum		12.45	0.18		99	
47731512	**Low discrimination**					**47731512**
					GRA	
Bifidobacterium adolescentis		86.89	0.77	XYL 87	99	
Clostridium clostridioforme		11.40	0.59		50	
Leptotrichia buccalis		0.78	0.24		1	
47731513	**Low discrimination**					**47731513**
					GRA	
Bifidobacterium adolescentis		87.71	0.43	XYL 87	99	
Clostridium clostridioforme		11.50	0.25	URE 1	50	
Leptotrichia buccalis		0.79	-.11		1	
47731520	**Low discrimination**					**47731520**
					GRA	
Bifidobacterium longum subsp. longum		35.86	0.43	SAL 1	99	
Clostridium perfringens		17.53	0.33		99	
Bacteroides vulgatus		11.64	0.21		1	
47731521	**Acceptable identification**					**47731521**
Actinomyces naeslundii		91.47	0.44	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. longum		8.53	0.09	ARA 1		
47731522	**Low discrimination**					**47731522**
Parabacteroides distasonis		79.38	0.72	XYL 90		
Clostridium clostridioforme		6.62	0.48			
Prevotella oralis		3.52	0.36			
47731523	**Low discrimination**					**47731523**
					GRA	
Parabacteroides distasonis		61.11	0.38	XYL 90	1	
Actinomyces naeslundii		24.89	0.44	URE 1	99	
Clostridium clostridioforme		5.09	0.14		50	
47731530	**Low discrimination**					**47731530**
					GRA	
Bifidobacterium longum subsp. longum		85.94	0.47	SAL 1	99	
Parabacteroides distasonis		7.23	0.33		1	
Clostridium clostridioforme		5.10	0.25		50	
47731532	**Low discrimination**					**47731532**
					GRA	
Parabacteroides distasonis		52.71	0.67	XYL 90	1	
Clostridium clostridioforme		37.18	0.59		50	
Bifidobacterium longum subsp. longum		6.33	0.47		99	
47731542	**Good identification**					**47731542**
					SPO	
Bifidobacterium adolescentis		94.17	0.60	XYL 87	1	
Actinomyces israelii		3.14	0.41	SOR 11	1	
Clostridium butyricum		1.53	0.15		99	
47731552	**Very good identification**					**47731552**
Bifidobacterium adolescentis		97.84	0.62	XYL 87		
Clostridium clostridioforme		1.05	0.25	SOR 11		
Bifidobacterium breve		0.57	0.24			
47731562	**Low discrimination**					**47731562**
					GRA	
Parabacteroides distasonis		45.24	0.38	XYL 90	1	
Bifidobacterium breve		16.58	0.29	SOR 1	99	
Bifidobacterium adolescentis		13.25	0.26		99	
47731602	**Good identification**					**47731602**
					GRA	
Clostridium clostridioforme		99.45	0.54	XYL 91	50	
Actinomyces israelii		0.26	0.19		99	
Parabacteroides distasonis		0.17	0.12		1	
47731612	**Very good identification**					**47731612**
Clostridium clostridioforme		99.97	0.65	XYL 91		
Bifidobacterium adolescentis		0.04	0.09			
47731622	**Low discrimination**					**47731622**
Clostridium clostridioforme		85.17	0.54	XYL 91		
Parabacteroides distasonis		14.13	0.46			
Prevotella oralis		0.57	0.09			
47731632	**Very good identification**					**47731632**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		98.08	0.65	XYL 91		
Parabacteroides distasonis		1.92	0.41			
47731702	**Good identification**					**47731702**
					GRA	
Clostridium clostridioforme		91.05	0.63	XYL 91	50	
Parabacteroides distasonis		4.51	0.46		1	
Bifidobacterium adolescentis		3.56	0.42		99	
47731710	**Good identification**					**47731710**
					SPO	
Clostridium clostridioforme		98.17	0.40	XYL 91	66	
Bifidobacterium adolescentis		1.03	0.09	BGL 99	1	
Clostridium butyricum		0.80	-.08		99	
47731712	**Very good identification**					**47731712**
					GRA	
Clostridium clostridioforme		98.37	0.74	XYL 91	50	
Bifidobacterium adolescentis		1.03	0.43		99	
Parabacteroides distasonis		0.58	0.41		1	
47731713	**Good identification**					**47731713**
Clostridium clostridioforme		98.96	0.40	XYL 91		
Bifidobacterium adolescentis		1.04	0.09	URE 1		
47731720	**Low discrimination**					**47731720**
Bacteroides vulgatus		73.34	0.37	TRE 4		
Parabacteroides distasonis		20.03	0.46	XYL 98		
Clostridium clostridioforme		4.08	0.28			
47731722	**Low discrimination**					**47731722**
Parabacteroides distasonis		82.17	0.80	XYL 90		
Clostridium clostridioforme		16.74	0.63			
Prevotella oralis		0.52	0.30			
47731723	**Low discrimination**					**47731723**
Parabacteroides distasonis		83.07	0.46	XYL 90		
Clostridium clostridioforme		16.93	0.28	URE 1		
47731730	**Low discrimination**					**47731730**
Clostridium clostridioforme		60.86	0.40	XYL 91		
Parabacteroides distasonis		35.30	0.41	BGL 99		
Bacteroides vulgatus		2.42	0.03			
47731732	**Low discrimination**					**47731732**
Clostridium clostridioforme		63.19	0.74	XYL 91		
Parabacteroides distasonis		36.65	0.75			
Bacteroides thetaiotaomicron		0.12	0.17			
47731733	**Low discrimination**					**47731733**
Clostridium clostridioforme		63.29	0.40	XYL 91		
Parabacteroides distasonis		36.71	0.41	URE 1		
47731752	**Low discrimination**					**47731752**
Clostridium clostridioforme		88.61	0.40	XYL 91		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		11.39	0.28	SOR 1			
47731762	**Low discrimination**						**47731762**
Parabacteroides distasonis		83.07	0.46	XYL 90			
Clostridium clostridioforme		16.93	0.28	SOR 1			
47731772	**Low discrimination**						**47731772**
Clostridium clostridioforme		63.29	0.40	XYL 91			
Parabacteroides distasonis		36.71	0.41	SOR 1			
47732000	**Low discrimination**						**47732000**
					GRA	CAT	
Actinomyces naeslundii		55.67	0.49	ESL 12	99	25	
Fusobacterium mortiferum		28.08	0.32	GAL 96	1	5	
Propionibacterium granulosum		14.15	0.24		99	90	
47732001	**Good identification**						**47732001**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47732002	**Low discrimination**						**47732002**
					CAT		
Actinomyces naeslundii		85.01	0.49	ESL 12	25		
Propionibacterium granulosum		5.40	0.13	GAL 96	90		
Actinomyces israelii		4.83	0.18		1		
47732003	**Good identification**						**47732003**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47732020	**Acceptable identification**						**47732020**
Actinomyces naeslundii		94.03	0.49	ESL 12			
Propionibacterium granulosum		5.97	0.13	GAL 96			
47732021	**Good identification**						**47732021**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47732022	**Good identification**						**47732022**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47732023	**Good identification**						**47732023**
Actinomyces naeslundii		100.0	0.52	ESL 12			
47732040	**Good identification**						**47732040**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47732041	**Good identification**						**47732041**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47732042	**Good identification**						**47732042**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47732043	**Good identification**						**47732043**
Actinomyces naeslundii		100.0	0.39	ESL 12			
47732060	**Good identification**						**47732060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	ESL 12		
47732061	**Good identification**					**47732061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47732062	**Good identification**					**47732062**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47732063	**Good identification**					**47732063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
47732100	**Low discrimination**					**47732100**
					GRA	
Actinomyces naeslundii		68.18	0.57	ESL 12	99	
Fusobacterium mortiferum		28.19	0.38	GAL 96	1	
Actinomyces israelii		3.63	0.25		99	
47732101	**Very good identification**					**47732101**
Actinomyces naeslundii		100.0	0.60	ESL 12		
47732102	**Low discrimination**					**47732102**
Actinomyces naeslundii		73.31	0.57	ESL 12		
Bifidobacterium adolescentis		9.77	0.26	GAL 96		
Bifidobacterium breve		6.95	0.24			
47732103	**Very good identification**					**47732103**
Actinomyces naeslundii		100.0	0.60	ESL 12		
47732111	**Good identification**					**47732111**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47732113	**Good identification**					**47732113**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47732120	**Low discrimination**					**47732120**
					SPO	
Actinomyces naeslundii		82.06	0.57	ESL 12	1	
Clostridium ramosum		16.58	0.19	GAL 96	92	
Bifidobacterium breve		1.37	0.12		1	
47732121	**Very good identification**					**47732121**
Actinomyces naeslundii		100.0	0.60	ESL 12		
47732122	**Low discrimination**					**47732122**
					SPO	
Actinomyces naeslundii		49.63	0.57	ESL 12	1	
Clostridium ramosum		40.10	0.30	GAL 96	92	
Bifidobacterium breve		8.36	0.29		1	
47732123	**Very good identification**					**47732123**
Actinomyces naeslundii		100.0	0.60	ESL 12		
47732131	**Good identification**					**47732131**
Actinomyces naeslundii		100.0	0.38	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47732133	**Good identification**					**47732133**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47732140	**Low discrimination**					**47732140**
Actinomyces naeslundii		89.50	0.44	ESL 12		
Actinomyces israelii		10.50	0.18	GAL 96		
47732141	**Good identification**					**47732141**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47732142	**Low discrimination**					**47732142**
Actinomyces naeslundii		55.36	0.44	ESL 12		
Bifidobacterium breve		29.74	0.24	GAL 96		
Actinomyces israelii		9.74	0.21			
47732143	**Good identification**					**47732143**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47732160	**Acceptable identification**					**47732160**
Actinomyces naeslundii		91.37	0.44	ESL 12		
Bifidobacterium breve		8.63	0.12	GAL 96		
47732161	**Good identification**					**47732161**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47732162	**Low discrimination**					**47732162**
Actinomyces naeslundii		51.15	0.44	ESL 12		
Bifidobacterium breve		48.85	0.29	GAL 96		
47732163	**Good identification**					**47732163**
Actinomyces naeslundii		100.0	0.47	ESL 12		
47732502	**Acceptable identification**					**47732502**
Bifidobacterium adolescentis		91.00	0.40	GAL 99		
Actinomyces israelii		6.67	0.27	XYL 87		
Clostridium clostridioforme		1.31	0.05			
47732512	**Acceptable identification**					**47732512**
Bifidobacterium adolescentis		93.43	0.42	GAL 99		
Clostridium clostridioforme		4.99	0.17	XYL 87		
Bifidobacterium dentium		1.58	-.07			
47733000	**Low discrimination**					**47733000**
					GRA	
Actinomyces naeslundii		83.17	0.72	ESL 12	99	
Actinomyces israelii		12.99	0.49		99	
Fusobacterium mortiferum		2.62	0.35		1	
47733001	**Very good identification**					**47733001**
Actinomyces naeslundii		99.90	0.75	ESL 12		
Actinomyces israelii		0.11	0.15			
47733002	**Low discrimination**					**47733002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		78.39	0.72	ESL 12		
Actinomyces israelii		18.37	0.52			
Bifidobacterium adolescentis		1.24	0.26			
47733003	**Very good identification**					**47733003**
Actinomyces naeslundii		99.84	0.75	ESL 12		
Actinomyces israelii		0.16	0.18			
47733010	**Low discrimination**					**47733010**
					CAT	
Actinomyces naeslundii		66.82	0.50	ESL 12	25	
Actinomyces israelii		29.64	0.35	MLZ 5	1	
Propionibacterium granulosum		1.83	0.09		90	
47733011	**Good identification**					**47733011**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47733012	**Low discrimination**					**47733012**
Actinomyces naeslundii		44.00	0.50	ESL 12		
Actinomyces israelii		29.27	0.38	MLZ 5		
Bifidobacterium adolescentis		16.14	0.28			
47733013	**Good identification**					**47733013**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47733020	**Very good identification**					**47733020**
					SPO	
Actinomyces naeslundii		95.04	0.72	ESL 12	1	
Clostridium ramosum		2.28	0.19		92	
Clostridium perfringens		1.30	0.18		84	
47733021	**Very good identification**					**47733021**
Actinomyces naeslundii		100.0	0.75	ESL 12		
47733022	**Low discrimination**					**47733022**
					SPO	
Actinomyces naeslundii		86.30	0.72	ESL 12	1	
Clostridium ramosum		8.27	0.30		92	
Bifidobacterium breve		1.73	0.29		1	
47733023	**Very good identification**					**47733023**
Actinomyces naeslundii		100.0	0.75	ESL 12		
47733030	**Acceptable identification**					**47733030**
Actinomyces naeslundii		93.82	0.50	ESL 12		
Bifidobacterium breve		3.52	0.12	MLZ 5		
Actinomyces israelii		2.66	0.15			
47733031	**Good identification**					**47733031**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47733032	**Low discrimination**					**47733032**
Actinomyces naeslundii		70.32	0.50	ESL 12		
Bifidobacterium breve		26.70	0.29	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		2.99	0.18			
47733033	**Good identification**					**47733033**
Actinomyces naeslundii		100.0	0.53	ESL 12		
47733040	**Low discrimination**					**47733040**
Actinomyces naeslundii		74.39	0.59	ESL 12		
Actinomyces israelii		25.61	0.42			
47733041	**Very good identification**					**47733041**
Actinomyces naeslundii		100.0	0.62	ESL 12		
47733042	**Low discrimination**					**47733042**
Actinomyces naeslundii		62.85	0.59	ESL 12		
Actinomyces israelii		32.45	0.45			
Bifidobacterium breve		4.00	0.24			
47733043	**Very good identification**					**47733043**
Actinomyces naeslundii		100.0	0.62	ESL 12		
47733050	**Low discrimination**					**47733050**
Actinomyces naeslundii		50.57	0.38	ESL 12		
Actinomyces israelii		49.43	0.28	MLZ 5		
47733051	**Good identification**					**47733051**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733053	**Good identification**					**47733053**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733060	**Very good identification**					**47733060**
Actinomyces naeslundii		95.43	0.59	ESL 12		
Actinomyces israelii		2.10	0.22			
Clostridium perfringens		1.41	0.06			
47733061	**Very good identification**					**47733061**
Actinomyces naeslundii		100.0	0.62	ESL 12		
47733062	**Low discrimination**					**47733062**
Actinomyces naeslundii		86.72	0.59	ESL 12		
Bifidobacterium breve		9.82	0.29			
Actinomyces israelii		2.86	0.25			
47733063	**Very good identification**					**47733063**
Actinomyces naeslundii		100.0	0.62	ESL 12		
47733070	**Low discrimination**					**47733070**
Actinomyces naeslundii		82.46	0.38	ESL 12		
Bifidobacterium breve		17.54	0.12	MLZ 5		
47733071	**Good identification**					**47733071**
Actinomyces naeslundii		100.0	0.41	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47733073	**Good identification**					**47733073**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733100	**Low discrimination**					**47733100**
Actinomyces naeslundii		76.00	0.80	ESL 12		
Actinomyces israelii		16.69	0.60			
Bifidobacterium breve		2.94	0.41			
47733101	**Very good identification**					**47733101**
Actinomyces naeslundii		99.85	0.83	ESL 12		
Actinomyces israelii		0.15	0.25			
47733102	**Low discrimination**					**47733102**
Actinomyces naeslundii		42.05	0.80	ESL 12		
Bifidobacterium adolescentis		23.10	0.60			
Bifidobacterium breve		16.44	0.59			
47733103	**Very good identification**					**47733103**
Actinomyces naeslundii		99.12	0.83	ESL 12		
Bifidobacterium adolescentis		0.37	0.26			
Bifidobacterium breve		0.26	0.24			
47733110	**Low discrimination**					**47733110**
Actinomyces naeslundii		39.21	0.58	ESL 12		
Bifidobacterium breve		28.81	0.41	MLZ 5		
Actinomyces israelii		24.44	0.46			
47733111	**Very good identification**					**47733111**
Actinomyces naeslundii		100.0	0.61	ESL 12		
47733112	**Low discrimination**					**47733112**
Bifidobacterium adolescentis		54.87	0.62	XYL 87		
Bifidobacterium breve		31.94	0.59	ARA 87		
Actinomyces naeslundii		4.30	0.58			
47733113	**Low discrimination**					**47733113**
Actinomyces naeslundii		87.07	0.61	ESL 12		
Bifidobacterium adolescentis		7.48	0.28	MLZ 5		
Bifidobacterium breve		4.36	0.24			
47733120	**Low discrimination**					**47733120**
					SPO	
Actinomyces naeslundii		50.71	0.80	ESL 12	1	
Clostridium ramosum		42.26	0.53		92	
Bifidobacterium breve		3.49	0.46		1	
47733121	**Very good identification**					**47733121**
					SPO	
Actinomyces naeslundii		99.40	0.83	ESL 12	1	
Clostridium ramosum		0.56	0.19		92	
Bifidobacterium breve		0.05	0.12		1	
47733122	**Low discrimination**					**47733122**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		62.96	0.64	CEL 99	92	
Actinomyces naeslundii		18.89	0.80		1	
Bifidobacterium breve		13.13	0.63		1	
47733123	**Very good identification**					**47733123**
					SPO	
Actinomyces naeslundii		97.26	0.83	ESL 12	1	
Clostridium ramosum		2.18	0.30		92	
Bifidobacterium breve		0.46	0.29		1	
47733130	**Low discrimination**					**47733130**
					SPO	
Bifidobacterium breve		48.95	0.46	CEL 99	1	
Actinomyces naeslundii		37.48	0.58	BGL 91	1	
Clostridium tertium		6.09	0.21		99	
47733131	**Very good identification**					**47733131**
Actinomyces naeslundii		99.13	0.61	ESL 12		
Bifidobacterium breve		0.87	0.12	MLZ 5		
47733132	**Low discrimination**					**47733132**
					SPO	
Bifidobacterium breve		82.48	0.63	CEL 99	1	
Actinomyces naeslundii		6.25	0.58		1	
Clostridium ramosum		4.00	0.30		92	
47733133	**Good identification**					**47733133**
Actinomyces naeslundii		91.83	0.61	ESL 12		
Bifidobacterium breve		8.17	0.29	MLZ 5		
47733140	**Low discrimination**					**47733140**
Actinomyces naeslundii		58.59	0.67	ESL 12		
Actinomyces israelii		28.35	0.53			
Bifidobacterium breve		12.84	0.41			
47733141	**Very good identification**					**47733141**
Actinomyces naeslundii		99.68	0.70	ESL 12		
Actinomyces israelii		0.33	0.18			
47733142	**Low discrimination**					**47733142**
Bifidobacterium breve		51.05	0.59	CEL 99		
Actinomyces naeslundii		23.04	0.67			
Actinomyces israelii		16.72	0.56			
47733143	**Very good identification**					**47733143**
Actinomyces naeslundii		97.81	0.70	ESL 12		
Bifidobacterium breve		1.46	0.24			
Actinomyces israelii		0.48	0.21			
47733150	**Low discrimination**					**47733150**
Bifidobacterium breve		62.06	0.41	CEL 99		
Actinomyces israelii		20.47	0.38	BGL 91		
Actinomyces naeslundii		14.91	0.45			
47733151	**Good identification**					**47733151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.48	ESL 12		
47733152	**Low discrimination**					**47733152**
Bifidobacterium breve		77.74	0.59	CEL 99		
Bifidobacterium adolescentis		16.51	0.46			
Actinomyces israelii		3.81	0.42			
47733153	**Low discrimination**					**47733153**
Actinomyces naeslundii		74.42	0.48	ESL 12		
Bifidobacterium breve		21.10	0.24	MLZ 5		
Bifidobacterium adolescentis		4.48	0.12			
47733160	**Low discrimination**					**47733160**
					SPO	
Actinomyces naeslundii		66.39	0.67	ESL 12	1	
Bifidobacterium breve		25.87	0.46		1	
Clostridium ramosum		3.17	0.19		92	
47733161	**Very good identification**					**47733161**
Actinomyces naeslundii		99.74	0.70	ESL 12		
Bifidobacterium breve		0.26	0.12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47733162	**Low discrimination**					**47733162**
					SPO	
Bifidobacterium breve		75.48	0.63	CEL 99	1	
Actinomyces naeslundii		19.16	0.67		1	
Clostridium ramosum		3.66	0.30		92	
47733163	**Very good identification**					**47733163**
Actinomyces naeslundii		97.42	0.70	ESL 12		
Bifidobacterium breve		2.58	0.29			
47733170	**Low discrimination**					**47733170**
Bifidobacterium breve		87.19	0.46	CEL 99		
Actinomyces naeslundii		11.78	0.45	BGL 91		
Actinomyces israelii		1.03	0.18			
47733171	**Good identification**					**47733171**
Actinomyces naeslundii		95.25	0.48	ESL 12		
Bifidobacterium breve		4.75	0.12	MLZ 5		
47733172	**Very good identification**					**47733172**
Bifidobacterium breve		98.39	0.63	CEL 99		
Actinomyces naeslundii		1.32	0.45			
Actinomyces israelii		0.17	0.21			
47733173	**Low discrimination**					**47733173**
Actinomyces naeslundii		66.49	0.48	ESL 12		
Bifidobacterium breve		33.51	0.29	MLZ 5		
47733201	**Good identification**					**47733201**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733203	**Good identification**					**47733203**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733221	**Good identification**					**47733221**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733223	**Acceptable identification**					**47733223**
					SPO	
Actinomyces naeslundii		92.19	0.41	ESL 12	1	
Clostridium ramosum		7.81	-.03	RHA 1	92	
47733300	**Low discrimination**					**47733300**
					SPO	
Actinomyces israelii		43.74	0.40	CEL 93	1	
Clostridium ramosum		29.53	0.21	XYL 92	92	
Actinomyces naeslundii		26.73	0.46	RHA 7	1	
47733301	**Good identification**					**47733301**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47733303	**Good identification**					**47733303**
					SPO	
Actinomyces naeslundii		97.11	0.49	ESL 12	1	
Clostridium ramosum		2.89	-.03	RHA 1	92	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47733320	**Good identification**						**47733320**
					SPO	GRA	
Clostridium ramosum		98.69	0.56	CEL 99	92	50	
Actinomyces naeslundii		0.90	0.46		1	99	
Prevotella oralis		0.17	0.08		1	1	
47733321	**Low discrimination**						**47733321**
					SPO		
Actinomyces naeslundii		57.59	0.49	ESL 12	1		
Clostridium ramosum		42.41	0.21	RHA 1	92		
47733322	**Very good identification**						**47733322**
					SPO		
Clostridium ramosum		96.69	0.66	CEL 99	92		
Parabacteroides distasonis		2.05	0.50		1		
Prevotella oralis		0.79	0.30		1		
47733332	**Low discrimination**						**47733332**
					SPO	GRA	
Parabacteroides distasonis		47.61	0.45	MAN 1	1	1	
Clostridium ramosum		42.07	0.32	XYL 90	92	50	
Bifidobacterium breve		6.62	0.29		1	99	
47733341	**Good identification**						**47733341**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47733342	**Low discrimination**						**47733342**
Actinomyces israelii		58.39	0.36	CEL 93			
Bifidobacterium breve		23.92	0.24	XYL 92			
Actinomyces naeslundii		10.80	0.33	RHA 7			
47733343	**Good identification**						**47733343**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47733361	**Good identification**						**47733361**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47733363	**Low discrimination**						**47733363**
					SPO		
Actinomyces naeslundii		85.57	0.36	ESL 12	1		
Clostridium ramosum		14.43	-.03	RHA 1	92		
47733400	**Acceptable identification**						**47733400**
					SPO		
Actinomyces israelii		90.37	0.48	CEL 93	1		
Actinomyces naeslundii		7.14	0.38	XYL 92	1		
Clostridium butyricum		2.48	0.00		99		
47733401	**Acceptable identification**						**47733401**
Actinomyces naeslundii		92.15	0.41	ESL 12			
Actinomyces israelii		7.85	0.14	ARA 1			
47733402	**Low discrimination**						**47733402**
Actinomyces israelii		61.79	0.51	CEL 93			
Bifidobacterium adolescentis		34.08	0.40	XYL 92			
Actinomyces naeslundii		3.26	0.38				
47733403	**Low discrimination**						**47733403**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		88.67	0.41	ESL 12		
Actinomyces israelii		11.33	0.17	ARA 1		
47733412	**Low discrimination**					**47733412**
Bifidobacterium adolescentis		77.73	0.42	XYL 87		
Actinomyces israelii		17.23	0.37	LAC 99		
Clostridium clostridioforme		3.72	0.16			
47733420	**Low discrimination**					**47733420**
Actinomyces naeslundii		55.32	0.38	ESL 12		
Actinomyces israelii		44.68	0.27	ARA 1		
47733421	**Good identification**					**47733421**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733423	**Good identification**					**47733423**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47733440	**Good identification**					**47733440**
Actinomyces israelii		96.54	0.41	CEL 93		
Actinomyces naeslundii		3.46	0.25	XYL 92		
47733442	**Low discrimination**					**47733442**
Actinomyces israelii		83.39	0.44	CEL 93		
Bifidobacterium adolescentis		14.62	0.25	XYL 92		
Actinomyces naeslundii		1.99	0.25			
47733500	**Low discrimination**					**47733500**
					SPO	
Actinomyces israelii		70.03	0.58	CEL 93	1	
Bifidobacterium adolescentis		14.48	0.40	XYL 92	1	
Clostridium butyricum		11.55	0.24		99	
47733501	**Low discrimination**					**47733501**
Actinomyces naeslundii		89.31	0.49	ESL 12		
Actinomyces israelii		10.69	0.24	ARA 1		
47733502	**Good identification**					**47733502**
					SPO	
Bifidobacterium adolescentis		92.51	0.74	XYL 87	1	
Actinomyces israelii		6.78	0.61		1	
Clostridium butyricum		0.29	0.17		99	
47733503	**Low discrimination**					**47733503**
Bifidobacterium adolescentis		67.51	0.40	XYL 87		
Actinomyces naeslundii		27.54	0.49	URE 1		
Actinomyces israelii		4.95	0.27			
47733510	**Low discrimination**					**47733510**
Bifidobacterium adolescentis		54.97	0.42	XYL 87		
Actinomyces israelii		32.50	0.44	BGL 99		
Bifidobacterium dentium		9.09	0.10			
47733512	**Very good identification**					**47733512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		97.21	0.76	XYL 87			
Bifidobacterium dentium		1.64	0.27				
Actinomyces israelii		0.87	0.47				
47733513	**Good identification**						**47733513**
Bifidobacterium adolescentis		96.67	0.42	XYL 87			
Actinomyces naeslundii		1.70	0.27	URE 1			
Bifidobacterium dentium		1.63	-.07				
47733520	**Low discrimination**						**47733520**
					SPO		
Actinomyces israelii		33.74	0.38	CEL 93	1		
Actinomyces naeslundii		29.73	0.46	XYL 92	1		
Clostridium ramosum		24.77	0.19	NAG 6	92		
47733521	**Good identification**						**47733521**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47733522	**Low discrimination**						**47733522**
					SPO	GRA	
Bifidobacterium adolescentis		28.44	0.40	XYL 87	1	99	
Clostridium ramosum		25.78	0.30	NAG 1	92	50	
Parabacteroides distasonis		14.67	0.38		1	1	
47733523	**Good identification**						**47733523**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47733540	**Low discrimination**						**47733540**
					COC		
Actinomyces israelii		67.34	0.51	CEL 93	1		
Blautia producta		24.04	0.24	XYL 92	99		
Bifidobacterium adolescentis		4.43	0.25		1		
47733541	**Low discrimination**						**47733541**
Actinomyces naeslundii		79.12	0.36	ESL 12			
Actinomyces israelii		20.88	0.17	ARA 1			
47733542	**Low discrimination**						**47733542**
Bifidobacterium adolescentis		79.91	0.59	XYL 87			
Actinomyces israelii		18.42	0.54	SOR 11			
Bifidobacterium breve		0.69	0.24				
47733552	**Very good identification**						**47733552**
					COC		
Bifidobacterium adolescentis		95.09	0.60	XYL 87	1		
Actinomyces israelii		2.68	0.40	SOR 11	1		
Blautia producta		1.42	0.15		99		
47733561	**Good identification**						**47733561**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47733563	**Good identification**						**47733563**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47733700	**Acceptable identification**						**47733700**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		93.66	0.39	CEL 93	1			
Clostridium butyricum		6.34	-.02	XYL 92	99			
47733702	**Low discrimination**							**47733702**
Bifidobacterium adolescentis		52.93	0.40	XYL 87				
Actinomyces israelii		28.90	0.42	RHA 1				
Clostridium clostridioforme		16.70	0.28					
47733712	**Low discrimination**							**47733712**
Clostridium clostridioforme		52.04	0.40	MAN 1				
Bifidobacterium adolescentis		44.26	0.42	XYL 91				
Actinomyces israelii		2.96	0.28					
47733722	**Low discrimination**							**47733722**
					SPO			
Parabacteroides distasonis		49.61	0.46	MAN 1	1			
Clostridium ramosum		38.51	0.32	XYL 90	92			
Clostridium clostridioforme		10.11	0.28		66			
47733732	**Low discrimination**							**47733732**
					GRA			
Clostridium clostridioforme		61.28	0.40	MAN 1	50			
Parabacteroides distasonis		35.54	0.41	XYL 91	1			
Bifidobacterium dentium		1.56	-.03		99			
47734000	**Good identification**							**47734000**
					GRA			
Fusobacterium mortiferum		98.00	0.47	MLT 11	1			
Actinomyces israelii		2.00	0.20	XYL 7	99			
47734012	**Good identification**							**47734012**
Clostridium clostridioforme		98.80	0.39	RHA 88				
Bifidobacterium adolescentis		1.20	0.09	ARA 94				
47734032	**Good identification**							**47734032**
Clostridium clostridioforme		96.80	0.39	RHA 88				
Parabacteroides distasonis		3.20	0.19	ARA 94				
47734100	**Low discrimination**							**47734100**
					COC	GRA	SPO	
Fusobacterium mortiferum		79.98	0.53	MLT 11	1	1	1	
Atopobium parvulum		8.78	0.25	XYL 7	99	99	1	
Clostridium butyricum		6.03	0.15		1	99	99	
47734102	**Low discrimination**							**47734102**
Bifidobacterium adolescentis		31.40	0.42	GAL 99				
Bifidobacterium longum subsp. infantis		23.86	0.26	ARA 87				
Clostridium clostridioforme		23.54	0.36					
47734112	**Low discrimination**							**47734112**
					GRA			
Clostridium clostridioforme		72.59	0.48	RHA 88	50			
Bifidobacterium adolescentis		25.99	0.43	ARA 94	99			
Parabacteroides distasonis		0.72	0.19		1			
47734122	**Low discrimination**							**47734122**
					GRA			
Parabacteroides distasonis		72.15	0.58	GAL 99	1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. infantis		15.72	0.30		99	
Clostridium clostridioforme		8.72	0.36		50	
47734132	**Low discrimination**					**47734132**
					GRA	
Clostridium clostridioforme		50.37	0.48	RHA 88	50	
Parabacteroides distasonis		49.24	0.53	ARA 94	1	
Bifidobacterium longum subsp. infantis		0.20	-.04		99	
47734202	**Good identification**					**47734202**
Clostridium clostridioforme		100.0	0.42	ARA 94		
47734212	**Good identification**					**47734212**
Clostridium clostridioforme		100.0	0.53	ARA 94		
47734222	**Low discrimination**					**47734222**
Clostridium clostridioforme		89.73	0.42	ARA 94		
Parabacteroides distasonis		10.27	0.32			
47734232	**Good identification**					**47734232**
Clostridium clostridioforme		98.67	0.53	ARA 94		
Parabacteroides distasonis		1.33	0.27			
47734302	**Good identification**					**47734302**
					GRA	
Clostridium clostridioforme		95.44	0.51	ARA 94	50	
Parabacteroides distasonis		3.26	0.32		1	
Eubacterium contortum		0.93	0.11		99	
47734312	**Very good identification**					**47734312**
					GRA	
Clostridium clostridioforme		99.55	0.62	ARA 94	50	
Parabacteroides distasonis		0.40	0.27		1	
Bifidobacterium adolescentis		0.05	0.09		99	
47734322	**Low discrimination**					**47734322**
Parabacteroides distasonis		76.80	0.66	GAL 99		
Clostridium clostridioforme		22.70	0.51			
Prevotella oralis		0.41	0.14			
47734332	**Low discrimination**					**47734332**
Clostridium clostridioforme		71.43	0.62	ARA 94		
Parabacteroides distasonis		28.57	0.61			
47734402	**Good identification**					**47734402**
					SPO	
Clostridium clostridioforme		96.45	0.48	RHA 88	66	
Bifidobacterium adolescentis		1.86	0.22		1	
Clostridium butyricum		0.85	0.01		99	
47734412	**Very good identification**					**47734412**
Clostridium clostridioforme		99.49	0.59	RHA 88		
Bifidobacterium adolescentis		0.51	0.23			
47734422	**Good identification**					**47734422**
Clostridium clostridioforme		98.92	0.48	RHA 88		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		1.08	0.20			
47734432	**Very good identification**					**47734432**
Clostridium clostridioforme		99.87	0.59	RHA 88		
Parabacteroides distasonis		0.13	0.15			
47734502	**Low discrimination**					**47734502**
					SPO	
Clostridium clostridioforme		60.68	0.57	RHA 88	66	
Bifidobacterium adolescentis		34.58	0.56		1	
Clostridium butyricum		3.85	0.25		99	
47734512	**Low discrimination**					**47734512**
					SPO	
Clostridium clostridioforme		86.57	0.68	RHA 88	66	
Bifidobacterium adolescentis		13.24	0.57		1	
Clostridium butyricum		0.06	0.03		99	
47734522	**Low discrimination**					**47734522**
					GRA	
Clostridium clostridioforme		74.57	0.57	RHA 88	50	
Parabacteroides distasonis		24.13	0.54		1	
Bifidobacterium adolescentis		0.43	0.22		99	
47734532	**Very good identification**					**47734532**
					GRA	
Clostridium clostridioforme		95.98	0.68	RHA 88	50	
Parabacteroides distasonis		3.67	0.49		1	
Bifidobacterium adolescentis		0.15	0.23		99	
47734542	**Low discrimination**					**47734542**
					SPO	
Bifidobacterium adolescentis		78.19	0.40	GAL 99	1	
Clostridium clostridioforme		11.21	0.23	SOR 11	66	
Clostridium butyricum		6.13	0.07		99	
47734552	**Low discrimination**					**47734552**
Bifidobacterium adolescentis		65.17	0.42	GAL 99		
Clostridium clostridioforme		34.83	0.34	SOR 11		
47734602	**Very good identification**					**47734602**
Clostridium clostridioforme		100.0	0.63			
47734610	**Good identification**					**47734610**
Clostridium clostridioforme		100.0	0.40	BGL 99		
47734612	**Very good identification**					**47734612**
Clostridium clostridioforme		100.0	0.74			
47734613	**Good identification**					**47734613**
Clostridium clostridioforme		100.0	0.40	URE 1		
47734622	**Very good identification**					**47734622**
Clostridium clostridioforme		99.55	0.63			
Parabacteroides distasonis		0.45	0.28			
47734630	**Good identification**					**47734630**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		100.0	0.40	BGL 99		
47734632	**Very good identification**					**47734632**
Clostridium clostridioforme		99.95	0.74			
Parabacteroides distasonis		0.05	0.23			
47734633	**Good identification**					**47734633**
Clostridium clostridioforme		100.0	0.40	URE 1		
47734652	**Good identification**					**47734652**
Clostridium clostridioforme		100.0	0.40	SOR 1		
47734672	**Good identification**					**47734672**
Clostridium clostridioforme		100.0	0.40	SOR 1		
47734700	**Low discrimination**					**47734700**
					SPO	
Clostridium clostridioforme		72.38	0.37	BGL 99	66	
Eubacterium contortum		20.76	0.23		1	
Clostridium butyricum		4.93	0.06		99	
47734702	**Very good identification**					**47734702**
					GRA	
Clostridium clostridioforme		99.71	0.72		50	
Parabacteroides distasonis		0.13	0.28		1	
Bifidobacterium adolescentis		0.08	0.22		99	
47734703	**Good identification**					**47734703**
Clostridium clostridioforme		100.0	0.37	URE 1		
47734710	**Good identification**					**47734710**
Clostridium clostridioforme		100.0	0.49	BGL 99		
47734712	**Very good identification**					**47734712**
					GRA	
Clostridium clostridioforme		99.96	0.83		50	
Bifidobacterium adolescentis		0.02	0.23		99	
Parabacteroides distasonis		0.02	0.23		1	
47734713	**Good identification**					**47734713**
Clostridium clostridioforme		100.0	0.49	URE 1		
47734722	**Low discrimination**					**47734722**
Clostridium clostridioforme		88.12	0.72			
Parabacteroides distasonis		11.66	0.62			
Bacteroides thetaiotaomicron		0.14	0.13			
47734723	**Low discrimination**					**47734723**
Clostridium clostridioforme		88.31	0.37	URE 1		
Parabacteroides distasonis		11.69	0.28			
47734730	**Good identification**					**47734730**
Clostridium clostridioforme		97.90	0.49	BGL 99		
Parabacteroides distasonis		1.53	0.23			
Bacteroides vulgatus		0.57	-.02			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47734732	**Very good identification**						**47734732**
Clostridium clostridioforme		98.44	0.83				
Parabacteroides distasonis		1.54	0.58				
Bacteroides thetaiotaomicron		0.01	0.06				
47734733	**Good identification**						**47734733**
Clostridium clostridioforme		98.46	0.49	URE 1			
Parabacteroides distasonis		1.54	0.23				
47734742	**Good identification**						**47734742**
Clostridium clostridioforme		100.0	0.37	SOR 1			
47734752	**Good identification**						**47734752**
Clostridium clostridioforme		99.74	0.49	SOR 1			
Bifidobacterium adolescentis		0.26	0.08				
47734762	**Low discrimination**						**47734762**
Clostridium clostridioforme		88.31	0.37	SOR 1			
Parabacteroides distasonis		11.69	0.28				
47734772	**Good identification**						**47734772**
Clostridium clostridioforme		98.46	0.49	SOR 1			
Parabacteroides distasonis		1.54	0.23				
47735000	**Low discrimination**						**47735000**
					GRA	SPO	
Actinomyces israelii		43.68	0.55	CEL 93	99	1	
Fusobacterium mortiferum		32.36	0.50		1	1	
Clostridium butyricum		15.64	0.26		99	99	
47735001	**Low discrimination**						**47735001**
					GRA		
Actinomyces naeslundii		58.34	0.36	ESL 12	99		
Actinomyces israelii		23.93	0.20	XYL 1	99		
Fusobacterium mortiferum		17.73	0.16		1		
47735002	**Low discrimination**						**47735002**
Actinomyces israelii		53.26	0.58	CEL 93			
Bifidobacterium adolescentis		14.47	0.42				
Bifidobacterium longum subsp. infantis		11.00	0.26				
47735003	**Low discrimination**						**47735003**
Actinomyces naeslundii		56.10	0.36	ESL 12			
Actinomyces israelii		34.52	0.23	XYL 1			
Bifidobacterium adolescentis		9.38	0.07				
47735010	**Low discrimination**						**47735010**
					SPO		
Actinomyces israelii		75.61	0.40	CEL 93	1		
Clostridium butyricum		9.54	0.04	MLZ 13	99		
Clostridium clostridioforme		6.37	0.13		66		
47735012	**Low discrimination**						**47735012**
Clostridium clostridioforme		62.51	0.47	RHA 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		24.97	0.43	ARA 94			
Actinomyces israelii		11.24	0.43				
47735021	**Good identification**						**47735021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47735022	**Low discrimination**						**47735022**
					GRA		
Parabacteroides distasonis		61.52	0.58	LAC 99	1		
Prevotella oralis		14.35	0.34		1		
Bifidobacterium longum subsp. infantis		13.41	0.30		99		
47735032	**Low discrimination**						**47735032**
					GRA		
Parabacteroides distasonis		51.56	0.53	LAC 99	1		
Clostridium clostridioforme		47.27	0.47		50		
Actinomyces israelii		0.54	0.23		99		
47735040	**Acceptable identification**						**47735040**
					GRA	SPO	
Actinomyces israelii		90.97	0.48	CEL 93	99	1	
Clostridium butyricum		7.29	0.08		99	99	
Fusobacterium mortiferum		1.75	0.16		1	1	
47735042	**Acceptable identification**						**47735042**
					SPO		
Actinomyces israelii		90.33	0.51	CEL 93	1		
Bifidobacterium adolescentis		7.80	0.26		1		
Clostridium butyricum		1.88	0.01		99		
47735052	**Low discrimination**						**47735052**
Actinomyces israelii		54.03	0.36	CEL 93			
Bifidobacterium adolescentis		38.17	0.28	MLZ 13			
Clostridium clostridioforme		7.81	0.13				
47735100	**Low discrimination**						**47735100**
					COC	SPO	
Atopobium parvulum		41.00	0.60	RAF 1	99	1	
Clostridium butyricum		28.14	0.50		1	99	
Actinomyces israelii		13.10	0.65		1	1	
47735102	**Low discrimination**						**47735102**
Bifidobacterium adolescentis		46.46	0.76	ARA 87			
Bifidobacterium longum subsp. infantis		35.30	0.60				
Actinomyces israelii		6.91	0.68				
47735103	**Low discrimination**						**47735103**
Bifidobacterium adolescentis		43.19	0.42	ARA 87			
Bifidobacterium longum subsp. infantis		32.81	0.26	URE 1			
Actinomyces naeslundii		7.43	0.44				
47735110	**Low discrimination**						**47735110**
					SPO		
Actinomyces israelii		33.36	0.51	CEL 93	1		
Bifidobacterium adolescentis		27.81	0.43	MLZ 13	1		
Clostridium butyricum		25.25	0.28		99		
47735112	**Low discrimination**						**47735112**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		86.89	0.77	ARA 87	99		
Clostridium clostridioforme		7.36	0.56		50		
Parabacteroides distasonis		2.40	0.53		1		
47735113	**Low discrimination**						**47735113**
					GRA		
Bifidobacterium adolescentis		87.77	0.43	ARA 87	99		
Clostridium clostridioforme		7.43	0.22	URE 1	50		
Parabacteroides distasonis		2.42	0.19		1		
47735120	**Low discrimination**						**47735120**
					COC	GRA	
Bifidobacterium longum subsp. infantis		41.62	0.47	TRE 1	1	99	
Parabacteroides distasonis		19.51	0.58	BGL 91	1	1	
Atopobium parvulum		17.06	0.47		99	99	
47735121	**Low discrimination**						**47735121**
					GRA		
Actinomyces naeslundii		36.38	0.44	ESL 12	99		
Bifidobacterium longum subsp. infantis		28.26	0.13	XYL 1	99		
Parabacteroides distasonis		13.25	0.24		1		
47735122	**Low discrimination**						**47735122**
					GRA		
Parabacteroides distasonis		78.63	0.92		1		
Bifidobacterium longum subsp. infantis		17.13	0.64		99		
Prevotella oralis		2.90	0.55		1		
47735123	**Low discrimination**						**47735123**
					GRA		
Parabacteroides distasonis		77.13	0.58	URE 1	1		
Bifidobacterium longum subsp. infantis		16.81	0.30		99		
Prevotella oralis		2.85	0.20		1		
47735130	**Low discrimination**						**47735130**
					GRA		
Parabacteroides distasonis		84.28	0.53	BGL 99	1		
Bifidobacterium longum subsp. infantis		3.37	0.13		99		
Clostridium clostridioforme		2.61	0.22		50		
47735132	**Very good identification**						**47735132**
					GRA		
Parabacteroides distasonis		96.03	0.87		1		
Clostridium clostridioforme		2.98	0.56		50		
Bifidobacterium longum subsp. infantis		0.39	0.30		99		
47735133	**Good identification**						**47735133**
					GRA		
Parabacteroides distasonis		96.27	0.53	URE 1	1		
Clostridium clostridioforme		2.98	0.22		50		
Bifidobacterium longum subsp. infantis		0.39	-.04		99		
47735140	**Low discrimination**						**47735140**
					COC	SPO	
Actinomyces israelii		61.33	0.58	CEL 93	1	1	
Clostridium butyricum		29.47	0.31		1	99	
Atopobium parvulum		4.99	0.25		99	1	
47735142	**Low discrimination**						**47735142**
Bifidobacterium adolescentis		62.13	0.60	ARA 87			
Actinomyces israelii		29.07	0.61	SOR 11			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium longum subsp. infantis		3.86	0.26			
47735150	**Low discrimination**					**47735150**
					SPO	
Actinomyces israelii		68.34	0.44	CEL 93	1	
Bifidobacterium adolescentis		18.10	0.28	MLZ 13	1	
Clostridium butyricum		11.57	0.10		99	
47735152	**Good identification**					**47735152**
Bifidobacterium adolescentis		92.99	0.62	ARA 87		
Actinomyces israelii		5.32	0.47	SOR 11		
Clostridium clostridioforme		0.64	0.22			
47735160	**Low discrimination**					**47735160**
					SPO	
Actinomyces israelii		28.05	0.37	CEL 93	1	
Clostridium butyricum		18.36	0.13	NAG 6	99	
Clostridium perfringens		17.02	0.21		84	
47735162	**Low discrimination**					**47735162**
					GRA	
Parabacteroides distasonis		71.58	0.58	SOR 1	1	
Bifidobacterium longum subsp. infantis		15.60	0.30		99	
Actinomyces israelii		4.22	0.40		99	
47735172	**Low discrimination**					**47735172**
					GRA	
Parabacteroides distasonis		87.49	0.53	SOR 1	1	
Bifidobacterium breve		4.05	0.29		99	
Bifidobacterium adolescentis		3.96	0.28		99	
47735202	**Low discrimination**					**47735202**
					GRA	
Clostridium clostridioforme		89.60	0.50	ARA 94	50	
Actinomyces israelii		5.04	0.38		99	
Parabacteroides distasonis		3.42	0.32		1	
47735212	**Very good identification**					**47735212**
					GRA	
Clostridium clostridioforme		99.31	0.62	ARA 94	50	
Parabacteroides distasonis		0.45	0.27		1	
Actinomyces israelii		0.18	0.24		99	
47735222	**Low discrimination**					**47735222**
Parabacteroides distasonis		76.90	0.66	LAC 99		
Clostridium clostridioforme		20.37	0.50			
Prevotella oralis		2.56	0.28			
47735232	**Low discrimination**					**47735232**
Clostridium clostridioforme		69.14	0.62	ARA 94		
Parabacteroides distasonis		30.86	0.61			
47735300	**Good identification**					**47735300**
					SPO	
Eubacterium contortum		92.68	0.58	TRE 1	1	
Actinomyces israelii		2.27	0.46		1	
Clostridium butyricum		2.00	0.24		99	
47735302	**Low discrimination**					**47735302**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		42.17	0.66	NAG 99	1	
Clostridium clostridioforme		37.39	0.59		50	
Eubacterium contortum		12.08	0.45		99	
47735310	**Low discrimination**					**47735310**
Clostridium clostridioforme		57.40	0.36	ARA 94		
Eubacterium contortum		23.06	0.24	BGL 99		
Actinomyces israelii		8.35	0.32			
47735312	**Low discrimination**					**47735312**
					GRA	
Clostridium clostridioforme		86.74	0.71	ARA 94	50	
Parabacteroides distasonis		11.56	0.61		1	
Bifidobacterium adolescentis		1.41	0.43		99	
47735313	**Low discrimination**					**47735313**
					GRA	
Clostridium clostridioforme		86.99	0.36	ARA 94	50	
Parabacteroides distasonis		11.60	0.27	URE 1	1	
Bifidobacterium adolescentis		1.42	0.09		99	
47735320	**Low discrimination**					**47735320**
					GRA	
Parabacteroides distasonis		44.24	0.66	BGL 99	1	
Bacteroides vulgatus		40.68	0.47		1	
Eubacterium contortum		12.67	0.45		99	
47735322	**Very good identification**					**47735322**
Parabacteroides distasonis		98.44	1.00			
Clostridium clostridioforme		0.88	0.59			
Prevotella oralis		0.52	0.48			
47735323	**Very good identification**					**47735323**
Parabacteroides distasonis		98.48	0.66	URE 1		
Clostridium clostridioforme		0.88	0.25			
Prevotella oralis		0.52	0.14			
47735330	**Good identification**					**47735330**
Parabacteroides distasonis		91.04	0.61	BGL 99		
Clostridium clostridioforme		6.90	0.36			
Bacteroides vulgatus		1.57	0.13			
47735332	**Good identification**					**47735332**
Parabacteroides distasonis		92.93	0.95			
Clostridium clostridioforme		7.04	0.71			
Prevotella oralis		0.01	0.14			
47735333	**Good identification**					**47735333**
Parabacteroides distasonis		92.96	0.61	URE 1		
Clostridium clostridioforme		7.04	0.36			
47735342	**Low discrimination**					**47735342**
					GRA	
Actinomyces israelii		43.76	0.42	CEL 93	99	
Parabacteroides distasonis		19.08	0.32	RHA 7	1	
Clostridium clostridioforme		16.91	0.25		50	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47735352	**Low discrimination**						**47735352**
					GRA		
Clostridium clostridioforme		70.56	0.36	ARA 94	50		
Bifidobacterium adolescentis		14.05	0.28	SOR 1	99		
Parabacteroides distasonis		9.41	0.27		1		
47735362	**Very good identification**						**47735362**
Parabacteroides distasonis		98.33	0.66	SOR 1			
Clostridium clostridioforme		0.88	0.25				
Prevotella oralis		0.52	0.14				
47735372	**Good identification**						**47735372**
Parabacteroides distasonis		92.96	0.61	SOR 1			
Clostridium clostridioforme		7.04	0.36				
47735400	**Low discrimination**						**47735400**
					SPO		
Clostridium butyricum		78.15	0.42	CEL 99	99		
Actinomyces israelii		19.84	0.53	LAC 96	1		
Clostridium clostridioforme		1.05	0.22		66		
47735402	**Low discrimination**						**47735402**
					SPO		
Clostridium clostridioforme		45.06	0.56	RHA 88	66		
Bifidobacterium adolescentis		28.66	0.56		1		
Clostridium butyricum		13.17	0.35		99		
47735412	**Low discrimination**						**47735412**
Clostridium clostridioforme		84.42	0.67	RHA 88			
Bifidobacterium adolescentis		14.41	0.57				
Actinomyces israelii		0.79	0.42				
47735422	**Low discrimination**						**47735422**
					GRA	SPO	
Clostridium clostridioforme		70.40	0.56	RHA 88	50	66	
Parabacteroides distasonis		25.42	0.54		1	1	
Clostridium butyricum		1.79	0.17		99	99	
47735432	**Very good identification**						**47735432**
					GRA		
Clostridium clostridioforme		95.45	0.67	RHA 88	50		
Parabacteroides distasonis		4.07	0.49		1		
Bifidobacterium adolescentis		0.17	0.23		99		
47735440	**Low discrimination**						**47735440**
					SPO		
Actinomyces israelii		53.16	0.46	CEL 93	1		
Clostridium butyricum		46.84	0.24		99		
47735442	**Low discrimination**						**47735442**
					SPO		
Actinomyces israelii		49.37	0.49	CEL 93	1		
Bifidobacterium adolescentis		34.87	0.40		1		
Clostridium butyricum		11.28	0.17		99		
47735452	**Low discrimination**						**47735452**
Bifidobacterium adolescentis		60.03	0.42	LAC 99			
Clostridium clostridioforme		28.75	0.33	SOR 11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		10.39	0.35				
47735500	**Good identification**						**47735500**
					SPO		
Clostridium butyricum		91.61	0.66	CEL 99	99		
Actinomyces israelii		3.88	0.63		1		
Bifidobacterium adolescentis		3.23	0.56		1		
47735502	**Low discrimination**						**47735502**
					SPO		
Bifidobacterium adolescentis		83.92	0.90		1		
Clostridium butyricum		9.35	0.59		99		
Clostridium clostridioforme		4.46	0.65		66		
47735503	**Low discrimination**						**47735503**
					SPO		
Bifidobacterium adolescentis		84.01	0.56	URE 1	1		
Clostridium butyricum		9.36	0.25		99		
Clostridium clostridioforme		4.47	0.31		66		
47735510	**Low discrimination**						**47735510**
					SPO		
Clostridium butyricum		42.94	0.44	CEL 99	99		
Bifidobacterium adolescentis		35.16	0.57	MLZ 5	1		
Bifidobacterium longum subsp. longum		8.71	0.45		1		
47735512	**Low discrimination**						**47735512**
					SPO		
Bifidobacterium adolescentis		82.49	0.91		1		
Clostridium clostridioforme		16.35	0.76		66		
Clostridium butyricum		0.40	0.37		99		
47735513	**Low discrimination**						**47735513**
					SPO		
Bifidobacterium adolescentis		82.49	0.57	URE 1	1		
Clostridium clostridioforme		16.35	0.42		66		
Clostridium butyricum		0.40	0.03		99		
47735520	**Low discrimination**						**47735520**
					GRA	SPO	
Bacteroides vulgatus		48.98	0.50	TRE 4	1	1	
Clostridium butyricum		32.03	0.48	RHA 90	99	99	
Bifidobacterium longum subsp. longum		9.24	0.52		99	1	
47735522	**Low discrimination**						**47735522**
					GRA		
Parabacteroides distasonis		86.82	0.88		1		
Clostridium clostridioforme		8.13	0.65		50		
Bifidobacterium adolescentis		1.55	0.56		99		
47735523	**Low discrimination**						**47735523**
					GRA		
Parabacteroides distasonis		87.15	0.54	URE 1	1		
Clostridium clostridioforme		8.16	0.31		50		
Bifidobacterium adolescentis		1.55	0.22		99		
47735530	**Low discrimination**						**47735530**
					GRA		
Bifidobacterium longum subsp. longum		60.67	0.56	SAL 1	99		
Parabacteroides distasonis		15.31	0.49		1		
Clostridium clostridioforme		12.13	0.42		50		
47735532	**Low discrimination**						**47735532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		52.84	0.84		1		
Clostridium clostridioforme		41.87	0.76		50		
Bifidobacterium adolescentis		2.13	0.57		99		
47735533	**Low discrimination**						**47735533**
					GRA		
Parabacteroides distasonis		52.98	0.49	URE 1	1		
Clostridium clostridioforme		41.98	0.42		50		
Bifidobacterium adolescentis		2.14	0.23		99		
47735540	**Low discrimination**						**47735540**
					SPO		
Clostridium butyricum		78.01	0.48	CEL 99	99		
Actinomyces israelii		14.76	0.56	SOR 8	1		
Bifidobacterium adolescentis		3.91	0.40		1		
47735542	**Low discrimination**						**47735542**
					SPO		
Bifidobacterium adolescentis		87.61	0.74	SOR 11	1		
Clostridium butyricum		6.87	0.41		99		
Actinomyces israelii		5.01	0.59		1		
47735543	**Low discrimination**						**47735543**
					SPO		
Bifidobacterium adolescentis		88.06	0.40	SOR 11	1		
Clostridium butyricum		6.90	0.07	URE 1	99		
Actinomyces israelii		5.04	0.25		1		
47735552	**Very good identification**						**47735552**
Bifidobacterium adolescentis		97.18	0.76	SOR 11			
Clostridium clostridioforme		1.57	0.42				
Actinomyces israelii		0.68	0.45				
47735553	**Good identification**						**47735553**
Bifidobacterium adolescentis		98.41	0.42	SOR 11			
Clostridium clostridioforme		1.59	0.08	URE 1			
47735562	**Low discrimination**						**47735562**
					GRA	SPO	
Parabacteroides distasonis		64.15	0.54	SOR 1	1	1	
Bifidobacterium adolescentis		13.97	0.40		99	1	
Clostridium butyricum		9.43	0.23		99	99	
47735572	**Low discrimination**						**47735572**
					GRA		
Parabacteroides distasonis		41.89	0.49	SOR 1	1		
Clostridium clostridioforme		33.19	0.42		50		
Bifidobacterium adolescentis		20.70	0.42		99		
47735600	**Low discrimination**						**47735600**
					SPO		
Clostridium clostridioforme		53.84	0.37	BGL 99	66		
Eubacterium contortum		17.23	0.23		1		
Clostridium butyricum		16.90	0.16		99		
47735602	**Very good identification**						**47735602**
					GRA		
Clostridium clostridioforme		99.29	0.71		50		
Actinomyces israelii		0.29	0.37		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		0.15	0.28		1	
47735603	**Good identification**					**47735603**
Clostridium clostridioforme		100.0	0.37	URE 1		
47735610	**Good identification**					**47735610**
					SPO	
Clostridium clostridioforme		99.01	0.48	BGL 99	66	
Actinomyces israelii		0.63	0.20		1	
Clostridium butyricum		0.36	-.06		99	
47735612	**Very good identification**					**47735612**
					GRA	
Clostridium clostridioforme		99.95	0.82		50	
Bifidobacterium adolescentis		0.02	0.23		99	
Parabacteroides distasonis		0.02	0.23		1	
47735613	**Good identification**					**47735613**
Clostridium clostridioforme		100.0	0.48	URE 1		
47735622	**Low discrimination**					**47735622**
Clostridium clostridioforme		86.78	0.71			
Parabacteroides distasonis		12.82	0.62			
Bacteroides thetaiotaomicron		0.31	0.18			
47735623	**Low discrimination**					**47735623**
Clostridium clostridioforme		87.13	0.37	URE 1		
Parabacteroides distasonis		12.87	0.28			
47735630	**Good identification**					**47735630**
Clostridium clostridioforme		97.66	0.48	BGL 99		
Parabacteroides distasonis		1.71	0.23			
Bacteroides vulgatus		0.64	-.02			
47735632	**Very good identification**					**47735632**
Clostridium clostridioforme		98.25	0.82			
Parabacteroides distasonis		1.72	0.58			
Bacteroides thetaiotaomicron		0.03	0.11			
47735633	**Good identification**					**47735633**
Clostridium clostridioforme		98.28	0.48	URE 1		
Parabacteroides distasonis		1.72	0.23			
47735642	**Low discrimination**					**47735642**
Clostridium clostridioforme		89.84	0.37	SOR 1		
Actinomyces israelii		10.16	0.30			
47735652	**Good identification**					**47735652**
Clostridium clostridioforme		99.35	0.48	SOR 1		
Actinomyces israelii		0.37	0.16			
Bifidobacterium adolescentis		0.29	0.08			
47735662	**Low discrimination**					**47735662**
Clostridium clostridioforme		87.13	0.37	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		12.87	0.28			
47735672	**Good identification**					**47735672**
Clostridium clostridioforme		98.28	0.48	SOR 1		
Parabacteroides distasonis		1.72	0.23			
47735700	**Low discrimination**					**47735700**
					SPO	
Eubacterium contortum		67.59	0.57	TRE 1	1	
Clostridium butyricum		16.07	0.40		99	
Clostridium clostridioforme		7.14	0.46		66	
47735702	**Good identification**					**47735702**
					GRA	
Clostridium clostridioforme		90.89	0.80		50	
Parabacteroides distasonis		4.01	0.62		1	
Bifidobacterium adolescentis		2.36	0.56		99	
47735703	**Acceptable identification**					**47735703**
					GRA	
Clostridium clostridioforme		91.26	0.46	URE 1	50	
Parabacteroides distasonis		4.03	0.28		1	
Bifidobacterium adolescentis		2.36	0.22		99	
47735710	**Good identification**					**47735710**
					SPO	
Clostridium clostridioforme		92.66	0.57	BGL 99	66	
Clostridium butyricum		2.41	0.18		99	
Eubacterium contortum		1.94	0.23		1	
47735712	**Very good identification**					**47735712**
					GRA	
Clostridium clostridioforme		98.76	0.91		50	
Bifidobacterium adolescentis		0.69	0.57		99	
Parabacteroides distasonis		0.52	0.58		1	
47735713	**Very good identification**					**47735713**
					GRA	
Clostridium clostridioforme		98.80	0.57	URE 1	50	
Bifidobacterium adolescentis		0.69	0.23		99	
Parabacteroides distasonis		0.52	0.23		1	
47735720	**Good identification**					**47735720**
					GRA	
Bacteroides vulgatus		92.20	0.66	TRE 4	1	
Parabacteroides distasonis		4.63	0.62		1	
Eubacterium contortum		1.77	0.44		99	
47735722	**Low discrimination**					**47735722**
Parabacteroides distasonis		80.11	0.96			
Clostridium clostridioforme		18.34	0.80			
Bacteroides thetaiotaomicron		0.95	0.47			
47735723	**Low discrimination**					**47735723**
Parabacteroides distasonis		80.18	0.62	URE 1		
Clostridium clostridioforme		18.36	0.46			
Bacteroides thetaiotaomicron		0.95	0.13			
47735730	**Low discrimination**					**47735730**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		57.76	0.57	BGL 99			
Parabacteroides distasonis		29.82	0.58				
Bacteroides vulgatus		11.15	0.32				
47735732	**Low discrimination**						**47735732**
Clostridium clostridioforme		65.63	0.91				
Parabacteroides distasonis		33.88	0.92				
Bacteroides thetaiotaomicron		0.29	0.40				
47735733	**Low discrimination**						**47735733**
Clostridium clostridioforme		65.64	0.57	URE 1			
Parabacteroides distasonis		33.89	0.58				
Bacteroides thetaiotaomicron		0.29	0.06				
47735742	**Low discrimination**						**47735742**
Clostridium clostridioforme		61.58	0.46	SOR 1			
Bifidobacterium adolescentis		19.52	0.40				
Actinomyces israelii		8.32	0.40				
47735752	**Low discrimination**						**47735752**
					COC		
Clostridium clostridioforme		89.94	0.57	SOR 1	1		
Bifidobacterium adolescentis		7.65	0.42		1		
Blautia producta		1.39	0.15		99		
47735762	**Low discrimination**						**47735762**
Parabacteroides distasonis		79.10	0.62	SOR 1			
Clostridium clostridioforme		18.11	0.46				
Bacteroides ovatus		1.42	0.16				
47735772	**Low discrimination**						**47735772**
Clostridium clostridioforme		65.16	0.57	SOR 1			
Parabacteroides distasonis		33.64	0.58				
Bacteroides ovatus		0.75	0.13				
47736002	**Good identification**						**47736002**
Actinomyces israelii		100.0	0.36	CEL 93			
47736100	**Acceptable identification**						**47736100**
					GRA	SPO	
Actinomyces israelii		90.13	0.44	CEL 93	99	1	
Fusobacterium mortiferum		4.58	0.19	GAL 99	1	1	
Clostridium butyricum		3.80	-.01		99	99	
47736102	**Low discrimination**						**47736102**
Actinomyces israelii		50.45	0.47	CEL 93			
Bifidobacterium adolescentis		48.99	0.40	GAL 99			
Actinomyces naeslundii		0.56	0.22				
47736112	**Low discrimination**						**47736112**
Bifidobacterium adolescentis		84.94	0.42	GAL 99			
Actinomyces israelii		10.70	0.33	ARA 87			
Clostridium clostridioforme		2.93	0.13				
47736140	**Good identification**						**47736140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.37	CEL 93		
47736142	**Low discrimination**					**47736142**
Actinomyces israelii		76.42	0.40	CEL 93		
Bifidobacterium adolescentis		23.58	0.25	GAL 99		
47736500	**Low discrimination**					**47736500**
					SPO	
Actinomyces israelii		63.14	0.42	CEL 93	1	
Clostridium butyricum		29.27	0.15	GAL 99	99	
Bifidobacterium adolescentis		7.60	0.20		1	
47736502	**Low discrimination**					**47736502**
Bifidobacterium adolescentis		85.78	0.54	GAL 99		
Actinomyces israelii		10.80	0.45			
Clostridium clostridioforme		1.86	0.23			
47736512	**Low discrimination**					**47736512**
Bifidobacterium adolescentis		74.30	0.56	GAL 99		
Bifidobacterium dentium		18.56	0.27			
Clostridium clostridioforme		6.00	0.34			
47736542	**Low discrimination**					**47736542**
					COC	
Bifidobacterium adolescentis		67.28	0.39	GAL 99	1	
Actinomyces israelii		26.65	0.38	SOR 11	1	
Blautia producta		6.07	0.07		99	
47736552	**Low discrimination**					**47736552**
					COC	
Bifidobacterium adolescentis		77.51	0.40	GAL 99	1	
Blautia producta		17.16	0.15	SOR 11	99	
Actinomyces israelii		3.75	0.24		1	
47736612	**Good identification**					**47736612**
Clostridium clostridioforme		100.0	0.40	MAN 1		
47736632	**Good identification**					**47736632**
Clostridium clostridioforme		100.0	0.40	MAN 1		
47736702	**Low discrimination**					**47736702**
Clostridium clostridioforme		89.03	0.37	MAN 1		
Bifidobacterium adolescentis		5.66	0.20			
Actinomyces israelii		5.31	0.26			
47736712	**Good identification**					**47736712**
Clostridium clostridioforme		97.91	0.49	MAN 1		
Bifidobacterium adolescentis		1.67	0.22			
Bifidobacterium dentium		0.42	-.07			
47736722	**Low discrimination**					**47736722**
Clostridium clostridioforme		86.85	0.37	MAN 1		
Parabacteroides distasonis		11.50	0.28			
Bacteroides ovatus		1.65	-.02			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47736732	**Good identification**					**47736732**
					GRA	
Clostridium clostridioforme		97.47	0.49	MAN 1	50	
Parabacteroides distasonis		1.53	0.23		1	
Bifidobacterium dentium		0.74	-.03		99	
47737000	**Very good identification**					**47737000**
					SPO	
Actinomyces israelii		98.62	0.68	CEL 93	1	
Clostridium butyricum		0.69	0.10		99	
Actinomyces naeslundii		0.56	0.38		1	
47737002	**Very good identification**					**47737002**
Actinomyces israelii		95.71	0.71	CEL 93		
Bifidobacterium adolescentis		3.76	0.40			
Actinomyces naeslundii		0.36	0.38			
47737003	**Low discrimination**					**47737003**
Actinomyces israelii		64.24	0.36	CEL 93		
Actinomyces naeslundii		35.76	0.41	URE 1		
47737010	**Good identification**					**47737010**
Actinomyces israelii		99.52	0.53	CEL 93		
Bifidobacterium adolescentis		0.48	0.07	MLZ 13		
47737012	**Low discrimination**					**47737012**
Actinomyces israelii		74.84	0.56	CEL 93		
Bifidobacterium adolescentis		24.02	0.42	MLZ 13		
Clostridium clostridioforme		0.74	0.13			
47737020	**Acceptable identification**					**47737020**
Actinomyces israelii		91.90	0.47	CEL 93		
Actinomyces naeslundii		8.10	0.38	NAG 6		
47737021	**Good identification**					**47737021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47737022	**Low discrimination**					**47737022**
					GRA	
Actinomyces israelii		88.73	0.50	CEL 93	99	
Actinomyces naeslundii		5.21	0.38	NAG 6	99	
Parabacteroides distasonis		4.17	0.24		1	
47737023	**Low discrimination**					**47737023**
Actinomyces naeslundii		89.71	0.41	ESL 12		
Actinomyces israelii		10.29	0.16	XYL 1		
47737032	**Low discrimination**					**47737032**
					GRA	
Actinomyces israelii		65.51	0.36	CEL 93	99	
Parabacteroides distasonis		11.10	0.19	MLZ 13	1	
Clostridium clostridioforme		10.18	0.13	NAG 6	50	
47737040	**Very good identification**					**47737040**
Actinomyces israelii		99.75	0.60	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.26	0.25			
47737042	**Very good identification**					**47737042**
Actinomyces israelii		98.60	0.63	CEL 93		
Bifidobacterium adolescentis		1.23	0.25			
Actinomyces naeslundii		0.17	0.25			
47737050	**Good identification**					**47737050**
					COC	
Actinomyces israelii		97.98	0.46	CEL 93	1	
Blautia producta		2.02	-.03	MLZ 13	99	
47737052	**Acceptable identification**					**47737052**
Actinomyces israelii		90.75	0.49	CEL 93		
Bifidobacterium adolescentis		9.26	0.26	MLZ 13		
47737060	**Good identification**					**47737060**
Actinomyces israelii		96.16	0.40	CEL 93		
Actinomyces naeslundii		3.84	0.25	NAG 6		
47737062	**Good identification**					**47737062**
Actinomyces israelii		97.40	0.43	CEL 93		
Actinomyces naeslundii		2.60	0.25	NAG 6		
47737100	**Good identification**					**47737100**
					SPO	
Actinomyces israelii		93.51	0.78	CEL 93	1	
Clostridium butyricum		3.94	0.33		99	
Bifidobacterium adolescentis		1.38	0.40		1	
47737101	**Low discrimination**					**47737101**
					SPO	
Actinomyces israelii		61.12	0.44	CEL 93	1	
Actinomyces naeslundii		36.31	0.49	URE 1	1	
Clostridium butyricum		2.58	-.01		99	
47737102	**Low discrimination**					**47737102**
					SPO	
Actinomyces israelii		50.04	0.81	CEL 93	1	
Bifidobacterium adolescentis		48.59	0.74		1	
Clostridium butyricum		0.55	0.26		99	
47737103	**Low discrimination**					**47737103**
Actinomyces israelii		42.25	0.47	CEL 93		
Bifidobacterium adolescentis		41.02	0.40	URE 1		
Actinomyces naeslundii		16.73	0.49			
47737110	**Low discrimination**					**47737110**
Actinomyces israelii		85.55	0.64	CEL 93		
Bifidobacterium adolescentis		10.30	0.42	MLZ 13		
Bifidobacterium dentium		1.70	0.10			
47737112	**Low discrimination**					**47737112**
Bifidobacterium adolescentis		87.08	0.76	ARA 87		
Actinomyces israelii		10.96	0.67			
Bifidobacterium dentium		1.47	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47737113	**Low discrimination**						**47737113**
Bifidobacterium adolescentis		86.18	0.42	ARA 87			
Actinomyces israelii		10.85	0.33	URE 1			
Actinomyces naeslundii		1.51	0.27				
47737120	**Low discrimination**						**47737120**
					SPO		
Actinomyces israelii		77.08	0.57	CEL 93	1		
Actinomyces naeslundii		4.83	0.46	NAG 6	1		
Clostridium butyricum		4.43	0.15		99		
47737121	**Acceptable identification**						**47737121**
Actinomyces naeslundii		90.30	0.49	ESL 12			
Actinomyces israelii		9.70	0.23	XYL 1			
47737122	**Low discrimination**						**47737122**
					GRA		
Parabacteroides distasonis		39.60	0.58	MAN 1	1		
Actinomyces israelii		34.01	0.60		99		
Bifidobacterium longum subsp. infantis		8.63	0.30		99		
47737123	**Low discrimination**						**47737123**
					GRA		
Actinomyces naeslundii		71.96	0.49	ESL 12	99		
Parabacteroides distasonis		13.50	0.24	XYL 1	1		
Actinomyces israelii		11.60	0.26		99		
47737130	**Low discrimination**						**47737130**
					SPO		
Actinomyces israelii		45.50	0.43	CEL 93	1		
Bifidobacterium dentium		25.21	0.15	MLZ 13	1		
Clostridium tertium		22.30	0.24	NAG 6	99		
47737132	**Low discrimination**						**47737132**
					GRA		
Parabacteroides distasonis		36.41	0.53	MAN 1	1		
Bifidobacterium dentium		32.41	0.32		99		
Bifidobacterium adolescentis		10.91	0.42		99		
47737140	**Very good identification**						**47737140**
					COC	SPO	
Actinomyces israelii		96.03	0.71	CEL 93	1	1	
Blautia producta		2.44	0.24		99	1	
Clostridium butyricum		0.91	0.15		1	99	
47737141	**Low discrimination**						**47737141**
Actinomyces israelii		78.77	0.37	CEL 93			
Actinomyces naeslundii		21.24	0.36	URE 1			
47737142	**Low discrimination**						**47737142**
Actinomyces israelii		75.94	0.74	CEL 93			
Bifidobacterium adolescentis		23.44	0.59				
Bifidobacterium breve		0.20	0.24				
47737143	**Low discrimination**						**47737143**
Actinomyces israelii		67.19	0.40	CEL 93			
Bifidobacterium adolescentis		20.74	0.25	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		12.08	0.36			
47737150	**Low discrimination**					**47737150**
					COC	
Actinomyces israelii		64.46	0.57	CEL 93	1	
Blautia producta		32.86	0.32	MLZ 13	99	
Bifidobacterium adolescentis		2.47	0.26		1	
47737152	**Low discrimination**					**47737152**
					COC	
Bifidobacterium adolescentis		70.44	0.60	ARA 87	1	
Actinomyces israelii		27.91	0.60	SOR 11	1	
Blautia producta		1.05	0.15		99	
47737160	**Acceptable identification**					**47737160**
					COC	
Actinomyces israelii		91.19	0.50	CEL 93	1	
Blautia producta		4.03	0.07	NAG 6	99	
Actinomyces naeslundii		2.59	0.33		1	
47737161	**Low discrimination**					**47737161**
Actinomyces naeslundii		80.86	0.36	ESL 12		
Actinomyces israelii		19.15	0.16	XYL 1		
47737162	**Low discrimination**					**47737162**
Actinomyces israelii		84.80	0.53	CEL 93		
Bifidobacterium breve		6.33	0.29	NAG 6		
Bifidobacterium adolescentis		4.14	0.25			
47737163	**Low discrimination**					**47737163**
Actinomyces naeslundii		73.80	0.36	ESL 12		
Actinomyces israelii		26.21	0.19	XYL 1		
47737170	**Low discrimination**					**47737170**
					COC	
Actinomyces israelii		50.98	0.36	CEL 93	1	
Blautia producta		45.25	0.15	MLZ 13	99	
Bifidobacterium breve		3.78	0.12	NAG 6	1	
47737172	**Low discrimination**					**47737172**
Actinomyces israelii		46.06	0.39	CEL 93		
Bifidobacterium breve		23.02	0.29	MLZ 13		
Bifidobacterium adolescentis		18.40	0.26	NAG 6		
47737200	**Good identification**					**47737200**
Actinomyces israelii		100.0	0.48	CEL 93		
47737202	**Good identification**					**47737202**
Actinomyces israelii		96.93	0.51	CEL 93		
Clostridium clostridioforme		3.07	0.16	RHA 7		
47737212	**Low discrimination**					**47737212**
Actinomyces israelii		49.77	0.37	CEL 93		
Clostridium clostridioforme		48.09	0.27	RHA 7		
Bifidobacterium adolescentis		2.14	0.07	MLZ 13		
47737240	**Good identification**					**47737240**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.41	CEL 93			
47737242	**Good identification**						**47737242**
Actinomyces israelii		100.0	0.44	CEL 93			
47737300	**Good identification**						**47737300**
					SPO		
Actinomyces israelii		91.73	0.59	CEL 93	1		
Eubacterium contortum		6.68	0.24	RHA 7	1		
Clostridium butyricum		1.59	0.07		99		
47737302	**Low discrimination**						**47737302**
					GRA		
Actinomyces israelii		83.24	0.62	CEL 93	99		
Bifidobacterium adolescentis		10.85	0.40	RHA 7	99		
Parabacteroides distasonis		2.49	0.32		1		
47737310	**Good identification**						**47737310**
					COC		
Actinomyces israelii		95.90	0.44	CEL 93	1		
Blautia producta		2.55	-.03	RHA 7	99		
Bifidobacterium adolescentis		1.55	0.07	MLZ 13	1		
47737312	**Low discrimination**						**47737312**
Bifidobacterium adolescentis		35.53	0.42	RHA 1			
Actinomyces israelii		33.33	0.47	ARA 87			
Clostridium clostridioforme		26.95	0.36				
47737320	**Low discrimination**						**47737320**
					SPO	GRA	
Actinomyces israelii		28.81	0.38	CEL 93	1	99	
Clostridium ramosum		26.50	0.21	RHA 7	92	50	
Parabacteroides distasonis		20.26	0.32	NAG 6	1	1	
47737322	**Good identification**						**47737322**
					SPO	GRA	
Parabacteroides distasonis		91.52	0.66	MAN 1	1	1	
Clostridium ramosum		4.84	0.32		92	50	
Actinomyces israelii		1.97	0.41		1	99	
47737332	**Good identification**						**47737332**
					GRA		
Parabacteroides distasonis		91.73	0.61	MAN 1	1		
Clostridium clostridioforme		6.95	0.36		50		
Actinomyces israelii		0.55	0.27		99		
47737340	**Low discrimination**						**47737340**
					COC		
Actinomyces israelii		74.77	0.52	CEL 93	1		
Blautia producta		25.23	0.24	RHA 7	99		
47737342	**Acceptable identification**						**47737342**
					COC		
Actinomyces israelii		93.77	0.55	CEL 93	1		
Bifidobacterium adolescentis		3.88	0.25	RHA 7	1		
Blautia producta		2.34	0.07		99		
47737352	**Low discrimination**						**47737352**
					COC		
Actinomyces israelii		54.33	0.40	CEL 93	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		27.26	0.15	RHA 7	99		
Bifidobacterium adolescentis		18.41	0.26	MLZ 13	1		
47737400	**Good identification**						**47737400**
					SPO		
Actinomyces israelii		92.41	0.66	CEL 93	1		
Clostridium butyricum		7.14	0.26		99		
Bifidobacterium adolescentis		0.45	0.20		1		
47737402	**Low discrimination**						**47737402**
					SPO		
Actinomyces israelii		74.18	0.69	CEL 93	1		
Bifidobacterium adolescentis		23.80	0.54		1		
Clostridium butyricum		1.49	0.19		99		
47737410	**Low discrimination**						**47737410**
Actinomyces israelii		85.18	0.52	CEL 93			
Bifidobacterium dentium		8.28	0.10	MLZ 13			
Bifidobacterium adolescentis		3.39	0.22				
47737412	**Low discrimination**						**47737412**
Bifidobacterium adolescentis		58.63	0.56	LAC 99			
Actinomyces israelii		22.34	0.55				
Bifidobacterium dentium		14.64	0.27				
47737420	**Acceptable identification**						**47737420**
					SPO		
Actinomyces israelii		90.48	0.46	CEL 93	1		
Clostridium butyricum		9.52	0.08	NAG 6	99		
47737422	**Low discrimination**						**47737422**
Actinomyces israelii		80.75	0.49	CEL 93			
Clostridium clostridioforme		7.88	0.22	NAG 6			
Bifidobacterium adolescentis		4.10	0.20				
47737440	**Low discrimination**						**47737440**
					COC	SPO	
Actinomyces israelii		87.61	0.59	CEL 93	1	1	
Blautia producta		10.88	0.24		99	1	
Clostridium butyricum		1.51	0.08		1	99	
47737442	**Low discrimination**						**47737442**
					COC		
Actinomyces israelii		89.63	0.62	CEL 93	1		
Bifidobacterium adolescentis		9.14	0.39		1		
Blautia producta		0.82	0.07		99		
47737452	**Low discrimination**						**47737452**
					COC		
Actinomyces israelii		49.12	0.48	CEL 93	1		
Bifidobacterium adolescentis		40.98	0.40	MLZ 13	1		
Blautia producta		9.07	0.15		99		
47737460	**Low discrimination**						**47737460**
					COC		
Actinomyces israelii		82.23	0.39	CEL 93	1		
Blautia producta		17.77	0.07	NAG 6	99		
47737462	**Good identification**						**47737462**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	100.0	0.42	CEL 93		
47737500	**Low discrimination**					**47737500**
					SPO	
	Actinomyces israelii	62.22	0.76	CEL 93	1	
	Clostridium butyricum	28.84	0.50		99	
	Bifidobacterium adolescentis	7.49	0.54		1	
47737501	**Low discrimination**					**47737501**
					SPO	
	Actinomyces israelii	60.08	0.42	CEL 93	1	
	Clostridium butyricum	27.85	0.15	URE 1	99	
	Bifidobacterium adolescentis	7.23	0.20		1	
47737502	**Low discrimination**					**47737502**
					SPO	
	Bifidobacterium adolescentis	87.33	0.88		1	
	Actinomyces israelii	11.00	0.79		1	
	Clostridium butyricum	1.32	0.43		99	
47737503	**Low discrimination**					**47737503**
					SPO	
	Bifidobacterium adolescentis	87.40	0.54	URE 1	1	
	Actinomyces israelii	11.01	0.45		1	
	Clostridium butyricum	1.32	0.08		99	
47737510	**Low discrimination**					**47737510**
	Bifidobacterium dentium	49.95	0.44	CEL 99		
	Actinomyces israelii	20.75	0.62	BGL 91		
	Bifidobacterium adolescentis	20.43	0.56			
47737512	**Low discrimination**					**47737512**
	Bifidobacterium adolescentis	78.83	0.90			
	Bifidobacterium dentium	19.69	0.62			
	Actinomyces israelii	1.21	0.65			
47737513	**Low discrimination**					**47737513**
	Bifidobacterium adolescentis	78.89	0.56	URE 1		
	Bifidobacterium dentium	19.70	0.27			
	Actinomyces israelii	1.21	0.31			
47737520	**Low discrimination**					**47737520**
					SPO	
	Actinomyces israelii	52.11	0.56	CEL 93	1	
	Clostridium butyricum	32.91	0.31	NAG 6	99	
	Bifidobacterium dentium	5.27	0.15		1	
47737522	**Low discrimination**					**47737522**
					GRA	
	Bifidobacterium adolescentis	30.71	0.54	NAG 1	99	
	Actinomyces israelii	24.43	0.59		99	
	Parabacteroides distasonis	21.31	0.54		1	
47737530	**Good identification**					**47737530**
					COC	
	Bifidobacterium dentium	96.91	0.49	CEL 99	1	
	Actinomyces israelii	1.45	0.42	BGL 91	1	
	Blautia producta	0.61	0.15		99	
47737532	**Very good identification**					**47737532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium dentium		96.33	0.66	CEL 99	99		
Bifidobacterium adolescentis		2.19	0.56		99		
Parabacteroides distasonis		0.67	0.49		1		
47737540	**Low discrimination**						**47737540**
					COC	SPO	
Blautia producta		72.58	0.58	CEL 99	99	1	
Actinomyces israelii		23.62	0.69		1	1	
Clostridium butyricum		2.45	0.31		1	99	
47737542	**Low discrimination**						**47737542**
					COC		
Bifidobacterium adolescentis		66.77	0.73	SOR 11	1		
Actinomyces israelii		26.45	0.72		1		
Blautia producta		6.02	0.41		99		
47737543	**Low discrimination**						**47737543**
					COC		
Bifidobacterium adolescentis		67.28	0.39	SOR 11	1		
Actinomyces israelii		26.65	0.38	URE 1	1		
Blautia producta		6.07	0.07		99		
47737550	**Very good identification**						**47737550**
					COC		
Blautia producta		97.70	0.66	CEL 99	99		
Actinomyces israelii		1.58	0.55		1		
Bifidobacterium adolescentis		0.50	0.40		1		
47737552	**Low discrimination**						**47737552**
					COC		
Bifidobacterium adolescentis		77.46	0.74	SOR 11	1		
Blautia producta		17.15	0.50		99		
Actinomyces israelii		3.75	0.58		1		
47737553	**Low discrimination**						**47737553**
					COC		
Bifidobacterium adolescentis		77.51	0.40	SOR 11	1		
Blautia producta		17.16	0.15	URE 1	99		
Actinomyces israelii		3.75	0.24		1		
47737560	**Low discrimination**						**47737560**
					COC	SPO	
Blautia producta		81.49	0.41	CEL 99	99	1	
Actinomyces israelii		15.24	0.49	NAG 10	1	1	
Clostridium butyricum		2.15	0.13		1	99	
47737562	**Low discrimination**						**47737562**
					COC		
Actinomyces israelii		53.40	0.52	CEL 93	1		
Bifidobacterium adolescentis		21.33	0.39	NAG 6	1		
Blautia producta		21.16	0.25		99		
47737570	**Good identification**						**47737570**
					COC		
Blautia producta		97.37	0.50	CEL 99	99		
Bifidobacterium dentium		1.58	0.15	NAG 10	1		
Actinomyces israelii		0.91	0.35		1		
47737600	**Good identification**						**47737600**
					SPO		
Actinomyces israelii		96.92	0.47	CEL 93	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		3.08	0.00	RHA 7	99		
47737602	**Low discrimination**						**47737602**
Actinomyces israelii		60.31	0.50	CEL 93			
Clostridium clostridioforme		36.60	0.37	RHA 7			
Bifidobacterium adolescentis		2.60	0.20				
47737612	**Acceptable identification**						**47737612**
Clostridium clostridioforme		92.78	0.48	MAN 1			
Actinomyces israelii		5.02	0.36				
Bifidobacterium adolescentis		1.77	0.22				
47737622	**Low discrimination**						**47737622**
					GRA		
Clostridium clostridioforme		78.49	0.37	MAN 1	50		
Parabacteroides distasonis		11.59	0.28		1		
Actinomyces israelii		8.26	0.29		99		
47737632	**Good identification**						**47737632**
					GRA		
Clostridium clostridioforme		96.86	0.48	MAN 1	50		
Parabacteroides distasonis		1.69	0.23		1		
Bifidobacterium dentium		0.82	-.03		99		
47737642	**Low discrimination**						**47737642**
					COC		
Actinomyces israelii		89.11	0.43	CEL 93	1		
Blautia producta		10.89	0.07	RHA 7	99		
47737700	**Low discrimination**						**47737700**
					GRA	SPO	
Actinomyces israelii		61.17	0.57	CEL 93	99	1	
Clostridium butyricum		11.65	0.24	RHA 7	99	99	
Mitsuokella multacida		11.04	0.32		1	1	
47737702	**Low discrimination**						**47737702**
Bifidobacterium adolescentis		39.68	0.54	RHA 1			
Actinomyces israelii		37.23	0.60				
Clostridium clostridioforme		18.91	0.46				
47737712	**Low discrimination**						**47737712**
Clostridium clostridioforme		55.78	0.57	MAN 1			
Bifidobacterium adolescentis		31.40	0.56				
Bifidobacterium dentium		7.84	0.27				
47737722	**Low discrimination**						**47737722**
Parabacteroides distasonis		70.07	0.62	MAN 1			
Clostridium clostridioforme		16.04	0.46				
Bacteroides ovatus		10.07	0.32				
47737732	**Low discrimination**						**47737732**
					GRA		
Clostridium clostridioforme		53.36	0.57	MAN 1	50		
Parabacteroides distasonis		27.55	0.58		1		
Bifidobacterium dentium		13.34	0.32		99		
47737740	**Very good identification**						**47737740**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		96.44	0.58	CEL 99	99	99	
Actinomyces israelii		2.36	0.50		1	99	
Mitsuokella multacida		1.10	0.32		1	1	
47737742	**Low discrimination**						**47737742**
					COC		
Blautia producta		68.40	0.41	CEL 99	99		
Actinomyces israelii		22.62	0.53	BGL 10	1		
Bifidobacterium adolescentis		7.66	0.39		1		
47737750	**Very good identification**						**47737750**
					COC	GRA	
Blautia producta		99.75	0.66	CEL 99	99	99	
Mitsuokella multacida		0.13	0.24		1	1	
Actinomyces israelii		0.12	0.36		1	99	
47737752	**Acceptable identification**						**47737752**
					COC		
Blautia producta		93.38	0.50	CEL 99	99		
Bifidobacterium adolescentis		4.26	0.40	BGL 10	1		
Actinomyces israelii		1.54	0.39		1		
47737760	**Good identification**						**47737760**
					COC	GRA	
Blautia producta		95.83	0.41	CEL 99	99	99	
Mitsuokella multacida		2.45	0.21	NAG 10	1	1	
Actinomyces israelii		1.35	0.30		1	99	
47737770	**Good identification**						**47737770**
					COC	GRA	
Blautia producta		99.65	0.50	CEL 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
Actinomyces israelii		0.07	0.15		1	99	
47740100	**Low discrimination**						**47740100**
					GRA		
Fusobacterium mortiferum		85.23	0.39	MLT 11	1		
Collinsella aerofaciens		8.76	-.03	FRU 99	99		
Actinomyces naeslundii		3.42	0.27		99		
47741001	**Good identification**						**47741001**
Actinomyces naeslundii		100.0	0.46	ESL 12			
47741003	**Good identification**						**47741003**
Actinomyces naeslundii		100.0	0.46	ESL 12			
47741021	**Good identification**						**47741021**
Actinomyces naeslundii		100.0	0.46	ESL 12			
47741022	**Low discrimination**						**47741022**
					GRA	SPO	
Prevotella oralis		84.72	0.45	FRU 99	1	1	
Clostridium barati		5.14	0.24	LAC 94	99	99	
Bifidobacterium breve		3.00	0.32		99	1	
47741023	**Low discrimination**						**47741023**
					GRA		
Actinomyces naeslundii		76.43	0.46	ESL 12	99		
Prevotella oralis		23.57	0.11	FRU 95	1		
47741101	**Acceptable identification**						**47741101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		91.56	0.53	ESL 12			
Collinsella aerofaciens		6.51	-.03	FRU 95			
Bifidobacterium breve		1.93	0.10				
47741102	**Low discrimination**						**47741102**
					GRA		
Bifidobacterium breve		53.90	0.62	FRU 99	99		
Prevotella oralis		22.57	0.44		1		
Lactobacillus acidophilus		8.84	0.38		99		
47741103	**Low discrimination**						**47741103**
					GRA		
Actinomyces naeslundii		73.91	0.53	ESL 12	99		
Bifidobacterium breve		15.77	0.27	FRU 95	99		
Prevotella oralis		6.60	0.10		1		
47741110	**Acceptable identification**						**47741110**
Bifidobacterium breve		94.39	0.44	FRU 99			
Collinsella aerofaciens		3.21	-.03	BGL 91			
Actinomyces naeslundii		1.59	0.29				
47741112	**Low discrimination**						**47741112**
					GRA		
Bifidobacterium breve		87.91	0.62	FRU 99	99		
Leptotrichia buccalis		7.40	0.24		1		
Bifidobacterium adolescentis		3.96	0.38		99		
47741120	**Low discrimination**						**47741120**
					GRA	SPO	
Clostridium paraputrificum		62.25	0.47	RAF 7	99	99	
Prevotella oralis		18.75	0.44	FRU 99	1	1	
Bifidobacterium breve		7.87	0.49		99	1	
47741121	**Low discrimination**						**47741121**
					GRA	SPO	
Actinomyces naeslundii		69.16	0.53	ESL 12	99	1	
Clostridium paraputrificum		20.52	0.13	FRU 95	99	99	
Prevotella oralis		6.18	0.10		1	1	
47741122	**Low discrimination**						**47741122**
					GRA	SPO	
Prevotella oralis		73.23	0.66	FRU 99	1	1	
Bifidobacterium breve		16.36	0.66		99	1	
Clostridium paraputrificum		5.48	0.40		99	99	
47741130	**Low discrimination**						**47741130**
					GRA	SPO	
Bifidobacterium breve		89.33	0.49	FRU 99	99	1	
Clostridium paraputrificum		7.14	0.13	BGL 91	99	99	
Prevotella oralis		2.15	0.10		1	1	
47741132	**Good identification**						**47741132**
					GRA		
Bifidobacterium breve		94.37	0.66	FRU 99	99		
Prevotella oralis		4.27	0.32		1		
Leptotrichia buccalis		0.50	0.07		1		
47741140	**Low discrimination**						**47741140**
Bifidobacterium breve		89.34	0.44	FRU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		5.03	0.38	BGL 91			
Collinsella aerofaciens		3.04	-.03				
47741141	**Low discrimination**						**47741141**
Actinomyces naeslundii		89.32	0.41	ESL 12			
Bifidobacterium breve		10.68	0.10	FRU 95			
47741142	**Very good identification**						**47741142**
Bifidobacterium breve		96.21	0.62	FRU 99			
Lactobacillus acidophilus		1.95	0.23				
Actinomyces naeslundii		0.54	0.38				
47741150	**Good identification**						**47741150**
Bifidobacterium breve		100.0	0.44	FRU 99			
47741152	**Very good identification**						**47741152**
					GRA		
Bifidobacterium breve		99.36	0.62	FRU 99	99		
Bifidobacterium adolescentis		0.55	0.22		99		
Leptotrichia buccalis		0.08	-.11		1		
47741160	**Low discrimination**						**47741160**
					SPO		
Bifidobacterium breve		59.80	0.49	FRU 99	1		
Clostridium paraputrificum		35.60	0.27	BGL 91	99		
Actinomyces naeslundii		1.89	0.38		1		
47741161	**Low discrimination**						**47741161**
					SPO		
Actinomyces naeslundii		74.68	0.41	ESL 12	1		
Bifidobacterium breve		15.87	0.15	FRU 95	1		
Clostridium paraputrificum		9.45	-.07		99		
47741162	**Good identification**						**47741162**
					GRA	SPO	
Bifidobacterium breve		91.66	0.66	FRU 99	99	1	
Prevotella oralis		4.14	0.32		1	1	
Clostridium paraputrificum		2.31	0.21		99	99	
47741170	**Good identification**						**47741170**
					SPO		
Bifidobacterium breve		99.40	0.49	FRU 99	1		
Clostridium paraputrificum		0.60	-.07	BGL 91	99		
47741172	**Very good identification**						**47741172**
					GRA		
Bifidobacterium breve		99.95	0.66	FRU 99	99		
Prevotella oralis		0.05	-.03		1		
47741222	**Good identification**						**47741222**
					GRA	SPO	
Prevotella oralis		97.93	0.39	FRU 99	1	1	
Clostridium barati		1.21	0.05	LAC 94	99	99	
Lactobacillus acidophilus		0.48	0.01		99	1	
47741302	**Low discrimination**						**47741302**
					GRA		
Prevotella oralis		87.05	0.38	FRU 99	1		
Lactobacillus acidophilus		5.99	0.19	NAG 95	99		
Bifidobacterium breve		4.90	0.27		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47741320	**Low discrimination**						**47741320**
					SPO	GRA	
Prevotella oralis		81.49	0.38	FRU 99	1	1	
Clostridium ramosum		9.63	0.22	BGL 95	92	50	
Clostridium paraputrificum		6.38	0.13		99	99	
47741322	**Very good identification**						**47741322**
					SPO		
Prevotella oralis		95.80	0.60	FRU 99	1		
Clostridium ramosum		2.38	0.33		92		
Parabacteroides distasonis		0.53	0.34		1		
47741502	**Low discrimination**						**47741502**
					SPO		
Bifidobacterium adolescentis		88.48	0.50	FRU 99	1		
Clostridium butyricum		4.05	0.13	XYL 87	99		
Bifidobacterium breve		3.62	0.27		1		
47741512	**Acceptable identification**						**47741512**
Bifidobacterium adolescentis		94.78	0.52	FRU 99			
Bifidobacterium breve		3.18	0.27	XYL 87			
Clostridium clostridioforme		1.78	0.19				
47741552	**Low discrimination**						**47741552**
Bifidobacterium adolescentis		78.68	0.36	FRU 99			
Bifidobacterium breve		21.32	0.27	XYL 87			
47743000	**Low discrimination**						**47743000**
Actinomyces naeslundii		80.40	0.48	ESL 12			
Actinomyces israelii		12.47	0.25	FRU 95			
Bifidobacterium breve		3.73	0.10				
47743001	**Good identification**						**47743001**
Actinomyces naeslundii		100.0	0.51	ESL 12			
47743002	**Low discrimination**						**47743002**
Actinomyces naeslundii		54.93	0.48	ESL 12			
Bifidobacterium breve		25.77	0.27	FRU 95			
Actinomyces israelii		12.78	0.28				
47743003	**Good identification**						**47743003**
Actinomyces naeslundii		100.0	0.51	ESL 12			
47743020	**Low discrimination**						**47743020**
					SPO		
Actinomyces naeslundii		85.72	0.48	ESL 12	1		
Bifidobacterium breve		7.07	0.15	FRU 95	1		
Clostridium ramosum		3.65	-.01		92		
47743021	**Good identification**						**47743021**
Actinomyces naeslundii		100.0	0.51	ESL 12			
47743022	**Low discrimination**						**47743022**
					GRA		
Actinomyces naeslundii		41.63	0.48	ESL 12	99		
Bifidobacterium breve		34.72	0.32	FRU 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		9.92	0.11		1	
47743023	**Good identification**					**47743023**
Actinomyces naeslundii		100.0	0.51	ESL 12		
47743041	**Good identification**					**47743041**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47743043	**Good identification**					**47743043**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47743061	**Good identification**					**47743061**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47743063	**Good identification**					**47743063**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47743100	**Low discrimination**					**47743100**
Bifidobacterium breve		55.34	0.44	FRU 99		
Actinomyces naeslundii		34.29	0.55	BGL 91		
Actinomyces israelii		7.47	0.35			
47743101	**Very good identification**					**47743101**
Actinomyces naeslundii		98.93	0.58	ESL 12		
Bifidobacterium breve		1.08	0.10	FRU 95		
47743102	**Low discrimination**					**47743102**
Bifidobacterium breve		88.31	0.62	FRU 99		
Actinomyces naeslundii		5.41	0.55			
Bifidobacterium adolescentis		2.66	0.35			
47743103	**Good identification**					**47743103**
Actinomyces naeslundii		90.10	0.58	ESL 12		
Bifidobacterium breve		9.90	0.27	FRU 95		
47743110	**Acceptable identification**					**47743110**
Bifidobacterium breve		94.99	0.44	FRU 99		
Actinomyces naeslundii		3.10	0.34	BGL 91		
Actinomyces israelii		1.92	0.21			
47743111	**Low discrimination**					**47743111**
Actinomyces naeslundii		82.88	0.37	ESL 12		
Bifidobacterium breve		17.12	0.10	FRU 95		
47743112	**Very good identification**					**47743112**
Bifidobacterium breve		95.64	0.62	FRU 99		
Bifidobacterium adolescentis		3.53	0.36			
Actinomyces naeslundii		0.31	0.34			
47743120	**Low discrimination**					**47743120**
					SPO	
Bifidobacterium breve		50.01	0.49	FRU 99	1	
Clostridium ramosum		25.80	0.33	BGL 91	92	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		17.43	0.55		1		
47743121	**Very good identification**						**47743121**
					SPO		
Actinomyces naeslundii		97.16	0.58	ESL 12	1		
Bifidobacterium breve		1.88	0.15	FRU 95	1		
Clostridium ramosum		0.97	-.01		92		
47743122	**Low discrimination**						**47743122**
					SPO	GRA	
Bifidobacterium breve		76.95	0.66	FRU 99	1	99	
Clostridium ramosum		15.70	0.43		92	50	
Prevotella oralis		3.48	0.32		1	1	
47743123	**Low discrimination**						**47743123**
					SPO		
Actinomyces naeslundii		80.38	0.58	ESL 12	1		
Bifidobacterium breve		15.70	0.32	FRU 95	1		
Clostridium ramosum		3.20	0.09		92		
47743130	**Good identification**						**47743130**
					SPO		
Bifidobacterium breve		96.83	0.49	FRU 99	1		
Actinomyces naeslundii		1.78	0.34	BGL 91	1		
Clostridium tertium		0.89	0.05		99		
47743131	**Low discrimination**						**47743131**
Actinomyces naeslundii		73.15	0.37	ESL 12			
Bifidobacterium breve		26.85	0.15	FRU 95			
47743132	**Very good identification**						**47743132**
					SPO		
Bifidobacterium breve		99.30	0.66	FRU 99	1		
Clostridium tertium		0.27	0.13		99		
Clostridium ramosum		0.21	0.09		92		
47743140	**Low discrimination**						**47743140**
Bifidobacterium breve		86.07	0.44	FRU 99			
Actinomyces naeslundii		9.41	0.43	BGL 91			
Actinomyces israelii		4.52	0.28				
47743141	**Acceptable identification**						**47743141**
Actinomyces naeslundii		94.20	0.46	ESL 12			
Bifidobacterium breve		5.80	0.10	FRU 95			
47743142	**Very good identification**						**47743142**
Bifidobacterium breve		97.71	0.62	FRU 99			
Actinomyces naeslundii		1.06	0.43				
Actinomyces israelii		0.76	0.31				
47743143	**Low discrimination**						**47743143**
Actinomyces naeslundii		61.63	0.46	ESL 12			
Bifidobacterium breve		38.38	0.27	FRU 95			
47743150	**Good identification**						**47743150**
Bifidobacterium breve		99.22	0.44	FRU 99			
Actinomyces israelii		0.78	0.14	BGL 91			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47743152	**Very good identification**						**47743152**
Bifidobacterium breve		99.43	0.62	FRU 99			
Bifidobacterium adolescentis		0.45	0.21				
Actinomyces israelii		0.12	0.17				
47743160	**Acceptable identification**						**47743160**
					SPO		
Bifidobacterium breve		93.22	0.49	FRU 99	1		
Actinomyces naeslundii		5.73	0.43	BGL 91	1		
Clostridium paraputrificum		0.56	-.07		99		
47743161	**Acceptable identification**						**47743161**
Actinomyces naeslundii		90.13	0.46	ESL 12			
Bifidobacterium breve		9.87	0.15	FRU 95			
47743162	**Very good identification**						**47743162**
					SPO		
Bifidobacterium breve		99.09	0.66	FRU 99	1		
Actinomyces naeslundii		0.60	0.43		1		
Clostridium ramosum		0.20	0.09		92		
47743170	**Good identification**						**47743170**
Bifidobacterium breve		100.0	0.49	FRU 99			
47743172	**Very good identification**						**47743172**
Bifidobacterium breve		100.0	0.66	FRU 99			
47743320	**Good identification**						**47743320**
					SPO	GRA	
Clostridium ramosum		97.10	0.35	FRU 99	92	50	
Prevotella oralis		1.47	0.04	TRE 94	1	1	
Bifidobacterium breve		1.43	0.15		1	99	
47743322	**Acceptable identification**						**47743322**
					SPO	GRA	
Clostridium ramosum		90.17	0.45	FRU 99	92	50	
Prevotella oralis		6.46	0.25	TRE 94	1	1	
Bifidobacterium breve		3.37	0.32		1	99	
47743502	**Low discrimination**						**47743502**
Bifidobacterium adolescentis		87.92	0.49	FRU 99			
Actinomyces israelii		7.14	0.36	XYL 87			
Bifidobacterium breve		4.40	0.27				
47743512	**Acceptable identification**						**47743512**
Bifidobacterium adolescentis		94.68	0.50	FRU 99			
Bifidobacterium breve		3.88	0.27	XYL 87			
Actinomyces israelii		0.94	0.22				
47744712	**Good identification**						**47744712**
Clostridium clostridioforme		100.0	0.43	FRU 99			
47744732	**Good identification**						**47744732**
Clostridium clostridioforme		100.0	0.43	FRU 99			
47745102	**Low discrimination**						**47745102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium adolescentis		54.56	0.50	FRU 99	99		
Bifidobacterium longum subsp. infantis		13.00	0.26	ARA 87	99		
Prevotella oralis		11.45	0.28		1		
47745112	**Acceptable identification**						**47745112**
Bifidobacterium adolescentis		92.97	0.52	FRU 99			
Bifidobacterium breve		3.11	0.27	ARA 87			
Actinomyces israelii		1.87	0.29				
47745122	**Low discrimination**						**47745122**
					CAT	GRA	
Prevotella oralis		74.39	0.50	FRU 99	1	1	
Bacteroides fragilis		8.69	0.25	XYL 11	99	1	
Bifidobacterium longum subsp. infantis		7.91	0.30		1	99	
47745132	**Low discrimination**						**47745132**
					GRA		
Parabacteroides distasonis		45.34	0.37	FRU 99	1		
Bifidobacterium breve		26.66	0.32		99		
Prevotella oralis		14.75	0.16		1		
47745152	**Low discrimination**						**47745152**
Bifidobacterium adolescentis		74.94	0.36	FRU 99			
Bifidobacterium breve		20.31	0.27	ARA 87			
Actinomyces israelii		4.75	0.22	SOR 11			
47745322	**Low discrimination**						**47745322**
Prevotella oralis		66.43	0.44	FRU 99			
Parabacteroides distasonis		26.81	0.50	XYL 11			
Bacteroides uniformis		5.04	0.19				
47745332	**Low discrimination**						**47745332**
Parabacteroides distasonis		74.21	0.45	FRU 99			
Clostridium clostridioforme		21.72	0.31				
Prevotella oralis		3.45	0.10				
47745500	**Good identification**						**47745500**
					SPO		
Clostridium butyricum		96.02	0.45	FRU 99	99		
Actinomyces israelii		2.27	0.39	TRE 94	1		
Bifidobacterium adolescentis		1.71	0.30		1		
47745502	**Low discrimination**						**47745502**
					SPO		
Bifidobacterium adolescentis		79.98	0.64	FRU 99	1		
Clostridium butyricum		17.67	0.38		99		
Actinomyces israelii		1.61	0.42		1		
47745512	**Very good identification**						**47745512**
					SPO		
Bifidobacterium adolescentis		96.06	0.66	FRU 99	1		
Clostridium clostridioforme		2.73	0.36		66		
Clostridium butyricum		0.91	0.17		99		
47745532	**Low discrimination**						**47745532**
					GRA		
Clostridium clostridioforme		53.43	0.36	FRU 99	50		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		19.02	0.32	TRE 88	99	
Parabacteroides distasonis		17.46	0.34	RHA 88	1	
47745542	**Low discrimination**					**47745542**
					SPO	
Bifidobacterium adolescentis		82.05	0.49	FRU 99	1	
Clostridium butyricum		12.75	0.20	SOR 11	99	
Actinomyces israelii		5.20	0.35		1	
47745552	**Good identification**					**47745552**
					SPO	
Bifidobacterium adolescentis		98.58	0.50	FRU 99	1	
Actinomyces israelii		0.76	0.21	SOR 11	1	
Clostridium butyricum		0.66	-.02		99	
47745612	**Good identification**					**47745612**
Clostridium clostridioforme		100.0	0.42	FRU 99		
47745632	**Good identification**					**47745632**
Clostridium clostridioforme		100.0	0.42	FRU 99		
47745702	**Low discrimination**					**47745702**
					SPO	
Clostridium clostridioforme		70.47	0.40	FRU 99	66	
Bifidobacterium adolescentis		12.75	0.30	TRE 88	1	
Clostridium butyricum		8.62	0.13		99	
47745712	**Good identification**					**47745712**
Clostridium clostridioforme		95.37	0.51	FRU 99		
Bifidobacterium adolescentis		4.63	0.32	TRE 88		
47745722	**Low discrimination**					**47745722**
Bacteroides uniformis		60.19	0.37	IND 98		
Parabacteroides distasonis		17.06	0.46	FRU 99		
Clostridium clostridioforme		15.09	0.40			
47745732	**Low discrimination**					**47745732**
Clostridium clostridioforme		83.18	0.51	FRU 99		
Parabacteroides distasonis		11.12	0.42	TRE 88		
Bacteroides ovatus		4.97	0.20			
47747000	**Good identification**					**47747000**
Actinomyces israelii		100.0	0.43	FRU 99		
47747002	**Good identification**					**47747002**
Actinomyces israelii		96.58	0.46	FRU 99		
Bifidobacterium adolescentis		3.42	0.15			
47747040	**Good identification**					**47747040**
Actinomyces israelii		100.0	0.36	FRU 99		
47747042	**Good identification**					**47747042**
Actinomyces israelii		100.0	0.39	FRU 99		
47747100	**Acceptable identification**					**47747100**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		91.31	0.53	FRU 99	1	
Clostridium butyricum		6.88	0.13		99	
Bifidobacterium adolescentis		1.21	0.15		1	
47747102	**Low discrimination**					**47747102**
Actinomyces israelii		51.41	0.56	FRU 99		
Bifidobacterium adolescentis		45.05	0.49			
Bifidobacterium breve		2.25	0.27			
47747110	**Low discrimination**					**47747110**
Actinomyces israelii		84.41	0.39	FRU 99		
Bifidobacterium adolescentis		9.17	0.16	MLZ 13		
Bifidobacterium breve		3.68	0.10			
47747112	**Low discrimination**					**47747112**
Bifidobacterium adolescentis		83.70	0.50	FRU 99		
Actinomyces israelii		11.68	0.42	ARA 87		
Bifidobacterium breve		3.43	0.27			
47747140	**Good identification**					**47747140**
					SPO	
Actinomyces israelii		96.75	0.46	FRU 99	1	
Clostridium butyricum		1.63	-.05		99	
Bifidobacterium breve		1.62	0.10		1	
47747142	**Low discrimination**					**47747142**
Actinomyces israelii		71.88	0.49	FRU 99		
Bifidobacterium adolescentis		20.02	0.33			
Bifidobacterium breve		8.11	0.27			
47747302	**Low discrimination**					**47747302**
Actinomyces israelii		89.48	0.37	FRU 99		
Bifidobacterium adolescentis		10.52	0.15	RHA 7		
47747400	**Low discrimination**					**47747400**
					SPO	
Actinomyces israelii		87.86	0.41	FRU 99	1	
Clostridium butyricum		12.14	0.05		99	
47747402	**Low discrimination**					**47747402**
					SPO	
Actinomyces israelii		75.44	0.44	FRU 99	1	
Bifidobacterium adolescentis		21.85	0.29		1	
Clostridium butyricum		2.70	-.02		99	
47747442	**Acceptable identification**					**47747442**
Actinomyces israelii		91.57	0.37	FRU 99		
Bifidobacterium adolescentis		8.43	0.13			
47747500	**Low discrimination**					**47747500**
					SPO	
Actinomyces israelii		51.60	0.52	FRU 99	1	
Clostridium butyricum		42.80	0.29		99	
Bifidobacterium adolescentis		5.61	0.29		1	
47747502	**Low discrimination**					**47747502**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		85.44	0.63	FRU 99	1		
Actinomyces israelii		11.92	0.55		1		
Clostridium butyricum		2.56	0.22		99		
47747510	**Low discrimination**						**47747510**
Actinomyces israelii		34.12	0.37	FRU 99			
Bifidobacterium adolescentis		30.31	0.30	MLZ 13			
Bifidobacterium dentium		23.26	0.10				
47747512	**Good identification**						**47747512**
Bifidobacterium adolescentis		91.18	0.64	FRU 99			
Bifidobacterium dentium		7.14	0.27				
Actinomyces israelii		1.56	0.40				
47747540	**Low discrimination**						**47747540**
					COC	SPO	
Actinomyces israelii		47.85	0.45	FRU 99	1	1	
Blautia producta		41.63	0.24		99	1	
Clostridium butyricum		8.87	0.11		1	99	
47747542	**Low discrimination**						**47747542**
					COC		
Bifidobacterium adolescentis		67.19	0.47	FRU 99	1		
Actinomyces israelii		29.49	0.48	SOR 11	1		
Blautia producta		1.90	0.07		99		
47747552	**Low discrimination**						**47747552**
					COC		
Bifidobacterium adolescentis		88.53	0.49	FRU 99	1		
Blautia producta		6.15	0.15	SOR 11	99		
Actinomyces israelii		4.75	0.33		1		
47747702	**Low discrimination**						**47747702**
					SPO		
Actinomyces israelii		47.26	0.35	FRU 99	1		
Bifidobacterium adolescentis		45.46	0.29	RHA 7	1		
Clostridium butyricum		4.18	-.04		99		
47750000	**Good identification**						**47750000**
					GRA	SPO	
Fusobacterium mortiferum		94.67	0.67	MLT 11	1	1	
Lactobacillus catenaformis		2.08	0.29		99	1	
Clostridium barati		1.22	0.19		99	99	
47750002	**Low discrimination**						**47750002**
					GRA	SPO	
Fusobacterium mortiferum		49.26	0.51	MLT 11	1	1	
Clostridium barati		17.07	0.28	BGL 10	99	99	
Lactobacillus catenaformis		9.74	0.29		99	1	
47750003	**Low discrimination**						**47750003**
					GRA		
Actinomyces naeslundii		84.44	0.44	ESL 12	99		
Fusobacterium mortiferum		10.01	0.17	GAL 96	1		
Actinomyces viscosus		5.55	-.01		99		
47750021	**Good identification**						**47750021**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47750022	**Low discrimination**						**47750022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Prevotella oralis		69.64	0.45	GAL 99	1	1	
Clostridium barati		22.02	0.36	LAC 94	99	99	
Bifidobacterium breve		2.46	0.32		99	1	
47750023	**Low discrimination**						**47750023**
					GRA	SPO	
Actinomyces naeslundii		66.11	0.44	ESL 12	99	1	
Prevotella oralis		25.75	0.11	GAL 96	1	1	
Clostridium barati		8.14	0.02		99	99	
47750100	**Low discrimination**						**47750100**
					GRA		
Fusobacterium mortiferum		84.24	0.73	MLT 11	1		
Collinsella aerofaciens		8.65	0.32		99		
Bifidobacterium breve		2.57	0.44		99		
47750101	**Low discrimination**						**47750101**
					GRA		
Actinomyces naeslundii		50.23	0.52	ESL 12	99		
Fusobacterium mortiferum		43.92	0.39	GAL 96	1		
Collinsella aerofaciens		4.51	-.03		99		
47750102	**Low discrimination**						**47750102**
					GRA		
Bifidobacterium breve		42.72	0.62	GAL 99	99		
Prevotella oralis		17.89	0.44		1		
Fusobacterium mortiferum		15.40	0.57		1		
47750103	**Low discrimination**						**47750103**
					GRA		
Actinomyces naeslundii		61.45	0.52	ESL 12	99		
Bifidobacterium breve		16.56	0.27	GAL 96	99		
Prevotella oralis		6.94	0.10		1		
47750110	**Low discrimination**						**47750110**
					GRA		
Bifidobacterium breve		72.08	0.44	GAL 99	99		
Fusobacterium mortiferum		23.89	0.39	BGL 91	1		
Collinsella aerofaciens		2.45	-.03		99		
47750112	**Low discrimination**						**47750112**
					GRA		
Bifidobacterium breve		86.63	0.62	GAL 99	99		
Leptotrichia buccalis		7.29	0.24		1		
Bifidobacterium adolescentis		3.90	0.38		99		
47750120	**Low discrimination**						**47750120**
					GRA	SPO	
Clostridium paraputrificum		60.60	0.47	RAF 7	99	99	
Prevotella oralis		18.25	0.44	GAL 99	1	1	
Bifidobacterium breve		7.66	0.49		99	1	
47750121	**Low discrimination**						**47750121**
					GRA	SPO	
Actinomyces naeslundii		63.36	0.52	ESL 12	99	1	
Clostridium paraputrificum		23.74	0.13	GAL 96	99	99	
Prevotella oralis		7.15	0.10		1	1	
47750122	**Low discrimination**						**47750122**
					GRA	SPO	
Prevotella oralis		70.81	0.66	GAL 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		15.82	0.66		99	1	
Clostridium paraputrificum		5.30	0.40		99	99	
47750130	**Low discrimination**						**47750130**
					GRA	SPO	
Bifidobacterium breve		89.23	0.49	GAL 99	99	1	
Clostridium paraputrificum		7.13	0.13	BGL 91	99	99	
Prevotella oralis		2.15	0.10		1	1	
47750132	**Good identification**						**47750132**
					GRA		
Bifidobacterium breve		93.60	0.66	GAL 99	99		
Prevotella oralis		4.23	0.32		1		
Clostridium clostridioforme		0.63	0.25		50		
47750140	**Low discrimination**						**47750140**
					GRA	SPO	
Bifidobacterium breve		67.70	0.44	GAL 99	99	1	
Fusobacterium mortiferum		22.44	0.39	BGL 91	1	1	
Clostridium barati		3.19	0.10		99	99	
47750141	**Low discrimination**						**47750141**
Actinomyces naeslundii		86.88	0.39	ESL 12			
Bifidobacterium breve		13.12	0.10	GAL 96			
47750142	**Good identification**						**47750142**
					SPO		
Bifidobacterium breve		94.95	0.62	GAL 99	1		
Lactobacillus acidophilus		1.93	0.23		1		
Clostridium barati		1.33	0.18		99		
47750150	**Good identification**						**47750150**
Bifidobacterium breve		100.0	0.44	GAL 99			
47750152	**Very good identification**						**47750152**
					GRA		
Bifidobacterium breve		99.36	0.62	GAL 99	99		
Bifidobacterium adolescentis		0.55	0.22		99		
Leptotrichia buccalis		0.08	-.11		1		
47750160	**Low discrimination**						**47750160**
					SPO		
Bifidobacterium breve		58.01	0.49	GAL 99	1		
Clostridium paraputrificum		34.54	0.27	BGL 91	99		
Clostridium barati		4.61	0.18		99		
47750161	**Low discrimination**						**47750161**
					SPO		
Actinomyces naeslundii		70.01	0.39	ESL 12	1		
Bifidobacterium breve		18.80	0.15	GAL 96	1		
Clostridium paraputrificum		11.19	-.07		99		
47750162	**Good identification**						**47750162**
					GRA	SPO	
Bifidobacterium breve		90.14	0.66	GAL 99	99	1	
Prevotella oralis		4.08	0.32		1	1	
Clostridium paraputrificum		2.28	0.21		99	99	
47750170	**Good identification**						**47750170**
					SPO		
Bifidobacterium breve		99.40	0.49	GAL 99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		0.60	-.07	BGL 91	99		
47750172	**Very good identification**						**47750172**
					GRA		
Bifidobacterium breve		99.95	0.66	GAL 99	99		
Prevotella oralis		0.05	-.03		1		
47750222	**Acceptable identification**						**47750222**
					GRA	SPO	
Prevotella oralis		90.96	0.39	GAL 99	1	1	
Clostridium barati		5.84	0.18	LAC 94	99	99	
Clostridium clostridioforme		2.40	0.19		50	66	
47750300	**Low discrimination**						**47750300**
					GRA	SPO	
Fusobacterium mortiferum		63.96	0.39	MLT 11	1	1	
Prevotella oralis		18.43	0.16	RHA 1	1	1	
Clostridium barati		9.08	0.10		99	99	
47750302	**Low discrimination**						**47750302**
					GRA	SPO	
Prevotella oralis		74.68	0.38	GAL 99	1	1	
Clostridium clostridioforme		7.99	0.28	NAG 95	50	66	
Clostridium barati		5.81	0.18		99	99	
47750312	**Low discrimination**						**47750312**
					GRA		
Clostridium clostridioforme		87.27	0.39	TRE 88	50		
Bifidobacterium breve		10.07	0.27	XYL 91	99		
Prevotella oralis		1.81	0.04	ARA 94	1		
47750320	**Low discrimination**						**47750320**
					SPO	GRA	
Prevotella oralis		77.87	0.38	GAL 99	1	1	
Clostridium ramosum		9.20	0.22	BGL 95	92	50	
Clostridium paraputrificum		6.09	0.13		99	99	
47750322	**Good identification**						**47750322**
					SPO	GRA	
Prevotella oralis		94.40	0.60	GAL 99	1	1	
Clostridium ramosum		2.35	0.33		92	50	
Clostridium barati		1.16	0.26		99	99	
47750332	**Low discrimination**						**47750332**
					GRA		
Clostridium clostridioforme		57.97	0.39	TRE 88	50		
Prevotella oralis		22.82	0.25	XYL 91	1		
Bifidobacterium breve		11.90	0.32	ARA 94	99		
47750412	**Acceptable identification**						**47750412**
Clostridium clostridioforme		94.83	0.36	TRE 88			
Bifidobacterium adolescentis		5.17	0.18	XYL 91			
47750432	**Good identification**						**47750432**
Clostridium clostridioforme		100.0	0.36	TRE 88			
47750500	**Low discrimination**						**47750500**
					GRA	SPO	
Fusobacterium mortiferum		46.45	0.39	MLT 11	1	1	
Clostridium butyricum		41.10	0.20	ARA 1	99	99	
Collinsella aerofaciens		4.77	-.03		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47750502	**Low discrimination**					**47750502**
					SPO	
Bifidobacterium adolescentis		76.17	0.50	GAL 99	1	
Clostridium clostridioforme		12.67	0.34	XYL 87	66	
Clostridium butyricum		3.48	0.13		99	
47750512	**Low discrimination**					**47750512**
Bifidobacterium adolescentis		60.38	0.52	GAL 99		
Clostridium clostridioforme		37.43	0.45	XYL 87		
Bifidobacterium breve		2.02	0.27			
47750532	**Low discrimination**					**47750532**
Clostridium clostridioforme		88.67	0.45	TRE 88		
Bifidobacterium breve		8.52	0.32	XYL 91		
Bifidobacterium adolescentis		1.45	0.18	RHA 88		
47750552	**Low discrimination**					**47750552**
Bifidobacterium adolescentis		75.66	0.36	GAL 99		
Bifidobacterium breve		20.51	0.27	XYL 87		
Clostridium clostridioforme		3.83	0.11	SOR 11		
47750602	**Good identification**					**47750602**
Clostridium clostridioforme		100.0	0.40	TRE 88		
47750612	**Good identification**					**47750612**
Clostridium clostridioforme		100.0	0.51	TRE 88		
47750622	**Good identification**					**47750622**
Clostridium clostridioforme		97.61	0.40	TRE 88		
Prevotella oralis		2.39	0.05	XYL 91		
47750632	**Good identification**					**47750632**
Clostridium clostridioforme		100.0	0.51	TRE 88		
47750702	**Good identification**					**47750702**
					GRA	
Clostridium clostridioforme		98.59	0.49	TRE 88	50	
Bifidobacterium adolescentis		0.82	0.16	XYL 91	99	
Prevotella oralis		0.59	0.04		1	
47750712	**Very good identification**					**47750712**
Clostridium clostridioforme		99.78	0.60	TRE 88		
Bifidobacterium adolescentis		0.22	0.18	XYL 91		
47750722	**Low discrimination**					**47750722**
Clostridium clostridioforme		85.60	0.49	TRE 88		
Prevotella oralis		9.80	0.25	XYL 91		
Parabacteroides distasonis		3.30	0.30			
47750732	**Very good identification**					**47750732**
Clostridium clostridioforme		99.55	0.60	TRE 88		
Parabacteroides distasonis		0.45	0.25	XYL 91		
47751000	**Low discrimination**					**47751000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Lactobacillus catenaformis		38.49	0.63	RAF 1	99		
Fusobacterium mortiferum		26.56	0.70		1		
Lactobacillus jensenii		19.32	0.58		99		
47751001	**Low discrimination**						**47751001**
					GRA		
Actinomyces naeslundii		80.59	0.68	ESL 12	99		
Lactobacillus catenaformis		7.79	0.29		99		
Fusobacterium mortiferum		5.37	0.36		1		
47751002	**Low discrimination**						**47751002**
					GRA		
Lactobacillus catenaformis		27.98	0.63	RAF 1	99		
Prevotella oralis		24.50	0.58		1		
Lactobacillus acidophilus		13.07	0.54		99		
47751003	**Low discrimination**						**47751003**
					GRA		
Actinomyces naeslundii		74.87	0.68	ESL 12	99		
Lactobacillus catenaformis		7.23	0.29		99		
Prevotella oralis		6.33	0.24		1		
47751010	**Low discrimination**						**47751010**
Lactobacillus jensenii		73.83	0.50	RAF 1			
Bifidobacterium breve		14.43	0.44				
Lactobacillus catenaformis		4.46	0.29				
47751011	**Low discrimination**						**47751011**
Actinomyces naeslundii		73.14	0.46	ESL 12			
Lactobacillus jensenii		22.47	0.15	MLZ 5			
Bifidobacterium breve		4.39	0.10				
47751012	**Low discrimination**						**47751012**
					GRA		
Bifidobacterium breve		63.04	0.62	LAC 99	99		
Lactobacillus jensenii		21.26	0.47		99		
Leptotrichia buccalis		5.31	0.24		1		
47751013	**Low discrimination**						**47751013**
Actinomyces naeslundii		53.68	0.46	ESL 12			
Bifidobacterium breve		32.59	0.27	MLZ 5			
Lactobacillus jensenii		10.99	0.12				
47751020	**Low discrimination**						**47751020**
					GRA	SPO	
Prevotella oralis		39.44	0.58	LAC 94	1	1	
Clostridium paraputrificum		20.72	0.47	BGL 95	99	99	
Clostridium barati		15.16	0.50		99	99	
47751021	**Low discrimination**						**47751021**
					GRA	SPO	
Actinomyces naeslundii		83.10	0.68	ESL 12	99	1	
Prevotella oralis		7.03	0.24		1	1	
Clostridium paraputrificum		3.69	0.13		99	99	
47751022	**Low discrimination**						**47751022**
					GRA	SPO	
Prevotella oralis		85.84	0.80	LAC 94	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		5.21	0.58		99	99	
Bifidobacterium breve		3.04	0.66		99	1	
47751023	**Low discrimination**						**47751023**
					GRA	SPO	
Prevotella oralis		56.15	0.45	LAC 94	1	1	
Actinomyces naeslundii		34.94	0.68	URE 1	99	1	
Clostridium barati		3.41	0.24		99	99	
47751030	**Low discrimination**						**47751030**
					GRA		
Bifidobacterium breve		47.25	0.49	LAC 99	99		
Lactobacillus jensenii		33.99	0.39	BGL 91	99		
Prevotella oralis		7.19	0.24		1		
47751031	**Low discrimination**						**47751031**
Actinomyces naeslundii		84.49	0.46	ESL 12			
Bifidobacterium breve		9.02	0.15	MLZ 5			
Lactobacillus jensenii		6.49	0.05				
47751032	**Low discrimination**						**47751032**
					GRA		
Bifidobacterium breve		72.07	0.66	LAC 99	99		
Prevotella oralis		20.59	0.45		1		
Lactobacillus jensenii		3.42	0.36		99		
47751040	**Low discrimination**						**47751040**
Bifidobacterium breve		37.47	0.44	LAC 99			
Actinomyces naeslundii		14.09	0.52	BGL 91			
Lactobacillus catenaformis		11.57	0.29				
47751041	**Good identification**						**47751041**
Actinomyces naeslundii		98.24	0.55	ESL 12			
Bifidobacterium breve		1.76	0.10				
47751042	**Low discrimination**						**47751042**
					SPO		
Bifidobacterium breve		70.96	0.62	LAC 99	1		
Lactobacillus acidophilus		12.39	0.39		1		
Clostridium barati		6.28	0.31		99		
47751043	**Low discrimination**						**47751043**
Actinomyces naeslundii		81.39	0.55	ESL 12			
Bifidobacterium breve		14.74	0.27				
Lactobacillus acidophilus		2.57	0.05				
47751050	**Acceptable identification**						**47751050**
Bifidobacterium breve		90.44	0.44	LAC 99			
Lactobacillus jensenii		4.67	0.15	BGL 91			
Actinomyces israelii		3.10	0.25				
47751052	**Very good identification**						**47751052**
Bifidobacterium breve		98.09	0.62	LAC 99			
Bifidobacterium adolescentis		0.55	0.22				
Actinomyces israelii		0.50	0.28				
47751060	**Low discrimination**						**47751060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Bifidobacterium breve		38.98	0.49	LAC 99	1		
Clostridium paraputrificum		23.21	0.27	BGL 91	99		
Clostridium barati		19.61	0.31		99		
47751061	**Acceptable identification**						**47751061**
					SPO		
Actinomyces naeslundii		93.74	0.55	ESL 12	1		
Bifidobacterium breve		2.98	0.15		1		
Clostridium paraputrificum		1.78	-.07		99		
47751062	**Low discrimination**						**47751062**
					GRA	SPO	
Bifidobacterium breve		62.94	0.66	LAC 99	99	1	
Prevotella oralis		17.98	0.45		1	1	
Clostridium barati		9.40	0.40		99	99	
47751063	**Low discrimination**						**47751063**
					GRA		
Actinomyces naeslundii		66.95	0.55	ESL 12	99		
Bifidobacterium breve		21.55	0.32		99		
Prevotella oralis		6.16	0.11		1		
47751070	**Good identification**						**47751070**
Bifidobacterium breve		97.14	0.49	LAC 99			
Actinomyces naeslundii		1.08	0.30	BGL 91			
Lactobacillus jensenii		0.71	0.05				
47751072	**Very good identification**						**47751072**
					GRA	SPO	
Bifidobacterium breve		99.31	0.66	LAC 99	99	1	
Prevotella oralis		0.29	0.11		1	1	
Clostridium barati		0.15	0.05		99	99	
47751100	**Low discrimination**						**47751100**
					GRA		
Collinsella aerofaciens		56.00	0.66	RAF 1	99		
Bifidobacterium breve		16.61	0.79		99		
Fusobacterium mortiferum		8.26	0.76		1		
47751101	**Low discrimination**						**47751101**
Actinomyces naeslundii		60.48	0.75	ESL 12			
Collinsella aerofaciens		22.41	0.32				
Bifidobacterium breve		6.65	0.44				
47751102	**Low discrimination**						**47751102**
					GRA		
Bifidobacterium breve		54.62	0.96		99		
Prevotella oralis		22.88	0.78		1		
Lactobacillus acidophilus		8.96	0.72		99		
47751103	**Low discrimination**						**47751103**
					GRA		
Bifidobacterium breve		36.70	0.62	URE 1	99		
Actinomyces naeslundii		33.02	0.75		99		
Prevotella oralis		15.37	0.44		1		
47751110	**Good identification**						**47751110**
					GRA		
Bifidobacterium breve		93.30	0.79	BGL 91	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Collinsella aerofaciens		3.18	0.32		99		
Leptotrichia buccalis		0.80	0.24		1		
47751111	**Low discrimination**						**47751111**
Bifidobacterium breve		65.73	0.44	URE 1			
Actinomyces naeslundii		31.47	0.53	BGL 91			
Collinsella aerofaciens		2.24	-.03				
47751112	**Low discrimination**						**47751112**
					GRA		
Bifidobacterium breve		87.82	0.96		99		
Leptotrichia buccalis		7.39	0.58		1		
Bifidobacterium adolescentis		3.96	0.72		99		
47751113	**Low discrimination**						**47751113**
					GRA		
Bifidobacterium breve		84.52	0.62	URE 1	99		
Leptotrichia buccalis		7.11	0.24		1		
Actinomyces naeslundii		4.00	0.53		99		
47751120	**Low discrimination**						**47751120**
					GRA	SPO	
Clostridium paraputrificum		65.76	0.81	RAF 7	99	99	
Prevotella oralis		19.80	0.78		1	1	
Bifidobacterium breve		8.31	0.83		99	1	
47751121	**Low discrimination**						**47751121**
					GRA	SPO	
Clostridium paraputrificum		46.25	0.47	RAF 7	99	99	
Actinomyces naeslundii		29.92	0.75	URE 1	99	1	
Prevotella oralis		13.93	0.44		1	1	
47751122	**Low discrimination**						**47751122**
					GRA	SPO	
Prevotella oralis		73.51	1.00		1	1	
Bifidobacterium breve		16.42	1.00		99	1	
Clostridium paraputrificum		5.51	0.75		99	99	
47751123	**Low discrimination**						**47751123**
					GRA		
Prevotella oralis		67.91	0.66	URE 1	1		
Bifidobacterium breve		15.17	0.66		99		
Actinomyces naeslundii		7.68	0.75		99		
47751130	**Low discrimination**						**47751130**
					GRA	SPO	
Bifidobacterium breve		89.71	0.83	BGL 91	99	1	
Clostridium paraputrificum		7.17	0.47		99	99	
Prevotella oralis		2.16	0.44		1	1	
47751131	**Low discrimination**						**47751131**
					SPO		
Bifidobacterium breve		72.82	0.49	URE 1	1		
Actinomyces naeslundii		19.61	0.53	BGL 91	1		
Clostridium paraputrificum		5.82	0.13		99		
47751132	**Good identification**						**47751132**
					GRA		
Bifidobacterium breve		94.35	1.00		99		
Prevotella oralis		4.27	0.66		1		
Leptotrichia buccalis		0.50	0.41		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47751133	**Good identification**						**47751133**
					GRA		
Bifidobacterium breve		92.13	0.66	URE 1	99		
Prevotella oralis		4.17	0.32		1		
Actinomyces naeslundii		2.45	0.53		99		
47751140	**Good identification**						**47751140**
Bifidobacterium breve		91.56	0.79	BGL 91			
Collinsella aerofaciens		3.12	0.32				
Actinomyces naeslundii		0.99	0.59				
47751141	**Low discrimination**						**47751141**
Actinomyces naeslundii		60.16	0.62	ESL 12			
Bifidobacterium breve		37.48	0.44				
Collinsella aerofaciens		1.28	-.03				
47751142	**Very good identification**						**47751142**
Bifidobacterium breve		96.48	0.96				
Lactobacillus acidophilus		1.96	0.57				
Bifidobacterium adolescentis		0.44	0.55				
47751143	**Low discrimination**						**47751143**
Bifidobacterium breve		83.85	0.62	URE 1			
Actinomyces naeslundii		13.31	0.62				
Lactobacillus acidophilus		1.70	0.23				
47751150	**Very good identification**						**47751150**
Bifidobacterium breve		99.67	0.79	BGL 91			
Actinomyces israelii		0.14	0.35				
Actinomyces naeslundii		0.06	0.38				
47751151	**Acceptable identification**						**47751151**
Bifidobacterium breve		92.21	0.44	URE 1			
Actinomyces naeslundii		7.79	0.41	BGL 91			
47751152	**Excellent identification**						**47751152**
					GRA		
Bifidobacterium breve		99.31	0.96		99		
Bifidobacterium adolescentis		0.55	0.56		99		
Leptotrichia buccalis		0.08	0.24		1		
47751153	**Very good identification**						**47751153**
Bifidobacterium breve		98.54	0.62	URE 1			
Actinomyces naeslundii		0.82	0.41				
Bifidobacterium adolescentis		0.55	0.22				
47751160	**Low discrimination**						**47751160**
					GRA	SPO	
Bifidobacterium breve		60.62	0.83	BGL 91	99	1	
Clostridium paraputrificum		36.09	0.62		99	99	
Prevotella oralis		1.46	0.44		1	1	
47751161	**Low discrimination**						**47751161**
					SPO		
Bifidobacterium breve		39.41	0.49	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		35.58	0.62	BGL 91	1		
Clostridium paraputrificum		23.46	0.27		99		
47751162	**Good identification**						**47751162**
					GRA	SPO	
Bifidobacterium breve		91.81	1.00		99	1	
Prevotella oralis		4.15	0.66		1	1	
Clostridium paraputrificum		2.32	0.55		99	99	
47751163	**Low discrimination**						**47751163**
					GRA		
Bifidobacterium breve		84.95	0.66	URE 1	99		
Actinomyces naeslundii		7.59	0.62		99		
Prevotella oralis		3.84	0.32		1		
47751170	**Very good identification**						**47751170**
					SPO		
Bifidobacterium breve		99.32	0.83	BGL 91	1		
Clostridium paraputrificum		0.60	0.27		99		
Actinomyces naeslundii		0.03	0.38		1		
47751171	**Acceptable identification**						**47751171**
					SPO		
Bifidobacterium breve		94.92	0.49	URE 1	1		
Actinomyces naeslundii		4.51	0.41	BGL 91	1		
Clostridium paraputrificum		0.57	-.07		99		
47751172	**Excellent identification**						**47751172**
					GRA	SPO	
Bifidobacterium breve		99.89	1.00		99	1	
Prevotella oralis		0.05	0.32		1	1	
Clostridium paraputrificum		0.03	0.21		99	99	
47751173	**Very good identification**						**47751173**
					GRA		
Bifidobacterium breve		99.49	0.66	URE 1	99		
Actinomyces naeslundii		0.47	0.41		99		
Prevotella oralis		0.05	-.03		1		
47751202	**Low discrimination**						**47751202**
					GRA	SPO	
Prevotella oralis		80.46	0.51	LAC 94	1	1	
Lactobacillus acidophilus		7.54	0.35	NAG 95	99	1	
Clostridium barati		6.27	0.31		99	99	
47751212	**Low discrimination**						**47751212**
					GRA		
Clostridium clostridioforme		73.16	0.39	TRE 88	50		
Prevotella oralis		10.69	0.17	XYL 91	1		
Bifidobacterium breve		9.42	0.27	ARA 94	99		
47751220	**Low discrimination**						**47751220**
					SPO	GRA	
Prevotella oralis		89.00	0.51	LAC 94	1	1	
Clostridium barati		6.94	0.31	BGL 95	99	99	
Clostridium ramosum		1.67	0.22		92	50	
47751222	**Very good identification**						**47751222**
					GRA	SPO	
Prevotella oralis		97.60	0.73	LAC 94	1	1	
Clostridium barati		1.20	0.40		99	99	
Lactobacillus acidophilus		0.48	0.35		99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
47751223	**Good identification**						**47751223**
					GRA	SPO	
Prevotella oralis		96.54	0.39	LAC 94	1	1	
Actinomyces naeslundii		1.42	0.33	URE 1	99	1	
Clostridium barati		1.19	0.05		99	99	
47751232	**Low discrimination**						**47751232**
					GRA		
Prevotella oralis		66.03	0.39	LAC 94	1		
Clostridium clostridioforme		23.78	0.39	MLZ 1	50		
Bifidobacterium breve		5.45	0.32		99		
47751262	**Low discrimination**						**47751262**
					GRA	SPO	
Prevotella oralis		79.82	0.39	LAC 94	1	1	
Clostridium barati		8.46	0.21	SOR 1	99	99	
Bifidobacterium breve		6.59	0.32		99	1	
47751300	**Low discrimination**						**47751300**
					GRA		
Prevotella oralis		56.25	0.50	NAG 95	1		
Collinsella aerofaciens		20.05	0.32	BGL 95	99		
Bifidobacterium breve		5.95	0.44		99		
47751302	**Low discrimination**						**47751302**
					GRA		
Prevotella oralis		85.99	0.72	NAG 95	1		
Lactobacillus acidophilus		5.92	0.53		99		
Bifidobacterium breve		4.84	0.62		99		
47751303	**Low discrimination**						**47751303**
					GRA		
Prevotella oralis		83.37	0.38	NAG 95	1		
Lactobacillus acidophilus		5.74	0.19	URE 1	99		
Bifidobacterium breve		4.69	0.27		99		
47751310	**Low discrimination**						**47751310**
					GRA		
Bifidobacterium breve		83.30	0.44	RHA 1	99		
Prevotella oralis		7.96	0.16	BGL 91	1		
Collinsella aerofaciens		2.84	-.03		99		
47751312	**Low discrimination**						**47751312**
					GRA		
Bifidobacterium breve		62.42	0.62	RHA 1	99		
Clostridium clostridioforme		16.39	0.48		50		
Prevotella oralis		11.21	0.38		1		
47751320	**Low discrimination**						**47751320**
					SPO	GRA	
Prevotella oralis		81.32	0.72	BGL 95	1	1	
Clostridium ramosum		9.61	0.56		92	50	
Clostridium paraputrificum		6.36	0.47		99	99	
47751321	**Low discrimination**						**47751321**
					SPO	GRA	
Prevotella oralis		78.62	0.38	URE 1	1	1	
Clostridium ramosum		9.29	0.22	BGL 95	92	50	
Clostridium paraputrificum		6.15	0.13		99	99	
47751322	**Very good identification**						**47751322**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		95.77	0.94		1		
Clostridium ramosum		2.38	0.67		92		
Parabacteroides distasonis		0.53	0.68		1		
47751323	**Very good identification**						**47751323**
					SPO		
Prevotella oralis		95.56	0.60	URE 1	1		
Clostridium ramosum		2.38	0.33		92		
Parabacteroides distasonis		0.53	0.34		1		
47751330	**Low discrimination**						**47751330**
					SPO	GRA	
Prevotella oralis		44.20	0.38	MLZ 1	1	1	
Bifidobacterium breve		43.30	0.49	BGL 95	1	99	
Clostridium ramosum		5.22	0.22		92	50	
47751332	**Low discrimination**						**47751332**
					GRA		
Prevotella oralis		51.81	0.60	MLZ 1	1		
Bifidobacterium breve		27.01	0.66		99		
Parabacteroides distasonis		15.31	0.63		1		
47751340	**Low discrimination**						**47751340**
					GRA	SPO	
Bifidobacterium breve		75.64	0.44	RHA 1	99	1	
Prevotella oralis		7.23	0.16	BGL 91	1	1	
Clostridium barati		5.88	0.13		99	99	
47751342	**Low discrimination**						**47751342**
					GRA		
Bifidobacterium breve		72.93	0.62	RHA 1	99		
Prevotella oralis		13.09	0.38		1		
Lactobacillus acidophilus		11.02	0.38		99		
47751350	**Good identification**						**47751350**
Bifidobacterium breve		98.98	0.44	RHA 1			
Actinomyces israelii		1.02	0.16	BGL 91			
47751352	**Very good identification**						**47751352**
Bifidobacterium breve		98.63	0.62	RHA 1			
Bifidobacterium adolescentis		0.55	0.22				
Clostridium clostridioforme		0.26	0.13				
47751360	**Low discrimination**						**47751360**
					GRA	SPO	
Prevotella oralis		35.41	0.38	SOR 1	1	1	
Bifidobacterium breve		34.69	0.49	BGL 95	99	1	
Clostridium paraputrificum		20.65	0.27		99	99	
47751362	**Low discrimination**						**47751362**
					GRA		
Prevotella oralis		61.33	0.60	SOR 1	1		
Bifidobacterium breve		31.97	0.66		99		
Lactobacillus acidophilus		2.72	0.38		99		
47751370	**Good identification**						**47751370**
					GRA	SPO	
Bifidobacterium breve		98.39	0.49	RHA 1	99	1	
Prevotella oralis		1.02	0.04	BGL 91	1	1	
Clostridium paraputrificum		0.59	-.07		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47751372	**Very good identification**						**47751372**
					GRA		
Bifidobacterium breve		97.28	0.66	RHA 1	99		
Prevotella oralis		1.89	0.25		1		
Parabacteroides distasonis		0.56	0.29		1		
47751402	**Low discrimination**						**47751402**
					SPO		
Bifidobacterium adolescentis		48.48	0.50	XYL 87	1		
Actinomyces israelii		14.06	0.47	LAC 99	1		
Clostridium butyricum		9.14	0.23		99		
47751412	**Low discrimination**						**47751412**
Bifidobacterium adolescentis		60.73	0.52	XYL 87			
Clostridium clostridioforme		33.73	0.44	LAC 99			
Actinomyces israelii		2.15	0.33				
47751422	**Low discrimination**						**47751422**
					GRA	SPO	
Prevotella oralis		78.14	0.45	LAC 94	1	1	
Clostridium clostridioforme		5.66	0.33	ARA 1	50	66	
Clostridium barati		4.74	0.24		99	99	
47751432	**Low discrimination**						**47751432**
					GRA		
Clostridium clostridioforme		85.22	0.44	TRE 88	50		
Bifidobacterium breve		9.14	0.32	XYL 91	99		
Prevotella oralis		2.61	0.11	RHA 88	1		
47751440	**Low discrimination**						**47751440**
					SPO		
Actinomyces israelii		61.94	0.37	XYL 92	1		
Clostridium butyricum		34.73	0.12		99		
Bifidobacterium breve		3.34	0.10		1		
47751452	**Low discrimination**						**47751452**
Bifidobacterium adolescentis		70.03	0.36	XYL 87			
Bifidobacterium breve		18.98	0.27	LAC 99			
Actinomyces israelii		7.82	0.26	SOR 11			
47751500	**Low discrimination**						**47751500**
					SPO		
Clostridium butyricum		73.58	0.54	TRE 94	99		
Collinsella aerofaciens		8.54	0.32	XYL 97	1		
Bifidobacterium adolescentis		6.32	0.50		1		
47751502	**Low discrimination**						**47751502**
					SPO		
Bifidobacterium adolescentis		87.95	0.84	XYL 87	1		
Clostridium butyricum		4.02	0.47		99		
Bifidobacterium breve		3.60	0.62		1		
47751503	**Low discrimination**						**47751503**
					SPO		
Bifidobacterium adolescentis		85.69	0.50	XYL 87	1		
Clostridium butyricum		3.92	0.13	URE 1	99		
Bifidobacterium breve		3.51	0.27		1		
47751510	**Low discrimination**						**47751510**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		50.93	0.52	XYL 87	1		
Clostridium butyricum		25.52	0.32	BGL 99	99		
Bifidobacterium breve		16.70	0.44		1		
47751512	**Good identification**						**47751512**
Bifidobacterium adolescentis		94.43	0.86	XYL 87			
Bifidobacterium breve		3.16	0.62				
Clostridium clostridioforme		1.77	0.53				
47751513	**Acceptable identification**						**47751513**
Bifidobacterium adolescentis		94.78	0.52	XYL 87			
Bifidobacterium breve		3.18	0.27	URE 1			
Clostridium clostridioforme		1.78	0.19				
47751520	**Low discrimination**						**47751520**
					GRA	SPO	
Clostridium paraputrificum		56.43	0.47	RAF 7	99	99	
Prevotella oralis		17.00	0.44	ARA 1	1	1	
Clostridium butyricum		10.13	0.36		99	99	
47751522	**Low discrimination**						**47751522**
					GRA	SPO	
Prevotella oralis		65.39	0.66	ARA 1	1	1	
Bifidobacterium breve		14.61	0.66		99	1	
Clostridium paraputrificum		4.90	0.40		99	99	
47751530	**Low discrimination**						**47751530**
					SPO		
Bifidobacterium breve		76.09	0.49	ARA 1	1		
Clostridium paraputrificum		6.08	0.13	BGL 91	99		
Clostridium butyricum		5.69	0.14		99		
47751532	**Low discrimination**						**47751532**
Bifidobacterium breve		59.39	0.66	ARA 1			
Clostridium clostridioforme		18.74	0.53				
Bifidobacterium adolescentis		10.08	0.52				
47751540	**Low discrimination**						**47751540**
					SPO		
Clostridium butyricum		54.32	0.36	TRE 94	99		
Bifidobacterium breve		21.51	0.44	XYL 97	1		
Actinomyces israelii		16.15	0.48	SOR 8	1		
47751542	**Low discrimination**						**47751542**
Bifidobacterium adolescentis		70.86	0.69	XYL 87			
Bifidobacterium breve		23.47	0.62	SOR 11			
Actinomyces israelii		2.61	0.51				
47751550	**Low discrimination**						**47751550**
					SPO		
Bifidobacterium breve		61.17	0.44	ARA 1	1		
Bifidobacterium adolescentis		23.05	0.36	BGL 91	1		
Clostridium butyricum		8.13	0.14		99		
47751552	**Low discrimination**						**47751552**
Bifidobacterium adolescentis		78.21	0.70	XYL 87			
Bifidobacterium breve		21.19	0.62	SOR 11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.35	0.37				
47751553	**Low discrimination**						**47751553**
Bifidobacterium adolescentis		78.68	0.36	XYL 87			
Bifidobacterium breve		21.32	0.27	SOR 11			
47751560	**Low discrimination**						**47751560**
					SPO		
Bifidobacterium breve		55.34	0.49	ARA 1	1		
Clostridium paraputrificum		32.95	0.27	BGL 91	99		
Clostridium butyricum		6.83	0.17		99		
47751562	**Low discrimination**						**47751562**
					GRA	SPO	
Bifidobacterium breve		88.33	0.66	ARA 1	99	1	
Prevotella oralis		3.99	0.32		1	1	
Clostridium paraputrificum		2.23	0.21		99	99	
47751570	**Good identification**						**47751570**
					SPO		
Bifidobacterium breve		98.76	0.49	ARA 1	1		
Clostridium butyricum		0.64	-.05	BGL 91	99		
Clostridium paraputrificum		0.59	-.07		99		
47751572	**Very good identification**						**47751572**
Bifidobacterium breve		97.28	0.66	ARA 1			
Bifidobacterium adolescentis		2.04	0.36				
Clostridium clostridioforme		0.31	0.19				
47751602	**Acceptable identification**						**47751602**
					GRA		
Clostridium clostridioforme		92.22	0.48	TRE 88	50		
Prevotella oralis		3.92	0.17	XYL 91	1		
Actinomyces israelii		1.84	0.28		99		
47751612	**Very good identification**						**47751612**
Clostridium clostridioforme		99.69	0.59	TRE 88			
Bifidobacterium adolescentis		0.25	0.18	XYL 91			
Actinomyces israelii		0.07	0.14				
47751622	**Low discrimination**						**47751622**
Clostridium clostridioforme		52.77	0.48	TRE 88			
Prevotella oralis		42.60	0.39	XYL 91			
Parabacteroides distasonis		2.27	0.30				
47751632	**Very good identification**						**47751632**
Clostridium clostridioforme		99.32	0.59	TRE 88			
Parabacteroides distasonis		0.50	0.25	XYL 91			
Prevotella oralis		0.18	0.05				
47751702	**Low discrimination**						**47751702**
					GRA		
Clostridium clostridioforme		63.11	0.57	TRE 88	50		
Bifidobacterium adolescentis		17.25	0.50	XYL 91	99		
Prevotella oralis		12.55	0.38		1		
47751712	**Good identification**						**47751712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		92.66	0.68	TRE 88		
Bifidobacterium adolescentis		6.79	0.52	XYL 91		
Bifidobacterium breve		0.23	0.27			
47751720	**Low discrimination**					**47751720**
					SPO	
Bacteroides vulgatus		63.94	0.39	CEL 5	1	
Prevotella oralis		25.32	0.38	XYL 98	1	
Clostridium ramosum		2.99	0.22		92	
47751722	**Low discrimination**					**47751722**
Prevotella oralis		56.68	0.60	ARA 1		
Parabacteroides distasonis		19.06	0.64			
Clostridium clostridioforme		15.01	0.57			
47751732	**Low discrimination**					**47751732**
Clostridium clostridioforme		85.36	0.68	TRE 88		
Parabacteroides distasonis		12.82	0.60	XYL 91		
Prevotella oralis		0.72	0.25			
47752000	**Low discrimination**					**47752000**
					GRA	
Actinomyces naeslundii		44.98	0.46	ESL 12	99	
Fusobacterium mortiferum		37.07	0.33	GAL 96	1	
Actinomyces israelii		8.81	0.25		99	
47752001	**Good identification**					**47752001**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47752002	**Low discrimination**					**47752002**
Actinomyces naeslundii		42.58	0.46	ESL 12		
Bifidobacterium breve		25.23	0.27	GAL 96		
Actinomyces israelii		12.51	0.28			
47752003	**Good identification**					**47752003**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47752020	**Low discrimination**					**47752020**
					SPO	
Actinomyces naeslundii		69.88	0.46	ESL 12	1	
Clostridium barati		19.08	0.10	GAL 96	99	
Bifidobacterium breve		7.28	0.15		1	
47752021	**Good identification**					**47752021**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47752023	**Good identification**					**47752023**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47752041	**Good identification**					**47752041**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47752043	**Good identification**					**47752043**
Actinomyces naeslundii		100.0	0.36	ESL 12		
47752061	**Good identification**					**47752061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.36	ESL 12			
47752063	**Good identification**						**47752063**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47752100	**Low discrimination**						**47752100**
					GRA		
Bifidobacterium breve		48.84	0.44	GAL 99	99		
Actinomyces naeslundii		23.96	0.54	BGL 91	99		
Fusobacterium mortiferum		16.19	0.39		1		
47752101	**Very good identification**						**47752101**
Actinomyces naeslundii		98.65	0.57	ESL 12			
Bifidobacterium breve		1.36	0.10	GAL 96			
47752102	**Low discrimination**						**47752102**
Bifidobacterium breve		88.15	0.62	GAL 99			
Actinomyces naeslundii		4.28	0.54				
Bifidobacterium adolescentis		2.66	0.35				
47752103	**Low discrimination**						**47752103**
Actinomyces naeslundii		87.81	0.57	ESL 12			
Bifidobacterium breve		12.19	0.27	GAL 96			
47752110	**Good identification**						**47752110**
Bifidobacterium breve		95.60	0.44	GAL 99			
Actinomyces naeslundii		2.47	0.32	BGL 91			
Actinomyces israelii		1.93	0.21				
47752112	**Very good identification**						**47752112**
Bifidobacterium breve		95.70	0.62	GAL 99			
Bifidobacterium adolescentis		3.53	0.36				
Actinomyces israelii		0.29	0.24				
47752120	**Low discrimination**						**47752120**
					SPO		
Bifidobacterium breve		50.38	0.49	GAL 99	1		
Clostridium ramosum		25.98	0.33	BGL 91	92		
Actinomyces naeslundii		13.90	0.54		1		
47752121	**Very good identification**						**47752121**
					SPO		
Actinomyces naeslundii		96.43	0.57	ESL 12	1		
Bifidobacterium breve		2.35	0.15	GAL 96	1		
Clostridium ramosum		1.21	-.01		92		
47752122	**Low discrimination**						**47752122**
					SPO	GRA	
Bifidobacterium breve		76.26	0.66	GAL 99	1	99	
Clostridium ramosum		15.56	0.43		92	50	
Prevotella oralis		3.45	0.32		1	1	
47752123	**Low discrimination**						**47752123**
					SPO		
Actinomyces naeslundii		76.44	0.57	ESL 12	1		
Bifidobacterium breve		18.86	0.32	GAL 96	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		3.85	0.09		92	
47752130	**Good identification**					**47752130**
					SPO	
Bifidobacterium breve		97.19	0.49	GAL 99	1	
Actinomyces naeslundii		1.41	0.32	BGL 91	1	
Clostridium tertium		0.90	0.05		99	
47752132	**Very good identification**					**47752132**
					SPO	
Bifidobacterium breve		99.34	0.66	GAL 99	1	
Clostridium tertium		0.27	0.13		99	
Clostridium ramosum		0.21	0.09		92	
47752140	**Low discrimination**					**47752140**
Bifidobacterium breve		87.79	0.44	GAL 99		
Actinomyces naeslundii		7.60	0.41	BGL 91		
Actinomyces israelii		4.61	0.28			
47752141	**Acceptable identification**					**47752141**
Actinomyces naeslundii		92.78	0.44	ESL 12		
Bifidobacterium breve		7.22	0.10	GAL 96		
47752142	**Very good identification**					**47752142**
Bifidobacterium breve		97.81	0.62	GAL 99		
Actinomyces naeslundii		0.84	0.41			
Actinomyces israelii		0.76	0.31			
47752143	**Low discrimination**					**47752143**
Actinomyces naeslundii		55.97	0.44	ESL 12		
Bifidobacterium breve		44.03	0.27	GAL 96		
47752150	**Good identification**					**47752150**
Bifidobacterium breve		99.22	0.44	GAL 99		
Actinomyces israelii		0.78	0.14	BGL 91		
47752152	**Very good identification**					**47752152**
Bifidobacterium breve		99.43	0.62	GAL 99		
Bifidobacterium adolescentis		0.45	0.21			
Actinomyces israelii		0.12	0.17			
47752160	**Acceptable identification**					**47752160**
					SPO	
Bifidobacterium breve		93.74	0.49	GAL 99	1	
Actinomyces naeslundii		4.56	0.41	BGL 91	1	
Clostridium barati		0.65	-.01		99	
47752161	**Low discrimination**					**47752161**
Actinomyces naeslundii		87.85	0.44	ESL 12		
Bifidobacterium breve		12.15	0.15	GAL 96		
47752162	**Very good identification**					**47752162**
					SPO	
Bifidobacterium breve		99.01	0.66	GAL 99	1	
Actinomyces naeslundii		0.48	0.41		1	
Clostridium ramosum		0.20	0.09		92	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47752170	**Good identification**						**47752170**
Bifidobacterium breve		100.0	0.49	GAL 99			
47752172	**Very good identification**						**47752172**
Bifidobacterium breve		100.0	0.66	GAL 99			
47752320	**Good identification**						**47752320**
					SPO	GRA	
Clostridium ramosum		96.16	0.35	TRE 94	92	50	
Prevotella oralis		1.45	0.04	GAL 99	1	1	
Bifidobacterium breve		1.42	0.15		1	99	
47752322	**Low discrimination**						**47752322**
					SPO	GRA	
Clostridium ramosum		89.56	0.45	TRE 94	92	50	
Prevotella oralis		6.42	0.25	GAL 99	1	1	
Bifidobacterium breve		3.35	0.32		1	99	
47752502	**Low discrimination**						**47752502**
Bifidobacterium adolescentis		87.92	0.49	GAL 99			
Actinomyces israelii		7.14	0.36	XYL 87			
Bifidobacterium breve		4.40	0.27				
47752512	**Acceptable identification**						**47752512**
Bifidobacterium adolescentis		94.00	0.50	GAL 99			
Bifidobacterium breve		3.85	0.27	XYL 87			
Actinomyces israelii		0.93	0.22				
47753000	**Low discrimination**						**47753000**
Actinomyces naeslundii		42.61	0.70	ESL 12			
Actinomyces israelii		34.42	0.59				
Bifidobacterium breve		10.30	0.44				
47753001	**Very good identification**						**47753001**
Actinomyces naeslundii		99.30	0.73	ESL 12			
Actinomyces israelii		0.54	0.25				
Bifidobacterium breve		0.16	0.10				
47753002	**Low discrimination**						**47753002**
Bifidobacterium breve		45.66	0.62	LAC 99			
Actinomyces israelii		22.64	0.62				
Actinomyces naeslundii		18.68	0.70				
47753003	**Very good identification**						**47753003**
Actinomyces naeslundii		97.34	0.73	ESL 12			
Bifidobacterium breve		1.60	0.27				
Actinomyces israelii		0.79	0.28				
47753010	**Low discrimination**						**47753010**
Bifidobacterium breve		56.31	0.44	LAC 99			
Actinomyces israelii		28.12	0.45	BGL 91			
Actinomyces naeslundii		12.26	0.48				
47753011	**Good identification**						**47753011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		97.00	0.51	ESL 12		
Bifidobacterium breve		3.00	0.10	MLZ 5		
47753012	**Low discrimination**					**47753012**
Bifidobacterium breve		87.66	0.62	LAC 99		
Actinomyces israelii		6.49	0.48			
Bifidobacterium adolescentis		3.23	0.36			
47753013	**Low discrimination**					**47753013**
Actinomyces naeslundii		74.85	0.51	ESL 12		
Bifidobacterium breve		23.41	0.27	MLZ 5		
Actinomyces israelii		1.73	0.13			
47753020	**Low discrimination**					**47753020**
					SPO	
Actinomyces naeslundii		48.76	0.70	ESL 12	1	
Bifidobacterium breve		20.96	0.49		1	
Clostridium ramosum		10.81	0.33		92	
47753021	**Very good identification**					**47753021**
					SPO	
Actinomyces naeslundii		99.42	0.73	ESL 12	1	
Bifidobacterium breve		0.29	0.15		1	
Clostridium ramosum		0.15	-.01		92	
47753022	**Low discrimination**					**47753022**
					GRA	
Bifidobacterium breve		51.54	0.66	LAC 99	99	
Prevotella oralis		14.73	0.45		1	
Actinomyces naeslundii		11.86	0.70		99	
47753023	**Very good identification**					**47753023**
					GRA	
Actinomyces naeslundii		95.31	0.73	ESL 12	99	
Bifidobacterium breve		2.79	0.32		99	
Prevotella oralis		0.80	0.11		1	
47753030	**Low discrimination**					**47753030**
Bifidobacterium breve		85.68	0.49	LAC 99		
Actinomyces naeslundii		10.49	0.48	BGL 91		
Actinomyces israelii		1.54	0.24			
47753031	**Acceptable identification**					**47753031**
Actinomyces naeslundii		94.79	0.51	ESL 12		
Bifidobacterium breve		5.21	0.15	MLZ 5		
47753032	**Very good identification**					**47753032**
					GRA	
Bifidobacterium breve		97.57	0.66	LAC 99	99	
Actinomyces naeslundii		1.18	0.48		99	
Prevotella oralis		0.28	0.11		1	
47753033	**Low discrimination**					**47753033**
Actinomyces naeslundii		64.27	0.51	ESL 12		
Bifidobacterium breve		35.73	0.32	MLZ 5		
47753040	**Low discrimination**					**47753040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		42.53	0.52	XYL 92		
Bifidobacterium breve		32.73	0.44			
Actinomyces naeslundii		23.89	0.57			
47753041	**Very good identification**					**47753041**
Actinomyces naeslundii		97.92	0.60	ESL 12		
Actinomyces israelii		1.17	0.18			
Bifidobacterium breve		0.90	0.10			
47753042	**Low discrimination**					**47753042**
Bifidobacterium breve		77.79	0.62	LAC 99		
Actinomyces israelii		15.00	0.55			
Actinomyces naeslundii		5.62	0.57			
47753043	**Low discrimination**					**47753043**
Actinomyces naeslundii		89.99	0.60	ESL 12		
Bifidobacterium breve		8.39	0.27			
Actinomyces israelii		1.62	0.21			
47753050	**Low discrimination**					**47753050**
Bifidobacterium breve		81.13	0.44	LAC 99		
Actinomyces israelii		15.75	0.38	BGL 91		
Actinomyces naeslundii		3.12	0.35			
47753051	**Low discrimination**					**47753051**
Actinomyces naeslundii		85.08	0.38	ESL 12		
Bifidobacterium breve		14.92	0.10	MLZ 5		
47753052	**Very good identification**					**47753052**
Bifidobacterium breve		96.42	0.62	LAC 99		
Actinomyces israelii		2.78	0.41			
Bifidobacterium adolescentis		0.44	0.21			
47753060	**Low discrimination**					**47753060**
Bifidobacterium breve		66.13	0.49	LAC 99		
Actinomyces naeslundii		27.15	0.57	BGL 91		
Actinomyces israelii		3.09	0.31			
47753061	**Very good identification**					**47753061**
Actinomyces naeslundii		98.39	0.60	ESL 12		
Bifidobacterium breve		1.61	0.15			
47753062	**Good identification**					**47753062**
					SPO	
Bifidobacterium breve		93.40	0.66	LAC 99	1	
Actinomyces naeslundii		3.79	0.57		1	
Clostridium barati		1.21	0.21		99	
47753063	**Low discrimination**					**47753063**
Actinomyces naeslundii		85.78	0.60	ESL 12		
Bifidobacterium breve		14.23	0.32			
47753070	**Good identification**					**47753070**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		97.22	0.49	LAC 99		
Actinomyces naeslundii		2.10	0.35	BGL 91		
Actinomyces israelii		0.68	0.17			
47753071	**Low discrimination**					**47753071**
Actinomyces naeslundii		76.24	0.38	ESL 12		
Bifidobacterium breve		23.76	0.15	MLZ 5		
47753072	**Very good identification**					**47753072**
Bifidobacterium breve		99.68	0.66	LAC 99		
Actinomyces naeslundii		0.21	0.35			
Actinomyces israelii		0.10	0.20			
47753100	**Low discrimination**					**47753100**
Bifidobacterium breve		75.13	0.79	BGL 91		
Actinomyces israelii		10.14	0.69			
Actinomyces naeslundii		8.93	0.77			
47753101	**Good identification**					**47753101**
Actinomyces naeslundii		93.70	0.80	ESL 12		
Bifidobacterium breve		5.31	0.44			
Actinomyces israelii		0.72	0.35			
47753102	**Good identification**					**47753102**
Bifidobacterium breve		92.26	0.96			
Bifidobacterium adolescentis		2.78	0.69			
Actinomyces israelii		1.85	0.72			
47753103	**Low discrimination**					**47753103**
Actinomyces naeslundii		61.99	0.80	ESL 12		
Bifidobacterium breve		35.49	0.62			
Bifidobacterium adolescentis		1.07	0.35			
47753110	**Very good identification**					**47753110**
Bifidobacterium breve		96.48	0.79	BGL 91		
Actinomyces israelii		1.95	0.55			
Actinomyces naeslundii		0.60	0.55			
47753111	**Low discrimination**					**47753111**
Bifidobacterium breve		51.29	0.44	URE 1		
Actinomyces naeslundii		47.67	0.58	BGL 91		
Actinomyces israelii		1.04	0.21			
47753112	**Very good identification**					**47753112**
Bifidobacterium breve		95.86	0.96			
Bifidobacterium adolescentis		3.53	0.70			
Actinomyces israelii		0.29	0.58			
47753113	**Low discrimination**					**47753113**
Bifidobacterium breve		88.16	0.62	URE 1		
Actinomyces naeslundii		8.10	0.58			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		3.25	0.36				
47753120	**Low discrimination**						**47753120**
					SPO		
Bifidobacterium breve		58.25	0.83	BGL 91	1		
Clostridium ramosum		30.04	0.67		92		
Clostridium paraputrificum		4.65	0.47		99		
47753121	**Low discrimination**						**47753121**
					SPO		
Actinomyces naeslundii		85.79	0.80	ESL 12	1		
Bifidobacterium breve		8.64	0.49		1		
Clostridium ramosum		4.46	0.33		92		
47753122	**Low discrimination**						**47753122**
					SPO	GRA	
Bifidobacterium breve		78.62	1.00		1	99	
Clostridium ramosum		16.04	0.77		92	50	
Prevotella oralis		3.56	0.66		1	1	
47753123	**Low discrimination**						**47753123**
					SPO		
Bifidobacterium breve		44.50	0.66	URE 1	1		
Actinomyces naeslundii		43.71	0.80		1		
Clostridium ramosum		9.08	0.43		92		
47753130	**Very good identification**						**47753130**
					SPO		
Bifidobacterium breve		98.05	0.83	BGL 91	1		
Clostridium tertium		0.90	0.39		99		
Clostridium ramosum		0.51	0.33		92		
47753131	**Low discrimination**						**47753131**
					SPO		
Bifidobacterium breve		65.05	0.49	URE 1	1		
Actinomyces naeslundii		34.01	0.58	BGL 91	1		
Clostridium tertium		0.60	0.05		99		
47753132	**Excellent identification**						**47753132**
					SPO		
Bifidobacterium breve		99.39	1.00		1		
Clostridium tertium		0.27	0.47		99		
Clostridium ramosum		0.21	0.43		92		
47753133	**Good identification**						**47753133**
					SPO		
Bifidobacterium breve		94.61	0.66	URE 1	1		
Actinomyces naeslundii		4.89	0.58		1		
Clostridium tertium		0.26	0.13		99		
47753140	**Good identification**						**47753140**
Bifidobacterium breve		92.97	0.79	BGL 91			
Actinomyces israelii		4.88	0.62				
Actinomyces naeslundii		1.95	0.64				
47753141	**Low discrimination**						**47753141**
Actinomyces naeslundii		74.75	0.67	ESL 12			
Bifidobacterium breve		23.99	0.44				
Actinomyces israelii		1.26	0.28				
47753142	**Very good identification**						**47753142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		98.52	0.96			
Actinomyces israelii		0.77	0.65			
Bifidobacterium adolescentis		0.37	0.53			
47753143	**Low discrimination**					**47753143**
Bifidobacterium breve		75.71	0.62	URE 1		
Actinomyces naeslundii		23.33	0.67			
Actinomyces israelii		0.59	0.31			
47753150	**Very good identification**					**47753150**
Bifidobacterium breve		99.05	0.79	BGL 91		
Actinomyces israelii		0.78	0.48			
Actinomyces naeslundii		0.11	0.43			
47753151	**Low discrimination**					**47753151**
Bifidobacterium breve		85.33	0.44	URE 1		
Actinomyces naeslundii		14.00	0.46	BGL 91		
Actinomyces israelii		0.67	0.14			
47753152	**Excellent identification**					**47753152**
Bifidobacterium breve		99.42	0.96			
Bifidobacterium adolescentis		0.45	0.55			
Actinomyces israelii		0.12	0.51			
47753153	**Very good identification**					**47753153**
Bifidobacterium breve		97.85	0.62	URE 1		
Actinomyces naeslundii		1.59	0.46			
Bifidobacterium adolescentis		0.45	0.21			
47753160	**Very good identification**					**47753160**
					SPO	
Bifidobacterium breve		97.41	0.83	BGL 91	1	
Actinomyces naeslundii		1.15	0.64		1	
Clostridium paraputrificum		0.59	0.27		99	
47753161	**Low discrimination**					**47753161**
					SPO	
Actinomyces naeslundii		63.42	0.67	ESL 12	1	
Bifidobacterium breve		36.18	0.49		1	
Clostridium paraputrificum		0.22	-.07		99	
47753162	**Excellent identification**					**47753162**
					SPO	
Bifidobacterium breve		99.48	1.00		1	
Clostridium ramosum		0.21	0.43		92	
Actinomyces naeslundii		0.12	0.64		1	
47753163	**Low discrimination**					**47753163**
					SPO	
Bifidobacterium breve		85.00	0.66	URE 1	1	
Actinomyces naeslundii		14.74	0.67		1	
Clostridium ramosum		0.18	0.09		92	
47753170	**Very good identification**					**47753170**
Bifidobacterium breve		99.89	0.83	BGL 91		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.06	0.43				
Actinomyces israelii		0.03	0.27				
47753171	**Acceptable identification**						**47753171**
Bifidobacterium breve		91.55	0.49	URE 1			
Actinomyces naeslundii		8.45	0.46	BGL 91			
47753172	**Excellent identification**						**47753172**
Bifidobacterium breve		99.98	1.00				
Actinomyces naeslundii		0.01	0.43				
Actinomyces israelii		0.00	0.30				
47753173	**Very good identification**						**47753173**
Bifidobacterium breve		99.10	0.66	URE 1			
Actinomyces naeslundii		0.90	0.46				
47753200	**Low discrimination**						**47753200**
					SPO		
Actinomyces israelii		73.63	0.39	XYL 92	1		
Actinomyces naeslundii		12.23	0.35	RHA 7	1		
Clostridium barati		7.59	0.05		99		
47753201	**Good identification**						**47753201**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47753202	**Low discrimination**						**47753202**
					GRA		
Actinomyces israelii		45.98	0.43	XYL 92	99		
Prevotella oralis		14.12	0.17	RHA 7	1		
Bifidobacterium breve		12.45	0.27		99		
47753203	**Good identification**						**47753203**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47753220	**Low discrimination**						**47753220**
					SPO	GRA	
Clostridium ramosum		81.86	0.35	TRE 94	92	50	
Prevotella oralis		7.80	0.17	LAC 99	1	1	
Clostridium barati		5.24	0.13		99	99	
47753221	**Low discrimination**						**47753221**
					SPO		
Actinomyces naeslundii		83.62	0.38	ESL 12	1		
Clostridium ramosum		16.38	0.01	RHA 1	92		
47753222	**Low discrimination**						**47753222**
					SPO	GRA	
Clostridium ramosum		63.97	0.45	TRE 94	92	50	
Prevotella oralis		28.96	0.39	LAC 99	1	1	
Clostridium barati		3.07	0.21		99	99	
47753223	**Low discrimination**						**47753223**
					SPO	GRA	
Actinomyces naeslundii		46.76	0.38	ESL 12	1	99	
Clostridium ramosum		36.65	0.11	RHA 1	92	50	
Prevotella oralis		16.59	0.05		1	1	
47753242	**Low discrimination**						**47753242**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		57.26	0.35	XYL 92			
Bifidobacterium breve		39.86	0.27	RHA 7			
Actinomyces naeslundii		2.88	0.22				
47753300	**Low discrimination**						**47753300**
					SPO		
Clostridium ramosum		34.15	0.35	TRE 94	92		
Actinomyces israelii		28.27	0.50	NAG 99	1		
Bifidobacterium breve		28.10	0.44		1		
47753301	**Low discrimination**						**47753301**
					SPO		
Actinomyces naeslundii		84.57	0.46	ESL 12	1		
Clostridium ramosum		5.82	0.01	RHA 1	92		
Actinomyces israelii		4.82	0.16		1		
47753302	**Low discrimination**						**47753302**
					SPO	GRA	
Bifidobacterium breve		51.53	0.62	RHA 1	1	99	
Clostridium ramosum		24.78	0.45		92	50	
Prevotella oralis		9.25	0.38		1	1	
47753303	**Low discrimination**						**47753303**
					SPO		
Actinomyces naeslundii		49.11	0.46	ESL 12	1		
Bifidobacterium breve		28.12	0.27	RHA 1	1		
Clostridium ramosum		13.52	0.11		92		
47753310	**Low discrimination**						**47753310**
					SPO		
Bifidobacterium breve		86.01	0.44	RHA 1	1		
Actinomyces israelii		12.93	0.36	BGL 91	1		
Clostridium ramosum		1.06	0.01		92		
47753312	**Good identification**						**47753312**
Bifidobacterium breve		93.37	0.62	RHA 1			
Bifidobacterium adolescentis		3.44	0.36				
Actinomyces israelii		2.08	0.39				
47753320	**Very good identification**						**47753320**
					SPO	GRA	
Clostridium ramosum		96.66	0.69	TRE 94	92	50	
Prevotella oralis		1.46	0.38		1	1	
Bifidobacterium breve		1.43	0.49		1	99	
47753321	**Low discrimination**						**47753321**
					SPO	GRA	
Clostridium ramosum		84.99	0.35	TRE 94	92	50	
Actinomyces naeslundii		12.47	0.46	URE 1	1	99	
Prevotella oralis		1.28	0.04		1	1	
47753322	**Low discrimination**						**47753322**
					SPO	GRA	
Clostridium ramosum		89.86	0.80	TRE 94	92	50	
Prevotella oralis		6.44	0.60		1	1	
Bifidobacterium breve		3.36	0.66		1	99	
47753323	**Low discrimination**						**47753323**
					SPO	GRA	
Clostridium ramosum		87.29	0.45	TRE 94	92	50	
Prevotella oralis		6.25	0.25	URE 1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		3.26	0.32		1	99	
47753330	**Low discrimination**						**47753330**
					SPO	GRA	
Bifidobacterium breve		58.53	0.49	RHA 1	1	99	
Clostridium ramosum		40.02	0.35	BGL 91	92	50	
Prevotella oralis		0.60	0.04		1	1	
47753332	**Low discrimination**						**47753332**
					SPO	GRA	
Bifidobacterium breve		76.89	0.66	RHA 1	1	99	
Clostridium ramosum		20.80	0.45		92	50	
Prevotella oralis		1.49	0.25		1	1	
47753340	**Low discrimination**						**47753340**
Bifidobacterium breve		70.20	0.44	RHA 1			
Actinomyces israelii		27.47	0.43	BGL 91			
Actinomyces naeslundii		1.47	0.30				
47753342	**Good identification**						**47753342**
Bifidobacterium breve		92.55	0.62	RHA 1			
Actinomyces israelii		5.37	0.46				
Lactobacillus acidophilus		0.74	0.16				
47753350	**Acceptable identification**						**47753350**
					COC		
Bifidobacterium breve		93.33	0.44	RHA 1	1		
Actinomyces israelii		5.46	0.29	BGL 91	1		
Blautia producta		1.22	-.03		99		
47753352	**Very good identification**						**47753352**
Bifidobacterium breve		98.69	0.62	RHA 1			
Actinomyces israelii		0.86	0.32				
Bifidobacterium adolescentis		0.45	0.21				
47753360	**Low discrimination**						**47753360**
					SPO		
Bifidobacterium breve		57.56	0.49	RHA 1	1		
Clostridium ramosum		39.36	0.35	BGL 91	92		
Actinomyces israelii		0.81	0.22		1		
47753362	**Low discrimination**						**47753362**
					SPO	GRA	
Bifidobacterium breve		76.87	0.66	RHA 1	1	99	
Clostridium ramosum		20.79	0.45		92	50	
Prevotella oralis		1.49	0.25		1	1	
47753370	**Good identification**						**47753370**
					SPO		
Bifidobacterium breve		99.31	0.49	RHA 1	1		
Clostridium ramosum		0.69	0.01	BGL 91	92		
47753372	**Very good identification**						**47753372**
					SPO		
Bifidobacterium breve		99.73	0.66	RHA 1	1		
Clostridium ramosum		0.27	0.11		92		
47753400	**Good identification**						**47753400**
					SPO		
Actinomyces israelii		92.00	0.57	XYL 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		4.52	0.14		99	
Actinomyces naeslundii		1.41	0.35		1	
47753401	**Low discrimination**					**47753401**
Actinomyces naeslundii		69.41	0.38	ESL 12		
Actinomyces israelii		30.59	0.23	ARA 1		
47753402	**Low discrimination**					**47753402**
Actinomyces israelii		64.22	0.60	XYL 92		
Bifidobacterium adolescentis		31.96	0.49			
Bifidobacterium breve		1.60	0.27			
47753403	**Low discrimination**					**47753403**
Actinomyces naeslundii		50.25	0.38	ESL 12		
Actinomyces israelii		33.22	0.26	ARA 1		
Bifidobacterium adolescentis		16.53	0.15			
47753410	**Acceptable identification**					**47753410**
Actinomyces israelii		92.05	0.43	XYL 92		
Bifidobacterium adolescentis		5.68	0.16	MLZ 13		
Bifidobacterium breve		2.28	0.10			
47753412	**Low discrimination**					**47753412**
Bifidobacterium adolescentis		77.00	0.50	XYL 87		
Actinomyces israelii		18.91	0.46	LAC 99		
Bifidobacterium breve		3.15	0.27			
47753420	**Low discrimination**					**47753420**
Actinomyces israelii		66.05	0.37	XYL 92		
Actinomyces naeslundii		15.81	0.35	NAG 6		
Bifidobacterium breve		6.80	0.15			
47753421	**Good identification**					**47753421**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47753422	**Low discrimination**					**47753422**
					GRA	
Actinomyces israelii		41.59	0.40	XYL 92	99	
Bifidobacterium breve		28.85	0.32	NAG 6	99	
Prevotella oralis		8.24	0.11		1	
47753423	**Good identification**					**47753423**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47753440	**Good identification**					**47753440**
					SPO	
Actinomyces israelii		97.33	0.50	XYL 92	1	
Clostridium butyricum		1.07	-.04		99	
Bifidobacterium breve		0.93	0.10		1	
47753442	**Low discrimination**					**47753442**
Actinomyces israelii		81.51	0.53	XYL 92		
Bifidobacterium adolescentis		12.89	0.33			
Bifidobacterium breve		5.22	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47753450	**Low discrimination**					**47753450**
					COC	
Actinomyces israelii		87.29	0.36	XYL 92	1	
Blautia producta		7.16	-.03	MLZ 13	99	
Bifidobacterium breve		5.55	0.10		1	
47753500	**Low discrimination**					**47753500**
					SPO	
Actinomyces israelii		62.94	0.68	XYL 92	1	
Clostridium butyricum		18.57	0.37		99	
Bifidobacterium adolescentis		11.75	0.49		1	
47753501	**Low discrimination**					**47753501**
					SPO	
Actinomyces naeslundii		50.65	0.46	ESL 12	1	
Actinomyces israelii		31.37	0.33	ARA 1	1	
Clostridium butyricum		9.25	0.03		99	
47753502	**Low discrimination**					**47753502**
Bifidobacterium adolescentis		87.78	0.83	XYL 87		
Actinomyces israelii		7.13	0.71			
Bifidobacterium breve		4.39	0.62			
47753503	**Low discrimination**					**47753503**
Bifidobacterium adolescentis		81.65	0.49	XYL 87		
Actinomyces naeslundii		7.14	0.46	URE 1		
Actinomyces israelii		6.63	0.36			
47753510	**Low discrimination**					**47753510**
Bifidobacterium adolescentis		45.85	0.50	XYL 87		
Actinomyces israelii		30.03	0.53	BGL 99		
Bifidobacterium breve		18.38	0.44			
47753512	**Good identification**					**47753512**
Bifidobacterium adolescentis		94.63	0.84	XYL 87		
Bifidobacterium breve		3.87	0.62			
Actinomyces israelii		0.94	0.56			
47753513	**Acceptable identification**					**47753513**
Bifidobacterium adolescentis		94.35	0.50	XYL 87		
Bifidobacterium breve		3.86	0.27	URE 1		
Actinomyces israelii		0.94	0.22			
47753520	**Low discrimination**					**47753520**
					SPO	
Bifidobacterium breve		44.07	0.49	ARA 1	1	
Clostridium ramosum		22.73	0.33	BGL 91	92	
Actinomyces israelii		17.31	0.47		1	
47753521	**Low discrimination**					**47753521**
					SPO	
Actinomyces naeslundii		86.76	0.46	ESL 12	1	
Bifidobacterium breve		8.73	0.15	ARA 1	1	
Clostridium ramosum		4.51	-.01		92	
47753522	**Low discrimination**					**47753522**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		68.69	0.66	ARA 1	1	
Clostridium ramosum		14.02	0.43		92	
Bifidobacterium adolescentis		7.80	0.49		1	
47753530	**Low discrimination**					**47753530**
Bifidobacterium breve		81.64	0.49	ARA 1		
Bifidobacterium dentium		10.56	0.15	BGL 91		
Actinomyces israelii		4.79	0.33			
47753532	**Low discrimination**					**47753532**
Bifidobacterium breve		77.73	0.66	ARA 1		
Bifidobacterium adolescentis		10.79	0.50			
Bifidobacterium dentium		10.06	0.32			
47753540	**Low discrimination**					**47753540**
					COC	
Actinomyces israelii		68.19	0.61	XYL 92	1	
Bifidobacterium breve		16.03	0.44		1	
Blautia producta		6.89	0.24		99	
47753542	**Low discrimination**					**47753542**
Bifidobacterium adolescentis		59.96	0.67	XYL 87		
Bifidobacterium breve		24.27	0.62	SOR 11		
Actinomyces israelii		15.32	0.64			
47753552	**Low discrimination**					**47753552**
Bifidobacterium adolescentis		73.12	0.69	XYL 87		
Bifidobacterium breve		24.22	0.62	SOR 11		
Actinomyces israelii		2.28	0.49			
47753560	**Low discrimination**					**47753560**
					COC	
Bifidobacterium breve		82.22	0.49	ARA 1	1	
Actinomyces israelii		12.55	0.40	BGL 91	1	
Blautia producta		2.21	0.07		99	
47753562	**Very good identification**					**47753562**
Bifidobacterium breve		96.02	0.66	ARA 1		
Actinomyces israelii		2.18	0.43			
Bifidobacterium adolescentis		1.35	0.33			
47753570	**Acceptable identification**					**47753570**
					COC	
Bifidobacterium breve		90.63	0.49	ARA 1	1	
Blautia producta		7.31	0.15	BGL 91	99	
Actinomyces israelii		2.07	0.26		1	
47753572	**Very good identification**					**47753572**
Bifidobacterium breve		97.78	0.66	ARA 1		
Bifidobacterium adolescentis		1.68	0.35			
Actinomyces israelii		0.33	0.29			
47753600	**Good identification**					**47753600**
Actinomyces israelii		100.0	0.38	XYL 92		
47753602	**Low discrimination**					**47753602**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		86.50	0.41	XYL 92			
Clostridium clostridioforme		7.72	0.14	RHA 7			
Bifidobacterium adolescentis		5.78	0.15				
47753700	**Low discrimination**						**47753700**
					SPO		
Actinomyces israelii		85.22	0.48	XYL 92	1		
Clostridium butyricum		10.33	0.12	RHA 7	99		
Bifidobacterium adolescentis		2.14	0.15		1		
47753702	**Low discrimination**						**47753702**
Bifidobacterium adolescentis		57.07	0.49	XYL 87			
Actinomyces israelii		34.52	0.51	RHA 1			
Bifidobacterium breve		2.86	0.27				
47753712	**Low discrimination**						**47753712**
Bifidobacterium adolescentis		77.61	0.50	XYL 87			
Clostridium clostridioforme		13.07	0.34	RHA 1			
Actinomyces israelii		5.74	0.37				
47753720	**Low discrimination**						**47753720**
					SPO	GRA	
Clostridium ramosum		89.42	0.35	TRE 94	92	50	
Actinomyces israelii		3.87	0.28	ARA 1	1	99	
Bacteroides vulgatus		3.41	0.05		1	1	
47753722	**Low discrimination**						**47753722**
					SPO	GRA	
Clostridium ramosum		82.90	0.45	TRE 94	92	50	
Prevotella oralis		5.94	0.25	ARA 1	1	1	
Bifidobacterium breve		3.10	0.32		1	99	
47753742	**Low discrimination**						**47753742**
Actinomyces israelii		54.42	0.44	XYL 92			
Bifidobacterium adolescentis		28.59	0.33	RHA 7			
Bifidobacterium breve		11.58	0.27				
47754000	**Acceptable identification**						**47754000**
					GRA	SPO	
Fusobacterium mortiferum		90.40	0.48	MLT 11	1	1	
Actinomyces israelii		5.85	0.30	XYL 7	99	1	
Clostridium butyricum		3.75	0.05		99	99	
47754100	**Low discrimination**						**47754100**
					GRA	SPO	
Fusobacterium mortiferum		62.48	0.54	MLT 11	1	1	
Clostridium butyricum		26.65	0.29	XYL 7	99	99	
Actinomyces israelii		6.93	0.40		99	1	
47754102	**Low discrimination**						**47754102**
					GRA		
Bifidobacterium adolescentis		48.70	0.50	GAL 99	99		
Bifidobacterium longum subsp. infantis		11.61	0.26	ARA 87	99		
Prevotella oralis		10.22	0.28		1		
47754112	**Low discrimination**						**47754112**
Bifidobacterium adolescentis		68.32	0.52	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		27.34	0.42	ARA 87			
Bifidobacterium breve		2.29	0.27				
47754122	**Low discrimination**						**47754122**
					CAT	GRA	
Prevotella oralis		72.76	0.50	GAL 99	1	1	
Bacteroides fragilis		8.50	0.25	XYL 11	99	1	
Bifidobacterium longum subsp. infantis		7.73	0.30		1	99	
47754132	**Low discrimination**						**47754132**
					GRA		
Clostridium clostridioforme		65.45	0.42	TRE 88	50		
Parabacteroides distasonis		16.57	0.37	RHA 88	1		
Bifidobacterium breve		9.74	0.32	ARA 94	99		
47754152	**Low discrimination**						**47754152**
Bifidobacterium adolescentis		73.15	0.36	GAL 99			
Bifidobacterium breve		19.82	0.27	ARA 87			
Actinomyces israelii		4.64	0.22	SOR 11			
47754202	**Good identification**						**47754202**
					GRA		
Clostridium clostridioforme		96.38	0.36	TRE 88	50		
Prevotella oralis		2.35	0.02	ARA 94	1		
Actinomyces israelii		1.27	0.14		99		
47754212	**Good identification**						**47754212**
Clostridium clostridioforme		100.0	0.48	TRE 88			
47754222	**Low discrimination**						**47754222**
Clostridium clostridioforme		66.96	0.36	TRE 88			
Prevotella oralis		31.06	0.24	ARA 94			
Parabacteroides distasonis		1.98	0.16				
47754232	**Good identification**						**47754232**
Clostridium clostridioforme		99.65	0.48	TRE 88			
Parabacteroides distasonis		0.35	0.11	ARA 94			
47754302	**Low discrimination**						**47754302**
					GRA		
Clostridium clostridioforme		84.78	0.45	TRE 88	50		
Prevotella oralis		9.68	0.22	ARA 94	1		
Eubacterium contortum		1.82	0.11		99		
47754312	**Very good identification**						**47754312**
					GRA		
Clostridium clostridioforme		99.55	0.57	TRE 88	50		
Bifidobacterium adolescentis		0.34	0.18	ARA 94	99		
Parabacteroides distasonis		0.10	0.11		1		
47754322	**Low discrimination**						**47754322**
Prevotella oralis		51.23	0.44	GAL 99			
Clostridium clostridioforme		23.60	0.45	XYL 11			
Parabacteroides distasonis		20.68	0.50				
47754332	**Good identification**						**47754332**
Clostridium clostridioforme		90.15	0.57	TRE 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		9.33	0.45	ARA 94			
Prevotella oralis		0.43	0.10				
47754402	**Low discrimination**						**47754402**
					SPO		
Clostridium clostridioforme		75.51	0.42	TRE 88	66		
Bifidobacterium adolescentis		10.16	0.30	RHA 88	1		
Clostridium butyricum		9.26	0.15		99		
47754412	**Good identification**						**47754412**
Clostridium clostridioforme		96.18	0.53	TRE 88			
Bifidobacterium adolescentis		3.47	0.32	RHA 88			
Actinomyces israelii		0.21	0.17				
47754422	**Acceptable identification**						**47754422**
					GRA	SPO	
Clostridium clostridioforme		94.83	0.42	TRE 88	50	66	
Bacteroides uniformis		4.16	0.06	RHA 88	1	1	
Clostridium butyricum		1.01	-.03		99	99	
47754432	**Good identification**						**47754432**
Clostridium clostridioforme		100.0	0.53	TRE 88			
47754500	**Good identification**						**47754500**
					SPO		
Clostridium butyricum		95.37	0.45	TRE 94	99		
Actinomyces israelii		2.26	0.39	GAL 99	1		
Bifidobacterium adolescentis		1.70	0.30		1		
47754502	**Low discrimination**						**47754502**
					SPO		
Bifidobacterium adolescentis		66.94	0.64	GAL 99	1		
Clostridium clostridioforme		16.82	0.51		66		
Clostridium butyricum		14.79	0.38		99		
47754512	**Low discrimination**						**47754512**
					SPO		
Bifidobacterium adolescentis		51.30	0.66	GAL 99	1		
Clostridium clostridioforme		48.05	0.62		66		
Clostridium butyricum		0.49	0.17		99		
47754522	**Low discrimination**						**47754522**
Clostridium clostridioforme		53.15	0.51	TRE 88			
Bacteroides uniformis		34.12	0.35	RHA 88			
Parabacteroides distasonis		4.45	0.38				
47754532	**Very good identification**						**47754532**
					GRA		
Clostridium clostridioforme		97.57	0.62	TRE 88	50		
Bifidobacterium adolescentis		1.05	0.32	RHA 88	99		
Parabacteroides distasonis		0.97	0.34		1		
47754542	**Low discrimination**						**47754542**
					SPO		
Bifidobacterium adolescentis		80.69	0.49	GAL 99	1		
Clostridium butyricum		12.54	0.20	SOR 11	99		
Actinomyces israelii		5.11	0.35		1		
47754552	**Acceptable identification**						**47754552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		91.66	0.50	GAL 99		
Clostridium clostridioforme		7.02	0.28	SOR 11		
Actinomyces israelii		0.71	0.21			
47754602	**Very good identification**					**47754602**
Clostridium clostridioforme		100.0	0.57	TRE 88		
47754612	**Very good identification**					**47754612**
Clostridium clostridioforme		100.0	0.68	TRE 88		
47754622	**Very good identification**					**47754622**
Clostridium clostridioforme		99.07	0.57	TRE 88		
Bacteroides uniformis		0.82	0.08			
Parabacteroides distasonis		0.12	0.12			
47754632	**Very good identification**					**47754632**
Clostridium clostridioforme		100.0	0.68	TRE 88		
47754702	**Very good identification**					**47754702**
					SPO	
Clostridium clostridioforme		98.75	0.66	TRE 88	66	
Bifidobacterium adolescentis		0.54	0.30		1	
Clostridium butyricum		0.37	0.13		99	
47754710	**Good identification**					**47754710**
Clostridium clostridioforme		98.92	0.43	TRE 88		
Clostridium butyricum		1.08	-.02	BGL 99		
47754712	**Very good identification**					**47754712**
Clostridium clostridioforme		99.85	0.77	TRE 88		
Bifidobacterium adolescentis		0.15	0.32			
47754713	**Good identification**					**47754713**
Clostridium clostridioforme		100.0	0.43	TRE 88		
47754722	**Low discrimination**					**47754722**
Clostridium clostridioforme		85.43	0.66	TRE 88		
Bacteroides uniformis		10.33	0.37			
Parabacteroides distasonis		2.93	0.46			
47754730	**Good identification**					**47754730**
Clostridium clostridioforme		98.41	0.43	TRE 88		
Bacteroides vulgatus		1.59	-.01	BGL 99		
47754732	**Very good identification**					**47754732**
Clostridium clostridioforme		99.39	0.77	TRE 88		
Parabacteroides distasonis		0.40	0.42			
Bacteroides ovatus		0.18	0.20			
47754733	**Good identification**					**47754733**
Clostridium clostridioforme		100.0	0.43	TRE 88		
47754752	**Good identification**					**47754752**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		98.23	0.43	TRE 88			
Bifidobacterium adolescentis		1.77	0.16	SOR 1			
47754772	**Good identification**						**47754772**
Clostridium clostridioforme		99.26	0.43	TRE 88			
Bacteroides ovatus		0.74	-.04	SOR 1			
47755000	**Low discrimination**						**47755000**
					GRA	SPO	
Actinomyces israelii		50.10	0.64		99	1	
Clostridium butyricum		32.10	0.40		99	99	
Fusobacterium mortiferum		11.73	0.51		1	1	
47755002	**Low discrimination**						**47755002**
					GRA		
Actinomyces israelii		52.24	0.67		99		
Prevotella oralis		16.99	0.42		1		
Bifidobacterium adolescentis		12.81	0.50		99		
47755010	**Low discrimination**						**47755010**
					SPO		
Actinomyces israelii		74.42	0.50	MLZ 13	1		
Clostridium butyricum		16.80	0.18		99		
Lactobacillus jensenii		3.80	0.15		1		
47755012	**Low discrimination**						**47755012**
Bifidobacterium adolescentis		51.42	0.52	LAC 99			
Actinomyces israelii		25.63	0.53	ARA 87			
Clostridium clostridioforme		18.44	0.41				
47755020	**Low discrimination**						**47755020**
					GRA	SPO	
Prevotella oralis		70.07	0.42	XYL 11	1	1	
Actinomyces israelii		9.17	0.44	LAC 94	99	1	
Clostridium butyricum		8.01	0.21	BGL 95	99	99	
47755022	**Good identification**						**47755022**
					CAT	GRA	
Prevotella oralis		92.38	0.64	XYL 11	1	1	
Bacteroides fragilis		1.71	0.25	LAC 94	99	1	
Bifidobacterium longum subsp. infantis		1.55	0.30		1	99	
47755032	**Low discrimination**						**47755032**
Clostridium clostridioforme		45.84	0.41	TRE 88			
Prevotella oralis		26.62	0.30	RHA 88			
Parabacteroides distasonis		12.95	0.37	ARA 94			
47755040	**Low discrimination**						**47755040**
					GRA	SPO	
Actinomyces israelii		86.71	0.57		99	1	
Clostridium butyricum		12.43	0.21		99	99	
Fusobacterium mortiferum		0.53	0.17		1	1	
47755042	**Low discrimination**						**47755042**
					SPO		
Actinomyces israelii		86.49	0.60		1		
Bifidobacterium adolescentis		6.74	0.35		1		
Clostridium butyricum		3.21	0.14		99		
47755050	**Acceptable identification**						**47755050**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		92.93	0.43	MLZ 13	1		
Clostridium butyricum		4.69	-.01		99		
Bifidobacterium breve		2.38	0.10		1		
47755052	**Low discrimination**						**47755052**
Actinomyces israelii		54.29	0.46	MLZ 13			
Bifidobacterium adolescentis		34.61	0.36				
Bifidobacterium breve		9.38	0.27				
47755060	**Low discrimination**						**47755060**
					GRA	SPO	
Actinomyces israelii		61.82	0.37	NAG 6	99	1	
Prevotella oralis		12.27	0.08		1	1	
Clostridium butyricum		12.07	0.03		99	99	
47755100	**Low discrimination**						**47755100**
					SPO		
Clostridium butyricum		68.50	0.63	TRE 94	99		
Actinomyces israelii		17.82	0.74	ARA 90	1		
Bifidobacterium longum subsp. infantis		3.82	0.43		1		
47755102	**Low discrimination**						**47755102**
					GRA		
Bifidobacterium adolescentis		54.20	0.84	ARA 87	99		
Bifidobacterium longum subsp. infantis		12.92	0.60		99		
Prevotella oralis		11.37	0.63		1		
47755103	**Low discrimination**						**47755103**
					GRA		
Bifidobacterium adolescentis		53.49	0.50	ARA 87	99		
Bifidobacterium longum subsp. infantis		12.75	0.26	URE 1	99		
Prevotella oralis		11.22	0.28		1		
47755110	**Low discrimination**						**47755110**
					SPO		
Clostridium butyricum		39.77	0.41	TRE 94	99		
Actinomyces israelii		29.37	0.60	ARA 90	1		
Bifidobacterium adolescentis		22.09	0.52	MLZ 5	1		
47755112	**Good identification**						**47755112**
Bifidobacterium adolescentis		92.53	0.86	ARA 87			
Bifidobacterium breve		3.10	0.62				
Actinomyces israelii		1.86	0.63				
47755113	**Acceptable identification**						**47755113**
Bifidobacterium adolescentis		92.97	0.52	ARA 87			
Bifidobacterium breve		3.11	0.27	URE 1			
Actinomyces israelii		1.87	0.29				
47755120	**Low discrimination**						**47755120**
					GRA	SPO	
Prevotella oralis		52.19	0.63	XYL 11	1	1	
Clostridium paraputrificum		14.16	0.47	BGL 95	99	99	
Bifidobacterium longum subsp. infantis		10.42	0.47		99	1	
47755122	**Low discrimination**						**47755122**
					CAT	GRA	
Prevotella oralis		74.34	0.85	XYL 11	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides fragilis		8.69	0.60		99	1	
Bifidobacterium longum subsp. infantis		7.90	0.64		1	99	
47755123	**Low discrimination**						**47755123**
					CAT	GRA	
Prevotella oralis		73.89	0.50	XYL 11	1	1	
Bacteroides fragilis		8.63	0.25	URE 1	99	1	
Bifidobacterium longum subsp. infantis		7.85	0.30		1	99	
47755130	**Low discrimination**						**47755130**
					GRA	SPO	
Bifidobacterium breve		46.75	0.49	XYL 1	99	1	
Prevotella oralis		13.76	0.28	BGL 91	1	1	
Clostridium butyricum		12.55	0.23		99	99	
47755132	**Low discrimination**						**47755132**
					GRA		
Parabacteroides distasonis		44.37	0.71		1		
Bifidobacterium breve		26.09	0.66		99		
Prevotella oralis		14.43	0.50		1		
47755133	**Low discrimination**						**47755133**
					GRA		
Parabacteroides distasonis		45.34	0.37	URE 1	1		
Bifidobacterium breve		26.66	0.32		99		
Prevotella oralis		14.75	0.16		1		
47755140	**Low discrimination**						**47755140**
					SPO		
Actinomyces israelii		49.96	0.67		1		
Clostridium butyricum		42.95	0.45		99		
Bifidobacterium breve		4.74	0.44		1		
47755142	**Low discrimination**						**47755142**
Bifidobacterium adolescentis		49.71	0.69	ARA 87			
Actinomyces israelii		25.77	0.70	SOR 11			
Bifidobacterium breve		16.47	0.62				
47755150	**Low discrimination**						**47755150**
					SPO		
Actinomyces israelii		45.78	0.53	MLZ 13	1		
Bifidobacterium breve		29.04	0.44		1		
Clostridium butyricum		13.86	0.23		99		
47755152	**Low discrimination**						**47755152**
Bifidobacterium adolescentis		74.60	0.70	ARA 87			
Bifidobacterium breve		20.22	0.62	SOR 11			
Actinomyces israelii		4.73	0.56				
47755153	**Low discrimination**						**47755153**
Bifidobacterium adolescentis		74.94	0.36	ARA 87			
Bifidobacterium breve		20.31	0.27	SOR 11			
Actinomyces israelii		4.75	0.22	URE 1			
47755160	**Low discrimination**						**47755160**
					SPO		
Bifidobacterium breve		34.97	0.49	XYL 1	1		
Clostridium paraputrificum		20.82	0.27	BGL 91	99		
Clostridium butyricum		15.51	0.27		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47755162	**Low discrimination**						**47755162**
					CAT	GRA	
Bifidobacterium breve		53.36	0.66	XYL 1	1	99	
Prevotella oralis		29.52	0.50		1	1	
Bacteroides fragilis		3.45	0.25		99	1	
47755170	**Acceptable identification**						**47755170**
					SPO		
Bifidobacterium breve		92.08	0.49	XYL 1	1		
Actinomyces israelii		5.21	0.33	BGL 91	1		
Clostridium butyricum		2.15	0.05		99		
47755172	**Good identification**						**47755172**
					GRA		
Bifidobacterium breve		94.47	0.66	XYL 1	99		
Bifidobacterium adolescentis		1.98	0.36		99		
Parabacteroides distasonis		1.62	0.37		1		
47755200	**Low discrimination**						**47755200**
					SPO		
Actinomyces israelii		51.46	0.45	RHA 7	1		
Eubacterium contortum		24.05	0.24		1		
Clostridium butyricum		13.55	0.14		99		
47755202	**Low discrimination**						**47755202**
					GRA		
Clostridium clostridioforme		42.92	0.45	TRE 88	50		
Prevotella oralis		34.58	0.36	ARA 94	1		
Actinomyces israelii		18.68	0.48		99		
47755212	**Good identification**						**47755212**
Clostridium clostridioforme		97.94	0.56	TRE 88			
Actinomyces israelii		1.40	0.34	ARA 94			
Bifidobacterium adolescentis		0.38	0.18				
47755220	**Low discrimination**						**47755220**
					GRA		
Prevotella oralis		82.47	0.36	XYL 11	1		
Bacteroides vulgatus		9.94	0.14	LAC 94	1		
Eubacterium contortum		2.45	0.11	BGL 95	99		
47755222	**Low discrimination**						**47755222**
Prevotella oralis		86.31	0.58	XYL 11			
Clostridium clostridioforme		5.64	0.45	LAC 94			
Parabacteroides distasonis		5.51	0.50				
47755232	**Low discrimination**						**47755232**
Clostridium clostridioforme		86.85	0.56	TRE 88			
Parabacteroides distasonis		10.04	0.45	ARA 94			
Prevotella oralis		2.95	0.24				
47755240	**Acceptable identification**						**47755240**
					SPO		
Actinomyces israelii		94.44	0.38	RHA 7	1		
Clostridium butyricum		5.56	-.04		99		
47755242	**Acceptable identification**						**47755242**
					GRA		
Actinomyces israelii		90.27	0.41	RHA 7	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		5.39	0.10		50	
Prevotella oralis		4.34	0.02		1	
47755300	**Low discrimination**					**47755300**
					SPO	
Eubacterium contortum		77.50	0.58	CEL 1	1	
Clostridium butyricum		10.59	0.37		99	
Actinomyces israelii		6.70	0.55		1	
47755302	**Low discrimination**					**47755302**
					GRA	
Prevotella oralis		52.87	0.56	XYL 11	1	
Clostridium clostridioforme		14.03	0.54	NAG 95	50	
Eubacterium contortum		9.92	0.45		99	
47755310	**Low discrimination**					**47755310**
Actinomyces israelii		29.15	0.41	RHA 7		
Clostridium clostridioforme		25.90	0.31	MLZ 13		
Eubacterium contortum		22.80	0.24			
47755312	**Low discrimination**					**47755312**
					GRA	
Clostridium clostridioforme		84.60	0.65	TRE 88	50	
Bifidobacterium adolescentis		9.61	0.52	ARA 94	99	
Parabacteroides distasonis		2.92	0.45		1	
47755320	**Low discrimination**					**47755320**
					GRA	
Prevotella oralis		47.89	0.56	XYL 11	1	
Bacteroides vulgatus		36.45	0.49	BGL 95	1	
Eubacterium contortum		8.99	0.45		99	
47755322	**Low discrimination**					**47755322**
Prevotella oralis		66.33	0.78	XYL 11		
Parabacteroides distasonis		26.77	0.84			
Bacteroides uniformis		5.03	0.53			
47755323	**Low discrimination**					**47755323**
Prevotella oralis		66.43	0.44	XYL 11		
Parabacteroides distasonis		26.81	0.50	URE 1		
Bacteroides uniformis		5.04	0.19			
47755330	**Low discrimination**					**47755330**
Parabacteroides distasonis		54.66	0.45	BGL 99		
Clostridium clostridioforme		16.00	0.31			
Prevotella oralis		13.24	0.22			
47755332	**Low discrimination**					**47755332**
Parabacteroides distasonis		73.84	0.80			
Clostridium clostridioforme		21.61	0.65			
Prevotella oralis		3.43	0.44			
47755333	**Low discrimination**					**47755333**
Parabacteroides distasonis		74.21	0.45	URE 1		
Clostridium clostridioforme		21.72	0.31			
Prevotella oralis		3.45	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47755340	**Low discrimination**						**47755340**
					SPO		
Actinomyces israelii		59.48	0.48	RHA 7	1		
Clostridium butyricum		21.01	0.19		99		
Eubacterium contortum		17.87	0.24		1		
47755342	**Low discrimination**						**47755342**
					GRA		
Actinomyces israelii		57.56	0.51	RHA 7	99		
Bifidobacterium adolescentis		14.90	0.35		99		
Prevotella oralis		10.84	0.22		1		
47755352	**Low discrimination**						**47755352**
Bifidobacterium adolescentis		40.20	0.36	RHA 1			
Clostridium clostridioforme		28.92	0.31	ARA 87			
Actinomyces israelii		18.99	0.37	SOR 11			
47755362	**Low discrimination**						**47755362**
Prevotella oralis		63.28	0.44	XYL 11			
Parabacteroides distasonis		25.54	0.50	SOR 1			
Bacteroides uniformis		4.80	0.19				
47755372	**Low discrimination**						**47755372**
					GRA		
Parabacteroides distasonis		63.20	0.45	SOR 1	1		
Clostridium clostridioforme		18.50	0.31		50		
Bifidobacterium breve		12.39	0.32		99		
47755400	**Low discrimination**						**47755400**
					SPO		
Clostridium butyricum		87.14	0.56	TRE 94	99		
Actinomyces israelii		12.36	0.62	LAC 96	1		
Bifidobacterium adolescentis		0.38	0.30		1		
47755402	**Low discrimination**						**47755402**
					SPO		
Bifidobacterium adolescentis		37.88	0.64	LAC 99	1		
Clostridium butyricum		34.51	0.49		99		
Actinomyces israelii		18.88	0.66		1		
47755412	**Low discrimination**						**47755412**
Bifidobacterium adolescentis		51.54	0.66	LAC 99			
Clostridium clostridioforme		43.26	0.61				
Actinomyces israelii		3.14	0.51				
47755420	**Low discrimination**						**47755420**
					GRA	SPO	
Clostridium butyricum		82.28	0.38	TRE 94	99	99	
Actinomyces israelii		8.57	0.42	LAC 96	99	1	
Bacteroides vulgatus		6.10	0.17	NAG 8	1	1	
47755422	**Low discrimination**						**47755422**
					GRA	SPO	
Bacteroides uniformis		44.41	0.40	IND 98	1	1	
Clostridium clostridioforme		30.68	0.50	LAC 98	50	66	
Clostridium butyricum		10.80	0.31		99	99	
47755432	**Very good identification**						**47755432**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		96.42	0.61	TRE 88	50		
Bifidobacterium adolescentis		1.16	0.32	RHA 88	99		
Parabacteroides distasonis		1.07	0.34		1		
47755440	**Low discrimination**						**47755440**
					SPO		
Clostridium butyricum		60.96	0.38	TRE 94	99		
Actinomyces israelii		38.67	0.55	LAC 96	1		
Bifidobacterium adolescentis		0.37	0.15	SOR 8	1		
47755442	**Low discrimination**						**47755442**
					SPO		
Actinomyces israelii		48.59	0.58		1		
Bifidobacterium adolescentis		30.98	0.49		1		
Clostridium butyricum		19.86	0.31		99		
47755450	**Low discrimination**						**47755450**
					SPO		
Actinomyces israelii		59.72	0.41	MLZ 13	1		
Clostridium butyricum		33.15	0.16		99		
Bifidobacterium adolescentis		4.72	0.16		1		
47755452	**Low discrimination**						**47755452**
Bifidobacterium adolescentis		77.64	0.50	LAC 99			
Actinomyces israelii		14.89	0.44	SOR 11			
Clostridium clostridioforme		5.33	0.27				
47755462	**Low discrimination**						**47755462**
					GRA	SPO	
Actinomyces israelii		47.45	0.38	NAG 6	99	1	
Clostridium butyricum		26.42	0.13		99	99	
Bacteroides uniformis		12.63	0.06		1	1	
47755500	**Very good identification**						**47755500**
					SPO		
Clostridium butyricum		95.93	0.80	TRE 94	99		
Actinomyces israelii		2.27	0.73		1		
Bifidobacterium adolescentis		1.71	0.64		1		
47755501	**Good identification**						**47755501**
					SPO		
Clostridium butyricum		95.90	0.45	TRE 94	99		
Actinomyces israelii		2.27	0.39	URE 1	1		
Bifidobacterium adolescentis		1.71	0.30		1		
47755502	**Low discrimination**						**47755502**
					SPO		
Bifidobacterium adolescentis		79.92	0.99		1		
Clostridium butyricum		17.65	0.73		99		
Actinomyces israelii		1.61	0.76		1		
47755503	**Low discrimination**						**47755503**
					SPO		
Bifidobacterium adolescentis		79.98	0.64	URE 1	1		
Clostridium butyricum		17.67	0.38		99		
Actinomyces israelii		1.61	0.42		1		
47755510	**Low discrimination**						**47755510**
					SPO		
Clostridium butyricum		66.80	0.58	TRE 94	99		
Bifidobacterium adolescentis		27.59	0.66	MLZ 5	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		4.48	0.59		1		
47755512	**Very good identification**						**47755512**
					SPO		
Bifidobacterium adolescentis		96.04	1.00		1		
Clostridium clostridioforme		2.73	0.70		66		
Clostridium butyricum		0.91	0.51		99		
47755513	**Very good identification**						**47755513**
					SPO		
Bifidobacterium adolescentis		96.06	0.66	URE 1	1		
Clostridium clostridioforme		2.73	0.36		66		
Clostridium butyricum		0.91	0.17		99		
47755520	**Low discrimination**						**47755520**
					GRA	SPO	
Clostridium butyricum		72.57	0.61	TRE 94	99	99	
Bacteroides vulgatus		22.18	0.52	NAG 8	1	1	
Bacteroides uniformis		2.39	0.35		1	1	
47755522	**Low discrimination**						**47755522**
					GRA	SPO	
Bacteroides uniformis		70.15	0.69	IND 98	1	1	
Parabacteroides distasonis		9.15	0.72		1	1	
Clostridium butyricum		8.35	0.54		99	99	
47755530	**Low discrimination**						**47755530**
					SPO		
Clostridium butyricum		68.32	0.40	TRE 94	99		
Clostridium clostridioforme		9.21	0.36	MLZ 5	66		
Bifidobacterium longum subsp. longum		4.08	0.32	NAG 8	1		
47755532	**Low discrimination**						**47755532**
					GRA		
Clostridium clostridioforme		54.36	0.70	TRE 88	50		
Bifidobacterium adolescentis		19.35	0.66	RHA 88	99		
Parabacteroides distasonis		17.76	0.68		1		
47755533	**Low discrimination**						**47755533**
					GRA		
Clostridium clostridioforme		54.87	0.36	TRE 88	50		
Bifidobacterium adolescentis		19.53	0.32	RHA 88	99		
Parabacteroides distasonis		17.93	0.34	URE 1	1		
47755540	**Low discrimination**						**47755540**
					SPO		
Clostridium butyricum		87.89	0.61	TRE 94	99		
Actinomyces israelii		9.29	0.66	SOR 8	1		
Bifidobacterium adolescentis		2.22	0.49		1		
47755542	**Low discrimination**						**47755542**
					SPO		
Bifidobacterium adolescentis		81.95	0.83	SOR 11	1		
Clostridium butyricum		12.73	0.54		99		
Actinomyces israelii		5.19	0.69		1		
47755543	**Low discrimination**						**47755543**
					SPO		
Bifidobacterium adolescentis		82.05	0.49	SOR 11	1		
Clostridium butyricum		12.75	0.20	URE 1	99		
Actinomyces israelii		5.20	0.35		1		
47755550	**Low discrimination**						**47755550**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium butyricum		45.49	0.40	TRE 94	99		
Bifidobacterium adolescentis		26.70	0.50	MLZ 5	1		
Actinomyces israelii		13.66	0.52	SOR 8	1		
47755552	**Very good identification**						**47755552**
					SPO		
Bifidobacterium adolescentis		98.25	0.85	SOR 11	1		
Actinomyces israelii		0.76	0.55		1		
Clostridium butyricum		0.66	0.33		99		
47755553	**Good identification**						**47755553**
					SPO		
Bifidobacterium adolescentis		98.58	0.50	SOR 11	1		
Actinomyces israelii		0.76	0.21	URE 1	1		
Clostridium butyricum		0.66	-.02		99		
47755560	**Low discrimination**						**47755560**
					GRA	SPO	
Clostridium butyricum		88.14	0.43	TRE 94	99	99	
Actinomyces israelii		6.84	0.45	NAG 8	99	1	
Bacteroides vulgatus		3.13	0.17	SOR 8	1	1	
47755562	**Low discrimination**						**47755562**
					GRA	SPO	
Clostridium butyricum		29.77	0.36	TRE 94	99	99	
Bacteroides uniformis		29.05	0.35	NAG 8	1	1	
Bifidobacterium adolescentis		22.25	0.49	SOR 8	99	1	
47755572	**Low discrimination**						**47755572**
					GRA		
Bifidobacterium adolescentis		65.30	0.50	NAG 1	99		
Clostridium clostridioforme		14.99	0.36	SOR 11	50		
Parabacteroides distasonis		4.90	0.34		1		
47755602	**Very good identification**						**47755602**
					SPO		
Clostridium clostridioforme		95.17	0.65	TRE 88	66		
Actinomyces israelii		2.16	0.46		1		
Clostridium butyricum		1.62	0.23		99		
47755610	**Acceptable identification**						**47755610**
					SPO		
Clostridium clostridioforme		90.96	0.42	TRE 88	66		
Clostridium butyricum		4.57	0.08	BGL 99	99		
Actinomyces israelii		4.47	0.29		1		
47755612	**Very good identification**						**47755612**
Clostridium clostridioforme		99.74	0.76	TRE 88			
Bifidobacterium adolescentis		0.16	0.32				
Actinomyces israelii		0.07	0.32				
47755613	**Good identification**						**47755613**
Clostridium clostridioforme		100.0	0.42	TRE 88			
47755622	**Low discrimination**						**47755622**
Clostridium clostridioforme		74.66	0.65	TRE 88			
Bacteroides uniformis		20.35	0.42				
Parabacteroides distasonis		2.86	0.46				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47755630	**Good identification**						**47755630**
Clostridium clostridioforme		98.23	0.42	TRE 88			
Bacteroides vulgatus		1.77	-.01	BGL 99			
47755632	**Very good identification**						**47755632**
Clostridium clostridioforme		99.28	0.76	TRE 88			
Parabacteroides distasonis		0.45	0.42				
Bacteroides ovatus		0.20	0.20				
47755633	**Good identification**						**47755633**
Clostridium clostridioforme		100.0	0.42	TRE 88			
47755640	**Low discrimination**						**47755640**
					SPO		
Actinomyces israelii		60.69	0.36	RHA 7	1		
Clostridium butyricum		39.31	0.12		99		
47755652	**Good identification**						**47755652**
Clostridium clostridioforme		95.35	0.42	TRE 88			
Actinomyces israelii		2.74	0.25	SOR 1			
Bifidobacterium adolescentis		1.92	0.16				
47755672	**Good identification**						**47755672**
Clostridium clostridioforme		99.17	0.42	TRE 88			
Bacteroides ovatus		0.83	-.04	SOR 1			
47755700	**Low discrimination**						**47755700**
					GRA	SPO	
Clostridium butyricum		54.10	0.54	TRE 94	99	99	
Eubacterium contortum		36.01	0.57	RHA 3	99	1	
Bacteroides vulgatus		4.23	0.34		1	1	
47755702	**Low discrimination**						**47755702**
					SPO		
Clostridium clostridioforme		70.11	0.74	TRE 88	66		
Bifidobacterium adolescentis		12.68	0.64		1		
Clostridium butyricum		8.58	0.47		99		
47755703	**Low discrimination**						**47755703**
					SPO		
Clostridium clostridioforme		70.47	0.40	TRE 88	66		
Bifidobacterium adolescentis		12.75	0.30	URE 1	1		
Clostridium butyricum		8.62	0.13		99		
47755710	**Low discrimination**						**47755710**
					SPO		
Clostridium clostridioforme		65.03	0.51	TRE 88	66		
Clostridium butyricum		23.40	0.32	BGL 99	99		
Actinomyces israelii		3.82	0.39		1		
47755712	**Good identification**						**47755712**
					GRA		
Clostridium clostridioforme		94.97	0.85	TRE 88	50		
Bifidobacterium adolescentis		4.61	0.66		99		
Parabacteroides distasonis		0.13	0.42		1		
47755713	**Good identification**						**47755713**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		95.37	0.51	TRE 88			
Bifidobacterium adolescentis		4.63	0.32	URE 1			
47755720	**Good identification**						**47755720**
					GRA		
Bacteroides vulgatus		94.76	0.68	CEL 5	1		
Bacteroides uniformis		1.57	0.37		1		
Eubacterium contortum		1.44	0.44		99		
47755722	**Low discrimination**						**47755722**
Bacteroides uniformis		60.10	0.71	IND 98			
Parabacteroides distasonis		17.04	0.81				
Clostridium clostridioforme		15.06	0.74				
47755723	**Low discrimination**						**47755723**
Bacteroides uniformis		60.19	0.37	IND 98			
Parabacteroides distasonis		17.06	0.46	URE 1			
Clostridium clostridioforme		15.09	0.40				
47755730	**Low discrimination**						**47755730**
Clostridium clostridioforme		56.07	0.51	TRE 88			
Bacteroides vulgatus		29.95	0.34	BGL 99			
Parabacteroides distasonis		7.49	0.42				
47755732	**Low discrimination**						**47755732**
Clostridium clostridioforme		83.01	0.85	TRE 88			
Parabacteroides distasonis		11.10	0.76				
Bacteroides ovatus		4.96	0.54				
47755733	**Low discrimination**						**47755733**
Clostridium clostridioforme		83.18	0.51	TRE 88			
Parabacteroides distasonis		11.12	0.42	URE 1			
Bacteroides ovatus		4.97	0.20				
47755740	**Low discrimination**						**47755740**
					COC	SPO	
Clostridium butyricum		62.19	0.36	TRE 94	1	99	
Actinomyces israelii		16.00	0.47	RHA 3	1	1	
Blautia producta		10.59	0.24	SOR 8	99	1	
47755742	**Low discrimination**						**47755742**
					SPO		
Bifidobacterium adolescentis		40.29	0.49	RHA 1	1		
Clostridium butyricum		19.17	0.29	SOR 11	99		
Actinomyces israelii		19.03	0.50		1		
47755752	**Low discrimination**						**47755752**
Clostridium clostridioforme		59.58	0.51	TRE 88			
Bifidobacterium adolescentis		35.38	0.50	SOR 1			
Actinomyces israelii		2.04	0.35				
47755762	**Low discrimination**						**47755762**
Bacteroides uniformis		49.42	0.37	IND 98			
Bacteroides ovatus		20.85	0.33	SOR 1			
Parabacteroides distasonis		14.01	0.46				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47755772	**Low discrimination**						**47755772**
Clostridium clostridioforme		71.39	0.51	TRE 88			
Bacteroides ovatus		17.58	0.30	SOR 1			
Parabacteroides distasonis		9.54	0.42				
47756000	**Good identification**						**47756000**
					GRA		
Actinomyces israelii		97.32	0.43	GAL 99	99		
Fusobacterium mortiferum		2.68	0.14		1		
47756002	**Good identification**						**47756002**
Actinomyces israelii		96.58	0.46	GAL 99			
Bifidobacterium adolescentis		3.42	0.15				
47756040	**Good identification**						**47756040**
Actinomyces israelii		100.0	0.36	GAL 99			
47756042	**Good identification**						**47756042**
Actinomyces israelii		100.0	0.39	GAL 99			
47756100	**Low discrimination**						**47756100**
					GRA	SPO	
Actinomyces israelii		89.99	0.53	GAL 99	99	1	
Clostridium butyricum		6.79	0.13		99	99	
Fusobacterium mortiferum		1.45	0.20		1	1	
47756102	**Low discrimination**						**47756102**
Actinomyces israelii		51.41	0.56	GAL 99			
Bifidobacterium adolescentis		45.05	0.49				
Bifidobacterium breve		2.25	0.27				
47756110	**Low discrimination**						**47756110**
Actinomyces israelii		84.41	0.39	GAL 99			
Bifidobacterium adolescentis		9.17	0.16	MLZ 13			
Bifidobacterium breve		3.68	0.10				
47756112	**Low discrimination**						**47756112**
Bifidobacterium adolescentis		83.35	0.50	GAL 99			
Actinomyces israelii		11.63	0.42	ARA 87			
Bifidobacterium breve		3.41	0.27				
47756140	**Good identification**						**47756140**
					SPO		
Actinomyces israelii		96.75	0.46	GAL 99	1		
Clostridium butyricum		1.63	-.05		99		
Bifidobacterium breve		1.62	0.10		1		
47756142	**Low discrimination**						**47756142**
Actinomyces israelii		71.88	0.49	GAL 99			
Bifidobacterium adolescentis		20.02	0.33				
Bifidobacterium breve		8.11	0.27				
47756302	**Low discrimination**						**47756302**
Actinomyces israelii		81.25	0.37	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		9.56	0.15	RHA 7			
Clostridium clostridioforme		9.20	0.11				
47756400	**Low discrimination**						**47756400**
					SPO		
Actinomyces israelii		87.86	0.41	GAL 99	1		
Clostridium butyricum		12.14	0.05		99		
47756402	**Low discrimination**						**47756402**
					SPO		
Actinomyces israelii		73.96	0.44	GAL 99	1		
Bifidobacterium adolescentis		21.42	0.29		1		
Clostridium butyricum		2.65	-.02		99		
47756442	**Acceptable identification**						**47756442**
Actinomyces israelii		91.57	0.37	GAL 99			
Bifidobacterium adolescentis		8.43	0.13				
47756500	**Low discrimination**						**47756500**
					SPO		
Actinomyces israelii		51.60	0.52	GAL 99	1		
Clostridium butyricum		42.80	0.29		99		
Bifidobacterium adolescentis		5.61	0.29		1		
47756502	**Low discrimination**						**47756502**
					SPO		
Bifidobacterium adolescentis		85.21	0.63	GAL 99	1		
Actinomyces israelii		11.89	0.55		1		
Clostridium butyricum		2.56	0.22		99		
47756510	**Low discrimination**						**47756510**
Actinomyces israelii		34.12	0.37	GAL 99			
Bifidobacterium adolescentis		30.31	0.30	MLZ 13			
Bifidobacterium dentium		23.26	0.10				
47756512	**Good identification**						**47756512**
Bifidobacterium adolescentis		90.23	0.64	GAL 99			
Bifidobacterium dentium		7.07	0.27				
Actinomyces israelii		1.54	0.40				
47756540	**Low discrimination**						**47756540**
					COC	SPO	
Actinomyces israelii		47.85	0.45	GAL 99	1	1	
Blautia producta		41.63	0.24		99	1	
Clostridium butyricum		8.87	0.11		1	99	
47756542	**Low discrimination**						**47756542**
					COC		
Bifidobacterium adolescentis		67.19	0.47	GAL 99	1		
Actinomyces israelii		29.49	0.48	SOR 11	1		
Blautia producta		1.90	0.07		99		
47756552	**Low discrimination**						**47756552**
					COC		
Bifidobacterium adolescentis		88.53	0.49	GAL 99	1		
Blautia producta		6.15	0.15	SOR 11	99		
Actinomyces israelii		4.75	0.33		1		
47756712	**Low discrimination**						**47756712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		87.44	0.43	TRE 88		
Bifidobacterium adolescentis		10.42	0.30	MAN 1		
Actinomyces israelii		1.32	0.21			
47756732	**Acceptable identification**					**47756732**
					GRA	
Clostridium clostridioforme		92.91	0.43	TRE 88	50	
Bacteroides ovatus		5.55	0.12	MAN 1	1	
Bifidobacterium dentium		1.54	-.03		99	
47757000	**Very good identification**					**47757000**
					SPO	
Actinomyces israelii		98.51	0.77		1	
Clostridium butyricum		1.24	0.23		99	
Actinomyces naeslundii		0.11	0.35		1	
47757001	**Low discrimination**					**47757001**
Actinomyces israelii		86.10	0.43	URE 1		
Actinomyces naeslundii		13.90	0.38			
47757002	**Very good identification**					**47757002**
					SPO	
Actinomyces israelii		95.93	0.80		1	
Bifidobacterium adolescentis		3.40	0.49		1	
Clostridium butyricum		0.31	0.16		99	
47757003	**Low discrimination**					**47757003**
Actinomyces israelii		87.49	0.46	URE 1		
Actinomyces naeslundii		9.42	0.38			
Bifidobacterium adolescentis		3.10	0.15			
47757010	**Very good identification**					**47757010**
					SPO	
Actinomyces israelii		98.83	0.63	MLZ 13	1	
Clostridium butyricum		0.44	0.01		99	
Bifidobacterium adolescentis		0.43	0.16		1	
47757012	**Low discrimination**					**47757012**
Actinomyces israelii		76.45	0.66	MLZ 13		
Bifidobacterium adolescentis		22.14	0.50			
Bifidobacterium breve		0.91	0.27			
47757020	**Good identification**					**47757020**
					SPO	
Actinomyces israelii		93.66	0.56	NAG 6	1	
Clostridium butyricum		1.60	0.05		99	
Actinomyces naeslundii		1.60	0.35		1	
47757021	**Low discrimination**					**47757021**
Actinomyces naeslundii		71.67	0.38	ESL 12		
Actinomyces israelii		28.33	0.22	XYL 1		
47757022	**Low discrimination**					**47757022**
					GRA	
Actinomyces israelii		78.17	0.59	NAG 6	99	
Prevotella oralis		13.49	0.30		1	
Bifidobacterium breve		3.86	0.32		99	
47757023	**Low discrimination**					**47757023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		62.77	0.38	ESL 12		
Actinomyces israelii		37.23	0.25	XYL 1		
47757030	**Acceptable identification**					**47757030**
					SPO	
Actinomyces israelii		90.03	0.42	MLZ 13	1	
Clostridium tertium		5.56	0.07	NAG 6	99	
Bifidobacterium breve		4.41	0.15		1	
47757032	**Low discrimination**					**47757032**
					SPO	
Actinomyces israelii		64.01	0.45	MLZ 13	1	
Bifidobacterium breve		21.14	0.32	NAG 6	1	
Clostridium tertium		7.90	0.15		99	
47757040	**Very good identification**					**47757040**
					SPO	
Actinomyces israelii		99.60	0.70		1	
Clostridium butyricum		0.28	0.05		99	
Bifidobacterium breve		0.07	0.10		1	
47757041	**Acceptable identification**					**47757041**
Actinomyces israelii		93.17	0.36	URE 1		
Actinomyces naeslundii		6.83	0.25			
47757042	**Very good identification**					**47757042**
Actinomyces israelii		98.34	0.73			
Bifidobacterium adolescentis		1.11	0.33			
Bifidobacterium breve		0.45	0.27			
47757043	**Good identification**					**47757043**
Actinomyces israelii		95.34	0.39	URE 1		
Actinomyces naeslundii		4.66	0.25			
47757050	**Good identification**					**47757050**
					COC	
Actinomyces israelii		98.98	0.56	MLZ 13	1	
Blautia producta		0.58	-.03		99	
Bifidobacterium breve		0.45	0.10		1	
47757052	**Low discrimination**					**47757052**
Actinomyces israelii		89.08	0.59	MLZ 13		
Bifidobacterium adolescentis		8.20	0.35			
Bifidobacterium breve		2.72	0.27			
47757060	**Good identification**					**47757060**
Actinomyces israelii		97.41	0.49	NAG 6		
Bifidobacterium breve		1.83	0.15			
Actinomyces naeslundii		0.75	0.22			
47757062	**Low discrimination**					**47757062**
Actinomyces israelii		88.34	0.52	NAG 6		
Bifidobacterium breve		11.21	0.32			
Actinomyces naeslundii		0.46	0.22			
47757070	**Low discrimination**					**47757070**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		88.81	0.35	MLZ 13		
Bifidobacterium breve		11.19	0.15	NAG 6		
47757072	**Low discrimination**					**47757072**
Actinomyces israelii		54.08	0.38	MLZ 13		
Bifidobacterium breve		45.92	0.32	NAG 6		
47757100	**Good identification**					**47757100**
					SPO	
Actinomyces israelii		90.92	0.87		1	
Clostridium butyricum		6.86	0.47		99	
Bifidobacterium adolescentis		1.21	0.49		1	
47757101	**Low discrimination**					**47757101**
					SPO	
Actinomyces israelii		82.64	0.53	URE 1	1	
Actinomyces naeslundii		9.49	0.46		1	
Clostridium butyricum		6.23	0.13		99	
47757102	**Low discrimination**					**47757102**
Actinomyces israelii		51.25	0.90			
Bifidobacterium adolescentis		44.91	0.83			
Bifidobacterium breve		2.25	0.62			
47757103	**Low discrimination**					**47757103**
Actinomyces israelii		49.46	0.56	URE 1		
Bifidobacterium adolescentis		43.35	0.49			
Actinomyces naeslundii		3.79	0.46			
47757110	**Low discrimination**					**47757110**
Actinomyces israelii		82.14	0.73	MLZ 13		
Bifidobacterium adolescentis		8.92	0.50			
Bifidobacterium breve		3.58	0.44			
47757111	**Low discrimination**					**47757111**
Actinomyces israelii		81.63	0.39	MLZ 13		
Bifidobacterium adolescentis		8.87	0.16	URE 1		
Bifidobacterium breve		3.56	0.10			
47757112	**Low discrimination**					**47757112**
Bifidobacterium adolescentis		83.62	0.84	ARA 87		
Actinomyces israelii		11.66	0.76			
Bifidobacterium breve		3.42	0.62			
47757113	**Low discrimination**					**47757113**
Bifidobacterium adolescentis		83.43	0.50	ARA 87		
Actinomyces israelii		11.64	0.42	URE 1		
Bifidobacterium breve		3.42	0.27			
47757120	**Low discrimination**					**47757120**
					SPO	
Actinomyces israelii		62.14	0.67	NAG 6	1	
Bifidobacterium breve		11.26	0.49		1	
Clostridium tertium		7.98	0.37		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47757121	**Low discrimination**					**47757121**
Actinomyces naeslundii		54.45	0.46	ESL 12		
Actinomyces israelii		30.25	0.33	XYL 1		
Bifidobacterium breve		5.48	0.15			
47757122	**Low discrimination**					**47757122**
					GRA	
Bifidobacterium breve		32.11	0.66	XYL 1	99	
Actinomyces israelii		26.29	0.70		99	
Prevotella oralis		17.77	0.50		1	
47757130	**Low discrimination**					**47757130**
					SPO	
Clostridium tertium		38.60	0.41	RAF 1	99	
Bifidobacterium breve		30.65	0.49	TRE 90	1	
Actinomyces israelii		25.28	0.53		1	
47757132	**Low discrimination**					**47757132**
					SPO	
Bifidobacterium breve		55.74	0.66	XYL 1	1	
Clostridium tertium		20.82	0.50		99	
Bifidobacterium adolescentis		7.74	0.50		1	
47757140	**Very good identification**					**47757140**
					SPO	
Actinomyces israelii		95.66	0.80		1	
Clostridium butyricum		1.61	0.29		99	
Bifidobacterium breve		1.60	0.44		1	
47757141	**Acceptable identification**					**47757141**
					SPO	
Actinomyces israelii		92.11	0.46	URE 1	1	
Actinomyces naeslundii		4.80	0.33		1	
Clostridium butyricum		1.55	-.05		99	
47757142	**Low discrimination**					**47757142**
Actinomyces israelii		71.60	0.83			
Bifidobacterium adolescentis		19.94	0.67			
Bifidobacterium breve		8.07	0.62			
47757143	**Low discrimination**					**47757143**
Actinomyces israelii		70.12	0.49	URE 1		
Bifidobacterium adolescentis		19.53	0.33			
Bifidobacterium breve		7.91	0.27			
47757150	**Low discrimination**					**47757150**
					COC	
Actinomyces israelii		77.06	0.66	MLZ 13	1	
Blautia producta		11.13	0.32		99	
Bifidobacterium breve		8.63	0.44		1	
47757152	**Low discrimination**					**47757152**
Bifidobacterium adolescentis		56.28	0.69	ARA 87		
Actinomyces israelii		24.71	0.69	SOR 11		
Bifidobacterium breve		18.64	0.62			
47757160	**Low discrimination**					**47757160**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		65.99	0.60	NAG 6	1		
Bifidobacterium breve		30.74	0.49		1		
Clostridium butyricum		1.52	0.11		99		
47757162	**Low discrimination**						**47757162**
Bifidobacterium breve		74.23	0.66	XYL 1			
Actinomyces israelii		23.64	0.63				
Bifidobacterium adolescentis		1.04	0.33				
47757170	**Low discrimination**						**47757170**
					COC		
Bifidobacterium breve		70.72	0.49	XYL 1	1		
Actinomyces israelii		22.68	0.46	BGL 91	1		
Blautia producta		5.70	0.15		99		
47757172	**Good identification**						**47757172**
Bifidobacterium breve		93.40	0.66	XYL 1			
Actinomyces israelii		4.44	0.49				
Bifidobacterium adolescentis		1.60	0.35				
47757200	**Very good identification**						**47757200**
					SPO		
Actinomyces israelii		99.49	0.58	RHA 7	1		
Clostridium butyricum		0.51	-.03		99		
47757202	**Very good identification**						**47757202**
Actinomyces israelii		98.80	0.61	RHA 7			
Bifidobacterium adolescentis		0.47	0.15				
Clostridium clostridioforme		0.41	0.10				
47757210	**Good identification**						**47757210**
Actinomyces israelii		100.0	0.44	RHA 7			
47757212	**Low discrimination**						**47757212**
Actinomyces israelii		85.93	0.47	RHA 7			
Clostridium clostridioforme		10.73	0.22	MLZ 13			
Bifidobacterium adolescentis		3.34	0.16				
47757220	**Low discrimination**						**47757220**
					SPO	GRA	
Actinomyces israelii		87.41	0.37	RHA 7	1	99	
Prevotella oralis		6.78	0.02	NAG 6	1	1	
Clostridium ramosum		5.81	0.01		92	50	
47757222	**Low discrimination**						**47757222**
					SPO	GRA	
Actinomyces israelii		43.73	0.40	RHA 7	1	99	
Prevotella oralis		42.96	0.24	NAG 6	1	1	
Clostridium ramosum		7.76	0.11		92	50	
47757240	**Good identification**						**47757240**
Actinomyces israelii		100.0	0.51	RHA 7			
47757242	**Good identification**						**47757242**
Actinomyces israelii		100.0	0.54	RHA 7			
47757250	**Acceptable identification**						**47757250**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces israelii		92.81	0.36	RHA 7	1		
Blautia producta		7.19	-.03	MLZ 13	99		
47757252	**Good identification**						**47757252**
Actinomyces israelii		100.0	0.40	RHA 7			
47757300	**Good identification**						**47757300**
					SPO		
Actinomyces israelii		94.67	0.68	RHA 7	1		
Clostridium butyricum		2.93	0.21		99		
Eubacterium contortum		1.95	0.24		1		
47757302	**Low discrimination**						**47757302**
					GRA		
Actinomyces israelii		86.59	0.71	RHA 7	99		
Bifidobacterium adolescentis		10.18	0.49		99		
Prevotella oralis		1.12	0.22		1		
47757303	**Low discrimination**						**47757303**
Actinomyces israelii		89.48	0.37	RHA 7			
Bifidobacterium adolescentis		10.52	0.15	URE 1			
47757310	**Good identification**						**47757310**
					SPO		
Actinomyces israelii		95.86	0.54	RHA 7	1		
Bifidobacterium adolescentis		1.40	0.16	MLZ 13	1		
Clostridium butyricum		1.05	-.01		99		
47757312	**Low discrimination**						**47757312**
Actinomyces israelii		47.03	0.57	RHA 7			
Bifidobacterium adolescentis		45.25	0.50	MLZ 13			
Clostridium clostridioforme		4.92	0.31				
47757320	**Low discrimination**						**47757320**
					SPO	GRA	
Clostridium ramosum		49.20	0.35	TRE 94	92	50	
Actinomyces israelii		29.90	0.48	XYL 1	1	99	
Prevotella oralis		9.08	0.22		1	1	
47757322	**Low discrimination**						**47757322**
					SPO		
Clostridium ramosum		37.38	0.45	TRE 94	92		
Prevotella oralis		32.77	0.44	XYL 1	1		
Parabacteroides distasonis		13.23	0.50		1		
47757332	**Low discrimination**						**47757332**
Parabacteroides distasonis		46.04	0.45	MAN 1			
Clostridium clostridioforme		13.47	0.31				
Bacteroides ovatus		12.87	0.20				
47757340	**Good identification**						**47757340**
					COC	SPO	
Actinomyces israelii		90.52	0.61	RHA 7	1	1	
Blautia producta		8.65	0.24		99	1	
Clostridium butyricum		0.63	0.03		1	99	
47757342	**Good identification**						**47757342**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		94.22	0.64	RHA 7		
Bifidobacterium adolescentis		3.52	0.33			
Bifidobacterium breve		1.43	0.27			
47757352	**Low discrimination**					**47757352**
					COC	
Actinomyces israelii		64.55	0.50	RHA 7	1	
Bifidobacterium adolescentis		19.74	0.35	MLZ 13	1	
Blautia producta		9.17	0.15		99	
47757360	**Low discrimination**					**47757360**
					COC	
Actinomyces israelii		78.64	0.40	RHA 7	1	
Blautia producta		13.08	0.07	NAG 6	99	
Bifidobacterium breve		4.92	0.15		1	
47757362	**Low discrimination**					**47757362**
					SPO	
Actinomyces israelii		58.37	0.43	RHA 7	1	
Bifidobacterium breve		24.60	0.32	NAG 6	1	
Clostridium ramosum		6.65	0.11		92	
47757400	**Low discrimination**					**47757400**
					SPO	
Actinomyces israelii		87.52	0.75		1	
Clostridium butyricum		12.10	0.40		99	
Bifidobacterium adolescentis		0.38	0.29		1	
47757401	**Low discrimination**					**47757401**
					SPO	
Actinomyces israelii		87.86	0.41	URE 1	1	
Clostridium butyricum		12.14	0.05		99	
47757402	**Low discrimination**					**47757402**
					SPO	
Actinomyces israelii		75.38	0.78		1	
Bifidobacterium adolescentis		21.83	0.63		1	
Clostridium butyricum		2.70	0.33		99	
47757403	**Low discrimination**					**47757403**
					SPO	
Actinomyces israelii		75.44	0.44	URE 1	1	
Bifidobacterium adolescentis		21.85	0.29		1	
Clostridium butyricum		2.70	-.02		99	
47757410	**Low discrimination**					**47757410**
					SPO	
Actinomyces israelii		89.70	0.61	MLZ 13	1	
Clostridium butyricum		4.37	0.18		99	
Bifidobacterium adolescentis		3.22	0.30		1	
47757412	**Low discrimination**					**47757412**
Bifidobacterium adolescentis		65.96	0.64	LAC 99		
Actinomyces israelii		27.84	0.64			
Bifidobacterium dentium		5.17	0.27			
47757420	**Low discrimination**					**47757420**
					SPO	
Actinomyces israelii		84.15	0.55	NAG 6	1	
Clostridium butyricum		15.85	0.21		99	
47757422	**Low discrimination**					**47757422**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		88.24	0.58	NAG 6	1		
Clostridium butyricum		4.31	0.14		99		
Bifidobacterium adolescentis		4.04	0.29		1		
47757430	**Low discrimination**						**47757430**
					SPO		
Actinomyces israelii		54.55	0.41	MLZ 13	1		
Bifidobacterium dentium		41.83	0.15	NAG 6	1		
Clostridium butyricum		3.62	-.01		99		
47757440	**Good identification**						**47757440**
					COC	SPO	
Actinomyces israelii		93.68	0.68		1	1	
Blautia producta		3.29	0.24		99	1	
Clostridium butyricum		2.90	0.21		1	99	
47757442	**Good identification**						**47757442**
					SPO		
Actinomyces israelii		90.69	0.71		1		
Bifidobacterium adolescentis		8.35	0.47		1		
Clostridium butyricum		0.73	0.14		99		
47757443	**Acceptable identification**						**47757443**
Actinomyces israelii		91.57	0.37	URE 1			
Bifidobacterium adolescentis		8.43	0.13				
47757450	**Low discrimination**						**47757450**
					COC		
Actinomyces israelii		57.87	0.54	MLZ 13	1		
Blautia producta		40.84	0.32		99		
Bifidobacterium adolescentis		0.66	0.15		1		
47757452	**Low discrimination**						**47757452**
					COC		
Actinomyces israelii		55.16	0.57	MLZ 13	1		
Bifidobacterium adolescentis		41.53	0.49		1		
Blautia producta		2.88	0.15		99		
47757460	**Acceptable identification**						**47757460**
					COC	SPO	
Actinomyces israelii		90.64	0.48	NAG 6	1	1	
Blautia producta		5.55	0.07		99	1	
Clostridium butyricum		3.82	0.03		1	99	
47757462	**Good identification**						**47757462**
					SPO		
Actinomyces israelii		97.52	0.51	NAG 6	1		
Bifidobacterium adolescentis		1.42	0.13		1		
Clostridium butyricum		1.07	-.04		99		
47757472	**Low discrimination**						**47757472**
Actinomyces israelii		74.38	0.37	MLZ 13			
Bifidobacterium dentium		9.99	-.03	NAG 6			
Bifidobacterium adolescentis		8.86	0.15				
47757500	**Low discrimination**						**47757500**
					SPO		
Actinomyces israelii		51.43	0.86		1		
Clostridium butyricum		42.65	0.63		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		5.59	0.63		1		
47757501	**Low discrimination**						**47757501**
					SPO		
Actinomyces israelii		51.00	0.52	URE 1	1		
Clostridium butyricum		42.30	0.29		99		
Bifidobacterium adolescentis		5.54	0.29		1		
47757502	**Low discrimination**						**47757502**
					SPO		
Bifidobacterium adolescentis		85.42	0.97		1		
Actinomyces israelii		11.92	0.89		1		
Clostridium butyricum		2.56	0.56		99		
47757503	**Low discrimination**						**47757503**
					SPO		
Bifidobacterium adolescentis		85.44	0.63	URE 1	1		
Actinomyces israelii		11.92	0.55		1		
Clostridium butyricum		2.56	0.22		99		
47757510	**Low discrimination**						**47757510**
Actinomyces israelii		34.06	0.72	MLZ 13			
Bifidobacterium adolescentis		30.26	0.64				
Bifidobacterium dentium		23.21	0.44				
47757511	**Low discrimination**						**47757511**
Actinomyces israelii		33.54	0.37	MLZ 13			
Bifidobacterium adolescentis		29.80	0.30	URE 1			
Bifidobacterium dentium		22.86	0.10				
47757512	**Good identification**						**47757512**
Bifidobacterium adolescentis		91.14	0.99				
Bifidobacterium dentium		7.14	0.62				
Actinomyces israelii		1.55	0.75				
47757513	**Good identification**						**47757513**
Bifidobacterium adolescentis		91.18	0.64	URE 1			
Bifidobacterium dentium		7.14	0.27				
Actinomyces israelii		1.56	0.40				
47757520	**Low discrimination**						**47757520**
					GRA	SPO	
Clostridium butyricum		50.68	0.45	TRE 94	99	99	
Actinomyces israelii		44.85	0.65	MAN 10	99	1	
Bacteroides vulgatus		1.41	0.17	NAG 8	1	1	
47757522	**Low discrimination**						**47757522**
					SPO		
Bifidobacterium adolescentis		38.19	0.63	NAG 1	1		
Actinomyces israelii		33.67	0.68		1		
Clostridium butyricum		9.86	0.38		99		
47757530	**Acceptable identification**						**47757530**
					SPO		
Bifidobacterium dentium		91.74	0.49	TRE 99	1		
Actinomyces israelii		4.83	0.51	BGL 91	1		
Clostridium butyricum		1.92	0.23		99		
47757532	**Good identification**						**47757532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		91.94	0.66	TRE 99			
Bifidobacterium adolescentis		6.67	0.64				
Actinomyces israelii		0.72	0.54				
47757540	**Low discrimination**						**47757540**
					COC	SPO	
Actinomyces israelii		47.72	0.79		1	1	
Blautia producta		41.52	0.58		99	1	
Clostridium butyricum		8.85	0.45		1	99	
47757541	**Low discrimination**						**47757541**
					COC	SPO	
Actinomyces israelii		46.54	0.45	URE 1	1	1	
Blautia producta		40.50	0.24		99	1	
Clostridium butyricum		8.63	0.11		1	99	
47757542	**Low discrimination**						**47757542**
					COC		
Bifidobacterium adolescentis		67.15	0.82	SOR 11	1		
Actinomyces israelii		29.47	0.82		1		
Blautia producta		1.90	0.41		99		
47757543	**Low discrimination**						**47757543**
					COC		
Bifidobacterium adolescentis		67.19	0.47	SOR 11	1		
Actinomyces israelii		29.49	0.48	URE 1	1		
Blautia producta		1.90	0.07		99		
47757550	**Good identification**						**47757550**
					COC		
Blautia producta		92.69	0.66	TRE 99	99		
Actinomyces israelii		5.31	0.65		1		
Bifidobacterium adolescentis		1.50	0.49		1		
47757552	**Low discrimination**						**47757552**
					COC		
Bifidobacterium adolescentis		88.42	0.83	SOR 11	1		
Blautia producta		6.14	0.50		99		
Actinomyces israelii		4.75	0.68		1		
47757553	**Low discrimination**						**47757553**
					COC		
Bifidobacterium adolescentis		88.53	0.49	SOR 11	1		
Blautia producta		6.15	0.15	URE 1	99		
Actinomyces israelii		4.75	0.33		1		
47757560	**Low discrimination**						**47757560**
					COC	SPO	
Blautia producta		54.11	0.41	TRE 99	99	1	
Actinomyces israelii		35.72	0.58	NAG 10	1	1	
Clostridium butyricum		9.03	0.27		1	99	
47757562	**Low discrimination**						**47757562**
					COC		
Actinomyces israelii		62.75	0.61	NAG 6	1		
Bifidobacterium adolescentis		22.62	0.47		1		
Blautia producta		7.04	0.25		99		
47757570	**Acceptable identification**						**47757570**
					COC		
Blautia producta		94.67	0.50	TRE 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		3.11	0.44	NAG 10	1		
Bifidobacterium dentium		1.54	0.15		1		
47757600	**Good identification**						**47757600**
					SPO		
Actinomyces israelii		94.63	0.56	RHA 7	1		
Clostridium butyricum		5.37	0.14		99		
47757602	**Low discrimination**						**47757602**
Actinomyces israelii		88.34	0.59	RHA 7			
Clostridium clostridioforme		6.93	0.31				
Bifidobacterium adolescentis		3.43	0.29				
47757610	**Acceptable identification**						**47757610**
					COC		
Actinomyces israelii		93.19	0.42	RHA 7	1		
Blautia producta		3.43	-.03	MLZ 13	99		
Clostridium clostridioforme		3.38	0.08		1		
47757612	**Low discrimination**						**47757612**
Clostridium clostridioforme		64.04	0.42	TRE 88			
Actinomyces israelii		26.78	0.45	MAN 1			
Bifidobacterium adolescentis		8.51	0.30				
47757620	**Low discrimination**						**47757620**
					GRA	SPO	
Actinomyces israelii		81.36	0.36	RHA 7	99	1	
Bacteroides vulgatus		12.34	-.01	NAG 6	1	1	
Clostridium butyricum		6.30	-.04		99	99	
47757632	**Low discrimination**						**47757632**
					GRA		
Clostridium clostridioforme		89.94	0.42	TRE 88	50		
Bacteroides ovatus		5.99	0.12	MAN 1	1		
Actinomyces israelii		2.40	0.25		99		
47757640	**Low discrimination**						**47757640**
					COC	SPO	
Actinomyces israelii		67.58	0.49	RHA 7	1	1	
Blautia producta		31.56	0.24		99	1	
Clostridium butyricum		0.86	-.04		1	99	
47757642	**Good identification**						**47757642**
					COC		
Actinomyces israelii		95.52	0.52	RHA 7	1		
Blautia producta		3.31	0.07		99		
Bifidobacterium adolescentis		1.18	0.13		1		
47757652	**Low discrimination**						**47757652**
					COC		
Actinomyces israelii		53.83	0.38	RHA 7	1		
Blautia producta		37.39	0.15	MLZ 13	99		
Bifidobacterium adolescentis		5.44	0.15		1		
47757700	**Low discrimination**						**47757700**
					GRA	SPO	
Actinomyces israelii		67.76	0.66	RHA 7	99	1	
Clostridium butyricum		23.09	0.37		99	99	
Mitsuokella multacida		3.46	0.32		1	1	
47757702	**Low discrimination**						**47757702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces israelii		46.75	0.69	RHA 7	1		
Bifidobacterium adolescentis		44.98	0.63		1		
Clostridium butyricum		4.13	0.30		99		
47757703	**Low discrimination**						**47757703**
					SPO		
Actinomyces israelii		47.26	0.35	RHA 7	1		
Bifidobacterium adolescentis		45.46	0.29	URE 1	1		
Clostridium butyricum		4.18	-.04		99		
47757710	**Low discrimination**						**47757710**
					COC	SPO	
Actinomyces israelii		41.90	0.52	RHA 7	1	1	
Blautia producta		38.20	0.32	MLZ 13	99	1	
Clostridium butyricum		5.03	0.16		1	99	
47757712	**Low discrimination**						**47757712**
Bifidobacterium adolescentis		67.64	0.64	RHA 1			
Clostridium clostridioforme		17.21	0.51				
Actinomyces israelii		8.60	0.55				
47757722	**Low discrimination**						**47757722**
Bacteroides ovatus		64.11	0.49	IND 97			
Bacteroides uniformis		18.99	0.37	TRE 91			
Parabacteroides distasonis		5.38	0.46				
47757732	**Low discrimination**						**47757732**
					GRA		
Bacteroides ovatus		52.75	0.46	IND 97	1		
Clostridium clostridioforme		26.77	0.51	TRE 91	50		
Bifidobacterium dentium		14.66	0.32		99		
47757740	**Good identification**						**47757740**
					COC	GRA	
Blautia producta		90.51	0.58	TRE 99	99	99	
Actinomyces israelii		7.83	0.59		1	99	
Mitsuokella multacida		1.03	0.32		1	1	
47757742	**Low discrimination**						**47757742**
					COC		
Actinomyces israelii		45.60	0.62	RHA 7	1		
Blautia producta		39.04	0.41		99		
Bifidobacterium adolescentis		13.94	0.47		1		
47757750	**Very good identification**						**47757750**
					COC	GRA	
Blautia producta		99.42	0.66	TRE 99	99	99	
Actinomyces israelii		0.43	0.45		1	99	
Mitsuokella multacida		0.13	0.24		1	1	
47757752	**Low discrimination**						**47757752**
					COC		
Blautia producta		82.72	0.50	TRE 99	99		
Bifidobacterium adolescentis		12.03	0.49	BGL 10	1		
Actinomyces israelii		4.81	0.48		1		
47757760	**Acceptable identification**						**47757760**
					COC	GRA	
Blautia producta		92.14	0.41	TRE 99	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		4.58	0.39	NAG 10	1	99	
Mitsuokella multacida		2.36	0.21		1	1	
47757770	**Good identification**						**47757770**
					COC	GRA	
Blautia producta		99.47	0.50	TRE 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
Actinomyces israelii		0.25	0.25		1	99	
47760712	**Good identification**						**47760712**
Clostridium clostridioforme		100.0	0.41	FRU 99			
47760732	**Good identification**						**47760732**
Clostridium clostridioforme		100.0	0.41	FRU 99			
47761001	**Good identification**						**47761001**
Actinomyces naeslundii		100.0	0.51	ESL 12			
47761003	**Good identification**						**47761003**
Actinomyces naeslundii		100.0	0.51	ESL 12			
47761020	**Low discrimination**						**47761020**
					SPO		
Actinomyces naeslundii		49.98	0.48	ESL 12	1		
Lactobacillus jensenii		26.69	0.21	FRU 95	1		
Clostridium paraputrificum		8.67	0.03		99		
47761021	**Good identification**						**47761021**
Actinomyces naeslundii		100.0	0.51	ESL 12			
47761023	**Good identification**						**47761023**
Actinomyces naeslundii		100.0	0.51	ESL 12			
47761041	**Good identification**						**47761041**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47761043	**Good identification**						**47761043**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47761061	**Good identification**						**47761061**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47761063	**Good identification**						**47761063**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47761100	**Low discrimination**						**47761100**
					SPO		
Bifidobacterium breve		26.40	0.41	FRU 99	1		
Actinomyces naeslundii		24.53	0.55	BGL 91	1		
Clostridium butyricum		20.22	0.24		99		
47761101	**Very good identification**						**47761101**
Actinomyces naeslundii		100.0	0.58	ESL 12			
47761102	**Low discrimination**						**47761102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium breve		45.42	0.59	FRU 99	99		
Lactobacillus acidophilus		37.43	0.47		99		
Leptotrichia buccalis		5.74	0.24		1		
47761103	**Low discrimination**						**47761103**
Actinomyces naeslundii		87.50	0.58	ESL 12			
Bifidobacterium breve		6.41	0.24	FRU 95			
Lactobacillus acidophilus		5.28	0.13				
47761111	**Low discrimination**						**47761111**
					GRA		
Actinomyces naeslundii		85.05	0.37	ESL 12	99		
Leptotrichia buccalis		14.95	-.11	FRU 95	1		
47761112	**Good identification**						**47761112**
					GRA		
Leptotrichia buccalis		92.14	0.58	FRU 99	1		
Bifidobacterium breve		7.37	0.59		99		
Bifidobacterium adolescentis		0.33	0.35		99		
47761120	**Low discrimination**						**47761120**
					SPO		
Clostridium paraputrificum		43.57	0.37	RAF 7	99		
Clostridium ramosum		29.55	0.41	FRU 99	92		
Bifidobacterium breve		13.82	0.46		1		
47761121	**Good identification**						**47761121**
					SPO		
Actinomyces naeslundii		92.50	0.58	ESL 12	1		
Clostridium paraputrificum		3.76	0.03	FRU 95	99		
Clostridium ramosum		2.55	0.06		92		
47761122	**Low discrimination**						**47761122**
					SPO		
Bifidobacterium breve		33.70	0.63	FRU 99	1		
Clostridium ramosum		28.52	0.51		92		
Lactobacillus acidophilus		15.62	0.47		1		
47761123	**Low discrimination**						**47761123**
					SPO		
Actinomyces naeslundii		72.78	0.58	ESL 12	1		
Bifidobacterium breve		9.48	0.29	FRU 95	1		
Clostridium ramosum		8.02	0.17		92		
47761130	**Low discrimination**						**47761130**
					GRA	SPO	
Bifidobacterium breve		86.15	0.46	FRU 99	99	1	
Leptotrichia buccalis		6.87	0.07	BGL 91	1	1	
Clostridium paraputrificum		2.74	0.03		99	99	
47761131	**Low discrimination**						**47761131**
Actinomyces naeslundii		80.34	0.37	ESL 12			
Bifidobacterium breve		19.66	0.12	FRU 95			
47761132	**Low discrimination**						**47761132**
					GRA		
Bifidobacterium breve		55.00	0.63	FRU 99	99		
Leptotrichia buccalis		42.97	0.41		1		
Parabacteroides distasonis		0.72	0.32		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47761140	**Low discrimination**					**47761140**
					SPO	
Bifidobacterium breve		77.95	0.41	FRU 99	1	
Actinomyces naeslundii		12.79	0.43	BGL 91	1	
Clostridium butyricum		5.19	0.06		99	
47761141	**Good identification**					**47761141**
Actinomyces naeslundii		100.0	0.46	ESL 12		
47761142	**Low discrimination**					**47761142**
Bifidobacterium breve		88.17	0.59	FRU 99		
Lactobacillus acidophilus		8.98	0.32			
Actinomyces naeslundii		1.43	0.43			
47761143	**Low discrimination**					**47761143**
Actinomyces naeslundii		70.67	0.46	ESL 12		
Bifidobacterium breve		29.34	0.24	FRU 95		
47761150	**Good identification**					**47761150**
					GRA	
Bifidobacterium breve		98.73	0.41	FRU 99	99	
Leptotrichia buccalis		1.27	-.11	BGL 91	1	
47761152	**Low discrimination**					**47761152**
					GRA	
Bifidobacterium breve		88.26	0.59	FRU 99	99	
Leptotrichia buccalis		11.15	0.24		1	
Bifidobacterium adolescentis		0.49	0.19		99	
47761160	**Low discrimination**					**47761160**
					SPO	
Bifidobacterium breve		74.01	0.46	FRU 99	1	
Clostridium paraputrificum		17.57	0.18	BGL 91	99	
Actinomyces naeslundii		6.83	0.43		1	
47761161	**Acceptable identification**					**47761161**
Actinomyces naeslundii		93.20	0.46	ESL 12		
Bifidobacterium breve		6.80	0.12	FRU 95		
47761162	**Good identification**					**47761162**
					SPO	
Bifidobacterium breve		91.80	0.63	FRU 99	1	
Lactobacillus acidophilus		5.26	0.32		1	
Clostridium paraputrificum		0.92	0.11		99	
47761163	**Low discrimination**					**47761163**
Actinomyces naeslundii		57.54	0.46	ESL 12		
Bifidobacterium breve		42.46	0.29	FRU 95		
47761170	**Good identification**					**47761170**
Bifidobacterium breve		100.0	0.46	FRU 99		
47761172	**Very good identification**					**47761172**
					GRA	
Bifidobacterium breve		99.22	0.63	FRU 99	99	
Leptotrichia buccalis		0.78	0.07		1	
47761320	**Good identification**					**47761320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO	GRA	
Clostridium ramosum		95.07	0.43	FRU 99	92	50	
Prevotella oralis		3.28	0.17		1	1	
Clostridium paraputrificum		1.07	0.03		99	99	
47761322	**Low discrimination**						**47761322**
					SPO		
Clostridium ramosum		78.55	0.53	FRU 99	92		
Prevotella oralis		12.86	0.39		1		
Parabacteroides distasonis		5.09	0.45		1		
47761332	**Low discrimination**						**47761332**
					SPO		
Parabacteroides distasonis		45.12	0.40	FRU 99	1		
Clostridium clostridioforme		18.92	0.28	XYL 90	66		
Clostridium ramosum		13.07	0.19		92		
47761500	**Good identification**						**47761500**
					SPO		
Clostridium butyricum		98.08	0.40	FRU 99	99		
Actinomyces israelii		1.57	0.30	XYL 97	1		
Bifidobacterium adolescentis		0.36	0.13		1		
47761512	**Low discrimination**						**47761512**
					GRA		
Bifidobacterium adolescentis		58.15	0.49	FRU 99	99		
Leptotrichia buccalis		24.35	0.24	XYL 87	1		
Clostridium clostridioforme		12.02	0.34		50		
47761612	**Good identification**						**47761612**
Clostridium clostridioforme		100.0	0.40	FRU 99			
47761632	**Good identification**						**47761632**
Clostridium clostridioforme		100.0	0.40	FRU 99			
47761702	**Low discrimination**						**47761702**
					SPO		
Clostridium clostridioforme		89.38	0.37	FRU 99	66		
Clostridium butyricum		7.31	0.07	XYL 91	99		
Bifidobacterium adolescentis		2.22	0.13		1		
47761712	**Good identification**						**47761712**
					GRA		
Clostridium clostridioforme		99.06	0.49	FRU 99	50		
Bifidobacterium adolescentis		0.66	0.15	XYL 91	99		
Leptotrichia buccalis		0.28	-.11		1		
47761722	**Low discrimination**						**47761722**
					SPO		
Clostridium clostridioforme		47.75	0.37	FRU 99	66		
Parabacteroides distasonis		37.69	0.41	XYL 91	1		
Clostridium ramosum		9.59	0.19		92		
47761732	**Acceptable identification**						**47761732**
Clostridium clostridioforme		90.77	0.49	FRU 99			
Parabacteroides distasonis		8.47	0.37	XYL 91			
Bacteroides ovatus		0.76	0.03				
47762100	**Good identification**						**47762100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ESL 12		
47762101	**Good identification**					**47762101**
Actinomyces naeslundii		100.0	0.40	ESL 12		
47762103	**Good identification**					**47762103**
Actinomyces naeslundii		100.0	0.40	ESL 12		
47762121	**Good identification**					**47762121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
47762123	**Good identification**					**47762123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
47763000	**Low discrimination**					**47763000**
Actinomyces naeslundii		76.90	0.53	ESL 12		
Actinomyces israelii		23.11	0.34	FRU 95		
47763001	**Good identification**					**47763001**
Actinomyces naeslundii		100.0	0.56	ESL 12		
47763002	**Low discrimination**					**47763002**
Actinomyces naeslundii		59.81	0.53	ESL 12		
Actinomyces israelii		26.96	0.37	FRU 95		
Bifidobacterium breve		9.64	0.24			
47763003	**Good identification**					**47763003**
Actinomyces naeslundii		100.0	0.56	ESL 12		
47763020	**Low discrimination**					**47763020**
					SPO	
Actinomyces naeslundii		71.90	0.53	ESL 12	1	
Clostridium ramosum		24.69	0.19	FRU 95	92	
Bifidobacterium breve		2.04	0.12		1	
47763021	**Good identification**					**47763021**
Actinomyces naeslundii		100.0	0.56	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47763023	**Good identification**					**47763023**
Actinomyces naeslundii		100.0	0.56	ESL 12		
47763040	**Low discrimination**					**47763040**
Actinomyces naeslundii		60.16	0.40	ESL 12		
Actinomyces israelii		39.84	0.27	FRU 95		
47763041	**Good identification**					**47763041**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47763042	**Low discrimination**					**47763042**
Actinomyces naeslundii		34.41	0.40	ESL 12		
Actinomyces israelii		34.18	0.30	FRU 95		
Bifidobacterium breve		31.42	0.24			
47763043	**Good identification**					**47763043**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47763060	**Low discrimination**					**47763060**
Actinomyces naeslundii		86.17	0.40	ESL 12		
Bifidobacterium breve		13.83	0.12	FRU 95		
47763061	**Good identification**					**47763061**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47763063	**Good identification**					**47763063**
Actinomyces naeslundii		100.0	0.43	ESL 12		
47763100	**Low discrimination**					**47763100**
Actinomyces naeslundii		46.14	0.60	ESL 12		
Bifidobacterium breve		25.57	0.41	FRU 95		
Actinomyces israelii		19.49	0.45			
47763101	**Very good identification**					**47763101**
Actinomyces naeslundii		100.0	0.63	ESL 12		
47763102	**Low discrimination**					**47763102**
Bifidobacterium breve		68.12	0.59	FRU 99		
Actinomyces naeslundii		12.16	0.60			
Actinomyces israelii		7.70	0.48			
47763103	**Very good identification**					**47763103**
Actinomyces naeslundii		95.97	0.63	ESL 12		
Bifidobacterium breve		3.62	0.24	FRU 95		
Actinomyces israelii		0.41	0.14			
47763110	**Low discrimination**					**47763110**
Bifidobacterium breve		77.20	0.41	FRU 99		
Actinomyces israelii		8.79	0.31	BGL 91		
Actinomyces naeslundii		7.33	0.38			
47763111	**Good identification**					**47763111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
47763112	**Low discrimination**					**47763112**
					GRA	
Bifidobacterium breve		82.42	0.59	FRU 99	99	
Leptotrichia buccalis		10.41	0.24		1	
Bifidobacterium adolescentis		3.04	0.33		99	
47763113	**Low discrimination**					**47763113**
					GRA	
Actinomyces naeslundii		55.32	0.41	ESL 12	99	
Bifidobacterium breve		39.67	0.24	FRU 95	99	
Leptotrichia buccalis		5.01	-.11	MLZ 5	1	
47763120	**Low discrimination**					**47763120**
					SPO	
Clostridium ramosum		84.97	0.53	FRU 99	92	
Actinomyces naeslundii		7.11	0.60		1	
Bifidobacterium breve		7.01	0.46		1	
47763121	**Good identification**					**47763121**
					SPO	
Actinomyces naeslundii		91.99	0.63	ESL 12	1	
Clostridium ramosum		7.40	0.19	FRU 95	92	
Bifidobacterium breve		0.61	0.12		1	
47763122	**Low discrimination**					**47763122**
					SPO	
Clostridium ramosum		80.59	0.64	FRU 99	92	
Bifidobacterium breve		16.81	0.63		1	
Actinomyces naeslundii		1.69	0.60		1	
47763123	**Low discrimination**					**47763123**
					SPO	
Actinomyces naeslundii		72.00	0.63	ESL 12	1	
Clostridium ramosum		23.17	0.30	FRU 95	92	
Bifidobacterium breve		4.83	0.29		1	
47763130	**Low discrimination**					**47763130**
					SPO	
Bifidobacterium breve		76.91	0.46	FRU 99	1	
Clostridium tertium		9.56	0.21	BGL 91	99	
Clostridium ramosum		9.42	0.19		92	
47763131	**Low discrimination**					**47763131**
Actinomyces naeslundii		88.81	0.41	ESL 12		
Bifidobacterium breve		11.19	0.12	FRU 95		
47763132	**Good identification**					**47763132**
					SPO	
Bifidobacterium breve		90.82	0.63	FRU 99	1	
Clostridium ramosum		4.40	0.30		92	
Clostridium tertium		3.35	0.29		99	
47763140	**Low discrimination**					**47763140**
Bifidobacterium breve		61.93	0.41	FRU 99		
Actinomyces naeslundii		19.72	0.47	BGL 91		
Actinomyces israelii		18.35	0.38			
47763141	**Good identification**					**47763141**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	ESL 12		
47763142	**Good identification**					**47763142**
Bifidobacterium breve		92.21	0.59	FRU 99		
Actinomyces israelii		4.05	0.41			
Actinomyces naeslundii		2.90	0.47			
47763143	**Low discrimination**					**47763143**
Actinomyces naeslundii		82.38	0.50	ESL 12		
Bifidobacterium breve		17.62	0.24	FRU 95		
47763150	**Acceptable identification**					**47763150**
					COC	
Bifidobacterium breve		92.54	0.41	FRU 99	1	
Actinomyces israelii		4.10	0.24	BGL 91	1	
Blautia producta		1.81	-.03		99	
47763152	**Very good identification**					**47763152**
Bifidobacterium breve		98.61	0.59	FRU 99		
Actinomyces israelii		0.65	0.27			
Bifidobacterium adolescentis		0.45	0.18			
47763160	**Low discrimination**					**47763160**
					SPO	
Bifidobacterium breve		76.21	0.46	FRU 99	1	
Actinomyces naeslundii		13.65	0.47	BGL 91	1	
Clostridium ramosum		9.33	0.19		92	
47763161	**Good identification**					**47763161**
Actinomyces naeslundii		96.38	0.50	ESL 12		
Bifidobacterium breve		3.62	0.12	FRU 95		
47763162	**Good identification**					**47763162**
					SPO	
Bifidobacterium breve		93.39	0.63	FRU 99	1	
Clostridium ramosum		4.52	0.30		92	
Actinomyces naeslundii		1.65	0.47		1	
47763163	**Low discrimination**					**47763163**
Actinomyces naeslundii		72.45	0.50	ESL 12		
Bifidobacterium breve		27.55	0.29	FRU 95		
47763170	**Good identification**					**47763170**
Bifidobacterium breve		99.07	0.46	FRU 99		
Actinomyces naeslundii		0.93	0.26	BGL 91		
47763172	**Very good identification**					**47763172**
Bifidobacterium breve		99.91	0.63	FRU 99		
Actinomyces naeslundii		0.09	0.26			
47763320	**Good identification**					**47763320**
					SPO	
Clostridium ramosum		99.87	0.56	FRU 99	92	
Actinomyces naeslundii		0.06	0.26		1	
Bifidobacterium breve		0.06	0.12		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47763322	**Very good identification**					**47763322**
					SPO	
Clostridium ramosum		99.76	0.66	FRU 99	92	
Bifidobacterium breve		0.16	0.29		1	
Lactobacillus acidophilus		0.03	0.06		1	
47763402	**Low discrimination**					**47763402**
					SPO	
Actinomyces israelii		87.62	0.36	FRU 99	1	
Bifidobacterium adolescentis		7.73	0.12	XYL 92	1	
Clostridium butyricum		4.65	-.07		99	
47763502	**Low discrimination**					**47763502**
					SPO	
Bifidobacterium adolescentis		59.79	0.46	FRU 99	1	
Actinomyces israelii		27.39	0.46	XYL 87	1	
Clostridium butyricum		8.72	0.17		99	
47763512	**Low discrimination**					**47763512**
Bifidobacterium adolescentis		52.77	0.47	FRU 99		
Bifidobacterium dentium		41.51	0.27	XYL 87		
Actinomyces israelii		2.96	0.32			
47763542	**Low discrimination**					**47763542**
Actinomyces israelii		44.59	0.39	FRU 99		
Bifidobacterium adolescentis		30.93	0.30	XYL 92		
Bifidobacterium breve		12.52	0.24			
47764312	**Good identification**					**47764312**
Clostridium clostridioforme		100.0	0.37	FRU 99		
47764332	**Acceptable identification**					**47764332**
Clostridium clostridioforme		93.96	0.37	FRU 99		
Parabacteroides distasonis		6.04	0.23	ARA 94		
47764512	**Low discrimination**					**47764512**
					SPO	
Clostridium clostridioforme		88.65	0.43	FRU 99	66	
Bifidobacterium adolescentis		8.60	0.29	RHA 88	1	
Clostridium butyricum		1.92	0.03		99	
47764532	**Good identification**					**47764532**
					GRA	
Clostridium clostridioforme		97.78	0.43	FRU 99	50	
Bifidobacterium dentium		1.62	-.03	RHA 88	99	
Parabacteroides distasonis		0.60	0.11		1	
47764602	**Good identification**					**47764602**
Clostridium clostridioforme		100.0	0.37	FRU 99		
47764612	**Good identification**					**47764612**
Clostridium clostridioforme		100.0	0.49	FRU 99		
47764622	**Good identification**					**47764622**
Clostridium clostridioforme		100.0	0.37	FRU 99		
47764632	**Good identification**					**47764632**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.49	FRU 99			
47764702	**Good identification**						**47764702**
Clostridium clostridioforme		99.21	0.46	FRU 99			
Clostridium butyricum		0.79	-.01				
47764712	**Very good identification**						**47764712**
Clostridium clostridioforme		100.0	0.58	FRU 99			
47764722	**Good identification**						**47764722**
Clostridium clostridioforme		96.31	0.46	FRU 99			
Parabacteroides distasonis		2.05	0.23				
Bacteroides ovatus		1.64	0.06				
47764732	**Very good identification**						**47764732**
Clostridium clostridioforme		99.50	0.58	FRU 99			
Parabacteroides distasonis		0.25	0.19				
Bacteroides ovatus		0.25	0.03				
47765002	**Low discrimination**						**47765002**
					SPO		
Actinomyces israelii		56.14	0.43	FRU 99	1		
Clostridium butyricum		38.84	0.19		99		
Clostridium clostridioforme		2.58	0.10		66		
47765042	**Low discrimination**						**47765042**
					SPO		
Actinomyces israelii		86.60	0.36	FRU 99	1		
Clostridium butyricum		13.40	0.01		99		
47765100	**Acceptable identification**						**47765100**
					COC	SPO	
Clostridium butyricum		91.04	0.50	FRU 99	1	99	
Actinomyces israelii		5.69	0.50	ARA 90	1	1	
Atopobium parvulum		3.13	0.32		99	1	
47765102	**Low discrimination**						**47765102**
					SPO		
Clostridium butyricum		64.32	0.43	FRU 99	99		
Bifidobacterium adolescentis		16.66	0.47	ARA 90	1		
Actinomyces israelii		15.49	0.53		1		
47765112	**Low discrimination**						**47765112**
					GRA	SPO	
Bifidobacterium adolescentis		53.22	0.49	FRU 99	99	1	
Leptotrichia buccalis		22.28	0.24	ARA 87	1	1	
Clostridium butyricum		8.85	0.21		99	99	
47765122	**Low discrimination**						**47765122**
					GRA	SPO	
Parabacteroides distasonis		60.52	0.53	FRU 99	1	1	
Prevotella oralis		14.71	0.30		1	1	
Clostridium butyricum		12.93	0.24		99	99	
47765132	**Low discrimination**						**47765132**
					GRA		
Parabacteroides distasonis		70.99	0.49	FRU 99	1		
Clostridium clostridioforme		13.68	0.31		50		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		6.11	0.29		99	
47765142	**Low discrimination**					**47765142**
					SPO	
Actinomyces israelii		41.96	0.46	FRU 99	1	
Clostridium butyricum		38.95	0.24		99	
Bifidobacterium adolescentis		14.34	0.32		1	
47765212	**Good identification**					**47765212**
Clostridium clostridioforme		100.0	0.36	FRU 99		
47765232	**Acceptable identification**					**47765232**
Clostridium clostridioforme		93.30	0.36	FRU 99		
Parabacteroides distasonis		6.70	0.23	ARA 94		
47765312	**Good identification**					**47765312**
					GRA	
Clostridium clostridioforme		95.23	0.45	FRU 99	50	
Parabacteroides distasonis		2.04	0.23	ARA 94	1	
Bifidobacterium adolescentis		0.98	0.15		99	
47765322	**Low discrimination**					**47765322**
Parabacteroides distasonis		88.81	0.61	FRU 99		
Clostridium clostridioforme		4.95	0.34			
Prevotella oralis		3.08	0.24			
47765332	**Low discrimination**					**47765332**
Parabacteroides distasonis		67.12	0.57	FRU 99		
Clostridium clostridioforme		31.62	0.45			
Bacteroides ovatus		1.26	0.11			
47765400	**Good identification**					**47765400**
					SPO	
Clostridium butyricum		96.71	0.42	FRU 99	99	
Actinomyces israelii		3.29	0.38	LAC 96	1	
47765402	**Low discrimination**					**47765402**
					SPO	
Clostridium butyricum		77.29	0.35	FRU 99	99	
Actinomyces israelii		10.16	0.41	LAC 96	1	
Clostridium clostridioforme		8.94	0.31		66	
47765412	**Low discrimination**					**47765412**
					SPO	
Clostridium clostridioforme		79.63	0.42	FRU 99	66	
Bifidobacterium adolescentis		8.63	0.29	RHA 88	1	
Clostridium butyricum		7.95	0.13		99	
47765432	**Good identification**					**47765432**
					SPO	
Clostridium clostridioforme		96.71	0.42	FRU 99	66	
Bifidobacterium dentium		1.79	-.03	RHA 88	1	
Clostridium butyricum		0.84	-.05		99	
47765500	**Very good identification**					**47765500**
					SPO	
Clostridium butyricum		99.28	0.66	FRU 99	99	
Actinomyces israelii		0.56	0.48		1	
Bifidobacterium adolescentis		0.08	0.27		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
47765502	**Low discrimination**						**47765502**
					SPO		
Clostridium butyricum		81.21	0.59	FRU 99	99		
Bifidobacterium adolescentis		15.65	0.61		1		
Actinomyces israelii		1.78	0.52		1		
47765510	**Acceptable identification**						**47765510**
					SPO		
Clostridium butyricum		94.15	0.44	FRU 99	99		
Bifidobacterium adolescentis		1.66	0.29	MLZ 5	1		
Bifidobacterium dentium		1.56	0.10		1		
47765512	**Low discrimination**						**47765512**
					SPO		
Bifidobacterium adolescentis		60.72	0.63	FRU 99	1		
Clostridium clostridioforme		18.96	0.51		66		
Clostridium butyricum		13.57	0.37		99		
47765520	**Good identification**						**47765520**
					GRA	SPO	
Clostridium butyricum		99.10	0.48	FRU 99	99	99	
Actinomyces israelii		0.41	0.28	NAG 8	99	1	
Mitsuokella multacida		0.21	0.11		1	1	
47765522	**Low discrimination**						**47765522**
					GRA	SPO	
Clostridium butyricum		62.64	0.41	FRU 99	99	99	
Parabacteroides distasonis		19.96	0.49	NAG 8	1	1	
Clostridium clostridioforme		11.63	0.40		50	66	
47765532	**Low discrimination**						**47765532**
					GRA		
Clostridium clostridioforme		53.20	0.51	FRU 99	50		
Bifidobacterium dentium		29.13	0.32	RHA 88	99		
Parabacteroides distasonis		10.80	0.45		1		
47765540	**Acceptable identification**						**47765540**
					COC	SPO	
Clostridium butyricum		93.60	0.48	FRU 99	1	99	
Blautia producta		3.12	0.24	SOR 8	99	1	
Actinomyces israelii		2.38	0.41		1	1	
47765542	**Low discrimination**						**47765542**
					SPO		
Clostridium butyricum		72.17	0.41	FRU 99	99		
Bifidobacterium adolescentis		19.76	0.46	SOR 8	1		
Actinomyces israelii		7.07	0.44		1		
47765552	**Low discrimination**						**47765552**
					COC	SPO	
Bifidobacterium adolescentis		75.78	0.47	FRU 99	1	1	
Clostridium butyricum		11.92	0.19	SOR 11	1	99	
Blautia producta		6.46	0.15		99	1	
47765602	**Good identification**						**47765602**
					SPO		
Clostridium clostridioforme		95.41	0.46	FRU 99	66		
Clostridium butyricum		3.48	0.09		99		
Actinomyces israelii		1.11	0.22		1		
47765612	**Very good identification**						**47765612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.57	FRU 99			
47765622	**Good identification**						**47765622**
Clostridium clostridioforme		95.90	0.46	FRU 99			
Parabacteroides distasonis		2.28	0.23				
Bacteroides ovatus		1.82	0.06				
47765632	**Very good identification**						**47765632**
Clostridium clostridioforme		99.44	0.57	FRU 99			
Parabacteroides distasonis		0.28	0.19				
Bacteroides ovatus		0.28	0.03				
47765700	**Acceptable identification**						**47765700**
					GRA	SPO	
Clostridium butyricum		93.11	0.40	FRU 99	99	99	
Mitsuokella multacida		4.20	0.25	RHA 3	1	1	
Clostridium clostridioforme		1.40	0.20		50	66	
47765702	**Low discrimination**						**47765702**
					SPO		
Clostridium clostridioforme		76.46	0.55	FRU 99	66		
Clostridium butyricum		19.98	0.33		99		
Bifidobacterium adolescentis		1.26	0.27		1		
47765712	**Very good identification**						**47765712**
					SPO		
Clostridium clostridioforme		98.99	0.66	FRU 99	66		
Bifidobacterium adolescentis		0.44	0.29		1		
Clostridium butyricum		0.30	0.11		99		
47765722	**Low discrimination**						**47765722**
Clostridium clostridioforme		42.99	0.55	FRU 99			
Parabacteroides distasonis		30.20	0.58				
Bacteroides ovatus		24.19	0.40				
47765732	**Low discrimination**						**47765732**
Clostridium clostridioforme		85.53	0.66	FRU 99			
Parabacteroides distasonis		7.10	0.53				
Bacteroides ovatus		7.04	0.37				
47767000	**Good identification**						**47767000**
					SPO		
Actinomyces israelii		95.03	0.53	FRU 99	1		
Clostridium butyricum		4.97	0.10		99		
47767002	**Good identification**						**47767002**
					SPO		
Actinomyces israelii		98.05	0.56	FRU 99	1		
Clostridium butyricum		1.33	0.03		99		
Bifidobacterium adolescentis		0.62	0.12		1		
47767010	**Good identification**						**47767010**
Actinomyces israelii		100.0	0.38	FRU 99			
47767012	**Acceptable identification**						**47767012**
Actinomyces israelii		91.60	0.41	FRU 99			
Bifidobacterium adolescentis		4.70	0.13	MLZ 13			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		3.70	-.07				
47767022	**Good identification**						**47767022**
Actinomyces israelii		100.0	0.35	FRU 99			
47767040	**Good identification**						**47767040**
Actinomyces israelii		100.0	0.46	FRU 99			
47767042	**Good identification**						**47767042**
Actinomyces israelii		100.0	0.49	FRU 99			
47767100	**Low discrimination**						**47767100**
					SPO		
Actinomyces israelii		75.47	0.63	FRU 99	1		
Clostridium butyricum		23.70	0.33		99		
Clostridium tertium		0.49	0.11		99		
47767102	**Low discrimination**						**47767102**
					SPO		
Actinomyces israelii		79.34	0.66	FRU 99	1		
Bifidobacterium adolescentis		12.32	0.46		1		
Clostridium butyricum		6.46	0.26		99		
47767110	**Low discrimination**						**47767110**
					SPO		
Actinomyces israelii		72.96	0.49	FRU 99	1		
Bifidobacterium dentium		10.81	0.10	MLZ 13	1		
Clostridium butyricum		8.07	0.11		99		
47767112	**Low discrimination**						**47767112**
Bifidobacterium adolescentis		36.17	0.47	FRU 99			
Actinomyces israelii		28.47	0.52	ARA 87			
Bifidobacterium dentium		28.45	0.27				
47767120	**Low discrimination**						**47767120**
					SPO		
Actinomyces israelii		45.69	0.42	FRU 99	1		
Clostridium butyricum		19.54	0.15	NAG 6	99		
Clostridium ramosum		17.78	0.19		92		
47767140	**Low discrimination**						**47767140**
					COC	SPO	
Actinomyces israelii		79.40	0.56	FRU 99	1	1	
Blautia producta		15.03	0.24		99	1	
Clostridium butyricum		5.58	0.15		1	99	
47767142	**Good identification**						**47767142**
Actinomyces israelii		90.78	0.59	FRU 99			
Bifidobacterium adolescentis		4.48	0.30				
Bifidobacterium breve		1.81	0.24				
47767152	**Low discrimination**						**47767152**
					COC		
Actinomyces israelii		54.20	0.45	FRU 99	1		
Bifidobacterium adolescentis		21.89	0.32	MLZ 13	1		
Blautia producta		15.26	0.15		99		
47767160	**Low discrimination**						**47767160**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		66.33	0.35	FRU 99	1	1	
Blautia producta		21.85	0.07	NAG 6	99	1	
Clostridium butyricum		6.34	-.03		1	99	
47767162	**Low discrimination**						**47767162**
Actinomyces israelii		64.24	0.38	FRU 99			
Bifidobacterium breve		35.76	0.29	NAG 6			
47767202	**Good identification**						**47767202**
Actinomyces israelii		100.0	0.36	FRU 99			
47767300	**Low discrimination**						**47767300**
					SPO		
Actinomyces israelii		88.57	0.44	FRU 99	1		
Clostridium butyricum		11.43	0.07	RHA 7	99		
47767302	**Acceptable identification**						**47767302**
					SPO		
Actinomyces israelii		93.80	0.47	FRU 99	1		
Clostridium butyricum		3.14	0.00	RHA 7	99		
Bifidobacterium adolescentis		1.96	0.12		1		
47767342	**Low discrimination**						**47767342**
					COC		
Actinomyces israelii		84.30	0.40	FRU 99	1		
Blautia producta		15.70	0.07	RHA 7	99		
47767400	**Low discrimination**						**47767400**
					SPO		
Actinomyces israelii		63.47	0.51	FRU 99	1		
Clostridium butyricum		36.53	0.26		99		
47767402	**Low discrimination**						**47767402**
					SPO		
Actinomyces israelii		82.79	0.54	FRU 99	1		
Clostridium butyricum		12.36	0.19		99		
Bifidobacterium adolescentis		4.25	0.26		1		
47767410	**Low discrimination**						**47767410**
					SPO		
Actinomyces israelii		49.99	0.37	FRU 99	1		
Bifidobacterium dentium		36.23	0.10	MLZ 13	1		
Clostridium butyricum		10.14	0.04		99		
47767440	**Low discrimination**						**47767440**
					COC	SPO	
Actinomyces israelii		48.69	0.44	FRU 99	1	1	
Blautia producta		45.05	0.24		99	1	
Clostridium butyricum		6.27	0.08		1	99	
47767442	**Low discrimination**						**47767442**
					COC	SPO	
Actinomyces israelii		89.43	0.47	FRU 99	1	1	
Blautia producta		6.13	0.07		99	1	
Clostridium butyricum		2.98	0.01		1	99	
47767500	**Low discrimination**						**47767500**
					COC	SPO	
Clostridium butyricum		74.36	0.50	FRU 99	1	99	
Actinomyces israelii		21.53	0.61	MAN 10	1	1	
Blautia producta		1.94	0.24		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47767502	**Low discrimination**						**47767502**
					SPO		
Bifidobacterium adolescentis		37.81	0.60	FRU 99	1		
Actinomyces israelii		29.76	0.64		1		
Clostridium butyricum		26.65	0.43		99		
47767510	**Low discrimination**						**47767510**
					COC	SPO	
Bifidobacterium dentium		80.66	0.44	FRU 99	1	1	
Blautia producta		8.12	0.32	BGL 91	99	1	
Clostridium butyricum		5.47	0.28		1	99	
47767512	**Good identification**						**47767512**
Bifidobacterium dentium		91.05	0.62	FRU 99			
Bifidobacterium adolescentis		7.83	0.61				
Actinomyces israelii		0.75	0.50				
47767530	**Good identification**						**47767530**
					COC	SPO	
Bifidobacterium dentium		98.76	0.49	FRU 99	1	1	
Blautia producta		0.62	0.15	BGL 91	99	1	
Clostridium butyricum		0.33	0.10		1	99	
47767532	**Very good identification**						**47767532**
Bifidobacterium dentium		99.86	0.66	FRU 99			
Bifidobacterium adolescentis		0.05	0.27				
Actinomyces israelii		0.03	0.30				
47767540	**Good identification**						**47767540**
					COC	SPO	
Blautia producta		92.27	0.58	FRU 99	99	1	
Actinomyces israelii		4.03	0.54		1	1	
Clostridium butyricum		3.11	0.31		1	99	
47767542	**Low discrimination**						**47767542**
					COC		
Blautia producta		51.16	0.41	FRU 99	99		
Actinomyces israelii		30.16	0.57	BGL 10	1		
Bifidobacterium adolescentis		12.18	0.44		1		
47767550	**Very good identification**						**47767550**
					COC		
Blautia producta		99.55	0.66	FRU 99	99		
Actinomyces israelii		0.22	0.40		1		
Bifidobacterium dentium		0.10	0.10		1		
47767552	**Low discrimination**						**47767552**
					COC		
Blautia producta		81.88	0.50	FRU 99	99		
Bifidobacterium adolescentis		7.94	0.46	BGL 10	1		
Bifidobacterium dentium		7.55	0.27		1		
47767560	**Acceptable identification**						**47767560**
					COC	SPO	
Blautia producta		93.87	0.41	FRU 99	99	1	
Clostridium butyricum		2.48	0.13	NAG 10	1	99	
Actinomyces israelii		2.36	0.34		1	1	
47767570	**Good identification**						**47767570**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		98.14	0.50	FRU 99	99	99	
Bifidobacterium dentium		1.59	0.15	NAG 10	1	99	
Mitsuokella multacida		0.15	0.09		1	1	
47767702	**Low discrimination**						**47767702**
					SPO		
Actinomyces israelii		52.55	0.45	FRU 99	1		
Clostridium butyricum		19.33	0.17	RHA 7	99		
Bifidobacterium adolescentis		8.96	0.26		1		
47767740	**Very good identification**						**47767740**
					COC	GRA	
Blautia producta		98.45	0.58	FRU 99	99	99	
Mitsuokella multacida		1.12	0.32		1	1	
Actinomyces israelii		0.32	0.35		1	99	
47767742	**Acceptable identification**						**47767742**
					COC	GRA	
Blautia producta		94.18	0.41	FRU 99	99	99	
Actinomyces israelii		4.18	0.38	BGL 10	1	99	
Mitsuokella multacida		1.07	0.15		1	1	
47767750	**Very good identification**						**47767750**
					COC	GRA	
Blautia producta		99.86	0.66	FRU 99	99	99	
Mitsuokella multacida		0.13	0.24		1	1	
Actinomyces israelii		0.02	0.21		1	99	
47767752	**Good identification**						**47767752**
					COC	GRA	
Blautia producta		99.46	0.50	FRU 99	99	99	
Actinomyces israelii		0.22	0.24	BGL 10	1	99	
Mitsuokella multacida		0.13	0.07		1	1	
47767760	**Good identification**						**47767760**
					COC	GRA	
Blautia producta		97.33	0.41	FRU 99	99	99	
Mitsuokella multacida		2.49	0.21	NAG 10	1	1	
Actinomyces israelii		0.18	0.15		1	99	
47767770	**Good identification**						**47767770**
					COC	GRA	
Blautia producta		99.72	0.50	FRU 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
47770000	**Low discrimination**						**47770000**
					GRA		
Fusobacterium mortiferum		85.77	0.58	MLT 11	1		
Lactobacillus jensenii		9.50	0.32		99		
Actinomyces naeslundii		3.52	0.46		99		
47770001	**Low discrimination**						**47770001**
					GRA		
Actinomyces naeslundii		85.90	0.49	ESL 12	99		
Fusobacterium mortiferum		14.10	0.24	GAL 96	1		
47770003	**Good identification**						**47770003**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47770020	**Low discrimination**						**47770020**
					SPO		
Actinomyces naeslundii		36.62	0.46	ESL 12	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		24.70	0.21	GAL 96	1		
Clostridium barati		10.00	0.10		99		
47770021	**Good identification**						**47770021**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47770023	**Good identification**						**47770023**
Actinomyces naeslundii		100.0	0.49	ESL 12			
47770041	**Good identification**						**47770041**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47770043	**Good identification**						**47770043**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47770061	**Good identification**						**47770061**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47770063	**Good identification**						**47770063**
Actinomyces naeslundii		100.0	0.36	ESL 12			
47770100	**Low discrimination**						**47770100**
					GRA	SPO	
Fusobacterium mortiferum		81.09	0.64	MLT 11	1	1	
Bifidobacterium breve		5.52	0.41		99	1	
Clostridium butyricum		4.22	0.24		99	99	
47770101	**Low discrimination**						**47770101**
					GRA		
Actinomyces naeslundii		88.14	0.57	ESL 12	99		
Fusobacterium mortiferum		11.86	0.30	GAL 96	1		
47770102	**Low discrimination**						**47770102**
					GRA		
Bifidobacterium breve		41.89	0.59	GAL 99	99		
Lactobacillus acidophilus		34.51	0.47		99		
Fusobacterium mortiferum		6.77	0.48		1		
47770103	**Low discrimination**						**47770103**
Actinomyces naeslundii		83.65	0.57	ESL 12			
Bifidobacterium breve		7.74	0.24	GAL 96			
Lactobacillus acidophilus		6.38	0.13				
47770112	**Good identification**						**47770112**
					GRA		
Leptotrichia buccalis		91.28	0.58	GAL 99	1		
Bifidobacterium breve		7.30	0.59		99		
Clostridium clostridioforme		0.96	0.39		50		
47770120	**Low discrimination**						**47770120**
					SPO		
Clostridium paraputrificum		43.20	0.37	RAF 7	99		
Clostridium ramosum		29.30	0.41	GAL 99	92		
Bifidobacterium breve		13.70	0.46		1		
47770121	**Good identification**						**47770121**
					SPO		
Actinomyces naeslundii		90.71	0.57	ESL 12	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		4.66	0.03	GAL 96	99		
Clostridium ramosum		3.16	0.06		92		
47770122	**Low discrimination**						**47770122**
					SPO		
Bifidobacterium breve		33.31	0.63	GAL 99	1		
Clostridium ramosum		28.18	0.51		92		
Lactobacillus acidophilus		15.44	0.47		1		
47770123	**Low discrimination**						**47770123**
					SPO		
Actinomyces naeslundii		67.91	0.57	ESL 12	1		
Bifidobacterium breve		11.17	0.29	GAL 96	1		
Clostridium ramosum		9.45	0.17		92		
47770130	**Low discrimination**						**47770130**
					GRA	SPO	
Bifidobacterium breve		86.02	0.46	GAL 99	99	1	
Leptotrichia buccalis		6.86	0.07	BGL 91	1	1	
Clostridium paraputrificum		2.74	0.03		99	99	
47770132	**Low discrimination**						**47770132**
					GRA		
Bifidobacterium breve		52.93	0.63	GAL 99	99		
Leptotrichia buccalis		41.36	0.41		1		
Clostridium clostridioforme		3.91	0.39		50		
47770140	**Low discrimination**						**47770140**
					GRA		
Bifidobacterium breve		71.57	0.41	GAL 99	99		
Fusobacterium mortiferum		10.63	0.30	BGL 91	1		
Actinomyces naeslundii		9.29	0.41		99		
47770141	**Good identification**						**47770141**
Actinomyces naeslundii		100.0	0.44	ESL 12			
47770142	**Low discrimination**						**47770142**
Bifidobacterium breve		88.16	0.59	GAL 99			
Lactobacillus acidophilus		8.98	0.32				
Actinomyces naeslundii		1.13	0.41				
47770143	**Low discrimination**						**47770143**
Actinomyces naeslundii		65.60	0.44	ESL 12			
Bifidobacterium breve		34.40	0.24	GAL 96			
47770150	**Good identification**						**47770150**
					GRA		
Bifidobacterium breve		98.73	0.41	GAL 99	99		
Leptotrichia buccalis		1.27	-.11	BGL 91	1		
47770152	**Low discrimination**						**47770152**
					GRA		
Bifidobacterium breve		88.16	0.59	GAL 99	99		
Leptotrichia buccalis		11.13	0.24		1		
Bifidobacterium adolescentis		0.49	0.19		99		
47770160	**Low discrimination**						**47770160**
					SPO		
Bifidobacterium breve		74.49	0.46	GAL 99	1		
Clostridium paraputrificum		17.68	0.18	BGL 91	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		5.44	0.41		1		
47770161	**Acceptable identification**						**47770161**
Actinomyces naeslundii		91.56	0.44	ESL 12			
Bifidobacterium breve		8.44	0.12	GAL 96			
47770162	**Good identification**						**47770162**
					SPO		
Bifidobacterium breve		91.70	0.63	GAL 99	1		
Lactobacillus acidophilus		5.25	0.32		1		
Clostridium paraputrificum		0.92	0.11		99		
47770163	**Low discrimination**						**47770163**
Actinomyces naeslundii		51.75	0.44	ESL 12			
Bifidobacterium breve		48.25	0.29	GAL 96			
47770170	**Good identification**						**47770170**
Bifidobacterium breve		100.0	0.46	GAL 99			
47770172	**Very good identification**						**47770172**
					GRA		
Bifidobacterium breve		99.14	0.63	GAL 99	99		
Leptotrichia buccalis		0.78	0.07		1		
Clostridium clostridioforme		0.07	0.05		50		
47770212	**Good identification**						**47770212**
Clostridium clostridioforme		99.56	0.45	XYL 91			
Leptotrichia buccalis		0.44	-.11	ARA 94			
47770232	**Good identification**						**47770232**
Clostridium clostridioforme		100.0	0.45	XYL 91			
47770302	**Low discrimination**						**47770302**
					SPO		
Clostridium clostridioforme		64.37	0.43	XYL 91	66		
Lactobacillus acidophilus		18.89	0.28	ARA 94	1		
Clostridium ramosum		6.14	0.19		92		
47770312	**Low discrimination**						**47770312**
					GRA		
Clostridium clostridioforme		87.59	0.54	XYL 91	50		
Leptotrichia buccalis		11.49	0.24	ARA 94	1		
Bifidobacterium breve		0.92	0.24		99		
47770320	**Acceptable identification**						**47770320**
					SPO	GRA	
Clostridium ramosum		94.36	0.43	GAL 99	92	50	
Prevotella oralis		3.25	0.17		1	1	
Clostridium paraputrificum		1.06	0.03		99	99	
47770322	**Low discrimination**						**47770322**
					SPO		
Clostridium ramosum		72.54	0.53	GAL 99	92		
Prevotella oralis		11.88	0.39		1		
Clostridium clostridioforme		7.69	0.43		66		
47770332	**Low discrimination**						**47770332**
					SPO		
Clostridium clostridioforme		88.51	0.54	XYL 91	66		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		6.40	0.40	ARA 94	1		
Clostridium ramosum		1.85	0.19		92		
47770402	**Low discrimination**						**47770402**
					SPO		
Clostridium clostridioforme		88.45	0.40	XYL 91	66		
Clostridium butyricum		7.24	0.09	RHA 88	99		
Actinomyces israelii		2.68	0.23		1		
47770412	**Good identification**						**47770412**
					GRA		
Clostridium clostridioforme		99.30	0.51	XYL 91	50		
Bifidobacterium adolescentis		0.49	0.15	RHA 88	99		
Leptotrichia buccalis		0.21	-.11		1		
47770422	**Good identification**						**47770422**
Clostridium clostridioforme		100.0	0.40	XYL 91			
47770432	**Good identification**						**47770432**
Clostridium clostridioforme		100.0	0.51	XYL 91			
47770500	**Good identification**						**47770500**
					GRA	SPO	
Clostridium butyricum		95.43	0.40	GAL 99	99	99	
Fusobacterium mortiferum		2.06	0.30	XYL 97	1	1	
Actinomyces israelii		1.52	0.30		99	1	
47770502	**Low discrimination**						**47770502**
					SPO		
Clostridium clostridioforme		45.10	0.49	XYL 91	66		
Clostridium butyricum		26.48	0.33	RHA 88	99		
Bifidobacterium adolescentis		24.65	0.47		1		
47770512	**Low discrimination**						**47770512**
					GRA		
Clostridium clostridioforme		81.84	0.60	XYL 91	50		
Bifidobacterium adolescentis		12.00	0.49	RHA 88	99		
Leptotrichia buccalis		5.03	0.24		1		
47770522	**Low discrimination**						**47770522**
					GRA	SPO	
Clostridium clostridioforme		81.35	0.49	XYL 91	50	66	
Parabacteroides distasonis		4.76	0.33	RHA 88	1	1	
Clostridium butyricum		4.15	0.15		99	99	
47770532	**Very good identification**						**47770532**
					GRA		
Clostridium clostridioforme		97.44	0.60	XYL 91	50		
Bifidobacterium breve		0.85	0.29	RHA 88	99		
Parabacteroides distasonis		0.67	0.29		1		
47770602	**Good identification**						**47770602**
Clostridium clostridioforme		100.0	0.54	XYL 91			
47770612	**Very good identification**						**47770612**
Clostridium clostridioforme		100.0	0.66	XYL 91			
47770622	**Good identification**						**47770622**
Clostridium clostridioforme		100.0	0.54	XYL 91			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47770632	**Very good identification**					**47770632**
Clostridium clostridioforme		100.0	0.66	XYL 91		
47770702	**Very good identification**					**47770702**
					SPO	
Clostridium clostridioforme		99.64	0.63	XYL 91	66	
Clostridium butyricum		0.25	0.07		99	
Bifidobacterium adolescentis		0.08	0.13		1	
47770710	**Good identification**					**47770710**
Clostridium clostridioforme		99.28	0.41	XYL 91		
Clostridium butyricum		0.72	-.08	BGL 99		
47770712	**Very good identification**					**47770712**
					GRA	
Clostridium clostridioforme		99.97	0.75	XYL 91	50	
Bifidobacterium adolescentis		0.02	0.15		99	
Leptotrichia buccalis		0.01	-.11		1	
47770713	**Good identification**					**47770713**
Clostridium clostridioforme		100.0	0.41	XYL 91		
47770722	**Very good identification**					**47770722**
					SPO	
Clostridium clostridioforme		96.79	0.63	XYL 91	66	
Parabacteroides distasonis		2.32	0.41		1	
Clostridium ramosum		0.59	0.19		92	
47770730	**Good identification**					**47770730**
Clostridium clostridioforme		100.0	0.41	XYL 91		
47770732	**Very good identification**					**47770732**
Clostridium clostridioforme		99.69	0.75	XYL 91		
Parabacteroides distasonis		0.28	0.37			
Bacteroides ovatus		0.03	0.03			
47770733	**Good identification**					**47770733**
Clostridium clostridioforme		100.0	0.41	XYL 91		
47770752	**Good identification**					**47770752**
Clostridium clostridioforme		100.0	0.41	XYL 91		
47770772	**Good identification**					**47770772**
Clostridium clostridioforme		100.0	0.41	XYL 91		
47771000	**Low discrimination**					**47771000**
					GRA	
Lactobacillus jensenii		73.00	0.66	RAF 1	99	
Fusobacterium mortiferum		9.99	0.61		1	
Actinomyces naeslundii		6.56	0.70		99	
47771001	**Good identification**					**47771001**
					GRA	
Actinomyces naeslundii		91.55	0.73	ESL 12	99	
Lactobacillus jensenii		6.86	0.32		99	
Fusobacterium mortiferum		0.94	0.27		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47771002	**Low discrimination**					**47771002**
Lactobacillus acidophilus		48.49	0.63	LAC 92		
Lactobacillus jensenii		31.13	0.63			
Bifidobacterium breve		6.84	0.59			
47771003	**Low discrimination**					**47771003**
Actinomyces naeslundii		86.87	0.73	ESL 12		
Lactobacillus acidophilus		6.76	0.29			
Lactobacillus jensenii		4.34	0.29			
47771010	**Low discrimination**					**47771010**
					GRA	
Lactobacillus jensenii		87.22	0.58	RAF 1	99	
Leptotrichia buccalis		4.84	0.24		1	
Bifidobacterium breve		3.79	0.41		99	
47771011	**Low discrimination**					**47771011**
					GRA	
Actinomyces naeslundii		66.62	0.51	ESL 12	99	
Lactobacillus jensenii		31.63	0.24	MLZ 5	99	
Leptotrichia buccalis		1.75	-.11		1	
47771012	**Low discrimination**					**47771012**
					GRA	
Leptotrichia buccalis		81.34	0.58	LAC 99	1	
Lactobacillus jensenii		9.88	0.55		99	
Bifidobacterium breve		6.51	0.59		99	
47771020	**Low discrimination**					**47771020**
					SPO	
Lactobacillus jensenii		42.53	0.56	RAF 1	1	
Actinomyces naeslundii		15.28	0.70		1	
Clostridium paraputrificum		13.81	0.37		99	
47771021	**Very good identification**					**47771021**
					SPO	
Actinomyces naeslundii		96.61	0.73	ESL 12	1	
Lactobacillus jensenii		1.81	0.21		1	
Clostridium paraputrificum		0.59	0.03		99	
47771022	**Low discrimination**					**47771022**
					GRA	
Lactobacillus acidophilus		39.98	0.63	LAC 92	99	
Prevotella oralis		27.14	0.59		1	
Bifidobacterium breve		10.02	0.63		99	
47771023	**Low discrimination**					**47771023**
					GRA	
Actinomyces naeslundii		84.31	0.73	ESL 12	99	
Lactobacillus acidophilus		6.56	0.29		99	
Prevotella oralis		4.45	0.25		1	
47771030	**Low discrimination**					**47771030**
Lactobacillus jensenii		70.24	0.47	RAF 1		
Bifidobacterium breve		21.70	0.46			
Actinomyces naeslundii		3.99	0.48			
47771031	**Low discrimination**					**47771031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		86.56	0.51	ESL 12		
Lactobacillus jensenii		10.27	0.13	MLZ 5		
Bifidobacterium breve		3.17	0.12			
47771032	**Low discrimination**					**47771032**
					GRA	
Bifidobacterium breve		45.99	0.63	LAC 99	99	
Leptotrichia buccalis		35.93	0.41		1	
Lactobacillus jensenii		9.82	0.44		99	
47771033	**Low discrimination**					**47771033**
					GRA	
Actinomyces naeslundii		56.69	0.51	ESL 12	99	
Bifidobacterium breve		21.01	0.29	MLZ 5	99	
Leptotrichia buccalis		16.42	0.07		1	
47771040	**Low discrimination**					**47771040**
Actinomyces israelii		28.22	0.49	XYL 92		
Actinomyces naeslundii		23.88	0.57			
Bifidobacterium breve		21.81	0.41			
47771041	**Very good identification**					**47771041**
Actinomyces naeslundii		99.21	0.60	ESL 12		
Actinomyces israelii		0.79	0.14			
47771042	**Low discrimination**					**47771042**
Bifidobacterium breve		44.38	0.59	LAC 99		
Lactobacillus acidophilus		38.90	0.48			
Actinomyces israelii		8.52	0.52			
47771043	**Low discrimination**					**47771043**
Actinomyces naeslundii		88.60	0.60	ESL 12		
Bifidobacterium breve		5.51	0.24			
Lactobacillus acidophilus		4.83	0.14			
47771050	**Low discrimination**					**47771050**
Bifidobacterium breve		66.19	0.41	LAC 99		
Lactobacillus jensenii		15.39	0.24	BGL 91		
Actinomyces israelii		12.80	0.35			
47771051	**Good identification**					**47771051**
Actinomyces naeslundii		100.0	0.38	ESL 12		
47771052	**Low discrimination**					**47771052**
					GRA	
Bifidobacterium breve		83.92	0.59	LAC 99	99	
Leptotrichia buccalis		10.60	0.24		1	
Actinomyces israelii		2.41	0.38		99	
47771060	**Low discrimination**					**47771060**
					SPO	
Bifidobacterium breve		45.08	0.46	LAC 99	1	
Actinomyces naeslundii		27.76	0.57	BGL 91	1	
Clostridium paraputrificum		10.70	0.18		99	
47771061	**Very good identification**					**47771061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		98.92	0.60	ESL 12			
Bifidobacterium breve		1.08	0.12				
47771062	**Low discrimination**						**47771062**
Bifidobacterium breve		61.15	0.63	LAC 99			
Lactobacillus acidophilus		30.15	0.48				
Actinomyces naeslundii		3.73	0.57				
47771063	**Low discrimination**						**47771063**
Actinomyces naeslundii		85.83	0.60	ESL 12			
Bifidobacterium breve		9.49	0.29				
Lactobacillus acidophilus		4.68	0.14				
47771070	**Acceptable identification**						**47771070**
Bifidobacterium breve		93.28	0.46	LAC 99			
Lactobacillus jensenii		3.05	0.13	BGL 91			
Actinomyces naeslundii		3.02	0.35				
47771071	**Low discrimination**						**47771071**
Actinomyces naeslundii		82.80	0.38	ESL 12			
Bifidobacterium breve		17.20	0.12	MLZ 5			
47771072	**Very good identification**						**47771072**
					GRA		
Bifidobacterium breve		98.05	0.63	LAC 99	99		
Leptotrichia buccalis		0.77	0.07		1		
Lactobacillus acidophilus		0.49	0.14		99		
47771100	**Low discrimination**						**47771100**
					COC	SPO	
Bifidobacterium breve		32.70	0.76	BGL 91	1	1	
Clostridium butyricum		25.04	0.58		1	99	
Atopobium parvulum		14.97	0.61		99	1	
47771101	**Good identification**						**47771101**
					SPO		
Actinomyces naeslundii		90.26	0.80	ESL 12	1		
Bifidobacterium breve		3.41	0.41		1		
Clostridium butyricum		2.61	0.24		99		
47771102	**Low discrimination**						**47771102**
					GRA		
Bifidobacterium breve		46.94	0.93		99		
Lactobacillus acidophilus		38.67	0.81		99		
Leptotrichia buccalis		5.93	0.58		1		
47771103	**Low discrimination**						**47771103**
Actinomyces naeslundii		55.39	0.80	ESL 12			
Bifidobacterium breve		21.14	0.59				
Lactobacillus acidophilus		17.42	0.47				
47771110	**Low discrimination**						**47771110**
					GRA	SPO	
Leptotrichia buccalis		54.00	0.58	BGL 99	1	1	
Bifidobacterium breve		42.30	0.76		99	1	
Clostridium butyricum		1.71	0.36		99	99	
47771111	**Low discrimination**						**47771111**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		37.36	0.58	ESL 12	99		
Leptotrichia buccalis		34.22	0.24	MLZ 5	1		
Bifidobacterium breve		26.80	0.41		99		
47771112	**Good identification**						**47771112**
					GRA		
Leptotrichia buccalis		92.16	0.92		1		
Bifidobacterium breve		7.37	0.93		99		
Bifidobacterium adolescentis		0.33	0.69		99		
47771113	**Good identification**						**47771113**
					GRA		
Leptotrichia buccalis		91.24	0.58	URE 1	1		
Bifidobacterium breve		7.30	0.59		99		
Actinomyces naeslundii		1.01	0.58		99		
47771120	**Low discrimination**						**47771120**
					SPO		
Clostridium paraputrificum		46.22	0.71	RAF 7	99		
Clostridium ramosum		31.35	0.75		92		
Bifidobacterium breve		14.66	0.80		1		
47771121	**Low discrimination**						**47771121**
					SPO		
Actinomyces naeslundii		68.96	0.80	ESL 12	1		
Clostridium paraputrificum		14.60	0.37		99		
Clostridium ramosum		9.90	0.41		92		
47771122	**Low discrimination**						**47771122**
					SPO		
Bifidobacterium breve		34.16	0.97		1		
Clostridium ramosum		28.91	0.85		92		
Lactobacillus acidophilus		15.83	0.81		1		
47771123	**Low discrimination**						**47771123**
					SPO		
Actinomyces naeslundii		33.57	0.80	ESL 12	1		
Bifidobacterium breve		22.78	0.63		1		
Clostridium ramosum		19.28	0.51		92		
47771130	**Low discrimination**						**47771130**
					GRA	SPO	
Bifidobacterium breve		87.10	0.80	BGL 91	99	1	
Leptotrichia buccalis		6.95	0.41		1	1	
Clostridium paraputrificum		2.78	0.37		99	99	
47771131	**Low discrimination**						**47771131**
					GRA		
Bifidobacterium breve		52.15	0.46	URE 1	99		
Actinomyces naeslundii		40.90	0.58	BGL 91	99		
Leptotrichia buccalis		4.16	0.07		1		
47771132	**Low discrimination**						**47771132**
					GRA		
Bifidobacterium breve		55.04	0.97		99		
Leptotrichia buccalis		43.00	0.76		1		
Parabacteroides distasonis		0.72	0.66		1		
47771133	**Low discrimination**						**47771133**
					GRA		
Bifidobacterium breve		52.83	0.63	URE 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Leptotrichia buccalis		41.27	0.41		1	
Actinomyces naeslundii		4.10	0.58		99	
47771140	**Low discrimination**					**47771140**
					SPO	
Bifidobacterium breve		85.11	0.76	BGL 91	1	
Clostridium butyricum		5.67	0.40		99	
Actinomyces israelii		4.45	0.59		1	
47771141	**Low discrimination**					**47771141**
					SPO	
Actinomyces naeslundii		80.69	0.67	ESL 12	1	
Bifidobacterium breve		17.26	0.41		1	
Clostridium butyricum		1.15	0.06		99	
47771142	**Low discrimination**					**47771142**
Bifidobacterium breve		89.10	0.93			
Lactobacillus acidophilus		9.07	0.66			
Actinomyces israelii		0.69	0.62			
47771143	**Low discrimination**					**47771143**
Bifidobacterium breve		63.28	0.59	URE 1		
Actinomyces naeslundii		29.25	0.67			
Lactobacillus acidophilus		6.44	0.32			
47771150	**Very good identification**					**47771150**
					GRA	
Bifidobacterium breve		97.41	0.76	BGL 91	99	
Leptotrichia buccalis		1.26	0.24		1	
Actinomyces israelii		0.76	0.45		99	
47771151	**Low discrimination**					**47771151**
					GRA	
Bifidobacterium breve		79.43	0.41	URE 1	99	
Actinomyces naeslundii		19.54	0.46	BGL 91	99	
Leptotrichia buccalis		1.02	-.11		1	
47771152	**Low discrimination**					**47771152**
					GRA	
Bifidobacterium breve		88.16	0.93		99	
Leptotrichia buccalis		11.13	0.58		1	
Bifidobacterium adolescentis		0.49	0.53		99	
47771153	**Low discrimination**					**47771153**
					GRA	
Bifidobacterium breve		86.41	0.59	URE 1	99	
Leptotrichia buccalis		10.91	0.24		1	
Actinomyces naeslundii		2.10	0.46		99	
47771160	**Low discrimination**					**47771160**
					SPO	
Bifidobacterium breve		77.12	0.80	BGL 91	1	
Clostridium paraputrificum		18.31	0.52		99	
Clostridium ramosum		1.67	0.41		92	
47771161	**Low discrimination**					**47771161**
					SPO	
Actinomyces naeslundii		67.62	0.67	ESL 12	1	
Bifidobacterium breve		25.72	0.46		1	
Clostridium paraputrificum		6.11	0.18		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47771162	**Good identification**						**47771162**
					SPO		
Bifidobacterium breve		92.29	0.97		1		
Lactobacillus acidophilus		5.29	0.66		1		
Clostridium paraputrificum		0.93	0.45		99		
47771163	**Low discrimination**						**47771163**
Bifidobacterium breve		74.61	0.63	URE 1			
Actinomyces naeslundii		19.40	0.67				
Lactobacillus acidophilus		4.27	0.32				
47771170	**Very good identification**						**47771170**
					SPO		
Bifidobacterium breve		99.52	0.80	BGL 91	1		
Clostridium paraputrificum		0.24	0.18		99		
Actinomyces naeslundii		0.09	0.43		1		
47771171	**Low discrimination**						**47771171**
Bifidobacterium breve		87.84	0.46	URE 1			
Actinomyces naeslundii		12.16	0.46	BGL 91			
47771172	**Excellent identification**						**47771172**
					GRA		
Bifidobacterium breve		99.11	0.97		99		
Leptotrichia buccalis		0.78	0.41		1		
Lactobacillus acidophilus		0.06	0.32		99		
47771173	**Very good identification**						**47771173**
					GRA		
Bifidobacterium breve		97.89	0.63	URE 1	99		
Actinomyces naeslundii		1.34	0.46		99		
Leptotrichia buccalis		0.77	0.07		1		
47771201	**Good identification**						**47771201**
Actinomyces naeslundii		100.0	0.38	ESL 12			
47771202	**Low discrimination**						**47771202**
					GRA		
Lactobacillus acidophilus		55.08	0.44	LAC 92	99		
Clostridium clostridioforme		19.53	0.42	RHA 7	50		
Prevotella oralis		11.20	0.31		1		
47771203	**Low discrimination**						**47771203**
Actinomyces naeslundii		52.53	0.38	ESL 12			
Lactobacillus acidophilus		30.47	0.10	RHA 1			
Clostridium clostridioforme		10.81	0.08				
47771212	**Low discrimination**						**47771212**
					GRA		
Clostridium clostridioforme		84.05	0.53	XYL 91	50		
Leptotrichia buccalis		12.30	0.24	ARA 94	1		
Lactobacillus jensenii		1.49	0.21		99		
47771220	**Low discrimination**						**47771220**
					SPO	GRA	
Clostridium ramosum		75.57	0.43	LAC 99	92	50	
Prevotella oralis		16.46	0.31		1	1	
Lactobacillus jensenii		2.62	0.21		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47771221	**Low discrimination**						**47771221**
					SPO	GRA	
Actinomyces naeslundii		60.26	0.38	ESL 12	1	99	
Clostridium ramosum		32.63	0.09	RHA 1	92	50	
Prevotella oralis		7.11	-.03		1	1	
47771222	**Low discrimination**						**47771222**
					SPO	GRA	
Prevotella oralis		41.39	0.53	TRE 6	1	1	
Clostridium ramosum		40.00	0.53	LAC 94	92	50	
Lactobacillus acidophilus		10.71	0.44		1	99	
47771232	**Low discrimination**						**47771232**
Clostridium clostridioforme		85.04	0.53	XYL 91			
Parabacteroides distasonis		6.86	0.40	ARA 94			
Prevotella oralis		2.06	0.19				
47771300	**Low discrimination**						**47771300**
					SPO		
Clostridium ramosum		41.68	0.43	NAG 99	92		
Clostridium butyricum		19.39	0.32		99		
Actinomyces israelii		8.28	0.47		1		
47771301	**Low discrimination**						**47771301**
					SPO		
Actinomyces naeslundii		78.20	0.46	ESL 12	1		
Clostridium ramosum		14.88	0.09	RHA 1	92		
Clostridium butyricum		6.92	-.02		99		
47771302	**Low discrimination**						**47771302**
					SPO	GRA	
Lactobacillus acidophilus		50.88	0.62	RHA 7	1	99	
Clostridium ramosum		16.53	0.53		92	50	
Prevotella oralis		14.10	0.51		1	1	
47771312	**Low discrimination**						**47771312**
					GRA		
Leptotrichia buccalis		75.17	0.58	RHA 1	1		
Clostridium clostridioforme		17.37	0.62		50		
Bifidobacterium breve		6.01	0.59		99		
47771320	**Good identification**						**47771320**
					SPO	GRA	
Clostridium ramosum		94.77	0.77		92	50	
Prevotella oralis		3.27	0.51		1	1	
Clostridium paraputrificum		1.07	0.37		99	99	
47771321	**Acceptable identification**						**47771321**
					SPO	GRA	
Clostridium ramosum		90.50	0.43	URE 1	92	50	
Actinomyces naeslundii		4.80	0.46		1	99	
Prevotella oralis		3.12	0.17		1	1	
47771322	**Low discrimination**						**47771322**
					SPO		
Clostridium ramosum		78.50	0.87		92		
Prevotella oralis		12.85	0.73		1		
Parabacteroides distasonis		5.09	0.79		1		
47771323	**Low discrimination**						**47771323**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		77.74	0.53	URE 1	92		
Prevotella oralis		12.73	0.39		1		
Parabacteroides distasonis		5.04	0.45		1		
47771330	**Low discrimination**						**47771330**
					SPO	GRA	
Clostridium ramosum		61.56	0.43	MLZ 1	92	50	
Bifidobacterium breve		21.71	0.46		1	99	
Parabacteroides distasonis		8.59	0.40		1	1	
47771332	**Low discrimination**						**47771332**
					SPO		
Parabacteroides distasonis		44.89	0.75	XYL 90	1		
Clostridium clostridioforme		18.83	0.62		66		
Clostridium ramosum		13.00	0.53		92		
47771333	**Low discrimination**						**47771333**
					SPO		
Parabacteroides distasonis		44.72	0.40	XYL 90	1		
Clostridium clostridioforme		18.75	0.28	URE 1	66		
Clostridium ramosum		12.95	0.19		92		
47771340	**Low discrimination**						**47771340**
					SPO		
Bifidobacterium breve		57.03	0.41	RHA 1	1		
Actinomyces israelii		22.22	0.40	BGL 91	1		
Clostridium butyricum		11.63	0.14		99		
47771342	**Low discrimination**						**47771342**
Bifidobacterium breve		53.15	0.59	RHA 1			
Lactobacillus acidophilus		40.33	0.47				
Actinomyces israelii		3.07	0.43				
47771350	**Acceptable identification**						**47771350**
					COC		
Bifidobacterium breve		91.69	0.41	RHA 1	1		
Actinomyces israelii		5.34	0.26	BGL 91	1		
Blautia producta		1.79	-.03		99		
47771352	**Low discrimination**						**47771352**
					GRA		
Bifidobacterium breve		84.94	0.59	RHA 1	99		
Leptotrichia buccalis		10.73	0.24		1		
Clostridium clostridioforme		2.48	0.28		50		
47771360	**Low discrimination**						**47771360**
					SPO		
Clostridium ramosum		66.24	0.43	SOR 1	92		
Bifidobacterium breve		23.36	0.46		1		
Clostridium paraputrificum		5.55	0.18		99		
47771362	**Low discrimination**						**47771362**
					SPO		
Clostridium ramosum		39.67	0.53	SOR 1	92		
Bifidobacterium breve		35.37	0.63		1		
Lactobacillus acidophilus		15.10	0.47		1		
47771370	**Good identification**						**47771370**
					SPO		
Bifidobacterium breve		97.22	0.46	RHA 1	1		
Clostridium ramosum		2.78	0.09	BGL 91	92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47771372	**Good identification**					**47771372**
					GRA	
Bifidobacterium breve		92.51	0.63	RHA 1	99	
Parabacteroides distasonis		3.62	0.40		1	
Clostridium clostridioforme		1.52	0.28		50	
47771400	**Low discrimination**					**47771400**
					SPO	
Clostridium butyricum		88.64	0.51	XYL 97	99	
Actinomyces israelii		8.49	0.54	LAC 96	1	
Lactobacillus jensenii		2.18	0.32		1	
47771402	**Low discrimination**					**47771402**
					SPO	
Clostridium butyricum		47.80	0.44	XYL 97	99	
Clostridium clostridioforme		17.68	0.48	LAC 96	66	
Actinomyces israelii		17.66	0.57		1	
47771412	**Low discrimination**					**47771412**
					GRA	
Clostridium clostridioforme		76.28	0.59	XYL 91	50	
Bifidobacterium adolescentis		12.48	0.49	RHA 88	99	
Leptotrichia buccalis		5.23	0.24		1	
47771421	**Low discrimination**					**47771421**
					SPO	
Actinomyces naeslundii		79.01	0.38	ESL 12	1	
Clostridium butyricum		20.99	-.02	ARA 1	99	
47771422	**Low discrimination**					**47771422**
					SPO	
Clostridium clostridioforme		55.28	0.48	XYL 91	66	
Clostridium butyricum		12.99	0.25	RHA 88	99	
Lactobacillus acidophilus		9.79	0.29		1	
47771423	**Low discrimination**					**47771423**
Actinomyces naeslundii		69.47	0.38	ESL 12		
Clostridium clostridioforme		30.53	0.14	ARA 1		
47771432	**Very good identification**					**47771432**
					GRA	
Clostridium clostridioforme		96.57	0.59	XYL 91	50	
Bifidobacterium breve		0.94	0.29	RHA 88	99	
Parabacteroides distasonis		0.75	0.29		1	
47771442	**Low discrimination**					**47771442**
					SPO	
Actinomyces israelii		51.20	0.50	XYL 92	1	
Clostridium butyricum		30.99	0.25		99	
Bifidobacterium adolescentis		9.94	0.32		1	
47771500	**Very good identification**					**47771500**
					SPO	
Clostridium butyricum		97.73	0.74	XYL 97	99	
Actinomyces israelii		1.56	0.64		1	
Bifidobacterium adolescentis		0.36	0.47		1	
47771501	**Acceptable identification**					**47771501**
					SPO	
Clostridium butyricum		94.24	0.40	XYL 97	99	
Actinomyces naeslundii		3.66	0.46	URE 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.50	0.30		1		
47771502	**Low discrimination**						**47771502**
					SPO		
Clostridium butyricum		47.11	0.67	XYL 97	99		
Bifidobacterium adolescentis		43.85	0.81		1		
Actinomyces israelii		2.90	0.68		1		
47771510	**Low discrimination**						**47771510**
					SPO		
Clostridium butyricum		82.06	0.52	XYL 97	99		
Bifidobacterium adolescentis		6.97	0.49	MLZ 5	1		
Actinomyces israelii		3.72	0.50		1		
47771512	**Low discrimination**						**47771512**
					GRA		
Bifidobacterium adolescentis		58.13	0.83	XYL 87	99		
Leptotrichia buccalis		24.34	0.58		1		
Clostridium clostridioforme		12.01	0.68		50		
47771513	**Low discrimination**						**47771513**
					GRA		
Bifidobacterium adolescentis		57.99	0.49	XYL 87	99		
Leptotrichia buccalis		24.28	0.24	URE 1	1		
Clostridium clostridioforme		11.98	0.34		50		
47771520	**Low discrimination**						**47771520**
Clostridium butyricum		81.74	0.56	XYL 97			
Clostridium paraputrificum		7.73	0.37	NAG 8			
Clostridium ramosum		5.24	0.41				
47771522	**Low discrimination**						**47771522**
					GRA	SPO	
Parabacteroides distasonis		22.13	0.67	XYL 90	1	1	
Clostridium butyricum		19.33	0.49		99	99	
Bifidobacterium breve		15.06	0.63		99	1	
47771523	**Low discrimination**						**47771523**
					GRA	SPO	
Actinomyces naeslundii		18.42	0.46	ESL 12	99	1	
Parabacteroides distasonis		18.37	0.33	ARA 1	1	1	
Clostridium butyricum		16.05	0.15		99	99	
47771532	**Low discrimination**						**47771532**
					GRA		
Clostridium clostridioforme		53.25	0.68	XYL 91	50		
Bifidobacterium breve		15.35	0.63	RHA 88	99		
Parabacteroides distasonis		12.14	0.63		1		
47771540	**Acceptable identification**						**47771540**
					SPO		
Clostridium butyricum		90.28	0.56	XYL 97	99		
Actinomyces israelii		6.45	0.57	SOR 8	1		
Bifidobacterium breve		1.52	0.41		1		
47771542	**Low discrimination**						**47771542**
					SPO		
Bifidobacterium adolescentis		42.77	0.66	XYL 87	1		
Clostridium butyricum		32.33	0.49	SOR 11	99		
Bifidobacterium breve		14.17	0.59		1		
47771552	**Low discrimination**						**47771552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium adolescentis		72.13	0.67	XYL 87	99		
Bifidobacterium breve		19.55	0.59	SOR 11	99		
Leptotrichia buccalis		2.47	0.24		1		
47771560	**Low discrimination**						**47771560**
					SPO		
Clostridium butyricum		66.04	0.38	XYL 97	99		
Bifidobacterium breve		22.76	0.46	NAG 8	1		
Clostridium paraputrificum		5.40	0.18	SOR 8	99		
47771562	**Low discrimination**						**47771562**
					SPO		
Bifidobacterium breve		79.58	0.63	ARA 1	1		
Clostridium butyricum		8.88	0.31		99		
Lactobacillus acidophilus		4.56	0.32		1		
47771570	**Low discrimination**						**47771570**
					COC	SPO	
Bifidobacterium breve		76.68	0.46	ARA 1	1	1	
Clostridium butyricum		11.71	0.16	BGL 91	1	99	
Blautia producta		9.27	0.15		99	1	
47771572	**Good identification**						**47771572**
Bifidobacterium breve		92.21	0.63	ARA 1			
Clostridium clostridioforme		3.23	0.34				
Bifidobacterium adolescentis		1.93	0.33				
47771602	**Very good identification**						**47771602**
					SPO		
Clostridium clostridioforme		97.59	0.63	XYL 91	66		
Clostridium butyricum		1.11	0.18		99		
Actinomyces israelii		1.00	0.38		1		
47771610	**Acceptable identification**						**47771610**
					SPO		
Clostridium clostridioforme		94.72	0.40	XYL 91	66		
Clostridium butyricum		3.18	0.03	BGL 99	99		
Actinomyces israelii		2.10	0.21		1		
47771612	**Very good identification**						**47771612**
Clostridium clostridioforme		99.92	0.74	XYL 91			
Actinomyces israelii		0.03	0.24				
Bifidobacterium adolescentis		0.02	0.15				
47771613	**Good identification**						**47771613**
Clostridium clostridioforme		100.0	0.40	XYL 91			
47771622	**Very good identification**						**47771622**
Clostridium clostridioforme		95.52	0.63	XYL 91			
Parabacteroides distasonis		2.55	0.41				
Prevotella oralis		0.67	0.19				
47771630	**Good identification**						**47771630**
Clostridium clostridioforme		100.0	0.40	XYL 91			
47771632	**Very good identification**						**47771632**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		99.65	0.74	XYL 91		
Parabacteroides distasonis		0.31	0.37			
Bacteroides ovatus		0.03	0.03			
47771633	**Good identification**					**47771633**
Clostridium clostridioforme		100.0	0.40	XYL 91		
47771652	**Good identification**					**47771652**
Clostridium clostridioforme		98.72	0.40	XYL 91		
Actinomyces israelii		1.28	0.17	SOR 1		
47771672	**Good identification**					**47771672**
Clostridium clostridioforme		100.0	0.40	XYL 91		
47771700	**Acceptable identification**					**47771700**
					SPO	
Clostridium butyricum		90.48	0.48	XYL 97	99	
Clostridium clostridioforme		4.35	0.37	RHA 3	66	
Actinomyces israelii		3.52	0.45		1	
47771702	**Low discrimination**					**47771702**
					SPO	
Clostridium clostridioforme		87.80	0.72	XYL 91	66	
Clostridium butyricum		7.18	0.41		99	
Bifidobacterium adolescentis		2.18	0.47		1	
47771703	**Low discrimination**					**47771703**
					SPO	
Clostridium clostridioforme		89.38	0.37	XYL 91	66	
Clostridium butyricum		7.31	0.07	URE 1	99	
Bifidobacterium adolescentis		2.22	0.13		1	
47771710	**Low discrimination**					**47771710**
					SPO	
Clostridium clostridioforme		76.75	0.49	XYL 91	66	
Clostridium butyricum		18.45	0.26	BGL 99	99	
Actinomyces israelii		2.03	0.31		1	
47771712	**Very good identification**					**47771712**
					GRA	
Clostridium clostridioforme		98.80	0.83	XYL 91	50	
Bifidobacterium adolescentis		0.66	0.49		99	
Leptotrichia buccalis		0.28	0.24		1	
47771713	**Good identification**					**47771713**
					GRA	
Clostridium clostridioforme		99.06	0.49	XYL 91	50	
Bifidobacterium adolescentis		0.66	0.15	URE 1	99	
Leptotrichia buccalis		0.28	-.11		1	
47771720	**Low discrimination**					**47771720**
Clostridium ramosum		53.47	0.43	ARA 1		
Clostridium butyricum		19.46	0.30			
Clostridium clostridioforme		10.76	0.37			
47771722	**Low discrimination**					**47771722**
					SPO	
Clostridium clostridioforme		47.37	0.72	XYL 91	66	
Parabacteroides distasonis		37.39	0.75		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium ramosum		9.51	0.53		92		
47771723	**Low discrimination**						**47771723**
					SPO		
Clostridium clostridioforme		47.75	0.37	XYL 91	66		
Parabacteroides distasonis		37.69	0.41	URE 1	1		
Clostridium ramosum		9.59	0.19		92		
47771730	**Low discrimination**						**47771730**
					GRA	SPO	
Clostridium clostridioforme		87.48	0.49	XYL 91	50	66	
Parabacteroides distasonis		8.16	0.37	BGL 99	1	1	
Clostridium butyricum		1.83	0.08		99	99	
47771732	**Good identification**						**47771732**
Clostridium clostridioforme		90.55	0.83	XYL 91			
Parabacteroides distasonis		8.45	0.71				
Bacteroides ovatus		0.76	0.37				
47771733	**Acceptable identification**						**47771733**
Clostridium clostridioforme		90.77	0.49	XYL 91			
Parabacteroides distasonis		8.47	0.37	URE 1			
Bacteroides ovatus		0.76	0.03				
47771742	**Low discrimination**						**47771742**
					SPO		
Clostridium clostridioforme		36.22	0.37	XYL 91	66		
Clostridium butyricum		25.48	0.23	SOR 1	99		
Actinomyces israelii		17.08	0.41		1		
47771752	**Low discrimination**						**47771752**
					COC		
Clostridium clostridioforme		84.81	0.49	XYL 91	1		
Bifidobacterium adolescentis		6.92	0.33	SOR 1	1		
Blautia producta		3.95	0.15		99		
47771762	**Low discrimination**						**47771762**
Clostridium clostridioforme		38.16	0.37	XYL 91			
Parabacteroides distasonis		30.12	0.41	SOR 1			
Bacteroides ovatus		9.05	0.16				
47771772	**Low discrimination**						**47771772**
					GRA		
Clostridium clostridioforme		85.29	0.49	XYL 91	50		
Parabacteroides distasonis		7.96	0.37	SOR 1	1		
Bifidobacterium breve		3.35	0.29		99		
47772000	**Low discrimination**						**47772000**
					GRA		
Actinomyces naeslundii		66.38	0.51	ESL 12	99		
Actinomyces israelii		25.20	0.34	GAL 96	99		
Fusobacterium mortiferum		8.42	0.24		1		
47772001	**Good identification**						**47772001**
Actinomyces naeslundii		100.0	0.54	ESL 12			
47772002	**Low discrimination**						**47772002**
Actinomyces naeslundii		54.09	0.51	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		30.79	0.37	GAL 96		
Bifidobacterium breve		11.01	0.24			
47772003	**Good identification**					**47772003**
Actinomyces naeslundii		100.0	0.54	ESL 12		
47772020	**Low discrimination**					**47772020**
					SPO	
Actinomyces naeslundii		66.95	0.51	ESL 12	1	
Clostridium ramosum		29.04	0.19	GAL 96	92	
Bifidobacterium breve		2.40	0.12		1	
47772021	**Good identification**					**47772021**
Actinomyces naeslundii		100.0	0.54	ESL 12		
47772023	**Good identification**					**47772023**
Actinomyces naeslundii		100.0	0.54	ESL 12		
47772040	**Low discrimination**					**47772040**
Actinomyces naeslundii		54.45	0.38	ESL 12		
Actinomyces israelii		45.55	0.27	GAL 96		
47772041	**Good identification**					**47772041**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47772043	**Good identification**					**47772043**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47772060	**Low discrimination**					**47772060**
Actinomyces naeslundii		83.15	0.38	ESL 12		
Bifidobacterium breve		16.85	0.12	GAL 96		
47772061	**Good identification**					**47772061**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47772063	**Good identification**					**47772063**
Actinomyces naeslundii		100.0	0.41	ESL 12		
47772100	**Low discrimination**					**47772100**
Actinomyces naeslundii		38.78	0.59	ESL 12		
Bifidobacterium breve		27.15	0.41	GAL 96		
Actinomyces israelii		20.69	0.45			
47772101	**Very good identification**					**47772101**
Actinomyces naeslundii		100.0	0.62	ESL 12		
47772102	**Low discrimination**					**47772102**
Bifidobacterium breve		69.72	0.59	GAL 99		
Actinomyces naeslundii		9.85	0.59			
Actinomyces israelii		7.88	0.48			
47772103	**Good identification**					**47772103**
Actinomyces naeslundii		94.96	0.62	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		4.53	0.24	GAL 96		
Actinomyces israelii		0.51	0.14			
47772110	**Low discrimination**					**47772110**
Bifidobacterium breve		78.40	0.41	GAL 99		
Actinomyces israelii		8.93	0.31	BGL 91		
Actinomyces naeslundii		5.89	0.37			
47772111	**Good identification**					**47772111**
Actinomyces naeslundii		100.0	0.40	ESL 12		
47772112	**Low discrimination**					**47772112**
					GRA	
Bifidobacterium breve		82.47	0.59	GAL 99	99	
Leptotrichia buccalis		10.41	0.24		1	
Bifidobacterium adolescentis		3.04	0.33		99	
47772113	**Low discrimination**					**47772113**
					GRA	
Actinomyces naeslundii		49.50	0.40	ESL 12	99	
Bifidobacterium breve		44.84	0.24	GAL 96	99	
Leptotrichia buccalis		5.66	-.11	MLZ 5	1	
47772120	**Low discrimination**					**47772120**
					SPO	
Clostridium ramosum		86.19	0.53	GAL 99	92	
Bifidobacterium breve		7.11	0.46		1	
Actinomyces naeslundii		5.71	0.59		1	
47772121	**Good identification**					**47772121**
					SPO	
Actinomyces naeslundii		90.09	0.62	ESL 12	1	
Clostridium ramosum		9.15	0.19	GAL 96	92	
Bifidobacterium breve		0.76	0.12		1	
47772122	**Low discrimination**					**47772122**
					SPO	
Clostridium ramosum		80.83	0.64	GAL 99	92	
Bifidobacterium breve		16.86	0.63		1	
Actinomyces naeslundii		1.34	0.59		1	
47772123	**Low discrimination**					**47772123**
					SPO	
Actinomyces naeslundii		67.06	0.62	ESL 12	1	
Clostridium ramosum		27.25	0.30	GAL 96	92	
Bifidobacterium breve		5.68	0.29		1	
47772130	**Low discrimination**					**47772130**
					SPO	
Bifidobacterium breve		77.58	0.46	GAL 99	1	
Clostridium tertium		9.65	0.21	BGL 91	99	
Clostridium ramosum		9.50	0.19		92	
47772131	**Low discrimination**					**47772131**
Actinomyces naeslundii		86.27	0.40	ESL 12		
Bifidobacterium breve		13.73	0.12	GAL 96		
47772132	**Good identification**					**47772132**
					SPO	
Bifidobacterium breve		90.85	0.63	GAL 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		4.40	0.30		92	
Clostridium tertium		3.35	0.29		99	
47772140	**Low discrimination**					**47772140**
Bifidobacterium breve		64.59	0.41	GAL 99		
Actinomyces israelii		19.14	0.38	BGL 91		
Actinomyces naeslundii		16.28	0.46			
47772141	**Good identification**					**47772141**
Actinomyces naeslundii		100.0	0.49	ESL 12		
47772142	**Good identification**					**47772142**
Bifidobacterium breve		92.77	0.59	GAL 99		
Actinomyces israelii		4.08	0.41			
Actinomyces naeslundii		2.31	0.46			
47772143	**Low discrimination**					**47772143**
Actinomyces naeslundii		78.73	0.49	ESL 12		
Bifidobacterium breve		21.27	0.24	GAL 96		
47772150	**Acceptable identification**					**47772150**
					COC	
Bifidobacterium breve		92.84	0.41	GAL 99	1	
Actinomyces israelii		4.11	0.24	BGL 91	1	
Blautia producta		1.81	-.03		99	
47772152	**Very good identification**					**47772152**
Bifidobacterium breve		98.65	0.59	GAL 99		
Actinomyces israelii		0.65	0.27			
Bifidobacterium adolescentis		0.45	0.18			
47772160	**Low discrimination**					**47772160**
					SPO	
Bifidobacterium breve		78.44	0.46	GAL 99	1	
Actinomyces naeslundii		11.12	0.46	BGL 91	1	
Clostridium ramosum		9.60	0.19		92	
47772161	**Good identification**					**47772161**
Actinomyces naeslundii		95.47	0.49	ESL 12		
Bifidobacterium breve		4.53	0.12	GAL 96		
47772162	**Good identification**					**47772162**
					SPO	
Bifidobacterium breve		93.72	0.63	GAL 99	1	
Clostridium ramosum		4.54	0.30		92	
Actinomyces naeslundii		1.31	0.46		1	
47772163	**Low discrimination**					**47772163**
Actinomyces naeslundii		67.56	0.49	ESL 12		
Bifidobacterium breve		32.44	0.29	GAL 96		
47772170	**Good identification**					**47772170**
Bifidobacterium breve		99.26	0.46	GAL 99		
Actinomyces naeslundii		0.74	0.24	BGL 91		
47772172	**Very good identification**					**47772172**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		99.93	0.63	GAL 99		
Actinomyces naeslundii		0.07	0.24			
47772320	**Good identification**					**47772320**
					SPO	
Clostridium ramosum		99.89	0.56	GAL 99	92	
Bifidobacterium breve		0.06	0.12		1	
Actinomyces naeslundii		0.05	0.24		1	
47772322	**Very good identification**					**47772322**
					SPO	
Clostridium ramosum		99.74	0.66	GAL 99	92	
Bifidobacterium breve		0.16	0.29		1	
Lactobacillus acidophilus		0.03	0.06		1	
47772402	**Low discrimination**					**47772402**
Actinomyces israelii		83.32	0.36	GAL 99		
Bifidobacterium adolescentis		7.35	0.12	XYL 92		
Clostridium clostridioforme		4.91	0.05			
47772502	**Low discrimination**					**47772502**
					SPO	
Bifidobacterium adolescentis		59.05	0.46	GAL 99	1	
Actinomyces israelii		27.05	0.46	XYL 87	1	
Clostridium butyricum		8.62	0.17		99	
47772512	**Low discrimination**					**47772512**
Bifidobacterium adolescentis		50.53	0.47	GAL 99		
Bifidobacterium dentium		39.74	0.27	XYL 87		
Clostridium clostridioforme		4.25	0.26			
47772542	**Low discrimination**					**47772542**
Actinomyces israelii		44.59	0.39	GAL 99		
Bifidobacterium adolescentis		30.93	0.30	XYL 92		
Bifidobacterium breve		12.52	0.24			
47772712	**Good identification**					**47772712**
Clostridium clostridioforme		97.16	0.41	MAN 1		
Bifidobacterium adolescentis		1.59	0.13	XYL 91		
Bifidobacterium dentium		1.25	-.07			
47772732	**Good identification**					**47772732**
					GRA	
Clostridium clostridioforme		96.26	0.41	MAN 1	50	
Bifidobacterium dentium		2.20	-.03	XYL 91	99	
Bacteroides ovatus		0.81	-.05		1	
47773000	**Low discrimination**					**47773000**
Actinomyces israelii		56.34	0.69	XYL 92		
Actinomyces naeslundii		35.98	0.74			
Bifidobacterium breve		2.99	0.41			
47773001	**Very good identification**					**47773001**
Actinomyces naeslundii		98.96	0.77	ESL 12		
Actinomyces israelii		1.04	0.34			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
47773002	**Low discrimination**					**47773002**
Actinomyces israelii		50.18	0.72	XYL 92		
Actinomyces naeslundii		21.37	0.74			
Bifidobacterium breve		17.94	0.59			
47773003	**Very good identification**					**47773003**
Actinomyces naeslundii		97.70	0.77	ESL 12		
Actinomyces israelii		1.55	0.37			
Bifidobacterium breve		0.55	0.24			
47773010	**Low discrimination**					**47773010**
Actinomyces israelii		58.84	0.54	XYL 92		
Bifidobacterium breve		20.88	0.41	MLZ 13		
Actinomyces naeslundii		13.24	0.53			
47773011	**Good identification**					**47773011**
Actinomyces naeslundii		97.09	0.56	ESL 12		
Actinomyces israelii		2.91	0.20	MLZ 5		
47773012	**Low discrimination**					**47773012**
					GRA	
Bifidobacterium breve		59.24	0.59	LAC 99	99	
Actinomyces israelii		24.76	0.57		99	
Leptotrichia buccalis		7.48	0.24		1	
47773013	**Low discrimination**					**47773013**
Actinomyces naeslundii		85.77	0.56	ESL 12		
Bifidobacterium breve		9.21	0.24	MLZ 5		
Actinomyces israelii		3.85	0.23			
47773020	**Low discrimination**					**47773020**
					SPO	
Clostridium ramosum		58.02	0.53	LAC 99	92	
Actinomyces naeslundii		32.43	0.74		1	
Bifidobacterium breve		4.79	0.46		1	
47773021	**Very good identification**					**47773021**
					SPO	
Actinomyces naeslundii		98.65	0.77	ESL 12	1	
Clostridium ramosum		1.19	0.19		92	
Bifidobacterium breve		0.10	0.12		1	
47773022	**Low discrimination**					**47773022**
					SPO	
Clostridium ramosum		69.99	0.64	LAC 99	92	
Bifidobacterium breve		14.60	0.63		1	
Actinomyces naeslundii		9.78	0.74		1	
47773023	**Good identification**					**47773023**
					SPO	
Actinomyces naeslundii		94.22	0.77	ESL 12	1	
Clostridium ramosum		4.54	0.30		92	
Bifidobacterium breve		0.95	0.29		1	
47773030	**Low discrimination**					**47773030**
					SPO	
Bifidobacterium breve		57.56	0.46	LAC 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		20.53	0.53	BGL 91	1	
Clostridium tertium		7.16	0.21		99	
47773031	**Good identification**					**47773031**
Actinomyces naeslundii		98.15	0.56	ESL 12		
Bifidobacterium breve		1.85	0.12	MLZ 5		
47773032	**Low discrimination**					**47773032**
					SPO	
Bifidobacterium breve		86.93	0.63	LAC 99	1	
Clostridium ramosum		4.21	0.30		92	
Clostridium tertium		3.21	0.29		99	
47773033	**Low discrimination**					**47773033**
Actinomyces naeslundii		83.97	0.56	ESL 12		
Bifidobacterium breve		16.03	0.29	MLZ 5		
47773040	**Low discrimination**					**47773040**
Actinomyces israelii		69.91	0.62	XYL 92		
Actinomyces naeslundii		20.26	0.62			
Bifidobacterium breve		9.53	0.41			
47773041	**Very good identification**					**47773041**
Actinomyces naeslundii		97.73	0.65	ESL 12		
Actinomyces israelii		2.27	0.27			
47773042	**Low discrimination**					**47773042**
Actinomyces israelii		46.31	0.65	XYL 92		
Bifidobacterium breve		42.57	0.59			
Actinomyces naeslundii		8.95	0.62			
47773043	**Good identification**					**47773043**
Actinomyces naeslundii		93.73	0.65	ESL 12		
Actinomyces israelii		3.27	0.30			
Bifidobacterium breve		3.00	0.24			
47773050	**Low discrimination**					**47773050**
Actinomyces israelii		49.20	0.47	XYL 92		
Bifidobacterium breve		44.90	0.41	MLZ 13		
Actinomyces naeslundii		5.02	0.40			
47773051	**Acceptable identification**					**47773051**
Actinomyces naeslundii		93.81	0.43	ESL 12		
Actinomyces israelii		6.19	0.13	MLZ 5		
47773052	**Low discrimination**					**47773052**
Bifidobacterium breve		84.79	0.59	LAC 99		
Actinomyces israelii		13.78	0.50			
Actinomyces naeslundii		0.94	0.40			
47773053	**Low discrimination**					**47773053**
Actinomyces naeslundii		58.56	0.43	ESL 12		
Bifidobacterium breve		35.64	0.24	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		5.80	0.16			
47773060	**Low discrimination**					**47773060**
Actinomyces naeslundii		46.32	0.62	ESL 12		
Bifidobacterium breve		38.74	0.46			
Actinomyces israelii		10.20	0.41			
47773061	**Very good identification**					**47773061**
Actinomyces naeslundii		99.44	0.65	ESL 12		
Bifidobacterium breve		0.56	0.12			
47773062	**Low discrimination**					**47773062**
					SPO	
Bifidobacterium breve		81.08	0.63	LAC 99	1	
Actinomyces naeslundii		9.59	0.62		1	
Clostridium ramosum		3.93	0.30		92	
47773063	**Good identification**					**47773063**
Actinomyces naeslundii		94.61	0.65	ESL 12		
Bifidobacterium breve		5.39	0.29			
47773070	**Acceptable identification**					**47773070**
Bifidobacterium breve		90.72	0.46	LAC 99		
Actinomyces naeslundii		5.71	0.40	BGL 91		
Actinomyces israelii		3.57	0.27			
47773071	**Acceptable identification**					**47773071**
Actinomyces naeslundii		90.34	0.43	ESL 12		
Bifidobacterium breve		9.66	0.12	MLZ 5		
47773072	**Very good identification**					**47773072**
Bifidobacterium breve		98.81	0.63	LAC 99		
Actinomyces naeslundii		0.62	0.40			
Actinomyces israelii		0.58	0.30			
47773100	**Low discrimination**					**47773100**
Bifidobacterium breve		40.50	0.76	BGL 91		
Actinomyces israelii		30.85	0.79			
Actinomyces naeslundii		14.02	0.82			
47773101	**Very good identification**					**47773101**
Actinomyces naeslundii		96.07	0.85	ESL 12		
Bifidobacterium breve		1.87	0.41			
Actinomyces israelii		1.42	0.45			
47773102	**Low discrimination**					**47773102**
					SPO	
Bifidobacterium breve		75.49	0.93		1	
Actinomyces israelii		8.53	0.82		1	
Clostridium ramosum		6.50	0.64		92	
47773103	**Low discrimination**					**47773103**
Actinomyces naeslundii		79.77	0.85	ESL 12		
Bifidobacterium breve		15.69	0.59			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		1.77	0.48			
47773110	**Low discrimination**					**47773110**
					SPO	
Bifidobacterium breve		81.22	0.76	BGL 91	1	
Actinomyces israelii		9.25	0.65		1	
Clostridium tertium		5.99	0.47		99	
47773111	**Low discrimination**					**47773111**
Actinomyces naeslundii		69.27	0.63	ESL 12		
Bifidobacterium breve		25.60	0.41	MLZ 5		
Actinomyces israelii		2.91	0.31			
47773112	**Low discrimination**					**47773112**
					GRA	
Bifidobacterium breve		82.84	0.93		99	
Leptotrichia buccalis		10.46	0.58		1	
Bifidobacterium adolescentis		3.05	0.67		99	
47773113	**Low discrimination**					**47773113**
					GRA	
Bifidobacterium breve		67.96	0.59	URE 1	99	
Actinomyces naeslundii		18.19	0.63		99	
Leptotrichia buccalis		8.58	0.24		1	
47773120	**Good identification**					**47773120**
					SPO	
Clostridium ramosum		90.07	0.88		92	
Bifidobacterium breve		7.43	0.80		1	
Actinomyces naeslundii		1.45	0.82		1	
47773121	**Low discrimination**					**47773121**
					SPO	
Actinomyces naeslundii		68.58	0.85	ESL 12	1	
Clostridium ramosum		28.74	0.53		92	
Bifidobacterium breve		2.37	0.46		1	
47773122	**Low discrimination**					**47773122**
					SPO	
Clostridium ramosum		81.68	0.98		92	
Bifidobacterium breve		17.03	0.97		1	
Lactobacillus acidophilus		0.42	0.60		1	
47773123	**Low discrimination**					**47773123**
					SPO	
Clostridium ramosum		55.05	0.64	URE 1	92	
Actinomyces naeslundii		32.84	0.85		1	
Bifidobacterium breve		11.48	0.63		1	
47773130	**Low discrimination**					**47773130**
					SPO	
Bifidobacterium breve		79.09	0.80	BGL 91	1	
Clostridium tertium		9.83	0.55		99	
Clostridium ramosum		9.68	0.53		92	
47773131	**Low discrimination**					**47773131**
					SPO	
Actinomyces naeslundii		54.97	0.63	ESL 12	1	
Bifidobacterium breve		36.11	0.46	MLZ 5	1	
Clostridium tertium		4.49	0.21		99	
47773132	**Good identification**					**47773132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Bifidobacterium breve		91.12	0.97		1	
Clostridium ramosum		4.41	0.64		92	
Clostridium tertium		3.36	0.63		99	
47773133	**Low discrimination**					**47773133**
					SPO	
Bifidobacterium breve		80.24	0.63	URE 1	1	
Actinomyces naeslundii		12.08	0.63		1	
Clostridium ramosum		3.89	0.30		92	
47773140	**Low discrimination**					**47773140**
Bifidobacterium breve		72.74	0.76	BGL 91		
Actinomyces israelii		21.55	0.72			
Actinomyces naeslundii		4.44	0.69			
47773141	**Low discrimination**					**47773141**
Actinomyces naeslundii		87.50	0.72	ESL 12		
Bifidobacterium breve		9.64	0.41			
Actinomyces israelii		2.86	0.38			
47773142	**Good identification**					**47773142**
Bifidobacterium breve		94.30	0.93			
Actinomyces israelii		4.15	0.75			
Actinomyces naeslundii		0.57	0.69			
47773143	**Low discrimination**					**47773143**
Bifidobacterium breve		51.27	0.59	URE 1		
Actinomyces naeslundii		46.01	0.72			
Actinomyces israelii		2.25	0.41			
47773150	**Good identification**					**47773150**
					COC	
Bifidobacterium breve		93.57	0.76	BGL 91	1	
Actinomyces israelii		4.14	0.58		1	
Blautia producta		1.83	0.32		99	
47773151	**Low discrimination**					**47773151**
Bifidobacterium breve		64.88	0.41	URE 1		
Actinomyces naeslundii		30.98	0.50	BGL 91		
Actinomyces israelii		2.87	0.24			
47773152	**Very good identification**					**47773152**
Bifidobacterium breve		98.69	0.93			
Actinomyces israelii		0.65	0.61			
Bifidobacterium adolescentis		0.45	0.52			
47773153	**Good identification**					**47773153**
Bifidobacterium breve		94.37	0.59	URE 1		
Actinomyces naeslundii		4.46	0.50			
Actinomyces israelii		0.62	0.27			
47773160	**Low discrimination**					**47773160**
					SPO	
Bifidobacterium breve		85.34	0.80	BGL 91	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		10.45	0.53		92	
Actinomyces naeslundii		2.93	0.69		1	
47773161	**Low discrimination**					**47773161**
					SPO	
Actinomyces naeslundii		81.83	0.72	ESL 12	1	
Bifidobacterium breve		16.03	0.46		1	
Clostridium ramosum		1.96	0.19		92	
47773162	**Good identification**					**47773162**
					SPO	
Bifidobacterium breve		94.62	0.97		1	
Clostridium ramosum		4.58	0.64		92	
Actinomyces naeslundii		0.32	0.69		1	
47773163	**Low discrimination**					**47773163**
					SPO	
Bifidobacterium breve		64.19	0.63	URE 1	1	
Actinomyces naeslundii		32.40	0.72		1	
Clostridium ramosum		3.11	0.30		92	
47773170	**Very good identification**					**47773170**
Bifidobacterium breve		99.29	0.80	BGL 91		
Actinomyces naeslundii		0.18	0.47			
Actinomyces israelii		0.16	0.37			
47773171	**Low discrimination**					**47773171**
Bifidobacterium breve		78.83	0.46	URE 1		
Actinomyces naeslundii		21.18	0.50	BGL 91		
47773172	**Excellent identification**					**47773172**
					SPO	
Bifidobacterium breve		99.86	0.97		1	
Clostridium ramosum		0.05	0.30		92	
Clostridium tertium		0.04	0.29		99	
47773173	**Very good identification**					**47773173**
Bifidobacterium breve		97.41	0.63	URE 1		
Actinomyces naeslundii		2.59	0.50			
47773200	**Low discrimination**					**47773200**
					SPO	
Actinomyces israelii		77.58	0.49	XYL 92	1	
Clostridium ramosum		15.77	0.21	RHA 7	92	
Actinomyces naeslundii		6.65	0.40		1	
47773201	**Acceptable identification**					**47773201**
Actinomyces naeslundii		92.72	0.43	ESL 12		
Actinomyces israelii		7.29	0.15	RHA 1		
47773202	**Low discrimination**					**47773202**
					SPO	
Actinomyces israelii		55.71	0.52	XYL 92	1	
Clostridium ramosum		30.20	0.32	RHA 7	92	
Lactobacillus acidophilus		7.44	0.22		1	
47773203	**Low discrimination**					**47773203**
					SPO	
Actinomyces naeslundii		84.62	0.43	ESL 12	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		9.97	0.18	RHA 1	1		
Clostridium ramosum		5.41	-.03		92		
47773210	**Good identification**						**47773210**
Actinomyces israelii		100.0	0.35	XYL 92			
47773212	**Low discrimination**						**47773212**
Actinomyces israelii		59.70	0.38	XYL 92			
Bifidobacterium breve		19.16	0.24	RHA 7			
Clostridium clostridioforme		16.53	0.19	MLZ 13			
47773220	**Good identification**						**47773220**
					SPO		
Clostridium ramosum		99.16	0.56	LAC 99	92		
Actinomyces naeslundii		0.42	0.40		1		
Actinomyces israelii		0.31	0.29		1		
47773222	**Very good identification**						**47773222**
					SPO	GRA	
Clostridium ramosum		99.14	0.66	LAC 99	92	50	
Lactobacillus acidophilus		0.25	0.22		1	99	
Prevotella oralis		0.18	0.19		1	1	
47773240	**Good identification**						**47773240**
Actinomyces israelii		96.26	0.42	XYL 92			
Actinomyces naeslundii		3.74	0.27	RHA 7			
47773242	**Low discrimination**						**47773242**
Actinomyces israelii		82.93	0.45	XYL 92			
Bifidobacterium breve		10.23	0.24	RHA 7			
Lactobacillus acidophilus		3.52	0.07				
47773300	**Low discrimination**						**47773300**
					SPO		
Clostridium ramosum		79.52	0.56	NAG 99	92		
Actinomyces israelii		15.80	0.60		1		
Bifidobacterium breve		2.78	0.41		1		
47773301	**Low discrimination**						**47773301**
					SPO		
Actinomyces naeslundii		60.03	0.51	ESL 12	1		
Clostridium ramosum		33.35	0.21	RHA 1	92		
Actinomyces israelii		6.63	0.25		1		
47773302	**Low discrimination**						**47773302**
					SPO		
Clostridium ramosum		83.20	0.66	NAG 99	92		
Bifidobacterium breve		7.36	0.59		1		
Actinomyces israelii		6.20	0.63		1		
47773310	**Low discrimination**						**47773310**
					SPO		
Bifidobacterium breve		43.96	0.41	RHA 1	1		
Actinomyces israelii		37.29	0.46	BGL 91	1		
Clostridium ramosum		12.68	0.21		92		
47773312	**Low discrimination**						**47773312**
					GRA		
Bifidobacterium breve		68.44	0.59	RHA 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		8.64	0.24		1		
Actinomyces israelii		8.61	0.49		99		
47773320	**Very good identification**						**47773320**
					SPO		
Clostridium ramosum		99.90	0.90		92		
Bifidobacterium breve		0.06	0.46		1		
Actinomyces naeslundii		0.01	0.48		1		
47773321	**Good identification**						**47773321**
					SPO		
Clostridium ramosum		98.15	0.56	URE 1	92		
Actinomyces naeslundii		1.79	0.51		1		
Bifidobacterium breve		0.06	0.12		1		
47773322	**Excellent identification**						**47773322**
					SPO		
Clostridium ramosum		99.76	1.00		92		
Bifidobacterium breve		0.16	0.63		1		
Lactobacillus acidophilus		0.03	0.40		1		
47773323	**Very good identification**						**47773323**
					SPO		
Clostridium ramosum		99.32	0.66	URE 1	92		
Actinomyces naeslundii		0.45	0.51		1		
Bifidobacterium breve		0.16	0.29		1		
47773330	**Acceptable identification**						**47773330**
					SPO		
Clostridium ramosum		93.24	0.56	MLZ 1	92		
Bifidobacterium breve		5.80	0.46		1		
Clostridium tertium		0.72	0.21		99		
47773332	**Low discrimination**						**47773332**
					SPO	GRA	
Clostridium ramosum		85.09	0.66	MLZ 1	92	50	
Bifidobacterium breve		13.39	0.63		1	99	
Parabacteroides distasonis		0.52	0.40		1	1	
47773340	**Low discrimination**						**47773340**
					COC		
Actinomyces israelii		52.26	0.53	XYL 92	1		
Bifidobacterium breve		23.67	0.41	RHA 7	1		
Blautia producta		15.26	0.24		99		
47773341	**Low discrimination**						**47773341**
Actinomyces naeslundii		80.43	0.38	ESL 12			
Actinomyces israelii		19.57	0.18	RHA 1			
47773342	**Low discrimination**						**47773342**
					SPO		
Bifidobacterium breve		66.73	0.59	RHA 1	1		
Actinomyces israelii		21.86	0.56		1		
Clostridium ramosum		7.62	0.32		92		
47773352	**Good identification**						**47773352**
					COC		
Bifidobacterium breve		92.84	0.59	RHA 1	1		
Actinomyces israelii		4.54	0.42		1		
Blautia producta		1.97	0.15		99		
47773360	**Acceptable identification**						**47773360**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium ramosum		93.30	0.56	SOR 1	92		
Bifidobacterium breve		5.81	0.46		1		
Actinomyces israelii		0.46	0.32		1		
47773362	**Low discrimination**						**47773362**
					SPO		
Clostridium ramosum		85.94	0.66	SOR 1	92		
Bifidobacterium breve		13.52	0.63		1		
Lactobacillus acidophilus		0.30	0.25		1		
47773370	**Low discrimination**						**47773370**
					COC	SPO	
Bifidobacterium breve		77.22	0.46	RHA 1	1	1	
Clostridium ramosum		12.53	0.21	BGL 91	1	92	
Blautia producta		9.34	0.15		99	1	
47773372	**Good identification**						**47773372**
					SPO		
Bifidobacterium breve		93.70	0.63	RHA 1	1		
Clostridium ramosum		6.02	0.32		92		
Actinomyces israelii		0.17	0.21		1		
47773400	**Low discrimination**						**47773400**
					SPO		
Actinomyces israelii		82.30	0.67	XYL 92	1		
Clostridium butyricum		16.85	0.34		99		
Actinomyces naeslundii		0.65	0.40		1		
47773401	**Low discrimination**						**47773401**
					SPO		
Actinomyces naeslundii		49.29	0.43	ESL 12	1		
Actinomyces israelii		42.09	0.33	ARA 1	1		
Clostridium butyricum		8.62	0.00		99		
47773402	**Low discrimination**						**47773402**
					SPO		
Actinomyces israelii		86.51	0.70	XYL 92	1		
Bifidobacterium adolescentis		7.63	0.46		1		
Clostridium butyricum		4.59	0.27		99		
47773403	**Low discrimination**						**47773403**
Actinomyces israelii		52.03	0.36	XYL 92			
Actinomyces naeslundii		40.62	0.43	URE 1			
Bifidobacterium adolescentis		4.59	0.12				
47773410	**Low discrimination**						**47773410**
					SPO		
Actinomyces israelii		85.06	0.53	XYL 92	1		
Bifidobacterium dentium		7.16	0.10	MLZ 13	1		
Clostridium butyricum		6.13	0.12		99		
47773412	**Low discrimination**						**47773412**
Actinomyces israelii		41.75	0.56	XYL 92			
Bifidobacterium adolescentis		30.12	0.47	MLZ 13			
Bifidobacterium dentium		23.69	0.27				
47773420	**Low discrimination**						**47773420**
					SPO		
Actinomyces israelii		60.91	0.47	XYL 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		16.99	0.16	NAG 6	99		
Clostridium ramosum		13.46	0.19		92		
47773421	**Good identification**						**47773421**
Actinomyces naeslundii		100.0	0.43	ESL 12			
47773422	**Low discrimination**						**47773422**
					SPO		
Actinomyces israelii		51.06	0.50	XYL 92	1		
Clostridium ramosum		30.09	0.30	NAG 6	92		
Bifidobacterium breve		6.28	0.29		1		
47773423	**Acceptable identification**						**47773423**
Actinomyces naeslundii		92.44	0.43	ESL 12			
Actinomyces israelii		7.56	0.15	ARA 1			
47773440	**Low discrimination**						**47773440**
					COC	SPO	
Actinomyces israelii		86.44	0.60	XYL 92	1	1	
Blautia producta		9.29	0.24		99	1	
Clostridium butyricum		3.96	0.16		1	99	
47773442	**Good identification**						**47773442**
					SPO		
Actinomyces israelii		94.20	0.63	XYL 92	1		
Bifidobacterium adolescentis		2.64	0.30		1		
Clostridium butyricum		1.12	0.09		99		
47773452	**Low discrimination**						**47773452**
					COC		
Actinomyces israelii		67.58	0.49	XYL 92	1		
Bifidobacterium adolescentis		15.50	0.32	MLZ 13	1		
Blautia producta		10.80	0.15		99		
47773460	**Low discrimination**						**47773460**
					COC	SPO	
Actinomyces israelii		73.94	0.40	XYL 92	1	1	
Blautia producta		13.83	0.07	NAG 6	99	1	
Clostridium butyricum		4.61	-.02		1	99	
47773462	**Low discrimination**						**47773462**
Actinomyces israelii		73.89	0.43	XYL 92			
Bifidobacterium breve		23.35	0.29	NAG 6			
Actinomyces naeslundii		2.76	0.27				
47773500	**Low discrimination**						**47773500**
					SPO		
Clostridium butyricum		53.24	0.58	MAN 10	99		
Actinomyces israelii		43.34	0.77	XYL 97	1		
Bifidobacterium adolescentis		1.43	0.46		1		
47773502	**Low discrimination**						**47773502**
					SPO		
Bifidobacterium adolescentis		59.74	0.80	XYL 87	1		
Actinomyces israelii		27.37	0.80		1		
Clostridium butyricum		8.72	0.51		99		
47773503	**Low discrimination**						**47773503**
Bifidobacterium adolescentis		52.13	0.46	XYL 87			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		23.88	0.46	URE 1			
Actinomyces naeslundii		13.27	0.51				
47773510	**Low discrimination**						**47773510**
					SPO		
Bifidobacterium dentium		50.44	0.44	XYL 99	1		
Actinomyces israelii		24.21	0.63	BGL 91	1		
Clostridium butyricum		10.48	0.36		99		
47773512	**Low discrimination**						**47773512**
Bifidobacterium adolescentis		52.65	0.81	XYL 87			
Bifidobacterium dentium		41.41	0.62				
Actinomyces israelii		2.95	0.66				
47773513	**Low discrimination**						**47773513**
Bifidobacterium adolescentis		52.47	0.47	XYL 87			
Bifidobacterium dentium		41.27	0.27	URE 1			
Actinomyces israelii		2.94	0.32				
47773520	**Low discrimination**						**47773520**
					SPO		
Clostridium ramosum		61.71	0.53	ARA 1	92		
Clostridium butyricum		18.88	0.40		99		
Actinomyces israelii		11.28	0.57		1		
47773521	**Low discrimination**						**47773521**
					SPO		
Actinomyces naeslundii		60.30	0.51	ESL 12	1		
Clostridium ramosum		25.27	0.19	ARA 1	92		
Clostridium butyricum		7.73	0.06		99		
47773522	**Low discrimination**						**47773522**
					SPO		
Clostridium ramosum		71.87	0.64	ARA 1	92		
Bifidobacterium breve		14.99	0.63		1		
Actinomyces israelii		4.93	0.60		1		
47773530	**Acceptable identification**						**47773530**
Bifidobacterium dentium		90.84	0.49	XYL 99			
Bifidobacterium breve		4.73	0.46	BGL 91			
Actinomyces israelii		1.57	0.43				
47773532	**Good identification**						**47773532**
Bifidobacterium dentium		93.49	0.66	XYL 99			
Bifidobacterium breve		4.87	0.63				
Bifidobacterium adolescentis		0.68	0.47				
47773540	**Low discrimination**						**47773540**
					COC	SPO	
Blautia producta		66.40	0.58	XYL 99	99	1	
Actinomyces israelii		24.97	0.70		1	1	
Clostridium butyricum		6.86	0.40		1	99	
47773542	**Low discrimination**						**47773542**
Actinomyces israelii		44.49	0.73	XYL 92			
Bifidobacterium adolescentis		30.86	0.64				
Bifidobacterium breve		12.50	0.59				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47773543	**Low discrimination**					**47773543**
Actinomyces israelii		40.08	0.39	XYL 92		
Bifidobacterium adolescentis		27.81	0.30	URE 1		
Bifidobacterium breve		11.26	0.24			
47773550	**Very good identification**					**47773550**
					COC	
Blautia producta		97.18	0.66	XYL 99	99	
Actinomyces israelii		1.82	0.56		1	
Bifidobacterium breve		0.51	0.41		1	
47773552	**Low discrimination**					**47773552**
					COC	
Bifidobacterium adolescentis		43.97	0.66	XYL 87	1	
Blautia producta		30.65	0.50	SOR 11	99	
Bifidobacterium breve		14.57	0.59		1	
47773560	**Low discrimination**					**47773560**
					COC	
Blautia producta		62.48	0.41	XYL 99	99	
Bifidobacterium breve		15.66	0.46	NAG 10	1	
Actinomyces israelii		13.49	0.50		1	
47773561	**Low discrimination**					**47773561**
					COC	
Actinomyces naeslundii		46.59	0.38	ESL 12	1	
Blautia producta		36.42	0.07	ARA 1	99	
Bifidobacterium breve		9.13	0.12		1	
47773562	**Low discrimination**					**47773562**
					SPO	
Bifidobacterium breve		79.69	0.63	ARA 1	1	
Actinomyces israelii		10.19	0.53		1	
Clostridium ramosum		3.86	0.30		92	
47773570	**Low discrimination**					**47773570**
					COC	
Blautia producta		89.81	0.50	XYL 99	99	
Bifidobacterium breve		7.50	0.46	NAG 10	1	
Bifidobacterium dentium		1.46	0.15		1	
47773572	**Low discrimination**					**47773572**
					COC	
Bifidobacterium breve		73.40	0.63	ARA 1	1	
Bifidobacterium dentium		14.24	0.32		1	
Blautia producta		9.65	0.33		99	
47773600	**Acceptable identification**					**47773600**
					SPO	
Actinomyces israelii		92.24	0.48	XYL 92	1	
Clostridium butyricum		7.76	0.08	RHA 7	99	
47773602	**Low discrimination**					**47773602**
					SPO	
Actinomyces israelii		82.35	0.51	XYL 92	1	
Clostridium clostridioforme		14.32	0.28	RHA 7	66	
Clostridium butyricum		1.80	0.01		99	
47773612	**Low discrimination**					**47773612**
Clostridium clostridioforme		81.89	0.40	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		15.44	0.37	XYL 91			
Bifidobacterium adolescentis		1.50	0.13				
47773632	**Acceptable identification**						**47773632**
Clostridium clostridioforme		94.75	0.40	MAN 1			
Bifidobacterium dentium		2.42	-.03	XYL 91			
Actinomyces israelii		1.14	0.16				
47773642	**Acceptable identification**						**47773642**
					COC		
Actinomyces israelii		90.44	0.44	XYL 92	1		
Blautia producta		9.57	0.07	RHA 7	99		
47773700	**Low discrimination**						**47773700**
					GRA	SPO	
Actinomyces israelii		53.58	0.58	XYL 92	99	1	
Clostridium butyricum		27.05	0.32	RHA 7	99	99	
Mitsuokella multacida		8.37	0.32		1	1	
47773702	**Low discrimination**						**47773702**
					SPO		
Actinomyces israelii		56.12	0.61	XYL 92	1		
Bifidobacterium adolescentis		16.44	0.46	RHA 7	1		
Clostridium ramosum		9.30	0.32		92		
47773712	**Low discrimination**						**47773712**
Clostridium clostridioforme		43.81	0.49	MAN 1			
Bifidobacterium adolescentis		23.65	0.47	XYL 91			
Bifidobacterium dentium		18.60	0.27				
47773720	**Good identification**						**47773720**
					SPO		
Clostridium ramosum		97.31	0.56	ARA 1	92		
Actinomyces israelii		1.01	0.38		1		
Clostridium butyricum		0.70	0.14		99		
47773722	**Very good identification**						**47773722**
					SPO		
Clostridium ramosum		96.13	0.66	ARA 1	92		
Bacteroides ovatus		1.62	0.32		1		
Clostridium clostridioforme		0.85	0.37		66		
47773732	**Low discrimination**						**47773732**
					GRA		
Clostridium clostridioforme		40.99	0.49	MAN 1	50		
Bifidobacterium dentium		30.94	0.32	XYL 91	99		
Bacteroides ovatus		11.37	0.29		1		
47773740	**Very good identification**						**47773740**
					COC	GRA	
Blautia producta		95.89	0.58	XYL 99	99	99	
Actinomyces israelii		2.71	0.51		1	99	
Mitsuokella multacida		1.09	0.32		1	1	
47773742	**Low discrimination**						**47773742**
					COC		
Blautia producta		68.67	0.41	XYL 99	99		
Actinomyces israelii		26.24	0.54	BGL 10	1		
Bifidobacterium adolescentis		2.44	0.30		1		
47773750	**Very good identification**						**47773750**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Blautia producta		99.73	0.66	XYL 99	99	99		
Actinomyces israelii		0.14	0.37		1	99		
Mitsuokella multacida		0.13	0.24		1	1		
47773752	**Good identification**							**47773752**
					COC			
Blautia producta		95.90	0.50	XYL 99	99			
Actinomyces israelii		1.83	0.40	BGL 10	1			
Bifidobacterium adolescentis		1.39	0.32		1			
47773760	**Acceptable identification**							**47773760**
					COC	SPO	GRA	
Blautia producta		91.95	0.41	XYL 99	99	1	99	
Clostridium ramosum		3.74	0.21	NAG 10	1	92	50	
Mitsuokella multacida		2.35	0.21		1	1	1	
47773770	**Good identification**							**47773770**
					COC	GRA		
Blautia producta		99.55	0.50	XYL 99	99	99		
Mitsuokella multacida		0.28	0.13	NAG 10	1	1		
Actinomyces israelii		0.08	0.17		1	99		
47774002	**Low discrimination**							**47774002**
					SPO			
Clostridium clostridioforme		45.38	0.36	RHA 88	66			
Actinomyces israelii		29.91	0.43	ARA 94	1			
Clostridium butyricum		20.69	0.19		99			
47774012	**Good identification**							**47774012**
Clostridium clostridioforme		96.36	0.48	RHA 88				
Actinomyces israelii		2.08	0.29	ARA 94				
Bifidobacterium adolescentis		0.74	0.15					
47774022	**Low discrimination**							**47774022**
Clostridium clostridioforme		83.63	0.36	RHA 88				
Prevotella oralis		5.78	0.09	ARA 94				
Parabacteroides distasonis		3.76	0.19					
47774032	**Good identification**							**47774032**
Clostridium clostridioforme		99.47	0.48	RHA 88				
Parabacteroides distasonis		0.53	0.14	ARA 94				
47774042	**Low discrimination**							**47774042**
					SPO			
Actinomyces israelii		86.60	0.36	GAL 99	1			
Clostridium butyricum		13.40	0.01		99			
47774100	**Low discrimination**							**47774100**
					GRA	SPO		
Clostridium butyricum		87.44	0.50	GAL 99	99	99		
Actinomyces israelii		5.46	0.50	ARA 90	99	1		
Fusobacterium mortiferum		3.91	0.45		1	1		
47774102	**Low discrimination**							**47774102**
					SPO			
Clostridium butyricum		53.64	0.43	GAL 99	99			
Clostridium clostridioforme		16.41	0.45	ARA 90	66			
Bifidobacterium adolescentis		13.90	0.47		1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47774112	**Low discrimination**					**47774112**
					GRA	
Clostridium clostridioforme		71.60	0.57	RHA 88	50	
Bifidobacterium adolescentis		16.27	0.49	ARA 94	99	
Leptotrichia buccalis		6.81	0.24		1	
47774122	**Low discrimination**					**47774122**
Parabacteroides distasonis		42.00	0.53	GAL 99		
Clostridium clostridioforme		31.57	0.45			
Prevotella oralis		10.21	0.30			
47774132	**Low discrimination**					**47774132**
					GRA	
Clostridium clostridioforme		83.95	0.57	RHA 88	50	
Parabacteroides distasonis		13.20	0.49	ARA 94	1	
Bifidobacterium breve		1.14	0.29		99	
47774142	**Low discrimination**					**47774142**
					SPO	
Actinomyces israelii		41.38	0.46	GAL 99	1	
Clostridium butyricum		38.42	0.24		99	
Bifidobacterium adolescentis		14.15	0.32		1	
47774202	**Good identification**					**47774202**
					SPO	
Clostridium clostridioforme		99.14	0.51	ARA 94	66	
Actinomyces israelii		0.67	0.23		1	
Clostridium butyricum		0.19	-.07		99	
47774212	**Very good identification**					**47774212**
Clostridium clostridioforme		100.0	0.62	ARA 94		
47774222	**Good identification**					**47774222**
Clostridium clostridioforme		97.81	0.51	ARA 94		
Parabacteroides distasonis		1.80	0.27			
Prevotella oralis		0.40	0.03			
47774232	**Very good identification**					**47774232**
Clostridium clostridioforme		99.78	0.62	ARA 94		
Parabacteroides distasonis		0.22	0.23			
47774302	**Very good identification**					**47774302**
					SPO	
Clostridium clostridioforme		97.13	0.60	ARA 94	66	
Clostridium butyricum		1.34	0.17		99	
Actinomyces israelii		0.79	0.34		1	
47774310	**Acceptable identification**					**47774310**
					SPO	
Clostridium clostridioforme		94.52	0.37	ARA 94	66	
Clostridium butyricum		3.83	0.02	BGL 99	99	
Actinomyces israelii		1.65	0.17		1	
47774312	**Very good identification**					**47774312**
					GRA	
Clostridium clostridioforme		99.85	0.71	ARA 94	50	
Parabacteroides distasonis		0.07	0.23		1	
Actinomyces israelii		0.03	0.20		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
47774313	**Good identification**					**47774313**
Clostridium clostridioforme		100.0	0.37	ARA 94		
47774322	**Low discrimination**					**47774322**
Clostridium clostridioforme		63.19	0.60	ARA 94		
Parabacteroides distasonis		34.39	0.61			
Prevotella oralis		1.19	0.24			
47774330	**Acceptable identification**					**47774330**
Clostridium clostridioforme		93.96	0.37	ARA 94		
Parabacteroides distasonis		6.04	0.23	BGL 99		
47774332	**Good identification**					**47774332**
Clostridium clostridioforme		93.85	0.71	ARA 94		
Parabacteroides distasonis		6.04	0.57			
Bacteroides ovatus		0.11	0.11			
47774333	**Acceptable identification**					**47774333**
Clostridium clostridioforme		93.96	0.37	ARA 94		
Parabacteroides distasonis		6.04	0.23	URE 1		
47774352	**Good identification**					**47774352**
Clostridium clostridioforme		100.0	0.37	ARA 94		
47774372	**Acceptable identification**					**47774372**
Clostridium clostridioforme		93.96	0.37	ARA 94		
Parabacteroides distasonis		6.04	0.23	SOR 1		
47774400	**Good identification**					**47774400**
					SPO	
Clostridium butyricum		95.32	0.42	GAL 99	99	
Actinomyces israelii		3.25	0.38	LAC 96	1	
Clostridium clostridioforme		1.43	0.23		66	
47774402	**Low discrimination**					**47774402**
					SPO	
Clostridium clostridioforme		76.42	0.57	RHA 88	66	
Clostridium butyricum		20.02	0.35		99	
Actinomyces israelii		2.63	0.41		1	
47774412	**Very good identification**					**47774412**
					SPO	
Clostridium clostridioforme		99.23	0.68	RHA 88	66	
Bifidobacterium adolescentis		0.33	0.29		1	
Clostridium butyricum		0.30	0.13		99	
47774422	**Very good identification**					**47774422**
					SPO	
Clostridium clostridioforme		97.40	0.57	RHA 88	66	
Clostridium butyricum		2.22	0.17		99	
Actinomyces israelii		0.21	0.21		1	
47774432	**Very good identification**					**47774432**
					SPO	
Clostridium clostridioforme		99.90	0.68	RHA 88	66	
Bifidobacterium dentium		0.06	-.03		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		0.03	-.05		99	
47774500	**Very good identification**					**47774500**
					SPO	
Clostridium butyricum		99.08	0.66	GAL 99	99	
Actinomyces israelii		0.56	0.48		1	
Clostridium clostridioforme		0.21	0.32		66	
47774502	**Low discrimination**					**47774502**
					SPO	
Clostridium butyricum		57.21	0.59	GAL 99	99	
Clostridium clostridioforme		30.47	0.66		66	
Bifidobacterium adolescentis		11.02	0.61		1	
47774510	**Low discrimination**					**47774510**
					SPO	
Clostridium butyricum		80.79	0.44	GAL 99	99	
Clostridium clostridioforme		14.63	0.43	MLZ 5	66	
Bifidobacterium adolescentis		1.42	0.29		1	
47774512	**Low discrimination**					**47774512**
					SPO	
Clostridium clostridioforme		88.53	0.77	RHA 88	66	
Bifidobacterium adolescentis		8.59	0.63		1	
Clostridium butyricum		1.92	0.37		99	
47774513	**Low discrimination**					**47774513**
					SPO	
Clostridium clostridioforme		88.65	0.43	RHA 88	66	
Bifidobacterium adolescentis		8.60	0.29	URE 1	1	
Clostridium butyricum		1.92	0.03		99	
47774520	**Good identification**					**47774520**
					SPO	
Clostridium butyricum		96.86	0.48	GAL 99	99	
Clostridium clostridioforme		2.33	0.32	NAG 8	66	
Actinomyces israelii		0.40	0.28		1	
47774522	**Low discrimination**					**47774522**
					GRA SPO	
Clostridium clostridioforme		81.28	0.66	RHA 88	50 66	
Clostridium butyricum		13.27	0.41		99 99	
Parabacteroides distasonis		4.23	0.49		1 1	
47774530	**Low discrimination**					**47774530**
					SPO	
Clostridium clostridioforme		60.20	0.43	RHA 88	66	
Clostridium butyricum		28.91	0.26	BGL 99	99	
Bifidobacterium dentium		9.78	0.15		1	
47774532	**Very good identification**					**47774532**
					GRA	
Clostridium clostridioforme		97.40	0.77	RHA 88	50	
Bifidobacterium dentium		1.62	0.32		99	
Parabacteroides distasonis		0.60	0.45		1	
47774533	**Good identification**					**47774533**
					GRA	
Clostridium clostridioforme		97.78	0.43	RHA 88	50	
Bifidobacterium dentium		1.62	-.03	URE 1	99	
Parabacteroides distasonis		0.60	0.11		1	
47774540	**Acceptable identification**					**47774540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium butyricum		93.60	0.48	GAL 99	1	99	
Blautia producta		3.12	0.24	SOR 8	99	1	
Actinomyces israelii		2.38	0.41		1	1	
47774542	**Low discrimination**						**47774542**
					SPO		
Clostridium butyricum		69.18	0.41	GAL 99	99		
Bifidobacterium adolescentis		18.94	0.46	SOR 8	1		
Actinomyces israelii		6.77	0.44		1		
47774552	**Low discrimination**						**47774552**
					SPO		
Bifidobacterium adolescentis		46.82	0.47	GAL 99	1		
Clostridium clostridioforme		39.42	0.43	SOR 11	66		
Clostridium butyricum		7.36	0.19		99		
47774572	**Acceptable identification**						**47774572**
					SPO		
Clostridium clostridioforme		94.14	0.43	RHA 88	66		
Bifidobacterium dentium		1.56	-.03	SOR 1	1		
Clostridium butyricum		1.53	0.01		99		
47774600	**Low discrimination**						**47774600**
					SPO		
Clostridium clostridioforme		76.66	0.37	BGL 99	66		
Clostridium butyricum		21.56	0.16		99		
Actinomyces israelii		1.79	0.19		1		
47774602	**Very good identification**						**47774602**
					SPO		
Clostridium clostridioforme		99.85	0.72		66		
Clostridium butyricum		0.11	0.09		99		
Actinomyces israelii		0.04	0.22		1		
47774603	**Good identification**						**47774603**
Clostridium clostridioforme		100.0	0.37	URE 1			
47774610	**Good identification**						**47774610**
Clostridium clostridioforme		99.68	0.49	BGL 99			
Clostridium butyricum		0.32	-.06				
47774612	**Very good identification**						**47774612**
Clostridium clostridioforme		100.0	0.83				
47774613	**Good identification**						**47774613**
Clostridium clostridioforme		100.0	0.49	URE 1			
47774620	**Good identification**						**47774620**
Clostridium clostridioforme		97.61	0.37	BGL 99			
Clostridium butyricum		2.39	-.02				
47774622	**Very good identification**						**47774622**
Clostridium clostridioforme		99.87	0.72				
Parabacteroides distasonis		0.07	0.23				
Bacteroides ovatus		0.06	0.06				
47774623	**Good identification**						**47774623**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.37	URE 1			
47774630	**Good identification**						**47774630**
Clostridium clostridioforme		100.0	0.49	BGL 99			
47774632	**Very good identification**						**47774632**
Clostridium clostridioforme		99.98	0.83				
Parabacteroides distasonis		0.01	0.19				
Bacteroides ovatus		0.01	0.03				
47774633	**Good identification**						**47774633**
Clostridium clostridioforme		100.0	0.49	URE 1			
47774642	**Good identification**						**47774642**
Clostridium clostridioforme		98.66	0.37	SOR 1			
Actinomyces israelii		1.34	0.15				
47774652	**Good identification**						**47774652**
Clostridium clostridioforme		100.0	0.49	SOR 1			
47774662	**Good identification**						**47774662**
Clostridium clostridioforme		100.0	0.37	SOR 1			
47774672	**Good identification**						**47774672**
Clostridium clostridioforme		100.0	0.49	SOR 1			
47774700	**Low discrimination**						**47774700**
					GRA	SPO	
Clostridium butyricum		64.31	0.40	GAL 99	99	99	
Clostridium clostridioforme		31.90	0.46	RHA 3	50	66	
Mitsuokella multacida		2.90	0.25		1	1	
47774702	**Very good identification**						**47774702**
					SPO		
Clostridium clostridioforme		99.08	0.81		66		
Clostridium butyricum		0.79	0.33		99		
Bifidobacterium adolescentis		0.05	0.27		1		
47774703	**Good identification**						**47774703**
Clostridium clostridioforme		99.21	0.46	URE 1			
Clostridium butyricum		0.79	-.01				
47774710	**Very good identification**						**47774710**
					GRA	SPO	
Clostridium clostridioforme		96.89	0.58	BGL 99	50	66	
Clostridium butyricum		2.26	0.18		99	99	
Mitsuokella multacida		0.65	0.17		1	1	
47774712	**Excellent identification**						**47774712**
					SPO		
Clostridium clostridioforme		99.97	0.92		66		
Bifidobacterium adolescentis		0.01	0.29		1		
Clostridium butyricum		0.01	0.11		99		
47774713	**Very good identification**						**47774713**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		100.0	0.58	URE 1			
47774720	**Low discrimination**						**47774720**
					GRA	SPO	
Clostridium clostridioforme		78.26	0.46	BGL 99	50	66	
Clostridium butyricum		13.72	0.22		99	99	
Bacteroides vulgatus		3.25	0.10		1	1	
47774722	**Very good identification**						**47774722**
Clostridium clostridioforme		96.14	0.81				
Parabacteroides distasonis		2.05	0.58				
Bacteroides ovatus		1.64	0.40				
47774723	**Good identification**						**47774723**
Clostridium clostridioforme		96.31	0.46	URE 1			
Parabacteroides distasonis		2.05	0.23				
Bacteroides ovatus		1.64	0.06				
47774730	**Very good identification**						**47774730**
Clostridium clostridioforme		99.14	0.58	BGL 99			
Parabacteroides distasonis		0.25	0.19				
Bacteroides ovatus		0.25	0.03				
47774732	**Excellent identification**						**47774732**
Clostridium clostridioforme		99.49	0.92				
Parabacteroides distasonis		0.25	0.53				
Bacteroides ovatus		0.25	0.37				
47774733	**Very good identification**						**47774733**
Clostridium clostridioforme		99.50	0.58	URE 1			
Parabacteroides distasonis		0.25	0.19				
Bacteroides ovatus		0.25	0.03				
47774742	**Low discrimination**						**47774742**
					COC	SPO	
Clostridium clostridioforme		89.16	0.46	SOR 1	1	66	
Clostridium butyricum		6.08	0.15		1	99	
Blautia producta		1.87	0.07		99	1	
47774752	**Very good identification**						**47774752**
					COC		
Clostridium clostridioforme		98.33	0.58	SOR 1	1		
Blautia producta		1.36	0.15		99		
Bifidobacterium adolescentis		0.16	0.13		1		
47774762	**Acceptable identification**						**47774762**
Clostridium clostridioforme		91.11	0.46	SOR 1			
Bacteroides ovatus		6.41	0.16				
Parabacteroides distasonis		1.94	0.23				
47774772	**Very good identification**						**47774772**
Clostridium clostridioforme		98.59	0.58	SOR 1			
Bacteroides ovatus		1.02	0.13				
Parabacteroides distasonis		0.25	0.19				
47775000	**Low discrimination**						**47775000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Clostridium butyricum		71.53	0.60	LAC 96	1	99	
Actinomyces israelii		26.80	0.74	ARA 90	1	1	
Atopobium parvulum		0.60	0.32		99	1	
47775002	**Low discrimination**						**47775002**
					SPO		
Actinomyces israelii		55.08	0.77		1		
Clostridium butyricum		38.11	0.53		99		
Clostridium clostridioforme		2.53	0.45		66		
47775003	**Low discrimination**						**47775003**
					SPO		
Actinomyces israelii		51.45	0.43	URE 1	1		
Clostridium butyricum		35.60	0.19		99		
Actinomyces naeslundii		8.34	0.38		1		
47775010	**Low discrimination**						**47775010**
					SPO		
Actinomyces israelii		49.76	0.60	MLZ 13	1		
Clostridium butyricum		46.78	0.38		99		
Lactobacillus jensenii		2.02	0.24		1		
47775012	**Low discrimination**						**47775012**
Clostridium clostridioforme		44.02	0.56	RHA 88			
Actinomyces israelii		31.40	0.63	ARA 94			
Bifidobacterium adolescentis		11.16	0.49				
47775020	**Low discrimination**						**47775020**
					GRA	SPO	
Clostridium butyricum		72.68	0.42	LAC 96	99	99	
Actinomyces israelii		19.99	0.53	ARA 90	99	1	
Prevotella oralis		2.59	0.22	NAG 8	1	1	
47775021	**Low discrimination**						**47775021**
					SPO		
Actinomyces naeslundii		45.11	0.38	ESL 12	1		
Clostridium butyricum		43.05	0.08	XYL 1	99		
Actinomyces israelii		11.84	0.19		1		
47775022	**Low discrimination**						**47775022**
					GRA		
Prevotella oralis		28.54	0.44	TRE 6	1		
Parabacteroides distasonis		18.59	0.53	XYL 11	1		
Actinomyces israelii		17.38	0.56	LAC 94	99		
47775023	**Low discrimination**						**47775023**
					GRA		
Actinomyces naeslundii		32.09	0.38	ESL 12	99		
Prevotella oralis		20.75	0.09	XYL 1	1		
Parabacteroides distasonis		13.51	0.19		1		
47775032	**Low discrimination**						**47775032**
					GRA		
Clostridium clostridioforme		78.32	0.56	RHA 88	50		
Parabacteroides distasonis		13.74	0.49	ARA 94	1		
Actinomyces israelii		3.57	0.42		99		
47775040	**Low discrimination**						**47775040**
					SPO		
Actinomyces israelii		62.60	0.67		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		37.36	0.42		99		
Actinomyces naeslundii		0.05	0.22		1		
47775042	**Low discrimination**						**47775042**
					SPO		
Actinomyces israelii		84.83	0.70		1		
Clostridium butyricum		13.13	0.35		99		
Bifidobacterium adolescentis		1.17	0.32		1		
47775043	**Low discrimination**						**47775043**
					SPO		
Actinomyces israelii		81.42	0.36	URE 1	1		
Clostridium butyricum		12.59	0.01		99		
Actinomyces naeslundii		5.99	0.25		1		
47775050	**Low discrimination**						**47775050**
					COC	SPO	
Actinomyces israelii		82.04	0.53	MLZ 13	1	1	
Clostridium butyricum		17.24	0.20		1	99	
Blautia producta		0.72	-.03		99	1	
47775052	**Low discrimination**						**47775052**
					SPO		
Actinomyces israelii		80.75	0.56	MLZ 13	1		
Bifidobacterium adolescentis		9.12	0.33		1		
Clostridium butyricum		4.40	0.13		99		
47775060	**Low discrimination**						**47775060**
					SPO		
Actinomyces israelii		54.24	0.46	NAG 6	1		
Clostridium butyricum		44.10	0.24		99		
Bifidobacterium breve		1.03	0.12		1		
47775062	**Low discrimination**						**47775062**
					SPO		
Actinomyces israelii		66.76	0.49	NAG 6	1		
Clostridium butyricum		14.07	0.17		99		
Bifidobacterium breve		8.51	0.29		1		
47775072	**Low discrimination**						**47775072**
Actinomyces israelii		38.50	0.35	MLZ 13			
Bifidobacterium breve		32.83	0.29	NAG 6			
Clostridium clostridioforme		21.96	0.22				
47775100	**Good identification**						**47775100**
					COC	SPO	
Clostridium butyricum		90.96	0.84	ARA 90	1	99	
Actinomyces israelii		5.68	0.84		1	1	
Atopobium parvulum		3.12	0.66		99	1	
47775101	**Acceptable identification**						**47775101**
					COC	SPO	
Clostridium butyricum		90.09	0.50	ARA 90	1	99	
Actinomyces israelii		5.63	0.50	URE 1	1	1	
Atopobium parvulum		3.09	0.32		99	1	
47775102	**Low discrimination**						**47775102**
					SPO		
Clostridium butyricum		64.24	0.77	ARA 90	99		
Bifidobacterium adolescentis		16.65	0.81		1		
Actinomyces israelii		15.47	0.87		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47775103	**Low discrimination**						**47775103**
					SPO		
Clostridium butyricum		63.19	0.43	ARA 90	99		
Bifidobacterium adolescentis		16.37	0.47	URE 1	1		
Actinomyces israelii		15.22	0.53		1		
47775110	**Low discrimination**						**47775110**
					SPO		
Clostridium butyricum		81.20	0.62	ARA 90	99		
Actinomyces israelii		14.40	0.70	MLZ 5	1		
Bifidobacterium adolescentis		1.92	0.49		1		
47775112	**Low discrimination**						**47775112**
					GRA	SPO	
Bifidobacterium adolescentis		53.17	0.83	ARA 87	99	1	
Leptotrichia buccalis		22.26	0.58		1	1	
Clostridium butyricum		8.84	0.55		99	99	
47775113	**Low discrimination**						**47775113**
					GRA	SPO	
Bifidobacterium adolescentis		53.09	0.49	ARA 87	99	1	
Leptotrichia buccalis		22.23	0.24	URE 1	1	1	
Clostridium butyricum		8.82	0.21		99	99	
47775120	**Low discrimination**						**47775120**
					COC	SPO	
Clostridium butyricum		82.33	0.66	ARA 90	1	99	
Atopobium parvulum		5.74	0.53	NAG 8	99	1	
Actinomyces israelii		3.77	0.64		1	1	
47775122	**Low discrimination**						**47775122**
					GRA	SPO	
Parabacteroides distasonis		60.30	0.87		1	1	
Prevotella oralis		14.65	0.64		1	1	
Clostridium butyricum		12.89	0.59		99	99	
47775123	**Low discrimination**						**47775123**
					GRA	SPO	
Parabacteroides distasonis		58.12	0.53	URE 1	1	1	
Prevotella oralis		14.12	0.30		1	1	
Clostridium butyricum		12.42	0.24		99	99	
47775130	**Low discrimination**						**47775130**
					GRA	SPO	
Clostridium butyricum		61.96	0.44	ARA 90	99	99	
Parabacteroides distasonis		11.65	0.49	MLZ 5	1	1	
Bifidobacterium breve		9.82	0.46	NAG 8	99	1	
47775132	**Low discrimination**						**47775132**
					GRA		
Parabacteroides distasonis		70.32	0.83		1		
Clostridium clostridioforme		13.55	0.65		50		
Bifidobacterium breve		6.05	0.63		99		
47775133	**Low discrimination**						**47775133**
					GRA		
Parabacteroides distasonis		70.65	0.49	URE 1	1		
Clostridium clostridioforme		13.62	0.31		50		
Bifidobacterium breve		6.08	0.29		99		
47775140	**Low discrimination**						**47775140**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		77.31	0.66	ARA 90	99		
Actinomyces israelii		21.59	0.77	SOR 8	1		
Bifidobacterium breve		0.36	0.41		1		
47775142	**Low discrimination**						**47775142**
					SPO		
Actinomyces israelii		41.66	0.80		1		
Clostridium butyricum		38.68	0.59		99		
Bifidobacterium adolescentis		14.24	0.66		1		
47775143	**Low discrimination**						**47775143**
					SPO		
Actinomyces israelii		41.06	0.46	URE 1	1		
Clostridium butyricum		38.11	0.24		99		
Bifidobacterium adolescentis		14.04	0.32		1		
47775150	**Low discrimination**						**47775150**
					COC	SPO	
Clostridium butyricum		47.75	0.44	ARA 90	1	99	
Actinomyces israelii		37.86	0.63	MLZ 5	1	1	
Blautia producta		8.23	0.32	SOR 8	99	1	
47775152	**Low discrimination**						**47775152**
Bifidobacterium adolescentis		55.63	0.67	ARA 87			
Actinomyces israelii		19.90	0.66	SOR 11			
Bifidobacterium breve		15.08	0.59				
47775160	**Low discrimination**						**47775160**
					SPO		
Clostridium butyricum		74.27	0.47	ARA 90	99		
Actinomyces israelii		15.23	0.57	NAG 8	1		
Bifidobacterium breve		7.12	0.46	SOR 8	1		
47775162	**Low discrimination**						**47775162**
					SPO		
Bifidobacterium breve		47.75	0.63	XYL 1	1		
Clostridium butyricum		19.15	0.40		99		
Actinomyces israelii		15.14	0.60		1		
47775170	**Low discrimination**						**47775170**
					SPO		
Bifidobacterium breve		49.78	0.46	XYL 1	1		
Clostridium butyricum		27.32	0.26	BGL 91	99		
Actinomyces israelii		15.90	0.43		1		
47775172	**Low discrimination**						**47775172**
					GRA		
Bifidobacterium breve		80.51	0.63	XYL 1	99		
Parabacteroides distasonis		9.45	0.49		1		
Actinomyces israelii		3.82	0.46		99		
47775202	**Low discrimination**						**47775202**
					SPO		
Clostridium clostridioforme		76.69	0.59	ARA 94	66		
Actinomyces israelii		17.12	0.58		1		
Clostridium butyricum		4.87	0.27		99		
47775210	**Low discrimination**						**47775210**
					SPO		
Clostridium clostridioforme		59.89	0.36	ARA 94	66		
Actinomyces israelii		28.94	0.40	BGL 99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		11.18	0.12		99		
47775212	**Very good identification**						**47775212**
					GRA		
Clostridium clostridioforme		99.08	0.71	ARA 94	50		
Actinomyces israelii		0.73	0.43		99		
Parabacteroides distasonis		0.07	0.23		1		
47775213	**Good identification**						**47775213**
Clostridium clostridioforme		100.0	0.36	ARA 94			
47775222	**Low discrimination**						**47775222**
Clostridium clostridioforme		56.09	0.59	ARA 94			
Parabacteroides distasonis		34.06	0.61				
Prevotella oralis		7.47	0.37				
47775230	**Low discrimination**						**47775230**
					GRA		
Clostridium clostridioforme		89.38	0.36	ARA 94	50		
Parabacteroides distasonis		6.42	0.23	BGL 99	1		
Actinomyces israelii		2.76	0.20		99		
47775232	**Good identification**						**47775232**
Clostridium clostridioforme		93.12	0.71	ARA 94			
Parabacteroides distasonis		6.68	0.57				
Bacteroides ovatus		0.13	0.11				
47775233	**Acceptable identification**						**47775233**
Clostridium clostridioforme		93.30	0.36	ARA 94			
Parabacteroides distasonis		6.70	0.23	URE 1			
47775240	**Low discrimination**						**47775240**
					SPO		
Actinomyces israelii		80.31	0.48	RHA 7	1		
Clostridium butyricum		19.69	0.16		99		
47775242	**Low discrimination**						**47775242**
					SPO		
Actinomyces israelii		84.75	0.51	RHA 7	1		
Clostridium clostridioforme		9.86	0.25		66		
Clostridium butyricum		5.39	0.09		99		
47775252	**Low discrimination**						**47775252**
Clostridium clostridioforme		78.01	0.36	ARA 94			
Actinomyces israelii		21.99	0.36	SOR 1			
47775272	**Acceptable identification**						**47775272**
					GRA		
Clostridium clostridioforme		91.76	0.36	ARA 94	50		
Parabacteroides distasonis		6.59	0.23	SOR 1	1		
Actinomyces israelii		1.65	0.16		99		
47775300	**Low discrimination**						**47775300**
					COC	SPO	
Clostridium butyricum		84.39	0.58	RHA 3	1	99	
Actinomyces israelii		12.83	0.65	ARA 90	1	1	
Atopobium parvulum		0.95	0.32		99	1	
47775302	**Low discrimination**						**47775302**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium clostridioforme		50.11	0.68	ARA 94	66		
Clostridium butyricum		22.80	0.51		99		
Actinomyces israelii		13.36	0.68		1		
47775310	**Low discrimination**						**47775310**
					SPO		
Clostridium butyricum		44.71	0.36	RHA 3	99		
Clostridium clostridioforme		33.41	0.45	ARA 90	66		
Actinomyces israelii		19.29	0.51	MLZ 5	1		
47775312	**Very good identification**						**47775312**
					GRA		
Clostridium clostridioforme		95.15	0.80	ARA 94	50		
Parabacteroides distasonis		2.04	0.57		1		
Bifidobacterium adolescentis		0.98	0.49		99		
47775313	**Good identification**						**47775313**
					GRA		
Clostridium clostridioforme		95.23	0.45	ARA 94	50		
Parabacteroides distasonis		2.04	0.23	URE 1	1		
Bifidobacterium adolescentis		0.98	0.15		99		
47775320	**Low discrimination**						**47775320**
					SPO	GRA	
Parabacteroides distasonis		41.24	0.61	BGL 99	1	1	
Clostridium butyricum		23.13	0.40		99	99	
Clostridium ramosum		17.68	0.43		92	50	
47775322	**Low discrimination**						**47775322**
Parabacteroides distasonis		88.69	0.95				
Clostridium clostridioforme		4.94	0.68				
Prevotella oralis		3.08	0.58				
47775323	**Low discrimination**						**47775323**
Parabacteroides distasonis		88.81	0.61	URE 1			
Clostridium clostridioforme		4.95	0.34				
Prevotella oralis		3.08	0.24				
47775330	**Low discrimination**						**47775330**
					GRA	SPO	
Parabacteroides distasonis		63.27	0.57	BGL 99	1	1	
Clostridium clostridioforme		29.81	0.45		50	66	
Clostridium butyricum		3.47	0.18		99	99	
47775332	**Low discrimination**						**47775332**
Parabacteroides distasonis		67.02	0.91				
Clostridium clostridioforme		31.57	0.80				
Bacteroides ovatus		1.26	0.45				
47775333	**Low discrimination**						**47775333**
Parabacteroides distasonis		67.12	0.57	URE 1			
Clostridium clostridioforme		31.62	0.45				
Bacteroides ovatus		1.26	0.11				
47775340	**Low discrimination**						**47775340**
					COC	SPO	
Clostridium butyricum		54.80	0.40	RHA 3	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		37.25	0.58	ARA 90	1	1	
Blautia producta		5.36	0.24	SOR 8	99	1	
47775342	**Low discrimination**						**47775342**
					SPO		
Actinomyces israelii		62.97	0.61	RHA 7	1		
Clostridium butyricum		24.02	0.33		99		
Clostridium clostridioforme		6.13	0.34		66		
47775352	**Low discrimination**						**47775352**
Clostridium clostridioforme		60.81	0.45	ARA 94			
Actinomyces israelii		20.48	0.47	SOR 1			
Bifidobacterium adolescentis		7.69	0.33				
47775362	**Low discrimination**						**47775362**
Parabacteroides distasonis		80.14	0.61	SOR 1			
Bacteroides ovatus		5.02	0.25				
Clostridium clostridioforme		4.46	0.34				
47775372	**Low discrimination**						**47775372**
Parabacteroides distasonis		62.77	0.57	SOR 1			
Clostridium clostridioforme		29.57	0.45				
Bacteroides ovatus		4.87	0.22				
47775400	**Very good identification**						**47775400**
					SPO		
Clostridium butyricum		96.63	0.76	LAC 96	99		
Actinomyces israelii		3.29	0.72		1		
Clostridium clostridioforme		0.04	0.31		66		
47775401	**Good identification**						**47775401**
					GRA	SPO	
Clostridium butyricum		96.52	0.42	LAC 96	99	99	
Actinomyces israelii		3.29	0.38	URE 1	99	1	
Mitsuokella multacida		0.19	0.04		1	1	
47775402	**Low discrimination**						**47775402**
					SPO		
Clostridium butyricum		77.28	0.69	LAC 96	99		
Actinomyces israelii		10.15	0.75		1		
Clostridium clostridioforme		8.94	0.65		66		
47775403	**Low discrimination**						**47775403**
					SPO		
Clostridium butyricum		77.29	0.35	LAC 96	99		
Actinomyces israelii		10.16	0.41	URE 1	1		
Clostridium clostridioforme		8.94	0.31		66		
47775410	**Low discrimination**						**47775410**
					SPO		
Clostridium butyricum		87.27	0.55	LAC 96	99		
Actinomyces israelii		8.44	0.58	MLZ 5	1		
Clostridium clostridioforme		3.43	0.42		66		
47775412	**Low discrimination**						**47775412**
					SPO		
Clostridium clostridioforme		79.62	0.76	RHA 88	66		
Bifidobacterium adolescentis		8.62	0.63		1		
Clostridium butyricum		7.95	0.48		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47775413	**Low discrimination**						**47775413**
					SPO		
Clostridium clostridioforme		79.63	0.42	RHA 88	66		
Bifidobacterium adolescentis		8.63	0.29	URE 1	1		
Clostridium butyricum		7.95	0.13		99		
47775420	**Very good identification**						**47775420**
					SPO		
Clostridium butyricum		96.97	0.58	LAC 96	99		
Actinomyces israelii		2.43	0.52	NAG 8	1		
Clostridium clostridioforme		0.51	0.31		66		
47775422	**Low discrimination**						**47775422**
					SPO		
Clostridium clostridioforme		52.65	0.65	RHA 88	66		
Clostridium butyricum		39.58	0.51		99		
Actinomyces israelii		3.82	0.55		1		
47775430	**Low discrimination**						**47775430**
					SPO		
Clostridium butyricum		62.04	0.36	LAC 96	99		
Clostridium clostridioforme		28.07	0.42	MLZ 5	66		
Bifidobacterium dentium		5.09	0.15	NAG 8	1		
47775432	**Very good identification**						**47775432**
					SPO		
Clostridium clostridioforme		96.27	0.76	RHA 88	66		
Bifidobacterium dentium		1.78	0.32		1		
Clostridium butyricum		0.84	0.29		99		
47775433	**Good identification**						**47775433**
					SPO		
Clostridium clostridioforme		96.71	0.42	RHA 88	66		
Bifidobacterium dentium		1.79	-.03	URE 1	1		
Clostridium butyricum		0.84	-.05		99		
47775440	**Low discrimination**						**47775440**
					COC	SPO	
Clostridium butyricum		86.00	0.58	LAC 96	1	99	
Actinomyces israelii		13.10	0.65	SOR 8	1	1	
Blautia producta		0.69	0.24		99	1	
47775442	**Low discrimination**						**47775442**
					SPO		
Clostridium butyricum		59.87	0.51	LAC 96	99		
Actinomyces israelii		35.18	0.68	SOR 8	1		
Bifidobacterium adolescentis		3.97	0.46		1		
47775450	**Low discrimination**						**47775450**
					COC	SPO	
Clostridium butyricum		52.19	0.36	LAC 96	1	99	
Blautia producta		24.00	0.32	MLZ 5	99	1	
Actinomyces israelii		22.57	0.51	SOR 8	1	1	
47775452	**Low discrimination**						**47775452**
Actinomyces israelii		30.24	0.54	MLZ 13			
Bifidobacterium adolescentis		27.95	0.47				
Clostridium clostridioforme		21.09	0.42				
47775460	**Low discrimination**						**47775460**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		88.64	0.40	LAC 96	1	99	
Actinomyces israelii		9.91	0.45	NAG 8	1	1	
Blautia producta		0.91	0.07	SOR 8	99	1	
47775472	**Low discrimination**						**47775472**
					SPO		
Clostridium clostridioforme		82.13	0.42	RHA 88	66		
Actinomyces israelii		7.52	0.34	SOR 1	1		
Clostridium butyricum		6.14	0.11		99		
47775500	**Excellent identification**						**47775500**
					SPO		
Clostridium butyricum		99.27	1.00		99		
Actinomyces israelii		0.56	0.83		1		
Bifidobacterium adolescentis		0.08	0.61		1		
47775501	**Very good identification**						**47775501**
					GRA	SPO	
Clostridium butyricum		98.55	0.66	URE 1	99	99	
Mitsuokella multacida		0.82	0.39		1	1	
Actinomyces israelii		0.56	0.48		99	1	
47775502	**Low discrimination**						**47775502**
					SPO		
Clostridium butyricum		81.20	0.93		99		
Bifidobacterium adolescentis		15.64	0.96		1		
Actinomyces israelii		1.78	0.86		1		
47775503	**Low discrimination**						**47775503**
					SPO		
Clostridium butyricum		81.07	0.59	URE 1	99		
Bifidobacterium adolescentis		15.62	0.61		1		
Actinomyces israelii		1.78	0.52		1		
47775510	**Good identification**						**47775510**
					SPO		
Clostridium butyricum		94.15	0.78	MLZ 5	99		
Bifidobacterium adolescentis		1.66	0.63		1		
Bifidobacterium dentium		1.56	0.44		1		
47775511	**Acceptable identification**						**47775511**
					GRA	SPO	
Clostridium butyricum		90.11	0.44	MLZ 5	99	99	
Mitsuokella multacida		4.72	0.30	URE 1	1	1	
Bifidobacterium adolescentis		1.58	0.29		99	1	
47775512	**Low discrimination**						**47775512**
					SPO		
Bifidobacterium adolescentis		60.69	0.97		1		
Clostridium clostridioforme		18.95	0.85		66		
Clostridium butyricum		13.57	0.71		99		
47775513	**Low discrimination**						**47775513**
					SPO		
Bifidobacterium adolescentis		60.59	0.63	URE 1	1		
Clostridium clostridioforme		18.92	0.51		66		
Clostridium butyricum		13.54	0.37		99		
47775520	**Very good identification**						**47775520**
					GRA	SPO	
Clostridium butyricum		99.03	0.82	NAG 8	99	99	
Actinomyces israelii		0.41	0.62		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Mitsuokella multacida		0.21	0.45		1	1		
47775521	**Good identification**							**47775521**
					GRA	SPO		
Clostridium butyricum		97.02	0.48	NAG 8	99	99		
Mitsuokella multacida		2.31	0.28	URE 1	1	1		
Actinomyces israelii		0.40	0.28		99	1		
47775522	**Low discrimination**							**47775522**
					GRA	SPO		
Clostridium butyricum		62.53	0.75	NAG 8	99	99		
Parabacteroides distasonis		19.93	0.84		1	1		
Clostridium clostridioforme		11.61	0.74		50	66		
47775523	**Low discrimination**							**47775523**
					GRA	SPO		
Clostridium butyricum		62.37	0.41	NAG 8	99	99		
Parabacteroides distasonis		19.88	0.49	URE 1	1	1		
Clostridium clostridioforme		11.58	0.40		50	66		
47775530	**Low discrimination**							**47775530**
					SPO			
Clostridium butyricum		69.10	0.60	MLZ 5	99			
Bifidobacterium dentium		23.38	0.49	NAG 8	1			
Clostridium clostridioforme		4.36	0.51		66			
47775532	**Low discrimination**							**47775532**
					GRA			
Clostridium clostridioforme		53.13	0.85	RHA 88	50			
Bifidobacterium dentium		29.09	0.66		99			
Parabacteroides distasonis		10.78	0.79		1			
47775533	**Low discrimination**							**47775533**
					GRA			
Clostridium clostridioforme		53.12	0.51	RHA 88	50			
Bifidobacterium dentium		29.09	0.32	URE 1	99			
Parabacteroides distasonis		10.78	0.45		1			
47775540	**Good identification**							**47775540**
					COC	SPO		
Clostridium butyricum		93.60	0.82	SOR 8	1	99		
Blautia producta		3.12	0.58		99	1		
Actinomyces israelii		2.38	0.76		1	1		
47775541	**Low discrimination**							**47775541**
					COC	GRA	SPO	
Clostridium butyricum		86.60	0.48	SOR 8	1	99	99	
Mitsuokella multacida		8.23	0.39	URE 1	1	1	1	
Blautia producta		2.88	0.24		99	99	1	
47775542	**Low discrimination**							**47775542**
					SPO			
Clostridium butyricum		72.16	0.75	SOR 8	99			
Bifidobacterium adolescentis		19.76	0.80		1			
Actinomyces israelii		7.07	0.79		1			
47775543	**Low discrimination**							**47775543**
					SPO			
Clostridium butyricum		70.91	0.41	SOR 8	99			
Bifidobacterium adolescentis		19.42	0.46	URE 1	1			
Actinomyces israelii		6.94	0.44		1			
47775550	**Low discrimination**							**47775550**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Blautia producta		62.17	0.66	MAN 99	99	1		
Clostridium butyricum		32.78	0.60		1	99		
Actinomyces israelii		2.36	0.61		1	1		
47775552	**Low discrimination**							**47775552**
					COC	SPO		
Bifidobacterium adolescentis		75.61	0.81	SOR 11	1	1		
Clostridium butyricum		11.89	0.53		1	99		
Blautia producta		6.44	0.50		99	1		
47775553	**Low discrimination**							**47775553**
					COC	SPO		
Bifidobacterium adolescentis		74.26	0.47	SOR 11	1	1		
Clostridium butyricum		11.68	0.19	URE 1	1	99		
Blautia producta		6.33	0.15		99	1		
47775560	**Good identification**							**47775560**
					COC	GRA	SPO	
Clostridium butyricum		92.03	0.64	NAG 8	1	99	99	
Blautia producta		3.91	0.41	SOR 8	99	99	1	
Mitsuokella multacida		2.29	0.45		1	1	1	
47775562	**Low discrimination**							**47775562**
					GRA	SPO		
Clostridium butyricum		82.97	0.57	NAG 8	99	99		
Actinomyces israelii		5.97	0.58	SOR 8	99	1		
Parabacteroides distasonis		3.07	0.49		1	1		
47775570	**Low discrimination**							**47775570**
					COC	GRA	SPO	
Blautia producta		65.74	0.50	MAN 99	99	99	1	
Clostridium butyricum		27.12	0.42	NAG 10	1	99	99	
Mitsuokella multacida		4.27	0.37		1	1	1	
47775572	**Low discrimination**							**47775572**
					SPO			
Clostridium clostridioforme		29.88	0.51	RHA 88	66			
Bifidobacterium dentium		16.36	0.32	SOR 1	1			
Clostridium butyricum		16.02	0.35		99			
47775600	**Low discrimination**							**47775600**
					SPO			
Clostridium butyricum		83.22	0.51	LAC 96	99			
Clostridium clostridioforme		8.97	0.46	RHA 3	66			
Actinomyces israelii		6.90	0.53		1			
47775602	**Very good identification**							**47775602**
					SPO			
Clostridium clostridioforme		95.31	0.80		66			
Clostridium butyricum		3.48	0.44		99			
Actinomyces israelii		1.11	0.56		1			
47775603	**Good identification**							**47775603**
					SPO			
Clostridium clostridioforme		95.41	0.46	URE 1	66			
Clostridium butyricum		3.48	0.09		99			
Actinomyces israelii		1.11	0.22		1			
47775610	**Low discrimination**							**47775610**
					SPO			
Clostridium clostridioforme		87.62	0.57	BGL 99	66			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		9.40	0.29		99		
Actinomyces israelii		2.21	0.39		1		
47775612	**Very good identification**						**47775612**
					SPO		
Clostridium clostridioforme		99.90	0.91		66		
Clostridium butyricum		0.04	0.22		99		
Actinomyces israelii		0.04	0.42		1		
47775613	**Very good identification**						**47775613**
Clostridium clostridioforme		100.0	0.57	URE 1			
47775620	**Low discrimination**						**47775620**
					SPO		
Clostridium clostridioforme		50.75	0.46	BGL 99	66		
Clostridium butyricum		40.95	0.32		99		
Actinomyces israelii		2.49	0.33		1		
47775622	**Very good identification**						**47775622**
Clostridium clostridioforme		95.29	0.80				
Parabacteroides distasonis		2.26	0.58				
Bacteroides ovatus		1.81	0.40				
47775623	**Good identification**						**47775623**
Clostridium clostridioforme		95.90	0.46	URE 1			
Parabacteroides distasonis		2.28	0.23				
Bacteroides ovatus		1.82	0.06				
47775630	**Very good identification**						**47775630**
					GRA	SPO	
Clostridium clostridioforme		98.20	0.57	BGL 99	50	66	
Clostridium butyricum		0.92	0.11		99	99	
Parabacteroides distasonis		0.28	0.19		1	1	
47775632	**Very good identification**						**47775632**
Clostridium clostridioforme		99.42	0.91				
Parabacteroides distasonis		0.28	0.53				
Bacteroides ovatus		0.28	0.37				
47775633	**Very good identification**						**47775633**
Clostridium clostridioforme		99.44	0.57	URE 1			
Parabacteroides distasonis		0.28	0.19				
Bacteroides ovatus		0.28	0.03				
47775642	**Low discrimination**						**47775642**
					SPO		
Clostridium clostridioforme		55.43	0.46	SOR 1	66		
Actinomyces israelii		24.88	0.49		1		
Clostridium butyricum		17.40	0.25		99		
47775652	**Very good identification**						**47775652**
					COC		
Clostridium clostridioforme		96.49	0.57	SOR 1	1		
Blautia producta		1.49	0.15		99		
Actinomyces israelii		1.42	0.35		1		
47775662	**Low discrimination**						**47775662**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		86.34	0.46	SOR 1	50		
Bacteroides ovatus		6.78	0.16		1		
Actinomyces israelii		2.47	0.29		99		
47775672	**Very good identification**						**47775672**
Clostridium clostridioforme		98.33	0.57	SOR 1			
Bacteroides ovatus		1.13	0.13				
Parabacteroides distasonis		0.28	0.19				
47775700	**Good identification**						**47775700**
					GRA	SPO	
Clostridium butyricum		92.94	0.74	RHA 3	99	99	
Mitsuokella multacida		4.19	0.60		1	1	
Clostridium clostridioforme		1.40	0.55		50	66	
47775701	**Low discrimination**						**47775701**
					GRA	SPO	
Clostridium butyricum		65.56	0.40	RHA 3	99	99	
Mitsuokella multacida		32.55	0.43	URE 1	1	1	
Clostridium clostridioforme		0.99	0.20		50	66	
47775702	**Low discrimination**						**47775702**
					SPO		
Clostridium clostridioforme		76.43	0.89		66		
Clostridium butyricum		19.97	0.67		99		
Bifidobacterium adolescentis		1.26	0.61		1		
47775703	**Low discrimination**						**47775703**
					GRA	SPO	
Clostridium clostridioforme		74.54	0.55	URE 1	50	66	
Clostridium butyricum		19.48	0.33		99	99	
Mitsuokella multacida		2.76	0.27		1	1	
47775710	**Low discrimination**						**47775710**
					GRA	SPO	
Clostridium clostridioforme		48.31	0.66	BGL 99	50	66	
Clostridium butyricum		37.13	0.52		99	99	
Mitsuokella multacida		10.61	0.51		1	1	
47775711	**Low discrimination**						**47775711**
					GRA	SPO	
Mitsuokella multacida		56.77	0.35	NIT 99	1	1	
Clostridium clostridioforme		23.49	0.32	URE 10	50	66	
Clostridium butyricum		18.06	0.18		99	99	
47775712	**Very good identification**						**47775712**
					SPO		
Clostridium clostridioforme		98.98	1.00		66		
Bifidobacterium adolescentis		0.44	0.63		1		
Clostridium butyricum		0.30	0.45		99		
47775713	**Very good identification**						**47775713**
					SPO		
Clostridium clostridioforme		98.75	0.66	URE 1	66		
Bifidobacterium adolescentis		0.44	0.29		1		
Clostridium butyricum		0.30	0.11		99		
47775720	**Low discrimination**						**47775720**
					GRA	SPO	
Clostridium butyricum		56.25	0.56	RHA 3	99	99	
Bacteroides vulgatus		13.31	0.44	NAG 8	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium clostridioforme		9.72	0.55		50	66		
47775722	**Low discrimination**							**47775722**
Clostridium clostridioforme		42.94	0.89					
Parabacteroides distasonis		30.17	0.92					
Bacteroides ovatus		24.16	0.74					
47775723	**Low discrimination**							**47775723**
Clostridium clostridioforme		42.82	0.55	URE 1				
Parabacteroides distasonis		30.08	0.58					
Bacteroides ovatus		24.09	0.40					
47775730	**Low discrimination**							**47775730**
Clostridium clostridioforme		76.52	0.66	BGL 99				
Parabacteroides distasonis		6.35	0.53					
Bacteroides ovatus		6.30	0.37					
47775732	**Low discrimination**							**47775732**
Clostridium clostridioforme		85.52	1.00					
Parabacteroides distasonis		7.10	0.87					
Bacteroides ovatus		7.04	0.71					
47775733	**Low discrimination**							**47775733**
Clostridium clostridioforme		85.48	0.66	URE 1				
Parabacteroides distasonis		7.10	0.53					
Bacteroides ovatus		7.04	0.37					
47775740	**Low discrimination**							**47775740**
					COC	GRA	SPO	
Blautia producta		40.47	0.58	MAN 99	99	99	1	
Clostridium butyricum		37.61	0.56		1	99	99	
Mitsuokella multacida		19.52	0.60		1	1	1	
47775741	**Low discrimination**							**47775741**
					COC	GRA	SPO	
Mitsuokella multacida		72.76	0.43	NIT 99	1	1	1	
Blautia producta		13.71	0.24	URE 10	99	99	1	
Clostridium butyricum		12.75	0.22		1	99	99	
47775742	**Low discrimination**							**47775742**
					COC	SPO		
Clostridium butyricum		44.61	0.49	RHA 3	1	99		
Clostridium clostridioforme		19.83	0.55	SOR 8	1	66		
Blautia producta		13.71	0.41		99	1		
47775750	**Good identification**							**47775750**
					COC	GRA	SPO	
Blautia producta		92.99	0.66	MAN 99	99	99	1	
Mitsuokella multacida		4.98	0.51		1	1	1	
Clostridium butyricum		1.52	0.34		1	99	99	
47775752	**Low discrimination**							**47775752**
					COC			
Clostridium clostridioforme		63.21	0.66	SOR 1	1			
Blautia producta		28.79	0.50		99			
Bifidobacterium adolescentis		3.41	0.47		1			
47775760	**Low discrimination**							**47775760**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA	SPO	
Mitsuokella multacida		37.11	0.49	NIT 99	1	1	1	
Blautia producta		34.20	0.41		99	99	1	
Clostridium butyricum		24.87	0.38		1	99	99	
47775762	**Low discrimination**							**47775762**
Bacteroides ovatus		52.11	0.51	IND 97				
Clostridium clostridioforme		22.45	0.55	SOR 4				
Parabacteroides distasonis		15.77	0.58					
47775770	**Low discrimination**							**47775770**
					COC	GRA		
Blautia producta		85.72	0.50	MAN 99	99	99		
Mitsuokella multacida		10.34	0.41	NAG 10	1	1		
Clostridium clostridioforme		1.90	0.32		1	50		
47775772	**Low discrimination**							**47775772**
Clostridium clostridioforme		67.26	0.66	SOR 1				
Bacteroides ovatus		22.84	0.48					
Parabacteroides distasonis		5.59	0.53					
47776000	**Good identification**							**47776000**
					SPO			
Actinomyces israelii		95.03	0.53	GAL 99	1			
Clostridium butyricum		4.97	0.10		99			
47776002	**Good identification**							**47776002**
					SPO			
Actinomyces israelii		98.05	0.56	GAL 99	1			
Clostridium butyricum		1.33	0.03		99			
Bifidobacterium adolescentis		0.62	0.12		1			
47776010	**Good identification**							**47776010**
Actinomyces israelii		100.0	0.38	GAL 99				
47776012	**Low discrimination**							**47776012**
Actinomyces israelii		85.17	0.41	GAL 99				
Clostridium clostridioforme		7.02	0.13	MLZ 13				
Bifidobacterium adolescentis		4.37	0.13					
47776022	**Good identification**							**47776022**
Actinomyces israelii		100.0	0.35	GAL 99				
47776040	**Good identification**							**47776040**
Actinomyces israelii		100.0	0.46	GAL 99				
47776042	**Good identification**							**47776042**
Actinomyces israelii		100.0	0.49	GAL 99				
47776100	**Low discrimination**							**47776100**
					SPO			
Actinomyces israelii		75.43	0.63	GAL 99	1			
Clostridium butyricum		23.68	0.33		99			
Clostridium tertium		0.49	0.11		99			
47776102	**Low discrimination**							**47776102**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		79.21	0.66	GAL 99	1		
Bifidobacterium adolescentis		12.30	0.46		1		
Clostridium butyricum		6.45	0.26		99		
47776110	**Low discrimination**						**47776110**
					SPO		
Actinomyces israelii		72.96	0.49	GAL 99	1		
Bifidobacterium dentium		10.81	0.10	MLZ 13	1		
Clostridium butyricum		8.07	0.11		99		
47776112	**Low discrimination**						**47776112**
Bifidobacterium adolescentis		35.47	0.47	GAL 99			
Actinomyces israelii		27.92	0.52	ARA 87			
Bifidobacterium dentium		27.90	0.27				
47776120	**Low discrimination**						**47776120**
					SPO		
Actinomyces israelii		45.83	0.42	GAL 99	1		
Clostridium butyricum		19.61	0.15	NAG 6	99		
Clostridium ramosum		17.84	0.19		92		
47776140	**Low discrimination**						**47776140**
					COC	SPO	
Actinomyces israelii		79.40	0.56	GAL 99	1	1	
Blautia producta		15.03	0.24		99	1	
Clostridium butyricum		5.58	0.15		1	99	
47776142	**Good identification**						**47776142**
Actinomyces israelii		90.78	0.59	GAL 99			
Bifidobacterium adolescentis		4.48	0.30				
Bifidobacterium breve		1.81	0.24				
47776152	**Low discrimination**						**47776152**
					COC		
Actinomyces israelii		54.20	0.45	GAL 99	1		
Bifidobacterium adolescentis		21.89	0.32	MLZ 13	1		
Blautia producta		15.26	0.15		99		
47776160	**Low discrimination**						**47776160**
					COC	SPO	
Actinomyces israelii		66.33	0.35	GAL 99	1	1	
Blautia producta		21.85	0.07	NAG 6	99	1	
Clostridium butyricum		6.34	-.03		1	99	
47776162	**Low discrimination**						**47776162**
Actinomyces israelii		64.24	0.38	GAL 99			
Bifidobacterium breve		35.76	0.29	NAG 6			
47776202	**Low discrimination**						**47776202**
Actinomyces israelii		79.14	0.36	GAL 99			
Clostridium clostridioforme		20.86	0.17	RHA 7			
47776300	**Low discrimination**						**47776300**
					SPO		
Actinomyces israelii		88.57	0.44	GAL 99	1		
Clostridium butyricum		11.43	0.07	RHA 7	99		
47776302	**Low discrimination**						**47776302**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		77.72	0.47	GAL 99	1		
Clostridium clostridioforme		17.14	0.26	RHA 7	66		
Clostridium butyricum		2.60	0.00		99		
47776312	**Low discrimination**						**47776312**
Clostridium clostridioforme		83.75	0.37	MAN 1			
Actinomyces israelii		12.45	0.33	ARA 94			
Bifidobacterium adolescentis		2.12	0.13				
47776332	**Low discrimination**						**47776332**
Clostridium clostridioforme		86.85	0.37	MAN 1			
Parabacteroides distasonis		5.59	0.23	ARA 94			
Bacteroides ovatus		3.47	0.03				
47776342	**Low discrimination**						**47776342**
					COC		
Actinomyces israelii		84.30	0.40	GAL 99	1		
Blautia producta		15.70	0.07	RHA 7	99		
47776400	**Low discrimination**						**47776400**
					SPO		
Actinomyces israelii		63.47	0.51	GAL 99	1		
Clostridium butyricum		36.53	0.26		99		
47776402	**Low discrimination**						**47776402**
					SPO		
Actinomyces israelii		79.38	0.54	GAL 99	1		
Clostridium butyricum		11.85	0.19		99		
Clostridium clostridioforme		4.11	0.23		66		
47776410	**Low discrimination**						**47776410**
					SPO		
Actinomyces israelii		49.99	0.37	GAL 99	1		
Bifidobacterium dentium		36.23	0.10	MLZ 13	1		
Clostridium butyricum		10.14	0.04		99		
47776440	**Low discrimination**						**47776440**
					COC	SPO	
Actinomyces israelii		48.69	0.44	GAL 99	1	1	
Blautia producta		45.05	0.24		99	1	
Clostridium butyricum		6.27	0.08		1	99	
47776442	**Low discrimination**						**47776442**
					COC	SPO	
Actinomyces israelii		89.43	0.47	GAL 99	1	1	
Blautia producta		6.13	0.07		99	1	
Clostridium butyricum		2.98	0.01		1	99	
47776500	**Low discrimination**						**47776500**
					COC	SPO	
Clostridium butyricum		74.36	0.50	GAL 99	1	99	
Actinomyces israelii		21.53	0.61	MAN 10	1	1	
Blautia producta		1.94	0.24		99	1	
47776502	**Low discrimination**						**47776502**
					SPO		
Bifidobacterium adolescentis		37.34	0.60	GAL 99	1		
Actinomyces israelii		29.39	0.64		1		
Clostridium butyricum		26.32	0.43		99		
47776510	**Low discrimination**						**47776510**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Bifidobacterium dentium		80.58	0.44	GAL 99	1	1	
Blautia producta		8.11	0.32	BGL 91	99	1	
Clostridium butyricum		5.47	0.28		1	99	
47776512	**Good identification**						**47776512**
Bifidobacterium dentium		90.18	0.62	GAL 99			
Bifidobacterium adolescentis		7.75	0.61				
Clostridium clostridioforme		0.99	0.43				
47776530	**Good identification**						**47776530**
					COC	SPO	
Bifidobacterium dentium		98.70	0.49	GAL 99	1	1	
Blautia producta		0.62	0.15	BGL 91	99	1	
Clostridium butyricum		0.33	0.10		1	99	
47776532	**Very good identification**						**47776532**
Bifidobacterium dentium		99.27	0.66	GAL 99			
Clostridium clostridioforme		0.61	0.43				
Bifidobacterium adolescentis		0.05	0.27				
47776540	**Good identification**						**47776540**
					COC	SPO	
Blautia producta		92.27	0.58	GAL 99	99	1	
Actinomyces israelii		4.03	0.54		1	1	
Clostridium butyricum		3.11	0.31		1	99	
47776542	**Low discrimination**						**47776542**
					COC		
Blautia producta		51.16	0.41	GAL 99	99		
Actinomyces israelii		30.16	0.57	BGL 10	1		
Bifidobacterium adolescentis		12.18	0.44		1		
47776550	**Very good identification**						**47776550**
					COC		
Blautia producta		99.55	0.66	GAL 99	99		
Actinomyces israelii		0.22	0.40		1		
Bifidobacterium dentium		0.10	0.10		1		
47776552	**Low discrimination**						**47776552**
					COC		
Blautia producta		81.81	0.50	GAL 99	99		
Bifidobacterium adolescentis		7.93	0.46	BGL 10	1		
Bifidobacterium dentium		7.54	0.27		1		
47776560	**Acceptable identification**						**47776560**
					COC	SPO	
Blautia producta		93.87	0.41	GAL 99	99	1	
Clostridium butyricum		2.48	0.13	NAG 10	1	99	
Actinomyces israelii		2.36	0.34		1	1	
47776570	**Good identification**						**47776570**
					COC	GRA	
Blautia producta		98.14	0.50	GAL 99	99	99	
Bifidobacterium dentium		1.59	0.15	NAG 10	1	99	
Mitsuokella multacida		0.15	0.09		1	1	
47776602	**Low discrimination**						**47776602**
					SPO		
Clostridium clostridioforme		82.62	0.37	MAN 1	66		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		16.37	0.35		1		
Clostridium butyricum		1.00	-.07		99		
47776612	**Good identification**						**47776612**
Clostridium clostridioforme		98.94	0.49	MAN 1			
Actinomyces israelii		0.64	0.21				
Bifidobacterium dentium		0.42	-.07				
47776622	**Good identification**						**47776622**
					GRA		
Clostridium clostridioforme		96.93	0.37	MAN 1	50		
Bacteroides ovatus		1.84	-.02		1		
Actinomyces israelii		1.23	0.14		99		
47776632	**Good identification**						**47776632**
					GRA		
Clostridium clostridioforme		98.98	0.49	MAN 1	50		
Bifidobacterium dentium		0.75	-.03		99		
Bacteroides ovatus		0.28	-.05		1		
47776702	**Low discrimination**						**47776702**
					SPO		
Clostridium clostridioforme		70.41	0.46	MAN 1	66		
Actinomyces israelii		16.67	0.45		1		
Clostridium butyricum		6.13	0.17		99		
47776712	**Low discrimination**						**47776712**
					COC		
Clostridium clostridioforme		86.06	0.58	MAN 1	1		
Bifidobacterium dentium		10.84	0.27		1		
Blautia producta		1.19	0.15		99		
47776722	**Low discrimination**						**47776722**
Clostridium clostridioforme		61.49	0.46	MAN 1			
Bacteroides ovatus		34.60	0.32				
Parabacteroides distasonis		1.31	0.23				
47776732	**Low discrimination**						**47776732**
					GRA		
Clostridium clostridioforme		76.31	0.58	MAN 1	50		
Bifidobacterium dentium		17.09	0.32		99		
Bacteroides ovatus		6.28	0.29		1		
47776740	**Very good identification**						**47776740**
					COC	GRA	
Blautia producta		98.45	0.58	GAL 99	99	99	
Mitsuokella multacida		1.12	0.32		1	1	
Actinomyces israelii		0.32	0.35		1	99	
47776742	**Acceptable identification**						**47776742**
					COC	GRA	
Blautia producta		93.75	0.41	GAL 99	99	99	
Actinomyces israelii		4.16	0.38	BGL 10	1	99	
Mitsuokella multacida		1.07	0.15		1	1	
47776750	**Very good identification**						**47776750**
					COC	GRA	
Blautia producta		99.86	0.66	GAL 99	99	99	
Mitsuokella multacida		0.13	0.24		1	1	
Actinomyces israelii		0.02	0.21		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47776752	**Good identification**						**47776752**
					COC		
Blautia producta		98.74	0.50	GAL 99	99		
Clostridium clostridioforme		0.73	0.24	BGL 10	1		
Actinomyces israelii		0.22	0.24		1		
47776760	**Good identification**						**47776760**
					COC	GRA	
Blautia producta		97.33	0.41	GAL 99	99	99	
Mitsuokella multacida		2.49	0.21	NAG 10	1	1	
Actinomyces israelii		0.18	0.15		1	99	
47776770	**Good identification**						**47776770**
					COC	GRA	
Blautia producta		99.72	0.50	GAL 99	99	99	
Mitsuokella multacida		0.28	0.13	NAG 10	1	1	
47777000	**Good identification**						**47777000**
					SPO		
Actinomyces israelii		94.94	0.87		1		
Clostridium butyricum		4.97	0.44		99		
Actinomyces naeslundii		0.05	0.40		1		
47777001	**Low discrimination**						**47777001**
					SPO		
Actinomyces israelii		88.06	0.53	URE 1	1		
Actinomyces naeslundii		7.34	0.43		1		
Clostridium butyricum		4.61	0.10		99		
47777002	**Very good identification**						**47777002**
					SPO		
Actinomyces israelii		97.91	0.90		1		
Clostridium butyricum		1.33	0.37		99		
Bifidobacterium adolescentis		0.61	0.46		1		
47777003	**Acceptable identification**						**47777003**
					SPO		
Actinomyces israelii		92.99	0.56	URE 1	1		
Actinomyces naeslundii		5.17	0.43		1		
Clostridium butyricum		1.26	0.03		99		
47777010	**Very good identification**						**47777010**
					SPO		
Actinomyces israelii		97.19	0.73	MLZ 13	1		
Clostridium butyricum		1.79	0.22		99		
Bifidobacterium dentium		0.58	0.10		1		
47777011	**Good identification**						**47777011**
Actinomyces israelii		100.0	0.38	MLZ 13			
47777012	**Good identification**						**47777012**
Actinomyces israelii		90.28	0.76	MLZ 13			
Bifidobacterium adolescentis		4.63	0.47				
Bifidobacterium dentium		3.65	0.27				
47777013	**Acceptable identification**						**47777013**
Actinomyces israelii		91.60	0.41	MLZ 13			
Bifidobacterium adolescentis		4.70	0.13	URE 1			
Bifidobacterium dentium		3.70	-.07				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47777020	**Good identification**						**47777020**
					SPO		
Actinomyces israelii		90.13	0.66	NAG 6	1		
Clostridium butyricum		6.43	0.26		99		
Clostridium ramosum		1.42	0.19		92		
47777021	**Low discrimination**						**47777021**
Actinomyces naeslundii		56.62	0.43	ESL 12			
Actinomyces israelii		43.38	0.32	XYL 1			
47777022	**Good identification**						**47777022**
					SPO		
Actinomyces israelii		90.05	0.69	NAG 6	1		
Clostridium ramosum		3.78	0.30		92		
Clostridium tertium		2.23	0.27		99		
47777023	**Low discrimination**						**47777023**
Actinomyces israelii		53.47	0.35	NAG 6			
Actinomyces naeslundii		46.53	0.43	URE 1			
47777030	**Low discrimination**						**47777030**
					SPO		
Actinomyces israelii		74.17	0.52	MLZ 13	1		
Bifidobacterium dentium		12.37	0.15	NAG 6	1		
Clostridium tertium		10.95	0.24		99		
47777040	**Very good identification**						**47777040**
					COC	SPO	
Actinomyces israelii		98.08	0.80		1	1	
Clostridium butyricum		1.15	0.26		1	99	
Blautia producta		0.75	0.24		99	1	
47777041	**Good identification**						**47777041**
Actinomyces israelii		96.36	0.46	URE 1			
Actinomyces naeslundii		3.64	0.30				
47777042	**Very good identification**						**47777042**
					SPO		
Actinomyces israelii		99.35	0.83		1		
Clostridium butyricum		0.30	0.19		99		
Bifidobacterium adolescentis		0.20	0.30		1		
47777043	**Good identification**						**47777043**
Actinomyces israelii		97.54	0.49	URE 1			
Actinomyces naeslundii		2.46	0.30				
47777050	**Low discrimination**						**47777050**
					COC	SPO	
Actinomyces israelii		86.38	0.66	MLZ 13	1	1	
Blautia producta		13.26	0.32		99	1	
Clostridium butyricum		0.36	0.04		1	99	
47777052	**Very good identification**						**47777052**
					COC		
Actinomyces israelii		96.60	0.69	MLZ 13	1		
Bifidobacterium adolescentis		1.58	0.32		1		
Blautia producta		1.10	0.15		99		
47777060	**Very good identification**						**47777060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		96.47	0.59	NAG 6	1	1	
Clostridium butyricum		1.54	0.07		1	99	
Blautia producta		1.28	0.07		99	1	
47777062	**Very good identification**						**47777062**
					SPO		
Actinomyces israelii		97.16	0.62	NAG 6	1		
Bifidobacterium breve		2.19	0.29		1		
Clostridium butyricum		0.40	0.00		99		
47777070	**Low discrimination**						**47777070**
					COC		
Actinomyces israelii		77.55	0.45	MLZ 13	1		
Blautia producta		20.72	0.15	NAG 6	99		
Bifidobacterium breve		1.73	0.12		1		
47777072	**Low discrimination**						**47777072**
Actinomyces israelii		83.37	0.48	MLZ 13			
Bifidobacterium breve		12.55	0.29	NAG 6			
Bifidobacterium dentium		2.44	-.03				
47777100	**Low discrimination**						**47777100**
					SPO		
Actinomyces israelii		75.35	0.97		1		
Clostridium butyricum		23.66	0.67		99		
Clostridium tertium		0.49	0.45		99		
47777101	**Low discrimination**						**47777101**
					SPO		
Actinomyces israelii		72.09	0.63	URE 1	1		
Clostridium butyricum		22.64	0.33		99		
Actinomyces naeslundii		4.27	0.51		1		
47777102	**Low discrimination**						**47777102**
					SPO		
Actinomyces israelii		79.32	1.00		1		
Bifidobacterium adolescentis		12.32	0.80		1		
Clostridium butyricum		6.46	0.60		99		
47777103	**Low discrimination**						**47777103**
					SPO		
Actinomyces israelii		77.01	0.66	URE 1	1		
Bifidobacterium adolescentis		11.96	0.46		1		
Clostridium butyricum		6.27	0.26		99		
47777110	**Low discrimination**						**47777110**
					SPO		
Actinomyces israelii		72.48	0.83	MLZ 13	1		
Bifidobacterium dentium		10.74	0.44		1		
Clostridium butyricum		8.02	0.45		99		
47777111	**Low discrimination**						**47777111**
					SPO		
Actinomyces israelii		71.29	0.49	MLZ 13	1		
Bifidobacterium dentium		10.57	0.10	URE 1	1		
Clostridium butyricum		7.89	0.11		99		
47777112	**Low discrimination**						**47777112**
Bifidobacterium adolescentis		36.13	0.81	ARA 87			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		28.44	0.86				
Bifidobacterium dentium		28.42	0.62				
47777113	**Low discrimination**						**47777113**
Bifidobacterium adolescentis		36.03	0.47	ARA 87			
Actinomyces israelii		28.36	0.52	URE 1			
Bifidobacterium dentium		28.34	0.27				
47777120	**Low discrimination**						**47777120**
					SPO		
Actinomyces israelii		45.93	0.77	NAG 6	1		
Clostridium butyricum		19.65	0.49		99		
Clostridium ramosum		17.88	0.53		92		
47777121	**Low discrimination**						**47777121**
					SPO		
Actinomyces israelii		32.42	0.42	NAG 6	1		
Actinomyces naeslundii		30.11	0.51	URE 1	1		
Clostridium butyricum		13.87	0.15		99		
47777122	**Low discrimination**						**47777122**
					SPO		
Clostridium ramosum		33.27	0.64	XYL 1	92		
Actinomyces israelii		32.05	0.80		1		
Clostridium tertium		19.68	0.62		99		
47777130	**Low discrimination**						**47777130**
					SPO		
Bifidobacterium dentium		44.82	0.49	ARA 99	1		
Clostridium tertium		39.66	0.58	BGL 91	99		
Actinomyces israelii		10.86	0.62		1		
47777132	**Low discrimination**						**47777132**
					SPO		
Bifidobacterium dentium		72.99	0.66	ARA 99	1		
Clostridium tertium		19.16	0.66		99		
Bifidobacterium breve		3.80	0.63		1		
47777140	**Low discrimination**						**47777140**
					COC	SPO	
Actinomyces israelii		79.07	0.90		1	1	
Blautia producta		14.96	0.58		99	1	
Clostridium butyricum		5.55	0.49		1	99	
47777141	**Low discrimination**						**47777141**
					COC	SPO	
Actinomyces israelii		76.98	0.56	URE 1	1	1	
Blautia producta		14.57	0.24		99	1	
Clostridium butyricum		5.41	0.15		1	99	
47777142	**Good identification**						**47777142**
Actinomyces israelii		90.72	0.93				
Bifidobacterium adolescentis		4.48	0.64				
Bifidobacterium breve		1.81	0.59				
47777143	**Low discrimination**						**47777143**
Actinomyces israelii		89.33	0.59	URE 1			
Bifidobacterium adolescentis		4.41	0.30				
Bifidobacterium breve		1.79	0.24				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
47777150	**Low discrimination**						**47777150**
					COC	SPO	
Blautia producta		78.20	0.66	ARA 99	99	1	
Actinomyces israelii		20.58	0.76		1	1	
Clostridium butyricum		0.51	0.27		1	99	
47777152	**Low discrimination**						**47777152**
					COC		
Actinomyces israelii		53.89	0.79	MLZ 13	1		
Bifidobacterium adolescentis		21.76	0.66		1		
Blautia producta		15.17	0.50		99		
47777153	**Low discrimination**						**47777153**
					COC		
Actinomyces israelii		54.20	0.45	MLZ 13	1		
Bifidobacterium adolescentis		21.89	0.32	URE 1	1		
Blautia producta		15.26	0.15		99		
47777160	**Low discrimination**						**47777160**
					COC	SPO	
Actinomyces israelii		65.23	0.70	NAG 6	1	1	
Blautia producta		21.49	0.41		99	1	
Clostridium butyricum		6.24	0.31		1	99	
47777161	**Low discrimination**						**47777161**
					COC		
Actinomyces israelii		51.84	0.35	NAG 6	1		
Actinomyces naeslundii		21.85	0.38	URE 1	1		
Blautia producta		17.08	0.07		99		
47777162	**Low discrimination**						**47777162**
					SPO		
Actinomyces israelii		60.14	0.73	NAG 6	1		
Bifidobacterium breve		33.47	0.63		1		
Clostridium ramosum		1.62	0.30		92		
47777163	**Low discrimination**						**47777163**
Actinomyces israelii		54.42	0.38	NAG 6			
Bifidobacterium breve		30.29	0.29	URE 1			
Actinomyces naeslundii		15.29	0.38				
47777170	**Low discrimination**						**47777170**
					COC		
Blautia producta		78.62	0.50	ARA 99	99		
Actinomyces israelii		11.89	0.55	NAG 10	1		
Bifidobacterium breve		6.57	0.46		1		
47777172	**Low discrimination**						**47777172**
Bifidobacterium breve		60.04	0.63	XYL 1			
Actinomyces israelii		16.12	0.58				
Bifidobacterium dentium		11.65	0.32				
47777200	**Very good identification**						**47777200**
					SPO		
Actinomyces israelii		97.90	0.68	RHA 7	1		
Clostridium butyricum		2.11	0.18		99		
47777202	**Very good identification**						**47777202**
					SPO		
Actinomyces israelii		98.53	0.71	RHA 7	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.79	0.25		66		
Clostridium butyricum		0.55	0.11		99		
47777203	**Good identification**						**47777203**
Actinomyces israelii		100.0	0.36	RHA 7			
47777210	**Good identification**						**47777210**
					COC	SPO	
Actinomyces israelii		98.47	0.53	RHA 7	1	1	
Blautia producta		0.79	-.03	MLZ 13	99	1	
Clostridium butyricum		0.75	-.04		1	99	
47777212	**Low discrimination**						**47777212**
Actinomyces israelii		79.63	0.56	RHA 7			
Clostridium clostridioforme		19.39	0.36	MLZ 13			
Bifidobacterium adolescentis		0.55	0.13				
47777220	**Low discrimination**						**47777220**
					SPO		
Actinomyces israelii		76.56	0.47	RHA 7	1		
Clostridium ramosum		21.20	0.21	NAG 6	92		
Clostridium butyricum		2.24	0.00		99		
47777222	**Low discrimination**						**47777222**
					SPO		
Actinomyces israelii		49.95	0.50	RHA 7	1		
Clostridium ramosum		36.89	0.32	NAG 6	92		
Clostridium clostridioforme		6.25	0.25		66		
47777232	**Low discrimination**						**47777232**
					GRA		
Clostridium clostridioforme		69.88	0.36	MAN 1	50		
Actinomyces israelii		18.32	0.36	ARA 94	99		
Parabacteroides distasonis		5.02	0.23		1		
47777240	**Good identification**						**47777240**
					COC	SPO	
Actinomyces israelii		90.38	0.60	RHA 7	1	1	
Blautia producta		9.18	0.24		99	1	
Clostridium butyricum		0.44	0.00		1	99	
47777242	**Very good identification**						**47777242**
					COC	SPO	
Actinomyces israelii		99.13	0.63	RHA 7	1	1	
Blautia producta		0.75	0.07		99	1	
Clostridium butyricum		0.12	-.07		1	99	
47777252	**Low discrimination**						**47777252**
					COC		
Actinomyces israelii		86.88	0.49	RHA 7	1		
Blautia producta		13.13	0.15	MLZ 13	99		
47777260	**Low discrimination**						**47777260**
					COC		
Actinomyces israelii		84.97	0.40	RHA 7	1		
Blautia producta		15.03	0.07	NAG 6	99		
47777262	**Good identification**						**47777262**
					SPO		
Actinomyces israelii		98.12	0.43	RHA 7	1		
Clostridium ramosum		1.88	-.03	NAG 6	92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47777300	**Low discrimination**						**47777300**
					GRA	SPO	
Actinomyces israelii		86.53	0.78	RHA 7	99	1	
Clostridium butyricum		11.16	0.42		99	99	
Mitsuokella multacida		0.96	0.32		1	1	
47777301	**Low discrimination**						**47777301**
					GRA	SPO	
Actinomyces israelii		79.92	0.44	RHA 7	99	1	
Clostridium butyricum		10.31	0.07	URE 1	99	99	
Mitsuokella multacida		9.77	0.15		1	1	
47777302	**Good identification**						**47777302**
					SPO		
Actinomyces israelii		92.62	0.81	RHA 7	1		
Clostridium butyricum		3.10	0.35		99		
Bifidobacterium adolescentis		1.93	0.46		1		
47777303	**Acceptable identification**						**47777303**
					SPO		
Actinomyces israelii		93.80	0.47	RHA 7	1		
Clostridium butyricum		3.14	0.00	URE 1	99		
Bifidobacterium adolescentis		1.96	0.12		1		
47777310	**Low discrimination**						**47777310**
					COC	SPO	
Actinomyces israelii		76.66	0.64	RHA 7	1	1	
Blautia producta		15.20	0.32	MLZ 13	99	1	
Clostridium butyricum		3.48	0.20		1	99	
47777312	**Low discrimination**						**47777312**
Actinomyces israelii		63.56	0.67	RHA 7			
Clostridium clostridioforme		12.95	0.45	MLZ 13			
Bifidobacterium adolescentis		10.84	0.47				
47777320	**Low discrimination**						**47777320**
					SPO		
Clostridium ramosum		83.75	0.56	XYL 1	92		
Actinomyces israelii		12.22	0.57		1		
Clostridium butyricum		2.15	0.23		99		
47777322	**Low discrimination**						**47777322**
					SPO	GRA	
Clostridium ramosum		81.44	0.66	XYL 1	92	50	
Parabacteroides distasonis		8.38	0.61		1	1	
Actinomyces israelii		4.46	0.60		1	99	
47777330	**Low discrimination**						**47777330**
					SPO		
Actinomyces israelii		30.72	0.43	RHA 7	1		
Bifidobacterium dentium		17.02	0.15	MLZ 13	1		
Clostridium tertium		15.06	0.24	NAG 6	99		
47777332	**Low discrimination**						**47777332**
					GRA		
Parabacteroides distasonis		31.69	0.57	MAN 1	1		
Bacteroides ovatus		19.65	0.37		1		
Bifidobacterium dentium		17.46	0.32		99		
47777340	**Low discrimination**						**47777340**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		70.39	0.58	ARA 99	99	1	
Actinomyces israelii		28.00	0.71		1	1	
Clostridium butyricum		0.81	0.23		1	99	
47777342	**Low discrimination**						**47777342**
					COC	SPO	
Actinomyces israelii		82.96	0.74	RHA 7	1	1	
Blautia producta		15.45	0.41		99	1	
Clostridium butyricum		0.62	0.16		1	99	
47777343	**Low discrimination**						**47777343**
					COC		
Actinomyces israelii		84.30	0.40	RHA 7	1		
Blautia producta		15.70	0.07	URE 1	99		
47777350	**Very good identification**						**47777350**
					COC	GRA	
Blautia producta		97.92	0.66	ARA 99	99	99	
Actinomyces israelii		1.94	0.57		1	99	
Mitsuokella multacida		0.12	0.24		1	1	
47777352	**Low discrimination**						**47777352**
					COC		
Blautia producta		77.46	0.50	ARA 99	99		
Actinomyces israelii		20.72	0.60	BGL 10	1		
Bifidobacterium adolescentis		1.12	0.32		1		
47777360	**Low discrimination**						**47777360**
					COC	SPO	
Blautia producta		76.56	0.41	ARA 99	99	1	
Actinomyces israelii		17.49	0.50	NAG 10	1	1	
Clostridium ramosum		3.11	0.21		1	92	
47777362	**Low discrimination**						**47777362**
					COC	SPO	
Actinomyces israelii		48.97	0.53	RHA 7	1	1	
Clostridium ramosum		23.25	0.32	NAG 6	1	92	
Blautia producta		15.88	0.25		99	1	
47777370	**Good identification**						**47777370**
					COC	GRA	
Blautia producta		98.52	0.50	ARA 99	99	99	
Actinomyces israelii		1.12	0.36	NAG 10	1	99	
Mitsuokella multacida		0.28	0.13		1	1	
47777400	**Low discrimination**						**47777400**
					COC	SPO	
Actinomyces israelii		63.16	0.85		1	1	
Clostridium butyricum		36.36	0.60		1	99	
Blautia producta		0.23	0.24		99	1	
47777401	**Low discrimination**						**47777401**
					GRA	SPO	
Actinomyces israelii		62.51	0.51	URE 1	99	1	
Clostridium butyricum		35.98	0.26		99	99	
Mitsuokella multacida		1.51	0.11		1	1	
47777402	**Low discrimination**						**47777402**
					SPO		
Actinomyces israelii		82.65	0.88		1		
Clostridium butyricum		12.34	0.53		99		
Bifidobacterium adolescentis		4.24	0.60		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
47777403	**Low discrimination**						**47777403**
					SPO		
Actinomyces israelii		82.79	0.54	URE 1	1		
Clostridium butyricum		12.36	0.19		99		
Bifidobacterium adolescentis		4.25	0.26		1		
47777410	**Low discrimination**						**47777410**
					SPO		
Actinomyces israelii		49.69	0.71	MLZ 13	1		
Bifidobacterium dentium		36.01	0.44		1		
Clostridium butyricum		10.08	0.38		99		
47777411	**Low discrimination**						**47777411**
					SPO		
Actinomyces israelii		48.68	0.37	MLZ 13	1		
Bifidobacterium dentium		35.27	0.10	URE 1	1		
Clostridium butyricum		9.87	0.04		99		
47777412	**Low discrimination**						**47777412**
Bifidobacterium dentium		76.20	0.62	LAC 99			
Actinomyces israelii		15.60	0.74				
Bifidobacterium adolescentis		6.55	0.61				
47777420	**Low discrimination**						**47777420**
					SPO		
Actinomyces israelii		54.66	0.65	NAG 6	1		
Clostridium butyricum		42.86	0.42		99		
Bifidobacterium dentium		1.67	0.15		1		
47777422	**Low discrimination**						**47777422**
					SPO		
Actinomyces israelii		69.03	0.68	NAG 6	1		
Bifidobacterium dentium		14.17	0.32		1		
Clostridium butyricum		14.03	0.35		99		
47777430	**Acceptable identification**						**47777430**
					SPO		
Bifidobacterium dentium		93.37	0.49	LAC 99	1		
Actinomyces israelii		4.63	0.51	BGL 91	1		
Clostridium butyricum		1.28	0.20		99		
47777432	**Very good identification**						**47777432**
Bifidobacterium dentium		98.60	0.66	LAC 99			
Actinomyces israelii		0.73	0.54				
Clostridium clostridioforme		0.54	0.42				
47777440	**Low discrimination**						**47777440**
					COC	SPO	
Actinomyces israelii		48.55	0.78		1	1	
Blautia producta		44.92	0.58		99	1	
Clostridium butyricum		6.25	0.42		1	99	
47777441	**Low discrimination**						**47777441**
					COC	SPO	
Actinomyces israelii		47.26	0.44	URE 1	1	1	
Blautia producta		43.72	0.24		99	1	
Clostridium butyricum		6.08	0.08		1	99	
47777442	**Low discrimination**						**47777442**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		89.38	0.81		1	1	
Blautia producta		6.13	0.41		99	1	
Clostridium butyricum		2.98	0.35		1	99	
47777443	**Low discrimination**						**47777443**
					COC	SPO	
Actinomyces israelii		89.43	0.47	URE 1	1	1	
Blautia producta		6.13	0.07		99	1	
Clostridium butyricum		2.98	0.01		1	99	
47777450	**Good identification**						**47777450**
					COC	SPO	
Blautia producta		94.51	0.66	LAC 99	99	1	
Actinomyces israelii		5.09	0.64		1	1	
Clostridium butyricum		0.23	0.20		1	99	
47777452	**Low discrimination**						**47777452**
					COC		
Blautia producta		51.92	0.50	LAC 99	99		
Actinomyces israelii		37.73	0.67	BGL 10	1		
Bifidobacterium adolescentis		5.04	0.46		1		
47777460	**Low discrimination**						**47777460**
					COC	SPO	
Blautia producta		57.35	0.41	LAC 99	99	1	
Actinomyces israelii		35.61	0.58	NAG 10	1	1	
Clostridium butyricum		6.25	0.24		1	99	
47777462	**Low discrimination**						**47777462**
					COC	SPO	
Actinomyces israelii		85.28	0.61	NAG 6	1	1	
Blautia producta		10.17	0.25		99	1	
Clostridium butyricum		3.88	0.17		1	99	
47777470	**Good identification**						**47777470**
					COC		
Blautia producta		95.19	0.50	LAC 99	99		
Actinomyces israelii		2.94	0.44	NAG 10	1		
Bifidobacterium dentium		1.54	0.15		1		
47777500	**Low discrimination**						**47777500**
					COC	SPO	
Clostridium butyricum		74.36	0.84	MAN 10	1	99	
Actinomyces israelii		21.53	0.96		1	1	
Blautia producta		1.94	0.58		99	1	
47777501	**Low discrimination**						**47777501**
					GRA	SPO	
Clostridium butyricum		66.55	0.50	MAN 10	99	99	
Actinomyces israelii		19.27	0.61	URE 1	99	1	
Mitsuokella multacida		11.55	0.45		1	1	
47777502	**Low discrimination**						**47777502**
					SPO		
Bifidobacterium adolescentis		37.80	0.94		1		
Actinomyces israelii		29.76	0.99		1		
Clostridium butyricum		26.65	0.77		99		
47777503	**Low discrimination**						**47777503**
					SPO		
Bifidobacterium adolescentis		37.36	0.60	URE 1	1		
Actinomyces israelii		29.41	0.64		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		26.33	0.43		99		
47777510	**Low discrimination**						**47777510**
					COC	SPO	
Bifidobacterium dentium		80.65	0.79	BGL 91	1	1	
Blautia producta		8.12	0.66		99	1	
Clostridium butyricum		5.47	0.62		1	99	
47777511	**Low discrimination**						**47777511**
					COC	GRA	
Bifidobacterium dentium		76.47	0.44	URE 1	1	99	
Blautia producta		7.70	0.32	BGL 91	99	99	
Mitsuokella multacida		5.70	0.37		1	1	
47777512	**Good identification**						**47777512**
Bifidobacterium dentium		91.04	0.96				
Bifidobacterium adolescentis		7.83	0.96				
Actinomyces israelii		0.75	0.84				
47777513	**Good identification**						**47777513**
Bifidobacterium dentium		90.99	0.62	URE 1			
Bifidobacterium adolescentis		7.82	0.61				
Actinomyces israelii		0.75	0.50				
47777520	**Low discrimination**						**47777520**
					SPO		
Clostridium butyricum		68.74	0.66	MAN 10	99		
Actinomyces israelii		14.61	0.75	NAG 8	1		
Bifidobacterium dentium		11.02	0.49		1		
47777522	**Low discrimination**						**47777522**
					SPO		
Bifidobacterium dentium		65.51	0.66	MLZ 99	1		
Clostridium butyricum		15.72	0.59		99		
Actinomyces israelii		12.89	0.78		1		
47777530	**Very good identification**						**47777530**
					COC	SPO	
Bifidobacterium dentium		98.74	0.83	BGL 91	1	1	
Blautia producta		0.62	0.50		99	1	
Clostridium butyricum		0.33	0.44		1	99	
47777531	**Good identification**						**47777531**
					COC	GRA	
Bifidobacterium dentium		97.84	0.49	URE 1	1	99	
Mitsuokella multacida		1.03	0.26	BGL 91	1	1	
Blautia producta		0.62	0.15		99	99	
47777532	**Excellent identification**						**47777532**
Bifidobacterium dentium		99.86	1.00				
Bifidobacterium adolescentis		0.05	0.61				
Actinomyces israelii		0.03	0.64				
47777533	**Very good identification**						**47777533**
Bifidobacterium dentium		99.85	0.66	URE 1			
Bifidobacterium adolescentis		0.05	0.27				
Actinomyces israelii		0.03	0.30				
47777540	**Good identification**						**47777540**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	SPO		
Blautia producta		92.27	0.92		99	1		
Actinomyces israelii		4.03	0.89		1	1		
Clostridium butyricum		3.11	0.66		1	99		
47777541	**Low discrimination**							**47777541**
					COC	GRA		
Blautia producta		87.34	0.58	URE 1	99	99		
Mitsuokella multacida		5.88	0.45		1	1		
Actinomyces israelii		3.81	0.54		1	99		
47777542	**Low discrimination**							**47777542**
					COC			
Blautia producta		51.16	0.76	BGL 10	99			
Actinomyces israelii		30.16	0.92		1			
Bifidobacterium adolescentis		12.18	0.79		1			
47777543	**Low discrimination**							**47777543**
					COC			
Blautia producta		49.61	0.41	URE 1	99			
Actinomyces israelii		29.25	0.57	BGL 10	1			
Bifidobacterium adolescentis		11.81	0.44		1			
47777550	**Excellent identification**							**47777550**
					COC			
Blautia producta		99.55	1.00		99			
Actinomyces israelii		0.22	0.74		1			
Bifidobacterium dentium		0.10	0.44		1			
47777551	**Very good identification**							**47777551**
					COC	GRA		
Blautia producta		98.88	0.66	URE 1	99	99		
Mitsuokella multacida		0.74	0.37		1	1		
Actinomyces israelii		0.22	0.40		1	99		
47777552	**Low discrimination**							**47777552**
					COC			
Blautia producta		81.87	0.84	BGL 10	99			
Bifidobacterium adolescentis		7.94	0.80		1			
Bifidobacterium dentium		7.55	0.62		1			
47777553	**Low discrimination**							**47777553**
					COC			
Blautia producta		81.43	0.50	URE 1	99			
Bifidobacterium adolescentis		7.90	0.46	BGL 10	1			
Bifidobacterium dentium		7.51	0.27		1			
47777560	**Good identification**							**47777560**
					COC	SPO		
Blautia producta		93.83	0.76	NAG 10	99	1		
Clostridium butyricum		2.48	0.47		1	99		
Actinomyces israelii		2.35	0.68		1	1		
47777561	**Low discrimination**							**47777561**
					COC	GRA	SPO	
Blautia producta		83.13	0.41	NAG 10	99	99	1	
Mitsuokella multacida		12.59	0.35	URE 1	1	1	1	
Clostridium butyricum		2.19	0.13		1	99	99	
47777562	**Low discrimination**							**47777562**
					COC	SPO		
Blautia producta		65.82	0.59	NAG 10	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		22.29	0.71	BGL 10	1	1	
Clostridium butyricum		6.08	0.40		1	99	
47777570	**Very good identification**						**47777570**
					COC	GRA	
Blautia producta		98.09	0.84	NAG 10	99	99	
Bifidobacterium dentium		1.59	0.49		1	99	
Mitsuokella multacida		0.15	0.43		1	1	
47777571	**Good identification**						**47777571**
					COC	GRA	
Blautia producta		96.68	0.50	NAG 10	99	99	
Mitsuokella multacida		1.63	0.26	URE 1	1	1	
Bifidobacterium dentium		1.57	0.15		1	99	
47777572	**Low discrimination**						**47777572**
					COC		
Bifidobacterium dentium		58.86	0.66	SOR 1	1		
Blautia producta		39.90	0.67		99		
Actinomyces israelii		0.67	0.57		1		
47777600	**Low discrimination**						**47777600**
					GRA	SPO	
Actinomyces israelii		74.55	0.66	RHA 7	99	1	
Clostridium butyricum		17.63	0.34		99	99	
Mitsuokella multacida		4.05	0.32		1	1	
47777602	**Low discrimination**						**47777602**
					SPO		
Actinomyces israelii		81.26	0.69	RHA 7	1		
Clostridium clostridioforme		12.43	0.46		66		
Clostridium butyricum		4.98	0.27		99		
47777610	**Low discrimination**						**47777610**
					COC	GRA	
Actinomyces israelii		42.70	0.52	RHA 7	1	99	
Blautia producta		41.40	0.32	MLZ 13	99	99	
Mitsuokella multacida		5.18	0.24		1	1	
47777612	**Low discrimination**						**47777612**
Clostridium clostridioforme		71.94	0.57	MAN 1			
Actinomyces israelii		15.43	0.55				
Bifidobacterium dentium		10.12	0.27				
47777620	**Low discrimination**						**47777620**
					GRA	SPO	
Actinomyces israelii		59.59	0.46	RHA 7	99	1	
Clostridium butyricum		19.20	0.16	NAG 6	99	99	
Mitsuokella multacida		12.68	0.21		1	1	
47777622	**Low discrimination**						**47777622**
					GRA		
Clostridium clostridioforme		47.04	0.46	MAN 1	50		
Bacteroides ovatus		29.54	0.32		1		
Actinomyces israelii		19.64	0.49		99		
47777632	**Low discrimination**						**47777632**
					GRA		
Clostridium clostridioforme		73.53	0.57	MAN 1	50		
Bifidobacterium dentium		18.38	0.32		99		
Bacteroides ovatus		6.76	0.29		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
47777640	**Good identification**							**47777640**
					COC	GRA		
Blautia producta		91.16	0.58	LAC 99	99	99		
Actinomyces israelii		7.42	0.59		1	99		
Mitsuokella multacida		1.04	0.32		1	1		
47777642	**Low discrimination**							**47777642**
					COC	SPO		
Actinomyces israelii		51.53	0.62	RHA 7	1	1		
Blautia producta		46.92	0.41		99	1		
Clostridium butyricum		0.71	0.09		1	99		
47777650	**Very good identification**							**47777650**
					COC	GRA		
Blautia producta		99.46	0.66	LAC 99	99	99		
Actinomyces israelii		0.40	0.45		1	99		
Mitsuokella multacida		0.13	0.24		1	1		
47777652	**Acceptable identification**							**47777652**
					COC			
Blautia producta		93.94	0.50	LAC 99	99			
Actinomyces israelii		5.14	0.48	BGL 10	1			
Clostridium clostridioforme		0.62	0.23		1			
47777660	**Acceptable identification**							**47777660**
					COC	GRA		
Blautia producta		92.96	0.41	LAC 99	99	99		
Actinomyces israelii		4.35	0.39	NAG 10	1	99		
Mitsuokella multacida		2.38	0.21		1	1		
47777670	**Good identification**							**47777670**
					COC	GRA		
Blautia producta		99.49	0.50	LAC 99	99	99		
Mitsuokella multacida		0.28	0.13	NAG 10	1	1		
Actinomyces israelii		0.23	0.24		1	99		
47777700	**Low discrimination**							**47777700**
					COC	GRA	SPO	
Clostridium butyricum		28.57	0.58	MAN 10	1	99	99	
Mitsuokella multacida		27.07	0.66	RHA 3	1	1	1	
Blautia producta		24.06	0.58		99	99	1	
47777701	**Low discrimination**							**47777701**
					COC	GRA	SPO	
Mitsuokella multacida		80.37	0.50	NIT 99	1	1	1	
Clostridium butyricum		7.71	0.24	URE 10	1	99	99	
Blautia producta		6.49	0.24		99	99	1	
47777702	**Low discrimination**							**47777702**
					SPO			
Actinomyces israelii		52.52	0.79	RHA 7	1			
Clostridium butyricum		19.32	0.51		99			
Bifidobacterium adolescentis		8.95	0.60		1			
47777710	**Low discrimination**							**47777710**
					COC	GRA		
Blautia producta		77.44	0.66	SOR 99	99	99		
Mitsuokella multacida		9.68	0.58		1	1		
Bifidobacterium dentium		7.77	0.44		1	99		
47777711	**Low discrimination**							**47777711**
					COC	GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Mitsuokella multacida		54.19	0.41	NIT 99	1	1		
Blautia producta		39.40	0.32	URE 10	99	99		
Bifidobacterium dentium		3.95	0.10		1	99		
47777712	**Low discrimination**							**47777712**
					COC			
Bifidobacterium dentium		65.53	0.62	RHA 1	1			
Clostridium clostridioforme		15.76	0.66		1			
Blautia producta		7.18	0.50		99			
47777720	**Low discrimination**							**47777720**
					COC	GRA	SPO	
Mitsuokella multacida		47.35	0.56	NIT 99	1	1	1	
Blautia producta		18.70	0.41		99	99	1	
Clostridium butyricum		17.38	0.40		1	99	99	
47777721	**Acceptable identification**							**47777721**
					COC	GRA	SPO	
Mitsuokella multacida		91.22	0.39	NIT 99	1	1	1	
Blautia producta		3.28	0.07	URE 10	99	99	1	
Clostridium butyricum		3.04	0.06		1	99	99	
47777722	**Low discrimination**							**47777722**
Bacteroides ovatus		85.19	0.66	IND 97				
Clostridium clostridioforme		4.59	0.55					
Parabacteroides distasonis		3.22	0.58					
47777730	**Low discrimination**							**47777730**
					COC	GRA		
Bifidobacterium dentium		53.31	0.49	RHA 1	1	99		
Blautia producta		33.21	0.50	BGL 91	99	99		
Mitsuokella multacida		9.34	0.47		1	1		
47777732	**Low discrimination**							**47777732**
					GRA			
Bifidobacterium dentium		65.52	0.66	RHA 1	99			
Bacteroides ovatus		24.08	0.63		1			
Clostridium clostridioforme		8.86	0.66		50			
47777740	**Very good identification**							**47777740**
					COC	GRA		
Blautia producta		98.45	0.92		99	99		
Mitsuokella multacida		1.12	0.66		1	1		
Actinomyces israelii		0.32	0.69		1	99		
47777741	**Low discrimination**							**47777741**
					COC	GRA		
Blautia producta		88.54	0.58	URE 1	99	99		
Mitsuokella multacida		11.07	0.50		1	1		
Actinomyces israelii		0.29	0.35		1	99		
47777742	**Good identification**							**47777742**
					COC	GRA		
Blautia producta		94.15	0.76	BGL 10	99	99		
Actinomyces israelii		4.18	0.72		1	99		
Mitsuokella multacida		1.07	0.50		1	1		
47777743	**Low discrimination**							**47777743**
					COC	GRA		
Blautia producta		85.07	0.41	URE 1	99	99		
Mitsuokella multacida		10.64	0.33	BGL 10	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		3.78	0.38		1	99	
47777750	**Excellent identification**						**47777750**
					COC	GRA	
Blautia producta		99.86	1.00		99	99	
Mitsuokella multacida		0.13	0.58		1	1	
Actinomyces israelii		0.02	0.55		1	99	
47777751	**Very good identification**						**47777751**
					COC	GRA	
Blautia producta		98.61	0.66	URE 1	99	99	
Mitsuokella multacida		1.37	0.41		1	1	
Actinomyces israelii		0.02	0.21		1	99	
47777752	**Very good identification**						**47777752**
					COC	GRA	
Blautia producta		99.43	0.84	BGL 10	99	99	
Actinomyces israelii		0.22	0.58		1	99	
Mitsuokella multacida		0.13	0.41		1	1	
47777753	**Good identification**						**47777753**
					COC	GRA	
Blautia producta		98.23	0.50	URE 1	99	99	
Mitsuokella multacida		1.37	0.25	BGL 10	1	1	
Actinomyces israelii		0.22	0.24		1	99	
47777760	**Very good identification**						**47777760**
					COC	GRA	
Blautia producta		97.24	0.76	NAG 10	99	99	
Mitsuokella multacida		2.49	0.56		1	1	
Actinomyces israelii		0.18	0.49		1	99	
47777761	**Low discrimination**						**47777761**
					COC	GRA	
Blautia producta		77.93	0.41	NAG 10	99	99	
Mitsuokella multacida		21.93	0.39	URE 1	1	1	
Actinomyces israelii		0.15	0.15		1	99	
47777762	**Low discrimination**						**47777762**
					COC	GRA	
Blautia producta		86.30	0.59	NAG 10	99	99	
Bacteroides ovatus		8.78	0.42	BGL 10	1	1	
Mitsuokella multacida		2.21	0.39		1	1	
47777770	**Very good identification**						**47777770**
					COC	GRA	
Blautia producta		99.69	0.84	NAG 10	99	99	
Mitsuokella multacida		0.28	0.47		1	1	
Bifidobacterium dentium		0.02	0.15		1	99	
47777771	**Good identification**						**47777771**
					COC	GRA	
Blautia producta		96.97	0.50	NAG 10	99	99	
Mitsuokella multacida		3.03	0.31	URE 1	1	1	
47777772	**Very good identification**						**47777772**
					COC	GRA	
Blautia producta		95.79	0.67	NAG 10	99	99	
Bacteroides ovatus		2.17	0.39	BGL 10	1	1	
Bifidobacterium dentium		1.43	0.32		1	99	
50000000	**Low discrimination**						**50000000**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Fusobacterium nucleatum subsp. nucleatum		21.16	0.66	ESL 1	1	1		
Peptoniphilus asaccharolyticus		21.16	0.66		99	99		
Porphyromonas asaccharolytica		20.94	0.66		1	1		
50000001	**Low discrimination**							**50000001**
					COC	GRA	SPO	
Anaerococcus prevotii		16.58	0.41	IND 9	99	99	1	
Clostridium sordellii		13.50	0.40	ESL 1	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		12.84	0.32		1	1	1	
50000002	**Low discrimination**							**50000002**
					GRA			
Porphyromonas asaccharolytica		39.43	0.42	ESL 1	1			
Atopobium minutum		10.18	0.53	BGL 4	99			
Fusobacterium nucleatum subsp. nucleatum		9.66	0.32		1			
50000020	**Low discrimination**							**50000020**
					GRA	SPO		
Clostridium bifermentans		36.12	0.54	GLU 92	99	99		
Clostridium cadaveris		13.17	0.44	NAG 6	99	99		
Fusobacterium nucleatum subsp. nucleatum		9.84	0.32		1	1		
50000022	**Very good identification**							**50000022**
					GRA			
Pseudoflavonifractor capillosus		98.84	0.66	IND 1	1			
Atopobium minutum		0.82	0.44		99			
Porphyromonas asaccharolytica		0.10	0.08		1			
50000040	**Low discrimination**							**50000040**
					COC	GRA	SPO	
Clostridium bifermentans		33.70	0.50	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		14.07	0.32	SOR 4	1	1	1	
Peptoniphilus asaccharolyticus		14.07	0.32		99	99	1	
50001000	**Good identification**							**50001000**
					COC	GRA		
Fusobacterium necrophorum subsp. necrophorum		90.47	0.66	ESL 1	1	1		
Fusobacterium nucleatum subsp. nucleatum		2.00	0.32		1	1		
Peptoniphilus asaccharolyticus		2.00	0.32		99	99		
50001022	**Good identification**							**50001022**
					GRA			
Pseudoflavonifractor capillosus		97.47	0.50	IND 1	1			
Atopobium minutum		2.44	0.36	GAL 10	99			
Fusobacterium necrophorum subsp. necrophorum		0.10	-.03		1			
50010000	**Low discrimination**							**50010000**
					GRA	SPO		
Fusobacterium varium		34.13	0.50	ESL 1	1	1		
Clostridium bifermentans		21.75	0.52		99	99		
Atopobium minutum		10.49	0.55		99	1		
50010001	**Low discrimination**							**50010001**
					GRA	SPO		
Clostridium sordellii		86.97	0.40	ESL 5	99	99		
Fusobacterium varium		3.97	0.16	GLU 95	1	1		
Clostridium bifermentans		2.53	0.18	MLT 90	99	99		
50010002	**Low discrimination**							**50010002**
					GRA	SPO		
Atopobium minutum		74.53	0.52	IND 1	99	1		
Clostridium sporogenes		10.32	0.27		99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium varium		3.67	0.16		1	1	
50010020	**Low discrimination**						**50010020**
					SPO		
Clostridium cadaveris		61.86	0.44	ESL 1	99		
Atopobium minutum		22.49	0.47	GLU 95	1		
Clostridium bifermentans		8.93	0.32		99		
50010022	**Acceptable identification**						**50010022**
					GRA	SPO	
Pseudoflavonifractor capillosus		93.24	0.50	IND 1	1	1	
Atopobium minutum		6.45	0.44	FRU 10	99	1	
Clostridium cadaveris		0.27	0.10		99	99	
50011000	**Low discrimination**						**50011000**
					GRA		
Atopobium minutum		45.50	0.47	IND 1	99		
Fusobacterium necrophorum subsp. necrophorum		42.86	0.32		1		
Fusobacterium varium		4.49	0.16		1		
50011002	**Good identification**						**50011002**
					GRA		
Atopobium minutum		97.04	0.44	IND 1	99		
Pseudoflavonifractor capillosus		1.57	-.01		1		
Fusobacterium necrophorum subsp. necrophorum		1.39	-.03		1		
50011020	**Low discrimination**						**50011020**
					SPO	CAT	
Atopobium minutum		84.08	0.39	IND 1	1	1	
Clostridium cadaveris		7.01	0.10		99	1	
Propionibacterium acnes		2.77	0.06		1	95	
50100000	**Good identification**						**50100000**
Clostridium bifermentans		92.91	0.80	GLU 92			
Clostridium sporogenes		2.95	0.53				
Clostridium botulinum biovar A		1.11	0.48				
50100001	**Low discrimination**						**50100001**
Clostridium sordellii		86.45	0.56	ESL 5			
Clostridium bifermentans		12.38	0.46	GLU 95			
Clostridium sporogenes		0.39	0.19				
50100002	**Low discrimination**						**50100002**
					GRA	SPO	
Clostridium sporogenes		49.27	0.45	IND 1	99	99	
Clostridium bifermentans		47.09	0.46		99	99	
Porphyromonas asaccharolytica		1.43	0.08		1	1	
50100010	**Acceptable identification**						**50100010**
Clostridium bifermentans		92.62	0.46	GLU 92			
Clostridium sporogenes		2.94	0.19	MLZ 1			
Clostridium botulinum biovar A		1.11	0.14				
50100020	**Very good identification**						**50100020**
Clostridium bifermentans		97.92	0.60	GLU 92			
Clostridium cadaveris		0.81	0.22	NAG 6			
Clostridium sporogenes		0.49	0.19				
50100040	**Good identification**						**50100040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		90.70	0.57	GLU 92			
Clostridium botulinum biovar A		7.78	0.39	SOR 4			
Clostridium sporogenes		0.70	0.19				
50100060	**Good identification**						**50100060**
Clostridium bifermentans		98.66	0.36	GLU 92			
Clostridium botulinum biovar A		1.34	0.05	NAG 6			
50100100	**Acceptable identification**						**50100100**
Clostridium bifermentans		93.03	0.46	GLU 92			
Clostridium sporogenes		2.95	0.19	LAC 1			
Clostridium botulinum biovar A		1.11	0.14				
50100200	**Acceptable identification**						**50100200**
Clostridium bifermentans		93.03	0.46	GLU 92			
Clostridium sporogenes		2.95	0.19	RHA 1			
Clostridium botulinum biovar A		1.11	0.14				
50100400	**Acceptable identification**						**50100400**
Clostridium bifermentans		93.03	0.46	GLU 92			
Clostridium sporogenes		2.95	0.19	ARA 1			
Clostridium botulinum biovar A		1.11	0.14				
50101000	**Low discrimination**						**50101000**
					GRA	SPO	
Clostridium bifermentans		69.92	0.46	GLU 92	99	99	
Fusobacterium necrophorum subsp. necrophorum		23.65	0.32	GAL 1	1	1	
Clostridium sporogenes		2.22	0.19		99	99	
50101001	**Good identification**						**50101001**
					GRA	SPO	
Clostridium sordellii		98.92	0.43	ESL 5	99	99	
Clostridium bifermentans		0.81	0.12	GLU 95	99	99	
Fusobacterium necrophorum subsp. necrophorum		0.27	-.03		1	1	
50102000	**Acceptable identification**						**50102000**
Clostridium bifermentans		93.03	0.46	GLU 92			
Clostridium sporogenes		2.95	0.19	MAN 1			
Clostridium botulinum biovar A		1.11	0.14				
50104000	**Acceptable identification**						**50104000**
Clostridium bifermentans		93.03	0.46	GLU 92			
Clostridium sporogenes		2.95	0.19	XYL 1			
Clostridium botulinum biovar A		1.11	0.14				
50110000	**Low discrimination**						**50110000**
					GRA	SPO	
Clostridium bifermentans		85.16	0.59	GLU 92	99	99	
Clostridium sporogenes		9.05	0.40	FRU 5	99	99	
Prevotella intermedia		2.13	0.34		1	1	
50110001	**Very good identification**						**50110001**
Clostridium sordellii		99.17	0.56	ESL 5			
Clostridium bifermentans		0.75	0.24	GLU 95			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium sporogenes		0.08	0.06					
50110020	**Low discrimination**							**50110020**
Clostridium bifermentans		84.46	0.38	GLU 92				
Clostridium cadaveris		13.20	0.22	FRU 5				
Clostridium sporogenes		1.42	0.06	NAG 6				
50111001	**Good identification**							**50111001**
Clostridium sordellii		100.0	0.43	ESL 5				
50120000	**Low discrimination**							**50120000**
Clostridium sporogenes		44.08	0.45	IND 1				
Clostridium bifermentans		42.13	0.46					
Clostridium botulinum biovar A		12.47	0.38					
50120002	**Good identification**							**50120002**
Clostridium sporogenes		96.38	0.37	IND 1				
Clostridium bifermentans		2.79	0.12					
Clostridium botulinum biovar A		0.83	0.04					
50140000	**Acceptable identification**							**50140000**
Clostridium bifermentans		93.03	0.46	GLU 92				
Clostridium sporogenes		2.95	0.19	CEL 1				
Clostridium botulinum biovar A		1.11	0.14					
50200000	**Low discrimination**							**50200000**
					COC	GRA	SPO	
Clostridium bifermentans		60.66	0.58	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		9.50	0.32	SAL 10	1	1	1	
Peptoniphilus asaccharolyticus		9.50	0.32		99	99	1	
50200020	**Low discrimination**							**50200020**
					GRA	SPO		
Clostridium bifermentans		86.49	0.37	GLU 92	99	99		
Clostridium cadaveris		2.87	0.10	SAL 10	99	99		
Fusobacterium nucleatum subsp. nucleatum		2.14	-.03	NAG 6	1	1		
50210000	**Low discrimination**							**50210000**
					GRA	SPO		
Clostridium bifermentans		75.80	0.36	GLU 92	99	99		
Fusobacterium varium		10.81	0.16	SAL 10	1	1		
Atopobium minutum		3.32	0.21	FRU 5	99	1		
50300000	**Very good identification**							**50300000**
Clostridium bifermentans		99.32	0.64	GLU 92				
Clostridium sporogenes		0.29	0.19	SAL 10				
Clostridium botulinum biovar A		0.11	0.14					
50300020	**Good identification**							**50300020**
Clostridium bifermentans		100.0	0.44	GLU 92				
50300040	**Good identification**							**50300040**
Clostridium bifermentans		99.23	0.40	GLU 92				
Clostridium botulinum biovar A		0.77	0.05	SAL 10				
50310000	**Good identification**							**50310000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		99.04	0.42	GLU 92			
Clostridium sporogenes		0.96	0.06	SAL 10			
50500000	**Acceptable identification**						**50500000**
Clostridium bifermentans		92.86	0.46	GLU 92			
Clostridium sporogenes		2.94	0.19	RAF 1			
Clostridium botulinum biovar A		1.11	0.14				
50510000	**Low discrimination**						**50510000**
					GRA	SPO	
Prevotella intermedia		77.08	0.37	ESL 1	1	1	
Clostridium bifermentans		20.72	0.24	GLU 90	99	99	
Clostridium sporogenes		2.20	0.06		99	99	
50511000	**Good identification**						**50511000**
Prevotella intermedia		96.79	0.37	ESL 1			
Eubacterium saburreum		3.22	0.21	GLU 90			
51000000	**Good identification**						**51000000**
Clostridium bifermentans		93.66	0.92				
Clostridium botulinum biovar A		2.50	0.66				
Clostridium sporogenes		1.84	0.61				
51000001	**Low discrimination**						**51000001**
Clostridium sordellii		71.97	0.62	ESL 5			
Clostridium bifermentans		24.07	0.58	MLT 90			
Clostridium novyi biovar A		2.30	0.53				
51000002	**Low discrimination**						**51000002**
					SPO		
Clostridium bifermentans		50.47	0.58	BGL 1	99		
Clostridium sporogenes		32.69	0.53		99		
Atopobium minutum		13.56	0.57		1		
51000010	**Low discrimination**						**51000010**
Clostridium bifermentans		85.92	0.58	MLZ 1			
Clostridium novyi biovar A		8.20	0.53				
Clostridium botulinum biovar A		2.29	0.32				
51000011	**Low discrimination**						**51000011**
Clostridium novyi biovar A		70.30	0.53	IND 1			
Clostridium sordellii		22.26	0.28				
Clostridium bifermentans		7.44	0.24				
51000012	**Low discrimination**						**51000012**
Clostridium novyi biovar A		46.51	0.40	IND 1			
Clostridium bifermentans		27.91	0.24				
Clostridium sporogenes		18.08	0.19				
51000013	**Good identification**						**51000013**
Clostridium novyi biovar A		98.22	0.40	IND 1			
Clostridium sordellii		1.78	-.07				
51000020	**Low discrimination**						**51000020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium bifermentans		61.08	0.72	NAG 6	99		
Clostridium cadaveris		36.78	0.66		99		
Atopobium minutum		1.30	0.51		1		
51000021	**Low discrimination**						**51000021**
Clostridium bifermentans		40.48	0.38	NAG 6			
Clostridium cadaveris		24.38	0.32	URE 1			
Clostridium sordellii		19.16	0.28				
51000022	**Low discrimination**						**51000022**
					GRA	SPO	
Atopobium minutum		33.62	0.48	IND 1	99	1	
Clostridium bifermentans		23.96	0.38		99	99	
Pseudoflavonifractor capillosus		23.95	0.32		1	1	
51000030	**Low discrimination**						**51000030**
Clostridium bifermentans		50.07	0.38	MLZ 1			
Clostridium cadaveris		30.16	0.32	NAG 6			
Clostridium novyi biovar A		18.71	0.43				
51000031	**Acceptable identification**						**51000031**
Clostridium novyi biovar A		94.69	0.43	IND 1			
Clostridium bifermentans		2.56	0.03				
Clostridium cadaveris		1.54	-.03				
51000040	**Low discrimination**						**51000040**
Clostridium bifermentans		81.93	0.69	SOR 4			
Clostridium botulinum biovar A		15.68	0.57				
Clostridium novyi biovar A		1.26	0.50				
51000041	**Low discrimination**						**51000041**
Clostridium sordellii		52.39	0.38	ESL 5			
Clostridium novyi biovar A		26.74	0.50	MLT 90			
Clostridium bifermentans		17.52	0.34	SOR 4			
51000043	**Low discrimination**						**51000043**
Clostridium novyi biovar A		86.98	0.37	IND 1			
Clostridium sordellii		9.76	0.04				
Clostridium bifermentans		3.26	0.00				
51000050	**Low discrimination**						**51000050**
Clostridium novyi biovar A		56.16	0.50	IND 1			
Clostridium bifermentans		36.80	0.34				
Clostridium botulinum biovar A		7.04	0.23				
51000051	**Good identification**						**51000051**
Clostridium novyi biovar A		97.43	0.50	IND 1			
Clostridium sordellii		1.93	0.04				
Clostridium bifermentans		0.65	0.00				
51000052	**Good identification**						**51000052**
Clostridium novyi biovar A		96.38	0.37	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		3.62	0.00			
51000053	**Good identification**					**51000053**
Clostridium novyi biovar A		100.0	0.37	IND 1		
51000060	**Low discrimination**					**51000060**
Clostridium bifermentans		80.53	0.48	NAG 6		
Clostridium cadaveris		11.76	0.32	SOR 4		
Clostridium novyi biovar A		4.86	0.40			
51000061	**Low discrimination**					**51000061**
Clostridium novyi biovar A		78.69	0.40	IND 1		
Clostridium bifermentans		13.16	0.14			
Clostridium sordellii		6.23	0.04			
51000070	**Low discrimination**					**51000070**
Clostridium novyi biovar A		83.91	0.40	IND 1		
Clostridium bifermentans		14.04	0.14			
Clostridium cadaveris		2.05	-.03			
51000071	**Good identification**					**51000071**
Clostridium novyi biovar A		100.0	0.40	IND 1		
51000100	**Good identification**					**51000100**
Clostridium bifermentans		93.65	0.58	LAC 1		
Clostridium botulinum biovar A		2.50	0.32			
Clostridium sporogenes		1.84	0.27			
51000120	**Low discrimination**					**51000120**
					SPO	
Clostridium bifermentans		61.36	0.38	LAC 1	99	
Clostridium cadaveris		36.95	0.32	NAG 6	99	
Atopobium minutum		1.30	0.17		1	
51000200	**Good identification**					**51000200**
Clostridium bifermentans		93.70	0.58	RHA 1		
Clostridium botulinum biovar A		2.50	0.32			
Clostridium sporogenes		1.84	0.27			
51000220	**Low discrimination**					**51000220**
					SPO	
Clostridium bifermentans		61.60	0.38	RHA 1	99	
Clostridium cadaveris		37.10	0.32	NAG 6	99	
Atopobium minutum		1.31	0.17		1	
51000400	**Good identification**					**51000400**
Clostridium bifermentans		93.70	0.58	ARA 1		
Clostridium botulinum biovar A		2.50	0.32			
Clostridium sporogenes		1.84	0.27			
51000420	**Low discrimination**					**51000420**
					SPO	
Clostridium bifermentans		61.36	0.38	ARA 1	99	
Clostridium cadaveris		36.95	0.32	NAG 6	99	
Atopobium minutum		1.30	0.17		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
51001000	**Low discrimination**						**51001000**
					GRA	SPO	
Clostridium bifermentans		46.51	0.58	GAL 1	99	99	
Fusobacterium necrophorum subsp. necrophorum		43.09	0.51		1	1	
Atopobium minutum		6.25	0.51		99	1	
51001001	**Acceptable identification**						**51001001**
Clostridium sordellii		94.38	0.49	ESL 5			
Clostridium novyi biovar A		1.90	0.37	MLT 90			
Clostridium bifermentans		1.81	0.24				
51001002	**Low discrimination**						**51001002**
					GRA	SPO	
Atopobium minutum		74.02	0.48	IND 1	99	1	
Clostridium bifermentans		8.35	0.24		99	99	
Fusobacterium necrophorum subsp. necrophorum		7.73	0.17		1	1	
51001010	**Low discrimination**						**51001010**
					SPO		
Eubacterium saburreum		34.50	0.40	SUC 99	1		
Clostridium novyi biovar A		22.10	0.37		99		
Clostridium bifermentans		21.06	0.24		99		
51001011	**Low discrimination**						**51001011**
Clostridium novyi biovar A		66.54	0.37	IND 1			
Clostridium sordellii		33.46	0.15	GAL 10			
51001020	**Low discrimination**						**51001020**
					SPO		
Clostridium bifermentans		38.96	0.38	GAL 1	99		
Atopobium minutum		27.34	0.43	NAG 6	1		
Clostridium cadaveris		23.46	0.32		99		
51001022	**Low discrimination**						**51001022**
					GRA	SPO	
Atopobium minutum		76.41	0.40	IND 1	99	1	
Pseudoflavonifractor capillosus		18.14	0.15		1	1	
Clostridium bifermentans		1.65	0.03		99	99	
51001100	**Low discrimination**						**51001100**
					SPO	GRA	
Eubacterium saburreum		68.51	0.47	SUC 99	1	85	
Clostridium bifermentans		14.66	0.24		99	99	
Fusobacterium necrophorum subsp. necrophorum		13.58	0.17		1	1	
51001110	**Good identification**						**51001110**
Eubacterium saburreum		100.0	0.46	SUC 99			
51001520	**Very good identification**						**51001520**
Odoribacter splanchnicus		100.0	0.61	MNS 99			
51001560	**Good identification**						**51001560**
Odoribacter splanchnicus		100.0	0.45	MNS 99			
51002000	**Good identification**						**51002000**
					SPO		
Clostridium bifermentans		90.53	0.58	MAN 1	99		
Eubacterium limosum		2.94	0.29		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		2.42	0.32		99		
51002020	**Low discrimination**						**51002020**
					SPO		
Clostridium bifermentans		61.14	0.38	MAN 1	99		
Clostridium cadaveris		36.82	0.32	NAG 6	99		
Atopobium minutum		1.30	0.17		1		
51004000	**Good identification**						**51004000**
Clostridium bifermentans		92.19	0.58	XYL 1			
Clostridium botulinum biovar A		2.46	0.32				
Clostridium sporogenes		1.81	0.27				
51004020	**Low discrimination**						**51004020**
					SPO		
Clostridium bifermentans		61.60	0.38	XYL 1	99		
Clostridium cadaveris		37.10	0.32	NAG 6	99		
Atopobium minutum		1.31	0.17		1		
51004040	**Low discrimination**						**51004040**
Clostridium glycolicum		86.17	0.49	IND 1			
Clostridium bifermentans		11.46	0.34	FRU 90			
Clostridium botulinum biovar A		2.19	0.23				
51005000	**Low discrimination**						**51005000**
					SPO	GRA	
Eubacterium saburreum		32.47	0.37	SUC 99	1	85	
Clostridium bifermentans		30.12	0.24		99	99	
Fusobacterium necrophorum subsp. necrophorum		27.90	0.17		1	1	
51005010	**Good identification**						**51005010**
Eubacterium saburreum		100.0	0.35	SUC 99			
51005100	**Good identification**						**51005100**
Eubacterium saburreum		100.0	0.43	SUC 99			
51005110	**Good identification**						**51005110**
Eubacterium saburreum		100.0	0.41	SUC 99			
51010000	**Low discrimination**						**51010000**
					GRA	SPO	
Clostridium bifermentans		56.50	0.70	FRU 5	99	99	
Fusobacterium varium		25.81	0.59		1	1	
Eubacterium limosum		5.68	0.49		99	1	
51010001	**Very good identification**						**51010001**
					GRA	SPO	
Clostridium sordellii		96.67	0.62	ESL 5	99	99	
Clostridium bifermentans		1.70	0.36	MLT 90	99	99	
Fusobacterium varium		0.78	0.25		1	1	
51010002	**Low discrimination**						**51010002**
					SPO		
Atopobium minutum		54.26	0.56	IND 1	1		
Clostridium sporogenes		25.01	0.40		99		
Clostridium bifermentans		11.51	0.36		99		
51010010	**Low discrimination**						**51010010**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		67.85	0.45	IND 1	99	99	
Clostridium bifermentans		16.38	0.36	MAN 90	99	99	
Fusobacterium varium		7.48	0.25		1	1	
51010020	**Low discrimination**						**51010020**
					SPO		
Clostridium cadaveris		86.68	0.66	ESL 1	99		
Clostridium bifermentans		7.58	0.50		99		
Atopobium minutum		2.83	0.51		1		
51010022	**Low discrimination**						**51010022**
					GRA	SPO	
Atopobium minutum		61.14	0.48	IND 1	99	1	
Clostridium cadaveris		28.43	0.32		99	99	
Pseudoflavonifractor capillosus		5.24	0.15		1	1	
51010040	**Low discrimination**						**51010040**
Clostridium bifermentans		44.88	0.47	FRU 5			
Clostridium glycolicum		30.68	0.43	SOR 4			
Clostridium botulinum biovar A		10.41	0.36				
51010041	**Acceptable identification**						**51010041**
Clostridium sordellii		91.71	0.38	ESL 5			
Clostridium novyi biovar A		5.20	0.34	MLT 90			
Clostridium bifermentans		1.61	0.13	SOR 4			
51010100	**Low discrimination**						**51010100**
					GRA	SPO	
Clostridium bifermentans		51.73	0.36	FRU 5	99	99	
Fusobacterium varium		23.63	0.25	LAC 1	1	1	
Fusobacterium mortiferum		7.15	0.35		1	1	
51010200	**Low discrimination**						**51010200**
					GRA	SPO	
Clostridium bifermentans		56.79	0.36	FRU 5	99	99	
Fusobacterium varium		25.94	0.25	RHA 1	1	1	
Eubacterium limosum		5.71	0.15		99	1	
51010400	**Low discrimination**						**51010400**
					GRA	SPO	
Clostridium bifermentans		56.79	0.36	FRU 5	99	99	
Fusobacterium varium		25.94	0.25	ARA 1	1	1	
Eubacterium limosum		5.71	0.15		99	1	
51011000	**Low discrimination**						**51011000**
					SPO		
Atopobium minutum		41.63	0.51	IND 1	1		
Eubacterium saburreum		20.37	0.49		1		
Clostridium bifermentans		17.67	0.36		99		
51011001	**Good identification**						**51011001**
					SPO		
Clostridium sordellii		99.33	0.49	ESL 5	99		
Atopobium minutum		0.24	0.17	MLT 90	1		
Clostridium novyi biovar A		0.22	0.20		99		
51011002	**Low discrimination**						**51011002**
					SPO		
Atopobium minutum		86.49	0.48	IND 1	1		
Eubacterium saburreum		11.20	0.36		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sporogenes		1.21	0.06		99		
51011010	**Acceptable identification**						**51011010**
					SPO		
Eubacterium saburreum		90.58	0.48	SUC 99	1		
Clostridium difficile		4.02	0.10		99		
Atopobium minutum		2.29	0.17		1		
51011020	**Low discrimination**						**51011020**
					SPO	CAT	
Propionibacterium acnes		45.51	0.40	NIT 92	1	95	
Atopobium minutum		23.74	0.43	ESL 1	1	1	
Clostridium cadaveris		22.08	0.32		99	1	
51011022	**Low discrimination**						**51011022**
					CAT		
Atopobium minutum		87.70	0.40	IND 1	1		
Eubacterium saburreum		6.01	0.23		1		
Propionibacterium acnes		2.55	0.06		95		
51011100	**Acceptable identification**						**51011100**
					GRA		
Eubacterium saburreum		91.01	0.56	SUC 99	85		
Fusobacterium mortiferum		7.01	0.38		1		
Atopobium minutum		0.81	0.17		99		
51011102	**Acceptable identification**						**51011102**
					GRA		
Eubacterium saburreum		92.43	0.43	SUC 99	85		
Fusobacterium mortiferum		4.49	0.22		1		
Atopobium minutum		3.09	0.14		99		
51011110	**Good identification**						**51011110**
Eubacterium saburreum		100.0	0.54	SUC 99			
51011112	**Good identification**						**51011112**
Eubacterium saburreum		100.0	0.41	SUC 99			
51011130	**Good identification**						**51011130**
Eubacterium saburreum		96.49	0.41	SUC 99			
Odoribacter splanchnicus		3.52	-.03				
51011520	**Very good identification**						**51011520**
Odoribacter splanchnicus		100.0	0.66	MNS 99			
51011560	**Good identification**						**51011560**
Odoribacter splanchnicus		100.0	0.50	MNS 99			
51012000	**Low discrimination**						**51012000**
					SPO		
Eubacterium limosum		85.88	0.63	IND 1	1		
Clostridium difficile		10.72	0.49		99		
Clostridium bifermentans		1.40	0.36		99		
51012010	**Very good identification**						**51012010**
					SPO		
Clostridium difficile		98.33	0.61	IND 1	99		
Eubacterium limosum		1.63	0.29		1		
Clostridium bifermentans		0.03	0.02		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
51012020	**Low discrimination**						**51012020**
					SPO	CAT	
Propionibacterium acnes		34.23	0.36	NIT 92	1	95	
Clostridium cadaveris		29.52	0.32	ESL 1	99	1	
Eubacterium limosum		24.97	0.29		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
51012050	**Good identification**						**51012050**
					SPO		
Clostridium difficile		99.68	0.39	IND 1	99		
Eubacterium limosum		0.32	-.06	SOR 5	1		
51013020	**Good identification**						**51013020**
					SPO	CAT	
Propionibacterium acnes		97.52	0.36	NIT 92	1	95	
Atopobium minutum		0.91	0.09	ESL 1	1	1	
Clostridium cadaveris		0.85	-.03		99	1	
51014000	**Low discrimination**						**51014000**
					GRA	SPO	
Clostridium glycolicum		46.61	0.43	IND 1	99	99	
Fusobacterium varium		31.14	0.35	SOR 95	1	1	
Clostridium bifermentans		16.53	0.36		99	99	
51014040	**Very good identification**						**51014040**
					GRA	SPO	
Clostridium glycolicum		99.87	0.65	IND 1	99	99	
Clostridium bifermentans		0.08	0.13		99	99	
Fusobacterium varium		0.04	0.01		1	1	
51015000	**Low discrimination**						**51015000**
					SPO		
Eubacterium saburreum		62.47	0.45	SUC 99	1		
Clostridium glycolicum		31.53	0.27		99		
Atopobium minutum		2.40	0.17		1		
51015010	**Good identification**						**51015010**
Eubacterium saburreum		100.0	0.43	SUC 99			
51015040	**Good identification**						**51015040**
					SPO		
Clostridium glycolicum		99.90	0.49	IND 1	99		
Eubacterium saburreum		0.11	0.11	GAL 10	1		
51015100	**Good identification**						**51015100**
					GRA		
Eubacterium saburreum		97.59	0.51	SUC 99	85		
Actinomyces meyeri		1.36	0.12		99		
Fusobacterium mortiferum		1.05	0.19		1		
51015102	**Good identification**						**51015102**
Eubacterium saburreum		100.0	0.38	SUC 99			
51015110	**Good identification**						**51015110**
Eubacterium saburreum		100.0	0.49	SUC 99			
51015112	**Good identification**						**51015112**
Eubacterium saburreum		100.0	0.37	SUC 99			
51015120	**Acceptable identification**						**51015120**
Eubacterium saburreum		94.76	0.38	SUC 99			
Odoribacter splanchnicus		5.25	-.03				
51015130	**Good identification**						**51015130**
Eubacterium saburreum		100.0	0.37	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
51016010	**Good identification**						**51016010**
					SPO		
Clostridium difficile		99.68	0.39	IND 1	99		
Eubacterium limosum		0.32	-.06	XYL 5	1		
51020000	**Low discrimination**						**51020000**
Clostridium bifermentans		42.90	0.58	TRE 1			
Clostridium botulinum biovar A		28.34	0.56				
Clostridium sporogenes		27.79	0.53				
51020002	**Acceptable identification**						**51020002**
Clostridium sporogenes		91.72	0.45	IND 1			
Clostridium bifermentans		4.29	0.24				
Clostridium botulinum biovar A		2.84	0.21				
51020020	**Low discrimination**						**51020020**
Clostridium cadaveris		71.87	0.44	ESL 1			
Clostridium bifermentans		22.90	0.38	TRE 5			
Clostridium botulinum biovar A		2.39	0.21				
51020040	**Low discrimination**						**51020040**
Clostridium botulinum biovar A		80.16	0.47	IND 1			
Clostridium bifermentans		16.93	0.34				
Clostridium sporogenes		2.66	0.19				
51021000	**Low discrimination**						**51021000**
					SPO	GRA	
Eubacterium saburreum		28.38	0.38	SUC 99	1	85	
Clostridium bifermentans		21.26	0.24		99	99	
Fusobacterium necrophorum subsp. necrophorum		19.69	0.17		1	1	
51021010	**Good identification**						**51021010**
Eubacterium saburreum		100.0	0.37	SUC 99			
51021100	**Good identification**						**51021100**
Eubacterium saburreum		100.0	0.44	SUC 99			
51021110	**Good identification**						**51021110**
Eubacterium saburreum		100.0	0.43	SUC 99			
51025100	**Good identification**						**51025100**
Eubacterium saburreum		100.0	0.40	SUC 99			
51025110	**Good identification**						**51025110**
Eubacterium saburreum		100.0	0.38	SUC 99			
51030000	**Low discrimination**						**51030000**
Clostridium sporogenes		45.75	0.40	IND 1			
Clostridium bifermentans		21.06	0.36				
Clostridium botulinum biovar A		16.88	0.35				
51030020	**Good identification**						**51030020**
					SPO		
Clostridium cadaveris		96.63	0.44	ESL 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		1.62	0.16	TRE 5	99	
Atopobium minutum		0.60	0.17		1	
51031000	**Low discrimination**					**51031000**
					GRA	
Eubacterium saburreum		89.47	0.46	SUC 99	85	
Fusobacterium mortiferum		3.09	0.23		1	
Atopobium minutum		2.77	0.17		99	
51031010	**Good identification**					**51031010**
Eubacterium saburreum		100.0	0.45	SUC 99		
51031100	**Good identification**					**51031100**
Eubacterium saburreum		96.66	0.53	SUC 99		
Fusobacterium mortiferum		3.34	0.29			
51031102	**Good identification**					**51031102**
Eubacterium saburreum		97.87	0.40	SUC 99		
Fusobacterium mortiferum		2.13	0.13			
51031110	**Good identification**					**51031110**
Eubacterium saburreum		100.0	0.51	SUC 99		
51031112	**Good identification**					**51031112**
Eubacterium saburreum		100.0	0.38	SUC 99		
51031120	**Good identification**					**51031120**
Eubacterium saburreum		95.72	0.40	SUC 99		
Odoribacter splanchnicus		4.28	-.03			
51031130	**Good identification**					**51031130**
Eubacterium saburreum		100.0	0.38	SUC 99		
51032010	**Good identification**					**51032010**
					SPO	
Clostridium difficile		99.78	0.42	IND 1	99	
Eubacterium limosum		0.22	-.06	TRE 7	1	
51035000	**Good identification**					**51035000**
Eubacterium saburreum		100.0	0.42	SUC 99		
51035010	**Good identification**					**51035010**
Eubacterium saburreum		100.0	0.40	SUC 99		
51035100	**Good identification**					**51035100**
Eubacterium saburreum		100.0	0.48	SUC 99		
51035102	**Good identification**					**51035102**
Eubacterium saburreum		100.0	0.35	SUC 99		
51035110	**Good identification**					**51035110**
Eubacterium saburreum		100.0	0.46	SUC 99		
51035120	**Good identification**					**51035120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.35	SUC 99			
51040000	**Good identification**						**51040000**
Clostridium bifermentans		93.70	0.58	CEL 1			
Clostridium botulinum biovar A		2.50	0.32				
Clostridium sporogenes		1.84	0.27				
51040020	**Low discrimination**						**51040020**
					SPO		
Clostridium bifermentans		61.60	0.38	CEL 1	99		
Clostridium cadaveris		37.10	0.32	NAG 6	99		
Atopobium minutum		1.31	0.17		1		
51050000	**Low discrimination**						**51050000**
					GRA	SPO	
Clostridium bifermentans		55.05	0.36	FRU 5	99	99	
Fusobacterium varium		25.14	0.25	CEL 1	1	1	
Eubacterium limosum		5.54	0.15		99	1	
51052010	**Good identification**						**51052010**
					SPO		
Clostridium difficile		99.68	0.39	IND 1	99		
Eubacterium limosum		0.32	-.06	CEL 5	1		
51100000	**Very good identification**						**51100000**
Clostridium bifermentans		97.38	0.99				
Clostridium sporogenes		1.52	0.66				
Clostridium botulinum biovar A		0.91	0.64				
51100001	**Good identification**						**51100001**
Clostridium sordellii		91.54	0.78	ESL 5			
Clostridium bifermentans		7.94	0.64				
Clostridium novyi biovar A		0.32	0.53				
51100002	**Low discrimination**						**51100002**
Clostridium bifermentans		65.16	0.64	BGL 1			
Clostridium sporogenes		33.60	0.58				
Clostridium botulinum biovar A		0.61	0.30				
51100003	**Low discrimination**						**51100003**
Clostridium sordellii		83.80	0.44	ESL 5			
Clostridium bifermentans		7.27	0.30	BGL 1			
Clostridium novyi biovar A		5.19	0.40				
51100010	**Good identification**						**51100010**
Clostridium bifermentans		93.68	0.64	MLZ 1			
Clostridium novyi biovar A		3.83	0.53				
Clostridium sporogenes		1.46	0.32				
51100011	**Low discrimination**						**51100011**
Clostridium sordellii		69.50	0.44	ESL 5			
Clostridium novyi biovar A		24.39	0.53	MLZ 1			
Clostridium bifermentans		6.03	0.30				
51100013	**Low discrimination**						**51100013**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		85.98	0.40	IND 1		
Clostridium sordellii		14.03	0.10			
51100020	**Very good identification**					**51100020**
Clostridium bifermentans		98.09	0.78	NAG 6		
Clostridium cadaveris		1.33	0.44			
Clostridium sporogenes		0.24	0.32			
51100021	**Low discrimination**					**51100021**
Clostridium sordellii		60.81	0.44	ESL 5		
Clostridium bifermentans		33.32	0.44	NAG 1		
Clostridium novyi biovar A		5.33	0.43			
51100022	**Low discrimination**					**51100022**
Clostridium bifermentans		88.20	0.44	NAG 6		
Clostridium sporogenes		7.20	0.24	BGL 1		
Clostridium novyi biovar A		2.49	0.30			
51100030	**Low discrimination**					**51100030**
Clostridium bifermentans		85.02	0.44	MLZ 1		
Clostridium novyi biovar A		13.61	0.43	NAG 6		
Clostridium cadaveris		1.16	0.10			
51100031	**Low discrimination**					**51100031**
Clostridium novyi biovar A		84.87	0.43	IND 1		
Clostridium sordellii		9.77	0.10			
Clostridium bifermentans		5.36	0.10			
51100040	**Good identification**					**51100040**
Clostridium bifermentans		92.54	0.75	SOR 4		
Clostridium botulinum biovar A		6.21	0.55			
Clostridium novyi biovar A		0.61	0.50			
51100041	**Low discrimination**					**51100041**
Clostridium sordellii		86.99	0.55	ESL 5		
Clostridium bifermentans		7.54	0.41	SOR 4		
Clostridium novyi biovar A		4.93	0.50			
51100042	**Low discrimination**					**51100042**
Clostridium bifermentans		76.48	0.41	SOR 4		
Clostridium sporogenes		9.56	0.24	BGL 1		
Clostridium novyi biovar A		8.83	0.37			
51100050	**Low discrimination**					**51100050**
Clostridium bifermentans		57.97	0.41	MLZ 1		
Clostridium novyi biovar A		37.92	0.50	SOR 4		
Clostridium botulinum biovar A		3.89	0.21			
51100051	**Low discrimination**					**51100051**
Clostridium novyi biovar A		83.78	0.50	IND 1		
Clostridium sordellii		14.92	0.20			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		1.29	0.07				
51100052	**Acceptable identification**						**51100052**
Clostridium novyi biovar A		91.95	0.37	IND 1			
Clostridium bifermentans		8.05	0.07				
51100053	**Good identification**						**51100053**
Clostridium novyi biovar A		100.0	0.37	IND 1			
51100060	**Good identification**						**51100060**
Clostridium bifermentans		96.12	0.54	NAG 6			
Clostridium novyi biovar A		2.49	0.40	SOR 4			
Clostridium botulinum biovar A		1.02	0.21				
51100061	**Low discrimination**						**51100061**
Clostridium novyi biovar A		47.55	0.40	IND 1			
Clostridium sordellii		33.88	0.20				
Clostridium bifermentans		18.56	0.20				
51100070	**Low discrimination**						**51100070**
Clostridium novyi biovar A		71.92	0.40	IND 1			
Clostridium bifermentans		28.08	0.20				
51100071	**Good identification**						**51100071**
Clostridium novyi biovar A		100.0	0.40	IND 1			
51100100	**Very good identification**						**51100100**
Clostridium bifermentans		97.30	0.64	LAC 1			
Clostridium sporogenes		1.52	0.32				
Clostridium botulinum biovar A		0.91	0.30				
51100101	**Acceptable identification**						**51100101**
Clostridium sordellii		91.61	0.44	ESL 5			
Clostridium bifermentans		7.94	0.30	LAC 1			
Clostridium novyi biovar A		0.33	0.19				
51100120	**Good identification**						**51100120**
					GRA	SPO	
Clostridium bifermentans		97.43	0.44	LAC 1	99	99	
Clostridium cadaveris		1.32	0.10	NAG 6	99	99	
Prevotella bivia		0.79	-.03		1	1	
51100140	**Acceptable identification**						**51100140**
Clostridium bifermentans		92.81	0.41	LAC 1			
Clostridium botulinum biovar A		6.23	0.21	SOR 4			
Clostridium novyi biovar A		0.61	0.16				
51100200	**Very good identification**						**51100200**
Clostridium bifermentans		97.40	0.64	RHA 1			
Clostridium sporogenes		1.52	0.32				
Clostridium botulinum biovar A		0.91	0.30				
51100201	**Acceptable identification**						**51100201**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sordellii		91.61	0.44	ESL 5		
Clostridium bifermentans		7.94	0.30	RHA 1		
Clostridium novyi biovar A		0.33	0.19			
51100220	**Good identification**					**51100220**
Clostridium bifermentans		98.42	0.44	RHA 1		
Clostridium cadaveris		1.34	0.10	NAG 6		
Clostridium sporogenes		0.24	-.03			
51100240	**Acceptable identification**					**51100240**
Clostridium bifermentans		92.81	0.41	RHA 1		
Clostridium botulinum biovar A		6.23	0.21	SOR 4		
Clostridium novyi biovar A		0.61	0.16			
51100400	**Very good identification**					**51100400**
Clostridium bifermentans		97.40	0.64	ARA 1		
Clostridium sporogenes		1.52	0.32			
Clostridium botulinum biovar A		0.91	0.30			
51100401	**Acceptable identification**					**51100401**
Clostridium sordellii		91.61	0.44	ESL 5		
Clostridium bifermentans		7.94	0.30	ARA 1		
Clostridium novyi biovar A		0.33	0.19			
51100420	**Good identification**					**51100420**
Clostridium bifermentans		98.42	0.44	ARA 1		
Clostridium cadaveris		1.34	0.10	NAG 6		
Clostridium sporogenes		0.24	-.03			
51100440	**Acceptable identification**					**51100440**
Clostridium bifermentans		92.81	0.41	ARA 1		
Clostridium botulinum biovar A		6.23	0.21	SOR 4		
Clostridium novyi biovar A		0.61	0.16			
51101000	**Good identification**					**51101000**
					SPO	
Clostridium bifermentans		91.68	0.64	GAL 1	99	
Eubacterium saburreum		3.18	0.51		1	
Clostridium sordellii		1.89	0.31		99	
51101001	**Very good identification**					**51101001**
Clostridium sordellii		99.26	0.65	ESL 5		
Clostridium bifermentans		0.49	0.30			
Clostridium novyi biovar A		0.22	0.37			
51101010	**Low discrimination**					**51101010**
					SPO	
Eubacterium saburreum		65.40	0.50	SUC 99	1	
Clostridium bifermentans		23.29	0.30		99	
Clostridium novyi biovar A		10.47	0.37		99	
51101012	**Low discrimination**					**51101012**
					SPO	
Eubacterium saburreum		86.20	0.37	SUC 99	1	
Clostridium novyi biovar A		13.80	0.24		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
51101020	**Low discrimination**						**51101020**
					SPO		
Clostridium bifermentans		88.03	0.44	GAL 1	99		
Eubacterium saburreum		8.44	0.39	NAG 6	1		
Clostridium novyi biovar A		1.57	0.26		99		
51101030	**Low discrimination**						**51101030**
					SPO		
Eubacterium saburreum		73.77	0.37	SUC 99	1		
Clostridium novyi biovar A		16.73	0.26		99		
Clostridium bifermentans		9.50	0.10		99		
51101040	**Low discrimination**						**51101040**
Clostridium bifermentans		83.83	0.41	GAL 1			
Clostridium novyi biovar A		6.09	0.34	SOR 4			
Clostridium botulinum biovar A		5.62	0.21				
51101041	**Good identification**						**51101041**
Clostridium sordellii		96.09	0.42	ESL 5			
Clostridium novyi biovar A		3.43	0.34	SOR 4			
Clostridium bifermentans		0.48	0.07				
51101100	**Low discrimination**						**51101100**
					COC	SPO	
Eubacterium saburreum		72.54	0.58	SUC 99	1	1	
Blautia hansenii		18.07	0.32		99	1	
Clostridium bifermentans		9.06	0.30		1	99	
51101102	**Low discrimination**						**51101102**
					COC		
Eubacterium saburreum		86.44	0.45	SUC 99	1		
Blautia hansenii		13.56	0.15		99		
51101110	**Very good identification**						**51101110**
					COC		
Eubacterium saburreum		99.69	0.56	SUC 99	1		
Blautia hansenii		0.31	-.03		99		
51101112	**Good identification**						**51101112**
Eubacterium saburreum		100.0	0.43	SUC 99			
51101120	**Low discrimination**						**51101120**
					COC	SPO	
Eubacterium saburreum		82.37	0.45	SUC 99	1	1	
Blautia hansenii		12.92	0.15		99	1	
Clostridium bifermentans		3.72	0.10		1	99	
51101130	**Good identification**						**51101130**
Eubacterium saburreum		100.0	0.43	SUC 99			
51101722	**Good identification**						**51101722**
Bacteroides eggerthii		100.0	0.36	MNS 99			
51102000	**Very good identification**						**51102000**
Clostridium bifermentans		97.36	0.64	MAN 1			
Clostridium sporogenes		1.52	0.32				
Clostridium botulinum biovar A		0.91	0.30				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
51102001	**Acceptable identification**					**51102001**
Clostridium sordellii		91.61	0.44	ESL 5		
Clostridium bifermentans		7.94	0.30	MAN 1		
Clostridium novyi biovar A		0.33	0.19			
51102020	**Good identification**					**51102020**
Clostridium bifermentans		98.42	0.44	MAN 1		
Clostridium cadaveris		1.34	0.10	NAG 6		
Clostridium sporogenes		0.24	-.03			
51102040	**Acceptable identification**					**51102040**
Clostridium bifermentans		92.50	0.41	MAN 1		
Clostridium botulinum biovar A		6.21	0.21	SOR 4		
Clostridium novyi biovar A		0.61	0.16			
51104000	**Very good identification**					**51104000**
Clostridium bifermentans		96.66	0.64	XYL 1		
Clostridium sporogenes		1.51	0.32			
Clostridium botulinum biovar A		0.91	0.30			
51104001	**Acceptable identification**					**51104001**
Clostridium sordellii		91.61	0.44	ESL 5		
Clostridium bifermentans		7.94	0.30	XYL 1		
Clostridium novyi biovar A		0.33	0.19			
51104020	**Good identification**					**51104020**
Clostridium bifermentans		98.42	0.44	XYL 1		
Clostridium cadaveris		1.34	0.10	NAG 6		
Clostridium sporogenes		0.24	-.03			
51104040	**Low discrimination**					**51104040**
Clostridium glycolicum		76.42	0.50	IND 1		
Clostridium bifermentans		21.89	0.41	FRU 90		
Clostridium botulinum biovar A		1.47	0.21			
51105000	**Low discrimination**					**51105000**
					SPO	
Eubacterium saburreum		62.26	0.47	SUC 99	1	
Clostridium bifermentans		33.69	0.30		99	
Clostridium glycolicum		2.84	0.11		99	
51105010	**Good identification**					**51105010**
Eubacterium saburreum		100.0	0.45	SUC 99		
51105100	**Acceptable identification**					**51105100**
					COC	
Eubacterium saburreum		92.45	0.53	SUC 99	1	
Blautia hansenii		7.55	0.19		99	
51105102	**Good identification**					**51105102**
					COC	
Eubacterium saburreum		95.11	0.40	SUC 99	1	
Blautia hansenii		4.89	0.03		99	
51105110	**Good identification**					**51105110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.52	SUC 99		
51105112	**Good identification**					**51105112**
Eubacterium saburreum		100.0	0.39	SUC 99		
51105120	**Good identification**					**51105120**
					COC	
Eubacterium saburreum		95.11	0.40	SUC 99	1	
Blautia hansenii		4.89	0.03		99	
51105130	**Good identification**					**51105130**
Eubacterium saburreum		100.0	0.39	SUC 99		
51105722	**Good identification**					**51105722**
Bacteroides eggerthii		100.0	0.56	MNS 99		
51110000	**Low discrimination**					**51110000**
Clostridium bifermentans		89.86	0.77	FRU 5		
Clostridium sporogenes		4.71	0.53			
Clostridium sordellii		2.01	0.44			
51110001	**Very good identification**					**51110001**
Clostridium sordellii		99.47	0.78	ESL 5		
Clostridium bifermentans		0.45	0.43			
Clostridium novyi biovar A		0.04	0.37			
51110002	**Low discrimination**					**51110002**
Clostridium sporogenes		61.08	0.45	IND 1		
Clostridium bifermentans		35.33	0.43			
Clostridium sordellii		0.79	0.10			
51110003	**Good identification**					**51110003**
Clostridium sordellii		98.10	0.44	ESL 5		
Clostridium sporogenes		0.77	0.11	BGL 1		
Clostridium novyi biovar A		0.68	0.24			
51110010	**Low discrimination**					**51110010**
Clostridium bifermentans		81.55	0.43	FRU 5		
Clostridium novyi biovar A		7.04	0.37	MLZ 1		
Clostridium sporogenes		4.27	0.19			
51110011	**Good identification**					**51110011**
Clostridium sordellii		95.83	0.44	ESL 5		
Clostridium novyi biovar A		3.74	0.37	MLZ 1		
Clostridium bifermentans		0.44	0.08			
51110020	**Low discrimination**					**51110020**
Clostridium bifermentans		78.04	0.56	FRU 5		
Clostridium cadaveris		20.14	0.44	NAG 6		
Clostridium sporogenes		0.65	0.19			
51110021	**Good identification**					**51110021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium sordellii		95.60	0.44	ESL 5			
Clostridium bifermentans		2.76	0.22	NAG 1			
Clostridium novyi biovar A		0.93	0.26				
51110040	**Low discrimination**						**51110040**
Clostridium bifermentans		68.23	0.53	FRU 5			
Clostridium glycolicum		21.66	0.44	SOR 4			
Clostridium botulinum biovar A		5.55	0.35				
51110041	**Good identification**						**51110041**
Clostridium sordellii		98.75	0.55	ESL 5			
Clostridium novyi biovar A		0.62	0.34	SOR 4			
Clostridium bifermentans		0.45	0.19				
51110100	**Low discrimination**						**51110100**
					SPO		
Clostridium bifermentans		85.99	0.43	FRU 5	99		
Clostridium sporogenes		4.50	0.19	LAC 1	99		
Eubacterium saburreum		4.01	0.32		1		
51110101	**Good identification**						**51110101**
Clostridium sordellii		99.55	0.44	ESL 5			
Clostridium bifermentans		0.45	0.08	LAC 1			
51110200	**Acceptable identification**						**51110200**
Clostridium bifermentans		90.44	0.43	FRU 5			
Clostridium sporogenes		4.74	0.19	RHA 1			
Clostridium sordellii		2.02	0.10				
51110201	**Good identification**						**51110201**
Clostridium sordellii		99.55	0.44	ESL 5			
Clostridium bifermentans		0.45	0.08	RHA 1			
51110400	**Acceptable identification**						**51110400**
Clostridium bifermentans		90.44	0.43	FRU 5			
Clostridium sporogenes		4.74	0.19	ARA 1			
Clostridium sordellii		2.02	0.10				
51110401	**Good identification**						**51110401**
Clostridium sordellii		99.55	0.44	ESL 5			
Clostridium bifermentans		0.45	0.08	ARA 1			
51111000	**Low discrimination**						**51111000**
					SPO	GRA	
Eubacterium saburreum		36.57	0.60	SUC 99	1	85	
Prevotella intermedia		35.92	0.50		1	1	
Clostridium bifermentans		18.50	0.43		99	99	
51111001	**Very good identification**						**51111001**
					SPO	GRA	
Clostridium sordellii		99.85	0.65	ESL 5	99	99	
Eubacterium saburreum		0.05	0.25		1	85	
Prevotella intermedia		0.05	0.16		1	1	
51111002	**Low discrimination**						**51111002**
					SPO	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		85.76	0.47	SUC 99	1	85	
Prevotella intermedia		4.82	0.16		1	1	
Clostridium sporogenes		4.29	0.11		99	99	
51111010	**Very good identification**						**51111010**
					SPO	GRA	
Eubacterium saburreum		97.40	0.58	SUC 99	1	85	
Prevotella intermedia		1.18	0.16		1	1	
Clostridium bifermentans		0.61	0.08		99	99	
51111012	**Good identification**						**51111012**
Eubacterium saburreum		100.0	0.45	SUC 99			
51111020	**Low discrimination**						**51111020**
					SPO	GRA	
Eubacterium saburreum		75.01	0.47	SUC 99	1	85	
Clostridium bifermentans		13.73	0.22		99	99	
Prevotella intermedia		4.22	0.16		1	1	
51111030	**Good identification**						**51111030**
					SPO		
Eubacterium saburreum		99.17	0.45	SUC 99	1		
Clostridium novyi biovar A		0.83	0.10		99		
51111041	**Good identification**						**51111041**
Clostridium sordellii		99.61	0.42	ESL 5			
Clostridium novyi biovar A		0.40	0.17	SOR 4			
51111100	**Very good identification**						**51111100**
Eubacterium saburreum		98.79	0.66	SUC 99			
Prevotella intermedia		0.42	0.16				
Fusobacterium mortiferum		0.24	0.23				
51111102	**Good identification**						**51111102**
Eubacterium saburreum		99.66	0.53	SUC 99			
Lactobacillus fermentum		0.34	0.14				
51111110	**Very good identification**						**51111110**
Eubacterium saburreum		100.0	0.64	SUC 99			
51111112	**Good identification**						**51111112**
Eubacterium saburreum		100.0	0.52	SUC 99			
51111120	**Good identification**						**51111120**
					SPO	GRA	
Eubacterium saburreum		97.65	0.53	SUC 99	1	85	
Clostridium septicum		1.23	0.05		99	99	
Odoribacter splanchnicus		0.73	-.03		1	1	
51111122	**Good identification**						**51111122**
Eubacterium saburreum		100.0	0.40	SUC 99			
51111130	**Good identification**						**51111130**
Eubacterium saburreum		100.0	0.52	SUC 99			
51111132	**Good identification**						**51111132**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium saburreum		100.0	0.39	SUC 99				
51112000	**Low discrimination**							**51112000**
					COC	SPO	GRA	
Clostridium bifermentans		67.22	0.43	FRU 5	1	99	99	
Eubacterium limosum		17.80	0.29	MAN 1	1	1	99	
Megasphaera elsdenii		5.47	0.29		99	1	1	
51112001	**Good identification**							**51112001**
					SPO			
Clostridium sordellii		99.43	0.44	ESL 5	99			
Clostridium bifermentans		0.45	0.08	MAN 1	99			
Eubacterium limosum		0.12	-.06		1			
51114000	**Low discrimination**							**51114000**
Clostridium glycolicum		53.79	0.44	IND 1				
Clostridium bifermentans		41.09	0.43	SOR 95				
Clostridium sporogenes		2.15	0.19					
51114001	**Good identification**							**51114001**
Clostridium sordellii		98.96	0.44	ESL 5				
Clostridium glycolicum		0.59	0.10	XYL 1				
Clostridium bifermentans		0.45	0.08					
51114040	**Very good identification**							**51114040**
Clostridium glycolicum		99.82	0.66	IND 1				
Clostridium bifermentans		0.17	0.19					
Clostridium botulinum biovar A		0.01	0.01					
51115000	**Low discrimination**							**51115000**
					SPO	GRA		
Eubacterium saburreum		85.43	0.55	SUC 99	1	85		
Clostridium glycolicum		11.68	0.28		99	99		
Prevotella intermedia		1.57	0.16		1	1		
51115002	**Good identification**							**51115002**
Eubacterium saburreum		100.0	0.42	SUC 99				
51115010	**Good identification**							**51115010**
Eubacterium saburreum		100.0	0.53	SUC 99				
51115012	**Good identification**							**51115012**
Eubacterium saburreum		100.0	0.41	SUC 99				
51115020	**Good identification**							**51115020**
Eubacterium saburreum		100.0	0.42	SUC 99				
51115030	**Good identification**							**51115030**
Eubacterium saburreum		100.0	0.41	SUC 99				
51115040	**Good identification**							**51115040**
					SPO			
Clostridium glycolicum		99.61	0.50	IND 1	99			
Eubacterium saburreum		0.39	0.21	GAL 10	1			
51115100	**Very good identification**							**51115100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.61	0.61	SUC 99		
Actinomyces meyeri		0.39	0.13			
51115102	**Good identification**					**51115102**
Eubacterium saburreum		100.0	0.48	SUC 99		
51115110	**Very good identification**					**51115110**
Eubacterium saburreum		100.0	0.60	SUC 99		
51115112	**Good identification**					**51115112**
Eubacterium saburreum		100.0	0.47	SUC 99		
51115120	**Good identification**					**51115120**
Eubacterium saburreum		100.0	0.48	SUC 99		
51115122	**Good identification**					**51115122**
Eubacterium saburreum		100.0	0.35	SUC 99		
51115130	**Good identification**					**51115130**
Eubacterium saburreum		100.0	0.47	SUC 99		
51120000	**Low discrimination**					**51120000**
Clostridium bifermentans		57.17	0.64	TRE 1		
Clostridium sporogenes		29.47	0.58			
Clostridium botulinum biovar A		13.24	0.54			
51120001	**Low discrimination**					**51120001**
Clostridium sordellii		86.58	0.44	ESL 5		
Clostridium bifermentans		7.51	0.30	TRE 1		
Clostridium sporogenes		3.87	0.24			
51120002	**Acceptable identification**					**51120002**
Clostridium sporogenes		93.25	0.50	IND 1		
Clostridium bifermentans		5.48	0.30			
Clostridium botulinum biovar A		1.27	0.20			
51120020	**Low discrimination**					**51120020**
Clostridium bifermentans		84.10	0.44	TRE 1		
Clostridium sporogenes		6.86	0.24	NAG 6		
Clostridium cadaveris		5.95	0.22			
51120040	**Low discrimination**					**51120040**
Clostridium botulinum biovar A		59.47	0.45	IND 1		
Clostridium bifermentans		35.81	0.41			
Clostridium sporogenes		4.48	0.24			
51121000	**Low discrimination**					**51121000**
					SPO	
Eubacterium saburreum		56.42	0.48	SUC 99	1	
Clostridium bifermentans		24.65	0.30		99	
Clostridium sporogenes		12.71	0.24		99	
51121002	**Low discrimination**					**51121002**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		70.15	0.36	SUC 99	1	
Clostridium sporogenes		29.85	0.15		99	
51121010	**Good identification**					**51121010**
Eubacterium saburreum		100.0	0.47	SUC 99		
51121020	**Low discrimination**					**51121020**
					SPO	
Eubacterium saburreum		86.35	0.36	SUC 99	1	
Clostridium bifermentans		13.65	0.10		99	
51121100	**Good identification**					**51121100**
					COC	
Eubacterium saburreum		99.62	0.55	SUC 99	1	
Blautia hansenii		0.38	-.03		99	
51121102	**Good identification**					**51121102**
Eubacterium saburreum		100.0	0.42	SUC 99		
51121110	**Good identification**					**51121110**
Eubacterium saburreum		100.0	0.53	SUC 99		
51121112	**Good identification**					**51121112**
Eubacterium saburreum		100.0	0.40	SUC 99		
51121120	**Good identification**					**51121120**
Eubacterium saburreum		100.0	0.42	SUC 99		
51121130	**Good identification**					**51121130**
Eubacterium saburreum		100.0	0.40	SUC 99		
51125000	**Good identification**					**51125000**
Eubacterium saburreum		100.0	0.44	SUC 99		
51125010	**Good identification**					**51125010**
Eubacterium saburreum		100.0	0.42	SUC 99		
51125100	**Good identification**					**51125100**
Eubacterium saburreum		100.0	0.50	SUC 99		
51125102	**Good identification**					**51125102**
Eubacterium saburreum		100.0	0.37	SUC 99		
51125110	**Good identification**					**51125110**
Eubacterium saburreum		100.0	0.49	SUC 99		
51125112	**Good identification**					**51125112**
Eubacterium saburreum		100.0	0.36	SUC 99		
51125120	**Good identification**					**51125120**
Eubacterium saburreum		100.0	0.37	SUC 99		
51125130	**Good identification**					**51125130**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.36	SUC 99			
51130000	**Low discrimination**						**51130000**
Clostridium sporogenes		56.40	0.45	IND 1			
Clostridium bifermentans		32.63	0.43				
Clostridium botulinum biovar A		9.17	0.34				
51130001	**Good identification**						**51130001**
Clostridium sordellii		98.65	0.44	ESL 5			
Clostridium sporogenes		0.78	0.11	TRE 1			
Clostridium bifermentans		0.45	0.08				
51130002	**Good identification**						**51130002**
Clostridium sporogenes		97.42	0.37	IND 1			
Clostridium bifermentans		1.71	0.08				
Clostridium botulinum biovar A		0.48	-.01				
51131000	**Very good identification**						**51131000**
					SPO	GRA	
Eubacterium saburreum		96.07	0.57	SUC 99	1	85	
Prevotella intermedia		1.43	0.16		1	1	
Clostridium sporogenes		1.27	0.11		99	99	
51131002	**Good identification**						**51131002**
					SPO		
Eubacterium saburreum		97.56	0.44	SUC 99	1		
Clostridium sporogenes		2.44	0.03		99		
51131010	**Good identification**						**51131010**
Eubacterium saburreum		100.0	0.55	SUC 99			
51131012	**Good identification**						**51131012**
Eubacterium saburreum		100.0	0.42	SUC 99			
51131020	**Good identification**						**51131020**
Eubacterium saburreum		100.0	0.44	SUC 99			
51131030	**Good identification**						**51131030**
Eubacterium saburreum		100.0	0.42	SUC 99			
51131100	**Very good identification**						**51131100**
Eubacterium saburreum		99.89	0.63	SUC 99			
Fusobacterium mortiferum		0.11	0.14				
51131102	**Good identification**						**51131102**
Eubacterium saburreum		100.0	0.50	SUC 99			
51131110	**Very good identification**						**51131110**
Eubacterium saburreum		100.0	0.61	SUC 99			
51131112	**Good identification**						**51131112**
Eubacterium saburreum		100.0	0.48	SUC 99			
51131120	**Acceptable identification**						**51131120**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		94.01	0.50	SUC 99	1	
Clostridium septicum		5.32	0.13		99	
Clostridium perfringens		0.67	0.04		84	
51131122	**Good identification**					**51131122**
Eubacterium saburreum		100.0	0.37	SUC 99		
51131130	**Good identification**					**51131130**
Eubacterium saburreum		100.0	0.48	SUC 99		
51131132	**Good identification**					**51131132**
Eubacterium saburreum		100.0	0.36	SUC 99		
51135000	**Good identification**					**51135000**
Eubacterium saburreum		100.0	0.52	SUC 99		
51135002	**Good identification**					**51135002**
Eubacterium saburreum		100.0	0.39	SUC 99		
51135010	**Good identification**					**51135010**
Eubacterium saburreum		100.0	0.50	SUC 99		
51135012	**Good identification**					**51135012**
Eubacterium saburreum		100.0	0.38	SUC 99		
51135020	**Good identification**					**51135020**
Eubacterium saburreum		100.0	0.39	SUC 99		
51135030	**Good identification**					**51135030**
Eubacterium saburreum		100.0	0.38	SUC 99		
51135100	**Very good identification**					**51135100**
					COC	
Eubacterium saburreum		99.86	0.58	SUC 99	1	
Atopobium parvulum		0.14	0.00		99	
51135102	**Good identification**					**51135102**
Eubacterium saburreum		100.0	0.45	SUC 99		
51135110	**Very good identification**					**51135110**
Eubacterium saburreum		100.0	0.57	SUC 99		
51135112	**Good identification**					**51135112**
Eubacterium saburreum		100.0	0.44	SUC 99		
51135120	**Good identification**					**51135120**
Eubacterium saburreum		100.0	0.45	SUC 99		
51135130	**Good identification**					**51135130**
Eubacterium saburreum		100.0	0.44	SUC 99		
51140000	**Very good identification**					**51140000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		97.40	0.64	CEL 1		
Clostridium sporogenes		1.52	0.32			
Clostridium botulinum biovar A		0.91	0.30			
51140001	**Acceptable identification**					**51140001**
Clostridium sordellii		91.61	0.44	ESL 5		
Clostridium bifermentans		7.94	0.30	CEL 1		
Clostridium novyi biovar A		0.33	0.19			
51140020	**Good identification**					**51140020**
Clostridium bifermentans		98.42	0.44	CEL 1		
Clostridium cadaveris		1.34	0.10	NAG 6		
Clostridium sporogenes		0.24	-.03			
51140040	**Acceptable identification**					**51140040**
Clostridium bifermentans		92.81	0.41	CEL 1		
Clostridium botulinum biovar A		6.23	0.21	SOR 4		
Clostridium novyi biovar A		0.61	0.16			
51141722	**Good identification**					**51141722**
Bacteroides eggerthii		100.0	0.47	MNS 99		
51145722	**Very good identification**					**51145722**
Bacteroides eggerthii		100.0	0.66	MNS 99		
51150000	**Acceptable identification**					**51150000**
Clostridium bifermentans		90.44	0.43	FRU 5		
Clostridium sporogenes		4.74	0.19	CEL 1		
Clostridium sordellii		2.02	0.10			
51150001	**Good identification**					**51150001**
Clostridium sordellii		99.55	0.44	ESL 5		
Clostridium bifermentans		0.45	0.08	CEL 1		
51200000	**Very good identification**					**51200000**
Clostridium bifermentans		99.32	0.76	SAL 10		
Clostridium botulinum biovar A		0.24	0.32			
Clostridium sporogenes		0.18	0.27			
51200001	**Low discrimination**					**51200001**
Clostridium bifermentans		74.27	0.42	SAL 10		
Clostridium sordellii		20.19	0.28	URE 1		
Clostridium novyi biovar A		5.54	0.35			
51200002	**Acceptable identification**					**51200002**
					SPO	
Clostridium bifermentans		91.20	0.42	SAL 10	99	
Clostridium sporogenes		5.37	0.19	BGL 1	99	
Atopobium minutum		2.23	0.22		1	
51200010	**Acceptable identification**					**51200010**
Clostridium bifermentans		92.57	0.42	SAL 10		
Clostridium novyi biovar A		6.91	0.35	MLZ 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		0.52	0.01				
51200020	**Acceptable identification**						**51200020**
Clostridium bifermentans		94.37	0.55	SAL 10			
Clostridium cadaveris		5.17	0.32	NAG 6			
Clostridium novyi biovar A		0.28	0.25				
51200040	**Good identification**						**51200040**
Clostridium bifermentans		97.14	0.52	SAL 10			
Clostridium botulinum biovar A		1.69	0.23	SOR 4			
Clostridium novyi biovar A		1.17	0.32				
51200100	**Good identification**						**51200100**
Clostridium bifermentans		100.0	0.42	SAL 10			
51200200	**Good identification**						**51200200**
Clostridium bifermentans		100.0	0.42	SAL 10			
51200400	**Good identification**						**51200400**
Clostridium bifermentans		100.0	0.42	SAL 10			
51201000	**Low discrimination**						**51201000**
					GRA	SPO	
Clostridium bifermentans		66.99	0.42	SAL 10	99	99	
Eubacterium tortuosum		25.99	0.32	GAL 1	99	1	
Fusobacterium necrophorum subsp. necrophorum		5.64	0.17		1	1	
51202000	**Good identification**						**51202000**
					SPO		
Clostridium bifermentans		98.69	0.42	SAL 10	99		
Clostridium difficile		1.02	0.06	MAN 1	99		
Eubacterium limosum		0.29	-.06		1		
51204000	**Good identification**						**51204000**
Clostridium bifermentans		100.0	0.42	SAL 10			
51210000	**Acceptable identification**						**51210000**
					GRA	SPO	
Clostridium bifermentans		90.73	0.54	SAL 10	99	99	
Fusobacterium varium		3.77	0.25	FRU 5	1	1	
Clostridium difficile		1.97	0.23		99	99	
51210010	**Acceptable identification**						**51210010**
Clostridium difficile		90.08	0.35	IND 1			
Clostridium bifermentans		8.57	0.20	MAN 90			
Clostridium novyi biovar A		1.35	0.19				
51210100	**Low discrimination**						**51210100**
					GRA	SPO	
Fusobacterium mortiferum		70.71	0.40	IND 1	1	1	
Clostridium bifermentans		29.29	0.20		99	99	
51211000	**Low discrimination**						**51211000**
					GRA	SPO	
Fusobacterium mortiferum		52.69	0.37	IND 1	1	1	
Clostridium bifermentans		33.95	0.20		99	99	
Atopobium minutum		7.27	0.17		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
51211100	**Acceptable identification**					**51211100**
Fusobacterium mortiferum		93.68	0.43	IND 1		
Eubacterium saburreum		6.33	0.21			
51212000	**Low discrimination**					**51212000**
					SPO	
Clostridium difficile		74.55	0.40	IND 1	99	
Eubacterium limosum		21.38	0.29		1	
Clostridium bifermentans		3.85	0.20		99	
51212010	**Good identification**					**51212010**
					SPO	
Clostridium difficile		99.94	0.52	IND 1	99	
Eubacterium limosum		0.06	-.06		1	
51220000	**Low discrimination**					**51220000**
Clostridium bifermentans		89.37	0.42	SAL 10		
Clostridium botulinum biovar A		5.37	0.21	TRE 1		
Clostridium sporogenes		5.26	0.19			
51240000	**Good identification**					**51240000**
Clostridium bifermentans		100.0	0.42	SAL 10		
51300000	**Very good identification**					**51300000**
Clostridium bifermentans		99.73	0.82	SAL 10		
Clostridium sporogenes		0.14	0.32			
Clostridium botulinum biovar A		0.09	0.30			
51300001	**Low discrimination**					**51300001**
Clostridium sordellii		50.36	0.44	ESL 5		
Clostridium bifermentans		48.03	0.48	SAL 1		
Clostridium novyi biovar A		1.54	0.35			
51300002	**Good identification**					**51300002**
Clostridium bifermentans		95.01	0.48	SAL 10		
Clostridium sporogenes		4.45	0.24	BGL 1		
Clostridium novyi biovar A		0.54	0.22			
51300010	**Good identification**					**51300010**
Clostridium bifermentans		96.77	0.48	SAL 10		
Clostridium novyi biovar A		3.10	0.35	MLZ 1		
Clostridium sporogenes		0.14	-.03			
51300020	**Very good identification**					**51300020**
Clostridium bifermentans		99.73	0.62	SAL 10		
Clostridium novyi biovar A		0.13	0.25	NAG 6		
Clostridium cadaveris		0.12	0.10			
51300040	**Very good identification**					**51300040**
Clostridium bifermentans		98.85	0.59	SAL 10		
Clostridium botulinum biovar A		0.60	0.21	SOR 4		
Clostridium novyi biovar A		0.51	0.32			
51300060	**Good identification**					**51300060**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		98.02	0.38	SAL 10		
Clostridium novyi biovar A		1.99	0.22	NAG 6		
51300100	**Good identification**					**51300100**
Clostridium bifermentans		99.86	0.48	SAL 10		
Clostridium sporogenes		0.14	-.03	LAC 1		
51300200	**Good identification**					**51300200**
Clostridium bifermentans		99.86	0.48	SAL 10		
Clostridium sporogenes		0.14	-.03	RHA 1		
51300400	**Good identification**					**51300400**
Clostridium bifermentans		99.86	0.48	SAL 10		
Clostridium sporogenes		0.14	-.03	ARA 1		
51301000	**Good identification**					**51301000**
					SPO	
Clostridium bifermentans		98.85	0.48	SAL 10	99	
Clostridium novyi biovar A		0.35	0.19	GAL 1	99	
Eubacterium saburreum		0.31	0.17		1	
51302000	**Good identification**					**51302000**
Clostridium bifermentans		99.86	0.48	SAL 10		
Clostridium sporogenes		0.14	-.03	MAN 1		
51304000	**Good identification**					**51304000**
Clostridium bifermentans		99.86	0.48	SAL 10		
Clostridium sporogenes		0.14	-.03	XYL 1		
51310000	**Very good identification**					**51310000**
Clostridium bifermentans		98.93	0.60	SAL 10		
Clostridium sporogenes		0.47	0.19	FRU 5		
Clostridium sordellii		0.20	0.10			
51310001	**Acceptable identification**					**51310001**
Clostridium sordellii		94.91	0.44	ESL 5		
Clostridium bifermentans		4.76	0.26	SAL 1		
Clostridium novyi biovar A		0.32	0.19			
51310020	**Good identification**					**51310020**
Clostridium bifermentans		97.71	0.40	SAL 10		
Clostridium cadaveris		2.29	0.10	FRU 5		
51310040	**Good identification**					**51310040**
Clostridium bifermentans		95.50	0.37	SAL 10		
Clostridium glycolicum		2.76	0.10	FRU 5		
Clostridium novyi biovar A		1.04	0.15	SOR 4		
51320000	**Acceptable identification**					**51320000**
Clostridium bifermentans		93.64	0.48	SAL 10		
Clostridium sporogenes		4.39	0.24	TRE 1		
Clostridium botulinum biovar A		1.97	0.20			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
51340000	**Good identification**						**51340000**
Clostridium bifermentans		99.86	0.48	SAL 10			
Clostridium sporogenes		0.14	-.03	CEL 1			
51400000	**Good identification**						**51400000**
Clostridium bifermentans		93.70	0.58	RAF 1			
Clostridium botulinum biovar A		2.50	0.32				
Clostridium sporogenes		1.84	0.27				
51400020	**Low discrimination**						**51400020**
					SPO		
Clostridium bifermentans		61.60	0.38	RAF 1	99		
Clostridium cadaveris		37.10	0.32	NAG 6	99		
Atopobium minutum		1.31	0.17		1		
51401000	**Low discrimination**						**51401000**
					SPO	GRA	
Eubacterium saburreum		33.84	0.37	SUC 99	1	85	
Clostridium bifermentans		31.39	0.24		99	99	
Fusobacterium necrophorum subsp. necrophorum		29.08	0.17		1	1	
51401010	**Good identification**						**51401010**
Eubacterium saburreum		100.0	0.35	SUC 99			
51401110	**Good identification**						**51401110**
					COC		
Eubacterium saburreum		97.77	0.41	SUC 99	1		
Blautia hansenii		2.23	-.03		99		
51405100	**Low discrimination**						**51405100**
					COC		
Eubacterium saburreum		62.25	0.38	SUC 99	1		
Blautia hansenii		37.75	0.19		99		
51405110	**Good identification**						**51405110**
Eubacterium saburreum		100.0	0.37	SUC 99			
51410000	**Low discrimination**						**51410000**
					GRA	SPO	
Clostridium bifermentans		51.26	0.36	RAF 1	99	99	
Fusobacterium varium		23.42	0.25	FRU 5	1	1	
Fusobacterium mortiferum		5.89	0.34		1	1	
51410100	**Acceptable identification**						**51410100**
					SPO	GRA	
Fusobacterium mortiferum		91.56	0.40	IND 1	1	1	
Eubacterium saburreum		4.99	0.17		1	85	
Clostridium bifermentans		3.45	0.02		99	99	
51411000	**Low discrimination**						**51411000**
Eubacterium saburreum		69.30	0.45	SUC 99			
Fusobacterium mortiferum		19.25	0.37				
Prevotella intermedia		7.66	0.19				
51411010	**Good identification**						**51411010**
Eubacterium saburreum		100.0	0.43	SUC 99			
51411100	**Low discrimination**						**51411100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Eubacterium saburreum		77.88	0.51	SUC 99	1	85	
Fusobacterium mortiferum		21.64	0.43		1	1	
Blautia hansenii		0.49	-.03		99	99	
51411102	**Low discrimination**						**51411102**
Eubacterium saburreum		85.11	0.38	SUC 99			
Fusobacterium mortiferum		14.89	0.27				
51411110	**Good identification**						**51411110**
Eubacterium saburreum		100.0	0.49	SUC 99			
51411112	**Good identification**						**51411112**
Eubacterium saburreum		100.0	0.37	SUC 99			
51411120	**Acceptable identification**						**51411120**
Eubacterium saburreum		94.76	0.38	SUC 99			
Odoribacter splanchnicus		5.25	-.03				
51411130	**Good identification**						**51411130**
Eubacterium saburreum		100.0	0.37	SUC 99			
51415000	**Good identification**						**51415000**
Eubacterium saburreum		96.26	0.40	SUC 99			
Fusobacterium mortiferum		3.74	0.18				
51415010	**Good identification**						**51415010**
Eubacterium saburreum		100.0	0.39	SUC 99			
51415100	**Good identification**						**51415100**
Eubacterium saburreum		96.26	0.46	SUC 99			
Fusobacterium mortiferum		3.74	0.24				
51415110	**Good identification**						**51415110**
Eubacterium saburreum		100.0	0.45	SUC 99			
51421100	**Good identification**						**51421100**
					COC		
Eubacterium saburreum		97.27	0.40	SUC 99	1		
Blautia hansenii		2.73	-.03		99		
51421110	**Good identification**						**51421110**
Eubacterium saburreum		100.0	0.38	SUC 99			
51425100	**Good identification**						**51425100**
Eubacterium saburreum		100.0	0.35	SUC 99			
51431000	**Low discrimination**						**51431000**
Eubacterium saburreum		88.93	0.42	SUC 99			
Fusobacterium mortiferum		11.07	0.28				
51431010	**Good identification**						**51431010**
Eubacterium saburreum		100.0	0.40	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
51431100	**Low discrimination**					**51431100**
Eubacterium saburreum		88.93	0.48	SUC 99		
Fusobacterium mortiferum		11.07	0.34			
51431102	**Acceptable identification**					**51431102**
Eubacterium saburreum		92.73	0.35	SUC 99		
Fusobacterium mortiferum		7.27	0.18			
51431110	**Good identification**					**51431110**
Eubacterium saburreum		100.0	0.46	SUC 99		
51431120	**Good identification**					**51431120**
Eubacterium saburreum		100.0	0.35	SUC 99		
51435000	**Good identification**					**51435000**
Eubacterium saburreum		100.0	0.37	SUC 99		
51435010	**Good identification**					**51435010**
Eubacterium saburreum		100.0	0.36	SUC 99		
51435100	**Good identification**					**51435100**
Eubacterium saburreum		98.29	0.43	SUC 99		
Fusobacterium mortiferum		1.71	0.15			
51435110	**Good identification**					**51435110**
Eubacterium saburreum		100.0	0.42	SUC 99		
51500000	**Very good identification**					**51500000**
Clostridium bifermentans		97.21	0.64	RAF 1		
Clostridium sporogenes		1.52	0.32			
Clostridium botulinum biovar A		0.91	0.30			
51500001	**Acceptable identification**					**51500001**
Clostridium sordellii		91.61	0.44	ESL 5		
Clostridium bifermentans		7.94	0.30	RAF 1		
Clostridium novyi biovar A		0.33	0.19			
51500020	**Good identification**					**51500020**
Clostridium bifermentans		98.42	0.44	RAF 1		
Clostridium cadaveris		1.34	0.10	NAG 6		
Clostridium sporogenes		0.24	-.03			
51500040	**Acceptable identification**					**51500040**
Clostridium bifermentans		92.81	0.41	RAF 1		
Clostridium botulinum biovar A		6.23	0.21	SOR 4		
Clostridium novyi biovar A		0.61	0.16			
51501010	**Good identification**					**51501010**
					COC	
Eubacterium saburreum		98.69	0.45	SUC 99	1	
Blautia hansenii		1.32	-.03		99	
51501100	**Very good identification**					**51501100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Blautia hansenii		97.86	0.66	IND 1	99		
Eubacterium saburreum		2.14	0.53		1		
51501102	**Good identification**						**51501102**
					COC		
Blautia hansenii		96.65	0.50	IND 1	99		
Eubacterium saburreum		3.35	0.40	BGL 10	1		
51501110	**Low discrimination**						**51501110**
					COC		
Eubacterium saburreum		63.88	0.52	SUC 99	1		
Blautia hansenii		36.12	0.32		99		
51501112	**Low discrimination**						**51501112**
					COC		
Eubacterium saburreum		73.74	0.39	SUC 99	1		
Blautia hansenii		26.26	0.15		99		
51501120	**Good identification**						**51501120**
					COC	GRA	
Blautia hansenii		96.58	0.50	IND 1	99	99	
Eubacterium saburreum		3.35	0.40	NAG 10	1	85	
Prevotella melaninogenica		0.08	-.04		1	1	
51501130	**Low discrimination**						**51501130**
					COC		
Eubacterium saburreum		73.74	0.39	SUC 99	1		
Blautia hansenii		26.26	0.15		99		
51501500	**Good identification**						**51501500**
					COC		
Blautia hansenii		99.88	0.53	IND 1	99		
Eubacterium saburreum		0.13	0.19		1		
51501502	**Good identification**						**51501502**
Blautia hansenii		100.0	0.37	IND 1			
51501520	**Good identification**						**51501520**
					COC	GRA	
Blautia hansenii		99.56	0.37	IND 1	99	99	
Odoribacter splanchnicus		0.44	-.07	NAG 10	1	1	
51505000	**Low discrimination**						**51505000**
					COC		
Eubacterium saburreum		73.87	0.42	SUC 99	1		
Blautia hansenii		26.14	0.19		99		
51505010	**Good identification**						**51505010**
Eubacterium saburreum		100.0	0.41	SUC 99			
51505100	**Acceptable identification**						**51505100**
					COC		
Blautia hansenii		93.76	0.53	IND 1	99		
Eubacterium saburreum		6.25	0.49		1		
51505102	**Acceptable identification**						**51505102**
					COC		
Blautia hansenii		90.43	0.37	IND 1	99		
Eubacterium saburreum		9.57	0.36	BGL 10	1		
51505110	**Low discrimination**						**51505110**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		84.37	0.47	SUC 99	1		
Blautia hansenii		15.64	0.19		99		
51505120	**Acceptable identification**						**51505120**
					COC		
Blautia hansenii		90.43	0.37	IND 1	99		
Eubacterium saburreum		9.57	0.36	NAG 10	1		
51505500	**Good identification**						**51505500**
					COC		
Blautia hansenii		99.62	0.40	IND 1	99		
Eubacterium saburreum		0.38	0.14		1		
51510000	**Low discrimination**						**51510000**
					GRA	SPO	
Prevotella intermedia		72.53	0.53	ESL 1	1	1	
Clostridium bifermentans		24.91	0.43		99	99	
Clostridium sporogenes		1.31	0.19		99	99	
51510001	**Good identification**						**51510001**
					GRA	SPO	
Clostridium sordellii		98.25	0.44	ESL 5	99	99	
Prevotella intermedia		1.31	0.19	RAF 1	1	1	
Clostridium bifermentans		0.45	0.08		99	99	
51511000	**Low discrimination**						**51511000**
					SPO	GRA	
Prevotella intermedia		72.69	0.53	ESL 1	1	1	
Eubacterium saburreum		26.56	0.55		1	85	
Clostridium bifermentans		0.25	0.08		99	99	
51511002	**Low discrimination**						**51511002**
					GRA		
Eubacterium saburreum		85.37	0.42	SUC 99	85		
Prevotella intermedia		13.37	0.19		1		
Lactobacillus fermentum		1.26	0.14		99		
51511010	**Good identification**						**51511010**
Eubacterium saburreum		96.73	0.53	SUC 99			
Prevotella intermedia		3.27	0.19				
51511012	**Good identification**						**51511012**
Eubacterium saburreum		100.0	0.41	SUC 99			
51511020	**Low discrimination**						**51511020**
Eubacterium saburreum		86.46	0.42	SUC 99			
Prevotella intermedia		13.54	0.19				
51511030	**Good identification**						**51511030**
Eubacterium saburreum		100.0	0.41	SUC 99			
51511100	**Low discrimination**						**51511100**
					COC	GRA	
Eubacterium saburreum		85.00	0.61	SUC 99	1	85	
Blautia hansenii		13.11	0.32		99	99	
Prevotella intermedia		1.01	0.19		1	1	
51511102	**Low discrimination**						**51511102**
					COC		
Eubacterium saburreum		89.86	0.48	SUC 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		8.73	0.15		99		
Lactobacillus fermentum		0.93	0.18		1		
51511110	**Very good identification**						**51511110**
					COC		
Eubacterium saburreum		99.81	0.60	SUC 99	1		
Blautia hansenii		0.19	-.03		99		
51511112	**Good identification**						**51511112**
Eubacterium saburreum		100.0	0.47	SUC 99			
51511120	**Low discrimination**						**51511120**
					COC	GRA	
Eubacterium saburreum		85.89	0.48	SUC 99	1	85	
Blautia hansenii		8.34	0.15		99	99	
Prevotella melaninogenica		5.77	0.12		1	1	
51511122	**Low discrimination**						**51511122**
					CAT		
Eubacterium saburreum		71.10	0.35	SUC 99	1		
Capnocytophaga ochracea		16.26	0.12		90		
Prevotella melaninogenica		5.16	0.00		1		
51511130	**Good identification**						**51511130**
Eubacterium saburreum		100.0	0.47	SUC 99			
51515000	**Good identification**						**51515000**
Eubacterium saburreum		95.12	0.50	SUC 99			
Prevotella intermedia		4.88	0.19				
51515002	**Good identification**						**51515002**
Eubacterium saburreum		100.0	0.37	SUC 99			
51515010	**Good identification**						**51515010**
Eubacterium saburreum		100.0	0.49	SUC 99			
51515012	**Good identification**						**51515012**
Eubacterium saburreum		100.0	0.36	SUC 99			
51515020	**Good identification**						**51515020**
Eubacterium saburreum		100.0	0.37	SUC 99			
51515030	**Good identification**						**51515030**
Eubacterium saburreum		100.0	0.36	SUC 99			
51515100	**Very good identification**						**51515100**
					COC		
Eubacterium saburreum		95.01	0.57	SUC 99	1		
Blautia hansenii		4.80	0.19		99		
Lactobacillus fermentum		0.19	0.14		1		
51515102	**Good identification**						**51515102**
					COC		
Eubacterium saburreum		95.87	0.44	SUC 99	1		
Blautia hansenii		3.05	0.03		99		
Lactobacillus fermentum		1.08	0.14		1		
51515110	**Good identification**						**51515110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.55	SUC 99		
51515112	**Good identification**					**51515112**
Eubacterium saburreum		100.0	0.42	SUC 99		
51515120	**Good identification**					**51515120**
					COC	
Eubacterium saburreum		96.92	0.44	SUC 99	1	
Blautia hansenii		3.08	0.03		99	
51515130	**Good identification**					**51515130**
Eubacterium saburreum		100.0	0.42	SUC 99		
51521000	**Good identification**					**51521000**
					COC	
Eubacterium saburreum		98.39	0.44	SUC 99	1	
Blautia hansenii		1.61	-.03		99	
51521010	**Good identification**					**51521010**
Eubacterium saburreum		100.0	0.42	SUC 99		
51521100	**Low discrimination**					**51521100**
					COC	
Eubacterium saburreum		59.03	0.50	SUC 99	1	
Blautia hansenii		40.97	0.32		99	
51521102	**Low discrimination**					**51521102**
					COC	
Eubacterium saburreum		69.59	0.37	SUC 99	1	
Blautia hansenii		30.41	0.15		99	
51521110	**Good identification**					**51521110**
					COC	
Eubacterium saburreum		99.15	0.49	SUC 99	1	
Blautia hansenii		0.85	-.03		99	
51521112	**Good identification**					**51521112**
Eubacterium saburreum		100.0	0.36	SUC 99		
51521120	**Low discrimination**					**51521120**
					COC	
Eubacterium saburreum		69.59	0.37	SUC 99	1	
Blautia hansenii		30.41	0.15		99	
51521130	**Good identification**					**51521130**
Eubacterium saburreum		100.0	0.36	SUC 99		
51525000	**Good identification**					**51525000**
Eubacterium saburreum		100.0	0.39	SUC 99		
51525010	**Good identification**					**51525010**
Eubacterium saburreum		100.0	0.38	SUC 99		
51525100	**Low discrimination**					**51525100**
					COC	
Eubacterium saburreum		81.47	0.46	SUC 99	1	
Blautia hansenii		18.53	0.19		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
51525110	**Good identification**						**51525110**
Eubacterium saburreum		100.0	0.44	SUC 99			
51531000	**Good identification**						**51531000**
Eubacterium saburreum		95.67	0.52	SUC 99			
Prevotella intermedia		3.97	0.19				
Fusobacterium mortiferum		0.37	0.12				
51531002	**Good identification**						**51531002**
Eubacterium saburreum		100.0	0.39	SUC 99			
51531010	**Good identification**						**51531010**
Eubacterium saburreum		100.0	0.50	SUC 99			
51531012	**Good identification**						**51531012**
Eubacterium saburreum		100.0	0.38	SUC 99			
51531020	**Good identification**						**51531020**
Eubacterium saburreum		100.0	0.39	SUC 99			
51531030	**Good identification**						**51531030**
Eubacterium saburreum		100.0	0.38	SUC 99			
51531100	**Very good identification**						**51531100**
					COC	GRA	
Eubacterium saburreum		99.39	0.58	SUC 99	1	85	
Fusobacterium mortiferum		0.38	0.19		1	1	
Blautia hansenii		0.23	-.03		99	99	
51531102	**Good identification**						**51531102**
Eubacterium saburreum		100.0	0.45	SUC 99			
51531110	**Very good identification**						**51531110**
Eubacterium saburreum		100.0	0.57	SUC 99			
51531112	**Good identification**						**51531112**
Eubacterium saburreum		100.0	0.44	SUC 99			
51531120	**Good identification**						**51531120**
Eubacterium saburreum		100.0	0.45	SUC 99			
51531130	**Good identification**						**51531130**
Eubacterium saburreum		100.0	0.44	SUC 99			
51535000	**Good identification**						**51535000**
Eubacterium saburreum		100.0	0.47	SUC 99			
51535010	**Good identification**						**51535010**
Eubacterium saburreum		100.0	0.46	SUC 99			
51535100	**Good identification**						**51535100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.54	SUC 99			
51535102	**Good identification**						**51535102**
Eubacterium saburreum		100.0	0.41	SUC 99			
51535110	**Good identification**						**51535110**
Eubacterium saburreum		100.0	0.52	SUC 99			
51535112	**Good identification**						**51535112**
Eubacterium saburreum		100.0	0.39	SUC 99			
51535120	**Good identification**						**51535120**
Eubacterium saburreum		100.0	0.41	SUC 99			
51535130	**Good identification**						**51535130**
Eubacterium saburreum		100.0	0.39	SUC 99			
51575722	**Acceptable identification**						**51575722**
Bacteroides thetaiotaomicron		92.96	0.37	SUC 98			
Bacteroides ovatus		6.22	0.17	MNS 99			
Parabacteroides distasonis		0.82	0.18				
51600000	**Good identification**						**51600000**
Clostridium bifermentans		100.0	0.42	SAL 10			
51610000	**Low discrimination**						**51610000**
					GRA	SPO	
Fusobacterium mortiferum		66.76	0.39	IND 1	1	1	
Clostridium bifermentans		33.24	0.20		99	99	
51610100	**Good identification**						**51610100**
Fusobacterium mortiferum		100.0	0.45	IND 1			
51611000	**Good identification**						**51611000**
Fusobacterium mortiferum		98.16	0.42	IND 1			
Eubacterium saburreum		1.84	0.11				
51611100	**Good identification**						**51611100**
Fusobacterium mortiferum		98.16	0.48	IND 1			
Eubacterium saburreum		1.84	0.17				
51630100	**Good identification**						**51630100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
51631100	**Good identification**						**51631100**
Fusobacterium mortiferum		95.99	0.39	IND 1			
Eubacterium saburreum		4.01	0.14				
51650100	**Good identification**						**51650100**
Fusobacterium mortiferum		100.0	0.37	IND 1			
51651100	**Good identification**						**51651100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.40	IND 1		
51700000	**Good identification**					**51700000**
Clostridium bifermentans		99.86	0.48	SAL 10		
Clostridium sporogenes		0.14	-.03	RAF 1		
52001010	**Good identification**					**52001010**
Eubacterium saburreum		95.98	0.40	GLU 99		
Fusobacterium necrophorum subsp. necrophorum		4.02	-.03			
52001100	**Good identification**					**52001100**
Eubacterium saburreum		98.55	0.47	GLU 99		
Fusobacterium necrophorum subsp. necrophorum		1.45	-.03			
52001110	**Good identification**					**52001110**
Eubacterium saburreum		100.0	0.46	GLU 99		
52005000	**Acceptable identification**					**52005000**
Eubacterium saburreum		94.02	0.37	GLU 99		
Fusobacterium necrophorum subsp. necrophorum		5.98	-.03			
52005010	**Good identification**					**52005010**
Eubacterium saburreum		100.0	0.35	GLU 99		
52005100	**Good identification**					**52005100**
Eubacterium saburreum		100.0	0.43	GLU 99		
52005110	**Good identification**					**52005110**
Eubacterium saburreum		100.0	0.41	GLU 99		
52011000	**Acceptable identification**					**52011000**
					GRA	
Eubacterium saburreum		93.06	0.49	GLU 99	85	
Atopobium minutum		5.88	0.25		99	
Fusobacterium necrophorum subsp. necrophorum		1.06	-.03		1	
52011002	**Low discrimination**					**52011002**
Eubacterium saburreum		80.73	0.36	GLU 99		
Atopobium minutum		19.27	0.22			
52011010	**Good identification**					**52011010**
Eubacterium saburreum		100.0	0.48	GLU 99		
52011020	**Low discrimination**					**52011020**
Eubacterium saburreum		89.34	0.36	GLU 99		
Atopobium minutum		10.66	0.17			
52011100	**Good identification**					**52011100**
					GRA	
Eubacterium saburreum		99.18	0.56	GLU 99	85	
Fusobacterium mortiferum		0.44	0.17		1	
Actinomyces meyeri		0.38	0.07		99	
52011102	**Good identification**					**52011102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.43	GLU 99		
52011110	**Good identification**					**52011110**
Eubacterium saburreum		100.0	0.54	GLU 99		
52011112	**Good identification**					**52011112**
Eubacterium saburreum		100.0	0.41	GLU 99		
52011120	**Good identification**					**52011120**
Eubacterium saburreum		100.0	0.43	GLU 99		
52011130	**Good identification**					**52011130**
Eubacterium saburreum		100.0	0.41	GLU 99		
52015000	**Good identification**					**52015000**
Eubacterium saburreum		100.0	0.45	GLU 99		
52015010	**Good identification**					**52015010**
Eubacterium saburreum		100.0	0.43	GLU 99		
52015100	**Good identification**					**52015100**
Eubacterium saburreum		98.63	0.51	GLU 99		
Actinomyces meyeri		1.38	0.12			
52015102	**Good identification**					**52015102**
Eubacterium saburreum		100.0	0.38	GLU 99		
52015110	**Good identification**					**52015110**
Eubacterium saburreum		100.0	0.49	GLU 99		
52015112	**Good identification**					**52015112**
Eubacterium saburreum		100.0	0.37	GLU 99		
52015120	**Good identification**					**52015120**
Eubacterium saburreum		100.0	0.38	GLU 99		
52015130	**Good identification**					**52015130**
Eubacterium saburreum		100.0	0.37	GLU 99		
52021000	**Good identification**					**52021000**
Eubacterium saburreum		95.11	0.38	GLU 99		
Fusobacterium necrophorum subsp. necrophorum		4.89	-.03			
52021010	**Good identification**					**52021010**
Eubacterium saburreum		100.0	0.37	GLU 99		
52021100	**Good identification**					**52021100**
Eubacterium saburreum		100.0	0.44	GLU 99		
52021110	**Good identification**					**52021110**
Eubacterium saburreum		100.0	0.43	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
52025100	**Good identification**					**52025100**
Eubacterium saburreum		100.0	0.40	GLU 99		
52025110	**Good identification**					**52025110**
Eubacterium saburreum		100.0	0.38	GLU 99		
52031000	**Good identification**					**52031000**
Eubacterium saburreum		100.0	0.46	GLU 99		
52031010	**Good identification**					**52031010**
Eubacterium saburreum		100.0	0.45	GLU 99		
52031100	**Good identification**					**52031100**
Eubacterium saburreum		100.0	0.53	GLU 99		
52031102	**Good identification**					**52031102**
Eubacterium saburreum		100.0	0.40	GLU 99		
52031110	**Good identification**					**52031110**
Eubacterium saburreum		100.0	0.51	GLU 99		
52031112	**Good identification**					**52031112**
Eubacterium saburreum		100.0	0.38	GLU 99		
52031120	**Good identification**					**52031120**
Eubacterium saburreum		100.0	0.40	GLU 99		
52031130	**Good identification**					**52031130**
Eubacterium saburreum		100.0	0.38	GLU 99		
52035000	**Good identification**					**52035000**
Eubacterium saburreum		100.0	0.42	GLU 99		
52035010	**Good identification**					**52035010**
Eubacterium saburreum		100.0	0.40	GLU 99		
52035100	**Good identification**					**52035100**
Eubacterium saburreum		100.0	0.48	GLU 99		
52035102	**Good identification**					**52035102**
Eubacterium saburreum		100.0	0.35	GLU 99		
52035110	**Good identification**					**52035110**
Eubacterium saburreum		100.0	0.46	GLU 99		
52035120	**Good identification**					**52035120**
Eubacterium saburreum		100.0	0.35	GLU 99		
52100000	**Low discrimination**					**52100000**
Clostridium bifermentans		86.50	0.46	GLU 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		7.71	0.29	SUC 1			
Clostridium sporogenes		2.74	0.19				
52101000	**Good identification**						**52101000**
					SPO	GRA	
Eubacterium saburreum		96.72	0.51	GLU 99	1	85	
Clostridium bifermentans		2.45	0.12		99	99	
Fusobacterium necrophorum subsp. necrophorum		0.83	-.03		1	1	
52101002	**Good identification**						**52101002**
Eubacterium saburreum		100.0	0.39	GLU 99			
52101010	**Good identification**						**52101010**
Eubacterium saburreum		100.0	0.50	GLU 99			
52101012	**Good identification**						**52101012**
Eubacterium saburreum		100.0	0.37	GLU 99			
52101020	**Good identification**						**52101020**
Eubacterium saburreum		100.0	0.39	GLU 99			
52101030	**Good identification**						**52101030**
Eubacterium saburreum		100.0	0.37	GLU 99			
52101100	**Very good identification**						**52101100**
Eubacterium saburreum		100.0	0.58	GLU 99			
52101102	**Good identification**						**52101102**
Eubacterium saburreum		100.0	0.45	GLU 99			
52101110	**Very good identification**						**52101110**
Eubacterium saburreum		100.0	0.56	GLU 99			
52101112	**Good identification**						**52101112**
Eubacterium saburreum		100.0	0.43	GLU 99			
52101120	**Good identification**						**52101120**
Eubacterium saburreum		98.55	0.45	GLU 99			
Prevotella melaninogenica		1.45	-.03				
52101130	**Good identification**						**52101130**
Eubacterium saburreum		100.0	0.43	GLU 99			
52105000	**Good identification**						**52105000**
Eubacterium saburreum		100.0	0.47	GLU 99			
52105010	**Good identification**						**52105010**
Eubacterium saburreum		100.0	0.45	GLU 99			
52105100	**Good identification**						**52105100**
Eubacterium saburreum		100.0	0.53	GLU 99			
52105102	**Good identification**						**52105102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.40	GLU 99		
52105110	**Good identification**					**52105110**
Eubacterium saburreum		100.0	0.52	GLU 99		
52105112	**Good identification**					**52105112**
Eubacterium saburreum		100.0	0.39	GLU 99		
52105120	**Good identification**					**52105120**
Eubacterium saburreum		100.0	0.40	GLU 99		
52105130	**Good identification**					**52105130**
Eubacterium saburreum		100.0	0.39	GLU 99		
52111000	**Good identification**					**52111000**
Eubacterium saburreum		94.90	0.60	GLU 99		
Prevotella intermedia		5.10	0.28			
52111002	**Good identification**					**52111002**
Eubacterium saburreum		100.0	0.47	GLU 99		
52111010	**Very good identification**					**52111010**
Eubacterium saburreum		100.0	0.58	GLU 99		
52111012	**Good identification**					**52111012**
Eubacterium saburreum		100.0	0.45	GLU 99		
52111020	**Good identification**					**52111020**
Eubacterium saburreum		100.0	0.47	GLU 99		
52111030	**Good identification**					**52111030**
Eubacterium saburreum		100.0	0.45	GLU 99		
52111100	**Very good identification**					**52111100**
Eubacterium saburreum		99.89	0.66	GLU 99		
Actinomyces meyeri		0.11	0.08			
52111102	**Good identification**					**52111102**
Eubacterium saburreum		100.0	0.53	GLU 99		
52111110	**Very good identification**					**52111110**
Eubacterium saburreum		100.0	0.64	GLU 99		
52111112	**Good identification**					**52111112**
Eubacterium saburreum		100.0	0.52	GLU 99		
52111120	**Good identification**					**52111120**
Eubacterium saburreum		95.77	0.53	GLU 99		
Prevotella melaninogenica		4.23	0.14			
52111122	**Good identification**					**52111122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		95.45	0.40	GLU 99		
Prevotella melaninogenica		4.56	0.02			
52111130	**Good identification**					**52111130**
Eubacterium saburreum		100.0	0.52	GLU 99		
52111132	**Good identification**					**52111132**
Eubacterium saburreum		100.0	0.39	GLU 99		
52115000	**Good identification**					**52115000**
Eubacterium saburreum		99.77	0.55	GLU 99		
Actinomyces meyeri		0.23	0.02			
52115002	**Good identification**					**52115002**
Eubacterium saburreum		100.0	0.42	GLU 99		
52115010	**Good identification**					**52115010**
Eubacterium saburreum		100.0	0.53	GLU 99		
52115012	**Good identification**					**52115012**
Eubacterium saburreum		100.0	0.41	GLU 99		
52115020	**Good identification**					**52115020**
Eubacterium saburreum		100.0	0.42	GLU 99		
52115030	**Good identification**					**52115030**
Eubacterium saburreum		100.0	0.41	GLU 99		
52115100	**Very good identification**					**52115100**
Eubacterium saburreum		99.61	0.61	GLU 99		
Actinomyces meyeri		0.39	0.13			
52115102	**Good identification**					**52115102**
Eubacterium saburreum		100.0	0.48	GLU 99		
52115110	**Very good identification**					**52115110**
Eubacterium saburreum		100.0	0.60	GLU 99		
52115112	**Good identification**					**52115112**
Eubacterium saburreum		100.0	0.47	GLU 99		
52115120	**Good identification**					**52115120**
Eubacterium saburreum		100.0	0.48	GLU 99		
52115122	**Good identification**					**52115122**
Eubacterium saburreum		100.0	0.35	GLU 99		
52115130	**Good identification**					**52115130**
Eubacterium saburreum		100.0	0.47	GLU 99		
52121000	**Good identification**					**52121000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.48	GLU 99		
52121002	**Good identification**					**52121002**
Eubacterium saburreum		100.0	0.36	GLU 99		
52121010	**Good identification**					**52121010**
Eubacterium saburreum		100.0	0.47	GLU 99		
52121020	**Good identification**					**52121020**
Eubacterium saburreum		100.0	0.36	GLU 99		
52121100	**Good identification**					**52121100**
Eubacterium saburreum		100.0	0.55	GLU 99		
52121102	**Good identification**					**52121102**
Eubacterium saburreum		100.0	0.42	GLU 99		
52121110	**Good identification**					**52121110**
Eubacterium saburreum		100.0	0.53	GLU 99		
52121112	**Good identification**					**52121112**
Eubacterium saburreum		100.0	0.40	GLU 99		
52121120	**Good identification**					**52121120**
Eubacterium saburreum		100.0	0.42	GLU 99		
52121130	**Good identification**					**52121130**
Eubacterium saburreum		100.0	0.40	GLU 99		
52125000	**Good identification**					**52125000**
Eubacterium saburreum		100.0	0.44	GLU 99		
52125010	**Good identification**					**52125010**
Eubacterium saburreum		100.0	0.42	GLU 99		
52125100	**Good identification**					**52125100**
Eubacterium saburreum		100.0	0.50	GLU 99		
52125102	**Good identification**					**52125102**
Eubacterium saburreum		100.0	0.37	GLU 99		
52125110	**Good identification**					**52125110**
Eubacterium saburreum		100.0	0.49	GLU 99		
52125112	**Good identification**					**52125112**
Eubacterium saburreum		100.0	0.36	GLU 99		
52125120	**Good identification**					**52125120**
Eubacterium saburreum		100.0	0.37	GLU 99		
52125130	**Good identification**					**52125130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.36	GLU 99		
52131000	**Very good identification**					**52131000**
Eubacterium saburreum		100.0	0.57	GLU 99		
52131002	**Good identification**					**52131002**
Eubacterium saburreum		100.0	0.44	GLU 99		
52131010	**Good identification**					**52131010**
Eubacterium saburreum		100.0	0.55	GLU 99		
52131012	**Good identification**					**52131012**
Eubacterium saburreum		100.0	0.42	GLU 99		
52131020	**Good identification**					**52131020**
Eubacterium saburreum		100.0	0.44	GLU 99		
52131030	**Good identification**					**52131030**
Eubacterium saburreum		100.0	0.42	GLU 99		
52131100	**Very good identification**					**52131100**
Eubacterium saburreum		100.0	0.63	GLU 99		
52131102	**Good identification**					**52131102**
Eubacterium saburreum		100.0	0.50	GLU 99		
52131110	**Very good identification**					**52131110**
Eubacterium saburreum		100.0	0.61	GLU 99		
52131112	**Good identification**					**52131112**
Eubacterium saburreum		100.0	0.48	GLU 99		
52131120	**Good identification**					**52131120**
Eubacterium saburreum		100.0	0.50	GLU 99		
52131122	**Good identification**					**52131122**
Eubacterium saburreum		100.0	0.37	GLU 99		
52131130	**Good identification**					**52131130**
Eubacterium saburreum		100.0	0.48	GLU 99		
52131132	**Good identification**					**52131132**
Eubacterium saburreum		100.0	0.36	GLU 99		
52135000	**Good identification**					**52135000**
Eubacterium saburreum		100.0	0.52	GLU 99		
52135002	**Good identification**					**52135002**
Eubacterium saburreum		100.0	0.39	GLU 99		
52135010	**Good identification**					**52135010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.50	GLU 99		
52135012	**Good identification**					**52135012**
Eubacterium saburreum		100.0	0.38	GLU 99		
52135020	**Good identification**					**52135020**
Eubacterium saburreum		100.0	0.39	GLU 99		
52135030	**Good identification**					**52135030**
Eubacterium saburreum		100.0	0.38	GLU 99		
52135100	**Very good identification**					**52135100**
Eubacterium saburreum		100.0	0.58	GLU 99		
52135102	**Good identification**					**52135102**
Eubacterium saburreum		100.0	0.45	GLU 99		
52135110	**Very good identification**					**52135110**
Eubacterium saburreum		100.0	0.57	GLU 99		
52135112	**Good identification**					**52135112**
Eubacterium saburreum		100.0	0.44	GLU 99		
52135120	**Good identification**					**52135120**
Eubacterium saburreum		100.0	0.45	GLU 99		
52135130	**Good identification**					**52135130**
Eubacterium saburreum		100.0	0.44	GLU 99		
52401000	**Acceptable identification**					**52401000**
Eubacterium saburreum		94.02	0.37	GLU 99		
Fusobacterium necrophorum subsp. necrophorum		5.98	-.03			
52401010	**Good identification**					**52401010**
Eubacterium saburreum		100.0	0.35	GLU 99		
52401100	**Good identification**					**52401100**
Eubacterium saburreum		100.0	0.43	GLU 99		
52401110	**Good identification**					**52401110**
Eubacterium saburreum		100.0	0.41	GLU 99		
52405100	**Good identification**					**52405100**
Eubacterium saburreum		100.0	0.38	GLU 99		
52405110	**Good identification**					**52405110**
Eubacterium saburreum		100.0	0.37	GLU 99		
52411000	**Good identification**					**52411000**
Eubacterium saburreum		98.44	0.45	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		1.57	0.16			
52411010	**Good identification**					**52411010**
Eubacterium saburreum		100.0	0.43	GLU 99		
52411100	**Good identification**					**52411100**
Eubacterium saburreum		98.44	0.51	GLU 99		
Fusobacterium mortiferum		1.57	0.22			
52411102	**Good identification**					**52411102**
Eubacterium saburreum		100.0	0.38	GLU 99		
52411110	**Good identification**					**52411110**
Eubacterium saburreum		100.0	0.49	GLU 99		
52411112	**Good identification**					**52411112**
Eubacterium saburreum		100.0	0.37	GLU 99		
52411120	**Good identification**					**52411120**
Eubacterium saburreum		97.10	0.38	GLU 99		
Prevotella melaninogenica		2.90	-.04			
52411130	**Good identification**					**52411130**
Eubacterium saburreum		100.0	0.37	GLU 99		
52415000	**Good identification**					**52415000**
Eubacterium saburreum		100.0	0.40	GLU 99		
52415010	**Good identification**					**52415010**
Eubacterium saburreum		100.0	0.39	GLU 99		
52415100	**Good identification**					**52415100**
Eubacterium saburreum		100.0	0.46	GLU 99		
52415110	**Good identification**					**52415110**
Eubacterium saburreum		100.0	0.45	GLU 99		
52421100	**Good identification**					**52421100**
Eubacterium saburreum		100.0	0.40	GLU 99		
52421110	**Good identification**					**52421110**
Eubacterium saburreum		100.0	0.38	GLU 99		
52425100	**Good identification**					**52425100**
Eubacterium saburreum		100.0	0.35	GLU 99		
52431000	**Good identification**					**52431000**
Eubacterium saburreum		100.0	0.42	GLU 99		
52431010	**Good identification**					**52431010**
Eubacterium saburreum		100.0	0.40	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
52431100	**Good identification**					**52431100**
Eubacterium saburreum		99.29	0.48	GLU 99		
Fusobacterium mortiferum		0.71	0.13			
52431102	**Good identification**					**52431102**
Eubacterium saburreum		100.0	0.35	GLU 99		
52431110	**Good identification**					**52431110**
Eubacterium saburreum		100.0	0.46	GLU 99		
52431120	**Good identification**					**52431120**
Eubacterium saburreum		100.0	0.35	GLU 99		
52435000	**Good identification**					**52435000**
Eubacterium saburreum		100.0	0.37	GLU 99		
52435010	**Good identification**					**52435010**
Eubacterium saburreum		100.0	0.36	GLU 99		
52435100	**Good identification**					**52435100**
Eubacterium saburreum		100.0	0.43	GLU 99		
52435110	**Good identification**					**52435110**
Eubacterium saburreum		100.0	0.42	GLU 99		
52501000	**Good identification**					**52501000**
Eubacterium saburreum		100.0	0.47	GLU 99		
52501010	**Good identification**					**52501010**
Eubacterium saburreum		100.0	0.45	GLU 99		
52501100	**Good identification**					**52501100**
					COC	
Eubacterium saburreum		99.54	0.53	GLU 99	1	
Blautia hansenii		0.47	-.03		99	
52501102	**Good identification**					**52501102**
Eubacterium saburreum		100.0	0.40	GLU 99		
52501110	**Good identification**					**52501110**
Eubacterium saburreum		100.0	0.52	GLU 99		
52501112	**Good identification**					**52501112**
Eubacterium saburreum		100.0	0.39	GLU 99		
52501120	**Low discrimination**					**52501120**
Eubacterium saburreum		80.24	0.40	GLU 99		
Prevotella melaninogenica		19.76	0.14			
52501130	**Good identification**					**52501130**
Eubacterium saburreum		100.0	0.39	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
52505000	**Good identification**					**52505000**
Eubacterium saburreum		100.0	0.42	GLU 99		
52505010	**Good identification**					**52505010**
Eubacterium saburreum		100.0	0.41	GLU 99		
52505100	**Good identification**					**52505100**
Eubacterium saburreum		100.0	0.49	GLU 99		
52505102	**Good identification**					**52505102**
Eubacterium saburreum		100.0	0.36	GLU 99		
52505110	**Good identification**					**52505110**
Eubacterium saburreum		100.0	0.47	GLU 99		
52505120	**Good identification**					**52505120**
Eubacterium saburreum		100.0	0.36	GLU 99		
52511000	**Low discrimination**					**52511000**
Eubacterium saburreum		86.98	0.55	GLU 99		
Prevotella intermedia		13.03	0.31			
52511002	**Good identification**					**52511002**
Eubacterium saburreum		100.0	0.42	GLU 99		
52511010	**Good identification**					**52511010**
Eubacterium saburreum		100.0	0.53	GLU 99		
52511012	**Good identification**					**52511012**
Eubacterium saburreum		100.0	0.41	GLU 99		
52511020	**Good identification**					**52511020**
Eubacterium saburreum		98.29	0.42	GLU 99		
Prevotella melaninogenica		1.71	-.04			
52511030	**Good identification**					**52511030**
Eubacterium saburreum		100.0	0.41	GLU 99		
52511100	**Very good identification**					**52511100**
Eubacterium saburreum		99.87	0.61	GLU 99		
Prevotella melaninogenica		0.13	-.04			
52511102	**Good identification**					**52511102**
Eubacterium saburreum		100.0	0.48	GLU 99		
52511110	**Very good identification**					**52511110**
Eubacterium saburreum		100.0	0.60	GLU 99		
52511112	**Good identification**					**52511112**
Eubacterium saburreum		100.0	0.47	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
52511120	**Low discrimination**					**52511120**
Eubacterium saburreum		57.51	0.48	GLU 99		
Prevotella melaninogenica		42.50	0.30			
52511122	**Low discrimination**					**52511122**
					CAT	
Eubacterium saburreum		49.35	0.35	GLU 99	1	
Prevotella melaninogenica		39.36	0.18		1	
Capnocytophaga ochracea		11.29	0.12		90	
52511130	**Good identification**					**52511130**
Eubacterium saburreum		99.10	0.47	GLU 99		
Prevotella melaninogenica		0.90	-.04			
52515000	**Good identification**					**52515000**
Eubacterium saburreum		100.0	0.50	GLU 99		
52515002	**Good identification**					**52515002**
Eubacterium saburreum		100.0	0.37	GLU 99		
52515010	**Good identification**					**52515010**
Eubacterium saburreum		100.0	0.49	GLU 99		
52515012	**Good identification**					**52515012**
Eubacterium saburreum		100.0	0.36	GLU 99		
52515020	**Good identification**					**52515020**
Eubacterium saburreum		100.0	0.37	GLU 99		
52515030	**Good identification**					**52515030**
Eubacterium saburreum		100.0	0.36	GLU 99		
52515100	**Very good identification**					**52515100**
Eubacterium saburreum		100.0	0.57	GLU 99		
52515102	**Good identification**					**52515102**
Eubacterium saburreum		100.0	0.44	GLU 99		
52515110	**Good identification**					**52515110**
Eubacterium saburreum		100.0	0.55	GLU 99		
52515112	**Good identification**					**52515112**
Eubacterium saburreum		100.0	0.42	GLU 99		
52515120	**Good identification**					**52515120**
Eubacterium saburreum		98.63	0.44	GLU 99		
Prevotella melaninogenica		1.37	-.04			
52515130	**Good identification**					**52515130**
Eubacterium saburreum		100.0	0.42	GLU 99		
52521000	**Good identification**					**52521000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.44	GLU 99		
52521010	**Good identification**					**52521010**
Eubacterium saburreum		100.0	0.42	GLU 99		
52521100	**Good identification**					**52521100**
Eubacterium saburreum		100.0	0.50	GLU 99		
52521102	**Good identification**					**52521102**
Eubacterium saburreum		100.0	0.37	GLU 99		
52521110	**Good identification**					**52521110**
Eubacterium saburreum		100.0	0.49	GLU 99		
52521112	**Good identification**					**52521112**
Eubacterium saburreum		100.0	0.36	GLU 99		
52521120	**Good identification**					**52521120**
Eubacterium saburreum		100.0	0.37	GLU 99		
52521130	**Good identification**					**52521130**
Eubacterium saburreum		100.0	0.36	GLU 99		
52525000	**Good identification**					**52525000**
Eubacterium saburreum		100.0	0.39	GLU 99		
52525010	**Good identification**					**52525010**
Eubacterium saburreum		100.0	0.38	GLU 99		
52525100	**Good identification**					**52525100**
Eubacterium saburreum		100.0	0.46	GLU 99		
52525110	**Good identification**					**52525110**
Eubacterium saburreum		100.0	0.44	GLU 99		
52531000	**Good identification**					**52531000**
Eubacterium saburreum		100.0	0.52	GLU 99		
52531002	**Good identification**					**52531002**
Eubacterium saburreum		100.0	0.39	GLU 99		
52531010	**Good identification**					**52531010**
Eubacterium saburreum		100.0	0.50	GLU 99		
52531012	**Good identification**					**52531012**
Eubacterium saburreum		100.0	0.38	GLU 99		
52531020	**Good identification**					**52531020**
Eubacterium saburreum		100.0	0.39	GLU 99		
52531030	**Good identification**					**52531030**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	GLU 99		
52531100	**Very good identification**					**52531100**
Eubacterium saburreum		100.0	0.58	GLU 99		
52531102	**Good identification**					**52531102**
Eubacterium saburreum		100.0	0.45	GLU 99		
52531110	**Very good identification**					**52531110**
Eubacterium saburreum		100.0	0.57	GLU 99		
52531112	**Good identification**					**52531112**
Eubacterium saburreum		100.0	0.44	GLU 99		
52531120	**Good identification**					**52531120**
Eubacterium saburreum		98.89	0.45	GLU 99		
Prevotella melaninogenica		1.11	-.04			
52531130	**Good identification**					**52531130**
Eubacterium saburreum		100.0	0.44	GLU 99		
52535000	**Good identification**					**52535000**
Eubacterium saburreum		100.0	0.47	GLU 99		
52535010	**Good identification**					**52535010**
Eubacterium saburreum		100.0	0.46	GLU 99		
52535100	**Good identification**					**52535100**
Eubacterium saburreum		100.0	0.54	GLU 99		
52535102	**Good identification**					**52535102**
Eubacterium saburreum		100.0	0.41	GLU 99		
52535110	**Good identification**					**52535110**
Eubacterium saburreum		100.0	0.52	GLU 99		
52535112	**Good identification**					**52535112**
Eubacterium saburreum		100.0	0.39	GLU 99		
52535120	**Good identification**					**52535120**
Eubacterium saburreum		100.0	0.41	GLU 99		
52535130	**Good identification**					**52535130**
Eubacterium saburreum		100.0	0.39	GLU 99		
52575722	**Good identification**					**52575722**
Bacteroides thetaiotaomicron		95.77	0.40	GLU 97		
Bacteroides ovatus		4.23	0.17	MNS 99		
53000000	**Low discrimination**					**53000000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		78.11	0.58	SUC 1	99	
Clostridium botulinum biovar A		15.53	0.47		99	
Atopobium minutum		1.66	0.38		1	
53000010	**Low discrimination**					**53000010**
					SPO	
Eubacterium saburreum		55.87	0.40	GAL 99	1	
Clostridium bifermentans		34.10	0.24		99	
Clostridium botulinum biovar A		6.78	0.13		99	
53000020	**Low discrimination**					**53000020**
					SPO	
Clostridium cadaveris		78.93	0.47	ESL 1	99	
Clostridium bifermentans		17.59	0.38	SUC 7	99	
Atopobium minutum		1.95	0.30		1	
53000040	**Low discrimination**					**53000040**
Clostridium botulinum biovar A		58.21	0.38	IND 1		
Clostridium bifermentans		40.82	0.34	SUC 7		
Clostridium novyi biovar A		0.63	0.16			
53000100	**Low discrimination**					**53000100**
					SPO	
Eubacterium saburreum		79.02	0.47	GAL 99	1	
Clostridium bifermentans		16.91	0.24		99	
Clostridium botulinum biovar A		3.36	0.13		99	
53000110	**Good identification**					**53000110**
Eubacterium saburreum		100.0	0.46	GAL 99		
53001000	**Very good identification**					**53001000**
					SPO	
Eubacterium saburreum		96.55	0.75		1	
Eubacterium tortuosum		2.08	0.32		1	
Clostridium bifermentans		0.49	0.24		99	
53001001	**Low discrimination**					**53001001**
					SPO	
Eubacterium saburreum		77.82	0.41	URE 1	1	
Clostridium sordellii		20.51	0.15		99	
Eubacterium tortuosum		1.68	-.03		1	
53001002	**Very good identification**					**53001002**
Eubacterium saburreum		96.63	0.62			
Eubacterium tortuosum		2.08	0.19			
Atopobium minutum		1.29	0.27			
53001010	**Very good identification**					**53001010**
Eubacterium saburreum		99.97	0.74			
Eubacterium tortuosum		0.03	-.03			
53001011	**Good identification**					**53001011**
Eubacterium saburreum		100.0	0.40	URE 1		
53001012	**Very good identification**					**53001012**
Eubacterium saburreum		100.0	0.61			
53001020	**Very good identification**					**53001020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Eubacterium saburreum		96.32	0.62		1	
Eubacterium tortuosum		2.07	0.19		1	
Clostridium cadaveris		0.79	0.13		99	
53001022	**Good identification**					**53001022**
Eubacterium saburreum		95.54	0.49			
Atopobium minutum		2.41	0.18			
Eubacterium tortuosum		2.06	0.06			
53001030	**Very good identification**					**53001030**
Eubacterium saburreum		100.0	0.61			
53001032	**Good identification**					**53001032**
Eubacterium saburreum		100.0	0.48			
53001040	**Acceptable identification**					**53001040**
					SPO	
Eubacterium saburreum		93.28	0.41	SOR 1	1	
Clostridium botulinum biovar A		2.77	0.04		99	
Eubacterium tortuosum		2.01	-.03		1	
53001050	**Good identification**					**53001050**
Eubacterium saburreum		100.0	0.40	SOR 1		
53001100	**Very good identification**					**53001100**
					GRA	
Eubacterium saburreum		99.97	0.82		85	
Fusobacterium mortiferum		0.01	0.17		1	
Actinomyces meyeri		0.01	0.04		99	
53001101	**Good identification**					**53001101**
					SPO	
Eubacterium saburreum		98.59	0.47	URE 1	1	
Clostridium chauvoei		1.42	0.06		99	
53001102	**Very good identification**					**53001102**
Eubacterium saburreum		100.0	0.69			
53001110	**Very good identification**					**53001110**
Eubacterium saburreum		100.0	0.80			
53001111	**Good identification**					**53001111**
Eubacterium saburreum		100.0	0.46	URE 1		
53001112	**Very good identification**					**53001112**
Eubacterium saburreum		100.0	0.67			
53001120	**Very good identification**					**53001120**
Eubacterium saburreum		99.95	0.69			
Odoribacter splanchnicus		0.05	-.07			
53001122	**Good identification**					**53001122**
Eubacterium saburreum		100.0	0.56			
53001130	**Very good identification**					**53001130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.67			
53001132	**Good identification**					**53001132**
Eubacterium saburreum		100.0	0.54			
53001140	**Good identification**					**53001140**
Eubacterium saburreum		100.0	0.47	SOR 1		
53001150	**Good identification**					**53001150**
Eubacterium saburreum		100.0	0.46	SOR 1		
53001200	**Good identification**					**53001200**
Eubacterium saburreum		97.89	0.41	RHA 1		
Eubacterium tortuosum		2.11	-.03			
53001210	**Good identification**					**53001210**
Eubacterium saburreum		100.0	0.40	RHA 1		
53001300	**Good identification**					**53001300**
Eubacterium saburreum		100.0	0.47	RHA 1		
53001310	**Good identification**					**53001310**
Eubacterium saburreum		100.0	0.46	RHA 1		
53001400	**Good identification**					**53001400**
Eubacterium saburreum		97.89	0.41	ARA 1		
Eubacterium tortuosum		2.11	-.03			
53001410	**Good identification**					**53001410**
Eubacterium saburreum		100.0	0.40	ARA 1		
53001500	**Good identification**					**53001500**
Eubacterium saburreum		99.16	0.47	ARA 1		
Odoribacter splanchnicus		0.84	-.07			
53001510	**Good identification**					**53001510**
Eubacterium saburreum		100.0	0.46	ARA 1		
53003000	**Good identification**					**53003000**
Eubacterium saburreum		97.89	0.41	MAN 1		
Eubacterium tortuosum		2.11	-.03			
53003010	**Good identification**					**53003010**
Eubacterium saburreum		100.0	0.40	MAN 1		
53003100	**Good identification**					**53003100**
Eubacterium saburreum		100.0	0.47	MAN 1		
53003110	**Good identification**					**53003110**
Eubacterium saburreum		100.0	0.46	MAN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53004000	**Low discrimination**					**53004000**
					SPO	
Eubacterium saburreum		47.35	0.37	GAL 99	1	
Clostridium bifermentans		43.92	0.24		99	
Clostridium botulinum biovar A		8.74	0.13		99	
53004010	**Good identification**					**53004010**
Eubacterium saburreum		100.0	0.35	GAL 99		
53004100	**Good identification**					**53004100**
Eubacterium saburreum		98.09	0.43	GAL 99		
Actinomyces meyeri		1.91	0.06			
53004110	**Good identification**					**53004110**
Eubacterium saburreum		100.0	0.41	GAL 99		
53005000	**Very good identification**					**53005000**
Eubacterium saburreum		99.96	0.71			
Eubacterium tortuosum		0.04	-.03			
53005001	**Good identification**					**53005001**
Eubacterium saburreum		100.0	0.37	URE 1		
53005002	**Very good identification**					**53005002**
Eubacterium saburreum		100.0	0.58			
53005010	**Very good identification**					**53005010**
Eubacterium saburreum		100.0	0.69			
53005011	**Good identification**					**53005011**
Eubacterium saburreum		100.0	0.35	URE 1		
53005012	**Very good identification**					**53005012**
Eubacterium saburreum		100.0	0.56			
53005020	**Very good identification**					**53005020**
Eubacterium saburreum		100.0	0.58			
53005022	**Good identification**					**53005022**
Eubacterium saburreum		100.0	0.45			
53005030	**Very good identification**					**53005030**
Eubacterium saburreum		100.0	0.56			
53005032	**Good identification**					**53005032**
Eubacterium saburreum		100.0	0.43			
53005040	**Good identification**					**53005040**
					SPO	
Eubacterium saburreum		96.87	0.37	SOR 1	1	
Clostridium glycolicum		3.13	-.02		99	
53005050	**Good identification**					**53005050**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.35	SOR 1		
53005100	**Very good identification**					**53005100**
Eubacterium saburreum		99.97	0.77			
Actinomyces meyeri		0.03	0.09			
53005101	**Good identification**					**53005101**
Eubacterium saburreum		100.0	0.43	URE 1		
53005102	**Very good identification**					**53005102**
Eubacterium saburreum		100.0	0.64			
53005110	**Very good identification**					**53005110**
Eubacterium saburreum		100.0	0.75			
53005111	**Good identification**					**53005111**
Eubacterium saburreum		100.0	0.41	URE 1		
53005112	**Very good identification**					**53005112**
Eubacterium saburreum		100.0	0.63			
53005120	**Very good identification**					**53005120**
Eubacterium saburreum		100.0	0.64			
53005122	**Good identification**					**53005122**
Eubacterium saburreum		100.0	0.51			
53005130	**Very good identification**					**53005130**
Eubacterium saburreum		100.0	0.63			
53005132	**Good identification**					**53005132**
Eubacterium saburreum		100.0	0.50			
53005140	**Good identification**					**53005140**
Eubacterium saburreum		100.0	0.43	SOR 1		
53005150	**Good identification**					**53005150**
Eubacterium saburreum		100.0	0.41	SOR 1		
53005200	**Good identification**					**53005200**
Eubacterium saburreum		100.0	0.37	RHA 1		
53005210	**Good identification**					**53005210**
Eubacterium saburreum		100.0	0.35	RHA 1		
53005300	**Good identification**					**53005300**
Eubacterium saburreum		100.0	0.43	RHA 1		
53005310	**Good identification**					**53005310**
Eubacterium saburreum		100.0	0.41	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53005400	**Good identification**						**53005400**
Eubacterium saburreum		100.0	0.37	ARA 1			
53005410	**Good identification**						**53005410**
Eubacterium saburreum		100.0	0.35	ARA 1			
53005500	**Good identification**						**53005500**
Eubacterium saburreum		100.0	0.43	ARA 1			
53005510	**Good identification**						**53005510**
Eubacterium saburreum		100.0	0.41	ARA 1			
53007000	**Good identification**						**53007000**
Eubacterium saburreum		100.0	0.37	MAN 1			
53007010	**Good identification**						**53007010**
Eubacterium saburreum		100.0	0.35	MAN 1			
53007100	**Good identification**						**53007100**
Eubacterium saburreum		100.0	0.43	MAN 1			
53007110	**Good identification**						**53007110**
Eubacterium saburreum		100.0	0.41	MAN 1			
53010000	**Low discrimination**						**53010000**
					SPO	GRA	
Eubacterium saburreum		28.52	0.49	GAL 99	1	85	
Clostridium bifermentans		24.74	0.36		99	99	
Fusobacterium varium		11.30	0.25		1	1	
53010010	**Acceptable identification**						**53010010**
					SPO		
Eubacterium saburreum		94.78	0.48	GAL 99	1		
Clostridium difficile		4.21	0.10		99		
Clostridium bifermentans		1.02	0.02		99		
53010020	**Acceptable identification**						**53010020**
					SPO		
Clostridium cadaveris		93.05	0.47	ESL 1	99		
Eubacterium saburreum		3.48	0.36	SUC 7	1		
Atopobium minutum		2.12	0.29		1		
53010100	**Low discrimination**						**53010100**
					GRA		
Eubacterium saburreum		75.24	0.56	GAL 99	85		
Fusobacterium mortiferum		21.90	0.48		1		
Actinomyces meyeri		2.58	0.23		99		
53010102	**Low discrimination**						**53010102**
Eubacterium saburreum		84.51	0.43	GAL 99			
Fusobacterium mortiferum		15.49	0.32				
53010110	**Good identification**						**53010110**
Eubacterium saburreum		99.64	0.54	GAL 99			
Fusobacterium mortiferum		0.36	0.14				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53010112	**Good identification**					**53010112**
Eubacterium saburreum		100.0	0.41	GAL 99		
53010120	**Low discrimination**					**53010120**
					SPO	
Eubacterium saburreum		86.29	0.43	GAL 99	1	
Clostridium cadaveris		9.99	0.13		99	
Actinomyces meyeri		2.28	0.08		1	
53010130	**Good identification**					**53010130**
Eubacterium saburreum		100.0	0.41	GAL 99		
53011000	**Very good identification**					**53011000**
					GRA	
Eubacterium saburreum		99.38	0.83		85	
Fusobacterium mortiferum		0.44	0.45		1	
Atopobium minutum		0.11	0.29		99	
53011001	**Acceptable identification**					**53011001**
					SPO	
Eubacterium saburreum		91.93	0.49	URE 1	1	
Clostridium sordellii		8.08	0.15		99	
53011002	**Very good identification**					**53011002**
					GRA	
Eubacterium saburreum		99.27	0.71		85	
Atopobium minutum		0.41	0.26		99	
Fusobacterium mortiferum		0.28	0.28		1	
53011003	**Good identification**					**53011003**
Eubacterium saburreum		100.0	0.36	URE 1		
53011010	**Very good identification**					**53011010**
Eubacterium saburreum		100.0	0.82			
53011011	**Good identification**					**53011011**
Eubacterium saburreum		100.0	0.48	URE 1		
53011012	**Very good identification**					**53011012**
Eubacterium saburreum		100.0	0.69			
53011020	**Very good identification**					**53011020**
					SPO	
Eubacterium saburreum		99.45	0.71		1	
Clostridium cadaveris		0.27	0.13		99	
Atopobium minutum		0.20	0.21		1	
53011021	**Good identification**					**53011021**
Eubacterium saburreum		100.0	0.36	URE 1		
53011022	**Very good identification**					**53011022**
Eubacterium saburreum		99.23	0.58			
Atopobium minutum		0.77	0.18			
53011030	**Very good identification**					**53011030**
Eubacterium saburreum		100.0	0.69			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53011032	**Very good identification**						**53011032**
Eubacterium saburreum		100.0	0.56				
53011040	**Good identification**						**53011040**
Eubacterium saburreum		100.0	0.49	SOR 1			
53011042	**Good identification**						**53011042**
Eubacterium saburreum		100.0	0.36	SOR 1			
53011050	**Good identification**						**53011050**
Eubacterium saburreum		100.0	0.48	SOR 1			
53011060	**Good identification**						**53011060**
Eubacterium saburreum		100.0	0.36	SOR 1			
53011100	**Very good identification**						**53011100**
					GRA		
Eubacterium saburreum		99.50	0.90		85		
Fusobacterium mortiferum		0.44	0.51		1		
Actinomyces meyeri		0.05	0.26		99		
53011101	**Good identification**						**53011101**
					SPO	GRA	
Eubacterium saburreum		98.47	0.56	URE 1	1	85	
Clostridium chauvoei		1.10	0.12		99	99	
Fusobacterium mortiferum		0.43	0.17		1	1	
53011102	**Very good identification**						**53011102**
					GRA		
Eubacterium saburreum		99.69	0.77		85		
Fusobacterium mortiferum		0.28	0.35		1		
Lactobacillus fermentum		0.03	0.20		99		
53011103	**Good identification**						**53011103**
					SPO		
Eubacterium saburreum		98.90	0.43	URE 1	1		
Clostridium chauvoei		1.10	-.01		99		
53011110	**Very good identification**						**53011110**
Eubacterium saburreum		100.0	0.88				
Fusobacterium mortiferum		0.01	0.17				
53011111	**Good identification**						**53011111**
Eubacterium saburreum		100.0	0.54	URE 1			
53011112	**Very good identification**						**53011112**
Eubacterium saburreum		100.0	0.75				
53011113	**Good identification**						**53011113**
Eubacterium saburreum		100.0	0.41	URE 1			
53011120	**Very good identification**						**53011120**
					GRA		
Eubacterium saburreum		99.89	0.77		85		
Actinomyces meyeri		0.04	0.11		99		
Odoribacter splanchnicus		0.03	-.03		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53011121	**Good identification**					**53011121**
					SPO	
Eubacterium saburreum		97.94	0.43	URE 1	1	
Clostridium chauvoei		2.07	0.04		99	
53011122	**Very good identification**					**53011122**
Eubacterium saburreum		100.0	0.64			
53011130	**Very good identification**					**53011130**
Eubacterium saburreum		100.0	0.75			
53011131	**Good identification**					**53011131**
Eubacterium saburreum		100.0	0.41	URE 1		
53011132	**Very good identification**					**53011132**
Eubacterium saburreum		100.0	0.62			
53011140	**Good identification**					**53011140**
Eubacterium saburreum		99.56	0.56	SOR 1		
Fusobacterium mortiferum		0.44	0.17			
53011142	**Good identification**					**53011142**
Eubacterium saburreum		100.0	0.43	SOR 1		
53011150	**Good identification**					**53011150**
Eubacterium saburreum		100.0	0.54	SOR 1		
53011152	**Good identification**					**53011152**
Eubacterium saburreum		100.0	0.41	SOR 1		
53011160	**Good identification**					**53011160**
Eubacterium saburreum		100.0	0.43	SOR 1		
53011170	**Good identification**					**53011170**
Eubacterium saburreum		100.0	0.41	SOR 1		
53011200	**Good identification**					**53011200**
Eubacterium saburreum		100.0	0.49	RHA 1		
53011202	**Good identification**					**53011202**
Eubacterium saburreum		100.0	0.36	RHA 1		
53011210	**Good identification**					**53011210**
Eubacterium saburreum		100.0	0.48	RHA 1		
53011220	**Good identification**					**53011220**
Eubacterium saburreum		100.0	0.36	RHA 1		
53011300	**Good identification**					**53011300**
Eubacterium saburreum		99.56	0.56	RHA 1		
Fusobacterium mortiferum		0.44	0.17			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53011302	**Good identification**					**53011302**
Eubacterium saburreum		100.0	0.43	RHA 1		
53011310	**Good identification**					**53011310**
Eubacterium saburreum		100.0	0.54	RHA 1		
53011312	**Good identification**					**53011312**
Eubacterium saburreum		100.0	0.41	RHA 1		
53011320	**Good identification**					**53011320**
Eubacterium saburreum		100.0	0.43	RHA 1		
53011330	**Good identification**					**53011330**
Eubacterium saburreum		100.0	0.41	RHA 1		
53011400	**Good identification**					**53011400**
Eubacterium saburreum		98.51	0.49	ARA 1		
Lactobacillus fermentum		0.76	0.16			
Actinomyces meyeri		0.74	0.05			
53011402	**Good identification**					**53011402**
Eubacterium saburreum		95.83	0.36	ARA 1		
Lactobacillus fermentum		4.17	0.16			
53011410	**Good identification**					**53011410**
Eubacterium saburreum		100.0	0.48	ARA 1		
53011420	**Acceptable identification**					**53011420**
Eubacterium saburreum		93.50	0.36	ARA 1		
Odoribacter splanchnicus		6.50	-.03			
53011500	**Good identification**					**53011500**
Eubacterium saburreum		97.29	0.56	ARA 1		
Actinomyces meyeri		1.25	0.16			
Lactobacillus fermentum		0.52	0.20			
53011502	**Good identification**					**53011502**
Eubacterium saburreum		97.05	0.43	ARA 1		
Lactobacillus fermentum		2.95	0.20			
53011510	**Good identification**					**53011510**
Eubacterium saburreum		100.0	0.54	ARA 1		
53011512	**Good identification**					**53011512**
Eubacterium saburreum		100.0	0.41	ARA 1		
53011530	**Good identification**					**53011530**
Eubacterium saburreum		96.49	0.41	ARA 1		
Odoribacter splanchnicus		3.52	-.03			
53013000	**Good identification**					**53013000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.46	0.49	MAN 1		
Eubacterium limosum		0.54	-.06			
53013002	**Good identification**					**53013002**
Eubacterium saburreum		100.0	0.36	MAN 1		
53013010	**Good identification**					**53013010**
Eubacterium saburreum		100.0	0.48	MAN 1		
53013020	**Good identification**					**53013020**
					CAT	
Eubacterium saburreum		95.96	0.36	MAN 1	1	
Propionibacterium acnes		4.04	0.02		95	
53013100	**Good identification**					**53013100**
Eubacterium saburreum		99.56	0.56	MAN 1		
Fusobacterium mortiferum		0.44	0.17			
53013102	**Good identification**					**53013102**
Eubacterium saburreum		100.0	0.43	MAN 1		
53013110	**Good identification**					**53013110**
Eubacterium saburreum		100.0	0.54	MAN 1		
53013112	**Good identification**					**53013112**
Eubacterium saburreum		100.0	0.41	MAN 1		
53013120	**Good identification**					**53013120**
Eubacterium saburreum		100.0	0.43	MAN 1		
53013130	**Good identification**					**53013130**
Eubacterium saburreum		100.0	0.41	MAN 1		
53014000	**Low discrimination**					**53014000**
					SPO	
Eubacterium saburreum		82.95	0.45	GAL 99	1	
Actinomyces meyeri		5.98	0.18		1	
Clostridium glycolicum		3.81	0.09		99	
53014010	**Good identification**					**53014010**
Eubacterium saburreum		100.0	0.43	GAL 99		
53014100	**Low discrimination**					**53014100**
					GRA	
Eubacterium saburreum		85.90	0.51	GAL 99	85	
Actinomyces meyeri		10.61	0.28		99	
Fusobacterium mortiferum		3.50	0.29		1	
53014102	**Good identification**					**53014102**
					GRA	
Eubacterium saburreum		95.49	0.38	GAL 99	85	
Fusobacterium mortiferum		2.45	0.13		1	
Actinomyces meyeri		2.07	0.02		99	
53014110	**Good identification**					**53014110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.49	GAL 99		
53014112	**Good identification**					**53014112**
Eubacterium saburreum		100.0	0.37	GAL 99		
53014120	**Acceptable identification**					**53014120**
Eubacterium saburreum		91.29	0.38	GAL 99		
Actinomyces meyeri		8.71	0.13			
53014130	**Good identification**					**53014130**
Eubacterium saburreum		100.0	0.37	GAL 99		
53015000	**Very good identification**					**53015000**
					GRA	
Eubacterium saburreum		99.82	0.79		85	
Actinomyces meyeri		0.11	0.21		99	
Fusobacterium mortiferum		0.06	0.26		1	
53015001	**Good identification**					**53015001**
Eubacterium saburreum		100.0	0.45	URE 1		
53015002	**Very good identification**					**53015002**
Eubacterium saburreum		99.95	0.66			
Lactobacillus fermentum		0.05	0.12			
53015010	**Very good identification**					**53015010**
Eubacterium saburreum		100.0	0.77			
53015011	**Good identification**					**53015011**
Eubacterium saburreum		100.0	0.43	URE 1		
53015012	**Very good identification**					**53015012**
Eubacterium saburreum		100.0	0.64			
53015020	**Very good identification**					**53015020**
Eubacterium saburreum		99.92	0.66			
Actinomyces meyeri		0.08	0.06			
53015022	**Good identification**					**53015022**
Eubacterium saburreum		100.0	0.53			
53015030	**Very good identification**					**53015030**
Eubacterium saburreum		100.0	0.64			
53015032	**Good identification**					**53015032**
Eubacterium saburreum		100.0	0.52			
53015040	**Acceptable identification**					**53015040**
					SPO	
Eubacterium saburreum		91.17	0.45	SOR 1	1	
Clostridium glycolicum		8.83	0.15		99	
53015050	**Good identification**					**53015050**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.43	SOR 1		
53015100	**Very good identification**					**53015100**
					GRA	
Eubacterium saburreum		99.75	0.85		85	
Actinomyces meyeri		0.19	0.31		99	
Fusobacterium mortiferum		0.06	0.32		1	
53015101	**Good identification**					**53015101**
Eubacterium saburreum		100.0	0.51	URE 1		
53015102	**Very good identification**					**53015102**
					GRA	
Eubacterium saburreum		99.90	0.72		85	
Fusobacterium mortiferum		0.04	0.16		1	
Lactobacillus fermentum		0.03	0.16		99	
53015103	**Good identification**					**53015103**
Eubacterium saburreum		100.0	0.38	URE 1		
53015110	**Very good identification**					**53015110**
Eubacterium saburreum		100.0	0.84			
53015111	**Good identification**					**53015111**
Eubacterium saburreum		100.0	0.49	URE 1		
53015112	**Very good identification**					**53015112**
Eubacterium saburreum		100.0	0.71			
53015113	**Good identification**					**53015113**
Eubacterium saburreum		100.0	0.37	URE 1		
53015120	**Very good identification**					**53015120**
Eubacterium saburreum		99.86	0.72			
Actinomyces meyeri		0.14	0.16			
53015121	**Good identification**					**53015121**
Eubacterium saburreum		100.0	0.38	URE 1		
53015122	**Very good identification**					**53015122**
Eubacterium saburreum		100.0	0.59			
53015130	**Very good identification**					**53015130**
Eubacterium saburreum		100.0	0.71			
53015131	**Good identification**					**53015131**
Eubacterium saburreum		100.0	0.37	URE 1		
53015132	**Very good identification**					**53015132**
Eubacterium saburreum		100.0	0.58			
53015140	**Good identification**					**53015140**
Eubacterium saburreum		100.0	0.51	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53015142	**Good identification**					**53015142**
Eubacterium saburreum		100.0	0.38	SOR 1		
53015150	**Good identification**					**53015150**
Eubacterium saburreum		100.0	0.49	SOR 1		
53015152	**Good identification**					**53015152**
Eubacterium saburreum		100.0	0.37	SOR 1		
53015160	**Good identification**					**53015160**
Eubacterium saburreum		100.0	0.38	SOR 1		
53015170	**Good identification**					**53015170**
Eubacterium saburreum		100.0	0.37	SOR 1		
53015200	**Good identification**					**53015200**
Eubacterium saburreum		100.0	0.45	RHA 1		
53015210	**Good identification**					**53015210**
Eubacterium saburreum		100.0	0.43	RHA 1		
53015300	**Good identification**					**53015300**
Eubacterium saburreum		100.0	0.51	RHA 1		
53015302	**Good identification**					**53015302**
Eubacterium saburreum		100.0	0.38	RHA 1		
53015310	**Good identification**					**53015310**
Eubacterium saburreum		100.0	0.49	RHA 1		
53015312	**Good identification**					**53015312**
Eubacterium saburreum		100.0	0.37	RHA 1		
53015320	**Good identification**					**53015320**
Eubacterium saburreum		100.0	0.38	RHA 1		
53015330	**Good identification**					**53015330**
Eubacterium saburreum		100.0	0.37	RHA 1		
53015400	**Good identification**					**53015400**
Eubacterium saburreum		96.58	0.45	ARA 1		
Actinomyces meyeri		2.61	0.10			
Lactobacillus fermentum		0.81	0.12			
53015410	**Good identification**					**53015410**
Eubacterium saburreum		100.0	0.43	ARA 1		
53015500	**Good identification**					**53015500**
Eubacterium saburreum		95.04	0.51	ARA 1		
Actinomyces meyeri		4.40	0.21			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus fermentum		0.56	0.16			
53015502	**Good identification**					**53015502**
Eubacterium saburreum		96.79	0.38	ARA 1		
Lactobacillus fermentum		3.21	0.16			
53015510	**Good identification**					**53015510**
Eubacterium saburreum		100.0	0.49	ARA 1		
53015512	**Good identification**					**53015512**
Eubacterium saburreum		100.0	0.37	ARA 1		
53015520	**Acceptable identification**					**53015520**
					GRA	
Eubacterium saburreum		91.65	0.38	ARA 1	85	
Odoribacter splanchnicus		5.07	-.03		1	
Actinomyces meyeri		3.28	0.06		99	
53015530	**Good identification**					**53015530**
Eubacterium saburreum		100.0	0.37	ARA 1		
53017000	**Good identification**					**53017000**
Eubacterium saburreum		100.0	0.45	MAN 1		
53017010	**Good identification**					**53017010**
Eubacterium saburreum		100.0	0.43	MAN 1		
53017100	**Good identification**					**53017100**
Eubacterium saburreum		100.0	0.51	MAN 1		
53017102	**Good identification**					**53017102**
Eubacterium saburreum		100.0	0.38	MAN 1		
53017110	**Good identification**					**53017110**
Eubacterium saburreum		100.0	0.49	MAN 1		
53017112	**Good identification**					**53017112**
Eubacterium saburreum		100.0	0.37	MAN 1		
53017120	**Good identification**					**53017120**
Eubacterium saburreum		100.0	0.38	MAN 1		
53017130	**Good identification**					**53017130**
Eubacterium saburreum		100.0	0.37	MAN 1		
53020000	**Low discrimination**					**53020000**
					SPO	
Clostridium botulinum biovar A		62.27	0.36	IND 1	99	
Eubacterium saburreum		16.89	0.38	SUC 7	1	
Clostridium bifermentans		12.65	0.24		99	
53020010	**Good identification**					**53020010**
					SPO	
Eubacterium saburreum		95.65	0.37	GAL 99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Clostridium botulinum biovar A	4.36	0.02		99	
53020100	**Good identification**					**53020100**
					SPO	
	Eubacterium saburreum	98.43	0.44	GAL 99	1	
	Clostridium botulinum biovar A	1.57	0.02		99	
53020110	**Good identification**					**53020110**
	Eubacterium saburreum	100.0	0.43	GAL 99		
53021000	**Very good identification**					**53021000**
					SPO	
	Eubacterium saburreum	99.93	0.72		1	
	Clostridium botulinum biovar A	0.04	0.02		99	
	Eubacterium tortuosum	0.03	-.03		1	
53021001	**Good identification**					**53021001**
	Eubacterium saburreum	100.0	0.38	URE 1		
53021002	**Very good identification**					**53021002**
	Eubacterium saburreum	100.0	0.59			
53021010	**Very good identification**					**53021010**
	Eubacterium saburreum	100.0	0.71			
53021011	**Good identification**					**53021011**
	Eubacterium saburreum	100.0	0.37	URE 1		
53021012	**Very good identification**					**53021012**
	Eubacterium saburreum	100.0	0.58			
53021020	**Very good identification**					**53021020**
	Eubacterium saburreum	100.0	0.59			
53021022	**Good identification**					**53021022**
	Eubacterium saburreum	100.0	0.46			
53021030	**Very good identification**					**53021030**
	Eubacterium saburreum	100.0	0.58			
53021032	**Good identification**					**53021032**
	Eubacterium saburreum	100.0	0.45			
53021040	**Good identification**					**53021040**
	Eubacterium saburreum	100.0	0.38	SOR 1		
53021050	**Good identification**					**53021050**
	Eubacterium saburreum	100.0	0.37	SOR 1		
53021100	**Very good identification**					**53021100**
	Eubacterium saburreum	100.0	0.79			
53021101	**Good identification**					**53021101**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Eubacterium saburreum	100.0	0.44	URE 1		
53021102	**Very good identification**					**53021102**
	Eubacterium saburreum	100.0	0.66			
53021110	**Very good identification**					**53021110**
	Eubacterium saburreum	100.0	0.77			
53021111	**Good identification**					**53021111**
	Eubacterium saburreum	100.0	0.43	URE 1		
53021112	**Very good identification**					**53021112**
	Eubacterium saburreum	100.0	0.64			
53021120	**Very good identification**					**53021120**
	Eubacterium saburreum	100.0	0.66			
53021122	**Good identification**					**53021122**
	Eubacterium saburreum	100.0	0.53			
53021130	**Very good identification**					**53021130**
	Eubacterium saburreum	100.0	0.64			
53021132	**Good identification**					**53021132**
	Eubacterium saburreum	100.0	0.51			
53021140	**Good identification**					**53021140**
	Eubacterium saburreum	100.0	0.44	SOR 1		
53021150	**Good identification**					**53021150**
	Eubacterium saburreum	100.0	0.43	SOR 1		
53021200	**Good identification**					**53021200**
	Eubacterium saburreum	100.0	0.38	RHA 1		
53021210	**Good identification**					**53021210**
	Eubacterium saburreum	100.0	0.37	RHA 1		
53021300	**Good identification**					**53021300**
	Eubacterium saburreum	100.0	0.44	RHA 1		
53021310	**Good identification**					**53021310**
	Eubacterium saburreum	100.0	0.43	RHA 1		
53021400	**Good identification**					**53021400**
	Eubacterium saburreum	100.0	0.38	ARA 1		
53021410	**Good identification**					**53021410**
	Eubacterium saburreum	100.0	0.37	ARA 1		
53021500	**Good identification**					**53021500**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Eubacterium saburreum	100.0	0.44	ARA 1		
53021510	**Good identification**					**53021510**
	Eubacterium saburreum	100.0	0.43	ARA 1		
53023000	**Good identification**					**53023000**
	Eubacterium saburreum	100.0	0.38	MAN 1		
53023010	**Good identification**					**53023010**
	Eubacterium saburreum	100.0	0.37	MAN 1		
53023100	**Good identification**					**53023100**
	Eubacterium saburreum	100.0	0.44	MAN 1		
53023110	**Good identification**					**53023110**
	Eubacterium saburreum	100.0	0.43	MAN 1		
53024100	**Good identification**					**53024100**
	Eubacterium saburreum	100.0	0.40	GAL 99		
53024110	**Good identification**					**53024110**
	Eubacterium saburreum	100.0	0.38	GAL 99		
53025000	**Very good identification**					**53025000**
	Eubacterium saburreum	100.0	0.68			
53025002	**Good identification**					**53025002**
	Eubacterium saburreum	100.0	0.55			
53025010	**Very good identification**					**53025010**
	Eubacterium saburreum	100.0	0.66			
53025012	**Good identification**					**53025012**
	Eubacterium saburreum	100.0	0.53			
53025020	**Good identification**					**53025020**
	Eubacterium saburreum	100.0	0.55			
53025022	**Good identification**					**53025022**
	Eubacterium saburreum	100.0	0.42			
53025030	**Good identification**					**53025030**
	Eubacterium saburreum	100.0	0.53			
53025032	**Good identification**					**53025032**
	Eubacterium saburreum	100.0	0.40			
53025100	**Very good identification**					**53025100**
	Eubacterium saburreum	100.0	0.74			
53025101	**Good identification**					**53025101**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Eubacterium saburreum	100.0	0.40	URE 1		
53025102	**Very good identification**					**53025102**
	Eubacterium saburreum	100.0	0.61			
53025110	**Very good identification**					**53025110**
	Eubacterium saburreum	100.0	0.72			
53025111	**Good identification**					**53025111**
	Eubacterium saburreum	100.0	0.38	URE 1		
53025112	**Very good identification**					**53025112**
	Eubacterium saburreum	100.0	0.60			
53025120	**Very good identification**					**53025120**
	Eubacterium saburreum	100.0	0.61			
53025122	**Good identification**					**53025122**
	Eubacterium saburreum	100.0	0.48			
53025130	**Very good identification**					**53025130**
	Eubacterium saburreum	100.0	0.60			
53025132	**Good identification**					**53025132**
	Eubacterium saburreum	100.0	0.47			
53025140	**Good identification**					**53025140**
	Eubacterium saburreum	100.0	0.40	SOR 1		
53025150	**Good identification**					**53025150**
	Eubacterium saburreum	100.0	0.38	SOR 1		
53025300	**Good identification**					**53025300**
	Eubacterium saburreum	100.0	0.40	RHA 1		
53025310	**Good identification**					**53025310**
	Eubacterium saburreum	100.0	0.38	RHA 1		
53025500	**Good identification**					**53025500**
	Eubacterium saburreum	100.0	0.40	ARA 1		
53025510	**Good identification**					**53025510**
	Eubacterium saburreum	100.0	0.38	ARA 1		
53027100	**Good identification**					**53027100**
	Eubacterium saburreum	100.0	0.40	MAN 1		
53027110	**Good identification**					**53027110**
	Eubacterium saburreum	100.0	0.38	MAN 1		
53030000	**Low discrimination**					**53030000**

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Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		79.96	0.46	GAL 99	1	85	
Fusobacterium mortiferum		10.43	0.33		1	1	
Clostridium botulinum biovar A		6.27	0.16		99	99	
53030010	**Good identification**						**53030010**
Eubacterium saburreum		100.0	0.45	GAL 99			
53030100	**Low discrimination**						**53030100**
Eubacterium saburreum		88.46	0.53	GAL 99			
Fusobacterium mortiferum		11.54	0.39				
53030102	**Acceptable identification**						**53030102**
Eubacterium saburreum		92.41	0.40	GAL 99			
Fusobacterium mortiferum		7.59	0.23				
53030110	**Good identification**						**53030110**
Eubacterium saburreum		100.0	0.51	GAL 99			
53030112	**Good identification**						**53030112**
Eubacterium saburreum		100.0	0.38	GAL 99			
53030120	**Good identification**						**53030120**
Eubacterium saburreum		100.0	0.40	GAL 99			
53030130	**Good identification**						**53030130**
Eubacterium saburreum		100.0	0.38	GAL 99			
53031000	**Very good identification**						**53031000**
Eubacterium saburreum		99.80	0.80				
Fusobacterium mortiferum		0.20	0.36				
53031001	**Good identification**						**53031001**
Eubacterium saburreum		100.0	0.46	URE 1			
53031002	**Very good identification**						**53031002**
Eubacterium saburreum		99.88	0.68				
Fusobacterium mortiferum		0.12	0.19				
53031010	**Very good identification**						**53031010**
Eubacterium saburreum		100.0	0.79				
53031011	**Good identification**						**53031011**
Eubacterium saburreum		100.0	0.45	URE 1			
53031012	**Very good identification**						**53031012**
Eubacterium saburreum		100.0	0.66				
53031020	**Very good identification**						**53031020**
Eubacterium saburreum		100.0	0.68				
53031022	**Good identification**						**53031022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.55			
53031030	**Very good identification**					**53031030**
Eubacterium saburreum		100.0	0.66			
53031032	**Good identification**					**53031032**
Eubacterium saburreum		100.0	0.53			
53031040	**Good identification**					**53031040**
Eubacterium saburreum		100.0	0.46	SOR 1		
53031050	**Good identification**					**53031050**
Eubacterium saburreum		100.0	0.45	SOR 1		
53031100	**Very good identification**					**53031100**
Eubacterium saburreum		99.80	0.87			
Fusobacterium mortiferum		0.20	0.42			
53031101	**Good identification**					**53031101**
Eubacterium saburreum		100.0	0.53	URE 1		
53031102	**Very good identification**					**53031102**
Eubacterium saburreum		99.88	0.74			
Fusobacterium mortiferum		0.12	0.26			
53031103	**Good identification**					**53031103**
Eubacterium saburreum		100.0	0.40	URE 1		
53031110	**Very good identification**					**53031110**
Eubacterium saburreum		100.0	0.85			
53031111	**Good identification**					**53031111**
Eubacterium saburreum		100.0	0.51	URE 1		
53031112	**Very good identification**					**53031112**
Eubacterium saburreum		100.0	0.72			
53031113	**Good identification**					**53031113**
Eubacterium saburreum		100.0	0.38	URE 1		
53031120	**Very good identification**					**53031120**
Eubacterium saburreum		100.0	0.74			
53031121	**Good identification**					**53031121**
Eubacterium saburreum		100.0	0.40	URE 1		
53031122	**Very good identification**					**53031122**
Eubacterium saburreum		100.0	0.61			
53031130	**Very good identification**					**53031130**
Eubacterium saburreum		100.0	0.72			

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
53031131	**Good identification**					**53031131**
Eubacterium saburreum		100.0	0.38	URE 1		
53031132	**Very good identification**					**53031132**
Eubacterium saburreum		100.0	0.59			
53031140	**Good identification**					**53031140**
Eubacterium saburreum		100.0	0.53	SOR 1		
53031142	**Good identification**					**53031142**
Eubacterium saburreum		100.0	0.40	SOR 1		
53031150	**Good identification**					**53031150**
Eubacterium saburreum		100.0	0.51	SOR 1		
53031152	**Good identification**					**53031152**
Eubacterium saburreum		100.0	0.38	SOR 1		
53031160	**Good identification**					**53031160**
Eubacterium saburreum		100.0	0.40	SOR 1		
53031170	**Good identification**					**53031170**
Eubacterium saburreum		100.0	0.38	SOR 1		
53031200	**Good identification**					**53031200**
Eubacterium saburreum		100.0	0.46	RHA 1		
53031210	**Good identification**					**53031210**
Eubacterium saburreum		100.0	0.45	RHA 1		
53031300	**Good identification**					**53031300**
Eubacterium saburreum		100.0	0.53	RHA 1		
53031302	**Good identification**					**53031302**
Eubacterium saburreum		100.0	0.40	RHA 1		
53031310	**Good identification**					**53031310**
Eubacterium saburreum		100.0	0.51	RHA 1		
53031312	**Good identification**					**53031312**
Eubacterium saburreum		100.0	0.38	RHA 1		
53031320	**Good identification**					**53031320**
Eubacterium saburreum		100.0	0.40	RHA 1		
53031330	**Good identification**					**53031330**
Eubacterium saburreum		100.0	0.38	RHA 1		
53031400	**Good identification**					**53031400**
Eubacterium saburreum		100.0	0.46	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53031410	**Good identification**					**53031410**
Eubacterium saburreum		100.0	0.45	ARA 1		
53031500	**Good identification**					**53031500**
Eubacterium saburreum		100.0	0.53	ARA 1		
53031502	**Good identification**					**53031502**
Eubacterium saburreum		100.0	0.40	ARA 1		
53031510	**Good identification**					**53031510**
Eubacterium saburreum		100.0	0.51	ARA 1		
53031512	**Good identification**					**53031512**
Eubacterium saburreum		100.0	0.38	ARA 1		
53031520	**Good identification**					**53031520**
Eubacterium saburreum		95.72	0.40	ARA 1		
Odoribacter splanchnicus		4.28	-.03			
53031530	**Good identification**					**53031530**
Eubacterium saburreum		100.0	0.38	ARA 1		
53033000	**Good identification**					**53033000**
Eubacterium saburreum		100.0	0.46	MAN 1		
53033010	**Good identification**					**53033010**
Eubacterium saburreum		100.0	0.45	MAN 1		
53033100	**Good identification**					**53033100**
Eubacterium saburreum		100.0	0.53	MAN 1		
53033102	**Good identification**					**53033102**
Eubacterium saburreum		100.0	0.40	MAN 1		
53033110	**Good identification**					**53033110**
Eubacterium saburreum		100.0	0.51	MAN 1		
53033112	**Good identification**					**53033112**
Eubacterium saburreum		100.0	0.38	MAN 1		
53033120	**Good identification**					**53033120**
Eubacterium saburreum		100.0	0.40	MAN 1		
53033130	**Good identification**					**53033130**
Eubacterium saburreum		100.0	0.38	MAN 1		
53034000	**Good identification**					**53034000**
Eubacterium saburreum		98.21	0.42	GAL 99		
Fusobacterium mortiferum		1.79	0.14			
53034010	**Good identification**					**53034010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.40	GAL 99		
53034100	**Good identification**					**53034100**
Eubacterium saburreum		98.21	0.48	GAL 99		
Fusobacterium mortiferum		1.79	0.20			
53034102	**Good identification**					**53034102**
Eubacterium saburreum		100.0	0.35	GAL 99		
53034110	**Good identification**					**53034110**
Eubacterium saburreum		100.0	0.46	GAL 99		
53034120	**Good identification**					**53034120**
Eubacterium saburreum		100.0	0.35	GAL 99		
53035000	**Very good identification**					**53035000**
Eubacterium saburreum		99.97	0.76			
Fusobacterium mortiferum		0.03	0.17			
53035001	**Good identification**					**53035001**
Eubacterium saburreum		100.0	0.42	URE 1		
53035002	**Very good identification**					**53035002**
Eubacterium saburreum		100.0	0.63			
53035010	**Very good identification**					**53035010**
Eubacterium saburreum		100.0	0.74			
53035011	**Good identification**					**53035011**
Eubacterium saburreum		100.0	0.40	URE 1		
53035012	**Very good identification**					**53035012**
Eubacterium saburreum		100.0	0.61			
53035020	**Very good identification**					**53035020**
Eubacterium saburreum		100.0	0.63			
53035022	**Good identification**					**53035022**
Eubacterium saburreum		100.0	0.50			
53035030	**Very good identification**					**53035030**
Eubacterium saburreum		100.0	0.61			
53035032	**Good identification**					**53035032**
Eubacterium saburreum		100.0	0.49			
53035040	**Good identification**					**53035040**
Eubacterium saburreum		100.0	0.42	SOR 1		
53035050	**Good identification**					**53035050**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.40	SOR 1		
53035100	**Very good identification**					**53035100**
Eubacterium saburreum		99.97	0.82			
Fusobacterium mortiferum		0.03	0.23			
53035101	**Good identification**					**53035101**
Eubacterium saburreum		100.0	0.48	URE 1		
53035102	**Very good identification**					**53035102**
Eubacterium saburreum		100.0	0.69			
53035103	**Good identification**					**53035103**
Eubacterium saburreum		100.0	0.35	URE 1		
53035110	**Very good identification**					**53035110**
Eubacterium saburreum		100.0	0.81			
53035111	**Good identification**					**53035111**
Eubacterium saburreum		100.0	0.46	URE 1		
53035112	**Very good identification**					**53035112**
Eubacterium saburreum		100.0	0.68			
53035120	**Very good identification**					**53035120**
Eubacterium saburreum		100.0	0.69			
53035121	**Good identification**					**53035121**
Eubacterium saburreum		100.0	0.35	URE 1		
53035122	**Very good identification**					**53035122**
Eubacterium saburreum		100.0	0.56			
53035130	**Very good identification**					**53035130**
Eubacterium saburreum		100.0	0.68			
53035132	**Good identification**					**53035132**
Eubacterium saburreum		100.0	0.55			
53035140	**Good identification**					**53035140**
Eubacterium saburreum		100.0	0.48	SOR 1		
53035142	**Good identification**					**53035142**
Eubacterium saburreum		100.0	0.35	SOR 1		
53035150	**Good identification**					**53035150**
Eubacterium saburreum		100.0	0.46	SOR 1		
53035160	**Good identification**					**53035160**
Eubacterium saburreum		100.0	0.35	SOR 1		
53035200	**Good identification**					**53035200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.42	RHA 1		
53035210	**Good identification**					**53035210**
Eubacterium saburreum		100.0	0.40	RHA 1		
53035300	**Good identification**					**53035300**
Eubacterium saburreum		100.0	0.48	RHA 1		
53035302	**Good identification**					**53035302**
Eubacterium saburreum		100.0	0.35	RHA 1		
53035310	**Good identification**					**53035310**
Eubacterium saburreum		100.0	0.46	RHA 1		
53035320	**Good identification**					**53035320**
Eubacterium saburreum		100.0	0.35	RHA 1		
53035400	**Good identification**					**53035400**
Eubacterium saburreum		100.0	0.42	ARA 1		
53035410	**Good identification**					**53035410**
Eubacterium saburreum		100.0	0.40	ARA 1		
53035500	**Good identification**					**53035500**
Eubacterium saburreum		100.0	0.48	ARA 1		
53035502	**Good identification**					**53035502**
Eubacterium saburreum		100.0	0.35	ARA 1		
53035510	**Good identification**					**53035510**
Eubacterium saburreum		100.0	0.46	ARA 1		
53035520	**Good identification**					**53035520**
Eubacterium saburreum		100.0	0.35	ARA 1		
53037000	**Good identification**					**53037000**
Eubacterium saburreum		100.0	0.42	MAN 1		
53037010	**Good identification**					**53037010**
Eubacterium saburreum		100.0	0.40	MAN 1		
53037100	**Good identification**					**53037100**
Eubacterium saburreum		100.0	0.48	MAN 1		
53037102	**Good identification**					**53037102**
Eubacterium saburreum		100.0	0.35	MAN 1		
53037110	**Good identification**					**53037110**
Eubacterium saburreum		100.0	0.46	MAN 1		
53037120	**Good identification**					**53037120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.35	MAN 1		
53041000	**Low discrimination**					**53041000**
Eubacterium saburreum		80.85	0.41	CEL 1		
Eubacterium tortuosum		19.15	0.15			
53041010	**Good identification**					**53041010**
Eubacterium saburreum		100.0	0.40	CEL 1		
53041100	**Good identification**					**53041100**
Eubacterium saburreum		100.0	0.47	CEL 1		
53041110	**Good identification**					**53041110**
Eubacterium saburreum		100.0	0.46	CEL 1		
53045000	**Good identification**					**53045000**
Eubacterium saburreum		100.0	0.37	CEL 1		
53045010	**Good identification**					**53045010**
Eubacterium saburreum		100.0	0.35	CEL 1		
53045100	**Good identification**					**53045100**
Eubacterium saburreum		100.0	0.43	CEL 1		
53045110	**Good identification**					**53045110**
Eubacterium saburreum		100.0	0.41	CEL 1		
53050100	**Acceptable identification**					**53050100**
Fusobacterium mortiferum		90.57	0.40	IND 1		
Eubacterium saburreum		9.43	0.21			
53051000	**Low discrimination**					**53051000**
Eubacterium saburreum		87.29	0.49	CEL 1		
Fusobacterium mortiferum		12.71	0.37			
53051002	**Acceptable identification**					**53051002**
Eubacterium saburreum		91.60	0.36	CEL 1		
Fusobacterium mortiferum		8.40	0.20			
53051010	**Good identification**					**53051010**
Eubacterium saburreum		100.0	0.48	CEL 1		
53051020	**Good identification**					**53051020**
Eubacterium saburreum		100.0	0.36	CEL 1		
53051100	**Low discrimination**					**53051100**
Eubacterium saburreum		87.29	0.56	CEL 1		
Fusobacterium mortiferum		12.71	0.43			
53051102	**Acceptable identification**					**53051102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		91.60	0.43	CEL 1		
Fusobacterium mortiferum		8.40	0.27			
53051110	**Good identification**					**53051110**
Eubacterium saburreum		100.0	0.54	CEL 1		
53051112	**Good identification**					**53051112**
Eubacterium saburreum		100.0	0.41	CEL 1		
53051120	**Good identification**					**53051120**
Eubacterium saburreum		100.0	0.43	CEL 1		
53051130	**Good identification**					**53051130**
Eubacterium saburreum		100.0	0.41	CEL 1		
53055000	**Good identification**					**53055000**
Eubacterium saburreum		98.01	0.45	CEL 1		
Fusobacterium mortiferum		1.99	0.17			
53055010	**Good identification**					**53055010**
Eubacterium saburreum		100.0	0.43	CEL 1		
53055100	**Good identification**					**53055100**
Eubacterium saburreum		98.01	0.51	CEL 1		
Fusobacterium mortiferum		1.99	0.24			
53055102	**Good identification**					**53055102**
Eubacterium saburreum		100.0	0.38	CEL 1		
53055110	**Good identification**					**53055110**
Eubacterium saburreum		100.0	0.49	CEL 1		
53055112	**Good identification**					**53055112**
Eubacterium saburreum		100.0	0.37	CEL 1		
53055120	**Good identification**					**53055120**
Eubacterium saburreum		100.0	0.38	CEL 1		
53055130	**Good identification**					**53055130**
Eubacterium saburreum		100.0	0.37	CEL 1		
53061000	**Good identification**					**53061000**
Eubacterium saburreum		100.0	0.38	CEL 1		
53061010	**Good identification**					**53061010**
Eubacterium saburreum		100.0	0.37	CEL 1		
53061100	**Good identification**					**53061100**
Eubacterium saburreum		100.0	0.44	CEL 1		
53061110	**Good identification**					**53061110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.43	CEL 1		
53065100	**Good identification**					**53065100**
Eubacterium saburreum		100.0	0.40	CEL 1		
53065110	**Good identification**					**53065110**
Eubacterium saburreum		100.0	0.38	CEL 1		
53071000	**Acceptable identification**					**53071000**
Eubacterium saburreum		93.88	0.46	CEL 1		
Fusobacterium mortiferum		6.12	0.28			
53071010	**Good identification**					**53071010**
Eubacterium saburreum		100.0	0.45	CEL 1		
53071100	**Acceptable identification**					**53071100**
Eubacterium saburreum		93.88	0.53	CEL 1		
Fusobacterium mortiferum		6.12	0.34			
53071102	**Good identification**					**53071102**
Eubacterium saburreum		96.06	0.40	CEL 1		
Fusobacterium mortiferum		3.95	0.18			
53071110	**Good identification**					**53071110**
Eubacterium saburreum		100.0	0.51	CEL 1		
53071112	**Good identification**					**53071112**
Eubacterium saburreum		100.0	0.38	CEL 1		
53071120	**Good identification**					**53071120**
Eubacterium saburreum		100.0	0.40	CEL 1		
53071130	**Good identification**					**53071130**
Eubacterium saburreum		100.0	0.38	CEL 1		
53075000	**Good identification**					**53075000**
Eubacterium saburreum		100.0	0.42	CEL 1		
53075010	**Good identification**					**53075010**
Eubacterium saburreum		100.0	0.40	CEL 1		
53075100	**Good identification**					**53075100**
Eubacterium saburreum		99.10	0.48	CEL 1		
Fusobacterium mortiferum		0.90	0.15			
53075102	**Good identification**					**53075102**
Eubacterium saburreum		100.0	0.35	CEL 1		
53075110	**Good identification**					**53075110**
Eubacterium saburreum		100.0	0.46	CEL 1		
53075120	**Good identification**					**53075120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.35	CEL 1		
53100000	**Low discrimination**					**53100000**
					SPO	
Clostridium bifermentans		89.08	0.64	SUC 1	99	
Clostridium botulinum biovar A		6.21	0.45		99	
Eubacterium saburreum		3.09	0.51		1	
53100001	**Acceptable identification**					**53100001**
Clostridium sordellii		90.86	0.44	ESL 5		
Clostridium bifermentans		7.88	0.30	SUC 1		
Clostridium botulinum biovar A		0.55	0.11			
53100010	**Low discrimination**					**53100010**
					SPO	
Eubacterium saburreum		71.38	0.50	GAL 99	1	
Clostridium bifermentans		25.42	0.30		99	
Clostridium botulinum biovar A		1.77	0.11		99	
53100012	**Good identification**					**53100012**
Eubacterium saburreum		100.0	0.37	GAL 99		
53100020	**Low discrimination**					**53100020**
					SPO	
Clostridium bifermentans		82.60	0.44	SUC 1	99	
Clostridium cadaveris		8.36	0.25	NAG 6	99	
Eubacterium saburreum		7.92	0.39		1	
53100030	**Low discrimination**					**53100030**
					SPO	
Eubacterium saburreum		88.59	0.37	GAL 99	1	
Clostridium bifermentans		11.41	0.10		99	
53100040	**Low discrimination**					**53100040**
					SPO	
Clostridium bifermentans		65.61	0.41	SUC 1	99	
Clostridium botulinum biovar A		32.80	0.36	SOR 4	99	
Eubacterium saburreum		0.55	0.17		1	
53100100	**Low discrimination**					**53100100**
					SPO	
Eubacterium saburreum		87.95	0.58	GAL 99	1	
Clostridium bifermentans		10.98	0.30		99	
Clostridium botulinum biovar A		0.77	0.11		99	
53100102	**Good identification**					**53100102**
Eubacterium saburreum		100.0	0.45	GAL 99		
53100110	**Very good identification**					**53100110**
Eubacterium saburreum		100.0	0.56	GAL 99		
53100112	**Good identification**					**53100112**
Eubacterium saburreum		100.0	0.43	GAL 99		
53100120	**Good identification**					**53100120**
					SPO	
Eubacterium saburreum		95.68	0.45	GAL 99	1	
Clostridium bifermentans		4.32	0.10		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53100130	**Good identification**					**53100130**
Eubacterium saburreum		100.0	0.43	GAL 99		
53101000	**Very good identification**					**53101000**
					SPO	
Eubacterium saburreum		99.66	0.86		1	
Clostridium bifermentans		0.29	0.30		99	
Clostridium botulinum biovar A		0.02	0.11		99	
53101001	**Low discrimination**					**53101001**
					SPO	
Eubacterium saburreum		62.45	0.51	URE 1	1	
Clostridium sordellii		37.03	0.31		99	
Clostridium chauvoei		0.52	0.06		99	
53101002	**Very good identification**					**53101002**
Eubacterium saburreum		100.0	0.73			
53101003	**Good identification**					**53101003**
					SPO	
Eubacterium saburreum		96.72	0.39	URE 1	1	
Clostridium sordellii		3.28	-.03		99	
53101010	**Very good identification**					**53101010**
Eubacterium saburreum		100.0	0.84			
53101011	**Good identification**					**53101011**
					SPO	
Eubacterium saburreum		99.27	0.50	URE 1	1	
Clostridium sordellii		0.73	-.03		99	
53101012	**Very good identification**					**53101012**
Eubacterium saburreum		100.0	0.71			
53101013	**Good identification**					**53101013**
Eubacterium saburreum		100.0	0.37	URE 1		
53101020	**Very good identification**					**53101020**
					SPO	
Eubacterium saburreum		99.89	0.73		1	
Clostridium bifermentans		0.11	0.10		99	
53101021	**Good identification**					**53101021**
					SPO	
Eubacterium saburreum		96.72	0.39	URE 1	1	
Clostridium sordellii		3.28	-.03		99	
53101022	**Very good identification**					**53101022**
Eubacterium saburreum		100.0	0.60			
53101030	**Very good identification**					**53101030**
Eubacterium saburreum		100.0	0.71			
53101031	**Good identification**					**53101031**
Eubacterium saburreum		100.0	0.37	URE 1		
53101032	**Very good identification**					**53101032**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.58			
53101040	**Good identification**					**53101040**
					SPO	
Eubacterium saburreum		98.21	0.51	SOR 1	1	
Clostridium bifermentans		1.19	0.07		99	
Clostridium botulinum biovar A		0.60	0.02		99	
53101042	**Good identification**					**53101042**
Eubacterium saburreum		100.0	0.39	SOR 1		
53101050	**Good identification**					**53101050**
Eubacterium saburreum		100.0	0.50	SOR 1		
53101052	**Good identification**					**53101052**
Eubacterium saburreum		100.0	0.37	SOR 1		
53101060	**Good identification**					**53101060**
Eubacterium saburreum		100.0	0.39	SOR 1		
53101070	**Good identification**					**53101070**
Eubacterium saburreum		100.0	0.37	SOR 1		
53101100	**Excellent identification**					**53101100**
					SPO	
Eubacterium saburreum		99.99	0.92		1	
Clostridium chauvoei		0.00	0.06		99	
Actinomyces meyeri		0.00	0.05		1	
53101101	**Low discrimination**					**53101101**
					SPO	
Eubacterium saburreum		73.65	0.58	URE 1	1	
Clostridium chauvoei		26.17	0.40		99	
Clostridium sordellii		0.19	-.03		99	
53101102	**Very good identification**					**53101102**
Eubacterium saburreum		100.0	0.79			
53101103	**Low discrimination**					**53101103**
					SPO	
Eubacterium saburreum		73.79	0.45	URE 1	1	
Clostridium chauvoei		26.21	0.27		99	
53101110	**Very good identification**					**53101110**
Eubacterium saburreum		100.0	0.90			
53101111	**Very good identification**					**53101111**
					SPO	
Eubacterium saburreum		99.56	0.56	URE 1	1	
Clostridium chauvoei		0.44	0.06		99	
53101112	**Very good identification**					**53101112**
Eubacterium saburreum		100.0	0.77			
53101113	**Good identification**					**53101113**
Eubacterium saburreum		100.0	0.43	URE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53101120	**Very good identification**					**53101120**
Eubacterium saburreum		99.87	0.79			
Prevotella melaninogenica		0.13	0.14			
53101121	**Low discrimination**					**53101121**
					SPO	
Eubacterium saburreum		59.84	0.45	URE 1	1	
Clostridium chauvoei		40.16	0.32		99	
53101122	**Very good identification**					**53101122**
Eubacterium saburreum		99.86	0.66			
Prevotella melaninogenica		0.14	0.02			
53101130	**Very good identification**					**53101130**
Eubacterium saburreum		100.0	0.77			
53101131	**Good identification**					**53101131**
Eubacterium saburreum		100.0	0.43	URE 1		
53101132	**Very good identification**					**53101132**
Eubacterium saburreum		100.0	0.65			
53101140	**Very good identification**					**53101140**
Eubacterium saburreum		100.0	0.58	SOR 1		
53101142	**Good identification**					**53101142**
Eubacterium saburreum		100.0	0.45	SOR 1		
53101150	**Very good identification**					**53101150**
Eubacterium saburreum		100.0	0.56	SOR 1		
53101152	**Good identification**					**53101152**
Eubacterium saburreum		100.0	0.43	SOR 1		
53101160	**Good identification**					**53101160**
Eubacterium saburreum		100.0	0.45	SOR 1		
53101170	**Good identification**					**53101170**
Eubacterium saburreum		100.0	0.43	SOR 1		
53101200	**Good identification**					**53101200**
Eubacterium saburreum		100.0	0.51	RHA 1		
53101202	**Good identification**					**53101202**
Eubacterium saburreum		100.0	0.39	RHA 1		
53101210	**Good identification**					**53101210**
Eubacterium saburreum		100.0	0.50	RHA 1		
53101212	**Good identification**					**53101212**
Eubacterium saburreum		100.0	0.37	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53101220	**Good identification**					**53101220**
Eubacterium saburreum		100.0	0.39	RHA 1		
53101230	**Good identification**					**53101230**
Eubacterium saburreum		100.0	0.37	RHA 1		
53101300	**Very good identification**					**53101300**
Eubacterium saburreum		100.0	0.58	RHA 1		
53101302	**Good identification**					**53101302**
Eubacterium saburreum		100.0	0.45	RHA 1		
53101310	**Very good identification**					**53101310**
Eubacterium saburreum		100.0	0.56	RHA 1		
53101312	**Good identification**					**53101312**
Eubacterium saburreum		100.0	0.43	RHA 1		
53101320	**Good identification**					**53101320**
Eubacterium saburreum		100.0	0.45	RHA 1		
53101330	**Good identification**					**53101330**
Eubacterium saburreum		100.0	0.43	RHA 1		
53101400	**Good identification**					**53101400**
Eubacterium saburreum		100.0	0.51	ARA 1		
53101402	**Good identification**					**53101402**
Eubacterium saburreum		100.0	0.39	ARA 1		
53101410	**Good identification**					**53101410**
Eubacterium saburreum		100.0	0.50	ARA 1		
53101412	**Good identification**					**53101412**
Eubacterium saburreum		100.0	0.37	ARA 1		
53101420	**Good identification**					**53101420**
Eubacterium saburreum		100.0	0.39	ARA 1		
53101430	**Good identification**					**53101430**
Eubacterium saburreum		100.0	0.37	ARA 1		
53101500	**Very good identification**					**53101500**
Eubacterium saburreum		100.0	0.58	ARA 1		
53101502	**Good identification**					**53101502**
Eubacterium saburreum		100.0	0.45	ARA 1		
53101510	**Very good identification**					**53101510**
Eubacterium saburreum		100.0	0.56	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53101512	**Good identification**					**53101512**
Eubacterium saburreum		100.0	0.43	ARA 1		
53101520	**Good identification**					**53101520**
Eubacterium saburreum		98.81	0.45	ARA 1		
Odoribacter splanchnicus		1.19	-.07			
53101530	**Good identification**					**53101530**
Eubacterium saburreum		100.0	0.43	ARA 1		
53103000	**Good identification**					**53103000**
Eubacterium saburreum		100.0	0.51	MAN 1		
53103002	**Good identification**					**53103002**
Eubacterium saburreum		100.0	0.39	MAN 1		
53103010	**Good identification**					**53103010**
Eubacterium saburreum		100.0	0.50	MAN 1		
53103012	**Good identification**					**53103012**
Eubacterium saburreum		100.0	0.37	MAN 1		
53103020	**Good identification**					**53103020**
Eubacterium saburreum		100.0	0.39	MAN 1		
53103030	**Good identification**					**53103030**
Eubacterium saburreum		100.0	0.37	MAN 1		
53103100	**Very good identification**					**53103100**
Eubacterium saburreum		100.0	0.58	MAN 1		
53103102	**Good identification**					**53103102**
Eubacterium saburreum		100.0	0.45	MAN 1		
53103110	**Very good identification**					**53103110**
Eubacterium saburreum		100.0	0.56	MAN 1		
53103112	**Good identification**					**53103112**
Eubacterium saburreum		100.0	0.43	MAN 1		
53103120	**Good identification**					**53103120**
Eubacterium saburreum		100.0	0.45	MAN 1		
53103130	**Good identification**					**53103130**
Eubacterium saburreum		100.0	0.43	MAN 1		
53104000	**Low discrimination**					**53104000**
					SPO	
Eubacterium saburreum		63.00	0.47	GAL 99	1	
Clostridium bifermentans		34.09	0.30		99	
Clostridium botulinum biovar A		2.38	0.11		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53104010	**Good identification**					**53104010**
Eubacterium saburreum		100.0	0.45	GAL 99		
53104100	**Good identification**					**53104100**
Eubacterium saburreum		99.45	0.53	GAL 99		
Actinomyces meyeri		0.55	0.07			
53104102	**Good identification**					**53104102**
Eubacterium saburreum		100.0	0.40	GAL 99		
53104110	**Good identification**					**53104110**
Eubacterium saburreum		100.0	0.52	GAL 99		
53104112	**Good identification**					**53104112**
Eubacterium saburreum		100.0	0.39	GAL 99		
53104120	**Good identification**					**53104120**
Eubacterium saburreum		100.0	0.40	GAL 99		
53104130	**Good identification**					**53104130**
Eubacterium saburreum		100.0	0.39	GAL 99		
53105000	**Very good identification**					**53105000**
Eubacterium saburreum		100.0	0.81			
53105001	**Good identification**					**53105001**
					SPO	
Eubacterium saburreum		98.90	0.47	URE 1	1	
Clostridium sordellii		1.10	-.03		99	
53105002	**Very good identification**					**53105002**
Eubacterium saburreum		100.0	0.68			
53105010	**Very good identification**					**53105010**
Eubacterium saburreum		100.0	0.79			
53105011	**Good identification**					**53105011**
Eubacterium saburreum		100.0	0.45	URE 1		
53105012	**Very good identification**					**53105012**
Eubacterium saburreum		100.0	0.67			
53105020	**Very good identification**					**53105020**
Eubacterium saburreum		100.0	0.68			
53105022	**Good identification**					**53105022**
Eubacterium saburreum		100.0	0.55			
53105030	**Very good identification**					**53105030**
Eubacterium saburreum		100.0	0.67			
53105032	**Good identification**					**53105032**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.54			
53105040	**Good identification**					**53105040**
					SPO	
Eubacterium saburreum		99.13	0.47	SOR 1	1	
Clostridium glycolicum		0.87	-.01		99	
53105050	**Good identification**					**53105050**
Eubacterium saburreum		100.0	0.45	SOR 1		
53105100	**Very good identification**					**53105100**
Eubacterium saburreum		99.99	0.87			
Actinomyces meyeri		0.01	0.10			
53105101	**Good identification**					**53105101**
					SPO	
Eubacterium saburreum		99.34	0.53	URE 1	1	
Clostridium chauvoei		0.66	0.06		99	
53105102	**Very good identification**					**53105102**
Eubacterium saburreum		100.0	0.74			
53105103	**Good identification**					**53105103**
Eubacterium saburreum		100.0	0.40	URE 1		
53105110	**Very good identification**					**53105110**
Eubacterium saburreum		100.0	0.86			
53105111	**Good identification**					**53105111**
Eubacterium saburreum		100.0	0.52	URE 1		
53105112	**Very good identification**					**53105112**
Eubacterium saburreum		100.0	0.73			
53105113	**Good identification**					**53105113**
Eubacterium saburreum		100.0	0.39	URE 1		
53105120	**Very good identification**					**53105120**
Eubacterium saburreum		100.0	0.74			
53105121	**Good identification**					**53105121**
Eubacterium saburreum		100.0	0.40	URE 1		
53105122	**Very good identification**					**53105122**
Eubacterium saburreum		100.0	0.61			
53105130	**Very good identification**					**53105130**
Eubacterium saburreum		100.0	0.73			
53105131	**Good identification**					**53105131**
Eubacterium saburreum		100.0	0.39	URE 1		
53105132	**Very good identification**					**53105132**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.60			
53105140	**Good identification**					**53105140**
Eubacterium saburreum		100.0	0.53	SOR 1		
53105142	**Good identification**					**53105142**
Eubacterium saburreum		100.0	0.40	SOR 1		
53105150	**Good identification**					**53105150**
Eubacterium saburreum		100.0	0.52	SOR 1		
53105152	**Good identification**					**53105152**
Eubacterium saburreum		100.0	0.39	SOR 1		
53105160	**Good identification**					**53105160**
Eubacterium saburreum		100.0	0.40	SOR 1		
53105170	**Good identification**					**53105170**
Eubacterium saburreum		100.0	0.39	SOR 1		
53105200	**Good identification**					**53105200**
Eubacterium saburreum		100.0	0.47	RHA 1		
53105210	**Good identification**					**53105210**
Eubacterium saburreum		100.0	0.45	RHA 1		
53105300	**Good identification**					**53105300**
Eubacterium saburreum		100.0	0.53	RHA 1		
53105302	**Good identification**					**53105302**
Eubacterium saburreum		100.0	0.40	RHA 1		
53105310	**Good identification**					**53105310**
Eubacterium saburreum		100.0	0.52	RHA 1		
53105312	**Good identification**					**53105312**
Eubacterium saburreum		100.0	0.39	RHA 1		
53105320	**Good identification**					**53105320**
Eubacterium saburreum		100.0	0.40	RHA 1		
53105330	**Good identification**					**53105330**
Eubacterium saburreum		100.0	0.39	RHA 1		
53105400	**Good identification**					**53105400**
Eubacterium saburreum		100.0	0.47	ARA 1		
53105410	**Good identification**					**53105410**
Eubacterium saburreum		100.0	0.45	ARA 1		
53105500	**Good identification**					**53105500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.53	ARA 1			
53105502	**Good identification**						**53105502**
Eubacterium saburreum		100.0	0.40	ARA 1			
53105510	**Good identification**						**53105510**
Eubacterium saburreum		100.0	0.52	ARA 1			
53105512	**Good identification**						**53105512**
Eubacterium saburreum		100.0	0.39	ARA 1			
53105520	**Good identification**						**53105520**
Eubacterium saburreum		98.43	0.40	ARA 1			
Bifidobacterium longum subsp. longum		1.57	0.06				
53105530	**Good identification**						**53105530**
Eubacterium saburreum		96.78	0.39	ARA 1			
Bifidobacterium longum subsp. longum		3.22	0.10				
53107000	**Good identification**						**53107000**
Eubacterium saburreum		100.0	0.47	MAN 1			
53107010	**Good identification**						**53107010**
Eubacterium saburreum		100.0	0.45	MAN 1			
53107100	**Good identification**						**53107100**
Eubacterium saburreum		100.0	0.53	MAN 1			
53107102	**Good identification**						**53107102**
Eubacterium saburreum		100.0	0.40	MAN 1			
53107110	**Good identification**						**53107110**
Eubacterium saburreum		100.0	0.52	MAN 1			
53107112	**Good identification**						**53107112**
Eubacterium saburreum		100.0	0.39	MAN 1			
53107120	**Good identification**						**53107120**
Eubacterium saburreum		100.0	0.40	MAN 1			
53107130	**Good identification**						**53107130**
Eubacterium saburreum		100.0	0.39	MAN 1			
53110000	**Low discrimination**						**53110000**
					SPO	GRA	
Eubacterium saburreum		46.58	0.60	GAL 99	1	85	
Clostridium bifermentans		23.57	0.43		99	99	
Prevotella intermedia		22.54	0.45		1	1	
53110001	**Good identification**						**53110001**
					SPO		
Clostridium sordellii		98.24	0.44	ESL 5	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		0.89	0.25	SUC 1	1		
Clostridium bifermentans		0.45	0.08		99		
53110002	**Low discrimination**						**53110002**
					COC	SPO	
Eubacterium saburreum		87.29	0.47	GAL 99	1	1	
Clostridium sporogenes		4.37	0.11		1	99	
Megasphaera elsdenii		3.40	0.15		99	1	
53110010	**Very good identification**						**53110010**
					SPO	GRA	
Eubacterium saburreum		98.79	0.58	GAL 99	1	85	
Clostridium bifermentans		0.62	0.08		99	99	
Prevotella intermedia		0.59	0.11		1	1	
53110012	**Good identification**						**53110012**
Eubacterium saburreum		100.0	0.45	GAL 99			
53110020	**Low discrimination**						**53110020**
					SPO		
Eubacterium saburreum		63.99	0.47	GAL 99	1		
Clostridium cadaveris		22.53	0.25		99		
Clostridium bifermentans		11.71	0.22		99		
53110030	**Good identification**						**53110030**
Eubacterium saburreum		100.0	0.45	GAL 99			
53110100	**Very good identification**						**53110100**
					GRA		
Eubacterium saburreum		97.76	0.66	GAL 99	85		
Actinomyces meyeri		0.94	0.24		99		
Fusobacterium mortiferum		0.88	0.33		1		
53110102	**Good identification**						**53110102**
Eubacterium saburreum		99.44	0.53	GAL 99			
Fusobacterium mortiferum		0.56	0.16				
53110110	**Very good identification**						**53110110**
Eubacterium saburreum		100.0	0.64	GAL 99			
53110112	**Good identification**						**53110112**
Eubacterium saburreum		100.0	0.52	GAL 99			
53110120	**Good identification**						**53110120**
					GRA		
Eubacterium saburreum		98.87	0.53	GAL 99	85		
Actinomyces meyeri		0.74	0.09		99		
Prevotella melaninogenica		0.40	-.04		1		
53110122	**Good identification**						**53110122**
Eubacterium saburreum		100.0	0.40	GAL 99			
53110130	**Good identification**						**53110130**
Eubacterium saburreum		100.0	0.52	GAL 99			
53110132	**Good identification**						**53110132**
Eubacterium saburreum		100.0	0.39	GAL 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53111000	**Excellent identification**					**53111000**
Eubacterium saburreum		99.48	0.94			
Prevotella intermedia		0.49	0.45			
Fusobacterium mortiferum		0.01	0.29			
53111001	**Low discrimination**					**53111001**
					SPO	
Eubacterium saburreum		82.71	0.60	URE 1	1	
Clostridium sordellii		16.35	0.31		99	
Clostridium chauvoei		0.54	0.12		99	
53111002	**Very good identification**					**53111002**
					GRA	
Eubacterium saburreum		99.93	0.81		85	
Lactobacillus fermentum		0.03	0.24		99	
Prevotella intermedia		0.03	0.11		1	
53111003	**Good identification**					**53111003**
					SPO	
Eubacterium saburreum		98.25	0.47	URE 1	1	
Clostridium sordellii		1.11	-.03		99	
Clostridium chauvoei		0.64	-.01		99	
53111010	**Excellent identification**					**53111010**
Eubacterium saburreum		99.99	0.92			
Prevotella intermedia		0.01	0.11			
53111011	**Very good identification**					**53111011**
					SPO	
Eubacterium saburreum		99.76	0.58	URE 1	1	
Clostridium sordellii		0.24	-.03		99	
53111012	**Very good identification**					**53111012**
Eubacterium saburreum		100.0	0.79			
53111013	**Good identification**					**53111013**
Eubacterium saburreum		100.0	0.45	URE 1		
53111020	**Very good identification**					**53111020**
					GRA	
Eubacterium saburreum		99.94	0.81		85	
Prevotella intermedia		0.03	0.11		1	
Lactobacillus fermentum		0.01	0.16		99	
53111021	**Good identification**					**53111021**
					SPO	
Eubacterium saburreum		97.69	0.47	URE 1	1	
Clostridium chauvoei		1.20	0.04		99	
Clostridium sordellii		1.11	-.03		99	
53111022	**Very good identification**					**53111022**
Eubacterium saburreum		99.94	0.68			
Lactobacillus fermentum		0.06	0.16			
53111030	**Very good identification**					**53111030**
Eubacterium saburreum		100.0	0.79			
53111031	**Good identification**					**53111031**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.45	URE 1			
53111032	**Very good identification**						**53111032**
Eubacterium saburreum		100.0	0.66				
53111040	**Very good identification**						**53111040**
Eubacterium saburreum		99.51	0.60	SOR 1			
Prevotella intermedia		0.49	0.11				
53111042	**Good identification**						**53111042**
Eubacterium saburreum		100.0	0.47	SOR 1			
53111050	**Very good identification**						**53111050**
Eubacterium saburreum		100.0	0.58	SOR 1			
53111052	**Good identification**						**53111052**
Eubacterium saburreum		100.0	0.45	SOR 1			
53111060	**Good identification**						**53111060**
Eubacterium saburreum		100.0	0.47	SOR 1			
53111070	**Good identification**						**53111070**
Eubacterium saburreum		100.0	0.45	SOR 1			
53111100	**Excellent identification**						**53111100**
					GRA		
Eubacterium saburreum		99.96	1.00		85		
Actinomyces meyeri		0.02	0.27		99		
Fusobacterium mortiferum		0.01	0.36		1		
53111101	**Low discrimination**						**53111101**
					SPO		
Eubacterium saburreum		78.30	0.66	URE 1	1		
Clostridium chauvoei		21.64	0.47		99		
Clostridium sordellii		0.07	-.03		99		
53111102	**Very good identification**						**53111102**
					CAT	GRA	
Eubacterium saburreum		99.94	0.87		1	85	
Capnocytophaga ochracea		0.03	0.14		90	1	
Lactobacillus fermentum		0.02	0.28		1	99	
53111103	**Low discrimination**						**53111103**
					SPO		
Eubacterium saburreum		78.35	0.53	URE 1	1		
Clostridium chauvoei		21.65	0.34		99		
53111110	**Excellent identification**						**53111110**
Eubacterium saburreum		100.0	0.99				
53111111	**Very good identification**						**53111111**
					SPO		
Eubacterium saburreum		99.66	0.64	URE 1	1		
Clostridium chauvoei		0.34	0.12		99		
53111112	**Very good identification**						**53111112**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.86		1		
Capnocytophaga ochracea		0.00	-.03		90		
53111113	**Good identification**						**53111113**
					SPO		
Eubacterium saburreum		99.66	0.52	URE 1	1		
Clostridium chauvoei		0.34	-.01		99		
53111120	**Very good identification**						**53111120**
					GRA		
Eubacterium saburreum		99.52	0.87		85		
Prevotella melaninogenica		0.40	0.30		1		
Clostridium perfringens		0.03	0.17		99		
53111121	**Low discrimination**						**53111121**
					SPO	GRA	
Eubacterium saburreum		65.53	0.53	URE 1	1	85	
Clostridium chauvoei		34.20	0.38		99	99	
Prevotella melaninogenica		0.26	-.04		1	1	
53111122	**Very good identification**						**53111122**
					CAT		
Eubacterium saburreum		98.10	0.74		1		
Capnocytophaga ochracea		1.34	0.30		90		
Prevotella melaninogenica		0.43	0.18		1		
53111123	**Low discrimination**						**53111123**
					SPO		
Eubacterium saburreum		65.71	0.40	URE 1	1		
Clostridium chauvoei		34.30	0.25		99		
53111130	**Very good identification**						**53111130**
					CAT		
Eubacterium saburreum		99.99	0.86		1		
Prevotella melaninogenica		0.01	-.04		1		
Capnocytophaga ochracea		0.00	-.03		90		
53111131	**Good identification**						**53111131**
					SPO		
Eubacterium saburreum		99.36	0.52	URE 1	1		
Clostridium chauvoei		0.64	0.04		99		
53111132	**Very good identification**						**53111132**
					CAT		
Eubacterium saburreum		99.82	0.73		1		
Capnocytophaga ochracea		0.19	0.14		90		
53111133	**Good identification**						**53111133**
Eubacterium saburreum		100.0	0.39	URE 1			
53111140	**Very good identification**						**53111140**
Eubacterium saburreum		100.0	0.66	SOR 1			
53111142	**Good identification**						**53111142**
					CAT		
Eubacterium saburreum		99.71	0.53	SOR 1	1		
Capnocytophaga ochracea		0.29	-.03		90		
53111150	**Very good identification**						**53111150**
Eubacterium saburreum		100.0	0.64	SOR 1			
53111152	**Good identification**						**53111152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.52	SOR 1		
53111160	**Good identification**					**53111160**
					GRA	
Eubacterium saburreum		98.82	0.53	SOR 1	85	
Clostridium perfringens		0.49	0.04		99	
Prevotella melaninogenica		0.40	-.04		1	
53111162	**Low discrimination**					**53111162**
					CAT	
Eubacterium saburreum		86.92	0.40	SOR 1	1	
Capnocytophaga ochracea		13.08	0.14		90	
53111170	**Good identification**					**53111170**
Eubacterium saburreum		100.0	0.52	SOR 1		
53111172	**Good identification**					**53111172**
					CAT	
Eubacterium saburreum		98.00	0.39	SOR 1	1	
Capnocytophaga ochracea		2.00	-.03		90	
53111200	**Very good identification**					**53111200**
Eubacterium saburreum		99.51	0.60	RHA 1		
Prevotella intermedia		0.49	0.11			
53111202	**Good identification**					**53111202**
Eubacterium saburreum		100.0	0.47	RHA 1		
53111210	**Very good identification**					**53111210**
Eubacterium saburreum		100.0	0.58	RHA 1		
53111212	**Good identification**					**53111212**
Eubacterium saburreum		100.0	0.45	RHA 1		
53111220	**Good identification**					**53111220**
Eubacterium saburreum		100.0	0.47	RHA 1		
53111230	**Good identification**					**53111230**
Eubacterium saburreum		100.0	0.45	RHA 1		
53111300	**Very good identification**					**53111300**
Eubacterium saburreum		100.0	0.66	RHA 1		
53111302	**Good identification**					**53111302**
Eubacterium saburreum		100.0	0.53	RHA 1		
53111310	**Very good identification**					**53111310**
Eubacterium saburreum		100.0	0.64	RHA 1		
53111312	**Good identification**					**53111312**
Eubacterium saburreum		100.0	0.52	RHA 1		
53111320	**Good identification**					**53111320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		99.60	0.53	RHA 1			
Prevotella melaninogenica		0.40	-.04				
53111322	**Good identification**						**53111322**
Eubacterium saburreum		100.0	0.40	RHA 1			
53111330	**Good identification**						**53111330**
Eubacterium saburreum		100.0	0.52	RHA 1			
53111332	**Good identification**						**53111332**
Eubacterium saburreum		100.0	0.39	RHA 1			
53111400	**Very good identification**						**53111400**
					GRA		
Eubacterium saburreum		98.74	0.60	ARA 1	85		
Lactobacillus fermentum		0.57	0.24		99		
Prevotella intermedia		0.48	0.11		1		
53111402	**Good identification**						**53111402**
Eubacterium saburreum		96.84	0.47	ARA 1			
Lactobacillus fermentum		3.16	0.24				
53111410	**Very good identification**						**53111410**
Eubacterium saburreum		100.0	0.58	ARA 1			
53111412	**Good identification**						**53111412**
Eubacterium saburreum		100.0	0.45	ARA 1			
53111420	**Good identification**						**53111420**
Eubacterium saburreum		98.93	0.47	ARA 1			
Lactobacillus fermentum		1.08	0.16				
53111430	**Good identification**						**53111430**
Eubacterium saburreum		100.0	0.45	ARA 1			
53111500	**Very good identification**						**53111500**
Eubacterium saburreum		99.24	0.66	ARA 1			
Lactobacillus fermentum		0.40	0.28				
Actinomyces meyeri		0.36	0.16				
53111502	**Good identification**						**53111502**
					CAT	GRA	
Eubacterium saburreum		97.33	0.53	ARA 1	1	85	
Lactobacillus fermentum		2.22	0.28		1	99	
Capnocytophaga ochracea		0.46	0.01		90	1	
53111510	**Very good identification**						**53111510**
Eubacterium saburreum		99.92	0.64	ARA 1			
Bifidobacterium longum subsp. longum		0.08	0.07				
53111512	**Good identification**						**53111512**
Eubacterium saburreum		99.55	0.52	ARA 1			
Bifidobacterium longum subsp. longum		0.45	0.07				
53111520	**Good identification**						**53111520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		96.60	0.53	ARA 1			
Bifidobacterium longum subsp. longum		0.83	0.14				
Lactobacillus fermentum		0.73	0.20				
53111522	**Low discrimination**						**53111522**
					CAT	GRA	
Eubacterium saburreum		75.14	0.40	ARA 1	1	85	
Capnocytophaga ochracea		17.96	0.17		90	1	
Bifidobacterium longum subsp. longum		3.67	0.14		1	99	
53111530	**Good identification**						**53111530**
Eubacterium saburreum		98.24	0.52	ARA 1			
Bifidobacterium longum subsp. longum		1.76	0.18				
53111532	**Low discrimination**						**53111532**
					CAT	GRA	
Eubacterium saburreum		88.18	0.39	ARA 1	1	85	
Bifidobacterium longum subsp. longum		8.95	0.18		1	99	
Capnocytophaga ochracea		2.86	0.01		90	1	
53113000	**Very good identification**						**53113000**
Eubacterium saburreum		99.51	0.60	MAN 1			
Prevotella intermedia		0.49	0.11				
53113002	**Good identification**						**53113002**
Eubacterium saburreum		100.0	0.47	MAN 1			
53113010	**Very good identification**						**53113010**
Eubacterium saburreum		100.0	0.58	MAN 1			
53113012	**Good identification**						**53113012**
Eubacterium saburreum		100.0	0.45	MAN 1			
53113020	**Good identification**						**53113020**
Eubacterium saburreum		100.0	0.47	MAN 1			
53113030	**Good identification**						**53113030**
Eubacterium saburreum		100.0	0.45	MAN 1			
53113100	**Very good identification**						**53113100**
Eubacterium saburreum		99.93	0.66	MAN 1			
Propionibacterium propionicum		0.07	0.15				
53113102	**Good identification**						**53113102**
Eubacterium saburreum		99.38	0.53	MAN 1			
Propionibacterium propionicum		0.62	0.18				
53113110	**Very good identification**						**53113110**
Eubacterium saburreum		100.0	0.64	MAN 1			
53113112	**Good identification**						**53113112**
Eubacterium saburreum		100.0	0.52	MAN 1			
53113120	**Good identification**						**53113120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.60	0.53	MAN 1		
Prevotella melaninogenica		0.40	-.04			
53113122	**Good identification**					**53113122**
Eubacterium saburreum		100.0	0.40	MAN 1		
53113130	**Good identification**					**53113130**
Eubacterium saburreum		100.0	0.52	MAN 1		
53113132	**Good identification**					**53113132**
Eubacterium saburreum		100.0	0.39	MAN 1		
53114000	**Good identification**					**53114000**
					SPO	
Eubacterium saburreum		95.12	0.55	GAL 99	1	
Actinomyces meyeri		1.93	0.18		1	
Clostridium glycolicum		1.18	0.10		99	
53114002	**Good identification**					**53114002**
Eubacterium saburreum		100.0	0.42	GAL 99		
53114010	**Good identification**					**53114010**
Eubacterium saburreum		100.0	0.53	GAL 99		
53114012	**Good identification**					**53114012**
Eubacterium saburreum		100.0	0.41	GAL 99		
53114020	**Good identification**					**53114020**
Eubacterium saburreum		98.46	0.42	GAL 99		
Actinomyces meyeri		1.55	0.04			
53114030	**Good identification**					**53114030**
Eubacterium saburreum		100.0	0.41	GAL 99		
53114100	**Very good identification**					**53114100**
					GRA	
Eubacterium saburreum		96.52	0.61	GAL 99	85	
Actinomyces meyeri		3.36	0.29		99	
Fusobacterium mortiferum		0.12	0.13		1	
53114102	**Good identification**					**53114102**
Eubacterium saburreum		99.39	0.48	GAL 99		
Actinomyces meyeri		0.61	0.03			
53114110	**Very good identification**					**53114110**
Eubacterium saburreum		100.0	0.60	GAL 99		
53114112	**Good identification**					**53114112**
Eubacterium saburreum		100.0	0.47	GAL 99		
53114120	**Good identification**					**53114120**
Eubacterium saburreum		97.38	0.48	GAL 99		
Actinomyces meyeri		2.62	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53114122	**Good identification**					**53114122**
Eubacterium saburreum		100.0	0.35	GAL 99		
53114130	**Good identification**					**53114130**
Eubacterium saburreum		100.0	0.47	GAL 99		
53115000	**Very good identification**					**53115000**
Eubacterium saburreum		99.95	0.89			
Actinomyces meyeri		0.03	0.21			
Lactobacillus fermentum		0.01	0.20			
53115001	**Good identification**					**53115001**
					SPO	
Eubacterium saburreum		99.63	0.55	URE 1	1	
Clostridium sordellii		0.37	-.03		99	
53115002	**Very good identification**					**53115002**
Eubacterium saburreum		99.96	0.76			
Lactobacillus fermentum		0.04	0.20			
53115003	**Good identification**					**53115003**
Eubacterium saburreum		100.0	0.42	URE 1		
53115010	**Very good identification**					**53115010**
Eubacterium saburreum		100.0	0.88			
53115011	**Good identification**					**53115011**
Eubacterium saburreum		100.0	0.53	URE 1		
53115012	**Very good identification**					**53115012**
Eubacterium saburreum		100.0	0.75			
53115013	**Good identification**					**53115013**
Eubacterium saburreum		100.0	0.41	URE 1		
53115020	**Very good identification**					**53115020**
Eubacterium saburreum		99.96	0.76			
Actinomyces meyeri		0.02	0.07			
Lactobacillus fermentum		0.01	0.12			
53115021	**Good identification**					**53115021**
Eubacterium saburreum		100.0	0.42	URE 1		
53115022	**Very good identification**					**53115022**
Eubacterium saburreum		99.93	0.63			
Lactobacillus fermentum		0.07	0.12			
53115030	**Very good identification**					**53115030**
Eubacterium saburreum		100.0	0.75			
53115031	**Good identification**					**53115031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.41	URE 1		
53115032	**Very good identification**					**53115032**
Eubacterium saburreum		100.0	0.62			
53115040	**Good identification**					**53115040**
					SPO	
Eubacterium saburreum		97.44	0.55	SOR 1	1	
Clostridium glycolicum		2.56	0.15		99	
53115042	**Good identification**					**53115042**
Eubacterium saburreum		100.0	0.42	SOR 1		
53115050	**Good identification**					**53115050**
Eubacterium saburreum		100.0	0.53	SOR 1		
53115052	**Good identification**					**53115052**
Eubacterium saburreum		100.0	0.41	SOR 1		
53115060	**Good identification**					**53115060**
Eubacterium saburreum		100.0	0.42	SOR 1		
53115070	**Good identification**					**53115070**
Eubacterium saburreum		100.0	0.41	SOR 1		
53115100	**Excellent identification**					**53115100**
Eubacterium saburreum		99.94	0.95			
Actinomyces meyeri		0.05	0.32			
Lactobacillus fermentum		0.00	0.24			
53115101	**Very good identification**					**53115101**
					SPO	
Eubacterium saburreum		99.48	0.61	URE 1	1	
Clostridium chauvoei		0.52	0.12		99	
53115102	**Very good identification**					**53115102**
Eubacterium saburreum		99.95	0.83			
Lactobacillus fermentum		0.03	0.24			
Actinomyces meyeri		0.01	0.06			
53115103	**Good identification**					**53115103**
					SPO	
Eubacterium saburreum		99.48	0.48	URE 1	1	
Clostridium chauvoei		0.52	-.01		99	
53115110	**Excellent identification**					**53115110**
Eubacterium saburreum		100.0	0.94			
53115111	**Very good identification**					**53115111**
Eubacterium saburreum		100.0	0.60	URE 1		
53115112	**Very good identification**					**53115112**
Eubacterium saburreum		100.0	0.81			
53115113	**Good identification**					**53115113**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.47	URE 1		
53115120	**Very good identification**					**53115120**
Eubacterium saburreum		99.93	0.83			
Actinomyces meyeri		0.04	0.17			
Lactobacillus fermentum		0.01	0.16			
53115121	**Good identification**					**53115121**
					SPO	
Eubacterium saburreum		99.03	0.48	URE 1	1	
Clostridium chauvoei		0.97	0.04		99	
53115122	**Very good identification**					**53115122**
					CAT	
Eubacterium saburreum		99.21	0.70		1	
Capnocytophaga ochracea		0.63	0.20		90	
Bacteroides fragilis		0.11	-.03		99	
53115123	**Good identification**					**53115123**
Eubacterium saburreum		100.0	0.35	URE 1		
53115130	**Very good identification**					**53115130**
Eubacterium saburreum		100.0	0.81			
53115131	**Good identification**					**53115131**
Eubacterium saburreum		100.0	0.47	URE 1		
53115132	**Very good identification**					**53115132**
					CAT	
Eubacterium saburreum		99.91	0.68		1	
Capnocytophaga ochracea		0.09	0.04		90	
53115140	**Very good identification**					**53115140**
Eubacterium saburreum		100.0	0.61	SOR 1		
53115142	**Good identification**					**53115142**
Eubacterium saburreum		100.0	0.48	SOR 1		
53115150	**Very good identification**					**53115150**
Eubacterium saburreum		100.0	0.60	SOR 1		
53115152	**Good identification**					**53115152**
Eubacterium saburreum		100.0	0.47	SOR 1		
53115160	**Good identification**					**53115160**
Eubacterium saburreum		100.0	0.48	SOR 1		
53115162	**Acceptable identification**					**53115162**
					CAT	
Eubacterium saburreum		93.47	0.35	SOR 1	1	
Capnocytophaga ochracea		6.53	0.04		90	
53115170	**Good identification**					**53115170**
Eubacterium saburreum		100.0	0.47	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53115200	**Good identification**					**53115200**
Eubacterium saburreum		100.0	0.55	RHA 1		
53115202	**Good identification**					**53115202**
Eubacterium saburreum		100.0	0.42	RHA 1		
53115210	**Good identification**					**53115210**
Eubacterium saburreum		100.0	0.53	RHA 1		
53115212	**Good identification**					**53115212**
Eubacterium saburreum		100.0	0.41	RHA 1		
53115220	**Good identification**					**53115220**
Eubacterium saburreum		100.0	0.42	RHA 1		
53115230	**Good identification**					**53115230**
Eubacterium saburreum		100.0	0.41	RHA 1		
53115300	**Good identification**					**53115300**
Eubacterium saburreum		92.45	0.61	RHA 1		
Eubacterium contortum		7.55	0.32			
53115302	**Acceptable identification**					**53115302**
Eubacterium saburreum		92.45	0.48	RHA 1		
Eubacterium contortum		7.55	0.19			
53115310	**Very good identification**					**53115310**
Eubacterium saburreum		100.0	0.60	RHA 1		
53115312	**Good identification**					**53115312**
Eubacterium saburreum		100.0	0.47	RHA 1		
53115320	**Acceptable identification**					**53115320**
Eubacterium saburreum		92.45	0.48	RHA 1		
Eubacterium contortum		7.55	0.19			
53115322	**Acceptable identification**					**53115322**
Eubacterium saburreum		92.45	0.35	RHA 1		
Eubacterium contortum		7.55	0.06			
53115330	**Good identification**					**53115330**
Eubacterium saburreum		100.0	0.47	RHA 1		
53115400	**Good identification**					**53115400**
Eubacterium saburreum		98.63	0.55	ARA 1		
Actinomyces meyeri		0.75	0.11			
Lactobacillus fermentum		0.62	0.20			
53115402	**Good identification**					**53115402**
Eubacterium saburreum		96.57	0.42	ARA 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Lactobacillus fermentum	3.43	0.20				
53115410	**Good identification**						**53115410**
	Eubacterium saburreum	100.0	0.53	ARA 1			
53115412	**Good identification**						**53115412**
	Eubacterium saburreum	100.0	0.41	ARA 1			
53115420	**Good identification**						**53115420**
	Eubacterium saburreum	98.83	0.42	ARA 1			
	Lactobacillus fermentum	1.17	0.12				
53115430	**Good identification**						**53115430**
	Eubacterium saburreum	100.0	0.41	ARA 1			
53115500	**Very good identification**						**53115500**
	Eubacterium saburreum	98.08	0.61	ARA 1			
	Actinomyces meyeri	1.28	0.21				
	Lactobacillus fermentum	0.43	0.24				
53115502	**Good identification**						**53115502**
	Eubacterium saburreum	95.29	0.48	ARA 1			
	Lactobacillus fermentum	2.37	0.24				
	Bifidobacterium adolescentis	1.20	0.14				
53115510	**Very good identification**						**53115510**
	Eubacterium saburreum	99.56	0.60	ARA 1			
	Bifidobacterium longum subsp. longum	0.44	0.16				
53115512	**Good identification**						**53115512**
	Eubacterium saburreum	95.81	0.47	ARA 1			
	Bifidobacterium longum subsp. longum	2.39	0.16				
	Bifidobacterium adolescentis	1.80	0.15				
53115520	**Acceptable identification**						**53115520**
	Eubacterium saburreum	93.78	0.48	ARA 1			
	Bifidobacterium longum subsp. longum	4.50	0.22				
	Actinomyces meyeri	0.95	0.07				
53115522	**Low discrimination**						**53115522**
					CAT	GRA	
	Eubacterium saburreum	69.94	0.35	ARA 1	1	85	
	Bifidobacterium longum subsp. longum	19.01	0.22		1	99	
	Capnocytophaga ochracea	7.76	0.07		90	1	
53115530	**Acceptable identification**						**53115530**
	Eubacterium saburreum	90.92	0.47	ARA 1			
	Bifidobacterium longum subsp. longum	9.08	0.26				
53117000	**Good identification**						**53117000**
	Eubacterium saburreum	100.0	0.55	MAN 1			
53117002	**Good identification**						**53117002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.42	MAN 1		
53117010	**Good identification**					**53117010**
Eubacterium saburreum		100.0	0.53	MAN 1		
53117012	**Good identification**					**53117012**
Eubacterium saburreum		100.0	0.41	MAN 1		
53117020	**Good identification**					**53117020**
Eubacterium saburreum		100.0	0.42	MAN 1		
53117030	**Good identification**					**53117030**
Eubacterium saburreum		100.0	0.41	MAN 1		
53117100	**Very good identification**					**53117100**
Eubacterium saburreum		100.0	0.61	MAN 1		
53117102	**Good identification**					**53117102**
Eubacterium saburreum		100.0	0.48	MAN 1		
53117110	**Very good identification**					**53117110**
Eubacterium saburreum		100.0	0.60	MAN 1		
53117112	**Good identification**					**53117112**
Eubacterium saburreum		100.0	0.47	MAN 1		
53117120	**Good identification**					**53117120**
Eubacterium saburreum		100.0	0.48	MAN 1		
53117122	**Good identification**					**53117122**
Eubacterium saburreum		100.0	0.35	MAN 1		
53117130	**Good identification**					**53117130**
Eubacterium saburreum		100.0	0.47	MAN 1		
53120000	**Low discrimination**					**53120000**
					SPO	
Eubacterium saburreum		41.38	0.48	GAL 99	1	
Clostridium botulinum biovar A		31.21	0.35		99	
Clostridium bifermentans		18.08	0.30		99	
53120002	**Low discrimination**					**53120002**
					SPO	
Eubacterium saburreum		68.09	0.36	GAL 99	1	
Clostridium sporogenes		28.97	0.15		99	
Clostridium botulinum biovar A		2.94	0.01		99	
53120010	**Good identification**					**53120010**
					SPO	
Eubacterium saburreum		99.08	0.47	GAL 99	1	
Clostridium botulinum biovar A		0.92	0.01		99	
53120020	**Low discrimination**					**53120020**
					SPO	
Eubacterium saburreum		77.85	0.36	GAL 99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		12.30	0.10		99	
Clostridium cadaveris		6.49	0.03		99	
53120100	**Good identification**					**53120100**
					SPO	
Eubacterium saburreum		99.68	0.55	GAL 99	1	
Clostridium botulinum biovar A		0.33	0.01		99	
53120102	**Good identification**					**53120102**
Eubacterium saburreum		100.0	0.42	GAL 99		
53120110	**Good identification**					**53120110**
Eubacterium saburreum		100.0	0.53	GAL 99		
53120112	**Good identification**					**53120112**
Eubacterium saburreum		100.0	0.40	GAL 99		
53120120	**Good identification**					**53120120**
Eubacterium saburreum		100.0	0.42	GAL 99		
53120130	**Good identification**					**53120130**
Eubacterium saburreum		100.0	0.40	GAL 99		
53121000	**Very good identification**					**53121000**
					SPO	
Eubacterium saburreum		99.99	0.83		1	
Clostridium botulinum biovar A		0.01	0.01		99	
53121001	**Good identification**					**53121001**
					SPO	
Eubacterium saburreum		99.11	0.48	URE 1	1	
Clostridium sordellii		0.89	-.03		99	
53121002	**Very good identification**					**53121002**
Eubacterium saburreum		100.0	0.70			
53121003	**Good identification**					**53121003**
Eubacterium saburreum		100.0	0.36	URE 1		
53121010	**Very good identification**					**53121010**
Eubacterium saburreum		100.0	0.81			
53121011	**Good identification**					**53121011**
Eubacterium saburreum		100.0	0.47	URE 1		
53121012	**Very good identification**					**53121012**
Eubacterium saburreum		100.0	0.68			
53121020	**Very good identification**					**53121020**
Eubacterium saburreum		100.0	0.70			
53121021	**Good identification**					**53121021**
Eubacterium saburreum		100.0	0.36	URE 1		
53121022	**Very good identification**					**53121022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.57			
53121030	**Very good identification**					**53121030**
Eubacterium saburreum		100.0	0.68			
53121032	**Good identification**					**53121032**
Eubacterium saburreum		100.0	0.55			
53121040	**Good identification**					**53121040**
Eubacterium saburreum		100.0	0.48	SOR 1		
53121042	**Good identification**					**53121042**
Eubacterium saburreum		100.0	0.36	SOR 1		
53121050	**Good identification**					**53121050**
Eubacterium saburreum		100.0	0.47	SOR 1		
53121060	**Good identification**					**53121060**
Eubacterium saburreum		100.0	0.36	SOR 1		
53121100	**Very good identification**					**53121100**
Eubacterium saburreum		100.0	0.89			
53121101	**Acceptable identification**					**53121101**
					SPO	
Eubacterium saburreum		94.41	0.55	URE 1	1	
Clostridium chauvoei		5.59	0.24		99	
53121102	**Very good identification**					**53121102**
Eubacterium saburreum		100.0	0.76			
53121103	**Acceptable identification**					**53121103**
					SPO	
Eubacterium saburreum		94.41	0.42	URE 1	1	
Clostridium chauvoei		5.59	0.11		99	
53121110	**Very good identification**					**53121110**
Eubacterium saburreum		100.0	0.87			
53121111	**Good identification**					**53121111**
Eubacterium saburreum		100.0	0.53	URE 1		
53121112	**Very good identification**					**53121112**
Eubacterium saburreum		100.0	0.74			
53121113	**Good identification**					**53121113**
Eubacterium saburreum		100.0	0.40	URE 1		
53121120	**Very good identification**					**53121120**
Eubacterium saburreum		100.0	0.76			
53121121	**Low discrimination**					**53121121**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		89.94	0.42	URE 1	1	
Clostridium chauvoei		10.06	0.16		99	
53121122	**Very good identification**					**53121122**
Eubacterium saburreum		100.0	0.63			
53121130	**Very good identification**					**53121130**
Eubacterium saburreum		100.0	0.74			
53121131	**Good identification**					**53121131**
Eubacterium saburreum		100.0	0.40	URE 1		
53121132	**Very good identification**					**53121132**
Eubacterium saburreum		100.0	0.62			
53121140	**Good identification**					**53121140**
Eubacterium saburreum		100.0	0.55	SOR 1		
53121142	**Good identification**					**53121142**
Eubacterium saburreum		100.0	0.42	SOR 1		
53121150	**Good identification**					**53121150**
Eubacterium saburreum		100.0	0.53	SOR 1		
53121152	**Good identification**					**53121152**
Eubacterium saburreum		100.0	0.40	SOR 1		
53121160	**Good identification**					**53121160**
Eubacterium saburreum		100.0	0.42	SOR 1		
53121170	**Good identification**					**53121170**
Eubacterium saburreum		100.0	0.40	SOR 1		
53121200	**Good identification**					**53121200**
Eubacterium saburreum		100.0	0.48	RHA 1		
53121202	**Good identification**					**53121202**
Eubacterium saburreum		100.0	0.36	RHA 1		
53121210	**Good identification**					**53121210**
Eubacterium saburreum		100.0	0.47	RHA 1		
53121220	**Good identification**					**53121220**
Eubacterium saburreum		100.0	0.36	RHA 1		
53121300	**Good identification**					**53121300**
Eubacterium saburreum		100.0	0.55	RHA 1		
53121302	**Good identification**					**53121302**
Eubacterium saburreum		100.0	0.42	RHA 1		
53121310	**Good identification**					**53121310**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.53	RHA 1		
53121312	**Good identification**					**53121312**
Eubacterium saburreum		100.0	0.40	RHA 1		
53121320	**Good identification**					**53121320**
Eubacterium saburreum		100.0	0.42	RHA 1		
53121330	**Good identification**					**53121330**
Eubacterium saburreum		100.0	0.40	RHA 1		
53121400	**Good identification**					**53121400**
Eubacterium saburreum		100.0	0.48	ARA 1		
53121402	**Good identification**					**53121402**
Eubacterium saburreum		100.0	0.36	ARA 1		
53121410	**Good identification**					**53121410**
Eubacterium saburreum		100.0	0.47	ARA 1		
53121420	**Good identification**					**53121420**
Eubacterium saburreum		100.0	0.36	ARA 1		
53121500	**Good identification**					**53121500**
Eubacterium saburreum		100.0	0.55	ARA 1		
53121502	**Good identification**					**53121502**
Eubacterium saburreum		100.0	0.42	ARA 1		
53121510	**Good identification**					**53121510**
Eubacterium saburreum		100.0	0.53	ARA 1		
53121512	**Good identification**					**53121512**
Eubacterium saburreum		100.0	0.40	ARA 1		
53121520	**Good identification**					**53121520**
Eubacterium saburreum		100.0	0.42	ARA 1		
53121530	**Good identification**					**53121530**
Eubacterium saburreum		100.0	0.40	ARA 1		
53123000	**Good identification**					**53123000**
Eubacterium saburreum		100.0	0.48	MAN 1		
53123002	**Good identification**					**53123002**
Eubacterium saburreum		100.0	0.36	MAN 1		
53123010	**Good identification**					**53123010**
Eubacterium saburreum		100.0	0.47	MAN 1		
53123020	**Good identification**					**53123020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.36	MAN 1		
53123100	**Good identification**					**53123100**
Eubacterium saburreum		100.0	0.55	MAN 1		
53123102	**Good identification**					**53123102**
Eubacterium saburreum		100.0	0.42	MAN 1		
53123110	**Good identification**					**53123110**
Eubacterium saburreum		100.0	0.53	MAN 1		
53123112	**Good identification**					**53123112**
Eubacterium saburreum		100.0	0.40	MAN 1		
53123120	**Good identification**					**53123120**
Eubacterium saburreum		100.0	0.42	MAN 1		
53123130	**Good identification**					**53123130**
Eubacterium saburreum		100.0	0.40	MAN 1		
53124000	**Good identification**					**53124000**
					SPO	
Eubacterium saburreum		98.61	0.44	GAL 99	1	
Clostridium botulinum biovar A		1.40	0.01		99	
53124010	**Good identification**					**53124010**
Eubacterium saburreum		100.0	0.42	GAL 99		
53124100	**Good identification**					**53124100**
Eubacterium saburreum		100.0	0.50	GAL 99		
53124102	**Good identification**					**53124102**
Eubacterium saburreum		100.0	0.37	GAL 99		
53124110	**Good identification**					**53124110**
Eubacterium saburreum		100.0	0.49	GAL 99		
53124112	**Good identification**					**53124112**
Eubacterium saburreum		100.0	0.36	GAL 99		
53124120	**Good identification**					**53124120**
Eubacterium saburreum		100.0	0.37	GAL 99		
53124130	**Good identification**					**53124130**
Eubacterium saburreum		100.0	0.36	GAL 99		
53125000	**Very good identification**					**53125000**
Eubacterium saburreum		100.0	0.78			
53125001	**Good identification**					**53125001**
Eubacterium saburreum		100.0	0.44	URE 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53125002	**Very good identification**					**53125002**
Eubacterium saburreum		100.0	0.65			
53125010	**Very good identification**					**53125010**
Eubacterium saburreum		100.0	0.76			
53125011	**Good identification**					**53125011**
Eubacterium saburreum		100.0	0.42	URE 1		
53125012	**Very good identification**					**53125012**
Eubacterium saburreum		100.0	0.64			
53125020	**Very good identification**					**53125020**
Eubacterium saburreum		100.0	0.65			
53125022	**Good identification**					**53125022**
Eubacterium saburreum		100.0	0.52			
53125030	**Very good identification**					**53125030**
Eubacterium saburreum		100.0	0.64			
53125032	**Good identification**					**53125032**
Eubacterium saburreum		100.0	0.51			
53125040	**Good identification**					**53125040**
Eubacterium saburreum		100.0	0.44	SOR 1		
53125050	**Good identification**					**53125050**
Eubacterium saburreum		100.0	0.42	SOR 1		
53125100	**Very good identification**					**53125100**
Eubacterium saburreum		100.0	0.84			
53125101	**Good identification**					**53125101**
Eubacterium saburreum		100.0	0.50	URE 1		
53125102	**Very good identification**					**53125102**
Eubacterium saburreum		100.0	0.71			
53125103	**Good identification**					**53125103**
Eubacterium saburreum		100.0	0.37	URE 1		
53125110	**Very good identification**					**53125110**
Eubacterium saburreum		100.0	0.83			
53125111	**Good identification**					**53125111**
Eubacterium saburreum		100.0	0.49	URE 1		
53125112	**Very good identification**					**53125112**
Eubacterium saburreum		100.0	0.70			

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
53125113	**Good identification**					**53125113**
Eubacterium saburreum		100.0	0.36	URE 1		
53125120	**Very good identification**					**53125120**
Eubacterium saburreum		100.0	0.71			
53125121	**Good identification**					**53125121**
Eubacterium saburreum		100.0	0.37	URE 1		
53125122	**Very good identification**					**53125122**
Eubacterium saburreum		100.0	0.58			
53125130	**Very good identification**					**53125130**
Eubacterium saburreum		100.0	0.70			
53125131	**Good identification**					**53125131**
Eubacterium saburreum		100.0	0.36	URE 1		
53125132	**Very good identification**					**53125132**
Eubacterium saburreum		100.0	0.57			
53125140	**Good identification**					**53125140**
Eubacterium saburreum		100.0	0.50	SOR 1		
53125142	**Good identification**					**53125142**
Eubacterium saburreum		100.0	0.37	SOR 1		
53125150	**Good identification**					**53125150**
Eubacterium saburreum		100.0	0.49	SOR 1		
53125152	**Good identification**					**53125152**
Eubacterium saburreum		100.0	0.36	SOR 1		
53125160	**Good identification**					**53125160**
Eubacterium saburreum		100.0	0.37	SOR 1		
53125170	**Good identification**					**53125170**
Eubacterium saburreum		100.0	0.36	SOR 1		
53125200	**Good identification**					**53125200**
Eubacterium saburreum		100.0	0.44	RHA 1		
53125210	**Good identification**					**53125210**
Eubacterium saburreum		100.0	0.42	RHA 1		
53125300	**Good identification**					**53125300**
Eubacterium saburreum		100.0	0.50	RHA 1		
53125302	**Good identification**					**53125302**
Eubacterium saburreum		100.0	0.37	RHA 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53125310	**Good identification**					**53125310**
	Eubacterium saburreum	100.0	0.49	RHA 1		
53125312	**Good identification**					**53125312**
	Eubacterium saburreum	100.0	0.36	RHA 1		
53125320	**Good identification**					**53125320**
	Eubacterium saburreum	100.0	0.37	RHA 1		
53125330	**Good identification**					**53125330**
	Eubacterium saburreum	100.0	0.36	RHA 1		
53125400	**Good identification**					**53125400**
	Eubacterium saburreum	100.0	0.44	ARA 1		
53125410	**Good identification**					**53125410**
	Eubacterium saburreum	100.0	0.42	ARA 1		
53125500	**Good identification**					**53125500**
	Eubacterium saburreum	100.0	0.50	ARA 1		
53125502	**Good identification**					**53125502**
	Eubacterium saburreum	100.0	0.37	ARA 1		
53125510	**Good identification**					**53125510**
	Eubacterium saburreum	100.0	0.49	ARA 1		
53125512	**Good identification**					**53125512**
	Eubacterium saburreum	100.0	0.36	ARA 1		
53125520	**Good identification**					**53125520**
	Eubacterium saburreum	100.0	0.37	ARA 1		
53125530	**Good identification**					**53125530**
	Eubacterium saburreum	100.0	0.36	ARA 1		
53127000	**Good identification**					**53127000**
	Eubacterium saburreum	100.0	0.44	MAN 1		
53127010	**Good identification**					**53127010**
	Eubacterium saburreum	100.0	0.42	MAN 1		
53127100	**Good identification**					**53127100**
	Eubacterium saburreum	100.0	0.50	MAN 1		
53127102	**Good identification**					**53127102**
	Eubacterium saburreum	100.0	0.37	MAN 1		
53127110	**Good identification**					**53127110**
	Eubacterium saburreum	100.0	0.49	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53127112	**Good identification**					**53127112**
Eubacterium saburreum		100.0	0.36	MAN 1		
53127120	**Good identification**					**53127120**
Eubacterium saburreum		100.0	0.37	MAN 1		
53127130	**Good identification**					**53127130**
Eubacterium saburreum		100.0	0.36	MAN 1		
53130000	**Very good identification**					**53130000**
					SPO	
Eubacterium saburreum		95.39	0.57	GAL 99	1	
Clostridium botulinum biovar A		1.53	0.14		99	
Clostridium sporogenes		1.26	0.11		99	
53130002	**Good identification**					**53130002**
					SPO	
Eubacterium saburreum		97.56	0.44	GAL 99	1	
Clostridium sporogenes		2.44	0.03		99	
53130010	**Good identification**					**53130010**
Eubacterium saburreum		100.0	0.55	GAL 99		
53130012	**Good identification**					**53130012**
Eubacterium saburreum		100.0	0.42	GAL 99		
53130020	**Good identification**					**53130020**
					SPO	
Eubacterium saburreum		97.30	0.44	GAL 99	1	
Clostridium cadaveris		2.70	0.03		99	
53130030	**Good identification**					**53130030**
Eubacterium saburreum		100.0	0.42	GAL 99		
53130100	**Very good identification**					**53130100**
Eubacterium saburreum		99.60	0.63	GAL 99		
Fusobacterium mortiferum		0.40	0.24			
53130102	**Good identification**					**53130102**
Eubacterium saburreum		100.0	0.50	GAL 99		
53130110	**Very good identification**					**53130110**
Eubacterium saburreum		100.0	0.61	GAL 99		
53130112	**Good identification**					**53130112**
Eubacterium saburreum		100.0	0.48	GAL 99		
53130120	**Good identification**					**53130120**
Eubacterium saburreum		100.0	0.50	GAL 99		
53130122	**Good identification**					**53130122**
Eubacterium saburreum		100.0	0.37	GAL 99		
53130130	**Good identification**					**53130130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.48	GAL 99		
53130132	**Good identification**					**53130132**
Eubacterium saburreum		100.0	0.36	GAL 99		
53131000	**Very good identification**					**53131000**
Eubacterium saburreum		99.99	0.91			
Prevotella intermedia		0.01	0.11			
Fusobacterium mortiferum		0.01	0.20			
53131001	**Very good identification**					**53131001**
					SPO	
Eubacterium saburreum		99.70	0.57	URE 1	1	
Clostridium sordellii		0.30	-.03		99	
53131002	**Very good identification**					**53131002**
Eubacterium saburreum		100.0	0.78			
53131003	**Good identification**					**53131003**
Eubacterium saburreum		100.0	0.44	URE 1		
53131010	**Very good identification**					**53131010**
Eubacterium saburreum		100.0	0.89			
53131011	**Good identification**					**53131011**
Eubacterium saburreum		100.0	0.55	URE 1		
53131012	**Very good identification**					**53131012**
Eubacterium saburreum		100.0	0.76			
53131013	**Good identification**					**53131013**
Eubacterium saburreum		100.0	0.42	URE 1		
53131020	**Very good identification**					**53131020**
Eubacterium saburreum		99.98	0.78			
Clostridium perfringens		0.02	0.07			
53131021	**Good identification**					**53131021**
Eubacterium saburreum		100.0	0.44	URE 1		
53131022	**Very good identification**					**53131022**
Eubacterium saburreum		99.94	0.65			
Clostridium perfringens		0.06	0.01			
53131030	**Very good identification**					**53131030**
Eubacterium saburreum		100.0	0.76			
53131031	**Good identification**					**53131031**
Eubacterium saburreum		100.0	0.42	URE 1		
53131032	**Very good identification**					**53131032**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.63				
53131040	**Very good identification**						**53131040**
Eubacterium saburreum		100.0	0.57	SOR 1			
53131042	**Good identification**						**53131042**
Eubacterium saburreum		100.0	0.44	SOR 1			
53131050	**Good identification**						**53131050**
Eubacterium saburreum		100.0	0.55	SOR 1			
53131052	**Good identification**						**53131052**
Eubacterium saburreum		100.0	0.42	SOR 1			
53131060	**Good identification**						**53131060**
Eubacterium saburreum		100.0	0.44	SOR 1			
53131070	**Good identification**						**53131070**
Eubacterium saburreum		100.0	0.42	SOR 1			
53131100	**Excellent identification**						**53131100**
					COC	GRA	
Eubacterium saburreum		99.98	0.97		1	85	
Atopobium parvulum		0.01	0.19		99	99	
Fusobacterium mortiferum		0.01	0.27		1	1	
53131101	**Very good identification**						**53131101**
					SPO		
Eubacterium saburreum		95.60	0.63	URE 1	1		
Clostridium chauvoei		4.40	0.30		99		
53131102	**Very good identification**						**53131102**
					COC		
Eubacterium saburreum		99.99	0.84		1		
Atopobium parvulum		0.01	0.06		99		
53131103	**Good identification**						**53131103**
					SPO		
Eubacterium saburreum		95.60	0.50	URE 1	1		
Clostridium chauvoei		4.40	0.17		99		
53131110	**Excellent identification**						**53131110**
Eubacterium saburreum		100.0	0.96				
53131111	**Very good identification**						**53131111**
Eubacterium saburreum		100.0	0.61	URE 1			
53131112	**Very good identification**						**53131112**
Eubacterium saburreum		100.0	0.83				
53131113	**Good identification**						**53131113**
Eubacterium saburreum		100.0	0.48	URE 1			
53131120	**Very good identification**						**53131120**
					COC		
Eubacterium saburreum		99.91	0.84		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		0.07	0.21		1	
Atopobium parvulum		0.01	0.06		99	
53131121	**Acceptable identification**					**53131121**
					SPO	
Eubacterium saburreum		92.00	0.50	URE	1	
Clostridium chauvoei		8.00	0.22		99	
53131122	**Very good identification**					**53131122**
Eubacterium saburreum		99.81	0.71			
Clostridium perfringens		0.20	0.16			
53131123	**Acceptable identification**					**53131123**
					SPO	
Eubacterium saburreum		92.00	0.37	URE	1	
Clostridium chauvoei		8.00	0.09		99	
53131130	**Very good identification**					**53131130**
Eubacterium saburreum		100.0	0.83			
53131131	**Good identification**					**53131131**
Eubacterium saburreum		100.0	0.48	URE	1	
53131132	**Very good identification**					**53131132**
Eubacterium saburreum		100.0	0.70			
53131133	**Good identification**					**53131133**
Eubacterium saburreum		100.0	0.36	URE	1	
53131140	**Very good identification**					**53131140**
Eubacterium saburreum		100.0	0.63	SOR	1	
53131142	**Good identification**					**53131142**
Eubacterium saburreum		100.0	0.50	SOR	1	
53131150	**Very good identification**					**53131150**
Eubacterium saburreum		100.0	0.61	SOR	1	
53131152	**Good identification**					**53131152**
Eubacterium saburreum		100.0	0.48	SOR	1	
53131160	**Good identification**					**53131160**
Eubacterium saburreum		98.64	0.50	SOR	1	
Clostridium perfringens		1.36	0.09			
53131162	**Good identification**					**53131162**
Eubacterium saburreum		96.45	0.37	SOR	1	
Clostridium perfringens		3.55	0.03			
53131170	**Good identification**					**53131170**
Eubacterium saburreum		100.0	0.48	SOR	1	
53131172	**Good identification**					**53131172**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Eubacterium saburreum	100.0	0.36	SOR 1		
53131200	**Very good identification**					**53131200**
	Eubacterium saburreum	100.0	0.57	RHA 1		
53131202	**Good identification**					**53131202**
	Eubacterium saburreum	100.0	0.44	RHA 1		
53131210	**Good identification**					**53131210**
	Eubacterium saburreum	100.0	0.55	RHA 1		
53131212	**Good identification**					**53131212**
	Eubacterium saburreum	100.0	0.42	RHA 1		
53131220	**Good identification**					**53131220**
	Eubacterium saburreum	100.0	0.44	RHA 1		
53131230	**Good identification**					**53131230**
	Eubacterium saburreum	100.0	0.42	RHA 1		
53131300	**Very good identification**					**53131300**
	Eubacterium saburreum	100.0	0.63	RHA 1		
53131302	**Good identification**					**53131302**
	Eubacterium saburreum	100.0	0.50	RHA 1		
53131310	**Very good identification**					**53131310**
	Eubacterium saburreum	100.0	0.61	RHA 1		
53131312	**Good identification**					**53131312**
	Eubacterium saburreum	100.0	0.48	RHA 1		
53131320	**Good identification**					**53131320**
	Eubacterium saburreum	100.0	0.50	RHA 1		
53131322	**Good identification**					**53131322**
	Eubacterium saburreum	100.0	0.37	RHA 1		
53131330	**Good identification**					**53131330**
	Eubacterium saburreum	100.0	0.48	RHA 1		
53131332	**Good identification**					**53131332**
	Eubacterium saburreum	100.0	0.36	RHA 1		
53131400	**Very good identification**					**53131400**
	Eubacterium saburreum	100.0	0.57	ARA 1		
53131402	**Good identification**					**53131402**
	Eubacterium saburreum	100.0	0.44	ARA 1		
53131410	**Good identification**					**53131410**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.55	ARA 1		
53131412	**Good identification**					**53131412**
Eubacterium saburreum		100.0	0.42	ARA 1		
53131420	**Good identification**					**53131420**
Eubacterium saburreum		100.0	0.44	ARA 1		
53131430	**Good identification**					**53131430**
Eubacterium saburreum		100.0	0.42	ARA 1		
53131500	**Very good identification**					**53131500**
Eubacterium saburreum		100.0	0.63	ARA 1		
53131502	**Good identification**					**53131502**
Eubacterium saburreum		100.0	0.50	ARA 1		
53131510	**Very good identification**					**53131510**
Eubacterium saburreum		100.0	0.61	ARA 1		
53131512	**Good identification**					**53131512**
Eubacterium saburreum		100.0	0.48	ARA 1		
53131520	**Good identification**					**53131520**
Eubacterium saburreum		100.0	0.50	ARA 1		
53131522	**Good identification**					**53131522**
Eubacterium saburreum		100.0	0.37	ARA 1		
53131530	**Good identification**					**53131530**
Eubacterium saburreum		99.33	0.48	ARA 1		
Bifidobacterium longum subsp. longum		0.67	0.07			
53131532	**Good identification**					**53131532**
Eubacterium saburreum		96.33	0.36	ARA 1		
Bifidobacterium longum subsp. longum		3.67	0.07			
53133000	**Very good identification**					**53133000**
Eubacterium saburreum		100.0	0.57	MAN 1		
53133002	**Good identification**					**53133002**
Eubacterium saburreum		100.0	0.44	MAN 1		
53133010	**Good identification**					**53133010**
Eubacterium saburreum		100.0	0.55	MAN 1		
53133012	**Good identification**					**53133012**
Eubacterium saburreum		100.0	0.42	MAN 1		
53133020	**Good identification**					**53133020**
Eubacterium saburreum		100.0	0.44	MAN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53133030	**Good identification**					**53133030**
Eubacterium saburreum		100.0	0.42	MAN 1		
53133100	**Very good identification**					**53133100**
Eubacterium saburreum		99.91	0.63	MAN 1		
Propionibacterium propionicum		0.09	0.14			
53133102	**Good identification**					**53133102**
Eubacterium saburreum		99.24	0.50	MAN 1		
Propionibacterium propionicum		0.76	0.17			
53133110	**Very good identification**					**53133110**
Eubacterium saburreum		100.0	0.61	MAN 1		
53133112	**Good identification**					**53133112**
Eubacterium saburreum		100.0	0.48	MAN 1		
53133120	**Good identification**					**53133120**
Eubacterium saburreum		100.0	0.50	MAN 1		
53133122	**Good identification**					**53133122**
Eubacterium saburreum		100.0	0.37	MAN 1		
53133130	**Good identification**					**53133130**
Eubacterium saburreum		100.0	0.48	MAN 1		
53133132	**Good identification**					**53133132**
Eubacterium saburreum		100.0	0.36	MAN 1		
53134000	**Good identification**					**53134000**
Eubacterium saburreum		100.0	0.52	GAL 99		
53134002	**Good identification**					**53134002**
Eubacterium saburreum		100.0	0.39	GAL 99		
53134010	**Good identification**					**53134010**
Eubacterium saburreum		100.0	0.50	GAL 99		
53134012	**Good identification**					**53134012**
Eubacterium saburreum		100.0	0.38	GAL 99		
53134020	**Good identification**					**53134020**
Eubacterium saburreum		100.0	0.39	GAL 99		
53134030	**Good identification**					**53134030**
Eubacterium saburreum		100.0	0.38	GAL 99		
53134100	**Very good identification**					**53134100**
Eubacterium saburreum		100.0	0.58	GAL 99		
53134102	**Good identification**					**53134102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.45	GAL 99		
53134110	**Very good identification**					**53134110**
Eubacterium saburreum		100.0	0.57	GAL 99		
53134112	**Good identification**					**53134112**
Eubacterium saburreum		100.0	0.44	GAL 99		
53134120	**Good identification**					**53134120**
Eubacterium saburreum		100.0	0.45	GAL 99		
53134130	**Good identification**					**53134130**
Eubacterium saburreum		100.0	0.44	GAL 99		
53135000	**Very good identification**					**53135000**
Eubacterium saburreum		100.0	0.86			
53135001	**Good identification**					**53135001**
Eubacterium saburreum		100.0	0.52	URE 1		
53135002	**Very good identification**					**53135002**
Eubacterium saburreum		100.0	0.73			
53135003	**Good identification**					**53135003**
Eubacterium saburreum		100.0	0.39	URE 1		
53135010	**Very good identification**					**53135010**
Eubacterium saburreum		100.0	0.85			
53135011	**Good identification**					**53135011**
Eubacterium saburreum		100.0	0.50	URE 1		
53135012	**Very good identification**					**53135012**
Eubacterium saburreum		100.0	0.72			
53135013	**Good identification**					**53135013**
Eubacterium saburreum		100.0	0.38	URE 1		
53135020	**Very good identification**					**53135020**
Eubacterium saburreum		100.0	0.73			
53135021	**Good identification**					**53135021**
Eubacterium saburreum		100.0	0.39	URE 1		
53135022	**Very good identification**					**53135022**
Eubacterium saburreum		100.0	0.60			
53135030	**Very good identification**					**53135030**
Eubacterium saburreum		100.0	0.72			
53135031	**Good identification**					**53135031**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	URE 1		
53135032	**Very good identification**					**53135032**
Eubacterium saburreum		100.0	0.59			
53135040	**Good identification**					**53135040**
Eubacterium saburreum		100.0	0.52	SOR 1		
53135042	**Good identification**					**53135042**
Eubacterium saburreum		100.0	0.39	SOR 1		
53135050	**Good identification**					**53135050**
Eubacterium saburreum		100.0	0.50	SOR 1		
53135052	**Good identification**					**53135052**
Eubacterium saburreum		100.0	0.38	SOR 1		
53135060	**Good identification**					**53135060**
Eubacterium saburreum		100.0	0.39	SOR 1		
53135070	**Good identification**					**53135070**
Eubacterium saburreum		100.0	0.38	SOR 1		
53135100	**Excellent identification**					**53135100**
					COC	
Eubacterium saburreum		99.97	0.92		1	
Atopobium parvulum		0.03	0.23		99	
53135101	**Very good identification**					**53135101**
Eubacterium saburreum		100.0	0.58	URE 1		
53135102	**Very good identification**					**53135102**
					COC	
Eubacterium saburreum		99.97	0.80		1	
Atopobium parvulum		0.03	0.10		99	
53135103	**Good identification**					**53135103**
Eubacterium saburreum		100.0	0.45	URE 1		
53135110	**Very good identification**					**53135110**
Eubacterium saburreum		100.0	0.91			
53135111	**Very good identification**					**53135111**
Eubacterium saburreum		100.0	0.57	URE 1		
53135112	**Very good identification**					**53135112**
Eubacterium saburreum		100.0	0.78			
53135113	**Good identification**					**53135113**
Eubacterium saburreum		100.0	0.44	URE 1		
53135120	**Very good identification**					**53135120**
					COC	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.97	0.80		1	
Atopobium parvulum		0.03	0.10		99	
53135121	**Good identification**					**53135121**
Eubacterium saburreum		100.0	0.45	URE	1	
53135122	**Very good identification**					**53135122**
Eubacterium saburreum		100.0	0.67			
53135130	**Very good identification**					**53135130**
Eubacterium saburreum		100.0	0.78			
53135131	**Good identification**					**53135131**
Eubacterium saburreum		100.0	0.44	URE	1	
53135132	**Very good identification**					**53135132**
Eubacterium saburreum		100.0	0.65			
53135140	**Very good identification**					**53135140**
Eubacterium saburreum		100.0	0.58	SOR	1	
53135142	**Good identification**					**53135142**
Eubacterium saburreum		100.0	0.45	SOR	1	
53135150	**Very good identification**					**53135150**
Eubacterium saburreum		100.0	0.57	SOR	1	
53135152	**Good identification**					**53135152**
Eubacterium saburreum		100.0	0.44	SOR	1	
53135160	**Good identification**					**53135160**
Eubacterium saburreum		100.0	0.45	SOR	1	
53135170	**Good identification**					**53135170**
Eubacterium saburreum		100.0	0.44	SOR	1	
53135200	**Good identification**					**53135200**
Eubacterium saburreum		100.0	0.52	RHA	1	
53135202	**Good identification**					**53135202**
Eubacterium saburreum		100.0	0.39	RHA	1	
53135210	**Good identification**					**53135210**
Eubacterium saburreum		100.0	0.50	RHA	1	
53135212	**Good identification**					**53135212**
Eubacterium saburreum		100.0	0.38	RHA	1	
53135220	**Good identification**					**53135220**
Eubacterium saburreum		100.0	0.39	RHA	1	
53135230	**Good identification**					**53135230**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	RHA 1		
53135300	**Very good identification**					**53135300**
Eubacterium saburreum		100.0	0.58	RHA 1		
53135302	**Good identification**					**53135302**
Eubacterium saburreum		100.0	0.45	RHA 1		
53135310	**Very good identification**					**53135310**
Eubacterium saburreum		100.0	0.57	RHA 1		
53135312	**Good identification**					**53135312**
Eubacterium saburreum		100.0	0.44	RHA 1		
53135320	**Good identification**					**53135320**
Eubacterium saburreum		100.0	0.45	RHA 1		
53135330	**Good identification**					**53135330**
Eubacterium saburreum		100.0	0.44	RHA 1		
53135400	**Good identification**					**53135400**
Eubacterium saburreum		100.0	0.52	ARA 1		
53135402	**Good identification**					**53135402**
Eubacterium saburreum		100.0	0.39	ARA 1		
53135410	**Good identification**					**53135410**
Eubacterium saburreum		100.0	0.50	ARA 1		
53135412	**Good identification**					**53135412**
Eubacterium saburreum		100.0	0.38	ARA 1		
53135420	**Good identification**					**53135420**
Eubacterium saburreum		100.0	0.39	ARA 1		
53135430	**Good identification**					**53135430**
Eubacterium saburreum		100.0	0.38	ARA 1		
53135500	**Very good identification**					**53135500**
Eubacterium saburreum		100.0	0.58	ARA 1		
53135502	**Good identification**					**53135502**
Eubacterium saburreum		98.76	0.45	ARA 1		
Bifidobacterium adolescentis		1.24	0.11			
53135510	**Very good identification**					**53135510**
Eubacterium saburreum		99.84	0.57	ARA 1		
Bifidobacterium longum subsp. longum		0.17	0.05			
53135512	**Good identification**					**53135512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		97.27	0.44	ARA 1		
Bifidobacterium adolescentis		1.82	0.12			
Bifidobacterium longum subsp. longum		0.91	0.05			
53135520	**Good identification**					**53135520**
Eubacterium saburreum		98.23	0.45	ARA 1		
Bifidobacterium longum subsp. longum		1.77	0.12			
53135530	**Good identification**					**53135530**
Eubacterium saburreum		96.39	0.44	ARA 1		
Bifidobacterium longum subsp. longum		3.61	0.16			
53137000	**Good identification**					**53137000**
Eubacterium saburreum		100.0	0.52	MAN 1		
53137002	**Good identification**					**53137002**
Eubacterium saburreum		100.0	0.39	MAN 1		
53137010	**Good identification**					**53137010**
Eubacterium saburreum		100.0	0.50	MAN 1		
53137012	**Good identification**					**53137012**
Eubacterium saburreum		100.0	0.38	MAN 1		
53137020	**Good identification**					**53137020**
Eubacterium saburreum		100.0	0.39	MAN 1		
53137030	**Good identification**					**53137030**
Eubacterium saburreum		100.0	0.38	MAN 1		
53137100	**Very good identification**					**53137100**
Eubacterium saburreum		100.0	0.58	MAN 1		
53137102	**Good identification**					**53137102**
Eubacterium saburreum		100.0	0.45	MAN 1		
53137110	**Very good identification**					**53137110**
Eubacterium saburreum		100.0	0.57	MAN 1		
53137112	**Good identification**					**53137112**
Eubacterium saburreum		100.0	0.44	MAN 1		
53137120	**Good identification**					**53137120**
Eubacterium saburreum		100.0	0.45	MAN 1		
53137130	**Good identification**					**53137130**
Eubacterium saburreum		100.0	0.44	MAN 1		
53141000	**Good identification**					**53141000**
Eubacterium saburreum		100.0	0.51	CEL 1		
53141002	**Good identification**					**53141002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.39	CEL 1		
53141010	**Good identification**					**53141010**
Eubacterium saburreum		100.0	0.50	CEL 1		
53141012	**Good identification**					**53141012**
Eubacterium saburreum		100.0	0.37	CEL 1		
53141020	**Good identification**					**53141020**
Eubacterium saburreum		100.0	0.39	CEL 1		
53141030	**Good identification**					**53141030**
Eubacterium saburreum		100.0	0.37	CEL 1		
53141100	**Very good identification**					**53141100**
Eubacterium saburreum		100.0	0.58	CEL 1		
53141102	**Good identification**					**53141102**
Eubacterium saburreum		100.0	0.45	CEL 1		
53141110	**Very good identification**					**53141110**
Eubacterium saburreum		100.0	0.56	CEL 1		
53141112	**Good identification**					**53141112**
Eubacterium saburreum		100.0	0.43	CEL 1		
53141120	**Good identification**					**53141120**
Eubacterium saburreum		98.55	0.45	CEL 1		
Prevotella melaninogenica		1.45	-.03			
53141130	**Good identification**					**53141130**
Eubacterium saburreum		100.0	0.43	CEL 1		
53145000	**Good identification**					**53145000**
Eubacterium saburreum		100.0	0.47	CEL 1		
53145010	**Good identification**					**53145010**
Eubacterium saburreum		100.0	0.45	CEL 1		
53145100	**Good identification**					**53145100**
Eubacterium saburreum		100.0	0.53	CEL 1		
53145102	**Good identification**					**53145102**
Eubacterium saburreum		100.0	0.40	CEL 1		
53145110	**Good identification**					**53145110**
Eubacterium saburreum		100.0	0.52	CEL 1		
53145112	**Good identification**					**53145112**
Eubacterium saburreum		100.0	0.39	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53145120	**Good identification**						**53145120**
Eubacterium saburreum		100.0	0.40	CEL 1			
53145130	**Good identification**						**53145130**
Eubacterium saburreum		100.0	0.39	CEL 1			
53151000	**Very good identification**						**53151000**
Eubacterium saburreum		98.95	0.60	CEL 1			
Prevotella intermedia		0.48	0.11				
Fusobacterium mortiferum		0.45	0.21				
53151002	**Good identification**						**53151002**
Eubacterium saburreum		100.0	0.47	CEL 1			
53151010	**Very good identification**						**53151010**
Eubacterium saburreum		100.0	0.58	CEL 1			
53151012	**Good identification**						**53151012**
Eubacterium saburreum		100.0	0.45	CEL 1			
53151020	**Good identification**						**53151020**
Eubacterium saburreum		100.0	0.47	CEL 1			
53151030	**Good identification**						**53151030**
Eubacterium saburreum		100.0	0.45	CEL 1			
53151100	**Very good identification**						**53151100**
					GRA		
Eubacterium saburreum		99.33	0.66	CEL 1	85		
Fusobacterium mortiferum		0.45	0.27		1		
Collinsella aerofaciens		0.22	-.03		99		
53151102	**Good identification**						**53151102**
					CAT		
Eubacterium saburreum		99.06	0.53	CEL 1	1		
Capnocytophaga ochracea		0.66	0.04		90		
Fusobacterium mortiferum		0.28	0.11		5		
53151110	**Very good identification**						**53151110**
Eubacterium saburreum		100.0	0.64	CEL 1			
53151112	**Good identification**						**53151112**
Eubacterium saburreum		100.0	0.52	CEL 1			
53151120	**Acceptable identification**						**53151120**
					SPO	GRA	
Eubacterium saburreum		94.52	0.53	CEL 1	1	85	
Prevotella melaninogenica		4.18	0.14		1	1	
Clostridium paraputrificum		0.68	-.03		99	99	
53151122	**Low discrimination**						**53151122**
					CAT		
Eubacterium saburreum		66.30	0.40	CEL 1	1		
Capnocytophaga ochracea		22.45	0.20		90		
Prevotella oralis		5.60	0.09		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53151130	**Good identification**					**53151130**
Eubacterium saburreum		100.0	0.52	CEL 1		
53151132	**Good identification**					**53151132**
					CAT	
Eubacterium saburreum		95.60	0.39	CEL 1	1	
Capnocytophaga ochracea		4.40	0.04		90	
53155000	**Good identification**					**53155000**
Eubacterium saburreum		100.0	0.55	CEL 1		
53155002	**Good identification**					**53155002**
Eubacterium saburreum		100.0	0.42	CEL 1		
53155010	**Good identification**					**53155010**
Eubacterium saburreum		100.0	0.53	CEL 1		
53155012	**Good identification**					**53155012**
Eubacterium saburreum		100.0	0.41	CEL 1		
53155020	**Good identification**					**53155020**
Eubacterium saburreum		100.0	0.42	CEL 1		
53155030	**Good identification**					**53155030**
Eubacterium saburreum		100.0	0.41	CEL 1		
53155100	**Very good identification**					**53155100**
Eubacterium saburreum		100.0	0.61	CEL 1		
53155102	**Good identification**					**53155102**
Eubacterium saburreum		100.0	0.48	CEL 1		
53155110	**Very good identification**					**53155110**
Eubacterium saburreum		100.0	0.60	CEL 1		
53155112	**Good identification**					**53155112**
Eubacterium saburreum		100.0	0.47	CEL 1		
53155120	**Good identification**					**53155120**
Eubacterium saburreum		100.0	0.48	CEL 1		
53155122	**Low discrimination**					**53155122**
					CAT	
Eubacterium saburreum		75.45	0.35	CEL 1	1	
Capnocytophaga ochracea		11.86	0.10		90	
Bacteroides uniformis		9.19	0.01		5	
53155130	**Good identification**					**53155130**
Eubacterium saburreum		100.0	0.47	CEL 1		
53161000	**Good identification**					**53161000**
Eubacterium saburreum		100.0	0.48	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53161002	**Good identification**					**53161002**
Eubacterium saburreum		100.0	0.36	CEL 1		
53161010	**Good identification**					**53161010**
Eubacterium saburreum		100.0	0.47	CEL 1		
53161020	**Good identification**					**53161020**
Eubacterium saburreum		100.0	0.36	CEL 1		
53161100	**Good identification**					**53161100**
Eubacterium saburreum		100.0	0.55	CEL 1		
53161102	**Good identification**					**53161102**
Eubacterium saburreum		100.0	0.42	CEL 1		
53161110	**Good identification**					**53161110**
Eubacterium saburreum		100.0	0.53	CEL 1		
53161112	**Good identification**					**53161112**
Eubacterium saburreum		100.0	0.40	CEL 1		
53161120	**Good identification**					**53161120**
Eubacterium saburreum		100.0	0.42	CEL 1		
53161130	**Good identification**					**53161130**
Eubacterium saburreum		100.0	0.40	CEL 1		
53165000	**Good identification**					**53165000**
Eubacterium saburreum		100.0	0.44	CEL 1		
53165010	**Good identification**					**53165010**
Eubacterium saburreum		100.0	0.42	CEL 1		
53165100	**Good identification**					**53165100**
Eubacterium saburreum		100.0	0.50	CEL 1		
53165102	**Good identification**					**53165102**
Eubacterium saburreum		100.0	0.37	CEL 1		
53165110	**Good identification**					**53165110**
Eubacterium saburreum		100.0	0.49	CEL 1		
53165112	**Good identification**					**53165112**
Eubacterium saburreum		100.0	0.36	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53165120	**Good identification**						**53165120**
Eubacterium saburreum		100.0	0.37	CEL 1			
53165130	**Good identification**						**53165130**
Eubacterium saburreum		100.0	0.36	CEL 1			
53171000	**Very good identification**						**53171000**
					GRA		
Eubacterium saburreum		99.24	0.57	CEL 1	85		
Lactobacillus jensenii		0.56	0.10		99		
Fusobacterium mortiferum		0.20	0.12		1		
53171002	**Good identification**						**53171002**
Eubacterium saburreum		97.90	0.44	CEL 1			
Lactobacillus jensenii		2.10	0.07				
53171010	**Good identification**						**53171010**
Eubacterium saburreum		99.77	0.55	CEL 1			
Lactobacillus jensenii		0.23	0.02				
53171012	**Good identification**						**53171012**
Eubacterium saburreum		100.0	0.42	CEL 1			
53171020	**Good identification**						**53171020**
Eubacterium saburreum		100.0	0.44	CEL 1			
53171030	**Good identification**						**53171030**
Eubacterium saburreum		100.0	0.42	CEL 1			
53171100	**Very good identification**						**53171100**
					COC	GRA	
Eubacterium saburreum		97.57	0.63	CEL 1	1	85	
Atopobium parvulum		2.23	0.25		99	99	
Fusobacterium mortiferum		0.20	0.18		1	1	
53171102	**Good identification**						**53171102**
					COC		
Eubacterium saburreum		97.77	0.50	CEL 1	1		
Atopobium parvulum		2.23	0.12		99		
53171110	**Very good identification**						**53171110**
Eubacterium saburreum		100.0	0.61	CEL 1			
53171112	**Good identification**						**53171112**
Eubacterium saburreum		98.26	0.48	CEL 1			
Leptotrichia buccalis		1.74	-.03				
53171120	**Good identification**						**53171120**
					COC		
Eubacterium saburreum		97.77	0.50	CEL 1	1		
Atopobium parvulum		2.23	0.12		99		
53171122	**Good identification**						**53171122**
Eubacterium saburreum		100.0	0.37	CEL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53171130	**Good identification**					**53171130**
Eubacterium saburreum		100.0	0.48	CEL 1		
53171132	**Good identification**					**53171132**
Eubacterium saburreum		100.0	0.36	CEL 1		
53175000	**Good identification**					**53175000**
Eubacterium saburreum		100.0	0.52	CEL 1		
53175002	**Good identification**					**53175002**
Eubacterium saburreum		100.0	0.39	CEL 1		
53175010	**Good identification**					**53175010**
Eubacterium saburreum		100.0	0.50	CEL 1		
53175012	**Good identification**					**53175012**
Eubacterium saburreum		100.0	0.38	CEL 1		
53175020	**Good identification**					**53175020**
Eubacterium saburreum		100.0	0.39	CEL 1		
53175030	**Good identification**					**53175030**
Eubacterium saburreum		100.0	0.38	CEL 1		
53175100	**Good identification**					**53175100**
					COC	
Eubacterium saburreum		92.54	0.58	CEL 1	1	
Atopobium parvulum		7.29	0.30		99	
Actinomyces israelii		0.17	0.15		1	
53175102	**Acceptable identification**					**53175102**
					COC	
Eubacterium saburreum		91.40	0.45	CEL 1	1	
Atopobium parvulum		7.20	0.17		99	
Actinomyces israelii		1.39	0.18		1	
53175110	**Very good identification**					**53175110**
Eubacterium saburreum		100.0	0.57	CEL 1		
53175112	**Good identification**					**53175112**
Eubacterium saburreum		100.0	0.44	CEL 1		
53175120	**Acceptable identification**					**53175120**
					COC	
Eubacterium saburreum		92.69	0.45	CEL 1	1	
Atopobium parvulum		7.31	0.17		99	
53175130	**Good identification**					**53175130**
Eubacterium saburreum		100.0	0.44	CEL 1		
53200000	**Low discrimination**					**53200000**
					SPO	
Clostridium bifermentans		70.96	0.42	SUC 1	99	
Eubacterium tortuosum		27.53	0.32	SAL 10	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		1.28	0.13		99		
53201000	**Very good identification**						**53201000**
					SPO		
Eubacterium tortuosum		99.49	0.66	IND 1	1		
Eubacterium saburreum		0.47	0.41		1		
Clostridium bifermentans		0.03	0.08		99		
53201002	**Good identification**						**53201002**
Eubacterium tortuosum		99.53	0.53	IND 1			
Eubacterium saburreum		0.47	0.28				
53201020	**Good identification**						**53201020**
Eubacterium tortuosum		99.53	0.53	IND 1			
Eubacterium saburreum		0.47	0.28				
53201022	**Good identification**						**53201022**
Eubacterium tortuosum		99.53	0.40	IND 1			
Eubacterium saburreum		0.47	0.15				
53201100	**Low discrimination**						**53201100**
					GRA		
Eubacterium saburreum		51.57	0.47	SAL 1	85		
Eubacterium tortuosum		47.11	0.32		99		
Fusobacterium mortiferum		1.33	0.22		1		
53201110	**Good identification**						**53201110**
Eubacterium saburreum		98.89	0.46	SAL 1			
Eubacterium tortuosum		1.12	-.03				
53205010	**Good identification**						**53205010**
Eubacterium saburreum		95.34	0.35	SAL 1			
Eubacterium tortuosum		4.66	-.03				
53205100	**Good identification**						**53205100**
Eubacterium saburreum		98.32	0.43	SAL 1			
Eubacterium tortuosum		1.69	-.03				
53205110	**Good identification**						**53205110**
Eubacterium saburreum		100.0	0.41	SAL 1			
53210000	**Low discrimination**						**53210000**
					SPO	GRA	
Fusobacterium mortiferum		83.26	0.47	IND 1	1	1	
Clostridium bifermentans		14.20	0.20		99	99	
Eubacterium saburreum		1.49	0.15		1	85	
53210100	**Good identification**						**53210100**
Fusobacterium mortiferum		98.24	0.53	IND 1			
Eubacterium saburreum		1.76	0.21				
53210102	**Good identification**						**53210102**
Fusobacterium mortiferum		100.0	0.37	IND 1			
53211000	**Low discrimination**						**53211000**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		39.09	0.49	SAL 1	85	
Fusobacterium mortiferum		33.14	0.50		1	
Eubacterium tortuosum		27.77	0.32		99	
53211002	**Low discrimination**					**53211002**
					GRA	
Eubacterium saburreum		44.56	0.36	SAL 1	85	
Eubacterium tortuosum		31.66	0.19		99	
Fusobacterium mortiferum		23.78	0.33		1	
53211010	**Good identification**					**53211010**
					GRA	
Eubacterium saburreum		98.11	0.48	SAL 1	85	
Fusobacterium mortiferum		1.03	0.16		1	
Eubacterium tortuosum		0.86	-.03		99	
53211020	**Low discrimination**					**53211020**
					GRA	
Eubacterium saburreum		56.85	0.36	SAL 1	85	
Eubacterium tortuosum		40.39	0.19		99	
Fusobacterium mortiferum		2.76	0.16		1	
53211100	**Low discrimination**					**53211100**
					GRA	
Eubacterium saburreum		54.03	0.56	SAL 1	85	
Fusobacterium mortiferum		45.80	0.56		1	
Eubacterium tortuosum		0.17	-.03		99	
53211102	**Low discrimination**					**53211102**
Eubacterium saburreum		65.20	0.43	SAL 1		
Fusobacterium mortiferum		34.80	0.40			
53211110	**Good identification**					**53211110**
Eubacterium saburreum		98.96	0.54	SAL 1		
Fusobacterium mortiferum		1.04	0.22			
53211112	**Good identification**					**53211112**
Eubacterium saburreum		100.0	0.41	SAL 1		
53211120	**Good identification**					**53211120**
Eubacterium saburreum		95.37	0.43	SAL 1		
Fusobacterium mortiferum		4.63	0.22			
53211130	**Good identification**					**53211130**
Eubacterium saburreum		100.0	0.41	SAL 1		
53215000	**Low discrimination**					**53215000**
					GRA	
Eubacterium saburreum		88.35	0.45	SAL 1	85	
Fusobacterium mortiferum		10.47	0.30		1	
Eubacterium tortuosum		1.18	-.03		99	
53215010	**Good identification**					**53215010**
Eubacterium saburreum		100.0	0.43	SAL 1		
53215100	**Low discrimination**					**53215100**
Eubacterium saburreum		89.41	0.51	SAL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		10.60	0.37			
53215102	**Acceptable identification**					**53215102**
Eubacterium saburreum		93.06	0.38	SAL 1		
Fusobacterium mortiferum		6.94	0.20			
53215110	**Good identification**					**53215110**
Eubacterium saburreum		100.0	0.49	SAL 1		
53215112	**Good identification**					**53215112**
Eubacterium saburreum		100.0	0.37	SAL 1		
53215120	**Good identification**					**53215120**
Eubacterium saburreum		100.0	0.38	SAL 1		
53215130	**Good identification**					**53215130**
Eubacterium saburreum		100.0	0.37	SAL 1		
53221010	**Good identification**					**53221010**
Eubacterium saburreum		96.20	0.37	SAL 1		
Eubacterium tortuosum		3.80	-.03			
53221100	**Good identification**					**53221100**
					GRA	
Eubacterium saburreum		97.53	0.44	SAL 1	85	
Eubacterium tortuosum		1.35	-.03		99	
Fusobacterium mortiferum		1.12	0.13		1	
53221110	**Good identification**					**53221110**
Eubacterium saburreum		100.0	0.43	SAL 1		
53225100	**Good identification**					**53225100**
Eubacterium saburreum		100.0	0.40	SAL 1		
53225110	**Good identification**					**53225110**
Eubacterium saburreum		100.0	0.38	SAL 1		
53230000	**Good identification**					**53230000**
Fusobacterium mortiferum		96.16	0.38	IND 1		
Eubacterium saburreum		3.84	0.12			
53230100	**Good identification**					**53230100**
Fusobacterium mortiferum		96.16	0.44	IND 1		
Eubacterium saburreum		3.84	0.18			
53231000	**Low discrimination**					**53231000**
					GRA	
Eubacterium saburreum		71.91	0.46	SAL 1	85	
Fusobacterium mortiferum		27.31	0.41		1	
Eubacterium tortuosum		0.77	-.03		99	
53231010	**Good identification**					**53231010**
Eubacterium saburreum		100.0	0.45	SAL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53231100	**Low discrimination**					**53231100**
Eubacterium saburreum		72.47	0.53	SAL 1		
Fusobacterium mortiferum		27.53	0.47			
53231102	**Low discrimination**					**53231102**
Eubacterium saburreum		80.70	0.40	SAL 1		
Fusobacterium mortiferum		19.30	0.31			
53231110	**Good identification**					**53231110**
Eubacterium saburreum		99.53	0.51	SAL 1		
Fusobacterium mortiferum		0.47	0.13			
53231112	**Good identification**					**53231112**
Eubacterium saburreum		100.0	0.38	SAL 1		
53231120	**Good identification**					**53231120**
Eubacterium saburreum		97.87	0.40	SAL 1		
Fusobacterium mortiferum		2.13	0.13			
53231130	**Good identification**					**53231130**
Eubacterium saburreum		100.0	0.38	SAL 1		
53235000	**Acceptable identification**					**53235000**
Eubacterium saburreum		94.96	0.42	SAL 1		
Fusobacterium mortiferum		5.04	0.22			
53235010	**Good identification**					**53235010**
Eubacterium saburreum		100.0	0.40	SAL 1		
53235100	**Acceptable identification**					**53235100**
Eubacterium saburreum		94.96	0.48	SAL 1		
Fusobacterium mortiferum		5.04	0.28			
53235102	**Good identification**					**53235102**
Eubacterium saburreum		96.77	0.35	SAL 1		
Fusobacterium mortiferum		3.23	0.11			
53235110	**Good identification**					**53235110**
Eubacterium saburreum		100.0	0.46	SAL 1		
53235120	**Good identification**					**53235120**
Eubacterium saburreum		100.0	0.35	SAL 1		
53241000	**Good identification**					**53241000**
Eubacterium tortuosum		100.0	0.50	IND 1		
53241002	**Good identification**					**53241002**
Eubacterium tortuosum		100.0	0.37	IND 1		
53241020	**Good identification**					**53241020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.37	IND 1		
53250000	**Good identification**					**53250000**
Fusobacterium mortiferum		100.0	0.39	IND 1		
53250100	**Good identification**					**53250100**
Fusobacterium mortiferum		100.0	0.45	IND 1		
53251000	**Low discrimination**					**53251000**
					GRA	
Fusobacterium mortiferum		71.03	0.42	IND 1	1	
Eubacterium tortuosum		19.84	0.15		99	
Lactobacillus catenaformis		5.33	0.12		99	
53251100	**Low discrimination**					**53251100**
					GRA	
Fusobacterium mortiferum		88.05	0.48	IND 1	1	
Lactobacillus catenaformis		4.25	0.15		99	
Eubacterium saburreum		3.15	0.21		85	
53270100	**Good identification**					**53270100**
Fusobacterium mortiferum		100.0	0.36	IND 1		
53271100	**Acceptable identification**					**53271100**
Fusobacterium mortiferum		92.61	0.39	IND 1		
Eubacterium saburreum		7.39	0.18			
53300000	**Good identification**					**53300000**
					SPO	
Clostridium bifermentans		98.76	0.48	SUC 1	99	
Clostridium botulinum biovar A		0.63	0.11	SAL 10	99	
Eubacterium saburreum		0.31	0.17		1	
53301000	**Low discrimination**					**53301000**
					SPO	
Eubacterium saburreum		63.89	0.51	SAL 1	1	
Eubacterium tortuosum		34.04	0.32		1	
Clostridium bifermentans		2.07	0.14		99	
53301002	**Low discrimination**					**53301002**
Eubacterium saburreum		65.24	0.39	SAL 1		
Eubacterium tortuosum		34.76	0.19			
53301010	**Good identification**					**53301010**
Eubacterium saburreum		99.35	0.50	SAL 1		
Eubacterium tortuosum		0.65	-.03			
53301012	**Good identification**					**53301012**
Eubacterium saburreum		100.0	0.37	SAL 1		
53301020	**Low discrimination**					**53301020**
Eubacterium saburreum		65.24	0.39	SAL 1		
Eubacterium tortuosum		34.76	0.19			
53301030	**Good identification**					**53301030**
Eubacterium saburreum		100.0	0.37	SAL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53301100	**Very good identification**					**53301100**
Eubacterium saburreum		99.77	0.58	SAL 1		
Eubacterium tortuosum		0.23	-.03			
53301102	**Good identification**					**53301102**
Eubacterium saburreum		100.0	0.45	SAL 1		
53301110	**Very good identification**					**53301110**
Eubacterium saburreum		100.0	0.56	SAL 1		
53301112	**Good identification**					**53301112**
Eubacterium saburreum		100.0	0.43	SAL 1		
53301120	**Good identification**					**53301120**
Eubacterium saburreum		100.0	0.45	SAL 1		
53301130	**Good identification**					**53301130**
Eubacterium saburreum		100.0	0.43	SAL 1		
53305000	**Good identification**					**53305000**
Eubacterium saburreum		99.01	0.47	SAL 1		
Eubacterium tortuosum		0.99	-.03			
53305010	**Good identification**					**53305010**
Eubacterium saburreum		100.0	0.45	SAL 1		
53305100	**Good identification**					**53305100**
Eubacterium saburreum		100.0	0.53	SAL 1		
53305102	**Good identification**					**53305102**
Eubacterium saburreum		100.0	0.40	SAL 1		
53305110	**Good identification**					**53305110**
Eubacterium saburreum		100.0	0.52	SAL 1		
53305112	**Good identification**					**53305112**
Eubacterium saburreum		100.0	0.39	SAL 1		
53305120	**Good identification**					**53305120**
Eubacterium saburreum		100.0	0.40	SAL 1		
53305130	**Good identification**					**53305130**
Eubacterium saburreum		100.0	0.39	SAL 1		
53310100	**Low discrimination**					**53310100**
Fusobacterium mortiferum		63.35	0.37	IND 1		
Eubacterium saburreum		36.65	0.32	MLT 11		
53311000	**Very good identification**					**53311000**
Eubacterium saburreum		96.70	0.60	SAL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		2.53	0.34			
Prevotella intermedia		0.47	0.11			
53311002	**Good identification**					**53311002**
Eubacterium saburreum		98.38	0.47	SAL 1		
Fusobacterium mortiferum		1.62	0.18			
53311010	**Very good identification**					**53311010**
Eubacterium saburreum		100.0	0.58	SAL 1		
53311012	**Good identification**					**53311012**
Eubacterium saburreum		100.0	0.45	SAL 1		
53311020	**Good identification**					**53311020**
Eubacterium saburreum		100.0	0.47	SAL 1		
53311030	**Good identification**					**53311030**
Eubacterium saburreum		100.0	0.45	SAL 1		
53311100	**Very good identification**					**53311100**
					GRA	
Eubacterium saburreum		97.23	0.66	SAL 1	85	
Fusobacterium mortiferum		2.55	0.40		1	
Collinsella aerofaciens		0.22	-.03		99	
53311102	**Good identification**					**53311102**
					GRA	
Eubacterium saburreum		97.01	0.53	SAL 1	85	
Fusobacterium mortiferum		1.60	0.24		1	
Bifidobacterium longum subsp. infantis		1.39	0.06		99	
53311110	**Very good identification**					**53311110**
Eubacterium saburreum		100.0	0.64	SAL 1		
53311112	**Good identification**					**53311112**
Eubacterium saburreum		100.0	0.52	SAL 1		
53311120	**Good identification**					**53311120**
					SPO	
Eubacterium saburreum		98.26	0.53	SAL 1	1	
Clostridium paraputrificum		0.70	-.03		99	
Clostridium perfringens		0.64	0.07		84	
53311122	**Low discrimination**					**53311122**
					GRA	
Eubacterium saburreum		80.54	0.40	SAL 1	85	
Bifidobacterium longum subsp. infantis		11.64	0.10		99	
Prevotella oralis		3.41	0.04		1	
53311130	**Good identification**					**53311130**
Eubacterium saburreum		100.0	0.52	SAL 1		
53311132	**Good identification**					**53311132**
Eubacterium saburreum		100.0	0.39	SAL 1		
53315000	**Good identification**					**53315000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.64	0.55	SAL 1		
Fusobacterium mortiferum		0.37	0.15			
53315002	**Good identification**					**53315002**
Eubacterium saburreum		100.0	0.42	SAL 1		
53315010	**Good identification**					**53315010**
Eubacterium saburreum		100.0	0.53	SAL 1		
53315012	**Good identification**					**53315012**
Eubacterium saburreum		100.0	0.41	SAL 1		
53315020	**Good identification**					**53315020**
Eubacterium saburreum		100.0	0.42	SAL 1		
53315030	**Good identification**					**53315030**
Eubacterium saburreum		100.0	0.41	SAL 1		
53315100	**Good identification**					**53315100**
					GRA	
Eubacterium saburreum		92.14	0.61	SAL 1	85	
Eubacterium contortum		7.53	0.32		99	
Fusobacterium mortiferum		0.34	0.21		1	
53315102	**Low discrimination**					**53315102**
Eubacterium saburreum		89.76	0.48	SAL 1		
Eubacterium contortum		7.33	0.19			
Bifidobacterium longum subsp. infantis		1.96	0.04			
53315110	**Very good identification**					**53315110**
Eubacterium saburreum		100.0	0.60	SAL 1		
53315112	**Good identification**					**53315112**
Eubacterium saburreum		98.44	0.47	SAL 1		
Bifidobacterium adolescentis		1.56	0.14			
53315120	**Acceptable identification**					**53315120**
Eubacterium saburreum		92.45	0.48	SAL 1		
Eubacterium contortum		7.55	0.19			
53315122	**Low discrimination**					**53315122**
Eubacterium saburreum		76.85	0.35	SAL 1		
Bifidobacterium longum subsp. infantis		16.87	0.08			
Eubacterium contortum		6.28	0.06			
53315130	**Good identification**					**53315130**
Eubacterium saburreum		100.0	0.47	SAL 1		
53315300	**Very good identification**					**53315300**
Eubacterium contortum		99.88	0.66	IND 1		
Eubacterium saburreum		0.13	0.27			
53315302	**Good identification**					**53315302**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum	99.88	0.53	IND 1		
Eubacterium saburreum	0.13	0.14			
53315320 Good identification					**53315320**
Eubacterium contortum	99.88	0.53	IND 1		
Eubacterium saburreum	0.13	0.14			
53315322 Good identification					**53315322**
Eubacterium contortum	100.0	0.40	IND 1		
53315700 Very good identification					**53315700**
Eubacterium contortum	100.0	0.64	IND 1		
53315702 Good identification					**53315702**
Eubacterium contortum	100.0	0.52	IND 1		
53315720 Good identification					**53315720**
				GRA	
Eubacterium contortum	99.86	0.52	IND 1	99	
Bacteroides vulgatus	0.14	-.05		1	
53315722 Good identification					**53315722**
				GRA CAT	
Eubacterium contortum	96.28	0.39	IND 1	99 1	
Bacteroides uniformis	3.27	0.06		1 5	
Bacteroides thetaiotaomicron	0.46	-.06		1 89	
53321000 Good identification					**53321000**
Eubacterium saburreum	99.20	0.48	SAL 1		
Eubacterium tortuosum	0.80	-.03			
53321002 Good identification					**53321002**
Eubacterium saburreum	100.0	0.36	SAL 1		
53321010 Good identification					**53321010**
Eubacterium saburreum	100.0	0.47	SAL 1		
53321020 Good identification					**53321020**
Eubacterium saburreum	100.0	0.36	SAL 1		
53321100 Good identification					**53321100**
Eubacterium saburreum	100.0	0.55	SAL 1		
53321102 Good identification					**53321102**
Eubacterium saburreum	100.0	0.42	SAL 1		
53321110 Good identification					**53321110**
Eubacterium saburreum	100.0	0.53	SAL 1		
53321112 Good identification					**53321112**
Eubacterium saburreum	100.0	0.40	SAL 1		
53321120 Good identification					**53321120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.42	SAL 1		
53321130	**Good identification**					**53321130**
Eubacterium saburreum		100.0	0.40	SAL 1		
53325000	**Good identification**					**53325000**
Eubacterium saburreum		100.0	0.44	SAL 1		
53325010	**Good identification**					**53325010**
Eubacterium saburreum		100.0	0.42	SAL 1		
53325100	**Good identification**					**53325100**
Eubacterium saburreum		100.0	0.50	SAL 1		
53325102	**Good identification**					**53325102**
Eubacterium saburreum		100.0	0.37	SAL 1		
53325110	**Good identification**					**53325110**
Eubacterium saburreum		100.0	0.49	SAL 1		
53325112	**Good identification**					**53325112**
Eubacterium saburreum		100.0	0.36	SAL 1		
53325120	**Good identification**					**53325120**
Eubacterium saburreum		100.0	0.37	SAL 1		
53325130	**Good identification**					**53325130**
Eubacterium saburreum		100.0	0.36	SAL 1		
53331000	**Very good identification**					**53331000**
					GRA	
Eubacterium saburreum		98.29	0.57	SAL 1	85	
Fusobacterium mortiferum		1.15	0.25		1	
Lactobacillus jensenii		0.56	0.10		99	
53331002	**Good identification**					**53331002**
Eubacterium saburreum		97.90	0.44	SAL 1		
Lactobacillus jensenii		2.10	0.07			
53331010	**Good identification**					**53331010**
Eubacterium saburreum		99.77	0.55	SAL 1		
Lactobacillus jensenii		0.23	0.02			
53331012	**Good identification**					**53331012**
Eubacterium saburreum		100.0	0.42	SAL 1		
53331020	**Good identification**					**53331020**
Eubacterium saburreum		100.0	0.44	SAL 1		
53331030	**Good identification**					**53331030**
Eubacterium saburreum		100.0	0.42	SAL 1		
53331100	**Very good identification**					**53331100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Eubacterium saburreum		97.59	0.63	SAL 1	1	85	
Atopobium parvulum		1.27	0.21		99	99	
Fusobacterium mortiferum		1.15	0.32		1	1	
53331102	**Good identification**						**53331102**
					COC	GRA	
Eubacterium saburreum		98.00	0.50	SAL 1	1	85	
Atopobium parvulum		1.27	0.08		99	99	
Fusobacterium mortiferum		0.72	0.15		1	1	
53331110	**Very good identification**						**53331110**
Eubacterium saburreum		100.0	0.61	SAL 1			
53331112	**Good identification**						**53331112**
Eubacterium saburreum		98.26	0.48	SAL 1			
Leptotrichia buccalis		1.74	-.03				
53331120	**Good identification**						**53331120**
					COC		
Eubacterium saburreum		96.98	0.50	SAL 1	1		
Clostridium perfringens		1.76	0.11		1		
Atopobium parvulum		1.26	0.08		99		
53331122	**Good identification**						**53331122**
Eubacterium saburreum		95.39	0.37	SAL 1			
Clostridium perfringens		4.61	0.06				
53331130	**Good identification**						**53331130**
Eubacterium saburreum		100.0	0.48	SAL 1			
53331132	**Good identification**						**53331132**
Eubacterium saburreum		100.0	0.36	SAL 1			
53335000	**Good identification**						**53335000**
Eubacterium saburreum		100.0	0.52	SAL 1			
53335002	**Good identification**						**53335002**
Eubacterium saburreum		100.0	0.39	SAL 1			
53335010	**Good identification**						**53335010**
Eubacterium saburreum		100.0	0.50	SAL 1			
53335012	**Good identification**						**53335012**
Eubacterium saburreum		100.0	0.38	SAL 1			
53335020	**Good identification**						**53335020**
Eubacterium saburreum		100.0	0.39	SAL 1			
53335030	**Good identification**						**53335030**
Eubacterium saburreum		100.0	0.38	SAL 1			
53335100	**Very good identification**						**53335100**
					COC	GRA	
Eubacterium saburreum		95.42	0.58	SAL 1	1	85	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		4.27	0.25		99	99	
Fusobacterium mortiferum		0.16	0.12		1	1	
53335102	**Acceptable identification**						**53335102**
					COC		
Eubacterium saburreum		93.58	0.45	SAL 1	1		
Atopobium parvulum		4.19	0.13		99		
Actinomyces israelii		1.24	0.17		1		
53335110	**Very good identification**						**53335110**
Eubacterium saburreum		100.0	0.57	SAL 1			
53335112	**Good identification**						**53335112**
Eubacterium saburreum		98.44	0.44	SAL 1			
Bifidobacterium adolescentis		1.56	0.11				
53335120	**Good identification**						**53335120**
					COC		
Eubacterium saburreum		95.71	0.45	SAL 1	1		
Atopobium parvulum		4.29	0.13		99		
53335130	**Good identification**						**53335130**
Eubacterium saburreum		100.0	0.44	SAL 1			
53355522	**Good identification**						**53355522**
Bacteroides uniformis		100.0	0.38	MNS 99			
53355722	**Good identification**						**53355722**
					GRA		
Bacteroides uniformis		97.27	0.40	MNS 99	1		
Bacteroides ovatus		2.14	0.15	RAF 97	1		
Eubacterium contortum		0.29	0.04		99		
53357100	**Good identification**						**53357100**
Actinomyces israelii		100.0	0.37	IND 1			
53357102	**Low discrimination**						**53357102**
					SPO		
Actinomyces israelii		86.57	0.40	IND 1	1		
Bifidobacterium adolescentis		9.08	0.16		1		
Clostridium tertium		4.35	0.03		99		
53357500	**Good identification**						**53357500**
Actinomyces israelii		100.0	0.35	IND 1			
53357502	**Low discrimination**						**53357502**
Actinomyces israelii		53.83	0.38	IND 1			
Bifidobacterium adolescentis		46.18	0.31				
53375102	**Low discrimination**						**53375102**
					COC		
Actinomyces israelii		54.22	0.36	IND 1	1		
Atopobium parvulum		32.30	0.19		99		
Bifidobacterium adolescentis		6.98	0.15		1		
53377000	**Good identification**						**53377000**
Actinomyces israelii		100.0	0.36	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53377002	**Good identification**					**53377002**
Actinomyces israelii		100.0	0.39	IND 1		
53377100	**Acceptable identification**					**53377100**
					SPO	
Actinomyces israelii		94.33	0.46	IND 1	1	
Clostridium tertium		5.68	0.11		99	
53377102	**Low discrimination**					**53377102**
					SPO	
Actinomyces israelii		87.80	0.49	IND 1	1	
Clostridium tertium		10.57	0.19		99	
Bifidobacterium adolescentis		1.63	0.13		1	
53377140	**Good identification**					**53377140**
Actinomyces israelii		100.0	0.39	IND 1		
53377142	**Good identification**					**53377142**
Actinomyces israelii		100.0	0.42	IND 1		
53377402	**Good identification**					**53377402**
Actinomyces israelii		100.0	0.38	IND 1		
53377500	**Good identification**					**53377500**
					SPO	
Actinomyces israelii		95.08	0.45	IND 1	1	
Clostridium butyricum		4.92	0.02		99	
53377502	**Low discrimination**					**53377502**
					SPO	
Actinomyces israelii		85.80	0.48	IND 1	1	
Bifidobacterium adolescentis		13.05	0.28		1	
Clostridium butyricum		1.15	-.05		99	
53377540	**Good identification**					**53377540**
Actinomyces israelii		100.0	0.38	IND 1		
53377542	**Good identification**					**53377542**
Actinomyces israelii		95.39	0.41	IND 1		
Bifidobacterium adolescentis		4.61	0.12			
53400000	**Low discrimination**					**53400000**
					SPO	
Eubacterium saburreum		46.65	0.37	GAL 99	1	
Clostridium bifermentans		43.27	0.24		99	
Clostridium botulinum biovar A		8.61	0.13		99	
53400010	**Good identification**					**53400010**
Eubacterium saburreum		100.0	0.35	GAL 99		
53400100	**Good identification**					**53400100**
Eubacterium saburreum		96.92	0.43	GAL 99		
Fusobacterium mortiferum		3.08	0.19			
53400110	**Good identification**					**53400110**
Eubacterium saburreum		100.0	0.41	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53401000	**Very good identification**						**53401000**
					GRA		
Eubacterium saburreum		98.64	0.71		85		
Eubacterium tortuosum		1.32	0.24		99		
Fusobacterium mortiferum		0.05	0.16		1		
53401001	**Good identification**						**53401001**
Eubacterium saburreum		100.0	0.37	URE 1			
53401002	**Very good identification**						**53401002**
Eubacterium saburreum		98.69	0.58				
Eubacterium tortuosum		1.32	0.11				
53401010	**Very good identification**						**53401010**
Eubacterium saburreum		100.0	0.69				
53401011	**Good identification**						**53401011**
Eubacterium saburreum		100.0	0.35	URE 1			
53401012	**Very good identification**						**53401012**
Eubacterium saburreum		100.0	0.56				
53401020	**Very good identification**						**53401020**
Eubacterium saburreum		98.69	0.58				
Eubacterium tortuosum		1.32	0.11				
53401022	**Good identification**						**53401022**
Eubacterium saburreum		98.69	0.45				
Eubacterium tortuosum		1.32	-.02				
53401030	**Very good identification**						**53401030**
Eubacterium saburreum		100.0	0.56				
53401032	**Good identification**						**53401032**
Eubacterium saburreum		100.0	0.43				
53401040	**Good identification**						**53401040**
Eubacterium saburreum		100.0	0.37	SOR 1			
53401050	**Good identification**						**53401050**
Eubacterium saburreum		100.0	0.35	SOR 1			
53401100	**Very good identification**						**53401100**
					COC	GRA	
Eubacterium saburreum		99.93	0.77		1	85	
Fusobacterium mortiferum		0.05	0.22		1	1	
Blautia hansenii		0.02	-.03		99	99	
53401101	**Good identification**						**53401101**
Eubacterium saburreum		100.0	0.43	URE 1			
53401102	**Very good identification**						**53401102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.64			
53401110	**Very good identification**					**53401110**
Eubacterium saburreum		100.0	0.75			
53401111	**Good identification**					**53401111**
Eubacterium saburreum		100.0	0.41	URE 1		
53401112	**Very good identification**					**53401112**
Eubacterium saburreum		100.0	0.63			
53401120	**Very good identification**					**53401120**
Eubacterium saburreum		99.91	0.64			
Prevotella melaninogenica		0.09	-.04			
53401122	**Good identification**					**53401122**
Eubacterium saburreum		100.0	0.51			
53401130	**Very good identification**					**53401130**
Eubacterium saburreum		100.0	0.63			
53401132	**Good identification**					**53401132**
Eubacterium saburreum		100.0	0.50			
53401140	**Good identification**					**53401140**
Eubacterium saburreum		100.0	0.43	SOR 1		
53401150	**Good identification**					**53401150**
Eubacterium saburreum		100.0	0.41	SOR 1		
53401200	**Good identification**					**53401200**
Eubacterium saburreum		100.0	0.37	RHA 1		
53401210	**Good identification**					**53401210**
Eubacterium saburreum		100.0	0.35	RHA 1		
53401300	**Good identification**					**53401300**
Eubacterium saburreum		100.0	0.43	RHA 1		
53401310	**Good identification**					**53401310**
Eubacterium saburreum		100.0	0.41	RHA 1		
53401400	**Good identification**					**53401400**
Eubacterium saburreum		100.0	0.37	ARA 1		
53401410	**Good identification**					**53401410**
Eubacterium saburreum		100.0	0.35	ARA 1		
53401500	**Good identification**					**53401500**
Eubacterium saburreum		100.0	0.43	ARA 1		
53401510	**Good identification**					**53401510**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.41	ARA 1		
53403000	**Good identification**					**53403000**
Eubacterium saburreum		100.0	0.37	MAN 1		
53403010	**Good identification**					**53403010**
Eubacterium saburreum		100.0	0.35	MAN 1		
53403100	**Good identification**					**53403100**
Eubacterium saburreum		100.0	0.43	MAN 1		
53403110	**Good identification**					**53403110**
Eubacterium saburreum		100.0	0.41	MAN 1		
53404100	**Good identification**					**53404100**
Eubacterium saburreum		100.0	0.38	GAL 99		
53404110	**Good identification**					**53404110**
Eubacterium saburreum		100.0	0.37	GAL 99		
53405000	**Very good identification**					**53405000**
Eubacterium saburreum		100.0	0.66			
53405002	**Good identification**					**53405002**
Eubacterium saburreum		100.0	0.53			
53405010	**Very good identification**					**53405010**
Eubacterium saburreum		100.0	0.65			
53405012	**Good identification**					**53405012**
Eubacterium saburreum		100.0	0.52			
53405020	**Good identification**					**53405020**
Eubacterium saburreum		100.0	0.53			
53405022	**Good identification**					**53405022**
Eubacterium saburreum		100.0	0.40			
53405030	**Good identification**					**53405030**
Eubacterium saburreum		100.0	0.52			
53405032	**Good identification**					**53405032**
Eubacterium saburreum		100.0	0.39			
53405100	**Very good identification**					**53405100**
Eubacterium saburreum		100.0	0.72			
53405101	**Good identification**					**53405101**
Eubacterium saburreum		100.0	0.38	URE 1		
53405102	**Very good identification**					**53405102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.59			
53405110	**Very good identification**					**53405110**
Eubacterium saburreum		100.0	0.71			
53405111	**Good identification**					**53405111**
Eubacterium saburreum		100.0	0.37	URE 1		
53405112	**Very good identification**					**53405112**
Eubacterium saburreum		100.0	0.58			
53405120	**Very good identification**					**53405120**
Eubacterium saburreum		100.0	0.59			
53405122	**Good identification**					**53405122**
Eubacterium saburreum		100.0	0.47			
53405130	**Very good identification**					**53405130**
Eubacterium saburreum		100.0	0.58			
53405132	**Good identification**					**53405132**
Eubacterium saburreum		100.0	0.45			
53405140	**Good identification**					**53405140**
Eubacterium saburreum		100.0	0.38	SOR 1		
53405150	**Good identification**					**53405150**
Eubacterium saburreum		100.0	0.37	SOR 1		
53405300	**Good identification**					**53405300**
Eubacterium saburreum		100.0	0.38	RHA 1		
53405310	**Good identification**					**53405310**
Eubacterium saburreum		100.0	0.37	RHA 1		
53405500	**Good identification**					**53405500**
Eubacterium saburreum		100.0	0.38	ARA 1		
53405510	**Good identification**					**53405510**
Eubacterium saburreum		100.0	0.37	ARA 1		
53407100	**Good identification**					**53407100**
Eubacterium saburreum		100.0	0.38	MAN 1		
53407110	**Good identification**					**53407110**
Eubacterium saburreum		100.0	0.37	MAN 1		
53410000	**Low discrimination**					**53410000**
Fusobacterium mortiferum		49.50	0.47	IND 1		
Eubacterium saburreum		47.16	0.45			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Prevotella intermedia	2.57	0.14			
53410010	**Good identification**					**53410010**
	Eubacterium saburreum	98.72	0.43	GAL 99		
	Fusobacterium mortiferum	1.28	0.13			
53410100	**Low discrimination**					**53410100**
	Fusobacterium mortiferum	51.21	0.53	IND 1		
	Eubacterium saburreum	48.79	0.51			
53410102	**Low discrimination**					**53410102**
	Eubacterium saburreum	60.21	0.38	GAL 99		
	Fusobacterium mortiferum	39.79	0.37			
53410110	**Good identification**					**53410110**
	Eubacterium saburreum	98.72	0.49	GAL 99		
	Fusobacterium mortiferum	1.28	0.19			
53410112	**Good identification**					**53410112**
	Eubacterium saburreum	100.0	0.37	GAL 99		
53410120	**Acceptable identification**					**53410120**
	Eubacterium saburreum	94.33	0.38	GAL 99		
	Fusobacterium mortiferum	5.67	0.19			
53410130	**Good identification**					**53410130**
	Eubacterium saburreum	100.0	0.37	GAL 99		
53411000	**Very good identification**					**53411000**
	Eubacterium saburreum	98.36	0.79			
	Fusobacterium mortiferum	1.56	0.50			
	Prevotella intermedia	0.05	0.14			
53411001	**Good identification**					**53411001**
	Eubacterium saburreum	98.43	0.45	URE 1		
	Fusobacterium mortiferum	1.57	0.16			
53411002	**Very good identification**					**53411002**
					GRA	
	Eubacterium saburreum	98.88	0.66		85	
	Fusobacterium mortiferum	0.99	0.33		1	
	Lactobacillus fermentum	0.13	0.20		99	
53411010	**Very good identification**					**53411010**
	Eubacterium saburreum	99.98	0.77			
	Fusobacterium mortiferum	0.02	0.16			
53411011	**Good identification**					**53411011**
	Eubacterium saburreum	100.0	0.43	URE 1		
53411012	**Very good identification**					**53411012**
	Eubacterium saburreum	100.0	0.64			
53411020	**Very good identification**					**53411020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.91	0.66			
Fusobacterium mortiferum		0.09	0.16			
53411022	**Good identification**					**53411022**
Eubacterium saburreum		100.0	0.53			
53411030	**Very good identification**					**53411030**
Eubacterium saburreum		100.0	0.64			
53411032	**Good identification**					**53411032**
Eubacterium saburreum		100.0	0.52			
53411040	**Good identification**					**53411040**
Eubacterium saburreum		98.43	0.45	SOR 1		
Fusobacterium mortiferum		1.57	0.16			
53411050	**Good identification**					**53411050**
Eubacterium saburreum		100.0	0.43	SOR 1		
53411100	**Very good identification**					**53411100**
					GRA	
Eubacterium saburreum		98.42	0.85		85	
Fusobacterium mortiferum		1.57	0.56		1	
Lactobacillus fermentum		0.02	0.24		99	
53411101	**Good identification**					**53411101**
Eubacterium saburreum		98.44	0.51	URE 1		
Fusobacterium mortiferum		1.57	0.22			
53411102	**Very good identification**					**53411102**
					GRA	
Eubacterium saburreum		98.92	0.72		85	
Fusobacterium mortiferum		0.99	0.40		1	
Lactobacillus fermentum		0.09	0.24		99	
53411103	**Good identification**					**53411103**
Eubacterium saburreum		100.0	0.38	URE 1		
53411110	**Very good identification**					**53411110**
Eubacterium saburreum		99.98	0.84			
Fusobacterium mortiferum		0.02	0.22			
53411111	**Good identification**					**53411111**
Eubacterium saburreum		100.0	0.49	URE 1		
53411112	**Very good identification**					**53411112**
Eubacterium saburreum		100.0	0.71			
53411113	**Good identification**					**53411113**
Eubacterium saburreum		100.0	0.37	URE 1		
53411120	**Very good identification**					**53411120**
Eubacterium saburreum		99.61	0.72			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella melaninogenica		0.27	0.12			
Fusobacterium mortiferum		0.09	0.22			
53411121	**Good identification**					**53411121**
Eubacterium saburreum		100.0	0.38	URE 1		
53411122	**Very good identification**					**53411122**
					CAT	
Eubacterium saburreum		98.63	0.59		1	
Capnocytophaga ochracea		0.91	0.12		90	
Prevotella melaninogenica		0.29	0.00		1	
53411130	**Very good identification**					**53411130**
Eubacterium saburreum		100.0	0.71			
53411131	**Good identification**					**53411131**
Eubacterium saburreum		100.0	0.37	URE 1		
53411132	**Very good identification**					**53411132**
Eubacterium saburreum		100.0	0.58			
53411140	**Good identification**					**53411140**
Eubacterium saburreum		98.44	0.51	SOR 1		
Fusobacterium mortiferum		1.57	0.22			
53411142	**Good identification**					**53411142**
Eubacterium saburreum		100.0	0.38	SOR 1		
53411150	**Good identification**					**53411150**
Eubacterium saburreum		100.0	0.49	SOR 1		
53411152	**Good identification**					**53411152**
Eubacterium saburreum		100.0	0.37	SOR 1		
53411160	**Good identification**					**53411160**
Eubacterium saburreum		100.0	0.38	SOR 1		
53411170	**Good identification**					**53411170**
Eubacterium saburreum		100.0	0.37	SOR 1		
53411200	**Good identification**					**53411200**
Eubacterium saburreum		98.43	0.45	RHA 1		
Fusobacterium mortiferum		1.57	0.16			
53411210	**Good identification**					**53411210**
Eubacterium saburreum		100.0	0.43	RHA 1		
53411300	**Good identification**					**53411300**
Eubacterium saburreum		98.44	0.51	RHA 1		
Fusobacterium mortiferum		1.57	0.22			
53411302	**Good identification**					**53411302**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	RHA 1		
53411310	**Good identification**					**53411310**
Eubacterium saburreum		100.0	0.49	RHA 1		
53411312	**Good identification**					**53411312**
Eubacterium saburreum		100.0	0.37	RHA 1		
53411320	**Good identification**					**53411320**
Eubacterium saburreum		100.0	0.38	RHA 1		
53411330	**Good identification**					**53411330**
Eubacterium saburreum		100.0	0.37	RHA 1		
53411400	**Good identification**					**53411400**
					GRA	
Eubacterium saburreum		96.23	0.45	ARA 1	85	
Lactobacillus fermentum		2.24	0.20		99	
Fusobacterium mortiferum		1.53	0.16		1	
53411410	**Good identification**					**53411410**
Eubacterium saburreum		100.0	0.43	ARA 1		
53411500	**Good identification**					**53411500**
					GRA	
Eubacterium saburreum		96.88	0.51	ARA 1	85	
Lactobacillus fermentum		1.58	0.24		99	
Fusobacterium mortiferum		1.54	0.22		1	
53411502	**Acceptable identification**					**53411502**
Eubacterium saburreum		91.56	0.38	ARA 1		
Lactobacillus fermentum		8.44	0.24			
53411510	**Good identification**					**53411510**
Eubacterium saburreum		100.0	0.49	ARA 1		
53411512	**Good identification**					**53411512**
Eubacterium saburreum		100.0	0.37	ARA 1		
53411520	**Acceptable identification**					**53411520**
					GRA	
Eubacterium saburreum		90.34	0.38	ARA 1	85	
Odoribacter splanchnicus		5.00	-.03		1	
Lactobacillus fermentum		2.78	0.15		99	
53411530	**Good identification**					**53411530**
Eubacterium saburreum		95.83	0.37	ARA 1		
Bifidobacterium longum subsp. longum		4.17	0.09			
53413000	**Good identification**					**53413000**
Eubacterium saburreum		98.43	0.45	MAN 1		
Fusobacterium mortiferum		1.57	0.16			
53413010	**Good identification**					**53413010**
Eubacterium saburreum		100.0	0.43	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53413100	**Good identification**					**53413100**
Eubacterium saburreum		98.44	0.51	MAN 1		
Fusobacterium mortiferum		1.57	0.22			
53413102	**Good identification**					**53413102**
Eubacterium saburreum		100.0	0.38	MAN 1		
53413110	**Good identification**					**53413110**
Eubacterium saburreum		100.0	0.49	MAN 1		
53413112	**Good identification**					**53413112**
Eubacterium saburreum		100.0	0.37	MAN 1		
53413120	**Good identification**					**53413120**
Eubacterium saburreum		100.0	0.38	MAN 1		
53413130	**Good identification**					**53413130**
Eubacterium saburreum		100.0	0.37	MAN 1		
53414000	**Low discrimination**					**53414000**
Eubacterium saburreum		87.21	0.40	GAL 99		
Fusobacterium mortiferum		12.80	0.27			
53414010	**Good identification**					**53414010**
Eubacterium saburreum		100.0	0.39	GAL 99		
53414100	**Low discrimination**					**53414100**
Eubacterium saburreum		87.21	0.46	GAL 99		
Fusobacterium mortiferum		12.79	0.34			
53414110	**Good identification**					**53414110**
Eubacterium saburreum		100.0	0.45	GAL 99		
53415000	**Very good identification**					**53415000**
					GRA	
Eubacterium saburreum		99.75	0.74		85	
Fusobacterium mortiferum		0.22	0.30		1	
Lactobacillus fermentum		0.03	0.16		99	
53415001	**Good identification**					**53415001**
Eubacterium saburreum		100.0	0.40	URE 1		
53415002	**Very good identification**					**53415002**
					GRA	
Eubacterium saburreum		99.72	0.61		85	
Lactobacillus fermentum		0.15	0.16		99	
Fusobacterium mortiferum		0.14	0.14		1	
53415010	**Very good identification**					**53415010**
Eubacterium saburreum		100.0	0.73			
53415011	**Good identification**					**53415011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.39	URE 1		
53415012	**Very good identification**					**53415012**
Eubacterium saburreum		100.0	0.60			
53415020	**Very good identification**					**53415020**
Eubacterium saburreum		100.0	0.61			
53415022	**Good identification**					**53415022**
Eubacterium saburreum		100.0	0.48			
53415030	**Very good identification**					**53415030**
Eubacterium saburreum		100.0	0.60			
53415032	**Good identification**					**53415032**
Eubacterium saburreum		100.0	0.47			
53415040	**Good identification**					**53415040**
Eubacterium saburreum		100.0	0.40	SOR 1		
53415050	**Good identification**					**53415050**
Eubacterium saburreum		100.0	0.39	SOR 1		
53415100	**Very good identification**					**53415100**
					GRA	
Eubacterium saburreum		99.76	0.81		85	
Fusobacterium mortiferum		0.22	0.37		1	
Lactobacillus fermentum		0.02	0.20		99	
53415101	**Good identification**					**53415101**
Eubacterium saburreum		100.0	0.46	URE 1		
53415102	**Very good identification**					**53415102**
					GRA	
Eubacterium saburreum		99.76	0.68		85	
Fusobacterium mortiferum		0.14	0.20		1	
Lactobacillus fermentum		0.10	0.20		99	
53415110	**Very good identification**					**53415110**
Eubacterium saburreum		100.0	0.79			
53415111	**Good identification**					**53415111**
Eubacterium saburreum		100.0	0.45	URE 1		
53415112	**Very good identification**					**53415112**
Eubacterium saburreum		100.0	0.66			
53415120	**Very good identification**					**53415120**
Eubacterium saburreum		100.0	0.68			
53415122	**Good identification**					**53415122**
					CAT	
Eubacterium saburreum		98.78	0.55		1	
Bacteroides fragilis		0.80	-.03		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Capnocytophaga ochracea		0.42	0.02		90	
53415130	**Very good identification**					**53415130**
Eubacterium saburreum		100.0	0.66			
53415132	**Good identification**					**53415132**
Eubacterium saburreum		100.0	0.53			
53415140	**Good identification**					**53415140**
Eubacterium saburreum		100.0	0.46	SOR 1		
53415150	**Good identification**					**53415150**
Eubacterium saburreum		100.0	0.45	SOR 1		
53415200	**Good identification**					**53415200**
Eubacterium saburreum		100.0	0.40	RHA 1		
53415210	**Good identification**					**53415210**
Eubacterium saburreum		100.0	0.39	RHA 1		
53415300	**Good identification**					**53415300**
Eubacterium saburreum		100.0	0.46	RHA 1		
53415310	**Good identification**					**53415310**
Eubacterium saburreum		100.0	0.45	RHA 1		
53415400	**Good identification**					**53415400**
Eubacterium saburreum		97.53	0.40	ARA 1		
Lactobacillus fermentum		2.47	0.16			
53415410	**Good identification**					**53415410**
Eubacterium saburreum		100.0	0.39	ARA 1		
53415500	**Good identification**					**53415500**
Eubacterium saburreum		98.26	0.46	ARA 1		
Lactobacillus fermentum		1.74	0.20			
53415510	**Good identification**					**53415510**
Eubacterium saburreum		98.94	0.45	ARA 1		
Bifidobacterium longum subsp. longum		1.06	0.07			
53417000	**Good identification**					**53417000**
Eubacterium saburreum		100.0	0.40	MAN 1		
53417010	**Good identification**					**53417010**
Eubacterium saburreum		100.0	0.39	MAN 1		
53417100	**Good identification**					**53417100**
Eubacterium saburreum		100.0	0.46	MAN 1		
53417110	**Good identification**					**53417110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.45	MAN 1		
53420100	**Good identification**					**53420100**
Eubacterium saburreum		100.0	0.40	GAL 99		
53420110	**Good identification**					**53420110**
Eubacterium saburreum		100.0	0.38	GAL 99		
53421000	**Very good identification**					**53421000**
Eubacterium saburreum		100.0	0.68			
53421002	**Good identification**					**53421002**
Eubacterium saburreum		100.0	0.55			
53421010	**Very good identification**					**53421010**
Eubacterium saburreum		100.0	0.66			
53421012	**Good identification**					**53421012**
Eubacterium saburreum		100.0	0.53			
53421020	**Good identification**					**53421020**
Eubacterium saburreum		100.0	0.55			
53421022	**Good identification**					**53421022**
Eubacterium saburreum		100.0	0.42			
53421030	**Good identification**					**53421030**
Eubacterium saburreum		100.0	0.53			
53421032	**Good identification**					**53421032**
Eubacterium saburreum		100.0	0.40			
53421100	**Very good identification**					**53421100**
Eubacterium saburreum		99.98	0.74			
Fusobacterium mortiferum		0.02	0.13			
53421101	**Good identification**					**53421101**
Eubacterium saburreum		100.0	0.40	URE 1		
53421102	**Very good identification**					**53421102**
Eubacterium saburreum		100.0	0.61			
53421110	**Very good identification**					**53421110**
Eubacterium saburreum		100.0	0.72			
53421111	**Good identification**					**53421111**
Eubacterium saburreum		100.0	0.38	URE 1		
53421112	**Very good identification**					**53421112**
Eubacterium saburreum		100.0	0.60			
53421120	**Very good identification**					**53421120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.61			
53421122	**Good identification**					**53421122**
Eubacterium saburreum		100.0	0.48			
53421130	**Very good identification**					**53421130**
Eubacterium saburreum		100.0	0.60			
53421132	**Good identification**					**53421132**
Eubacterium saburreum		100.0	0.47			
53421140	**Good identification**					**53421140**
Eubacterium saburreum		100.0	0.40	SOR 1		
53421150	**Good identification**					**53421150**
Eubacterium saburreum		100.0	0.38	SOR 1		
53421300	**Good identification**					**53421300**
Eubacterium saburreum		100.0	0.40	RHA 1		
53421310	**Good identification**					**53421310**
Eubacterium saburreum		100.0	0.38	RHA 1		
53421500	**Good identification**					**53421500**
Eubacterium saburreum		100.0	0.40	ARA 1		
53421510	**Good identification**					**53421510**
Eubacterium saburreum		100.0	0.38	ARA 1		
53423100	**Good identification**					**53423100**
Eubacterium saburreum		100.0	0.40	MAN 1		
53423110	**Good identification**					**53423110**
Eubacterium saburreum		100.0	0.38	MAN 1		
53424100	**Good identification**					**53424100**
Eubacterium saburreum		100.0	0.35	GAL 99		
53425000	**Very good identification**					**53425000**
Eubacterium saburreum		100.0	0.63			
53425002	**Good identification**					**53425002**
Eubacterium saburreum		100.0	0.50			
53425010	**Very good identification**					**53425010**
Eubacterium saburreum		100.0	0.62			
53425012	**Good identification**					**53425012**
Eubacterium saburreum		100.0	0.49			
53425020	**Good identification**					**53425020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.50			
53425022	**Good identification**					**53425022**
Eubacterium saburreum		100.0	0.37			
53425030	**Good identification**					**53425030**
Eubacterium saburreum		100.0	0.49			
53425032	**Good identification**					**53425032**
Eubacterium saburreum		100.0	0.36			
53425100	**Very good identification**					**53425100**
Eubacterium saburreum		100.0	0.69			
53425101	**Good identification**					**53425101**
Eubacterium saburreum		100.0	0.35	URE 1		
53425102	**Very good identification**					**53425102**
Eubacterium saburreum		100.0	0.56			
53425110	**Very good identification**					**53425110**
Eubacterium saburreum		100.0	0.68			
53425112	**Good identification**					**53425112**
Eubacterium saburreum		100.0	0.55			
53425120	**Very good identification**					**53425120**
Eubacterium saburreum		100.0	0.56			
53425122	**Good identification**					**53425122**
Eubacterium saburreum		100.0	0.44			
53425130	**Good identification**					**53425130**
Eubacterium saburreum		100.0	0.55			
53425132	**Good identification**					**53425132**
Eubacterium saburreum		100.0	0.42			
53425140	**Good identification**					**53425140**
Eubacterium saburreum		100.0	0.35	SOR 1		
53425300	**Good identification**					**53425300**
Eubacterium saburreum		100.0	0.35	RHA 1		
53425500	**Good identification**					**53425500**
Eubacterium saburreum		100.0	0.35	ARA 1		
53427100	**Good identification**					**53427100**
Eubacterium saburreum		100.0	0.35	MAN 1		
53430000	**Low discrimination**					**53430000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		68.02	0.42	GAL 99		
Fusobacterium mortiferum		31.99	0.38			
53430010	**Good identification**					**53430010**
Eubacterium saburreum		100.0	0.40	GAL 99		
53430100	**Low discrimination**					**53430100**
Eubacterium saburreum		68.02	0.48	GAL 99		
Fusobacterium mortiferum		31.99	0.44			
53430102	**Low discrimination**					**53430102**
Eubacterium saburreum		77.16	0.35	GAL 99		
Fusobacterium mortiferum		22.85	0.28			
53430110	**Good identification**					**53430110**
Eubacterium saburreum		100.0	0.46	GAL 99		
53430120	**Good identification**					**53430120**
Eubacterium saburreum		100.0	0.35	GAL 99		
53431000	**Very good identification**					**53431000**
Eubacterium saburreum		99.29	0.76			
Fusobacterium mortiferum		0.71	0.41			
53431001	**Good identification**					**53431001**
Eubacterium saburreum		100.0	0.42	URE 1		
53431002	**Very good identification**					**53431002**
Eubacterium saburreum		99.55	0.63			
Fusobacterium mortiferum		0.45	0.24			
53431010	**Very good identification**					**53431010**
Eubacterium saburreum		100.0	0.74			
53431011	**Good identification**					**53431011**
Eubacterium saburreum		100.0	0.40	URE 1		
53431012	**Very good identification**					**53431012**
Eubacterium saburreum		100.0	0.61			
53431020	**Very good identification**					**53431020**
Eubacterium saburreum		100.0	0.63			
53431022	**Good identification**					**53431022**
Eubacterium saburreum		100.0	0.50			
53431030	**Very good identification**					**53431030**
Eubacterium saburreum		100.0	0.61			
53431032	**Good identification**					**53431032**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.49			
53431040	**Good identification**					**53431040**
Eubacterium saburreum		100.0	0.42	SOR 1		
53431050	**Good identification**					**53431050**
Eubacterium saburreum		100.0	0.40	SOR 1		
53431100	**Very good identification**					**53431100**
Eubacterium saburreum		99.29	0.82			
Fusobacterium mortiferum		0.71	0.47			
53431101	**Good identification**					**53431101**
Eubacterium saburreum		99.29	0.48	URE 1		
Fusobacterium mortiferum		0.71	0.13			
53431102	**Very good identification**					**53431102**
Eubacterium saburreum		99.55	0.69			
Fusobacterium mortiferum		0.45	0.31			
53431103	**Good identification**					**53431103**
Eubacterium saburreum		100.0	0.35	URE 1		
53431110	**Very good identification**					**53431110**
Eubacterium saburreum		99.99	0.81			
Fusobacterium mortiferum		0.01	0.13			
53431111	**Good identification**					**53431111**
Eubacterium saburreum		100.0	0.46	URE 1		
53431112	**Very good identification**					**53431112**
Eubacterium saburreum		100.0	0.68			
53431120	**Very good identification**					**53431120**
Eubacterium saburreum		99.96	0.69			
Fusobacterium mortiferum		0.04	0.13			
53431121	**Good identification**					**53431121**
Eubacterium saburreum		100.0	0.35	URE 1		
53431122	**Very good identification**					**53431122**
Eubacterium saburreum		100.0	0.56			
53431130	**Very good identification**					**53431130**
Eubacterium saburreum		100.0	0.68			
53431132	**Good identification**					**53431132**
Eubacterium saburreum		100.0	0.55			
53431140	**Good identification**					**53431140**
Eubacterium saburreum		99.29	0.48	SOR 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		0.71	0.13			
53431142	**Good identification**					**53431142**
Eubacterium saburreum		100.0	0.35	SOR 1		
53431150	**Good identification**					**53431150**
Eubacterium saburreum		100.0	0.46	SOR 1		
53431160	**Good identification**					**53431160**
Eubacterium saburreum		100.0	0.35	SOR 1		
53431200	**Good identification**					**53431200**
Eubacterium saburreum		100.0	0.42	RHA 1		
53431210	**Good identification**					**53431210**
Eubacterium saburreum		100.0	0.40	RHA 1		
53431300	**Good identification**					**53431300**
Eubacterium saburreum		99.29	0.48	RHA 1		
Fusobacterium mortiferum		0.71	0.13			
53431302	**Good identification**					**53431302**
Eubacterium saburreum		100.0	0.35	RHA 1		
53431310	**Good identification**					**53431310**
Eubacterium saburreum		100.0	0.46	RHA 1		
53431320	**Good identification**					**53431320**
Eubacterium saburreum		100.0	0.35	RHA 1		
53431400	**Good identification**					**53431400**
Eubacterium saburreum		100.0	0.42	ARA 1		
53431410	**Good identification**					**53431410**
Eubacterium saburreum		100.0	0.40	ARA 1		
53431500	**Good identification**					**53431500**
Eubacterium saburreum		99.29	0.48	ARA 1		
Fusobacterium mortiferum		0.71	0.13			
53431502	**Good identification**					**53431502**
Eubacterium saburreum		100.0	0.35	ARA 1		
53431510	**Good identification**					**53431510**
Eubacterium saburreum		100.0	0.46	ARA 1		
53431520	**Good identification**					**53431520**
Eubacterium saburreum		100.0	0.35	ARA 1		
53433000	**Good identification**					**53433000**
Eubacterium saburreum		100.0	0.42	MAN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53433010	**Good identification**					**53433010**
Eubacterium saburreum		100.0	0.40	MAN 1		
53433100	**Good identification**					**53433100**
Eubacterium saburreum		99.29	0.48	MAN 1		
Fusobacterium mortiferum		0.71	0.13			
53433102	**Good identification**					**53433102**
Eubacterium saburreum		100.0	0.35	MAN 1		
53433110	**Good identification**					**53433110**
Eubacterium saburreum		100.0	0.46	MAN 1		
53433120	**Good identification**					**53433120**
Eubacterium saburreum		100.0	0.35	MAN 1		
53434000	**Acceptable identification**					**53434000**
Eubacterium saburreum		93.83	0.37	GAL 99		
Fusobacterium mortiferum		6.17	0.19			
53434010	**Good identification**					**53434010**
Eubacterium saburreum		100.0	0.36	GAL 99		
53434100	**Acceptable identification**					**53434100**
Eubacterium saburreum		93.83	0.43	GAL 99		
Fusobacterium mortiferum		6.17	0.25			
53434110	**Good identification**					**53434110**
Eubacterium saburreum		100.0	0.42	GAL 99		
53435000	**Very good identification**					**53435000**
Eubacterium saburreum		99.90	0.71			
Fusobacterium mortiferum		0.10	0.22			
53435001	**Good identification**					**53435001**
Eubacterium saburreum		100.0	0.37	URE 1		
53435002	**Very good identification**					**53435002**
Eubacterium saburreum		100.0	0.58			
53435010	**Very good identification**					**53435010**
Eubacterium saburreum		100.0	0.70			
53435011	**Good identification**					**53435011**
Eubacterium saburreum		100.0	0.36	URE 1		
53435012	**Very good identification**					**53435012**
Eubacterium saburreum		100.0	0.57			
53435020	**Very good identification**					**53435020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.58			
53435022	**Good identification**					**53435022**
Eubacterium saburreum		100.0	0.45			
53435030	**Very good identification**					**53435030**
Eubacterium saburreum		100.0	0.57			
53435032	**Good identification**					**53435032**
Eubacterium saburreum		100.0	0.44			
53435040	**Good identification**					**53435040**
Eubacterium saburreum		100.0	0.37	SOR 1		
53435050	**Good identification**					**53435050**
Eubacterium saburreum		100.0	0.36	SOR 1		
53435100	**Very good identification**					**53435100**
Eubacterium saburreum		99.90	0.78			
Fusobacterium mortiferum		0.10	0.28			
53435101	**Good identification**					**53435101**
Eubacterium saburreum		100.0	0.43	URE 1		
53435102	**Very good identification**					**53435102**
Eubacterium saburreum		99.94	0.65			
Fusobacterium mortiferum		0.06	0.11			
53435110	**Very good identification**					**53435110**
Eubacterium saburreum		100.0	0.76			
53435111	**Good identification**					**53435111**
Eubacterium saburreum		100.0	0.42	URE 1		
53435112	**Very good identification**					**53435112**
Eubacterium saburreum		100.0	0.63			
53435120	**Very good identification**					**53435120**
Eubacterium saburreum		100.0	0.65			
53435122	**Good identification**					**53435122**
Eubacterium saburreum		98.87	0.52			
Parabacteroides distasonis		1.13	0.19			
53435130	**Very good identification**					**53435130**
Eubacterium saburreum		100.0	0.63			
53435132	**Good identification**					**53435132**
Eubacterium saburreum		99.25	0.50			
Parabacteroides distasonis		0.75	0.14			
53435140	**Good identification**					**53435140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.43	SOR 1		
53435150	**Good identification**					**53435150**
Eubacterium saburreum		100.0	0.42	SOR 1		
53435200	**Good identification**					**53435200**
Eubacterium saburreum		100.0	0.37	RHA 1		
53435210	**Good identification**					**53435210**
Eubacterium saburreum		100.0	0.36	RHA 1		
53435300	**Good identification**					**53435300**
Eubacterium saburreum		100.0	0.43	RHA 1		
53435310	**Good identification**					**53435310**
Eubacterium saburreum		100.0	0.42	RHA 1		
53435400	**Good identification**					**53435400**
Eubacterium saburreum		100.0	0.37	ARA 1		
53435410	**Good identification**					**53435410**
Eubacterium saburreum		100.0	0.36	ARA 1		
53435500	**Good identification**					**53435500**
Eubacterium saburreum		100.0	0.43	ARA 1		
53435510	**Good identification**					**53435510**
Eubacterium saburreum		100.0	0.42	ARA 1		
53435722	**Good identification**					**53435722**
Bacteroides thetaiotaomicron		98.76	0.40	MNS 99		
Parabacteroides distasonis		1.24	0.23	MLT 94		
53437000	**Good identification**					**53437000**
Eubacterium saburreum		100.0	0.37	MAN 1		
53437010	**Good identification**					**53437010**
Eubacterium saburreum		100.0	0.36	MAN 1		
53437100	**Good identification**					**53437100**
Eubacterium saburreum		100.0	0.43	MAN 1		
53437110	**Good identification**					**53437110**
Eubacterium saburreum		100.0	0.42	MAN 1		
53441000	**Low discrimination**					**53441000**
Eubacterium saburreum		87.21	0.37	CEL 1		
Eubacterium tortuosum		12.79	0.07			
53441010	**Good identification**					**53441010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.35	CEL 1		
53441100	**Good identification**					**53441100**
Eubacterium saburreum		98.44	0.43	CEL 1		
Fusobacterium mortiferum		1.57	0.14			
53441110	**Good identification**					**53441110**
Eubacterium saburreum		100.0	0.41	CEL 1		
53445100	**Good identification**					**53445100**
Eubacterium saburreum		100.0	0.38	CEL 1		
53445110	**Good identification**					**53445110**
Eubacterium saburreum		100.0	0.37	CEL 1		
53450000	**Good identification**					**53450000**
Fusobacterium mortiferum		97.19	0.39	IND 1		
Eubacterium saburreum		2.81	0.11			
53450100	**Good identification**					**53450100**
Fusobacterium mortiferum		97.19	0.45	IND 1		
Eubacterium saburreum		2.81	0.17			
53451000	**Low discrimination**					**53451000**
Eubacterium saburreum		65.58	0.45	CEL 1		
Fusobacterium mortiferum		34.42	0.42			
53451010	**Good identification**					**53451010**
Eubacterium saburreum		100.0	0.43	CEL 1		
53451100	**Low discrimination**					**53451100**
Eubacterium saburreum		65.58	0.51	CEL 1		
Fusobacterium mortiferum		34.42	0.48			
53451102	**Low discrimination**					**53451102**
Eubacterium saburreum		75.16	0.38	CEL 1		
Fusobacterium mortiferum		24.84	0.32			
53451110	**Good identification**					**53451110**
Eubacterium saburreum		99.36	0.49	CEL 1		
Fusobacterium mortiferum		0.64	0.14			
53451112	**Good identification**					**53451112**
Eubacterium saburreum		100.0	0.37	CEL 1		
53451120	**Acceptable identification**					**53451120**
Eubacterium saburreum		94.35	0.38	CEL 1		
Fusobacterium mortiferum		2.83	0.14			
Prevotella melaninogenica		2.82	-.04			
53451130	**Good identification**					**53451130**
Eubacterium saburreum		100.0	0.37	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53455000	**Acceptable identification**					**53455000**
Eubacterium saburreum		93.17	0.40	CEL 1		
Fusobacterium mortiferum		6.84	0.22			
53455010	**Good identification**					**53455010**
Eubacterium saburreum		100.0	0.39	CEL 1		
53455100	**Acceptable identification**					**53455100**
Eubacterium saburreum		93.17	0.46	CEL 1		
Fusobacterium mortiferum		6.84	0.29			
53455110	**Good identification**					**53455110**
Eubacterium saburreum		100.0	0.45	CEL 1		
53461100	**Good identification**					**53461100**
Eubacterium saburreum		100.0	0.40	CEL 1		
53461110	**Good identification**					**53461110**
Eubacterium saburreum		100.0	0.38	CEL 1		
53465100	**Good identification**					**53465100**
Eubacterium saburreum		100.0	0.35	CEL 1		
53470100	**Acceptable identification**					**53470100**
Fusobacterium mortiferum		93.95	0.36	IND 1		
Eubacterium saburreum		6.06	0.14			
53471000	**Low discrimination**					**53471000**
Eubacterium saburreum		80.96	0.42	CEL 1		
Fusobacterium mortiferum		19.04	0.33			
53471010	**Good identification**					**53471010**
Eubacterium saburreum		100.0	0.40	CEL 1		
53471100	**Low discrimination**					**53471100**
Eubacterium saburreum		80.96	0.48	CEL 1		
Fusobacterium mortiferum		19.04	0.39			
53471102	**Low discrimination**					**53471102**
Eubacterium saburreum		87.11	0.35	CEL 1		
Fusobacterium mortiferum		12.90	0.23			
53471110	**Good identification**					**53471110**
Eubacterium saburreum		100.0	0.46	CEL 1		
53471120	**Good identification**					**53471120**
Eubacterium saburreum		100.0	0.35	CEL 1		
53475000	**Good identification**					**53475000**
Eubacterium saburreum		96.82	0.37	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		3.18	0.13				
53475010	**Good identification**						**53475010**
Eubacterium saburreum		100.0	0.36	CEL 1			
53475100	**Good identification**						**53475100**
Eubacterium saburreum		96.82	0.43	CEL 1			
Fusobacterium mortiferum		3.18	0.20				
53475110	**Good identification**						**53475110**
Eubacterium saburreum		100.0	0.42	CEL 1			
53475722	**Good identification**						**53475722**
Bacteroides thetaiotaomicron		97.44	0.45	MNS 99			
Bacteroides ovatus		2.08	0.17	MLT 94			
Parabacteroides distasonis		0.31	0.19				
53475732	**Good identification**						**53475732**
Bacteroides thetaiotaomicron		96.06	0.38	MNS 99			
Bacteroides ovatus		3.52	0.14	MLT 94			
Parabacteroides distasonis		0.42	0.14				
53500000	**Low discrimination**						**53500000**
					SPO	GRA	
Eubacterium saburreum		61.42	0.47	GAL 99	1	85	
Clostridium bifermentans		33.23	0.30		99	99	
Prevotella intermedia		2.51	0.14		1	1	
53500010	**Good identification**						**53500010**
Eubacterium saburreum		100.0	0.45	GAL 99			
53500100	**Good identification**						**53500100**
					COC		
Eubacterium saburreum		99.54	0.53	GAL 99	1		
Blautia hansenii		0.47	-.03		99		
53500102	**Good identification**						**53500102**
Eubacterium saburreum		100.0	0.40	GAL 99			
53500110	**Good identification**						**53500110**
Eubacterium saburreum		100.0	0.52	GAL 99			
53500112	**Good identification**						**53500112**
Eubacterium saburreum		100.0	0.39	GAL 99			
53500120	**Good identification**						**53500120**
Eubacterium saburreum		97.81	0.40	GAL 99			
Prevotella melaninogenica		2.19	-.04				
53500130	**Good identification**						**53500130**
Eubacterium saburreum		100.0	0.39	GAL 99			
53501000	**Very good identification**						**53501000**
					COC	GRA	
Eubacterium saburreum		99.95	0.81		1	85	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella intermedia		0.04	0.14		1	1	
Blautia hansenii		0.01	-.03		99	99	
53501001	**Good identification**						**53501001**
					SPO		
Eubacterium saburreum		98.90	0.47	URE 1	1		
Clostridium sordellii		1.10	-.03		99		
53501002	**Very good identification**						**53501002**
Eubacterium saburreum		100.0	0.68				
53501010	**Very good identification**						**53501010**
Eubacterium saburreum		100.0	0.79				
53501011	**Good identification**						**53501011**
Eubacterium saburreum		100.0	0.45	URE 1			
53501012	**Very good identification**						**53501012**
Eubacterium saburreum		100.0	0.67				
53501020	**Very good identification**						**53501020**
Eubacterium saburreum		99.95	0.68				
Prevotella melaninogenica		0.05	-.04				
53501022	**Good identification**						**53501022**
Eubacterium saburreum		100.0	0.55				
53501030	**Very good identification**						**53501030**
Eubacterium saburreum		100.0	0.67				
53501032	**Good identification**						**53501032**
Eubacterium saburreum		100.0	0.54				
53501040	**Good identification**						**53501040**
Eubacterium saburreum		100.0	0.47	SOR 1			
53501050	**Good identification**						**53501050**
Eubacterium saburreum		100.0	0.45	SOR 1			
53501100	**Very good identification**						**53501100**
					COC	GRA	
Eubacterium saburreum		99.53	0.87		1	85	
Blautia hansenii		0.47	0.32		99	99	
Prevotella melaninogenica		0.00	-.04		1	1	
53501101	**Good identification**						**53501101**
					COC	SPO	
Eubacterium saburreum		98.88	0.53	URE 1	1	1	
Clostridium chauvoei		0.66	0.06		1	99	
Blautia hansenii		0.46	-.03		99	1	
53501102	**Very good identification**						**53501102**
					COC		
Eubacterium saburreum		99.71	0.74		1		
Blautia hansenii		0.29	0.15		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53501103	**Good identification**						**53501103**
Eubacterium saburreum		100.0	0.40	URE 1			
53501110	**Very good identification**						**53501110**
					COC		
Eubacterium saburreum		99.99	0.86		1		
Blautia hansenii		0.01	-.03		99		
53501111	**Good identification**						**53501111**
Eubacterium saburreum		100.0	0.52	URE 1			
53501112	**Very good identification**						**53501112**
Eubacterium saburreum		100.0	0.73				
53501113	**Good identification**						**53501113**
Eubacterium saburreum		100.0	0.39	URE 1			
53501120	**Very good identification**						**53501120**
					COC	GRA	
Eubacterium saburreum		97.53	0.74		1	85	
Prevotella melaninogenica		2.18	0.30		1	1	
Blautia hansenii		0.29	0.15		99	99	
53501121	**Good identification**						**53501121**
Eubacterium saburreum		97.81	0.40	URE 1			
Prevotella melaninogenica		2.19	-.04				
53501122	**Very good identification**						**53501122**
					CAT		
Eubacterium saburreum		96.81	0.61		1		
Prevotella melaninogenica		2.34	0.18		1		
Capnocytophaga ochracea		0.67	0.12		90		
53501130	**Very good identification**						**53501130**
Eubacterium saburreum		99.97	0.73				
Prevotella melaninogenica		0.03	-.04				
53501131	**Good identification**						**53501131**
Eubacterium saburreum		100.0	0.39	URE 1			
53501132	**Very good identification**						**53501132**
Eubacterium saburreum		100.0	0.60				
53501140	**Good identification**						**53501140**
					COC		
Eubacterium saburreum		99.54	0.53	SOR 1	1		
Blautia hansenii		0.47	-.03		99		
53501142	**Good identification**						**53501142**
Eubacterium saburreum		100.0	0.40	SOR 1			
53501150	**Good identification**						**53501150**
Eubacterium saburreum		100.0	0.52	SOR 1			
53501152	**Good identification**						**53501152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.39	SOR 1		
53501160	**Good identification**					**53501160**
Eubacterium saburreum		97.81	0.40	SOR 1		
Prevotella melaninogenica		2.19	-.04			
53501170	**Good identification**					**53501170**
Eubacterium saburreum		100.0	0.39	SOR 1		
53501200	**Good identification**					**53501200**
Eubacterium saburreum		100.0	0.47	RHA 1		
53501210	**Good identification**					**53501210**
Eubacterium saburreum		100.0	0.45	RHA 1		
53501300	**Good identification**					**53501300**
					COC	
Eubacterium saburreum		99.54	0.53	RHA 1	1	
Blautia hansenii		0.47	-.03		99	
53501302	**Good identification**					**53501302**
Eubacterium saburreum		100.0	0.40	RHA 1		
53501310	**Good identification**					**53501310**
Eubacterium saburreum		100.0	0.52	RHA 1		
53501312	**Good identification**					**53501312**
Eubacterium saburreum		100.0	0.39	RHA 1		
53501320	**Good identification**					**53501320**
Eubacterium saburreum		97.81	0.40	RHA 1		
Prevotella melaninogenica		2.19	-.04			
53501330	**Good identification**					**53501330**
Eubacterium saburreum		100.0	0.39	RHA 1		
53501400	**Good identification**					**53501400**
Eubacterium saburreum		100.0	0.47	ARA 1		
53501410	**Good identification**					**53501410**
Eubacterium saburreum		100.0	0.45	ARA 1		
53501500	**Acceptable identification**					**53501500**
					COC	
Eubacterium saburreum		91.81	0.53	ARA 1	1	
Blautia hansenii		7.50	0.19		99	
Bifidobacterium longum subsp. longum		0.70	0.13		1	
53501502	**Acceptable identification**					**53501502**
					COC	
Eubacterium saburreum		91.37	0.40	ARA 1	1	
Blautia hansenii		4.70	0.03		99	
Bifidobacterium longum subsp. longum		3.94	0.13		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53501510	**Good identification**					**53501510**
Eubacterium saburreum		98.44	0.52	ARA 1		
Bifidobacterium longum subsp. longum		1.56	0.17			
53501512	**Acceptable identification**					**53501512**
Eubacterium saburreum		91.77	0.39	ARA 1		
Bifidobacterium longum subsp. longum		8.23	0.17			
53501520	**Low discrimination**					**53501520**
					COC	
Eubacterium saburreum		80.25	0.40	ARA 1	1	
Bifidobacterium longum subsp. longum		13.83	0.23		1	
Blautia hansenii		4.13	0.03		99	
53501530	**Low discrimination**					**53501530**
Eubacterium saburreum		73.60	0.39	ARA 1		
Bifidobacterium longum subsp. longum		26.40	0.27			
53503000	**Good identification**					**53503000**
Eubacterium saburreum		100.0	0.47	MAN 1		
53503010	**Good identification**					**53503010**
Eubacterium saburreum		100.0	0.45	MAN 1		
53503100	**Good identification**					**53503100**
					COC	
Eubacterium saburreum		99.54	0.53	MAN 1	1	
Blautia hansenii		0.47	-.03		99	
53503102	**Good identification**					**53503102**
Eubacterium saburreum		100.0	0.40	MAN 1		
53503110	**Good identification**					**53503110**
Eubacterium saburreum		100.0	0.52	MAN 1		
53503112	**Good identification**					**53503112**
Eubacterium saburreum		100.0	0.39	MAN 1		
53503120	**Good identification**					**53503120**
Eubacterium saburreum		97.81	0.40	MAN 1		
Prevotella melaninogenica		2.19	-.04			
53503130	**Good identification**					**53503130**
Eubacterium saburreum		100.0	0.39	MAN 1		
53504000	**Good identification**					**53504000**
Eubacterium saburreum		100.0	0.42	GAL 99		
53504010	**Good identification**					**53504010**
Eubacterium saburreum		100.0	0.41	GAL 99		
53504100	**Good identification**					**53504100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.49	GAL 99		
53504102	**Good identification**					**53504102**
Eubacterium saburreum		100.0	0.36	GAL 99		
53504110	**Good identification**					**53504110**
Eubacterium saburreum		100.0	0.47	GAL 99		
53504120	**Good identification**					**53504120**
Eubacterium saburreum		100.0	0.36	GAL 99		
53505000	**Very good identification**					**53505000**
Eubacterium saburreum		100.0	0.76			
53505001	**Good identification**					**53505001**
Eubacterium saburreum		100.0	0.42	URE 1		
53505002	**Very good identification**					**53505002**
Eubacterium saburreum		100.0	0.63			
53505010	**Very good identification**					**53505010**
Eubacterium saburreum		100.0	0.75			
53505011	**Good identification**					**53505011**
Eubacterium saburreum		100.0	0.41	URE 1		
53505012	**Very good identification**					**53505012**
Eubacterium saburreum		100.0	0.62			
53505020	**Very good identification**					**53505020**
Eubacterium saburreum		100.0	0.63			
53505022	**Good identification**					**53505022**
Eubacterium saburreum		100.0	0.51			
53505030	**Very good identification**					**53505030**
Eubacterium saburreum		100.0	0.62			
53505032	**Good identification**					**53505032**
Eubacterium saburreum		100.0	0.49			
53505040	**Good identification**					**53505040**
Eubacterium saburreum		100.0	0.42	SOR 1		
53505050	**Good identification**					**53505050**
Eubacterium saburreum		100.0	0.41	SOR 1		
53505100	**Very good identification**					**53505100**
					COC	
Eubacterium saburreum		99.85	0.83		1	
Blautia hansenii		0.15	0.19		99	
53505101	**Good identification**					**53505101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.49	URE 1			
53505102	**Very good identification**						**53505102**
					COC		
Eubacterium saburreum		99.90	0.70		1		
Blautia hansenii		0.10	0.03		99		
53505103	**Good identification**						**53505103**
Eubacterium saburreum		100.0	0.36	URE 1			
53505110	**Very good identification**						**53505110**
Eubacterium saburreum		100.0	0.81				
53505111	**Good identification**						**53505111**
Eubacterium saburreum		100.0	0.47	URE 1			
53505112	**Very good identification**						**53505112**
Eubacterium saburreum		100.0	0.68				
53505120	**Very good identification**						**53505120**
					COC	GRA	
Eubacterium saburreum		99.86	0.70		1	85	
Blautia hansenii		0.10	0.03		99	99	
Prevotella melaninogenica		0.04	-.04		1	1	
53505121	**Good identification**						**53505121**
Eubacterium saburreum		100.0	0.36	URE 1			
53505122	**Very good identification**						**53505122**
					CAT		
Eubacterium saburreum		99.08	0.57		1		
Bacteroides fragilis		0.60	-.03		99		
Capnocytophaga ochracea		0.32	0.02		90		
53505130	**Very good identification**						**53505130**
Eubacterium saburreum		100.0	0.68				
53505132	**Good identification**						**53505132**
Eubacterium saburreum		100.0	0.55				
53505140	**Good identification**						**53505140**
Eubacterium saburreum		100.0	0.49	SOR 1			
53505142	**Good identification**						**53505142**
Eubacterium saburreum		100.0	0.36	SOR 1			
53505150	**Good identification**						**53505150**
Eubacterium saburreum		100.0	0.47	SOR 1			
53505160	**Good identification**						**53505160**
Eubacterium saburreum		100.0	0.36	SOR 1			
53505200	**Good identification**						**53505200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.42	RHA 1		
53505210	**Good identification**					**53505210**
Eubacterium saburreum		100.0	0.41	RHA 1		
53505300	**Good identification**					**53505300**
Eubacterium saburreum		100.0	0.49	RHA 1		
53505302	**Good identification**					**53505302**
Eubacterium saburreum		100.0	0.36	RHA 1		
53505310	**Good identification**					**53505310**
Eubacterium saburreum		100.0	0.47	RHA 1		
53505320	**Good identification**					**53505320**
Eubacterium saburreum		100.0	0.36	RHA 1		
53505400	**Good identification**					**53505400**
Eubacterium saburreum		100.0	0.42	ARA 1		
53505410	**Good identification**					**53505410**
Eubacterium saburreum		100.0	0.41	ARA 1		
53505500	**Acceptable identification**					**53505500**
					COC	
Eubacterium saburreum		93.54	0.49	ARA 1	1	
Bifidobacterium longum subsp. longum		3.96	0.21		1	
Blautia hansenii		2.50	0.06		99	
53505502	**Low discrimination**					**53505502**
Eubacterium saburreum		80.64	0.36	ARA 1		
Bifidobacterium longum subsp. longum		19.36	0.21			
53505510	**Acceptable identification**					**53505510**
Eubacterium saburreum		91.90	0.47	ARA 1		
Bifidobacterium longum subsp. longum		8.10	0.25			
53505520	**Low discrimination**					**53505520**
Eubacterium saburreum		51.02	0.36	ARA 1		
Bifidobacterium longum subsp. longum		48.99	0.31			
53505530	**Low discrimination**					**53505530**
Bifidobacterium longum subsp. longum		66.65	0.35	IND 1		
Eubacterium saburreum		33.35	0.34	MNS 87		
53505532	**Acceptable identification**					**53505532**
Bifidobacterium longum subsp. longum		91.89	0.35	IND 1		
Eubacterium saburreum		8.12	0.21	MNS 87		
53507000	**Good identification**					**53507000**
Eubacterium saburreum		100.0	0.42	MAN 1		
53507010	**Good identification**					**53507010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.41	MAN 1		
53507100	**Good identification**					**53507100**
Eubacterium saburreum		100.0	0.49	MAN 1		
53507102	**Good identification**					**53507102**
Eubacterium saburreum		100.0	0.36	MAN 1		
53507110	**Good identification**					**53507110**
Eubacterium saburreum		100.0	0.47	MAN 1		
53507120	**Good identification**					**53507120**
Eubacterium saburreum		100.0	0.36	MAN 1		
53510000	**Low discrimination**					**53510000**
Prevotella intermedia		56.40	0.48	ESL 1		
Eubacterium saburreum		41.85	0.55			
Fusobacterium mortiferum		1.36	0.31			
53510002	**Acceptable identification**					**53510002**
Eubacterium saburreum		91.11	0.42	GAL 99		
Prevotella intermedia		7.03	0.14			
Fusobacterium mortiferum		1.86	0.15			
53510010	**Good identification**					**53510010**
Eubacterium saburreum		98.36	0.53	GAL 99		
Prevotella intermedia		1.64	0.14			
53510012	**Good identification**					**53510012**
Eubacterium saburreum		100.0	0.41	GAL 99		
53510020	**Acceptable identification**					**53510020**
Eubacterium saburreum		92.84	0.42	GAL 99		
Prevotella intermedia		7.16	0.14			
53510030	**Good identification**					**53510030**
Eubacterium saburreum		100.0	0.41	GAL 99		
53510100	**Very good identification**					**53510100**
Eubacterium saburreum		96.31	0.61	GAL 99		
Fusobacterium mortiferum		3.12	0.37			
Prevotella intermedia		0.56	0.14			
53510102	**Good identification**					**53510102**
Eubacterium saburreum		98.00	0.48	GAL 99		
Fusobacterium mortiferum		2.00	0.21			
53510110	**Very good identification**					**53510110**
Eubacterium saburreum		100.0	0.60	GAL 99		
53510112	**Good identification**					**53510112**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.47	GAL 99			
53510120	**Acceptable identification**						**53510120**
Eubacterium saburreum		93.71	0.48	GAL 99			
Prevotella melaninogenica		6.30	0.12				
53510122	**Low discrimination**						**53510122**
					CAT		
Eubacterium saburreum		76.85	0.35	GAL 99	1		
Capnocytophaga ochracea		17.57	0.12		90		
Prevotella melaninogenica		5.57	0.00		1		
53510130	**Good identification**						**53510130**
Eubacterium saburreum		100.0	0.47	GAL 99			
53511000	**Very good identification**						**53511000**
Eubacterium saburreum		98.59	0.89				
Prevotella intermedia		1.34	0.48				
Fusobacterium mortiferum		0.05	0.34				
53511001	**Good identification**						**53511001**
					SPO	GRA	
Eubacterium saburreum		98.30	0.55	URE 1	1	85	
Prevotella intermedia		1.34	0.14		1	1	
Clostridium sordellii		0.36	-.03		99	99	
53511002	**Very good identification**						**53511002**
					GRA		
Eubacterium saburreum		99.79	0.76		85		
Lactobacillus fermentum		0.10	0.28		99		
Prevotella intermedia		0.08	0.14		1		
53511003	**Good identification**						**53511003**
Eubacterium saburreum		100.0	0.42	URE 1			
53511010	**Very good identification**						**53511010**
Eubacterium saburreum		99.98	0.88				
Prevotella intermedia		0.02	0.14				
53511011	**Good identification**						**53511011**
Eubacterium saburreum		100.0	0.53	URE 1			
53511012	**Very good identification**						**53511012**
Eubacterium saburreum		100.0	0.75				
53511013	**Good identification**						**53511013**
Eubacterium saburreum		100.0	0.41	URE 1			
53511020	**Very good identification**						**53511020**
Eubacterium saburreum		99.73	0.76				
Prevotella melaninogenica		0.16	0.12				
Prevotella intermedia		0.08	0.14				
53511021	**Good identification**						**53511021**
Eubacterium saburreum		100.0	0.42	URE 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53511022	**Very good identification**						**53511022**
					CAT	GRA	
Eubacterium saburreum		99.03	0.63		1	85	
Capnocytophaga ochracea		0.53	0.12		90	1	
Lactobacillus fermentum		0.19	0.20		1	99	
53511030	**Very good identification**						**53511030**
Eubacterium saburreum		100.0	0.75				
53511031	**Good identification**						**53511031**
Eubacterium saburreum		100.0	0.41	URE 1			
53511032	**Very good identification**						**53511032**
Eubacterium saburreum		100.0	0.62				
53511040	**Good identification**						**53511040**
Eubacterium saburreum		98.66	0.55	SOR 1			
Prevotella intermedia		1.34	0.14				
53511042	**Good identification**						**53511042**
Eubacterium saburreum		100.0	0.42	SOR 1			
53511050	**Good identification**						**53511050**
Eubacterium saburreum		100.0	0.53	SOR 1			
53511052	**Good identification**						**53511052**
Eubacterium saburreum		100.0	0.41	SOR 1			
53511060	**Good identification**						**53511060**
Eubacterium saburreum		100.0	0.42	SOR 1			
53511070	**Good identification**						**53511070**
Eubacterium saburreum		100.0	0.41	SOR 1			
53511100	**Excellent identification**						**53511100**
					GRA		
Eubacterium saburreum		99.91	0.95		85		
Fusobacterium mortiferum		0.05	0.40		1		
Lactobacillus fermentum		0.01	0.32		99		
53511101	**Very good identification**						**53511101**
					SPO		
Eubacterium saburreum		99.48	0.61	URE 1	1		
Clostridium chauvoei		0.52	0.12		99		
53511102	**Very good identification**						**53511102**
					CAT	GRA	
Eubacterium saburreum		99.44	0.83		1	85	
Capnocytophaga ochracea		0.45	0.30		90	1	
Lactobacillus fermentum		0.07	0.32		1	99	
53511103	**Good identification**						**53511103**
					SPO		
Eubacterium saburreum		99.48	0.48	URE 1	1		
Clostridium chauvoei		0.52	-.01		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53511110	**Excellent identification**						**53511110**
					CAT		
Eubacterium saburreum		100.0	0.94		1		
Capnocytophaga ochracea		0.00	-.03		90		
53511111	**Very good identification**						**53511111**
Eubacterium saburreum		100.0	0.60	URE 1			
53511112	**Very good identification**						**53511112**
					CAT		
Eubacterium saburreum		99.94	0.81		1		
Capnocytophaga ochracea		0.06	0.14		90		
53511113	**Good identification**						**53511113**
Eubacterium saburreum		100.0	0.47	URE 1			
53511120	**Good identification**						**53511120**
					CAT		
Eubacterium saburreum		93.28	0.83		1		
Prevotella melaninogenica		6.27	0.47		1		
Capnocytophaga ochracea		0.42	0.30		90		
53511121	**Acceptable identification**						**53511121**
					SPO	GRA	
Eubacterium saburreum		92.85	0.48	URE 1	1	85	
Prevotella melaninogenica		6.24	0.12		1	1	
Clostridium chauvoei		0.91	0.04		99	99	
53511122	**Low discrimination**						**53511122**
					CAT		
Eubacterium saburreum		76.36	0.70		1		
Capnocytophaga ochracea		17.46	0.47		90		
Prevotella melaninogenica		5.54	0.34		1		
53511123	**Low discrimination**						**53511123**
					CAT		
Eubacterium saburreum		76.85	0.35	URE 1	1		
Capnocytophaga ochracea		17.57	0.12		90		
Prevotella melaninogenica		5.57	0.00		1		
53511130	**Very good identification**						**53511130**
					CAT		
Eubacterium saburreum		99.85	0.81		1		
Prevotella melaninogenica		0.08	0.12		1		
Capnocytophaga ochracea		0.06	0.14		90		
53511131	**Good identification**						**53511131**
Eubacterium saburreum		100.0	0.47	URE 1			
53511132	**Very good identification**						**53511132**
					CAT		
Eubacterium saburreum		96.85	0.68		1		
Capnocytophaga ochracea		3.01	0.30		90		
Prevotella melaninogenica		0.09	0.00		1		
53511140	**Very good identification**						**53511140**
					CAT		
Eubacterium saburreum		99.90	0.61	SOR 1	1		
Capnocytophaga ochracea		0.10	-.03		90		
53511142	**Good identification**						**53511142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Eubacterium saburreum		95.30	0.48	SOR 1	1	
Capnocytophaga ochracea		4.70	0.14		90	
53511150	**Very good identification**					**53511150**
Eubacterium saburreum		100.0	0.60	SOR 1		
53511152	**Good identification**					**53511152**
					CAT	
Eubacterium saburreum		99.34	0.47	SOR 1	1	
Capnocytophaga ochracea		0.67	-.03		90	
53511160	**Low discrimination**					**53511160**
					CAT	
Eubacterium saburreum		89.57	0.48	SOR 1	1	
Prevotella melaninogenica		6.02	0.12		1	
Capnocytophaga ochracea		4.42	0.14		90	
53511170	**Good identification**					**53511170**
					CAT	
Eubacterium saburreum		99.34	0.47	SOR 1	1	
Capnocytophaga ochracea		0.67	-.03		90	
53511200	**Good identification**					**53511200**
Eubacterium saburreum		98.66	0.55	RHA 1		
Prevotella intermedia		1.34	0.14			
53511202	**Good identification**					**53511202**
Eubacterium saburreum		100.0	0.42	RHA 1		
53511210	**Good identification**					**53511210**
Eubacterium saburreum		100.0	0.53	RHA 1		
53511212	**Good identification**					**53511212**
Eubacterium saburreum		100.0	0.41	RHA 1		
53511220	**Good identification**					**53511220**
Eubacterium saburreum		100.0	0.42	RHA 1		
53511230	**Good identification**					**53511230**
Eubacterium saburreum		100.0	0.41	RHA 1		
53511300	**Very good identification**					**53511300**
Eubacterium saburreum		100.0	0.61	RHA 1		
53511302	**Good identification**					**53511302**
Eubacterium saburreum		100.0	0.48	RHA 1		
53511310	**Very good identification**					**53511310**
Eubacterium saburreum		100.0	0.60	RHA 1		
53511312	**Good identification**					**53511312**
Eubacterium saburreum		100.0	0.47	RHA 1		
53511320	**Acceptable identification**					**53511320**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		93.71	0.48	RHA 1			
Prevotella melaninogenica		6.30	0.12				
53511322	**Low discrimination**						**53511322**
					CAT		
Eubacterium saburreum		65.82	0.35	RHA 1	1		
Capnocytophaga ochracea		15.05	0.12		90		
Prevotella oralis		14.36	0.12		1		
53511330	**Good identification**						**53511330**
Eubacterium saburreum		100.0	0.47	RHA 1			
53511400	**Good identification**						**53511400**
					GRA		
Eubacterium saburreum		96.99	0.55	ARA 1	85		
Lactobacillus fermentum		1.69	0.28		99		
Prevotella intermedia		1.32	0.14		1		
53511402	**Acceptable identification**						**53511402**
Eubacterium saburreum		91.01	0.42	ARA 1			
Lactobacillus fermentum		8.99	0.28				
53511410	**Good identification**						**53511410**
Eubacterium saburreum		100.0	0.53	ARA 1			
53511412	**Good identification**						**53511412**
Eubacterium saburreum		100.0	0.41	ARA 1			
53511420	**Good identification**						**53511420**
Eubacterium saburreum		95.68	0.42	ARA 1			
Lactobacillus fermentum		3.15	0.20				
Bifidobacterium longum subsp. longum		1.17	0.05				
53511430	**Good identification**						**53511430**
Eubacterium saburreum		97.53	0.41	ARA 1			
Bifidobacterium longum subsp. longum		2.47	0.09				
53511500	**Very good identification**						**53511500**
Eubacterium saburreum		96.48	0.61	ARA 1			
Bifidobacterium longum subsp. longum		2.20	0.29				
Lactobacillus fermentum		1.18	0.32				
53511502	**Low discrimination**						**53511502**
					CAT	GRA	
Eubacterium saburreum		77.03	0.48	ARA 1	1	85	
Bifidobacterium longum subsp. longum		9.96	0.29		1	99	
Capnocytophaga ochracea		6.03	0.17		90	1	
53511510	**Very good identification**						**53511510**
Eubacterium saburreum		95.47	0.60	ARA 1			
Bifidobacterium longum subsp. longum		4.53	0.33				
53511512	**Low discrimination**						**53511512**
Eubacterium saburreum		76.23	0.47	ARA 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		20.51	0.33				
Bifidobacterium adolescentis		2.46	0.19				
53511520	**Low discrimination**						**53511520**
					CAT	GRA	
Eubacterium saburreum		59.33	0.48	ARA 1	1	85	
Bifidobacterium longum subsp. longum		30.67	0.40		1	99	
Capnocytophaga ochracea		4.65	0.17		90	1	
53511530	**Low discrimination**						**53511530**
					CAT	GRA	
Bifidobacterium longum subsp. longum		51.57	0.44	IND 1	1	99	
Eubacterium saburreum		47.93	0.47	MNS 87	1	85	
Capnocytophaga ochracea		0.51	0.01		90	1	
53511532	**Low discrimination**						**53511532**
					CAT	GRA	
Bifidobacterium longum subsp. longum		79.81	0.44	IND 1	1	99	
Eubacterium saburreum		13.09	0.34	MNS 87	1	85	
Capnocytophaga ochracea		7.10	0.17		90	1	
53513000	**Good identification**						**53513000**
Eubacterium saburreum		98.66	0.55	MAN 1			
Prevotella intermedia		1.34	0.14				
53513002	**Good identification**						**53513002**
Eubacterium saburreum		100.0	0.42	MAN 1			
53513010	**Good identification**						**53513010**
Eubacterium saburreum		100.0	0.53	MAN 1			
53513012	**Good identification**						**53513012**
Eubacterium saburreum		100.0	0.41	MAN 1			
53513020	**Good identification**						**53513020**
Eubacterium saburreum		100.0	0.42	MAN 1			
53513030	**Good identification**						**53513030**
Eubacterium saburreum		100.0	0.41	MAN 1			
53513100	**Very good identification**						**53513100**
Eubacterium saburreum		99.91	0.61	MAN 1			
Propionibacterium propionicum		0.09	0.12				
53513102	**Good identification**						**53513102**
Eubacterium saburreum		99.24	0.48	MAN 1			
Propionibacterium propionicum		0.76	0.15				
53513110	**Very good identification**						**53513110**
Eubacterium saburreum		100.0	0.60	MAN 1			
53513112	**Good identification**						**53513112**
Eubacterium saburreum		100.0	0.47	MAN 1			
53513120	**Acceptable identification**						**53513120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		93.71	0.48	MAN 1		
Prevotella melaninogenica		6.30	0.12			
53513122	**Low discrimination**					**53513122**
					CAT	
Eubacterium saburreum		76.85	0.35	MAN 1	1	
Capnocytophaga ochracea		17.57	0.12		90	
Prevotella melaninogenica		5.57	0.00		1	
53513130	**Good identification**					**53513130**
Eubacterium saburreum		100.0	0.47	MAN 1		
53514000	**Good identification**					**53514000**
Eubacterium saburreum		97.11	0.50	GAL 99		
Prevotella intermedia		2.46	0.14			
Fusobacterium mortiferum		0.44	0.12			
53514002	**Good identification**					**53514002**
Eubacterium saburreum		100.0	0.37	GAL 99		
53514010	**Good identification**					**53514010**
Eubacterium saburreum		100.0	0.49	GAL 99		
53514012	**Good identification**					**53514012**
Eubacterium saburreum		100.0	0.36	GAL 99		
53514020	**Good identification**					**53514020**
Eubacterium saburreum		100.0	0.37	GAL 99		
53514030	**Good identification**					**53514030**
Eubacterium saburreum		100.0	0.36	GAL 99		
53514100	**Very good identification**					**53514100**
Eubacterium saburreum		99.55	0.57	GAL 99		
Fusobacterium mortiferum		0.45	0.18			
53514102	**Good identification**					**53514102**
Eubacterium saburreum		100.0	0.44	GAL 99		
53514110	**Good identification**					**53514110**
Eubacterium saburreum		100.0	0.55	GAL 99		
53514112	**Good identification**					**53514112**
Eubacterium saburreum		100.0	0.42	GAL 99		
53514120	**Good identification**					**53514120**
Eubacterium saburreum		100.0	0.44	GAL 99		
53514130	**Good identification**					**53514130**
Eubacterium saburreum		100.0	0.42	GAL 99		
53515000	**Very good identification**					**53515000**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.95	0.85		85	
Prevotella intermedia		0.03	0.14		1	
Lactobacillus fermentum		0.02	0.24		99	
53515001	**Good identification**					**53515001**
Eubacterium saburreum		100.0	0.50	URE 1		
53515002	**Very good identification**					**53515002**
Eubacterium saburreum		99.89	0.72			
Lactobacillus fermentum		0.11	0.24			
53515003	**Good identification**					**53515003**
Eubacterium saburreum		100.0	0.37	URE 1		
53515010	**Very good identification**					**53515010**
Eubacterium saburreum		100.0	0.83			
53515011	**Good identification**					**53515011**
Eubacterium saburreum		100.0	0.49	URE 1		
53515012	**Very good identification**					**53515012**
Eubacterium saburreum		100.0	0.70			
53515013	**Good identification**					**53515013**
Eubacterium saburreum		100.0	0.36	URE 1		
53515020	**Very good identification**					**53515020**
Eubacterium saburreum		99.96	0.72			
Lactobacillus fermentum		0.04	0.16			
53515021	**Good identification**					**53515021**
Eubacterium saburreum		100.0	0.37	URE 1		
53515022	**Very good identification**					**53515022**
					CAT	
Eubacterium saburreum		99.08	0.59		1	
Bacteroides fragilis		0.47	-.03		99	
Capnocytophaga ochracea		0.25	0.02		90	
53515030	**Very good identification**					**53515030**
Eubacterium saburreum		100.0	0.70			
53515031	**Good identification**					**53515031**
Eubacterium saburreum		100.0	0.36	URE 1		
53515032	**Very good identification**					**53515032**
Eubacterium saburreum		100.0	0.57			
53515040	**Good identification**					**53515040**
Eubacterium saburreum		100.0	0.50	SOR 1		
53515042	**Good identification**					**53515042**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.37	SOR 1			
53515050	**Good identification**						**53515050**
Eubacterium saburreum		100.0	0.49	SOR 1			
53515052	**Good identification**						**53515052**
Eubacterium saburreum		100.0	0.36	SOR 1			
53515060	**Good identification**						**53515060**
Eubacterium saburreum		100.0	0.37	SOR 1			
53515070	**Good identification**						**53515070**
Eubacterium saburreum		100.0	0.36	SOR 1			
53515100	**Very good identification**						**53515100**
					GRA		
Eubacterium saburreum		99.98	0.91		85		
Lactobacillus fermentum		0.01	0.28		99		
Fusobacterium mortiferum		0.01	0.21		1		
53515101	**Very good identification**						**53515101**
Eubacterium saburreum		100.0	0.57	URE 1			
53515102	**Very good identification**						**53515102**
					CAT	GRA	
Eubacterium saburreum		99.64	0.78		1	85	
Capnocytophaga ochracea		0.21	0.20		90	1	
Lactobacillus fermentum		0.08	0.28		1	99	
53515103	**Good identification**						**53515103**
Eubacterium saburreum		100.0	0.44	URE 1			
53515110	**Very good identification**						**53515110**
Eubacterium saburreum		100.0	0.89				
Bifidobacterium longum subsp. longum		0.00	0.07				
53515111	**Good identification**						**53515111**
Eubacterium saburreum		100.0	0.55	URE 1			
53515112	**Very good identification**						**53515112**
					CAT	GRA	
Eubacterium saburreum		99.90	0.76		1	85	
Bifidobacterium adolescentis		0.06	0.19		1	99	
Capnocytophaga ochracea		0.03	0.04		90	1	
53515113	**Good identification**						**53515113**
Eubacterium saburreum		100.0	0.42	URE 1			
53515120	**Very good identification**						**53515120**
					CAT		
Eubacterium saburreum		99.41	0.78		1		
Capnocytophaga ochracea		0.21	0.20		90		
Bacteroides fragilis		0.19	0.10		99		
53515121	**Good identification**						**53515121**
Eubacterium saburreum		100.0	0.44	URE 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
53515122	**Low discrimination**						**53515122**
					CAT		
Eubacterium saburreum		76.01	0.65		1		
Bacteroides fragilis		15.26	0.32		99		
Capnocytophaga ochracea		8.07	0.36		90		
53515130	**Very good identification**						**53515130**
					CAT	GRA	
Eubacterium saburreum		99.91	0.76		1	85	
Bifidobacterium longum subsp. longum		0.06	0.18		1	99	
Capnocytophaga ochracea		0.03	0.04		90	1	
53515131	**Good identification**						**53515131**
Eubacterium saburreum		100.0	0.42	URE 1			
53515132	**Very good identification**						**53515132**
					CAT	GRA	
Eubacterium saburreum		97.91	0.64		1	85	
Capnocytophaga ochracea		1.41	0.20		90	1	
Bifidobacterium longum subsp. longum		0.34	0.18		1	99	
53515140	**Very good identification**						**53515140**
Eubacterium saburreum		100.0	0.57	SOR 1			
53515142	**Good identification**						**53515142**
					CAT		
Eubacterium saburreum		97.76	0.44	SOR 1	1		
Capnocytophaga ochracea		2.24	0.04		90		
53515150	**Good identification**						**53515150**
Eubacterium saburreum		100.0	0.55	SOR 1			
53515152	**Good identification**						**53515152**
Eubacterium saburreum		100.0	0.42	SOR 1			
53515160	**Good identification**						**53515160**
					CAT		
Eubacterium saburreum		97.76	0.44	SOR 1	1		
Capnocytophaga ochracea		2.24	0.04		90		
53515170	**Good identification**						**53515170**
Eubacterium saburreum		100.0	0.42	SOR 1			
53515200	**Good identification**						**53515200**
Eubacterium saburreum		100.0	0.50	RHA 1			
53515202	**Good identification**						**53515202**
Eubacterium saburreum		100.0	0.37	RHA 1			
53515210	**Good identification**						**53515210**
Eubacterium saburreum		100.0	0.49	RHA 1			
53515212	**Good identification**						**53515212**
Eubacterium saburreum		100.0	0.36	RHA 1			
53515220	**Good identification**						**53515220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.37	RHA 1		
53515230	**Good identification**					**53515230**
Eubacterium saburreum		100.0	0.36	RHA 1		
53515300	**Good identification**					**53515300**
Eubacterium saburreum		92.45	0.57	RHA 1		
Eubacterium contortum		7.55	0.27			
53515302	**Acceptable identification**					**53515302**
Eubacterium saburreum		92.45	0.44	RHA 1		
Eubacterium contortum		7.55	0.14			
53515310	**Good identification**					**53515310**
Eubacterium saburreum		100.0	0.55	RHA 1		
53515312	**Good identification**					**53515312**
Eubacterium saburreum		100.0	0.42	RHA 1		
53515320	**Low discrimination**					**53515320**
					GRA	
Eubacterium saburreum		79.92	0.44	RHA 1	85	
Bacteroides vulgatus		13.55	0.13		1	
Eubacterium contortum		6.53	0.14		99	
53515330	**Good identification**					**53515330**
Eubacterium saburreum		100.0	0.42	RHA 1		
53515400	**Good identification**					**53515400**
Eubacterium saburreum		98.13	0.50	ARA 1		
Lactobacillus fermentum		1.87	0.24			
53515402	**Acceptable identification**					**53515402**
Eubacterium saburreum		90.27	0.37	ARA 1		
Lactobacillus fermentum		9.73	0.24			
53515410	**Good identification**					**53515410**
Eubacterium saburreum		99.38	0.49	ARA 1		
Bifidobacterium longum subsp. longum		0.62	0.07			
53515412	**Good identification**					**53515412**
Eubacterium saburreum		96.59	0.36	ARA 1		
Bifidobacterium longum subsp. longum		3.41	0.07			
53515420	**Acceptable identification**					**53515420**
Eubacterium saburreum		90.60	0.37	ARA 1		
Bifidobacterium longum subsp. longum		6.15	0.14			
Lactobacillus fermentum		3.25	0.16			
53515430	**Low discrimination**					**53515430**
Eubacterium saburreum		87.62	0.36	ARA 1		
Bifidobacterium longum subsp. longum		12.38	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53515500	**Low discrimination**						**53515500**
Eubacterium saburreum		87.69	0.57	ARA 1			
Bifidobacterium longum subsp. longum		11.14	0.37				
Lactobacillus fermentum		1.17	0.28				
53515502	**Low discrimination**						**53515502**
Eubacterium saburreum		47.62	0.44	ARA 1			
Bifidobacterium longum subsp. longum		34.30	0.37				
Bifidobacterium adolescentis		12.76	0.32				
53515510	**Low discrimination**						**53515510**
Eubacterium saburreum		79.08	0.55	ARA 1			
Bifidobacterium longum subsp. longum		20.92	0.41				
53515512	**Low discrimination**						**53515512**
Bifidobacterium longum subsp. longum		51.70	0.41	IND 1			
Eubacterium saburreum		34.50	0.42	MNS 87			
Bifidobacterium adolescentis		13.81	0.33				
53515520	**Low discrimination**						**53515520**
					GRA		
Bifidobacterium longum subsp. longum		68.71	0.48	IND 1	99		
Eubacterium saburreum		23.85	0.44	MNS 87	85		
Bacteroides vulgatus		5.97	0.15		1		
53515522	**Low discrimination**						**53515522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		71.24	0.48	IND 1	99	1	
Bacteroides thetaiotaomicron		10.71	0.22	MNS 87	1	89	
Capnocytophaga ochracea		8.10	0.23		1	90	
53515530	**Low discrimination**						**53515530**
Bifidobacterium longum subsp. longum		85.70	0.52	IND 1			
Eubacterium saburreum		14.30	0.42	MNS 87			
53515532	**Acceptable identification**						**53515532**
					CAT	GRA	
Bifidobacterium longum subsp. longum		93.36	0.52	IND 1	1	99	
Bacteroides thetaiotaomicron		3.20	0.15	MNS 87	89	1	
Eubacterium saburreum		2.75	0.29		1	85	
53515722	**Acceptable identification**						**53515722**
					CAT		
Bacteroides thetaiotaomicron		94.41	0.41	MNS 99	89		
Bacteroides uniformis		3.40	0.14	TRE 93	5		
Parabacteroides distasonis		0.78	0.21		76		
53517000	**Good identification**						**53517000**
Eubacterium saburreum		100.0	0.50	MAN 1			
53517002	**Good identification**						**53517002**
Eubacterium saburreum		100.0	0.37	MAN 1			
53517010	**Good identification**						**53517010**
Eubacterium saburreum		100.0	0.49	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53517012	**Good identification**					**53517012**
Eubacterium saburreum		100.0	0.36	MAN 1		
53517020	**Good identification**					**53517020**
Eubacterium saburreum		100.0	0.37	MAN 1		
53517030	**Good identification**					**53517030**
Eubacterium saburreum		100.0	0.36	MAN 1		
53517100	**Very good identification**					**53517100**
Eubacterium saburreum		100.0	0.57	MAN 1		
53517102	**Good identification**					**53517102**
Eubacterium saburreum		96.83	0.44	MAN 1		
Bifidobacterium adolescentis		3.17	0.16			
53517110	**Good identification**					**53517110**
Eubacterium saburreum		100.0	0.55	MAN 1		
53517112	**Good identification**					**53517112**
Eubacterium saburreum		95.33	0.42	MAN 1		
Bifidobacterium adolescentis		4.67	0.18			
53517120	**Good identification**					**53517120**
Eubacterium saburreum		100.0	0.44	MAN 1		
53517130	**Good identification**					**53517130**
Eubacterium saburreum		100.0	0.42	MAN 1		
53520000	**Good identification**					**53520000**
					SPO	
Eubacterium saburreum		98.61	0.44	GAL 99	1	
Clostridium botulinum biovar A		1.40	0.01		99	
53520010	**Good identification**					**53520010**
Eubacterium saburreum		100.0	0.42	GAL 99		
53520100	**Good identification**					**53520100**
Eubacterium saburreum		100.0	0.50	GAL 99		
53520102	**Good identification**					**53520102**
Eubacterium saburreum		100.0	0.37	GAL 99		
53520110	**Good identification**					**53520110**
Eubacterium saburreum		100.0	0.49	GAL 99		
53520112	**Good identification**					**53520112**
Eubacterium saburreum		100.0	0.36	GAL 99		
53520120	**Good identification**					**53520120**
Eubacterium saburreum		100.0	0.37	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53520130	**Good identification**					**53520130**
Eubacterium saburreum		100.0	0.36	GAL 99		
53521000	**Very good identification**					**53521000**
Eubacterium saburreum		100.0	0.78			
53521001	**Good identification**					**53521001**
Eubacterium saburreum		100.0	0.44	URE 1		
53521002	**Very good identification**					**53521002**
Eubacterium saburreum		100.0	0.65			
53521010	**Very good identification**					**53521010**
Eubacterium saburreum		100.0	0.76			
53521011	**Good identification**					**53521011**
Eubacterium saburreum		100.0	0.42	URE 1		
53521012	**Very good identification**					**53521012**
Eubacterium saburreum		100.0	0.64			
53521020	**Very good identification**					**53521020**
Eubacterium saburreum		100.0	0.65			
53521022	**Good identification**					**53521022**
Eubacterium saburreum		100.0	0.52			
53521030	**Very good identification**					**53521030**
Eubacterium saburreum		100.0	0.64			
53521032	**Good identification**					**53521032**
Eubacterium saburreum		100.0	0.51			
53521040	**Good identification**					**53521040**
Eubacterium saburreum		100.0	0.44	SOR 1		
53521050	**Good identification**					**53521050**
Eubacterium saburreum		100.0	0.42	SOR 1		
53521100	**Very good identification**					**53521100**
					COC	
Eubacterium saburreum		99.99	0.84		1	
Blautia hansenii		0.01	-.03		99	
53521101	**Good identification**					**53521101**
Eubacterium saburreum		100.0	0.50	URE 1		
53521102	**Very good identification**					**53521102**
Eubacterium saburreum		100.0	0.71			
53521103	**Good identification**					**53521103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.37	URE 1		
53521110	**Very good identification**					**53521110**
Eubacterium saburreum		100.0	0.83			
53521111	**Good identification**					**53521111**
Eubacterium saburreum		100.0	0.49	URE 1		
53521112	**Very good identification**					**53521112**
Eubacterium saburreum		100.0	0.70			
53521113	**Good identification**					**53521113**
Eubacterium saburreum		100.0	0.36	URE 1		
53521120	**Very good identification**					**53521120**
Eubacterium saburreum		99.97	0.71			
Prevotella melaninogenica		0.03	-.04			
53521121	**Good identification**					**53521121**
Eubacterium saburreum		100.0	0.37	URE 1		
53521122	**Very good identification**					**53521122**
Eubacterium saburreum		100.0	0.58			
53521130	**Very good identification**					**53521130**
Eubacterium saburreum		100.0	0.70			
53521131	**Good identification**					**53521131**
Eubacterium saburreum		100.0	0.36	URE 1		
53521132	**Very good identification**					**53521132**
Eubacterium saburreum		100.0	0.57			
53521140	**Good identification**					**53521140**
Eubacterium saburreum		100.0	0.50	SOR 1		
53521142	**Good identification**					**53521142**
Eubacterium saburreum		100.0	0.37	SOR 1		
53521150	**Good identification**					**53521150**
Eubacterium saburreum		100.0	0.49	SOR 1		
53521152	**Good identification**					**53521152**
Eubacterium saburreum		100.0	0.36	SOR 1		
53521160	**Good identification**					**53521160**
Eubacterium saburreum		100.0	0.37	SOR 1		
53521170	**Good identification**					**53521170**
Eubacterium saburreum		100.0	0.36	SOR 1		
53521200	**Good identification**					**53521200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.44	RHA 1		
53521210	**Good identification**					**53521210**
Eubacterium saburreum		100.0	0.42	RHA 1		
53521300	**Good identification**					**53521300**
Eubacterium saburreum		100.0	0.50	RHA 1		
53521302	**Good identification**					**53521302**
Eubacterium saburreum		100.0	0.37	RHA 1		
53521310	**Good identification**					**53521310**
Eubacterium saburreum		100.0	0.49	RHA 1		
53521312	**Good identification**					**53521312**
Eubacterium saburreum		100.0	0.36	RHA 1		
53521320	**Good identification**					**53521320**
Eubacterium saburreum		100.0	0.37	RHA 1		
53521330	**Good identification**					**53521330**
Eubacterium saburreum		100.0	0.36	RHA 1		
53521400	**Good identification**					**53521400**
Eubacterium saburreum		100.0	0.44	ARA 1		
53521410	**Good identification**					**53521410**
Eubacterium saburreum		100.0	0.42	ARA 1		
53521500	**Good identification**					**53521500**
Eubacterium saburreum		100.0	0.50	ARA 1		
53521502	**Good identification**					**53521502**
Eubacterium saburreum		100.0	0.37	ARA 1		
53521510	**Good identification**					**53521510**
Eubacterium saburreum		99.41	0.49	ARA 1		
Bifidobacterium longum subsp. longum		0.59	0.07			
53521512	**Good identification**					**53521512**
Eubacterium saburreum		96.75	0.36	ARA 1		
Bifidobacterium longum subsp. longum		3.25	0.07			
53521520	**Acceptable identification**					**53521520**
Eubacterium saburreum		93.93	0.37	ARA 1		
Bifidobacterium longum subsp. longum		6.07	0.13			
53521530	**Low discrimination**					**53521530**
Eubacterium saburreum		88.15	0.36	ARA 1		
Bifidobacterium longum subsp. longum		11.86	0.17			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
53523000	**Good identification**					**53523000**
	Eubacterium saburreum	100.0	0.44	MAN 1		
53523010	**Good identification**					**53523010**
	Eubacterium saburreum	100.0	0.42	MAN 1		
53523100	**Good identification**					**53523100**
	Eubacterium saburreum	100.0	0.50	MAN 1		
53523102	**Good identification**					**53523102**
	Eubacterium saburreum	100.0	0.37	MAN 1		
53523110	**Good identification**					**53523110**
	Eubacterium saburreum	100.0	0.49	MAN 1		
53523112	**Good identification**					**53523112**
	Eubacterium saburreum	100.0	0.36	MAN 1		
53523120	**Good identification**					**53523120**
	Eubacterium saburreum	100.0	0.37	MAN 1		
53523130	**Good identification**					**53523130**
	Eubacterium saburreum	100.0	0.36	MAN 1		
53524000	**Good identification**					**53524000**
	Eubacterium saburreum	100.0	0.39	GAL 99		
53524010	**Good identification**					**53524010**
	Eubacterium saburreum	100.0	0.38	GAL 99		
53524100	**Good identification**					**53524100**
	Eubacterium saburreum	100.0	0.46	GAL 99		
53524110	**Good identification**					**53524110**
	Eubacterium saburreum	100.0	0.44	GAL 99		
53525000	**Very good identification**					**53525000**
	Eubacterium saburreum	100.0	0.73			
53525001	**Good identification**					**53525001**
	Eubacterium saburreum	100.0	0.39	URE 1		
53525002	**Very good identification**					**53525002**
	Eubacterium saburreum	100.0	0.60			
53525010	**Very good identification**					**53525010**
	Eubacterium saburreum	100.0	0.72			
53525011	**Good identification**					**53525011**
	Eubacterium saburreum	100.0	0.38	URE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53525012	**Very good identification**					**53525012**
Eubacterium saburreum		100.0	0.59			
53525020	**Very good identification**					**53525020**
Eubacterium saburreum		100.0	0.60			
53525022	**Good identification**					**53525022**
Eubacterium saburreum		100.0	0.48			
53525030	**Very good identification**					**53525030**
Eubacterium saburreum		100.0	0.59			
53525032	**Good identification**					**53525032**
Eubacterium saburreum		100.0	0.46			
53525040	**Good identification**					**53525040**
Eubacterium saburreum		100.0	0.39	SOR 1		
53525050	**Good identification**					**53525050**
Eubacterium saburreum		100.0	0.38	SOR 1		
53525100	**Very good identification**					**53525100**
Eubacterium saburreum		100.0	0.80			
53525101	**Good identification**					**53525101**
Eubacterium saburreum		100.0	0.46	URE 1		
53525102	**Very good identification**					**53525102**
Eubacterium saburreum		100.0	0.67			
53525110	**Very good identification**					**53525110**
Eubacterium saburreum		100.0	0.78			
53525111	**Good identification**					**53525111**
Eubacterium saburreum		100.0	0.44	URE 1		
53525112	**Very good identification**					**53525112**
Eubacterium saburreum		100.0	0.65			
53525120	**Very good identification**					**53525120**
Eubacterium saburreum		100.0	0.67			
53525122	**Good identification**					**53525122**
Eubacterium saburreum		100.0	0.54			
53525130	**Very good identification**					**53525130**
Eubacterium saburreum		100.0	0.65			
53525132	**Good identification**					**53525132**
Eubacterium saburreum		100.0	0.52			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53525140	**Good identification**					**53525140**
Eubacterium saburreum		100.0	0.46	SOR 1		
53525150	**Good identification**					**53525150**
Eubacterium saburreum		100.0	0.44	SOR 1		
53525200	**Good identification**					**53525200**
Eubacterium saburreum		100.0	0.39	RHA 1		
53525210	**Good identification**					**53525210**
Eubacterium saburreum		100.0	0.38	RHA 1		
53525300	**Good identification**					**53525300**
Eubacterium saburreum		100.0	0.46	RHA 1		
53525310	**Good identification**					**53525310**
Eubacterium saburreum		100.0	0.44	RHA 1		
53525400	**Good identification**					**53525400**
Eubacterium saburreum		100.0	0.39	ARA 1		
53525410	**Good identification**					**53525410**
Eubacterium saburreum		100.0	0.38	ARA 1		
53525500	**Good identification**					**53525500**
Eubacterium saburreum		98.44	0.46	ARA 1		
Bifidobacterium longum subsp. longum		1.56	0.11			
53525510	**Good identification**					**53525510**
Eubacterium saburreum		96.80	0.44	ARA 1		
Bifidobacterium longum subsp. longum		3.20	0.15			
53527000	**Good identification**					**53527000**
Eubacterium saburreum		100.0	0.39	MAN 1		
53527010	**Good identification**					**53527010**
Eubacterium saburreum		100.0	0.38	MAN 1		
53527100	**Good identification**					**53527100**
Eubacterium saburreum		100.0	0.46	MAN 1		
53527110	**Good identification**					**53527110**
Eubacterium saburreum		100.0	0.44	MAN 1		
53530000	**Good identification**					**53530000**
Eubacterium saburreum		96.62	0.52	GAL 99		
Prevotella intermedia		1.97	0.14			
Fusobacterium mortiferum		1.40	0.22			
53530002	**Good identification**					**53530002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.39	GAL 99		
53530010	**Good identification**					**53530010**
Eubacterium saburreum		100.0	0.50	GAL 99		
53530012	**Good identification**					**53530012**
Eubacterium saburreum		100.0	0.38	GAL 99		
53530020	**Good identification**					**53530020**
Eubacterium saburreum		100.0	0.39	GAL 99		
53530030	**Good identification**					**53530030**
Eubacterium saburreum		100.0	0.38	GAL 99		
53530100	**Very good identification**					**53530100**
Eubacterium saburreum		98.57	0.58	GAL 99		
Fusobacterium mortiferum		1.43	0.28			
53530102	**Good identification**					**53530102**
Eubacterium saburreum		99.09	0.45	GAL 99		
Fusobacterium mortiferum		0.91	0.12			
53530110	**Very good identification**					**53530110**
Eubacterium saburreum		100.0	0.57	GAL 99		
53530112	**Good identification**					**53530112**
Eubacterium saburreum		100.0	0.44	GAL 99		
53530120	**Good identification**					**53530120**
Eubacterium saburreum		100.0	0.45	GAL 99		
53530130	**Good identification**					**53530130**
Eubacterium saburreum		100.0	0.44	GAL 99		
53531000	**Very good identification**					**53531000**
Eubacterium saburreum		99.96	0.86			
Fusobacterium mortiferum		0.02	0.25			
Prevotella intermedia		0.02	0.14			
53531001	**Good identification**					**53531001**
Eubacterium saburreum		100.0	0.52	URE 1		
53531002	**Very good identification**					**53531002**
Eubacterium saburreum		100.0	0.73			
53531003	**Good identification**					**53531003**
Eubacterium saburreum		100.0	0.39	URE 1		
53531010	**Very good identification**					**53531010**
Eubacterium saburreum		100.0	0.85			
53531011	**Good identification**					**53531011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.50	URE 1		
53531012	**Very good identification**					**53531012**
Eubacterium saburreum		100.0	0.72			
53531013	**Good identification**					**53531013**
Eubacterium saburreum		100.0	0.38	URE 1		
53531020	**Very good identification**					**53531020**
Eubacterium saburreum		100.0	0.73			
53531021	**Good identification**					**53531021**
Eubacterium saburreum		100.0	0.39	URE 1		
53531022	**Very good identification**					**53531022**
Eubacterium saburreum		100.0	0.60			
53531030	**Very good identification**					**53531030**
Eubacterium saburreum		100.0	0.72			
53531031	**Good identification**					**53531031**
Eubacterium saburreum		100.0	0.38	URE 1		
53531032	**Very good identification**					**53531032**
Eubacterium saburreum		100.0	0.59			
53531040	**Good identification**					**53531040**
Eubacterium saburreum		100.0	0.52	SOR 1		
53531042	**Good identification**					**53531042**
Eubacterium saburreum		100.0	0.39	SOR 1		
53531050	**Good identification**					**53531050**
Eubacterium saburreum		100.0	0.50	SOR 1		
53531052	**Good identification**					**53531052**
Eubacterium saburreum		100.0	0.38	SOR 1		
53531060	**Good identification**					**53531060**
Eubacterium saburreum		100.0	0.39	SOR 1		
53531070	**Good identification**					**53531070**
Eubacterium saburreum		100.0	0.38	SOR 1		
53531100	**Excellent identification**					**53531100**
Eubacterium saburreum		99.98	0.92			
Fusobacterium mortiferum		0.02	0.32			
53531101	**Very good identification**					**53531101**
Eubacterium saburreum		100.0	0.58	URE 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53531102	**Very good identification**					**53531102**
Eubacterium saburreum		99.99	0.80			
Fusobacterium mortiferum		0.01	0.15			
53531103	**Good identification**					**53531103**
Eubacterium saburreum		100.0	0.45	URE 1		
53531110	**Very good identification**					**53531110**
Eubacterium saburreum		100.0	0.91			
53531111	**Very good identification**					**53531111**
Eubacterium saburreum		100.0	0.57	URE 1		
53531112	**Very good identification**					**53531112**
Eubacterium saburreum		100.0	0.78			
53531113	**Good identification**					**53531113**
Eubacterium saburreum		100.0	0.44	URE 1		
53531120	**Very good identification**					**53531120**
					GRA	
Eubacterium saburreum		99.87	0.80		85	
Prevotella melaninogenica		0.10	0.12		1	
Clostridium perfringens		0.03	0.09		99	
53531121	**Good identification**					**53531121**
Eubacterium saburreum		100.0	0.45	URE 1		
53531122	**Very good identification**					**53531122**
					CAT	
Eubacterium saburreum		99.37	0.67		1	
Capnocytophaga ochracea		0.34	0.12		90	
Prevotella melaninogenica		0.11	0.00		1	
53531130	**Very good identification**					**53531130**
Eubacterium saburreum		100.0	0.78			
53531131	**Good identification**					**53531131**
Eubacterium saburreum		100.0	0.44	URE 1		
53531132	**Very good identification**					**53531132**
Eubacterium saburreum		100.0	0.65			
53531140	**Very good identification**					**53531140**
Eubacterium saburreum		100.0	0.58	SOR 1		
53531142	**Good identification**					**53531142**
Eubacterium saburreum		100.0	0.45	SOR 1		
53531150	**Very good identification**					**53531150**
Eubacterium saburreum		100.0	0.57	SOR 1		
53531152	**Good identification**					**53531152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.44	SOR 1		
53531160	**Good identification**					**53531160**
Eubacterium saburreum		100.0	0.45	SOR 1		
53531170	**Good identification**					**53531170**
Eubacterium saburreum		100.0	0.44	SOR 1		
53531200	**Good identification**					**53531200**
Eubacterium saburreum		100.0	0.52	RHA 1		
53531202	**Good identification**					**53531202**
Eubacterium saburreum		100.0	0.39	RHA 1		
53531210	**Good identification**					**53531210**
Eubacterium saburreum		100.0	0.50	RHA 1		
53531212	**Good identification**					**53531212**
Eubacterium saburreum		100.0	0.38	RHA 1		
53531220	**Good identification**					**53531220**
Eubacterium saburreum		100.0	0.39	RHA 1		
53531230	**Good identification**					**53531230**
Eubacterium saburreum		100.0	0.38	RHA 1		
53531300	**Very good identification**					**53531300**
Eubacterium saburreum		100.0	0.58	RHA 1		
53531302	**Good identification**					**53531302**
Eubacterium saburreum		100.0	0.45	RHA 1		
53531310	**Very good identification**					**53531310**
Eubacterium saburreum		100.0	0.57	RHA 1		
53531312	**Good identification**					**53531312**
Eubacterium saburreum		100.0	0.44	RHA 1		
53531320	**Good identification**					**53531320**
Eubacterium saburreum		100.0	0.45	RHA 1		
53531330	**Good identification**					**53531330**
Eubacterium saburreum		100.0	0.44	RHA 1		
53531400	**Good identification**					**53531400**
Eubacterium saburreum		100.0	0.52	ARA 1		
53531402	**Good identification**					**53531402**
Eubacterium saburreum		100.0	0.39	ARA 1		
53531410	**Good identification**					**53531410**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.50	ARA 1		
53531412	**Good identification**					**53531412**
Eubacterium saburreum		100.0	0.38	ARA 1		
53531420	**Good identification**					**53531420**
Eubacterium saburreum		100.0	0.39	ARA 1		
53531430	**Good identification**					**53531430**
Eubacterium saburreum		100.0	0.38	ARA 1		
53531500	**Very good identification**					**53531500**
Eubacterium saburreum		99.15	0.58	ARA 1		
Bifidobacterium longum subsp. longum		0.85	0.19			
53531502	**Acceptable identification**					**53531502**
Eubacterium saburreum		93.46	0.45	ARA 1		
Bifidobacterium longum subsp. longum		4.53	0.19			
Bifidobacterium adolescentis		2.02	0.15			
53531510	**Very good identification**					**53531510**
Eubacterium saburreum		98.25	0.57	ARA 1		
Bifidobacterium longum subsp. longum		1.75	0.23			
53531512	**Low discrimination**					**53531512**
Eubacterium saburreum		88.26	0.44	ARA 1		
Bifidobacterium longum subsp. longum		8.90	0.23			
Bifidobacterium adolescentis		2.84	0.16			
53531520	**Low discrimination**					**53531520**
Eubacterium saburreum		83.76	0.45	ARA 1		
Bifidobacterium longum subsp. longum		16.24	0.29			
53531530	**Low discrimination**					**53531530**
Eubacterium saburreum		71.25	0.44	ARA 1		
Bifidobacterium longum subsp. longum		28.75	0.33			
53531722	**Good identification**					**53531722**
Bacteroides thetaiotaomicron		99.25	0.37	MNS 99		
Parabacteroides distasonis		0.75	0.16	XYL 96		
53533000	**Good identification**					**53533000**
Eubacterium saburreum		100.0	0.52	MAN 1		
53533002	**Good identification**					**53533002**
Eubacterium saburreum		100.0	0.39	MAN 1		
53533010	**Good identification**					**53533010**
Eubacterium saburreum		100.0	0.50	MAN 1		
53533012	**Good identification**					**53533012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	MAN 1		
53533020	**Good identification**					**53533020**
Eubacterium saburreum		100.0	0.39	MAN 1		
53533030	**Good identification**					**53533030**
Eubacterium saburreum		100.0	0.38	MAN 1		
53533100	**Very good identification**					**53533100**
Eubacterium saburreum		100.0	0.58	MAN 1		
53533102	**Good identification**					**53533102**
Eubacterium saburreum		99.07	0.45	MAN 1		
Propionibacterium propionicum		0.93	0.14			
53533110	**Very good identification**					**53533110**
Eubacterium saburreum		100.0	0.57	MAN 1		
53533112	**Good identification**					**53533112**
Eubacterium saburreum		100.0	0.44	MAN 1		
53533120	**Good identification**					**53533120**
Eubacterium saburreum		100.0	0.45	MAN 1		
53533130	**Good identification**					**53533130**
Eubacterium saburreum		100.0	0.44	MAN 1		
53534000	**Good identification**					**53534000**
Eubacterium saburreum		100.0	0.47	GAL 99		
53534010	**Good identification**					**53534010**
Eubacterium saburreum		100.0	0.46	GAL 99		
53534100	**Good identification**					**53534100**
Eubacterium saburreum		100.0	0.54	GAL 99		
53534102	**Good identification**					**53534102**
Eubacterium saburreum		100.0	0.41	GAL 99		
53534110	**Good identification**					**53534110**
Eubacterium saburreum		100.0	0.52	GAL 99		
53534112	**Good identification**					**53534112**
Eubacterium saburreum		100.0	0.39	GAL 99		
53534120	**Good identification**					**53534120**
Eubacterium saburreum		100.0	0.41	GAL 99		
53534130	**Good identification**					**53534130**
Eubacterium saburreum		100.0	0.39	GAL 99		
53535000	**Very good identification**					**53535000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.82			
53535001	**Good identification**					**53535001**
Eubacterium saburreum		100.0	0.47	URE 1		
53535002	**Very good identification**					**53535002**
Eubacterium saburreum		100.0	0.69			
53535010	**Very good identification**					**53535010**
Eubacterium saburreum		100.0	0.80			
53535011	**Good identification**					**53535011**
Eubacterium saburreum		100.0	0.46	URE 1		
53535012	**Very good identification**					**53535012**
Eubacterium saburreum		100.0	0.67			
53535020	**Very good identification**					**53535020**
Eubacterium saburreum		100.0	0.69			
53535022	**Good identification**					**53535022**
Eubacterium saburreum		100.0	0.56			
53535030	**Very good identification**					**53535030**
Eubacterium saburreum		100.0	0.67			
53535032	**Good identification**					**53535032**
Eubacterium saburreum		100.0	0.54			
53535040	**Good identification**					**53535040**
Eubacterium saburreum		100.0	0.47	SOR 1		
53535050	**Good identification**					**53535050**
Eubacterium saburreum		100.0	0.46	SOR 1		
53535100	**Very good identification**					**53535100**
Eubacterium saburreum		100.0	0.88			
Fusobacterium mortiferum		0.00	0.12			
53535101	**Good identification**					**53535101**
Eubacterium saburreum		100.0	0.54	URE 1		
53535102	**Very good identification**					**53535102**
Eubacterium saburreum		99.96	0.75			
Bifidobacterium adolescentis		0.04	0.15			
53535103	**Good identification**					**53535103**
Eubacterium saburreum		100.0	0.41	URE 1		
53535110	**Very good identification**					**53535110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.86			
53535111	**Good identification**					**53535111**
Eubacterium saburreum		100.0	0.52	URE 1		
53535112	**Very good identification**					**53535112**
Eubacterium saburreum		99.94	0.73			
Bifidobacterium adolescentis		0.06	0.16			
53535113	**Good identification**					**53535113**
Eubacterium saburreum		100.0	0.39	URE 1		
53535120	**Very good identification**					**53535120**
Eubacterium saburreum		100.0	0.75			
53535121	**Good identification**					**53535121**
Eubacterium saburreum		100.0	0.41	URE 1		
53535122	**Very good identification**					**53535122**
					CAT	
Eubacterium saburreum		98.06	0.62		1	
Parabacteroides distasonis		1.00	0.28		76	
Bacteroides thetaiotaomicron		0.49	0.07		89	
53535130	**Very good identification**					**53535130**
Eubacterium saburreum		99.98	0.73			
Bifidobacterium longum subsp. longum		0.02	0.07			
53535131	**Good identification**					**53535131**
Eubacterium saburreum		100.0	0.39	URE 1		
53535132	**Very good identification**					**53535132**
					CAT	
Eubacterium saburreum		98.98	0.61		1	
Parabacteroides distasonis		0.66	0.24		76	
Bacteroides thetaiotaomicron		0.24	0.00		89	
53535140	**Good identification**					**53535140**
Eubacterium saburreum		100.0	0.54	SOR 1		
53535142	**Good identification**					**53535142**
Eubacterium saburreum		100.0	0.41	SOR 1		
53535150	**Good identification**					**53535150**
Eubacterium saburreum		100.0	0.52	SOR 1		
53535152	**Good identification**					**53535152**
Eubacterium saburreum		100.0	0.39	SOR 1		
53535160	**Good identification**					**53535160**
Eubacterium saburreum		100.0	0.41	SOR 1		
53535170	**Good identification**					**53535170**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.39	SOR 1		
53535200	**Good identification**					**53535200**
Eubacterium saburreum		100.0	0.47	RHA 1		
53535210	**Good identification**					**53535210**
Eubacterium saburreum		100.0	0.46	RHA 1		
53535300	**Good identification**					**53535300**
Eubacterium saburreum		100.0	0.54	RHA 1		
53535302	**Good identification**					**53535302**
Eubacterium saburreum		100.0	0.41	RHA 1		
53535310	**Good identification**					**53535310**
Eubacterium saburreum		100.0	0.52	RHA 1		
53535312	**Good identification**					**53535312**
Eubacterium saburreum		100.0	0.39	RHA 1		
53535320	**Good identification**					**53535320**
Eubacterium saburreum		100.0	0.41	RHA 1		
53535330	**Good identification**					**53535330**
Eubacterium saburreum		100.0	0.39	RHA 1		
53535400	**Good identification**					**53535400**
Eubacterium saburreum		100.0	0.47	ARA 1		
53535410	**Good identification**					**53535410**
Eubacterium saburreum		100.0	0.46	ARA 1		
53535500	**Good identification**					**53535500**
Eubacterium saburreum		95.45	0.54	ARA 1		
Bifidobacterium longum subsp. longum		4.55	0.27			
53535502	**Low discrimination**					**53535502**
Eubacterium saburreum		61.53	0.41	ARA 1		
Bifidobacterium longum subsp. longum		16.62	0.27			
Bifidobacterium adolescentis		16.49	0.29			
53535510	**Acceptable identification**					**53535510**
Eubacterium saburreum		90.98	0.52	ARA 1		
Bifidobacterium longum subsp. longum		9.02	0.31			
53535512	**Low discrimination**					**53535512**
Eubacterium saburreum		49.91	0.39	ARA 1		
Bifidobacterium longum subsp. longum		28.05	0.31			
Bifidobacterium adolescentis		19.98	0.30			
53535520	**Low discrimination**					**53535520**

CAT GRA

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		49.48	0.37	IND 1	1	99	
Eubacterium saburreum		45.81	0.41	MNS 87	1	85	
Bacteroides thetaiotaomicron		3.99	0.07		89	1	
53535522	**Low discrimination**						**53535522**
					CAT	GRA	
Bacteroides thetaiotaomicron		86.19	0.41	MNS 99	89	1	
Bifidobacterium longum subsp. longum		10.79	0.37	RHA 93	1	99	
Eubacterium saburreum		1.76	0.28		1	85	
53535530	**Low discrimination**						**53535530**
					CAT	GRA	
Bifidobacterium longum subsp. longum		68.34	0.41	IND 1	1	99	
Eubacterium saburreum		30.40	0.39	MNS 87	1	85	
Bacteroides thetaiotaomicron		1.26	0.00		89	1	
53535722	**Very good identification**						**53535722**
Bacteroides thetaiotaomicron		99.43	0.60	MNS 99			
Bacteroides ovatus		0.29	0.17				
Parabacteroides distasonis		0.28	0.33				
53535732	**Good identification**						**53535732**
Bacteroides thetaiotaomicron		99.02	0.53	MNS 99			
Bacteroides ovatus		0.49	0.14				
Parabacteroides distasonis		0.39	0.28				
53537000	**Good identification**						**53537000**
Eubacterium saburreum		100.0	0.47	MAN 1			
53537010	**Good identification**						**53537010**
Eubacterium saburreum		100.0	0.46	MAN 1			
53537100	**Good identification**						**53537100**
Eubacterium saburreum		99.47	0.54	MAN 1			
Actinomyces israelii		0.53	0.19				
53537102	**Acceptable identification**						**53537102**
Eubacterium saburreum		92.75	0.41	MAN 1			
Actinomyces israelii		4.22	0.22				
Bifidobacterium adolescentis		3.04	0.13				
53537110	**Good identification**						**53537110**
Eubacterium saburreum		100.0	0.52	MAN 1			
53537112	**Good identification**						**53537112**
Eubacterium saburreum		95.33	0.39	MAN 1			
Bifidobacterium adolescentis		4.67	0.15				
53537120	**Good identification**						**53537120**
Eubacterium saburreum		100.0	0.41	MAN 1			
53537130	**Good identification**						**53537130**
Eubacterium saburreum		100.0	0.39	MAN 1			
53541000	**Good identification**						**53541000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.47	CEL 1		
53541010	**Good identification**					**53541010**
Eubacterium saburreum		100.0	0.45	CEL 1		
53541100	**Good identification**					**53541100**
					COC	
Eubacterium saburreum		99.54	0.53	CEL 1	1	
Blautia hansenii		0.47	-.03		99	
53541102	**Good identification**					**53541102**
Eubacterium saburreum		100.0	0.40	CEL 1		
53541110	**Good identification**					**53541110**
Eubacterium saburreum		100.0	0.52	CEL 1		
53541112	**Good identification**					**53541112**
Eubacterium saburreum		100.0	0.39	CEL 1		
53541120	**Low discrimination**					**53541120**
Eubacterium saburreum		80.24	0.40	CEL 1		
Prevotella melaninogenica		19.76	0.14			
53541130	**Good identification**					**53541130**
Eubacterium saburreum		100.0	0.39	CEL 1		
53545000	**Good identification**					**53545000**
Eubacterium saburreum		100.0	0.42	CEL 1		
53545010	**Good identification**					**53545010**
Eubacterium saburreum		100.0	0.41	CEL 1		
53545100	**Good identification**					**53545100**
Eubacterium saburreum		100.0	0.49	CEL 1		
53545102	**Good identification**					**53545102**
Eubacterium saburreum		100.0	0.36	CEL 1		
53545110	**Good identification**					**53545110**
Eubacterium saburreum		100.0	0.47	CEL 1		
53545120	**Good identification**					**53545120**
Eubacterium saburreum		100.0	0.36	CEL 1		
53551000	**Good identification**					**53551000**
Eubacterium saburreum		97.10	0.55	CEL 1		
Fusobacterium mortiferum		1.58	0.26			
Prevotella intermedia		1.32	0.14			
53551002	**Good identification**					**53551002**
Eubacterium saburreum		100.0	0.42	CEL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53551010	**Good identification**						**53551010**
Eubacterium saburreum		100.0	0.53	CEL 1			
53551012	**Good identification**						**53551012**
Eubacterium saburreum		100.0	0.41	CEL 1			
53551020	**Good identification**						**53551020**
Eubacterium saburreum		98.29	0.42	CEL 1			
Prevotella melaninogenica		1.71	-.04				
53551030	**Good identification**						**53551030**
Eubacterium saburreum		100.0	0.41	CEL 1			
53551100	**Very good identification**						**53551100**
					CAT		
Eubacterium saburreum		98.07	0.61	CEL 1	1		
Fusobacterium mortiferum		1.59	0.32		5		
Capnocytophaga ochracea		0.21	0.04		90		
53551102	**Low discrimination**						**53551102**
					CAT		
Eubacterium saburreum		87.35	0.48	CEL 1	1		
Capnocytophaga ochracea		9.69	0.20		90		
Prevotella oralis		1.46	0.06		1		
53551110	**Very good identification**						**53551110**
Eubacterium saburreum		100.0	0.60	CEL 1			
53551112	**Good identification**						**53551112**
					CAT	GRA	
Eubacterium saburreum		97.53	0.47	CEL 1	1	85	
Capnocytophaga ochracea		1.47	0.04		90	1	
Bifidobacterium adolescentis		1.00	0.11		1	99	
53551120	**Low discrimination**						**53551120**
					CAT		
Eubacterium saburreum		53.57	0.48	CEL 1	1		
Prevotella melaninogenica		39.59	0.30		1		
Capnocytophaga ochracea		5.95	0.20		90		
53551122	**Low discrimination**						**53551122**
					CAT		
Capnocytophaga ochracea		60.55	0.36	IND 1	90		
Prevotella oralis		19.30	0.27	MNS 90	1		
Eubacterium saburreum		10.70	0.35		1		
53551130	**Good identification**						**53551130**
					CAT		
Eubacterium saburreum		97.64	0.47	CEL 1	1		
Capnocytophaga ochracea		1.47	0.04		90		
Prevotella melaninogenica		0.89	-.04		1		
53555000	**Good identification**						**53555000**
Eubacterium saburreum		100.0	0.50	CEL 1			
53555002	**Good identification**						**53555002**
Eubacterium saburreum		100.0	0.37	CEL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53555010	**Good identification**						**53555010**
Eubacterium saburreum		100.0	0.49	CEL 1			
53555012	**Good identification**						**53555012**
Eubacterium saburreum		100.0	0.36	CEL 1			
53555020	**Good identification**						**53555020**
Eubacterium saburreum		100.0	0.37	CEL 1			
53555030	**Good identification**						**53555030**
Eubacterium saburreum		100.0	0.36	CEL 1			
53555100	**Very good identification**						**53555100**
					GRA		
Eubacterium saburreum		99.55	0.57	CEL 1	85		
Fusobacterium mortiferum		0.23	0.13		1		
Actinomyces israelii		0.22	0.15		99		
53555102	**Low discrimination**						**53555102**
					CAT	GRA	
Eubacterium saburreum		84.48	0.44	CEL 1	1	85	
Bifidobacterium adolescentis		7.19	0.23		1	99	
Capnocytophaga ochracea		4.35	0.10		90	1	
53555110	**Good identification**						**53555110**
Eubacterium saburreum		100.0	0.55	CEL 1			
53555112	**Low discrimination**						**53555112**
Eubacterium saburreum		88.72	0.42	CEL 1			
Bifidobacterium adolescentis		11.28	0.25				
53555120	**Low discrimination**						**53555120**
					CAT		
Eubacterium saburreum		86.39	0.44	CEL 1	1		
Bacteroides fragilis		6.83	0.04		99		
Capnocytophaga ochracea		4.45	0.10		90		
53555130	**Good identification**						**53555130**
Eubacterium saburreum		100.0	0.42	CEL 1			
53555502	**Low discrimination**						**53555502**
					GRA		
Bifidobacterium adolescentis		76.44	0.37	IND 1	99		
Bacteroides uniformis		20.13	0.11	MNS 87	1		
Actinomyces israelii		2.07	0.17		99		
53555512	**Good identification**						**53555512**
Bifidobacterium adolescentis		98.25	0.39	IND 1			
Bifidobacterium longum subsp. longum		1.75	0.07	MNS 87			
53555522	**Acceptable identification**						**53555522**
					CAT		
Bacteroides uniformis		92.75	0.45	MNS 99	5		
Bacteroides thetaiotaomicron		5.75	0.27	SAL 92	89		
Bacteroides ovatus		0.71	0.12		68		
53555722	**Low discrimination**						**53555722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Bacteroides uniformis		58.71	0.48	MNS 99	5	
Bacteroides thetaiotaomicron		35.00	0.47	SAL 92	89	
Bacteroides ovatus		6.21	0.34		68	
53555732	**Low discrimination**					**53555732**
					CAT	
Bacteroides thetaiotaomicron		74.03	0.40	MNS 99	89	
Bacteroides ovatus		22.53	0.31	TRE 93	68	
Bacteroides uniformis		3.23	0.14		5	
53557502	**Low discrimination**					**53557502**
Bifidobacterium adolescentis		84.18	0.36	IND 1		
Actinomyces israelii		15.82	0.30	MNS 87		
53557512	**Good identification**					**53557512**
Bifidobacterium adolescentis		97.75	0.37	IND 1		
Actinomyces israelii		2.25	0.16	MNS 87		
53561000	**Good identification**					**53561000**
Eubacterium saburreum		100.0	0.44	CEL 1		
53561010	**Good identification**					**53561010**
Eubacterium saburreum		100.0	0.42	CEL 1		
53561100	**Good identification**					**53561100**
Eubacterium saburreum		100.0	0.50	CEL 1		
53561102	**Good identification**					**53561102**
Eubacterium saburreum		100.0	0.37	CEL 1		
53561110	**Good identification**					**53561110**
Eubacterium saburreum		100.0	0.49	CEL 1		
53561112	**Good identification**					**53561112**
Eubacterium saburreum		100.0	0.36	CEL 1		
53561120	**Good identification**					**53561120**
Eubacterium saburreum		100.0	0.37	CEL 1		
53561130	**Good identification**					**53561130**
Eubacterium saburreum		100.0	0.36	CEL 1		
53565000	**Good identification**					**53565000**
Eubacterium saburreum		100.0	0.39	CEL 1		
53565010	**Good identification**					**53565010**
Eubacterium saburreum		100.0	0.38	CEL 1		
53565100	**Good identification**					**53565100**
Eubacterium saburreum		100.0	0.46	CEL 1		
53565110	**Good identification**					**53565110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.44	CEL 1		
53571000	**Good identification**					**53571000**
Eubacterium saburreum		99.28	0.52	CEL 1		
Fusobacterium mortiferum		0.72	0.17			
53571002	**Good identification**					**53571002**
Eubacterium saburreum		100.0	0.39	CEL 1		
53571010	**Good identification**					**53571010**
Eubacterium saburreum		100.0	0.50	CEL 1		
53571012	**Good identification**					**53571012**
Eubacterium saburreum		100.0	0.38	CEL 1		
53571020	**Good identification**					**53571020**
Eubacterium saburreum		100.0	0.39	CEL 1		
53571030	**Good identification**					**53571030**
Eubacterium saburreum		100.0	0.38	CEL 1		
53571100	**Very good identification**					**53571100**
Eubacterium saburreum		99.28	0.58	CEL 1		
Fusobacterium mortiferum		0.72	0.23			
53571102	**Good identification**					**53571102**
Eubacterium saburreum		100.0	0.45	CEL 1		
53571110	**Very good identification**					**53571110**
Eubacterium saburreum		100.0	0.57	CEL 1		
53571112	**Good identification**					**53571112**
					GRA	
Eubacterium saburreum		97.92	0.44	CEL 1	85	
Leptotrichia buccalis		1.08	-.11		1	
Bifidobacterium adolescentis		1.00	0.08		99	
53571120	**Good identification**					**53571120**
Eubacterium saburreum		98.89	0.45	CEL 1		
Prevotella melaninogenica		1.11	-.04			
53571130	**Good identification**					**53571130**
Eubacterium saburreum		100.0	0.44	CEL 1		
53571722	**Good identification**					**53571722**
Bacteroides thetaiotaomicron		96.65	0.42	MNS 99		
Bacteroides ovatus		3.17	0.17	XYL 96		
Parabacteroides distasonis		0.19	0.12			
53571732	**Acceptable identification**					**53571732**
Bacteroides thetaiotaomicron		94.68	0.35	MNS 99		
Bacteroides ovatus		5.32	0.14	XYL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
53575000	**Good identification**						**53575000**
Eubacterium saburreum		99.27	0.47	CEL 1			
Actinomyces israelii		0.73	0.15				
53575010	**Good identification**						**53575010**
Eubacterium saburreum		100.0	0.46	CEL 1			
53575100	**Good identification**						**53575100**
Eubacterium saburreum		98.76	0.54	CEL 1			
Actinomyces israelii		1.24	0.25				
53575102	**Low discrimination**						**53575102**
Eubacterium saburreum		83.92	0.41	CEL 1			
Actinomyces israelii		8.94	0.28				
Bifidobacterium adolescentis		7.14	0.20				
53575110	**Good identification**						**53575110**
Eubacterium saburreum		100.0	0.52	CEL 1			
53575112	**Low discrimination**						**53575112**
Eubacterium saburreum		87.22	0.39	CEL 1			
Bifidobacterium adolescentis		11.09	0.22				
Actinomyces israelii		1.70	0.14				
53575120	**Good identification**						**53575120**
Eubacterium saburreum		100.0	0.41	CEL 1			
53575130	**Good identification**						**53575130**
Eubacterium saburreum		100.0	0.39	CEL 1			
53575512	**Acceptable identification**						**53575512**
					GRA	CAT	
Bifidobacterium adolescentis		90.62	0.36	IND 1	99	1	
Bacteroides thetaiotaomicron		9.38	0.05	MNS 87	1	89	
53575522	**Acceptable identification**						**53575522**
					CAT		
Bacteroides thetaiotaomicron		90.10	0.47	MNS 99	89		
Bacteroides ovatus		8.51	0.29	RHA 93	68		
Bacteroides uniformis		1.11	0.11		5		
53575532	**Low discrimination**						**53575532**
Bacteroides thetaiotaomicron		85.55	0.40	MNS 99			
Bacteroides ovatus		13.85	0.26	RHA 93			
Parabacteroides distasonis		0.38	0.15				
53575622	**Acceptable identification**						**53575622**
Bacteroides thetaiotaomicron		93.73	0.37	MNS 99			
Bacteroides ovatus		6.27	0.17	LAC 98			
53575722	**Low discrimination**						**53575722**
					CAT		
Bacteroides thetaiotaomicron		87.94	0.66	MNS 99	89		
Bacteroides ovatus		11.88	0.51		68		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		0.11	0.14		5	
53575732	**Low discrimination**					**53575732**
Bacteroides thetaiotaomicron		81.07	0.59	MNS 99		
Bacteroides ovatus		18.78	0.48			
Parabacteroides distasonis		0.08	0.24			
53577100	**Low discrimination**					**53577100**
Actinomyces israelii		87.55	0.38	IND 1		
Eubacterium saburreum		12.45	0.20	SAL 92		
53577102	**Low discrimination**					**53577102**
Actinomyces israelii		89.66	0.41	IND 1		
Bifidobacterium adolescentis		10.34	0.19	SAL 92		
53577500	**Good identification**					**53577500**
Actinomyces israelii		100.0	0.37	IND 1		
53577502	**Low discrimination**					**53577502**
Actinomyces israelii		51.46	0.40	IND 1		
Bifidobacterium adolescentis		48.54	0.33	SAL 92		
53577722	**Low discrimination**					**53577722**
Bacteroides ovatus		81.68	0.43	MNS 99		
Bacteroides thetaiotaomicron		18.32	0.32	SAL 88		
53577732	**Low discrimination**					**53577732**
Bacteroides ovatus		88.43	0.40	MNS 99		
Bacteroides thetaiotaomicron		11.57	0.25	SAL 88		
53601000	**Very good identification**					**53601000**
					GRA	
Eubacterium tortuosum		99.17	0.58	IND 1	99	
Eubacterium saburreum		0.76	0.37		85	
Fusobacterium mortiferum		0.07	0.20		1	
53601002	**Good identification**					**53601002**
Eubacterium tortuosum		99.24	0.45	IND 1		
Eubacterium saburreum		0.76	0.24			
53601020	**Good identification**					**53601020**
Eubacterium tortuosum		99.24	0.45	IND 1		
Eubacterium saburreum		0.76	0.24			
53601100	**Low discrimination**					**53601100**
					GRA	
Eubacterium saburreum		60.31	0.43	SAL 1	85	
Eubacterium tortuosum		34.11	0.24		99	
Fusobacterium mortiferum		5.59	0.27		1	
53601110	**Good identification**					**53601110**
Eubacterium saburreum		100.0	0.41	SAL 1		
53605100	**Good identification**					**53605100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	SAL 1		
53605110	**Good identification**					**53605110**
Eubacterium saburreum		100.0	0.37	SAL 1		
53610000	**Good identification**					**53610000**
Fusobacterium mortiferum		99.51	0.52	IND 1		
Eubacterium saburreum		0.49	0.11			
53610002	**Good identification**					**53610002**
Fusobacterium mortiferum		100.0	0.35	IND 1		
53610100	**Very good identification**					**53610100**
Fusobacterium mortiferum		99.51	0.58	IND 1		
Eubacterium saburreum		0.49	0.17			
53610102	**Good identification**					**53610102**
Fusobacterium mortiferum		100.0	0.42	IND 1		
53611000	**Low discrimination**					**53611000**
					GRA	
Fusobacterium mortiferum		67.98	0.55	IND 1	1	
Eubacterium saburreum		22.24	0.45		85	
Eubacterium tortuosum		9.78	0.24		99	
53611002	**Low discrimination**					**53611002**
					GRA	
Fusobacterium mortiferum		57.20	0.38	IND 1	1	
Eubacterium saburreum		29.73	0.32	BGL 10	85	
Eubacterium tortuosum		13.07	0.11		99	
53611010	**Good identification**					**53611010**
Eubacterium saburreum		96.36	0.43	SAL 1		
Fusobacterium mortiferum		3.64	0.20			
53611100	**Low discrimination**					**53611100**
Fusobacterium mortiferum		75.35	0.61	IND 1		
Eubacterium saburreum		24.65	0.51			
53611102	**Low discrimination**					**53611102**
					GRA	
Fusobacterium mortiferum		65.35	0.45	IND 1	1	
Eubacterium saburreum		33.96	0.38	BGL 10	85	
Bifidobacterium longum subsp. infantis		0.69	-.07		99	
53611110	**Good identification**					**53611110**
Eubacterium saburreum		96.36	0.49	SAL 1		
Fusobacterium mortiferum		3.64	0.27			
53611112	**Good identification**					**53611112**
Eubacterium saburreum		100.0	0.37	SAL 1		
53611120	**Low discrimination**					**53611120**
Eubacterium saburreum		85.11	0.38	SAL 1		
Fusobacterium mortiferum		14.89	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53611130	**Good identification**					**53611130**
Eubacterium saburreum		100.0	0.37	SAL 1		
53614100	**Good identification**					**53614100**
Fusobacterium mortiferum		96.57	0.39	IND 1		
Eubacterium saburreum		3.43	0.12	XYL 7		
53615000	**Low discrimination**					**53615000**
Eubacterium saburreum		70.07	0.40	SAL 1		
Fusobacterium mortiferum		29.93	0.35			
53615010	**Good identification**					**53615010**
Eubacterium saburreum		100.0	0.39	SAL 1		
53615100	**Low discrimination**					**53615100**
Eubacterium saburreum		70.07	0.46	SAL 1		
Fusobacterium mortiferum		29.93	0.42			
53615110	**Good identification**					**53615110**
Eubacterium saburreum		100.0	0.45	SAL 1		
53621100	**Good identification**					**53621100**
Eubacterium saburreum		96.02	0.40	SAL 1		
Fusobacterium mortiferum		3.98	0.18			
53621110	**Good identification**					**53621110**
Eubacterium saburreum		100.0	0.38	SAL 1		
53625100	**Good identification**					**53625100**
Eubacterium saburreum		100.0	0.35	SAL 1		
53630000	**Good identification**					**53630000**
Fusobacterium mortiferum		100.0	0.43	IND 1		
53630100	**Good identification**					**53630100**
Fusobacterium mortiferum		98.91	0.49	IND 1		
Eubacterium saburreum		1.10	0.14			
53631000	**Low discrimination**					**53631000**
Fusobacterium mortiferum		57.79	0.46	IND 1		
Eubacterium saburreum		42.21	0.42			
53631010	**Good identification**					**53631010**
Eubacterium saburreum		98.34	0.40	SAL 1		
Fusobacterium mortiferum		1.66	0.12			
53631100	**Low discrimination**					**53631100**
Fusobacterium mortiferum		57.79	0.52	IND 1		
Eubacterium saburreum		42.21	0.48			
53631102	**Low discrimination**					**53631102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		53.70	0.35	SAL 1		
Fusobacterium mortiferum		46.30	0.36			
53631110	**Good identification**					**53631110**
Eubacterium saburreum		98.34	0.46	SAL 1		
Fusobacterium mortiferum		1.66	0.18			
53631120	**Acceptable identification**					**53631120**
Eubacterium saburreum		92.73	0.35	SAL 1		
Fusobacterium mortiferum		7.27	0.18			
53635000	**Low discrimination**					**53635000**
Eubacterium saburreum		83.93	0.37	SAL 1		
Fusobacterium mortiferum		16.07	0.26			
53635010	**Good identification**					**53635010**
Eubacterium saburreum		100.0	0.36	SAL 1		
53635100	**Low discrimination**					**53635100**
Eubacterium saburreum		83.93	0.43	SAL 1		
Fusobacterium mortiferum		16.07	0.33			
53635110	**Good identification**					**53635110**
Eubacterium saburreum		100.0	0.42	SAL 1		
53641000	**Good identification**					**53641000**
					GRA	
Eubacterium tortuosum		99.79	0.41	IND 1	99	
Fusobacterium mortiferum		0.21	0.12	CEL 10	1	
53650000	**Good identification**					**53650000**
Fusobacterium mortiferum		100.0	0.43	IND 1		
53650100	**Good identification**					**53650100**
Fusobacterium mortiferum		100.0	0.50	IND 1		
53651000	**Acceptable identification**					**53651000**
					GRA	
Fusobacterium mortiferum		94.53	0.46	IND 1	1	
Eubacterium tortuosum		4.54	0.07		99	
Eubacterium saburreum		0.94	0.11		85	
53651100	**Good identification**					**53651100**
Fusobacterium mortiferum		99.02	0.53	IND 1		
Eubacterium saburreum		0.98	0.17			
53651102	**Good identification**					**53651102**
Fusobacterium mortiferum		100.0	0.36	IND 1		
53655522	**Good identification**					**53655522**
					CAT	
Bacteroides uniformis		99.93	0.45	MNS 99	5	
Bacteroides thetaiotaomicron		0.07	-.06	MLT 92	89	
53655722	**Good identification**					**53655722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Bacteroides uniformis		98.57	0.48	MNS 99	5		
Bacteroides ovatus		0.77	0.15	MLT 92	68		
Bacteroides thetaiotaomicron		0.66	0.13		89		
53670100	**Good identification**						**53670100**
Fusobacterium mortiferum		100.0	0.41	IND 1			
53671000	**Good identification**						**53671000**
Fusobacterium mortiferum		100.0	0.38	IND 1			
53671100	**Good identification**						**53671100**
Fusobacterium mortiferum		97.84	0.44	IND 1			
Eubacterium saburreum		2.17	0.14				
53701000	**Low discrimination**						**53701000**
Eubacterium saburreum		75.20	0.47	SAL 1			
Eubacterium tortuosum		24.81	0.24				
53701010	**Good identification**						**53701010**
Eubacterium saburreum		100.0	0.45	SAL 1			
53701100	**Good identification**						**53701100**
					COC	GRA	
Eubacterium saburreum		99.25	0.53	SAL 1	1	85	
Blautia hansenii		0.46	-.03		99	99	
Fusobacterium mortiferum		0.28	0.11		1	1	
53701102	**Good identification**						**53701102**
Eubacterium saburreum		98.49	0.40	SAL 1			
Bifidobacterium longum subsp. infantis		1.51	-.07				
53701110	**Good identification**						**53701110**
Eubacterium saburreum		100.0	0.52	SAL 1			
53701112	**Good identification**						**53701112**
Eubacterium saburreum		100.0	0.39	SAL 1			
53701120	**Good identification**						**53701120**
Eubacterium saburreum		97.81	0.40	SAL 1			
Prevotella melaninogenica		2.19	-.04				
53701130	**Good identification**						**53701130**
Eubacterium saburreum		100.0	0.39	SAL 1			
53705000	**Good identification**						**53705000**
Eubacterium saburreum		100.0	0.42	SAL 1			
53705010	**Good identification**						**53705010**
Eubacterium saburreum		100.0	0.41	SAL 1			
53705100	**Good identification**						**53705100**
Eubacterium saburreum		100.0	0.49	SAL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
53705102	**Good identification**					**53705102**
Eubacterium saburreum		100.0	0.36	SAL 1		
53705110	**Good identification**					**53705110**
Eubacterium saburreum		100.0	0.47	SAL 1		
53705120	**Good identification**					**53705120**
Eubacterium saburreum		100.0	0.36	SAL 1		
53710000	**Low discrimination**					**53710000**
Fusobacterium mortiferum		72.64	0.36	IND 1		
Prevotella intermedia		15.71	0.14	MLT 11		
Eubacterium saburreum		11.65	0.21			
53710100	**Low discrimination**					**53710100**
Fusobacterium mortiferum		86.17	0.42	IND 1		
Eubacterium saburreum		13.83	0.27	MLT 11		
53711000	**Acceptable identification**					**53711000**
Eubacterium saburreum		90.25	0.55	SAL 1		
Fusobacterium mortiferum		8.52	0.39			
Prevotella intermedia		1.23	0.14			
53711002	**Acceptable identification**					**53711002**
					GRA	
Eubacterium saburreum		93.34	0.42	SAL 1	85	
Fusobacterium mortiferum		5.55	0.23		1	
Bifidobacterium longum subsp. infantis		1.11	-.07		99	
53711010	**Good identification**					**53711010**
Eubacterium saburreum		100.0	0.53	SAL 1		
53711012	**Good identification**					**53711012**
Eubacterium saburreum		100.0	0.41	SAL 1		
53711020	**Good identification**					**53711020**
Eubacterium saburreum		100.0	0.42	SAL 1		
53711030	**Good identification**					**53711030**
Eubacterium saburreum		100.0	0.41	SAL 1		
53711100	**Good identification**					**53711100**
					GRA	
Eubacterium saburreum		90.64	0.61	SAL 1	85	
Fusobacterium mortiferum		8.56	0.45		1	
Bifidobacterium longum subsp. infantis		0.80	0.10		99	
53711102	**Low discrimination**					**53711102**
					GRA	
Eubacterium saburreum		62.80	0.48	SAL 1	85	
Bifidobacterium longum subsp. infantis		31.79	0.27		99	
Fusobacterium mortiferum		3.73	0.29		1	
53711110	**Very good identification**					**53711110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.88	0.60	SAL 1		
Fusobacterium mortiferum		0.12	0.11			
53711112	**Good identification**					**53711112**
Eubacterium saburreum		96.76	0.47	SAL 1		
Bifidobacterium adolescentis		2.64	0.18			
Bifidobacterium longum subsp. infantis		0.60	-.07			
53711120	**Low discrimination**					**53711120**
					GRA	
Eubacterium saburreum		85.47	0.48	SAL 1	85	
Bifidobacterium longum subsp. infantis		7.61	0.15		99	
Prevotella melaninogenica		5.74	0.12		1	
53711130	**Good identification**					**53711130**
Eubacterium saburreum		100.0	0.47	SAL 1		
53715000	**Good identification**					**53715000**
Eubacterium saburreum		98.70	0.50	SAL 1		
Fusobacterium mortiferum		1.30	0.20			
53715002	**Good identification**					**53715002**
Eubacterium saburreum		100.0	0.37	SAL 1		
53715010	**Good identification**					**53715010**
Eubacterium saburreum		100.0	0.49	SAL 1		
53715012	**Good identification**					**53715012**
Eubacterium saburreum		100.0	0.36	SAL 1		
53715020	**Good identification**					**53715020**
Eubacterium saburreum		100.0	0.37	SAL 1		
53715030	**Good identification**					**53715030**
Eubacterium saburreum		100.0	0.36	SAL 1		
53715100	**Good identification**					**53715100**
Eubacterium saburreum		90.07	0.57	SAL 1		
Eubacterium contortum		7.36	0.27			
Bifidobacterium longum subsp. infantis		1.21	0.09			
53715102	**Low discrimination**					**53715102**
Eubacterium saburreum		47.76	0.44	SAL 1		
Bifidobacterium longum subsp. infantis		36.73	0.26			
Bifidobacterium adolescentis		10.84	0.31			
53715110	**Good identification**					**53715110**
Eubacterium saburreum		100.0	0.55	SAL 1		
53715112	**Low discrimination**					**53715112**
Eubacterium saburreum		74.69	0.42	SAL 1		
Bifidobacterium adolescentis		25.31	0.32			
53715120	**Low discrimination**					**53715120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		82.18	0.44	SAL 1		
Bifidobacterium longum subsp. infantis		11.11	0.13			
Eubacterium contortum		6.71	0.14			
53715130	**Good identification**					**53715130**
Eubacterium saburreum		100.0	0.42	SAL 1		
53715300	**Very good identification**					**53715300**
Eubacterium contortum		99.88	0.61	IND 1		
Eubacterium saburreum		0.13	0.23			
53715302	**Good identification**					**53715302**
Eubacterium contortum		99.88	0.48	IND 1		
Eubacterium saburreum		0.13	0.10			
53715320	**Good identification**					**53715320**
					GRA	
Eubacterium contortum		99.36	0.48	IND 1	99	
Bacteroides vulgatus		0.52	0.02		1	
Eubacterium saburreum		0.12	0.10		85	
53715322	**Low discrimination**					**53715322**
					GRA	
Eubacterium contortum		81.17	0.35	IND 1	99	
Bacteroides uniformis		11.78	0.14		1	
Parabacteroides distasonis		4.80	0.26		1	
53715502	**Acceptable identification**					**53715502**
					GRA	
Bifidobacterium adolescentis		93.38	0.45	IND 1	99	
Eubacterium contortum		4.11	0.13	MNS 87	99	
Bacteroides uniformis		1.07	-.05		1	
53715512	**Good identification**					**53715512**
Bifidobacterium adolescentis		99.34	0.46	IND 1		
Bifidobacterium longum subsp. longum		0.66	0.07	MNS 87		
53715700	**Very good identification**					**53715700**
Eubacterium contortum		100.0	0.60	IND 1		
53715702	**Good identification**					**53715702**
					GRA	
Eubacterium contortum		99.26	0.47	IND 1	99	
Bacteroides uniformis		0.36	-.03		1	
Bifidobacterium adolescentis		0.23	0.11		99	
53715720	**Acceptable identification**					**53715720**
					GRA	
Eubacterium contortum		91.72	0.47	IND 1	99	
Bacteroides vulgatus		7.81	0.21		1	
Bacteroides uniformis		0.33	-.03		1	
53717502	**Acceptable identification**					**53717502**
Bifidobacterium adolescentis		94.25	0.43	IND 1		
Actinomyces israelii		5.75	0.29	MNS 87		
53717512	**Good identification**					**53717512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		99.26	0.45	IND 1		
Actinomyces israelii		0.74	0.15	MNS 87		
53721000	**Good identification**					**53721000**
Eubacterium saburreum		100.0	0.44	SAL 1		
53721010	**Good identification**					**53721010**
Eubacterium saburreum		100.0	0.42	SAL 1		
53721100	**Good identification**					**53721100**
Eubacterium saburreum		100.0	0.50	SAL 1		
53721102	**Good identification**					**53721102**
Eubacterium saburreum		100.0	0.37	SAL 1		
53721110	**Good identification**					**53721110**
Eubacterium saburreum		100.0	0.49	SAL 1		
53721112	**Good identification**					**53721112**
Eubacterium saburreum		100.0	0.36	SAL 1		
53721120	**Good identification**					**53721120**
Eubacterium saburreum		100.0	0.37	SAL 1		
53721130	**Good identification**					**53721130**
Eubacterium saburreum		100.0	0.36	SAL 1		
53725000	**Good identification**					**53725000**
Eubacterium saburreum		100.0	0.39	SAL 1		
53725010	**Good identification**					**53725010**
Eubacterium saburreum		100.0	0.38	SAL 1		
53725100	**Good identification**					**53725100**
Eubacterium saburreum		100.0	0.46	SAL 1		
53725110	**Good identification**					**53725110**
Eubacterium saburreum		100.0	0.44	SAL 1		
53731000	**Good identification**					**53731000**
Eubacterium saburreum		95.94	0.52	SAL 1		
Fusobacterium mortiferum		4.06	0.30			
53731002	**Good identification**					**53731002**
Eubacterium saburreum		97.41	0.39	SAL 1		
Fusobacterium mortiferum		2.60	0.14			
53731010	**Good identification**					**53731010**
Eubacterium saburreum		100.0	0.50	SAL 1		
53731012	**Good identification**					**53731012**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	SAL 1		
53731020	**Good identification**					**53731020**
Eubacterium saburreum		100.0	0.39	SAL 1		
53731030	**Good identification**					**53731030**
Eubacterium saburreum		100.0	0.38	SAL 1		
53731100	**Very good identification**					**53731100**
					GRA	
Eubacterium saburreum		95.73	0.58	SAL 1	85	
Fusobacterium mortiferum		4.05	0.36		1	
Actinomyces naeslundii		0.22	0.26		99	
53731102	**Acceptable identification**					**53731102**
					GRA	
Eubacterium saburreum		93.87	0.45	SAL 1	85	
Fusobacterium mortiferum		2.50	0.20		1	
Bifidobacterium adolescentis		1.71	0.13		99	
53731110	**Very good identification**					**53731110**
Eubacterium saburreum		100.0	0.57	SAL 1		
53731112	**Good identification**					**53731112**
					GRA	
Eubacterium saburreum		96.32	0.44	SAL 1	85	
Bifidobacterium adolescentis		2.63	0.15		99	
Leptotrichia buccalis		1.06	-.11		1	
53731120	**Good identification**					**53731120**
Eubacterium saburreum		98.74	0.45	SAL 1		
Actinomyces naeslundii		1.26	0.26			
53731130	**Good identification**					**53731130**
Eubacterium saburreum		100.0	0.44	SAL 1		
53735000	**Good identification**					**53735000**
					GRA	
Eubacterium saburreum		98.79	0.47	SAL 1	85	
Actinomyces israelii		0.63	0.14		99	
Fusobacterium mortiferum		0.58	0.11		1	
53735010	**Good identification**					**53735010**
Eubacterium saburreum		100.0	0.46	SAL 1		
53735100	**Good identification**					**53735100**
					GRA	
Eubacterium saburreum		98.35	0.54	SAL 1	85	
Actinomyces israelii		1.07	0.24		99	
Fusobacterium mortiferum		0.58	0.17		1	
53735102	**Low discrimination**					**53735102**
Eubacterium saburreum		75.81	0.41	SAL 1		
Bifidobacterium adolescentis		17.20	0.28			
Actinomyces israelii		6.99	0.27			
53735110	**Good identification**					**53735110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.52	SAL 1		
53735112	**Low discrimination**					**53735112**
Eubacterium saburreum		74.69	0.39	SAL 1		
Bifidobacterium adolescentis		25.31	0.29			
53735120	**Good identification**					**53735120**
Eubacterium saburreum		100.0	0.41	SAL 1		
53735130	**Good identification**					**53735130**
Eubacterium saburreum		100.0	0.39	SAL 1		
53735322	**Low discrimination**					**53735322**
Parabacteroides distasonis		65.67	0.37	IND 1		
Bacteroides thetaiotaomicron		23.28	0.13	MNS 98		
Bacteroides ovatus		8.59	0.06			
53735502	**Acceptable identification**					**53735502**
					SPO	
Bifidobacterium adolescentis		92.35	0.42	IND 1	1	
Actinomyces israelii		4.59	0.26	MNS 87	1	
Clostridium butyricum		2.13	-.01		99	
53735512	**Good identification**					**53735512**
Bifidobacterium adolescentis		98.68	0.43	IND 1		
Clostridium clostridioforme		1.32	0.08	MNS 87		
53735722	**Low discrimination**					**53735722**
Bacteroides thetaiotaomicron		87.52	0.47	MNS 99		
Bacteroides ovatus		10.55	0.32			
Parabacteroides distasonis		1.53	0.33			
53735732	**Low discrimination**					**53735732**
Bacteroides thetaiotaomicron		79.75	0.40	MNS 99		
Bacteroides ovatus		16.47	0.29			
Parabacteroides distasonis		1.93	0.29			
53737100	**Low discrimination**					**53737100**
Actinomyces israelii		85.89	0.37	IND 1		
Eubacterium saburreum		14.11	0.20	CEL 93		
53737102	**Low discrimination**					**53737102**
Actinomyces israelii		73.79	0.40	IND 1		
Bifidobacterium adolescentis		26.21	0.26	CEL 93		
53737500	**Good identification**					**53737500**
Actinomyces israelii		100.0	0.35	IND 1		
53737502	**Low discrimination**					**53737502**
Bifidobacterium adolescentis		74.40	0.40	IND 1		
Actinomyces israelii		25.60	0.39	MNS 87		
53737512	**Acceptable identification**					**53737512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		94.43	0.42	IND 1		
Actinomyces israelii		3.97	0.24	MNS 87		
Bifidobacterium dentium		1.59	-.07			
53751100	**Low discrimination**					**53751100**
					GRA	
Fusobacterium mortiferum		63.94	0.37	IND 1	1	
Eubacterium saburreum		20.52	0.27	MLT 11	85	
Collinsella aerofaciens		8.50	-.03		99	
53751502	**Acceptable identification**					**53751502**
					GRA	
Bifidobacterium adolescentis		93.62	0.36	IND 1	99	
Bacteroides uniformis		3.38	-.05	MNS 87	1	
Actinomyces israelii		3.00	0.17	XYL 87	99	
53751512	**Good identification**					**53751512**
Bifidobacterium adolescentis		100.0	0.38	IND 1		
53753512	**Good identification**					**53753512**
Bifidobacterium adolescentis		97.36	0.36	IND 1		
Actinomyces israelii		2.64	0.16	MNS 87		
53755102	**Low discrimination**					**53755102**
					GRA	
Bifidobacterium adolescentis		55.06	0.36	IND 1	99	
Actinomyces israelii		24.80	0.36	MNS 87	99	
Bacteroides uniformis		17.12	0.11	ARA 87	1	
53755112	**Acceptable identification**					**53755112**
Bifidobacterium adolescentis		94.78	0.38	IND 1		
Actinomyces israelii		5.22	0.22	MNS 87		
53755122	**Good identification**					**53755122**
Bacteroides uniformis		98.58	0.45	MNS 99		
Prevotella oralis		0.86	0.16	ARA 92		
Bacteroides ovatus		0.17	0.01			
53755322	**Good identification**					**53755322**
Bacteroides uniformis		97.02	0.48	MNS 99		
Bacteroides ovatus		2.33	0.23	ARA 92		
Prevotella oralis		0.26	0.10			
53755502	**Low discrimination**					**53755502**
					GRA	
Bifidobacterium adolescentis		61.16	0.50	IND 1	99	
Bacteroides uniformis		32.68	0.29	MNS 87	1	
Actinomyces israelii		3.37	0.35		99	
53755512	**Good identification**					**53755512**
					GRA	
Bifidobacterium adolescentis		98.90	0.52	IND 1	99	
Actinomyces israelii		0.67	0.21	MNS 87	99	
Bacteroides uniformis		0.44	-.05		1	
53755522	**Very good identification**					**53755522**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		99.39	0.63	MNS 99	5	
Bacteroides ovatus		0.49	0.27		68	
Bacteroides thetaiotaomicron		0.10	0.15		89	
53755552	**Good identification**					**53755552**
Bifidobacterium adolescentis		97.93	0.36	IND 1		
Actinomyces israelii		2.07	0.14	MNS 87		
53755622	**Good identification**					**53755622**
					CAT	
Bacteroides uniformis		95.92	0.37	MNS 99	5	
Bacteroides ovatus		3.20	0.15	LAC 98	68	
Bacteroides thetaiotaomicron		0.88	0.05		89	
53755722	**Good identification**					**53755722**
					CAT	
Bacteroides uniformis		92.86	0.66	MNS 99	5	
Bacteroides ovatus		6.27	0.49		68	
Bacteroides thetaiotaomicron		0.85	0.34		89	
53755732	**Low discrimination**					**53755732**
					CAT	
Bacteroides ovatus		76.32	0.46	MNS 99	68	
Bacteroides uniformis		17.14	0.32	TRE 91	5	
Bacteroides thetaiotaomicron		6.04	0.27		89	
53757000	**Good identification**					**53757000**
Actinomyces israelii		100.0	0.36	IND 1		
53757002	**Good identification**					**53757002**
Actinomyces israelii		100.0	0.39	IND 1		
53757100	**Good identification**					**53757100**
Actinomyces israelii		100.0	0.46	IND 1		
53757102	**Low discrimination**					**53757102**
Actinomyces israelii		75.72	0.49	IND 1		
Bifidobacterium adolescentis		24.28	0.35			
53757112	**Low discrimination**					**53757112**
Bifidobacterium adolescentis		72.40	0.36	IND 1		
Actinomyces israelii		27.61	0.35	MNS 87		
53757140	**Good identification**					**53757140**
Actinomyces israelii		100.0	0.39	IND 1		
53757142	**Acceptable identification**					**53757142**
Actinomyces israelii		90.75	0.42	IND 1		
Bifidobacterium adolescentis		9.25	0.19			
53757402	**Acceptable identification**					**53757402**
Actinomyces israelii		90.42	0.38	IND 1		
Bifidobacterium adolescentis		9.58	0.15			
53757500	**Acceptable identification**					**53757500**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		90.70	0.45	IND 1	1	
Clostridium butyricum		5.70	0.03		99	
Bifidobacterium adolescentis		3.60	0.15		1	
53757502	**Low discrimination**					**53757502**
					GRA	
Bifidobacterium adolescentis		71.73	0.49	IND 1	99	
Actinomyces israelii		27.35	0.48	MNS 87	99	
Bacteroides uniformis		0.47	-.05		1	
53757512	**Good identification**					**53757512**
Bifidobacterium adolescentis		95.07	0.50	IND 1		
Actinomyces israelii		4.43	0.34	MNS 87		
Bifidobacterium dentium		0.50	-.07			
53757540	**Good identification**					**53757540**
Actinomyces israelii		100.0	0.38	IND 1		
53757542	**Low discrimination**					**53757542**
Actinomyces israelii		54.54	0.41	IND 1		
Bifidobacterium adolescentis		45.46	0.33			
53757722	**Low discrimination**					**53757722**
					CAT	
Bacteroides ovatus		68.82	0.41	MNS 99	68	
Bacteroides uniformis		30.90	0.32	TRE 91	5	
Bacteroides thetaiotaomicron		0.28	0.00		89	
53757732	**Good identification**					**53757732**
Bacteroides ovatus		99.32	0.38	MNS 99		
Bacteroides uniformis		0.68	-.03	TRE 91		
53773100	**Low discrimination**					**53773100**
Actinomyces israelii		86.83	0.38	IND 1		
Actinomyces naeslundii		13.17	0.33	XYL 92		
53773101	**Good identification**					**53773101**
Actinomyces naeslundii		100.0	0.36	IND 1		
53773102	**Low discrimination**					**53773102**
Actinomyces israelii		70.54	0.41	IND 1		
Bifidobacterium breve		15.44	0.24	XYL 92		
Actinomyces naeslundii		7.14	0.33			
53773103	**Good identification**					**53773103**
Actinomyces naeslundii		100.0	0.36	IND 1		
53773121	**Good identification**					**53773121**
Actinomyces naeslundii		100.0	0.36	IND 1		
53773123	**Good identification**					**53773123**
Actinomyces naeslundii		100.0	0.36	IND 1		
53773500	**Acceptable identification**					**53773500**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		91.49	0.37	IND 1	1	
Clostridium butyricum		8.51	-.02	XYL 92	99	
53773502	**Low discrimination**					**53773502**
Actinomyces israelii		55.60	0.40	IND 1		
Bifidobacterium adolescentis		44.40	0.32	XYL 92		
53775002	**Good identification**					**53775002**
					SPO	
Actinomyces israelii		95.02	0.36	IND 1	1	
Clostridium butyricum		4.98	-.07		99	
53775102	**Low discrimination**					**53775102**
					SPO	
Actinomyces israelii		58.55	0.46	IND 1	1	
Bifidobacterium adolescentis		23.04	0.33		1	
Clostridium butyricum		18.41	0.17		99	
53775140	**Low discrimination**					**53775140**
					SPO	
Actinomyces israelii		78.67	0.36	IND 1	1	
Clostridium butyricum		21.33	0.06		99	
53775142	**Low discrimination**					**53775142**
					SPO	
Actinomyces israelii		83.66	0.39	IND 1	1	
Bifidobacterium adolescentis		10.46	0.18		1	
Clostridium butyricum		5.88	-.02		99	
53775322	**Low discrimination**					**53775322**
					CAT	
Bacteroides ovatus		87.61	0.40	MNS 99	68	
Bacteroides thetaiotaomicron		5.10	0.19	ARA 97	89	
Bacteroides uniformis		3.65	0.14		5	
53775332	**Acceptable identification**					**53775332**
Bacteroides ovatus		92.99	0.37	MNS 99		
Bacteroides thetaiotaomicron		3.16	0.12	ARA 97		
Parabacteroides distasonis		3.12	0.28			
53775500	**Acceptable identification**					**53775500**
					SPO	
Clostridium butyricum		92.71	0.40	IND 3	99	
Actinomyces israelii		6.95	0.42	MNS 99	1	
Bifidobacterium adolescentis		0.34	0.13		1	
53775512	**Low discrimination**					**53775512**
					SPO	
Bifidobacterium adolescentis		88.22	0.49	IND 1	1	
Clostridium butyricum		4.08	0.11	MNS 87	99	
Actinomyces israelii		3.35	0.31		1	
53775522	**Low discrimination**					**53775522**
					CAT	
Bacteroides ovatus		66.84	0.44	MNS 99	68	
Bacteroides thetaiotaomicron		17.03	0.34	RHA 95	89	
Bacteroides uniformis		13.61	0.29		5	
53775532	**Low discrimination**					**53775532**
Bacteroides ovatus		84.93	0.41	MNS 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		12.62	0.27	RHA 95		
Clostridium clostridioforme		0.84	0.17			
53775722	**Low discrimination**					**53775722**
					CAT	
Bacteroides ovatus		83.73	0.66	MNS 99	68	
Bacteroides thetaiotaomicron		14.92	0.53		89	
Bacteroides uniformis		1.24	0.32		5	
53775732	**Good identification**					**53775732**
Bacteroides ovatus		90.21	0.63	MNS 99		
Bacteroides thetaiotaomicron		9.37	0.46			
Clostridium clostridioforme		0.34	0.32			
53775762	**Good identification**					**53775762**
					CAT	
Bacteroides ovatus		95.53	0.42	MNS 99	68	
Bacteroides thetaiotaomicron		4.13	0.19	SOR 4	89	
Bacteroides uniformis		0.34	-.03		5	
53775772	**Good identification**					**53775772**
Bacteroides ovatus		97.54	0.39	MNS 99		
Bacteroides thetaiotaomicron		2.46	0.12	SOR 4		
53777000	**Good identification**					**53777000**
Actinomyces israelii		100.0	0.46	IND 1		
53777002	**Good identification**					**53777002**
Actinomyces israelii		100.0	0.49	IND 1		
53777040	**Good identification**					**53777040**
Actinomyces israelii		100.0	0.39	IND 1		
53777042	**Good identification**					**53777042**
Actinomyces israelii		100.0	0.42	IND 1		
53777100	**Very good identification**					**53777100**
					SPO	
Actinomyces israelii		97.68	0.56	IND 1	1	
Clostridium butyricum		2.32	0.07		99	
53777102	**Good identification**					**53777102**
					SPO	
Actinomyces israelii		94.08	0.59	IND 1	1	
Bifidobacterium adolescentis		5.35	0.32		1	
Clostridium butyricum		0.58	0.00		99	
53777110	**Good identification**					**53777110**
Actinomyces israelii		100.0	0.42	IND 1		
53777112	**Low discrimination**					**53777112**
Actinomyces israelii		67.14	0.45	IND 1		
Bifidobacterium adolescentis		31.20	0.33	MLZ 13		
Bifidobacterium dentium		1.66	-.07			
53777120	**Good identification**					**53777120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.36	IND 1		
53777122	**Low discrimination**					**53777122**
					GRA	
Actinomyces israelii		77.20	0.39	IND 1	99	
Bacteroides ovatus		22.81	0.10	NAG 6	1	
53777140	**Good identification**					**53777140**
Actinomyces israelii		100.0	0.49	IND 1		
53777142	**Good identification**					**53777142**
Actinomyces israelii		98.23	0.52	IND 1		
Bifidobacterium adolescentis		1.77	0.16			
53777150	**Acceptable identification**					**53777150**
					COC	
Actinomyces israelii		91.41	0.35	IND 1	1	
Blautia producta		8.59	-.03	MLZ 13	99	
53777152	**Low discrimination**					**53777152**
Actinomyces israelii		87.13	0.38	IND 1		
Bifidobacterium adolescentis		12.87	0.18	MLZ 13		
53777300	**Good identification**					**53777300**
Actinomyces israelii		100.0	0.37	IND 1		
53777302	**Good identification**					**53777302**
					GRA	
Actinomyces israelii		95.41	0.40	IND 1	99	
Bacteroides ovatus		4.59	-.02	RHA 7	1	
53777400	**Good identification**					**53777400**
					SPO	
Actinomyces israelii		95.82	0.44	IND 1	1	
Clostridium butyricum		4.18	0.00		99	
53777402	**Good identification**					**53777402**
					SPO	
Actinomyces israelii		97.08	0.47	IND 1	1	
Bifidobacterium adolescentis		1.82	0.12		1	
Clostridium butyricum		1.10	-.07		99	
53777440	**Good identification**					**53777440**
Actinomyces israelii		100.0	0.37	IND 1		
53777442	**Good identification**					**53777442**
Actinomyces israelii		100.0	0.40	IND 1		
53777500	**Low discrimination**					**53777500**
					SPO	
Actinomyces israelii		78.83	0.55	IND 1	1	
Clostridium butyricum		20.62	0.24		99	
Bifidobacterium adolescentis		0.56	0.12		1	
53777502	**Low discrimination**					**53777502**
					SPO	
Actinomyces israelii		64.74	0.58	IND 1	1	
Bifidobacterium adolescentis		30.09	0.46		1	
Clostridium butyricum		4.39	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
53777510	**Low discrimination**							**53777510**
					SPO			
Actinomyces israelii		61.05	0.41	IND 1	1			
Bifidobacterium dentium		27.08	0.10	MLZ 13	1			
Clostridium butyricum		5.62	0.02		99			
53777512	**Low discrimination**							**53777512**
Bifidobacterium adolescentis		48.43	0.47	IND 1				
Bifidobacterium dentium		38.09	0.27	MNS 87				
Actinomyces israelii		12.74	0.44					
53777522	**Low discrimination**							**53777522**
					GRA			
Bacteroides ovatus		88.93	0.36	MNS 99	1			
Actinomyces israelii		7.62	0.37	RHA 95	99			
Bifidobacterium dentium		0.96	-.03		99			
53777540	**Low discrimination**							**53777540**
					COC	SPO		
Actinomyces israelii		61.55	0.48	IND 1	1	1		
Blautia producta		34.85	0.24		99	1		
Clostridium butyricum		3.60	0.06		1	99		
53777542	**Low discrimination**							**53777542**
					COC			
Actinomyces israelii		83.00	0.51	IND 1	1			
Bifidobacterium adolescentis		12.26	0.30		1			
Blautia producta		3.48	0.07		99			
53777700	**Acceptable identification**							**53777700**
					SPO			
Actinomyces israelii		90.30	0.35	IND 1	1			
Clostridium butyricum		9.70	-.02	RHA 7	99			
53777722	**Very good identification**							**53777722**
					CAT			
Bacteroides ovatus		99.39	0.58	MNS 99	68			
Bacteroides thetaiotaomicron		0.54	0.19		89			
Bacteroides uniformis		0.05	-.03		5			
53777732	**Good identification**							**53777732**
					GRA	CAT		
Bacteroides ovatus		99.60	0.55	MNS 99	1	68		
Bacteroides thetaiotaomicron		0.31	0.12		1	89		
Bifidobacterium dentium		0.09	-.03		99	1		
54000000	**Good identification**							**54000000**
					COC	GRA	SPO	
Clostridium bifermentans		92.90	0.76	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.32	0.32		1	1	1	
Peptoniphilus asaccharolyticus		1.32	0.32		99	99	1	
54000001	**Low discrimination**							**54000001**
					COC	SPO		
Anaerococcus prevotii		82.75	0.49	IND 9	99	1		
Clostridium bifermentans		15.16	0.41	ESL 1	1	99		
Clostridium sordellii		0.94	0.16		1	99		
54000002	**Low discrimination**							**54000002**
					COC	SPO		
Clostridium bifermentans		58.53	0.41	GLU 92	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Atopobium minutum		24.51	0.43	BGL 1	1	1		
Anaerococcus prevotii		8.87	0.22		99	1		
54000003	**Good identification**							**54000003**
					COC	SPO		
Anaerococcus prevotii		98.96	0.38	IND 9	99	1		
Clostridium bifermentans		0.73	0.07	ESL 1	1	99		
Atopobium minutum		0.31	0.09		1	1		
54000010	**Acceptable identification**							**54000010**
					COC	GRA	SPO	
Clostridium bifermentans		93.49	0.41	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.33	-.03	MLZ 1	1	1	1	
Peptoniphilus asaccharolyticus		1.33	-.03		99	99	1	
54000020	**Low discrimination**							**54000020**
					SPO			
Clostridium bifermentans		80.42	0.55	GLU 92	99			
Clostridium cadaveris		15.99	0.41	NAG 6	99			
Atopobium minutum		2.67	0.38		1			
54000040	**Good identification**							**54000040**
					GRA	SPO		
Clostridium bifermentans		96.91	0.52	GLU 92	99	99		
Clostridium botulinum biovar A		1.46	0.21	SOR 4	99	99		
Porphyromonas asaccharolytica		0.33	-.03		1	1		
54000100	**Acceptable identification**							**54000100**
					COC	GRA	SPO	
Clostridium bifermentans		93.49	0.41	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.33	-.03	LAC 1	1	1	1	
Peptoniphilus asaccharolyticus		1.33	-.03		99	99	1	
54000200	**Acceptable identification**							**54000200**
					COC	GRA	SPO	
Clostridium bifermentans		93.49	0.41	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.33	-.03	RHA 1	1	1	1	
Peptoniphilus asaccharolyticus		1.33	-.03		99	99	1	
54000400	**Acceptable identification**							**54000400**
					COC	GRA	SPO	
Clostridium bifermentans		93.49	0.41	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.33	-.03	ARA 1	1	1	1	
Peptoniphilus asaccharolyticus		1.33	-.03		99	99	1	
54001000	**Low discrimination**							**54001000**
					GRA	SPO		
Clostridium bifermentans		52.31	0.41	GLU 92	99	99		
Fusobacterium necrophorum subsp. necrophorum		33.77	0.32	GAL 1	1	1		
Atopobium minutum		10.95	0.38		99	1		
54001002	**Low discrimination**							**54001002**
					GRA	SPO		
Atopobium minutum		89.36	0.35	IND 1	99	1		
Clostridium bifermentans		6.47	0.07		99	99		
Fusobacterium necrophorum subsp. necrophorum		4.18	-.03		1	1		
54001520	**Very good identification**							**54001520**
Odoribacter splanchnicus		100.0	0.61	GLU 99				
54001560	**Good identification**							**54001560**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Odoribacter splanchnicus		100.0	0.45	GLU 99				
54002000	**Acceptable identification**							**54002000**
					COC	GRA	SPO	
Clostridium bifermentans		93.09	0.41	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.33	-.03	MAN 1	1	1	1	
Peptoniphilus asaccharolyticus		1.33	-.03		99	99	1	
54004000	**Acceptable identification**							**54004000**
					COC	GRA	SPO	
Clostridium bifermentans		92.86	0.41	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.32	-.03	XYL 1	1	1	1	
Peptoniphilus asaccharolyticus		1.32	-.03		99	99	1	
54010000	**Low discrimination**							**54010000**
					GRA	SPO		
Fusobacterium varium		74.20	0.57	ESL 1	1	1		
Clostridium bifermentans		22.48	0.54		99	99		
Atopobium minutum		2.50	0.46		99	1		
54010002	**Low discrimination**							**54010002**
					GRA	SPO		
Atopobium minutum		63.07	0.43	IND 1	99	1		
Fusobacterium varium		28.35	0.23		1	1		
Clostridium bifermentans		8.59	0.20		99	99		
54010020	**Low discrimination**							**54010020**
					SPO			
Clostridium cadaveris		63.57	0.41	ESL 1	99			
Clostridium bifermentans		16.83	0.33	GLU 95	99			
Atopobium minutum		9.78	0.37		1			
54011000	**Low discrimination**							**54011000**
					GRA	SPO		
Atopobium minutum		43.88	0.37	IND 1	99	1		
Fusobacterium varium		39.45	0.23		1	1		
Clostridium bifermentans		11.95	0.20		99	99		
54011520	**Very good identification**							**54011520**
Odoribacter splanchnicus		100.0	0.66	GLU 99				
54011560	**Good identification**							**54011560**
Odoribacter splanchnicus		100.0	0.50	GLU 99				
54020000	**Low discrimination**							**54020000**
					GRA	SPO		
Clostridium bifermentans		88.30	0.41	GLU 92	99	99		
Clostridium botulinum biovar A		4.59	0.20	TRE 1	99	99		
Fusobacterium nucleatum subsp. nucleatum		1.26	-.03		1	1		
54040000	**Acceptable identification**							**54040000**
					COC	GRA	SPO	
Clostridium bifermentans		93.49	0.41	GLU 92	1	99	99	
Fusobacterium nucleatum subsp. nucleatum		1.33	-.03	CEL 1	1	1	1	
Peptoniphilus asaccharolyticus		1.33	-.03		99	99	1	
54100000	**Very good identification**							**54100000**
Clostridium bifermentans		99.86	0.82	GLU 92				
Clostridium botulinum biovar A		0.07	0.29					
Clostridium sporogenes		0.03	0.19					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
54100001	**Low discrimination**						**54100001**
					COC	SPO	
Clostridium bifermentans		79.26	0.48	GLU 92	1	99	
Clostridium sordellii		18.86	0.33	URE 1	1	99	
Anaerococcus prevotii		1.87	0.15		99	1	
54100002	**Good identification**						**54100002**
					SPO		
Clostridium bifermentans		98.97	0.48	GLU 92	99		
Clostridium sporogenes		0.86	0.11	BGL 1	99		
Atopobium minutum		0.18	0.09		1		
54100010	**Good identification**						**54100010**
Clostridium bifermentans		100.0	0.48	GLU 92			
54100020	**Very good identification**						**54100020**
Clostridium bifermentans		99.55	0.61	GLU 92			
Clostridium cadaveris		0.45	0.19	NAG 6			
54100040	**Very good identification**						**54100040**
Clostridium bifermentans		99.48	0.58	GLU 92			
Clostridium botulinum biovar A		0.53	0.20	SOR 4			
54100060	**Good identification**						**54100060**
Clostridium bifermentans		100.0	0.38	GLU 92			
54100100	**Good identification**						**54100100**
Clostridium bifermentans		100.0	0.48	GLU 92			
54100200	**Good identification**						**54100200**
Clostridium bifermentans		100.0	0.48	GLU 92			
54100400	**Good identification**						**54100400**
Clostridium bifermentans		100.0	0.48	GLU 92			
54101000	**Good identification**						**54101000**
					GRA	SPO	
Clostridium bifermentans		99.12	0.48	GLU 92	99	99	
Prevotella intermedia		0.61	0.12	GAL 1	1	1	
Fusobacterium necrophorum subsp. necrophorum		0.28	-.03		1	1	
54101722	**Good identification**						**54101722**
Bacteroides eggerthii		100.0	0.36	GLU 99			
54102000	**Good identification**						**54102000**
Clostridium bifermentans		100.0	0.48	GLU 92			
54104000	**Good identification**						**54104000**
Clostridium bifermentans		100.0	0.48	GLU 92			
54105722	**Good identification**						**54105722**
Bacteroides eggerthii		100.0	0.56	GLU 99			
54110000	**Low discrimination**						**54110000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium bifermentans		88.21	0.60	GLU 92	99	99	
Prevotella intermedia		10.25	0.47	FRU 5	1	1	
Fusobacterium varium		1.26	0.23		1	1	
54110020	**Acceptable identification**						**54110020**
					GRA	SPO	
Clostridium bifermentans		90.61	0.40	GLU 92	99	99	
Clostridium cadaveris		7.72	0.19	FRU 5	99	99	
Prevotella intermedia		1.67	0.12	NAG 6	1	1	
54110040	**Good identification**						**54110040**
					GRA	SPO	
Clostridium bifermentans		96.66	0.36	GLU 92	99	99	
Prevotella intermedia		2.72	0.12	FRU 5	1	1	
Clostridium botulinum biovar A		0.62	-.01	SOR 4	99	99	
54111000	**Acceptable identification**						**54111000**
					SPO	GRA	
Prevotella intermedia		91.66	0.47	ESL 1	1	1	
Clostridium bifermentans		7.97	0.26	GLU 90	99	99	
Eubacterium saburreum		0.37	0.15		1	85	
54120000	**Good identification**						**54120000**
Clostridium bifermentans		97.38	0.48	GLU 92			
Clostridium botulinum biovar A		1.78	0.19	TRE 1			
Clostridium sporogenes		0.84	0.11				
54140000	**Good identification**						**54140000**
Clostridium bifermentans		100.0	0.48	GLU 92			
54141722	**Good identification**						**54141722**
Bacteroides eggerthii		100.0	0.47	GLU 99			
54145722	**Very good identification**						**54145722**
Bacteroides eggerthii		100.0	0.66	GLU 99			
54200000	**Very good identification**						**54200000**
					GRA	SPO	
Clostridium bifermentans		99.37	0.59	GLU 92	99	99	
Fusobacterium nucleatum subsp. nucleatum		0.13	-.03	SAL 10	1	1	
Porphyromonas asaccharolytica		0.13	-.03		1	1	
54200020	**Good identification**						**54200020**
Clostridium bifermentans		98.22	0.39	GLU 92			
Clostridium cadaveris		1.78	0.07	SAL 10			
54200040	**Good identification**						**54200040**
Clostridium bifermentans		100.0	0.36	GLU 92			
54210000	**Low discrimination**						**54210000**
					GRA	SPO	
Clostridium bifermentans		76.32	0.37	GLU 92	99	99	
Fusobacterium varium		22.91	0.23	SAL 10	1	1	
Atopobium minutum		0.77	0.11	FRU 5	99	1	
54300000	**Very good identification**						**54300000**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium bifermentans		100.0	0.66	GLU 92				
54300020	**Good identification**							**54300020**
Clostridium bifermentans		100.0	0.45	GLU 92				
54300040	**Good identification**							**54300040**
Clostridium bifermentans		100.0	0.42	GLU 92				
54310000	**Good identification**							**54310000**
					GRA	SPO		
Clostridium bifermentans		98.96	0.44	GLU 92	99	99		
Prevotella intermedia		1.05	0.12	SAL 10	1	1		
54400000	**Low discrimination**							**54400000**
					COC	GRA	SPO	
Clostridium bifermentans		79.01	0.41	GLU 92	1	99	99	
Anaerococcus prevotii		15.97	0.24	RAF 1	99	99	1	
Fusobacterium nucleatum subsp. nucleatum		1.13	-.03		1	1	1	
54400001	**Good identification**							**54400001**
					COC	SPO		
Anaerococcus prevotii		99.45	0.41	IND 9	99	1		
Clostridium bifermentans		0.55	0.07	ESL 1	1	99		
54500000	**Good identification**							**54500000**
					GRA	SPO		
Clostridium bifermentans		99.09	0.48	GLU 92	99	99		
Prevotella intermedia		0.91	0.15	RAF 1	1	1		
54510000	**Acceptable identification**							**54510000**
					GRA	SPO		
Prevotella intermedia		94.52	0.50	ESL 1	1	1		
Clostridium bifermentans		5.48	0.26	GLU 90	99	99		
54511000	**Good identification**							**54511000**
Prevotella intermedia		99.85	0.50	ESL 1				
Eubacterium saburreum		0.15	0.11	GLU 90				
54535722	**Good identification**							**54535722**
Bacteroides thetaiotaomicron		99.44	0.40	GLU 97				
Parabacteroides distasonis		0.56	0.17	SUC 98				
54575722	**Good identification**							**54575722**
Bacteroides thetaiotaomicron		97.73	0.45	GLU 97				
Bacteroides ovatus		2.13	0.17	SUC 98				
Parabacteroides distasonis		0.14	0.13					
54575732	**Good identification**							**54575732**
Bacteroides thetaiotaomicron		96.39	0.38	GLU 97				
Bacteroides ovatus		3.61	0.14	SUC 98				
55000000	**Excellent identification**							**55000000**
					SPO			
Clostridium bifermentans		99.63	0.94		99			
Clostridium botulinum biovar A		0.16	0.47		99			
Atopobium minutum		0.09	0.50		1			
55000001	**Good identification**							**55000001**
					COC	SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		90.04	0.60	URE 1	1	99	
Clostridium sordellii		9.18	0.38		1	99	
Anaerococcus prevotii		0.43	0.15		99	1	
55000002	**Good identification**						**55000002**
					SPO		
Clostridium bifermentans		93.50	0.60	BGL 1	99		
Atopobium minutum		5.80	0.47		1		
Clostridium sporogenes		0.50	0.19		99		
55000010	**Very good identification**						**55000010**
Clostridium bifermentans		99.27	0.60	MLZ 1			
Clostridium difficile		0.35	0.15				
Clostridium botulinum biovar A		0.16	0.13				
55000020	**Low discrimination**						**55000020**
					SPO		
Clostridium bifermentans		74.96	0.73	NAG 6	99		
Clostridium cadaveris		24.62	0.63		99		
Atopobium minutum		0.37	0.42		1		
55000021	**Low discrimination**						**55000021**
Clostridium bifermentans		74.10	0.39	NAG 6			
Clostridium cadaveris		24.34	0.29	URE 1			
Clostridium sordellii		1.20	0.04				
55000022	**Low discrimination**						**55000022**
					SPO		
Clostridium bifermentans		60.22	0.39	NAG 6	99		
Clostridium cadaveris		19.78	0.29	BGL 1	99		
Atopobium minutum		19.50	0.39		1		
55000030	**Low discrimination**						**55000030**
					SPO		
Clostridium bifermentans		75.00	0.39	MLZ 1	99		
Clostridium cadaveris		24.64	0.29	NAG 6	99		
Atopobium minutum		0.37	0.08		1		
55000040	**Very good identification**						**55000040**
					SPO		
Clostridium bifermentans		98.79	0.70	SOR 4	99		
Clostridium botulinum biovar A		1.16	0.38		99		
Atopobium minutum		0.02	0.16		1		
55000041	**Low discrimination**						**55000041**
Clostridium bifermentans		88.78	0.36	SOR 4			
Clostridium sordellii		9.05	0.15	URE 1			
Clostridium novyi biovar A		1.12	0.16				
55000042	**Good identification**						**55000042**
					SPO		
Clostridium bifermentans		97.39	0.36	SOR 4	99		
Atopobium minutum		1.46	0.13	BGL 1	1		
Clostridium botulinum biovar A		1.15	0.04		99		
55000050	**Good identification**						**55000050**
Clostridium bifermentans		97.62	0.36	MLZ 1			
Clostridium novyi biovar A		1.23	0.16	SOR 4			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		1.15	0.04				
55000060	**Acceptable identification**						**55000060**
Clostridium bifermentans		92.24	0.50	NAG 6			
Clostridium cadaveris		7.35	0.29	SOR 4			
Clostridium botulinum biovar A		0.17	0.04				
55000100	**Very good identification**						**55000100**
					SPO		
Clostridium bifermentans		99.66	0.60	LAC 1	99		
Clostridium botulinum biovar A		0.16	0.13		99		
Atopobium minutum		0.09	0.16		1		
55000120	**Low discrimination**						**55000120**
					GRA	SPO	
Clostridium bifermentans		53.83	0.39	LAC 1	99	99	
Odoribacter splanchnicus		27.39	0.27	NAG 6	1	1	
Clostridium cadaveris		17.68	0.29		99	99	
55000140	**Good identification**						**55000140**
Clostridium bifermentans		98.84	0.36	LAC 1			
Clostridium botulinum biovar A		1.17	0.04	SOR 4			
55000200	**Very good identification**						**55000200**
					SPO		
Clostridium bifermentans		99.69	0.60	RHA 1	99		
Clostridium botulinum biovar A		0.16	0.13		99		
Atopobium minutum		0.09	0.16		1		
55000220	**Low discrimination**						**55000220**
					SPO		
Clostridium bifermentans		75.00	0.39	RHA 1	99		
Clostridium cadaveris		24.64	0.29	NAG 6	99		
Atopobium minutum		0.37	0.08		1		
55000240	**Good identification**						**55000240**
Clostridium bifermentans		98.84	0.36	RHA 1			
Clostridium botulinum biovar A		1.17	0.04	SOR 4			
55000400	**Very good identification**						**55000400**
					SPO		
Clostridium bifermentans		99.64	0.60	ARA 1	99		
Clostridium botulinum biovar A		0.16	0.13		99		
Atopobium minutum		0.09	0.16		1		
55000420	**Low discrimination**						**55000420**
					GRA	SPO	
Clostridium bifermentans		54.28	0.39	ARA 1	99	99	
Odoribacter splanchnicus		27.62	0.27	NAG 6	1	1	
Clostridium cadaveris		17.83	0.29		99	99	
55000440	**Good identification**						**55000440**
Clostridium bifermentans		98.84	0.36	ARA 1			
Clostridium botulinum biovar A		1.17	0.04	SOR 4			
55000520	**Very good identification**						**55000520**
					GRA	SPO	
Odoribacter splanchnicus		99.98	0.61	GAL 99	1	1	
Clostridium bifermentans		0.02	0.05		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55000560	**Good identification**						**55000560**
Odoribacter splanchnicus		100.0	0.45	GAL 99			
55001000	**Very good identification**						**55001000**
					GRA	SPO	
Clostridium bifermentans		95.56	0.60	GAL 1	99	99	
Atopobium minutum		2.96	0.42		99	1	
Fusobacterium necrophorum subsp. necrophorum		0.73	0.17		1	1	
55001002	**Low discrimination**						**55001002**
					SPO		
Atopobium minutum		65.62	0.39	IND 1	1		
Clostridium bifermentans		32.07	0.25		99		
Eubacterium saburreum		2.32	0.18		1		
55001020	**Low discrimination**						**55001020**
					GRA	SPO	
Clostridium bifermentans		49.06	0.39	GAL 1	99	99	
Odoribacter splanchnicus		24.96	0.27	NAG 6	1	1	
Clostridium cadaveris		16.12	0.29		99	99	
55001040	**Good identification**						**55001040**
					SPO		
Clostridium bifermentans		98.11	0.36	GAL 1	99		
Clostridium botulinum biovar A		1.16	0.04	SOR 4	99		
Atopobium minutum		0.74	0.08		1		
55001110	**Good identification**						**55001110**
Eubacterium saburreum		96.01	0.36	SUC 99			
Odoribacter splanchnicus		3.99	-.07				
55001120	**Very good identification**						**55001120**
					SPO	GRA	
Odoribacter splanchnicus		99.93	0.61	ARA 99	1	1	
Eubacterium saburreum		0.05	0.24		1	85	
Clostridium bifermentans		0.02	0.05		99	99	
55001160	**Good identification**						**55001160**
Odoribacter splanchnicus		100.0	0.45	ARA 99			
55001420	**Very good identification**						**55001420**
					GRA	SPO	
Odoribacter splanchnicus		99.98	0.61	LAC 99	1	1	
Clostridium bifermentans		0.02	0.05		99	99	
55001460	**Good identification**						**55001460**
Odoribacter splanchnicus		100.0	0.45	LAC 99			
55001500	**Very good identification**						**55001500**
Odoribacter splanchnicus		100.0	0.61	NAG 99			
55001520	**Excellent identification**						**55001520**
Odoribacter splanchnicus		100.0	0.95				
55001521	**Very good identification**						**55001521**
Odoribacter splanchnicus		100.0	0.61	URE 1			
55001522	**Very good identification**						**55001522**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Odoribacter splanchnicus		99.95	0.61	BGL 1	1		
Bacteroides eggerthii		0.05	0.02		93		
55001530	**Very good identification**						**55001530**
Odoribacter splanchnicus		100.0	0.61	MLZ 1			
55001540	**Good identification**						**55001540**
Odoribacter splanchnicus		100.0	0.45	NAG 99			
55001560	**Very good identification**						**55001560**
Odoribacter splanchnicus		100.0	0.79	SOR 10			
55001561	**Good identification**						**55001561**
Odoribacter splanchnicus		100.0	0.45	SOR 10			
55001562	**Good identification**						**55001562**
Odoribacter splanchnicus		100.0	0.45	SOR 10			
55001570	**Good identification**						**55001570**
Odoribacter splanchnicus		100.0	0.45	MLZ 1			
55001720	**Very good identification**						**55001720**
					CAT		
Odoribacter splanchnicus		99.95	0.61	RHA 1	1		
Bacteroides eggerthii		0.05	0.02		93		
55001722	**Low discrimination**						**55001722**
					CAT		
Bacteroides eggerthii		83.19	0.36	MLT 99	93		
Odoribacter splanchnicus		16.81	0.27	XYL 93	1		
55001760	**Good identification**						**55001760**
Odoribacter splanchnicus		100.0	0.45	RHA 1			
55002000	**Very good identification**						**55002000**
					SPO		
Clostridium bifermentans		95.29	0.60	MAN 1	99		
Eubacterium limosum		3.80	0.32		1		
Clostridium difficile		0.61	0.20		99		
55002020	**Low discrimination**						**55002020**
					SPO	CAT	
Clostridium bifermentans		73.78	0.39	MAN 1	99	1	
Clostridium cadaveris		24.24	0.29	NAG 6	99	1	
Propionibacterium acnes		1.15	0.09		1	95	
55002040	**Good identification**						**55002040**
					SPO		
Clostridium bifermentans		97.13	0.36	MAN 1	99		
Clostridium botulinum biovar A		1.14	0.04	SOR 4	99		
Eubacterium limosum		0.94	-.03		1		
55003520	**Very good identification**						**55003520**
Odoribacter splanchnicus		100.0	0.61	MAN 1			
55003560	**Good identification**						**55003560**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Odoribacter splanchnicus		100.0	0.45	MAN 1			
55004000	**Very good identification**						**55004000**
					GRA	SPO	
Clostridium bifermentans		99.51	0.60	XYL 1	99	99	
Fusobacterium varium		0.21	0.08		1	1	
Clostridium botulinum biovar A		0.16	0.13		99	99	
55004020	**Low discrimination**						**55004020**
					SPO		
Clostridium bifermentans		75.00	0.39	XYL 1	99		
Clostridium cadaveris		24.64	0.29	NAG 6	99		
Atopobium minutum		0.37	0.08		1		
55004040	**Acceptable identification**						**55004040**
Clostridium bifermentans		93.12	0.36	XYL 1			
Clostridium glycolicum		5.79	0.15	SOR 4			
Clostridium botulinum biovar A		1.10	0.04				
55005520	**Very good identification**						**55005520**
Odoribacter splanchnicus		100.0	0.61	XYL 1			
55005560	**Good identification**						**55005560**
Odoribacter splanchnicus		100.0	0.45	XYL 1			
55005722	**Good identification**						**55005722**
					CAT		
Bacteroides eggerthii		99.99	0.56	MLT 99	93		
Odoribacter splanchnicus		0.02	-.07		1		
55010000	**Low discrimination**						**55010000**
					GRA	SPO	
Clostridium bifermentans		47.05	0.72	FRU 5	99	99	
Fusobacterium varium		45.21	0.66		1	1	
Eubacterium limosum		5.81	0.52		99	1	
55010001	**Low discrimination**						**55010001**
					GRA	SPO	
Clostridium sordellii		47.81	0.38	ESL 5	99	99	
Clostridium bifermentans		24.68	0.38	MNS 4	99	99	
Fusobacterium varium		23.72	0.32	MLT 90	1	1	
55010002	**Low discrimination**						**55010002**
					GRA	SPO	
Atopobium minutum		33.84	0.47	IND 1	99	1	
Clostridium bifermentans		31.11	0.38		99	99	
Fusobacterium varium		29.90	0.32		1	1	
55010010	**Low discrimination**						**55010010**
					GRA	SPO	
Clostridium difficile		75.80	0.50	IND 1	99	99	
Clostridium bifermentans		11.52	0.38	MAN 90	99	99	
Fusobacterium varium		11.07	0.32		1	1	
55010020	**Low discrimination**						**55010020**
					SPO	CAT	
Clostridium cadaveris		77.72	0.63	ESL 1	99	1	
Clostridium bifermentans		12.45	0.51		99	1	
Propionibacterium acnes		6.56	0.48		1	95	
55010040	**Low discrimination**						**55010040**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	SPO		
Clostridium bifermentans		76.58	0.48	FRU 5	99	99		
Fusobacterium varium		17.84	0.32	SOR 4	1	1		
Eubacterium limosum		2.29	0.18		99	1		
55010060	**Low discrimination**							**55010060**
					SPO	CAT		
Propionibacterium acnes		51.93	0.36	NIT 92	1	95		
Clostridium cadaveris		28.29	0.29	ESL 1	99	1		
Clostridium bifermentans		18.70	0.28		99	1		
55010100	**Low discrimination**							**55010100**
					GRA	SPO		
Clostridium bifermentans		42.53	0.38	FRU 5	99	99		
Fusobacterium varium		40.87	0.32	LAC 1	1	1		
Fusobacterium mortiferum		9.34	0.40		1	1		
55010200	**Low discrimination**							**55010200**
					GRA	SPO		
Clostridium bifermentans		47.29	0.38	FRU 5	99	99		
Fusobacterium varium		45.45	0.32	RHA 1	1	1		
Eubacterium limosum		5.84	0.18		99	1		
55010400	**Low discrimination**							**55010400**
					GRA	SPO		
Clostridium bifermentans		46.91	0.38	FRU 5	99	99		
Fusobacterium varium		45.08	0.32	ARA 1	1	1		
Eubacterium limosum		5.79	0.18		99	1		
55010520	**Very good identification**							**55010520**
Odoribacter splanchnicus		100.0	0.66	GAL 99				
55010560	**Good identification**							**55010560**
Odoribacter splanchnicus		100.0	0.50	GAL 99				
55011000	**Low discrimination**							**55011000**
					GRA	SPO		
Clostridium bifermentans		30.28	0.38	FRU 5	99	99		
Fusobacterium varium		29.10	0.32	GAL 1	1	1		
Atopobium minutum		16.46	0.41		99	1		
55011002	**Low discrimination**							**55011002**
					GRA			
Atopobium minutum		82.46	0.38	IND 1	99			
Eubacterium saburreum		9.47	0.26		85			
Fusobacterium mortiferum		3.57	0.20		1			
55011010	**Low discrimination**							**55011010**
					SPO			
Eubacterium saburreum		67.77	0.37	SUC 99	1			
Clostridium difficile		23.34	0.15		99			
Clostridium bifermentans		3.55	0.04		99			
55011020	**Low discrimination**							**55011020**
					GRA	SPO	CAT	
Propionibacterium acnes		64.41	0.48	NIT 92	99	1	95	
Odoribacter splanchnicus		22.16	0.32	ESL 1	1	1	1	
Clostridium cadaveris		7.70	0.29		99	99	1	
55011060	**Low discrimination**							**55011060**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		85.17	0.36	NIT 92	99	95	
Odoribacter splanchnicus		14.83	0.15	ESL 1	1	1	
55011110	**Good identification**						**55011110**
Eubacterium saburreum		97.49	0.44	SUC 99			
Odoribacter splanchnicus		2.51	-.03				
55011120	**Very good identification**						**55011120**
					SPO	GRA	
Odoribacter splanchnicus		99.78	0.66	ARA 99	1	1	
Clostridium septicum		0.11	0.18		99	99	
Eubacterium saburreum		0.09	0.32		1	85	
55011160	**Good identification**						**55011160**
					GRA	CAT	
Odoribacter splanchnicus		99.94	0.50	ARA 99	1	1	
Propionibacterium acnes		0.06	0.02	SOR 10	99	95	
55011420	**Very good identification**						**55011420**
					GRA	CAT	
Odoribacter splanchnicus		99.97	0.66	LAC 99	1	1	
Propionibacterium acnes		0.03	0.14		99	95	
55011460	**Good identification**						**55011460**
					GRA	CAT	
Odoribacter splanchnicus		99.94	0.50	LAC 99	1	1	
Propionibacterium acnes		0.06	0.02	SOR 10	99	95	
55011500	**Very good identification**						**55011500**
Odoribacter splanchnicus		100.0	0.66	NAG 99			
Eubacterium saburreum		0.01	0.11				
55011520	**Excellent identification**						**55011520**
Odoribacter splanchnicus		100.0	1.00				
55011521	**Very good identification**						**55011521**
Odoribacter splanchnicus		100.0	0.66	URE 1			
55011522	**Very good identification**						**55011522**
Odoribacter splanchnicus		100.0	0.66	BGL 1			
55011530	**Very good identification**						**55011530**
Odoribacter splanchnicus		100.0	0.66	MLZ 1			
55011540	**Good identification**						**55011540**
Odoribacter splanchnicus		100.0	0.50	NAG 99			
55011560	**Very good identification**						**55011560**
Odoribacter splanchnicus		100.0	0.84	SOR 10			
55011561	**Good identification**						**55011561**
Odoribacter splanchnicus		100.0	0.50	SOR 10			
55011562	**Good identification**						**55011562**
Odoribacter splanchnicus		100.0	0.50	SOR 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
55011570	**Good identification**						**55011570**
Odoribacter splanchnicus		100.0	0.50	MLZ 1			
55011720	**Very good identification**						**55011720**
Odoribacter splanchnicus		100.0	0.66	RHA 1			
55011760	**Good identification**						**55011760**
Odoribacter splanchnicus		100.0	0.50	RHA 1			
55012000	**Low discrimination**						**55012000**
					SPO		
Eubacterium limosum		84.16	0.66	IND 1	1		
Clostridium difficile		13.60	0.54		99		
Clostridium bifermentans		1.12	0.38		99		
55012010	**Very good identification**						**55012010**
					SPO		
Clostridium difficile		98.70	0.66	IND 1	99		
Eubacterium limosum		1.26	0.32		1		
Clostridium bifermentans		0.02	0.04		99		
55012020	**Low discrimination**						**55012020**
					SPO	CAT	
Propionibacterium acnes		57.98	0.43	NIT 92	1	95	
Eubacterium limosum		23.46	0.32	ESL 1	1	1	
Clostridium cadaveris		12.33	0.29		99	1	
55012050	**Good identification**						**55012050**
					SPO		
Clostridium difficile		99.76	0.44	IND 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		
55013020	**Good identification**						**55013020**
					GRA	CAT	
Propionibacterium acnes		98.98	0.43	NIT 92	99	95	
Odoribacter splanchnicus		0.61	-.03	ESL 1	1	1	
Eubacterium limosum		0.41	-.03		99	1	
55013520	**Very good identification**						**55013520**
Odoribacter splanchnicus		100.0	0.66	MAN 1			
55013560	**Good identification**						**55013560**
Odoribacter splanchnicus		100.0	0.50	MAN 1			
55014000	**Low discrimination**						**55014000**
					GRA	SPO	
Fusobacterium varium		76.08	0.42	ESL 1	1	1	
Clostridium bifermentans		19.19	0.38	XYL 4	99	99	
Eubacterium limosum		2.37	0.18		99	1	
55015100	**Low discrimination**						**55015100**
Eubacterium saburreum		89.07	0.41	SUC 99			
Fusobacterium mortiferum		7.45	0.24				
Odoribacter splanchnicus		3.48	-.03				
55015110	**Good identification**						**55015110**
Eubacterium saburreum		100.0	0.39	SUC 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
55015520	**Very good identification**						**55015520**
Odoribacter splanchnicus		100.0	0.66	XYL 1			
55015560	**Good identification**						**55015560**
Odoribacter splanchnicus		100.0	0.50	XYL 1			
55016010	**Good identification**						**55016010**
					SPO		
Clostridium difficile		99.76	0.44	IND 1	99		
Eubacterium limosum		0.25	-.03	XYL 5	1		
55020000	**Very good identification**						**55020000**
Clostridium bifermentans		95.37	0.60	TRE 1			
Clostridium botulinum biovar A		3.88	0.36				
Clostridium sporogenes		0.51	0.19				
55020020	**Low discrimination**						**55020020**
Clostridium cadaveris		62.86	0.41	ESL 1			
Clostridium bifermentans		36.72	0.39	TRE 5			
Clostridium botulinum biovar A		0.24	0.02				
55020040	**Low discrimination**						**55020040**
Clostridium bifermentans		77.42	0.36	TRE 1			
Clostridium botulinum biovar A		22.58	0.27	SOR 4			
55021520	**Very good identification**						**55021520**
Odoribacter splanchnicus		100.0	0.61	TRE 1			
55021560	**Good identification**						**55021560**
Odoribacter splanchnicus		100.0	0.45	TRE 1			
55030000	**Low discrimination**						**55030000**
					GRA	SPO	
Clostridium bifermentans		43.01	0.38	FRU 5	99	99	
Fusobacterium varium		41.33	0.32	TRE 1	1	1	
Eubacterium limosum		5.31	0.18		99	1	
55030020	**Acceptable identification**						**55030020**
					SPO	CAT	
Clostridium cadaveris		94.85	0.41	ESL 1	99	1	
Clostridium bifermentans		2.92	0.17	TRE 5	99	1	
Propionibacterium acnes		1.54	0.14		1	95	
55031000	**Low discrimination**						**55031000**
					SPO	GRA	
Eubacterium saburreum		68.00	0.36	SUC 99	1	85	
Fusobacterium mortiferum		18.23	0.28		1	1	
Clostridium bifermentans		4.37	0.04		99	99	
55031100	**Low discrimination**						**55031100**
Eubacterium saburreum		76.94	0.42	SUC 99			
Fusobacterium mortiferum		20.63	0.34				
Odoribacter splanchnicus		2.43	-.03				
55031110	**Good identification**						**55031110**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.41	SUC 99			
55031520	**Very good identification**						**55031520**
Odoribacter splanchnicus		100.0	0.66	TRE 1			
55031560	**Good identification**						**55031560**
Odoribacter splanchnicus		100.0	0.50	TRE 1			
55032010	**Good identification**						**55032010**
					SPO		
Clostridium difficile		99.83	0.47	IND 1	99		
Eubacterium limosum		0.17	-.03	TRE 7	1		
55035100	**Acceptable identification**						**55035100**
					COC	GRA	
Eubacterium saburreum		94.34	0.38	SUC 99	1	85	
Fusobacterium mortiferum		3.54	0.15		1	1	
Atopobium parvulum		2.13	0.00		99	99	
55035110	**Good identification**						**55035110**
Eubacterium saburreum		100.0	0.36	SUC 99			
55040000	**Very good identification**						**55040000**
					SPO		
Clostridium bifermentans		99.69	0.60	CEL 1	99		
Clostridium botulinum biovar A		0.16	0.13		99		
Atopobium minutum		0.09	0.16		1		
55040020	**Low discrimination**						**55040020**
					SPO		
Clostridium bifermentans		75.00	0.39	CEL 1	99		
Clostridium cadaveris		24.64	0.29	NAG 6	99		
Atopobium minutum		0.37	0.08		1		
55040040	**Good identification**						**55040040**
Clostridium bifermentans		98.84	0.36	CEL 1			
Clostridium botulinum biovar A		1.17	0.04	SOR 4			
55041520	**Very good identification**						**55041520**
Odoribacter splanchnicus		100.0	0.61	CEL 1			
55041560	**Good identification**						**55041560**
Odoribacter splanchnicus		100.0	0.45	CEL 1			
55041722	**Good identification**						**55041722**
					CAT		
Bacteroides eggerthii		99.95	0.47	MLT 99	93		
Odoribacter splanchnicus		0.05	-.07	XYL 93	1		
55045722	**Very good identification**						**55045722**
Bacteroides eggerthii		100.0	0.66	MLT 99			
55050000	**Low discrimination**						**55050000**
					GRA	SPO	
Clostridium bifermentans		45.39	0.38	FRU 5	99	99	
Fusobacterium varium		43.62	0.32	CEL 1	1	1	
Eubacterium limosum		5.60	0.18		99	1	
55051520	**Very good identification**						**55051520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Odoribacter splanchnicus		100.0	0.66	CEL 1			
55051560	**Good identification**						**55051560**
Odoribacter splanchnicus		100.0	0.50	CEL 1			
55052010	**Good identification**						**55052010**
					SPO		
Clostridium difficile		99.76	0.44	IND 1	99		
Eubacterium limosum		0.25	-.03	CEL 5	1		
55071120	**Good identification**						**55071120**
					GRA	SPO	
Clostridium septicum		99.02	0.35	IND 1	99	99	
Odoribacter splanchnicus		0.98	-.03	MLT 94	1	1	
55100000	**Excellent identification**						**55100000**
					GRA	SPO	
Clostridium bifermentans		99.89	1.00		99	99	
Clostridium botulinum biovar A		0.06	0.45		99	99	
Prevotella disiens		0.02	0.32		1	1	
55100001	**Low discrimination**						**55100001**
Clostridium bifermentans		71.71	0.66	URE 1			
Clostridium sordellii		28.20	0.55				
Clostridium botulinum biovar A		0.04	0.11				
55100002	**Very good identification**						**55100002**
Clostridium bifermentans		99.47	0.66	BGL 1			
Clostridium sporogenes		0.42	0.24				
Clostridium botulinum biovar A		0.06	0.11				
55100010	**Very good identification**						**55100010**
Clostridium bifermentans		99.88	0.66	MLZ 1			
Clostridium botulinum biovar A		0.06	0.11				
Clostridium novyi biovar A		0.03	0.19				
55100020	**Very good identification**						**55100020**
					COC	SPO	
Clostridium bifermentans		99.23	0.80	NAG 6	1	99	
Clostridium cadaveris		0.74	0.41		1	99	
Gemella morbillorum		0.01	0.14		99	1	
55100021	**Acceptable identification**						**55100021**
Clostridium bifermentans		93.49	0.45	NAG 6			
Clostridium sordellii		5.82	0.20	URE 1			
Clostridium cadaveris		0.69	0.07				
55100022	**Good identification**						**55100022**
Clostridium bifermentans		99.26	0.45	NAG 6			
Clostridium cadaveris		0.74	0.07	BGL 1			
55100030	**Good identification**						**55100030**
Clostridium bifermentans		99.26	0.45	MLZ 1			
Clostridium cadaveris		0.74	0.07	NAG 6			
55100040	**Very good identification**						**55100040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		99.57	0.76	SOR 4			
Clostridium botulinum biovar A		0.41	0.36				
Clostridium novyi biovar A		0.01	0.16				
55100041	**Low discrimination**						**55100041**
Clostridium bifermentans		71.29	0.42	SOR 4			
Clostridium sordellii		28.03	0.31	URE 1			
Clostridium novyi biovar A		0.39	0.16				
55100042	**Good identification**						**55100042**
Clostridium bifermentans		99.59	0.42	SOR 4			
Clostridium botulinum biovar A		0.41	0.02	BGL 1			
55100050	**Good identification**						**55100050**
Clostridium bifermentans		99.06	0.42	MLZ 1			
Clostridium novyi biovar A		0.54	0.16	SOR 4			
Clostridium botulinum biovar A		0.41	0.02				
55100060	**Good identification**						**55100060**
Clostridium bifermentans		99.76	0.56	NAG 6			
Clostridium cadaveris		0.18	0.07	SOR 4			
Clostridium botulinum biovar A		0.07	0.02				
55100100	**Very good identification**						**55100100**
					GRA	SPO	
Clostridium bifermentans		99.85	0.66	LAC 1	99	99	
Clostridium botulinum biovar A		0.06	0.11		99	99	
Prevotella bivia		0.04	-.03		1	1	
55100120	**Low discrimination**						**55100120**
					GRA	SPO	
Clostridium bifermentans		60.06	0.45	LAC 1	99	99	
Prevotella bivia		39.36	0.32	NAG 6	1	1	
Clostridium cadaveris		0.45	0.07		99	99	
55100140	**Good identification**						**55100140**
Clostridium bifermentans		99.59	0.42	LAC 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
55100200	**Very good identification**						**55100200**
					GRA	SPO	
Clostridium bifermentans		99.91	0.66	RHA 1	99	99	
Clostridium botulinum biovar A		0.06	0.11		99	99	
Prevotella disiens		0.02	-.03		1	1	
55100220	**Good identification**						**55100220**
Clostridium bifermentans		99.26	0.45	RHA 1			
Clostridium cadaveris		0.74	0.07	NAG 6			
55100240	**Good identification**						**55100240**
Clostridium bifermentans		99.59	0.42	RHA 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
55100400	**Very good identification**						**55100400**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		99.85	0.66	ARA 1	1	99	
Gemella morbillorum		0.06	0.14		99	1	
Clostridium botulinum biovar A		0.06	0.11		1	99	
55100420	**Good identification**						**55100420**
					GRA	SPO	
Clostridium bifermentans		99.05	0.45	ARA 1	99	99	
Clostridium cadaveris		0.73	0.07	NAG 6	99	99	
Odoribacter splanchnicus		0.22	-.07		1	1	
55100440	**Good identification**						**55100440**
Clostridium bifermentans		99.59	0.42	ARA 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
55100722	**Good identification**						**55100722**
Bacteroides eggerthii		100.0	0.36	GAL 99			
55101000	**Very good identification**						**55101000**
					SPO	GRA	
Clostridium bifermentans		98.38	0.66	GAL 1	99	99	
Eubacterium saburreum		0.70	0.41		1	85	
Prevotella intermedia		0.47	0.29		1	1	
55101001	**Low discrimination**						**55101001**
Clostridium sordellii		87.29	0.42	ESL 5			
Clostridium bifermentans		12.71	0.32	MNS 4			
55101020	**Good identification**						**55101020**
					SPO		
Clostridium bifermentans		96.70	0.45	GAL 1	99		
Eubacterium saburreum		1.90	0.28	NAG 6	1		
Clostridium cadaveris		0.72	0.07		99		
55101040	**Good identification**						**55101040**
Clostridium bifermentans		99.59	0.42	GAL 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
55101100	**Low discrimination**						**55101100**
					COC	SPO	
Eubacterium saburreum		61.39	0.47	SUC 99	1	1	
Clostridium bifermentans		37.47	0.32		1	99	
Blautia hansenii		0.62	-.03		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55101110	**Good identification**						**55101110**
					SPO		
Eubacterium saburreum		99.25	0.46	SUC 99	1		
Clostridium bifermentans		0.75	-.03		99		
55101322	**Good identification**						**55101322**
Bacteroides eggerthii		100.0	0.36	XYL 93			
55101520	**Very good identification**						**55101520**
					CAT		
Odoribacter splanchnicus		99.95	0.61	MLT 1	1		
Bacteroides eggerthii		0.05	0.02		93		
55101522	**Low discrimination**						**55101522**
					CAT		
Bacteroides eggerthii		83.19	0.36	XYL 93	93		
Odoribacter splanchnicus		16.81	0.27	RHA 99	1		
55101560	**Good identification**						**55101560**
Odoribacter splanchnicus		100.0	0.45	MLT 1			
55101622	**Good identification**						**55101622**
Bacteroides eggerthii		100.0	0.36	XYL 93			
55101702	**Good identification**						**55101702**
Bacteroides eggerthii		100.0	0.36	XYL 93			
55101720	**Low discrimination**						**55101720**
					CAT		
Bacteroides eggerthii		83.19	0.36	XYL 93	93		
Odoribacter splanchnicus		16.81	0.27	BGL 99	1		
55101722	**Very good identification**						**55101722**
					CAT		
Bacteroides eggerthii		100.0	0.71	XYL 93	93		
Odoribacter splanchnicus		0.00	-.07		1		
55101723	**Good identification**						**55101723**
Bacteroides eggerthii		100.0	0.36	XYL 93			
55101732	**Good identification**						**55101732**
Bacteroides eggerthii		100.0	0.36	XYL 93			
55101762	**Good identification**						**55101762**
Bacteroides eggerthii		100.0	0.36	XYL 93			
55102000	**Very good identification**						**55102000**
					GRA	SPO	
Clostridium bifermentans		99.89	0.66	MAN 1	99	99	
Clostridium botulinum biovar A		0.06	0.11		99	99	
Prevotella disiens		0.02	-.03		1	1	
55102020	**Good identification**						**55102020**
Clostridium bifermentans		99.26	0.45	MAN 1			
Clostridium cadaveris		0.74	0.07	NAG 6			
55102040	**Good identification**						**55102040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		99.59	0.42	MAN 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
55103722	**Good identification**						**55103722**
Bacteroides eggerthii		100.0	0.36	MAN 1			
55104000	**Very good identification**						**55104000**
					COC	SPO	
Clostridium bifermentans		99.85	0.66	XYL 1	1	99	
Gemella morbillorum		0.06	0.14		99	1	
Clostridium botulinum biovar A		0.06	0.11		1	99	
55104020	**Good identification**						**55104020**
Clostridium bifermentans		99.26	0.45	XYL 1			
Clostridium cadaveris		0.74	0.07	NAG 6			
55104040	**Good identification**						**55104040**
Clostridium bifermentans		96.81	0.42	XYL 1			
Clostridium glycolicum		2.79	0.15	SOR 4			
Clostridium botulinum biovar A		0.40	0.02				
55104722	**Good identification**						**55104722**
Bacteroides eggerthii		100.0	0.56	GAL 99			
55105010	**Good identification**						**55105010**
					SPO		
Eubacterium saburreum		96.84	0.35	SUC 99	1		
Clostridium bifermentans		3.16	-.03		99		
55105100	**Good identification**						**55105100**
					SPO		
Eubacterium saburreum		98.87	0.43	SUC 99	1		
Clostridium bifermentans		1.13	-.03		99		
55105110	**Good identification**						**55105110**
Eubacterium saburreum		100.0	0.41	SUC 99			
55105322	**Good identification**						**55105322**
Bacteroides eggerthii		100.0	0.56	ARA 99			
55105522	**Good identification**						**55105522**
					CAT		
Bacteroides eggerthii		99.99	0.56	RHA 99	93		
Odoribacter splanchnicus		0.02	-.07		1		
55105622	**Good identification**						**55105622**
Bacteroides eggerthii		100.0	0.56	LAC 99			
55105702	**Good identification**						**55105702**
Bacteroides eggerthii		100.0	0.56	NAG 99			
55105720	**Good identification**						**55105720**
					CAT		
Bacteroides eggerthii		99.99	0.56	BGL 99	93		
Odoribacter splanchnicus		0.02	-.07		1		
55105722	**Very good identification**						**55105722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides eggerthii		100.0	0.90				
55105723	**Good identification**						**55105723**
Bacteroides eggerthii		100.0	0.56	URE 1			
55105732	**Good identification**						**55105732**
Bacteroides eggerthii		100.0	0.56	MLZ 1			
55105762	**Good identification**						**55105762**
Bacteroides eggerthii		100.0	0.56	SOR 1			
55107722	**Good identification**						**55107722**
Bacteroides eggerthii		100.0	0.56	MAN 1			
55110000	**Good identification**						**55110000**
					GRA	SPO	
Clostridium bifermentans		90.67	0.78	FRU 5	99	99	
Prevotella intermedia		8.24	0.63		1	1	
Fusobacterium varium		0.38	0.32		1	1	
55110001	**Low discrimination**						**55110001**
					GRA	SPO	
Clostridium sordellii		87.18	0.55	ESL 5	99	99	
Clostridium bifermentans		11.67	0.44	MNS 4	99	99	
Prevotella intermedia		1.06	0.29		1	1	
55110002	**Low discrimination**						**55110002**
					GRA	SPO	
Clostridium bifermentans		89.45	0.44	FRU 5	99	99	
Prevotella intermedia		8.13	0.29	BGL 1	1	1	
Clostridium sporogenes		1.28	0.11		99	99	
55110010	**Low discrimination**						**55110010**
					GRA	SPO	
Clostridium bifermentans		88.43	0.44	FRU 5	99	99	
Prevotella intermedia		8.04	0.29	MLZ 1	1	1	
Clostridium difficile		2.52	0.15		99	99	
55110020	**Low discrimination**						**55110020**
					GRA	SPO	
Clostridium bifermentans		86.12	0.58	FRU 5	99	99	
Clostridium cadaveris		12.12	0.41	NAG 6	99	99	
Prevotella intermedia		1.24	0.29		1	1	
55110040	**Good identification**						**55110040**
					GRA	SPO	
Clostridium bifermentans		96.64	0.55	FRU 5	99	99	
Prevotella intermedia		2.13	0.29	SOR 4	1	1	
Clostridium botulinum biovar A		0.48	0.16		99	99	
55110100	**Low discrimination**						**55110100**
					GRA	SPO	
Clostridium bifermentans		89.28	0.44	FRU 5	99	99	
Prevotella intermedia		8.12	0.29	LAC 1	1	1	
Fusobacterium mortiferum		1.04	0.25		1	1	
55110200	**Acceptable identification**						**55110200**
					GRA	SPO	
Clostridium bifermentans		90.99	0.44	FRU 5	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella intermedia		8.27	0.29	RHA 1	1	1		
Fusobacterium varium		0.38	-.03		1	1		
55110400	**Low discrimination**							**55110400**
					COC	GRA	SPO	
Clostridium bifermentans		89.89	0.44	FRU 5	1	99	99	
Prevotella intermedia		8.17	0.29	ARA 1	1	1	1	
Gemella morbillorum		1.21	0.15		99	99	1	
55111000	**Low discrimination**							**55111000**
					SPO	GRA		
Prevotella intermedia		85.71	0.63	ESL 1	1	1		
Clostridium bifermentans		9.52	0.44		99	99		
Eubacterium saburreum		3.85	0.49		1	85		
55111001	**Acceptable identification**							**55111001**
					GRA	SPO		
Clostridium sordellii		92.62	0.42	ESL 5	99	99		
Prevotella intermedia		6.39	0.29	MNS 4	1	1		
Clostridium bifermentans		0.71	0.10		99	99		
55111010	**Low discrimination**							**55111010**
					SPO	GRA		
Eubacterium saburreum		76.60	0.48	SUC 99	1	85		
Prevotella intermedia		21.06	0.29		1	1		
Clostridium bifermentans		2.34	0.10		99	99		
55111100	**Low discrimination**							**55111100**
Eubacterium saburreum		87.32	0.56	SUC 99				
Prevotella intermedia		8.42	0.29					
Fusobacterium mortiferum		1.61	0.28					
55111102	**Good identification**							**55111102**
					GRA			
Eubacterium saburreum		97.42	0.43	SUC 99	85			
Clostridium perfringens		1.45	0.02		99			
Fusobacterium mortiferum		1.13	0.11		1			
55111110	**Good identification**							**55111110**
Eubacterium saburreum		100.0	0.54	SUC 99				
55111112	**Good identification**							**55111112**
Eubacterium saburreum		100.0	0.41	SUC 99				
55111120	**Low discrimination**							**55111120**
					GRA	SPO		
Clostridium septicum		48.87	0.39	IND 1	99	99		
Odoribacter splanchnicus		28.97	0.32		1	1		
Clostridium perfringens		10.00	0.34		99	84		
55111130	**Acceptable identification**							**55111130**
					SPO	GRA		
Eubacterium saburreum		91.08	0.41	SUC 99	1	85		
Clostridium septicum		5.60	0.05		99	99		
Odoribacter splanchnicus		3.32	-.03		1	1		
55111520	**Very good identification**							**55111520**
					GRA	SPO		
Odoribacter splanchnicus		99.98	0.66	MLT 1	1	1		
Clostridium septicum		0.02	0.05		99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
55111560	**Good identification**						**55111560**
Odoribacter splanchnicus		100.0	0.50	MLT 1			
55111722	**Good identification**						**55111722**
					CAT		
Bacteroides eggerthii		99.62	0.36	FRU 1	93		
Odoribacter splanchnicus		0.38	-.03	XYL 93	1		
55112000	**Low discrimination**						**55112000**
					GRA	SPO	
Clostridium bifermentans		67.73	0.44	FRU 5	99	99	
Eubacterium limosum		22.01	0.32	MAN 1	99	1	
Prevotella intermedia		6.16	0.29		1	1	
55114000	**Low discrimination**						**55114000**
					GRA	SPO	
Clostridium bifermentans		87.84	0.44	FRU 5	99	99	
Prevotella intermedia		7.99	0.29	XYL 1	1	1	
Fusobacterium varium		1.51	0.08		1	1	
55115000	**Low discrimination**						**55115000**
					SPO	GRA	
Eubacterium saburreum		66.78	0.45	SUC 99	1	85	
Prevotella intermedia		27.90	0.29		1	1	
Clostridium bifermentans		3.10	0.10		99	99	
55115010	**Good identification**						**55115010**
Eubacterium saburreum		100.0	0.43	SUC 99			
55115100	**Good identification**						**55115100**
					COC		
Eubacterium saburreum		99.63	0.51	SUC 99	1		
Atopobium parvulum		0.37	0.00		99		
55115102	**Good identification**						**55115102**
Eubacterium saburreum		100.0	0.38	SUC 99			
55115110	**Good identification**						**55115110**
Eubacterium saburreum		100.0	0.49	SUC 99			
55115112	**Good identification**						**55115112**
Eubacterium saburreum		100.0	0.37	SUC 99			
55115120	**Low discrimination**						**55115120**
					SPO	GRA	
Eubacterium saburreum		87.05	0.38	SUC 99	1	85	
Clostridium septicum		8.13	0.05		99	99	
Odoribacter splanchnicus		4.82	-.03		1	1	
55115130	**Good identification**						**55115130**
Eubacterium saburreum		100.0	0.37	SUC 99			
55115722	**Good identification**						**55115722**
Bacteroides eggerthii		99.87	0.56	FRU 1			
Bacteroides thetaiotaomicron		0.13	0.12				
55120000	**Very good identification**						**55120000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		98.16	0.66	TRE 1			
Clostridium botulinum biovar A		1.40	0.35				
Clostridium sporogenes		0.42	0.24				
55120020	**Good identification**						**55120020**
Clostridium bifermentans		96.07	0.45	TRE 1			
Clostridium cadaveris		3.71	0.19	NAG 6			
Clostridium botulinum biovar A		0.22	0.01				
55120040	**Low discrimination**						**55120040**
Clostridium bifermentans		89.98	0.42	TRE 1			
Clostridium botulinum biovar A		9.20	0.26	SOR 4			
Clostridium botulinum biovar B		0.82	0.05				
55121010	**Good identification**						**55121010**
					SPO		
Eubacterium saburreum		97.43	0.37	SUC 99	1		
Clostridium bifermentans		2.57	-.03		99		
55121100	**Good identification**						**55121100**
					SPO		
Eubacterium saburreum		99.08	0.44	SUC 99	1		
Clostridium bifermentans		0.92	-.03		99		
55121110	**Good identification**						**55121110**
Eubacterium saburreum		100.0	0.43	SUC 99			
55121722	**Good identification**						**55121722**
Bacteroides eggerthii		100.0	0.36	TRE 1			
55125100	**Good identification**						**55125100**
					COC		
Eubacterium saburreum		98.34	0.40	SUC 99	1		
Atopobium parvulum		1.66	0.00		99		
55125110	**Good identification**						**55125110**
Eubacterium saburreum		100.0	0.38	SUC 99			
55125722	**Good identification**						**55125722**
Bacteroides eggerthii		99.98	0.56	TRE 1			
Bacteroides thetaiotaomicron		0.02	-.03				
55130000	**Low discrimination**						**55130000**
					GRA	SPO	
Clostridium bifermentans		88.24	0.44	FRU 5	99	99	
Prevotella intermedia		8.02	0.29	TRE 1	1	1	
Clostridium botulinum biovar A		1.53	0.14		99	99	
55131000	**Low discrimination**						**55131000**
					SPO	GRA	
Eubacterium saburreum		71.62	0.46	SUC 99	1	85	
Prevotella intermedia		24.16	0.29		1	1	
Clostridium bifermentans		2.68	0.10		99	99	
55131010	**Good identification**						**55131010**
Eubacterium saburreum		100.0	0.45	SUC 99			
55131100	**Low discrimination**						**55131100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Eubacterium saburreum		81.61	0.53	SUC 99	1	1	
Atopobium parvulum		13.19	0.29		99	1	
Clostridium septicum		3.26	0.13		1	99	
55131102	**Low discrimination**						**55131102**
					COC		
Eubacterium saburreum		83.13	0.40	SUC 99	1		
Atopobium parvulum		13.43	0.16		99		
Clostridium perfringens		3.43	0.07		1		
55131110	**Good identification**						**55131110**
Eubacterium saburreum		100.0	0.51	SUC 99			
55131112	**Good identification**						**55131112**
Eubacterium saburreum		100.0	0.38	SUC 99			
55131120	**Low discrimination**						**55131120**
					SPO		
Clostridium septicum		84.71	0.47	IND 1	99		
Clostridium perfringens		10.73	0.38		84		
Eubacterium saburreum		3.78	0.40		1		
55131130	**Low discrimination**						**55131130**
					SPO		
Eubacterium saburreum		76.24	0.38	SUC 99	1		
Clostridium septicum		21.09	0.13		99		
Clostridium perfringens		2.67	0.04		84		
55135000	**Good identification**						**55135000**
					COC		
Eubacterium saburreum		98.70	0.42	SUC 99	1		
Atopobium parvulum		1.30	0.00		99		
55135010	**Good identification**						**55135010**
Eubacterium saburreum		100.0	0.40	SUC 99			
55135100	**Low discrimination**						**55135100**
					COC		
Eubacterium saburreum		64.21	0.48	SUC 99	1		
Atopobium parvulum		35.79	0.34		99		
55135102	**Low discrimination**						**55135102**
					COC		
Eubacterium saburreum		64.21	0.35	SUC 99	1		
Atopobium parvulum		35.79	0.21		99		
55135110	**Good identification**						**55135110**
					COC		
Eubacterium saburreum		99.32	0.46	SUC 99	1		
Atopobium parvulum		0.68	0.00		99		
55135120	**Low discrimination**						**55135120**
					COC	SPO	
Eubacterium saburreum		49.24	0.35	SUC 99	1	1	
Atopobium parvulum		27.45	0.21		99	1	
Clostridium septicum		20.69	0.13		1	99	
55140000	**Very good identification**						**55140000**
					GRA	SPO	
Clostridium bifermentans		99.91	0.66	CEL 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		0.06	0.11		99	99	
Prevotella disiens		0.02	-.03		1	1	
55140020	**Good identification**						**55140020**
Clostridium bifermentans		99.26	0.45	CEL 1			
Clostridium cadaveris		0.74	0.07	NAG 6			
55140040	**Good identification**						**55140040**
Clostridium bifermentans		99.59	0.42	CEL 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
55140722	**Good identification**						**55140722**
Bacteroides eggerthii		100.0	0.47	GAL 99			
55141322	**Good identification**						**55141322**
Bacteroides eggerthii		100.0	0.47	XYL 93			
55141522	**Good identification**						**55141522**
					CAT		
Bacteroides eggerthii		99.95	0.47	XYL 93	93		
Odoribacter splanchnicus		0.05	-.07	RHA 99	1		
55141622	**Good identification**						**55141622**
Bacteroides eggerthii		100.0	0.47	XYL 93			
55141702	**Good identification**						**55141702**
Bacteroides eggerthii		100.0	0.47	XYL 93			
55141720	**Good identification**						**55141720**
					CAT		
Bacteroides eggerthii		99.95	0.47	XYL 93	93		
Odoribacter splanchnicus		0.05	-.07	BGL 99	1		
55141722	**Very good identification**						**55141722**
Bacteroides eggerthii		100.0	0.81	XYL 93			
55141723	**Good identification**						**55141723**
Bacteroides eggerthii		100.0	0.47	XYL 93			
55141732	**Good identification**						**55141732**
Bacteroides eggerthii		100.0	0.47	XYL 93			
55141762	**Good identification**						**55141762**
Bacteroides eggerthii		100.0	0.47	XYL 93			
55143722	**Good identification**						**55143722**
Bacteroides eggerthii		100.0	0.47	MAN 1			
55144722	**Very good identification**						**55144722**
Bacteroides eggerthii		100.0	0.66	GAL 99			
55145322	**Very good identification**						**55145322**
Bacteroides eggerthii		100.0	0.66	ARA 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55145522	**Very good identification**						**55145522**
Bacteroides eggerthii		100.0	0.66	RHA 99			
55145622	**Very good identification**						**55145622**
Bacteroides eggerthii		100.0	0.66	LAC 99			
55145702	**Very good identification**						**55145702**
Bacteroides eggerthii		100.0	0.66	NAG 99			
55145720	**Very good identification**						**55145720**
Bacteroides eggerthii		100.0	0.66	BGL 99			
55145722	**Excellent identification**						**55145722**
Bacteroides eggerthii		100.0	1.00				
55145723	**Very good identification**						**55145723**
Bacteroides eggerthii		100.0	0.66	URE 1			
55145732	**Very good identification**						**55145732**
Bacteroides eggerthii		100.0	0.66	MLZ 1			
55145762	**Very good identification**						**55145762**
Bacteroides eggerthii		100.0	0.66	SOR 1			
55147722	**Very good identification**						**55147722**
Bacteroides eggerthii		100.0	0.66	MAN 1			
55150000	**Acceptable identification**						**55150000**
					GRA	SPO	
Clostridium bifermentans		90.99	0.44	FRU 5	99	99	
Prevotella intermedia		8.27	0.29	CEL 1	1	1	
Fusobacterium varium		0.38	-.03		1	1	
55151120	**Good identification**						**55151120**
Clostridium septicum		97.76	0.47	IND 1			
Clostridium paraputrificum		1.53	0.14				
Clostridium perfringens		0.26	0.10				
55151722	**Good identification**						**55151722**
Bacteroides eggerthii		99.96	0.47	FRU 1			
Bacteroides thetaiotaomicron		0.04	-.06	XYL 93			
55155722	**Very good identification**						**55155722**
					CAT		
Bacteroides eggerthii		99.75	0.66	FRU 1	93		
Bacteroides uniformis		0.18	0.22		5		
Bacteroides thetaiotaomicron		0.07	0.18		89		
55161722	**Good identification**						**55161722**
Bacteroides eggerthii		100.0	0.47	TRE 1			
55165722	**Very good identification**						**55165722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides eggerthii		99.99	0.66	TRE 1			
Bacteroides thetaiotaomicron		0.01	0.03				
55171100	**Low discrimination**						**55171100**
					COC	SPO	
Atopobium parvulum		73.41	0.36	IND 1	99	1	
Clostridium septicum		24.62	0.21	SUC 96	1	99	
Eubacterium saburreum		1.97	0.18		1	1	
55171120	**Good identification**						**55171120**
					COC	SPO	
Clostridium septicum		99.17	0.56	IND 1	1	99	
Atopobium parvulum		0.53	0.23		99	1	
Clostridium perfringens		0.17	0.15		1	84	
55175100	**Good identification**						**55175100**
					COC		
Atopobium parvulum		99.23	0.40	IND 1	99		
Eubacterium saburreum		0.77	0.14	SUC 96	1		
55200000	**Very good identification**						**55200000**
					SPO		
Clostridium bifermentans		99.97	0.77	SAL 10	99		
Clostridium botulinum biovar A		0.02	0.13		99		
Atopobium minutum		0.01	0.16		1		
55200001	**Good identification**						**55200001**
Clostridium bifermentans		99.08	0.43	SAL 10			
Clostridium sordellii		0.92	0.04	URE 1			
55200002	**Good identification**						**55200002**
					SPO		
Clostridium bifermentans		99.44	0.43	SAL 10	99		
Atopobium minutum		0.56	0.13	BGL 1	1		
55200010	**Good identification**						**55200010**
Clostridium bifermentans		99.12	0.43	SAL 10			
Clostridium difficile		0.88	0.06	MLZ 1			
55200020	**Very good identification**						**55200020**
					SPO		
Clostridium bifermentans		97.06	0.57	SAL 10	99		
Clostridium cadaveris		2.90	0.29	NAG 6	99		
Atopobium minutum		0.04	0.08		1		
55200040	**Good identification**						**55200040**
Clostridium bifermentans		99.89	0.54	SAL 10			
Clostridium botulinum biovar A		0.11	0.04	SOR 4			
55200100	**Good identification**						**55200100**
					GRA	SPO	
Clostridium bifermentans		99.80	0.43	SAL 10	99	99	
Fusobacterium mortiferum		0.20	0.11	LAC 1	1	1	
55200200	**Good identification**						**55200200**
Clostridium bifermentans		100.0	0.43	SAL 10			
55200400	**Good identification**						**55200400**
Clostridium bifermentans		100.0	0.43	SAL 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
55201000	**Good identification**						**55201000**
					SPO		
Clostridium bifermentans		99.40	0.43	SAL 10	99		
Eubacterium tortuosum		0.32	-.03	GAL 1	1		
Atopobium minutum		0.28	0.08		1		
55201520	**Very good identification**						**55201520**
Odoribacter splanchnicus		100.0	0.61	SAL 1			
55201560	**Good identification**						**55201560**
Odoribacter splanchnicus		100.0	0.45	SAL 1			
55202000	**Good identification**						**55202000**
					SPO		
Clostridium bifermentans		98.04	0.43	SAL 10	99		
Clostridium difficile		1.61	0.11	MAN 1	99		
Eubacterium limosum		0.36	-.03		1		
55204000	**Good identification**						**55204000**
Clostridium bifermentans		100.0	0.43	SAL 10			
55210000	**Low discrimination**						**55210000**
					GRA	SPO	
Clostridium bifermentans		86.86	0.56	SAL 10	99	99	
Fusobacterium varium		7.59	0.32	FRU 5	1	1	
Clostridium difficile		3.00	0.28		99	99	
55210010	**Acceptable identification**						**55210010**
					GRA	SPO	
Clostridium difficile		93.89	0.40	IND 1	99	99	
Clostridium bifermentans		5.62	0.21	MAN 90	99	99	
Fusobacterium varium		0.49	-.03		1	1	
55210020	**Low discrimination**						**55210020**
					SPO	CAT	
Clostridium bifermentans		61.09	0.35	SAL 10	99	1	
Clostridium cadaveris		34.66	0.29	FRU 5	99	1	
Propionibacterium acnes		2.93	0.14	NAG 6	1	95	
55210100	**Low discrimination**						**55210100**
					GRA	SPO	
Fusobacterium mortiferum		77.91	0.45	IND 1	1	1	
Clostridium bifermentans		20.32	0.21		99	99	
Fusobacterium varium		1.78	-.03		1	1	
55211000	**Low discrimination**						**55211000**
					GRA	SPO	
Fusobacterium mortiferum		68.44	0.42	IND 1	1	1	
Clostridium bifermentans		27.76	0.21		99	99	
Fusobacterium varium		2.43	-.03		1	1	
55211100	**Good identification**						**55211100**
Fusobacterium mortiferum		96.66	0.48	IND 1			
Odoribacter splanchnicus		1.75	-.03				
Eubacterium saburreum		0.84	0.11				
55211520	**Very good identification**						**55211520**
					GRA	SPO	
Odoribacter splanchnicus		100.0	0.66	SAL 1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		0.00	-.06		99	99	
55211560	**Good identification**						**55211560**
Odoribacter splanchnicus		100.0	0.50	SAL 1			
55212000	**Low discrimination**						**55212000**
					SPO		
Clostridium difficile		79.56	0.45	IND 1	99		
Eubacterium limosum		17.63	0.32		1		
Clostridium bifermentans		2.58	0.21		99		
55212010	**Very good identification**						**55212010**
					SPO		
Clostridium difficile		99.95	0.56	IND 1	99		
Eubacterium limosum		0.05	-.03		1		
55220000	**Good identification**						**55220000**
Clostridium bifermentans		99.63	0.43	SAL 10			
Clostridium botulinum biovar A		0.37	0.02	TRE 1			
55230100	**Good identification**						**55230100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
55231100	**Acceptable identification**						**55231100**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	IND 1	1	1	
Clostridium septicum		7.27	0.03		99	99	
55231120	**Good identification**						**55231120**
					GRA	SPO	
Clostridium septicum		99.22	0.37	IND 1	99	99	
Odoribacter splanchnicus		0.78	-.03	MLT 94	1	1	
55232010	**Good identification**						**55232010**
Clostridium difficile		100.0	0.37	IND 1			
55240000	**Good identification**						**55240000**
Clostridium bifermentans		100.0	0.43	SAL 10			
55250100	**Good identification**						**55250100**
Fusobacterium mortiferum		100.0	0.37	IND 1			
55251100	**Low discrimination**						**55251100**
					GRA		
Fusobacterium mortiferum		73.20	0.40	IND 1	1		
Lactobacillus catenaformis		10.30	0.15		99		
Collinsella aerofaciens		6.82	-.03		99		
55251120	**Acceptable identification**						**55251120**
Clostridium septicum		91.15	0.37	IND 1			
Clostridium paraputrificum		5.57	0.14	MLT 94			
Clostridium barati		2.17	0.12				
55253000	**Good identification**						**55253000**
Clostridium innocuum		100.0	0.49	IND 1			
55253002	**Good identification**						**55253002**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		100.0	0.49	IND 1			
55253400	**Good identification**						**55253400**
Clostridium innocuum		100.0	0.36	IND 1			
55253402	**Good identification**						**55253402**
Clostridium innocuum		100.0	0.36	IND 1			
55271120	**Good identification**						**55271120**
Clostridium septicum		99.46	0.45	IND 1			
Clostridium paraputrificum		0.54	0.04	MLT 94			
55273000	**Good identification**						**55273000**
Clostridium innocuum		100.0	0.41	IND 1			
55273002	**Good identification**						**55273002**
Clostridium innocuum		100.0	0.41	IND 1			
55300000	**Very good identification**						**55300000**
					GRA	SPO	
Clostridium bifermentans		99.99	0.84	SAL 10	99	99	
Clostridium botulinum biovar A		0.01	0.11		99	99	
Prevotella disiens		0.00	-.03		1	1	
55300001	**Good identification**						**55300001**
Clostridium bifermentans		96.55	0.50	SAL 10			
Clostridium sordellii		3.45	0.20	URE 1			
55300002	**Good identification**						**55300002**
Clostridium bifermentans		100.0	0.50	SAL 10			
55300010	**Good identification**						**55300010**
Clostridium bifermentans		100.0	0.50	SAL 10			
55300020	**Very good identification**						**55300020**
Clostridium bifermentans		99.93	0.63	SAL 10			
Clostridium cadaveris		0.07	0.07	NAG 6			
55300040	**Very good identification**						**55300040**
Clostridium bifermentans		99.96	0.60	SAL 10			
Clostridium botulinum biovar A		0.04	0.02	SOR 4			
55300060	**Good identification**						**55300060**
Clostridium bifermentans		100.0	0.40	SAL 10			
55300100	**Good identification**						**55300100**
Clostridium bifermentans		100.0	0.50	SAL 10			
55300200	**Good identification**						**55300200**
Clostridium bifermentans		100.0	0.50	SAL 10			
55300400	**Good identification**						**55300400**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		100.0	0.50	SAL 10			
55301000	**Good identification**						**55301000**
Clostridium bifermentans		100.0	0.50	SAL 10			
55301722	**Good identification**						**55301722**
Bacteroides eggerthii		100.0	0.36	SAL 1			
55302000	**Good identification**						**55302000**
Clostridium bifermentans		100.0	0.50	SAL 10			
55304000	**Good identification**						**55304000**
Clostridium bifermentans		100.0	0.50	SAL 10			
55305722	**Good identification**						**55305722**
Bacteroides eggerthii		100.0	0.56	SAL 1			
55310000	**Very good identification**						**55310000**
					GRA	SPO	
Clostridium bifermentans		99.02	0.62	SAL 10	99	99	
Prevotella intermedia		0.82	0.29	FRU 5	1	1	
Fusobacterium mortiferum		0.09	0.23		1	1	
55310020	**Good identification**						**55310020**
Clostridium bifermentans		98.74	0.41	SAL 10			
Clostridium cadaveris		1.26	0.07	FRU 5			
55310040	**Good identification**						**55310040**
Clostridium bifermentans		100.0	0.38	SAL 10			
55311120	**Good identification**						**55311120**
Clostridium septicum		97.34	0.49	IND 1			
Clostridium perfringens		1.25	0.23				
Clostridium paraputrificum		1.20	0.14				
55320000	**Good identification**						**55320000**
Clostridium bifermentans		99.87	0.50	SAL 10			
Clostridium botulinum biovar A		0.13	0.01	TRE 1			
55331120	**Very good identification**						**55331120**
					COC	SPO	
Clostridium septicum		98.87	0.57	IND 1	1	99	
Clostridium perfringens		0.78	0.28		1	84	
Atopobium parvulum		0.24	0.18		99	1	
55335100	**Good identification**						**55335100**
					COC		
Atopobium parvulum		98.65	0.36	IND 1	99		
Eubacterium saburreum		1.35	0.14	SUC 96	1		
55340000	**Good identification**						**55340000**
Clostridium bifermentans		100.0	0.50	SAL 10			
55341722	**Good identification**						**55341722**
Bacteroides eggerthii		100.0	0.47	SAL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55345722	**Very good identification**						**55345722**
					CAT		
Bacteroides eggerthii		99.98	0.66	SAL 1	93		
Bacteroides uniformis		0.02	0.06		5		
55351000	**Low discrimination**						**55351000**
					SPO		
Lactobacillus jensenii		81.12	0.41	IND 1	1		
Lactobacillus catenaformis		14.69	0.29	SUC 90	1		
Clostridium barati		1.65	0.07		99		
55351002	**Low discrimination**						**55351002**
					SPO		
Lactobacillus jensenii		73.01	0.38	IND 1	1		
Lactobacillus catenaformis		19.83	0.29	SUC 90	1		
Clostridium barati		6.67	0.15		99		
55351120	**Low discrimination**						**55351120**
					SPO		
Clostridium septicum		71.11	0.58	IND 1	99		
Clostridium paraputrificum		27.48	0.48		99		
Collinsella aerofaciens		0.63	0.15		1		
55351122	**Low discrimination**						**55351122**
Clostridium paraputrificum		72.39	0.41	IND 1			
Clostridium barati		9.72	0.32	SUC 92			
Clostridium septicum		8.92	0.29				
55355522	**Good identification**						**55355522**
					CAT		
Bacteroides uniformis		99.31	0.38	SUC 99	5		
Bacteroides eggerthii		0.69	-.03	RAF 97	93		
55355722	**Low discrimination**						**55355722**
					CAT		
Bacteroides uniformis		65.47	0.40	SUC 99	5		
Bacteroides eggerthii		32.69	0.32	RAF 97	93		
Bacteroides ovatus		1.44	0.15		68		
55371000	**Good identification**						**55371000**
					COC	SPO	
Lactobacillus jensenii		99.50	0.50	IND 1	1	1	
Clostridium septicum		0.25	-.03	SUC 90	1	99	
Atopobium parvulum		0.25	0.04		99	1	
55371002	**Good identification**						**55371002**
					SPO		
Lactobacillus jensenii		98.99	0.47	IND 1	1		
Lactobacillus acidophilus		0.74	0.07	SUC 90	1		
Clostridium barati		0.26	-.03		99		
55371010	**Good identification**						**55371010**
Lactobacillus jensenii		100.0	0.41	IND 1			
55371012	**Good identification**						**55371012**
					GRA		
Lactobacillus jensenii		97.78	0.38	IND 1	99		
Leptotrichia buccalis		2.22	-.03	SUC 90	1		
55371020	**Low discrimination**						**55371020**
					SPO		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		49.09	0.39	IND 1	1		
Clostridium septicum		48.89	0.32	SUC 90	99		
Clostridium paraputrificum		1.67	0.04		99		
55371022	**Low discrimination**						**55371022**
					SPO		
Lactobacillus jensenii		89.75	0.36	IND 1	1		
Clostridium barati		2.85	0.05	SUC 90	99		
Clostridium septicum		2.74	0.03		99		
55371120	**Very good identification**						**55371120**
					COC	SPO	
Clostridium septicum		96.48	0.66	IND 1	1	99	
Clostridium paraputrificum		3.30	0.38		1	99	
Atopobium parvulum		0.17	0.25		99	1	
55371122	**Low discrimination**						**55371122**
					SPO		
Clostridium septicum		52.30	0.37	IND 1	99		
Clostridium paraputrificum		37.61	0.32	BGL 2	99		
Lactobacillus acidophilus		5.99	0.25		1		
55375100	**Good identification**						**55375100**
					COC	SPO	
Atopobium parvulum		98.33	0.42	IND 1	99	1	
Clostridium butyricum		1.13	0.02	SUC 96	1	99	
Clostridium septicum		0.54	-.03		1	99	
55400000	**Very good identification**						**55400000**
					SPO		
Clostridium bifermentans		99.69	0.60	RAF 1	99		
Clostridium botulinum biovar A		0.16	0.13		99		
Atopobium minutum		0.09	0.16		1		
55400020	**Low discrimination**						**55400020**
					SPO		
Clostridium bifermentans		75.00	0.39	RAF 1	99		
Clostridium cadaveris		24.64	0.29	NAG 6	99		
Atopobium minutum		0.37	0.08		1		
55400040	**Good identification**						**55400040**
Clostridium bifermentans		98.84	0.36	RAF 1			
Clostridium botulinum biovar A		1.17	0.04	SOR 4			
55401520	**Very good identification**						**55401520**
Odoribacter splanchnicus		100.0	0.61	RAF 1			
55401560	**Good identification**						**55401560**
Odoribacter splanchnicus		100.0	0.45	RAF 1			
55410000	**Low discrimination**						**55410000**
					GRA	SPO	
Clostridium bifermentans		38.24	0.38	RAF 1	99	99	
Fusobacterium varium		36.75	0.32	FRU 5	1	1	
Prevotella intermedia		12.17	0.32		1	1	
55410100	**Good identification**						**55410100**
					GRA	SPO	
Fusobacterium mortiferum		95.56	0.45	IND 1	1	1	
Clostridium bifermentans		2.27	0.04		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium varium		2.18	-.03		1	1	
55411100	**Low discrimination**						**55411100**
Fusobacterium mortiferum		67.49	0.48	IND 1			
Eubacterium saburreum		31.29	0.41				
Odoribacter splanchnicus		1.22	-.03				
55411110	**Good identification**						**55411110**
Eubacterium saburreum		97.41	0.39	SUC 99			
Fusobacterium mortiferum		2.59	0.14				
55411520	**Very good identification**						**55411520**
Odoribacter splanchnicus		100.0	0.66	RAF 1			
55411560	**Good identification**						**55411560**
Odoribacter splanchnicus		100.0	0.50	RAF 1			
55415100	**Low discrimination**						**55415100**
Eubacterium saburreum		76.83	0.36	SUC 99			
Fusobacterium mortiferum		23.17	0.29				
55430100	**Good identification**						**55430100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
55431100	**Low discrimination**						**55431100**
Eubacterium saburreum		50.85	0.38	SUC 99			
Fusobacterium mortiferum		49.15	0.39				
55431110	**Good identification**						**55431110**
Eubacterium saburreum		100.0	0.36	SUC 99			
55435122	**Good identification**						**55435122**
Parabacteroides distasonis		100.0	0.37	IND 1			
55435322	**Acceptable identification**						**55435322**
Parabacteroides distasonis		92.49	0.46	IND 1			
Bacteroides thetaiotaomicron		7.51	0.11				
55435332	**Acceptable identification**						**55435332**
Parabacteroides distasonis		94.46	0.41	IND 1			
Bacteroides thetaiotaomicron		5.54	0.04				
55435722	**Acceptable identification**						**55435722**
Bacteroides thetaiotaomicron		92.92	0.45	SUC 98			
Parabacteroides distasonis		7.08	0.42	MLT 94			
55435732	**Acceptable identification**						**55435732**
Bacteroides thetaiotaomicron		90.46	0.38	SUC 98			
Parabacteroides distasonis		9.55	0.37	MLT 94			
55450100	**Good identification**						**55450100**
Fusobacterium mortiferum		100.0	0.37	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55451100	**Good identification**						**55451100**
Fusobacterium mortiferum		100.0	0.40	IND 1			
55475322	**Low discrimination**						**55475322**
Parabacteroides distasonis		75.72	0.41	IND 1			
Bacteroides thetaiotaomicron		24.28	0.17				
55475332	**Low discrimination**						**55475332**
Parabacteroides distasonis		81.20	0.36	IND 1			
Bacteroides thetaiotaomicron		18.80	0.09				
55475722	**Good identification**						**55475722**
Bacteroides thetaiotaomicron		97.02	0.51	SUC 98			
Parabacteroides distasonis		1.87	0.37	MLT 94			
Bacteroides ovatus		1.03	0.17				
55475732	**Good identification**						**55475732**
Bacteroides thetaiotaomicron		95.71	0.44	SUC 98			
Parabacteroides distasonis		2.56	0.33	MLT 94			
Bacteroides ovatus		1.74	0.14				
55500000	**Very good identification**						**55500000**
					GRA	SPO	
Clostridium bifermentans		99.20	0.66	RAF 1	99	99	
Prevotella intermedia		0.71	0.32		1	1	
Clostridium botulinum biovar A		0.06	0.11		99	99	
55500020	**Good identification**						**55500020**
Clostridium bifermentans		99.26	0.45	RAF 1			
Clostridium cadaveris		0.74	0.07	NAG 6			
55500040	**Good identification**						**55500040**
Clostridium bifermentans		99.59	0.42	RAF 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
55501010	**Good identification**						**55501010**
					SPO		
Eubacterium saburreum		96.84	0.35	SUC 99	1		
Clostridium bifermentans		3.16	-.03		99		
55501110	**Good identification**						**55501110**
					COC		
Eubacterium saburreum		97.77	0.41	SUC 99	1		
Blautia hansenii		2.23	-.03		99		
55501722	**Good identification**						**55501722**
Bacteroides eggerthii		100.0	0.36	RAF 1			
55505100	**Low discrimination**						**55505100**
					COC		
Eubacterium saburreum		62.25	0.38	SUC 99	1		
Blautia hansenii		37.75	0.19		99		
55505110	**Good identification**						**55505110**
Eubacterium saburreum		100.0	0.37	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55505722	**Good identification**						**55505722**
Bacteroides eggerthii		99.87	0.56	RAF 1			
Bacteroides thetaiotaomicron		0.13	0.12				
55510000	**Good identification**						**55510000**
					GRA	SPO	
Prevotella intermedia		92.98	0.66	ESL 1	1	1	
Clostridium bifermentans		6.89	0.44		99	99	
Fusobacterium mortiferum		0.07	0.23		1	1	
55511000	**Very good identification**						**55511000**
Prevotella intermedia		98.24	0.66	ESL 1			
Eubacterium saburreum		1.58	0.45				
Fusobacterium mortiferum		0.11	0.26				
55511010	**Low discrimination**						**55511010**
Eubacterium saburreum		56.63	0.43	SUC 99			
Prevotella intermedia		43.37	0.32				
55511100	**Low discrimination**						**55511100**
Eubacterium saburreum		74.31	0.51	SUC 99			
Prevotella intermedia		19.95	0.32				
Fusobacterium mortiferum		4.95	0.33				
55511102	**Low discrimination**						**55511102**
					CAT		
Eubacterium saburreum		72.12	0.38	SUC 99	1		
Prevotella oralis		11.74	0.12		1		
Capnocytophaga ochracea		11.64	0.12		90		
55511110	**Good identification**						**55511110**
Eubacterium saburreum		100.0	0.49	SUC 99			
55511112	**Good identification**						**55511112**
Eubacterium saburreum		100.0	0.37	SUC 99			
55511130	**Good identification**						**55511130**
Eubacterium saburreum		97.10	0.37	SUC 99			
Prevotella melaninogenica		2.90	-.06				
55515010	**Good identification**						**55515010**
Eubacterium saburreum		100.0	0.39	SUC 99			
55515100	**Good identification**						**55515100**
Eubacterium saburreum		99.08	0.46	SUC 99			
Fusobacterium mortiferum		0.92	0.13				
55515110	**Good identification**						**55515110**
Eubacterium saburreum		100.0	0.45	SUC 99			
55515322	**Low discrimination**						**55515322**
					CAT		
Parabacteroides distasonis		79.44	0.44	IND 1	76		
Bacteroides thetaiotaomicron		9.78	0.12		89		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Prevotella oralis	6.34	0.12		1	
55515332	**Acceptable identification**					**55515332**
	Parabacteroides distasonis	91.84	0.39	IND 1		
	Bacteroides thetaiotaomicron	8.16	0.05			
55515720	**Good identification**					**55515720**
					CAT	
	Bacteroides vulgatus	98.75	0.42	IND 1	3	
	Bacteroides thetaiotaomicron	1.25	0.12	SUC 96	89	
55515722	**Low discrimination**					**55515722**
	Bacteroides thetaiotaomicron	86.48	0.46	SUC 98		
	Bacteroides eggerthii	6.77	0.21	TRE 93		
	Parabacteroides distasonis	4.35	0.40			
55515732	**Acceptable identification**					**55515732**
	Bacteroides thetaiotaomicron	93.21	0.39	SUC 98		
	Parabacteroides distasonis	6.49	0.35	TRE 93		
	Bacteroides ovatus	0.30	-.03			
55521100	**Good identification**					**55521100**
					COC	
	Eubacterium saburreum	97.27	0.40	SUC 99	1	
	Blautia hansenii	2.73	-.03		99	
55521110	**Good identification**					**55521110**
	Eubacterium saburreum	100.0	0.38	SUC 99		
55525100	**Good identification**					**55525100**
	Eubacterium saburreum	100.0	0.35	SUC 99		
55531000	**Low discrimination**					**55531000**
	Eubacterium saburreum	50.77	0.42	SUC 99		
	Prevotella intermedia	47.71	0.32			
	Fusobacterium mortiferum	1.52	0.17			
55531010	**Good identification**					**55531010**
	Eubacterium saburreum	100.0	0.40	SUC 99		
55531100	**Good identification**					**55531100**
					GRA	
	Eubacterium saburreum	96.59	0.48	SUC 99	85	
	Fusobacterium mortiferum	2.89	0.24		1	
	Clostridium perfringens	0.53	0.00		99	
55531102	**Good identification**					**55531102**
	Eubacterium saburreum	100.0	0.35	SUC 99		
55531110	**Good identification**					**55531110**
	Eubacterium saburreum	100.0	0.46	SUC 99		
55531322	**Low discrimination**					**55531322**
					CAT	
	Parabacteroides distasonis	82.24	0.39	IND 1	76	
	Bacteroides thetaiotaomicron	11.07	0.08	XYL 90	89	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		6.69	0.07		1		
55531722	**Good identification**						**55531722**
Bacteroides thetaiotaomicron		95.60	0.42	SUC 98			
Parabacteroides distasonis		4.40	0.35	XYL 96			
55535000	**Good identification**						**55535000**
Eubacterium saburreum		100.0	0.37	SUC 99			
55535010	**Good identification**						**55535010**
Eubacterium saburreum		100.0	0.36	SUC 99			
55535100	**Good identification**						**55535100**
					COC		
Eubacterium saburreum		98.97	0.43	SUC 99	1		
Atopobium parvulum		1.04	0.00		99		
55535110	**Good identification**						**55535110**
Eubacterium saburreum		100.0	0.42	SUC 99			
55535122	**Low discrimination**						**55535122**
Parabacteroides distasonis		88.21	0.47	IND 1			
Bacteroides thetaiotaomicron		7.15	0.12				
Bacteroides fragilis		2.16	-.03				
55535132	**Acceptable identification**						**55535132**
					CAT		
Parabacteroides distasonis		91.79	0.42	IND 1	76		
Bacteroides thetaiotaomicron		5.37	0.05		89		
Eubacterium saburreum		2.83	0.16		1		
55535322	**Low discrimination**						**55535322**
Parabacteroides distasonis		73.59	0.55	IND 1			
Bacteroides thetaiotaomicron		26.42	0.31				
55535332	**Low discrimination**						**55535332**
Parabacteroides distasonis		79.41	0.50	IND 1			
Bacteroides thetaiotaomicron		20.59	0.24				
55535522	**Acceptable identification**						**55535522**
					GRA	CAT	
Bacteroides thetaiotaomicron		92.46	0.46	SUC 98	1	89	
Parabacteroides distasonis		7.06	0.43	RHA 93	1	76	
Bifidobacterium longum subsp. longum		0.39	0.17		99	1	
55535532	**Low discrimination**						**55535532**
					GRA	CAT	
Bacteroides thetaiotaomicron		88.96	0.39	SUC 98	1	89	
Parabacteroides distasonis		9.41	0.39	RHA 93	1	76	
Bifidobacterium longum subsp. longum		1.63	0.21		99	1	
55535622	**Good identification**						**55535622**
Bacteroides thetaiotaomicron		99.15	0.37	SUC 98			
Parabacteroides distasonis		0.85	0.17	LAC 98			
55535722	**Very good identification**						**55535722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		98.17	0.66	SUC 98		
Parabacteroides distasonis		1.69	0.51			
Bacteroides ovatus		0.14	0.17			
55535732	**Very good identification**					**55535732**
Bacteroides thetaiotaomicron		97.41	0.58	SUC 98		
Parabacteroides distasonis		2.33	0.47			
Bacteroides ovatus		0.24	0.14			
55541722	**Good identification**					**55541722**
Bacteroides eggerthii		99.96	0.47	RAF 1		
Bacteroides thetaiotaomicron		0.04	-.06	XYL 93		
55545722	**Very good identification**					**55545722**
					CAT	
Bacteroides eggerthii		99.87	0.66	RAF 1	93	
Bacteroides thetaiotaomicron		0.07	0.18		89	
Bacteroides uniformis		0.06	0.14		5	
55551122	**Good identification**					**55551122**
					CAT	
Prevotella oralis		95.50	0.43	IND 1	1	
Capnocytophaga ochracea		3.16	0.18	SUC 92	90	
Prevotella melaninogenica		0.45	0.00		1	
55551322	**Good identification**					**55551322**
Prevotella oralis		96.23	0.37	IND 1		
Parabacteroides distasonis		2.52	0.23	SUC 92		
Bacteroides uniformis		0.79	-.05			
55555522	**Low discrimination**					**55555522**
					CAT	
Bacteroides uniformis		87.45	0.45	SUC 99	5	
Bacteroides thetaiotaomicron		10.95	0.33	SAL 92	89	
Bacteroides ovatus		0.67	0.12		68	
55555722	**Low discrimination**					**55555722**
					CAT	
Bacteroides thetaiotaomicron		48.12	0.52	SUC 98	89	
Bacteroides uniformis		39.95	0.48	TRE 93	5	
Bacteroides eggerthii		7.09	0.32		93	
55555732	**Low discrimination**					**55555732**
					CAT	
Bacteroides thetaiotaomicron		83.78	0.45	SUC 98	89	
Bacteroides ovatus		12.62	0.31	TRE 93	68	
Bacteroides uniformis		1.81	0.14		5	
55571722	**Good identification**					**55571722**
Bacteroides thetaiotaomicron		97.29	0.47	SUC 98		
Bacteroides ovatus		1.58	0.17	XYL 96		
Parabacteroides distasonis		1.13	0.30			
55571732	**Good identification**					**55571732**
Bacteroides thetaiotaomicron		95.79	0.40	SUC 98		
Bacteroides ovatus		2.66	0.14	XYL 96		
Parabacteroides distasonis		1.55	0.26			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
55574722	**Acceptable identification**					**55574722**
Bacteroides thetaiotaomicron		93.35	0.37	SUC 98		
Bacteroides ovatus		6.24	0.17	GAL 99		
Parabacteroides distasonis		0.41	0.13			
55575122	**Low discrimination**					**55575122**
					CAT	
Parabacteroides distasonis		68.81	0.42	IND 1	76	
Bacteroides thetaiotaomicron		22.01	0.18		89	
Prevotella oralis		3.54	0.07		1	
55575132	**Low discrimination**					**55575132**
Parabacteroides distasonis		77.66	0.38	IND 1		
Bacteroides thetaiotaomicron		17.94	0.11			
Bacteroides ovatus		4.40	0.00			
55575322	**Low discrimination**					**55575322**
Bacteroides thetaiotaomicron		52.05	0.37	SUC 98		
Parabacteroides distasonis		36.74	0.50	ARA 99		
Bacteroides ovatus		10.66	0.25			
55575522	**Acceptable identification**					**55575522**
Bacteroides thetaiotaomicron		93.27	0.52	SUC 98		
Bacteroides ovatus		4.36	0.29	RHA 93		
Parabacteroides distasonis		1.81	0.39			
55575532	**Acceptable identification**					**55575532**
Bacteroides thetaiotaomicron		90.34	0.45	SUC 98		
Bacteroides ovatus		7.24	0.26	RHA 93		
Parabacteroides distasonis		2.42	0.34			
55575622	**Good identification**					**55575622**
Bacteroides thetaiotaomicron		96.59	0.42	SUC 98		
Bacteroides ovatus		3.20	0.17	LAC 98		
Parabacteroides distasonis		0.21	0.13			
55575632	**Acceptable identification**					**55575632**
Bacteroides thetaiotaomicron		94.06	0.35	SUC 98		
Bacteroides ovatus		5.34	0.14	LAC 98		
Clostridium clostridioforme		0.60	0.08			
55575702	**Acceptable identification**					**55575702**
Bacteroides thetaiotaomicron		93.03	0.37	SUC 98		
Bacteroides ovatus		6.22	0.17	NAG 99		
Parabacteroides distasonis		0.41	0.13			
55575720	**Acceptable identification**					**55575720**
					CAT	
Bacteroides thetaiotaomicron		92.82	0.37	SUC 98	89	
Bacteroides ovatus		6.21	0.17	BGL 99	68	
Bacteroides vulgatus		0.57	-.03		3	
55575722	**Good identification**					**55575722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		93.28	0.71	SUC 98			
Bacteroides ovatus		6.24	0.51				
Parabacteroides distasonis		0.41	0.47				
55575723	**Acceptable identification**						**55575723**
Bacteroides thetaiotaomicron		93.35	0.37	SUC 98			
Bacteroides ovatus		6.24	0.17	URE 1			
Parabacteroides distasonis		0.41	0.13				
55575732	**Low discrimination**						**55575732**
Bacteroides thetaiotaomicron		89.20	0.64	SUC 98			
Bacteroides ovatus		10.23	0.48				
Parabacteroides distasonis		0.54	0.42				
55575762	**Low discrimination**						**55575762**
Bacteroides thetaiotaomicron		78.11	0.37	SUC 98			
Bacteroides ovatus		21.55	0.27	SOR 1			
Parabacteroides distasonis		0.34	0.13				
55577722	**Low discrimination**						**55577722**
Bacteroides ovatus		68.72	0.43	SUC 99			
Bacteroides thetaiotaomicron		31.14	0.37	SAL 88			
Parabacteroides distasonis		0.14	0.13				
55577732	**Low discrimination**						**55577732**
Bacteroides ovatus		79.09	0.40	SUC 99			
Bacteroides thetaiotaomicron		20.91	0.30	SAL 88			
55600000	**Good identification**						**55600000**
Clostridium bifermentans		100.0	0.43	SAL 10			
55610000	**Low discrimination**						**55610000**
					GRA	SPO	
Fusobacterium mortiferum		74.58	0.44	IND 1	1	1	
Clostridium bifermentans		23.38	0.21		99	99	
Fusobacterium varium		2.04	-.03		1	1	
55610100	**Good identification**						**55610100**
Fusobacterium mortiferum		100.0	0.50	IND 1			
55611000	**Good identification**						**55611000**
Fusobacterium mortiferum		100.0	0.47	IND 1			
55611100	**Good identification**						**55611100**
Fusobacterium mortiferum		100.0	0.53	IND 1			
55611102	**Good identification**						**55611102**
					GRA		
Fusobacterium mortiferum		96.99	0.37	IND 1	1		
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99		
55630100	**Good identification**						**55630100**
Fusobacterium mortiferum		100.0	0.41	IND 1			
55631000	**Good identification**						**55631000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.38	IND 1		
55631100	**Good identification**					**55631100**
Fusobacterium mortiferum		100.0	0.44	IND 1		
55635122	**Good identification**					**55635122**
Parabacteroides distasonis		100.0	0.38	IND 1		
55635322	**Good identification**					**55635322**
Parabacteroides distasonis		98.69	0.46	IND 1		
Bacteroides thetaiotaomicron		1.31	-.02			
55635332	**Good identification**					**55635332**
Parabacteroides distasonis		100.0	0.42	IND 1		
55650000	**Good identification**					**55650000**
Fusobacterium mortiferum		100.0	0.36	IND 1		
55650100	**Good identification**					**55650100**
Fusobacterium mortiferum		100.0	0.42	IND 1		
55651000	**Good identification**					**55651000**
Fusobacterium mortiferum		100.0	0.39	IND 1		
55651100	**Good identification**					**55651100**
Fusobacterium mortiferum		100.0	0.45	IND 1		
55655522	**Good identification**					**55655522**
					CAT	
Bacteroides uniformis		99.64	0.45	SUC 99	5	
Parabacteroides distasonis		0.22	0.18	MLT 92	76	
Bacteroides thetaiotaomicron		0.14	-.01		89	
55655722	**Good identification**					**55655722**
					CAT	
Bacteroides uniformis		97.46	0.48	SUC 99	5	
Bacteroides thetaiotaomicron		1.32	0.19	MLT 92	89	
Bacteroides ovatus		0.76	0.15		68	
55671100	**Good identification**					**55671100**
Fusobacterium mortiferum		100.0	0.36	IND 1		
55675322	**Low discrimination**					**55675322**
Parabacteroides distasonis		87.13	0.42	IND 1		
Bacteroides ovatus		6.13	0.06			
Bacteroides thetaiotaomicron		4.55	0.04			
55675332	**Low discrimination**					**55675332**
Parabacteroides distasonis		88.91	0.37	IND 1		
Bacteroides ovatus		7.74	0.03			
Bacteroides thetaiotaomicron		3.35	-.04			
55675722	**Low discrimination**					**55675722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		61.95	0.38	SUC 98		
Bacteroides ovatus		27.24	0.32	MLT 94		
Parabacteroides distasonis		7.34	0.38			
55700000	**Good identification**					**55700000**
Clostridium bifermentans		100.0	0.50	SAL 10		
55711100	**Low discrimination**					**55711100**
					GRA	
Fusobacterium mortiferum		74.01	0.37	IND 1	1	
Bifidobacterium longum subsp. infantis		20.21	0.10	MLT 11	99	
Eubacterium saburreum		5.78	0.17		85	
55711122	**Low discrimination**					**55711122**
					GRA	
Prevotella oralis		59.62	0.38	IND 1	1	
Bifidobacterium longum subsp. infantis		39.03	0.32	SUC 92	99	
Parabacteroides distasonis		0.89	0.20		1	
55715322	**Low discrimination**					**55715322**
Parabacteroides distasonis		72.78	0.44	IND 1		
Bacteroides uniformis		14.59	0.14			
Prevotella oralis		9.53	0.16			
55715332	**Good identification**					**55715332**
Parabacteroides distasonis		100.0	0.40	IND 1		
55731322	**Low discrimination**					**55731322**
					CAT	
Parabacteroides distasonis		86.55	0.39	IND 1	76	
Prevotella oralis		11.56	0.12	XYL 90	1	
Bacteroides thetaiotaomicron		1.90	-.05		89	
55735122	**Good identification**					**55735122**
					CAT	
Parabacteroides distasonis		96.63	0.47	IND 1	76	
Bacteroides thetaiotaomicron		1.28	-.01		89	
Prevotella oralis		1.24	0.02		1	
55735132	**Good identification**					**55735132**
Parabacteroides distasonis		100.0	0.43	IND 1		
55735322	**Acceptable identification**					**55735322**
Parabacteroides distasonis		93.53	0.56	IND 1		
Bacteroides thetaiotaomicron		5.47	0.18			
Bacteroides ovatus		1.00	0.06			
55735332	**Acceptable identification**					**55735332**
Parabacteroides distasonis		94.75	0.51	IND 1		
Bacteroides thetaiotaomicron		4.00	0.11			
Bacteroides ovatus		1.25	0.03			
55735722	**Low discrimination**					**55735722**
Bacteroides thetaiotaomicron		85.66	0.53	SUC 98		
Parabacteroides distasonis		9.07	0.52			
Bacteroides ovatus		5.11	0.32			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55735732	**Low discrimination**						**55735732**
Bacteroides thetaiotaomicron		79.35	0.46	SUC 98			
Parabacteroides distasonis		11.63	0.47				
Bacteroides ovatus		8.11	0.29				
55751122	**Low discrimination**						**55751122**
					GRA	SPO	
Prevotella oralis		88.77	0.48	IND 1	1	1	
Clostridium paraputrificum		6.65	0.22	SUC 92	99	99	
Bifidobacterium breve		2.30	0.32		99	1	
55751322	**Acceptable identification**						**55751322**
Prevotella oralis		93.27	0.41	IND 1			
Bacteroides uniformis		4.98	0.14	SUC 92			
Parabacteroides distasonis		1.49	0.23				
55755122	**Acceptable identification**						**55755122**
Bacteroides uniformis		91.61	0.45	SUC 99			
Prevotella oralis		6.90	0.32	ARA 92			
Parabacteroides distasonis		1.14	0.32				
55755322	**Acceptable identification**						**55755322**
Bacteroides uniformis		92.88	0.48	SUC 99			
Parabacteroides distasonis		2.52	0.40	ARA 92			
Bacteroides ovatus		2.23	0.23				
55755522	**Very good identification**						**55755522**
					CAT		
Bacteroides uniformis		99.24	0.63	SUC 99	5		
Bacteroides ovatus		0.49	0.27		68		
Bacteroides thetaiotaomicron		0.19	0.20		89		
55755622	**Good identification**						**55755622**
					CAT		
Bacteroides uniformis		95.07	0.37	SUC 99	5		
Bacteroides ovatus		3.18	0.15	LAC 98	68		
Bacteroides thetaiotaomicron		1.76	0.10		89		
55755722	**Good identification**						**55755722**
					CAT		
Bacteroides uniformis		91.94	0.66	SUC 99	5		
Bacteroides ovatus		6.21	0.49		68		
Bacteroides thetaiotaomicron		1.70	0.39		89		
55755732	**Low discrimination**						**55755732**
					CAT		
Bacteroides ovatus		71.08	0.46	SUC 99	68		
Bacteroides uniformis		15.96	0.32	TRE 91	5		
Bacteroides thetaiotaomicron		11.36	0.32		89		
55757722	**Low discrimination**						**55757722**
					CAT		
Bacteroides ovatus		68.62	0.41	SUC 99	68		
Bacteroides uniformis		30.81	0.32	TRE 91	5		
Bacteroides thetaiotaomicron		0.57	0.05		89		
55757732	**Good identification**						**55757732**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		98.85	0.38	SUC 99	68	
Bacteroides uniformis		0.67	-.03	TRE 91	5	
Bacteroides thetaiotaomicron		0.48	-.02		89	
55775122	**Low discrimination**					**55775122**
Parabacteroides distasonis		60.59	0.43	IND 1		
Bacteroides ovatus		18.78	0.18			
Bacteroides uniformis		10.75	0.11			
55775132	**Low discrimination**					**55775132**
Parabacteroides distasonis		70.35	0.38	IND 1		
Bacteroides ovatus		27.00	0.15			
Bacteroides thetaiotaomicron		2.65	-.02			
55775322	**Low discrimination**					**55775322**
Bacteroides ovatus		59.71	0.40	SUC 99		
Parabacteroides distasonis		30.42	0.51	ARA 97		
Bacteroides thetaiotaomicron		7.02	0.24			
55775332	**Low discrimination**					**55775332**
Bacteroides ovatus		67.21	0.37	SUC 99		
Parabacteroides distasonis		27.65	0.46	ARA 97		
Bacteroides thetaiotaomicron		4.61	0.17			
55775500	**Good identification**					**55775500**
					SPO	
Clostridium butyricum		99.63	0.40	IND 3	99	
Actinomyces israelii		0.37	0.20	SUC 99	1	
55775522	**Low discrimination**					**55775522**
					CAT	
Bacteroides ovatus		55.41	0.44	SUC 99	68	
Bacteroides thetaiotaomicron		28.52	0.39	RHA 95	89	
Bacteroides uniformis		11.29	0.29		5	
55775532	**Low discrimination**					**55775532**
Bacteroides ovatus		73.00	0.41	SUC 99		
Bacteroides thetaiotaomicron		21.92	0.32	RHA 95		
Parabacteroides distasonis		3.61	0.35			
55775722	**Low discrimination**					**55775722**
					CAT	
Bacteroides ovatus		72.21	0.66	SUC 99	68	
Bacteroides thetaiotaomicron		25.99	0.58		89	
Bacteroides uniformis		1.07	0.32		5	
55775732	**Low discrimination**					**55775732**
Bacteroides ovatus		81.85	0.63	SUC 99		
Bacteroides thetaiotaomicron		17.18	0.51			
Parabacteroides distasonis		0.64	0.43			
55775762	**Acceptable identification**					**55775762**
					CAT	
Bacteroides ovatus		91.48	0.42	SUC 99	68	
Bacteroides thetaiotaomicron		7.98	0.24	SOR 4	89	
Bacteroides uniformis		0.33	-.03		5	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
55775772	**Good identification**						**55775772**
Bacteroides ovatus		95.16	0.39	SUC 99			
Bacteroides thetaiotaomicron		4.84	0.17	SOR 4			
55777102	**Low discrimination**						**55777102**
					SPO		
Actinomyces israelii		83.32	0.37	IND 1	1		
Clostridium butyricum		10.28	0.00	SUC 98	99		
Bifidobacterium adolescentis		6.40	0.12		1		
55777522	**Good identification**						**55777522**
					GRA	CAT	
Bacteroides ovatus		95.72	0.36	SUC 99	1	68	
Bacteroides thetaiotaomicron		1.49	0.05	RHA 95	1	89	
Bifidobacterium dentium		1.03	-.03		99	1	
55777722	**Very good identification**						**55777722**
					CAT		
Bacteroides ovatus		98.85	0.58	SUC 99	68		
Bacteroides thetaiotaomicron		1.08	0.24		89		
Bacteroides uniformis		0.04	-.03		5		
55777732	**Good identification**						**55777732**
					GRA	CAT	
Bacteroides ovatus		99.28	0.55	SUC 99	1	68	
Bacteroides thetaiotaomicron		0.63	0.17		1	89	
Bifidobacterium dentium		0.08	-.03		99	1	
56000000	**Low discrimination**						**56000000**
					SPO		
Clostridium bifermentans		89.93	0.41	GLU 92	99		
Atopobium minutum		2.97	0.24	SUC 1	1		
Clostridium botulinum biovar A		1.41	0.11		99		
56001100	**Good identification**						**56001100**
Eubacterium saburreum		100.0	0.37	GLU 99			
56001110	**Good identification**						**56001110**
Eubacterium saburreum		100.0	0.36	GLU 99			
56011000	**Low discrimination**						**56011000**
					GRA		
Eubacterium saburreum		86.60	0.39	GLU 99	85		
Atopobium minutum		6.17	0.16		99		
Prevotella intermedia		4.26	0.07		1		
56011010	**Good identification**						**56011010**
Eubacterium saburreum		100.0	0.37	GLU 99			
56011100	**Good identification**						**56011100**
Eubacterium saburreum		96.69	0.45	GLU 99			
Fusobacterium mortiferum		3.31	0.22				
56011110	**Good identification**						**56011110**
Eubacterium saburreum		100.0	0.44	GLU 99			
56015100	**Good identification**						**56015100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.41	GLU 99			
56015110	**Good identification**						**56015110**
Eubacterium saburreum		100.0	0.39	GLU 99			
56031000	**Good identification**						**56031000**
Eubacterium saburreum		100.0	0.36	GLU 99			
56031100	**Good identification**						**56031100**
Eubacterium saburreum		98.49	0.42	GLU 99			
Fusobacterium mortiferum		1.51	0.13				
56031110	**Good identification**						**56031110**
Eubacterium saburreum		100.0	0.41	GLU 99			
56035100	**Good identification**						**56035100**
Eubacterium saburreum		100.0	0.38	GLU 99			
56035110	**Good identification**						**56035110**
Eubacterium saburreum		100.0	0.36	GLU 99			
56100000	**Good identification**						**56100000**
					GRA	SPO	
Clostridium bifermentans		99.16	0.48	GLU 92	99	99	
Clostridium botulinum biovar A		0.54	0.10	SUC 1	99	99	
Prevotella intermedia		0.30	0.07		1	1	
56101000	**Low discrimination**						**56101000**
					SPO	GRA	
Eubacterium saburreum		86.15	0.41	GLU 99	1	85	
Clostridium bifermentans		10.67	0.14		99	99	
Prevotella intermedia		3.18	0.07		1	1	
56101010	**Good identification**						**56101010**
Eubacterium saburreum		100.0	0.40	GLU 99			
56101100	**Good identification**						**56101100**
Eubacterium saburreum		100.0	0.47	GLU 99			
56101110	**Good identification**						**56101110**
Eubacterium saburreum		100.0	0.46	GLU 99			
56105000	**Good identification**						**56105000**
Eubacterium saburreum		100.0	0.37	GLU 99			
56105010	**Good identification**						**56105010**
Eubacterium saburreum		100.0	0.35	GLU 99			
56105100	**Good identification**						**56105100**
Eubacterium saburreum		100.0	0.43	GLU 99			
56105110	**Good identification**						**56105110**
Eubacterium saburreum		100.0	0.41	GLU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
56110000	**Low discrimination**						**56110000**
					SPO	GRA	
Prevotella intermedia		84.40	0.41	ESL 1	1	1	
Clostridium bifermentans		14.90	0.26	GLU 90	99	99	
Eubacterium saburreum		0.70	0.15		1	85	
56111000	**Low discrimination**						**56111000**
Prevotella intermedia		54.93	0.41	ESL 1			
Eubacterium saburreum		45.08	0.49	GLU 90			
56111002	**Acceptable identification**						**56111002**
Eubacterium saburreum		93.48	0.36	GLU 99			
Prevotella intermedia		6.52	0.07				
56111010	**Good identification**						**56111010**
Eubacterium saburreum		98.52	0.48	GLU 99			
Prevotella intermedia		1.48	0.07				
56111020	**Low discrimination**						**56111020**
Eubacterium saburreum		87.69	0.36	GLU 99			
Prevotella intermedia		6.12	0.07				
Prevotella melaninogenica		3.29	-.04				
56111100	**Good identification**						**56111100**
Eubacterium saburreum		99.20	0.56	GLU 99			
Prevotella intermedia		0.52	0.07				
Prevotella melaninogenica		0.28	-.04				
56111102	**Good identification**						**56111102**
Eubacterium saburreum		100.0	0.43	GLU 99			
56111110	**Good identification**						**56111110**
Eubacterium saburreum		100.0	0.54	GLU 99			
56111112	**Good identification**						**56111112**
Eubacterium saburreum		100.0	0.41	GLU 99			
56111130	**Good identification**						**56111130**
Eubacterium saburreum		98.07	0.41	GLU 99			
Prevotella melaninogenica		1.93	-.04				
56115000	**Good identification**						**56115000**
Eubacterium saburreum		97.77	0.45	GLU 99			
Prevotella intermedia		2.24	0.07				
56115010	**Good identification**						**56115010**
Eubacterium saburreum		100.0	0.43	GLU 99			
56115100	**Good identification**						**56115100**
Eubacterium saburreum		100.0	0.51	GLU 99			
56115102	**Good identification**						**56115102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	GLU 99		
56115110	**Good identification**					**56115110**
Eubacterium saburreum		100.0	0.49	GLU 99		
56115112	**Good identification**					**56115112**
Eubacterium saburreum		100.0	0.37	GLU 99		
56115120	**Good identification**					**56115120**
Eubacterium saburreum		97.10	0.38	GLU 99		
Prevotella melaninogenica		2.90	-.04			
56115130	**Good identification**					**56115130**
Eubacterium saburreum		100.0	0.37	GLU 99		
56121000	**Good identification**					**56121000**
Eubacterium saburreum		100.0	0.38	GLU 99		
56121010	**Good identification**					**56121010**
Eubacterium saburreum		100.0	0.37	GLU 99		
56121100	**Good identification**					**56121100**
Eubacterium saburreum		100.0	0.44	GLU 99		
56121110	**Good identification**					**56121110**
Eubacterium saburreum		100.0	0.43	GLU 99		
56125100	**Good identification**					**56125100**
Eubacterium saburreum		100.0	0.40	GLU 99		
56125110	**Good identification**					**56125110**
Eubacterium saburreum		100.0	0.38	GLU 99		
56131000	**Good identification**					**56131000**
Eubacterium saburreum		98.19	0.46	GLU 99		
Prevotella intermedia		1.81	0.07			
56131010	**Good identification**					**56131010**
Eubacterium saburreum		100.0	0.45	GLU 99		
56131100	**Good identification**					**56131100**
					COC	
Eubacterium saburreum		96.23	0.53	GLU 99	1	
Atopobium parvulum		3.77	0.19		99	
56131102	**Good identification**					**56131102**
					COC	
Eubacterium saburreum		96.23	0.40	GLU 99	1	
Atopobium parvulum		3.77	0.06		99	
56131110	**Good identification**					**56131110**
Eubacterium saburreum		100.0	0.51	GLU 99		
56131112	**Good identification**					**56131112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.38	GLU 99		
56131120	**Low discrimination**					**56131120**
					COC	
Eubacterium saburreum		73.90	0.40	GLU 99	1	
Clostridium perfringens		21.42	0.21		1	
Atopobium parvulum		2.90	0.06		99	
56131130	**Good identification**					**56131130**
Eubacterium saburreum		100.0	0.38	GLU 99		
56135000	**Good identification**					**56135000**
Eubacterium saburreum		100.0	0.42	GLU 99		
56135010	**Good identification**					**56135010**
Eubacterium saburreum		100.0	0.40	GLU 99		
56135100	**Low discrimination**					**56135100**
					COC	
Eubacterium saburreum		88.10	0.48	GLU 99	1	
Atopobium parvulum		11.90	0.23		99	
56135102	**Low discrimination**					**56135102**
					COC	
Eubacterium saburreum		88.10	0.35	GLU 99	1	
Atopobium parvulum		11.90	0.10		99	
56135110	**Good identification**					**56135110**
Eubacterium saburreum		100.0	0.46	GLU 99		
56135120	**Low discrimination**					**56135120**
					COC	
Eubacterium saburreum		88.10	0.35	GLU 99	1	
Atopobium parvulum		11.90	0.10		99	
56175722	**Good identification**					**56175722**
Bacteroides thetaiotaomicron		95.58	0.40	GLU 97		
Bacteroides ovatus		4.22	0.17	RAF 99		
Clostridium clostridioforme		0.20	0.05			
56355522	**Good identification**					**56355522**
Bacteroides uniformis		100.0	0.38	GLU 99		
56355722	**Good identification**					**56355722**
					CAT	
Bacteroides uniformis		96.84	0.40	GLU 99	5	
Bacteroides ovatus		2.14	0.15	RAF 97	68	
Bacteroides thetaiotaomicron		0.89	0.08		89	
56411100	**Low discrimination**					**56411100**
Eubacterium saburreum		89.01	0.41	GLU 99		
Fusobacterium mortiferum		10.99	0.27			
56411110	**Good identification**					**56411110**
Eubacterium saburreum		100.0	0.39	GLU 99		
56415100	**Good identification**					**56415100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.36	GLU 99		
56431100	**Acceptable identification**					**56431100**
Eubacterium saburreum		94.76	0.38	GLU 99		
Fusobacterium mortiferum		5.24	0.18			
56431110	**Good identification**					**56431110**
Eubacterium saburreum		100.0	0.36	GLU 99		
56435722	**Good identification**					**56435722**
Bacteroides thetaiotaomicron		99.80	0.48	GLU 97		
Parabacteroides distasonis		0.20	0.18	MLT 94		
56435732	**Good identification**					**56435732**
Bacteroides thetaiotaomicron		99.72	0.41	GLU 97		
Parabacteroides distasonis		0.28	0.13	MLT 94		
56475722	**Good identification**					**56475722**
					CAT	
Bacteroides thetaiotaomicron		99.20	0.54	GLU 97	89	
Bacteroides ovatus		0.69	0.17	MLT 94	68	
Bacteroides uniformis		0.06	-.05		5	
56475732	**Good identification**					**56475732**
Bacteroides thetaiotaomicron		98.82	0.47	GLU 97		
Bacteroides ovatus		1.18	0.14	MLT 94		
56501000	**Acceptable identification**					**56501000**
Eubacterium saburreum		90.67	0.37	GLU 99		
Prevotella intermedia		9.33	0.10			
56501010	**Good identification**					**56501010**
Eubacterium saburreum		100.0	0.35	GLU 99		
56501100	**Good identification**					**56501100**
Eubacterium saburreum		98.44	0.43	GLU 99		
Prevotella melaninogenica		1.56	-.04			
56501110	**Good identification**					**56501110**
Eubacterium saburreum		100.0	0.41	GLU 99		
56505100	**Good identification**					**56505100**
Eubacterium saburreum		100.0	0.38	GLU 99		
56505110	**Good identification**					**56505110**
Eubacterium saburreum		100.0	0.37	GLU 99		
56510000	**Good identification**					**56510000**
Prevotella intermedia		99.70	0.44	ESL 1		
Eubacterium saburreum		0.30	0.11	GLU 90		
56511000	**Low discrimination**					**56511000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella intermedia		77.24	0.44	ESL 1		
Eubacterium saburreum		22.76	0.45	GLU 90		
56511010	**Good identification**					**56511010**
Eubacterium saburreum		95.98	0.43	GLU 99		
Prevotella intermedia		4.02	0.10			
56511100	**Acceptable identification**					**56511100**
Eubacterium saburreum		93.82	0.51	GLU 99		
Prevotella melaninogenica		4.45	0.12			
Prevotella intermedia		1.38	0.10			
56511102	**Low discrimination**					**56511102**
					CAT	
Eubacterium saburreum		81.10	0.38	GLU 99	1	
Capnocytophaga ochracea		13.09	0.12		90	
Prevotella melaninogenica		4.15	0.00		1	
56511110	**Good identification**					**56511110**
Eubacterium saburreum		100.0	0.49	GLU 99		
56511112	**Good identification**					**56511112**
Eubacterium saburreum		100.0	0.37	GLU 99		
56511120	**Good identification**					**56511120**
					CAT	
Prevotella melaninogenica		95.69	0.47	IND 1	1	
Eubacterium saburreum		3.60	0.38	GLU 90	1	
Capnocytophaga ochracea		0.58	0.12		90	
56511130	**Low discrimination**					**56511130**
Eubacterium saburreum		75.28	0.37	GLU 99		
Prevotella melaninogenica		24.72	0.12			
56515000	**Acceptable identification**					**56515000**
Eubacterium saburreum		94.01	0.40	GLU 99		
Prevotella intermedia		5.99	0.10			
56515010	**Good identification**					**56515010**
Eubacterium saburreum		100.0	0.39	GLU 99		
56515100	**Good identification**					**56515100**
Eubacterium saburreum		100.0	0.46	GLU 99		
56515110	**Good identification**					**56515110**
Eubacterium saburreum		100.0	0.45	GLU 99		
56515722	**Good identification**					**56515722**
					CAT	
Bacteroides thetaiotaomicron		98.46	0.49	GLU 97	89	
Bacteroides uniformis		1.16	0.14	TRE 93	5	
Parabacteroides distasonis		0.13	0.16		76	
56515732	**Good identification**					**56515732**
Bacteroides thetaiotaomicron		99.16	0.42	GLU 97		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		0.44	0.12	TRE 93		
Bacteroides ovatus		0.21	-.03			
56521100	**Good identification**					**56521100**
Eubacterium saburreum		100.0	0.40	GLU 99		
56521110	**Good identification**					**56521110**
Eubacterium saburreum		100.0	0.38	GLU 99		
56525100	**Good identification**					**56525100**
Eubacterium saburreum		100.0	0.35	GLU 99		
56531000	**Good identification**					**56531000**
Eubacterium saburreum		95.11	0.42	GLU 99		
Prevotella intermedia		4.89	0.10			
56531010	**Good identification**					**56531010**
Eubacterium saburreum		100.0	0.40	GLU 99		
56531100	**Good identification**					**56531100**
Eubacterium saburreum		100.0	0.48	GLU 99		
56531102	**Good identification**					**56531102**
Eubacterium saburreum		100.0	0.35	GLU 99		
56531110	**Good identification**					**56531110**
Eubacterium saburreum		100.0	0.46	GLU 99		
56531120	**Low discrimination**					**56531120**
					GRA	
Eubacterium saburreum		66.42	0.35	GLU 99	85	
Prevotella melaninogenica		26.77	0.12		1	
Clostridium perfringens		6.81	0.09		99	
56531722	**Good identification**					**56531722**
Bacteroides thetaiotaomicron		99.88	0.45	GLU 97		
Parabacteroides distasonis		0.12	0.11	XYL 96		
56531732	**Good identification**					**56531732**
Bacteroides thetaiotaomicron		99.42	0.38	GLU 97		
Clostridium clostridioforme		0.58	0.10	XYL 96		
56535000	**Good identification**					**56535000**
Eubacterium saburreum		100.0	0.37	GLU 99		
56535010	**Good identification**					**56535010**
Eubacterium saburreum		100.0	0.36	GLU 99		
56535100	**Good identification**					**56535100**
Eubacterium saburreum		100.0	0.43	GLU 99		
56535110	**Good identification**					**56535110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.42	GLU 99			
56535522	**Good identification**						**56535522**
					GRA	CAT	
Bacteroides thetaiotaomicron		99.46	0.49	GLU 97	1	89	
Bifidobacterium longum subsp. longum		0.28	0.17	RHA 93	99	1	
Parabacteroides distasonis		0.20	0.19		1	76	
56535532	**Good identification**						**56535532**
					GRA	CAT	
Bacteroides thetaiotaomicron		98.10	0.42	GLU 97	1	89	
Bifidobacterium longum subsp. longum		1.19	0.21	RHA 93	99	1	
Clostridium clostridioforme		0.44	0.12		50	33	
56535622	**Good identification**						**56535622**
Bacteroides thetaiotaomicron		99.69	0.40	GLU 97			
Clostridium clostridioforme		0.31	0.07	LAC 98			
56535722	**Very good identification**						**56535722**
Bacteroides thetaiotaomicron		99.84	0.69	GLU 97			
Bacteroides ovatus		0.10	0.17				
Parabacteroides distasonis		0.05	0.27				
56535732	**Very good identification**						**56535732**
Bacteroides thetaiotaomicron		99.53	0.62	GLU 97			
Clostridium clostridioforme		0.25	0.27				
Bacteroides ovatus		0.16	0.14				
56555522	**Low discrimination**						**56555522**
					CAT		
Bacteroides uniformis		83.44	0.45	GLU 99	5		
Bacteroides thetaiotaomicron		15.83	0.36	SAL 92	89		
Bacteroides ovatus		0.64	0.12		68		
56555722	**Low discrimination**						**56555722**
					CAT		
Bacteroides thetaiotaomicron		62.24	0.55	GLU 97	89		
Bacteroides uniformis		34.09	0.48	TRE 93	5		
Bacteroides ovatus		3.61	0.34		68		
56555732	**Low discrimination**						**56555732**
					CAT		
Bacteroides thetaiotaomicron		89.24	0.48	GLU 97	89		
Bacteroides ovatus		8.87	0.31	TRE 93	68		
Bacteroides uniformis		1.27	0.14		5		
56565722	**Good identification**						**56565722**
Bacteroides thetaiotaomicron		95.77	0.40	GLU 97			
Bacteroides ovatus		4.23	0.17	FRU 99			
56571722	**Good identification**						**56571722**
Bacteroides thetaiotaomicron		98.87	0.51	GLU 97			
Bacteroides ovatus		1.06	0.17	XYL 96			
Clostridium clostridioforme		0.08	0.08				
56571732	**Good identification**						**56571732**
Bacteroides thetaiotaomicron		97.32	0.44	GLU 97			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		1.79	0.14	XYL 96		
Clostridium clostridioforme		0.90	0.19			
56574722	**Acceptable identification**					**56574722**
Bacteroides thetaiotaomicron		94.78	0.40	GLU 97		
Bacteroides ovatus		4.18	0.17	GAL 99		
Clostridium clostridioforme		1.04	0.17			
56575322	**Low discrimination**					**56575322**
Bacteroides thetaiotaomicron		86.24	0.40	GLU 97		
Bacteroides ovatus		11.65	0.25	ARA 99		
Parabacteroides distasonis		1.62	0.27			
56575522	**Good identification**					**56575522**
					CAT	
Bacteroides thetaiotaomicron		96.53	0.55	GLU 97	89	
Bacteroides ovatus		2.98	0.29	RHA 93	68	
Bacteroides uniformis		0.39	0.11		5	
56575532	**Acceptable identification**					**56575532**
Bacteroides thetaiotaomicron		94.28	0.48	GLU 97		
Bacteroides ovatus		4.99	0.26	RHA 93		
Clostridium clostridioforme		0.66	0.21			
56575622	**Good identification**					**56575622**
Bacteroides thetaiotaomicron		97.40	0.45	GLU 97		
Bacteroides ovatus		2.13	0.17	LAC 98		
Clostridium clostridioforme		0.47	0.16			
56575632	**Acceptable identification**					**56575632**
Bacteroides thetaiotaomicron		91.40	0.38	GLU 97		
Clostridium clostridioforme		5.18	0.27	LAC 98		
Bacteroides ovatus		3.42	0.14			
56575702	**Acceptable identification**					**56575702**
Bacteroides thetaiotaomicron		92.86	0.40	GLU 97		
Bacteroides ovatus		4.10	0.17	NAG 99		
Clostridium clostridioforme		3.04	0.25			
56575720	**Good identification**					**56575720**
Bacteroides thetaiotaomicron		95.77	0.40	GLU 97		
Bacteroides ovatus		4.23	0.17	BGL 99		
56575722	**Very good identification**					**56575722**
					CAT	
Bacteroides thetaiotaomicron		95.70	0.74	GLU 97	89	
Bacteroides ovatus		4.22	0.51		68	
Bacteroides uniformis		0.04	0.14		5	
56575723	**Good identification**					**56575723**
Bacteroides thetaiotaomicron		95.77	0.40	GLU 97		
Bacteroides ovatus		4.23	0.17	URE 1		
56575732	**Good identification**					**56575732**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		92.62	0.67	GLU 97		
Bacteroides ovatus		7.01	0.48			
Clostridium clostridioforme		0.36	0.36			
56575762	**Low discrimination**					**56575762**
Bacteroides thetaiotaomicron		84.60	0.40	GLU 97		
Bacteroides ovatus		15.40	0.27	SOR 1		
56577722	**Low discrimination**					**56577722**
Bacteroides ovatus		59.28	0.43	GLU 99		
Bacteroides thetaiotaomicron		40.72	0.40	SAL 88		
56577732	**Low discrimination**					**56577732**
Bacteroides ovatus		71.40	0.40	GLU 99		
Bacteroides thetaiotaomicron		28.60	0.33	SAL 88		
56635722	**Good identification**					**56635722**
Bacteroides thetaiotaomicron		98.16	0.35	GLU 97		
Parabacteroides distasonis		1.22	0.19	MLT 94		
Bacteroides ovatus		0.61	-.03			
56655522	**Good identification**					**56655522**
					CAT	
Bacteroides uniformis		99.79	0.45	GLU 99	5	
Bacteroides thetaiotaomicron		0.21	0.02	MLT 92	89	
56655722	**Good identification**					**56655722**
					CAT	
Bacteroides uniformis		97.24	0.48	GLU 99	5	
Bacteroides thetaiotaomicron		2.00	0.22	MLT 92	89	
Bacteroides ovatus		0.76	0.15		68	
56675722	**Low discrimination**					**56675722**
					CAT	
Bacteroides thetaiotaomicron		75.05	0.41	GLU 97	89	
Bacteroides ovatus		21.78	0.32	MLT 94	68	
Bacteroides uniformis		2.78	0.14		5	
56715722	**Low discrimination**					**56715722**
					CAT	
Bacteroides thetaiotaomicron		54.29	0.36	GLU 97	89	
Bacteroides uniformis		41.60	0.32	TRE 93	5	
Bacteroides ovatus		2.81	0.15		68	
56735522	**Acceptable identification**					**56735522**
Bacteroides thetaiotaomicron		94.21	0.36	GLU 97		
Bacteroides ovatus		2.59	0.10	RHA 93		
Clostridium clostridioforme		1.49	0.16			
56735712	**Acceptable identification**					**56735712**
Clostridium clostridioforme		90.46	0.42	IND 1		
Bacteroides thetaiotaomicron		8.94	0.15	GLU 88		
Bacteroides ovatus		0.60	-.06			
56735722	**Good identification**					**56735722**
Bacteroides thetaiotaomicron		95.10	0.56	GLU 97		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides ovatus		3.74	0.32			
Clostridium clostridioforme		0.83	0.31			
56735732	**Low discrimination**					**56735732**
Bacteroides thetaiotaomicron		85.21	0.49	GLU 97		
Clostridium clostridioforme		8.71	0.42			
Bacteroides ovatus		5.75	0.29			
56755122	**Good identification**					**56755122**
Bacteroides uniformis		98.77	0.45	GLU 99		
Prevotella oralis		0.87	0.16	ARA 92		
Bacteroides ovatus		0.17	0.01			
56755322	**Good identification**					**56755322**
					CAT	
Bacteroides uniformis		96.94	0.48	GLU 99	5	
Bacteroides ovatus		2.33	0.23	ARA 92	68	
Bacteroides thetaiotaomicron		0.32	0.08		89	
56755522	**Very good identification**					**56755522**
					CAT	
Bacteroides uniformis		99.20	0.63	GLU 99	5	
Bacteroides ovatus		0.49	0.27		68	
Bacteroides thetaiotaomicron		0.29	0.23		89	
56755622	**Acceptable identification**					**56755622**
					CAT	
Bacteroides uniformis		93.32	0.37	GLU 99	5	
Bacteroides ovatus		3.12	0.15	LAC 98	68	
Bacteroides thetaiotaomicron		2.61	0.13		89	
56755712	**Low discrimination**					**56755712**
Clostridium clostridioforme		82.66	0.36	IND 1		
Bacteroides ovatus		11.82	0.12	GLU 88		
Bacteroides thetaiotaomicron		2.86	0.01	TRE 88		
56755722	**Good identification**					**56755722**
					CAT	
Bacteroides uniformis		91.22	0.66	GLU 99	5	
Bacteroides ovatus		6.16	0.49		68	
Bacteroides thetaiotaomicron		2.56	0.42		89	
56755732	**Low discrimination**					**56755732**
					CAT	
Bacteroides ovatus		65.05	0.46	GLU 99	68	
Bacteroides thetaiotaomicron		15.75	0.35	TRE 91	89	
Bacteroides uniformis		14.60	0.32		5	
56757722	**Low discrimination**					**56757722**
					CAT	
Bacteroides ovatus		68.42	0.41	GLU 99	68	
Bacteroides uniformis		30.72	0.32	TRE 91	5	
Bacteroides thetaiotaomicron		0.86	0.08		89	
56757732	**Good identification**					**56757732**
					CAT	
Bacteroides ovatus		98.61	0.38	GLU 99	68	
Bacteroides thetaiotaomicron		0.72	0.01	TRE 91	89	
Bacteroides uniformis		0.67	-.03		5	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
56771722	**Low discrimination**					**56771722**
Bacteroides thetaiotaomicron		67.40	0.38	GLU 97		
Bacteroides ovatus		29.96	0.32	XYL 96		
Clostridium clostridioforme		2.20	0.23			
56774712	**Good identification**					**56774712**
Clostridium clostridioforme		99.41	0.43	IND 1		
Bacteroides ovatus		0.59	-.06	GLU 88		
56774732	**Low discrimination**					**56774732**
Clostridium clostridioforme		56.19	0.43	IND 1		
Bacteroides ovatus		33.24	0.29	GLU 88		
Bacteroides thetaiotaomicron		10.58	0.20			
56775322	**Low discrimination**					**56775322**
					CAT	
Bacteroides ovatus		79.63	0.40	GLU 99	68	
Bacteroides thetaiotaomicron		14.18	0.27	ARA 97	89	
Bacteroides uniformis		3.31	0.14		5	
56775332	**Low discrimination**					**56775332**
Bacteroides ovatus		81.55	0.37	GLU 99		
Clostridium clostridioforme		8.62	0.31	ARA 97		
Bacteroides thetaiotaomicron		8.47	0.20			
56775500	**Good identification**					**56775500**
					SPO	
Clostridium butyricum		99.82	0.40	IND 3	99	
Actinomyces israelii		0.19	0.14	GLU 99	1	
56775512	**Low discrimination**					**56775512**
					SPO	
Clostridium clostridioforme		65.22	0.36	IND 1	66	
Bifidobacterium adolescentis		15.48	0.29	GLU 88	1	
Clostridium butyricum		10.59	0.11	RHA 88	99	
56775522	**Low discrimination**					**56775522**
					CAT	
Bacteroides ovatus		49.28	0.44	GLU 99	68	
Bacteroides thetaiotaomicron		38.44	0.42	RHA 95	89	
Bacteroides uniformis		10.04	0.29		5	
56775532	**Low discrimination**					**56775532**
Bacteroides ovatus		62.50	0.41	GLU 99		
Bacteroides thetaiotaomicron		28.44	0.35	RHA 95		
Clostridium clostridioforme		8.30	0.36			
56775612	**Good identification**					**56775612**
Clostridium clostridioforme		99.34	0.42	IND 1		
Bacteroides ovatus		0.66	-.06	GLU 88		
56775632	**Low discrimination**					**56775632**
Clostridium clostridioforme		47.97	0.42	IND 1		
Bacteroides ovatus		31.67	0.29	GLU 88		
Bacteroides thetaiotaomicron		20.36	0.25			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
56775712	**Low discrimination**						**56775712**
Clostridium clostridioforme		79.37	0.51	IND 1			
Bacteroides ovatus		15.65	0.29	GLU 88			
Bacteroides thetaiotaomicron		4.98	0.20				
56775722	**Low discrimination**						**56775722**
					CAT		
Bacteroides ovatus		63.77	0.66	GLU 99	68		
Bacteroides thetaiotaomicron		34.78	0.61		89		
Bacteroides uniformis		0.94	0.32		5		
56775732	**Low discrimination**						**56775732**
Bacteroides ovatus		73.00	0.63	GLU 99			
Bacteroides thetaiotaomicron		23.22	0.54				
Clostridium clostridioforme		3.74	0.51				
56775762	**Low discrimination**						**56775762**
					CAT		
Bacteroides ovatus		87.90	0.42	GLU 99	68		
Bacteroides thetaiotaomicron		11.62	0.27	SOR 4	89		
Bacteroides uniformis		0.32	-.03		5		
56775772	**Acceptable identification**						**56775772**
Bacteroides ovatus		91.78	0.39	GLU 99			
Bacteroides thetaiotaomicron		7.08	0.20	SOR 4			
Clostridium clostridioforme		1.14	0.17				
56777522	**Good identification**						**56777522**
					GRA	CAT	
Bacteroides ovatus		95.38	0.36	GLU 99	1	68	
Bacteroides thetaiotaomicron		2.25	0.08	RHA 95	1	89	
Bifidobacterium dentium		1.03	-.03		99	1	
56777722	**Very good identification**						**56777722**
					CAT		
Bacteroides ovatus		98.31	0.58	GLU 99	68		
Bacteroides thetaiotaomicron		1.63	0.27		89		
Bacteroides uniformis		0.04	-.03		5		
56777732	**Good identification**						**56777732**
Bacteroides ovatus		98.81	0.55	GLU 99			
Bacteroides thetaiotaomicron		0.95	0.20				
Clostridium clostridioforme		0.15	0.17				
57000000	**Very good identification**						**57000000**
					SPO		
Clostridium bifermentans		97.49	0.60	SUC 1	99		
Clostridium botulinum biovar A		1.19	0.27		99		
Atopobium minutum		0.48	0.28		1		
57000020	**Low discrimination**						**57000020**
					SPO		
Clostridium cadaveris		70.17	0.44	ESL 1	99		
Clostridium bifermentans		28.66	0.39	SUC 7	99		
Atopobium minutum		0.73	0.20		1		
57000040	**Acceptable identification**						**57000040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium bifermentans		91.93	0.36	SUC 1		
Clostridium botulinum biovar A		8.07	0.18	SOR 4		
57000110	**Good identification**					**57000110**
Eubacterium saburreum		100.0	0.36	GAL 99		
57001000	**Very good identification**					**57001000**
					SPO	
Eubacterium saburreum		96.77	0.65		1	
Clostridium bifermentans		2.39	0.25		99	
Atopobium minutum		0.39	0.20		1	
57001002	**Good identification**					**57001002**
Eubacterium saburreum		98.52	0.52			
Atopobium minutum		1.48	0.17			
57001010	**Very good identification**					**57001010**
Eubacterium saburreum		100.0	0.63			
57001012	**Good identification**					**57001012**
Eubacterium saburreum		100.0	0.51			
57001020	**Good identification**					**57001020**
					SPO	
Eubacterium saburreum		95.89	0.52		1	
Clostridium cadaveris		2.09	0.09		99	
Clostridium bifermentans		0.86	0.05		99	
57001022	**Good identification**					**57001022**
Eubacterium saburreum		97.24	0.39			
Atopobium minutum		2.77	0.09			
57001030	**Good identification**					**57001030**
Eubacterium saburreum		100.0	0.51			
57001032	**Good identification**					**57001032**
Eubacterium saburreum		100.0	0.38			
57001100	**Very good identification**					**57001100**
Eubacterium saburreum		99.86	0.71			
Fusobacterium mortiferum		0.10	0.22			
Odoribacter splanchnicus		0.03	-.07			
57001101	**Low discrimination**					**57001101**
					SPO	
Eubacterium saburreum		65.93	0.37	URE 1	1	
Clostridium chauvoei		34.07	0.22		99	
57001102	**Very good identification**					**57001102**
Eubacterium saburreum		100.0	0.58			
57001110	**Very good identification**					**57001110**
Eubacterium saburreum		100.0	0.70			
57001111	**Good identification**					**57001111**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.36	URE 1		
57001112	**Very good identification**					**57001112**
Eubacterium saburreum		100.0	0.57			
57001120	**Low discrimination**					**57001120**
Eubacterium saburreum		83.72	0.58			
Odoribacter splanchnicus		16.12	0.27			
Prevotella melaninogenica		0.16	-.04			
57001122	**Good identification**					**57001122**
Eubacterium saburreum		98.91	0.45			
Odoribacter splanchnicus		1.09	-.07			
57001130	**Very good identification**					**57001130**
Eubacterium saburreum		99.76	0.57			
Odoribacter splanchnicus		0.24	-.07			
57001132	**Good identification**					**57001132**
Eubacterium saburreum		100.0	0.44			
57001140	**Good identification**					**57001140**
Eubacterium saburreum		100.0	0.37	SOR 1		
57001150	**Good identification**					**57001150**
Eubacterium saburreum		100.0	0.36	SOR 1		
57001300	**Good identification**					**57001300**
Eubacterium saburreum		100.0	0.37	RHA 1		
57001310	**Good identification**					**57001310**
Eubacterium saburreum		100.0	0.36	RHA 1		
57001510	**Good identification**					**57001510**
Eubacterium saburreum		96.01	0.36	ARA 1		
Odoribacter splanchnicus		3.99	-.07			
57001520	**Very good identification**					**57001520**
Odoribacter splanchnicus		99.95	0.61	SUC 1		
Eubacterium saburreum		0.05	0.24			
57001560	**Good identification**					**57001560**
Odoribacter splanchnicus		100.0	0.45	SUC 1		
57003100	**Good identification**					**57003100**
Eubacterium saburreum		100.0	0.37	MAN 1		
57003110	**Good identification**					**57003110**
Eubacterium saburreum		100.0	0.36	MAN 1		
57005000	**Very good identification**					**57005000**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.60				
57005002	**Good identification**						**57005002**
Eubacterium saburreum		100.0	0.47				
57005010	**Very good identification**						**57005010**
Eubacterium saburreum		100.0	0.59				
57005012	**Good identification**						**57005012**
Eubacterium saburreum		100.0	0.46				
57005020	**Good identification**						**57005020**
Eubacterium saburreum		100.0	0.47				
57005030	**Good identification**						**57005030**
Eubacterium saburreum		100.0	0.46				
57005100	**Very good identification**						**57005100**
Eubacterium saburreum		100.0	0.67				
57005102	**Good identification**						**57005102**
Eubacterium saburreum		100.0	0.54				
57005110	**Very good identification**						**57005110**
Eubacterium saburreum		100.0	0.65				
57005112	**Good identification**						**57005112**
Eubacterium saburreum		100.0	0.52				
57005120	**Good identification**						**57005120**
Eubacterium saburreum		99.64	0.54				
Odoribacter splanchnicus		0.36	-.07				
57005122	**Good identification**						**57005122**
Eubacterium saburreum		100.0	0.41				
57005130	**Good identification**						**57005130**
Eubacterium saburreum		100.0	0.52				
57005132	**Good identification**						**57005132**
Eubacterium saburreum		100.0	0.39				
57010000	**Low discrimination**						**57010000**
					GRA	SPO	
Clostridium bifermentans		31.44	0.38	SUC 1	99	99	
Fusobacterium varium		30.21	0.32	FRU 5	1	1	
Fusobacterium mortiferum		16.76	0.47		1	1	
57010010	**Low discrimination**						**57010010**
					SPO		
Eubacterium saburreum		67.80	0.37	GAL 99	1		
Clostridium difficile		23.35	0.15		99		
Clostridium bifermentans		3.55	0.04		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57010020	**Acceptable identification**						**57010020**
					SPO		
Clostridium cadaveris		93.82	0.44	ESL 1	99		
Clostridium bifermentans		2.02	0.17	SUC 7	99		
Eubacterium saburreum		1.32	0.26		1		
57010100	**Low discrimination**						**57010100**
					SPO	GRA	
Fusobacterium mortiferum		68.57	0.53	IND 1	1	1	
Eubacterium saburreum		30.33	0.45		1	85	
Clostridium bifermentans		0.56	0.04		99	99	
57010102	**Low discrimination**						**57010102**
Fusobacterium mortiferum		58.73	0.37	IND 1			
Eubacterium saburreum		41.27	0.32	BGL 10			
57010110	**Good identification**						**57010110**
Eubacterium saburreum		97.29	0.44	GAL 99			
Fusobacterium mortiferum		2.72	0.19				
57011000	**Very good identification**						**57011000**
Eubacterium saburreum		95.86	0.73				
Fusobacterium mortiferum		3.28	0.50				
Prevotella intermedia		0.43	0.23				
57011001	**Good identification**						**57011001**
Eubacterium saburreum		96.69	0.39	URE 1			
Fusobacterium mortiferum		3.31	0.16				
57011002	**Very good identification**						**57011002**
					GRA		
Eubacterium saburreum		97.28	0.60		85		
Fusobacterium mortiferum		2.10	0.33		1		
Atopobium minutum		0.45	0.17		99		
57011010	**Very good identification**						**57011010**
Eubacterium saburreum		99.96	0.72				
Fusobacterium mortiferum		0.04	0.16				
57011011	**Good identification**						**57011011**
Eubacterium saburreum		100.0	0.37	URE 1			
57011012	**Very good identification**						**57011012**
Eubacterium saburreum		100.0	0.59				
57011020	**Very good identification**						**57011020**
					SPO	CAT	
Eubacterium saburreum		97.50	0.60		1	1	
Propionibacterium acnes		0.80	0.14		1	95	
Clostridium cadaveris		0.71	0.09		99	1	
57011022	**Good identification**						**57011022**
Eubacterium saburreum		99.13	0.47				
Atopobium minutum		0.87	0.08				
57011030	**Very good identification**						**57011030**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.59				
57011032	**Good identification**						**57011032**
Eubacterium saburreum		100.0	0.46				
57011040	**Good identification**						**57011040**
Eubacterium saburreum		96.69	0.39	SOR 1			
Fusobacterium mortiferum		3.31	0.16				
57011050	**Good identification**						**57011050**
Eubacterium saburreum		100.0	0.37	SOR 1			
57011100	**Very good identification**						**57011100**
Eubacterium saburreum		96.67	0.79				
Fusobacterium mortiferum		3.31	0.56				
Odoribacter splanchnicus		0.02	-.03				
57011101	**Low discrimination**						**57011101**
					SPO	GRA	
Eubacterium saburreum		69.63	0.45	URE 1	1	85	
Clostridium chauvoei		27.99	0.29		99	99	
Fusobacterium mortiferum		2.39	0.22		1	1	
57011102	**Very good identification**						**57011102**
Eubacterium saburreum		97.89	0.67				
Fusobacterium mortiferum		2.11	0.40				
57011110	**Very good identification**						**57011110**
Eubacterium saburreum		99.96	0.78				
Fusobacterium mortiferum		0.04	0.22				
57011111	**Good identification**						**57011111**
Eubacterium saburreum		100.0	0.44	URE 1			
57011112	**Very good identification**						**57011112**
Eubacterium saburreum		100.0	0.65				
57011120	**Low discrimination**						**57011120**
Eubacterium saburreum		88.40	0.67				
Odoribacter splanchnicus		10.54	0.32				
Prevotella melaninogenica		0.52	0.12				
57011122	**Good identification**						**57011122**
					CAT	GRA	
Eubacterium saburreum		95.41	0.54		1	85	
Capnocytophaga ochracea		1.90	0.12		90	1	
Clostridium perfringens		1.07	0.11		1	99	
57011130	**Very good identification**						**57011130**
Eubacterium saburreum		99.85	0.65				
Odoribacter splanchnicus		0.15	-.03				
57011132	**Good identification**						**57011132**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.52				
57011140	**Good identification**						**57011140**
Eubacterium saburreum		96.69	0.45	SOR 1			
Fusobacterium mortiferum		3.31	0.22				
57011150	**Good identification**						**57011150**
Eubacterium saburreum		100.0	0.44	SOR 1			
57011200	**Good identification**						**57011200**
Eubacterium saburreum		96.69	0.39	RHA 1			
Fusobacterium mortiferum		3.31	0.16				
57011210	**Good identification**						**57011210**
Eubacterium saburreum		100.0	0.37	RHA 1			
57011300	**Good identification**						**57011300**
Eubacterium saburreum		96.69	0.45	RHA 1			
Fusobacterium mortiferum		3.31	0.22				
57011310	**Good identification**						**57011310**
Eubacterium saburreum		100.0	0.44	RHA 1			
57011400	**Acceptable identification**						**57011400**
Eubacterium saburreum		92.31	0.39	ARA 1			
Odoribacter splanchnicus		4.53	-.03				
Fusobacterium mortiferum		3.16	0.16				
57011410	**Good identification**						**57011410**
Eubacterium saburreum		100.0	0.37	ARA 1			
57011510	**Good identification**						**57011510**
Eubacterium saburreum		97.49	0.44	ARA 1			
Odoribacter splanchnicus		2.51	-.03				
57011520	**Very good identification**						**57011520**
Odoribacter splanchnicus		99.91	0.66	SUC 1			
Eubacterium saburreum		0.09	0.32				
57011560	**Good identification**						**57011560**
Odoribacter splanchnicus		100.0	0.50	SUC 1			
57013000	**Low discrimination**						**57013000**
					CAT	GRA	
Eubacterium saburreum		86.27	0.39	MAN 1	1	85	
Propionibacterium granulosum		7.97	0.20		90	99	
Fusobacterium mortiferum		2.96	0.16		5	1	
57013010	**Good identification**						**57013010**
					SPO		
Eubacterium saburreum		96.96	0.37	MAN 1	1		
Clostridium difficile		3.04	-.03		99		
57013100	**Good identification**						**57013100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		96.69	0.45	MAN 1		
Fusobacterium mortiferum		3.31	0.22			
57013110	**Good identification**					**57013110**
Eubacterium saburreum		100.0	0.44	MAN 1		
57014100	**Low discrimination**					**57014100**
Eubacterium saburreum		75.99	0.41	GAL 99		
Fusobacterium mortiferum		24.01	0.34			
57014110	**Good identification**					**57014110**
Eubacterium saburreum		100.0	0.39	GAL 99		
57015000	**Very good identification**					**57015000**
Eubacterium saburreum		99.52	0.69			
Fusobacterium mortiferum		0.48	0.30			
57015002	**Good identification**					**57015002**
Eubacterium saburreum		99.70	0.56			
Fusobacterium mortiferum		0.30	0.14			
57015010	**Very good identification**					**57015010**
Eubacterium saburreum		100.0	0.67			
57015012	**Good identification**					**57015012**
Eubacterium saburreum		100.0	0.54			
57015020	**Good identification**					**57015020**
Eubacterium saburreum		100.0	0.56			
57015022	**Good identification**					**57015022**
Eubacterium saburreum		100.0	0.43			
57015030	**Good identification**					**57015030**
Eubacterium saburreum		100.0	0.54			
57015032	**Good identification**					**57015032**
Eubacterium saburreum		100.0	0.41			
57015100	**Very good identification**					**57015100**
Eubacterium saburreum		99.52	0.75			
Fusobacterium mortiferum		0.48	0.37			
57015101	**Good identification**					**57015101**
Eubacterium saburreum		100.0	0.41	URE 1		
57015102	**Very good identification**					**57015102**
Eubacterium saburreum		99.70	0.62			
Fusobacterium mortiferum		0.30	0.20			
57015110	**Very good identification**					**57015110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.73			
57015111	**Good identification**					**57015111**
Eubacterium saburreum		100.0	0.39	URE 1		
57015112	**Very good identification**					**57015112**
Eubacterium saburreum		100.0	0.60			
57015120	**Very good identification**					**57015120**
Eubacterium saburreum		99.78	0.62			
Odoribacter splanchnicus		0.22	-.03			
57015122	**Good identification**					**57015122**
					CAT	
Eubacterium saburreum		97.40	0.49		1	
Bacteroides fragilis		1.70	-.03		99	
Capnocytophaga ochracea		0.90	0.02		90	
57015130	**Very good identification**					**57015130**
Eubacterium saburreum		100.0	0.60			
57015132	**Good identification**					**57015132**
Eubacterium saburreum		100.0	0.48			
57015140	**Good identification**					**57015140**
Eubacterium saburreum		100.0	0.41	SOR 1		
57015150	**Good identification**					**57015150**
Eubacterium saburreum		100.0	0.39	SOR 1		
57015300	**Good identification**					**57015300**
Eubacterium saburreum		100.0	0.41	RHA 1		
57015310	**Good identification**					**57015310**
Eubacterium saburreum		100.0	0.39	RHA 1		
57015500	**Good identification**					**57015500**
Eubacterium saburreum		96.24	0.41	ARA 1		
Odoribacter splanchnicus		3.76	-.03			
57015510	**Good identification**					**57015510**
Eubacterium saburreum		100.0	0.39	ARA 1		
57017100	**Good identification**					**57017100**
Eubacterium saburreum		100.0	0.41	MAN 1		
57017110	**Good identification**					**57017110**
Eubacterium saburreum		100.0	0.39	MAN 1		
57021000	**Very good identification**					**57021000**
Eubacterium saburreum		100.0	0.62			
57021002	**Good identification**					**57021002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.49			
57021010	**Very good identification**					**57021010**
Eubacterium saburreum		100.0	0.60			
57021012	**Good identification**					**57021012**
Eubacterium saburreum		100.0	0.48			
57021020	**Good identification**					**57021020**
Eubacterium saburreum		100.0	0.49			
57021022	**Good identification**					**57021022**
Eubacterium saburreum		100.0	0.36			
57021030	**Good identification**					**57021030**
Eubacterium saburreum		100.0	0.48			
57021100	**Very good identification**					**57021100**
Eubacterium saburreum		99.95	0.68			
Fusobacterium mortiferum		0.05	0.13			
57021102	**Good identification**					**57021102**
Eubacterium saburreum		100.0	0.55			
57021110	**Very good identification**					**57021110**
Eubacterium saburreum		100.0	0.67			
57021112	**Good identification**					**57021112**
Eubacterium saburreum		100.0	0.54			
57021120	**Good identification**					**57021120**
Eubacterium saburreum		99.71	0.55			
Odoribacter splanchnicus		0.29	-.07			
57021122	**Good identification**					**57021122**
Eubacterium saburreum		100.0	0.42			
57021130	**Good identification**					**57021130**
Eubacterium saburreum		100.0	0.54			
57021132	**Good identification**					**57021132**
Eubacterium saburreum		100.0	0.41			
57025000	**Very good identification**					**57025000**
Eubacterium saburreum		100.0	0.57			
57025002	**Good identification**					**57025002**
Eubacterium saburreum		100.0	0.44			
57025010	**Good identification**					**57025010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.56				
57025012	**Good identification**						**57025012**
Eubacterium saburreum		100.0	0.43				
57025020	**Good identification**						**57025020**
Eubacterium saburreum		100.0	0.44				
57025030	**Good identification**						**57025030**
Eubacterium saburreum		100.0	0.43				
57025100	**Very good identification**						**57025100**
Eubacterium saburreum		100.0	0.64				
57025102	**Good identification**						**57025102**
Eubacterium saburreum		100.0	0.51				
57025110	**Very good identification**						**57025110**
Eubacterium saburreum		100.0	0.62				
57025112	**Good identification**						**57025112**
Eubacterium saburreum		100.0	0.49				
57025120	**Good identification**						**57025120**
Eubacterium saburreum		100.0	0.51				
57025122	**Good identification**						**57025122**
Eubacterium saburreum		100.0	0.38				
57025130	**Good identification**						**57025130**
Eubacterium saburreum		100.0	0.49				
57025132	**Good identification**						**57025132**
Eubacterium saburreum		100.0	0.36				
57030000	**Low discrimination**						**57030000**
					CAT	GRA	
Fusobacterium mortiferum		37.66	0.38	IND 1	5	1	
Eubacterium saburreum		37.18	0.36		1	85	
Propionibacterium granulosum		20.49	0.30		90	99	
57030100	**Low discrimination**						**57030100**
Fusobacterium mortiferum		50.32	0.44	IND 1			
Eubacterium saburreum		49.68	0.42				
57030110	**Good identification**						**57030110**
Eubacterium saburreum		100.0	0.41	GAL 99			
57031000	**Very good identification**						**57031000**
					CAT	GRA	
Eubacterium saburreum		97.95	0.70		1	85	
Fusobacterium mortiferum		1.50	0.41		5	1	
Propionibacterium granulosum		0.55	0.30		90	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57031001	**Good identification**						**57031001**
Eubacterium saburreum		100.0	0.36	URE 1			
57031002	**Very good identification**						**57031002**
					CAT	GRA	
Eubacterium saburreum		98.28	0.57		1	85	
Fusobacterium mortiferum		0.95	0.24		5	1	
Propionibacterium granulosum		0.78	0.20		90	99	
57031010	**Very good identification**						**57031010**
					CAT		
Eubacterium saburreum		99.91	0.69		1		
Propionibacterium granulosum		0.09	0.15		90		
57031012	**Good identification**						**57031012**
Eubacterium saburreum		100.0	0.56				
57031020	**Very good identification**						**57031020**
					CAT		
Eubacterium saburreum		98.85	0.57		1		
Propionibacterium granulosum		0.78	0.20		90		
Clostridium perfringens		0.37	0.07		1		
57031022	**Good identification**						**57031022**
					CAT		
Eubacterium saburreum		97.93	0.44		1		
Propionibacterium granulosum		1.09	0.09		90		
Clostridium perfringens		0.97	0.01		1		
57031030	**Good identification**						**57031030**
Eubacterium saburreum		100.0	0.56				
57031032	**Good identification**						**57031032**
Eubacterium saburreum		100.0	0.43				
57031040	**Good identification**						**57031040**
Eubacterium saburreum		100.0	0.36	SOR 1			
57031100	**Very good identification**						**57031100**
					COC	GRA	
Eubacterium saburreum		98.34	0.76		1	85	
Fusobacterium mortiferum		1.51	0.47		1	1	
Atopobium parvulum		0.16	0.19		99	99	
57031101	**Acceptable identification**						**57031101**
					SPO	GRA	
Eubacterium saburreum		92.39	0.42	URE 1	1	85	
Clostridium chauvoei		6.19	0.12		99	99	
Fusobacterium mortiferum		1.42	0.13		1	1	
57031102	**Very good identification**						**57031102**
					COC	GRA	
Eubacterium saburreum		98.89	0.64		1	85	
Fusobacterium mortiferum		0.96	0.31		1	1	
Atopobium parvulum		0.16	0.06		99	99	
57031110	**Very good identification**						**57031110**
Eubacterium saburreum		99.98	0.75				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		0.02	0.13			
57031111	**Good identification**					**57031111**
Eubacterium saburreum		100.0	0.41	URE 1		
57031112	**Very good identification**					**57031112**
Eubacterium saburreum		100.0	0.62			
57031120	**Very good identification**					**57031120**
					GRA	
Eubacterium saburreum		98.43	0.64		85	
Clostridium perfringens		1.15	0.21		99	
Odoribacter splanchnicus		0.18	-.03		1	
57031122	**Good identification**					**57031122**
Eubacterium saburreum		96.97	0.51			
Clostridium perfringens		3.03	0.16			
57031130	**Very good identification**					**57031130**
Eubacterium saburreum		100.0	0.62			
57031132	**Good identification**					**57031132**
Eubacterium saburreum		100.0	0.49			
57031140	**Good identification**					**57031140**
Eubacterium saburreum		98.49	0.42	SOR 1		
Fusobacterium mortiferum		1.51	0.13			
57031150	**Good identification**					**57031150**
Eubacterium saburreum		100.0	0.41	SOR 1		
57031200	**Good identification**					**57031200**
Eubacterium saburreum		100.0	0.36	RHA 1		
57031300	**Good identification**					**57031300**
Eubacterium saburreum		98.49	0.42	RHA 1		
Fusobacterium mortiferum		1.51	0.13			
57031310	**Good identification**					**57031310**
Eubacterium saburreum		100.0	0.41	RHA 1		
57031400	**Good identification**					**57031400**
Eubacterium saburreum		100.0	0.36	ARA 1		
57031500	**Good identification**					**57031500**
Eubacterium saburreum		95.52	0.42	ARA 1		
Odoribacter splanchnicus		3.01	-.03			
Fusobacterium mortiferum		1.47	0.13			
57031510	**Good identification**					**57031510**
Eubacterium saburreum		100.0	0.41	ARA 1		
57033000	**Low discrimination**					**57033000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Eubacterium saburreum		70.64	0.36	MAN 1	1		
Propionibacterium granulosum		29.36	0.28		90		
57033100	**Good identification**						**57033100**
Eubacterium saburreum		98.49	0.42	MAN 1			
Fusobacterium mortiferum		1.51	0.13				
57033110	**Good identification**						**57033110**
Eubacterium saburreum		100.0	0.41	MAN 1			
57034100	**Low discrimination**						**57034100**
Eubacterium saburreum		87.60	0.38	GAL 99			
Fusobacterium mortiferum		12.40	0.25				
57034110	**Good identification**						**57034110**
Eubacterium saburreum		100.0	0.36	GAL 99			
57035000	**Very good identification**						**57035000**
Eubacterium saburreum		99.79	0.66				
Fusobacterium mortiferum		0.21	0.22				
57035002	**Good identification**						**57035002**
Eubacterium saburreum		100.0	0.53				
57035010	**Very good identification**						**57035010**
Eubacterium saburreum		100.0	0.64				
57035012	**Good identification**						**57035012**
Eubacterium saburreum		100.0	0.51				
57035020	**Good identification**						**57035020**
Eubacterium saburreum		100.0	0.53				
57035022	**Good identification**						**57035022**
Eubacterium saburreum		100.0	0.40				
57035030	**Good identification**						**57035030**
Eubacterium saburreum		100.0	0.51				
57035032	**Good identification**						**57035032**
Eubacterium saburreum		100.0	0.38				
57035100	**Very good identification**						**57035100**
					COC	GRA	
Eubacterium saburreum		99.25	0.72		1	85	
Atopobium parvulum		0.54	0.23		99	99	
Fusobacterium mortiferum		0.21	0.28		1	1	
57035101	**Good identification**						**57035101**
Eubacterium saburreum		100.0	0.38	URE 1			
57035102	**Very good identification**						**57035102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Eubacterium saburreum		99.32	0.59		1	85	
Atopobium parvulum		0.54	0.10		99	99	
Fusobacterium mortiferum		0.13	0.11		1	1	
57035110	**Very good identification**						**57035110**
Eubacterium saburreum		100.0	0.70				
57035111	**Good identification**						**57035111**
Eubacterium saburreum		100.0	0.36	URE 1			
57035112	**Very good identification**						**57035112**
Eubacterium saburreum		100.0	0.57				
57035120	**Very good identification**						**57035120**
					COC		
Eubacterium saburreum		99.46	0.59		1		
Atopobium parvulum		0.54	0.10		99		
57035122	**Good identification**						**57035122**
Eubacterium saburreum		98.79	0.46				
Parabacteroides distasonis		1.21	0.14				
57035130	**Very good identification**						**57035130**
Eubacterium saburreum		100.0	0.57				
57035132	**Good identification**						**57035132**
Eubacterium saburreum		100.0	0.45				
57035140	**Good identification**						**57035140**
Eubacterium saburreum		100.0	0.38	SOR 1			
57035150	**Good identification**						**57035150**
Eubacterium saburreum		100.0	0.36	SOR 1			
57035300	**Good identification**						**57035300**
Eubacterium saburreum		100.0	0.38	RHA 1			
57035310	**Good identification**						**57035310**
Eubacterium saburreum		100.0	0.36	RHA 1			
57035500	**Good identification**						**57035500**
Eubacterium saburreum		100.0	0.38	ARA 1			
57035510	**Good identification**						**57035510**
Eubacterium saburreum		100.0	0.36	ARA 1			
57035722	**Good identification**						**57035722**
Bacteroides thetaiotaomicron		99.38	0.40	MLT 94			
Parabacteroides distasonis		0.62	0.18	RAF 99			
57037100	**Good identification**						**57037100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.38	MAN 1			
57037110	**Good identification**						**57037110**
Eubacterium saburreum		100.0	0.36	MAN 1			
57041100	**Good identification**						**57041100**
Eubacterium saburreum		96.69	0.37	CEL 1			
Fusobacterium mortiferum		3.31	0.14				
57041110	**Good identification**						**57041110**
Eubacterium saburreum		100.0	0.36	CEL 1			
57050000	**Low discrimination**						**57050000**
					GRA	SPO	
Fusobacterium mortiferum		89.97	0.39	IND 1	1	1	
Clostridium bifermentans		5.11	0.04		99	99	
Fusobacterium varium		4.92	-.03		1	1	
57050100	**Good identification**						**57050100**
Fusobacterium mortiferum		98.68	0.45	IND 1			
Eubacterium saburreum		1.32	0.11				
57051000	**Low discrimination**						**57051000**
					GRA		
Fusobacterium mortiferum		48.72	0.42	IND 1	1		
Eubacterium saburreum		43.10	0.39		85		
Lactobacillus jensenii		4.53	0.14		99		
57051010	**Good identification**						**57051010**
Eubacterium saburreum		95.90	0.37	CEL 1			
Lactobacillus jensenii		4.10	0.06				
57051100	**Low discrimination**						**57051100**
					GRA		
Fusobacterium mortiferum		50.38	0.48	IND 1	1		
Eubacterium saburreum		44.58	0.45		85		
Lactobacillus catenaformis		2.43	0.15		99		
57051110	**Good identification**						**57051110**
Eubacterium saburreum		98.62	0.44	CEL 1			
Fusobacterium mortiferum		1.38	0.14				
57055100	**Low discrimination**						**57055100**
Eubacterium saburreum		86.36	0.41	CEL 1			
Fusobacterium mortiferum		13.64	0.29				
57055110	**Good identification**						**57055110**
Eubacterium saburreum		100.0	0.39	CEL 1			
57055522	**Good identification**						**57055522**
					CAT		
Bacteroides uniformis		98.54	0.35	MLT 92	5		
Bacteroides thetaiotaomicron		1.46	0.07	SAL 92	89		
57055722	**Low discrimination**						**57055722**
					CAT		
Bacteroides uniformis		86.68	0.38	MLT 92	5		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		12.39	0.26	SAL 92	89		
Bacteroides eggerthii		0.58	-.03	RAF 97	93		
57070100	**Good identification**						**57070100**
Fusobacterium mortiferum		100.0	0.36	IND 1			
57071000	**Low discrimination**						**57071000**
					GRA		
Eubacterium saburreum		50.53	0.36	CEL 1	85		
Fusobacterium mortiferum		25.59	0.33		1		
Lactobacillus jensenii		23.88	0.22		99		
57071100	**Low discrimination**						**57071100**
					COC	GRA	
Eubacterium saburreum		53.42	0.42	CEL 1	1	85	
Fusobacterium mortiferum		27.05	0.39		1	1	
Atopobium parvulum		19.53	0.25		99	99	
57071110	**Good identification**						**57071110**
Eubacterium saburreum		100.0	0.41	CEL 1			
57075110	**Good identification**						**57075110**
Eubacterium saburreum		100.0	0.36	CEL 1			
57075722	**Good identification**						**57075722**
					CAT		
Bacteroides thetaiotaomicron		97.25	0.45	MLT 94	89		
Bacteroides ovatus		2.08	0.17	RAF 99	68		
Bacteroides uniformis		0.52	0.04		5		
57075732	**Good identification**						**57075732**
Bacteroides thetaiotaomicron		96.47	0.38	MLT 94			
Bacteroides ovatus		3.54	0.14	RAF 99			
57100000	**Very good identification**						**57100000**
					COC	SPO	
Clostridium bifermentans		98.11	0.66	SUC 1	1	99	
Eubacterium saburreum		0.70	0.41		1	1	
Gemella morbillorum		0.51	0.30		99	1	
57100020	**Acceptable identification**						**57100020**
					SPO		
Clostridium bifermentans		92.26	0.45	SUC 1	99		
Clostridium cadaveris		5.09	0.22	NAG 6	99		
Eubacterium saburreum		1.81	0.28		1		
57100040	**Acceptable identification**						**57100040**
Clostridium bifermentans		94.51	0.42	SUC 1			
Clostridium botulinum biovar A		2.91	0.17	SOR 4			
Clostridium botulinum biovar B		2.58	0.13				
57100100	**Low discrimination**						**57100100**
					SPO		
Eubacterium saburreum		62.10	0.47	GAL 99	1		
Clostridium bifermentans		37.90	0.32		99		
57100110	**Good identification**						**57100110**
					SPO		
Eubacterium saburreum		99.25	0.46	GAL 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium bifermentans		0.75	-.03		99		
57101000	**Very good identification**						**57101000**
					COC	SPO	
Eubacterium saburreum		97.87	0.75		1	1	
Clostridium bifermentans		1.41	0.32		1	99	
Gemella morbillorum		0.39	0.26		99	1	
57101001	**Low discrimination**						**57101001**
					SPO		
Eubacterium saburreum		70.69	0.41	URE 1	1		
Clostridium chauvoei		21.31	0.22		99		
Clostridium sordellii		6.99	0.07		99		
57101002	**Very good identification**						**57101002**
					SPO		
Eubacterium saburreum		99.92	0.62		1		
Clostridium bifermentans		0.08	-.03		99		
57101010	**Very good identification**						**57101010**
					SPO		
Eubacterium saburreum		99.98	0.74		1		
Clostridium bifermentans		0.02	-.03		99		
57101011	**Good identification**						**57101011**
Eubacterium saburreum		100.0	0.40	URE 1			
57101012	**Very good identification**						**57101012**
Eubacterium saburreum		100.0	0.61				
57101020	**Very good identification**						**57101020**
					COC	SPO	
Eubacterium saburreum		99.02	0.62		1	1	
Clostridium bifermentans		0.52	0.11		1	99	
Gemella morbillorum		0.25	0.09		99	1	
57101022	**Good identification**						**57101022**
Eubacterium saburreum		100.0	0.49				
57101030	**Very good identification**						**57101030**
Eubacterium saburreum		100.0	0.61				
57101032	**Good identification**						**57101032**
Eubacterium saburreum		100.0	0.48				
57101040	**Acceptable identification**						**57101040**
					SPO		
Eubacterium saburreum		94.40	0.41	SOR 1	1		
Clostridium bifermentans		5.60	0.08		99		
57101050	**Good identification**						**57101050**
Eubacterium saburreum		100.0	0.40	SOR 1			
57101100	**Very good identification**						**57101100**
					SPO		
Eubacterium saburreum		99.86	0.82		1		
Clostridium chauvoei		0.13	0.22		99		
Clostridium bifermentans		0.01	-.03		99		
57101101	**Good identification**						**57101101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium chauvoei		92.75	0.57	IND 1	99		
Eubacterium saburreum		7.25	0.47		1		
57101102	**Very good identification**						**57101102**
					SPO		
Eubacterium saburreum		99.87	0.69		1		
Clostridium chauvoei		0.13	0.10		99		
57101103	**Acceptable identification**						**57101103**
					SPO		
Clostridium chauvoei		92.75	0.44	IND 1	99		
Eubacterium saburreum		7.25	0.35		1		
57101110	**Very good identification**						**57101110**
Eubacterium saburreum		100.0	0.80				
57101111	**Low discrimination**						**57101111**
					SPO		
Eubacterium saburreum		86.36	0.46	URE 1	1		
Clostridium chauvoei		13.64	0.22		99		
57101112	**Very good identification**						**57101112**
Eubacterium saburreum		100.0	0.67				
57101120	**Good identification**						**57101120**
					GRA		
Eubacterium saburreum		94.85	0.69		85		
Prevotella melaninogenica		4.58	0.30		1		
Clostridium perfringens		0.30	0.17		99		
57101121	**Good identification**						**57101121**
					SPO	GRA	
Clostridium chauvoei		95.84	0.48	IND 1	99	99	
Eubacterium saburreum		3.97	0.35		1	85	
Prevotella melaninogenica		0.19	-.04		1	1	
57101122	**Acceptable identification**						**57101122**
					CAT		
Eubacterium saburreum		92.53	0.56		1		
Prevotella melaninogenica		4.82	0.18		1		
Capnocytophaga ochracea		1.38	0.12		90		
57101123	**Good identification**						**57101123**
					SPO		
Clostridium chauvoei		96.03	0.36	IND 1	99		
Eubacterium saburreum		3.98	0.22		1		
57101130	**Very good identification**						**57101130**
Eubacterium saburreum		99.94	0.67				
Prevotella melaninogenica		0.06	-.04				
57101132	**Good identification**						**57101132**
Eubacterium saburreum		100.0	0.54				
57101140	**Good identification**						**57101140**
Eubacterium saburreum		100.0	0.47	SOR 1			
57101150	**Good identification**						**57101150**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.46	SOR 1			
57101200	**Good identification**						**57101200**
					SPO		
Eubacterium saburreum		98.58	0.41	RHA 1	1		
Clostridium bifermentans		1.42	-.03		99		
57101210	**Good identification**						**57101210**
Eubacterium saburreum		100.0	0.40	RHA 1			
57101300	**Good identification**						**57101300**
Eubacterium saburreum		100.0	0.47	RHA 1			
57101310	**Good identification**						**57101310**
Eubacterium saburreum		100.0	0.46	RHA 1			
57101400	**Acceptable identification**						**57101400**
					COC	SPO	
Eubacterium saburreum		94.49	0.41	ARA 1	1	1	
Gemella morbillorum		4.15	0.09		99	1	
Clostridium bifermentans		1.36	-.03		1	99	
57101410	**Good identification**						**57101410**
Eubacterium saburreum		100.0	0.40	ARA 1			
57101500	**Good identification**						**57101500**
Eubacterium saburreum		99.16	0.47	ARA 1			
Odoribacter splanchnicus		0.84	-.07				
57101510	**Good identification**						**57101510**
Eubacterium saburreum		99.30	0.46	ARA 1			
Bifidobacterium longum subsp. longum		0.70	0.05				
57101722	**Good identification**						**57101722**
Bacteroides eggerthii		100.0	0.36	SUC 1			
57103000	**Good identification**						**57103000**
					SPO		
Eubacterium saburreum		98.58	0.41	MAN 1	1		
Clostridium bifermentans		1.42	-.03		99		
57103010	**Good identification**						**57103010**
Eubacterium saburreum		100.0	0.40	MAN 1			
57103100	**Good identification**						**57103100**
Eubacterium saburreum		100.0	0.47	MAN 1			
57103110	**Good identification**						**57103110**
Eubacterium saburreum		100.0	0.46	MAN 1			
57104010	**Good identification**						**57104010**
					SPO		
Eubacterium saburreum		96.84	0.35	GAL 99	1		
Clostridium bifermentans		3.16	-.03		99		
57104100	**Good identification**						**57104100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Eubacterium saburreum		98.87	0.43	GAL 99	1		
Clostridium bifermentans		1.13	-.03		99		
57104110	**Good identification**						**57104110**
Eubacterium saburreum		100.0	0.41	GAL 99			
57105000	**Very good identification**						**57105000**
					COC	SPO	
Eubacterium saburreum		99.89	0.71		1	1	
Gemella morbillorum		0.08	0.09		99	1	
Clostridium bifermentans		0.03	-.03		1	99	
57105001	**Good identification**						**57105001**
Eubacterium saburreum		100.0	0.37	URE 1			
57105002	**Very good identification**						**57105002**
Eubacterium saburreum		100.0	0.58				
57105010	**Very good identification**						**57105010**
Eubacterium saburreum		100.0	0.69				
57105011	**Good identification**						**57105011**
Eubacterium saburreum		100.0	0.35	URE 1			
57105012	**Very good identification**						**57105012**
Eubacterium saburreum		100.0	0.56				
57105020	**Very good identification**						**57105020**
Eubacterium saburreum		100.0	0.58				
57105022	**Good identification**						**57105022**
Eubacterium saburreum		100.0	0.45				
57105030	**Very good identification**						**57105030**
Eubacterium saburreum		100.0	0.56				
57105032	**Good identification**						**57105032**
Eubacterium saburreum		100.0	0.43				
57105040	**Good identification**						**57105040**
Eubacterium saburreum		100.0	0.37	SOR 1			
57105050	**Good identification**						**57105050**
Eubacterium saburreum		100.0	0.35	SOR 1			
57105100	**Very good identification**						**57105100**
Eubacterium saburreum		100.0	0.77				
57105101	**Low discrimination**						**57105101**
					SPO		
Eubacterium saburreum		80.65	0.43	URE 1	1		
Clostridium chauvoei		19.35	0.22		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57105102	**Very good identification**					**57105102**
Eubacterium saburreum		100.0	0.64			
57105110	**Very good identification**					**57105110**
Eubacterium saburreum		100.0	0.75			
57105111	**Good identification**					**57105111**
Eubacterium saburreum		100.0	0.41	URE 1		
57105112	**Very good identification**					**57105112**
Eubacterium saburreum		100.0	0.63			
57105120	**Very good identification**					**57105120**
Eubacterium saburreum		99.91	0.64			
Prevotella melaninogenica		0.09	-.04			
57105122	**Good identification**					**57105122**
					CAT	
Eubacterium saburreum		98.04	0.51		1	
Bacteroides fragilis		1.28	-.03		99	
Capnocytophaga ochracea		0.68	0.02		90	
57105130	**Very good identification**					**57105130**
Eubacterium saburreum		100.0	0.63			
57105132	**Good identification**					**57105132**
Eubacterium saburreum		100.0	0.50			
57105140	**Good identification**					**57105140**
Eubacterium saburreum		100.0	0.43	SOR 1		
57105150	**Good identification**					**57105150**
Eubacterium saburreum		100.0	0.41	SOR 1		
57105200	**Good identification**					**57105200**
Eubacterium saburreum		100.0	0.37	RHA 1		
57105210	**Good identification**					**57105210**
Eubacterium saburreum		100.0	0.35	RHA 1		
57105300	**Good identification**					**57105300**
Eubacterium saburreum		100.0	0.43	RHA 1		
57105310	**Good identification**					**57105310**
Eubacterium saburreum		100.0	0.41	RHA 1		
57105400	**Good identification**					**57105400**
Eubacterium saburreum		100.0	0.37	ARA 1		
57105410	**Good identification**					**57105410**
Eubacterium saburreum		100.0	0.35	ARA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57105500	**Good identification**						**57105500**
Eubacterium saburreum		98.15	0.43	ARA 1			
Bifidobacterium longum subsp. longum		1.85	0.09				
57105510	**Good identification**						**57105510**
Eubacterium saburreum		96.22	0.41	ARA 1			
Bifidobacterium longum subsp. longum		3.78	0.13				
57105722	**Good identification**						**57105722**
Bacteroides eggerthii		99.94	0.56	SUC 1			
Bacteroides thetaiotaomicron		0.06	0.07				
57107000	**Good identification**						**57107000**
Eubacterium saburreum		100.0	0.37	MAN 1			
57107010	**Good identification**						**57107010**
Eubacterium saburreum		100.0	0.35	MAN 1			
57107100	**Good identification**						**57107100**
Eubacterium saburreum		100.0	0.43	MAN 1			
57107110	**Good identification**						**57107110**
Eubacterium saburreum		100.0	0.41	MAN 1			
57110000	**Low discrimination**						**57110000**
					SPO	GRA	
Prevotella intermedia		73.82	0.58	ESL 1	1	1	
Clostridium bifermentans		16.65	0.44		99	99	
Eubacterium saburreum		6.73	0.49		1	85	
57110002	**Low discrimination**						**57110002**
					SPO	GRA	
Eubacterium saburreum		55.15	0.36	GAL 99	1	85	
Prevotella intermedia		34.62	0.23		1	1	
Clostridium bifermentans		7.81	0.10		99	99	
57110010	**Low discrimination**						**57110010**
					SPO	GRA	
Eubacterium saburreum		85.77	0.48	GAL 99	1	85	
Prevotella intermedia		11.61	0.23		1	1	
Clostridium bifermentans		2.62	0.10		99	99	
57110020	**Low discrimination**						**57110020**
					SPO		
Eubacterium saburreum		27.15	0.36	GAL 99	1		
Clostridium cadaveris		25.49	0.22		99		
Clostridium bifermentans		24.30	0.24		99		
57110100	**Low discrimination**						**57110100**
Eubacterium saburreum		88.65	0.56	GAL 99			
Fusobacterium mortiferum		6.19	0.37				
Prevotella intermedia		4.21	0.23				
57110102	**Good identification**						**57110102**
Eubacterium saburreum		95.79	0.43	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		4.21	0.21				
57110110	**Good identification**						**57110110**
Eubacterium saburreum		100.0	0.54	GAL 99			
57110112	**Good identification**						**57110112**
Eubacterium saburreum		100.0	0.41	GAL 99			
57110120	**Low discrimination**						**57110120**
					GRA		
Eubacterium saburreum		80.08	0.43	GAL 99	85		
Prevotella melaninogenica		11.59	0.12		1		
Clostridium perfringens		8.33	0.17		99		
57110130	**Good identification**						**57110130**
Eubacterium saburreum		100.0	0.41	GAL 99			
57111000	**Low discrimination**						**57111000**
					COC	GRA	
Eubacterium saburreum		89.74	0.83		1	85	
Prevotella intermedia		9.94	0.58		1	1	
Gemella morbillorum		0.15	0.27		99	99	
57111001	**Low discrimination**						**57111001**
					SPO	GRA	
Eubacterium saburreum		72.56	0.49	URE 1	1	85	
Clostridium chauvoei		17.01	0.29		99	99	
Prevotella intermedia		8.04	0.23		1	1	
57111002	**Very good identification**						**57111002**
					GRA		
Eubacterium saburreum		99.12	0.71		85		
Prevotella intermedia		0.63	0.23		1		
Lactobacillus jensenii		0.10	0.11		99		
57111003	**Low discrimination**						**57111003**
					SPO		
Eubacterium saburreum		81.01	0.36	URE 1	1		
Clostridium chauvoei		18.99	0.16		99		
57111010	**Very good identification**						**57111010**
					GRA		
Eubacterium saburreum		99.85	0.82		85		
Prevotella intermedia		0.14	0.23		1		
Lactobacillus jensenii		0.01	0.06		99		
57111011	**Good identification**						**57111011**
Eubacterium saburreum		100.0	0.48	URE 1			
57111012	**Very good identification**						**57111012**
Eubacterium saburreum		99.96	0.69				
Lactobacillus jensenii		0.04	0.03				
57111020	**Very good identification**						**57111020**
					GRA		
Eubacterium saburreum		95.76	0.71		85		
Clostridium perfringens		3.17	0.36		99		
Prevotella intermedia		0.61	0.23		1		
57111021	**Low discrimination**						**57111021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Eubacterium saburreum		67.75	0.36	URE 1	1		
Clostridium chauvoei		30.01	0.21		99		
Clostridium perfringens		2.24	0.02		84		
57111022	**Low discrimination**						**57111022**
					GRA		
Eubacterium saburreum		89.46	0.58		85		
Clostridium perfringens		7.90	0.31		99		
Prevotella oralis		1.27	0.14		1		
57111030	**Very good identification**						**57111030**
Eubacterium saburreum		99.96	0.69				
Clostridium perfringens		0.04	0.02				
57111032	**Very good identification**						**57111032**
Eubacterium saburreum		100.0	0.56				
57111040	**Acceptable identification**						**57111040**
Eubacterium saburreum		90.03	0.49	SOR 1			
Prevotella intermedia		9.97	0.23				
57111042	**Good identification**						**57111042**
Eubacterium saburreum		100.0	0.36	SOR 1			
57111050	**Good identification**						**57111050**
Eubacterium saburreum		100.0	0.48	SOR 1			
57111060	**Low discrimination**						**57111060**
Eubacterium saburreum		61.57	0.36	SOR 1			
Clostridium perfringens		38.44	0.24				
57111100	**Very good identification**						**57111100**
					SPO	GRA	
Eubacterium saburreum		99.60	0.90		1	85	
Fusobacterium mortiferum		0.11	0.40		1	1	
Clostridium chauvoei		0.10	0.29		99	99	
57111101	**Good identification**						**57111101**
					SPO		
Clostridium chauvoei		90.87	0.63	IND 1	99		
Eubacterium saburreum		9.13	0.56		1		
57111102	**Very good identification**						**57111102**
					CAT	GRA	
Eubacterium saburreum		98.51	0.77		1	85	
Capnocytophaga ochracea		0.95	0.30		90	1	
Clostridium perfringens		0.15	0.20		1	99	
57111103	**Acceptable identification**						**57111103**
					SPO		
Clostridium chauvoei		90.87	0.50	IND 1	99		
Eubacterium saburreum		9.13	0.43		1		
57111110	**Very good identification**						**57111110**
					CAT		
Eubacterium saburreum		100.0	0.88		1		
Capnocytophaga ochracea		0.00	-.03		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57111111	**Low discrimination**						**57111111**
					SPO		
Eubacterium saburreum		89.06	0.54	URE 1	1		
Clostridium chauvoei		10.94	0.29		99		
57111112	**Very good identification**						**57111112**
					CAT		
Eubacterium saburreum		99.87	0.75		1		
Capnocytophaga ochracea		0.13	0.14		90		
57111113	**Low discrimination**						**57111113**
					SPO		
Eubacterium saburreum		89.06	0.41	URE 1	1		
Clostridium chauvoei		10.94	0.16		99		
57111120	**Low discrimination**						**57111120**
					GRA		
Eubacterium saburreum		79.20	0.77		85		
Prevotella melaninogenica		11.46	0.47		1		
Clostridium perfringens		8.24	0.51		99		
57111121	**Acceptable identification**						**57111121**
					SPO	GRA	
Clostridium chauvoei		93.77	0.55	IND 1	99	99	
Eubacterium saburreum		4.99	0.43		1	85	
Prevotella melaninogenica		0.72	0.12		1	1	
57111122	**Low discrimination**						**57111122**
					CAT	GRA	
Eubacterium saburreum		49.19	0.64		1	85	
Capnocytophaga ochracea		24.23	0.47		90	1	
Clostridium perfringens		13.65	0.45		1	99	
57111123	**Acceptable identification**						**57111123**
					SPO	CAT	
Clostridium chauvoei		90.29	0.42	IND 1	99	1	
Eubacterium saburreum		4.81	0.30		1	1	
Capnocytophaga ochracea		2.37	0.12		1	90	
57111130	**Very good identification**						**57111130**
					GRA		
Eubacterium saburreum		99.56	0.75		85		
Prevotella melaninogenica		0.18	0.12		1		
Clostridium perfringens		0.13	0.17		99		
57111131	**Low discrimination**						**57111131**
					SPO		
Eubacterium saburreum		81.17	0.41	URE 1	1		
Clostridium chauvoei		18.83	0.21		99		
57111132	**Good identification**						**57111132**
					CAT	GRA	
Eubacterium saburreum		93.16	0.62		1	85	
Capnocytophaga ochracea		6.23	0.30		90	1	
Clostridium perfringens		0.32	0.11		1	99	
57111140	**Good identification**						**57111140**
					CAT	GRA	
Eubacterium saburreum		98.74	0.56	SOR 1	1	85	
Clostridium perfringens		1.06	0.13		1	99	
Capnocytophaga ochracea		0.21	-.03		90	1	
57111142	**Low discrimination**						**57111142**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				CAT	GRA	
Eubacterium saburreum	88.12	0.43	SOR 1	1	85	
Capnocytophaga ochracea	9.36	0.14		90	1	
Clostridium perfringens	2.52	0.07		1	99	
57111150 Good identification						**57111150**
Eubacterium saburreum	100.0	0.54	SOR 1			
57111152 Good identification						**57111152**
				CAT		
Eubacterium saburreum	98.58	0.41	SOR 1	1		
Capnocytophaga ochracea	1.42	-.03		90		
57111160 Low discrimination						**57111160**
				GRA		
Clostridium perfringens	61.07	0.39	IND 1	99		
Eubacterium saburreum	31.12	0.43		85		
Prevotella melaninogenica	4.50	0.12		1		
57111170 Good identification						**57111170**
				CAT	GRA	
Eubacterium saburreum	96.28	0.41	SOR 1	1	85	
Clostridium perfringens	2.33	0.04		1	99	
Capnocytophaga ochracea	1.39	-.03		90	1	
57111200 Acceptable identification						**57111200**
Eubacterium saburreum	90.03	0.49	RHA 1			
Prevotella intermedia	9.97	0.23				
57111202 Good identification						**57111202**
Eubacterium saburreum	100.0	0.36	RHA 1			
57111210 Good identification						**57111210**
Eubacterium saburreum	100.0	0.48	RHA 1			
57111220 Good identification						**57111220**
Eubacterium saburreum	96.80	0.36	RHA 1			
Clostridium perfringens	3.21	0.02				
57111300 Good identification						**57111300**
Eubacterium saburreum	100.0	0.56	RHA 1			
57111302 Good identification						**57111302**
Eubacterium saburreum	96.38	0.43	RHA 1			
Prevotella oralis	3.62	0.06				
57111310 Good identification						**57111310**
Eubacterium saburreum	100.0	0.54	RHA 1			
57111312 Good identification						**57111312**
Eubacterium saburreum	100.0	0.41	RHA 1			
57111320 Low discrimination						**57111320**
				GRA		
Eubacterium saburreum	77.01	0.43	RHA 1	85		
Prevotella melaninogenica	11.14	0.12		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		8.01	0.17		99		
57111330	**Good identification**						**57111330**
Eubacterium saburreum		100.0	0.41	RHA 1			
57111400	**Low discrimination**						**57111400**
					COC	GRA	
Eubacterium saburreum		88.00	0.49	ARA 1	1	85	
Prevotella intermedia		9.75	0.23		1	1	
Gemella morbillorum		1.58	0.11		99	99	
57111402	**Good identification**						**57111402**
Eubacterium saburreum		95.83	0.36	ARA 1			
Lactobacillus fermentum		4.17	0.16				
57111410	**Good identification**						**57111410**
Eubacterium saburreum		100.0	0.48	ARA 1			
57111420	**Acceptable identification**						**57111420**
					GRA		
Eubacterium saburreum		90.69	0.36	ARA 1	85		
Odoribacter splanchnicus		6.31	-.03		1		
Clostridium perfringens		3.00	0.02		99		
57111500	**Good identification**						**57111500**
Eubacterium saburreum		97.65	0.56	ARA 1			
Bifidobacterium longum subsp. longum		0.99	0.18				
Lactobacillus fermentum		0.52	0.20				
57111502	**Low discrimination**						**57111502**
					CAT	GRA	
Eubacterium saburreum		77.90	0.43	ARA 1	1	85	
Capnocytophaga ochracea		13.14	0.17		90	1	
Bifidobacterium longum subsp. longum		4.49	0.18		1	99	
57111510	**Good identification**						**57111510**
Eubacterium saburreum		97.93	0.54	ARA 1			
Bifidobacterium longum subsp. longum		2.07	0.22				
57111512	**Low discrimination**						**57111512**
Eubacterium saburreum		84.52	0.41	ARA 1			
Bifidobacterium longum subsp. longum		10.14	0.22				
Bifidobacterium adolescentis		3.41	0.15				
57111530	**Low discrimination**						**57111530**
					GRA		
Eubacterium saburreum		64.98	0.41	ARA 1	85		
Bifidobacterium longum subsp. longum		31.17	0.32		99		
Odoribacter splanchnicus		2.37	-.03		1		
57113000	**Low discrimination**						**57113000**
					CAT	GRA	
Eubacterium saburreum		89.19	0.49	MAN 1	1	85	
Prevotella intermedia		9.88	0.23		1	1	
Propionibacterium granulosum		0.93	0.14		90	99	
57113002	**Good identification**						**57113002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.36	MAN 1			
57113010	**Good identification**						**57113010**
Eubacterium saburreum		100.0	0.48	MAN 1			
57113020	**Good identification**						**57113020**
Eubacterium saburreum		96.80	0.36	MAN 1			
Clostridium perfringens		3.21	0.02				
57113100	**Good identification**						**57113100**
Eubacterium saburreum		99.81	0.56	MAN 1			
Propionibacterium propionicum		0.19	0.12				
57113102	**Good identification**						**57113102**
Eubacterium saburreum		98.37	0.43	MAN 1			
Propionibacterium propionicum		1.63	0.15				
57113110	**Good identification**						**57113110**
Eubacterium saburreum		100.0	0.54	MAN 1			
57113112	**Good identification**						**57113112**
Eubacterium saburreum		100.0	0.41	MAN 1			
57113120	**Low discrimination**						**57113120**
					GRA		
Eubacterium saburreum		80.08	0.43	MAN 1	85		
Prevotella melaninogenica		11.59	0.12		1		
Clostridium perfringens		8.33	0.17		99		
57113130	**Good identification**						**57113130**
Eubacterium saburreum		100.0	0.41	MAN 1			
57114000	**Low discrimination**						**57114000**
					COC	GRA	
Eubacterium saburreum		75.55	0.45	GAL 99	1	85	
Prevotella intermedia		15.54	0.23		1	1	
Gemella morbillorum		4.67	0.15		99	99	
57114010	**Good identification**						**57114010**
Eubacterium saburreum		100.0	0.43	GAL 99			
57114100	**Good identification**						**57114100**
Eubacterium saburreum		99.03	0.51	GAL 99			
Fusobacterium mortiferum		0.97	0.18				
57114102	**Good identification**						**57114102**
Eubacterium saburreum		100.0	0.38	GAL 99			
57114110	**Good identification**						**57114110**
Eubacterium saburreum		100.0	0.49	GAL 99			
57114112	**Good identification**						**57114112**
Eubacterium saburreum		100.0	0.37	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57114120	**Good identification**						**57114120**
Eubacterium saburreum		100.0	0.38	GAL 99			
57114130	**Good identification**						**57114130**
Eubacterium saburreum		100.0	0.37	GAL 99			
57115000	**Very good identification**						**57115000**
					COC	GRA	
Eubacterium saburreum		99.74	0.79		1	85	
Prevotella intermedia		0.21	0.23		1	1	
Gemella morbillorum		0.03	0.11		99	99	
57115001	**Good identification**						**57115001**
Eubacterium saburreum		100.0	0.45	URE 1			
57115002	**Very good identification**						**57115002**
Eubacterium saburreum		99.95	0.66				
Lactobacillus fermentum		0.05	0.12				
57115010	**Very good identification**						**57115010**
Eubacterium saburreum		100.0	0.77				
57115011	**Good identification**						**57115011**
Eubacterium saburreum		100.0	0.43	URE 1			
57115012	**Very good identification**						**57115012**
Eubacterium saburreum		100.0	0.64				
57115020	**Very good identification**						**57115020**
Eubacterium saburreum		99.94	0.66				
Clostridium perfringens		0.06	0.02				
57115022	**Good identification**						**57115022**
					CAT		
Eubacterium saburreum		98.15	0.53		1		
Bacteroides fragilis		1.00	-.03		99		
Capnocytophaga ochracea		0.53	0.02		90		
57115030	**Very good identification**						**57115030**
Eubacterium saburreum		100.0	0.64				
57115032	**Good identification**						**57115032**
Eubacterium saburreum		100.0	0.52				
57115040	**Good identification**						**57115040**
Eubacterium saburreum		100.0	0.45	SOR 1			
57115050	**Good identification**						**57115050**
Eubacterium saburreum		100.0	0.43	SOR 1			
57115100	**Very good identification**						**57115100**
					COC	GRA	
Eubacterium saburreum		99.88	0.85		1	85	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		0.09	0.23		99	99	
Fusobacterium mortiferum		0.02	0.21		1	1	
57115101	**Low discrimination**						**57115101**
					SPO		
Eubacterium saburreum		84.27	0.51	URE 1	1		
Clostridium chauvoei		15.73	0.29		99		
57115102	**Very good identification**						**57115102**
					COC	CAT	
Eubacterium saburreum		99.22	0.72		1	1	
Capnocytophaga ochracea		0.45	0.20		1	90	
Atopobium parvulum		0.09	0.10		99	1	
57115103	**Low discrimination**						**57115103**
					SPO		
Eubacterium saburreum		84.27	0.38	URE 1	1		
Clostridium chauvoei		15.73	0.16		99		
57115110	**Very good identification**						**57115110**
Eubacterium saburreum		100.0	0.84				
57115111	**Good identification**						**57115111**
Eubacterium saburreum		100.0	0.49	URE 1			
57115112	**Very good identification**						**57115112**
					CAT	GRA	
Eubacterium saburreum		99.86	0.71		1	85	
Bifidobacterium adolescentis		0.08	0.15		1	99	
Capnocytophaga ochracea		0.06	0.04		90	1	
57115113	**Good identification**						**57115113**
Eubacterium saburreum		100.0	0.37	URE 1			
57115120	**Very good identification**						**57115120**
					CAT		
Eubacterium saburreum		98.49	0.72		1		
Capnocytophaga ochracea		0.44	0.20		90		
Bacteroides fragilis		0.40	0.10		99		
57115121	**Low discrimination**						**57115121**
					SPO		
Eubacterium saburreum		73.94	0.38	URE 1	1		
Clostridium chauvoei		26.06	0.21		99		
57115122	**Low discrimination**						**57115122**
					CAT		
Eubacterium saburreum		58.60	0.59		1		
Bacteroides fragilis		25.34	0.32		99		
Capnocytophaga ochracea		13.40	0.36		90		
57115130	**Very good identification**						**57115130**
					CAT	GRA	
Eubacterium saburreum		99.91	0.71		1	85	
Capnocytophaga ochracea		0.06	0.04		90	1	
Bifidobacterium longum subsp. longum		0.03	0.06		1	99	
57115131	**Good identification**						**57115131**
Eubacterium saburreum		100.0	0.37	URE 1			
57115132	**Very good identification**						**57115132**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Eubacterium saburreum		96.35	0.58		1		
Capnocytophaga ochracea		2.99	0.20		90		
Bacteroides fragilis		0.51	-.03		99		
57115140	**Good identification**						**57115140**
Eubacterium saburreum		100.0	0.51	SOR 1			
57115142	**Good identification**						**57115142**
					CAT		
Eubacterium saburreum		95.30	0.38	SOR 1	1		
Capnocytophaga ochracea		4.70	0.04		90		
57115150	**Good identification**						**57115150**
Eubacterium saburreum		100.0	0.49	SOR 1			
57115152	**Good identification**						**57115152**
Eubacterium saburreum		100.0	0.37	SOR 1			
57115160	**Acceptable identification**						**57115160**
					CAT	GRA	
Eubacterium saburreum		92.07	0.38	SOR 1	1	85	
Capnocytophaga ochracea		4.54	0.04		90	1	
Clostridium perfringens		3.39	0.04		1	99	
57115170	**Good identification**						**57115170**
Eubacterium saburreum		100.0	0.37	SOR 1			
57115200	**Good identification**						**57115200**
Eubacterium saburreum		100.0	0.45	RHA 1			
57115210	**Good identification**						**57115210**
Eubacterium saburreum		100.0	0.43	RHA 1			
57115300	**Low discrimination**						**57115300**
Eubacterium saburreum		82.11	0.51	RHA 1			
Eubacterium contortum		17.89	0.29				
57115302	**Low discrimination**						**57115302**
Eubacterium saburreum		82.11	0.38	RHA 1			
Eubacterium contortum		17.89	0.16				
57115310	**Good identification**						**57115310**
Eubacterium saburreum		100.0	0.49	RHA 1			
57115312	**Good identification**						**57115312**
Eubacterium saburreum		100.0	0.37	RHA 1			
57115330	**Good identification**						**57115330**
Eubacterium saburreum		100.0	0.37	RHA 1			
57115400	**Good identification**						**57115400**
Eubacterium saburreum		99.17	0.45	ARA 1			
Lactobacillus fermentum		0.83	0.12				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57115410	**Good identification**						**57115410**
Eubacterium saburreum		100.0	0.43	ARA 1			
57115500	**Acceptable identification**						**57115500**
Eubacterium saburreum		94.12	0.51	ARA 1			
Bifidobacterium longum subsp. longum		5.33	0.26				
Lactobacillus fermentum		0.55	0.16				
57115502	**Low discrimination**						**57115502**
Eubacterium saburreum		56.55	0.38	ARA 1			
Bifidobacterium adolescentis		18.99	0.28				
Bifidobacterium longum subsp. longum		18.15	0.26				
57115510	**Low discrimination**						**57115510**
Eubacterium saburreum		89.45	0.49	ARA 1			
Bifidobacterium longum subsp. longum		10.55	0.30				
57115512	**Low discrimination**						**57115512**
Eubacterium saburreum		46.09	0.37	ARA 1			
Bifidobacterium longum subsp. longum		30.79	0.30				
Bifidobacterium adolescentis		23.12	0.29				
57115522	**Low discrimination**						**57115522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		30.85	0.36	IND 1	99	1	
Bacteroides uniformis		23.75	0.19	RAF 97	1	5	
Bacteroides thetaiotaomicron		22.40	0.22		1	89	
57115530	**Low discrimination**						**57115530**
Bifidobacterium longum subsp. longum		72.77	0.40	IND 1			
Eubacterium saburreum		27.23	0.37	RAF 97			
57115532	**Low discrimination**						**57115532**
					CAT	GRA	
Bifidobacterium longum subsp. longum		78.88	0.40	IND 1	1	99	
Bacteroides thetaiotaomicron		13.08	0.15	RAF 97	89	1	
Eubacterium saburreum		5.21	0.24		1	85	
57115720	**Good identification**						**57115720**
					GRA	CAT	
Bacteroides vulgatus		98.00	0.40	IND 1	1	3	
Eubacterium contortum		1.18	0.14	RAF 97	99	1	
Bacteroides thetaiotaomicron		0.83	0.07		1	89	
57115722	**Low discrimination**						**57115722**
					CAT		
Bacteroides thetaiotaomicron		77.46	0.41	RAF 99	89		
Bacteroides eggerthii		12.26	0.21	TRE 93	93		
Bacteroides uniformis		8.54	0.22		5		
57117000	**Good identification**						**57117000**
Eubacterium saburreum		100.0	0.45	MAN 1			
57117010	**Good identification**						**57117010**
Eubacterium saburreum		100.0	0.43	MAN 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
57117100	**Good identification**					**57117100**
Eubacterium saburreum		100.0	0.51	MAN 1		
57117102	**Good identification**					**57117102**
Eubacterium saburreum		96.06	0.38	MAN 1		
Bifidobacterium adolescentis		3.95	0.12			
57117110	**Good identification**					**57117110**
Eubacterium saburreum		100.0	0.49	MAN 1		
57117112	**Acceptable identification**					**57117112**
Eubacterium saburreum		94.22	0.37	MAN 1		
Bifidobacterium adolescentis		5.78	0.14			
57117120	**Good identification**					**57117120**
Eubacterium saburreum		100.0	0.38	MAN 1		
57117130	**Good identification**					**57117130**
Eubacterium saburreum		100.0	0.37	MAN 1		
57120010	**Good identification**					**57120010**
					SPO	
Eubacterium saburreum		97.43	0.37	GAL 99	1	
Clostridium bifermentans		2.57	-.03		99	
57120100	**Good identification**					**57120100**
					SPO	
Eubacterium saburreum		99.08	0.44	GAL 99	1	
Clostridium bifermentans		0.92	-.03		99	
57120110	**Good identification**					**57120110**
Eubacterium saburreum		100.0	0.43	GAL 99		
57121000	**Very good identification**					**57121000**
					SPO	
Eubacterium saburreum		99.98	0.72		1	
Clostridium bifermentans		0.02	-.03		99	
57121001	**Good identification**					**57121001**
					SPO	
Eubacterium saburreum		95.22	0.38	URE 1	1	
Clostridium chauvoei		4.78	0.06		99	
57121002	**Very good identification**					**57121002**
Eubacterium saburreum		100.0	0.59			
57121010	**Very good identification**					**57121010**
Eubacterium saburreum		100.0	0.71			
57121011	**Good identification**					**57121011**
Eubacterium saburreum		100.0	0.37	URE 1		
57121012	**Very good identification**					**57121012**
Eubacterium saburreum		100.0	0.58			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57121020	**Very good identification**						**57121020**
Eubacterium saburreum		99.72	0.59				
Clostridium perfringens		0.28	0.07				
57121022	**Good identification**						**57121022**
Eubacterium saburreum		99.26	0.46				
Clostridium perfringens		0.74	0.01				
57121030	**Very good identification**						**57121030**
Eubacterium saburreum		100.0	0.58				
57121032	**Good identification**						**57121032**
Eubacterium saburreum		100.0	0.45				
57121040	**Good identification**						**57121040**
Eubacterium saburreum		100.0	0.38	SOR 1			
57121050	**Good identification**						**57121050**
Eubacterium saburreum		100.0	0.37	SOR 1			
57121100	**Very good identification**						**57121100**
					COC	SPO	
Eubacterium saburreum		99.86	0.79		1	1	
Atopobium parvulum		0.12	0.19		99	1	
Clostridium chauvoei		0.02	0.06		1	99	
57121101	**Low discrimination**						**57121101**
					SPO		
Clostridium chauvoei		68.07	0.40	IND 1	99		
Eubacterium saburreum		31.93	0.44	TRE 10	1		
57121102	**Very good identification**						**57121102**
					COC		
Eubacterium saburreum		99.88	0.66		1		
Atopobium parvulum		0.12	0.06		99		
57121110	**Very good identification**						**57121110**
Eubacterium saburreum		100.0	0.77				
57121111	**Good identification**						**57121111**
					SPO		
Eubacterium saburreum		97.44	0.43	URE 1	1		
Clostridium chauvoei		2.56	0.06		99		
57121112	**Very good identification**						**57121112**
Eubacterium saburreum		100.0	0.64				
57121120	**Very good identification**						**57121120**
					COC		
Eubacterium saburreum		98.94	0.66		1		
Clostridium perfringens		0.87	0.21		1		
Atopobium parvulum		0.12	0.06		99		
57121122	**Good identification**						**57121122**
Eubacterium saburreum		97.71	0.53				
Clostridium perfringens		2.29	0.16				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
57121130	**Very good identification**					**57121130**
Eubacterium saburreum		100.0	0.64			
57121132	**Good identification**					**57121132**
Eubacterium saburreum		100.0	0.51			
57121140	**Good identification**					**57121140**
Eubacterium saburreum		100.0	0.44	SOR 1		
57121150	**Good identification**					**57121150**
Eubacterium saburreum		100.0	0.43	SOR 1		
57121200	**Good identification**					**57121200**
Eubacterium saburreum		100.0	0.38	RHA 1		
57121210	**Good identification**					**57121210**
Eubacterium saburreum		100.0	0.37	RHA 1		
57121300	**Good identification**					**57121300**
Eubacterium saburreum		100.0	0.44	RHA 1		
57121310	**Good identification**					**57121310**
Eubacterium saburreum		100.0	0.43	RHA 1		
57121400	**Good identification**					**57121400**
Eubacterium saburreum		100.0	0.38	ARA 1		
57121410	**Good identification**					**57121410**
Eubacterium saburreum		100.0	0.37	ARA 1		
57121500	**Good identification**					**57121500**
Eubacterium saburreum		100.0	0.44	ARA 1		
57121510	**Good identification**					**57121510**
Eubacterium saburreum		100.0	0.43	ARA 1		
57123000	**Good identification**					**57123000**
Eubacterium saburreum		100.0	0.38	MAN 1		
57123010	**Good identification**					**57123010**
Eubacterium saburreum		100.0	0.37	MAN 1		
57123100	**Good identification**					**57123100**
Eubacterium saburreum		100.0	0.44	MAN 1		
57123110	**Good identification**					**57123110**
Eubacterium saburreum		100.0	0.43	MAN 1		
57124100	**Good identification**					**57124100**
Eubacterium saburreum		100.0	0.40	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57124110	**Good identification**					**57124110**
Eubacterium saburreum		100.0	0.38	GAL 99		
57125000	**Very good identification**					**57125000**
Eubacterium saburreum		100.0	0.68			
57125002	**Good identification**					**57125002**
Eubacterium saburreum		100.0	0.55			
57125010	**Very good identification**					**57125010**
Eubacterium saburreum		100.0	0.66			
57125012	**Good identification**					**57125012**
Eubacterium saburreum		100.0	0.53			
57125020	**Good identification**					**57125020**
Eubacterium saburreum		100.0	0.55			
57125022	**Good identification**					**57125022**
Eubacterium saburreum		100.0	0.42			
57125030	**Good identification**					**57125030**
Eubacterium saburreum		100.0	0.53			
57125032	**Good identification**					**57125032**
Eubacterium saburreum		100.0	0.40			
57125100	**Very good identification**					**57125100**
					COC	
Eubacterium saburreum		99.59	0.74		1	
Atopobium parvulum		0.41	0.23		99	
57125101	**Good identification**					**57125101**
					SPO	
Eubacterium saburreum		96.16	0.40	URE 1	1	
Clostridium chauvoei		3.85	0.06		99	
57125102	**Very good identification**					**57125102**
					COC	
Eubacterium saburreum		99.59	0.61		1	
Atopobium parvulum		0.41	0.10		99	
57125110	**Very good identification**					**57125110**
Eubacterium saburreum		100.0	0.72			
57125111	**Good identification**					**57125111**
Eubacterium saburreum		100.0	0.38	URE 1		
57125112	**Very good identification**					**57125112**
Eubacterium saburreum		100.0	0.60			
57125120	**Very good identification**					**57125120**
					COC	
Eubacterium saburreum		99.59	0.61		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		0.41	0.10		99		
57125122	**Good identification**						**57125122**
Eubacterium saburreum		100.0	0.48				
57125130	**Very good identification**						**57125130**
Eubacterium saburreum		100.0	0.60				
57125132	**Good identification**						**57125132**
Eubacterium saburreum		100.0	0.47				
57125140	**Good identification**						**57125140**
Eubacterium saburreum		100.0	0.40	SOR 1			
57125150	**Good identification**						**57125150**
Eubacterium saburreum		100.0	0.38	SOR 1			
57125300	**Good identification**						**57125300**
Eubacterium saburreum		100.0	0.40	RHA 1			
57125310	**Good identification**						**57125310**
Eubacterium saburreum		100.0	0.38	RHA 1			
57125500	**Good identification**						**57125500**
Eubacterium saburreum		100.0	0.40	ARA 1			
57125510	**Good identification**						**57125510**
Eubacterium saburreum		100.0	0.38	ARA 1			
57127100	**Good identification**						**57127100**
Eubacterium saburreum		100.0	0.40	MAN 1			
57127110	**Good identification**						**57127110**
Eubacterium saburreum		100.0	0.38	MAN 1			
57130000	**Low discrimination**						**57130000**
					CAT	GRA	
Eubacterium saburreum		76.54	0.46	GAL 99	1	85	
Prevotella intermedia		12.72	0.23		1	1	
Propionibacterium granulosum		4.74	0.24		90	99	
57130010	**Good identification**						**57130010**
					CAT		
Eubacterium saburreum		98.98	0.45	GAL 99	1		
Propionibacterium granulosum		1.02	0.09		90		
57130100	**Acceptable identification**						**57130100**
					COC	GRA	
Eubacterium saburreum		93.42	0.53	GAL 99	1	85	
Atopobium parvulum		3.66	0.19		99	99	
Fusobacterium mortiferum		2.92	0.28		1	1	
57130102	**Acceptable identification**						**57130102**
					COC	GRA	
Eubacterium saburreum		94.44	0.40	GAL 99	1	85	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		3.70	0.06		99	99	
Fusobacterium mortiferum		1.86	0.12		1	1	
57130110	**Good identification**						**57130110**
Eubacterium saburreum		100.0	0.51	GAL 99			
57130112	**Good identification**						**57130112**
Eubacterium saburreum		100.0	0.38	GAL 99			
57130120	**Low discrimination**						**57130120**
					COC		
Eubacterium saburreum		75.24	0.40	GAL 99	1		
Clostridium perfringens		21.81	0.21		1		
Atopobium parvulum		2.95	0.06		99		
57130130	**Good identification**						**57130130**
Eubacterium saburreum		100.0	0.38	GAL 99			
57131000	**Very good identification**						**57131000**
					GRA		
Eubacterium saburreum		99.46	0.80		85		
Prevotella intermedia		0.17	0.23		1		
Lactobacillus jensenii		0.12	0.22		99		
57131001	**Good identification**						**57131001**
					SPO		
Eubacterium saburreum		96.24	0.46	URE 1	1		
Clostridium chauvoei		3.76	0.12		99		
57131002	**Very good identification**						**57131002**
Eubacterium saburreum		99.24	0.68				
Lactobacillus jensenii		0.45	0.19				
Clostridium perfringens		0.13	0.09				
57131010	**Very good identification**						**57131010**
					CAT		
Eubacterium saburreum		99.94	0.79		1		
Lactobacillus jensenii		0.05	0.14		1		
Propionibacterium granulosum		0.01	0.09		90		
57131011	**Good identification**						**57131011**
Eubacterium saburreum		100.0	0.45	URE 1			
57131012	**Very good identification**						**57131012**
Eubacterium saburreum		99.82	0.66				
Lactobacillus jensenii		0.18	0.11				
57131020	**Good identification**						**57131020**
Eubacterium saburreum		91.26	0.68				
Clostridium perfringens		8.42	0.41				
Lactobacillus jensenii		0.15	0.12				
57131022	**Low discrimination**						**57131022**
Eubacterium saburreum		79.85	0.55				
Clostridium perfringens		19.64	0.35				
Lactobacillus jensenii		0.51	0.09				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57131030	**Very good identification**						**57131030**
Eubacterium saburreum		99.82	0.66				
Clostridium perfringens		0.11	0.07				
Lactobacillus jensenii		0.07	0.04				
57131032	**Good identification**						**57131032**
Eubacterium saburreum		99.70	0.53				
Clostridium perfringens		0.30	0.01				
57131040	**Good identification**						**57131040**
					SPO		
Eubacterium saburreum		98.11	0.46	SOR 1	1		
Clostridium botulinum biovar B		0.96	-.03		99		
Clostridium perfringens		0.93	0.03		84		
57131050	**Good identification**						**57131050**
Eubacterium saburreum		100.0	0.45	SOR 1			
57131100	**Very good identification**						**57131100**
					COC		
Eubacterium saburreum		96.03	0.87		1		
Atopobium parvulum		3.76	0.53		99		
Clostridium perfringens		0.15	0.30		1		
57131101	**Low discrimination**						**57131101**
					COC	SPO	
Clostridium chauvoei		61.47	0.47	IND 1	1	99	
Eubacterium saburreum		37.08	0.53	TRE 10	1	1	
Atopobium parvulum		1.45	0.19		99	1	
57131102	**Very good identification**						**57131102**
					COC		
Eubacterium saburreum		95.80	0.74		1		
Atopobium parvulum		3.75	0.40		99		
Clostridium perfringens		0.40	0.24		1		
57131110	**Very good identification**						**57131110**
					COC		
Eubacterium saburreum		99.95	0.85		1		
Atopobium parvulum		0.05	0.19		99		
57131111	**Good identification**						**57131111**
					SPO		
Eubacterium saburreum		97.99	0.51	URE 1	1		
Clostridium chauvoei		2.01	0.12		99		
57131112	**Very good identification**						**57131112**
					COC	GRA	
Eubacterium saburreum		99.88	0.72		1	85	
Leptotrichia buccalis		0.07	-.03		1	1	
Atopobium parvulum		0.05	0.06		99	99	
57131113	**Good identification**						**57131113**
					SPO		
Eubacterium saburreum		97.99	0.38	URE 1	1		
Clostridium chauvoei		2.01	-.01		99		
57131120	**Low discrimination**						**57131120**
					COC		
Eubacterium saburreum		74.97	0.74		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		21.73	0.56		1	
Atopobium parvulum		2.94	0.40		99	
57131121	**Low discrimination**					**57131121**
					SPO	
Clostridium chauvoei		70.21	0.38	IND 1	99	
Eubacterium saburreum		22.42	0.40	TRE 10	1	
Clostridium perfringens		6.50	0.21		84	
57131122	**Low discrimination**					**57131122**
					COC	
Eubacterium saburreum		54.61	0.61		1	
Clostridium perfringens		42.21	0.50		1	
Atopobium parvulum		2.14	0.27		99	
57131130	**Very good identification**					**57131130**
					COC	
Eubacterium saburreum		99.60	0.72		1	
Clostridium perfringens		0.36	0.21		1	
Atopobium parvulum		0.05	0.06		99	
57131131	**Good identification**					**57131131**
					SPO	
Eubacterium saburreum		96.28	0.38	URE 1	1	
Clostridium chauvoei		3.72	0.04		99	
57131132	**Very good identification**					**57131132**
Eubacterium saburreum		99.06	0.59			
Clostridium perfringens		0.95	0.16			
57131140	**Acceptable identification**					**57131140**
					COC	
Eubacterium saburreum		93.55	0.53	SOR 1	1	
Atopobium parvulum		3.67	0.19		99	
Clostridium perfringens		2.79	0.17		1	
57131142	**Low discrimination**					**57131142**
					COC	
Eubacterium saburreum		89.39	0.40	SOR 1	1	
Clostridium perfringens		7.11	0.12		1	
Atopobium parvulum		3.50	0.06		99	
57131150	**Good identification**					**57131150**
Eubacterium saburreum		100.0	0.51	SOR 1		
57131152	**Good identification**					**57131152**
Eubacterium saburreum		100.0	0.38	SOR 1		
57131160	**Low discrimination**					**57131160**
					COC	
Clostridium perfringens		84.03	0.43	IND 1	1	
Eubacterium saburreum		15.37	0.40		1	
Atopobium parvulum		0.60	0.06		99	
57131162	**Acceptable identification**					**57131162**
Clostridium perfringens		93.58	0.38	IND 1		
Eubacterium saburreum		6.42	0.27			
57131170	**Acceptable identification**					**57131170**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		93.68	0.38	SOR 1		
Clostridium perfringens		6.32	0.09			
57131200	**Good identification**					**57131200**
Eubacterium saburreum		100.0	0.46	RHA 1		
57131210	**Good identification**					**57131210**
Eubacterium saburreum		100.0	0.45	RHA 1		
57131300	**Good identification**					**57131300**
					COC	
Eubacterium saburreum		96.23	0.53	RHA 1	1	
Atopobium parvulum		3.77	0.19		99	
57131302	**Good identification**					**57131302**
					COC	
Eubacterium saburreum		96.23	0.40	RHA 1	1	
Atopobium parvulum		3.77	0.06		99	
57131310	**Good identification**					**57131310**
Eubacterium saburreum		100.0	0.51	RHA 1		
57131312	**Good identification**					**57131312**
Eubacterium saburreum		100.0	0.38	RHA 1		
57131320	**Low discrimination**					**57131320**
					COC	
Eubacterium saburreum		75.24	0.40	RHA 1	1	
Clostridium perfringens		21.81	0.21		1	
Atopobium parvulum		2.95	0.06		99	
57131330	**Good identification**					**57131330**
Eubacterium saburreum		100.0	0.38	RHA 1		
57131400	**Good identification**					**57131400**
Eubacterium saburreum		100.0	0.46	ARA 1		
57131410	**Good identification**					**57131410**
Eubacterium saburreum		100.0	0.45	ARA 1		
57131500	**Good identification**					**57131500**
					COC	
Eubacterium saburreum		95.88	0.53	ARA 1	1	
Atopobium parvulum		3.76	0.19		99	
Bifidobacterium longum subsp. longum		0.37	0.07		1	
57131502	**Acceptable identification**					**57131502**
					COC	
Eubacterium saburreum		91.93	0.40	ARA 1	1	
Atopobium parvulum		3.60	0.06		99	
Bifidobacterium adolescentis		2.49	0.11		1	
57131510	**Good identification**					**57131510**
Eubacterium saburreum		99.21	0.51	ARA 1		
Bifidobacterium longum subsp. longum		0.79	0.11			
57131512	**Acceptable identification**					**57131512**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		92.14	0.38	ARA 1		
Bifidobacterium longum subsp. longum		4.14	0.11			
Bifidobacterium adolescentis		3.72	0.12			
57131520	**Low discrimination**					**57131520**
Eubacterium saburreum		68.49	0.40	ARA 1		
Clostridium perfringens		19.85	0.21			
Bifidobacterium longum subsp. longum		5.92	0.18			
57131530	**Low discrimination**					**57131530**
Eubacterium saburreum		84.76	0.38	ARA 1		
Bifidobacterium longum subsp. longum		15.25	0.22			
57131722	**Good identification**					**57131722**
Bacteroides thetaiotaomicron		99.63	0.37	RAF 99		
Parabacteroides distasonis		0.37	0.11	XYL 96		
57133000	**Good identification**					**57133000**
					CAT	
Eubacterium saburreum		95.54	0.46	MAN 1	1	
Propionibacterium granulosum		4.46	0.22		90	
57133010	**Good identification**					**57133010**
Eubacterium saburreum		100.0	0.45	MAN 1		
57133100	**Good identification**					**57133100**
					COC	
Eubacterium saburreum		96.23	0.53	MAN 1	1	
Atopobium parvulum		3.77	0.19		99	
57133102	**Acceptable identification**					**57133102**
					COC	
Eubacterium saburreum		94.39	0.40	MAN 1	1	
Atopobium parvulum		3.70	0.06		99	
Propionibacterium propionicum		1.92	0.14		1	
57133110	**Good identification**					**57133110**
Eubacterium saburreum		100.0	0.51	MAN 1		
57133112	**Good identification**					**57133112**
Eubacterium saburreum		100.0	0.38	MAN 1		
57133120	**Low discrimination**					**57133120**
					COC	
Eubacterium saburreum		75.24	0.40	MAN 1	1	
Clostridium perfringens		21.81	0.21		1	
Atopobium parvulum		2.95	0.06		99	
57133130	**Good identification**					**57133130**
Eubacterium saburreum		100.0	0.38	MAN 1		
57134000	**Good identification**					**57134000**
Eubacterium saburreum		100.0	0.42	GAL 99		
57134010	**Good identification**					**57134010**
Eubacterium saburreum		100.0	0.40	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57134100	**Low discrimination**					**57134100**
					COC	
Eubacterium saburreum		88.10	0.48	GAL 99	1	
Atopobium parvulum		11.90	0.23		99	
57134102	**Low discrimination**					**57134102**
					COC	
Eubacterium saburreum		88.10	0.35	GAL 99	1	
Atopobium parvulum		11.90	0.10		99	
57134110	**Good identification**					**57134110**
Eubacterium saburreum		100.0	0.46	GAL 99		
57134120	**Low discrimination**					**57134120**
					COC	
Eubacterium saburreum		88.10	0.35	GAL 99	1	
Atopobium parvulum		11.90	0.10		99	
57135000	**Very good identification**					**57135000**
					COC	
Eubacterium saburreum		99.68	0.76		1	
Atopobium parvulum		0.32	0.23		99	
57135001	**Good identification**					**57135001**
Eubacterium saburreum		100.0	0.42	URE 1		
57135002	**Very good identification**					**57135002**
					COC	
Eubacterium saburreum		99.68	0.63		1	
Atopobium parvulum		0.32	0.10		99	
57135010	**Very good identification**					**57135010**
Eubacterium saburreum		100.0	0.74			
57135011	**Good identification**					**57135011**
Eubacterium saburreum		100.0	0.40	URE 1		
57135012	**Very good identification**					**57135012**
Eubacterium saburreum		100.0	0.61			
57135020	**Very good identification**					**57135020**
					COC	
Eubacterium saburreum		99.51	0.63		1	
Atopobium parvulum		0.32	0.10		99	
Clostridium perfringens		0.17	0.07		1	
57135022	**Good identification**					**57135022**
Eubacterium saburreum		99.54	0.50			
Clostridium perfringens		0.46	0.01			
57135030	**Very good identification**					**57135030**
Eubacterium saburreum		100.0	0.61			
57135032	**Good identification**					**57135032**
Eubacterium saburreum		100.0	0.49			
57135040	**Good identification**					**57135040**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.42	SOR 1			
57135050	**Good identification**						**57135050**
Eubacterium saburreum		100.0	0.40	SOR 1			
57135100	**Low discrimination**						**57135100**
					COC	GRA	
Eubacterium saburreum		88.09	0.82		1	85	
Atopobium parvulum		11.90	0.57		99	99	
Fusobacterium mortiferum		0.01	0.12		1	1	
57135101	**Low discrimination**						**57135101**
					COC	SPO	
Eubacterium saburreum		85.75	0.48	URE 1	1	1	
Atopobium parvulum		11.59	0.23		99	1	
Clostridium chauvoei		2.67	0.12		1	99	
57135102	**Low discrimination**						**57135102**
					COC		
Eubacterium saburreum		88.06	0.69		1		
Atopobium parvulum		11.90	0.45		99		
Bifidobacterium adolescentis		0.05	0.11		1		
57135103	**Low discrimination**						**57135103**
					COC	SPO	
Eubacterium saburreum		85.75	0.35	URE 1	1	1	
Atopobium parvulum		11.59	0.10		99	1	
Clostridium chauvoei		2.67	-.01		1	99	
57135110	**Very good identification**						**57135110**
					COC		
Eubacterium saburreum		99.83	0.81		1		
Atopobium parvulum		0.17	0.23		99		
57135111	**Good identification**						**57135111**
Eubacterium saburreum		100.0	0.46	URE 1			
57135112	**Very good identification**						**57135112**
					COC		
Eubacterium saburreum		99.76	0.68		1		
Atopobium parvulum		0.17	0.10		99		
Bifidobacterium adolescentis		0.08	0.12		1		
57135120	**Low discrimination**						**57135120**
					COC		
Eubacterium saburreum		87.68	0.69		1		
Atopobium parvulum		11.85	0.45		99		
Clostridium perfringens		0.48	0.21		1		
57135121	**Low discrimination**						**57135121**
					COC	SPO	
Eubacterium saburreum		83.76	0.35	URE 1	1	1	
Atopobium parvulum		11.32	0.10		99	1	
Clostridium chauvoei		4.92	0.04		1	99	
57135122	**Low discrimination**						**57135122**
					COC		
Eubacterium saburreum		84.51	0.56		1		
Atopobium parvulum		11.42	0.32		99		
Clostridium perfringens		1.23	0.16		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57135130	**Very good identification**						**57135130**
					COC		
Eubacterium saburreum		99.83	0.68		1		
Atopobium parvulum		0.17	0.10		99		
57135132	**Good identification**						**57135132**
					CAT		
Eubacterium saburreum		98.79	0.55		1		
Parabacteroides distasonis		0.70	0.18		76		
Bacteroides thetaiotaomicron		0.50	0.00		89		
57135140	**Low discrimination**						**57135140**
					COC		
Eubacterium saburreum		88.10	0.48	SOR 1	1		
Atopobium parvulum		11.90	0.23		99		
57135142	**Low discrimination**						**57135142**
					COC		
Eubacterium saburreum		88.10	0.35	SOR 1	1		
Atopobium parvulum		11.90	0.10		99		
57135150	**Good identification**						**57135150**
Eubacterium saburreum		100.0	0.46	SOR 1			
57135160	**Low discrimination**						**57135160**
					COC		
Eubacterium saburreum		80.80	0.35	SOR 1	1		
Atopobium parvulum		10.92	0.10		99		
Clostridium perfringens		8.29	0.09		1		
57135200	**Good identification**						**57135200**
Eubacterium saburreum		100.0	0.42	RHA 1			
57135210	**Good identification**						**57135210**
Eubacterium saburreum		100.0	0.40	RHA 1			
57135300	**Low discrimination**						**57135300**
					COC		
Eubacterium saburreum		88.10	0.48	RHA 1	1		
Atopobium parvulum		11.90	0.23		99		
57135302	**Low discrimination**						**57135302**
					COC		
Eubacterium saburreum		88.10	0.35	RHA 1	1		
Atopobium parvulum		11.90	0.10		99		
57135310	**Good identification**						**57135310**
Eubacterium saburreum		100.0	0.46	RHA 1			
57135320	**Low discrimination**						**57135320**
					COC	GRA	
Eubacterium saburreum		82.99	0.35	RHA 1	1	85	
Atopobium parvulum		11.21	0.10		99	99	
Bacteroides vulgatus		5.80	-.03		1	1	
57135400	**Good identification**						**57135400**
Eubacterium saburreum		100.0	0.42	ARA 1			
57135410	**Good identification**						**57135410**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.40	ARA 1			
57135500	**Low discrimination**						**57135500**
					COC		
Eubacterium saburreum		86.48	0.48	ARA 1	1		
Atopobium parvulum		11.68	0.23		99		
Bifidobacterium longum subsp. longum		1.84	0.15		1		
57135502	**Low discrimination**						**57135502**
					CAT	GRA	
Eubacterium saburreum		56.21	0.35	ARA 1	1	85	
Bifidobacterium adolescentis		18.88	0.25		1	99	
Bacteroides thetaiotaomicron		10.55	0.07		89	1	
57135510	**Good identification**						**57135510**
Eubacterium saburreum		95.77	0.46	ARA 1			
Bifidobacterium longum subsp. longum		4.23	0.19				
57135520	**Low discrimination**						**57135520**
					CAT	GRA	
Eubacterium saburreum		52.42	0.35	ARA 1	1	85	
Bifidobacterium longum subsp. longum		25.25	0.26		1	99	
Bacteroides thetaiotaomicron		9.84	0.07		89	1	
57135522	**Good identification**						**57135522**
					CAT	GRA	
Bacteroides thetaiotaomicron		95.82	0.41	RAF 99	89	1	
Bifidobacterium longum subsp. longum		2.48	0.26	RHA 93	1	99	
Eubacterium saburreum		0.91	0.22		1	85	
57135722	**Very good identification**						**57135722**
Bacteroides thetaiotaomicron		99.54	0.60	RAF 99			
Bacteroides ovatus		0.29	0.17				
Parabacteroides distasonis		0.14	0.27				
57135732	**Good identification**						**57135732**
Bacteroides thetaiotaomicron		98.97	0.53	RAF 99			
Bacteroides ovatus		0.49	0.14				
Clostridium clostridioforme		0.35	0.22				
57137000	**Good identification**						**57137000**
Eubacterium saburreum		100.0	0.42	MAN 1			
57137010	**Good identification**						**57137010**
Eubacterium saburreum		100.0	0.40	MAN 1			
57137100	**Low discrimination**						**57137100**
					COC		
Eubacterium saburreum		87.52	0.48	MAN 1	1		
Atopobium parvulum		11.83	0.23		99		
Actinomyces israelii		0.66	0.16		1		
57137102	**Low discrimination**						**57137102**
					COC		
Eubacterium saburreum		80.65	0.35	MAN 1	1		
Atopobium parvulum		10.90	0.10		99		
Actinomyces israelii		5.14	0.19		1		
57137110	**Good identification**						**57137110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.46	MAN 1		
57137120	**Low discrimination**					**57137120**
					COC	
Eubacterium saburreum		88.10	0.35	MAN 1	1	
Atopobium parvulum		11.90	0.10		99	
57141000	**Good identification**					**57141000**
					SPO	
Eubacterium saburreum		97.13	0.41	CEL 1	1	
Lactobacillus jensenii		1.47	0.02		1	
Clostridium bifermentans		1.40	-.03		99	
57141010	**Good identification**					**57141010**
Eubacterium saburreum		100.0	0.40	CEL 1		
57141100	**Good identification**					**57141100**
Eubacterium saburreum		97.37	0.47	CEL 1		
Collinsella aerofaciens		2.63	-.03			
57141110	**Good identification**					**57141110**
Eubacterium saburreum		100.0	0.46	CEL 1		
57141722	**Good identification**					**57141722**
Bacteroides eggerthii		100.0	0.47	SUC 1		
57145000	**Good identification**					**57145000**
Eubacterium saburreum		100.0	0.37	CEL 1		
57145010	**Good identification**					**57145010**
Eubacterium saburreum		100.0	0.35	CEL 1		
57145100	**Good identification**					**57145100**
Eubacterium saburreum		100.0	0.43	CEL 1		
57145110	**Good identification**					**57145110**
Eubacterium saburreum		100.0	0.41	CEL 1		
57145722	**Very good identification**					**57145722**
					CAT	
Bacteroides eggerthii		99.78	0.66	SUC 1	93	
Bacteroides uniformis		0.18	0.22		5	
Bacteroides thetaiotaomicron		0.03	0.13		89	
57151000	**Low discrimination**					**57151000**
Eubacterium saburreum		53.26	0.49	CEL 1		
Lactobacillus jensenii		26.57	0.36			
Lactobacillus catenaformis		10.16	0.29			
57151010	**Low discrimination**					**57151010**
Eubacterium saburreum		83.11	0.48	CEL 1		
Lactobacillus jensenii		16.89	0.28			
57151020	**Low discrimination**					**57151020**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium saburreum		38.06	0.36	CEL 1	1			
Lactobacillus jensenii		26.90	0.26		1			
Clostridium barati		13.84	0.15		99			
57151100	**Low discrimination**							**57151100**
Eubacterium saburreum		46.34	0.56	CEL 1				
Collinsella aerofaciens		41.34	0.32					
Lactobacillus catenaformis		5.68	0.32					
57151102	**Low discrimination**							**57151102**
Eubacterium saburreum		28.56	0.43	CEL 1				
Lactobacillus catenaformis		19.85	0.32					
Collinsella aerofaciens		16.05	0.15					
57151110	**Good identification**							**57151110**
Eubacterium saburreum		98.91	0.54	CEL 1				
Collinsella aerofaciens		1.09	-.03					
57151112	**Acceptable identification**							**57151112**
					CAT			
Eubacterium saburreum		92.61	0.41	CEL 1	1			
Leptotrichia buccalis		4.38	-.03		1			
Capnocytophaga ochracea		3.01	0.04		90			
57151122	**Low discrimination**							**57151122**
					GRA	CAT	SPO	
Prevotella oralis		50.04	0.43	IND 1	1	1	1	
Capnocytophaga ochracea		18.23	0.36	RAF 92	1	90	1	
Clostridium barati		11.89	0.32		99	1	99	
57151130	**Acceptable identification**							**57151130**
					SPO	CAT		
Eubacterium saburreum		91.99	0.41	CEL 1	1	1		
Clostridium paraputrificum		3.21	-.03		99	1		
Capnocytophaga ochracea		2.99	0.04		1	90		
57151322	**Acceptable identification**							**57151322**
					GRA	SPO		
Prevotella oralis		92.01	0.37	IND 1	1	1		
Clostridium barati		4.44	0.14	RAF 92	99	99		
Bacteroides uniformis		2.33	0.04		1	1		
57155000	**Good identification**							**57155000**
Eubacterium saburreum		99.07	0.45	CEL 1				
Lactobacillus jensenii		0.93	0.02					
57155010	**Good identification**							**57155010**
Eubacterium saburreum		100.0	0.43	CEL 1				
57155100	**Low discrimination**							**57155100**
					COC			
Eubacterium saburreum		81.18	0.51	CEL 1	1			
Atopobium parvulum		17.06	0.30		99			
Collinsella aerofaciens		1.36	-.03		1			
57155102	**Low discrimination**							**57155102**
					COC	CAT		
Eubacterium saburreum		60.82	0.38	CEL 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		12.78	0.17		99	1	
Capnocytophaga ochracea		6.75	0.10		1	90	
57155110	**Good identification**						**57155110**
Eubacterium saburreum		100.0	0.49	CEL 1			
57155112	**Low discrimination**						**57155112**
Eubacterium saburreum		86.26	0.37	CEL 1			
Bifidobacterium adolescentis		13.74	0.21				
57155120	**Low discrimination**						**57155120**
					CAT	COC	
Eubacterium saburreum		57.75	0.38	CEL 1	1	1	
Atopobium parvulum		12.14	0.17		1	99	
Bacteroides fragilis		9.84	0.04		99	1	
57155122	**Low discrimination**						**57155122**
					CAT		
Bacteroides uniformis		59.25	0.35	SAL 92	5		
Bacteroides fragilis		22.53	0.25	RAF 97	99		
Prevotella oralis		9.43	0.28	ARA 92	1		
57155130	**Good identification**						**57155130**
Eubacterium saburreum		100.0	0.37	CEL 1			
57155322	**Acceptable identification**						**57155322**
					CAT		
Bacteroides uniformis		91.11	0.38	SAL 92	5		
Prevotella oralis		4.50	0.22	RAF 97	1		
Bacteroides thetaiotaomicron		2.06	0.13	ARA 92	89		
57155522	**Good identification**						**57155522**
					CAT		
Bacteroides uniformis		97.50	0.54	SAL 92	5		
Bacteroides thetaiotaomicron		1.97	0.27	RAF 97	89		
Bacteroides ovatus		0.25	0.12		68		
57155722	**Low discrimination**						**57155722**
					CAT		
Bacteroides uniformis		77.66	0.56	SAL 92	5		
Bacteroides thetaiotaomicron		15.12	0.47	RAF 97	89		
Bacteroides eggerthii		4.50	0.32		93		
57155732	**Low discrimination**						**57155732**
					CAT		
Bacteroides thetaiotaomicron		68.71	0.40	RAF 99	89		
Bacteroides ovatus		20.91	0.31	TRE 93	68		
Bacteroides uniformis		9.17	0.22		5		
57161000	**Acceptable identification**						**57161000**
Eubacterium saburreum		93.63	0.38	CEL 1			
Lactobacillus jensenii		6.37	0.10				
57161010	**Good identification**						**57161010**
Eubacterium saburreum		97.30	0.37	CEL 1			
Lactobacillus jensenii		2.70	0.02				
57161100	**Low discrimination**						**57161100**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		78.48	0.44	CEL 1	1		
Atopobium parvulum		21.52	0.25		99		
57161110	**Good identification**						**57161110**
Eubacterium saburreum		100.0	0.43	CEL 1			
57165100	**Low discrimination**						**57165100**
					COC		
Eubacterium saburreum		51.39	0.40	CEL 1	1		
Atopobium parvulum		48.61	0.30		99		
57165110	**Good identification**						**57165110**
Eubacterium saburreum		100.0	0.38	CEL 1			
57171000	**Low discrimination**						**57171000**
					COC		
Lactobacillus jensenii		64.63	0.44	IND 1	1		
Eubacterium saburreum		28.79	0.46	SAL 95	1		
Atopobium parvulum		6.14	0.25		99		
57171002	**Low discrimination**						**57171002**
					COC		
Lactobacillus jensenii		85.87	0.41	IND 1	1		
Eubacterium saburreum		10.13	0.33	SAL 95	1		
Atopobium parvulum		2.16	0.12		99		
57171010	**Low discrimination**						**57171010**
Eubacterium saburreum		52.23	0.45	CEL 1			
Lactobacillus jensenii		47.77	0.36				
57171100	**Low discrimination**						**57171100**
					COC	GRA	
Atopobium parvulum		89.61	0.59	IND 1	99	99	
Eubacterium saburreum		9.90	0.53		1	85	
Fusobacterium mortiferum		0.16	0.23		1	1	
57171102	**Low discrimination**						**57171102**
					COC		
Atopobium parvulum		82.91	0.46	IND 1	99		
Eubacterium saburreum		9.16	0.40		1		
Lactobacillus acidophilus		6.07	0.25		1		
57171110	**Low discrimination**						**57171110**
					COC	GRA	
Eubacterium saburreum		88.64	0.51	CEL 1	1	85	
Atopobium parvulum		9.90	0.25		99	99	
Leptotrichia buccalis		1.11	-.03		1	1	
57171120	**Low discrimination**						**57171120**
					COC	SPO	
Atopobium parvulum		68.43	0.46	IND 1	99	1	
Clostridium perfringens		9.04	0.32		1	84	
Clostridium paraputrificum		8.54	0.22		1	99	
57171130	**Low discrimination**						**57171130**
					COC		
Eubacterium saburreum		89.95	0.38	CEL 1	1		
Atopobium parvulum		10.05	0.12		99		
57171722	**Good identification**						**57171722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		96.83	0.42	RAF 99			
Bacteroides ovatus		3.17	0.17	XYL 96			
57171732	**Acceptable identification**						**57171732**
Bacteroides thetaiotaomicron		93.52	0.35	RAF 99			
Bacteroides ovatus		5.25	0.14	XYL 96			
Clostridium clostridioforme		1.23	0.13				
57175000	**Low discrimination**						**57175000**
					COC		
Eubacterium saburreum		56.25	0.42	CEL 1	1		
Atopobium parvulum		41.38	0.30		99		
Lactobacillus jensenii		2.37	0.10		1		
57175010	**Good identification**						**57175010**
Eubacterium saburreum		98.31	0.40	CEL 1			
Lactobacillus jensenii		1.69	0.02				
57175100	**Very good identification**						**57175100**
					COC	SPO	
Atopobium parvulum		96.78	0.64	IND 1	99	1	
Eubacterium saburreum		3.10	0.48		1	1	
Clostridium butyricum		0.06	0.02		1	99	
57175102	**Good identification**						**57175102**
					COC		
Atopobium parvulum		96.00	0.51	IND 1	99		
Eubacterium saburreum		3.08	0.35		1		
Actinomyces israelii		0.46	0.25		1		
57175110	**Low discrimination**						**57175110**
					COC		
Eubacterium saburreum		72.19	0.46	CEL 1	1		
Atopobium parvulum		27.82	0.30		99		
57175120	**Good identification**						**57175120**
					COC		
Atopobium parvulum		96.90	0.51	IND 1	99		
Eubacterium saburreum		3.11	0.35		1		
57175122	**Low discrimination**						**57175122**
					COC	CAT	
Atopobium parvulum		83.35	0.38	IND 1	99	1	
Bacteroides thetaiotaomicron		6.04	0.13		1	89	
Eubacterium saburreum		2.67	0.22		1	1	
57175522	**Low discrimination**						**57175522**
					CAT		
Bacteroides thetaiotaomicron		87.97	0.47	RAF 99	89		
Bacteroides ovatus		8.31	0.29	RHA 93	68		
Bacteroides uniformis		3.31	0.19		5		
57175532	**Low discrimination**						**57175532**
Bacteroides thetaiotaomicron		85.17	0.40	RAF 99			
Bacteroides ovatus		13.79	0.26	RHA 93			
Clostridium clostridioforme		0.86	0.16				
57175622	**Acceptable identification**						**57175622**
Bacteroides thetaiotaomicron		92.79	0.37	RAF 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		6.20	0.17	LAC 98			
Clostridium clostridioforme		0.64	0.10				
57175722	**Low discrimination**						**57175722**
					CAT		
Bacteroides thetaiotaomicron		87.72	0.66	RAF 99	89		
Bacteroides ovatus		11.85	0.51		68		
Bacteroides uniformis		0.34	0.22		5		
57175732	**Low discrimination**						**57175732**
Bacteroides thetaiotaomicron		80.80	0.59	RAF 99			
Bacteroides ovatus		18.71	0.48				
Clostridium clostridioforme		0.45	0.31				
57177102	**Low discrimination**						**57177102**
					COC	SPO	
Actinomyces israelii		49.00	0.38	IND 1	1	1	
Clostridium tertium		27.70	0.19	SAL 92	1	99	
Atopobium parvulum		18.25	0.17		99	1	
57177502	**Low discrimination**						**57177502**
					SPO		
Actinomyces israelii		52.46	0.36	IND 1	1		
Bifidobacterium adolescentis		44.24	0.29	SAL 92	1		
Clostridium butyricum		3.31	-.05		99		
57177722	**Low discrimination**						**57177722**
Bacteroides ovatus		81.68	0.43	SAL 88			
Bacteroides thetaiotaomicron		18.32	0.32	RAF 99			
57177732	**Low discrimination**						**57177732**
Bacteroides ovatus		88.43	0.40	SAL 88			
Bacteroides thetaiotaomicron		11.57	0.25	RAF 99			
57200000	**Good identification**						**57200000**
					GRA	SPO	
Clostridium bifermentans		99.19	0.43	SUC 1	99	99	
Fusobacterium mortiferum		0.49	0.17	SAL 10	1	1	
Eubacterium tortuosum		0.32	-.03		99	1	
57201100	**Low discrimination**						**57201100**
					GRA		
Eubacterium saburreum		80.88	0.37	SAL 1	85		
Fusobacterium mortiferum		16.13	0.27		1		
Eubacterium tortuosum		2.99	-.03		99		
57201110	**Good identification**						**57201110**
Eubacterium saburreum		100.0	0.36	SAL 1			
57210000	**Low discrimination**						**57210000**
					GRA	SPO	
Fusobacterium mortiferum		89.55	0.52	IND 1	1	1	
Clostridium bifermentans		9.62	0.21		99	99	
Fusobacterium varium		0.84	-.03		1	1	
57210002	**Good identification**						**57210002**
Fusobacterium mortiferum		100.0	0.35	IND 1			
57210100	**Very good identification**						**57210100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		99.77	0.58	IND 1			
Eubacterium saburreum		0.23	0.11				
57210102	**Good identification**						**57210102**
Fusobacterium mortiferum		100.0	0.42	IND 1			
57211000	**Low discrimination**						**57211000**
					GRA		
Fusobacterium mortiferum		84.38	0.55	IND 1	1		
Eubacterium saburreum		12.82	0.39		85		
Lactobacillus jensenii		1.35	0.14		99		
57211002	**Low discrimination**						**57211002**
					GRA		
Fusobacterium mortiferum		65.58	0.38	IND 1	1		
Eubacterium saburreum		15.82	0.26	BGL 10	85		
Lactobacillus catenaformis		7.60	0.12		99		
57211010	**Low discrimination**						**57211010**
					GRA		
Eubacterium saburreum		88.96	0.37	SAL 1	85		
Fusobacterium mortiferum		7.23	0.20		1		
Lactobacillus jensenii		3.81	0.06		99		
57211100	**Low discrimination**						**57211100**
					GRA		
Fusobacterium mortiferum		85.54	0.61	IND 1	1		
Eubacterium saburreum		13.00	0.45		85		
Lactobacillus catenaformis		0.71	0.15		99		
57211102	**Low discrimination**						**57211102**
					SPO	GRA	
Fusobacterium mortiferum		68.33	0.45	IND 1	1	1	
Eubacterium saburreum		16.49	0.32	BGL 10	1	85	
Clostridium barati		6.31	0.12		99	99	
57211110	**Acceptable identification**						**57211110**
Eubacterium saburreum		92.48	0.44	SAL 1			
Fusobacterium mortiferum		7.52	0.27				
57214100	**Good identification**						**57214100**
Fusobacterium mortiferum		100.0	0.39	IND 1			
57215100	**Low discrimination**						**57215100**
Eubacterium saburreum		52.08	0.41	SAL 1			
Fusobacterium mortiferum		47.92	0.42				
57215110	**Good identification**						**57215110**
Eubacterium saburreum		100.0	0.39	SAL 1			
57230000	**Good identification**						**57230000**
Fusobacterium mortiferum		100.0	0.43	IND 1			
57230100	**Good identification**						**57230100**
Fusobacterium mortiferum		100.0	0.49	IND 1			
57231000	**Low discrimination**						**57231000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Fusobacterium mortiferum		66.70	0.46	IND 1	1		
Eubacterium saburreum		22.62	0.36		85		
Lactobacillus jensenii		10.69	0.22		99		
57231100	**Low discrimination**						**57231100**
					COC	GRA	
Fusobacterium mortiferum		70.95	0.52	IND 1	1	1	
Eubacterium saburreum		24.06	0.42		1	85	
Atopobium parvulum		5.00	0.21		99	99	
57231102	**Low discrimination**						**57231102**
					COC	GRA	
Fusobacterium mortiferum		57.42	0.36	IND 1	1	1	
Eubacterium saburreum		30.92	0.29	BGL 10	1	85	
Atopobium parvulum		6.42	0.08		99	99	
57231110	**Good identification**						**57231110**
Eubacterium saburreum		96.49	0.41	SAL 1			
Fusobacterium mortiferum		3.51	0.18				
57235100	**Low discrimination**						**57235100**
					COC	GRA	
Eubacterium saburreum		46.98	0.38	SAL 1	1	85	
Atopobium parvulum		33.66	0.25		99	99	
Fusobacterium mortiferum		19.37	0.33		1	1	
57235110	**Good identification**						**57235110**
Eubacterium saburreum		100.0	0.36	SAL 1			
57250000	**Low discrimination**						**57250000**
					GRA	SPO	
Fusobacterium mortiferum		86.98	0.43	IND 1	1	1	
Clostridium barati		5.43	0.08		99	99	
Lactobacillus catenaformis		5.04	0.12		99	1	
57250100	**Low discrimination**						**57250100**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.50	IND 1	1	1	
Clostridium barati		7.03	0.16		99	99	
Lactobacillus catenaformis		3.26	0.15		99	1	
57251000	**Low discrimination**						**57251000**
					GRA		
Lactobacillus catenaformis		56.58	0.47	IND 1	99		
Fusobacterium mortiferum		14.79	0.46	MLT 90	1		
Lactobacillus jensenii		13.46	0.36		99		
57251002	**Low discrimination**						**57251002**
					SPO		
Lactobacillus catenaformis		53.38	0.47	IND 1	1		
Clostridium barati		33.10	0.38	MLT 90	99		
Lactobacillus jensenii		8.46	0.33		1		
57251020	**Low discrimination**						**57251020**
					SPO		
Clostridium barati		66.13	0.38	IND 1	99		
Lactobacillus catenaformis		26.66	0.36		1		
Lactobacillus jensenii		6.34	0.26		1		
57251022	**Low discrimination**						**57251022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium barati		85.87	0.46	IND 1	99		
Lactobacillus catenaformis		11.54	0.36		1		
Lactobacillus jensenii		1.83	0.23		1		
57251100	**Low discrimination**						**57251100**
					SPO		
Lactobacillus catenaformis		40.18	0.50	IND 1	1		
Collinsella aerofaciens		26.58	0.32	MLT 90	1		
Clostridium barati		16.61	0.38		99		
57251102	**Low discrimination**						**57251102**
					SPO		
Clostridium barati		50.22	0.46	IND 1	99		
Lactobacillus catenaformis		40.49	0.50		1		
Lactobacillus acidophilus		4.42	0.30		1		
57251120	**Low discrimination**						**57251120**
					SPO		
Clostridium barati		63.82	0.46	IND 1	99		
Clostridium paraputrificum		19.06	0.32		99		
Lactobacillus catenaformis		12.86	0.39		1		
57251122	**Low discrimination**						**57251122**
					SPO		
Clostridium barati		87.17	0.54	IND 1	99		
Lactobacillus catenaformis		5.86	0.39		1		
Clostridium paraputrificum		3.72	0.25		99		
57251162	**Acceptable identification**						**57251162**
					SPO		
Clostridium barati		91.56	0.36	IND 1	99		
Lactobacillus acidophilus		3.82	0.15	SOR 8	1		
Clostridium paraputrificum		3.38	0.06		99		
57251322	**Acceptable identification**						**57251322**
					GRA	SPO	
Clostridium barati		92.39	0.36	IND 1	99	99	
Bacteroides uniformis		2.39	0.04	RHA 8	1	1	
Lactobacillus acidophilus		2.35	0.11		99	1	
57253000	**Good identification**						**57253000**
					SPO		
Clostridium innocuum		99.35	0.56	IND 1	99		
Clostridium barati		0.35	0.11		99		
Lactobacillus catenaformis		0.20	0.12		1		
57253002	**Good identification**						**57253002**
					SPO		
Clostridium innocuum		98.76	0.56	IND 1	99		
Clostridium barati		1.05	0.20		99		
Lactobacillus catenaformis		0.20	0.12		1		
57253122	**Acceptable identification**						**57253122**
					SPO		
Clostridium barati		94.29	0.36	IND 1	99		
Lactobacillus acidophilus		1.67	0.08	MAN 8	1		
Clostridium tertium		1.58	0.09		99		
57253400	**Good identification**						**57253400**
Clostridium innocuum		100.0	0.43	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
57253402	**Good identification**							**57253402**
Clostridium innocuum		100.0	0.43	IND 1				
57255122	**Acceptable identification**							**57255122**
					GRA	SPO		
Bacteroides uniformis		93.25	0.35	MLT 92	1	1		
Clostridium barati		5.84	0.20	RAF 97	99	99		
Lactobacillus catenaformis		0.39	0.05	ARA 92	99	1		
57255322	**Good identification**							**57255322**
					GRA	SPO		
Bacteroides uniformis		99.61	0.38	MLT 92	1	1		
Clostridium barati		0.39	0.02	RAF 97	99	99		
57255522	**Good identification**							**57255522**
					CAT			
Bacteroides uniformis		99.98	0.54	MLT 92	5			
Bacteroides thetaiotaomicron		0.02	-.06	RAF 97	89			
57255722	**Good identification**							**57255722**
					CAT			
Bacteroides uniformis		99.53	0.56	MLT 92	5			
Bacteroides ovatus		0.26	0.15	RAF 97	68			
Bacteroides thetaiotaomicron		0.22	0.13		89			
57270100	**Good identification**							**57270100**
					GRA	SPO		
Fusobacterium mortiferum		97.72	0.41	IND 1	1	1		
Clostridium barati		2.28	-.02		99	99		
57271000	**Low discrimination**							**57271000**
					GRA	SPO		
Lactobacillus jensenii		85.17	0.44	IND 1	99	1		
Fusobacterium mortiferum		9.32	0.38	MLT 95	1	1		
Clostridium innocuum		2.16	0.14		99	99		
57271002	**Low discrimination**							**57271002**
					SPO			
Lactobacillus jensenii		77.46	0.41	IND 1	1			
Clostridium barati		8.78	0.20	MLT 95	99			
Lactobacillus acidophilus		7.76	0.21		1			
57271010	**Good identification**							**57271010**
Lactobacillus jensenii		100.0	0.36	IND 1				
57271100	**Low discrimination**							**57271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		50.08	0.44	IND 1	1	1	1	
Atopobium parvulum		24.69	0.27		99	99	1	
Clostridium barati		14.82	0.20		1	99	99	
57271102	**Low discrimination**							**57271102**
					GRA	SPO		
Lactobacillus acidophilus		70.85	0.39	IND 1	99	1		
Clostridium barati		20.92	0.28	MLT 94	99	99		
Fusobacterium mortiferum		2.62	0.27		1	1		
57271122	**Low discrimination**							**57271122**
					SPO			
Lactobacillus acidophilus		49.22	0.39	IND 1	1			
Clostridium barati		43.60	0.36	MLT 94	99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium paraputrificum		5.69	0.16		99	
57273000	**Good identification**					**57273000**
					SPO	
Clostridium innocuum		99.60	0.48	IND 1	99	
Lactobacillus jensenii		0.40	0.10		1	
57273002	**Good identification**					**57273002**
					SPO	
Clostridium innocuum		99.33	0.48	IND 1	99	
Lactobacillus jensenii		0.27	0.07		1	
Clostridium barati		0.26	0.01		99	
57273400	**Good identification**					**57273400**
Clostridium innocuum		100.0	0.35	IND 1		
57273402	**Good identification**					**57273402**
Clostridium innocuum		100.0	0.35	IND 1		
57300000	**Good identification**					**57300000**
Clostridium bifermentans		100.0	0.50	SUC 1		
57301000	**Low discrimination**					**57301000**
					SPO	
Eubacterium saburreum		83.69	0.41	SAL 1	1	
Clostridium bifermentans		13.24	0.15		99	
Eubacterium tortuosum		1.80	-.03		1	
57301010	**Good identification**					**57301010**
Eubacterium saburreum		100.0	0.40	SAL 1		
57301100	**Good identification**					**57301100**
					GRA	
Eubacterium saburreum		96.79	0.47	SAL 1	85	
Collinsella aerofaciens		2.62	-.03		99	
Fusobacterium mortiferum		0.60	0.11		1	
57301110	**Good identification**					**57301110**
Eubacterium saburreum		100.0	0.46	SAL 1		
57305000	**Good identification**					**57305000**
Eubacterium saburreum		100.0	0.37	SAL 1		
57305010	**Good identification**					**57305010**
Eubacterium saburreum		100.0	0.35	SAL 1		
57305100	**Good identification**					**57305100**
Eubacterium saburreum		100.0	0.43	SAL 1		
57305110	**Good identification**					**57305110**
Eubacterium saburreum		100.0	0.41	SAL 1		
57310100	**Low discrimination**					**57310100**
					GRA	
Fusobacterium mortiferum		87.65	0.42	IND 1	1	
Eubacterium saburreum		6.53	0.21	MLT 11	85	
Collinsella aerofaciens		5.83	-.03		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57311000	**Low discrimination**						**57311000**
					GRA		
Eubacterium saburreum		48.87	0.49	SAL 1	85		
Lactobacillus jensenii		24.38	0.36		99		
Fusobacterium mortiferum		9.94	0.39		1		
57311010	**Low discrimination**						**57311010**
Eubacterium saburreum		83.11	0.48	SAL 1			
Lactobacillus jensenii		16.89	0.28				
57311020	**Low discrimination**						**57311020**
Eubacterium saburreum		30.99	0.36	SAL 1			
Clostridium perfringens		25.39	0.26				
Lactobacillus jensenii		21.91	0.26				
57311100	**Low discrimination**						**57311100**
					GRA		
Eubacterium saburreum		41.65	0.56	SAL 1	85		
Collinsella aerofaciens		37.17	0.32		99		
Fusobacterium mortiferum		8.47	0.45		1		
57311102	**Low discrimination**						**57311102**
Bifidobacterium longum subsp. infantis		63.63	0.40	IND 1			
Eubacterium saburreum		11.20	0.43	RAF 95			
Lactobacillus catenaformis		7.78	0.32				
57311110	**Good identification**						**57311110**
					GRA		
Eubacterium saburreum		98.67	0.54	SAL 1	85		
Collinsella aerofaciens		1.09	-.03		99		
Fusobacterium mortiferum		0.25	0.11		1		
57311112	**Low discrimination**						**57311112**
					GRA		
Eubacterium saburreum		86.84	0.41	SAL 1	85		
Bifidobacterium longum subsp. infantis		6.09	0.06		99		
Leptotrichia buccalis		4.11	-.03		1		
57311122	**Low discrimination**						**57311122**
					GRA	SPO	
Bifidobacterium longum subsp. infantis		57.51	0.44	IND 1	99	1	
Prevotella oralis		16.86	0.38	RAF 95	1	1	
Clostridium barati		7.99	0.32		99	99	
57311130	**Acceptable identification**						**57311130**
					SPO		
Eubacterium saburreum		93.74	0.41	SAL 1	1		
Clostridium paraputrificum		3.28	-.03		99		
Clostridium perfringens		2.98	0.07		84		
57315000	**Good identification**						**57315000**
					GRA		
Eubacterium saburreum		96.36	0.45	SAL 1	85		
Fusobacterium mortiferum		2.74	0.20		1		
Lactobacillus jensenii		0.90	0.02		99		
57315010	**Good identification**						**57315010**
Eubacterium saburreum		100.0	0.43	SAL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57315100	**Low discrimination**					**57315100**
Eubacterium saburreum		65.23	0.51	SAL 1		
Eubacterium contortum		14.21	0.29			
Bifidobacterium longum subsp. infantis		9.83	0.21			
57315102	**Low discrimination**					**57315102**
Bifidobacterium longum subsp. infantis		83.70	0.38	IND 1		
Eubacterium saburreum		9.70	0.38	RAF 95		
Bifidobacterium adolescentis		2.76	0.27			
57315110	**Good identification**					**57315110**
Eubacterium saburreum		100.0	0.49	SAL 1		
57315112	**Low discrimination**					**57315112**
Eubacterium saburreum		65.30	0.37	SAL 1		
Bifidobacterium adolescentis		27.74	0.28			
Bifidobacterium longum subsp. infantis		6.96	0.04			
57315122	**Low discrimination**					**57315122**
					GRA	
Bifidobacterium longum subsp. infantis		87.85	0.43	IND 1	99	
Bacteroides uniformis		5.66	0.19	RAF 95	1	
Prevotella oralis		3.89	0.23		1	
57315130	**Good identification**					**57315130**
Eubacterium saburreum		100.0	0.37	SAL 1		
57315300	**Very good identification**					**57315300**
Eubacterium contortum		99.95	0.63	IND 1		
Eubacterium saburreum		0.05	0.17			
57315302	**Good identification**					**57315302**
Eubacterium contortum		99.60	0.50	IND 1		
Bifidobacterium longum subsp. infantis		0.40	0.04			
57315320	**Good identification**					**57315320**
					GRA	
Eubacterium contortum		98.72	0.50	IND 1	99	
Bacteroides vulgatus		1.28	0.11		1	
57315322	**Low discrimination**					**57315322**
					GRA	
Eubacterium contortum		66.42	0.37	IND 1	99	
Bacteroides uniformis		23.83	0.22		1	
Prevotella oralis		5.08	0.16		1	
57315502	**Low discrimination**					**57315502**
					GRA	
Bifidobacterium adolescentis		83.35	0.41	IND 1	99	
Eubacterium contortum		7.81	0.14	RAF 92	99	
Bacteroides uniformis		5.03	0.03		1	
57315512	**Good identification**					**57315512**
Bifidobacterium adolescentis		98.70	0.42	IND 1		
Clostridium clostridioforme		1.31	0.07	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57315522	**Good identification**						**57315522**
					GRA	CAT	
Bacteroides uniformis		97.01	0.38	RAF 97	1	5	
Bacteroides thetaiotaomicron		1.40	0.09	CEL 99	1	89	
Bifidobacterium longum subsp. infantis		1.32	0.08		99	1	
57315700	**Very good identification**						**57315700**
					GRA		
Eubacterium contortum		99.96	0.61	IND 1	99		
Bacteroides vulgatus		0.04	-.05		1		
57315702	**Good identification**						**57315702**
					GRA		
Eubacterium contortum		98.71	0.48	IND 1	99		
Bacteroides uniformis		0.89	0.06		1		
Clostridium clostridioforme		0.28	0.10		50		
57315720	**Low discrimination**						**57315720**
					GRA		
Eubacterium contortum		81.92	0.48	IND 1	99		
Bacteroides vulgatus		17.24	0.30		1		
Bacteroides uniformis		0.74	0.06		1		
57315722	**Low discrimination**						**57315722**
					GRA	CAT	
Bacteroides uniformis		73.07	0.40	RAF 97	1	5	
Eubacterium contortum		14.49	0.36	CEL 99	99	1	
Bacteroides thetaiotaomicron		10.17	0.28		1	89	
57317502	**Acceptable identification**						**57317502**
Bifidobacterium adolescentis		93.62	0.39	IND 1			
Actinomyces israelii		6.39	0.25	RAF 92			
57317512	**Good identification**						**57317512**
Bifidobacterium adolescentis		100.0	0.41	IND 1			
57321000	**Acceptable identification**						**57321000**
Eubacterium saburreum		93.63	0.38	SAL 1			
Lactobacillus jensenii		6.37	0.10				
57321010	**Good identification**						**57321010**
Eubacterium saburreum		97.30	0.37	SAL 1			
Lactobacillus jensenii		2.70	0.02				
57321100	**Low discrimination**						**57321100**
					COC		
Eubacterium saburreum		86.52	0.44	SAL 1	1		
Atopobium parvulum		13.48	0.21		99		
57321110	**Good identification**						**57321110**
Eubacterium saburreum		100.0	0.43	SAL 1			
57325100	**Low discrimination**						**57325100**
					COC		
Eubacterium saburreum		65.05	0.40	SAL 1	1		
Atopobium parvulum		34.95	0.25		99		
57325110	**Good identification**						**57325110**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.38	SAL 1			
57331000	**Low discrimination**						**57331000**
					COC		
Lactobacillus jensenii		64.70	0.44	IND 1	1		
Eubacterium saburreum		28.82	0.46	CEL 95	1		
Atopobium parvulum		3.49	0.21		99		
57331002	**Low discrimination**						**57331002**
Lactobacillus jensenii		86.18	0.41	IND 1			
Eubacterium saburreum		10.16	0.33	CEL 95			
Lactobacillus acidophilus		1.37	0.07				
57331010	**Low discrimination**						**57331010**
Eubacterium saburreum		52.23	0.45	SAL 1			
Lactobacillus jensenii		47.77	0.36				
57331100	**Low discrimination**						**57331100**
					COC	GRA	
Atopobium parvulum		81.59	0.55	IND 1	99	99	
Eubacterium saburreum		15.87	0.53		1	85	
Fusobacterium mortiferum		1.45	0.36		1	1	
57331102	**Low discrimination**						**57331102**
					COC		
Atopobium parvulum		70.93	0.42	IND 1	99		
Eubacterium saburreum		13.80	0.40		1		
Lactobacillus acidophilus		9.14	0.25		1		
57331110	**Acceptable identification**						**57331110**
					COC	GRA	
Eubacterium saburreum		92.60	0.51	SAL 1	1	85	
Atopobium parvulum		5.88	0.21		99	99	
Leptotrichia buccalis		1.16	-.03		1	1	
57331120	**Low discrimination**						**57331120**
					COC	SPO	
Clostridium perfringens		46.44	0.45	IND 1	1	84	
Atopobium parvulum		33.28	0.42		99	1	
Clostridium paraputrificum		7.31	0.22		1	99	
57331122	**Low discrimination**						**57331122**
					COC		
Clostridium perfringens		53.26	0.40	IND 1	1		
Atopobium parvulum		14.31	0.29		99		
Lactobacillus acidophilus		10.46	0.25		1		
57331130	**Low discrimination**						**57331130**
					COC		
Eubacterium saburreum		86.80	0.38	SAL 1	1		
Clostridium perfringens		7.69	0.11		1		
Atopobium parvulum		5.51	0.08		99		
57335000	**Low discrimination**						**57335000**
					COC		
Eubacterium saburreum		67.90	0.42	SAL 1	1		
Atopobium parvulum		28.38	0.25		99		
Lactobacillus jensenii		2.86	0.10		1		
57335010	**Good identification**						**57335010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		98.31	0.40	SAL 1			
Lactobacillus jensenii		1.69	0.02				
57335100	**Good identification**						**57335100**
					COC	SPO	
Atopobium parvulum		94.42	0.60	IND 1	99	1	
Eubacterium saburreum		5.33	0.48		1	1	
Clostridium butyricum		0.11	0.02		1	99	
57335102	**Acceptable identification**						**57335102**
					COC		
Atopobium parvulum		91.78	0.47	IND 1	99		
Eubacterium saburreum		5.18	0.35		1		
Bifidobacterium adolescentis		1.47	0.23		1		
57335110	**Low discrimination**						**57335110**
					COC		
Eubacterium saburreum		82.04	0.46	SAL 1	1		
Atopobium parvulum		17.96	0.25		99		
57335120	**Acceptable identification**						**57335120**
					COC		
Atopobium parvulum		93.99	0.47	IND 1	99		
Eubacterium saburreum		5.30	0.35		1		
Clostridium perfringens		0.71	0.11		1		
57335502	**Low discrimination**						**57335502**
					COC	SPO	
Bifidobacterium adolescentis		70.79	0.38	IND 1	1	1	
Clostridium butyricum		14.62	0.11	RAF 92	1	99	
Atopobium parvulum		6.66	0.13		99	1	
57335512	**Low discrimination**						**57335512**
Bifidobacterium adolescentis		87.30	0.39	IND 1			
Clostridium clostridioforme		12.70	0.22	RAF 92			
57335712	**Acceptable identification**						**57335712**
Clostridium clostridioforme		92.32	0.36	IND 1			
Bacteroides thetaiotaomicron		6.37	0.06	RAF 94			
Bacteroides ovatus		1.32	-.06				
57335722	**Low discrimination**						**57335722**
Bacteroides thetaiotaomicron		87.09	0.47	RAF 99			
Bacteroides ovatus		10.50	0.32				
Clostridium clostridioforme		1.09	0.25				
57335732	**Low discrimination**						**57335732**
Bacteroides thetaiotaomicron		73.26	0.40	RAF 99			
Bacteroides ovatus		15.13	0.29				
Clostridium clostridioforme		10.73	0.36				
57337102	**Low discrimination**						**57337102**
					SPO		
Actinomyces israelii		45.14	0.37	IND 1	1		
Clostridium tertium		29.48	0.19	CEL 93	99		
Bifidobacterium adolescentis		14.34	0.22		1		
57337502	**Low discrimination**						**57337502**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		70.77	0.36	IND 1	1	
Actinomyces israelii		27.24	0.35	RAF 92	1	
Clostridium butyricum		1.99	-.05		99	
57337512	**Low discrimination**					**57337512**
Bifidobacterium adolescentis		83.46	0.38	IND 1		
Bifidobacterium dentium		12.62	0.06	RAF 92		
Actinomyces israelii		3.93	0.21			
57345522	**Good identification**					**57345522**
					CAT	
Bacteroides uniformis		99.31	0.38	RAF 97	5	
Bacteroides eggerthii		0.69	-.03	FRU 99	93	
57345722	**Low discrimination**					**57345722**
					CAT	
Bacteroides uniformis		65.60	0.40	RAF 97	5	
Bacteroides eggerthii		32.76	0.32	FRU 99	93	
Bacteroides ovatus		1.45	0.15		68	
57350022	**Acceptable identification**					**57350022**
					SPO	
Clostridium barati		93.18	0.36	IND 1	99	
Lactobacillus catenaformis		4.43	0.18	GAL 95	1	
Lactobacillus jensenii		1.48	0.10		1	
57350102	**Low discrimination**					**57350102**
					SPO	
Clostridium barati		63.02	0.36	IND 1	99	
Lactobacillus catenaformis		17.98	0.32	GAL 95	1	
Collinsella aerofaciens		14.54	0.15		1	
57350122	**Low discrimination**					**57350122**
					SPO	
Clostridium barati		81.11	0.44	IND 1	99	
Clostridium paraputrificum		13.47	0.25	GAL 95	99	
Lactobacillus catenaformis		1.93	0.21		1	
57351000	**Low discrimination**					**57351000**
					SPO	
Lactobacillus catenaformis		57.82	0.63	IND 1	1	
Lactobacillus jensenii		29.03	0.58		1	
Clostridium barati		6.48	0.41		99	
57351002	**Low discrimination**					**57351002**
					SPO	
Lactobacillus catenaformis		58.49	0.63	IND 1	1	
Clostridium barati		19.68	0.50		99	
Lactobacillus jensenii		19.58	0.55		1	
57351010	**Acceptable identification**					**57351010**
					SPO	
Lactobacillus jensenii		92.90	0.50	IND 1	1	
Lactobacillus catenaformis		5.61	0.29		1	
Clostridium barati		0.63	0.07		99	
57351012	**Low discrimination**					**57351012**
					SPO	
Lactobacillus jensenii		88.43	0.47	IND 1	1	
Lactobacillus catenaformis		8.01	0.29		1	
Clostridium barati		2.69	0.15		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57351020	**Low discrimination**					**57351020**
					SPO	
Clostridium barati		42.74	0.50	IND 1	99	
Lactobacillus catenaformis		31.76	0.53		1	
Lactobacillus jensenii		15.94	0.47		1	
57351022	**Low discrimination**					**57351022**
					SPO	
Clostridium barati		71.56	0.58	IND 1	99	
Lactobacillus catenaformis		17.72	0.53		1	
Lactobacillus jensenii		5.93	0.44		1	
57351030	**Low discrimination**					**57351030**
					SPO	
Lactobacillus jensenii		86.47	0.39	IND 1	1	
Clostridium barati		7.02	0.15		99	
Lactobacillus catenaformis		5.22	0.18		1	
57351032	**Low discrimination**					**57351032**
					SPO	
Lactobacillus jensenii		67.30	0.36	IND 1	1	
Clostridium barati		24.60	0.24		99	
Lactobacillus catenaformis		6.09	0.18		1	
57351062	**Acceptable identification**					**57351062**
					SPO	
Clostridium barati		90.95	0.40	IND 1	99	
Lactobacillus acidophilus		3.17	0.17	SOR 8	1	
Lactobacillus catenaformis		2.62	0.18		1	
57351100	**Low discrimination**					**57351100**
					SPO	
Collinsella aerofaciens		84.99	0.66	IND 1	1	
Lactobacillus catenaformis		11.68	0.66		1	
Clostridium barati		2.62	0.50		99	
57351102	**Low discrimination**					**57351102**
					SPO	
Lactobacillus catenaformis		36.83	0.66	IND 1	1	
Collinsella aerofaciens		29.78	0.50		1	
Clostridium barati		24.78	0.58		99	
57351120	**Low discrimination**					**57351120**
					SPO	
Clostridium paraputrificum		69.86	0.66	IND 1	99	
Collinsella aerofaciens		13.86	0.50		1	
Clostridium barati		11.54	0.58		99	
57351122	**Low discrimination**					**57351122**
					SPO	
Clostridium barati		45.02	0.66	IND 1	99	
Clostridium paraputrificum		38.94	0.60		99	
Lactobacillus catenaformis		5.58	0.56		1	
57351132	**Low discrimination**					**57351132**
					SPO	
Bifidobacterium breve		28.26	0.44	IND 1	1	
Clostridium barati		27.93	0.32	RAF 95	99	
Clostridium paraputrificum		24.16	0.25		99	
57351142	**Low discrimination**					**57351142**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		51.69	0.40	IND 1	99		
Lactobacillus acidophilus		20.74	0.35	SOR 8	1		
Bifidobacterium breve		10.25	0.40		1		
57351152	**Low discrimination**						**57351152**
					GRA	SPO	
Bifidobacterium breve		89.30	0.40	IND 1	99	1	
Clostridium barati		4.55	0.05	RAF 95	99	99	
Leptotrichia buccalis		4.33	-.03		1	1	
57351160	**Low discrimination**						**57351160**
					SPO		
Clostridium paraputrificum		81.03	0.47	IND 1	99		
Clostridium barati		15.46	0.40	SOR 7	99		
Collinsella aerofaciens		2.16	0.15		1		
57351162	**Low discrimination**						**57351162**
					SPO		
Clostridium barati		49.30	0.48	IND 1	99		
Clostridium paraputrificum		36.91	0.40	SOR 8	99		
Lactobacillus acidophilus		6.59	0.35		1		
57351172	**Low discrimination**						**57351172**
					SPO		
Bifidobacterium breve		86.08	0.44	IND 1	1		
Clostridium barati		7.40	0.14	RAF 95	99		
Clostridium paraputrificum		5.54	0.06		99		
57351222	**Low discrimination**						**57351222**
					GRA	SPO	
Clostridium barati		87.89	0.40	IND 1	99	99	
Prevotella oralis		6.27	0.21	RHA 8	1	1	
Lactobacillus catenaformis		2.53	0.18		99	1	
57351302	**Low discrimination**						**57351302**
					SPO		
Clostridium barati		61.79	0.40	IND 1	99		
Lactobacillus acidophilus		15.10	0.31	RHA 8	1		
Lactobacillus catenaformis		10.67	0.32		1		
57351320	**Low discrimination**						**57351320**
					SPO		
Clostridium barati		50.72	0.40	IND 1	99		
Clostridium paraputrificum		35.68	0.32	RHA 8	99		
Collinsella aerofaciens		7.08	0.15		1		
57351322	**Low discrimination**						**57351322**
					GRA	SPO	
Clostridium barati		59.82	0.48	IND 1	99	99	
Prevotella oralis		22.29	0.41	RHA 8	1	1	
Clostridium paraputrificum		6.01	0.25		99	99	
57351522	**Low discrimination**						**57351522**
					GRA	SPO	
Bacteroides uniformis		66.36	0.38	RAF 97	1	1	
Clostridium barati		15.20	0.32	XYL 99	99	99	
Clostridium paraputrificum		13.15	0.25		99	99	
57351722	**Acceptable identification**						**57351722**
					GRA	SPO	
Bacteroides uniformis		94.73	0.40	RAF 97	1	1	
Bacteroides ovatus		2.09	0.15	XYL 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		1.37	0.14		99	99	
57353022	**Acceptable identification**						**57353022**
					SPO		
Clostridium barati		93.04	0.40	IND 1	99		
Lactobacillus catenaformis		2.68	0.18	MAN 8	1		
Lactobacillus acidophilus		1.38	0.11		1		
57353102	**Low discrimination**						**57353102**
					SPO		
Clostridium barati		48.48	0.40	IND 1	99		
Clostridium tertium		16.48	0.35	MAN 8	99		
Bifidobacterium breve		9.62	0.40		1		
57353112	**Low discrimination**						**57353112**
					SPO		
Clostridium tertium		72.16	0.39	IND 1	99		
Bifidobacterium breve		23.69	0.40	TRE 90	1		
Bifidobacterium adolescentis		1.44	0.18		1		
57353120	**Low discrimination**						**57353120**
Clostridium barati		42.04	0.40	IND 1			
Clostridium paraputrificum		29.57	0.32	MAN 8			
Clostridium tertium		14.29	0.35				
57353122	**Low discrimination**						**57353122**
Clostridium barati		55.67	0.48	IND 1			
Clostridium tertium		18.92	0.43	MAN 8			
Clostridium ramosum		8.46	0.35				
57353130	**Acceptable identification**						**57353130**
					SPO		
Clostridium tertium		92.06	0.39	IND 1	99		
Bifidobacterium breve		5.31	0.27	TRE 90	1		
Clostridium barati		1.54	0.05		99		
57353132	**Low discrimination**						**57353132**
					SPO		
Clostridium tertium		82.27	0.47	IND 1	99		
Bifidobacterium breve		16.01	0.44	TRE 90	1		
Clostridium barati		1.38	0.14		99		
57353142	**Low discrimination**						**57353142**
					SPO		
Bifidobacterium breve		62.44	0.40	IND 1	1		
Clostridium barati		27.37	0.21	RAF 95	99		
Lactobacillus acidophilus		6.65	0.13		1		
57353152	**Good identification**						**57353152**
					SPO		
Bifidobacterium breve		97.02	0.40	IND 1	1		
Clostridium tertium		2.99	0.05	RAF 95	99		
57353162	**Low discrimination**						**57353162**
					SPO		
Bifidobacterium breve		52.46	0.44	IND 1	1		
Clostridium barati		38.81	0.30	RAF 95	99		
Clostridium paraputrificum		3.38	0.06		99		
57353172	**Good identification**						**57353172**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		95.06	0.44	IND 1	1		
Clostridium tertium		4.94	0.13	RAF 95	99		
57353322	**Low discrimination**						**57353322**
					SPO		
Clostridium ramosum		66.68	0.37	IND 1	92		
Clostridium barati		28.79	0.30	TRE 94	99		
Lactobacillus acidophilus		1.42	0.09		1		
57354522	**Good identification**						**57354522**
Bacteroides uniformis		100.0	0.38	RAF 97			
57354722	**Good identification**						**57354722**
Bacteroides uniformis		97.22	0.40	RAF 97			
Bacteroides ovatus		2.14	0.15	GAL 99			
Clostridium clostridioforme		0.34	0.11				
57355122	**Acceptable identification**						**57355122**
					GRA	SPO	
Bacteroides uniformis		92.12	0.54	RAF 97	1	1	
Clostridium barati		2.45	0.32	ARA 92	99	99	
Prevotella oralis		2.27	0.32		1	1	
57355322	**Good identification**						**57355322**
Bacteroides uniformis		98.20	0.56	RAF 97			
Bacteroides ovatus		0.77	0.23	ARA 92			
Prevotella oralis		0.75	0.26				
57355422	**Good identification**						**57355422**
Bacteroides uniformis		100.0	0.43	RAF 97			
57355502	**Low discrimination**						**57355502**
					GRA	SPO	
Bacteroides uniformis		65.57	0.38	RAF 97	1	1	
Bifidobacterium adolescentis		23.32	0.46	NAG 99	99	1	
Clostridium butyricum		9.55	0.25		99	99	
57355512	**Acceptable identification**						**57355512**
					GRA		
Bifidobacterium adolescentis		93.53	0.48	IND 1	99		
Bacteroides uniformis		2.17	0.03	RAF 92	1		
Clostridium clostridioforme		1.95	0.16		50		
57355520	**Good identification**						**57355520**
					GRA	SPO	
Bacteroides uniformis		96.23	0.38	RAF 97	1	1	
Clostridium butyricum		3.13	0.14	BGL 99	99	99	
Clostridium paraputrificum		0.45	-.03		99	99	
57355522	**Very good identification**						**57355522**
					CAT		
Bacteroides uniformis		99.78	0.72	RAF 97	5		
Bacteroides ovatus		0.16	0.27		68		
Bacteroides thetaiotaomicron		0.03	0.15		89		
57355523	**Good identification**						**57355523**
Bacteroides uniformis		100.0	0.38	RAF 97			
57355532	**Low discrimination**						**57355532**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Bacteroides uniformis		88.41	0.38	RAF 97	5	
Bacteroides ovatus		9.35	0.24	MLZ 1	68	
Bacteroides thetaiotaomicron		1.06	0.08		89	
57355562	**Good identification**					**57355562**
Bacteroides uniformis		99.34	0.38	RAF 97		
Bacteroides ovatus		0.66	0.03	SOR 1		
57355622	**Good identification**					**57355622**
					CAT	
Bacteroides uniformis		98.48	0.45	RAF 97	5	
Bacteroides ovatus		1.07	0.15	LAC 98	68	
Bacteroides thetaiotaomicron		0.29	0.05		89	
57355702	**Acceptable identification**					**57355702**
					GRA	
Bacteroides uniformis		94.99	0.40	RAF 97	1	
Bacteroides ovatus		2.09	0.15	NAG 99	1	
Eubacterium contortum		1.08	0.14		99	
57355720	**Good identification**					**57355720**
Bacteroides uniformis		95.35	0.40	RAF 97		
Bacteroides ovatus		2.10	0.15	BGL 99		
Bacteroides vulgatus		1.19	0.08			
57355722	**Very good identification**					**57355722**
					CAT	
Bacteroides uniformis		97.54	0.74	RAF 97	5	
Bacteroides ovatus		2.15	0.49		68	
Bacteroides thetaiotaomicron		0.29	0.34		89	
57355723	**Good identification**					**57355723**
					CAT	
Bacteroides uniformis		97.56	0.40	RAF 97	5	
Bacteroides ovatus		2.15	0.15	URE 1	68	
Bacteroides thetaiotaomicron		0.29	0.00		89	
57355732	**Low discrimination**					**57355732**
					CAT	
Bacteroides ovatus		55.57	0.46	RAF 99	68	
Bacteroides uniformis		38.20	0.40	TRE 91	5	
Bacteroides thetaiotaomicron		4.39	0.27		89	
57355762	**Acceptable identification**					**57355762**
					CAT	
Bacteroides uniformis		91.42	0.40	RAF 97	5	
Bacteroides ovatus		8.31	0.25	SOR 1	68	
Bacteroides thetaiotaomicron		0.27	0.00		89	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57357002	**Low discrimination**						**57357002**
					SPO		
Actinomyces israelii		83.34	0.36	IND 1	1		
Clostridium tertium		6.85	0.03		99		
Clostridium innocuum		4.95	0.00		99		
57357102	**Low discrimination**						**57357102**
					SPO		
Clostridium tertium		60.24	0.37	IND 1	99		
Actinomyces israelii		29.62	0.46	TRE 90	1		
Bifidobacterium adolescentis		8.49	0.31		1		
57357112	**Low discrimination**						**57357112**
					SPO		
Clostridium tertium		87.86	0.41	IND 1	99		
Bifidobacterium adolescentis		8.51	0.32	TRE 90	1		
Actinomyces israelii		3.63	0.32		1		
57357120	**Acceptable identification**						**57357120**
					SPO		
Clostridium tertium		94.56	0.37	IND 1	99		
Clostridium barati		2.04	0.05	TRE 90	99		
Actinomyces israelii		1.98	0.23		1		
57357122	**Low discrimination**						**57357122**
					GRA	SPO	
Clostridium tertium		88.01	0.45	IND 1	99	99	
Bacteroides uniformis		8.27	0.19	TRE 90	1	1	
Clostridium barati		1.89	0.14		99	99	
57357130	**Good identification**						**57357130**
Clostridium tertium		100.0	0.41	IND 1			
57357132	**Good identification**						**57357132**
					SPO		
Clostridium tertium		99.86	0.50	IND 1	99		
Bifidobacterium breve		0.14	0.10	TRE 90	1		
57357140	**Good identification**						**57357140**
Actinomyces israelii		100.0	0.36	IND 1			
57357142	**Low discrimination**						**57357142**
					SPO		
Actinomyces israelii		87.42	0.39	IND 1	1		
Bifidobacterium adolescentis		7.97	0.15		1		
Clostridium tertium		4.62	0.03		99		
57357500	**Low discrimination**						**57357500**
					SPO		
Actinomyces israelii		65.01	0.42	IND 1	1		
Clostridium butyricum		32.68	0.16		99		
Bifidobacterium adolescentis		2.31	0.11		1		
57357502	**Low discrimination**						**57357502**
					SPO		
Bifidobacterium adolescentis		65.47	0.45	IND 1	1		
Actinomyces israelii		27.92	0.45	RAF 92	1		
Clostridium butyricum		3.64	0.09		99		
57357512	**Low discrimination**						**57357512**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		89.68	0.46	IND 1			
Actinomyces israelii		4.68	0.31	RAF 92			
Bifidobacterium dentium		4.25	0.06				
57357522	**Acceptable identification**						**57357522**
					GRA	SPO	
Bacteroides uniformis		93.19	0.38	RAF 97	1	1	
Bacteroides ovatus		4.93	0.19	MAN 1	1	1	
Clostridium tertium		0.87	0.11		99	99	
57357542	**Low discrimination**						**57357542**
Actinomyces israelii		57.30	0.38	IND 1			
Bifidobacterium adolescentis		42.70	0.29				
57357722	**Low discrimination**						**57357722**
					CAT		
Bacteroides uniformis		57.79	0.40	RAF 97	5		
Bacteroides ovatus		42.04	0.41	MAN 1	68		
Bacteroides thetaiotaomicron		0.17	0.00		89		
57357732	**Good identification**						**57357732**
Bacteroides ovatus		97.96	0.38	RAF 99			
Bacteroides uniformis		2.04	0.06	TRE 91			
57371000	**Very good identification**						**57371000**
					COC		
Lactobacillus jensenii		97.84	0.66	IND 1	1		
Lactobacillus catenaformis		0.66	0.29		1		
Atopobium parvulum		0.65	0.27		99		
57371002	**Low discrimination**						**57371002**
					SPO		
Lactobacillus jensenii		88.91	0.63	IND 1	1		
Lactobacillus acidophilus		7.35	0.41		1		
Clostridium barati		2.59	0.31		99		
57371010	**Very good identification**						**57371010**
					GRA		
Lactobacillus jensenii		99.77	0.58	IND 1	99		
Leptotrichia buccalis		0.17	-.03		1		
Eubacterium saburreum		0.06	0.11		85		
57371012	**Low discrimination**						**57371012**
					GRA		
Lactobacillus jensenii		79.81	0.55	IND 1	99		
Leptotrichia buccalis		19.92	0.32		1		
Lactobacillus acidophilus		0.20	0.07		99		
57371020	**Low discrimination**						**57371020**
					SPO		
Lactobacillus jensenii		87.85	0.56	IND 1	1		
Clostridium barati		6.83	0.31		99		
Clostridium paraputrificum		3.83	0.22		99		
57371022	**Low discrimination**						**57371022**
					SPO		
Lactobacillus jensenii		57.85	0.53	IND 1	1		
Clostridium barati		20.23	0.40		99		
Lactobacillus acidophilus		19.12	0.41		1		
57371030	**Good identification**						**57371030**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Lactobacillus jensenii		99.77	0.47	IND 1	1		
Clostridium barati		0.24	-.03		99		
57371032	**Low discrimination**						**57371032**
					GRA	SPO	
Lactobacillus jensenii		88.38	0.44	IND 1	99	1	
Leptotrichia buccalis		9.80	0.15		1	1	
Clostridium barati		0.94	0.05		99	99	
57371100	**Low discrimination**						**57371100**
					COC	SPO	
Atopobium parvulum		82.71	0.61	IND 1	99	1	
Collinsella aerofaciens		9.23	0.32		1	1	
Clostridium barati		2.45	0.31		1	99	
57371102	**Low discrimination**						**57371102**
					COC	SPO	
Lactobacillus acidophilus		69.94	0.60	IND 1	1	1	
Atopobium parvulum		12.79	0.48		99	1	
Clostridium barati		6.44	0.40		1	99	
57371112	**Very good identification**						**57371112**
					GRA		
Leptotrichia buccalis		99.60	0.66	IND 1	1		
Bifidobacterium breve		0.14	0.37		99		
Lactobacillus acidophilus		0.12	0.25		99		
57371120	**Low discrimination**						**57371120**
					COC	SPO	
Clostridium paraputrificum		81.40	0.56	IND 1	1	99	
Atopobium parvulum		8.74	0.48		99	1	
Clostridium barati		4.40	0.40		1	99	
57371122	**Low discrimination**						**57371122**
					SPO		
Lactobacillus acidophilus		45.03	0.60	IND 1	1		
Clostridium paraputrificum		32.87	0.50		99		
Clostridium barati		12.43	0.48		99		
57371132	**Acceptable identification**						**57371132**
					GRA	SPO	
Leptotrichia buccalis		94.19	0.50	IND 1	1	1	
Bifidobacterium breve		2.12	0.41	NAG 10	99	1	
Clostridium tertium		1.48	0.29		99	99	
57371142	**Low discrimination**						**57371142**
					SPO		
Lactobacillus acidophilus		83.78	0.44	IND 1	1		
Bifidobacterium breve		8.25	0.37	SOR 11	1		
Clostridium barati		5.42	0.21		99		
57371160	**Low discrimination**						**57371160**
					SPO		
Clostridium paraputrificum		89.63	0.37	IND 1	99		
Clostridium barati		5.59	0.21	SOR 7	99		
Bifidobacterium breve		1.50	0.24		1		
57371162	**Low discrimination**						**57371162**
					SPO		
Lactobacillus acidophilus		54.67	0.44	IND 1	1		
Clostridium paraputrificum		24.30	0.31	SOR 11	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		10.62	0.30		99		
57371172	**Low discrimination**						**57371172**
					GRA		
Bifidobacterium breve		65.22	0.41	IND 1	99		
Leptotrichia buccalis		29.34	0.15	RAF 95	1		
Lactobacillus acidophilus		3.77	0.10		99		
57371302	**Low discrimination**						**57371302**
					COC	SPO	
Lactobacillus acidophilus		85.63	0.40	IND 1	1	1	
Clostridium barati		9.10	0.21	RHA 7	1	99	
Atopobium parvulum		2.10	0.14		99	1	
57371322	**Low discrimination**						**57371322**
					SPO		
Clostridium ramosum		57.09	0.45	IND 1	92		
Lactobacillus acidophilus		27.97	0.40		1		
Clostridium barati		8.92	0.30		99		
57373100	**Low discrimination**						**57373100**
					COC	SPO	
Clostridium tertium		80.09	0.43	IND 1	1	99	
Atopobium parvulum		8.93	0.27		99	1	
Actinomyces israelii		3.95	0.35		1	1	
57373102	**Low discrimination**						**57373102**
					SPO		
Clostridium tertium		75.69	0.51	IND 1	99		
Lactobacillus acidophilus		14.15	0.38		1		
Bifidobacterium breve		3.27	0.37		1		
57373110	**Good identification**						**57373110**
					GRA	SPO	
Clostridium tertium		98.36	0.47	IND 1	99	99	
Bifidobacterium breve		0.71	0.20		99	1	
Leptotrichia buccalis		0.52	-.03		1	1	
57373112	**Low discrimination**						**57373112**
					GRA	SPO	
Clostridium tertium		83.06	0.55	IND 1	99	99	
Leptotrichia buccalis		14.54	0.32		1	1	
Bifidobacterium breve		2.02	0.37		99	1	
57373120	**Low discrimination**						**57373120**
Clostridium tertium		59.03	0.51	IND 1			
Clostridium ramosum		34.44	0.45				
Clostridium paraputrificum		3.61	0.22				
57373122	**Low discrimination**						**57373122**
					SPO		
Clostridium tertium		52.50	0.59	IND 1	99		
Clostridium ramosum		40.84	0.56		92		
Lactobacillus acidophilus		3.27	0.38		1		
57373130	**Good identification**						**57373130**
					SPO		
Clostridium tertium		99.25	0.55	IND 1	99		
Bifidobacterium breve		0.42	0.24		1		
Clostridium ramosum		0.33	0.11		92		
57373132	**Very good identification**						**57373132**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium tertium		97.50	0.63	IND 1	99	99	
Bifidobacterium breve		1.41	0.41		99	1	
Leptotrichia buccalis		0.63	0.15		1	1	
57373142	**Low discrimination**						**57373142**
					SPO		
Bifidobacterium breve		48.84	0.37	IND 1	1		
Lactobacillus acidophilus		26.11	0.22	RAF 95	1		
Clostridium tertium		11.41	0.17		99		
57373152	**Low discrimination**						**57373152**
					GRA	SPO	
Bifidobacterium breve		65.74	0.37	IND 1	99	1	
Clostridium tertium		27.31	0.21	RAF 95	99	99	
Leptotrichia buccalis		4.78	-.03		1	1	
57373162	**Low discrimination**						**57373162**
					SPO		
Bifidobacterium breve		47.08	0.41	IND 1	1		
Clostridium tertium		18.56	0.25	RAF 95	99		
Clostridium ramosum		14.44	0.21		92		
57373172	**Low discrimination**						**57373172**
					SPO		
Bifidobacterium breve		58.79	0.41	IND 1	1		
Clostridium tertium		41.21	0.29	RAF 95	99		
57373320	**Good identification**						**57373320**
Clostridium ramosum		98.40	0.47	IND 1			
Clostridium tertium		1.29	0.17				
Clostridium barati		0.31	0.03				
57373322	**Very good identification**						**57373322**
					SPO		
Clostridium ramosum		98.30	0.58	IND 1	92		
Clostridium tertium		0.96	0.25		99		
Lactobacillus acidophilus		0.45	0.18		1		
57373502	**Low discrimination**						**57373502**
					SPO		
Actinomyces israelii		39.98	0.36	IND 1	1		
Bifidobacterium adolescentis		28.55	0.28	XYL 92	1		
Clostridium tertium		20.00	0.17		99		
57374712	**Good identification**						**57374712**
Clostridium clostridioforme		98.74	0.37	IND 1			
Bacteroides ovatus		1.26	-.06	RAF 94			
57375100	**Good identification**						**57375100**
					COC	SPO	
Atopobium parvulum		94.88	0.66	IND 1	99	1	
Clostridium butyricum		4.50	0.36		1	99	
Actinomyces israelii		0.46	0.40		1	1	
57375102	**Low discrimination**						**57375102**
					COC	SPO	
Atopobium parvulum		82.83	0.53	IND 1	99	1	
Clostridium butyricum		8.65	0.29		1	99	
Actinomyces israelii		3.44	0.43		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
57375120	**Acceptable identification**							**57375120**
					COC	SPO		
Atopobium parvulum		91.55	0.53	IND 1	99	1		
Clostridium paraputrificum		4.64	0.22		1	99		
Clostridium butyricum		2.14	0.18		1	99		
57375122	**Low discrimination**							**57375122**
					COC	GRA	SPO	
Atopobium parvulum		50.10	0.40	IND 1	99	99	1	
Bacteroides uniformis		10.18	0.19		1	1	1	
Clostridium tertium		9.85	0.27		1	99	99	
57375142	**Low discrimination**							**57375142**
					COC	SPO		
Actinomyces israelii		39.56	0.36	IND 1	1	1		
Atopobium parvulum		24.75	0.19		99	1		
Clostridium butyricum		22.26	0.11		1	99		
57375322	**Low discrimination**							**57375322**
					CAT			
Bacteroides ovatus		80.54	0.40	RAF 99	68			
Bacteroides uniformis		10.26	0.22	ARA 97	5			
Bacteroides thetaiotaomicron		4.69	0.19		89			
57375332	**Acceptable identification**							**57375332**
Bacteroides ovatus		90.91	0.37	RAF 99				
Clostridium clostridioforme		4.50	0.25	ARA 97				
Bacteroides thetaiotaomicron		3.09	0.12					
57375500	**Good identification**							**57375500**
					COC	SPO		
Clostridium butyricum		96.76	0.52	IND 3	1	99		
Atopobium parvulum		2.29	0.32	RAF 95	99	1		
Actinomyces israelii		0.91	0.39		1	1		
57375502	**Low discrimination**							**57375502**
					SPO			
Clostridium butyricum		85.43	0.45	IND 3	99			
Bifidobacterium adolescentis		8.88	0.43	RAF 95	1			
Actinomyces israelii		3.09	0.42		1			
57375512	**Low discrimination**							**57375512**
					SPO			
Bifidobacterium adolescentis		55.46	0.45	IND 1	1			
Clostridium butyricum		22.97	0.23	RAF 92	99			
Clostridium clostridioforme		12.71	0.31		66			
57375522	**Low discrimination**							**57375522**
					CAT			
Bacteroides ovatus		49.48	0.44	RAF 99	68			
Bacteroides uniformis		30.86	0.38	RHA 95	5			
Bacteroides thetaiotaomicron		12.60	0.34		89			
57375532	**Low discrimination**							**57375532**
Bacteroides ovatus		79.20	0.41	RAF 99				
Bacteroides thetaiotaomicron		11.77	0.27	RHA 95				
Clostridium clostridioforme		4.92	0.31					
57375612	**Good identification**							**57375612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		98.60	0.36	IND 1			
Bacteroides ovatus		1.41	-.06	RAF 94			
57375712	**Low discrimination**						**57375712**
Clostridium clostridioforme		67.63	0.45	IND 1			
Bacteroides ovatus		28.49	0.29	RAF 94			
Bacteroides thetaiotaomicron		2.96	0.12				
57375722	**Low discrimination**						**57375722**
					CAT		
Bacteroides ovatus		81.46	0.66	RAF 99	68		
Bacteroides thetaiotaomicron		14.51	0.53		89		
Bacteroides uniformis		3.69	0.40		5		
57375732	**Low discrimination**						**57375732**
Bacteroides ovatus		88.58	0.63	RAF 99			
Bacteroides thetaiotaomicron		9.20	0.46				
Clostridium clostridioforme		2.12	0.45				
57375762	**Acceptable identification**						**57375762**
					CAT		
Bacteroides ovatus		94.86	0.42	RAF 99	68		
Bacteroides thetaiotaomicron		4.10	0.19	SOR 4	89		
Bacteroides uniformis		1.04	0.06		5		
57375772	**Good identification**						**57375772**
Bacteroides ovatus		96.99	0.39	RAF 99			
Bacteroides thetaiotaomicron		2.44	0.12	SOR 4			
Clostridium clostridioforme		0.56	0.11				
57377000	**Low discrimination**						**57377000**
					SPO		
Actinomyces israelii		88.49	0.43	IND 1	1		
Clostridium tertium		8.70	0.11		99		
Clostridium butyricum		2.81	-.04		99		
57377002	**Low discrimination**						**57377002**
					SPO		
Actinomyces israelii		83.57	0.46	IND 1	1		
Clostridium tertium		16.43	0.19		99		
57377040	**Good identification**						**57377040**
Actinomyces israelii		100.0	0.36	IND 1			
57377042	**Good identification**						**57377042**
Actinomyces israelii		100.0	0.39	IND 1			
57377100	**Low discrimination**						**57377100**
					COC	SPO	
Clostridium tertium		61.00	0.45	IND 1	1	99	
Actinomyces israelii		25.07	0.53		1	1	
Atopobium parvulum		9.15	0.32		99	1	
57377102	**Low discrimination**						**57377102**
					SPO		
Clostridium tertium		80.89	0.53	IND 1	99		
Actinomyces israelii		16.62	0.56		1		
Bifidobacterium adolescentis		0.85	0.28		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57377110	**Good identification**						**57377110**
					SPO		
Clostridium tertium		96.45	0.50	IND 1	99		
Actinomyces israelii		3.33	0.39		1		
Clostridium butyricum		0.22	-.02		99		
57377112	**Very good identification**						**57377112**
					SPO		
Clostridium tertium		97.17	0.58	IND 1	99		
Actinomyces israelii		1.68	0.42		1		
Bifidobacterium adolescentis		0.70	0.29		1		
57377120	**Good identification**						**57377120**
					COC	SPO	
Clostridium tertium		97.65	0.53	IND 1	1	99	
Atopobium parvulum		0.86	0.19		99	1	
Actinomyces israelii		0.85	0.33		1	1	
57377122	**Very good identification**						**57377122**
					SPO		
Clostridium tertium		98.69	0.62	IND 1	99		
Clostridium ramosum		0.56	0.21		92		
Actinomyces israelii		0.43	0.36		1		
57377130	**Very good identification**						**57377130**
					SPO		
Clostridium tertium		99.87	0.58	IND 1	99		
Actinomyces israelii		0.07	0.18		1		
Bifidobacterium dentium		0.06	-.07		1		
57377132	**Very good identification**						**57377132**
					GRA	SPO	
Clostridium tertium		99.69	0.66	IND 1	99	99	
Bifidobacterium dentium		0.20	0.10		99	1	
Bacteroides ovatus		0.07	0.07		1	1	
57377140	**Acceptable identification**						**57377140**
					SPO		
Actinomyces israelii		90.44	0.46	IND 1	1		
Clostridium tertium		5.72	0.11		99		
Clostridium butyricum		3.85	0.01		99		
57377142	**Low discrimination**						**57377142**
					SPO		
Actinomyces israelii		86.69	0.49	IND 1	1		
Clostridium tertium		10.96	0.19		99		
Bifidobacterium adolescentis		1.40	0.12		1		
57377302	**Low discrimination**						**57377302**
					GRA	SPO	
Actinomyces israelii		57.91	0.37	IND 1	99	1	
Clostridium tertium		37.81	0.19	RHA 7	99	99	
Bacteroides ovatus		4.28	-.02		1	1	
57377400	**Low discrimination**						**57377400**
					SPO		
Actinomyces israelii		74.13	0.41	IND 1	1		
Clostridium butyricum		25.87	0.12		99		
57377402	**Acceptable identification**						**57377402**
					SPO		
Actinomyces israelii		90.31	0.44	IND 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		8.17	0.05		99		
Bifidobacterium adolescentis		1.52	0.08		1		
57377442	**Good identification**						**57377442**
Actinomyces israelii		100.0	0.37	IND 1			
57377500	**Low discrimination**						**57377500**
					SPO		
Clostridium butyricum		66.89	0.36	IND 3	99		
Actinomyces israelii		31.95	0.52	RAF 95	1		
Clostridium tertium		0.96	0.11	MAN 10	99		
57377502	**Low discrimination**						**57377502**
					SPO		
Actinomyces israelii		48.40	0.55	IND 1	1		
Clostridium butyricum		26.27	0.29		99		
Bifidobacterium adolescentis		20.11	0.42		1		
57377512	**Low discrimination**						**57377512**
Bifidobacterium dentium		81.71	0.40	IND 1			
Bifidobacterium adolescentis		11.60	0.43	RAF 95			
Actinomyces israelii		3.41	0.40				
57377522	**Low discrimination**						**57377522**
					GRA	SPO	
Bacteroides ovatus		74.59	0.36	RAF 99	1	1	
Clostridium tertium		11.74	0.27	RHA 95	99	99	
Bifidobacterium dentium		4.18	0.10		99	1	
57377532	**Low discrimination**						**57377532**
					GRA	SPO	
Bifidobacterium dentium		85.27	0.44	IND 1	99	1	
Bacteroides ovatus		10.24	0.33	RAF 95	1	1	
Clostridium tertium		4.30	0.32		99	99	
57377540	**Low discrimination**						**57377540**
					COC	SPO	
Actinomyces israelii		42.77	0.45	IND 1	1	1	
Blautia producta		37.21	0.24		99	1	
Clostridium butyricum		20.02	0.18		1	99	
57377542	**Low discrimination**						**57377542**
					SPO		
Actinomyces israelii		75.88	0.48	IND 1	1		
Bifidobacterium adolescentis		10.02	0.26		1		
Clostridium butyricum		9.21	0.11		99		
57377722	**Very good identification**						**57377722**
					CAT		
Bacteroides ovatus		99.31	0.58	RAF 99	68		
Bacteroides thetaiotaomicron		0.54	0.19		89		
Bacteroides uniformis		0.14	0.06		5		
57377732	**Good identification**						**57377732**
					GRA	CAT	
Bacteroides ovatus		99.18	0.55	RAF 99	1	68	
Bifidobacterium dentium		0.44	0.10		99	1	
Bacteroides thetaiotaomicron		0.31	0.12		1	89	
57401000	**Very good identification**						**57401000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		99.63	0.60			
Fusobacterium mortiferum		0.37	0.20			
57401002	**Good identification**					**57401002**
Eubacterium saburreum		100.0	0.47			
57401010	**Very good identification**					**57401010**
Eubacterium saburreum		100.0	0.59			
57401012	**Good identification**					**57401012**
Eubacterium saburreum		100.0	0.46			
57401020	**Good identification**					**57401020**
Eubacterium saburreum		100.0	0.47			
57401030	**Good identification**					**57401030**
Eubacterium saburreum		100.0	0.46			
57401100	**Very good identification**					**57401100**
Eubacterium saburreum		99.63	0.67			
Fusobacterium mortiferum		0.37	0.27			
57401102	**Good identification**					**57401102**
Eubacterium saburreum		100.0	0.54			
57401110	**Very good identification**					**57401110**
Eubacterium saburreum		100.0	0.65			
57401112	**Good identification**					**57401112**
Eubacterium saburreum		100.0	0.52			
57401120	**Good identification**					**57401120**
Eubacterium saburreum		96.51	0.54			
Prevotella melaninogenica		3.14	0.12			
Odoribacter splanchnicus		0.35	-.07			
57401122	**Good identification**					**57401122**
Eubacterium saburreum		96.60	0.41			
Prevotella melaninogenica		3.40	0.00			
57401130	**Good identification**					**57401130**
Eubacterium saburreum		100.0	0.52			
57401132	**Good identification**					**57401132**
Eubacterium saburreum		100.0	0.39			
57405000	**Good identification**					**57405000**
Eubacterium saburreum		100.0	0.56			
57405002	**Good identification**					**57405002**
Eubacterium saburreum		100.0	0.43			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57405010	**Good identification**						**57405010**
Eubacterium saburreum		100.0	0.54				
57405012	**Good identification**						**57405012**
Eubacterium saburreum		100.0	0.41				
57405020	**Good identification**						**57405020**
Eubacterium saburreum		100.0	0.43				
57405030	**Good identification**						**57405030**
Eubacterium saburreum		100.0	0.41				
57405100	**Very good identification**						**57405100**
Eubacterium saburreum		100.0	0.62				
57405102	**Good identification**						**57405102**
Eubacterium saburreum		100.0	0.49				
57405110	**Very good identification**						**57405110**
Eubacterium saburreum		100.0	0.61				
57405112	**Good identification**						**57405112**
Eubacterium saburreum		100.0	0.48				
57405120	**Good identification**						**57405120**
Eubacterium saburreum		100.0	0.49				
57405122	**Acceptable identification**						**57405122**
					CAT		
Eubacterium saburreum		91.13	0.36		1		
Bacteroides fragilis		8.87	-.03		99		
57405130	**Good identification**						**57405130**
Eubacterium saburreum		100.0	0.48				
57410000	**Low discrimination**						**57410000**
Fusobacterium mortiferum		75.26	0.52	IND 1			
Prevotella intermedia		11.40	0.26				
Eubacterium saburreum		9.23	0.34				
57410002	**Low discrimination**						**57410002**
					CAT	GRA	
Fusobacterium mortiferum		78.44	0.35	IND 1	5	1	
Eubacterium saburreum		15.29	0.21	BGL 10	1	85	
Propionibacterium granulosum		6.27	0.13		90	99	
57410100	**Low discrimination**						**57410100**
Fusobacterium mortiferum		89.07	0.58	IND 1			
Eubacterium saburreum		10.93	0.41				
57410102	**Low discrimination**						**57410102**
Fusobacterium mortiferum		83.69	0.42	IND 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		16.31	0.28	BGL 10			
57410110	**Acceptable identification**						**57410110**
Eubacterium saburreum		90.86	0.39	GAL 99			
Fusobacterium mortiferum		9.14	0.24				
57411000	**Low discrimination**						**57411000**
Eubacterium saburreum		87.81	0.69				
Fusobacterium mortiferum		10.84	0.55				
Prevotella intermedia		1.10	0.26				
57411002	**Acceptable identification**						**57411002**
					CAT	GRA	
Eubacterium saburreum		92.43	0.56		1	85	
Fusobacterium mortiferum		7.19	0.38		5	1	
Propionibacterium granulosum		0.38	0.13		90	99	
57411010	**Very good identification**						**57411010**
					CAT	GRA	
Eubacterium saburreum		99.80	0.67		1	85	
Fusobacterium mortiferum		0.15	0.20		5	1	
Propionibacterium granulosum		0.05	0.09		90	99	
57411012	**Good identification**						**57411012**
Eubacterium saburreum		100.0	0.54				
57411020	**Good identification**						**57411020**
					CAT	GRA	
Eubacterium saburreum		98.67	0.56		1	85	
Fusobacterium mortiferum		0.70	0.20		5	1	
Propionibacterium granulosum		0.41	0.13		90	99	
57411022	**Good identification**						**57411022**
Eubacterium saburreum		98.79	0.43				
Prevotella oralis		1.21	-.03				
57411030	**Good identification**						**57411030**
Eubacterium saburreum		100.0	0.54				
57411032	**Good identification**						**57411032**
Eubacterium saburreum		100.0	0.41				
57411100	**Low discrimination**						**57411100**
					GRA		
Eubacterium saburreum		88.98	0.75		85		
Fusobacterium mortiferum		10.99	0.61		1		
Lactobacillus fermentum		0.02	0.15		99		
57411101	**Low discrimination**						**57411101**
Eubacterium saburreum		89.01	0.41	URE 1			
Fusobacterium mortiferum		10.99	0.27				
57411102	**Good identification**						**57411102**
					CAT		
Eubacterium saburreum		92.05	0.62		1		
Fusobacterium mortiferum		7.16	0.45		5		
Capnocytophaga ochracea		0.60	0.12		90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57411110	**Very good identification**					**57411110**
Eubacterium saburreum		99.85	0.73			
Fusobacterium mortiferum		0.15	0.27			
57411111	**Good identification**					**57411111**
Eubacterium saburreum		100.0	0.39	URE 1		
57411112	**Very good identification**					**57411112**
Eubacterium saburreum		100.0	0.60			
57411120	**Low discrimination**					**57411120**
Eubacterium saburreum		89.68	0.62			
Prevotella melaninogenica		8.76	0.29			
Fusobacterium mortiferum		0.63	0.27			
57411122	**Low discrimination**					**57411122**
					CAT	
Eubacterium saburreum		63.87	0.49		1	
Capnocytophaga ochracea		21.24	0.29		90	
Prevotella melaninogenica		6.74	0.16		1	
57411130	**Very good identification**					**57411130**
Eubacterium saburreum		99.88	0.60			
Prevotella melaninogenica		0.12	-.06			
57411132	**Acceptable identification**					**57411132**
					CAT	
Eubacterium saburreum		94.51	0.48		1	
Capnocytophaga ochracea		4.27	0.12		90	
Parabacteroides distasonis		1.22	0.16		76	
57411140	**Low discrimination**					**57411140**
Eubacterium saburreum		89.01	0.41	SOR 1		
Fusobacterium mortiferum		10.99	0.27			
57411150	**Good identification**					**57411150**
Eubacterium saburreum		100.0	0.39	SOR 1		
57411300	**Low discrimination**					**57411300**
Eubacterium saburreum		89.01	0.41	RHA 1		
Fusobacterium mortiferum		10.99	0.27			
57411310	**Good identification**					**57411310**
Eubacterium saburreum		100.0	0.39	RHA 1		
57411500	**Low discrimination**					**57411500**
Eubacterium saburreum		82.72	0.41	ARA 1		
Fusobacterium mortiferum		10.21	0.27			
Odoribacter splanchnicus		3.23	-.03			
57411510	**Good identification**					**57411510**
Eubacterium saburreum		95.12	0.39	ARA 1		
Bifidobacterium longum subsp. longum		4.88	0.13			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57413100	**Low discrimination**					**57413100**
Eubacterium saburreum		89.01	0.41	MAN 1		
Fusobacterium mortiferum		10.99	0.27			
57413110	**Good identification**					**57413110**
Eubacterium saburreum		100.0	0.39	MAN 1		
57414100	**Low discrimination**					**57414100**
Fusobacterium mortiferum		53.25	0.39	IND 1		
Eubacterium saburreum		46.75	0.36	XYL 7		
57415000	**Very good identification**					**57415000**
Eubacterium saburreum		98.30	0.64			
Fusobacterium mortiferum		1.70	0.35			
57415002	**Good identification**					**57415002**
Eubacterium saburreum		98.93	0.51			
Fusobacterium mortiferum		1.08	0.19			
57415010	**Very good identification**					**57415010**
Eubacterium saburreum		100.0	0.62			
57415012	**Good identification**					**57415012**
Eubacterium saburreum		100.0	0.50			
57415020	**Good identification**					**57415020**
Eubacterium saburreum		100.0	0.51			
57415022	**Acceptable identification**					**57415022**
					CAT	
Eubacterium saburreum		92.96	0.38		1	
Bacteroides fragilis		7.04	-.03		99	
57415030	**Good identification**					**57415030**
Eubacterium saburreum		100.0	0.50			
57415032	**Good identification**					**57415032**
Eubacterium saburreum		100.0	0.37			
57415100	**Very good identification**					**57415100**
Eubacterium saburreum		98.30	0.70			
Fusobacterium mortiferum		1.70	0.42			
57415101	**Good identification**					**57415101**
Eubacterium saburreum		100.0	0.36	URE 1		
57415102	**Very good identification**					**57415102**
					CAT	
Eubacterium saburreum		98.08	0.57		1	
Fusobacterium mortiferum		1.07	0.25		5	
Bacteroides fragilis		0.56	-.03		99	
57415110	**Very good identification**					**57415110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.69			
57415112	**Good identification**					**57415112**
Eubacterium saburreum		100.0	0.56			
57415120	**Very good identification**					**57415120**
					CAT	
Eubacterium saburreum		96.36	0.57		1	
Bacteroides fragilis		2.87	0.10		99	
Capnocytophaga ochracea		0.29	0.02		90	
57415130	**Good identification**					**57415130**
Eubacterium saburreum		100.0	0.56			
57415132	**Low discrimination**					**57415132**
					CAT	
Eubacterium saburreum		77.59	0.43		1	
Parabacteroides distasonis		16.77	0.32		76	
Bacteroides fragilis		3.08	-.03		99	
57415140	**Good identification**					**57415140**
Eubacterium saburreum		100.0	0.36	SOR 1		
57415300	**Good identification**					**57415300**
Eubacterium saburreum		100.0	0.36	RHA 1		
57415322	**Low discrimination**					**57415322**
					CAT	
Parabacteroides distasonis		65.43	0.45	IND 1	76	
Bacteroides thetaiotaomicron		22.33	0.21		89	
Bacteroides uniformis		9.43	0.11		5	
57415332	**Low discrimination**					**57415332**
Parabacteroides distasonis		80.23	0.40	IND 1		
Bacteroides thetaiotaomicron		19.77	0.14			
57415500	**Low discrimination**					**57415500**
Eubacterium saburreum		87.92	0.36	ARA 1		
Bifidobacterium longum subsp. longum		12.09	0.17			
57415522	**Low discrimination**					**57415522**
					CAT	
Bacteroides thetaiotaomicron		60.56	0.36	MLT 94	89	
Bacteroides uniformis		28.58	0.27	TRE 93	5	
Parabacteroides distasonis		4.87	0.33	RHA 93	76	
57415722	**Acceptable identification**					**57415722**
					CAT	
Bacteroides thetaiotaomicron		93.60	0.55	MLT 94	89	
Bacteroides uniformis		4.59	0.30	TRE 93	5	
Parabacteroides distasonis		1.70	0.41		76	
57415732	**Good identification**					**57415732**
					CAT	
Bacteroides thetaiotaomicron		97.33	0.48	MLT 94	89	
Parabacteroides distasonis		2.45	0.36	TRE 93	76	
Bacteroides uniformis		0.12	-.05		5	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57417100	**Good identification**					**57417100**
Eubacterium saburreum		100.0	0.36	MAN 1		
57421000	**Very good identification**					**57421000**
Eubacterium saburreum		99.83	0.57			
Fusobacterium mortiferum		0.17	0.12			
57421002	**Good identification**					**57421002**
Eubacterium saburreum		100.0	0.44			
57421010	**Good identification**					**57421010**
Eubacterium saburreum		100.0	0.56			
57421012	**Good identification**					**57421012**
Eubacterium saburreum		100.0	0.43			
57421020	**Good identification**					**57421020**
Eubacterium saburreum		100.0	0.44			
57421030	**Good identification**					**57421030**
Eubacterium saburreum		100.0	0.43			
57421100	**Very good identification**					**57421100**
Eubacterium saburreum		99.83	0.64			
Fusobacterium mortiferum		0.17	0.18			
57421102	**Good identification**					**57421102**
Eubacterium saburreum		100.0	0.51			
57421110	**Very good identification**					**57421110**
Eubacterium saburreum		100.0	0.62			
57421112	**Good identification**					**57421112**
Eubacterium saburreum		100.0	0.49			
57421120	**Good identification**					**57421120**
Eubacterium saburreum		100.0	0.51			
57421122	**Good identification**					**57421122**
Eubacterium saburreum		100.0	0.38			
57421130	**Good identification**					**57421130**
Eubacterium saburreum		100.0	0.49			
57421132	**Good identification**					**57421132**
Eubacterium saburreum		100.0	0.36			
57425000	**Good identification**					**57425000**
Eubacterium saburreum		100.0	0.53			
57425002	**Good identification**					**57425002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.40				
57425010	**Good identification**						**57425010**
Eubacterium saburreum		100.0	0.51				
57425012	**Good identification**						**57425012**
Eubacterium saburreum		100.0	0.38				
57425020	**Good identification**						**57425020**
Eubacterium saburreum		100.0	0.40				
57425030	**Good identification**						**57425030**
Eubacterium saburreum		100.0	0.38				
57425100	**Very good identification**						**57425100**
Eubacterium saburreum		100.0	0.59				
57425102	**Good identification**						**57425102**
Eubacterium saburreum		100.0	0.46				
57425110	**Very good identification**						**57425110**
Eubacterium saburreum		100.0	0.58				
57425112	**Good identification**						**57425112**
Eubacterium saburreum		100.0	0.45				
57425120	**Good identification**						**57425120**
Eubacterium saburreum		100.0	0.46				
57425130	**Good identification**						**57425130**
Eubacterium saburreum		100.0	0.45				
57425722	**Good identification**						**57425722**
Bacteroides thetaiotaomicron		99.38	0.40	MLT 94			
Parabacteroides distasonis		0.62	0.18	FRU 99			
57430000	**Low discrimination**						**57430000**
					CAT	GRA	
Fusobacterium mortiferum		61.33	0.43	IND 1	5	1	
Propionibacterium granulosum		21.88	0.32		90	99	
Eubacterium saburreum		16.80	0.31		1	85	
57430100	**Low discrimination**						**57430100**
Fusobacterium mortiferum		78.50	0.49	IND 1			
Eubacterium saburreum		21.50	0.38				
57430110	**Good identification**						**57430110**
Eubacterium saburreum		95.69	0.36	GAL 99			
Fusobacterium mortiferum		4.31	0.15				
57431000	**Good identification**						**57431000**
					CAT	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		93.59	0.66		1	85	
Fusobacterium mortiferum		5.18	0.46		5	1	
Propionibacterium granulosum		1.23	0.32		90	99	
57431002	**Acceptable identification**						**57431002**
					CAT	GRA	
Eubacterium saburreum		94.92	0.53		1	85	
Fusobacterium mortiferum		3.31	0.29		5	1	
Propionibacterium granulosum		1.77	0.21		90	99	
57431010	**Very good identification**						**57431010**
					CAT	GRA	
Eubacterium saburreum		99.71	0.64		1	85	
Propionibacterium granulosum		0.22	0.17		90	99	
Fusobacterium mortiferum		0.07	0.12		5	1	
57431012	**Good identification**						**57431012**
Eubacterium saburreum		100.0	0.51				
57431020	**Good identification**						**57431020**
					CAT	GRA	
Eubacterium saburreum		97.87	0.53		1	85	
Propionibacterium granulosum		1.82	0.21		90	99	
Fusobacterium mortiferum		0.31	0.12		5	1	
57431022	**Good identification**						**57431022**
					CAT		
Eubacterium saburreum		97.43	0.40		1		
Propionibacterium granulosum		2.57	0.11		90		
57431030	**Good identification**						**57431030**
Eubacterium saburreum		100.0	0.51				
57431032	**Good identification**						**57431032**
Eubacterium saburreum		100.0	0.38				
57431100	**Good identification**						**57431100**
Eubacterium saburreum		94.76	0.72				
Fusobacterium mortiferum		5.24	0.52				
57431101	**Acceptable identification**						**57431101**
Eubacterium saburreum		94.76	0.38	URE 1			
Fusobacterium mortiferum		5.24	0.18				
57431102	**Very good identification**						**57431102**
Eubacterium saburreum		96.63	0.59				
Fusobacterium mortiferum		3.37	0.36				
57431110	**Very good identification**						**57431110**
Eubacterium saburreum		99.93	0.70				
Fusobacterium mortiferum		0.07	0.18				
57431111	**Good identification**						**57431111**
Eubacterium saburreum		100.0	0.36	URE 1			
57431112	**Very good identification**						**57431112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.57			
57431120	**Very good identification**					**57431120**
					GRA	
Eubacterium saburreum		99.13	0.59		85	
Clostridium perfringens		0.41	0.09		99	
Fusobacterium mortiferum		0.31	0.18		1	
57431122	**Low discrimination**					**57431122**
					CAT	
Eubacterium saburreum		85.87	0.46		1	
Parabacteroides distasonis		11.53	0.31		76	
Bacteroides thetaiotaomicron		0.97	-.03		89	
57431130	**Very good identification**					**57431130**
Eubacterium saburreum		100.0	0.57			
57431132	**Acceptable identification**					**57431132**
Eubacterium saburreum		91.88	0.45			
Parabacteroides distasonis		8.12	0.27			
57431140	**Acceptable identification**					**57431140**
Eubacterium saburreum		94.76	0.38	SOR 1		
Fusobacterium mortiferum		5.24	0.18			
57431150	**Good identification**					**57431150**
Eubacterium saburreum		100.0	0.36	SOR 1		
57431300	**Acceptable identification**					**57431300**
Eubacterium saburreum		94.76	0.38	RHA 1		
Fusobacterium mortiferum		5.24	0.18			
57431310	**Good identification**					**57431310**
Eubacterium saburreum		100.0	0.36	RHA 1		
57431322	**Low discrimination**					**57431322**
					CAT	
Parabacteroides distasonis		71.52	0.40	IND 1	76	
Bacteroides thetaiotaomicron		26.69	0.16	XYL 90	89	
Eubacterium saburreum		1.79	0.12		1	
57431500	**Acceptable identification**					**57431500**
Eubacterium saburreum		94.76	0.38	ARA 1		
Fusobacterium mortiferum		5.24	0.18			
57431510	**Good identification**					**57431510**
Eubacterium saburreum		100.0	0.36	ARA 1		
57431722	**Good identification**					**57431722**
Bacteroides thetaiotaomicron		98.37	0.50	MLT 94		
Parabacteroides distasonis		1.63	0.36	XYL 96		
57431732	**Good identification**					**57431732**
Bacteroides thetaiotaomicron		97.76	0.43	MLT 94		
Parabacteroides distasonis		2.25	0.31	XYL 96		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57433100	**Acceptable identification**					**57433100**
Eubacterium saburreum		94.76	0.38	MAN 1		
Fusobacterium mortiferum		5.24	0.18			
57433110	**Good identification**					**57433110**
Eubacterium saburreum		100.0	0.36	MAN 1		
57434722	**Good identification**					**57434722**
Bacteroides thetaiotaomicron		99.38	0.40	MLT 94		
Parabacteroides distasonis		0.62	0.18	GAL 99		
57435000	**Very good identification**					**57435000**
Eubacterium saburreum		99.23	0.61			
Fusobacterium mortiferum		0.77	0.26			
57435002	**Good identification**					**57435002**
Eubacterium saburreum		100.0	0.48			
57435010	**Very good identification**					**57435010**
Eubacterium saburreum		100.0	0.59			
57435012	**Good identification**					**57435012**
Eubacterium saburreum		100.0	0.47			
57435020	**Good identification**					**57435020**
Eubacterium saburreum		100.0	0.48			
57435022	**Acceptable identification**					**57435022**
Eubacterium saburreum		94.98	0.35			
Parabacteroides distasonis		5.02	0.14			
57435030	**Good identification**					**57435030**
Eubacterium saburreum		100.0	0.47			
57435100	**Very good identification**					**57435100**
Eubacterium saburreum		99.23	0.67			
Fusobacterium mortiferum		0.77	0.33			
57435102	**Good identification**					**57435102**
Eubacterium saburreum		99.12	0.54			
Fusobacterium mortiferum		0.48	0.16			
Parabacteroides distasonis		0.40	0.14			
57435110	**Very good identification**					**57435110**
Eubacterium saburreum		100.0	0.66			
57435112	**Good identification**					**57435112**
Eubacterium saburreum		100.0	0.53			
57435120	**Good identification**					**57435120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		99.60	0.54				
Parabacteroides distasonis		0.40	0.14				
57435122	**Low discrimination**						**57435122**
					CAT		
Parabacteroides distasonis		59.10	0.48	IND 1	76		
Eubacterium saburreum		26.34	0.41		1		
Bacteroides thetaiotaomicron		13.28	0.21		89		
57435130	**Good identification**						**57435130**
Eubacterium saburreum		100.0	0.53				
57435132	**Low discrimination**						**57435132**
					CAT		
Parabacteroides distasonis		54.37	0.43	IND 1	76		
Eubacterium saburreum		36.81	0.40		1		
Bacteroides thetaiotaomicron		8.82	0.14		89		
57435322	**Low discrimination**						**57435322**
Parabacteroides distasonis		50.12	0.56	IND 1			
Bacteroides thetaiotaomicron		49.88	0.40				
57435332	**Low discrimination**						**57435332**
Parabacteroides distasonis		58.18	0.51	IND 1			
Bacteroides thetaiotaomicron		41.82	0.33				
57435522	**Good identification**						**57435522**
					GRA	CAT	
Bacteroides thetaiotaomicron		97.13	0.55	MLT 94	1	89	
Parabacteroides distasonis		2.68	0.44	RHA 93	1	76	
Bifidobacterium longum subsp. longum		0.13	0.17		99	1	
57435532	**Good identification**						**57435532**
					GRA	CAT	
Bacteroides thetaiotaomicron		95.78	0.48	MLT 94	1	89	
Parabacteroides distasonis		3.65	0.39	RHA 93	1	76	
Bifidobacterium longum subsp. longum		0.56	0.21		99	1	
57435622	**Good identification**						**57435622**
Bacteroides thetaiotaomicron		99.69	0.45	MLT 94			
Parabacteroides distasonis		0.31	0.18	LAC 98			
57435632	**Good identification**						**57435632**
Bacteroides thetaiotaomicron		99.58	0.38	MLT 94			
Parabacteroides distasonis		0.43	0.13	LAC 98			
57435702	**Good identification**						**57435702**
Bacteroides thetaiotaomicron		99.38	0.40	MLT 94			
Parabacteroides distasonis		0.62	0.18	NAG 99			
57435720	**Good identification**						**57435720**
					CAT		
Bacteroides thetaiotaomicron		97.52	0.40	MLT 94	89		
Bacteroides vulgatus		1.88	0.08	BGL 99	3		
Parabacteroides distasonis		0.61	0.18		76		
57435722	**Very good identification**						**57435722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		99.33	0.74	MLT 94		
Parabacteroides distasonis		0.62	0.52			
Bacteroides ovatus		0.05	0.17			
57435723	**Good identification**					**57435723**
Bacteroides thetaiotaomicron		99.38	0.40	MLT 94		
Parabacteroides distasonis		0.62	0.18	URE 1		
57435732	**Very good identification**					**57435732**
Bacteroides thetaiotaomicron		99.06	0.67	MLT 94		
Parabacteroides distasonis		0.85	0.48			
Bacteroides ovatus		0.08	0.14			
57435762	**Good identification**					**57435762**
Bacteroides thetaiotaomicron		99.38	0.40	MLT 94		
Parabacteroides distasonis		0.62	0.18	SOR 1		
57437722	**Good identification**					**57437722**
Bacteroides thetaiotaomicron		97.91	0.40	MLT 94		
Bacteroides ovatus		1.48	0.09	MAN 1		
Parabacteroides distasonis		0.61	0.18			
57450000	**Good identification**					**57450000**
Fusobacterium mortiferum		100.0	0.43	IND 1		
57450100	**Good identification**					**57450100**
Fusobacterium mortiferum		100.0	0.50	IND 1		
57451000	**Low discrimination**					**57451000**
Fusobacterium mortiferum		80.30	0.46	IND 1		
Eubacterium saburreum		19.71	0.34			
57451100	**Low discrimination**					**57451100**
Fusobacterium mortiferum		80.30	0.53	IND 1		
Eubacterium saburreum		19.71	0.41			
57451102	**Low discrimination**					**57451102**
Fusobacterium mortiferum		64.22	0.36	IND 1		
Eubacterium saburreum		25.03	0.28	BGL 10		
Prevotella oralis		6.71	0.06			
57451110	**Good identification**					**57451110**
Eubacterium saburreum		95.21	0.39	CEL 1		
Fusobacterium mortiferum		4.79	0.19			
57451722	**Low discrimination**					**57451722**
					CAT	
Bacteroides thetaiotaomicron		63.57	0.37	MLT 94	89	
Bacteroides uniformis		35.22	0.30	TRE 93	5	
Parabacteroides distasonis		0.78	0.20	XYL 96	76	
57455100	**Low discrimination**					**57455100**
Eubacterium saburreum		63.71	0.36	CEL 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		36.29	0.34			
57455122	**Low discrimination**					**57455122**
					CAT	
Bacteroides uniformis		85.52	0.43	MLT 92	5	
Bacteroides fragilis		11.57	0.25	SAL 92	99	
Parabacteroides distasonis		1.49	0.32	ARA 92	76	
57455322	**Acceptable identification**					**57455322**
					CAT	
Bacteroides uniformis		91.24	0.46	MLT 92	5	
Bacteroides thetaiotaomicron		4.64	0.26	SAL 92	89	
Parabacteroides distasonis		3.44	0.40	ARA 92	76	
57455332	**Low discrimination**					**57455332**
					CAT	
Parabacteroides distasonis		38.27	0.35	IND 1	76	
Bacteroides thetaiotaomicron		37.22	0.19		89	
Bacteroides uniformis		19.02	0.11		5	
57455522	**Very good identification**					**57455522**
					CAT	
Bacteroides uniformis		95.47	0.61	MLT 92	5	
Bacteroides thetaiotaomicron		4.34	0.41	SAL 92	89	
Parabacteroides distasonis		0.09	0.28		76	
57455722	**Low discrimination**					**57455722**
					CAT	
Bacteroides uniformis		68.87	0.64	MLT 92	5	
Bacteroides thetaiotaomicron		30.14	0.60	SAL 92	89	
Bacteroides ovatus		0.85	0.34		68	
57455732	**Low discrimination**					**57455732**
					CAT	
Bacteroides thetaiotaomicron		89.78	0.53	MLT 94	89	
Bacteroides uniformis		5.33	0.30	TRE 93	5	
Bacteroides ovatus		4.32	0.31		68	
57465722	**Good identification**					**57465722**
					CAT	
Bacteroides thetaiotaomicron		97.59	0.45	MLT 94	89	
Bacteroides ovatus		2.09	0.17	FRU 99	68	
Bacteroides uniformis		0.17	-.05		5	
57465732	**Good identification**					**57465732**
Bacteroides thetaiotaomicron		96.47	0.38	MLT 94		
Bacteroides ovatus		3.54	0.14	FRU 99		
57470100	**Good identification**					**57470100**
Fusobacterium mortiferum		100.0	0.41	IND 1		
57471000	**Low discrimination**					**57471000**
Fusobacterium mortiferum		64.61	0.38	IND 1		
Eubacterium saburreum		35.39	0.31			
57471100	**Low discrimination**					**57471100**
Fusobacterium mortiferum		64.61	0.44	IND 1		
Eubacterium saburreum		35.39	0.38			
57471110	**Good identification**					**57471110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.36	CEL 1		
57471522	**Good identification**					**57471522**
					CAT	
Bacteroides thetaiotaomicron		97.44	0.37	MLT 94	89	
Parabacteroides distasonis		1.81	0.23	XYL 96	76	
Bacteroides uniformis		0.40	-.07	RHA 93	5	
57471722	**Good identification**					**57471722**
Bacteroides thetaiotaomicron		99.03	0.56	MLT 94		
Bacteroides ovatus		0.51	0.17	XYL 96		
Parabacteroides distasonis		0.42	0.31			
57471732	**Good identification**					**57471732**
Bacteroides thetaiotaomicron		98.55	0.49	MLT 94		
Bacteroides ovatus		0.88	0.14	XYL 96		
Parabacteroides distasonis		0.57	0.27			
57474722	**Good identification**					**57474722**
					CAT	
Bacteroides thetaiotaomicron		97.59	0.45	MLT 94	89	
Bacteroides ovatus		2.09	0.17	GAL 99	68	
Bacteroides uniformis		0.17	-.05		5	
57474732	**Good identification**					**57474732**
Bacteroides thetaiotaomicron		96.04	0.38	MLT 94		
Bacteroides ovatus		3.52	0.14	GAL 99		
Clostridium clostridioforme		0.44	0.09			
57475122	**Low discrimination**					**57475122**
					CAT	
Parabacteroides distasonis		48.26	0.43	IND 1	76	
Bacteroides thetaiotaomicron		42.79	0.26		89	
Bacteroides uniformis		6.16	0.09		5	
57475132	**Low discrimination**					**57475132**
Parabacteroides distasonis		59.15	0.39	IND 1		
Bacteroides thetaiotaomicron		37.88	0.19			
Bacteroides ovatus		2.97	0.00			
57475322	**Low discrimination**					**57475322**
Bacteroides thetaiotaomicron		74.90	0.45	MLT 94		
Parabacteroides distasonis		19.07	0.51	ARA 99		
Bacteroides ovatus		4.90	0.25			
57475332	**Low discrimination**					**57475332**
Bacteroides thetaiotaomicron		68.27	0.38	MLT 94		
Parabacteroides distasonis		24.07	0.47	ARA 99		
Bacteroides ovatus		7.66	0.22			
57475522	**Very good identification**					**57475522**
					CAT	
Bacteroides thetaiotaomicron		96.28	0.60	MLT 94	89	
Bacteroides uniformis		1.61	0.27	RHA 93	5	
Bacteroides ovatus		1.44	0.29		68	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57475532	**Good identification**					**57475532**
Bacteroides thetaiotaomicron		96.55	0.53	MLT 94		
Bacteroides ovatus		2.47	0.26	RHA 93		
Parabacteroides distasonis		0.93	0.35			
57475622	**Good identification**					**57475622**
					CAT	
Bacteroides thetaiotaomicron		98.71	0.51	MLT 94	89	
Bacteroides ovatus		1.04	0.17	LAC 98	68	
Bacteroides uniformis		0.17	0.01		5	
57475632	**Good identification**					**57475632**
Bacteroides thetaiotaomicron		98.02	0.44	MLT 94		
Bacteroides ovatus		1.78	0.14	LAC 98		
Clostridium clostridioforme		0.20	0.08			
57475702	**Good identification**					**57475702**
					CAT	
Bacteroides thetaiotaomicron		97.48	0.45	MLT 94	89	
Bacteroides ovatus		2.08	0.17	NAG 99	68	
Bacteroides uniformis		0.17	-.05		5	
57475712	**Good identification**					**57475712**
Bacteroides thetaiotaomicron		95.20	0.38	MLT 94		
Bacteroides ovatus		3.49	0.14	NAG 99		
Clostridium clostridioforme		1.31	0.17			
57475720	**Good identification**					**57475720**
					CAT	
Bacteroides thetaiotaomicron		97.59	0.45	MLT 94	89	
Bacteroides ovatus		2.09	0.17	BGL 99	68	
Bacteroides uniformis		0.17	-.05		5	
57475722	**Very good identification**					**57475722**
					CAT	
Bacteroides thetaiotaomicron		97.59	0.80	MLT 94	89	
Bacteroides ovatus		2.09	0.51		68	
Bacteroides uniformis		0.17	0.30		5	
57475723	**Good identification**					**57475723**
					CAT	
Bacteroides thetaiotaomicron		97.59	0.45	MLT 94	89	
Bacteroides ovatus		2.09	0.17	URE 1	68	
Bacteroides uniformis		0.17	-.05		5	
57475730	**Good identification**					**57475730**
Bacteroides thetaiotaomicron		96.47	0.38	MLT 94		
Bacteroides ovatus		3.54	0.14	BGL 99		
57475732	**Very good identification**					**57475732**
Bacteroides thetaiotaomicron		96.25	0.73	MLT 94		
Bacteroides ovatus		3.53	0.48			
Parabacteroides distasonis		0.21	0.43			
57475733	**Good identification**					**57475733**
Bacteroides thetaiotaomicron		96.47	0.38	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		3.54	0.14	URE 1			
57475762	**Acceptable identification**						**57475762**
					CAT		
Bacteroides thetaiotaomicron		91.62	0.45	MLT 94	89		
Bacteroides ovatus		8.08	0.27	SOR 1	68		
Bacteroides uniformis		0.16	-.05		5		
57475772	**Low discrimination**						**57475772**
Bacteroides thetaiotaomicron		86.87	0.38	MLT 94			
Bacteroides ovatus		13.13	0.24	SOR 1			
57477722	**Low discrimination**						**57477722**
					CAT		
Bacteroides thetaiotaomicron		58.52	0.45	MLT 94	89		
Bacteroides ovatus		41.29	0.43	MAN 1	68		
Bacteroides uniformis		0.10	-.05		5		
57477732	**Low discrimination**						**57477732**
Bacteroides ovatus		54.74	0.40	MLT 99			
Bacteroides thetaiotaomicron		45.26	0.38	SAL 88			
57500010	**Good identification**						**57500010**
					SPO		
Eubacterium saburreum		96.84	0.35	GAL 99	1		
Clostridium bifermentans		3.16	-.03		99		
57500100	**Good identification**						**57500100**
					SPO		
Eubacterium saburreum		98.87	0.43	GAL 99	1		
Clostridium bifermentans		1.13	-.03		99		
57500110	**Good identification**						**57500110**
Eubacterium saburreum		100.0	0.41	GAL 99			
57501000	**Very good identification**						**57501000**
					SPO	GRA	
Eubacterium saburreum		99.05	0.71		1	85	
Prevotella intermedia		0.93	0.26		1	1	
Clostridium bifermentans		0.03	-.03		99	99	
57501001	**Good identification**						**57501001**
Eubacterium saburreum		100.0	0.37	URE 1			
57501002	**Very good identification**						**57501002**
Eubacterium saburreum		100.0	0.58				
57501010	**Very good identification**						**57501010**
Eubacterium saburreum		100.0	0.69				
57501011	**Good identification**						**57501011**
Eubacterium saburreum		100.0	0.35	URE 1			
57501012	**Very good identification**						**57501012**
Eubacterium saburreum		100.0	0.56				
57501020	**Very good identification**						**57501020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		98.14	0.58				
Prevotella melaninogenica		1.87	0.12				
57501022	**Good identification**						**57501022**
Eubacterium saburreum		97.12	0.45				
Prevotella melaninogenica		1.99	0.00				
Prevotella oralis		0.89	-.03				
57501030	**Very good identification**						**57501030**
Eubacterium saburreum		100.0	0.56				
57501032	**Good identification**						**57501032**
Eubacterium saburreum		100.0	0.43				
57501040	**Good identification**						**57501040**
Eubacterium saburreum		100.0	0.37	SOR 1			
57501050	**Good identification**						**57501050**
Eubacterium saburreum		100.0	0.35	SOR 1			
57501100	**Very good identification**						**57501100**
					COC	GRA	
Eubacterium saburreum		99.83	0.77		1	85	
Prevotella melaninogenica		0.14	0.12		1	1	
Blautia hansenii		0.02	-.03		99	99	
57501101	**Low discrimination**						**57501101**
					SPO		
Eubacterium saburreum		80.65	0.43	URE 1	1		
Clostridium chauvoei		19.35	0.22		99		
57501102	**Very good identification**						**57501102**
					CAT		
Eubacterium saburreum		99.30	0.64		1		
Capnocytophaga ochracea		0.49	0.12		90		
Prevotella melaninogenica		0.15	0.00		1		
57501110	**Very good identification**						**57501110**
Eubacterium saburreum		100.0	0.75				
57501111	**Good identification**						**57501111**
Eubacterium saburreum		100.0	0.41	URE 1			
57501112	**Very good identification**						**57501112**
Eubacterium saburreum		100.0	0.63				
57501120	**Low discrimination**						**57501120**
					CAT		
Eubacterium saburreum		55.18	0.64		1		
Prevotella melaninogenica		44.49	0.47		1		
Capnocytophaga ochracea		0.27	0.12		90		
57501122	**Low discrimination**						**57501122**
					CAT		
Eubacterium saburreum		45.44	0.51		1		
Prevotella melaninogenica		39.54	0.34		1		
Capnocytophaga ochracea		11.34	0.29		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57501130	**Very good identification**						**57501130**
					GRA		
Eubacterium saburreum		98.92	0.63		85		
Prevotella melaninogenica		0.98	0.12		1		
Bifidobacterium longum subsp. longum		0.10	0.07		99		
57501132	**Good identification**						**57501132**
					CAT		
Eubacterium saburreum		95.22	0.50		1		
Capnocytophaga ochracea		3.23	0.12		90		
Prevotella melaninogenica		1.02	0.00		1		
57501140	**Good identification**						**57501140**
Eubacterium saburreum		100.0	0.43	SOR 1			
57501150	**Good identification**						**57501150**
Eubacterium saburreum		100.0	0.41	SOR 1			
57501200	**Good identification**						**57501200**
Eubacterium saburreum		100.0	0.37	RHA 1			
57501210	**Good identification**						**57501210**
Eubacterium saburreum		100.0	0.35	RHA 1			
57501300	**Good identification**						**57501300**
Eubacterium saburreum		100.0	0.43	RHA 1			
57501310	**Good identification**						**57501310**
Eubacterium saburreum		100.0	0.41	RHA 1			
57501400	**Good identification**						**57501400**
Eubacterium saburreum		100.0	0.37	ARA 1			
57501410	**Good identification**						**57501410**
Eubacterium saburreum		100.0	0.35	ARA 1			
57501500	**Low discrimination**						**57501500**
Eubacterium saburreum		83.09	0.43	ARA 1			
Bifidobacterium longum subsp. longum		16.91	0.27				
57501510	**Low discrimination**						**57501510**
Eubacterium saburreum		70.25	0.41	ARA 1			
Bifidobacterium longum subsp. longum		29.75	0.31				
57501520	**Low discrimination**						**57501520**
					GRA		
Bifidobacterium longum subsp. longum		69.96	0.37	IND 1	99		
Eubacterium saburreum		15.16	0.30	FRU 90	85		
Prevotella melaninogenica		12.22	0.12		1		
57501522	**Low discrimination**						**57501522**
					CAT	GRA	
Bifidobacterium longum subsp. longum		79.15	0.37	IND 1	1	99	
Capnocytophaga ochracea		13.19	0.16	FRU 90	90	1	
Eubacterium saburreum		3.03	0.17		1	85	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
57501530	**Acceptable identification**						**57501530**
Bifidobacterium longum subsp. longum		90.57	0.41	IND 1			
Eubacterium saburreum		9.43	0.28	FRU 90			
57501532	**Good identification**						**57501532**
					CAT	GRA	
Bifidobacterium longum subsp. longum		97.16	0.41	IND 1	1	99	
Eubacterium saburreum		1.79	0.16	FRU 90	1	85	
Capnocytophaga ochracea		1.06	-.01		90	1	
57503000	**Good identification**						**57503000**
Eubacterium saburreum		100.0	0.37	MAN 1			
57503010	**Good identification**						**57503010**
Eubacterium saburreum		100.0	0.35	MAN 1			
57503100	**Good identification**						**57503100**
Eubacterium saburreum		100.0	0.43	MAN 1			
57503110	**Good identification**						**57503110**
Eubacterium saburreum		100.0	0.41	MAN 1			
57504100	**Good identification**						**57504100**
Eubacterium saburreum		100.0	0.38	GAL 99			
57504110	**Good identification**						**57504110**
Eubacterium saburreum		100.0	0.37	GAL 99			
57505000	**Very good identification**						**57505000**
Eubacterium saburreum		100.0	0.66				
57505002	**Good identification**						**57505002**
Eubacterium saburreum		100.0	0.53				
57505010	**Very good identification**						**57505010**
Eubacterium saburreum		100.0	0.65				
57505012	**Good identification**						**57505012**
Eubacterium saburreum		100.0	0.52				
57505020	**Good identification**						**57505020**
Eubacterium saburreum		100.0	0.53				
57505022	**Acceptable identification**						**57505022**
					CAT		
Eubacterium saburreum		94.63	0.40		1		
Bacteroides fragilis		5.37	-.03		99		
57505030	**Good identification**						**57505030**
Eubacterium saburreum		100.0	0.52				
57505032	**Good identification**						**57505032**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.39				
57505100	**Very good identification**						**57505100**
Eubacterium saburreum		100.0	0.72				
57505101	**Good identification**						**57505101**
Eubacterium saburreum		100.0	0.38	URE 1			
57505102	**Very good identification**						**57505102**
					CAT		
Eubacterium saburreum		99.35	0.59		1		
Bacteroides fragilis		0.43	-.03		99		
Capnocytophaga ochracea		0.23	0.02		90		
57505110	**Very good identification**						**57505110**
Eubacterium saburreum		100.0	0.71				
57505111	**Good identification**						**57505111**
Eubacterium saburreum		100.0	0.37	URE 1			
57505112	**Very good identification**						**57505112**
Eubacterium saburreum		100.0	0.58				
57505120	**Very good identification**						**57505120**
					CAT		
Eubacterium saburreum		95.72	0.59		1		
Bacteroides fragilis		2.14	0.10		99		
Prevotella melaninogenica		1.45	0.12		1		
57505130	**Very good identification**						**57505130**
Eubacterium saburreum		99.46	0.58				
Bifidobacterium longum subsp. longum		0.54	0.15				
57505132	**Acceptable identification**						**57505132**
					CAT	GRA	
Eubacterium saburreum		92.90	0.45		1	85	
Bifidobacterium longum subsp. longum		2.87	0.15		1	99	
Bacteroides fragilis		2.76	-.03		99	1	
57505140	**Good identification**						**57505140**
Eubacterium saburreum		100.0	0.38	SOR 1			
57505150	**Good identification**						**57505150**
Eubacterium saburreum		100.0	0.37	SOR 1			
57505300	**Good identification**						**57505300**
Eubacterium saburreum		100.0	0.38	RHA 1			
57505310	**Good identification**						**57505310**
Eubacterium saburreum		100.0	0.37	RHA 1			
57505500	**Low discrimination**						**57505500**
Bifidobacterium longum subsp. longum		53.14	0.35	IND 1			
Eubacterium saburreum		46.86	0.38	FRU 90			

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
57505502 Low discrimination						**57505502**
Bifidobacterium longum subsp. longum	84.07	0.35	IND 1			
Eubacterium saburreum	13.08	0.25	FRU 90			
Bifidobacterium adolescentis	2.84	0.12				
57505510 Low discrimination						**57505510**
Bifidobacterium longum subsp. longum	70.24	0.39	IND 1			
Eubacterium saburreum	29.76	0.37	FRU 90			
57505512 Acceptable identification						**57505512**
Bifidobacterium longum subsp. longum	90.99	0.39	IND 1			
Eubacterium saburreum	6.80	0.24	FRU 90			
Bifidobacterium adolescentis	2.21	0.13				
57505520 Low discrimination						**57505520**
				GRA		
Bifidobacterium longum subsp. longum	86.52	0.46	IND 1	99		
Bacteroides vulgatus	10.11	0.15	FRU 90	1		
Eubacterium saburreum	3.37	0.25		85		
57505522 Low discrimination						**57505522**
				GRA	CAT	
Bifidobacterium longum subsp. longum	76.37	0.46	IND 1	99	1	
Bacteroides thetaiotaomicron	15.44	0.22	FRU 90	1	89	
Bacteroides uniformis	5.35	0.11		1	5	
57505530 Good identification						**57505530**
Bifidobacterium longum subsp. longum	98.17	0.50	IND 1			
Eubacterium saburreum	1.84	0.24	FRU 90			
57505532 Good identification						**57505532**
				CAT	GRA	
Bifidobacterium longum subsp. longum	95.29	0.50	IND 1	1	99	
Bacteroides thetaiotaomicron	4.40	0.15	FRU 90	89	1	
Eubacterium saburreum	0.31	0.11		1	85	
57505722 Low discrimination						**57505722**
				CAT		
Bacteroides thetaiotaomicron	82.65	0.41	FRU 99	89		
Bacteroides eggerthii	13.08	0.21	TRE 93	93		
Bacteroides uniformis	2.98	0.14		5		
57507100 Good identification						**57507100**
Eubacterium saburreum	100.0	0.38	MAN 1			
57507110 Good identification						**57507110**
Eubacterium saburreum	100.0	0.37	MAN 1			
57510000 Very good identification						**57510000**
Prevotella intermedia	95.83	0.61	ESL 1			
Eubacterium saburreum	3.14	0.45				
Fusobacterium mortiferum	0.79	0.36				
57510010 Low discrimination						**57510010**
Eubacterium saburreum	72.61	0.43	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella intermedia		27.39	0.26			
57510100	**Low discrimination**					**57510100**
Eubacterium saburreum		72.03	0.51	GAL 99		
Fusobacterium mortiferum		18.14	0.42			
Prevotella intermedia		9.53	0.26			
57510102	**Low discrimination**					**57510102**
					CAT	
Eubacterium saburreum		74.60	0.38	GAL 99	1	
Capnocytophaga ochracea		12.04	0.12		90	
Fusobacterium mortiferum		11.83	0.26		5	
57510110	**Good identification**					**57510110**
Eubacterium saburreum		100.0	0.49	GAL 99		
57510112	**Good identification**					**57510112**
Eubacterium saburreum		100.0	0.37	GAL 99		
57510130	**Good identification**					**57510130**
Eubacterium saburreum		97.10	0.37	GAL 99		
Prevotella melaninogenica		2.90	-.06			
57511000	**Low discrimination**					**57511000**
Eubacterium saburreum		76.16	0.79			
Prevotella intermedia		23.50	0.61			
Fusobacterium mortiferum		0.29	0.39			
57511001	**Low discrimination**					**57511001**
Eubacterium saburreum		76.42	0.45	URE 1		
Prevotella intermedia		23.58	0.26			
57511002	**Very good identification**					**57511002**
					CAT	
Eubacterium saburreum		97.23	0.66		1	
Prevotella intermedia		1.72	0.26		1	
Capnocytophaga ochracea		0.37	0.12		90	
57511010	**Very good identification**					**57511010**
Eubacterium saburreum		99.62	0.77			
Prevotella intermedia		0.38	0.26			
57511011	**Good identification**					**57511011**
Eubacterium saburreum		100.0	0.43	URE 1		
57511012	**Very good identification**					**57511012**
Eubacterium saburreum		100.0	0.64			
57511020	**Good identification**					**57511020**
Eubacterium saburreum		91.49	0.66			
Prevotella melaninogenica		5.22	0.29			
Prevotella intermedia		1.62	0.26			
57511022	**Low discrimination**					**57511022**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		62.31	0.53		1		
Prevotella oralis		18.89	0.32		1		
Capnocytophaga ochracea		12.09	0.29		90		
57511030	**Very good identification**						**57511030**
Eubacterium saburreum		99.93	0.64				
Prevotella melaninogenica		0.07	-.06				
57511032	**Good identification**						**57511032**
					CAT		
Eubacterium saburreum		97.08	0.52		1		
Capnocytophaga ochracea		2.56	0.12		90		
Prevotella oralis		0.36	-.03		1		
57511040	**Low discrimination**						**57511040**
Eubacterium saburreum		76.42	0.45	SOR 1			
Prevotella intermedia		23.58	0.26				
57511050	**Good identification**						**57511050**
Eubacterium saburreum		100.0	0.43	SOR 1			
57511100	**Very good identification**						**57511100**
Eubacterium saburreum		98.65	0.85				
Prevotella melaninogenica		0.43	0.29				
Fusobacterium mortiferum		0.38	0.45				
57511101	**Low discrimination**						**57511101**
					SPO		
Eubacterium saburreum		83.03	0.51	URE 1	1		
Clostridium chauvoei		15.50	0.29		99		
Actinomyces naeslundii		0.80	0.30		1		
57511102	**Low discrimination**						**57511102**
					CAT	GRA	
Eubacterium saburreum		82.53	0.72		1	85	
Capnocytophaga ochracea		13.32	0.47		90	1	
Bifidobacterium longum subsp. infantis		1.69	0.27		1	99	
57511103	**Low discrimination**						**57511103**
					SPO	CAT	
Eubacterium saburreum		70.28	0.38	URE 1	1	1	
Clostridium chauvoei		13.12	0.16		99	1	
Capnocytophaga ochracea		11.34	0.12		1	90	
57511110	**Very good identification**						**57511110**
					CAT	GRA	
Eubacterium saburreum		99.93	0.84		1	85	
Capnocytophaga ochracea		0.04	0.14		90	1	
Bifidobacterium longum subsp. longum		0.01	0.13		1	99	
57511111	**Good identification**						**57511111**
Eubacterium saburreum		100.0	0.49	URE 1			
57511112	**Very good identification**						**57511112**
					CAT	GRA	
Eubacterium saburreum		97.64	0.71		1	85	
Capnocytophaga ochracea		2.14	0.30		90	1	
Bifidobacterium adolescentis		0.13	0.19		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57511113	**Good identification**					**57511113**
Eubacterium saburreum		100.0	0.37	URE 1		
57511120	**Low discrimination**					**57511120**
					CAT	
Prevotella melaninogenica		66.33	0.63	IND 1	1	
Eubacterium saburreum		27.43	0.72		1	
Capnocytophaga ochracea		4.43	0.47		90	
57511122	**Low discrimination**					**57511122**
					CAT	
Capnocytophaga ochracea		56.20	0.63	IND 1	90	
Prevotella melaninogenica		17.82	0.51		1	
Prevotella oralis		13.90	0.52		1	
57511130	**Good identification**					**57511130**
					CAT	
Eubacterium saburreum		94.77	0.71		1	
Prevotella melaninogenica		2.83	0.29		1	
Capnocytophaga ochracea		2.08	0.30		90	
57511131	**Good identification**					**57511131**
Eubacterium saburreum		97.10	0.37	URE 1		
Prevotella melaninogenica		2.90	-.06			
57511132	**Low discrimination**					**57511132**
					CAT	
Capnocytophaga ochracea		50.56	0.47	IND 1	90	
Eubacterium saburreum		45.22	0.58	MLZ 10	1	
Prevotella melaninogenica		1.46	0.16		1	
57511140	**Good identification**					**57511140**
					CAT	
Eubacterium saburreum		95.54	0.51	SOR 1	1	
Capnocytophaga ochracea		3.33	0.14		90	
Prevotella melaninogenica		0.41	-.06		1	
57511150	**Good identification**					**57511150**
					CAT	
Eubacterium saburreum		99.53	0.49	SOR 1	1	
Capnocytophaga ochracea		0.47	-.03		90	
57511152	**Low discrimination**					**57511152**
					CAT	
Eubacterium saburreum		80.57	0.37	SOR 1	1	
Capnocytophaga ochracea		19.43	0.14		90	
57511162	**Acceptable identification**					**57511162**
					CAT	
Capnocytophaga ochracea		91.76	0.47	IND 1	90	
Prevotella melaninogenica		2.65	0.16	SOR 10	1	
Prevotella oralis		2.06	0.18		1	
57511170	**Low discrimination**					**57511170**
					CAT	
Eubacterium saburreum		78.68	0.37	SOR 1	1	
Capnocytophaga ochracea		18.97	0.14		90	
Prevotella melaninogenica		2.35	-.06		1	
57511200	**Low discrimination**					**57511200**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		76.42	0.45	RHA 1			
Prevotella intermedia		23.58	0.26				
57511210	**Good identification**						**57511210**
Eubacterium saburreum		100.0	0.43	RHA 1			
57511300	**Good identification**						**57511300**
Eubacterium saburreum		98.47	0.51	RHA 1			
Prevotella oralis		0.73	0.02				
Prevotella melaninogenica		0.42	-.06				
57511302	**Low discrimination**						**57511302**
					CAT		
Eubacterium saburreum		50.40	0.38	RHA 1	1		
Prevotella oralis		40.43	0.24		1		
Capnocytophaga ochracea		8.14	0.12		90		
57511310	**Good identification**						**57511310**
Eubacterium saburreum		100.0	0.49	RHA 1			
57511312	**Good identification**						**57511312**
Eubacterium saburreum		100.0	0.37	RHA 1			
57511322	**Low discrimination**						**57511322**
					CAT		
Prevotella oralis		80.59	0.46	IND 1	1		
Capnocytophaga ochracea		7.68	0.29		90		
Parabacteroides distasonis		4.83	0.38		76		
57511330	**Good identification**						**57511330**
Eubacterium saburreum		97.10	0.37	RHA 1			
Prevotella melaninogenica		2.90	-.06				
57511400	**Low discrimination**						**57511400**
					GRA		
Eubacterium saburreum		74.28	0.45	ARA 1	85		
Prevotella intermedia		22.92	0.26		1		
Lactobacillus fermentum		1.73	0.20		99		
57511410	**Good identification**						**57511410**
Eubacterium saburreum		97.09	0.43	ARA 1			
Bifidobacterium longum subsp. longum		2.91	0.13				
57511500	**Low discrimination**						**57511500**
					CAT	GRA	
Eubacterium saburreum		59.16	0.51	ARA 1	1	85	
Bifidobacterium longum subsp. longum		36.12	0.43		1	99	
Capnocytophaga ochracea		3.27	0.17		90	1	
57511502	**Low discrimination**						**57511502**
					CAT	GRA	
Bifidobacterium longum subsp. longum		43.20	0.43	IND 1	1	99	
Capnocytophaga ochracea		35.21	0.34		90	1	
Eubacterium saburreum		12.49	0.38		1	85	
57511510	**Low discrimination**						**57511510**
					CAT	GRA	
Bifidobacterium longum subsp. longum		55.78	0.47	IND 1	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		43.89	0.49		1	85	
Capnocytophaga ochracea		0.33	0.01		90	1	
57511512	**Low discrimination**						**57511512**
Bifidobacterium longum subsp. longum		76.23	0.47	IND 1			
Eubacterium saburreum		10.59	0.37				
Bifidobacterium adolescentis		9.13	0.33				
57511520	**Low discrimination**						**57511520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		64.89	0.54	IND 1	99	1	
Capnocytophaga ochracea		13.22	0.34		1	90	
Prevotella melaninogenica		11.34	0.29		1	1	
57511522	**Low discrimination**						**57511522**
					GRA	CAT	
Capnocytophaga ochracea		56.76	0.50	IND 1	1	90	
Bifidobacterium longum subsp. longum		30.95	0.54		99	1	
Bacteroides thetaiotaomicron		8.60	0.32		1	89	
57511530	**Very good identification**						**57511530**
					CAT	GRA	
Bifidobacterium longum subsp. longum		95.32	0.58	IND 1	1	99	
Eubacterium saburreum		3.31	0.37		1	85	
Capnocytophaga ochracea		1.27	0.17		90	1	
57511532	**Low discrimination**						**57511532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		83.79	0.58	IND 1	99	1	
Capnocytophaga ochracea		10.03	0.34		1	90	
Bacteroides thetaiotaomicron		5.32	0.25		1	89	
57511720	**Acceptable identification**						**57511720**
					GRA	CAT	
Bacteroides vulgatus		93.26	0.37	IND 1	1	3	
Bacteroides thetaiotaomicron		4.91	0.17	XYL 98	1	89	
Bifidobacterium longum subsp. longum		1.33	0.20		99	1	
57511722	**Good identification**						**57511722**
					CAT		
Bacteroides thetaiotaomicron		96.78	0.52	TRE 93	89		
Parabacteroides distasonis		1.06	0.34	XYL 96	76		
Bacteroides uniformis		0.85	0.14		5		
57511732	**Good identification**						**57511732**
					GRA	CAT	
Bacteroides thetaiotaomicron		96.85	0.45	TRE 93	1	89	
Parabacteroides distasonis		1.47	0.29	XYL 96	1	76	
Bifidobacterium longum subsp. longum		1.16	0.24		99	1	
57513000	**Low discrimination**						**57513000**
					CAT	GRA	
Eubacterium saburreum		74.44	0.45	MAN 1	1	85	
Prevotella intermedia		22.97	0.26		1	1	
Propionibacterium granulosum		1.83	0.16		90	99	
57513010	**Good identification**						**57513010**
Eubacterium saburreum		100.0	0.43	MAN 1			
57513100	**Good identification**						**57513100**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		97.99	0.51	MAN 1	85		
Actinomyces naeslundii		1.22	0.32		99		
Prevotella melaninogenica		0.42	-.06		1		
57513102	**Low discrimination**						**57513102**
					GRA	CAT	
Eubacterium saburreum		74.44	0.38	MAN 1	85	1	
Capnocytophaga ochracea		12.02	0.12		1	90	
Actinomyces naeslundii		5.25	0.32		99	25	
57513110	**Good identification**						**57513110**
Eubacterium saburreum		100.0	0.49	MAN 1			
57513112	**Acceptable identification**						**57513112**
Eubacterium saburreum		90.46	0.37	MAN 1			
Bifidobacterium adolescentis		9.54	0.18				
57513130	**Good identification**						**57513130**
Eubacterium saburreum		97.10	0.37	MAN 1			
Prevotella melaninogenica		2.90	-.06				
57514000	**Low discrimination**						**57514000**
Eubacterium saburreum		62.18	0.40	GAL 99			
Prevotella intermedia		35.63	0.26				
Fusobacterium mortiferum		2.19	0.17				
57514010	**Good identification**						**57514010**
Eubacterium saburreum		100.0	0.39	GAL 99			
57514100	**Good identification**						**57514100**
Eubacterium saburreum		96.60	0.46	GAL 99			
Fusobacterium mortiferum		3.40	0.23				
57514110	**Good identification**						**57514110**
Eubacterium saburreum		100.0	0.45	GAL 99			
57514722	**Good identification**						**57514722**
					CAT		
Bacteroides thetaiotaomicron		95.17	0.41	TRE 93	89		
Bacteroides uniformis		3.43	0.14	GAL 99	5		
Parabacteroides distasonis		0.39	0.16		76		
57515000	**Very good identification**						**57515000**
Eubacterium saburreum		99.35	0.74				
Prevotella intermedia		0.58	0.26				
Fusobacterium mortiferum		0.05	0.20				
57515001	**Good identification**						**57515001**
Eubacterium saburreum		100.0	0.40	URE 1			
57515002	**Very good identification**						**57515002**
					CAT		
Eubacterium saburreum		99.35	0.61		1		
Bacteroides fragilis		0.33	-.03		99		
Capnocytophaga ochracea		0.18	0.02		90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57515010	**Very good identification**						**57515010**
Eubacterium saburreum		100.0	0.73				
57515011	**Good identification**						**57515011**
Eubacterium saburreum		100.0	0.39	URE 1			
57515012	**Very good identification**						**57515012**
Eubacterium saburreum		100.0	0.60				
57515020	**Very good identification**						**57515020**
					CAT		
Eubacterium saburreum		97.85	0.61		1		
Bacteroides fragilis		1.70	0.10		99		
Capnocytophaga ochracea		0.17	0.02		90		
57515030	**Very good identification**						**57515030**
Eubacterium saburreum		100.0	0.60				
57515032	**Good identification**						**57515032**
					CAT		
Eubacterium saburreum		96.59	0.47		1		
Bacteroides fragilis		2.23	-.03		99		
Capnocytophaga ochracea		1.18	0.02		90		
57515040	**Good identification**						**57515040**
Eubacterium saburreum		100.0	0.40	SOR 1			
57515050	**Good identification**						**57515050**
Eubacterium saburreum		100.0	0.39	SOR 1			
57515100	**Very good identification**						**57515100**
					CAT		
Eubacterium saburreum		99.54	0.81		1		
Capnocytophaga ochracea		0.15	0.20		90		
Bacteroides fragilis		0.13	0.10		99		
57515101	**Good identification**						**57515101**
Eubacterium saburreum		100.0	0.46	URE 1			
57515102	**Low discrimination**						**57515102**
					CAT		
Eubacterium saburreum		78.93	0.68		1		
Bacteroides fragilis		11.18	0.32		99		
Capnocytophaga ochracea		5.92	0.36		90		
57515110	**Very good identification**						**57515110**
					CAT	GRA	
Eubacterium saburreum		99.91	0.79		1	85	
Bifidobacterium longum subsp. longum		0.07	0.21		1	99	
Capnocytophaga ochracea		0.02	0.04		90	1	
57515111	**Good identification**						**57515111**
Eubacterium saburreum		100.0	0.45	URE 1			
57515112	**Very good identification**						**57515112**
					CAT	GRA	
Eubacterium saburreum		96.88	0.66		1	85	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		1.57	0.33		1	99	
Capnocytophaga ochracea		0.99	0.20		90	1	
57515120	**Low discrimination**						**57515120**
					CAT		
Eubacterium saburreum		51.19	0.68		1		
Bacteroides fragilis		37.80	0.44		99		
Bacteroides vulgatus		3.86	0.30		3		
57515122	**Good identification**						**57515122**
					CAT		
Bacteroides fragilis		92.56	0.66	IND 1	99		
Capnocytophaga ochracea		4.45	0.53		90		
Eubacterium saburreum		1.17	0.55		1		
57515130	**Very good identification**						**57515130**
					CAT	GRA	
Eubacterium saburreum		96.42	0.66		1	85	
Bifidobacterium longum subsp. longum		1.58	0.32		1	99	
Capnocytophaga ochracea		0.98	0.20		90	1	
57515132	**Low discrimination**						**57515132**
					CAT		
Eubacterium saburreum		35.15	0.53		1		
Bacteroides fragilis		34.49	0.32		99		
Capnocytophaga ochracea		18.24	0.36		90		
57515140	**Good identification**						**57515140**
					CAT		
Eubacterium saburreum		98.41	0.46	SOR 1	1		
Capnocytophaga ochracea		1.59	0.04		90		
57515150	**Good identification**						**57515150**
Eubacterium saburreum		100.0	0.45	SOR 1			
57515200	**Good identification**						**57515200**
Eubacterium saburreum		100.0	0.40	RHA 1			
57515210	**Good identification**						**57515210**
Eubacterium saburreum		100.0	0.39	RHA 1			
57515300	**Low discrimination**						**57515300**
					GRA		
Eubacterium saburreum		74.77	0.46	RHA 1	85		
Eubacterium contortum		16.29	0.24		99		
Bacteroides vulgatus		8.95	0.13		1		
57515310	**Good identification**						**57515310**
Eubacterium saburreum		100.0	0.45	RHA 1			
57515320	**Good identification**						**57515320**
					CAT		
Bacteroides vulgatus		96.11	0.47	IND 1	3		
Eubacterium saburreum		1.43	0.33	ARA 93	1		
Bacteroides fragilis		1.06	0.10		99		
57515322	**Low discrimination**						**57515322**
Bacteroides thetaiotaomicron		35.50	0.41	TRE 93			
Parabacteroides distasonis		23.54	0.54	ARA 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides fragilis		21.64	0.32				
57515400	**Acceptable identification**						**57515400**
Eubacterium saburreum		90.45	0.40	ARA 1			
Bifidobacterium longum subsp. longum		7.25	0.17				
Lactobacillus fermentum		2.29	0.16				
57515410	**Low discrimination**						**57515410**
Eubacterium saburreum		85.70	0.39	ARA 1			
Bifidobacterium longum subsp. longum		14.30	0.21				
57515500	**Low discrimination**						**57515500**
					GRA		
Bifidobacterium longum subsp. longum		73.40	0.52	IND 1	99		
Eubacterium saburreum		21.58	0.46		85		
Bacteroides vulgatus		3.81	0.15		1		
57515502	**Low discrimination**						**57515502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		61.52	0.52	IND 1	99	1	
Bifidobacterium adolescentis		22.89	0.46		99	1	
Bacteroides thetaiotaomicron		5.53	0.22		1	89	
57515510	**Low discrimination**						**57515510**
Bifidobacterium longum subsp. longum		87.42	0.56	IND 1			
Eubacterium saburreum		12.35	0.45				
Bifidobacterium adolescentis		0.24	0.13				
57515512	**Low discrimination**						**57515512**
Bifidobacterium longum subsp. longum		75.72	0.56	IND 1			
Bifidobacterium adolescentis		20.22	0.47				
Eubacterium saburreum		1.89	0.32				
57515520	**Low discrimination**						**57515520**
					GRA	CAT	
Bacteroides vulgatus		54.56	0.50	IND 1	1	3	
Bifidobacterium longum subsp. longum		42.45	0.62	RHA 90	99	1	
Bacteroides thetaiotaomicron		0.95	0.22		1	89	
57515522	**Low discrimination**						**57515522**
					GRA	CAT	
Bacteroides thetaiotaomicron		50.62	0.56	TRE 93	1	89	
Bifidobacterium longum subsp. longum		22.77	0.62	RHA 93	99	1	
Bacteroides uniformis		17.53	0.45		1	5	
57515530	**Very good identification**						**57515530**
					GRA		
Bifidobacterium longum subsp. longum		98.03	0.66	IND 1	99		
Bacteroides vulgatus		0.75	0.15		1		
Eubacterium saburreum		0.61	0.32		85		
57515532	**Low discrimination**						**57515532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		64.89	0.66	IND 1	99	1	
Bacteroides thetaiotaomicron		32.95	0.49		1	89	
Parabacteroides distasonis		0.83	0.38		1	76	
57515622	**Good identification**						**57515622**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		95.86	0.46	TRE 93	89		
Bacteroides uniformis		3.45	0.19	LAC 98	5		
Parabacteroides distasonis		0.20	0.16		76		
57515632	**Good identification**						**57515632**
Bacteroides thetaiotaomicron		97.96	0.39	TRE 93			
Clostridium clostridioforme		1.45	0.18	LAC 98			
Bacteroides ovatus		0.32	-.03				
57515702	**Acceptable identification**						**57515702**
					GRA	CAT	
Bacteroides thetaiotaomicron		92.96	0.41	TRE 93	1	89	
Bacteroides uniformis		3.35	0.14	NAG 99	1	5	
Bifidobacterium longum subsp. longum		0.79	0.17		99	1	
57515720	**Very good identification**						**57515720**
					CAT		
Bacteroides vulgatus		97.28	0.66	IND 1	3		
Bacteroides thetaiotaomicron		2.51	0.41		89		
Bacteroides uniformis		0.09	0.14		5		
57515722	**Very good identification**						**57515722**
					CAT		
Bacteroides thetaiotaomicron		95.38	0.75	TRE 93	89		
Bacteroides uniformis		3.43	0.48		5		
Parabacteroides distasonis		0.39	0.50		76		
57515723	**Good identification**						**57515723**
					CAT		
Bacteroides thetaiotaomicron		95.43	0.41	TRE 93	89		
Bacteroides uniformis		3.44	0.14	URE 1	5		
Parabacteroides distasonis		0.39	0.16		76		
57515732	**Very good identification**						**57515732**
Bacteroides thetaiotaomicron		98.45	0.68	TRE 93			
Bacteroides ovatus		0.64	0.31				
Parabacteroides distasonis		0.56	0.46				
57515762	**Acceptable identification**						**57515762**
					CAT		
Bacteroides thetaiotaomicron		94.36	0.41	TRE 93	89		
Bacteroides uniformis		3.40	0.14	SOR 1	5		
Bacteroides ovatus		1.48	0.10		68		
57517000	**Good identification**						**57517000**
Eubacterium saburreum		100.0	0.40	MAN 1			
57517010	**Good identification**						**57517010**
Eubacterium saburreum		100.0	0.39	MAN 1			
57517100	**Good identification**						**57517100**
Eubacterium saburreum		99.08	0.46	MAN 1			
Actinomyces israelii		0.92	0.16				
57517110	**Good identification**						**57517110**
Eubacterium saburreum		100.0	0.45	MAN 1			
57517502	**Good identification**						**57517502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		95.77	0.44	IND 1		
Bifidobacterium longum subsp. longum		3.18	0.17			
Actinomyces israelii		1.05	0.17			
57517512	**Good identification**					**57517512**
Bifidobacterium adolescentis		95.58	0.46	IND 1		
Bifidobacterium longum subsp. longum		4.42	0.21			
57517722	**Low discrimination**					**57517722**
					CAT	
Bacteroides thetaiotaomicron		85.48	0.41	TRE 93	89	
Bacteroides ovatus		10.75	0.26	MAN 1	68	
Bacteroides uniformis		3.08	0.14		5	
57520100	**Good identification**					**57520100**
Eubacterium saburreum		100.0	0.40	GAL 99		
57520110	**Good identification**					**57520110**
Eubacterium saburreum		100.0	0.38	GAL 99		
57521000	**Very good identification**					**57521000**
Eubacterium saburreum		100.0	0.68			
57521002	**Good identification**					**57521002**
Eubacterium saburreum		100.0	0.55			
57521010	**Very good identification**					**57521010**
Eubacterium saburreum		100.0	0.66			
57521012	**Good identification**					**57521012**
Eubacterium saburreum		100.0	0.53			
57521020	**Good identification**					**57521020**
Eubacterium saburreum		100.0	0.55			
57521022	**Good identification**					**57521022**
Eubacterium saburreum		100.0	0.42			
57521030	**Good identification**					**57521030**
Eubacterium saburreum		100.0	0.53			
57521032	**Good identification**					**57521032**
Eubacterium saburreum		100.0	0.40			
57521100	**Very good identification**					**57521100**
Eubacterium saburreum		100.0	0.74			
57521101	**Good identification**					**57521101**
					SPO	
Eubacterium saburreum		96.16	0.40	URE 1	1	
Clostridium chauvoei		3.85	0.06		99	
57521102	**Very good identification**					**57521102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.61			
57521110	**Very good identification**					**57521110**
Eubacterium saburreum		100.0	0.72			
57521111	**Good identification**					**57521111**
Eubacterium saburreum		100.0	0.38	URE 1		
57521112	**Very good identification**					**57521112**
Eubacterium saburreum		100.0	0.60			
57521120	**Very good identification**					**57521120**
					GRA	
Eubacterium saburreum		98.49	0.61		85	
Prevotella melaninogenica		1.20	0.12		1	
Clostridium perfringens		0.31	0.09		99	
57521122	**Good identification**					**57521122**
					GRA	
Eubacterium saburreum		97.34	0.48		85	
Prevotella melaninogenica		1.28	0.00		1	
Clostridium perfringens		0.81	0.03		99	
57521130	**Very good identification**					**57521130**
Eubacterium saburreum		100.0	0.60			
57521132	**Good identification**					**57521132**
Eubacterium saburreum		100.0	0.47			
57521140	**Good identification**					**57521140**
Eubacterium saburreum		100.0	0.40	SOR 1		
57521150	**Good identification**					**57521150**
Eubacterium saburreum		100.0	0.38	SOR 1		
57521300	**Good identification**					**57521300**
Eubacterium saburreum		100.0	0.40	RHA 1		
57521310	**Good identification**					**57521310**
Eubacterium saburreum		100.0	0.38	RHA 1		
57521500	**Acceptable identification**					**57521500**
Eubacterium saburreum		92.91	0.40	ARA 1		
Bifidobacterium longum subsp. longum		7.09	0.17			
57521510	**Low discrimination**					**57521510**
Eubacterium saburreum		86.29	0.38	ARA 1		
Bifidobacterium longum subsp. longum		13.71	0.21			
57521722	**Good identification**					**57521722**
Bacteroides thetaiotaomicron		99.63	0.37	FRU 99		
Parabacteroides distasonis		0.37	0.11	XYL 96		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
57523100	**Good identification**					**57523100**
	Eubacterium saburreum	100.0	0.40	MAN 1		
57523110	**Good identification**					**57523110**
	Eubacterium saburreum	100.0	0.38	MAN 1		
57524100	**Good identification**					**57524100**
	Eubacterium saburreum	100.0	0.35	GAL 99		
57525000	**Very good identification**					**57525000**
	Eubacterium saburreum	100.0	0.63			
57525002	**Good identification**					**57525002**
	Eubacterium saburreum	100.0	0.50			
57525010	**Very good identification**					**57525010**
	Eubacterium saburreum	100.0	0.62			
57525012	**Good identification**					**57525012**
	Eubacterium saburreum	100.0	0.49			
57525020	**Good identification**					**57525020**
	Eubacterium saburreum	100.0	0.50			
57525022	**Good identification**					**57525022**
	Eubacterium saburreum	100.0	0.37			
57525030	**Good identification**					**57525030**
	Eubacterium saburreum	100.0	0.49			
57525032	**Good identification**					**57525032**
	Eubacterium saburreum	100.0	0.36			
57525100	**Very good identification**					**57525100**
	Eubacterium saburreum	100.0	0.69			
57525101	**Good identification**					**57525101**
	Eubacterium saburreum	100.0	0.35	URE 1		
57525102	**Very good identification**					**57525102**
	Eubacterium saburreum	100.0	0.56			
57525110	**Very good identification**					**57525110**
	Eubacterium saburreum	100.0	0.68			
57525112	**Good identification**					**57525112**
	Eubacterium saburreum	100.0	0.55			
57525120	**Very good identification**					**57525120**
	Eubacterium saburreum	100.0	0.56			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57525122	**Low discrimination**						**57525122**
					CAT		
Eubacterium saburreum		86.46	0.44		1		
Parabacteroides distasonis		5.21	0.23		76		
Bacteroides thetaiotaomicron		5.18	0.07		89		
57525130	**Good identification**						**57525130**
Eubacterium saburreum		100.0	0.55				
57525132	**Acceptable identification**						**57525132**
					CAT		
Eubacterium saburreum		93.62	0.42		1		
Parabacteroides distasonis		3.71	0.18		76		
Bacteroides thetaiotaomicron		2.66	0.00		89		
57525140	**Good identification**						**57525140**
Eubacterium saburreum		100.0	0.35	SOR 1			
57525300	**Good identification**						**57525300**
Eubacterium saburreum		100.0	0.35	RHA 1			
57525500	**Low discrimination**						**57525500**
Eubacterium saburreum		70.16	0.35	ARA 1			
Bifidobacterium longum subsp. longum		29.84	0.25				
57525520	**Low discrimination**						**57525520**
					CAT	GRA	
Bifidobacterium longum subsp. longum		81.19	0.35	IND 1	1	99	
Bacteroides thetaiotaomicron		8.81	0.07	FRU 90	89	1	
Eubacterium saburreum		8.42	0.22		1	85	
57525522	**Acceptable identification**						**57525522**
					GRA	CAT	
Bacteroides thetaiotaomicron		90.66	0.41	FRU 99	1	89	
Bifidobacterium longum subsp. longum		8.44	0.35	RHA 93	99	1	
Parabacteroides distasonis		0.57	0.19		1	76	
57525530	**Acceptable identification**						**57525530**
					CAT	GRA	
Bifidobacterium longum subsp. longum		93.06	0.39	IND 1	1	99	
Eubacterium saburreum		4.64	0.21	FRU 90	1	85	
Bacteroides thetaiotaomicron		2.31	0.00		89	1	
57525722	**Very good identification**						**57525722**
Bacteroides thetaiotaomicron		99.57	0.60	FRU 99			
Bacteroides ovatus		0.29	0.17				
Parabacteroides distasonis		0.14	0.27				
57525732	**Good identification**						**57525732**
Bacteroides thetaiotaomicron		99.26	0.53	FRU 99			
Bacteroides ovatus		0.49	0.14				
Parabacteroides distasonis		0.19	0.23				
57527100	**Good identification**						**57527100**
Eubacterium saburreum		100.0	0.35	MAN 1			
57530000	**Low discrimination**						**57530000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT	GRA	
Eubacterium saburreum		58.07	0.42	GAL 99	1	85	
Prevotella intermedia		26.88	0.26		1	1	
Propionibacterium granulosum		8.50	0.26		90	99	
57530010	**Good identification**						**57530010**
					CAT		
Eubacterium saburreum		97.62	0.40	GAL 99	1		
Propionibacterium granulosum		2.38	0.11		90		
57530100	**Low discrimination**						**57530100**
Eubacterium saburreum		89.86	0.48	GAL 99			
Fusobacterium mortiferum		10.14	0.33				
57530102	**Acceptable identification**						**57530102**
Eubacterium saburreum		93.37	0.35	GAL 99			
Fusobacterium mortiferum		6.63	0.17				
57530110	**Good identification**						**57530110**
Eubacterium saburreum		100.0	0.46	GAL 99			
57530120	**Low discrimination**						**57530120**
					GRA		
Eubacterium saburreum		87.78	0.35	GAL 99	85		
Clostridium perfringens		9.00	0.09		99		
Prevotella melaninogenica		3.22	-.06		1		
57530722	**Good identification**						**57530722**
Bacteroides thetaiotaomicron		99.26	0.37	GAL 99			
Parabacteroides distasonis		0.37	0.11	XYL 96			
Clostridium clostridioforme		0.37	0.05				
57531000	**Very good identification**						**57531000**
Eubacterium saburreum		99.19	0.76				
Prevotella intermedia		0.46	0.26				
Fusobacterium mortiferum		0.17	0.30				
57531001	**Good identification**						**57531001**
Eubacterium saburreum		97.76	0.42	URE 1			
Actinomyces naeslundii		2.24	0.27				
57531002	**Very good identification**						**57531002**
					CAT	GRA	
Eubacterium saburreum		99.60	0.63		1	85	
Propionibacterium granulosum		0.21	0.15		90	99	
Fusobacterium mortiferum		0.11	0.14		5	1	
57531010	**Very good identification**						**57531010**
					CAT		
Eubacterium saburreum		99.98	0.74		1		
Propionibacterium granulosum		0.03	0.11		90		
57531011	**Good identification**						**57531011**
Eubacterium saburreum		100.0	0.40	URE 1			
57531012	**Very good identification**						**57531012**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.61				
57531020	**Very good identification**						**57531020**
					CAT		
Eubacterium saburreum		96.48	0.63		1		
Clostridium perfringens		3.15	0.28		1		
Propionibacterium granulosum		0.20	0.15		90		
57531022	**Low discrimination**						**57531022**
					GRA		
Eubacterium saburreum		89.21	0.50		85		
Clostridium perfringens		7.76	0.23		99		
Prevotella oralis		2.59	0.11		1		
57531030	**Very good identification**						**57531030**
Eubacterium saburreum		100.0	0.61				
57531032	**Good identification**						**57531032**
Eubacterium saburreum		100.0	0.49				
57531040	**Good identification**						**57531040**
Eubacterium saburreum		100.0	0.42	SOR 1			
57531050	**Good identification**						**57531050**
Eubacterium saburreum		100.0	0.40	SOR 1			
57531100	**Very good identification**						**57531100**
					COC	GRA	
Eubacterium saburreum		99.68	0.82		1	85	
Fusobacterium mortiferum		0.17	0.36		1	1	
Atopobium parvulum		0.07	0.19		99	99	
57531101	**Acceptable identification**						**57531101**
					SPO		
Eubacterium saburreum		94.42	0.48	URE 1	1		
Clostridium chauvoei		2.94	0.12		99		
Actinomyces naeslundii		2.64	0.35		1		
57531102	**Very good identification**						**57531102**
					CAT		
Eubacterium saburreum		99.03	0.69		1		
Capnocytophaga ochracea		0.24	0.12		90		
Prevotella oralis		0.18	0.10		1		
57531103	**Low discrimination**						**57531103**
					SPO		
Eubacterium saburreum		84.06	0.35	URE 1	1		
Actinomyces naeslundii		13.33	0.35		1		
Clostridium chauvoei		2.62	-.01		99		
57531110	**Very good identification**						**57531110**
Eubacterium saburreum		100.0	0.81				
57531111	**Good identification**						**57531111**
Eubacterium saburreum		100.0	0.46	URE 1			
57531112	**Very good identification**						**57531112**
					GRA		
Eubacterium saburreum		99.83	0.68		85		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		0.13	0.16		99	
Leptotrichia buccalis		0.04	-.11		1	
57531120	**Low discrimination**					**57531120**
					GRA	
Eubacterium saburreum		87.28	0.69		85	
Clostridium perfringens		8.95	0.43		99	
Prevotella melaninogenica		3.20	0.29		1	
57531121	**Low discrimination**					**57531121**
Eubacterium saburreum		73.72	0.35	URE 1		
Actinomyces naeslundii		11.69	0.35			
Clostridium perfringens		7.56	0.09			
57531122	**Low discrimination**					**57531122**
					GRA	
Eubacterium saburreum		55.11	0.56		85	
Clostridium perfringens		15.07	0.38		99	
Prevotella oralis		10.74	0.32		1	
57531130	**Very good identification**					**57531130**
Eubacterium saburreum		99.72	0.68			
Clostridium perfringens		0.13	0.09			
Bifidobacterium longum subsp. longum		0.11	0.13			
57531132	**Low discrimination**					**57531132**
					CAT	
Eubacterium saburreum		88.63	0.55		1	
Parabacteroides distasonis		6.94	0.36		76	
Bacteroides thetaiotaomicron		1.87	0.10		89	
57531140	**Good identification**					**57531140**
Eubacterium saburreum		98.96	0.48	SOR 1		
Clostridium perfringens		1.04	0.05			
57531142	**Good identification**					**57531142**
Eubacterium saburreum		100.0	0.35	SOR 1		
57531150	**Good identification**					**57531150**
Eubacterium saburreum		100.0	0.46	SOR 1		
57531200	**Good identification**					**57531200**
Eubacterium saburreum		100.0	0.42	RHA 1		
57531210	**Good identification**					**57531210**
Eubacterium saburreum		100.0	0.40	RHA 1		
57531300	**Good identification**					**57531300**
Eubacterium saburreum		100.0	0.48	RHA 1		
57531302	**Low discrimination**					**57531302**
					CAT	
Eubacterium saburreum		80.41	0.35	RHA 1	1	
Bacteroides thetaiotaomicron		8.35	0.02		89	
Prevotella oralis		6.18	0.04		1	
57531310	**Good identification**					**57531310**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.46	RHA 1			
57531320	**Low discrimination**						**57531320**
					CAT	GRA	
Eubacterium saburreum		69.98	0.35	RHA 1	1	85	
Bacteroides thetaiotaomicron		7.27	0.02		89	1	
Clostridium perfringens		7.18	0.09		1	99	
57531322	**Low discrimination**						**57531322**
					CAT		
Bacteroides thetaiotaomicron		56.49	0.37	XYL 96	89		
Parabacteroides distasonis		34.26	0.49	ARA 99	76		
Prevotella oralis		8.02	0.25		1		
57531400	**Good identification**						**57531400**
Eubacterium saburreum		100.0	0.42	ARA 1			
57531410	**Good identification**						**57531410**
Eubacterium saburreum		100.0	0.40	ARA 1			
57531500	**Low discrimination**						**57531500**
Eubacterium saburreum		81.37	0.48	ARA 1			
Bifidobacterium longum subsp. longum		18.63	0.33				
57531510	**Low discrimination**						**57531510**
Eubacterium saburreum		67.73	0.46	ARA 1			
Bifidobacterium longum subsp. longum		32.27	0.37				
57531512	**Low discrimination**						**57531512**
Bifidobacterium longum subsp. longum		53.81	0.37	IND 1			
Eubacterium saburreum		19.93	0.34				
Bifidobacterium adolescentis		17.18	0.30				
57531520	**Low discrimination**						**57531520**
					CAT	GRA	
Bifidobacterium longum subsp. longum		71.94	0.43	IND 1	1	99	
Eubacterium saburreum		13.86	0.35		1	85	
Bacteroides thetaiotaomicron		10.73	0.17		89	1	
57531522	**Acceptable identification**						**57531522**
					GRA	CAT	
Bacteroides thetaiotaomicron		91.47	0.52	XYL 96	1	89	
Bifidobacterium longum subsp. longum		6.19	0.43	RHA 93	99	1	
Parabacteroides distasonis		1.52	0.37		1	76	
57531530	**Low discrimination**						**57531530**
					CAT	GRA	
Bifidobacterium longum subsp. longum		88.76	0.47	IND 1	1	99	
Eubacterium saburreum		8.22	0.34		1	85	
Bacteroides thetaiotaomicron		3.02	0.10		89	1	
57531532	**Low discrimination**						**57531532**
					GRA	CAT	
Bacteroides thetaiotaomicron		75.29	0.45	XYL 96	1	89	
Bifidobacterium longum subsp. longum		22.32	0.47	RHA 93	99	1	
Parabacteroides distasonis		1.73	0.33		1	76	
57531622	**Good identification**						**57531622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		99.65	0.42	XYL 96			
Parabacteroides distasonis		0.19	0.11	LAC 98			
Clostridium clostridioforme		0.17	0.05				
57531702	**Good identification**						**57531702**
Bacteroides thetaiotaomicron		98.53	0.37	XYL 96			
Clostridium clostridioforme		1.10	0.14	NAG 99			
Parabacteroides distasonis		0.37	0.11				
57531720	**Acceptable identification**						**57531720**
					GRA	CAT	
Bacteroides thetaiotaomicron		93.60	0.37	XYL 96	1	89	
Bacteroides vulgatus		5.58	0.13	BGL 99	1	3	
Bifidobacterium longum subsp. longum		0.48	0.09		99	1	
57531722	**Very good identification**						**57531722**
Bacteroides thetaiotaomicron		99.54	0.71	XYL 96			
Parabacteroides distasonis		0.37	0.45				
Bacteroides ovatus		0.07	0.17				
57531723	**Good identification**						**57531723**
Bacteroides thetaiotaomicron		99.63	0.37	XYL 96			
Parabacteroides distasonis		0.37	0.11	URE 1			
57531732	**Very good identification**						**57531732**
Bacteroides thetaiotaomicron		99.21	0.64	XYL 96			
Parabacteroides distasonis		0.52	0.41				
Clostridium clostridioforme		0.13	0.25				
57531762	**Good identification**						**57531762**
Bacteroides thetaiotaomicron		99.63	0.37	XYL 96			
Parabacteroides distasonis		0.37	0.11	SOR 1			
57533000	**Low discrimination**						**57533000**
					CAT		
Eubacterium saburreum		87.72	0.42	MAN 1	1		
Propionibacterium granulosum		9.68	0.24		90		
Actinomyces naeslundii		2.60	0.29		25		
57533010	**Good identification**						**57533010**
					CAT		
Eubacterium saburreum		98.19	0.40	MAN 1	1		
Propionibacterium granulosum		1.81	0.09		90		
57533100	**Good identification**						**57533100**
Eubacterium saburreum		96.51	0.48	MAN 1			
Actinomyces naeslundii		3.50	0.37				
57533101	**Low discrimination**						**57533101**
Actinomyces naeslundii		84.32	0.40	IND 1			
Eubacterium saburreum		15.68	0.14	ESL 12			
57533102	**Low discrimination**						**57533102**
Eubacterium saburreum		78.38	0.35	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		16.09	0.37				
Bifidobacterium adolescentis		5.53	0.13				
57533103	**Good identification**						**57533103**
Actinomyces naeslundii		100.0	0.40	IND 1			
57533110	**Good identification**						**57533110**
Eubacterium saburreum		100.0	0.46	MAN 1			
57533120	**Low discrimination**						**57533120**
Eubacterium saburreum		74.38	0.35	MAN 1			
Actinomyces naeslundii		15.27	0.37				
Clostridium perfringens		7.63	0.09				
57533121	**Good identification**						**57533121**
Actinomyces naeslundii		100.0	0.40	IND 1			
57533122	**Low discrimination**						**57533122**
Actinomyces naeslundii		44.20	0.37	IND 1			
Eubacterium saburreum		38.00	0.22	ESL 12			
Clostridium perfringens		10.39	0.03				
57533123	**Good identification**						**57533123**
Actinomyces naeslundii		100.0	0.40	IND 1			
57533722	**Good identification**						**57533722**
Bacteroides thetaiotaomicron		97.38	0.37	MAN 1			
Bacteroides ovatus		2.26	0.09	XYL 96			
Parabacteroides distasonis		0.37	0.11				
57534000	**Good identification**						**57534000**
Eubacterium saburreum		100.0	0.37	GAL 99			
57534010	**Good identification**						**57534010**
Eubacterium saburreum		100.0	0.36	GAL 99			
57534100	**Good identification**						**57534100**
Eubacterium saburreum		98.45	0.43	GAL 99			
Fusobacterium mortiferum		1.55	0.14				
57534110	**Good identification**						**57534110**
Eubacterium saburreum		100.0	0.42	GAL 99			
57534522	**Good identification**						**57534522**
					GRA	CAT	
Bacteroides thetaiotaomicron		98.08	0.41	GAL 99	1	89	
Bifidobacterium longum subsp. longum		0.83	0.17	RHA 93	99	1	
Parabacteroides distasonis		0.61	0.19		1	76	
57534722	**Very good identification**						**57534722**
Bacteroides thetaiotaomicron		99.42	0.60	GAL 99			
Bacteroides ovatus		0.29	0.17				
Clostridium clostridioforme		0.16	0.23				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57534732	**Good identification**						**57534732**
Bacteroides thetaiotaomicron		97.53	0.53	GAL 99			
Clostridium clostridioforme		1.80	0.34				
Bacteroides ovatus		0.49	0.14				
57535000	**Very good identification**						**57535000**
Eubacterium saburreum		99.98	0.71				
Fusobacterium mortiferum		0.02	0.11				
57535001	**Good identification**						**57535001**
Eubacterium saburreum		100.0	0.37	URE 1			
57535002	**Very good identification**						**57535002**
Eubacterium saburreum		100.0	0.58				
57535010	**Very good identification**						**57535010**
Eubacterium saburreum		100.0	0.70				
57535011	**Good identification**						**57535011**
Eubacterium saburreum		100.0	0.36	URE 1			
57535012	**Very good identification**						**57535012**
Eubacterium saburreum		100.0	0.57				
57535020	**Very good identification**						**57535020**
Eubacterium saburreum		100.0	0.58				
57535022	**Low discrimination**						**57535022**
					CAT		
Eubacterium saburreum		85.52	0.45		1		
Bacteroides thetaiotaomicron		8.04	0.12		89		
Parabacteroides distasonis		4.01	0.23		76		
57535030	**Very good identification**						**57535030**
Eubacterium saburreum		100.0	0.57				
57535032	**Acceptable identification**						**57535032**
					CAT		
Eubacterium saburreum		92.98	0.44		1		
Bacteroides thetaiotaomicron		4.16	0.05		89		
Parabacteroides distasonis		2.87	0.18		76		
57535040	**Good identification**						**57535040**
Eubacterium saburreum		100.0	0.37	SOR 1			
57535050	**Good identification**						**57535050**
Eubacterium saburreum		100.0	0.36	SOR 1			
57535100	**Very good identification**						**57535100**
					COC	GRA	
Eubacterium saburreum		99.71	0.78		1	85	
Atopobium parvulum		0.25	0.23		99	99	
Fusobacterium mortiferum		0.02	0.17		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
57535101	**Good identification**					**57535101**
Eubacterium saburreum		100.0	0.43	URE 1		
57535102	**Very good identification**					**57535102**
					GRA	
Eubacterium saburreum		97.54	0.65		85	
Bifidobacterium adolescentis		1.06	0.29		99	
Parabacteroides distasonis		0.35	0.23		1	
57535110	**Very good identification**					**57535110**
Eubacterium saburreum		99.97	0.76			
Bifidobacterium longum subsp. longum		0.03	0.11			
57535111	**Good identification**					**57535111**
Eubacterium saburreum		100.0	0.42	URE 1		
57535112	**Very good identification**					**57535112**
					GRA	
Eubacterium saburreum		97.77	0.63		85	
Bifidobacterium adolescentis		1.58	0.30		99	
Parabacteroides distasonis		0.23	0.18		1	
57535120	**Very good identification**					**57535120**
					CAT	
Eubacterium saburreum		96.88	0.65		1	
Bacteroides fragilis		1.08	0.10		99	
Bacteroides vulgatus		0.46	0.07		3	
57535122	**Low discrimination**					**57535122**
Parabacteroides distasonis		31.50	0.57	IND 1		
Bacteroides thetaiotaomicron		31.28	0.41			
Bacteroides fragilis		19.07	0.32			
57535130	**Very good identification**					**57535130**
					GRA	
Eubacterium saburreum		99.00	0.63		85	
Bifidobacterium longum subsp. longum		0.61	0.21		99	
Parabacteroides distasonis		0.23	0.18		1	
57535132	**Low discrimination**					**57535132**
					CAT	
Parabacteroides distasonis		39.41	0.53	IND 1	76	
Eubacterium saburreum		30.11	0.50		1	
Bacteroides thetaiotaomicron		28.27	0.34		89	
57535140	**Good identification**					**57535140**
Eubacterium saburreum		100.0	0.43	SOR 1		
57535150	**Good identification**					**57535150**
Eubacterium saburreum		100.0	0.42	SOR 1		
57535200	**Good identification**					**57535200**
Eubacterium saburreum		100.0	0.37	RHA 1		
57535210	**Good identification**					**57535210**
Eubacterium saburreum		100.0	0.36	RHA 1		

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
57535300	**Good identification**						**57535300**
Eubacterium saburreum		100.0	0.43	RHA 1			
57535310	**Good identification**						**57535310**
Eubacterium saburreum		100.0	0.42	RHA 1			
57535322	**Low discrimination**						**57535322**
Bacteroides thetaiotaomicron		80.75	0.60	ARA 99			
Parabacteroides distasonis		18.36	0.65				
Bacteroides ovatus		0.72	0.25				
57535332	**Low discrimination**						**57535332**
Bacteroides thetaiotaomicron		74.94	0.53	ARA 99			
Parabacteroides distasonis		23.60	0.61				
Bacteroides ovatus		1.14	0.22				
57535400	**Good identification**						**57535400**
Eubacterium saburreum		97.08	0.37	ARA 1			
Bifidobacterium longum subsp. longum		2.92	0.07				
57535410	**Acceptable identification**						**57535410**
Eubacterium saburreum		94.11	0.36	ARA 1			
Bifidobacterium longum subsp. longum		5.89	0.11				
57535422	**Good identification**						**57535422**
					GRA	CAT	
Bacteroides thetaiotaomicron		98.95	0.46	LAC 98	1	89	
Bifidobacterium longum subsp. longum		0.42	0.17	RHA 93	99	1	
Parabacteroides distasonis		0.31	0.19		1	76	
57535432	**Good identification**						**57535432**
					GRA	CAT	
Bacteroides thetaiotaomicron		96.18	0.39	LAC 98	1	89	
Bifidobacterium longum subsp. longum		1.76	0.21	RHA 93	99	1	
Clostridium clostridioforme		1.42	0.18		50	33	
57535500	**Low discrimination**						**57535500**
					SPO		
Bifidobacterium longum subsp. longum		52.62	0.41	IND 1	1		
Eubacterium saburreum		41.24	0.43		1		
Clostridium butyricum		2.62	0.06		99		
57535502	**Low discrimination**						**57535502**
					GRA	CAT	
Bacteroides thetaiotaomicron		68.07	0.41	RHA 93	1	89	
Bifidobacterium longum subsp. longum		14.26	0.41	NAG 99	99	1	
Bifidobacterium adolescentis		14.15	0.43		99	1	
57535510	**Low discrimination**						**57535510**
					CAT	GRA	
Bifidobacterium longum subsp. longum		71.69	0.45	IND 1	1	99	
Eubacterium saburreum		27.00	0.42		1	85	
Bacteroides thetaiotaomicron		0.79	0.00		89	1	
57535520	**Low discrimination**						**57535520**
					GRA	CAT	
Bacteroides thetaiotaomicron		48.67	0.41	RHA 93	1	89	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		40.78	0.52	BGL 99	99	1	
Bacteroides vulgatus		8.74	0.26		1	3	
57535522	**Very good identification**						**57535522**
					GRA	CAT	
Bacteroides thetaiotaomicron		98.30	0.75	RHA 93	1	89	
Bifidobacterium longum subsp. longum		0.83	0.52		99	1	
Parabacteroides distasonis		0.61	0.53		1	76	
57535523	**Good identification**						**57535523**
					GRA	CAT	
Bacteroides thetaiotaomicron		98.36	0.41	RHA 93	1	89	
Bifidobacterium longum subsp. longum		0.83	0.17	URE 1	99	1	
Parabacteroides distasonis		0.61	0.19		1	76	
57535530	**Low discrimination**						**57535530**
					CAT	GRA	
Bifidobacterium longum subsp. longum		77.35	0.56	IND 1	1	99	
Bacteroides thetaiotaomicron		21.08	0.34		89	1	
Eubacterium saburreum		1.29	0.29		1	85	
57535532	**Very good identification**						**57535532**
					GRA	CAT	
Bacteroides thetaiotaomicron		95.21	0.68	RHA 93	1	89	
Bifidobacterium longum subsp. longum		3.53	0.56		99	1	
Parabacteroides distasonis		0.82	0.49		1	76	
57535562	**Good identification**						**57535562**
					GRA	CAT	
Bacteroides thetaiotaomicron		97.75	0.41	RHA 93	1	89	
Bifidobacterium longum subsp. longum		0.83	0.17	SOR 1	99	1	
Bacteroides ovatus		0.82	0.06		1	68	
57535622	**Very good identification**						**57535622**
Bacteroides thetaiotaomicron		99.72	0.66	LAC 98			
Bacteroides ovatus		0.14	0.17				
Parabacteroides distasonis		0.07	0.27				
57535632	**Very good identification**						**57535632**
Bacteroides thetaiotaomicron		98.85	0.58	LAC 98			
Clostridium clostridioforme		0.81	0.33				
Bacteroides ovatus		0.24	0.14				
57535702	**Very good identification**						**57535702**
Bacteroides thetaiotaomicron		99.08	0.60	NAG 99			
Clostridium clostridioforme		0.47	0.31				
Bacteroides ovatus		0.29	0.17				
57535712	**Acceptable identification**						**57535712**
Bacteroides thetaiotaomicron		94.04	0.53	NAG 99			
Clostridium clostridioforme		5.19	0.42				
Bacteroides ovatus		0.47	0.14				
57535720	**Low discrimination**						**57535720**
					CAT		
Bacteroides thetaiotaomicron		88.77	0.60	BGL 99	89		
Bacteroides vulgatus		10.80	0.42		3		
Bacteroides ovatus		0.26	0.17		68		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57535722	**Excellent identification**						**57535722**
Bacteroides thetaiotaomicron		99.56	0.94				
Bacteroides ovatus		0.29	0.51				
Parabacteroides distasonis		0.14	0.62				
57535723	**Very good identification**						**57535723**
Bacteroides thetaiotaomicron		99.57	0.60	URE 1			
Bacteroides ovatus		0.29	0.17				
Parabacteroides distasonis		0.14	0.27				
57535730	**Good identification**						**57535730**
					CAT		
Bacteroides thetaiotaomicron		98.68	0.53	BGL 99	89		
Bacteroides ovatus		0.49	0.14		68		
Bacteroides vulgatus		0.31	0.08		3		
57535732	**Very good identification**						**57535732**
Bacteroides thetaiotaomicron		99.26	0.87				
Bacteroides ovatus		0.49	0.48				
Parabacteroides distasonis		0.19	0.57				
57535733	**Good identification**						**57535733**
Bacteroides thetaiotaomicron		99.26	0.53	URE 1			
Bacteroides ovatus		0.49	0.14				
Parabacteroides distasonis		0.19	0.23				
57535762	**Very good identification**						**57535762**
Bacteroides thetaiotaomicron		98.68	0.60	SOR 1			
Bacteroides ovatus		1.18	0.27				
Parabacteroides distasonis		0.14	0.27				
57535772	**Good identification**						**57535772**
Bacteroides thetaiotaomicron		97.75	0.53	SOR 1			
Bacteroides ovatus		2.00	0.24				
Parabacteroides distasonis		0.19	0.23				
57537000	**Good identification**						**57537000**
Eubacterium saburreum		97.04	0.37	MAN 1			
Actinomyces israelii		2.96	0.15				
57537010	**Good identification**						**57537010**
Eubacterium saburreum		100.0	0.36	MAN 1			
57537100	**Good identification**						**57537100**
Eubacterium saburreum		95.03	0.43	MAN 1			
Actinomyces israelii		4.98	0.25				
57537110	**Good identification**						**57537110**
Eubacterium saburreum		100.0	0.42	MAN 1			
57537502	**Low discrimination**						**57537502**
					GRA	CAT	
Bifidobacterium adolescentis		88.19	0.41	IND 1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		5.47	0.27		99	1	
Bacteroides thetaiotaomicron		5.24	0.07		1	89	
57537512	**Good identification**						**57537512**
					GRA	CAT	
Bifidobacterium adolescentis		95.19	0.43	IND 1	99	1	
Bacteroides thetaiotaomicron		1.80	0.00		1	89	
Bifidobacterium longum subsp. longum		1.65	0.11		99	1	
57537522	**Acceptable identification**						**57537522**
					GRA	CAT	
Bacteroides thetaiotaomicron		92.31	0.41	MAN 1	1	89	
Bacteroides ovatus		6.18	0.21	RHA 93	1	68	
Bifidobacterium longum subsp. longum		0.78	0.17		99	1	
57537722	**Good identification**						**57537722**
Bacteroides thetaiotaomicron		91.15	0.60	MAN 1			
Bacteroides ovatus		8.72	0.43				
Parabacteroides distasonis		0.13	0.27				
57537732	**Low discrimination**						**57537732**
Bacteroides thetaiotaomicron		85.72	0.53	MAN 1			
Bacteroides ovatus		14.06	0.40				
Parabacteroides distasonis		0.17	0.23				
57541000	**Good identification**						**57541000**
Eubacterium saburreum		100.0	0.37	CEL 1			
57541010	**Good identification**						**57541010**
Eubacterium saburreum		100.0	0.35	CEL 1			
57541100	**Good identification**						**57541100**
Eubacterium saburreum		98.44	0.43	CEL 1			
Prevotella melaninogenica		1.56	-.04				
57541110	**Good identification**						**57541110**
Eubacterium saburreum		100.0	0.41	CEL 1			
57545100	**Good identification**						**57545100**
Eubacterium saburreum		100.0	0.38	CEL 1			
57545110	**Good identification**						**57545110**
Eubacterium saburreum		100.0	0.37	CEL 1			
57545522	**Acceptable identification**						**57545522**
					CAT		
Bacteroides uniformis		93.06	0.45	SAL 92	5		
Bacteroides thetaiotaomicron		5.77	0.27	FRU 99	89		
Bacteroides ovatus		0.72	0.12		68		
57545722	**Low discrimination**						**57545722**
					CAT		
Bacteroides uniformis		53.19	0.48	SAL 92	5		
Bacteroides thetaiotaomicron		31.71	0.47	FRU 99	89		
Bacteroides eggerthii		9.45	0.32		93		
57545732	**Low discrimination**						**57545732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Bacteroides thetaiotaomicron		73.77	0.40	FRU 99	89		
Bacteroides ovatus		22.45	0.31	TRE 93	68		
Bacteroides uniformis		3.21	0.14		5		
57551000	**Low discrimination**						**57551000**
Eubacterium saburreum		68.82	0.45	CEL 1			
Prevotella intermedia		21.24	0.26				
Fusobacterium mortiferum		8.67	0.31				
57551010	**Good identification**						**57551010**
Eubacterium saburreum		100.0	0.43	CEL 1			
57551022	**Acceptable identification**						**57551022**
					CAT		
Prevotella oralis		93.28	0.41	IND 1	1		
Capnocytophaga ochracea		4.21	0.18	LAC 94	90		
Eubacterium saburreum		0.88	0.19		1		
57551100	**Low discrimination**						**57551100**
					CAT		
Eubacterium saburreum		74.43	0.51	CEL 1	1		
Fusobacterium mortiferum		9.37	0.37		5		
Capnocytophaga ochracea		5.83	0.20		90		
57551102	**Low discrimination**						**57551102**
					CAT		
Prevotella oralis		53.49	0.40	IND 1	1		
Capnocytophaga ochracea		32.20	0.36	NAG 95	90		
Eubacterium saburreum		8.06	0.38		1		
57551110	**Good identification**						**57551110**
					CAT	GRA	
Eubacterium saburreum		98.36	0.49	CEL 1	1	85	
Capnocytophaga ochracea		1.05	0.04		90	1	
Bifidobacterium breve		0.60	0.10		1	99	
57551112	**Low discrimination**						**57551112**
					CAT	GRA	
Eubacterium saburreum		43.95	0.37	CEL 1	1	85	
Capnocytophaga ochracea		23.85	0.20		90	1	
Bifidobacterium breve		15.28	0.27		1	99	
57551120	**Low discrimination**						**57551120**
					CAT		
Prevotella melaninogenica		68.08	0.47	IND 1	1		
Prevotella oralis		16.98	0.40	CEL 10	1		
Capnocytophaga ochracea		10.22	0.36		90		
57551122	**Low discrimination**						**57551122**
					CAT		
Prevotella oralis		72.28	0.62	IND 1	1		
Capnocytophaga ochracea		20.61	0.53		90		
Prevotella melaninogenica		2.91	0.34		1		
57551130	**Low discrimination**						**57551130**
					CAT		
Eubacterium saburreum		49.65	0.37	CEL 1	1		
Capnocytophaga ochracea		26.94	0.20		90		
Prevotella melaninogenica		16.31	0.12		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57551132	**Low discrimination**						**57551132**
					GRA	CAT	
Capnocytophaga ochracea		64.95	0.36	IND 1	1	90	
Prevotella oralis		20.71	0.27	MLZ 10	1	1	
Bifidobacterium breve		8.22	0.32		99	1	
57551162	**Low discrimination**						**57551162**
					GRA	CAT	
Capnocytophaga ochracea		67.87	0.36	IND 1	1	90	
Prevotella oralis		21.64	0.27	SOR 10	1	1	
Bifidobacterium breve		8.59	0.32		99	1	
57551322	**Low discrimination**						**57551322**
					CAT		
Prevotella oralis		89.94	0.55	IND 1	1		
Bacteroides uniformis		6.39	0.30		5		
Bacteroides thetaiotaomicron		1.83	0.23		89		
57551502	**Low discrimination**						**57551502**
					GRA	CAT	
Bifidobacterium adolescentis		50.29	0.37	IND 1	99	1	
Capnocytophaga ochracea		28.91	0.23	XYL 87	1	90	
Bacteroides uniformis		13.24	0.11		1	5	
57551512	**Low discrimination**						**57551512**
					GRA	CAT	
Bifidobacterium adolescentis		88.66	0.39	IND 1	99	1	
Capnocytophaga ochracea		4.63	0.07	XYL 87	1	90	
Bifidobacterium longum subsp. longum		3.52	0.13		99	1	
57551522	**Low discrimination**						**57551522**
					CAT		
Bacteroides uniformis		66.15	0.45	SAL 92	5		
Bacteroides thetaiotaomicron		16.91	0.38	XYL 99	89		
Capnocytophaga ochracea		13.13	0.40		90		
57551722	**Low discrimination**						**57551722**
					CAT		
Bacteroides thetaiotaomicron		68.57	0.57	TRE 93	89		
Bacteroides uniformis		27.88	0.48	XYL 96	5		
Bacteroides ovatus		2.95	0.34		68		
57551732	**Acceptable identification**						**57551732**
					CAT		
Bacteroides thetaiotaomicron		91.58	0.50	TRE 93	89		
Bacteroides ovatus		6.76	0.31	XYL 96	68		
Bacteroides uniformis		0.97	0.14		5		
57553101	**Good identification**						**57553101**
Actinomyces naeslundii		100.0	0.37	IND 1			
57553103	**Good identification**						**57553103**
Actinomyces naeslundii		100.0	0.37	IND 1			
57553121	**Good identification**						**57553121**
Actinomyces naeslundii		100.0	0.37	IND 1			
57553123	**Good identification**						**57553123**
Actinomyces naeslundii		100.0	0.37	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57553502	**Good identification**						**57553502**
Bifidobacterium adolescentis		96.15	0.36	IND 1			
Actinomyces israelii		3.85	0.18	XYL 87			
57553512	**Good identification**						**57553512**
Bifidobacterium adolescentis		100.0	0.37	IND 1			
57554522	**Acceptable identification**						**57554522**
					CAT		
Bacteroides uniformis		93.40	0.45	SAL 92	5		
Bacteroides thetaiotaomicron		5.79	0.27	GAL 99	89		
Bacteroides ovatus		0.72	0.12		68		
57554722	**Low discrimination**						**57554722**
					CAT		
Bacteroides uniformis		58.64	0.48	SAL 92	5		
Bacteroides thetaiotaomicron		34.96	0.47	GAL 99	89		
Bacteroides ovatus		6.21	0.34		68		
57554732	**Low discrimination**						**57554732**
Bacteroides thetaiotaomicron		71.41	0.40	TRE 93			
Bacteroides ovatus		21.73	0.31	GAL 99			
Clostridium clostridioforme		3.75	0.28				
57555000	**Good identification**						**57555000**
Eubacterium saburreum		98.27	0.40	CEL 1			
Fusobacterium mortiferum		1.73	0.12				
57555010	**Good identification**						**57555010**
Eubacterium saburreum		100.0	0.39	CEL 1			
57555100	**Low discrimination**						**57555100**
					CAT		
Eubacterium saburreum		84.41	0.46	CEL 1	1		
Bacteroides fragilis		4.71	0.04		99		
Capnocytophaga ochracea		3.07	0.10		90		
57555110	**Good identification**						**57555110**
Eubacterium saburreum		100.0	0.45	CEL 1			
57555112	**Low discrimination**						**57555112**
					CAT	GRA	
Bifidobacterium adolescentis		71.00	0.39	IND 1	1	99	
Eubacterium saburreum		20.87	0.32	ARA 87	1	85	
Capnocytophaga ochracea		5.26	0.10		90	1	
57555120	**Low discrimination**						**57555120**
					CAT		
Bacteroides fragilis		75.02	0.38	IND 1	99		
Bacteroides uniformis		12.37	0.27	BGL 95	5		
Capnocytophaga ochracea		4.44	0.26		90		
57555122	**Low discrimination**						**57555122**
					CAT		
Bacteroides fragilis		51.30	0.60	IND 1	99		
Bacteroides uniformis		44.06	0.61		5		
Prevotella oralis		2.50	0.46		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57555322	**Low discrimination**						**57555322**
					CAT		
Bacteroides uniformis		87.34	0.64	SAL 92	5		
Bacteroides thetaiotaomicron		6.05	0.47	ARA 92	89		
Bacteroides ovatus		3.29	0.42		68		
57555332	**Low discrimination**						**57555332**
					CAT		
Bacteroides thetaiotaomicron		39.07	0.40	TRE 93	89		
Bacteroides ovatus		36.41	0.39	ARA 99	68		
Bacteroides uniformis		14.65	0.30		5		
57555422	**Acceptable identification**						**57555422**
					CAT		
Bacteroides uniformis		93.78	0.51	SAL 92	5		
Bacteroides thetaiotaomicron		5.81	0.33	LAC 98	89		
Bacteroides ovatus		0.36	0.12		68		
57555502	**Low discrimination**						**57555502**
					GRA	CAT	
Bacteroides uniformis		74.06	0.45	SAL 92	1	5	
Bifidobacterium adolescentis		19.01	0.52	NAG 99	99	1	
Bacteroides thetaiotaomicron		4.59	0.27		1	89	
57555512	**Low discrimination**						**57555512**
					GRA	CAT	
Bifidobacterium adolescentis		85.83	0.53	IND 1	99	1	
Bacteroides thetaiotaomicron		6.59	0.20		1	89	
Bacteroides uniformis		2.76	0.11		1	5	
57555520	**Low discrimination**						**57555520**
					CAT		
Bacteroides uniformis		84.48	0.45	SAL 92	5		
Bacteroides vulgatus		7.42	0.28	BGL 99	3		
Bacteroides thetaiotaomicron		5.24	0.27		89		
57555522	**Good identification**						**57555522**
					CAT		
Bacteroides uniformis		93.31	0.79	SAL 92	5		
Bacteroides thetaiotaomicron		5.78	0.62		89		
Bacteroides ovatus		0.72	0.46		68		
57555523	**Acceptable identification**						**57555523**
					CAT		
Bacteroides uniformis		93.40	0.45	SAL 92	5		
Bacteroides thetaiotaomicron		5.79	0.27	URE 1	89		
Bacteroides ovatus		0.72	0.12		68		
57555532	**Low discrimination**						**57555532**
					CAT		
Bacteroides thetaiotaomicron		60.44	0.55	TRE 93	89		
Bacteroides uniformis		25.33	0.45	RHA 93	5		
Bacteroides ovatus		12.86	0.43		68		
57555542	**Low discrimination**						**57555542**
					GRA		
Bifidobacterium adolescentis		72.63	0.36	IND 1	99		
Bacteroides uniformis		23.12	0.11	SOR 11	1		
Actinomyces israelii		2.27	0.16		99		
57555552	**Good identification**						**57555552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		100.0	0.37	IND 1		
57555562	**Acceptable identification**					**57555562**
					CAT	
Bacteroides uniformis		90.89	0.45	SAL 92	5	
Bacteroides thetaiotaomicron		5.63	0.27	SOR 1	89	
Bacteroides ovatus		2.88	0.23		68	
57555622	**Low discrimination**					**57555622**
					CAT	
Bacteroides uniformis		60.60	0.53	SAL 92	5	
Bacteroides thetaiotaomicron		36.13	0.52	LAC 98	89	
Bacteroides ovatus		3.18	0.34		68	
57555632	**Low discrimination**					**57555632**
					CAT	
Bacteroides thetaiotaomicron		82.14	0.45	TRE 93	89	
Bacteroides ovatus		12.37	0.31	LAC 98	68	
Bacteroides uniformis		3.58	0.19		5	
57555702	**Low discrimination**					**57555702**
					CAT	
Bacteroides uniformis		58.39	0.48	SAL 92	5	
Bacteroides thetaiotaomicron		34.81	0.47	NAG 99	89	
Bacteroides ovatus		6.18	0.34		68	
57555712	**Low discrimination**					**57555712**
Bacteroides thetaiotaomicron		65.99	0.40	TRE 93		
Bacteroides ovatus		20.08	0.31	NAG 99		
Clostridium clostridioforme		10.40	0.36			
57555720	**Low discrimination**					**57555720**
					CAT	
Bacteroides uniformis		43.95	0.48	SAL 92	5	
Bacteroides thetaiotaomicron		26.20	0.47	BGL 99	89	
Bacteroides vulgatus		25.17	0.44		3	
57555722	**Low discrimination**					**57555722**
					CAT	
Bacteroides uniformis		58.73	0.82	SAL 92	5	
Bacteroides thetaiotaomicron		35.01	0.81		89	
Bacteroides ovatus		6.22	0.68		68	
57555723	**Low discrimination**					**57555723**
					CAT	
Bacteroides uniformis		58.73	0.48	SAL 92	5	
Bacteroides thetaiotaomicron		35.02	0.47	URE 1	89	
Bacteroides ovatus		6.22	0.34		68	
57555730	**Low discrimination**					**57555730**
					CAT	
Bacteroides thetaiotaomicron		72.84	0.40	TRE 93	89	
Bacteroides ovatus		22.17	0.31	BGL 99	68	
Bacteroides uniformis		3.17	0.14		5	
57555732	**Low discrimination**					**57555732**
					CAT	
Bacteroides thetaiotaomicron		74.02	0.74	TRE 93	89	
Bacteroides ovatus		22.53	0.65		68	
Bacteroides uniformis		3.23	0.48		5	
57555733	**Low discrimination**					**57555733**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Bacteroides thetaiotaomicron		74.19	0.40	TRE 93	89		
Bacteroides ovatus		22.58	0.31	URE 1	68		
Bacteroides uniformis		3.23	0.14		5		
57555762	**Low discrimination**						**57555762**
					CAT		
Bacteroides uniformis		49.18	0.48	SAL 92	5		
Bacteroides thetaiotaomicron		29.32	0.47	SOR 1	89		
Bacteroides ovatus		21.47	0.44		68		
57555772	**Low discrimination**						**57555772**
					CAT		
Bacteroides ovatus		54.61	0.41	SAL 88	68		
Bacteroides thetaiotaomicron		43.50	0.40	TRE 91	89		
Bacteroides uniformis		1.90	0.14	SOR 4	5		
57557102	**Low discrimination**						**57557102**
					CAT	GRA	
Bifidobacterium adolescentis		61.62	0.36	IND 1	1	99	
Actinomyces israelii		34.65	0.38	ARA 87	1	99	
Bacteroides fragilis		2.01	-.09		99	1	
57557112	**Acceptable identification**						**57557112**
Bifidobacterium adolescentis		93.57	0.37	IND 1			
Actinomyces israelii		6.43	0.24	ARA 87			
57557502	**Low discrimination**						**57557502**
					GRA		
Bifidobacterium adolescentis		88.16	0.50	IND 1	99		
Actinomyces israelii		6.06	0.36		99		
Bacteroides uniformis		4.24	0.11		1		
57557512	**Good identification**						**57557512**
					GRA		
Bifidobacterium adolescentis		98.08	0.52	IND 1	99		
Actinomyces israelii		0.82	0.22		99		
Bacteroides ovatus		0.65	0.01		1		
57557522	**Low discrimination**						**57557522**
					CAT		
Bacteroides uniformis		75.76	0.45	SAL 92	5		
Bacteroides ovatus		19.23	0.38	MAN 1	68		
Bacteroides thetaiotaomicron		4.70	0.27		89		
57557552	**Good identification**						**57557552**
Bifidobacterium adolescentis		97.42	0.36	IND 1			
Actinomyces israelii		2.58	0.15	SOR 11			
57557722	**Low discrimination**						**57557722**
					CAT		
Bacteroides ovatus		68.62	0.60	SAL 88	68		
Bacteroides uniformis		19.65	0.48	TRE 91	5		
Bacteroides thetaiotaomicron		11.72	0.47		89		
57557732	**Good identification**						**57557732**
					CAT		
Bacteroides ovatus		90.59	0.57	SAL 88	68		
Bacteroides thetaiotaomicron		9.02	0.40	TRE 91	89		
Bacteroides uniformis		0.39	0.14		5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57557762	**Acceptable identification**					**57557762**
					CAT	
Bacteroides ovatus		90.03	0.36	SAL 88	68	
Bacteroides uniformis		6.25	0.14	TRE 91	5	
Bacteroides thetaiotaomicron		3.73	0.13	SOR 4	89	
57561100	**Good identification**					**57561100**
Eubacterium saburreum		100.0	0.40	CEL 1		
57561110	**Good identification**					**57561110**
Eubacterium saburreum		100.0	0.38	CEL 1		
57561722	**Good identification**					**57561722**
Bacteroides thetaiotaomicron		96.83	0.42	FRU 99		
Bacteroides ovatus		3.17	0.17	XYL 96		
57561732	**Acceptable identification**					**57561732**
Bacteroides thetaiotaomicron		94.68	0.35	FRU 99		
Bacteroides ovatus		5.32	0.14	XYL 96		
57565100	**Good identification**					**57565100**
Eubacterium saburreum		100.0	0.35	CEL 1		
57565522	**Acceptable identification**					**57565522**
					CAT	
Bacteroides thetaiotaomicron		90.23	0.47	FRU 99	89	
Bacteroides ovatus		8.52	0.29	RHA 93	68	
Bacteroides uniformis		1.11	0.11		5	
57565532	**Low discrimination**					**57565532**
Bacteroides thetaiotaomicron		85.90	0.40	FRU 99		
Bacteroides ovatus		13.91	0.26	RHA 93		
Parabacteroides distasonis		0.19	0.10			
57565622	**Acceptable identification**					**57565622**
Bacteroides thetaiotaomicron		93.73	0.37	FRU 99		
Bacteroides ovatus		6.27	0.17	LAC 98		
57565722	**Low discrimination**					**57565722**
					CAT	
Bacteroides thetaiotaomicron		87.95	0.66	FRU 99	89	
Bacteroides ovatus		11.88	0.51		68	
Bacteroides uniformis		0.11	0.14		5	
57565732	**Low discrimination**					**57565732**
Bacteroides thetaiotaomicron		81.11	0.59	FRU 99		
Bacteroides ovatus		18.78	0.48			
Clostridium clostridioforme		0.07	0.17			
57567722	**Low discrimination**					**57567722**
Bacteroides ovatus		81.68	0.43	SAL 88		
Bacteroides thetaiotaomicron		18.32	0.32	FRU 99		
57567732	**Low discrimination**					**57567732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		88.43	0.40	SAL 88			
Bacteroides thetaiotaomicron		11.57	0.25	FRU 99			
57570722	**Good identification**						**57570722**
Bacteroides thetaiotaomicron		96.28	0.42	GAL 99			
Bacteroides ovatus		3.15	0.17	XYL 96			
Clostridium clostridioforme		0.57	0.14				
57570732	**Low discrimination**						**57570732**
Bacteroides thetaiotaomicron		88.89	0.35	GAL 99			
Clostridium clostridioforme		6.12	0.26	XYL 96			
Bacteroides ovatus		4.99	0.14				
57571000	**Low discrimination**						**57571000**
					GRA		
Eubacterium saburreum		89.38	0.42	CEL 1	85		
Fusobacterium mortiferum		5.04	0.22		1		
Lactobacillus jensenii		3.77	0.10		99		
57571010	**Good identification**						**57571010**
Eubacterium saburreum		98.31	0.40	CEL 1			
Lactobacillus jensenii		1.69	0.02				
57571100	**Low discrimination**						**57571100**
					COC	GRA	
Eubacterium saburreum		78.48	0.48	CEL 1	1	85	
Atopobium parvulum		13.32	0.25		99	99	
Fusobacterium mortiferum		4.43	0.28		1	1	
57571101	**Low discrimination**						**57571101**
Actinomyces naeslundii		78.60	0.37	IND 1			
Eubacterium saburreum		21.40	0.14	ESL 12			
57571102	**Low discrimination**						**57571102**
					GRA		
Eubacterium saburreum		35.36	0.35	CEL 1	85		
Prevotella oralis		22.46	0.19		1		
Bifidobacterium breve		10.06	0.24		99		
57571103	**Good identification**						**57571103**
Actinomyces naeslundii		100.0	0.37	IND 1			
57571110	**Good identification**						**57571110**
Eubacterium saburreum		99.23	0.46	CEL 1			
Leptotrichia buccalis		0.77	-.11				
57571120	**Low discrimination**						**57571120**
					GRA		
Eubacterium saburreum		31.78	0.35	CEL 1	85		
Prevotella oralis		20.19	0.19		1		
Clostridium perfringens		13.44	0.20		99		
57571121	**Good identification**						**57571121**
Actinomyces naeslundii		100.0	0.37	IND 1			
57571122	**Low discrimination**						**57571122**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		69.31	0.41	IND 1	1		
Bacteroides thetaiotaomicron		9.45	0.23	TRE 6	89		
Parabacteroides distasonis		6.43	0.36		76		
57571123	**Low discrimination**						**57571123**
					GRA		
Actinomyces naeslundii		63.29	0.37	IND 1	99		
Prevotella oralis		36.71	0.07	ESL 12	1		
57571322	**Low discrimination**						**57571322**
					CAT		
Bacteroides thetaiotaomicron		65.82	0.42	XYL 96	89		
Prevotella oralis		15.58	0.35	ARA 99	1		
Parabacteroides distasonis		10.11	0.44		76		
57571332	**Low discrimination**						**57571332**
Bacteroides thetaiotaomicron		71.23	0.35	XYL 96			
Parabacteroides distasonis		15.15	0.40	ARA 99			
Bacteroides ovatus		12.24	0.22				
57571512	**Low discrimination**						**57571512**
					GRA	CAT	
Bifidobacterium adolescentis		62.09	0.36	IND 1	99	1	
Bacteroides thetaiotaomicron		26.50	0.16	XYL 87	1	89	
Clostridium clostridioforme		9.92	0.19		50	33	
57571522	**Very good identification**						**57571522**
Bacteroides thetaiotaomicron		96.89	0.57	XYL 96			
Bacteroides ovatus		2.22	0.29	RHA 93			
Parabacteroides distasonis		0.41	0.33				
57571532	**Good identification**						**57571532**
Bacteroides thetaiotaomicron		95.21	0.50	XYL 96			
Bacteroides ovatus		3.74	0.26	RHA 93			
Parabacteroides distasonis		0.56	0.28				
57571622	**Good identification**						**57571622**
Bacteroides thetaiotaomicron		98.15	0.47	XYL 96			
Bacteroides ovatus		1.59	0.17	LAC 98			
Clostridium clostridioforme		0.26	0.14				
57571632	**Acceptable identification**						**57571632**
Bacteroides thetaiotaomicron		94.49	0.40	XYL 96			
Clostridium clostridioforme		2.89	0.25	LAC 98			
Bacteroides ovatus		2.63	0.14				
57571702	**Good identification**						**57571702**
Bacteroides thetaiotaomicron		95.20	0.42	XYL 96			
Bacteroides ovatus		3.12	0.17	NAG 99			
Clostridium clostridioforme		1.68	0.23				
57571712	**Low discrimination**						**57571712**
Bacteroides thetaiotaomicron		79.20	0.35	XYL 96			
Clostridium clostridioforme		16.36	0.34	NAG 99			
Bacteroides ovatus		4.45	0.14				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57571720	**Good identification**					**57571720**
					CAT	
Bacteroides thetaiotaomicron		96.69	0.42	XYL 96	89	
Bacteroides ovatus		3.17	0.17	BGL 99	68	
Bacteroides vulgatus		0.14	-.09		3	
57571722	**Very good identification**					**57571722**
Bacteroides thetaiotaomicron		96.69	0.76	XYL 96		
Bacteroides ovatus		3.17	0.51			
Parabacteroides distasonis		0.09	0.41			
57571723	**Good identification**					**57571723**
Bacteroides thetaiotaomicron		96.83	0.42	XYL 96		
Bacteroides ovatus		3.17	0.17	URE 1		
57571730	**Acceptable identification**					**57571730**
Bacteroides thetaiotaomicron		94.68	0.35	XYL 96		
Bacteroides ovatus		5.32	0.14	BGL 99		
57571732	**Good identification**					**57571732**
Bacteroides thetaiotaomicron		94.38	0.69	XYL 96		
Bacteroides ovatus		5.30	0.48			
Clostridium clostridioforme		0.20	0.34			
57571733	**Acceptable identification**					**57571733**
Bacteroides thetaiotaomicron		94.68	0.35	XYL 96		
Bacteroides ovatus		5.32	0.14	URE 1		
57571762	**Low discrimination**					**57571762**
Bacteroides thetaiotaomicron		88.10	0.42	XYL 96		
Bacteroides ovatus		11.90	0.27	SOR 1		
57571772	**Low discrimination**					**57571772**
Bacteroides thetaiotaomicron		81.20	0.35	XYL 96		
Bacteroides ovatus		18.81	0.24	SOR 1		
57573100	**Low discrimination**					**57573100**
Actinomyces naeslundii		52.94	0.39	IND 1		
Actinomyces israelii		35.92	0.27	ESL 12		
Eubacterium saburreum		11.14	0.14			
57573101	**Good identification**					**57573101**
Actinomyces naeslundii		100.0	0.42	IND 1		
57573103	**Good identification**					**57573103**
Actinomyces naeslundii		100.0	0.42	IND 1		
57573121	**Good identification**					**57573121**
Actinomyces naeslundii		100.0	0.42	IND 1		
57573123	**Good identification**					**57573123**
Actinomyces naeslundii		100.0	0.42	IND 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
57573722	**Low discrimination**						**57573722**
					SPO	CAT	
Bacteroides ovatus		51.90	0.43	SAL 88	1	68	
Bacteroides thetaiotaomicron		48.03	0.42	XYL 99	1	89	
Clostridium ramosum		0.07	-.03		92	1	
57573732	**Low discrimination**						**57573732**
Bacteroides ovatus		64.95	0.40	SAL 88			
Bacteroides thetaiotaomicron		35.05	0.35	XYL 99			
57574522	**Low discrimination**						**57574522**
					CAT		
Bacteroides thetaiotaomicron		89.86	0.47	GAL 99	89		
Bacteroides ovatus		8.49	0.29	RHA 93	68		
Bacteroides uniformis		1.10	0.11		5		
57574532	**Low discrimination**						**57574532**
Bacteroides thetaiotaomicron		82.19	0.40	GAL 99			
Bacteroides ovatus		13.31	0.26	RHA 93			
Clostridium clostridioforme		4.32	0.28				
57574622	**Acceptable identification**						**57574622**
Bacteroides thetaiotaomicron		90.65	0.37	GAL 99			
Bacteroides ovatus		6.06	0.17	LAC 98			
Clostridium clostridioforme		3.29	0.23				
57574712	**Low discrimination**						**57574712**
Clostridium clostridioforme		69.98	0.43	IND 1			
Bacteroides thetaiotaomicron		24.38	0.25	SAL 88			
Bacteroides ovatus		5.65	0.14				
57574722	**Low discrimination**						**57574722**
Bacteroides thetaiotaomicron		87.78	0.66	GAL 99			
Bacteroides ovatus		11.86	0.51				
Clostridium clostridioforme		0.22	0.32				
57574732	**Low discrimination**						**57574732**
Bacteroides thetaiotaomicron		79.30	0.59	GAL 99			
Bacteroides ovatus		18.37	0.48				
Clostridium clostridioforme		2.30	0.43				
57575000	**Low discrimination**						**57575000**
					SPO		
Eubacterium saburreum		87.75	0.37	CEL 1	1		
Actinomyces israelii		6.28	0.22		1		
Clostridium butyricum		5.96	0.00		99		
57575010	**Good identification**						**57575010**
Eubacterium saburreum		100.0	0.36	CEL 1			
57575100	**Low discrimination**						**57575100**
					COC	SPO	
Eubacterium saburreum		41.54	0.43	CEL 1	1	1	
Clostridium butyricum		29.04	0.24		1	99	
Atopobium parvulum		24.32	0.30		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57575110	**Acceptable identification**						**57575110**
					SPO		
Eubacterium saburreum		93.69	0.42	CEL 1	1		
Clostridium butyricum		4.21	0.02		99		
Actinomyces israelii		2.10	0.18		1		
57575112	**Low discrimination**						**57575112**
					CAT	GRA	
Bifidobacterium adolescentis		58.91	0.36	IND 1	1	99	
Eubacterium saburreum		17.31	0.29	ARA 87	1	85	
Bacteroides thetaiotaomicron		6.10	0.05		89	1	
57575122	**Low discrimination**						**57575122**
Bacteroides thetaiotaomicron		54.90	0.47	RHA 93			
Bacteroides ovatus		15.88	0.38	ARA 99			
Parabacteroides distasonis		14.01	0.53				
57575132	**Low discrimination**						**57575132**
Bacteroides thetaiotaomicron		53.47	0.40	RHA 93			
Bacteroides ovatus		26.51	0.34	ARA 99			
Parabacteroides distasonis		18.89	0.48				
57575222	**Low discrimination**						**57575222**
Bacteroides thetaiotaomicron		79.68	0.37	LAC 98			
Bacteroides ovatus		16.31	0.25	ARA 99			
Parabacteroides distasonis		2.27	0.27				
57575322	**Low discrimination**						**57575322**
Bacteroides thetaiotaomicron		67.38	0.66	ARA 99			
Bacteroides ovatus		27.87	0.59				
Parabacteroides distasonis		3.88	0.61				
57575332	**Low discrimination**						**57575332**
Bacteroides thetaiotaomicron		55.72	0.59	ARA 99			
Bacteroides ovatus		39.51	0.56				
Parabacteroides distasonis		4.45	0.56				
57575362	**Low discrimination**						**57575362**
Bacteroides ovatus		61.49	0.36	SAL 88			
Bacteroides thetaiotaomicron		36.04	0.32	ARA 97			
Parabacteroides distasonis		2.08	0.27	SOR 4			
57575422	**Acceptable identification**						**57575422**
					CAT		
Bacteroides thetaiotaomicron		94.18	0.52	LAC 98	89		
Bacteroides ovatus		4.40	0.29	RHA 93	68		
Bacteroides uniformis		1.16	0.16		5		
57575432	**Acceptable identification**						**57575432**
Bacteroides thetaiotaomicron		90.54	0.45	LAC 98			
Bacteroides ovatus		7.26	0.26	RHA 93			
Clostridium clostridioforme		2.11	0.27				
57575500	**Good identification**						**57575500**
					GRA	SPO CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium butyricum		95.54	0.40	IND 3	99	99	1	
Bacteroides thetaiotaomicron		1.99	0.13	SAL 99	1	1	89	
Actinomyces israelii		1.53	0.30		99	1	1	
57575502	**Low discrimination**							**57575502**
					GRA	SPO	CAT	
Bacteroides thetaiotaomicron		65.30	0.47	RHA 93	1	1	89	
Bifidobacterium adolescentis		13.58	0.48	NAG 99	99	1	1	
Clostridium butyricum		12.35	0.33		99	99	1	
57575512	**Low discrimination**							**57575512**
					GRA	CAT		
Bacteroides thetaiotaomicron		49.68	0.40	RHA 93	1	89		
Bifidobacterium adolescentis		32.45	0.50	NAG 99	99	1		
Bacteroides ovatus		8.04	0.26		1	68		
57575520	**Low discrimination**							**57575520**
					GRA	SPO		
Bacteroides thetaiotaomicron		86.28	0.47	RHA 93	1	1		
Bacteroides ovatus		8.15	0.29	BGL 99	1	1		
Clostridium butyricum		3.65	0.22		99	99		
57575522	**Good identification**							**57575522**
					CAT			
Bacteroides thetaiotaomicron		90.20	0.81	RHA 93	89			
Bacteroides ovatus		8.52	0.63		68			
Bacteroides uniformis		1.11	0.45		5			
57575523	**Acceptable identification**							**57575523**
					CAT			
Bacteroides thetaiotaomicron		90.23	0.47	RHA 93	89			
Bacteroides ovatus		8.52	0.29	URE 1	68			
Bacteroides uniformis		1.11	0.11		5			
57575530	**Low discrimination**							**57575530**
					GRA	CAT		
Bacteroides thetaiotaomicron		84.23	0.40	RHA 93	1	89		
Bacteroides ovatus		13.64	0.26	BGL 99	1	68		
Bifidobacterium longum subsp. longum		1.47	0.21		99	1		
57575532	**Low discrimination**							**57575532**
Bacteroides thetaiotaomicron		85.74	0.74	RHA 93				
Bacteroides ovatus		13.88	0.60					
Parabacteroides distasonis		0.19	0.44					
57575533	**Low discrimination**							**57575533**
Bacteroides thetaiotaomicron		85.90	0.40	RHA 93				
Bacteroides ovatus		13.91	0.26	URE 1				
Parabacteroides distasonis		0.19	0.10					
57575562	**Low discrimination**							**57575562**
					CAT			
Bacteroides thetaiotaomicron		71.18	0.47	RHA 93	89			
Bacteroides ovatus		27.74	0.40	SOR 1	68			
Bacteroides uniformis		0.87	0.11		5			
57575572	**Low discrimination**							**57575572**
Bacteroides thetaiotaomicron		59.87	0.40	RHA 93				
Bacteroides ovatus		40.00	0.37	SOR 1				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		0.13	0.10			
57575602	**Low discrimination**					**57575602**
Bacteroides thetaiotaomicron		85.07	0.37	LAC 98		
Clostridium clostridioforme		9.25	0.31	NAG 99		
Bacteroides ovatus		5.69	0.17			
57575612	**Low discrimination**					**57575612**
Clostridium clostridioforme		53.32	0.42	IND 1		
Bacteroides thetaiotaomicron		41.88	0.30	SAL 88		
Bacteroides ovatus		4.80	0.14			
57575620	**Acceptable identification**					**57575620**
Bacteroides thetaiotaomicron		93.73	0.37	LAC 98		
Bacteroides ovatus		6.27	0.17	BGL 99		
57575622	**Good identification**					**57575622**
					CAT	
Bacteroides thetaiotaomicron		93.51	0.71	LAC 98	89	
Bacteroides ovatus		6.25	0.51		68	
Bacteroides uniformis		0.12	0.19		5	
57575623	**Acceptable identification**					**57575623**
Bacteroides thetaiotaomicron		93.73	0.37	LAC 98		
Bacteroides ovatus		6.27	0.17	URE 1		
57575632	**Low discrimination**					**57575632**
Bacteroides thetaiotaomicron		88.67	0.64	LAC 98		
Bacteroides ovatus		10.16	0.48			
Clostridium clostridioforme		1.14	0.42			
57575662	**Low discrimination**					**57575662**
Bacteroides thetaiotaomicron		78.38	0.37	LAC 98		
Bacteroides ovatus		21.62	0.27	SOR 1		
57575702	**Low discrimination**					**57575702**
Bacteroides thetaiotaomicron		87.35	0.66	NAG 99		
Bacteroides ovatus		11.80	0.51			
Clostridium clostridioforme		0.65	0.40			
57575712	**Low discrimination**					**57575712**
Bacteroides thetaiotaomicron		75.78	0.59	NAG 99		
Bacteroides ovatus		17.55	0.48			
Clostridium clostridioforme		6.59	0.51			
57575720	**Low discrimination**					**57575720**
					CAT	
Bacteroides thetaiotaomicron		87.73	0.66	BGL 99	89	
Bacteroides ovatus		11.85	0.51		68	
Bacteroides vulgatus		0.26	0.20		3	
57575722	**Low discrimination**					**57575722**
					CAT	
Bacteroides thetaiotaomicron		87.97	1.00		89	
Bacteroides ovatus		11.88	0.85		68	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		0.11	0.48		5	
57575723	**Low discrimination**					**57575723**
					CAT	
Bacteroides thetaiotaomicron		87.97	0.66	URE 1	89	
Bacteroides ovatus		11.88	0.51		68	
Bacteroides uniformis		0.11	0.14		5	
57575730	**Low discrimination**					**57575730**
Bacteroides thetaiotaomicron		81.11	0.59	BGL 99		
Bacteroides ovatus		18.78	0.48			
Clostridium clostridioforme		0.07	0.17			
57575732	**Low discrimination**					**57575732**
Bacteroides thetaiotaomicron		81.10	0.93			
Bacteroides ovatus		18.78	0.82			
Clostridium clostridioforme		0.07	0.51			
57575733	**Low discrimination**					**57575733**
Bacteroides thetaiotaomicron		81.11	0.59	URE 1		
Bacteroides ovatus		18.78	0.48			
Clostridium clostridioforme		0.07	0.17			
57575762	**Low discrimination**					**57575762**
					CAT	
Bacteroides thetaiotaomicron		64.14	0.66	SOR 1	89	
Bacteroides ovatus		35.75	0.62		68	
Bacteroides uniformis		0.08	0.14		5	
57575772	**Low discrimination**					**57575772**
Bacteroides thetaiotaomicron		51.11	0.59	SOR 1		
Bacteroides ovatus		48.82	0.59			
Clostridium clostridioforme		0.05	0.17			
57576722	**Low discrimination**					**57576722**
Bacteroides ovatus		81.68	0.43	SAL 88		
Bacteroides thetaiotaomicron		18.32	0.32	GAL 99		
57576732	**Low discrimination**					**57576732**
Bacteroides ovatus		88.13	0.40	SAL 88		
Bacteroides thetaiotaomicron		11.53	0.25	GAL 99		
Clostridium clostridioforme		0.33	0.09			
57577002	**Good identification**					**57577002**
Actinomyces israelii		100.0	0.37	IND 1		
57577100	**Low discrimination**					**57577100**
					SPO	
Actinomyces israelii		88.80	0.45	IND 1	1	
Clostridium butyricum		9.92	0.07	SAL 92	99	
Eubacterium saburreum		1.29	0.09		1	
57577102	**Low discrimination**					**57577102**
					SPO	
Actinomyces israelii		73.73	0.48	IND 1	1	
Bifidobacterium adolescentis		23.24	0.33	SAL 92	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Clostridium butyricum	2.14	0.00		99		
57577140	**Good identification**						**57577140**
	Actinomyces israelii	100.0	0.38	IND 1			
57577142	**Acceptable identification**						**57577142**
	Actinomyces israelii	90.90	0.41	IND 1			
	Bifidobacterium adolescentis	9.11	0.17	SAL 92			
57577322	**Acceptable identification**						**57577322**
	Bacteroides ovatus	92.70	0.51	SAL 88			
	Bacteroides thetaiotaomicron	6.79	0.32	ARA 97			
	Parabacteroides distasonis	0.39	0.27				
57577332	**Good identification**						**57577332**
	Bacteroides ovatus	95.59	0.48	SAL 88			
	Bacteroides thetaiotaomicron	4.09	0.25	ARA 97			
	Parabacteroides distasonis	0.33	0.22				
57577402	**Low discrimination**						**57577402**
					SPO		
	Actinomyces israelii	86.41	0.36	IND 1	1		
	Bifidobacterium adolescentis	9.00	0.13	SAL 92	1		
	Clostridium butyricum	4.59	-.07		99		
57577502	**Low discrimination**						**57577502**
					GRA		
	Bifidobacterium adolescentis	56.68	0.47	IND 1	99		
	Actinomyces israelii	21.99	0.46		99		
	Bacteroides ovatus	10.49	0.21		1		
57577512	**Low discrimination**						**57577512**
					GRA		
	Bifidobacterium adolescentis	54.39	0.48	IND 1	99		
	Bifidobacterium dentium	36.22	0.27		99		
	Bacteroides ovatus	5.49	0.18		1		
57577522	**Low discrimination**						**57577522**
					CAT		
	Bacteroides ovatus	75.27	0.55	SAL 88	68		
	Bacteroides thetaiotaomicron	24.15	0.47	RHA 95	89		
	Bacteroides uniformis	0.30	0.11		5		
57577532	**Low discrimination**						**57577532**
					GRA	CAT	
	Bacteroides ovatus	76.51	0.52	SAL 88	1	68	
	Bacteroides thetaiotaomicron	14.32	0.40	RHA 95	1	89	
	Bifidobacterium dentium	9.06	0.32		99	1	
57577542	**Low discrimination**						**57577542**
					COC		
	Actinomyces israelii	46.76	0.39	IND 1	1		
	Bifidobacterium adolescentis	38.31	0.31	SAL 92	1		
	Blautia producta	9.21	0.07		99		
57577622	**Low discrimination**						**57577622**
	Bacteroides ovatus	68.81	0.43	SAL 88			
	Bacteroides thetaiotaomicron	31.19	0.37	LAC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57577632	**Low discrimination**						**57577632**
Bacteroides ovatus		78.88	0.40	SAL 88			
Bacteroides thetaiotaomicron		20.85	0.30	LAC 99			
Clostridium clostridioforme		0.27	0.08				
57577702	**Low discrimination**						**57577702**
					GRA	CAT	
Bacteroides ovatus		80.83	0.43	SAL 88	1	68	
Bacteroides thetaiotaomicron		18.13	0.32	NAG 99	1	89	
Actinomyces israelii		0.67	0.27		99	1	
57577712	**Low discrimination**						**57577712**
Bacteroides ovatus		86.88	0.40	SAL 88			
Bacteroides thetaiotaomicron		11.37	0.25	NAG 99			
Clostridium clostridioforme		0.99	0.17				
57577720	**Low discrimination**						**57577720**
Bacteroides ovatus		81.68	0.43	SAL 88			
Bacteroides thetaiotaomicron		18.32	0.32	BGL 99			
57577722	**Low discrimination**						**57577722**
					CAT		
Bacteroides ovatus		81.65	0.77	SAL 88	68		
Bacteroides thetaiotaomicron		18.32	0.66		89		
Bacteroides uniformis		0.02	0.14		5		
57577723	**Low discrimination**						**57577723**
Bacteroides ovatus		81.68	0.43	SAL 88			
Bacteroides thetaiotaomicron		18.32	0.32	URE 1			
57577730	**Low discrimination**						**57577730**
Bacteroides ovatus		88.43	0.40	SAL 88			
Bacteroides thetaiotaomicron		11.57	0.25	BGL 99			
57577732	**Low discrimination**						**57577732**
Bacteroides ovatus		88.41	0.74	SAL 88			
Bacteroides thetaiotaomicron		11.57	0.59				
Parabacteroides distasonis		0.01	0.18				
57577733	**Low discrimination**						**57577733**
Bacteroides ovatus		88.43	0.40	SAL 88			
Bacteroides thetaiotaomicron		11.57	0.25	URE 1			
57577762	**Acceptable identification**						**57577762**
Bacteroides ovatus		94.84	0.53	SAL 88			
Bacteroides thetaiotaomicron		5.16	0.32	SOR 4			
57577772	**Good identification**						**57577772**
					COC	GRA	
Bacteroides ovatus		96.83	0.50	SAL 88	1	1	
Bacteroides thetaiotaomicron		3.07	0.25	SOR 4	1	1	
Blautia producta		0.10	-.01		99	99	
57610000	**Very good identification**						**57610000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		100.0	0.57	IND 1		
57610002	**Good identification**					**57610002**
Fusobacterium mortiferum		100.0	0.40	IND 1		
57610100	**Very good identification**					**57610100**
Fusobacterium mortiferum		100.0	0.63	IND 1		
57610102	**Good identification**					**57610102**
					GRA	
Fusobacterium mortiferum		99.19	0.47	IND 1	1	
Bifidobacterium longum subsp. infantis		0.81	-.07	BGL 10	99	
57611000	**Very good identification**					**57611000**
Fusobacterium mortiferum		95.96	0.60	IND 1		
Eubacterium saburreum		4.04	0.34			
57611002	**Acceptable identification**					**57611002**
					GRA	
Fusobacterium mortiferum		92.62	0.43	IND 1	1	
Eubacterium saburreum		6.20	0.21	BGL 10	85	
Bifidobacterium longum subsp. infantis		1.18	-.07		99	
57611100	**Very good identification**					**57611100**
					GRA	
Fusobacterium mortiferum		95.41	0.66	IND 1	1	
Eubacterium saburreum		4.02	0.41		85	
Bifidobacterium longum subsp. infantis		0.57	0.10		99	
57611102	**Low discrimination**					**57611102**
					GRA	
Fusobacterium mortiferum		61.81	0.50	IND 1	1	
Bifidobacterium longum subsp. infantis		33.50	0.27	BGL 10	99	
Eubacterium saburreum		4.14	0.28		85	
57611110	**Low discrimination**					**57611110**
Eubacterium saburreum		77.34	0.39	SAL 1		
Fusobacterium mortiferum		22.66	0.32			
57614000	**Good identification**					**57614000**
Fusobacterium mortiferum		100.0	0.37	IND 1		
57614100	**Good identification**					**57614100**
Fusobacterium mortiferum		100.0	0.44	IND 1		
57615000	**Low discrimination**					**57615000**
Fusobacterium mortiferum		76.84	0.40	IND 1		
Eubacterium saburreum		23.16	0.30	XYL 7		
57615100	**Low discrimination**					**57615100**
					GRA	
Fusobacterium mortiferum		73.20	0.47	IND 1	1	
Eubacterium saburreum		22.07	0.36	XYL 7	85	
Bifidobacterium longum subsp. infantis		4.74	0.09		99	
57615332	**Acceptable identification**					**57615332**
					CAT	
Parabacteroides distasonis		93.56	0.41	IND 1	76	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		3.76	0.01		89	
Bacteroides uniformis		2.68	-.05		5	
57615522	**Good identification**					**57615522**
					CAT	
Bacteroides uniformis		95.28	0.45	MLT 92	5	
Bacteroides thetaiotaomicron		3.10	0.23	CEL 99	89	
Parabacteroides distasonis		1.53	0.33		76	
57615722	**Low discrimination**					**57615722**
					CAT	
Bacteroides uniformis		73.77	0.48	MLT 92	5	
Bacteroides thetaiotaomicron		23.08	0.42	CEL 99	89	
Parabacteroides distasonis		2.57	0.42		76	
57630000	**Good identification**					**57630000**
Fusobacterium mortiferum		100.0	0.48	IND 1		
57630100	**Good identification**					**57630100**
Fusobacterium mortiferum		100.0	0.54	IND 1		
57630102	**Good identification**					**57630102**
Fusobacterium mortiferum		100.0	0.38	IND 1		
57631000	**Acceptable identification**					**57631000**
Fusobacterium mortiferum		91.40	0.51	IND 1		
Eubacterium saburreum		8.60	0.31			
57631100	**Good identification**					**57631100**
Fusobacterium mortiferum		91.40	0.57	IND 1		
Eubacterium saburreum		8.60	0.38			
57631102	**Low discrimination**					**57631102**
					GRA	
Fusobacterium mortiferum		85.64	0.41	IND 1	1	
Eubacterium saburreum		12.79	0.25	BGL 10	85	
Bifidobacterium longum subsp. infantis		1.57	-.07		99	
57631110	**Low discrimination**					**57631110**
Eubacterium saburreum		88.40	0.36	SAL 1		
Fusobacterium mortiferum		11.60	0.23			
57631322	**Acceptable identification**					**57631322**
Parabacteroides distasonis		94.27	0.40	IND 1		
Bacteroides thetaiotaomicron		5.73	0.03	XYL 90		
57631332	**Good identification**					**57631332**
Parabacteroides distasonis		95.79	0.36	IND 1		
Bacteroides thetaiotaomicron		4.21	-.04	XYL 90		
57631722	**Acceptable identification**					**57631722**
Bacteroides thetaiotaomicron		90.38	0.37	MLT 94		
Parabacteroides distasonis		9.20	0.37	XYL 96		
Bacteroides ovatus		0.42	-.03			
57635100	**Low discrimination**					**57635100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		59.78	0.38	IND 1			
Eubacterium saburreum		40.22	0.33	XYL 7			
57635122	**Good identification**						**57635122**
					GRA	CAT	
Parabacteroides distasonis		95.08	0.48	IND 1	1	76	
Bacteroides thetaiotaomicron		3.48	0.08		1	89	
Bifidobacterium longum subsp. infantis		0.74	-.04		99	1	
57635132	**Good identification**						**57635132**
Parabacteroides distasonis		97.43	0.44	IND 1			
Bacteroides thetaiotaomicron		2.58	0.01				
57635322	**Low discrimination**						**57635322**
Parabacteroides distasonis		85.11	0.56	IND 1			
Bacteroides thetaiotaomicron		13.80	0.27				
Bacteroides ovatus		0.81	0.06				
57635332	**Low discrimination**						**57635332**
Parabacteroides distasonis		88.59	0.52	IND 1			
Bacteroides thetaiotaomicron		10.37	0.20				
Bacteroides ovatus		1.04	0.03				
57635522	**Low discrimination**						**57635522**
					CAT		
Bacteroides thetaiotaomicron		82.93	0.42	MLT 94	89		
Parabacteroides distasonis		14.03	0.45	RHA 93	76		
Bacteroides uniformis		1.94	0.11		5		
57635722	**Good identification**						**57635722**
Bacteroides thetaiotaomicron		94.34	0.61	MLT 94			
Parabacteroides distasonis		3.60	0.53				
Bacteroides ovatus		1.80	0.32				
57635732	**Acceptable identification**						**57635732**
Bacteroides thetaiotaomicron		91.81	0.54	MLT 94			
Parabacteroides distasonis		4.86	0.48				
Bacteroides ovatus		3.00	0.29				
57645522	**Good identification**						**57645522**
					CAT		
Bacteroides uniformis		99.93	0.45	MLT 92	5		
Bacteroides thetaiotaomicron		0.07	-.06	FRU 99	89		
57645722	**Good identification**						**57645722**
					CAT		
Bacteroides uniformis		98.57	0.48	MLT 92	5		
Bacteroides ovatus		0.77	0.15	FRU 99	68		
Bacteroides thetaiotaomicron		0.66	0.13		89		
57650000	**Good identification**						**57650000**
Fusobacterium mortiferum		100.0	0.48	IND 1			
57650100	**Good identification**						**57650100**
Fusobacterium mortiferum		100.0	0.55	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57650102	**Good identification**						**57650102**
Fusobacterium mortiferum		100.0	0.38	IND 1			
57651000	**Good identification**						**57651000**
					GRA		
Fusobacterium mortiferum		97.60	0.51	IND 1	1		
Lactobacillus catenaformis		1.94	0.12		99		
Lactobacillus jensenii		0.46	0.02		99		
57651002	**Low discrimination**						**57651002**
					GRA	SPO	
Fusobacterium mortiferum		74.72	0.35	IND 1	1	1	
Lactobacillus catenaformis		13.39	0.12	BGL 10	99	1	
Clostridium barati		8.30	0.04		99	99	
57651100	**Very good identification**						**57651100**
					GRA		
Fusobacterium mortiferum		97.18	0.58	IND 1	1		
Lactobacillus catenaformis		1.24	0.15		99		
Collinsella aerofaciens		0.82	-.03		99		
57651102	**Low discrimination**						**57651102**
					GRA		
Fusobacterium mortiferum		53.82	0.41	IND 1	1		
Lactobacillus acidophilus		12.75	0.18	BGL 10	99		
Bifidobacterium breve		12.30	0.27		99		
57651522	**Good identification**						**57651522**
					CAT		
Bacteroides uniformis		99.55	0.45	MLT 92	5		
Bacteroides thetaiotaomicron		0.29	0.05	XYL 99	89		
Parabacteroides distasonis		0.10	0.12		76		
57651722	**Good identification**						**57651722**
					CAT		
Bacteroides uniformis		96.37	0.48	MLT 92	5		
Bacteroides thetaiotaomicron		2.67	0.24	XYL 99	89		
Bacteroides ovatus		0.76	0.15		68		
57654100	**Good identification**						**57654100**
Fusobacterium mortiferum		100.0	0.35	IND 1			
57654522	**Good identification**						**57654522**
					CAT		
Bacteroides uniformis		99.93	0.45	MLT 92	5		
Bacteroides thetaiotaomicron		0.07	-.06	GAL 99	89		
57654722	**Good identification**						**57654722**
					CAT		
Bacteroides uniformis		98.57	0.48	MLT 92	5		
Bacteroides ovatus		0.77	0.15	GAL 99	68		
Bacteroides thetaiotaomicron		0.66	0.13		89		
57655100	**Low discrimination**						**57655100**
					GRA	SPO	
Fusobacterium mortiferum		87.65	0.38	IND 1	1	1	
Clostridium butyricum		9.43	0.03	XYL 7	99	99	
Bacteroides uniformis		2.92	-.07		1	1	
57655122	**Very good identification**						**57655122**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides uniformis		99.69	0.61	MLT 92				
Parabacteroides distasonis		0.16	0.32	ARA 92				
Prevotella oralis		0.10	0.16					
57655322	**Very good identification**							**57655322**
Bacteroides uniformis		99.26	0.64	MLT 92				
Parabacteroides distasonis		0.35	0.41	ARA 92				
Bacteroides ovatus		0.28	0.23					
57655422	**Good identification**							**57655422**
					CAT			
Bacteroides uniformis		99.93	0.51	MLT 92	5			
Bacteroides thetaiotaomicron		0.07	-.01	LAC 98	89			
57655502	**Good identification**							**57655502**
					GRA	SPO		
Bacteroides uniformis		97.53	0.45	MLT 92	1	1		
Bifidobacterium adolescentis		1.43	0.30	NAG 99	99	1		
Clostridium butyricum		0.97	0.13		99	99		
57655520	**Good identification**							**57655520**
					GRA	CAT	SPO	
Bacteroides uniformis		99.71	0.45	MLT 92	1	5	1	
Clostridium butyricum		0.22	0.01	BGL 99	99	1	99	
Bacteroides thetaiotaomicron		0.07	-.06		1	89	1	
57655522	**Very good identification**							**57655522**
					CAT			
Bacteroides uniformis		99.86	0.79	MLT 92	5			
Bacteroides thetaiotaomicron		0.07	0.28		89			
Bacteroides ovatus		0.06	0.27		68			
57655523	**Good identification**							**57655523**
					CAT			
Bacteroides uniformis		99.93	0.45	MLT 92	5			
Bacteroides thetaiotaomicron		0.07	-.06	URE 1	89			
57655532	**Acceptable identification**							**57655532**
					CAT			
Bacteroides uniformis		93.53	0.45	MLT 92	5			
Bacteroides ovatus		3.52	0.24	MLZ 1	68			
Bacteroides thetaiotaomicron		2.51	0.21		89			
57655562	**Good identification**							**57655562**
					CAT			
Bacteroides uniformis		99.70	0.45	MLT 92	5			
Bacteroides ovatus		0.23	0.03	SOR 1	68			
Bacteroides thetaiotaomicron		0.07	-.06		89			
57655622	**Good identification**							**57655622**
					CAT			
Bacteroides uniformis		98.95	0.53	MLT 92	5			
Bacteroides thetaiotaomicron		0.67	0.19	LAC 98	89			
Bacteroides ovatus		0.38	0.15		68			
57655702	**Good identification**							**57655702**
					CAT			
Bacteroides uniformis		98.51	0.48	MLT 92	5			
Bacteroides ovatus		0.77	0.15	NAG 99	68			
Bacteroides thetaiotaomicron		0.66	0.13		89			
57655720	**Good identification**							**57655720**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
				CAT	
Bacteroides uniformis	98.43	0.48	MLT 92	5	
Bacteroides ovatus	0.77	0.15	BGL 99	68	
Bacteroides thetaiotaomicron	0.66	0.13		89	
57655722 Very good identification					**57655722**
				CAT	
Bacteroides uniformis	98.55	0.82	MLT 92	5	
Bacteroides ovatus	0.77	0.49		68	
Bacteroides thetaiotaomicron	0.66	0.47		89	
57655723 Good identification					**57655723**
				CAT	
Bacteroides uniformis	98.57	0.48	MLT 92	5	
Bacteroides ovatus	0.77	0.15	URE 1	68	
Bacteroides thetaiotaomicron	0.66	0.13		89	
57655732 Low discrimination					**57655732**
				CAT	
Bacteroides uniformis	55.90	0.48	MLT 92	5	
Bacteroides ovatus	28.93	0.46	MLZ 1	68	
Bacteroides thetaiotaomicron	14.45	0.40		89	
57655762 Good identification					**57655762**
				CAT	
Bacteroides uniformis	96.24	0.48	MLT 92	5	
Bacteroides ovatus	3.11	0.25	SOR 1	68	
Bacteroides thetaiotaomicron	0.65	0.13		89	
57657522 Good identification					**57657522**
				CAT	
Bacteroides uniformis	98.09	0.45	MLT 92	5	
Bacteroides ovatus	1.84	0.19	MAN 1	68	
Bacteroides thetaiotaomicron	0.07	-.06		89	
57657722 Low discrimination					**57657722**
				CAT	
Bacteroides uniformis	79.03	0.48	MLT 92	5	
Bacteroides ovatus	20.44	0.41	MAN 1	68	
Bacteroides thetaiotaomicron	0.53	0.13		89	
57657732 Acceptable identification					**57657732**
				CAT	
Bacteroides ovatus	93.14	0.38	MLT 99	68	
Bacteroides uniformis	5.46	0.14	TRE 91	5	
Bacteroides thetaiotaomicron	1.41	0.06		89	
57670000 Good identification					**57670000**
Fusobacterium mortiferum	100.0	0.39	IND 1		
57670100 Good identification					**57670100**
Fusobacterium mortiferum	100.0	0.46	IND 1		
57671000 Good identification					**57671000**
				GRA	
Fusobacterium mortiferum	95.46	0.42	IND 1	1	
Lactobacillus jensenii	4.54	0.10		99	
57671100 Good identification					**57671100**
				GRA SPO	
Fusobacterium mortiferum	98.09	0.49	IND 1	1 1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		1.29	-.02		99	99	
Actinomyces naeslundii		0.62	0.23		99	1	
57671322	**Low discrimination**						**57671322**
					SPO	CAT	
Parabacteroides distasonis		45.31	0.36	IND 1	1	76	
Clostridium ramosum		25.05	0.19	XYL 90	92	1	
Bacteroides thetaiotaomicron		10.87	0.09		1	89	
57671722	**Low discrimination**						**57671722**
					CAT		
Bacteroides thetaiotaomicron		78.83	0.43	MLT 94	89		
Bacteroides ovatus		16.98	0.32	XYL 96	68		
Bacteroides uniformis		2.16	0.14		5		
57671732	**Low discrimination**						**57671732**
Bacteroides thetaiotaomicron		70.49	0.36	MLT 94			
Bacteroides ovatus		26.03	0.29	XYL 96			
Parabacteroides distasonis		2.52	0.27				
57675132	**Low discrimination**						**57675132**
Parabacteroides distasonis		68.03	0.39	IND 1			
Bacteroides ovatus		23.14	0.15				
Bacteroides thetaiotaomicron		7.10	0.06				
57675322	**Low discrimination**						**57675322**
Bacteroides ovatus		43.47	0.40	MLT 99			
Parabacteroides distasonis		24.98	0.52	ARA 97			
Bacteroides thetaiotaomicron		15.98	0.33				
57675332	**Low discrimination**						**57675332**
Bacteroides ovatus		59.11	0.37	MLT 99			
Parabacteroides distasonis		27.43	0.47	ARA 97			
Bacteroides thetaiotaomicron		12.68	0.26				
57675500	**Good identification**						**57675500**
					SPO		
Clostridium butyricum		99.82	0.40	IND 3	99		
Actinomyces israelii		0.19	0.14	MLT 99	1		
57675522	**Low discrimination**						**57675522**
					CAT		
Bacteroides uniformis		39.34	0.45	MLT 92	5		
Bacteroides thetaiotaomicron		36.12	0.47	TRE 1	89		
Bacteroides ovatus		22.44	0.44		68		
57675532	**Low discrimination**						**57675532**
Bacteroides ovatus		49.06	0.41	MLT 99			
Bacteroides thetaiotaomicron		46.07	0.40	RHA 95			
Parabacteroides distasonis		2.73	0.35				
57675622	**Low discrimination**						**57675622**
					CAT		
Bacteroides thetaiotaomicron		63.90	0.38	MLT 94	89		
Bacteroides ovatus		28.10	0.32	LAC 98	68		
Bacteroides uniformis		7.23	0.19		5		
57675722	**Low discrimination**						**57675722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Bacteroides thetaiotaomicron		49.71	0.67	MLT 94	89		
Bacteroides ovatus		44.16	0.66		68		
Bacteroides uniformis		5.63	0.48		5		
57675732	**Low discrimination**						**57675732**
Bacteroides ovatus		59.84	0.63	MLT 99			
Bacteroides thetaiotaomicron		39.29	0.60				
Parabacteroides distasonis		0.53	0.44				
57675762	**Low discrimination**						**57675762**
					CAT		
Bacteroides ovatus		76.55	0.42	MLT 99	68		
Bacteroides thetaiotaomicron		20.89	0.33	SOR 4	89		
Bacteroides uniformis		2.37	0.14		5		
57675772	**Low discrimination**						**57675772**
Bacteroides ovatus		86.11	0.39	MLT 99			
Bacteroides thetaiotaomicron		13.71	0.26	SOR 4			
Parabacteroides distasonis		0.18	0.09				
57677522	**Low discrimination**						**57677522**
					CAT		
Bacteroides ovatus		89.24	0.36	MLT 99	68		
Bacteroides uniformis		4.74	0.11	RHA 95	5		
Bacteroides thetaiotaomicron		4.35	0.13		89		
57677722	**Very good identification**						**57677722**
					CAT		
Bacteroides ovatus		96.31	0.58	MLT 99	68		
Bacteroides thetaiotaomicron		3.29	0.33		89		
Bacteroides uniformis		0.37	0.14		5		
57677732	**Good identification**						**57677732**
					GRA	CAT	
Bacteroides ovatus		97.94	0.55	MLT 99	1	68	
Bacteroides thetaiotaomicron		1.95	0.26		1	89	
Bifidobacterium dentium		0.08	-.03		99	1	
57701000	**Good identification**						**57701000**
Eubacterium saburreum		100.0	0.37	SAL 1			
57701010	**Good identification**						**57701010**
Eubacterium saburreum		100.0	0.35	SAL 1			
57701100	**Low discrimination**						**57701100**
					GRA		
Eubacterium saburreum		88.64	0.43	SAL 1	85		
Bifidobacterium longum subsp. infantis		9.40	0.10		99		
Fusobacterium mortiferum		1.97	0.16		1		
57701110	**Good identification**						**57701110**
Eubacterium saburreum		100.0	0.41	SAL 1			
57705100	**Low discrimination**						**57705100**
Eubacterium saburreum		86.13	0.38	SAL 1			
Bifidobacterium longum subsp. infantis		13.87	0.09				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
57705110	**Good identification**						**57705110**
Eubacterium saburreum		100.0	0.37	SAL 1			
57710000	**Low discrimination**						**57710000**
Fusobacterium mortiferum		60.53	0.41	IND 1			
Prevotella intermedia		38.22	0.26	MLT 11			
Eubacterium saburreum		1.25	0.11				
57710100	**Acceptable identification**						**57710100**
					GRA		
Fusobacterium mortiferum		91.50	0.47	IND 1	1		
Bifidobacterium longum subsp. infantis		6.61	0.10	MLT 11	99		
Eubacterium saburreum		1.89	0.17		85		
57711000	**Low discrimination**						**57711000**
Eubacterium saburreum		46.46	0.45	SAL 1			
Fusobacterium mortiferum		34.07	0.44				
Prevotella intermedia		14.34	0.26				
57711010	**Good identification**						**57711010**
Eubacterium saburreum		100.0	0.43	SAL 1			
57711022	**Low discrimination**						**57711022**
					GRA		
Prevotella oralis		51.64	0.36	IND 1	1		
Bifidobacterium longum subsp. infantis		46.06	0.32	LAC 94	99		
Eubacterium saburreum		0.97	0.19		85		
57711100	**Low discrimination**						**57711100**
					GRA		
Bifidobacterium longum subsp. infantis		65.80	0.44	IND 1	99		
Eubacterium saburreum		18.81	0.51	BGL 91	85		
Fusobacterium mortiferum		13.79	0.50		1		
57711102	**Very good identification**						**57711102**
					GRA		
Bifidobacterium longum subsp. infantis		97.14	0.62	IND 1	99		
Prevotella oralis		1.61	0.35		1		
Eubacterium saburreum		0.49	0.38		85		
57711110	**Acceptable identification**						**57711110**
					GRA		
Eubacterium saburreum		94.49	0.49	SAL 1	85		
Bifidobacterium longum subsp. infantis		4.08	0.10		99		
Fusobacterium mortiferum		0.86	0.16		1		
57711120	**Low discrimination**						**57711120**
					GRA		
Bifidobacterium longum subsp. infantis		80.65	0.49	IND 1	99		
Prevotella oralis		7.62	0.35	BGL 91	1		
Prevotella melaninogenica		5.54	0.29		1		
57711122	**Low discrimination**						**57711122**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		84.10	0.66	IND 1	99	1	
Prevotella oralis		14.92	0.56		1	1	
Capnocytophaga ochracea		0.34	0.29		1	90	
57711130	**Low discrimination**						**57711130**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		63.81	0.37	SAL 1		
Bifidobacterium longum subsp. infantis		27.76	0.15			
Bifidobacterium breve		3.90	0.15			
57711322	**Low discrimination**					**57711322**
					GRA	
Prevotella oralis		83.98	0.50	IND 1	1	
Bifidobacterium longum subsp. infantis		11.16	0.32		99	
Parabacteroides distasonis		3.07	0.38		1	
57711502	**Low discrimination**					**57711502**
					GRA	
Bifidobacterium adolescentis		60.24	0.45	IND 1	99	
Bifidobacterium longum subsp. infantis		37.29	0.27	XYL 87	99	
Bacteroides uniformis		0.69	-.05		1	
57711512	**Good identification**					**57711512**
Bifidobacterium adolescentis		96.91	0.46	IND 1		
Bifidobacterium longum subsp. longum		1.44	0.13	XYL 87		
Clostridium clostridioforme		1.16	0.10			
57711722	**Low discrimination**					**57711722**
					CAT	
Bacteroides thetaiotaomicron		58.14	0.39	TRE 93	89	
Bacteroides uniformis		33.07	0.32	XYL 96	5	
Parabacteroides distasonis		3.91	0.35		76	
57713001	**Good identification**					**57713001**
Actinomyces naeslundii		100.0	0.36	IND 1		
57713003	**Good identification**					**57713003**
Actinomyces naeslundii		100.0	0.36	IND 1		
57713021	**Good identification**					**57713021**
Actinomyces naeslundii		100.0	0.36	IND 1		
57713023	**Good identification**					**57713023**
Actinomyces naeslundii		100.0	0.36	IND 1		
57713100	**Low discrimination**					**57713100**
Actinomyces naeslundii		41.25	0.41	IND 1		
Bifidobacterium longum subsp. infantis		33.05	0.10	ESL 12		
Eubacterium saburreum		9.45	0.17			
57713101	**Good identification**					**57713101**
Actinomyces naeslundii		100.0	0.44	IND 1		
57713103	**Acceptable identification**					**57713103**
Actinomyces naeslundii		94.83	0.44	IND 1		
Bifidobacterium longum subsp. infantis		5.17	-.07	ESL 12		
57713121	**Good identification**					**57713121**
Actinomyces naeslundii		100.0	0.44	IND 1		
57713123	**Acceptable identification**					**57713123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		91.16	0.44	IND 1		
Bifidobacterium longum subsp. infantis		8.84	-.03	ESL 12		
57713502	**Good identification**					**57713502**
Bifidobacterium adolescentis		97.98	0.43	IND 1		
Actinomyces israelii		1.27	0.17	XYL 87		
Bifidobacterium longum subsp. infantis		0.75	-.07			
57713512	**Good identification**					**57713512**
Bifidobacterium adolescentis		100.0	0.45	IND 1		
57715000	**Low discrimination**					**57715000**
					GRA	
Eubacterium saburreum		81.45	0.40	SAL 1	85	
Bifidobacterium longum subsp. infantis		10.20	0.09		99	
Fusobacterium mortiferum		8.35	0.25		1	
57715010	**Good identification**					**57715010**
Eubacterium saburreum		100.0	0.39	SAL 1		
57715100	**Low discrimination**					**57715100**
Bifidobacterium longum subsp. infantis		79.31	0.43	IND 1		
Eubacterium saburreum		14.92	0.46	BGL 91		
Eubacterium contortum		3.25	0.24			
57715102	**Very good identification**					**57715102**
Bifidobacterium longum subsp. infantis		97.06	0.60	IND 1		
Bifidobacterium adolescentis		1.94	0.45			
Eubacterium saburreum		0.32	0.33			
57715110	**Acceptable identification**					**57715110**
Eubacterium saburreum		92.44	0.45	SAL 1		
Bifidobacterium longum subsp. infantis		6.07	0.09			
Bifidobacterium adolescentis		1.50	0.12			
57715112	**Low discrimination**					**57715112**
Bifidobacterium adolescentis		65.26	0.46	IND 1		
Bifidobacterium longum subsp. infantis		27.04	0.26	ARA 87		
Eubacterium saburreum		7.19	0.32			
57715120	**Acceptable identification**					**57715120**
					GRA	
Bifidobacterium longum subsp. infantis		90.90	0.47	IND 1	99	
Bacteroides vulgatus		3.17	0.20	BGL 91	1	
Eubacterium saburreum		1.70	0.33		85	
57715122	**Low discrimination**					**57715122**
					GRA	
Bifidobacterium longum subsp. infantis		85.10	0.64	IND 1	99	
Bacteroides uniformis		9.33	0.45		1	
Prevotella oralis		2.28	0.41		1	
57715300	**Very good identification**					**57715300**
					GRA	
Eubacterium contortum		99.57	0.58	IND 1	99	

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis	0.25	0.09		99		
Bacteroides vulgatus	0.14	0.02		1		
57715302 Low discrimination						**57715302**
				GRA		
Eubacterium contortum	82.44	0.45	IND 1	99		
Bifidobacterium longum subsp. infantis	11.76	0.26		99		
Bacteroides uniformis	3.17	0.14		1		
57715320 Low discrimination						**57715320**
				GRA		
Eubacterium contortum	53.57	0.45	IND 1	99		
Bacteroides vulgatus	41.70	0.36		1		
Bacteroides uniformis	2.06	0.14		1		
57715322 Low discrimination						**57715322**
Bacteroides uniformis	68.08	0.48	CEL 99			
Parabacteroides distasonis	13.72	0.55	ARA 92			
Prevotella oralis	5.16	0.35				
57715332 Low discrimination						**57715332**
Parabacteroides distasonis	67.45	0.50	IND 1			
Bacteroides thetaiotaomicron	11.97	0.21				
Bacteroides ovatus	9.94	0.20				
57715502 Low discrimination						**57715502**
				GRA		
Bifidobacterium adolescentis	77.84	0.59	IND 1	99		
Bacteroides uniformis	13.21	0.29		1		
Bifidobacterium longum subsp. infantis	5.89	0.26		99		
57715512 Very good identification						**57715512**
Bifidobacterium adolescentis	97.23	0.60	IND 1			
Clostridium clostridioforme	1.75	0.27				
Bifidobacterium longum subsp. longum	0.65	0.21				
57715520 Low discrimination						**57715520**
				GRA		
Bacteroides vulgatus	73.72	0.39	IND 1	1		
Bacteroides uniformis	20.52	0.29	RHA 90	1		
Bifidobacterium longum subsp. longum	2.32	0.28		99		
57715522 Good identification						**57715522**
				GRA	CAT	
Bacteroides uniformis	93.88	0.63	CEL 99	1	5	
Bacteroides thetaiotaomicron	4.16	0.43		1	89	
Bifidobacterium longum subsp. infantis	0.75	0.30		99	1	
57715532 Low discrimination						**57715532**
				CAT		
Bacteroides thetaiotaomicron	46.15	0.36	TRE 93	89		
Bacteroides uniformis	27.06	0.29	RHA 93	5		
Bacteroides ovatus	8.76	0.24		68		
57715542 Good identification						**57715542**
				GRA	SPO	
Bifidobacterium adolescentis	96.69	0.43	IND 1	99	1	
Bacteroides uniformis	1.34	-.05	SOR 11	1	1	
Clostridium butyricum	0.99	-.06		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57715552	**Good identification**						**57715552**
Bifidobacterium adolescentis		100.0	0.45	IND 1			
57715622	**Low discrimination**						**57715622**
					CAT		
Bacteroides uniformis		67.24	0.37	CEL 99	5		
Bacteroides thetaiotaomicron		28.66	0.33	LAC 98	89		
Bacteroides ovatus		2.25	0.15		68		
57715700	**Very good identification**						**57715700**
					GRA		
Eubacterium contortum		97.62	0.57	IND 1	99		
Bacteroides vulgatus		2.20	0.21		1		
Bacteroides uniformis		0.09	-.03		1		
57715702	**Low discrimination**						**57715702**
					GRA	CAT	
Eubacterium contortum		52.16	0.44	IND 1	99	1	
Bacteroides uniformis		28.16	0.32		1	5	
Bacteroides thetaiotaomicron		12.00	0.28		1	89	
57715712	**Low discrimination**						**57715712**
					GRA	CAT	
Clostridium clostridioforme		69.81	0.42	IND 1	50	33	
Bacteroides thetaiotaomicron		16.79	0.21	TRE 88	1	89	
Bifidobacterium adolescentis		5.34	0.26		99	1	
57715720	**Low discrimination**						**57715720**
					GRA		
Bacteroides vulgatus		87.46	0.56	IND 1	1		
Eubacterium contortum		6.92	0.44		99		
Bacteroides uniformis		3.74	0.32		1		
57715722	**Low discrimination**						**57715722**
					CAT		
Bacteroides uniformis		66.15	0.66	CEL 99	5		
Bacteroides thetaiotaomicron		28.19	0.62		89		
Bacteroides ovatus		4.46	0.49		68		
57715732	**Low discrimination**						**57715732**
					CAT		
Bacteroides thetaiotaomicron		70.96	0.55	TRE 93	89		
Bacteroides ovatus		19.26	0.46		68		
Bacteroides uniformis		4.32	0.32		5		
57717102	**Low discrimination**						**57717102**
Bifidobacterium adolescentis		55.51	0.43	IND 1			
Bifidobacterium longum subsp. infantis		34.36	0.26	ARA 87			
Actinomyces israelii		10.13	0.37				
57717112	**Good identification**						**57717112**
Bifidobacterium adolescentis		97.82	0.45	IND 1			
Actinomyces israelii		2.18	0.23	ARA 87			
57717502	**Very good identification**						**57717502**
					GRA		
Bifidobacterium adolescentis		97.43	0.57	IND 1	99		
Actinomyces israelii		2.17	0.35		99		
Bacteroides uniformis		0.20	-.05		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
57717512	**Very good identification**					**57717512**
Bifidobacterium adolescentis		99.56	0.59	IND 1		
Actinomyces israelii		0.27	0.21			
Bifidobacterium dentium		0.17	-.07			
57717542	**Acceptable identification**					**57717542**
Bifidobacterium adolescentis		93.44	0.42	IND 1		
Actinomyces israelii		6.56	0.28	SOR 11		
57717552	**Good identification**					**57717552**
Bifidobacterium adolescentis		99.15	0.43	IND 1		
Actinomyces israelii		0.85	0.14	SOR 11		
57717722	**Low discrimination**					**57717722**
					CAT	
Bacteroides ovatus		60.78	0.41	TRE 91	68	
Bacteroides uniformis		27.29	0.32	CEL 99	5	
Bacteroides thetaiotaomicron		11.63	0.28		89	
57717732	**Low discrimination**					**57717732**
					CAT	
Bacteroides ovatus		88.73	0.38	TRE 91	68	
Bacteroides thetaiotaomicron		9.91	0.21	CEL 99	89	
Bacteroides uniformis		0.60	-.03		5	
57721100	**Good identification**					**57721100**
Eubacterium saburreum		100.0	0.40	SAL 1		
57721110	**Good identification**					**57721110**
Eubacterium saburreum		100.0	0.38	SAL 1		
57725100	**Good identification**					**57725100**
Eubacterium saburreum		100.0	0.35	SAL 1		
57725722	**Low discrimination**					**57725722**
Bacteroides thetaiotaomicron		88.27	0.47	FRU 99		
Bacteroides ovatus		10.64	0.32			
Parabacteroides distasonis		0.76	0.28			
57725732	**Low discrimination**					**57725732**
Bacteroides thetaiotaomicron		80.53	0.40	FRU 99		
Bacteroides ovatus		16.64	0.29			
Clostridium clostridioforme		1.87	0.23			
57730100	**Good identification**					**57730100**
Fusobacterium mortiferum		95.59	0.38	IND 1		
Eubacterium saburreum		4.41	0.14	MLT 11		
57731000	**Low discrimination**					**57731000**
					GRA	
Eubacterium saburreum		70.18	0.42	SAL 1	85	
Fusobacterium mortiferum		23.06	0.35		1	
Actinomyces naeslundii		3.81	0.33		99	
57731001	**Good identification**					**57731001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.36	IND 1			
57731003	**Good identification**						**57731003**
Actinomyces naeslundii		100.0	0.36	IND 1			
57731010	**Good identification**						**57731010**
Eubacterium saburreum		98.31	0.40	SAL 1			
Lactobacillus jensenii		1.69	0.02				
57731021	**Good identification**						**57731021**
Actinomyces naeslundii		100.0	0.36	IND 1			
57731023	**Good identification**						**57731023**
Actinomyces naeslundii		100.0	0.36	IND 1			
57731100	**Low discrimination**						**57731100**
					COC	GRA	
Eubacterium saburreum		63.71	0.48	SAL 1	1	85	
Fusobacterium mortiferum		20.93	0.41		1	1	
Atopobium parvulum		6.14	0.21		99	99	
57731101	**Acceptable identification**						**57731101**
Actinomyces naeslundii		90.76	0.44	IND 1			
Eubacterium saburreum		9.24	0.14	ESL 12			
57731103	**Acceptable identification**						**57731103**
Actinomyces naeslundii		94.83	0.44	IND 1			
Bifidobacterium longum subsp. infantis		5.17	-.07	ESL 12			
57731110	**Good identification**						**57731110**
Eubacterium saburreum		99.23	0.46	SAL 1			
Leptotrichia buccalis		0.77	-.11				
57731121	**Good identification**						**57731121**
Actinomyces naeslundii		100.0	0.44	IND 1			
57731122	**Low discrimination**						**57731122**
					GRA		
Prevotella oralis		35.98	0.36	IND 1	1		
Bifidobacterium longum subsp. infantis		32.09	0.32	TRE 6	99		
Parabacteroides distasonis		13.39	0.41		1		
57731123	**Low discrimination**						**57731123**
					GRA		
Actinomyces naeslundii		82.94	0.44	IND 1	99		
Prevotella oralis		9.02	0.02	ESL 12	1		
Bifidobacterium longum subsp. infantis		8.04	-.03		99		
57731132	**Low discrimination**						**57731132**
					GRA		
Parabacteroides distasonis		49.29	0.37	IND 1	1		
Bifidobacterium breve		20.56	0.29	XYL 90	99		
Leptotrichia buccalis		16.06	0.07		1		
57731322	**Low discrimination**						**57731322**
					CAT		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Parabacteroides distasonis		56.29	0.50	IND 1	76			
Prevotella oralis		21.61	0.30	XYL 90	1			
Bacteroides thetaiotaomicron		15.12	0.24		89			
57731332	**Low discrimination**							**57731332**
Parabacteroides distasonis		77.45	0.45	IND 1				
Bacteroides thetaiotaomicron		15.03	0.17	XYL 90				
Clostridium clostridioforme		5.22	0.19					
57731502	**Low discrimination**							**57731502**
					GRA	SPO	CAT	
Bifidobacterium adolescentis		85.65	0.42	IND 1	99	1	1	
Clostridium butyricum		6.05	0.07	XYL 87	99	99	1	
Bacteroides thetaiotaomicron		3.58	0.05		1	1	89	
57731512	**Low discrimination**							**57731512**
					GRA	CAT		
Bifidobacterium adolescentis		86.70	0.43	IND 1	99	1		
Clostridium clostridioforme		11.37	0.25	XYL 87	50	33		
Bacteroides thetaiotaomicron		1.15	-.03		1	89		
57731522	**Low discrimination**							**57731522**
Bacteroides thetaiotaomicron		86.95	0.39	XYL 96				
Parabacteroides distasonis		8.88	0.38	RHA 93				
Bacteroides ovatus		1.78	0.10					
57731712	**Low discrimination**							**57731712**
					GRA	CAT		
Clostridium clostridioforme		83.11	0.40	IND 1	50	33		
Bacteroides thetaiotaomicron		15.26	0.17	XYL 91	1	89		
Bifidobacterium adolescentis		0.87	0.09		99	1		
57731722	**Good identification**							**57731722**
Bacteroides thetaiotaomicron		94.57	0.58	XYL 96				
Bacteroides ovatus		2.76	0.32					
Parabacteroides distasonis		2.18	0.46					
57731732	**Low discrimination**							**57731732**
Bacteroides thetaiotaomicron		87.95	0.51	XYL 96				
Clostridium clostridioforme		4.84	0.40					
Bacteroides ovatus		4.40	0.29					
57733000	**Low discrimination**							**57733000**
Actinomyces naeslundii		86.49	0.38	IND 1				
Actinomyces israelii		13.51	0.15	ESL 12				
57733001	**Good identification**							**57733001**
Actinomyces naeslundii		100.0	0.41	IND 1				
57733002	**Low discrimination**							**57733002**
Actinomyces naeslundii		81.02	0.38	IND 1				
Actinomyces israelii		18.98	0.18	ESL 12				
57733003	**Good identification**							**57733003**
Actinomyces naeslundii		100.0	0.41	IND 1				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57733020	**Good identification**					**57733020**
Actinomyces naeslundii		100.0	0.38	IND 1		
57733021	**Good identification**					**57733021**
Actinomyces naeslundii		100.0	0.41	IND 1		
57733022	**Good identification**					**57733022**
Actinomyces naeslundii		100.0	0.38	IND 1		
57733023	**Good identification**					**57733023**
Actinomyces naeslundii		100.0	0.41	IND 1		
57733100	**Low discrimination**					**57733100**
Actinomyces naeslundii		77.03	0.46	IND 1		
Actinomyces israelii		16.91	0.25	ESL 12		
Eubacterium saburreum		6.06	0.14			
57733101	**Good identification**					**57733101**
Actinomyces naeslundii		100.0	0.49	IND 1		
57733102	**Low discrimination**					**57733102**
Actinomyces naeslundii		43.25	0.46	IND 1		
Bifidobacterium adolescentis		23.77	0.26	ESL 12		
Bifidobacterium breve		16.91	0.24			
57733103	**Good identification**					**57733103**
Actinomyces naeslundii		100.0	0.49	IND 1		
57733120	**Low discrimination**					**57733120**
					SPO	
Actinomyces naeslundii		52.58	0.46	IND 1	1	
Clostridium ramosum		43.81	0.19	ESL 12	92	
Bifidobacterium breve		3.62	0.12		1	
57733121	**Good identification**					**57733121**
Actinomyces naeslundii		100.0	0.49	IND 1		
57733123	**Good identification**					**57733123**
Actinomyces naeslundii		100.0	0.49	IND 1		
57733141	**Good identification**					**57733141**
Actinomyces naeslundii		100.0	0.36	IND 1		
57733143	**Good identification**					**57733143**
Actinomyces naeslundii		100.0	0.36	IND 1		
57733161	**Good identification**					**57733161**
Actinomyces naeslundii		100.0	0.36	IND 1		
57733163	**Good identification**					**57733163**
Actinomyces naeslundii		100.0	0.36	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57733502	**Acceptable identification**						**57733502**
Bifidobacterium adolescentis		93.17	0.40	IND 1			
Actinomyces israelii		6.83	0.27	XYL 87			
57733512	**Good identification**						**57733512**
Bifidobacterium adolescentis		98.34	0.42	IND 1			
Bifidobacterium dentium		1.66	-.07	XYL 87			
57734612	**Good identification**						**57734612**
Clostridium clostridioforme		100.0	0.40	IND 1			
57734632	**Acceptable identification**						**57734632**
Clostridium clostridioforme		91.04	0.40	IND 1			
Bacteroides thetaiotaomicron		8.13	0.11				
Bacteroides ovatus		0.83	-.06				
57734702	**Low discrimination**						**57734702**
Clostridium clostridioforme		85.23	0.37	IND 1			
Bacteroides thetaiotaomicron		13.18	0.13				
Bacteroides ovatus		1.59	-.03				
57734712	**Good identification**						**57734712**
Clostridium clostridioforme		98.43	0.49	IND 1			
Bacteroides thetaiotaomicron		1.30	0.06				
Bacteroides ovatus		0.27	-.06				
57734722	**Low discrimination**						**57734722**
Bacteroides thetaiotaomicron		83.60	0.47	GAL 99			
Bacteroides ovatus		10.07	0.32				
Clostridium clostridioforme		5.46	0.37				
57734732	**Low discrimination**						**57734732**
Bacteroides thetaiotaomicron		50.42	0.40	GAL 99			
Clostridium clostridioforme		38.57	0.49				
Bacteroides ovatus		10.42	0.29				
57735000	**Low discrimination**						**57735000**
					SPO		
Eubacterium saburreum		85.04	0.37	SAL 1	1		
Clostridium butyricum		5.78	0.00		99		
Actinomyces israelii		5.27	0.20		1		
57735010	**Good identification**						**57735010**
Eubacterium saburreum		100.0	0.36	SAL 1			
57735100	**Low discrimination**						**57735100**
					COC	SPO	
Eubacterium saburreum		43.95	0.43	SAL 1	1	1	
Clostridium butyricum		30.73	0.24		1	99	
Atopobium parvulum		14.62	0.25		99	1	
57735102	**Low discrimination**						**57735102**
					SPO		
Bifidobacterium adolescentis		40.18	0.42	IND 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Bifidobacterium longum subsp. infantis		30.52	0.26	ARA 87	1			
Clostridium butyricum		10.19	0.17		99			
57735110	**Acceptable identification**							**57735110**
					SPO			
Eubacterium saburreum		92.55	0.42	SAL 1	1			
Clostridium butyricum		4.16	0.02		99			
Actinomyces israelii		1.80	0.17		1			
57735112	**Low discrimination**							**57735112**
Bifidobacterium adolescentis		79.33	0.43	IND 1				
Eubacterium saburreum		8.74	0.29	ARA 87				
Clostridium clostridioforme		6.72	0.22					
57735122	**Low discrimination**							**57735122**
					GRA	CAT		
Parabacteroides distasonis		66.47	0.58	IND 1	1	76		
Bifidobacterium longum subsp. infantis		14.48	0.30		99	1		
Bacteroides thetaiotaomicron		10.75	0.28		1	89		
57735132	**Low discrimination**							**57735132**
Parabacteroides distasonis		82.35	0.53	IND 1				
Bacteroides thetaiotaomicron		9.62	0.21					
Bacteroides ovatus		4.25	0.15					
57735312	**Low discrimination**							**57735312**
Clostridium clostridioforme		79.23	0.36	IND 1				
Parabacteroides distasonis		10.56	0.27	ARA 94				
Bacteroides thetaiotaomicron		5.46	0.06					
57735322	**Low discrimination**							**57735322**
Parabacteroides distasonis		49.82	0.66	IND 1				
Bacteroides thetaiotaomicron		35.69	0.47					
Bacteroides ovatus		13.17	0.40					
57735332	**Low discrimination**							**57735332**
Parabacteroides distasonis		52.07	0.61	IND 1				
Bacteroides thetaiotaomicron		26.94	0.40					
Bacteroides ovatus		17.04	0.37					
57735500	**Good identification**							**57735500**
					SPO			
Clostridium butyricum		97.02	0.40	IND 3	99			
Actinomyces israelii		1.34	0.29	CEL 99	1			
Bifidobacterium adolescentis		1.12	0.22		1			
57735502	**Low discrimination**							**57735502**
					GRA	SPO	CAT	
Bifidobacterium adolescentis		63.09	0.56	IND 1	99	1	1	
Clostridium butyricum		21.52	0.33		99	99	1	
Bacteroides thetaiotaomicron		9.45	0.28		1	1	89	
57735512	**Low discrimination**							**57735512**
					GRA	CAT		
Bifidobacterium adolescentis		78.33	0.57	IND 1	99	1		
Clostridium clostridioforme		15.52	0.42		50	33		
Bacteroides thetaiotaomicron		3.73	0.21		1	89		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57735522	**Low discrimination**					**57735522**
Bacteroides thetaiotaomicron		87.22	0.62	RHA 93		
Bacteroides ovatus		7.35	0.44			
Parabacteroides distasonis		3.34	0.54			
57735532	**Low discrimination**					**57735532**
Bacteroides thetaiotaomicron		80.31	0.55	RHA 93		
Bacteroides ovatus		11.60	0.41			
Parabacteroides distasonis		4.26	0.49			
57735542	**Low discrimination**					**57735542**
					SPO	
Bifidobacterium adolescentis		76.37	0.40	IND 1	1	
Clostridium butyricum		18.33	0.15	SOR 11	99	
Actinomyces israelii		4.37	0.25		1	
57735552	**Good identification**					**57735552**
					SPO	
Bifidobacterium adolescentis		97.42	0.42	IND 1	1	
Clostridium clostridioforme		1.58	0.08	SOR 11	66	
Clostridium butyricum		1.01	-.07		99	
57735602	**Low discrimination**					**57735602**
Clostridium clostridioforme		73.02	0.37	IND 1		
Bacteroides thetaiotaomicron		25.47	0.18			
Bacteroides ovatus		1.52	-.03			
57735612	**Good identification**					**57735612**
Clostridium clostridioforme		96.82	0.48	IND 1		
Bacteroides thetaiotaomicron		2.88	0.11			
Bacteroides ovatus		0.30	-.06			
57735622	**Acceptable identification**					**57735622**
Bacteroides thetaiotaomicron		91.35	0.53	LAC 98		
Bacteroides ovatus		5.45	0.32			
Clostridium clostridioforme		2.65	0.37			
57735632	**Low discrimination**					**57735632**
Bacteroides thetaiotaomicron		69.09	0.46	LAC 98		
Clostridium clostridioforme		23.44	0.48			
Bacteroides ovatus		7.06	0.29			
57735702	**Low discrimination**					**57735702**
Bacteroides thetaiotaomicron		74.60	0.47	NAG 99		
Clostridium clostridioforme		14.61	0.46			
Bacteroides ovatus		8.99	0.32			
57735712	**Low discrimination**					**57735712**
Clostridium clostridioforme		65.02	0.57	IND 1		
Bacteroides thetaiotaomicron		28.34	0.40			
Bacteroides ovatus		5.85	0.29			
57735720	**Low discrimination**					**57735720**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		76.36	0.47	BGL 99	89	
Bacteroides vulgatus		13.15	0.32		3	
Bacteroides ovatus		9.20	0.32		68	
57735722	**Low discrimination**					**57735722**
Bacteroides thetaiotaomicron		88.27	0.82			
Bacteroides ovatus		10.64	0.66			
Parabacteroides distasonis		0.76	0.62			
57735723	**Low discrimination**					**57735723**
Bacteroides thetaiotaomicron		88.27	0.47	URE 1		
Bacteroides ovatus		10.64	0.32			
Parabacteroides distasonis		0.76	0.28			
57735730	**Low discrimination**					**57735730**
Bacteroides thetaiotaomicron		80.24	0.40	BGL 99		
Bacteroides ovatus		16.58	0.29			
Clostridium clostridioforme		1.86	0.23			
57735732	**Low discrimination**					**57735732**
Bacteroides thetaiotaomicron		80.53	0.75			
Bacteroides ovatus		16.64	0.63			
Clostridium clostridioforme		1.87	0.57			
57735733	**Low discrimination**					**57735733**
Bacteroides thetaiotaomicron		80.53	0.40	URE 1		
Bacteroides ovatus		16.64	0.29			
Clostridium clostridioforme		1.87	0.23			
57735762	**Low discrimination**					**57735762**
Bacteroides thetaiotaomicron		66.25	0.47	SOR 1		
Bacteroides ovatus		32.93	0.42			
Parabacteroides distasonis		0.57	0.28			
57735772	**Low discrimination**					**57735772**
Bacteroides thetaiotaomicron		52.99	0.40	SOR 1		
Bacteroides ovatus		45.15	0.39			
Clostridium clostridioforme		1.23	0.23			
57737002	**Good identification**					**57737002**
Actinomyces israelii		100.0	0.36	IND 1		
57737100	**Low discrimination**					**57737100**
					SPO	
Actinomyces israelii		87.28	0.44	IND 1	1	
Clostridium butyricum		11.26	0.07	CEL 93	99	
Eubacterium saburreum		1.46	0.09		1	
57737102	**Low discrimination**					**57737102**
					SPO	
Actinomyces israelii		49.89	0.47	IND 1	1	
Bifidobacterium adolescentis		48.44	0.40	CEL 93	1	
Clostridium butyricum		1.67	0.00		99	
57737112	**Low discrimination**					**57737112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		87.51	0.42	IND 1			
Actinomyces israelii		11.02	0.33	ARA 87			
Bifidobacterium dentium		1.48	-.07				
57737140	**Good identification**						**57737140**
Actinomyces israelii		100.0	0.37	IND 1			
57737142	**Low discrimination**						**57737142**
Actinomyces israelii		76.42	0.40	IND 1			
Bifidobacterium adolescentis		23.58	0.25	CEL 93			
57737502	**Low discrimination**						**57737502**
					SPO		
Bifidobacterium adolescentis		84.58	0.54	IND 1	1		
Actinomyces israelii		10.65	0.45		1		
Clostridium butyricum		3.92	0.17		99		
57737512	**Low discrimination**						**57737512**
Bifidobacterium adolescentis		78.59	0.56	IND 1			
Bifidobacterium dentium		19.63	0.27				
Actinomyces israelii		1.21	0.31				
57737522	**Low discrimination**						**57737522**
					GRA	CAT	
Bacteroides ovatus		69.74	0.36	CEL 99	1	68	
Bacteroides thetaiotaomicron		25.08	0.28	RHA 95	1	89	
Bifidobacterium adolescentis		1.38	0.20		99	1	
57737542	**Low discrimination**						**57737542**
					COC		
Bifidobacterium adolescentis		65.84	0.39	IND 1	1		
Actinomyces israelii		26.08	0.38	SOR 11	1		
Blautia producta		5.94	0.07		99		
57737552	**Low discrimination**						**57737552**
					COC		
Bifidobacterium adolescentis		77.51	0.40	IND 1	1		
Blautia producta		17.16	0.15	SOR 11	99		
Actinomyces israelii		3.75	0.24		1		
57737722	**Low discrimination**						**57737722**
Bacteroides ovatus		79.71	0.58	CEL 99			
Bacteroides thetaiotaomicron		20.04	0.47				
Parabacteroides distasonis		0.17	0.28				
57737732	**Low discrimination**						**57737732**
Bacteroides ovatus		86.75	0.55	CEL 99			
Bacteroides thetaiotaomicron		12.73	0.40				
Clostridium clostridioforme		0.30	0.23				
57745122	**Good identification**						**57745122**
Bacteroides uniformis		98.77	0.45	FRU 99			
Prevotella oralis		0.87	0.16	ARA 92			
Bacteroides ovatus		0.17	0.01				
57745322	**Good identification**						**57745322**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides uniformis		97.19	0.48	FRU 99			
Bacteroides ovatus		2.33	0.23	ARA 92			
Prevotella oralis		0.26	0.10				
57745522	**Very good identification**						**57745522**
					CAT		
Bacteroides uniformis		99.41	0.63	FRU 99	5		
Bacteroides ovatus		0.49	0.27		68		
Bacteroides thetaiotaomicron		0.10	0.15		89		
57745622	**Good identification**						**57745622**
					CAT		
Bacteroides uniformis		95.92	0.37	FRU 99	5		
Bacteroides ovatus		3.20	0.15	LAC 98	68		
Bacteroides thetaiotaomicron		0.88	0.05		89		
57745722	**Good identification**						**57745722**
					CAT		
Bacteroides uniformis		92.86	0.66	FRU 99	5		
Bacteroides ovatus		6.27	0.49		68		
Bacteroides thetaiotaomicron		0.85	0.34		89		
57745732	**Low discrimination**						**57745732**
					CAT		
Bacteroides ovatus		76.41	0.46	FRU 99	68		
Bacteroides uniformis		17.15	0.32	TRE 91	5		
Bacteroides thetaiotaomicron		6.04	0.27		89		
57747722	**Low discrimination**						**57747722**
					CAT		
Bacteroides ovatus		68.82	0.41	FRU 99	68		
Bacteroides uniformis		30.90	0.32	TRE 91	5		
Bacteroides thetaiotaomicron		0.28	0.00		89		
57747732	**Good identification**						**57747732**
Bacteroides ovatus		99.32	0.38	FRU 99			
Bacteroides uniformis		0.68	-.03	TRE 91			
57750100	**Low discrimination**						**57750100**
					GRA		
Fusobacterium mortiferum		88.25	0.39	IND 1	1		
Collinsella aerofaciens		9.07	-.03	MLT 11	99		
Bifidobacterium breve		2.69	0.10		99		
57751022	**Low discrimination**						**57751022**
					GRA	SPO	
Prevotella oralis		82.76	0.45	IND 1	1	1	
Clostridium barati		5.02	0.24	LAC 94	99	99	
Bacteroides uniformis		3.77	0.16		1	1	
57751101	**Low discrimination**						**57751101**
Actinomyces naeslundii		67.55	0.41	IND 1			
Collinsella aerofaciens		25.03	-.03	ESL 12			
Bifidobacterium breve		7.43	0.10				
57751102	**Low discrimination**						**57751102**
					GRA		
Bifidobacterium breve		54.23	0.62	IND 1	99		
Prevotella oralis		22.71	0.44		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		8.90	0.38		99		
57751110	**Acceptable identification**						**57751110**
Bifidobacterium breve		94.39	0.44	IND 1			
Collinsella aerofaciens		3.21	-.03	BGL 91			
Eubacterium saburreum		1.59	0.15				
57751112	**Low discrimination**						**57751112**
					GRA		
Bifidobacterium breve		88.04	0.62	IND 1	99		
Leptotrichia buccalis		7.41	0.24		1		
Bifidobacterium adolescentis		3.97	0.38		99		
57751120	**Low discrimination**						**57751120**
					GRA	SPO	
Clostridium paraputrificum		65.42	0.47	IND 1	99	99	
Prevotella oralis		19.70	0.44	RAF 7	1	1	
Bifidobacterium breve		8.27	0.49		99	1	
57751122	**Low discrimination**						**57751122**
					GRA		
Prevotella oralis		66.53	0.66	IND 1	1		
Bifidobacterium breve		14.86	0.66		99		
Bacteroides uniformis		9.49	0.45		1		
57751130	**Acceptable identification**						**57751130**
					GRA	SPO	
Bifidobacterium breve		90.58	0.49	IND 1	99	1	
Clostridium paraputrificum		7.24	0.13	BGL 91	99	99	
Prevotella oralis		2.18	0.10		1	1	
57751132	**Good identification**						**57751132**
					GRA		
Bifidobacterium breve		93.81	0.66	IND 1	99		
Prevotella oralis		4.24	0.32		1		
Bacteroides uniformis		0.61	0.11		1		
57751140	**Acceptable identification**						**57751140**
Bifidobacterium breve		92.38	0.44	IND 1			
Collinsella aerofaciens		3.15	-.03	BGL 91			
Actinomyces naeslundii		1.00	0.25				
57751142	**Very good identification**						**57751142**
Bifidobacterium breve		96.63	0.62	IND 1			
Lactobacillus acidophilus		1.96	0.23				
Bifidobacterium adolescentis		0.44	0.21				
57751150	**Good identification**						**57751150**
Bifidobacterium breve		100.0	0.44	IND 1			
57751152	**Very good identification**						**57751152**
					GRA		
Bifidobacterium breve		99.36	0.62	IND 1	99		
Bifidobacterium adolescentis		0.55	0.22		99		
Leptotrichia buccalis		0.08	-.11		1		
57751160	**Low discrimination**						**57751160**
					GRA	SPO	
Bifidobacterium breve		60.95	0.49	IND 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium paraputrificum		36.29	0.27	BGL 91	99	99	
Prevotella oralis		1.47	0.10		1	1	
57751162	**Good identification**						**57751162**
					GRA	SPO	
Bifidobacterium breve		91.33	0.66	IND 1	99	1	
Prevotella oralis		4.13	0.32		1	1	
Clostridium paraputrificum		2.31	0.21		99	99	
57751170	**Good identification**						**57751170**
					SPO		
Bifidobacterium breve		99.40	0.49	IND 1	1		
Clostridium paraputrificum		0.60	-.07	BGL 91	99		
57751172	**Very good identification**						**57751172**
					GRA		
Bifidobacterium breve		99.95	0.66	IND 1	99		
Prevotella oralis		0.05	-.03		1		
57751222	**Low discrimination**						**57751222**
					GRA	SPO	
Prevotella oralis		85.61	0.39	IND 1	1	1	
Bacteroides uniformis		12.58	0.19	LAC 94	1	1	
Clostridium barati		1.05	0.05		99	99	
57751302	**Low discrimination**						**57751302**
					GRA		
Prevotella oralis		80.84	0.38	IND 1	1		
Bacteroides uniformis		7.13	0.14	NAG 95	1		
Lactobacillus acidophilus		5.56	0.19		99		
57751320	**Low discrimination**						**57751320**
					SPO		
Prevotella oralis		76.37	0.38	IND 1	1		
Clostridium ramosum		9.03	0.22	BGL 95	92		
Bacteroides uniformis		6.74	0.14		1		
57751322	**Low discrimination**						**57751322**
					SPO		
Prevotella oralis		65.94	0.60	IND 1	1		
Bacteroides uniformis		30.31	0.48		1		
Clostridium ramosum		1.64	0.33		92		
57751502	**Low discrimination**						**57751502**
					GRA	SPO	
Bifidobacterium adolescentis		56.86	0.50	IND 1	99	1	
Bacteroides uniformis		30.38	0.29	XYL 87	1	1	
Clostridium butyricum		7.96	0.21		99	99	
57751512	**Acceptable identification**						**57751512**
Bifidobacterium adolescentis		93.85	0.52	IND 1			
Bifidobacterium breve		3.14	0.27	XYL 87			
Clostridium clostridioforme		1.76	0.19				
57751522	**Very good identification**						**57751522**
Bacteroides uniformis		98.22	0.63	XYL 99			
Prevotella oralis		0.61	0.32				
Bacteroides ovatus		0.48	0.27				
57751552	**Low discrimination**						**57751552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		78.68	0.36	IND 1		
Bifidobacterium breve		21.32	0.27	XYL 87		
57751622	**Acceptable identification**					**57751622**
					CAT	
Bacteroides uniformis		92.21	0.37	XYL 99	5	
Bacteroides thetaiotaomicron		3.48	0.15	LAC 98	89	
Bacteroides ovatus		3.08	0.15		68	
57751722	**Good identification**					**57751722**
					CAT	
Bacteroides uniformis		90.23	0.66	XYL 99	5	
Bacteroides ovatus		6.09	0.49		68	
Bacteroides thetaiotaomicron		3.41	0.44		89	
57751732	**Low discrimination**					**57751732**
					CAT	
Bacteroides ovatus		62.13	0.46	TRE 91	68	
Bacteroides thetaiotaomicron		20.27	0.37	XYL 99	89	
Bacteroides uniformis		13.95	0.32		5	
57753000	**Low discrimination**					**57753000**
Actinomyces naeslundii		48.79	0.35	IND 1		
Actinomyces israelii		39.42	0.25	ESL 12		
Bifidobacterium breve		11.80	0.10			
57753001	**Good identification**					**57753001**
Actinomyces naeslundii		100.0	0.38	IND 1		
57753003	**Good identification**					**57753003**
Actinomyces naeslundii		100.0	0.38	IND 1		
57753020	**Low discrimination**					**57753020**
					SPO	
Actinomyces naeslundii		53.54	0.35	IND 1	1	
Bifidobacterium breve		23.01	0.15	ESL 12	1	
Clostridium ramosum		11.87	-.01		92	
57753021	**Good identification**					**57753021**
Actinomyces naeslundii		100.0	0.38	IND 1		
57753023	**Good identification**					**57753023**
Actinomyces naeslundii		100.0	0.38	IND 1		
57753100	**Low discrimination**					**57753100**
Bifidobacterium breve		75.91	0.44	IND 1		
Actinomyces israelii		10.25	0.35	BGL 91		
Actinomyces naeslundii		9.03	0.43			
57753101	**Acceptable identification**					**57753101**
Actinomyces naeslundii		94.64	0.46	IND 1		
Bifidobacterium breve		5.36	0.10	ESL 12		
57753102	**Good identification**					**57753102**
Bifidobacterium breve		92.35	0.62	IND 1		
Bifidobacterium adolescentis		2.79	0.35			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.85	0.38				
57753103	**Low discrimination**						**57753103**
Actinomyces naeslundii		63.59	0.46	IND 1			
Bifidobacterium breve		36.41	0.27	ESL 12			
57753110	**Good identification**						**57753110**
Bifidobacterium breve		98.02	0.44	IND 1			
Actinomyces israelii		1.98	0.21	BGL 91			
57753112	**Very good identification**						**57753112**
Bifidobacterium breve		95.94	0.62	IND 1			
Bifidobacterium adolescentis		3.54	0.36				
Actinomyces israelii		0.29	0.24				
57753120	**Low discrimination**						**57753120**
					SPO		
Bifidobacterium breve		58.41	0.49	IND 1	1		
Clostridium ramosum		30.13	0.33	BGL 91	92		
Clostridium paraputrificum		4.67	0.13		99		
57753121	**Low discrimination**						**57753121**
					SPO		
Actinomyces naeslundii		86.76	0.46	IND 1	1		
Bifidobacterium breve		8.73	0.15	ESL 12	1		
Clostridium ramosum		4.51	-.01		92		
57753122	**Low discrimination**						**57753122**
					SPO	GRA	
Bifidobacterium breve		78.24	0.66	IND 1	1	99	
Clostridium ramosum		15.97	0.43		92	50	
Prevotella oralis		3.54	0.32		1	1	
57753130	**Good identification**						**57753130**
					SPO		
Bifidobacterium breve		98.58	0.49	IND 1	1		
Clostridium tertium		0.91	0.05	BGL 91	99		
Clostridium ramosum		0.51	-.01		92		
57753132	**Very good identification**						**57753132**
					SPO		
Bifidobacterium breve		99.48	0.66	IND 1	1		
Clostridium tertium		0.27	0.13		99		
Clostridium ramosum		0.21	0.09		92		
57753140	**Acceptable identification**						**57753140**
Bifidobacterium breve		93.15	0.44	IND 1			
Actinomyces israelii		4.89	0.28	BGL 91			
Actinomyces naeslundii		1.96	0.30				
57753142	**Very good identification**						**57753142**
Bifidobacterium breve		98.56	0.62	IND 1			
Actinomyces israelii		0.77	0.31				
Bifidobacterium adolescentis		0.37	0.19				
57753150	**Good identification**						**57753150**
Bifidobacterium breve		99.22	0.44	IND 1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.78	0.14	BGL 91			
57753152	**Very good identification**						**57753152**
Bifidobacterium breve		99.43	0.62	IND 1			
Bifidobacterium adolescentis		0.45	0.21				
Actinomyces israelii		0.12	0.17				
57753160	**Good identification**						**57753160**
					SPO		
Bifidobacterium breve		97.75	0.49	IND 1	1		
Actinomyces naeslundii		1.15	0.30	BGL 91	1		
Clostridium paraputrificum		0.59	-.07		99		
57753162	**Very good identification**						**57753162**
					SPO		
Bifidobacterium breve		99.57	0.66	IND 1	1		
Clostridium ramosum		0.21	0.09		92		
Actinomyces naeslundii		0.12	0.30		1		
57753170	**Good identification**						**57753170**
Bifidobacterium breve		100.0	0.49	IND 1			
57753172	**Very good identification**						**57753172**
Bifidobacterium breve		100.0	0.66	IND 1			
57753320	**Good identification**						**57753320**
					SPO	GRA	
Clostridium ramosum		97.10	0.35	IND 1	92	50	
Prevotella oralis		1.47	0.04	TRE 94	1	1	
Bifidobacterium breve		1.43	0.15		1	99	
57753322	**Low discrimination**						**57753322**
					SPO	GRA	
Clostridium ramosum		85.62	0.45	IND 1	92	50	
Prevotella oralis		6.13	0.25	TRE 94	1	1	
Bifidobacterium breve		3.20	0.32		1	99	
57753502	**Low discrimination**						**57753502**
Bifidobacterium adolescentis		86.44	0.49	IND 1			
Actinomyces israelii		7.02	0.36	XYL 87			
Bifidobacterium breve		4.33	0.27				
57753512	**Acceptable identification**						**57753512**
Bifidobacterium adolescentis		94.68	0.50	IND 1			
Bifidobacterium breve		3.88	0.27	XYL 87			
Actinomyces israelii		0.94	0.22				
57753722	**Low discrimination**						**57753722**
					CAT		
Bacteroides ovatus		67.66	0.41	TRE 91	68		
Bacteroides uniformis		30.38	0.32	XYL 99	5		
Bacteroides thetaiotaomicron		1.15	0.10		89		
57753732	**Good identification**						**57753732**
					CAT		
Bacteroides ovatus		98.36	0.38	TRE 91	68		
Bacteroides thetaiotaomicron		0.97	0.03	XYL 99	89		
Bacteroides uniformis		0.67	-.03		5		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57754122	**Good identification**						**57754122**
Bacteroides uniformis		98.77	0.45	GAL 99			
Prevotella oralis		0.87	0.16	ARA 92			
Bacteroides ovatus		0.17	0.01				
57754322	**Good identification**						**57754322**
Bacteroides uniformis		97.08	0.48	GAL 99			
Bacteroides ovatus		2.33	0.23	ARA 92			
Prevotella oralis		0.26	0.10				
57754522	**Very good identification**						**57754522**
					CAT		
Bacteroides uniformis		99.38	0.63	GAL 99	5		
Bacteroides ovatus		0.49	0.27		68		
Bacteroides thetaiotaomicron		0.10	0.15		89		
57754622	**Acceptable identification**						**57754622**
Bacteroides uniformis		93.68	0.37	GAL 99			
Bacteroides ovatus		3.13	0.15	LAC 98			
Clostridium clostridioforme		2.34	0.23				
57754712	**Acceptable identification**						**57754712**
Clostridium clostridioforme		93.32	0.43	IND 1			
Bacteroides ovatus		5.46	0.12	TRE 88			
Bacteroides uniformis		1.23	-.03				
57754722	**Good identification**						**57754722**
					CAT		
Bacteroides uniformis		92.73	0.66	GAL 99	5		
Bacteroides ovatus		6.26	0.49		68		
Bacteroides thetaiotaomicron		0.85	0.34		89		
57754732	**Low discrimination**						**57754732**
Bacteroides ovatus		67.74	0.46	TRE 91			
Bacteroides uniformis		15.21	0.32	GAL 99			
Clostridium clostridioforme		11.70	0.43				
57755022	**Good identification**						**57755022**
Bacteroides uniformis		97.16	0.51	LAC 98			
Prevotella oralis		2.66	0.30	ARA 92			
Bacteroides ovatus		0.08	0.01				
57755100	**Low discrimination**						**57755100**
					GRA	SPO	
Clostridium butyricum		84.45	0.37	IND 3	99	99	
Actinomyces israelii		7.17	0.40	TRE 94	99	1	
Bacteroides uniformis		3.04	0.11	ARA 90	1	1	
57755102	**Low discrimination**						**57755102**
					GRA	SPO	
Bacteroides uniformis		67.85	0.45	ARA 92	1	1	
Bifidobacterium adolescentis		14.75	0.50	NAG 99	99	1	
Clostridium butyricum		7.42	0.30		99	99	
57755112	**Low discrimination**						**57755112**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		88.29	0.52	IND 1	99		
Bacteroides uniformis		3.36	0.11	ARA 87	1		
Bifidobacterium breve		2.96	0.27		99		
57755120	**Acceptable identification**						**57755120**
					GRA	SPO	
Bacteroides uniformis		90.63	0.45	ARA 92	1	1	
Prevotella oralis		4.13	0.28	BGL 99	1	1	
Clostridium butyricum		2.22	0.19		99	99	
57755122	**Very good identification**						**57755122**
Bacteroides uniformis		98.67	0.79	ARA 92			
Prevotella oralis		0.86	0.50				
Bacteroides ovatus		0.17	0.35				
57755123	**Good identification**						**57755123**
Bacteroides uniformis		98.77	0.45	ARA 92			
Prevotella oralis		0.87	0.16	URE 1			
Bacteroides ovatus		0.17	0.01				
57755132	**Low discrimination**						**57755132**
Bacteroides uniformis		84.84	0.45	ARA 92			
Bacteroides ovatus		9.77	0.32	MLZ 1			
Parabacteroides distasonis		2.28	0.37				
57755152	**Low discrimination**						**57755152**
Bifidobacterium adolescentis		74.94	0.36	IND 1			
Bifidobacterium breve		20.31	0.27	ARA 87			
Actinomyces israelii		4.75	0.22	SOR 11			
57755162	**Good identification**						**57755162**
					GRA		
Bacteroides uniformis		96.58	0.45	ARA 92	1		
Bifidobacterium breve		1.53	0.32	SOR 1	99		
Prevotella oralis		0.85	0.16		1		
57755222	**Good identification**						**57755222**
Bacteroides uniformis		97.80	0.53	LAC 98			
Bacteroides ovatus		1.16	0.23	ARA 92			
Prevotella oralis		0.83	0.24				
57755302	**Acceptable identification**						**57755302**
Bacteroides uniformis		94.99	0.48	ARA 92			
Bacteroides ovatus		2.28	0.23	NAG 99			
Prevotella oralis		1.35	0.22				
57755320	**Acceptable identification**						**57755320**
Bacteroides uniformis		94.85	0.48	ARA 92			
Bacteroides ovatus		2.28	0.23	BGL 99			
Prevotella oralis		1.34	0.22				
57755322	**Very good identification**						**57755322**
Bacteroides uniformis		97.19	0.82	ARA 92			
Bacteroides ovatus		2.33	0.57				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		0.26	0.44				
57755323	**Good identification**						**57755323**
Bacteroides uniformis		97.19	0.48	ARA 92			
Bacteroides ovatus		2.33	0.23	URE 1			
Prevotella oralis		0.26	0.10				
57755332	**Low discrimination**						**57755332**
Bacteroides ovatus		58.60	0.54	TRE 91			
Bacteroides uniformis		36.99	0.48	ARA 97			
Parabacteroides distasonis		2.16	0.45				
57755362	**Acceptable identification**						**57755362**
Bacteroides uniformis		90.59	0.48	ARA 92			
Bacteroides ovatus		8.97	0.33	SOR 1			
Prevotella oralis		0.25	0.10				
57755422	**Very good identification**						**57755422**
					CAT		
Bacteroides uniformis		99.63	0.69	LAC 98	5		
Bacteroides ovatus		0.24	0.27		68		
Bacteroides thetaiotaomicron		0.10	0.20		89		
57755500	**Acceptable identification**						**57755500**
					GRA	SPO	
Clostridium butyricum		94.39	0.54	IND 3	99	99	
Bacteroides uniformis		4.33	0.29	TRE 94	1	1	
Actinomyces israelii		0.73	0.39		99	1	
57755502	**Low discrimination**						**57755502**
					GRA	SPO	
Bacteroides uniformis		81.87	0.63	NAG 99	1	1	
Bifidobacterium adolescentis		10.36	0.64		99	1	
Clostridium butyricum		7.00	0.47		99	99	
57755512	**Low discrimination**						**57755512**
					GRA	SPO	
Bifidobacterium adolescentis		87.00	0.66	IND 1	99	1	
Bacteroides uniformis		5.68	0.29		1	1	
Clostridium butyricum		2.53	0.25		99	99	
57755520	**Very good identification**						**57755520**
					GRA	SPO	
Bacteroides uniformis		97.37	0.63	BGL 99	1	1	
Clostridium butyricum		1.86	0.36		99	99	
Bacteroides ovatus		0.48	0.27		1	1	
57755522	**Excellent identification**						**57755522**
					CAT		
Bacteroides uniformis		99.40	0.98		5		
Bacteroides ovatus		0.49	0.61		68		
Bacteroides thetaiotaomicron		0.10	0.49		89		
57755523	**Very good identification**						**57755523**
					CAT		
Bacteroides uniformis		99.41	0.63	URE 1	5		
Bacteroides ovatus		0.49	0.27		68		
Bacteroides thetaiotaomicron		0.10	0.15		89		
57755532	**Low discrimination**						**57755532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Bacteroides uniformis		73.08	0.63	MLZ 1	5		
Bacteroides ovatus		23.66	0.58		68		
Bacteroides thetaiotaomicron		2.68	0.42		89		
57755540	**Good identification**						**57755540**
					SPO		
Clostridium butyricum		95.41	0.36	IND 3	99		
Actinomyces israelii		3.30	0.32	TRE 94	1		
Bifidobacterium adolescentis		0.79	0.15	SOR 8	1		
57755542	**Low discrimination**						**57755542**
					GRA	SPO	
Bifidobacterium adolescentis		45.49	0.49	IND 1	99	1	
Bacteroides uniformis		29.39	0.29	SOR 11	1	1	
Clostridium butyricum		21.64	0.29		99	99	
57755552	**Good identification**						**57755552**
					SPO		
Bifidobacterium adolescentis		96.09	0.50	IND 1	1		
Clostridium butyricum		1.97	0.07	SOR 11	99		
Actinomyces israelii		0.75	0.21		1		
57755562	**Very good identification**						**57755562**
					CAT		
Bacteroides uniformis		97.84	0.63	SOR 1	5		
Bacteroides ovatus		1.98	0.37		68		
Bacteroides thetaiotaomicron		0.09	0.15		89		
57755602	**Low discrimination**						**57755602**
Bacteroides uniformis		89.18	0.37	LAC 98			
Clostridium clostridioforme		6.68	0.31	NAG 99			
Bacteroides ovatus		2.98	0.15				
57755612	**Acceptable identification**						**57755612**
Clostridium clostridioforme		90.47	0.42	IND 1			
Bacteroides ovatus		5.91	0.12	TRE 88			
Bacteroides uniformis		2.68	0.03				
57755620	**Good identification**						**57755620**
					CAT		
Bacteroides uniformis		95.36	0.37	LAC 98	5		
Bacteroides ovatus		3.19	0.15	BGL 99	68		
Bacteroides thetaiotaomicron		0.87	0.05		89		
57755622	**Very good identification**						**57755622**
					CAT		
Bacteroides uniformis		95.85	0.71	LAC 98	5		
Bacteroides ovatus		3.20	0.49		68		
Bacteroides thetaiotaomicron		0.88	0.39		89		
57755623	**Good identification**						**57755623**
					CAT		
Bacteroides uniformis		95.92	0.37	LAC 98	5		
Bacteroides ovatus		3.20	0.15	URE 1	68		
Bacteroides thetaiotaomicron		0.88	0.05		89		
57755632	**Low discrimination**						**57755632**
					CAT		
Bacteroides ovatus		56.56	0.46	TRE 91	68		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		25.66	0.37	LAC 99	5	
Bacteroides thetaiotaomicron		9.04	0.32		89	
57755662	**Low discrimination**					**57755662**
					CAT	
Bacteroides uniformis		87.19	0.37	LAC 98	5	
Bacteroides ovatus		12.01	0.25	SOR 1	68	
Bacteroides thetaiotaomicron		0.80	0.05		89	
57755702	**Good identification**					**57755702**
					CAT	
Bacteroides uniformis		92.17	0.66	NAG 99	5	
Bacteroides ovatus		6.22	0.49		68	
Bacteroides thetaiotaomicron		0.84	0.34		89	
57755712	**Low discrimination**					**57755712**
Bacteroides ovatus		54.09	0.46	TRE 91		
Clostridium clostridioforme		28.01	0.51	NAG 99		
Bacteroides uniformis		12.14	0.32			
57755720	**Good identification**					**57755720**
Bacteroides uniformis		91.76	0.66	BGL 99		
Bacteroides ovatus		6.19	0.49			
Bacteroides vulgatus		1.14	0.34			
57755722	**Good identification**					**57755722**
					CAT	
Bacteroides uniformis		92.87	1.00		5	
Bacteroides ovatus		6.27	0.83		68	
Bacteroides thetaiotaomicron		0.85	0.68		89	
57755723	**Good identification**					**57755723**
					CAT	
Bacteroides uniformis		92.87	0.66	URE 1	5	
Bacteroides ovatus		6.27	0.49		68	
Bacteroides thetaiotaomicron		0.85	0.34		89	
57755730	**Low discrimination**					**57755730**
					CAT	
Bacteroides ovatus		76.24	0.46	TRE 91	68	
Bacteroides uniformis		17.12	0.32	BGL 99	5	
Bacteroides thetaiotaomicron		6.03	0.27		89	
57755732	**Low discrimination**					**57755732**
					CAT	
Bacteroides ovatus		76.36	0.80	TRE 91	68	
Bacteroides uniformis		17.14	0.66		5	
Bacteroides thetaiotaomicron		6.04	0.61		89	
57755733	**Low discrimination**					**57755733**
					CAT	
Bacteroides ovatus		76.41	0.46	TRE 91	68	
Bacteroides uniformis		17.15	0.32	URE 1	5	
Bacteroides thetaiotaomicron		6.04	0.27		89	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57755762	**Low discrimination**					**57755762**
					CAT	
Bacteroides uniformis		77.66	0.66	SOR 1	5	
Bacteroides ovatus		21.62	0.59		68	
Bacteroides thetaiotaomicron		0.71	0.34		89	
57755772	**Good identification**					**57755772**
					CAT	
Bacteroides ovatus		93.04	0.56	TRE 91	68	
Bacteroides uniformis		5.06	0.32	SOR 4	5	
Bacteroides thetaiotaomicron		1.78	0.27		89	
57756722	**Low discrimination**					**57756722**
					CAT	
Bacteroides ovatus		68.82	0.41	TRE 91	68	
Bacteroides uniformis		30.90	0.32	GAL 99	5	
Bacteroides thetaiotaomicron		0.28	0.00		89	
57756732	**Good identification**					**57756732**
Bacteroides ovatus		98.81	0.38	TRE 91		
Bacteroides uniformis		0.67	-.03	GAL 99		
Clostridium clostridioforme		0.52	0.09			
57757000	**Good identification**					**57757000**
					SPO	
Actinomyces israelii		96.30	0.43	IND 1	1	
Clostridium butyricum		3.71	-.03		99	
57757002	**Good identification**					**57757002**
Actinomyces israelii		96.58	0.46	IND 1		
Bifidobacterium adolescentis		3.42	0.15			
57757040	**Good identification**					**57757040**
Actinomyces israelii		100.0	0.36	IND 1		
57757042	**Good identification**					**57757042**
Actinomyces israelii		100.0	0.39	IND 1		
57757100	**Low discrimination**					**57757100**
					SPO	
Actinomyces israelii		79.96	0.53	IND 1	1	
Clostridium butyricum		18.46	0.21		99	
Bifidobacterium adolescentis		1.06	0.15		1	
57757102	**Low discrimination**					**57757102**
					SPO	
Actinomyces israelii		49.13	0.56	IND 1	1	
Bifidobacterium adolescentis		43.06	0.49		1	
Clostridium butyricum		2.94	0.14		99	
57757110	**Low discrimination**					**57757110**
					SPO	
Actinomyces israelii		78.99	0.39	IND 1	1	
Bifidobacterium adolescentis		8.58	0.16	MLZ 13	1	
Clostridium butyricum		6.42	-.01		99	
57757112	**Low discrimination**					**57757112**
Bifidobacterium adolescentis		83.49	0.50	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		11.65	0.42	ARA 87		
Bifidobacterium breve		3.42	0.27			
57757122	**Acceptable identification**					**57757122**
					GRA	
Bacteroides uniformis		90.38	0.45	MAN 1	1	
Bacteroides ovatus		5.20	0.27	ARA 92	1	
Bifidobacterium breve		1.43	0.32		99	
57757140	**Acceptable identification**					**57757140**
					SPO	
Actinomyces israelii		93.60	0.46	IND 1	1	
Clostridium butyricum		4.83	0.03		99	
Bifidobacterium breve		1.57	0.10		1	
57757142	**Low discrimination**					**57757142**
Actinomyces israelii		71.19	0.49	IND 1		
Bifidobacterium adolescentis		19.83	0.33			
Bifidobacterium breve		8.03	0.27			
57757302	**Low discrimination**					**57757302**
					GRA	
Actinomyces israelii		36.31	0.37	IND 1	99	
Bacteroides uniformis		33.16	0.14	RHA 7	1	
Bacteroides ovatus		26.26	0.15		1	
57757322	**Low discrimination**					**57757322**
					SPO	
Bacteroides uniformis		55.53	0.48	MAN 1	1	
Bacteroides ovatus		43.99	0.49	ARA 92	1	
Clostridium ramosum		0.17	0.11		92	
57757332	**Good identification**					**57757332**
Bacteroides ovatus		98.02	0.46	TRE 91		
Bacteroides uniformis		1.88	0.14	ARA 97		
Parabacteroides distasonis		0.11	0.11			
57757400	**Low discrimination**					**57757400**
					SPO	
Actinomyces israelii		70.26	0.41	IND 1	1	
Clostridium butyricum		29.74	0.14		99	
57757402	**Low discrimination**					**57757402**
					SPO	
Actinomyces israelii		68.66	0.44	IND 1	1	
Bifidobacterium adolescentis		19.89	0.29		1	
Clostridium butyricum		7.53	0.07		99	
57757442	**Acceptable identification**					**57757442**
Actinomyces israelii		91.57	0.37	IND 1		
Bifidobacterium adolescentis		8.43	0.13			
57757500	**Low discrimination**					**57757500**
					SPO	
Clostridium butyricum		69.41	0.37	IND 3	99	
Actinomyces israelii		27.33	0.52	TRE 94	1	
Bifidobacterium adolescentis		2.97	0.29	MAN 10	1	
57757502	**Low discrimination**					**57757502**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		74.31	0.63	IND 1	99		
Actinomyces israelii		10.37	0.55		99		
Bacteroides uniformis		7.25	0.29		1		
57757510	**Low discrimination**						**57757510**
					SPO		
Actinomyces israelii		28.31	0.37	IND 1	1		
Clostridium butyricum		25.32	0.16	MLZ 13	99		
Bifidobacterium adolescentis		25.15	0.30		1		
57757512	**Good identification**						**57757512**
Bifidobacterium adolescentis		90.19	0.64	IND 1			
Bifidobacterium dentium		7.07	0.27				
Actinomyces israelii		1.54	0.40				
57757522	**Low discrimination**						**57757522**
					GRA		
Bacteroides uniformis		85.78	0.63	MAN 1	1		
Bacteroides ovatus		13.89	0.53		1		
Bifidobacterium adolescentis		0.09	0.29		99		
57757532	**Low discrimination**						**57757532**
					GRA		
Bacteroides ovatus		78.53	0.50	TRE 91	1		
Bifidobacterium dentium		12.82	0.32	RHA 95	99		
Bacteroides uniformis		7.35	0.29		1		
57757540	**Low discrimination**						**57757540**
					COC	SPO	
Actinomyces israelii		40.45	0.45	IND 1	1	1	
Blautia producta		35.19	0.24		99	1	
Clostridium butyricum		22.97	0.19		1	99	
57757542	**Low discrimination**						**57757542**
					SPO		
Bifidobacterium adolescentis		64.72	0.47	IND 1	1		
Actinomyces israelii		28.41	0.48	SOR 11	1		
Clostridium butyricum		4.18	0.12		99		
57757552	**Low discrimination**						**57757552**
					COC		
Bifidobacterium adolescentis		88.53	0.49	IND 1	1		
Blautia producta		6.15	0.15	SOR 11	99		
Actinomyces israelii		4.75	0.33		1		
57757622	**Low discrimination**						**57757622**
					CAT		
Bacteroides ovatus		52.21	0.41	TRE 91	68		
Bacteroides uniformis		47.36	0.37	LAC 99	5		
Bacteroides thetaiotaomicron		0.43	0.05		89		
57757632	**Good identification**						**57757632**
					CAT		
Bacteroides ovatus		97.73	0.38	TRE 91	68		
Bacteroides uniformis		1.34	0.03	LAC 99	5		
Bacteroides thetaiotaomicron		0.47	-.02		89		
57757702	**Low discrimination**						**57757702**
					GRA		
Bacteroides ovatus		65.21	0.41	TRE 91	1		
Bacteroides uniformis		29.28	0.32	NAG 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		2.28	0.35		99		
57757712	**Acceptable identification**						**57757712**
					GRA		
Bacteroides ovatus		91.17	0.38	TRE 91	1		
Bifidobacterium adolescentis		5.63	0.30	NAG 99	99		
Clostridium clostridioforme		1.43	0.17		50		
57757720	**Low discrimination**						**57757720**
Bacteroides ovatus		68.46	0.41	TRE 91			
Bacteroides uniformis		30.74	0.32	BGL 99			
Bacteroides vulgatus		0.38	-.01				
57757722	**Low discrimination**						**57757722**
					CAT		
Bacteroides ovatus		68.81	0.75	TRE 91	68		
Bacteroides uniformis		30.90	0.66		5		
Bacteroides thetaiotaomicron		0.28	0.34		89		
57757723	**Low discrimination**						**57757723**
					CAT		
Bacteroides ovatus		68.82	0.41	TRE 91	68		
Bacteroides uniformis		30.90	0.32	URE 1	5		
Bacteroides thetaiotaomicron		0.28	0.00		89		
57757730	**Good identification**						**57757730**
Bacteroides ovatus		99.32	0.38	TRE 91			
Bacteroides uniformis		0.68	-.03	BGL 99			
57757732	**Very good identification**						**57757732**
					CAT		
Bacteroides ovatus		99.06	0.72	TRE 91	68		
Bacteroides uniformis		0.67	0.32		5		
Bacteroides thetaiotaomicron		0.24	0.27		89		
57757733	**Good identification**						**57757733**
Bacteroides ovatus		99.32	0.38	TRE 91			
Bacteroides uniformis		0.68	-.03	URE 1			
57757762	**Acceptable identification**						**57757762**
					CAT		
Bacteroides ovatus		90.10	0.51	TRE 91	68		
Bacteroides uniformis		9.81	0.32	SOR 4	5		
Bacteroides thetaiotaomicron		0.09	0.00		89		
57757772	**Good identification**						**57757772**
					COC	GRA	
Bacteroides ovatus		99.70	0.48	TRE 91	1	1	
Bacteroides uniformis		0.16	-.03	SOR 4	1	1	
Blautia producta		0.14	-.01		99	99	
57765322	**Low discrimination**						**57765322**
					CAT		
Bacteroides ovatus		89.26	0.40	FRU 99	68		
Bacteroides thetaiotaomicron		5.19	0.19	ARA 97	89		
Bacteroides uniformis		3.72	0.14		5		
57765332	**Acceptable identification**						**57765332**
Bacteroides ovatus		94.48	0.37	FRU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		3.21	0.12	ARA 97			
Parabacteroides distasonis		1.57	0.23				
57765500	**Good identification**						**57765500**
					SPO		
Clostridium butyricum		99.82	0.40	IND 3	99		
Actinomyces israelii		0.19	0.14	FRU 99	1		
57765522	**Low discrimination**						**57765522**
					CAT		
Bacteroides ovatus		67.29	0.44	FRU 99	68		
Bacteroides thetaiotaomicron		17.14	0.34	RHA 95	89		
Bacteroides uniformis		13.71	0.29		5		
57765532	**Low discrimination**						**57765532**
Bacteroides ovatus		85.41	0.41	FRU 99			
Bacteroides thetaiotaomicron		12.69	0.27	RHA 95			
Clostridium clostridioforme		0.84	0.17				
57765722	**Low discrimination**						**57765722**
					CAT		
Bacteroides ovatus		83.76	0.66	FRU 99	68		
Bacteroides thetaiotaomicron		14.92	0.53		89		
Bacteroides uniformis		1.24	0.32		5		
57765732	**Good identification**						**57765732**
Bacteroides ovatus		90.23	0.63	FRU 99			
Bacteroides thetaiotaomicron		9.38	0.46				
Clostridium clostridioforme		0.34	0.32				
57765762	**Good identification**						**57765762**
					CAT		
Bacteroides ovatus		95.53	0.42	FRU 99	68		
Bacteroides thetaiotaomicron		4.13	0.19	SOR 4	89		
Bacteroides uniformis		0.34	-.03		5		
57765772	**Good identification**						**57765772**
Bacteroides ovatus		97.54	0.39	FRU 99			
Bacteroides thetaiotaomicron		2.46	0.12	SOR 4			
57767522	**Good identification**						**57767522**
					GRA	SPO	
Bacteroides ovatus		96.85	0.36	FRU 99	1	1	
Bifidobacterium dentium		1.04	-.03	RHA 95	99	1	
Clostridium butyricum		0.77	-.02		99	99	
57767722	**Very good identification**						**57767722**
					CAT		
Bacteroides ovatus		99.42	0.58	FRU 99	68		
Bacteroides thetaiotaomicron		0.54	0.19		89		
Bacteroides uniformis		0.05	-.03		5		
57767732	**Good identification**						**57767732**
					GRA	CAT	
Bacteroides ovatus		99.60	0.55	FRU 99	1	68	
Bacteroides thetaiotaomicron		0.31	0.12		1	89	
Bifidobacterium dentium		0.09	-.03		99	1	
57770712	**Good identification**						**57770712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		99.18	0.41	IND 1		
Bacteroides ovatus		0.82	-.06	XYL 91		
57770732	**Low discrimination**					**57770732**
Clostridium clostridioforme		46.18	0.41	IND 1		
Bacteroides ovatus		37.67	0.29	XYL 91		
Bacteroides thetaiotaomicron		16.15	0.22			
57771001	**Good identification**					**57771001**
Actinomyces naeslundii		100.0	0.38	IND 1		
57771003	**Good identification**					**57771003**
Actinomyces naeslundii		100.0	0.38	IND 1		
57771021	**Good identification**					**57771021**
Actinomyces naeslundii		100.0	0.38	IND 1		
57771023	**Good identification**					**57771023**
Actinomyces naeslundii		100.0	0.38	IND 1		
57771101	**Acceptable identification**					**57771101**
					SPO	
Actinomyces naeslundii		91.87	0.46	IND 1	1	
Clostridium butyricum		8.13	-.02	ESL 12	99	
57771102	**Low discrimination**					**57771102**
					GRA	
Bifidobacterium breve		45.70	0.59	IND 1	99	
Lactobacillus acidophilus		37.66	0.47		99	
Leptotrichia buccalis		5.77	0.24		1	
57771103	**Low discrimination**					**57771103**
Actinomyces naeslundii		57.32	0.46	IND 1		
Bifidobacterium breve		21.88	0.24	ESL 12		
Lactobacillus acidophilus		18.03	0.13			
57771112	**Good identification**					**57771112**
					GRA	
Leptotrichia buccalis		92.15	0.58	IND 1	1	
Bifidobacterium breve		7.37	0.59		99	
Bifidobacterium adolescentis		0.33	0.35		99	
57771120	**Low discrimination**					**57771120**
					SPO	
Clostridium paraputrificum		45.82	0.37	IND 1	99	
Clostridium ramosum		31.07	0.41	RAF 7	92	
Bifidobacterium breve		14.53	0.46		1	
57771121	**Low discrimination**					**57771121**
					SPO	
Actinomyces naeslundii		70.31	0.46	IND 1	1	
Clostridium paraputrificum		14.88	0.03	ESL 12	99	
Clostridium ramosum		10.09	0.06		92	
57771122	**Low discrimination**					**57771122**
					SPO	
Bifidobacterium breve		33.77	0.63	IND 1	1	
Clostridium ramosum		28.58	0.51		92	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		15.65	0.47		1		
57771123	**Low discrimination**						**57771123**
					SPO		
Actinomyces naeslundii		33.91	0.46	IND 1	1		
Bifidobacterium breve		23.01	0.29	ESL 12	1		
Clostridium ramosum		19.47	0.17		92		
57771130	**Low discrimination**						**57771130**
					GRA	SPO	
Bifidobacterium breve		88.24	0.46	IND 1	99	1	
Leptotrichia buccalis		7.04	0.07	BGL 91	1	1	
Clostridium paraputrificum		2.81	0.03		99	99	
57771132	**Low discrimination**						**57771132**
					GRA		
Bifidobacterium breve		54.68	0.63	IND 1	99		
Leptotrichia buccalis		42.72	0.41		1		
Parabacteroides distasonis		0.71	0.32		1		
57771140	**Low discrimination**						**57771140**
					SPO		
Bifidobacterium breve		77.66	0.41	IND 1	1		
Clostridium butyricum		15.83	0.14	BGL 91	99		
Actinomyces israelii		4.06	0.25		1		
57771142	**Low discrimination**						**57771142**
					SPO		
Bifidobacterium breve		88.78	0.59	IND 1	1		
Lactobacillus acidophilus		9.04	0.32		1		
Clostridium butyricum		0.70	0.07		99		
57771150	**Good identification**						**57771150**
					GRA		
Bifidobacterium breve		98.73	0.41	IND 1	99		
Leptotrichia buccalis		1.27	-.11	BGL 91	1		
57771152	**Low discrimination**						**57771152**
					GRA		
Bifidobacterium breve		88.26	0.59	IND 1	99		
Leptotrichia buccalis		11.15	0.24		1		
Bifidobacterium adolescentis		0.49	0.19		99		
57771160	**Low discrimination**						**57771160**
					SPO		
Bifidobacterium breve		77.72	0.46	IND 1	1		
Clostridium paraputrificum		18.45	0.18	BGL 91	99		
Clostridium ramosum		1.68	0.06		92		
57771162	**Good identification**						**57771162**
					SPO		
Bifidobacterium breve		92.36	0.63	IND 1	1		
Lactobacillus acidophilus		5.29	0.32		1		
Clostridium paraputrificum		0.93	0.11		99		
57771170	**Good identification**						**57771170**
Bifidobacterium breve		100.0	0.46	IND 1			
57771172	**Very good identification**						**57771172**
					GRA		
Bifidobacterium breve		99.22	0.63	IND 1	99		
Leptotrichia buccalis		0.78	0.07		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
57771320	**Acceptable identification**					**57771320**
					SPO	
Clostridium ramosum		93.79	0.43	IND 1	92	
Prevotella oralis		3.23	0.17		1	
Bacteroides ovatus		1.08	0.06		1	
57771322	**Low discrimination**					**57771322**
					SPO	
Clostridium ramosum		60.98	0.53	IND 1	92	
Bacteroides ovatus		17.45	0.40		1	
Prevotella oralis		9.99	0.39		1	
57771332	**Low discrimination**					**57771332**
Bacteroides ovatus		64.72	0.37	XYL 99		
Parabacteroides distasonis		11.83	0.40	ARA 97		
Bacteroides thetaiotaomicron		9.06	0.22			
57771500	**Good identification**					**57771500**
					SPO	
Clostridium butyricum		99.36	0.48	IND 3	99	
Actinomyces israelii		0.52	0.30	XYL 97	1	
Bifidobacterium adolescentis		0.12	0.13		1	
57771502	**Low discrimination**					**57771502**
					SPO	
Clostridium butyricum		71.81	0.41	IND 3	99	
Bifidobacterium adolescentis		21.83	0.47	XYL 97	1	
Actinomyces israelii		1.44	0.33		1	
57771512	**Low discrimination**					**57771512**
					GRA	
Bifidobacterium adolescentis		54.05	0.49	IND 1	99	
Leptotrichia buccalis		22.63	0.24	XYL 87	1	
Clostridium clostridioforme		11.17	0.34		50	
57771522	**Low discrimination**					**57771522**
					CAT	
Bacteroides thetaiotaomicron		43.35	0.44	XYL 96	89	
Bacteroides ovatus		41.25	0.44	RHA 93	68	
Bacteroides uniformis		8.40	0.29		5	
57771532	**Low discrimination**					**57771532**
Bacteroides ovatus		55.57	0.41	XYL 99		
Bacteroides thetaiotaomicron		34.06	0.37	RHA 95		
Clostridium clostridioforme		5.35	0.34			
57771612	**Good identification**					**57771612**
Clostridium clostridioforme		98.31	0.40	IND 1		
Bacteroides ovatus		0.90	-.06	XYL 91		
Bacteroides thetaiotaomicron		0.78	-.07			
57771632	**Low discrimination**					**57771632**
Clostridium clostridioforme		37.06	0.40	IND 1		
Bacteroides ovatus		33.73	0.29	XYL 91		
Bacteroides thetaiotaomicron		29.21	0.28			
57771712	**Low discrimination**					**57771712**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium clostridioforme		71.39	0.49	IND 1		
Bacteroides ovatus		19.41	0.29	XYL 91		
Bacteroides thetaiotaomicron		8.32	0.22			
57771722	**Low discrimination**					**57771722**
					CAT	
Bacteroides ovatus		56.80	0.66	XYL 99	68	
Bacteroides thetaiotaomicron		41.73	0.64		89	
Bacteroides uniformis		0.84	0.32		5	
57771732	**Low discrimination**					**57771732**
Bacteroides ovatus		68.05	0.63	XYL 99		
Bacteroides thetaiotaomicron		29.17	0.57			
Clostridium clostridioforme		2.53	0.49			
57771762	**Low discrimination**					**57771762**
					CAT	
Bacteroides ovatus		84.62	0.42	XYL 99	68	
Bacteroides thetaiotaomicron		15.07	0.29	SOR 4	89	
Bacteroides uniformis		0.30	-.03		5	
57771772	**Low discrimination**					**57771772**
Bacteroides ovatus		89.85	0.39	XYL 99		
Bacteroides thetaiotaomicron		9.34	0.22	SOR 4		
Clostridium clostridioforme		0.81	0.15			
57773001	**Good identification**					**57773001**
Actinomyces naeslundii		100.0	0.43	IND 1		
57773002	**Low discrimination**					**57773002**
Actinomyces israelii		52.17	0.37	IND 1		
Actinomyces naeslundii		22.22	0.40	XYL 92		
Bifidobacterium breve		18.65	0.24			
57773003	**Good identification**					**57773003**
Actinomyces naeslundii		100.0	0.43	IND 1		
57773021	**Good identification**					**57773021**
Actinomyces naeslundii		100.0	0.43	IND 1		
57773023	**Good identification**					**57773023**
Actinomyces naeslundii		100.0	0.43	IND 1		
57773100	**Low discrimination**					**57773100**
Bifidobacterium breve		38.06	0.41	IND 1		
Actinomyces israelii		28.99	0.45	BGL 91		
Actinomyces naeslundii		13.18	0.48			
57773101	**Good identification**					**57773101**
Actinomyces naeslundii		100.0	0.51	IND 1		
57773102	**Low discrimination**					**57773102**
					SPO	
Bifidobacterium breve		75.16	0.59	IND 1	1	
Actinomyces israelii		8.50	0.48		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		6.47	0.30		92	
57773103	**Low discrimination**					**57773103**
Actinomyces naeslundii		82.04	0.51	IND 1		
Bifidobacterium breve		16.13	0.24	ESL 12		
Actinomyces israelii		1.82	0.14			
57773110	**Low discrimination**					**57773110**
					SPO	
Bifidobacterium breve		81.15	0.41	IND 1	1	
Actinomyces israelii		9.24	0.31	BGL 91	1	
Clostridium tertium		5.98	0.13		99	
57773112	**Low discrimination**					**57773112**
					GRA	
Bifidobacterium breve		82.94	0.59	IND 1	99	
Leptotrichia buccalis		10.47	0.24		1	
Bifidobacterium adolescentis		3.06	0.33		99	
57773120	**Acceptable identification**					**57773120**
					SPO	
Clostridium ramosum		90.07	0.53	IND 1	92	
Bifidobacterium breve		7.43	0.46		1	
Actinomyces naeslundii		1.45	0.48		1	
57773121	**Low discrimination**					**57773121**
					SPO	
Actinomyces naeslundii		68.79	0.51	IND 1	1	
Clostridium ramosum		28.83	0.19	ESL 12	92	
Bifidobacterium breve		2.38	0.12		1	
57773122	**Low discrimination**					**57773122**
					SPO	
Clostridium ramosum		81.62	0.64	IND 1	92	
Bifidobacterium breve		17.02	0.63		1	
Lactobacillus acidophilus		0.42	0.25		1	
57773130	**Low discrimination**					**57773130**
					SPO	
Bifidobacterium breve		79.55	0.46	IND 1	1	
Clostridium tertium		9.89	0.21	BGL 91	99	
Clostridium ramosum		9.74	0.19		92	
57773132	**Good identification**					**57773132**
					SPO	
Bifidobacterium breve		90.86	0.63	IND 1	1	
Clostridium ramosum		4.40	0.30		92	
Clostridium tertium		3.35	0.29		99	
57773140	**Low discrimination**					**57773140**
Bifidobacterium breve		72.46	0.41	IND 1		
Actinomyces israelii		21.47	0.38	BGL 91		
Actinomyces naeslundii		4.43	0.35			
57773141	**Good identification**					**57773141**
Actinomyces naeslundii		100.0	0.38	IND 1		
57773142	**Good identification**					**57773142**
Bifidobacterium breve		94.42	0.59	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		4.15	0.41				
Actinomyces naeslundii		0.57	0.35				
57773150	**Acceptable identification**						**57773150**
					COC		
Bifidobacterium breve		94.00	0.41	IND 1	1		
Actinomyces israelii		4.16	0.24	BGL 91	1		
Blautia producta		1.84	-.03		99		
57773152	**Very good identification**						**57773152**
Bifidobacterium breve		98.78	0.59	IND 1			
Actinomyces israelii		0.65	0.27				
Bifidobacterium adolescentis		0.45	0.18				
57773160	**Low discrimination**						**57773160**
					SPO		
Bifidobacterium breve		85.66	0.46	IND 1	1		
Clostridium ramosum		10.49	0.19	BGL 91	92		
Actinomyces naeslundii		2.94	0.35		1		
57773161	**Low discrimination**						**57773161**
Actinomyces naeslundii		83.62	0.38	IND 1			
Bifidobacterium breve		16.38	0.12	ESL 12			
57773162	**Good identification**						**57773162**
					SPO		
Bifidobacterium breve		94.66	0.63	IND 1	1		
Clostridium ramosum		4.59	0.30		92		
Actinomyces naeslundii		0.32	0.35		1		
57773170	**Good identification**						**57773170**
Bifidobacterium breve		100.0	0.46	IND 1			
57773172	**Very good identification**						**57773172**
Bifidobacterium breve		100.0	0.63	IND 1			
57773320	**Good identification**						**57773320**
					SPO	GRA	
Clostridium ramosum		99.87	0.56	IND 1	92	50	
Bacteroides ovatus		0.07	-.02		1	1	
Bifidobacterium breve		0.06	0.12		1	99	
57773322	**Very good identification**						**57773322**
					SPO	GRA	
Clostridium ramosum		98.11	0.66	IND 1	92	50	
Bacteroides ovatus		1.65	0.32		1	1	
Bifidobacterium breve		0.16	0.29		1	99	
57773402	**Low discrimination**						**57773402**
					SPO		
Actinomyces israelii		79.95	0.36	IND 1	1		
Clostridium butyricum		13.00	0.01	XYL 92	99		
Bifidobacterium adolescentis		7.05	0.12		1		
57773502	**Low discrimination**						**57773502**
					SPO		
Bifidobacterium adolescentis		50.49	0.46	IND 1	1		
Actinomyces israelii		23.13	0.46	XYL 87	1		
Clostridium butyricum		22.55	0.25		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57773512	**Low discrimination**						**57773512**
Bifidobacterium adolescentis		52.17	0.47	IND 1			
Bifidobacterium dentium		41.04	0.27	XYL 87			
Actinomyces israelii		2.92	0.32				
57773522	**Low discrimination**						**57773522**
					SPO	GRA	
Bacteroides ovatus		70.47	0.36	XYL 99	1	1	
Clostridium ramosum		18.75	0.30	RHA 95	92	50	
Bifidobacterium breve		3.91	0.29		1	99	
57773542	**Low discrimination**						**57773542**
Actinomyces israelii		41.85	0.39	IND 1			
Bifidobacterium adolescentis		29.03	0.30	XYL 92			
Bifidobacterium breve		11.75	0.24				
57773722	**Very good identification**						**57773722**
					SPO	CAT	
Bacteroides ovatus		96.04	0.58	XYL 99	1	68	
Bacteroides thetaiotaomicron		2.14	0.29		1	89	
Clostridium ramosum		1.78	0.32		92	1	
57773732	**Good identification**						**57773732**
Bacteroides ovatus		98.50	0.55	XYL 99			
Bacteroides thetaiotaomicron		1.28	0.22				
Clostridium clostridioforme		0.11	0.15				
57774312	**Good identification**						**57774312**
Clostridium clostridioforme		96.24	0.37	IND 1			
Bacteroides ovatus		3.76	0.03	ARA 94			
57774322	**Low discrimination**						**57774322**
					CAT		
Bacteroides ovatus		86.34	0.40	GAL 99	68		
Bacteroides thetaiotaomicron		5.02	0.19	ARA 97	89		
Bacteroides uniformis		3.59	0.14		5		
57774332	**Low discrimination**						**57774332**
Bacteroides ovatus		76.40	0.37	GAL 99			
Clostridium clostridioforme		19.74	0.37	ARA 97			
Bacteroides thetaiotaomicron		2.59	0.12				
57774500	**Good identification**						**57774500**
					SPO		
Clostridium butyricum		99.82	0.40	IND 3	99		
Actinomyces israelii		0.19	0.14	GAL 99	1		
57774512	**Low discrimination**						**57774512**
					SPO		
Clostridium clostridioforme		83.06	0.43	IND 1	66		
Bifidobacterium adolescentis		8.06	0.29	RHA 88	1		
Clostridium butyricum		5.52	0.11		99		
57774522	**Low discrimination**						**57774522**
					CAT		
Bacteroides ovatus		65.27	0.44	GAL 99	68		
Bacteroides thetaiotaomicron		16.63	0.34	RHA 95	89		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		13.29	0.29		5	
57774532	**Low discrimination**					**57774532**
Bacteroides ovatus		67.32	0.41	GAL 99		
Clostridium clostridioforme		21.84	0.43	RHA 95		
Bacteroides thetaiotaomicron		10.00	0.27			
57774602	**Good identification**					**57774602**
Clostridium clostridioforme		98.17	0.37	IND 1		
Bacteroides ovatus		1.83	-.03			
57774612	**Good identification**					**57774612**
Clostridium clostridioforme		99.73	0.49	IND 1		
Bacteroides ovatus		0.27	-.06			
57774632	**Low discrimination**					**57774632**
Clostridium clostridioforme		75.37	0.49	IND 1		
Bacteroides ovatus		20.36	0.29			
Bacteroides thetaiotaomicron		4.27	0.17			
57774702	**Low discrimination**					**57774702**
Clostridium clostridioforme		59.46	0.46	IND 1		
Bacteroides ovatus		32.78	0.32			
Bacteroides thetaiotaomicron		5.84	0.19			
57774712	**Good identification**					**57774712**
Clostridium clostridioforme		91.82	0.58	IND 1		
Bacteroides ovatus		7.41	0.29			
Bacteroides thetaiotaomicron		0.77	0.12			
57774722	**Low discrimination**					**57774722**
Bacteroides ovatus		82.53	0.66	GAL 99		
Bacteroides thetaiotaomicron		14.70	0.53			
Clostridium clostridioforme		1.51	0.46			
57774732	**Low discrimination**					**57774732**
Bacteroides ovatus		81.32	0.63	GAL 99		
Clostridium clostridioforme		10.18	0.58			
Bacteroides thetaiotaomicron		8.45	0.46			
57774762	**Good identification**					**57774762**
Bacteroides ovatus		95.13	0.42	GAL 99		
Bacteroides thetaiotaomicron		4.11	0.19	SOR 4		
Clostridium clostridioforme		0.42	0.12			
57774772	**Acceptable identification**					**57774772**
Bacteroides ovatus		94.74	0.39	GAL 99		
Clostridium clostridioforme		2.88	0.24	SOR 4		
Bacteroides thetaiotaomicron		2.39	0.12			
57775042	**Low discrimination**					**57775042**
					SPO	
Actinomyces israelii		67.86	0.36	IND 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		32.15	0.09		99		
57775100	**Very good identification**						**57775100**
					COC	SPO	
Clostridium butyricum		96.87	0.58	IND 3	1	99	
Actinomyces israelii		1.98	0.50	ARA 90	1	1	
Atopobium parvulum		1.09	0.32		99	1	
57775102	**Low discrimination**						**57775102**
					SPO		
Clostridium butyricum		83.45	0.51	IND 3	99		
Bifidobacterium adolescentis		7.06	0.47	ARA 90	1		
Actinomyces israelii		6.57	0.53		1		
57775110	**Acceptable identification**						**57775110**
					SPO		
Clostridium butyricum		93.56	0.36	IND 3	99		
Actinomyces israelii		5.42	0.36	ARA 90	1		
Bifidobacterium adolescentis		0.72	0.15	MLZ 5	1		
57775112	**Low discrimination**						**57775112**
					GRA	SPO	
Bifidobacterium adolescentis		43.71	0.49	IND 1	99	1	
Clostridium butyricum		22.24	0.29	ARA 87	99	99	
Leptotrichia buccalis		18.30	0.24		1	1	
57775120	**Low discrimination**						**57775120**
					GRA	SPO	
Clostridium butyricum		87.36	0.40	IND 3	99	99	
Bacteroides ovatus		4.02	0.18	ARA 90	1	1	
Bacteroides uniformis		2.30	0.11	NAG 8	1	1	
57775122	**Low discrimination**						**57775122**
Bacteroides ovatus		51.85	0.52	RHA 95			
Bacteroides uniformis		29.69	0.45	ARA 97			
Parabacteroides distasonis		6.76	0.53				
57775132	**Low discrimination**						**57775132**
Bacteroides ovatus		82.64	0.49	RHA 95			
Parabacteroides distasonis		8.70	0.49	ARA 97			
Bacteroides thetaiotaomicron		4.01	0.27				
57775140	**Acceptable identification**						**57775140**
					SPO		
Clostridium butyricum		91.64	0.40	IND 3	99		
Actinomyces israelii		8.36	0.43	ARA 90	1		
57775212	**Good identification**						**57775212**
Clostridium clostridioforme		95.82	0.36	IND 1			
Bacteroides ovatus		4.18	0.03	ARA 94			
57775222	**Low discrimination**						**57775222**
					CAT		
Bacteroides ovatus		79.33	0.40	LAC 99	68		
Bacteroides thetaiotaomicron		9.32	0.24	ARA 97	89		
Bacteroides uniformis		6.67	0.19		5		
57775232	**Low discrimination**						**57775232**
Bacteroides ovatus		75.94	0.37	LAC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		17.59	0.36	ARA 97			
Bacteroides thetaiotaomicron		5.21	0.17				
57775302	**Low discrimination**						**57775302**
					GRA	SPO	
Bacteroides ovatus		69.92	0.40	ARA 97	1	1	
Clostridium butyricum		11.05	0.25	NAG 99	99	99	
Clostridium clostridioforme		7.93	0.34		50	66	
57775312	**Low discrimination**						**57775312**
Bacteroides ovatus		53.87	0.37	ARA 97			
Clostridium clostridioforme		41.76	0.45	NAG 99			
Bacteroides thetaiotaomicron		1.83	0.12				
57775320	**Low discrimination**						**57775320**
					CAT		
Bacteroides ovatus		85.60	0.40	ARA 97	68		
Bacteroides thetaiotaomicron		4.98	0.19	BGL 99	89		
Bacteroides uniformis		3.56	0.14		5		
57775322	**Low discrimination**						**57775322**
					CAT		
Bacteroides ovatus		89.07	0.74	ARA 97	68		
Bacteroides thetaiotaomicron		5.18	0.53		89		
Bacteroides uniformis		3.71	0.48		5		
57775323	**Low discrimination**						**57775323**
					CAT		
Bacteroides ovatus		89.26	0.40	ARA 97	68		
Bacteroides thetaiotaomicron		5.19	0.19	URE 1	89		
Bacteroides uniformis		3.72	0.14		5		
57775330	**Acceptable identification**						**57775330**
Bacteroides ovatus		94.48	0.37	ARA 97			
Bacteroides thetaiotaomicron		3.21	0.12	BGL 99			
Parabacteroides distasonis		1.57	0.23				
57775332	**Good identification**						**57775332**
Bacteroides ovatus		94.43	0.71	ARA 97			
Bacteroides thetaiotaomicron		3.20	0.46				
Parabacteroides distasonis		1.57	0.57				
57775333	**Acceptable identification**						**57775333**
Bacteroides ovatus		94.48	0.37	ARA 97			
Bacteroides thetaiotaomicron		3.21	0.12	URE 1			
Parabacteroides distasonis		1.57	0.23				
57775362	**Good identification**						**57775362**
					CAT		
Bacteroides ovatus		97.17	0.51	ARA 97	68		
Bacteroides thetaiotaomicron		1.37	0.19	SOR 4	89		
Bacteroides uniformis		0.98	0.14		5		
57775372	**Good identification**						**57775372**
Bacteroides ovatus		98.60	0.48	ARA 97			
Bacteroides thetaiotaomicron		0.81	0.12	SOR 4			
Parabacteroides distasonis		0.40	0.23				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57775400	**Good identification**						**57775400**
					SPO		
Clostridium butyricum		98.90	0.51	IND 3	99		
Actinomyces israelii		1.10	0.38	LAC 96	1		
57775402	**Low discrimination**						**57775402**
					SPO		
Clostridium butyricum		89.83	0.44	IND 3	99		
Actinomyces israelii		3.86	0.41	LAC 96	1		
Clostridium clostridioforme		3.39	0.31		66		
57775412	**Low discrimination**						**57775412**
					SPO		
Clostridium clostridioforme		66.36	0.42	IND 1	66		
Clostridium butyricum		20.29	0.22	RHA 88	99		
Bifidobacterium adolescentis		7.19	0.29		1		
57775422	**Low discrimination**						**57775422**
					CAT		
Bacteroides ovatus		48.26	0.44	LAC 99	68		
Bacteroides thetaiotaomicron		24.84	0.39	RHA 95	89		
Bacteroides uniformis		19.86	0.35		5		
57775432	**Low discrimination**						**57775432**
Bacteroides ovatus		62.02	0.41	LAC 99			
Bacteroides thetaiotaomicron		18.62	0.32	RHA 95			
Clostridium clostridioforme		18.03	0.42				
57775500	**Very good identification**						**57775500**
					SPO		
Clostridium butyricum		99.74	0.74	IND 3	99		
Actinomyces israelii		0.19	0.48		1		
Bifidobacterium adolescentis		0.03	0.27		1		
57775501	**Good identification**						**57775501**
					GRA	SPO	
Clostridium butyricum		99.55	0.40	IND 3	99	99	
Mitsuokella multacida		0.27	0.04	URE 1	1	1	
Actinomyces israelii		0.19	0.14		99	1	
57775502	**Low discrimination**						**57775502**
					GRA	SPO	
Clostridium butyricum		88.53	0.67	IND 3	99	99	
Bifidobacterium adolescentis		5.57	0.61		99	1	
Bacteroides ovatus		3.28	0.44		1	1	
57775510	**Good identification**						**57775510**
					SPO		
Clostridium butyricum		97.83	0.52	IND 3	99		
Bifidobacterium adolescentis		0.56	0.29	MLZ 5	1		
Bifidobacterium dentium		0.53	0.10		1		
57775512	**Low discrimination**						**57775512**
					GRA	SPO	
Bifidobacterium adolescentis		40.37	0.63	IND 1	99	1	
Clostridium butyricum		27.63	0.45		99	99	
Bacteroides ovatus		12.95	0.41		1	1	
57775520	**Low discrimination**						**57775520**
					GRA	SPO	
Clostridium butyricum		80.36	0.56	IND 3	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Bacteroides ovatus		13.30	0.44	NAG 8	1	1		
Bacteroides thetaiotaomicron		3.39	0.34		1	1		
57775522	**Low discrimination**							**57775522**
					CAT			
Bacteroides ovatus		67.28	0.78	RHA 95	68			
Bacteroides thetaiotaomicron		17.14	0.68		89			
Bacteroides uniformis		13.70	0.63		5			
57775523	**Low discrimination**							**57775523**
					CAT			
Bacteroides ovatus		67.29	0.44	RHA 95	68			
Bacteroides thetaiotaomicron		17.14	0.34	URE 1	89			
Bacteroides uniformis		13.71	0.29		5			
57775530	**Low discrimination**							**57775530**
					GRA	SPO		
Bacteroides ovatus		58.92	0.41	RHA 95	1	1		
Clostridium butyricum		28.11	0.34	BGL 99	99	99		
Bacteroides thetaiotaomicron		8.76	0.27		1	1		
57775532	**Low discrimination**							**57775532**
Bacteroides ovatus		85.39	0.75	RHA 95				
Bacteroides thetaiotaomicron		12.69	0.61					
Clostridium clostridioforme		0.84	0.51					
57775533	**Low discrimination**							**57775533**
Bacteroides ovatus		85.41	0.41	RHA 95				
Bacteroides thetaiotaomicron		12.69	0.27	URE 1				
Clostridium clostridioforme		0.84	0.17					
57775540	**Good identification**							**57775540**
					COC	SPO		
Clostridium butyricum		97.82	0.56	IND 3	1	99		
Blautia producta		1.06	0.24	SOR 8	99	1		
Actinomyces israelii		0.81	0.41		1	1		
57775542	**Low discrimination**							**57775542**
					SPO			
Clostridium butyricum		87.29	0.49	IND 3	99			
Bifidobacterium adolescentis		7.81	0.46	SOR 8	1			
Actinomyces israelii		2.79	0.44		1			
57775552	**Low discrimination**							**57775552**
					GRA	SPO		
Bifidobacterium adolescentis		57.08	0.47	IND 1	99	1		
Clostridium butyricum		27.48	0.27	SOR 11	99	99		
Bacteroides ovatus		6.17	0.17		1	1		
57775560	**Low discrimination**							**57775560**
					COC	GRA	SPO	
Clostridium butyricum		89.59	0.38	IND 3	1	99	99	
Bacteroides ovatus		7.10	0.20	NAG 8	1	1	1	
Blautia producta		1.24	0.07	SOR 8	99	99	1	
57775562	**Low discrimination**							**57775562**
					GRA	SPO		
Bacteroides ovatus		85.96	0.55	RHA 95	1	1		
Bacteroides thetaiotaomicron		5.31	0.34	SOR 4	1	1		
Clostridium butyricum		4.26	0.31		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
57775572	**Good identification**						**57775572**
					GRA	SPO	
Bacteroides ovatus		95.54	0.52	RHA 95	1	1	
Bacteroides thetaiotaomicron		3.44	0.27	SOR 4	1	1	
Clostridium butyricum		0.37	0.09		99	99	
57775602	**Low discrimination**						**57775602**
Clostridium clostridioforme		50.55	0.46	IND 1			
Bacteroides ovatus		31.10	0.32				
Bacteroides thetaiotaomicron		11.19	0.24				
57775612	**Good identification**						**57775612**
Clostridium clostridioforme		90.07	0.57	IND 1			
Bacteroides ovatus		8.11	0.29				
Bacteroides thetaiotaomicron		1.70	0.17				
57775622	**Low discrimination**						**57775622**
					CAT		
Bacteroides ovatus		71.09	0.66	LAC 99	68		
Bacteroides thetaiotaomicron		25.58	0.58		89		
Bacteroides uniformis		2.13	0.37		5		
57775632	**Low discrimination**						**57775632**
Bacteroides ovatus		75.59	0.63	LAC 99			
Bacteroides thetaiotaomicron		15.87	0.51				
Clostridium clostridioforme		8.48	0.57				
57775662	**Acceptable identification**						**57775662**
					CAT		
Bacteroides ovatus		91.04	0.42	LAC 99	68		
Bacteroides thetaiotaomicron		7.94	0.24	SOR 4	89		
Bacteroides uniformis		0.66	0.03		5		
57775672	**Acceptable identification**						**57775672**
Bacteroides ovatus		92.76	0.39	LAC 99			
Bacteroides thetaiotaomicron		4.72	0.17	SOR 4			
Clostridium clostridioforme		2.52	0.23				
57775700	**Low discrimination**						**57775700**
					GRA	SPO	
Clostridium butyricum		88.44	0.48	IND 3	99	99	
Bacteroides ovatus		7.90	0.32	RHA 3	1	1	
Bacteroides thetaiotaomicron		1.41	0.19		1	1	
57775702	**Low discrimination**						**57775702**
Bacteroides ovatus		77.27	0.66	NAG 99			
Bacteroides thetaiotaomicron		13.76	0.53				
Clostridium clostridioforme		4.25	0.55				
57775712	**Low discrimination**						**57775712**
Bacteroides ovatus		67.31	0.63	NAG 99			
Clostridium clostridioforme		25.29	0.66				
Bacteroides thetaiotaomicron		6.99	0.46				
57775720	**Low discrimination**						**57775720**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		83.00	0.66	BGL 99	68		
Bacteroides thetaiotaomicron		14.78	0.53		89		
Bacteroides uniformis		1.23	0.32		5		
57775722	**Low discrimination**						**57775722**
					CAT		
Bacteroides ovatus		83.76	1.00		68		
Bacteroides thetaiotaomicron		14.92	0.87		89		
Bacteroides uniformis		1.24	0.66		5		
57775723	**Low discrimination**						**57775723**
					CAT		
Bacteroides ovatus		83.76	0.66	URE 1	68		
Bacteroides thetaiotaomicron		14.92	0.53		89		
Bacteroides uniformis		1.24	0.32		5		
57775730	**Good identification**						**57775730**
Bacteroides ovatus		90.15	0.63	BGL 99			
Bacteroides thetaiotaomicron		9.37	0.46				
Clostridium clostridioforme		0.34	0.32				
57775732	**Good identification**						**57775732**
Bacteroides ovatus		90.23	0.97				
Bacteroides thetaiotaomicron		9.38	0.80				
Clostridium clostridioforme		0.34	0.66				
57775733	**Good identification**						**57775733**
Bacteroides ovatus		90.23	0.63	URE 1			
Bacteroides thetaiotaomicron		9.38	0.46				
Clostridium clostridioforme		0.34	0.32				
57775742	**Low discrimination**						**57775742**
					GRA	SPO	
Bacteroides ovatus		85.08	0.42	NAG 99	1	1	
Clostridium butyricum		7.81	0.23	SOR 4	99	99	
Bacteroides thetaiotaomicron		3.67	0.19		1	1	
57775752	**Low discrimination**						**57775752**
					COC	GRA	
Bacteroides ovatus		85.32	0.39	NAG 99	1	1	
Clostridium clostridioforme		7.77	0.32	SOR 4	1	50	
Blautia producta		3.54	0.15		99	99	
57775760	**Acceptable identification**						**57775760**
					GRA	SPO	
Bacteroides ovatus		92.04	0.42	SOR 4	1	1	
Bacteroides thetaiotaomicron		3.97	0.19	BGL 99	1	1	
Clostridium butyricum		1.89	0.12		99	99	
57775762	**Very good identification**						**57775762**
					CAT		
Bacteroides ovatus		95.50	0.76	SOR 4	68		
Bacteroides thetaiotaomicron		4.12	0.53		89		
Bacteroides uniformis		0.34	0.32		5		
57775763	**Good identification**						**57775763**
					CAT		
Bacteroides ovatus		95.53	0.42	SOR 4	68		
Bacteroides thetaiotaomicron		4.13	0.19	URE 1	89		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides uniformis		0.34	-.03		5		
57775770	**Acceptable identification**						**57775770**
					COC	GRA	
Bacteroides ovatus		93.31	0.39	SOR 4	1	1	
Blautia producta		3.87	0.15	BGL 99	99	99	
Bacteroides thetaiotaomicron		2.35	0.12		1	1	
57775772	**Very good identification**						**57775772**
Bacteroides ovatus		97.44	0.73	SOR 4			
Bacteroides thetaiotaomicron		2.45	0.46				
Clostridium clostridioforme		0.09	0.32				
57775773	**Good identification**						**57775773**
Bacteroides ovatus		97.54	0.39	SOR 4			
Bacteroides thetaiotaomicron		2.46	0.12	URE 1			
57776522	**Good identification**						**57776522**
					GRA	SPO	
Bacteroides ovatus		96.85	0.36	GAL 99	1	1	
Bifidobacterium dentium		1.04	-.03	RHA 95	99	1	
Clostridium butyricum		0.77	-.02		99	99	
57776722	**Very good identification**						**57776722**
Bacteroides ovatus		99.36	0.58	GAL 99			
Bacteroides thetaiotaomicron		0.54	0.19				
Clostridium clostridioforme		0.06	0.12				
57776732	**Good identification**						**57776732**
Bacteroides ovatus		99.23	0.55	GAL 99			
Clostridium clostridioforme		0.38	0.24				
Bacteroides thetaiotaomicron		0.31	0.12				
57777000	**Low discrimination**						**57777000**
					SPO		
Actinomyces israelii		86.19	0.53	IND 1	1		
Clostridium butyricum		13.81	0.18		99		
57777002	**Good identification**						**57777002**
					SPO		
Actinomyces israelii		95.44	0.56	IND 1	1		
Clostridium butyricum		3.97	0.11		99		
Bifidobacterium adolescentis		0.60	0.12		1		
57777010	**Acceptable identification**						**57777010**
					SPO		
Actinomyces israelii		94.66	0.38	IND 1	1		
Clostridium butyricum		5.34	-.04	MLZ 13	99		
57777012	**Acceptable identification**						**57777012**
Actinomyces israelii		91.60	0.41	IND 1			
Bifidobacterium adolescentis		4.70	0.13	MLZ 13			
Bifidobacterium dentium		3.70	-.07				
57777022	**Low discrimination**						**57777022**
					GRA	SPO	
Actinomyces israelii		64.97	0.35	IND 1	99	1	
Bacteroides ovatus		31.36	0.10	NAG 6	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		3.68	-.07		99	99	
57777040	**Good identification**						**57777040**
					SPO		
Actinomyces israelii		96.54	0.46	IND 1	1		
Clostridium butyricum		3.46	0.00		99		
57777042	**Good identification**						**57777042**
					SPO		
Actinomyces israelii		99.08	0.49	IND 1	1		
Clostridium butyricum		0.92	-.07		99		
57777100	**Low discrimination**						**57777100**
					SPO		
Actinomyces israelii		50.75	0.63	IND 1	1		
Clostridium butyricum		48.80	0.42		99		
Clostridium tertium		0.33	0.11		99		
57777102	**Low discrimination**						**57777102**
					SPO		
Actinomyces israelii		69.71	0.66	IND 1	1		
Clostridium butyricum		17.37	0.35		99		
Bifidobacterium adolescentis		10.83	0.46		1		
57777110	**Low discrimination**						**57777110**
					SPO		
Actinomyces israelii		62.55	0.49	IND 1	1		
Clostridium butyricum		21.18	0.20	MLZ 13	99		
Bifidobacterium dentium		9.27	0.10		1		
57777112	**Low discrimination**						**57777112**
Bifidobacterium adolescentis		35.23	0.47	IND 1			
Actinomyces israelii		27.73	0.52	ARA 87			
Bifidobacterium dentium		27.71	0.27				
57777122	**Low discrimination**						**57777122**
					SPO	GRA	
Bacteroides ovatus		77.05	0.44	RHA 95	1	1	
Clostridium ramosum		6.69	0.30	ARA 97	92	50	
Actinomyces israelii		6.45	0.45		1	99	
57777132	**Low discrimination**						**57777132**
					GRA	SPO	
Bacteroides ovatus		58.33	0.41	RHA 95	1	1	
Bifidobacterium dentium		30.46	0.32	ARA 97	99	1	
Clostridium tertium		8.00	0.32		99	99	
57777140	**Low discrimination**						**57777140**
					COC	SPO	
Actinomyces israelii		71.22	0.56	IND 1	1	1	
Clostridium butyricum		15.31	0.23		1	99	
Blautia producta		13.48	0.24		99	1	
57777142	**Low discrimination**						**57777142**
					SPO		
Actinomyces israelii		87.79	0.59	IND 1	1		
Clostridium butyricum		4.89	0.16		99		
Bifidobacterium adolescentis		4.33	0.30		1		
57777152	**Low discrimination**						**57777152**
					COC		
Actinomyces israelii		53.63	0.45	IND 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		21.65	0.32	MLZ 13	1		
Blautia producta		15.09	0.15		99		
57777160	**Low discrimination**						**57777160**
					COC	SPO	
Actinomyces israelii		58.66	0.35	IND 1	1	1	
Blautia producta		19.32	0.07	NAG 6	99	1	
Clostridium butyricum		17.18	0.05		1	99	
57777202	**Acceptable identification**						**57777202**
					GRA		
Actinomyces israelii		92.72	0.36	IND 1	99		
Bacteroides ovatus		7.28	-.02	RHA 7	1		
57777300	**Low discrimination**						**57777300**
					GRA	SPO	
Actinomyces israelii		70.20	0.44	IND 1	99	1	
Clostridium butyricum		27.73	0.16	RHA 7	99	99	
Bacteroides ovatus		2.07	-.02		1	1	
57777322	**Very good identification**						**57777322**
					SPO	CAT	
Bacteroides ovatus		99.01	0.66	ARA 97	1	68	
Clostridium ramosum		0.60	0.32		92	1	
Bacteroides thetaiotaomicron		0.18	0.19		1	89	
57777332	**Very good identification**						**57777332**
Bacteroides ovatus		99.79	0.63	ARA 97			
Bacteroides thetaiotaomicron		0.10	0.12				
Parabacteroides distasonis		0.05	0.23				
57777342	**Low discrimination**						**57777342**
					COC	GRA	
Actinomyces israelii		71.71	0.40	IND 1	1	99	
Bacteroides ovatus		14.94	0.08	RHA 7	1	1	
Blautia producta		13.35	0.07		99	99	
57777362	**Good identification**						**57777362**
					SPO	GRA	
Bacteroides ovatus		99.55	0.42	ARA 97	1	1	
Actinomyces israelii		0.31	0.19	SOR 4	1	99	
Clostridium ramosum		0.15	-.03		92	50	
57777372	**Good identification**						**57777372**
					COC	GRA	
Bacteroides ovatus		99.55	0.39	ARA 97	1	1	
Blautia producta		0.45	-.01	SOR 4	99	99	
57777402	**Low discrimination**						**57777402**
					SPO		
Actinomyces israelii		65.45	0.54	IND 1	1		
Clostridium butyricum		29.91	0.27		99		
Bifidobacterium adolescentis		3.36	0.26		1		
57777410	**Low discrimination**						**57777410**
					SPO		
Actinomyces israelii		41.35	0.37	IND 1	1		
Bifidobacterium dentium		29.96	0.10	MLZ 13	1		
Clostridium butyricum		25.67	0.12		99		
57777422	**Low discrimination**						**57777422**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		88.98	0.36	LAC 99	1	1	
Actinomyces israelii		4.67	0.34	RHA 95	99	1	
Clostridium butyricum		2.90	0.09		99	99	
57777440	**Low discrimination**						**57777440**
					COC	SPO	
Actinomyces israelii		43.12	0.44	IND 1	1	1	
Blautia producta		39.89	0.24		99	1	
Clostridium butyricum		16.99	0.16		1	99	
57777442	**Low discrimination**						**57777442**
					COC	SPO	
Actinomyces israelii		84.25	0.47	IND 1	1	1	
Clostridium butyricum		8.61	0.09		1	99	
Blautia producta		5.77	0.07		99	1	
57777500	**Low discrimination**						**57777500**
					COC	SPO	
Clostridium butyricum		89.84	0.58	IND 3	1	99	
Actinomyces israelii		8.50	0.61	MAN 10	1	1	
Blautia producta		0.76	0.24		99	1	
57777502	**Low discrimination**						**57777502**
					SPO		
Clostridium butyricum		49.71	0.51	IND 3	99		
Bifidobacterium adolescentis		23.03	0.60	MAN 10	1		
Actinomyces israelii		18.13	0.64		1		
57777510	**Low discrimination**						**57777510**
					COC	SPO	
Bifidobacterium dentium		72.42	0.44	IND 1	1	1	
Clostridium butyricum		15.04	0.36	BGL 91	1	99	
Blautia producta		7.29	0.32		99	1	
57777512	**Low discrimination**						**57777512**
					GRA		
Bifidobacterium dentium		89.69	0.62	IND 1	99		
Bifidobacterium adolescentis		7.71	0.61		99		
Bacteroides ovatus		1.01	0.33		1		
57777520	**Low discrimination**						**57777520**
					GRA	SPO	
Clostridium butyricum		60.58	0.40	IND 3	99	99	
Bacteroides ovatus		30.08	0.36	MAN 10	1	1	
Actinomyces israelii		4.21	0.41	NAG 8	99	1	
57777522	**Very good identification**						**57777522**
					GRA	SPO	
Bacteroides ovatus		96.60	0.70	RHA 95	1	1	
Bifidobacterium dentium		1.04	0.32		99	1	
Clostridium butyricum		0.76	0.33		99	99	
57777523	**Good identification**						**57777523**
					GRA	SPO	
Bacteroides ovatus		96.85	0.36	RHA 95	1	1	
Bifidobacterium dentium		1.04	-.03	URE 1	99	1	
Clostridium butyricum		0.77	-.02		99	99	
57777530	**Acceptable identification**						**57777530**
					GRA	SPO	
Bifidobacterium dentium		92.31	0.49	IND 1	99	1	
Bacteroides ovatus		5.90	0.33	BGL 91	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.94	0.18		99	99	
57777532	**Low discrimination**						**57777532**
					GRA	CAT	
Bifidobacterium dentium		61.36	0.66	IND 1	99	1	
Bacteroides ovatus		38.38	0.67		1	68	
Bacteroides thetaiotaomicron		0.17	0.27		1	89	
57777540	**Low discrimination**						**57777540**
					COC	SPO	
Blautia producta		86.71	0.58	IND 1	99	1	
Clostridium butyricum		8.95	0.40		1	99	
Actinomyces israelii		3.79	0.54		1	1	
57777542	**Low discrimination**						**57777542**
					COC	SPO	
Blautia producta		45.10	0.41	IND 1	99	1	
Actinomyces israelii		26.59	0.57	BGL 10	1	1	
Clostridium butyricum		16.30	0.33		1	99	
57777550	**Very good identification**						**57777550**
					COC	SPO	
Blautia producta		99.42	0.66	IND 1	99	1	
Actinomyces israelii		0.22	0.40		1	1	
Clostridium butyricum		0.18	0.18		1	99	
57777552	**Low discrimination**						**57777552**
					COC		
Blautia producta		81.31	0.50	IND 1	99		
Bifidobacterium adolescentis		7.89	0.46	BGL 10	1		
Bifidobacterium dentium		7.50	0.27		1		
57777560	**Low discrimination**						**57777560**
					COC	SPO	
Blautia producta		87.80	0.41	IND 1	99	1	
Clostridium butyricum		7.10	0.22	NAG 10	1	99	
Actinomyces israelii		2.20	0.34		1	1	
57777562	**Acceptable identification**						**57777562**
					COC	GRA	
Bacteroides ovatus		90.75	0.46	RHA 95	1	1	
Blautia producta		5.30	0.25	SOR 4	99	99	
Actinomyces israelii		1.79	0.37		1	99	
57777570	**Good identification**						**57777570**
					COC	GRA	
Blautia producta		97.59	0.50	IND 1	99	99	
Bifidobacterium dentium		1.58	0.15	NAG 10	1	99	
Bacteroides ovatus		0.42	0.09		1	1	
57777572	**Low discrimination**						**57777572**
					COC	GRA	
Bacteroides ovatus		60.30	0.43	RHA 95	1	1	
Bifidobacterium dentium		23.37	0.32	SOR 4	1	99	
Blautia producta		15.84	0.33		99	99	
57777622	**Very good identification**						**57777622**
					CAT		
Bacteroides ovatus		98.76	0.58	LAC 99	68		
Bacteroides thetaiotaomicron		1.08	0.24		89		
Bacteroides uniformis		0.09	0.03		5		
57777632	**Good identification**						**57777632**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		98.95	0.55	LAC 99			
Bacteroides thetaiotaomicron		0.63	0.17				
Clostridium clostridioforme		0.34	0.23				
57777702	**Very good identification**						**57777702**
					GRA	SPO	
Bacteroides ovatus		96.14	0.58	NAG 99	1	1	
Clostridium butyricum		1.41	0.25		99	99	
Actinomyces israelii		1.25	0.45		99	1	
57777712	**Acceptable identification**						**57777712**
					GRA		
Bacteroides ovatus		92.95	0.55	NAG 99	1		
Bifidobacterium dentium		4.40	0.27		99		
Clostridium clostridioforme		1.06	0.32		50		
57777720	**Very good identification**						**57777720**
					GRA	SPO	
Bacteroides ovatus		98.64	0.58	BGL 99	1	1	
Bacteroides thetaiotaomicron		0.53	0.19		1	1	
Clostridium butyricum		0.32	0.14		99	99	
57777722	**Excellent identification**						**57777722**
					CAT		
Bacteroides ovatus		99.41	0.92		68		
Bacteroides thetaiotaomicron		0.54	0.53		89		
Bacteroides uniformis		0.05	0.32		5		
57777723	**Very good identification**						**57777723**
					CAT		
Bacteroides ovatus		99.42	0.58	URE 1	68		
Bacteroides thetaiotaomicron		0.54	0.19		89		
Bacteroides uniformis		0.05	-.03		5		
57777730	**Good identification**						**57777730**
					COC	GRA	
Bacteroides ovatus		98.22	0.55	BGL 99	1	1	
Bifidobacterium dentium		0.82	0.15		1	99	
Blautia producta		0.51	0.15		99	99	
57777732	**Very good identification**						**57777732**
					GRA	CAT	
Bacteroides ovatus		99.59	0.89		1	68	
Bacteroides thetaiotaomicron		0.31	0.46		1	89	
Bifidobacterium dentium		0.09	0.32		99	1	
57777733	**Good identification**						**57777733**
					GRA	CAT	
Bacteroides ovatus		99.60	0.55	URE 1	1	68	
Bacteroides thetaiotaomicron		0.31	0.12		1	89	
Bifidobacterium dentium		0.09	-.03		99	1	
57777740	**Very good identification**						**57777740**
					COC	GRA	
Blautia producta		98.21	0.58	IND 1	99	99	
Mitsuokella multacida		1.12	0.32		1	1	
Actinomyces israelii		0.32	0.35		1	99	
57777742	**Low discrimination**						**57777742**
					COC	GRA	
Blautia producta		69.70	0.41	IND 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		25.46	0.34	BGL 10	1	1	
Actinomyces israelii		3.09	0.38		1	99	
57777750	**Very good identification**						**57777750**
					COC	GRA	
Blautia producta		99.85	0.66	IND 1	99	99	
Mitsuokella multacida		0.13	0.24		1	1	
Actinomyces israelii		0.02	0.21		1	99	
57777752	**Acceptable identification**						**57777752**
					COC	GRA	
Blautia producta		92.03	0.50	IND 1	99	99	
Bacteroides ovatus		7.47	0.31	BGL 10	1	1	
Actinomyces israelii		0.20	0.24		1	99	
57777760	**Low discrimination**						**57777760**
					COC	GRA	
Blautia producta		71.67	0.41	IND 1	99	99	
Bacteroides ovatus		26.18	0.34	NAG 10	1	1	
Mitsuokella multacida		1.83	0.21		1	1	
57777762	**Very good identification**						**57777762**
					COC	GRA	
Bacteroides ovatus		99.54	0.68	SOR 4	1	1	
Blautia producta		0.31	0.25		99	99	
Bacteroides thetaiotaomicron		0.13	0.19		1	1	
57777770	**Acceptable identification**						**57777770**
					COC	GRA	
Blautia producta		92.25	0.50	IND 1	99	99	
Bacteroides ovatus		7.49	0.31	NAG 10	1	1	
Mitsuokella multacida		0.26	0.13		1	1	
57777772	**Very good identification**						**57777772**
					COC	GRA	
Bacteroides ovatus		98.54	0.65	SOR 4	1	1	
Blautia producta		1.36	0.33		99	99	
Bacteroides thetaiotaomicron		0.08	0.12		1	1	
60000000	**Low discrimination**						**60000000**
					COC	GRA	
Veillonella parvula		49.91	0.66	ESL 1	99	1	
Campylobacter gracilis		36.67	0.66		1	1	
Eggerthella lenta		6.05	0.51		1	99	
60000001	**Good identification**						**60000001**
					COC		
Campylobacter ureolyticus		93.19	0.60	ESL 1	1		
Veillonella parvula		3.13	0.32		99		
Campylobacter gracilis		2.30	0.32		1		
60000002	**Low discrimination**						**60000002**
					COC	GRA	
Campylobacter gracilis		68.46	0.50	ESL 1	1	1	
Atopobium minutum		8.75	0.53	BGL 10	1	99	
Veillonella parvula		8.47	0.32		99	1	
60000020	**Low discrimination**						**60000020**
					COC	GRA	
Campylobacter gracilis		88.94	0.56	ESL 1	1	1	
Veillonella parvula		4.89	0.32		99	1	
Atopobium minutum		2.53	0.47		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
60000022	**Good identification**						**60000022**
					GRA		
Pseudoflavonifractor capillosus		94.52	0.66	NIT 1	1		
Campylobacter gracilis		4.62	0.39		1		
Atopobium minutum		0.79	0.44		99		
60001002	**Low discrimination**						**60001002**
					GRA		
Atopobium minutum		70.26	0.44	NIT 1	99		
Campylobacter gracilis		16.66	0.15		1		
Pseudoflavonifractor capillosus		9.47	0.15		1		
60001022	**Good identification**						**60001022**
					GRA		
Pseudoflavonifractor capillosus		97.14	0.50	NIT 1	1		
Atopobium minutum		2.43	0.36	GAL 10	99		
Campylobacter gracilis		0.43	0.05		1		
60010000	**Low discrimination**						**60010000**
					COC	GRA	
Atopobium minutum		36.15	0.55	NIT 1	1	99	
Veillonella parvula		25.29	0.32		99	1	
Campylobacter gracilis		18.58	0.32		1	1	
60010002	**Low discrimination**						**60010002**
					GRA	SPO	
Atopobium minutum		77.92	0.52	NIT 1	99	1	
Clostridium sporogenes		10.79	0.27		99	99	
Campylobacter gracilis		6.67	0.15		1	1	
60010020	**Low discrimination**						**60010020**
					GRA		
Atopobium minutum		65.31	0.47	NIT 1	99		
Campylobacter gracilis		25.17	0.21		1		
Pseudoflavonifractor capillosus		6.36	0.15		1		
60010022	**Acceptable identification**						**60010022**
					GRA		
Pseudoflavonifractor capillosus		93.14	0.50	NIT 1	1		
Atopobium minutum		6.45	0.44	FRU 10	99		
Campylobacter gracilis		0.41	0.05		1		
60011000	**Good identification**						**60011000**
					COC	GRA	
Atopobium minutum		95.68	0.47	NIT 1	1	99	
Veillonella parvula		2.03	-.03		99	1	
Campylobacter gracilis		1.49	-.03		1	1	
60011002	**Good identification**						**60011002**
					GRA		
Atopobium minutum		98.40	0.44	NIT 1	99		
Pseudoflavonifractor capillosus		1.60	-.01		1		
60011020	**Acceptable identification**						**60011020**
					GRA	CAT	
Atopobium minutum		90.84	0.39	NIT 1	99	1	
Propionibacterium acnes		5.15	0.10		99	95	
Pseudoflavonifractor capillosus		2.95	-.01		1	1	
60100000	**Low discrimination**						**60100000**
					COC	GRA	
Clostridium sporogenes		51.91	0.53	NIT 1	1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		19.58	0.48		1	99	
Veillonella parvula		12.46	0.32		99	1	
60100002	**Acceptable identification**						**60100002**
					GRA	SPO	
Clostridium sporogenes		91.80	0.45	NIT 1	99	99	
Campylobacter gracilis		5.40	0.15		1	1	
Clostridium botulinum biovar A		1.05	0.14		99	99	
60100040	**Low discrimination**						**60100040**
					COC	GRA	
Clostridium botulinum biovar A		86.74	0.39	NIT 1	1	99	
Clostridium sporogenes		7.78	0.19	GLU 90	1	99	
Veillonella parvula		1.87	-.03		99	1	
60110000	**Low discrimination**						**60110000**
					SPO		
Clostridium sporogenes		83.83	0.40	NIT 1	99		
Clostridium botulinum biovar A		11.44	0.28		99		
Atopobium minutum		1.65	0.21		1		
60120000	**Low discrimination**						**60120000**
					COC	GRA	
Clostridium sporogenes		77.11	0.45	NIT 1	1	99	
Clostridium botulinum biovar A		21.82	0.38		1	99	
Veillonella parvula		0.56	-.03		99	1	
60120002	**Good identification**						**60120002**
Clostridium sporogenes		99.15	0.37	NIT 1			
Clostridium botulinum biovar A		0.85	0.04				
61000000	**Low discrimination**						**61000000**
					SPO		
Clostridium botulinum biovar A		47.61	0.66	NIT 1	99		
Clostridium sporogenes		35.01	0.61		99		
Atopobium minutum		7.26	0.60		1		
61000001	**Low discrimination**						**61000001**
					GRA	SPO	
Clostridium novyi biovar A		48.59	0.53	NIT 1	99	99	
Campylobacter ureolyticus		23.39	0.26		1	1	
Clostridium botulinum biovar A		13.60	0.32		99	99	
61000002	**Low discrimination**						**61000002**
					SPO		
Clostridium sporogenes		66.35	0.53	NIT 1	99		
Atopobium minutum		27.52	0.57		1		
Clostridium botulinum biovar A		2.73	0.32		99		
61000003	**Low discrimination**						**61000003**
					SPO		
Clostridium novyi biovar A		63.39	0.40	NIT 1	99		
Clostridium sporogenes		24.64	0.19		99		
Atopobium minutum		10.22	0.22		1		
61000010	**Low discrimination**						**61000010**
Clostridium novyi biovar A		62.64	0.53	NIT 1			
Clostridium botulinum biovar A		17.53	0.32				
Clostridium sporogenes		12.89	0.27				
61000011	**Good identification**						**61000011**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium novyi biovar A		99.52	0.53	NIT 1	99	99	
Campylobacter ureolyticus		0.48	-.08		1	1	
61000012	**Low discrimination**						**61000012**
					SPO		
Clostridium novyi biovar A		64.52	0.40	NIT 1	99		
Clostridium sporogenes		25.08	0.19		99		
Atopobium minutum		10.40	0.22		1		
61000013	**Good identification**						**61000013**
Clostridium novyi biovar A		100.0	0.40	NIT 1			
61000020	**Low discrimination**						**61000020**
					GRA	SPO	
Atopobium minutum		54.94	0.51	NIT 1	99	1	
Campylobacter gracilis		11.48	0.21		1	1	
Clostridium botulinum biovar A		10.92	0.32		99	99	
61000021	**Acceptable identification**						**61000021**
					GRA	SPO	
Clostridium novyi biovar A		91.89	0.43	NIT 1	99	99	
Atopobium minutum		5.23	0.17		99	1	
Campylobacter ureolyticus		1.79	-.08		1	1	
61000022	**Low discrimination**						**61000022**
					GRA	SPO	
Atopobium minutum		53.56	0.48	NIT 1	99	1	
Pseudoflavonifractor capillosus		38.15	0.32		1	1	
Clostridium sporogenes		3.91	0.19		99	99	
61000030	**Acceptable identification**						**61000030**
					GRA	SPO	
Clostridium novyi biovar A		93.56	0.43	NIT 1	99	99	
Atopobium minutum		5.32	0.17		99	1	
Campylobacter gracilis		1.11	-.13		1	1	
61000031	**Good identification**						**61000031**
Clostridium novyi biovar A		100.0	0.43	NIT 1			
61000040	**Low discrimination**						**61000040**
Clostridium botulinum biovar A		87.61	0.57	NIT 1			
Clostridium novyi biovar A		7.06	0.50				
Clostridium sporogenes		2.18	0.27				
61000041	**Low discrimination**						**61000041**
					GRA	SPO	
Clostridium novyi biovar A		88.29	0.50	NIT 1	99	99	
Clostridium botulinum biovar A		11.07	0.23		99	99	
Campylobacter ureolyticus		0.64	-.08		1	1	
61000042	**Low discrimination**						**61000042**
Clostridium novyi biovar A		39.44	0.37	NIT 1			
Clostridium botulinum biovar A		28.02	0.23				
Clostridium sporogenes		23.00	0.19				
61000043	**Good identification**						**61000043**
Clostridium novyi biovar A		100.0	0.37	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
61000050	**Low discrimination**						**61000050**
Clostridium novyi biovar A		88.86	0.50	NIT 1			
Clostridium botulinum biovar A		11.14	0.23				
61000051	**Good identification**						**61000051**
Clostridium novyi biovar A		100.0	0.50	NIT 1			
61000052	**Good identification**						**61000052**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61000053	**Good identification**						**61000053**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61000060	**Low discrimination**						**61000060**
					SPO		
Clostridium novyi biovar A		60.89	0.40	NIT 1	99		
Clostridium botulinum biovar A		30.54	0.23		99		
Atopobium minutum		5.20	0.17		1		
61000061	**Good identification**						**61000061**
Clostridium novyi biovar A		100.0	0.40	NIT 1			
61000070	**Good identification**						**61000070**
Clostridium novyi biovar A		100.0	0.40	NIT 1			
61000071	**Good identification**						**61000071**
Clostridium novyi biovar A		100.0	0.40	NIT 1			
61001000	**Low discrimination**						**61001000**
					SPO		
Atopobium minutum		58.20	0.51	NIT 1	1		
Eubacterium tortuosum		15.28	0.30		1		
Clostridium botulinum biovar A		11.56	0.32		99		
61001001	**Low discrimination**						**61001001**
					GRA	SPO	
Clostridium novyi biovar A		82.96	0.37	NIT 1	99	99	
Atopobium minutum		10.62	0.17	GAL 10	99	1	
Campylobacter ureolyticus		3.63	-.08		1	1	
61001002	**Low discrimination**						**61001002**
					SPO		
Atopobium minutum		85.95	0.48	NIT 1	1		
Clostridium sporogenes		6.28	0.19		99		
Eubacterium tortuosum		5.97	0.17		1		
61001010	**Low discrimination**						**61001010**
					SPO		
Clostridium novyi biovar A		86.08	0.37	NIT 1	99		
Atopobium minutum		11.02	0.17	GAL 10	1		
Eubacterium tortuosum		2.89	-.04		1		
61001011	**Good identification**						**61001011**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61001020	**Low discrimination**						**61001020**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Atopobium minutum		80.96	0.43	NIT 1	1	
Eubacterium tortuosum		11.25	0.17		1	
Clostridium novyi biovar A		4.79	0.26		99	
61001022	**Low discrimination**					**61001022**
					GRA	
Atopobium minutum		77.53	0.40	NIT 1	99	
Pseudoflavonifractor capillosus		18.41	0.15		1	
Eubacterium tortuosum		2.85	0.04		99	
61004040	**Good identification**					**61004040**
Clostridium glycolicum		97.32	0.49	NIT 1		
Clostridium botulinum biovar A		2.48	0.23	FRU 90		
Clostridium novyi biovar A		0.20	0.16			
61010000	**Low discrimination**					**61010000**
					SPO	
Eubacterium limosum		33.51	0.49	NIT 1	1	
Atopobium minutum		23.79	0.59	MAN 86	1	
Clostridium sporogenes		21.93	0.48		99	
61010001	**Low discrimination**					**61010001**
					SPO	
Clostridium novyi biovar A		39.73	0.37	NIT 1	99	
Eubacterium limosum		19.84	0.15	FRU 10	1	
Atopobium minutum		14.09	0.25		1	
61010002	**Low discrimination**					**61010002**
					SPO	
Atopobium minutum		65.31	0.56	NIT 1	1	
Clostridium sporogenes		30.10	0.40		99	
Pseudoramibacter alactolyticus		1.85	0.15		1	
61010010	**Low discrimination**					**61010010**
					SPO	
Clostridium difficile		89.55	0.45	NIT 1	99	
Clostridium novyi biovar A		4.35	0.37	MAN 90	99	
Eubacterium limosum		2.17	0.15		1	
61010011	**Low discrimination**					**61010011**
Clostridium novyi biovar A		82.79	0.37	NIT 1		
Clostridium difficile		17.21	0.10	FRU 10		
61010020	**Low discrimination**					**61010020**
					CAT	
Propionibacterium acnes		47.56	0.44	IND 87	95	
Atopobium minutum		43.31	0.51	ESL 1	1	
Pseudoramibacter alactolyticus		3.90	0.19		1	
61010022	**Low discrimination**					**61010022**
					GRA CAT	
Atopobium minutum		87.95	0.48	NIT 1	99 1	
Pseudoflavonifractor capillosus		7.54	0.15		1 1	
Propionibacterium acnes		1.46	0.10		99 95	
61010040	**Low discrimination**					**61010040**
					SPO	
Clostridium glycolicum		61.51	0.43	NIT 1	99	
Clostridium botulinum biovar A		20.88	0.36	XYL 95	99	
Pseudoramibacter alactolyticus		6.56	0.21		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
61010100	**Low discrimination**						**61010100**
					GRA		
Fusobacterium mortiferum		31.70	0.35	NIT 1	1		
Eubacterium limosum		23.07	0.15		99		
Atopobium minutum		16.38	0.25		99		
61011000	**Low discrimination**						**61011000**
					GRA		
Atopobium minutum		86.11	0.51	NIT 1	99		
Eubacterium limosum		3.68	0.15		99		
Fusobacterium mortiferum		3.25	0.32		1		
61011002	**Good identification**						**61011002**
					GRA	SPO	
Atopobium minutum		98.02	0.48	NIT 1	99	1	
Clostridium sporogenes		1.37	0.06		99	99	
Fusobacterium mortiferum		0.62	0.16		1	1	
61011020	**Low discrimination**						**61011020**
					SPO	CAT	
Propionibacterium acnes		76.59	0.44	IND 87	1	95	
Atopobium minutum		23.25	0.43	ESL 1	1	1	
Clostridium novyi biovar A		0.17	0.10		99	1	
61011022	**Acceptable identification**						**61011022**
					GRA	CAT	
Atopobium minutum		92.72	0.40	NIT 1	99	1	
Propionibacterium acnes		4.63	0.10		99	95	
Pseudoflavonifractor capillosus		2.65	-.01		1	1	
61011100	**Low discrimination**						**61011100**
					GRA		
Fusobacterium mortiferum		85.84	0.38	NIT 1	1		
Atopobium minutum		9.86	0.17		99		
Actinomyces meyeri		4.31	0.07		99		
61012000	**Low discrimination**						**61012000**
					SPO		
Eubacterium limosum		87.68	0.63	NIT 1	1		
Clostridium difficile		10.95	0.49		99		
Pseudoramibacter alactolyticus		0.93	0.27		1		
61012010	**Very good identification**						**61012010**
					SPO		
Clostridium difficile		98.35	0.61	NIT 1	99		
Eubacterium limosum		1.63	0.29		1		
Pseudoramibacter alactolyticus		0.02	-.07		1		
61012020	**Low discrimination**						**61012020**
					CAT		
Propionibacterium acnes		63.60	0.40	IND 87	95		
Eubacterium limosum		26.99	0.29	ESL 1	1		
Pseudoramibacter alactolyticus		4.99	0.14		1		
61012050	**Good identification**						**61012050**
					SPO		
Clostridium difficile		99.68	0.39	NIT 1	99		
Eubacterium limosum		0.32	-.06	SOR 5	1		
61013020	**Good identification**						**61013020**
					CAT		
Propionibacterium acnes		99.04	0.40	IND 87	95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Atopobium minutum		0.54	0.09	ESL 1	1	
Eubacterium limosum		0.43	-.06		1	
61014000	**Low discrimination**					**61014000**
					SPO	
Clostridium glycolicum		89.41	0.43	NIT 1	99	
Eubacterium limosum		3.19	0.15	SOR 95	1	
Atopobium minutum		2.26	0.25		1	
61014040	**Very good identification**					**61014040**
Clostridium glycolicum		99.98	0.65	NIT 1		
Clostridium botulinum biovar A		0.02	0.02			
61015040	**Good identification**					**61015040**
Clostridium glycolicum		100.0	0.49	NIT 1		
61016010	**Good identification**					**61016010**
					SPO	
Clostridium difficile		99.68	0.39	NIT 1	99	
Eubacterium limosum		0.32	-.06	XYL 5	1	
61020000	**Low discrimination**					**61020000**
					SPO	
Clostridium botulinum biovar A		50.13	0.56	NIT 1	99	
Clostridium sporogenes		49.16	0.53		99	
Atopobium minutum		0.31	0.25		1	
61020002	**Good identification**					**61020002**
					SPO	
Clostridium sporogenes		95.83	0.45	NIT 1	99	
Clostridium botulinum biovar A		2.96	0.21		99	
Atopobium minutum		1.21	0.22		1	
61020040	**Good identification**					**61020040**
Clostridium botulinum biovar A		96.49	0.47	NIT 1		
Clostridium sporogenes		3.20	0.19			
Clostridium novyi biovar A		0.31	0.16			
61030000	**Low discrimination**					**61030000**
					SPO	
Clostridium sporogenes		67.06	0.40	NIT 1	99	
Clostridium botulinum biovar A		24.74	0.35		99	
Eubacterium limosum		3.11	0.15		1	
61032010	**Good identification**					**61032010**
					SPO	
Clostridium difficile		99.78	0.42	NIT 1	99	
Eubacterium limosum		0.22	-.06	TRE 7	1	
61052010	**Good identification**					**61052010**
					SPO	
Clostridium difficile		99.68	0.39	NIT 1	99	
Eubacterium limosum		0.32	-.06	CEL 5	1	
61100000	**Low discrimination**					**61100000**
Clostridium sporogenes		58.35	0.66	NIT 1		
Clostridium botulinum biovar A		34.95	0.64			
Clostridium bifermentans		4.19	0.48			
61100001	**Low discrimination**					**61100001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium novyi biovar A		59.68	0.53	NIT 1		
Clostridium sporogenes		22.81	0.32			
Clostridium botulinum biovar A		13.66	0.30			
61100002	**Very good identification**					**61100002**
Clostridium sporogenes		96.45	0.58	NIT 1		
Clostridium botulinum biovar A		1.75	0.30			
Clostridium novyi biovar A		1.35	0.40			
61100003	**Low discrimination**					**61100003**
Clostridium novyi biovar A		58.08	0.40	NIT 1		
Clostridium sporogenes		41.92	0.24			
61100010	**Low discrimination**					**61100010**
Clostridium novyi biovar A		60.90	0.53	NIT 1		
Clostridium sporogenes		23.27	0.32			
Clostridium botulinum biovar A		13.94	0.30			
61100011	**Good identification**					**61100011**
Clostridium novyi biovar A		99.62	0.53	NIT 1		
Clostridium sporogenes		0.39	-.03			
61100012	**Low discrimination**					**61100012**
Clostridium novyi biovar A		58.08	0.40	NIT 1		
Clostridium sporogenes		41.92	0.24			
61100013	**Good identification**					**61100013**
Clostridium novyi biovar A		100.0	0.40	NIT 1		
61100021	**Good identification**					**61100021**
Clostridium novyi biovar A		98.48	0.43	NIT 1		
Clostridium sporogenes		1.52	-.03			
61100030	**Good identification**					**61100030**
Clostridium novyi biovar A		98.48	0.43	NIT 1		
Clostridium sporogenes		1.52	-.03			
61100031	**Good identification**					**61100031**
Clostridium novyi biovar A		100.0	0.43	NIT 1		
61100040	**Low discrimination**					**61100040**
Clostridium botulinum biovar A		83.31	0.55	NIT 1		
Clostridium novyi biovar A		8.20	0.50			
Clostridium sporogenes		4.70	0.32			
61100041	**Acceptable identification**					**61100041**
Clostridium novyi biovar A		90.22	0.50	NIT 1		
Clostridium botulinum biovar A		9.26	0.21			
Clostridium sporogenes		0.52	-.03			
61100043	**Good identification**					**61100043**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61100050	**Acceptable identification**						**61100050**
Clostridium novyi biovar A		90.22	0.50	NIT 1			
Clostridium botulinum biovar A		9.26	0.21				
Clostridium sporogenes		0.52	-.03				
61100051	**Good identification**						**61100051**
Clostridium novyi biovar A		100.0	0.50	NIT 1			
61100052	**Good identification**						**61100052**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61100053	**Good identification**						**61100053**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61100060	**Low discrimination**						**61100060**
Clostridium novyi biovar A		67.71	0.40	NIT 1			
Clostridium botulinum biovar A		27.78	0.21				
Clostridium bifermentans		2.94	0.04				
61100061	**Good identification**						**61100061**
Clostridium novyi biovar A		100.0	0.40	NIT 1			
61100070	**Good identification**						**61100070**
Clostridium novyi biovar A		100.0	0.40	NIT 1			
61100071	**Good identification**						**61100071**
Clostridium novyi biovar A		100.0	0.40	NIT 1			
61101001	**Low discrimination**						**61101001**
Clostridium novyi biovar A		85.01	0.37	NIT 1			
Clostridium chauvoei		8.15	0.09	GAL 10			
Clostridium sordellii		3.89	-.03				
61101010	**Good identification**						**61101010**
Clostridium novyi biovar A		96.64	0.37	NIT 1			
Clostridium sporogenes		3.36	-.03	GAL 10			
61101011	**Good identification**						**61101011**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61101101	**Good identification**						**61101101**
					COC	SPO	
Clostridium chauvoei		99.53	0.43	SUC 99	1	99	
Blautia hansenii		0.47	-.03	MNS 90	99	1	
61101121	**Good identification**						**61101121**
Clostridium chauvoei		100.0	0.35	SUC 99			
61104040	**Good identification**						**61104040**
Clostridium glycolicum		97.83	0.50	NIT 1			
Clostridium botulinum biovar A		1.88	0.21	FRU 90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium novyi biovar A		0.19	0.16				
61110000	**Low discrimination**						**61110000**
					GRA	SPO	
Clostridium sporogenes		74.94	0.53	NIT 1	99	99	
Clostridium botulinum biovar A		16.24	0.44		99	99	
Prevotella disiens		2.16	0.23		1	1	
61110001	**Low discrimination**						**61110001**
Clostridium novyi biovar A		50.06	0.37	NIT 1			
Clostridium sporogenes		30.39	0.19	FRU 10			
Clostridium sordellii		12.97	0.10				
61110002	**Good identification**						**61110002**
					SPO		
Clostridium sporogenes		96.17	0.45	NIT 1	99		
Atopobium minutum		1.14	0.22		1		
Clostridium novyi biovar A		0.85	0.24		99		
61110010	**Low discrimination**						**61110010**
Clostridium novyi biovar A		51.38	0.37	NIT 1			
Clostridium sporogenes		31.19	0.19	FRU 10			
Clostridium difficile		10.68	0.10				
61110011	**Good identification**						**61110011**
Clostridium novyi biovar A		100.0	0.37	NIT 1			
61110040	**Low discrimination**						**61110040**
Clostridium glycolicum		72.19	0.44	NIT 1			
Clostridium botulinum biovar A		18.51	0.35	XYL 95			
Clostridium novyi biovar A		3.17	0.34				
61111101	**Good identification**						**61111101**
Clostridium chauvoei		100.0	0.50	SUC 99			
61111103	**Good identification**						**61111103**
Clostridium chauvoei		100.0	0.37	SUC 99			
61111121	**Good identification**						**61111121**
Clostridium chauvoei		100.0	0.41	SUC 99			
61114000	**Acceptable identification**						**61114000**
Clostridium glycolicum		94.64	0.44	NIT 1			
Clostridium sporogenes		3.79	0.19	SOR 95			
Clostridium botulinum biovar A		0.82	0.10				
61114040	**Very good identification**						**61114040**
Clostridium glycolicum		99.99	0.66	NIT 1			
Clostridium botulinum biovar A		0.01	0.01				
61115040	**Good identification**						**61115040**
Clostridium glycolicum		100.0	0.50	NIT 1			
61120000	**Low discrimination**						**61120000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sporogenes		68.84	0.58	NIT 1		
Clostridium botulinum biovar A		30.93	0.54			
Clostridium bifermentans		0.15	0.14			
61120002	**Good identification**					**61120002**
Clostridium sporogenes		98.66	0.50	NIT 1		
Clostridium botulinum biovar A		1.34	0.20			
61120040	**Acceptable identification**					**61120040**
Clostridium botulinum biovar A		92.66	0.45	NIT 1		
Clostridium sporogenes		6.97	0.24			
Clostridium novyi biovar A		0.37	0.16			
61130000	**Low discrimination**					**61130000**
Clostridium sporogenes		86.02	0.45	NIT 1		
Clostridium botulinum biovar A		13.98	0.34			
61130002	**Good identification**					**61130002**
Clostridium sporogenes		99.51	0.37	NIT 1		
Clostridium botulinum biovar A		0.49	-.01			
61201000	**Very good identification**					**61201000**
					SPO	
Eubacterium tortuosum		99.94	0.64	SUC 99	1	
Atopobium minutum		0.04	0.17		1	
Clostridium novyi biovar A		0.03	0.19		99	
61201002	**Good identification**					**61201002**
Eubacterium tortuosum		99.85	0.52	SUC 99		
Atopobium minutum		0.15	0.14			
61201020	**Good identification**					**61201020**
Eubacterium tortuosum		99.93	0.52	SUC 99		
Atopobium minutum		0.07	0.09			
61201022	**Good identification**					**61201022**
Eubacterium tortuosum		99.72	0.39	SUC 99		
Atopobium minutum		0.28	0.06			
61210010	**Good identification**					**61210010**
Clostridium difficile		98.52	0.35	NIT 1		
Clostridium novyi biovar A		1.48	0.19	MAN 90		
61210100	**Good identification**					**61210100**
Fusobacterium mortiferum		100.0	0.40	NIT 1		
61211100	**Good identification**					**61211100**
					GRA	
Fusobacterium mortiferum		98.35	0.43	NIT 1	1	
Eubacterium tortuosum		1.65	-.04		99	
61212000	**Low discrimination**					**61212000**
					SPO	
Clostridium difficile		77.53	0.40	NIT 1	99	
Eubacterium limosum		22.24	0.29		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Pseudoramibacter alactolyticus		0.24	-.07		1	
61212010	**Good identification**					**61212010**
					SPO	
Clostridium difficile		99.94	0.52	NIT 1	99	
Eubacterium limosum		0.06	-.06		1	
61241000	**Good identification**					**61241000**
Eubacterium tortuosum		100.0	0.48	SUC 99		
61241002	**Good identification**					**61241002**
Eubacterium tortuosum		100.0	0.35	SUC 99		
61241020	**Good identification**					**61241020**
Eubacterium tortuosum		100.0	0.35	SUC 99		
61357100	**Good identification**					**61357100**
Actinomyces israelii		100.0	0.36	SUC 98		
61357102	**Good identification**					**61357102**
Actinomyces israelii		100.0	0.39	SUC 98		
61357502	**Good identification**					**61357502**
Actinomyces israelii		100.0	0.38	SUC 98		
61375102	**Good identification**					**61375102**
Actinomyces israelii		100.0	0.36	SUC 98		
61377000	**Good identification**					**61377000**
Actinomyces israelii		100.0	0.36	SUC 98		
61377002	**Good identification**					**61377002**
Actinomyces israelii		100.0	0.39	SUC 98		
61377100	**Good identification**					**61377100**
Actinomyces israelii		100.0	0.46	SUC 98		
61377102	**Good identification**					**61377102**
Actinomyces israelii		100.0	0.49	SUC 98		
61377112	**Good identification**					**61377112**
Actinomyces israelii		100.0	0.35	SUC 98		
61377140	**Good identification**					**61377140**
Actinomyces israelii		100.0	0.39	SUC 98		
61377142	**Good identification**					**61377142**
Actinomyces israelii		100.0	0.42	SUC 98		
61377402	**Good identification**					**61377402**
Actinomyces israelii		100.0	0.38	SUC 98		
61377500	**Good identification**					**61377500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		100.0	0.45	SUC 98				
61377502	**Good identification**							**61377502**
Actinomyces israelii		100.0	0.48	SUC 98				
61377540	**Good identification**							**61377540**
Actinomyces israelii		100.0	0.38	SUC 98				
61377542	**Good identification**							**61377542**
Actinomyces israelii		100.0	0.41	SUC 98				
61410100	**Good identification**							**61410100**
Fusobacterium mortiferum		100.0	0.40	NIT 1				
61411000	**Low discrimination**							**61411000**
					GRA			
Fusobacterium mortiferum		87.87	0.37	NIT 1	1			
Atopobium minutum		12.13	0.17		99			
61411100	**Good identification**							**61411100**
					COC	GRA		
Fusobacterium mortiferum		97.81	0.43	NIT 1	1	1		
Blautia hansenii		2.19	-.03		99	99		
61501100	**Very good identification**							**61501100**
Blautia hansenii		100.0	0.66	NIT 1				
61501102	**Good identification**							**61501102**
					COC	GRA		
Blautia hansenii		99.93	0.50	NIT 1	99	99		
Capnocytophaga ochracea		0.07	-.01	BGL 10	1	1		
61501120	**Good identification**							**61501120**
					COC	CAT	GRA	
Blautia hansenii		99.85	0.50	NIT 1	99	1	99	
Prevotella melaninogenica		0.08	-.04	NAG 10	1	1	1	
Capnocytophaga ochracea		0.07	-.01		1	90	1	
61501500	**Good identification**							**61501500**
Blautia hansenii		100.0	0.53	NIT 1				
61501502	**Good identification**							**61501502**
Blautia hansenii		100.0	0.37	NIT 1				
61501520	**Good identification**							**61501520**
Blautia hansenii		100.0	0.37	NIT 1				
61505100	**Good identification**							**61505100**
Blautia hansenii		100.0	0.53	NIT 1				
61505102	**Good identification**							**61505102**
Blautia hansenii		100.0	0.37	NIT 1				
61505120	**Good identification**							**61505120**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia hansenii		100.0	0.37	NIT 1			
61505500	**Good identification**						**61505500**
Blautia hansenii		100.0	0.40	NIT 1			
61511122	**Good identification**						**61511122**
					GRA	CAT	
Capnocytophaga ochracea		99.26	0.50	SUC 99	1	90	
Clostridium perfringens		0.22	0.07	MNS 90	99	1	
Prevotella melaninogenica		0.21	0.00		1	1	
61511522	**Good identification**						**61511522**
					GRA	CAT	
Capnocytophaga ochracea		99.51	0.37	SUC 99	1	90	
Bifidobacterium longum subsp. longum		0.49	0.05	MNS 90	99	1	
61515122	**Good identification**						**61515122**
Capnocytophaga ochracea		96.25	0.39	SUC 99			
Bacteroides fragilis		3.75	0.06	MNS 90			
61551122	**Good identification**						**61551122**
					CAT		
Capnocytophaga ochracea		98.19	0.39	SUC 99	90		
Prevotella oralis		1.82	0.09	MNS 90	1		
61577100	**Good identification**						**61577100**
Actinomyces israelii		100.0	0.38	SUC 98			
61577102	**Good identification**						**61577102**
Actinomyces israelii		100.0	0.41	SUC 98			
61577500	**Good identification**						**61577500**
Actinomyces israelii		100.0	0.37	SUC 98			
61577502	**Good identification**						**61577502**
Actinomyces israelii		100.0	0.40	SUC 98			
61601000	**Very good identification**						**61601000**
Eubacterium tortuosum		100.0	0.56	SUC 99			
61601002	**Good identification**						**61601002**
Eubacterium tortuosum		100.0	0.43	SUC 99			
61601020	**Good identification**						**61601020**
Eubacterium tortuosum		100.0	0.43	SUC 99			
61610000	**Good identification**						**61610000**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
61610100	**Good identification**						**61610100**
Fusobacterium mortiferum		100.0	0.45	NIT 1			
61611000	**Low discrimination**						**61611000**
					GRA		
Fusobacterium mortiferum		59.98	0.42	NIT 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		40.02	0.22		99	
61611100	**Good identification**					**61611100**
Fusobacterium mortiferum		100.0	0.48	NIT 1		
61630100	**Good identification**					**61630100**
Fusobacterium mortiferum		100.0	0.36	NIT 1		
61631100	**Good identification**					**61631100**
Fusobacterium mortiferum		100.0	0.39	NIT 1		
61641000	**Good identification**					**61641000**
Eubacterium tortuosum		100.0	0.40	SUC 99		
61650100	**Good identification**					**61650100**
Fusobacterium mortiferum		100.0	0.37	NIT 1		
61651100	**Good identification**					**61651100**
Fusobacterium mortiferum		100.0	0.40	NIT 1		
61737100	**Good identification**					**61737100**
Actinomyces israelii		100.0	0.37	SUC 98		
61737102	**Good identification**					**61737102**
Actinomyces israelii		100.0	0.40	SUC 98		
61737500	**Good identification**					**61737500**
Actinomyces israelii		100.0	0.35	SUC 98		
61737502	**Good identification**					**61737502**
Actinomyces israelii		100.0	0.38	SUC 98		
61755102	**Good identification**					**61755102**
Actinomyces israelii		100.0	0.36	SUC 98		
61757000	**Good identification**					**61757000**
Actinomyces israelii		100.0	0.36	SUC 98		
61757002	**Good identification**					**61757002**
Actinomyces israelii		100.0	0.39	SUC 98		
61757100	**Good identification**					**61757100**
Actinomyces israelii		100.0	0.46	SUC 98		
61757102	**Good identification**					**61757102**
Actinomyces israelii		100.0	0.49	SUC 98		
61757112	**Good identification**					**61757112**
Actinomyces israelii		100.0	0.35	SUC 98		
61757140	**Good identification**					**61757140**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.39	SUC 98		
61757142	**Good identification**					**61757142**
Actinomyces israelii		100.0	0.42	SUC 98		
61757402	**Good identification**					**61757402**
Actinomyces israelii		100.0	0.38	SUC 98		
61757500	**Good identification**					**61757500**
Actinomyces israelii		100.0	0.45	SUC 98		
61757502	**Good identification**					**61757502**
Actinomyces israelii		97.41	0.48	SUC 98		
Bifidobacterium adolescentis		2.59	0.15			
61757540	**Good identification**					**61757540**
Actinomyces israelii		100.0	0.38	SUC 98		
61757542	**Good identification**					**61757542**
Actinomyces israelii		100.0	0.41	SUC 98		
61773100	**Low discrimination**					**61773100**
Actinomyces israelii		86.77	0.38	SUC 98		
Actinomyces naeslundii		13.23	0.33	XYL 92		
61773101	**Good identification**					**61773101**
Actinomyces naeslundii		100.0	0.36	ESL 12		
61773102	**Acceptable identification**					**61773102**
Actinomyces israelii		90.77	0.41	SUC 98		
Actinomyces naeslundii		9.23	0.33	XYL 92		
61773103	**Good identification**					**61773103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
61773121	**Good identification**					**61773121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
61773123	**Good identification**					**61773123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
61773500	**Good identification**					**61773500**
Actinomyces israelii		100.0	0.37	SUC 98		
61773502	**Good identification**					**61773502**
Actinomyces israelii		100.0	0.40	SUC 98		
61775002	**Good identification**					**61775002**
Actinomyces israelii		100.0	0.36	SUC 98		
61775100	**Good identification**					**61775100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.43	SUC 98			
61775102	**Good identification**						**61775102**
Actinomyces israelii		100.0	0.46	SUC 98			
61775140	**Good identification**						**61775140**
Actinomyces israelii		100.0	0.36	SUC 98			
61775142	**Good identification**						**61775142**
Actinomyces israelii		100.0	0.39	SUC 98			
61775500	**Low discrimination**						**61775500**
					GRA	SPO	
Actinomyces israelii		73.00	0.42	SUC 98	99	1	
Mitsuokella multacida		23.77	0.21		1	1	
Clostridium butyricum		3.23	-.03		99	99	
61775502	**Acceptable identification**						**61775502**
					GRA		
Actinomyces israelii		94.63	0.45	SUC 98	99		
Bifidobacterium adolescentis		3.09	0.13		99		
Mitsuokella multacida		2.28	0.04		1		
61775542	**Acceptable identification**						**61775542**
					GRA		
Actinomyces israelii		94.16	0.38	SUC 98	99		
Mitsuokella multacida		5.84	0.04		1		
61777000	**Good identification**						**61777000**
Actinomyces israelii		100.0	0.46	SUC 98			
61777002	**Good identification**						**61777002**
Actinomyces israelii		100.0	0.49	SUC 98			
61777040	**Good identification**						**61777040**
Actinomyces israelii		100.0	0.39	SUC 98			
61777042	**Good identification**						**61777042**
Actinomyces israelii		100.0	0.42	SUC 98			
61777100	**Very good identification**						**61777100**
Actinomyces israelii		100.0	0.56	SUC 98			
61777102	**Very good identification**						**61777102**
Actinomyces israelii		100.0	0.59	SUC 98			
61777110	**Good identification**						**61777110**
Actinomyces israelii		100.0	0.42	SUC 98			
61777112	**Good identification**						**61777112**
Actinomyces israelii		100.0	0.45	SUC 98			
61777120	**Good identification**						**61777120**
Actinomyces israelii		100.0	0.36	SUC 98			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
61777122	**Good identification**					**61777122**
Actinomyces israelii		100.0	0.39	SUC 98		
61777140	**Good identification**					**61777140**
Actinomyces israelii		100.0	0.49	SUC 98		
61777142	**Good identification**					**61777142**
Actinomyces israelii		100.0	0.52	SUC 98		
61777150	**Good identification**					**61777150**
Actinomyces israelii		100.0	0.35	SUC 98		
61777152	**Good identification**					**61777152**
Actinomyces israelii		100.0	0.38	SUC 98		
61777300	**Good identification**					**61777300**
Actinomyces israelii		100.0	0.37	SUC 98		
61777302	**Good identification**					**61777302**
Actinomyces israelii		100.0	0.40	SUC 98		
61777400	**Good identification**					**61777400**
Actinomyces israelii		100.0	0.44	SUC 98		
61777402	**Good identification**					**61777402**
Actinomyces israelii		100.0	0.47	SUC 98		
61777440	**Good identification**					**61777440**
Actinomyces israelii		100.0	0.37	SUC 98		
61777442	**Good identification**					**61777442**
Actinomyces israelii		100.0	0.40	SUC 98		
61777500	**Low discrimination**					**61777500**
					GRA	
Actinomyces israelii		88.18	0.55	SUC 98	99	
Mitsuokella multacida		11.83	0.27		1	
61777502	**Very good identification**					**61777502**
					GRA	
Actinomyces israelii		98.56	0.58	SUC 98	99	
Mitsuokella multacida		0.98	0.11		1	
Bifidobacterium adolescentis		0.47	0.12		99	
61777510	**Low discrimination**					**61777510**
					GRA	
Actinomyces israelii		76.97	0.41	SUC 98	99	
Mitsuokella multacida		23.03	0.19	MLZ 13	1	
61777512	**Acceptable identification**					**61777512**
Actinomyces israelii		91.65	0.44	SUC 98		
Bifidobacterium adolescentis		3.54	0.13	MLZ 13		
Bifidobacterium dentium		2.78	-.07			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
61777522	**Good identification**					**61777522**
Actinomyces israelii		100.0	0.37	SUC 98		
61777540	**Low discrimination**					**61777540**
					GRA	
Actinomyces israelii		74.36	0.48	SUC 98	99	
Mitsuokella multacida		25.64	0.27		1	
61777542	**Good identification**					**61777542**
					GRA	
Actinomyces israelii		97.51	0.51	SUC 98	99	
Mitsuokella multacida		2.49	0.11		1	
61777552	**Acceptable identification**					**61777552**
					GRA	
Actinomyces israelii		94.61	0.37	SUC 98	99	
Mitsuokella multacida		5.39	0.03	MLZ 13	1	
61777702	**Low discrimination**					**61777702**
					GRA	
Actinomyces israelii		80.31	0.38	SUC 98	99	
Mitsuokella multacida		19.69	0.15	RHA 7	1	
62101101	**Good identification**					**62101101**
Clostridium chauvoei		100.0	0.43	GLU 99		
62101121	**Good identification**					**62101121**
Clostridium chauvoei		100.0	0.35	GLU 99		
62111101	**Good identification**					**62111101**
Clostridium chauvoei		100.0	0.50	GLU 99		
62111103	**Good identification**					**62111103**
Clostridium chauvoei		100.0	0.37	GLU 99		
62111121	**Good identification**					**62111121**
Clostridium chauvoei		100.0	0.41	GLU 99		
62201000	**Very good identification**					**62201000**
Eubacterium tortuosum		100.0	0.64	GLU 99		
62201002	**Good identification**					**62201002**
Eubacterium tortuosum		100.0	0.52	GLU 99		
62201020	**Good identification**					**62201020**
Eubacterium tortuosum		100.0	0.52	GLU 99		
62201022	**Good identification**					**62201022**
Eubacterium tortuosum		100.0	0.39	GLU 99		
62241000	**Good identification**					**62241000**
Eubacterium tortuosum		100.0	0.48	GLU 99		
62241002	**Good identification**					**62241002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium tortuosum		100.0	0.35	GLU 99			
62241020	**Good identification**						**62241020**
Eubacterium tortuosum		100.0	0.35	GLU 99			
62377100	**Good identification**						**62377100**
Actinomyces israelii		100.0	0.41	GLU 99			
62377102	**Good identification**						**62377102**
Actinomyces israelii		100.0	0.44	GLU 99			
62377142	**Good identification**						**62377142**
Actinomyces israelii		100.0	0.37	GLU 99			
62377500	**Good identification**						**62377500**
Actinomyces israelii		100.0	0.40	GLU 99			
62377502	**Good identification**						**62377502**
Actinomyces israelii		100.0	0.43	GLU 99			
62377542	**Good identification**						**62377542**
Actinomyces israelii		100.0	0.36	GLU 99			
62511122	**Good identification**						**62511122**
					CAT		
Capnocytophaga ochracea		97.71	0.50	GLU 99	90		
Prevotella melaninogenica		2.30	0.18	MNS 90	1		
62511522	**Good identification**						**62511522**
					GRA	CAT	
Capnocytophaga ochracea		99.51	0.37	GLU 99	1	90	
Bifidobacterium longum subsp. longum		0.49	0.05	MNS 90	99	1	
62515122	**Good identification**						**62515122**
Capnocytophaga ochracea		96.25	0.39	GLU 99			
Bacteroides fragilis		3.75	0.06	MNS 90			
62551122	**Good identification**						**62551122**
					CAT		
Capnocytophaga ochracea		98.97	0.39	GLU 99	90		
Prevotella melaninogenica		1.03	0.02	MNS 90	1		
62577102	**Good identification**						**62577102**
Actinomyces israelii		100.0	0.36	GLU 99			
62601000	**Very good identification**						**62601000**
Eubacterium tortuosum		100.0	0.56	GLU 99			
62601002	**Good identification**						**62601002**
Eubacterium tortuosum		100.0	0.43	GLU 99			
62601020	**Good identification**						**62601020**
Eubacterium tortuosum		100.0	0.43	GLU 99			
62641000	**Good identification**						**62641000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium tortuosum		100.0	0.40	GLU 99			
62757100	**Good identification**						**62757100**
Actinomyces israelii		100.0	0.41	GLU 99			
62757102	**Good identification**						**62757102**
Actinomyces israelii		100.0	0.44	GLU 99			
62757142	**Good identification**						**62757142**
Actinomyces israelii		100.0	0.37	GLU 99			
62757500	**Good identification**						**62757500**
Actinomyces israelii		100.0	0.40	GLU 99			
62757502	**Acceptable identification**						**62757502**
Actinomyces israelii		94.90	0.43	GLU 99			
Bifidobacterium adolescentis		5.10	0.15				
62757542	**Good identification**						**62757542**
Actinomyces israelii		100.0	0.36	GLU 99			
62773101	**Good identification**						**62773101**
Actinomyces naeslundii		100.0	0.36	ESL 12			
62773102	**Low discrimination**						**62773102**
Actinomyces israelii		82.96	0.36	GLU 99			
Actinomyces naeslundii		17.04	0.33	XYL 92			
62773103	**Good identification**						**62773103**
Actinomyces naeslundii		100.0	0.36	ESL 12			
62773121	**Good identification**						**62773121**
Actinomyces naeslundii		100.0	0.36	ESL 12			
62773123	**Good identification**						**62773123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
62775100	**Good identification**						**62775100**
Actinomyces israelii		100.0	0.38	GLU 99			
62775102	**Good identification**						**62775102**
Actinomyces israelii		100.0	0.41	GLU 99			
62775500	**Low discrimination**						**62775500**
					GRA	SPO	
Actinomyces israelii		57.23	0.37	GLU 99	99	1	
Mitsuokella multacida		37.66	0.21		1	1	
Clostridium butyricum		5.11	-.03		99	99	
62775502	**Low discrimination**						**62775502**
					GRA		
Actinomyces israelii		89.71	0.40	GLU 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		5.92	0.13		99	
Mitsuokella multacida		4.37	0.04		1	
62777000	**Good identification**					**62777000**
Actinomyces israelii		100.0	0.41	GLU 99		
62777002	**Good identification**					**62777002**
Actinomyces israelii		100.0	0.44	GLU 99		
62777042	**Good identification**					**62777042**
Actinomyces israelii		100.0	0.37	GLU 99		
62777100	**Good identification**					**62777100**
Actinomyces israelii		100.0	0.51	GLU 99		
62777102	**Good identification**					**62777102**
Actinomyces israelii		100.0	0.54	GLU 99		
62777110	**Good identification**					**62777110**
Actinomyces israelii		100.0	0.37	GLU 99		
62777112	**Good identification**					**62777112**
Actinomyces israelii		100.0	0.40	GLU 99		
62777140	**Good identification**					**62777140**
Actinomyces israelii		100.0	0.44	GLU 99		
62777142	**Good identification**					**62777142**
Actinomyces israelii		100.0	0.47	GLU 99		
62777400	**Good identification**					**62777400**
Actinomyces israelii		100.0	0.39	GLU 99		
62777402	**Good identification**					**62777402**
Actinomyces israelii		100.0	0.42	GLU 99		
62777442	**Good identification**					**62777442**
Actinomyces israelii		100.0	0.35	GLU 99		
62777500	**Low discrimination**					**62777500**
					GRA	
Actinomyces israelii		78.68	0.49	GLU 99	99	
Mitsuokella multacida		21.32	0.27		1	
62777502	**Good identification**					**62777502**
					GRA	
Actinomyces israelii		97.13	0.52	GLU 99	99	
Mitsuokella multacida		1.95	0.11		1	
Bifidobacterium adolescentis		0.93	0.12		99	
62777510	**Low discrimination**					**62777510**
					GRA	
Actinomyces israelii		62.33	0.35	GLU 99	99	
Mitsuokella multacida		37.67	0.19	MLZ 13	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
62777512	**Low discrimination**						**62777512**
Actinomyces israelii		84.46	0.38	GLU 99			
Bifidobacterium adolescentis		6.59	0.13	MLZ 13			
Bifidobacterium dentium		5.18	-.07				
62777540	**Low discrimination**						**62777540**
					GRA		
Actinomyces israelii		58.94	0.42	GLU 99	99		
Mitsuokella multacida		41.06	0.27		1		
62777542	**Good identification**						**62777542**
					GRA		
Actinomyces israelii		95.09	0.45	GLU 99	99		
Mitsuokella multacida		4.91	0.11		1		
63000000	**Low discrimination**						**63000000**
					SPO		
Clostridium botulinum biovar A		70.83	0.47	NIT 1	99		
Eubacterium tortuosum		12.56	0.30	SUC 7	1		
Atopobium minutum		7.56	0.38		1		
63000040	**Good identification**						**63000040**
					SPO		
Clostridium botulinum biovar A		97.78	0.38	NIT 1	99		
Clostridium novyi biovar A		1.06	0.16	SUC 7	99		
Eubacterium tortuosum		0.59	-.04		1		
63001000	**Very good identification**						**63001000**
					SPO		
Eubacterium tortuosum		99.74	0.64	SAL 99	1		
Atopobium minutum		0.20	0.30		1		
Clostridium botulinum biovar A		0.06	0.13		99		
63001002	**Good identification**						**63001002**
Eubacterium tortuosum		99.24	0.52	SAL 99			
Atopobium minutum		0.76	0.27				
63001020	**Good identification**						**63001020**
Eubacterium tortuosum		99.62	0.52	SAL 99			
Atopobium minutum		0.38	0.21				
63001022	**Good identification**						**63001022**
Eubacterium tortuosum		98.58	0.39	SAL 99			
Atopobium minutum		1.42	0.18				
63001101	**Good identification**						**63001101**
					SPO		
Clostridium chauvoei		99.65	0.43	MNS 90	99		
Eubacterium tortuosum		0.35	-.04	MLT 99	1		
63001121	**Good identification**						**63001121**
Clostridium chauvoei		100.0	0.35	MNS 90			
63010000	**Low discrimination**						**63010000**
					GRA	SPO	
Atopobium minutum		26.48	0.37	NIT 1	99	1	
Fusobacterium mortiferum		23.89	0.42	SUC 5	1	1	
Clostridium botulinum biovar A		17.17	0.26		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63010100	**Low discrimination**						**63010100**
					GRA		
Fusobacterium mortiferum		87.92	0.48	NIT 1	1		
Actinomyces meyeri		10.34	0.23		99		
Actinomyces odontolyticus		1.74	0.13		99		
63011100	**Low discrimination**						**63011100**
					GRA	SPO	
Fusobacterium mortiferum		82.43	0.51	NIT 1	1	1	
Actinomyces meyeri		9.70	0.26		99	1	
Clostridium chauvoei		3.16	0.15		99	99	
63011101	**Good identification**						**63011101**
					GRA	SPO	
Clostridium chauvoei		99.74	0.50	MNS 90	99	99	
Fusobacterium mortiferum		0.27	0.17	MLT 99	1	1	
63011103	**Good identification**						**63011103**
Clostridium chauvoei		100.0	0.37	MNS 90			
63011121	**Good identification**						**63011121**
Clostridium chauvoei		100.0	0.41	MNS 90			
63020000	**Low discrimination**						**63020000**
					SPO		
Clostridium botulinum biovar A		87.82	0.36	NIT 1	99		
Clostridium sporogenes		11.56	0.19	SUC 7	99		
Eubacterium tortuosum		0.63	-.04		1		
63030100	**Good identification**						**63030100**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
63031000	**Acceptable identification**						**63031000**
					GRA		
Fusobacterium mortiferum		91.19	0.36	NIT 1	1		
Eubacterium saburreum		4.71	0.12		85		
Eubacterium tortuosum		4.10	-.04		99		
63031100	**Good identification**						**63031100**
Fusobacterium mortiferum		95.09	0.42	NIT 1			
Eubacterium saburreum		4.91	0.18				
63041000	**Good identification**						**63041000**
Eubacterium tortuosum		100.0	0.48	SAL 99			
63041002	**Good identification**						**63041002**
Eubacterium tortuosum		100.0	0.35	SAL 99			
63041020	**Good identification**						**63041020**
Eubacterium tortuosum		100.0	0.35	SAL 99			
63050100	**Good identification**						**63050100**
Fusobacterium mortiferum		100.0	0.40	NIT 1			
63051000	**Low discrimination**						**63051000**
					GRA		
Fusobacterium mortiferum		69.27	0.37	NIT 1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Eubacterium tortuosum		30.73	0.14		99			
63051100	**Good identification**							**63051100**
Fusobacterium mortiferum		100.0	0.43	NIT 1				
63100000	**Low discrimination**							**63100000**
					SPO			
Clostridium botulinum biovar A		71.85	0.45	NIT 1	99			
Clostridium sporogenes		16.09	0.32	SUC 7	99			
Actinomyces odontolyticus		9.82	0.38		1			
63100040	**Good identification**							**63100040**
					COC	GRA		
Clostridium botulinum biovar A		96.94	0.36	NIT 1	1	99		
Clostridium novyi biovar A		1.28	0.16	SUC 7	1	99		
Megasphaera elsdenii		1.05	0.07		99	1		
63100101	**Good identification**							**63100101**
Clostridium chauvoei		100.0	0.43	MNS 90				
63100121	**Good identification**							**63100121**
Clostridium chauvoei		100.0	0.35	MNS 90				
63100200	**Good identification**							**63100200**
					SPO			
Actinomyces odontolyticus		95.55	0.43	ESL 1	1			
Clostridium botulinum biovar A		3.64	0.11	FRU 92	99			
Clostridium sporogenes		0.82	-.03		99			
63100202	**Good identification**							**63100202**
Actinomyces odontolyticus		100.0	0.39	ESL 1				
63101001	**Good identification**							**63101001**
					SPO			
Clostridium chauvoei		99.65	0.43	MNS 90	99			
Eubacterium tortuosum		0.35	-.04	LAC 99	1			
63101021	**Good identification**							**63101021**
Clostridium chauvoei		100.0	0.35	MNS 90				
63101100	**Acceptable identification**							**63101100**
					SPO			
Clostridium chauvoei		92.38	0.43	MNS 90	99			
Actinomyces odontolyticus		4.30	0.26	URE 99	1			
Eubacterium saburreum		1.75	0.24		1			
63101101	**Very good identification**							**63101101**
Clostridium chauvoei		100.0	0.77	MNS 90				
63101103	**Very good identification**							**63101103**
Clostridium chauvoei		100.0	0.65	MNS 90				
63101111	**Good identification**							**63101111**
Clostridium chauvoei		100.0	0.43	MNS 90				
63101120	**Low discrimination**							**63101120**
					GRA	CAT	SPO	
Clostridium chauvoei		76.47	0.35	MNS 90	99	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Prevotella melaninogenica		10.08	0.14	URE 99	1	1	1	
Capnocytophaga ochracea		9.08	0.17		1	90	1	
63101121	**Very good identification**							**63101121**
Clostridium chauvoei		100.0	0.69	MNS 90				
63101123	**Very good identification**							**63101123**
					GRA	SPO		
Clostridium chauvoei		99.94	0.56	MNS 90	99	99		
Capnocytophaga ochracea		0.06	-.01		1	1		
63101131	**Good identification**							**63101131**
Clostridium chauvoei		100.0	0.35	MNS 90				
63101141	**Good identification**							**63101141**
Clostridium chauvoei		100.0	0.43	MNS 90				
63101161	**Good identification**							**63101161**
Clostridium chauvoei		100.0	0.35	MNS 90				
63101200	**Good identification**							**63101200**
Actinomyces odontolyticus		98.78	0.40	ESL 1				
Eubacterium tortuosum		1.22	-.04	FRU 92				
63101202	**Good identification**							**63101202**
Actinomyces odontolyticus		100.0	0.36	ESL 1				
63101301	**Good identification**							**63101301**
Clostridium chauvoei		100.0	0.43	MNS 90				
63101321	**Good identification**							**63101321**
Clostridium chauvoei		100.0	0.35	MNS 90				
63101501	**Good identification**							**63101501**
Clostridium chauvoei		100.0	0.43	MNS 90				
63101521	**Good identification**							**63101521**
Clostridium chauvoei		100.0	0.35	MNS 90				
63103101	**Good identification**							**63103101**
					SPO			
Clostridium chauvoei		99.72	0.43	MNS 90	99			
Propionibacterium propionicum		0.28	0.12	MAN 1	1			
63103121	**Good identification**							**63103121**
Clostridium chauvoei		100.0	0.35	MNS 90				
63104200	**Good identification**							**63104200**
Actinomyces odontolyticus		100.0	0.38	ESL 1				
63105101	**Good identification**							**63105101**
Clostridium chauvoei		100.0	0.43	MNS 90				
63105121	**Good identification**							**63105121**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium chauvoei		100.0	0.35	MNS 90				
63110000	**Low discrimination**							**63110000**
					COC	SPO	GRA	
Actinomyces odontolyticus		87.86	0.56	ESL 1	1	1	99	
Megasphaera elsdenii		5.15	0.32		99	1	1	
Clostridium botulinum biovar A		3.57	0.25		1	99	99	
63110002	**Good identification**							**63110002**
					SPO			
Actinomyces odontolyticus		95.45	0.52	ESL 1	1			
Actinomyces viscosus		2.23	0.18		1			
Clostridium sporogenes		1.31	0.11		99			
63110020	**Good identification**							**63110020**
Actinomyces odontolyticus		100.0	0.43	ESL 1				
63110022	**Good identification**							**63110022**
					GRA	CAT		
Actinomyces odontolyticus		98.60	0.39	ESL 1	99	10		
Capnocytophaga ochracea		1.40	-.01	NAG 14	1	90		
63110100	**Low discrimination**							**63110100**
					GRA			
Actinomyces odontolyticus		83.55	0.47	ESL 1	99			
Actinomyces meyeri		5.65	0.24		99			
Fusobacterium mortiferum		5.27	0.33		1			
63110101	**Good identification**							**63110101**
					SPO			
Clostridium chauvoei		99.53	0.50	MNS 90	99			
Actinomyces odontolyticus		0.35	0.13	GAL 99	1			
Propionibacterium propionicum		0.12	0.12		1			
63110102	**Low discrimination**							**63110102**
					GRA	CAT		
Actinomyces odontolyticus		76.40	0.44	ESL 1	99	10		
Actinomyces viscosus		10.06	0.22		99	76		
Capnocytophaga ochracea		6.14	0.17		1	90		
63110103	**Acceptable identification**							**63110103**
					SPO			
Clostridium chauvoei		93.04	0.37	MNS 90	99			
Actinomyces viscosus		4.88	0.14	GAL 99	1			
Actinomyces odontolyticus		1.12	0.10		1			
63110121	**Good identification**							**63110121**
Clostridium chauvoei		100.0	0.41	MNS 90				
63110200	**Very good identification**							**63110200**
Actinomyces odontolyticus		100.0	0.61	ESL 1				
63110202	**Very good identification**							**63110202**
Actinomyces odontolyticus		100.0	0.57	ESL 1				
63110220	**Good identification**							**63110220**
Actinomyces odontolyticus		100.0	0.47	ESL 1				
63110222	**Good identification**							**63110222**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces odontolyticus		100.0	0.44	ESL 1				
63110300	**Good identification**							**63110300**
Actinomyces odontolyticus		100.0	0.52	ESL 1				
63110302	**Good identification**							**63110302**
Actinomyces odontolyticus		100.0	0.49	ESL 1				
63110320	**Good identification**							**63110320**
Actinomyces odontolyticus		100.0	0.39	ESL 1				
63110322	**Good identification**							**63110322**
					GRA	CAT		
Actinomyces odontolyticus		97.74	0.35	ESL 1	99	10		
Capnocytophaga ochracea		2.26	-.01	NAG 14	1	90		
63111000	**Acceptable identification**							**63111000**
					GRA			
Actinomyces odontolyticus		93.08	0.53	ESL 1	99			
Fusobacterium mortiferum		1.79	0.29		1			
Eubacterium saburreum		1.34	0.25		85			
63111001	**Good identification**							**63111001**
					SPO			
Clostridium chauvoei		99.15	0.50	MNS 90	99			
Actinomyces odontolyticus		0.73	0.19	LAC 99	1			
Actinomyces naeslundii		0.13	0.23		1			
63111002	**Acceptable identification**							**63111002**
					GRA	CAT		
Actinomyces odontolyticus		92.20	0.49	ESL 1	99	10		
Capnocytophaga ochracea		3.51	0.17		1	90		
Actinomyces viscosus		1.98	0.14		99	76		
63111003	**Good identification**							**63111003**
					SPO			
Clostridium chauvoei		95.16	0.37	MNS 90	99			
Actinomyces odontolyticus		2.43	0.15	LAC 99	1			
Actinomyces viscosus		1.72	0.06		1			
63111020	**Low discrimination**							**63111020**
					GRA	CAT		
Actinomyces odontolyticus		61.43	0.40	ESL 1	99	10		
Clostridium perfringens		24.52	0.28	NAG 14	99	1		
Capnocytophaga ochracea		8.81	0.17		1	90		
63111021	**Good identification**							**63111021**
					SPO			
Clostridium chauvoei		99.26	0.41	MNS 90	99			
Actinomyces naeslundii		0.38	0.23	LAC 99	1			
Actinomyces odontolyticus		0.36	0.05		1			
63111100	**Low discrimination**							**63111100**
					GRA	SPO	CAT	
Clostridium chauvoei		64.80	0.50	MNS 90	99	99	1	
Actinomyces odontolyticus		14.87	0.44	URE 99	99	1	10	
Capnocytophaga ochracea		12.09	0.35		1	1	90	
63111101	**Very good identification**							**63111101**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium chauvoei		99.99	0.84	MNS 90	99			
Actinomyces naeslundii		0.00	0.31		1			
Propionibacterium propionicum		0.00	0.12		1			
63111102	**Low discrimination**							**63111102**
					GRA	SPO	CAT	
Capnocytophaga ochracea		81.73	0.51	MNS 90	1	1	90	
Clostridium chauvoei		8.59	0.37	RAF 90	99	99	1	
Actinomyces odontolyticus		6.85	0.41	NAG 90	99	1	10	
63111103	**Very good identification**							**63111103**
					GRA	SPO		
Clostridium chauvoei		99.83	0.71	MNS 90	99	99		
Capnocytophaga ochracea		0.10	0.17		1	1		
Actinomyces viscosus		0.03	0.10		99	1		
63111111	**Good identification**							**63111111**
Clostridium chauvoei		100.0	0.50	MNS 90				
63111113	**Good identification**							**63111113**
					GRA	SPO		
Clostridium chauvoei		98.94	0.37	MNS 90	99	99		
Capnocytophaga ochracea		1.06	0.01	MLZ 1	1	1		
63111120	**Low discrimination**							**63111120**
					GRA	CAT		
Capnocytophaga ochracea		65.08	0.51	MNS 90	1	90		
Clostridium perfringens		13.42	0.42	RAF 90	99	1		
Clostridium chauvoei		12.92	0.41	BGL 90	99	1		
63111121	**Very good identification**							**63111121**
					GRA	SPO		
Clostridium chauvoei		99.92	0.76	MNS 90	99	99		
Capnocytophaga ochracea		0.05	0.17		1	1		
Actinomyces naeslundii		0.01	0.31		99	1		
63111122	**Very good identification**							**63111122**
					GRA	CAT		
Capnocytophaga ochracea		98.15	0.67	MNS 90	1	90		
Clostridium perfringens		1.06	0.37	RAF 90	99	1		
Clostridium chauvoei		0.38	0.28		99	1		
63111123	**Very good identification**							**63111123**
					GRA	SPO		
Clostridium chauvoei		97.36	0.63	MNS 90	99	99		
Capnocytophaga ochracea		2.55	0.33		1	1		
Actinomyces naeslundii		0.06	0.31		99	1		
63111131	**Good identification**							**63111131**
					GRA	SPO		
Clostridium chauvoei		99.44	0.41	MNS 90	99	99		
Capnocytophaga ochracea		0.56	0.01	MLZ 1	1	1		
63111132	**Good identification**							**63111132**
					GRA	CAT		
Capnocytophaga ochracea		99.90	0.51	MNS 90	1	90		
Clostridium perfringens		0.10	0.03	RAF 90	99	1		
63111141	**Good identification**							**63111141**
					SPO			
Clostridium chauvoei		99.88	0.50	MNS 90	99			
Propionibacterium propionicum		0.12	0.12	SOR 1	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
63111143	**Good identification**							**63111143**
					GRA	SPO	CAT	
Clostridium chauvoei		97.95	0.37	MNS 90	99	99	1	
Capnocytophaga ochracea		1.05	0.01	SOR 1	1	1	90	
Propionibacterium propionicum		1.00	0.15		99	1	1	
63111161	**Good identification**							**63111161**
					GRA	SPO		
Clostridium chauvoei		99.44	0.41	MNS 90	99	99		
Capnocytophaga ochracea		0.56	0.01	SOR 1	1	1		
63111162	**Good identification**							**63111162**
					GRA	CAT		
Capnocytophaga ochracea		98.19	0.51	MNS 90	1	90		
Clostridium perfringens		1.82	0.24	RAF 90	99	1		
63111200	**Very good identification**							**63111200**
Actinomyces odontolyticus		100.0	0.58	ESL 1				
63111202	**Good identification**							**63111202**
Actinomyces odontolyticus		100.0	0.54	ESL 1				
63111220	**Good identification**							**63111220**
Actinomyces odontolyticus		100.0	0.44	ESL 1				
63111222	**Good identification**							**63111222**
					GRA	CAT		
Actinomyces odontolyticus		98.92	0.41	ESL 1	99	10		
Capnocytophaga ochracea		1.08	-.01	NAG 14	1	90		
63111300	**Good identification**							**63111300**
					GRA	SPO	CAT	
Actinomyces odontolyticus		97.38	0.49	ESL 1	99	1	10	
Clostridium chauvoei		2.21	0.15		99	99	1	
Capnocytophaga ochracea		0.41	0.01		1	1	90	
63111301	**Good identification**							**63111301**
					SPO			
Clostridium chauvoei		99.55	0.50	MNS 90	99			
Actinomyces odontolyticus		0.45	0.15	RHA 1	1			
63111302	**Acceptable identification**							**63111302**
					GRA	SPO	CAT	
Actinomyces odontolyticus		93.58	0.46	ESL 1	99	1	10	
Capnocytophaga ochracea		5.81	0.17		1	1	90	
Clostridium chauvoei		0.61	0.03		99	99	1	
63111303	**Good identification**							**63111303**
					SPO			
Clostridium chauvoei		98.46	0.37	MNS 90	99			
Actinomyces odontolyticus		1.54	0.12	RHA 1	1			
63111320	**Low discrimination**							**63111320**
					GRA	CAT		
Actinomyces odontolyticus		73.96	0.36	ESL 1	99	10		
Capnocytophaga ochracea		17.30	0.17	NAG 14	1	90		
Clostridium perfringens		3.57	0.08		99	1		
63111321	**Good identification**							**63111321**
Clostridium chauvoei		100.0	0.41	MNS 90				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
63111501	**Good identification**							**63111501**
Clostridium chauvoei		100.0	0.50	MNS 90				
63111502	**Good identification**							**63111502**
					GRA	SPO	CAT	
Capnocytophaga ochracea		95.83	0.38	MNS 90	1	1	90	
Lactobacillus fermentum		3.14	0.28	RAF 90	99	1	1	
Clostridium chauvoei		0.58	0.03	NAG 90	99	99	1	
63111503	**Good identification**							**63111503**
					GRA	SPO		
Clostridium chauvoei		98.33	0.37	MNS 90	99	99		
Capnocytophaga ochracea		1.67	0.04	ARA 1	1	1		
63111520	**Acceptable identification**							**63111520**
					GRA	CAT		
Capnocytophaga ochracea		94.67	0.38	MNS 90	1	90		
Bifidobacterium longum subsp. longum		1.17	0.14	RAF 90	99	1		
Clostridium perfringens		1.12	0.08	BGL 90	99	1		
63111521	**Good identification**							**63111521**
					GRA	SPO		
Clostridium chauvoei		99.11	0.41	MNS 90	99	99		
Capnocytophaga ochracea		0.89	0.04	ARA 1	1	1		
63111522	**Good identification**							**63111522**
					GRA	CAT		
Capnocytophaga ochracea		99.68	0.54	MNS 90	1	90		
Bifidobacterium longum subsp. longum		0.14	0.14	RAF 90	99	1		
Lactobacillus fermentum		0.12	0.20		99	1		
63111532	**Good identification**							**63111532**
					GRA	CAT		
Capnocytophaga ochracea		97.94	0.38	MNS 90	1	90		
Bifidobacterium longum subsp. longum		2.06	0.18	RAF 90	99	1		
63111562	**Good identification**							**63111562**
Capnocytophaga ochracea		100.0	0.38	MNS 90				
63112100	**Very good identification**							**63112100**
Propionibacterium propionicum		99.68	0.63	ESL 1				
Actinomyces odontolyticus		0.32	0.13					
63112101	**Acceptable identification**							**63112101**
					SPO			
Propionibacterium propionicum		92.17	0.47	ESL 1	1			
Clostridium chauvoei		7.83	0.15	URE 10	99			
63112102	**Very good identification**							**63112102**
Propionibacterium propionicum		99.87	0.66	ESL 1				
Actinomyces odontolyticus		0.13	0.10					
63112103	**Good identification**							**63112103**
					SPO			
Propionibacterium propionicum		99.01	0.50	ESL 1	1			
Clostridium chauvoei		0.99	0.03	URE 10	99			
63112140	**Very good identification**							**63112140**
Propionibacterium propionicum		100.0	0.63	ESL 1				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
63112141	**Good identification**						**63112141**
Propionibacterium propionicum		100.0	0.47	ESL 1			
63112142	**Very good identification**						**63112142**
Propionibacterium propionicum		100.0	0.66	ESL 1			
63112143	**Good identification**						**63112143**
Propionibacterium propionicum		100.0	0.50	ESL 1			
63113100	**Very good identification**						**63113100**
					SPO		
Propionibacterium propionicum		98.25	0.63	ESL 1	1		
Clostridium chauvoei		0.93	0.15		99		
Actinomyces naeslundii		0.44	0.33		1		
63113101	**Low discrimination**						**63113101**
					SPO		
Clostridium chauvoei		88.80	0.50	MNS 90	99		
Propionibacterium propionicum		10.56	0.47	MAN 1	1		
Actinomyces naeslundii		0.63	0.36		1		
63113102	**Very good identification**						**63113102**
					GRA	CAT	
Propionibacterium propionicum		98.47	0.66	ESL 1	99	1	
Capnocytophaga ochracea		1.04	0.17		1	90	
Actinomyces naeslundii		0.29	0.33		99	25	
63113103	**Low discrimination**						**63113103**
					SPO		
Propionibacterium propionicum		49.29	0.50	ESL 1	1		
Clostridium chauvoei		48.75	0.37	URE 10	99		
Actinomyces naeslundii		1.97	0.36		1		
63113121	**Good identification**						**63113121**
					SPO		
Clostridium chauvoei		97.57	0.41	MNS 90	99		
Actinomyces naeslundii		2.08	0.36	MAN 1	1		
Propionibacterium propionicum		0.35	0.12		1		
63113140	**Very good identification**						**63113140**
Propionibacterium propionicum		100.0	0.63	ESL 1			
63113141	**Acceptable identification**						**63113141**
					SPO		
Propionibacterium propionicum		91.29	0.47	ESL 1	1		
Clostridium chauvoei		7.75	0.15	URE 10	99		
Actinomyces naeslundii		0.96	0.23		1		
63113142	**Very good identification**						**63113142**
					GRA	CAT	
Propionibacterium propionicum		99.88	0.66	ESL 1	99	1	
Capnocytophaga ochracea		0.12	0.01		1	90	
63113143	**Good identification**						**63113143**
					SPO		
Propionibacterium propionicum		98.33	0.50	ESL 1	1		
Clostridium chauvoei		0.98	0.03	URE 10	99		
Actinomyces naeslundii		0.69	0.23		1		
63114000	**Good identification**						**63114000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces odontolyticus		96.72	0.51	ESL 1	1		
Actinomyces meyeri		2.04	0.18		1		
Clostridium glycolicum		1.25	0.10		99		
63114002	**Good identification**						**63114002**
Actinomyces odontolyticus		100.0	0.47	ESL 1			
63114020	**Good identification**						**63114020**
Actinomyces odontolyticus		98.27	0.37	ESL 1			
Actinomyces meyeri		1.73	0.04	NAG 14			
63114100	**Low discrimination**						**63114100**
					GRA		
Actinomyces odontolyticus		78.35	0.42	ESL 1	99		
Actinomyces meyeri		20.90	0.29		99		
Fusobacterium mortiferum		0.76	0.13		1		
63114102	**Acceptable identification**						**63114102**
					GRA	CAT	
Actinomyces odontolyticus		94.85	0.39	ESL 1	99	10	
Capnocytophaga ochracea		3.87	0.07		1	90	
Actinomyces meyeri		1.28	0.03		99	1	
63114200	**Good identification**						**63114200**
Actinomyces odontolyticus		100.0	0.56	ESL 1			
63114202	**Good identification**						**63114202**
Actinomyces odontolyticus		100.0	0.52	ESL 1			
63114220	**Good identification**						**63114220**
Actinomyces odontolyticus		100.0	0.42	ESL 1			
63114222	**Good identification**						**63114222**
Actinomyces odontolyticus		100.0	0.39	ESL 1			
63114300	**Good identification**						**63114300**
Actinomyces odontolyticus		100.0	0.47	ESL 1			
63114302	**Good identification**						**63114302**
Actinomyces odontolyticus		100.0	0.43	ESL 1			
63115000	**Acceptable identification**						**63115000**
Actinomyces odontolyticus		93.21	0.48	ESL 1			
Actinomyces meyeri		4.42	0.21				
Eubacterium saburreum		1.46	0.21				
63115002	**Good identification**						**63115002**
					GRA	CAT	
Actinomyces odontolyticus		96.59	0.44	ESL 1	99	10	
Capnocytophaga ochracea		1.87	0.07		1	90	
Lactobacillus fermentum		1.54	0.20		99	1	
63115100	**Low discrimination**						**63115100**
					GRA	CAT	
Actinomyces odontolyticus		43.41	0.39	ESL 1	99	10	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces meyeri		26.05	0.32		99	1	
Capnocytophaga ochracea		17.91	0.24		1	90	
63115101	**Good identification**						**63115101**
Clostridium chauvoei		100.0	0.50	MNS 90			
63115102	**Low discrimination**						**63115102**
					GRA	CAT	
Capnocytophaga ochracea		83.42	0.41	MNS 90	1	90	
Actinomyces odontolyticus		13.77	0.36	RAF 90	99	10	
Lactobacillus fermentum		1.13	0.24	NAG 90	99	1	
63115103	**Good identification**						**63115103**
					GRA	SPO	
Clostridium chauvoei		97.65	0.37	MNS 90	99	99	
Capnocytophaga ochracea		2.35	0.07	XYL 1	1	1	
63115120	**Acceptable identification**						**63115120**
					GRA	CAT	
Capnocytophaga ochracea		91.21	0.41	MNS 90	1	90	
Actinomyces odontolyticus		4.00	0.26	RAF 90	99	10	
Actinomyces meyeri		2.01	0.17	BGL 90	99	1	
63115121	**Good identification**						**63115121**
					GRA	SPO	
Clostridium chauvoei		98.74	0.41	MNS 90	99	99	
Capnocytophaga ochracea		1.26	0.07	XYL 1	1	1	
63115122	**Very good identification**						**63115122**
					GRA	CAT	
Capnocytophaga ochracea		99.26	0.57	MNS 90	1	90	
Bacteroides fragilis		0.35	0.06	RAF 90	1	99	
Actinomyces odontolyticus		0.30	0.22		99	10	
63115132	**Good identification**						**63115132**
Capnocytophaga ochracea		100.0	0.41	MNS 90			
63115162	**Good identification**						**63115162**
Capnocytophaga ochracea		100.0	0.41	MNS 90			
63115200	**Good identification**						**63115200**
Actinomyces odontolyticus		100.0	0.53	ESL 1			
63115202	**Good identification**						**63115202**
Actinomyces odontolyticus		100.0	0.49	ESL 1			
63115220	**Good identification**						**63115220**
Actinomyces odontolyticus		100.0	0.39	ESL 1			
63115222	**Good identification**						**63115222**
Actinomyces odontolyticus		100.0	0.36	ESL 1			
63115300	**Low discrimination**						**63115300**
Actinomyces odontolyticus		67.40	0.44	ESL 1			
Eubacterium contortum		32.60	0.32				
63115302	**Low discrimination**						**63115302**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		85.41	0.40	ESL 1	99	10	
Eubacterium contortum		11.89	0.19		99	1	
Capnocytophaga ochracea		2.69	0.07		1	90	
63115522	**Good identification**						**63115522**
					GRA	CAT	
Capnocytophaga ochracea		98.10	0.44	MNS 90	1	90	
Bifidobacterium longum subsp. longum		1.62	0.22	RAF 90	99	1	
Lactobacillus fermentum		0.28	0.16		99	1	
63121101	**Very good identification**						**63121101**
Clostridium chauvoei		100.0	0.61	MNS 90			
63121103	**Good identification**						**63121103**
Clostridium chauvoei		100.0	0.48	MNS 90			
63121121	**Good identification**						**63121121**
Clostridium chauvoei		100.0	0.53	MNS 90			
63121123	**Good identification**						**63121123**
Clostridium chauvoei		100.0	0.40	MNS 90			
63131101	**Very good identification**						**63131101**
					SPO		
Clostridium chauvoei		99.94	0.67	MNS 90	99		
Actinomyces naeslundii		0.07	0.36	TRE 10	1		
63131103	**Good identification**						**63131103**
					SPO		
Clostridium chauvoei		99.56	0.54	MNS 90	99		
Actinomyces naeslundii		0.37	0.36	TRE 10	1		
Propionibacterium propionicum		0.08	0.14		1		
63131120	**Acceptable identification**						**63131120**
					GRA	CAT	
Clostridium perfringens		90.68	0.47	MNS 99	99	1	
Clostridium chauvoei		5.22	0.25		99	1	
Capnocytophaga ochracea		2.39	0.17		1	90	
63131121	**Very good identification**						**63131121**
					SPO		
Clostridium chauvoei		99.63	0.59	MNS 90	99		
Actinomyces naeslundii		0.19	0.36	TRE 10	1		
Clostridium perfringens		0.18	0.13		84		
63131122	**Low discrimination**						**63131122**
					GRA	CAT	
Clostridium perfringens		64.86	0.41	MNS 99	99	1	
Capnocytophaga ochracea		32.72	0.33		1	90	
Clostridium chauvoei		1.40	0.12		99	1	
63131123	**Good identification**						**63131123**
					SPO		
Clostridium chauvoei		98.22	0.46	MNS 90	99		
Actinomyces naeslundii		1.08	0.36	TRE 10	1		
Clostridium perfringens		0.46	0.07		84		
63132100	**Very good identification**						**63132100**
Propionibacterium propionicum		100.0	0.61	ESL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63132101	**Good identification**					**63132101**
Propionibacterium propionicum		100.0	0.45	ESL 1		
63132102	**Very good identification**					**63132102**
Propionibacterium propionicum		100.0	0.64	ESL 1		
63132103	**Good identification**					**63132103**
Propionibacterium propionicum		100.0	0.48	ESL 1		
63132140	**Very good identification**					**63132140**
Propionibacterium propionicum		100.0	0.61	ESL 1		
63132141	**Good identification**					**63132141**
Propionibacterium propionicum		100.0	0.45	ESL 1		
63132142	**Very good identification**					**63132142**
Propionibacterium propionicum		100.0	0.64	ESL 1		
63132143	**Good identification**					**63132143**
Propionibacterium propionicum		100.0	0.48	ESL 1		
63133100	**Very good identification**					**63133100**
Propionibacterium propionicum		98.65	0.61	ESL 1		
Actinomyces naeslundii		1.04	0.38			
Actinomyces israelii		0.32	0.20			
63133102	**Very good identification**					**63133102**
Propionibacterium propionicum		98.99	0.64	ESL 1		
Actinomyces naeslundii		0.69	0.38			
Actinomyces israelii		0.32	0.23			
63133103	**Low discrimination**					**63133103**
					SPO	
Propionibacterium propionicum		81.38	0.48	ESL 1	1	
Clostridium chauvoei		10.93	0.20	URE 10	99	
Actinomyces naeslundii		7.70	0.41		1	
63133120	**Low discrimination**					**63133120**
Actinomyces naeslundii		37.50	0.38	ESL 12		
Propionibacterium propionicum		36.04	0.27	MNS 88		
Clostridium perfringens		26.46	0.13	RAF 92		
63133123	**Low discrimination**					**63133123**
					SPO	
Actinomyces naeslundii		63.29	0.41	ESL 12	1	
Clostridium chauvoei		29.96	0.12	MNS 88	99	
Propionibacterium propionicum		6.76	0.14	RAF 92	1	
63133140	**Very good identification**					**63133140**
Propionibacterium propionicum		99.82	0.61	ESL 1		
Actinomyces naeslundii		0.19	0.25			
63133141	**Good identification**					**63133141**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		97.56	0.45	ESL 1			
Actinomyces naeslundii		2.44	0.28	URE 10			
63133142	**Very good identification**						**63133142**
Propionibacterium propionicum		99.75	0.64	ESL 1			
Actinomyces israelii		0.12	0.16				
Actinomyces naeslundii		0.12	0.25				
63133143	**Good identification**						**63133143**
Propionibacterium propionicum		98.36	0.48	ESL 1			
Actinomyces naeslundii		1.64	0.28	URE 10			
63137100	**Low discrimination**						**63137100**
Actinomyces israelii		78.51	0.38	SAL 92			
Propionibacterium propionicum		21.50	0.27	CEL 93			
63137102	**Low discrimination**						**63137102**
Actinomyces israelii		78.51	0.41	SAL 92			
Propionibacterium propionicum		21.50	0.30	CEL 93			
63137500	**Good identification**						**63137500**
Actinomyces israelii		100.0	0.36	SAL 92			
63137502	**Good identification**						**63137502**
Actinomyces israelii		96.20	0.39	SAL 92			
Bifidobacterium adolescentis		3.81	0.09	CEL 93			
63141101	**Good identification**						**63141101**
Clostridium chauvoei		100.0	0.43	MNS 90			
63141121	**Good identification**						**63141121**
Clostridium chauvoei		100.0	0.35	MNS 90			
63151101	**Good identification**						**63151101**
					SPO		
Clostridium chauvoei		99.52	0.50	MNS 90	99		
Actinomyces naeslundii		0.48	0.33	CEL 1	1		
63151102	**Good identification**						**63151102**
					GRA		
Capnocytophaga ochracea		96.59	0.41	MNS 90	1		
Actinomyces viscosus		1.04	0.08	RAF 90	99		
Actinomyces naeslundii		0.75	0.30	NAG 90	99		
63151103	**Acceptable identification**						**63151103**
					GRA	SPO	
Clostridium chauvoei		94.33	0.37	MNS 90	99	99	
Actinomyces naeslundii		2.60	0.33	CEL 1	99	1	
Capnocytophaga ochracea		2.27	0.07		1	1	
63151120	**Acceptable identification**						**63151120**
					GRA	CAT	
Capnocytophaga ochracea		90.73	0.41	MNS 90	1	90	
Prevotella melaninogenica		4.07	0.14	RAF 90	1	1	
Clostridium perfringens		3.12	0.19	BGL 90	99	1	
63151121	**Good identification**						**63151121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Clostridium chauvoei		97.34	0.41	MNS 90	99	99	
Actinomyces naeslundii		1.42	0.33	CEL 1	99	1	
Capnocytophaga ochracea		1.24	0.07		1	1	
63151122	**Very good identification**						**63151122**
					GRA	CAT	
Capnocytophaga ochracea		99.40	0.57	MNS 90	1	90	
Clostridium perfringens		0.18	0.13	RAF 90	99	1	
Prevotella oralis		0.17	0.09		1	1	
63151132	**Good identification**						**63151132**
Capnocytophaga ochracea		100.0	0.41	MNS 90			
63151162	**Good identification**						**63151162**
					GRA	CAT	
Capnocytophaga ochracea		99.69	0.41	MNS 90	1	90	
Clostridium perfringens		0.31	0.01	RAF 90	99	1	
63151522	**Good identification**						**63151522**
Capnocytophaga ochracea		100.0	0.44	MNS 90			
63153103	**Low discrimination**						**63153103**
					SPO		
Actinomyces naeslundii		72.45	0.38	ESL 12	1		
Propionibacterium propionicum		13.85	0.15	MNS 88	1		
Clostridium chauvoei		13.70	0.03	RAF 92	99		
63153121	**Low discrimination**						**63153121**
					SPO		
Actinomyces naeslundii		73.69	0.38	ESL 12	1		
Clostridium chauvoei		26.31	0.07	MNS 88	99		
63153123	**Good identification**						**63153123**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63155122	**Good identification**						**63155122**
					GRA	CAT	
Capnocytophaga ochracea		99.20	0.47	MNS 90	1	90	
Bacteroides fragilis		0.60	-.01	RAF 90	1	99	
Actinomyces israelii		0.20	0.17		99	1	
63155502	**Low discrimination**						**63155502**
					GRA	CAT	
Actinomyces israelii		50.09	0.36	SAL 92	99	1	
Capnocytophaga ochracea		37.53	0.18		1	90	
Bifidobacterium adolescentis		12.39	0.19		99	1	
63157000	**Good identification**						**63157000**
Actinomyces israelii		100.0	0.37	SAL 92			
63157002	**Good identification**						**63157002**
Actinomyces israelii		100.0	0.40	SAL 92			
63157100	**Good identification**						**63157100**
Actinomyces israelii		100.0	0.47	SAL 92			
63157102	**Good identification**						**63157102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.50	SAL 92		
63157112	**Good identification**					**63157112**
Actinomyces israelii		100.0	0.36	SAL 92		
63157140	**Good identification**					**63157140**
Actinomyces israelii		100.0	0.40	SAL 92		
63157142	**Good identification**					**63157142**
Actinomyces israelii		100.0	0.43	SAL 92		
63157400	**Good identification**					**63157400**
Actinomyces israelii		100.0	0.35	SAL 92		
63157402	**Good identification**					**63157402**
Actinomyces israelii		100.0	0.38	SAL 92		
63157500	**Good identification**					**63157500**
Actinomyces israelii		100.0	0.46	SAL 92		
63157502	**Good identification**					**63157502**
Actinomyces israelii		96.55	0.49	SAL 92		
Bifidobacterium adolescentis		3.45	0.18			
63157540	**Good identification**					**63157540**
Actinomyces israelii		100.0	0.39	SAL 92		
63157542	**Good identification**					**63157542**
Actinomyces israelii		100.0	0.42	SAL 92		
63171123	**Low discrimination**					**63171123**
					SPO	
Actinomyces naeslundii		59.06	0.38	ESL 12	1	
Clostridium chauvoei		40.94	0.12	MNS 88	99	
63173001	**Good identification**					**63173001**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63173003	**Good identification**					**63173003**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63173021	**Good identification**					**63173021**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63173023	**Good identification**					**63173023**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63173100	**Low discrimination**					**63173100**
Actinomyces israelii		63.95	0.39	SAL 92		
Actinomyces naeslundii		20.90	0.40	XYL 92		
Propionibacterium propionicum		15.16	0.27			
63173101	**Good identification**					**63173101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ESL 12		
63173102	**Low discrimination**					**63173102**
Actinomyces israelii		68.73	0.42	SAL 92		
Propionibacterium propionicum		16.29	0.30	XYL 92		
Actinomyces naeslundii		14.98	0.40			
63173103	**Acceptable identification**					**63173103**
Actinomyces naeslundii		92.54	0.43	ESL 12		
Propionibacterium propionicum		7.46	0.14	MNS 88		
63173120	**Low discrimination**					**63173120**
Actinomyces naeslundii		83.66	0.40	ESL 12		
Actinomyces israelii		16.34	0.18	MNS 88		
63173121	**Good identification**					**63173121**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63173122	**Low discrimination**					**63173122**
Actinomyces naeslundii		77.35	0.40	ESL 12		
Actinomyces israelii		22.66	0.22	MNS 88		
63173123	**Good identification**					**63173123**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63173500	**Good identification**					**63173500**
Actinomyces israelii		100.0	0.37	SAL 92		
63173502	**Good identification**					**63173502**
Actinomyces israelii		100.0	0.40	SAL 92		
63175002	**Good identification**					**63175002**
Actinomyces israelii		100.0	0.37	SAL 92		
63175100	**Low discrimination**					**63175100**
					COC	
Actinomyces israelii		52.61	0.44	SAL 92	1	
Atopobium parvulum		47.39	0.30		99	
63175102	**Acceptable identification**					**63175102**
					COC	
Actinomyces israelii		90.42	0.47	SAL 92	1	
Atopobium parvulum		9.58	0.17		99	
63175140	**Good identification**					**63175140**
Actinomyces israelii		100.0	0.37	SAL 92		
63175142	**Good identification**					**63175142**
Actinomyces israelii		100.0	0.40	SAL 92		
63175402	**Good identification**					**63175402**
Actinomyces israelii		100.0	0.35	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63175500	**Good identification**					**63175500**
Actinomyces israelii		100.0	0.43	SAL 92		
63175502	**Good identification**					**63175502**
Actinomyces israelii		95.80	0.46	SAL 92		
Bifidobacterium adolescentis		4.20	0.16			
63175540	**Good identification**					**63175540**
Actinomyces israelii		100.0	0.36	SAL 92		
63175542	**Good identification**					**63175542**
Actinomyces israelii		100.0	0.39	SAL 92		
63177000	**Good identification**					**63177000**
Actinomyces israelii		100.0	0.47	SAL 92		
63177002	**Good identification**					**63177002**
Actinomyces israelii		100.0	0.50	SAL 92		
63177012	**Good identification**					**63177012**
Actinomyces israelii		100.0	0.36	SAL 92		
63177040	**Good identification**					**63177040**
Actinomyces israelii		100.0	0.40	SAL 92		
63177042	**Good identification**					**63177042**
Actinomyces israelii		100.0	0.43	SAL 92		
63177100	**Very good identification**					**63177100**
Actinomyces israelii		100.0	0.57	SAL 92		
63177102	**Very good identification**					**63177102**
Actinomyces israelii		100.0	0.60	SAL 92		
63177110	**Good identification**					**63177110**
Actinomyces israelii		100.0	0.43	SAL 92		
63177112	**Good identification**					**63177112**
Actinomyces israelii		100.0	0.46	SAL 92		
63177120	**Good identification**					**63177120**
Actinomyces israelii		100.0	0.37	SAL 92		
63177122	**Good identification**					**63177122**
Actinomyces israelii		100.0	0.40	SAL 92		
63177140	**Good identification**					**63177140**
Actinomyces israelii		100.0	0.50	SAL 92		
63177142	**Good identification**					**63177142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.53	SAL 92		
63177150	**Good identification**					**63177150**
Actinomyces israelii		100.0	0.36	SAL 92		
63177152	**Good identification**					**63177152**
Actinomyces israelii		100.0	0.39	SAL 92		
63177300	**Good identification**					**63177300**
Actinomyces israelii		100.0	0.38	SAL 92		
63177302	**Good identification**					**63177302**
Actinomyces israelii		100.0	0.41	SAL 92		
63177400	**Good identification**					**63177400**
Actinomyces israelii		100.0	0.45	SAL 92		
63177402	**Good identification**					**63177402**
Actinomyces israelii		100.0	0.48	SAL 92		
63177440	**Good identification**					**63177440**
Actinomyces israelii		100.0	0.38	SAL 92		
63177442	**Good identification**					**63177442**
Actinomyces israelii		100.0	0.41	SAL 92		
63177500	**Good identification**					**63177500**
Actinomyces israelii		100.0	0.56	SAL 92		
63177502	**Very good identification**					**63177502**
Actinomyces israelii		99.37	0.59	SAL 92		
Bifidobacterium adolescentis		0.63	0.15			
63177510	**Good identification**					**63177510**
Actinomyces israelii		100.0	0.41	SAL 92		
63177512	**Good identification**					**63177512**
Actinomyces israelii		95.08	0.45	SAL 92		
Bifidobacterium adolescentis		4.92	0.16	MLZ 13		
63177520	**Good identification**					**63177520**
Actinomyces israelii		100.0	0.35	SAL 92		
63177522	**Good identification**					**63177522**
Actinomyces israelii		100.0	0.38	SAL 92		
63177540	**Good identification**					**63177540**
Actinomyces israelii		100.0	0.49	SAL 92		
63177542	**Good identification**					**63177542**
Actinomyces israelii		100.0	0.52	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63177552	**Good identification**						**63177552**
Actinomyces israelii		100.0	0.37	SAL 92			
63177700	**Good identification**						**63177700**
Actinomyces israelii		100.0	0.36	SAL 92			
63177702	**Good identification**						**63177702**
Actinomyces israelii		100.0	0.39	SAL 92			
63200000	**Very good identification**						**63200000**
					GRA	SPO	
Eubacterium tortuosum		99.92	0.64	GAL 99	99	1	
Clostridium botulinum biovar A		0.06	0.13		99	99	
Fusobacterium mortiferum		0.01	0.13		1	1	
63200002	**Good identification**						**63200002**
Eubacterium tortuosum		100.0	0.52	GAL 99			
63200020	**Good identification**						**63200020**
Eubacterium tortuosum		100.0	0.52	GAL 99			
63200022	**Good identification**						**63200022**
Eubacterium tortuosum		100.0	0.39	GAL 99			
63201000	**Excellent identification**						**63201000**
					GRA		
Eubacterium tortuosum		100.0	0.99		99		
Fusobacterium mortiferum		0.00	0.16		1		
Actinomyces odontolyticus		0.00	0.06		99		
63201001	**Very good identification**						**63201001**
Eubacterium tortuosum		100.0	0.64	URE 1			
63201002	**Very good identification**						**63201002**
Eubacterium tortuosum		100.0	0.86				
63201003	**Good identification**						**63201003**
Eubacterium tortuosum		100.0	0.52	URE 1			
63201010	**Very good identification**						**63201010**
Eubacterium tortuosum		100.0	0.64	MLZ 1			
63201012	**Good identification**						**63201012**
Eubacterium tortuosum		100.0	0.52	MLZ 1			
63201020	**Very good identification**						**63201020**
Eubacterium tortuosum		100.0	0.86				
63201021	**Good identification**						**63201021**
Eubacterium tortuosum		100.0	0.52	URE 1			
63201022	**Very good identification**						**63201022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.73			
63201023	**Good identification**					**63201023**
Eubacterium tortuosum		100.0	0.39	URE 1		
63201030	**Good identification**					**63201030**
Eubacterium tortuosum		100.0	0.52	MLZ 1		
63201032	**Good identification**					**63201032**
Eubacterium tortuosum		100.0	0.39	MLZ 1		
63201040	**Very good identification**					**63201040**
Eubacterium tortuosum		100.0	0.64	SOR 1		
63201042	**Good identification**					**63201042**
Eubacterium tortuosum		100.0	0.52	SOR 1		
63201060	**Good identification**					**63201060**
Eubacterium tortuosum		100.0	0.52	SOR 1		
63201062	**Good identification**					**63201062**
Eubacterium tortuosum		100.0	0.39	SOR 1		
63201100	**Very good identification**					**63201100**
					GRA	
Eubacterium tortuosum		99.97	0.64	LAC 1	99	
Fusobacterium mortiferum		0.04	0.22		1	
63201102	**Good identification**					**63201102**
Eubacterium tortuosum		100.0	0.52	LAC 1		
63201120	**Good identification**					**63201120**
Eubacterium tortuosum		100.0	0.52	LAC 1		
63201122	**Good identification**					**63201122**
Eubacterium tortuosum		100.0	0.39	LAC 1		
63201200	**Very good identification**					**63201200**
Eubacterium tortuosum		99.98	0.64	RHA 1		
Actinomyces odontolyticus		0.02	0.11			
63201202	**Good identification**					**63201202**
Eubacterium tortuosum		99.94	0.52	RHA 1		
Actinomyces odontolyticus		0.06	0.07			
63201220	**Good identification**					**63201220**
Eubacterium tortuosum		100.0	0.52	RHA 1		
63201222	**Good identification**					**63201222**
Eubacterium tortuosum		100.0	0.39	RHA 1		
63201400	**Very good identification**					**63201400**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.64	ARA 1		
63201402	**Good identification**					**63201402**
Eubacterium tortuosum		100.0	0.52	ARA 1		
63201420	**Good identification**					**63201420**
Eubacterium tortuosum		100.0	0.52	ARA 1		
63201422	**Good identification**					**63201422**
Eubacterium tortuosum		100.0	0.39	ARA 1		
63203000	**Very good identification**					**63203000**
Eubacterium tortuosum		100.0	0.64	MAN 1		
63203002	**Good identification**					**63203002**
Eubacterium tortuosum		100.0	0.52	MAN 1		
63203020	**Good identification**					**63203020**
Eubacterium tortuosum		100.0	0.52	MAN 1		
63203022	**Good identification**					**63203022**
Eubacterium tortuosum		100.0	0.39	MAN 1		
63205000	**Very good identification**					**63205000**
Eubacterium tortuosum		100.0	0.64	XYL 1		
63205002	**Good identification**					**63205002**
Eubacterium tortuosum		100.0	0.52	XYL 1		
63205020	**Good identification**					**63205020**
Eubacterium tortuosum		100.0	0.52	XYL 1		
63205022	**Good identification**					**63205022**
Eubacterium tortuosum		100.0	0.39	XYL 1		
63210100	**Good identification**					**63210100**
					GRA	
Fusobacterium mortiferum		97.63	0.53	NIT 1	1	
Actinomyces odontolyticus		1.93	0.18		99	
Eubacterium tortuosum		0.44	-.04		99	
63210102	**Acceptable identification**					**63210102**
					GRA	
Fusobacterium mortiferum		90.15	0.37	NIT 1	1	
Actinomyces odontolyticus		9.85	0.15	BGL 10	99	
63211000	**Very good identification**					**63211000**
					GRA	
Eubacterium tortuosum		98.46	0.64	FRU 1	99	
Fusobacterium mortiferum		1.45	0.50		1	
Actinomyces odontolyticus		0.09	0.24		99	
63211002	**Good identification**					**63211002**
					GRA	
Eubacterium tortuosum		98.76	0.52	FRU 1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Fusobacterium mortiferum		0.92	0.33		1	
Actinomyces odontolyticus		0.33	0.20		99	
63211020	**Good identification**					**63211020**
					GRA	
Eubacterium tortuosum		99.83	0.52	FRU 1	99	
Actinomyces odontolyticus		0.09	0.10		99	
Fusobacterium mortiferum		0.08	0.16		1	
63211022	**Good identification**					**63211022**
Eubacterium tortuosum		99.69	0.39	FRU 1		
Actinomyces odontolyticus		0.31	0.07			
63211100	**Low discrimination**					**63211100**
					GRA	
Fusobacterium mortiferum		76.76	0.56	NIT 1	1	
Eubacterium tortuosum		22.56	0.30		99	
Actinomyces odontolyticus		0.68	0.15		99	
63211102	**Low discrimination**					**63211102**
					GRA	
Fusobacterium mortiferum		65.99	0.40	NIT 1	1	
Eubacterium tortuosum		30.80	0.17	BGL 10	99	
Actinomyces odontolyticus		3.20	0.12		99	
63215100	**Acceptable identification**					**63215100**
					GRA	
Fusobacterium mortiferum		91.16	0.37	NIT 1	1	
Actinomyces odontolyticus		5.25	0.10	XYL 7	99	
Eubacterium tortuosum		3.59	-.04		99	
63221000	**Very good identification**					**63221000**
Eubacterium tortuosum		100.0	0.64	TRE 1		
63221002	**Good identification**					**63221002**
Eubacterium tortuosum		100.0	0.52	TRE 1		
63221020	**Good identification**					**63221020**
Eubacterium tortuosum		100.0	0.52	TRE 1		
63221022	**Good identification**					**63221022**
Eubacterium tortuosum		100.0	0.39	TRE 1		
63230000	**Good identification**					**63230000**
					GRA	
Fusobacterium mortiferum		96.64	0.38	NIT 1	1	
Eubacterium tortuosum		3.36	-.04		99	
63230100	**Good identification**					**63230100**
Fusobacterium mortiferum		100.0	0.44	NIT 1		
63231100	**Good identification**					**63231100**
					GRA	
Fusobacterium mortiferum		99.02	0.47	NIT 1	1	
Eubacterium tortuosum		0.98	-.04		99	
63240000	**Good identification**					**63240000**
Eubacterium tortuosum		100.0	0.48	CEL 10		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63240002	**Good identification**					**63240002**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63240020	**Good identification**					**63240020**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241000	**Very good identification**					**63241000**
Eubacterium tortuosum		100.0	0.82	CEL 10		
63241001	**Good identification**					**63241001**
Eubacterium tortuosum		100.0	0.48	CEL 10		
63241002	**Very good identification**					**63241002**
Eubacterium tortuosum		100.0	0.69	CEL 10		
63241003	**Good identification**					**63241003**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241010	**Good identification**					**63241010**
Eubacterium tortuosum		100.0	0.48	CEL 10		
63241012	**Good identification**					**63241012**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241020	**Very good identification**					**63241020**
Eubacterium tortuosum		100.0	0.69	CEL 10		
63241021	**Good identification**					**63241021**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241022	**Very good identification**					**63241022**
Eubacterium tortuosum		100.0	0.56	CEL 10		
63241030	**Good identification**					**63241030**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241040	**Good identification**					**63241040**
Eubacterium tortuosum		100.0	0.48	CEL 10		
63241042	**Good identification**					**63241042**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241060	**Good identification**					**63241060**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241100	**Good identification**					**63241100**
					GRA	
Eubacterium tortuosum		99.90	0.48	CEL 10	99	
Fusobacterium mortiferum		0.10	0.14	LAC 1	1	
63241102	**Good identification**					**63241102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241120	**Good identification**					**63241120**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241200	**Good identification**					**63241200**
Eubacterium tortuosum		100.0	0.48	CEL 10		
63241202	**Good identification**					**63241202**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241220	**Good identification**					**63241220**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241400	**Good identification**					**63241400**
Eubacterium tortuosum		100.0	0.48	CEL 10		
63241402	**Good identification**					**63241402**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63241420	**Good identification**					**63241420**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63243000	**Good identification**					**63243000**
					SPO	
Eubacterium tortuosum		99.91	0.48	CEL 10	1	
Clostridium innocuum		0.09	0.00	MAN 1	99	
63243002	**Good identification**					**63243002**
					SPO	
Eubacterium tortuosum		99.50	0.35	CEL 10	1	
Clostridium innocuum		0.50	0.00	MAN 1	99	
63243020	**Good identification**					**63243020**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63245000	**Good identification**					**63245000**
Eubacterium tortuosum		100.0	0.48	CEL 10		
63245002	**Good identification**					**63245002**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63245020	**Good identification**					**63245020**
Eubacterium tortuosum		100.0	0.35	CEL 10		
63250000	**Low discrimination**					**63250000**
					GRA	
Fusobacterium mortiferum		74.47	0.39	NIT 1	1	
Eubacterium tortuosum		25.53	0.14		99	
63250100	**Good identification**					**63250100**
Fusobacterium mortiferum		100.0	0.45	NIT 1		
63251000	**Good identification**					**63251000**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		95.39	0.48	FRU 1	99	
Fusobacterium mortiferum		4.22	0.42	CEL 10	1	
Lactobacillus catenaformis		0.32	0.12		99	
63251002	**Acceptable identification**					**63251002**
					GRA	
Eubacterium tortuosum		94.50	0.35	FRU 1	99	
Fusobacterium mortiferum		2.63	0.25	CEL 10	1	
Lactobacillus catenaformis		1.78	0.12		99	
63251020	**Good identification**					**63251020**
					SPO	
Eubacterium tortuosum		98.39	0.35	FRU 1	1	
Clostridium barati		1.15	0.04	CEL 10	99	
Lactobacillus catenaformis		0.46	0.02		1	
63251100	**Low discrimination**					**63251100**
					GRA	
Fusobacterium mortiferum		83.47	0.48	NIT 1	1	
Eubacterium tortuosum		8.18	0.14		99	
Lactobacillus catenaformis		4.02	0.15		99	
63261000	**Good identification**					**63261000**
Eubacterium tortuosum		100.0	0.48	TRE 1		
63261002	**Good identification**					**63261002**
Eubacterium tortuosum		100.0	0.35	TRE 1		
63261020	**Good identification**					**63261020**
Eubacterium tortuosum		100.0	0.35	TRE 1		
63270100	**Good identification**					**63270100**
Fusobacterium mortiferum		100.0	0.36	NIT 1		
63271100	**Good identification**					**63271100**
					GRA	
Fusobacterium mortiferum		97.28	0.39	NIT 1	1	
Actinomyces naeslundii		2.72	0.24		99	
63277100	**Good identification**					**63277100**
Actinomyces israelii		100.0	0.41	MLT 99		
63277102	**Good identification**					**63277102**
Actinomyces israelii		100.0	0.44	MLT 99		
63277142	**Good identification**					**63277142**
Actinomyces israelii		100.0	0.37	MLT 99		
63277500	**Good identification**					**63277500**
Actinomyces israelii		100.0	0.40	MLT 99		
63277502	**Good identification**					**63277502**
Actinomyces israelii		100.0	0.43	MLT 99		
63277542	**Good identification**					**63277542**
Actinomyces israelii		100.0	0.36	MLT 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63300000	**Low discrimination**					**63300000**
					SPO	
Actinomyces odontolyticus		53.48	0.43	ESL 1	1	
Eubacterium tortuosum		43.67	0.30	FRU 92	1	
Clostridium botulinum biovar A		2.04	0.11		99	
63300002	**Low discrimination**					**63300002**
Actinomyces odontolyticus		80.96	0.39	ESL 1		
Eubacterium tortuosum		19.04	0.17	FRU 92		
63300200	**Good identification**					**63300200**
Actinomyces odontolyticus		99.58	0.48	ESL 1		
Eubacterium tortuosum		0.42	-.04	FRU 92		
63300202	**Good identification**					**63300202**
Actinomyces odontolyticus		100.0	0.44	ESL 1		
63300300	**Good identification**					**63300300**
Actinomyces odontolyticus		100.0	0.39	ESL 1		
63300302	**Good identification**					**63300302**
Actinomyces odontolyticus		100.0	0.36	ESL 1		
63301000	**Very good identification**					**63301000**
Eubacterium tortuosum		99.18	0.64	MLT 1		
Actinomyces odontolyticus		0.82	0.40			
63301002	**Good identification**					**63301002**
Eubacterium tortuosum		97.22	0.52	MLT 1		
Actinomyces odontolyticus		2.78	0.36			
63301020	**Good identification**					**63301020**
Eubacterium tortuosum		99.25	0.52	MLT 1		
Actinomyces odontolyticus		0.76	0.26			
63301022	**Good identification**					**63301022**
Eubacterium tortuosum		97.43	0.39	MLT 1		
Actinomyces odontolyticus		2.57	0.23			
63301101	**Good identification**					**63301101**
					SPO	
Clostridium chauvoei		99.65	0.43	MNS 90	99	
Eubacterium tortuosum		0.35	-.04	SAL 1	1	
63301121	**Good identification**					**63301121**
Clostridium chauvoei		100.0	0.35	MNS 90		
63301200	**Low discrimination**					**63301200**
Actinomyces odontolyticus		61.31	0.45	ESL 1		
Eubacterium tortuosum		38.69	0.30	FRU 92		
63301202	**Low discrimination**					**63301202**
Actinomyces odontolyticus		84.62	0.41	ESL 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		15.38	0.17	FRU 92		
63301300	**Good identification**					**63301300**
Actinomyces odontolyticus		98.02	0.36	ESL 1		
Eubacterium tortuosum		1.98	-.04	FRU 92		
63304000	**Good identification**					**63304000**
Actinomyces odontolyticus		98.35	0.38	ESL 1		
Eubacterium tortuosum		1.65	-.04	FRU 92		
63304200	**Good identification**					**63304200**
Actinomyces odontolyticus		100.0	0.42	ESL 1		
63304202	**Good identification**					**63304202**
Actinomyces odontolyticus		100.0	0.39	ESL 1		
63305200	**Good identification**					**63305200**
Actinomyces odontolyticus		98.72	0.39	ESL 1		
Eubacterium tortuosum		1.28	-.04	FRU 92		
63305202	**Good identification**					**63305202**
Actinomyces odontolyticus		100.0	0.36	ESL 1		
63310000	**Very good identification**					**63310000**
					GRA	
Actinomyces odontolyticus		99.08	0.61	ESL 1	99	
Fusobacterium mortiferum		0.85	0.31		1	
Eubacterium tortuosum		0.07	-.04		99	
63310002	**Very good identification**					**63310002**
					GRA	
Actinomyces odontolyticus		99.53	0.57	ESL 1	99	
Actinomyces viscosus		0.32	0.08		99	
Fusobacterium mortiferum		0.15	0.15		1	
63310020	**Good identification**					**63310020**
Actinomyces odontolyticus		100.0	0.47	ESL 1		
63310022	**Good identification**					**63310022**
Actinomyces odontolyticus		100.0	0.44	ESL 1		
63310100	**Acceptable identification**					**63310100**
					GRA	
Actinomyces odontolyticus		94.07	0.52	ESL 1	99	
Fusobacterium mortiferum		5.93	0.37		1	
63310102	**Good identification**					**63310102**
					GRA	
Actinomyces odontolyticus		97.14	0.49	ESL 1	99	
Actinomyces viscosus		1.75	0.12		99	
Fusobacterium mortiferum		1.11	0.21		1	
63310120	**Good identification**					**63310120**
Actinomyces odontolyticus		100.0	0.39	ESL 1		
63310122	**Good identification**					**63310122**
					GRA CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces odontolyticus		97.74	0.35	ESL 1	99	10	
Capnocytophaga ochracea		2.26	-.01	NAG 14	1	90	
63310200	**Very good identification**						**63310200**
Actinomyces odontolyticus		100.0	0.66	ESL 1			
63310202	**Very good identification**						**63310202**
Actinomyces odontolyticus		100.0	0.62	ESL 1			
63310220	**Good identification**						**63310220**
Actinomyces odontolyticus		100.0	0.52	ESL 1			
63310222	**Good identification**						**63310222**
Actinomyces odontolyticus		100.0	0.49	ESL 1			
63310300	**Very good identification**						**63310300**
Actinomyces odontolyticus		100.0	0.57	ESL 1			
63310302	**Good identification**						**63310302**
Actinomyces odontolyticus		100.0	0.54	ESL 1			
63310320	**Good identification**						**63310320**
Actinomyces odontolyticus		100.0	0.44	ESL 1			
63310322	**Good identification**						**63310322**
Actinomyces odontolyticus		100.0	0.40	ESL 1			
63311000	**Low discrimination**						**63311000**
					GRA		
Actinomyces odontolyticus		88.59	0.58	ESL 1	99		
Eubacterium tortuosum		9.44	0.30		99		
Fusobacterium mortiferum		1.70	0.34		1		
63311002	**Good identification**						**63311002**
Actinomyces odontolyticus		96.10	0.54	ESL 1			
Eubacterium tortuosum		2.95	0.17				
Actinomyces naeslundii		0.34	0.30				
63311020	**Low discrimination**						**63311020**
Actinomyces odontolyticus		84.74	0.44	ESL 1			
Eubacterium tortuosum		9.79	0.17	NAG 14			
Clostridium perfringens		4.36	0.17				
63311022	**Acceptable identification**						**63311022**
Actinomyces odontolyticus		90.48	0.41	ESL 1			
Clostridium perfringens		3.57	0.12	NAG 14			
Eubacterium tortuosum		3.01	0.04				
63311100	**Low discrimination**						**63311100**
					GRA	SPO	
Actinomyces odontolyticus		82.43	0.49	ESL 1	99	1	
Fusobacterium mortiferum		11.69	0.40		1	1	
Clostridium chauvoei		1.87	0.15		99	99	
63311101	**Good identification**						**63311101**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium chauvoei		98.28	0.50	MNS 90	99		
Actinomyces naeslundii		1.28	0.40	SAL 1	1		
Actinomyces odontolyticus		0.44	0.15		1		
63311102	**Low discrimination**						**63311102**
					GRA	CAT	
Actinomyces odontolyticus		84.89	0.46	ESL 1	99	10	
Capnocytophaga ochracea		5.27	0.17		1	90	
Actinomyces naeslundii		2.69	0.37		99	25	
63311103	**Acceptable identification**						**63311103**
					SPO		
Clostridium chauvoei		91.09	0.37	MNS 90	99		
Actinomyces naeslundii		6.71	0.40	SAL 1	1		
Actinomyces odontolyticus		1.42	0.12		1		
63311121	**Good identification**						**63311121**
					SPO		
Clostridium chauvoei		96.25	0.41	MNS 90	99		
Actinomyces naeslundii		3.75	0.40	SAL 1	1		
63311200	**Very good identification**						**63311200**
Actinomyces odontolyticus		99.95	0.63	ESL 1			
Eubacterium tortuosum		0.06	-.04				
63311202	**Very good identification**						**63311202**
Actinomyces odontolyticus		100.0	0.59	ESL 1			
63311220	**Good identification**						**63311220**
Actinomyces odontolyticus		100.0	0.49	ESL 1			
63311222	**Good identification**						**63311222**
Actinomyces odontolyticus		100.0	0.46	ESL 1			
63311300	**Low discrimination**						**63311300**
Actinomyces odontolyticus		89.07	0.54	ESL 1			
Eubacterium contortum		10.93	0.32				
63311302	**Good identification**						**63311302**
Actinomyces odontolyticus		96.59	0.51	ESL 1			
Eubacterium contortum		3.41	0.19				
63311320	**Low discrimination**						**63311320**
Actinomyces odontolyticus		88.26	0.41	ESL 1			
Eubacterium contortum		11.74	0.19	NAG 14			
63311322	**Acceptable identification**						**63311322**
					GRA	CAT	
Actinomyces odontolyticus		93.29	0.37	ESL 1	99	10	
Eubacterium contortum		3.57	0.06	NAG 14	99	1	
Capnocytophaga ochracea		1.67	-.01		1	90	
63313001	**Good identification**						**63313001**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63313003	**Good identification**						**63313003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ESL 12		
63313021	**Good identification**					**63313021**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63313023	**Good identification**					**63313023**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63313100	**Low discrimination**					**63313100**
Actinomyces naeslundii		41.29	0.42	ESL 12		
Propionibacterium propionicum		26.56	0.29	MNS 88		
Actinomyces israelii		21.10	0.28	RAF 92		
63313101	**Low discrimination**					**63313101**
					SPO	
Actinomyces naeslundii		69.05	0.45	ESL 12	1	
Clostridium chauvoei		27.66	0.15	MNS 88	99	
Propionibacterium propionicum		3.29	0.12	RAF 92	1	
63313102	**Low discrimination**					**63313102**
Actinomyces naeslundii		34.54	0.42	ESL 12		
Propionibacterium propionicum		33.32	0.32	MNS 88		
Actinomyces israelii		26.47	0.31	RAF 92		
63313103	**Low discrimination**					**63313103**
					SPO	
Actinomyces naeslundii		87.55	0.45	ESL 12	1	
Propionibacterium propionicum		6.26	0.15	MNS 88	1	
Clostridium chauvoei		6.19	0.03	RAF 92	99	
63313120	**Good identification**					**63313120**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63313121	**Low discrimination**					**63313121**
					SPO	
Actinomyces naeslundii		88.22	0.45	ESL 12	1	
Clostridium chauvoei		11.78	0.07	MNS 88	99	
63313122	**Acceptable identification**					**63313122**
					GRA	
Actinomyces naeslundii		91.60	0.42	ESL 12	99	
Capnocytophaga ochracea		8.40	-.01	MNS 88	1	
63313123	**Good identification**					**63313123**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63314000	**Good identification**					**63314000**
					GRA	
Actinomyces odontolyticus		99.87	0.56	ESL 1	99	
Fusobacterium mortiferum		0.13	0.12		1	
63314002	**Good identification**					**63314002**
Actinomyces odontolyticus		100.0	0.52	ESL 1		
63314020	**Good identification**					**63314020**
Actinomyces odontolyticus		100.0	0.42	ESL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63314022	**Good identification**					**63314022**
Actinomyces odontolyticus		100.0	0.39	ESL 1		
63314100	**Good identification**					**63314100**
					GRA	
Actinomyces odontolyticus		99.05	0.47	ESL 1	99	
Fusobacterium mortiferum		0.95	0.18		1	
63314102	**Good identification**					**63314102**
Actinomyces odontolyticus		100.0	0.43	ESL 1		
63314200	**Very good identification**					**63314200**
Actinomyces odontolyticus		100.0	0.61	ESL 1		
63314202	**Very good identification**					**63314202**
Actinomyces odontolyticus		100.0	0.57	ESL 1		
63314220	**Good identification**					**63314220**
Actinomyces odontolyticus		100.0	0.47	ESL 1		
63314222	**Good identification**					**63314222**
Actinomyces odontolyticus		100.0	0.43	ESL 1		
63314300	**Low discrimination**					**63314300**
Actinomyces odontolyticus		85.76	0.52	ESL 1		
Eubacterium contortum		14.24	0.32			
63314302	**Good identification**					**63314302**
Actinomyces odontolyticus		95.44	0.48	ESL 1		
Eubacterium contortum		4.56	0.19			
63314320	**Low discrimination**					**63314320**
Actinomyces odontolyticus		84.74	0.39	ESL 1		
Eubacterium contortum		15.26	0.19	NAG 14		
63315000	**Good identification**					**63315000**
					GRA	
Actinomyces odontolyticus		98.87	0.53	ESL 1	99	
Actinomyces israelii		0.62	0.23		99	
Fusobacterium mortiferum		0.29	0.15		1	
63315002	**Good identification**					**63315002**
Actinomyces odontolyticus		98.49	0.49	ESL 1		
Actinomyces israelii		1.51	0.26			
63315020	**Good identification**					**63315020**
Actinomyces odontolyticus		100.0	0.39	ESL 1		
63315022	**Good identification**					**63315022**
Actinomyces odontolyticus		100.0	0.36	ESL 1		
63315100	**Low discrimination**					**63315100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces odontolyticus		63.10	0.44	ESL 1		
Eubacterium contortum		30.52	0.32			
Actinomyces israelii		5.01	0.33			
63315102	**Low discrimination**					**63315102**
Actinomyces odontolyticus		70.40	0.40	ESL 1		
Actinomyces israelii		13.69	0.36			
Eubacterium contortum		9.80	0.19			
63315200	**Good identification**					**63315200**
Actinomyces odontolyticus		92.71	0.58	ESL 1		
Eubacterium contortum		7.29	0.32			
63315202	**Good identification**					**63315202**
Actinomyces odontolyticus		97.79	0.54	ESL 1		
Eubacterium contortum		2.22	0.19			
63315220	**Acceptable identification**					**63315220**
Actinomyces odontolyticus		92.14	0.44	ESL 1		
Eubacterium contortum		7.86	0.19	NAG 14		
63315222	**Good identification**					**63315222**
Actinomyces odontolyticus		97.60	0.40	ESL 1		
Eubacterium contortum		2.40	0.06	NAG 14		
63315300	**Very good identification**					**63315300**
Eubacterium contortum		96.09	0.66	NIT 1		
Actinomyces odontolyticus		3.90	0.49			
Actinomyces israelii		0.01	0.14			
63315302	**Low discrimination**					**63315302**
Eubacterium contortum		87.58	0.53	NIT 1		
Actinomyces odontolyticus		12.33	0.45			
Actinomyces israelii		0.09	0.17			
63315320	**Good identification**					**63315320**
Eubacterium contortum		96.40	0.53	NIT 1		
Actinomyces odontolyticus		3.61	0.36			
63315322	**Low discrimination**					**63315322**
Eubacterium contortum		88.50	0.40	NIT 1		
Actinomyces odontolyticus		11.50	0.32			
63315700	**Very good identification**					**63315700**
Eubacterium contortum		99.95	0.64	NIT 1		
Actinomyces odontolyticus		0.05	0.15			
63315702	**Good identification**					**63315702**
Eubacterium contortum		99.72	0.52	NIT 1		
Actinomyces odontolyticus		0.17	0.11			
Actinomyces israelii		0.11	0.16			
63315720	**Good identification**					**63315720**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Eubacterium contortum		99.86	0.52	NIT 1	99	
Bacteroides vulgatus		0.14	-.05		1	
63315722	**Good identification**					**63315722**
Eubacterium contortum		100.0	0.39	NIT 1		
63317000	**Low discrimination**					**63317000**
Actinomyces israelii		77.87	0.36	CEL 93		
Actinomyces odontolyticus		22.13	0.19			
63317002	**Low discrimination**					**63317002**
Actinomyces israelii		89.60	0.39	CEL 93		
Actinomyces odontolyticus		10.40	0.15			
63317100	**Good identification**					**63317100**
Actinomyces israelii		97.81	0.46	CEL 93		
Actinomyces odontolyticus		2.19	0.10			
63317102	**Good identification**					**63317102**
Actinomyces israelii		97.79	0.49	CEL 93		
Bifidobacterium adolescentis		1.32	0.11			
Actinomyces odontolyticus		0.90	0.06			
63317112	**Acceptable identification**					**63317112**
Actinomyces israelii		90.09	0.35	CEL 93		
Bifidobacterium adolescentis		9.91	0.12	MLZ 13		
63317140	**Good identification**					**63317140**
Actinomyces israelii		100.0	0.39	CEL 93		
63317142	**Good identification**					**63317142**
Actinomyces israelii		100.0	0.42	CEL 93		
63317402	**Good identification**					**63317402**
Actinomyces israelii		100.0	0.37	CEL 93		
63317500	**Good identification**					**63317500**
Actinomyces israelii		100.0	0.45	CEL 93		
63317502	**Acceptable identification**					**63317502**
Actinomyces israelii		90.09	0.48	CEL 93		
Bifidobacterium adolescentis		9.91	0.25			
63317540	**Good identification**					**63317540**
Actinomyces israelii		100.0	0.38	CEL 93		
63317542	**Good identification**					**63317542**
Actinomyces israelii		96.62	0.41	CEL 93		
Bifidobacterium adolescentis		3.38	0.09			
63331001	**Good identification**					**63331001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ESL 12		
63331003	**Good identification**					**63331003**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63331021	**Good identification**					**63331021**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63331023	**Good identification**					**63331023**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63331102	**Low discrimination**					**63331102**
					COC	
Actinomyces naeslundii		46.39	0.42	ESL 12	1	
Actinomyces israelii		23.60	0.28	MNS 88	1	
Atopobium parvulum		10.17	0.08	RAF 92	99	
63331103	**Low discrimination**					**63331103**
					SPO	
Actinomyces naeslundii		56.26	0.45	ESL 12	1	
Clostridium chauvoei		43.74	0.20	MNS 88	99	
63331120	**Low discrimination**					**63331120**
					COC	
Clostridium perfringens		88.69	0.37	MNS 99	1	
Actinomyces naeslundii		9.28	0.42		1	
Atopobium parvulum		2.03	0.08		99	
63331123	**Low discrimination**					**63331123**
					SPO	
Actinomyces naeslundii		79.42	0.45	ESL 12	1	
Clostridium chauvoei		20.58	0.12	MNS 88	99	
63333000	**Low discrimination**					**63333000**
Actinomyces naeslundii		58.67	0.39	ESL 12		
Actinomyces israelii		41.33	0.28	MNS 88		
63333001	**Good identification**					**63333001**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63333003	**Good identification**					**63333003**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63333020	**Good identification**					**63333020**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63333021	**Good identification**					**63333021**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63333022	**Good identification**					**63333022**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63333023	**Good identification**					**63333023**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63333100	**Low discrimination**					**63333100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		44.22	0.47	ESL 12		
Actinomyces israelii		43.79	0.38	MNS 88		
Propionibacterium propionicum		11.99	0.27	RAF 92		
63333101	**Good identification**					**63333101**
Actinomyces naeslundii		100.0	0.50	ESL 12		
63333102	**Low discrimination**					**63333102**
Actinomyces israelii		51.36	0.41	CEL 93		
Actinomyces naeslundii		34.58	0.47	XYL 92		
Propionibacterium propionicum		14.06	0.30			
63333103	**Good identification**					**63333103**
Actinomyces naeslundii		97.08	0.50	ESL 12		
Propionibacterium propionicum		2.92	0.14	MNS 88		
63333120	**Low discrimination**					**63333120**
Actinomyces naeslundii		89.85	0.47	ESL 12		
Actinomyces israelii		5.68	0.17	MNS 88		
Clostridium perfringens		4.47	0.03	RAF 92		
63333121	**Good identification**					**63333121**
Actinomyces naeslundii		100.0	0.50	ESL 12		
63333122	**Acceptable identification**					**63333122**
Actinomyces naeslundii		91.34	0.47	ESL 12		
Actinomyces israelii		8.66	0.20	MNS 88		
63333123	**Good identification**					**63333123**
Actinomyces naeslundii		100.0	0.50	ESL 12		
63333141	**Good identification**					**63333141**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63333143	**Low discrimination**					**63333143**
Actinomyces naeslundii		85.42	0.37	ESL 12		
Propionibacterium propionicum		14.58	0.14	MNS 88		
63333161	**Good identification**					**63333161**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63333163	**Good identification**					**63333163**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63333500	**Good identification**					**63333500**
Actinomyces israelii		100.0	0.36	CEL 93		
63333502	**Good identification**					**63333502**
Actinomyces israelii		96.76	0.39	CEL 93		
Bifidobacterium adolescentis		3.24	0.08	XYL 92		
63335002	**Low discrimination**					**63335002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		85.11	0.36	CEL 93		
Actinomyces odontolyticus		14.89	0.15			
63335100	**Low discrimination**					**63335100**
					COC	
Actinomyces israelii		60.73	0.43	CEL 93	1	
Atopobium parvulum		35.91	0.25		99	
Actinomyces odontolyticus		2.05	0.10		1	
63335102	**Acceptable identification**					**63335102**
					COC	
Actinomyces israelii		90.92	0.46	CEL 93	1	
Atopobium parvulum		6.33	0.13		99	
Bifidobacterium adolescentis		1.50	0.09		1	
63335140	**Good identification**					**63335140**
Actinomyces israelii		100.0	0.36	CEL 93		
63335142	**Good identification**					**63335142**
Actinomyces israelii		100.0	0.39	CEL 93		
63335500	**Good identification**					**63335500**
Actinomyces israelii		100.0	0.42	CEL 93		
63335502	**Low discrimination**					**63335502**
Actinomyces israelii		88.11	0.45	CEL 93		
Bifidobacterium adolescentis		11.90	0.23			
63335542	**Good identification**					**63335542**
Actinomyces israelii		95.89	0.38	CEL 93		
Bifidobacterium adolescentis		4.11	0.08			
63337000	**Good identification**					**63337000**
Actinomyces israelii		100.0	0.46	CEL 93		
63337002	**Good identification**					**63337002**
Actinomyces israelii		100.0	0.49	CEL 93		
63337040	**Good identification**					**63337040**
Actinomyces israelii		100.0	0.39	CEL 93		
63337042	**Good identification**					**63337042**
Actinomyces israelii		100.0	0.42	CEL 93		
63337100	**Good identification**					**63337100**
Actinomyces israelii		100.0	0.56	CEL 93		
63337102	**Very good identification**					**63337102**
Actinomyces israelii		99.76	0.59	CEL 93		
Bifidobacterium adolescentis		0.24	0.08			
63337110	**Good identification**					**63337110**
Actinomyces israelii		100.0	0.42	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63337112	**Good identification**					**63337112**
Actinomyces israelii		98.09	0.45	CEL 93		
Bifidobacterium adolescentis		1.91	0.09	MLZ 13		
63337120	**Good identification**					**63337120**
Actinomyces israelii		100.0	0.36	CEL 93		
63337122	**Good identification**					**63337122**
Actinomyces israelii		100.0	0.39	CEL 93		
63337140	**Good identification**					**63337140**
Actinomyces israelii		100.0	0.49	CEL 93		
63337142	**Good identification**					**63337142**
Actinomyces israelii		100.0	0.52	CEL 93		
63337152	**Good identification**					**63337152**
Actinomyces israelii		100.0	0.38	CEL 93		
63337300	**Good identification**					**63337300**
Actinomyces israelii		100.0	0.37	CEL 93		
63337302	**Good identification**					**63337302**
Actinomyces israelii		100.0	0.40	CEL 93		
63337400	**Good identification**					**63337400**
Actinomyces israelii		100.0	0.44	CEL 93		
63337402	**Good identification**					**63337402**
Actinomyces israelii		100.0	0.47	CEL 93		
63337440	**Good identification**					**63337440**
Actinomyces israelii		100.0	0.37	CEL 93		
63337442	**Good identification**					**63337442**
Actinomyces israelii		100.0	0.40	CEL 93		
63337500	**Good identification**					**63337500**
Actinomyces israelii		100.0	0.55	CEL 93		
63337502	**Very good identification**					**63337502**
Actinomyces israelii		98.09	0.58	CEL 93		
Bifidobacterium adolescentis		1.91	0.22			
63337510	**Good identification**					**63337510**
Actinomyces israelii		100.0	0.40	CEL 93		
63337512	**Low discrimination**					**63337512**
Actinomyces israelii		86.25	0.43	CEL 93		
Bifidobacterium adolescentis		13.75	0.23	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63337522	**Good identification**					**63337522**
Actinomyces israelii		100.0	0.37	CEL 93		
63337540	**Good identification**					**63337540**
Actinomyces israelii		100.0	0.48	CEL 93		
63337542	**Good identification**					**63337542**
Actinomyces israelii		100.0	0.51	CEL 93		
63337552	**Good identification**					**63337552**
Actinomyces israelii		95.18	0.36	CEL 93		
Bifidobacterium adolescentis		4.82	0.08	MLZ 13		
63337700	**Good identification**					**63337700**
Actinomyces israelii		100.0	0.35	CEL 93		
63337702	**Good identification**					**63337702**
Actinomyces israelii		100.0	0.38	CEL 93		
63341000	**Good identification**					**63341000**
Eubacterium tortuosum		99.93	0.48	MLT 1		
Actinomyces odontolyticus		0.08	0.06	CEL 10		
63341002	**Good identification**					**63341002**
Eubacterium tortuosum		100.0	0.35	MLT 1		
63341020	**Good identification**					**63341020**
Eubacterium tortuosum		100.0	0.35	MLT 1		
63351101	**Low discrimination**					**63351101**
					SPO	
Actinomyces naeslundii		52.57	0.42	ESL 12	1	
Clostridium chauvoei		30.83	0.15	MNS 88	99	
Collinsella aerofaciens		16.59	-.03	RAF 92	1	
63351103	**Acceptable identification**					**63351103**
					SPO	
Actinomyces naeslundii		90.62	0.42	ESL 12	1	
Clostridium chauvoei		9.38	0.03	MNS 88	99	
63351121	**Low discrimination**					**63351121**
					SPO	
Actinomyces naeslundii		72.88	0.42	ESL 12	1	
Clostridium chauvoei		14.25	0.07	MNS 88	99	
Clostridium paraputrificum		12.88	-.03	RAF 92	99	
63351123	**Low discrimination**					**63351123**
					SPO	
Actinomyces naeslundii		85.96	0.42	ESL 12	1	
Clostridium barati		7.53	-.03	MNS 88	99	
Clostridium paraputrificum		6.51	-.09	RAF 92	99	
63351502	**Low discrimination**					**63351502**
Actinomyces israelii		82.69	0.36	XYL 92		
Bifidobacterium adolescentis		17.31	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63353000	**Low discrimination**					**63353000**
Actinomyces israelii		78.47	0.37	XYL 92		
Actinomyces naeslundii		21.54	0.37			
63353001	**Good identification**					**63353001**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63353002	**Low discrimination**					**63353002**
					SPO	
Actinomyces israelii		82.19	0.40	XYL 92	1	
Actinomyces naeslundii		15.04	0.37		1	
Clostridium barati		2.77	-.03		99	
63353003	**Good identification**					**63353003**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63353020	**Low discrimination**					**63353020**
					SPO	
Actinomyces naeslundii		70.58	0.37	ESL 12	1	
Actinomyces israelii		16.42	0.17	MNS 88	1	
Clostridium barati		13.00	-.03	RAF 92	99	
63353021	**Good identification**					**63353021**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63353022	**Low discrimination**					**63353022**
					SPO	
Actinomyces naeslundii		52.59	0.37	ESL 12	1	
Clostridium barati		29.06	0.05	MNS 88	99	
Actinomyces israelii		18.35	0.20	RAF 92	1	
63353023	**Good identification**					**63353023**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63353100	**Low discrimination**					**63353100**
Actinomyces israelii		79.65	0.47	XYL 92		
Actinomyces naeslundii		15.55	0.44			
Collinsella aerofaciens		3.79	-.03			
63353101	**Good identification**					**63353101**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63353102	**Low discrimination**					**63353102**
					SPO	
Actinomyces israelii		85.91	0.50	XYL 92	1	
Actinomyces naeslundii		11.18	0.44		1	
Clostridium barati		2.17	0.05		99	
63353103	**Good identification**					**63353103**
Actinomyces naeslundii		95.08	0.47	ESL 12		
Actinomyces israelii		4.92	0.16	MNS 88		
63353112	**Low discrimination**					**63353112**
					SPO	
Actinomyces israelii		87.10	0.36	XYL 92	1	
Clostridium tertium		8.90	0.05	MLZ 13	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		4.00	0.22		1	
63353120	**Low discrimination**					**63353120**
					SPO	
Actinomyces naeslundii		58.02	0.44	ESL 12	1	
Actinomyces israelii		18.96	0.27	MNS 88	1	
Clostridium barati		11.27	0.05	RAF 92	99	
63353121	**Good identification**					**63353121**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63353122	**Low discrimination**					**63353122**
					SPO	
Actinomyces naeslundii		40.23	0.44	ESL 12	1	
Clostridium barati		23.43	0.14	MNS 88	99	
Actinomyces israelii		19.72	0.30	RAF 92	1	
63353123	**Good identification**					**63353123**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63353140	**Acceptable identification**					**63353140**
Actinomyces israelii		91.86	0.40	XYL 92		
Actinomyces naeslundii		8.14	0.31			
63353142	**Acceptable identification**					**63353142**
Actinomyces israelii		94.42	0.43	XYL 92		
Actinomyces naeslundii		5.58	0.31			
63353400	**Good identification**					**63353400**
Actinomyces israelii		100.0	0.35	XYL 92		
63353402	**Good identification**					**63353402**
Actinomyces israelii		100.0	0.38	XYL 92		
63353500	**Good identification**					**63353500**
Actinomyces israelii		100.0	0.46	XYL 92		
63353502	**Good identification**					**63353502**
Actinomyces israelii		97.07	0.49	XYL 92		
Bifidobacterium adolescentis		2.94	0.16			
63353540	**Good identification**					**63353540**
Actinomyces israelii		100.0	0.39	XYL 92		
63353542	**Good identification**					**63353542**
Actinomyces israelii		100.0	0.42	XYL 92		
63355000	**Low discrimination**					**63355000**
Actinomyces israelii		89.19	0.42			
Actinomyces odontolyticus		10.81	0.19			
63355002	**Good identification**					**63355002**
Actinomyces israelii		95.28	0.45			
Actinomyces odontolyticus		4.72	0.15			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
63355040	**Good identification**							**63355040**
Actinomyces israelii		100.0	0.35					
63355042	**Good identification**							**63355042**
Actinomyces israelii		100.0	0.38					
63355100	**Good identification**							**63355100**
Actinomyces israelii		96.16	0.53					
Collinsella aerofaciens		2.26	-.03					
Actinomyces odontolyticus		0.92	0.10					
63355102	**Good identification**							**63355102**
Actinomyces israelii		98.15	0.56					
Bifidobacterium adolescentis		1.46	0.18					
Actinomyces odontolyticus		0.38	0.06					
63355110	**Good identification**							**63355110**
Actinomyces israelii		100.0	0.38	MLZ 13				
63355112	**Low discrimination**							**63355112**
Actinomyces israelii		89.14	0.41	MLZ 13				
Bifidobacterium adolescentis		10.86	0.19					
63355122	**Low discrimination**							**63355122**
					GRA	SPO	CAT	
Actinomyces israelii		61.73	0.35	NAG 6	99	1	1	
Capnocytophaga ochracea		26.56	0.13		1	1	90	
Clostridium barati		4.20	-.03		99	99	1	
63355140	**Good identification**							**63355140**
Actinomyces israelii		100.0	0.46					
63355142	**Good identification**							**63355142**
Actinomyces israelii		100.0	0.49					
63355302	**Low discrimination**							**63355302**
Actinomyces israelii		55.04	0.36	RHA 7				
Eubacterium contortum		39.41	0.19					
Actinomyces odontolyticus		5.55	0.11					
63355400	**Good identification**							**63355400**
Actinomyces israelii		100.0	0.41					
63355402	**Good identification**							**63355402**
Actinomyces israelii		100.0	0.44					
63355442	**Good identification**							**63355442**
Actinomyces israelii		100.0	0.37					
63355500	**Good identification**							**63355500**
Actinomyces israelii		100.0	0.51					

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63355502	**Low discrimination**					**63355502**
Actinomyces israelii		89.14	0.54			
Bifidobacterium adolescentis		10.86	0.32			
63355510	**Good identification**					**63355510**
Actinomyces israelii		100.0	0.37	MLZ 13		
63355512	**Low discrimination**					**63355512**
Actinomyces israelii		50.08	0.40	MLZ 13		
Bifidobacterium adolescentis		49.92	0.34			
63355540	**Good identification**					**63355540**
Actinomyces israelii		100.0	0.44			
63355542	**Good identification**					**63355542**
Actinomyces israelii		96.27	0.47			
Bifidobacterium adolescentis		3.73	0.17			
63357000	**Good identification**					**63357000**
Actinomyces israelii		100.0	0.55			
63357002	**Very good identification**					**63357002**
Actinomyces israelii		100.0	0.58			
63357010	**Good identification**					**63357010**
Actinomyces israelii		100.0	0.41	MLZ 13		
63357012	**Good identification**					**63357012**
Actinomyces israelii		100.0	0.44	MLZ 13		
63357022	**Good identification**					**63357022**
Actinomyces israelii		100.0	0.38	NAG 6		
63357040	**Good identification**					**63357040**
Actinomyces israelii		100.0	0.48			
63357042	**Good identification**					**63357042**
Actinomyces israelii		100.0	0.51			
63357052	**Good identification**					**63357052**
Actinomyces israelii		100.0	0.37	MLZ 13		
63357100	**Very good identification**					**63357100**
Actinomyces israelii		100.0	0.65			
63357102	**Very good identification**					**63357102**
					SPO	
Actinomyces israelii		99.68	0.68		1	
Bifidobacterium adolescentis		0.21	0.16		1	
Clostridium tertium		0.10	0.03		99	
63357110	**Good identification**					**63357110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces israelii		99.39	0.51	MLZ 13	1	
Clostridium tertium		0.61	-.01		99	
63357112	**Good identification**					**63357112**
					SPO	
Actinomyces israelii		97.10	0.54	MLZ 13	1	
Bifidobacterium adolescentis		1.71	0.18		1	
Clostridium tertium		1.19	0.07		99	
63357120	**Good identification**					**63357120**
					SPO	
Actinomyces israelii		97.63	0.45	NAG 6	1	
Clostridium tertium		2.37	0.03		99	
63357122	**Good identification**					**63357122**
					SPO	
Actinomyces israelii		95.38	0.48	NAG 6	1	
Clostridium tertium		4.62	0.11		99	
63357140	**Very good identification**					**63357140**
Actinomyces israelii		100.0	0.58			
63357142	**Very good identification**					**63357142**
Actinomyces israelii		100.0	0.61			
63357150	**Good identification**					**63357150**
Actinomyces israelii		100.0	0.44	MLZ 13		
63357152	**Good identification**					**63357152**
Actinomyces israelii		100.0	0.47	MLZ 13		
63357160	**Good identification**					**63357160**
Actinomyces israelii		100.0	0.38	NAG 6		
63357162	**Good identification**					**63357162**
Actinomyces israelii		100.0	0.41	NAG 6		
63357200	**Good identification**					**63357200**
Actinomyces israelii		100.0	0.36	RHA 7		
63357202	**Good identification**					**63357202**
Actinomyces israelii		100.0	0.39	RHA 7		
63357300	**Good identification**					**63357300**
Actinomyces israelii		100.0	0.46	RHA 7		
63357302	**Good identification**					**63357302**
Actinomyces israelii		100.0	0.49	RHA 7		
63357312	**Good identification**					**63357312**
Actinomyces israelii		100.0	0.35	RHA 7		
63357340	**Good identification**					**63357340**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.39	RHA 7		
63357342	**Good identification**					**63357342**
Actinomyces israelii		100.0	0.42	RHA 7		
63357400	**Good identification**					**63357400**
Actinomyces israelii		100.0	0.54			
63357402	**Very good identification**					**63357402**
Actinomyces israelii		100.0	0.57			
63357410	**Good identification**					**63357410**
Actinomyces israelii		100.0	0.39	MLZ 13		
63357412	**Good identification**					**63357412**
Actinomyces israelii		100.0	0.43	MLZ 13		
63357422	**Good identification**					**63357422**
Actinomyces israelii		100.0	0.36	NAG 6		
63357440	**Good identification**					**63357440**
Actinomyces israelii		100.0	0.47			
63357442	**Good identification**					**63357442**
Actinomyces israelii		100.0	0.50			
63357452	**Good identification**					**63357452**
Actinomyces israelii		100.0	0.35	MLZ 13		
63357500	**Very good identification**					**63357500**
Actinomyces israelii		100.0	0.64			
63357502	**Very good identification**					**63357502**
Actinomyces israelii		98.27	0.67			
Bifidobacterium adolescentis		1.73	0.31			
63357510	**Good identification**					**63357510**
Actinomyces israelii		100.0	0.50	MLZ 13		
63357512	**Low discrimination**					**63357512**
Actinomyces israelii		87.42	0.53	MLZ 13		
Bifidobacterium adolescentis		12.58	0.32			
63357520	**Good identification**					**63357520**
Actinomyces israelii		100.0	0.43	NAG 6		
63357522	**Good identification**					**63357522**
Actinomyces israelii		100.0	0.46	NAG 6		
63357540	**Very good identification**					**63357540**
Actinomyces israelii		100.0	0.57			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63357542	**Very good identification**					**63357542**
Actinomyces israelii		99.44	0.60			
Bifidobacterium adolescentis		0.56	0.15			
63357550	**Good identification**					**63357550**
Actinomyces israelii		100.0	0.43	MLZ 13		
63357552	**Good identification**					**63357552**
Actinomyces israelii		95.63	0.46	MLZ 13		
Bifidobacterium adolescentis		4.37	0.17			
63357560	**Good identification**					**63357560**
Actinomyces israelii		100.0	0.36	NAG 6		
63357562	**Good identification**					**63357562**
Actinomyces israelii		100.0	0.39	NAG 6		
63357602	**Good identification**					**63357602**
Actinomyces israelii		100.0	0.37	RHA 7		
63357700	**Good identification**					**63357700**
Actinomyces israelii		100.0	0.45	RHA 7		
63357702	**Good identification**					**63357702**
Actinomyces israelii		100.0	0.48	RHA 7		
63357740	**Good identification**					**63357740**
Actinomyces israelii		100.0	0.38	RHA 7		
63357742	**Good identification**					**63357742**
Actinomyces israelii		100.0	0.41	RHA 7		
63367100	**Good identification**					**63367100**
Actinomyces israelii		100.0	0.41	FRU 99		
63367102	**Good identification**					**63367102**
Actinomyces israelii		100.0	0.44	FRU 99		
63367142	**Good identification**					**63367142**
Actinomyces israelii		100.0	0.37	FRU 99		
63367500	**Good identification**					**63367500**
Actinomyces israelii		100.0	0.40	FRU 99		
63367502	**Good identification**					**63367502**
Actinomyces israelii		100.0	0.43	FRU 99		
63367542	**Good identification**					**63367542**
Actinomyces israelii		100.0	0.36	FRU 99		
63371001	**Good identification**					**63371001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	ESL 12		
63371003	**Good identification**					**63371003**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63371021	**Good identification**					**63371021**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63371023	**Good identification**					**63371023**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63371100	**Low discrimination**					**63371100**
					COC	
Actinomyces israelii		47.37	0.44	XYL 92	1	
Atopobium parvulum		30.46	0.27		99	
Actinomyces naeslundii		13.94	0.44		1	
63371101	**Good identification**					**63371101**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63371102	**Low discrimination**					**63371102**
					GRA	
Actinomyces israelii		47.12	0.47	XYL 92	99	
Lactobacillus acidophilus		19.50	0.25		99	
Leptotrichia buccalis		18.78	0.15		1	
63371103	**Good identification**					**63371103**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63371112	**Good identification**					**63371112**
					GRA	
Leptotrichia buccalis		99.60	0.50	NIT 10	1	
Actinomyces israelii		0.38	0.33	MNS 99	99	
Actinomyces naeslundii		0.03	0.22		99	
63371121	**Good identification**					**63371121**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63371123	**Good identification**					**63371123**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63371140	**Low discrimination**					**63371140**
Actinomyces israelii		88.22	0.37	XYL 92		
Actinomyces naeslundii		11.78	0.31			
63371142	**Low discrimination**					**63371142**
Actinomyces israelii		81.93	0.40	XYL 92		
Lactobacillus acidophilus		10.78	0.10			
Actinomyces naeslundii		7.29	0.31			
63371402	**Good identification**					**63371402**
Actinomyces israelii		100.0	0.35	XYL 92		
63371500	**Good identification**					**63371500**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	100.0	0.43	XYL 92		
63371502	**Good identification**					**63371502**
	Actinomyces israelii	96.42	0.46	XYL 92		
	Bifidobacterium adolescentis	3.58	0.15			
63371540	**Good identification**					**63371540**
	Actinomyces israelii	100.0	0.36	XYL 92		
63371542	**Good identification**					**63371542**
	Actinomyces israelii	100.0	0.39	XYL 92		
63373000	**Low discrimination**					**63373000**
	Actinomyces israelii	87.60	0.47	XYL 92		
	Actinomyces naeslundii	12.41	0.41			
63373001	**Good identification**					**63373001**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
63373002	**Acceptable identification**					**63373002**
	Actinomyces israelii	91.37	0.50	XYL 92		
	Actinomyces naeslundii	8.63	0.41			
63373003	**Acceptable identification**					**63373003**
	Actinomyces naeslundii	93.34	0.44	ESL 12		
	Actinomyces israelii	6.66	0.16	MNS 88		
63373012	**Good identification**					**63373012**
	Actinomyces israelii	100.0	0.36	XYL 92		
63373020	**Low discrimination**					**63373020**
	Actinomyces naeslundii	68.93	0.41	ESL 12		
	Actinomyces israelii	31.07	0.26	MNS 88		
63373021	**Good identification**					**63373021**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
63373022	**Low discrimination**					**63373022**
	Actinomyces naeslundii	59.66	0.41	ESL 12		
	Actinomyces israelii	40.34	0.29	MNS 88		
63373023	**Good identification**					**63373023**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
63373040	**Acceptable identification**					**63373040**
	Actinomyces israelii	93.96	0.40	XYL 92		
	Actinomyces naeslundii	6.04	0.29			
63373042	**Good identification**					**63373042**
	Actinomyces israelii	95.89	0.43	XYL 92		
	Actinomyces naeslundii	4.11	0.29			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63373100	**Good identification**						**63373100**
					SPO		
Actinomyces israelii		90.01	0.57	XYL 92	1		
Actinomyces naeslundii		9.07	0.49		1		
Clostridium tertium		0.93	0.09		99		
63373101	**Acceptable identification**						**63373101**
Actinomyces naeslundii		93.74	0.52	ESL 12			
Actinomyces israelii		6.26	0.23	MNS 88			
63373102	**Good identification**						**63373102**
					SPO		
Actinomyces israelii		91.61	0.60	XYL 92	1		
Actinomyces naeslundii		6.15	0.49		1		
Clostridium tertium		1.88	0.17		99		
63373103	**Acceptable identification**						**63373103**
Actinomyces naeslundii		90.89	0.52	ESL 12			
Actinomyces israelii		9.11	0.26	MNS 88			
63373110	**Low discrimination**						**63373110**
					SPO		
Actinomyces israelii		86.38	0.43	XYL 92	1		
Clostridium tertium		10.56	0.13	MLZ 13	99		
Actinomyces naeslundii		3.07	0.27		1		
63373112	**Low discrimination**						**63373112**
					GRA	SPO	
Actinomyces israelii		57.51	0.46	XYL 92	99	1	
Leptotrichia buccalis		27.07	0.15	MLZ 13	1	1	
Clostridium tertium		14.06	0.21		99	99	
63373120	**Low discrimination**						**63373120**
					SPO		
Actinomyces naeslundii		47.22	0.49	ESL 12	1		
Actinomyces israelii		29.91	0.37	MNS 88	1		
Clostridium tertium		14.44	0.17	RAF 92	99		
63373121	**Good identification**						**63373121**
Actinomyces naeslundii		100.0	0.52	ESL 12			
63373122	**Low discrimination**						**63373122**
					SPO		
Actinomyces naeslundii		27.48	0.49	ESL 12	1		
Actinomyces israelii		26.11	0.40	MNS 88	1		
Clostridium tertium		25.22	0.25	RAF 92	99		
63373123	**Good identification**						**63373123**
Actinomyces naeslundii		100.0	0.52	ESL 12			
63373140	**Good identification**						**63373140**
Actinomyces israelii		95.63	0.50	XYL 92			
Actinomyces naeslundii		4.37	0.36				
63373141	**Low discrimination**						**63373141**
Actinomyces naeslundii		87.17	0.39	ESL 12			
Actinomyces israelii		12.84	0.16	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63373142	**Good identification**					**63373142**
Actinomyces israelii		97.04	0.53	XYL 92		
Actinomyces naeslundii		2.96	0.36			
63373143	**Low discrimination**					**63373143**
Actinomyces naeslundii		81.91	0.39	ESL 12		
Actinomyces israelii		18.09	0.19	MNS 88		
63373150	**Good identification**					**63373150**
Actinomyces israelii		100.0	0.36	XYL 92		
63373152	**Good identification**					**63373152**
Actinomyces israelii		100.0	0.39	XYL 92		
63373161	**Good identification**					**63373161**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63373163	**Good identification**					**63373163**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63373300	**Good identification**					**63373300**
Actinomyces israelii		100.0	0.38	XYL 92		
63373302	**Good identification**					**63373302**
Actinomyces israelii		100.0	0.41	XYL 92		
63373400	**Good identification**					**63373400**
Actinomyces israelii		100.0	0.45	XYL 92		
63373402	**Good identification**					**63373402**
Actinomyces israelii		100.0	0.48	XYL 92		
63373440	**Good identification**					**63373440**
Actinomyces israelii		100.0	0.38	XYL 92		
63373442	**Good identification**					**63373442**
Actinomyces israelii		100.0	0.41	XYL 92		
63373500	**Good identification**					**63373500**
Actinomyces israelii		100.0	0.56	XYL 92		
63373502	**Very good identification**					**63373502**
Actinomyces israelii		99.47	0.59	XYL 92		
Bifidobacterium adolescentis		0.53	0.13			
63373510	**Good identification**					**63373510**
Actinomyces israelii		100.0	0.41	XYL 92		
63373512	**Good identification**					**63373512**
Actinomyces israelii		95.80	0.45	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		4.20	0.15	MLZ 13			
63373520	**Good identification**						**63373520**
Actinomyces israelii		100.0	0.35	XYL 92			
63373522	**Good identification**						**63373522**
Actinomyces israelii		100.0	0.38	XYL 92			
63373540	**Good identification**						**63373540**
Actinomyces israelii		100.0	0.49	XYL 92			
63373542	**Good identification**						**63373542**
Actinomyces israelii		100.0	0.52	XYL 92			
63373552	**Good identification**						**63373552**
Actinomyces israelii		100.0	0.37	XYL 92			
63373700	**Good identification**						**63373700**
Actinomyces israelii		100.0	0.36	XYL 92			
63373702	**Good identification**						**63373702**
Actinomyces israelii		100.0	0.39	XYL 92			
63375000	**Good identification**						**63375000**
Actinomyces israelii		100.0	0.52				
63375002	**Good identification**						**63375002**
Actinomyces israelii		100.0	0.55				
63375010	**Good identification**						**63375010**
Actinomyces israelii		100.0	0.38	MLZ 13			
63375012	**Good identification**						**63375012**
Actinomyces israelii		100.0	0.41	MLZ 13			
63375040	**Good identification**						**63375040**
Actinomyces israelii		100.0	0.45				
63375042	**Good identification**						**63375042**
Actinomyces israelii		100.0	0.48				
63375100	**Good identification**						**63375100**
					COC	SPO	
Actinomyces israelii		90.46	0.62		1	1	
Atopobium parvulum		9.39	0.32		99	1	
Clostridium butyricum		0.15	-.07		1	99	
63375102	**Very good identification**						**63375102**
					COC		
Actinomyces israelii		98.54	0.65		1		
Atopobium parvulum		1.20	0.19		99		
Bifidobacterium adolescentis		0.26	0.15		1		
63375110	**Good identification**						**63375110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.48	MLZ 13		
63375112	**Low discrimination**					**63375112**
					GRA	
Actinomyces israelii		79.77	0.51	MLZ 13	99	
Leptotrichia buccalis		18.50	0.15		1	
Bifidobacterium adolescentis		1.72	0.16		99	
63375120	**Low discrimination**					**63375120**
					COC	
Actinomyces israelii		77.69	0.42	NAG 6	1	
Atopobium parvulum		22.31	0.19		99	
63375122	**Good identification**					**63375122**
					COC	
Actinomyces israelii		96.73	0.45	NAG 6	1	
Atopobium parvulum		3.27	0.06		99	
63375140	**Good identification**					**63375140**
Actinomyces israelii		100.0	0.55			
63375142	**Very good identification**					**63375142**
Actinomyces israelii		100.0	0.58			
63375150	**Good identification**					**63375150**
Actinomyces israelii		100.0	0.41	MLZ 13		
63375152	**Good identification**					**63375152**
Actinomyces israelii		100.0	0.44	MLZ 13		
63375162	**Good identification**					**63375162**
Actinomyces israelii		100.0	0.38	NAG 6		
63375202	**Good identification**					**63375202**
Actinomyces israelii		100.0	0.36	RHA 7		
63375300	**Good identification**					**63375300**
Actinomyces israelii		100.0	0.43	RHA 7		
63375302	**Good identification**					**63375302**
Actinomyces israelii		100.0	0.46	RHA 7		
63375340	**Good identification**					**63375340**
Actinomyces israelii		100.0	0.36	RHA 7		
63375342	**Good identification**					**63375342**
Actinomyces israelii		100.0	0.39	RHA 7		
63375400	**Good identification**					**63375400**
Actinomyces israelii		100.0	0.51			
63375402	**Good identification**					**63375402**
Actinomyces israelii		100.0	0.54			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63375410	**Good identification**						**63375410**
Actinomyces israelii		100.0	0.36	MLZ 13			
63375412	**Good identification**						**63375412**
Actinomyces israelii		100.0	0.39	MLZ 13			
63375440	**Good identification**						**63375440**
Actinomyces israelii		100.0	0.44				
63375442	**Good identification**						**63375442**
Actinomyces israelii		100.0	0.47				
63375500	**Very good identification**						**63375500**
					GRA	SPO	
Actinomyces israelii		95.91	0.61		99	1	
Mitsuokella multacida		2.40	0.21		1	1	
Clostridium butyricum		1.70	0.10		99	99	
63375502	**Very good identification**						**63375502**
					SPO		
Actinomyces israelii		97.27	0.64		1		
Bifidobacterium adolescentis		2.10	0.29		1		
Clostridium butyricum		0.45	0.03		99		
63375510	**Acceptable identification**						**63375510**
					GRA		
Actinomyces israelii		94.72	0.47	MLZ 13	99		
Mitsuokella multacida		5.28	0.13		1		
63375512	**Low discrimination**						**63375512**
Actinomyces israelii		84.99	0.50	MLZ 13			
Bifidobacterium adolescentis		15.01	0.31				
63375520	**Acceptable identification**						**63375520**
					GRA		
Actinomyces israelii		91.08	0.40	NAG 6	99		
Mitsuokella multacida		8.92	0.11		1		
63375522	**Good identification**						**63375522**
Actinomyces israelii		100.0	0.43	NAG 6			
63375540	**Acceptable identification**						**63375540**
					GRA		
Actinomyces israelii		93.96	0.54		99		
Mitsuokella multacida		6.04	0.21		1		
63375542	**Very good identification**						**63375542**
					GRA		
Actinomyces israelii		98.85	0.57		99		
Bifidobacterium adolescentis		0.68	0.14		99		
Mitsuokella multacida		0.47	0.04		1		
63375550	**Low discrimination**						**63375550**
					GRA		
Actinomyces israelii		87.46	0.40	MLZ 13	99		
Mitsuokella multacida		12.54	0.13		1		
63375552	**Acceptable identification**						**63375552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		94.68	0.43	MLZ 13		
Bifidobacterium adolescentis		5.32	0.15			
63375562	**Good identification**					**63375562**
Actinomyces israelii		100.0	0.36	NAG 6		
63375700	**Low discrimination**					**63375700**
					GRA	
Actinomyces israelii		61.85	0.42	RHA 7	99	
Mitsuokella multacida		38.15	0.25		1	
63375702	**Good identification**					**63375702**
					GRA	
Actinomyces israelii		95.63	0.45	RHA 7	99	
Mitsuokella multacida		4.37	0.09		1	
63375742	**Low discrimination**					**63375742**
					GRA	
Actinomyces israelii		89.49	0.38	RHA 7	99	
Mitsuokella multacida		10.52	0.09		1	
63376100	**Good identification**					**63376100**
Actinomyces israelii		100.0	0.41	GAL 99		
63376102	**Good identification**					**63376102**
Actinomyces israelii		100.0	0.44	GAL 99		
63376142	**Good identification**					**63376142**
Actinomyces israelii		100.0	0.37	GAL 99		
63376500	**Good identification**					**63376500**
Actinomyces israelii		100.0	0.40	GAL 99		
63376502	**Good identification**					**63376502**
Actinomyces israelii		100.0	0.43	GAL 99		
63376542	**Good identification**					**63376542**
Actinomyces israelii		100.0	0.36	GAL 99		
63377000	**Very good identification**					**63377000**
Actinomyces israelii		100.0	0.65			
63377002	**Very good identification**					**63377002**
Actinomyces israelii		100.0	0.68			
63377010	**Good identification**					**63377010**
Actinomyces israelii		100.0	0.51	MLZ 13		
63377012	**Good identification**					**63377012**
Actinomyces israelii		100.0	0.54	MLZ 13		
63377020	**Good identification**					**63377020**
Actinomyces israelii		100.0	0.45	NAG 6		
63377022	**Good identification**					**63377022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.48	NAG 6		
63377040	**Very good identification**					**63377040**
Actinomyces israelii		100.0	0.58			
63377042	**Very good identification**					**63377042**
Actinomyces israelii		100.0	0.61			
63377050	**Good identification**					**63377050**
Actinomyces israelii		100.0	0.44	MLZ 13		
63377052	**Good identification**					**63377052**
Actinomyces israelii		100.0	0.47	MLZ 13		
63377060	**Good identification**					**63377060**
Actinomyces israelii		100.0	0.37	NAG 6		
63377062	**Good identification**					**63377062**
Actinomyces israelii		100.0	0.40	NAG 6		
63377100	**Very good identification**					**63377100**
					SPO	
Actinomyces israelii		99.88	0.75		1	
Clostridium tertium		0.12	0.11		99	
63377101	**Good identification**					**63377101**
Actinomyces israelii		100.0	0.41	URE 1		
63377102	**Very good identification**					**63377102**
					SPO	
Actinomyces israelii		99.72	0.78		1	
Clostridium tertium		0.25	0.19		99	
Bifidobacterium adolescentis		0.04	0.13		1	
63377103	**Good identification**					**63377103**
Actinomyces israelii		100.0	0.44	URE 1		
63377110	**Very good identification**					**63377110**
					SPO	
Actinomyces israelii		98.55	0.61	MLZ 13	1	
Clostridium tertium		1.45	0.15		99	
63377112	**Very good identification**					**63377112**
					SPO	
Actinomyces israelii		96.85	0.64	MLZ 13	1	
Clostridium tertium		2.84	0.24		99	
Bifidobacterium adolescentis		0.30	0.15		1	
63377120	**Acceptable identification**					**63377120**
					SPO	
Actinomyces israelii		94.52	0.55	NAG 6	1	
Clostridium tertium		5.48	0.19		99	
63377122	**Low discrimination**					**63377122**
					SPO	
Actinomyces israelii		89.61	0.58	NAG 6	1	
Clostridium tertium		10.39	0.27		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
63377130	**Low discrimination**					**63377130**
					SPO	
Actinomyces israelii		59.17	0.41	MLZ 13	1	
Clostridium tertium		40.83	0.24	NAG 6	99	
63377140	**Very good identification**					**63377140**
Actinomyces israelii		100.0	0.68			
63377142	**Very good identification**					**63377142**
Actinomyces israelii		100.0	0.71			
63377143	**Good identification**					**63377143**
Actinomyces israelii		100.0	0.37	URE 1		
63377150	**Good identification**					**63377150**
Actinomyces israelii		100.0	0.54	MLZ 13		
63377152	**Very good identification**					**63377152**
Actinomyces israelii		100.0	0.57	MLZ 13		
63377160	**Good identification**					**63377160**
Actinomyces israelii		100.0	0.48	NAG 6		
63377162	**Good identification**					**63377162**
Actinomyces israelii		100.0	0.51	NAG 6		
63377172	**Good identification**					**63377172**
Actinomyces israelii		100.0	0.37	MLZ 13		
63377200	**Good identification**					**63377200**
Actinomyces israelii		100.0	0.46	RHA 7		
63377202	**Good identification**					**63377202**
Actinomyces israelii		100.0	0.49	RHA 7		
63377240	**Good identification**					**63377240**
Actinomyces israelii		100.0	0.39	RHA 7		
63377242	**Good identification**					**63377242**
Actinomyces israelii		100.0	0.42	RHA 7		
63377300	**Good identification**					**63377300**
Actinomyces israelii		100.0	0.56	RHA 7		
63377302	**Very good identification**					**63377302**
Actinomyces israelii		100.0	0.59	RHA 7		
63377310	**Good identification**					**63377310**
Actinomyces israelii		100.0	0.42	RHA 7		
63377312	**Good identification**					**63377312**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	100.0	0.45	RHA 7		
63377320	**Good identification**					**63377320**
	Actinomyces israelii	100.0	0.36	RHA 7		
63377322	**Good identification**					**63377322**
	Actinomyces israelii	100.0	0.39	RHA 7		
63377340	**Good identification**					**63377340**
	Actinomyces israelii	100.0	0.49	RHA 7		
63377342	**Good identification**					**63377342**
	Actinomyces israelii	100.0	0.52	RHA 7		
63377352	**Good identification**					**63377352**
	Actinomyces israelii	100.0	0.38	RHA 7		
63377400	**Very good identification**					**63377400**
	Actinomyces israelii	100.0	0.63			
63377402	**Very good identification**					**63377402**
	Actinomyces israelii	100.0	0.66			
63377410	**Good identification**					**63377410**
	Actinomyces israelii	100.0	0.49	MLZ 13		
63377412	**Good identification**					**63377412**
	Actinomyces israelii	100.0	0.52	MLZ 13		
63377420	**Good identification**					**63377420**
	Actinomyces israelii	100.0	0.43	NAG 6		
63377422	**Good identification**					**63377422**
	Actinomyces israelii	100.0	0.46	NAG 6		
63377440	**Very good identification**					**63377440**
	Actinomyces israelii	100.0	0.56			
63377442	**Very good identification**					**63377442**
	Actinomyces israelii	100.0	0.59			
63377450	**Good identification**					**63377450**
	Actinomyces israelii	100.0	0.42	MLZ 13		
63377452	**Good identification**					**63377452**
	Actinomyces israelii	100.0	0.45	MLZ 13		
63377460	**Good identification**					**63377460**
	Actinomyces israelii	100.0	0.36	NAG 6		
63377462	**Good identification**					**63377462**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.39	NAG 6			
63377500	**Very good identification**						**63377500**
					GRA	SPO	
Actinomyces israelii		98.95	0.74		99	1	
Mitsuokella multacida		1.02	0.27		1	1	
Clostridium butyricum		0.03	-.07		99	99	
63377501	**Low discrimination**						**63377501**
					GRA		
Actinomyces israelii		89.83	0.40	URE 1	99		
Mitsuokella multacida		10.17	0.11		1		
63377502	**Very good identification**						**63377502**
					GRA		
Actinomyces israelii		99.61	0.77		99		
Bifidobacterium adolescentis		0.31	0.28		99		
Mitsuokella multacida		0.08	0.11		1		
63377503	**Good identification**						**63377503**
Actinomyces israelii		100.0	0.43	URE 1			
63377510	**Very good identification**						**63377510**
					GRA		
Actinomyces israelii		97.76	0.60	MLZ 13	99		
Mitsuokella multacida		2.25	0.19		1		
63377512	**Very good identification**						**63377512**
Actinomyces israelii		96.22	0.63	MLZ 13			
Bifidobacterium adolescentis		2.45	0.29				
Bifidobacterium dentium		1.17	0.06				
63377520	**Good identification**						**63377520**
					GRA		
Actinomyces israelii		96.12	0.53	NAG 6	99		
Mitsuokella multacida		3.88	0.17		1		
63377522	**Very good identification**						**63377522**
Actinomyces israelii		100.0	0.56	NAG 6			
63377530	**Low discrimination**						**63377530**
					GRA		
Actinomyces israelii		87.71	0.39	MLZ 13	99		
Mitsuokella multacida		7.89	0.09	NAG 6	1		
Bifidobacterium dentium		4.40	-.07		99		
63377532	**Low discrimination**						**63377532**
Actinomyces israelii		74.73	0.42	MLZ 13			
Bifidobacterium dentium		25.27	0.10	NAG 6			
63377540	**Very good identification**						**63377540**
					GRA		
Actinomyces israelii		97.42	0.67		99		
Mitsuokella multacida		2.58	0.27		1		
63377542	**Very good identification**						**63377542**
					GRA		
Actinomyces israelii		99.71	0.70		99		
Mitsuokella multacida		0.20	0.11		1		
Bifidobacterium adolescentis		0.10	0.12		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63377543	**Good identification**						**63377543**
Actinomyces israelii		100.0	0.36	URE 1			
63377550	**Acceptable identification**						**63377550**
					COC	GRA	
Actinomyces israelii		93.64	0.53	MLZ 13	1	99	
Mitsuokella multacida		5.53	0.19		1	1	
Blautia producta		0.83	-.03		99	99	
63377552	**Good identification**						**63377552**
					GRA		
Actinomyces israelii		98.77	0.56	MLZ 13	99		
Bifidobacterium adolescentis		0.80	0.14		99		
Mitsuokella multacida		0.43	0.03		1		
63377560	**Acceptable identification**						**63377560**
					GRA		
Actinomyces israelii		90.61	0.46	NAG 6	99		
Mitsuokella multacida		9.40	0.17		1		
63377562	**Good identification**						**63377562**
Actinomyces israelii		100.0	0.49	NAG 6			
63377572	**Good identification**						**63377572**
Actinomyces israelii		100.0	0.35	MLZ 13			
63377600	**Good identification**						**63377600**
Actinomyces israelii		100.0	0.44	RHA 7			
63377602	**Good identification**						**63377602**
Actinomyces israelii		100.0	0.47	RHA 7			
63377640	**Good identification**						**63377640**
Actinomyces israelii		100.0	0.37	RHA 7			
63377642	**Good identification**						**63377642**
Actinomyces israelii		100.0	0.40	RHA 7			
63377700	**Low discrimination**						**63377700**
					GRA		
Actinomyces israelii		79.74	0.55	RHA 7	99		
Mitsuokella multacida		20.26	0.32		1		
63377702	**Very good identification**						**63377702**
					GRA		
Actinomyces israelii		98.15	0.58	RHA 7	99		
Mitsuokella multacida		1.85	0.15		1		
63377710	**Low discrimination**						**63377710**
					GRA		
Actinomyces israelii		63.83	0.40	RHA 7	99		
Mitsuokella multacida		36.17	0.24	MLZ 13	1		
63377712	**Good identification**						**63377712**
					GRA		
Actinomyces israelii		95.97	0.43	RHA 7	99		
Mitsuokella multacida		4.03	0.07	MLZ 13	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
63377722	**Acceptable identification**						**63377722**
					GRA		
Actinomyces israelii		93.14	0.37	RHA 7	99		
Mitsuokella multacida		6.86	0.05	NAG 6	1		
63377740	**Low discrimination**						**63377740**
					GRA		
Actinomyces israelii		60.49	0.48	RHA 7	99		
Mitsuokella multacida		39.51	0.32		1		
63377742	**Good identification**						**63377742**
					GRA		
Actinomyces israelii		95.38	0.51	RHA 7	99		
Mitsuokella multacida		4.62	0.15		1		
63377752	**Acceptable identification**						**63377752**
					GRA		
Actinomyces israelii		90.26	0.36	RHA 7	99		
Mitsuokella multacida		9.74	0.07	MLZ 13	1		
63401000	**Very good identification**						**63401000**
					GRA		
Eubacterium tortuosum		99.96	0.56	SAL 99	99		
Fusobacterium mortiferum		0.05	0.16		1		
63401002	**Good identification**						**63401002**
Eubacterium tortuosum		100.0	0.43	SAL 99			
63401020	**Good identification**						**63401020**
Eubacterium tortuosum		100.0	0.43	SAL 99			
63410000	**Good identification**						**63410000**
Fusobacterium mortiferum		100.0	0.47	NIT 1			
63410100	**Good identification**						**63410100**
Fusobacterium mortiferum		100.0	0.53	NIT 1			
63410102	**Low discrimination**						**63410102**
					GRA	CAT	
Fusobacterium mortiferum		86.44	0.37	NIT 1	1	5	
Actinomyces viscosus		10.04	0.08	BGL 10	99	76	
Capnocytophaga ochracea		3.52	-.01		1	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
63411000	**Low discrimination**						**63411000**
					GRA		
Fusobacterium mortiferum		80.02	0.50	NIT 1	1		
Eubacterium tortuosum		18.29	0.22		99		
Lactobacillus fermentum		1.18	0.20		99		
63411100	**Acceptable identification**						**63411100**
					GRA	CAT	
Fusobacterium mortiferum		94.93	0.56	NIT 1	1	5	
Capnocytophaga ochracea		3.15	0.17		1	90	
Lactobacillus fermentum		0.98	0.24		99	1	
63411122	**Good identification**						**63411122**
					GRA	CAT	
Capnocytophaga ochracea		99.53	0.50	MNS 90	1	90	
Prevotella melaninogenica		0.21	0.00	MLT 99	1	1	
Actinomyces naeslundii		0.13	0.26		99	25	
63411522	**Good identification**						**63411522**
					GRA	CAT	
Capnocytophaga ochracea		98.80	0.37	MNS 90	1	90	
Lactobacillus fermentum		0.72	0.15	MLT 99	99	1	
Bifidobacterium longum subsp. longum		0.49	0.05		99	1	
63415100	**Low discrimination**						**63415100**
					GRA	CAT	
Fusobacterium mortiferum		80.86	0.37	NIT 1	1	5	
Capnocytophaga ochracea		8.91	0.07	XYL 7	1	90	
Lactobacillus fermentum		6.52	0.20		99	1	
63415122	**Good identification**						**63415122**
Capnocytophaga ochracea		96.25	0.39	MNS 90			
Bacteroides fragilis		3.75	0.06	MLT 99			
63430000	**Good identification**						**63430000**
Fusobacterium mortiferum		100.0	0.38	NIT 1			
63430100	**Good identification**						**63430100**
Fusobacterium mortiferum		100.0	0.44	NIT 1			
63431000	**Good identification**						**63431000**
					GRA		
Fusobacterium mortiferum		98.17	0.41	NIT 1	1		
Actinomyces naeslundii		1.83	0.23		99		
63431100	**Good identification**						**63431100**
					GRA		
Fusobacterium mortiferum		96.42	0.47	NIT 1	1		
Actinomyces naeslundii		2.20	0.31		99		
Eubacterium saburreum		1.38	0.14		85		
63433100	**Low discrimination**						**63433100**
					GRA		
Actinomyces naeslundii		47.59	0.35	ESL 12	99		
Propionibacterium propionicum		41.54	0.24	MNS 88	99		
Fusobacterium mortiferum		10.86	0.13	MLT 94	1		
63433101	**Good identification**						**63433101**
Actinomyces naeslundii		100.0	0.38	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63433103	**Good identification**						**63433103**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63433120	**Good identification**						**63433120**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63433121	**Good identification**						**63433121**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63433122	**Good identification**						**63433122**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63433123	**Good identification**						**63433123**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63441000	**Good identification**						**63441000**
Eubacterium tortuosum		100.0	0.40	SAL 99			
63450000	**Good identification**						**63450000**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
63450100	**Good identification**						**63450100**
Fusobacterium mortiferum		100.0	0.45	NIT 1			
63451000	**Acceptable identification**						**63451000**
					GRA		
Fusobacterium mortiferum		92.92	0.42	NIT 1	1		
Eubacterium tortuosum		7.08	0.06		99		
63451100	**Good identification**						**63451100**
					GRA	CAT	
Fusobacterium mortiferum		96.26	0.48	NIT 1	1	5	
Capnocytophaga ochracea		2.40	0.07		1	90	
Actinomyces naeslundii		1.34	0.28		99	25	
63451122	**Good identification**						**63451122**
					GRA		
Capnocytophaga ochracea		99.31	0.39	MNS 90	1		
Actinomyces naeslundii		0.69	0.28	MLT 99	99		
63453101	**Good identification**						**63453101**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63453103	**Good identification**						**63453103**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63453121	**Good identification**						**63453121**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63453123	**Good identification**						**63453123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63470100	**Good identification**						**63470100**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.36	NIT 1			
63471100	**Acceptable identification**						**63471100**
					GRA		
Fusobacterium mortiferum		91.69	0.39	NIT 1	1		
Actinomyces naeslundii		8.31	0.33		99		
63471101	**Good identification**						**63471101**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63471103	**Good identification**						**63471103**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63471121	**Good identification**						**63471121**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63471123	**Good identification**						**63471123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63473100	**Low discrimination**						**63473100**
Actinomyces naeslundii		86.33	0.38	ESL 12			
Actinomyces israelii		13.67	0.15	MNS 88			
63473101	**Good identification**						**63473101**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63473102	**Low discrimination**						**63473102**
Actinomyces naeslundii		80.81	0.38	ESL 12			
Actinomyces israelii		19.20	0.18	MNS 88			
63473103	**Good identification**						**63473103**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63473120	**Good identification**						**63473120**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63473121	**Good identification**						**63473121**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63473122	**Good identification**						**63473122**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63473123	**Good identification**						**63473123**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63477102	**Good identification**						**63477102**
Actinomyces israelii		100.0	0.36	MLT 99			
63501101	**Good identification**						**63501101**
					COC	SPO	
Clostridium chauvoei		99.02	0.43	MNS 90	1	99	
Actinomyces naeslundii		0.51	0.27	RAF 1	1	1	
Blautia hansenii		0.47	-.03		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
63501121	**Good identification**							**63501121**
					GRA	SPO		
Clostridium chauvoei		96.31	0.35	MNS 90	99	99		
Actinomyces naeslundii		1.50	0.27	RAF 1	99	1		
Prevotella melaninogenica		1.15	-.04		1	1		
63501122	**Good identification**							**63501122**
					COC	CAT	GRA	
Capnocytophaga ochracea		97.43	0.50	MNS 90	1	90	1	
Prevotella melaninogenica		2.29	0.18	FRU 99	1	1	1	
Blautia hansenii		0.18	-.01		99	1	99	
63501522	**Acceptable identification**							**63501522**
					GRA	CAT		
Capnocytophaga ochracea		94.85	0.37	MNS 90	1	90		
Bifidobacterium longum subsp. longum		5.15	0.23	FRU 99	99	1		
63505122	**Good identification**							**63505122**
Capnocytophaga ochracea		96.25	0.39	MNS 90				
Bacteroides fragilis		3.75	0.06	FRU 99				
63505530	**Good identification**							**63505530**
Bifidobacterium longum subsp. longum		100.0	0.35	NIT 1				
63505532	**Acceptable identification**							**63505532**
					GRA	CAT		
Bifidobacterium longum subsp. longum		90.90	0.35	NIT 1	99	1		
Capnocytophaga ochracea		9.10	0.10	MNS 87	1	90		
63510002	**Acceptable identification**							**63510002**
					GRA			
Actinomyces viscosus		91.00	0.38	ESL 1	99			
Actinomyces odontolyticus		7.68	0.27	MNS 96	99			
Fusobacterium mortiferum		0.75	0.15		1			
63510100	**Low discrimination**							**63510100**
					GRA	CAT		
Fusobacterium mortiferum		51.06	0.37	NIT 1	1	5		
Capnocytophaga ochracea		20.56	0.17	MLT 11	1	90		
Actinomyces odontolyticus		12.90	0.22		99	10		
63510102	**Low discrimination**							**63510102**
					GRA	CAT		
Actinomyces viscosus		71.23	0.42	ESL 1	99	76		
Capnocytophaga ochracea		24.97	0.33	MNS 96	1	90		
Propionibacterium propionicum		1.77	0.29		99	1		
63510122	**Good identification**							**63510122**
					GRA	CAT		
Capnocytophaga ochracea		99.39	0.50	MNS 90	1	90		
Actinomyces viscosus		0.32	0.08	GAL 99	99	76		
Prevotella melaninogenica		0.21	0.00		1	1		
63510200	**Good identification**							**63510200**
Actinomyces odontolyticus		100.0	0.35	ESL 1				
63510522	**Good identification**							**63510522**
					GRA	CAT		
Capnocytophaga ochracea		99.51	0.37	MNS 90	1	90		
Bifidobacterium longum subsp. longum		0.49	0.05	GAL 99	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63511001	**Low discrimination**						**63511001**
					SPO		
Actinomyces naeslundii		59.45	0.41	ESL 12	1		
Clostridium chauvoei		40.56	0.15	MNS 88	99		
63511021	**Low discrimination**						**63511021**
					GRA	SPO	
Actinomyces naeslundii		75.10	0.41	ESL 12	99	1	
Clostridium chauvoei		17.08	0.07	MNS 88	99	99	
Capnocytophaga ochracea		7.82	-.01		1	1	
63511022	**Good identification**						**63511022**
					GRA	CAT	
Capnocytophaga ochracea		98.19	0.50	MNS 90	1	90	
Actinomyces naeslundii		0.71	0.38	LAC 99	99	25	
Clostridium perfringens		0.30	0.10		99	1	
63511023	**Low discrimination**						**63511023**
					GRA		
Actinomyces naeslundii		51.62	0.41	ESL 12	99		
Capnocytophaga ochracea		48.38	0.15	MNS 88	1		
63511100	**Acceptable identification**						**63511100**
					GRA	CAT	
Capnocytophaga ochracea		90.96	0.51	MNS 90	1	90	
Fusobacterium mortiferum		3.42	0.40	NAG 90	1	5	
Actinomyces naeslundii		1.52	0.46	BGL 90	99	25	
63511101	**Acceptable identification**						**63511101**
					GRA	SPO	
Clostridium chauvoei		94.42	0.50	MNS 90	99	99	
Actinomyces naeslundii		3.98	0.49	RAF 1	99	1	
Capnocytophaga ochracea		1.60	0.17		1	1	
63511102	**Very good identification**						**63511102**
					GRA		
Capnocytophaga ochracea		97.79	0.67	MNS 90	1		
Actinomyces viscosus		1.73	0.39	NAG 90	99		
Actinomyces naeslundii		0.18	0.46		99		
63511103	**Low discrimination**						**63511103**
					GRA	SPO	
Clostridium chauvoei		38.25	0.37	MNS 90	99	99	
Capnocytophaga ochracea		33.08	0.33	RAF 1	1	1	
Actinomyces viscosus		19.28	0.30		99	1	
63511112	**Good identification**						**63511112**
					GRA	CAT	
Capnocytophaga ochracea		99.64	0.51	MNS 90	1	90	
Actinomyces viscosus		0.16	0.04	MLZ 10	99	76	
Eubacterium saburreum		0.11	0.13	NAG 90	85	1	
63511120	**Good identification**						**63511120**
					GRA	CAT	
Capnocytophaga ochracea		90.26	0.67	MNS 90	1	90	
Prevotella melaninogenica		9.11	0.47	BGL 90	1	1	
Clostridium perfringens		0.39	0.30		99	1	
63511121	**Low discrimination**						**63511121**
					GRA	SPO	
Clostridium chauvoei		61.33	0.41	MNS 90	99	99	
Capnocytophaga ochracea		28.09	0.33	RAF 1	1	1	

Code / Taxon		ID score	T index	tests against		Differentiation tests		Code
Actinomyces naeslundii		7.75	0.49			99	1	
63511122	**Very good identification**							**63511122**
						GRA	CAT	
Capnocytophaga ochracea		99.71	0.84	MNS	90	1	90	
Prevotella melaninogenica		0.21	0.34			1	1	
Actinomyces naeslundii		0.02	0.46			99	25	
63511123	**Acceptable identification**							**63511123**
						GRA	SPO	
Capnocytophaga ochracea		92.97	0.50	MNS	90	1	1	
Clostridium chauvoei		3.98	0.28	URE	1	99	99	
Actinomyces naeslundii		2.85	0.49			99	1	
63511130	**Good identification**							**63511130**
						GRA	CAT	
Capnocytophaga ochracea		98.78	0.51	MNS	90	1	90	
Prevotella melaninogenica		0.91	0.12	MLZ	10	1	1	
Bifidobacterium longum subsp. longum		0.12	0.09	BGL	90	99	1	
63511132	**Very good identification**							**63511132**
						GRA	CAT	
Capnocytophaga ochracea		99.96	0.67	MNS	90	1	90	
Prevotella melaninogenica		0.02	0.00	MLZ	10	1	1	
Actinomyces naeslundii		0.01	0.24			99	25	
63511142	**Good identification**							**63511142**
						GRA	CAT	
Capnocytophaga ochracea		98.91	0.51	MNS	90	1	90	
Propionibacterium propionicum		0.64	0.29	NAG	90	99	1	
Actinomyces naeslundii		0.29	0.33	SOR	10	99	25	
63511160	**Good identification**							**63511160**
						GRA	CAT	
Capnocytophaga ochracea		98.08	0.51	MNS	90	1	90	
Prevotella melaninogenica		0.90	0.12	SOR	10	1	1	
Clostridium perfringens		0.73	0.18	BGL	90	99	1	
63511162	**Very good identification**							**63511162**
						GRA	CAT	
Capnocytophaga ochracea		99.91	0.67	MNS	90	1	90	
Clostridium perfringens		0.04	0.12	SOR	10	99	1	
Actinomyces naeslundii		0.03	0.33			99	25	
63511172	**Good identification**							**63511172**
Capnocytophaga ochracea		100.0	0.51	MNS	90			
63511322	**Good identification**							**63511322**
						CAT		
Capnocytophaga ochracea		99.05	0.50	MNS	90	90		
Prevotella oralis		0.64	0.12	RHA	1	1		
Prevotella melaninogenica		0.21	0.00			1		
63511422	**Good identification**							**63511422**
						GRA	CAT	
Capnocytophaga ochracea		98.20	0.37	MNS	90	1	90	
Lactobacillus fermentum		1.31	0.20	LAC	99	99	1	
Bifidobacterium longum subsp. longum		0.49	0.05			99	1	
63511500	**Low discrimination**							**63511500**
						GRA	CAT	
Capnocytophaga ochracea		86.70	0.38	MNS	90	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		8.67	0.29	NAG 90	99	1	
Lactobacillus fermentum		4.63	0.32	BGL 90	99	1	
63511502	**Good identification**						**63511502**
					GRA	CAT	
Capnocytophaga ochracea		98.05	0.54	MNS 90	1	90	
Bifidobacterium longum subsp. longum		1.09	0.29	NAG 90	99	1	
Lactobacillus fermentum		0.58	0.32		99	1	
63511512	**Low discrimination**						**63511512**
					GRA	CAT	
Capnocytophaga ochracea		83.99	0.38	MNS 90	1	90	
Bifidobacterium longum subsp. longum		14.30	0.33	MLZ 10	99	1	
Bifidobacterium adolescentis		1.71	0.19	NAG 90	99	1	
63511520	**Good identification**						**63511520**
					GRA	CAT	
Capnocytophaga ochracea		95.04	0.54	MNS 90	1	90	
Bifidobacterium longum subsp. longum		4.22	0.40	BGL 90	99	1	
Prevotella melaninogenica		0.55	0.12		1	1	
63511522	**Very good identification**						**63511522**
					GRA	CAT	
Capnocytophaga ochracea		99.48	0.71	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.49	0.40		99	1	
Lactobacillus fermentum		0.02	0.24		99	1	
63511523	**Good identification**						**63511523**
					GRA	CAT	
Capnocytophaga ochracea		99.51	0.37	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.49	0.05	URE 1	99	1	
63511530	**Low discrimination**						**63511530**
					GRA	CAT	
Capnocytophaga ochracea		59.49	0.38	MNS 90	1	90	
Bifidobacterium longum subsp. longum		40.52	0.44	MLZ 10	99	1	
63511532	**Acceptable identification**						**63511532**
					GRA	CAT	
Capnocytophaga ochracea		92.97	0.54	MNS 90	1	90	
Bifidobacterium longum subsp. longum		7.04	0.44	MLZ 10	99	1	
63511542	**Good identification**						**63511542**
Capnocytophaga ochracea		100.0	0.38	MNS 90			
63511560	**Good identification**						**63511560**
					GRA	CAT	
Capnocytophaga ochracea		99.60	0.38	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.40	0.05	SOR 10	99	1	
63511562	**Good identification**						**63511562**
					GRA	CAT	
Capnocytophaga ochracea		99.96	0.54	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.05	0.05	SOR 10	99	1	
63511572	**Good identification**						**63511572**
					GRA	CAT	
Capnocytophaga ochracea		99.32	0.38	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.68	0.09	MLZ 10	99	1	
63511722	**Good identification**						**63511722**
					GRA	CAT	
Capnocytophaga ochracea		99.51	0.37	MNS 90	1	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		0.49	0.05	RHA 1	99	1	
63512100	**Very good identification**						**63512100**
Propionibacterium propionicum		99.68	0.60	ESL 1			
Actinomyces naeslundii		0.32	0.27				
63512101	**Good identification**						**63512101**
Propionibacterium propionicum		95.88	0.44	ESL 1			
Actinomyces naeslundii		4.12	0.30	URE 10			
63512102	**Very good identification**						**63512102**
Propionibacterium propionicum		99.24	0.63	ESL 1			
Actinomyces viscosus		0.41	0.08				
Actinomyces naeslundii		0.21	0.27				
63512103	**Good identification**						**63512103**
Propionibacterium propionicum		96.06	0.47	ESL 1			
Actinomyces naeslundii		2.75	0.30	URE 10			
Actinomyces viscosus		1.18	0.00				
63512140	**Very good identification**						**63512140**
Propionibacterium propionicum		100.0	0.60	ESL 1			
63512141	**Good identification**						**63512141**
Propionibacterium propionicum		100.0	0.44	ESL 1			
63512142	**Very good identification**						**63512142**
Propionibacterium propionicum		100.0	0.63	ESL 1			
63512143	**Good identification**						**63512143**
Propionibacterium propionicum		100.0	0.47	ESL 1			
63513000	**Low discrimination**						**63513000**
Actinomyces naeslundii		72.66	0.43	ESL 12			
Propionibacterium propionicum		27.34	0.26	MNS 88			
63513001	**Good identification**						**63513001**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63513002	**Low discrimination**						**63513002**
					GRA	CAT	
Actinomyces naeslundii		58.01	0.43	ESL 12	99	25	
Propionibacterium propionicum		32.74	0.29	MNS 88	99	1	
Capnocytophaga ochracea		4.67	-.01		1	90	
63513003	**Good identification**						**63513003**
Actinomyces naeslundii		95.99	0.46	ESL 12			
Propionibacterium propionicum		4.01	0.12	MNS 88			
63513020	**Acceptable identification**						**63513020**
					GRA		
Actinomyces naeslundii		92.55	0.43	ESL 12	99		
Capnocytophaga ochracea		7.45	-.01	MNS 88	1		
63513021	**Good identification**						**63513021**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.46	ESL 12			
63513022	**Low discrimination**						**63513022**
					GRA		
Actinomyces naeslundii		58.00	0.43	ESL 12	99		
Capnocytophaga ochracea		42.00	0.15	MNS 88	1		
63513023	**Good identification**						**63513023**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63513100	**Good identification**						**63513100**
					GRA	CAT	
Propionibacterium propionicum		90.57	0.60	ESL 1	99	1	
Actinomyces naeslundii		6.92	0.51		99	25	
Capnocytophaga ochracea		2.15	0.17		1	90	
63513101	**Low discrimination**						**63513101**
					SPO		
Actinomyces naeslundii		47.78	0.54	ESL 12	1		
Propionibacterium propionicum		46.32	0.44	MNS 88	1		
Clostridium chauvoei		5.90	0.15		99		
63513102	**Low discrimination**						**63513102**
					GRA	CAT	
Propionibacterium propionicum		83.34	0.63	ESL 1	99	1	
Capnocytophaga ochracea		11.88	0.33		1	90	
Actinomyces naeslundii		4.25	0.51		99	25	
63513103	**Low discrimination**						**63513103**
					SPO		
Propionibacterium propionicum		58.29	0.47	ESL 1	1		
Actinomyces naeslundii		40.08	0.54	URE 10	1		
Clostridium chauvoei		0.87	0.03		99		
63513121	**Acceptable identification**						**63513121**
					GRA	SPO	
Actinomyces naeslundii		94.34	0.54	ESL 12	99	1	
Clostridium chauvoei		3.88	0.07	MNS 88	99	99	
Capnocytophaga ochracea		1.78	-.01		1	1	
63513122	**Good identification**						**63513122**
					GRA	CAT	
Capnocytophaga ochracea		95.27	0.50	MNS 90	1	90	
Actinomyces naeslundii		3.78	0.51	MAN 1	99	25	
Propionibacterium propionicum		0.75	0.29		99	1	
63513123	**Low discrimination**						**63513123**
					GRA	CAT	
Actinomyces naeslundii		84.44	0.54	ESL 12	99	25	
Capnocytophaga ochracea		14.32	0.15	MNS 88	1	90	
Propionibacterium propionicum		1.24	0.12		99	1	
63513140	**Very good identification**						**63513140**
					GRA	CAT	
Propionibacterium propionicum		98.41	0.60	ESL 1	99	1	
Actinomyces naeslundii		1.33	0.38		99	25	
Capnocytophaga ochracea		0.26	0.01		1	90	
63513141	**Low discrimination**						**63513141**
Propionibacterium propionicum		84.60	0.44	ESL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		15.40	0.41	URE 10			
63513142	**Very good identification**						**63513142**
					GRA	CAT	
Propionibacterium propionicum		97.43	0.63	ESL 1	99	1	
Capnocytophaga ochracea		1.54	0.17		1	90	
Actinomyces naeslundii		0.88	0.38		99	25	
63513143	**Low discrimination**						**63513143**
Propionibacterium propionicum		89.18	0.47	ESL 1			
Actinomyces naeslundii		10.82	0.41	URE 10			
63513161	**Good identification**						**63513161**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63513163	**Low discrimination**						**63513163**
					GRA	CAT	
Actinomyces naeslundii		84.03	0.41	ESL 12	99	25	
Capnocytophaga ochracea		8.97	-.01	MNS 88	1	90	
Propionibacterium propionicum		6.99	0.12		99	1	
63513522	**Good identification**						**63513522**
					GRA	CAT	
Capnocytophaga ochracea		99.51	0.37	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.49	0.05	MAN 1	99	1	
63514122	**Good identification**						**63514122**
Capnocytophaga ochracea		96.25	0.39	MNS 90			
Bacteroides fragilis		3.75	0.06	GAL 99			
63515022	**Good identification**						**63515022**
					GRA	CAT	
Capnocytophaga ochracea		95.74	0.39	MNS 90	1	90	
Bacteroides fragilis		3.73	0.06	LAC 99	1	99	
Lactobacillus fermentum		0.53	0.16		99	1	
63515100	**Acceptable identification**						**63515100**
					CAT	GRA	
Capnocytophaga ochracea		93.26	0.41	MNS 90	90	1	
Lactobacillus fermentum		2.07	0.28	NAG 90	1	99	
Eubacterium saburreum		1.57	0.23	BGL 90	1	85	
63515102	**Very good identification**						**63515102**
					GRA	CAT	
Capnocytophaga ochracea		98.89	0.57	MNS 90	1	90	
Bacteroides fragilis		0.35	0.06	NAG 90	1	99	
Lactobacillus fermentum		0.24	0.28		99	1	
63515112	**Good identification**						**63515112**
					GRA	CAT	
Capnocytophaga ochracea		97.94	0.41	MNS 90	1	90	
Bifidobacterium adolescentis		1.41	0.19	MLZ 10	99	1	
Bifidobacterium longum subsp. longum		0.36	0.07	NAG 90	99	1	
63515120	**Very good identification**						**63515120**
					CAT		
Capnocytophaga ochracea		97.54	0.57	MNS 90	90		
Bacteroides fragilis		1.80	0.18	BGL 90	99		
Prevotella melaninogenica		0.40	0.12		1		
63515122	**Very good identification**						**63515122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		96.20	0.73	MNS 90	1	90	
Bacteroides fragilis		3.75	0.40		1	99	
Lactobacillus fermentum		0.01	0.20		99	1	
63515123	**Good identification**						**63515123**
Capnocytophaga ochracea		96.25	0.39	MNS 90			
Bacteroides fragilis		3.75	0.06	URE 1			
63515130	**Good identification**						**63515130**
					GRA	CAT	
Capnocytophaga ochracea		98.56	0.41	MNS 90	1	90	
Bifidobacterium longum subsp. longum		1.44	0.18	MLZ 10	99	1	
63515132	**Very good identification**						**63515132**
					GRA	CAT	
Capnocytophaga ochracea		99.44	0.57	MNS 90	1	90	
Bacteroides fragilis		0.35	0.06	MLZ 10	1	99	
Bifidobacterium longum subsp. longum		0.16	0.18		99	1	
63515142	**Good identification**						**63515142**
					GRA	CAT	
Capnocytophaga ochracea		99.21	0.41	MNS 90	1	90	
Actinomyces israelii		0.79	0.21	NAG 90	99	1	
63515160	**Good identification**						**63515160**
Capnocytophaga ochracea		100.0	0.41	MNS 90			
63515162	**Very good identification**						**63515162**
Capnocytophaga ochracea		99.65	0.57	MNS 90			
Bacteroides fragilis		0.35	0.06	SOR 10			
63515172	**Good identification**						**63515172**
Capnocytophaga ochracea		100.0	0.41	MNS 90			
63515322	**Acceptable identification**						**63515322**
Capnocytophaga ochracea		93.27	0.39	MNS 90			
Bacteroides fragilis		3.64	0.06	RHA 1			
Parabacteroides distasonis		2.61	0.25				
63515500	**Low discrimination**						**63515500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		50.24	0.37	NIT 1	99	1	
Capnocytophaga ochracea		41.87	0.28	MNS 87	1	90	
Lactobacillus fermentum		5.26	0.28		99	1	
63515502	**Low discrimination**						**63515502**
					GRA	CAT	
Capnocytophaga ochracea		82.82	0.44	MNS 90	1	90	
Bifidobacterium longum subsp. longum		11.04	0.37	NAG 90	99	1	
Bifidobacterium adolescentis		4.11	0.32		99	1	
63515510	**Acceptable identification**						**63515510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		94.84	0.41	NIT 1	99	1	
Capnocytophaga ochracea		5.16	0.12	MNS 87	1	90	
63515512	**Low discrimination**						**63515512**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		56.93	0.41	NIT 1	99	1	
Capnocytophaga ochracea		27.86	0.28	MNS 87	1	90	
Bifidobacterium adolescentis		15.21	0.33		99	1	
63515520	**Low discrimination**						**63515520**
					GRA	CAT	
Capnocytophaga ochracea		63.12	0.44	MNS 90	1	90	
Bifidobacterium longum subsp. longum		33.66	0.48	BGL 90	99	1	
Bacteroides vulgatus		2.93	0.15		1	3	
63515522	**Good identification**						**63515522**
					GRA	CAT	
Capnocytophaga ochracea		94.13	0.61	MNS 90	1	90	
Bifidobacterium longum subsp. longum		5.58	0.48		99	1	
Bacteroides fragilis		0.21	0.06		1	99	
63515530	**Low discrimination**						**63515530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		89.10	0.52	NIT 1	99	1	
Capnocytophaga ochracea		10.90	0.28	MNS 87	1	90	
63515532	**Low discrimination**						**63515532**
					GRA	CAT	
Capnocytophaga ochracea		52.41	0.44	MNS 90	1	90	
Bifidobacterium longum subsp. longum		47.59	0.52	MLZ 10	99	1	
63515562	**Good identification**						**63515562**
					GRA	CAT	
Capnocytophaga ochracea		99.46	0.44	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.54	0.14	SOR 10	99	1	
63517100	**Low discrimination**						**63517100**
					GRA	CAT	
Actinomyces israelii		73.84	0.38	SAL 92	99	1	
Propionibacterium propionicum		16.47	0.26	CEL 93	99	1	
Capnocytophaga ochracea		9.69	0.07		1	90	
63517102	**Low discrimination**						**63517102**
					GRA	CAT	
Actinomyces israelii		47.76	0.41	SAL 92	99	1	
Capnocytophaga ochracea		37.60	0.23	CEL 93	1	90	
Propionibacterium propionicum		10.66	0.29		99	1	
63517122	**Good identification**						**63517122**
					GRA	CAT	
Capnocytophaga ochracea		95.42	0.39	MNS 90	1	90	
Bacteroides fragilis		3.72	0.06	MAN 1	1	99	
Actinomyces israelii		0.86	0.20		99	1	
63517500	**Good identification**						**63517500**
Actinomyces israelii		100.0	0.36	SAL 92			
63517502	**Low discrimination**						**63517502**
					GRA	CAT	
Actinomyces israelii		53.99	0.39	SAL 92	99	1	
Bifidobacterium adolescentis		36.84	0.30	CEL 93	99	1	
Capnocytophaga ochracea		9.17	0.10		1	90	
63523101	**Good identification**						**63523101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63523103	**Good identification**						**63523103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	ESL 12			
63523121	**Good identification**						**63523121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63523123	**Good identification**						**63523123**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63531000	**Low discrimination**						**63531000**
					GRA		
Actinomyces naeslundii		61.81	0.43	ESL 12	99		
Fusobacterium mortiferum		26.10	0.25	MNS 88	1		
Eubacterium saburreum		12.09	0.18		85		
63531001	**Good identification**						**63531001**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63531002	**Low discrimination**						**63531002**
					GRA		
Actinomyces naeslundii		59.88	0.43	ESL 12	99		
Actinomyces viscosus		35.31	0.15	MNS 88	99		
Capnocytophaga ochracea		4.82	-.01		1		
63531003	**Low discrimination**						**63531003**
Actinomyces naeslundii		88.42	0.46	ESL 12			
Actinomyces viscosus		11.58	0.07	MNS 88			
63531020	**Low discrimination**						**63531020**
					GRA	CAT	
Actinomyces naeslundii		51.23	0.43	ESL 12	99	25	
Clostridium perfringens		44.65	0.20	MNS 88	99	1	
Capnocytophaga ochracea		4.12	-.01		1	90	
63531021	**Good identification**						**63531021**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63531022	**Low discrimination**						**63531022**
					GRA	CAT	
Actinomyces naeslundii		46.85	0.43	ESL 12	99	25	
Capnocytophaga ochracea		33.93	0.15	MNS 88	1	90	
Clostridium perfringens		19.22	0.14		99	1	
63531023	**Good identification**						**63531023**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63531100	**Low discrimination**						**63531100**
					GRA	CAT	
Actinomyces naeslundii		49.28	0.51	ESL 12	99	25	
Fusobacterium mortiferum		17.06	0.32	MNS 88	1	5	
Capnocytophaga ochracea		15.33	0.17		1	90	
63531103	**Low discrimination**						**63531103**
					SPO		
Actinomyces naeslundii		74.61	0.54	ESL 12	1		
Clostridium chauvoei		17.88	0.20	MNS 88	99		
Actinomyces viscosus		6.11	0.11		1		
63531121	**Low discrimination**						**63531121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Actinomyces naeslundii		67.26	0.54	ESL 12	99	1	
Clostridium chauvoei		30.45	0.25	MNS 88	99	99	
Capnocytophaga ochracea		1.27	-.01		1	1	
63531122	**Acceptable identification**						**63531122**
					GRA	CAT	
Capnocytophaga ochracea		92.09	0.50	MNS 90	1	90	
Clostridium perfringens		3.86	0.29	TRE 1	99	1	
Actinomyces naeslundii		3.66	0.51		99	25	
63531123	**Low discrimination**						**63531123**
					GRA	SPO	
Actinomyces naeslundii		80.03	0.54	ESL 12	99	1	
Capnocytophaga ochracea		13.58	0.15	MNS 88	1	1	
Clostridium chauvoei		6.39	0.12		99	99	
63531140	**Low discrimination**						**63531140**
					GRA	CAT	
Actinomyces naeslundii		55.29	0.38	ESL 12	99	25	
Propionibacterium propionicum		33.89	0.24	MNS 88	99	1	
Capnocytophaga ochracea		10.82	0.01		1	90	
63531141	**Good identification**						**63531141**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63531143	**Good identification**						**63531143**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63531161	**Good identification**						**63531161**
Actinomyces naeslundii		100.0	0.41	ESL 12			
63531163	**Acceptable identification**						**63531163**
					GRA		
Actinomyces naeslundii		90.35	0.41	ESL 12	99		
Capnocytophaga ochracea		9.65	-.01	MNS 88	1		
63531522	**Low discrimination**						**63531522**
					GRA	CAT	
Capnocytophaga ochracea		89.11	0.37	MNS 90	1	90	
Bifidobacterium longum subsp. longum		10.89	0.29	TRE 1	99	1	
63532100	**Very good identification**						**63532100**
Propionibacterium propionicum		99.25	0.58	ESL 1			
Actinomyces naeslundii		0.75	0.32				
63532101	**Acceptable identification**						**63532101**
Propionibacterium propionicum		90.75	0.42	ESL 1			
Actinomyces naeslundii		9.25	0.35	URE 10			
63532102	**Very good identification**						**63532102**
Propionibacterium propionicum		99.50	0.61	ESL 1			
Actinomyces naeslundii		0.50	0.32				
63532103	**Acceptable identification**						**63532103**
Propionibacterium propionicum		93.64	0.45	ESL 1			
Actinomyces naeslundii		6.37	0.35	URE 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
63532121	**Good identification**					**63532121**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63532123	**Good identification**					**63532123**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63532140	**Very good identification**					**63532140**
Propionibacterium propionicum		100.0	0.58	ESL 1		
63532141	**Good identification**					**63532141**
Propionibacterium propionicum		98.23	0.42	ESL 1		
Actinomyces naeslundii		1.77	0.22	URE 10		
63532142	**Very good identification**					**63532142**
Propionibacterium propionicum		100.0	0.61	ESL 1		
63532143	**Good identification**					**63532143**
Propionibacterium propionicum		98.82	0.45	ESL 1		
Actinomyces naeslundii		1.19	0.22	URE 10		
63533000	**Low discrimination**					**63533000**
Actinomyces naeslundii		81.33	0.48	ESL 12		
Propionibacterium propionicum		12.90	0.24	MNS 88		
Actinomyces israelii		5.78	0.19			
63533001	**Good identification**					**63533001**
Actinomyces naeslundii		100.0	0.51	ESL 12		
63533002	**Low discrimination**					**63533002**
Actinomyces naeslundii		74.38	0.48	ESL 12		
Propionibacterium propionicum		17.69	0.27	MNS 88		
Actinomyces israelii		7.93	0.22			
63533003	**Good identification**					**63533003**
Actinomyces naeslundii		100.0	0.51	ESL 12		
63533020	**Good identification**					**63533020**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63533021	**Good identification**					**63533021**
Actinomyces naeslundii		100.0	0.51	ESL 12		
63533022	**Good identification**					**63533022**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63533023	**Good identification**					**63533023**
Actinomyces naeslundii		100.0	0.51	ESL 12		
63533040	**Low discrimination**					**63533040**
Actinomyces naeslundii		52.67	0.35	ESL 12		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		47.33	0.24	MNS 88			
63533041	**Good identification**						**63533041**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63533043	**Good identification**						**63533043**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63533060	**Good identification**						**63533060**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63533061	**Good identification**						**63533061**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63533062	**Good identification**						**63533062**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63533063	**Good identification**						**63533063**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63533100	**Low discrimination**						**63533100**
Propionibacterium propionicum		83.38	0.58	ESL 1			
Actinomyces naeslundii		15.11	0.56				
Actinomyces israelii		1.51	0.30				
63533101	**Low discrimination**						**63533101**
Actinomyces naeslundii		70.99	0.59	ESL 12			
Propionibacterium propionicum		29.01	0.42	MNS 88			
63533102	**Low discrimination**						**63533102**
Propionibacterium propionicum		87.66	0.61	ESL 1			
Actinomyces naeslundii		10.60	0.56				
Actinomyces israelii		1.59	0.33				
63533103	**Low discrimination**						**63533103**
Actinomyces naeslundii		62.00	0.59	ESL 12			
Propionibacterium propionicum		38.00	0.45	MNS 88			
63533111	**Good identification**						**63533111**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63533113	**Good identification**						**63533113**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63533120	**Acceptable identification**						**63533120**
					GRA	CAT	
Actinomyces naeslundii		92.43	0.56	ESL 12	99	25	
Propionibacterium propionicum		5.15	0.24	MNS 88	99	1	
Capnocytophaga ochracea		1.35	-.01		1	90	
63533121	**Very good identification**						**63533121**
Actinomyces naeslundii		100.0	0.59	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63533122	**Low discrimination**						**63533122**
					GRA	CAT	
Actinomyces naeslundii		82.33	0.56	ESL 12	99	25	
Capnocytophaga ochracea		10.79	0.15	MNS 88	1	90	
Propionibacterium propionicum		6.88	0.27		99	1	
63533123	**Very good identification**						**63533123**
Actinomyces naeslundii		100.0	0.59	ESL 12			
63533131	**Good identification**						**63533131**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63533133	**Good identification**						**63533133**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63533140	**Very good identification**						**63533140**
Propionibacterium propionicum		96.24	0.58	ESL 1			
Actinomyces naeslundii		3.08	0.43				
Actinomyces israelii		0.68	0.23				
63533141	**Low discrimination**						**63533141**
Propionibacterium propionicum		69.84	0.42	ESL 1			
Actinomyces naeslundii		30.16	0.46	URE 10			
63533142	**Very good identification**						**63533142**
Propionibacterium propionicum		97.24	0.61	ESL 1			
Actinomyces naeslundii		2.07	0.43				
Actinomyces israelii		0.69	0.26				
63533143	**Low discrimination**						**63533143**
Propionibacterium propionicum		77.65	0.45	ESL 1			
Actinomyces naeslundii		22.36	0.46	URE 10			
63533160	**Low discrimination**						**63533160**
Actinomyces naeslundii		76.00	0.43	ESL 12			
Propionibacterium propionicum		24.00	0.24	MNS 88			
63533161	**Good identification**						**63533161**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63533162	**Low discrimination**						**63533162**
					GRA	CAT	
Actinomyces naeslundii		64.26	0.43	ESL 12	99	25	
Propionibacterium propionicum		30.44	0.27	MNS 88	99	1	
Capnocytophaga ochracea		5.30	-.01		1	90	
63533163	**Good identification**						**63533163**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63535122	**Acceptable identification**						**63535122**
Capnocytophaga ochracea		91.96	0.39	MNS 90			
Parabacteroides distasonis		3.91	0.28	TRE 1			
Bacteroides fragilis		3.59	0.06				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63535322	**Acceptable identification**						**63535322**
Parabacteroides distasonis		92.66	0.36	NIT 1			
Capnocytophaga ochracea		7.34	0.05	MNS 98			
63535502	**Low discrimination**						**63535502**
Actinomyces israelii		34.04	0.36	SAL 92			
Bifidobacterium longum subsp. longum		28.74	0.27	CEL 93			
Bifidobacterium adolescentis		28.51	0.29				
63535520	**Acceptable identification**						**63535520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		91.72	0.37	NIT 1	99	1	
Capnocytophaga ochracea		6.95	0.10	MNS 87	1	90	
Bacteroides vulgatus		1.33	-.08		1	3	
63535522	**Low discrimination**						**63535522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		55.52	0.37	NIT 1	99	1	
Capnocytophaga ochracea		37.85	0.26	MNS 87	1	90	
Parabacteroides distasonis		5.59	0.25		1	76	
63535530	**Good identification**						**63535530**
Bifidobacterium longum subsp. longum		100.0	0.41	NIT 1			
63535532	**Acceptable identification**						**63535532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.91	0.41	NIT 1	99	1	
Capnocytophaga ochracea		4.13	0.10	MNS 87	1	90	
Parabacteroides distasonis		2.96	0.20		1	76	
63537000	**Good identification**						**63537000**
Actinomyces israelii		100.0	0.37	SAL 92			
63537002	**Good identification**						**63537002**
Actinomyces israelii		100.0	0.40	SAL 92			
63537100	**Good identification**						**63537100**
Actinomyces israelii		95.37	0.48	SAL 92			
Propionibacterium propionicum		4.63	0.24	CEL 93			
63537102	**Acceptable identification**						**63537102**
Actinomyces israelii		94.05	0.51	SAL 92			
Propionibacterium propionicum		4.56	0.27	CEL 93			
Bifidobacterium adolescentis		1.39	0.13				
63537112	**Low discrimination**						**63537112**
Actinomyces israelii		89.21	0.37	SAL 92			
Bifidobacterium adolescentis		10.79	0.15	CEL 93			
63537140	**Low discrimination**						**63537140**
Actinomyces israelii		88.91	0.41	SAL 92			
Propionibacterium propionicum		11.09	0.24	CEL 93			
63537142	**Low discrimination**						**63537142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		88.91	0.44	SAL 92			
Propionibacterium propionicum		11.09	0.27	CEL 93			
63537400	**Good identification**						**63537400**
Actinomyces israelii		100.0	0.36	SAL 92			
63537402	**Good identification**						**63537402**
Actinomyces israelii		100.0	0.39	SAL 92			
63537500	**Good identification**						**63537500**
Actinomyces israelii		100.0	0.46	SAL 92			
63537502	**Low discrimination**						**63537502**
Actinomyces israelii		89.21	0.49	SAL 92			
Bifidobacterium adolescentis		10.79	0.27	CEL 93			
63537512	**Low discrimination**						**63537512**
Actinomyces israelii		50.27	0.35	SAL 92			
Bifidobacterium adolescentis		49.73	0.29	CEL 93			
63537540	**Good identification**						**63537540**
Actinomyces israelii		100.0	0.39	SAL 92			
63537542	**Good identification**						**63537542**
Actinomyces israelii		96.30	0.42	SAL 92			
Bifidobacterium adolescentis		3.70	0.12	CEL 93			
63541122	**Good identification**						**63541122**
					GRA	CAT	
Capnocytophaga ochracea		98.41	0.39	MNS 90	1	90	
Prevotella melaninogenica		1.03	0.02	FRU 99	1	1	
Actinomyces naeslundii		0.56	0.26		99	25	
63550122	**Good identification**						**63550122**
					GRA		
Capnocytophaga ochracea		99.21	0.39	MNS 90	1		
Actinomyces naeslundii		0.45	0.25	GAL 99	99		
Actinomyces viscosus		0.34	-.02		99		
63551000	**Low discrimination**						**63551000**
					GRA	CAT	
Actinomyces naeslundii		46.73	0.40	ESL 12	99	25	
Fusobacterium mortiferum		32.24	0.26	MNS 88	1	5	
Capnocytophaga ochracea		15.14	0.07		1	90	
63551001	**Good identification**						**63551001**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63551003	**Low discrimination**						**63551003**
Actinomyces naeslundii		59.61	0.43	ESL 12			
Actinomyces viscosus		40.39	0.16	MNS 88			
63551021	**Good identification**						**63551021**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63551022	**Good identification**						**63551022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		95.09	0.39	MNS 90	1	90	
Actinomyces naeslundii		3.62	0.40	LAC 99	99	25	
Prevotella oralis		1.29	0.07		1	1	
63551023	**Low discrimination**						**63551023**
					GRA		
Actinomyces naeslundii		84.98	0.43	ESL 12	99		
Capnocytophaga ochracea		15.02	0.05	MNS 88	1		
63551100	**Low discrimination**						**63551100**
					GRA	CAT	
Capnocytophaga ochracea		86.31	0.41	MNS 90	1	90	
Actinomyces naeslundii		7.66	0.48	NAG 90	99	25	
Fusobacterium mortiferum		4.33	0.32	BGL 90	1	5	
63551101	**Low discrimination**						**63551101**
					GRA	SPO	
Actinomyces naeslundii		79.57	0.51	ESL 12	99	1	
Clostridium chauvoei		14.39	0.15	MNS 88	99	99	
Capnocytophaga ochracea		6.04	0.07		1	1	
63551102	**Very good identification**						**63551102**
					GRA		
Capnocytophaga ochracea		96.78	0.57	MNS 90	1		
Actinomyces viscosus		1.82	0.29	NAG 90	99		
Actinomyces naeslundii		0.95	0.48		99		
63551103	**Low discrimination**						**63551103**
					GRA		
Actinomyces naeslundii		46.77	0.51	ESL 12	99		
Capnocytophaga ochracea		31.95	0.23	MNS 88	1		
Actinomyces viscosus		19.80	0.21		99		
63551112	**Good identification**						**63551112**
					GRA	CAT	
Capnocytophaga ochracea		98.74	0.41	MNS 90	1	90	
Actinomyces naeslundii		0.46	0.26	MLZ 10	99	25	
Bifidobacterium adolescentis		0.45	0.11	NAG 90	99	1	
63551120	**Good identification**						**63551120**
					GRA	CAT	
Capnocytophaga ochracea		94.66	0.57	MNS 90	1	90	
Prevotella melaninogenica		4.25	0.30	BGL 90	1	1	
Actinomyces naeslundii		0.93	0.48		99	25	
63551121	**Low discrimination**						**63551121**
					GRA	SPO	
Actinomyces naeslundii		56.37	0.51	ESL 12	99	1	
Capnocytophaga ochracea		38.51	0.23	MNS 88	1	1	
Clostridium chauvoei		3.40	0.07		99	99	
63551122	**Very good identification**						**63551122**
					GRA	CAT	
Capnocytophaga ochracea		99.56	0.73	MNS 90	1	90	
Prevotella oralis		0.21	0.27		1	1	
Actinomyces naeslundii		0.11	0.48		99	25	
63551123	**Low discrimination**						**63551123**
					GRA		
Capnocytophaga ochracea		86.01	0.39	MNS 90	1		
Actinomyces naeslundii		13.99	0.51	URE 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63551130	**Good identification**						**63551130**
					GRA	CAT	
Capnocytophaga ochracea		99.13	0.41	MNS 90	1	90	
Actinomyces naeslundii		0.46	0.26	MLZ 10	99	25	
Prevotella melaninogenica		0.40	-.04	BGL 90	1	1	
63551132	**Very good identification**						**63551132**
					GRA		
Capnocytophaga ochracea		99.95	0.57	MNS 90	1		
Actinomyces naeslundii		0.05	0.26	MLZ 10	99		
63551141	**Good identification**						**63551141**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63551142	**Good identification**						**63551142**
					GRA	CAT	
Capnocytophaga ochracea		97.58	0.41	MNS 90	1	90	
Actinomyces naeslundii		1.53	0.35	NAG 90	99	25	
Actinomyces israelii		0.89	0.22	SOR 10	99	1	
63551143	**Low discrimination**						**63551143**
					GRA		
Actinomyces naeslundii		69.92	0.38	ESL 12	99		
Capnocytophaga ochracea		30.08	0.07	MNS 88	1		
63551160	**Good identification**						**63551160**
					GRA	CAT	
Capnocytophaga ochracea		98.07	0.41	MNS 90	1	90	
Actinomyces naeslundii		1.54	0.35	SOR 10	99	25	
Prevotella melaninogenica		0.40	-.04	BGL 90	1	1	
63551161	**Low discrimination**						**63551161**
					GRA		
Actinomyces naeslundii		69.92	0.38	ESL 12	99		
Capnocytophaga ochracea		30.08	0.07	MNS 88	1		
63551162	**Very good identification**						**63551162**
					GRA		
Capnocytophaga ochracea		99.83	0.57	MNS 90	1		
Actinomyces naeslundii		0.17	0.35	SOR 10	99		
63551172	**Good identification**						**63551172**
Capnocytophaga ochracea		100.0	0.41	MNS 90			
63551322	**Acceptable identification**						**63551322**
					CAT		
Capnocytophaga ochracea		91.65	0.39	MNS 90	90		
Prevotella oralis		8.35	0.21	RHA 1	1		
63551502	**Good identification**						**63551502**
					GRA	CAT	
Capnocytophaga ochracea		97.29	0.44	MNS 90	1	90	
Bifidobacterium adolescentis		1.53	0.23	NAG 90	99	1	
Actinomyces israelii		1.18	0.28		99	1	
63551520	**Good identification**						**63551520**
					GRA	CAT	
Capnocytophaga ochracea		99.57	0.44	MNS 90	1	90	
Prevotella melaninogenica		0.26	-.04	BGL 90	1	1	
Bifidobacterium longum subsp. longum		0.18	0.05		99	1	
63551522	**Very good identification**						**63551522**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		99.98	0.61	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.02	0.05		99	1	
63551532	**Good identification**						**63551532**
					GRA	CAT	
Capnocytophaga ochracea		99.70	0.44	MNS 90	1	90	
Bifidobacterium longum subsp. longum		0.31	0.09	MLZ 10	99	1	
63551562	**Good identification**						**63551562**
Capnocytophaga ochracea		100.0	0.44	MNS 90			
63553000	**Low discrimination**						**63553000**
Actinomyces naeslundii		73.13	0.45	ESL 12			
Actinomyces israelii		26.87	0.29	MNS 88			
63553001	**Good identification**						**63553001**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63553002	**Low discrimination**						**63553002**
Actinomyces naeslundii		64.47	0.45	ESL 12			
Actinomyces israelii		35.53	0.32	MNS 88			
63553003	**Good identification**						**63553003**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63553020	**Good identification**						**63553020**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63553021	**Good identification**						**63553021**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63553022	**Low discrimination**						**63553022**
					GRA		
Actinomyces naeslundii		87.98	0.45	ESL 12	99		
Capnocytophaga ochracea		12.02	0.05	MNS 88	1		
63553023	**Good identification**						**63553023**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63553041	**Good identification**						**63553041**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63553043	**Good identification**						**63553043**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63553061	**Good identification**						**63553061**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63553063	**Good identification**						**63553063**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63553100	**Low discrimination**						**63553100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		59.71	0.53	ESL 12			
Actinomyces israelii		30.84	0.39	MNS 88			
Propionibacterium propionicum		5.95	0.26				
63553101	**Good identification**						**63553101**
Actinomyces naeslundii		100.0	0.56	ESL 12			
63553102	**Low discrimination**						**63553102**
					GRA	CAT	
Actinomyces naeslundii		40.49	0.53	ESL 12	99	25	
Actinomyces israelii		31.36	0.42	MNS 88	99	1	
Capnocytophaga ochracea		21.37	0.23		1	90	
63553103	**Good identification**						**63553103**
Actinomyces naeslundii		98.90	0.56	ESL 12			
Propionibacterium propionicum		1.10	0.12	MNS 88			
63553120	**Low discrimination**						**63553120**
					GRA	CAT	
Actinomyces naeslundii		63.11	0.53	ESL 12	99	25	
Capnocytophaga ochracea		33.32	0.23	MNS 88	1	90	
Actinomyces israelii		2.08	0.18		99	1	
63553121	**Good identification**						**63553121**
Actinomyces naeslundii		100.0	0.56	ESL 12			
63553122	**Low discrimination**						**63553122**
					GRA	CAT	
Capnocytophaga ochracea		81.91	0.39	MNS 90	1	90	
Actinomyces naeslundii		17.24	0.53	MAN 1	99	25	
Actinomyces israelii		0.85	0.22		99	1	
63553123	**Good identification**						**63553123**
					GRA		
Actinomyces naeslundii		96.90	0.56	ESL 12	99		
Capnocytophaga ochracea		3.10	0.05	MNS 88	1		
63553141	**Good identification**						**63553141**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63553143	**Acceptable identification**						**63553143**
Actinomyces naeslundii		94.09	0.43	ESL 12			
Propionibacterium propionicum		5.91	0.12	MNS 88			
63553160	**Low discrimination**						**63553160**
					GRA		
Actinomyces naeslundii		75.05	0.40	ESL 12	99		
Capnocytophaga ochracea		24.95	0.07	MNS 88	1		
63553161	**Good identification**						**63553161**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63553163	**Good identification**						**63553163**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63553500	**Good identification**						**63553500**
Actinomyces israelii		100.0	0.37	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63553502	**Low discrimination**						**63553502**
					GRA	CAT	
Actinomyces israelii		74.93	0.40	SAL 92	99	1	
Bifidobacterium adolescentis		14.06	0.22	XYL 92	99	1	
Capnocytophaga ochracea		11.01	0.10		1	90	
63555002	**Low discrimination**						**63555002**
					GRA	CAT	
Actinomyces israelii		74.86	0.37	SAL 92	99	1	
Capnocytophaga ochracea		25.14	0.13		1	90	
63555102	**Low discrimination**						**63555102**
					GRA	CAT	
Capnocytophaga ochracea		87.91	0.47	MNS 90	1	90	
Actinomyces israelii		10.58	0.47	NAG 90	99	1	
Bifidobacterium adolescentis		0.98	0.23		99	1	
63555120	**Good identification**						**63555120**
					GRA	CAT	
Capnocytophaga ochracea		96.29	0.47	MNS 90	1	90	
Bacteroides fragilis		3.05	0.12	BGL 90	1	99	
Actinomyces israelii		0.49	0.24		99	1	
63555122	**Good identification**						**63555122**
					CAT		
Capnocytophaga ochracea		93.55	0.63	MNS 90	90		
Bacteroides fragilis		6.25	0.34		99		
Prevotella oralis		0.10	0.12		1		
63555132	**Good identification**						**63555132**
Capnocytophaga ochracea		99.40	0.47	MNS 90			
Bacteroides fragilis		0.60	-.01	MLZ 10			
63555140	**Low discrimination**						**63555140**
					GRA	CAT	
Actinomyces israelii		71.65	0.37	SAL 92	99	1	
Capnocytophaga ochracea		28.35	0.14		1	90	
63555162	**Good identification**						**63555162**
					GRA	CAT	
Capnocytophaga ochracea		99.10	0.47	MNS 90	1	90	
Bacteroides fragilis		0.60	-.01	SOR 10	1	99	
Actinomyces israelii		0.30	0.20		99	1	
63555402	**Acceptable identification**						**63555402**
					GRA	CAT	
Actinomyces israelii		93.25	0.35	SAL 92	99	1	
Capnocytophaga ochracea		6.75	0.00		1	90	
63555500	**Low discrimination**						**63555500**
					GRA	CAT	
Actinomyces israelii		77.00	0.43	SAL 92	99	1	
Capnocytophaga ochracea		23.00	0.18		1	90	
63555512	**Low discrimination**						**63555512**
					GRA	CAT	
Bifidobacterium adolescentis		71.68	0.39	NIT 1	99	1	
Capnocytophaga ochracea		15.45	0.18	MNS 87	1	90	
Actinomyces israelii		11.59	0.32		99	1	
63555522	**Good identification**						**63555522**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		98.99	0.50	MNS 90	1	90	
Actinomyces israelii		0.39	0.25		99	1	
Bacteroides fragilis		0.38	-.01		1	99	
63555540	**Acceptable identification**						**63555540**
					GRA	CAT	
Actinomyces israelii		92.14	0.36	SAL 92	99	1	
Capnocytophaga ochracea		7.87	0.01		1	90	
63555542	**Low discrimination**						**63555542**
					GRA	CAT	
Actinomyces israelii		57.07	0.39	SAL 92	99	1	
Capnocytophaga ochracea		29.23	0.18		1	90	
Bifidobacterium adolescentis		13.71	0.22		99	1	
63557000	**Good identification**						**63557000**
Actinomyces israelii		100.0	0.47	SAL 92			
63557002	**Good identification**						**63557002**
Actinomyces israelii		100.0	0.50	SAL 92			
63557012	**Good identification**						**63557012**
Actinomyces israelii		100.0	0.36	SAL 92			
63557040	**Good identification**						**63557040**
Actinomyces israelii		100.0	0.40	SAL 92			
63557042	**Good identification**						**63557042**
Actinomyces israelii		100.0	0.43	SAL 92			
63557100	**Very good identification**						**63557100**
Actinomyces israelii		100.0	0.57	SAL 92			
63557102	**Very good identification**						**63557102**
					GRA	CAT	
Actinomyces israelii		97.26	0.60	SAL 92	99	1	
Capnocytophaga ochracea		1.44	0.13		1	90	
Bifidobacterium adolescentis		1.30	0.22		99	1	
63557110	**Good identification**						**63557110**
Actinomyces israelii		100.0	0.43	SAL 92			
63557112	**Acceptable identification**						**63557112**
Actinomyces israelii		90.16	0.46	SAL 92			
Bifidobacterium adolescentis		9.84	0.23	MLZ 13			
63557120	**Low discrimination**						**63557120**
					GRA	CAT	
Actinomyces israelii		74.18	0.37	SAL 92	99	1	
Capnocytophaga ochracea		25.82	0.13	NAG 6	1	90	
63557140	**Good identification**						**63557140**
Actinomyces israelii		100.0	0.50	SAL 92			
63557142	**Good identification**						**63557142**
Actinomyces israelii		100.0	0.53	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63557150	**Good identification**						**63557150**
Actinomyces israelii		100.0	0.36	SAL 92			
63557152	**Good identification**						**63557152**
Actinomyces israelii		96.65	0.39	SAL 92			
Bifidobacterium adolescentis		3.35	0.08	MLZ 13			
63557300	**Good identification**						**63557300**
Actinomyces israelii		100.0	0.38	SAL 92			
63557302	**Good identification**						**63557302**
Actinomyces israelii		100.0	0.41	SAL 92			
63557400	**Good identification**						**63557400**
Actinomyces israelii		100.0	0.45	SAL 92			
63557402	**Good identification**						**63557402**
Actinomyces israelii		100.0	0.48	SAL 92			
63557440	**Good identification**						**63557440**
Actinomyces israelii		100.0	0.38	SAL 92			
63557442	**Good identification**						**63557442**
Actinomyces israelii		100.0	0.41	SAL 92			
63557500	**Good identification**						**63557500**
Actinomyces israelii		100.0	0.56	SAL 92			
63557502	**Low discrimination**						**63557502**
					GRA	CAT	
Actinomyces israelii		89.90	0.59	SAL 92	99	1	
Bifidobacterium adolescentis		9.81	0.36		99	1	
Capnocytophaga ochracea		0.29	0.00		1	90	
63557510	**Good identification**						**63557510**
Actinomyces israelii		100.0	0.41	SAL 92			
63557512	**Low discrimination**						**63557512**
Actinomyces israelii		52.83	0.45	SAL 92			
Bifidobacterium adolescentis		47.17	0.37	MLZ 13			
63557520	**Acceptable identification**						**63557520**
					GRA	CAT	
Actinomyces israelii		93.02	0.35	SAL 92	99	1	
Capnocytophaga ochracea		6.99	0.00	NAG 6	1	90	
63557522	**Low discrimination**						**63557522**
					GRA	CAT	
Actinomyces israelii		68.94	0.38	SAL 92	99	1	
Capnocytophaga ochracea		31.06	0.16	NAG 6	1	90	
63557540	**Good identification**						**63557540**
Actinomyces israelii		100.0	0.49	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63557542	**Good identification**					**63557542**
Actinomyces israelii		96.65	0.52	SAL 92		
Bifidobacterium adolescentis		3.35	0.20			
63557552	**Low discrimination**					**63557552**
Actinomyces israelii		77.90	0.37	SAL 92		
Bifidobacterium adolescentis		22.10	0.22	MLZ 13		
63557700	**Good identification**					**63557700**
Actinomyces israelii		100.0	0.36	SAL 92		
63557702	**Good identification**					**63557702**
Actinomyces israelii		100.0	0.39	SAL 92		
63563100	**Low discrimination**					**63563100**
Actinomyces naeslundii		83.89	0.36	ESL 12		
Actinomyces israelii		16.11	0.15	MNS 88		
63563101	**Good identification**					**63563101**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63563102	**Low discrimination**					**63563102**
Actinomyces naeslundii		77.64	0.36	ESL 12		
Actinomyces israelii		22.37	0.18	MNS 88		
63563103	**Good identification**					**63563103**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63563120	**Good identification**					**63563120**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63563121	**Good identification**					**63563121**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63563122	**Good identification**					**63563122**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63563123	**Good identification**					**63563123**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63567102	**Good identification**					**63567102**
Actinomyces israelii		100.0	0.36	SAL 92		
63571000	**Low discrimination**					**63571000**
					GRA	
Actinomyces naeslundii		74.07	0.45	ESL 12	99	
Actinomyces israelii		18.07	0.26	MNS 88	99	
Fusobacterium mortiferum		7.86	0.17		1	
63571001	**Good identification**					**63571001**
Actinomyces naeslundii		100.0	0.48	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63571002	**Low discrimination**						**63571002**
Actinomyces naeslundii		67.38	0.45	ESL 12			
Actinomyces israelii		24.66	0.29	MNS 88			
Actinomyces viscosus		7.97	0.05				
63571003	**Good identification**						**63571003**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63571020	**Good identification**						**63571020**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63571021	**Good identification**						**63571021**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63571022	**Low discrimination**						**63571022**
					GRA		
Actinomyces naeslundii		87.98	0.45	ESL 12	99		
Capnocytophaga ochracea		12.02	0.05	MNS 88	1		
63571023	**Good identification**						**63571023**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63571041	**Good identification**						**63571041**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63571043	**Good identification**						**63571043**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63571061	**Good identification**						**63571061**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63571063	**Good identification**						**63571063**
Actinomyces naeslundii		100.0	0.35	ESL 12			
63571100	**Low discrimination**						**63571100**
					GRA		
Actinomyces naeslundii		67.18	0.53	ESL 12	99		
Actinomyces israelii		23.03	0.36	MNS 88	99		
Fusobacterium mortiferum		5.85	0.23		1		
63571101	**Good identification**						**63571101**
Actinomyces naeslundii		100.0	0.56	ESL 12			
63571102	**Low discrimination**						**63571102**
					GRA	CAT	
Actinomyces naeslundii		47.26	0.53	ESL 12	99	25	
Capnocytophaga ochracea		24.95	0.23	MNS 88	1	90	
Actinomyces israelii		24.30	0.39		99	1	
63571103	**Good identification**						**63571103**
Actinomyces naeslundii		98.39	0.56	ESL 12			
Actinomyces viscosus		1.62	0.01	MNS 88			
63571120	**Low discrimination**						**63571120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces naeslundii		60.83	0.53	ESL 12	99	25	
Capnocytophaga ochracea		32.11	0.23	MNS 88	1	90	
Clostridium perfringens		4.29	0.11		99	1	
63571121	**Good identification**						**63571121**
Actinomyces naeslundii		100.0	0.56	ESL 12			
63571122	**Low discrimination**						**63571122**
					GRA	CAT	
Capnocytophaga ochracea		80.78	0.39	MNS 90	1	90	
Actinomyces naeslundii		17.00	0.53	TRE 1	99	25	
Prevotella oralis		1.10	0.07		1	1	
63571123	**Good identification**						**63571123**
					GRA		
Actinomyces naeslundii		96.90	0.56	ESL 12	99		
Capnocytophaga ochracea		3.10	0.05	MNS 88	1		
63571140	**Low discrimination**						**63571140**
Actinomyces naeslundii		56.96	0.40	ESL 12			
Actinomyces israelii		43.04	0.29	MNS 88			
63571141	**Good identification**						**63571141**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63571143	**Good identification**						**63571143**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63571160	**Low discrimination**						**63571160**
					GRA		
Actinomyces naeslundii		75.05	0.40	ESL 12	99		
Capnocytophaga ochracea		24.95	0.07	MNS 88	1		
63571161	**Good identification**						**63571161**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63571163	**Good identification**						**63571163**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63571502	**Low discrimination**						**63571502**
					GRA	CAT	
Actinomyces israelii		68.89	0.37	SAL 92	99	1	
Bifidobacterium adolescentis		15.86	0.20	XYL 92	99	1	
Capnocytophaga ochracea		15.25	0.10		1	90	
63572101	**Good identification**						**63572101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63572103	**Good identification**						**63572103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63572121	**Good identification**						**63572121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63572123	**Good identification**						**63572123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ESL 12		
63573000	**Low discrimination**					**63573000**
Actinomyces naeslundii		58.41	0.50	ESL 12		
Actinomyces israelii		41.59	0.38	MNS 88		
63573001	**Good identification**					**63573001**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63573002	**Low discrimination**					**63573002**
Actinomyces israelii		51.65	0.42	SAL 92		
Actinomyces naeslundii		48.35	0.50	XYL 92		
63573003	**Good identification**					**63573003**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63573020	**Good identification**					**63573020**
Actinomyces naeslundii		95.65	0.50	ESL 12		
Actinomyces israelii		4.35	0.18	MNS 88		
63573021	**Good identification**					**63573021**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63573022	**Acceptable identification**					**63573022**
Actinomyces naeslundii		93.62	0.50	ESL 12		
Actinomyces israelii		6.38	0.21	MNS 88		
63573023	**Good identification**					**63573023**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63573041	**Good identification**					**63573041**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63573043	**Good identification**					**63573043**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63573060	**Good identification**					**63573060**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63573061	**Good identification**					**63573061**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63573062	**Low discrimination**					**63573062**
Actinomyces naeslundii		86.94	0.37	ESL 12		
Actinomyces israelii		13.06	0.14	MNS 88		
63573063	**Good identification**					**63573063**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63573100	**Low discrimination**					**63573100**
Actinomyces israelii		48.99	0.49	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		48.95	0.58	XYL 92			
Propionibacterium propionicum		2.06	0.24				
63573101	**Very good identification**						**63573101**
Actinomyces naeslundii		99.33	0.61	ESL 12			
Actinomyces israelii		0.67	0.15	MNS 88			
63573102	**Low discrimination**						**63573102**
Actinomyces israelii		58.54	0.52	SAL 92			
Actinomyces naeslundii		39.00	0.58	XYL 92			
Propionibacterium propionicum		2.46	0.27				
63573103	**Very good identification**						**63573103**
Actinomyces naeslundii		99.00	0.61	ESL 12			
Actinomyces israelii		1.00	0.18	MNS 88			
63573111	**Good identification**						**63573111**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63573112	**Low discrimination**						**63573112**
Actinomyces israelii		81.00	0.38	SAL 92			
Actinomyces naeslundii		19.00	0.36	XYL 92			
63573113	**Good identification**						**63573113**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63573120	**Good identification**						**63573120**
Actinomyces naeslundii		94.00	0.58	ESL 12			
Actinomyces israelii		6.01	0.28	MNS 88			
63573121	**Very good identification**						**63573121**
Actinomyces naeslundii		100.0	0.61	ESL 12			
63573122	**Low discrimination**						**63573122**
					GRA	CAT	
Actinomyces naeslundii		89.24	0.58	ESL 12	99	25	
Actinomyces israelii		8.55	0.31	MNS 88	99	1	
Capnocytophaga ochracea		2.21	0.05		1	90	
63573123	**Very good identification**						**63573123**
Actinomyces naeslundii		100.0	0.61	ESL 12			
63573130	**Low discrimination**						**63573130**
Actinomyces naeslundii		84.65	0.36	ESL 12			
Actinomyces israelii		15.35	0.14	MNS 88			
63573131	**Good identification**						**63573131**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63573132	**Low discrimination**						**63573132**
Actinomyces naeslundii		78.61	0.36	ESL 12			
Actinomyces israelii		21.39	0.17	MNS 88			
63573133	**Good identification**						**63573133**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	ESL 12		
63573140	**Low discrimination**					**63573140**
Actinomyces israelii		64.05	0.42	SAL 92		
Actinomyces naeslundii		29.04	0.45	XYL 92		
Propionibacterium propionicum		6.92	0.24			
63573141	**Good identification**					**63573141**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63573142	**Low discrimination**					**63573142**
Actinomyces israelii		70.91	0.45	SAL 92		
Actinomyces naeslundii		21.43	0.45	XYL 92		
Propionibacterium propionicum		7.66	0.27			
63573143	**Good identification**					**63573143**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63573160	**Low discrimination**					**63573160**
Actinomyces naeslundii		87.66	0.45	ESL 12		
Actinomyces israelii		12.34	0.21	MNS 88		
63573161	**Good identification**					**63573161**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63573162	**Low discrimination**					**63573162**
Actinomyces naeslundii		82.57	0.45	ESL 12		
Actinomyces israelii		17.44	0.24	MNS 88		
63573163	**Good identification**					**63573163**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63573400	**Good identification**					**63573400**
Actinomyces israelii		100.0	0.37	SAL 92		
63573402	**Good identification**					**63573402**
Actinomyces israelii		100.0	0.40	SAL 92		
63573500	**Good identification**					**63573500**
Actinomyces israelii		98.78	0.47	SAL 92		
Actinomyces naeslundii		1.22	0.24	XYL 92		
63573502	**Good identification**					**63573502**
Actinomyces israelii		96.02	0.50	SAL 92		
Bifidobacterium adolescentis		3.19	0.19	XYL 92		
Actinomyces naeslundii		0.79	0.24			
63573512	**Low discrimination**					**63573512**
Actinomyces israelii		78.62	0.36	SAL 92		
Bifidobacterium adolescentis		21.38	0.20	XYL 92		
63573540	**Good identification**					**63573540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	SAL 92			
63573542	**Good identification**						**63573542**
Actinomyces israelii		100.0	0.43	SAL 92			
63575000	**Good identification**						**63575000**
Actinomyces israelii		100.0	0.44	SAL 92			
63575002	**Good identification**						**63575002**
Actinomyces israelii		100.0	0.47	SAL 92			
63575040	**Good identification**						**63575040**
Actinomyces israelii		100.0	0.37	SAL 92			
63575042	**Good identification**						**63575042**
Actinomyces israelii		100.0	0.40	SAL 92			
63575100	**Good identification**						**63575100**
Actinomyces israelii		100.0	0.54	SAL 92			
63575102	**Very good identification**						**63575102**
					GRA	CAT	
Actinomyces israelii		96.28	0.57	SAL 92	99	1	
Capnocytophaga ochracea		2.15	0.13		1	90	
Bifidobacterium adolescentis		1.58	0.20		99	1	
63575110	**Good identification**						**63575110**
Actinomyces israelii		100.0	0.40	SAL 92			
63575112	**Low discrimination**						**63575112**
Actinomyces israelii		88.19	0.43	SAL 92			
Bifidobacterium adolescentis		11.81	0.22	MLZ 13			
63575140	**Good identification**						**63575140**
Actinomyces israelii		100.0	0.47	SAL 92			
63575142	**Good identification**						**63575142**
Actinomyces israelii		100.0	0.50	SAL 92			
63575152	**Good identification**						**63575152**
Actinomyces israelii		100.0	0.36	SAL 92			
63575302	**Good identification**						**63575302**
Actinomyces israelii		100.0	0.38	SAL 92			
63575400	**Good identification**						**63575400**
Actinomyces israelii		100.0	0.42	SAL 92			
63575402	**Good identification**						**63575402**
Actinomyces israelii		100.0	0.45	SAL 92			
63575440	**Good identification**						**63575440**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.35	SAL 92			
63575442	**Good identification**						**63575442**
Actinomyces israelii		100.0	0.38	SAL 92			
63575500	**Acceptable identification**						**63575500**
					GRA	SPO	
Actinomyces israelii		92.01	0.53	SAL 92	99	1	
Mitsuokella multacida		7.03	0.21		1	1	
Clostridium butyricum		0.96	-.03		99	99	
63575502	**Low discrimination**						**63575502**
					GRA		
Actinomyces israelii		87.38	0.56	SAL 92	99		
Bifidobacterium adolescentis		11.71	0.34		99		
Mitsuokella multacida		0.50	0.04		1		
63575510	**Low discrimination**						**63575510**
					GRA		
Actinomyces israelii		85.43	0.38	SAL 92	99		
Mitsuokella multacida		14.57	0.13	MLZ 13	1		
63575512	**Low discrimination**						**63575512**
Bifidobacterium adolescentis		52.29	0.36	NIT 1			
Actinomyces israelii		47.72	0.41	MNS 87			
63575522	**Low discrimination**						**63575522**
					GRA	CAT	
Actinomyces israelii		52.79	0.35	SAL 92	99	1	
Capnocytophaga ochracea		35.82	0.16	NAG 6	1	90	
Parabacteroides distasonis		11.39	0.20		1	76	
63575540	**Low discrimination**						**63575540**
					GRA		
Actinomyces israelii		83.57	0.46	SAL 92	99		
Mitsuokella multacida		16.43	0.21		1		
63575542	**Acceptable identification**						**63575542**
					GRA		
Actinomyces israelii		94.60	0.49	SAL 92	99		
Bifidobacterium adolescentis		4.03	0.19		99		
Mitsuokella multacida		1.38	0.04		1		
63575702	**Low discrimination**						**63575702**
					GRA		
Actinomyces israelii		83.56	0.36	SAL 92	99		
Mitsuokella multacida		11.67	0.09	RHA 7	1		
Clostridium clostridioforme		4.77	0.06		50		
63576102	**Good identification**						**63576102**
Actinomyces israelii		100.0	0.36	SAL 92			
63577000	**Very good identification**						**63577000**
Actinomyces israelii		100.0	0.57	SAL 92			
63577002	**Very good identification**						**63577002**
Actinomyces israelii		100.0	0.60	SAL 92			
63577010	**Good identification**						**63577010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.43	SAL 92		
63577012	**Good identification**					**63577012**
Actinomyces israelii		100.0	0.46	SAL 92		
63577020	**Good identification**					**63577020**
Actinomyces israelii		100.0	0.36	SAL 92		
63577022	**Good identification**					**63577022**
Actinomyces israelii		100.0	0.39	SAL 92		
63577040	**Good identification**					**63577040**
Actinomyces israelii		100.0	0.50	SAL 92		
63577042	**Good identification**					**63577042**
Actinomyces israelii		100.0	0.53	SAL 92		
63577050	**Good identification**					**63577050**
Actinomyces israelii		100.0	0.36	SAL 92		
63577052	**Good identification**					**63577052**
Actinomyces israelii		100.0	0.39	SAL 92		
63577100	**Very good identification**					**63577100**
Actinomyces israelii		99.91	0.67	SAL 92		
Actinomyces naeslundii		0.09	0.24			
63577102	**Very good identification**					**63577102**
Actinomyces israelii		99.71	0.70	SAL 92		
Bifidobacterium adolescentis		0.24	0.19			
Actinomyces naeslundii		0.06	0.24			
63577103	**Acceptable identification**					**63577103**
Actinomyces israelii		92.01	0.36	SAL 92		
Actinomyces naeslundii		7.99	0.27	URE 1		
63577110	**Good identification**					**63577110**
Actinomyces israelii		100.0	0.53	SAL 92		
63577112	**Good identification**					**63577112**
Actinomyces israelii		98.10	0.56	SAL 92		
Bifidobacterium adolescentis		1.90	0.20	MLZ 13		
63577120	**Good identification**					**63577120**
Actinomyces israelii		98.64	0.47	SAL 92		
Actinomyces naeslundii		1.36	0.24	NAG 6		
63577122	**Good identification**					**63577122**
Actinomyces israelii		99.09	0.50	SAL 92		
Actinomyces naeslundii		0.91	0.24	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63577132	**Good identification**					**63577132**
Actinomyces israelii		100.0	0.35	SAL 92		
63577140	**Very good identification**					**63577140**
Actinomyces israelii		100.0	0.60	SAL 92		
63577142	**Very good identification**					**63577142**
Actinomyces israelii		100.0	0.63	SAL 92		
63577150	**Good identification**					**63577150**
Actinomyces israelii		100.0	0.46	SAL 92		
63577152	**Good identification**					**63577152**
Actinomyces israelii		100.0	0.49	SAL 92		
63577160	**Good identification**					**63577160**
Actinomyces israelii		100.0	0.39	SAL 92		
63577162	**Good identification**					**63577162**
Actinomyces israelii		100.0	0.42	SAL 92		
63577200	**Good identification**					**63577200**
Actinomyces israelii		100.0	0.37	SAL 92		
63577202	**Good identification**					**63577202**
Actinomyces israelii		100.0	0.40	SAL 92		
63577300	**Good identification**					**63577300**
Actinomyces israelii		100.0	0.48	SAL 92		
63577302	**Good identification**					**63577302**
Actinomyces israelii		100.0	0.51	SAL 92		
63577312	**Good identification**					**63577312**
Actinomyces israelii		100.0	0.37	SAL 92		
63577340	**Good identification**					**63577340**
Actinomyces israelii		100.0	0.41	SAL 92		
63577342	**Good identification**					**63577342**
Actinomyces israelii		100.0	0.44	SAL 92		
63577400	**Good identification**					**63577400**
Actinomyces israelii		100.0	0.55	SAL 92		
63577402	**Very good identification**					**63577402**
Actinomyces israelii		100.0	0.58	SAL 92		
63577410	**Good identification**					**63577410**
Actinomyces israelii		100.0	0.41	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63577412	**Good identification**						**63577412**
Actinomyces israelii		100.0	0.44	SAL 92			
63577422	**Good identification**						**63577422**
Actinomyces israelii		100.0	0.38	SAL 92			
63577440	**Good identification**						**63577440**
Actinomyces israelii		100.0	0.48	SAL 92			
63577442	**Good identification**						**63577442**
Actinomyces israelii		100.0	0.51	SAL 92			
63577452	**Good identification**						**63577452**
Actinomyces israelii		100.0	0.37	SAL 92			
63577500	**Very good identification**						**63577500**
					GRA		
Actinomyces israelii		96.95	0.65	SAL 92	99		
Mitsuokella multacida		3.05	0.27		1		
63577502	**Very good identification**						**63577502**
					GRA		
Actinomyces israelii		97.88	0.68	SAL 92	99		
Bifidobacterium adolescentis		1.89	0.33		99		
Mitsuokella multacida		0.23	0.11		1		
63577510	**Acceptable identification**						**63577510**
					GRA		
Actinomyces israelii		93.44	0.51	SAL 92	99		
Mitsuokella multacida		6.56	0.19	MLZ 13	1		
63577512	**Low discrimination**						**63577512**
Actinomyces israelii		85.43	0.54	SAL 92			
Bifidobacterium adolescentis		13.52	0.34	MLZ 13			
Bifidobacterium dentium		0.61	-.07				
63577520	**Low discrimination**						**63577520**
					GRA		
Actinomyces israelii		89.03	0.45	SAL 92	99		
Mitsuokella multacida		10.97	0.17	NAG 6	1		
63577522	**Good identification**						**63577522**
Actinomyces israelii		100.0	0.48	SAL 92			
63577540	**Good identification**						**63577540**
					GRA		
Actinomyces israelii		92.51	0.58	SAL 92	99		
Mitsuokella multacida		7.49	0.27		1		
63577542	**Very good identification**						**63577542**
					GRA		
Actinomyces israelii		98.80	0.61	SAL 92	99		
Bifidobacterium adolescentis		0.61	0.17		99		
Mitsuokella multacida		0.59	0.11		1		
63577550	**Low discrimination**						**63577550**
					COC	GRA	
Actinomyces israelii		82.81	0.44	SAL 92	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		14.95	0.19	MLZ 13	1	1	
Blautia producta		2.24	-.03		99	99	
63577552	**Acceptable identification**						**63577552**
					GRA		
Actinomyces israelii		94.02	0.47	SAL 92	99		
Bifidobacterium adolescentis		4.73	0.19	MLZ 13	99		
Mitsuokella multacida		1.26	0.03		1		
63577560	**Low discrimination**						**63577560**
					GRA		
Actinomyces israelii		75.93	0.38	SAL 92	99		
Mitsuokella multacida		24.07	0.17	NAG 6	1		
63577562	**Good identification**						**63577562**
Actinomyces israelii		100.0	0.41	SAL 92			
63577600	**Good identification**						**63577600**
Actinomyces israelii		100.0	0.36	SAL 92			
63577602	**Good identification**						**63577602**
Actinomyces israelii		100.0	0.39	SAL 92			
63577700	**Low discrimination**						**63577700**
					GRA		
Actinomyces israelii		56.29	0.46	SAL 92	99		
Mitsuokella multacida		43.71	0.32	RHA 7	1		
63577702	**Acceptable identification**						**63577702**
					GRA		
Actinomyces israelii		94.56	0.49	SAL 92	99		
Mitsuokella multacida		5.44	0.15	RHA 7	1		
63577712	**Low discrimination**						**63577712**
					GRA		
Actinomyces israelii		88.63	0.35	SAL 92	99		
Mitsuokella multacida		11.37	0.07	RHA 7	1		
63577742	**Low discrimination**						**63577742**
					GRA		
Actinomyces israelii		87.11	0.42	SAL 92	99		
Mitsuokella multacida		12.89	0.15	RHA 7	1		
63600000	**Very good identification**						**63600000**
					GRA		
Eubacterium tortuosum		99.94	0.56	GAL 99	99		
Fusobacterium mortiferum		0.06	0.17		1		
63600002	**Good identification**						**63600002**
Eubacterium tortuosum		100.0	0.43	GAL 99			
63600020	**Good identification**						**63600020**
Eubacterium tortuosum		100.0	0.43	GAL 99			
63601000	**Very good identification**						**63601000**
					GRA		
Eubacterium tortuosum		100.0	0.90		99		
Fusobacterium mortiferum		0.00	0.20		1		
63601001	**Very good identification**						**63601001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.56	URE 1		
63601002	**Very good identification**					**63601002**
Eubacterium tortuosum		100.0	0.77			
63601003	**Good identification**					**63601003**
Eubacterium tortuosum		100.0	0.43	URE 1		
63601010	**Very good identification**					**63601010**
Eubacterium tortuosum		100.0	0.56	MLZ 1		
63601012	**Good identification**					**63601012**
Eubacterium tortuosum		100.0	0.43	MLZ 1		
63601020	**Very good identification**					**63601020**
Eubacterium tortuosum		100.0	0.77			
63601021	**Good identification**					**63601021**
Eubacterium tortuosum		100.0	0.43	URE 1		
63601022	**Very good identification**					**63601022**
Eubacterium tortuosum		100.0	0.65			
63601030	**Good identification**					**63601030**
Eubacterium tortuosum		100.0	0.43	MLZ 1		
63601040	**Very good identification**					**63601040**
Eubacterium tortuosum		100.0	0.56	SOR 1		
63601042	**Good identification**					**63601042**
Eubacterium tortuosum		100.0	0.43	SOR 1		
63601060	**Good identification**					**63601060**
Eubacterium tortuosum		100.0	0.43	SOR 1		
63601100	**Very good identification**					**63601100**
					GRA	
Eubacterium tortuosum		99.80	0.56	LAC 1	99	
Fusobacterium mortiferum		0.20	0.27		1	
63601102	**Good identification**					**63601102**
Eubacterium tortuosum		100.0	0.43	LAC 1		
63601120	**Good identification**					**63601120**
Eubacterium tortuosum		100.0	0.43	LAC 1		
63601200	**Very good identification**					**63601200**
Eubacterium tortuosum		100.0	0.56	RHA 1		
63601202	**Good identification**					**63601202**
Eubacterium tortuosum		100.0	0.43	RHA 1		
63601220	**Good identification**					**63601220**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.43	RHA 1		
63601400	**Very good identification**					**63601400**
Eubacterium tortuosum		100.0	0.56	ARA 1		
63601402	**Good identification**					**63601402**
Eubacterium tortuosum		100.0	0.43	ARA 1		
63601420	**Good identification**					**63601420**
Eubacterium tortuosum		100.0	0.43	ARA 1		
63603000	**Very good identification**					**63603000**
Eubacterium tortuosum		100.0	0.56	MAN 1		
63603002	**Good identification**					**63603002**
Eubacterium tortuosum		100.0	0.43	MAN 1		
63603020	**Good identification**					**63603020**
Eubacterium tortuosum		100.0	0.43	MAN 1		
63605000	**Very good identification**					**63605000**
Eubacterium tortuosum		100.0	0.56	XYL 1		
63605002	**Good identification**					**63605002**
Eubacterium tortuosum		100.0	0.43	XYL 1		
63605020	**Good identification**					**63605020**
Eubacterium tortuosum		100.0	0.43	XYL 1		
63610000	**Low discrimination**					**63610000**
					GRA	
Fusobacterium mortiferum		84.99	0.52	NIT 1	1	
Eubacterium tortuosum		15.01	0.22		99	
63610002	**Low discrimination**					**63610002**
					GRA	
Fusobacterium mortiferum		78.09	0.35	NIT 1	1	
Eubacterium tortuosum		21.91	0.09	BGL 10	99	
63610100	**Very good identification**					**63610100**
Fusobacterium mortiferum		100.0	0.58	NIT 1		
63610102	**Good identification**					**63610102**
					GRA	
Fusobacterium mortiferum		98.44	0.42	NIT 1	1	
Actinomyces viscosus		1.57	-.02	BGL 10	99	
63611000	**Good identification**					**63611000**
					GRA	
Eubacterium tortuosum		92.06	0.56	FRU 1	99	
Fusobacterium mortiferum		7.90	0.55		1	
Actinomyces naeslundii		0.04	0.27		99	
63611002	**Acceptable identification**					**63611002**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		94.65	0.43	FRU 1	99	
Fusobacterium mortiferum		5.11	0.38		1	
Actinomyces naeslundii		0.24	0.27		99	
63611020	**Good identification**					**63611020**
					GRA	
Eubacterium tortuosum		99.26	0.43	FRU 1	99	
Fusobacterium mortiferum		0.49	0.20		1	
Actinomyces naeslundii		0.25	0.27		99	
63611100	**Good identification**					**63611100**
					GRA	
Fusobacterium mortiferum		94.62	0.61	NIT 1	1	
Eubacterium tortuosum		4.78	0.22		99	
Actinomyces naeslundii		0.61	0.35		99	
63611102	**Low discrimination**					**63611102**
					GRA	
Fusobacterium mortiferum		86.03	0.45	NIT 1	1	
Eubacterium tortuosum		6.90	0.09	BGL 10	99	
Actinomyces naeslundii		4.96	0.35		99	
63611103	**Good identification**					**63611103**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63611121	**Good identification**					**63611121**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63611123	**Good identification**					**63611123**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63613001	**Good identification**					**63613001**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63613003	**Good identification**					**63613003**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63613021	**Good identification**					**63613021**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63613023	**Good identification**					**63613023**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63613100	**Low discrimination**					**63613100**
					GRA	
Actinomyces naeslundii		55.18	0.40	ESL 12	99	
Fusobacterium mortiferum		44.82	0.27	MNS 88	1	
63613101	**Good identification**					**63613101**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63613102	**Good identification**					**63613102**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63613103	**Good identification**					**63613103**
Actinomyces naeslundii		100.0	0.43	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63613120	**Good identification**					**63613120**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63613121	**Good identification**					**63613121**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63613122	**Good identification**					**63613122**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63613123	**Good identification**					**63613123**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63614100	**Good identification**					**63614100**
Fusobacterium mortiferum		100.0	0.39	NIT 1		
63615100	**Good identification**					**63615100**
Fusobacterium mortiferum		100.0	0.42	NIT 1		
63621000	**Very good identification**					**63621000**
					GRA	
Eubacterium tortuosum		99.97	0.56	TRE 1	99	
Fusobacterium mortiferum		0.03	0.12		1	
63621002	**Good identification**					**63621002**
Eubacterium tortuosum		100.0	0.43	TRE 1		
63621020	**Good identification**					**63621020**
Eubacterium tortuosum		100.0	0.43	TRE 1		
63630000	**Good identification**					**63630000**
Fusobacterium mortiferum		100.0	0.43	NIT 1		
63630100	**Good identification**					**63630100**
Fusobacterium mortiferum		100.0	0.49	NIT 1		
63631000	**Low discrimination**					**63631000**
					GRA	
Fusobacterium mortiferum		70.01	0.46	NIT 1	1	
Eubacterium tortuosum		27.60	0.22		99	
Actinomyces naeslundii		2.39	0.32		99	
63631001	**Low discrimination**					**63631001**
					GRA	
Actinomyces naeslundii		83.52	0.35	ESL 12	99	
Fusobacterium mortiferum		16.48	0.12	MNS 88	1	
63631003	**Good identification**					**63631003**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63631021	**Good identification**					**63631021**
Actinomyces naeslundii		100.0	0.35	ESL 12		
63631023	**Good identification**					**63631023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	ESL 12		
63631100	**Good identification**					**63631100**
					GRA	
Fusobacterium mortiferum		96.00	0.52	NIT 1	1	
Actinomyces naeslundii		4.00	0.40		99	
63631101	**Low discrimination**					**63631101**
					GRA	
Actinomyces naeslundii		86.08	0.43	ESL 12	99	
Fusobacterium mortiferum		13.92	0.18	MNS 88	1	
63631102	**Low discrimination**					**63631102**
					GRA	
Fusobacterium mortiferum		72.75	0.36	NIT 1	1	
Actinomyces naeslundii		27.26	0.40	BGL 10	99	
63631103	**Good identification**					**63631103**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63631120	**Low discrimination**					**63631120**
					GRA	
Actinomyces naeslundii		80.47	0.40	ESL 12	99	
Fusobacterium mortiferum		19.53	0.18	MNS 88	1	
63631121	**Good identification**					**63631121**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63631122	**Good identification**					**63631122**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63631123	**Good identification**					**63631123**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63633000	**Low discrimination**					**63633000**
					GRA	
Actinomyces naeslundii		86.77	0.37	ESL 12	99	
Fusobacterium mortiferum		13.23	0.12	MNS 88	1	
63633001	**Good identification**					**63633001**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63633002	**Good identification**					**63633002**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63633003	**Good identification**					**63633003**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63633020	**Good identification**					**63633020**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63633021	**Good identification**					**63633021**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63633022	**Good identification**					**63633022**
Actinomyces naeslundii		100.0	0.37	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63633023	**Good identification**					**63633023**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63633100	**Low discrimination**					**63633100**
					GRA	
Actinomyces naeslundii		85.02	0.45	ESL 12	99	
Fusobacterium mortiferum		10.63	0.18	MNS 88	1	
Actinomyces israelii		4.36	0.14	MLT 94	99	
63633101	**Good identification**					**63633101**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63633102	**Acceptable identification**					**63633102**
Actinomyces naeslundii		92.86	0.45	ESL 12		
Actinomyces israelii		7.14	0.17	MNS 88		
63633103	**Good identification**					**63633103**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63633120	**Good identification**					**63633120**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63633121	**Good identification**					**63633121**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63633122	**Good identification**					**63633122**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63633123	**Good identification**					**63633123**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63640000	**Good identification**					**63640000**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63641000	**Very good identification**					**63641000**
					GRA	
Eubacterium tortuosum		100.0	0.74	CEL 10	99	
Fusobacterium mortiferum		0.00	0.12		1	
63641001	**Good identification**					**63641001**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63641002	**Very good identification**					**63641002**
Eubacterium tortuosum		100.0	0.61	CEL 10		
63641010	**Good identification**					**63641010**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63641020	**Very good identification**					**63641020**
Eubacterium tortuosum		100.0	0.61	CEL 10		
63641022	**Good identification**					**63641022**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.48	CEL 10		
63641040	**Good identification**					**63641040**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63641100	**Good identification**					**63641100**
					GRA	
Eubacterium tortuosum		99.40	0.40	CEL 10	99	
Fusobacterium mortiferum		0.60	0.19	LAC 1	1	
63641200	**Good identification**					**63641200**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63641400	**Good identification**					**63641400**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63643000	**Good identification**					**63643000**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63645000	**Good identification**					**63645000**
Eubacterium tortuosum		100.0	0.40	CEL 10		
63650000	**Acceptable identification**					**63650000**
					GRA	
Fusobacterium mortiferum		94.44	0.43	NIT 1	1	
Eubacterium tortuosum		5.56	0.06		99	
63650100	**Good identification**					**63650100**
Fusobacterium mortiferum		100.0	0.50	NIT 1		
63651000	**Low discrimination**					**63651000**
					GRA	
Eubacterium tortuosum		79.19	0.40	FRU 1	99	
Fusobacterium mortiferum		20.38	0.46	CEL 10	1	
Actinomyces naeslundii		0.43	0.29		99	
63651100	**Good identification**					**63651100**
					GRA	
Fusobacterium mortiferum		95.94	0.53	NIT 1	1	
Actinomyces naeslundii		2.44	0.37		99	
Eubacterium tortuosum		1.61	0.06		99	
63651101	**Low discrimination**					**63651101**
					GRA	
Actinomyces naeslundii		79.09	0.40	ESL 12	99	
Fusobacterium mortiferum		20.91	0.19	MNS 88	1	
63651102	**Low discrimination**					**63651102**
					GRA	
Fusobacterium mortiferum		81.35	0.36	NIT 1	1	
Actinomyces naeslundii		18.65	0.37	BGL 10	99	
63651103	**Good identification**					**63651103**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63651120	**Low discrimination**					**63651120**
					GRA	
Actinomyces naeslundii		71.60	0.37	ESL 12	99	
Fusobacterium mortiferum		28.40	0.19	MNS 88	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63651121	**Good identification**						**63651121**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63651122	**Low discrimination**						**63651122**
					GRA	CAT	
Actinomyces naeslundii		63.81	0.37	ESL 12	99	25	
Capnocytophaga ochracea		26.27	0.05	MNS 88	1	90	
Prevotella oralis		9.93	-.03	MLT 94	1	1	
63651123	**Good identification**						**63651123**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63653001	**Good identification**						**63653001**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63653003	**Good identification**						**63653003**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63653021	**Good identification**						**63653021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63653023	**Good identification**						**63653023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63653100	**Low discrimination**						**63653100**
					GRA		
Actinomyces naeslundii		68.06	0.42	ESL 12	99		
Actinomyces israelii		18.04	0.23	MNS 88	99		
Fusobacterium mortiferum		13.90	0.19	MLT 94	1		
63653101	**Good identification**						**63653101**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63653102	**Low discrimination**						**63653102**
Actinomyces naeslundii		71.55	0.42	ESL 12			
Actinomyces israelii		28.45	0.26	MNS 88			
63653103	**Good identification**						**63653103**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63653120	**Good identification**						**63653120**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63653121	**Good identification**						**63653121**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63653122	**Good identification**						**63653122**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63653123	**Good identification**						**63653123**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63657100	**Good identification**						**63657100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.41	MLT 99		
63657102	**Good identification**					**63657102**
Actinomyces israelii		100.0	0.44	MLT 99		
63657142	**Good identification**					**63657142**
Actinomyces israelii		100.0	0.37	MLT 99		
63657500	**Good identification**					**63657500**
Actinomyces israelii		100.0	0.40	MLT 99		
63657502	**Acceptable identification**					**63657502**
Actinomyces israelii		94.90	0.43	MLT 99		
Bifidobacterium adolescentis		5.10	0.15			
63657542	**Good identification**					**63657542**
Actinomyces israelii		100.0	0.36	MLT 99		
63661000	**Good identification**					**63661000**
Eubacterium tortuosum		100.0	0.40	TRE 1		
63670100	**Good identification**					**63670100**
Fusobacterium mortiferum		100.0	0.41	NIT 1		
63671000	**Low discrimination**					**63671000**
					GRA	
Fusobacterium mortiferum		78.92	0.38	NIT 1	1	
Actinomyces naeslundii		10.71	0.34		99	
Eubacterium tortuosum		10.37	0.06		99	
63671001	**Good identification**					**63671001**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63671003	**Good identification**					**63671003**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63671021	**Good identification**					**63671021**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63671023	**Good identification**					**63671023**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63671100	**Low discrimination**					**63671100**
					GRA	
Fusobacterium mortiferum		83.71	0.44	NIT 1	1	
Actinomyces naeslundii		13.86	0.42		99	
Actinomyces israelii		2.44	0.20		99	
63671101	**Good identification**					**63671101**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63671102	**Low discrimination**					**63671102**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		51.68	0.42	ESL 12	99	
Fusobacterium mortiferum		34.69	0.27	MNS 88	1	
Actinomyces israelii		13.64	0.23	MLT 94	99	
63671103	**Good identification**					**63671103**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63671120	**Good identification**					**63671120**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63671121	**Good identification**					**63671121**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63671122	**Good identification**					**63671122**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63671123	**Good identification**					**63671123**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63673000	**Low discrimination**					**63673000**
Actinomyces naeslundii		73.23	0.39	ESL 12		
Actinomyces israelii		26.77	0.22	MNS 88		
63673001	**Good identification**					**63673001**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63673002	**Low discrimination**					**63673002**
Actinomyces naeslundii		64.59	0.39	ESL 12		
Actinomyces israelii		35.41	0.26	MNS 88		
63673003	**Good identification**					**63673003**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63673020	**Good identification**					**63673020**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63673021	**Good identification**					**63673021**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63673022	**Good identification**					**63673022**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63673023	**Good identification**					**63673023**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63673100	**Low discrimination**					**63673100**
Actinomyces naeslundii		66.06	0.47	ESL 12		
Actinomyces israelii		33.94	0.33	MNS 88		
63673101	**Good identification**					**63673101**
Actinomyces naeslundii		100.0	0.50	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63673102	**Low discrimination**						**63673102**
Actinomyces naeslundii		56.48	0.47	ESL 12			
Actinomyces israelii		43.52	0.36	MNS 88			
63673103	**Good identification**						**63673103**
Actinomyces naeslundii		100.0	0.50	ESL 12			
63673120	**Good identification**						**63673120**
Actinomyces naeslundii		100.0	0.47	ESL 12			
63673121	**Good identification**						**63673121**
Actinomyces naeslundii		100.0	0.50	ESL 12			
63673122	**Good identification**						**63673122**
Actinomyces naeslundii		95.31	0.47	ESL 12			
Actinomyces israelii		4.69	0.15	MNS 88			
63673123	**Good identification**						**63673123**
Actinomyces naeslundii		100.0	0.50	ESL 12			
63673141	**Good identification**						**63673141**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63673143	**Good identification**						**63673143**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63673161	**Good identification**						**63673161**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63673163	**Good identification**						**63673163**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63675100	**Low discrimination**						**63675100**
					GRA		
Actinomyces israelii		81.66	0.38	MLT 99	99		
Fusobacterium mortiferum		18.35	0.25		1		
63675102	**Good identification**						**63675102**
Actinomyces israelii		100.0	0.41	MLT 99			
63675500	**Low discrimination**						**63675500**
					GRA	SPO	
Actinomyces israelii		57.23	0.37	MLT 99	99	1	
Mitsuokella multacida		37.66	0.21		1	1	
Clostridium butyricum		5.11	-.03		99	99	
63675502	**Low discrimination**						**63675502**
					GRA		
Actinomyces israelii		89.71	0.40	MLT 99	99		
Bifidobacterium adolescentis		5.92	0.13		99		
Mitsuokella multacida		4.37	0.04		1		
63677000	**Good identification**						**63677000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.41	MLT 99		
63677002	**Good identification**					**63677002**
Actinomyces israelii		100.0	0.44	MLT 99		
63677042	**Good identification**					**63677042**
Actinomyces israelii		100.0	0.37	MLT 99		
63677100	**Good identification**					**63677100**
Actinomyces israelii		100.0	0.51	MLT 99		
63677102	**Good identification**					**63677102**
Actinomyces israelii		100.0	0.54	MLT 99		
63677110	**Good identification**					**63677110**
Actinomyces israelii		100.0	0.37	MLT 99		
63677112	**Good identification**					**63677112**
Actinomyces israelii		100.0	0.40	MLT 99		
63677140	**Good identification**					**63677140**
Actinomyces israelii		100.0	0.44	MLT 99		
63677142	**Good identification**					**63677142**
Actinomyces israelii		100.0	0.47	MLT 99		
63677400	**Good identification**					**63677400**
Actinomyces israelii		100.0	0.39	MLT 99		
63677402	**Good identification**					**63677402**
Actinomyces israelii		100.0	0.42	MLT 99		
63677442	**Good identification**					**63677442**
Actinomyces israelii		100.0	0.35	MLT 99		
63677500	**Low discrimination**					**63677500**
					GRA	
Actinomyces israelii		78.68	0.49	MLT 99	99	
Mitsuokella multacida		21.32	0.27		1	
63677502	**Good identification**					**63677502**
					GRA	
Actinomyces israelii		97.13	0.52	MLT 99	99	
Mitsuokella multacida		1.95	0.11		1	
Bifidobacterium adolescentis		0.93	0.12		99	
63677510	**Low discrimination**					**63677510**
					GRA	
Actinomyces israelii		62.33	0.35	MLT 99	99	
Mitsuokella multacida		37.67	0.19	MLZ 13	1	
63677512	**Low discrimination**					**63677512**
Actinomyces israelii		84.46	0.38	MLT 99		
Bifidobacterium adolescentis		6.59	0.13	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium dentium		5.18	-.07			
63677540	**Low discrimination**					**63677540**
					GRA	
Actinomyces israelii		58.94	0.42	MLT 99	99	
Mitsuokella multacida		41.06	0.27		1	
63677542	**Good identification**					**63677542**
					GRA	
Actinomyces israelii		95.09	0.45	MLT 99	99	
Mitsuokella multacida		4.91	0.11		1	
63701000	**Very good identification**					**63701000**
Eubacterium tortuosum		99.89	0.56	MLT 1		
Actinomyces odontolyticus		0.08	0.14			
Actinomyces naeslundii		0.04	0.26			
63701002	**Good identification**					**63701002**
Eubacterium tortuosum		99.29	0.43	MLT 1		
Actinomyces odontolyticus		0.26	0.10			
Actinomyces viscosus		0.24	0.03			
63701020	**Good identification**					**63701020**
Eubacterium tortuosum		99.79	0.43	MLT 1		
Actinomyces naeslundii		0.21	0.26			
63701101	**Low discrimination**					**63701101**
					SPO	
Actinomyces naeslundii		64.50	0.37	ESL 12	1	
Clostridium chauvoei		35.50	0.09	MNS 88	99	
63701103	**Low discrimination**					**63701103**
Actinomyces naeslundii		86.33	0.37	ESL 12		
Actinomyces viscosus		13.67	-.01	MNS 88		
63701121	**Low discrimination**					**63701121**
					SPO	
Actinomyces naeslundii		84.50	0.37	ESL 12	1	
Clostridium chauvoei		15.50	0.01	MNS 88	99	
63701123	**Good identification**					**63701123**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63703100	**Good identification**					**63703100**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63703101	**Good identification**					**63703101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63703102	**Good identification**					**63703102**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63703103	**Good identification**					**63703103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63703120	**Good identification**					**63703120**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	ESL 12			
63703121	**Good identification**						**63703121**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63703122	**Good identification**						**63703122**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63703123	**Good identification**						**63703123**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63710000	**Low discrimination**						**63710000**
					GRA		
Actinomyces odontolyticus		64.19	0.35	ESL 1	99		
Fusobacterium mortiferum		34.38	0.36	RAF 3	1		
Actinomyces naeslundii		1.43	0.24		99		
63710100	**Low discrimination**						**63710100**
					GRA		
Fusobacterium mortiferum		75.89	0.42	NIT 1	1		
Actinomyces odontolyticus		19.18	0.27	MLT 11	99		
Actinomyces naeslundii		3.85	0.32		99		
63710200	**Good identification**						**63710200**
Actinomyces odontolyticus		100.0	0.40	ESL 1			
63710202	**Good identification**						**63710202**
Actinomyces odontolyticus		100.0	0.36	ESL 1			
63711000	**Low discrimination**						**63711000**
					GRA		
Fusobacterium mortiferum		28.77	0.39	NIT 1	1		
Eubacterium tortuosum		27.41	0.22	MLT 11	99		
Actinomyces odontolyticus		23.88	0.32		99		
63711001	**Good identification**						**63711001**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63711003	**Low discrimination**						**63711003**
Actinomyces naeslundii		79.79	0.51	ESL 12			
Actinomyces viscosus		20.22	0.16	MNS 88			
63711020	**Low discrimination**						**63711020**
Actinomyces naeslundii		63.95	0.48	ESL 12			
Eubacterium tortuosum		16.15	0.09	MNS 88			
Actinomyces odontolyticus		12.98	0.19				
63711021	**Good identification**						**63711021**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63711022	**Low discrimination**						**63711022**
					GRA	CAT	
Actinomyces naeslundii		59.11	0.48	ESL 12	99	25	
Capnocytophaga ochracea		23.44	0.15	MNS 88	1	90	
Actinomyces odontolyticus		7.35	0.15		99	10	
63711023	**Good identification**						**63711023**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.51	ESL 12			
63711041	**Good identification**						**63711041**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63711043	**Good identification**						**63711043**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63711061	**Good identification**						**63711061**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63711063	**Good identification**						**63711063**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63711100	**Low discrimination**						**63711100**
					GRA	CAT	
Fusobacterium mortiferum		45.40	0.45	NIT 1	1	5	
Actinomyces naeslundii		36.87	0.55	MLT 11	99	25	
Capnocytophaga ochracea		6.28	0.17		1	90	
63711101	**Good identification**						**63711101**
					GRA	SPO	
Actinomyces naeslundii		92.95	0.58	ESL 12	99	1	
Clostridium chauvoei		6.28	0.15	MNS 88	99	99	
Fusobacterium mortiferum		0.77	0.11		1	1	
63711103	**Low discrimination**						**63711103**
					SPO		
Actinomyces naeslundii		84.15	0.58	ESL 12	1		
Actinomyces viscosus		13.32	0.21	MNS 88	1		
Clostridium chauvoei		1.00	0.03		99		
63711111	**Good identification**						**63711111**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63711113	**Good identification**						**63711113**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63711121	**Very good identification**						**63711121**
					GRA	SPO	
Actinomyces naeslundii		96.82	0.58	ESL 12	99	1	
Clostridium chauvoei		2.18	0.07	MNS 88	99	99	
Capnocytophaga ochracea		1.00	-.01		1	1	
63711122	**Low discrimination**						**63711122**
					GRA	CAT	
Capnocytophaga ochracea		79.49	0.50	MNS 90	1	90	
Bifidobacterium longum subsp. infantis		11.94	0.32	SAL 1	99	1	
Actinomyces naeslundii		5.76	0.55		99	25	
63711123	**Good identification**						**63711123**
					GRA	CAT	
Actinomyces naeslundii		90.35	0.58	ESL 12	99	25	
Capnocytophaga ochracea		8.39	0.15	MNS 88	1	90	
Bifidobacterium longum subsp. infantis		1.26	-.03		99	1	
63711131	**Good identification**						**63711131**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.37	ESL 12			
63711133	**Low discrimination**						**63711133**
					GRA		
Actinomyces naeslundii		83.61	0.37	ESL 12	99		
Capnocytophaga ochracea		16.39	-.01	MNS 88	1		
63711140	**Low discrimination**						**63711140**
					GRA	CAT	
Actinomyces naeslundii		76.70	0.43	ESL 12	99	25	
Actinomyces israelii		9.68	0.18	MNS 88	99	1	
Capnocytophaga ochracea		8.22	0.01		1	90	
63711141	**Good identification**						**63711141**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63711142	**Low discrimination**						**63711142**
					GRA	CAT	
Actinomyces naeslundii		45.03	0.43	ESL 12	99	25	
Capnocytophaga ochracea		43.45	0.17	MNS 88	1	90	
Actinomyces israelii		8.52	0.21		99	1	
63711143	**Good identification**						**63711143**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63711160	**Low discrimination**						**63711160**
					GRA	CAT	
Actinomyces naeslundii		46.65	0.43	ESL 12	99	25	
Capnocytophaga ochracea		45.01	0.17	MNS 88	1	90	
Clostridium perfringens		8.33	0.07		99	1	
63711161	**Good identification**						**63711161**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63711163	**Acceptable identification**						**63711163**
					GRA		
Actinomyces naeslundii		94.48	0.46	ESL 12	99		
Capnocytophaga ochracea		5.52	-.01	MNS 88	1		
63711200	**Good identification**						**63711200**
Actinomyces odontolyticus		100.0	0.37	ESL 1			
63711522	**Good identification**						**63711522**
					GRA	CAT	
Capnocytophaga ochracea		98.67	0.37	MNS 90	1	90	
Bifidobacterium longum subsp. infantis		0.85	-.03	SAL 1	99	1	
Bifidobacterium longum subsp. longum		0.49	0.05		99	1	
63712100	**Low discrimination**						**63712100**
Actinomyces naeslundii		52.78	0.37	ESL 12			
Propionibacterium propionicum		47.22	0.26	MNS 88			
63712101	**Good identification**						**63712101**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63712103	**Acceptable identification**						**63712103**
Actinomyces naeslundii		90.96	0.40	ESL 12			
Propionibacterium propionicum		9.04	0.12	MNS 88			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63712120	**Good identification**					**63712120**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63712121	**Good identification**					**63712121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63712122	**Good identification**					**63712122**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63712123	**Good identification**					**63712123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63713000	**Low discrimination**					**63713000**
Actinomyces naeslundii		89.37	0.53	ESL 12		
Actinomyces israelii		10.63	0.28	MNS 88		
63713001	**Good identification**					**63713001**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63713002	**Low discrimination**					**63713002**
Actinomyces naeslundii		84.86	0.53	ESL 12		
Actinomyces israelii		15.14	0.31	MNS 88		
63713003	**Good identification**					**63713003**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63713020	**Good identification**					**63713020**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63713021	**Good identification**					**63713021**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63713022	**Good identification**					**63713022**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63713023	**Good identification**					**63713023**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63713040	**Low discrimination**					**63713040**
Actinomyces naeslundii		79.24	0.40	ESL 12		
Actinomyces israelii		20.77	0.21	MNS 88		
63713041	**Good identification**					**63713041**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63713042	**Low discrimination**					**63713042**
Actinomyces naeslundii		71.78	0.40	ESL 12		
Actinomyces israelii		28.22	0.24	MNS 88		
63713043	**Good identification**					**63713043**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.43	ESL 12			
63713060	**Good identification**						**63713060**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63713061	**Good identification**						**63713061**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63713062	**Good identification**						**63713062**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63713063	**Good identification**						**63713063**
Actinomyces naeslundii		100.0	0.43	ESL 12			
63713100	**Low discrimination**						**63713100**
Actinomyces naeslundii		82.59	0.60	ESL 12			
Actinomyces israelii		13.80	0.38	MNS 88			
Propionibacterium propionicum		3.08	0.26				
63713101	**Very good identification**						**63713101**
Actinomyces naeslundii		100.0	0.63	ESL 12			
63713102	**Low discrimination**						**63713102**
Actinomyces naeslundii		74.72	0.60	ESL 12			
Actinomyces israelii		18.73	0.41	MNS 88			
Propionibacterium propionicum		4.18	0.29				
63713103	**Very good identification**						**63713103**
Actinomyces naeslundii		99.59	0.63	ESL 12			
Propionibacterium propionicum		0.41	0.12	MNS 88			
63713110	**Low discrimination**						**63713110**
Actinomyces naeslundii		67.82	0.38	ESL 12			
Actinomyces israelii		32.18	0.24	MNS 88			
63713111	**Good identification**						**63713111**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63713112	**Low discrimination**						**63713112**
Actinomyces naeslundii		47.10	0.38	ESL 12			
Actinomyces israelii		33.52	0.27	MNS 88			
Bifidobacterium adolescentis		19.37	0.16	MLZ 5			
63713113	**Good identification**						**63713113**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63713120	**Very good identification**						**63713120**
					GRA	CAT	
Actinomyces naeslundii		98.17	0.60	ESL 12	99	25	
Actinomyces israelii		1.05	0.17	MNS 88	99	1	
Capnocytophaga ochracea		0.78	-.01		1	90	
63713121	**Very good identification**						**63713121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.63	ESL 12			
63713122	**Good identification**						**63713122**
					GRA	CAT	
Actinomyces naeslundii		91.03	0.60	ESL 12	99	25	
Capnocytophaga ochracea		6.53	0.15	MNS 88	1	90	
Actinomyces israelii		1.46	0.20		99	1	
63713123	**Very good identification**						**63713123**
Actinomyces naeslundii		100.0	0.63	ESL 12			
63713130	**Good identification**						**63713130**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63713131	**Good identification**						**63713131**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63713132	**Low discrimination**						**63713132**
					GRA		
Actinomyces naeslundii		86.85	0.38	ESL 12	99		
Capnocytophaga ochracea		13.16	-.01	MNS 88	1		
63713133	**Good identification**						**63713133**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63713140	**Low discrimination**						**63713140**
Actinomyces naeslundii		63.31	0.47	ESL 12			
Actinomyces israelii		23.32	0.31	MNS 88			
Propionibacterium propionicum		13.38	0.26				
63713141	**Good identification**						**63713141**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63713142	**Low discrimination**						**63713142**
Actinomyces naeslundii		53.49	0.47	ESL 12			
Actinomyces israelii		29.55	0.34	MNS 88			
Propionibacterium propionicum		16.95	0.29				
63713143	**Good identification**						**63713143**
Actinomyces naeslundii		97.71	0.51	ESL 12			
Propionibacterium propionicum		2.29	0.12	MNS 88			
63713160	**Good identification**						**63713160**
Actinomyces naeslundii		100.0	0.47	ESL 12			
63713161	**Good identification**						**63713161**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63713162	**Acceptable identification**						**63713162**
					GRA	CAT	
Actinomyces naeslundii		92.56	0.47	ESL 12	99	25	
Capnocytophaga ochracea		4.18	-.01	MNS 88	1	90	
Actinomyces israelii		3.26	0.13		99	1	
63713163	**Good identification**						**63713163**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Actinomyces naeslundii	100.0	0.51	ESL 12			
63713500	**Acceptable identification**						**63713500**
	Actinomyces israelii	93.12	0.36	CEL 93			
	Actinomyces naeslundii	6.88	0.26	XYL 92			
63713502	**Low discrimination**						**63713502**
	Actinomyces israelii	61.46	0.39	CEL 93			
	Bifidobacterium adolescentis	35.51	0.29	XYL 92			
	Actinomyces naeslundii	3.03	0.26				
63715002	**Low discrimination**						**63715002**
	Actinomyces israelii	65.12	0.36	CEL 93			
	Actinomyces odontolyticus	34.88	0.23				
63715100	**Low discrimination**						**63715100**
	Actinomyces israelii	43.04	0.43	CEL 93			
	Eubacterium contortum	37.50	0.27				
	Bifidobacterium longum subsp. infantis	6.16	0.09				
63715102	**Low discrimination**						**63715102**
					GRA	CAT	
	Actinomyces israelii	35.94	0.46	CEL 93	99	1	
	Bifidobacterium longum subsp. infantis	34.68	0.26		99	1	
	Capnocytophaga ochracea	13.94	0.23		1	90	
63715122	**Low discrimination**						**63715122**
					GRA	CAT	
	Capnocytophaga ochracea	63.41	0.39	MNS 90	1	90	
	Bifidobacterium longum subsp. infantis	31.16	0.30	SAL 1	99	1	
	Bacteroides fragilis	2.47	0.06		1	99	
63715140	**Good identification**						**63715140**
	Actinomyces israelii	100.0	0.36	CEL 93			
63715142	**Low discrimination**						**63715142**
					GRA	CAT	
	Actinomyces israelii	83.24	0.39	CEL 93	99	1	
	Capnocytophaga ochracea	9.23	0.07		1	90	
	Bifidobacterium adolescentis	7.53	0.15		99	1	
63715300	**Very good identification**						**63715300**
	Eubacterium contortum	99.68	0.61	NIT 1			
	Actinomyces odontolyticus	0.23	0.23				
	Actinomyces israelii	0.09	0.24				
63715302	**Good identification**						**63715302**
	Eubacterium contortum	98.47	0.48	NIT 1			
	Actinomyces odontolyticus	0.80	0.20				
	Actinomyces israelii	0.73	0.27				
63715320	**Good identification**						**63715320**
					GRA		
	Eubacterium contortum	99.27	0.48	NIT 1	99		
	Bacteroides vulgatus	0.52	0.02		1		
	Actinomyces odontolyticus	0.21	0.10		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63715322	**Acceptable identification**						**63715322**
					GRA	CAT	
Eubacterium contortum		90.34	0.35	NIT 1	99	1	
Parabacteroides distasonis		5.34	0.26		1	76	
Capnocytophaga ochracea		1.78	0.05		1	90	
63715500	**Low discrimination**						**63715500**
Actinomyces israelii		52.45	0.42	CEL 93			
Eubacterium contortum		45.70	0.26				
Bifidobacterium adolescentis		1.85	0.11				
63715502	**Low discrimination**						**63715502**
Bifidobacterium adolescentis		66.25	0.45	NIT 1			
Actinomyces israelii		28.45	0.45	MNS 87			
Eubacterium contortum		2.92	0.13				
63715512	**Acceptable identification**						**63715512**
Bifidobacterium adolescentis		94.41	0.46	NIT 1			
Actinomyces israelii		4.96	0.31	MNS 87			
Bifidobacterium longum subsp. longum		0.63	0.07				
63715542	**Low discrimination**						**63715542**
Actinomyces israelii		57.47	0.38	CEL 93			
Bifidobacterium adolescentis		42.53	0.29				
63715700	**Very good identification**						**63715700**
Eubacterium contortum		99.91	0.60	NIT 1			
Actinomyces israelii		0.09	0.22				
63715702	**Good identification**						**63715702**
Eubacterium contortum		99.04	0.47	NIT 1			
Actinomyces israelii		0.74	0.25				
Bifidobacterium adolescentis		0.23	0.11				
63715720	**Acceptable identification**						**63715720**
					GRA		
Eubacterium contortum		92.16	0.47	NIT 1	99		
Bacteroides vulgatus		7.84	0.21		1		
63717000	**Good identification**						**63717000**
Actinomyces israelii		100.0	0.46	CEL 93			
63717002	**Good identification**						**63717002**
Actinomyces israelii		100.0	0.49	CEL 93			
63717040	**Good identification**						**63717040**
Actinomyces israelii		100.0	0.39	CEL 93			
63717042	**Good identification**						**63717042**
Actinomyces israelii		100.0	0.42	CEL 93			
63717100	**Good identification**						**63717100**
Actinomyces israelii		99.48	0.56	CEL 93			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.52	0.26				
63717102	**Very good identification**						**63717102**
Actinomyces israelii		95.73	0.59	CEL 93			
Bifidobacterium adolescentis		3.94	0.29				
Actinomyces naeslundii		0.34	0.26				
63717110	**Good identification**						**63717110**
Actinomyces israelii		100.0	0.42	CEL 93			
63717112	**Low discrimination**						**63717112**
Actinomyces israelii		74.84	0.45	CEL 93			
Bifidobacterium adolescentis		25.16	0.31	MLZ 13			
63717120	**Acceptable identification**						**63717120**
Actinomyces israelii		92.39	0.36	CEL 93			
Actinomyces naeslundii		7.61	0.26	NAG 6			
63717122	**Low discrimination**						**63717122**
					GRA	CAT	
Actinomyces israelii		83.31	0.39	CEL 93	99	1	
Capnocytophaga ochracea		8.12	0.05	NAG 6	1	90	
Actinomyces naeslundii		4.57	0.26		99	25	
63717140	**Good identification**						**63717140**
Actinomyces israelii		100.0	0.49	CEL 93			
63717142	**Good identification**						**63717142**
Actinomyces israelii		98.71	0.52	CEL 93			
Bifidobacterium adolescentis		1.29	0.13				
63717152	**Acceptable identification**						**63717152**
Actinomyces israelii		90.35	0.38	CEL 93			
Bifidobacterium adolescentis		9.66	0.15	MLZ 13			
63717302	**Low discrimination**						**63717302**
Actinomyces israelii		80.63	0.40	CEL 93			
Eubacterium contortum		19.38	0.14	RHA 7			
63717400	**Good identification**						**63717400**
Actinomyces israelii		100.0	0.44	CEL 93			
63717402	**Good identification**						**63717402**
Actinomyces israelii		98.66	0.47	CEL 93			
Bifidobacterium adolescentis		1.34	0.09				
63717440	**Good identification**						**63717440**
Actinomyces israelii		100.0	0.37	CEL 93			
63717442	**Good identification**						**63717442**
Actinomyces israelii		100.0	0.40	CEL 93			
63717500	**Good identification**						**63717500**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		99.49	0.55	CEL 93		
Bifidobacterium adolescentis		0.51	0.09			
63717502	**Low discrimination**					**63717502**
Actinomyces israelii		74.84	0.58	CEL 93		
Bifidobacterium adolescentis		25.16	0.43			
63717510	**Good identification**					**63717510**
Actinomyces israelii		96.00	0.40	CEL 93		
Bifidobacterium adolescentis		4.00	0.11	MLZ 13		
63717512	**Low discrimination**					**63717512**
Bifidobacterium adolescentis		73.34	0.45	NIT 1		
Actinomyces israelii		26.66	0.43	MNS 87		
63717522	**Acceptable identification**					**63717522**
Actinomyces israelii		94.95	0.37	CEL 93		
Bifidobacterium adolescentis		5.05	0.09	NAG 6		
63717540	**Good identification**					**63717540**
Actinomyces israelii		100.0	0.48	CEL 93		
63717542	**Acceptable identification**					**63717542**
Actinomyces israelii		90.35	0.51	CEL 93		
Bifidobacterium adolescentis		9.66	0.28			
63717552	**Low discrimination**					**63717552**
Actinomyces israelii		53.36	0.36	CEL 93		
Bifidobacterium adolescentis		46.64	0.29	MLZ 13		
63717702	**Low discrimination**					**63717702**
Actinomyces israelii		77.80	0.38	CEL 93		
Eubacterium contortum		18.70	0.13	RHA 7		
Bifidobacterium adolescentis		3.51	0.09			
63721100	**Good identification**					**63721100**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63721101	**Good identification**					**63721101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63721102	**Good identification**					**63721102**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63721103	**Good identification**					**63721103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63721120	**Good identification**					**63721120**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63721121	**Good identification**					**63721121**
Actinomyces naeslundii		100.0	0.42	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63721122	**Good identification**					**63721122**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63721123	**Good identification**					**63721123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63723000	**Good identification**					**63723000**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63723001	**Good identification**					**63723001**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63723002	**Good identification**					**63723002**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63723003	**Good identification**					**63723003**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63723020	**Good identification**					**63723020**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63723021	**Good identification**					**63723021**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63723022	**Good identification**					**63723022**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63723023	**Good identification**					**63723023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63723100	**Acceptable identification**					**63723100**
Actinomyces naeslundii		94.15	0.43	ESL 12		
Actinomyces israelii		5.85	0.14	MNS 88		
63723101	**Good identification**					**63723101**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63723102	**Acceptable identification**					**63723102**
Actinomyces naeslundii		91.47	0.43	ESL 12		
Actinomyces israelii		8.53	0.17	MNS 88		
63723103	**Good identification**					**63723103**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63723120	**Good identification**					**63723120**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63723121	**Good identification**					**63723121**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63723122	**Good identification**					**63723122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ESL 12		
63723123	**Good identification**					**63723123**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63730101	**Good identification**					**63730101**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63730103	**Low discrimination**					**63730103**
Actinomyces naeslundii		80.63	0.40	ESL 12		
Actinomyces viscosus		19.37	0.05	MNS 88		
63730120	**Good identification**					**63730120**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63730121	**Good identification**					**63730121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63730122	**Good identification**					**63730122**
Actinomyces naeslundii		100.0	0.37	ESL 12		
63730123	**Good identification**					**63730123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63731000	**Low discrimination**					**63731000**
					GRA	
Actinomyces naeslundii		76.33	0.53	ESL 12	99	
Fusobacterium mortiferum		17.64	0.30	MNS 88	1	
Actinomyces israelii		6.03	0.25		99	
63731001	**Good identification**					**63731001**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63731002	**Low discrimination**					**63731002**
Actinomyces naeslundii		84.15	0.53	ESL 12		
Actinomyces israelii		9.97	0.28	MNS 88		
Actinomyces viscosus		3.72	0.05			
63731003	**Good identification**					**63731003**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63731020	**Acceptable identification**					**63731020**
Actinomyces naeslundii		94.21	0.53	ESL 12		
Clostridium perfringens		5.79	0.10	MNS 88		
63731021	**Good identification**					**63731021**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63731022	**Good identification**					**63731022**
Actinomyces naeslundii		97.19	0.53	ESL 12		
Clostridium perfringens		2.81	0.04	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63731023	**Good identification**					**63731023**
Actinomyces naeslundii		100.0	0.56	ESL 12		
63731040	**Low discrimination**					**63731040**
Actinomyces naeslundii		85.18	0.40	ESL 12		
Actinomyces israelii		14.82	0.18	MNS 88		
63731041	**Good identification**					**63731041**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63731042	**Low discrimination**					**63731042**
Actinomyces naeslundii		79.30	0.40	ESL 12		
Actinomyces israelii		20.70	0.21	MNS 88		
63731043	**Good identification**					**63731043**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63731060	**Good identification**					**63731060**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63731061	**Good identification**					**63731061**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63731062	**Good identification**					**63731062**
Actinomyces naeslundii		100.0	0.40	ESL 12		
63731063	**Good identification**					**63731063**
Actinomyces naeslundii		100.0	0.43	ESL 12		
63731100	**Low discrimination**					**63731100**
					GRA	
Actinomyces naeslundii		76.90	0.60	ESL 12	99	
Fusobacterium mortiferum		14.57	0.36	MNS 88	1	
Actinomyces israelii		8.53	0.35		99	
63731101	**Very good identification**					**63731101**
Actinomyces naeslundii		100.0	0.63	ESL 12		
63731102	**Low discrimination**					**63731102**
Actinomyces naeslundii		80.41	0.60	ESL 12		
Actinomyces israelii		13.38	0.38	MNS 88		
Actinomyces viscosus		2.22	0.10			
63731103	**Very good identification**					**63731103**
Actinomyces naeslundii		99.39	0.63	ESL 12		
Actinomyces viscosus		0.61	0.01	MNS 88		
63731110	**Low discrimination**					**63731110**
Actinomyces naeslundii		76.05	0.38	ESL 12		
Actinomyces israelii		23.95	0.21	MNS 88		
63731111	**Good identification**					**63731111**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.42	ESL 12			
63731113	**Good identification**						**63731113**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63731120	**Low discrimination**						**63731120**
					GRA	CAT	
Actinomyces naeslundii		85.22	0.60	ESL 12	99	25	
Clostridium perfringens		13.50	0.24	MNS 88	99	1	
Capnocytophaga ochracea		0.68	-.01		1	90	
63731121	**Very good identification**						**63731121**
Actinomyces naeslundii		100.0	0.63	ESL 12			
63731122	**Low discrimination**						**63731122**
					GRA	CAT	
Actinomyces naeslundii		84.12	0.60	ESL 12	99	25	
Clostridium perfringens		6.27	0.19	MNS 88	99	1	
Capnocytophaga ochracea		6.04	0.15		1	90	
63731123	**Very good identification**						**63731123**
Actinomyces naeslundii		100.0	0.63	ESL 12			
63731130	**Good identification**						**63731130**
Actinomyces naeslundii		100.0	0.38	ESL 12			
63731131	**Good identification**						**63731131**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63731132	**Low discrimination**						**63731132**
					GRA	CAT	
Actinomyces naeslundii		75.97	0.38	ESL 12	99	25	
Leptotrichia buccalis		12.52	-.09	MNS 88	1	1	
Capnocytophaga ochracea		11.51	-.01	MLZ 5	1	90	
63731133	**Good identification**						**63731133**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63731140	**Low discrimination**						**63731140**
Actinomyces naeslundii		80.35	0.47	ESL 12			
Actinomyces israelii		19.65	0.28	MNS 88			
63731141	**Good identification**						**63731141**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63731142	**Low discrimination**						**63731142**
Actinomyces naeslundii		73.17	0.47	ESL 12			
Actinomyces israelii		26.83	0.31	MNS 88			
63731143	**Good identification**						**63731143**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63731160	**Low discrimination**						**63731160**
Actinomyces naeslundii		85.40	0.47	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		14.60	0.12	MNS 88			
63731161	**Good identification**						**63731161**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63731162	**Low discrimination**						**63731162**
					GRA	CAT	
Actinomyces naeslundii		88.84	0.47	ESL 12	99	25	
Clostridium perfringens		7.15	0.06	MNS 88	99	1	
Capnocytophaga ochracea		4.01	-.01		1	90	
63731163	**Good identification**						**63731163**
Actinomyces naeslundii		100.0	0.51	ESL 12			
63731502	**Low discrimination**						**63731502**
Actinomyces israelii		56.07	0.36	CEL 93			
Bifidobacterium adolescentis		39.77	0.28	XYL 92			
Actinomyces naeslundii		4.16	0.26				
63732001	**Good identification**						**63732001**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63732003	**Good identification**						**63732003**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63732021	**Good identification**						**63732021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63732023	**Good identification**						**63732023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63732100	**Low discrimination**						**63732100**
Actinomyces naeslundii		68.70	0.42	ESL 12			
Propionibacterium propionicum		25.91	0.24	MNS 88			
Actinomyces israelii		5.39	0.14	GAL 96			
63732101	**Good identification**						**63732101**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63732102	**Low discrimination**						**63732102**
Actinomyces naeslundii		59.40	0.42	ESL 12			
Propionibacterium propionicum		33.61	0.27	MNS 88			
Actinomyces israelii		7.00	0.17	GAL 96			
63732103	**Good identification**						**63732103**
Actinomyces naeslundii		100.0	0.45	ESL 12			
63732120	**Good identification**						**63732120**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63732121	**Good identification**						**63732121**
Actinomyces naeslundii		100.0	0.45	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63732122	**Good identification**					**63732122**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63732123	**Good identification**					**63732123**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63733000	**Low discrimination**					**63733000**
Actinomyces naeslundii		81.27	0.58	ESL 12		
Actinomyces israelii		18.73	0.37	MNS 88		
63733001	**Very good identification**					**63733001**
Actinomyces naeslundii		100.0	0.61	ESL 12		
63733002	**Low discrimination**					**63733002**
Actinomyces naeslundii		74.31	0.58	ESL 12		
Actinomyces israelii		25.69	0.40	MNS 88		
63733003	**Very good identification**					**63733003**
Actinomyces naeslundii		100.0	0.61	ESL 12		
63733010	**Low discrimination**					**63733010**
Actinomyces naeslundii		60.45	0.36	ESL 12		
Actinomyces israelii		39.55	0.23	MNS 88		
63733011	**Good identification**					**63733011**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63733012	**Low discrimination**					**63733012**
Actinomyces naeslundii		50.47	0.36	ESL 12		
Actinomyces israelii		49.53	0.26	MNS 88		
63733013	**Good identification**					**63733013**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63733020	**Very good identification**					**63733020**
Actinomyces naeslundii		98.55	0.58	ESL 12		
Actinomyces israelii		1.45	0.17	MNS 88		
63733021	**Very good identification**					**63733021**
Actinomyces naeslundii		100.0	0.61	ESL 12		
63733022	**Very good identification**					**63733022**
Actinomyces naeslundii		97.84	0.58	ESL 12		
Actinomyces israelii		2.16	0.20	MNS 88		
63733023	**Very good identification**					**63733023**
Actinomyces naeslundii		100.0	0.61	ESL 12		
63733030	**Good identification**					**63733030**
Actinomyces naeslundii		100.0	0.36	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63733031	**Good identification**					**63733031**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63733032	**Good identification**					**63733032**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63733033	**Good identification**					**63733033**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63733040	**Low discrimination**					**63733040**
Actinomyces naeslundii		66.32	0.45	ESL 12		
Actinomyces israelii		33.68	0.30	MNS 88		
63733041	**Good identification**					**63733041**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63733042	**Low discrimination**					**63733042**
Actinomyces naeslundii		56.76	0.45	ESL 12		
Actinomyces israelii		43.24	0.33	MNS 88		
63733043	**Good identification**					**63733043**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63733060	**Good identification**					**63733060**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63733061	**Good identification**					**63733061**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63733062	**Good identification**					**63733062**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63733063	**Good identification**					**63733063**
Actinomyces naeslundii		100.0	0.48	ESL 12		
63733100	**Low discrimination**					**63733100**
Actinomyces naeslundii		74.65	0.65	ESL 12		
Actinomyces israelii		24.18	0.48	MNS 88		
Propionibacterium propionicum		1.17	0.24			
63733101	**Very good identification**					**63733101**
Actinomyces naeslundii		99.78	0.68	ESL 12		
Actinomyces israelii		0.22	0.14	MNS 88		
63733102	**Low discrimination**					**63733102**
Actinomyces naeslundii		65.99	0.65	ESL 12		
Actinomyces israelii		32.06	0.51	MNS 88		
Propionibacterium propionicum		1.56	0.27			
63733103	**Very good identification**					**63733103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.67	0.68	ESL 12		
Actinomyces israelii		0.33	0.17	MNS 88		
63733110	**Low discrimination**					**63733110**
Actinomyces naeslundii		52.10	0.43	ESL 12		
Actinomyces israelii		47.91	0.34	MNS 88		
63733111	**Good identification**					**63733111**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63733112	**Low discrimination**					**63733112**
Actinomyces israelii		54.72	0.37	CEL 93		
Actinomyces naeslundii		39.67	0.43	XYL 92		
Bifidobacterium adolescentis		5.60	0.13	MLZ 13		
63733113	**Good identification**					**63733113**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63733120	**Very good identification**					**63733120**
Actinomyces naeslundii		97.97	0.65	ESL 12		
Actinomyces israelii		2.03	0.27	MNS 88		
63733121	**Very good identification**					**63733121**
Actinomyces naeslundii		100.0	0.68	ESL 12		
63733122	**Very good identification**					**63733122**
Actinomyces naeslundii		96.99	0.65	ESL 12		
Actinomyces israelii		3.01	0.30	MNS 88		
63733123	**Very good identification**					**63733123**
Actinomyces naeslundii		100.0	0.68	ESL 12		
63733130	**Acceptable identification**					**63733130**
Actinomyces naeslundii		94.46	0.43	ESL 12		
Actinomyces israelii		5.54	0.13	MNS 88		
63733131	**Good identification**					**63733131**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63733132	**Acceptable identification**					**63733132**
Actinomyces naeslundii		91.91	0.43	ESL 12		
Actinomyces israelii		8.09	0.16	MNS 88		
63733133	**Good identification**					**63733133**
Actinomyces naeslundii		100.0	0.46	ESL 12		
63733140	**Low discrimination**					**63733140**
Actinomyces naeslundii		55.47	0.52	ESL 12		
Actinomyces israelii		39.59	0.41	MNS 88		
Propionibacterium propionicum		4.94	0.24			
63733141	**Good identification**					**63733141**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	ESL 12		
63733142	**Low discrimination**					**63733142**
Actinomyces israelii		48.57	0.44	CEL 93		
Actinomyces naeslundii		45.37	0.52	XYL 92		
Propionibacterium propionicum		6.06	0.27			
63733143	**Good identification**					**63733143**
Actinomyces naeslundii		100.0	0.55	ESL 12		
63733160	**Good identification**					**63733160**
Actinomyces naeslundii		95.64	0.52	ESL 12		
Actinomyces israelii		4.36	0.20	MNS 88		
63733161	**Good identification**					**63733161**
Actinomyces naeslundii		100.0	0.55	ESL 12		
63733162	**Acceptable identification**					**63733162**
Actinomyces naeslundii		93.60	0.52	ESL 12		
Actinomyces israelii		6.40	0.23	MNS 88		
63733163	**Good identification**					**63733163**
Actinomyces naeslundii		100.0	0.55	ESL 12		
63733400	**Acceptable identification**					**63733400**
Actinomyces israelii		94.92	0.36	CEL 93		
Actinomyces naeslundii		5.08	0.23	XYL 92		
63733402	**Good identification**					**63733402**
Actinomyces israelii		96.55	0.39	CEL 93		
Actinomyces naeslundii		3.45	0.23	XYL 92		
63733500	**Good identification**					**63733500**
Actinomyces israelii		96.33	0.46	CEL 93		
Actinomyces naeslundii		3.67	0.31	XYL 92		
63733502	**Low discrimination**					**63733502**
Actinomyces israelii		88.67	0.49	CEL 93		
Bifidobacterium adolescentis		9.08	0.26	XYL 92		
Actinomyces naeslundii		2.25	0.31			
63733512	**Low discrimination**					**63733512**
Actinomyces israelii		54.42	0.35	CEL 93		
Bifidobacterium adolescentis		45.58	0.28	XYL 92		
63733540	**Good identification**					**63733540**
Actinomyces israelii		100.0	0.39	CEL 93		
63733542	**Good identification**					**63733542**
Actinomyces israelii		96.85	0.42	CEL 93		
Bifidobacterium adolescentis		3.15	0.10	XYL 92		
63735000	**Good identification**					**63735000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		98.12	0.43	CEL 93	99		
Fusobacterium mortiferum		1.88	0.11		1		
63735002	**Good identification**						**63735002**
Actinomyces israelii		100.0	0.46	CEL 93			
63735040	**Good identification**						**63735040**
Actinomyces israelii		100.0	0.36	CEL 93			
63735042	**Good identification**						**63735042**
Actinomyces israelii		100.0	0.39	CEL 93			
63735100	**Good identification**						**63735100**
					GRA		
Actinomyces israelii		98.13	0.53	CEL 93	99		
Fusobacterium mortiferum		1.10	0.17		1		
Actinomyces naeslundii		0.78	0.26		99		
63735102	**Acceptable identification**						**63735102**
Actinomyces israelii		94.72	0.56	CEL 93			
Bifidobacterium adolescentis		4.78	0.28				
Actinomyces naeslundii		0.50	0.26				
63735110	**Good identification**						**63735110**
Actinomyces israelii		100.0	0.39	CEL 93			
63735112	**Low discrimination**						**63735112**
Actinomyces israelii		70.79	0.42	CEL 93			
Bifidobacterium adolescentis		29.21	0.29	MLZ 13			
63735122	**Low discrimination**						**63735122**
					GRA	CAT	
Actinomyces israelii		50.74	0.36	CEL 93	99	1	
Parabacteroides distasonis		33.95	0.29	NAG 6	1	76	
Capnocytophaga ochracea		7.45	0.05		1	90	
63735140	**Good identification**						**63735140**
Actinomyces israelii		100.0	0.46	CEL 93			
63735142	**Good identification**						**63735142**
Actinomyces israelii		98.42	0.49	CEL 93			
Bifidobacterium adolescentis		1.58	0.12				
63735302	**Low discrimination**						**63735302**
Actinomyces israelii		73.42	0.37	CEL 93			
Eubacterium contortum		26.58	0.14	RHA 7			
63735322	**Acceptable identification**						**63735322**
					GRA		
Parabacteroides distasonis		93.02	0.37	NIT 1	1		
Eubacterium contortum		3.49	0.01	MNS 98	99		
Actinomyces israelii		3.49	0.16		99		
63735400	**Good identification**						**63735400**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.41	CEL 93			
63735402	**Good identification**						**63735402**
Actinomyces israelii		98.36	0.44	CEL 93			
Bifidobacterium adolescentis		1.64	0.07				
63735442	**Good identification**						**63735442**
Actinomyces israelii		100.0	0.37	CEL 93			
63735500	**Acceptable identification**						**63735500**
					GRA	SPO	
Actinomyces israelii		90.37	0.51	CEL 93	99	1	
Mitsuokella multacida		7.98	0.21		1	1	
Clostridium butyricum		1.08	-.03		99	99	
63735502	**Low discrimination**						**63735502**
					GRA		
Actinomyces israelii		70.46	0.55	CEL 93	99		
Bifidobacterium adolescentis		29.08	0.42		99		
Mitsuokella multacida		0.46	0.04		1		
63735510	**Low discrimination**						**63735510**
					GRA		
Actinomyces israelii		80.12	0.37	CEL 93	99		
Mitsuokella multacida		15.78	0.13	MLZ 13	1		
Bifidobacterium adolescentis		4.10	0.09		99		
63735512	**Low discrimination**						**63735512**
Bifidobacterium adolescentis		76.36	0.43	NIT 1			
Actinomyces israelii		22.62	0.40	MNS 87			
Clostridium clostridioforme		1.02	0.08				
63735540	**Low discrimination**						**63735540**
					GRA		
Actinomyces israelii		81.50	0.44	CEL 93	99		
Mitsuokella multacida		18.51	0.21		1		
63735542	**Low discrimination**						**63735542**
					GRA		
Actinomyces israelii		87.11	0.47	CEL 93	99		
Bifidobacterium adolescentis		11.43	0.26		99		
Mitsuokella multacida		1.47	0.04		1		
63735702	**Low discrimination**						**63735702**
					GRA		
Actinomyces israelii		58.03	0.35	CEL 93	99		
Eubacterium contortum		21.01	0.13	RHA 7	99		
Mitsuokella multacida		9.37	0.09		1		
63737000	**Good identification**						**63737000**
Actinomyces israelii		99.62	0.56	CEL 93			
Actinomyces naeslundii		0.38	0.23				
63737002	**Very good identification**						**63737002**
Actinomyces israelii		99.75	0.59	CEL 93			
Actinomyces naeslundii		0.25	0.23				
63737010	**Good identification**						**63737010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.41	CEL 93		
63737012	**Good identification**					**63737012**
Actinomyces israelii		100.0	0.44	CEL 93		
63737020	**Acceptable identification**					**63737020**
Actinomyces israelii		94.37	0.35	CEL 93		
Actinomyces naeslundii		5.63	0.23	NAG 6		
63737022	**Good identification**					**63737022**
Actinomyces israelii		96.17	0.38	CEL 93		
Actinomyces naeslundii		3.83	0.23	NAG 6		
63737040	**Good identification**					**63737040**
Actinomyces israelii		100.0	0.49	CEL 93		
63737042	**Good identification**					**63737042**
Actinomyces israelii		100.0	0.52	CEL 93		
63737052	**Good identification**					**63737052**
Actinomyces israelii		100.0	0.37	CEL 93		
63737100	**Very good identification**					**63737100**
Actinomyces israelii		99.73	0.66	CEL 93		
Actinomyces naeslundii		0.27	0.31			
63737102	**Very good identification**					**63737102**
Actinomyces israelii		99.10	0.69	CEL 93		
Bifidobacterium adolescentis		0.72	0.26			
Actinomyces naeslundii		0.18	0.31			
63737110	**Good identification**					**63737110**
Actinomyces israelii		100.0	0.52	CEL 93		
63737112	**Acceptable identification**					**63737112**
Actinomyces israelii		94.38	0.55	CEL 93		
Bifidobacterium adolescentis		5.62	0.28	MLZ 13		
63737120	**Good identification**					**63737120**
Actinomyces israelii		95.92	0.45	CEL 93		
Actinomyces naeslundii		4.08	0.31	NAG 6		
63737122	**Good identification**					**63737122**
Actinomyces israelii		97.25	0.48	CEL 93		
Actinomyces naeslundii		2.76	0.31	NAG 6		
63737140	**Very good identification**					**63737140**
Actinomyces israelii		100.0	0.59	CEL 93		
63737142	**Very good identification**					**63737142**
Actinomyces israelii		99.77	0.62	CEL 93		
Bifidobacterium adolescentis		0.23	0.10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63737150	**Good identification**					**63737150**
Actinomyces israelii		100.0	0.45	CEL 93		
63737152	**Good identification**					**63737152**
Actinomyces israelii		98.14	0.48	CEL 93		
Bifidobacterium adolescentis		1.86	0.12	MLZ 13		
63737160	**Good identification**					**63737160**
Actinomyces israelii		100.0	0.38	CEL 93		
63737162	**Good identification**					**63737162**
Actinomyces israelii		100.0	0.41	CEL 93		
63737200	**Good identification**					**63737200**
Actinomyces israelii		100.0	0.36	CEL 93		
63737202	**Good identification**					**63737202**
Actinomyces israelii		100.0	0.39	CEL 93		
63737300	**Good identification**					**63737300**
Actinomyces israelii		100.0	0.47	CEL 93		
63737302	**Good identification**					**63737302**
Actinomyces israelii		100.0	0.50	CEL 93		
63737312	**Good identification**					**63737312**
Actinomyces israelii		100.0	0.36	CEL 93		
63737340	**Good identification**					**63737340**
Actinomyces israelii		100.0	0.40	CEL 93		
63737342	**Good identification**					**63737342**
Actinomyces israelii		100.0	0.43	CEL 93		
63737400	**Good identification**					**63737400**
Actinomyces israelii		100.0	0.54	CEL 93		
63737402	**Very good identification**					**63737402**
Actinomyces israelii		100.0	0.57	CEL 93		
63737410	**Good identification**					**63737410**
Actinomyces israelii		100.0	0.40	CEL 93		
63737412	**Good identification**					**63737412**
Actinomyces israelii		98.07	0.43	CEL 93		
Bifidobacterium adolescentis		1.93	0.07	MLZ 13		
63737422	**Good identification**					**63737422**
Actinomyces israelii		100.0	0.37	CEL 93		
63737440	**Good identification**					**63737440**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.47	CEL 93			
63737442	**Good identification**						**63737442**
Actinomyces israelii		100.0	0.50	CEL 93			
63737452	**Good identification**						**63737452**
Actinomyces israelii		100.0	0.36	CEL 93			
63737500	**Very good identification**						**63737500**
					GRA		
Actinomyces israelii		96.49	0.64	CEL 93	99		
Mitsuokella multacida		3.51	0.27		1		
63737502	**Good identification**						**63737502**
					GRA		
Actinomyces israelii		94.14	0.67	CEL 93	99		
Bifidobacterium adolescentis		5.61	0.40		99		
Mitsuokella multacida		0.25	0.11		1		
63737510	**Acceptable identification**						**63737510**
					GRA		
Actinomyces israelii		91.87	0.50	CEL 93	99		
Mitsuokella multacida		7.45	0.19	MLZ 13	1		
Bifidobacterium adolescentis		0.68	0.07		99		
63737512	**Low discrimination**						**63737512**
Actinomyces israelii		66.59	0.53	CEL 93			
Bifidobacterium adolescentis		32.46	0.42	MLZ 13			
Bifidobacterium dentium		0.55	-.07				
63737520	**Low discrimination**						**63737520**
					GRA		
Actinomyces israelii		87.53	0.44	CEL 93	99		
Mitsuokella multacida		12.47	0.17	NAG 6	1		
63737522	**Good identification**						**63737522**
Actinomyces israelii		100.0	0.47	CEL 93			
63737540	**Good identification**						**63737540**
					GRA		
Actinomyces israelii		91.45	0.57	CEL 93	99		
Mitsuokella multacida		8.55	0.27		1		
63737542	**Very good identification**						**63737542**
					GRA		
Actinomyces israelii		97.48	0.60	CEL 93	99		
Bifidobacterium adolescentis		1.85	0.25		99		
Mitsuokella multacida		0.68	0.11		1		
63737550	**Low discrimination**						**63737550**
					COC	GRA	
Actinomyces israelii		80.66	0.43	CEL 93	1	99	
Mitsuokella multacida		16.82	0.19	MLZ 13	1	1	
Blautia producta		2.52	-.03		99	99	
63737552	**Low discrimination**						**63737552**
					GRA		
Actinomyces israelii		85.44	0.46	CEL 93	99		
Bifidobacterium adolescentis		13.24	0.26	MLZ 13	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		1.32	0.03		1		
63737560	**Low discrimination**						**63737560**
					GRA		
Actinomyces israelii		73.20	0.37	CEL 93	99		
Mitsuokella multacida		26.80	0.17	NAG 6	1		
63737562	**Good identification**						**63737562**
Actinomyces israelii		100.0	0.40	CEL 93			
63737602	**Good identification**						**63737602**
Actinomyces israelii		100.0	0.38	CEL 93			
63737700	**Low discrimination**						**63737700**
					GRA		
Actinomyces israelii		52.71	0.45	CEL 93	99		
Mitsuokella multacida		47.29	0.32	RHA 7	1		
63737702	**Acceptable identification**						**63737702**
					GRA		
Actinomyces israelii		93.77	0.48	CEL 93	99		
Mitsuokella multacida		6.23	0.15	RHA 7	1		
63737742	**Low discrimination**						**63737742**
					GRA		
Actinomyces israelii		85.41	0.41	CEL 93	99		
Mitsuokella multacida		14.60	0.15	RHA 7	1		
63741000	**Good identification**						**63741000**
Eubacterium tortuosum		99.56	0.40	MLT 1			
Actinomyces naeslundii		0.44	0.28	CEL 10			
63741100	**Low discrimination**						**63741100**
Actinomyces naeslundii		55.53	0.36	ESL 12			
Eubacterium tortuosum		44.47	0.06	MNS 88			
63741101	**Good identification**						**63741101**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63741102	**Good identification**						**63741102**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63741103	**Good identification**						**63741103**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63741120	**Good identification**						**63741120**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63741121	**Good identification**						**63741121**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63741122	**Low discrimination**						**63741122**
					GRA	CAT	
Actinomyces naeslundii		59.25	0.36	ESL 12	99	25	
Capnocytophaga ochracea		29.58	0.05	MNS 88	1	90	
Prevotella oralis		11.18	-.03	FRU 95	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63741123	**Good identification**					**63741123**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63743001	**Good identification**					**63743001**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63743003	**Good identification**					**63743003**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63743021	**Good identification**					**63743021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63743023	**Good identification**					**63743023**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63743100	**Low discrimination**					**63743100**
Actinomyces naeslundii		75.67	0.41	ESL 12		
Actinomyces israelii		24.33	0.23	MNS 88		
63743101	**Good identification**					**63743101**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63743102	**Low discrimination**					**63743102**
Actinomyces naeslundii		67.47	0.41	ESL 12		
Actinomyces israelii		32.53	0.26	MNS 88		
63743103	**Good identification**					**63743103**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63743120	**Good identification**					**63743120**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63743121	**Good identification**					**63743121**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63743122	**Good identification**					**63743122**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63743123	**Good identification**					**63743123**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63747100	**Good identification**					**63747100**
Actinomyces israelii		100.0	0.41	FRU 99		
63747102	**Good identification**					**63747102**
Actinomyces israelii		100.0	0.44	FRU 99		
63747142	**Good identification**					**63747142**
Actinomyces israelii		100.0	0.37	FRU 99		
63747500	**Good identification**					**63747500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	FRU 99			
63747502	**Acceptable identification**						**63747502**
Actinomyces israelii		94.90	0.43	FRU 99			
Bifidobacterium adolescentis		5.10	0.15				
63747542	**Good identification**						**63747542**
Actinomyces israelii		100.0	0.36	FRU 99			
63750101	**Good identification**						**63750101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63750121	**Good identification**						**63750121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63750123	**Good identification**						**63750123**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63751000	**Low discrimination**						**63751000**
					GRA		
Actinomyces naeslundii		52.47	0.50	ESL 12	99		
Actinomyces israelii		21.42	0.34	MNS 88	99		
Fusobacterium mortiferum		19.82	0.31		1		
63751001	**Good identification**						**63751001**
Actinomyces naeslundii		100.0	0.53	ESL 12			
63751002	**Low discrimination**						**63751002**
Actinomyces naeslundii		53.11	0.50	ESL 12			
Actinomyces israelii		32.52	0.37	MNS 88			
Actinomyces viscosus		12.14	0.15				
63751003	**Good identification**						**63751003**
Actinomyces naeslundii		95.16	0.53	ESL 12			
Actinomyces viscosus		4.84	0.06	MNS 88			
63751020	**Good identification**						**63751020**
Actinomyces naeslundii		97.46	0.50	ESL 12			
Actinomyces israelii		2.54	0.13	MNS 88			
63751021	**Good identification**						**63751021**
Actinomyces naeslundii		100.0	0.53	ESL 12			
63751022	**Low discrimination**						**63751022**
					GRA	CAT	
Actinomyces naeslundii		77.38	0.50	ESL 12	99	25	
Prevotella oralis		13.82	0.11	MNS 88	1	1	
Capnocytophaga ochracea		5.79	0.05		1	90	
63751023	**Good identification**						**63751023**
Actinomyces naeslundii		100.0	0.53	ESL 12			
63751040	**Low discrimination**						**63751040**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		52.64	0.37	ESL 12			
Actinomyces israelii		47.36	0.27	MNS 88			
63751041	**Good identification**						**63751041**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63751043	**Good identification**						**63751043**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63751060	**Good identification**						**63751060**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63751061	**Good identification**						**63751061**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63751062	**Good identification**						**63751062**
Actinomyces naeslundii		100.0	0.37	ESL 12			
63751063	**Good identification**						**63751063**
Actinomyces naeslundii		100.0	0.40	ESL 12			
63751100	**Low discrimination**						**63751100**
					GRA		
Actinomyces naeslundii		50.79	0.58	ESL 12	99		
Actinomyces israelii		29.14	0.44	MNS 88	99		
Fusobacterium mortiferum		15.73	0.37		1		
63751101	**Very good identification**						**63751101**
Actinomyces naeslundii		100.0	0.61	ESL 12			
63751102	**Low discrimination**						**63751102**
					GRA	CAT	
Actinomyces naeslundii		38.48	0.58	ESL 12	99	25	
Actinomyces israelii		33.12	0.47	MNS 88	99	1	
Capnocytophaga ochracea		11.12	0.23		1	90	
63751103	**Very good identification**						**63751103**
Actinomyces naeslundii		96.92	0.61	ESL 12			
Actinomyces viscosus		3.08	0.11	MNS 88			
63751111	**Good identification**						**63751111**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63751113	**Good identification**						**63751113**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63751120	**Low discrimination**						**63751120**
					GRA	SPO	
Actinomyces naeslundii		63.14	0.58	ESL 12	99	1	
Capnocytophaga ochracea		18.25	0.23	MNS 88	1	1	
Clostridium paraputrificum		10.84	0.13		99	99	
63751121	**Very good identification**						**63751121**
Actinomyces naeslundii		100.0	0.61	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63751122	**Low discrimination**						**63751122**
					GRA	CAT	
Capnocytophaga ochracea		52.24	0.39	MNS 90	1	90	
Actinomyces naeslundii		20.08	0.58	SAL 1	99	25	
Prevotella oralis		19.74	0.32		1	1	
63751123	**Very good identification**						**63751123**
					GRA	CAT	
Actinomyces naeslundii		97.64	0.61	ESL 12	99	25	
Capnocytophaga ochracea		1.71	0.05	MNS 88	1	90	
Prevotella oralis		0.65	-.03		1	1	
63751130	**Low discrimination**						**63751130**
					GRA	CAT	
Actinomyces naeslundii		49.44	0.36	ESL 12	99	25	
Capnocytophaga ochracea		30.17	0.07	MNS 88	1	90	
Bifidobacterium breve		20.40	0.15	MLZ 5	99	1	
63751131	**Good identification**						**63751131**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63751133	**Good identification**						**63751133**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63751140	**Low discrimination**						**63751140**
Actinomyces israelii		54.19	0.37	XYL 92			
Actinomyces naeslundii		42.85	0.45				
Bifidobacterium breve		2.97	0.10				
63751141	**Good identification**						**63751141**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63751142	**Low discrimination**						**63751142**
Actinomyces israelii		50.20	0.40	XYL 92			
Actinomyces naeslundii		26.46	0.45				
Bifidobacterium breve		18.52	0.27				
63751143	**Good identification**						**63751143**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63751160	**Low discrimination**						**63751160**
					GRA	CAT	
Actinomyces naeslundii		68.54	0.45	ESL 12	99	25	
Capnocytophaga ochracea		12.47	0.07	MNS 88	1	90	
Bifidobacterium breve		8.43	0.15		99	1	
63751161	**Good identification**						**63751161**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63751163	**Good identification**						**63751163**
Actinomyces naeslundii		100.0	0.48	ESL 12			
63751402	**Good identification**						**63751402**
Actinomyces israelii		100.0	0.35	XYL 92			
63751500	**Good identification**						**63751500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		97.89	0.43	XYL 92			
Actinomyces naeslundii		2.11	0.23				
63751502	**Low discrimination**						**63751502**
					GRA	CAT	
Actinomyces israelii		57.91	0.46	XYL 92	99	1	
Bifidobacterium adolescentis		37.07	0.36		99	1	
Capnocytophaga ochracea		4.19	0.10		1	90	
63751512	**Low discrimination**						**63751512**
Bifidobacterium adolescentis		83.96	0.38	NIT 1			
Actinomyces israelii		16.04	0.32	MNS 87			
63751540	**Good identification**						**63751540**
Actinomyces israelii		100.0	0.36	XYL 92			
63751542	**Low discrimination**						**63751542**
Actinomyces israelii		83.10	0.39	XYL 92			
Bifidobacterium adolescentis		16.90	0.21				
63752100	**Low discrimination**						**63752100**
Actinomyces naeslundii		71.12	0.39	ESL 12			
Actinomyces israelii		28.88	0.23	MNS 88			
63752101	**Good identification**						**63752101**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63752102	**Low discrimination**						**63752102**
Actinomyces naeslundii		62.15	0.39	ESL 12			
Actinomyces israelii		37.85	0.26	MNS 88			
63752103	**Good identification**						**63752103**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63752120	**Good identification**						**63752120**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63752121	**Good identification**						**63752121**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63752122	**Good identification**						**63752122**
Actinomyces naeslundii		100.0	0.39	ESL 12			
63752123	**Good identification**						**63752123**
Actinomyces naeslundii		100.0	0.42	ESL 12			
63753000	**Low discrimination**						**63753000**
Actinomyces israelii		54.38	0.47	XYL 92			
Actinomyces naeslundii		45.62	0.55				
63753001	**Very good identification**						**63753001**
Actinomyces naeslundii		100.0	0.58	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63753002	**Low discrimination**					**63753002**
Actinomyces israelii		64.13	0.50	XYL 92		
Actinomyces naeslundii		35.87	0.55			
63753003	**Very good identification**					**63753003**
Actinomyces naeslundii		98.81	0.58	ESL 12		
Actinomyces israelii		1.19	0.16	MNS 88		
63753011	**Good identification**					**63753011**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63753012	**Low discrimination**					**63753012**
Actinomyces israelii		83.54	0.36	XYL 92		
Actinomyces naeslundii		16.46	0.33	MLZ 13		
63753013	**Good identification**					**63753013**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63753020	**Acceptable identification**					**63753020**
Actinomyces naeslundii		92.93	0.55	ESL 12		
Actinomyces israelii		7.07	0.26	MNS 88		
63753021	**Very good identification**					**63753021**
Actinomyces naeslundii		100.0	0.58	ESL 12		
63753022	**Low discrimination**					**63753022**
Actinomyces naeslundii		89.76	0.55	ESL 12		
Actinomyces israelii		10.24	0.29	MNS 88		
63753023	**Very good identification**					**63753023**
Actinomyces naeslundii		100.0	0.58	ESL 12		
63753031	**Good identification**					**63753031**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63753033	**Good identification**					**63753033**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63753040	**Low discrimination**					**63753040**
Actinomyces israelii		72.43	0.40	XYL 92		
Actinomyces naeslundii		27.57	0.42			
63753041	**Good identification**					**63753041**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63753042	**Low discrimination**					**63753042**
Actinomyces israelii		79.76	0.43	XYL 92		
Actinomyces naeslundii		20.24	0.42			
63753043	**Good identification**					**63753043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.45	ESL 12		
63753060	**Low discrimination**					**63753060**
Actinomyces naeslundii		85.64	0.42	ESL 12		
Actinomyces israelii		14.36	0.19	MNS 88		
63753061	**Good identification**					**63753061**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63753062	**Low discrimination**					**63753062**
Actinomyces naeslundii		79.91	0.42	ESL 12		
Actinomyces israelii		20.10	0.22	MNS 88		
63753063	**Good identification**					**63753063**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63753100	**Low discrimination**					**63753100**
Actinomyces israelii		62.47	0.57	XYL 92		
Actinomyces naeslundii		37.29	0.62			
Bifidobacterium breve		0.24	0.10			
63753101	**Very good identification**					**63753101**
Actinomyces naeslundii		98.89	0.65	ESL 12		
Actinomyces israelii		1.12	0.23	MNS 88		
63753102	**Low discrimination**					**63753102**
Actinomyces israelii		69.71	0.60	XYL 92		
Actinomyces naeslundii		27.74	0.62			
Bifidobacterium breve		1.77	0.27			
63753103	**Very good identification**					**63753103**
Actinomyces naeslundii		98.34	0.65	ESL 12		
Actinomyces israelii		1.66	0.26	MNS 88		
63753110	**Low discrimination**					**63753110**
Actinomyces israelii		80.95	0.43	XYL 92		
Actinomyces naeslundii		17.02	0.41	MLZ 13		
Bifidobacterium breve		2.03	0.10			
63753111	**Good identification**					**63753111**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63753112	**Low discrimination**					**63753112**
Actinomyces israelii		71.32	0.46	XYL 92		
Bifidobacterium breve		12.09	0.27	MLZ 13		
Actinomyces naeslundii		10.00	0.41			
63753113	**Good identification**					**63753113**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63753120	**Low discrimination**					**63753120**
Actinomyces naeslundii		88.98	0.62	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		9.51	0.37	MNS 88		
Bifidobacterium breve		1.00	0.15			
63753121	**Very good identification**					**63753121**
Actinomyces naeslundii		100.0	0.65	ESL 12		
63753122	**Low discrimination**					**63753122**
Actinomyces naeslundii		76.03	0.62	ESL 12		
Actinomyces israelii		12.20	0.40	MNS 88		
Bifidobacterium breve		8.60	0.32			
63753123	**Very good identification**					**63753123**
Actinomyces naeslundii		100.0	0.65	ESL 12		
63753130	**Low discrimination**					**63753130**
Actinomyces naeslundii		65.96	0.41	ESL 12		
Actinomyces israelii		20.02	0.23	MNS 88		
Bifidobacterium breve		14.02	0.15	MLZ 5		
63753131	**Good identification**					**63753131**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63753133	**Good identification**					**63753133**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63753140	**Low discrimination**					**63753140**
Actinomyces israelii		78.09	0.50	XYL 92		
Actinomyces naeslundii		21.16	0.50			
Bifidobacterium breve		0.75	0.10			
63753141	**Good identification**					**63753141**
Actinomyces naeslundii		97.58	0.53	ESL 12		
Actinomyces israelii		2.43	0.16	MNS 88		
63753142	**Low discrimination**					**63753142**
Actinomyces israelii		80.28	0.53	XYL 92		
Actinomyces naeslundii		14.50	0.50			
Bifidobacterium breve		5.23	0.27			
63753143	**Good identification**					**63753143**
Actinomyces naeslundii		96.41	0.53	ESL 12		
Actinomyces israelii		3.60	0.19	MNS 88		
63753150	**Low discrimination**					**63753150**
Actinomyces israelii		86.21	0.36	XYL 92		
Actinomyces naeslundii		8.23	0.28	MLZ 13		
Bifidobacterium breve		5.57	0.10			
63753152	**Low discrimination**					**63753152**
Actinomyces israelii		66.70	0.39	XYL 92		
Bifidobacterium breve		29.06	0.27	MLZ 13		
Actinomyces naeslundii		4.24	0.28			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63753160	**Low discrimination**					**63753160**
Actinomyces naeslundii		76.98	0.50	ESL 12		
Actinomyces israelii		18.14	0.30	MNS 88		
Bifidobacterium breve		4.88	0.15			
63753161	**Good identification**					**63753161**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63753162	**Low discrimination**					**63753162**
Actinomyces naeslundii		50.14	0.50	ESL 12		
Bifidobacterium breve		32.14	0.32	MNS 88		
Actinomyces israelii		17.72	0.33			
63753163	**Good identification**					**63753163**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63753300	**Acceptable identification**					**63753300**
Actinomyces israelii		92.58	0.38	XYL 92		
Actinomyces naeslundii		7.42	0.28	RHA 7		
63753302	**Acceptable identification**					**63753302**
Actinomyces israelii		94.93	0.41	XYL 92		
Actinomyces naeslundii		5.07	0.28	RHA 7		
63753400	**Good identification**					**63753400**
Actinomyces israelii		100.0	0.45	XYL 92		
63753402	**Good identification**					**63753402**
Actinomyces israelii		100.0	0.48	XYL 92		
63753440	**Good identification**					**63753440**
Actinomyces israelii		100.0	0.38	XYL 92		
63753442	**Good identification**					**63753442**
Actinomyces israelii		100.0	0.41	XYL 92		
63753500	**Good identification**					**63753500**
Actinomyces israelii		99.27	0.56	XYL 92		
Actinomyces naeslundii		0.73	0.28			
63753502	**Good identification**					**63753502**
Actinomyces israelii		91.13	0.59	XYL 92		
Bifidobacterium adolescentis		8.42	0.35			
Actinomyces naeslundii		0.45	0.28			
63753510	**Good identification**					**63753510**
Actinomyces israelii		100.0	0.41	XYL 92		
63753512	**Low discrimination**					**63753512**
Actinomyces israelii		56.95	0.45	XYL 92		
Bifidobacterium adolescentis		43.05	0.36	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63753520	**Low discrimination**						**63753520**
Actinomyces israelii		89.65	0.35	XYL 92			
Actinomyces naeslundii		10.35	0.28	NAG 6			
63753522	**Acceptable identification**						**63753522**
Actinomyces israelii		92.85	0.38	XYL 92			
Actinomyces naeslundii		7.15	0.28	NAG 6			
63753540	**Good identification**						**63753540**
Actinomyces israelii		100.0	0.49	XYL 92			
63753542	**Good identification**						**63753542**
Actinomyces israelii		97.15	0.52	XYL 92			
Bifidobacterium adolescentis		2.85	0.19				
63753552	**Low discrimination**						**63753552**
Actinomyces israelii		80.63	0.37	XYL 92			
Bifidobacterium adolescentis		19.37	0.21	MLZ 13			
63753700	**Good identification**						**63753700**
Actinomyces israelii		100.0	0.36	XYL 92			
63753702	**Good identification**						**63753702**
Actinomyces israelii		100.0	0.39	XYL 92			
63755000	**Good identification**						**63755000**
					GRA		
Actinomyces israelii		99.40	0.52		99		
Fusobacterium mortiferum		0.60	0.12		1		
63755002	**Good identification**						**63755002**
Actinomyces israelii		100.0	0.55				
63755010	**Good identification**						**63755010**
Actinomyces israelii		100.0	0.38	MLZ 13			
63755012	**Good identification**						**63755012**
Actinomyces israelii		100.0	0.41	MLZ 13			
63755040	**Good identification**						**63755040**
Actinomyces israelii		100.0	0.45				
63755042	**Good identification**						**63755042**
Actinomyces israelii		100.0	0.48				
63755100	**Very good identification**						**63755100**
					GRA	SPO	
Actinomyces israelii		99.30	0.62		99	1	
Fusobacterium mortiferum		0.35	0.18		1	1	
Clostridium butyricum		0.19	-.05		99	99	
63755102	**Good identification**						**63755102**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		94.89	0.65		99	1	
Bifidobacterium adolescentis		4.32	0.36		99	1	
Capnocytophaga ochracea		0.69	0.13		1	90	
63755110	**Good identification**						**63755110**
Actinomyces israelii		100.0	0.48	MLZ 13			
63755112	**Low discrimination**						**63755112**
Actinomyces israelii		72.86	0.51	MLZ 13			
Bifidobacterium adolescentis		27.14	0.38				
63755120	**Low discrimination**						**63755120**
					GRA	CAT	
Actinomyces israelii		83.65	0.42	NAG 6	99	1	
Capnocytophaga ochracea		14.35	0.13		1	90	
Actinomyces naeslundii		2.01	0.23		99	25	
63755140	**Good identification**						**63755140**
Actinomyces israelii		100.0	0.55				
63755142	**Very good identification**						**63755142**
Actinomyces israelii		98.57	0.58				
Bifidobacterium adolescentis		1.43	0.21				
63755150	**Good identification**						**63755150**
Actinomyces israelii		100.0	0.41	MLZ 13			
63755152	**Low discrimination**						**63755152**
Actinomyces israelii		89.41	0.44	MLZ 13			
Bifidobacterium adolescentis		10.59	0.22				
63755162	**Low discrimination**						**63755162**
					GRA	CAT	
Actinomyces israelii		77.28	0.38	NAG 6	99	1	
Capnocytophaga ochracea		22.72	0.13		1	90	
63755202	**Good identification**						**63755202**
Actinomyces israelii		100.0	0.36	RHA 7			
63755300	**Low discrimination**						**63755300**
Actinomyces israelii		53.44	0.43	RHA 7			
Eubacterium contortum		46.56	0.27				
63755302	**Acceptable identification**						**63755302**
Actinomyces israelii		90.70	0.46	RHA 7			
Eubacterium contortum		9.30	0.14				
63755340	**Good identification**						**63755340**
Actinomyces israelii		100.0	0.36	RHA 7			
63755342	**Good identification**						**63755342**
Actinomyces israelii		100.0	0.39	RHA 7			
63755400	**Good identification**						**63755400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.51				
63755402	**Good identification**						**63755402**
Actinomyces israelii		98.52	0.54				
Bifidobacterium adolescentis		1.48	0.16				
63755410	**Good identification**						**63755410**
Actinomyces israelii		100.0	0.36	MLZ 13			
63755412	**Low discrimination**						**63755412**
Actinomyces israelii		89.04	0.39	MLZ 13			
Bifidobacterium adolescentis		10.96	0.18				
63755440	**Good identification**						**63755440**
Actinomyces israelii		100.0	0.44				
63755442	**Good identification**						**63755442**
Actinomyces israelii		100.0	0.47				
63755500	**Very good identification**						**63755500**
					GRA	SPO	
Actinomyces israelii		95.05	0.61		99	1	
Mitsuokella multacida		2.38	0.21		1	1	
Clostridium butyricum		2.04	0.11		99	99	
63755502	**Low discrimination**						**63755502**
					SPO		
Actinomyces israelii		72.39	0.64		1		
Bifidobacterium adolescentis		26.96	0.50		1		
Clostridium butyricum		0.40	0.04		99		
63755510	**Acceptable identification**						**63755510**
					GRA		
Actinomyces israelii		90.75	0.47	MLZ 13	99		
Mitsuokella multacida		5.06	0.13		1		
Bifidobacterium adolescentis		4.19	0.18		99		
63755512	**Low discrimination**						**63755512**
Bifidobacterium adolescentis		75.29	0.52	NIT 1			
Actinomyces israelii		24.71	0.50	MNS 87			
63755520	**Low discrimination**						**63755520**
					GRA	CAT	
Actinomyces israelii		85.90	0.40	NAG 6	99	1	
Mitsuokella multacida		8.41	0.11		1	1	
Capnocytophaga ochracea		3.18	0.00		1	90	
63755522	**Low discrimination**						**63755522**
					GRA	CAT	
Actinomyces israelii		74.37	0.43	NAG 6	99	1	
Capnocytophaga ochracea		16.51	0.16		1	90	
Bacteroides uniformis		4.73	0.00		1	5	
63755540	**Acceptable identification**						**63755540**
					GRA	SPO	
Actinomyces israelii		93.54	0.54		99	1	
Mitsuokella multacida		6.01	0.21		1	1	
Clostridium butyricum		0.45	-.07		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63755542	**Low discrimination**						**63755542**
					GRA		
Actinomyces israelii		89.04	0.57		99		
Bifidobacterium adolescentis		10.54	0.35		99		
Mitsuokella multacida		0.42	0.04		1		
63755550	**Low discrimination**						**63755550**
					GRA		
Actinomyces israelii		87.46	0.40	MLZ 13	99		
Mitsuokella multacida		12.54	0.13		1		
63755552	**Low discrimination**						**63755552**
Actinomyces israelii		50.80	0.43	MLZ 13			
Bifidobacterium adolescentis		49.20	0.36				
63755562	**Acceptable identification**						**63755562**
					GRA	CAT	
Actinomyces israelii		94.04	0.36	NAG 6	99	1	
Capnocytophaga ochracea		5.97	0.00		1	90	
63755700	**Low discrimination**						**63755700**
					GRA		
Actinomyces israelii		40.19	0.42	RHA 7	99		
Eubacterium contortum		35.02	0.26		99		
Mitsuokella multacida		24.80	0.25		1		
63755702	**Low discrimination**						**63755702**
Actinomyces israelii		82.18	0.45	RHA 7			
Eubacterium contortum		8.42	0.13				
Bifidobacterium adolescentis		4.11	0.16				
63755742	**Low discrimination**						**63755742**
					GRA		
Actinomyces israelii		89.49	0.38	RHA 7	99		
Mitsuokella multacida		10.52	0.09		1		
63756100	**Good identification**						**63756100**
Actinomyces israelii		100.0	0.41	GAL 99			
63756102	**Good identification**						**63756102**
Actinomyces israelii		100.0	0.44	GAL 99			
63756142	**Good identification**						**63756142**
Actinomyces israelii		100.0	0.37	GAL 99			
63756500	**Good identification**						**63756500**
Actinomyces israelii		100.0	0.40	GAL 99			
63756502	**Acceptable identification**						**63756502**
Actinomyces israelii		94.90	0.43	GAL 99			
Bifidobacterium adolescentis		5.10	0.15				
63756542	**Good identification**						**63756542**
Actinomyces israelii		100.0	0.36	GAL 99			
63757000	**Very good identification**						**63757000**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces israelii	100.0	0.65			
63757002	**Very good identification**					**63757002**
	Actinomyces israelii	100.0	0.68			
63757010	**Good identification**					**63757010**
	Actinomyces israelii	100.0	0.51	MLZ 13		
63757012	**Good identification**					**63757012**
	Actinomyces israelii	100.0	0.54	MLZ 13		
63757020	**Good identification**					**63757020**
	Actinomyces israelii	100.0	0.45	NAG 6		
63757022	**Good identification**					**63757022**
	Actinomyces israelii	100.0	0.48	NAG 6		
63757040	**Very good identification**					**63757040**
	Actinomyces israelii	100.0	0.58			
63757042	**Very good identification**					**63757042**
	Actinomyces israelii	100.0	0.61			
63757050	**Good identification**					**63757050**
	Actinomyces israelii	100.0	0.44	MLZ 13		
63757052	**Good identification**					**63757052**
	Actinomyces israelii	100.0	0.47	MLZ 13		
63757060	**Good identification**					**63757060**
	Actinomyces israelii	100.0	0.37	NAG 6		
63757062	**Good identification**					**63757062**
	Actinomyces israelii	100.0	0.40	NAG 6		
63757100	**Very good identification**					**63757100**
	Actinomyces israelii	99.95	0.75			
	Actinomyces naeslundii	0.05	0.28			
63757101	**Acceptable identification**					**63757101**
	Actinomyces israelii	92.78	0.41	URE 1		
	Actinomyces naeslundii	7.22	0.31			
63757102	**Very good identification**					**63757102**
	Actinomyces israelii	99.31	0.78			
	Bifidobacterium adolescentis	0.65	0.35			
	Actinomyces naeslundii	0.04	0.28			
63757103	**Good identification**					**63757103**
	Actinomyces israelii	95.07	0.44	URE 1		
	Actinomyces naeslundii	4.94	0.31			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63757110	**Very good identification**					**63757110**
Actinomyces israelii		100.0	0.61	MLZ 13		
63757112	**Good identification**					**63757112**
Actinomyces israelii		94.90	0.64	MLZ 13		
Bifidobacterium adolescentis		5.10	0.36			
63757120	**Good identification**					**63757120**
Actinomyces israelii		99.19	0.55	NAG 6		
Actinomyces naeslundii		0.82	0.28			
63757122	**Very good identification**					**63757122**
Actinomyces israelii		99.46	0.58	NAG 6		
Actinomyces naeslundii		0.55	0.28			
63757130	**Good identification**					**63757130**
Actinomyces israelii		100.0	0.41	MLZ 13		
63757132	**Good identification**					**63757132**
Actinomyces israelii		100.0	0.44	MLZ 13		
63757140	**Very good identification**					**63757140**
Actinomyces israelii		100.0	0.68			
63757142	**Very good identification**					**63757142**
Actinomyces israelii		99.79	0.71			
Bifidobacterium adolescentis		0.21	0.19			
63757143	**Good identification**					**63757143**
Actinomyces israelii		100.0	0.37	URE 1		
63757150	**Good identification**					**63757150**
Actinomyces israelii		100.0	0.54	MLZ 13		
63757152	**Very good identification**					**63757152**
Actinomyces israelii		98.32	0.57	MLZ 13		
Bifidobacterium adolescentis		1.68	0.21			
63757160	**Good identification**					**63757160**
Actinomyces israelii		100.0	0.48	NAG 6		
63757162	**Good identification**					**63757162**
Actinomyces israelii		100.0	0.51	NAG 6		
63757172	**Good identification**					**63757172**
Actinomyces israelii		100.0	0.37	MLZ 13		
63757200	**Good identification**					**63757200**
Actinomyces israelii		100.0	0.46	RHA 7		
63757202	**Good identification**					**63757202**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.49	RHA 7		
63757240	**Good identification**					**63757240**
Actinomyces israelii		100.0	0.39	RHA 7		
63757242	**Good identification**					**63757242**
Actinomyces israelii		100.0	0.42	RHA 7		
63757300	**Good identification**					**63757300**
Actinomyces israelii		100.0	0.56	RHA 7		
63757302	**Very good identification**					**63757302**
Actinomyces israelii		100.0	0.59	RHA 7		
63757310	**Good identification**					**63757310**
Actinomyces israelii		100.0	0.42	RHA 7		
63757312	**Good identification**					**63757312**
Actinomyces israelii		100.0	0.45	RHA 7		
63757320	**Good identification**					**63757320**
Actinomyces israelii		100.0	0.36	RHA 7		
63757322	**Good identification**					**63757322**
Actinomyces israelii		100.0	0.39	RHA 7		
63757340	**Good identification**					**63757340**
Actinomyces israelii		100.0	0.49	RHA 7		
63757342	**Good identification**					**63757342**
Actinomyces israelii		100.0	0.52	RHA 7		
63757352	**Good identification**					**63757352**
Actinomyces israelii		100.0	0.38	RHA 7		
63757400	**Very good identification**					**63757400**
Actinomyces israelii		100.0	0.63			
63757402	**Very good identification**					**63757402**
Actinomyces israelii		99.78	0.66			
Bifidobacterium adolescentis		0.22	0.15			
63757410	**Good identification**					**63757410**
Actinomyces israelii		100.0	0.49	MLZ 13		
63757412	**Good identification**					**63757412**
Actinomyces israelii		98.25	0.52	MLZ 13		
Bifidobacterium adolescentis		1.75	0.16			
63757420	**Good identification**					**63757420**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.43	NAG 6		
63757422	**Good identification**					**63757422**
Actinomyces israelii		100.0	0.46	NAG 6		
63757440	**Very good identification**					**63757440**
Actinomyces israelii		100.0	0.56			
63757442	**Very good identification**					**63757442**
Actinomyces israelii		100.0	0.59			
63757450	**Good identification**					**63757450**
Actinomyces israelii		100.0	0.42	MLZ 13		
63757452	**Good identification**					**63757452**
Actinomyces israelii		100.0	0.45	MLZ 13		
63757460	**Good identification**					**63757460**
Actinomyces israelii		100.0	0.36	NAG 6		
63757462	**Good identification**					**63757462**
Actinomyces israelii		100.0	0.39	NAG 6		
63757500	**Very good identification**					**63757500**
					GRA	
Actinomyces israelii		98.86	0.74		99	
Mitsuokella multacida		1.02	0.27		1	
Bifidobacterium adolescentis		0.08	0.15		99	
63757501	**Low discrimination**					**63757501**
					GRA	
Actinomyces israelii		89.83	0.40	URE 1	99	
Mitsuokella multacida		10.17	0.11		1	
63757502	**Good identification**					**63757502**
					GRA	
Actinomyces israelii		94.83	0.77		99	
Bifidobacterium adolescentis		5.10	0.49		99	
Mitsuokella multacida		0.07	0.11		1	
63757503	**Acceptable identification**					**63757503**
Actinomyces israelii		94.90	0.43	URE 1		
Bifidobacterium adolescentis		5.10	0.15			
63757510	**Very good identification**					**63757510**
					GRA	
Actinomyces israelii		97.12	0.60	MLZ 13	99	
Mitsuokella multacida		2.23	0.19		1	
Bifidobacterium adolescentis		0.65	0.16		99	
63757512	**Low discrimination**					**63757512**
Actinomyces israelii		69.26	0.63	MLZ 13		
Bifidobacterium adolescentis		30.47	0.50			
Bifidobacterium dentium		0.16	-.07			
63757520	**Good identification**					**63757520**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		96.12	0.53	NAG 6	99		
Mitsuokella multacida		3.88	0.17		1		
63757522	**Very good identification**						**63757522**
Actinomyces israelii		99.16	0.56	NAG 6			
Bifidobacterium adolescentis		0.84	0.15				
63757530	**Acceptable identification**						**63757530**
					GRA		
Actinomyces israelii		91.75	0.39	MLZ 13	99		
Mitsuokella multacida		8.25	0.09	NAG 6	1		
63757532	**Low discrimination**						**63757532**
Actinomyces israelii		88.14	0.42	MLZ 13			
Bifidobacterium adolescentis		6.14	0.16	NAG 6			
Bifidobacterium dentium		5.72	-.03				
63757540	**Very good identification**						**63757540**
					GRA		
Actinomyces israelii		97.42	0.67		99		
Mitsuokella multacida		2.58	0.27		1		
63757542	**Very good identification**						**63757542**
					GRA		
Actinomyces israelii		98.13	0.70		99		
Bifidobacterium adolescentis		1.68	0.33		99		
Mitsuokella multacida		0.19	0.11		1		
63757543	**Good identification**						**63757543**
Actinomyces israelii		100.0	0.36	URE 1			
63757550	**Acceptable identification**						**63757550**
					COC	GRA	
Actinomyces israelii		93.64	0.53	MLZ 13	1	99	
Mitsuokella multacida		5.53	0.19		1	1	
Blautia producta		0.83	-.03		99	99	
63757552	**Low discrimination**						**63757552**
					GRA		
Actinomyces israelii		87.40	0.56	MLZ 13	99		
Bifidobacterium adolescentis		12.22	0.35		99		
Mitsuokella multacida		0.38	0.03		1		
63757560	**Acceptable identification**						**63757560**
					GRA		
Actinomyces israelii		90.61	0.46	NAG 6	99		
Mitsuokella multacida		9.40	0.17		1		
63757562	**Good identification**						**63757562**
Actinomyces israelii		100.0	0.49	NAG 6			
63757572	**Good identification**						**63757572**
Actinomyces israelii		100.0	0.35	MLZ 13			
63757600	**Good identification**						**63757600**
Actinomyces israelii		100.0	0.44	RHA 7			
63757602	**Good identification**						**63757602**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.47	RHA 7		
63757640	**Good identification**					**63757640**
Actinomyces israelii		100.0	0.37	RHA 7		
63757642	**Good identification**					**63757642**
Actinomyces israelii		100.0	0.40	RHA 7		
63757700	**Low discrimination**					**63757700**
					GRA	
Actinomyces israelii		79.74	0.55	RHA 7	99	
Mitsuokella multacida		20.26	0.32		1	
63757702	**Very good identification**					**63757702**
					GRA	
Actinomyces israelii		97.46	0.58	RHA 7	99	
Mitsuokella multacida		1.83	0.15		1	
Bifidobacterium adolescentis		0.70	0.15		99	
63757710	**Low discrimination**					**63757710**
					GRA	
Actinomyces israelii		63.83	0.40	RHA 7	99	
Mitsuokella multacida		36.17	0.24	MLZ 13	1	
63757712	**Acceptable identification**					**63757712**
					GRA	
Actinomyces israelii		90.83	0.43	RHA 7	99	
Bifidobacterium adolescentis		5.36	0.16	MLZ 13	99	
Mitsuokella multacida		3.81	0.07		1	
63757722	**Acceptable identification**					**63757722**
					GRA	
Actinomyces israelii		93.14	0.37	RHA 7	99	
Mitsuokella multacida		6.86	0.05	NAG 6	1	
63757740	**Low discrimination**					**63757740**
					GRA	
Actinomyces israelii		60.49	0.48	RHA 7	99	
Mitsuokella multacida		39.51	0.32		1	
63757742	**Good identification**					**63757742**
					GRA	
Actinomyces israelii		95.38	0.51	RHA 7	99	
Mitsuokella multacida		4.62	0.15		1	
63757752	**Acceptable identification**					**63757752**
					GRA	
Actinomyces israelii		90.26	0.36	RHA 7	99	
Mitsuokella multacida		9.74	0.07	MLZ 13	1	
63761001	**Good identification**					**63761001**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63761003	**Good identification**					**63761003**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63761021	**Good identification**					**63761021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63761023	**Good identification**					**63761023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	ESL 12		
63761100	**Low discrimination**					**63761100**
Actinomyces naeslundii		82.41	0.41	ESL 12		
Actinomyces israelii		17.59	0.20	MNS 88		
63761101	**Good identification**					**63761101**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63761102	**Low discrimination**					**63761102**
Actinomyces naeslundii		75.75	0.41	ESL 12		
Actinomyces israelii		24.25	0.23	MNS 88		
63761103	**Good identification**					**63761103**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63761120	**Good identification**					**63761120**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63761121	**Good identification**					**63761121**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63761122	**Good identification**					**63761122**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63761123	**Good identification**					**63761123**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63763000	**Low discrimination**					**63763000**
Actinomyces naeslundii		69.29	0.38	ESL 12		
Actinomyces israelii		30.72	0.22	MNS 88		
63763001	**Good identification**					**63763001**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63763002	**Low discrimination**					**63763002**
Actinomyces naeslundii		60.06	0.38	ESL 12		
Actinomyces israelii		39.94	0.26	MNS 88		
63763003	**Good identification**					**63763003**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63763020	**Good identification**					**63763020**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63763021	**Good identification**					**63763021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63763022	**Good identification**					**63763022**
Actinomyces naeslundii		100.0	0.38	ESL 12		
63763023	**Good identification**					**63763023**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.41	ESL 12			
63763100	**Low discrimination**						**63763100**
Actinomyces naeslundii		61.61	0.46	ESL 12			
Actinomyces israelii		38.39	0.33	MNS 88			
63763101	**Good identification**						**63763101**
Actinomyces naeslundii		100.0	0.49	ESL 12			
63763102	**Low discrimination**						**63763102**
Actinomyces naeslundii		51.69	0.46	ESL 12			
Actinomyces israelii		48.31	0.36	MNS 88			
63763103	**Good identification**						**63763103**
Actinomyces naeslundii		100.0	0.49	ESL 12			
63763120	**Good identification**						**63763120**
Actinomyces naeslundii		100.0	0.46	ESL 12			
63763121	**Good identification**						**63763121**
Actinomyces naeslundii		100.0	0.49	ESL 12			
63763122	**Acceptable identification**						**63763122**
Actinomyces naeslundii		94.37	0.46	ESL 12			
Actinomyces israelii		5.63	0.15	MNS 88			
63763123	**Good identification**						**63763123**
Actinomyces naeslundii		100.0	0.49	ESL 12			
63763141	**Good identification**						**63763141**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63763143	**Good identification**						**63763143**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63763161	**Good identification**						**63763161**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63763163	**Good identification**						**63763163**
Actinomyces naeslundii		100.0	0.36	ESL 12			
63765100	**Good identification**						**63765100**
Actinomyces israelii		100.0	0.38	FRU 99			
63765102	**Good identification**						**63765102**
Actinomyces israelii		100.0	0.41	FRU 99			
63765500	**Low discrimination**						**63765500**
					GRA	SPO	
Actinomyces israelii		57.23	0.37	FRU 99	99	1	
Mitsuokella multacida		37.66	0.21		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium butyricum		5.11	-.03		99 99	
63765502	**Low discrimination**					**63765502**
					GRA	
Actinomyces israelii		89.71	0.40	FRU 99	99	
Bifidobacterium adolescentis		5.92	0.13		99	
Mitsuokella multacida		4.37	0.04		1	
63767000	**Good identification**					**63767000**
Actinomyces israelii		100.0	0.41	FRU 99		
63767002	**Good identification**					**63767002**
Actinomyces israelii		100.0	0.44	FRU 99		
63767042	**Good identification**					**63767042**
Actinomyces israelii		100.0	0.37	FRU 99		
63767100	**Good identification**					**63767100**
Actinomyces israelii		100.0	0.51	FRU 99		
63767102	**Good identification**					**63767102**
Actinomyces israelii		100.0	0.54	FRU 99		
63767110	**Good identification**					**63767110**
Actinomyces israelii		100.0	0.37	FRU 99		
63767112	**Good identification**					**63767112**
Actinomyces israelii		100.0	0.40	FRU 99		
63767140	**Good identification**					**63767140**
Actinomyces israelii		100.0	0.44	FRU 99		
63767142	**Good identification**					**63767142**
Actinomyces israelii		100.0	0.47	FRU 99		
63767400	**Good identification**					**63767400**
Actinomyces israelii		100.0	0.39	FRU 99		
63767402	**Good identification**					**63767402**
Actinomyces israelii		100.0	0.42	FRU 99		
63767442	**Good identification**					**63767442**
Actinomyces israelii		100.0	0.35	FRU 99		
63767500	**Low discrimination**					**63767500**
					GRA	
Actinomyces israelii		78.68	0.49	FRU 99	99	
Mitsuokella multacida		21.32	0.27		1	
63767502	**Good identification**					**63767502**
					GRA	
Actinomyces israelii		97.13	0.52	FRU 99	99	
Mitsuokella multacida		1.95	0.11		1	
Bifidobacterium adolescentis		0.93	0.12		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63767510	**Low discrimination**					**63767510**
					GRA	
Actinomyces israelii		62.33	0.35	FRU 99	99	
Mitsuokella multacida		37.67	0.19	MLZ 13	1	
63767512	**Low discrimination**					**63767512**
Actinomyces israelii		84.46	0.38	FRU 99		
Bifidobacterium adolescentis		6.59	0.13	MLZ 13		
Bifidobacterium dentium		5.18	-.07			
63767540	**Low discrimination**					**63767540**
					GRA	
Actinomyces israelii		58.94	0.42	FRU 99	99	
Mitsuokella multacida		41.06	0.27		1	
63767542	**Good identification**					**63767542**
					GRA	
Actinomyces israelii		95.09	0.45	FRU 99	99	
Mitsuokella multacida		4.91	0.11		1	
63770100	**Low discrimination**					**63770100**
					GRA	
Actinomyces naeslundii		49.21	0.39	ESL 12	99	
Fusobacterium mortiferum		37.52	0.25	MNS 88	1	
Actinomyces israelii		13.27	0.20	GAL 96	99	
63770101	**Good identification**					**63770101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63770102	**Low discrimination**					**63770102**
Actinomyces naeslundii		61.68	0.39	ESL 12		
Actinomyces israelii		24.94	0.23	MNS 88		
Actinomyces viscosus		13.38	0.03	GAL 96		
63770103	**Good identification**					**63770103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63770120	**Good identification**					**63770120**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63770121	**Good identification**					**63770121**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63770122	**Good identification**					**63770122**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63770123	**Good identification**					**63770123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
63771000	**Low discrimination**					**63771000**
					GRA	
Actinomyces naeslundii		54.07	0.55	ESL 12	99	
Actinomyces israelii		42.79	0.44	MNS 88	99	
Fusobacterium mortiferum		3.14	0.22		1	
63771001	**Very good identification**					**63771001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.58	ESL 12		
63771002	**Low discrimination**					**63771002**
Actinomyces israelii		54.27	0.47	XYL 92		
Actinomyces naeslundii		45.73	0.55			
63771003	**Very good identification**					**63771003**
Actinomyces naeslundii		100.0	0.58	ESL 12		
63771011	**Good identification**					**63771011**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63771013	**Good identification**					**63771013**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63771020	**Good identification**					**63771020**
Actinomyces naeslundii		95.19	0.55	ESL 12		
Actinomyces israelii		4.81	0.23	MNS 88		
63771021	**Very good identification**					**63771021**
Actinomyces naeslundii		100.0	0.58	ESL 12		
63771022	**Acceptable identification**					**63771022**
Actinomyces naeslundii		92.96	0.55	ESL 12		
Actinomyces israelii		7.04	0.26	MNS 88		
63771023	**Very good identification**					**63771023**
Actinomyces naeslundii		100.0	0.58	ESL 12		
63771031	**Good identification**					**63771031**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63771033	**Good identification**					**63771033**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63771040	**Low discrimination**					**63771040**
Actinomyces israelii		63.55	0.37	XYL 92		
Actinomyces naeslundii		36.45	0.42			
63771041	**Good identification**					**63771041**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63771042	**Low discrimination**					**63771042**
Actinomyces israelii		72.34	0.40	XYL 92		
Actinomyces naeslundii		27.66	0.42			
63771043	**Good identification**					**63771043**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63771060	**Low discrimination**					**63771060**
Actinomyces naeslundii		89.98	0.42	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		10.02	0.16	MNS 88		
63771061	**Good identification**					**63771061**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63771062	**Low discrimination**					**63771062**
Actinomyces naeslundii		85.69	0.42	ESL 12		
Actinomyces israelii		14.31	0.19	MNS 88		
63771063	**Good identification**					**63771063**
Actinomyces naeslundii		100.0	0.45	ESL 12		
63771100	**Low discrimination**					**63771100**
					GRA	
Actinomyces israelii		51.49	0.54	XYL 92	99	
Actinomyces naeslundii		46.30	0.62		99	
Fusobacterium mortiferum		2.21	0.28		1	
63771101	**Very good identification**					**63771101**
Actinomyces naeslundii		99.26	0.65	ESL 12		
Actinomyces israelii		0.74	0.20	MNS 88		
63771102	**Low discrimination**					**63771102**
					GRA	
Actinomyces israelii		58.77	0.57	XYL 92	99	
Actinomyces naeslundii		35.23	0.62		99	
Leptotrichia buccalis		2.08	0.07		1	
63771103	**Very good identification**					**63771103**
Actinomyces naeslundii		98.89	0.65	ESL 12		
Actinomyces israelii		1.11	0.23	MNS 88		
63771110	**Low discrimination**					**63771110**
					GRA	
Actinomyces israelii		59.84	0.40	XYL 92	99	
Leptotrichia buccalis		21.21	0.07	MLZ 13	1	
Actinomyces naeslundii		18.95	0.41		99	
63771111	**Good identification**					**63771111**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63771112	**Acceptable identification**					**63771112**
					GRA	
Leptotrichia buccalis		94.00	0.41	NIT 10	1	
Actinomyces israelii		4.02	0.43	MNS 99	99	
Actinomyces naeslundii		0.85	0.41		99	
63771113	**Low discrimination**					**63771113**
					GRA	
Actinomyces naeslundii		57.27	0.44	ESL 12	99	
Leptotrichia buccalis		42.73	0.07	MNS 88	1	
63771120	**Low discrimination**					**63771120**
					SPO	
Actinomyces naeslundii		89.06	0.62	ESL 12	1	
Actinomyces israelii		6.32	0.34	MNS 88	1	
Clostridium paraputrificum		2.09	0.03		99	
63771121	**Very good identification**					**63771121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.65	ESL 12		
63771122	**Low discrimination**					**63771122**
Actinomyces naeslundii		74.45	0.62	ESL 12		
Actinomyces israelii		7.93	0.37	MNS 88		
Bifidobacterium breve		5.61	0.29			
63771123	**Very good identification**					**63771123**
Actinomyces naeslundii		100.0	0.65	ESL 12		
63771130	**Low discrimination**					**63771130**
Actinomyces naeslundii		68.14	0.41	ESL 12		
Actinomyces israelii		13.73	0.19	MNS 88		
Bifidobacterium breve		9.66	0.12	MLZ 5		
63771131	**Good identification**					**63771131**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63771133	**Acceptable identification**					**63771133**
					GRA	
Actinomyces naeslundii		92.35	0.44	ESL 12	99	
Leptotrichia buccalis		7.66	-.09	MNS 88	1	
63771140	**Low discrimination**					**63771140**
Actinomyces israelii		71.02	0.47	XYL 92		
Actinomyces naeslundii		28.98	0.50			
63771141	**Good identification**					**63771141**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63771142	**Low discrimination**					**63771142**
Actinomyces israelii		74.77	0.50	XYL 92		
Actinomyces naeslundii		20.34	0.50			
Bifidobacterium breve		4.89	0.24			
63771143	**Good identification**					**63771143**
Actinomyces naeslundii		97.59	0.53	ESL 12		
Actinomyces israelii		2.42	0.16	MNS 88		
63771152	**Low discrimination**					**63771152**
					GRA	
Actinomyces israelii		46.70	0.36	XYL 92	99	
Leptotrichia buccalis		28.39	0.07	MLZ 13	1	
Bifidobacterium breve		20.44	0.24		99	
63771160	**Low discrimination**					**63771160**
Actinomyces naeslundii		83.43	0.50	ESL 12		
Actinomyces israelii		13.05	0.27	MNS 88		
Bifidobacterium breve		3.53	0.12			
63771161	**Good identification**					**63771161**
Actinomyces naeslundii		100.0	0.53	ESL 12		
63771162	**Low discrimination**					**63771162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		60.17	0.50	ESL 12			
Bifidobacterium breve		25.71	0.29	MNS 88			
Actinomyces israelii		14.12	0.30				
63771163	**Good identification**						**63771163**
Actinomyces naeslundii		100.0	0.53	ESL 12			
63771302	**Acceptable identification**						**63771302**
Actinomyces israelii		92.56	0.38	XYL 92			
Actinomyces naeslundii		7.45	0.28	RHA 7			
63771400	**Good identification**						**63771400**
Actinomyces israelii		100.0	0.42	XYL 92			
63771402	**Good identification**						**63771402**
Actinomyces israelii		100.0	0.45	XYL 92			
63771440	**Good identification**						**63771440**
Actinomyces israelii		100.0	0.35	XYL 92			
63771442	**Good identification**						**63771442**
Actinomyces israelii		100.0	0.38	XYL 92			
63771500	**Low discrimination**						**63771500**
					GRA	SPO	
Actinomyces israelii		89.34	0.53	XYL 92	99	1	
Mitsuokella multacida		6.83	0.21		1	1	
Clostridium butyricum		2.84	0.06		99	99	
63771502	**Low discrimination**						**63771502**
					SPO		
Actinomyces israelii		88.13	0.56	XYL 92	1		
Bifidobacterium adolescentis		10.00	0.33		1		
Clostridium butyricum		0.73	-.01		99		
63771510	**Low discrimination**						**63771510**
					GRA		
Actinomyces israelii		85.43	0.38	XYL 92	99		
Mitsuokella multacida		14.57	0.13	MLZ 13	1		
63771512	**Low discrimination**						**63771512**
					GRA		
Actinomyces israelii		45.11	0.41	XYL 92	99		
Bifidobacterium adolescentis		41.86	0.35	MLZ 13	99		
Leptotrichia buccalis		13.03	0.07		1		
63771522	**Low discrimination**						**63771522**
Actinomyces israelii		89.61	0.35	XYL 92			
Actinomyces naeslundii		10.39	0.28	NAG 6			
63771540	**Low discrimination**						**63771540**
					GRA		
Actinomyces israelii		83.57	0.46	XYL 92	99		
Mitsuokella multacida		16.43	0.21		1		
63771542	**Good identification**						**63771542**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		95.18	0.49	XYL 92	99	
Bifidobacterium adolescentis		3.43	0.18		99	
Mitsuokella multacida		1.39	0.04		1	
63771702	**Low discrimination**					**63771702**
					GRA	
Actinomyces israelii		87.74	0.36	XYL 92	99	
Mitsuokella multacida		12.26	0.09	RHA 7	1	
63772000	**Low discrimination**					**63772000**
Actinomyces naeslundii		64.11	0.36	ESL 12		
Actinomyces israelii		35.90	0.22	MNS 88		
63772001	**Good identification**					**63772001**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63772002	**Low discrimination**					**63772002**
Actinomyces naeslundii		54.35	0.36	ESL 12		
Actinomyces israelii		45.65	0.26	MNS 88		
63772003	**Good identification**					**63772003**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63772020	**Good identification**					**63772020**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63772021	**Good identification**					**63772021**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63772022	**Good identification**					**63772022**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63772023	**Good identification**					**63772023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
63772100	**Low discrimination**					**63772100**
Actinomyces naeslundii		55.96	0.44	ESL 12		
Actinomyces israelii		44.04	0.33	MNS 88		
63772101	**Good identification**					**63772101**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63772102	**Low discrimination**					**63772102**
Actinomyces israelii		54.14	0.36	GAL 99		
Actinomyces naeslundii		45.86	0.44	XYL 92		
63772103	**Good identification**					**63772103**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63772120	**Good identification**					**63772120**
Actinomyces naeslundii		100.0	0.44	ESL 12		
63772121	**Good identification**					**63772121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	ESL 12		
63772122	**Acceptable identification**					**63772122**
Actinomyces naeslundii		93.00	0.44	ESL 12		
Actinomyces israelii		7.01	0.15	MNS 88		
63772123	**Good identification**					**63772123**
Actinomyces naeslundii		100.0	0.47	ESL 12		
63773000	**Low discrimination**					**63773000**
Actinomyces israelii		69.79	0.57	XYL 92		
Actinomyces naeslundii		30.21	0.60			
63773001	**Very good identification**					**63773001**
Actinomyces naeslundii		98.47	0.63	ESL 12		
Actinomyces israelii		1.53	0.22	MNS 88		
63773002	**Low discrimination**					**63773002**
Actinomyces israelii		77.60	0.60	XYL 92		
Actinomyces naeslundii		22.40	0.60			
63773003	**Very good identification**					**63773003**
Actinomyces naeslundii		97.72	0.63	ESL 12		
Actinomyces israelii		2.28	0.26	MNS 88		
63773010	**Low discrimination**					**63773010**
Actinomyces israelii		86.77	0.43	XYL 92		
Actinomyces naeslundii		13.23	0.38	MLZ 13		
63773011	**Good identification**					**63773011**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63773012	**Acceptable identification**					**63773012**
Actinomyces israelii		90.77	0.46	XYL 92		
Actinomyces naeslundii		9.23	0.38	MLZ 13		
63773013	**Good identification**					**63773013**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63773020	**Low discrimination**					**63773020**
Actinomyces naeslundii		87.15	0.60	ESL 12		
Actinomyces israelii		12.85	0.36	MNS 88		
63773021	**Very good identification**					**63773021**
Actinomyces naeslundii		100.0	0.63	ESL 12		
63773022	**Low discrimination**					**63773022**
Actinomyces naeslundii		81.89	0.60	ESL 12		
Actinomyces israelii		18.11	0.39	MNS 88		
63773023	**Very good identification**					**63773023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.63	ESL 12		
63773030	**Low discrimination**					**63773030**
Actinomyces naeslundii		70.49	0.38	ESL 12		
Actinomyces israelii		29.51	0.22	MNS 88		
63773031	**Good identification**					**63773031**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63773032	**Low discrimination**					**63773032**
Actinomyces naeslundii		61.43	0.38	ESL 12		
Actinomyces israelii		38.57	0.25	MNS 88		
63773033	**Good identification**					**63773033**
Actinomyces naeslundii		100.0	0.41	ESL 12		
63773040	**Low discrimination**					**63773040**
Actinomyces israelii		83.58	0.50	XYL 92		
Actinomyces naeslundii		16.42	0.47			
63773041	**Good identification**					**63773041**
Actinomyces naeslundii		96.69	0.50	ESL 12		
Actinomyces israelii		3.31	0.15	MNS 88		
63773042	**Low discrimination**					**63773042**
Actinomyces israelii		88.42	0.53	XYL 92		
Actinomyces naeslundii		11.58	0.47			
63773043	**Good identification**					**63773043**
Actinomyces naeslundii		95.11	0.50	ESL 12		
Actinomyces israelii		4.89	0.18	MNS 88		
63773050	**Acceptable identification**					**63773050**
Actinomyces israelii		93.53	0.36	XYL 92		
Actinomyces naeslundii		6.47	0.25	MLZ 13		
63773052	**Good identification**					**63773052**
Actinomyces israelii		95.59	0.39	XYL 92		
Actinomyces naeslundii		4.41	0.25	MLZ 13		
63773060	**Low discrimination**					**63773060**
Actinomyces naeslundii		75.48	0.47	ESL 12		
Actinomyces israelii		24.52	0.29	MNS 88		
63773061	**Good identification**					**63773061**
Actinomyces naeslundii		100.0	0.50	ESL 12		
63773062	**Low discrimination**					**63773062**
Actinomyces naeslundii		67.23	0.47	ESL 12		
Actinomyces israelii		32.77	0.32	MNS 88		
63773063	**Good identification**					**63773063**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	ESL 12		
63773100	**Low discrimination**					**63773100**
Actinomyces israelii		76.45	0.67	XYL 92		
Actinomyces naeslundii		23.55	0.67			
63773101	**Very good identification**					**63773101**
Actinomyces naeslundii		97.86	0.70	ESL 12		
Actinomyces israelii		2.14	0.33	MNS 88		
63773102	**Low discrimination**					**63773102**
Actinomyces israelii		82.52	0.70	XYL 92		
Actinomyces naeslundii		16.95	0.67			
Bifidobacterium breve		0.37	0.24			
63773103	**Very good identification**					**63773103**
Actinomyces naeslundii		96.83	0.70	ESL 12		
Actinomyces israelii		3.18	0.36	MNS 88		
63773110	**Acceptable identification**					**63773110**
Actinomyces israelii		90.21	0.53	XYL 92		
Actinomyces naeslundii		9.79	0.46	MLZ 13		
63773111	**Acceptable identification**					**63773111**
Actinomyces naeslundii		94.16	0.49	ESL 12		
Actinomyces israelii		5.85	0.19	MNS 88		
63773112	**Low discrimination**					**63773112**
					GRA	
Actinomyces israelii		86.17	0.56	XYL 92	99	
Actinomyces naeslundii		6.23	0.46	MLZ 13	99	
Leptotrichia buccalis		3.59	0.07		1	
63773113	**Acceptable identification**					**63773113**
Actinomyces naeslundii		91.48	0.49	ESL 12		
Actinomyces israelii		8.52	0.22	MNS 88		
63773120	**Low discrimination**					**63773120**
					SPO	
Actinomyces naeslundii		79.52	0.67	ESL 12	1	
Actinomyces israelii		16.48	0.47	MNS 88	1	
Clostridium ramosum		3.70	0.19		92	
63773121	**Very good identification**					**63773121**
Actinomyces naeslundii		100.0	0.70	ESL 12		
63773122	**Low discrimination**					**63773122**
					SPO	
Actinomyces naeslundii		65.11	0.67	ESL 12	1	
Actinomyces israelii		20.24	0.50	MNS 88	1	
Clostridium ramosum		12.13	0.30		92	
63773123	**Very good identification**					**63773123**
Actinomyces naeslundii		99.79	0.70	ESL 12		
Actinomyces israelii		0.21	0.15	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63773130	**Low discrimination**					**63773130**
Actinomyces naeslundii		60.19	0.46	ESL 12		
Actinomyces israelii		35.41	0.32	MNS 88		
Bifidobacterium breve		4.39	0.12	MLZ 5		
63773131	**Good identification**					**63773131**
Actinomyces naeslundii		100.0	0.49	ESL 12		
63773132	**Low discrimination**					**63773132**
Actinomyces naeslundii		37.25	0.46	ESL 12		
Actinomyces israelii		32.87	0.35	MNS 88		
Bifidobacterium breve		27.49	0.29	MLZ 5		
63773133	**Good identification**					**63773133**
Actinomyces naeslundii		100.0	0.49	ESL 12		
63773140	**Low discrimination**					**63773140**
Actinomyces israelii		87.74	0.60	XYL 92		
Actinomyces naeslundii		12.26	0.55			
63773141	**Very good identification**					**63773141**
Actinomyces naeslundii		95.40	0.58	ESL 12		
Actinomyces israelii		4.60	0.26	MNS 88		
63773142	**Good identification**					**63773142**
Actinomyces israelii		90.52	0.63	XYL 92		
Actinomyces naeslundii		8.44	0.55			
Bifidobacterium breve		1.04	0.24			
63773143	**Good identification**					**63773143**
Actinomyces naeslundii		93.26	0.58	ESL 12		
Actinomyces israelii		6.74	0.29	MNS 88		
63773150	**Good identification**					**63773150**
Actinomyces israelii		95.31	0.46	XYL 92		
Actinomyces naeslundii		4.69	0.33	MLZ 13		
63773151	**Good identification**					**63773151**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63773152	**Acceptable identification**					**63773152**
Actinomyces israelii		90.09	0.49	XYL 92		
Bifidobacterium breve		6.96	0.24	MLZ 13		
Actinomyces naeslundii		2.96	0.33			
63773153	**Low discrimination**					**63773153**
Actinomyces naeslundii		82.98	0.36	ESL 12		
Actinomyces israelii		17.02	0.15	MNS 88		
63773160	**Low discrimination**					**63773160**
Actinomyces naeslundii		67.64	0.55	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		30.89	0.39	MNS 88		
Bifidobacterium breve		1.47	0.12			
63773161	**Very good identification**					**63773161**
Actinomyces naeslundii		100.0	0.58	ESL 12		
63773162	**Low discrimination**					**63773162**
Actinomyces naeslundii		52.49	0.55	ESL 12		
Actinomyces israelii		35.95	0.42	MNS 88		
Bifidobacterium breve		11.55	0.29			
63773163	**Very good identification**					**63773163**
Actinomyces naeslundii		100.0	0.58	ESL 12		
63773171	**Good identification**					**63773171**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63773173	**Good identification**					**63773173**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63773200	**Acceptable identification**					**63773200**
Actinomyces israelii		94.51	0.37	XYL 92		
Actinomyces naeslundii		5.49	0.25	RHA 7		
63773202	**Good identification**					**63773202**
Actinomyces israelii		96.27	0.40	XYL 92		
Actinomyces naeslundii		3.73	0.25	RHA 7		
63773300	**Good identification**					**63773300**
Actinomyces israelii		96.03	0.48	XYL 92		
Actinomyces naeslundii		3.97	0.33	RHA 7		
63773301	**Low discrimination**					**63773301**
Actinomyces naeslundii		85.99	0.36	ESL 12		
Actinomyces israelii		14.01	0.14	MNS 88		
63773302	**Good identification**					**63773302**
					SPO	
Actinomyces israelii		96.68	0.51	XYL 92	1	
Actinomyces naeslundii		2.67	0.33	RHA 7	1	
Clostridium ramosum		0.66	-.03		92	
63773303	**Low discrimination**					**63773303**
Actinomyces naeslundii		80.36	0.36	ESL 12		
Actinomyces israelii		19.64	0.17	MNS 88		
63773312	**Good identification**					**63773312**
Actinomyces israelii		100.0	0.37	XYL 92		
63773321	**Good identification**					**63773321**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63773323	**Low discrimination**					**63773323**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		85.87	0.36	ESL 12	1	
Clostridium ramosum		14.13	-.03	MNS 88	92	
63773340	**Good identification**					**63773340**
Actinomyces israelii		100.0	0.41	XYL 92		
63773342	**Good identification**					**63773342**
Actinomyces israelii		100.0	0.44	XYL 92		
63773400	**Good identification**					**63773400**
Actinomyces israelii		99.47	0.55	XYL 92		
Actinomyces naeslundii		0.53	0.25			
63773402	**Very good identification**					**63773402**
Actinomyces israelii		99.65	0.58	XYL 92		
Actinomyces naeslundii		0.36	0.25			
63773410	**Good identification**					**63773410**
Actinomyces israelii		100.0	0.41	XYL 92		
63773412	**Good identification**					**63773412**
Actinomyces israelii		100.0	0.44	XYL 92		
63773422	**Acceptable identification**					**63773422**
Actinomyces israelii		94.71	0.38	XYL 92		
Actinomyces naeslundii		5.29	0.25	NAG 6		
63773440	**Good identification**					**63773440**
Actinomyces israelii		100.0	0.48	XYL 92		
63773442	**Good identification**					**63773442**
Actinomyces israelii		100.0	0.51	XYL 92		
63773452	**Good identification**					**63773452**
Actinomyces israelii		100.0	0.37	XYL 92		
63773500	**Very good identification**					**63773500**
					GRA	
Actinomyces israelii		96.59	0.65	XYL 92	99	
Mitsuokella multacida		3.04	0.27		1	
Actinomyces naeslundii		0.37	0.33		99	
63773502	**Very good identification**					**63773502**
Actinomyces israelii		97.92	0.68	XYL 92		
Bifidobacterium adolescentis		1.60	0.32			
Actinomyces naeslundii		0.25	0.33			
63773510	**Acceptable identification**					**63773510**
					GRA	
Actinomyces israelii		93.44	0.51	XYL 92	99	
Mitsuokella multacida		6.56	0.19	MLZ 13	1	
63773512	**Low discrimination**					**63773512**
Actinomyces israelii		87.24	0.54	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		11.69	0.33	MLZ 13		
Bifidobacterium dentium		0.62	-.07			
63773520	**Low discrimination**					**63773520**
					GRA	
Actinomyces israelii		84.54	0.45	XYL 92	99	
Mitsuokella multacida		10.42	0.17	NAG 6	1	
Actinomyces naeslundii		5.04	0.33		99	
63773521	**Good identification**					**63773521**
Actinomyces naeslundii		100.0	0.36	ESL 12		
63773522	**Good identification**					**63773522**
Actinomyces israelii		96.18	0.48	XYL 92		
Actinomyces naeslundii		3.82	0.33	NAG 6		
63773523	**Low discrimination**					**63773523**
Actinomyces naeslundii		85.51	0.36	ESL 12		
Actinomyces israelii		14.49	0.14	MNS 88		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63773540	**Good identification**						**63773540**
					GRA		
Actinomyces israelii		92.51	0.58	XYL 92	99		
Mitsuokella multacida		7.49	0.27		1		
63773542	**Very good identification**						**63773542**
					GRA		
Actinomyces israelii		98.89	0.61	XYL 92	99		
Mitsuokella multacida		0.59	0.11		1		
Bifidobacterium adolescentis		0.52	0.16		99		
63773550	**Low discrimination**						**63773550**
					COC	GRA	
Actinomyces israelii		82.81	0.44	XYL 92	1	99	
Mitsuokella multacida		14.95	0.19	MLZ 13	1	1	
Blautia producta		2.24	-.03		99	99	
63773552	**Acceptable identification**						**63773552**
					GRA		
Actinomyces israelii		94.70	0.47	XYL 92	99		
Bifidobacterium adolescentis		4.03	0.18	MLZ 13	99		
Mitsuokella multacida		1.27	0.03		1		
63773560	**Low discrimination**						**63773560**
					GRA		
Actinomyces israelii		75.93	0.38	XYL 92	99		
Mitsuokella multacida		24.07	0.17	NAG 6	1		
63773562	**Good identification**						**63773562**
Actinomyces israelii		100.0	0.41	XYL 92			
63773600	**Good identification**						**63773600**
Actinomyces israelii		100.0	0.36	XYL 92			
63773602	**Good identification**						**63773602**
Actinomyces israelii		100.0	0.39	XYL 92			
63773700	**Low discrimination**						**63773700**
					GRA		
Actinomyces israelii		56.29	0.46	XYL 92	99		
Mitsuokella multacida		43.71	0.32	RHA 7	1		
63773702	**Acceptable identification**						**63773702**
					GRA		
Actinomyces israelii		94.56	0.49	XYL 92	99		
Mitsuokella multacida		5.44	0.15	RHA 7	1		
63773712	**Low discrimination**						**63773712**
					GRA		
Actinomyces israelii		88.63	0.35	XYL 92	99		
Mitsuokella multacida		11.37	0.07	RHA 7	1		
63773742	**Low discrimination**						**63773742**
					GRA		
Actinomyces israelii		87.11	0.42	XYL 92	99		
Mitsuokella multacida		12.89	0.15	RHA 7	1		
63774100	**Good identification**						**63774100**
Actinomyces israelii		100.0	0.38	GAL 99			
63774102	**Good identification**						**63774102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.41	GAL 99			
63774500	**Low discrimination**						**63774500**
					GRA	SPO	
Actinomyces israelii		57.23	0.37	GAL 99	99	1	
Mitsuokella multacida		37.66	0.21		1	1	
Clostridium butyricum		5.11	-.03		99	99	
63774502	**Low discrimination**						**63774502**
					GRA		
Actinomyces israelii		89.71	0.40	GAL 99	99		
Bifidobacterium adolescentis		5.92	0.13		99		
Mitsuokella multacida		4.37	0.04		1		
63775000	**Very good identification**						**63775000**
Actinomyces israelii		100.0	0.62				
63775002	**Very good identification**						**63775002**
Actinomyces israelii		100.0	0.65				
63775010	**Good identification**						**63775010**
Actinomyces israelii		100.0	0.48	MLZ 13			
63775012	**Good identification**						**63775012**
Actinomyces israelii		100.0	0.51	MLZ 13			
63775020	**Good identification**						**63775020**
Actinomyces israelii		100.0	0.41	NAG 6			
63775022	**Good identification**						**63775022**
Actinomyces israelii		100.0	0.44	NAG 6			
63775040	**Good identification**						**63775040**
Actinomyces israelii		100.0	0.55				
63775042	**Very good identification**						**63775042**
Actinomyces israelii		100.0	0.58				
63775050	**Good identification**						**63775050**
Actinomyces israelii		100.0	0.41	MLZ 13			
63775052	**Good identification**						**63775052**
Actinomyces israelii		100.0	0.44	MLZ 13			
63775062	**Good identification**						**63775062**
Actinomyces israelii		100.0	0.37	NAG 6			
63775100	**Very good identification**						**63775100**
					GRA	SPO	
Actinomyces israelii		98.59	0.72		99	1	
Clostridium butyricum		0.80	0.15		99	99	
Mitsuokella multacida		0.54	0.21		1	1	
63775101	**Low discrimination**						**63775101**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		84.95	0.38	URE 1	99		
Actinomyces naeslundii		9.96	0.31		99		
Mitsuokella multacida		5.08	0.04		1		
63775102	**Very good identification**						**63775102**
					SPO		
Actinomyces israelii		98.90	0.75		1		
Bifidobacterium adolescentis		0.80	0.33		1		
Clostridium butyricum		0.21	0.08		99		
63775103	**Acceptable identification**						**63775103**
Actinomyces israelii		92.75	0.41	URE 1			
Actinomyces naeslundii		7.25	0.31				
63775110	**Very good identification**						**63775110**
					GRA	SPO	
Actinomyces israelii		98.52	0.58	MLZ 13	99	1	
Mitsuokella multacida		1.20	0.13		1	1	
Clostridium butyricum		0.28	-.07		99	99	
63775112	**Good identification**						**63775112**
					GRA		
Actinomyces israelii		92.03	0.61	MLZ 13	99		
Bifidobacterium adolescentis		6.08	0.35		99		
Leptotrichia buccalis		1.89	0.07		1		
63775120	**Good identification**						**63775120**
					GRA		
Actinomyces israelii		95.72	0.52	NAG 6	99		
Mitsuokella multacida		2.04	0.11		1		
Actinomyces naeslundii		1.18	0.28		99		
63775122	**Good identification**						**63775122**
					GRA		
Actinomyces israelii		96.58	0.55	NAG 6	99		
Parabacteroides distasonis		2.62	0.24		1		
Actinomyces naeslundii		0.80	0.28		99		
63775130	**Good identification**						**63775130**
					GRA		
Actinomyces israelii		95.47	0.38	MLZ 13	99		
Mitsuokella multacida		4.54	0.02	NAG 6	1		
63775132	**Low discrimination**						**63775132**
					GRA		
Actinomyces israelii		88.22	0.41	MLZ 13	99		
Parabacteroides distasonis		8.62	0.20	NAG 6	1		
Leptotrichia buccalis		3.16	-.09		1		
63775140	**Very good identification**						**63775140**
					GRA	SPO	
Actinomyces israelii		98.45	0.65		99	1	
Mitsuokella multacida		1.38	0.21		1	1	
Clostridium butyricum		0.18	-.03		99	99	
63775142	**Very good identification**						**63775142**
					GRA		
Actinomyces israelii		99.64	0.68		99		
Bifidobacterium adolescentis		0.26	0.18		99		
Mitsuokella multacida		0.10	0.04		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63775150	**Good identification**					**63775150**
					GRA	
Actinomyces israelii		96.98	0.51	MLZ 13	99	
Mitsuokella multacida		3.03	0.13		1	
63775152	**Good identification**					**63775152**
Actinomyces israelii		97.95	0.54	MLZ 13		
Bifidobacterium adolescentis		2.06	0.19			
63775160	**Acceptable identification**					**63775160**
					GRA	
Actinomyces israelii		94.81	0.45	NAG 6	99	
Mitsuokella multacida		5.19	0.11		1	
63775162	**Good identification**					**63775162**
Actinomyces israelii		100.0	0.48	NAG 6		
63775200	**Good identification**					**63775200**
Actinomyces israelii		100.0	0.43	RHA 7		
63775202	**Good identification**					**63775202**
Actinomyces israelii		100.0	0.46	RHA 7		
63775240	**Good identification**					**63775240**
Actinomyces israelii		100.0	0.36	RHA 7		
63775242	**Good identification**					**63775242**
Actinomyces israelii		100.0	0.39	RHA 7		
63775300	**Low discrimination**					**63775300**
					GRA	
Actinomyces israelii		88.17	0.53	RHA 7	99	
Mitsuokella multacida		11.83	0.25		1	
63775302	**Good identification**					**63775302**
					GRA	
Actinomyces israelii		99.02	0.56	RHA 7	99	
Mitsuokella multacida		0.98	0.09		1	
63775310	**Low discrimination**					**63775310**
					GRA	
Actinomyces israelii		76.96	0.39	RHA 7	99	
Mitsuokella multacida		23.04	0.17	MLZ 13	1	
63775312	**Acceptable identification**					**63775312**
					GRA	
Actinomyces israelii		92.58	0.42	RHA 7	99	
Clostridium clostridioforme		5.37	0.11	MLZ 13	50	
Mitsuokella multacida		2.05	0.01		1	
63775340	**Low discrimination**					**63775340**
					GRA	
Actinomyces israelii		74.35	0.46	RHA 7	99	
Mitsuokella multacida		25.65	0.25		1	
63775342	**Good identification**					**63775342**
					GRA	
Actinomyces israelii		97.51	0.49	RHA 7	99	
Mitsuokella multacida		2.49	0.09		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63775400	**Very good identification**						**63775400**
					GRA	SPO	
Actinomyces israelii		96.02	0.60		99	1	
Mitsuokella multacida		2.55	0.21		1	1	
Clostridium butyricum		1.43	0.08		99	99	
63775402	**Very good identification**						**63775402**
					SPO		
Actinomyces israelii		99.16	0.63		1		
Clostridium butyricum		0.38	0.01		99		
Bifidobacterium adolescentis		0.27	0.13		1		
63775410	**Acceptable identification**						**63775410**
					GRA		
Actinomyces israelii		94.40	0.46	MLZ 13	99		
Mitsuokella multacida		5.60	0.13		1		
63775412	**Good identification**						**63775412**
Actinomyces israelii		96.58	0.49	MLZ 13			
Bifidobacterium adolescentis		2.11	0.15				
Clostridium clostridioforme		1.32	0.08				
63775420	**Acceptable identification**						**63775420**
					GRA		
Actinomyces israelii		90.57	0.40	NAG 6	99		
Mitsuokella multacida		9.43	0.11		1		
63775422	**Good identification**						**63775422**
Actinomyces israelii		100.0	0.43	NAG 6			
63775440	**Acceptable identification**						**63775440**
					GRA		
Actinomyces israelii		93.60	0.53		99		
Mitsuokella multacida		6.40	0.21		1		
63775442	**Very good identification**						**63775442**
					GRA		
Actinomyces israelii		99.50	0.56		99		
Mitsuokella multacida		0.50	0.04		1		
63775450	**Low discrimination**						**63775450**
					GRA		
Actinomyces israelii		86.77	0.39	MLZ 13	99		
Mitsuokella multacida		13.23	0.13		1		
63775452	**Good identification**						**63775452**
Actinomyces israelii		100.0	0.42	MLZ 13			
63775462	**Good identification**						**63775462**
Actinomyces israelii		100.0	0.36	NAG 6			
63775500	**Low discrimination**						**63775500**
					GRA	SPO	
Actinomyces israelii		57.20	0.71		99	1	
Mitsuokella multacida		37.64	0.55		1	1	
Clostridium butyricum		5.11	0.32		99	99	
63775501	**Low discrimination**						**63775501**
					GRA	SPO	
Mitsuokella multacida		86.92	0.39	MNS 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		12.01	0.37	URE 10	99	1	
Clostridium butyricum		1.07	-.03		99	99	
63775502	**Low discrimination**						**63775502**
					GRA		
Actinomyces israelii		87.85	0.74		99		
Bifidobacterium adolescentis		5.80	0.47		99		
Mitsuokella multacida		4.28	0.39		1		
63775503	**Low discrimination**						**63775503**
					GRA		
Actinomyces israelii		62.42	0.40	URE 1	99		
Mitsuokella multacida		33.46	0.22		1		
Bifidobacterium adolescentis		4.12	0.13		99		
63775510	**Low discrimination**						**63775510**
					GRA	SPO	
Mitsuokella multacida		58.54	0.47	MNS 99	1	1	
Actinomyces israelii		39.88	0.57		99	1	
Clostridium butyricum		1.26	0.10		99	99	
63775512	**Low discrimination**						**63775512**
					GRA		
Actinomyces israelii		59.82	0.60	MLZ 13	99		
Bifidobacterium adolescentis		32.30	0.49		99		
Mitsuokella multacida		6.50	0.30		1		
63775520	**Low discrimination**						**63775520**
					GRA	SPO	
Mitsuokella multacida		69.68	0.45	MNS 99	1	1	
Actinomyces israelii		27.04	0.50		99	1	
Clostridium butyricum		3.29	0.14		99	99	
63775522	**Low discrimination**						**63775522**
					GRA	SPO	
Actinomyces israelii		79.42	0.53	NAG 6	99	1	
Mitsuokella multacida		15.16	0.28		1	1	
Clostridium butyricum		2.51	0.07		99	99	
63775530	**Low discrimination**						**63775530**
					GRA		
Mitsuokella multacida		85.18	0.37	MNS 99	1		
Actinomyces israelii		14.82	0.36		99		
63775532	**Low discrimination**						**63775532**
					GRA		
Actinomyces israelii		53.74	0.39	MLZ 13	99		
Mitsuokella multacida		22.89	0.20	NAG 6	1		
Clostridium clostridioforme		9.59	0.17		50		
63775540	**Low discrimination**						**63775540**
					GRA	SPO	
Mitsuokella multacida		62.39	0.55	MNS 99	1	1	
Actinomyces israelii		36.87	0.64		99	1	
Clostridium butyricum		0.74	0.14		99	99	
63775541	**Acceptable identification**						**63775541**
					GRA		
Mitsuokella multacida		94.90	0.39	MNS 99	1		
Actinomyces israelii		5.10	0.30	URE 10	99		
63775542	**Low discrimination**						**63775542**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		86.84	0.67		99		
Mitsuokella multacida		10.89	0.39		1		
Bifidobacterium adolescentis		1.82	0.32		99		
63775550	**Low discrimination**						**63775550**
					COC	GRA	
Mitsuokella multacida		78.84	0.47	MNS 99	1	1	
Actinomyces israelii		20.89	0.50		1	99	
Blautia producta		0.28	-.03		99	99	
63775552	**Low discrimination**						**63775552**
					GRA		
Actinomyces israelii		68.91	0.53	MLZ 13	99		
Mitsuokella multacida		19.27	0.30		1		
Bifidobacterium adolescentis		11.83	0.33		99		
63775560	**Low discrimination**						**63775560**
					GRA	SPO	
Mitsuokella multacida		86.58	0.45	MNS 99	1	1	
Actinomyces israelii		13.07	0.43		99	1	
Clostridium butyricum		0.36	-.05		99	99	
63775562	**Low discrimination**						**63775562**
					GRA		
Actinomyces israelii		67.07	0.46	NAG 6	99		
Mitsuokella multacida		32.93	0.28		1		
63775570	**Acceptable identification**						**63775570**
					GRA		
Mitsuokella multacida		93.66	0.37	MNS 99	1		
Actinomyces israelii		6.34	0.29		99		
63775600	**Low discrimination**						**63775600**
					GRA		
Actinomyces israelii		60.39	0.41	RHA 7	99		
Mitsuokella multacida		39.61	0.25		1		
63775602	**Acceptable identification**						**63775602**
					GRA		
Actinomyces israelii		91.57	0.44	RHA 7	99		
Mitsuokella multacida		4.45	0.09		1		
Clostridium clostridioforme		3.98	0.11		50		
63775642	**Low discrimination**						**63775642**
					GRA		
Actinomyces israelii		88.89	0.37	RHA 7	99		
Mitsuokella multacida		11.11	0.09		1		
63775700	**Good identification**						**63775700**
					GRA	SPO	
Mitsuokella multacida		94.00	0.60	MNS 99	1	1	
Actinomyces israelii		5.79	0.51		99	1	
Clostridium butyricum		0.21	0.06		99	99	
63775701	**Good identification**						**63775701**
					GRA		
Mitsuokella multacida		99.44	0.43	MNS 99	1		
Actinomyces israelii		0.56	0.17	URE 10	99		
63775702	**Low discrimination**						**63775702**
					GRA		
Mitsuokella multacida		53.27	0.43	MNS 99	1		
Actinomyces israelii		44.30	0.55	BGL 10	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		1.61	0.20		50		
63775710	**Good identification**						**63775710**
					GRA		
Mitsuokella multacida		97.31	0.51	MNS 99	1		
Actinomyces israelii		2.69	0.37		99		
63775711	**Good identification**						**63775711**
Mitsuokella multacida		100.0	0.35	MNS 99			
63775712	**Low discrimination**						**63775712**
					GRA		
Mitsuokella multacida		55.14	0.35	MNS 99	1		
Clostridium clostridioforme		22.82	0.32	BGL 10	50		
Actinomyces israelii		20.55	0.40		99		
63775720	**Good identification**						**63775720**
					GRA		
Mitsuokella multacida		98.45	0.49	MNS 99	1		
Actinomyces israelii		1.55	0.31		99		
63775730	**Good identification**						**63775730**
					GRA		
Mitsuokella multacida		99.30	0.41	MNS 99	1		
Actinomyces israelii		0.70	0.17		99		
63775740	**Very good identification**						**63775740**
					GRA		
Mitsuokella multacida		97.66	0.60	MNS 99	1		
Actinomyces israelii		2.34	0.44		99		
63775741	**Good identification**						**63775741**
Mitsuokella multacida		100.0	0.43	MNS 99			
63775742	**Low discrimination**						**63775742**
					GRA		
Mitsuokella multacida		75.57	0.43	MNS 99	1		
Actinomyces israelii		24.44	0.47	BGL 10	99		
63775750	**Good identification**						**63775750**
					COC	GRA	
Mitsuokella multacida		98.75	0.51	MNS 99	1	1	
Actinomyces israelii		1.06	0.30		1	99	
Blautia producta		0.19	-.03		99	99	
63775751	**Good identification**						**63775751**
Mitsuokella multacida		100.0	0.35	MNS 99			
63775752	**Low discrimination**						**63775752**
					GRA		
Mitsuokella multacida		87.34	0.35	MNS 99	1		
Actinomyces israelii		12.66	0.33	BGL 10	99		
63775760	**Good identification**						**63775760**
					GRA		
Mitsuokella multacida		99.39	0.49	MNS 99	1		
Actinomyces israelii		0.61	0.24		99		
63775770	**Good identification**						**63775770**
Mitsuokella multacida		100.0	0.41	MNS 99			
63776000	**Good identification**						**63776000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.41	GAL 99		
63776002	**Good identification**					**63776002**
Actinomyces israelii		100.0	0.44	GAL 99		
63776042	**Good identification**					**63776042**
Actinomyces israelii		100.0	0.37	GAL 99		
63776100	**Good identification**					**63776100**
Actinomyces israelii		100.0	0.51	GAL 99		
63776102	**Good identification**					**63776102**
Actinomyces israelii		100.0	0.54	GAL 99		
63776110	**Good identification**					**63776110**
Actinomyces israelii		100.0	0.37	GAL 99		
63776112	**Good identification**					**63776112**
Actinomyces israelii		100.0	0.40	GAL 99		
63776140	**Good identification**					**63776140**
Actinomyces israelii		100.0	0.44	GAL 99		
63776142	**Good identification**					**63776142**
Actinomyces israelii		100.0	0.47	GAL 99		
63776400	**Good identification**					**63776400**
Actinomyces israelii		100.0	0.39	GAL 99		
63776402	**Good identification**					**63776402**
Actinomyces israelii		100.0	0.42	GAL 99		
63776442	**Good identification**					**63776442**
Actinomyces israelii		100.0	0.35	GAL 99		
63776500	**Low discrimination**					**63776500**
					GRA	
Actinomyces israelii		78.68	0.49	GAL 99	99	
Mitsuokella multacida		21.32	0.27		1	
63776502	**Good identification**					**63776502**
					GRA	
Actinomyces israelii		97.13	0.52	GAL 99	99	
Mitsuokella multacida		1.95	0.11		1	
Bifidobacterium adolescentis		0.93	0.12		99	
63776510	**Low discrimination**					**63776510**
					GRA	
Actinomyces israelii		62.33	0.35	GAL 99	99	
Mitsuokella multacida		37.67	0.19	MLZ 13	1	
63776512	**Low discrimination**					**63776512**
Actinomyces israelii		84.46	0.38	GAL 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		6.59	0.13	MLZ 13		
Bifidobacterium dentium		5.18	-.07			
63776540	**Low discrimination**					**63776540**
					GRA	
Actinomyces israelii		58.94	0.42	GAL 99	99	
Mitsuokella multacida		41.06	0.27		1	
63776542	**Good identification**					**63776542**
					GRA	
Actinomyces israelii		95.09	0.45	GAL 99	99	
Mitsuokella multacida		4.91	0.11		1	
63777000	**Very good identification**					**63777000**
Actinomyces israelii		99.96	0.75			
Actinomyces naeslundii		0.04	0.25			
63777001	**Acceptable identification**					**63777001**
Actinomyces israelii		94.66	0.41	URE 1		
Actinomyces naeslundii		5.35	0.28			
63777002	**Very good identification**					**63777002**
Actinomyces israelii		99.98	0.78			
Actinomyces naeslundii		0.03	0.25			
63777003	**Good identification**					**63777003**
Actinomyces israelii		96.37	0.44	URE 1		
Actinomyces naeslundii		3.63	0.28			
63777010	**Very good identification**					**63777010**
Actinomyces israelii		100.0	0.61	MLZ 13		
63777012	**Very good identification**					**63777012**
Actinomyces israelii		100.0	0.64	MLZ 13		
63777020	**Good identification**					**63777020**
Actinomyces israelii		99.41	0.54	NAG 6		
Actinomyces naeslundii		0.59	0.25			
63777022	**Very good identification**					**63777022**
Actinomyces israelii		99.60	0.57	NAG 6		
Actinomyces naeslundii		0.40	0.25			
63777030	**Good identification**					**63777030**
Actinomyces israelii		100.0	0.40	MLZ 13		
63777032	**Good identification**					**63777032**
Actinomyces israelii		100.0	0.43	MLZ 13		
63777040	**Very good identification**					**63777040**
Actinomyces israelii		100.0	0.68			
63777042	**Very good identification**					**63777042**
Actinomyces israelii		100.0	0.71			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
63777043	**Good identification**					**63777043**
Actinomyces israelii		100.0	0.37	URE 1		
63777050	**Good identification**					**63777050**
Actinomyces israelii		100.0	0.54	MLZ 13		
63777052	**Very good identification**					**63777052**
Actinomyces israelii		100.0	0.57	MLZ 13		
63777060	**Good identification**					**63777060**
Actinomyces israelii		100.0	0.47	NAG 6		
63777062	**Good identification**					**63777062**
Actinomyces israelii		100.0	0.50	NAG 6		
63777072	**Good identification**					**63777072**
Actinomyces israelii		100.0	0.36	MLZ 13		
63777100	**Very good identification**					**63777100**
					GRA	
Actinomyces israelii		99.73	0.85		99	
Mitsuokella multacida		0.22	0.27		1	
Actinomyces naeslundii		0.03	0.33		99	
63777101	**Acceptable identification**					**63777101**
					GRA	
Actinomyces israelii		93.91	0.51	URE 1	99	
Actinomyces naeslundii		3.77	0.36		99	
Mitsuokella multacida		2.31	0.11		1	
63777102	**Very good identification**					**63777102**
Actinomyces israelii		99.85	0.88			
Bifidobacterium adolescentis		0.12	0.32			
Actinomyces naeslundii		0.02	0.33			
63777103	**Good identification**					**63777103**
Actinomyces israelii		97.39	0.54	URE 1		
Actinomyces naeslundii		2.61	0.36			
63777110	**Very good identification**					**63777110**
					GRA	
Actinomyces israelii		99.50	0.71	MLZ 13	99	
Mitsuokella multacida		0.50	0.19		1	
63777111	**Acceptable identification**					**63777111**
					GRA	
Actinomyces israelii		94.79	0.37	MLZ 13	99	
Mitsuokella multacida		5.21	0.03	URE 1	1	
63777112	**Very good identification**					**63777112**
Actinomyces israelii		98.97	0.74	MLZ 13		
Bifidobacterium adolescentis		0.94	0.33			
Bifidobacterium dentium		0.05	-.07			
63777113	**Good identification**					**63777113**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.40	MLZ 13			
63777120	**Very good identification**						**63777120**
					GRA		
Actinomyces israelii		98.72	0.65	NAG 6	99		
Mitsuokella multacida		0.87	0.17		1		
Actinomyces naeslundii		0.42	0.33		99		
63777122	**Very good identification**						**63777122**
Actinomyces israelii		99.72	0.68	NAG 6			
Actinomyces naeslundii		0.28	0.33				
63777130	**Good identification**						**63777130**
					GRA		
Actinomyces israelii		98.08	0.51	MLZ 13	99		
Mitsuokella multacida		1.92	0.09	NAG 6	1		
63777132	**Good identification**						**63777132**
Actinomyces israelii		98.61	0.54	MLZ 13			
Bifidobacterium dentium		1.39	-.03	NAG 6			
63777140	**Very good identification**						**63777140**
					GRA		
Actinomyces israelii		99.43	0.78		99		
Mitsuokella multacida		0.57	0.27		1		
63777141	**Acceptable identification**						**63777141**
					GRA		
Actinomyces israelii		92.46	0.44	URE 1	99		
Mitsuokella multacida		5.86	0.11		1		
Actinomyces naeslundii		1.69	0.23		99		
63777142	**Very good identification**						**63777142**
					GRA		
Actinomyces israelii		99.92	0.81		99		
Mitsuokella multacida		0.04	0.11		1		
Bifidobacterium adolescentis		0.04	0.16		99		
63777143	**Good identification**						**63777143**
Actinomyces israelii		98.80	0.47	URE 1			
Actinomyces naeslundii		1.20	0.23				
63777150	**Very good identification**						**63777150**
					COC	GRA	
Actinomyces israelii		98.54	0.64	MLZ 13	1	99	
Mitsuokella multacida		1.27	0.19		1	1	
Blautia producta		0.19	-.03		99	99	
63777152	**Very good identification**						**63777152**
					GRA		
Actinomyces israelii		99.60	0.67	MLZ 13	99		
Bifidobacterium adolescentis		0.30	0.18		99		
Mitsuokella multacida		0.10	0.03		1		
63777160	**Very good identification**						**63777160**
					GRA		
Actinomyces israelii		97.80	0.58	NAG 6	99		
Mitsuokella multacida		2.21	0.17		1		
63777162	**Very good identification**						**63777162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.61	NAG 6		
63777170	**Good identification**					**63777170**
					GRA	
Actinomyces israelii		95.21	0.44	MLZ 13	99	
Mitsuokella multacida		4.79	0.09	NAG 6	1	
63777172	**Good identification**					**63777172**
Actinomyces israelii		100.0	0.47	MLZ 13		
63777200	**Good identification**					**63777200**
Actinomyces israelii		100.0	0.56	RHA 7		
63777202	**Very good identification**					**63777202**
Actinomyces israelii		100.0	0.59	RHA 7		
63777210	**Good identification**					**63777210**
Actinomyces israelii		100.0	0.41	RHA 7		
63777212	**Good identification**					**63777212**
Actinomyces israelii		100.0	0.44	RHA 7		
63777220	**Good identification**					**63777220**
Actinomyces israelii		100.0	0.35	RHA 7		
63777222	**Good identification**					**63777222**
Actinomyces israelii		100.0	0.38	RHA 7		
63777240	**Good identification**					**63777240**
Actinomyces israelii		100.0	0.49	RHA 7		
63777242	**Good identification**					**63777242**
Actinomyces israelii		100.0	0.52	RHA 7		
63777252	**Good identification**					**63777252**
Actinomyces israelii		100.0	0.37	RHA 7		
63777300	**Good identification**					**63777300**
					GRA	
Actinomyces israelii		94.76	0.66	RHA 7	99	
Mitsuokella multacida		5.24	0.32		1	
63777302	**Very good identification**					**63777302**
					GRA	
Actinomyces israelii		99.59	0.69	RHA 7	99	
Mitsuokella multacida		0.41	0.15		1	
63777310	**Low discrimination**					**63777310**
					GRA	
Actinomyces israelii		89.03	0.52	RHA 7	99	
Mitsuokella multacida		10.97	0.24	MLZ 13	1	
63777312	**Good identification**					**63777312**
					GRA	
Actinomyces israelii		99.10	0.55	RHA 7	99	
Mitsuokella multacida		0.91	0.07	MLZ 13	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63777320	**Low discrimination**						**63777320**
					GRA		
Actinomyces israelii		82.21	0.45	RHA 7	99		
Mitsuokella multacida		17.79	0.21	NAG 6	1		
63777322	**Good identification**						**63777322**
					SPO	GRA	
Actinomyces israelii		97.53	0.48	RHA 7	1	99	
Mitsuokella multacida		1.56	0.05	NAG 6	1	1	
Clostridium ramosum		0.90	-.03		92	50	
63777340	**Low discrimination**						**63777340**
					GRA		
Actinomyces israelii		87.56	0.59	RHA 7	99		
Mitsuokella multacida		12.44	0.32		1		
63777342	**Very good identification**						**63777342**
					GRA		
Actinomyces israelii		98.96	0.62	RHA 7	99		
Mitsuokella multacida		1.04	0.15		1		
63777350	**Low discrimination**						**63777350**
					COC	GRA	
Actinomyces israelii		74.49	0.45	RHA 7	1	99	
Mitsuokella multacida		23.61	0.24	MLZ 13	1	1	
Blautia producta		1.91	-.03		99	99	
63777352	**Good identification**						**63777352**
					GRA		
Actinomyces israelii		97.71	0.48	RHA 7	99		
Mitsuokella multacida		2.29	0.07	MLZ 13	1		
63777360	**Low discrimination**						**63777360**
					GRA		
Actinomyces israelii		64.25	0.38	RHA 7	99		
Mitsuokella multacida		35.75	0.21	NAG 6	1		
63777362	**Good identification**						**63777362**
					GRA		
Actinomyces israelii		96.04	0.41	RHA 7	99		
Mitsuokella multacida		3.96	0.05	NAG 6	1		
63777400	**Very good identification**						**63777400**
					GRA		
Actinomyces israelii		98.92	0.73		99		
Mitsuokella multacida		1.08	0.27		1		
63777401	**Low discrimination**						**63777401**
					GRA		
Actinomyces israelii		89.25	0.39	URE 1	99		
Mitsuokella multacida		10.75	0.11		1		
63777402	**Very good identification**						**63777402**
					GRA		
Actinomyces israelii		99.88	0.76		99		
Mitsuokella multacida		0.08	0.11		1		
Bifidobacterium adolescentis		0.04	0.12		99		
63777403	**Good identification**						**63777403**
Actinomyces israelii		100.0	0.42	URE 1			
63777410	**Very good identification**						**63777410**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		97.62	0.59	MLZ 13	99		
Mitsuokella multacida		2.38	0.19		1		
63777412	**Very good identification**						**63777412**
Actinomyces israelii		99.26	0.62	MLZ 13			
Bifidobacterium adolescentis		0.31	0.13				
Bifidobacterium dentium		0.25	-.07				
63777420	**Good identification**						**63777420**
					GRA		
Actinomyces israelii		95.89	0.53	NAG 6	99		
Mitsuokella multacida		4.11	0.17		1		
63777422	**Good identification**						**63777422**
Actinomyces israelii		100.0	0.56	NAG 6			
63777430	**Acceptable identification**						**63777430**
					GRA		
Actinomyces israelii		91.27	0.39	MLZ 13	99		
Mitsuokella multacida		8.73	0.09	NAG 6	1		
63777432	**Acceptable identification**						**63777432**
Actinomyces israelii		93.55	0.42	MLZ 13			
Bifidobacterium dentium		6.46	-.03	NAG 6			
63777440	**Very good identification**						**63777440**
					GRA		
Actinomyces israelii		97.26	0.66		99		
Mitsuokella multacida		2.74	0.27		1		
63777442	**Very good identification**						**63777442**
					GRA		
Actinomyces israelii		99.79	0.69		99		
Mitsuokella multacida		0.21	0.11		1		
63777443	**Good identification**						**63777443**
Actinomyces israelii		100.0	0.35	URE 1			
63777450	**Acceptable identification**						**63777450**
					COC	GRA	
Actinomyces israelii		93.27	0.52	MLZ 13	1	99	
Mitsuokella multacida		5.86	0.19		1	1	
Blautia producta		0.88	-.03		99	99	
63777452	**Good identification**						**63777452**
					GRA		
Actinomyces israelii		99.54	0.55	MLZ 13	99		
Mitsuokella multacida		0.46	0.03		1		
63777460	**Acceptable identification**						**63777460**
					GRA		
Actinomyces israelii		90.07	0.46	NAG 6	99		
Mitsuokella multacida		9.93	0.17		1		
63777462	**Good identification**						**63777462**
Actinomyces israelii		100.0	0.49	NAG 6			
63777500	**Low discrimination**						**63777500**
					GRA	SPO	
Actinomyces israelii		78.56	0.84		99	1	

Code / Taxon		ID score	T index	tests against		Differentiation tests		Code
Mitsuokella multacida		21.29	0.61			1	1	
Clostridium butyricum		0.14	0.15			99	99	
63777501	**Low discrimination**							**63777501**
						GRA		
Mitsuokella multacida		74.88	0.45	MNS	99	1		
Actinomyces israelii		25.12	0.49	URE	10	99		
63777502	**Very good identification**							**63777502**
						GRA		
Actinomyces israelii		97.07	0.87			99		
Mitsuokella multacida		1.95	0.45			1		
Bifidobacterium adolescentis		0.93	0.46			99		
63777503	**Low discrimination**							**63777503**
						GRA		
Actinomyces israelii		81.28	0.52	URE	1	99		
Mitsuokella multacida		17.94	0.29			1		
Bifidobacterium adolescentis		0.78	0.12			99		
63777510	**Low discrimination**							**63777510**
						GRA		
Actinomyces israelii		61.87	0.69	MLZ	13	99		
Mitsuokella multacida		37.40	0.53			1		
Bifidobacterium dentium		0.56	0.10			99		
63777511	**Low discrimination**							**63777511**
						GRA		
Mitsuokella multacida		86.93	0.37	MNS	99	1		
Actinomyces israelii		13.07	0.35	URE	10	99		
63777512	**Low discrimination**							**63777512**
Actinomyces israelii		84.46	0.73	MLZ	13			
Bifidobacterium adolescentis		6.58	0.47					
Bifidobacterium dentium		5.18	0.27					
63777513	**Low discrimination**							**63777513**
						GRA		
Actinomyces israelii		61.28	0.38	MLZ	13	99		
Mitsuokella multacida		30.18	0.20	URE	1	1		
Bifidobacterium adolescentis		4.78	0.13			99		
63777520	**Low discrimination**							**63777520**
						GRA	SPO	
Mitsuokella multacida		51.43	0.51	MNS	99	1	1	
Actinomyces israelii		48.46	0.63			99	1	
Clostridium butyricum		0.12	-.03			99	99	
63777522	**Good identification**							**63777522**
						GRA		
Actinomyces israelii		92.36	0.66	NAG	6	99		
Mitsuokella multacida		7.26	0.35			1		
Bifidobacterium dentium		0.24	-.03			99		
63777530	**Low discrimination**							**63777530**
						GRA		
Mitsuokella multacida		65.38	0.43	MNS	99	1		
Actinomyces israelii		27.62	0.49			99		
Bifidobacterium dentium		7.00	0.15			99		
63777540	**Low discrimination**							**63777540**
						COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		58.52	0.77		1	99	
Mitsuokella multacida		40.77	0.61		1	1	
Blautia producta		0.68	0.24		99	99	
63777541	**Low discrimination**						**63777541**
					GRA		
Mitsuokella multacida		88.46	0.45	MNS 99	1		
Actinomyces israelii		11.54	0.42	URE 10	99		
63777542	**Good identification**						**63777542**
					GRA		
Actinomyces israelii		94.74	0.80		99		
Mitsuokella multacida		4.89	0.45		1		
Bifidobacterium adolescentis		0.29	0.30		99		
63777543	**Low discrimination**						**63777543**
					GRA		
Actinomyces israelii		63.79	0.45	URE 1	99		
Mitsuokella multacida		36.21	0.29		1		
63777550	**Low discrimination**						**63777550**
					COC	GRA	
Mitsuokella multacida		55.76	0.53	MNS 99	1	1	
Actinomyces israelii		35.88	0.62		1	99	
Blautia producta		8.37	0.32		99	99	
63777551	**Acceptable identification**						**63777551**
					COC	GRA	
Mitsuokella multacida		93.27	0.37	MNS 99	1	1	
Actinomyces israelii		5.46	0.28	URE 10	1	99	
Blautia producta		1.27	-.03		99	99	
63777552	**Low discrimination**						**63777552**
					GRA		
Actinomyces israelii		86.30	0.65	MLZ 13	99		
Mitsuokella multacida		9.94	0.37		1		
Bifidobacterium adolescentis		2.14	0.32		99		
63777560	**Low discrimination**						**63777560**
					COC	GRA	
Mitsuokella multacida		72.79	0.51	MNS 99	1	1	
Actinomyces israelii		26.67	0.56		1	99	
Blautia producta		0.54	0.07		99	99	
63777562	**Low discrimination**						**63777562**
					GRA		
Actinomyces israelii		83.19	0.59	NAG 6	99		
Mitsuokella multacida		16.82	0.35		1		
63777570	**Low discrimination**						**63777570**
					COC	GRA	
Mitsuokella multacida		81.24	0.43	MNS 99	1	1	
Actinomyces israelii		13.35	0.42		1	99	
Blautia producta		5.42	0.15		99	99	
63777572	**Low discrimination**						**63777572**
					GRA		
Actinomyces israelii		65.56	0.45	MLZ 13	99		
Mitsuokella multacida		29.56	0.26	NAG 6	1		
Bifidobacterium dentium		2.91	-.03		99		
63777600	**Low discrimination**						**63777600**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		78.73	0.54	RHA 7	99		
Mitsuokella multacida		21.27	0.32		1		
63777602	**Very good identification**						**63777602**
					GRA		
Actinomyces israelii		98.04	0.57	RHA 7	99		
Mitsuokella multacida		1.96	0.15		1		
63777610	**Low discrimination**						**63777610**
					GRA		
Actinomyces israelii		62.40	0.40	RHA 7	99		
Mitsuokella multacida		37.60	0.24	MLZ 13	1		
63777612	**Good identification**						**63777612**
					GRA		
Actinomyces israelii		95.73	0.43	RHA 7	99		
Mitsuokella multacida		4.27	0.07	MLZ 13	1		
63777622	**Acceptable identification**						**63777622**
					GRA		
Actinomyces israelii		92.73	0.37	RHA 7	99		
Mitsuokella multacida		7.27	0.05	NAG 6	1		
63777640	**Low discrimination**						**63777640**
					GRA		
Actinomyces israelii		59.01	0.47	RHA 7	99		
Mitsuokella multacida		40.99	0.32		1		
63777642	**Good identification**						**63777642**
					GRA		
Actinomyces israelii		95.11	0.50	RHA 7	99		
Mitsuokella multacida		4.89	0.15		1		
63777652	**Low discrimination**						**63777652**
					GRA		
Actinomyces israelii		89.70	0.36	RHA 7	99		
Mitsuokella multacida		10.30	0.07	MLZ 13	1		
63777700	**Low discrimination**						**63777700**
					GRA		
Mitsuokella multacida		86.99	0.66	MNS 99	1		
Actinomyces israelii		13.01	0.64		99		
63777701	**Good identification**						**63777701**
					GRA		
Mitsuokella multacida		98.66	0.50	MNS 99	1		
Actinomyces israelii		1.34	0.30	URE 10	99		
63777702	**Low discrimination**						**63777702**
					GRA		
Actinomyces israelii		66.82	0.67	RHA 7	99		
Mitsuokella multacida		33.09	0.50		1		
Bifidobacterium adolescentis		0.09	0.12		99		
63777710	**Good identification**						**63777710**
					COC	GRA	
Mitsuokella multacida		93.64	0.58	MNS 99	1	1	
Actinomyces israelii		6.28	0.50		1	99	
Blautia producta		0.08	-.03		99	99	
63777711	**Good identification**						**63777711**
					GRA		
Mitsuokella multacida		99.39	0.41	MNS 99	1		
Actinomyces israelii		0.61	0.16	URE 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
63777712	**Low discrimination**						**63777712**
					GRA		
Mitsuokella multacida		52.03	0.41	MNS 99	1		
Actinomyces israelii		47.10	0.53	BGL 10	99		
Bifidobacterium adolescentis		0.49	0.13		99		
63777720	**Good identification**						**63777720**
					GRA		
Mitsuokella multacida		96.32	0.56	MNS 99	1		
Actinomyces israelii		3.68	0.44		99		
63777721	**Good identification**						**63777721**
Mitsuokella multacida		100.0	0.39	MNS 99			
63777722	**Low discrimination**						**63777722**
					GRA		
Mitsuokella multacida		65.56	0.39	MNS 99	1		
Actinomyces israelii		33.80	0.47	BGL 10	99		
Bacteroides ovatus		0.64	-.02		1		
63777730	**Good identification**						**63777730**
					GRA		
Mitsuokella multacida		98.32	0.47	MNS 99	1		
Actinomyces israelii		1.68	0.30		99		
63777740	**Good identification**						**63777740**
					COC	GRA	
Mitsuokella multacida		93.71	0.66	MNS 99	1	1	
Actinomyces israelii		5.45	0.57		1	99	
Blautia producta		0.84	0.24		99	99	
63777741	**Good identification**						**63777741**
					GRA		
Mitsuokella multacida		99.47	0.50	MNS 99	1		
Actinomyces israelii		0.53	0.23	URE 10	99		
63777742	**Low discrimination**						**63777742**
					COC	GRA	
Mitsuokella multacida		55.73	0.50	MNS 99	1	1	
Actinomyces israelii		43.77	0.60	BGL 10	1	99	
Blautia producta		0.50	0.07		99	99	
63777750	**Good identification**						**63777750**
					COC	GRA	
Mitsuokella multacida		90.35	0.58	MNS 99	1	1	
Blautia producta		7.30	0.32		99	99	
Actinomyces israelii		2.36	0.43		1	99	
63777751	**Good identification**						**63777751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	MNS 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
63777752	**Low discrimination**						**63777752**
					COC	GRA	
Mitsuokella multacida		69.79	0.41	MNS 99	1	1	
Actinomyces israelii		24.57	0.46	BGL 10	1	99	
Blautia producta		5.64	0.15		99	99	
63777760	**Good identification**						**63777760**
					COC	GRA	
Mitsuokella multacida		98.15	0.56	MNS 99	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.46	0.37		1	99	
Blautia producta		0.39	0.07		99	99	
63777761	**Good identification**						**63777761**
Mitsuokella multacida		100.0	0.39	MNS 99			
63777762	**Low discrimination**						**63777762**
					GRA		
Mitsuokella multacida		83.30	0.39	MNS 99	1		
Actinomyces israelii		16.70	0.40	BGL 10	99		
63777770	**Good identification**						**63777770**
					COC	GRA	
Mitsuokella multacida		95.92	0.47	MNS 99	1	1	
Blautia producta		3.44	0.15		99	99	
Actinomyces israelii		0.64	0.23		1	99	
64000002	**Low discrimination**						**64000002**
					COC	GRA	
Atopobium minutum		69.42	0.43	NIT 1	1	99	
Campylobacter gracilis		19.46	0.15		1	1	
Anaerococcus prevotii		2.57	0.05		99	99	
64001000	**Low discrimination**						**64001000**
					COC	GRA	
Atopobium minutum		89.26	0.38	NIT 1	1	99	
Veillonella parvula		6.19	-.03		99	1	
Campylobacter gracilis		4.55	-.03		1	1	
64001002	**Good identification**						**64001002**
Atopobium minutum		100.0	0.35	NIT 1			
64010000	**Low discrimination**						**64010000**
					GRA		
Atopobium minutum		74.62	0.46	NIT 1	99		
Eubacterium limosum		9.61	0.18		99		
Fusobacterium varium		8.27	0.15		1		
64010002	**Good identification**						**64010002**
Atopobium minutum		100.0	0.43	NIT 1			
64010020	**Low discrimination**						**64010020**
					GRA	CAT	
Atopobium minutum		83.69	0.37	NIT 1	99	1	
Propionibacterium acnes		15.15	0.18		99	95	
Campylobacter gracilis		1.16	-.13		1	1	
64011000	**Good identification**						**64011000**
Atopobium minutum		100.0	0.37	NIT 1			
65000000	**Low discrimination**						**65000000**
					SPO		
Clostridium botulinum biovar A		40.84	0.47	NIT 1	99		
Clostridium bifermentans		27.89	0.43	MNS 7	99		
Atopobium minutum		23.35	0.50		1		
65000002	**Low discrimination**						**65000002**
					SPO		
Atopobium minutum		88.42	0.47	NIT 1	1		
Clostridium sporogenes		7.63	0.19		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium botulinum biovar A		2.34	0.13		99		
65000020	**Low discrimination**						**65000020**
					SPO	CAT	
Atopobium minutum		70.98	0.42	NIT 1	1	1	
Clostridium bifermentans		16.24	0.23		99	1	
Propionibacterium acnes		6.97	0.18		1	95	
65000022	**Good identification**						**65000022**
					GRA		
Atopobium minutum		97.51	0.39	NIT 1	99		
Pseudoflavonifractor capillosus		2.49	-.03		1		
65000040	**Low discrimination**						**65000040**
					SPO		
Clostridium botulinum biovar A		88.86	0.38	NIT 1	99		
Clostridium bifermentans		8.46	0.20	MNS 7	99		
Atopobium minutum		1.72	0.16		1		
65001000	**Low discrimination**						**65001000**
					COC	SPO	
Atopobium minutum		88.04	0.42	NIT 1	1	1	
Clostridium botulinum biovar A		4.67	0.13		1	99	
Gemella morbillorum		3.28	0.13		99	1	
65001002	**Good identification**						**65001002**
Atopobium minutum		100.0	0.39	NIT 1			
65001520	**Good identification**						**65001520**
Odoribacter splanchnicus		100.0	0.51	IND 99			
65010000	**Low discrimination**						**65010000**
					SPO		
Eubacterium limosum		75.63	0.52	NIT 1	1		
Atopobium minutum		10.10	0.50	MAN 86	1		
Clostridium difficile		8.34	0.38		99		
65010002	**Low discrimination**						**65010002**
					SPO		
Atopobium minutum		86.80	0.47	NIT 1	1		
Eubacterium limosum		9.85	0.18		1		
Clostridium sporogenes		1.43	0.06		99		
65010010	**Good identification**						**65010010**
					SPO		
Clostridium difficile		97.92	0.50	NIT 1	99		
Eubacterium limosum		1.84	0.18	MAN 90	1		
Atopobium minutum		0.25	0.15		1		
65010020	**Low discrimination**						**65010020**
					CAT		
Propionibacterium acnes		89.08	0.52	IND 87	95		
Atopobium minutum		8.46	0.41	ESL 1	1		
Eubacterium limosum		1.92	0.18		1		
65010022	**Low discrimination**						**65010022**
					CAT		
Atopobium minutum		86.24	0.38	NIT 1	1		
Propionibacterium acnes		13.76	0.18		95		
65010060	**Good identification**						**65010060**
					CAT		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		99.57	0.40	IND 87	95		
Atopobium minutum		0.44	0.07	ESL 1	1		
65010100	**Low discrimination**						**65010100**
					GRA		
Fusobacterium mortiferum		58.85	0.40	NIT 1	1		
Eubacterium limosum		33.09	0.18		99		
Atopobium minutum		4.42	0.15		99		
65011000	**Low discrimination**						**65011000**
					GRA		
Atopobium minutum		59.98	0.41	NIT 1	99		
Fusobacterium mortiferum		15.57	0.37		1		
Eubacterium limosum		13.61	0.18		99		
65011002	**Good identification**						**65011002**
					GRA		
Atopobium minutum		95.85	0.38	NIT 1	99		
Fusobacterium mortiferum		4.15	0.20		1		
65011020	**Good identification**						**65011020**
					CAT		
Propionibacterium acnes		96.65	0.52	IND 87	95		
Atopobium minutum		3.06	0.33	ESL 1	1		
Clostridium perfringens		0.29	0.10		1		
65011060	**Good identification**						**65011060**
Propionibacterium acnes		100.0	0.40	IND 87			
65011100	**Good identification**						**65011100**
					GRA		
Fusobacterium mortiferum		98.36	0.43	NIT 1	1		
Atopobium minutum		1.64	0.07		99		
65011520	**Good identification**						**65011520**
Odoribacter splanchnicus		100.0	0.56	IND 99			
65011560	**Good identification**						**65011560**
Odoribacter splanchnicus		100.0	0.39	IND 99			
65012000	**Low discrimination**						**65012000**
					SPO	CAT	
Eubacterium limosum		86.04	0.66	NIT 1	1	1	
Clostridium difficile		13.90	0.54		99	1	
Propionibacterium acnes		0.04	0.13		1	95	
65012010	**Very good identification**						**65012010**
					SPO		
Clostridium difficile		98.74	0.66	NIT 1	99		
Eubacterium limosum		1.26	0.32		1		
65012020	**Low discrimination**						**65012020**
					SPO	CAT	
Propionibacterium acnes		78.42	0.47	IND 87	1	95	
Eubacterium limosum		18.47	0.32	ESL 1	1	1	
Clostridium difficile		2.98	0.20		99	1	
65012050	**Good identification**						**65012050**
					SPO		
Clostridium difficile		99.76	0.44	NIT 1	99		
Eubacterium limosum		0.25	-.03	SOR 5	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
65012060	**Good identification**						**65012060**
					SPO	CAT	
Propionibacterium acnes		98.04	0.36	IND 87	1	95	
Eubacterium limosum		1.06	-.03	ESL 1	1	1	
Clostridium difficile		0.89	-.02		99	1	
65013020	**Good identification**						**65013020**
					CAT		
Propionibacterium acnes		99.76	0.47	IND 87	95		
Eubacterium limosum		0.24	-.03	ESL 1	1		
65013060	**Good identification**						**65013060**
Propionibacterium acnes		100.0	0.36	IND 87			
65016010	**Good identification**						**65016010**
					SPO		
Clostridium difficile		99.76	0.44	NIT 1	99		
Eubacterium limosum		0.25	-.03	XYL 5	1		
65020000	**Low discrimination**						**65020000**
Clostridium botulinum biovar A		84.58	0.36	NIT 1			
Clostridium sporogenes		11.13	0.19	MNS 7			
Clostridium bifermentans		2.33	0.09				
65032010	**Good identification**						**65032010**
					SPO		
Clostridium difficile		99.83	0.47	NIT 1	99		
Eubacterium limosum		0.17	-.03	TRE 7	1		
65052010	**Good identification**						**65052010**
					SPO		
Clostridium difficile		99.76	0.44	NIT 1	99		
Eubacterium limosum		0.25	-.03	CEL 5	1		
65071120	**Good identification**						**65071120**
Clostridium septicum		99.18	0.35	NIT 1			
Clostridium perfringens		0.82	0.06	MLT 94			
65100000	**Low discrimination**						**65100000**
					GRA	SPO	
Clostridium bifermentans		53.48	0.50	IND 90	99	99	
Clostridium botulinum biovar A		27.47	0.45	NIT 1	99	99	
Prevotella disiens		9.80	0.32		1	1	
65100040	**Low discrimination**						**65100040**
Clostridium botulinum biovar A		76.23	0.36	NIT 1			
Clostridium bifermentans		20.71	0.26	MNS 7			
Clostridium novyi biovar A		1.01	0.16				
65101101	**Very good identification**						**65101101**
Clostridium chauvoei		100.0	0.60	SUC 99			
65101103	**Good identification**						**65101103**
Clostridium chauvoei		100.0	0.47	SUC 99			
65101121	**Good identification**						**65101121**
Clostridium chauvoei		100.0	0.51	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
65101123	**Good identification**						**65101123**
Clostridium chauvoei		100.0	0.39	SUC 99			
65111020	**Acceptable identification**						**65111020**
					CAT	SPO	
Clostridium perfringens		94.41	0.45	SUC 91	1	84	
Propionibacterium acnes		3.24	0.18	LAC 88	95	1	
Clostridium septicum		1.09	0.05		1	99	
65111022	**Acceptable identification**						**65111022**
					GRA	CAT	
Clostridium perfringens		94.00	0.39	SUC 91	99	1	
Capnocytophaga ochracea		6.00	0.15	LAC 88	1	90	
65111101	**Very good identification**						**65111101**
Clostridium chauvoei		100.0	0.66	SUC 99			
65111103	**Good identification**						**65111103**
					GRA	SPO	
Clostridium chauvoei		99.82	0.53	SUC 99	99	99	
Capnocytophaga ochracea		0.10	-.01		1	1	
Actinomyces viscosus		0.09	-.01		99	1	
65111120	**Low discrimination**						**65111120**
					GRA	CAT	
Clostridium perfringens		82.12	0.59	SUC 91	99	1	
Clostridium septicum		12.84	0.39		99	1	
Capnocytophaga ochracea		3.70	0.33		1	90	
65111121	**Very good identification**						**65111121**
Clostridium chauvoei		98.65	0.58	SUC 99			
Clostridium perfringens		1.13	0.25				
Clostridium septicum		0.18	0.05				
65111122	**Low discrimination**						**65111122**
					GRA	CAT	
Clostridium perfringens		53.11	0.54	SUC 91	99	1	
Capnocytophaga ochracea		45.74	0.50		1	90	
Prevotella oralis		0.51	0.16		1	1	
65111123	**Acceptable identification**						**65111123**
					GRA	CAT	
Clostridium chauvoei		94.64	0.45	SUC 99	99	1	
Clostridium perfringens		2.88	0.20		99	1	
Capnocytophaga ochracea		2.48	0.15		1	90	
65111160	**Good identification**						**65111160**
					GRA	CAT	
Clostridium perfringens		96.66	0.47	SUC 91	99	1	
Capnocytophaga ochracea		2.54	0.17		1	90	
Clostridium septicum		0.80	0.05		99	1	
65111162	**Low discrimination**						**65111162**
					GRA	CAT	
Clostridium perfringens		66.56	0.41	SUC 91	99	1	
Capnocytophaga ochracea		33.44	0.33		1	90	
65111522	**Acceptable identification**						**65111522**
					GRA	CAT	
Capnocytophaga ochracea		93.77	0.37	SUC 99	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		6.23	0.20	RAF 90	99	1	
65115122	**Acceptable identification**						**65115122**
					GRA	CAT	
Capnocytophaga ochracea		91.62	0.39	SUC 99	1	90	
Clostridium perfringens		4.30	0.20	RAF 90	99	1	
Bacteroides fragilis		3.57	0.06		1	99	
65121101	**Good identification**						**65121101**
Clostridium chauvoei		100.0	0.43	SUC 99			
65121121	**Good identification**						**65121121**
Clostridium chauvoei		100.0	0.35	SUC 99			
65131020	**Good identification**						**65131020**
Clostridium perfringens		98.16	0.49	SUC 91			
Clostridium septicum		1.84	0.13	LAC 88			
65131022	**Good identification**						**65131022**
Clostridium perfringens		100.0	0.44	SUC 91			
65131060	**Good identification**						**65131060**
Clostridium perfringens		100.0	0.37	SUC 91			
65131100	**Low discrimination**						**65131100**
					COC	SPO	
Clostridium perfringens		67.33	0.38	SUC 91	1	84	
Atopobium parvulum		22.47	0.29	NAG 97	99	1	
Clostridium septicum		5.55	0.13		1	99	
65131101	**Good identification**						**65131101**
Clostridium chauvoei		99.80	0.50	SUC 99			
Clostridium perfringens		0.20	0.04	TRE 10			
65131103	**Good identification**						**65131103**
Clostridium chauvoei		100.0	0.37	SUC 99			
65131120	**Low discrimination**						**65131120**
					COC	SPO	
Clostridium perfringens		79.68	0.64	SUC 91	1	84	
Clostridium septicum		20.12	0.47		1	99	
Atopobium parvulum		0.15	0.16		99	1	
65131121	**Low discrimination**						**65131121**
Clostridium chauvoei		80.71	0.41	SUC 99			
Clostridium perfringens		15.40	0.30	TRE 10			
Clostridium septicum		3.89	0.13				
65131122	**Very good identification**						**65131122**
					GRA	CAT	
Clostridium perfringens		98.40	0.58	SUC 91	99	1	
Clostridium septicum		1.08	0.18		99	1	
Capnocytophaga ochracea		0.46	0.15		1	90	
65131160	**Good identification**						**65131160**
Clostridium perfringens		98.68	0.52	SUC 91			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium septicum		1.32	0.13				
65131162	**Good identification**						**65131162**
					GRA	CAT	
Clostridium perfringens		99.73	0.46	SUC 91	99	1	
Capnocytophaga ochracea		0.27	-.01		1	90	
65151120	**Low discrimination**						**65151120**
					GRA	CAT	
Clostridium septicum		88.87	0.47	NIT 1	99	1	
Clostridium perfringens		7.48	0.36		99	1	
Capnocytophaga ochracea		2.02	0.23		1	90	
65151122	**Low discrimination**						**65151122**
					GRA	CAT	
Capnocytophaga ochracea		67.14	0.39	SUC 99	1	90	
Clostridium perfringens		12.99	0.30	RAF 90	99	1	
Prevotella oralis		10.68	0.25		1	1	
65171100	**Low discrimination**						**65171100**
					COC	SPO	
Atopobium parvulum		72.00	0.36	NIT 1	99	1	
Clostridium septicum		24.15	0.21	SUC 96	1	99	
Clostridium perfringens		3.85	0.15		1	84	
65171120	**Acceptable identification**						**65171120**
					COC	SPO	
Clostridium septicum		94.45	0.56	NIT 1	1	99	
Clostridium perfringens		4.92	0.40		1	84	
Atopobium parvulum		0.50	0.23		99	1	
65175100	**Good identification**						**65175100**
					COC		
Atopobium parvulum		98.67	0.40	NIT 1	99		
Actinomyces israelii		1.33	0.22	SUC 96	1		
65177102	**Good identification**						**65177102**
Actinomyces israelii		100.0	0.38	SUC 98			
65177502	**Good identification**						**65177502**
Actinomyces israelii		100.0	0.36	SUC 98			
65210010	**Good identification**						**65210010**
Clostridium difficile		100.0	0.40	NIT 1			
65210100	**Good identification**						**65210100**
Fusobacterium mortiferum		100.0	0.45	NIT 1			
65211000	**Good identification**						**65211000**
					GRA		
Fusobacterium mortiferum		96.13	0.42	NIT 1	1		
Eubacterium tortuosum		1.94	-.04		99		
Atopobium minutum		1.93	0.07		99		
65211100	**Good identification**						**65211100**
					GRA	SPO	
Fusobacterium mortiferum		99.23	0.48	NIT 1	1	1	
Clostridium septicum		0.77	-.06		99	99	
65212000	**Low discrimination**						**65212000**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium difficile		81.86	0.45	NIT 1	99		
Eubacterium limosum		18.14	0.32		1		
65212010	**Very good identification**						**65212010**
					SPO		
Clostridium difficile		99.95	0.56	NIT 1	99		
Eubacterium limosum		0.05	-.03		1		
65230100	**Good identification**						**65230100**
Fusobacterium mortiferum		100.0	0.36	NIT 1			
65231100	**Acceptable identification**						**65231100**
					GRA	SPO	
Fusobacterium mortiferum		92.73	0.39	NIT 1	1	1	
Clostridium septicum		7.27	0.03		99	99	
65231120	**Good identification**						**65231120**
Clostridium septicum		96.23	0.37	NIT 1			
Clostridium perfringens		3.77	0.20	MLT 94			
65232010	**Good identification**						**65232010**
Clostridium difficile		100.0	0.37	NIT 1			
65250100	**Good identification**						**65250100**
Fusobacterium mortiferum		100.0	0.37	NIT 1			
65251100	**Low discrimination**						**65251100**
					GRA		
Fusobacterium mortiferum		73.20	0.40	NIT 1	1		
Lactobacillus catenaformis		10.30	0.15		99		
Collinsella aerofaciens		6.82	-.03		99		
65251120	**Acceptable identification**						**65251120**
Clostridium septicum		91.77	0.37	NIT 1			
Clostridium paraputrificum		5.61	0.14	MLT 94			
Clostridium barati		2.18	0.12				
65253000	**Good identification**						**65253000**
Clostridium innocuum		100.0	0.49	NIT 1			
65253002	**Good identification**						**65253002**
Clostridium innocuum		100.0	0.49	NIT 1			
65253400	**Good identification**						**65253400**
Clostridium innocuum		100.0	0.36	NIT 1			
65253402	**Good identification**						**65253402**
Clostridium innocuum		100.0	0.36	NIT 1			
65271120	**Good identification**						**65271120**
Clostridium septicum		99.46	0.45	NIT 1			
Clostridium paraputrificum		0.54	0.04	MLT 94			
65273000	**Good identification**						**65273000**
Clostridium innocuum		100.0	0.41	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
65273002	**Good identification**						**65273002**
Clostridium innocuum		100.0	0.41	NIT 1			
65311120	**Low discrimination**						**65311120**
Clostridium septicum		70.74	0.49	NIT 1			
Clostridium perfringens		28.29	0.49				
Clostridium paraputrificum		0.87	0.14				
65311122	**Low discrimination**						**65311122**
					GRA	SPO	
Clostridium perfringens		78.67	0.43	SUC 91	99	84	
Clostridium septicum		8.53	0.20		99	99	
Prevotella oralis		5.41	0.20		1	1	
65311160	**Low discrimination**						**65311160**
Clostridium perfringens		87.35	0.37	SUC 91			
Clostridium septicum		11.58	0.15				
Clostridium paraputrificum		1.07	-.06				
65331020	**Low discrimination**						**65331020**
					SPO		
Clostridium perfringens		74.11	0.39	SUC 91	84		
Clostridium septicum		22.18	0.23	LAC 88	99		
Lactobacillus jensenii		3.71	0.17		1		
65331120	**Low discrimination**						**65331120**
					COC	SPO	
Clostridium septicum		79.92	0.57	NIT 1	1	99	
Clostridium perfringens		19.79	0.54		1	84	
Atopobium parvulum		0.19	0.18		99	1	
65331122	**Low discrimination**						**65331122**
Clostridium perfringens		84.25	0.48	SUC 91			
Clostridium septicum		14.76	0.28				
Clostridium paraputrificum		0.34	-.03				
65331160	**Low discrimination**						**65331160**
Clostridium perfringens		82.36	0.41	SUC 91			
Clostridium septicum		17.64	0.23				
65331162	**Good identification**						**65331162**
Clostridium perfringens		99.08	0.36	SUC 91			
Clostridium septicum		0.92	-.06				
65335100	**Good identification**						**65335100**
					COC		
Atopobium parvulum		97.99	0.36	NIT 1	99		
Actinomyces israelii		2.01	0.21	SUC 96	1		
65337102	**Good identification**						**65337102**
Actinomyces israelii		100.0	0.37	SUC 98			
65337502	**Good identification**						**65337502**
Actinomyces israelii		100.0	0.35	SUC 98			
65351000	**Low discrimination**						**65351000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Lactobacillus jensenii		81.12	0.41	NIT 1	1		
Lactobacillus catenaformis		14.69	0.29	SUC 90	1		
Clostridium barati		1.65	0.07		99		
65351002	**Low discrimination**						**65351002**
					SPO		
Lactobacillus jensenii		73.01	0.38	NIT 1	1		
Lactobacillus catenaformis		19.83	0.29	SUC 90	1		
Clostridium barati		6.67	0.15		99		
65351120	**Low discrimination**						**65351120**
					SPO		
Clostridium septicum		70.84	0.58	NIT 1	99		
Clostridium paraputrificum		27.38	0.48		99		
Collinsella aerofaciens		0.63	0.15		1		
65351122	**Low discrimination**						**65351122**
Clostridium paraputrificum		71.45	0.41	NIT 1			
Clostridium barati		9.60	0.32	SUC 92			
Clostridium septicum		8.80	0.29				
65357002	**Good identification**						**65357002**
Actinomyces israelii		100.0	0.36	SUC 98			
65357100	**Good identification**						**65357100**
Actinomyces israelii		100.0	0.43	SUC 98			
65357102	**Good identification**						**65357102**
					SPO		
Actinomyces israelii		97.98	0.46	SUC 98	1		
Clostridium tertium		2.02	0.03		99		
65357140	**Good identification**						**65357140**
Actinomyces israelii		100.0	0.36	SUC 98			
65357142	**Good identification**						**65357142**
Actinomyces israelii		100.0	0.39	SUC 98			
65357500	**Good identification**						**65357500**
Actinomyces israelii		100.0	0.42	SUC 98			
65357502	**Good identification**						**65357502**
Actinomyces israelii		97.68	0.45	SUC 98			
Bifidobacterium adolescentis		2.33	0.11				
65357542	**Good identification**						**65357542**
Actinomyces israelii		100.0	0.38	SUC 98			
65371000	**Good identification**						**65371000**
					COC	SPO	
Lactobacillus jensenii		99.50	0.50	NIT 1	1	1	
Clostridium septicum		0.25	-.03	SUC 90	1	99	
Atopobium parvulum		0.25	0.04		99	1	
65371002	**Good identification**						**65371002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus jensenii		98.74	0.47	NIT 1			
Lactobacillus acidophilus		0.74	0.07	SUC 90			
Actinomyces israelii		0.26	0.15				
65371010	**Good identification**						**65371010**
Lactobacillus jensenii		100.0	0.41	NIT 1			
65371012	**Low discrimination**						**65371012**
					GRA		
Lactobacillus jensenii		80.03	0.38	NIT 1	99		
Leptotrichia buccalis		19.97	0.15	SUC 90	1		
65371020	**Low discrimination**						**65371020**
					SPO		
Lactobacillus jensenii		48.06	0.39	NIT 1	1		
Clostridium septicum		47.86	0.32	SUC 90	99		
Clostridium perfringens		2.11	0.15		84		
65371022	**Low discrimination**						**65371022**
					SPO		
Lactobacillus jensenii		87.32	0.36	NIT 1	1		
Clostridium barati		2.78	0.05	SUC 90	99		
Clostridium perfringens		2.70	0.10		84		
65371112	**Good identification**						**65371112**
					GRA		
Leptotrichia buccalis		99.96	0.50	NIT 10	1		
Lactobacillus jensenii		0.04	0.04	SUC 99	99		
65371120	**Very good identification**						**65371120**
Clostridium septicum		96.18	0.66	NIT 1			
Clostridium paraputrificum		3.29	0.38				
Clostridium perfringens		0.31	0.30				
65371122	**Low discrimination**						**65371122**
					SPO		
Clostridium septicum		49.91	0.37	NIT 1	99		
Clostridium paraputrificum		35.89	0.32	BGL 2	99		
Lactobacillus acidophilus		5.71	0.25		1		
65373102	**Low discrimination**						**65373102**
					SPO		
Actinomyces israelii		62.98	0.38	SUC 98	1		
Clostridium tertium		25.90	0.17	XYL 92	99		
Actinomyces naeslundii		6.27	0.30		1		
65373502	**Good identification**						**65373502**
Actinomyces israelii		100.0	0.36	SUC 98			
65375100	**Low discrimination**						**65375100**
					COC	SPO	
Atopobium parvulum		88.82	0.42	NIT 1	99	1	
Actinomyces israelii		10.36	0.40	SUC 96	1	1	
Clostridium septicum		0.49	-.03		1	99	
65375142	**Good identification**						**65375142**
Actinomyces israelii		100.0	0.36	SUC 98			
65375500	**Low discrimination**						**65375500**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		51.01	0.39	SUC 98	99	1	
Mitsuokella multacida		25.53	0.21		1	1	
Clostridium butyricum		18.06	0.10		99	99	
65375502	**Low discrimination**						**65375502**
					GRA	SPO	
Actinomyces israelii		86.35	0.42	SUC 98	99	1	
Clostridium butyricum		7.93	0.03		99	99	
Mitsuokella multacida		3.20	0.04		1	1	
65377000	**Good identification**						**65377000**
Actinomyces israelii		100.0	0.43	SUC 98			
65377002	**Good identification**						**65377002**
Actinomyces israelii		100.0	0.46	SUC 98			
65377040	**Good identification**						**65377040**
Actinomyces israelii		100.0	0.36	SUC 98			
65377042	**Good identification**						**65377042**
Actinomyces israelii		100.0	0.39	SUC 98			
65377100	**Good identification**						**65377100**
					COC	SPO	
Actinomyces israelii		96.16	0.53	SUC 98	1	1	
Clostridium tertium		2.38	0.11		1	99	
Atopobium parvulum		1.47	0.08		99	1	
65377102	**Good identification**						**65377102**
					SPO		
Actinomyces israelii		95.29	0.56	SUC 98	1		
Clostridium tertium		4.71	0.19		99		
65377110	**Low discrimination**						**65377110**
					SPO		
Actinomyces israelii		77.29	0.39	SUC 98	1		
Clostridium tertium		22.71	0.15	MLZ 13	99		
65377112	**Low discrimination**						**65377112**
					SPO		
Actinomyces israelii		62.99	0.42	SUC 98	1		
Clostridium tertium		37.01	0.24	MLZ 13	99		
65377140	**Good identification**						**65377140**
Actinomyces israelii		100.0	0.46	SUC 98			
65377142	**Good identification**						**65377142**
Actinomyces israelii		100.0	0.49	SUC 98			
65377302	**Good identification**						**65377302**
Actinomyces israelii		100.0	0.37	SUC 98			
65377400	**Good identification**						**65377400**
Actinomyces israelii		100.0	0.41	SUC 98			
65377402	**Good identification**						**65377402**
Actinomyces israelii		100.0	0.44	SUC 98			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
65377442	**Good identification**						**65377442**
Actinomyces israelii		100.0	0.37	SUC 98			
65377500	**Low discrimination**						**65377500**
					GRA	SPO	
Actinomyces israelii		82.44	0.51	SUC 98	99	1	
Mitsuokella multacida		16.99	0.27		1	1	
Clostridium butyricum		0.57	-.07		99	99	
65377502	**Good identification**						**65377502**
					GRA		
Actinomyces israelii		98.09	0.55	SUC 98	99		
Mitsuokella multacida		1.50	0.11		1		
Bifidobacterium adolescentis		0.41	0.08		99		
65377510	**Low discrimination**						**65377510**
					GRA		
Actinomyces israelii		68.51	0.37	SUC 98	99		
Mitsuokella multacida		31.49	0.19	MLZ 13	1		
65377512	**Low discrimination**						**65377512**
Actinomyces israelii		76.25	0.40	SUC 98			
Bifidobacterium dentium		18.53	0.06	MLZ 13			
Bifidobacterium adolescentis		2.63	0.09				
65377540	**Low discrimination**						**65377540**
					GRA		
Actinomyces israelii		65.37	0.44	SUC 98	99		
Mitsuokella multacida		34.64	0.27		1		
65377542	**Good identification**						**65377542**
					GRA		
Actinomyces israelii		96.22	0.47	SUC 98	99		
Mitsuokella multacida		3.78	0.11		1		
65377702	**Low discrimination**						**65377702**
					GRA		
Actinomyces israelii		72.64	0.35	SUC 98	99		
Mitsuokella multacida		27.36	0.15	RHA 7	1		
65410000	**Low discrimination**						**65410000**
					GRA		
Fusobacterium mortiferum		54.33	0.39	NIT 1	1		
Eubacterium limosum		36.72	0.18		99		
Atopobium minutum		4.90	0.15		99		
65410100	**Good identification**						**65410100**
Fusobacterium mortiferum		100.0	0.45	NIT 1			
65411000	**Good identification**						**65411000**
					GRA		
Fusobacterium mortiferum		98.03	0.42	NIT 1	1		
Atopobium minutum		1.97	0.07		99		
65411100	**Good identification**						**65411100**
					CAT		
Fusobacterium mortiferum		99.13	0.48	NIT 1	5		
Capnocytophaga ochracea		0.87	-.01		90		
65430100	**Good identification**						**65430100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		100.0	0.36	NIT 1			
65431100	**Good identification**						**65431100**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
65435122	**Good identification**						**65435122**
Parabacteroides distasonis		100.0	0.37	NIT 1			
65435322	**Good identification**						**65435322**
Parabacteroides distasonis		100.0	0.46	NIT 1			
65435332	**Good identification**						**65435332**
Parabacteroides distasonis		100.0	0.41	NIT 1			
65435722	**Good identification**						**65435722**
Parabacteroides distasonis		100.0	0.42	NIT 1			
65435732	**Good identification**						**65435732**
Parabacteroides distasonis		100.0	0.37	NIT 1			
65450100	**Good identification**						**65450100**
Fusobacterium mortiferum		100.0	0.37	NIT 1			
65451100	**Good identification**						**65451100**
Fusobacterium mortiferum		100.0	0.40	NIT 1			
65475322	**Good identification**						**65475322**
Parabacteroides distasonis		100.0	0.41	NIT 1			
65475332	**Good identification**						**65475332**
Parabacteroides distasonis		100.0	0.36	NIT 1			
65475722	**Good identification**						**65475722**
Parabacteroides distasonis		100.0	0.37	NIT 1			
65511102	**Acceptable identification**						**65511102**
					GRA	CAT	
Capnocytophaga ochracea		93.91	0.50	SUC 99	1	90	
Actinomyces viscosus		4.42	0.28	NAG 90	99	76	
Clostridium perfringens		0.64	0.16		99	1	
65511120	**Low discrimination**						**65511120**
					GRA	CAT	
Capnocytophaga ochracea		63.17	0.50	SUC 99	1	90	
Clostridium perfringens		29.69	0.47	BGL 90	99	1	
Prevotella melaninogenica		6.37	0.29		1	1	
65511122	**Very good identification**						**65511122**
					GRA	CAT	
Capnocytophaga ochracea		95.89	0.66	SUC 99	1	90	
Clostridium perfringens		2.36	0.41		99	1	
Prevotella oralis		1.38	0.34		1	1	
65511132	**Good identification**						**65511132**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		99.48	0.50	SUC 99	1	90	
Clostridium perfringens		0.22	0.07	MLZ 10	99	1	
Parabacteroides distasonis		0.17	0.15		1	76	
65511162	**Good identification**						**65511162**
					GRA	CAT	
Capnocytophaga ochracea		95.84	0.50	SUC 99	1	90	
Clostridium perfringens		4.04	0.29	SOR 10	99	1	
Prevotella oralis		0.13	0.00		1	1	
65511502	**Good identification**						**65511502**
					GRA	CAT	
Capnocytophaga ochracea		99.18	0.37	SUC 99	1	90	
Bifidobacterium longum subsp. longum		0.82	0.09	NAG 90	99	1	
65511520	**Acceptable identification**						**65511520**
					GRA	CAT	
Capnocytophaga ochracea		92.90	0.37	SUC 99	1	90	
Bifidobacterium longum subsp. longum		3.07	0.20	BGL 90	99	1	
Clostridium perfringens		2.50	0.13		99	1	
65511522	**Good identification**						**65511522**
					GRA	CAT	
Capnocytophaga ochracea		99.29	0.53	SUC 99	1	90	
Bifidobacterium longum subsp. longum		0.37	0.20		99	1	
Clostridium perfringens		0.14	0.07		99	1	
65511532	**Acceptable identification**						**65511532**
					GRA	CAT	
Capnocytophaga ochracea		94.14	0.37	SUC 99	1	90	
Bifidobacterium longum subsp. longum		5.30	0.24	MLZ 10	99	1	
Parabacteroides distasonis		0.56	0.11		1	76	
65511562	**Good identification**						**65511562**
Capnocytophaga ochracea		100.0	0.37	SUC 99			
65515102	**Good identification**						**65515102**
Capnocytophaga ochracea		96.25	0.39	SUC 99			
Bacteroides fragilis		3.75	0.06	NAG 90			
65515120	**Low discrimination**						**65515120**
					CAT		
Capnocytophaga ochracea		79.74	0.39	SUC 99	90		
Bacteroides fragilis		16.20	0.18	BGL 90	99		
Bacteroides vulgatus		2.23	0.07		3		
65515122	**Low discrimination**						**65515122**
Capnocytophaga ochracea		68.96	0.56	SUC 99			
Bacteroides fragilis		29.57	0.40				
Parabacteroides distasonis		0.88	0.36				
65515132	**Acceptable identification**						**65515132**
Capnocytophaga ochracea		90.87	0.39	SUC 99			
Parabacteroides distasonis		5.59	0.31	MLZ 10			
Bacteroides fragilis		3.54	0.06				
65515162	**Good identification**						**65515162**
Capnocytophaga ochracea		96.25	0.39	SUC 99			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides fragilis		3.75	0.06	SOR 10			
65515322	**Low discrimination**						**65515322**
Parabacteroides distasonis		68.55	0.44	NIT 1			
Capnocytophaga ochracea		18.18	0.21				
Bacteroides fragilis		7.80	0.06				
65515332	**Acceptable identification**						**65515332**
Parabacteroides distasonis		94.81	0.39	NIT 1			
Capnocytophaga ochracea		5.19	0.05				
65515522	**Low discrimination**						**65515522**
					GRA	CAT	
Capnocytophaga ochracea		89.68	0.43	SUC 99	1	90	
Parabacteroides distasonis		3.95	0.32		1	76	
Bifidobacterium longum subsp. longum		3.95	0.28		99	1	
65515720	**Good identification**						**65515720**
Bacteroides vulgatus		100.0	0.42	NIT 1			
65515722	**Low discrimination**						**65515722**
					CAT		
Parabacteroides distasonis		80.84	0.40	NIT 1	76		
Bacteroides vulgatus		12.98	0.08		3		
Capnocytophaga ochracea		6.17	0.09		90		
65515732	**Good identification**						**65515732**
Parabacteroides distasonis		100.0	0.35	NIT 1			
65531020	**Good identification**						**65531020**
Clostridium perfringens		99.14	0.37	SUC 91			
Actinomyces naeslundii		0.86	0.24	LAC 88			
65531120	**Good identification**						**65531120**
					GRA	CAT	
Clostridium perfringens		97.14	0.52	SUC 91	99	1	
Clostridium septicum		1.30	0.13		99	1	
Capnocytophaga ochracea		1.12	0.15		1	90	
65531122	**Low discrimination**						**65531122**
					GRA	CAT	
Clostridium perfringens		77.90	0.46	SUC 91	99	1	
Capnocytophaga ochracea		17.24	0.32		1	90	
Parabacteroides distasonis		2.74	0.30		1	76	
65531160	**Good identification**						**65531160**
					GRA	CAT	
Clostridium perfringens		99.33	0.39	SUC 91	99	1	
Capnocytophaga ochracea		0.67	-.01		1	90	
65531322	**Low discrimination**						**65531322**
					GRA		
Parabacteroides distasonis		83.46	0.39	NIT 1	1		
Clostridium perfringens		7.98	0.12	XYL 90	99		
Prevotella oralis		6.79	0.07		1		
65531722	**Good identification**						**65531722**
Parabacteroides distasonis		100.0	0.35	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
65533100	**Low discrimination**						**65533100**
Actinomyces naeslundii		66.60	0.37	ESL 12			
Propionibacterium propionicum		33.40	0.21	SUC 99			
65533101	**Good identification**						**65533101**
Actinomyces naeslundii		100.0	0.40	ESL 12			
65533102	**Low discrimination**						**65533102**
Actinomyces naeslundii		57.07	0.37	ESL 12			
Propionibacterium propionicum		42.93	0.24	SUC 99			
65533103	**Good identification**						**65533103**
Actinomyces naeslundii		100.0	0.40	ESL 12			
65533121	**Good identification**						**65533121**
Actinomyces naeslundii		100.0	0.40	ESL 12			
65533122	**Low discrimination**						**65533122**
					GRA	CAT	
Actinomyces naeslundii		52.97	0.37	ESL 12	99	25	
Clostridium perfringens		38.50	0.12	SUC 99	99	1	
Capnocytophaga ochracea		8.52	-.03		1	90	
65533123	**Good identification**						**65533123**
Actinomyces naeslundii		100.0	0.40	ESL 12			
65535122	**Low discrimination**						**65535122**
Parabacteroides distasonis		77.56	0.47	NIT 1			
Capnocytophaga ochracea		13.55	0.21				
Bacteroides fragilis		5.81	0.06				
65535132	**Good identification**						**65535132**
Parabacteroides distasonis		96.52	0.42	NIT 1			
Capnocytophaga ochracea		3.48	0.05				
65535322	**Good identification**						**65535322**
Parabacteroides distasonis		100.0	0.55	NIT 1			
65535332	**Good identification**						**65535332**
Parabacteroides distasonis		100.0	0.50	NIT 1			
65535522	**Acceptable identification**						**65535522**
					GRA	CAT	
Parabacteroides distasonis		90.49	0.43	NIT 1	1	76	
Bifidobacterium longum subsp. longum		4.96	0.17		99	1	
Capnocytophaga ochracea		4.55	0.09		1	90	
65535532	**Low discrimination**						**65535532**
					GRA		
Parabacteroides distasonis		85.22	0.39	NIT 1	1		
Bifidobacterium longum subsp. longum		14.78	0.21		99		
65535722	**Good identification**						**65535722**
Parabacteroides distasonis		99.41	0.51	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		0.59	-.03				
65535732	**Good identification**						**65535732**
Parabacteroides distasonis		98.86	0.47	NIT 1			
Clostridium clostridioforme		1.14	0.08				
65551102	**Low discrimination**						**65551102**
					GRA	CAT	
Capnocytophaga ochracea		86.63	0.39	SUC 99	1	90	
Prevotella oralis		8.34	0.22	NAG 90	1	1	
Actinomyces viscosus		4.34	0.18		99	76	
65551120	**Low discrimination**						**65551120**
					GRA	CAT	
Capnocytophaga ochracea		77.97	0.39	SUC 99	1	90	
Prevotella oralis		7.51	0.22	BGL 90	1	1	
Clostridium perfringens		6.11	0.23		99	1	
65551122	**Low discrimination**						**65551122**
					GRA	CAT	
Capnocytophaga ochracea		82.52	0.56	SUC 99	1	90	
Prevotella oralis		16.78	0.43		1	1	
Clostridium perfringens		0.34	0.18		99	1	
65551132	**Good identification**						**65551132**
					CAT		
Capnocytophaga ochracea		97.83	0.39	SUC 99	90		
Prevotella oralis		1.81	0.09	MLZ 10	1		
Parabacteroides distasonis		0.36	0.10		76		
65551162	**Good identification**						**65551162**
					GRA	CAT	
Capnocytophaga ochracea		97.51	0.39	SUC 99	1	90	
Prevotella oralis		1.80	0.09	SOR 10	1	1	
Clostridium perfringens		0.69	0.05		99	1	
65551322	**Low discrimination**						**65551322**
					CAT		
Prevotella oralis		87.56	0.37	NIT 1	1		
Capnocytophaga ochracea		10.15	0.21	SUC 92	90		
Parabacteroides distasonis		2.29	0.23		76		
65551522	**Good identification**						**65551522**
					CAT		
Capnocytophaga ochracea		98.59	0.43	SUC 99	90		
Prevotella oralis		1.15	0.09		1		
Parabacteroides distasonis		0.26	0.11		76		
65553101	**Good identification**						**65553101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
65553103	**Good identification**						**65553103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
65553121	**Good identification**						**65553121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
65553123	**Good identification**						**65553123**
Actinomyces naeslundii		100.0	0.37	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
65555122	**Low discrimination**					**65555122**
					CAT	
Capnocytophaga ochracea		53.68	0.45	SUC 99	90	
Bacteroides fragilis		39.46	0.34		99	
Prevotella oralis		5.40	0.28		1	
65555322	**Low discrimination**					**65555322**
					CAT	
Parabacteroides distasonis		57.51	0.39	NIT 1	76	
Prevotella oralis		30.20	0.22		1	
Capnocytophaga ochracea		7.08	0.11		90	
65555722	**Good identification**					**65555722**
Parabacteroides distasonis		96.58	0.35	NIT 1		
Capnocytophaga ochracea		3.43	-.02			
65557102	**Good identification**					**65557102**
Actinomyces israelii		100.0	0.38	SUC 98		
65557502	**Low discrimination**					**65557502**
Actinomyces israelii		87.13	0.36	SUC 98		
Bifidobacterium adolescentis		12.87	0.16	SAL 92		
65571101	**Good identification**					**65571101**
Actinomyces naeslundii		100.0	0.37	ESL 12		
65571103	**Good identification**					**65571103**
Actinomyces naeslundii		100.0	0.37	ESL 12		
65571121	**Good identification**					**65571121**
Actinomyces naeslundii		100.0	0.37	ESL 12		
65571123	**Good identification**					**65571123**
Actinomyces naeslundii		100.0	0.37	ESL 12		
65573100	**Low discrimination**					**65573100**
Actinomyces naeslundii		59.70	0.39	ESL 12		
Actinomyces israelii		40.30	0.27	SUC 99		
65573101	**Good identification**					**65573101**
Actinomyces naeslundii		100.0	0.42	ESL 12		
65573103	**Good identification**					**65573103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
65573120	**Good identification**					**65573120**
Actinomyces naeslundii		100.0	0.39	ESL 12		
65573121	**Good identification**					**65573121**
Actinomyces naeslundii		100.0	0.42	ESL 12		
65573122	**Good identification**					**65573122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	ESL 12		
65573123	**Good identification**					**65573123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
65575122	**Low discrimination**					**65575122**
Parabacteroides distasonis		82.89	0.42	NIT 1		
Capnocytophaga ochracea		6.72	0.11			
Bacteroides fragilis		4.94	-.01			
65575132	**Good identification**					**65575132**
Parabacteroides distasonis		100.0	0.38	NIT 1		
65575322	**Good identification**					**65575322**
Parabacteroides distasonis		99.27	0.50	NIT 1		
Prevotella oralis		0.73	0.01			
65575332	**Good identification**					**65575332**
Parabacteroides distasonis		100.0	0.46	NIT 1		
65575522	**Good identification**					**65575522**
Parabacteroides distasonis		97.72	0.39	NIT 1		
Capnocytophaga ochracea		2.28	-.02			
65575722	**Good identification**					**65575722**
Parabacteroides distasonis		96.91	0.47	NIT 1		
Bacteroides thetaiotaomicron		2.26	0.03			
Clostridium clostridioforme		0.82	0.06			
65575732	**Acceptable identification**					**65575732**
Parabacteroides distasonis		91.85	0.42	NIT 1		
Clostridium clostridioforme		6.60	0.17			
Bacteroides thetaiotaomicron		1.55	-.04			
65577002	**Good identification**					**65577002**
Actinomyces israelii		100.0	0.37	SUC 98		
65577100	**Good identification**					**65577100**
Actinomyces israelii		100.0	0.45	SUC 98		
65577102	**Good identification**					**65577102**
Actinomyces israelii		100.0	0.48	SUC 98		
65577140	**Good identification**					**65577140**
Actinomyces israelii		100.0	0.38	SUC 98		
65577142	**Good identification**					**65577142**
Actinomyces israelii		100.0	0.41	SUC 98		
65577402	**Good identification**					**65577402**
Actinomyces israelii		100.0	0.36	SUC 98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
65577500	**Low discrimination**					**65577500**
					GRA	
Actinomyces israelii		61.35	0.43	SUC 98	99	
Mitsuokella multacida		38.65	0.27	SAL 92	1	
65577502	**Acceptable identification**					**65577502**
					GRA	
Actinomyces israelii		93.21	0.46	SUC 98	99	
Mitsuokella multacida		4.35	0.11	SAL 92	1	
Bifidobacterium adolescentis		2.44	0.13		99	
65577542	**Low discrimination**					**65577542**
					GRA	
Actinomyces israelii		89.29	0.39	SUC 98	99	
Mitsuokella multacida		10.71	0.11	SAL 92	1	
65610000	**Good identification**					**65610000**
Fusobacterium mortiferum		100.0	0.44	NIT 1		
65610100	**Good identification**					**65610100**
Fusobacterium mortiferum		100.0	0.50	NIT 1		
65611000	**Good identification**					**65611000**
Fusobacterium mortiferum		100.0	0.47	NIT 1		
65611100	**Good identification**					**65611100**
Fusobacterium mortiferum		100.0	0.53	NIT 1		
65611102	**Good identification**					**65611102**
					GRA	
Fusobacterium mortiferum		96.99	0.37	NIT 1	1	
Bifidobacterium longum subsp. infantis		3.01	-.07	BGL 10	99	
65630100	**Good identification**					**65630100**
Fusobacterium mortiferum		100.0	0.41	NIT 1		
65631000	**Good identification**					**65631000**
Fusobacterium mortiferum		100.0	0.38	NIT 1		
65631100	**Good identification**					**65631100**
Fusobacterium mortiferum		100.0	0.44	NIT 1		
65635122	**Good identification**					**65635122**
Parabacteroides distasonis		100.0	0.38	NIT 1		
65635322	**Good identification**					**65635322**
Parabacteroides distasonis		100.0	0.46	NIT 1		
65635332	**Good identification**					**65635332**
Parabacteroides distasonis		100.0	0.42	NIT 1		
65635722	**Good identification**					**65635722**
Parabacteroides distasonis		100.0	0.43	NIT 1		
65635732	**Good identification**					**65635732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		100.0	0.38	NIT 1			
65650000	**Good identification**						**65650000**
Fusobacterium mortiferum		100.0	0.36	NIT 1			
65650100	**Good identification**						**65650100**
Fusobacterium mortiferum		100.0	0.42	NIT 1			
65651000	**Good identification**						**65651000**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
65651100	**Good identification**						**65651100**
Fusobacterium mortiferum		100.0	0.45	NIT 1			
65671100	**Good identification**						**65671100**
					GRA		
Fusobacterium mortiferum		96.54	0.36	NIT 1	1		
Actinomyces naeslundii		3.46	0.23		99		
65675322	**Good identification**						**65675322**
Parabacteroides distasonis		100.0	0.42	NIT 1			
65675332	**Good identification**						**65675332**
Parabacteroides distasonis		100.0	0.37	NIT 1			
65675722	**Good identification**						**65675722**
Parabacteroides distasonis		100.0	0.38	NIT 1			
65711100	**Low discrimination**						**65711100**
					GRA		
Fusobacterium mortiferum		62.34	0.37	NIT 1	1		
Bifidobacterium longum subsp. infantis		17.02	0.10	MLT 11	99		
Actinomyces naeslundii		10.95	0.36		99		
65711101	**Good identification**						**65711101**
Actinomyces naeslundii		100.0	0.39	ESL 12			
65711103	**Low discrimination**						**65711103**
Actinomyces naeslundii		61.59	0.39	ESL 12			
Actinomyces viscosus		31.90	0.10	SUC 99			
Bifidobacterium longum subsp. infantis		6.52	-.07				
65711120	**Low discrimination**						**65711120**
					SPO		
Clostridium perfringens		66.04	0.37	SUC 91	84		
Clostridium septicum		8.76	0.15		99		
Bifidobacterium longum subsp. infantis		8.35	0.15		1		
65711121	**Low discrimination**						**65711121**
Actinomyces naeslundii		87.16	0.39	ESL 12			
Clostridium perfringens		12.84	0.03	SUC 99			
65711122	**Low discrimination**						**65711122**
					GRA	CAT	
Prevotella oralis		42.91	0.38	NIT 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis		28.09	0.32	SUC 92	99	1	
Capnocytophaga ochracea		17.01	0.32		1	90	
65711123	**Low discrimination**						**65711123**
					GRA		
Actinomyces naeslundii		62.90	0.39	ESL 12	99		
Prevotella oralis		18.09	0.04	SUC 99	1		
Bifidobacterium longum subsp. infantis		11.84	-.03		99		
65713001	**Good identification**						**65713001**
Actinomyces naeslundii		100.0	0.36	ESL 12			
65713003	**Good identification**						**65713003**
Actinomyces naeslundii		100.0	0.36	ESL 12			
65713021	**Good identification**						**65713021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
65713023	**Good identification**						**65713023**
Actinomyces naeslundii		100.0	0.36	ESL 12			
65713100	**Low discrimination**						**65713100**
Actinomyces naeslundii		89.87	0.41	ESL 12			
Actinomyces israelii		10.13	0.16	SUC 99			
65713101	**Good identification**						**65713101**
Actinomyces naeslundii		100.0	0.44	ESL 12			
65713102	**Low discrimination**						**65713102**
Actinomyces naeslundii		79.94	0.41	ESL 12			
Actinomyces israelii		13.52	0.19	SUC 99			
Bifidobacterium longum subsp. infantis		6.54	-.07				
65713103	**Good identification**						**65713103**
Actinomyces naeslundii		100.0	0.44	ESL 12			
65713120	**Low discrimination**						**65713120**
Actinomyces naeslundii		89.78	0.41	ESL 12			
Clostridium perfringens		10.22	0.03	SUC 99			
65713121	**Good identification**						**65713121**
Actinomyces naeslundii		100.0	0.44	ESL 12			
65713122	**Low discrimination**						**65713122**
					GRA		
Actinomyces naeslundii		68.69	0.41	ESL 12	99		
Prevotella oralis		15.27	0.04	SUC 99	1		
Bifidobacterium longum subsp. infantis		9.99	-.03		99		
65713123	**Good identification**						**65713123**
Actinomyces naeslundii		100.0	0.44	ESL 12			
65715322	**Low discrimination**						**65715322**
					GRA		
Parabacteroides distasonis		86.69	0.44	NIT 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		11.35	0.16		1	
Bifidobacterium longum subsp. infantis		1.16	-.04		99	
65715332	**Good identification**					**65715332**
Parabacteroides distasonis		100.0	0.40	NIT 1		
65715722	**Good identification**					**65715722**
Parabacteroides distasonis		96.43	0.41	NIT 1		
Bacteroides vulgatus		3.57	-.02			
65715732	**Good identification**					**65715732**
Parabacteroides distasonis		95.36	0.36	NIT 1		
Clostridium clostridioforme		4.64	0.08			
65717102	**Acceptable identification**					**65717102**
Actinomyces israelii		94.73	0.37	SUC 98		
Bifidobacterium adolescentis		5.27	0.09	CEL 93		
65717502	**Low discrimination**					**65717502**
Actinomyces israelii		68.73	0.35	SUC 98		
Bifidobacterium adolescentis		31.27	0.23	CEL 93		
65731001	**Good identification**					**65731001**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65731003	**Good identification**					**65731003**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65731021	**Good identification**					**65731021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65731023	**Good identification**					**65731023**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65731100	**Low discrimination**					**65731100**
					GRA	
Actinomyces naeslundii		39.62	0.41	ESL 12	99	
Fusobacterium mortiferum		34.73	0.28	SUC 99	1	
Clostridium perfringens		25.65	0.16		99	
65731101	**Good identification**					**65731101**
Actinomyces naeslundii		100.0	0.44	ESL 12		
65731102	**Low discrimination**					**65731102**
Actinomyces naeslundii		59.30	0.41	ESL 12		
Clostridium perfringens		18.06	0.10	SUC 99		
Actinomyces israelii		6.66	0.16			
65731103	**Good identification**					**65731103**
Actinomyces naeslundii		100.0	0.44	ESL 12		
65731120	**Low discrimination**					**65731120**
					SPO	
Clostridium perfringens		79.24	0.41	SUC 91	84	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium septicum		16.98	0.23		99	
Actinomyces naeslundii		3.79	0.41		1	
65731121	**Low discrimination**					**65731121**
Actinomyces naeslundii		87.65	0.44	ESL 12		
Clostridium perfringens		12.36	0.07	SUC 99		
65731122	**Low discrimination**					**65731122**
					GRA	
Clostridium perfringens		69.95	0.36	SUC 91	99	
Parabacteroides distasonis		10.67	0.31		1	
Prevotella oralis		9.97	0.18		1	
65731123	**Acceptable identification**					**65731123**
Actinomyces naeslundii		93.78	0.44	ESL 12		
Clostridium perfringens		6.22	0.02	SUC 99		
65731322	**Low discrimination**					**65731322**
					GRA	
Parabacteroides distasonis		86.54	0.39	NIT 1	1	
Prevotella oralis		11.56	0.12	XYL 90	1	
Clostridium perfringens		1.91	0.02		99	
65731722	**Good identification**					**65731722**
Parabacteroides distasonis		100.0	0.36	NIT 1		
65733000	**Low discrimination**					**65733000**
Actinomyces naeslundii		86.55	0.38	ESL 12		
Actinomyces israelii		13.45	0.15	SUC 99		
65733001	**Good identification**					**65733001**
Actinomyces naeslundii		100.0	0.41	ESL 12		
65733002	**Low discrimination**					**65733002**
Actinomyces naeslundii		81.09	0.38	ESL 12		
Actinomyces israelii		18.91	0.18	SUC 99		
65733003	**Good identification**					**65733003**
Actinomyces naeslundii		100.0	0.41	ESL 12		
65733020	**Good identification**					**65733020**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65733021	**Good identification**					**65733021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
65733022	**Good identification**					**65733022**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65733023	**Good identification**					**65733023**
Actinomyces naeslundii		100.0	0.41	ESL 12		
65733100	**Low discrimination**					**65733100**
Actinomyces naeslundii		82.07	0.46	ESL 12		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		17.93	0.25	SUC 99		
65733101	**Good identification**					**65733101**
Actinomyces naeslundii		100.0	0.49	ESL 12		
65733102	**Low discrimination**					**65733102**
Actinomyces naeslundii		75.32	0.46	ESL 12		
Actinomyces israelii		24.68	0.28	SUC 99		
65733103	**Good identification**					**65733103**
Actinomyces naeslundii		100.0	0.49	ESL 12		
65733120	**Acceptable identification**					**65733120**
Actinomyces naeslundii		90.18	0.46	ESL 12		
Clostridium perfringens		9.82	0.07	SUC 99		
65733121	**Good identification**					**65733121**
Actinomyces naeslundii		100.0	0.49	ESL 12		
65733122	**Good identification**					**65733122**
Actinomyces naeslundii		95.13	0.46	ESL 12		
Clostridium perfringens		4.88	0.02	SUC 99		
65733123	**Good identification**					**65733123**
Actinomyces naeslundii		100.0	0.49	ESL 12		
65733141	**Good identification**					**65733141**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65733143	**Good identification**					**65733143**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65733161	**Good identification**					**65733161**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65733163	**Good identification**					**65733163**
Actinomyces naeslundii		100.0	0.36	ESL 12		
65735122	**Good identification**					**65735122**
					GRA	
Parabacteroides distasonis		96.61	0.47	NIT 1	1	
Prevotella oralis		1.24	0.02		1	
Bifidobacterium longum subsp. infantis		0.85	-.04		99	
65735132	**Good identification**					**65735132**
Parabacteroides distasonis		100.0	0.43	NIT 1		
65735322	**Good identification**					**65735322**
Parabacteroides distasonis		100.0	0.56	NIT 1		
65735332	**Good identification**					**65735332**
Parabacteroides distasonis		100.0	0.51	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
65735522	**Good identification**					**65735522**
Parabacteroides distasonis		100.0	0.44	NIT 1		
65735532	**Good identification**					**65735532**
Parabacteroides distasonis		96.90	0.39	NIT 1		
Clostridium clostridioforme		3.10	0.08			
65735722	**Good identification**					**65735722**
Parabacteroides distasonis		99.08	0.52	NIT 1		
Clostridium clostridioforme		0.92	0.11			
65735732	**Acceptable identification**					**65735732**
Parabacteroides distasonis		92.74	0.47	NIT 1		
Clostridium clostridioforme		7.26	0.23			
65737002	**Good identification**					**65737002**
Actinomyces israelii		100.0	0.36	SUC 98		
65737100	**Good identification**					**65737100**
Actinomyces israelii		100.0	0.44	SUC 98		
65737102	**Good identification**					**65737102**
Actinomyces israelii		100.0	0.47	SUC 98		
65737140	**Good identification**					**65737140**
Actinomyces israelii		100.0	0.37	SUC 98		
65737142	**Good identification**					**65737142**
Actinomyces israelii		100.0	0.40	SUC 98		
65737500	**Low discrimination**					**65737500**
					GRA	
Actinomyces israelii		57.88	0.42	SUC 98	99	
Mitsuokella multacida		42.12	0.27	CEL 93	1	
65737502	**Low discrimination**					**65737502**
					GRA	
Actinomyces israelii		88.14	0.45	SUC 98	99	
Bifidobacterium adolescentis		7.11	0.20	CEL 93	99	
Mitsuokella multacida		4.75	0.11		1	
65737542	**Low discrimination**					**65737542**
					GRA	
Actinomyces israelii		87.83	0.38	SUC 98	99	
Mitsuokella multacida		12.17	0.11	CEL 93	1	
65751101	**Good identification**					**65751101**
Actinomyces naeslundii		100.0	0.41	ESL 12		
65751103	**Acceptable identification**					**65751103**
Actinomyces naeslundii		90.59	0.41	ESL 12		
Actinomyces viscosus		9.41	0.00	SUC 99		
65751121	**Low discrimination**					**65751121**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Actinomyces naeslundii		88.15	0.41	ESL 12	1			
Clostridium paraputrificum		11.85	-.06	SUC 99	99			
65751122	**Low discrimination**							**65751122**
					GRA	CAT	SPO	
Prevotella oralis		86.99	0.48	NIT 1	1	1	1	
Clostridium paraputrificum		6.52	0.22	SUC 92	99	1	99	
Capnocytophaga ochracea		2.43	0.21		1	90	1	
65751123	**Low discrimination**							**65751123**
					GRA			
Actinomyces naeslundii		56.53	0.41	ESL 12	99			
Prevotella oralis		43.48	0.14	SUC 99	1			
65751322	**Good identification**							**65751322**
					SPO			
Prevotella oralis		98.15	0.41	NIT 1	1			
Parabacteroides distasonis		1.56	0.23	SUC 92	1			
Clostridium ramosum		0.28	-.02		92			
65753000	**Low discrimination**							**65753000**
Actinomyces naeslundii		55.43	0.35	ESL 12				
Actinomyces israelii		44.57	0.25	SUC 99				
65753001	**Good identification**							**65753001**
Actinomyces naeslundii		100.0	0.38	ESL 12				
65753003	**Good identification**							**65753003**
Actinomyces naeslundii		100.0	0.38	ESL 12				
65753020	**Good identification**							**65753020**
Actinomyces naeslundii		100.0	0.35	ESL 12				
65753021	**Good identification**							**65753021**
Actinomyces naeslundii		100.0	0.38	ESL 12				
65753022	**Good identification**							**65753022**
Actinomyces naeslundii		100.0	0.35	ESL 12				
65753023	**Good identification**							**65753023**
Actinomyces naeslundii		100.0	0.38	ESL 12				
65753101	**Good identification**							**65753101**
Actinomyces naeslundii		100.0	0.46	ESL 12				
65753102	**Low discrimination**							**65753102**
Actinomyces israelii		47.69	0.38	SUC 98				
Actinomyces naeslundii		28.14	0.43	XYL 92				
Bifidobacterium breve		24.17	0.27					
65753103	**Good identification**							**65753103**
Actinomyces naeslundii		100.0	0.46	ESL 12				
65753120	**Low discrimination**							**65753120**
					SPO			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		71.18	0.43	ESL 12	1		
Bifidobacterium breve		10.75	0.15	SUC 99	1		
Clostridium paraputrificum		7.39	-.06		99		
65753121	**Good identification**						**65753121**
Actinomyces naeslundii		100.0	0.46	ESL 12			
65753123	**Good identification**						**65753123**
Actinomyces naeslundii		100.0	0.46	ESL 12			
65753502	**Low discrimination**						**65753502**
Actinomyces israelii		88.89	0.36	SUC 98			
Bifidobacterium adolescentis		11.12	0.15	XYL 92			
65755100	**Good identification**						**65755100**
					SPO		
Actinomyces israelii		96.24	0.40	SUC 98	1		
Clostridium butyricum		3.76	-.05		99		
65755102	**Low discrimination**						**65755102**
					GRA		
Actinomyces israelii		85.26	0.43	SUC 98	99		
Prevotella oralis		9.49	0.10		1		
Bifidobacterium adolescentis		5.25	0.16		99		
65755142	**Good identification**						**65755142**
Actinomyces israelii		100.0	0.36	SUC 98			
65755322	**Low discrimination**						**65755322**
Parabacteroides distasonis		53.71	0.40	NIT 1			
Prevotella oralis		46.29	0.26				
65755332	**Good identification**						**65755332**
Parabacteroides distasonis		100.0	0.35	NIT 1			
65755500	**Low discrimination**						**65755500**
					GRA	SPO	
Actinomyces israelii		51.82	0.39	SUC 98	99	1	
Mitsuokella multacida		25.93	0.21		1	1	
Clostridium butyricum		22.25	0.11		99	99	
65755502	**Low discrimination**						**65755502**
					SPO		
Actinomyces israelii		60.52	0.42	SUC 98	1		
Bifidobacterium adolescentis		30.50	0.30		1		
Clostridium butyricum		6.74	0.04		99		
65755722	**Low discrimination**						**65755722**
Parabacteroides distasonis		84.87	0.36	NIT 1			
Bacteroides uniformis		12.10	0.03				
Clostridium clostridioforme		3.03	0.06				
65757000	**Good identification**						**65757000**
Actinomyces israelii		100.0	0.43	SUC 98			
65757002	**Good identification**						**65757002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.46	SUC 98			
65757040	**Good identification**						**65757040**
Actinomyces israelii		100.0	0.36	SUC 98			
65757042	**Good identification**						**65757042**
Actinomyces israelii		100.0	0.39	SUC 98			
65757100	**Good identification**						**65757100**
Actinomyces israelii		100.0	0.53	SUC 98			
65757102	**Good identification**						**65757102**
Actinomyces israelii		99.12	0.56	SUC 98			
Bifidobacterium adolescentis		0.88	0.15				
65757110	**Good identification**						**65757110**
Actinomyces israelii		100.0	0.39	SUC 98			
65757112	**Acceptable identification**						**65757112**
Actinomyces israelii		93.22	0.42	SUC 98			
Bifidobacterium adolescentis		6.78	0.16	MLZ 13			
65757122	**Acceptable identification**						**65757122**
					GRA		
Actinomyces israelii		94.43	0.36	SUC 98	99		
Prevotella oralis		5.57	-.02	NAG 6	1		
65757140	**Good identification**						**65757140**
Actinomyces israelii		100.0	0.46	SUC 98			
65757142	**Good identification**						**65757142**
Actinomyces israelii		100.0	0.49	SUC 98			
65757302	**Good identification**						**65757302**
Actinomyces israelii		100.0	0.37	SUC 98			
65757400	**Good identification**						**65757400**
Actinomyces israelii		100.0	0.41	SUC 98			
65757402	**Good identification**						**65757402**
Actinomyces israelii		100.0	0.44	SUC 98			
65757442	**Good identification**						**65757442**
Actinomyces israelii		100.0	0.37	SUC 98			
65757500	**Low discrimination**						**65757500**
					GRA	SPO	
Actinomyces israelii		82.34	0.51	SUC 98	99	1	
Mitsuokella multacida		16.97	0.27		1	1	
Clostridium butyricum		0.69	-.05		99	99	
65757502	**Acceptable identification**						**65757502**
					GRA		
Actinomyces israelii		91.91	0.55	SUC 98	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		6.69	0.29		99	
Mitsuokella multacida		1.40	0.11		1	
65757510	**Low discrimination**					**65757510**
					GRA	
Actinomyces israelii		68.51	0.37	SUC 98	99	
Mitsuokella multacida		31.49	0.19	MLZ 13	1	
65757512	**Low discrimination**					**65757512**
Actinomyces israelii		59.67	0.40	SUC 98		
Bifidobacterium adolescentis		35.52	0.30	MLZ 13		
Bifidobacterium dentium		2.78	-.07			
65757540	**Low discrimination**					**65757540**
					GRA	
Actinomyces israelii		65.37	0.44	SUC 98	99	
Mitsuokella multacida		34.64	0.27		1	
65757542	**Acceptable identification**					**65757542**
					GRA	
Actinomyces israelii		94.13	0.47	SUC 98	99	
Mitsuokella multacida		3.70	0.11		1	
Bifidobacterium adolescentis		2.18	0.13		99	
65757702	**Low discrimination**					**65757702**
					GRA	
Actinomyces israelii		72.64	0.35	SUC 98	99	
Mitsuokella multacida		27.36	0.15	RHA 7	1	
65771001	**Good identification**					**65771001**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65771002	**Low discrimination**					**65771002**
Actinomyces naeslundii		38.09	0.35	ESL 12		
Lactobacillus jensenii		31.41	0.12	SUC 99		
Actinomyces israelii		30.50	0.24			
65771003	**Good identification**					**65771003**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65771020	**Low discrimination**					**65771020**
					SPO	
Actinomyces naeslundii		53.40	0.35	ESL 12	1	
Lactobacillus jensenii		16.51	0.05	SUC 99	1	
Clostridium septicum		16.45	-.03		99	
65771021	**Good identification**					**65771021**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65771022	**Low discrimination**					**65771022**
					GRA	
Actinomyces naeslundii		52.95	0.35	ESL 12	99	
Prevotella oralis		36.13	0.07	SUC 99	1	
Lactobacillus jensenii		10.92	0.02		99	
65771023	**Good identification**					**65771023**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65771100	**Low discrimination**					**65771100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		45.75	0.43	ESL 12	99	
Actinomyces israelii		34.32	0.32	SUC 99	99	
Fusobacterium mortiferum		10.08	0.20		1	
65771101	**Good identification**					**65771101**
Actinomyces naeslundii		100.0	0.46	ESL 12		
65771103	**Good identification**					**65771103**
Actinomyces naeslundii		100.0	0.46	ESL 12		
65771112	**Good identification**					**65771112**
					GRA	
Leptotrichia buccalis		99.07	0.41	NIT 10	1	
Bifidobacterium breve		0.72	0.24	SUC 99	99	
Actinomyces israelii		0.21	0.21		99	
65771121	**Acceptable identification**					**65771121**
					SPO	
Actinomyces naeslundii		93.27	0.46	ESL 12	1	
Clostridium septicum		6.73	-.03	SUC 99	99	
65771123	**Good identification**					**65771123**
Actinomyces naeslundii		100.0	0.46	ESL 12		
65773001	**Good identification**					**65773001**
Actinomyces naeslundii		100.0	0.43	ESL 12		
65773002	**Low discrimination**					**65773002**
Actinomyces israelii		70.03	0.37	SUC 98		
Actinomyces naeslundii		29.97	0.40	XYL 92		
65773003	**Good identification**					**65773003**
Actinomyces naeslundii		100.0	0.43	ESL 12		
65773020	**Acceptable identification**					**65773020**
Actinomyces naeslundii		90.96	0.40	ESL 12		
Actinomyces israelii		9.04	0.14	SUC 99		
65773021	**Good identification**					**65773021**
Actinomyces naeslundii		100.0	0.43	ESL 12		
65773022	**Low discrimination**					**65773022**
Actinomyces naeslundii		87.02	0.40	ESL 12		
Actinomyces israelii		12.98	0.17	SUC 99		
65773023	**Good identification**					**65773023**
Actinomyces naeslundii		100.0	0.43	ESL 12		
65773100	**Low discrimination**					**65773100**
Actinomyces israelii		68.65	0.45	SUC 98		
Actinomyces naeslundii		31.35	0.48	XYL 92		
65773101	**Good identification**					**65773101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.51	ESL 12		
65773102	**Low discrimination**					**65773102**
Actinomyces israelii		71.72	0.48	SUC 98		
Actinomyces naeslundii		21.84	0.48	XYL 92		
Bifidobacterium breve		6.44	0.24			
65773103	**Good identification**					**65773103**
Actinomyces naeslundii		97.84	0.51	ESL 12		
Actinomyces israelii		2.17	0.14	SUC 99		
65773120	**Low discrimination**					**65773120**
					SPO	
Actinomyces naeslundii		53.30	0.48	ESL 12	1	
Clostridium ramosum		33.51	0.19	SUC 99	92	
Actinomyces israelii		7.45	0.24		1	
65773121	**Good identification**					**65773121**
Actinomyces naeslundii		100.0	0.51	ESL 12		
65773123	**Good identification**					**65773123**
Actinomyces naeslundii		100.0	0.51	ESL 12		
65773140	**Low discrimination**					**65773140**
Actinomyces israelii		82.83	0.38	SUC 98		
Actinomyces naeslundii		17.17	0.35	XYL 92		
65773141	**Good identification**					**65773141**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65773142	**Low discrimination**					**65773142**
Actinomyces israelii		73.04	0.41	SUC 98		
Bifidobacterium breve		16.87	0.24	XYL 92		
Actinomyces naeslundii		10.09	0.35			
65773143	**Good identification**					**65773143**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65773160	**Low discrimination**					**65773160**
Actinomyces naeslundii		62.43	0.35	ESL 12		
Actinomyces israelii		19.23	0.17	SUC 99		
Bifidobacterium breve		18.35	0.12			
65773161	**Good identification**					**65773161**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65773163	**Good identification**					**65773163**
Actinomyces naeslundii		100.0	0.38	ESL 12		
65773402	**Good identification**					**65773402**
Actinomyces israelii		100.0	0.36	SUC 98		
65773500	**Low discrimination**					**65773500**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		61.35	0.43	SUC 98	99		
Mitsuokella multacida		38.65	0.27	XYL 92	1		
65773502	**Acceptable identification**						**65773502**
					GRA		
Actinomyces israelii		93.56	0.46	SUC 98	99		
Mitsuokella multacida		4.37	0.11	XYL 92	1		
Bifidobacterium adolescentis		2.07	0.12		99		
65773542	**Low discrimination**						**65773542**
					GRA		
Actinomyces israelii		89.29	0.39	SUC 98	99		
Mitsuokella multacida		10.71	0.11	XYL 92	1		
65775000	**Good identification**						**65775000**
Actinomyces israelii		100.0	0.40	SUC 98			
65775002	**Good identification**						**65775002**
Actinomyces israelii		100.0	0.43	SUC 98			
65775042	**Good identification**						**65775042**
Actinomyces israelii		100.0	0.36	SUC 98			
65775100	**Low discrimination**						**65775100**
					GRA	SPO	
Actinomyces israelii		77.26	0.50	SUC 98	99	1	
Clostridium butyricum		12.56	0.15		99	99	
Mitsuokella multacida		8.41	0.21		1	1	
65775102	**Acceptable identification**						**65775102**
					SPO		
Actinomyces israelii		94.24	0.53	SUC 98	1		
Clostridium butyricum		3.97	0.08		99		
Bifidobacterium adolescentis		1.03	0.13		1		
65775110	**Low discrimination**						**65775110**
					GRA	SPO	
Actinomyces israelii		76.92	0.36	SUC 98	99	1	
Mitsuokella multacida		18.68	0.13	MLZ 13	1	1	
Clostridium butyricum		4.40	-.07		99	99	
65775112	**Low discrimination**						**65775112**
					GRA		
Actinomyces israelii		66.64	0.39	SUC 98	99		
Leptotrichia buccalis		27.41	0.07	MLZ 13	1		
Bifidobacterium adolescentis		5.95	0.15		99		
65775122	**Low discrimination**						**65775122**
					GRA		
Parabacteroides distasonis		80.98	0.43	NIT 1	1		
Actinomyces israelii		12.18	0.33		99		
Prevotella oralis		6.84	0.12		1		
65775132	**Acceptable identification**						**65775132**
					GRA		
Parabacteroides distasonis		93.32	0.38	NIT 1	1		
Actinomyces israelii		3.90	0.18		99		
Leptotrichia buccalis		2.79	-.09		1		
65775140	**Low discrimination**						**65775140**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		75.98	0.43	SUC 98	99	1	
Mitsuokella multacida		21.26	0.21		1	1	
Clostridium butyricum		2.76	-.03		99	99	
65775142	**Good identification**						**65775142**
					GRA		
Actinomyces israelii		97.97	0.46	SUC 98	99		
Mitsuokella multacida		2.03	0.04		1		
65775322	**Good identification**						**65775322**
					GRA		
Parabacteroides distasonis		98.44	0.51	NIT 1	1		
Prevotella oralis		1.19	0.05		1		
Actinomyces israelii		0.37	0.13		99		
65775332	**Good identification**						**65775332**
Parabacteroides distasonis		98.13	0.46	NIT 1			
Clostridium clostridioforme		1.87	0.11				
65775400	**Low discrimination**						**65775400**
					GRA	SPO	
Actinomyces israelii		54.64	0.38	SUC 98	99	1	
Mitsuokella multacida		29.08	0.21		1	1	
Clostridium butyricum		16.28	0.08		99	99	
65775402	**Low discrimination**						**65775402**
					GRA	SPO	
Actinomyces israelii		89.55	0.41	SUC 98	99	1	
Clostridium butyricum		6.92	0.01		99	99	
Mitsuokella multacida		3.53	0.04		1	1	
65775500	**Low discrimination**						**65775500**
					GRA	SPO	
Mitsuokella multacida		82.53	0.55	SUC 99	1	1	
Clostridium butyricum		11.21	0.32		99	99	
Actinomyces israelii		6.27	0.48		99	1	
65775501	**Good identification**						**65775501**
					GRA	SPO	
Mitsuokella multacida		98.11	0.39	SUC 99	1	1	
Clostridium butyricum		1.21	-.03	URE 10	99	99	
Actinomyces israelii		0.68	0.14		99	1	
65775502	**Low discrimination**						**65775502**
					GRA	SPO	
Actinomyces israelii		39.43	0.51	SUC 98	99	1	
Mitsuokella multacida		38.47	0.39		1	1	
Clostridium butyricum		18.28	0.25		99	99	
65775510	**Acceptable identification**						**65775510**
					GRA	SPO	
Mitsuokella multacida		94.74	0.47	SUC 99	1	1	
Actinomyces israelii		3.23	0.34		99	1	
Clostridium butyricum		2.03	0.10		99	99	
65775520	**Acceptable identification**						**65775520**
					GRA	SPO	
Mitsuokella multacida		93.76	0.45	SUC 99	1	1	
Clostridium butyricum		4.43	0.14		99	99	
Actinomyces israelii		1.82	0.28		99	1	
65775522	**Low discrimination**						**65775522**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Parabacteroides distasonis		47.21	0.39	NIT 1	1		
Mitsuokella multacida		36.21	0.28		1		
Actinomyces israelii		9.48	0.31		99		
65775530	**Good identification**						**65775530**
					GRA		
Mitsuokella multacida		99.14	0.37	SUC 99	1		
Actinomyces israelii		0.86	0.14		99		
65775540	**Good identification**						**65775540**
					GRA	SPO	
Mitsuokella multacida		96.03	0.55	SUC 99	1	1	
Actinomyces israelii		2.84	0.41		99	1	
Clostridium butyricum		1.13	0.14		99	99	
65775541	**Good identification**						**65775541**
Mitsuokella multacida		100.0	0.39	SUC 99			
65775542	**Low discrimination**						**65775542**
					GRA	SPO	
Mitsuokella multacida		68.91	0.39	SUC 99	1	1	
Actinomyces israelii		27.47	0.44	BGL 10	99	1	
Clostridium butyricum		2.85	0.07		99	99	
65775550	**Good identification**						**65775550**
					COC	GRA	
Mitsuokella multacida		98.35	0.47	SUC 99	1	1	
Actinomyces israelii		1.30	0.27		1	99	
Blautia producta		0.35	-.03		99	99	
65775560	**Good identification**						**65775560**
					GRA	SPO	
Mitsuokella multacida		98.85	0.45	SUC 99	1	1	
Actinomyces israelii		0.75	0.21		99	1	
Clostridium butyricum		0.41	-.05		99	99	
65775570	**Good identification**						**65775570**
Mitsuokella multacida		100.0	0.37	SUC 99			
65775700	**Very good identification**						**65775700**
					GRA	SPO	
Mitsuokella multacida		99.47	0.60	SUC 99	1	1	
Actinomyces israelii		0.31	0.29		99	1	
Clostridium butyricum		0.23	0.06		99	99	
65775701	**Good identification**						**65775701**
Mitsuokella multacida		100.0	0.43	SUC 99			
65775702	**Acceptable identification**						**65775702**
					GRA		
Mitsuokella multacida		92.16	0.43	SUC 99	1		
Actinomyces israelii		3.83	0.32	BGL 10	99		
Clostridium clostridioforme		2.79	0.20		50		
65775710	**Good identification**						**65775710**
					GRA		
Mitsuokella multacida		99.86	0.51	SUC 99	1		
Actinomyces israelii		0.14	0.15		99		
65775711	**Good identification**						**65775711**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.35	SUC 99			
65775712	**Low discrimination**						**65775712**
					GRA		
Mitsuokella multacida		69.81	0.35	SUC 99	1		
Clostridium clostridioforme		28.89	0.32	BGL 10	50		
Actinomyces israelii		1.30	0.18		99		
65775720	**Good identification**						**65775720**
Mitsuokella multacida		99.76	0.49	SUC 99			
Parabacteroides distasonis		0.24	0.13				
65775722	**Low discrimination**						**65775722**
Parabacteroides distasonis		63.91	0.47	NIT 1			
Mitsuokella multacida		30.35	0.33				
Clostridium clostridioforme		3.68	0.20				
65775730	**Good identification**						**65775730**
Mitsuokella multacida		100.0	0.41	SUC 99			
65775732	**Low discrimination**						**65775732**
Parabacteroides distasonis		54.93	0.43	NIT 1			
Clostridium clostridioforme		26.73	0.32				
Mitsuokella multacida		16.15	0.25				
65775740	**Very good identification**						**65775740**
					GRA		
Mitsuokella multacida		99.88	0.60	SUC 99	1		
Actinomyces israelii		0.12	0.22		99		
65775741	**Good identification**						**65775741**
Mitsuokella multacida		100.0	0.43	SUC 99			
65775742	**Good identification**						**65775742**
					GRA		
Mitsuokella multacida		98.41	0.43	SUC 99	1		
Actinomyces israelii		1.59	0.25	BGL 10	99		
65775750	**Good identification**						**65775750**
					COC	GRA	
Mitsuokella multacida		99.81	0.51	SUC 99	1	1	
Blautia producta		0.19	-.03		99	99	
65775751	**Good identification**						**65775751**
Mitsuokella multacida		100.0	0.35	SUC 99			
65775752	**Good identification**						**65775752**
Mitsuokella multacida		100.0	0.35	SUC 99			
65775760	**Good identification**						**65775760**
Mitsuokella multacida		100.0	0.49	SUC 99			
65775770	**Good identification**						**65775770**
Mitsuokella multacida		100.0	0.41	SUC 99			
65777000	**Good identification**						**65777000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.53	SUC 98			
65777002	**Good identification**						**65777002**
Actinomyces israelii		100.0	0.56	SUC 98			
65777010	**Good identification**						**65777010**
Actinomyces israelii		100.0	0.38	SUC 98			
65777012	**Good identification**						**65777012**
Actinomyces israelii		100.0	0.41	SUC 98			
65777022	**Good identification**						**65777022**
Actinomyces israelii		100.0	0.35	SUC 98			
65777040	**Good identification**						**65777040**
Actinomyces israelii		100.0	0.46	SUC 98			
65777042	**Good identification**						**65777042**
Actinomyces israelii		100.0	0.49	SUC 98			
65777100	**Very good identification**						**65777100**
					GRA	SPO	
Actinomyces israelii		95.42	0.63	SUC 98	99	1	
Mitsuokella multacida		4.28	0.27		1	1	
Clostridium butyricum		0.30	-.01		99	99	
65777102	**Very good identification**						**65777102**
					GRA		
Actinomyces israelii		99.51	0.66	SUC 98	99		
Mitsuokella multacida		0.33	0.11		1		
Bifidobacterium adolescentis		0.16	0.12		99		
65777110	**Acceptable identification**						**65777110**
					GRA		
Actinomyces israelii		90.91	0.49	SUC 98	99		
Mitsuokella multacida		9.09	0.19	MLZ 13	1		
65777112	**Good identification**						**65777112**
Actinomyces israelii		97.05	0.52	SUC 98			
Bifidobacterium adolescentis		1.25	0.13	MLZ 13			
Bifidobacterium dentium		0.98	-.07				
65777120	**Low discrimination**						**65777120**
					GRA		
Actinomyces israelii		85.07	0.42	SUC 98	99		
Mitsuokella multacida		14.93	0.17	NAG 6	1		
65777122	**Good identification**						**65777122**
Actinomyces israelii		100.0	0.45	SUC 98			
65777140	**Low discrimination**						**65777140**
					GRA		
Actinomyces israelii		89.67	0.56	SUC 98	99		
Mitsuokella multacida		10.33	0.27		1		
65777142	**Very good identification**						**65777142**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		99.15	0.59	SUC 98	99		
Mitsuokella multacida		0.85	0.11		1		
65777150	**Low discrimination**						**65777150**
					COC	GRA	
Actinomyces israelii		77.18	0.42	SUC 98	1	99	
Mitsuokella multacida		19.84	0.19	MLZ 13	1	1	
Blautia producta		2.98	-.03		99	99	
65777152	**Good identification**						**65777152**
					GRA		
Actinomyces israelii		98.13	0.45	SUC 98	99		
Mitsuokella multacida		1.87	0.03	MLZ 13	1		
65777160	**Low discrimination**						**65777160**
					GRA		
Actinomyces israelii		68.90	0.35	SUC 98	99		
Mitsuokella multacida		31.10	0.17	NAG 6	1		
65777162	**Good identification**						**65777162**
Actinomyces israelii		100.0	0.38	SUC 98			
65777202	**Good identification**						**65777202**
Actinomyces israelii		100.0	0.36	SUC 98			
65777302	**Acceptable identification**						**65777302**
					GRA		
Actinomyces israelii		92.43	0.47	SUC 98	99		
Mitsuokella multacida		7.57	0.15	RHA 7	1		
65777342	**Low discrimination**						**65777342**
					GRA		
Actinomyces israelii		82.60	0.40	SUC 98	99		
Mitsuokella multacida		17.40	0.15	RHA 7	1		
65777400	**Low discrimination**						**65777400**
					GRA		
Actinomyces israelii		82.03	0.51	SUC 98	99		
Mitsuokella multacida		17.97	0.27		1		
65777402	**Good identification**						**65777402**
					GRA		
Actinomyces israelii		98.40	0.54	SUC 98	99		
Mitsuokella multacida		1.60	0.11		1		
65777410	**Low discrimination**						**65777410**
					GRA		
Actinomyces israelii		67.17	0.37	SUC 98	99		
Mitsuokella multacida		32.83	0.19	MLZ 13	1		
65777412	**Acceptable identification**						**65777412**
					GRA		
Actinomyces israelii		92.10	0.40	SUC 98	99		
Bifidobacterium dentium		4.57	-.07	MLZ 13	99		
Mitsuokella multacida		3.34	0.03		1		
65777440	**Low discrimination**						**65777440**
					GRA		
Actinomyces israelii		63.96	0.44	SUC 98	99		
Mitsuokella multacida		36.04	0.27		1		
65777442	**Good identification**						**65777442**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		95.99	0.47	SUC 98	99		
Mitsuokella multacida		4.01	0.11		1		
65777500	**Low discrimination**						**65777500**
					GRA	SPO	
Mitsuokella multacida		83.97	0.61	SUC 99	1	1	
Actinomyces israelii		15.49	0.61		99	1	
Clostridium butyricum		0.54	0.15		99	99	
65777501	**Good identification**						**65777501**
					GRA		
Mitsuokella multacida		98.35	0.45	SUC 99	1		
Actinomyces israelii		1.65	0.27	URE 10	99		
65777502	**Low discrimination**						**65777502**
					GRA		
Actinomyces israelii		70.15	0.64	SUC 98	99		
Mitsuokella multacida		28.18	0.45		1		
Bifidobacterium adolescentis		0.91	0.26		99		
65777510	**Acceptable identification**						**65777510**
					GRA		
Mitsuokella multacida		90.89	0.53	SUC 99	1		
Actinomyces israelii		7.51	0.47		99		
Bifidobacterium dentium		1.37	0.10		99		
65777511	**Good identification**						**65777511**
Mitsuokella multacida		100.0	0.37	SUC 99			
65777520	**Good identification**						**65777520**
					GRA	SPO	
Mitsuokella multacida		95.30	0.51	SUC 99	1	1	
Actinomyces israelii		4.49	0.41		99	1	
Clostridium butyricum		0.21	-.03		99	99	
65777530	**Low discrimination**						**65777530**
					GRA		
Mitsuokella multacida		88.64	0.43	SUC 99	1		
Bifidobacterium dentium		9.49	0.15		99		
Actinomyces israelii		1.87	0.27		99		
65777540	**Good identification**						**65777540**
					COC	GRA	
Mitsuokella multacida		91.83	0.61	SUC 99	1	1	
Actinomyces israelii		6.59	0.54		1	99	
Blautia producta		1.53	0.24		99	99	
65777541	**Good identification**						**65777541**
					GRA		
Mitsuokella multacida		99.35	0.45	SUC 99	1		
Actinomyces israelii		0.65	0.20	URE 10	99		
65777542	**Low discrimination**						**65777542**
					COC	GRA	
Mitsuokella multacida		50.28	0.45	SUC 99	1	1	
Actinomyces israelii		48.68	0.57	BGL 10	1	99	
Blautia producta		0.84	0.07		99	99	
65777550	**Low discrimination**						**65777550**
					COC	GRA	
Mitsuokella multacida		84.59	0.53	SUC 99	1	1	
Blautia producta		12.69	0.32		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		2.72	0.40		1	99	
65777551	**Good identification**						**65777551**
					COC	GRA	
Mitsuokella multacida		98.66	0.37	SUC 99	1	1	
Blautia producta		1.35	-.03	URE 10	99	99	
65777552	**Low discrimination**						**65777552**
					COC	GRA	
Mitsuokella multacida		62.02	0.37	SUC 99	1	1	
Actinomyces israelii		26.92	0.43	BGL 10	1	99	
Blautia producta		9.30	0.15		99	99	
65777560	**Good identification**						**65777560**
					COC	GRA	
Mitsuokella multacida		97.49	0.51	SUC 99	1	1	
Actinomyces israelii		1.79	0.34		1	99	
Blautia producta		0.72	0.07		99	99	
65777570	**Acceptable identification**						**65777570**
					COC	GRA	
Mitsuokella multacida		93.03	0.43	SUC 99	1	1	
Blautia producta		6.20	0.15		99	99	
Actinomyces israelii		0.76	0.20		1	99	
65777700	**Very good identification**						**65777700**
					GRA		
Mitsuokella multacida		99.26	0.66	SUC 99	1		
Actinomyces israelii		0.74	0.42		99		
65777701	**Good identification**						**65777701**
Mitsuokella multacida		100.0	0.50	SUC 99			
65777702	**Acceptable identification**						**65777702**
					GRA		
Mitsuokella multacida		90.84	0.50	SUC 99	1		
Actinomyces israelii		9.17	0.45	BGL 10	99		
65777710	**Very good identification**						**65777710**
					COC	GRA	
Mitsuokella multacida		99.59	0.58	SUC 99	1	1	
Actinomyces israelii		0.33	0.28		1	99	
Blautia producta		0.08	-.03		99	99	
65777711	**Good identification**						**65777711**
Mitsuokella multacida		100.0	0.41	SUC 99			
65777712	**Acceptable identification**						**65777712**
					GRA		
Mitsuokella multacida		95.00	0.41	SUC 99	1		
Actinomyces israelii		4.30	0.31	BGL 10	99		
Bifidobacterium dentium		0.71	-.07		99		
65777720	**Good identification**						**65777720**
					GRA		
Mitsuokella multacida		99.81	0.56	SUC 99	1		
Actinomyces israelii		0.19	0.22		99		
65777721	**Good identification**						**65777721**
Mitsuokella multacida		100.0	0.39	SUC 99			
65777722	**Good identification**						**65777722**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Mitsuokella multacida		95.73	0.39	SUC 99	1		
Actinomyces israelii		2.47	0.25	BGL 10	99		
Bacteroides ovatus		0.93	-.02		1		
65777730	**Good identification**						**65777730**
Mitsuokella multacida		100.0	0.47	SUC 99			
65777740	**Very good identification**						**65777740**
					COC	GRA	
Mitsuokella multacida		98.83	0.66	SUC 99	1	1	
Blautia producta		0.89	0.24		99	99	
Actinomyces israelii		0.29	0.35		1	99	
65777741	**Good identification**						**65777741**
Mitsuokella multacida		100.0	0.50	SUC 99			
65777742	**Good identification**						**65777742**
					COC	GRA	
Mitsuokella multacida		95.40	0.50	SUC 99	1	1	
Actinomyces israelii		3.74	0.38	BGL 10	1	99	
Blautia producta		0.86	0.07		99	99	
65777750	**Good identification**						**65777750**
					COC	GRA	
Mitsuokella multacida		92.42	0.58	SUC 99	1	1	
Blautia producta		7.47	0.32		99	99	
Actinomyces israelii		0.12	0.21		1	99	
65777751	**Good identification**						**65777751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	SUC 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
65777752	**Acceptable identification**						**65777752**
					COC	GRA	
Mitsuokella multacida		91.04	0.41	SUC 99	1	1	
Blautia producta		7.35	0.15	BGL 10	99	99	
Actinomyces israelii		1.60	0.24		1	99	
65777760	**Good identification**						**65777760**
					COC	GRA	
Mitsuokella multacida		99.53	0.56	SUC 99	1	1	
Blautia producta		0.40	0.07		99	99	
Actinomyces israelii		0.07	0.15		1	99	
65777761	**Good identification**						**65777761**
Mitsuokella multacida		100.0	0.39	SUC 99			
65777762	**Good identification**						**65777762**
					GRA		
Mitsuokella multacida		99.01	0.39	SUC 99	1		
Actinomyces israelii		0.99	0.18	BGL 10	99		
65777770	**Good identification**						**65777770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	SUC 99	1	1	
Blautia producta		3.47	0.15		99	99	
66101101	**Very good identification**						**66101101**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		100.0	0.60	GLU 99			
66101103	**Good identification**						**66101103**
Clostridium chauvoei		100.0	0.47	GLU 99			
66101121	**Good identification**						**66101121**
Clostridium chauvoei		100.0	0.51	GLU 99			
66101123	**Good identification**						**66101123**
Clostridium chauvoei		100.0	0.39	GLU 99			
66111101	**Very good identification**						**66111101**
Clostridium chauvoei		100.0	0.66	GLU 99			
66111103	**Good identification**						**66111103**
					GRA	SPO	
Clostridium chauvoei		99.82	0.53	GLU 99	99	99	
Capnocytophaga ochracea		0.10	-.01		1	1	
Actinomyces viscosus		0.09	-.01		99	1	
66111120	**Low discrimination**						**66111120**
					GRA	CAT	
Clostridium perfringens		49.56	0.42	GLU 99	99	1	
Prevotella melaninogenica		24.25	0.30		1	1	
Capnocytophaga ochracea		21.85	0.33		1	90	
66111121	**Very good identification**						**66111121**
					GRA	SPO	
Clostridium chauvoei		99.78	0.58	GLU 99	99	99	
Clostridium perfringens		0.12	0.08		99	84	
Prevotella melaninogenica		0.06	-.04		1	1	
66111122	**Low discrimination**						**66111122**
					GRA	CAT	
Capnocytophaga ochracea		87.16	0.50	GLU 99	1	90	
Clostridium perfringens		10.34	0.37	RAF 90	99	1	
Prevotella melaninogenica		2.05	0.18		1	1	
66111123	**Good identification**						**66111123**
					GRA	CAT	
Clostridium chauvoei		97.15	0.45	GLU 99	99	1	
Capnocytophaga ochracea		2.55	0.15		1	90	
Clostridium perfringens		0.30	0.03		99	1	
66111522	**Good identification**						**66111522**
					GRA	CAT	
Capnocytophaga ochracea		99.33	0.37	GLU 99	1	90	
Clostridium perfringens		0.67	0.03	RAF 90	99	1	
66115122	**Good identification**						**66115122**
					GRA	CAT	
Capnocytophaga ochracea		95.81	0.39	GLU 99	1	90	
Bacteroides fragilis		3.73	0.06	RAF 90	1	99	
Clostridium perfringens		0.46	0.03		99	1	
66121101	**Good identification**						**66121101**
Clostridium chauvoei		100.0	0.43	GLU 99			
66121121	**Good identification**						**66121121**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		100.0	0.35	GLU 99			
66131101	**Good identification**						**66131101**
Clostridium chauvoei		100.0	0.50	GLU 99			
66131103	**Good identification**						**66131103**
Clostridium chauvoei		100.0	0.37	GLU 99			
66131120	**Good identification**						**66131120**
					COC	SPO	
Clostridium perfringens		98.56	0.47	GLU 99	1	84	
Clostridium chauvoei		0.52	0.07		1	99	
Atopobium parvulum		0.43	0.06		99	1	
66131121	**Good identification**						**66131121**
Clostridium chauvoei		98.09	0.41	GLU 99			
Clostridium perfringens		1.91	0.13	TRE 10			
66131122	**Good identification**						**66131122**
					GRA	CAT	
Clostridium perfringens		95.62	0.41	GLU 99	99	1	
Capnocytophaga ochracea		4.39	0.15		1	90	
66151122	**Acceptable identification**						**66151122**
					GRA	CAT	
Capnocytophaga ochracea		94.62	0.39	GLU 99	1	90	
Clostridium perfringens		1.87	0.13	RAF 90	99	1	
Prevotella oralis		1.75	0.09		1	1	
66331120	**Good identification**						**66331120**
					COC		
Clostridium perfringens		97.02	0.37	GLU 99	1		
Atopobium parvulum		2.22	0.08		99		
Actinomyces naeslundii		0.75	0.23		1		
66357100	**Good identification**						**66357100**
Actinomyces israelii		100.0	0.38	GLU 99			
66357102	**Good identification**						**66357102**
					SPO		
Actinomyces israelii		96.00	0.41	GLU 99	1		
Clostridium tertium		4.00	0.03		99		
66357500	**Good identification**						**66357500**
Actinomyces israelii		100.0	0.36	GLU 99			
66357502	**Good identification**						**66357502**
Actinomyces israelii		95.41	0.39	GLU 99			
Bifidobacterium adolescentis		4.59	0.11				
66371112	**Good identification**						**66371112**
Leptotrichia buccalis		100.0	0.50	NIT 10			
66375102	**Low discrimination**						**66375102**
					COC		
Actinomyces israelii		66.93	0.38	GLU 99	1		
Atopobium parvulum		33.07	0.19		99		
66375502	**Low discrimination**						**66375502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Actinomyces israelii		73.08	0.36	GLU 99	99	1	
Clostridium butyricum		13.55	0.03		99	99	
Mitsuokella multacida		5.47	0.04		1	1	
66377000	**Good identification**						**66377000**
Actinomyces israelii		100.0	0.37	GLU 99			
66377002	**Good identification**						**66377002**
Actinomyces israelii		100.0	0.40	GLU 99			
66377100	**Good identification**						**66377100**
					SPO		
Actinomyces israelii		95.25	0.48	GLU 99	1		
Clostridium tertium		4.75	0.11		99		
66377102	**Acceptable identification**						**66377102**
					SPO		
Actinomyces israelii		90.93	0.51	GLU 99	1		
Clostridium tertium		9.07	0.19		99		
66377140	**Good identification**						**66377140**
Actinomyces israelii		100.0	0.41	GLU 99			
66377142	**Good identification**						**66377142**
Actinomyces israelii		100.0	0.44	GLU 99			
66377400	**Good identification**						**66377400**
Actinomyces israelii		100.0	0.36	GLU 99			
66377402	**Good identification**						**66377402**
Actinomyces israelii		100.0	0.39	GLU 99			
66377500	**Low discrimination**						**66377500**
					GRA	SPO	
Actinomyces israelii		69.91	0.46	GLU 99	99	1	
Mitsuokella multacida		29.11	0.27		1	1	
Clostridium butyricum		0.98	-.07		99	99	
66377502	**Good identification**						**66377502**
					GRA		
Actinomyces israelii		96.21	0.49	GLU 99	99		
Mitsuokella multacida		2.97	0.11		1		
Bifidobacterium adolescentis		0.82	0.08		99		
66377512	**Low discrimination**						**66377512**
Actinomyces israelii		61.37	0.35	GLU 99			
Bifidobacterium dentium		30.13	0.06	MLZ 13			
Bifidobacterium adolescentis		4.28	0.09				
66377542	**Acceptable identification**						**66377542**
					GRA		
Actinomyces israelii		92.65	0.42	GLU 99	99		
Mitsuokella multacida		7.35	0.11		1		
66511102	**Good identification**						**66511102**
					GRA	CAT	
Capnocytophaga ochracea		95.09	0.50	GLU 99	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		4.48	0.28	NAG 90	99	76	
Prevotella melaninogenica		0.20	0.00		1	1	
66511120	**Low discrimination**						**66511120**
					GRA	CAT	
Prevotella melaninogenica		51.40	0.47	NIT 1	1	1	
Capnocytophaga ochracea		46.31	0.50	GLU 90	1	90	
Clostridium perfringens		2.22	0.30		99	1	
66511122	**Very good identification**						**66511122**
					GRA	CAT	
Capnocytophaga ochracea		97.17	0.66	GLU 99	1	90	
Prevotella melaninogenica		2.28	0.34		1	1	
Clostridium perfringens		0.24	0.24		99	1	
66511132	**Good identification**						**66511132**
					CAT		
Capnocytophaga ochracea		99.79	0.50	GLU 99	90		
Prevotella melaninogenica		0.21	0.00	MLZ 10	1		
66511162	**Good identification**						**66511162**
					GRA	CAT	
Capnocytophaga ochracea		99.36	0.50	GLU 99	1	90	
Clostridium perfringens		0.43	0.12	SOR 10	99	1	
Prevotella melaninogenica		0.21	0.00		1	1	
66511502	**Good identification**						**66511502**
					GRA	CAT	
Capnocytophaga ochracea		99.18	0.37	GLU 99	1	90	
Bifidobacterium longum subsp. longum		0.82	0.09	NAG 90	99	1	
66511520	**Acceptable identification**						**66511520**
					GRA	CAT	
Capnocytophaga ochracea		91.19	0.37	GLU 99	1	90	
Prevotella melaninogenica		5.79	0.12	BGL 90	1	1	
Bifidobacterium longum subsp. longum		3.01	0.20		99	1	
66511522	**Good identification**						**66511522**
					GRA	CAT	
Capnocytophaga ochracea		99.50	0.53	GLU 99	1	90	
Bifidobacterium longum subsp. longum		0.37	0.20		99	1	
Prevotella melaninogenica		0.13	0.00		1	1	
66511532	**Acceptable identification**						**66511532**
					GRA	CAT	
Capnocytophaga ochracea		94.67	0.37	GLU 99	1	90	
Bifidobacterium longum subsp. longum		5.33	0.24	MLZ 10	99	1	
66511562	**Good identification**						**66511562**
Capnocytophaga ochracea		100.0	0.37	GLU 99			
66515102	**Good identification**						**66515102**
Capnocytophaga ochracea		96.25	0.39	GLU 99			
Bacteroides fragilis		3.75	0.06	NAG 90			
66515120	**Low discrimination**						**66515120**
					CAT		
Capnocytophaga ochracea		79.70	0.39	GLU 99	90		
Bacteroides fragilis		16.19	0.18	BGL 90	99		
Prevotella melaninogenica		3.57	0.12		1		
66515122	**Low discrimination**						**66515122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Capnocytophaga ochracea		69.85	0.56	GLU 99	90		
Bacteroides fragilis		29.95	0.40		99		
Prevotella melaninogenica		0.07	0.00		1		
66515132	**Good identification**						**66515132**
Capnocytophaga ochracea		96.25	0.39	GLU 99			
Bacteroides fragilis		3.75	0.06	MLZ 10			
66515162	**Good identification**						**66515162**
Capnocytophaga ochracea		96.25	0.39	GLU 99			
Bacteroides fragilis		3.75	0.06	SOR 10			
66515522	**Acceptable identification**						**66515522**
					GRA	CAT	
Capnocytophaga ochracea		93.58	0.43	GLU 99	1	90	
Bifidobacterium longum subsp. longum		4.12	0.28		99	1	
Bacteroides fragilis		2.30	0.06		1	99	
66533100	**Low discrimination**						**66533100**
Actinomyces naeslundii		66.60	0.37	ESL 12			
Propionibacterium propionicum		33.40	0.21	GLU 99			
66533101	**Good identification**						**66533101**
Actinomyces naeslundii		100.0	0.40	ESL 12			
66533102	**Low discrimination**						**66533102**
Actinomyces naeslundii		57.07	0.37	ESL 12			
Propionibacterium propionicum		42.93	0.24	GLU 99			
66533103	**Good identification**						**66533103**
Actinomyces naeslundii		100.0	0.40	ESL 12			
66533120	**Low discrimination**						**66533120**
Actinomyces naeslundii		86.38	0.37	ESL 12			
Clostridium perfringens		13.62	0.01	GLU 99			
66533121	**Good identification**						**66533121**
Actinomyces naeslundii		100.0	0.40	ESL 12			
66533122	**Low discrimination**						**66533122**
					GRA		
Actinomyces naeslundii		86.14	0.37	ESL 12	99		
Capnocytophaga ochracea		13.86	-.03	GLU 99	1		
66533123	**Good identification**						**66533123**
Actinomyces naeslundii		100.0	0.40	ESL 12			
66551102	**Acceptable identification**						**66551102**
					GRA	CAT	
Capnocytophaga ochracea		93.52	0.39	GLU 99	1	90	
Actinomyces viscosus		4.68	0.18	NAG 90	99	76	
Prevotella oralis		1.05	0.06		1	1	
66551120	**Low discrimination**						**66551120**
					CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Capnocytophaga ochracea		65.77	0.39	GLU 99	90	
Prevotella melaninogenica		32.44	0.30	BGL 90	1	
Prevotella oralis		0.74	0.06		1	
66551122	**Good identification**					**66551122**
					CAT	
Capnocytophaga ochracea		96.41	0.56	GLU 99	90	
Prevotella oralis		2.28	0.27		1	
Prevotella melaninogenica		1.01	0.18		1	
66551132	**Good identification**					**66551132**
Capnocytophaga ochracea		100.0	0.39	GLU 99		
66551162	**Good identification**					**66551162**
Capnocytophaga ochracea		100.0	0.39	GLU 99		
66551522	**Good identification**					**66551522**
Capnocytophaga ochracea		100.0	0.43	GLU 99		
66553101	**Good identification**					**66553101**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66553103	**Good identification**					**66553103**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66553121	**Good identification**					**66553121**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66553123	**Good identification**					**66553123**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66555122	**Low discrimination**					**66555122**
					CAT	
Capnocytophaga ochracea		57.25	0.45	GLU 99	90	
Bacteroides fragilis		42.08	0.34		99	
Prevotella oralis		0.67	0.12		1	
66571101	**Good identification**					**66571101**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66571103	**Good identification**					**66571103**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66571121	**Good identification**					**66571121**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66571123	**Good identification**					**66571123**
Actinomyces naeslundii		100.0	0.37	ESL 12		
66573100	**Low discrimination**					**66573100**
Actinomyces naeslundii		74.96	0.39	ESL 12		
Actinomyces israelii		25.05	0.21	GLU 99		
66573101	**Good identification**					**66573101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	ESL 12		
66573102	**Low discrimination**					**66573102**
Actinomyces naeslundii		66.61	0.39	ESL 12		
Actinomyces israelii		33.39	0.24	GLU 99		
66573103	**Good identification**					**66573103**
Actinomyces naeslundii		100.0	0.42	ESL 12		
66573120	**Good identification**					**66573120**
Actinomyces naeslundii		100.0	0.39	ESL 12		
66573121	**Good identification**					**66573121**
Actinomyces naeslundii		100.0	0.42	ESL 12		
66573122	**Good identification**					**66573122**
Actinomyces naeslundii		100.0	0.39	ESL 12		
66573123	**Good identification**					**66573123**
Actinomyces naeslundii		100.0	0.42	ESL 12		
66575712	**Good identification**					**66575712**
Clostridium clostridioforme		98.69	0.36	NIT 1		
Mitsuokella multacida		1.31	0.01	GLU 88		
66575732	**Acceptable identification**					**66575732**
Clostridium clostridioforme		93.64	0.36	NIT 1		
Parabacteroides distasonis		3.90	0.18	GLU 88		
Bacteroides thetaiotaomicron		2.47	-.01	SAL 88		
66577100	**Good identification**					**66577100**
Actinomyces israelii		100.0	0.39	GLU 99		
66577102	**Good identification**					**66577102**
Actinomyces israelii		100.0	0.42	GLU 99		
66577142	**Good identification**					**66577142**
Actinomyces israelii		100.0	0.35	GLU 99		
66577502	**Low discrimination**					**66577502**
					GRA	
Actinomyces israelii		87.17	0.41	GLU 99	99	
Mitsuokella multacida		8.22	0.11	SAL 92	1	
Bifidobacterium adolescentis		4.61	0.13		99	
66711101	**Good identification**					**66711101**
Actinomyces naeslundii		100.0	0.39	ESL 12		
66711103	**Low discrimination**					**66711103**
Actinomyces naeslundii		61.59	0.39	ESL 12		
Actinomyces viscosus		31.90	0.10	GLU 99		
Bifidobacterium longum subsp. infantis		6.52	-.07			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
66711121	**Good identification**						**66711121**
Actinomyces naeslundii		100.0	0.39	ESL 12			
66711123	**Low discrimination**						**66711123**
					GRA	CAT	
Actinomyces naeslundii		76.79	0.39	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		14.46	-.03	GLU 99	99	1	
Capnocytophaga ochracea		8.75	-.03		1	90	
66713001	**Good identification**						**66713001**
Actinomyces naeslundii		100.0	0.36	ESL 12			
66713003	**Good identification**						**66713003**
Actinomyces naeslundii		100.0	0.36	ESL 12			
66713021	**Good identification**						**66713021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
66713023	**Good identification**						**66713023**
Actinomyces naeslundii		100.0	0.36	ESL 12			
66713100	**Good identification**						**66713100**
Actinomyces naeslundii		100.0	0.41	ESL 12			
66713101	**Good identification**						**66713101**
Actinomyces naeslundii		100.0	0.44	ESL 12			
66713102	**Low discrimination**						**66713102**
Actinomyces naeslundii		85.80	0.41	ESL 12			
Actinomyces israelii		7.18	0.13	GLU 99			
Bifidobacterium longum subsp. infantis		7.02	-.07				
66713103	**Good identification**						**66713103**
Actinomyces naeslundii		100.0	0.44	ESL 12			
66713120	**Good identification**						**66713120**
Actinomyces naeslundii		100.0	0.41	ESL 12			
66713121	**Good identification**						**66713121**
Actinomyces naeslundii		100.0	0.44	ESL 12			
66713122	**Low discrimination**						**66713122**
					GRA	CAT	
Actinomyces naeslundii		81.07	0.41	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		11.79	-.03	GLU 99	99	1	
Capnocytophaga ochracea		7.14	-.03		1	90	
66713123	**Good identification**						**66713123**
Actinomyces naeslundii		100.0	0.44	ESL 12			
66731001	**Good identification**						**66731001**
Actinomyces naeslundii		100.0	0.36	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
66731003	**Good identification**					**66731003**
Actinomyces naeslundii		100.0	0.36	ESL 12		
66731021	**Good identification**					**66731021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
66731023	**Good identification**					**66731023**
Actinomyces naeslundii		100.0	0.36	ESL 12		
66731100	**Good identification**					**66731100**
Actinomyces naeslundii		100.0	0.41	ESL 12		
66731101	**Good identification**					**66731101**
Actinomyces naeslundii		100.0	0.44	ESL 12		
66731102	**Low discrimination**					**66731102**
Actinomyces naeslundii		85.31	0.41	ESL 12		
Actinomyces viscosus		7.71	-.01	GLU 99		
Bifidobacterium longum subsp. infantis		6.98	-.07			
66731103	**Good identification**					**66731103**
Actinomyces naeslundii		100.0	0.44	ESL 12		
66731121	**Good identification**					**66731121**
Actinomyces naeslundii		100.0	0.44	ESL 12		
66731123	**Good identification**					**66731123**
Actinomyces naeslundii		100.0	0.44	ESL 12		
66733000	**Good identification**					**66733000**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66733001	**Good identification**					**66733001**
Actinomyces naeslundii		100.0	0.41	ESL 12		
66733002	**Good identification**					**66733002**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66733003	**Good identification**					**66733003**
Actinomyces naeslundii		100.0	0.41	ESL 12		
66733020	**Good identification**					**66733020**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66733021	**Good identification**					**66733021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
66733022	**Good identification**					**66733022**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66733023	**Good identification**					**66733023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
66733100	**Acceptable identification**					**66733100**
Actinomyces naeslundii		90.24	0.46	ESL 12		
Actinomyces israelii		9.76	0.20	GLU 99		
66733101	**Good identification**					**66733101**
Actinomyces naeslundii		100.0	0.49	ESL 12		
66733102	**Low discrimination**					**66733102**
Actinomyces naeslundii		86.04	0.46	ESL 12		
Actinomyces israelii		13.96	0.23	GLU 99		
66733103	**Good identification**					**66733103**
Actinomyces naeslundii		100.0	0.49	ESL 12		
66733120	**Good identification**					**66733120**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66733121	**Good identification**					**66733121**
Actinomyces naeslundii		100.0	0.49	ESL 12		
66733122	**Good identification**					**66733122**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66733123	**Good identification**					**66733123**
Actinomyces naeslundii		100.0	0.49	ESL 12		
66733141	**Good identification**					**66733141**
Actinomyces naeslundii		100.0	0.36	ESL 12		
66733143	**Good identification**					**66733143**
Actinomyces naeslundii		100.0	0.36	ESL 12		
66733161	**Good identification**					**66733161**
Actinomyces naeslundii		100.0	0.36	ESL 12		
66733163	**Good identification**					**66733163**
Actinomyces naeslundii		100.0	0.36	ESL 12		
66735712	**Good identification**					**66735712**
Clostridium clostridioforme		99.40	0.42	NIT 1		
Mitsuokella multacida		0.60	0.01	GLU 88		
66735732	**Good identification**					**66735732**
Clostridium clostridioforme		96.32	0.42	NIT 1		
Parabacteroides distasonis		3.68	0.23	GLU 88		
66737100	**Good identification**					**66737100**
Actinomyces israelii		100.0	0.38	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
66737102	**Good identification**					**66737102**
Actinomyces israelii		100.0	0.41	GLU 99		
66737502	**Low discrimination**					**66737502**
					GRA	
Actinomyces israelii		78.63	0.40	GLU 99	99	
Bifidobacterium adolescentis		12.81	0.20	CEL 93	99	
Mitsuokella multacida		8.56	0.11		1	
66751100	**Low discrimination**					**66751100**
Actinomyces naeslundii		52.29	0.38	ESL 12		
Collinsella aerofaciens		29.07	-.03	GLU 99		
Actinomyces israelii		10.02	0.17			
66751101	**Good identification**					**66751101**
Actinomyces naeslundii		100.0	0.41	ESL 12		
66751103	**Acceptable identification**					**66751103**
Actinomyces naeslundii		90.59	0.41	ESL 12		
Actinomyces viscosus		9.41	0.00	GLU 99		
66751121	**Good identification**					**66751121**
Actinomyces naeslundii		100.0	0.41	ESL 12		
66751123	**Acceptable identification**					**66751123**
					GRA	
Actinomyces naeslundii		91.80	0.41	ESL 12	99	
Prevotella oralis		8.20	-.03	GLU 99	1	
66753000	**Low discrimination**					**66753000**
Actinomyces naeslundii		71.53	0.35	ESL 12		
Actinomyces israelii		28.47	0.19	GLU 99		
66753001	**Good identification**					**66753001**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66753002	**Low discrimination**					**66753002**
Actinomyces naeslundii		62.62	0.35	ESL 12		
Actinomyces israelii		37.38	0.22	GLU 99		
66753003	**Good identification**					**66753003**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66753020	**Good identification**					**66753020**
Actinomyces naeslundii		100.0	0.35	ESL 12		
66753021	**Good identification**					**66753021**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66753022	**Good identification**					**66753022**
Actinomyces naeslundii		100.0	0.35	ESL 12		
66753023	**Good identification**					**66753023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	ESL 12		
66753100	**Low discrimination**					**66753100**
Actinomyces naeslundii		60.82	0.43	ESL 12		
Actinomyces israelii		34.01	0.30	GLU 99		
Bifidobacterium breve		5.17	0.10			
66753101	**Good identification**					**66753101**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66753102	**Low discrimination**					**66753102**
Actinomyces naeslundii		37.07	0.43	ESL 12		
Bifidobacterium breve		31.84	0.27	GLU 99		
Actinomyces israelii		31.09	0.33			
66753103	**Good identification**					**66753103**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66753120	**Low discrimination**					**66753120**
					SPO	
Actinomyces naeslundii		81.38	0.43	ESL 12	1	
Bifidobacterium breve		12.29	0.15	GLU 99	1	
Clostridium ramosum		6.34	-.01		92	
66753121	**Good identification**					**66753121**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66753123	**Good identification**					**66753123**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66755102	**Low discrimination**					**66755102**
Actinomyces israelii		88.93	0.38	GLU 99		
Bifidobacterium adolescentis		11.07	0.16			
66755712	**Good identification**					**66755712**
Clostridium clostridioforme		98.69	0.36	NIT 1		
Mitsuokella multacida		1.31	0.01	GLU 88		
66755732	**Good identification**					**66755732**
Clostridium clostridioforme		100.0	0.36	NIT 1		
66757000	**Good identification**					**66757000**
Actinomyces israelii		100.0	0.37	GLU 99		
66757002	**Good identification**					**66757002**
Actinomyces israelii		100.0	0.40	GLU 99		
66757100	**Good identification**					**66757100**
Actinomyces israelii		100.0	0.48	GLU 99		
66757102	**Good identification**					**66757102**
Actinomyces israelii		98.23	0.51	GLU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		1.77	0.15				
66757112	**Low discrimination**						**66757112**
Actinomyces israelii		87.18	0.37	GLU 99			
Bifidobacterium adolescentis		12.82	0.16	MLZ 13			
66757140	**Good identification**						**66757140**
Actinomyces israelii		100.0	0.41	GLU 99			
66757142	**Good identification**						**66757142**
Actinomyces israelii		100.0	0.44	GLU 99			
66757400	**Good identification**						**66757400**
Actinomyces israelii		100.0	0.36	GLU 99			
66757402	**Good identification**						**66757402**
Actinomyces israelii		100.0	0.39	GLU 99			
66757500	**Low discrimination**						**66757500**
					GRA	SPO	
Actinomyces israelii		69.77	0.46	GLU 99	99	1	
Mitsuokella multacida		29.05	0.27		1	1	
Clostridium butyricum		1.19	-.05		99	99	
66757502	**Low discrimination**						**66757502**
					GRA		
Actinomyces israelii		84.90	0.49	GLU 99	99		
Bifidobacterium adolescentis		12.48	0.29		99		
Mitsuokella multacida		2.62	0.11		1		
66757542	**Low discrimination**						**66757542**
					GRA		
Actinomyces israelii		88.81	0.42	GLU 99	99		
Mitsuokella multacida		7.04	0.11		1		
Bifidobacterium adolescentis		4.15	0.13		99		
66771000	**Low discrimination**						**66771000**
Actinomyces naeslundii		79.11	0.35	ESL 12			
Actinomyces israelii		20.90	0.16	GLU 99			
66771001	**Good identification**						**66771001**
Actinomyces naeslundii		100.0	0.38	ESL 12			
66771002	**Low discrimination**						**66771002**
Actinomyces naeslundii		71.62	0.35	ESL 12			
Actinomyces israelii		28.38	0.19	GLU 99			
66771003	**Good identification**						**66771003**
Actinomyces naeslundii		100.0	0.38	ESL 12			
66771020	**Good identification**						**66771020**
Actinomyces naeslundii		100.0	0.35	ESL 12			
66771021	**Good identification**						**66771021**
Actinomyces naeslundii		100.0	0.38	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
66771022	**Good identification**					**66771022**
Actinomyces naeslundii		100.0	0.35	ESL 12		
66771023	**Good identification**					**66771023**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66771100	**Low discrimination**					**66771100**
Actinomyces naeslundii		72.93	0.43	ESL 12		
Actinomyces israelii		27.07	0.27	GLU 99		
66771101	**Good identification**					**66771101**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66771102	**Low discrimination**					**66771102**
					GRA	
Actinomyces naeslundii		29.44	0.43	ESL 12	99	
Leptotrichia buccalis		23.42	0.07	GLU 99	1	
Bifidobacterium breve		16.86	0.24		99	
66771103	**Good identification**					**66771103**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66771112	**Good identification**					**66771112**
					GRA	
Leptotrichia buccalis		99.17	0.41	NIT 10	1	
Bifidobacterium breve		0.72	0.24	GLU 99	99	
Actinomyces israelii		0.11	0.15		99	
66771120	**Low discrimination**					**66771120**
					SPO	
Actinomyces naeslundii		58.80	0.43	ESL 12	1	
Clostridium paraputrificum		18.67	0.03	GLU 99	99	
Clostridium ramosum		12.66	0.06		92	
66771121	**Good identification**					**66771121**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66771123	**Good identification**					**66771123**
Actinomyces naeslundii		100.0	0.46	ESL 12		
66773000	**Low discrimination**					**66773000**
Actinomyces naeslundii		56.46	0.40	ESL 12		
Actinomyces israelii		43.54	0.29	GLU 99		
66773001	**Good identification**					**66773001**
Actinomyces naeslundii		100.0	0.43	ESL 12		
66773003	**Good identification**					**66773003**
Actinomyces naeslundii		100.0	0.43	ESL 12		
66773020	**Good identification**					**66773020**
Actinomyces naeslundii		100.0	0.40	ESL 12		
66773021	**Good identification**					**66773021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ESL 12		
66773022	**Good identification**					**66773022**
Actinomyces naeslundii		100.0	0.40	ESL 12		
66773023	**Good identification**					**66773023**
Actinomyces naeslundii		100.0	0.43	ESL 12		
66773100	**Low discrimination**					**66773100**
Actinomyces israelii		52.01	0.39	GLU 99		
Actinomyces naeslundii		47.99	0.48	XYL 92		
66773101	**Good identification**					**66773101**
Actinomyces naeslundii		100.0	0.51	ESL 12		
66773102	**Low discrimination**					**66773102**
Actinomyces israelii		55.66	0.42	GLU 99		
Actinomyces naeslundii		34.24	0.48	XYL 92		
Bifidobacterium breve		10.10	0.24			
66773103	**Good identification**					**66773103**
Actinomyces naeslundii		100.0	0.51	ESL 12		
66773120	**Low discrimination**					**66773120**
					SPO	
Actinomyces naeslundii		57.15	0.48	ESL 12	1	
Clostridium ramosum		35.93	0.19	GLU 99	92	
Actinomyces israelii		3.95	0.19		1	
66773121	**Good identification**					**66773121**
Actinomyces naeslundii		100.0	0.51	ESL 12		
66773123	**Good identification**					**66773123**
Actinomyces naeslundii		100.0	0.51	ESL 12		
66773141	**Good identification**					**66773141**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66773142	**Low discrimination**					**66773142**
Actinomyces israelii		57.28	0.35	GLU 99		
Bifidobacterium breve		26.73	0.24	XYL 92		
Actinomyces naeslundii		15.99	0.35			
66773143	**Good identification**					**66773143**
Actinomyces naeslundii		100.0	0.38	ESL 12		
66773160	**Low discrimination**					**66773160**
Actinomyces naeslundii		77.29	0.35	ESL 12		
Bifidobacterium breve		22.71	0.12	GLU 99		
66773161	**Good identification**					**66773161**
Actinomyces naeslundii		100.0	0.38	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
66773163	**Good identification**						**66773163**
Actinomyces naeslundii		100.0	0.38	ESL 12			
66773502	**Low discrimination**						**66773502**
					GRA		
Actinomyces israelii		87.79	0.41	GLU 99	99		
Mitsuokella multacida		8.28	0.11	XYL 92	1		
Bifidobacterium adolescentis		3.93	0.12		99		
66774712	**Good identification**						**66774712**
Clostridium clostridioforme		99.46	0.43	NIT 1			
Mitsuokella multacida		0.54	0.01	GLU 88			
66774732	**Good identification**						**66774732**
Clostridium clostridioforme		100.0	0.43	NIT 1			
66775002	**Good identification**						**66775002**
Actinomyces israelii		100.0	0.37	GLU 99			
66775100	**Low discrimination**						**66775100**
					GRA	SPO	
Actinomyces israelii		64.59	0.45	GLU 99	99	1	
Clostridium butyricum		21.21	0.15		99	99	
Mitsuokella multacida		14.20	0.21		1	1	
66775102	**Low discrimination**						**66775102**
					SPO		
Actinomyces israelii		88.17	0.48	GLU 99	1		
Clostridium butyricum		7.51	0.08		99		
Bifidobacterium adolescentis		1.95	0.13		1		
66775140	**Low discrimination**						**66775140**
					GRA	SPO	
Actinomyces israelii		61.02	0.38	GLU 99	99	1	
Mitsuokella multacida		34.50	0.21		1	1	
Clostridium butyricum		4.48	-.03		99	99	
66775142	**Good identification**						**66775142**
					GRA		
Actinomyces israelii		95.98	0.41	GLU 99	99		
Mitsuokella multacida		4.02	0.04		1		
66775402	**Low discrimination**						**66775402**
					SPO		
Actinomyces israelii		67.59	0.36	GLU 99	1		
Clostridium clostridioforme		16.48	0.16		66		
Clostridium butyricum		10.55	0.01		99		
66775500	**Low discrimination**						**66775500**
					GRA	SPO	
Mitsuokella multacida		85.23	0.55	GLU 99	1	1	
Clostridium butyricum		11.57	0.32		99	99	
Actinomyces israelii		3.20	0.43		99	1	
66775501	**Good identification**						**66775501**
					GRA	SPO	
Mitsuokella multacida		98.78	0.39	GLU 99	1	1	
Clostridium butyricum		1.22	-.03	URE 10	99	99	
66775502	**Low discrimination**						**66775502**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				GRA	SPO	
Mitsuokella multacida	45.92	0.39	GLU 99	1	1	
Actinomyces israelii	23.30	0.46	BGL 10	99	1	
Clostridium butyricum	21.82	0.25		99	99	
66775510 Good identification						**66775510**
				GRA	SPO	
Mitsuokella multacida	96.31	0.47	GLU 99	1	1	
Clostridium butyricum	2.07	0.10		99	99	
Actinomyces israelii	1.62	0.29		99	1	
66775512 Low discrimination						**66775512**
				GRA		
Clostridium clostridioforme	45.86	0.36	NIT 1	50		
Mitsuokella multacida	32.41	0.30	GLU 88	1		
Bifidobacterium adolescentis	10.88	0.29	RHA 88	99		
66775520 Acceptable identification						**66775520**
				GRA	SPO	
Mitsuokella multacida	94.62	0.45	GLU 99	1	1	
Clostridium butyricum	4.47	0.14		99	99	
Actinomyces israelii	0.91	0.23		99	1	
66775530 Good identification						**66775530**
Mitsuokella multacida	100.0	0.37	GLU 99			
66775532 Low discrimination						**66775532**
				GRA		
Clostridium clostridioforme	81.15	0.36	NIT 1	50		
Mitsuokella multacida	14.34	0.20	GLU 88	1		
Bifidobacterium dentium	3.29	-.03	RHA 88	99		
66775540 Good identification						**66775540**
				GRA	SPO	
Mitsuokella multacida	97.43	0.55	GLU 99	1	1	
Actinomyces israelii	1.42	0.36		99	1	
Clostridium butyricum	1.15	0.14		99	99	
66775541 Good identification						**66775541**
Mitsuokella multacida	100.0	0.39	GLU 99			
66775542 Low discrimination						**66775542**
				GRA	SPO	
Mitsuokella multacida	80.00	0.39	GLU 99	1	1	
Actinomyces israelii	15.79	0.39	BGL 10	99	1	
Clostridium butyricum	3.31	0.07		99	99	
66775550 Good identification						**66775550**
				COC	GRA	
Mitsuokella multacida	99.00	0.47	GLU 99	1	1	
Actinomyces israelii	0.65	0.22		1	99	
Blautia producta	0.35	-.03		99	99	
66775560 Good identification						**66775560**
				GRA	SPO	
Mitsuokella multacida	99.22	0.45	GLU 99	1	1	
Clostridium butyricum	0.41	-.05		99	99	
Actinomyces israelii	0.37	0.16		99	1	
66775570 Good identification						**66775570**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.37	GLU 99			
66775612	**Good identification**						**66775612**
Clostridium clostridioforme		99.40	0.42	NIT 1			
Mitsuokella multacida		0.60	0.01	GLU 88			
66775632	**Good identification**						**66775632**
Clostridium clostridioforme		100.0	0.42	NIT 1			
66775700	**Very good identification**						**66775700**
					GRA	SPO	
Mitsuokella multacida		99.58	0.60	GLU 99	1	1	
Clostridium butyricum		0.23	0.06		99	99	
Actinomyces israelii		0.15	0.24		99	1	
66775701	**Good identification**						**66775701**
Mitsuokella multacida		100.0	0.43	GLU 99			
66775702	**Low discrimination**						**66775702**
					GRA		
Mitsuokella multacida		69.58	0.43	GLU 99	1		
Clostridium clostridioforme		28.44	0.40	BGL 10	50		
Actinomyces israelii		1.43	0.27		99		
66775710	**Good identification**						**66775710**
Mitsuokella multacida		99.38	0.51	GLU 99			
Clostridium clostridioforme		0.62	0.17				
66775711	**Good identification**						**66775711**
Mitsuokella multacida		100.0	0.35	GLU 99			
66775712	**Low discrimination**						**66775712**
Clostridium clostridioforme		84.82	0.51	NIT 1			
Mitsuokella multacida		15.18	0.35	GLU 88			
66775720	**Good identification**						**66775720**
Mitsuokella multacida		99.82	0.49	GLU 99			
Clostridium clostridioforme		0.18	0.06				
66775722	**Low discrimination**						**66775722**
Clostridium clostridioforme		58.64	0.40	NIT 1			
Mitsuokella multacida		35.86	0.33	GLU 88			
Parabacteroides distasonis		3.05	0.23				
66775730	**Good identification**						**66775730**
Mitsuokella multacida		97.55	0.41	GLU 99			
Clostridium clostridioforme		2.45	0.17				
66775732	**Acceptable identification**						**66775732**
Clostridium clostridioforme		94.61	0.51	NIT 1			
Mitsuokella multacida		4.23	0.25	GLU 88			
Parabacteroides distasonis		0.58	0.19				
66775740	**Very good identification**						**66775740**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		99.94	0.60	GLU 99	1		
Actinomyces israelii		0.06	0.17		99		
66775741	**Good identification**						**66775741**
Mitsuokella multacida		100.0	0.43	GLU 99			
66775742	**Good identification**						**66775742**
					GRA		
Mitsuokella multacida		98.80	0.43	GLU 99	1		
Actinomyces israelii		0.79	0.20	BGL 10	99		
Clostridium clostridioforme		0.41	0.06		50		
66775750	**Good identification**						**66775750**
					COC	GRA	
Mitsuokella multacida		99.81	0.51	GLU 99	1	1	
Blautia producta		0.19	-.03		99	99	
66775751	**Good identification**						**66775751**
Mitsuokella multacida		100.0	0.35	GLU 99			
66775752	**Acceptable identification**						**66775752**
Mitsuokella multacida		94.66	0.35	GLU 99			
Clostridium clostridioforme		5.34	0.17	BGL 10			
66775760	**Good identification**						**66775760**
Mitsuokella multacida		100.0	0.49	GLU 99			
66775770	**Good identification**						**66775770**
Mitsuokella multacida		100.0	0.41	GLU 99			
66777000	**Good identification**						**66777000**
Actinomyces israelii		100.0	0.47	GLU 99			
66777002	**Good identification**						**66777002**
Actinomyces israelii		100.0	0.50	GLU 99			
66777012	**Good identification**						**66777012**
Actinomyces israelii		100.0	0.36	GLU 99			
66777040	**Good identification**						**66777040**
Actinomyces israelii		100.0	0.40	GLU 99			
66777042	**Good identification**						**66777042**
Actinomyces israelii		100.0	0.43	GLU 99			
66777100	**Good identification**						**66777100**
					GRA	SPO	
Actinomyces israelii		91.16	0.58	GLU 99	99	1	
Mitsuokella multacida		8.25	0.27		1	1	
Clostridium butyricum		0.59	-.01		99	99	
66777102	**Very good identification**						**66777102**
					GRA		
Actinomyces israelii		99.02	0.61	GLU 99	99		
Mitsuokella multacida		0.66	0.11		1		
Bifidobacterium adolescentis		0.32	0.12		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
66777110	**Low discrimination**						**66777110**
					GRA		
Actinomyces israelii		83.20	0.43	GLU 99	99		
Mitsuokella multacida		16.80	0.19	MLZ 13	1		
66777112	**Acceptable identification**						**66777112**
Actinomyces israelii		94.21	0.46	GLU 99			
Bifidobacterium adolescentis		2.46	0.13	MLZ 13			
Bifidobacterium dentium		1.93	-.07				
66777120	**Low discrimination**						**66777120**
					GRA		
Actinomyces israelii		73.82	0.37	GLU 99	99		
Mitsuokella multacida		26.18	0.17	NAG 6	1		
66777122	**Good identification**						**66777122**
Actinomyces israelii		100.0	0.40	GLU 99			
66777140	**Low discrimination**						**66777140**
					GRA		
Actinomyces israelii		81.11	0.51	GLU 99	99		
Mitsuokella multacida		18.89	0.27		1		
66777142	**Good identification**						**66777142**
					GRA		
Actinomyces israelii		98.30	0.54	GLU 99	99		
Mitsuokella multacida		1.70	0.11		1		
66777150	**Low discrimination**						**66777150**
					COC	GRA	
Actinomyces israelii		62.61	0.36	GLU 99	1	99	
Mitsuokella multacida		32.52	0.19	MLZ 13	1	1	
Blautia producta		4.88	-.03		99	99	
66777152	**Good identification**						**66777152**
					GRA		
Actinomyces israelii		96.29	0.39	GLU 99	99		
Mitsuokella multacida		3.71	0.03	MLZ 13	1		
66777302	**Low discrimination**						**66777302**
					GRA		
Actinomyces israelii		85.80	0.41	GLU 99	99		
Mitsuokella multacida		14.20	0.15	RHA 7	1		
66777400	**Low discrimination**						**66777400**
					GRA		
Actinomyces israelii		69.32	0.46	GLU 99	99		
Mitsuokella multacida		30.68	0.27		1		
66777402	**Good identification**						**66777402**
					GRA		
Actinomyces israelii		96.83	0.49	GLU 99	99		
Mitsuokella multacida		3.18	0.11		1		
66777442	**Acceptable identification**						**66777442**
					GRA		
Actinomyces israelii		92.22	0.42	GLU 99	99		
Mitsuokella multacida		7.78	0.11		1		
66777500	**Good identification**						**66777500**
					GRA	SPO	
Mitsuokella multacida		91.10	0.61	GLU 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		8.32	0.56		99	1	
Clostridium butyricum		0.59	0.15		99	99	
66777501	**Good identification**						**66777501**
					GRA		
Mitsuokella multacida		99.18	0.45	GLU 99	1		
Actinomyces israelii		0.82	0.22	URE 10	99		
66777502	**Low discrimination**						**66777502**
					GRA		
Actinomyces israelii		53.77	0.59	GLU 99	99		
Mitsuokella multacida		43.64	0.45		1		
Bifidobacterium adolescentis		1.40	0.26		99		
66777510	**Acceptable identification**						**66777510**
					GRA		
Mitsuokella multacida		94.47	0.53	GLU 99	1		
Actinomyces israelii		3.87	0.42		99		
Bifidobacterium dentium		1.42	0.10		99		
66777511	**Good identification**						**66777511**
Mitsuokella multacida		100.0	0.37	GLU 99			
66777520	**Good identification**						**66777520**
					GRA	SPO	
Mitsuokella multacida		97.51	0.51	GLU 99	1	1	
Actinomyces israelii		2.27	0.36		99	1	
Clostridium butyricum		0.22	-.03		99	99	
66777530	**Low discrimination**						**66777530**
					GRA		
Mitsuokella multacida		89.49	0.43	GLU 99	1		
Bifidobacterium dentium		9.58	0.15		99		
Actinomyces israelii		0.94	0.22		99		
66777540	**Good identification**						**66777540**
					COC	GRA	
Mitsuokella multacida		94.99	0.61	GLU 99	1	1	
Actinomyces israelii		3.37	0.49		1	99	
Blautia producta		1.58	0.24		99	99	
66777541	**Good identification**						**66777541**
					GRA		
Mitsuokella multacida		99.68	0.45	GLU 99	1		
Actinomyces israelii		0.32	0.15	URE 10	99		
66777542	**Low discrimination**						**66777542**
					COC	GRA	
Mitsuokella multacida		66.68	0.45	GLU 99	1	1	
Actinomyces israelii		31.95	0.52	BGL 10	1	99	
Blautia producta		1.11	0.07		99	99	
66777550	**Low discrimination**						**66777550**
					COC	GRA	
Mitsuokella multacida		85.77	0.53	GLU 99	1	1	
Blautia producta		12.87	0.32		99	99	
Actinomyces israelii		1.37	0.35		1	99	
66777551	**Good identification**						**66777551**
					COC	GRA	
Mitsuokella multacida		98.66	0.37	GLU 99	1	1	
Blautia producta		1.35	-.03	URE 10	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
66777552	**Low discrimination**						**66777552**
					COC	GRA	
Mitsuokella multacida		71.78	0.37	GLU 99	1	1	
Actinomyces israelii		15.42	0.38	BGL 10	1	99	
Blautia producta		10.77	0.15		99	99	
66777560	**Good identification**						**66777560**
					COC	GRA	
Mitsuokella multacida		98.38	0.51	GLU 99	1	1	
Actinomyces israelii		0.89	0.29		1	99	
Blautia producta		0.73	0.07		99	99	
66777570	**Acceptable identification**						**66777570**
					COC	GRA	
Mitsuokella multacida		93.39	0.43	GLU 99	1	1	
Blautia producta		6.23	0.15		99	99	
Actinomyces israelii		0.38	0.15		1	99	
66777700	**Very good identification**						**66777700**
					GRA		
Mitsuokella multacida		99.63	0.66	GLU 99	1		
Actinomyces israelii		0.37	0.37		99		
66777701	**Good identification**						**66777701**
Mitsuokella multacida		100.0	0.50	GLU 99			
66777702	**Good identification**						**66777702**
					GRA		
Mitsuokella multacida		95.08	0.50	GLU 99	1		
Actinomyces israelii		4.75	0.40	BGL 10	99		
Clostridium clostridioforme		0.17	0.06		50		
66777710	**Very good identification**						**66777710**
					COC	GRA	
Mitsuokella multacida		99.75	0.58	GLU 99	1	1	
Actinomyces israelii		0.17	0.23		1	99	
Blautia producta		0.08	-.03		99	99	
66777711	**Good identification**						**66777711**
Mitsuokella multacida		100.0	0.41	GLU 99			
66777712	**Acceptable identification**						**66777712**
					GRA		
Mitsuokella multacida		94.88	0.41	GLU 99	1		
Clostridium clostridioforme		2.30	0.17	BGL 10	50		
Actinomyces israelii		2.12	0.26		99		
66777720	**Good identification**						**66777720**
					GRA		
Mitsuokella multacida		99.91	0.56	GLU 99	1		
Actinomyces israelii		0.09	0.16		99		
66777721	**Good identification**						**66777721**
Mitsuokella multacida		100.0	0.39	GLU 99			
66777722	**Good identification**						**66777722**
					GRA		
Mitsuokella multacida		97.13	0.39	GLU 99	1		
Actinomyces israelii		1.24	0.19	BGL 10	99		
Bacteroides ovatus		0.95	-.02		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
66777730	**Good identification**						**66777730**
Mitsuokella multacida		100.0	0.47	GLU 99			
66777740	**Very good identification**						**66777740**
					COC	GRA	
Mitsuokella multacida		98.97	0.66	GLU 99	1	1	
Blautia producta		0.89	0.24		99	99	
Actinomyces israelii		0.14	0.30		1	99	
66777741	**Good identification**						**66777741**
Mitsuokella multacida		100.0	0.50	GLU 99			
66777742	**Good identification**						**66777742**
					COC	GRA	
Mitsuokella multacida		97.24	0.50	GLU 99	1	1	
Actinomyces israelii		1.89	0.33	BGL 10	1	99	
Blautia producta		0.87	0.07		99	99	
66777750	**Good identification**						**66777750**
					COC	GRA	
Mitsuokella multacida		92.47	0.58	GLU 99	1	1	
Blautia producta		7.47	0.32		99	99	
Actinomyces israelii		0.06	0.16		1	99	
66777751	**Good identification**						**66777751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	GLU 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
66777752	**Acceptable identification**						**66777752**
					COC	GRA	
Mitsuokella multacida		91.79	0.41	GLU 99	1	1	
Blautia producta		7.41	0.15	BGL 10	99	99	
Actinomyces israelii		0.80	0.19		1	99	
66777760	**Good identification**						**66777760**
					COC	GRA	
Mitsuokella multacida		99.60	0.56	GLU 99	1	1	
Blautia producta		0.40	0.07		99	99	
66777761	**Good identification**						**66777761**
Mitsuokella multacida		100.0	0.39	GLU 99			
66777762	**Good identification**						**66777762**
Mitsuokella multacida		100.0	0.39	GLU 99			
66777770	**Good identification**						**66777770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	GLU 99	1	1	
Blautia producta		3.47	0.15		99	99	
67001101	**Very good identification**						**67001101**
Clostridium chauvoei		100.0	0.60	MLT 99			
67001103	**Good identification**						**67001103**
Clostridium chauvoei		100.0	0.47	MLT 99			
67001121	**Good identification**						**67001121**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Clostridium chauvoei		100.0	0.51	MLT 99				
67001123	**Good identification**							**67001123**
Clostridium chauvoei		100.0	0.39	MLT 99				
67010000	**Low discrimination**							**67010000**
					GRA			
Fusobacterium mortiferum		62.75	0.47	NIT 1	1			
Eubacterium limosum		14.53	0.18		99			
Atopobium minutum		10.11	0.28		99			
67010100	**Good identification**							**67010100**
Fusobacterium mortiferum		100.0	0.53	NIT 1				
67010102	**Low discrimination**							**67010102**
					GRA	CAT		
Fusobacterium mortiferum		82.06	0.37	NIT 1	1	5		
Actinomyces viscosus		14.60	0.11	BGL 10	99	76		
Capnocytophaga ochracea		3.34	-.01		1	90		
67011000	**Low discrimination**							**67011000**
					GRA	CAT		
Fusobacterium mortiferum		89.36	0.50	NIT 1	1	5		
Propionibacterium granulosum		3.23	0.22		99	90		
Atopobium minutum		3.20	0.20		99	1		
67011100	**Low discrimination**							**67011100**
					GRA	CAT	SPO	
Fusobacterium mortiferum		80.98	0.56	NIT 1	1	5	1	
Clostridium chauvoei		14.40	0.32		99	1	99	
Capnocytophaga ochracea		2.69	0.17		1	90	1	
67011101	**Very good identification**							**67011101**
					GRA	SPO		
Clostridium chauvoei		99.92	0.66	MLT 99	99	99		
Fusobacterium mortiferum		0.06	0.22		1	1		
Actinomyces naeslundii		0.02	0.25		99	1		
67011103	**Good identification**							**67011103**
					GRA	SPO		
Clostridium chauvoei		99.71	0.53	MLT 99	99	99		
Actinomyces naeslundii		0.11	0.25		99	1		
Capnocytophaga ochracea		0.10	-.01		1	1		
67011120	**Low discrimination**							**67011120**
					GRA	CAT		
Clostridium perfringens		62.98	0.42	MLT 99	99	1		
Capnocytophaga ochracea		27.77	0.33		1	90		
Clostridium chauvoei		5.51	0.24		99	1		
67011121	**Very good identification**							**67011121**
					SPO			
Clostridium chauvoei		99.78	0.58	MLT 99	99			
Clostridium perfringens		0.12	0.08		84			
Actinomyces naeslundii		0.06	0.25		1			
67011122	**Low discrimination**							**67011122**
					GRA	CAT		
Capnocytophaga ochracea		88.82	0.50	MLT 99	1	90		
Clostridium perfringens		10.53	0.37	RAF 90	99	1		
Clostridium chauvoei		0.35	0.11		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67011123	**Good identification**						**67011123**
					GRA	SPO	
Clostridium chauvoei		96.85	0.45	MLT 99	99	99	
Capnocytophaga ochracea		2.54	0.15		1	1	
Actinomyces naeslundii		0.31	0.25		99	1	
67011522	**Good identification**						**67011522**
					GRA	CAT	
Capnocytophaga ochracea		99.33	0.37	MLT 99	1	90	
Clostridium perfringens		0.67	0.03	RAF 90	99	1	
67015100	**Acceptable identification**						**67015100**
					CAT		
Fusobacterium mortiferum		90.08	0.37	NIT 1	5		
Capnocytophaga ochracea		9.93	0.07	XYL 7	90		
67015122	**Good identification**						**67015122**
					GRA	CAT	
Capnocytophaga ochracea		95.81	0.39	MLT 99	1	90	
Bacteroides fragilis		3.73	0.06	RAF 90	1	99	
Clostridium perfringens		0.46	0.03		99	1	
67021101	**Good identification**						**67021101**
Clostridium chauvoei		100.0	0.43	MLT 99			
67021121	**Good identification**						**67021121**
Clostridium chauvoei		100.0	0.35	MLT 99			
67030000	**Low discrimination**						**67030000**
					GRA	CAT	
Fusobacterium mortiferum		64.77	0.38	NIT 1	1	5	
Propionibacterium granulosum		35.23	0.30		99	90	
67030100	**Good identification**						**67030100**
Fusobacterium mortiferum		100.0	0.44	NIT 1			
67031000	**Low discrimination**						**67031000**
					GRA	CAT	
Fusobacterium mortiferum		71.22	0.41	NIT 1	1	5	
Propionibacterium granulosum		25.83	0.30		99	90	
Clostridium perfringens		2.95	0.06		99	1	
67031100	**Low discrimination**						**67031100**
					COC	GRA	
Fusobacterium mortiferum		76.10	0.47	NIT 1	1	1	
Clostridium perfringens		9.91	0.21		1	99	
Atopobium parvulum		7.85	0.19		99	99	
67031101	**Good identification**						**67031101**
					GRA	SPO	
Clostridium chauvoei		99.52	0.50	MLT 99	99	99	
Actinomyces naeslundii		0.33	0.30	TRE 10	99	1	
Fusobacterium mortiferum		0.15	0.13		1	1	
67031103	**Good identification**						**67031103**
					SPO		
Clostridium chauvoei		98.15	0.37	MLT 99	99		
Actinomyces naeslundii		1.85	0.30	TRE 10	1		
67031120	**Good identification**						**67031120**
					COC	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		98.25	0.47	MLT 99	1	84	
Clostridium chauvoei		0.51	0.07		1	99	
Atopobium parvulum		0.43	0.06		99	1	
67031121	**Good identification**						**67031121**
					SPO		
Clostridium chauvoei		97.14	0.41	MLT 99	99		
Clostridium perfringens		1.89	0.13	TRE 10	84		
Actinomyces naeslundii		0.97	0.30		1		
67031122	**Acceptable identification**						**67031122**
					GRA	CAT	
Clostridium perfringens		94.95	0.41	MLT 99	99	1	
Capnocytophaga ochracea		4.35	0.15		1	90	
Actinomyces naeslundii		0.70	0.27		99	25	
67033103	**Low discrimination**						**67033103**
					SPO		
Actinomyces naeslundii		78.38	0.35	ESL 12	1		
Clostridium chauvoei		21.62	0.03	MLT 94	99		
67033121	**Low discrimination**						**67033121**
					SPO		
Actinomyces naeslundii		65.75	0.35	ESL 12	1		
Clostridium chauvoei		34.25	0.07	MLT 94	99		
67033123	**Good identification**						**67033123**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67050000	**Good identification**						**67050000**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
67050100	**Good identification**						**67050100**
Fusobacterium mortiferum		100.0	0.45	NIT 1			
67051000	**Low discrimination**						**67051000**
					GRA		
Fusobacterium mortiferum		85.62	0.42	NIT 1	1		
Lactobacillus jensenii		7.96	0.14		99		
Lactobacillus catenaformis		6.42	0.12		99		
67051100	**Low discrimination**						**67051100**
					GRA		
Fusobacterium mortiferum		88.20	0.48	NIT 1	1		
Lactobacillus catenaformis		4.25	0.15		99		
Collinsella aerofaciens		2.81	-.03		99		
67051122	**Low discrimination**						**67051122**
					GRA	CAT	
Capnocytophaga ochracea		87.61	0.39	MLT 99	1	90	
Clostridium barati		7.80	0.20	RAF 90	99	1	
Clostridium perfringens		1.73	0.13		99	1	
67070100	**Good identification**						**67070100**
Fusobacterium mortiferum		100.0	0.36	NIT 1			
67071100	**Low discrimination**						**67071100**
					COC	GRA	
Fusobacterium mortiferum		56.19	0.39	NIT 1	1	1	
Atopobium parvulum		40.57	0.25		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		3.25	0.29		1	99	
67073101	**Good identification**						**67073101**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67073103	**Good identification**						**67073103**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67073121	**Good identification**						**67073121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67073123	**Good identification**						**67073123**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67100101	**Very good identification**						**67100101**
Clostridium chauvoei		100.0	0.60	GAL 99			
67100103	**Good identification**						**67100103**
					SPO		
Clostridium chauvoei		98.31	0.47	GAL 99	99		
Actinomyces viscosus		1.69	0.15		1		
67100121	**Good identification**						**67100121**
Clostridium chauvoei		100.0	0.51	GAL 99			
67100123	**Good identification**						**67100123**
Clostridium chauvoei		100.0	0.39	GAL 99			
67101001	**Very good identification**						**67101001**
Clostridium chauvoei		100.0	0.60	LAC 99			
67101003	**Good identification**						**67101003**
					SPO		
Clostridium chauvoei		99.41	0.47	LAC 99	99		
Actinomyces viscosus		0.59	0.07		1		
67101021	**Good identification**						**67101021**
Clostridium chauvoei		100.0	0.51	LAC 99			
67101023	**Good identification**						**67101023**
Clostridium chauvoei		100.0	0.39	LAC 99			
67101100	**Very good identification**						**67101100**
					GRA	CAT	
Clostridium chauvoei		99.14	0.60	URE 99	99	1	
Capnocytophaga ochracea		0.44	0.17		1	90	
Clostridium perfringens		0.28	0.17		99	1	
67101101	**Excellent identification**						**67101101**
					SPO		
Clostridium chauvoei		100.0	0.94		99		
Actinomyces naeslundii		0.00	0.24		1		
67101102	**Low discrimination**						**67101102**
					GRA	SPO	
Clostridium chauvoei		79.05	0.47	URE 99	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		17.73	0.33		1	1	
Actinomyces viscosus		2.50	0.20		99	1	
67101103	**Very good identification**						**67101103**
					SPO		
Clostridium chauvoei		99.99	0.81		99		
Actinomyces viscosus		0.01	0.12		1		
Actinomyces naeslundii		0.00	0.24		1		
67101111	**Very good identification**						**67101111**
Clostridium chauvoei		100.0	0.60	MLZ 1			
67101113	**Good identification**						**67101113**
Clostridium chauvoei		100.0	0.47	MLZ 1			
67101120	**Low discrimination**						**67101120**
					GRA	SPO	
Clostridium chauvoei		65.80	0.51	URE 99	99	99	
Clostridium perfringens		17.72	0.42		99	84	
Prevotella melaninogenica		8.67	0.30		1	1	
67101121	**Very good identification**						**67101121**
					SPO		
Clostridium chauvoei		99.99	0.86		99		
Clostridium perfringens		0.00	0.08		84		
Actinomyces naeslundii		0.00	0.24		1		
67101122	**Low discrimination**						**67101122**
					GRA	CAT	
Capnocytophaga ochracea		76.42	0.50	RAF 90	1	90	
Clostridium chauvoei		12.62	0.39	FRU 99	99	1	
Clostridium perfringens		9.06	0.37		99	1	
67101123	**Very good identification**						**67101123**
					GRA	SPO	
Clostridium chauvoei		99.93	0.73		99	99	
Capnocytophaga ochracea		0.06	0.15		1	1	
Actinomyces naeslundii		0.01	0.24		99	1	
67101131	**Good identification**						**67101131**
Clostridium chauvoei		100.0	0.51	MLZ 1			
67101133	**Good identification**						**67101133**
					GRA	SPO	
Clostridium chauvoei		99.33	0.39	MLZ 1	99	99	
Capnocytophaga ochracea		0.68	-.01		1	1	
67101141	**Very good identification**						**67101141**
Clostridium chauvoei		100.0	0.60	SOR 1			
67101143	**Good identification**						**67101143**
Clostridium chauvoei		100.0	0.47	SOR 1			
67101161	**Good identification**						**67101161**
Clostridium chauvoei		100.0	0.51	SOR 1			
67101163	**Good identification**						**67101163**
					GRA	SPO	
Clostridium chauvoei		99.33	0.39	SOR 1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Capnocytophaga ochracea		0.68	-.01		1	1		
67101301	**Very good identification**							**67101301**
Clostridium chauvoei		100.0	0.60	RHA 1				
67101303	**Good identification**							**67101303**
Clostridium chauvoei		100.0	0.47	RHA 1				
67101321	**Good identification**							**67101321**
Clostridium chauvoei		100.0	0.51	RHA 1				
67101323	**Good identification**							**67101323**
Clostridium chauvoei		100.0	0.39	RHA 1				
67101501	**Very good identification**							**67101501**
Clostridium chauvoei		100.0	0.60	ARA 1				
67101503	**Good identification**							**67101503**
Clostridium chauvoei		100.0	0.47	ARA 1				
67101521	**Good identification**							**67101521**
Clostridium chauvoei		100.0	0.51	ARA 1				
67101522	**Good identification**							**67101522**
					GRA	SPO	CAT	
Capnocytophaga ochracea		97.33	0.37	RAF 90	1	1	90	
Bifidobacterium longum subsp. longum		1.09	0.12	FRU 99	99	1	1	
Clostridium chauvoei		0.92	0.04		99	99	1	
67101523	**Good identification**							**67101523**
					GRA	SPO		
Clostridium chauvoei		98.93	0.39	ARA 1	99	99		
Capnocytophaga ochracea		1.07	0.02		1	1		
67103101	**Very good identification**							**67103101**
					SPO			
Clostridium chauvoei		99.93	0.60	MAN 1	99			
Actinomyces naeslundii		0.07	0.29		1			
67103103	**Good identification**							**67103103**
					SPO			
Clostridium chauvoei		99.42	0.47	MAN 1	99			
Actinomyces naeslundii		0.40	0.29		1			
Propionibacterium propionicum		0.18	0.12		1			
67103121	**Good identification**							**67103121**
					SPO			
Clostridium chauvoei		99.79	0.51	MAN 1	99			
Actinomyces naeslundii		0.21	0.29		1			
67103123	**Good identification**							**67103123**
					SPO			
Clostridium chauvoei		98.80	0.39	MAN 1	99			
Actinomyces naeslundii		1.20	0.29		1			
67105101	**Very good identification**							**67105101**
Clostridium chauvoei		100.0	0.60	XYL 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67105103	**Good identification**						**67105103**
Clostridium chauvoei		100.0	0.47	XYL 1			
67105121	**Good identification**						**67105121**
Clostridium chauvoei		100.0	0.51	XYL 1			
67105122	**Good identification**						**67105122**
					GRA	SPO	
Capnocytophaga ochracea		95.20	0.39	RAF 90	1	1	
Bacteroides fragilis		3.71	0.06	FRU 99	1	1	
Clostridium chauvoei		0.64	0.04		99	99	
67105123	**Good identification**						**67105123**
					GRA	SPO	
Clostridium chauvoei		98.49	0.39	XYL 1	99	99	
Capnocytophaga ochracea		1.51	0.05		1	1	
67110000	**Low discrimination**						**67110000**
					COC	GRA	
Actinomyces odontolyticus		48.09	0.39	ESL 1	1	99	
Gemella morbillorum		34.52	0.32	MNS 9	99	99	
Fusobacterium mortiferum		8.06	0.31		1	1	
67110002	**Low discrimination**						**67110002**
					GRA		
Actinomyces viscosus		84.35	0.41	ESL 1	99		
Actinomyces odontolyticus		14.86	0.35	RAF 94	99		
Fusobacterium mortiferum		0.45	0.15		1		
67110101	**Very good identification**						**67110101**
					SPO		
Clostridium chauvoei		99.98	0.66	GAL 99	99		
Actinomyces viscosus		0.03	0.03		1		
67110102	**Low discrimination**						**67110102**
					GRA	CAT	
Actinomyces viscosus		76.05	0.45	ESL 1	99	76	
Capnocytophaga ochracea		17.40	0.33	RAF 94	1	90	
Actinomyces odontolyticus		2.38	0.27		99	10	
67110103	**Low discrimination**						**67110103**
					GRA	SPO	
Clostridium chauvoei		87.59	0.53	GAL 99	99	99	
Actinomyces viscosus		12.26	0.37		99	1	
Capnocytophaga ochracea		0.09	-.01		1	1	
67110120	**Low discrimination**						**67110120**
					GRA	CAT	
Clostridium perfringens		62.94	0.42	GAL 99	99	1	
Capnocytophaga ochracea		27.75	0.33		1	90	
Clostridium chauvoei		5.51	0.24		99	1	
67110121	**Very good identification**						**67110121**
					GRA	CAT	
Clostridium chauvoei		99.83	0.58	GAL 99	99	1	
Clostridium perfringens		0.12	0.08		99	1	
Capnocytophaga ochracea		0.05	-.01		1	90	
67110122	**Low discrimination**						**67110122**
					GRA	CAT	
Capnocytophaga ochracea		88.24	0.50	RAF 90	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		10.47	0.37	GAL 99	99	1	
Actinomyces viscosus		0.43	0.11		99	76	
67110123	**Good identification**						**67110123**
					GRA	SPO	
Clostridium chauvoei		96.75	0.45	GAL 99	99	99	
Capnocytophaga ochracea		2.54	0.15		1	1	
Actinomyces viscosus		0.41	0.03		99	1	
67110200	**Good identification**						**67110200**
Actinomyces odontolyticus		100.0	0.44	ESL 1			
67110202	**Good identification**						**67110202**
Actinomyces odontolyticus		97.13	0.40	ESL 1			
Actinomyces viscosus		2.87	0.07	MNS 9			
67110300	**Good identification**						**67110300**
Actinomyces odontolyticus		100.0	0.35	ESL 1			
67110522	**Good identification**						**67110522**
					GRA	CAT	
Capnocytophaga ochracea		99.33	0.37	RAF 90	1	90	
Clostridium perfringens		0.67	0.03	GAL 99	99	1	
67111000	**Low discrimination**						**67111000**
					SPO		
Clostridium perfringens		34.18	0.36	LAC 88	84		
Clostridium chauvoei		21.50	0.32	NAG 97	99		
Actinomyces odontolyticus		17.00	0.36		1		
67111001	**Very good identification**						**67111001**
					SPO		
Clostridium chauvoei		99.88	0.66	LAC 99	99		
Actinomyces naeslundii		0.11	0.38		1		
Clostridium perfringens		0.02	0.02		84		
67111002	**Low discrimination**						**67111002**
					GRA	CAT	
Actinomyces viscosus		43.67	0.38	ESL 1	99	76	
Capnocytophaga ochracea		28.99	0.33	RAF 94	1	90	
Clostridium perfringens		12.92	0.30		99	1	
67111003	**Acceptable identification**						**67111003**
					SPO		
Clostridium chauvoei		94.77	0.53	LAC 99	99		
Actinomyces viscosus		4.57	0.29		1		
Actinomyces naeslundii		0.56	0.38		1		
67111020	**Very good identification**						**67111020**
					GRA	CAT	
Clostridium perfringens		95.45	0.62	LAC 88	99	1	
Capnocytophaga ochracea		3.12	0.33		1	90	
Clostridium chauvoei		0.62	0.24		99	1	
67111021	**Very good identification**						**67111021**
					SPO		
Clostridium chauvoei		98.10	0.58	LAC 99	99		
Clostridium perfringens		1.54	0.28		84		
Actinomyces naeslundii		0.31	0.38		1		
67111022	**Low discrimination**						**67111022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Clostridium perfringens		60.95	0.56	LAC 88	99	1	
Capnocytophaga ochracea		38.07	0.50		1	90	
Prevotella oralis		0.31	0.14		1	1	
67111023	**Acceptable identification**						**67111023**
					GRA	CAT	
Clostridium chauvoei		92.08	0.45	LAC 99	99	1	
Clostridium perfringens		3.86	0.22		99	1	
Capnocytophaga ochracea		2.41	0.15		1	90	
67111060	**Good identification**						**67111060**
					GRA	CAT	
Clostridium perfringens		98.13	0.49	LAC 88	99	1	
Capnocytophaga ochracea		1.87	0.17		1	90	
67111062	**Low discrimination**						**67111062**
					GRA	CAT	
Clostridium perfringens		73.30	0.44	LAC 88	99	1	
Capnocytophaga ochracea		26.70	0.33		1	90	
67111100	**Low discrimination**						**67111100**
					GRA	CAT	
Clostridium chauvoei		75.63	0.66	URE 99	99	1	
Capnocytophaga ochracea		14.11	0.51		1	90	
Clostridium perfringens		8.91	0.51		99	1	
67111101	**Excellent identification**						**67111101**
					GRA	SPO	
Clostridium chauvoei		99.99	1.00		99	99	
Actinomyces naeslundii		0.00	0.46		99	1	
Capnocytophaga ochracea		0.00	0.17		1	1	
67111102	**Low discrimination**						**67111102**
					GRA	CAT	
Capnocytophaga ochracea		85.47	0.67	RAF 90	1	90	
Clostridium chauvoei		8.98	0.53	NAG 90	99	1	
Clostridium perfringens		2.82	0.45		99	1	
67111103	**Very good identification**						**67111103**
					GRA	SPO	
Clostridium chauvoei		99.80	0.87		99	99	
Capnocytophaga ochracea		0.10	0.33		1	1	
Actinomyces viscosus		0.09	0.34		99	1	
67111111	**Very good identification**						**67111111**
					GRA	SPO	
Clostridium chauvoei		99.96	0.66	MLZ 1	99	99	
Capnocytophaga ochracea		0.02	0.01		1	1	
Actinomyces naeslundii		0.02	0.24		99	1	
67111112	**Good identification**						**67111112**
					GRA	CAT	
Capnocytophaga ochracea		98.52	0.51	RAF 90	1	90	
Clostridium chauvoei		0.94	0.19	MLZ 10	99	1	
Clostridium perfringens		0.30	0.11	NAG 90	99	1	
67111113	**Good identification**						**67111113**
					GRA	SPO	
Clostridium chauvoei		98.77	0.53	MLZ 1	99	99	
Capnocytophaga ochracea		1.06	0.17		1	1	
Actinomyces naeslundii		0.09	0.24		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67111120	**Low discrimination**						**67111120**
					GRA	CAT	
Clostridium perfringens		63.53	0.77		99	1	
Capnocytophaga ochracea		28.01	0.67		1	90	
Clostridium chauvoei		5.56	0.58		99	1	
67111121	**Excellent identification**						**67111121**
					GRA	CAT	
Clostridium chauvoei		99.82	0.92		99	1	
Clostridium perfringens		0.12	0.42		99	1	
Capnocytophaga ochracea		0.05	0.33		1	90	
67111122	**Low discrimination**						**67111122**
					GRA	CAT	
Capnocytophaga ochracea		88.79	0.84	RAF 90	1	90	
Clostridium perfringens		10.53	0.71		99	1	
Clostridium chauvoei		0.35	0.45		99	1	
67111123	**Very good identification**						**67111123**
					GRA	CAT	
Clostridium chauvoei		97.09	0.79		99	1	
Capnocytophaga ochracea		2.55	0.50		1	90	
Clostridium perfringens		0.30	0.37		99	1	
67111130	**Low discrimination**						**67111130**
					GRA	CAT	
Capnocytophaga ochracea		81.08	0.51	RAF 90	1	90	
Clostridium perfringens		16.72	0.42	MLZ 10	99	1	
Clostridium chauvoei		1.46	0.24	BGL 90	99	1	
67111131	**Very good identification**						**67111131**
					GRA	CAT	
Clostridium chauvoei		99.28	0.58	MLZ 1	99	1	
Capnocytophaga ochracea		0.56	0.17		1	90	
Clostridium perfringens		0.12	0.08		99	1	
67111132	**Very good identification**						**67111132**
					GRA	CAT	
Capnocytophaga ochracea		98.87	0.67	RAF 90	1	90	
Clostridium perfringens		1.07	0.37	MLZ 10	99	1	
Clostridium chauvoei		0.04	0.11		99	1	
67111133	**Low discrimination**						**67111133**
					GRA	CAT	
Clostridium chauvoei		77.27	0.45	MLZ 1	99	1	
Capnocytophaga ochracea		22.28	0.33		1	90	
Clostridium perfringens		0.24	0.03		99	1	
67111140	**Low discrimination**						**67111140**
					GRA	CAT	
Clostridium perfringens		41.23	0.38	NAG 97	99	1	
Capnocytophaga ochracea		38.09	0.35		1	90	
Clostridium chauvoei		18.56	0.32		99	1	
67111141	**Very good identification**						**67111141**
					SPO		
Clostridium chauvoei		99.90	0.66	SOR 1	99		
Actinomyces naeslundii		0.05	0.33		1		
Clostridium perfringens		0.02	0.04		84		
67111142	**Acceptable identification**						**67111142**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		92.86	0.51	RAF 90	1	90	
Clostridium perfringens		5.26	0.33	NAG 90	99	1	
Clostridium chauvoei		0.89	0.19	SOR 10	99	1	
67111143	**Good identification**						**67111143**
					GRA	SPO	
Clostridium chauvoei		98.49	0.53	SOR 1	99	99	
Capnocytophaga ochracea		1.05	0.17		1	1	
Actinomyces naeslundii		0.29	0.33		99	1	
67111160	**Low discrimination**						**67111160**
					GRA	CAT	
Clostridium perfringens		79.07	0.64		99	1	
Capnocytophaga ochracea		20.34	0.51		1	90	
Clostridium chauvoei		0.37	0.24		99	1	
67111161	**Very good identification**						**67111161**
					GRA	CAT	
Clostridium chauvoei		97.16	0.58	SOR 1	99	1	
Clostridium perfringens		2.14	0.30		99	1	
Capnocytophaga ochracea		0.55	0.17		1	90	
67111162	**Low discrimination**						**67111162**
					GRA	CAT	
Capnocytophaga ochracea		83.04	0.67	RAF 90	1	90	
Clostridium perfringens		16.88	0.59	SOR 10	99	1	
Clostridium chauvoei		0.03	0.11		99	1	
67111163	**Low discrimination**						**67111163**
					GRA	CAT	
Clostridium chauvoei		73.75	0.45	SOR 1	99	1	
Capnocytophaga ochracea		21.27	0.33		1	90	
Clostridium perfringens		4.32	0.24		99	1	
67111172	**Good identification**						**67111172**
					GRA	CAT	
Capnocytophaga ochracea		98.19	0.51	RAF 90	1	90	
Clostridium perfringens		1.82	0.24	MLZ 10	99	1	
67111200	**Good identification**						**67111200**
Actinomyces odontolyticus		98.96	0.41	ESL 1			
Clostridium perfringens		1.04	0.02	MNS 9			
67111202	**Good identification**						**67111202**
					GRA	CAT	
Actinomyces odontolyticus		95.68	0.37	ESL 1	99	10	
Actinomyces viscosus		2.60	0.03	MNS 9	99	76	
Capnocytophaga ochracea		1.73	-.01		1	90	
67111301	**Very good identification**						**67111301**
Clostridium chauvoei		100.0	0.66	RHA 1			
67111303	**Good identification**						**67111303**
					GRA	SPO	
Clostridium chauvoei		99.82	0.53	RHA 1	99	99	
Capnocytophaga ochracea		0.10	-.01		1	1	
Actinomyces viscosus		0.09	-.01		99	1	
67111320	**Low discrimination**						**67111320**
					GRA	CAT	
Clostridium perfringens		62.15	0.42	RHA 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Capnocytophaga ochracea		27.40	0.33		1	90		
Clostridium chauvoei		5.44	0.24		99	1		
67111321	**Very good identification**							**67111321**
					GRA	CAT		
Clostridium chauvoei		99.83	0.58	RHA 1	99	1		
Clostridium perfringens		0.12	0.08		99	1		
Capnocytophaga ochracea		0.05	-.01		1	90		
67111322	**Low discrimination**							**67111322**
					GRA	CAT		
Capnocytophaga ochracea		84.52	0.50	RAF 90	1	90		
Clostridium perfringens		10.02	0.37	RHA 1	99	1		
Prevotella oralis		4.67	0.28		1	1		
67111323	**Good identification**							**67111323**
					GRA	CAT		
Clostridium chauvoei		97.15	0.45	RHA 1	99	1		
Capnocytophaga ochracea		2.55	0.15		1	90		
Clostridium perfringens		0.30	0.03		99	1		
67111422	**Acceptable identification**							**67111422**
					GRA	CAT		
Capnocytophaga ochracea		91.61	0.37	RAF 90	1	90		
Clostridium perfringens		8.40	0.22	LAC 99	99	1		
67111500	**Low discrimination**							**67111500**
					GRA	CAT		
Capnocytophaga ochracea		72.75	0.38	RAF 90	1	90		
Clostridium chauvoei		22.32	0.32	NAG 90	99	1		
Clostridium perfringens		2.63	0.17	BGL 90	99	1		
67111501	**Very good identification**							**67111501**
					GRA	SPO		
Clostridium chauvoei		99.97	0.66	ARA 1	99	99		
Capnocytophaga ochracea		0.03	0.04		1	1		
67111502	**Good identification**							**67111502**
					GRA	SPO	CAT	
Capnocytophaga ochracea		98.62	0.54	RAF 90	1	1	90	
Clostridium chauvoei		0.59	0.19	NAG 90	99	99	1	
Bifidobacterium longum subsp. longum		0.23	0.18		99	1	1	
67111503	**Good identification**							**67111503**
					GRA	SPO		
Clostridium chauvoei		98.25	0.53	ARA 1	99	99		
Capnocytophaga ochracea		1.67	0.20		1	1		
Actinomyces viscosus		0.09	-.01		99	1		
67111512	**Good identification**							**67111512**
					GRA	CAT		
Capnocytophaga ochracea		95.50	0.38	RAF 90	1	90		
Bifidobacterium longum subsp. longum		3.37	0.22	MLZ 10	99	1		
Bifidobacterium adolescentis		1.13	0.15	NAG 90	99	1		
67111520	**Low discrimination**							**67111520**
					GRA	CAT		
Capnocytophaga ochracea		86.44	0.54	RAF 90	1	90		
Clostridium perfringens		11.22	0.42	BGL 90	99	1		
Clostridium chauvoei		0.98	0.24		99	1		
67111521	**Very good identification**							**67111521**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		99.00	0.58	ARA 1	99	1	
Capnocytophaga ochracea		0.89	0.20		1	90	
Clostridium perfringens		0.12	0.08		99	1	
67111522	**Very good identification**						**67111522**
					GRA	CAT	
Capnocytophaga ochracea		99.18	0.71	RAF 90	1	90	
Clostridium perfringens		0.67	0.37		99	1	
Bifidobacterium longum subsp. longum		0.10	0.28		99	1	
67111523	**Low discrimination**						**67111523**
					GRA	CAT	
Clostridium chauvoei		68.44	0.45	ARA 1	99	1	
Capnocytophaga ochracea		31.34	0.37		1	90	
Clostridium perfringens		0.21	0.03		99	1	
67111530	**Low discrimination**						**67111530**
					GRA	CAT	
Capnocytophaga ochracea		86.75	0.38	RAF 90	1	90	
Bifidobacterium longum subsp. longum		12.23	0.32	MLZ 10	99	1	
Clostridium perfringens		1.02	0.08	BGL 90	99	1	
67111532	**Good identification**						**67111532**
					GRA	CAT	
Capnocytophaga ochracea		98.40	0.54	RAF 90	1	90	
Bifidobacterium longum subsp. longum		1.54	0.32	MLZ 10	99	1	
Clostridium perfringens		0.06	0.03		99	1	
67111542	**Good identification**						**67111542**
Capnocytophaga ochracea		100.0	0.38	RAF 90			
67111560	**Low discrimination**						**67111560**
					GRA	CAT	
Capnocytophaga ochracea		81.80	0.38	RAF 90	1	90	
Clostridium perfringens		18.21	0.30	SOR 10	99	1	
67111562	**Good identification**						**67111562**
					GRA	CAT	
Capnocytophaga ochracea		98.85	0.54	RAF 90	1	90	
Clostridium perfringens		1.15	0.24	SOR 10	99	1	
67111572	**Good identification**						**67111572**
Capnocytophaga ochracea		100.0	0.38	RAF 90			
67111722	**Good identification**						**67111722**
					GRA	CAT	
Capnocytophaga ochracea		99.33	0.37	RAF 90	1	90	
Clostridium perfringens		0.67	0.03	RHA 1	99	1	
67112100	**Very good identification**						**67112100**
Propionibacterium propionicum		99.80	0.60	ESL 1			
Actinomyces naeslundii		0.20	0.24				
67112102	**Very good identification**						**67112102**
					GRA	CAT	
Propionibacterium propionicum		99.10	0.63	ESL 1	99	1	
Actinomyces viscosus		0.62	0.11		99	76	
Capnocytophaga ochracea		0.14	-.01		1	90	
67112103	**Low discrimination**						**67112103**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		85.32	0.47	ESL 1	1		
Clostridium chauvoei		11.51	0.19	URE 10	99		
Actinomyces viscosus		1.61	0.03		1		
67112140	**Very good identification**						**67112140**
Propionibacterium propionicum		100.0	0.60	ESL 1			
67112141	**Good identification**						**67112141**
Propionibacterium propionicum		100.0	0.44	ESL 1			
67112142	**Very good identification**						**67112142**
Propionibacterium propionicum		100.0	0.63	ESL 1			
67112143	**Good identification**						**67112143**
Propionibacterium propionicum		100.0	0.47	ESL 1			
67113000	**Low discrimination**						**67113000**
					CAT		
Actinomyces naeslundii		51.84	0.40	ESL 12	25		
Propionibacterium propionicum		30.59	0.26	RAF 92	1		
Propionibacterium granulosum		11.38	0.14		90		
67113002	**Low discrimination**						**67113002**
Actinomyces naeslundii		45.43	0.40	ESL 12			
Propionibacterium propionicum		40.21	0.29	RAF 92			
Actinomyces viscosus		8.63	0.03				
67113003	**Low discrimination**						**67113003**
					SPO		
Actinomyces naeslundii		51.52	0.43	ESL 12	1		
Clostridium chauvoei		45.10	0.19	RAF 92	99		
Propionibacterium propionicum		3.38	0.12		1		
67113023	**Low discrimination**						**67113023**
					SPO		
Actinomyces naeslundii		77.41	0.43	ESL 12	1		
Clostridium chauvoei		22.59	0.11	RAF 92	99		
67113100	**Low discrimination**						**67113100**
					SPO		
Propionibacterium propionicum		82.14	0.60	ESL 1	1		
Clostridium chauvoei		10.46	0.32		99		
Actinomyces naeslundii		4.00	0.48		1		
67113101	**Very good identification**						**67113101**
					SPO		
Clostridium chauvoei		98.56	0.66	MAN 1	99		
Propionibacterium propionicum		0.87	0.44		1		
Actinomyces naeslundii		0.57	0.51		1		
67113102	**Low discrimination**						**67113102**
					GRA	CAT	
Propionibacterium propionicum		83.25	0.63	ESL 1	99	1	
Capnocytophaga ochracea		11.87	0.33		1	90	
Actinomyces naeslundii		2.70	0.48		99	25	
67113103	**Acceptable identification**						**67113103**
					SPO		
Clostridium chauvoei		90.12	0.53	MAN 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		6.75	0.47		1		
Actinomyces naeslundii		2.96	0.51		1		
67113120	**Low discrimination**						**67113120**
					GRA	CAT	
Clostridium perfringens		59.03	0.42	MAN 1	99	1	
Capnocytophaga ochracea		26.02	0.33		1	90	
Actinomyces naeslundii		5.93	0.48		99	25	
67113121	**Very good identification**						**67113121**
					SPO		
Clostridium chauvoei		98.13	0.58	MAN 1	99		
Actinomyces naeslundii		1.71	0.51		1		
Clostridium perfringens		0.11	0.08		84		
67113122	**Low discrimination**						**67113122**
					GRA	CAT	
Capnocytophaga ochracea		86.27	0.50	RAF 90	1	90	
Clostridium perfringens		10.23	0.37	MAN 1	99	1	
Actinomyces naeslundii		2.18	0.48		99	25	
67113123	**Low discrimination**						**67113123**
					GRA	SPO	
Clostridium chauvoei		88.49	0.45	MAN 1	99	99	
Actinomyces naeslundii		8.72	0.51		99	1	
Capnocytophaga ochracea		2.32	0.15		1	1	
67113140	**Very good identification**						**67113140**
Propionibacterium propionicum		98.61	0.60	ESL 1			
Actinomyces naeslundii		0.85	0.35				
Clostridium perfringens		0.28	0.04				
67113142	**Very good identification**						**67113142**
					GRA	CAT	
Propionibacterium propionicum		97.89	0.63	ESL 1	99	1	
Capnocytophaga ochracea		1.55	0.17		1	90	
Actinomyces naeslundii		0.56	0.35		99	25	
67113143	**Low discrimination**						**67113143**
					SPO		
Propionibacterium propionicum		82.49	0.47	ESL 1	1		
Clostridium chauvoei		11.13	0.19	URE 10	99		
Actinomyces naeslundii		6.38	0.38		1		
67113163	**Low discrimination**						**67113163**
					GRA	SPO	
Actinomyces naeslundii		53.22	0.38	ESL 12	99	1	
Clostridium chauvoei		30.92	0.11	RAF 92	99	99	
Capnocytophaga ochracea		8.91	-.01		1	1	
67113522	**Good identification**						**67113522**
					GRA	CAT	
Capnocytophaga ochracea		99.33	0.37	RAF 90	1	90	
Clostridium perfringens		0.67	0.03	MAN 1	99	1	
67114122	**Good identification**						**67114122**
					GRA	CAT	
Capnocytophaga ochracea		95.36	0.39	RAF 90	1	90	
Bacteroides fragilis		3.72	0.06	GAL 99	1	99	
Actinomyces odontolyticus		0.47	0.08		99	10	
67114200	**Good identification**						**67114200**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces odontolyticus		100.0	0.38	ESL 1				
67115022	**Low discrimination**							**67115022**
					GRA	CAT		
Capnocytophaga ochracea		89.44	0.39	RAF 90	1	90		
Clostridium perfringens		5.79	0.22	LAC 99	99	1		
Bacteroides fragilis		3.49	0.06		1	99		
67115100	**Low discrimination**							**67115100**
					COC	GRA	SPO	
Capnocytophaga ochracea		72.95	0.41	RAF 90	1	1	1	
Clostridium chauvoei		15.80	0.32	NAG 90	1	99	99	
Atopobium parvulum		5.08	0.23	BGL 90	99	99	1	
67115101	**Very good identification**							**67115101**
					GRA	SPO		
Clostridium chauvoei		99.95	0.66	XYL 1	99	99		
Capnocytophaga ochracea		0.05	0.07		1	1		
67115102	**Very good identification**							**67115102**
					GRA	SPO		
Capnocytophaga ochracea		98.28	0.57	RAF 90	1	1		
Clostridium chauvoei		0.42	0.19	NAG 90	99	99		
Bacteroides fragilis		0.35	0.06		1	1		
67115103	**Good identification**							**67115103**
					GRA	SPO		
Clostridium chauvoei		97.57	0.53	XYL 1	99	99		
Capnocytophaga ochracea		2.34	0.23		1	1		
Actinomyces viscosus		0.09	-.01		99	1		
67115112	**Good identification**							**67115112**
					GRA	CAT		
Capnocytophaga ochracea		99.17	0.41	RAF 90	1	90		
Bifidobacterium adolescentis		0.83	0.15	MLZ 10	99	1		
67115120	**Low discrimination**							**67115120**
					GRA	CAT		
Capnocytophaga ochracea		88.82	0.57	RAF 90	1	90		
Clostridium perfringens		8.14	0.42	BGL 90	99	1		
Bacteroides fragilis		1.64	0.18		1	99		
67115121	**Very good identification**							**67115121**
					GRA	CAT		
Clostridium chauvoei		98.63	0.58	XYL 1	99	1		
Capnocytophaga ochracea		1.26	0.23		1	90		
Clostridium perfringens		0.12	0.08		99	1		
67115122	**Very good identification**							**67115122**
					GRA	CAT		
Capnocytophaga ochracea		95.70	0.73	RAF 90	1	90		
Bacteroides fragilis		3.73	0.40		1	99		
Clostridium perfringens		0.46	0.37		99	1		
67115123	**Low discrimination**							**67115123**
					GRA	SPO		
Clostridium chauvoei		59.63	0.45	XYL 1	99	99		
Capnocytophaga ochracea		38.68	0.39		1	1		
Bacteroides fragilis		1.51	0.06		1	1		
67115130	**Good identification**							**67115130**
					GRA	CAT		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		98.88	0.41	RAF 90	1	90	
Clostridium perfringens		0.82	0.08	MLZ 10	99	1	
Bifidobacterium longum subsp. longum		0.30	0.06	BGL 90	99	1	
67115132	**Very good identification**						**67115132**
					GRA	CAT	
Capnocytophaga ochracea		99.57	0.57	RAF 90	1	90	
Bacteroides fragilis		0.35	0.06	MLZ 10	1	99	
Clostridium perfringens		0.04	0.03		99	1	
67115142	**Good identification**						**67115142**
					GRA	CAT	
Capnocytophaga ochracea		99.49	0.41	RAF 90	1	90	
Actinomyces israelii		0.51	0.18	NAG 90	99	1	
67115160	**Low discrimination**						**67115160**
					GRA	CAT	
Capnocytophaga ochracea		86.42	0.41	RAF 90	1	90	
Clostridium perfringens		13.58	0.30	SOR 10	99	1	
67115162	**Very good identification**						**67115162**
					GRA	CAT	
Capnocytophaga ochracea		98.84	0.57	RAF 90	1	90	
Clostridium perfringens		0.81	0.24	SOR 10	99	1	
Bacteroides fragilis		0.35	0.06		1	99	
67115172	**Good identification**						**67115172**
Capnocytophaga ochracea		100.0	0.41	RAF 90			
67115200	**Good identification**						**67115200**
Actinomyces odontolyticus		100.0	0.35	ESL 1			
67115322	**Acceptable identification**						**67115322**
					CAT		
Capnocytophaga ochracea		91.10	0.39	RAF 90	90		
Bacteroides fragilis		3.55	0.06	RHA 1	99		
Prevotella oralis		2.49	0.12		1		
67115502	**Acceptable identification**						**67115502**
					GRA	CAT	
Capnocytophaga ochracea		93.80	0.44	RAF 90	1	90	
Bifidobacterium adolescentis		2.71	0.28	NAG 90	99	1	
Bifidobacterium longum subsp. longum		2.59	0.26		99	1	
67115520	**Low discrimination**						**67115520**
					GRA	CAT	
Capnocytophaga ochracea		79.52	0.44	RAF 90	1	90	
Bacteroides vulgatus		11.28	0.24	BGL 90	1	3	
Bifidobacterium longum subsp. longum		8.78	0.36		99	1	
67115522	**Very good identification**						**67115522**
					GRA	CAT	
Capnocytophaga ochracea		98.54	0.61	RAF 90	1	90	
Bifidobacterium longum subsp. longum		1.21	0.36		99	1	
Bacteroides fragilis		0.22	0.06		1	99	
67115530	**Low discrimination**						**67115530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		62.85	0.40	NIT 1	99	1	
Capnocytophaga ochracea		37.15	0.28	RAF 97	1	90	
67115532	**Low discrimination**						**67115532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		84.18	0.44	RAF 90	1	90	
Bifidobacterium longum subsp. longum		15.82	0.40	MLZ 10	99	1	
67115562	**Good identification**						**67115562**
Capnocytophaga ochracea		100.0	0.44	RAF 90			
67115720	**Good identification**						**67115720**
					GRA	CAT	
Bacteroides vulgatus		98.05	0.40	NIT 1	1	3	
Eubacterium contortum		1.18	0.14	RAF 97	99	1	
Capnocytophaga ochracea		0.78	0.10		1	90	
67117122	**Good identification**						**67117122**
					GRA	CAT	
Capnocytophaga ochracea		95.27	0.39	RAF 90	1	90	
Bacteroides fragilis		3.71	0.06	MAN 1	1	99	
Actinomyces israelii		0.56	0.17		99	1	
67117502	**Low discrimination**						**67117502**
					GRA	CAT	
Actinomyces israelii		53.45	0.36	SAL 92	99	1	
Bifidobacterium adolescentis		32.61	0.26	CEL 93	99	1	
Capnocytophaga ochracea		13.94	0.10		1	90	
67120101	**Good identification**						**67120101**
Clostridium chauvoei		100.0	0.43	TRE 10			
67120121	**Good identification**						**67120121**
Clostridium chauvoei		100.0	0.35	TRE 10			
67121001	**Good identification**						**67121001**
Clostridium chauvoei		100.0	0.43	TRE 10			
67121021	**Good identification**						**67121021**
Clostridium chauvoei		100.0	0.35	TRE 10			
67121100	**Acceptable identification**						**67121100**
					COC	SPO	
Clostridium chauvoei		91.69	0.43	TRE 10	1	99	
Clostridium perfringens		4.26	0.21	URE 99	1	84	
Atopobium parvulum		3.37	0.19		99	1	
67121101	**Very good identification**						**67121101**
					SPO		
Clostridium chauvoei		99.99	0.77	TRE 10	99		
Actinomyces naeslundii		0.01	0.29		1		
67121103	**Very good identification**						**67121103**
					SPO		
Clostridium chauvoei		99.96	0.65	TRE 10	99		
Actinomyces naeslundii		0.04	0.29		1		
67121111	**Good identification**						**67121111**
Clostridium chauvoei		100.0	0.43	TRE 10			
67121120	**Low discrimination**						**67121120**
					COC	SPO	
Clostridium perfringens		81.01	0.47	FRU 99	1	84	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		18.00	0.35		1	99	
Atopobium parvulum		0.35	0.06		99	1	
67121121	**Very good identification**						**67121121**
					SPO		
Clostridium chauvoei		99.94	0.69	TRE 10	99		
Clostridium perfringens		0.05	0.13		84		
Actinomyces naeslundii		0.02	0.29		1		
67121122	**Low discrimination**						**67121122**
					GRA	CAT	
Clostridium perfringens		88.09	0.41	FRU 99	99	1	
Clostridium chauvoei		7.34	0.22		99	1	
Capnocytophaga ochracea		4.04	0.15		1	90	
67121123	**Very good identification**						**67121123**
					SPO		
Clostridium chauvoei		99.77	0.56	TRE 10	99		
Clostridium perfringens		0.12	0.07		84		
Actinomyces naeslundii		0.11	0.29		1		
67121131	**Good identification**						**67121131**
Clostridium chauvoei		100.0	0.35	TRE 10			
67121141	**Good identification**						**67121141**
Clostridium chauvoei		100.0	0.43	TRE 10			
67121161	**Good identification**						**67121161**
Clostridium chauvoei		99.14	0.35	TRE 10			
Clostridium perfringens		0.86	0.01	SOR 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67121301	**Good identification**						**67121301**
Clostridium chauvoei		100.0	0.43	TRE 10			
67121321	**Good identification**						**67121321**
Clostridium chauvoei		100.0	0.35	TRE 10			
67121501	**Good identification**						**67121501**
Clostridium chauvoei		100.0	0.43	TRE 10			
67121521	**Good identification**						**67121521**
Clostridium chauvoei		100.0	0.35	TRE 10			
67123101	**Good identification**						**67123101**
					SPO		
Clostridium chauvoei		98.77	0.43	TRE 10	99		
Actinomyces naeslundii		1.23	0.34	MAN 1	1		
67123121	**Good identification**						**67123121**
					SPO		
Clostridium chauvoei		96.40	0.35	TRE 10	99		
Actinomyces naeslundii		3.60	0.34	MAN 1	1		
67125101	**Good identification**						**67125101**
Clostridium chauvoei		100.0	0.43	TRE 10			
67125121	**Good identification**						**67125121**
Clostridium chauvoei		100.0	0.35	TRE 10			
67130101	**Good identification**						**67130101**
					SPO		
Clostridium chauvoei		99.78	0.50	TRE 10	99		
Actinomyces naeslundii		0.22	0.27	GAL 99	1		
67130103	**Acceptable identification**						**67130103**
					SPO		
Clostridium chauvoei		90.32	0.37	TRE 10	99		
Actinomyces viscosus		8.57	0.18	GAL 99	1		
Actinomyces naeslundii		1.11	0.27		1		
67130120	**Good identification**						**67130120**
					COC	SPO	
Clostridium perfringens		98.60	0.47	GAL 99	1	84	
Clostridium chauvoei		0.52	0.07		1	99	
Atopobium parvulum		0.43	0.06		99	1	
67130121	**Good identification**						**67130121**
					SPO		
Clostridium chauvoei		97.47	0.41	TRE 10	99		
Clostridium perfringens		1.90	0.13	GAL 99	84		
Actinomyces naeslundii		0.64	0.27		1		
67130122	**Good identification**						**67130122**
					GRA	CAT	
Clostridium perfringens		95.18	0.41	GAL 99	99	1	
Capnocytophaga ochracea		4.37	0.15		1	90	
Actinomyces naeslundii		0.46	0.24		99	25	
67131000	**Low discrimination**						**67131000**
					COC		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		77.28	0.41	LAC 88	1		
Lactobacillus jensenii		5.86	0.22	NAG 97	1		
Atopobium parvulum		4.53	0.19		99		
67131001	**Good identification**						**67131001**
					SPO		
Clostridium chauvoei		97.94	0.50	TRE 10	99		
Actinomyces naeslundii		1.80	0.43	LAC 99	1		
Clostridium perfringens		0.27	0.06		84		
67131003	**Low discrimination**						**67131003**
					SPO		
Clostridium chauvoei		87.43	0.37	TRE 10	99		
Actinomyces naeslundii		9.08	0.43	LAC 99	1		
Actinomyces viscosus		2.86	0.10		1		
67131020	**Very good identification**						**67131020**
Clostridium perfringens		99.68	0.66	LAC 88			
Actinomyces naeslundii		0.14	0.40				
Lactobacillus jensenii		0.06	0.12				
67131021	**Low discrimination**						**67131021**
					SPO		
Clostridium chauvoei		75.87	0.41	TRE 10	99		
Clostridium perfringens		19.96	0.32	LAC 99	84		
Actinomyces naeslundii		4.17	0.43		1		
67131022	**Very good identification**						**67131022**
					GRA	CAT	
Clostridium perfringens		99.28	0.61	LAC 88	99	1	
Capnocytophaga ochracea		0.34	0.15		1	90	
Actinomyces naeslundii		0.30	0.40		99	25	
67131060	**Good identification**						**67131060**
Clostridium perfringens		99.87	0.54	LAC 88			
Actinomyces naeslundii		0.13	0.27				
67131062	**Good identification**						**67131062**
					GRA	CAT	
Clostridium perfringens		99.53	0.48	LAC 88	99	1	
Actinomyces naeslundii		0.28	0.27		99	25	
Capnocytophaga ochracea		0.20	-.01		1	90	
67131100	**Low discrimination**						**67131100**
					COC	SPO	
Clostridium perfringens		42.71	0.55	NAG 97	1	84	
Atopobium parvulum		33.83	0.53		99	1	
Clostridium chauvoei		21.69	0.50		1	99	
67131101	**Very good identification**						**67131101**
					SPO		
Clostridium chauvoei		99.91	0.84	TRE 10	99		
Actinomyces naeslundii		0.05	0.51		1		
Clostridium perfringens		0.02	0.21		84		
67131102	**Low discrimination**						**67131102**
					COC	SPO	
Clostridium perfringens		57.58	0.50	NAG 97	1	84	
Atopobium parvulum		17.10	0.40		99	1	
Clostridium chauvoei		10.97	0.37		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67131103	**Very good identification**						**67131103**
					SPO		
Clostridium chauvoei		99.57	0.71	TRE 10	99		
Actinomyces naeslundii		0.30	0.51		1		
Actinomyces viscosus		0.06	0.14		1		
67131111	**Good identification**						**67131111**
					SPO		
Clostridium chauvoei		99.73	0.50	TRE 10	99		
Actinomyces naeslundii		0.27	0.29	MLZ 1	1		
67131113	**Good identification**						**67131113**
					SPO		
Clostridium chauvoei		98.47	0.37	TRE 10	99		
Actinomyces naeslundii		1.53	0.29	MLZ 1	1		
67131120	**Very good identification**						**67131120**
					COC	SPO	
Clostridium perfringens		98.72	0.81		1	84	
Clostridium chauvoei		0.52	0.41		1	99	
Atopobium parvulum		0.43	0.40		99	1	
67131121	**Very good identification**						**67131121**
					SPO		
Clostridium chauvoei		97.92	0.76	TRE 10	99		
Clostridium perfringens		1.91	0.47		84		
Actinomyces naeslundii		0.16	0.51		1		
67131122	**Very good identification**						**67131122**
					GRA	CAT	
Clostridium perfringens		95.14	0.76		99	1	
Capnocytophaga ochracea		4.36	0.50		1	90	
Clostridium chauvoei		0.19	0.28		99	1	
67131123	**Good identification**						**67131123**
					SPO		
Clostridium chauvoei		94.05	0.63	TRE 10	99		
Clostridium perfringens		4.89	0.41		84		
Actinomyces naeslundii		0.84	0.51		1		
67131130	**Good identification**						**67131130**
					GRA	CAT	
Clostridium perfringens		96.27	0.47	MLZ 1	99	1	
Capnocytophaga ochracea		2.54	0.17		1	90	
Clostridium chauvoei		0.50	0.07		99	1	
67131131	**Good identification**						**67131131**
					SPO		
Clostridium chauvoei		97.30	0.41	TRE 10	99		
Clostridium perfringens		1.90	0.13	MLZ 1	84		
Actinomyces naeslundii		0.80	0.29		1		
67131132	**Low discrimination**						**67131132**
					GRA	CAT	
Clostridium perfringens		65.48	0.41	MLZ 1	99	1	
Capnocytophaga ochracea		33.03	0.33		1	90	
Leptotrichia buccalis		1.09	-.01		1	1	
67131140	**Acceptable identification**						**67131140**
					COC	SPO	
Clostridium perfringens		90.41	0.43	NAG 97	1	84	
Atopobium parvulum		3.80	0.19		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		2.44	0.15		1	99	
67131141	**Good identification**						**67131141**
					SPO		
Clostridium chauvoei		98.72	0.50	TRE 10	99		
Actinomyces naeslundii		0.91	0.38	SOR 1	1		
Clostridium perfringens		0.37	0.09		84		
67131142	**Low discrimination**						**67131142**
					GRA	CAT	
Clostridium perfringens		82.86	0.37	NAG 97	99	1	
Capnocytophaga ochracea		7.96	0.17		1	90	
Propionibacterium propionicum		4.15	0.27		99	1	
67131143	**Acceptable identification**						**67131143**
					SPO		
Clostridium chauvoei		94.14	0.37	TRE 10	99		
Actinomyces naeslundii		4.91	0.38	SOR 1	1		
Clostridium perfringens		0.95	0.03		84		
67131160	**Very good identification**						**67131160**
					GRA	CAT	
Clostridium perfringens		99.76	0.69		99	1	
Capnocytophaga ochracea		0.14	0.17		1	90	
Actinomyces naeslundii		0.05	0.35		99	25	
67131161	**Low discrimination**						**67131161**
					SPO		
Clostridium chauvoei		71.68	0.41	TRE 10	99		
Clostridium perfringens		26.34	0.35	SOR 1	84		
Actinomyces naeslundii		1.98	0.38		1		
67131162	**Very good identification**						**67131162**
					GRA	CAT	
Clostridium perfringens		97.29	0.63		99	1	
Capnocytophaga ochracea		2.60	0.33		1	90	
Actinomyces naeslundii		0.11	0.35		99	25	
67131301	**Good identification**						**67131301**
Clostridium chauvoei		100.0	0.50	TRE 10			
67131303	**Good identification**						**67131303**
Clostridium chauvoei		100.0	0.37	TRE 10			
67131320	**Good identification**						**67131320**
					COC	SPO	
Clostridium perfringens		98.82	0.47	RHA 1	1	84	
Clostridium chauvoei		0.52	0.07		1	99	
Atopobium parvulum		0.43	0.06		99	1	
67131321	**Good identification**						**67131321**
Clostridium chauvoei		98.09	0.41	TRE 10			
Clostridium perfringens		1.91	0.13	RHA 1			
67131322	**Acceptable identification**						**67131322**
					GRA	CAT	
Clostridium perfringens		93.47	0.41	RHA 1	99	1	
Capnocytophaga ochracea		4.29	0.15		1	90	
Prevotella oralis		1.50	0.07		1	1	
67131501	**Good identification**						**67131501**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		100.0	0.50	TRE 10			
67131503	**Good identification**						**67131503**
Clostridium chauvoei		100.0	0.37	TRE 10			
67131520	**Acceptable identification**						**67131520**
					GRA	CAT	
Clostridium perfringens		94.25	0.47	ARA 1	99	1	
Capnocytophaga ochracea		3.95	0.20		1	90	
Bifidobacterium longum subsp. longum		0.90	0.18		99	1	
67131521	**Good identification**						**67131521**
Clostridium chauvoei		98.09	0.41	TRE 10			
Clostridium perfringens		1.91	0.13	ARA 1			
67131522	**Low discrimination**						**67131522**
					GRA	CAT	
Clostridium perfringens		54.90	0.41	ARA 1	99	1	
Capnocytophaga ochracea		43.99	0.37		1	90	
Bifidobacterium longum subsp. longum		1.11	0.18		99	1	
67132100	**Very good identification**						**67132100**
Propionibacterium propionicum		99.52	0.58	ESL 1			
Actinomyces naeslundii		0.48	0.29				
67132101	**Low discrimination**						**67132101**
					SPO		
Propionibacterium propionicum		81.92	0.42	ESL 1	1		
Clostridium chauvoei		12.75	0.15	URE 10	99		
Actinomyces naeslundii		5.33	0.32		1		
67132102	**Very good identification**						**67132102**
Propionibacterium propionicum		99.68	0.61	ESL 1			
Actinomyces naeslundii		0.32	0.29				
67132103	**Acceptable identification**						**67132103**
					SPO		
Propionibacterium propionicum		94.19	0.45	ESL 1	1		
Actinomyces naeslundii		4.08	0.32	URE 10	1		
Clostridium chauvoei		1.73	0.03		99		
67132140	**Very good identification**						**67132140**
Propionibacterium propionicum		100.0	0.58	ESL 1			
67132141	**Good identification**						**67132141**
Propionibacterium propionicum		100.0	0.42	ESL 1			
67132142	**Very good identification**						**67132142**
Propionibacterium propionicum		100.0	0.61	ESL 1			
67132143	**Good identification**						**67132143**
Propionibacterium propionicum		100.0	0.45	ESL 1			
67133000	**Low discrimination**						**67133000**
					CAT		
Actinomyces naeslundii		56.35	0.45	ESL 12	25		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Propionibacterium granulosum		19.11	0.22	RAF 92	90	
Propionibacterium propionicum		14.01	0.24		1	
67133001	**Low discrimination**					**67133001**
					SPO	
Actinomyces naeslundii		77.89	0.48	ESL 12	1	
Clostridium chauvoei		22.11	0.15	RAF 92	99	
67133002	**Low discrimination**					**67133002**
Actinomyces naeslundii		61.71	0.45	ESL 12		
Propionibacterium propionicum		23.02	0.27	RAF 92		
Actinomyces israelii		6.71	0.19			
67133003	**Good identification**					**67133003**
					SPO	
Actinomyces naeslundii		95.23	0.48	ESL 12	1	
Clostridium chauvoei		4.77	0.03	RAF 92	99	
67133021	**Acceptable identification**					**67133021**
					SPO	
Actinomyces naeslundii		91.36	0.48	ESL 12	1	
Clostridium chauvoei		8.65	0.07	RAF 92	99	
67133023	**Good identification**					**67133023**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67133100	**Low discrimination**					**67133100**
Propionibacterium propionicum		82.94	0.58	ESL 1		
Actinomyces naeslundii		9.59	0.53			
Clostridium perfringens		2.82	0.21			
67133101	**Low discrimination**					**67133101**
					SPO	
Clostridium chauvoei		85.75	0.50	TRE 10	99	
Actinomyces naeslundii		8.68	0.56	MAN 1	1	
Propionibacterium propionicum		5.56	0.42		1	
67133102	**Good identification**					**67133102**
Propionibacterium propionicum		90.38	0.61	ESL 1		
Actinomyces naeslundii		6.97	0.53			
Actinomyces israelii		1.07	0.29			
67133103	**Low discrimination**					**67133103**
					SPO	
Clostridium chauvoei		47.05	0.37	TRE 10	99	
Actinomyces naeslundii		27.00	0.56	MAN 1	1	
Propionibacterium propionicum		25.95	0.45		1	
67133120	**Low discrimination**					**67133120**
Clostridium perfringens		88.80	0.47	MAN 1		
Actinomyces naeslundii		9.32	0.53			
Propionibacterium propionicum		0.82	0.24			
67133121	**Low discrimination**					**67133121**
					SPO	
Clostridium chauvoei		75.57	0.41	TRE 10	99	
Actinomyces naeslundii		22.96	0.56	MAN 1	1	
Clostridium perfringens		1.47	0.13		84	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
67133122	**Low discrimination**							**67133122**
					GRA	CAT		
Clostridium perfringens		77.03	0.41	MAN 1	99	1		
Actinomyces naeslundii		17.19	0.53		99	25		
Capnocytophaga ochracea		3.53	0.15		1	90		
67133123	**Low discrimination**							**67133123**
					SPO			
Actinomyces naeslundii		62.07	0.56	ESL 12	1			
Clostridium chauvoei		36.06	0.28	RAF 92	99			
Clostridium perfringens		1.87	0.07		84			
67133140	**Very good identification**							**67133140**
Propionibacterium propionicum		96.95	0.58	ESL 1				
Actinomyces naeslundii		1.98	0.40					
Clostridium perfringens		0.63	0.09					
67133141	**Low discrimination**							**67133141**
					SPO			
Propionibacterium propionicum		69.88	0.42	ESL 1	1			
Actinomyces naeslundii		19.24	0.43	URE 10	1			
Clostridium chauvoei		10.88	0.15		99			
67133142	**Very good identification**							**67133142**
Propionibacterium propionicum		98.02	0.61	ESL 1				
Actinomyces naeslundii		1.33	0.40					
Actinomyces israelii		0.45	0.22					
67133143	**Low discrimination**							**67133143**
					SPO			
Propionibacterium propionicum		83.20	0.45	ESL 1	1			
Actinomyces naeslundii		15.28	0.43	URE 10	1			
Clostridium chauvoei		1.52	0.03		99			
67133161	**Low discrimination**							**67133161**
					SPO			
Actinomyces naeslundii		79.52	0.43	ESL 12	1			
Clostridium chauvoei		14.98	0.07	RAF 92	99			
Clostridium perfringens		5.50	0.01		84			
67133163	**Good identification**							**67133163**
Actinomyces naeslundii		100.0	0.43	ESL 12				
67135100	**Very good identification**							**67135100**
					COC			
Atopobium parvulum		98.24	0.57	NIT 1	99			
Clostridium perfringens		0.68	0.21		1			
Actinomyces israelii		0.47	0.32		1			
67135101	**Good identification**							**67135101**
					COC	SPO		
Clostridium chauvoei		97.16	0.50	TRE 10	1	99		
Atopobium parvulum		2.84	0.23	XYL 1	99	1		
67135102	**Low discrimination**							**67135102**
					COC	GRA	CAT	
Atopobium parvulum		87.63	0.45	NIT 1	99	99	1	
Capnocytophaga ochracea		6.54	0.23		1	1	90	
Actinomyces israelii		3.59	0.35		1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
67135103	**Good identification**							**67135103**
					COC	SPO		
Clostridium chauvoei		97.16	0.37	TRE 10	1	99		
Atopobium parvulum		2.84	0.10	XYL 1	99	1		
67135120	**Low discrimination**							**67135120**
					COC	GRA	CAT	
Clostridium perfringens		53.78	0.47	XYL 1	1	99	1	
Atopobium parvulum		42.75	0.45		99	99	1	
Capnocytophaga ochracea		3.19	0.23		1	1	90	
67135121	**Good identification**							**67135121**
					COC	SPO		
Clostridium chauvoei		96.62	0.41	TRE 10	1	99		
Clostridium perfringens		1.88	0.13	XYL 1	1	84		
Atopobium parvulum		1.50	0.10		99	1		
67135122	**Low discrimination**							**67135122**
					COC	GRA	CAT	
Capnocytophaga ochracea		45.22	0.39	RAF 90	1	1	90	
Clostridium perfringens		39.84	0.41	TRE 1	1	99	1	
Atopobium parvulum		11.87	0.32		99	99	1	
67137002	**Good identification**							**67137002**
Actinomyces israelii		100.0	0.37	SAL 92				
67137100	**Low discrimination**							**67137100**
					COC			
Actinomyces israelii		69.24	0.45	SAL 92	1			
Atopobium parvulum		25.60	0.23	CEL 93	99			
Propionibacterium propionicum		5.16	0.24		1			
67137102	**Low discrimination**							**67137102**
					COC			
Actinomyces israelii		88.40	0.48	SAL 92	1			
Propionibacterium propionicum		6.59	0.27	CEL 93	1			
Atopobium parvulum		3.85	0.10		99			
67137140	**Low discrimination**							**67137140**
Actinomyces israelii		83.91	0.38	SAL 92				
Propionibacterium propionicum		16.09	0.24	CEL 93				
67137142	**Low discrimination**							**67137142**
Actinomyces israelii		83.91	0.41	SAL 92				
Propionibacterium propionicum		16.09	0.27	CEL 93				
67137402	**Good identification**							**67137402**
Actinomyces israelii		100.0	0.36	SAL 92				
67137500	**Good identification**							**67137500**
Actinomyces israelii		100.0	0.43	SAL 92				
67137502	**Acceptable identification**							**67137502**
Actinomyces israelii		90.24	0.46	SAL 92				
Bifidobacterium adolescentis		9.76	0.23	CEL 93				
67137540	**Good identification**							**67137540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.36	SAL 92			
67137542	**Good identification**						**67137542**
Actinomyces israelii		96.68	0.39	SAL 92			
Bifidobacterium adolescentis		3.32	0.08	CEL 93			
67141101	**Very good identification**						**67141101**
					SPO		
Clostridium chauvoei		99.95	0.60	CEL 1	99		
Actinomyces naeslundii		0.05	0.26		1		
67141103	**Good identification**						**67141103**
					SPO		
Clostridium chauvoei		99.45	0.47	CEL 1	99		
Actinomyces viscosus		0.28	0.02		1		
Actinomyces naeslundii		0.27	0.26		1		
67141121	**Good identification**						**67141121**
					SPO		
Clostridium chauvoei		99.85	0.51	CEL 1	99		
Actinomyces naeslundii		0.15	0.26		1		
67141122	**Acceptable identification**						**67141122**
					GRA	CAT	
Capnocytophaga ochracea		93.70	0.39	RAF 90	1	90	
Clostridium perfringens		1.85	0.13	FRU 99	99	1	
Prevotella oralis		1.73	0.09		1	1	
67141123	**Good identification**						**67141123**
					GRA	SPO	
Clostridium chauvoei		97.70	0.39	CEL 1	99	99	
Capnocytophaga ochracea		1.49	0.05		1	1	
Actinomyces naeslundii		0.81	0.26		99	1	
67150102	**Low discrimination**						**67150102**
					GRA	SPO	
Actinomyces viscosus		80.72	0.36	ESL 1	99	1	
Capnocytophaga ochracea		17.38	0.23	RAF 94	1	1	
Clostridium barati		1.19	0.02		99	99	
67150122	**Acceptable identification**						**67150122**
					GRA	CAT	
Capnocytophaga ochracea		93.45	0.39	RAF 90	1	90	
Clostridium barati		2.14	0.10	GAL 99	99	1	
Clostridium perfringens		1.85	0.13		99	1	
67151000	**Low discrimination**						**67151000**
					GRA		
Lactobacillus jensenii		59.47	0.36	NIT 1	99		
Lactobacillus catenaformis		22.73	0.29	SAL 95	99		
Fusobacterium mortiferum		4.16	0.26		1		
67151020	**Low discrimination**						**67151020**
					GRA	CAT	
Clostridium perfringens		62.97	0.38	CEL 4	99	1	
Capnocytophaga ochracea		12.34	0.23	LAC 88	1	90	
Lactobacillus jensenii		10.43	0.26		99	1	
67151022	**Low discrimination**						**67151022**
					GRA	CAT	
Capnocytophaga ochracea		60.29	0.39	RAF 90	1	90	
Clostridium perfringens		16.09	0.33	LAC 99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		8.74	0.24		99	1	
67151023	**Low discrimination**						**67151023**
					GRA	SPO	
Actinomyces naeslundii		58.67	0.40	ESL 12	99	1	
Clostridium chauvoei		25.07	0.11	RAF 92	99	99	
Capnocytophaga ochracea		16.26	0.05		1	1	
67151101	**Very good identification**						**67151101**
					SPO		
Clostridium chauvoei		99.50	0.66	CEL 1	99		
Actinomyces naeslundii		0.39	0.48		1		
Collinsella aerofaciens		0.06	-.03		1		
67151102	**Good identification**						**67151102**
					GRA	CAT	
Capnocytophaga ochracea		90.07	0.57	RAF 90	1	90	
Actinomyces viscosus		2.59	0.32	NAG 90	99	76	
Lactobacillus catenaformis		1.76	0.32		99	1	
67151103	**Acceptable identification**						**67151103**
					GRA	SPO	
Clostridium chauvoei		93.52	0.53	CEL 1	99	99	
Capnocytophaga ochracea		2.25	0.23		1	1	
Actinomyces viscosus		2.13	0.24		99	1	
67151112	**Acceptable identification**						**67151112**
					GRA	CAT	
Capnocytophaga ochracea		90.02	0.41	RAF 90	1	90	
Leptotrichia buccalis		9.71	0.15	MLZ 10	1	1	
Actinomyces naeslundii		0.27	0.23	NAG 90	99	25	
67151120	**Low discrimination**						**67151120**
					GRA	CAT	
Capnocytophaga ochracea		63.87	0.57	RAF 90	1	90	
Clostridium perfringens		24.14	0.53	BGL 90	99	1	
Clostridium paraputrificum		5.09	0.32		99	1	
67151121	**Very good identification**						**67151121**
					GRA	SPO	
Clostridium chauvoei		96.99	0.58	CEL 1	99	99	
Capnocytophaga ochracea		1.23	0.23		1	1	
Actinomyces naeslundii		1.15	0.48		99	1	
67151122	**Very good identification**						**67151122**
					GRA	CAT	
Capnocytophaga ochracea		95.27	0.73	RAF 90	1	90	
Clostridium perfringens		1.88	0.47		99	1	
Prevotella oralis		1.76	0.43		1	1	
67151123	**Low discrimination**						**67151123**
					GRA	SPO	
Clostridium chauvoei		57.27	0.45	CEL 1	99	99	
Capnocytophaga ochracea		37.15	0.39		1	1	
Actinomyces naeslundii		3.85	0.48		99	1	
67151130	**Good identification**						**67151130**
					GRA	CAT	
Capnocytophaga ochracea		95.36	0.41	RAF 90	1	90	
Clostridium perfringens		3.28	0.19	MLZ 10	99	1	
Clostridium paraputrificum		0.69	-.03	BGL 90	99	1	
67151132	**Very good identification**						**67151132**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					GRA	CAT		
Capnocytophaga ochracea		99.38	0.57	RAF 90	1	90		
Clostridium perfringens		0.18	0.13	MLZ 10	99	1		
Prevotella oralis		0.17	0.09		1	1		
67151142	**Good identification**							**67151142**
					GRA	SPO		
Capnocytophaga ochracea		96.17	0.41	RAF 90	1	1		
Clostridium barati		0.99	0.05	NAG 90	99	99		
Actinomyces naeslundii		0.96	0.32	SOR 10	99	1		
67151160	**Low discrimination**							**67151160**
					GRA	CAT		
Capnocytophaga ochracea		57.92	0.41	RAF 90	1	90		
Clostridium perfringens		37.54	0.41	SOR 10	99	1		
Clostridium paraputrificum		3.13	0.13	BGL 90	99	1		
67151162	**Very good identification**							**67151162**
					GRA	CAT		
Capnocytophaga ochracea		95.83	0.57	RAF 90	1	90		
Clostridium perfringens		3.25	0.35	SOR 10	99	1		
Clostridium barati		0.33	0.14		99	1		
67151172	**Good identification**							**67151172**
					GRA	CAT		
Capnocytophaga ochracea		99.33	0.41	RAF 90	1	90		
Bifidobacterium breve		0.36	0.10	MLZ 10	99	1		
Clostridium perfringens		0.31	0.01	SOR 10	99	1		
67151322	**Low discrimination**							**67151322**
					GRA	CAT	SPO	
Capnocytophaga ochracea		54.19	0.39	RAF 90	1	90	1	
Prevotella oralis		42.49	0.37	RHA 1	1	1	1	
Clostridium barati		2.05	0.14		99	1	99	
67151502	**Good identification**							**67151502**
					GRA	CAT		
Capnocytophaga ochracea		98.32	0.44	RAF 90	1	90		
Bifidobacterium adolescentis		0.90	0.19	NAG 90	99	1		
Actinomyces israelii		0.78	0.24		99	1		
67151520	**Good identification**							**67151520**
					GRA	CAT		
Capnocytophaga ochracea		97.20	0.44	RAF 90	1	90		
Clostridium perfringens		2.10	0.19	BGL 90	99	1		
Clostridium paraputrificum		0.44	-.03		99	1		
67151522	**Very good identification**							**67151522**
					GRA	CAT		
Capnocytophaga ochracea		99.74	0.61	RAF 90	1	90		
Clostridium perfringens		0.11	0.13		99	1		
Prevotella oralis		0.11	0.09		1	1		
67151532	**Good identification**							**67151532**
Capnocytophaga ochracea		100.0	0.44	RAF 90				
67151562	**Good identification**							**67151562**
					GRA	CAT		
Capnocytophaga ochracea		99.81	0.44	RAF 90	1	90		
Clostridium perfringens		0.19	0.01	SOR 10	99	1		
67153000	**Low discrimination**							**67153000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		68.71	0.42	ESL 12			
Actinomyces israelii		25.77	0.25	RAF 92			
Lactobacillus jensenii		5.53	0.02				
67153001	**Good identification**						**67153001**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67153002	**Low discrimination**						**67153002**
					SPO		
Actinomyces naeslundii		60.55	0.42	ESL 12	1		
Actinomyces israelii		34.06	0.28	RAF 92	1		
Clostridium barati		5.39	-.03		99		
67153003	**Good identification**						**67153003**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67153020	**Low discrimination**						**67153020**
					SPO		
Actinomyces naeslundii		82.61	0.42	ESL 12	1		
Clostridium perfringens		10.03	0.04	RAF 92	84		
Clostridium barati		7.36	-.03		99		
67153021	**Good identification**						**67153021**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67153022	**Low discrimination**						**67153022**
					GRA	SPO	
Actinomyces naeslundii		67.51	0.42	ESL 12	99	1	
Clostridium barati		18.04	0.05	RAF 92	99	99	
Capnocytophaga ochracea		14.46	0.05		1	1	
67153023	**Good identification**						**67153023**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67153100	**Low discrimination**						**67153100**
Actinomyces naeslundii		51.96	0.50	ESL 12			
Actinomyces israelii		27.39	0.36	RAF 92			
Propionibacterium propionicum		8.13	0.26				
67153102	**Low discrimination**						**67153102**
					GRA	CAT	
Actinomyces naeslundii		33.57	0.50	ESL 12	99	25	
Capnocytophaga ochracea		27.79	0.23	RAF 92	1	90	
Actinomyces israelii		26.54	0.39		99	1	
67153103	**Low discrimination**						**67153103**
					SPO		
Actinomyces naeslundii		80.04	0.53	ESL 12	1		
Clostridium chauvoei		18.57	0.19	RAF 92	99		
Propionibacterium propionicum		1.39	0.12		1		
67153120	**Low discrimination**						**67153120**
					GRA	CAT	
Actinomyces naeslundii		41.61	0.50	ESL 12	99	25	
Capnocytophaga ochracea		34.45	0.23	RAF 92	1	90	
Clostridium perfringens		13.02	0.19		99	1	
67153121	**Low discrimination**						**67153121**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					SPO			
Actinomyces naeslundii		69.53	0.53	ESL 12	1			
Clostridium chauvoei		30.47	0.24	RAF 92	99			
67153122	**Low discrimination**							**67153122**
					GRA	SPO		
Capnocytophaga ochracea		80.40	0.39	RAF 90	1	1		
Actinomyces naeslundii		10.79	0.50	MAN 1	99	1		
Clostridium barati		3.04	0.14		99	99		
67153123	**Low discrimination**							**67153123**
					GRA	SPO		
Actinomyces naeslundii		88.69	0.53	ESL 12	99	1		
Clostridium chauvoei		6.86	0.11	RAF 92	99	99		
Capnocytophaga ochracea		4.45	0.05		1	1		
67153141	**Good identification**							**67153141**
Actinomyces naeslundii		100.0	0.40	ESL 12				
67153143	**Acceptable identification**							**67153143**
Actinomyces naeslundii		91.04	0.40	ESL 12				
Propionibacterium propionicum		8.96	0.12	RAF 92				
67153160	**Low discrimination**							**67153160**
					GRA	CAT		
Actinomyces naeslundii		53.79	0.37	ESL 12	99	25		
Capnocytophaga ochracea		28.04	0.07	RAF 92	1	90		
Clostridium perfringens		18.17	0.06		99	1		
67153161	**Good identification**							**67153161**
Actinomyces naeslundii		100.0	0.40	ESL 12				
67153163	**Good identification**							**67153163**
Actinomyces naeslundii		100.0	0.40	ESL 12				
67153502	**Low discrimination**							**67153502**
					GRA	CAT		
Actinomyces israelii		71.76	0.37	SAL 92	99	1		
Capnocytophaga ochracea		16.21	0.10	XYL 92	1	90		
Bifidobacterium adolescentis		12.04	0.18		99	1		
67155102	**Acceptable identification**							**67155102**
					COC	GRA	CAT	
Capnocytophaga ochracea		90.26	0.47	RAF 90	1	1	90	
Actinomyces israelii		7.07	0.44	NAG 90	1	99	1	
Atopobium parvulum		1.15	0.17		99	99	1	
67155120	**Acceptable identification**							**67155120**
					GRA	CAT		
Capnocytophaga ochracea		93.25	0.47	RAF 90	1	90		
Bacteroides fragilis		2.95	0.12	BGL 90	1	99		
Clostridium perfringens		1.42	0.19		99	1		
67155122	**Good identification**							**67155122**
					CAT			
Capnocytophaga ochracea		92.77	0.63	RAF 90	90			
Bacteroides fragilis		6.20	0.34		99			
Prevotella oralis		0.85	0.28		1			
67155132	**Good identification**							**67155132**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		99.40	0.47	RAF 90			
Bacteroides fragilis		0.60	-.01	MLZ 10			
67155162	**Good identification**						**67155162**
					GRA	CAT	
Capnocytophaga ochracea		99.07	0.47	RAF 90	1	90	
Bacteroides fragilis		0.60	-.01	SOR 10	1	99	
Actinomyces israelii		0.19	0.17		99	1	
67155500	**Low discrimination**						**67155500**
					GRA	CAT	
Actinomyces israelii		68.54	0.40	SAL 92	99	1	
Capnocytophaga ochracea		31.46	0.18		1	90	
67155522	**Good identification**						**67155522**
					GRA	CAT	
Capnocytophaga ochracea		99.36	0.50	RAF 90	1	90	
Bacteroides fragilis		0.38	-.01		1	99	
Actinomyces israelii		0.26	0.22		99	1	
67155542	**Low discrimination**						**67155542**
					GRA	CAT	
Actinomyces israelii		49.96	0.36	SAL 92	99	1	
Capnocytophaga ochracea		39.31	0.18		1	90	
Bifidobacterium adolescentis		10.73	0.18		99	1	
67157000	**Good identification**						**67157000**
Actinomyces israelii		100.0	0.44	SAL 92			
67157002	**Good identification**						**67157002**
Actinomyces israelii		100.0	0.47	SAL 92			
67157040	**Good identification**						**67157040**
Actinomyces israelii		100.0	0.37	SAL 92			
67157042	**Good identification**						**67157042**
Actinomyces israelii		100.0	0.40	SAL 92			
67157100	**Good identification**						**67157100**
Actinomyces israelii		100.0	0.54	SAL 92			
67157102	**Very good identification**						**67157102**
					GRA	CAT	
Actinomyces israelii		96.20	0.57	SAL 92	99	1	
Capnocytophaga ochracea		2.19	0.13		1	90	
Bifidobacterium adolescentis		1.15	0.18		99	1	
67157110	**Good identification**						**67157110**
					SPO		
Actinomyces israelii		97.20	0.40	SAL 92	1		
Clostridium tertium		2.80	-.01	MLZ 13	99		
67157112	**Low discrimination**						**67157112**
					SPO		
Actinomyces israelii		86.56	0.43	SAL 92	1		
Bifidobacterium adolescentis		8.45	0.19	MLZ 13	1		
Clostridium tertium		4.99	0.07		99		
67157140	**Good identification**						**67157140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.47	SAL 92			
67157142	**Good identification**						**67157142**
Actinomyces israelii		100.0	0.50	SAL 92			
67157152	**Good identification**						**67157152**
Actinomyces israelii		100.0	0.36	SAL 92			
67157302	**Good identification**						**67157302**
Actinomyces israelii		100.0	0.38	SAL 92			
67157400	**Good identification**						**67157400**
Actinomyces israelii		100.0	0.42	SAL 92			
67157402	**Good identification**						**67157402**
Actinomyces israelii		100.0	0.45	SAL 92			
67157440	**Good identification**						**67157440**
Actinomyces israelii		100.0	0.35	SAL 92			
67157442	**Good identification**						**67157442**
Actinomyces israelii		100.0	0.38	SAL 92			
67157500	**Good identification**						**67157500**
Actinomyces israelii		100.0	0.52	SAL 92			
67157502	**Acceptable identification**						**67157502**
					GRA	CAT	
Actinomyces israelii		90.70	0.55	SAL 92	99	1	
Bifidobacterium adolescentis		8.85	0.32		99	1	
Capnocytophaga ochracea		0.45	0.00		1	90	
67157510	**Good identification**						**67157510**
Actinomyces israelii		100.0	0.38	SAL 92			
67157512	**Low discrimination**						**67157512**
Actinomyces israelii		55.61	0.41	SAL 92			
Bifidobacterium adolescentis		44.39	0.33	MLZ 13			
67157540	**Good identification**						**67157540**
Actinomyces israelii		100.0	0.45	SAL 92			
67157542	**Good identification**						**67157542**
Actinomyces israelii		96.99	0.48	SAL 92			
Bifidobacterium adolescentis		3.01	0.16				
67157702	**Good identification**						**67157702**
Actinomyces israelii		100.0	0.36	SAL 92			
67161101	**Good identification**						**67161101**
					SPO		
Clostridium chauvoei		99.16	0.43	TRE 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		0.84	0.31	CEL 1	1	
67161121	**Good identification**					**67161121**
					SPO	
Clostridium chauvoei		97.52	0.35	TRE 10	99	
Actinomyces naeslundii		2.48	0.31	CEL 1	1	
67163101	**Low discrimination**					**67163101**
					SPO	
Actinomyces naeslundii		62.01	0.36	ESL 12	1	
Clostridium chauvoei		37.99	0.09	RAF 92	99	
67163103	**Good identification**					**67163103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67163121	**Low discrimination**					**67163121**
					SPO	
Actinomyces naeslundii		83.04	0.36	ESL 12	1	
Clostridium chauvoei		16.96	0.01	RAF 92	99	
67163123	**Good identification**					**67163123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67171000	**Low discrimination**					**67171000**
					COC	
Lactobacillus jensenii		84.52	0.44	NIT 1	1	
Atopobium parvulum		8.03	0.25	SAL 95	99	
Actinomyces naeslundii		3.54	0.42		1	
67171001	**Low discrimination**					**67171001**
					SPO	
Actinomyces naeslundii		63.43	0.45	ESL 12	1	
Clostridium chauvoei		26.37	0.15	RAF 92	99	
Lactobacillus jensenii		10.20	0.10		1	
67171002	**Low discrimination**					**67171002**
					COC	
Lactobacillus jensenii		85.39	0.41	NIT 1	1	
Actinomyces naeslundii		5.36	0.42	SAL 95	1	
Atopobium parvulum		2.15	0.12		99	
67171003	**Low discrimination**					**67171003**
					SPO	
Actinomyces naeslundii		80.40	0.45	ESL 12	1	
Lactobacillus jensenii		8.62	0.07	RAF 92	1	
Clostridium chauvoei		5.90	0.03		99	
67171010	**Good identification**					**67171010**
Lactobacillus jensenii		100.0	0.36	NIT 1		
67171020	**Low discrimination**					**67171020**
Clostridium perfringens		74.96	0.43	CEL 4		
Lactobacillus jensenii		20.05	0.34	LAC 88		
Actinomyces naeslundii		3.36	0.42			
67171021	**Low discrimination**					**67171021**
					SPO	
Actinomyces naeslundii		77.58	0.45	ESL 12	1	
Clostridium perfringens		11.67	0.09	RAF 92	84	
Clostridium chauvoei		10.75	0.07		99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
67171022	**Low discrimination**							**67171022**
Clostridium perfringens		64.15	0.37	CEL 4				
Lactobacillus jensenii		24.30	0.31	LAC 88				
Actinomyces naeslundii		6.11	0.42					
67171023	**Acceptable identification**							**67171023**
Actinomyces naeslundii		93.39	0.45	ESL 12				
Clostridium perfringens		6.61	0.03	RAF 92				
67171100	**Very good identification**							**67171100**
					COC			
Atopobium parvulum		95.39	0.59	NIT 1	99			
Clostridium perfringens		2.15	0.32		1			
Actinomyces naeslundii		1.21	0.50		1			
67171101	**Acceptable identification**							**67171101**
					COC	SPO		
Clostridium chauvoei		90.42	0.50	TRE 10	1	99		
Actinomyces naeslundii		6.25	0.53	CEL 1	1	1		
Atopobium parvulum		3.32	0.25		99	1		
67171102	**Low discrimination**							**67171102**
					COC			
Atopobium parvulum		71.31	0.46	NIT 1	99			
Lactobacillus acidophilus		5.22	0.25		1			
Actinomyces naeslundii		5.12	0.50		1			
67171103	**Low discrimination**							**67171103**
					COC	SPO		
Clostridium chauvoei		69.25	0.37	TRE 10	1	99		
Actinomyces naeslundii		27.14	0.53	CEL 1	1	1		
Atopobium parvulum		2.55	0.12		99	1		
67171112	**Good identification**							**67171112**
					COC	CAT	GRA	
Leptotrichia buccalis		99.63	0.50	NIT 10	1	1	1	
Atopobium parvulum		0.14	0.12	SAL 99	99	1	99	
Capnocytophaga ochracea		0.09	0.07		1	90	1	
67171120	**Low discrimination**							**67171120**
					COC	SPO		
Clostridium perfringens		75.30	0.58	CEL 4	1	84		
Atopobium parvulum		18.23	0.46		99	1		
Clostridium paraputrificum		2.27	0.22		1	99		
67171121	**Low discrimination**							**67171121**
					SPO			
Clostridium chauvoei		76.49	0.41	TRE 10	99			
Actinomyces naeslundii		15.87	0.53	CEL 1	1			
Clostridium perfringens		6.15	0.23		84			
67171122	**Low discrimination**							**67171122**
					COC	GRA	CAT	
Clostridium perfringens		65.78	0.52	CEL 4	1	99	1	
Capnocytophaga ochracea		18.10	0.39		1	1	90	
Atopobium parvulum		5.97	0.33		99	99	1	
67171123	**Low discrimination**							**67171123**
					SPO			
Actinomyces naeslundii		48.00	0.53	ESL 12	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		40.84	0.28	RAF 92	99		
Clostridium perfringens		8.75	0.18		84		
67171141	**Low discrimination**						**67171141**
					SPO		
Actinomyces naeslundii		54.71	0.40	ESL 12	1		
Clostridium chauvoei		45.29	0.15	RAF 92	99		
67171143	**Low discrimination**						**67171143**
					SPO		
Actinomyces naeslundii		87.25	0.40	ESL 12	1		
Clostridium chauvoei		12.75	0.03	RAF 92	99		
67171160	**Good identification**						**67171160**
					COC		
Clostridium perfringens		95.31	0.45	CEL 4	1		
Actinomyces naeslundii		1.54	0.37		1		
Atopobium parvulum		1.22	0.12		99		
67171161	**Low discrimination**						**67171161**
					SPO		
Actinomyces naeslundii		59.03	0.40	ESL 12	1		
Clostridium perfringens		24.69	0.11	RAF 92	84		
Clostridium chauvoei		16.29	0.07		99		
67171162	**Low discrimination**						**67171162**
					GRA	CAT	
Clostridium perfringens		81.30	0.40	CEL 4	99	1	
Capnocytophaga ochracea		13.05	0.23		1	90	
Actinomyces naeslundii		2.78	0.37		99	25	
67171163	**Low discrimination**						**67171163**
Actinomyces naeslundii		83.56	0.40	ESL 12			
Clostridium perfringens		16.44	0.05	RAF 92			
67173000	**Low discrimination**						**67173000**
Actinomyces naeslundii		54.02	0.47	ESL 12			
Actinomyces israelii		39.26	0.35	RAF 92			
Lactobacillus jensenii		6.72	0.10				
67173001	**Good identification**						**67173001**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67173002	**Low discrimination**						**67173002**
Actinomyces israelii		50.17	0.38	SAL 92			
Actinomyces naeslundii		46.02	0.47	XYL 92			
Lactobacillus jensenii		3.81	0.07				
67173003	**Good identification**						**67173003**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67173020	**Low discrimination**						**67173020**
Actinomyces naeslundii		86.02	0.47	ESL 12			
Clostridium perfringens		10.00	0.09	RAF 92			
Actinomyces israelii		3.99	0.15				
67173021	**Good identification**						**67173021**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	ESL 12		
67173022	**Low discrimination**					**67173022**
Actinomyces naeslundii		88.95	0.47	ESL 12		
Actinomyces israelii		6.19	0.18	RAF 92		
Clostridium perfringens		4.86	0.03			
67173023	**Good identification**					**67173023**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67173041	**Good identification**					**67173041**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67173043	**Good identification**					**67173043**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67173061	**Good identification**					**67173061**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67173063	**Good identification**					**67173063**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67173100	**Low discrimination**					**67173100**
					COC	
Actinomyces israelii		40.10	0.46	SAL 92	1	
Actinomyces naeslundii		39.26	0.55	XYL 92	1	
Atopobium parvulum		16.12	0.25		99	
67173101	**Good identification**					**67173101**
					SPO	
Actinomyces naeslundii		93.00	0.58	ESL 12	1	
Clostridium chauvoei		7.00	0.15	RAF 92	99	
67173102	**Low discrimination**					**67173102**
					SPO	
Actinomyces israelii		53.22	0.49	SAL 92	1	
Actinomyces naeslundii		34.73	0.55	XYL 92	1	
Clostridium tertium		5.14	0.17		99	
67173103	**Very good identification**					**67173103**
					SPO	
Actinomyces naeslundii		97.70	0.58	ESL 12	1	
Clostridium chauvoei		1.30	0.03	RAF 92	99	
Actinomyces israelii		1.01	0.14		1	
67173111	**Good identification**					**67173111**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67173113	**Good identification**					**67173113**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67173120	**Low discrimination**					**67173120**
					SPO	
Actinomyces naeslundii		59.84	0.55	ESL 12	1	
Clostridium perfringens		17.91	0.23	RAF 92	84	
Clostridium tertium		8.85	0.17		99	
67173121	**Very good identification**					**67173121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		97.55	0.58	ESL 12	1	
Clostridium chauvoei		2.45	0.07	RAF 92	99	
67173122	**Low discrimination**					**67173122**
					SPO	
Actinomyces naeslundii		47.76	0.55	ESL 12	1	
Clostridium tertium		21.19	0.25	RAF 92	99	
Clostridium ramosum		16.49	0.21		92	
67173123	**Very good identification**					**67173123**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67173131	**Good identification**					**67173131**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67173133	**Good identification**					**67173133**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67173140	**Low discrimination**					**67173140**
Actinomyces israelii		62.10	0.39	SAL 92		
Actinomyces naeslundii		27.59	0.42	XYL 92		
Propionibacterium propionicum		10.31	0.24			
67173141	**Good identification**					**67173141**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67173142	**Low discrimination**					**67173142**
Actinomyces israelii		68.39	0.42	SAL 92		
Actinomyces naeslundii		20.26	0.42	XYL 92		
Propionibacterium propionicum		11.35	0.27			
67173143	**Good identification**					**67173143**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67173160	**Low discrimination**					**67173160**
Actinomyces naeslundii		68.18	0.42	ESL 12		
Clostridium perfringens		22.03	0.11	RAF 92		
Actinomyces israelii		9.79	0.18			
67173161	**Good identification**					**67173161**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67173162	**Low discrimination**					**67173162**
Actinomyces naeslundii		73.12	0.42	ESL 12		
Actinomyces israelii		15.76	0.21	RAF 92		
Clostridium perfringens		11.12	0.05			
67173163	**Good identification**					**67173163**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67173402	**Good identification**					**67173402**
Actinomyces israelii		100.0	0.37	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
67173500	**Good identification**							**67173500**
Actinomyces israelii		100.0	0.44	SAL 92				
67173502	**Good identification**							**67173502**
Actinomyces israelii		97.11	0.47	SAL 92				
Bifidobacterium adolescentis		2.89	0.15	XYL 92				
67173540	**Good identification**							**67173540**
Actinomyces israelii		100.0	0.37	SAL 92				
67173542	**Good identification**							**67173542**
Actinomyces israelii		100.0	0.40	SAL 92				
67175002	**Low discrimination**							**67175002**
					COC			
Actinomyces israelii		82.60	0.44	SAL 92	1			
Atopobium parvulum		14.30	0.17		99			
Lactobacillus jensenii		3.09	0.07		1			
67175042	**Good identification**							**67175042**
Actinomyces israelii		100.0	0.37	SAL 92				
67175100	**Very good identification**							**67175100**
					COC	SPO		
Atopobium parvulum		97.31	0.64	NIT 1	99	1		
Actinomyces israelii		2.67	0.51		1	1		
Clostridium butyricum		0.02	-.07		1	99		
67175102	**Low discrimination**							**67175102**
					COC	GRA	CAT	
Atopobium parvulum		80.34	0.51	NIT 1	99	99	1	
Actinomyces israelii		18.75	0.54		1	99	1	
Capnocytophaga ochracea		0.64	0.13		1	1	90	
67175120	**Good identification**							**67175120**
					COC			
Atopobium parvulum		96.12	0.51	NIT 1	99			
Clostridium perfringens		2.16	0.23		1			
Actinomyces israelii		0.95	0.30		1			
67175122	**Low discrimination**							**67175122**
					COC	GRA	CAT	
Atopobium parvulum		61.67	0.38	NIT 1	99	99	1	
Capnocytophaga ochracea		25.16	0.29		1	1	90	
Actinomyces israelii		5.20	0.33		1	99	1	
67175140	**Low discrimination**							**67175140**
					COC			
Actinomyces israelii		51.38	0.44	SAL 92	1			
Atopobium parvulum		48.62	0.30		99			
67175142	**Low discrimination**							**67175142**
					COC			
Actinomyces israelii		89.98	0.47	SAL 92	1			
Atopobium parvulum		10.02	0.17		99			
67175400	**Good identification**							**67175400**
Actinomyces israelii		100.0	0.39	SAL 92				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67175402	**Good identification**						**67175402**
Actinomyces israelii		100.0	0.42	SAL 92			
67175442	**Good identification**						**67175442**
Actinomyces israelii		100.0	0.35	SAL 92			
67175500	**Low discrimination**						**67175500**
					COC	GRA	
Actinomyces israelii		60.60	0.49	SAL 92	1	99	
Atopobium parvulum		27.25	0.30		99	99	
Mitsuokella multacida		7.12	0.21		1	1	
67175502	**Low discrimination**						**67175502**
					COC		
Actinomyces israelii		82.24	0.52	SAL 92	1		
Bifidobacterium adolescentis		9.85	0.30		1		
Atopobium parvulum		4.35	0.17		99		
67175510	**Low discrimination**						**67175510**
					GRA		
Actinomyces israelii		79.24	0.35	SAL 92	99		
Mitsuokella multacida		20.76	0.13	MLZ 13	1		
67175512	**Low discrimination**						**67175512**
Actinomyces israelii		46.44	0.38	SAL 92			
Bifidobacterium adolescentis		45.50	0.32	MLZ 13			
Clostridium clostridioforme		8.06	0.16				
67175540	**Low discrimination**						**67175540**
					GRA		
Actinomyces israelii		76.81	0.42	SAL 92	99		
Mitsuokella multacida		23.20	0.21		1		
67175542	**Acceptable identification**						**67175542**
					GRA		
Actinomyces israelii		94.30	0.45	SAL 92	99		
Bifidobacterium adolescentis		3.59	0.15		99		
Mitsuokella multacida		2.11	0.04		1		
67177000	**Good identification**						**67177000**
Actinomyces israelii		100.0	0.53	SAL 92			
67177002	**Very good identification**						**67177002**
Actinomyces israelii		100.0	0.56	SAL 92			
67177010	**Good identification**						**67177010**
Actinomyces israelii		100.0	0.39	SAL 92			
67177012	**Good identification**						**67177012**
Actinomyces israelii		100.0	0.42	SAL 92			
67177022	**Good identification**						**67177022**
Actinomyces israelii		100.0	0.36	SAL 92			
67177040	**Good identification**						**67177040**
Actinomyces israelii		100.0	0.46	SAL 92			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67177042	**Good identification**						**67177042**
Actinomyces israelii		100.0	0.49	SAL 92			
67177052	**Good identification**						**67177052**
Actinomyces israelii		100.0	0.35	SAL 92			
67177100	**Good identification**						**67177100**
					COC	SPO	
Actinomyces israelii		93.39	0.64	SAL 92	1	1	
Atopobium parvulum		6.07	0.30		99	1	
Clostridium tertium		0.54	0.11		1	99	
67177102	**Very good identification**						**67177102**
					COC	SPO	
Actinomyces israelii		97.91	0.67	SAL 92	1	1	
Clostridium tertium		1.14	0.19		1	99	
Atopobium parvulum		0.75	0.17		99	1	
67177110	**Acceptable identification**						**67177110**
					SPO		
Actinomyces israelii		93.55	0.50	SAL 92	1		
Clostridium tertium		6.45	0.15	MLZ 13	99		
67177112	**Low discrimination**						**67177112**
					SPO		
Actinomyces israelii		86.56	0.53	SAL 92	1		
Clostridium tertium		11.94	0.24	MLZ 13	99		
Bifidobacterium adolescentis		1.50	0.16		1		
67177120	**Low discrimination**						**67177120**
					COC	SPO	
Actinomyces israelii		68.87	0.43	SAL 92	1	1	
Clostridium tertium		18.76	0.19	NAG 6	1	99	
Atopobium parvulum		12.37	0.17		99	1	
67177122	**Low discrimination**						**67177122**
					COC	SPO	
Actinomyces israelii		63.86	0.46	SAL 92	1	1	
Clostridium tertium		34.79	0.27	NAG 6	1	99	
Atopobium parvulum		1.35	0.04		99	1	
67177140	**Very good identification**						**67177140**
Actinomyces israelii		100.0	0.57	SAL 92			
67177142	**Very good identification**						**67177142**
Actinomyces israelii		100.0	0.60	SAL 92			
67177150	**Good identification**						**67177150**
Actinomyces israelii		100.0	0.43	SAL 92			
67177152	**Good identification**						**67177152**
Actinomyces israelii		100.0	0.46	SAL 92			
67177160	**Good identification**						**67177160**
Actinomyces israelii		100.0	0.36	SAL 92			
67177162	**Good identification**						**67177162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.39	SAL 92			
67177202	**Good identification**						**67177202**
Actinomyces israelii		100.0	0.37	SAL 92			
67177300	**Good identification**						**67177300**
Actinomyces israelii		100.0	0.45	SAL 92			
67177302	**Good identification**						**67177302**
Actinomyces israelii		100.0	0.48	SAL 92			
67177340	**Good identification**						**67177340**
Actinomyces israelii		100.0	0.38	SAL 92			
67177342	**Good identification**						**67177342**
Actinomyces israelii		100.0	0.41	SAL 92			
67177400	**Good identification**						**67177400**
Actinomyces israelii		100.0	0.52	SAL 92			
67177402	**Good identification**						**67177402**
Actinomyces israelii		100.0	0.55	SAL 92			
67177410	**Good identification**						**67177410**
Actinomyces israelii		100.0	0.38	SAL 92			
67177412	**Good identification**						**67177412**
Actinomyces israelii		100.0	0.41	SAL 92			
67177440	**Good identification**						**67177440**
Actinomyces israelii		100.0	0.45	SAL 92			
67177442	**Good identification**						**67177442**
Actinomyces israelii		100.0	0.48	SAL 92			
67177500	**Very good identification**						**67177500**
					GRA	SPO	
Actinomyces israelii		95.24	0.62	SAL 92	99	1	
Mitsuokella multacida		4.61	0.27		1	1	
Clostridium butyricum		0.16	-.07		99	99	
67177502	**Very good identification**						**67177502**
					GRA		
Actinomyces israelii		97.96	0.65	SAL 92	99		
Bifidobacterium adolescentis		1.69	0.29		99		
Mitsuokella multacida		0.35	0.11		1		
67177510	**Acceptable identification**						**67177510**
					GRA		
Actinomyces israelii		90.26	0.48	SAL 92	99		
Mitsuokella multacida		9.74	0.19	MLZ 13	1		
67177512	**Low discrimination**						**67177512**
Actinomyces israelii		82.89	0.51	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		11.73	0.30	MLZ 13			
Bifidobacterium dentium		4.73	0.06				
67177520	**Low discrimination**						**67177520**
					GRA		
Actinomyces israelii		84.08	0.42	SAL 92	99		
Mitsuokella multacida		15.93	0.17	NAG 6	1		
67177522	**Good identification**						**67177522**
Actinomyces israelii		100.0	0.45	SAL 92			
67177540	**Low discrimination**						**67177540**
					GRA		
Actinomyces israelii		88.94	0.55	SAL 92	99		
Mitsuokella multacida		11.06	0.27		1		
67177542	**Very good identification**						**67177542**
					GRA		
Actinomyces israelii		98.55	0.58	SAL 92	99		
Mitsuokella multacida		0.91	0.11		1		
Bifidobacterium adolescentis		0.54	0.13		99		
67177550	**Low discrimination**						**67177550**
					COC	GRA	
Actinomyces israelii		75.81	0.41	SAL 92	1	99	
Mitsuokella multacida		21.03	0.19	MLZ 13	1	1	
Blautia producta		3.16	-.03		99	99	
67177552	**Acceptable identification**						**67177552**
					GRA		
Actinomyces israelii		93.85	0.44	SAL 92	99		
Bifidobacterium adolescentis		4.22	0.15	MLZ 13	99		
Mitsuokella multacida		1.93	0.03		1		
67177562	**Good identification**						**67177562**
Actinomyces israelii		100.0	0.38	SAL 92			
67177602	**Good identification**						**67177602**
Actinomyces israelii		100.0	0.36	SAL 92			
67177702	**Acceptable identification**						**67177702**
					GRA		
Actinomyces israelii		91.88	0.46	SAL 92	99		
Mitsuokella multacida		8.12	0.15	RHA 7	1		
67177742	**Low discrimination**						**67177742**
					GRA		
Actinomyces israelii		81.48	0.39	SAL 92	99		
Mitsuokella multacida		18.52	0.15	RHA 7	1		
67201000	**Very good identification**						**67201000**
					GRA		
Eubacterium tortuosum		99.97	0.64	MNS 1	99		
Fusobacterium mortiferum		0.03	0.20		1		
67201002	**Good identification**						**67201002**
Eubacterium tortuosum		100.0	0.52	MNS 1			
67201020	**Good identification**						**67201020**
Eubacterium tortuosum		100.0	0.52	MNS 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67201022	**Good identification**						**67201022**
Eubacterium tortuosum		100.0	0.39	MNS 1			
67210000	**Good identification**						**67210000**
					GRA		
Fusobacterium mortiferum		98.74	0.52	NIT 1	1		
Actinomyces odontolyticus		0.74	0.10		99		
Eubacterium tortuosum		0.53	-.04		99		
67210002	**Good identification**						**67210002**
					GRA		
Fusobacterium mortiferum		96.05	0.35	NIT 1	1		
Actinomyces odontolyticus		3.95	0.06	BGL 10	99		
67210100	**Very good identification**						**67210100**
Fusobacterium mortiferum		100.0	0.58	NIT 1			
67210102	**Good identification**						**67210102**
					GRA		
Fusobacterium mortiferum		97.62	0.42	NIT 1	1		
Actinomyces viscosus		2.38	0.01	BGL 10	99		
67211000	**Low discrimination**						**67211000**
					GRA		
Fusobacterium mortiferum		72.01	0.55	NIT 1	1		
Eubacterium tortuosum		25.44	0.30		99		
Lactobacillus jensenii		1.15	0.14		99		
67211002	**Low discrimination**						**67211002**
					GRA		
Fusobacterium mortiferum		53.34	0.38	NIT 1	1		
Eubacterium tortuosum		29.93	0.17	BGL 10	99		
Lactobacillus catenaformis		6.18	0.12		99		
67211100	**Very good identification**						**67211100**
					GRA		
Fusobacterium mortiferum		97.52	0.61	NIT 1	1		
Lactobacillus catenaformis		0.81	0.15		99		
Collinsella aerofaciens		0.53	-.03		99		
67211102	**Low discrimination**						**67211102**
					GRA	SPO	
Fusobacterium mortiferum		77.85	0.45	NIT 1	1	1	
Clostridium barati		7.19	0.12	BGL 10	99	99	
Lactobacillus catenaformis		5.80	0.15		99	1	
67213101	**Good identification**						**67213101**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67213102	**Good identification**						**67213102**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67213103	**Good identification**						**67213103**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67213120	**Good identification**						**67213120**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67213121	**Good identification**						**67213121**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.40	ESL 12			
67213122	**Low discrimination**						**67213122**
					SPO		
Actinomyces naeslundii		74.75	0.37	ESL 12	1		
Clostridium barati		25.25	0.02	MLT 94	99		
67213123	**Good identification**						**67213123**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67214100	**Good identification**						**67214100**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
67215000	**Good identification**						**67215000**
					GRA		
Fusobacterium mortiferum		95.48	0.35	NIT 1	1		
Eubacterium tortuosum		4.53	-.04	XYL 7	99		
67215100	**Good identification**						**67215100**
Fusobacterium mortiferum		100.0	0.42	NIT 1			
67230000	**Good identification**						**67230000**
Fusobacterium mortiferum		100.0	0.43	NIT 1			
67230100	**Good identification**						**67230100**
Fusobacterium mortiferum		100.0	0.49	NIT 1			
67231000	**Low discrimination**						**67231000**
					GRA		
Fusobacterium mortiferum		83.75	0.46	NIT 1	1		
Lactobacillus jensenii		13.42	0.22		99		
Actinomyces naeslundii		1.82	0.29		99		
67231100	**Low discrimination**						**67231100**
					COC	GRA	
Fusobacterium mortiferum		89.79	0.52	NIT 1	1	1	
Atopobium parvulum		6.32	0.21		99	99	
Actinomyces naeslundii		2.38	0.37		1	99	
67231101	**Low discrimination**						**67231101**
					GRA	SPO	
Actinomyces naeslundii		47.41	0.40	ESL 12	99	1	
Clostridium chauvoei		40.57	0.15	MLT 94	99	99	
Fusobacterium mortiferum		12.03	0.18	RAF 92	1	1	
67231102	**Low discrimination**						**67231102**
					COC	GRA	
Fusobacterium mortiferum		66.10	0.36	NIT 1	1	1	
Actinomyces naeslundii		15.79	0.37	BGL 10	1	99	
Atopobium parvulum		7.39	0.08		99	99	
67231103	**Low discrimination**						**67231103**
					SPO		
Actinomyces naeslundii		86.88	0.40	ESL 12	1		
Clostridium chauvoei		13.12	0.03	MLT 94	99		
67231120	**Low discrimination**						**67231120**
					SPO		
Clostridium perfringens		89.61	0.37	MLT 99	84		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		4.39	0.37		1	
Clostridium septicum		2.29	0.03		99	
67231121	**Low discrimination**					**67231121**
					SPO	
Actinomyces naeslundii		70.29	0.40	ESL 12	1	
Clostridium chauvoei		20.05	0.07	MLT 94	99	
Clostridium perfringens		9.67	0.03	RAF 92	84	
67231123	**Good identification**					**67231123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67233001	**Good identification**					**67233001**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67233003	**Good identification**					**67233003**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67233021	**Good identification**					**67233021**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67233023	**Good identification**					**67233023**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67233100	**Low discrimination**					**67233100**
					GRA	
Actinomyces naeslundii		83.61	0.42	ESL 12	99	
Fusobacterium mortiferum		16.39	0.18	MLT 94	1	
67233101	**Good identification**					**67233101**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67233102	**Acceptable identification**					**67233102**
Actinomyces naeslundii		92.72	0.42	ESL 12		
Actinomyces israelii		7.28	0.13	MLT 94		
67233103	**Good identification**					**67233103**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67233120	**Acceptable identification**					**67233120**
Actinomyces naeslundii		90.39	0.42	ESL 12		
Clostridium perfringens		9.61	0.03	MLT 94		
67233121	**Good identification**					**67233121**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67233122	**Good identification**					**67233122**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67233123	**Good identification**					**67233123**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67241000	**Good identification**					**67241000**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium tortuosum		99.42	0.48	MNS 1	99		
Lactobacillus catenaformis		0.33	0.12	CEL 10	99		
Fusobacterium mortiferum		0.09	0.12		1		
67241002	**Good identification**						**67241002**
					SPO		
Eubacterium tortuosum		96.58	0.35	MNS 1	1		
Lactobacillus catenaformis		1.81	0.12	CEL 10	1		
Clostridium barati		1.13	0.04		99		
67241020	**Good identification**						**67241020**
					SPO		
Eubacterium tortuosum		98.39	0.35	MNS 1	1		
Clostridium barati		1.15	0.04	CEL 10	99		
Lactobacillus catenaformis		0.46	0.02		1		
67250000	**Low discrimination**						**67250000**
					GRA	SPO	
Fusobacterium mortiferum		86.98	0.43	NIT 1	1	1	
Clostridium barati		5.43	0.08		99	99	
Lactobacillus catenaformis		5.04	0.12		99	1	
67250100	**Low discrimination**						**67250100**
					GRA	SPO	
Fusobacterium mortiferum		87.55	0.50	NIT 1	1	1	
Clostridium barati		7.03	0.16		99	99	
Lactobacillus catenaformis		3.26	0.15		99	1	
67251000	**Low discrimination**						**67251000**
					GRA		
Lactobacillus catenaformis		55.50	0.47	NIT 1	99		
Fusobacterium mortiferum		14.51	0.46	MLT 90	1		
Lactobacillus jensenii		13.20	0.36		99		
67251002	**Low discrimination**						**67251002**
					SPO		
Lactobacillus catenaformis		53.13	0.47	NIT 1	1		
Clostridium barati		32.95	0.38	MLT 90	99		
Lactobacillus jensenii		8.42	0.33		1		
67251020	**Low discrimination**						**67251020**
					SPO		
Clostridium barati		65.50	0.38	NIT 1	99		
Lactobacillus catenaformis		26.41	0.36		1		
Lactobacillus jensenii		6.28	0.26		1		
67251022	**Low discrimination**						**67251022**
					SPO		
Clostridium barati		85.73	0.46	NIT 1	99		
Lactobacillus catenaformis		11.52	0.36		1		
Lactobacillus jensenii		1.83	0.23		1		
67251100	**Low discrimination**						**67251100**
					SPO		
Lactobacillus catenaformis		40.10	0.50	NIT 1	1		
Collinsella aerofaciens		26.53	0.32	MLT 90	1		
Clostridium barati		16.58	0.38		99		
67251102	**Low discrimination**						**67251102**
					SPO		
Clostridium barati		50.09	0.46	NIT 1	99		
Lactobacillus catenaformis		40.38	0.50		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		4.41	0.30		1		
67251120	**Low discrimination**						**67251120**
					SPO		
Clostridium barati		63.45	0.46	NIT 1	99		
Clostridium paraputrificum		18.95	0.32		99		
Lactobacillus catenaformis		12.79	0.39		1		
67251122	**Low discrimination**						**67251122**
					SPO		
Clostridium barati		87.03	0.54	NIT 1	99		
Lactobacillus catenaformis		5.85	0.39		1		
Clostridium paraputrificum		3.71	0.25		99		
67251162	**Acceptable identification**						**67251162**
					SPO		
Clostridium barati		91.56	0.36	NIT 1	99		
Lactobacillus acidophilus		3.82	0.15	SOR 8	1		
Clostridium paraputrificum		3.38	0.06		99		
67251322	**Acceptable identification**						**67251322**
					GRA	SPO	
Clostridium barati		94.65	0.36	NIT 1	99	99	
Lactobacillus acidophilus		2.40	0.11	RHA 8	99	1	
Prevotella oralis		1.74	0.07		1	1	
67253000	**Good identification**						**67253000**
					SPO		
Clostridium innocuum		99.22	0.56	NIT 1	99		
Clostridium barati		0.35	0.11		99		
Lactobacillus catenaformis		0.20	0.12		1		
67253002	**Good identification**						**67253002**
					SPO		
Clostridium innocuum		98.63	0.56	NIT 1	99		
Clostridium barati		1.04	0.20		99		
Lactobacillus catenaformis		0.20	0.12		1		
67253101	**Good identification**						**67253101**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67253103	**Good identification**						**67253103**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67253121	**Good identification**						**67253121**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67253122	**Acceptable identification**						**67253122**
					SPO		
Clostridium barati		90.89	0.36	NIT 1	99		
Actinomyces naeslundii		3.60	0.39	MAN 8	1		
Lactobacillus acidophilus		1.61	0.08		1		
67253123	**Low discrimination**						**67253123**
					SPO		
Actinomyces naeslundii		85.48	0.42	ESL 12	1		
Clostridium barati		14.52	0.02	MLT 94	99		
67253400	**Good identification**						**67253400**
Clostridium innocuum		100.0	0.43	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
67253402	**Good identification**							**67253402**
Clostridium innocuum		100.0	0.43	NIT 1				
67257100	**Good identification**							**67257100**
					SPO			
Actinomyces israelii		95.67	0.38	MLT 99	1			
Clostridium innocuum		4.33	0.00		99			
67257102	**Acceptable identification**							**67257102**
					SPO			
Actinomyces israelii		93.30	0.41	MLT 99	1			
Clostridium tertium		3.89	0.03		99			
Clostridium innocuum		2.81	0.00		99			
67257500	**Good identification**							**67257500**
Actinomyces israelii		100.0	0.36	MLT 99				
67257502	**Good identification**							**67257502**
Actinomyces israelii		95.41	0.39	MLT 99				
Bifidobacterium adolescentis		4.59	0.11					
67270100	**Good identification**							**67270100**
					GRA	SPO		
Fusobacterium mortiferum		97.72	0.41	NIT 1	1	1		
Clostridium barati		2.28	-.02		99	99		
67271000	**Low discrimination**							**67271000**
					GRA	SPO		
Lactobacillus jensenii		84.49	0.44	NIT 1	99	1		
Fusobacterium mortiferum		9.25	0.38	MLT 95	1	1		
Clostridium innocuum		2.14	0.14		99	99		
67271002	**Low discrimination**							**67271002**
					SPO			
Lactobacillus jensenii		76.61	0.41	NIT 1	1			
Clostridium barati		8.69	0.20	MLT 95	99			
Lactobacillus acidophilus		7.68	0.21		1			
67271010	**Good identification**							**67271010**
Lactobacillus jensenii		100.0	0.36	NIT 1				
67271100	**Low discrimination**							**67271100**
					COC	GRA	SPO	
Fusobacterium mortiferum		47.14	0.44	NIT 1	1	1	1	
Atopobium parvulum		23.24	0.27		99	99	1	
Clostridium barati		13.95	0.20		1	99	99	
67271101	**Good identification**							**67271101**
Actinomyces naeslundii		100.0	0.42	ESL 12				
67271102	**Low discrimination**							**67271102**
					GRA	SPO		
Lactobacillus acidophilus		62.71	0.39	NIT 1	99	1		
Clostridium barati		18.52	0.28	MLT 94	99	99		
Leptotrichia buccalis		9.56	0.15		1	1		
67271103	**Low discrimination**							**67271103**
Actinomyces naeslundii		83.91	0.42	ESL 12				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Lactobacillus acidophilus		16.10	0.05	MLT 94		
67271112	**Good identification**					**67271112**
					GRA	
Leptotrichia buccalis		99.93	0.50	NIT 10	1	
Lactobacillus acidophilus		0.07	0.05	MLT 99	99	
67271121	**Good identification**					**67271121**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67271122	**Low discrimination**					**67271122**
					SPO	
Lactobacillus acidophilus		47.75	0.39	NIT 1	1	
Clostridium barati		42.29	0.36	MLT 94	99	
Clostridium paraputrificum		5.52	0.16		99	
67271123	**Low discrimination**					**67271123**
					SPO	
Actinomyces naeslundii		73.44	0.42	ESL 12	1	
Lactobacillus acidophilus		14.09	0.05	MLT 94	1	
Clostridium barati		12.48	0.02	RAF 92	99	
67273000	**Good identification**					**67273000**
					SPO	
Clostridium innocuum		98.61	0.48	NIT 1	99	
Actinomyces naeslundii		0.72	0.36		1	
Lactobacillus jensenii		0.40	0.10		1	
67273001	**Low discrimination**					**67273001**
					SPO	
Actinomyces naeslundii		52.13	0.39	ESL 12	1	
Clostridium innocuum		47.87	0.14	MLT 94	99	
67273002	**Good identification**					**67273002**
					SPO	
Clostridium innocuum		98.22	0.48	NIT 1	99	
Actinomyces naeslundii		0.72	0.36		1	
Actinomyces israelii		0.40	0.22		1	
67273003	**Low discrimination**					**67273003**
					SPO	
Actinomyces naeslundii		52.13	0.39	ESL 12	1	
Clostridium innocuum		47.87	0.14	MLT 94	99	
67273021	**Good identification**					**67273021**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67273023	**Good identification**					**67273023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67273100	**Low discrimination**					**67273100**
					SPO	
Actinomyces naeslundii		42.52	0.44	ESL 12	1	
Actinomyces israelii		22.29	0.30	MLT 94	1	
Clostridium innocuum		20.58	0.14	RAF 92	99	
67273101	**Good identification**					**67273101**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67273102	**Low discrimination**					**67273102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		24.55	0.44	ESL 12			
Actinomyces israelii		19.31	0.33	MLT 94			
Lactobacillus acidophilus		18.96	0.17	RAF 92			
67273103	**Good identification**						**67273103**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67273120	**Low discrimination**						**67273120**
					SPO		
Actinomyces naeslundii		41.47	0.44	ESL 12	1		
Clostridium tertium		27.10	0.17	MLT 94	99		
Clostridium ramosum		15.81	0.11	RAF 92	92		
67273121	**Good identification**						**67273121**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67273123	**Good identification**						**67273123**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67273400	**Good identification**						**67273400**
					SPO		
Clostridium innocuum		98.75	0.35	NIT 1	99		
Actinomyces israelii		1.25	0.18		1		
67273402	**Good identification**						**67273402**
					SPO		
Clostridium innocuum		98.13	0.35	NIT 1	99		
Actinomyces israelii		1.87	0.21		1		
67275102	**Low discrimination**						**67275102**
					COC		
Actinomyces israelii		63.02	0.38	MLT 99	1		
Atopobium parvulum		31.13	0.19		99		
Lactobacillus acidophilus		5.85	0.05		1		
67275502	**Low discrimination**						**67275502**
					GRA	SPO	
Actinomyces israelii		75.80	0.36	MLT 99	99	1	
Clostridium butyricum		14.06	0.03		99	99	
Mitsuokella multacida		5.68	0.04		1	1	
67277100	**Good identification**						**67277100**
					SPO		
Actinomyces israelii		95.25	0.48	MLT 99	1		
Clostridium tertium		4.75	0.11		99		
67277102	**Acceptable identification**						**67277102**
					SPO		
Actinomyces israelii		90.93	0.51	MLT 99	1		
Clostridium tertium		9.07	0.19		99		
67277140	**Good identification**						**67277140**
Actinomyces israelii		100.0	0.41	MLT 99			
67277142	**Good identification**						**67277142**
Actinomyces israelii		100.0	0.44	MLT 99			
67277400	**Low discrimination**						**67277400**
					SPO		
Actinomyces israelii		73.48	0.36	MLT 99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium innocuum		26.52	0.13		99		
67277402	**Low discrimination**						**67277402**
					SPO		
Actinomyces israelii		80.61	0.39	MLT 99	1		
Clostridium innocuum		19.39	0.13		99		
67277500	**Low discrimination**						**67277500**
					GRA	SPO	
Actinomyces israelii		69.91	0.46	MLT 99	99	1	
Mitsuokella multacida		29.11	0.27		1	1	
Clostridium butyricum		0.98	-.07		99	99	
67277502	**Good identification**						**67277502**
					GRA		
Actinomyces israelii		96.21	0.49	MLT 99	99		
Mitsuokella multacida		2.97	0.11		1		
Bifidobacterium adolescentis		0.82	0.08		99		
67277512	**Low discrimination**						**67277512**
Actinomyces israelii		61.37	0.35	MLT 99			
Bifidobacterium dentium		30.13	0.06	MLZ 13			
Bifidobacterium adolescentis		4.28	0.09				
67277542	**Acceptable identification**						**67277542**
					GRA		
Actinomyces israelii		92.65	0.42	MLT 99	99		
Mitsuokella multacida		7.35	0.11		1		
67301101	**Very good identification**						**67301101**
					SPO		
Clostridium chauvoei		99.87	0.60	SAL 1	99		
Actinomyces naeslundii		0.13	0.33		1		
67301103	**Good identification**						**67301103**
					SPO		
Clostridium chauvoei		98.99	0.47	SAL 1	99		
Actinomyces naeslundii		0.73	0.33		1		
Actinomyces viscosus		0.28	0.02		1		
67301121	**Good identification**						**67301121**
					SPO		
Clostridium chauvoei		99.61	0.51	SAL 1	99		
Actinomyces naeslundii		0.39	0.33		1		
67301123	**Good identification**						**67301123**
					SPO		
Clostridium chauvoei		97.84	0.39	SAL 1	99		
Actinomyces naeslundii		2.16	0.33		1		
67303100	**Good identification**						**67303100**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67303102	**Good identification**						**67303102**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67303103	**Low discrimination**						**67303103**
					SPO		
Actinomyces naeslundii		58.61	0.38	ESL 12	1		
Clostridium chauvoei		41.39	0.13	RAF 92	99		
67303120	**Good identification**						**67303120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	ESL 12		
67303122	**Good identification**					**67303122**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67303123	**Low discrimination**					**67303123**
					SPO	
Actinomyces naeslundii		80.94	0.38	ESL 12	1	
Clostridium chauvoei		19.06	0.04	RAF 92	99	
67310000	**Low discrimination**					**67310000**
					GRA	
Actinomyces odontolyticus		85.20	0.44	ESL 1	99	
Fusobacterium mortiferum		14.27	0.36	MNS 9	1	
Lactobacillus jensenii		0.53	0.02		99	
67310002	**Low discrimination**					**67310002**
					GRA	
Actinomyces odontolyticus		55.31	0.40	ESL 1	99	
Actinomyces viscosus		43.01	0.31	MNS 9	99	
Fusobacterium mortiferum		1.68	0.20		1	
67310100	**Low discrimination**					**67310100**
					GRA	
Fusobacterium mortiferum		50.98	0.42	NIT 1	1	
Actinomyces odontolyticus		41.20	0.35	MLT 11	99	
Collinsella aerofaciens		3.39	-.03		99	
67310102	**Low discrimination**					**67310102**
					GRA	
Actinomyces viscosus		73.55	0.36	ESL 1	99	
Actinomyces odontolyticus		16.80	0.32	RAF 94	99	
Fusobacterium mortiferum		3.77	0.26		1	
67310200	**Good identification**					**67310200**
Actinomyces odontolyticus		100.0	0.49	ESL 1		
67310202	**Good identification**					**67310202**
Actinomyces odontolyticus		100.0	0.45	ESL 1		
67310220	**Good identification**					**67310220**
Actinomyces odontolyticus		100.0	0.35	ESL 1		
67310300	**Good identification**					**67310300**
Actinomyces odontolyticus		100.0	0.40	ESL 1		
67310302	**Good identification**					**67310302**
Actinomyces odontolyticus		97.77	0.36	ESL 1		
Actinomyces viscosus		2.23	0.01	MNS 9		
67311000	**Low discrimination**					**67311000**
					GRA	
Actinomyces odontolyticus		31.62	0.41	ESL 1	99	
Lactobacillus jensenii		29.22	0.36	MNS 9	99	
Fusobacterium mortiferum		11.91	0.39		1	
67311003	**Low discrimination**					**67311003**
					SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		47.90	0.47	ESL 12	1		
Actinomyces viscosus		29.15	0.20	RAF 92	1		
Clostridium chauvoei		22.95	0.19		99		
67311020	**Acceptable identification**						**67311020**
Clostridium perfringens		91.34	0.51	LAC 88			
Lactobacillus jensenii		2.52	0.26				
Actinomyces odontolyticus		1.78	0.27				
67311021	**Low discrimination**						**67311021**
					SPO		
Actinomyces naeslundii		44.29	0.47	ESL 12	1		
Clostridium chauvoei		40.09	0.24	RAF 92	99		
Clostridium perfringens		15.62	0.17		84		
67311022	**Low discrimination**						**67311022**
					SPO		
Clostridium perfringens		76.69	0.46	LAC 88	84		
Clostridium barati		6.94	0.24		99		
Actinomyces naeslundii		3.11	0.44		1		
67311023	**Low discrimination**						**67311023**
					SPO		
Actinomyces naeslundii		75.43	0.47	ESL 12	1		
Clostridium perfringens		12.52	0.12	RAF 92	84		
Clostridium chauvoei		12.05	0.11		99		
67311060	**Good identification**						**67311060**
					SPO		
Clostridium perfringens		97.64	0.39	LAC 88	84		
Actinomyces naeslundii		1.73	0.31		1		
Clostridium barati		0.63	-.03		99		
67311101	**Very good identification**						**67311101**
					SPO		
Clostridium chauvoei		98.85	0.66	SAL 1	99		
Actinomyces naeslundii		1.05	0.55		1		
Collinsella aerofaciens		0.06	-.03		1		
67311102	**Low discrimination**						**67311102**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		45.19	0.40	NIT 1	99	1	
Capnocytophaga ochracea		11.41	0.33	RAF 95	1	90	
Clostridium perfringens		9.32	0.35		99	1	
67311103	**Acceptable identification**						**67311103**
					SPO		
Clostridium chauvoei		91.88	0.53	SAL 1	99		
Actinomyces naeslundii		5.51	0.55		1		
Actinomyces viscosus		2.10	0.24		1		
67311120	**Good identification**						**67311120**
					GRA	CAT	
Clostridium perfringens		91.00	0.66		99	1	
Clostridium paraputrificum		3.20	0.32		99	1	
Capnocytophaga ochracea		1.62	0.33		1	90	
67311121	**Good identification**						**67311121**
					SPO		
Clostridium chauvoei		94.15	0.58	SAL 1	99		
Actinomyces naeslundii		2.99	0.55		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		2.72	0.32		84		
67311122	**Low discrimination**						**67311122**
					GRA	CAT	
Clostridium perfringens		56.07	0.61		99	1	
Capnocytophaga ochracea		19.10	0.50		1	90	
Bifidobacterium longum subsp. infantis		14.95	0.44		99	1	
67311123	**Low discrimination**						**67311123**
					SPO		
Clostridium chauvoei		76.05	0.45	SAL 1	99		
Actinomyces naeslundii		13.69	0.55		1		
Clostridium perfringens		5.85	0.27		84		
67311143	**Low discrimination**						**67311143**
					SPO		
Actinomyces naeslundii		51.17	0.42	ESL 12	1		
Clostridium chauvoei		48.83	0.19	RAF 92	99		
67311160	**Good identification**						**67311160**
					GRA	CAT	
Clostridium perfringens		96.73	0.54		99	1	
Clostridium paraputrificum		1.34	0.13		99	1	
Capnocytophaga ochracea		1.01	0.17		1	90	
67311162	**Low discrimination**						**67311162**
					GRA	CAT	
Clostridium perfringens		79.03	0.48		99	1	
Capnocytophaga ochracea		15.71	0.33		1	90	
Clostridium barati		1.34	0.14		99	1	
67311163	**Low discrimination**						**67311163**
					SPO		
Actinomyces naeslundii		53.44	0.42	ESL 12	1		
Clostridium perfringens		24.66	0.14	RAF 92	84		
Clostridium chauvoei		17.00	0.11		99		
67311200	**Good identification**						**67311200**
Actinomyces odontolyticus		99.52	0.46	ESL 1			
Lactobacillus jensenii		0.48	0.02	MNS 9			
67311202	**Good identification**						**67311202**
					SPO		
Actinomyces odontolyticus		99.14	0.42	ESL 1	1		
Clostridium barati		0.86	-.03	MNS 9	99		
67311300	**Low discrimination**						**67311300**
Actinomyces odontolyticus		51.19	0.37	ESL 1			
Eubacterium contortum		42.35	0.29	MNS 9			
Collinsella aerofaciens		3.25	-.03				
67311522	**Low discrimination**						**67311522**
					GRA	CAT	
Capnocytophaga ochracea		80.81	0.37	SAL 1	1	90	
Clostridium perfringens		13.58	0.27	RAF 90	99	1	
Bifidobacterium longum subsp. infantis		3.62	0.10		99	1	
67312101	**Good identification**						**67312101**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67312103	**Low discrimination**						**67312103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		86.51	0.36	ESL 12		
Propionibacterium propionicum		13.49	0.12	RAF 92		
67312121	**Good identification**					**67312121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67312123	**Good identification**					**67312123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67313000	**Low discrimination**					**67313000**
Actinomyces naeslundii		84.46	0.49	ESL 12		
Actinomyces israelii		10.25	0.24	RAF 92		
Actinomyces odontolyticus		2.75	0.07			
67313001	**Good identification**					**67313001**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67313002	**Low discrimination**					**67313002**
					SPO	
Actinomyces naeslundii		82.28	0.49	ESL 12	1	
Actinomyces israelii		14.98	0.27	RAF 92	1	
Clostridium barati		2.74	-.03		99	
67313003	**Good identification**					**67313003**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67313020	**Low discrimination**					**67313020**
					SPO	
Actinomyces naeslundii		76.58	0.49	ESL 12	1	
Clostridium perfringens		20.87	0.17	RAF 92	84	
Clostridium barati		2.55	-.03		99	
67313021	**Good identification**					**67313021**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67313022	**Low discrimination**					**67313022**
					SPO	
Actinomyces naeslundii		81.43	0.49	ESL 12	1	
Clostridium perfringens		10.44	0.12	RAF 92	84	
Clostridium barati		8.13	0.05		99	
67313023	**Good identification**					**67313023**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67313040	**Low discrimination**					**67313040**
Actinomyces naeslundii		78.90	0.36	ESL 12		
Actinomyces israelii		21.10	0.17	RAF 92		
67313041	**Good identification**					**67313041**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67313042	**Low discrimination**					**67313042**
Actinomyces naeslundii		71.37	0.36	ESL 12		
Actinomyces israelii		28.63	0.20	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67313043	**Good identification**					**67313043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67313060	**Low discrimination**					**67313060**
Actinomyces naeslundii		77.27	0.36	ESL 12		
Clostridium perfringens		22.73	0.05	RAF 92		
67313061	**Good identification**					**67313061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67313062	**Good identification**					**67313062**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67313063	**Good identification**					**67313063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67313100	**Low discrimination**					**67313100**
Actinomyces naeslundii		75.95	0.57	ESL 12		
Actinomyces israelii		12.96	0.35	RAF 92		
Propionibacterium propionicum		4.44	0.26			
67313101	**Low discrimination**					**67313101**
					SPO	
Actinomyces naeslundii		67.05	0.60	ESL 12	1	
Clostridium chauvoei		32.96	0.32	RAF 92	99	
67313102	**Low discrimination**					**67313102**
Actinomyces naeslundii		67.61	0.57	ESL 12		
Actinomyces israelii		17.30	0.38	RAF 92		
Propionibacterium propionicum		5.93	0.29			
67313103	**Good identification**					**67313103**
					SPO	
Actinomyces naeslundii		91.47	0.60	ESL 12	1	
Clostridium chauvoei		7.93	0.19	RAF 92	99	
Propionibacterium propionicum		0.59	0.12		1	
67313110	**Low discrimination**					**67313110**
Actinomyces naeslundii		67.37	0.35	ESL 12		
Actinomyces israelii		32.63	0.21	RAF 92		
67313111	**Good identification**					**67313111**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67313112	**Low discrimination**					**67313112**
					SPO	
Actinomyces naeslundii		39.93	0.35	ESL 12	1	
Actinomyces israelii		29.00	0.24	RAF 92	1	
Clostridium tertium		16.09	0.05	MLZ 5	99	
67313113	**Good identification**					**67313113**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67313120	**Low discrimination**					**67313120**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		55.65	0.57	ESL 12	1		
Clostridium perfringens		39.06	0.32	RAF 92	84		
Clostridium barati		1.95	0.05		99		
67313121	**Low discrimination**						**67313121**
					SPO		
Actinomyces naeslundii		85.92	0.60	ESL 12	1		
Clostridium chauvoei		14.08	0.24	RAF 92	99		
67313122	**Low discrimination**						**67313122**
					GRA	CAT	
Actinomyces naeslundii		57.04	0.57	ESL 12	99	25	
Clostridium perfringens		18.84	0.27	RAF 92	99	1	
Capnocytophaga ochracea		6.42	0.15		1	90	
67313123	**Very good identification**						**67313123**
					SPO		
Actinomyces naeslundii		97.19	0.60	ESL 12	1		
Clostridium chauvoei		2.81	0.11	RAF 92	99		
67313130	**Low discrimination**						**67313130**
					SPO		
Actinomyces naeslundii		71.28	0.35	ESL 12	1		
Clostridium tertium		28.72	0.05	RAF 92	99		
67313131	**Good identification**						**67313131**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67313133	**Good identification**						**67313133**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67313140	**Low discrimination**						**67313140**
Actinomyces naeslundii		58.58	0.44	ESL 12			
Actinomyces israelii		22.02	0.28	RAF 92			
Propionibacterium propionicum		19.41	0.26				
67313141	**Good identification**						**67313141**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67313142	**Low discrimination**						**67313142**
Actinomyces naeslundii		48.52	0.44	ESL 12			
Actinomyces israelii		27.36	0.31	RAF 92			
Propionibacterium propionicum		24.12	0.29				
67313143	**Good identification**						**67313143**
Actinomyces naeslundii		96.45	0.47	ESL 12			
Propionibacterium propionicum		3.55	0.12	RAF 92			
67313160	**Low discrimination**						**67313160**
Actinomyces naeslundii		56.89	0.44	ESL 12			
Clostridium perfringens		43.11	0.20	RAF 92			
67313161	**Good identification**						**67313161**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67313162	**Low discrimination**						**67313162**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		66.78	0.44	ESL 12	99	25	
Clostridium perfringens		23.81	0.14	RAF 92	99	1	
Capnocytophaga ochracea		4.73	-.01		1	90	
67313163	**Good identification**						**67313163**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67313502	**Low discrimination**						**67313502**
Actinomyces israelii		63.90	0.36	CEL 93			
Bifidobacterium adolescentis		33.02	0.25	XYL 92			
Actinomyces naeslundii		3.08	0.23				
67314000	**Good identification**						**67314000**
					GRA		
Actinomyces odontolyticus		97.50	0.38	ESL 1	99		
Fusobacterium mortiferum		2.50	0.17	MNS 9	1		
67314200	**Good identification**						**67314200**
Actinomyces odontolyticus		100.0	0.43	ESL 1			
67314202	**Good identification**						**67314202**
Actinomyces odontolyticus		100.0	0.40	ESL 1			
67315000	**Low discrimination**						**67315000**
					GRA		
Actinomyces odontolyticus		81.18	0.35	ESL 1	99		
Actinomyces israelii		12.61	0.30	MNS 9	99		
Fusobacterium mortiferum		4.68	0.20		1		
67315102	**Low discrimination**						**67315102**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		76.21	0.38	NIT 1	99	1	
Actinomyces israelii		9.86	0.43	RAF 95	99	1	
Capnocytophaga ochracea		5.88	0.23		1	90	
67315122	**Low discrimination**						**67315122**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		66.12	0.43	NIT 1	99	1	
Capnocytophaga ochracea		25.82	0.39	RAF 95	1	90	
Clostridium perfringens		3.06	0.27		99	1	
67315142	**Low discrimination**						**67315142**
					GRA	CAT	
Actinomyces israelii		68.88	0.36	CEL 93	99	1	
Bifidobacterium longum subsp. infantis		13.82	0.04		99	1	
Capnocytophaga ochracea		11.73	0.07		1	90	
67315200	**Low discrimination**						**67315200**
Actinomyces odontolyticus		65.35	0.40	ESL 1			
Eubacterium contortum		34.66	0.29	MNS 9			
67315202	**Low discrimination**						**67315202**
Actinomyces odontolyticus		85.66	0.37	ESL 1			
Eubacterium contortum		13.08	0.16	MNS 9			
Actinomyces israelii		1.26	0.13				
67315300	**Very good identification**						**67315300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium contortum		99.36	0.63	NIT 1		
Actinomyces odontolyticus		0.60	0.32			
Actinomyces israelii		0.05	0.21			
67315302	**Good identification**					**67315302**
Eubacterium contortum		97.20	0.50	NIT 1		
Actinomyces odontolyticus		2.03	0.28			
Bifidobacterium longum subsp. infantis		0.39	0.04			
67315320	**Good identification**					**67315320**
					GRA	
Eubacterium contortum		98.18	0.50	NIT 1	99	
Bacteroides vulgatus		1.27	0.11		1	
Actinomyces odontolyticus		0.55	0.18		99	
67315322	**Low discrimination**					**67315322**
					GRA	
Eubacterium contortum		85.03	0.37	NIT 1	99	
Prevotella oralis		6.51	0.16		1	
Bifidobacterium longum subsp. infantis		3.46	0.08		99	
67315502	**Low discrimination**					**67315502**
Bifidobacterium adolescentis		59.48	0.41	NIT 1		
Actinomyces israelii		28.56	0.41	RAF 92		
Eubacterium contortum		5.57	0.14			
67315512	**Acceptable identification**					**67315512**
Bifidobacterium adolescentis		93.29	0.42	NIT 1		
Actinomyces israelii		5.48	0.27	RAF 92		
Clostridium clostridioforme		1.23	0.07			
67315700	**Very good identification**					**67315700**
					GRA	
Eubacterium contortum		99.92	0.61	NIT 1	99	
Actinomyces israelii		0.05	0.19		99	
Bacteroides vulgatus		0.04	-.05		1	
67315702	**Good identification**					**67315702**
Eubacterium contortum		99.33	0.48	NIT 1		
Actinomyces israelii		0.39	0.22			
Clostridium clostridioforme		0.28	0.10			
67315720	**Low discrimination**					**67315720**
					GRA	
Eubacterium contortum		82.61	0.48	NIT 1	99	
Bacteroides vulgatus		17.39	0.30		1	
67315722	**Good identification**					**67315722**
					GRA	
Eubacterium contortum		95.63	0.36	NIT 1	99	
Parabacteroides distasonis		1.69	0.17		1	
Clostridium clostridioforme		1.52	0.10		50	
67317000	**Good identification**					**67317000**
Actinomyces israelii		100.0	0.43	CEL 93		
67317002	**Good identification**					**67317002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.46	CEL 93		
67317040	**Good identification**					**67317040**
Actinomyces israelii		100.0	0.36	CEL 93		
67317042	**Good identification**					**67317042**
Actinomyces israelii		100.0	0.39	CEL 93		
67317100	**Good identification**					**67317100**
Actinomyces israelii		99.49	0.53	CEL 93		
Actinomyces naeslundii		0.51	0.23			
67317102	**Acceptable identification**					**67317102**
Actinomyces israelii		94.38	0.56	CEL 93		
Bifidobacterium adolescentis		3.47	0.25			
Bifidobacterium longum subsp. infantis		1.30	0.04			
67317110	**Good identification**					**67317110**
					SPO	
Actinomyces israelii		96.78	0.39	CEL 93	1	
Clostridium tertium		3.22	-.01	MLZ 13	99	
67317112	**Low discrimination**					**67317112**
					SPO	
Actinomyces israelii		73.14	0.42	CEL 93	1	
Bifidobacterium adolescentis		21.99	0.27	MLZ 13	1	
Clostridium tertium		4.87	0.07		99	
67317122	**Low discrimination**					**67317122**
					SPO	
Actinomyces israelii		54.05	0.35	CEL 93	1	
Bifidobacterium longum subsp. infantis		20.73	0.08	NAG 6	1	
Clostridium tertium		14.22	0.11		99	
67317140	**Good identification**					**67317140**
Actinomyces israelii		100.0	0.46	CEL 93		
67317142	**Good identification**					**67317142**
Actinomyces israelii		98.85	0.49	CEL 93		
Bifidobacterium adolescentis		1.16	0.09			
67317302	**Low discrimination**					**67317302**
Actinomyces israelii		68.63	0.37	CEL 93		
Eubacterium contortum		31.37	0.16	RHA 7		
67317400	**Good identification**					**67317400**
Actinomyces israelii		100.0	0.41	CEL 93		
67317402	**Good identification**					**67317402**
Actinomyces israelii		100.0	0.44	CEL 93		
67317442	**Good identification**					**67317442**
Actinomyces israelii		100.0	0.37	CEL 93		
67317500	**Good identification**					**67317500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.51	CEL 93		
67317502	**Low discrimination**					**67317502**
Actinomyces israelii		76.88	0.54	CEL 93		
Bifidobacterium adolescentis		23.12	0.39			
67317510	**Good identification**					**67317510**
Actinomyces israelii		100.0	0.37	CEL 93		
67317512	**Low discrimination**					**67317512**
Bifidobacterium adolescentis		71.09	0.41	NIT 1		
Actinomyces israelii		28.91	0.40	RAF 92		
67317540	**Good identification**					**67317540**
Actinomyces israelii		100.0	0.44	CEL 93		
67317542	**Acceptable identification**					**67317542**
Actinomyces israelii		91.28	0.47	CEL 93		
Bifidobacterium adolescentis		8.72	0.24			
67317702	**Low discrimination**					**67317702**
Actinomyces israelii		68.63	0.35	CEL 93		
Eubacterium contortum		31.38	0.14	RHA 7		
67321101	**Good identification**					**67321101**
					SPO	
Clostridium chauvoei		97.78	0.43	SAL 1	99	
Actinomyces naeslundii		2.22	0.38	TRE 10	1	
67321102	**Low discrimination**					**67321102**
					COC	
Actinomyces naeslundii		51.86	0.35	ESL 12	1	
Atopobium parvulum		29.45	0.08	RAF 92	99	
Clostridium perfringens		18.69	0.05	FRU 95	1	
67321120	**Acceptable identification**					**67321120**
					COC	
Clostridium perfringens		93.26	0.37	FRU 99	1	
Actinomyces naeslundii		3.77	0.35		1	
Atopobium parvulum		2.14	0.08		99	
67321121	**Acceptable identification**					**67321121**
					SPO	
Clostridium chauvoei		92.64	0.35	SAL 1	99	
Actinomyces naeslundii		6.31	0.38	TRE 10	1	
Clostridium perfringens		1.05	0.03		84	
67323001	**Good identification**					**67323001**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67323003	**Good identification**					**67323003**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67323021	**Good identification**					**67323021**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67323023	**Good identification**					**67323023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	ESL 12		
67323100	**Good identification**					**67323100**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67323101	**Low discrimination**					**67323101**
					SPO	
Actinomyces naeslundii		81.36	0.43	ESL 12	1	
Clostridium chauvoei		18.64	0.09	RAF 92	99	
67323102	**Acceptable identification**					**67323102**
Actinomyces naeslundii		91.31	0.40	ESL 12		
Actinomyces israelii		8.69	0.13	RAF 92		
67323103	**Good identification**					**67323103**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67323120	**Low discrimination**					**67323120**
Actinomyces naeslundii		88.58	0.40	ESL 12		
Clostridium perfringens		11.42	0.03	RAF 92		
67323121	**Acceptable identification**					**67323121**
					SPO	
Actinomyces naeslundii		92.91	0.43	ESL 12	1	
Clostridium chauvoei		7.09	0.01	RAF 92	99	
67323122	**Good identification**					**67323122**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67323123	**Good identification**					**67323123**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67330103	**Low discrimination**					**67330103**
					SPO	
Actinomyces naeslundii		55.29	0.36	ESL 12	1	
Actinomyces viscosus		31.92	0.08	RAF 92	1	
Clostridium chauvoei		12.79	0.03	GAL 96	99	
67330120	**Acceptable identification**					**67330120**
					COC	
Clostridium perfringens		94.80	0.37	GAL 99	1	
Actinomyces naeslundii		3.03	0.33		1	
Atopobium parvulum		2.17	0.08		99	
67330121	**Low discrimination**					**67330121**
					SPO	
Actinomyces naeslundii		60.69	0.36	ESL 12	1	
Clostridium chauvoei		26.52	0.07	RAF 92	99	
Clostridium perfringens		12.79	0.03	GAL 96	84	
67330123	**Good identification**					**67330123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67331000	**Low discrimination**					**67331000**
Lactobacillus jensenii		71.80	0.44	NIT 1		
Clostridium perfringens		12.46	0.30	CEL 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		8.04	0.49			
67331001	**Low discrimination**					**67331001**
					SPO	
Actinomyces naeslundii		82.27	0.52	ESL 12	1	
Clostridium chauvoei		12.79	0.15	RAF 92	99	
Lactobacillus jensenii		4.95	0.10		1	
67331002	**Low discrimination**					**67331002**
Lactobacillus jensenii		72.94	0.41	NIT 1		
Actinomyces naeslundii		12.25	0.49	CEL 95		
Clostridium perfringens		8.93	0.25			
67331003	**Acceptable identification**					**67331003**
					SPO	
Actinomyces naeslundii		91.65	0.52	ESL 12	1	
Lactobacillus jensenii		3.68	0.07	RAF 92	1	
Clostridium chauvoei		2.51	0.03		99	
67331010	**Good identification**					**67331010**
Lactobacillus jensenii		98.26	0.36	NIT 1		
Actinomyces naeslundii		1.74	0.27	CEL 95		
67331020	**Acceptable identification**					**67331020**
Clostridium perfringens		93.73	0.56	LAC 88		
Lactobacillus jensenii		4.18	0.34			
Actinomyces naeslundii		1.87	0.49			
67331021	**Low discrimination**					**67331021**
					SPO	
Actinomyces naeslundii		71.99	0.52	ESL 12	1	
Clostridium perfringens		24.28	0.22	RAF 92	84	
Clostridium chauvoei		3.73	0.07		99	
67331022	**Low discrimination**					**67331022**
Clostridium perfringens		89.68	0.50	LAC 88		
Lactobacillus jensenii		5.66	0.31			
Actinomyces naeslundii		3.80	0.49			
67331023	**Low discrimination**					**67331023**
Actinomyces naeslundii		86.30	0.52	ESL 12		
Clostridium perfringens		13.70	0.16	RAF 92		
67331041	**Good identification**					**67331041**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67331042	**Low discrimination**					**67331042**
Actinomyces naeslundii		41.77	0.36	ESL 12		
Clostridium perfringens		32.87	0.12	RAF 92		
Lactobacillus jensenii		14.24	0.07			
67331043	**Good identification**					**67331043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67331060	**Good identification**					**67331060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		98.18	0.44	LAC 88			
Actinomyces naeslundii		1.82	0.36				
67331061	**Low discrimination**						**67331061**
Actinomyces naeslundii		73.31	0.39	ESL 12			
Clostridium perfringens		26.69	0.10	RAF 92			
67331062	**Good identification**						**67331062**
Clostridium perfringens		96.22	0.38	LAC 88			
Actinomyces naeslundii		3.78	0.36				
67331063	**Low discrimination**						**67331063**
Actinomyces naeslundii		85.37	0.39	ESL 12			
Clostridium perfringens		14.63	0.04	RAF 92			
67331100	**Low discrimination**						**67331100**
					COC		
Atopobium parvulum		75.06	0.55	NIT 1	99		
Clostridium perfringens		17.87	0.45		1		
Actinomyces naeslundii		4.48	0.57		1		
67331101	**Low discrimination**						**67331101**
					COC	SPO	
Clostridium chauvoei		82.59	0.50	SAL 1	1	99	
Actinomyces naeslundii		15.28	0.60	TRE 10	1	1	
Atopobium parvulum		1.73	0.21		99	1	
67331102	**Low discrimination**						**67331102**
					COC		
Atopobium parvulum		42.26	0.42	NIT 1	99		
Clostridium perfringens		26.83	0.39		1		
Actinomyces naeslundii		14.28	0.57		1		
67331103	**Low discrimination**						**67331103**
					COC	SPO	
Actinomyces naeslundii		49.96	0.60	ESL 12	1	1	
Clostridium chauvoei		47.67	0.37	RAF 92	1	99	
Atopobium parvulum		1.00	0.08		99	1	
67331112	**Good identification**						**67331112**
					COC	GRA	
Leptotrichia buccalis		99.61	0.50	NIT 10	1	1	
Actinomyces naeslundii		0.14	0.35	CEL 99	1	99	
Atopobium parvulum		0.08	0.08		99	99	
67331120	**Very good identification**						**67331120**
					COC		
Clostridium perfringens		96.12	0.71		1		
Atopobium parvulum		2.20	0.42		99		
Actinomyces naeslundii		0.75	0.57		1		
67331121	**Low discrimination**						**67331121**
					SPO		
Clostridium chauvoei		48.82	0.41	SAL 1	99		
Actinomyces naeslundii		27.09	0.60	TRE 10	1		
Clostridium perfringens		23.55	0.37		84		
67331122	**Very good identification**						**67331122**
					COC		
Clostridium perfringens		95.66	0.65		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		1.58	0.57		1		
Atopobium parvulum		0.82	0.29		99		
67331123	**Low discrimination**						**67331123**
					SPO		
Actinomyces naeslundii		57.90	0.60	ESL 12	1		
Clostridium perfringens		23.68	0.31	RAF 92	84		
Clostridium chauvoei		18.42	0.28		99		
67331130	**Acceptable identification**						**67331130**
					GRA		
Clostridium perfringens		91.36	0.37	MLZ 1	99		
Actinomyces naeslundii		3.69	0.35		99		
Leptotrichia buccalis		2.86	-.01		1		
67331131	**Low discrimination**						**67331131**
					SPO		
Actinomyces naeslundii		66.11	0.38	ESL 12	1		
Clostridium chauvoei		22.87	0.07	RAF 92	99		
Clostridium perfringens		11.03	0.03	MLZ 5	84		
67331133	**Low discrimination**						**67331133**
					GRA		
Actinomyces naeslundii		65.92	0.38	ESL 12	99		
Leptotrichia buccalis		34.08	-.01	RAF 92	1		
67331141	**Low discrimination**						**67331141**
					SPO		
Actinomyces naeslundii		76.37	0.47	ESL 12	1		
Clostridium chauvoei		23.63	0.15	RAF 92	99		
67331143	**Acceptable identification**						**67331143**
					SPO		
Actinomyces naeslundii		94.82	0.47	ESL 12	1		
Clostridium chauvoei		5.18	0.03	RAF 92	99		
67331160	**Very good identification**						**67331160**
					SPO		
Clostridium perfringens		98.97	0.59		84		
Actinomyces naeslundii		0.71	0.44		1		
Clostridium paraputrificum		0.20	0.03		99		
67331161	**Low discrimination**						**67331161**
					SPO		
Actinomyces naeslundii		48.99	0.47	ESL 12	1		
Clostridium perfringens		45.96	0.24	RAF 92	84		
Clostridium chauvoei		5.05	0.07		99		
67331162	**Good identification**						**67331162**
Clostridium perfringens		97.83	0.53				
Actinomyces naeslundii		1.49	0.44				
Lactobacillus acidophilus		0.40	0.10				
67331163	**Low discrimination**						**67331163**
Actinomyces naeslundii		69.37	0.47	ESL 12			
Clostridium perfringens		30.63	0.19	RAF 92			
67331320	**Good identification**						**67331320**
					COC	SPO	
Clostridium perfringens		95.98	0.37	RHA 1	1	84	
Atopobium parvulum		2.20	0.08		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		1.08	0.00		1 92	
67331520	**Good identification**					**67331520**
					COC	
Clostridium perfringens		97.02	0.37	ARA 1	1	
Atopobium parvulum		2.22	0.08		99	
Actinomyces naeslundii		0.75	0.23		1	
67332100	**Low discrimination**					**67332100**
Actinomyces naeslundii		62.83	0.38	ESL 12		
Propionibacterium propionicum		37.17	0.24	RAF 92		
67332101	**Good identification**					**67332101**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67332102	**Low discrimination**					**67332102**
Actinomyces naeslundii		49.81	0.38	ESL 12		
Propionibacterium propionicum		44.20	0.27	RAF 92		
Actinomyces israelii		5.99	0.13	GAL 96		
67332103	**Good identification**					**67332103**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67332120	**Low discrimination**					**67332120**
Actinomyces naeslundii		86.00	0.38	ESL 12		
Clostridium perfringens		14.00	0.03	RAF 92		
67332121	**Good identification**					**67332121**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67332122	**Good identification**					**67332122**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67332123	**Good identification**					**67332123**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67333000	**Low discrimination**					**67333000**
Actinomyces naeslundii		78.02	0.54	ESL 12		
Actinomyces israelii		18.35	0.34	RAF 92		
Lactobacillus jensenii		3.63	0.10			
67333001	**Very good identification**					**67333001**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67333002	**Low discrimination**					**67333002**
Actinomyces naeslundii		72.27	0.54	ESL 12		
Actinomyces israelii		25.50	0.37	RAF 92		
Lactobacillus jensenii		2.24	0.07			
67333003	**Very good identification**					**67333003**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67333011	**Good identification**					**67333011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	ESL 12		
67333013	**Good identification**					**67333013**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67333020	**Low discrimination**					**67333020**
Actinomyces naeslundii		78.39	0.54	ESL 12		
Clostridium perfringens		20.43	0.22	RAF 92		
Actinomyces israelii		1.18	0.14			
67333021	**Very good identification**					**67333021**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67333022	**Low discrimination**					**67333022**
Actinomyces naeslundii		87.32	0.54	ESL 12		
Clostridium perfringens		10.71	0.16	RAF 92		
Actinomyces israelii		1.97	0.17			
67333023	**Very good identification**					**67333023**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67333031	**Good identification**					**67333031**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67333033	**Good identification**					**67333033**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67333040	**Low discrimination**					**67333040**
Actinomyces naeslundii		65.86	0.41	ESL 12		
Actinomyces israelii		34.14	0.27	RAF 92		
67333041	**Good identification**					**67333041**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67333042	**Low discrimination**					**67333042**
Actinomyces naeslundii		56.26	0.41	ESL 12		
Actinomyces israelii		43.74	0.30	RAF 92		
67333043	**Good identification**					**67333043**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67333060	**Low discrimination**					**67333060**
Actinomyces naeslundii		78.04	0.41	ESL 12		
Clostridium perfringens		21.96	0.10	RAF 92		
67333061	**Good identification**					**67333061**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67333062	**Low discrimination**					**67333062**
Actinomyces naeslundii		88.31	0.41	ESL 12		
Clostridium perfringens		11.69	0.04	RAF 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67333063	**Good identification**					**67333063**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67333100	**Low discrimination**					**67333100**
					COC	
Actinomyces naeslundii		67.49	0.62	ESL 12	1	
Actinomyces israelii		22.31	0.45	RAF 92	1	
Atopobium parvulum		5.89	0.21		99	
67333101	**Very good identification**					**67333101**
					SPO	
Actinomyces naeslundii		97.26	0.65	ESL 12	1	
Clostridium chauvoei		2.74	0.15	RAF 92	99	
67333102	**Low discrimination**					**67333102**
					SPO	
Actinomyces naeslundii		61.38	0.62	ESL 12	1	
Actinomyces israelii		30.44	0.48	RAF 92	1	
Clostridium tertium		3.39	0.17		99	
67333103	**Very good identification**					**67333103**
					SPO	
Actinomyces naeslundii		99.18	0.65	ESL 12	1	
Clostridium chauvoei		0.49	0.03	RAF 92	99	
Actinomyces israelii		0.33	0.13		1	
67333110	**Low discrimination**					**67333110**
					SPO	
Actinomyces naeslundii		39.05	0.40	ESL 12	1	
Actinomyces israelii		36.65	0.30	RAF 92	1	
Clostridium tertium		24.31	0.13	MLZ 5	99	
67333111	**Good identification**					**67333111**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67333113	**Good identification**					**67333113**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67333120	**Low discrimination**					**67333120**
					SPO	
Actinomyces naeslundii		55.69	0.62	ESL 12	1	
Clostridium perfringens		37.40	0.37	RAF 92	84	
Clostridium tertium		3.08	0.17		99	
67333121	**Very good identification**					**67333121**
					SPO	
Actinomyces naeslundii		98.63	0.65	ESL 12	1	
Clostridium chauvoei		0.93	0.07	RAF 92	99	
Clostridium perfringens		0.45	0.03		84	
67333122	**Low discrimination**					**67333122**
					SPO	
Actinomyces naeslundii		60.50	0.62	ESL 12	1	
Clostridium perfringens		19.12	0.31	RAF 92	84	
Clostridium tertium		10.04	0.25		99	
67333123	**Very good identification**					**67333123**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67333131	**Good identification**					**67333131**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ESL 12		
67333133	**Good identification**					**67333133**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67333140	**Low discrimination**					**67333140**
Actinomyces naeslundii		53.53	0.49	ESL 12		
Actinomyces israelii		39.00	0.38	RAF 92		
Propionibacterium propionicum		7.48	0.24			
67333141	**Good identification**					**67333141**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67333142	**Low discrimination**					**67333142**
Actinomyces israelii		47.46	0.41	CEL 93		
Actinomyces naeslundii		43.44	0.49	XYL 92		
Propionibacterium propionicum		9.10	0.27			
67333143	**Good identification**					**67333143**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67333160	**Low discrimination**					**67333160**
Actinomyces naeslundii		56.45	0.49	ESL 12		
Clostridium perfringens		40.92	0.24	RAF 92		
Actinomyces israelii		2.63	0.17			
67333161	**Good identification**					**67333161**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67333162	**Low discrimination**					**67333162**
Actinomyces naeslundii		70.88	0.49	ESL 12		
Clostridium perfringens		24.18	0.19	RAF 92		
Actinomyces israelii		4.94	0.20			
67333163	**Good identification**					**67333163**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67333402	**Good identification**					**67333402**
Actinomyces israelii		100.0	0.36	CEL 93		
67333500	**Good identification**					**67333500**
Actinomyces israelii		96.40	0.43	CEL 93		
Actinomyces naeslundii		3.60	0.28	XYL 92		
67333502	**Low discrimination**					**67333502**
Actinomyces israelii		89.57	0.46	CEL 93		
Bifidobacterium adolescentis		8.20	0.22	XYL 92		
Actinomyces naeslundii		2.23	0.28			
67333540	**Good identification**					**67333540**
Actinomyces israelii		100.0	0.36	CEL 93		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67333542	**Good identification**						**67333542**
Actinomyces israelii		100.0	0.39	CEL 93			
67335000	**Low discrimination**						**67335000**
					COC		
Actinomyces israelii		47.78	0.39	CEL 93	1		
Atopobium parvulum		46.16	0.25		99		
Lactobacillus jensenii		4.65	0.10		1		
67335002	**Low discrimination**						**67335002**
					COC		
Actinomyces israelii		86.44	0.43	CEL 93	1		
Atopobium parvulum		9.82	0.13		99		
Lactobacillus jensenii		3.74	0.07		1		
67335042	**Good identification**						**67335042**
Actinomyces israelii		100.0	0.35	CEL 93			
67335100	**Very good identification**						**67335100**
					COC		
Atopobium parvulum		95.74	0.60	NIT 1	99		
Actinomyces israelii		4.00	0.50		1		
Clostridium perfringens		0.12	0.11		1		
67335102	**Low discrimination**						**67335102**
					COC		
Atopobium parvulum		72.24	0.47	NIT 1	99		
Actinomyces israelii		25.68	0.53		1		
Bifidobacterium adolescentis		1.16	0.23		1		
67335120	**Low discrimination**						**67335120**
					COC		
Atopobium parvulum		79.73	0.47	NIT 1	99		
Clostridium perfringens		18.92	0.37		1		
Actinomyces israelii		1.21	0.29		1		
67335140	**Low discrimination**						**67335140**
					COC		
Actinomyces israelii		61.69	0.43	CEL 93	1		
Atopobium parvulum		38.31	0.25		99		
67335142	**Acceptable identification**						**67335142**
					COC		
Actinomyces israelii		91.96	0.46	CEL 93	1		
Atopobium parvulum		6.72	0.13		99		
Bifidobacterium adolescentis		1.32	0.08		1		
67335400	**Good identification**						**67335400**
Actinomyces israelii		100.0	0.38	CEL 93			
67335402	**Good identification**						**67335402**
Actinomyces israelii		100.0	0.41	CEL 93			
67335500	**Low discrimination**						**67335500**
					COC	GRA	
Actinomyces israelii		65.49	0.48	CEL 93	1	99	
Atopobium parvulum		19.33	0.25		99	99	
Mitsuokella multacida		8.89	0.21		1	1	
67335502	**Low discrimination**						**67335502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC		
Actinomyces israelii		68.82	0.51	CEL 93	1		
Bifidobacterium adolescentis		25.40	0.38		1		
Atopobium parvulum		2.39	0.13		99		
67335512	**Low discrimination**						**67335512**
Bifidobacterium adolescentis		67.72	0.39	NIT 1			
Actinomyces israelii		22.43	0.37	RAF 92			
Clostridium clostridioforme		9.85	0.22				
67335540	**Low discrimination**						**67335540**
					GRA		
Actinomyces israelii		74.13	0.41	CEL 93	99		
Mitsuokella multacida		25.87	0.21		1		
67335542	**Low discrimination**						**67335542**
					GRA		
Actinomyces israelii		87.48	0.44	CEL 93	99		
Bifidobacterium adolescentis		10.26	0.22		99		
Mitsuokella multacida		2.26	0.04		1		
67335712	**Good identification**						**67335712**
					GRA		
Clostridium clostridioforme		96.50	0.36	NIT 1	50		
Actinomyces israelii		2.26	0.18	RAF 94	99		
Mitsuokella multacida		1.25	0.01		1		
67335732	**Acceptable identification**						**67335732**
Clostridium clostridioforme		92.45	0.36	NIT 1			
Parabacteroides distasonis		7.55	0.23	RAF 94			
67337000	**Good identification**						**67337000**
Actinomyces israelii		100.0	0.52	CEL 93			
67337002	**Good identification**						**67337002**
Actinomyces israelii		100.0	0.55	CEL 93			
67337010	**Good identification**						**67337010**
Actinomyces israelii		100.0	0.38	CEL 93			
67337012	**Good identification**						**67337012**
Actinomyces israelii		100.0	0.41	CEL 93			
67337040	**Good identification**						**67337040**
Actinomyces israelii		100.0	0.45	CEL 93			
67337042	**Good identification**						**67337042**
Actinomyces israelii		100.0	0.48	CEL 93			
67337100	**Very good identification**						**67337100**
					COC	SPO	
Actinomyces israelii		95.06	0.63	CEL 93	1	1	
Atopobium parvulum		4.05	0.25		99	1	
Clostridium tertium		0.64	0.11		1	99	
67337102	**Very good identification**						**67337102**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		97.40	0.66	CEL 93	1		
Clostridium tertium		1.30	0.19		99		
Bifidobacterium adolescentis		0.63	0.22		1		
67337110	**Acceptable identification**						**67337110**
					SPO		
Actinomyces israelii		92.62	0.49	CEL 93	1		
Clostridium tertium		7.38	0.15	MLZ 13	99		
67337112	**Low discrimination**						**67337112**
					SPO		
Actinomyces israelii		82.47	0.52	CEL 93	1		
Clostridium tertium		13.14	0.24	MLZ 13	99		
Bifidobacterium adolescentis		4.39	0.23		1		
67337120	**Low discrimination**						**67337120**
					COC	SPO	
Actinomyces israelii		66.57	0.42	CEL 93	1	1	
Clostridium tertium		20.95	0.19	NAG 6	1	99	
Atopobium parvulum		7.84	0.13		99	1	
67337122	**Low discrimination**						**67337122**
					SPO		
Actinomyces israelii		59.84	0.45	CEL 93	1		
Clostridium tertium		37.67	0.27	NAG 6	99		
Actinomyces naeslundii		1.66	0.28		1		
67337140	**Good identification**						**67337140**
Actinomyces israelii		100.0	0.56	CEL 93			
67337142	**Very good identification**						**67337142**
Actinomyces israelii		100.0	0.59	CEL 93			
67337150	**Good identification**						**67337150**
Actinomyces israelii		100.0	0.42	CEL 93			
67337152	**Good identification**						**67337152**
Actinomyces israelii		98.34	0.45	CEL 93			
Bifidobacterium adolescentis		1.67	0.08	MLZ 13			
67337160	**Good identification**						**67337160**
Actinomyces israelii		100.0	0.35	CEL 93			
67337162	**Good identification**						**67337162**
Actinomyces israelii		100.0	0.38	CEL 93			
67337202	**Good identification**						**67337202**
Actinomyces israelii		100.0	0.36	CEL 93			
67337300	**Good identification**						**67337300**
Actinomyces israelii		100.0	0.43	CEL 93			
67337302	**Good identification**						**67337302**
Actinomyces israelii		100.0	0.46	CEL 93			
67337340	**Good identification**						**67337340**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.36	CEL 93			
67337342	**Good identification**						**67337342**
Actinomyces israelii		100.0	0.39	CEL 93			
67337400	**Good identification**						**67337400**
Actinomyces israelii		100.0	0.51	CEL 93			
67337402	**Good identification**						**67337402**
Actinomyces israelii		100.0	0.54	CEL 93			
67337410	**Good identification**						**67337410**
Actinomyces israelii		100.0	0.37	CEL 93			
67337412	**Good identification**						**67337412**
Actinomyces israelii		100.0	0.40	CEL 93			
67337440	**Good identification**						**67337440**
Actinomyces israelii		100.0	0.44	CEL 93			
67337442	**Good identification**						**67337442**
Actinomyces israelii		100.0	0.47	CEL 93			
67337500	**Good identification**						**67337500**
					GRA	SPO	
Actinomyces israelii		94.54	0.61	CEL 93	99	1	
Mitsuokella multacida		5.28	0.27		1	1	
Clostridium butyricum		0.18	-.07		99	99	
67337502	**Good identification**						**67337502**
					GRA		
Actinomyces israelii		94.57	0.64	CEL 93	99		
Bifidobacterium adolescentis		5.04	0.36		99		
Mitsuokella multacida		0.39	0.11		1		
67337510	**Low discrimination**						**67337510**
					GRA		
Actinomyces israelii		88.92	0.47	CEL 93	99		
Mitsuokella multacida		11.08	0.19	MLZ 13	1		
67337512	**Low discrimination**						**67337512**
Actinomyces israelii		66.18	0.50	CEL 93			
Bifidobacterium adolescentis		28.85	0.38	MLZ 13			
Bifidobacterium dentium		4.36	0.06				
67337520	**Low discrimination**						**67337520**
					GRA		
Actinomyces israelii		82.05	0.41	CEL 93	99		
Mitsuokella multacida		17.95	0.17	NAG 6	1		
67337522	**Good identification**						**67337522**
Actinomyces israelii		100.0	0.44	CEL 93			
67337540	**Low discrimination**						**67337540**
					GRA		
Actinomyces israelii		87.44	0.54	CEL 93	99		
Mitsuokella multacida		12.56	0.27		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67337542	**Very good identification**						**67337542**
					GRA		
Actinomyces israelii		97.32	0.57	CEL 93	99		
Bifidobacterium adolescentis		1.65	0.21		99		
Mitsuokella multacida		1.04	0.11		1		
67337550	**Low discrimination**						**67337550**
					COC	GRA	
Actinomyces israelii		73.07	0.40	CEL 93	1	99	
Mitsuokella multacida		23.42	0.19	MLZ 13	1	1	
Blautia producta		3.51	-.03		99	99	
67337552	**Low discrimination**						**67337552**
					GRA		
Actinomyces israelii		86.04	0.43	CEL 93	99		
Bifidobacterium adolescentis		11.92	0.22	MLZ 13	99		
Mitsuokella multacida		2.04	0.03		1		
67337562	**Good identification**						**67337562**
Actinomyces israelii		100.0	0.37	CEL 93			
67337702	**Acceptable identification**						**67337702**
					GRA		
Actinomyces israelii		90.74	0.45	CEL 93	99		
Mitsuokella multacida		9.27	0.15	RHA 7	1		
67337742	**Low discrimination**						**67337742**
					GRA		
Actinomyces israelii		79.20	0.38	CEL 93	99		
Mitsuokella multacida		20.80	0.15	RHA 7	1		
67341123	**Low discrimination**						**67341123**
					SPO		
Actinomyces naeslundii		56.57	0.35	ESL 12	1		
Clostridium chauvoei		19.50	0.04	RAF 92	99		
Clostridium barati		12.83	-.03	FRU 95	99		
67343100	**Low discrimination**						**67343100**
Actinomyces naeslundii		47.00	0.37	ESL 12			
Collinsella aerofaciens		29.70	-.03	RAF 92			
Actinomyces israelii		15.42	0.20	FRU 95			
67343101	**Good identification**						**67343101**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67343102	**Low discrimination**						**67343102**
					SPO		
Actinomyces naeslundii		46.16	0.37	ESL 12	1		
Clostridium barati		23.22	0.05	RAF 92	99		
Actinomyces israelii		22.72	0.23	FRU 95	1		
67343103	**Good identification**						**67343103**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67343120	**Low discrimination**						**67343120**
					SPO		
Actinomyces naeslundii		49.32	0.37	ESL 12	1		
Clostridium barati		24.81	0.05	RAF 92	99		
Clostridium paraputrificum		17.45	-.03	FRU 95	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67343121	**Good identification**					**67343121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67343123	**Good identification**					**67343123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67347100	**Good identification**					**67347100**
Actinomyces israelii		100.0	0.38	FRU 99		
67347102	**Good identification**					**67347102**
					SPO	
Actinomyces israelii		96.00	0.41	FRU 99	1	
Clostridium tertium		4.00	0.03		99	
67347500	**Good identification**					**67347500**
Actinomyces israelii		100.0	0.36	FRU 99		
67347502	**Good identification**					**67347502**
Actinomyces israelii		95.41	0.39	FRU 99		
Bifidobacterium adolescentis		4.59	0.11			
67350022	**Acceptable identification**					**67350022**
					SPO	
Clostridium barati		92.77	0.36	NIT 1	99	
Lactobacillus catenaformis		4.41	0.18	GAL 95	1	
Lactobacillus jensenii		1.48	0.10		1	
67350102	**Low discrimination**					**67350102**
					SPO	
Clostridium barati		56.12	0.36	NIT 1	99	
Lactobacillus catenaformis		16.01	0.32	GAL 95	1	
Collinsella aerofaciens		12.94	0.15		1	
67350122	**Low discrimination**					**67350122**
					SPO	
Clostridium barati		80.39	0.44	NIT 1	99	
Clostridium paraputrificum		13.35	0.25	GAL 95	99	
Lactobacillus catenaformis		1.91	0.21		1	
67351000	**Low discrimination**					**67351000**
					SPO	
Lactobacillus catenaformis		57.53	0.63	NIT 1	1	
Lactobacillus jensenii		28.89	0.58		1	
Clostridium barati		6.45	0.41		99	
67351002	**Low discrimination**					**67351002**
					SPO	
Lactobacillus catenaformis		58.05	0.63	NIT 1	1	
Clostridium barati		19.53	0.50		99	
Lactobacillus jensenii		19.43	0.55		1	
67351010	**Acceptable identification**					**67351010**
					SPO	
Lactobacillus jensenii		92.77	0.50	NIT 1	1	
Lactobacillus catenaformis		5.60	0.29		1	
Clostridium barati		0.63	0.07		99	
67351012	**Low discrimination**					**67351012**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Lactobacillus jensenii		82.35	0.47	NIT 1	99	
Lactobacillus catenaformis		7.46	0.29		99	
Leptotrichia buccalis		6.85	0.15		1	
67351020	**Low discrimination**					**67351020**
					SPO	
Clostridium barati		41.90	0.50	NIT 1	99	
Lactobacillus catenaformis		31.13	0.53		1	
Lactobacillus jensenii		15.63	0.47		1	
67351021	**Low discrimination**					**67351021**
					SPO	
Actinomyces naeslundii		53.78	0.49	ESL 12	1	
Clostridium barati		20.10	0.15	RAF 92	99	
Lactobacillus catenaformis		14.94	0.18		1	
67351022	**Low discrimination**					**67351022**
					SPO	
Clostridium barati		70.97	0.58	NIT 1	99	
Lactobacillus catenaformis		17.58	0.53		1	
Lactobacillus jensenii		5.88	0.44		1	
67351030	**Low discrimination**					**67351030**
					SPO	
Lactobacillus jensenii		85.90	0.39	NIT 1	1	
Clostridium barati		6.98	0.15		99	
Lactobacillus catenaformis		5.19	0.18		1	
67351032	**Low discrimination**					**67351032**
					SPO	
Lactobacillus jensenii		65.18	0.36	NIT 1	1	
Clostridium barati		23.82	0.24		99	
Lactobacillus catenaformis		5.90	0.18		1	
67351041	**Good identification**					**67351041**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67351043	**Low discrimination**					**67351043**
					SPO	
Actinomyces naeslundii		84.44	0.37	ESL 12	1	
Clostridium barati		15.56	-.03	RAF 92	99	
67351061	**Low discrimination**					**67351061**
					SPO	
Actinomyces naeslundii		84.44	0.37	ESL 12	1	
Clostridium barati		15.56	-.03	RAF 92	99	
67351062	**Low discrimination**					**67351062**
					SPO	
Clostridium barati		89.13	0.40	NIT 1	99	
Lactobacillus acidophilus		3.11	0.17	SOR 8	1	
Lactobacillus catenaformis		2.57	0.18		1	
67351063	**Low discrimination**					**67351063**
					SPO	
Actinomyces naeslundii		64.41	0.37	ESL 12	1	
Clostridium barati		35.59	0.05	RAF 92	99	
67351100	**Low discrimination**					**67351100**
					SPO	
Collinsella aerofaciens		84.80	0.66	NIT 1	1	
Lactobacillus catenaformis		11.65	0.66		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		2.61	0.50		99	
67351102	**Low discrimination**					**67351102**
					SPO	
Lactobacillus catenaformis		36.34	0.66	NIT 1	1	
Collinsella aerofaciens		29.38	0.50		1	
Clostridium barati		24.45	0.58		99	
67351103	**Low discrimination**					**67351103**
Actinomyces naeslundii		36.00	0.57	ESL 12		
Lactobacillus catenaformis		21.08	0.32	RAF 92		
Collinsella aerofaciens		17.04	0.15			
67351111	**Low discrimination**					**67351111**
Actinomyces naeslundii		55.01	0.35	ESL 12		
Collinsella aerofaciens		44.99	-.03	RAF 92		
67351112	**Acceptable identification**					**67351112**
					GRA	
Leptotrichia buccalis		93.38	0.50	NIT 10	1	
Bifidobacterium breve		1.77	0.40	TRE 99	99	
Lactobacillus catenaformis		1.54	0.32		99	
67351120	**Low discrimination**					**67351120**
					SPO	
Clostridium paraputrificum		69.10	0.66	NIT 1	99	
Collinsella aerofaciens		13.71	0.50		1	
Clostridium barati		11.41	0.58		99	
67351122	**Low discrimination**					**67351122**
					SPO	
Clostridium barati		44.29	0.66	NIT 1	99	
Clostridium paraputrificum		38.31	0.60		99	
Lactobacillus catenaformis		5.49	0.56		1	
67351131	**Low discrimination**					**67351131**
					SPO	
Actinomyces naeslundii		68.60	0.35	ESL 12	1	
Clostridium paraputrificum		31.40	-.03	RAF 92	99	
67351133	**Low discrimination**					**67351133**
					GRA	
Actinomyces naeslundii		41.50	0.35	ESL 12	99	
Leptotrichia buccalis		31.42	-.01	RAF 92	1	
Bifidobacterium breve		9.52	0.10	MLZ 5	99	
67351141	**Low discrimination**					**67351141**
					SPO	
Actinomyces naeslundii		76.42	0.44	ESL 12	1	
Collinsella aerofaciens		18.64	-.03	RAF 92	1	
Clostridium barati		4.95	-.03		99	
67351142	**Low discrimination**					**67351142**
					SPO	
Clostridium barati		48.87	0.40	NIT 1	99	
Lactobacillus acidophilus		19.61	0.35	SOR 8	1	
Bifidobacterium breve		9.69	0.40		1	
67351143	**Low discrimination**					**67351143**
					SPO	
Actinomyces naeslundii		78.61	0.44	ESL 12	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium barati		15.26	0.05	RAF 92	99		
Lactobacillus acidophilus		6.12	0.01		1		
67351152	**Low discrimination**						**67351152**
					GRA	SPO	
Bifidobacterium breve		60.42	0.40	NIT 1	99	1	
Leptotrichia buccalis		32.21	0.15	RAF 95	1	1	
Clostridium barati		3.08	0.05		99	99	
67351160	**Low discrimination**						**67351160**
Clostridium paraputrificum		78.59	0.47	NIT 1			
Clostridium barati		14.99	0.40	SOR 7			
Clostridium perfringens		2.38	0.30				
67351162	**Low discrimination**						**67351162**
					SPO		
Clostridium barati		47.71	0.48	NIT 1	99		
Clostridium paraputrificum		35.72	0.40	SOR 8	99		
Lactobacillus acidophilus		6.38	0.35		1		
67351163	**Low discrimination**						**67351163**
					SPO		
Actinomyces naeslundii		46.19	0.44	ESL 12	1		
Clostridium barati		26.90	0.14	RAF 92	99		
Clostridium paraputrificum		20.14	0.06		99		
67351172	**Low discrimination**						**67351172**
					SPO		
Bifidobacterium breve		81.76	0.44	NIT 1	1		
Clostridium barati		7.03	0.14	RAF 95	99		
Clostridium paraputrificum		5.26	0.06		99		
67351222	**Low discrimination**						**67351222**
					GRA	SPO	
Clostridium barati		88.18	0.40	NIT 1	99	99	
Prevotella oralis		6.29	0.21	RHA 8	1	1	
Lactobacillus catenaformis		2.54	0.18		99	1	
67351302	**Low discrimination**						**67351302**
					SPO		
Clostridium barati		61.30	0.40	NIT 1	99		
Lactobacillus acidophilus		14.98	0.31	RHA 8	1		
Lactobacillus catenaformis		10.58	0.32		1		
67351320	**Low discrimination**						**67351320**
					SPO		
Clostridium barati		50.51	0.40	NIT 1	99		
Clostridium paraputrificum		35.52	0.32	RHA 8	99		
Collinsella aerofaciens		7.05	0.15		1		
67351322	**Low discrimination**						**67351322**
					GRA	SPO	
Clostridium barati		61.93	0.48	NIT 1	99	99	
Prevotella oralis		23.07	0.41	RHA 8	1	1	
Clostridium paraputrificum		6.22	0.25		99	99	
67351542	**Low discrimination**						**67351542**
					SPO		
Actinomyces israelii		69.47	0.36	XYL 92	1		
Clostridium barati		12.77	0.05		99		
Bifidobacterium adolescentis		12.64	0.17		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67352101	**Good identification**					**67352101**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67352103	**Good identification**					**67352103**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67352121	**Good identification**					**67352121**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67352123	**Good identification**					**67352123**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67353000	**Low discrimination**					**67353000**
Actinomyces israelii		24.75	0.44	XYL 92		
Actinomyces naeslundii		20.35	0.51			
Lactobacillus catenaformis		17.47	0.29			
67353001	**Good identification**					**67353001**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67353003	**Good identification**					**67353003**
					SPO	
Actinomyces naeslundii		98.35	0.54	ESL 12	1	
Clostridium barati		1.65	-.03	RAF 92	99	
67353021	**Good identification**					**67353021**
					SPO	
Actinomyces naeslundii		98.35	0.54	ESL 12	1	
Clostridium barati		1.65	-.03	RAF 92	99	
67353022	**Low discrimination**					**67353022**
					SPO	
Clostridium barati		81.67	0.40	NIT 1	99	
Actinomyces naeslundii		10.95	0.51	MAN 8	1	
Lactobacillus catenaformis		2.35	0.18		1	
67353023	**Good identification**					**67353023**
					SPO	
Actinomyces naeslundii		95.22	0.54	ESL 12	1	
Clostridium barati		4.78	0.05	RAF 92	99	
67353040	**Low discrimination**					**67353040**
					SPO	
Actinomyces israelii		65.56	0.37	XYL 92	1	
Actinomyces naeslundii		24.46	0.38		1	
Clostridium barati		9.99	0.05		99	
67353041	**Good identification**					**67353041**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67353042	**Low discrimination**					**67353042**
					SPO	
Actinomyces israelii		64.38	0.40	XYL 92	1	
Clostridium barati		19.62	0.13		99	
Actinomyces naeslundii		16.01	0.38		1	
67353043	**Good identification**					**67353043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
67353061	**Good identification**					**67353061**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67353063	**Good identification**					**67353063**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67353101	**Very good identification**					**67353101**
Actinomyces naeslundii		96.19	0.62	ESL 12		
Collinsella aerofaciens		2.13	-.03	RAF 92		
Actinomyces israelii		1.11	0.20			
67353102	**Low discrimination**					**67353102**
					SPO	
Clostridium barati		29.42	0.40	NIT 1	99	
Actinomyces israelii		28.79	0.57	MAN 8	1	
Actinomyces naeslundii		11.23	0.59		1	
67353103	**Very good identification**					**67353103**
					SPO	
Actinomyces naeslundii		96.07	0.62	ESL 12	1	
Clostridium barati		1.70	0.05	RAF 92	99	
Actinomyces israelii		1.66	0.23		1	
67353111	**Good identification**					**67353111**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67353112	**Low discrimination**					**67353112**
					SPO	
Clostridium tertium		55.09	0.39	NIT 1	99	
Bifidobacterium breve		18.09	0.40	TRE 90	1	
Actinomyces israelii		13.33	0.43		1	
67353113	**Low discrimination**					**67353113**
					SPO	
Actinomyces naeslundii		83.15	0.40	ESL 12	1	
Clostridium tertium		16.85	0.05	RAF 92	99	
67353120	**Low discrimination**					**67353120**
					SPO	
Clostridium barati		35.63	0.40	NIT 1	99	
Clostridium paraputrificum		25.06	0.32	MAN 8	99	
Actinomyces naeslundii		13.60	0.59		1	
67353121	**Very good identification**					**67353121**
					SPO	
Actinomyces naeslundii		96.52	0.62	ESL 12	1	
Clostridium barati		1.70	0.05	RAF 92	99	
Clostridium paraputrificum		1.20	-.03		99	
67353122	**Low discrimination**					**67353122**
Clostridium barati		51.36	0.48	NIT 1		
Clostridium tertium		17.46	0.43	MAN 8		
Clostridium ramosum		7.80	0.35			
67353123	**Good identification**					**67353123**
					SPO	
Actinomyces naeslundii		91.70	0.62	ESL 12	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium barati		4.86	0.14	RAF 92	99	
Clostridium tertium		1.65	0.09		99	
67353130	**Low discrimination**					**67353130**
					SPO	
Clostridium tertium		88.52	0.39	NIT 1	99	
Bifidobacterium breve		5.11	0.27	TRE 90	1	
Actinomyces naeslundii		2.94	0.37		1	
67353131	**Low discrimination**					**67353131**
					SPO	
Actinomyces naeslundii		83.15	0.40	ESL 12	1	
Clostridium tertium		16.85	0.05	RAF 92	99	
67353132	**Low discrimination**					**67353132**
					SPO	
Clostridium tertium		80.76	0.47	NIT 1	99	
Bifidobacterium breve		15.71	0.44	TRE 90	1	
Clostridium barati		1.35	0.14		99	
67353133	**Low discrimination**					**67353133**
					SPO	
Actinomyces naeslundii		57.94	0.40	ESL 12	1	
Clostridium tertium		35.21	0.13	RAF 92	99	
Bifidobacterium breve		6.85	0.10	MLZ 5	1	
67353140	**Low discrimination**					**67353140**
					SPO	
Actinomyces israelii		66.36	0.47	XYL 92	1	
Actinomyces naeslundii		17.61	0.46		1	
Clostridium barati		7.58	0.13		99	
67353141	**Good identification**					**67353141**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67353142	**Low discrimination**					**67353142**
					SPO	
Actinomyces israelii		50.22	0.50	XYL 92	1	
Bifidobacterium breve		26.18	0.40		1	
Clostridium barati		11.48	0.21		99	
67353143	**Good identification**					**67353143**
Actinomyces naeslundii		96.34	0.49	ESL 12		
Actinomyces israelii		3.67	0.16	RAF 92		
67353152	**Low discrimination**					**67353152**
					SPO	
Bifidobacterium breve		74.89	0.40	NIT 1	1	
Actinomyces israelii		21.47	0.36	RAF 95	1	
Clostridium tertium		2.30	0.05		99	
67353161	**Good identification**					**67353161**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67353162	**Low discrimination**					**67353162**
					SPO	
Bifidobacterium breve		46.17	0.44	NIT 1	1	
Clostridium barati		34.15	0.30	RAF 95	99	
Actinomyces naeslundii		8.82	0.46		1	
67353163	**Good identification**					**67353163**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.59	0.49	ESL 12		
Bifidobacterium breve		3.41	0.10	RAF 92		
67353172	**Acceptable identification**					**67353172**
					SPO	
Bifidobacterium breve		93.26	0.44	NIT 1	1	
Clostridium tertium		4.84	0.13	RAF 95	99	
Actinomyces israelii		0.96	0.15		1	
67353322	**Low discrimination**					**67353322**
					SPO	
Clostridium ramosum		66.05	0.37	NIT 1	92	
Clostridium barati		28.52	0.30	TRE 94	99	
Lactobacillus acidophilus		1.41	0.09		1	
67353400	**Acceptable identification**					**67353400**
					SPO	
Actinomyces israelii		92.07	0.42	XYL 92	1	
Clostridium innocuum		7.93	0.09		99	
67353402	**Acceptable identification**					**67353402**
					SPO	
Actinomyces israelii		93.09	0.45	XYL 92	1	
Clostridium innocuum		5.35	0.09		99	
Clostridium barati		1.57	-.03		99	
67353440	**Good identification**					**67353440**
Actinomyces israelii		100.0	0.35	XYL 92		
67353442	**Good identification**					**67353442**
Actinomyces israelii		100.0	0.38	XYL 92		
67353500	**Good identification**					**67353500**
Actinomyces israelii		96.40	0.52	XYL 92		
Collinsella aerofaciens		2.29	-.03			
Actinomyces naeslundii		0.70	0.25			
67353502	**Acceptable identification**					**67353502**
					SPO	
Actinomyces israelii		90.55	0.55	XYL 92	1	
Bifidobacterium adolescentis		7.48	0.31		1	
Clostridium barati		1.14	0.05		99	
67353510	**Good identification**					**67353510**
Actinomyces israelii		100.0	0.38	XYL 92		
67353512	**Low discrimination**					**67353512**
					SPO	
Actinomyces israelii		57.91	0.41	XYL 92	1	
Bifidobacterium adolescentis		39.14	0.32	MLZ 13	1	
Clostridium tertium		2.95	0.05		99	
67353540	**Good identification**					**67353540**
Actinomyces israelii		100.0	0.45	XYL 92		
67353542	**Good identification**					**67353542**
Actinomyces israelii		97.44	0.48	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		2.56	0.15			
67353702	**Good identification**					**67353702**
Actinomyces israelii		100.0	0.36	XYL 92		
67355000	**Low discrimination**					**67355000**
Actinomyces israelii		62.15	0.49			
Lactobacillus catenaformis		21.61	0.29			
Lactobacillus jensenii		10.85	0.24			
67355002	**Low discrimination**					**67355002**
					SPO	
Actinomyces israelii		71.78	0.52		1	
Lactobacillus catenaformis		16.64	0.29		1	
Clostridium barati		5.60	0.15		99	
67355012	**Low discrimination**					**67355012**
Actinomyces israelii		85.24	0.38	MLZ 13		
Lactobacillus jensenii		14.76	0.12			
67355040	**Good identification**					**67355040**
Actinomyces israelii		100.0	0.42			
67355042	**Good identification**					**67355042**
					SPO	
Actinomyces israelii		98.29	0.45		1	
Clostridium barati		1.72	-.03		99	
67355100	**Low discrimination**					**67355100**
					COC	
Actinomyces israelii		43.58	0.59		1	
Collinsella aerofaciens		41.36	0.32		1	
Atopobium parvulum		6.95	0.32		99	
67355102	**Low discrimination**					**67355102**
Actinomyces israelii		75.45	0.62			
Lactobacillus catenaformis		6.56	0.32			
Collinsella aerofaciens		5.30	0.15			
67355110	**Acceptable identification**					**67355110**
					SPO	
Actinomyces israelii		92.73	0.45	MLZ 13	1	
Collinsella aerofaciens		5.95	-.03		1	
Clostridium tertium		1.32	-.01		99	
67355112	**Low discrimination**					**67355112**
					GRA	
Actinomyces israelii		58.21	0.48	MLZ 13	99	
Leptotrichia buccalis		20.75	0.15		1	
Bifidobacterium adolescentis		19.39	0.34		99	
67355140	**Good identification**					**67355140**
					SPO	
Actinomyces israelii		96.98	0.52		1	
Collinsella aerofaciens		2.39	-.03		1	
Clostridium barati		0.63	-.03		99	
67355142	**Good identification**					**67355142**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		96.97	0.55		1		
Clostridium barati		1.27	0.05		99		
Bifidobacterium adolescentis		1.26	0.17		1		
67355150	**Good identification**						**67355150**
Actinomyces israelii		100.0	0.38	MLZ 13			
67355152	**Acceptable identification**						**67355152**
Actinomyces israelii		90.43	0.41	MLZ 13			
Bifidobacterium adolescentis		9.57	0.18				
67355302	**Low discrimination**						**67355302**
					SPO		
Actinomyces israelii		76.27	0.43	RHA 7	1		
Eubacterium contortum		14.87	0.16		1		
Clostridium barati		5.16	0.05		99		
67355342	**Good identification**						**67355342**
Actinomyces israelii		100.0	0.36	RHA 7			
67355400	**Good identification**						**67355400**
					SPO		
Actinomyces israelii		97.22	0.47		1		
Clostridium butyricum		2.78	0.00		99		
67355402	**Good identification**						**67355402**
					SPO		
Actinomyces israelii		97.95	0.50		1		
Bifidobacterium adolescentis		1.32	0.12		1		
Clostridium butyricum		0.73	-.07		99		
67355412	**Low discrimination**						**67355412**
Actinomyces israelii		84.90	0.36	MLZ 13			
Bifidobacterium adolescentis		9.35	0.14				
Clostridium clostridioforme		5.76	0.07				
67355440	**Good identification**						**67355440**
Actinomyces israelii		100.0	0.40				
67355442	**Good identification**						**67355442**
Actinomyces israelii		100.0	0.43				
67355500	**Low discrimination**						**67355500**
					GRA	SPO	
Actinomyces israelii		81.50	0.58		99	1	
Clostridium butyricum		14.00	0.24		99	99	
Mitsuokella multacida		3.13	0.21		1	1	
67355502	**Low discrimination**						**67355502**
					SPO		
Actinomyces israelii		72.22	0.61		1		
Bifidobacterium adolescentis		24.05	0.46		1		
Clostridium butyricum		3.22	0.17		99		
67355510	**Low discrimination**						**67355510**
					GRA	SPO	
Actinomyces israelii		84.21	0.44	MLZ 13	99	1	
Mitsuokella multacida		7.22	0.13		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		5.09	0.02		99	99	
67355512	**Low discrimination**						**67355512**
Bifidobacterium adolescentis		71.75	0.48	NIT 1			
Actinomyces israelii		26.34	0.47	RAF 92			
Clostridium clostridioforme		1.50	0.16				
67355520	**Low discrimination**						**67355520**
					GRA	SPO	
Actinomyces israelii		63.05	0.37	NAG 6	99	1	
Clostridium butyricum		14.75	0.05		99	99	
Mitsuokella multacida		9.49	0.11		1	1	
67355522	**Low discrimination**						**67355522**
					GRA	CAT	
Actinomyces israelii		53.26	0.40	NAG 6	99	1	
Capnocytophaga ochracea		18.17	0.16		1	90	
Bacteroides uniformis		15.95	0.09		1	5	
67355540	**Low discrimination**						**67355540**
					GRA	SPO	
Actinomyces israelii		87.94	0.51		99	1	
Mitsuokella multacida		8.69	0.21		1	1	
Clostridium butyricum		3.38	0.05		99	99	
67355542	**Low discrimination**						**67355542**
					SPO		
Actinomyces israelii		89.04	0.54		1		
Bifidobacterium adolescentis		9.43	0.31		1		
Clostridium butyricum		0.89	-.02		99		
67355550	**Low discrimination**						**67355550**
					GRA		
Actinomyces israelii		81.94	0.37	MLZ 13	99		
Mitsuokella multacida		18.06	0.13		1		
67355552	**Low discrimination**						**67355552**
Actinomyces israelii		53.60	0.40	MLZ 13			
Bifidobacterium adolescentis		46.41	0.32				
67355702	**Low discrimination**						**67355702**
Actinomyces israelii		67.06	0.41	RHA 7			
Eubacterium contortum		13.08	0.14				
Clostridium clostridioforme		12.16	0.19				
67356100	**Good identification**						**67356100**
Actinomyces israelii		100.0	0.38	GAL 99			
67356102	**Good identification**						**67356102**
					SPO		
Actinomyces israelii		96.00	0.41	GAL 99	1		
Clostridium tertium		4.00	0.03		99		
67356500	**Good identification**						**67356500**
Actinomyces israelii		100.0	0.36	GAL 99			
67356502	**Good identification**						**67356502**
Actinomyces israelii		95.41	0.39	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium adolescentis		4.59	0.11			
67357000	**Very good identification**					**67357000**
					SPO	
Actinomyces israelii		99.82	0.62		1	
Clostridium innocuum		0.18	0.00		99	
67357002	**Very good identification**					**67357002**
					SPO	
Actinomyces israelii		99.59	0.65		1	
Clostridium tertium		0.17	0.03		99	
Clostridium innocuum		0.12	0.00		99	
67357010	**Good identification**					**67357010**
					SPO	
Actinomyces israelii		99.01	0.48	MLZ 13	1	
Clostridium tertium		0.99	-.01		99	
67357012	**Good identification**					**67357012**
					SPO	
Actinomyces israelii		98.03	0.51	MLZ 13	1	
Clostridium tertium		1.97	0.07		99	
67357020	**Acceptable identification**					**67357020**
					SPO	
Actinomyces israelii		93.66	0.41	NAG 6	1	
Clostridium tertium		3.71	0.03		99	
Clostridium barati		2.64	-.03		99	
67357022	**Low discrimination**					**67357022**
					SPO	
Actinomyces israelii		88.07	0.44	NAG 6	1	
Clostridium tertium		6.98	0.11		99	
Clostridium barati		4.96	0.05		99	
67357040	**Good identification**					**67357040**
Actinomyces israelii		100.0	0.55			
67357042	**Very good identification**					**67357042**
Actinomyces israelii		100.0	0.58			
67357050	**Good identification**					**67357050**
Actinomyces israelii		100.0	0.41	MLZ 13		
67357052	**Good identification**					**67357052**
Actinomyces israelii		100.0	0.44	MLZ 13		
67357062	**Good identification**					**67357062**
Actinomyces israelii		100.0	0.37	NAG 6		
67357100	**Very good identification**					**67357100**
					SPO	
Actinomyces israelii		97.70	0.72		1	
Clostridium tertium		2.04	0.29		99	
Collinsella aerofaciens		0.17	-.03		1	
67357101	**Acceptable identification**					**67357101**
Actinomyces israelii		92.91	0.38	URE 1		
Actinomyces naeslundii		7.09	0.28			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67357102	**Very good identification**					**67357102**
					SPO	
Actinomyces israelii		95.34	0.75		1	
Clostridium tertium		3.98	0.37		99	
Bifidobacterium adolescentis		0.56	0.31		1	
67357103	**Acceptable identification**					**67357103**
					SPO	
Actinomyces israelii		91.53	0.41	URE 1	1	
Actinomyces naeslundii		4.66	0.28		1	
Clostridium tertium		3.82	0.03		99	
67357110	**Low discrimination**					**67357110**
					SPO	
Actinomyces israelii		80.12	0.58	MLZ 13	1	
Clostridium tertium		19.88	0.33		99	
67357112	**Low discrimination**					**67357112**
					SPO	
Actinomyces israelii		64.75	0.61	MLZ 13	1	
Clostridium tertium		32.13	0.41		99	
Bifidobacterium adolescentis		3.11	0.32		1	
67357120	**Low discrimination**					**67357120**
					SPO	
Actinomyces israelii		49.40	0.52	NAG 6	1	
Clostridium tertium		48.43	0.37		99	
Clostridium barati		1.04	0.05		99	
67357122	**Low discrimination**					**67357122**
					SPO	
Clostridium tertium		64.74	0.45	NIT 1	99	
Actinomyces israelii		33.02	0.55	TRE 90	1	
Clostridium barati		1.39	0.14		99	
67357130	**Acceptable identification**					**67357130**
					SPO	
Clostridium tertium		92.10	0.41	NIT 1	99	
Actinomyces israelii		7.90	0.37	TRE 90	1	
67357132	**Good identification**					**67357132**
					SPO	
Clostridium tertium		95.76	0.50	NIT 1	99	
Actinomyces israelii		4.11	0.41	TRE 90	1	
Bifidobacterium breve		0.14	0.10		1	
67357140	**Very good identification**					**67357140**
Actinomyces israelii		100.0	0.65			
67357142	**Very good identification**					**67357142**
					SPO	
Actinomyces israelii		99.71	0.68		1	
Bifidobacterium adolescentis		0.19	0.15		1	
Clostridium tertium		0.11	0.03		99	
67357150	**Good identification**					**67357150**
					SPO	
Actinomyces israelii		99.36	0.51	MLZ 13	1	
Clostridium tertium		0.64	-.01		99	
67357152	**Good identification**					**67357152**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		97.26	0.54	MLZ 13	1	
Bifidobacterium adolescentis		1.49	0.17		1	
Clostridium tertium		1.25	0.07		99	
67357160	**Good identification**					**67357160**
					SPO	
Actinomyces israelii		97.52	0.45	NAG 6	1	
Clostridium tertium		2.48	0.03		99	
67357162	**Acceptable identification**					**67357162**
					SPO	
Actinomyces israelii		94.01	0.48	NAG 6	1	
Clostridium tertium		4.79	0.11		99	
Bifidobacterium breve		1.20	0.10		1	
67357200	**Good identification**					**67357200**
Actinomyces israelii		100.0	0.43	RHA 7		
67357202	**Good identification**					**67357202**
Actinomyces israelii		100.0	0.46	RHA 7		
67357240	**Good identification**					**67357240**
Actinomyces israelii		100.0	0.36	RHA 7		
67357242	**Good identification**					**67357242**
Actinomyces israelii		100.0	0.39	RHA 7		
67357300	**Good identification**					**67357300**
Actinomyces israelii		100.0	0.53	RHA 7		
67357302	**Good identification**					**67357302**
					SPO	
Actinomyces israelii		99.44	0.56	RHA 7	1	
Clostridium tertium		0.56	0.03		99	
67357310	**Good identification**					**67357310**
					SPO	
Actinomyces israelii		96.78	0.39	RHA 7	1	
Clostridium tertium		3.22	-.01	MLZ 13	99	
67357312	**Acceptable identification**					**67357312**
					SPO	
Actinomyces israelii		93.76	0.42	RHA 7	1	
Clostridium tertium		6.25	0.07	MLZ 13	99	
67357322	**Low discrimination**					**67357322**
					SPO	
Actinomyces israelii		72.68	0.35	RHA 7	1	
Clostridium tertium		19.12	0.11	NAG 6	99	
Clostridium ramosum		8.20	0.03		92	
67357340	**Good identification**					**67357340**
Actinomyces israelii		100.0	0.46	RHA 7		
67357342	**Good identification**					**67357342**
Actinomyces israelii		100.0	0.49	RHA 7		
67357400	**Very good identification**					**67357400**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.60				
67357402	**Very good identification**						**67357402**
Actinomyces israelii		99.81	0.63				
Bifidobacterium adolescentis		0.19	0.11				
67357410	**Good identification**						**67357410**
Actinomyces israelii		100.0	0.46	MLZ 13			
67357412	**Good identification**						**67357412**
Actinomyces israelii		98.44	0.49	MLZ 13			
Bifidobacterium adolescentis		1.57	0.12				
67357420	**Good identification**						**67357420**
Actinomyces israelii		100.0	0.40	NAG 6			
67357422	**Good identification**						**67357422**
Actinomyces israelii		100.0	0.43	NAG 6			
67357440	**Good identification**						**67357440**
Actinomyces israelii		100.0	0.53				
67357442	**Very good identification**						**67357442**
Actinomyces israelii		100.0	0.56				
67357450	**Good identification**						**67357450**
Actinomyces israelii		100.0	0.39	MLZ 13			
67357452	**Good identification**						**67357452**
Actinomyces israelii		100.0	0.42	MLZ 13			
67357462	**Good identification**						**67357462**
Actinomyces israelii		100.0	0.36	NAG 6			
67357500	**Very good identification**						**67357500**
					GRA	SPO	
Actinomyces israelii		98.05	0.71		99	1	
Mitsuokella multacida		1.55	0.27		1	1	
Clostridium butyricum		0.33	0.07		99	99	
67357501	**Low discrimination**						**67357501**
					GRA		
Actinomyces israelii		85.18	0.36	URE 1	99		
Mitsuokella multacida		14.82	0.11		1		
67357502	**Very good identification**						**67357502**
					GRA		
Actinomyces israelii		95.18	0.74		99		
Bifidobacterium adolescentis		4.58	0.45		99		
Mitsuokella multacida		0.11	0.11		1		
67357503	**Good identification**						**67357503**
Actinomyces israelii		95.41	0.39	URE 1			
Bifidobacterium adolescentis		4.59	0.11				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67357510	**Very good identification**						**67357510**
					GRA		
Actinomyces israelii		95.76	0.56	MLZ 13	99		
Mitsuokella multacida		3.38	0.19		1		
Bifidobacterium adolescentis		0.57	0.12		99		
67357512	**Low discrimination**						**67357512**
Actinomyces israelii		70.39	0.59	MLZ 13			
Bifidobacterium adolescentis		27.69	0.46				
Bifidobacterium dentium		1.31	0.06				
67357520	**Acceptable identification**						**67357520**
					GRA	SPO	
Actinomyces israelii		93.11	0.50	NAG 6	99	1	
Mitsuokella multacida		5.77	0.17		1	1	
Clostridium tertium		1.13	0.03		99	99	
67357522	**Good identification**						**67357522**
					SPO		
Actinomyces israelii		96.92	0.53	NAG 6	1		
Clostridium tertium		2.35	0.11		99		
Bifidobacterium adolescentis		0.74	0.11		1		
67357530	**Low discrimination**						**67357530**
					GRA	SPO	
Actinomyces israelii		73.57	0.36	MLZ 13	99	1	
Clostridium tertium		10.59	0.07	NAG 6	99	99	
Mitsuokella multacida		10.17	0.09		1	1	
67357532	**Low discrimination**						**67357532**
					SPO		
Actinomyces israelii		53.48	0.39	MLZ 13	1		
Bifidobacterium dentium		27.79	0.10	NAG 6	1		
Clostridium tertium		15.40	0.15		99		
67357540	**Very good identification**						**67357540**
					GRA		
Actinomyces israelii		96.09	0.64		99		
Mitsuokella multacida		3.91	0.27		1		
67357542	**Very good identification**						**67357542**
					GRA		
Actinomyces israelii		98.20	0.67		99		
Bifidobacterium adolescentis		1.50	0.29		99		
Mitsuokella multacida		0.30	0.11		1		
67357550	**Acceptable identification**						**67357550**
					COC	GRA	
Actinomyces israelii		90.55	0.49	MLZ 13	1	99	
Mitsuokella multacida		8.22	0.19		1	1	
Blautia producta		1.23	-.03		99	99	
67357552	**Low discrimination**						**67357552**
					GRA		
Actinomyces israelii		88.36	0.52	MLZ 13	99		
Bifidobacterium adolescentis		11.05	0.31		99		
Mitsuokella multacida		0.59	0.03		1		
67357560	**Low discrimination**						**67357560**
					GRA		
Actinomyces israelii		86.26	0.43	NAG 6	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Mitsuokella multacida		13.74	0.17		1	
67357562	**Good identification**					**67357562**
Actinomyces israelii		100.0	0.46	NAG 6		
67357600	**Good identification**					**67357600**
Actinomyces israelii		100.0	0.41	RHA 7		
67357602	**Good identification**					**67357602**
Actinomyces israelii		100.0	0.44	RHA 7		
67357642	**Good identification**					**67357642**
Actinomyces israelii		100.0	0.37	RHA 7		
67357700	**Low discrimination**					**67357700**
					GRA	
Actinomyces israelii		71.93	0.51	RHA 7	99	
Mitsuokella multacida		28.08	0.32		1	
67357702	**Good identification**					**67357702**
					GRA	
Actinomyces israelii		96.58	0.54	RHA 7	99	
Mitsuokella multacida		2.79	0.15		1	
Bifidobacterium adolescentis		0.62	0.11		99	
67357710	**Low discrimination**					**67357710**
					GRA	
Actinomyces israelii		53.46	0.37	RHA 7	99	
Mitsuokella multacida		46.55	0.24	MLZ 13	1	
67357712	**Low discrimination**					**67357712**
					GRA	
Actinomyces israelii		89.50	0.40	RHA 7	99	
Mitsuokella multacida		5.77	0.07	MLZ 13	1	
Bifidobacterium adolescentis		4.73	0.12		99	
67357742	**Acceptable identification**					**67357742**
					GRA	
Actinomyces israelii		93.08	0.47	RHA 7	99	
Mitsuokella multacida		6.92	0.15		1	
67361101	**Low discrimination**					**67361101**
					SPO	
Actinomyces naeslundii		74.88	0.40	ESL 12	1	
Clostridium chauvoei		25.12	0.09	RAF 92	99	
67361103	**Good identification**					**67361103**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67361112	**Good identification**					**67361112**
Leptotrichia buccalis		100.0	0.50	NIT 10		
67361121	**Low discrimination**					**67361121**
					SPO	
Actinomyces naeslundii		89.94	0.40	ESL 12	1	
Clostridium chauvoei		10.06	0.01	RAF 92	99	
67361123	**Good identification**					**67361123**
Actinomyces naeslundii		100.0	0.40	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67363001	**Good identification**						**67363001**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67363003	**Good identification**						**67363003**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67363021	**Good identification**						**67363021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67363023	**Good identification**						**67363023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67363100	**Low discrimination**						**67363100**
					SPO		
Actinomyces naeslundii		52.63	0.42	ESL 12	1		
Actinomyces israelii		33.47	0.30	RAF 92	1		
Clostridium tertium		13.90	0.09	FRU 95	99		
67363101	**Good identification**						**67363101**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67363102	**Low discrimination**						**67363102**
					SPO		
Actinomyces naeslundii		34.55	0.42	ESL 12	1		
Actinomyces israelii		32.95	0.33	RAF 92	1		
Clostridium tertium		27.38	0.17	FRU 95	99		
67363103	**Good identification**						**67363103**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67363120	**Low discrimination**						**67363120**
					SPO		
Actinomyces naeslundii		44.35	0.42	ESL 12	1		
Clostridium tertium		35.15	0.17	RAF 92	99		
Clostridium ramosum		20.51	0.11	FRU 95	92		
67363121	**Good identification**						**67363121**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67363123	**Good identification**						**67363123**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67365102	**Low discrimination**						**67365102**
					COC		
Actinomyces israelii		66.93	0.38	FRU 99	1		
Atopobium parvulum		33.07	0.19		99		
67365502	**Low discrimination**						**67365502**
					GRA	SPO	
Actinomyces israelii		75.80	0.36	FRU 99	99	1	
Clostridium butyricum		14.06	0.03		99	99	
Mitsuokella multacida		5.68	0.04		1	1	
67367000	**Good identification**						**67367000**
Actinomyces israelii		100.0	0.37	FRU 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67367002	**Good identification**						**67367002**
Actinomyces israelii		100.0	0.40	FRU 99			
67367100	**Good identification**						**67367100**
					SPO		
Actinomyces israelii		95.25	0.48	FRU 99	1		
Clostridium tertium		4.75	0.11		99		
67367102	**Acceptable identification**						**67367102**
					SPO		
Actinomyces israelii		90.93	0.51	FRU 99	1		
Clostridium tertium		9.07	0.19		99		
67367140	**Good identification**						**67367140**
Actinomyces israelii		100.0	0.41	FRU 99			
67367142	**Good identification**						**67367142**
Actinomyces israelii		100.0	0.44	FRU 99			
67367400	**Good identification**						**67367400**
Actinomyces israelii		100.0	0.36	FRU 99			
67367402	**Good identification**						**67367402**
Actinomyces israelii		100.0	0.39	FRU 99			
67367500	**Low discrimination**						**67367500**
					GRA	SPO	
Actinomyces israelii		69.91	0.46	FRU 99	99	1	
Mitsuokella multacida		29.11	0.27		1	1	
Clostridium butyricum		0.98	-.07		99	99	
67367502	**Good identification**						**67367502**
					GRA		
Actinomyces israelii		96.21	0.49	FRU 99	99		
Mitsuokella multacida		2.97	0.11		1		
Bifidobacterium adolescentis		0.82	0.08		99		
67367512	**Low discrimination**						**67367512**
Actinomyces israelii		61.37	0.35	FRU 99			
Bifidobacterium dentium		30.13	0.06	MLZ 13			
Bifidobacterium adolescentis		4.28	0.09				
67367542	**Acceptable identification**						**67367542**
					GRA		
Actinomyces israelii		92.65	0.42	FRU 99	99		
Mitsuokella multacida		7.35	0.11		1		
67370101	**Good identification**						**67370101**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67370103	**Low discrimination**						**67370103**
Actinomyces naeslundii		89.63	0.39	ESL 12			
Actinomyces viscosus		10.37	-.02	RAF 92			
67370112	**Good identification**						**67370112**
Leptotrichia buccalis		100.0	0.50	NIT 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67370121	**Good identification**					**67370121**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67370123	**Good identification**					**67370123**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67371000	**Very good identification**					**67371000**
Lactobacillus jensenii		96.50	0.66	NIT 1		
Actinomyces naeslundii		0.75	0.51			
Lactobacillus catenaformis		0.65	0.29			
67371001	**Low discrimination**					**67371001**
Actinomyces naeslundii		53.71	0.54	ESL 12		
Lactobacillus jensenii		46.30	0.32	RAF 92		
67371002	**Low discrimination**					**67371002**
					SPO	
Lactobacillus jensenii		86.26	0.63	NIT 1	1	
Lactobacillus acidophilus		7.13	0.41		1	
Clostridium barati		2.51	0.31		99	
67371003	**Low discrimination**					**67371003**
Actinomyces naeslundii		61.02	0.54	ESL 12		
Lactobacillus jensenii		35.07	0.29	RAF 92		
Lactobacillus acidophilus		2.90	0.07			
67371010	**Very good identification**					**67371010**
					GRA	
Lactobacillus jensenii		97.80	0.58	NIT 1	99	
Leptotrichia buccalis		1.81	0.15		1	
Actinomyces israelii		0.28	0.26		99	
67371012	**Low discrimination**					**67371012**
					GRA	
Leptotrichia buccalis		73.07	0.50	NIT 10	1	
Lactobacillus jensenii		26.62	0.55	LAC 99	99	
Actinomyces israelii		0.17	0.29		99	
67371020	**Low discrimination**					**67371020**
					SPO	
Lactobacillus jensenii		82.05	0.56	NIT 1	1	
Clostridium barati		6.38	0.31		99	
Clostridium perfringens		4.04	0.32		84	
67371021	**Low discrimination**					**67371021**
					SPO	
Actinomyces naeslundii		81.15	0.54	ESL 12	1	
Lactobacillus jensenii		17.49	0.21	RAF 92	1	
Clostridium barati		1.36	-.03		99	
67371022	**Low discrimination**					**67371022**
					SPO	
Lactobacillus jensenii		55.04	0.53	NIT 1	1	
Clostridium barati		19.24	0.40		99	
Lactobacillus acidophilus		18.19	0.41		1	
67371023	**Low discrimination**					**67371023**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		80.56	0.54	ESL 12	1		
Lactobacillus jensenii		11.57	0.18	RAF 92	1		
Clostridium barati		4.05	0.05		99		
67371030	**Good identification**						**67371030**
					GRA		
Lactobacillus jensenii		98.47	0.47	NIT 1	99		
Leptotrichia buccalis		0.81	-.01		1		
Actinomyces naeslundii		0.49	0.29		99		
67371041	**Good identification**						**67371041**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67371043	**Good identification**						**67371043**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67371061	**Good identification**						**67371061**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67371063	**Good identification**						**67371063**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67371100	**Low discrimination**						**67371100**
					COC		
Atopobium parvulum		77.49	0.61	NIT 1	99		
Collinsella aerofaciens		8.65	0.32		1		
Actinomyces israelii		2.98	0.51		1		
67371101	**Low discrimination**						**67371101**
					COC	SPO	
Actinomyces naeslundii		78.39	0.62	ESL 12	1	1	
Atopobium parvulum		15.58	0.27	RAF 92	99	1	
Clostridium chauvoei		3.23	0.15		1	99	
67371102	**Low discrimination**						**67371102**
					COC	GRA	
Lactobacillus acidophilus		41.66	0.60	NIT 1	1	99	
Leptotrichia buccalis		40.13	0.50		1	1	
Atopobium parvulum		7.62	0.48		99	99	
67371103	**Low discrimination**						**67371103**
					GRA		
Actinomyces naeslundii		69.06	0.62	ESL 12	99		
Lactobacillus acidophilus		13.25	0.25	RAF 92	99		
Leptotrichia buccalis		12.76	0.15		1		
67371110	**Good identification**						**67371110**
					COC	GRA	
Leptotrichia buccalis		97.18	0.50	NIT 10	1	1	
Atopobium parvulum		1.06	0.27	BGL 99	99	99	
Actinomyces israelii		0.60	0.37		1	99	
67371112	**Very good identification**						**67371112**
					GRA		
Leptotrichia buccalis		99.95	0.84	NIT 10	1		
Bifidobacterium breve		0.01	0.37		99		
Actinomyces israelii		0.01	0.40		99		
67371113	**Good identification**						**67371113**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		99.71	0.50	NIT 10	1		
Actinomyces naeslundii		0.29	0.40	URE 1	99		
67371120	**Low discrimination**						**67371120**
					COC	SPO	
Clostridium paraputrificum		75.08	0.56	NIT 1	1	99	
Atopobium parvulum		8.06	0.48		99	1	
Clostridium perfringens		6.28	0.47		1	84	
67371121	**Low discrimination**						**67371121**
					COC	SPO	
Actinomyces naeslundii		69.72	0.62	ESL 12	1	1	
Clostridium paraputrificum		22.78	0.22	RAF 92	1	99	
Atopobium parvulum		2.45	0.14		99	1	
67371122	**Low discrimination**						**67371122**
					SPO		
Lactobacillus acidophilus		41.28	0.60	NIT 1	1		
Clostridium paraputrificum		30.13	0.50		99		
Clostridium barati		11.40	0.48		99		
67371123	**Low discrimination**						**67371123**
					SPO		
Actinomyces naeslundii		69.13	0.62	ESL 12	1		
Lactobacillus acidophilus		13.26	0.25	RAF 92	1		
Clostridium paraputrificum		9.68	0.16		99		
67371131	**Low discrimination**						**67371131**
					GRA		
Actinomyces naeslundii		71.94	0.40	ESL 12	99		
Leptotrichia buccalis		28.06	-.01	RAF 92	1		
67371132	**Very good identification**						**67371132**
					GRA	SPO	
Leptotrichia buccalis		99.41	0.67	NIT 10	1	1	
Bifidobacterium breve		0.20	0.41	NAG 10	99	1	
Clostridium tertium		0.14	0.29		99	99	
67371140	**Low discrimination**						**67371140**
					COC		
Actinomyces israelii		38.19	0.44	XYL 92	1		
Atopobium parvulum		25.79	0.27		99		
Actinomyces naeslundii		15.27	0.46		1		
67371141	**Good identification**						**67371141**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67371142	**Low discrimination**						**67371142**
Lactobacillus acidophilus		66.55	0.44	NIT 1			
Actinomyces israelii		12.51	0.47	SOR 11			
Bifidobacterium breve		6.55	0.37				
67371143	**Low discrimination**						**67371143**
Actinomyces naeslundii		88.16	0.49	ESL 12			
Lactobacillus acidophilus		11.84	0.10	RAF 92			
67371152	**Good identification**						**67371152**
					GRA		
Leptotrichia buccalis		98.24	0.50	NIT 10	1		
Bifidobacterium breve		1.24	0.37	SOR 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.35	0.33		99		
67371160	**Low discrimination**						**67371160**
Clostridium paraputrificum		72.64	0.37	NIT 1			
Clostridium perfringens		15.37	0.35	SOR 7			
Clostridium barati		4.53	0.21				
67371161	**Low discrimination**						**67371161**
					SPO		
Actinomyces naeslundii		85.56	0.49	ESL 12	1		
Clostridium paraputrificum		11.92	0.03	RAF 92	99		
Clostridium perfringens		2.52	0.01		84		
67371162	**Low discrimination**						**67371162**
					SPO		
Lactobacillus acidophilus		50.00	0.44	NIT 1	1		
Clostridium paraputrificum		22.23	0.31	SOR 11	99		
Clostridium barati		9.71	0.30		99		
67371163	**Low discrimination**						**67371163**
					SPO		
Actinomyces naeslundii		83.75	0.49	ESL 12	1		
Lactobacillus acidophilus		11.25	0.10	RAF 92	1		
Clostridium paraputrificum		5.00	-.04		99		
67371302	**Low discrimination**						**67371302**
					GRA	SPO	
Lactobacillus acidophilus		74.09	0.40	NIT 1	99	1	
Leptotrichia buccalis		9.58	0.15	RHA 7	1	1	
Clostridium barati		7.88	0.21		99	99	
67371312	**Good identification**						**67371312**
					GRA		
Leptotrichia buccalis		99.81	0.50	NIT 10	1		
Lactobacillus acidophilus		0.08	0.06	RHA 1	99		
Actinomyces israelii		0.07	0.21		99		
67371322	**Low discrimination**						**67371322**
					SPO		
Clostridium ramosum		56.93	0.45	NIT 1	92		
Lactobacillus acidophilus		27.89	0.40		1		
Clostridium barati		8.89	0.30		99		
67371402	**Low discrimination**						**67371402**
Actinomyces israelii		50.84	0.42	XYL 92			
Lactobacillus jensenii		44.22	0.29				
Lactobacillus acidophilus		3.65	0.07				
67371442	**Good identification**						**67371442**
Actinomyces israelii		100.0	0.35	XYL 92			
67371500	**Low discrimination**						**67371500**
					COC	SPO	
Actinomyces israelii		57.17	0.49	XYL 92	1	1	
Atopobium parvulum		18.35	0.27		99	1	
Clostridium butyricum		14.55	0.18		1	99	
67371502	**Low discrimination**						**67371502**
					GRA		
Actinomyces israelii		60.77	0.52	XYL 92	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Lactobacillus acidophilus		12.55	0.25		99		
Leptotrichia buccalis		12.09	0.15		1		
67371512	**Good identification**						**67371512**
					GRA		
Leptotrichia buccalis		98.54	0.50	NIT 10	1		
Actinomyces israelii		0.75	0.38	ARA 1	99		
Bifidobacterium adolescentis		0.62	0.31		99		
67371540	**Low discrimination**						**67371540**
					GRA	SPO	
Actinomyces israelii		73.59	0.42	XYL 92	99	1	
Mitsuokella multacida		22.22	0.21		1	1	
Clostridium butyricum		4.19	0.00		99	99	
67371542	**Low discrimination**						**67371542**
Actinomyces israelii		88.11	0.45	XYL 92			
Lactobacillus acidophilus		5.78	0.10				
Bifidobacterium adolescentis		2.84	0.14				
67372001	**Good identification**						**67372001**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67372003	**Good identification**						**67372003**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67372021	**Good identification**						**67372021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67372023	**Good identification**						**67372023**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67372100	**Low discrimination**						**67372100**
					SPO		
Actinomyces naeslundii		46.80	0.40	ESL 12	1		
Actinomyces israelii		37.59	0.30	RAF 92	1		
Clostridium tertium		15.62	0.09	GAL 96	99		
67372101	**Good identification**						**67372101**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67372103	**Good identification**						**67372103**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67372121	**Good identification**						**67372121**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67372123	**Good identification**						**67372123**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67373000	**Low discrimination**						**67373000**
Actinomyces israelii		56.29	0.53	XYL 92			
Actinomyces naeslundii		23.88	0.56				
Lactobacillus jensenii		15.90	0.32				
67373001	**Very good identification**						**67373001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		98.44	0.59	ESL 12			
Actinomyces israelii		1.56	0.19	RAF 92			
67373002	**Low discrimination**						**67373002**
Actinomyces israelii		64.42	0.56	XYL 92			
Actinomyces naeslundii		18.22	0.56				
Lactobacillus jensenii		8.09	0.29				
67373003	**Very good identification**						**67373003**
Actinomyces naeslundii		97.67	0.59	ESL 12			
Actinomyces israelii		2.33	0.22	RAF 92			
67373010	**Low discrimination**						**67373010**
					SPO		
Actinomyces israelii		50.52	0.39	XYL 92	1		
Lactobacillus jensenii		31.84	0.24	MLZ 13	1		
Clostridium tertium		10.09	0.13		99		
67373011	**Good identification**						**67373011**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67373012	**Low discrimination**						**67373012**
					GRA	SPO	
Actinomyces israelii		39.24	0.42	XYL 92	99	1	
Leptotrichia buccalis		30.18	0.15	MLZ 13	1	1	
Clostridium tertium		15.68	0.21		99	99	
67373013	**Good identification**						**67373013**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67373020	**Low discrimination**						**67373020**
Actinomyces naeslundii		61.87	0.56	ESL 12			
Lactobacillus jensenii		10.30	0.21	RAF 92			
Actinomyces israelii		9.31	0.33				
67373021	**Very good identification**						**67373021**
Actinomyces naeslundii		100.0	0.59	ESL 12			
67373022	**Low discrimination**						**67373022**
					SPO		
Actinomyces naeslundii		40.08	0.56	ESL 12	1		
Clostridium tertium		14.27	0.25	RAF 92	99		
Clostridium barati		13.39	0.21		99		
67373023	**Very good identification**						**67373023**
Actinomyces naeslundii		100.0	0.59	ESL 12			
67373031	**Good identification**						**67373031**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67373033	**Good identification**						**67373033**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67373040	**Low discrimination**						**67373040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		83.86	0.46	XYL 92		
Actinomyces naeslundii		16.14	0.43			
67373041	**Good identification**					**67373041**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67373042	**Low discrimination**					**67373042**
Actinomyces israelii		87.30	0.49	XYL 92		
Actinomyces naeslundii		11.20	0.43			
Lactobacillus acidophilus		1.50	0.04			
67373043	**Good identification**					**67373043**
Actinomyces naeslundii		95.01	0.46	ESL 12		
Actinomyces israelii		4.99	0.15	RAF 92		
67373052	**Good identification**					**67373052**
Actinomyces israelii		100.0	0.35	XYL 92		
67373060	**Low discrimination**					**67373060**
Actinomyces naeslundii		75.10	0.43	ESL 12		
Actinomyces israelii		24.90	0.26	RAF 92		
67373061	**Good identification**					**67373061**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67373062	**Low discrimination**					**67373062**
					SPO	
Actinomyces naeslundii		55.68	0.43	ESL 12	1	
Actinomyces israelii		27.70	0.29	RAF 92	1	
Clostridium barati		9.17	0.03		99	
67373063	**Good identification**					**67373063**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67373100	**Low discrimination**					**67373100**
					SPO	
Actinomyces israelii		55.69	0.64	XYL 92	1	
Clostridium tertium		23.14	0.43		99	
Actinomyces naeslundii		16.81	0.64		1	
67373101	**Very good identification**					**67373101**
					SPO	
Actinomyces naeslundii		96.94	0.67	ESL 12	1	
Actinomyces israelii		2.16	0.30	RAF 92	1	
Clostridium tertium		0.90	0.09		99	
67373102	**Low discrimination**					**67373102**
					SPO	
Actinomyces israelii		43.43	0.67	XYL 92	1	
Clostridium tertium		36.09	0.51		99	
Actinomyces naeslundii		8.74	0.64		1	
67373103	**Good identification**					**67373103**
					SPO	
Actinomyces naeslundii		93.77	0.67	ESL 12	1	
Actinomyces israelii		3.14	0.33	RAF 92	1	
Clostridium tertium		2.61	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67373110	**Low discrimination**						**67373110**
					GRA	SPO	
Clostridium tertium		77.56	0.47	NIT 1	99	99	
Actinomyces israelii		15.69	0.50		99	1	
Leptotrichia buccalis		4.53	0.15		1	1	
67373111	**Low discrimination**						**67373111**
					SPO		
Actinomyces naeslundii		72.65	0.45	ESL 12	1		
Clostridium tertium		22.75	0.13	RAF 92	99		
Actinomyces israelii		4.60	0.15	MLZ 5	1		
67373112	**Low discrimination**						**67373112**
					GRA	SPO	
Leptotrichia buccalis		62.91	0.50	NIT 10	1	1	
Clostridium tertium		32.68	0.55	MAN 1	99	99	
Actinomyces israelii		3.31	0.53		99	1	
67373120	**Low discrimination**						**67373120**
					SPO		
Clostridium tertium		50.27	0.51	NIT 1	99		
Clostridium ramosum		29.33	0.45		92		
Actinomyces naeslundii		12.17	0.64		1		
67373121	**Very good identification**						**67373121**
					SPO		
Actinomyces naeslundii		95.78	0.67	ESL 12	1		
Clostridium tertium		2.66	0.17	RAF 92	99		
Clostridium ramosum		1.55	0.11		92		
67373122	**Low discrimination**						**67373122**
					SPO		
Clostridium tertium		49.70	0.59	NIT 1	99		
Clostridium ramosum		38.67	0.56		92		
Actinomyces naeslundii		4.01	0.64		1		
67373123	**Low discrimination**						**67373123**
					SPO		
Actinomyces naeslundii		86.69	0.67	ESL 12	1		
Clostridium tertium		7.23	0.25	RAF 92	99		
Clostridium ramosum		5.63	0.21		92		
67373130	**Good identification**						**67373130**
					SPO		
Clostridium tertium		97.92	0.55	NIT 1	99		
Actinomyces naeslundii		0.70	0.42		1		
Actinomyces israelii		0.42	0.29		1		
67373131	**Low discrimination**						**67373131**
					SPO		
Actinomyces naeslundii		51.56	0.45	ESL 12	1		
Clostridium tertium		48.44	0.21	RAF 92	99		
67373132	**Good identification**						**67373132**
					GRA	SPO	
Clostridium tertium		91.32	0.63	NIT 1	99	99	
Leptotrichia buccalis		6.51	0.33		1	1	
Bifidobacterium breve		1.32	0.41		99	1	
67373140	**Low discrimination**						**67373140**
Actinomyces israelii		86.10	0.57	XYL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		11.79	0.51			
Bifidobacterium breve		1.18	0.20			
67373141	**Good identification**					**67373141**
Actinomyces naeslundii		95.31	0.54	ESL 12		
Actinomyces israelii		4.69	0.23	RAF 92		
67373142	**Low discrimination**					**67373142**
Actinomyces israelii		79.37	0.60	XYL 92		
Bifidobacterium breve		7.33	0.37			
Actinomyces naeslundii		7.25	0.51			
67373143	**Acceptable identification**					**67373143**
Actinomyces naeslundii		93.13	0.54	ESL 12		
Actinomyces israelii		6.87	0.26	RAF 92		
67373150	**Low discrimination**					**67373150**
					SPO	
Actinomyces israelii		78.84	0.43	XYL 92	1	
Clostridium tertium		10.12	0.13	MLZ 13	99	
Bifidobacterium breve		7.23	0.20		1	
67373152	**Low discrimination**					**67373152**
					GRA	
Actinomyces israelii		41.64	0.46	XYL 92	99	
Bifidobacterium breve		25.74	0.37	MLZ 13	99	
Leptotrichia buccalis		20.59	0.15		1	
67373160	**Low discrimination**					**67373160**
					SPO	
Actinomyces naeslundii		45.32	0.51	ESL 12	1	
Actinomyces israelii		21.12	0.36	RAF 92	1	
Clostridium tertium		10.71	0.17		99	
67373161	**Good identification**					**67373161**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67373162	**Low discrimination**					**67373162**
					SPO	
Bifidobacterium breve		32.59	0.41	NIT 1	1	
Actinomyces naeslundii		18.12	0.51	RAF 95	1	
Clostridium tertium		12.85	0.25		99	
67373163	**Good identification**					**67373163**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67373172	**Low discrimination**					**67373172**
					SPO	
Bifidobacterium breve		54.40	0.41	NIT 1	1	
Clostridium tertium		38.13	0.29	RAF 95	99	
Actinomyces israelii		3.16	0.25		1	
67373202	**Acceptable identification**					**67373202**
Actinomyces israelii		94.87	0.37	XYL 92		
Lactobacillus acidophilus		5.13	0.00	RHA 7		
67373300	**Low discrimination**					**67373300**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		81.59	0.45	XYL 92	1		
Clostridium ramosum		10.56	0.13	RHA 7	92		
Clostridium tertium		4.55	0.09		99		
67373302	**Low discrimination**						**67373302**
					SPO		
Actinomyces israelii		60.02	0.48	XYL 92	1		
Clostridium ramosum		20.71	0.24	RHA 7	92		
Lactobacillus acidophilus		9.33	0.18		1		
67373320	**Good identification**						**67373320**
					SPO		
Clostridium ramosum		97.62	0.47	NIT 1	92		
Clostridium tertium		1.28	0.17		99		
Actinomyces israelii		0.49	0.24		1		
67373322	**Very good identification**						**67373322**
					SPO		
Clostridium ramosum		98.10	0.58	NIT 1	92		
Clostridium tertium		0.96	0.25		99		
Lactobacillus acidophilus		0.45	0.18		1		
67373340	**Good identification**						**67373340**
Actinomyces israelii		100.0	0.38	XYL 92			
67373342	**Good identification**						**67373342**
Actinomyces israelii		95.29	0.41	XYL 92			
Lactobacillus acidophilus		4.71	0.03	RHA 7			
67373400	**Good identification**						**67373400**
					SPO		
Actinomyces israelii		99.20	0.52	XYL 92	1		
Clostridium innocuum		0.80	0.01		99		
67373402	**Good identification**						**67373402**
					SPO		
Actinomyces israelii		99.47	0.55	XYL 92	1		
Clostridium innocuum		0.53	0.01		99		
67373410	**Good identification**						**67373410**
Actinomyces israelii		100.0	0.38	XYL 92			
67373412	**Good identification**						**67373412**
Actinomyces israelii		100.0	0.41	XYL 92			
67373440	**Good identification**						**67373440**
Actinomyces israelii		100.0	0.45	XYL 92			
67373442	**Good identification**						**67373442**
Actinomyces israelii		100.0	0.48	XYL 92			
67373500	**Good identification**						**67373500**
					GRA	SPO	
Actinomyces israelii		94.14	0.62	XYL 92	99	1	
Mitsuokella multacida		4.55	0.27		1	1	
Clostridium tertium		0.48	0.09		99	99	
67373502	**Very good identification**						**67373502**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		96.70	0.65	XYL 92	1		
Bifidobacterium adolescentis		1.42	0.28		1		
Clostridium tertium		0.99	0.17		99		
67373510	**Low discrimination**						**67373510**
					GRA	SPO	
Actinomyces israelii		85.55	0.48	XYL 92	99	1	
Mitsuokella multacida		9.23	0.19	MLZ 13	1	1	
Clostridium tertium		5.22	0.13		99	99	
67373512	**Low discrimination**						**67373512**
					GRA	SPO	
Actinomyces israelii		64.86	0.51	XYL 92	99	1	
Leptotrichia buccalis		15.24	0.15	MLZ 13	1	1	
Clostridium tertium		7.92	0.21		99	99	
67373520	**Low discrimination**						**67373520**
					GRA	SPO	
Actinomyces israelii		61.37	0.42	XYL 92	99	1	
Clostridium tertium		14.80	0.17	NAG 6	99	99	
Mitsuokella multacida		11.62	0.17		1	1	
67373522	**Low discrimination**						**67373522**
					SPO		
Actinomyces israelii		51.91	0.45	XYL 92	1		
Clostridium tertium		25.03	0.25	NAG 6	99		
Clostridium ramosum		19.48	0.21		92		
67373540	**Low discrimination**						**67373540**
					GRA		
Actinomyces israelii		88.94	0.55	XYL 92	99		
Mitsuokella multacida		11.06	0.27		1		
67373542	**Very good identification**						**67373542**
					GRA		
Actinomyces israelii		98.63	0.58	XYL 92	99		
Mitsuokella multacida		0.91	0.11		1		
Bifidobacterium adolescentis		0.46	0.12		99		
67373550	**Low discrimination**						**67373550**
					COC	GRA	
Actinomyces israelii		75.81	0.41	XYL 92	1	99	
Mitsuokella multacida		21.03	0.19	MLZ 13	1	1	
Blautia producta		3.16	-.03		99	99	
67373552	**Acceptable identification**						**67373552**
					GRA		
Actinomyces israelii		94.46	0.44	XYL 92	99		
Bifidobacterium adolescentis		3.60	0.14	MLZ 13	99		
Mitsuokella multacida		1.94	0.03		1		
67373562	**Good identification**						**67373562**
Actinomyces israelii		100.0	0.38	XYL 92			
67373602	**Good identification**						**67373602**
Actinomyces israelii		100.0	0.36	XYL 92			
67373702	**Acceptable identification**						**67373702**
					GRA		
Actinomyces israelii		91.88	0.46	XYL 92	99		
Mitsuokella multacida		8.12	0.15	RHA 7	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67373742	**Low discrimination**					**67373742**
					GRA	
Actinomyces israelii		81.48	0.39	XYL 92	99	
Mitsuokella multacida		18.52	0.15	RHA 7	1	
67374102	**Low discrimination**					**67374102**
					COC	
Actinomyces israelii		66.93	0.38	GAL 99	1	
Atopobium parvulum		33.07	0.19		99	
67374502	**Low discrimination**					**67374502**
					SPO	
Actinomyces israelii		69.49	0.36	GAL 99	1	
Clostridium butyricum		12.89	0.03		99	
Clostridium clostridioforme		8.32	0.11		66	
67374712	**Good identification**					**67374712**
Clostridium clostridioforme		98.85	0.37	NIT 1		
Mitsuokella multacida		1.15	0.01	RAF 94		
67374732	**Good identification**					**67374732**
Clostridium clostridioforme		100.0	0.37	NIT 1		
67375000	**Low discrimination**					**67375000**
					COC	
Actinomyces israelii		75.77	0.59		1	
Atopobium parvulum		12.86	0.32		99	
Lactobacillus jensenii		10.55	0.32		1	
67375002	**Good identification**					**67375002**
					COC	
Actinomyces israelii		91.61	0.62		1	
Lactobacillus jensenii		5.67	0.29		1	
Atopobium parvulum		1.83	0.19		99	
67375010	**Low discrimination**					**67375010**
Actinomyces israelii		76.30	0.45	MLZ 13		
Lactobacillus jensenii		23.70	0.24			
67375012	**Low discrimination**					**67375012**
					GRA	
Actinomyces israelii		65.92	0.48	MLZ 13	99	
Leptotrichia buccalis		24.98	0.15		1	
Lactobacillus jensenii		9.10	0.21		99	
67375020	**Low discrimination**					**67375020**
					COC	
Actinomyces israelii		48.62	0.38	NAG 6	1	
Lactobacillus jensenii		26.51	0.21		1	
Atopobium parvulum		22.81	0.19		99	
67375022	**Low discrimination**					**67375022**
					SPO	
Actinomyces israelii		68.38	0.41	NAG 6	1	
Lactobacillus jensenii		16.57	0.18		1	
Clostridium barati		5.80	0.05		99	
67375040	**Good identification**					**67375040**
Actinomyces israelii		100.0	0.52			

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
67375042	**Good identification**						**67375042**
Actinomyces israelii		100.0	0.55				
67375050	**Good identification**						**67375050**
Actinomyces israelii		100.0	0.38	MLZ 13			
67375052	**Good identification**						**67375052**
Actinomyces israelii		100.0	0.41	MLZ 13			
67375100	**Low discrimination**						**67375100**
					COC	SPO	
Atopobium parvulum		79.54	0.66	NIT 1	99	1	
Actinomyces israelii		18.94	0.69		1	1	
Clostridium butyricum		1.23	0.28		1	99	
67375102	**Low discrimination**						**67375102**
					COC	SPO	
Actinomyces israelii		64.30	0.72		1	1	
Atopobium parvulum		31.77	0.53		99	1	
Clostridium butyricum		1.08	0.21		1	99	
67375103	**Low discrimination**						**67375103**
					COC		
Actinomyces israelii		63.67	0.38	URE 1	1		
Atopobium parvulum		31.45	0.19		99		
Actinomyces naeslundii		4.88	0.28		1		
67375110	**Low discrimination**						**67375110**
					COC	GRA	
Actinomyces israelii		66.60	0.55	MLZ 13	1	99	
Atopobium parvulum		18.91	0.32		99	99	
Leptotrichia buccalis		9.47	0.15		1	1	
67375112	**Low discrimination**						**67375112**
					GRA	SPO	
Leptotrichia buccalis		88.89	0.50	NIT 10	1	1	
Actinomyces israelii		9.48	0.58	XYL 1	99	1	
Clostridium tertium		0.64	0.24		99	99	
67375120	**Low discrimination**						**67375120**
					COC	SPO	
Atopobium parvulum		85.54	0.53	NIT 1	99	1	
Actinomyces israelii		7.37	0.49		1	1	
Clostridium paraputrificum		4.33	0.22		1	99	
67375122	**Low discrimination**						**67375122**
					COC	SPO	
Atopobium parvulum		41.93	0.40	NIT 1	99	1	
Actinomyces israelii		30.70	0.52		1	1	
Clostridium tertium		8.24	0.27		1	99	
67375140	**Low discrimination**						**67375140**
					COC	GRA	
Actinomyces israelii		87.33	0.62		1	99	
Atopobium parvulum		9.53	0.32		99	99	
Mitsuokella multacida		1.88	0.21		1	1	
67375142	**Very good identification**						**67375142**
					COC		
Actinomyces israelii		97.55	0.65		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Atopobium parvulum		1.25	0.19		99		
Lactobacillus acidophilus		0.46	0.10		1		
67375150	**Good identification**						**67375150**
					GRA		
Actinomyces israelii		95.43	0.48	MLZ 13	99		
Mitsuokella multacida		4.57	0.13		1		
67375152	**Low discrimination**						**67375152**
					GRA		
Actinomyces israelii		79.22	0.51	MLZ 13	99		
Leptotrichia buccalis		19.30	0.15		1		
Bifidobacterium adolescentis		1.49	0.15		99		
67375160	**Low discrimination**						**67375160**
					COC	SPO	
Actinomyces israelii		65.63	0.42	NAG 6	1	1	
Atopobium parvulum		19.79	0.19		99	1	
Clostridium paraputrificum		7.47	0.03		1	99	
67375162	**Low discrimination**						**67375162**
					COC		
Actinomyces israelii		87.63	0.45	NAG 6	1		
Lactobacillus acidophilus		6.41	0.10		1		
Atopobium parvulum		3.11	0.06		99		
67375200	**Good identification**						**67375200**
Actinomyces israelii		100.0	0.39	RHA 7			
67375202	**Good identification**						**67375202**
Actinomyces israelii		97.84	0.43	RHA 7			
Clostridium clostridioforme		2.16	0.05				
67375242	**Good identification**						**67375242**
Actinomyces israelii		100.0	0.35	RHA 7			
67375300	**Low discrimination**						**67375300**
					COC	GRA	
Actinomyces israelii		55.67	0.50	RHA 7	1	99	
Atopobium parvulum		31.37	0.32		99	99	
Mitsuokella multacida		11.48	0.25		1	1	
67375302	**Low discrimination**						**67375302**
					COC		
Actinomyces israelii		89.15	0.53	RHA 7	1		
Atopobium parvulum		5.91	0.19		99		
Clostridium clostridioforme		1.65	0.14		1		
67375310	**Low discrimination**						**67375310**
					GRA		
Actinomyces israelii		68.50	0.36	RHA 7	99		
Mitsuokella multacida		31.50	0.17	MLZ 13	1		
67375340	**Low discrimination**						**67375340**
					GRA		
Actinomyces israelii		65.35	0.43	RHA 7	99		
Mitsuokella multacida		34.65	0.25		1		
67375342	**Good identification**						**67375342**
					GRA		
Actinomyces israelii		96.22	0.46	RHA 7	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		3.78	0.09		1		
67375400	**Low discrimination**						**67375400**
					GRA	SPO	
Actinomyces israelii		86.20	0.57		99	1	
Clostridium butyricum		10.28	0.20		99	99	
Mitsuokella multacida		3.52	0.21		1	1	
67375402	**Very good identification**						**67375402**
					SPO		
Actinomyces israelii		96.09	0.60		1		
Clostridium butyricum		2.97	0.13		99		
Clostridium clostridioforme		0.42	0.10		66		
67375410	**Low discrimination**						**67375410**
					GRA	SPO	
Actinomyces israelii		88.25	0.43	MLZ 13	99	1	
Mitsuokella multacida		8.04	0.13		1	1	
Clostridium butyricum		3.71	-.02		99	99	
67375412	**Low discrimination**						**67375412**
Actinomyces israelii		86.83	0.46	MLZ 13			
Clostridium clostridioforme		11.48	0.22				
Bifidobacterium adolescentis		1.69	0.11				
67375420	**Low discrimination**						**67375420**
					GRA	SPO	
Actinomyces israelii		75.62	0.37	NAG 6	99	1	
Clostridium butyricum		12.28	0.02		99	99	
Mitsuokella multacida		12.10	0.11		1	1	
67375422	**Acceptable identification**						**67375422**
					SPO		
Actinomyces israelii		90.09	0.40	NAG 6	1		
Clostridium clostridioforme		6.12	0.10		66		
Clostridium butyricum		3.79	-.05		99		
67375440	**Low discrimination**						**67375440**
					GRA	SPO	
Actinomyces israelii		88.36	0.50		99	1	
Mitsuokella multacida		9.28	0.21		1	1	
Clostridium butyricum		2.36	0.02		99	99	
67375442	**Good identification**						**67375442**
					GRA	SPO	
Actinomyces israelii		98.55	0.53		99	1	
Mitsuokella multacida		0.77	0.04		1	1	
Clostridium butyricum		0.68	-.05		99	99	
67375450	**Low discrimination**						**67375450**
					GRA		
Actinomyces israelii		81.02	0.36	MLZ 13	99		
Mitsuokella multacida		18.99	0.13		1		
67375452	**Good identification**						**67375452**
Actinomyces israelii		100.0	0.39	MLZ 13			
67375500	**Low discrimination**						**67375500**
					GRA	SPO	
Mitsuokella multacida		36.38	0.55	RAF 99	1	1	
Actinomyces israelii		35.98	0.68		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		25.74	0.44		99	99	
67375501	**Low discrimination**						**67375501**
					GRA	SPO	
Mitsuokella multacida		86.64	0.39	RAF 99	1	1	
Actinomyces israelii		7.79	0.33	URE 10	99	1	
Clostridium butyricum		5.57	0.10		99	99	
67375502	**Low discrimination**						**67375502**
					GRA	SPO	
Actinomyces israelii		75.24	0.71		99	1	
Clostridium butyricum		13.95	0.37		99	99	
Mitsuokella multacida		5.64	0.39		1	1	
67375503	**Low discrimination**						**67375503**
					GRA	SPO	
Actinomyces israelii		48.35	0.36	URE 1	99	1	
Mitsuokella multacida		39.83	0.22		1	1	
Clostridium butyricum		8.97	0.03		99	99	
67375510	**Low discrimination**						**67375510**
					GRA	SPO	
Mitsuokella multacida		64.17	0.47	RAF 99	1	1	
Actinomyces israelii		28.45	0.53		99	1	
Clostridium butyricum		7.17	0.22		99	99	
67375512	**Low discrimination**						**67375512**
					GRA		
Actinomyces israelii		50.77	0.56	MLZ 13	99		
Bifidobacterium adolescentis		24.51	0.45		99		
Mitsuokella multacida		8.48	0.30		1		
67375520	**Low discrimination**						**67375520**
					GRA	SPO	
Mitsuokella multacida		65.16	0.45	RAF 99	1	1	
Actinomyces israelii		16.45	0.47		99	1	
Clostridium butyricum		16.03	0.26		99	99	
67375522	**Low discrimination**						**67375522**
					GRA	SPO	
Actinomyces israelii		60.81	0.50	NAG 6	99	1	
Mitsuokella multacida		17.84	0.28		1	1	
Clostridium butyricum		15.36	0.19		99	99	
67375530	**Low discrimination**						**67375530**
					GRA	SPO	
Mitsuokella multacida		85.82	0.37	RAF 99	1	1	
Actinomyces israelii		9.72	0.33		99	1	
Clostridium butyricum		3.33	0.04		99	99	
67375540	**Low discrimination**						**67375540**
					GRA	SPO	
Mitsuokella multacida		69.15	0.55	RAF 99	1	1	
Actinomyces israelii		26.60	0.61		99	1	
Clostridium butyricum		4.25	0.26		99	99	
67375541	**Good identification**						**67375541**
					GRA		
Mitsuokella multacida		96.62	0.39	RAF 99	1		
Actinomyces israelii		3.38	0.26	URE 10	99		
67375542	**Low discrimination**						**67375542**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		79.82	0.64		99	1	
Mitsuokella multacida		15.37	0.39		1	1	
Clostridium butyricum		3.31	0.19		99	99	
67375550	**Low discrimination**						**67375550**
					GRA	SPO	
Mitsuokella multacida		84.34	0.47	RAF 99	1	1	
Actinomyces israelii		14.54	0.46		99	1	
Clostridium butyricum		0.82	0.04		99	99	
67375552	**Low discrimination**						**67375552**
					GRA		
Actinomyces israelii		62.59	0.49	MLZ 13	99		
Mitsuokella multacida		26.89	0.30		1		
Bifidobacterium adolescentis		9.60	0.29		99		
67375560	**Low discrimination**						**67375560**
					GRA	SPO	
Mitsuokella multacida		89.32	0.45	RAF 99	1	1	
Actinomyces israelii		8.77	0.40		99	1	
Clostridium butyricum		1.91	0.08		99	99	
67375562	**Low discrimination**						**67375562**
					GRA	SPO	
Actinomyces israelii		55.23	0.43	NAG 6	99	1	
Mitsuokella multacida		41.66	0.28		1	1	
Clostridium butyricum		3.12	0.01		99	99	
67375570	**Good identification**						**67375570**
					GRA		
Mitsuokella multacida		95.78	0.37	RAF 99	1		
Actinomyces israelii		4.22	0.26		99		
67375602	**Low discrimination**						**67375602**
					GRA		
Actinomyces israelii		66.80	0.41	RHA 7	99		
Clostridium clostridioforme		28.22	0.25		50		
Mitsuokella multacida		4.99	0.09		1		
67375612	**Acceptable identification**						**67375612**
					GRA		
Clostridium clostridioforme		91.70	0.36	NIT 1	50		
Actinomyces israelii		7.12	0.27	RAF 94	99		
Mitsuokella multacida		1.19	0.01		1		
67375632	**Good identification**						**67375632**
Clostridium clostridioforme		100.0	0.36	NIT 1			
67375700	**Very good identification**						**67375700**
					GRA	SPO	
Mitsuokella multacida		95.07	0.60	RAF 99	1	1	
Actinomyces israelii		3.81	0.48		99	1	
Clostridium butyricum		1.12	0.18		99	99	
67375701	**Good identification**						**67375701**
					GRA		
Mitsuokella multacida		99.64	0.43	RAF 99	1		
Actinomyces israelii		0.36	0.14	URE 10	99		
67375702	**Low discrimination**						**67375702**
					GRA		
Mitsuokella multacida		56.24	0.43	RAF 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		30.44	0.51	BGL 10	99		
Clostridium clostridioforme		10.76	0.34		50		
67375710	**Good identification**						**67375710**
					GRA		
Mitsuokella multacida		97.77	0.51	RAF 99	1		
Actinomyces israelii		1.76	0.34		99		
Clostridium clostridioforme		0.29	0.11		50		
67375711	**Good identification**						**67375711**
Mitsuokella multacida		100.0	0.35	RAF 99			
67375712	**Low discrimination**						**67375712**
					GRA		
Clostridium clostridioforme		67.51	0.45	NIT 1	50		
Mitsuokella multacida		25.82	0.35	RAF 94	1		
Actinomyces israelii		6.26	0.37		99		
67375720	**Good identification**						**67375720**
					GRA	SPO	
Mitsuokella multacida		98.59	0.49	RAF 99	1	1	
Actinomyces israelii		1.01	0.28		99	1	
Clostridium butyricum		0.40	0.00		99	99	
67375730	**Good identification**						**67375730**
					GRA		
Mitsuokella multacida		98.39	0.41	RAF 99	1		
Clostridium clostridioforme		1.16	0.11		50		
Actinomyces israelii		0.45	0.14		99		
67375732	**Low discrimination**						**67375732**
Clostridium clostridioforme		88.68	0.45	NIT 1			
Mitsuokella multacida		8.48	0.25	RAF 94			
Parabacteroides distasonis		1.17	0.19				
67375740	**Very good identification**						**67375740**
					GRA	SPO	
Mitsuokella multacida		98.37	0.60	RAF 99	1	1	
Actinomyces israelii		1.53	0.41		99	1	
Clostridium butyricum		0.10	0.00		99	99	
67375741	**Good identification**						**67375741**
Mitsuokella multacida		100.0	0.43	RAF 99			
67375742	**Low discrimination**						**67375742**
					GRA	SPO	
Mitsuokella multacida		82.37	0.43	RAF 99	1	1	
Actinomyces israelii		17.34	0.44	BGL 10	99	1	
Clostridium butyricum		0.30	-.07		99	99	
67375750	**Good identification**						**67375750**
					COC	GRA	
Mitsuokella multacida		99.12	0.51	RAF 99	1	1	
Actinomyces israelii		0.69	0.27		1	99	
Blautia producta		0.19	-.03		99	99	
67375751	**Good identification**						**67375751**
Mitsuokella multacida		100.0	0.35	RAF 99			
67375752	**Low discrimination**						**67375752**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Mitsuokella multacida		89.23	0.35	RAF 99	1		
Actinomyces israelii		8.42	0.30	BGL 10	99		
Clostridium clostridioforme		2.36	0.11		50		
67375760	**Good identification**						**67375760**
					GRA		
Mitsuokella multacida		99.60	0.49	RAF 99	1		
Actinomyces israelii		0.40	0.21		99		
67375770	**Good identification**						**67375770**
Mitsuokella multacida		100.0	0.41	RAF 99			
67376000	**Good identification**						**67376000**
Actinomyces israelii		100.0	0.37	GAL 99			
67376002	**Good identification**						**67376002**
Actinomyces israelii		100.0	0.40	GAL 99			
67376100	**Good identification**						**67376100**
					SPO		
Actinomyces israelii		95.25	0.48	GAL 99	1		
Clostridium tertium		4.75	0.11		99		
67376102	**Acceptable identification**						**67376102**
					SPO		
Actinomyces israelii		90.93	0.51	GAL 99	1		
Clostridium tertium		9.07	0.19		99		
67376140	**Good identification**						**67376140**
Actinomyces israelii		100.0	0.41	GAL 99			
67376142	**Good identification**						**67376142**
Actinomyces israelii		100.0	0.44	GAL 99			
67376400	**Good identification**						**67376400**
Actinomyces israelii		100.0	0.36	GAL 99			
67376402	**Good identification**						**67376402**
Actinomyces israelii		100.0	0.39	GAL 99			
67376500	**Low discrimination**						**67376500**
					GRA	SPO	
Actinomyces israelii		69.91	0.46	GAL 99	99	1	
Mitsuokella multacida		29.11	0.27		1	1	
Clostridium butyricum		0.98	-.07		99	99	
67376502	**Good identification**						**67376502**
					GRA		
Actinomyces israelii		96.21	0.49	GAL 99	99		
Mitsuokella multacida		2.97	0.11		1		
Bifidobacterium adolescentis		0.82	0.08		99		
67376512	**Low discrimination**						**67376512**
Actinomyces israelii		61.37	0.35	GAL 99			
Bifidobacterium dentium		30.13	0.06	MLZ 13			
Bifidobacterium adolescentis		4.28	0.09				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67376542	**Acceptable identification**					**67376542**
					GRA	
Actinomyces israelii		92.65	0.42	GAL 99	99	
Mitsuokella multacida		7.35	0.11		1	
67377000	**Very good identification**					**67377000**
					SPO	
Actinomyces israelii		99.80	0.72		1	
Clostridium tertium		0.20	0.11		99	
67377001	**Acceptable identification**					**67377001**
Actinomyces israelii		94.76	0.37	URE 1		
Actinomyces naeslundii		5.24	0.25			
67377002	**Very good identification**					**67377002**
					SPO	
Actinomyces israelii		99.60	0.75		1	
Clostridium tertium		0.40	0.19		99	
67377003	**Good identification**					**67377003**
Actinomyces israelii		96.44	0.40	URE 1		
Actinomyces naeslundii		3.56	0.25			
67377010	**Very good identification**					**67377010**
					SPO	
Actinomyces israelii		97.66	0.57	MLZ 13	1	
Clostridium tertium		2.34	0.15		99	
67377012	**Very good identification**					**67377012**
					SPO	
Actinomyces israelii		95.42	0.60	MLZ 13	1	
Clostridium tertium		4.58	0.24		99	
67377020	**Acceptable identification**					**67377020**
					SPO	
Actinomyces israelii		91.35	0.51	NAG 6	1	
Clostridium tertium		8.66	0.19		99	
67377022	**Low discrimination**					**67377022**
					SPO	
Actinomyces israelii		84.07	0.54	NAG 6	1	
Clostridium tertium		15.93	0.27		99	
67377040	**Very good identification**					**67377040**
Actinomyces israelii		100.0	0.65			
67377042	**Very good identification**					**67377042**
Actinomyces israelii		100.0	0.68			
67377050	**Good identification**					**67377050**
Actinomyces israelii		100.0	0.50	MLZ 13		
67377052	**Good identification**					**67377052**
Actinomyces israelii		100.0	0.53	MLZ 13		
67377060	**Good identification**					**67377060**
Actinomyces israelii		100.0	0.44	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67377062	**Good identification**						**67377062**
Actinomyces israelii		100.0	0.47	NAG 6			
67377100	**Good identification**						**67377100**
					COC	SPO	
Actinomyces israelii		94.13	0.82		1	1	
Clostridium tertium		4.70	0.45		1	99	
Atopobium parvulum		0.71	0.32		99	1	
67377101	**Low discrimination**						**67377101**
					SPO		
Actinomyces israelii		88.72	0.48	URE 1	1		
Clostridium tertium		4.43	0.11		99		
Actinomyces naeslundii		3.49	0.33		1		
67377102	**Good identification**						**67377102**
					SPO		
Actinomyces israelii		90.69	0.85		1		
Clostridium tertium		9.05	0.53		99		
Bifidobacterium adolescentis		0.09	0.28		1		
67377103	**Low discrimination**						**67377103**
					SPO		
Actinomyces israelii		88.81	0.51	URE 1	1		
Clostridium tertium		8.86	0.19		99		
Actinomyces naeslundii		2.33	0.33		1		
67377110	**Low discrimination**						**67377110**
					GRA	SPO	
Actinomyces israelii		62.45	0.68	MLZ 13	99	1	
Clostridium tertium		37.07	0.50		99	99	
Mitsuokella multacida		0.48	0.19		1	1	
67377112	**Low discrimination**						**67377112**
					GRA	SPO	
Clostridium tertium		53.55	0.58	NIT 1	99	99	
Actinomyces israelii		45.10	0.71		99	1	
Leptotrichia buccalis		0.75	0.15		1	1	
67377120	**Low discrimination**						**67377120**
					COC	SPO	
Clostridium tertium		69.07	0.53	NIT 1	1	99	
Actinomyces israelii		29.45	0.61		1	1	
Atopobium parvulum		0.61	0.19		99	1	
67377122	**Low discrimination**						**67377122**
					SPO		
Clostridium tertium		81.94	0.62	NIT 1	99		
Actinomyces israelii		17.47	0.64		1		
Clostridium ramosum		0.47	0.21		92		
67377130	**Very good identification**						**67377130**
					GRA	SPO	
Clostridium tertium		96.38	0.58	NIT 1	99	99	
Actinomyces israelii		3.46	0.47		99	1	
Mitsuokella multacida		0.10	0.09		1	1	
67377132	**Very good identification**						**67377132**
					SPO		
Clostridium tertium		97.99	0.66	NIT 1	99		
Actinomyces israelii		1.76	0.50		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		0.20	0.10		1		
67377140	**Very good identification**						**67377140**
					GRA	SPO	
Actinomyces israelii		98.97	0.75		99	1	
Mitsuokella multacida		0.88	0.27		1	1	
Clostridium tertium		0.13	0.11		99	99	
67377141	**Acceptable identification**						**67377141**
					GRA		
Actinomyces israelii		91.13	0.41	URE 1	99		
Mitsuokella multacida		8.87	0.11		1		
67377142	**Very good identification**						**67377142**
					GRA	SPO	
Actinomyces israelii		99.64	0.78		99	1	
Clostridium tertium		0.26	0.19		99	99	
Mitsuokella multacida		0.07	0.11		1	1	
67377143	**Good identification**						**67377143**
Actinomyces israelii		100.0	0.44	URE 1			
67377150	**Very good identification**						**67377150**
					GRA	SPO	
Actinomyces israelii		96.33	0.61	MLZ 13	99	1	
Mitsuokella multacida		1.90	0.19		1	1	
Clostridium tertium		1.49	0.15		99	99	
67377152	**Very good identification**						**67377152**
					SPO		
Actinomyces israelii		96.62	0.64	MLZ 13	1		
Clostridium tertium		2.98	0.24		99		
Bifidobacterium adolescentis		0.26	0.14		1		
67377160	**Acceptable identification**						**67377160**
					GRA	SPO	
Actinomyces israelii		91.28	0.54	NAG 6	99	1	
Clostridium tertium		5.56	0.19		99	99	
Mitsuokella multacida		3.16	0.17		1	1	
67377162	**Low discrimination**						**67377162**
					SPO		
Actinomyces israelii		89.14	0.57	NAG 6	1		
Clostridium tertium		10.86	0.27		99		
67377170	**Low discrimination**						**67377170**
					GRA	SPO	
Actinomyces israelii		55.49	0.40	MLZ 13	99	1	
Clostridium tertium		40.22	0.24	NAG 6	99	99	
Mitsuokella multacida		4.29	0.09		1	1	
67377200	**Good identification**						**67377200**
Actinomyces israelii		100.0	0.52	RHA 7			
67377202	**Good identification**						**67377202**
Actinomyces israelii		100.0	0.55	RHA 7			
67377210	**Good identification**						**67377210**
Actinomyces israelii		100.0	0.38	RHA 7			
67377212	**Good identification**						**67377212**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.41	RHA 7			
67377240	**Good identification**						**67377240**
Actinomyces israelii		100.0	0.45	RHA 7			
67377242	**Good identification**						**67377242**
Actinomyces israelii		100.0	0.48	RHA 7			
67377300	**Good identification**						**67377300**
					GRA	SPO	
Actinomyces israelii		91.61	0.63	RHA 7	99	1	
Mitsuokella multacida		7.78	0.32		1	1	
Clostridium tertium		0.61	0.11		99	99	
67377302	**Very good identification**						**67377302**
					GRA	SPO	
Actinomyces israelii		98.07	0.66	RHA 7	99	1	
Clostridium tertium		1.31	0.19		99	99	
Mitsuokella multacida		0.62	0.15		1	1	
67377310	**Low discrimination**						**67377310**
					GRA	SPO	
Actinomyces israelii		78.80	0.49	RHA 7	99	1	
Mitsuokella multacida		14.92	0.24	MLZ 13	1	1	
Clostridium tertium		6.28	0.15		99	99	
67377312	**Low discrimination**						**67377312**
					GRA	SPO	
Actinomyces israelii		85.23	0.52	RHA 7	99	1	
Clostridium tertium		13.58	0.24	MLZ 13	99	99	
Mitsuokella multacida		1.20	0.07		1	1	
67377320	**Low discrimination**						**67377320**
					GRA	SPO	
Actinomyces israelii		54.84	0.42	RHA 7	99	1	
Mitsuokella multacida		18.23	0.21	NAG 6	1	1	
Clostridium tertium		17.26	0.19		99	99	
67377322	**Low discrimination**						**67377322**
					SPO		
Actinomyces israelii		47.08	0.45	RHA 7	1		
Clostridium tertium		29.63	0.27	NAG 6	99		
Clostridium ramosum		22.13	0.24		92		
67377340	**Low discrimination**						**67377340**
					GRA		
Actinomyces israelii		82.08	0.56	RHA 7	99		
Mitsuokella multacida		17.92	0.32		1		
67377342	**Very good identification**						**67377342**
					GRA		
Actinomyces israelii		98.41	0.59	RHA 7	99		
Mitsuokella multacida		1.59	0.15		1		
67377350	**Low discrimination**						**67377350**
					COC	GRA	
Actinomyces israelii		65.52	0.42	RHA 7	1	99	
Mitsuokella multacida		31.91	0.24	MLZ 13	1	1	
Blautia producta		2.58	-.03		99	99	
67377352	**Good identification**						**67377352**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		96.52	0.45	RHA 7	99		
Mitsuokella multacida		3.48	0.07	MLZ 13	1		
67377360	**Low discrimination**						**67377360**
					GRA		
Actinomyces israelii		53.91	0.35	RHA 7	99		
Mitsuokella multacida		46.09	0.21	NAG 6	1		
67377362	**Acceptable identification**						**67377362**
					GRA		
Actinomyces israelii		94.04	0.38	RHA 7	99		
Mitsuokella multacida		5.96	0.05	NAG 6	1		
67377400	**Very good identification**						**67377400**
					GRA	SPO	
Actinomyces israelii		98.12	0.70		99	1	
Mitsuokella multacida		1.65	0.27		1	1	
Clostridium butyricum		0.23	0.04		99	99	
67377401	**Low discrimination**						**67377401**
					GRA		
Actinomyces israelii		84.39	0.36	URE 1	99		
Mitsuokella multacida		15.62	0.11		1		
67377402	**Very good identification**						**67377402**
					GRA	SPO	
Actinomyces israelii		99.78	0.73		99	1	
Mitsuokella multacida		0.12	0.11		1	1	
Clostridium butyricum		0.06	-.03		99	99	
67377403	**Good identification**						**67377403**
Actinomyces israelii		100.0	0.39	URE 1			
67377410	**Good identification**						**67377410**
					GRA		
Actinomyces israelii		96.38	0.56	MLZ 13	99		
Mitsuokella multacida		3.62	0.19		1		
67377412	**Very good identification**						**67377412**
Actinomyces israelii		97.52	0.59	MLZ 13			
Bifidobacterium dentium		1.93	0.06				
Bifidobacterium adolescentis		0.28	0.09				
67377420	**Acceptable identification**						**67377420**
					GRA		
Actinomyces israelii		93.82	0.50	NAG 6	99		
Mitsuokella multacida		6.18	0.17		1		
67377422	**Good identification**						**67377422**
Actinomyces israelii		100.0	0.53	NAG 6			
67377430	**Low discrimination**						**67377430**
					GRA		
Actinomyces israelii		81.37	0.36	MLZ 13	99		
Mitsuokella multacida		11.96	0.09	NAG 6	1		
Bifidobacterium dentium		6.67	-.07		99		
67377432	**Low discrimination**						**67377432**
Actinomyces israelii		64.42	0.39	MLZ 13			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		35.58	0.10	NAG 6			
67377440	**Very good identification**						**67377440**
					GRA		
Actinomyces israelii		95.85	0.63		99		
Mitsuokella multacida		4.15	0.27		1		
67377442	**Very good identification**						**67377442**
					GRA		
Actinomyces israelii		99.68	0.66		99		
Mitsuokella multacida		0.32	0.11		1		
67377450	**Acceptable identification**						**67377450**
					COC	GRA	
Actinomyces israelii		90.01	0.49	MLZ 13	1	99	
Mitsuokella multacida		8.69	0.19		1	1	
Blautia producta		1.30	-.03		99	99	
67377452	**Good identification**						**67377452**
					GRA		
Actinomyces israelii		99.29	0.52	MLZ 13	99		
Mitsuokella multacida		0.71	0.03		1		
67377460	**Low discrimination**						**67377460**
					GRA		
Actinomyces israelii		85.51	0.43	NAG 6	99		
Mitsuokella multacida		14.49	0.17		1		
67377462	**Good identification**						**67377462**
Actinomyces israelii		100.0	0.46	NAG 6			
67377500	**Low discrimination**						**67377500**
					GRA	SPO	
Actinomyces israelii		69.88	0.80		99	1	
Mitsuokella multacida		29.09	0.61		1	1	
Clostridium butyricum		0.98	0.28		99	99	
67377501	**Low discrimination**						**67377501**
					GRA	SPO	
Mitsuokella multacida		81.87	0.45	RAF 99	1	1	
Actinomyces israelii		17.88	0.46	URE 10	99	1	
Clostridium butyricum		0.25	-.07		99	99	
67377502	**Very good identification**						**67377502**
					GRA		
Actinomyces israelii		95.70	0.83		99		
Mitsuokella multacida		2.95	0.45		1		
Bifidobacterium adolescentis		0.82	0.42		99		
67377503	**Low discrimination**						**67377503**
					GRA		
Actinomyces israelii		74.20	0.49	URE 1	99		
Mitsuokella multacida		25.17	0.29		1		
Bifidobacterium adolescentis		0.63	0.08		99		
67377510	**Low discrimination**						**67377510**
					GRA		
Actinomyces israelii		49.60	0.66	MLZ 13	99		
Mitsuokella multacida		46.06	0.53		1		
Bifidobacterium dentium		3.61	0.23		99		
67377511	**Acceptable identification**						**67377511**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		91.08	0.37	RAF 99	1		
Actinomyces israelii		8.92	0.32	URE 10	99		
67377512	**Low discrimination**						**67377512**
Actinomyces israelii		60.78	0.69	MLZ 13			
Bifidobacterium dentium		29.84	0.40				
Bifidobacterium adolescentis		4.24	0.43				
67377513	**Low discrimination**						**67377513**
					GRA		
Actinomyces israelii		43.15	0.35	MLZ 13	99		
Mitsuokella multacida		32.66	0.20	URE 1	1		
Bifidobacterium dentium		21.18	0.06		99		
67377520	**Low discrimination**						**67377520**
					GRA	SPO	
Mitsuokella multacida		60.80	0.51	RAF 99	1	1	
Actinomyces israelii		37.29	0.60		99	1	
Clostridium tertium		1.08	0.19		99	99	
67377522	**Low discrimination**						**67377522**
					GRA	SPO	
Actinomyces israelii		82.95	0.63	NAG 6	99	1	
Mitsuokella multacida		10.02	0.35		1	1	
Clostridium tertium		4.80	0.27		99	99	
67377530	**Low discrimination**						**67377530**
					GRA		
Mitsuokella multacida		51.89	0.43	RAF 99	1		
Bifidobacterium dentium		28.93	0.27		99		
Actinomyces israelii		14.27	0.46		99		
67377532	**Low discrimination**						**67377532**
					SPO		
Bifidobacterium dentium		87.41	0.44	NIT 1	1		
Actinomyces israelii		6.39	0.49	RAF 95	1		
Clostridium tertium		4.40	0.32		99		
67377540	**Low discrimination**						**67377540**
					COC	GRA	
Mitsuokella multacida		51.19	0.61	RAF 99	1	1	
Actinomyces israelii		47.81	0.73		1	99	
Blautia producta		0.85	0.24		99	99	
67377541	**Acceptable identification**						**67377541**
					GRA		
Mitsuokella multacida		92.17	0.45	RAF 99	1		
Actinomyces israelii		7.83	0.39	URE 10	99		
67377542	**Good identification**						**67377542**
					GRA		
Actinomyces israelii		92.24	0.76		99		
Mitsuokella multacida		7.31	0.45		1		
Bifidobacterium adolescentis		0.25	0.26		99		
67377543	**Low discrimination**						**67377543**
					GRA		
Actinomyces israelii		53.41	0.42	URE 1	99		
Mitsuokella multacida		46.59	0.29		1		
67377550	**Low discrimination**						**67377550**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		63.75	0.53	RAF 99	1	1	
Actinomyces israelii		26.69	0.59		1	99	
Blautia producta		9.56	0.32		99	99	
67377551	**Good identification**						**67377551**
					COC	GRA	
Mitsuokella multacida		95.08	0.37	RAF 99	1	1	
Actinomyces israelii		3.62	0.25	URE 10	1	99	
Blautia producta		1.30	-.03		99	99	
67377552	**Low discrimination**						**67377552**
					COC	GRA	
Actinomyces israelii		80.75	0.62	MLZ 13	1	99	
Mitsuokella multacida		14.29	0.37		1	1	
Blautia producta		2.14	0.15		99	99	
67377560	**Low discrimination**						**67377560**
					COC	GRA	
Mitsuokella multacida		80.26	0.51	RAF 99	1	1	
Actinomyces israelii		19.14	0.53		1	99	
Blautia producta		0.60	0.07		99	99	
67377562	**Low discrimination**						**67377562**
					GRA		
Actinomyces israelii		76.30	0.56	NAG 6	99		
Mitsuokella multacida		23.70	0.35		1		
67377570	**Low discrimination**						**67377570**
					COC	GRA	
Mitsuokella multacida		84.80	0.43	RAF 99	1	1	
Actinomyces israelii		9.07	0.39		1	99	
Blautia producta		5.65	0.15		99	99	
67377572	**Low discrimination**						**67377572**
					GRA		
Actinomyces israelii		47.75	0.42	MLZ 13	99		
Mitsuokella multacida		33.09	0.26	NAG 6	1		
Bifidobacterium dentium		16.96	0.10		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67377600	**Low discrimination**						**67377600**
					GRA		
Actinomyces israelii		70.67	0.51	RHA 7	99		
Mitsuokella multacida		29.33	0.32		1		
67377602	**Good identification**						**67377602**
					GRA		
Actinomyces israelii		97.02	0.54	RHA 7	99		
Mitsuokella multacida		2.98	0.15		1		
67377610	**Low discrimination**						**67377610**
					GRA		
Actinomyces israelii		51.93	0.37	RHA 7	99		
Mitsuokella multacida		48.08	0.24	MLZ 13	1		
67377612	**Acceptable identification**						**67377612**
					GRA		
Actinomyces israelii		93.58	0.40	RHA 7	99		
Mitsuokella multacida		6.42	0.07	MLZ 13	1		
67377642	**Acceptable identification**						**67377642**
					GRA		
Actinomyces israelii		92.67	0.47	RHA 7	99		
Mitsuokella multacida		7.33	0.15		1		
67377700	**Good identification**						**67377700**
					GRA	SPO	
Mitsuokella multacida		91.08	0.66	RAF 99	1	1	
Actinomyces israelii		8.87	0.61		99	1	
Clostridium butyricum		0.05	0.02		99	99	
67377701	**Good identification**						**67377701**
					GRA		
Mitsuokella multacida		99.12	0.50	RAF 99	1		
Actinomyces israelii		0.88	0.27	URE 10	99		
67377702	**Low discrimination**						**67377702**
					GRA	SPO	
Actinomyces israelii		56.70	0.64	RHA 7	99	1	
Mitsuokella multacida		43.15	0.50		1	1	
Clostridium butyricum		0.09	-.05		99	99	
67377710	**Very good identification**						**67377710**
					COC	GRA	
Mitsuokella multacida		95.74	0.58	RAF 99	1	1	
Actinomyces israelii		4.18	0.47		1	99	
Blautia producta		0.08	-.03		99	99	
67377711	**Good identification**						**67377711**
Mitsuokella multacida		100.0	0.41	RAF 99			
67377712	**Low discrimination**						**67377712**
					GRA		
Mitsuokella multacida		60.80	0.41	RAF 99	1		
Actinomyces israelii		35.82	0.50	BGL 10	99		
Bifidobacterium dentium		2.36	0.06		99		
67377720	**Good identification**						**67377720**
					GRA		
Mitsuokella multacida		97.58	0.56	RAF 99	1		
Actinomyces israelii		2.43	0.41		99		
67377721	**Good identification**						**67377721**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.39	RAF 99			
67377722	**Low discrimination**						**67377722**
					GRA		
Mitsuokella multacida		74.33	0.39	RAF 99	1		
Actinomyces israelii		24.94	0.44	BGL 10	99		
Bacteroides ovatus		0.72	-.02		1		
67377730	**Good identification**						**67377730**
					GRA		
Mitsuokella multacida		98.60	0.47	RAF 99	1		
Actinomyces israelii		1.10	0.27		99		
Bifidobacterium dentium		0.30	-.07		99		
67377740	**Very good identification**						**67377740**
					COC	GRA	
Mitsuokella multacida		95.53	0.66	RAF 99	1	1	
Actinomyces israelii		3.62	0.54		1	99	
Blautia producta		0.86	0.24		99	99	
67377741	**Good identification**						**67377741**
					GRA		
Mitsuokella multacida		99.66	0.50	RAF 99	1		
Actinomyces israelii		0.34	0.20	URE 10	99		
67377742	**Low discrimination**						**67377742**
					COC	GRA	
Mitsuokella multacida		65.79	0.50	RAF 99	1	1	
Actinomyces israelii		33.62	0.57	BGL 10	1	99	
Blautia producta		0.59	0.07		99	99	
67377750	**Good identification**						**67377750**
					COC	GRA	
Mitsuokella multacida		91.10	0.58	RAF 99	1	1	
Blautia producta		7.36	0.32		99	99	
Actinomyces israelii		1.55	0.40		1	99	
67377751	**Good identification**						**67377751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	RAF 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67377752	**Low discrimination**						**67377752**
					COC	GRA	
Mitsuokella multacida		76.34	0.41	RAF 99	1	1	
Actinomyces israelii		17.49	0.43	BGL 10	1	99	
Blautia producta		6.17	0.15		99	99	
67377760	**Good identification**						**67377760**
					COC	GRA	
Mitsuokella multacida		98.65	0.56	RAF 99	1	1	
Actinomyces israelii		0.95	0.34		1	99	
Blautia producta		0.39	0.07		99	99	
67377761	**Good identification**						**67377761**
Mitsuokella multacida		100.0	0.39	RAF 99			
67377762	**Low discrimination**						**67377762**
					GRA		
Mitsuokella multacida		88.46	0.39	RAF 99	1		
Actinomyces israelii		11.54	0.37	BGL 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67377770	**Good identification**						**67377770**
					COC	GRA	
Mitsuokella multacida		96.13	0.47	RAF 99	1	1	
Blautia producta		3.45	0.15		99	99	
Actinomyces israelii		0.42	0.20		1	99	
67410000	**Good identification**						**67410000**
					GRA	CAT	
Fusobacterium mortiferum		96.57	0.52	NIT 1	1	5	
Propionibacterium granulosum		3.43	0.24		99	90	
67410100	**Very good identification**						**67410100**
					CAT		
Fusobacterium mortiferum		99.77	0.58	NIT 1	5		
Capnocytophaga ochracea		0.23	-.01		90		
67411000	**Good identification**						**67411000**
					GRA	CAT	
Fusobacterium mortiferum		96.32	0.55	NIT 1	1	5	
Propionibacterium granulosum		2.28	0.24		99	90	
Actinomyces naeslundii		1.05	0.33		99	25	
67411001	**Low discrimination**						**67411001**
					GRA		
Actinomyces naeslundii		61.73	0.36	ESL 12	99		
Fusobacterium mortiferum		38.27	0.20	MLT 94	1		
67411002	**Low discrimination**						**67411002**
					GRA	CAT	
Fusobacterium mortiferum		45.16	0.38	NIT 1	1	5	
Actinomyces viscosus		34.77	0.24	BGL 10	99	76	
Capnocytophaga ochracea		13.26	0.15		1	90	
67411021	**Good identification**						**67411021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67411023	**Low discrimination**						**67411023**
					GRA		
Actinomyces naeslundii		84.60	0.36	ESL 12	99		
Capnocytophaga ochracea		15.40	-.03	MLT 94	1		
67411100	**Low discrimination**						**67411100**
					GRA	CAT	
Fusobacterium mortiferum		85.30	0.61	NIT 1	1	5	
Capnocytophaga ochracea		13.12	0.33		1	90	
Actinomyces naeslundii		1.13	0.40		99	25	
67411102	**Low discrimination**						**67411102**
					GRA	CAT	
Capnocytophaga ochracea		87.79	0.50	MLT 99	1	90	
Fusobacterium mortiferum		7.05	0.45	NAG 90	1	5	
Actinomyces viscosus		4.13	0.28		99	76	
67411120	**Low discrimination**						**67411120**
					GRA	CAT	
Capnocytophaga ochracea		85.78	0.50	MLT 99	1	90	
Prevotella melaninogenica		8.66	0.29	BGL 90	1	1	
Clostridium perfringens		4.12	0.30		99	1	
67411122	**Very good identification**						**67411122**
					GRA	CAT	
Capnocytophaga ochracea		99.11	0.66	MLT 99	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.25	0.24		99	1	
Prevotella melaninogenica		0.21	0.16		1	1	
67411132	**Good identification**						**67411132**
Capnocytophaga ochracea		99.81	0.50	MLT 99			
Parabacteroides distasonis		0.19	0.16	MLZ 10			
67411162	**Good identification**						**67411162**
					GRA	CAT	
Capnocytophaga ochracea		99.40	0.50	MLT 99	1	90	
Clostridium perfringens		0.43	0.12	SOR 10	99	1	
Actinomyces naeslundii		0.17	0.28		99	25	
67411502	**Good identification**						**67411502**
					GRA	CAT	
Capnocytophaga ochracea		98.47	0.37	MLT 99	1	90	
Bifidobacterium longum subsp. longum		0.81	0.09	NAG 90	99	1	
Lactobacillus fermentum		0.72	0.15		99	1	
67411520	**Good identification**						**67411520**
					GRA	CAT	
Capnocytophaga ochracea		96.26	0.37	MLT 99	1	90	
Bifidobacterium longum subsp. longum		3.18	0.20	BGL 90	99	1	
Prevotella melaninogenica		0.56	-.06		1	1	
67411522	**Good identification**						**67411522**
					GRA	CAT	
Capnocytophaga ochracea		99.50	0.53	MLT 99	1	90	
Bifidobacterium longum subsp. longum		0.37	0.20		99	1	
Parabacteroides distasonis		0.14	0.16		1	76	
67411532	**Acceptable identification**						**67411532**
					GRA	CAT	
Capnocytophaga ochracea		94.08	0.37	MLT 99	1	90	
Bifidobacterium longum subsp. longum		5.29	0.24	MLZ 10	99	1	
Parabacteroides distasonis		0.63	0.12		1	76	
67411562	**Good identification**						**67411562**
Capnocytophaga ochracea		100.0	0.37	MLT 99			
67413000	**Low discrimination**						**67413000**
					GRA	CAT	
Actinomyces naeslundii		42.99	0.38	ESL 12	99	25	
Propionibacterium granulosum		36.42	0.21	MLT 94	99	90	
Fusobacterium mortiferum		20.59	0.20		1	5	
67413001	**Good identification**						**67413001**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67413002	**Low discrimination**						**67413002**
Actinomyces naeslundii		82.52	0.38	ESL 12			
Propionibacterium granulosum		17.48	0.11	MLT 94			
67413003	**Good identification**						**67413003**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67413020	**Low discrimination**						**67413020**
Actinomyces naeslundii		82.52	0.38	ESL 12			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		17.48	0.11	MLT 94			
67413021	**Good identification**						**67413021**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67413022	**Low discrimination**						**67413022**
					GRA		
Actinomyces naeslundii		87.67	0.38	ESL 12	99		
Capnocytophaga ochracea		12.33	-.03	MLT 94	1		
67413023	**Good identification**						**67413023**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67413100	**Low discrimination**						**67413100**
					GRA		
Actinomyces naeslundii		60.92	0.45	ESL 12	99		
Fusobacterium mortiferum		23.93	0.27	MLT 94	1		
Propionibacterium propionicum		11.47	0.23		99		
67413101	**Good identification**						**67413101**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67413102	**Low discrimination**						**67413102**
					GRA	CAT	
Actinomyces naeslundii		54.76	0.45	ESL 12	99	25	
Capnocytophaga ochracea		29.77	0.15	MLT 94	1	90	
Propionibacterium propionicum		15.47	0.26		99	1	
67413103	**Good identification**						**67413103**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67413120	**Low discrimination**						**67413120**
					GRA	CAT	
Actinomyces naeslundii		62.56	0.45	ESL 12	99	25	
Capnocytophaga ochracea		34.01	0.15	MLT 94	1	90	
Prevotella melaninogenica		3.43	-.06		1	1	
67413121	**Good identification**						**67413121**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67413123	**Good identification**						**67413123**
					GRA		
Actinomyces naeslundii		96.81	0.48	ESL 12	99		
Capnocytophaga ochracea		3.19	-.03	MLT 94	1		
67413141	**Good identification**						**67413141**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67413143	**Good identification**						**67413143**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67413161	**Good identification**						**67413161**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67413163	**Good identification**						**67413163**
Actinomyces naeslundii		100.0	0.35	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67414100	**Good identification**						**67414100**
Fusobacterium mortiferum		100.0	0.39	NIT 1			
67415000	**Good identification**						**67415000**
Fusobacterium mortiferum		100.0	0.35	NIT 1			
67415100	**Low discrimination**						**67415100**
					CAT		
Fusobacterium mortiferum		66.19	0.42	NIT 1	5		
Capnocytophaga ochracea		33.81	0.23	XYL 7	90		
67415102	**Acceptable identification**						**67415102**
					CAT		
Capnocytophaga ochracea		94.06	0.39	MLT 99	90		
Bacteroides fragilis		3.67	0.06	NAG 90	99		
Fusobacterium mortiferum		2.27	0.25		5		
67415120	**Low discrimination**						**67415120**
					CAT		
Capnocytophaga ochracea		82.38	0.39	MLT 99	90		
Bacteroides fragilis		16.73	0.18	BGL 90	99		
Bacteroides vulgatus		0.56	-.04		3		
67415122	**Low discrimination**						**67415122**
Capnocytophaga ochracea		69.23	0.56	MLT 99			
Bacteroides fragilis		29.69	0.40				
Parabacteroides distasonis		0.99	0.36				
67415132	**Acceptable identification**						**67415132**
Capnocytophaga ochracea		90.22	0.39	MLT 99			
Parabacteroides distasonis		6.26	0.32	MLZ 10			
Bacteroides fragilis		3.52	0.06				
67415162	**Good identification**						**67415162**
Capnocytophaga ochracea		96.25	0.39	MLT 99			
Bacteroides fragilis		3.75	0.06	SOR 10			
67415322	**Low discrimination**						**67415322**
Parabacteroides distasonis		74.85	0.45	NIT 1			
Capnocytophaga ochracea		17.60	0.21				
Bacteroides fragilis		7.55	0.06				
67415332	**Good identification**						**67415332**
Parabacteroides distasonis		95.37	0.40	NIT 1			
Capnocytophaga ochracea		4.63	0.05				
67415522	**Low discrimination**						**67415522**
					GRA	CAT	
Capnocytophaga ochracea		89.42	0.43	MLT 99	1	90	
Parabacteroides distasonis		4.45	0.33		1	76	
Bifidobacterium longum subsp. longum		3.94	0.28		99	1	
67415722	**Acceptable identification**						**67415722**
					CAT		
Parabacteroides distasonis		90.73	0.41	NIT 1	76		
Capnocytophaga ochracea		6.14	0.09		90		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		3.13	-.03		3		
67415732	**Good identification**						**67415732**
Parabacteroides distasonis		100.0	0.36	NIT 1			
67430000	**Low discrimination**						**67430000**
					GRA	CAT	
Fusobacterium mortiferum		73.71	0.43	NIT 1	1	5	
Propionibacterium granulosum		26.29	0.32		99	90	
67430100	**Good identification**						**67430100**
Fusobacterium mortiferum		100.0	0.49	NIT 1			
67431000	**Low discrimination**						**67431000**
					GRA	CAT	
Fusobacterium mortiferum		76.43	0.46	NIT 1	1	5	
Propionibacterium granulosum		18.17	0.32		99	90	
Actinomyces naeslundii		5.40	0.38		99	25	
67431001	**Acceptable identification**						**67431001**
					GRA		
Actinomyces naeslundii		91.29	0.41	ESL 12	99		
Fusobacterium mortiferum		8.71	0.12	MLT 94	1		
67431003	**Good identification**						**67431003**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67431021	**Good identification**						**67431021**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67431022	**Low discrimination**						**67431022**
					CAT		
Actinomyces naeslundii		44.89	0.38	ESL 12	25		
Clostridium perfringens		39.34	0.14	MLT 94	1		
Propionibacterium granulosum		9.45	0.11		90		
67431023	**Good identification**						**67431023**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67431100	**Acceptable identification**						**67431100**
					GRA		
Fusobacterium mortiferum		90.57	0.52	NIT 1	1		
Actinomyces naeslundii		7.80	0.45		99		
Clostridium perfringens		1.16	0.09		99		
67431101	**Low discrimination**						**67431101**
					GRA	SPO	
Actinomyces naeslundii		74.52	0.48	ESL 12	99	1	
Clostridium chauvoei		19.66	0.15	MLT 94	99	99	
Fusobacterium mortiferum		5.83	0.18		1	1	
67431102	**Low discrimination**						**67431102**
					GRA	CAT	
Fusobacterium mortiferum		41.70	0.36	NIT 1	1	5	
Actinomyces naeslundii		32.31	0.45	BGL 10	99	25	
Capnocytophaga ochracea		17.57	0.15		1	90	
67431103	**Acceptable identification**						**67431103**
					SPO		
Actinomyces naeslundii		91.83	0.48	ESL 12	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium chauvoei		4.28	0.03	MLT 94	99	
Actinomyces viscosus		3.89	0.01		1	
67431121	**Low discrimination**					**67431121**
					SPO	
Actinomyces naeslundii		89.27	0.48	ESL 12	1	
Clostridium chauvoei		7.85	0.07	MLT 94	99	
Clostridium perfringens		2.88	0.01		84	
67431123	**Good identification**					**67431123**
					GRA	
Actinomyces naeslundii		96.81	0.48	ESL 12	99	
Capnocytophaga ochracea		3.19	-.03	MLT 94	1	
67431141	**Good identification**					**67431141**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67431143	**Good identification**					**67431143**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67431161	**Good identification**					**67431161**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67431163	**Good identification**					**67431163**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67431322	**Good identification**					**67431322**
Parabacteroides distasonis		98.16	0.40	NIT 1		
Capnocytophaga ochracea		1.84	-.03	XYL 90		
67431722	**Good identification**					**67431722**
Parabacteroides distasonis		100.0	0.36	NIT 1		
67433001	**Good identification**					**67433001**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67433002	**Low discrimination**					**67433002**
Actinomyces naeslundii		75.34	0.42	ESL 12		
Propionibacterium granulosum		24.66	0.19	MLT 94		
67433003	**Good identification**					**67433003**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67433020	**Low discrimination**					**67433020**
Actinomyces naeslundii		75.34	0.42	ESL 12		
Propionibacterium granulosum		24.66	0.19	MLT 94		
67433021	**Good identification**					**67433021**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67433022	**Acceptable identification**					**67433022**
Actinomyces naeslundii		92.44	0.42	ESL 12		
Propionibacterium granulosum		7.56	0.09	MLT 94		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67433023	**Good identification**					**67433023**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67433100	**Low discrimination**					**67433100**
					GRA	
Actinomyces naeslundii		87.74	0.50	ESL 12	99	
Propionibacterium propionicum		6.96	0.21	MLT 94	99	
Fusobacterium mortiferum		5.30	0.18		1	
67433101	**Good identification**					**67433101**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67433102	**Low discrimination**					**67433102**
Actinomyces naeslundii		89.36	0.50	ESL 12		
Propionibacterium propionicum		10.64	0.24	MLT 94		
67433103	**Good identification**					**67433103**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67433120	**Good identification**					**67433120**
Actinomyces naeslundii		97.57	0.50	ESL 12		
Clostridium perfringens		2.44	0.01	MLT 94		
67433121	**Good identification**					**67433121**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67433122	**Good identification**					**67433122**
					GRA	
Actinomyces naeslundii		97.52	0.50	ESL 12	99	
Capnocytophaga ochracea		2.48	-.03	MLT 94	1	
67433123	**Good identification**					**67433123**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67433140	**Low discrimination**					**67433140**
Actinomyces naeslundii		68.98	0.37	ESL 12		
Propionibacterium propionicum		31.02	0.21	MLT 94		
67433141	**Good identification**					**67433141**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67433142	**Low discrimination**					**67433142**
Actinomyces naeslundii		59.72	0.37	ESL 12		
Propionibacterium propionicum		40.28	0.24	MLT 94		
67433143	**Good identification**					**67433143**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67433160	**Good identification**					**67433160**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67433161	**Good identification**					**67433161**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.40	ESL 12			
67433162	**Good identification**						**67433162**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67433163	**Good identification**						**67433163**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67435122	**Low discrimination**						**67435122**
Parabacteroides distasonis		81.89	0.48	NIT 1			
Capnocytophaga ochracea		12.68	0.21				
Bacteroides fragilis		5.44	0.06				
67435132	**Good identification**						**67435132**
Parabacteroides distasonis		96.90	0.43	NIT 1			
Capnocytophaga ochracea		3.10	0.05				
67435322	**Good identification**						**67435322**
Parabacteroides distasonis		100.0	0.56	NIT 1			
67435332	**Good identification**						**67435332**
Parabacteroides distasonis		100.0	0.51	NIT 1			
67435522	**Acceptable identification**						**67435522**
					GRA	CAT	
Parabacteroides distasonis		91.47	0.44	NIT 1	1	76	
Bifidobacterium longum subsp. longum		4.45	0.17		99	1	
Capnocytophaga ochracea		4.08	0.09		1	90	
67435532	**Low discrimination**						**67435532**
					GRA		
Parabacteroides distasonis		86.67	0.39	NIT 1	1		
Bifidobacterium longum subsp. longum		13.33	0.21		99		
67435722	**Good identification**						**67435722**
Parabacteroides distasonis		98.39	0.52	NIT 1			
Bacteroides thetaiotaomicron		1.61	0.06				
67435732	**Good identification**						**67435732**
Parabacteroides distasonis		97.84	0.48	NIT 1			
Bacteroides thetaiotaomicron		1.16	-.01				
Clostridium clostridioforme		1.00	0.08				
67450000	**Good identification**						**67450000**
Fusobacterium mortiferum		100.0	0.43	NIT 1			
67450100	**Good identification**						**67450100**
Fusobacterium mortiferum		100.0	0.50	NIT 1			
67451000	**Good identification**						**67451000**
					GRA		
Fusobacterium mortiferum		95.86	0.46	NIT 1	1		
Actinomyces naeslundii		4.14	0.35		99		
67451001	**Low discrimination**						**67451001**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		86.51	0.38	ESL 12	99		
Fusobacterium mortiferum		13.49	0.12	MLT 94	1		
67451003	**Low discrimination**						**67451003**
Actinomyces naeslundii		74.02	0.38	ESL 12			
Actinomyces viscosus		25.98	0.06	MLT 94			
67451021	**Good identification**						**67451021**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67451023	**Good identification**						**67451023**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67451100	**Low discrimination**						**67451100**
					GRA	CAT	
Fusobacterium mortiferum		85.61	0.53	NIT 1	1	5	
Capnocytophaga ochracea		9.88	0.23		1	90	
Actinomyces naeslundii		4.51	0.42		99	25	
67451101	**Low discrimination**						**67451101**
					GRA		
Actinomyces naeslundii		88.67	0.46	ESL 12	99		
Fusobacterium mortiferum		11.33	0.19	MLT 94	1		
67451102	**Low discrimination**						**67451102**
					GRA	CAT	
Capnocytophaga ochracea		82.03	0.39	MLT 99	1	90	
Fusobacterium mortiferum		8.78	0.36	NAG 90	1	5	
Actinomyces naeslundii		4.16	0.42		99	25	
67451103	**Low discrimination**						**67451103**
					GRA		
Actinomyces naeslundii		73.97	0.46	ESL 12	99		
Actinomyces viscosus		16.22	0.10	MLT 94	99		
Capnocytophaga ochracea		9.81	0.05		1		
67451120	**Low discrimination**						**67451120**
					GRA	CAT	
Capnocytophaga ochracea		88.93	0.39	MLT 99	1	90	
Actinomyces naeslundii		4.51	0.42	BGL 90	99	25	
Prevotella melaninogenica		3.99	0.12		1	1	
67451121	**Low discrimination**						**67451121**
					GRA		
Actinomyces naeslundii		88.29	0.46	ESL 12	99		
Capnocytophaga ochracea		11.71	0.05	MLT 94	1		
67451122	**Good identification**						**67451122**
					GRA	CAT	
Capnocytophaga ochracea		96.77	0.56	MLT 99	1	90	
Prevotella oralis		2.29	0.27		1	1	
Actinomyces naeslundii		0.55	0.42		99	25	
67451132	**Good identification**						**67451132**
Capnocytophaga ochracea		99.59	0.39	MLT 99			
Parabacteroides distasonis		0.41	0.11	MLZ 10			
67451162	**Good identification**						**67451162**
					GRA		
Capnocytophaga ochracea		99.11	0.39	MLT 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.89	0.30	SOR 10	99		
67451522	**Good identification**						**67451522**
Capnocytophaga ochracea		99.70	0.43	MLT 99			
Parabacteroides distasonis		0.30	0.12				
67453000	**Low discrimination**						**67453000**
					GRA		
Actinomyces naeslundii		89.25	0.40	ESL 12	99		
Fusobacterium mortiferum		10.75	0.12	MLT 94	1		
67453001	**Good identification**						**67453001**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67453002	**Good identification**						**67453002**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67453003	**Good identification**						**67453003**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67453020	**Good identification**						**67453020**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67453021	**Good identification**						**67453021**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67453022	**Good identification**						**67453022**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67453023	**Good identification**						**67453023**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67453100	**Acceptable identification**						**67453100**
					GRA		
Actinomyces naeslundii		91.01	0.47	ESL 12	99		
Fusobacterium mortiferum		8.99	0.19	MLT 94	1		
67453101	**Good identification**						**67453101**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67453102	**Low discrimination**						**67453102**
					GRA	CAT	
Actinomyces naeslundii		87.46	0.47	ESL 12	99	25	
Capnocytophaga ochracea		8.97	0.05	MLT 94	1	90	
Actinomyces israelii		3.58	0.14		99	1	
67453103	**Good identification**						**67453103**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67453120	**Acceptable identification**						**67453120**
					GRA		
Actinomyces naeslundii		90.70	0.47	ESL 12	99		
Capnocytophaga ochracea		9.30	0.05	MLT 94	1		
67453121	**Good identification**						**67453121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	ESL 12		
67453122	**Low discrimination**					**67453122**
					GRA	
Actinomyces naeslundii		52.01	0.47	ESL 12	99	
Capnocytophaga ochracea		47.99	0.21	MLT 94	1	
67453123	**Good identification**					**67453123**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67453141	**Good identification**					**67453141**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67453143	**Good identification**					**67453143**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67453161	**Good identification**					**67453161**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67453163	**Good identification**					**67453163**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67455122	**Low discrimination**					**67455122**
Capnocytophaga ochracea		56.26	0.45	MLT 99		
Bacteroides fragilis		41.35	0.34			
Parabacteroides distasonis		1.74	0.32			
67455322	**Low discrimination**					**67455322**
Parabacteroides distasonis		80.42	0.40	NIT 1		
Capnocytophaga ochracea		8.78	0.11			
Bacteroides fragilis		6.45	-.01			
67455332	**Good identification**					**67455332**
Parabacteroides distasonis		100.0	0.35	NIT 1		
67455722	**Low discrimination**					**67455722**
					CAT	
Parabacteroides distasonis		88.19	0.36	NIT 1	76	
Bacteroides uniformis		9.04	0.01		5	
Capnocytophaga ochracea		2.77	-.02		90	
67470100	**Good identification**					**67470100**
					GRA	
Fusobacterium mortiferum		97.91	0.41	NIT 1	1	
Actinomyces naeslundii		2.10	0.24		99	
67471000	**Low discrimination**					**67471000**
					GRA	
Fusobacterium mortiferum		78.08	0.38	NIT 1	1	
Actinomyces naeslundii		21.92	0.40		99	
67471001	**Good identification**					**67471001**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67471002	**Low discrimination**					**67471002**
					GRA	
Actinomyces naeslundii		71.64	0.40	ESL 12	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		28.36	0.21	MLT 94	1		
67471003	**Good identification**						**67471003**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67471020	**Good identification**						**67471020**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67471021	**Good identification**						**67471021**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67471022	**Good identification**						**67471022**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67471023	**Good identification**						**67471023**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67471100	**Low discrimination**						**67471100**
					GRA		
Fusobacterium mortiferum		74.49	0.44	NIT 1	1		
Actinomyces naeslundii		25.51	0.47		99		
67471101	**Good identification**						**67471101**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67471102	**Low discrimination**						**67471102**
					GRA	CAT	
Actinomyces naeslundii		68.25	0.47	ESL 12	99	25	
Fusobacterium mortiferum		22.15	0.27	MLT 94	1	5	
Capnocytophaga ochracea		7.00	0.05		1	90	
67471103	**Good identification**						**67471103**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67471120	**Low discrimination**						**67471120**
					GRA	CAT	
Actinomyces naeslundii		79.79	0.47	ESL 12	99	25	
Clostridium perfringens		12.03	0.11	MLT 94	99	1	
Capnocytophaga ochracea		8.18	0.05		1	90	
67471121	**Good identification**						**67471121**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67471122	**Low discrimination**						**67471122**
					GRA		
Actinomyces naeslundii		40.20	0.47	ESL 12	99		
Capnocytophaga ochracea		37.09	0.21	MLT 94	1		
Parabacteroides distasonis		14.33	0.27		1		
67471123	**Good identification**						**67471123**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67471141	**Good identification**						**67471141**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67471143	**Good identification**						**67471143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	ESL 12		
67471161	**Good identification**					**67471161**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67471163	**Good identification**					**67471163**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67473000	**Good identification**					**67473000**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67473001	**Good identification**					**67473001**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67473002	**Acceptable identification**					**67473002**
Actinomyces naeslundii		94.66	0.45	ESL 12		
Actinomyces israelii		5.34	0.14	MLT 94		
67473003	**Good identification**					**67473003**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67473020	**Good identification**					**67473020**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67473021	**Good identification**					**67473021**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67473022	**Good identification**					**67473022**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67473023	**Good identification**					**67473023**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67473100	**Acceptable identification**					**67473100**
Actinomyces naeslundii		94.98	0.52	ESL 12		
Actinomyces israelii		5.02	0.21	MLT 94		
67473101	**Good identification**					**67473101**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67473102	**Acceptable identification**					**67473102**
Actinomyces naeslundii		92.65	0.52	ESL 12		
Actinomyces israelii		7.35	0.24	MLT 94		
67473103	**Good identification**					**67473103**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67473120	**Good identification**					**67473120**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67473121	**Good identification**					**67473121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.55	ESL 12		
67473122	**Good identification**					**67473122**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67473123	**Good identification**					**67473123**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67473140	**Low discrimination**					**67473140**
Actinomyces naeslundii		89.56	0.39	ESL 12		
Actinomyces israelii		10.44	0.14	MLT 94		
67473141	**Good identification**					**67473141**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67473142	**Low discrimination**					**67473142**
Actinomyces naeslundii		85.12	0.39	ESL 12		
Actinomyces israelii		14.88	0.17	MLT 94		
67473143	**Good identification**					**67473143**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67473160	**Good identification**					**67473160**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67473161	**Good identification**					**67473161**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67473162	**Good identification**					**67473162**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67473163	**Good identification**					**67473163**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67475122	**Low discrimination**					**67475122**
Parabacteroides distasonis		88.91	0.43	NIT 1		
Capnocytophaga ochracea		6.39	0.11			
Bacteroides fragilis		4.70	-.01			
67475132	**Good identification**					**67475132**
Parabacteroides distasonis		100.0	0.39	NIT 1		
67475322	**Good identification**					**67475322**
Parabacteroides distasonis		100.0	0.51	NIT 1		
67475332	**Good identification**					**67475332**
Parabacteroides distasonis		100.0	0.47	NIT 1		
67475522	**Good identification**					**67475522**
Parabacteroides distasonis		97.97	0.39	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		2.03	-.02				
67475722	**Acceptable identification**						**67475722**
Parabacteroides distasonis		93.26	0.48	NIT 1			
Bacteroides thetaiotaomicron		6.04	0.11				
Clostridium clostridioforme		0.70	0.06				
67475732	**Acceptable identification**						**67475732**
Parabacteroides distasonis		90.05	0.43	NIT 1			
Clostridium clostridioforme		5.74	0.17				
Bacteroides thetaiotaomicron		4.21	0.04				
67477100	**Good identification**						**67477100**
Actinomyces israelii		100.0	0.39	MLT 99			
67477102	**Good identification**						**67477102**
Actinomyces israelii		100.0	0.42	MLT 99			
67477142	**Good identification**						**67477142**
Actinomyces israelii		100.0	0.35	MLT 99			
67477502	**Low discrimination**						**67477502**
					GRA		
Actinomyces israelii		87.17	0.41	MLT 99	99		
Mitsuokella multacida		8.22	0.11	SAL 92	1		
Bifidobacterium adolescentis		4.61	0.13		99		
67500002	**Good identification**						**67500002**
Actinomyces viscosus		100.0	0.40	ESL 1			
67500102	**Good identification**						**67500102**
					GRA		
Actinomyces viscosus		97.54	0.44	ESL 1	99		
Capnocytophaga ochracea		2.46	0.15	FRU 95	1		
67500103	**Good identification**						**67500103**
					SPO		
Actinomyces viscosus		96.39	0.36	ESL 1	1		
Clostridium chauvoei		3.62	0.13	FRU 95	99		
67501002	**Acceptable identification**						**67501002**
					GRA		
Actinomyces viscosus		91.47	0.36	ESL 1	99		
Capnocytophaga ochracea		6.69	0.15	FRU 95	1		
Actinomyces naeslundii		1.84	0.31		99		
67501101	**Very good identification**						**67501101**
					GRA	SPO	
Clostridium chauvoei		99.54	0.60	RAF 1	99	99	
Actinomyces naeslundii		0.42	0.42		99	1	
Capnocytophaga ochracea		0.04	-.01		1	1	
67501102	**Low discrimination**						**67501102**
					GRA		
Capnocytophaga ochracea		79.29	0.50	FRU 99	1		
Actinomyces viscosus		19.45	0.40	NAG 90	99		
Actinomyces naeslundii		0.63	0.39		99		
67501103	**Low discrimination**						**67501103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Clostridium chauvoei		82.69	0.47	RAF 1	99		
Actinomyces viscosus		13.65	0.32		1		
Actinomyces naeslundii		1.98	0.42		1		
67501120	**Low discrimination**						**67501120**
					GRA	CAT	
Prevotella melaninogenica		51.02	0.47	NIT 1	1	1	
Capnocytophaga ochracea		45.97	0.50	FRU 90	1	90	
Clostridium perfringens		2.21	0.30		99	1	
67501121	**Good identification**						**67501121**
					GRA	SPO	
Clostridium chauvoei		96.58	0.51	RAF 1	99	99	
Actinomyces naeslundii		1.22	0.42		99	1	
Prevotella melaninogenica		1.16	0.12		1	1	
67501122	**Very good identification**						**67501122**
					GRA	CAT	
Capnocytophaga ochracea		97.06	0.66	FRU 99	1	90	
Prevotella melaninogenica		2.28	0.34		1	1	
Clostridium perfringens		0.24	0.24		99	1	
67501123	**Low discrimination**						**67501123**
					GRA	SPO	
Clostridium chauvoei		61.16	0.39	RAF 1	99	99	
Capnocytophaga ochracea		33.67	0.32		1	1	
Actinomyces naeslundii		4.38	0.42		99	1	
67501132	**Good identification**						**67501132**
					GRA	CAT	
Capnocytophaga ochracea		99.68	0.50	FRU 99	1	90	
Prevotella melaninogenica		0.21	0.00	MLZ 10	1	1	
Bifidobacterium longum subsp. longum		0.11	0.07		99	1	
67501162	**Good identification**						**67501162**
					GRA	CAT	
Capnocytophaga ochracea		99.22	0.50	FRU 99	1	90	
Clostridium perfringens		0.43	0.12	SOR 10	99	1	
Prevotella melaninogenica		0.21	0.00		1	1	
67501502	**Acceptable identification**						**67501502**
					GRA	CAT	
Capnocytophaga ochracea		90.50	0.37	FRU 99	1	90	
Bifidobacterium longum subsp. longum		8.23	0.27	NAG 90	99	1	
Actinomyces viscosus		1.27	0.06		99	76	
67501520	**Low discrimination**						**67501520**
					GRA	CAT	
Capnocytophaga ochracea		70.07	0.37	FRU 99	1	90	
Bifidobacterium longum subsp. longum		25.48	0.37	BGL 90	99	1	
Prevotella melaninogenica		4.45	0.12		1	1	
67501522	**Good identification**						**67501522**
					GRA	CAT	
Capnocytophaga ochracea		95.99	0.53	FRU 99	1	90	
Bifidobacterium longum subsp. longum		3.88	0.37		99	1	
Prevotella melaninogenica		0.13	0.00		1	1	
67501530	**Low discrimination**						**67501530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		84.78	0.41	NIT 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		15.22	0.20	FRU 90	1	90	
67501532	**Low discrimination**						**67501532**
					GRA	CAT	
Capnocytophaga ochracea		61.77	0.37	FRU 99	1	90	
Bifidobacterium longum subsp. longum		38.23	0.41	MLZ 10	99	1	
67501562	**Good identification**						**67501562**
Capnocytophaga ochracea		100.0	0.37	FRU 99			
67503000	**Good identification**						**67503000**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67503001	**Good identification**						**67503001**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67503002	**Low discrimination**						**67503002**
Actinomyces naeslundii		79.43	0.36	ESL 12			
Actinomyces viscosus		20.57	0.02	FRU 95			
67503003	**Good identification**						**67503003**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67503020	**Good identification**						**67503020**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67503021	**Good identification**						**67503021**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67503022	**Low discrimination**						**67503022**
					GRA		
Actinomyces naeslundii		85.43	0.36	ESL 12	99		
Capnocytophaga ochracea		14.57	-.03	FRU 95	1		
67503023	**Good identification**						**67503023**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67503100	**Low discrimination**						**67503100**
					GRA	CAT	
Actinomyces naeslundii		76.83	0.44	ESL 12	99	25	
Propionibacterium propionicum		17.54	0.23	FRU 95	99	1	
Capnocytophaga ochracea		5.63	-.01		1	90	
67503102	**Low discrimination**						**67503102**
					GRA	CAT	
Actinomyces naeslundii		46.22	0.44	ESL 12	99	25	
Capnocytophaga ochracea		30.47	0.15	FRU 95	1	90	
Propionibacterium propionicum		15.83	0.26		99	1	
67503103	**Low discrimination**						**67503103**
					SPO		
Actinomyces naeslundii		82.12	0.47	ESL 12	1		
Clostridium chauvoei		17.88	0.13	FRU 95	99		
67503120	**Low discrimination**						**67503120**
					GRA	CAT	
Actinomyces naeslundii		41.83	0.44	ESL 12	99	25	
Prevotella melaninogenica		30.60	0.12	FRU 95	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		27.57	0.15		1	90	
67503121	**Low discrimination**						**67503121**
					SPO		
Actinomyces naeslundii		70.86	0.47	ESL 12	1		
Clostridium chauvoei		29.15	0.17	FRU 95	99		
67503123	**Low discrimination**						**67503123**
					GRA	SPO	
Actinomyces naeslundii		89.88	0.47	ESL 12	99	1	
Clostridium chauvoei		6.52	0.04	FRU 95	99	99	
Capnocytophaga ochracea		3.59	-.03		1	1	
67505102	**Good identification**						**67505102**
					GRA		
Capnocytophaga ochracea		95.34	0.39	FRU 99	1		
Bacteroides fragilis		3.72	0.06	NAG 90	1		
Actinomyces viscosus		0.95	0.06		99		
67505120	**Low discrimination**						**67505120**
					CAT		
Capnocytophaga ochracea		79.21	0.39	FRU 99	90		
Bacteroides fragilis		16.09	0.18	BGL 90	99		
Prevotella melaninogenica		3.55	0.12		1		
67505122	**Low discrimination**						**67505122**
					CAT		
Capnocytophaga ochracea		69.81	0.56	FRU 99	90		
Bacteroides fragilis		29.93	0.40		99		
Prevotella melaninogenica		0.07	0.00		1		
67505132	**Good identification**						**67505132**
					GRA	CAT	
Capnocytophaga ochracea		95.04	0.39	FRU 99	1	90	
Bacteroides fragilis		3.71	0.06	MLZ 10	1	99	
Bifidobacterium longum subsp. longum		1.26	0.15		99	1	
67505162	**Good identification**						**67505162**
Capnocytophaga ochracea		96.25	0.39	FRU 99			
Bacteroides fragilis		3.75	0.06	SOR 10			
67505500	**Acceptable identification**						**67505500**
					GRA	CAT	
Bifidobacterium longum subsp. longum		90.76	0.35	NIT 1	99	1	
Capnocytophaga ochracea		9.25	0.10	FRU 90	1	90	
67505502	**Low discrimination**						**67505502**
					GRA	CAT	
Bifidobacterium longum subsp. longum		51.26	0.35	NIT 1	99	1	
Capnocytophaga ochracea		47.00	0.26	FRU 90	1	90	
Bifidobacterium adolescentis		1.73	0.12		99	1	
67505510	**Good identification**						**67505510**
Bifidobacterium longum subsp. longum		100.0	0.39	NIT 1			
67505512	**Acceptable identification**						**67505512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.24	0.39	NIT 1	99	1	
Capnocytophaga ochracea		5.52	0.10	FRU 90	1	90	
Bifidobacterium adolescentis		2.24	0.13		99	1	
67505520	**Low discrimination**						**67505520**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.29	0.46	NIT 1	99	1	
Capnocytophaga ochracea		17.03	0.26	FRU 90	1	90	
Bacteroides vulgatus		8.68	0.15		1	3	
67505522	**Low discrimination**						**67505522**
					GRA	CAT	
Capnocytophaga ochracea		66.26	0.43	FRU 99	1	90	
Bifidobacterium longum subsp. longum		32.12	0.46		99	1	
Bacteroides fragilis		1.63	0.06		1	99	
67505530	**Good identification**						**67505530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.53	0.50	NIT 1	99	1	
Capnocytophaga ochracea		1.47	0.10	FRU 90	1	90	
67505532	**Low discrimination**						**67505532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		88.14	0.50	NIT 1	99	1	
Capnocytophaga ochracea		11.87	0.26	FRU 90	1	90	
67510002	**Very good identification**						**67510002**
					GRA	CAT	
Actinomyces viscosus		99.61	0.62	ESL 1	99	76	
Capnocytophaga ochracea		0.24	0.15		1	90	
Fusobacterium mortiferum		0.07	0.20		1	5	
67510003	**Good identification**						**67510003**
Actinomyces viscosus		99.77	0.53	ESL 1			
Actinomyces naeslundii		0.23	0.32				
67510102	**Low discrimination**						**67510102**
					GRA	CAT	
Actinomyces viscosus		88.19	0.66	ESL 1	99	76	
Capnocytophaga ochracea		11.59	0.50		1	90	
Fusobacterium mortiferum		0.08	0.26		1	5	
67510103	**Very good identification**						**67510103**
					GRA	SPO	
Actinomyces viscosus		98.79	0.58	ESL 1	99	1	
Clostridium chauvoei		0.46	0.19		99	99	
Capnocytophaga ochracea		0.39	0.15		1	1	
67510120	**Low discrimination**						**67510120**
					GRA	CAT	
Capnocytophaga ochracea		86.57	0.50	GAL 99	1	90	
Prevotella melaninogenica		8.73	0.29	BGL 90	1	1	
Clostridium perfringens		4.16	0.30		99	1	
67510122	**Very good identification**						**67510122**
					GRA	CAT	
Capnocytophaga ochracea		98.35	0.66	GAL 99	1	90	
Actinomyces viscosus		0.84	0.32		99	76	
Clostridium perfringens		0.25	0.24		99	1	
67510132	**Good identification**						**67510132**
Capnocytophaga ochracea		100.0	0.50	GAL 99			
67510162	**Good identification**						**67510162**
					GRA	CAT	
Capnocytophaga ochracea		99.46	0.50	GAL 99	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.43	0.12	SOR 10	99	1	
Actinomyces naeslundii		0.11	0.24		99	25	
67510502	**Low discrimination**						**67510502**
					GRA	CAT	
Capnocytophaga ochracea		69.27	0.37	GAL 99	1	90	
Actinomyces viscosus		30.16	0.32	NAG 90	99	76	
Bifidobacterium longum subsp. longum		0.57	0.09		99	1	
67510520	**Good identification**						**67510520**
					GRA	CAT	
Capnocytophaga ochracea		96.26	0.37	GAL 99	1	90	
Bifidobacterium longum subsp. longum		3.18	0.20	BGL 90	99	1	
Prevotella melaninogenica		0.56	-.06		1	1	
67510522	**Good identification**						**67510522**
					GRA	CAT	
Capnocytophaga ochracea		99.63	0.53	GAL 99	1	90	
Bifidobacterium longum subsp. longum		0.37	0.20		99	1	
67510532	**Acceptable identification**						**67510532**
					GRA	CAT	
Capnocytophaga ochracea		94.67	0.37	GAL 99	1	90	
Bifidobacterium longum subsp. longum		5.33	0.24	MLZ 10	99	1	
67510562	**Good identification**						**67510562**
Capnocytophaga ochracea		100.0	0.37	GAL 99			
67511001	**Low discrimination**						**67511001**
					SPO		
Actinomyces naeslundii		50.94	0.56	ESL 12	1		
Clostridium chauvoei		42.65	0.32		99		
Actinomyces viscosus		5.69	0.16		1		
67511002	**Low discrimination**						**67511002**
					GRA		
Actinomyces viscosus		70.91	0.58	ESL 1	99		
Capnocytophaga ochracea		27.04	0.50		1		
Actinomyces naeslundii		1.43	0.53		99		
67511003	**Low discrimination**						**67511003**
					SPO		
Actinomyces viscosus		89.66	0.50	ESL 1	1		
Actinomyces naeslundii		8.11	0.56		1		
Clostridium chauvoei		1.20	0.19		99		
67511020	**Low discrimination**						**67511020**
					GRA	CAT	
Capnocytophaga ochracea		55.35	0.50	LAC 99	1	90	
Clostridium perfringens		35.87	0.49	BGL 90	99	1	
Prevotella melaninogenica		5.59	0.29		1	1	
67511021	**Low discrimination**						**67511021**
					GRA	SPO	
Actinomyces naeslundii		66.55	0.56	ESL 12	99	1	
Clostridium chauvoei		18.57	0.24		99	99	
Capnocytophaga ochracea		8.51	0.15		1	1	
67511022	**Good identification**						**67511022**
					GRA	CAT	
Capnocytophaga ochracea		94.64	0.66	LAC 99	1	90	
Clostridium perfringens		3.21	0.44		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		1.00	0.32		1	1	
67511032	**Good identification**						**67511032**
					GRA	CAT	
Capnocytophaga ochracea		99.32	0.50	LAC 99	1	90	
Clostridium perfringens		0.31	0.10	MLZ 10	99	1	
Actinomyces naeslundii		0.28	0.31		99	25	
67511041	**Good identification**						**67511041**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67511043	**Low discrimination**						**67511043**
					GRA		
Actinomyces naeslundii		58.36	0.43	ESL 12	99		
Actinomyces viscosus		36.94	0.16		99		
Capnocytophaga ochracea		4.70	-.01		1		
67511060	**Low discrimination**						**67511060**
					GRA	CAT	
Clostridium perfringens		50.41	0.37	LAC 88	99	1	
Capnocytophaga ochracea		45.38	0.33		1	90	
Actinomyces naeslundii		3.80	0.40		99	25	
67511061	**Low discrimination**						**67511061**
					GRA	CAT	
Actinomyces naeslundii		85.48	0.43	ESL 12	99	25	
Clostridium perfringens		7.64	0.03		99	1	
Capnocytophaga ochracea		6.88	-.01		1	90	
67511062	**Acceptable identification**						**67511062**
					GRA	CAT	
Capnocytophaga ochracea		93.60	0.50	LAC 99	1	90	
Clostridium perfringens		5.44	0.32	SOR 10	99	1	
Actinomyces naeslundii		0.87	0.40		99	25	
67511063	**Low discrimination**						**67511063**
					GRA		
Actinomyces naeslundii		58.00	0.43	ESL 12	99		
Capnocytophaga ochracea		42.00	0.15		1		
67511100	**Good identification**						**67511100**
					GRA	CAT	
Capnocytophaga ochracea		94.62	0.67	NAG 90	1	90	
Actinomyces naeslundii		1.29	0.61	BGL 90	99	25	
Clostridium perfringens		1.26	0.38		99	1	
67511101	**Good identification**						**67511101**
					GRA	SPO	
Clostridium chauvoei		94.86	0.66	RAF 1	99	99	
Actinomyces naeslundii		3.26	0.64		99	1	
Capnocytophaga ochracea		1.61	0.33		1	1	
67511102	**Very good identification**						**67511102**
					GRA		
Capnocytophaga ochracea		95.09	0.84	NAG 90	1		
Actinomyces viscosus		4.48	0.62		99		
Actinomyces naeslundii		0.14	0.61		99		
67511103	**Low discrimination**						**67511103**
					GRA	SPO	
Actinomyces viscosus		39.48	0.54	ESL 1	99	1	
Clostridium chauvoei		29.37	0.53		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		25.41	0.50		1	1	
67511110	**Good identification**						**67511110**
					GRA	CAT	
Capnocytophaga ochracea		98.73	0.51	MLZ 10	1	90	
Actinomyces naeslundii		0.64	0.39	NAG 90	99	25	
Bifidobacterium longum subsp. longum		0.20	0.13	BGL 90	99	1	
67511112	**Very good identification**						**67511112**
					GRA		
Capnocytophaga ochracea		99.43	0.67	MLZ 10	1		
Actinomyces viscosus		0.43	0.28	NAG 90	99		
Actinomyces naeslundii		0.07	0.39		99		
67511120	**Low discrimination**						**67511120**
					GRA	CAT	
Capnocytophaga ochracea		86.79	0.84	BGL 90	1	90	
Prevotella melaninogenica		8.76	0.63		1	1	
Clostridium perfringens		4.17	0.64		99	1	
67511121	**Low discrimination**						**67511121**
					GRA	SPO	
Clostridium chauvoei		61.38	0.58	RAF 1	99	99	
Capnocytophaga ochracea		28.11	0.50		1	1	
Actinomyces naeslundii		6.32	0.64		99	1	
67511122	**Excellent identification**						**67511122**
					GRA	CAT	
Capnocytophaga ochracea		99.23	1.00		1	90	
Clostridium perfringens		0.25	0.59		99	1	
Prevotella melaninogenica		0.21	0.51		1	1	
67511123	**Good identification**						**67511123**
					GRA	SPO	
Capnocytophaga ochracea		92.84	0.66	URE 1	1	1	
Clostridium chauvoei		3.98	0.45		99	99	
Actinomyces naeslundii		2.32	0.64		99	1	
67511130	**Very good identification**						**67511130**
					GRA	CAT	
Capnocytophaga ochracea		98.51	0.67	MLZ 10	1	90	
Prevotella melaninogenica		0.90	0.29	BGL 90	1	1	
Clostridium perfringens		0.43	0.30		99	1	
67511132	**Very good identification**						**67511132**
					GRA	CAT	
Capnocytophaga ochracea		99.91	0.84	MLZ 10	1	90	
Clostridium perfringens		0.02	0.24		99	1	
Prevotella melaninogenica		0.02	0.16		1	1	
67511133	**Good identification**						**67511133**
					GRA	SPO	
Capnocytophaga ochracea		98.45	0.50	MLZ 10	1	1	
Actinomyces naeslundii		1.17	0.42	URE 1	99	1	
Clostridium chauvoei		0.38	0.11		99	99	
67511140	**Good identification**						**67511140**
					GRA	CAT	
Capnocytophaga ochracea		95.33	0.51	NAG 90	1	90	
Clostridium perfringens		2.18	0.26	SOR 10	99	1	
Actinomyces naeslundii		2.06	0.48	BGL 90	99	25	
67511142	**Very good identification**						**67511142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Capnocytophaga ochracea		99.16	0.67	NAG 90	1		
Actinomyces viscosus		0.42	0.28	SOR 10	99		
Actinomyces naeslundii		0.24	0.48		99		
67511152	**Good identification**						**67511152**
					GRA		
Capnocytophaga ochracea		99.89	0.51	MLZ 10	1		
Actinomyces naeslundii		0.11	0.26	NAG 90	99		
67511160	**Good identification**						**67511160**
					GRA	CAT	
Capnocytophaga ochracea		91.42	0.67	SOR 10	1	90	
Clostridium perfringens		7.52	0.52	BGL 90	99	1	
Prevotella melaninogenica		0.84	0.29		1	1	
67511162	**Very good identification**						**67511162**
					GRA	CAT	
Capnocytophaga ochracea		99.50	0.84	SOR 10	1	90	
Clostridium perfringens		0.43	0.46		99	1	
Actinomyces naeslundii		0.03	0.48		99	25	
67511163	**Good identification**						**67511163**
					GRA	CAT	
Capnocytophaga ochracea		95.43	0.50	SOR 10	1	90	
Actinomyces naeslundii		3.79	0.51	URE 1	99	25	
Clostridium perfringens		0.41	0.12		99	1	
67511170	**Good identification**						**67511170**
					GRA	CAT	
Capnocytophaga ochracea		99.06	0.51	MLZ 10	1	90	
Clostridium perfringens		0.74	0.18	SOR 10	99	1	
Actinomyces naeslundii		0.11	0.26	BGL 90	99	25	
67511172	**Very good identification**						**67511172**
					GRA	CAT	
Capnocytophaga ochracea		99.95	0.67	MLZ 10	1	90	
Clostridium perfringens		0.04	0.12	SOR 10	99	1	
Actinomyces naeslundii		0.01	0.26		99	25	
67511302	**Acceptable identification**						**67511302**
					GRA	CAT	
Capnocytophaga ochracea		92.34	0.50	RHA 1	1	90	
Actinomyces viscosus		4.35	0.28	NAG 90	99	76	
Prevotella oralis		3.09	0.24		1	1	
67511320	**Low discrimination**						**67511320**
					GRA	CAT	
Capnocytophaga ochracea		82.33	0.50	RHA 1	1	90	
Prevotella melaninogenica		8.31	0.29	BGL 90	1	1	
Clostridium perfringens		3.95	0.30		99	1	
67511322	**Good identification**						**67511322**
					CAT		
Capnocytophaga ochracea		92.51	0.66	RHA 1	90		
Prevotella oralis		6.54	0.46		1		
Parabacteroides distasonis		0.39	0.38		76		
67511332	**Good identification**						**67511332**
					CAT		
Capnocytophaga ochracea		97.38	0.50	RHA 1	90		
Parabacteroides distasonis		2.00	0.33	MLZ 10	76		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		0.63	0.12		1		
67511362	**Good identification**						**67511362**
					GRA	CAT	
Capnocytophaga ochracea		98.94	0.50	RHA 1	1	90	
Prevotella oralis		0.64	0.12	SOR 10	1	1	
Clostridium perfringens		0.43	0.12		99	1	
67511402	**Low discrimination**						**67511402**
					GRA	CAT	
Capnocytophaga ochracea		85.35	0.37	LAC 99	1	90	
Actinomyces viscosus		12.81	0.24	NAG 90	99	76	
Lactobacillus fermentum		1.14	0.20		99	1	
67511420	**Acceptable identification**						**67511420**
					GRA	CAT	
Capnocytophaga ochracea		92.94	0.37	LAC 99	1	90	
Clostridium perfringens		3.45	0.15	BGL 90	99	1	
Bifidobacterium longum subsp. longum		3.07	0.20		99	1	
67511422	**Good identification**						**67511422**
					GRA	CAT	
Capnocytophaga ochracea		99.38	0.53	LAC 99	1	90	
Bifidobacterium longum subsp. longum		0.37	0.20		99	1	
Clostridium perfringens		0.19	0.10		99	1	
67511432	**Acceptable identification**						**67511432**
					GRA	CAT	
Capnocytophaga ochracea		94.67	0.37	LAC 99	1	90	
Bifidobacterium longum subsp. longum		5.33	0.24	MLZ 10	99	1	
67511462	**Good identification**						**67511462**
Capnocytophaga ochracea		100.0	0.37	LAC 99			
67511500	**Acceptable identification**						**67511500**
					GRA	CAT	
Capnocytophaga ochracea		92.69	0.54	NAG 90	1	90	
Bifidobacterium longum subsp. longum		6.89	0.43	BGL 90	99	1	
Lactobacillus fermentum		0.18	0.24		99	1	
67511502	**Very good identification**						**67511502**
					GRA	CAT	
Capnocytophaga ochracea		98.74	0.71	NAG 90	1	90	
Bifidobacterium longum subsp. longum		0.82	0.43		99	1	
Actinomyces viscosus		0.27	0.28		99	76	
67511503	**Low discrimination**						**67511503**
					GRA	SPO	
Capnocytophaga ochracea		85.02	0.37	NAG 90	1	1	
Actinomyces viscosus		7.56	0.20	URE 1	99	1	
Clostridium chauvoei		5.63	0.19		99	99	
67511510	**Low discrimination**						**67511510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		53.26	0.47	NIT 1	99	1	
Capnocytophaga ochracea		46.74	0.38		1	90	
67511512	**Low discrimination**						**67511512**
					GRA	CAT	
Capnocytophaga ochracea		87.58	0.54	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		11.09	0.47	NAG 90	99	1	
Bifidobacterium adolescentis		1.33	0.33		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
67511520	**Very good identification**							**67511520**
					GRA	CAT		
Capnocytophaga ochracea		95.76	0.71	BGL 90	1	90		
Bifidobacterium longum subsp. longum		3.17	0.54		99	1		
Prevotella melaninogenica		0.55	0.29		1	1		
67511521	**Low discrimination**							**67511521**
					GRA	SPO	CAT	
Capnocytophaga ochracea		84.98	0.37	URE 1	1	1	90	
Clostridium chauvoei		10.62	0.24	BGL 90	99	99	1	
Bifidobacterium longum subsp. longum		2.81	0.20		99	1	1	
67511522	**Very good identification**							**67511522**
					GRA	CAT		
Capnocytophaga ochracea		99.59	0.87		1	90		
Bifidobacterium longum subsp. longum		0.37	0.54		99	1		
Clostridium perfringens		0.01	0.24		99	1		
67511523	**Good identification**							**67511523**
					GRA	SPO	CAT	
Capnocytophaga ochracea		99.25	0.53	URE 1	1	1	90	
Bifidobacterium longum subsp. longum		0.36	0.20		99	1	1	
Clostridium chauvoei		0.24	0.11		99	99	1	
67511530	**Low discrimination**							**67511530**
					GRA	CAT		
Capnocytophaga ochracea		66.36	0.54	MLZ 10	1	90		
Bifidobacterium longum subsp. longum		33.61	0.58	BGL 90	99	1		
Prevotella melaninogenica		0.04	-.06		1	1		
67511532	**Good identification**							**67511532**
					GRA	CAT		
Capnocytophaga ochracea		94.65	0.71	MLZ 10	1	90		
Bifidobacterium longum subsp. longum		5.33	0.58		99	1		
Parabacteroides distasonis		0.02	0.21		1	76		
67511533	**Acceptable identification**							**67511533**
					GRA	CAT		
Capnocytophaga ochracea		94.67	0.37	MLZ 10	1	90		
Bifidobacterium longum subsp. longum		5.33	0.24	URE 1	99	1		
67511540	**Good identification**							**67511540**
					GRA	CAT		
Capnocytophaga ochracea		99.33	0.38	NAG 90	1	90		
Bifidobacterium longum subsp. longum		0.67	0.09	SOR 10	99	1		
67511542	**Good identification**							**67511542**
					GRA	CAT		
Capnocytophaga ochracea		99.77	0.54	NAG 90	1	90		
Bifidobacterium adolescentis		0.15	0.16	SOR 10	99	1		
Bifidobacterium longum subsp. longum		0.08	0.09		99	1		
67511552	**Good identification**							**67511552**
					GRA	CAT		
Capnocytophaga ochracea		97.24	0.38	MLZ 10	1	90		
Bifidobacterium adolescentis		1.64	0.18	NAG 90	99	1		
Bifidobacterium longum subsp. longum		1.12	0.13	SOR 10	99	1		
67511560	**Good identification**							**67511560**
					GRA	CAT		
Capnocytophaga ochracea		99.18	0.54	SOR 10	1	90		
Clostridium perfringens		0.47	0.18	BGL 90	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		0.30	0.20		99	1	
67511562	**Very good identification**						**67511562**
					GRA	CAT	
Capnocytophaga ochracea		99.94	0.71	SOR 10	1	90	
Bifidobacterium longum subsp. longum		0.03	0.20		99	1	
Clostridium perfringens		0.03	0.12		99	1	
67511563	**Good identification**						**67511563**
Capnocytophaga ochracea		100.0	0.37	SOR 10			
67511570	**Good identification**						**67511570**
					GRA	CAT	
Capnocytophaga ochracea		95.60	0.38	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		4.40	0.24	SOR 10	99	1	
67511572	**Good identification**						**67511572**
					GRA	CAT	
Capnocytophaga ochracea		99.49	0.54	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		0.51	0.24	SOR 10	99	1	
67511702	**Good identification**						**67511702**
					GRA	CAT	
Capnocytophaga ochracea		99.18	0.37	RHA 1	1	90	
Bifidobacterium longum subsp. longum		0.82	0.09	NAG 90	99	1	
67511720	**Low discrimination**						**67511720**
					GRA	CAT	
Bacteroides vulgatus		69.05	0.37	NIT 1	1	3	
Capnocytophaga ochracea		29.80	0.37	XYL 98	1	90	
Bifidobacterium longum subsp. longum		0.99	0.20		99	1	
67511722	**Good identification**						**67511722**
					CAT		
Capnocytophaga ochracea		97.56	0.53	RHA 1	90		
Parabacteroides distasonis		1.43	0.34		76		
Prevotella oralis		0.40	0.12		1		
67511732	**Low discrimination**						**67511732**
					GRA	CAT	
Capnocytophaga ochracea		87.82	0.37	RHA 1	1	90	
Parabacteroides distasonis		6.25	0.29	MLZ 10	1	76	
Bifidobacterium longum subsp. longum		4.94	0.24		99	1	
67511762	**Good identification**						**67511762**
Capnocytophaga ochracea		100.0	0.37	RHA 1			
67512001	**Good identification**						**67512001**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67512021	**Good identification**						**67512021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67512023	**Good identification**						**67512023**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67512100	**Very good identification**						**67512100**
					GRA	CAT	
Propionibacterium propionicum		96.32	0.57	ESL 1	99	1	
Actinomyces naeslundii		3.37	0.42		99	25	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		0.31	-.01		1	90	
67512101	**Low discrimination**						**67512101**
Propionibacterium propionicum		67.90	0.41	ESL 1			
Actinomyces naeslundii		32.10	0.45	URE 10			
67512102	**Low discrimination**						**67512102**
Propionibacterium propionicum		83.99	0.60	ESL 1			
Actinomyces viscosus		12.42	0.32				
Actinomyces naeslundii		1.96	0.42				
67512103	**Low discrimination**						**67512103**
Propionibacterium propionicum		56.86	0.44	ESL 1			
Actinomyces viscosus		25.22	0.24	URE 10			
Actinomyces naeslundii		17.92	0.45				
67512120	**Low discrimination**						**67512120**
					GRA	CAT	
Actinomyces naeslundii		45.35	0.42	ESL 12	99	25	
Capnocytophaga ochracea		37.76	0.15	GAL 96	1	90	
Propionibacterium propionicum		13.08	0.23		99	1	
67512121	**Good identification**						**67512121**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67512123	**Good identification**						**67512123**
					GRA		
Actinomyces naeslundii		95.20	0.45	ESL 12	99		
Capnocytophaga ochracea		4.81	-.03	GAL 96	1		
67512140	**Very good identification**						**67512140**
Propionibacterium propionicum		99.39	0.57	ESL 1			
Actinomyces naeslundii		0.61	0.29				
67512141	**Acceptable identification**						**67512141**
Propionibacterium propionicum		92.30	0.41	ESL 1			
Actinomyces naeslundii		7.70	0.32	URE 10			
67512142	**Very good identification**						**67512142**
					GRA	CAT	
Propionibacterium propionicum		99.38	0.60	ESL 1	99	1	
Actinomyces naeslundii		0.41	0.29		99	25	
Capnocytophaga ochracea		0.22	-.01		1	90	
67512143	**Acceptable identification**						**67512143**
Propionibacterium propionicum		94.73	0.44	ESL 1			
Actinomyces naeslundii		5.27	0.32	URE 10			
67513000	**Good identification**						**67513000**
					CAT		
Actinomyces naeslundii		92.45	0.58	ESL 12	25		
Propionibacterium propionicum		3.16	0.23		1		
Propionibacterium granulosum		2.25	0.16		90		
67513001	**Very good identification**						**67513001**
Actinomyces naeslundii		100.0	0.61	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67513002	**Low discrimination**						**67513002**
					GRA		
Actinomyces naeslundii		70.06	0.58	ESL 12	99		
Actinomyces viscosus		18.14	0.24		99		
Capnocytophaga ochracea		6.92	0.15		1		
67513003	**Good identification**						**67513003**
Actinomyces naeslundii		94.56	0.61	ESL 12			
Actinomyces viscosus		5.44	0.16				
67513010	**Good identification**						**67513010**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67513011	**Good identification**						**67513011**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67513012	**Low discrimination**						**67513012**
					GRA		
Actinomyces naeslundii		82.75	0.36	ESL 12	99		
Capnocytophaga ochracea		17.25	-.01	MLZ 5	1		
67513013	**Good identification**						**67513013**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67513020	**Low discrimination**						**67513020**
					GRA	CAT	
Actinomyces naeslundii		85.27	0.58	ESL 12	99	25	
Capnocytophaga ochracea		8.42	0.15		1	90	
Clostridium perfringens		5.46	0.15		99	1	
67513021	**Very good identification**						**67513021**
Actinomyces naeslundii		100.0	0.61	ESL 12			
67513022	**Low discrimination**						**67513022**
					GRA	CAT	
Actinomyces naeslundii		51.86	0.58	ESL 12	99	25	
Capnocytophaga ochracea		46.09	0.32		1	90	
Clostridium perfringens		1.56	0.10		99	1	
67513023	**Very good identification**						**67513023**
					GRA		
Actinomyces naeslundii		99.41	0.61	ESL 12	99		
Capnocytophaga ochracea		0.60	-.03		1		
67513030	**Low discrimination**						**67513030**
					GRA		
Actinomyces naeslundii		82.75	0.36	ESL 12	99		
Capnocytophaga ochracea		17.25	-.01	MLZ 5	1		
67513031	**Good identification**						**67513031**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67513033	**Good identification**						**67513033**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67513040	**Low discrimination**						**67513040**
Actinomyces naeslundii		83.76	0.45	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		16.24	0.23				
67513041	**Good identification**						**67513041**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67513042	**Low discrimination**						**67513042**
					GRA	CAT	
Actinomyces naeslundii		73.92	0.45	ESL 12	99	25	
Propionibacterium propionicum		21.49	0.26		99	1	
Capnocytophaga ochracea		4.60	-.01		1	90	
67513043	**Good identification**						**67513043**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67513060	**Low discrimination**						**67513060**
					GRA	CAT	
Actinomyces naeslundii		88.40	0.45	ESL 12	99	25	
Clostridium perfringens		6.11	0.03		99	1	
Capnocytophaga ochracea		5.50	-.01		1	90	
67513061	**Good identification**						**67513061**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67513062	**Low discrimination**						**67513062**
					GRA		
Actinomyces naeslundii		64.12	0.45	ESL 12	99		
Capnocytophaga ochracea		35.88	0.15		1		
67513063	**Good identification**						**67513063**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67513100	**Low discrimination**						**67513100**
					GRA	CAT	
Propionibacterium propionicum		45.76	0.57	ESL 1	99	1	
Actinomyces naeslundii		38.46	0.66		99	25	
Capnocytophaga ochracea		14.68	0.33		1	90	
67513101	**Low discrimination**						**67513101**
					SPO		
Actinomyces naeslundii		80.50	0.69	ESL 12	1		
Clostridium chauvoei		12.20	0.32		99		
Propionibacterium propionicum		7.10	0.41		1		
67513102	**Low discrimination**						**67513102**
					GRA	CAT	
Capnocytophaga ochracea		53.50	0.50	MAN 1	1	90	
Propionibacterium propionicum		27.80	0.60	NAG 90	99	1	
Actinomyces naeslundii		15.58	0.66		99	25	
67513103	**Low discrimination**						**67513103**
Actinomyces naeslundii		82.10	0.69	ESL 12			
Propionibacterium propionicum		10.85	0.44				
Actinomyces viscosus		2.95	0.20				
67513110	**Low discrimination**						**67513110**
					GRA	CAT	
Actinomyces naeslundii		49.16	0.44	ESL 12	99	25	
Capnocytophaga ochracea		39.61	0.17	MLZ 5	1	90	
Propionibacterium propionicum		11.23	0.23		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67513111	**Good identification**						**67513111**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67513113	**Good identification**						**67513113**
					GRA		
Actinomyces naeslundii		95.35	0.47	ESL 12	99		
Capnocytophaga ochracea		4.66	-.01	MLZ 5	1		
67513120	**Low discrimination**						**67513120**
					GRA	CAT	
Capnocytophaga ochracea		69.28	0.50	MAN 1	1	90	
Actinomyces naeslundii		20.17	0.66	BGL 90	99	25	
Prevotella melaninogenica		6.99	0.29		1	1	
67513121	**Good identification**						**67513121**
					GRA	SPO	
Actinomyces naeslundii		92.94	0.69	ESL 12	99	1	
Clostridium chauvoei		4.70	0.24		99	99	
Capnocytophaga ochracea		2.15	0.15		1	1	
67513122	**Very good identification**						**67513122**
					GRA	CAT	
Capnocytophaga ochracea		96.11	0.66	MAN 1	1	90	
Actinomyces naeslundii		3.11	0.66		99	25	
Clostridium perfringens		0.24	0.24		99	1	
67513123	**Low discrimination**						**67513123**
					GRA	SPO	
Actinomyces naeslundii		82.16	0.69	ESL 12	99	1	
Capnocytophaga ochracea		17.11	0.32		1	1	
Clostridium chauvoei		0.73	0.11		99	99	
67513131	**Good identification**						**67513131**
					GRA		
Actinomyces naeslundii		95.35	0.47	ESL 12	99		
Capnocytophaga ochracea		4.66	-.01	MLZ 5	1		
67513132	**Good identification**						**67513132**
					GRA		
Capnocytophaga ochracea		98.49	0.50	MAN 1	1		
Actinomyces naeslundii		1.51	0.44	MLZ 10	99		
67513133	**Low discrimination**						**67513133**
					GRA		
Actinomyces naeslundii		69.47	0.47	ESL 12	99		
Capnocytophaga ochracea		30.53	0.15	MLZ 5	1		
67513140	**Low discrimination**						**67513140**
					GRA	CAT	
Propionibacterium propionicum		84.06	0.57	ESL 1	99	1	
Actinomyces naeslundii		12.47	0.53		99	25	
Capnocytophaga ochracea		3.00	0.17		1	90	
67513141	**Low discrimination**						**67513141**
Actinomyces naeslundii		66.69	0.56	ESL 12			
Propionibacterium propionicum		33.31	0.41				
67513142	**Low discrimination**						**67513142**
					GRA	CAT	
Propionibacterium propionicum		75.85	0.60	ESL 1	99	1	
Capnocytophaga ochracea		16.22	0.33		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		7.50	0.53		99	25	
67513143	**Low discrimination**						**67513143**
					GRA	CAT	
Actinomyces naeslundii		56.70	0.56	ESL 12	99	25	
Propionibacterium propionicum		42.48	0.44		99	1	
Capnocytophaga ochracea		0.83	-.01		1	90	
67513161	**Good identification**						**67513161**
					GRA		
Actinomyces naeslundii		98.57	0.56	ESL 12	99		
Capnocytophaga ochracea		1.44	-.01		1		
67513162	**Acceptable identification**						**67513162**
					GRA	CAT	
Capnocytophaga ochracea		94.26	0.50	MAN 1	1	90	
Actinomyces naeslundii		4.84	0.53	SOR 10	99	25	
Propionibacterium propionicum		0.50	0.26		99	1	
67513163	**Low discrimination**						**67513163**
					GRA		
Actinomyces naeslundii		88.41	0.56	ESL 12	99		
Capnocytophaga ochracea		11.59	0.15		1		
67513502	**Low discrimination**						**67513502**
					GRA	CAT	
Capnocytophaga ochracea		83.26	0.37	MAN 1	1	90	
Bifidobacterium adolescentis		9.29	0.30	NAG 90	99	1	
Actinomyces israelii		2.90	0.28		99	1	
67513520	**Acceptable identification**						**67513520**
					GRA	CAT	
Capnocytophaga ochracea		94.74	0.37	MAN 1	1	90	
Bifidobacterium longum subsp. longum		3.13	0.20	BGL 90	99	1	
Actinomyces naeslundii		1.58	0.32		99	25	
67513522	**Good identification**						**67513522**
					GRA	CAT	
Capnocytophaga ochracea		99.45	0.53	MAN 1	1	90	
Bifidobacterium longum subsp. longum		0.37	0.20		99	1	
Actinomyces naeslundii		0.18	0.32		99	25	
67513532	**Acceptable identification**						**67513532**
					GRA	CAT	
Capnocytophaga ochracea		94.67	0.37	MAN 1	1	90	
Bifidobacterium longum subsp. longum		5.33	0.24	MLZ 10	99	1	
67513562	**Good identification**						**67513562**
Capnocytophaga ochracea		100.0	0.37	MAN 1			
67514102	**Low discrimination**						**67514102**
					GRA		
Capnocytophaga ochracea		74.28	0.39	GAL 99	1		
Actinomyces viscosus		22.83	0.32	NAG 90	99		
Bacteroides fragilis		2.90	0.06		1		
67514120	**Low discrimination**						**67514120**
					CAT		
Capnocytophaga ochracea		82.38	0.39	GAL 99	90		
Bacteroides fragilis		16.73	0.18	BGL 90	99		
Bacteroides vulgatus		0.56	-.04		3		
67514122	**Low discrimination**						**67514122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Capnocytophaga ochracea		69.90	0.56	GAL 99	90		
Bacteroides fragilis		29.97	0.40		99		
Prevotella oralis		0.06	0.03		1		
67514132	**Good identification**						**67514132**
Capnocytophaga ochracea		96.25	0.39	GAL 99			
Bacteroides fragilis		3.75	0.06	MLZ 10			
67514162	**Good identification**						**67514162**
Capnocytophaga ochracea		96.25	0.39	GAL 99			
Bacteroides fragilis		3.75	0.06	SOR 10			
67514522	**Acceptable identification**						**67514522**
					GRA	CAT	
Capnocytophaga ochracea		93.58	0.43	GAL 99	1	90	
Bifidobacterium longum subsp. longum		4.12	0.28		99	1	
Bacteroides fragilis		2.30	0.06		1	99	
67515002	**Low discrimination**						**67515002**
					GRA		
Capnocytophaga ochracea		85.79	0.39	LAC 99	1		
Actinomyces viscosus		9.09	0.24	NAG 90	99		
Bacteroides fragilis		3.34	0.06		1		
67515020	**Low discrimination**						**67515020**
					GRA	CAT	
Capnocytophaga ochracea		80.64	0.39	LAC 99	1	90	
Bacteroides fragilis		16.38	0.18	BGL 90	1	99	
Clostridium perfringens		2.11	0.15		99	1	
67515022	**Low discrimination**						**67515022**
					CAT		
Capnocytophaga ochracea		69.62	0.56	LAC 99	90		
Bacteroides fragilis		29.85	0.40		99		
Prevotella oralis		0.36	0.16		1		
67515032	**Good identification**						**67515032**
Capnocytophaga ochracea		96.25	0.39	LAC 99			
Bacteroides fragilis		3.75	0.06	MLZ 10			
67515062	**Good identification**						**67515062**
Capnocytophaga ochracea		96.25	0.39	LAC 99			
Bacteroides fragilis		3.75	0.06	SOR 10			
67515100	**Very good identification**						**67515100**
					GRA	CAT	
Capnocytophaga ochracea		96.90	0.57	NAG 90	1	90	
Bacteroides fragilis		1.79	0.18	BGL 90	1	99	
Actinomyces israelii		0.36	0.32		99	1	
67515102	**Very good identification**						**67515102**
					GRA	CAT	
Capnocytophaga ochracea		95.60	0.73	NAG 90	1	90	
Bacteroides fragilis		3.73	0.40		1	99	
Bifidobacterium longum subsp. infantis		0.27	0.26		99	1	
67515103	**Low discrimination**						**67515103**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		86.39	0.39	NAG 90	1	1	
Actinomyces viscosus		5.42	0.20	URE 1	99	1	
Clostridium chauvoei		4.04	0.19		99	99	
67515110	**Good identification**						**67515110**
					GRA	CAT	
Capnocytophaga ochracea		96.83	0.41	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		2.36	0.21	NAG 90	99	1	
Actinomyces israelii		0.48	0.18	BGL 90	99	1	
67515112	**Very good identification**						**67515112**
					GRA	CAT	
Capnocytophaga ochracea		98.25	0.57	MLZ 10	1	90	
Bifidobacterium adolescentis		1.05	0.33	NAG 90	99	1	
Bacteroides fragilis		0.35	0.06		1	99	
67515120	**Low discrimination**						**67515120**
					CAT		
Capnocytophaga ochracea		82.13	0.73	BGL 90	90		
Bacteroides fragilis		16.68	0.52		99		
Bacteroides vulgatus		0.56	0.30		3		
67515121	**Low discrimination**						**67515121**
					GRA	SPO	
Capnocytophaga ochracea		76.26	0.39	URE 1	1	1	
Bacteroides fragilis		15.49	0.18	BGL 90	1	1	
Clostridium chauvoei		6.73	0.24		99	99	
67515122	**Low discrimination**						**67515122**
					CAT		
Capnocytophaga ochracea		69.88	0.90		90		
Bacteroides fragilis		29.97	0.74		99		
Prevotella oralis		0.06	0.37		1		
67515123	**Low discrimination**						**67515123**
					GRA	SPO	
Capnocytophaga ochracea		69.76	0.56	URE 1	1	1	
Bacteroides fragilis		29.91	0.40		1	1	
Clostridium chauvoei		0.12	0.11		99	99	
67515130	**Very good identification**						**67515130**
					GRA	CAT	
Capnocytophaga ochracea		97.06	0.57	MLZ 10	1	90	
Bacteroides fragilis		1.79	0.18	BGL 90	1	99	
Bifidobacterium longum subsp. longum		1.05	0.32		99	1	
67515132	**Very good identification**						**67515132**
Capnocytophaga ochracea		95.90	0.73	MLZ 10			
Bacteroides fragilis		3.74	0.40				
Parabacteroides distasonis		0.24	0.41				
67515133	**Good identification**						**67515133**
Capnocytophaga ochracea		96.25	0.39	MLZ 10			
Bacteroides fragilis		3.75	0.06	URE 1			
67515140	**Good identification**						**67515140**
					GRA	CAT	
Capnocytophaga ochracea		98.72	0.41	NAG 90	1	90	
Actinomyces israelii		1.28	0.25	SOR 10	99	1	
67515142	**Very good identification**						**67515142**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		99.33	0.57	NAG 90	1	90	
Bacteroides fragilis		0.35	0.06	SOR 10	1	99	
Actinomyces israelii		0.21	0.28		99	1	
67515152	**Good identification**						**67515152**
					GRA	CAT	
Capnocytophaga ochracea		98.54	0.41	MLZ 10	1	90	
Bifidobacterium adolescentis		1.17	0.18	NAG 90	99	1	
Actinomyces israelii		0.29	0.14	SOR 10	99	1	
67515160	**Very good identification**						**67515160**
					GRA	CAT	
Capnocytophaga ochracea		97.77	0.57	SOR 10	1	90	
Bacteroides fragilis		1.81	0.18	BGL 90	1	99	
Clostridium perfringens		0.33	0.18		99	1	
67515162	**Very good identification**						**67515162**
					GRA	CAT	
Capnocytophaga ochracea		96.22	0.73	SOR 10	1	90	
Bacteroides fragilis		3.75	0.40		1	99	
Clostridium perfringens		0.02	0.12		99	1	
67515163	**Good identification**						**67515163**
Capnocytophaga ochracea		96.25	0.39	SOR 10			
Bacteroides fragilis		3.75	0.06	URE 1			
67515170	**Good identification**						**67515170**
Capnocytophaga ochracea		100.0	0.41	MLZ 10			
67515172	**Very good identification**						**67515172**
Capnocytophaga ochracea		99.65	0.57	MLZ 10			
Bacteroides fragilis		0.35	0.06	SOR 10			
67515302	**Acceptable identification**						**67515302**
					GRA	CAT	
Capnocytophaga ochracea		91.45	0.39	RHA 1	1	90	
Bacteroides fragilis		3.57	0.06	NAG 90	1	99	
Eubacterium contortum		1.79	0.11		99	1	
67515320	**Low discrimination**						**67515320**
					CAT		
Bacteroides vulgatus		82.76	0.47	NIT 1	3		
Capnocytophaga ochracea		13.72	0.39	ARA 93	90		
Bacteroides fragilis		2.79	0.18		99		
67515322	**Low discrimination**						**67515322**
Capnocytophaga ochracea		61.59	0.56	RHA 1			
Bacteroides fragilis		26.41	0.40				
Parabacteroides distasonis		9.38	0.54				
67515332	**Low discrimination**						**67515332**
Capnocytophaga ochracea		56.11	0.39	RHA 1			
Parabacteroides distasonis		41.42	0.49	MLZ 10			
Bacteroides fragilis		2.19	0.06				
67515362	**Acceptable identification**						**67515362**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		94.98	0.39	RHA 1			
Bacteroides fragilis		3.70	0.06	SOR 10			
Parabacteroides distasonis		1.32	0.20				
67515422	**Acceptable identification**						**67515422**
					GRA	CAT	
Capnocytophaga ochracea		93.58	0.43	LAC 99	1	90	
Bifidobacterium longum subsp. longum		4.12	0.28		99	1	
Bacteroides fragilis		2.30	0.06		1	99	
67515500	**Low discrimination**						**67515500**
					GRA	CAT	
Capnocytophaga ochracea		50.92	0.44	NAG 90	1	90	
Bifidobacterium longum subsp. longum		45.44	0.52	BGL 90	99	1	
Bacteroides vulgatus		2.36	0.15		1	3	
67515502	**Low discrimination**						**67515502**
					GRA	CAT	
Capnocytophaga ochracea		87.59	0.61	NAG 90	1	90	
Bifidobacterium longum subsp. longum		8.69	0.52		99	1	
Bifidobacterium adolescentis		3.23	0.46		99	1	
67515510	**Acceptable identification**						**67515510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.81	0.56	NIT 1	99	1	
Capnocytophaga ochracea		6.79	0.28		1	90	
Bifidobacterium adolescentis		0.25	0.13		99	1	
67515512	**Low discrimination**						**67515512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		51.77	0.56	NIT 1	99	1	
Capnocytophaga ochracea		34.08	0.44		1	90	
Bifidobacterium adolescentis		13.83	0.47		99	1	
67515520	**Low discrimination**						**67515520**
					GRA	CAT	
Capnocytophaga ochracea		52.13	0.61	BGL 90	1	90	
Bacteroides vulgatus		26.58	0.50		1	3	
Bifidobacterium longum subsp. longum		20.68	0.62		99	1	
67515522	**Good identification**						**67515522**
					GRA	CAT	
Capnocytophaga ochracea		93.36	0.77		1	90	
Bifidobacterium longum subsp. longum		4.11	0.62		99	1	
Bacteroides fragilis		2.29	0.40		1	99	
67515523	**Acceptable identification**						**67515523**
					GRA	CAT	
Capnocytophaga ochracea		93.58	0.43	URE 1	1	90	
Bifidobacterium longum subsp. longum		4.12	0.28		99	1	
Bacteroides fragilis		2.30	0.06		1	99	
67515530	**Low discrimination**						**67515530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		85.31	0.66	NIT 1	99	1	
Capnocytophaga ochracea		14.04	0.44		1	90	
Bacteroides vulgatus		0.65	0.15		1	3	
67515532	**Low discrimination**						**67515532**
					GRA	CAT	
Capnocytophaga ochracea		59.27	0.61	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		40.02	0.66		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		0.51	0.38		1	76	
67515542	**Acceptable identification**						**67515542**
					GRA	CAT	
Capnocytophaga ochracea		94.34	0.44	NAG 90	1	90	
Bifidobacterium adolescentis		3.87	0.30	SOR 10	99	1	
Actinomyces israelii		0.94	0.26		99	1	
67515560	**Acceptable identification**						**67515560**
					GRA	CAT	
Capnocytophaga ochracea		92.39	0.44	SOR 10	1	90	
Bacteroides vulgatus		4.28	0.15	BGL 90	1	3	
Bifidobacterium longum subsp. longum		3.33	0.28		99	1	
67515562	**Very good identification**						**67515562**
					GRA	CAT	
Capnocytophaga ochracea		99.38	0.61	SOR 10	1	90	
Bifidobacterium longum subsp. longum		0.40	0.28		99	1	
Bacteroides fragilis		0.22	0.06		1	99	
67515572	**Acceptable identification**						**67515572**
					GRA	CAT	
Capnocytophaga ochracea		94.22	0.44	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		5.78	0.32	SOR 10	99	1	
67515720	**Very good identification**						**67515720**
					GRA	CAT	
Bacteroides vulgatus		99.66	0.66	NIT 1	1	3	
Capnocytophaga ochracea		0.22	0.26		1	90	
Bifidobacterium longum subsp. longum		0.09	0.28		99	1	
67515722	**Low discrimination**						**67515722**
					CAT		
Capnocytophaga ochracea		46.91	0.43	RHA 1	90		
Parabacteroides distasonis		24.82	0.50		76		
Bacteroides vulgatus		23.91	0.32		3		
67515732	**Low discrimination**						**67515732**
					GRA	CAT	
Parabacteroides distasonis		53.56	0.46	NIT 1	1	76	
Capnocytophaga ochracea		20.89	0.26		1	90	
Bifidobacterium longum subsp. longum		14.11	0.32		99	1	
67517002	**Low discrimination**						**67517002**
					GRA	CAT	
Actinomyces israelii		85.85	0.37	SAL 92	99	1	
Capnocytophaga ochracea		10.04	0.05	CEL 93	1	90	
Actinomyces naeslundii		4.11	0.24		99	25	
67517100	**Low discrimination**						**67517100**
					GRA	CAT	
Actinomyces israelii		62.73	0.45	SAL 92	99	1	
Capnocytophaga ochracea		30.26	0.23	CEL 93	1	90	
Propionibacterium propionicum		3.81	0.23		99	1	
67517102	**Low discrimination**						**67517102**
					GRA	CAT	
Capnocytophaga ochracea		66.84	0.39	MAN 1	1	90	
Actinomyces israelii		23.10	0.48	NAG 90	99	1	
Bifidobacterium adolescentis		5.27	0.30		99	1	
67517120	**Low discrimination**						**67517120**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		80.62	0.39	MAN 1	1	90	
Bacteroides fragilis		16.37	0.18	BGL 90	1	99	
Actinomyces israelii		1.19	0.24		99	1	
67517122	**Low discrimination**						**67517122**
					GRA	CAT	
Capnocytophaga ochracea		69.71	0.56	MAN 1	1	90	
Bacteroides fragilis		29.89	0.40		1	99	
Actinomyces israelii		0.17	0.27		99	1	
67517132	**Good identification**						**67517132**
Capnocytophaga ochracea		96.25	0.39	MAN 1			
Bacteroides fragilis		3.75	0.06	MLZ 10			
67517140	**Low discrimination**						**67517140**
					GRA	CAT	
Actinomyces israelii		77.28	0.38	SAL 92	99	1	
Propionibacterium propionicum		12.07	0.23	CEL 93	99	1	
Capnocytophaga ochracea		10.65	0.07		1	90	
67517142	**Low discrimination**						**67517142**
					GRA	CAT	
Actinomyces israelii		48.65	0.41	SAL 92	99	1	
Capnocytophaga ochracea		40.23	0.23	CEL 93	1	90	
Propionibacterium propionicum		7.60	0.26		99	1	
67517162	**Good identification**						**67517162**
					GRA	CAT	
Capnocytophaga ochracea		95.46	0.39	MAN 1	1	90	
Bacteroides fragilis		3.72	0.06	SOR 10	1	99	
Actinomyces israelii		0.82	0.20		99	1	
67517402	**Acceptable identification**						**67517402**
Actinomyces israelii		92.99	0.36	SAL 92			
Bifidobacterium adolescentis		7.01	0.10	CEL 93			
67517500	**Low discrimination**						**67517500**
					GRA	CAT	
Actinomyces israelii		81.63	0.43	SAL 92	99	1	
Capnocytophaga ochracea		8.49	0.10	CEL 93	1	90	
Bifidobacterium longum subsp. longum		7.58	0.17		99	1	
67517502	**Low discrimination**						**67517502**
					GRA	CAT	
Bifidobacterium adolescentis		52.52	0.44	NIT 1	99	1	
Actinomyces israelii		28.16	0.46		99	1	
Capnocytophaga ochracea		17.58	0.26		1	90	
67517512	**Low discrimination**						**67517512**
Bifidobacterium adolescentis		87.55	0.46	NIT 1			
Actinomyces israelii		5.74	0.32				
Bifidobacterium longum subsp. longum		4.05	0.21				
67517522	**Acceptable identification**						**67517522**
					GRA	CAT	
Capnocytophaga ochracea		92.31	0.43	MAN 1	1	90	
Bifidobacterium longum subsp. longum		4.07	0.28		99	1	
Bacteroides fragilis		2.27	0.06		1	99	
67517540	**Good identification**						**67517540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.36	SAL 92			
67517542	**Low discrimination**						**67517542**
					GRA	CAT	
Actinomyces israelii		56.46	0.39	SAL 92	99	1	
Bifidobacterium adolescentis		33.47	0.29	CEL 93	99	1	
Capnocytophaga ochracea		10.07	0.10		1	90	
67521000	**Good identification**						**67521000**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67521001	**Low discrimination**						**67521001**
					SPO		
Actinomyces naeslundii		71.93	0.39	ESL 12	1		
Clostridium chauvoei		28.07	0.09	FRU 95	99		
67521003	**Low discrimination**						**67521003**
Actinomyces naeslundii		69.99	0.39	ESL 12			
Actinomyces viscosus		30.01	0.09	FRU 95			
67521021	**Low discrimination**						**67521021**
					SPO		
Actinomyces naeslundii		88.49	0.39	ESL 12	1		
Clostridium chauvoei		11.51	0.01	FRU 95	99		
67521023	**Good identification**						**67521023**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67521100	**Low discrimination**						**67521100**
					SPO		
Actinomyces naeslundii		68.55	0.44	ESL 12	1		
Clostridium chauvoei		14.10	0.09	FRU 95	99		
Clostridium perfringens		12.34	0.09		84		
67521101	**Acceptable identification**						**67521101**
					SPO		
Clostridium chauvoei		93.14	0.43	RAF 1	99		
Actinomyces naeslundii		6.86	0.47	TRE 10	1		
67521121	**Low discrimination**						**67521121**
					SPO		
Clostridium chauvoei		81.32	0.35	RAF 1	99		
Actinomyces naeslundii		17.97	0.47	TRE 10	1		
Clostridium perfringens		0.70	0.01		84		
67521123	**Low discrimination**						**67521123**
					GRA	SPO	
Actinomyces naeslundii		54.40	0.47	ESL 12	99	1	
Clostridium chauvoei		43.43	0.22	FRU 95	99	99	
Capnocytophaga ochracea		2.17	-.03		1	1	
67523000	**Good identification**						**67523000**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67523001	**Good identification**						**67523001**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67523002	**Good identification**						**67523002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
67523003	**Good identification**					**67523003**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67523020	**Good identification**					**67523020**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67523021	**Good identification**					**67523021**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67523022	**Good identification**					**67523022**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67523023	**Good identification**					**67523023**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67523100	**Acceptable identification**					**67523100**
Actinomyces naeslundii		91.22	0.49	ESL 12		
Propionibacterium propionicum		8.78	0.21	FRU 95		
67523101	**Acceptable identification**					**67523101**
					SPO	
Actinomyces naeslundii		93.40	0.52	ESL 12	1	
Clostridium chauvoei		6.60	0.09	FRU 95	99	
67523102	**Low discrimination**					**67523102**
Actinomyces naeslundii		87.39	0.49	ESL 12		
Propionibacterium propionicum		12.62	0.24	FRU 95		
67523103	**Good identification**					**67523103**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67523120	**Good identification**					**67523120**
Actinomyces naeslundii		97.06	0.49	ESL 12		
Clostridium perfringens		2.94	0.01	FRU 95		
67523121	**Good identification**					**67523121**
					SPO	
Actinomyces naeslundii		97.70	0.52	ESL 12	1	
Clostridium chauvoei		2.30	0.01	FRU 95	99	
67523122	**Good identification**					**67523122**
					GRA	
Actinomyces naeslundii		97.01	0.49	ESL 12	99	
Capnocytophaga ochracea		3.00	-.03	FRU 95	1	
67523123	**Good identification**					**67523123**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67523140	**Low discrimination**					**67523140**
Actinomyces naeslundii		64.71	0.36	ESL 12		
Propionibacterium propionicum		35.29	0.21	FRU 95		
67523141	**Good identification**					**67523141**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.39	ESL 12			
67523142	**Low discrimination**						**67523142**
Actinomyces naeslundii		55.01	0.36	ESL 12			
Propionibacterium propionicum		44.99	0.24	FRU 95			
67523143	**Good identification**						**67523143**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67523160	**Good identification**						**67523160**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67523161	**Good identification**						**67523161**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67523162	**Good identification**						**67523162**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67523163	**Good identification**						**67523163**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67525520	**Good identification**						**67525520**
					GRA		
Bifidobacterium longum subsp. longum		98.09	0.35	NIT 1	99		
Bacteroides vulgatus		1.91	-.08	FRU 90	1		
67525522	**Low discrimination**						**67525522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		86.93	0.35	NIT 1	99	1	
Capnocytophaga ochracea		7.25	0.09	FRU 90	1	90	
Parabacteroides distasonis		5.82	0.19		1	76	
67525530	**Good identification**						**67525530**
Bifidobacterium longum subsp. longum		100.0	0.39	NIT 1			
67525532	**Good identification**						**67525532**
					GRA		
Bifidobacterium longum subsp. longum		97.93	0.39	NIT 1	99		
Parabacteroides distasonis		2.07	0.15	FRU 90	1		
67530001	**Low discrimination**						**67530001**
Actinomyces naeslundii		85.50	0.37	ESL 12			
Actinomyces viscosus		14.50	0.00	GAL 96			
67530002	**Good identification**						**67530002**
Actinomyces viscosus		97.70	0.42	ESL 1			
Actinomyces naeslundii		1.29	0.34	TRE 7			
Propionibacterium granulosum		0.75	0.15				
67530021	**Good identification**						**67530021**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67530023	**Low discrimination**						**67530023**
Actinomyces naeslundii		85.50	0.37	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		14.50	0.00	GAL 96		
67530101	**Low discrimination**					**67530101**
					SPO	
Actinomyces naeslundii		66.22	0.45	ESL 12	1	
Clostridium chauvoei		26.76	0.15	GAL 96	99	
Actinomyces viscosus		7.02	0.04		1	
67530102	**Good identification**					**67530102**
					GRA	
Actinomyces viscosus		95.05	0.47	ESL 1	99	
Actinomyces naeslundii		2.01	0.42	TRE 7	99	
Capnocytophaga ochracea		1.68	0.15		1	
67530103	**Acceptable identification**					**67530103**
					SPO	
Actinomyces viscosus		90.73	0.38	ESL 1	1	
Actinomyces naeslundii		8.65	0.45	TRE 7	1	
Clostridium chauvoei		0.62	0.03		99	
67530121	**Low discrimination**					**67530121**
					SPO	
Actinomyces naeslundii		84.45	0.45	ESL 12	1	
Clostridium chauvoei		11.38	0.07	GAL 96	99	
Clostridium perfringens		4.18	0.01		84	
67530123	**Low discrimination**					**67530123**
					GRA	
Actinomyces naeslundii		86.47	0.45	ESL 12	99	
Actinomyces viscosus		9.16	0.04	GAL 96	99	
Capnocytophaga ochracea		4.37	-.03		1	
67531000	**Low discrimination**					**67531000**
					GRA	
Actinomyces naeslundii		62.39	0.58	ESL 12	99	
Clostridium perfringens		22.71	0.28		99	
Fusobacterium mortiferum		6.97	0.30		1	
67531001	**Very good identification**					**67531001**
					SPO	
Actinomyces naeslundii		95.43	0.61	ESL 12	1	
Clostridium chauvoei		4.57	0.15		99	
67531002	**Low discrimination**					**67531002**
Actinomyces viscosus		59.40	0.39	ESL 1		
Actinomyces naeslundii		30.78	0.58	TRE 7		
Clostridium perfringens		5.27	0.23			
67531003	**Low discrimination**					**67531003**
					SPO	
Actinomyces naeslundii		69.58	0.61	ESL 12	1	
Actinomyces viscosus		29.84	0.31		1	
Clostridium chauvoei		0.59	0.03		99	
67531010	**Low discrimination**					**67531010**
Actinomyces naeslundii		79.98	0.36	ESL 12		
Propionibacterium granulosum		20.02	0.11	MLZ 5		
67531011	**Good identification**					**67531011**
Actinomyces naeslundii		100.0	0.39	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67531012	**Low discrimination**						**67531012**
					GRA		
Actinomyces naeslundii		63.34	0.36	ESL 12	99		
Actinomyces viscosus		23.46	0.05	MLZ 5	99		
Capnocytophaga ochracea		13.20	-.01		1		
67531013	**Good identification**						**67531013**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67531020	**Acceptable identification**						**67531020**
					GRA	CAT	
Clostridium perfringens		91.22	0.54	LAC 88	99	1	
Actinomyces naeslundii		7.75	0.58		99	25	
Capnocytophaga ochracea		0.77	0.15		1	90	
67531021	**Good identification**						**67531021**
					SPO		
Actinomyces naeslundii		91.31	0.61	ESL 12	1		
Clostridium perfringens		7.24	0.20		84		
Clostridium chauvoei		1.46	0.07		99		
67531022	**Low discrimination**						**67531022**
					GRA	CAT	
Clostridium perfringens		73.78	0.48	LAC 88	99	1	
Actinomyces naeslundii		13.33	0.58		99	25	
Capnocytophaga ochracea		11.84	0.32		1	90	
67531023	**Very good identification**						**67531023**
					GRA	CAT	
Actinomyces naeslundii		95.85	0.61	ESL 12	99	25	
Clostridium perfringens		3.57	0.14		99	1	
Capnocytophaga ochracea		0.57	-.03		1	90	
67531031	**Good identification**						**67531031**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67531033	**Good identification**						**67531033**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67531040	**Low discrimination**						**67531040**
Actinomyces naeslundii		71.80	0.45	ESL 12			
Clostridium perfringens		28.20	0.16				
67531041	**Good identification**						**67531041**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67531042	**Low discrimination**						**67531042**
Actinomyces naeslundii		73.67	0.45	ESL 12			
Clostridium perfringens		13.62	0.10				
Actinomyces viscosus		8.14	0.05				
67531043	**Good identification**						**67531043**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67531060	**Acceptable identification**						**67531060**
					GRA	CAT	
Clostridium perfringens		92.28	0.42	LAC 88	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		7.27	0.45		99	25	
Capnocytophaga ochracea		0.45	-.01		1	90	
67531061	**Acceptable identification**						**67531061**
Actinomyces naeslundii		92.12	0.48	ESL 12			
Clostridium perfringens		7.88	0.08				
67531062	**Low discrimination**						**67531062**
					GRA	CAT	
Clostridium perfringens		79.31	0.36	LAC 88	99	1	
Actinomyces naeslundii		13.27	0.45		99	25	
Capnocytophaga ochracea		7.43	0.15		1	90	
67531063	**Good identification**						**67531063**
Actinomyces naeslundii		96.13	0.48	ESL 12			
Clostridium perfringens		3.87	0.02				
67531100	**Low discrimination**						**67531100**
					GRA	CAT	
Actinomyces naeslundii		39.71	0.66	ESL 12	99	25	
Clostridium perfringens		37.24	0.43		99	1	
Capnocytophaga ochracea		15.16	0.33		1	90	
67531101	**Low discrimination**						**67531101**
					SPO		
Clostridium chauvoei		62.23	0.50	RAF 1	99		
Actinomyces naeslundii		37.33	0.69	TRE 10	1		
Clostridium perfringens		0.24	0.09		84		
67531102	**Low discrimination**						**67531102**
					GRA		
Capnocytophaga ochracea		55.73	0.50	TRE 1	1		
Actinomyces viscosus		19.56	0.43	NAG 90	99		
Actinomyces naeslundii		16.23	0.66		99		
67531103	**Low discrimination**						**67531103**
					SPO		
Actinomyces naeslundii		62.97	0.69	ESL 12	1		
Clostridium chauvoei		18.52	0.37		99		
Actinomyces viscosus		16.87	0.35		1		
67531110	**Low discrimination**						**67531110**
					GRA	CAT	
Actinomyces naeslundii		48.29	0.44	ESL 12	99	25	
Capnocytophaga ochracea		38.91	0.17	MLZ 5	1	90	
Clostridium perfringens		8.69	0.09		99	1	
67531111	**Low discrimination**						**67531111**
					SPO		
Actinomyces naeslundii		75.76	0.47	ESL 12	1		
Clostridium chauvoei		24.24	0.15	MLZ 5	99		
67531113	**Low discrimination**						**67531113**
					SPO		
Actinomyces naeslundii		86.45	0.47	ESL 12	1		
Clostridium chauvoei		4.88	0.03	MLZ 5	99		
Actinomyces viscosus		4.44	0.01		1		
67531120	**Low discrimination**						**67531120**
					GRA	CAT	
Clostridium perfringens		86.29	0.69		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		9.78	0.50		1	90	
Actinomyces naeslundii		2.85	0.66		99	25	
67531121	**Low discrimination**						**67531121**
					SPO		
Actinomyces naeslundii		56.01	0.69	ESL 12	1		
Clostridium chauvoei		31.12	0.41		99		
Clostridium perfringens		11.44	0.35		84		
67531122	**Low discrimination**						**67531122**
					GRA	CAT	
Capnocytophaga ochracea		66.02	0.66	TRE 1	1	90	
Clostridium perfringens		30.47	0.63		99	1	
Actinomyces naeslundii		2.14	0.66		99	25	
67531123	**Low discrimination**						**67531123**
					GRA	SPO	
Actinomyces naeslundii		71.06	0.69	ESL 12	99	1	
Capnocytophaga ochracea		14.79	0.32		1	1	
Clostridium chauvoei		6.97	0.28		99	99	
67531131	**Low discrimination**						**67531131**
					GRA	SPO	
Actinomyces naeslundii		83.71	0.47	ESL 12	99	1	
Clostridium chauvoei		8.93	0.07	MLZ 5	99	99	
Capnocytophaga ochracea		4.09	-.01		1	1	
67531132	**Acceptable identification**						**67531132**
					GRA	CAT	
Capnocytophaga ochracea		91.49	0.50	TRE 1	1	90	
Clostridium perfringens		3.84	0.29	MLZ 10	99	1	
Parabacteroides distasonis		2.85	0.36		1	76	
67531133	**Low discrimination**						**67531133**
					GRA		
Actinomyces naeslundii		69.47	0.47	ESL 12	99		
Capnocytophaga ochracea		30.53	0.15	MLZ 5	1		
67531141	**Acceptable identification**						**67531141**
					SPO		
Actinomyces naeslundii		91.29	0.56	ESL 12	1		
Clostridium chauvoei		8.71	0.15		99		
67531143	**Good identification**						**67531143**
					SPO		
Actinomyces naeslundii		95.54	0.56	ESL 12	1		
Clostridium chauvoei		1.61	0.03		99		
Actinomyces viscosus		1.46	0.01		1		
67531160	**Good identification**						**67531160**
					GRA	CAT	
Clostridium perfringens		91.14	0.57		99	1	
Capnocytophaga ochracea		6.02	0.33		1	90	
Actinomyces naeslundii		2.79	0.53		99	25	
67531161	**Low discrimination**						**67531161**
					SPO		
Actinomyces naeslundii		78.94	0.56	ESL 12	1		
Clostridium perfringens		17.40	0.22		84		
Clostridium chauvoei		2.51	0.07		99		
67531162	**Low discrimination**						**67531162**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		54.25	0.50	TRE 1	1	90	
Clostridium perfringens		42.92	0.51	SOR 10	99	1	
Actinomyces naeslundii		2.79	0.53		99	25	
67531163	**Low discrimination**						**67531163**
					GRA	CAT	
Actinomyces naeslundii		80.99	0.56	ESL 12	99	25	
Capnocytophaga ochracea		10.62	0.15		1	90	
Clostridium perfringens		8.40	0.17		99	1	
67531322	**Low discrimination**						**67531322**
					GRA	CAT	
Parabacteroides distasonis		49.58	0.49	NIT 1	1	76	
Capnocytophaga ochracea		25.98	0.32	XYL 90	1	90	
Clostridium perfringens		11.99	0.29		99	1	
67531332	**Low discrimination**						**67531332**
Parabacteroides distasonis		89.44	0.44	NIT 1			
Capnocytophaga ochracea		9.67	0.15	XYL 90			
Clostridium clostridioforme		0.89	0.04				
67531502	**Low discrimination**						**67531502**
					GRA	CAT	
Capnocytophaga ochracea		73.39	0.37	TRE 1	1	90	
Bifidobacterium longum subsp. longum		15.01	0.33	NAG 90	99	1	
Bifidobacterium adolescentis		6.68	0.29		99	1	
67531510	**Good identification**						**67531510**
					GRA	CAT	
Bifidobacterium longum subsp. longum		96.58	0.37	NIT 1	99	1	
Capnocytophaga ochracea		3.42	0.04		1	90	
67531512	**Low discrimination**						**67531512**
					GRA	CAT	
Bifidobacterium longum subsp. longum		59.85	0.37	NIT 1	99	1	
Bifidobacterium adolescentis		19.11	0.30		99	1	
Capnocytophaga ochracea		19.10	0.20		1	90	
67531520	**Low discrimination**						**67531520**
					GRA	CAT	
Capnocytophaga ochracea		42.44	0.37	TRE 1	1	90	
Bifidobacterium longum subsp. longum		34.71	0.43	BGL 90	99	1	
Clostridium perfringens		21.44	0.35		99	1	
67531522	**Low discrimination**						**67531522**
					GRA	CAT	
Capnocytophaga ochracea		87.56	0.53	TRE 1	1	90	
Bifidobacterium longum subsp. longum		7.96	0.43		99	1	
Clostridium perfringens		2.31	0.29		99	1	
67531530	**Acceptable identification**						**67531530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		92.30	0.47	NIT 1	99	1	
Capnocytophaga ochracea		7.36	0.20		1	90	
Clostridium perfringens		0.34	0.01		99	1	
67531532	**Low discrimination**						**67531532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		55.44	0.47	NIT 1	99	1	
Capnocytophaga ochracea		39.81	0.37		1	90	
Parabacteroides distasonis		4.31	0.33		1	76	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67531562	**Acceptable identification**						**67531562**
					GRA	CAT	
Capnocytophaga ochracea		94.92	0.37	TRE 1	1	90	
Clostridium perfringens		4.30	0.17	SOR 10	99	1	
Bifidobacterium longum subsp. longum		0.78	0.09		99	1	
67531722	**Low discrimination**						**67531722**
Parabacteroides distasonis		81.84	0.45	NIT 1			
Capnocytophaga ochracea		12.35	0.19	XYL 90			
Clostridium clostridioforme		2.46	0.14				
67531732	**Low discrimination**						**67531732**
					GRA		
Parabacteroides distasonis		74.14	0.41	NIT 1	1		
Clostridium clostridioforme		18.88	0.25	XYL 90	50		
Bifidobacterium longum subsp. longum		3.22	0.13		99		
67532000	**Low discrimination**						**67532000**
					CAT		
Actinomyces naeslundii		44.50	0.39	ESL 12	25		
Propionibacterium granulosum		40.10	0.24	GAL 96	90		
Propionibacterium propionicum		15.40	0.21		1		
67532001	**Good identification**						**67532001**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67532002	**Low discrimination**						**67532002**
Actinomyces naeslundii		46.79	0.39	ESL 12			
Propionibacterium propionicum		24.28	0.24	GAL 96			
Actinomyces viscosus		18.40	0.08				
67532003	**Acceptable identification**						**67532003**
Actinomyces naeslundii		91.97	0.42	ESL 12			
Actinomyces viscosus		8.03	0.00	GAL 96			
67532020	**Low discrimination**						**67532020**
Actinomyces naeslundii		81.62	0.39	ESL 12			
Propionibacterium granulosum		18.38	0.13	GAL 96			
67532021	**Good identification**						**67532021**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67532022	**Good identification**						**67532022**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67532023	**Good identification**						**67532023**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67532100	**Acceptable identification**						**67532100**
Propionibacterium propionicum		92.33	0.55	ESL 1			
Actinomyces naeslundii		7.67	0.47				
67532101	**Low discrimination**						**67532101**
Actinomyces naeslundii		52.87	0.50	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		47.13	0.39	GAL 96			
67532102	**Good identification**						**67532102**
Propionibacterium propionicum		93.55	0.58	ESL 1			
Actinomyces naeslundii		5.18	0.47				
Actinomyces viscosus		1.27	0.12				
67532103	**Low discrimination**						**67532103**
Propionibacterium propionicum		55.91	0.42	ESL 1			
Actinomyces naeslundii		41.81	0.50	URE 10			
Actinomyces viscosus		2.28	0.04				
67532120	**Low discrimination**						**67532120**
Actinomyces naeslundii		86.22	0.47	ESL 12			
Propionibacterium propionicum		10.48	0.21	GAL 96			
Clostridium perfringens		3.30	0.01				
67532121	**Good identification**						**67532121**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67532122	**Low discrimination**						**67532122**
					GRA	CAT	
Actinomyces naeslundii		81.88	0.47	ESL 12	99	25	
Propionibacterium propionicum		14.93	0.24	GAL 96	99	1	
Capnocytophaga ochracea		3.19	-.03		1	90	
67532123	**Good identification**						**67532123**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67532140	**Good identification**						**67532140**
Propionibacterium propionicum		98.56	0.55	ESL 1			
Actinomyces naeslundii		1.45	0.34				
67532141	**Low discrimination**						**67532141**
Propionibacterium propionicum		83.48	0.39	ESL 1			
Actinomyces naeslundii		16.52	0.37	URE 10			
67532142	**Very good identification**						**67532142**
Propionibacterium propionicum		99.03	0.58	ESL 1			
Actinomyces naeslundii		0.97	0.34				
67532143	**Low discrimination**						**67532143**
Propionibacterium propionicum		88.34	0.42	ESL 1			
Actinomyces naeslundii		11.66	0.37	URE 10			
67532161	**Good identification**						**67532161**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67532163	**Good identification**						**67532163**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67533000	**Good identification**						**67533000**
					CAT		
Actinomyces naeslundii		92.97	0.63	ESL 12	25		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium granulosum		3.49	0.24		90		
Actinomyces israelii		2.21	0.26		1		
67533001	**Very good identification**						**67533001**
Actinomyces naeslundii		100.0	0.66	ESL 12			
67533002	**Good identification**						**67533002**
Actinomyces naeslundii		92.88	0.63	ESL 12			
Actinomyces israelii		3.31	0.29				
Propionibacterium propionicum		2.01	0.24				
67533003	**Very good identification**						**67533003**
Actinomyces naeslundii		100.0	0.66	ESL 12			
67533010	**Acceptable identification**						**67533010**
Actinomyces naeslundii		91.14	0.41	ESL 12			
Propionibacterium granulosum		8.86	0.09	MLZ 5			
67533011	**Good identification**						**67533011**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67533012	**Acceptable identification**						**67533012**
Actinomyces naeslundii		90.83	0.41	ESL 12			
Actinomyces israelii		9.17	0.15	MLZ 5			
67533013	**Good identification**						**67533013**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67533020	**Good identification**						**67533020**
					CAT		
Actinomyces naeslundii		93.40	0.63	ESL 12	25		
Clostridium perfringens		5.72	0.20		1		
Propionibacterium granulosum		0.88	0.13		90		
67533021	**Very good identification**						**67533021**
Actinomyces naeslundii		100.0	0.66	ESL 12			
67533022	**Very good identification**						**67533022**
					GRA	CAT	
Actinomyces naeslundii		96.76	0.63	ESL 12	99	25	
Clostridium perfringens		2.79	0.14		99	1	
Capnocytophaga ochracea		0.45	-.03		1	90	
67533023	**Very good identification**						**67533023**
Actinomyces naeslundii		100.0	0.66	ESL 12			
67533030	**Good identification**						**67533030**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67533031	**Good identification**						**67533031**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67533032	**Good identification**						**67533032**
Actinomyces naeslundii		100.0	0.41	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67533033	**Good identification**					**67533033**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67533040	**Low discrimination**					**67533040**
Actinomyces naeslundii		88.19	0.50	ESL 12		
Propionibacterium propionicum		7.20	0.21			
Actinomyces israelii		4.61	0.19			
67533041	**Good identification**					**67533041**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67533042	**Low discrimination**					**67533042**
Actinomyces naeslundii		83.27	0.50	ESL 12		
Propionibacterium propionicum		10.20	0.24			
Actinomyces israelii		6.53	0.22			
67533043	**Good identification**					**67533043**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67533060	**Acceptable identification**					**67533060**
Actinomyces naeslundii		93.80	0.50	ESL 12		
Clostridium perfringens		6.20	0.08			
67533061	**Good identification**					**67533061**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67533062	**Good identification**					**67533062**
Actinomyces naeslundii		96.98	0.50	ESL 12		
Clostridium perfringens		3.02	0.02			
67533063	**Good identification**					**67533063**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67533100	**Low discrimination**					**67533100**
Actinomyces naeslundii		64.86	0.71	ESL 12		
Propionibacterium propionicum		32.53	0.55			
Actinomyces israelii		2.16	0.36			
67533101	**Very good identification**					**67533101**
					SPO	
Actinomyces naeslundii		95.62	0.74	ESL 12	1	
Propionibacterium propionicum		3.55	0.39		1	
Clostridium chauvoei		0.83	0.15		99	
67533102	**Low discrimination**					**67533102**
Actinomyces naeslundii		54.58	0.71	ESL 12		
Propionibacterium propionicum		41.06	0.58			
Actinomyces israelii		2.73	0.39			
67533103	**Good identification**					**67533103**
					SPO	
Actinomyces naeslundii		94.46	0.74	ESL 12	1	
Propionibacterium propionicum		5.26	0.42		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		0.15	0.03		99		
67533110	**Low discrimination**						**67533110**
Actinomyces naeslundii		83.97	0.49	ESL 12			
Propionibacterium propionicum		8.08	0.21	MLZ 5			
Actinomyces israelii		7.95	0.22				
67533111	**Good identification**						**67533111**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67533112	**Low discrimination**						**67533112**
Actinomyces naeslundii		71.19	0.49	ESL 12			
Propionibacterium propionicum		10.28	0.24	MLZ 5			
Actinomyces israelii		10.11	0.25				
67533113	**Good identification**						**67533113**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67533120	**Low discrimination**						**67533120**
					GRA	CAT	
Actinomyces naeslundii		84.41	0.71	ESL 12	99	25	
Clostridium perfringens		13.32	0.35		99	1	
Capnocytophaga ochracea		1.51	0.15		1	90	
67533121	**Very good identification**						**67533121**
					SPO		
Actinomyces naeslundii		99.61	0.74	ESL 12	1		
Clostridium chauvoei		0.29	0.07		99		
Clostridium perfringens		0.11	0.01		84		
67533122	**Low discrimination**						**67533122**
					GRA	CAT	
Actinomyces naeslundii		80.15	0.71	ESL 12	99	25	
Capnocytophaga ochracea		12.90	0.32		1	90	
Clostridium perfringens		5.95	0.29		99	1	
67533123	**Very good identification**						**67533123**
					GRA		
Actinomyces naeslundii		99.89	0.74	ESL 12	99		
Capnocytophaga ochracea		0.11	-.03		1		
67533130	**Acceptable identification**						**67533130**
					GRA	CAT	
Actinomyces naeslundii		93.63	0.49	ESL 12	99	25	
Capnocytophaga ochracea		3.53	-.01	MLZ 5	1	90	
Clostridium perfringens		2.83	0.01		99	1	
67533131	**Good identification**						**67533131**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67533132	**Low discrimination**						**67533132**
					GRA		
Actinomyces naeslundii		74.65	0.49	ESL 12	99		
Capnocytophaga ochracea		25.35	0.15	MLZ 5	1		
67533133	**Good identification**						**67533133**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67533140	**Low discrimination**						**67533140**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		72.58	0.55	ESL 1			
Actinomyces naeslundii		25.54	0.58				
Actinomyces israelii		1.88	0.29				
67533141	**Low discrimination**						**67533141**
Actinomyces naeslundii		82.61	0.61	ESL 12			
Propionibacterium propionicum		17.39	0.39				
67533142	**Low discrimination**						**67533142**
Propionibacterium propionicum		79.17	0.58	ESL 1			
Actinomyces naeslundii		18.57	0.58				
Actinomyces israelii		2.05	0.32				
67533143	**Low discrimination**						**67533143**
Actinomyces naeslundii		76.00	0.61	ESL 12			
Propionibacterium propionicum		24.00	0.42				
67533150	**Low discrimination**						**67533150**
Actinomyces naeslundii		57.02	0.36	ESL 12			
Propionibacterium propionicum		31.09	0.21	MLZ 5			
Actinomyces israelii		11.89	0.15				
67533151	**Good identification**						**67533151**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67533152	**Low discrimination**						**67533152**
Actinomyces naeslundii		46.93	0.36	ESL 12			
Propionibacterium propionicum		38.39	0.24	MLZ 5			
Actinomyces israelii		14.68	0.18				
67533153	**Good identification**						**67533153**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67533160	**Low discrimination**						**67533160**
Actinomyces naeslundii		82.63	0.58	ESL 12			
Clostridium perfringens		14.07	0.22				
Propionibacterium propionicum		2.37	0.21				
67533161	**Very good identification**						**67533161**
Actinomyces naeslundii		100.0	0.61	ESL 12			
67533162	**Low discrimination**						**67533162**
					GRA	CAT	
Actinomyces naeslundii		81.67	0.58	ESL 12	99	25	
Capnocytophaga ochracea		8.27	0.15		1	90	
Clostridium perfringens		6.54	0.17		99	1	
67533163	**Very good identification**						**67533163**
Actinomyces naeslundii		100.0	0.61	ESL 12			
67533170	**Good identification**						**67533170**
Actinomyces naeslundii		100.0	0.36	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67533171	**Good identification**					**67533171**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67533172	**Low discrimination**					**67533172**
					GRA	
Actinomyces naeslundii		82.39	0.36	ESL 12	99	
Capnocytophaga ochracea		17.61	-.01	MLZ 5	1	
67533173	**Good identification**					**67533173**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67533300	**Low discrimination**					**67533300**
Actinomyces naeslundii		57.15	0.37	ESL 12		
Propionibacterium propionicum		28.66	0.21	RHA 1		
Actinomyces israelii		14.20	0.17			
67533301	**Good identification**					**67533301**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67533302	**Low discrimination**					**67533302**
Actinomyces naeslundii		47.06	0.37	ESL 12		
Propionibacterium propionicum		35.40	0.24	RHA 1		
Actinomyces israelii		17.54	0.20			
67533303	**Good identification**					**67533303**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67533320	**Low discrimination**					**67533320**
Actinomyces naeslundii		86.38	0.37	ESL 12		
Clostridium perfringens		13.62	0.01	RHA 1		
67533321	**Good identification**					**67533321**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67533322	**Low discrimination**					**67533322**
					GRA	
Actinomyces naeslundii		61.50	0.37	ESL 12	99	
Parabacteroides distasonis		18.88	0.15	RHA 1	1	
Capnocytophaga ochracea		9.89	-.03		1	
67533323	**Good identification**					**67533323**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67533501	**Good identification**					**67533501**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67533502	**Low discrimination**					**67533502**
Actinomyces israelii		48.13	0.38	SAL 92		
Bifidobacterium adolescentis		27.34	0.27	CEL 93		
Actinomyces naeslundii		11.88	0.37	XYL 92		
67533503	**Good identification**					**67533503**
Actinomyces naeslundii		100.0	0.40	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
67533520	**Low discrimination**							**67533520**
					GRA	CAT		
Actinomyces naeslundii		52.69	0.37	ESL 12	99	25		
Capnocytophaga ochracea		16.45	0.02	ARA 1	1	90		
Bifidobacterium longum subsp. longum		13.46	0.09		99	1		
67533521	**Good identification**							**67533521**
Actinomyces naeslundii		100.0	0.40	ESL 12				
67533523	**Good identification**							**67533523**
Actinomyces naeslundii		100.0	0.40	ESL 12				
67535100	**Low discrimination**							**67535100**
					COC	GRA	CAT	
Actinomyces israelii		39.43	0.41	SAL 92	1	99	1	
Capnocytophaga ochracea		28.65	0.23	CEL 93	1	1	90	
Atopobium parvulum		21.96	0.23		99	99	1	
67535102	**Low discrimination**							**67535102**
					GRA	CAT		
Capnocytophaga ochracea		71.22	0.39	TRE 1	1	90		
Actinomyces israelii		16.34	0.45	NAG 90	99	1		
Bifidobacterium adolescentis		4.57	0.29		99	1		
67535120	**Low discrimination**							**67535120**
					GRA	CAT		
Capnocytophaga ochracea		60.00	0.39	TRE 1	1	90		
Clostridium perfringens		21.39	0.35	BGL 90	99	1		
Bacteroides fragilis		12.19	0.18		1	99		
67535122	**Low discrimination**							**67535122**
Capnocytophaga ochracea		59.19	0.56	TRE 1				
Bacteroides fragilis		25.38	0.40					
Parabacteroides distasonis		13.69	0.57					
67535132	**Low discrimination**							**67535132**
Parabacteroides distasonis		51.07	0.53	NIT 1				
Capnocytophaga ochracea		45.56	0.39					
Bacteroides fragilis		1.78	0.06					
67535162	**Acceptable identification**							**67535162**
					GRA	CAT		
Capnocytophaga ochracea		91.10	0.39	TRE 1	1	90		
Bacteroides fragilis		3.55	0.06	SOR 10	1	99		
Clostridium perfringens		2.91	0.17		99	1		
67535322	**Very good identification**							**67535322**
Parabacteroides distasonis		97.53	0.65	NIT 1				
Capnocytophaga ochracea		1.42	0.21					
Bacteroides fragilis		0.61	0.06					
67535332	**Very good identification**							**67535332**
Parabacteroides distasonis		98.60	0.61	NIT 1				
Clostridium clostridioforme		1.10	0.22					
Capnocytophaga ochracea		0.30	0.05					
67535500	**Low discrimination**							**67535500**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium longum subsp. longum		67.16	0.41	NIT 1	99		
Actinomyces israelii		19.41	0.40		99		
Mitsuokella multacida		8.05	0.21		1		
67535502	**Low discrimination**						**67535502**
Bifidobacterium longum subsp. longum		34.28	0.41	NIT 1			
Bifidobacterium adolescentis		34.02	0.43				
Actinomyces israelii		14.86	0.43				
67535510	**Acceptable identification**						**67535510**
					GRA		
Bifidobacterium longum subsp. longum		94.70	0.45	NIT 1	99		
Actinomyces israelii		2.40	0.26		99		
Mitsuokella multacida		2.22	0.13		1		
67535512	**Low discrimination**						**67535512**
Bifidobacterium longum subsp. longum		52.76	0.45	NIT 1			
Bifidobacterium adolescentis		37.58	0.44				
Clostridium clostridioforme		5.76	0.27				
67535520	**Low discrimination**						**67535520**
					GRA	CAT	
Bifidobacterium longum subsp. longum		74.75	0.52	NIT 1	99	1	
Bacteroides vulgatus		16.01	0.26		1	3	
Capnocytophaga ochracea		7.62	0.26		1	90	
67535522	**Low discrimination**						**67535522**
					GRA	CAT	
Bifidobacterium longum subsp. longum		36.90	0.52	NIT 1	99	1	
Capnocytophaga ochracea		33.84	0.43		1	90	
Parabacteroides distasonis		27.18	0.53		1	76	
67535530	**Good identification**						**67535530**
					GRA	CAT	
Bifidobacterium longum subsp. longum		98.84	0.56	NIT 1	99	1	
Capnocytophaga ochracea		0.66	0.10		1	90	
Parabacteroides distasonis		0.23	0.15		1	76	
67535532	**Low discrimination**						**67535532**
					GRA	CAT	
Bifidobacterium longum subsp. longum		75.50	0.56	NIT 1	99	1	
Parabacteroides distasonis		17.59	0.49		1	76	
Capnocytophaga ochracea		4.52	0.26		1	90	
67535542	**Low discrimination**						**67535542**
					GRA	CAT	
Actinomyces israelii		46.84	0.36	SAL 92	99	1	
Bifidobacterium adolescentis		34.08	0.27	CEL 93	99	1	
Capnocytophaga ochracea		12.58	0.10		1	90	
67535712	**Acceptable identification**						**67535712**
					GRA		
Clostridium clostridioforme		94.08	0.42	NIT 1	50		
Parabacteroides distasonis		3.32	0.23	SAL 88	1		
Bifidobacterium longum subsp. longum		1.19	0.11		99		
67535720	**Good identification**						**67535720**
Bacteroides vulgatus		97.65	0.42	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Parabacteroides distasonis		1.13	0.27	TRE 4		
Mitsuokella multacida		0.71	0.15			
67535722	**Low discrimination**					**67535722**
Parabacteroides distasonis		88.98	0.62	NIT 1		
Bacteroides thetaiotaomicron		6.45	0.26			
Clostridium clostridioforme		3.01	0.31			
67535732	**Low discrimination**					**67535732**
Parabacteroides distasonis		73.91	0.57	NIT 1		
Clostridium clostridioforme		21.15	0.42			
Bacteroides thetaiotaomicron		3.87	0.19			
67537000	**Good identification**					**67537000**
Actinomyces israelii		96.43	0.44	SAL 92		
Actinomyces naeslundii		3.57	0.29	CEL 93		
67537002	**Good identification**					**67537002**
Actinomyces israelii		97.59	0.47	SAL 92		
Actinomyces naeslundii		2.41	0.29	CEL 93		
67537040	**Good identification**					**67537040**
Actinomyces israelii		100.0	0.37	SAL 92		
67537042	**Good identification**					**67537042**
Actinomyces israelii		100.0	0.40	SAL 92		
67537100	**Good identification**					**67537100**
Actinomyces israelii		96.20	0.54	SAL 92		
Actinomyces naeslundii		2.54	0.37	CEL 93		
Propionibacterium propionicum		1.27	0.21			
67537101	**Low discrimination**					**67537101**
Actinomyces naeslundii		79.64	0.40	ESL 12		
Actinomyces israelii		20.36	0.20	XYL 1		
67537102	**Good identification**					**67537102**
Actinomyces israelii		92.68	0.57	SAL 92		
Bifidobacterium adolescentis		3.75	0.27	CEL 93		
Actinomyces naeslundii		1.63	0.37			
67537103	**Low discrimination**					**67537103**
Actinomyces naeslundii		72.29	0.40	ESL 12		
Actinomyces israelii		27.71	0.23	XYL 1		
67537110	**Good identification**					**67537110**
Actinomyces israelii		100.0	0.40	SAL 92		
67537112	**Low discrimination**					**67537112**
Actinomyces israelii		75.16	0.43	SAL 92		
Bifidobacterium adolescentis		24.84	0.29	CEL 93		
67537121	**Good identification**					**67537121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.40	ESL 12		
67537123	**Good identification**					**67537123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67537140	**Good identification**					**67537140**
Actinomyces israelii		95.61	0.47	SAL 92		
Propionibacterium propionicum		3.25	0.21	CEL 93		
Actinomyces naeslundii		1.14	0.24			
67537142	**Acceptable identification**					**67537142**
Actinomyces israelii		94.81	0.50	SAL 92		
Propionibacterium propionicum		3.22	0.24	CEL 93		
Bifidobacterium adolescentis		1.22	0.12			
67537152	**Acceptable identification**					**67537152**
Actinomyces israelii		90.49	0.36	SAL 92		
Bifidobacterium adolescentis		9.51	0.13	CEL 93		
67537300	**Good identification**					**67537300**
Actinomyces israelii		100.0	0.35	SAL 92		
67537302	**Good identification**					**67537302**
Actinomyces israelii		100.0	0.38	SAL 92		
67537400	**Good identification**					**67537400**
Actinomyces israelii		100.0	0.43	SAL 92		
67537402	**Good identification**					**67537402**
Actinomyces israelii		98.68	0.46	SAL 92		
Bifidobacterium adolescentis		1.32	0.07	CEL 93		
67537440	**Good identification**					**67537440**
Actinomyces israelii		100.0	0.36	SAL 92		
67537442	**Good identification**					**67537442**
Actinomyces israelii		100.0	0.39	SAL 92		
67537500	**Low discrimination**					**67537500**
					GRA	
Actinomyces israelii		84.60	0.53	SAL 92	99	
Mitsuokella multacida		14.45	0.27	CEL 93	1	
Bifidobacterium longum subsp. longum		0.52	0.07		99	
67537502	**Low discrimination**					**67537502**
					GRA	
Actinomyces israelii		74.22	0.56	SAL 92	99	
Bifidobacterium adolescentis		24.54	0.41	CEL 93	99	
Mitsuokella multacida		0.94	0.11		1	
67537510	**Low discrimination**					**67537510**
					GRA	
Actinomyces israelii		67.01	0.39	SAL 92	99	
Mitsuokella multacida		25.53	0.19	CEL 93	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. longum		4.71	0.11	MLZ 13	99		
67537512	**Low discrimination**						**67537512**
Bifidobacterium adolescentis		70.83	0.43	NIT 1			
Actinomyces israelii		26.20	0.42				
Bifidobacterium longum subsp. longum		1.23	0.11				
67537522	**Low discrimination**						**67537522**
					GRA	CAT	
Actinomyces israelii		57.60	0.35	SAL 92	99	1	
Bifidobacterium longum subsp. longum		14.85	0.17	CEL 93	99	1	
Capnocytophaga ochracea		13.61	0.09	NAG 6	1	90	
67537540	**Low discrimination**						**67537540**
					GRA		
Actinomyces israelii		69.48	0.46	SAL 92	99		
Mitsuokella multacida		30.52	0.27	CEL 93	1		
67537542	**Low discrimination**						**67537542**
					GRA		
Actinomyces israelii		87.91	0.49	SAL 92	99		
Bifidobacterium adolescentis		9.24	0.26	CEL 93	99		
Mitsuokella multacida		2.86	0.11		1		
67537702	**Low discrimination**						**67537702**
					GRA		
Actinomyces israelii		73.72	0.37	SAL 92	99		
Mitsuokella multacida		23.01	0.15	CEL 93	1		
Bifidobacterium adolescentis		3.27	0.07	RHA 7	99		
67541001	**Good identification**						**67541001**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67541021	**Good identification**						**67541021**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67541023	**Good identification**						**67541023**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67541102	**Low discrimination**						**67541102**
					GRA		
Capnocytophaga ochracea		76.11	0.39	FRU 99	1		
Actinomyces viscosus		19.86	0.31	NAG 90	99		
Actinomyces naeslundii		3.19	0.41		99		
67541120	**Low discrimination**						**67541120**
					GRA	CAT	
Capnocytophaga ochracea		64.34	0.39	FRU 99	1	90	
Prevotella melaninogenica		31.74	0.30	BGL 90	1	1	
Actinomyces naeslundii		2.69	0.41		99	25	
67541121	**Low discrimination**						**67541121**
					GRA	SPO	
Actinomyces naeslundii		54.27	0.44	ESL 12	99	1	
Clostridium chauvoei		32.69	0.17	FRU 95	99	99	
Capnocytophaga ochracea		8.73	0.05		1	1	
67541122	**Good identification**						**67541122**
					CAT		
Capnocytophaga ochracea		96.06	0.56	FRU 99	90		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		2.27	0.27		1		
Prevotella melaninogenica		1.00	0.18		1		
67541132	**Good identification**						**67541132**
Capnocytophaga ochracea		100.0	0.39	FRU 99			
67541162	**Good identification**						**67541162**
					GRA		
Capnocytophaga ochracea		99.27	0.39	FRU 99	1		
Actinomyces naeslundii		0.73	0.28	SOR 10	99		
67541522	**Good identification**						**67541522**
Capnocytophaga ochracea		100.0	0.43	FRU 99			
67543000	**Good identification**						**67543000**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67543001	**Good identification**						**67543001**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67543002	**Good identification**						**67543002**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67543003	**Good identification**						**67543003**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67543020	**Good identification**						**67543020**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67543021	**Good identification**						**67543021**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67543022	**Good identification**						**67543022**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67543023	**Good identification**						**67543023**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67543100	**Good identification**						**67543100**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67543101	**Good identification**						**67543101**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67543102	**Low discrimination**						**67543102**
					GRA	CAT	
Actinomyces naeslundii		85.18	0.46	ESL 12	99	25	
Capnocytophaga ochracea		10.59	0.05	FRU 95	1	90	
Actinomyces israelii		4.23	0.14		99	1	
67543103	**Good identification**						**67543103**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67543120	**Low discrimination**						**67543120**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces naeslundii		84.34	0.46	ESL 12	99	25	
Capnocytophaga ochracea		10.49	0.05	FRU 95	1	90	
Prevotella melaninogenica		5.17	-.04		1	1	
67543121	**Good identification**						**67543121**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67543123	**Good identification**						**67543123**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67543141	**Good identification**						**67543141**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67543143	**Good identification**						**67543143**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67543161	**Good identification**						**67543161**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67543163	**Good identification**						**67543163**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67545122	**Low discrimination**						**67545122**
					CAT		
Capnocytophaga ochracea		57.25	0.45	FRU 99	90		
Bacteroides fragilis		42.08	0.34		99		
Prevotella oralis		0.67	0.12		1		
67550002	**Good identification**						**67550002**
					GRA		
Actinomyces viscosus		99.44	0.52	ESL 1	99		
Actinomyces naeslundii		0.25	0.32		99		
Capnocytophaga ochracea		0.22	0.05		1		
67550003	**Good identification**						**67550003**
Actinomyces viscosus		98.86	0.44	ESL 1			
Actinomyces naeslundii		1.14	0.35				
67550101	**Low discrimination**						**67550101**
Actinomyces naeslundii		64.60	0.42	ESL 12			
Actinomyces viscosus		35.40	0.14	GAL 96			
67550102	**Low discrimination**						**67550102**
					GRA		
Actinomyces viscosus		88.48	0.56	ESL 1	99		
Capnocytophaga ochracea		10.94	0.39		1		
Actinomyces naeslundii		0.36	0.39		99		
67550103	**Good identification**						**67550103**
					GRA		
Actinomyces viscosus		97.83	0.48	ESL 1	99		
Actinomyces naeslundii		1.80	0.42		99		
Capnocytophaga ochracea		0.37	0.05		1		
67550120	**Acceptable identification**						**67550120**
					GRA	CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		91.15	0.39	GAL 99	1	90	
Prevotella melaninogenica		4.09	0.12	BGL 90	1	1	
Actinomyces naeslundii		3.02	0.39		99	25	
67550121	**Low discrimination**						**67550121**
					GRA		
Actinomyces naeslundii		83.11	0.42	ESL 12	99		
Capnocytophaga ochracea		16.89	0.05	GAL 96	1		
67550122	**Good identification**						**67550122**
					GRA	CAT	
Capnocytophaga ochracea		96.19	0.56	GAL 99	1	90	
Prevotella oralis		2.27	0.27		1	1	
Actinomyces viscosus		0.87	0.22		99	76	
67550132	**Good identification**						**67550132**
Capnocytophaga ochracea		100.0	0.39	GAL 99			
67550162	**Good identification**						**67550162**
					GRA		
Capnocytophaga ochracea		99.42	0.39	GAL 99	1		
Actinomyces naeslundii		0.58	0.26	SOR 10	99		
67550522	**Good identification**						**67550522**
Capnocytophaga ochracea		100.0	0.43	GAL 99			
67551000	**Low discrimination**						**67551000**
					GRA	CAT	
Actinomyces naeslundii		57.14	0.55	ESL 12	99	25	
Capnocytophaga ochracea		22.73	0.23		1	90	
Fusobacterium mortiferum		10.44	0.31		1	5	
67551001	**Very good identification**						**67551001**
Actinomyces naeslundii		97.81	0.58	ESL 12			
Actinomyces viscosus		2.19	0.06				
67551002	**Low discrimination**						**67551002**
					GRA		
Actinomyces viscosus		67.00	0.48	ESL 1	99		
Capnocytophaga ochracea		24.03	0.39		1		
Actinomyces naeslundii		6.71	0.55		99		
67551003	**Low discrimination**						**67551003**
					GRA		
Actinomyces viscosus		68.41	0.40	ESL 1	99		
Actinomyces naeslundii		30.85	0.58		99		
Capnocytophaga ochracea		0.74	0.05		1		
67551011	**Good identification**						**67551011**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67551013	**Low discrimination**						**67551013**
Actinomyces naeslundii		70.14	0.36	ESL 12			
Actinomyces viscosus		29.86	0.06	MLZ 5			
67551020	**Low discrimination**						**67551020**
					GRA	CAT	
Capnocytophaga ochracea		66.54	0.39	LAC 99	1	90	
Actinomyces naeslundii		18.59	0.55	BGL 90	99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		7.19	0.26		99	1	
67551021	**Very good identification**						**67551021**
					GRA		
Actinomyces naeslundii		97.65	0.58	ESL 12	99		
Capnocytophaga ochracea		2.35	0.05		1		
67551022	**Low discrimination**						**67551022**
					GRA	CAT	
Capnocytophaga ochracea		83.90	0.56	LAC 99	1	90	
Prevotella oralis		12.52	0.41		1	1	
Actinomyces naeslundii		2.61	0.55		99	25	
67551023	**Low discrimination**						**67551023**
					GRA	CAT	
Actinomyces naeslundii		78.64	0.58	ESL 12	99	25	
Capnocytophaga ochracea		17.06	0.21		1	90	
Prevotella oralis		2.55	0.07		1	1	
67551031	**Good identification**						**67551031**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67551032	**Good identification**						**67551032**
					GRA	CAT	
Capnocytophaga ochracea		97.25	0.39	LAC 99	1	90	
Actinomyces naeslundii		1.43	0.33	MLZ 10	99	25	
Prevotella oralis		1.32	0.07		1	1	
67551033	**Low discrimination**						**67551033**
					GRA		
Actinomyces naeslundii		68.59	0.36	ESL 12	99		
Capnocytophaga ochracea		31.41	0.05	MLZ 5	1		
67551040	**Low discrimination**						**67551040**
					GRA	CAT	
Actinomyces naeslundii		74.46	0.42	ESL 12	99	25	
Capnocytophaga ochracea		18.65	0.07		1	90	
Actinomyces israelii		6.90	0.15		99	1	
67551041	**Good identification**						**67551041**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67551043	**Low discrimination**						**67551043**
Actinomyces naeslundii		88.74	0.45	ESL 12			
Actinomyces viscosus		11.27	0.06				
67551061	**Good identification**						**67551061**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67551062	**Acceptable identification**						**67551062**
					GRA	CAT	
Capnocytophaga ochracea		93.24	0.39	LAC 99	1	90	
Actinomyces naeslundii		4.60	0.42	SOR 10	99	25	
Prevotella oralis		1.27	0.07		1	1	
67551063	**Low discrimination**						**67551063**
					GRA		
Actinomyces naeslundii		87.98	0.45	ESL 12	99		
Capnocytophaga ochracea		12.02	0.05		1		
67551100	**Good identification**						**67551100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Capnocytophaga ochracea		90.47	0.57	NAG 90	1	90	
Actinomyces naeslundii		6.54	0.63	BGL 90	99	25	
Fusobacterium mortiferum		0.98	0.37		1	5	
67551101	**Low discrimination**						**67551101**
					GRA	SPO	
Actinomyces naeslundii		75.24	0.66	ESL 12	99	1	
Clostridium chauvoei		16.70	0.32		99	99	
Capnocytophaga ochracea		7.01	0.23		1	1	
67551102	**Good identification**						**67551102**
					GRA	CAT	
Capnocytophaga ochracea		93.34	0.73	NAG 90	1	90	
Actinomyces viscosus		4.67	0.52		99	76	
Prevotella oralis		1.04	0.40		1	1	
67551103	**Low discrimination**						**67551103**
					GRA		
Actinomyces viscosus		42.34	0.44	ESL 1	99		
Actinomyces naeslundii		30.56	0.66		99		
Capnocytophaga ochracea		25.62	0.39		1		
67551110	**Good identification**						**67551110**
					GRA	CAT	
Capnocytophaga ochracea		95.80	0.41	MLZ 10	1	90	
Actinomyces naeslundii		3.28	0.41	NAG 90	99	25	
Actinomyces israelii		0.55	0.19	BGL 90	99	1	
67551111	**Low discrimination**						**67551111**
					GRA		
Actinomyces naeslundii		83.57	0.44	ESL 12	99		
Capnocytophaga ochracea		16.43	0.07	MLZ 5	1		
67551112	**Very good identification**						**67551112**
					GRA	CAT	
Capnocytophaga ochracea		97.84	0.57	MLZ 10	1	90	
Actinomyces viscosus		0.45	0.18	NAG 90	99	76	
Bifidobacterium breve		0.42	0.27		99	1	
67551120	**Good identification**						**67551120**
					CAT		
Capnocytophaga ochracea		93.14	0.73	BGL 90	90		
Prevotella melaninogenica		4.18	0.47		1		
Prevotella oralis		1.04	0.40		1		
67551121	**Low discrimination**						**67551121**
					GRA	SPO	
Actinomyces naeslundii		50.86	0.66	ESL 12	99	1	
Capnocytophaga ochracea		42.64	0.39		1	1	
Clostridium chauvoei		3.76	0.24		99	99	
67551122	**Very good identification**						**67551122**
					CAT		
Capnocytophaga ochracea		97.28	0.90		90		
Prevotella oralis		2.30	0.62		1		
Bacteroides fragilis		0.18	0.34		99		
67551123	**Low discrimination**						**67551123**
					GRA	CAT	
Capnocytophaga ochracea		85.99	0.56	URE 1	1	90	
Actinomyces naeslundii		11.40	0.66		99	25	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella oralis		2.03	0.27		1	1	
67551130	**Very good identification**						**67551130**
					GRA	CAT	
Capnocytophaga ochracea		98.97	0.57	MLZ 10	1	90	
Prevotella melaninogenica		0.40	0.12	BGL 90	1	1	
Actinomyces naeslundii		0.38	0.41		99	25	
67551132	**Very good identification**						**67551132**
					GRA	CAT	
Capnocytophaga ochracea		99.61	0.73	MLZ 10	1	90	
Prevotella oralis		0.21	0.27		1	1	
Bifidobacterium breve		0.09	0.32		99	1	
67551133	**Acceptable identification**						**67551133**
					GRA		
Capnocytophaga ochracea		94.09	0.39	MLZ 10	1		
Actinomyces naeslundii		5.91	0.44	URE 1	99		
67551140	**Low discrimination**						**67551140**
					GRA	CAT	
Capnocytophaga ochracea		87.92	0.41	NAG 90	1	90	
Actinomyces naeslundii		10.09	0.50	SOR 10	99	25	
Actinomyces israelii		1.31	0.26	BGL 90	99	1	
67551141	**Acceptable identification**						**67551141**
					GRA		
Actinomyces naeslundii		94.46	0.53	ESL 12	99		
Capnocytophaga ochracea		5.54	0.07		1		
67551142	**Very good identification**						**67551142**
					GRA		
Capnocytophaga ochracea		97.52	0.57	NAG 90	1		
Actinomyces naeslundii		1.24	0.50	SOR 10	99		
Actinomyces viscosus		0.44	0.18		99		
67551143	**Low discrimination**						**67551143**
					GRA		
Actinomyces naeslundii		62.22	0.53	ESL 12	99		
Capnocytophaga ochracea		32.85	0.23		1		
Actinomyces viscosus		4.93	0.10		99		
67551152	**Good identification**						**67551152**
					GRA	CAT	
Capnocytophaga ochracea		95.06	0.41	MLZ 10	1	90	
Bifidobacterium breve		3.69	0.27	NAG 90	99	1	
Actinomyces naeslundii		0.57	0.28	SOR 10	99	25	
67551160	**Very good identification**						**67551160**
					GRA	CAT	
Capnocytophaga ochracea		96.83	0.57	SOR 10	1	90	
Clostridium perfringens		1.33	0.28	BGL 90	99	1	
Actinomyces naeslundii		1.24	0.50		99	25	
67551161	**Low discrimination**						**67551161**
					GRA		
Actinomyces naeslundii		65.45	0.53	ESL 12	99		
Capnocytophaga ochracea		34.55	0.23		1		
67551162	**Very good identification**						**67551162**
					GRA	CAT	
Capnocytophaga ochracea		99.46	0.73	SOR 10	1	90	
Prevotella oralis		0.21	0.27		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		0.14	0.50		99	25	
67551163	**Low discrimination**						**67551163**
					GRA		
Capnocytophaga ochracea		82.61	0.39	SOR 10	1		
Actinomyces naeslundii		17.39	0.53	URE 1	99		
67551170	**Good identification**						**67551170**
					GRA	CAT	
Capnocytophaga ochracea		98.73	0.41	MLZ 10	1	90	
Bifidobacterium breve		0.67	0.15	SOR 10	99	1	
Actinomyces naeslundii		0.60	0.28	BGL 90	99	25	
67551172	**Very good identification**						**67551172**
					GRA	CAT	
Capnocytophaga ochracea		99.17	0.57	MLZ 10	1	90	
Bifidobacterium breve		0.76	0.32	SOR 10	99	1	
Actinomyces naeslundii		0.07	0.28		99	25	
67551302	**Low discrimination**						**67551302**
					GRA	CAT	
Capnocytophaga ochracea		65.02	0.39	RHA 1	1	90	
Prevotella oralis		30.86	0.33	NAG 90	1	1	
Actinomyces viscosus		3.26	0.18		99	76	
67551320	**Low discrimination**						**67551320**
					CAT		
Capnocytophaga ochracea		64.85	0.39	RHA 1	90		
Prevotella oralis		30.78	0.33	BGL 90	1		
Prevotella melaninogenica		2.91	0.12		1		
67551322	**Low discrimination**						**67551322**
					CAT		
Prevotella oralis		49.71	0.55	NIT 1	1		
Capnocytophaga ochracea		49.61	0.56		90		
Parabacteroides distasonis		0.45	0.33		76		
67551332	**Low discrimination**						**67551332**
					CAT		
Capnocytophaga ochracea		88.09	0.39	RHA 1	90		
Prevotella oralis		8.02	0.21	MLZ 10	1		
Parabacteroides distasonis		3.89	0.28		76		
67551362	**Acceptable identification**						**67551362**
					CAT		
Capnocytophaga ochracea		91.65	0.39	RHA 1	90		
Prevotella oralis		8.35	0.21	SOR 10	1		
67551422	**Good identification**						**67551422**
					CAT		
Capnocytophaga ochracea		99.15	0.43	LAC 99	90		
Prevotella oralis		0.85	0.07		1		
67551500	**Good identification**						**67551500**
					GRA	CAT	
Capnocytophaga ochracea		97.38	0.44	NAG 90	1	90	
Actinomyces israelii		1.93	0.31	BGL 90	99	1	
Actinomyces naeslundii		0.40	0.29		99	25	
67551502	**Very good identification**						**67551502**
					GRA	CAT	
Capnocytophaga ochracea		98.11	0.61	NAG 90	1	90	
Bifidobacterium adolescentis		1.15	0.37		99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.32	0.34		99	1	
67551512	**Low discrimination**						**67551512**
					GRA	CAT	
Capnocytophaga ochracea		87.70	0.44	MLZ 10	1	90	
Bifidobacterium adolescentis		11.30	0.39	NAG 90	99	1	
Bifidobacterium longum subsp. longum		0.45	0.13		99	1	
67551520	**Very good identification**						**67551520**
					GRA	CAT	
Capnocytophaga ochracea		99.40	0.61	BGL 90	1	90	
Prevotella melaninogenica		0.26	0.12		1	1	
Bifidobacterium longum subsp. longum		0.13	0.20		99	1	
67551522	**Very good identification**						**67551522**
					CAT		
Capnocytophaga ochracea		99.81	0.77		90		
Prevotella oralis		0.14	0.27		1		
Parabacteroides distasonis		0.01	0.21		76		
67551523	**Good identification**						**67551523**
					GRA		
Capnocytophaga ochracea		99.25	0.43	URE 1	1		
Actinomyces naeslundii		0.75	0.32		99		
67551530	**Good identification**						**67551530**
					GRA	CAT	
Capnocytophaga ochracea		98.00	0.44	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		2.01	0.24	BGL 90	99	1	
67551532	**Very good identification**						**67551532**
					GRA	CAT	
Capnocytophaga ochracea		99.70	0.61	MLZ 10	1	90	
Bifidobacterium longum subsp. longum		0.23	0.24		99	1	
Parabacteroides distasonis		0.05	0.17		1	76	
67551542	**Good identification**						**67551542**
					GRA	CAT	
Capnocytophaga ochracea		97.60	0.44	NAG 90	1	90	
Bifidobacterium adolescentis		1.27	0.22	SOR 10	99	1	
Actinomyces israelii		1.13	0.27		99	1	
67551560	**Good identification**						**67551560**
Capnocytophaga ochracea		100.0	0.44	SOR 10			
67551562	**Very good identification**						**67551562**
Capnocytophaga ochracea		100.0	0.61	SOR 10			
67551572	**Good identification**						**67551572**
Capnocytophaga ochracea		100.0	0.44	MLZ 10			
67551722	**Acceptable identification**						**67551722**
					CAT		
Capnocytophaga ochracea		91.52	0.43	RHA 1	90		
Prevotella oralis		5.25	0.21		1		
Parabacteroides distasonis		2.90	0.29		76		
67552000	**Good identification**						**67552000**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67552001	**Good identification**						**67552001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	ESL 12		
67552003	**Low discrimination**					**67552003**
Actinomyces naeslundii		68.88	0.39	ESL 12		
Actinomyces viscosus		31.12	0.09	GAL 96		
67552020	**Good identification**					**67552020**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67552021	**Good identification**					**67552021**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67552022	**Good identification**					**67552022**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67552023	**Good identification**					**67552023**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67552100	**Low discrimination**					**67552100**
Actinomyces naeslundii		82.13	0.44	ESL 12		
Propionibacterium propionicum		17.87	0.23	GAL 96		
67552101	**Good identification**					**67552101**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67552103	**Low discrimination**					**67552103**
Actinomyces naeslundii		77.99	0.47	ESL 12		
Actinomyces viscosus		22.01	0.14	GAL 96		
67552120	**Low discrimination**					**67552120**
					GRA	
Actinomyces naeslundii		86.43	0.44	ESL 12	99	
Capnocytophaga ochracea		13.57	0.05	GAL 96	1	
67552121	**Good identification**					**67552121**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67552123	**Good identification**					**67552123**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67553000	**Low discrimination**					**67553000**
Actinomyces naeslundii		89.07	0.60	ESL 12		
Actinomyces israelii		10.93	0.35			
67553001	**Very good identification**					**67553001**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67553002	**Low discrimination**					**67553002**
Actinomyces naeslundii		79.71	0.60	ESL 12		
Actinomyces israelii		14.67	0.38			
Actinomyces viscosus		4.14	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67553003	**Very good identification**						**67553003**
Actinomyces naeslundii		98.86	0.63	ESL 12			
Actinomyces viscosus		1.14	0.06				
67553010	**Low discrimination**						**67553010**
Actinomyces naeslundii		74.17	0.38	ESL 12			
Actinomyces israelii		25.83	0.21	MLZ 5			
67553011	**Good identification**						**67553011**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67553012	**Low discrimination**						**67553012**
Actinomyces naeslundii		65.68	0.38	ESL 12			
Actinomyces israelii		34.32	0.24	MLZ 5			
67553013	**Good identification**						**67553013**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67553020	**Very good identification**						**67553020**
					GRA	CAT	
Actinomyces naeslundii		97.42	0.60	ESL 12	99	25	
Capnocytophaga ochracea		1.81	0.05		1	90	
Actinomyces israelii		0.76	0.15		99	1	
67553021	**Very good identification**						**67553021**
Actinomyces naeslundii		100.0	0.63	ESL 12			
67553022	**Low discrimination**						**67553022**
					GRA	CAT	
Actinomyces naeslundii		83.03	0.60	ESL 12	99	25	
Capnocytophaga ochracea		13.92	0.21		1	90	
Prevotella oralis		2.08	0.07		1	1	
67553023	**Very good identification**						**67553023**
Actinomyces naeslundii		100.0	0.63	ESL 12			
67553030	**Good identification**						**67553030**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67553031	**Good identification**						**67553031**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67553032	**Low discrimination**						**67553032**
					GRA		
Actinomyces naeslundii		73.86	0.38	ESL 12	99		
Capnocytophaga ochracea		26.14	0.05	MLZ 5	1		
67553033	**Good identification**						**67553033**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67553040	**Low discrimination**						**67553040**
Actinomyces naeslundii		78.72	0.47	ESL 12			
Actinomyces israelii		21.28	0.28				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67553041	**Good identification**						**67553041**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67553042	**Low discrimination**						**67553042**
Actinomyces naeslundii		71.15	0.47	ESL 12			
Actinomyces israelii		28.85	0.31				
67553043	**Good identification**						**67553043**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67553060	**Good identification**						**67553060**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67553061	**Good identification**						**67553061**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67553062	**Acceptable identification**						**67553062**
					GRA		
Actinomyces naeslundii		90.45	0.47	ESL 12	99		
Capnocytophaga ochracea		9.55	0.05		1		
67553063	**Good identification**						**67553063**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67553100	**Low discrimination**						**67553100**
					GRA	CAT	
Actinomyces naeslundii		79.59	0.68	ESL 12	99	25	
Actinomyces israelii		13.72	0.46		99	1	
Capnocytophaga ochracea		5.73	0.23		1	90	
67553101	**Very good identification**						**67553101**
Actinomyces naeslundii		100.0	0.71	ESL 12			
67553102	**Low discrimination**						**67553102**
					GRA	CAT	
Actinomyces naeslundii		49.83	0.68	ESL 12	99	25	
Capnocytophaga ochracea		32.28	0.39		1	90	
Actinomyces israelii		12.89	0.49		99	1	
67553103	**Very good identification**						**67553103**
					GRA		
Actinomyces naeslundii		98.69	0.71	ESL 12	99		
Actinomyces viscosus		0.71	0.10		99		
Capnocytophaga ochracea		0.43	0.05		1		
67553110	**Low discrimination**						**67553110**
					GRA	CAT	
Actinomyces naeslundii		58.85	0.46	ESL 12	99	25	
Actinomyces israelii		28.81	0.31	MLZ 5	99	1	
Capnocytophaga ochracea		8.94	0.07		1	90	
67553111	**Good identification**						**67553111**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67553113	**Good identification**						**67553113**
Actinomyces naeslundii		100.0	0.49	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67553120	**Low discrimination**						**67553120**
					GRA	CAT	
Actinomyces naeslundii		58.53	0.68	ESL 12	99	25	
Capnocytophaga ochracea		37.92	0.39		1	90	
Prevotella melaninogenica		1.70	0.12		1	1	
67553121	**Very good identification**						**67553121**
					GRA		
Actinomyces naeslundii		99.57	0.71	ESL 12	99		
Capnocytophaga ochracea		0.43	0.05		1		
67553122	**Low discrimination**						**67553122**
					GRA	CAT	
Capnocytophaga ochracea		82.48	0.56	MAN 1	1	90	
Actinomyces naeslundii		14.15	0.68		99	25	
Prevotella oralis		1.95	0.27		1	1	
67553123	**Very good identification**						**67553123**
					GRA		
Actinomyces naeslundii		96.22	0.71	ESL 12	99		
Capnocytophaga ochracea		3.78	0.21		1		
67553131	**Good identification**						**67553131**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67553132	**Low discrimination**						**67553132**
					GRA	CAT	
Capnocytophaga ochracea		85.51	0.39	MAN 1	1	90	
Bifidobacterium breve		7.22	0.32	MLZ 10	99	1	
Actinomyces naeslundii		6.95	0.46		99	25	
67553133	**Acceptable identification**						**67553133**
					GRA		
Actinomyces naeslundii		92.35	0.49	ESL 12	99		
Capnocytophaga ochracea		7.65	0.05	MLZ 5	1		
67553140	**Low discrimination**						**67553140**
Actinomyces naeslundii		66.94	0.55	ESL 12			
Actinomyces israelii		25.43	0.39				
Propionibacterium propionicum		3.44	0.23				
67553141	**Very good identification**						**67553141**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67553142	**Low discrimination**						**67553142**
					GRA	CAT	
Actinomyces naeslundii		44.86	0.55	ESL 12	99	25	
Actinomyces israelii		25.57	0.42		99	1	
Capnocytophaga ochracea		18.30	0.23		1	90	
67553143	**Very good identification**						**67553143**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67553151	**Good identification**						**67553151**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67553153	**Good identification**						**67553153**
Actinomyces naeslundii		100.0	0.36	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67553160	**Low discrimination**						**67553160**
					GRA	CAT	
Actinomyces naeslundii		68.36	0.55	ESL 12	99	25	
Capnocytophaga ochracea		27.89	0.23		1	90	
Bifidobacterium breve		2.10	0.15		99	1	
67553161	**Very good identification**						**67553161**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67553162	**Low discrimination**						**67553162**
					GRA	CAT	
Capnocytophaga ochracea		73.17	0.39	MAN 1	1	90	
Actinomyces naeslundii		19.93	0.55	SOR 10	99	25	
Bifidobacterium breve		6.18	0.32		99	1	
67553163	**Very good identification**						**67553163**
					GRA		
Actinomyces naeslundii		97.59	0.58	ESL 12	99		
Capnocytophaga ochracea		2.41	0.05		1		
67553171	**Good identification**						**67553171**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67553173	**Good identification**						**67553173**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67553301	**Good identification**						**67553301**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67553303	**Good identification**						**67553303**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67553321	**Good identification**						**67553321**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67553323	**Good identification**						**67553323**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67553402	**Acceptable identification**						**67553402**
Actinomyces israelii		93.71	0.37	SAL 92			
Actinomyces naeslundii		6.29	0.26	XYL 92			
67553500	**Low discrimination**						**67553500**
					GRA	CAT	
Actinomyces israelii		86.09	0.44	SAL 92	99	1	
Capnocytophaga ochracea		7.75	0.10	XYL 92	1	90	
Actinomyces naeslundii		6.16	0.34		99	25	
67553501	**Good identification**						**67553501**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67553502	**Low discrimination**						**67553502**
					GRA	CAT	
Actinomyces israelii		47.60	0.47	SAL 92	99	1	
Capnocytophaga ochracea		25.72	0.26	XYL 92	1	90	
Bifidobacterium adolescentis		24.41	0.36		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67553503	**Good identification**						**67553503**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67553512	**Low discrimination**						**67553512**
					GRA	CAT	
Bifidobacterium adolescentis		74.95	0.37	NIT 1	99	1	
Actinomyces israelii		17.87	0.33	XYL 87	99	1	
Capnocytophaga ochracea		7.18	0.10		1	90	
67553521	**Good identification**						**67553521**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67553522	**Good identification**						**67553522**
					GRA	CAT	
Capnocytophaga ochracea		97.76	0.43	MAN 1	1	90	
Actinomyces israelii		1.28	0.27		99	1	
Actinomyces naeslundii		0.96	0.34		99	25	
67553523	**Low discrimination**						**67553523**
					GRA		
Actinomyces naeslundii		59.31	0.37	ESL 12	99		
Capnocytophaga ochracea		40.69	0.09	ARA 1	1		
67553540	**Good identification**						**67553540**
Actinomyces israelii		100.0	0.37	SAL 92			
67553542	**Low discrimination**						**67553542**
					GRA	CAT	
Actinomyces israelii		75.91	0.40	SAL 92	99	1	
Bifidobacterium adolescentis		12.37	0.20	XYL 92	99	1	
Capnocytophaga ochracea		11.72	0.10		1	90	
67554122	**Low discrimination**						**67554122**
					CAT		
Capnocytophaga ochracea		57.25	0.45	GAL 99	90		
Bacteroides fragilis		42.08	0.34		99		
Prevotella oralis		0.67	0.12		1		
67555000	**Low discrimination**						**67555000**
					GRA	CAT	
Actinomyces israelii		81.03	0.41	SAL 92	99	1	
Capnocytophaga ochracea		16.67	0.13		1	90	
Fusobacterium mortiferum		2.30	0.12		1	5	
67555022	**Low discrimination**						**67555022**
					CAT		
Capnocytophaga ochracea		55.11	0.45	LAC 99	90		
Bacteroides fragilis		40.51	0.34		99		
Prevotella oralis		4.07	0.26		1		
67555042	**Low discrimination**						**67555042**
					GRA	CAT	
Actinomyces israelii		73.93	0.37	SAL 92	99	1	
Capnocytophaga ochracea		26.07	0.13		1	90	
67555100	**Low discrimination**						**67555100**
					GRA	CAT	
Capnocytophaga ochracea		80.73	0.47	NAG 90	1	90	
Actinomyces israelii		15.86	0.51	BGL 90	99	1	
Bacteroides fragilis		2.56	0.12		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67555102	**Low discrimination**						**67555102**
					GRA	CAT	
Capnocytophaga ochracea		89.63	0.63	NAG 90	1	90	
Bacteroides fragilis		5.99	0.34		1	99	
Actinomyces israelii		2.93	0.54		99	1	
67555112	**Low discrimination**						**67555112**
					GRA	CAT	
Capnocytophaga ochracea		87.64	0.47	MLZ 10	1	90	
Bifidobacterium adolescentis		7.97	0.39	NAG 90	99	1	
Actinomyces israelii		3.86	0.40		99	1	
67555120	**Low discrimination**						**67555120**
					CAT		
Capnocytophaga ochracea		73.58	0.63	BGL 90	90		
Bacteroides fragilis		25.62	0.46		99		
Prevotella oralis		0.41	0.24		1		
67555122	**Low discrimination**						**67555122**
					CAT		
Capnocytophaga ochracea		57.20	0.79		90		
Bacteroides fragilis		42.05	0.68		99		
Prevotella oralis		0.67	0.46		1		
67555123	**Low discrimination**						**67555123**
					CAT		
Capnocytophaga ochracea		57.07	0.45	URE 1	90		
Bacteroides fragilis		41.95	0.34		99		
Prevotella oralis		0.67	0.12		1		
67555130	**Good identification**						**67555130**
					GRA	CAT	
Capnocytophaga ochracea		96.76	0.47	MLZ 10	1	90	
Bacteroides fragilis		3.06	0.12	BGL 90	1	99	
Actinomyces israelii		0.18	0.16		99	1	
67555132	**Good identification**						**67555132**
Capnocytophaga ochracea		93.15	0.63	MLZ 10			
Bacteroides fragilis		6.23	0.34				
Parabacteroides distasonis		0.50	0.37				
67555142	**Low discrimination**						**67555142**
					GRA	CAT	
Capnocytophaga ochracea		88.51	0.47	NAG 90	1	90	
Actinomyces israelii		10.14	0.47	SOR 10	99	1	
Bifidobacterium adolescentis		0.81	0.22		99	1	
67555160	**Good identification**						**67555160**
					GRA	CAT	
Capnocytophaga ochracea		96.48	0.47	SOR 10	1	90	
Bacteroides fragilis		3.05	0.12	BGL 90	1	99	
Actinomyces israelii		0.47	0.23		99	1	
67555162	**Good identification**						**67555162**
					CAT		
Capnocytophaga ochracea		93.57	0.63	SOR 10	90		
Bacteroides fragilis		6.25	0.34		99		
Prevotella oralis		0.10	0.12		1		
67555172	**Good identification**						**67555172**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		99.40	0.47	MLZ 10			
Bacteroides fragilis		0.60	-.01	SOR 10			
67555322	**Low discrimination**						**67555322**
					CAT		
Capnocytophaga ochracea		38.95	0.45	RHA 1	90		
Bacteroides fragilis		28.63	0.34		99		
Prevotella oralis		19.29	0.40		1		
67555332	**Low discrimination**						**67555332**
Parabacteroides distasonis		57.39	0.45	NIT 1			
Capnocytophaga ochracea		36.09	0.29				
Clostridium clostridioforme		2.48	0.16				
67555400	**Good identification**						**67555400**
					GRA	CAT	
Actinomyces israelii		95.75	0.39	SAL 92	99	1	
Capnocytophaga ochracea		4.25	0.00		1	90	
67555402	**Low discrimination**						**67555402**
					GRA	CAT	
Actinomyces israelii		74.09	0.42	SAL 92	99	1	
Capnocytophaga ochracea		19.72	0.16		1	90	
Bifidobacterium adolescentis		6.19	0.17		99	1	
67555442	**Acceptable identification**						**67555442**
					GRA	CAT	
Actinomyces israelii		92.93	0.35	SAL 92	99	1	
Capnocytophaga ochracea		7.07	0.00		1	90	
67555502	**Low discrimination**						**67555502**
					GRA	CAT	
Capnocytophaga ochracea		67.55	0.50	NAG 90	1	90	
Bifidobacterium adolescentis		21.18	0.52		99	1	
Actinomyces israelii		10.25	0.52		99	1	
67555510	**Low discrimination**						**67555510**
					GRA	CAT	
Actinomyces israelii		35.90	0.35	SAL 92	99	1	
Capnocytophaga ochracea		29.31	0.18	MLZ 13	1	90	
Bifidobacterium longum subsp. longum		16.20	0.21		99	1	
67555512	**Low discrimination**						**67555512**
					GRA	CAT	
Bifidobacterium adolescentis		72.01	0.53	NIT 1	99	1	
Capnocytophaga ochracea		20.87	0.34		1	90	
Actinomyces israelii		4.26	0.38		99	1	
67555520	**Low discrimination**						**67555520**
					CAT		
Capnocytophaga ochracea		86.69	0.50	BGL 90	90		
Bacteroides vulgatus		9.30	0.28		3		
Bacteroides fragilis		1.73	0.12		99		
67555522	**Good identification**						**67555522**
Capnocytophaga ochracea		94.98	0.67				
Bacteroides fragilis		4.00	0.34				
Parabacteroides distasonis		0.36	0.38				
67555532	**Acceptable identification**						**67555532**
					GRA	CAT	

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea	94.04	0.50	MLZ 10	1	90	
Bifidobacterium longum subsp. longum	2.57	0.32		99	1	
Parabacteroides distasonis	1.75	0.33		1	76	
67555540 Low discrimination						**67555540**
				GRA	CAT	
Actinomyces israelii	61.04	0.42	SAL 92	99	1	
Capnocytophaga ochracea	19.15	0.18		1	90	
Mitsuokella multacida	18.43	0.21		1	1	
67555552 Low discrimination						**67555552**
				GRA	CAT	
Bifidobacterium adolescentis	69.12	0.37	NIT 1	99	1	
Capnocytophaga ochracea	18.01	0.18	SOR 11	1	90	
Actinomyces israelii	12.87	0.31		99	1	
67555562 Good identification						**67555562**
				GRA	CAT	
Capnocytophaga ochracea	99.25	0.50	SOR 10	1	90	
Bacteroides fragilis	0.38	-.01		1	99	
Actinomyces israelii	0.37	0.25		99	1	
67555712 Low discrimination						**67555712**
Clostridium clostridioforme	89.13	0.36	NIT 1			
Bifidobacterium adolescentis	5.60	0.19	SAL 88			
Actinomyces israelii	2.47	0.19	TRE 88			
67555720 Good identification						**67555720**
				CAT		
Bacteroides vulgatus	98.30	0.44	NIT 1	3		
Capnocytophaga ochracea	1.03	0.16	CEL 5	90		
Mitsuokella multacida	0.56	0.15		1		
67555722 Low discrimination						**67555722**
				CAT		
Parabacteroides distasonis	37.77	0.46	NIT 1	76		
Capnocytophaga ochracea	33.14	0.32		90		
Bacteroides uniformis	12.55	0.19		5		
67555732 Low discrimination						**67555732**
Clostridium clostridioforme	45.61	0.36	NIT 1			
Parabacteroides distasonis	41.27	0.41	SAL 88			
Capnocytophaga ochracea	7.47	0.16	TRE 88			
67557000 Good identification						**67557000**
Actinomyces israelii	99.29	0.53	SAL 92			
Actinomyces naeslundii	0.71	0.26				
67557002 Very good identification						**67557002**
Actinomyces israelii	99.53	0.56	SAL 92			
Actinomyces naeslundii	0.48	0.26				
67557010 Good identification						**67557010**
Actinomyces israelii	100.0	0.39	SAL 92			
67557012 Good identification						**67557012**
Actinomyces israelii	100.0	0.42	SAL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67557022	**Low discrimination**						**67557022**
					GRA	CAT	
Actinomyces israelii		61.99	0.36	SAL 92	99	1	
Capnocytophaga ochracea		19.23	0.11	NAG 6	1	90	
Bacteroides fragilis		14.14	-.01		1	99	
67557040	**Good identification**						**67557040**
Actinomyces israelii		100.0	0.46	SAL 92			
67557042	**Good identification**						**67557042**
Actinomyces israelii		100.0	0.49	SAL 92			
67557052	**Good identification**						**67557052**
Actinomyces israelii		100.0	0.35	SAL 92			
67557100	**Very good identification**						**67557100**
					GRA	CAT	
Actinomyces israelii		98.60	0.64	SAL 92	99	1	
Capnocytophaga ochracea		0.90	0.13		1	90	
Actinomyces naeslundii		0.50	0.34		99	25	
67557102	**Good identification**						**67557102**
					GRA	CAT	
Actinomyces israelii		91.08	0.67	SAL 92	99	1	
Capnocytophaga ochracea		4.96	0.29		1	90	
Bifidobacterium adolescentis		3.32	0.36		99	1	
67557110	**Good identification**						**67557110**
Actinomyces israelii		100.0	0.50	SAL 92			
67557112	**Low discrimination**						**67557112**
					GRA	CAT	
Actinomyces israelii		74.69	0.53	SAL 92	99	1	
Bifidobacterium adolescentis		22.28	0.37	MLZ 13	99	1	
Capnocytophaga ochracea		3.03	0.13		1	90	
67557121	**Good identification**						**67557121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67557122	**Low discrimination**						**67557122**
					GRA	CAT	
Capnocytophaga ochracea		53.08	0.45	MAN 1	1	90	
Bacteroides fragilis		39.02	0.34		1	99	
Actinomyces israelii		6.91	0.46		99	1	
67557123	**Low discrimination**						**67557123**
					GRA		
Actinomyces naeslundii		37.23	0.37	ESL 12	99		
Capnocytophaga ochracea		36.18	0.11	XYL 1	1		
Bacteroides fragilis		26.59	-.01		1		
67557140	**Very good identification**						**67557140**
Actinomyces israelii		100.0	0.57	SAL 92			
67557142	**Very good identification**						**67557142**
					GRA	CAT	
Actinomyces israelii		97.36	0.60	SAL 92	99	1	
Capnocytophaga ochracea		1.52	0.13		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		1.13	0.20		99	1	
67557150	**Good identification**						**67557150**
Actinomyces israelii		100.0	0.43	SAL 92			
67557152	**Acceptable identification**						**67557152**
Actinomyces israelii		91.34	0.46	SAL 92			
Bifidobacterium adolescentis		8.66	0.22	MLZ 13			
67557160	**Low discrimination**						**67557160**
					GRA	CAT	
Actinomyces israelii		73.23	0.36	SAL 92	99	1	
Capnocytophaga ochracea		26.77	0.13	NAG 6	1	90	
67557202	**Good identification**						**67557202**
Actinomyces israelii		100.0	0.37	SAL 92			
67557300	**Good identification**						**67557300**
Actinomyces israelii		100.0	0.45	SAL 92			
67557302	**Good identification**						**67557302**
Actinomyces israelii		100.0	0.48	SAL 92			
67557340	**Good identification**						**67557340**
Actinomyces israelii		100.0	0.38	SAL 92			
67557342	**Good identification**						**67557342**
Actinomyces israelii		100.0	0.41	SAL 92			
67557400	**Good identification**						**67557400**
Actinomyces israelii		100.0	0.52	SAL 92			
67557402	**Good identification**						**67557402**
Actinomyces israelii		98.81	0.55	SAL 92			
Bifidobacterium adolescentis		1.19	0.16				
67557410	**Good identification**						**67557410**
Actinomyces israelii		100.0	0.38	SAL 92			
67557412	**Acceptable identification**						**67557412**
Actinomyces israelii		91.03	0.41	SAL 92			
Bifidobacterium adolescentis		8.98	0.17	MLZ 13			
67557440	**Good identification**						**67557440**
Actinomyces israelii		100.0	0.45	SAL 92			
67557442	**Good identification**						**67557442**
Actinomyces israelii		100.0	0.48	SAL 92			
67557500	**Good identification**						**67557500**
					GRA		
Actinomyces israelii		94.62	0.62	SAL 92	99		
Mitsuokella multacida		4.58	0.27		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		0.43	0.16		99		
67557502	**Low discrimination**						**67557502**
					GRA	CAT	
Actinomyces israelii		76.12	0.65	SAL 92	99	1	
Bifidobacterium adolescentis		22.71	0.50		99	1	
Capnocytophaga ochracea		0.89	0.16		1	90	
67557510	**Low discrimination**						**67557510**
					GRA		
Actinomyces israelii		87.35	0.48	SAL 92	99		
Mitsuokella multacida		9.42	0.19	MLZ 13	1		
Bifidobacterium adolescentis		3.23	0.17		99		
67557512	**Low discrimination**						**67557512**
Bifidobacterium adolescentis		70.37	0.52	NIT 1			
Actinomyces israelii		28.84	0.51				
Bifidobacterium dentium		0.32	-.07				
67557520	**Low discrimination**						**67557520**
					GRA	CAT	
Actinomyces israelii		66.89	0.42	SAL 92	99	1	
Capnocytophaga ochracea		18.46	0.16	NAG 6	1	90	
Mitsuokella multacida		12.67	0.17		1	1	
67557540	**Low discrimination**						**67557540**
					GRA		
Actinomyces israelii		88.94	0.55	SAL 92	99		
Mitsuokella multacida		11.06	0.27		1		
67557542	**Good identification**						**67557542**
					GRA		
Actinomyces israelii		90.30	0.58	SAL 92	99		
Bifidobacterium adolescentis		8.56	0.34		99		
Mitsuokella multacida		0.83	0.11		1		
67557550	**Low discrimination**						**67557550**
					COC	GRA	
Actinomyces israelii		75.81	0.41	SAL 92	1	99	
Mitsuokella multacida		21.03	0.19	MLZ 13	1	1	
Blautia producta		3.16	-.03		99	99	
67557552	**Low discrimination**						**67557552**
					GRA		
Actinomyces israelii		55.68	0.44	SAL 92	99		
Bifidobacterium adolescentis		43.18	0.36	MLZ 13	99		
Mitsuokella multacida		1.14	0.03		1		
67557562	**Low discrimination**						**67557562**
					GRA	CAT	
Actinomyces israelii		67.88	0.38	SAL 92	99	1	
Capnocytophaga ochracea		32.12	0.16	NAG 6	1	90	
67557602	**Good identification**						**67557602**
Actinomyces israelii		100.0	0.36	SAL 92			
67557702	**Low discrimination**						**67557702**
					GRA		
Actinomyces israelii		88.62	0.46	SAL 92	99		
Mitsuokella multacida		7.83	0.15	RHA 7	1		
Bifidobacterium adolescentis		3.55	0.16		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67557742	**Low discrimination**						**67557742**
					GRA		
Actinomyces israelii		81.48	0.39	SAL 92	99		
Mitsuokella multacida		18.52	0.15	RHA 7	1		
67561000	**Good identification**						**67561000**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67561001	**Good identification**						**67561001**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67561002	**Low discrimination**						**67561002**
Actinomyces naeslundii		72.10	0.38	ESL 12			
Actinomyces viscosus		27.90	0.07	FRU 95			
67561003	**Acceptable identification**						**67561003**
Actinomyces naeslundii		92.08	0.41	ESL 12			
Actinomyces viscosus		7.92	-.01	FRU 95			
67561020	**Good identification**						**67561020**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67561021	**Good identification**						**67561021**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67561022	**Good identification**						**67561022**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67561023	**Good identification**						**67561023**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67561100	**Good identification**						**67561100**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67561101	**Acceptable identification**						**67561101**
					SPO		
Actinomyces naeslundii		90.63	0.49	ESL 12	1		
Clostridium chauvoei		9.37	0.09	FRU 95	99		
67561102	**Low discrimination**						**67561102**
					GRA		
Actinomyces naeslundii		73.20	0.46	ESL 12	99		
Actinomyces viscosus		17.70	0.11	FRU 95	99		
Capnocytophaga ochracea		9.10	0.05		1		
67561103	**Acceptable identification**						**67561103**
Actinomyces naeslundii		94.90	0.49	ESL 12			
Actinomyces viscosus		5.10	0.03	FRU 95			
67561120	**Low discrimination**						**67561120**
					GRA	CAT	
Actinomyces naeslundii		73.07	0.46	ESL 12	99	25	
Clostridium perfringens		13.36	0.11	FRU 95	99	1	
Capnocytophaga ochracea		9.09	0.05		1	90	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67561121	**Good identification**					**67561121**
					SPO	
Actinomyces naeslundii		96.67	0.49	ESL 12	1	
Clostridium chauvoei		3.33	0.01	FRU 95	99	
67561123	**Good identification**					**67561123**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67561141	**Good identification**					**67561141**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67561143	**Good identification**					**67561143**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67561161	**Good identification**					**67561161**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67561163	**Good identification**					**67561163**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67563000	**Good identification**					**67563000**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67563001	**Good identification**					**67563001**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67563002	**Acceptable identification**					**67563002**
Actinomyces naeslundii		93.59	0.43	ESL 12		
Actinomyces israelii		6.41	0.14	FRU 95		
67563003	**Good identification**					**67563003**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67563020	**Good identification**					**67563020**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67563021	**Good identification**					**67563021**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67563022	**Good identification**					**67563022**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67563023	**Good identification**					**67563023**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67563100	**Acceptable identification**					**67563100**
Actinomyces naeslundii		93.97	0.51	ESL 12		
Actinomyces israelii		6.03	0.21	FRU 95		
67563101	**Good identification**					**67563101**
Actinomyces naeslundii		100.0	0.54	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67563102	**Acceptable identification**					**67563102**
Actinomyces naeslundii		91.23	0.51	ESL 12		
Actinomyces israelii		8.78	0.24	FRU 95		
67563103	**Good identification**					**67563103**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67563120	**Good identification**					**67563120**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67563121	**Good identification**					**67563121**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67563122	**Good identification**					**67563122**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67563123	**Good identification**					**67563123**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67563140	**Low discrimination**					**67563140**
Actinomyces naeslundii		87.62	0.38	ESL 12		
Actinomyces israelii		12.38	0.14	FRU 95		
67563141	**Good identification**					**67563141**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67563142	**Low discrimination**					**67563142**
Actinomyces naeslundii		82.51	0.38	ESL 12		
Actinomyces israelii		17.49	0.17	FRU 95		
67563143	**Good identification**					**67563143**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67563160	**Good identification**					**67563160**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67563161	**Good identification**					**67563161**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67563162	**Good identification**					**67563162**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67563163	**Good identification**					**67563163**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67567100	**Good identification**					**67567100**
Actinomyces israelii		100.0	0.39	SAL 92		
67567102	**Good identification**					**67567102**
Actinomyces israelii		100.0	0.42	SAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67567142	**Good identification**						**67567142**
Actinomyces israelii		100.0	0.35	SAL 92			
67567502	**Low discrimination**						**67567502**
					GRA		
Actinomyces israelii		87.17	0.41	SAL 92	99		
Mitsuokella multacida		8.22	0.11	FRU 99	1		
Bifidobacterium adolescentis		4.61	0.13		99		
67570000	**Low discrimination**						**67570000**
					GRA		
Actinomyces naeslundii		62.39	0.36	ESL 12	99		
Fusobacterium mortiferum		28.06	0.19	GAL 96	1		
Actinomyces viscosus		9.55	-.02		99		
67570001	**Good identification**						**67570001**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67570020	**Good identification**						**67570020**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67570021	**Good identification**						**67570021**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67570022	**Low discrimination**						**67570022**
Actinomyces naeslundii		86.73	0.36	ESL 12			
Actinomyces viscosus		13.27	-.02	GAL 96			
67570023	**Good identification**						**67570023**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67570100	**Low discrimination**						**67570100**
					GRA		
Actinomyces naeslundii		68.30	0.44	ESL 12	99		
Fusobacterium mortiferum		25.18	0.25	GAL 96	1		
Actinomyces viscosus		6.53	0.03		99		
67570101	**Good identification**						**67570101**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67570102	**Low discrimination**						**67570102**
					GRA		
Actinomyces viscosus		89.11	0.37	ESL 1	99		
Actinomyces naeslundii		9.41	0.44	TRE 7	99		
Capnocytophaga ochracea		1.48	0.05		1		
67570120	**Low discrimination**						**67570120**
					GRA	CAT	
Actinomyces naeslundii		72.04	0.44	ESL 12	99	25	
Clostridium perfringens		16.64	0.11	GAL 96	99	1	
Capnocytophaga ochracea		11.31	0.05		1	90	
67570121	**Good identification**						**67570121**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67570123	**Good identification**						**67570123**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.47	ESL 12			
67571000	**Low discrimination**						**67571000**
					GRA		
Actinomyces naeslundii		87.57	0.60	ESL 12	99		
Actinomyces israelii		7.13	0.32		99		
Fusobacterium mortiferum		2.46	0.22		1		
67571001	**Very good identification**						**67571001**
Actinomyces naeslundii		100.0	0.63	ESL 12			
67571002	**Low discrimination**						**67571002**
Actinomyces naeslundii		64.51	0.60	ESL 12			
Actinomyces viscosus		24.97	0.29				
Actinomyces israelii		7.88	0.35				
67571003	**Good identification**						**67571003**
Actinomyces naeslundii		92.08	0.63	ESL 12			
Actinomyces viscosus		7.92	0.21				
67571010	**Low discrimination**						**67571010**
Actinomyces naeslundii		73.31	0.38	ESL 12			
Actinomyces israelii		16.95	0.18	MLZ 5			
Lactobacillus jensenii		9.74	0.02				
67571011	**Good identification**						**67571011**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67571013	**Good identification**						**67571013**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67571020	**Low discrimination**						**67571020**
					GRA	CAT	
Actinomyces naeslundii		71.59	0.60	ESL 12	99	25	
Clostridium perfringens		26.48	0.30		99	1	
Capnocytophaga ochracea		1.33	0.05		1	90	
67571021	**Very good identification**						**67571021**
Actinomyces naeslundii		100.0	0.63	ESL 12			
67571022	**Low discrimination**						**67571022**
					GRA	CAT	
Actinomyces naeslundii		66.33	0.60	ESL 12	99	25	
Clostridium perfringens		11.55	0.25		99	1	
Capnocytophaga ochracea		11.12	0.21		1	90	
67571023	**Very good identification**						**67571023**
Actinomyces naeslundii		100.0	0.63	ESL 12			
67571030	**Good identification**						**67571030**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67571031	**Good identification**						**67571031**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67571032	**Low discrimination**						**67571032**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces naeslundii		65.58	0.38	ESL 12	99	25	
Capnocytophaga ochracea		23.21	0.05	MLZ 5	1	90	
Leptotrichia buccalis		11.22	-.09		1	1	
67571033	**Good identification**						**67571033**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67571040	**Low discrimination**						**67571040**
Actinomyces naeslundii		84.78	0.47	ESL 12			
Actinomyces israelii		15.22	0.25				
67571041	**Good identification**						**67571041**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67571042	**Low discrimination**						**67571042**
Actinomyces naeslundii		78.79	0.47	ESL 12			
Actinomyces israelii		21.21	0.28				
67571043	**Good identification**						**67571043**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67571060	**Low discrimination**						**67571060**
Actinomyces naeslundii		71.47	0.47	ESL 12			
Clostridium perfringens		28.53	0.18				
67571061	**Good identification**						**67571061**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67571062	**Low discrimination**						**67571062**
					GRA	CAT	
Actinomyces naeslundii		77.32	0.47	ESL 12	99	25	
Clostridium perfringens		14.53	0.12		99	1	
Capnocytophaga ochracea		8.16	0.05		1	90	
67571063	**Good identification**						**67571063**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67571100	**Low discrimination**						**67571100**
					GRA	CAT	
Actinomyces naeslundii		76.15	0.68	ESL 12	99	25	
Actinomyces israelii		8.72	0.43		99	1	
Capnocytophaga ochracea		5.48	0.23		1	90	
67571101	**Very good identification**						**67571101**
					SPO		
Actinomyces naeslundii		98.75	0.71	ESL 12	1		
Clostridium chauvoei		1.25	0.15		99		
67571102	**Low discrimination**						**67571102**
					GRA		
Actinomyces naeslundii		45.15	0.68	ESL 12	99		
Capnocytophaga ochracea		29.25	0.39		1		
Actinomyces viscosus		10.91	0.33		99		
67571103	**Good identification**						**67571103**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		94.31	0.71	ESL 12	99		
Actinomyces viscosus		5.07	0.25		99		
Capnocytophaga ochracea		0.41	0.05		1		
67571110	**Low discrimination**						**67571110**
					GRA		
Actinomyces naeslundii		49.55	0.46	ESL 12	99		
Leptotrichia buccalis		26.82	0.07	MLZ 5	1		
Actinomyces israelii		16.10	0.28		99		
67571111	**Good identification**						**67571111**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67571112	**Acceptable identification**						**67571112**
					GRA	CAT	
Leptotrichia buccalis		93.71	0.41	NIT 10	1	1	
Capnocytophaga ochracea		2.39	0.23	SAL 99	1	90	
Actinomyces naeslundii		1.75	0.46		99	25	
67571113	**Low discrimination**						**67571113**
					GRA		
Actinomyces naeslundii		73.49	0.49	ESL 12	99		
Leptotrichia buccalis		26.51	0.07	MLZ 5	1		
67571120	**Low discrimination**						**67571120**
					GRA	CAT	
Actinomyces naeslundii		36.79	0.68	ESL 12	99	25	
Clostridium perfringens		35.06	0.45		99	1	
Capnocytophaga ochracea		23.84	0.39		1	90	
67571121	**Very good identification**						**67571121**
					GRA	CAT	
Actinomyces naeslundii		98.52	0.71	ESL 12	99	25	
Clostridium perfringens		0.63	0.11		99	1	
Capnocytophaga ochracea		0.43	0.05		1	90	
67571122	**Low discrimination**						**67571122**
					GRA	CAT	
Capnocytophaga ochracea		69.81	0.56	TRE 1	1	90	
Actinomyces naeslundii		11.97	0.68		99	25	
Prevotella oralis		10.42	0.41		1	1	
67571123	**Very good identification**						**67571123**
					GRA	CAT	
Actinomyces naeslundii		95.41	0.71	ESL 12	99	25	
Capnocytophaga ochracea		3.75	0.21		1	90	
Prevotella oralis		0.56	0.07		1	1	
67571131	**Good identification**						**67571131**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67571132	**Low discrimination**						**67571132**
					GRA	CAT	
Capnocytophaga ochracea		58.52	0.39	TRE 1	1	90	
Leptotrichia buccalis		28.30	0.25	MLZ 10	1	1	
Actinomyces naeslundii		4.75	0.46		99	25	
67571133	**Low discrimination**						**67571133**
					GRA	CAT	
Actinomyces naeslundii		89.05	0.49	ESL 12	99	25	
Capnocytophaga ochracea		7.38	0.05	MLZ 5	1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		3.57	-.09		1	1	
67571140	**Low discrimination**						**67571140**
					GRA	CAT	
Actinomyces naeslundii		75.22	0.55	ESL 12	99	25	
Actinomyces israelii		18.97	0.36		99	1	
Capnocytophaga ochracea		3.41	0.07		1	90	
67571141	**Very good identification**						**67571141**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67571142	**Low discrimination**						**67571142**
					GRA	CAT	
Actinomyces naeslundii		51.78	0.55	ESL 12	99	25	
Capnocytophaga ochracea		21.12	0.23		1	90	
Actinomyces israelii		19.59	0.39		99	1	
67571143	**Very good identification**						**67571143**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67571151	**Good identification**						**67571151**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67571153	**Good identification**						**67571153**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67571161	**Very good identification**						**67571161**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67571162	**Low discrimination**						**67571162**
					GRA	CAT	
Capnocytophaga ochracea		67.54	0.39	TRE 1	1	90	
Actinomyces naeslundii		18.40	0.55	SOR 10	99	25	
Clostridium perfringens		8.91	0.27		99	1	
67571163	**Very good identification**						**67571163**
					GRA		
Actinomyces naeslundii		97.59	0.58	ESL 12	99		
Capnocytophaga ochracea		2.41	0.05		1		
67571171	**Good identification**						**67571171**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67571173	**Good identification**						**67571173**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67571301	**Good identification**						**67571301**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67571303	**Good identification**						**67571303**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67571321	**Good identification**						**67571321**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67571323	**Low discrimination**						**67571323**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		70.92	0.37	ESL 12	99		
Prevotella oralis		17.63	0.01	RHA 1	1		
Parabacteroides distasonis		11.45	0.10		1		
67571332	**Low discrimination**						**67571332**
Parabacteroides distasonis		85.81	0.40	NIT 1			
Clostridium clostridioforme		5.32	0.13	XYL 90			
Capnocytophaga ochracea		4.31	0.05				
67571500	**Low discrimination**						**67571500**
					GRA	SPO	
Actinomyces israelii		57.09	0.41	SAL 92	99	1	
Mitsuokella multacida		20.50	0.21	XYL 92	1	1	
Clostridium butyricum		8.52	0.06		99	99	
67571501	**Low discrimination**						**67571501**
					GRA		
Actinomyces naeslundii		80.22	0.37	ESL 12	99		
Mitsuokella multacida		19.78	0.04	ARA 1	1		
67571502	**Low discrimination**						**67571502**
					GRA	CAT	
Actinomyces israelii		38.10	0.44	SAL 92	99	1	
Capnocytophaga ochracea		31.01	0.26	XYL 92	1	90	
Bifidobacterium adolescentis		23.98	0.34		99	1	
67571503	**Good identification**						**67571503**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67571512	**Low discrimination**						**67571512**
					GRA		
Bifidobacterium adolescentis		57.63	0.36	NIT 1	99		
Leptotrichia buccalis		15.19	0.07	XYL 87	1		
Actinomyces israelii		11.20	0.30		99		
67571521	**Good identification**						**67571521**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67571522	**Acceptable identification**						**67571522**
					GRA		
Capnocytophaga ochracea		92.01	0.43	TRE 1	1		
Parabacteroides distasonis		4.42	0.33		1		
Actinomyces naeslundii		0.90	0.34		99		
67571523	**Low discrimination**						**67571523**
					GRA		
Actinomyces naeslundii		59.31	0.37	ESL 12	99		
Capnocytophaga ochracea		40.69	0.09	ARA 1	1		
67571542	**Low discrimination**						**67571542**
					GRA	CAT	
Actinomyces israelii		66.63	0.37	SAL 92	99	1	
Capnocytophaga ochracea		15.49	0.10	XYL 92	1	90	
Bifidobacterium adolescentis		13.32	0.19		99	1	
67571722	**Low discrimination**						**67571722**
Parabacteroides distasonis		71.95	0.41	NIT 1			
Clostridium clostridioforme		13.47	0.23	XYL 90			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		7.72	0.08			
67572000	**Good identification**					**67572000**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67572001	**Good identification**					**67572001**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67572002	**Low discrimination**					**67572002**
Actinomyces naeslundii		85.81	0.41	ESL 12		
Actinomyces israelii		7.42	0.14	GAL 96		
Actinomyces viscosus		6.77	-.02			
67572003	**Good identification**					**67572003**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67572020	**Good identification**					**67572020**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67572021	**Good identification**					**67572021**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67572022	**Good identification**					**67572022**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67572023	**Good identification**					**67572023**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67572100	**Low discrimination**					**67572100**
Actinomyces naeslundii		85.27	0.49	ESL 12		
Propionibacterium propionicum		7.82	0.21	GAL 96		
Actinomyces israelii		6.91	0.21			
67572101	**Good identification**					**67572101**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67572102	**Low discrimination**					**67572102**
Actinomyces naeslundii		76.43	0.49	ESL 12		
Propionibacterium propionicum		10.51	0.24	GAL 96		
Actinomyces israelii		9.29	0.24			
67572103	**Good identification**					**67572103**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67572120	**Good identification**					**67572120**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67572121	**Good identification**					**67572121**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67572122	**Good identification**					**67572122**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.49	ESL 12		
67572123	**Good identification**					**67572123**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67572140	**Low discrimination**					**67572140**
Actinomyces naeslundii		58.89	0.36	ESL 12		
Propionibacterium propionicum		30.60	0.21	GAL 96		
Actinomyces israelii		10.51	0.14			
67572141	**Good identification**					**67572141**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67572142	**Low discrimination**					**67572142**
Actinomyces naeslundii		48.85	0.36	ESL 12		
Propionibacterium propionicum		38.07	0.24	GAL 96		
Actinomyces israelii		13.08	0.17			
67572143	**Good identification**					**67572143**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67572160	**Good identification**					**67572160**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67572161	**Good identification**					**67572161**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67572162	**Good identification**					**67572162**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67572163	**Good identification**					**67572163**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67573000	**Low discrimination**					**67573000**
Actinomyces naeslundii		80.79	0.65	ESL 12		
Actinomyces israelii		19.21	0.45			
67573001	**Very good identification**					**67573001**
Actinomyces naeslundii		100.0	0.68	ESL 12		
67573002	**Low discrimination**					**67573002**
Actinomyces naeslundii		73.71	0.65	ESL 12		
Actinomyces israelii		26.29	0.48			
67573003	**Very good identification**					**67573003**
Actinomyces naeslundii		99.76	0.68	ESL 12		
Actinomyces israelii		0.24	0.14			
67573010	**Low discrimination**					**67573010**
Actinomyces naeslundii		59.70	0.43	ESL 12		
Actinomyces israelii		40.30	0.31	MLZ 5		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67573011	**Good identification**					**67573011**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67573013	**Good identification**					**67573013**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67573020	**Very good identification**					**67573020**
Actinomyces naeslundii		98.51	0.65	ESL 12		
Actinomyces israelii		1.50	0.25			
67573021	**Very good identification**					**67573021**
Actinomyces naeslundii		100.0	0.68	ESL 12		
67573022	**Very good identification**					**67573022**
Actinomyces naeslundii		97.77	0.65	ESL 12		
Actinomyces israelii		2.23	0.28			
67573023	**Very good identification**					**67573023**
Actinomyces naeslundii		100.0	0.68	ESL 12		
67573030	**Good identification**					**67573030**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67573031	**Good identification**					**67573031**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67573032	**Acceptable identification**					**67573032**
Actinomyces naeslundii		93.93	0.43	ESL 12		
Actinomyces israelii		6.07	0.14	MLZ 5		
67573033	**Good identification**					**67573033**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67573040	**Low discrimination**					**67573040**
Actinomyces naeslundii		65.62	0.52	ESL 12		
Actinomyces israelii		34.38	0.38			
67573041	**Good identification**					**67573041**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67573042	**Low discrimination**					**67573042**
Actinomyces naeslundii		55.99	0.52	ESL 12		
Actinomyces israelii		44.01	0.41			
67573043	**Good identification**					**67573043**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67573060	**Good identification**					**67573060**
Actinomyces naeslundii		96.76	0.52	ESL 12		
Actinomyces israelii		3.24	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67573061	**Good identification**					**67573061**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67573062	**Good identification**					**67573062**
Actinomyces naeslundii		95.22	0.52	ESL 12		
Actinomyces israelii		4.78	0.21			
67573063	**Good identification**					**67573063**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67573100	**Low discrimination**					**67573100**
Actinomyces naeslundii		74.74	0.73	ESL 12		
Actinomyces israelii		24.97	0.55			
Propionibacterium propionicum		0.29	0.21			
67573101	**Very good identification**					**67573101**
Actinomyces naeslundii		99.78	0.76	ESL 12		
Actinomyces israelii		0.22	0.21			
67573102	**Low discrimination**					**67573102**
Actinomyces naeslundii		65.46	0.73	ESL 12		
Actinomyces israelii		32.81	0.58			
Bifidobacterium breve		0.69	0.24			
67573103	**Very good identification**					**67573103**
Actinomyces naeslundii		99.66	0.76	ESL 12		
Actinomyces israelii		0.34	0.24			
67573110	**Low discrimination**					**67573110**
Actinomyces naeslundii		51.32	0.51	ESL 12		
Actinomyces israelii		48.68	0.41	MLZ 5		
67573111	**Good identification**					**67573111**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67573112	**Low discrimination**					**67573112**
					GRA	
Actinomyces israelii		46.94	0.44	SAL 92	99	
Actinomyces naeslundii		32.98	0.51	XYL 92	99	
Leptotrichia buccalis		9.20	0.07	MLZ 13	1	
67573113	**Good identification**					**67573113**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67573120	**Good identification**					**67573120**
					SPO	
Actinomyces naeslundii		94.87	0.73	ESL 12	1	
Clostridium ramosum		2.14	0.19		92	
Actinomyces israelii		2.02	0.35		1	
67573121	**Very good identification**					**67573121**
Actinomyces naeslundii		100.0	0.76	ESL 12		
67573122	**Low discrimination**					**67573122**

Code Taxon		ID score	T index	tests against	Differentiation tests SPO	Code
Actinomyces naeslundii		84.89	0.73	ESL 12	1	
Clostridium ramosum		7.64	0.30		92	
Actinomyces israelii		2.72	0.38		1	
67573123	**Very good identification**					**67573123**
Actinomyces naeslundii		100.0	0.76	ESL 12		
67573130	**Acceptable identification**					**67573130**
Actinomyces naeslundii		91.25	0.51	ESL 12		
Actinomyces israelii		5.53	0.21	MLZ 5		
Bifidobacterium breve		3.22	0.12			
67573131	**Good identification**					**67573131**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67573132	**Low discrimination**					**67573132**
Actinomyces naeslundii		64.82	0.51	ESL 12		
Bifidobacterium breve		23.13	0.29	MLZ 5		
Actinomyces israelii		5.89	0.24			
67573133	**Good identification**					**67573133**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67573140	**Low discrimination**					**67573140**
Actinomyces naeslundii		56.88	0.60	ESL 12		
Actinomyces israelii		41.89	0.48			
Propionibacterium propionicum		1.23	0.21			
67573141	**Very good identification**					**67573141**
Actinomyces naeslundii		99.51	0.63	ESL 12		
Actinomyces israelii		0.49	0.14			
67573142	**Low discrimination**					**67573142**
Actinomyces israelii		50.28	0.51	SAL 92		
Actinomyces naeslundii		45.52	0.60	XYL 92		
Bifidobacterium breve		2.73	0.24			
67573143	**Very good identification**					**67573143**
Actinomyces naeslundii		99.26	0.63	ESL 12		
Actinomyces israelii		0.74	0.17			
67573151	**Good identification**					**67573151**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67573152	**Low discrimination**					**67573152**
Actinomyces israelii		59.47	0.37	SAL 92		
Bifidobacterium breve		21.57	0.24	XYL 92		
Actinomyces naeslundii		18.96	0.38	MLZ 13		
67573153	**Good identification**					**67573153**
Actinomyces naeslundii		100.0	0.41	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67573160	**Good identification**					**67573160**
Actinomyces naeslundii		94.56	0.60	ESL 12		
Actinomyces israelii		4.44	0.28			
Bifidobacterium breve		1.00	0.12			
67573161	**Very good identification**					**67573161**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67573162	**Low discrimination**					**67573162**
Actinomyces naeslundii		83.61	0.60	ESL 12		
Bifidobacterium breve		8.90	0.29			
Actinomyces israelii		5.89	0.31			
67573163	**Very good identification**					**67573163**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67573170	**Low discrimination**					**67573170**
Actinomyces naeslundii		75.00	0.38	ESL 12		
Bifidobacterium breve		15.00	0.12	MLZ 5		
Actinomyces israelii		10.01	0.14			
67573171	**Good identification**					**67573171**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67573173	**Good identification**					**67573173**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67573300	**Low discrimination**					**67573300**
Actinomyces israelii		71.35	0.36	SAL 92		
Actinomyces naeslundii		28.65	0.39	XYL 92		
67573301	**Good identification**					**67573301**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67573302	**Low discrimination**					**67573302**
					SPO	
Actinomyces israelii		76.94	0.39	SAL 92	1	
Actinomyces naeslundii		20.60	0.39	XYL 92	1	
Clostridium ramosum		2.46	-.03	RHA 7	92	
67573303	**Good identification**					**67573303**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67573321	**Good identification**					**67573321**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67573323	**Acceptable identification**					**67573323**
					SPO	
Actinomyces naeslundii		92.63	0.42	ESL 12	1	
Clostridium ramosum		7.37	-.03	RHA 1	92	
67573400	**Good identification**					**67573400**
Actinomyces israelii		95.06	0.44	SAL 92		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		4.94	0.31	XYL 92			
67573402	**Good identification**						**67573402**
Actinomyces israelii		96.65	0.47	SAL 92			
Actinomyces naeslundii		3.35	0.31	XYL 92			
67573440	**Good identification**						**67573440**
Actinomyces israelii		100.0	0.37	SAL 92			
67573442	**Good identification**						**67573442**
Actinomyces israelii		100.0	0.40	SAL 92			
67573500	**Low discrimination**						**67573500**
					GRA		
Actinomyces israelii		84.40	0.54	SAL 92	99		
Mitsuokella multacida		12.48	0.27	XYL 92	1		
Actinomyces naeslundii		3.12	0.39		99		
67573501	**Low discrimination**						**67573501**
					GRA		
Actinomyces naeslundii		67.63	0.42	ESL 12	99		
Mitsuokella multacida		20.05	0.11	ARA 1	1		
Actinomyces israelii		12.33	0.20		99		
67573502	**Low discrimination**						**67573502**
Actinomyces israelii		88.78	0.57	SAL 92			
Bifidobacterium adolescentis		8.07	0.33	XYL 92			
Actinomyces naeslundii		2.19	0.39				
67573503	**Low discrimination**						**67573503**
Actinomyces naeslundii		78.53	0.42	ESL 12			
Actinomyces israelii		21.47	0.23	ARA 1			
67573510	**Low discrimination**						**67573510**
					GRA		
Actinomyces israelii		75.20	0.40	SAL 92	99		
Mitsuokella multacida		24.80	0.19	XYL 92	1		
67573512	**Low discrimination**						**67573512**
Actinomyces israelii		55.52	0.43	SAL 92			
Bifidobacterium adolescentis		41.26	0.34	XYL 92			
Bifidobacterium dentium		1.86	-.07	MLZ 13			
67573521	**Good identification**						**67573521**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67573522	**Low discrimination**						**67573522**
					GRA	CAT	
Actinomyces israelii		62.88	0.37	SAL 92	99	1	
Actinomyces naeslundii		24.26	0.39	XYL 92	99	25	
Capnocytophaga ochracea		12.86	0.09	NAG 6	1	90	
67573523	**Good identification**						**67573523**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67573540	**Low discrimination**						**67573540**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		71.59	0.47	SAL 92	99		
Mitsuokella multacida		27.21	0.27	XYL 92	1		
Actinomyces naeslundii		1.20	0.26		99		
67573542	**Acceptable identification**						**67573542**
					GRA		
Actinomyces israelii		93.61	0.50	SAL 92	99		
Bifidobacterium adolescentis		2.70	0.17	XYL 92	99		
Mitsuokella multacida		2.64	0.11		1		
67573552	**Low discrimination**						**67573552**
					GRA		
Actinomyces israelii		76.98	0.36	SAL 92	99		
Bifidobacterium adolescentis		18.18	0.19	XYL 92	99		
Mitsuokella multacida		4.84	0.03	MLZ 13	1		
67573702	**Low discrimination**						**67573702**
					GRA		
Actinomyces israelii		78.73	0.38	SAL 92	99		
Mitsuokella multacida		21.27	0.15	XYL 92	1		
67574712	**Good identification**						**67574712**
Clostridium clostridioforme		99.46	0.43	NIT 1			
Mitsuokella multacida		0.54	0.01	SAL 88			
67574732	**Good identification**						**67574732**
Clostridium clostridioforme		98.33	0.43	NIT 1			
Parabacteroides distasonis		1.68	0.18	SAL 88			
67575000	**Good identification**						**67575000**
Actinomyces israelii		98.93	0.50	SAL 92			
Actinomyces naeslundii		1.07	0.26				
67575002	**Good identification**						**67575002**
Actinomyces israelii		99.29	0.53	SAL 92			
Actinomyces naeslundii		0.71	0.26				
67575010	**Good identification**						**67575010**
Actinomyces israelii		100.0	0.36	SAL 92			
67575012	**Good identification**						**67575012**
Actinomyces israelii		95.69	0.39	SAL 92			
Clostridium clostridioforme		4.31	0.07	MLZ 13			
67575040	**Good identification**						**67575040**
Actinomyces israelii		100.0	0.43	SAL 92			
67575042	**Good identification**						**67575042**
Actinomyces israelii		100.0	0.46	SAL 92			
67575100	**Low discrimination**						**67575100**
					COC	SPO	
Actinomyces israelii		84.54	0.61	SAL 92	1	1	
Atopobium parvulum		8.27	0.30		99	1	
Clostridium butyricum		3.23	0.15		1	99	
67575101	**Low discrimination**						**67575101**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces naeslundii		47.08	0.37	ESL 12	99		
Actinomyces israelii		41.31	0.27	XYL 1	99		
Mitsuokella multacida		11.61	0.04		1		
67575102	**Low discrimination**						**67575102**
					GRA	CAT	
Actinomyces israelii		85.49	0.64	SAL 92	99	1	
Capnocytophaga ochracea		7.01	0.29		1	90	
Bifidobacterium adolescentis		3.83	0.34		99	1	
67575110	**Acceptable identification**						**67575110**
					GRA	SPO	
Actinomyces israelii		93.42	0.47	SAL 92	99	1	
Mitsuokella multacida		5.32	0.13	MLZ 13	1	1	
Clostridium butyricum		1.26	-.07		99	99	
67575112	**Low discrimination**						**67575112**
					GRA		
Actinomyces israelii		63.50	0.50	SAL 92	99		
Bifidobacterium adolescentis		23.25	0.36	MLZ 13	99		
Leptotrichia buccalis		6.13	0.07		1		
67575121	**Good identification**						**67575121**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67575122	**Low discrimination**						**67575122**
Capnocytophaga ochracea		41.53	0.45	TRE 1			
Bacteroides fragilis		30.53	0.34				
Parabacteroides distasonis		20.69	0.53				
67575123	**Low discrimination**						**67575123**
					GRA		
Actinomyces naeslundii		31.54	0.37	ESL 12	99		
Capnocytophaga ochracea		30.65	0.11	XYL 1	1		
Bacteroides fragilis		22.53	-.01		1		
67575132	**Low discrimination**						**67575132**
					GRA	CAT	
Parabacteroides distasonis		65.51	0.48	NIT 1	1	76	
Capnocytophaga ochracea		27.13	0.29		1	90	
Actinomyces israelii		3.16	0.29		99	1	
67575140	**Acceptable identification**						**67575140**
					GRA	SPO	
Actinomyces israelii		93.09	0.54	SAL 92	99	1	
Mitsuokella multacida		6.12	0.21		1	1	
Clostridium butyricum		0.79	-.03		99	99	
67575142	**Very good identification**						**67575142**
					GRA	CAT	
Actinomyces israelii		95.92	0.57	SAL 92	99	1	
Capnocytophaga ochracea		2.25	0.13		1	90	
Bifidobacterium adolescentis		1.37	0.19		99	1	
67575150	**Low discrimination**						**67575150**
					GRA		
Actinomyces israelii		87.22	0.40	SAL 92	99		
Mitsuokella multacida		12.78	0.13	MLZ 13	1		
67575152	**Low discrimination**						**67575152**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		89.58	0.43	SAL 92			
Bifidobacterium adolescentis		10.43	0.20	MLZ 13			
67575300	**Low discrimination**						**67575300**
					GRA		
Actinomyces israelii		61.34	0.41	SAL 92	99		
Mitsuokella multacida		38.66	0.25	RHA 7	1		
67575302	**Low discrimination**						**67575302**
					GRA		
Actinomyces israelii		75.09	0.45	SAL 92	99		
Parabacteroides distasonis		11.12	0.27	RHA 7	1		
Clostridium clostridioforme		9.06	0.19		50		
67575322	**Very good identification**						**67575322**
Parabacteroides distasonis		95.29	0.61	NIT 1			
Prevotella oralis		2.02	0.19				
Clostridium clostridioforme		0.78	0.19				
67575332	**Acceptable identification**						**67575332**
Parabacteroides distasonis		93.25	0.56	NIT 1			
Clostridium clostridioforme		6.49	0.31				
Bacteroides ovatus		0.26	-.04				
67575342	**Low discrimination**						**67575342**
					GRA		
Actinomyces israelii		89.28	0.37	SAL 92	99		
Mitsuokella multacida		10.72	0.09	RHA 7	1		
67575400	**Low discrimination**						**67575400**
					GRA	SPO	
Actinomyces israelii		83.70	0.49	SAL 92	99	1	
Mitsuokella multacida		10.45	0.21		1	1	
Clostridium butyricum		5.85	0.08		99	99	
67575402	**Acceptable identification**						**67575402**
					SPO		
Actinomyces israelii		93.41	0.52	SAL 92	1		
Clostridium clostridioforme		2.65	0.16		66		
Clostridium butyricum		1.69	0.01		99		
67575412	**Low discrimination**						**67575412**
Actinomyces israelii		50.39	0.38	SAL 92			
Clostridium clostridioforme		43.51	0.27	MLZ 13			
Bifidobacterium adolescentis		6.10	0.16				
67575440	**Low discrimination**						**67575440**
					GRA		
Actinomyces israelii		75.69	0.42	SAL 92	99		
Mitsuokella multacida		24.31	0.21		1		
67575442	**Good identification**						**67575442**
					GRA		
Actinomyces israelii		97.68	0.45	SAL 92	99		
Mitsuokella multacida		2.32	0.04		1		
67575500	**Low discrimination**						**67575500**
					GRA	SPO	
Mitsuokella multacida		68.36	0.55	SAL 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		22.12	0.59		99	1	
Clostridium butyricum		9.28	0.32		99	99	
67575501	**Good identification**						**67575501**
					GRA	SPO	
Mitsuokella multacida		95.99	0.39	SAL 99	1	1	
Actinomyces israelii		2.82	0.25	URE 10	99	1	
Clostridium butyricum		1.19	-.03		99	99	
67575502	**Low discrimination**						**67575502**
					GRA		
Actinomyces israelii		57.15	0.62	SAL 92	99		
Bifidobacterium adolescentis		20.93	0.48		99		
Mitsuokella multacida		13.08	0.39		1		
67575510	**Low discrimination**						**67575510**
					GRA	SPO	
Mitsuokella multacida		84.93	0.47	SAL 99	1	1	
Actinomyces israelii		12.32	0.45		99	1	
Clostridium butyricum		1.82	0.10		99	99	
67575512	**Low discrimination**						**67575512**
Bifidobacterium adolescentis		56.23	0.50	NIT 1			
Actinomyces israelii		18.77	0.48				
Clostridium clostridioforme		13.57	0.36				
67575520	**Low discrimination**						**67575520**
					GRA	SPO	
Mitsuokella multacida		83.44	0.45	SAL 99	1	1	
Actinomyces israelii		6.89	0.39		99	1	
Clostridium butyricum		3.94	0.14		99	99	
67575522	**Low discrimination**						**67575522**
					GRA	CAT	
Parabacteroides distasonis		43.78	0.49	NIT 1	1	76	
Capnocytophaga ochracea		25.30	0.32		1	90	
Actinomyces israelii		10.14	0.42		99	1	
67575530	**Low discrimination**						**67575530**
					GRA		
Mitsuokella multacida		88.75	0.37	SAL 99	1		
Bifidobacterium longum subsp. longum		6.18	0.21		99		
Actinomyces israelii		3.29	0.25		99		
67575532	**Low discrimination**						**67575532**
Parabacteroides distasonis		44.44	0.44	NIT 1			
Clostridium clostridioforme		32.34	0.36				
Mitsuokella multacida		5.72	0.20				
67575540	**Low discrimination**						**67575540**
					GRA	SPO	
Mitsuokella multacida		87.90	0.55	SAL 99	1	1	
Actinomyces israelii		11.06	0.52		99	1	
Clostridium butyricum		1.04	0.14		99	99	
67575541	**Good identification**						**67575541**
					GRA		
Mitsuokella multacida		98.87	0.39	SAL 99	1		
Actinomyces israelii		1.13	0.18	URE 10	99		
67575542	**Low discrimination**						**67575542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		57.65	0.55	SAL 92	99		
Mitsuokella multacida		33.94	0.39		1		
Bifidobacterium adolescentis		6.71	0.33		99		
67575550	**Acceptable identification**						**67575550**
					COC	GRA	
Mitsuokella multacida		94.35	0.47	SAL 99	1	1	
Actinomyces israelii		5.32	0.38		1	99	
Blautia producta		0.33	-.03		99	99	
67575560	**Good identification**						**67575560**
					GRA	SPO	
Mitsuokella multacida		96.15	0.45	SAL 99	1	1	
Actinomyces israelii		3.09	0.32		99	1	
Clostridium butyricum		0.40	-.05		99	99	
67575570	**Good identification**						**67575570**
					GRA		
Mitsuokella multacida		98.58	0.37	SAL 99	1		
Actinomyces israelii		1.42	0.18		99		
67575612	**Good identification**						**67575612**
					GRA		
Clostridium clostridioforme		98.24	0.42	NIT 1	50		
Actinomyces israelii		1.17	0.19	SAL 88	99		
Mitsuokella multacida		0.60	0.01		1		
67575632	**Good identification**						**67575632**
Clostridium clostridioforme		97.38	0.42	NIT 1			
Parabacteroides distasonis		1.85	0.18	SAL 88			
Bacteroides thetaiotaomicron		0.77	-.04				
67575700	**Very good identification**						**67575700**
					GRA	SPO	
Mitsuokella multacida		98.44	0.60	SAL 99	1	1	
Actinomyces israelii		1.29	0.40		99	1	
Clostridium butyricum		0.22	0.06		99	99	
67575701	**Good identification**						**67575701**
Mitsuokella multacida		100.0	0.43	SAL 99			
67575702	**Low discrimination**						**67575702**
					GRA		
Mitsuokella multacida		61.44	0.43	SAL 99	1		
Clostridium clostridioforme		25.12	0.40	BGL 10	50		
Actinomyces israelii		10.88	0.43		99		
67575710	**Good identification**						**67575710**
					GRA		
Mitsuokella multacida		98.80	0.51	SAL 99	1		
Clostridium clostridioforme		0.62	0.17		50		
Actinomyces israelii		0.58	0.26		99		
67575711	**Good identification**						**67575711**
Mitsuokella multacida		100.0	0.35	SAL 99			
67575712	**Low discrimination**						**67575712**
					GRA		
Clostridium clostridioforme		83.02	0.51	NIT 1	50		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Mitsuokella multacida		14.86	0.35	SAL 88	1	
Actinomyces israelii		1.18	0.29		99	
67575720	**Acceptable identification**					**67575720**
Mitsuokella multacida		91.74	0.49	SAL 99		
Bacteroides vulgatus		6.76	0.20			
Parabacteroides distasonis		0.80	0.23			
67575722	**Low discrimination**					**67575722**
Parabacteroides distasonis		57.11	0.57	NIT 1		
Bacteroides thetaiotaomicron		16.33	0.32			
Clostridium clostridioforme		12.01	0.40			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67575730	**Good identification**						**67575730**
Mitsuokella multacida		96.23	0.41	SAL 99			
Clostridium clostridioforme		2.41	0.17				
Parabacteroides distasonis		1.36	0.18				
67575732	**Low discrimination**						**67575732**
Clostridium clostridioforme		55.39	0.51	NIT 1			
Parabacteroides distasonis		31.14	0.52	SAL 88			
Bacteroides thetaiotaomicron		6.43	0.25				
67575740	**Very good identification**						**67575740**
					GRA		
Mitsuokella multacida		99.49	0.60	SAL 99	1		
Actinomyces israelii		0.51	0.33		99		
67575741	**Good identification**						**67575741**
Mitsuokella multacida		100.0	0.43	SAL 99			
67575742	**Acceptable identification**						**67575742**
					GRA		
Mitsuokella multacida		93.20	0.43	SAL 99	1		
Actinomyces israelii		6.42	0.36	BGL 10	99		
Clostridium clostridioforme		0.39	0.06		50		
67575750	**Good identification**						**67575750**
					COC	GRA	
Mitsuokella multacida		99.58	0.51	SAL 99	1	1	
Actinomyces israelii		0.23	0.19		1	99	
Blautia producta		0.19	-.03		99	99	
67575751	**Good identification**						**67575751**
Mitsuokella multacida		100.0	0.35	SAL 99			
67575752	**Acceptable identification**						**67575752**
					GRA		
Mitsuokella multacida		91.97	0.35	SAL 99	1		
Clostridium clostridioforme		5.19	0.17	BGL 10	50		
Actinomyces israelii		2.84	0.22		99		
67575760	**Good identification**						**67575760**
Mitsuokella multacida		100.0	0.49	SAL 99			
67575770	**Good identification**						**67575770**
Mitsuokella multacida		100.0	0.41	SAL 99			
67576100	**Good identification**						**67576100**
Actinomyces israelii		100.0	0.39	SAL 92			
67576102	**Good identification**						**67576102**
Actinomyces israelii		100.0	0.42	SAL 92			
67576142	**Good identification**						**67576142**
Actinomyces israelii		100.0	0.35	SAL 92			
67576502	**Low discrimination**						**67576502**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		87.17	0.41	SAL 92	99	
Mitsuokella multacida		8.22	0.11	GAL 99	1	
Bifidobacterium adolescentis		4.61	0.13		99	
67577000	**Very good identification**					**67577000**
Actinomyces israelii		99.63	0.63	SAL 92		
Actinomyces naeslundii		0.37	0.31			
67577002	**Very good identification**					**67577002**
Actinomyces israelii		99.75	0.66	SAL 92		
Actinomyces naeslundii		0.25	0.31			
67577010	**Good identification**					**67577010**
Actinomyces israelii		100.0	0.49	SAL 92		
67577012	**Good identification**					**67577012**
Actinomyces israelii		100.0	0.52	SAL 92		
67577020	**Acceptable identification**					**67577020**
Actinomyces israelii		94.53	0.43	SAL 92		
Actinomyces naeslundii		5.47	0.31	NAG 6		
67577022	**Good identification**					**67577022**
Actinomyces israelii		96.29	0.46	SAL 92		
Actinomyces naeslundii		3.72	0.31	NAG 6		
67577040	**Very good identification**					**67577040**
Actinomyces israelii		100.0	0.56	SAL 92		
67577042	**Very good identification**					**67577042**
Actinomyces israelii		100.0	0.59	SAL 92		
67577050	**Good identification**					**67577050**
Actinomyces israelii		100.0	0.42	SAL 92		
67577052	**Good identification**					**67577052**
Actinomyces israelii		100.0	0.45	SAL 92		
67577060	**Good identification**					**67577060**
Actinomyces israelii		100.0	0.36	SAL 92		
67577062	**Good identification**					**67577062**
Actinomyces israelii		100.0	0.39	SAL 92		
67577100	**Very good identification**					**67577100**
					GRA	
Actinomyces israelii		98.63	0.74	SAL 92	99	
Mitsuokella multacida		1.04	0.27		1	
Actinomyces naeslundii		0.26	0.39		99	
67577101	**Low discrimination**					**67577101**
					GRA	
Actinomyces israelii		66.40	0.39	SAL 92	99	
Actinomyces naeslundii		25.92	0.42	URE 1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		7.68	0.11		1		
67577102	**Very good identification**						**67577102**
Actinomyces israelii		99.11	0.77	SAL 92			
Bifidobacterium adolescentis		0.64	0.33				
Actinomyces naeslundii		0.17	0.39				
67577103	**Low discrimination**						**67577103**
Actinomyces israelii		79.35	0.42	SAL 92			
Actinomyces naeslundii		20.65	0.42	URE 1			
67577110	**Very good identification**						**67577110**
					GRA		
Actinomyces israelii		97.71	0.59	SAL 92	99		
Mitsuokella multacida		2.29	0.19	MLZ 13	1		
67577112	**Good identification**						**67577112**
Actinomyces israelii		94.61	0.62	SAL 92			
Bifidobacterium adolescentis		5.00	0.34	MLZ 13			
Bifidobacterium dentium		0.23	-.07				
67577120	**Acceptable identification**						**67577120**
					GRA		
Actinomyces israelii		92.39	0.53	SAL 92	99		
Mitsuokella multacida		3.81	0.17	NAG 6	1		
Actinomyces naeslundii		3.81	0.39		99		
67577121	**Low discrimination**						**67577121**
Actinomyces naeslundii		85.95	0.42	ESL 12			
Actinomyces israelii		14.05	0.19	XYL 1			
67577122	**Good identification**						**67577122**
					GRA	CAT	
Actinomyces israelii		93.15	0.56	SAL 92	99	1	
Actinomyces naeslundii		2.56	0.39	NAG 6	99	25	
Capnocytophaga ochracea		1.92	0.11		1	90	
67577123	**Low discrimination**						**67577123**
Actinomyces naeslundii		80.30	0.42	ESL 12			
Actinomyces israelii		19.70	0.22	XYL 1			
67577130	**Acceptable identification**						**67577130**
					GRA		
Actinomyces israelii		91.58	0.39	SAL 92	99		
Mitsuokella multacida		8.42	0.09	MLZ 13	1		
67577132	**Acceptable identification**						**67577132**
					GRA		
Actinomyces israelii		90.64	0.42	SAL 92	99		
Bifidobacterium dentium		6.01	-.03	MLZ 13	99		
Parabacteroides distasonis		3.36	0.14	NAG 6	1		
67577140	**Very good identification**						**67577140**
					GRA		
Actinomyces israelii		97.25	0.67	SAL 92	99		
Mitsuokella multacida		2.63	0.27		1		
Actinomyces naeslundii		0.12	0.26		99		
67577142	**Very good identification**						**67577142**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				GRA		
Actinomyces israelii	99.52	0.70	SAL 92	99		
Bifidobacterium adolescentis	0.20	0.17		99		
Mitsuokella multacida	0.20	0.11		1		
67577143 Low discrimination						**67577143**
Actinomyces israelii	89.44	0.35	SAL 92			
Actinomyces naeslundii	10.56	0.29	URE 1			
67577150 Acceptable identification						**67577150**
				COC	GRA	
Actinomyces israelii	93.51	0.52	SAL 92	1	99	
Mitsuokella multacida	5.64	0.19	MLZ 13	1	1	
Blautia producta	0.85	-.03		99	99	
67577152 Good identification						**67577152**
				GRA		
Actinomyces israelii	97.92	0.55	SAL 92	99		
Bifidobacterium adolescentis	1.65	0.19	MLZ 13	99		
Mitsuokella multacida	0.44	0.03		1		
67577160 Low discrimination						**67577160**
				GRA		
Actinomyces israelii	88.92	0.46	SAL 92	99		
Mitsuokella multacida	9.42	0.17	NAG 6	1		
Actinomyces naeslundii	1.66	0.26		99		
67577162 Good identification						**67577162**
Actinomyces israelii	98.77	0.49	SAL 92			
Actinomyces naeslundii	1.23	0.26	NAG 6			
67577172 Good identification						**67577172**
Actinomyces israelii	100.0	0.35	SAL 92			
67577200 Good identification						**67577200**
Actinomyces israelii	100.0	0.44	SAL 92			
67577202 Good identification						**67577202**
Actinomyces israelii	100.0	0.47	SAL 92			
67577240 Good identification						**67577240**
Actinomyces israelii	100.0	0.37	SAL 92			
67577242 Good identification						**67577242**
Actinomyces israelii	100.0	0.40	SAL 92			
67577300 Low discrimination						**67577300**
				GRA		
Actinomyces israelii	79.40	0.54	SAL 92	99		
Mitsuokella multacida	20.60	0.32	RHA 7	1		
67577302 Very good identification						**67577302**
				GRA		
Actinomyces israelii	98.11	0.57	SAL 92	99		
Mitsuokella multacida	1.89	0.15	RHA 7	1		
67577310 Low discrimination						**67577310**
				GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		63.33	0.40	SAL 92	99	
Mitsuokella multacida		36.67	0.24	RHA 7	1	
67577312	**Good identification**					**67577312**
					GRA	
Actinomyces israelii		95.89	0.43	SAL 92	99	
Mitsuokella multacida		4.11	0.07	RHA 7	1	
67577322	**Low discrimination**					**67577322**
					GRA	
Actinomyces israelii		65.44	0.37	SAL 92	99	
Parabacteroides distasonis		26.79	0.27	RHA 7	1	
Mitsuokella multacida		4.93	0.05	NAG 6	1	
67577340	**Low discrimination**					**67577340**
					GRA	
Actinomyces israelii		59.98	0.47	SAL 92	99	
Mitsuokella multacida		40.02	0.32	RHA 7	1	
67577342	**Good identification**					**67577342**
					GRA	
Actinomyces israelii		95.29	0.50	SAL 92	99	
Mitsuokella multacida		4.71	0.15	RHA 7	1	
67577352	**Acceptable identification**					**67577352**
					GRA	
Actinomyces israelii		90.07	0.36	SAL 92	99	
Mitsuokella multacida		9.93	0.07	RHA 7	1	
67577400	**Very good identification**					**67577400**
					GRA	
Actinomyces israelii		95.11	0.62	SAL 92	99	
Mitsuokella multacida		4.89	0.27		1	
67577402	**Very good identification**					**67577402**
					GRA	
Actinomyces israelii		99.41	0.65	SAL 92	99	
Mitsuokella multacida		0.38	0.11		1	
Bifidobacterium adolescentis		0.21	0.13		99	
67577410	**Low discrimination**					**67577410**
					GRA	
Actinomyces israelii		89.71	0.48	SAL 92	99	
Mitsuokella multacida		10.29	0.19	MLZ 13	1	
67577412	**Good identification**					**67577412**
Actinomyces israelii		96.38	0.51	SAL 92		
Bifidobacterium adolescentis		1.68	0.14	MLZ 13		
Bifidobacterium dentium		1.12	-.07			
67577420	**Low discrimination**					**67577420**
					GRA	
Actinomyces israelii		83.24	0.41	SAL 92	99	
Mitsuokella multacida		16.76	0.17	NAG 6	1	
67577422	**Good identification**					**67577422**
Actinomyces israelii		100.0	0.44	SAL 92		
67577440	**Low discrimination**					**67577440**
					GRA	
Actinomyces israelii		88.32	0.55	SAL 92	99	
Mitsuokella multacida		11.68	0.27		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67577442	**Very good identification**						**67577442**
					GRA		
Actinomyces israelii		99.03	0.58	SAL 92	99		
Mitsuokella multacida		0.97	0.11		1		
67577450	**Low discrimination**						**67577450**
					COC	GRA	
Actinomyces israelii		74.67	0.41	SAL 92	1	99	
Mitsuokella multacida		22.03	0.19	MLZ 13	1	1	
Blautia producta		3.30	-.03		99	99	
67577452	**Good identification**						**67577452**
					GRA		
Actinomyces israelii		97.86	0.44	SAL 92	99		
Mitsuokella multacida		2.14	0.03	MLZ 13	1		
67577462	**Good identification**						**67577462**
Actinomyces israelii		100.0	0.37	SAL 92			
67577500	**Low discrimination**						**67577500**
					GRA	SPO	
Mitsuokella multacida		55.78	0.61	SAL 99	1	1	
Actinomyces israelii		43.83	0.72		99	1	
Clostridium butyricum		0.36	0.15		99	99	
67577501	**Acceptable identification**						**67577501**
					GRA		
Mitsuokella multacida		93.33	0.45	SAL 99	1		
Actinomyces israelii		6.67	0.38	URE 10	99		
67577502	**Low discrimination**						**67577502**
					GRA		
Actinomyces israelii		86.98	0.75	SAL 92	99		
Mitsuokella multacida		8.20	0.45		1		
Bifidobacterium adolescentis		4.60	0.47		99		
67577510	**Low discrimination**						**67577510**
					GRA		
Mitsuokella multacida		72.88	0.53	SAL 99	1		
Actinomyces israelii		25.67	0.58		99		
Bifidobacterium dentium		1.10	0.10		99		
67577511	**Good identification**						**67577511**
					GRA		
Mitsuokella multacida		96.90	0.37	SAL 99	1		
Actinomyces israelii		3.10	0.24	URE 10	99		
67577512	**Low discrimination**						**67577512**
Actinomyces israelii		51.79	0.61	SAL 92			
Bifidobacterium adolescentis		22.40	0.48	MLZ 13			
Bifidobacterium dentium		14.92	0.27				
67577520	**Low discrimination**						**67577520**
					GRA	SPO	
Mitsuokella multacida		83.14	0.51	SAL 99	1	1	
Actinomyces israelii		16.68	0.52		99	1	
Clostridium butyricum		0.19	-.03		99	99	
67577522	**Low discrimination**						**67577522**
					GRA		
Actinomyces israelii		71.09	0.55	SAL 92	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		26.25	0.35	NAG 6	1		
Bifidobacterium dentium		0.86	-.03		99		
67577530	**Low discrimination**						**67577530**
					GRA		
Mitsuokella multacida		83.55	0.43	SAL 99	1		
Bifidobacterium dentium		8.94	0.15		99		
Actinomyces israelii		7.51	0.38		99		
67577540	**Low discrimination**						**67577540**
					COC	GRA	
Mitsuokella multacida		75.60	0.61	SAL 99	1	1	
Actinomyces israelii		23.10	0.65		1	99	
Blautia producta		1.26	0.24		99	99	
67577541	**Good identification**						**67577541**
					GRA		
Mitsuokella multacida		97.30	0.45	SAL 99	1		
Actinomyces israelii		2.70	0.31	URE 10	99		
67577542	**Low discrimination**						**67577542**
					GRA		
Actinomyces israelii		79.16	0.68	SAL 92	99		
Mitsuokella multacida		19.19	0.45		1		
Bifidobacterium adolescentis		1.33	0.31		99		
67577550	**Low discrimination**						**67577550**
					COC	GRA	
Mitsuokella multacida		77.70	0.53	SAL 99	1	1	
Blautia producta		11.66	0.32		99	99	
Actinomyces israelii		10.64	0.51		1	99	
67577551	**Good identification**						**67577551**
					COC	GRA	
Mitsuokella multacida		97.46	0.37	SAL 99	1	1	
Blautia producta		1.33	-.03	URE 10	99	99	
Actinomyces israelii		1.21	0.17		1	99	
67577552	**Low discrimination**						**67577552**
					GRA		
Actinomyces israelii		56.60	0.54	SAL 92	99		
Mitsuokella multacida		30.61	0.37	MLZ 13	1		
Bifidobacterium adolescentis		7.78	0.33		99		
67577560	**Acceptable identification**						**67577560**
					COC	GRA	
Mitsuokella multacida		92.13	0.51	SAL 99	1	1	
Actinomyces israelii		7.19	0.45		1	99	
Blautia producta		0.68	0.07		99	99	
67577562	**Low discrimination**						**67577562**
					GRA		
Actinomyces israelii		51.30	0.48	SAL 92	99		
Mitsuokella multacida		48.70	0.35	NAG 6	1		
67577570	**Acceptable identification**						**67577570**
					COC	GRA	
Mitsuokella multacida		90.77	0.43	SAL 99	1	1	
Blautia producta		6.05	0.15		99	99	
Actinomyces israelii		3.18	0.31		1	99	
67577602	**Acceptable identification**						**67577602**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		91.41	0.46	SAL 92	99		
Mitsuokella multacida		8.59	0.15	RHA 7	1		
67577642	**Low discrimination**						**67577642**
					GRA		
Actinomyces israelii		80.54	0.39	SAL 92	99		
Mitsuokella multacida		19.46	0.15	RHA 7	1		
67577700	**Very good identification**						**67577700**
					GRA		
Mitsuokella multacida		96.91	0.66	SAL 99	1		
Actinomyces israelii		3.09	0.53		99		
67577701	**Good identification**						**67577701**
					GRA		
Mitsuokella multacida		99.71	0.50	SAL 99	1		
Actinomyces israelii		0.29	0.19	URE 10	99		
67577702	**Low discrimination**						**67577702**
					GRA		
Mitsuokella multacida		69.70	0.50	SAL 99	1		
Actinomyces israelii		29.97	0.56	BGL 10	99		
Bifidobacterium adolescentis		0.21	0.13		99		
67577710	**Very good identification**						**67577710**
					COC	GRA	
Mitsuokella multacida		98.51	0.58	SAL 99	1	1	
Actinomyces israelii		1.41	0.39		1	99	
Blautia producta		0.08	-.03		99	99	
67577711	**Good identification**						**67577711**
Mitsuokella multacida		100.0	0.41	SAL 99			
67577712	**Low discrimination**						**67577712**
					GRA		
Mitsuokella multacida		80.94	0.41	SAL 99	1		
Actinomyces israelii		15.60	0.42	BGL 10	99		
Clostridium clostridioforme		1.96	0.17		50		
67577720	**Good identification**						**67577720**
					GRA		
Mitsuokella multacida		99.19	0.56	SAL 99	1		
Actinomyces israelii		0.81	0.32		99		
67577721	**Good identification**						**67577721**
Mitsuokella multacida		100.0	0.39	SAL 99			
67577722	**Low discrimination**						**67577722**
					GRA		
Mitsuokella multacida		77.43	0.39	SAL 99	1		
Bacteroides ovatus		10.18	0.17	BGL 10	1		
Actinomyces israelii		8.50	0.35		99		
67577730	**Good identification**						**67577730**
					GRA		
Mitsuokella multacida		99.64	0.47	SAL 99	1		
Actinomyces israelii		0.36	0.18		99		
67577740	**Very good identification**						**67577740**
					COC	GRA	
Mitsuokella multacida		97.91	0.66	SAL 99	1	1	
Actinomyces israelii		1.21	0.46		1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		0.88	0.24		99	99	
67577741	**Good identification**						**67577741**
Mitsuokella multacida		100.0	0.50	SAL 99			
67577742	**Low discrimination**						**67577742**
					COC	GRA	
Mitsuokella multacida		85.02	0.50	SAL 99	1	1	
Actinomyces israelii		14.22	0.49	BGL 10	1	99	
Blautia producta		0.76	0.07		99	99	
67577750	**Good identification**						**67577750**
					COC	GRA	
Mitsuokella multacida		92.05	0.58	SAL 99	1	1	
Blautia producta		7.44	0.32		99	99	
Actinomyces israelii		0.51	0.32		1	99	
67577751	**Good identification**						**67577751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	SAL 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67577752	**Low discrimination**						**67577752**
					COC	GRA	
Mitsuokella multacida		86.53	0.41	SAL 99	1	1	
Blautia producta		6.99	0.15	BGL 10	99	99	
Actinomyces israelii		6.49	0.35		1	99	
67577760	**Good identification**						**67577760**
					COC	GRA	
Mitsuokella multacida		99.29	0.56	SAL 99	1	1	
Blautia producta		0.40	0.07		99	99	
Actinomyces israelii		0.31	0.25		1	99	
67577761	**Good identification**						**67577761**
Mitsuokella multacida		100.0	0.39	SAL 99			
67577762	**Good identification**						**67577762**
					GRA		
Mitsuokella multacida		95.41	0.39	SAL 99	1		
Actinomyces israelii		4.07	0.28	BGL 10	99		
Bacteroides ovatus		0.52	-.07		1		
67577770	**Good identification**						**67577770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	SAL 99	1	1	
Blautia producta		3.47	0.15		99	99	
67601000	**Very good identification**						**67601000**
					GRA		
Eubacterium tortuosum		99.83	0.56	MNS 1	99		
Fusobacterium mortiferum		0.17	0.25		1		
67601002	**Good identification**						**67601002**
Eubacterium tortuosum		100.0	0.43	MNS 1			
67601020	**Good identification**						**67601020**
Eubacterium tortuosum		100.0	0.43	MNS 1			
67603101	**Good identification**						**67603101**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	ESL 12		
67603103	**Good identification**					**67603103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67603121	**Good identification**					**67603121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67603123	**Good identification**					**67603123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67610000	**Very good identification**					**67610000**
Fusobacterium mortiferum		100.0	0.57	NIT 1		
67610002	**Low discrimination**					**67610002**
					GRA	
Fusobacterium mortiferum		79.49	0.40	NIT 1	1	
Actinomyces viscosus		20.51	0.18	BGL 10	99	
67610100	**Very good identification**					**67610100**
					GRA	
Fusobacterium mortiferum		99.85	0.63	NIT 1	1	
Actinomyces naeslundii		0.15	0.26		99	
67610102	**Low discrimination**					**67610102**
					GRA	
Fusobacterium mortiferum		82.07	0.47	NIT 1	1	
Actinomyces viscosus		16.14	0.22	BGL 10	99	
Actinomyces naeslundii		1.12	0.26		99	
67611000	**Good identification**					**67611000**
					GRA	
Fusobacterium mortiferum		92.55	0.60	NIT 1	1	
Eubacterium tortuosum		5.61	0.22		99	
Actinomyces naeslundii		1.84	0.42		99	
67611001	**Low discrimination**					**67611001**
					GRA	
Actinomyces naeslundii		74.66	0.45	ESL 12	99	
Fusobacterium mortiferum		25.34	0.25	MLT 94	1	
67611002	**Low discrimination**					**67611002**
					GRA	
Fusobacterium mortiferum		71.78	0.43	NIT 1	1	
Actinomyces naeslundii		12.82	0.42	BGL 10	99	
Actinomyces viscosus		7.57	0.14		99	
67611003	**Low discrimination**					**67611003**
Actinomyces naeslundii		88.40	0.45	ESL 12		
Actinomyces viscosus		11.60	0.06	MLT 94		
67611020	**Low discrimination**					**67611020**
					GRA	
Actinomyces naeslundii		45.75	0.42	ESL 12	99	
Eubacterium tortuosum		24.68	0.09	MLT 94	99	
Fusobacterium mortiferum		23.29	0.25		1	
67611021	**Good identification**					**67611021**
Actinomyces naeslundii		100.0	0.45	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67611022	**Low discrimination**						**67611022**
					GRA		
Actinomyces naeslundii		69.35	0.42	ESL 12	99		
Prevotella oralis		9.89	0.02	MLT 94	1		
Bifidobacterium longum subsp. infantis		8.82	-.03		99		
67611023	**Good identification**						**67611023**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67611100	**Very good identification**						**67611100**
					GRA		
Fusobacterium mortiferum		97.00	0.66	NIT 1	1		
Actinomyces naeslundii		2.35	0.50		99		
Bifidobacterium longum subsp. infantis		0.58	0.10		99		
67611101	**Low discrimination**						**67611101**
					GRA		
Actinomyces naeslundii		78.23	0.53	ESL 12	99		
Fusobacterium mortiferum		21.77	0.32	MLT 94	1		
67611102	**Low discrimination**						**67611102**
					GRA		
Fusobacterium mortiferum		52.25	0.50	NIT 1	1		
Bifidobacterium longum subsp. infantis		28.32	0.27	BGL 10	99		
Actinomyces naeslundii		11.38	0.50		99		
67611103	**Low discrimination**						**67611103**
					GRA		
Actinomyces naeslundii		88.52	0.53	ESL 12	99		
Actinomyces viscosus		7.26	0.10	MLT 94	99		
Fusobacterium mortiferum		2.74	0.15		1		
67611120	**Low discrimination**						**67611120**
					GRA		
Actinomyces naeslundii		38.80	0.50	ESL 12	99		
Bifidobacterium longum subsp. infantis		16.98	0.15	MLT 94	99		
Fusobacterium mortiferum		16.19	0.32		1		
67611121	**Good identification**						**67611121**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67611123	**Good identification**						**67611123**
					GRA	CAT	
Actinomyces naeslundii		95.44	0.53	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		2.84	-.03	MLT 94	99	1	
Capnocytophaga ochracea		1.72	-.03		1	90	
67611141	**Good identification**						**67611141**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67611142	**Low discrimination**						**67611142**
					GRA	CAT	
Actinomyces naeslundii		62.79	0.37	ESL 12	99	25	
Fusobacterium mortiferum		16.50	0.15	MLT 94	1	5	
Capnocytophaga ochracea		11.76	-.01		1	90	
67611143	**Good identification**						**67611143**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67611160	**Low discrimination**						**67611160**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	CAT	
Actinomyces naeslundii		63.74	0.37	ESL 12	99	25	
Clostridium perfringens		24.32	0.07	MLT 94	99	1	
Capnocytophaga ochracea		11.94	-.01		1	90	
67611161	**Good identification**						**67611161**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67611163	**Good identification**						**67611163**
Actinomyces naeslundii		100.0	0.40	ESL 12			
67613000	**Low discrimination**						**67613000**
					GRA		
Actinomyces naeslundii		79.22	0.47	ESL 12	99		
Fusobacterium mortiferum		20.78	0.25	MLT 94	1		
67613001	**Good identification**						**67613001**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67613002	**Good identification**						**67613002**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67613003	**Good identification**						**67613003**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67613020	**Good identification**						**67613020**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67613021	**Good identification**						**67613021**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67613022	**Good identification**						**67613022**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67613023	**Good identification**						**67613023**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67613041	**Good identification**						**67613041**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67613043	**Good identification**						**67613043**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67613061	**Good identification**						**67613061**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67613063	**Good identification**						**67613063**
Actinomyces naeslundii		100.0	0.37	ESL 12			
67613100	**Low discrimination**						**67613100**
					GRA		
Actinomyces naeslundii		82.30	0.55	ESL 12	99		
Fusobacterium mortiferum		17.70	0.32	MLT 94	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67613101	**Very good identification**						**67613101**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67613102	**Good identification**						**67613102**
					GRA		
Actinomyces naeslundii		95.23	0.55	ESL 12	99		
Fusobacterium mortiferum		2.28	0.15	MLT 94	1		
Actinomyces israelii		1.26	0.13		99		
67613103	**Very good identification**						**67613103**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67613111	**Good identification**						**67613111**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67613113	**Good identification**						**67613113**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67613120	**Good identification**						**67613120**
Actinomyces naeslundii		100.0	0.55	ESL 12			
67613121	**Very good identification**						**67613121**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67613122	**Good identification**						**67613122**
					GRA	CAT	
Actinomyces naeslundii		96.44	0.55	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		2.22	-.03	MLT 94	99	1	
Capnocytophaga ochracea		1.34	-.03		1	90	
67613123	**Very good identification**						**67613123**
Actinomyces naeslundii		100.0	0.58	ESL 12			
67613131	**Good identification**						**67613131**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67613133	**Good identification**						**67613133**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67613140	**Good identification**						**67613140**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67613141	**Good identification**						**67613141**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67613142	**Good identification**						**67613142**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67613143	**Good identification**						**67613143**
Actinomyces naeslundii		100.0	0.45	ESL 12			
67613160	**Good identification**						**67613160**

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	ESL 12		
67613161	**Good identification**					**67613161**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67613162	**Good identification**					**67613162**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67613163	**Good identification**					**67613163**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67614000	**Good identification**					**67614000**
Fusobacterium mortiferum		100.0	0.37	NIT 1		
67614100	**Good identification**					**67614100**
Fusobacterium mortiferum		100.0	0.44	NIT 1		
67615000	**Good identification**					**67615000**
Fusobacterium mortiferum		100.0	0.40	NIT 1		
67615100	**Acceptable identification**					**67615100**
					GRA	
Fusobacterium mortiferum		92.94	0.47	NIT 1	1	
Bifidobacterium longum subsp. infantis		6.02	0.09	XYL 7	99	
Actinomyces israelii		1.04	0.16		99	
67615322	**Good identification**					**67615322**
					GRA	
Parabacteroides distasonis		96.75	0.45	NIT 1	1	
Prevotella oralis		1.30	0.00		1	
Bifidobacterium longum subsp. infantis		1.15	-.04		99	
67615332	**Good identification**					**67615332**
Parabacteroides distasonis		100.0	0.41	NIT 1		
67615722	**Good identification**					**67615722**
Parabacteroides distasonis		100.0	0.42	NIT 1		
67615732	**Good identification**					**67615732**
Parabacteroides distasonis		95.87	0.37	NIT 1		
Clostridium clostridioforme		4.13	0.08			
67621101	**Good identification**					**67621101**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67621103	**Good identification**					**67621103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67621121	**Good identification**					**67621121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67621123	**Good identification**					**67621123**
Actinomyces naeslundii		100.0	0.36	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67623100	**Good identification**					**67623100**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67623101	**Good identification**					**67623101**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67623102	**Good identification**					**67623102**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67623103	**Good identification**					**67623103**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67623120	**Good identification**					**67623120**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67623121	**Good identification**					**67623121**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67623122	**Good identification**					**67623122**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67623123	**Good identification**					**67623123**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67630000	**Good identification**					**67630000**
					GRA	
Fusobacterium mortiferum		99.20	0.48	NIT 1	1	
Actinomyces naeslundii		0.80	0.23		99	
67630100	**Good identification**					**67630100**
					GRA	
Fusobacterium mortiferum		99.03	0.54	NIT 1	1	
Actinomyces naeslundii		0.97	0.31		99	
67630102	**Low discrimination**					**67630102**
					GRA	
Fusobacterium mortiferum		87.87	0.38	NIT 1	1	
Actinomyces naeslundii		7.77	0.31	BGL 10	99	
Actinomyces viscosus		4.35	0.03		99	
67631000	**Low discrimination**					**67631000**
					GRA	
Fusobacterium mortiferum		88.58	0.51	NIT 1	1	
Actinomyces naeslundii		11.42	0.47		99	
67631001	**Good identification**					**67631001**
					GRA	
Actinomyces naeslundii		95.04	0.50	ESL 12	99	
Fusobacterium mortiferum		4.96	0.16	MLT 94	1	
67631002	**Low discrimination**					**67631002**
					GRA	
Actinomyces naeslundii		53.71	0.47	ESL 12	99	
Fusobacterium mortiferum		46.29	0.34	MLT 94	1	
67631003	**Good identification**					**67631003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.50	ESL 12		
67631020	**Low discrimination**					**67631020**
					GRA	
Actinomyces naeslundii		82.67	0.47	ESL 12	99	
Clostridium perfringens		10.85	0.10	MLT 94	99	
Fusobacterium mortiferum		6.48	0.16		1	
67631021	**Good identification**					**67631021**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67631022	**Acceptable identification**					**67631022**
Actinomyces naeslundii		94.18	0.47	ESL 12		
Clostridium perfringens		5.82	0.04	MLT 94		
67631023	**Good identification**					**67631023**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67631041	**Good identification**					**67631041**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67631043	**Good identification**					**67631043**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67631061	**Good identification**					**67631061**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67631063	**Good identification**					**67631063**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67631100	**Low discrimination**					**67631100**
					GRA	
Fusobacterium mortiferum		86.41	0.57	NIT 1	1	
Actinomyces naeslundii		13.59	0.55		99	
67631101	**Very good identification**					**67631101**
					GRA	
Actinomyces naeslundii		95.89	0.58	ESL 12	99	
Fusobacterium mortiferum		4.11	0.23	MLT 94	1	
67631102	**Low discrimination**					**67631102**
					GRA	
Actinomyces naeslundii		57.68	0.55	ESL 12	99	
Fusobacterium mortiferum		40.75	0.41	MLT 94	1	
Actinomyces viscosus		0.83	-.01		99	
67631103	**Very good identification**					**67631103**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67631111	**Good identification**					**67631111**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67631113	**Good identification**					**67631113**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67631120	**Low discrimination**					**67631120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		71.30	0.55	ESL 12	99	
Clostridium perfringens		24.12	0.24	MLT 94	99	
Fusobacterium mortiferum		4.58	0.23		1	
67631121	**Very good identification**					**67631121**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67631122	**Low discrimination**					**67631122**
					GRA	
Actinomyces naeslundii		67.11	0.55	ESL 12	99	
Parabacteroides distasonis		18.00	0.32	MLT 94	1	
Clostridium perfringens		10.68	0.19		99	
67631123	**Very good identification**					**67631123**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67631131	**Good identification**					**67631131**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67631133	**Good identification**					**67631133**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67631140	**Low discrimination**					**67631140**
					GRA	
Actinomyces naeslundii		73.32	0.42	ESL 12	99	
Fusobacterium mortiferum		26.69	0.23	MLT 94	1	
67631141	**Good identification**					**67631141**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67631142	**Good identification**					**67631142**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67631143	**Good identification**					**67631143**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67631160	**Low discrimination**					**67631160**
Actinomyces naeslundii		73.25	0.42	ESL 12		
Clostridium perfringens		26.75	0.12	MLT 94		
67631161	**Good identification**					**67631161**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67631162	**Low discrimination**					**67631162**
Actinomyces naeslundii		85.34	0.42	ESL 12		
Clostridium perfringens		14.66	0.06	MLT 94		
67631163	**Good identification**					**67631163**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67631322	**Good identification**					**67631322**
Parabacteroides distasonis		100.0	0.40	NIT 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67631332	**Good identification**					**67631332**
Parabacteroides distasonis		100.0	0.36	NIT 1		
67631722	**Good identification**					**67631722**
Parabacteroides distasonis		100.0	0.37	NIT 1		
67632100	**Low discrimination**					**67632100**
					GRA	
Actinomyces naeslundii		65.39	0.36	ESL 12	99	
Fusobacterium mortiferum		34.61	0.20	MLT 94	1	
67632101	**Good identification**					**67632101**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67632102	**Good identification**					**67632102**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67632103	**Good identification**					**67632103**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67632120	**Good identification**					**67632120**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67632121	**Good identification**					**67632121**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67632122	**Good identification**					**67632122**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67632123	**Good identification**					**67632123**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67633000	**Good identification**					**67633000**
					GRA	
Actinomyces naeslundii		96.12	0.52	ESL 12	99	
Fusobacterium mortiferum		3.88	0.16	MLT 94	1	
67633001	**Good identification**					**67633001**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67633002	**Good identification**					**67633002**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67633003	**Good identification**					**67633003**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67633020	**Good identification**					**67633020**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67633021	**Good identification**					**67633021**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67633022	**Good identification**					**67633022**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.52	ESL 12		
67633023	**Good identification**					**67633023**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67633040	**Good identification**					**67633040**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67633041	**Good identification**					**67633041**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67633042	**Good identification**					**67633042**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67633043	**Good identification**					**67633043**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67633060	**Good identification**					**67633060**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67633061	**Good identification**					**67633061**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67633062	**Good identification**					**67633062**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67633063	**Good identification**					**67633063**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67633100	**Very good identification**					**67633100**
					GRA	
Actinomyces naeslundii		95.22	0.60	ESL 12	99	
Fusobacterium mortiferum		3.15	0.23	MLT 94	1	
Actinomyces israelii		1.63	0.20		99	
67633101	**Very good identification**					**67633101**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67633102	**Very good identification**					**67633102**
Actinomyces naeslundii		97.50	0.60	ESL 12		
Actinomyces israelii		2.50	0.23	MLT 94		
67633103	**Very good identification**					**67633103**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67633110	**Good identification**					**67633110**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67633111	**Good identification**					**67633111**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67633112	**Good identification**					**67633112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	ESL 12		
67633113	**Good identification**					**67633113**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67633120	**Very good identification**					**67633120**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67633121	**Very good identification**					**67633121**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67633122	**Very good identification**					**67633122**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67633123	**Very good identification**					**67633123**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67633130	**Good identification**					**67633130**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67633131	**Good identification**					**67633131**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67633132	**Good identification**					**67633132**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67633133	**Good identification**					**67633133**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67633140	**Good identification**					**67633140**
Actinomyces naeslundii		96.37	0.47	ESL 12		
Actinomyces israelii		3.63	0.13	MLT 94		
67633141	**Good identification**					**67633141**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67633142	**Acceptable identification**					**67633142**
Actinomyces naeslundii		94.65	0.47	ESL 12		
Actinomyces israelii		5.35	0.16	MLT 94		
67633143	**Good identification**					**67633143**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67633160	**Good identification**					**67633160**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67633161	**Good identification**					**67633161**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67633162	**Good identification**					**67633162**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	ESL 12		
67633163	**Good identification**					**67633163**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67635100	**Low discrimination**					**67635100**
					GRA	
Fusobacterium mortiferum		87.65	0.38	NIT 1	1	
Actinomyces israelii		12.35	0.25	XYL 7	99	
67635122	**Good identification**					**67635122**
					GRA	
Parabacteroides distasonis		99.23	0.48	NIT 1	1	
Bifidobacterium longum subsp. infantis		0.77	-.04		99	
67635132	**Good identification**					**67635132**
Parabacteroides distasonis		100.0	0.44	NIT 1		
67635322	**Very good identification**					**67635322**
Parabacteroides distasonis		100.0	0.56	NIT 1		
67635332	**Good identification**					**67635332**
Parabacteroides distasonis		100.0	0.52	NIT 1		
67635522	**Good identification**					**67635522**
Parabacteroides distasonis		100.0	0.45	NIT 1		
67635532	**Good identification**					**67635532**
Parabacteroides distasonis		97.24	0.40	NIT 1		
Clostridium clostridioforme		2.76	0.08			
67635722	**Good identification**					**67635722**
Parabacteroides distasonis		99.19	0.53	NIT 1		
Clostridium clostridioforme		0.81	0.11			
67635732	**Acceptable identification**					**67635732**
Parabacteroides distasonis		93.51	0.48	NIT 1		
Clostridium clostridioforme		6.49	0.23			
67637100	**Good identification**					**67637100**
Actinomyces israelii		95.12	0.38	MLT 99		
Actinomyces naeslundii		4.88	0.26	CEL 93		
67637102	**Good identification**					**67637102**
Actinomyces israelii		96.69	0.41	MLT 99		
Actinomyces naeslundii		3.31	0.26	CEL 93		
67637502	**Low discrimination**					**67637502**
					GRA	
Actinomyces israelii		78.63	0.40	MLT 99	99	
Bifidobacterium adolescentis		12.81	0.20	CEL 93	99	
Mitsuokella multacida		8.56	0.11		1	
67641000	**Good identification**					**67641000**
					GRA	
Eubacterium tortuosum		99.50	0.40	MNS 1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Fusobacterium mortiferum		0.50	0.17	CEL 10	1		
67643101	**Good identification**						**67643101**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67643103	**Good identification**						**67643103**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67643121	**Good identification**						**67643121**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67643123	**Good identification**						**67643123**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67650000	**Good identification**						**67650000**
Fusobacterium mortiferum		100.0	0.48	NIT 1			
67650100	**Good identification**						**67650100**
					GRA		
Fusobacterium mortiferum		99.40	0.55	NIT 1	1		
Actinomyces naeslundii		0.60	0.28		99		
67650102	**Low discrimination**						**67650102**
					GRA		
Fusobacterium mortiferum		82.58	0.38	NIT 1	1		
Actinomyces viscosus		12.95	0.12	BGL 10	99		
Actinomyces naeslundii		4.47	0.28		99		
67651000	**Low discrimination**						**67651000**
					GRA		
Fusobacterium mortiferum		88.99	0.51	NIT 1	1		
Actinomyces naeslundii		7.02	0.44		99		
Eubacterium tortuosum		1.80	0.06		99		
67651001	**Acceptable identification**						**67651001**
					GRA		
Actinomyces naeslundii		92.14	0.47	ESL 12	99		
Fusobacterium mortiferum		7.86	0.17	MLT 94	1		
67651002	**Low discrimination**						**67651002**
					GRA		
Fusobacterium mortiferum		46.90	0.35	NIT 1	1		
Actinomyces naeslundii		33.30	0.44	BGL 10	99		
Lactobacillus catenaformis		8.40	0.12		99		
67651003	**Good identification**						**67651003**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67651020	**Low discrimination**						**67651020**
					GRA	SPO	
Actinomyces naeslundii		74.21	0.44	ESL 12	99	1	
Clostridium barati		11.61	0.04	MLT 94	99	99	
Fusobacterium mortiferum		9.50	0.17		1	1	
67651021	**Good identification**						**67651021**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67651022	**Low discrimination**						**67651022**
					GRA	SPO	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		50.47	0.44	ESL 12	99	1	
Clostridium barati		23.68	0.12	MLT 94	99	99	
Prevotella oralis		19.25	0.11		1	1	
67651023	**Good identification**						**67651023**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67651100	**Low discrimination**						**67651100**
					GRA		
Fusobacterium mortiferum		88.64	0.58	NIT 1	1		
Actinomyces naeslundii		8.53	0.52		99		
Lactobacillus catenaformis		1.13	0.15		99		
67651101	**Acceptable identification**						**67651101**
					GRA		
Actinomyces naeslundii		93.46	0.55	ESL 12	99		
Fusobacterium mortiferum		6.54	0.24	MLT 94	1		
67651102	**Low discrimination**						**67651102**
					GRA		
Fusobacterium mortiferum		35.04	0.41	NIT 1	1		
Actinomyces naeslundii		30.35	0.52	BGL 10	99		
Lactobacillus acidophilus		8.30	0.18		99		
67651103	**Good identification**						**67651103**
Actinomyces naeslundii		98.38	0.55	ESL 12			
Actinomyces viscosus		1.62	0.00	MLT 94			
67651120	**Low discrimination**						**67651120**
					SPO		
Actinomyces naeslundii		53.11	0.52	ESL 12	1		
Clostridium paraputrificum		19.49	0.13	MLT 94	99		
Clostridium barati		8.76	0.12		99		
67651121	**Good identification**						**67651121**
Actinomyces naeslundii		100.0	0.55	ESL 12			
67651123	**Good identification**						**67651123**
					GRA		
Actinomyces naeslundii		98.61	0.55	ESL 12	99		
Prevotella oralis		1.39	-.03	MLT 94	1		
67651140	**Low discrimination**						**67651140**
					GRA		
Actinomyces naeslundii		57.39	0.39	ESL 12	99		
Fusobacterium mortiferum		34.13	0.24	MLT 94	1		
Bifidobacterium breve		8.48	0.10		99		
67651141	**Good identification**						**67651141**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67651143	**Good identification**						**67651143**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67651160	**Low discrimination**						**67651160**
					SPO		
Actinomyces naeslundii		70.45	0.39	ESL 12	1		
Bifidobacterium breve		18.52	0.15	MLT 94	1		
Clostridium paraputrificum		11.03	-.07		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67651161	**Good identification**						**67651161**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67651163	**Good identification**						**67651163**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67652101	**Good identification**						**67652101**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67652103	**Good identification**						**67652103**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67652121	**Good identification**						**67652121**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67652123	**Good identification**						**67652123**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67653000	**Low discrimination**						**67653000**
					GRA	SPO	
Actinomyces naeslundii		55.84	0.49	ESL 12	99	1	
Clostridium innocuum		36.96	0.22	MLT 94	99	99	
Fusobacterium mortiferum		3.68	0.17		1	1	
67653001	**Good identification**						**67653001**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67653002	**Low discrimination**						**67653002**
					SPO		
Actinomyces naeslundii		56.94	0.49	ESL 12	1		
Clostridium innocuum		37.68	0.22	MLT 94	99		
Actinomyces israelii		5.38	0.22		1		
67653003	**Good identification**						**67653003**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67653020	**Good identification**						**67653020**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67653021	**Good identification**						**67653021**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67653022	**Good identification**						**67653022**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67653023	**Good identification**						**67653023**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67653040	**Good identification**						**67653040**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67653041	**Good identification**						**67653041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.39	ESL 12		
67653042	**Low discrimination**					**67653042**
Actinomyces naeslundii		82.77	0.36	ESL 12		
Actinomyces israelii		17.23	0.15	MLT 94		
67653043	**Good identification**					**67653043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67653060	**Good identification**					**67653060**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67653061	**Good identification**					**67653061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67653062	**Good identification**					**67653062**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67653063	**Good identification**					**67653063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67653100	**Low discrimination**					**67653100**
					GRA	
Actinomyces naeslundii		86.51	0.57	ESL 12	99	
Actinomyces israelii		7.66	0.30	MLT 94	99	
Fusobacterium mortiferum		4.68	0.24		1	
67653101	**Very good identification**					**67653101**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67653102	**Low discrimination**					**67653102**
Actinomyces naeslundii		78.82	0.57	ESL 12		
Bifidobacterium breve		10.71	0.27	MLT 94		
Actinomyces israelii		10.46	0.33			
67653103	**Very good identification**					**67653103**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67653111	**Good identification**					**67653111**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67653113	**Good identification**					**67653113**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67653120	**Very good identification**					**67653120**
					SPO	
Actinomyces naeslundii		96.50	0.57	ESL 12	1	
Bifidobacterium breve		2.31	0.15	MLT 94	1	
Clostridium ramosum		1.19	-.01		92	
67653121	**Very good identification**					**67653121**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67653122	**Low discrimination**					**67653122**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					SPO		
Actinomyces naeslundii		75.53	0.57	ESL 12	1		
Bifidobacterium breve		18.25	0.32	MLT 94	1		
Clostridium ramosum		3.72	0.09		92		
67653123	**Very good identification**						**67653123**
Actinomyces naeslundii		100.0	0.60	ESL 12			
67653131	**Good identification**						**67653131**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67653133	**Good identification**						**67653133**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67653140	**Low discrimination**						**67653140**
Actinomyces naeslundii		78.67	0.44	ESL 12			
Actinomyces israelii		15.34	0.23	MLT 94			
Bifidobacterium breve		5.99	0.10				
67653141	**Good identification**						**67653141**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67653142	**Low discrimination**						**67653142**
Actinomyces naeslundii		48.48	0.44	ESL 12			
Bifidobacterium breve		37.34	0.27	MLT 94			
Actinomyces israelii		14.18	0.26				
67653143	**Good identification**						**67653143**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67653160	**Low discrimination**						**67653160**
Actinomyces naeslundii		88.07	0.44	ESL 12			
Bifidobacterium breve		11.93	0.15	MLT 94			
67653161	**Good identification**						**67653161**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67653163	**Good identification**						**67653163**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67654100	**Good identification**						**67654100**
Fusobacterium mortiferum		100.0	0.35	NIT 1			
67655100	**Low discrimination**						**67655100**
					GRA	SPO	
Fusobacterium mortiferum		67.53	0.38	NIT 1	1	1	
Actinomyces israelii		30.10	0.35	XYL 7	99	1	
Clostridium butyricum		2.37	-.05		99	99	
67655102	**Low discrimination**						**67655102**
					GRA		
Actinomyces israelii		77.48	0.38	MLT 99	99		
Fusobacterium mortiferum		12.88	0.22		1		
Bifidobacterium adolescentis		9.64	0.16		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67655322	**Low discrimination**						**67655322**
Parabacteroides distasonis		87.21	0.41	NIT 1			
Prevotella oralis		7.74	0.10				
Bacteroides uniformis		5.06	0.01				
67655332	**Good identification**						**67655332**
Parabacteroides distasonis		100.0	0.36	NIT 1			
67657000	**Good identification**						**67657000**
					SPO		
Actinomyces israelii		95.41	0.37	MLT 99	1		
Clostridium innocuum		4.59	0.00		99		
67657002	**Good identification**						**67657002**
					SPO		
Actinomyces israelii		96.89	0.40	MLT 99	1		
Clostridium innocuum		3.11	0.00		99		
67657100	**Good identification**						**67657100**
Actinomyces israelii		99.02	0.48	MLT 99			
Actinomyces naeslundii		0.98	0.23				
67657102	**Good identification**						**67657102**
Actinomyces israelii		97.60	0.51	MLT 99			
Bifidobacterium adolescentis		1.75	0.15				
Actinomyces naeslundii		0.65	0.23				
67657112	**Low discrimination**						**67657112**
Actinomyces israelii		87.18	0.37	MLT 99			
Bifidobacterium adolescentis		12.82	0.16	MLZ 13			
67657140	**Good identification**						**67657140**
Actinomyces israelii		100.0	0.41	MLT 99			
67657142	**Good identification**						**67657142**
Actinomyces israelii		100.0	0.44	MLT 99			
67657400	**Good identification**						**67657400**
Actinomyces israelii		100.0	0.36	MLT 99			
67657402	**Good identification**						**67657402**
Actinomyces israelii		100.0	0.39	MLT 99			
67657500	**Low discrimination**						**67657500**
					GRA	SPO	
Actinomyces israelii		69.77	0.46	MLT 99	99	1	
Mitsuokella multacida		29.05	0.27		1	1	
Clostridium butyricum		1.19	-.05		99	99	
67657502	**Low discrimination**						**67657502**
					GRA		
Actinomyces israelii		84.90	0.49	MLT 99	99		
Bifidobacterium adolescentis		12.48	0.29		99		
Mitsuokella multacida		2.62	0.11		1		
67657542	**Low discrimination**						**67657542**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces israelii		88.81	0.42	MLT 99	99	
Mitsuokella multacida		7.04	0.11		1	
Bifidobacterium adolescentis		4.15	0.13		99	
67661101	**Good identification**					**67661101**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67661103	**Good identification**					**67661103**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67661121	**Good identification**					**67661121**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67661123	**Good identification**					**67661123**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67663001	**Good identification**					**67663001**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67663003	**Good identification**					**67663003**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67663021	**Good identification**					**67663021**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67663023	**Good identification**					**67663023**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67663100	**Good identification**					**67663100**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67663101	**Good identification**					**67663101**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67663102	**Good identification**					**67663102**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67663103	**Good identification**					**67663103**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67663120	**Good identification**					**67663120**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67663121	**Good identification**					**67663121**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67663122	**Good identification**					**67663122**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67663123	**Good identification**					**67663123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ESL 12		
67670000	**Good identification**					**67670000**
					GRA	
Fusobacterium mortiferum		96.90	0.39	NIT 1	1	
Actinomyces naeslundii		3.11	0.25		99	
67670100	**Good identification**					**67670100**
					GRA	
Fusobacterium mortiferum		96.24	0.46	NIT 1	1	
Actinomyces naeslundii		3.76	0.33		99	
67670101	**Low discrimination**					**67670101**
					GRA	
Actinomyces naeslundii		85.31	0.36	ESL 12	99	
Fusobacterium mortiferum		14.69	0.12	MLT 94	1	
67670103	**Good identification**					**67670103**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67670121	**Good identification**					**67670121**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67670123	**Good identification**					**67670123**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67671000	**Low discrimination**					**67671000**
					GRA	
Fusobacterium mortiferum		63.22	0.42	NIT 1	1	
Actinomyces naeslundii		32.42	0.49		99	
Lactobacillus jensenii		3.01	0.10		99	
67671001	**Good identification**					**67671001**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67671002	**Low discrimination**					**67671002**
					GRA	
Actinomyces naeslundii		68.59	0.49	ESL 12	99	
Fusobacterium mortiferum		14.86	0.26	MLT 94	1	
Lactobacillus acidophilus		8.01	0.09		99	
67671003	**Good identification**					**67671003**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67671020	**Good identification**					**67671020**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67671021	**Good identification**					**67671021**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67671022	**Low discrimination**					**67671022**
Actinomyces naeslundii		89.54	0.49	ESL 12		
Lactobacillus acidophilus		10.46	0.09	MLT 94		
67671023	**Good identification**					**67671023**
Actinomyces naeslundii		100.0	0.52	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67671040	**Good identification**					**67671040**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67671041	**Good identification**					**67671041**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67671042	**Good identification**					**67671042**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67671043	**Good identification**					**67671043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67671060	**Good identification**					**67671060**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67671061	**Good identification**					**67671061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67671062	**Good identification**					**67671062**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67671063	**Good identification**					**67671063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67671100	**Low discrimination**					**67671100**
					GRA	
Fusobacterium mortiferum		60.16	0.49	NIT 1	1	
Actinomyces naeslundii		37.63	0.57		99	
Actinomyces israelii		2.21	0.27		99	
67671101	**Very good identification**					**67671101**
					GRA	
Actinomyces naeslundii		98.94	0.60	ESL 12	99	
Fusobacterium mortiferum		1.07	0.15	MLT 94	1	
67671102	**Low discrimination**					**67671102**
					GRA	
Actinomyces naeslundii		51.18	0.57	ESL 12	99	
Lactobacillus acidophilus		24.14	0.27	MLT 94	99	
Fusobacterium mortiferum		9.09	0.32		1	
67671103	**Very good identification**					**67671103**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67671111	**Good identification**					**67671111**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67671112	**Good identification**					**67671112**
					GRA	
Leptotrichia buccalis		98.76	0.41	NIT 10	1	
Bifidobacterium breve		0.72	0.24	MLT 99	99	
Actinomyces naeslundii		0.42	0.35		99	
67671120	**Low discrimination**					**67671120**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		88.73	0.57	ESL 12	1	
Clostridium paraputrificum		4.46	0.03	MLT 94	99	
Clostridium ramosum		3.02	0.06		92	
67671121	**Very good identification**					**67671121**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67671122	**Low discrimination**					**67671122**
Actinomyces naeslundii		49.88	0.57	ESL 12		
Lactobacillus acidophilus		23.53	0.27	MLT 94		
Bifidobacterium breve		8.04	0.29			
67671123	**Very good identification**					**67671123**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67671131	**Good identification**					**67671131**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67671133	**Low discrimination**					**67671133**
					GRA	
Actinomyces naeslundii		84.95	0.38	ESL 12	99	
Leptotrichia buccalis		15.05	-.09	MLT 94	1	
67671140	**Low discrimination**					**67671140**
					GRA	
Actinomyces naeslundii		81.90	0.44	ESL 12	99	
Actinomyces israelii		10.60	0.20	MLT 94	99	
Fusobacterium mortiferum		7.50	0.15		1	
67671141	**Good identification**					**67671141**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67671142	**Low discrimination**					**67671142**
Actinomyces naeslundii		49.06	0.44	ESL 12		
Bifidobacterium breve		25.19	0.24	MLT 94		
Lactobacillus acidophilus		16.21	0.11			
67671143	**Good identification**					**67671143**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67671160	**Acceptable identification**					**67671160**
Actinomyces naeslundii		91.72	0.44	ESL 12		
Bifidobacterium breve		8.28	0.12	MLT 94		
67671161	**Good identification**					**67671161**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67671162	**Low discrimination**					**67671162**
Actinomyces naeslundii		44.58	0.44	ESL 12		
Bifidobacterium breve		40.69	0.29	MLT 94		
Lactobacillus acidophilus		14.73	0.11			
67671163	**Good identification**					**67671163**
Actinomyces naeslundii		100.0	0.47	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67671322	**Low discrimination**						**67671322**
					SPO	GRA	
Parabacteroides distasonis		56.09	0.36	NIT 1	1	1	
Clostridium ramosum		31.01	0.19	XYL 90	92	50	
Lactobacillus acidophilus		6.09	0.08		1	99	
67672100	**Low discrimination**						**67672100**
					GRA		
Actinomyces naeslundii		88.25	0.38	ESL 12	99		
Fusobacterium mortiferum		11.75	0.12	MLT 94	1		
67672101	**Good identification**						**67672101**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67672102	**Good identification**						**67672102**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67672103	**Good identification**						**67672103**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67672120	**Good identification**						**67672120**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67672121	**Good identification**						**67672121**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67672122	**Good identification**						**67672122**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67672123	**Good identification**						**67672123**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67673000	**Low discrimination**						**67673000**
					SPO		
Actinomyces naeslundii		80.53	0.54	ESL 12	1		
Actinomyces israelii		9.83	0.29	MLT 94	1		
Clostridium innocuum		9.65	0.14		99		
67673001	**Very good identification**						**67673001**
Actinomyces naeslundii		100.0	0.57	ESL 12			
67673002	**Low discrimination**						**67673002**
					SPO		
Actinomyces naeslundii		76.76	0.54	ESL 12	1		
Actinomyces israelii		14.05	0.32	MLT 94	1		
Clostridium innocuum		9.19	0.14		99		
67673003	**Very good identification**						**67673003**
Actinomyces naeslundii		100.0	0.57	ESL 12			
67673011	**Good identification**						**67673011**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67673013	**Good identification**						**67673013**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.35	ESL 12		
67673020	**Good identification**					**67673020**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67673021	**Very good identification**					**67673021**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67673022	**Good identification**					**67673022**
Actinomyces naeslundii		100.0	0.54	ESL 12		
67673023	**Very good identification**					**67673023**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67673031	**Good identification**					**67673031**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67673033	**Good identification**					**67673033**
Actinomyces naeslundii		100.0	0.35	ESL 12		
67673040	**Low discrimination**					**67673040**
Actinomyces naeslundii		78.81	0.41	ESL 12		
Actinomyces israelii		21.19	0.22	MLT 94		
67673041	**Good identification**					**67673041**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67673042	**Low discrimination**					**67673042**
Actinomyces naeslundii		71.26	0.41	ESL 12		
Actinomyces israelii		28.75	0.25	MLT 94		
67673043	**Good identification**					**67673043**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67673060	**Good identification**					**67673060**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67673061	**Good identification**					**67673061**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67673062	**Good identification**					**67673062**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67673063	**Good identification**					**67673063**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67673100	**Low discrimination**					**67673100**
					GRA	
Actinomyces naeslundii		84.76	0.62	ESL 12	99	
Actinomyces israelii		14.54	0.39	MLT 94	99	
Fusobacterium mortiferum		0.71	0.15		1	
67673101	**Very good identification**					**67673101**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.65	ESL 12		
67673102	**Low discrimination**					**67673102**
Actinomyces naeslundii		75.95	0.62	ESL 12		
Actinomyces israelii		19.54	0.42	MLT 94		
Bifidobacterium breve		3.55	0.24			
67673103	**Very good identification**					**67673103**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67673110	**Low discrimination**					**67673110**
Actinomyces naeslundii		67.25	0.40	ESL 12		
Actinomyces israelii		32.75	0.25	MLT 94		
67673111	**Good identification**					**67673111**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67673113	**Good identification**					**67673113**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67673120	**Low discrimination**					**67673120**
					SPO	
Actinomyces naeslundii		89.40	0.62	ESL 12	1	
Clostridium ramosum		8.89	0.19	MLT 94	92	
Actinomyces israelii		0.98	0.19		1	
67673121	**Very good identification**					**67673121**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67673122	**Low discrimination**					**67673122**
					SPO	
Actinomyces naeslundii		66.22	0.62	ESL 12	1	
Clostridium ramosum		26.35	0.30	MLT 94	92	
Bifidobacterium breve		5.50	0.29		1	
67673123	**Very good identification**					**67673123**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67673130	**Low discrimination**					**67673130**
Actinomyces naeslundii		86.51	0.40	ESL 12		
Bifidobacterium breve		13.49	0.12	MLT 94		
67673131	**Good identification**					**67673131**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67673133	**Good identification**					**67673133**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67673140	**Low discrimination**					**67673140**
Actinomyces naeslundii		72.57	0.49	ESL 12		
Actinomyces israelii		27.43	0.32	MLT 94		
67673141	**Good identification**					**67673141**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.52	ESL 12			
67673142	**Low discrimination**						**67673142**
Actinomyces naeslundii		54.60	0.49	ESL 12			
Actinomyces israelii		30.96	0.35	MLT 94			
Bifidobacterium breve		14.44	0.24				
67673143	**Good identification**						**67673143**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67673160	**Good identification**						**67673160**
Actinomyces naeslundii		95.56	0.49	ESL 12			
Bifidobacterium breve		4.44	0.12	MLT 94			
67673161	**Good identification**						**67673161**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67673162	**Low discrimination**						**67673162**
Actinomyces naeslundii		66.38	0.49	ESL 12			
Bifidobacterium breve		31.22	0.29	MLT 94			
Actinomyces israelii		2.40	0.15				
67673163	**Good identification**						**67673163**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67673502	**Low discrimination**						**67673502**
					GRA		
Actinomyces israelii		84.24	0.41	MLT 99	99		
Mitsuokella multacida		7.94	0.11	XYL 92	1		
Actinomyces naeslundii		4.04	0.28		99		
67675002	**Good identification**						**67675002**
Actinomyces israelii		100.0	0.37	MLT 99			
67675100	**Low discrimination**						**67675100**
					GRA	SPO	
Actinomyces israelii		57.43	0.45	MLT 99	99	1	
Clostridium butyricum		18.86	0.15		99	99	
Mitsuokella multacida		12.63	0.21		1	1	
67675102	**Low discrimination**						**67675102**
					SPO		
Actinomyces israelii		86.47	0.48	MLT 99	1		
Clostridium butyricum		7.36	0.08		99		
Bifidobacterium adolescentis		1.91	0.13		1		
67675122	**Acceptable identification**						**67675122**
					GRA		
Parabacteroides distasonis		92.91	0.44	NIT 1	1		
Actinomyces israelii		6.13	0.27		99		
Actinomyces naeslundii		0.96	0.23		99		
67675132	**Good identification**						**67675132**
					GRA		
Parabacteroides distasonis		95.71	0.39	NIT 1	1		
Leptotrichia buccalis		2.54	-.09		1		
Actinomyces israelii		1.75	0.13		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67675140	**Low discrimination**						**67675140**
					GRA	SPO	
Actinomyces israelii		61.02	0.38	MLT 99	99	1	
Mitsuokella multacida		34.50	0.21		1	1	
Clostridium butyricum		4.48	-.03		99	99	
67675142	**Good identification**						**67675142**
					GRA		
Actinomyces israelii		95.98	0.41	MLT 99	99		
Mitsuokella multacida		4.02	0.04		1		
67675322	**Good identification**						**67675322**
Parabacteroides distasonis		100.0	0.52	NIT 1			
67675332	**Good identification**						**67675332**
Parabacteroides distasonis		98.34	0.47	NIT 1			
Clostridium clostridioforme		1.66	0.11				
67675402	**Low discrimination**						**67675402**
					GRA	SPO	
Actinomyces israelii		80.92	0.36	MLT 99	99	1	
Clostridium butyricum		12.63	0.01		99	99	
Mitsuokella multacida		6.44	0.04		1	1	
67675500	**Low discrimination**						**67675500**
					GRA	SPO	
Mitsuokella multacida		85.23	0.55	MLT 99	1	1	
Clostridium butyricum		11.57	0.32		99	99	
Actinomyces israelii		3.20	0.43		99	1	
67675501	**Good identification**						**67675501**
					GRA	SPO	
Mitsuokella multacida		98.78	0.39	MLT 99	1	1	
Clostridium butyricum		1.22	-.03	URE 10	99	99	
67675502	**Low discrimination**						**67675502**
					GRA	SPO	
Mitsuokella multacida		48.04	0.39	MLT 99	1	1	
Actinomyces israelii		24.37	0.46	BGL 10	99	1	
Clostridium butyricum		22.83	0.25		99	99	
67675510	**Good identification**						**67675510**
					GRA	SPO	
Mitsuokella multacida		96.31	0.47	MLT 99	1	1	
Clostridium butyricum		2.07	0.10		99	99	
Actinomyces israelii		1.62	0.29		99	1	
67675520	**Acceptable identification**						**67675520**
					GRA	SPO	
Mitsuokella multacida		94.62	0.45	MLT 99	1	1	
Clostridium butyricum		4.47	0.14		99	99	
Actinomyces israelii		0.91	0.23		99	1	
67675522	**Low discrimination**						**67675522**
					GRA	SPO	
Parabacteroides distasonis		52.60	0.40	NIT 1	1	1	
Mitsuokella multacida		35.76	0.28		1	1	
Clostridium butyricum		5.91	0.07		99	99	
67675530	**Good identification**						**67675530**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.37	MLT 99			
67675532	**Low discrimination**						**67675532**
Parabacteroides distasonis		59.03	0.35	NIT 1			
Mitsuokella multacida		24.85	0.20				
Clostridium clostridioforme		10.42	0.17				
67675540	**Good identification**						**67675540**
					GRA	SPO	
Mitsuokella multacida		97.43	0.55	MLT 99	1	1	
Actinomyces israelii		1.42	0.36		99	1	
Clostridium butyricum		1.15	0.14		99	99	
67675541	**Good identification**						**67675541**
Mitsuokella multacida		100.0	0.39	MLT 99			
67675542	**Low discrimination**						**67675542**
					GRA	SPO	
Mitsuokella multacida		80.00	0.39	MLT 99	1	1	
Actinomyces israelii		15.79	0.39	BGL 10	99	1	
Clostridium butyricum		3.31	0.07		99	99	
67675550	**Good identification**						**67675550**
					COC	GRA	
Mitsuokella multacida		99.00	0.47	MLT 99	1	1	
Actinomyces israelii		0.65	0.22		1	99	
Blautia producta		0.35	-.03		99	99	
67675560	**Good identification**						**67675560**
					GRA	SPO	
Mitsuokella multacida		99.22	0.45	MLT 99	1	1	
Clostridium butyricum		0.41	-.05		99	99	
Actinomyces israelii		0.37	0.16		99	1	
67675570	**Good identification**						**67675570**
Mitsuokella multacida		100.0	0.37	MLT 99			
67675700	**Very good identification**						**67675700**
					GRA	SPO	
Mitsuokella multacida		99.62	0.60	MLT 99	1	1	
Clostridium butyricum		0.23	0.06		99	99	
Actinomyces israelii		0.15	0.24		99	1	
67675701	**Good identification**						**67675701**
Mitsuokella multacida		100.0	0.43	MLT 99			
67675702	**Acceptable identification**						**67675702**
					GRA		
Mitsuokella multacida		93.92	0.43	MLT 99	1		
Clostridium clostridioforme		2.84	0.20	BGL 10	50		
Actinomyces israelii		1.93	0.27		99		
67675710	**Good identification**						**67675710**
Mitsuokella multacida		100.0	0.51	MLT 99			
67675711	**Good identification**						**67675711**
Mitsuokella multacida		100.0	0.35	MLT 99			
67675712	**Low discrimination**						**67675712**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		70.25	0.35	MLT 99			
Clostridium clostridioforme		29.07	0.32	BGL 10			
Parabacteroides distasonis		0.68	0.09				
67675720	**Good identification**						**67675720**
Mitsuokella multacida		99.73	0.49	MLT 99			
Parabacteroides distasonis		0.27	0.14				
67675722	**Low discrimination**						**67675722**
Parabacteroides distasonis		66.18	0.48	NIT 1			
Mitsuokella multacida		27.85	0.33				
Clostridium clostridioforme		3.37	0.20				
67675730	**Good identification**						**67675730**
Mitsuokella multacida		99.57	0.41	MLT 99			
Parabacteroides distasonis		0.43	0.09				
67675732	**Low discrimination**						**67675732**
Parabacteroides distasonis		57.89	0.44	NIT 1			
Clostridium clostridioforme		24.97	0.32				
Mitsuokella multacida		15.08	0.25				
67675740	**Very good identification**						**67675740**
					GRA		
Mitsuokella multacida		99.94	0.60	MLT 99	1		
Actinomyces israelii		0.06	0.17		99		
67675741	**Good identification**						**67675741**
Mitsuokella multacida		100.0	0.43	MLT 99			
67675742	**Good identification**						**67675742**
					GRA		
Mitsuokella multacida		99.21	0.43	MLT 99	1		
Actinomyces israelii		0.79	0.20	BGL 10	99		
67675750	**Good identification**						**67675750**
					COC	GRA	
Mitsuokella multacida		99.81	0.51	MLT 99	1	1	
Blautia producta		0.19	-.03		99	99	
67675751	**Good identification**						**67675751**
Mitsuokella multacida		100.0	0.35	MLT 99			
67675752	**Good identification**						**67675752**
Mitsuokella multacida		100.0	0.35	MLT 99			
67675760	**Good identification**						**67675760**
Mitsuokella multacida		100.0	0.49	MLT 99			
67675770	**Good identification**						**67675770**
Mitsuokella multacida		100.0	0.41	MLT 99			
67677000	**Good identification**						**67677000**
Actinomyces israelii		100.0	0.47	MLT 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67677002	**Good identification**						**67677002**
Actinomyces israelii		100.0	0.50	MLT 99			
67677012	**Good identification**						**67677012**
Actinomyces israelii		100.0	0.36	MLT 99			
67677040	**Good identification**						**67677040**
Actinomyces israelii		100.0	0.40	MLT 99			
67677042	**Good identification**						**67677042**
Actinomyces israelii		100.0	0.43	MLT 99			
67677100	**Good identification**						**67677100**
					GRA	SPO	
Actinomyces israelii		90.74	0.58	MLT 99	99	1	
Mitsuokella multacida		8.22	0.27		1	1	
Clostridium butyricum		0.58	-.01		99	99	
67677102	**Very good identification**						**67677102**
					GRA		
Actinomyces israelii		98.69	0.61	MLT 99	99		
Mitsuokella multacida		0.66	0.11		1		
Actinomyces naeslundii		0.34	0.28		99		
67677110	**Low discrimination**						**67677110**
					GRA		
Actinomyces israelii		83.20	0.43	MLT 99	99		
Mitsuokella multacida		16.80	0.19	MLZ 13	1		
67677112	**Acceptable identification**						**67677112**
Actinomyces israelii		94.21	0.46	MLT 99			
Bifidobacterium adolescentis		2.46	0.13	MLZ 13			
Bifidobacterium dentium		1.93	-.07				
67677120	**Low discrimination**						**67677120**
					GRA		
Actinomyces israelii		69.69	0.37	MLT 99	99		
Mitsuokella multacida		24.72	0.17	NAG 6	1		
Actinomyces naeslundii		5.59	0.28		99		
67677122	**Acceptable identification**						**67677122**
					GRA		
Actinomyces israelii		92.55	0.40	MLT 99	99		
Actinomyces naeslundii		4.95	0.28	NAG 6	99		
Parabacteroides distasonis		2.50	0.10		1		
67677140	**Low discrimination**						**67677140**
					GRA		
Actinomyces israelii		81.11	0.51	MLT 99	99		
Mitsuokella multacida		18.89	0.27		1		
67677142	**Good identification**						**67677142**
					GRA		
Actinomyces israelii		98.30	0.54	MLT 99	99		
Mitsuokella multacida		1.70	0.11		1		
67677150	**Low discrimination**						**67677150**
					COC	GRA	
Actinomyces israelii		62.61	0.36	MLT 99	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		32.52	0.19	MLZ 13	1	1	
Blautia producta		4.88	-.03		99	99	
67677152	**Good identification**						**67677152**
					GRA		
Actinomyces israelii		96.29	0.39	MLT 99	99		
Mitsuokella multacida		3.71	0.03	MLZ 13	1		
67677302	**Low discrimination**						**67677302**
					GRA		
Actinomyces israelii		85.80	0.41	MLT 99	99		
Mitsuokella multacida		14.20	0.15	RHA 7	1		
67677400	**Low discrimination**						**67677400**
					GRA		
Actinomyces israelii		69.32	0.46	MLT 99	99		
Mitsuokella multacida		30.68	0.27		1		
67677402	**Good identification**						**67677402**
					GRA		
Actinomyces israelii		96.83	0.49	MLT 99	99		
Mitsuokella multacida		3.18	0.11		1		
67677442	**Acceptable identification**						**67677442**
					GRA		
Actinomyces israelii		92.22	0.42	MLT 99	99		
Mitsuokella multacida		7.78	0.11		1		
67677500	**Good identification**						**67677500**
					GRA	SPO	
Mitsuokella multacida		91.10	0.61	MLT 99	1	1	
Actinomyces israelii		8.32	0.56		99	1	
Clostridium butyricum		0.59	0.15		99	99	
67677501	**Good identification**						**67677501**
					GRA		
Mitsuokella multacida		99.18	0.45	MLT 99	1		
Actinomyces israelii		0.82	0.22	URE 10	99		
67677502	**Low discrimination**						**67677502**
					GRA		
Actinomyces israelii		53.77	0.59	MLT 99	99		
Mitsuokella multacida		43.64	0.45		1		
Bifidobacterium adolescentis		1.40	0.26		99		
67677510	**Acceptable identification**						**67677510**
					GRA		
Mitsuokella multacida		94.47	0.53	MLT 99	1		
Actinomyces israelii		3.87	0.42		99		
Bifidobacterium dentium		1.42	0.10		99		
67677511	**Good identification**						**67677511**
Mitsuokella multacida		100.0	0.37	MLT 99			
67677520	**Good identification**						**67677520**
					GRA	SPO	
Mitsuokella multacida		97.51	0.51	MLT 99	1	1	
Actinomyces israelii		2.27	0.36		99	1	
Clostridium butyricum		0.22	-.03		99	99	
67677530	**Low discrimination**						**67677530**
					GRA		
Mitsuokella multacida		89.49	0.43	MLT 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		9.58	0.15		99		
Actinomyces israelii		0.94	0.22		99		
67677540	**Good identification**						**67677540**
					COC	GRA	
Mitsuokella multacida		94.99	0.61	MLT 99	1	1	
Actinomyces israelii		3.37	0.49		1	99	
Blautia producta		1.58	0.24		99	99	
67677541	**Good identification**						**67677541**
					GRA		
Mitsuokella multacida		99.68	0.45	MLT 99	1		
Actinomyces israelii		0.32	0.15	URE 10	99		
67677542	**Low discrimination**						**67677542**
					COC	GRA	
Mitsuokella multacida		66.68	0.45	MLT 99	1	1	
Actinomyces israelii		31.95	0.52	BGL 10	1	99	
Blautia producta		1.11	0.07		99	99	
67677550	**Low discrimination**						**67677550**
					COC	GRA	
Mitsuokella multacida		85.77	0.53	MLT 99	1	1	
Blautia producta		12.87	0.32		99	99	
Actinomyces israelii		1.37	0.35		1	99	
67677551	**Good identification**						**67677551**
					COC	GRA	
Mitsuokella multacida		98.66	0.37	MLT 99	1	1	
Blautia producta		1.35	-.03	URE 10	99	99	
67677552	**Low discrimination**						**67677552**
					COC	GRA	
Mitsuokella multacida		71.78	0.37	MLT 99	1	1	
Actinomyces israelii		15.42	0.38	BGL 10	1	99	
Blautia producta		10.77	0.15		99	99	
67677560	**Good identification**						**67677560**
					COC	GRA	
Mitsuokella multacida		98.38	0.51	MLT 99	1	1	
Actinomyces israelii		0.89	0.29		1	99	
Blautia producta		0.73	0.07		99	99	
67677570	**Acceptable identification**						**67677570**
					COC	GRA	
Mitsuokella multacida		93.39	0.43	MLT 99	1	1	
Blautia producta		6.23	0.15		99	99	
Actinomyces israelii		0.38	0.15		1	99	
67677700	**Very good identification**						**67677700**
					GRA		
Mitsuokella multacida		99.63	0.66	MLT 99	1		
Actinomyces israelii		0.37	0.37		99		
67677701	**Good identification**						**67677701**
Mitsuokella multacida		100.0	0.50	MLT 99			
67677702	**Good identification**						**67677702**
					GRA		
Mitsuokella multacida		95.24	0.50	MLT 99	1		
Actinomyces israelii		4.76	0.40	BGL 10	99		
67677710	**Very good identification**						**67677710**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Mitsuokella multacida		99.75	0.58	MLT 99	1	1	
Actinomyces israelii		0.17	0.23		1	99	
Blautia producta		0.08	-.03		99	99	
67677711	**Good identification**						**67677711**
Mitsuokella multacida		100.0	0.41	MLT 99			
67677712	**Good identification**						**67677712**
					GRA		
Mitsuokella multacida		97.10	0.41	MLT 99	1		
Actinomyces israelii		2.17	0.26	BGL 10	99		
Bifidobacterium dentium		0.72	-.07		99		
67677720	**Good identification**						**67677720**
					GRA		
Mitsuokella multacida		99.91	0.56	MLT 99	1		
Actinomyces israelii		0.09	0.16		99		
67677721	**Good identification**						**67677721**
Mitsuokella multacida		100.0	0.39	MLT 99			
67677722	**Good identification**						**67677722**
					GRA		
Mitsuokella multacida		96.83	0.39	MLT 99	1		
Actinomyces israelii		1.24	0.19	BGL 10	99		
Parabacteroides distasonis		1.00	0.14		1		
67677730	**Good identification**						**67677730**
Mitsuokella multacida		100.0	0.47	MLT 99			
67677740	**Very good identification**						**67677740**
					COC	GRA	
Mitsuokella multacida		98.97	0.66	MLT 99	1	1	
Blautia producta		0.89	0.24		99	99	
Actinomyces israelii		0.14	0.30		1	99	
67677741	**Good identification**						**67677741**
Mitsuokella multacida		100.0	0.50	MLT 99			
67677742	**Good identification**						**67677742**
					COC	GRA	
Mitsuokella multacida		97.24	0.50	MLT 99	1	1	
Actinomyces israelii		1.89	0.33	BGL 10	1	99	
Blautia producta		0.87	0.07		99	99	
67677750	**Good identification**						**67677750**
					COC	GRA	
Mitsuokella multacida		92.47	0.58	MLT 99	1	1	
Blautia producta		7.47	0.32		99	99	
Actinomyces israelii		0.06	0.16		1	99	
67677751	**Good identification**						**67677751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	MLT 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67677752	**Acceptable identification**						**67677752**
					COC	GRA	
Mitsuokella multacida		91.79	0.41	MLT 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		7.41	0.15	BGL 10	99	99	
Actinomyces israelii		0.80	0.19		1	99	
67677760	**Good identification**						**67677760**
					COC	GRA	
Mitsuokella multacida		99.60	0.56	MLT 99	1	1	
Blautia producta		0.40	0.07		99	99	
67677761	**Good identification**						**67677761**
Mitsuokella multacida		100.0	0.39	MLT 99			
67677762	**Good identification**						**67677762**
Mitsuokella multacida		100.0	0.39	MLT 99			
67677770	**Good identification**						**67677770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	MLT 99	1	1	
Blautia producta		3.47	0.15		99	99	
67701001	**Good identification**						**67701001**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67701003	**Low discrimination**						**67701003**
Actinomyces naeslundii		54.67	0.44	ESL 12			
Actinomyces viscosus		45.33	0.18	FRU 95			
67701020	**Low discrimination**						**67701020**
Actinomyces naeslundii		54.92	0.41	ESL 12			
Eubacterium tortuosum		35.94	0.09	FRU 95			
Clostridium perfringens		9.14	0.05				
67701021	**Good identification**						**67701021**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67701022	**Low discrimination**						**67701022**
					GRA		
Actinomyces naeslundii		65.10	0.41	ESL 12	99		
Prevotella oralis		11.26	0.02	FRU 95	1		
Bifidobacterium longum subsp. infantis		10.04	-.03		99		
67701023	**Good identification**						**67701023**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67701100	**Low discrimination**						**67701100**
					GRA		
Actinomyces naeslundii		71.37	0.48	ESL 12	99		
Bifidobacterium longum subsp. infantis		21.30	0.10	FRU 95	99		
Fusobacterium mortiferum		4.47	0.16		1		
67701101	**Low discrimination**						**67701101**
					SPO		
Actinomyces naeslundii		59.68	0.51	ESL 12	1		
Clostridium chauvoei		40.32	0.25	FRU 95	99		
67701103	**Low discrimination**						**67701103**
					SPO		
Actinomyces naeslundii		60.33	0.51	ESL 12	1		
Actinomyces viscosus		31.25	0.22	FRU 95	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium chauvoei		7.19	0.13		99		
67701120	**Low discrimination**						**67701120**
Actinomyces naeslundii		36.09	0.48	ESL 12			
Bifidobacterium longum subsp. infantis		19.15	0.15	FRU 95			
Clostridium perfringens		15.48	0.20				
67701121	**Low discrimination**						**67701121**
					SPO		
Actinomyces naeslundii		81.62	0.51	ESL 12	1		
Clostridium chauvoei		18.38	0.17	FRU 95	99		
67701123	**Acceptable identification**						**67701123**
					SPO		
Actinomyces naeslundii		91.10	0.51	ESL 12	1		
Clostridium chauvoei		3.62	0.04	FRU 95	99		
Bifidobacterium longum subsp. infantis		3.29	-.03		1		
67701140	**Good identification**						**67701140**
Actinomyces naeslundii		100.0	0.35	ESL 12			
67701141	**Good identification**						**67701141**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67701142	**Low discrimination**						**67701142**
					GRA	CAT	
Actinomyces naeslundii		71.43	0.35	ESL 12	99	25	
Capnocytophaga ochracea		16.23	-.01	FRU 95	1	90	
Bifidobacterium longum subsp. infantis		12.34	-.07		99	1	
67701143	**Good identification**						**67701143**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67701160	**Low discrimination**						**67701160**
					GRA	CAT	
Actinomyces naeslundii		59.17	0.35	ESL 12	99	25	
Clostridium perfringens		27.39	0.07	FRU 95	99	1	
Capnocytophaga ochracea		13.44	-.01		1	90	
67701161	**Good identification**						**67701161**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67701163	**Good identification**						**67701163**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67703000	**Good identification**						**67703000**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67703001	**Good identification**						**67703001**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67703002	**Good identification**						**67703002**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67703003	**Good identification**						**67703003**
Actinomyces naeslundii		100.0	0.49	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67703020	**Good identification**						**67703020**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67703021	**Good identification**						**67703021**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67703022	**Good identification**						**67703022**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67703023	**Good identification**						**67703023**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67703041	**Good identification**						**67703041**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67703043	**Good identification**						**67703043**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67703061	**Good identification**						**67703061**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67703063	**Good identification**						**67703063**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67703100	**Good identification**						**67703100**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67703101	**Very good identification**						**67703101**
Actinomyces naeslundii		100.0	0.56	ESL 12			
67703102	**Good identification**						**67703102**
Actinomyces naeslundii		96.92	0.53	ESL 12			
Actinomyces israelii		1.56	0.13	FRU 95			
Bifidobacterium longum subsp. infantis		1.52	-.07				
67703103	**Very good identification**						**67703103**
Actinomyces naeslundii		100.0	0.56	ESL 12			
67703120	**Good identification**						**67703120**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67703121	**Very good identification**						**67703121**
Actinomyces naeslundii		100.0	0.56	ESL 12			
67703122	**Good identification**						**67703122**
					GRA	CAT	
Actinomyces naeslundii		95.71	0.53	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		2.67	-.03	FRU 95	99	1	
Capnocytophaga ochracea		1.62	-.03		1	90	
67703123	**Very good identification**						**67703123**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Actinomyces naeslundii	100.0	0.56	ESL 12		
67703140	**Good identification**					**67703140**
	Actinomyces naeslundii	100.0	0.40	ESL 12		
67703141	**Good identification**					**67703141**
	Actinomyces naeslundii	100.0	0.43	ESL 12		
67703142	**Good identification**					**67703142**
	Actinomyces naeslundii	100.0	0.40	ESL 12		
67703143	**Good identification**					**67703143**
	Actinomyces naeslundii	100.0	0.43	ESL 12		
67703160	**Good identification**					**67703160**
	Actinomyces naeslundii	100.0	0.40	ESL 12		
67703161	**Good identification**					**67703161**
	Actinomyces naeslundii	100.0	0.43	ESL 12		
67703162	**Good identification**					**67703162**
	Actinomyces naeslundii	100.0	0.40	ESL 12		
67703163	**Good identification**					**67703163**
	Actinomyces naeslundii	100.0	0.43	ESL 12		
67710000	**Low discrimination**					**67710000**
					GRA	
	Fusobacterium mortiferum	67.37	0.41	NIT 1	1	
	Actinomyces viscosus	15.63	0.18	MLT 11	99	
	Actinomyces naeslundii	10.59	0.39		99	
67710001	**Low discrimination**					**67710001**
	Actinomyces naeslundii	75.30	0.42	ESL 12		
	Actinomyces viscosus	24.70	0.09	GAL 96		
67710002	**Good identification**					**67710002**
					GRA	
	Actinomyces viscosus	98.53	0.52	ESL 1	99	
	Actinomyces naeslundii	0.67	0.39		99	
	Fusobacterium mortiferum	0.48	0.25		1	
67710003	**Good identification**					**67710003**
	Actinomyces viscosus	97.01	0.44	ESL 1		
	Actinomyces naeslundii	2.99	0.42			
67710020	**Low discrimination**					**67710020**
	Actinomyces naeslundii	82.63	0.39	ESL 12		
	Clostridium perfringens	17.37	0.05	GAL 96		
67710021	**Good identification**					**67710021**
	Actinomyces naeslundii	100.0	0.42	ESL 12		
67710023	**Low discrimination**					**67710023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		75.30	0.42	ESL 12		
Actinomyces viscosus		24.70	0.09	GAL 96		
67710100	**Low discrimination**					**67710100**
					GRA	
Fusobacterium mortiferum		68.34	0.47	NIT 1	1	
Actinomyces naeslundii		13.10	0.47	MLT 11	99	
Actinomyces viscosus		12.08	0.22		99	
67710101	**Low discrimination**					**67710101**
					GRA	
Actinomyces naeslundii		80.65	0.50	ESL 12	99	
Actinomyces viscosus		16.52	0.14	GAL 96	99	
Fusobacterium mortiferum		2.83	0.13		1	
67710102	**Acceptable identification**					**67710102**
Actinomyces viscosus		93.88	0.56	ESL 1		
Bifidobacterium longum subsp. infantis		3.92	0.27			
Actinomyces naeslundii		1.03	0.47			
67710103	**Good identification**					**67710103**
Actinomyces viscosus		95.19	0.48	ESL 1		
Actinomyces naeslundii		4.69	0.50			
Bifidobacterium longum subsp. infantis		0.12	-.07			
67710120	**Low discrimination**					**67710120**
Actinomyces naeslundii		35.34	0.47	ESL 12		
Bifidobacterium longum subsp. infantis		23.69	0.15	GAL 96		
Clostridium perfringens		19.14	0.20			
67710121	**Good identification**					**67710121**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67710123	**Low discrimination**					**67710123**
Actinomyces naeslundii		78.24	0.50	ESL 12		
Actinomyces viscosus		16.03	0.14	GAL 96		
Bifidobacterium longum subsp. infantis		3.57	-.03			
67710141	**Good identification**					**67710141**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67710161	**Good identification**					**67710161**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67710163	**Good identification**					**67710163**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67711000	**Low discrimination**					**67711000**
					GRA	
Actinomyces naeslundii		64.58	0.62	ESL 12	99	
Fusobacterium mortiferum		25.68	0.44		1	
Bifidobacterium longum subsp. infantis		2.89	0.10		99	
67711001	**Very good identification**					**67711001**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		99.17	0.66	ESL 12			
Actinomyces viscosus		0.83	0.06				
67711002	**Low discrimination**						**67711002**
Actinomyces viscosus		69.45	0.48	ESL 1			
Actinomyces naeslundii		18.61	0.62				
Bifidobacterium longum subsp. infantis		8.42	0.27				
67711003	**Low discrimination**						**67711003**
Actinomyces naeslundii		54.58	0.66	ESL 12			
Actinomyces viscosus		45.26	0.40				
Bifidobacterium longum subsp. infantis		0.17	-.07				
67711010	**Good identification**						**67711010**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67711011	**Good identification**						**67711011**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67711012	**Low discrimination**						**67711012**
					GRA		
Actinomyces naeslundii		52.17	0.41	ESL 12	99		
Actinomyces viscosus		37.36	0.14	MLZ 5	99		
Capnocytophaga ochracea		5.95	-.01		1		
67711013	**Low discrimination**						**67711013**
Actinomyces naeslundii		86.27	0.44	ESL 12			
Actinomyces viscosus		13.73	0.06	MLZ 5			
67711020	**Low discrimination**						**67711020**
Actinomyces naeslundii		48.69	0.62	ESL 12			
Clostridium perfringens		42.23	0.39				
Bifidobacterium longum subsp. infantis		3.87	0.15				
67711021	**Very good identification**						**67711021**
Actinomyces naeslundii		99.42	0.66	ESL 12			
Clostridium perfringens		0.58	0.05				
67711022	**Low discrimination**						**67711022**
					GRA		
Actinomyces naeslundii		27.49	0.62	ESL 12	99		
Prevotella oralis		24.78	0.36		1		
Bifidobacterium longum subsp. infantis		22.10	0.32		99		
67711023	**Very good identification**						**67711023**
					GRA		
Actinomyces naeslundii		97.74	0.66	ESL 12	99		
Actinomyces viscosus		0.82	0.06		99		
Prevotella oralis		0.59	0.02		1		
67711030	**Low discrimination**						**67711030**
					GRA	CAT	
Actinomyces naeslundii		78.09	0.41	ESL 12	99	25	
Clostridium perfringens		13.00	0.05	MLZ 5	99	1	
Capnocytophaga ochracea		8.91	-.01		1	90	
67711031	**Good identification**						**67711031**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.44	ESL 12			
67711033	**Good identification**						**67711033**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67711040	**Acceptable identification**						**67711040**
Actinomyces naeslundii		94.44	0.50	ESL 12			
Actinomyces israelii		2.83	0.14				
Clostridium perfringens		2.74	0.01				
67711041	**Good identification**						**67711041**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67711042	**Low discrimination**						**67711042**
Actinomyces naeslundii		75.85	0.50	ESL 12			
Actinomyces viscosus		16.20	0.14				
Actinomyces israelii		3.41	0.17				
67711043	**Good identification**						**67711043**
Actinomyces naeslundii		95.47	0.53	ESL 12			
Actinomyces viscosus		4.53	0.06				
67711060	**Low discrimination**						**67711060**
					GRA	CAT	
Actinomyces naeslundii		50.76	0.50	ESL 12	99	25	
Clostridium perfringens		47.51	0.27		99	1	
Capnocytophaga ochracea		1.73	-.01		1	90	
67711061	**Good identification**						**67711061**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67711062	**Low discrimination**						**67711062**
					GRA	CAT	
Actinomyces naeslundii		54.22	0.50	ESL 12	99	25	
Clostridium perfringens		23.88	0.21		99	1	
Capnocytophaga ochracea		16.61	0.15		1	90	
67711063	**Good identification**						**67711063**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67711100	**Low discrimination**						**67711100**
					GRA		
Bifidobacterium longum subsp. infantis		48.16	0.44	NIT 1	99		
Actinomyces naeslundii		30.97	0.70	BGL 91	99		
Fusobacterium mortiferum		10.10	0.50		1		
67711101	**Good identification**						**67711101**
					SPO		
Actinomyces naeslundii		90.72	0.73	ESL 12	1		
Clostridium chauvoei		7.53	0.32		99		
Bifidobacterium longum subsp. infantis		0.95	0.10		1		
67711102	**Low discrimination**						**67711102**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		73.62	0.62	NIT 1	99	1	
Actinomyces viscosus		10.91	0.52		99	76	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		8.81	0.50		1	90	
67711103	**Low discrimination**						**67711103**
Actinomyces naeslundii		60.50	0.73	ESL 12			
Actinomyces viscosus		31.34	0.44				
Bifidobacterium longum subsp. infantis		6.41	0.27				
67711110	**Low discrimination**						**67711110**
					GRA	CAT	
Actinomyces naeslundii		52.68	0.48	ESL 12	99	25	
Capnocytophaga ochracea		23.24	0.17	MLZ 5	1	90	
Bifidobacterium longum subsp. infantis		15.72	0.10		99	1	
67711111	**Good identification**						**67711111**
Actinomyces naeslundii		100.0	0.51	ESL 12			
67711113	**Low discrimination**						**67711113**
					GRA		
Actinomyces naeslundii		87.23	0.51	ESL 12	99		
Actinomyces viscosus		8.67	0.10	MLZ 5	99		
Capnocytophaga ochracea		2.33	-.01		1		
67711120	**Low discrimination**						**67711120**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		33.03	0.49	NIT 1	99	1	
Clostridium perfringens		26.69	0.54	BGL 91	99	1	
Capnocytophaga ochracea		22.47	0.50		1	90	
67711121	**Good identification**						**67711121**
					SPO		
Actinomyces naeslundii		92.85	0.73	ESL 12	1		
Clostridium chauvoei		2.57	0.24		99		
Bifidobacterium longum subsp. infantis		1.73	0.15		1		
67711122	**Low discrimination**						**67711122**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		53.70	0.66	NIT 1	99	1	
Capnocytophaga ochracea		32.51	0.66		1	90	
Prevotella oralis		9.53	0.56		1	1	
67711123	**Low discrimination**						**67711123**
					GRA	CAT	
Actinomyces naeslundii		73.92	0.73	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		13.92	0.32		99	1	
Capnocytophaga ochracea		8.43	0.32		1	90	
67711131	**Good identification**						**67711131**
					GRA		
Actinomyces naeslundii		97.40	0.51	ESL 12	99		
Capnocytophaga ochracea		2.60	-.01	MLZ 5	1		
67711132	**Low discrimination**						**67711132**
					GRA	CAT	
Capnocytophaga ochracea		80.64	0.50	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		12.11	0.32	MLZ 10	99	1	
Actinomyces naeslundii		2.26	0.48		99	25	
67711133	**Low discrimination**						**67711133**
					GRA	CAT	
Actinomyces naeslundii		78.32	0.51	ESL 12	99	25	
Capnocytophaga ochracea		18.85	0.15	MLZ 5	1	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium longum subsp. infantis		2.83	-.03		99	1	
67711140	**Low discrimination**						**67711140**
					GRA	CAT	
Actinomyces naeslundii		73.07	0.57	ESL 12	99	25	
Capnocytophaga ochracea		9.62	0.17		1	90	
Bifidobacterium longum subsp. infantis		6.51	0.10		99	1	
67711141	**Very good identification**						**67711141**
Actinomyces naeslundii		100.0	0.60	ESL 12			
67711143	**Very good identification**						**67711143**
					GRA		
Actinomyces naeslundii		95.82	0.60	ESL 12	99		
Actinomyces viscosus		2.84	0.10		99		
Capnocytophaga ochracea		0.76	-.01		1		
67711150	**Low discrimination**						**67711150**
					GRA	CAT	
Actinomyces naeslundii		65.99	0.35	ESL 12	99	25	
Capnocytophaga ochracea		18.33	0.01	MLZ 5	1	90	
Bifidobacterium breve		15.69	0.10		99	1	
67711151	**Good identification**						**67711151**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67711153	**Good identification**						**67711153**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67711160	**Low discrimination**						**67711160**
					GRA	CAT	
Clostridium perfringens		50.09	0.42		99	1	
Capnocytophaga ochracea		24.59	0.33		1	90	
Actinomyces naeslundii		20.77	0.57		99	25	
67711161	**Very good identification**						**67711161**
					GRA	CAT	
Actinomyces naeslundii		97.64	0.60	ESL 12	99	25	
Clostridium perfringens		1.59	0.07		99	1	
Capnocytophaga ochracea		0.78	-.01		1	90	
67711162	**Low discrimination**						**67711162**
					GRA	CAT	
Capnocytophaga ochracea		71.52	0.50	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		10.74	0.32	SOR 10	99	1	
Clostridium perfringens		7.62	0.36		99	1	
67711163	**Good identification**						**67711163**
					GRA	CAT	
Actinomyces naeslundii		91.73	0.60	ESL 12	99	25	
Capnocytophaga ochracea		6.58	0.15		1	90	
Bifidobacterium longum subsp. infantis		0.99	-.03		99	1	
67711171	**Good identification**						**67711171**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67711173	**Low discrimination**						**67711173**
					GRA		
Actinomyces naeslundii		86.85	0.38	ESL 12	99		
Capnocytophaga ochracea		13.16	-.01	MLZ 5	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67711301	**Good identification**						**67711301**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67711303	**Low discrimination**						**67711303**
Actinomyces naeslundii		61.59	0.39	ESL 12			
Actinomyces viscosus		31.90	0.10	RHA 1			
Bifidobacterium longum subsp. infantis		6.52	-.07				
67711321	**Good identification**						**67711321**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67711322	**Low discrimination**						**67711322**
					GRA	CAT	
Prevotella oralis		79.29	0.50	NIT 1	1	1	
Bifidobacterium longum subsp. infantis		10.53	0.32		99	1	
Capnocytophaga ochracea		6.38	0.32		1	90	
67711501	**Good identification**						**67711501**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67711502	**Low discrimination**						**67711502**
					GRA	CAT	
Capnocytophaga ochracea		40.98	0.37	SAL 1	1	90	
Bifidobacterium adolescentis		31.69	0.45	NAG 90	99	1	
Bifidobacterium longum subsp. infantis		19.61	0.27		99	1	
67711503	**Low discrimination**						**67711503**
					GRA		
Actinomyces naeslundii		49.60	0.39	ESL 12	99		
Actinomyces viscosus		25.69	0.10	ARA 1	99		
Capnocytophaga ochracea		10.97	0.02		1		
67711512	**Low discrimination**						**67711512**
					GRA	CAT	
Bifidobacterium adolescentis		86.16	0.46	NIT 1	99	1	
Capnocytophaga ochracea		10.13	0.20	XYL 87	1	90	
Bifidobacterium longum subsp. longum		1.28	0.13		99	1	
67711520	**Low discrimination**						**67711520**
					GRA	CAT	
Capnocytophaga ochracea		77.30	0.37	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		6.50	0.15	BGL 90	99	1	
Clostridium perfringens		5.26	0.20		99	1	
67711521	**Low discrimination**						**67711521**
					GRA		
Actinomyces naeslundii		81.88	0.39	ESL 12	99		
Capnocytophaga ochracea		18.12	0.02	ARA 1	1		
67711522	**Low discrimination**						**67711522**
					GRA	CAT	
Capnocytophaga ochracea		88.65	0.53	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		8.38	0.32		99	1	
Prevotella oralis		1.49	0.22		1	1	
67711532	**Low discrimination**						**67711532**
					GRA	CAT	
Capnocytophaga ochracea		89.98	0.37	SAL 1	1	90	
Bifidobacterium longum subsp. longum		5.06	0.24	MLZ 10	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		2.31	0.22		1	76	
67711562	**Good identification**						**67711562**
					GRA	CAT	
Capnocytophaga ochracea		98.03	0.37	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		0.84	-.03	SOR 10	99	1	
Clostridium perfringens		0.60	0.02		99	1	
67712000	**Good identification**						**67712000**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67712001	**Good identification**						**67712001**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67712002	**Low discrimination**						**67712002**
Actinomyces naeslundii		56.81	0.44	ESL 12			
Actinomyces viscosus		43.19	0.18	GAL 96			
67712003	**Low discrimination**						**67712003**
Actinomyces naeslundii		85.55	0.47	ESL 12			
Actinomyces viscosus		14.45	0.09	GAL 96			
67712020	**Good identification**						**67712020**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67712021	**Good identification**						**67712021**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67712022	**Good identification**						**67712022**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67712023	**Good identification**						**67712023**
Actinomyces naeslundii		100.0	0.47	ESL 12			
67712100	**Acceptable identification**						**67712100**
					GRA		
Actinomyces naeslundii		90.21	0.52	ESL 12	99		
Propionibacterium propionicum		7.34	0.23	GAL 96	99		
Fusobacterium mortiferum		2.45	0.13		1		
67712101	**Good identification**						**67712101**
Actinomyces naeslundii		100.0	0.55	ESL 12			
67712102	**Low discrimination**						**67712102**
Actinomyces naeslundii		61.09	0.52	ESL 12			
Actinomyces viscosus		29.01	0.22	GAL 96			
Propionibacterium propionicum		7.45	0.26				
67712103	**Acceptable identification**						**67712103**
Actinomyces naeslundii		90.45	0.55	ESL 12			
Actinomyces viscosus		9.55	0.14	GAL 96			
67712120	**Good identification**						**67712120**
Actinomyces naeslundii		100.0	0.52	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67712121	**Good identification**						**67712121**
Actinomyces naeslundii		100.0	0.55	ESL 12			
67712122	**Acceptable identification**						**67712122**
					GRA	CAT	
Actinomyces naeslundii		94.64	0.52	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		3.34	-.03	GAL 96	99	1	
Capnocytophaga ochracea		2.02	-.03		1	90	
67712123	**Good identification**						**67712123**
Actinomyces naeslundii		100.0	0.55	ESL 12			
67712140	**Low discrimination**						**67712140**
Actinomyces naeslundii		68.45	0.39	ESL 12			
Propionibacterium propionicum		31.55	0.23	GAL 96			
67712141	**Good identification**						**67712141**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67712142	**Low discrimination**						**67712142**
Actinomyces naeslundii		59.12	0.39	ESL 12			
Propionibacterium propionicum		40.88	0.26	GAL 96			
67712143	**Good identification**						**67712143**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67712160	**Good identification**						**67712160**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67712161	**Good identification**						**67712161**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67712162	**Good identification**						**67712162**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67712163	**Good identification**						**67712163**
Actinomyces naeslundii		100.0	0.42	ESL 12			
67713000	**Very good identification**						**67713000**
Actinomyces naeslundii		96.18	0.67	ESL 12			
Actinomyces israelii		3.82	0.34				
67713001	**Very good identification**						**67713001**
Actinomyces naeslundii		100.0	0.70	ESL 12			
67713002	**Good identification**						**67713002**
Actinomyces naeslundii		92.48	0.67	ESL 12			
Actinomyces israelii		5.51	0.37				
Actinomyces viscosus		1.80	0.14				
67713003	**Very good identification**						**67713003**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		99.57	0.70	ESL 12		
Actinomyces viscosus		0.43	0.06			
67713010	**Low discrimination**					**67713010**
Actinomyces naeslundii		89.87	0.46	ESL 12		
Actinomyces israelii		10.13	0.20	MLZ 5		
67713011	**Good identification**					**67713011**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67713012	**Low discrimination**					**67713012**
Actinomyces naeslundii		85.54	0.46	ESL 12		
Actinomyces israelii		14.46	0.23	MLZ 5		
67713013	**Good identification**					**67713013**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67713020	**Very good identification**					**67713020**
Actinomyces naeslundii		99.30	0.67	ESL 12		
Clostridium perfringens		0.45	0.05			
Actinomyces israelii		0.25	0.14			
67713021	**Very good identification**					**67713021**
Actinomyces naeslundii		100.0	0.70	ESL 12		
67713022	**Very good identification**					**67713022**
					GRA	
Actinomyces naeslundii		98.50	0.67	ESL 12	99	
Prevotella oralis		0.46	0.02		1	
Bifidobacterium longum subsp. infantis		0.41	-.03		99	
67713023	**Very good identification**					**67713023**
Actinomyces naeslundii		100.0	0.70	ESL 12		
67713030	**Good identification**					**67713030**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67713031	**Good identification**					**67713031**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67713032	**Good identification**					**67713032**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67713033	**Good identification**					**67713033**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67713040	**Acceptable identification**					**67713040**
Actinomyces naeslundii		91.96	0.55	ESL 12		
Actinomyces israelii		8.05	0.27			
67713041	**Very good identification**					**67713041**
Actinomyces naeslundii		100.0	0.58	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67713042	**Low discrimination**					**67713042**
Actinomyces naeslundii		88.40	0.55	ESL 12		
Actinomyces israelii		11.60	0.30			
67713043	**Very good identification**					**67713043**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67713051	**Good identification**					**67713051**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713053	**Good identification**					**67713053**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713060	**Good identification**					**67713060**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67713061	**Very good identification**					**67713061**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67713062	**Good identification**					**67713062**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67713063	**Very good identification**					**67713063**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67713071	**Good identification**					**67713071**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713073	**Good identification**					**67713073**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713100	**Good identification**					**67713100**
Actinomyces naeslundii		93.35	0.75	ESL 12		
Actinomyces israelii		5.21	0.45			
Bifidobacterium longum subsp. infantis		0.76	0.10			
67713101	**Very good identification**					**67713101**
Actinomyces naeslundii		100.0	0.78	ESL 12		
67713102	**Low discrimination**					**67713102**
Actinomyces naeslundii		81.86	0.75	ESL 12		
Actinomyces israelii		6.85	0.48			
Bifidobacterium longum subsp. infantis		6.70	0.27			
67713103	**Very good identification**					**67713103**
Actinomyces naeslundii		99.62	0.78	ESL 12		
Actinomyces viscosus		0.27	0.10			
Actinomyces israelii		0.06	0.13			
67713110	**Low discrimination**					**67713110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		84.74	0.53	ESL 12			
Actinomyces israelii		13.43	0.30	MLZ 5			
Bifidobacterium breve		1.83	0.10				
67713111	**Very good identification**						**67713111**
Actinomyces naeslundii		100.0	0.56	ESL 12			
67713112	**Low discrimination**						**67713112**
Actinomyces naeslundii		53.54	0.53	ESL 12			
Bifidobacterium adolescentis		20.09	0.31	MLZ 5			
Actinomyces israelii		12.72	0.33				
67713113	**Very good identification**						**67713113**
Actinomyces naeslundii		100.0	0.56	ESL 12			
67713120	**Very good identification**						**67713120**
Actinomyces naeslundii		95.72	0.75	ESL 12			
Bifidobacterium longum subsp. infantis		1.38	0.15				
Clostridium perfringens		1.11	0.20				
67713121	**Very good identification**						**67713121**
Actinomyces naeslundii		100.0	0.78	ESL 12			
67713122	**Low discrimination**						**67713122**
					GRA	CAT	
Actinomyces naeslundii		77.23	0.75	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		11.24	0.32		99	1	
Capnocytophaga ochracea		6.80	0.32		1	90	
67713123	**Very good identification**						**67713123**
					GRA	CAT	
Actinomyces naeslundii		99.84	0.78	ESL 12	99	25	
Bifidobacterium longum subsp. infantis		0.10	-.03		99	1	
Capnocytophaga ochracea		0.06	-.03		1	90	
67713130	**Acceptable identification**						**67713130**
					GRA	CAT	
Actinomyces naeslundii		94.42	0.53	ESL 12	99	25	
Bifidobacterium breve		3.63	0.15	MLZ 5	99	1	
Capnocytophaga ochracea		1.95	-.01		1	90	
67713131	**Very good identification**						**67713131**
Actinomyces naeslundii		100.0	0.56	ESL 12			
67713132	**Low discrimination**						**67713132**
					GRA	CAT	
Actinomyces naeslundii		62.41	0.53	ESL 12	99	25	
Bifidobacterium breve		24.24	0.32	MLZ 5	99	1	
Capnocytophaga ochracea		11.61	0.15		1	90	
67713133	**Very good identification**						**67713133**
Actinomyces naeslundii		100.0	0.56	ESL 12			
67713140	**Low discrimination**						**67713140**
Actinomyces naeslundii		87.06	0.62	ESL 12			
Actinomyces israelii		10.71	0.38				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium propionicum		1.67	0.23				
67713141	**Very good identification**						**67713141**
Actinomyces naeslundii		100.0	0.65	ESL 12			
67713142	**Low discrimination**						**67713142**
Actinomyces naeslundii		76.89	0.62	ESL 12			
Actinomyces israelii		14.18	0.41				
Bifidobacterium breve		5.01	0.27				
67713143	**Very good identification**						**67713143**
Actinomyces naeslundii		100.0	0.65	ESL 12			
67713150	**Low discrimination**						**67713150**
Actinomyces naeslundii		67.95	0.40	ESL 12			
Actinomyces israelii		23.72	0.23	MLZ 5			
Bifidobacterium breve		8.32	0.10				
67713151	**Good identification**						**67713151**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67713153	**Good identification**						**67713153**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67713160	**Very good identification**						**67713160**
Actinomyces naeslundii		96.34	0.62	ESL 12			
Clostridium perfringens		1.21	0.07				
Bifidobacterium breve		1.10	0.15				
67713161	**Very good identification**						**67713161**
Actinomyces naeslundii		100.0	0.65	ESL 12			
67713162	**Low discrimination**						**67713162**
					GRA	CAT	
Actinomyces naeslundii		83.52	0.62	ESL 12	99	25	
Bifidobacterium breve		9.68	0.32		99	1	
Capnocytophaga ochracea		4.63	0.15		1	90	
67713163	**Very good identification**						**67713163**
Actinomyces naeslundii		100.0	0.65	ESL 12			
67713170	**Low discrimination**						**67713170**
Actinomyces naeslundii		82.12	0.40	ESL 12			
Bifidobacterium breve		17.88	0.15	MLZ 5			
67713171	**Good identification**						**67713171**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67713173	**Good identification**						**67713173**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67713201	**Good identification**						**67713201**
Actinomyces naeslundii		100.0	0.36	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67713203	**Good identification**					**67713203**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713221	**Good identification**					**67713221**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713223	**Good identification**					**67713223**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713300	**Low discrimination**					**67713300**
Actinomyces naeslundii		70.63	0.41	ESL 12		
Actinomyces israelii		29.37	0.25	RHA 1		
67713301	**Good identification**					**67713301**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67713302	**Low discrimination**					**67713302**
Actinomyces naeslundii		58.64	0.41	ESL 12		
Actinomyces israelii		36.57	0.28	RHA 1		
Bifidobacterium longum subsp. infantis		4.80	-.07			
67713303	**Good identification**					**67713303**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67713320	**Low discrimination**					**67713320**
					SPO	
Actinomyces naeslundii		87.96	0.41	ESL 12	1	
Clostridium ramosum		12.04	0.01	RHA 1	92	
67713321	**Good identification**					**67713321**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67713323	**Good identification**					**67713323**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67713401	**Good identification**					**67713401**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713402	**Low discrimination**					**67713402**
Actinomyces israelii		78.67	0.36	CEL 93		
Actinomyces naeslundii		16.31	0.33	XYL 92		
Bifidobacterium adolescentis		5.02	0.09			
67713403	**Good identification**					**67713403**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713421	**Good identification**					**67713421**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67713423	**Good identification**					**67713423**
Actinomyces naeslundii		100.0	0.36	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67713500	**Low discrimination**						**67713500**
Actinomyces israelii		80.31	0.43	CEL 93			
Actinomyces naeslundii		17.77	0.41	XYL 92			
Bifidobacterium adolescentis		1.92	0.09				
67713501	**Good identification**						**67713501**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67713502	**Low discrimination**						**67713502**
Bifidobacterium adolescentis		57.14	0.43	NIT 1			
Actinomyces israelii		36.18	0.46	XYL 87			
Actinomyces naeslundii		5.34	0.41				
67713503	**Acceptable identification**						**67713503**
Actinomyces naeslundii		93.28	0.44	ESL 12			
Bifidobacterium adolescentis		6.72	0.09	ARA 1			
67713512	**Acceptable identification**						**67713512**
Bifidobacterium adolescentis		92.82	0.45	NIT 1			
Actinomyces israelii		7.18	0.32	XYL 87			
67713520	**Low discrimination**						**67713520**
					GRA	CAT	
Actinomyces naeslundii		68.52	0.41	ESL 12	99	25	
Actinomyces israelii		19.77	0.23	ARA 1	99	1	
Capnocytophaga ochracea		11.71	0.02		1	90	
67713521	**Good identification**						**67713521**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67713523	**Good identification**						**67713523**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67713540	**Acceptable identification**						**67713540**
Actinomyces israelii		90.88	0.36	CEL 93			
Actinomyces naeslundii		9.12	0.28	XYL 92			
67713542	**Low discrimination**						**67713542**
Actinomyces israelii		63.74	0.39	CEL 93			
Bifidobacterium adolescentis		32.00	0.28	XYL 92			
Actinomyces naeslundii		4.27	0.28				
67715000	**Low discrimination**						**67715000**
					GRA		
Actinomyces israelii		64.82	0.39	CEL 93	99		
Bifidobacterium longum subsp. infantis		15.16	0.09		99		
Fusobacterium mortiferum		12.40	0.25		1		
67715042	**Good identification**						**67715042**
Actinomyces israelii		100.0	0.35	CEL 93			
67715100	**Low discrimination**						**67715100**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		77.69	0.43	NIT 1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		13.42	0.50	BGL 91	99	1	
Capnocytophaga ochracea		3.19	0.23		1	90	
67715101	**Low discrimination**						**67715101**
Actinomyces naeslundii		50.13	0.39	ESL 12			
Bifidobacterium longum subsp. infantis		42.53	0.09	XYL 1			
Actinomyces israelii		7.35	0.16				
67715102	**Good identification**						**67715102**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		91.76	0.60	NIT 1	99	1	
Capnocytophaga ochracea		3.35	0.39		1	90	
Actinomyces israelii		2.35	0.53		99	1	
67715110	**Low discrimination**						**67715110**
					GRA	CAT	
Actinomyces israelii		60.08	0.36	CEL 93	99	1	
Bifidobacterium longum subsp. infantis		23.51	0.09	MLZ 13	99	1	
Capnocytophaga ochracea		10.62	0.07		1	90	
67715112	**Low discrimination**						**67715112**
					GRA	CAT	
Bifidobacterium adolescentis		57.28	0.46	NIT 1	99	1	
Bifidobacterium longum subsp. infantis		23.74	0.26	ARA 87	99	1	
Capnocytophaga ochracea		9.54	0.23		1	90	
67715120	**Low discrimination**						**67715120**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		75.43	0.47	NIT 1	99	1	
Capnocytophaga ochracea		15.68	0.39	BGL 91	1	90	
Bacteroides fragilis		3.18	0.18		1	99	
67715122	**Low discrimination**						**67715122**
					GRA	CAT	
Bifidobacterium longum subsp. infantis		76.73	0.64	NIT 1	99	1	
Capnocytophaga ochracea		14.20	0.56		1	90	
Bacteroides fragilis		6.09	0.40		1	99	
67715132	**Low discrimination**						**67715132**
					GRA	CAT	
Capnocytophaga ochracea		54.69	0.39	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		26.87	0.30	MLZ 10	99	1	
Parabacteroides distasonis		14.58	0.42		1	76	
67715140	**Low discrimination**						**67715140**
					GRA	CAT	
Actinomyces israelii		80.70	0.43	CEL 93	99	1	
Bifidobacterium longum subsp. infantis		12.14	0.09		99	1	
Capnocytophaga ochracea		5.48	0.07		1	90	
67715162	**Low discrimination**						**67715162**
					GRA	CAT	
Capnocytophaga ochracea		63.53	0.39	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		31.22	0.30	SOR 10	99	1	
Bacteroides fragilis		2.48	0.06		1	99	
67715300	**Very good identification**						**67715300**
Eubacterium contortum		99.26	0.58	NIT 1			
Actinomyces israelii		0.32	0.31				
Bifidobacterium longum subsp. infantis		0.25	0.09				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67715302	**Low discrimination**						**67715302**
Eubacterium contortum		82.98	0.45	NIT 1			
Bifidobacterium longum subsp. infantis		11.84	0.26				
Actinomyces israelii		2.26	0.34				
67715320	**Low discrimination**						**67715320**
					GRA		
Eubacterium contortum		54.60	0.45	NIT 1	99		
Bacteroides vulgatus		42.50	0.36		1		
Bifidobacterium longum subsp. infantis		1.37	0.13		99		
67715322	**Low discrimination**						**67715322**
					GRA		
Parabacteroides distasonis		48.95	0.55	NIT 1	1		
Prevotella oralis		18.42	0.35		1		
Bifidobacterium longum subsp. infantis		16.20	0.30		99		
67715332	**Acceptable identification**						**67715332**
Parabacteroides distasonis		92.83	0.50	NIT 1			
Clostridium clostridioforme		4.37	0.22				
Capnocytophaga ochracea		1.17	0.05				
67715400	**Good identification**						**67715400**
Actinomyces israelii		100.0	0.38	CEL 93			
67715402	**Low discrimination**						**67715402**
Actinomyces israelii		77.28	0.41	CEL 93			
Bifidobacterium adolescentis		19.88	0.25				
Clostridium clostridioforme		2.84	0.07				
67715500	**Low discrimination**						**67715500**
					GRA		
Actinomyces israelii		55.16	0.48	CEL 93	99		
Eubacterium contortum		13.09	0.23		99		
Mitsuokella multacida		7.48	0.21		1		
67715502	**Low discrimination**						**67715502**
Bifidobacterium adolescentis		76.84	0.59	NIT 1			
Actinomyces israelii		12.07	0.51				
Bifidobacterium longum subsp. infantis		5.82	0.26				
67715512	**Very good identification**						**67715512**
Bifidobacterium adolescentis		95.39	0.60	NIT 1			
Actinomyces israelii		1.83	0.37				
Clostridium clostridioforme		1.72	0.27				
67715520	**Low discrimination**						**67715520**
					GRA	CAT	
Bacteroides vulgatus		86.34	0.39	NIT 1	1	3	
Capnocytophaga ochracea		6.84	0.26	RHA 90	1	90	
Bifidobacterium longum subsp. longum		2.71	0.28		99	1	
67715522	**Low discrimination**						**67715522**
					GRA	CAT	
Capnocytophaga ochracea		60.36	0.43	SAL 1	1	90	
Bifidobacterium longum subsp. infantis		18.67	0.30		99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		11.53	0.43		1	76	
67715540	**Low discrimination**						**67715540**
					GRA		
Actinomyces israelii		71.15	0.41	CEL 93	99		
Mitsuokella multacida		24.82	0.21		1		
Bifidobacterium adolescentis		2.18	0.09		99		
67715542	**Low discrimination**						**67715542**
					GRA	CAT	
Bifidobacterium adolescentis		64.50	0.43	NIT 1	99	1	
Actinomyces israelii		31.88	0.44	SOR 11	99	1	
Capnocytophaga ochracea		2.80	0.10		1	90	
67715552	**Acceptable identification**						**67715552**
Bifidobacterium adolescentis		94.30	0.45	NIT 1			
Actinomyces israelii		5.70	0.30	SOR 11			
67715700	**Very good identification**						**67715700**
					GRA		
Eubacterium contortum		96.48	0.57	NIT 1	99		
Bacteroides vulgatus		2.17	0.21		1		
Mitsuokella multacida		1.04	0.25		1		
67715702	**Low discrimination**						**67715702**
Eubacterium contortum		87.36	0.44	NIT 1			
Clostridium clostridioforme		7.13	0.31				
Actinomyces israelii		2.38	0.32				
67715712	**Low discrimination**						**67715712**
Clostridium clostridioforme		88.57	0.42	NIT 1			
Bifidobacterium adolescentis		6.77	0.26	TRE 88			
Eubacterium contortum		2.41	0.10				
67715720	**Acceptable identification**						**67715720**
					GRA		
Bacteroides vulgatus		92.53	0.56	NIT 1	1		
Eubacterium contortum		7.32	0.44		99		
Mitsuokella multacida		0.11	0.15		1		
67715722	**Low discrimination**						**67715722**
					GRA		
Parabacteroides distasonis		57.99	0.51	NIT 1	1		
Eubacterium contortum		17.82	0.31		99		
Bacteroides vulgatus		12.89	0.21		1		
67715732	**Low discrimination**						**67715732**
Clostridium clostridioforme		54.61	0.42	NIT 1			
Parabacteroides distasonis		45.39	0.46	TRE 88			
67717000	**Good identification**						**67717000**
Actinomyces israelii		97.84	0.52	CEL 93			
Actinomyces naeslundii		2.16	0.33				
67717001	**Low discrimination**						**67717001**
Actinomyces naeslundii		76.67	0.36	ESL 12			
Actinomyces israelii		23.33	0.18	XYL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67717002	**Good identification**					**67717002**
Actinomyces israelii		98.11	0.55	CEL 93		
Actinomyces naeslundii		1.45	0.33			
Bifidobacterium adolescentis		0.45	0.09			
67717003	**Low discrimination**					**67717003**
Actinomyces naeslundii		68.65	0.36	ESL 12		
Actinomyces israelii		31.35	0.21	XYL 1		
67717010	**Good identification**					**67717010**
Actinomyces israelii		100.0	0.38	CEL 93		
67717012	**Good identification**					**67717012**
Actinomyces israelii		96.42	0.41	CEL 93		
Bifidobacterium adolescentis		3.58	0.11	MLZ 13		
67717021	**Good identification**					**67717021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67717023	**Good identification**					**67717023**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67717040	**Good identification**					**67717040**
Actinomyces israelii		100.0	0.45	CEL 93		
67717042	**Good identification**					**67717042**
Actinomyces israelii		100.0	0.48	CEL 93		
67717100	**Very good identification**					**67717100**
Actinomyces israelii		97.30	0.63	CEL 93		
Actinomyces naeslundii		1.53	0.41			
Bifidobacterium longum subsp. infantis		1.00	0.09			
67717101	**Low discrimination**					**67717101**
Actinomyces naeslundii		70.04	0.44	ESL 12		
Actinomyces israelii		29.96	0.29	XYL 1		
67717102	**Low discrimination**					**67717102**
Actinomyces israelii		83.68	0.66	CEL 93		
Bifidobacterium adolescentis		9.40	0.43			
Bifidobacterium longum subsp. infantis		5.82	0.26			
67717103	**Low discrimination**					**67717103**
Actinomyces naeslundii		58.35	0.44	ESL 12		
Actinomyces israelii		37.44	0.32	XYL 1		
Bifidobacterium adolescentis		4.21	0.09			
67717110	**Good identification**					**67717110**
Actinomyces israelii		98.63	0.49	CEL 93		
Bifidobacterium adolescentis		1.37	0.11	MLZ 13		
67717112	**Low discrimination**					**67717112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		52.11	0.52	CEL 93			
Bifidobacterium adolescentis		47.89	0.45	MLZ 13			
67717120	**Low discrimination**						**67717120**
Actinomyces israelii		62.75	0.42	CEL 93			
Bifidobacterium longum subsp. infantis		18.03	0.13	NAG 6			
Actinomyces naeslundii		15.47	0.41				
67717121	**Good identification**						**67717121**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67717123	**Acceptable identification**						**67717123**
Actinomyces naeslundii		92.65	0.44	ESL 12			
Bifidobacterium longum subsp. infantis		7.35	-.04	XYL 1			
67717140	**Good identification**						**67717140**
Actinomyces israelii		99.29	0.56	CEL 93			
Actinomyces naeslundii		0.71	0.28				
67717142	**Very good identification**						**67717142**
Actinomyces israelii		96.11	0.59	CEL 93			
Bifidobacterium adolescentis		3.43	0.28				
Actinomyces naeslundii		0.46	0.28				
67717150	**Good identification**						**67717150**
Actinomyces israelii		100.0	0.42	CEL 93			
67717152	**Low discrimination**						**67717152**
Actinomyces israelii		77.39	0.45	CEL 93			
Bifidobacterium adolescentis		22.61	0.29	MLZ 13			
67717160	**Low discrimination**						**67717160**
Actinomyces israelii		89.94	0.35	CEL 93			
Actinomyces naeslundii		10.06	0.28	NAG 6			
67717162	**Low discrimination**						**67717162**
					GRA	CAT	
Actinomyces israelii		81.48	0.38	CEL 93	99	1	
Capnocytophaga ochracea		8.34	0.05	NAG 6	1	90	
Actinomyces naeslundii		6.08	0.28		99	25	
67717202	**Good identification**						**67717202**
Actinomyces israelii		100.0	0.36	CEL 93			
67717300	**Low discrimination**						**67717300**
Actinomyces israelii		64.26	0.43	CEL 93			
Eubacterium contortum		35.74	0.24	RHA 7			
67717302	**Acceptable identification**						**67717302**
Actinomyces israelii		92.55	0.46	CEL 93			
Eubacterium contortum		6.06	0.11	RHA 7			
Bifidobacterium adolescentis		1.40	0.09				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67717340	**Good identification**					**67717340**
Actinomyces israelii		100.0	0.36	CEL 93		
67717342	**Good identification**					**67717342**
Actinomyces israelii		100.0	0.39	CEL 93		
67717400	**Good identification**					**67717400**
Actinomyces israelii		100.0	0.51	CEL 93		
67717402	**Good identification**					**67717402**
Actinomyces israelii		96.42	0.54	CEL 93		
Bifidobacterium adolescentis		3.58	0.23			
67717410	**Good identification**					**67717410**
Actinomyces israelii		100.0	0.37	CEL 93		
67717412	**Low discrimination**					**67717412**
Actinomyces israelii		76.70	0.40	CEL 93		
Bifidobacterium adolescentis		23.30	0.25	MLZ 13		
67717440	**Good identification**					**67717440**
Actinomyces israelii		100.0	0.44	CEL 93		
67717442	**Good identification**					**67717442**
Actinomyces israelii		98.83	0.47	CEL 93		
Bifidobacterium adolescentis		1.17	0.08			
67717500	**Good identification**					**67717500**
					GRA	
Actinomyces israelii		93.28	0.61	CEL 93	99	
Mitsuokella multacida		5.21	0.27		1	
Bifidobacterium adolescentis		1.30	0.23		99	
67717502	**Low discrimination**					**67717502**
					GRA	
Actinomyces israelii		52.00	0.64	CEL 93	99	
Bifidobacterium adolescentis		47.79	0.57		99	
Mitsuokella multacida		0.22	0.11		1	
67717510	**Low discrimination**					**67717510**
					GRA	
Actinomyces israelii		80.74	0.47	CEL 93	99	
Mitsuokella multacida		10.06	0.19	MLZ 13	1	
Bifidobacterium adolescentis		9.20	0.25		99	
67717512	**Low discrimination**					**67717512**
Bifidobacterium adolescentis		88.03	0.59	NIT 1		
Actinomyces israelii		11.71	0.50			
Bifidobacterium dentium		0.15	-.07			
67717520	**Low discrimination**					**67717520**
					GRA	
Actinomyces israelii		72.39	0.41	CEL 93	99	
Mitsuokella multacida		15.84	0.17	NAG 6	1	
Bacteroides vulgatus		11.77	0.05		1	

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
67717522	**Low discrimination**						**67717522**
					GRA	CAT	
Actinomyces israelii		80.21	0.44	CEL 93	99	1	
Bifidobacterium adolescentis		11.67	0.23	NAG 6	99	1	
Capnocytophaga ochracea		6.20	0.09		1	90	
67717540	**Low discrimination**						**67717540**
					GRA		
Actinomyces israelii		87.10	0.54	CEL 93	99		
Mitsuokella multacida		12.51	0.27		1		
Bifidobacterium adolescentis		0.39	0.08		99		
67717542	**Low discrimination**						**67717542**
					GRA		
Actinomyces israelii		76.76	0.57	CEL 93	99		
Bifidobacterium adolescentis		22.42	0.42		99		
Mitsuokella multacida		0.82	0.11		1		
67717550	**Low discrimination**						**67717550**
					COC	GRA	
Actinomyces israelii		71.19	0.40	CEL 93	1	99	
Mitsuokella multacida		22.81	0.19	MLZ 13	1	1	
Blautia producta		3.42	-.03		99	99	
67717552	**Low discrimination**						**67717552**
					GRA		
Bifidobacterium adolescentis		70.01	0.43	NIT 1	99		
Actinomyces israelii		29.30	0.43	SOR 11	99		
Mitsuokella multacida		0.70	0.03		1		
67717562	**Good identification**						**67717562**
Actinomyces israelii		95.58	0.37	CEL 93			
Bifidobacterium adolescentis		4.42	0.08	NAG 6			
67717702	**Low discrimination**						**67717702**
					GRA		
Actinomyces israelii		77.47	0.45	CEL 93	99		
Bifidobacterium adolescentis		9.56	0.23	RHA 7	99		
Mitsuokella multacida		7.91	0.15		1		
67717742	**Low discrimination**						**67717742**
					GRA		
Actinomyces israelii		76.82	0.38	CEL 93	99		
Mitsuokella multacida		20.17	0.15	RHA 7	1		
Bifidobacterium adolescentis		3.01	0.08		99		
67721000	**Good identification**						**67721000**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67721001	**Good identification**						**67721001**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67721002	**Low discrimination**						**67721002**
Actinomyces naeslundii		87.36	0.46	ESL 12			
Actinomyces viscosus		12.64	0.07	FRU 95			
67721003	**Good identification**						**67721003**
Actinomyces naeslundii		96.88	0.49	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces viscosus		3.12	-.01	FRU 95		
67721020	**Low discrimination**					**67721020**
Actinomyces naeslundii		86.26	0.46	ESL 12		
Clostridium perfringens		13.74	0.10	FRU 95		
67721021	**Good identification**					**67721021**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67721022	**Acceptable identification**					**67721022**
Actinomyces naeslundii		93.03	0.46	ESL 12		
Clostridium perfringens		6.97	0.04	FRU 95		
67721023	**Good identification**					**67721023**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67721041	**Good identification**					**67721041**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67721043	**Good identification**					**67721043**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67721061	**Good identification**					**67721061**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67721063	**Good identification**					**67721063**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67721100	**Good identification**					**67721100**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67721101	**Very good identification**					**67721101**
					SPO	
Actinomyces naeslundii		96.28	0.56	ESL 12	1	
Clostridium chauvoei		3.72	0.09	FRU 95	99	
67721102	**Acceptable identification**					**67721102**
Actinomyces naeslundii		90.41	0.53	ESL 12		
Actinomyces viscosus		8.17	0.11	FRU 95		
Bifidobacterium longum subsp. infantis		1.42	-.07			
67721103	**Very good identification**					**67721103**
Actinomyces naeslundii		98.03	0.56	ESL 12		
Actinomyces viscosus		1.97	0.03	FRU 95		
67721120	**Low discrimination**					**67721120**
Actinomyces naeslundii		70.91	0.53	ESL 12		
Clostridium perfringens		29.09	0.24	FRU 95		
67721121	**Very good identification**					**67721121**
					SPO	
Actinomyces naeslundii		98.73	0.56	ESL 12	1	
Clostridium chauvoei		1.27	0.01	FRU 95	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67721122	**Low discrimination**					**67721122**
					GRA	
Actinomyces naeslundii		78.79	0.53	ESL 12	99	
Clostridium perfringens		15.21	0.19	FRU 95	99	
Prevotella oralis		2.47	0.02		1	
67721123	**Very good identification**					**67721123**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67721140	**Good identification**					**67721140**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67721141	**Good identification**					**67721141**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67721142	**Good identification**					**67721142**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67721143	**Good identification**					**67721143**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67721160	**Low discrimination**					**67721160**
Actinomyces naeslundii		69.31	0.40	ESL 12		
Clostridium perfringens		30.69	0.12	FRU 95		
67721161	**Good identification**					**67721161**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67721162	**Low discrimination**					**67721162**
Actinomyces naeslundii		82.75	0.40	ESL 12		
Clostridium perfringens		17.25	0.06	FRU 95		
67721163	**Good identification**					**67721163**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67722101	**Good identification**					**67722101**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67722103	**Good identification**					**67722103**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67722121	**Good identification**					**67722121**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67722123	**Good identification**					**67722123**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67723000	**Good identification**					**67723000**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67723001	**Good identification**					**67723001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	ESL 12		
67723002	**Good identification**					**67723002**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67723003	**Good identification**					**67723003**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67723020	**Good identification**					**67723020**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67723021	**Good identification**					**67723021**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67723022	**Good identification**					**67723022**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67723023	**Good identification**					**67723023**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67723040	**Good identification**					**67723040**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67723041	**Good identification**					**67723041**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67723042	**Good identification**					**67723042**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67723043	**Good identification**					**67723043**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67723060	**Good identification**					**67723060**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67723061	**Good identification**					**67723061**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67723062	**Good identification**					**67723062**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67723063	**Good identification**					**67723063**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67723100	**Very good identification**					**67723100**
Actinomyces naeslundii		97.97	0.58	ESL 12		
Actinomyces israelii		2.03	0.20	FRU 95		
67723101	**Very good identification**					**67723101**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67723102	**Very good identification**					**67723102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.98	0.58	ESL 12		
Actinomyces israelii		3.02	0.23	FRU 95		
67723103	**Very good identification**					**67723103**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67723110	**Good identification**					**67723110**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67723111	**Good identification**					**67723111**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67723112	**Good identification**					**67723112**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67723113	**Good identification**					**67723113**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67723120	**Very good identification**					**67723120**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67723121	**Very good identification**					**67723121**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67723122	**Very good identification**					**67723122**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67723123	**Very good identification**					**67723123**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67723130	**Good identification**					**67723130**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67723131	**Good identification**					**67723131**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67723132	**Good identification**					**67723132**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67723133	**Good identification**					**67723133**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67723140	**Good identification**					**67723140**
Actinomyces naeslundii		95.63	0.45	ESL 12		
Actinomyces israelii		4.37	0.13	FRU 95		
67723141	**Good identification**					**67723141**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67723142	**Acceptable identification**					**67723142**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		93.58	0.45	ESL 12		
Actinomyces israelii		6.42	0.16	FRU 95		
67723143	**Good identification**					**67723143**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67723160	**Good identification**					**67723160**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67723161	**Good identification**					**67723161**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67723162	**Good identification**					**67723162**
Actinomyces naeslundii		100.0	0.45	ESL 12		
67723163	**Good identification**					**67723163**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67727100	**Good identification**					**67727100**
Actinomyces israelii		95.94	0.38	FRU 99		
Actinomyces naeslundii		4.06	0.24	CEL 93		
67727102	**Good identification**					**67727102**
Actinomyces israelii		97.26	0.41	FRU 99		
Actinomyces naeslundii		2.75	0.24	CEL 93		
67727502	**Low discrimination**					**67727502**
					GRA	
Actinomyces israelii		78.63	0.40	FRU 99	99	
Bifidobacterium adolescentis		12.81	0.20	CEL 93	99	
Mitsuokella multacida		8.56	0.11		1	
67730000	**Low discrimination**					**67730000**
					GRA	
Actinomyces naeslundii		49.11	0.44	ESL 12	99	
Fusobacterium mortiferum		48.08	0.32	GAL 96	1	
Actinomyces viscosus		2.81	-.02		99	
67730001	**Good identification**					**67730001**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67730020	**Low discrimination**					**67730020**
Actinomyces naeslundii		83.25	0.44	ESL 12		
Clostridium perfringens		16.75	0.10	GAL 96		
67730021	**Good identification**					**67730021**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67730022	**Low discrimination**					**67730022**
Actinomyces naeslundii		86.82	0.44	ESL 12		
Clostridium perfringens		8.22	0.04	GAL 96		
Actinomyces viscosus		4.97	-.02			
67730023	**Good identification**					**67730023**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.47	ESL 12		
67730100	**Low discrimination**					**67730100**
					GRA	
Actinomyces naeslundii		54.40	0.52	ESL 12	99	
Fusobacterium mortiferum		43.66	0.38	GAL 96	1	
Actinomyces viscosus		1.94	0.03		99	
67730101	**Good identification**					**67730101**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67730102	**Low discrimination**					**67730102**
					GRA	
Actinomyces viscosus		76.14	0.37	ESL 1	99	
Actinomyces naeslundii		21.52	0.52	TRE 7	99	
Fusobacterium mortiferum		1.92	0.22		1	
67730103	**Low discrimination**					**67730103**
Actinomyces naeslundii		55.98	0.55	ESL 12		
Actinomyces viscosus		44.02	0.29	GAL 96		
67730120	**Low discrimination**					**67730120**
Actinomyces naeslundii		65.87	0.52	ESL 12		
Clostridium perfringens		34.14	0.24	GAL 96		
67730121	**Good identification**					**67730121**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67730122	**Low discrimination**					**67730122**
					GRA	
Actinomyces naeslundii		72.69	0.52	ESL 12	99	
Clostridium perfringens		17.73	0.19	GAL 96	99	
Prevotella oralis		2.87	0.02		1	
67730123	**Good identification**					**67730123**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67730140	**Good identification**					**67730140**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67730141	**Good identification**					**67730141**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67730142	**Low discrimination**					**67730142**
Actinomyces naeslundii		83.16	0.39	ESL 12		
Actinomyces viscosus		16.84	0.03	GAL 96		
67730143	**Good identification**					**67730143**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67730160	**Low discrimination**					**67730160**
Actinomyces naeslundii		64.13	0.39	ESL 12		
Clostridium perfringens		35.87	0.12	GAL 96		
67730161	**Good identification**					**67730161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.42	ESL 12		
67730162	**Low discrimination**					**67730162**
Actinomyces naeslundii		79.16	0.39	ESL 12		
Clostridium perfringens		20.84	0.06	GAL 96		
67730163	**Good identification**					**67730163**
Actinomyces naeslundii		100.0	0.42	ESL 12		
67731000	**Low discrimination**					**67731000**
					GRA	
Actinomyces naeslundii		89.20	0.67	ESL 12	99	
Fusobacterium mortiferum		5.46	0.35		1	
Actinomyces israelii		2.35	0.31		99	
67731001	**Very good identification**					**67731001**
Actinomyces naeslundii		100.0	0.70	ESL 12		
67731002	**Low discrimination**					**67731002**
Actinomyces naeslundii		82.60	0.67	ESL 12		
Actinomyces viscosus		11.95	0.29			
Actinomyces israelii		3.27	0.34			
67731003	**Very good identification**					**67731003**
Actinomyces naeslundii		96.89	0.70	ESL 12		
Actinomyces viscosus		3.12	0.21			
67731010	**Low discrimination**					**67731010**
Actinomyces naeslundii		88.93	0.46	ESL 12		
Actinomyces israelii		6.66	0.17	MLZ 5		
Lactobacillus jensenii		4.42	0.02			
67731011	**Good identification**					**67731011**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67731012	**Low discrimination**					**67731012**
					GRA	
Actinomyces naeslundii		59.24	0.46	ESL 12	99	
Leptotrichia buccalis		34.11	0.07	MLZ 5	1	
Actinomyces israelii		6.65	0.20		99	
67731013	**Good identification**					**67731013**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67731020	**Low discrimination**					**67731020**
Actinomyces naeslundii		54.65	0.67	ESL 12		
Clostridium perfringens		45.35	0.44			
67731021	**Very good identification**					**67731021**
Actinomyces naeslundii		99.44	0.70	ESL 12		
Clostridium perfringens		0.56	0.10			
67731022	**Low discrimination**					**67731022**
					GRA	
Actinomyces naeslundii		69.96	0.67	ESL 12	99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		27.32	0.38		99	
Prevotella oralis		2.07	0.16		1	
67731023	**Very good identification**					**67731023**
Actinomyces naeslundii		99.74	0.70	ESL 12		
Clostridium perfringens		0.26	0.04			
67731030	**Low discrimination**					**67731030**
Actinomyces naeslundii		86.26	0.46	ESL 12		
Clostridium perfringens		13.74	0.10	MLZ 5		
67731031	**Good identification**					**67731031**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67731032	**Low discrimination**					**67731032**
					GRA	
Actinomyces naeslundii		87.80	0.46	ESL 12	99	
Clostridium perfringens		6.58	0.04	MLZ 5	99	
Leptotrichia buccalis		5.62	-.09		1	
67731033	**Good identification**					**67731033**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67731040	**Acceptable identification**					**67731040**
Actinomyces naeslundii		92.10	0.55	ESL 12		
Actinomyces israelii		5.35	0.24			
Clostridium perfringens		2.55	0.06			
67731041	**Very good identification**					**67731041**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67731042	**Acceptable identification**					**67731042**
Actinomyces naeslundii		91.99	0.55	ESL 12		
Actinomyces israelii		8.01	0.27			
67731043	**Very good identification**					**67731043**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67731051	**Good identification**					**67731051**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731053	**Good identification**					**67731053**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731060	**Low discrimination**					**67731060**
Actinomyces naeslundii		52.76	0.55	ESL 12		
Clostridium perfringens		47.25	0.31			
67731061	**Very good identification**					**67731061**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67731062	**Low discrimination**					**67731062**
Actinomyces naeslundii		70.35	0.55	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium perfringens		29.65	0.26			
67731063	**Very good identification**					**67731063**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67731071	**Good identification**					**67731071**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731073	**Good identification**					**67731073**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731100	**Low discrimination**					**67731100**
					GRA	
Actinomyces naeslundii		84.88	0.75	ESL 12	99	
Clostridium perfringens		5.61	0.33		99	
Fusobacterium mortiferum		4.26	0.41		1	
67731101	**Very good identification**					**67731101**
					SPO	
Actinomyces naeslundii		99.53	0.78	ESL 12	1	
Clostridium chauvoei		0.47	0.15		99	
67731102	**Low discrimination**					**67731102**
Actinomyces naeslundii		75.57	0.75	ESL 12		
Actinomyces viscosus		6.83	0.33			
Bifidobacterium longum subsp. infantis		6.18	0.27			
67731103	**Very good identification**					**67731103**
					SPO	
Actinomyces naeslundii		97.90	0.78	ESL 12	1	
Actinomyces viscosus		1.97	0.25		1	
Clostridium chauvoei		0.08	0.03		99	
67731110	**Low discrimination**					**67731110**
					GRA	
Actinomyces naeslundii		76.48	0.53	ESL 12	99	
Leptotrichia buccalis		15.48	0.07	MLZ 5	1	
Actinomyces israelii		8.04	0.27		99	
67731111	**Very good identification**					**67731111**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67731112	**Acceptable identification**					**67731112**
					GRA	
Leptotrichia buccalis		92.26	0.41	NIT 10	1	
Actinomyces naeslundii		4.61	0.53	CEL 99	99	
Bifidobacterium adolescentis		1.41	0.29		99	
67731113	**Low discrimination**					**67731113**
					GRA	
Actinomyces naeslundii		88.11	0.56	ESL 12	99	
Leptotrichia buccalis		11.89	0.07	MLZ 5	1	
67731120	**Low discrimination**					**67731120**
Clostridium perfringens		67.12	0.59			
Actinomyces naeslundii		31.40	0.75			
Bifidobacterium longum subsp. infantis		0.45	0.15			
67731121	**Very good identification**					**67731121**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		98.43	0.78	ESL 12	1	
Clostridium perfringens		1.42	0.24		84	
Clostridium chauvoei		0.16	0.07		99	
67731122	**Low discrimination**					**67731122**
					GRA	
Clostridium perfringens		40.20	0.53		99	
Actinomyces naeslundii		39.96	0.75		99	
Prevotella oralis		6.52	0.36		1	
67731123	**Very good identification**					**67731123**
					GRA	
Actinomyces naeslundii		99.06	0.78	ESL 12	99	
Clostridium perfringens		0.67	0.19		99	
Prevotella oralis		0.11	0.02		1	
67731130	**Low discrimination**					**67731130**
Actinomyces naeslundii		67.61	0.53	ESL 12		
Clostridium perfringens		27.74	0.24	MLZ 5		
Bifidobacterium breve		1.73	0.12			
67731131	**Very good identification**					**67731131**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67731133	**Very good identification**					**67731133**
					GRA	
Actinomyces naeslundii		98.52	0.56	ESL 12	99	
Leptotrichia buccalis		1.48	-.09	MLZ 5	1	
67731140	**Low discrimination**					**67731140**
Actinomyces naeslundii		86.73	0.62	ESL 12		
Actinomyces israelii		7.08	0.34			
Clostridium perfringens		6.19	0.20			
67731141	**Very good identification**					**67731141**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67731142	**Low discrimination**					**67731142**
Actinomyces naeslundii		81.65	0.62	ESL 12		
Actinomyces israelii		10.00	0.37			
Bifidobacterium breve		3.55	0.24			
67731143	**Very good identification**					**67731143**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67731150	**Low discrimination**					**67731150**
Actinomyces naeslundii		81.18	0.40	ESL 12		
Actinomyces israelii		18.82	0.20	MLZ 5		
67731151	**Good identification**					**67731151**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67731153	**Good identification**					**67731153**
Actinomyces naeslundii		100.0	0.43	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67731160	**Low discrimination**					**67731160**
Clostridium perfringens		69.37	0.46			
Actinomyces naeslundii		30.06	0.62			
Bifidobacterium breve		0.23	0.12			
67731161	**Very good identification**					**67731161**
Actinomyces naeslundii		98.47	0.65	ESL 12		
Clostridium perfringens		1.53	0.12			
67731162	**Low discrimination**					**67731162**
Clostridium perfringens		48.39	0.41			
Actinomyces naeslundii		44.56	0.62			
Bifidobacterium breve		3.44	0.29			
67731163	**Very good identification**					**67731163**
Actinomyces naeslundii		99.27	0.65	ESL 12		
Clostridium perfringens		0.73	0.06			
67731170	**Low discrimination**					**67731170**
Actinomyces naeslundii		62.97	0.40	ESL 12		
Clostridium perfringens		27.89	0.12	MLZ 5		
Bifidobacterium breve		9.14	0.12			
67731171	**Good identification**					**67731171**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67731173	**Good identification**					**67731173**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67731201	**Good identification**					**67731201**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731203	**Good identification**					**67731203**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731221	**Good identification**					**67731221**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731223	**Good identification**					**67731223**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67731300	**Low discrimination**					**67731300**
Actinomyces naeslundii		78.37	0.41	ESL 12		
Actinomyces israelii		21.63	0.22	RHA 1		
67731301	**Good identification**					**67731301**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67731302	**Low discrimination**					**67731302**
					GRA	
Actinomyces naeslundii		43.93	0.41	ESL 12	99	
Actinomyces israelii		18.19	0.25	RHA 1	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Prevotella oralis	16.00	0.08		1	
67731303	**Good identification**					**67731303**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
67731321	**Good identification**					**67731321**
	Actinomyces naeslundii	100.0	0.44	ESL 12		
67731322	**Low discrimination**					**67731322**
					SPO	
	Parabacteroides distasonis	62.50	0.50	NIT 1	1	
	Prevotella oralis	23.99	0.30	XYL 90	1	
	Clostridium ramosum	5.25	0.19		92	
67731323	**Low discrimination**					**67731323**
					GRA	
	Actinomyces naeslundii	89.18	0.44	ESL 12	99	
	Parabacteroides distasonis	10.83	0.15	RHA 1	1	
67731332	**Acceptable identification**					**67731332**
	Parabacteroides distasonis	92.63	0.45	NIT 1		
	Clostridium clostridioforme	6.25	0.19	XYL 90		
	Leptotrichia buccalis	1.12	-.09			
67731401	**Good identification**					**67731401**
	Actinomyces naeslundii	100.0	0.36	ESL 12		
67731403	**Good identification**					**67731403**
	Actinomyces naeslundii	100.0	0.36	ESL 12		
67731421	**Good identification**					**67731421**
	Actinomyces naeslundii	100.0	0.36	ESL 12		
67731423	**Good identification**					**67731423**
	Actinomyces naeslundii	100.0	0.36	ESL 12		
67731500	**Low discrimination**					**67731500**
					GRA	
	Actinomyces israelii	51.29	0.40	CEL 93	99	
	Mitsuokella multacida	21.27	0.21	XYL 92	1	
	Actinomyces naeslundii	17.09	0.41		99	
67731501	**Acceptable identification**					**67731501**
					GRA	
	Actinomyces naeslundii	91.56	0.44	ESL 12	99	
	Mitsuokella multacida	8.44	0.04	ARA 1	1	
67731502	**Low discrimination**					**67731502**
	Bifidobacterium adolescentis	57.34	0.42	NIT 1		
	Actinomyces israelii	29.58	0.43	XYL 87		
	Actinomyces naeslundii	6.57	0.41			
67731503	**Acceptable identification**					**67731503**
	Actinomyces naeslundii	94.45	0.44	ESL 12		
	Bifidobacterium adolescentis	5.55	0.07	ARA 1		
67731512	**Low discrimination**					**67731512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Bifidobacterium adolescentis		77.33	0.43	NIT 1	99		
Clostridium clostridioforme		10.14	0.25	XYL 87	50		
Leptotrichia buccalis		7.65	0.07		1		
67731521	**Good identification**						**67731521**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67731522	**Low discrimination**						**67731522**
					GRA	CAT	
Parabacteroides distasonis		44.06	0.38	NIT 1	1	76	
Capnocytophaga ochracea		18.41	0.19	XYL 90	1	90	
Clostridium perfringens		12.04	0.19		99	1	
67731523	**Good identification**						**67731523**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67731542	**Low discrimination**						**67731542**
Actinomyces israelii		55.68	0.36	CEL 93			
Bifidobacterium adolescentis		34.31	0.26	XYL 92			
Actinomyces naeslundii		5.61	0.28				
67731712	**Good identification**						**67731712**
					GRA		
Clostridium clostridioforme		98.15	0.40	NIT 1	50		
Bifidobacterium adolescentis		1.03	0.09	XYL 91	99		
Mitsuokella multacida		0.82	0.01		1		
67731722	**Low discrimination**						**67731722**
Parabacteroides distasonis		83.07	0.46	NIT 1			
Clostridium clostridioforme		16.93	0.28	XYL 90			
67731732	**Low discrimination**						**67731732**
Clostridium clostridioforme		63.29	0.40	NIT 1			
Parabacteroides distasonis		36.71	0.41	XYL 91			
67732000	**Good identification**						**67732000**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67732001	**Good identification**						**67732001**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67732002	**Good identification**						**67732002**
Actinomyces naeslundii		97.14	0.49	ESL 12			
Actinomyces viscosus		2.86	-.02	GAL 96			
67732003	**Good identification**						**67732003**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67732020	**Good identification**						**67732020**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67732021	**Good identification**						**67732021**
Actinomyces naeslundii		100.0	0.52	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67732022	**Good identification**					**67732022**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67732023	**Good identification**					**67732023**
Actinomyces naeslundii		100.0	0.52	ESL 12		
67732040	**Good identification**					**67732040**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67732041	**Good identification**					**67732041**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67732042	**Good identification**					**67732042**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67732043	**Good identification**					**67732043**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67732060	**Good identification**					**67732060**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67732061	**Good identification**					**67732061**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67732062	**Good identification**					**67732062**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67732063	**Good identification**					**67732063**
Actinomyces naeslundii		100.0	0.39	ESL 12		
67732100	**Good identification**					**67732100**
Actinomyces naeslundii		94.30	0.57	ESL 12		
Propionibacterium propionicum		3.23	0.21	GAL 96		
Actinomyces israelii		2.47	0.20			
67732101	**Very good identification**					**67732101**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67732102	**Good identification**					**67732102**
Actinomyces naeslundii		90.16	0.57	ESL 12		
Propionibacterium propionicum		4.64	0.24	GAL 96		
Actinomyces israelii		3.55	0.23			
67732103	**Very good identification**					**67732103**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67732111	**Good identification**					**67732111**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67732113	**Good identification**					**67732113**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	ESL 12		
67732120	**Very good identification**					**67732120**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67732121	**Very good identification**					**67732121**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67732122	**Very good identification**					**67732122**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67732123	**Very good identification**					**67732123**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67732131	**Good identification**					**67732131**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67732133	**Good identification**					**67732133**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67732140	**Low discrimination**					**67732140**
Actinomyces naeslundii		79.87	0.44	ESL 12		
Propionibacterium propionicum		15.52	0.21	GAL 96		
Actinomyces israelii		4.61	0.13			
67732141	**Good identification**					**67732141**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67732142	**Low discrimination**					**67732142**
Actinomyces naeslundii		72.56	0.44	ESL 12		
Propionibacterium propionicum		21.15	0.24	GAL 96		
Actinomyces israelii		6.29	0.16			
67732143	**Good identification**					**67732143**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67732160	**Good identification**					**67732160**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67732161	**Good identification**					**67732161**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67732162	**Good identification**					**67732162**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67732163	**Good identification**					**67732163**
Actinomyces naeslundii		100.0	0.47	ESL 12		
67733000	**Good identification**					**67733000**
Actinomyces naeslundii		92.86	0.72	ESL 12		
Actinomyces israelii		7.15	0.44			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67733001	**Very good identification**					**67733001**
Actinomyces naeslundii		100.0	0.75	ESL 12		
67733002	**Low discrimination**					**67733002**
Actinomyces naeslundii		89.65	0.72	ESL 12		
Actinomyces israelii		10.35	0.47			
67733003	**Very good identification**					**67733003**
Actinomyces naeslundii		100.0	0.75	ESL 12		
67733010	**Low discrimination**					**67733010**
Actinomyces naeslundii		82.07	0.50	ESL 12		
Actinomyces israelii		17.93	0.30	MLZ 5		
67733011	**Good identification**					**67733011**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67733012	**Low discrimination**					**67733012**
Actinomyces naeslundii		75.32	0.50	ESL 12		
Actinomyces israelii		24.68	0.33	MLZ 5		
67733013	**Good identification**					**67733013**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67733020	**Very good identification**					**67733020**
Actinomyces naeslundii		99.09	0.72	ESL 12		
Actinomyces israelii		0.49	0.24			
Clostridium perfringens		0.43	0.10			
67733021	**Very good identification**					**67733021**
Actinomyces naeslundii		100.0	0.75	ESL 12		
67733022	**Very good identification**					**67733022**
Actinomyces naeslundii		99.07	0.72	ESL 12		
Actinomyces israelii		0.73	0.27			
Clostridium perfringens		0.20	0.04			
67733023	**Very good identification**					**67733023**
Actinomyces naeslundii		100.0	0.75	ESL 12		
67733030	**Good identification**					**67733030**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67733031	**Good identification**					**67733031**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67733032	**Good identification**					**67733032**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67733033	**Good identification**					**67733033**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	ESL 12		
67733040	**Low discrimination**					**67733040**
Actinomyces naeslundii		85.50	0.59	ESL 12		
Actinomyces israelii		14.50	0.37			
67733041	**Very good identification**					**67733041**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67733042	**Low discrimination**					**67733042**
Actinomyces naeslundii		79.72	0.59	ESL 12		
Actinomyces israelii		20.28	0.40			
67733043	**Very good identification**					**67733043**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67733050	**Low discrimination**					**67733050**
Actinomyces naeslundii		67.51	0.38	ESL 12		
Actinomyces israelii		32.50	0.23	MLZ 5		
67733051	**Good identification**					**67733051**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733052	**Low discrimination**					**67733052**
Actinomyces naeslundii		58.07	0.38	ESL 12		
Actinomyces israelii		41.93	0.26	MLZ 5		
67733053	**Good identification**					**67733053**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733060	**Very good identification**					**67733060**
Actinomyces naeslundii		98.93	0.59	ESL 12		
Actinomyces israelii		1.07	0.17			
67733061	**Very good identification**					**67733061**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67733062	**Very good identification**					**67733062**
Actinomyces naeslundii		98.40	0.59	ESL 12		
Actinomyces israelii		1.60	0.20			
67733063	**Very good identification**					**67733063**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67733070	**Good identification**					**67733070**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67733071	**Good identification**					**67733071**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733072	**Good identification**					**67733072**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	ESL 12		
67733073	**Good identification**					**67733073**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733100	**Good identification**					**67733100**
Actinomyces naeslundii		90.13	0.80	ESL 12		
Actinomyces israelii		9.75	0.54			
Propionibacterium propionicum		0.13	0.21			
67733101	**Very good identification**					**67733101**
Actinomyces naeslundii		99.93	0.83	ESL 12		
Actinomyces israelii		0.07	0.20			
67733102	**Low discrimination**					**67733102**
Actinomyces naeslundii		85.13	0.80	ESL 12		
Actinomyces israelii		13.81	0.57			
Bifidobacterium adolescentis		0.47	0.26			
67733103	**Very good identification**					**67733103**
Actinomyces naeslundii		99.89	0.83	ESL 12		
Actinomyces israelii		0.11	0.23			
67733110	**Low discrimination**					**67733110**
Actinomyces naeslundii		76.51	0.58	ESL 12		
Actinomyces israelii		23.49	0.40	MLZ 5		
67733111	**Very good identification**					**67733111**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67733112	**Low discrimination**					**67733112**
Actinomyces naeslundii		56.54	0.58	ESL 12		
Actinomyces israelii		26.04	0.43	MLZ 5		
Bifidobacterium adolescentis		7.29	0.28			
67733113	**Very good identification**					**67733113**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67733120	**Very good identification**					**67733120**
					SPO	
Actinomyces naeslundii		97.36	0.80	ESL 12	1	
Clostridium perfringens		1.08	0.24		84	
Clostridium ramosum		0.82	0.19		92	
67733121	**Very good identification**					**67733121**
Actinomyces naeslundii		100.0	0.83	ESL 12		
67733122	**Good identification**					**67733122**
					SPO	
Actinomyces naeslundii		94.49	0.80	ESL 12	1	
Clostridium ramosum		3.18	0.30		92	
Actinomyces israelii		0.98	0.37		1	
67733123	**Very good identification**					**67733123**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.83	ESL 12		
67733130	**Very good identification**					**67733130**
Actinomyces naeslundii		96.83	0.58	ESL 12		
Actinomyces israelii		1.90	0.20	MLZ 5		
Bifidobacterium breve		1.28	0.12			
67733131	**Very good identification**					**67733131**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67733132	**Low discrimination**					**67733132**
Actinomyces naeslundii		85.15	0.58	ESL 12		
Bifidobacterium breve		11.36	0.29	MLZ 5		
Actinomyces israelii		2.50	0.23			
67733133	**Very good identification**					**67733133**
Actinomyces naeslundii		100.0	0.61	ESL 12		
67733140	**Low discrimination**					**67733140**
Actinomyces naeslundii		80.23	0.67	ESL 12		
Actinomyces israelii		19.12	0.47			
Propionibacterium propionicum		0.65	0.21			
67733141	**Very good identification**					**67733141**
Actinomyces naeslundii		99.84	0.70	ESL 12		
Actinomyces israelii		0.16	0.13			
67733142	**Low discrimination**					**67733142**
Actinomyces naeslundii		71.64	0.67	ESL 12		
Actinomyces israelii		25.61	0.50			
Bifidobacterium breve		1.60	0.24			
67733143	**Very good identification**					**67733143**
Actinomyces naeslundii		99.76	0.70	ESL 12		
Actinomyces israelii		0.24	0.16			
67733150	**Low discrimination**					**67733150**
Actinomyces naeslundii		59.65	0.45	ESL 12		
Actinomyces israelii		40.35	0.33	MLZ 5		
67733151	**Good identification**					**67733151**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67733152	**Low discrimination**					**67733152**
Actinomyces israelii		40.11	0.36	CEL 93		
Actinomyces naeslundii		39.52	0.45	XYL 92		
Bifidobacterium breve		16.81	0.24	MLZ 13		
67733153	**Good identification**					**67733153**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67733160	**Very good identification**					**67733160**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		96.98	0.67	ESL 12		
Actinomyces israelii		1.48	0.27			
Clostridium perfringens		1.16	0.12			
67733161	**Very good identification**					**67733161**
Actinomyces naeslundii		100.0	0.70	ESL 12		
67733162	**Good identification**					**67733162**
Actinomyces naeslundii		93.61	0.67	ESL 12		
Bifidobacterium breve		3.73	0.29			
Actinomyces israelii		2.14	0.30			
67733163	**Very good identification**					**67733163**
Actinomyces naeslundii		100.0	0.70	ESL 12		
67733170	**Acceptable identification**					**67733170**
Actinomyces naeslundii		93.04	0.45	ESL 12		
Bifidobacterium breve		6.96	0.12	MLZ 5		
67733171	**Good identification**					**67733171**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67733172	**Low discrimination**					**67733172**
Actinomyces naeslundii		54.92	0.45	ESL 12		
Bifidobacterium breve		41.52	0.29	MLZ 5		
Actinomyces israelii		3.56	0.16			
67733173	**Good identification**					**67733173**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67733200	**Low discrimination**					**67733200**
Actinomyces naeslundii		63.56	0.38	ESL 12		
Actinomyces israelii		36.44	0.25	RHA 1		
67733201	**Good identification**					**67733201**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733202	**Low discrimination**					**67733202**
Actinomyces naeslundii		53.76	0.38	ESL 12		
Actinomyces israelii		46.24	0.28	RHA 1		
67733203	**Good identification**					**67733203**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733220	**Good identification**					**67733220**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67733221	**Good identification**					**67733221**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733222	**Low discrimination**					**67733222**
Actinomyces naeslundii		88.73	0.38	ESL 12	SPO 1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium ramosum		11.27	-.03	RHA 1	92	
67733223	**Good identification**					**67733223**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733300	**Low discrimination**					**67733300**
Actinomyces naeslundii		55.38	0.46	ESL 12		
Actinomyces israelii		44.62	0.35	RHA 1		
67733301	**Good identification**					**67733301**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67733302	**Low discrimination**					**67733302**
					SPO	
Actinomyces israelii		53.64	0.38	CEL 93	1	
Actinomyces naeslundii		44.38	0.46	XYL 92	1	
Clostridium ramosum		1.98	-.03	RHA 7	92	
67733303	**Good identification**					**67733303**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67733321	**Good identification**					**67733321**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67733323	**Good identification**					**67733323**
					SPO	
Actinomyces naeslundii		97.11	0.49	ESL 12	1	
Clostridium ramosum		2.89	-.03	RHA 1	92	
67733341	**Good identification**					**67733341**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67733343	**Good identification**					**67733343**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67733361	**Good identification**					**67733361**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67733363	**Good identification**					**67733363**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67733400	**Low discrimination**					**67733400**
Actinomyces israelii		86.17	0.43	CEL 93		
Actinomyces naeslundii		13.83	0.38	XYL 92		
67733401	**Good identification**					**67733401**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733402	**Acceptable identification**					**67733402**
Actinomyces israelii		90.34	0.46	CEL 93		
Actinomyces naeslundii		9.66	0.38	XYL 92		
67733403	**Good identification**					**67733403**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733420	**Low discrimination**					**67733420**
Actinomyces naeslundii		71.54	0.38	ESL 12		
Actinomyces israelii		28.46	0.22	ARA 1		
67733421	**Good identification**					**67733421**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733422	**Low discrimination**					**67733422**
Actinomyces naeslundii		62.63	0.38	ESL 12		
Actinomyces israelii		37.37	0.25	ARA 1		
67733423	**Good identification**					**67733423**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67733440	**Acceptable identification**					**67733440**
Actinomyces israelii		93.21	0.36	CEL 93		
Actinomyces naeslundii		6.79	0.25	XYL 92		
67733442	**Good identification**					**67733442**
Actinomyces israelii		95.37	0.39	CEL 93		
Actinomyces naeslundii		4.63	0.25	XYL 92		
67733500	**Low discrimination**					**67733500**
					GRA	
Actinomyces israelii		77.82	0.53	CEL 93	99	
Mitsuokella multacida		13.29	0.27	XYL 92	1	
Actinomyces naeslundii		8.89	0.46		99	
67733501	**Low discrimination**					**67733501**
					GRA	
Actinomyces naeslundii		85.49	0.49	ESL 12	99	
Mitsuokella multacida		9.47	0.11	ARA 1	1	
Actinomyces israelii		5.04	0.19		99	
67733502	**Low discrimination**					**67733502**
Actinomyces israelii		73.06	0.56	CEL 93		
Bifidobacterium adolescentis		20.45	0.40	XYL 92		
Actinomyces naeslundii		5.56	0.46			
67733503	**Acceptable identification**					**67733503**
Actinomyces naeslundii		91.87	0.49	ESL 12		
Actinomyces israelii		8.13	0.22	ARA 1		
67733510	**Low discrimination**					**67733510**
					GRA	
Actinomyces israelii		68.69	0.39	CEL 93	99	
Mitsuokella multacida		26.17	0.19	XYL 92	1	
Actinomyces naeslundii		2.76	0.24	MLZ 13	99	
67733512	**Low discrimination**					**67733512**
Bifidobacterium adolescentis		67.67	0.42	NIT 1		
Actinomyces israelii		29.56	0.42	XYL 87		
Bifidobacterium dentium		1.14	-.07			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67733520	**Low discrimination**					**67733520**
					GRA	
Actinomyces naeslundii		51.73	0.46	ESL 12	99	
Actinomyces israelii		28.92	0.32	ARA 1	99	
Mitsuokella multacida		19.35	0.17		1	
67733521	**Good identification**					**67733521**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67733522	**Low discrimination**					**67733522**
Actinomyces naeslundii		54.39	0.46	ESL 12		
Actinomyces israelii		45.61	0.35	ARA 1		
67733523	**Good identification**					**67733523**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67733540	**Low discrimination**					**67733540**
					GRA	
Actinomyces israelii		67.07	0.46	CEL 93	99	
Mitsuokella multacida		29.46	0.27	XYL 92	1	
Actinomyces naeslundii		3.48	0.33		99	
67733541	**Low discrimination**					**67733541**
					GRA	
Actinomyces naeslundii		61.43	0.36	ESL 12	99	
Mitsuokella multacida		38.57	0.11	ARA 1	1	
67733542	**Low discrimination**					**67733542**
Actinomyces israelii		86.50	0.49	CEL 93		
Bifidobacterium adolescentis		7.70	0.25	XYL 92		
Actinomyces naeslundii		2.99	0.33			
67733543	**Low discrimination**					**67733543**
Actinomyces naeslundii		83.69	0.36	ESL 12		
Actinomyces israelii		16.31	0.15	ARA 1		
67733561	**Good identification**					**67733561**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67733563	**Good identification**					**67733563**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67733702	**Low discrimination**					**67733702**
					GRA	
Actinomyces israelii		76.21	0.37	CEL 93	99	
Mitsuokella multacida		23.79	0.15	XYL 92	1	
67734612	**Good identification**					**67734612**
Clostridium clostridioforme		100.0	0.40	NIT 1		
67734632	**Good identification**					**67734632**
Clostridium clostridioforme		100.0	0.40	NIT 1		
67734702	**Good identification**					**67734702**
Clostridium clostridioforme		96.73	0.37	NIT 1		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Mitsuokella multacida		3.27	0.09			
67734712	**Good identification**					**67734712**
Clostridium clostridioforme		99.75	0.49	NIT 1		
Mitsuokella multacida		0.25	0.01			
67734722	**Low discrimination**					**67734722**
Clostridium clostridioforme		88.31	0.37	NIT 1		
Parabacteroides distasonis		11.69	0.28			
67734732	**Good identification**					**67734732**
Clostridium clostridioforme		98.46	0.49	NIT 1		
Parabacteroides distasonis		1.54	0.23			
67735000	**Good identification**					**67735000**
					GRA	
Actinomyces israelii		95.37	0.49	CEL 93	99	
Actinomyces naeslundii		3.18	0.33		99	
Fusobacterium mortiferum		1.45	0.16		1	
67735001	**Low discrimination**					**67735001**
Actinomyces naeslundii		83.19	0.36	ESL 12		
Actinomyces israelii		16.81	0.15	XYL 1		
67735002	**Good identification**					**67735002**
Actinomyces israelii		97.30	0.52	CEL 93		
Actinomyces naeslundii		2.16	0.33			
Bifidobacterium adolescentis		0.54	0.07			
67735003	**Low discrimination**					**67735003**
Actinomyces naeslundii		76.74	0.36	ESL 12		
Actinomyces israelii		23.26	0.18	XYL 1		
67735010	**Good identification**					**67735010**
Actinomyces israelii		100.0	0.35	CEL 93		
67735012	**Low discrimination**					**67735012**
Actinomyces israelii		86.23	0.38	CEL 93		
Clostridium clostridioforme		9.84	0.13	MLZ 13		
Bifidobacterium adolescentis		3.93	0.09			
67735021	**Good identification**					**67735021**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67735023	**Good identification**					**67735023**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67735040	**Good identification**					**67735040**
Actinomyces israelii		100.0	0.42	CEL 93		
67735042	**Good identification**					**67735042**
Actinomyces israelii		100.0	0.45	CEL 93		
67735100	**Low discrimination**					**67735100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	SPO	
Actinomyces israelii		84.18	0.60	CEL 93	1	1	
Atopobium parvulum		5.40	0.25		99	1	
Clostridium butyricum		3.71	0.15		1	99	
67735101	**Low discrimination**						**67735101**
					GRA		
Actinomyces naeslundii		72.67	0.44	ESL 12	99		
Actinomyces israelii		20.64	0.25	XYL 1	99		
Mitsuokella multacida		6.70	0.04		1		
67735102	**Low discrimination**						**67735102**
Actinomyces israelii		77.02	0.63	CEL 93			
Bifidobacterium adolescentis		10.62	0.42				
Bifidobacterium longum subsp. infantis		8.07	0.26				
67735103	**Low discrimination**						**67735103**
Actinomyces naeslundii		67.35	0.44	ESL 12			
Actinomyces israelii		28.69	0.29	XYL 1			
Bifidobacterium adolescentis		3.96	0.07				
67735110	**Acceptable identification**						**67735110**
					GRA		
Actinomyces israelii		91.04	0.46	CEL 93	99		
Mitsuokella multacida		5.99	0.13	MLZ 13	1		
Bifidobacterium adolescentis		1.56	0.09		99		
67735112	**Low discrimination**						**67735112**
					GRA		
Bifidobacterium adolescentis		47.68	0.43	NIT 1	99		
Actinomyces israelii		42.26	0.49	ARA 87	99		
Leptotrichia buccalis		4.71	0.07		1		
67735120	**Low discrimination**						**67735120**
Actinomyces israelii		29.89	0.39	CEL 93			
Clostridium perfringens		23.74	0.24	NAG 6			
Bifidobacterium longum subsp. infantis		12.94	0.13				
67735121	**Good identification**						**67735121**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67735122	**Low discrimination**						**67735122**
					GRA		
Parabacteroides distasonis		69.88	0.58	NIT 1	1		
Bifidobacterium longum subsp. infantis		15.23	0.30		99		
Actinomyces israelii		5.22	0.42		99		
67735123	**Low discrimination**						**67735123**
					GRA		
Actinomyces naeslundii		69.28	0.44	ESL 12	99		
Parabacteroides distasonis		25.23	0.24	XYL 1	1		
Bifidobacterium longum subsp. infantis		5.50	-.04		99		
67735132	**Acceptable identification**						**67735132**
					GRA		
Parabacteroides distasonis		93.30	0.53	NIT 1	1		
Clostridium clostridioforme		2.89	0.22		50		
Actinomyces israelii		1.93	0.28		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67735140	**Acceptable identification**						**67735140**
					GRA		
Actinomyces israelii		91.20	0.53	CEL 93	99		
Mitsuokella multacida		6.92	0.21		1		
Actinomyces naeslundii		0.98	0.28		99		
67735142	**Acceptable identification**						**67735142**
Actinomyces israelii		94.64	0.56	CEL 93			
Bifidobacterium adolescentis		4.15	0.26				
Actinomyces naeslundii		0.68	0.28				
67735150	**Low discrimination**						**67735150**
					GRA		
Actinomyces israelii		85.52	0.38	CEL 93	99		
Mitsuokella multacida		14.48	0.13	MLZ 13	1		
67735152	**Low discrimination**						**67735152**
Actinomyces israelii		73.61	0.42	CEL 93			
Bifidobacterium adolescentis		26.39	0.28	MLZ 13			
67735162	**Low discrimination**						**67735162**
					GRA	CAT	
Actinomyces israelii		55.18	0.35	CEL 93	99	1	
Parabacteroides distasonis		19.19	0.24	NAG 6	1	76	
Capnocytophaga ochracea		8.51	0.05		1	90	
67735300	**Low discrimination**						**67735300**
					GRA		
Actinomyces israelii		38.97	0.40	CEL 93	99		
Eubacterium contortum		32.65	0.24	RHA 7	99		
Mitsuokella multacida		28.38	0.25		1		
67735302	**Low discrimination**						**67735302**
					GRA		
Actinomyces israelii		54.93	0.43	CEL 93	99		
Parabacteroides distasonis		18.91	0.32	RHA 7	1		
Clostridium clostridioforme		16.77	0.25		50		
67735312	**Low discrimination**						**67735312**
					GRA		
Clostridium clostridioforme		78.74	0.36	NIT 1	50		
Parabacteroides distasonis		10.50	0.27	ARA 94	1		
Actinomyces israelii		8.46	0.29		99		
67735322	**Very good identification**						**67735322**
Parabacteroides distasonis		98.30	0.66	NIT 1			
Clostridium clostridioforme		0.88	0.25				
Prevotella oralis		0.52	0.14				
67735332	**Good identification**						**67735332**
Parabacteroides distasonis		92.96	0.61	NIT 1			
Clostridium clostridioforme		7.04	0.36				
67735342	**Low discrimination**						**67735342**
					GRA		
Actinomyces israelii		87.82	0.36	CEL 93	99		
Mitsuokella multacida		12.18	0.09	RHA 7	1		
67735400	**Low discrimination**						**67735400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Actinomyces israelii		81.63	0.48	CEL 93	99	1	
Mitsuokella multacida		11.78	0.21		1	1	
Clostridium butyricum		6.60	0.08		99	99	
67735402	**Low discrimination**						**67735402**
Actinomyces israelii		87.04	0.51	CEL 93			
Clostridium clostridioforme		6.24	0.22				
Bifidobacterium adolescentis		3.97	0.22				
67735440	**Low discrimination**						**67735440**
					GRA		
Actinomyces israelii		72.94	0.41	CEL 93	99		
Mitsuokella multacida		27.06	0.21		1		
67735442	**Good identification**						**67735442**
					GRA		
Actinomyces israelii		97.33	0.44	CEL 93	99		
Mitsuokella multacida		2.68	0.04		1		
67735500	**Low discrimination**						**67735500**
					GRA	SPO	
Mitsuokella multacida		70.35	0.55	CEL 99	1	1	
Actinomyces israelii		19.70	0.58		99	1	
Clostridium butyricum		9.55	0.32		99	99	
67735501	**Good identification**						**67735501**
					GRA	SPO	
Mitsuokella multacida		96.36	0.39	CEL 99	1	1	
Actinomyces israelii		2.45	0.24	URE 10	99	1	
Clostridium butyricum		1.19	-.03		99	99	
67735502	**Low discrimination**						**67735502**
					GRA		
Bifidobacterium adolescentis		43.61	0.56	NIT 1	99		
Actinomyces israelii		38.66	0.61		99		
Mitsuokella multacida		10.23	0.39		1		
67735510	**Low discrimination**						**67735510**
					GRA	SPO	
Mitsuokella multacida		85.30	0.47	CEL 99	1	1	
Actinomyces israelii		10.71	0.44		99	1	
Clostridium butyricum		1.83	0.10		99	99	
67735512	**Low discrimination**						**67735512**
Bifidobacterium adolescentis		72.34	0.57	NIT 1			
Clostridium clostridioforme		14.34	0.42				
Actinomyces israelii		7.84	0.47				
67735520	**Low discrimination**						**67735520**
					GRA		
Mitsuokella multacida		82.28	0.45	CEL 99	1		
Bacteroides vulgatus		5.94	0.16		1		
Actinomyces israelii		5.88	0.38		99		
67735522	**Low discrimination**						**67735522**
					GRA		
Parabacteroides distasonis		73.42	0.54	NIT 1	1		
Mitsuokella multacida		7.58	0.28		1		
Actinomyces israelii		7.32	0.41		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67735530	**Low discrimination**						**67735530**
					GRA		
Mitsuokella multacida		88.25	0.37	CEL 99	1		
Bifidobacterium longum subsp. longum		6.14	0.21		99		
Actinomyces israelii		2.83	0.24		99		
67735532	**Low discrimination**						**67735532**
Parabacteroides distasonis		50.53	0.49	NIT 1			
Clostridium clostridioforme		40.04	0.42				
Mitsuokella multacida		3.23	0.20				
67735540	**Low discrimination**						**67735540**
					GRA	SPO	
Mitsuokella multacida		89.23	0.55	CEL 99	1	1	
Actinomyces israelii		9.72	0.51		99	1	
Clostridium butyricum		1.05	0.14		99	99	
67735541	**Good identification**						**67735541**
					GRA		
Mitsuokella multacida		99.02	0.39	CEL 99	1		
Actinomyces israelii		0.98	0.17	URE 10	99		
67735542	**Low discrimination**						**67735542**
					GRA		
Actinomyces israelii		48.38	0.54	CEL 93	99		
Mitsuokella multacida		32.91	0.39		1		
Bifidobacterium adolescentis		17.35	0.40		99		
67735550	**Acceptable identification**						**67735550**
					COC	GRA	
Mitsuokella multacida		94.83	0.47	CEL 99	1	1	
Actinomyces israelii		4.63	0.37		1	99	
Blautia producta		0.34	-.03		99	99	
67735552	**Low discrimination**						**67735552**
					GRA		
Bifidobacterium adolescentis		53.35	0.42	NIT 1	99		
Mitsuokella multacida		27.60	0.30	SOR 11	1		
Actinomyces israelii		18.19	0.40		99		
67735560	**Good identification**						**67735560**
					GRA	SPO	
Mitsuokella multacida		96.91	0.45	CEL 99	1	1	
Actinomyces israelii		2.70	0.31		99	1	
Clostridium butyricum		0.40	-.05		99	99	
67735570	**Good identification**						**67735570**
					GRA		
Mitsuokella multacida		98.77	0.37	CEL 99	1		
Actinomyces israelii		1.23	0.17		99		
67735602	**Low discrimination**						**67735602**
					GRA		
Clostridium clostridioforme		84.67	0.37	NIT 1	50		
Actinomyces israelii		12.13	0.32		99		
Mitsuokella multacida		3.20	0.09		1		
67735612	**Good identification**						**67735612**
					GRA		
Clostridium clostridioforme		99.26	0.48	NIT 1	50		
Actinomyces israelii		0.47	0.18		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Mitsuokella multacida		0.27	0.01		1	
67735622	**Low discrimination**					**67735622**
Clostridium clostridioforme		87.13	0.37	NIT 1		
Parabacteroides distasonis		12.87	0.28			
67735632	**Good identification**					**67735632**
Clostridium clostridioforme		98.28	0.48	NIT 1		
Parabacteroides distasonis		1.72	0.23			
67735700	**Very good identification**					**67735700**
					GRA	
Mitsuokella multacida		97.56	0.60	CEL 99	1	
Actinomyces israelii		1.11	0.39		99	
Eubacterium contortum		0.93	0.23		99	
67735701	**Good identification**					**67735701**
Mitsuokella multacida		100.0	0.43	CEL 99		
67735702	**Low discrimination**					**67735702**
					GRA	
Mitsuokella multacida		46.85	0.43	CEL 99	1	
Clostridium clostridioforme		41.95	0.46	BGL 10	50	
Actinomyces israelii		7.18	0.42		99	
67735710	**Good identification**					**67735710**
					GRA	
Mitsuokella multacida		98.15	0.51	CEL 99	1	
Clostridium clostridioforme		1.35	0.23		50	
Actinomyces israelii		0.50	0.25		99	
67735711	**Good identification**					**67735711**
Mitsuokella multacida		100.0	0.35	CEL 99		
67735712	**Good identification**					**67735712**
					GRA	
Clostridium clostridioforme		90.95	0.57	NIT 1	50	
Mitsuokella multacida		7.43	0.35		1	
Bifidobacterium adolescentis		0.63	0.23		99	
67735720	**Low discrimination**					**67735720**
Mitsuokella multacida		72.40	0.49	CEL 99		
Bacteroides vulgatus		25.34	0.32			
Parabacteroides distasonis		1.27	0.28			
67735722	**Low discrimination**					**67735722**
Parabacteroides distasonis		76.13	0.62	NIT 1		
Clostridium clostridioforme		17.43	0.46			
Mitsuokella multacida		4.87	0.33			
67735730	**Acceptable identification**					**67735730**
Mitsuokella multacida		91.42	0.41	CEL 99		
Clostridium clostridioforme		5.02	0.23			
Parabacteroides distasonis		2.59	0.23			
67735732	**Low discrimination**					**67735732**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		64.77	0.57	NIT 1			
Parabacteroides distasonis		33.44	0.58				
Mitsuokella multacida		1.32	0.25				
67735740	**Very good identification**						**67735740**
					GRA		
Mitsuokella multacida		99.56	0.60	CEL 99	1		
Actinomyces israelii		0.44	0.32		99		
67735741	**Good identification**						**67735741**
Mitsuokella multacida		100.0	0.43	CEL 99			
67735742	**Acceptable identification**						**67735742**
					GRA		
Mitsuokella multacida		93.58	0.43	CEL 99	1		
Actinomyces israelii		5.58	0.35	BGL 10	99		
Clostridium clostridioforme		0.85	0.11		50		
67735750	**Good identification**						**67735750**
					COC	GRA	
Mitsuokella multacida		99.61	0.51	CEL 99	1	1	
Actinomyces israelii		0.20	0.18		1	99	
Blautia producta		0.19	-.03		99	99	
67735751	**Good identification**						**67735751**
Mitsuokella multacida		100.0	0.35	CEL 99			
67735752	**Low discrimination**						**67735752**
					GRA		
Mitsuokella multacida		86.15	0.35	CEL 99	1		
Clostridium clostridioforme		10.65	0.23	BGL 10	50		
Actinomyces israelii		2.30	0.21		99		
67735760	**Good identification**						**67735760**
Mitsuokella multacida		99.65	0.49	CEL 99			
Bacteroides vulgatus		0.35	-.02				
67735770	**Good identification**						**67735770**
Mitsuokella multacida		100.0	0.41	CEL 99			
67736100	**Good identification**						**67736100**
Actinomyces israelii		96.76	0.38	CEL 93			
Actinomyces naeslundii		3.24	0.22	GAL 99			
67736102	**Good identification**						**67736102**
Actinomyces israelii		97.81	0.41	CEL 93			
Actinomyces naeslundii		2.19	0.22	GAL 99			
67736502	**Low discrimination**						**67736502**
					GRA		
Actinomyces israelii		78.63	0.40	CEL 93	99		
Bifidobacterium adolescentis		12.81	0.20	GAL 99	99		
Mitsuokella multacida		8.56	0.11		1		
67737000	**Very good identification**						**67737000**
Actinomyces israelii		98.87	0.62	CEL 93			
Actinomyces naeslundii		1.13	0.38				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67737001	**Low discrimination**					**67737001**
Actinomyces naeslundii		62.90	0.41	ESL 12		
Actinomyces israelii		37.10	0.28	XYL 1		
67737002	**Very good identification**					**67737002**
Actinomyces israelii		99.25	0.65	CEL 93		
Actinomyces naeslundii		0.76	0.38			
67737003	**Low discrimination**					**67737003**
Actinomyces naeslundii		53.05	0.41	ESL 12		
Actinomyces israelii		46.95	0.31	XYL 1		
67737010	**Good identification**					**67737010**
Actinomyces israelii		100.0	0.48	CEL 93		
67737012	**Good identification**					**67737012**
Actinomyces israelii		99.35	0.51	CEL 93		
Bifidobacterium adolescentis		0.65	0.07	MLZ 13		
67737020	**Low discrimination**					**67737020**
Actinomyces israelii		84.83	0.42	CEL 93		
Actinomyces naeslundii		15.17	0.38	NAG 6		
67737021	**Good identification**					**67737021**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67737022	**Low discrimination**					**67737022**
Actinomyces israelii		89.35	0.45	CEL 93		
Actinomyces naeslundii		10.65	0.38	NAG 6		
67737023	**Good identification**					**67737023**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67737040	**Good identification**					**67737040**
Actinomyces israelii		99.49	0.55	CEL 93		
Actinomyces naeslundii		0.52	0.25			
67737042	**Very good identification**					**67737042**
Actinomyces israelii		99.66	0.58	CEL 93		
Actinomyces naeslundii		0.34	0.25			
67737050	**Good identification**					**67737050**
Actinomyces israelii		100.0	0.41	CEL 93		
67737052	**Good identification**					**67737052**
Actinomyces israelii		100.0	0.44	CEL 93		
67737062	**Acceptable identification**					**67737062**
Actinomyces israelii		94.87	0.38	CEL 93		
Actinomyces naeslundii		5.13	0.25	NAG 6		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67737100	**Very good identification**					**67737100**
					GRA	
Actinomyces israelii		97.93	0.73	CEL 93	99	
Mitsuokella multacida		1.19	0.27		1	
Actinomyces naeslundii		0.80	0.46		99	
67737101	**Low discrimination**					**67737101**
					GRA	
Actinomyces naeslundii		51.55	0.49	ESL 12	99	
Actinomyces israelii		42.74	0.38	XYL 1	99	
Mitsuokella multacida		5.71	0.11		1	
67737102	**Very good identification**					**67737102**
Actinomyces israelii		97.44	0.76	CEL 93		
Bifidobacterium adolescentis		1.94	0.40			
Actinomyces naeslundii		0.53	0.46			
67737103	**Low discrimination**					**67737103**
Actinomyces israelii		55.43	0.41	CEL 93		
Actinomyces naeslundii		44.57	0.49	URE 1		
67737110	**Very good identification**					**67737110**
					GRA	
Actinomyces israelii		96.86	0.58	CEL 93	99	
Mitsuokella multacida		2.63	0.19	MLZ 13	1	
Actinomyces naeslundii		0.28	0.24		99	
67737112	**Low discrimination**					**67737112**
Actinomyces israelii		85.50	0.61	CEL 93		
Bifidobacterium adolescentis		13.93	0.42	MLZ 13		
Bifidobacterium dentium		0.24	-.07			
67737120	**Low discrimination**					**67737120**
					GRA	
Actinomyces israelii		85.12	0.52	CEL 93	99	
Actinomyces naeslundii		10.83	0.46	NAG 6	99	
Mitsuokella multacida		4.05	0.17		1	
67737121	**Acceptable identification**					**67737121**
Actinomyces naeslundii		94.98	0.49	ESL 12		
Actinomyces israelii		5.03	0.18	XYL 1		
67737122	**Low discrimination**					**67737122**
					GRA	
Actinomyces israelii		89.77	0.55	CEL 93	99	
Actinomyces naeslundii		7.62	0.46	NAG 6	99	
Parabacteroides distasonis		2.14	0.24		1	
67737123	**Acceptable identification**					**67737123**
Actinomyces naeslundii		92.65	0.49	ESL 12		
Actinomyces israelii		7.35	0.21	XYL 1		
67737130	**Low discrimination**					**67737130**
					GRA	
Actinomyces israelii		86.88	0.38	CEL 93	99	
Mitsuokella multacida		9.22	0.09	MLZ 13	1	
Actinomyces naeslundii		3.90	0.24	NAG 6	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67737132	**Low discrimination**						**67737132**
					GRA		
Actinomyces israelii		82.08	0.41	CEL 93	99		
Parabacteroides distasonis		7.06	0.19	MLZ 13	1		
Bifidobacterium dentium		6.29	-.03	NAG 6	99		
67737140	**Very good identification**						**67737140**
					GRA		
Actinomyces israelii		96.63	0.66	CEL 93	99		
Mitsuokella multacida		3.02	0.27		1		
Actinomyces naeslundii		0.36	0.33		99		
67737142	**Very good identification**						**67737142**
Actinomyces israelii		98.90	0.69	CEL 93			
Bifidobacterium adolescentis		0.63	0.25				
Actinomyces naeslundii		0.24	0.33				
67737150	**Acceptable identification**						**67737150**
					COC	GRA	
Actinomyces israelii		92.58	0.51	CEL 93	1	99	
Mitsuokella multacida		6.45	0.19	MLZ 13	1	1	
Blautia producta		0.97	-.03		99	99	
67737152	**Acceptable identification**						**67737152**
					GRA		
Actinomyces israelii		94.61	0.54	CEL 93	99		
Bifidobacterium adolescentis		4.90	0.26	MLZ 13	99		
Mitsuokella multacida		0.49	0.03		1		
67737160	**Low discrimination**						**67737160**
					GRA		
Actinomyces israelii		84.74	0.45	CEL 93	99		
Mitsuokella multacida		10.37	0.17	NAG 6	1		
Actinomyces naeslundii		4.89	0.33		99		
67737161	**Good identification**						**67737161**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67737162	**Good identification**						**67737162**
Actinomyces israelii		96.29	0.48	CEL 93			
Actinomyces naeslundii		3.71	0.33	NAG 6			
67737163	**Low discrimination**						**67737163**
Actinomyces naeslundii		85.11	0.36	ESL 12			
Actinomyces israelii		14.89	0.14	XYL 1			
67737200	**Good identification**						**67737200**
Actinomyces israelii		100.0	0.43	CEL 93			
67737202	**Good identification**						**67737202**
Actinomyces israelii		100.0	0.46	CEL 93			
67737240	**Good identification**						**67737240**
Actinomyces israelii		100.0	0.36	CEL 93			
67737242	**Good identification**						**67737242**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.39	CEL 93		
67737300	**Low discrimination**					**67737300**
					GRA	
Actinomyces israelii		76.93	0.53	CEL 93	99	
Mitsuokella multacida		23.07	0.32	RHA 7	1	
67737302	**Very good identification**					**67737302**
					GRA	
Actinomyces israelii		97.83	0.56	CEL 93	99	
Mitsuokella multacida		2.17	0.15	RHA 7	1	
67737310	**Low discrimination**					**67737310**
					GRA	
Actinomyces israelii		59.92	0.39	CEL 93	99	
Mitsuokella multacida		40.08	0.24	RHA 7	1	
67737312	**Acceptable identification**					**67737312**
					GRA	
Actinomyces israelii		93.34	0.42	CEL 93	99	
Mitsuokella multacida		4.62	0.07	RHA 7	1	
Bifidobacterium adolescentis		2.04	0.07	MLZ 13	99	
67737322	**Low discrimination**					**67737322**
					GRA	
Actinomyces israelii		47.87	0.36	CEL 93	99	
Parabacteroides distasonis		45.55	0.32	RHA 7	1	
Mitsuokella multacida		4.16	0.05	NAG 6	1	
67737340	**Low discrimination**					**67737340**
					GRA	
Actinomyces israelii		56.47	0.46	CEL 93	99	
Mitsuokella multacida		43.53	0.32	RHA 7	1	
67737342	**Acceptable identification**					**67737342**
					GRA	
Actinomyces israelii		94.60	0.49	CEL 93	99	
Mitsuokella multacida		5.40	0.15	RHA 7	1	
67737352	**Low discrimination**					**67737352**
					GRA	
Actinomyces israelii		88.70	0.35	CEL 93	99	
Mitsuokella multacida		11.30	0.07	RHA 7	1	
67737400	**Good identification**					**67737400**
					GRA	
Actinomyces israelii		94.39	0.61	CEL 93	99	
Mitsuokella multacida		5.61	0.27		1	
67737402	**Very good identification**					**67737402**
					GRA	
Actinomyces israelii		98.91	0.64	CEL 93	99	
Bifidobacterium adolescentis		0.65	0.20		99	
Mitsuokella multacida		0.44	0.11		1	
67737410	**Low discrimination**					**67737410**
					GRA	
Actinomyces israelii		88.30	0.47	CEL 93	99	
Mitsuokella multacida		11.70	0.19	MLZ 13	1	
67737412	**Acceptable identification**					**67737412**
Actinomyces israelii		92.84	0.50	CEL 93		
Bifidobacterium adolescentis		5.00	0.22	MLZ 13		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium dentium		1.25	-.07				
67737420	**Low discrimination**						**67737420**
					GRA		
Actinomyces israelii		81.13	0.40	CEL 93	99		
Mitsuokella multacida		18.88	0.17	NAG 6	1		
67737422	**Good identification**						**67737422**
Actinomyces israelii		100.0	0.43	CEL 93			
67737440	**Low discrimination**						**67737440**
					GRA		
Actinomyces israelii		86.75	0.54	CEL 93	99		
Mitsuokella multacida		13.25	0.27		1		
67737442	**Very good identification**						**67737442**
					GRA		
Actinomyces israelii		98.88	0.57	CEL 93	99		
Mitsuokella multacida		1.12	0.11		1		
67737450	**Low discrimination**						**67737450**
					COC	GRA	
Actinomyces israelii		71.85	0.40	CEL 93	1	99	
Mitsuokella multacida		24.48	0.19	MLZ 13	1	1	
Blautia producta		3.67	-.03		99	99	
67737452	**Good identification**						**67737452**
					GRA		
Actinomyces israelii		97.54	0.43	CEL 93	99		
Mitsuokella multacida		2.46	0.03	MLZ 13	1		
67737462	**Good identification**						**67737462**
Actinomyces israelii		100.0	0.36	CEL 93			
67737500	**Low discrimination**						**67737500**
					GRA	SPO	
Mitsuokella multacida		59.23	0.61	CEL 99	1	1	
Actinomyces israelii		40.29	0.71		99	1	
Clostridium butyricum		0.38	0.15		99	99	
67737501	**Acceptable identification**						**67737501**
					GRA		
Mitsuokella multacida		94.18	0.45	CEL 99	1		
Actinomyces israelii		5.82	0.37	URE 10	99		
67737502	**Low discrimination**						**67737502**
					GRA		
Actinomyces israelii		78.45	0.74	CEL 93	99		
Bifidobacterium adolescentis		12.78	0.54		99		
Mitsuokella multacida		8.54	0.45		1		
67737510	**Low discrimination**						**67737510**
					GRA		
Mitsuokella multacida		75.27	0.53	CEL 99	1		
Actinomyces israelii		22.95	0.57		99		
Bifidobacterium dentium		1.13	0.10		99		
67737511	**Good identification**						**67737511**
					GRA		
Mitsuokella multacida		97.30	0.37	CEL 99	1		
Actinomyces israelii		2.70	0.23	URE 10	99		
67737512	**Low discrimination**						**67737512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium adolescentis		45.76	0.56	NIT 1			
Actinomyces israelii		34.35	0.60				
Bifidobacterium dentium		11.43	0.27				
67737520	**Low discrimination**						**67737520**
					GRA	SPO	
Mitsuokella multacida		85.04	0.51	CEL 99	1	1	
Actinomyces israelii		14.77	0.51		99	1	
Clostridium butyricum		0.19	-.03		99	99	
67737522	**Low discrimination**						**67737522**
					GRA		
Actinomyces israelii		67.38	0.54	CEL 93	99		
Mitsuokella multacida		28.74	0.35	NAG 6	1		
Bifidobacterium adolescentis		1.74	0.20		99		
67737530	**Low discrimination**						**67737530**
					GRA		
Mitsuokella multacida		84.40	0.43	CEL 99	1		
Bifidobacterium dentium		9.03	0.15		99		
Actinomyces israelii		6.57	0.36		99		
67737540	**Low discrimination**						**67737540**
					COC	GRA	
Mitsuokella multacida		78.02	0.61	CEL 99	1	1	
Actinomyces israelii		20.64	0.64		1	99	
Blautia producta		1.30	0.24		99	99	
67737541	**Good identification**						**67737541**
					GRA		
Mitsuokella multacida		97.65	0.45	CEL 99	1		
Actinomyces israelii		2.35	0.30	URE 10	99		
67737542	**Low discrimination**						**67737542**
					GRA		
Actinomyces israelii		74.82	0.67	CEL 93	99		
Mitsuokella multacida		20.96	0.45		1		
Bifidobacterium adolescentis		3.87	0.39		99		
67737550	**Low discrimination**						**67737550**
					COC	GRA	
Mitsuokella multacida		78.83	0.53	CEL 99	1	1	
Blautia producta		11.83	0.32		99	99	
Actinomyces israelii		9.35	0.50		1	99	
67737551	**Good identification**						**67737551**
					COC	GRA	
Mitsuokella multacida		97.62	0.37	CEL 99	1	1	
Blautia producta		1.33	-.03	URE 10	99	99	
Actinomyces israelii		1.05	0.16		1	99	
67737552	**Low discrimination**						**67737552**
					GRA		
Actinomyces israelii		46.50	0.53	CEL 93	99		
Mitsuokella multacida		29.05	0.37	MLZ 13	1		
Bifidobacterium adolescentis		19.69	0.40		99		
67737560	**Acceptable identification**						**67737560**
					COC	GRA	
Mitsuokella multacida		93.03	0.51	CEL 99	1	1	
Actinomyces israelii		6.28	0.44		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		0.69	0.07		99	99	
67737570	**Acceptable identification**						**67737570**
					COC	GRA	
Mitsuokella multacida		91.16	0.43	CEL 99	1	1	
Blautia producta		6.08	0.15		99	99	
Actinomyces israelii		2.76	0.29		1	99	
67737602	**Acceptable identification**						**67737602**
					GRA		
Actinomyces israelii		90.21	0.45	CEL 93	99		
Mitsuokella multacida		9.79	0.15	RHA 7	1		
67737642	**Low discrimination**						**67737642**
					GRA		
Actinomyces israelii		78.17	0.38	CEL 93	99		
Mitsuokella multacida		21.83	0.15	RHA 7	1		
67737700	**Very good identification**						**67737700**
					GRA		
Mitsuokella multacida		97.32	0.66	CEL 99	1		
Actinomyces israelii		2.68	0.52		99		
67737701	**Good identification**						**67737701**
					GRA		
Mitsuokella multacida		99.75	0.50	CEL 99	1		
Actinomyces israelii		0.25	0.18	URE 10	99		
67737702	**Low discrimination**						**67737702**
					GRA		
Mitsuokella multacida		72.25	0.50	CEL 99	1		
Actinomyces israelii		26.89	0.55	BGL 10	99		
Bifidobacterium adolescentis		0.59	0.20		99		
67737710	**Very good identification**						**67737710**
					COC	GRA	
Mitsuokella multacida		98.70	0.58	CEL 99	1	1	
Actinomyces israelii		1.22	0.38		1	99	
Blautia producta		0.08	-.03		99	99	
67737711	**Good identification**						**67737711**
Mitsuokella multacida		100.0	0.41	CEL 99			
67737712	**Low discrimination**						**67737712**
					GRA		
Mitsuokella multacida		79.55	0.41	CEL 99	1		
Actinomyces israelii		13.27	0.41	BGL 10	99		
Clostridium clostridioforme		4.21	0.23		50		
67737720	**Good identification**						**67737720**
					GRA		
Mitsuokella multacida		99.15	0.56	CEL 99	1		
Actinomyces israelii		0.70	0.31		99		
Bacteroides vulgatus		0.15	-.02		1		
67737721	**Good identification**						**67737721**
Mitsuokella multacida		100.0	0.39	CEL 99			
67737722	**Low discrimination**						**67737722**
					GRA		
Mitsuokella multacida		84.18	0.39	CEL 99	1		
Actinomyces israelii		8.00	0.34	BGL 10	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Parabacteroides distasonis		5.70	0.28		1		
67737730	**Good identification**						**67737730**
					GRA		
Mitsuokella multacida		99.69	0.47	CEL 99	1		
Actinomyces israelii		0.31	0.17		99		
67737740	**Very good identification**						**67737740**
					COC	GRA	
Mitsuokella multacida		98.07	0.66	CEL 99	1	1	
Actinomyces israelii		1.05	0.45		1	99	
Blautia producta		0.88	0.24		99	99	
67737741	**Good identification**						**67737741**
Mitsuokella multacida		100.0	0.50	CEL 99			
67737742	**Low discrimination**						**67737742**
					COC	GRA	
Mitsuokella multacida		86.68	0.50	CEL 99	1	1	
Actinomyces israelii		12.54	0.48	BGL 10	1	99	
Blautia producta		0.78	0.07		99	99	
67737750	**Good identification**						**67737750**
					COC	GRA	
Mitsuokella multacida		92.12	0.58	CEL 99	1	1	
Blautia producta		7.44	0.32		99	99	
Actinomyces israelii		0.44	0.31		1	99	
67737751	**Good identification**						**67737751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	CEL 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67737752	**Low discrimination**						**67737752**
					COC	GRA	
Mitsuokella multacida		87.29	0.41	CEL 99	1	1	
Blautia producta		7.05	0.15	BGL 10	99	99	
Actinomyces israelii		5.66	0.34		1	99	
67737760	**Good identification**						**67737760**
					COC	GRA	
Mitsuokella multacida		99.33	0.56	CEL 99	1	1	
Blautia producta		0.40	0.07		99	99	
Actinomyces israelii		0.27	0.24		1	99	
67737761	**Good identification**						**67737761**
Mitsuokella multacida		100.0	0.39	CEL 99			
67737762	**Good identification**						**67737762**
					GRA		
Mitsuokella multacida		96.44	0.39	CEL 99	1		
Actinomyces israelii		3.56	0.27	BGL 10	99		
67737770	**Good identification**						**67737770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	CEL 99	1	1	
Blautia producta		3.47	0.15		99	99	
67741000	**Low discrimination**						**67741000**
Actinomyces naeslundii		76.29	0.43	ESL 12			
Eubacterium tortuosum		23.71	0.06	FRU 95			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67741001	**Good identification**						**67741001**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67741002	**Low discrimination**						**67741002**
Actinomyces naeslundii		57.20	0.43	ESL 12			
Actinomyces viscosus		42.80	0.16	FRU 95			
67741003	**Low discrimination**						**67741003**
Actinomyces naeslundii		85.74	0.46	ESL 12			
Actinomyces viscosus		14.26	0.08	FRU 95			
67741020	**Good identification**						**67741020**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67741021	**Good identification**						**67741021**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67741022	**Low discrimination**						**67741022**
					GRA		
Actinomyces naeslundii		68.37	0.43	ESL 12	99		
Prevotella oralis		31.63	0.11	FRU 95	1		
67741023	**Good identification**						**67741023**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67741100	**Low discrimination**						**67741100**
Actinomyces naeslundii		85.10	0.50	ESL 12			
Collinsella aerofaciens		9.08	-.03	FRU 95			
Actinomyces israelii		3.13	0.17				
67741101	**Good identification**						**67741101**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67741102	**Low discrimination**						**67741102**
Actinomyces naeslundii		47.17	0.50	ESL 12			
Actinomyces viscosus		22.05	0.21	FRU 95			
Bifidobacterium breve		15.09	0.27				
67741103	**Acceptable identification**						**67741103**
Actinomyces naeslundii		90.59	0.53	ESL 12			
Actinomyces viscosus		9.41	0.12	FRU 95			
67741120	**Low discrimination**						**67741120**
					GRA	SPO	
Actinomyces naeslundii		57.57	0.50	ESL 12	99	1	
Clostridium paraputrificum		25.62	0.13	FRU 95	99	99	
Prevotella oralis		7.72	0.10		1	1	
67741121	**Good identification**						**67741121**
Actinomyces naeslundii		100.0	0.53	ESL 12			
67741123	**Good identification**						**67741123**
					GRA		
Actinomyces naeslundii		98.31	0.53	ESL 12	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Prevotella oralis		1.69	-.03	FRU 95	1	
67741140	**Low discrimination**					**67741140**
Actinomyces naeslundii		84.79	0.38	ESL 12		
Bifidobacterium breve		15.21	0.10	FRU 95		
67741141	**Good identification**					**67741141**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67741143	**Good identification**					**67741143**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67741160	**Low discrimination**					**67741160**
					SPO	
Actinomyces naeslundii		66.29	0.38	ESL 12	1	
Bifidobacterium breve		21.13	0.15	FRU 95	1	
Clostridium paraputrificum		12.58	-.07		99	
67741161	**Good identification**					**67741161**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67741163	**Good identification**					**67741163**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67743000	**Acceptable identification**					**67743000**
Actinomyces naeslundii		92.91	0.48	ESL 12		
Actinomyces israelii		7.10	0.19	FRU 95		
67743001	**Good identification**					**67743001**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67743002	**Low discrimination**					**67743002**
Actinomyces naeslundii		89.72	0.48	ESL 12		
Actinomyces israelii		10.28	0.22	FRU 95		
67743003	**Good identification**					**67743003**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67743020	**Good identification**					**67743020**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67743021	**Good identification**					**67743021**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67743022	**Good identification**					**67743022**
Actinomyces naeslundii		100.0	0.48	ESL 12		
67743023	**Good identification**					**67743023**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67743041	**Good identification**					**67743041**
Actinomyces naeslundii		100.0	0.38	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67743043	**Good identification**					**67743043**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67743061	**Good identification**					**67743061**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67743063	**Good identification**					**67743063**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67743100	**Low discrimination**					**67743100**
Actinomyces naeslundii		89.00	0.55	ESL 12		
Actinomyces israelii		9.55	0.30	FRU 95		
Bifidobacterium breve		1.45	0.10			
67743101	**Very good identification**					**67743101**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67743102	**Low discrimination**					**67743102**
Actinomyces naeslundii		75.42	0.55	ESL 12		
Bifidobacterium breve		12.43	0.27	FRU 95		
Actinomyces israelii		12.14	0.33			
67743103	**Very good identification**					**67743103**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67743111	**Good identification**					**67743111**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67743113	**Good identification**					**67743113**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67743120	**Good identification**					**67743120**
					SPO	
Actinomyces naeslundii		95.79	0.55	ESL 12	1	
Bifidobacterium breve		2.78	0.15	FRU 95	1	
Clostridium ramosum		1.43	-.01		92	
67743121	**Very good identification**					**67743121**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67743122	**Low discrimination**					**67743122**
					SPO	
Actinomyces naeslundii		73.20	0.55	ESL 12	1	
Bifidobacterium breve		21.45	0.32	FRU 95	1	
Clostridium ramosum		4.38	0.09		92	
67743123	**Very good identification**					**67743123**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67743131	**Good identification**					**67743131**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67743133	**Good identification**					**67743133**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.37	ESL 12		
67743140	**Low discrimination**					**67743140**
Actinomyces naeslundii		75.25	0.43	ESL 12		
Actinomyces israelii		17.80	0.23	FRU 95		
Bifidobacterium breve		6.95	0.10			
67743141	**Good identification**					**67743141**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67743142	**Low discrimination**					**67743142**
Actinomyces naeslundii		43.69	0.43	ESL 12		
Bifidobacterium breve		40.81	0.27	FRU 95		
Actinomyces israelii		15.50	0.26			
67743143	**Good identification**					**67743143**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67743160	**Low discrimination**					**67743160**
Actinomyces naeslundii		85.90	0.43	ESL 12		
Bifidobacterium breve		14.11	0.15	FRU 95		
67743161	**Good identification**					**67743161**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67743163	**Good identification**					**67743163**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67745102	**Low discrimination**					**67745102**
Actinomyces israelii		88.93	0.38	FRU 99		
Bifidobacterium adolescentis		11.07	0.16			
67747000	**Good identification**					**67747000**
Actinomyces israelii		100.0	0.37	FRU 99		
67747002	**Good identification**					**67747002**
Actinomyces israelii		100.0	0.40	FRU 99		
67747100	**Good identification**					**67747100**
Actinomyces israelii		100.0	0.48	FRU 99		
67747102	**Good identification**					**67747102**
Actinomyces israelii		98.23	0.51	FRU 99		
Bifidobacterium adolescentis		1.77	0.15			
67747112	**Low discrimination**					**67747112**
Actinomyces israelii		87.18	0.37	FRU 99		
Bifidobacterium adolescentis		12.82	0.16	MLZ 13		
67747140	**Good identification**					**67747140**
Actinomyces israelii		100.0	0.41	FRU 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67747142	**Good identification**					**67747142**
Actinomyces israelii		100.0	0.44	FRU 99		
67747400	**Good identification**					**67747400**
Actinomyces israelii		100.0	0.36	FRU 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67747402	**Good identification**						**67747402**
Actinomyces israelii		100.0	0.39	FRU 99			
67747500	**Low discrimination**						**67747500**
					GRA	SPO	
Actinomyces israelii		69.77	0.46	FRU 99	99	1	
Mitsuokella multacida		29.05	0.27		1	1	
Clostridium butyricum		1.19	-.05		99	99	
67747502	**Low discrimination**						**67747502**
					GRA		
Actinomyces israelii		84.90	0.49	FRU 99	99		
Bifidobacterium adolescentis		12.48	0.29		99		
Mitsuokella multacida		2.62	0.11		1		
67747542	**Low discrimination**						**67747542**
					GRA		
Actinomyces israelii		88.81	0.42	FRU 99	99		
Mitsuokella multacida		7.04	0.11		1		
Bifidobacterium adolescentis		4.15	0.13		99		
67750001	**Acceptable identification**						**67750001**
Actinomyces naeslundii		93.83	0.44	ESL 12			
Actinomyces viscosus		6.17	-.01	GAL 96			
67750002	**Good identification**						**67750002**
					GRA		
Actinomyces viscosus		96.14	0.42	ESL 1	99		
Actinomyces naeslundii		3.28	0.41		99		
Fusobacterium mortiferum		0.58	0.17		1		
67750020	**Good identification**						**67750020**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67750021	**Good identification**						**67750021**
Actinomyces naeslundii		100.0	0.44	ESL 12			
67750022	**Low discrimination**						**67750022**
					GRA		
Actinomyces naeslundii		48.42	0.41	ESL 12	99		
Prevotella oralis		28.30	0.11	GAL 96	1		
Actinomyces viscosus		14.33	0.08		99		
67750023	**Acceptable identification**						**67750023**
Actinomyces naeslundii		93.83	0.44	ESL 12			
Actinomyces viscosus		6.17	-.01	GAL 96			
67750100	**Low discrimination**						**67750100**
					GRA		
Fusobacterium mortiferum		48.26	0.39	NIT 1	1		
Actinomyces naeslundii		36.80	0.49	MLT 11	99		
Actinomyces viscosus		6.80	0.12		99		
67750101	**Good identification**						**67750101**
Actinomyces naeslundii		96.05	0.52	ESL 12			
Actinomyces viscosus		3.95	0.04	GAL 96			
67750102	**Low discrimination**						**67750102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces viscosus		89.99	0.46	ESL 1			
Actinomyces naeslundii		4.92	0.49				
Bifidobacterium breve		1.99	0.27				
67750103	**Low discrimination**						**67750103**
Actinomyces viscosus		80.27	0.38	ESL 1			
Actinomyces naeslundii		19.74	0.52				
67750120	**Low discrimination**						**67750120**
					GRA	SPO	
Actinomyces naeslundii		51.19	0.49	ESL 12	99	1	
Clostridium paraputrificum		28.78	0.13	GAL 96	99	99	
Prevotella oralis		8.67	0.10		1	1	
67750121	**Good identification**						**67750121**
Actinomyces naeslundii		100.0	0.52	ESL 12			
67750123	**Acceptable identification**						**67750123**
					GRA		
Actinomyces naeslundii		94.10	0.52	ESL 12	99		
Actinomyces viscosus		3.87	0.04	GAL 96	99		
Prevotella oralis		2.04	-.03		1		
67750140	**Low discrimination**						**67750140**
Actinomyces naeslundii		81.53	0.36	ESL 12			
Bifidobacterium breve		18.47	0.10	GAL 96			
67750141	**Good identification**						**67750141**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67750143	**Low discrimination**						**67750143**
Actinomyces naeslundii		81.11	0.39	ESL 12			
Actinomyces viscosus		18.89	0.04	GAL 96			
67750160	**Low discrimination**						**67750160**
					SPO		
Actinomyces naeslundii		60.89	0.36	ESL 12	1		
Bifidobacterium breve		24.52	0.15	GAL 96	1		
Clostridium paraputrificum		14.60	-.07		99		
67750161	**Good identification**						**67750161**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67750163	**Good identification**						**67750163**
Actinomyces naeslundii		100.0	0.39	ESL 12			
67751000	**Low discrimination**						**67751000**
Actinomyces naeslundii		66.44	0.65	ESL 12			
Lactobacillus catenaformis		9.63	0.29				
Actinomyces israelii		9.06	0.41				
67751001	**Very good identification**						**67751001**
Actinomyces naeslundii		100.0	0.68	ESL 12			
67751002	**Low discrimination**						**67751002**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		40.62	0.65	ESL 12			
Actinomyces viscosus		30.39	0.38				
Actinomyces israelii		8.31	0.44				
67751003	**Low discrimination**						**67751003**
Actinomyces naeslundii		85.74	0.68	ESL 12			
Actinomyces viscosus		14.26	0.30				
67751010	**Low discrimination**						**67751010**
Actinomyces naeslundii		51.60	0.43	ESL 12			
Lactobacillus jensenii		23.78	0.15	MLZ 5			
Actinomyces israelii		19.97	0.26				
67751011	**Good identification**						**67751011**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67751012	**Low discrimination**						**67751012**
					GRA		
Actinomyces naeslundii		26.42	0.43	ESL 12	99		
Bifidobacterium breve		24.06	0.27	MLZ 5	99		
Leptotrichia buccalis		22.28	0.07		1		
67751013	**Good identification**						**67751013**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67751020	**Low discrimination**						**67751020**
					GRA	SPO	
Actinomyces naeslundii		74.00	0.65	ESL 12	99	1	
Prevotella oralis		9.39	0.24		1	1	
Clostridium paraputrificum		4.93	0.13		99	99	
67751021	**Very good identification**						**67751021**
Actinomyces naeslundii		100.0	0.68	ESL 12			
67751022	**Low discrimination**						**67751022**
					GRA	SPO	
Prevotella oralis		61.52	0.45	NIT 1	1	1	
Actinomyces naeslundii		25.53	0.65	LAC 94	99	1	
Clostridium barati		3.73	0.24		99	99	
67751023	**Very good identification**						**67751023**
					GRA		
Actinomyces naeslundii		98.40	0.68	ESL 12	99		
Prevotella oralis		1.60	0.11		1		
67751030	**Low discrimination**						**67751030**
Actinomyces naeslundii		78.41	0.43	ESL 12			
Bifidobacterium breve		12.56	0.15	MLZ 5			
Lactobacillus jensenii		9.03	0.05				
67751031	**Good identification**						**67751031**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67751033	**Good identification**						**67751033**
Actinomyces naeslundii		100.0	0.46	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67751040	**Low discrimination**					**67751040**
Actinomyces naeslundii		75.34	0.52	ESL 12		
Actinomyces israelii		22.63	0.34			
Bifidobacterium breve		2.02	0.10			
67751041	**Good identification**					**67751041**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67751042	**Low discrimination**					**67751042**
Actinomyces naeslundii		54.45	0.52	ESL 12		
Actinomyces israelii		24.54	0.37			
Bifidobacterium breve		14.79	0.27			
67751043	**Good identification**					**67751043**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67751060	**Low discrimination**					**67751060**
Actinomyces naeslundii		88.52	0.52	ESL 12		
Bifidobacterium breve		4.23	0.15			
Clostridium perfringens		2.61	0.03			
67751061	**Good identification**					**67751061**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67751062	**Low discrimination**					**67751062**
					GRA	
Actinomyces naeslundii		54.74	0.52	ESL 12	99	
Bifidobacterium breve		26.43	0.32		99	
Prevotella oralis		7.55	0.11		1	
67751063	**Good identification**					**67751063**
Actinomyces naeslundii		100.0	0.55	ESL 12		
67751100	**Low discrimination**					**67751100**
Actinomyces naeslundii		45.16	0.72	ESL 12		
Collinsella aerofaciens		25.10	0.32			
Actinomyces israelii		8.65	0.51			
67751101	**Very good identification**					**67751101**
Actinomyces naeslundii		99.29	0.75	ESL 12		
Collinsella aerofaciens		0.37	-.03			
Actinomyces israelii		0.13	0.17			
67751102	**Low discrimination**					**67751102**
					GRA	
Bifidobacterium breve		32.34	0.62	NIT 1	99	
Actinomyces naeslundii		19.39	0.72		99	
Prevotella oralis		13.54	0.44		1	
67751103	**Low discrimination**					**67751103**
Actinomyces naeslundii		88.75	0.75	ESL 12		
Actinomyces viscosus		9.22	0.34			
Bifidobacterium breve		1.00	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67751110	**Low discrimination**						**67751110**
Bifidobacterium breve		60.68	0.44	NIT 1			
Actinomyces naeslundii		19.37	0.50	BGL 91			
Actinomyces israelii		10.53	0.37				
67751111	**Good identification**						**67751111**
Actinomyces naeslundii		97.93	0.53	ESL 12			
Bifidobacterium breve		2.07	0.10	MLZ 5			
67751112	**Low discrimination**						**67751112**
					GRA		
Bifidobacterium breve		48.51	0.62	NIT 1	99		
Leptotrichia buccalis		44.92	0.41		1		
Bifidobacterium adolescentis		2.19	0.38		99		
67751113	**Low discrimination**						**67751113**
					GRA		
Actinomyces naeslundii		69.89	0.53	ESL 12	99		
Bifidobacterium breve		14.91	0.27	MLZ 5	99		
Leptotrichia buccalis		13.81	0.07		1		
67751120	**Low discrimination**						**67751120**
					GRA	SPO	
Clostridium paraputrificum		46.88	0.47	NIT 1	99	99	
Actinomyces naeslundii		20.22	0.72	RAF 7	99	1	
Prevotella oralis		14.12	0.44		1	1	
67751121	**Very good identification**						**67751121**
					GRA	SPO	
Actinomyces naeslundii		97.59	0.75	ESL 12	99	1	
Clostridium paraputrificum		1.52	0.13		99	99	
Prevotella oralis		0.46	0.10		1	1	
67751122	**Low discrimination**						**67751122**
					GRA	CAT	
Prevotella oralis		59.58	0.66	NIT 1	1	1	
Capnocytophaga ochracea		14.34	0.56		1	90	
Bifidobacterium breve		13.31	0.66		99	1	
67751123	**Low discrimination**						**67751123**
					GRA	CAT	
Actinomyces naeslundii		87.50	0.75	ESL 12	99	25	
Prevotella oralis		7.82	0.32		1	1	
Capnocytophaga ochracea		1.88	0.21		1	90	
67751130	**Low discrimination**						**67751130**
					GRA	CAT	
Bifidobacterium breve		69.93	0.49	NIT 1	99	1	
Actinomyces naeslundii		12.56	0.50	BGL 91	99	25	
Capnocytophaga ochracea		9.40	0.23		1	90	
67751131	**Good identification**						**67751131**
Actinomyces naeslundii		96.39	0.53	ESL 12			
Bifidobacterium breve		3.62	0.15	MLZ 5			
67751132	**Low discrimination**						**67751132**
					GRA	CAT	
Bifidobacterium breve		80.07	0.66	NIT 1	99	1	
Capnocytophaga ochracea		9.58	0.39		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		4.63	0.25		1	1	
67751133	**Low discrimination**						**67751133**
					GRA	CAT	
Actinomyces naeslundii		68.32	0.53	ESL 12	99	25	
Bifidobacterium breve		25.91	0.32	MLZ 5	99	1	
Capnocytophaga ochracea		3.10	0.05		1	90	
67751140	**Low discrimination**						**67751140**
Actinomyces naeslundii		41.13	0.59	ESL 12			
Bifidobacterium breve		38.43	0.44				
Actinomyces israelii		17.36	0.44				
67751141	**Very good identification**						**67751141**
Actinomyces naeslundii		99.38	0.62	ESL 12			
Bifidobacterium breve		0.63	0.10				
67751142	**Low discrimination**						**67751142**
Bifidobacterium breve		81.18	0.62	NIT 1			
Actinomyces naeslundii		8.59	0.59				
Actinomyces israelii		5.44	0.47				
67751143	**Good identification**						**67751143**
Actinomyces naeslundii		93.50	0.62	ESL 12			
Bifidobacterium breve		5.95	0.27				
Actinomyces viscosus		0.56	0.00				
67751150	**Low discrimination**						**67751150**
Bifidobacterium breve		88.98	0.44	NIT 1			
Actinomyces israelii		6.01	0.30	BGL 91			
Actinomyces naeslundii		5.01	0.38				
67751151	**Low discrimination**						**67751151**
Actinomyces naeslundii		89.32	0.41	ESL 12			
Bifidobacterium breve		10.68	0.10	MLZ 5			
67751152	**Very good identification**						**67751152**
					GRA		
Bifidobacterium breve		96.79	0.62	NIT 1	99		
Actinomyces israelii		0.97	0.33		99		
Leptotrichia buccalis		0.91	0.07		1		
67751160	**Low discrimination**						**67751160**
					SPO		
Bifidobacterium breve		41.10	0.49	NIT 1	1		
Actinomyces naeslundii		24.74	0.59	BGL 91	1		
Clostridium paraputrificum		24.47	0.27		99		
67751161	**Very good identification**						**67751161**
					SPO		
Actinomyces naeslundii		98.25	0.62	ESL 12	1		
Bifidobacterium breve		1.10	0.15		1		
Clostridium paraputrificum		0.65	-.07		99		
67751162	**Low discrimination**						**67751162**
					GRA	CAT	
Bifidobacterium breve		78.69	0.66	NIT 1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		9.42	0.39		1	90	
Actinomyces naeslundii		4.69	0.59		99	25	
67751163	**Low discrimination**						**67751163**
					GRA	CAT	
Actinomyces naeslundii		88.36	0.62	ESL 12	99	25	
Bifidobacterium breve		9.99	0.32		99	1	
Capnocytophaga ochracea		1.20	0.05		1	90	
67751170	**Acceptable identification**						**67751170**
					GRA	CAT	
Bifidobacterium breve		95.00	0.49	NIT 1	99	1	
Actinomyces naeslundii		3.01	0.38	BGL 91	99	25	
Capnocytophaga ochracea		1.42	0.07		1	90	
67751171	**Low discrimination**						**67751171**
Actinomyces naeslundii		82.47	0.41	ESL 12			
Bifidobacterium breve		17.53	0.15	MLZ 5			
67751172	**Very good identification**						**67751172**
					GRA	CAT	
Bifidobacterium breve		98.28	0.66	NIT 1	99	1	
Capnocytophaga ochracea		1.31	0.23		1	90	
Actinomyces naeslundii		0.31	0.38		99	25	
67751222	**Good identification**						**67751222**
					GRA	SPO	
Prevotella oralis		97.00	0.39	NIT 1	1	1	
Clostridium barati		1.19	0.05	LAC 94	99	99	
Actinomyces naeslundii		0.95	0.30		99	1	
67751301	**Good identification**						**67751301**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67751302	**Low discrimination**						**67751302**
					GRA		
Prevotella oralis		77.97	0.38	NIT 1	1		
Actinomyces israelii		5.64	0.35	NAG 95	99		
Lactobacillus acidophilus		5.36	0.19		99		
67751303	**Low discrimination**						**67751303**
					GRA		
Actinomyces naeslundii		76.72	0.41	ESL 12	99		
Prevotella oralis		15.31	0.04	RHA 1	1		
Actinomyces viscosus		7.97	0.00		99		
67751320	**Low discrimination**						**67751320**
					SPO	GRA	
Prevotella oralis		78.93	0.38	NIT 1	1	1	
Clostridium ramosum		9.33	0.22	BGL 95	92	50	
Clostridium paraputrificum		6.18	0.13		99	99	
67751321	**Low discrimination**						**67751321**
					GRA		
Actinomyces naeslundii		83.37	0.41	ESL 12	99		
Prevotella oralis		16.63	0.04	RHA 1	1		
67751322	**Very good identification**						**67751322**
					SPO	CAT	
Prevotella oralis		95.10	0.60	NIT 1	1	1	
Clostridium ramosum		2.37	0.33		92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea		0.54	0.21		1	90	
67751400	**Low discrimination**						**67751400**
					SPO		
Actinomyces israelii		87.57	0.39	XYL 92	1		
Actinomyces naeslundii		7.93	0.30		1		
Clostridium butyricum		4.50	-.04		99		
67751402	**Low discrimination**						**67751402**
Actinomyces israelii		85.02	0.42	XYL 92			
Bifidobacterium adolescentis		6.01	0.16				
Actinomyces naeslundii		5.13	0.30				
67751442	**Good identification**						**67751442**
Actinomyces israelii		100.0	0.35	XYL 92			
67751500	**Low discrimination**						**67751500**
					GRA	SPO	
Actinomyces israelii		62.20	0.49	XYL 92	99	1	
Clostridium butyricum		19.19	0.20		99	99	
Mitsuokella multacida		7.31	0.21		1	1	
67751501	**Low discrimination**						**67751501**
					GRA		
Actinomyces naeslundii		80.68	0.41	ESL 12	99		
Mitsuokella multacida		10.89	0.04	ARA 1	1		
Actinomyces israelii		8.43	0.15		99		
67751502	**Low discrimination**						**67751502**
					GRA	CAT	
Bifidobacterium adolescentis		53.10	0.50	NIT 1	99	1	
Actinomyces israelii		30.35	0.52	XYL 87	99	1	
Capnocytophaga ochracea		8.08	0.26		1	90	
67751503	**Low discrimination**						**67751503**
Actinomyces naeslundii		65.15	0.41	ESL 12			
Bifidobacterium adolescentis		17.87	0.16	ARA 1			
Actinomyces israelii		10.21	0.18				
67751510	**Low discrimination**						**67751510**
					GRA		
Actinomyces israelii		60.29	0.35	XYL 92	99		
Mitsuokella multacida		15.80	0.13	MLZ 13	1		
Bifidobacterium adolescentis		13.07	0.18		99		
67751512	**Low discrimination**						**67751512**
Bifidobacterium adolescentis		85.69	0.52	NIT 1			
Actinomyces israelii		5.99	0.38	XYL 87			
Bifidobacterium breve		2.87	0.27				
67751521	**Good identification**						**67751521**
Actinomyces naeslundii		100.0	0.41	ESL 12			
67751522	**Low discrimination**						**67751522**
					GRA	CAT	
Capnocytophaga ochracea		71.52	0.43	SAL 1	1	90	
Prevotella oralis		17.01	0.32		1	1	
Bifidobacterium breve		3.80	0.32		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67751523	**Low discrimination**						**67751523**
					GRA	CAT	
Actinomyces naeslundii		68.26	0.41	ESL 12	99	25	
Capnocytophaga ochracea		25.64	0.09	ARA 1	1	90	
Prevotella oralis		6.10	-.03		1	1	
67751540	**Low discrimination**						**67751540**
					GRA	SPO	
Actinomyces israelii		70.05	0.42	XYL 92	99	1	
Mitsuokella multacida		21.16	0.21		1	1	
Clostridium butyricum		4.83	0.01		99	99	
67751542	**Low discrimination**						**67751542**
Actinomyces israelii		53.30	0.45	XYL 92			
Bifidobacterium adolescentis		29.64	0.35				
Bifidobacterium breve		9.82	0.27				
67751552	**Low discrimination**						**67751552**
Bifidobacterium adolescentis		67.08	0.36	NIT 1			
Bifidobacterium breve		18.18	0.27	XYL 87			
Actinomyces israelii		14.75	0.31	SOR 11			
67752000	**Acceptable identification**						**67752000**
Actinomyces naeslundii		91.20	0.46	ESL 12			
Actinomyces israelii		8.80	0.19	GAL 96			
67752001	**Good identification**						**67752001**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67752002	**Low discrimination**						**67752002**
Actinomyces naeslundii		77.09	0.46	ESL 12			
Actinomyces viscosus		11.75	0.08	GAL 96			
Actinomyces israelii		11.15	0.22				
67752003	**Good identification**						**67752003**
Actinomyces naeslundii		96.72	0.49	ESL 12			
Actinomyces viscosus		3.28	-.01	GAL 96			
67752020	**Good identification**						**67752020**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67752021	**Good identification**						**67752021**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67752022	**Good identification**						**67752022**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67752023	**Good identification**						**67752023**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67752041	**Good identification**						**67752041**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67752043	**Good identification**						**67752043**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	ESL 12		
67752061	**Good identification**					**67752061**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67752063	**Good identification**					**67752063**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67752100	**Low discrimination**					**67752100**
Actinomyces naeslundii		86.49	0.54	ESL 12		
Actinomyces israelii		11.73	0.30	GAL 96		
Bifidobacterium breve		1.78	0.10			
67752101	**Very good identification**					**67752101**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67752102	**Low discrimination**					**67752102**
Actinomyces naeslundii		66.37	0.54	ESL 12		
Bifidobacterium breve		13.82	0.27	GAL 96		
Actinomyces israelii		13.50	0.33			
67752103	**Very good identification**					**67752103**
Actinomyces naeslundii		97.93	0.57	ESL 12		
Actinomyces viscosus		2.07	0.04	GAL 96		
67752120	**Acceptable identification**					**67752120**
					SPO	
Actinomyces naeslundii		94.74	0.54	ESL 12	1	
Bifidobacterium breve		3.47	0.15	GAL 96	1	
Clostridium ramosum		1.79	-.01		92	
67752121	**Very good identification**					**67752121**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67752122	**Low discrimination**					**67752122**
					SPO	
Actinomyces naeslundii		68.38	0.54	ESL 12	1	
Bifidobacterium breve		25.31	0.32	GAL 96	1	
Clostridium ramosum		5.17	0.09		92	
67752123	**Very good identification**					**67752123**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67752140	**Low discrimination**					**67752140**
Actinomyces naeslundii		70.65	0.41	ESL 12		
Actinomyces israelii		21.11	0.23	GAL 96		
Bifidobacterium breve		8.25	0.10			
67752141	**Good identification**					**67752141**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67752143	**Good identification**					**67752143**
Actinomyces naeslundii		100.0	0.44	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67752160	**Low discrimination**					**67752160**
Actinomyces naeslundii		82.82	0.41	ESL 12		
Bifidobacterium breve		17.18	0.15	GAL 96		
67752161	**Good identification**					**67752161**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67752163	**Good identification**					**67752163**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67753000	**Low discrimination**					**67753000**
Actinomyces naeslundii		71.41	0.70	ESL 12		
Actinomyces israelii		28.42	0.53			
Bifidobacterium breve		0.17	0.10			
67753001	**Very good identification**					**67753001**
Actinomyces naeslundii		99.73	0.73	ESL 12		
Actinomyces israelii		0.27	0.19			
67753002	**Low discrimination**					**67753002**
Actinomyces naeslundii		61.37	0.70	ESL 12		
Actinomyces israelii		36.63	0.56			
Bifidobacterium breve		1.52	0.27			
67753003	**Very good identification**					**67753003**
Actinomyces naeslundii		99.60	0.73	ESL 12		
Actinomyces israelii		0.40	0.22			
67753010	**Low discrimination**					**67753010**
Actinomyces israelii		51.91	0.39	XYL 92		
Actinomyces naeslundii		45.95	0.48	MLZ 13		
Bifidobacterium breve		2.13	0.10			
67753011	**Good identification**					**67753011**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67753012	**Low discrimination**					**67753012**
Actinomyces israelii		53.56	0.42	XYL 92		
Actinomyces naeslundii		31.61	0.48	MLZ 13		
Bifidobacterium breve		14.83	0.27			
67753013	**Good identification**					**67753013**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67753020	**Very good identification**					**67753020**
Actinomyces naeslundii		96.70	0.70	ESL 12		
Actinomyces israelii		2.46	0.33			
Bifidobacterium breve		0.42	0.15			
67753021	**Very good identification**					**67753021**
Actinomyces naeslundii		100.0	0.73	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67753022	**Low discrimination**					**67753022**
Actinomyces naeslundii		89.94	0.70	ESL 12		
Bifidobacterium breve		3.95	0.32			
Actinomyces israelii		3.43	0.36			
67753023	**Very good identification**					**67753023**
Actinomyces naeslundii		100.0	0.73	ESL 12		
67753030	**Low discrimination**					**67753030**
Actinomyces naeslundii		86.61	0.48	ESL 12		
Bifidobacterium breve		7.14	0.15	MLZ 5		
Actinomyces israelii		6.25	0.19			
67753031	**Good identification**					**67753031**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67753032	**Low discrimination**					**67753032**
Actinomyces naeslundii		51.49	0.48	ESL 12		
Bifidobacterium breve		42.94	0.32	MLZ 5		
Actinomyces israelii		5.57	0.22			
67753033	**Good identification**					**67753033**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67753040	**Low discrimination**					**67753040**
Actinomyces naeslundii		52.89	0.57	ESL 12		
Actinomyces israelii		46.38	0.46			
Bifidobacterium breve		0.73	0.10			
67753041	**Very good identification**					**67753041**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67753042	**Low discrimination**					**67753042**
Actinomyces israelii		53.57	0.49	XYL 92		
Actinomyces naeslundii		40.73	0.57			
Bifidobacterium breve		5.70	0.27			
67753043	**Very good identification**					**67753043**
Actinomyces naeslundii		99.12	0.60	ESL 12		
Actinomyces israelii		0.88	0.15			
67753051	**Good identification**					**67753051**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753052	**Low discrimination**					**67753052**
Actinomyces israelii		50.51	0.35	XYL 92		
Bifidobacterium breve		35.96	0.27	MLZ 13		
Actinomyces naeslundii		13.53	0.35			
67753053	**Good identification**					**67753053**
Actinomyces naeslundii		100.0	0.38	ESL 12		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
67753060	**Good identification**					**67753060**
Actinomyces naeslundii		92.54	0.57	ESL 12		
Actinomyces israelii		5.18	0.26			
Bifidobacterium breve		2.28	0.15			
67753061	**Very good identification**					**67753061**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67753062	**Low discrimination**					**67753062**
Actinomyces naeslundii		75.04	0.57	ESL 12		
Bifidobacterium breve		18.67	0.32			
Actinomyces israelii		6.30	0.29			
67753063	**Very good identification**					**67753063**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67753071	**Good identification**					**67753071**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753073	**Good identification**					**67753073**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753100	**Low discrimination**					**67753100**
Actinomyces naeslundii		60.66	0.77	ESL 12		
Actinomyces israelii		33.92	0.64			
Bifidobacterium breve		5.15	0.44			
67753101	**Very good identification**					**67753101**
Actinomyces naeslundii		99.57	0.80	ESL 12		
Actinomyces israelii		0.38	0.30			
Bifidobacterium breve		0.06	0.10			
67753102	**Low discrimination**					**67753102**
Actinomyces naeslundii		36.42	0.77	ESL 12		
Bifidobacterium breve		31.28	0.62			
Actinomyces israelii		30.55	0.67			
67753103	**Very good identification**					**67753103**
Actinomyces naeslundii		98.82	0.80	ESL 12		
Bifidobacterium breve		0.57	0.27			
Actinomyces israelii		0.56	0.33			
67753110	**Low discrimination**					**67753110**
Bifidobacterium breve		38.41	0.44	NIT 1		
Actinomyces israelii		37.79	0.50	BGL 91		
Actinomyces naeslundii		23.80	0.55			
67753111	**Very good identification**					**67753111**
Actinomyces naeslundii		97.89	0.58	ESL 12		
Bifidobacterium breve		1.06	0.10	MLZ 5		
Actinomyces israelii		1.05	0.15			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67753112	**Low discrimination**					**67753112**
Bifidobacterium breve		79.68	0.62	NIT 1		
Actinomyces israelii		11.63	0.53			
Actinomyces naeslundii		4.88	0.55			
67753113	**Low discrimination**					**67753113**
Actinomyces naeslundii		88.82	0.58	ESL 12		
Bifidobacterium breve		9.76	0.27	MLZ 5		
Actinomyces israelii		1.42	0.18			
67753120	**Low discrimination**					**67753120**
					SPO	
Actinomyces naeslundii		77.81	0.77	ESL 12	1	
Bifidobacterium breve		11.75	0.49		1	
Clostridium ramosum		6.06	0.33		92	
67753121	**Very good identification**					**67753121**
					SPO	
Actinomyces naeslundii		99.85	0.80	ESL 12	1	
Bifidobacterium breve		0.10	0.15		1	
Clostridium ramosum		0.05	-.01		92	
67753122	**Low discrimination**					**67753122**
					SPO	
Bifidobacterium breve		50.89	0.66	NIT 1	1	
Actinomyces naeslundii		33.33	0.77		1	
Clostridium ramosum		10.38	0.43		92	
67753123	**Very good identification**					**67753123**
					SPO	
Actinomyces naeslundii		98.73	0.80	ESL 12	1	
Bifidobacterium breve		1.02	0.32		1	
Clostridium ramosum		0.21	0.09		92	
67753130	**Low discrimination**					**67753130**
Bifidobacterium breve		71.52	0.49	NIT 1		
Actinomyces naeslundii		24.93	0.55	BGL 91		
Actinomyces israelii		2.53	0.29			
67753131	**Very good identification**					**67753131**
Actinomyces naeslundii		98.11	0.58	ESL 12		
Bifidobacterium breve		1.90	0.15	MLZ 5		
67753132	**Very good identification**					**67753132**
Bifidobacterium breve		95.53	0.66	NIT 1		
Actinomyces naeslundii		3.29	0.55			
Actinomyces israelii		0.50	0.32			
67753133	**Low discrimination**					**67753133**
Actinomyces naeslundii		83.66	0.58	ESL 12		
Bifidobacterium breve		16.34	0.32	MLZ 5		
67753140	**Low discrimination**					**67753140**
Actinomyces israelii		45.41	0.57	XYL 92		
Actinomyces naeslundii		36.85	0.64			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		17.74	0.44			
67753141	**Very good identification**					**67753141**
Actinomyces naeslundii		98.86	0.67	ESL 12		
Actinomyces israelii		0.82	0.23			
Bifidobacterium breve		0.32	0.10			
67753142	**Low discrimination**					**67753142**
Bifidobacterium breve		62.89	0.62	NIT 1		
Actinomyces israelii		23.89	0.60			
Actinomyces naeslundii		12.92	0.64			
67753143	**Very good identification**					**67753143**
Actinomyces naeslundii		95.67	0.67	ESL 12		
Bifidobacterium breve		3.14	0.27			
Actinomyces israelii		1.19	0.26			
67753150	**Low discrimination**					**67753150**
Bifidobacterium breve		67.03	0.44	NIT 1		
Actinomyces israelii		25.64	0.43	BGL 91		
Actinomyces naeslundii		7.33	0.43			
67753151	**Acceptable identification**					**67753151**
Actinomyces naeslundii		94.20	0.46	ESL 12		
Bifidobacterium breve		5.80	0.10	MLZ 5		
67753152	**Good identification**					**67753152**
Bifidobacterium breve		93.27	0.62	NIT 1		
Actinomyces israelii		5.29	0.46			
Actinomyces naeslundii		1.01	0.43			
67753153	**Low discrimination**					**67753153**
Actinomyces naeslundii		61.63	0.46	ESL 12		
Bifidobacterium breve		38.38	0.27	MLZ 5		
67753160	**Low discrimination**					**67753160**
Actinomyces naeslundii		51.44	0.64	ESL 12		
Bifidobacterium breve		44.02	0.49			
Actinomyces israelii		4.05	0.36			
67753161	**Very good identification**					**67753161**
Actinomyces naeslundii		99.43	0.67	ESL 12		
Bifidobacterium breve		0.57	0.15			
67753162	**Low discrimination**					**67753162**
Bifidobacterium breve		88.22	0.66	NIT 1		
Actinomyces naeslundii		10.20	0.64			
Actinomyces israelii		1.20	0.39			
67753163	**Good identification**					**67753163**
Actinomyces naeslundii		94.49	0.67	ESL 12		
Bifidobacterium breve		5.51	0.32			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67753170	**Acceptable identification**					**67753170**
Bifidobacterium breve		93.00	0.49	NIT 1		
Actinomyces naeslundii		5.72	0.43	BGL 91		
Actinomyces israelii		1.28	0.22			
67753171	**Acceptable identification**					**67753171**
Actinomyces naeslundii		90.13	0.46	ESL 12		
Bifidobacterium breve		9.87	0.15	MLZ 5		
67753172	**Very good identification**					**67753172**
Bifidobacterium breve		99.19	0.66	NIT 1		
Actinomyces naeslundii		0.60	0.43			
Actinomyces israelii		0.20	0.25			
67753201	**Good identification**					**67753201**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753202	**Low discrimination**					**67753202**
Actinomyces israelii		81.64	0.37	XYL 92		
Actinomyces naeslundii		18.36	0.35	RHA 7		
67753203	**Good identification**					**67753203**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753220	**Low discrimination**					**67753220**
					SPO	
Actinomyces naeslundii		67.42	0.35	ESL 12	1	
Clostridium ramosum		19.82	0.01	RHA 1	92	
Actinomyces israelii		12.76	0.14		1	
67753221	**Good identification**					**67753221**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753223	**Good identification**					**67753223**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753300	**Low discrimination**					**67753300**
					SPO	
Actinomyces israelii		77.81	0.45	XYL 92	1	
Actinomyces naeslundii		18.67	0.43	RHA 7	1	
Clostridium ramosum		1.93	0.01		92	
67753301	**Good identification**					**67753301**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67753302	**Low discrimination**					**67753302**
Actinomyces israelii		72.04	0.48	XYL 92		
Actinomyces naeslundii		11.53	0.43	RHA 7		
Bifidobacterium breve		9.90	0.27			
67753303	**Good identification**					**67753303**
Actinomyces naeslundii		95.96	0.46	ESL 12		
Actinomyces israelii		4.04	0.13	RHA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67753320	**Low discrimination**						**67753320**
					SPO		
Clostridium ramosum		86.68	0.35	NIT 1	92		
Actinomyces naeslundii		8.48	0.43	TRE 94	1		
Actinomyces israelii		2.26	0.24		1		
67753321	**Acceptable identification**						**67753321**
					SPO		
Actinomyces naeslundii		93.56	0.46	ESL 12	1		
Clostridium ramosum		6.44	0.01	RHA 1	92		
67753322	**Low discrimination**						**67753322**
					SPO	GRA	
Clostridium ramosum		87.47	0.45	NIT 1	92	50	
Prevotella oralis		6.27	0.25	TRE 94	1	1	
Bifidobacterium breve		3.27	0.32		1	99	
67753323	**Low discrimination**						**67753323**
					SPO		
Actinomyces naeslundii		78.41	0.46	ESL 12	1		
Clostridium ramosum		21.59	0.11	RHA 1	92		
67753340	**Low discrimination**						**67753340**
Actinomyces israelii		86.11	0.38	XYL 92			
Actinomyces naeslundii		9.38	0.30	RHA 7			
Bifidobacterium breve		4.52	0.10				
67753342	**Low discrimination**						**67753342**
Actinomyces israelii		70.13	0.41	XYL 92			
Bifidobacterium breve		24.78	0.27	RHA 7			
Actinomyces naeslundii		5.09	0.30				
67753400	**Good identification**						**67753400**
Actinomyces israelii		96.99	0.52	XYL 92			
Actinomyces naeslundii		3.01	0.35				
67753401	**Low discrimination**						**67753401**
Actinomyces naeslundii		82.17	0.38	ESL 12			
Actinomyces israelii		17.84	0.18	ARA 1			
67753402	**Good identification**						**67753402**
Actinomyces israelii		97.00	0.55	XYL 92			
Actinomyces naeslundii		2.01	0.35				
Bifidobacterium adolescentis		0.99	0.15				
67753403	**Low discrimination**						**67753403**
Actinomyces naeslundii		75.44	0.38	ESL 12			
Actinomyces israelii		24.56	0.21	ARA 1			
67753410	**Good identification**						**67753410**
Actinomyces israelii		100.0	0.38	XYL 92			
67753412	**Acceptable identification**						**67753412**
Actinomyces israelii		92.30	0.41	XYL 92			
Bifidobacterium adolescentis		7.71	0.16	MLZ 13			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67753421	**Good identification**					**67753421**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753423	**Good identification**					**67753423**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67753440	**Good identification**					**67753440**
Actinomyces israelii		98.61	0.45	XYL 92		
Actinomyces naeslundii		1.39	0.22			
67753442	**Good identification**					**67753442**
Actinomyces israelii		99.07	0.48	XYL 92		
Actinomyces naeslundii		0.93	0.22			
67753500	**Good identification**					**67753500**
					GRA	
Actinomyces israelii		92.41	0.62	XYL 92	99	
Mitsuokella multacida		4.47	0.27		1	
Actinomyces naeslundii		2.04	0.43		99	
67753501	**Low discrimination**					**67753501**
					GRA	
Actinomyces naeslundii		68.15	0.46	ESL 12	99	
Actinomyces israelii		20.79	0.28	ARA 1	99	
Mitsuokella multacida		11.06	0.11		1	
67753502	**Low discrimination**					**67753502**
Actinomyces israelii		77.81	0.65	XYL 92		
Bifidobacterium adolescentis		19.66	0.49			
Actinomyces naeslundii		1.15	0.43			
67753503	**Low discrimination**					**67753503**
Actinomyces naeslundii		63.57	0.46	ESL 12		
Actinomyces israelii		29.09	0.31	ARA 1		
Bifidobacterium adolescentis		7.35	0.15			
67753510	**Low discrimination**					**67753510**
					GRA	
Actinomyces israelii		86.83	0.48	XYL 92	99	
Mitsuokella multacida		9.37	0.19	MLZ 13	1	
Bifidobacterium adolescentis		2.72	0.16		99	
67753512	**Low discrimination**					**67753512**
Bifidobacterium adolescentis		65.19	0.50	NIT 1		
Actinomyces israelii		31.55	0.51	XYL 87		
Bifidobacterium breve		2.67	0.27			
67753520	**Low discrimination**					**67753520**
					GRA	
Actinomyces israelii		61.94	0.42	XYL 92	99	
Actinomyces naeslundii		21.42	0.43	NAG 6	99	
Mitsuokella multacida		11.73	0.17		1	
67753521	**Good identification**					**67753521**
Actinomyces naeslundii		100.0	0.46	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67753522	**Low discrimination**						**67753522**
Actinomyces israelii		56.27	0.45	XYL 92			
Bifidobacterium breve		19.81	0.32	NAG 6			
Actinomyces naeslundii		12.97	0.43				
67753523	**Good identification**						**67753523**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67753540	**Low discrimination**						**67753540**
					GRA		
Actinomyces israelii		87.78	0.55	XYL 92	99		
Mitsuokella multacida		10.92	0.27		1		
Actinomyces naeslundii		0.88	0.30		99		
67753542	**Low discrimination**						**67753542**
Actinomyces israelii		88.60	0.58	XYL 92			
Bifidobacterium adolescentis		7.11	0.33				
Bifidobacterium breve		2.88	0.27				
67753550	**Low discrimination**						**67753550**
					COC	GRA	
Actinomyces israelii		74.00	0.41	XYL 92	1	99	
Mitsuokella multacida		20.53	0.19	MLZ 13	1	1	
Blautia producta		3.08	-.03		99	99	
67753552	**Low discrimination**						**67753552**
Actinomyces israelii		52.78	0.44	XYL 92			
Bifidobacterium adolescentis		34.66	0.35	MLZ 13			
Bifidobacterium breve		11.48	0.27				
67753562	**Low discrimination**						**67753562**
Actinomyces israelii		49.75	0.38	XYL 92			
Bifidobacterium breve		45.04	0.32	NAG 6			
Actinomyces naeslundii		5.21	0.30				
67753602	**Good identification**						**67753602**
Actinomyces israelii		100.0	0.36	XYL 92			
67753702	**Low discrimination**						**67753702**
					GRA		
Actinomyces israelii		89.10	0.46	XYL 92	99		
Mitsuokella multacida		7.88	0.15	RHA 7	1		
Bifidobacterium adolescentis		3.02	0.15		99		
67753742	**Low discrimination**						**67753742**
					GRA		
Actinomyces israelii		81.48	0.39	XYL 92	99		
Mitsuokella multacida		18.52	0.15	RHA 7	1		
67754102	**Low discrimination**						**67754102**
Actinomyces israelii		73.79	0.38	GAL 99			
Actinomyces viscosus		17.03	0.12				
Bifidobacterium adolescentis		9.19	0.16				
67754712	**Good identification**						**67754712**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		99.46	0.43	NIT 1			
Mitsuokella multacida		0.54	0.01	TRE 88			
67754732	**Good identification**						**67754732**
Clostridium clostridioforme		100.0	0.43	NIT 1			
67755000	**Very good identification**						**67755000**
					SPO		
Actinomyces israelii		97.62	0.59		1		
Clostridium butyricum		1.28	0.05		99		
Actinomyces naeslundii		0.63	0.30		1		
67755002	**Very good identification**						**67755002**
					GRA		
Actinomyces israelii		97.79	0.62		99		
Prevotella oralis		0.65	0.08		1		
Bifidobacterium adolescentis		0.49	0.16		99		
67755010	**Good identification**						**67755010**
Actinomyces israelii		100.0	0.45	MLZ 13			
67755012	**Acceptable identification**						**67755012**
Actinomyces israelii		94.71	0.48	MLZ 13			
Bifidobacterium adolescentis		3.90	0.18				
Clostridium clostridioforme		1.40	0.07				
67755020	**Low discrimination**						**67755020**
					GRA		
Actinomyces israelii		79.51	0.38	NAG 6	99		
Prevotella oralis		12.46	0.08		1		
Actinomyces naeslundii		8.03	0.30		99		
67755040	**Good identification**						**67755040**
Actinomyces israelii		100.0	0.52				
67755042	**Good identification**						**67755042**
Actinomyces israelii		100.0	0.55				
67755050	**Good identification**						**67755050**
Actinomyces israelii		100.0	0.38	MLZ 13			
67755052	**Good identification**						**67755052**
Actinomyces israelii		100.0	0.41	MLZ 13			
67755100	**Good identification**						**67755100**
					GRA	SPO	
Actinomyces israelii		90.03	0.69		99	1	
Clostridium butyricum		7.10	0.29		99	99	
Mitsuokella multacida		0.75	0.21		1	1	
67755102	**Low discrimination**						**67755102**
Actinomyces israelii		80.60	0.72				
Bifidobacterium adolescentis		10.03	0.50				
Bifidobacterium longum subsp. infantis		2.39	0.26				
67755103	**Low discrimination**						**67755103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		61.52	0.38	URE 1			
Actinomyces naeslundii		27.92	0.41				
Bifidobacterium adolescentis		7.66	0.16				
67755110	**Acceptable identification**						**67755110**
					GRA	SPO	
Actinomyces israelii		93.73	0.55	MLZ 13	99	1	
Clostridium butyricum		2.60	0.07		99	99	
Mitsuokella multacida		1.75	0.13		1	1	
67755112	**Low discrimination**						**67755112**
Bifidobacterium adolescentis		47.95	0.52	NIT 1			
Actinomyces israelii		47.10	0.58	ARA 87			
Bifidobacterium breve		1.61	0.27				
67755120	**Low discrimination**						**67755120**
					GRA	CAT	
Actinomyces israelii		33.24	0.49	NAG 6	99	1	
Capnocytophaga ochracea		20.96	0.29		1	90	
Prevotella oralis		20.40	0.28		1	1	
67755121	**Acceptable identification**						**67755121**
Actinomyces naeslundii		91.43	0.41	ESL 12			
Actinomyces israelii		8.57	0.14	XYL 1			
67755122	**Low discrimination**						**67755122**
					CAT		
Prevotella oralis		45.44	0.50	NIT 1	1		
Capnocytophaga ochracea		22.12	0.45	XYL 11	90		
Bacteroides fragilis		16.26	0.34		99		
67755140	**Very good identification**						**67755140**
					GRA	SPO	
Actinomyces israelii		95.86	0.62		99	1	
Mitsuokella multacida		2.06	0.21		1	1	
Clostridium butyricum		1.69	0.11		99	99	
67755142	**Good identification**						**67755142**
Actinomyces israelii		93.64	0.65				
Bifidobacterium adolescentis		3.71	0.35				
Bifidobacterium breve		1.23	0.27				
67755150	**Acceptable identification**						**67755150**
					GRA		
Actinomyces israelii		94.26	0.48	MLZ 13	99		
Mitsuokella multacida		4.52	0.13		1		
Bifidobacterium breve		1.23	0.10		99		
67755152	**Low discrimination**						**67755152**
Actinomyces israelii		70.86	0.51	MLZ 13			
Bifidobacterium adolescentis		22.93	0.36				
Bifidobacterium breve		6.21	0.27				
67755160	**Low discrimination**						**67755160**
					GRA	CAT	
Actinomyces israelii		71.06	0.42	NAG 6	99	1	
Capnocytophaga ochracea		12.80	0.13		1	90	
Mitsuokella multacida		5.98	0.11		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67755200	**Good identification**						**67755200**
Actinomyces israelii		100.0	0.39	RHA 7			
67755202	**Acceptable identification**						**67755202**
					GRA		
Actinomyces israelii		92.16	0.43	RHA 7	99		
Clostridium clostridioforme		4.34	0.10		50		
Prevotella oralis		3.50	0.02		1		
67755242	**Good identification**						**67755242**
Actinomyces israelii		100.0	0.35	RHA 7			
67755300	**Low discrimination**						**67755300**
					GRA		
Actinomyces israelii		67.22	0.50	RHA 7	99		
Eubacterium contortum		15.95	0.24		99		
Mitsuokella multacida		13.86	0.25		1		
67755302	**Low discrimination**						**67755302**
					GRA		
Actinomyces israelii		78.87	0.53	RHA 7	99		
Prevotella oralis		11.73	0.22		1		
Clostridium clostridioforme		3.11	0.19		50		
67755310	**Low discrimination**						**67755310**
					GRA		
Actinomyces israelii		68.50	0.36	RHA 7	99		
Mitsuokella multacida		31.50	0.17	MLZ 13	1		
67755322	**Low discrimination**						**67755322**
Prevotella oralis		64.72	0.44	NIT 1			
Parabacteroides distasonis		26.12	0.50	XYL 11			
Bacteroides uniformis		4.91	0.19				
67755332	**Low discrimination**						**67755332**
Parabacteroides distasonis		73.36	0.45	NIT 1			
Clostridium clostridioforme		21.47	0.31				
Prevotella oralis		3.41	0.10				
67755340	**Low discrimination**						**67755340**
					GRA		
Actinomyces israelii		65.35	0.43	RHA 7	99		
Mitsuokella multacida		34.65	0.25		1		
67755342	**Good identification**						**67755342**
					GRA		
Actinomyces israelii		96.22	0.46	RHA 7	99		
Mitsuokella multacida		3.78	0.09		1		
67755400	**Low discrimination**						**67755400**
					GRA	SPO	
Actinomyces israelii		84.36	0.57		99	1	
Clostridium butyricum		12.20	0.22		99	99	
Mitsuokella multacida		3.45	0.21		1	1	
67755402	**Good identification**						**67755402**
					SPO		
Actinomyces israelii		91.67	0.60		1		
Bifidobacterium adolescentis		3.77	0.30		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		3.44	0.15		99		
67755410	**Low discrimination**						**67755410**
					GRA	SPO	
Actinomyces israelii		87.56	0.43	MLZ 13	99	1	
Mitsuokella multacida		7.98	0.13		1	1	
Clostridium butyricum		4.46	0.00		99	99	
67755412	**Low discrimination**						**67755412**
Actinomyces israelii		61.27	0.46	MLZ 13			
Bifidobacterium adolescentis		20.62	0.32				
Clostridium clostridioforme		17.30	0.27				
67755420	**Low discrimination**						**67755420**
					GRA	SPO	
Actinomyces israelii		73.70	0.37	NAG 6	99	1	
Clostridium butyricum		14.51	0.04		99	99	
Mitsuokella multacida		11.79	0.11		1	1	
67755422	**Low discrimination**						**67755422**
					GRA		
Actinomyces israelii		69.48	0.40	NAG 6	99		
Bacteroides uniformis		14.60	0.06		1		
Clostridium clostridioforme		10.08	0.16		50		
67755440	**Low discrimination**						**67755440**
					GRA	SPO	
Actinomyces israelii		87.92	0.50		99	1	
Mitsuokella multacida		9.24	0.21		1	1	
Clostridium butyricum		2.84	0.04		99	99	
67755442	**Good identification**						**67755442**
					SPO		
Actinomyces israelii		97.16	0.53		1		
Bifidobacterium adolescentis		1.27	0.15		1		
Clostridium butyricum		0.81	-.03		99		
67755450	**Low discrimination**						**67755450**
					GRA		
Actinomyces israelii		81.02	0.36	MLZ 13	99		
Mitsuokella multacida		18.99	0.13		1		
67755452	**Acceptable identification**						**67755452**
Actinomyces israelii		90.34	0.39	MLZ 13			
Bifidobacterium adolescentis		9.66	0.16				
67755500	**Low discrimination**						**67755500**
					GRA	SPO	
Mitsuokella multacida		34.93	0.55	TRE 99	1	1	
Actinomyces israelii		34.54	0.68		99	1	
Clostridium butyricum		29.96	0.45		99	99	
67755501	**Low discrimination**						**67755501**
					GRA	SPO	
Mitsuokella multacida		85.62	0.39	TRE 99	1	1	
Actinomyces israelii		7.70	0.33	URE 10	99	1	
Clostridium butyricum		6.68	0.11		99	99	
67755502	**Low discrimination**						**67755502**
					SPO		
Bifidobacterium adolescentis		43.64	0.64	NIT 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		42.86	0.71		1		
Clostridium butyricum		9.64	0.38		99		
67755510	**Low discrimination**						**67755510**
					GRA	SPO	
Mitsuokella multacida		61.17	0.47	TRE 99	1	1	
Actinomyces israelii		27.12	0.53		99	1	
Clostridium butyricum		8.29	0.24		99	99	
67755512	**Low discrimination**						**67755512**
Bifidobacterium adolescentis		84.80	0.66	NIT 1			
Actinomyces israelii		10.18	0.56				
Clostridium clostridioforme		2.41	0.36				
67755520	**Low discrimination**						**67755520**
					GRA	SPO	
Mitsuokella multacida		59.30	0.45	TRE 99	1	1	
Clostridium butyricum		17.70	0.27		99	99	
Actinomyces israelii		14.98	0.47		99	1	
67755530	**Low discrimination**						**67755530**
					GRA	SPO	
Mitsuokella multacida		85.34	0.37	TRE 99	1	1	
Actinomyces israelii		9.66	0.33		99	1	
Clostridium butyricum		4.02	0.05		99	99	
67755532	**Low discrimination**						**67755532**
					GRA		
Clostridium clostridioforme		41.10	0.36	NIT 1	50		
Bifidobacterium adolescentis		14.63	0.32	TRE 88	99		
Parabacteroides distasonis		13.43	0.34	RHA 88	1		
67755540	**Low discrimination**						**67755540**
					GRA	SPO	
Mitsuokella multacida		68.44	0.55	TRE 99	1	1	
Actinomyces israelii		26.32	0.61		99	1	
Clostridium butyricum		5.11	0.27		99	99	
67755541	**Good identification**						**67755541**
					GRA	SPO	
Mitsuokella multacida		95.99	0.39	TRE 99	1	1	
Actinomyces israelii		3.36	0.26	URE 10	99	1	
Clostridium butyricum		0.65	-.07		99	99	
67755542	**Low discrimination**						**67755542**
					GRA		
Actinomyces israelii		63.77	0.64		99		
Bifidobacterium adolescentis		20.64	0.49		99		
Mitsuokella multacida		12.28	0.39		1		
67755550	**Low discrimination**						**67755550**
					GRA	SPO	
Mitsuokella multacida		83.71	0.47	TRE 99	1	1	
Actinomyces israelii		14.43	0.46		99	1	
Clostridium butyricum		0.99	0.05		99	99	
67755552	**Low discrimination**						**67755552**
					GRA		
Bifidobacterium adolescentis		64.65	0.50	NIT 1	99		
Actinomyces israelii		24.43	0.49	SOR 11	99		
Mitsuokella multacida		10.49	0.30		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67755560	**Low discrimination**						**67755560**
					GRA	SPO	
Mitsuokella multacida		88.66	0.45	TRE 99	1	1	
Actinomyces israelii		8.71	0.40		99	1	
Clostridium butyricum		2.30	0.09		99	99	
67755562	**Low discrimination**						**67755562**
					GRA	CAT	
Actinomyces israelii		46.00	0.43	NAG 6	99	1	
Mitsuokella multacida		34.70	0.28		1	1	
Capnocytophaga ochracea		10.73	0.16		1	90	
67755570	**Good identification**						**67755570**
					GRA		
Mitsuokella multacida		95.78	0.37	TRE 99	1		
Actinomyces israelii		4.22	0.26		99		
67755602	**Low discrimination**						**67755602**
					GRA		
Actinomyces israelii		50.58	0.41	RHA 7	99		
Clostridium clostridioforme		45.65	0.31		50		
Mitsuokella multacida		3.78	0.09		1		
67755612	**Good identification**						**67755612**
					GRA		
Clostridium clostridioforme		95.93	0.42	NIT 1	50		
Actinomyces israelii		3.49	0.27	TRE 88	99		
Mitsuokella multacida		0.58	0.01		1		
67755632	**Good identification**						**67755632**
Clostridium clostridioforme		100.0	0.42	NIT 1			
67755700	**Good identification**						**67755700**
					GRA	SPO	
Mitsuokella multacida		93.86	0.60	TRE 99	1	1	
Actinomyces israelii		3.76	0.48		99	1	
Clostridium butyricum		1.34	0.20		99	99	
67755701	**Good identification**						**67755701**
					GRA		
Mitsuokella multacida		99.64	0.43	TRE 99	1		
Actinomyces israelii		0.36	0.14	URE 10	99		
67755702	**Low discrimination**						**67755702**
					GRA		
Mitsuokella multacida		47.39	0.43	TRE 99	1		
Actinomyces israelii		25.65	0.51	BGL 10	99		
Clostridium clostridioforme		19.37	0.40		50		
67755710	**Good identification**						**67755710**
					GRA		
Mitsuokella multacida		97.42	0.51	TRE 99	1		
Actinomyces israelii		1.75	0.34		99		
Clostridium clostridioforme		0.61	0.17		50		
67755711	**Good identification**						**67755711**
Mitsuokella multacida		100.0	0.35	TRE 99			
67755712	**Low discrimination**						**67755712**
					GRA		
Clostridium clostridioforme		78.68	0.51	NIT 1	50		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		14.08	0.35	TRE 88	1		
Bifidobacterium adolescentis		3.82	0.32		99		
67755720	**Low discrimination**						**67755720**
					GRA		
Mitsuokella multacida		67.79	0.49	TRE 99	1		
Bacteroides vulgatus		29.97	0.34		1		
Actinomyces israelii		0.69	0.28		99		
67755722	**Low discrimination**						**67755722**
Bacteroides uniformis		54.40	0.37	IND 98			
Parabacteroides distasonis		15.42	0.46	NIT 1			
Clostridium clostridioforme		13.64	0.40				
67755730	**Good identification**						**67755730**
Mitsuokella multacida		95.87	0.41	TRE 99			
Clostridium clostridioforme		2.41	0.17				
Bacteroides vulgatus		1.28	-.01				
67755732	**Low discrimination**						**67755732**
Clostridium clostridioforme		80.02	0.51	NIT 1			
Parabacteroides distasonis		10.70	0.42	TRE 88			
Bacteroides ovatus		4.78	0.20				
67755740	**Very good identification**						**67755740**
					GRA	SPO	
Mitsuokella multacida		98.35	0.60	TRE 99	1	1	
Actinomyces israelii		1.53	0.41		99	1	
Clostridium butyricum		0.12	0.01		99	99	
67755741	**Good identification**						**67755741**
Mitsuokella multacida		100.0	0.43	TRE 99			
67755742	**Low discrimination**						**67755742**
					GRA		
Mitsuokella multacida		81.43	0.43	TRE 99	1		
Actinomyces israelii		17.14	0.44	BGL 10	99		
Bifidobacterium adolescentis		0.74	0.15		99		
67755750	**Good identification**						**67755750**
					COC	GRA	
Mitsuokella multacida		99.12	0.51	TRE 99	1	1	
Actinomyces israelii		0.69	0.27		1	99	
Blautia producta		0.19	-.03		99	99	
67755751	**Good identification**						**67755751**
Mitsuokella multacida		100.0	0.35	TRE 99			
67755752	**Low discrimination**						**67755752**
					GRA		
Mitsuokella multacida		84.44	0.35	TRE 99	1		
Actinomyces israelii		7.97	0.30	BGL 10	99		
Clostridium clostridioforme		4.77	0.17		50		
67755760	**Good identification**						**67755760**
					GRA		
Mitsuokella multacida		99.16	0.49	TRE 99	1		
Bacteroides vulgatus		0.44	-.01		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.40	0.21		99		
67755770	**Good identification**						**67755770**
Mitsuokella multacida		100.0	0.41	TRE 99			
67756000	**Good identification**						**67756000**
Actinomyces israelii		100.0	0.37	GAL 99			
67756002	**Good identification**						**67756002**
Actinomyces israelii		100.0	0.40	GAL 99			
67756100	**Good identification**						**67756100**
Actinomyces israelii		100.0	0.48	GAL 99			
67756102	**Good identification**						**67756102**
Actinomyces israelii		98.23	0.51	GAL 99			
Bifidobacterium adolescentis		1.77	0.15				
67756112	**Low discrimination**						**67756112**
Actinomyces israelii		87.18	0.37	GAL 99			
Bifidobacterium adolescentis		12.82	0.16	MLZ 13			
67756140	**Good identification**						**67756140**
Actinomyces israelii		100.0	0.41	GAL 99			
67756142	**Good identification**						**67756142**
Actinomyces israelii		100.0	0.44	GAL 99			
67756400	**Good identification**						**67756400**
Actinomyces israelii		100.0	0.36	GAL 99			
67756402	**Good identification**						**67756402**
Actinomyces israelii		100.0	0.39	GAL 99			
67756500	**Low discrimination**						**67756500**
					GRA	SPO	
Actinomyces israelii		69.77	0.46	GAL 99	99	1	
Mitsuokella multacida		29.05	0.27		1	1	
Clostridium butyricum		1.19	-.05		99	99	
67756502	**Low discrimination**						**67756502**
					GRA		
Actinomyces israelii		84.90	0.49	GAL 99	99		
Bifidobacterium adolescentis		12.48	0.29		99		
Mitsuokella multacida		2.62	0.11		1		
67756542	**Low discrimination**						**67756542**
					GRA		
Actinomyces israelii		88.81	0.42	GAL 99	99		
Mitsuokella multacida		7.04	0.11		1		
Bifidobacterium adolescentis		4.15	0.13		99		
67757000	**Very good identification**						**67757000**
Actinomyces israelii		99.78	0.72				
Actinomyces naeslundii		0.22	0.35				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67757001	**Low discrimination**					**67757001**
Actinomyces israelii		75.31	0.37	URE 1		
Actinomyces naeslundii		24.69	0.38			
67757002	**Very good identification**					**67757002**
Actinomyces israelii		99.78	0.75			
Actinomyces naeslundii		0.15	0.35			
Bifidobacterium adolescentis		0.07	0.15			
67757003	**Low discrimination**					**67757003**
Actinomyces israelii		82.07	0.40	URE 1		
Actinomyces naeslundii		17.93	0.38			
67757010	**Very good identification**					**67757010**
Actinomyces israelii		100.0	0.57	MLZ 13		
67757012	**Very good identification**					**67757012**
Actinomyces israelii		99.41	0.60	MLZ 13		
Bifidobacterium adolescentis		0.59	0.16			
67757020	**Good identification**					**67757020**
Actinomyces israelii		96.66	0.51	NAG 6		
Actinomyces naeslundii		3.34	0.35			
67757021	**Low discrimination**					**67757021**
Actinomyces naeslundii		83.70	0.38	ESL 12		
Actinomyces israelii		16.30	0.17	XYL 1		
67757022	**Good identification**					**67757022**
Actinomyces israelii		97.75	0.54	NAG 6		
Actinomyces naeslundii		2.25	0.35			
67757023	**Low discrimination**					**67757023**
Actinomyces naeslundii		77.40	0.38	ESL 12		
Actinomyces israelii		22.60	0.20	XYL 1		
67757030	**Good identification**					**67757030**
Actinomyces israelii		100.0	0.37	MLZ 13		
67757032	**Good identification**					**67757032**
Actinomyces israelii		100.0	0.40	MLZ 13		
67757040	**Very good identification**					**67757040**
Actinomyces israelii		99.90	0.65			
Actinomyces naeslundii		0.10	0.22			
67757042	**Very good identification**					**67757042**
Actinomyces israelii		99.93	0.68			
Actinomyces naeslundii		0.07	0.22			
67757050	**Good identification**					**67757050**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		100.0	0.50	MLZ 13		
67757052	**Good identification**					**67757052**
Actinomyces israelii		100.0	0.53	MLZ 13		
67757060	**Good identification**					**67757060**
Actinomyces israelii		98.46	0.44	NAG 6		
Actinomyces naeslundii		1.55	0.22			
67757062	**Good identification**					**67757062**
Actinomyces israelii		98.97	0.47	NAG 6		
Actinomyces naeslundii		1.04	0.22			
67757100	**Very good identification**					**67757100**
					GRA	
Actinomyces israelii		99.31	0.82		99	
Mitsuokella multacida		0.34	0.27		1	
Actinomyces naeslundii		0.16	0.43		99	
67757101	**Low discrimination**					**67757101**
					GRA	
Actinomyces israelii		78.67	0.48	URE 1	99	
Actinomyces naeslundii		18.35	0.46		99	
Mitsuokella multacida		2.98	0.11		1	
67757102	**Very good identification**					**67757102**
Actinomyces israelii		97.97	0.85			
Bifidobacterium adolescentis		1.76	0.49			
Actinomyces naeslundii		0.10	0.43			
67757103	**Low discrimination**					**67757103**
Actinomyces israelii		85.22	0.51	URE 1		
Actinomyces naeslundii		13.25	0.46			
Bifidobacterium adolescentis		1.53	0.15			
67757110	**Very good identification**					**67757110**
					GRA	
Actinomyces israelii		98.87	0.68	MLZ 13	99	
Mitsuokella multacida		0.76	0.19		1	
Bifidobacterium adolescentis		0.22	0.16		99	
67757112	**Low discrimination**					**67757112**
Actinomyces israelii		86.53	0.71	MLZ 13		
Bifidobacterium adolescentis		12.72	0.50			
Bifidobacterium breve		0.52	0.27			
67757113	**Low discrimination**					**67757113**
Actinomyces israelii		83.21	0.37	MLZ 13		
Bifidobacterium adolescentis		12.23	0.16	URE 1		
Actinomyces naeslundii		4.56	0.24			
67757120	**Very good identification**					**67757120**
					GRA	
Actinomyces israelii		95.38	0.61	NAG 6	99	
Actinomyces naeslundii		2.35	0.43		99	
Mitsuokella multacida		1.29	0.17		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67757121	**Low discrimination**						**67757121**
Actinomyces naeslundii		78.51	0.46	ESL 12			
Actinomyces israelii		21.49	0.27	XYL 1			
67757122	**Good identification**						**67757122**
Actinomyces israelii		92.45	0.64	NAG 6			
Bifidobacterium breve		2.32	0.32				
Actinomyces naeslundii		1.52	0.43				
67757123	**Low discrimination**						**67757123**
Actinomyces naeslundii		70.90	0.46	ESL 12			
Actinomyces israelii		29.10	0.30	XYL 1			
67757130	**Acceptable identification**						**67757130**
					GRA	SPO	
Actinomyces israelii		92.06	0.47	MLZ 13	99	1	
Clostridium tertium		2.88	0.07	NAG 6	99	99	
Mitsuokella multacida		2.77	0.09		1	1	
67757132	**Low discrimination**						**67757132**
					SPO		
Actinomyces israelii		78.42	0.50	MLZ 13	1		
Bifidobacterium breve		13.15	0.32	NAG 6	1		
Clostridium tertium		4.91	0.15		99		
67757140	**Very good identification**						**67757140**
					GRA		
Actinomyces israelii		98.99	0.75		99		
Mitsuokella multacida		0.88	0.27		1		
Actinomyces naeslundii		0.07	0.30		99		
67757141	**Low discrimination**						**67757141**
					GRA		
Actinomyces israelii		83.11	0.41	URE 1	99		
Actinomyces naeslundii		8.80	0.33		99		
Mitsuokella multacida		8.09	0.11		1		
67757142	**Very good identification**						**67757142**
Actinomyces israelii		99.09	0.78				
Bifidobacterium adolescentis		0.57	0.33				
Bifidobacterium breve		0.23	0.27				
67757143	**Acceptable identification**						**67757143**
Actinomyces israelii		93.41	0.44	URE 1			
Actinomyces naeslundii		6.59	0.33				
67757150	**Very good identification**						**67757150**
					COC	GRA	
Actinomyces israelii		97.56	0.61	MLZ 13	1	99	
Mitsuokella multacida		1.93	0.19		1	1	
Blautia producta		0.29	-.03		99	99	
67757152	**Good identification**						**67757152**
Actinomyces israelii		94.02	0.64	MLZ 13			
Bifidobacterium adolescentis		4.39	0.35				
Bifidobacterium breve		1.46	0.27				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67757160	**Acceptable identification**					**67757160**
					GRA	
Actinomyces israelii		94.75	0.54	NAG 6	99	
Mitsuokella multacida		3.28	0.17		1	
Actinomyces naeslundii		1.06	0.30		99	
67757162	**Good identification**					**67757162**
Actinomyces israelii		93.30	0.57	NAG 6		
Bifidobacterium breve		6.01	0.32			
Actinomyces naeslundii		0.69	0.30			
67757170	**Low discrimination**					**67757170**
					GRA	
Actinomyces israelii		87.62	0.40	MLZ 13	99	
Mitsuokella multacida		6.77	0.09	NAG 6	1	
Bifidobacterium breve		5.60	0.15		99	
67757172	**Low discrimination**					**67757172**
Actinomyces israelii		69.88	0.43	MLZ 13		
Bifidobacterium breve		30.12	0.32	NAG 6		
67757200	**Good identification**					**67757200**
Actinomyces israelii		100.0	0.52	RHA 7		
67757202	**Good identification**					**67757202**
Actinomyces israelii		100.0	0.55	RHA 7		
67757210	**Good identification**					**67757210**
Actinomyces israelii		100.0	0.38	RHA 7		
67757212	**Good identification**					**67757212**
Actinomyces israelii		100.0	0.41	RHA 7		
67757240	**Good identification**					**67757240**
Actinomyces israelii		100.0	0.45	RHA 7		
67757242	**Good identification**					**67757242**
Actinomyces israelii		100.0	0.48	RHA 7		
67757300	**Good identification**					**67757300**
					GRA	
Actinomyces israelii		92.18	0.63	RHA 7	99	
Mitsuokella multacida		7.83	0.32		1	
67757302	**Very good identification**					**67757302**
					GRA	
Actinomyces israelii		99.14	0.66	RHA 7	99	
Mitsuokella multacida		0.62	0.15		1	
Bifidobacterium adolescentis		0.24	0.15		99	
67757310	**Low discrimination**					**67757310**
					GRA	
Actinomyces israelii		84.08	0.49	RHA 7	99	
Mitsuokella multacida		15.92	0.24	MLZ 13	1	
67757312	**Good identification**					**67757312**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		96.74	0.52	RHA 7	99		
Bifidobacterium adolescentis		1.91	0.16	MLZ 13	99		
Mitsuokella multacida		1.36	0.07		1		
67757320	**Low discrimination**						**67757320**
					SPO	GRA	
Actinomyces israelii		73.19	0.42	RHA 7	1	99	
Mitsuokella multacida		24.34	0.21	NAG 6	1	1	
Clostridium ramosum		2.47	0.01		92	50	
67757322	**Low discrimination**						**67757322**
					SPO	GRA	
Actinomyces israelii		81.61	0.45	RHA 7	1	99	
Clostridium ramosum		7.35	0.11	NAG 6	92	50	
Prevotella oralis		6.44	0.10		1	1	
67757340	**Low discrimination**						**67757340**
					GRA		
Actinomyces israelii		82.08	0.56	RHA 7	99		
Mitsuokella multacida		17.92	0.32		1		
67757342	**Very good identification**						**67757342**
					GRA		
Actinomyces israelii		98.41	0.59	RHA 7	99		
Mitsuokella multacida		1.59	0.15		1		
67757350	**Low discrimination**						**67757350**
					COC	GRA	
Actinomyces israelii		65.52	0.42	RHA 7	1	99	
Mitsuokella multacida		31.91	0.24	MLZ 13	1	1	
Blautia producta		2.58	-.03		99	99	
67757352	**Good identification**						**67757352**
					GRA		
Actinomyces israelii		96.52	0.45	RHA 7	99		
Mitsuokella multacida		3.48	0.07	MLZ 13	1		
67757360	**Low discrimination**						**67757360**
					GRA		
Actinomyces israelii		53.91	0.35	RHA 7	99		
Mitsuokella multacida		46.09	0.21	NAG 6	1		
67757362	**Acceptable identification**						**67757362**
					GRA		
Actinomyces israelii		94.04	0.38	RHA 7	99		
Mitsuokella multacida		5.96	0.05	NAG 6	1		
67757400	**Very good identification**						**67757400**
					GRA	SPO	
Actinomyces israelii		98.07	0.70		99	1	
Mitsuokella multacida		1.65	0.27		1	1	
Clostridium butyricum		0.28	0.05		99	99	
67757401	**Low discrimination**						**67757401**
					GRA		
Actinomyces israelii		84.39	0.36	URE 1	99		
Mitsuokella multacida		15.62	0.11		1		
67757402	**Very good identification**						**67757402**
					GRA		
Actinomyces israelii		99.21	0.73		99		
Bifidobacterium adolescentis		0.59	0.29		99		
Mitsuokella multacida		0.12	0.11		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67757403	**Good identification**						**67757403**
Actinomyces israelii		100.0	0.39	URE 1			
67757410	**Good identification**						**67757410**
					GRA		
Actinomyces israelii		96.38	0.56	MLZ 13	99		
Mitsuokella multacida		3.62	0.19		1		
67757412	**Good identification**						**67757412**
Actinomyces israelii		94.77	0.59	MLZ 13			
Bifidobacterium adolescentis		4.61	0.30				
Bifidobacterium dentium		0.36	-.07				
67757420	**Acceptable identification**						**67757420**
					GRA		
Actinomyces israelii		93.82	0.50	NAG 6	99		
Mitsuokella multacida		6.18	0.17		1		
67757422	**Good identification**						**67757422**
Actinomyces israelii		100.0	0.53	NAG 6			
67757430	**Low discrimination**						**67757430**
					GRA		
Actinomyces israelii		87.19	0.36	MLZ 13	99		
Mitsuokella multacida		12.81	0.09	NAG 6	1		
67757432	**Acceptable identification**						**67757432**
Actinomyces israelii		90.41	0.39	MLZ 13			
Bifidobacterium dentium		9.59	-.03	NAG 6			
67757440	**Very good identification**						**67757440**
					GRA		
Actinomyces israelii		95.85	0.63		99		
Mitsuokella multacida		4.15	0.27		1		
67757442	**Very good identification**						**67757442**
					GRA		
Actinomyces israelii		99.49	0.66		99		
Mitsuokella multacida		0.32	0.11		1		
Bifidobacterium adolescentis		0.19	0.13		99		
67757450	**Acceptable identification**						**67757450**
					COC	GRA	
Actinomyces israelii		90.01	0.49	MLZ 13	1	99	
Mitsuokella multacida		8.69	0.19		1	1	
Blautia producta		1.30	-.03		99	99	
67757452	**Good identification**						**67757452**
					GRA		
Actinomyces israelii		97.79	0.52	MLZ 13	99		
Bifidobacterium adolescentis		1.51	0.15		99		
Mitsuokella multacida		0.70	0.03		1		
67757460	**Low discrimination**						**67757460**
					GRA		
Actinomyces israelii		85.51	0.43	NAG 6	99		
Mitsuokella multacida		14.49	0.17		1		
67757462	**Good identification**						**67757462**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.46	NAG 6			
67757500	**Low discrimination**						**67757500**
					GRA	SPO	
Actinomyces israelii		69.66	0.80		99	1	
Mitsuokella multacida		29.00	0.61		1	1	
Clostridium butyricum		1.19	0.29		99	99	
67757501	**Low discrimination**						**67757501**
					GRA	SPO	
Mitsuokella multacida		81.83	0.45	TRE 99	1	1	
Actinomyces israelii		17.87	0.46	URE 10	99	1	
Clostridium butyricum		0.30	-.05		99	99	
67757502	**Low discrimination**						**67757502**
					GRA		
Actinomyces israelii		84.57	0.83		99		
Bifidobacterium adolescentis		12.43	0.63		99		
Mitsuokella multacida		2.61	0.45		1		
67757503	**Low discrimination**						**67757503**
					GRA		
Actinomyces israelii		67.28	0.49	URE 1	99		
Mitsuokella multacida		22.82	0.29		1		
Bifidobacterium adolescentis		9.89	0.29		99		
67757510	**Low discrimination**						**67757510**
					GRA		
Actinomyces israelii		50.81	0.66	MLZ 13	99		
Mitsuokella multacida		47.18	0.53		1		
Bifidobacterium adolescentis		0.93	0.30		99		
67757511	**Acceptable identification**						**67757511**
					GRA		
Mitsuokella multacida		91.08	0.37	TRE 99	1		
Actinomyces israelii		8.92	0.32	URE 10	99		
67757512	**Low discrimination**						**67757512**
Bifidobacterium adolescentis		50.80	0.64	NIT 1			
Actinomyces israelii		42.25	0.69				
Bifidobacterium dentium		3.98	0.27				
67757520	**Low discrimination**						**67757520**
					GRA	SPO	
Mitsuokella multacida		61.45	0.51	TRE 99	1	1	
Actinomyces israelii		37.68	0.60		99	1	
Clostridium butyricum		0.87	0.11		99	99	
67757522	**Low discrimination**						**67757522**
					GRA		
Actinomyces israelii		86.20	0.63	NAG 6	99		
Mitsuokella multacida		10.41	0.35		1		
Bifidobacterium adolescentis		2.01	0.29		99		
67757530	**Low discrimination**						**67757530**
					GRA		
Mitsuokella multacida		72.36	0.43	TRE 99	1		
Actinomyces israelii		19.89	0.46		99		
Bifidobacterium dentium		7.74	0.15		99		
67757540	**Low discrimination**						**67757540**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		51.15	0.61	TRE 99	1	1	
Actinomyces israelii		47.78	0.73		1	99	
Blautia producta		0.85	0.24		99	99	
67757541	**Acceptable identification**						**67757541**
					GRA		
Mitsuokella multacida		92.17	0.45	TRE 99	1		
Actinomyces israelii		7.83	0.39	URE 10	99		
67757542	**Low discrimination**						**67757542**
					GRA		
Actinomyces israelii		88.63	0.76		99		
Mitsuokella multacida		7.03	0.45		1		
Bifidobacterium adolescentis		4.14	0.47		99		
67757543	**Low discrimination**						**67757543**
					GRA		
Actinomyces israelii		52.11	0.42	URE 1	99		
Mitsuokella multacida		45.46	0.29		1		
Bifidobacterium adolescentis		2.44	0.13		99		
67757550	**Low discrimination**						**67757550**
					COC	GRA	
Mitsuokella multacida		63.65	0.53	TRE 99	1	1	
Actinomyces israelii		26.65	0.59		1	99	
Blautia producta		9.55	0.32		99	99	
67757551	**Good identification**						**67757551**
					COC	GRA	
Mitsuokella multacida		95.08	0.37	TRE 99	1	1	
Actinomyces israelii		3.62	0.25	URE 10	1	99	
Blautia producta		1.30	-.03		99	99	
67757552	**Low discrimination**						**67757552**
					GRA		
Actinomyces israelii		62.97	0.62	MLZ 13	99		
Bifidobacterium adolescentis		24.07	0.49		99		
Mitsuokella multacida		11.14	0.37		1		
67757560	**Low discrimination**						**67757560**
					COC	GRA	
Mitsuokella multacida		80.18	0.51	TRE 99	1	1	
Actinomyces israelii		19.12	0.53		1	99	
Blautia producta		0.59	0.07		99	99	
67757562	**Low discrimination**						**67757562**
					GRA		
Actinomyces israelii		75.87	0.56	NAG 6	99		
Mitsuokella multacida		23.57	0.35		1		
Bifidobacterium adolescentis		0.56	0.13		99		
67757570	**Low discrimination**						**67757570**
					COC	GRA	
Mitsuokella multacida		85.21	0.43	TRE 99	1	1	
Actinomyces israelii		9.11	0.39		1	99	
Blautia producta		5.68	0.15		99	99	
67757572	**Low discrimination**						**67757572**
					GRA		
Actinomyces israelii		53.54	0.42	MLZ 13	99		
Mitsuokella multacida		37.10	0.26	NAG 6	1		
Bifidobacterium dentium		3.65	-.03		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67757600	**Low discrimination**						**67757600**
					GRA		
Actinomyces israelii		70.67	0.51	RHA 7	99		
Mitsuokella multacida		29.33	0.32		1		
67757602	**Good identification**						**67757602**
					GRA		
Actinomyces israelii		97.02	0.54	RHA 7	99		
Mitsuokella multacida		2.98	0.15		1		
67757610	**Low discrimination**						**67757610**
					GRA		
Actinomyces israelii		51.93	0.37	RHA 7	99		
Mitsuokella multacida		48.08	0.24	MLZ 13	1		
67757612	**Low discrimination**						**67757612**
					GRA		
Actinomyces israelii		89.48	0.40	RHA 7	99		
Mitsuokella multacida		6.14	0.07	MLZ 13	1		
Clostridium clostridioforme		4.39	0.08		50		
67757642	**Acceptable identification**						**67757642**
					GRA		
Actinomyces israelii		92.67	0.47	RHA 7	99		
Mitsuokella multacida		7.33	0.15		1		
67757700	**Good identification**						**67757700**
					GRA	SPO	
Mitsuokella multacida		91.07	0.66	TRE 99	1	1	
Actinomyces israelii		8.87	0.61		99	1	
Clostridium butyricum		0.06	0.03		99	99	
67757701	**Good identification**						**67757701**
					GRA		
Mitsuokella multacida		99.12	0.50	TRE 99	1		
Actinomyces israelii		0.88	0.27	URE 10	99		
67757702	**Low discrimination**						**67757702**
					GRA		
Actinomyces israelii		56.06	0.64	RHA 7	99		
Mitsuokella multacida		42.66	0.50		1		
Bifidobacterium adolescentis		1.11	0.29		99		
67757710	**Very good identification**						**67757710**
					COC	GRA	
Mitsuokella multacida		95.74	0.58	TRE 99	1	1	
Actinomyces israelii		4.18	0.47		1	99	
Blautia producta		0.08	-.03		99	99	
67757711	**Good identification**						**67757711**
Mitsuokella multacida		100.0	0.41	TRE 99			
67757712	**Low discrimination**						**67757712**
					GRA		
Mitsuokella multacida		58.28	0.41	TRE 99	1		
Actinomyces israelii		34.34	0.50	BGL 10	99		
Bifidobacterium adolescentis		5.54	0.30		99		
67757720	**Good identification**						**67757720**
					GRA		
Mitsuokella multacida		97.39	0.56	TRE 99	1		
Actinomyces israelii		2.42	0.41		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides vulgatus		0.19	-.01		1		
67757721	**Good identification**						**67757721**
Mitsuokella multacida		100.0	0.39	TRE 99			
67757722	**Low discrimination**						**67757722**
					GRA		
Mitsuokella multacida		67.83	0.39	TRE 99	1		
Actinomyces israelii		22.76	0.44	BGL 10	99		
Bacteroides ovatus		6.47	0.15		1		
67757730	**Good identification**						**67757730**
					GRA		
Mitsuokella multacida		98.90	0.47	TRE 99	1		
Actinomyces israelii		1.10	0.27		99		
67757740	**Very good identification**						**67757740**
					COC	GRA	
Mitsuokella multacida		95.53	0.66	TRE 99	1	1	
Actinomyces israelii		3.62	0.54		1	99	
Blautia producta		0.86	0.24		99	99	
67757741	**Good identification**						**67757741**
					GRA		
Mitsuokella multacida		99.66	0.50	TRE 99	1		
Actinomyces israelii		0.34	0.20	URE 10	99		
67757742	**Low discrimination**						**67757742**
					COC	GRA	
Mitsuokella multacida		65.65	0.50	TRE 99	1	1	
Actinomyces israelii		33.55	0.57	BGL 10	1	99	
Blautia producta		0.59	0.07		99	99	
67757750	**Good identification**						**67757750**
					COC	GRA	
Mitsuokella multacida		91.10	0.58	TRE 99	1	1	
Blautia producta		7.36	0.32		99	99	
Actinomyces israelii		1.55	0.40		1	99	
67757751	**Good identification**						**67757751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	TRE 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67757752	**Low discrimination**						**67757752**
					COC	GRA	
Mitsuokella multacida		75.66	0.41	TRE 99	1	1	
Actinomyces israelii		17.34	0.43	BGL 10	1	99	
Blautia producta		6.11	0.15		99	99	
67757760	**Good identification**						**67757760**
					COC	GRA	
Mitsuokella multacida		98.65	0.56	TRE 99	1	1	
Actinomyces israelii		0.95	0.34		1	99	
Blautia producta		0.39	0.07		99	99	
67757761	**Good identification**						**67757761**
Mitsuokella multacida		100.0	0.39	TRE 99			
67757762	**Low discrimination**						**67757762**
					GRA		
Mitsuokella multacida		88.46	0.39	TRE 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		11.54	0.37	BGL 10	99		
67757770	**Good identification**						**67757770**
					COC	GRA	
Mitsuokella multacida		96.13	0.47	TRE 99	1	1	
Blautia producta		3.45	0.15		99	99	
Actinomyces israelii		0.42	0.20		1	99	
67761000	**Good identification**						**67761000**
Actinomyces naeslundii		95.18	0.48	ESL 12			
Actinomyces israelii		4.83	0.16	FRU 95			
67761001	**Good identification**						**67761001**
Actinomyces naeslundii		100.0	0.51	ESL 12			
67761002	**Acceptable identification**						**67761002**
Actinomyces naeslundii		92.93	0.48	ESL 12			
Actinomyces israelii		7.07	0.19	FRU 95			
67761003	**Good identification**						**67761003**
Actinomyces naeslundii		100.0	0.51	ESL 12			
67761020	**Good identification**						**67761020**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67761021	**Good identification**						**67761021**
Actinomyces naeslundii		100.0	0.51	ESL 12			
67761022	**Good identification**						**67761022**
Actinomyces naeslundii		100.0	0.48	ESL 12			
67761023	**Good identification**						**67761023**
Actinomyces naeslundii		100.0	0.51	ESL 12			
67761041	**Good identification**						**67761041**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67761043	**Good identification**						**67761043**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67761061	**Good identification**						**67761061**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67761063	**Good identification**						**67761063**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67761100	**Acceptable identification**						**67761100**
Actinomyces naeslundii		93.35	0.55	ESL 12			
Actinomyces israelii		6.65	0.27	FRU 95			
67761101	**Very good identification**						**67761101**
Actinomyces naeslundii		100.0	0.58	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67761102	**Low discrimination**					**67761102**
					GRA	
Actinomyces naeslundii		67.66	0.55	ESL 12	99	
Leptotrichia buccalis		10.33	0.07	FRU 95	1	
Bifidobacterium breve		7.44	0.24		99	
67761103	**Very good identification**					**67761103**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67761111	**Good identification**					**67761111**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67761112	**Good identification**					**67761112**
					GRA	
Leptotrichia buccalis		98.83	0.41	NIT 10	1	
Bifidobacterium breve		0.72	0.24	FRU 99	99	
Actinomyces naeslundii		0.34	0.34		99	
67761120	**Low discrimination**					**67761120**
					SPO	
Actinomyces naeslundii		88.15	0.55	ESL 12	1	
Clostridium paraputrificum		5.37	0.03	FRU 95	99	
Clostridium ramosum		3.64	0.06		92	
67761121	**Very good identification**					**67761121**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67761122	**Low discrimination**					**67761122**
					SPO	
Actinomyces naeslundii		63.37	0.55	ESL 12	1	
Bifidobacterium breve		12.38	0.29	FRU 95	1	
Clostridium ramosum		10.48	0.17		92	
67761123	**Very good identification**					**67761123**
Actinomyces naeslundii		100.0	0.58	ESL 12		
67761131	**Good identification**					**67761131**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67761133	**Low discrimination**					**67761133**
					GRA	
Actinomyces naeslundii		82.32	0.37	ESL 12	99	
Leptotrichia buccalis		17.68	-.09	FRU 95	1	
67761140	**Low discrimination**					**67761140**
Actinomyces naeslundii		86.43	0.43	ESL 12		
Actinomyces israelii		13.57	0.20	FRU 95		
67761141	**Good identification**					**67761141**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67761142	**Low discrimination**					**67761142**
Actinomyces naeslundii		53.81	0.43	ESL 12		
Bifidobacterium breve		33.51	0.24	FRU 95		
Actinomyces israelii		12.67	0.23			
67761143	**Good identification**					**67761143**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.46	ESL 12		
67761160	**Acceptable identification**					**67761160**
Actinomyces naeslundii		90.13	0.43	ESL 12		
Bifidobacterium breve		9.87	0.12	FRU 95		
67761161	**Good identification**					**67761161**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67761163	**Good identification**					**67761163**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67762100	**Good identification**					**67762100**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67762101	**Good identification**					**67762101**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67762102	**Good identification**					**67762102**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67762103	**Good identification**					**67762103**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67762120	**Good identification**					**67762120**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67762121	**Good identification**					**67762121**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67762122	**Good identification**					**67762122**
Actinomyces naeslundii		100.0	0.37	ESL 12		
67762123	**Good identification**					**67762123**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67763000	**Low discrimination**					**67763000**
Actinomyces naeslundii		87.11	0.53	ESL 12		
Actinomyces israelii		12.89	0.29	FRU 95		
67763001	**Good identification**					**67763001**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67763002	**Low discrimination**					**67763002**
Actinomyces naeslundii		81.83	0.53	ESL 12		
Actinomyces israelii		18.17	0.32	FRU 95		
67763003	**Good identification**					**67763003**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67763020	**Good identification**					**67763020**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.53	ESL 12		
67763021	**Good identification**					**67763021**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67763022	**Good identification**					**67763022**
Actinomyces naeslundii		100.0	0.53	ESL 12		
67763023	**Good identification**					**67763023**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67763040	**Low discrimination**					**67763040**
Actinomyces naeslundii		75.41	0.40	ESL 12		
Actinomyces israelii		24.59	0.22	FRU 95		
67763041	**Good identification**					**67763041**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67763042	**Low discrimination**					**67763042**
Actinomyces naeslundii		67.15	0.40	ESL 12		
Actinomyces israelii		32.85	0.25	FRU 95		
67763043	**Good identification**					**67763043**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67763060	**Good identification**					**67763060**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67763061	**Good identification**					**67763061**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67763062	**Good identification**					**67763062**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67763063	**Good identification**					**67763063**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67763100	**Low discrimination**					**67763100**
Actinomyces naeslundii		82.78	0.60	ESL 12		
Actinomyces israelii		17.22	0.39	FRU 95		
67763101	**Very good identification**					**67763101**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67763102	**Low discrimination**					**67763102**
Actinomyces naeslundii		73.07	0.60	ESL 12		
Actinomyces israelii		22.80	0.42	FRU 95		
Bifidobacterium breve		4.14	0.24			
67763103	**Very good identification**					**67763103**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.63	ESL 12		
67763110	**Low discrimination**					**67763110**
Actinomyces naeslundii		62.87	0.38	ESL 12		
Actinomyces israelii		37.13	0.25	FRU 95		
67763111	**Good identification**					**67763111**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67763113	**Good identification**					**67763113**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67763120	**Low discrimination**					**67763120**
					SPO	
Actinomyces naeslundii		87.42	0.60	ESL 12	1	
Clostridium ramosum		10.55	0.19	FRU 95	92	
Actinomyces israelii		1.16	0.19		1	
67763121	**Very good identification**					**67763121**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67763122	**Low discrimination**					**67763122**
					SPO	
Actinomyces naeslundii		62.38	0.60	ESL 12	1	
Clostridium ramosum		30.10	0.30	FRU 95	92	
Bifidobacterium breve		6.28	0.29		1	
67763123	**Very good identification**					**67763123**
Actinomyces naeslundii		100.0	0.63	ESL 12		
67763130	**Low discrimination**					**67763130**
Actinomyces naeslundii		84.10	0.38	ESL 12		
Bifidobacterium breve		15.91	0.12	FRU 95		
67763131	**Good identification**					**67763131**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67763133	**Good identification**					**67763133**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67763140	**Low discrimination**					**67763140**
Actinomyces naeslundii		68.57	0.47	ESL 12		
Actinomyces israelii		31.43	0.32	FRU 95		
67763141	**Good identification**					**67763141**
Actinomyces naeslundii		100.0	0.50	ESL 12		
67763142	**Low discrimination**					**67763142**
Actinomyces naeslundii		49.79	0.47	ESL 12		
Actinomyces israelii		34.23	0.35	FRU 95		
Bifidobacterium breve		15.97	0.24			
67763143	**Good identification**					**67763143**
Actinomyces naeslundii		100.0	0.50	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67763160	**Acceptable identification**						**67763160**
Actinomyces naeslundii		94.66	0.47	ESL 12			
Bifidobacterium breve		5.34	0.12	FRU 95			
67763161	**Good identification**						**67763161**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67763162	**Low discrimination**						**67763162**
Actinomyces naeslundii		61.95	0.47	ESL 12			
Bifidobacterium breve		35.33	0.29	FRU 95			
Actinomyces israelii		2.72	0.15				
67763163	**Good identification**						**67763163**
Actinomyces naeslundii		100.0	0.50	ESL 12			
67763502	**Low discrimination**						**67763502**
					GRA		
Actinomyces israelii		84.85	0.41	FRU 99	99		
Mitsuokella multacida		8.00	0.11	XYL 92	1		
Bifidobacterium adolescentis		3.80	0.12		99		
67765002	**Good identification**						**67765002**
Actinomyces israelii		100.0	0.37	FRU 99			
67765100	**Low discrimination**						**67765100**
					GRA	SPO	
Actinomyces israelii		64.59	0.45	FRU 99	99	1	
Clostridium butyricum		21.21	0.15		99	99	
Mitsuokella multacida		14.20	0.21		1	1	
67765102	**Low discrimination**						**67765102**
					SPO		
Actinomyces israelii		89.01	0.48	FRU 99	1		
Clostridium butyricum		7.58	0.08		99		
Bifidobacterium adolescentis		1.96	0.13		1		
67765140	**Low discrimination**						**67765140**
					GRA	SPO	
Actinomyces israelii		61.02	0.38	FRU 99	99	1	
Mitsuokella multacida		34.50	0.21		1	1	
Clostridium butyricum		4.48	-.03		99	99	
67765142	**Good identification**						**67765142**
					GRA		
Actinomyces israelii		95.98	0.41	FRU 99	99		
Mitsuokella multacida		4.02	0.04		1		
67765402	**Low discrimination**						**67765402**
					GRA	SPO	
Actinomyces israelii		80.92	0.36	FRU 99	99	1	
Clostridium butyricum		12.63	0.01		99	99	
Mitsuokella multacida		6.44	0.04		1	1	
67765500	**Low discrimination**						**67765500**
					GRA	SPO	
Mitsuokella multacida		85.23	0.55	FRU 99	1	1	
Clostridium butyricum		11.57	0.32		99	99	
Actinomyces israelii		3.20	0.43		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67765501	**Good identification**						**67765501**
					GRA	SPO	
Mitsuokella multacida		98.78	0.39	FRU 99	1	1	
Clostridium butyricum		1.22	-.03	URE 10	99	99	
67765502	**Low discrimination**						**67765502**
					GRA	SPO	
Mitsuokella multacida		48.04	0.39	FRU 99	1	1	
Actinomyces israelii		24.37	0.46	BGL 10	99	1	
Clostridium butyricum		22.83	0.25		99	99	
67765510	**Good identification**						**67765510**
					GRA	SPO	
Mitsuokella multacida		96.31	0.47	FRU 99	1	1	
Clostridium butyricum		2.07	0.10		99	99	
Actinomyces israelii		1.62	0.29		99	1	
67765520	**Acceptable identification**						**67765520**
					GRA	SPO	
Mitsuokella multacida		94.62	0.45	FRU 99	1	1	
Clostridium butyricum		4.47	0.14		99	99	
Actinomyces israelii		0.91	0.23		99	1	
67765530	**Good identification**						**67765530**
Mitsuokella multacida		100.0	0.37	FRU 99			
67765540	**Good identification**						**67765540**
					GRA	SPO	
Mitsuokella multacida		97.43	0.55	FRU 99	1	1	
Actinomyces israelii		1.42	0.36		99	1	
Clostridium butyricum		1.15	0.14		99	99	
67765541	**Good identification**						**67765541**
Mitsuokella multacida		100.0	0.39	FRU 99			
67765542	**Low discrimination**						**67765542**
					GRA	SPO	
Mitsuokella multacida		80.00	0.39	FRU 99	1	1	
Actinomyces israelii		15.79	0.39	BGL 10	99	1	
Clostridium butyricum		3.31	0.07		99	99	
67765550	**Good identification**						**67765550**
					COC	GRA	
Mitsuokella multacida		99.00	0.47	FRU 99	1	1	
Actinomyces israelii		0.65	0.22		1	99	
Blautia producta		0.35	-.03		99	99	
67765560	**Good identification**						**67765560**
					GRA	SPO	
Mitsuokella multacida		99.22	0.45	FRU 99	1	1	
Clostridium butyricum		0.41	-.05		99	99	
Actinomyces israelii		0.37	0.16		99	1	
67765570	**Good identification**						**67765570**
Mitsuokella multacida		100.0	0.37	FRU 99			
67765700	**Very good identification**						**67765700**
					GRA	SPO	
Mitsuokella multacida		99.62	0.60	FRU 99	1	1	
Clostridium butyricum		0.23	0.06		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		0.15	0.24		99	1	
67765701	**Good identification**						**67765701**
Mitsuokella multacida		100.0	0.43	FRU 99			
67765702	**Acceptable identification**						**67765702**
					GRA		
Mitsuokella multacida		94.45	0.43	FRU 99	1		
Clostridium clostridioforme		2.86	0.20	BGL 10	50		
Actinomyces israelii		1.94	0.27		99		
67765710	**Good identification**						**67765710**
Mitsuokella multacida		100.0	0.51	FRU 99			
67765711	**Good identification**						**67765711**
Mitsuokella multacida		100.0	0.35	FRU 99			
67765712	**Low discrimination**						**67765712**
Mitsuokella multacida		70.73	0.35	FRU 99			
Clostridium clostridioforme		29.27	0.32	BGL 10			
67765720	**Good identification**						**67765720**
Mitsuokella multacida		100.0	0.49	FRU 99			
67765730	**Good identification**						**67765730**
Mitsuokella multacida		100.0	0.41	FRU 99			
67765740	**Very good identification**						**67765740**
					GRA		
Mitsuokella multacida		99.94	0.60	FRU 99	1		
Actinomyces israelii		0.06	0.17		99		
67765741	**Good identification**						**67765741**
Mitsuokella multacida		100.0	0.43	FRU 99			
67765742	**Good identification**						**67765742**
					GRA		
Mitsuokella multacida		99.21	0.43	FRU 99	1		
Actinomyces israelii		0.79	0.20	BGL 10	99		
67765750	**Good identification**						**67765750**
					COC	GRA	
Mitsuokella multacida		99.81	0.51	FRU 99	1	1	
Blautia producta		0.19	-.03		99	99	
67765751	**Good identification**						**67765751**
Mitsuokella multacida		100.0	0.35	FRU 99			
67765752	**Good identification**						**67765752**
Mitsuokella multacida		100.0	0.35	FRU 99			
67765760	**Good identification**						**67765760**
Mitsuokella multacida		100.0	0.49	FRU 99			
67765770	**Good identification**						**67765770**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.41	FRU 99			
67767000	**Good identification**						**67767000**
Actinomyces israelii		100.0	0.47	FRU 99			
67767002	**Good identification**						**67767002**
Actinomyces israelii		100.0	0.50	FRU 99			
67767012	**Good identification**						**67767012**
Actinomyces israelii		100.0	0.36	FRU 99			
67767040	**Good identification**						**67767040**
Actinomyces israelii		100.0	0.40	FRU 99			
67767042	**Good identification**						**67767042**
Actinomyces israelii		100.0	0.43	FRU 99			
67767100	**Good identification**						**67767100**
					GRA	SPO	
Actinomyces israelii		90.81	0.58	FRU 99	99	1	
Mitsuokella multacida		8.22	0.27		1	1	
Clostridium butyricum		0.59	-.01		99	99	
67767102	**Very good identification**						**67767102**
					GRA		
Actinomyces israelii		98.75	0.61	FRU 99	99		
Mitsuokella multacida		0.66	0.11		1		
Bifidobacterium adolescentis		0.32	0.12		99		
67767110	**Low discrimination**						**67767110**
					GRA		
Actinomyces israelii		83.20	0.43	FRU 99	99		
Mitsuokella multacida		16.80	0.19	MLZ 13	1		
67767112	**Acceptable identification**						**67767112**
Actinomyces israelii		94.21	0.46	FRU 99			
Bifidobacterium adolescentis		2.46	0.13	MLZ 13			
Bifidobacterium dentium		1.93	-.07				
67767120	**Low discrimination**						**67767120**
					GRA		
Actinomyces israelii		70.38	0.37	FRU 99	99		
Mitsuokella multacida		24.96	0.17	NAG 6	1		
Actinomyces naeslundii		4.66	0.26		99		
67767122	**Good identification**						**67767122**
Actinomyces israelii		95.78	0.40	FRU 99			
Actinomyces naeslundii		4.22	0.26	NAG 6			
67767140	**Low discrimination**						**67767140**
					GRA		
Actinomyces israelii		81.11	0.51	FRU 99	99		
Mitsuokella multacida		18.89	0.27		1		
67767142	**Good identification**						**67767142**
					GRA		
Actinomyces israelii		98.30	0.54	FRU 99	99		
Mitsuokella multacida		1.70	0.11		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67767150	**Low discrimination**						**67767150**
					COC	GRA	
Actinomyces israelii		62.61	0.36	FRU 99	1	99	
Mitsuokella multacida		32.52	0.19	MLZ 13	1	1	
Blautia producta		4.88	-.03		99	99	
67767152	**Good identification**						**67767152**
					GRA		
Actinomyces israelii		96.29	0.39	FRU 99	99		
Mitsuokella multacida		3.71	0.03	MLZ 13	1		
67767302	**Low discrimination**						**67767302**
					GRA		
Actinomyces israelii		85.80	0.41	FRU 99	99		
Mitsuokella multacida		14.20	0.15	RHA 7	1		
67767400	**Low discrimination**						**67767400**
					GRA		
Actinomyces israelii		69.32	0.46	FRU 99	99		
Mitsuokella multacida		30.68	0.27		1		
67767402	**Good identification**						**67767402**
					GRA		
Actinomyces israelii		96.83	0.49	FRU 99	99		
Mitsuokella multacida		3.18	0.11		1		
67767442	**Acceptable identification**						**67767442**
					GRA		
Actinomyces israelii		92.22	0.42	FRU 99	99		
Mitsuokella multacida		7.78	0.11		1		
67767500	**Good identification**						**67767500**
					GRA	SPO	
Mitsuokella multacida		91.10	0.61	FRU 99	1	1	
Actinomyces israelii		8.32	0.56		99	1	
Clostridium butyricum		0.59	0.15		99	99	
67767501	**Good identification**						**67767501**
					GRA		
Mitsuokella multacida		99.18	0.45	FRU 99	1		
Actinomyces israelii		0.82	0.22	URE 10	99		
67767502	**Low discrimination**						**67767502**
					GRA		
Actinomyces israelii		53.77	0.59	FRU 99	99		
Mitsuokella multacida		43.64	0.45		1		
Bifidobacterium adolescentis		1.40	0.26		99		
67767510	**Acceptable identification**						**67767510**
					GRA		
Mitsuokella multacida		94.47	0.53	FRU 99	1		
Actinomyces israelii		3.87	0.42		99		
Bifidobacterium dentium		1.42	0.10		99		
67767511	**Good identification**						**67767511**
Mitsuokella multacida		100.0	0.37	FRU 99			
67767520	**Good identification**						**67767520**
					GRA	SPO	
Mitsuokella multacida		97.51	0.51	FRU 99	1	1	
Actinomyces israelii		2.27	0.36		99	1	
Clostridium butyricum		0.22	-.03		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67767530	**Low discrimination**						**67767530**
					GRA		
Mitsuokella multacida		89.49	0.43	FRU 99	1		
Bifidobacterium dentium		9.58	0.15		99		
Actinomyces israelii		0.94	0.22		99		
67767540	**Good identification**						**67767540**
					COC	GRA	
Mitsuokella multacida		94.99	0.61	FRU 99	1	1	
Actinomyces israelii		3.37	0.49		1	99	
Blautia producta		1.58	0.24		99	99	
67767541	**Good identification**						**67767541**
					GRA		
Mitsuokella multacida		99.68	0.45	FRU 99	1		
Actinomyces israelii		0.32	0.15	URE 10	99		
67767542	**Low discrimination**						**67767542**
					COC	GRA	
Mitsuokella multacida		66.68	0.45	FRU 99	1	1	
Actinomyces israelii		31.95	0.52	BGL 10	1	99	
Blautia producta		1.11	0.07		99	99	
67767550	**Low discrimination**						**67767550**
					COC	GRA	
Mitsuokella multacida		85.77	0.53	FRU 99	1	1	
Blautia producta		12.87	0.32		99	99	
Actinomyces israelii		1.37	0.35		1	99	
67767551	**Good identification**						**67767551**
					COC	GRA	
Mitsuokella multacida		98.66	0.37	FRU 99	1	1	
Blautia producta		1.35	-.03	URE 10	99	99	
67767552	**Low discrimination**						**67767552**
					COC	GRA	
Mitsuokella multacida		71.78	0.37	FRU 99	1	1	
Actinomyces israelii		15.42	0.38	BGL 10	1	99	
Blautia producta		10.77	0.15		99	99	
67767560	**Good identification**						**67767560**
					COC	GRA	
Mitsuokella multacida		98.38	0.51	FRU 99	1	1	
Actinomyces israelii		0.89	0.29		1	99	
Blautia producta		0.73	0.07		99	99	
67767570	**Acceptable identification**						**67767570**
					COC	GRA	
Mitsuokella multacida		93.39	0.43	FRU 99	1	1	
Blautia producta		6.23	0.15		99	99	
Actinomyces israelii		0.38	0.15		1	99	
67767700	**Very good identification**						**67767700**
					GRA		
Mitsuokella multacida		99.63	0.66	FRU 99	1		
Actinomyces israelii		0.37	0.37		99		
67767701	**Good identification**						**67767701**
Mitsuokella multacida		100.0	0.50	FRU 99			
67767702	**Good identification**						**67767702**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		95.24	0.50	FRU 99	1		
Actinomyces israelii		4.76	0.40	BGL 10	99		
67767710	**Very good identification**						**67767710**
					COC	GRA	
Mitsuokella multacida		99.75	0.58	FRU 99	1	1	
Actinomyces israelii		0.17	0.23		1	99	
Blautia producta		0.08	-.03		99	99	
67767711	**Good identification**						**67767711**
Mitsuokella multacida		100.0	0.41	FRU 99			
67767712	**Good identification**						**67767712**
					GRA		
Mitsuokella multacida		97.10	0.41	FRU 99	1		
Actinomyces israelii		2.17	0.26	BGL 10	99		
Bifidobacterium dentium		0.72	-.07		99		
67767720	**Good identification**						**67767720**
					GRA		
Mitsuokella multacida		99.91	0.56	FRU 99	1		
Actinomyces israelii		0.09	0.16		99		
67767721	**Good identification**						**67767721**
Mitsuokella multacida		100.0	0.39	FRU 99			
67767722	**Good identification**						**67767722**
					GRA		
Mitsuokella multacida		97.80	0.39	FRU 99	1		
Actinomyces israelii		1.25	0.19	BGL 10	99		
Bacteroides ovatus		0.95	-.02		1		
67767730	**Good identification**						**67767730**
Mitsuokella multacida		100.0	0.47	FRU 99			
67767740	**Very good identification**						**67767740**
					COC	GRA	
Mitsuokella multacida		98.97	0.66	FRU 99	1	1	
Blautia producta		0.89	0.24		99	99	
Actinomyces israelii		0.14	0.30		1	99	
67767741	**Good identification**						**67767741**
Mitsuokella multacida		100.0	0.50	FRU 99			
67767742	**Good identification**						**67767742**
					COC	GRA	
Mitsuokella multacida		97.24	0.50	FRU 99	1	1	
Actinomyces israelii		1.89	0.33	BGL 10	1	99	
Blautia producta		0.87	0.07		99	99	
67767750	**Good identification**						**67767750**
					COC	GRA	
Mitsuokella multacida		92.47	0.58	FRU 99	1	1	
Blautia producta		7.47	0.32		99	99	
Actinomyces israelii		0.06	0.16		1	99	
67767751	**Good identification**						**67767751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	FRU 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67767752	**Acceptable identification**						**67767752**
					COC	GRA	
Mitsuokella multacida		91.79	0.41	FRU 99	1	1	
Blautia producta		7.41	0.15	BGL 10	99	99	
Actinomyces israelii		0.80	0.19		1	99	
67767760	**Good identification**						**67767760**
					COC	GRA	
Mitsuokella multacida		99.60	0.56	FRU 99	1	1	
Blautia producta		0.40	0.07		99	99	
67767761	**Good identification**						**67767761**
Mitsuokella multacida		100.0	0.39	FRU 99			
67767762	**Good identification**						**67767762**
Mitsuokella multacida		100.0	0.39	FRU 99			
67767770	**Good identification**						**67767770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	FRU 99	1	1	
Blautia producta		3.47	0.15		99	99	
67770000	**Low discrimination**						**67770000**
					GRA		
Actinomyces naeslundii		76.32	0.46	ESL 12	99		
Fusobacterium mortiferum		18.79	0.24	GAL 96	1		
Actinomyces israelii		4.89	0.16		99		
67770001	**Good identification**						**67770001**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67770003	**Low discrimination**						**67770003**
Actinomyces naeslundii		79.84	0.49	ESL 12			
Actinomyces viscosus		20.16	0.15	GAL 96			
67770020	**Good identification**						**67770020**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67770021	**Good identification**						**67770021**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67770022	**Good identification**						**67770022**
Actinomyces naeslundii		100.0	0.46	ESL 12			
67770023	**Good identification**						**67770023**
Actinomyces naeslundii		100.0	0.49	ESL 12			
67770041	**Good identification**						**67770041**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67770043	**Good identification**						**67770043**
Actinomyces naeslundii		100.0	0.36	ESL 12			
67770061	**Good identification**						**67770061**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	ESL 12		
67770063	**Good identification**					**67770063**
Actinomyces naeslundii		100.0	0.36	ESL 12		
67770100	**Low discrimination**					**67770100**
					GRA	
Actinomyces naeslundii		77.41	0.54	ESL 12	99	
Fusobacterium mortiferum		15.62	0.30	GAL 96	1	
Actinomyces israelii		6.97	0.27		99	
67770101	**Very good identification**					**67770101**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67770102	**Low discrimination**					**67770102**
					GRA	
Actinomyces naeslundii		43.23	0.54	ESL 12	99	
Actinomyces viscosus		30.68	0.27	GAL 96	99	
Leptotrichia buccalis		8.34	0.07		1	
67770103	**Low discrimination**					**67770103**
Actinomyces naeslundii		86.38	0.57	ESL 12		
Actinomyces viscosus		13.62	0.19	GAL 96		
67770112	**Good identification**					**67770112**
					GRA	
Leptotrichia buccalis		98.81	0.41	NIT 10	1	
Bifidobacterium breve		0.72	0.24	GAL 99	99	
Actinomyces naeslundii		0.27	0.32		99	
67770120	**Low discrimination**					**67770120**
					SPO	
Actinomyces naeslundii		85.48	0.54	ESL 12	1	
Clostridium paraputrificum		6.58	0.03	GAL 96	99	
Clostridium ramosum		4.46	0.06		92	
67770121	**Very good identification**					**67770121**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67770122	**Low discrimination**					**67770122**
					SPO	
Actinomyces naeslundii		57.80	0.54	ESL 12	1	
Bifidobacterium breve		14.26	0.29	GAL 96	1	
Clostridium ramosum		12.07	0.17		92	
67770123	**Very good identification**					**67770123**
Actinomyces naeslundii		100.0	0.57	ESL 12		
67770140	**Low discrimination**					**67770140**
Actinomyces naeslundii		83.45	0.41	ESL 12		
Actinomyces israelii		16.55	0.20	GAL 96		
67770141	**Good identification**					**67770141**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67770142	**Low discrimination**					**67770142**
Actinomyces naeslundii		47.98	0.41	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		37.74	0.24	GAL 96		
Actinomyces israelii		14.28	0.23			
67770143	**Good identification**					**67770143**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67770160	**Low discrimination**					**67770160**
Actinomyces naeslundii		87.85	0.41	ESL 12		
Bifidobacterium breve		12.15	0.12	GAL 96		
67770161	**Good identification**					**67770161**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67770163	**Good identification**					**67770163**
Actinomyces naeslundii		100.0	0.44	ESL 12		
67770712	**Good identification**					**67770712**
Clostridium clostridioforme		99.26	0.41	NIT 1		
Mitsuokella multacida		0.74	0.01	XYL 91		
67770732	**Good identification**					**67770732**
Clostridium clostridioforme		100.0	0.41	NIT 1		
67771000	**Low discrimination**					**67771000**
Actinomyces naeslundii		71.58	0.70	ESL 12		
Actinomyces israelii		18.91	0.50			
Lactobacillus jensenii		8.05	0.32			
67771001	**Very good identification**					**67771001**
Actinomyces naeslundii		99.82	0.73	ESL 12		
Actinomyces israelii		0.18	0.16			
67771002	**Low discrimination**					**67771002**
Actinomyces naeslundii		60.07	0.70	ESL 12		
Actinomyces israelii		23.80	0.53			
Lactobacillus acidophilus		7.01	0.29			
67771003	**Very good identification**					**67771003**
Actinomyces naeslundii		99.10	0.73	ESL 12		
Actinomyces viscosus		0.64	0.11			
Actinomyces israelii		0.26	0.19			
67771010	**Low discrimination**					**67771010**
Actinomyces naeslundii		34.52	0.48	ESL 12		
Actinomyces israelii		25.89	0.36	MLZ 5		
Lactobacillus jensenii		24.59	0.24			
67771011	**Good identification**					**67771011**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67771012	**Acceptable identification**					**67771012**
Leptotrichia buccalis		93.58	0.41	NIT 10	GRA 1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces israelii		2.45	0.39	LAC 99	99	
Actinomyces naeslundii		2.18	0.48		99	
67771013	**Low discrimination**					**67771013**
					GRA	
Actinomyces naeslundii		77.54	0.51	ESL 12	99	
Leptotrichia buccalis		22.46	0.07	MLZ 5	1	
67771020	**Good identification**					**67771020**
Actinomyces naeslundii		91.27	0.70	ESL 12		
Lactobacillus jensenii		2.57	0.21			
Clostridium perfringens		2.38	0.20			
67771021	**Very good identification**					**67771021**
Actinomyces naeslundii		100.0	0.73	ESL 12		
67771022	**Low discrimination**					**67771022**
					GRA	
Actinomyces naeslundii		75.79	0.70	ESL 12	99	
Lactobacillus acidophilus		8.85	0.29		99	
Prevotella oralis		6.01	0.25		1	
67771023	**Very good identification**					**67771023**
Actinomyces naeslundii		100.0	0.73	ESL 12		
67771030	**Low discrimination**					**67771030**
Actinomyces naeslundii		75.24	0.48	ESL 12		
Lactobacillus jensenii		13.39	0.13	MLZ 5		
Bifidobacterium breve		4.14	0.12			
67771031	**Good identification**					**67771031**
Actinomyces naeslundii		100.0	0.51	ESL 12		
67771033	**Good identification**					**67771033**
					GRA	
Actinomyces naeslundii		96.88	0.51	ESL 12	99	
Leptotrichia buccalis		3.12	-.09	MLZ 5	1	
67771040	**Low discrimination**					**67771040**
Actinomyces naeslundii		63.21	0.57	ESL 12		
Actinomyces israelii		36.80	0.43			
67771041	**Very good identification**					**67771041**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67771042	**Low discrimination**					**67771042**
Actinomyces naeslundii		48.82	0.57	ESL 12		
Actinomyces israelii		42.63	0.46			
Bifidobacterium breve		4.55	0.24			
67771043	**Very good identification**					**67771043**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67771051	**Good identification**					**67771051**
Actinomyces naeslundii		100.0	0.38	ESL 12		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67771053	**Good identification**					**67771053**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67771060	**Good identification**					**67771060**
Actinomyces naeslundii		92.45	0.57	ESL 12		
Actinomyces israelii		3.44	0.23			
Clostridium perfringens		2.60	0.08			
67771061	**Very good identification**					**67771061**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67771062	**Low discrimination**					**67771062**
Actinomyces naeslundii		75.95	0.57	ESL 12		
Bifidobacterium breve		12.60	0.29			
Lactobacillus acidophilus		6.21	0.14			
67771063	**Very good identification**					**67771063**
Actinomyces naeslundii		100.0	0.60	ESL 12		
67771071	**Good identification**					**67771071**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67771073	**Good identification**					**67771073**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67771100	**Low discrimination**					**67771100**
Actinomyces naeslundii		64.75	0.77	ESL 12		
Actinomyces israelii		24.04	0.61			
Bifidobacterium breve		3.67	0.41			
67771101	**Very good identification**					**67771101**
					GRA	
Actinomyces naeslundii		99.68	0.80	ESL 12	99	
Actinomyces israelii		0.25	0.27		99	
Mitsuokella multacida		0.07	0.04		1	
67771102	**Low discrimination**					**67771102**
					GRA	
Actinomyces naeslundii		28.58	0.77	ESL 12	99	
Leptotrichia buccalis		22.73	0.41		1	
Bifidobacterium breve		16.37	0.59		99	
67771103	**Very good identification**					**67771103**
					GRA	
Actinomyces naeslundii		98.02	0.80	ESL 12	99	
Leptotrichia buccalis		0.53	0.07		1	
Actinomyces viscosus		0.40	0.15		99	
67771110	**Low discrimination**					**67771110**
					GRA	
Leptotrichia buccalis		82.37	0.41	NIT 10	1	
Bifidobacterium breve		5.87	0.41	BGL 99	99	
Actinomyces israelii		5.75	0.47		99	
67771111	**Good identification**					**67771111**
					GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		90.76	0.58	ESL 12	99	
Leptotrichia buccalis		9.24	0.07	MLZ 5	1	
67771112	**Very good identification**					**67771112**
					GRA	
Leptotrichia buccalis		99.07	0.76	NIT 10	1	
Bifidobacterium breve		0.72	0.59		99	
Actinomyces israelii		0.11	0.50		99	
67771113	**Acceptable identification**					**67771113**
					GRA	
Leptotrichia buccalis		90.29	0.41	NIT 10	1	
Actinomyces naeslundii		8.96	0.58	URE 1	99	
Bifidobacterium breve		0.66	0.24		99	
67771120	**Low discrimination**					**67771120**
					SPO	
Actinomyces naeslundii		56.53	0.77	ESL 12	1	
Clostridium paraputrificum		17.95	0.37		99	
Clostridium ramosum		12.17	0.41		92	
67771121	**Very good identification**					**67771121**
					SPO	
Actinomyces naeslundii		99.53	0.80	ESL 12	1	
Clostridium paraputrificum		0.21	0.03		99	
Clostridium ramosum		0.14	0.06		92	
67771122	**Low discrimination**					**67771122**
					SPO	
Bifidobacterium breve		24.64	0.63	NIT 1	1	
Actinomyces naeslundii		24.21	0.77		1	
Clostridium ramosum		20.85	0.51		92	
67771123	**Very good identification**					**67771123**
					SPO	
Actinomyces naeslundii		98.01	0.80	ESL 12	1	
Bifidobacterium breve		0.67	0.29		1	
Clostridium ramosum		0.57	0.17		92	
67771130	**Low discrimination**					**67771130**
					GRA	
Bifidobacterium breve		39.94	0.46	NIT 1	99	
Leptotrichia buccalis		35.06	0.25	BGL 91	1	
Actinomyces naeslundii		20.88	0.55		99	
67771131	**Very good identification**					**67771131**
					GRA	
Actinomyces naeslundii		97.64	0.58	ESL 12	99	
Bifidobacterium breve		1.26	0.12	MLZ 5	99	
Leptotrichia buccalis		1.10	-.09		1	
67771132	**Low discrimination**					**67771132**
					GRA	
Leptotrichia buccalis		88.72	0.59	NIT 10	1	
Bifidobacterium breve		10.32	0.63	NAG 10	99	
Actinomyces naeslundii		0.53	0.55		99	
67771140	**Low discrimination**					**67771140**
Actinomyces naeslundii		45.16	0.64	ESL 12		
Actinomyces israelii		36.95	0.54			
Bifidobacterium breve		14.49	0.41			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
67771141	**Very good identification**					**67771141**
					GRA	
Actinomyces naeslundii		99.06	0.67	ESL 12	99	
Actinomyces israelii		0.55	0.20		99	
Mitsuokella multacida		0.39	0.04		1	
67771142	**Low discrimination**					**67771142**
Bifidobacterium breve		55.21	0.59	NIT 1		
Actinomyces israelii		20.88	0.57			
Actinomyces naeslundii		17.02	0.64			
67771143	**Very good identification**					**67771143**
Actinomyces naeslundii		97.08	0.67	ESL 12		
Bifidobacterium breve		2.12	0.24			
Actinomyces israelii		0.80	0.23			
67771150	**Low discrimination**					**67771150**
Bifidobacterium breve		57.38	0.41	NIT 1		
Actinomyces israelii		21.86	0.40	BGL 91		
Actinomyces naeslundii		9.41	0.43			
67771151	**Good identification**					**67771151**
Actinomyces naeslundii		100.0	0.46	ESL 12		
67771152	**Low discrimination**					**67771152**
					GRA	
Leptotrichia buccalis		56.30	0.41	NIT 10	1	
Bifidobacterium breve		40.53	0.59	SOR 1	99	
Actinomyces israelii		2.29	0.43		99	
67771153	**Low discrimination**					**67771153**
					GRA	
Actinomyces naeslundii		50.21	0.46	ESL 12	99	
Leptotrichia buccalis		28.95	0.07	MLZ 5	1	
Bifidobacterium breve		20.84	0.24		99	
67771160	**Low discrimination**					**67771160**
					SPO	
Actinomyces naeslundii		53.87	0.64	ESL 12	1	
Bifidobacterium breve		30.73	0.46		1	
Clostridium paraputrificum		7.30	0.18		99	
67771161	**Very good identification**					**67771161**
Actinomyces naeslundii		99.62	0.67	ESL 12		
Bifidobacterium breve		0.38	0.12			
67771162	**Low discrimination**					**67771162**
Bifidobacterium breve		78.38	0.63	NIT 1		
Actinomyces naeslundii		13.59	0.64			
Lactobacillus acidophilus		4.49	0.32			
67771163	**Very good identification**					**67771163**
Actinomyces naeslundii		96.26	0.67	ESL 12		
Bifidobacterium breve		3.74	0.29			
67771170	**Low discrimination**					**67771170**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bifidobacterium breve		89.08	0.46	NIT 1			
Actinomyces naeslundii		8.22	0.43	BGL 91			
Actinomyces israelii		1.22	0.19				
67771171	**Acceptable identification**						**67771171**
Actinomyces naeslundii		93.20	0.46	ESL 12			
Bifidobacterium breve		6.80	0.12	MLZ 5			
67771172	**Good identification**						**67771172**
					GRA		
Bifidobacterium breve		91.08	0.63	NIT 1	99		
Leptotrichia buccalis		7.91	0.25		1		
Actinomyces naeslundii		0.83	0.43		99		
67771173	**Low discrimination**						**67771173**
					GRA		
Actinomyces naeslundii		55.49	0.46	ESL 12	99		
Bifidobacterium breve		40.95	0.29	MLZ 5	99		
Leptotrichia buccalis		3.56	-.09		1		
67771201	**Good identification**						**67771201**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67771203	**Good identification**						**67771203**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67771220	**Low discrimination**						**67771220**
					SPO	GRA	
Actinomyces naeslundii		50.27	0.35	ESL 12	1	99	
Clostridium ramosum		40.84	0.09	RHA 1	92	50	
Prevotella oralis		8.90	-.03		1	1	
67771221	**Good identification**						**67771221**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67771223	**Good identification**						**67771223**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67771300	**Low discrimination**						**67771300**
					GRA		
Actinomyces israelii		46.67	0.41	XYL 92	99		
Mitsuokella multacida		29.41	0.25	RHA 7	1		
Actinomyces naeslundii		16.87	0.43		99		
67771301	**Low discrimination**						**67771301**
					GRA		
Actinomyces naeslundii		88.56	0.46	ESL 12	99		
Mitsuokella multacida		11.44	0.09	RHA 1	1		
67771302	**Low discrimination**						**67771302**
					SPO		
Actinomyces israelii		32.47	0.45	XYL 92	1		
Lactobacillus acidophilus		27.51	0.28	RHA 7	1		
Clostridium ramosum		8.93	0.19		92		
67771303	**Good identification**						**67771303**
Actinomyces naeslundii		100.0	0.46	ESL 12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67771312	**Good identification**						**67771312**
					GRA		
Leptotrichia buccalis		96.44	0.41	NIT 10	1		
Clostridium clostridioforme		2.03	0.28	RHA 1	50		
Actinomyces israelii		0.76	0.30		99		
67771320	**Low discrimination**						**67771320**
					SPO	GRA	
Clostridium ramosum		89.98	0.43	NIT 1	92	50	
Actinomyces naeslundii		3.18	0.43		1	99	
Prevotella oralis		3.10	0.17		1	1	
67771321	**Low discrimination**						**67771321**
					SPO		
Actinomyces naeslundii		84.01	0.46	ESL 12	1		
Clostridium ramosum		15.99	0.09	RHA 1	92		
67771322	**Low discrimination**						**67771322**
					SPO		
Clostridium ramosum		77.82	0.53	NIT 1	92		
Prevotella oralis		12.74	0.39		1		
Parabacteroides distasonis		5.04	0.45		1		
67771323	**Low discrimination**						**67771323**
					SPO	GRA	
Actinomyces naeslundii		51.68	0.46	ESL 12	1	99	
Clostridium ramosum		39.34	0.19	RHA 1	92	50	
Prevotella oralis		6.44	0.05		1	1	
67771342	**Low discrimination**						**67771342**
Actinomyces israelii		53.95	0.37	XYL 92			
Bifidobacterium breve		19.14	0.24	RHA 7			
Lactobacillus acidophilus		14.53	0.12				
67771400	**Low discrimination**						**67771400**
					GRA	SPO	
Actinomyces israelii		72.16	0.49	XYL 92	99	1	
Clostridium butyricum		15.45	0.16		99	99	
Mitsuokella multacida		9.01	0.21		1	1	
67771401	**Low discrimination**						**67771401**
					GRA		
Actinomyces naeslundii		74.51	0.38	ESL 12	99		
Mitsuokella multacida		14.75	0.04	ARA 1	1		
Actinomyces israelii		10.74	0.15		99		
67771402	**Low discrimination**						**67771402**
					SPO		
Actinomyces israelii		88.58	0.52	XYL 92	1		
Clostridium butyricum		4.92	0.09		99		
Actinomyces naeslundii		2.76	0.35		1		
67771403	**Low discrimination**						**67771403**
Actinomyces naeslundii		82.23	0.38	ESL 12			
Actinomyces israelii		17.77	0.18	ARA 1			
67771412	**Low discrimination**						**67771412**
					GRA		
Actinomyces israelii		45.44	0.38	XYL 92	99		
Clostridium clostridioforme		28.45	0.25	MLZ 13	50		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Leptotrichia buccalis		21.45	0.07		1		
67771421	**Good identification**						**67771421**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67771423	**Good identification**						**67771423**
Actinomyces naeslundii		100.0	0.38	ESL 12			
67771440	**Low discrimination**						**67771440**
					GRA	SPO	
Actinomyces israelii		71.93	0.42	XYL 92	99	1	
Mitsuokella multacida		23.10	0.21		1	1	
Clostridium butyricum		3.44	-.02		99	99	
67771442	**Good identification**						**67771442**
					GRA		
Actinomyces israelii		96.35	0.45	XYL 92	99		
Mitsuokella multacida		2.29	0.04		1		
Actinomyces naeslundii		1.36	0.22		99		
67771500	**Low discrimination**						**67771500**
					GRA	SPO	
Mitsuokella multacida		57.09	0.55	XYL 99	1	1	
Clostridium butyricum		23.73	0.40		99	99	
Actinomyces israelii		18.47	0.59		99	1	
67771501	**Low discrimination**						**67771501**
					GRA	SPO	
Mitsuokella multacida		82.48	0.39	XYL 99	1	1	
Actinomyces naeslundii		11.98	0.46	URE 10	99	1	
Clostridium butyricum		3.12	0.06		99	99	
67771502	**Low discrimination**						**67771502**
					SPO		
Actinomyces israelii		51.22	0.62	XYL 92	1		
Clostridium butyricum		17.06	0.33		99		
Bifidobacterium adolescentis		15.88	0.47		1		
67771503	**Low discrimination**						**67771503**
					GRA		
Actinomyces naeslundii		44.17	0.46	ESL 12	99		
Mitsuokella multacida		33.79	0.22	ARA 1	1		
Actinomyces israelii		13.42	0.28		99		
67771510	**Low discrimination**						**67771510**
					GRA	SPO	
Mitsuokella multacida		80.52	0.47	XYL 99	1	1	
Actinomyces israelii		11.68	0.45		99	1	
Clostridium butyricum		5.29	0.18		99	99	
67771512	**Low discrimination**						**67771512**
					GRA		
Leptotrichia buccalis		70.92	0.41	NIT 10	1		
Bifidobacterium adolescentis		15.40	0.49	ARA 1	99		
Actinomyces israelii		6.07	0.48		99		
67771520	**Low discrimination**						**67771520**
					GRA	SPO	
Mitsuokella multacida		76.90	0.45	XYL 99	1	1	
Clostridium butyricum		11.12	0.22		99	99	
Actinomyces israelii		6.35	0.39		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67771522	**Low discrimination**						**67771522**
					GRA		
Actinomyces israelii		21.36	0.42	XYL 92	99		
Mitsuokella multacida		19.15	0.28	NAG 6	1		
Parabacteroides distasonis		11.10	0.33		1		
67771523	**Low discrimination**						**67771523**
					GRA		
Actinomyces naeslundii		83.95	0.46	ESL 12	99		
Mitsuokella multacida		16.06	0.12	ARA 1	1		
67771530	**Acceptable identification**						**67771530**
					GRA	SPO	
Mitsuokella multacida		92.23	0.37	XYL 99	1	1	
Actinomyces israelii		3.42	0.25		99	1	
Clostridium butyricum		2.11	0.00		99	99	
67771540	**Low discrimination**						**67771540**
					GRA	SPO	
Mitsuokella multacida		85.92	0.55	XYL 99	1	1	
Actinomyces israelii		10.81	0.52		99	1	
Clostridium butyricum		3.11	0.22		99	99	
67771541	**Good identification**						**67771541**
					GRA		
Mitsuokella multacida		96.43	0.39	XYL 99	1		
Actinomyces naeslundii		2.47	0.33	URE 10	99		
Actinomyces israelii		1.10	0.18		99		
67771542	**Low discrimination**						**67771542**
					GRA		
Actinomyces israelii		55.42	0.55	XYL 92	99		
Mitsuokella multacida		32.63	0.39		1		
Bifidobacterium adolescentis		5.46	0.32		99		
67771550	**Acceptable identification**						**67771550**
					GRA	SPO	
Mitsuokella multacida		93.84	0.47	XYL 99	1	1	
Actinomyces israelii		5.29	0.38		99	1	
Clostridium butyricum		0.54	0.00		99	99	
67771560	**Acceptable identification**						**67771560**
					GRA	SPO	
Mitsuokella multacida		94.64	0.45	XYL 99	1	1	
Actinomyces israelii		3.04	0.32		99	1	
Clostridium butyricum		1.19	0.04		99	99	
67771570	**Good identification**						**67771570**
					GRA		
Mitsuokella multacida		97.33	0.37	XYL 99	1		
Actinomyces israelii		1.40	0.18		99		
Bifidobacterium breve		1.27	0.12		99		
67771612	**Good identification**						**67771612**
					GRA		
Clostridium clostridioforme		97.59	0.40	NIT 1	50		
Actinomyces israelii		1.60	0.19	XYL 91	99		
Mitsuokella multacida		0.82	0.01		1		
67771632	**Good identification**						**67771632**
Clostridium clostridioforme		100.0	0.40	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67771700	**Very good identification**						**67771700**
					GRA	SPO	
Mitsuokella multacida		98.04	0.60	XYL 99	1	1	
Actinomyces israelii		1.29	0.40		99	1	
Clostridium butyricum		0.68	0.14		99	99	
67771701	**Good identification**						**67771701**
Mitsuokella multacida		100.0	0.43	XYL 99			
67771702	**Low discrimination**						**67771702**
					GRA		
Mitsuokella multacida		66.44	0.43	XYL 99	1		
Clostridium clostridioforme		19.70	0.37	BGL 10	50		
Actinomyces israelii		11.76	0.43		99		
67771710	**Good identification**						**67771710**
					GRA		
Mitsuokella multacida		98.86	0.51	XYL 99	1		
Actinomyces israelii		0.58	0.26		99		
Clostridium clostridioforme		0.45	0.15		50		
67771711	**Good identification**						**67771711**
Mitsuokella multacida		100.0	0.35	XYL 99			
67771712	**Low discrimination**						**67771712**
Clostridium clostridioforme		76.70	0.49	NIT 1			
Mitsuokella multacida		18.93	0.35	XYL 91			
Leptotrichia buccalis		2.36	0.07				
67771720	**Good identification**						**67771720**
					SPO	GRA	
Mitsuokella multacida		98.78	0.49	XYL 99	1	1	
Clostridium ramosum		0.65	0.09		92	50	
Actinomyces israelii		0.33	0.20		1	99	
67771722	**Low discrimination**						**67771722**
Clostridium clostridioforme		33.61	0.37	NIT 1			
Mitsuokella multacida		28.34	0.33	XYL 91			
Parabacteroides distasonis		26.53	0.41				
67771730	**Good identification**						**67771730**
Mitsuokella multacida		98.21	0.41	XYL 99			
Clostridium clostridioforme		1.79	0.15				
67771732	**Low discrimination**						**67771732**
Clostridium clostridioforme		85.70	0.49	NIT 1			
Parabacteroides distasonis		8.00	0.37	XYL 91			
Mitsuokella multacida		5.29	0.25				
67771740	**Very good identification**						**67771740**
					GRA	SPO	
Mitsuokella multacida		99.43	0.60	XYL 99	1	1	
Actinomyces israelii		0.51	0.33		99	1	
Clostridium butyricum		0.06	-.04		99	99	
67771741	**Good identification**						**67771741**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.43	XYL 99			
67771742	**Acceptable identification**						**67771742**
					GRA		
Mitsuokella multacida		93.56	0.43	XYL 99	1		
Actinomyces israelii		6.44	0.36	BGL 10	99		
67771750	**Good identification**						**67771750**
					COC	GRA	
Mitsuokella multacida		99.58	0.51	XYL 99	1	1	
Actinomyces israelii		0.23	0.19		1	99	
Blautia producta		0.19	-.03		99	99	
67771751	**Good identification**						**67771751**
Mitsuokella multacida		100.0	0.35	XYL 99			
67771752	**Acceptable identification**						**67771752**
					GRA		
Mitsuokella multacida		93.30	0.35	XYL 99	1		
Clostridium clostridioforme		3.82	0.15	BGL 10	50		
Actinomyces israelii		2.88	0.22		99		
67771760	**Good identification**						**67771760**
Mitsuokella multacida		100.0	0.49	XYL 99			
67771770	**Good identification**						**67771770**
Mitsuokella multacida		100.0	0.41	XYL 99			
67772000	**Low discrimination**						**67772000**
Actinomyces naeslundii		84.25	0.51	ESL 12			
Actinomyces israelii		15.75	0.29	GAL 96			
67772001	**Good identification**						**67772001**
Actinomyces naeslundii		100.0	0.54	ESL 12			
67772002	**Low discrimination**						**67772002**
Actinomyces naeslundii		78.10	0.51	ESL 12			
Actinomyces israelii		21.90	0.32	GAL 96			
67772003	**Good identification**						**67772003**
Actinomyces naeslundii		100.0	0.54	ESL 12			
67772020	**Good identification**						**67772020**
Actinomyces naeslundii		100.0	0.51	ESL 12			
67772021	**Good identification**						**67772021**
Actinomyces naeslundii		100.0	0.54	ESL 12			
67772022	**Good identification**						**67772022**
Actinomyces naeslundii		100.0	0.51	ESL 12			
67772023	**Good identification**						**67772023**
Actinomyces naeslundii		100.0	0.54	ESL 12			
67772040	**Low discrimination**						**67772040**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		70.82	0.38	ESL 12		
Actinomyces israelii		29.18	0.22	GAL 96		
67772041	**Good identification**					**67772041**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67772042	**Low discrimination**					**67772042**
Actinomyces naeslundii		61.81	0.38	ESL 12		
Actinomyces israelii		38.20	0.25	GAL 96		
67772043	**Good identification**					**67772043**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67772060	**Good identification**					**67772060**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67772061	**Good identification**					**67772061**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67772062	**Good identification**					**67772062**
Actinomyces naeslundii		100.0	0.38	ESL 12		
67772063	**Good identification**					**67772063**
Actinomyces naeslundii		100.0	0.41	ESL 12		
67772100	**Low discrimination**					**67772100**
Actinomyces naeslundii		79.19	0.59	ESL 12		
Actinomyces israelii		20.81	0.39	GAL 96		
67772101	**Very good identification**					**67772101**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67772102	**Low discrimination**					**67772102**
Actinomyces naeslundii		68.23	0.59	ESL 12		
Actinomyces israelii		26.89	0.42	GAL 96		
Bifidobacterium breve		4.88	0.24			
67772103	**Very good identification**					**67772103**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67772110	**Low discrimination**					**67772110**
Actinomyces naeslundii		57.28	0.37	ESL 12		
Actinomyces israelii		42.72	0.25	GAL 96		
67772111	**Good identification**					**67772111**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67772113	**Good identification**					**67772113**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67772120	**Low discrimination**					**67772120**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					SPO	
Actinomyces naeslundii		84.62	0.59	ESL 12	1	
Clostridium ramosum		12.90	0.19	GAL 96	92	
Actinomyces israelii		1.42	0.19		1	
67772121	**Very good identification**					**67772121**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67772122	**Low discrimination**					**67772122**
					SPO	
Actinomyces naeslundii		56.76	0.59	ESL 12	1	
Clostridium ramosum		34.60	0.30	GAL 96	92	
Bifidobacterium breve		7.22	0.29		1	
67772123	**Very good identification**					**67772123**
Actinomyces naeslundii		100.0	0.62	ESL 12		
67772130	**Low discrimination**					**67772130**
Actinomyces naeslundii		80.72	0.37	ESL 12		
Bifidobacterium breve		19.28	0.12	GAL 96		
67772131	**Good identification**					**67772131**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67772133	**Good identification**					**67772133**
Actinomyces naeslundii		100.0	0.40	ESL 12		
67772140	**Low discrimination**					**67772140**
Actinomyces naeslundii		63.33	0.46	ESL 12		
Actinomyces israelii		36.67	0.32	GAL 96		
67772141	**Good identification**					**67772141**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67772142	**Low discrimination**					**67772142**
Actinomyces naeslundii		43.98	0.46	ESL 12		
Actinomyces israelii		38.20	0.35	GAL 96		
Bifidobacterium breve		17.82	0.24			
67772143	**Good identification**					**67772143**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67772160	**Acceptable identification**					**67772160**
Actinomyces naeslundii		93.35	0.46	ESL 12		
Bifidobacterium breve		6.65	0.12	GAL 96		
67772161	**Good identification**					**67772161**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67772162	**Low discrimination**					**67772162**
Actinomyces naeslundii		56.31	0.46	ESL 12		
Bifidobacterium breve		40.57	0.29	GAL 96		
Actinomyces israelii		3.12	0.15			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67772163	**Good identification**					**67772163**
Actinomyces naeslundii		100.0	0.49	ESL 12		
67772502	**Low discrimination**					**67772502**
					GRA	
Actinomyces israelii		85.44	0.41	GAL 99	99	
Mitsuokella multacida		8.06	0.11	XYL 92	1	
Bifidobacterium adolescentis		3.83	0.12		99	
67773000	**Low discrimination**					**67773000**
Actinomyces naeslundii		56.46	0.74	ESL 12		
Actinomyces israelii		43.54	0.63			
67773001	**Very good identification**					**67773001**
Actinomyces naeslundii		99.48	0.77	ESL 12		
Actinomyces israelii		0.52	0.29			
67773002	**Low discrimination**					**67773002**
Actinomyces israelii		53.35	0.66	XYL 92		
Actinomyces naeslundii		46.12	0.74			
Bifidobacterium breve		0.39	0.24			
67773003	**Very good identification**					**67773003**
Actinomyces naeslundii		99.23	0.77	ESL 12		
Actinomyces israelii		0.77	0.32			
67773010	**Low discrimination**					**67773010**
Actinomyces israelii		68.65	0.49	XYL 92		
Actinomyces naeslundii		31.35	0.53	MLZ 13		
67773011	**Good identification**					**67773011**
Actinomyces naeslundii		98.55	0.56	ESL 12		
Actinomyces israelii		1.45	0.15	MLZ 5		
67773012	**Low discrimination**					**67773012**
					GRA	
Actinomyces israelii		70.34	0.52	XYL 92	99	
Actinomyces naeslundii		21.42	0.53	MLZ 13	99	
Leptotrichia buccalis		4.79	0.07		1	
67773013	**Good identification**					**67773013**
Actinomyces naeslundii		97.84	0.56	ESL 12		
Actinomyces israelii		2.16	0.18	MLZ 5		
67773020	**Good identification**					**67773020**
					SPO	
Actinomyces naeslundii		93.56	0.74	ESL 12	1	
Actinomyces israelii		4.61	0.43		1	
Clostridium ramosum		1.69	0.19		92	
67773021	**Very good identification**					**67773021**
Actinomyces naeslundii		100.0	0.77	ESL 12		
67773022	**Low discrimination**					**67773022**
					SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		85.88	0.74	ESL 12	1	
Actinomyces israelii		6.34	0.46		1	
Clostridium ramosum		6.21	0.30		92	
67773023	**Very good identification**					**67773023**
Actinomyces naeslundii		100.0	0.77	ESL 12		
67773030	**Low discrimination**					**67773030**
Actinomyces naeslundii		85.61	0.53	ESL 12		
Actinomyces israelii		11.96	0.29	MLZ 5		
Bifidobacterium breve		2.43	0.12			
67773031	**Good identification**					**67773031**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67773032	**Low discrimination**					**67773032**
Actinomyces naeslundii		65.75	0.53	ESL 12		
Bifidobacterium breve		18.83	0.29	MLZ 5		
Actinomyces israelii		13.78	0.32			
67773033	**Good identification**					**67773033**
Actinomyces naeslundii		100.0	0.56	ESL 12		
67773040	**Low discrimination**					**67773040**
Actinomyces israelii		62.96	0.56	XYL 92		
Actinomyces naeslundii		37.05	0.62			
67773041	**Very good identification**					**67773041**
Actinomyces naeslundii		98.87	0.65	ESL 12		
Actinomyces israelii		1.13	0.22			
67773042	**Low discrimination**					**67773042**
Actinomyces israelii		70.86	0.59	XYL 92		
Actinomyces naeslundii		27.80	0.62			
Bifidobacterium breve		1.34	0.24			
67773043	**Very good identification**					**67773043**
Actinomyces naeslundii		98.31	0.65	ESL 12		
Actinomyces israelii		1.69	0.25			
67773050	**Low discrimination**					**67773050**
Actinomyces israelii		82.83	0.42	XYL 92		
Actinomyces naeslundii		17.17	0.40	MLZ 13		
67773051	**Good identification**					**67773051**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67773052	**Low discrimination**					**67773052**
Actinomyces israelii		79.09	0.45	XYL 92		
Actinomyces naeslundii		10.93	0.40	MLZ 13		
Bifidobacterium breve		9.98	0.24			
67773053	**Good identification**					**67773053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.43	ESL 12		
67773060	**Low discrimination**					**67773060**
Actinomyces naeslundii		89.53	0.62	ESL 12		
Actinomyces israelii		9.71	0.36			
Bifidobacterium breve		0.76	0.12			
67773061	**Very good identification**					**67773061**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67773062	**Low discrimination**					**67773062**
Actinomyces naeslundii		80.12	0.62	ESL 12		
Actinomyces israelii		13.04	0.39			
Bifidobacterium breve		6.85	0.29			
67773063	**Very good identification**					**67773063**
Actinomyces naeslundii		100.0	0.65	ESL 12		
67773070	**Low discrimination**					**67773070**
Actinomyces naeslundii		68.10	0.40	ESL 12		
Actinomyces israelii		20.97	0.22	MLZ 5		
Bifidobacterium breve		10.93	0.12			
67773071	**Good identification**					**67773071**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67773073	**Good identification**					**67773073**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67773100	**Low discrimination**					**67773100**
Actinomyces israelii		50.78	0.74	XYL 92		
Actinomyces naeslundii		46.85	0.82			
Bifidobacterium breve		1.37	0.41			
67773101	**Very good identification**					**67773101**
					GRA	
Actinomyces naeslundii		99.19	0.85	ESL 12	99	
Actinomyces israelii		0.72	0.39		99	
Mitsuokella multacida		0.08	0.11		1	
67773102	**Low discrimination**					**67773102**
Actinomyces israelii		54.60	0.77	XYL 92		
Actinomyces naeslundii		33.58	0.82			
Bifidobacterium breve		9.91	0.59			
67773103	**Very good identification**					**67773103**
Actinomyces naeslundii		98.72	0.85	ESL 12		
Actinomyces israelii		1.08	0.42			
Bifidobacterium breve		0.20	0.24			
67773110	**Low discrimination**					**67773110**
Actinomyces israelii		63.80	0.59	XYL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		20.73	0.60	MLZ 13		
Bifidobacterium breve		11.49	0.41			
67773111	**Very good identification**					**67773111**
					GRA	
Actinomyces naeslundii		97.46	0.63	ESL 12	99	
Actinomyces israelii		2.02	0.25	MLZ 5	99	
Mitsuokella multacida		0.52	0.03		1	
67773112	**Low discrimination**					**67773112**
					GRA	
Leptotrichia buccalis		40.22	0.41	NIT 10	1	
Bifidobacterium breve		28.96	0.59	MAN 1	99	
Actinomyces israelii		23.85	0.62		99	
67773113	**Low discrimination**					**67773113**
					GRA	
Actinomyces naeslundii		89.19	0.63	ESL 12	99	
Leptotrichia buccalis		4.68	0.07	MLZ 5	1	
Bifidobacterium breve		3.37	0.24		99	
67773120	**Low discrimination**					**67773120**
					SPO	
Actinomyces naeslundii		56.88	0.82	ESL 12	1	
Clostridium ramosum		35.76	0.53		92	
Actinomyces israelii		3.94	0.53		1	
67773121	**Very good identification**					**67773121**
					SPO	
Actinomyces naeslundii		99.50	0.85	ESL 12	1	
Clostridium ramosum		0.42	0.19		92	
Actinomyces israelii		0.05	0.19		1	
67773122	**Low discrimination**					**67773122**
					SPO	
Clostridium ramosum		60.30	0.64	NIT 1	92	
Actinomyces naeslundii		23.98	0.82		1	
Bifidobacterium breve		12.58	0.63		1	
67773123	**Very good identification**					**67773123**
					SPO	
Actinomyces naeslundii		97.93	0.85	ESL 12	1	
Clostridium ramosum		1.66	0.30		92	
Bifidobacterium breve		0.35	0.29		1	
67773130	**Low discrimination**					**67773130**
Actinomyces naeslundii		40.79	0.60	ESL 12		
Bifidobacterium breve		40.19	0.46	MLZ 5		
Actinomyces israelii		8.01	0.39			
67773131	**Very good identification**					**67773131**
Actinomyces naeslundii		99.34	0.63	ESL 12		
Bifidobacterium breve		0.66	0.12	MLZ 5		
67773132	**Low discrimination**					**67773132**
					GRA	
Bifidobacterium breve		76.69	0.63	NIT 1	99	
Actinomyces naeslundii		7.70	0.60		99	
Leptotrichia buccalis		6.66	0.25		1	
67773133	**Good identification**					**67773133**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					GRA	
Actinomyces naeslundii		93.20	0.63	ESL 12	99	
Bifidobacterium breve		6.25	0.29	MLZ 5	99	
Leptotrichia buccalis		0.54	-.09		1	
67773140	**Low discrimination**					**67773140**
Actinomyces israelii		66.01	0.67	XYL 92		
Actinomyces naeslundii		27.64	0.69			
Bifidobacterium breve		4.57	0.41			
67773141	**Very good identification**					**67773141**
					GRA	
Actinomyces naeslundii		97.96	0.72	ESL 12	99	
Actinomyces israelii		1.58	0.32		99	
Mitsuokella multacida		0.47	0.11		1	
67773142	**Low discrimination**					**67773142**
Actinomyces israelii		57.08	0.70	XYL 92		
Bifidobacterium breve		26.63	0.59			
Actinomyces naeslundii		15.93	0.69			
67773143	**Very good identification**					**67773143**
Actinomyces naeslundii		96.58	0.72	ESL 12		
Actinomyces israelii		2.33	0.35			
Bifidobacterium breve		1.09	0.24			
67773150	**Low discrimination**					**67773150**
Actinomyces israelii		59.52	0.52	XYL 92		
Bifidobacterium breve		27.57	0.41	MLZ 13		
Actinomyces naeslundii		8.78	0.47			
67773151	**Acceptable identification**					**67773151**
					GRA	
Actinomyces naeslundii		92.94	0.50	ESL 12	99	
Actinomyces israelii		4.24	0.18	MLZ 5	99	
Mitsuokella multacida		2.82	0.03		1	
67773152	**Low discrimination**					**67773152**
Bifidobacterium breve		72.90	0.59	NIT 1		
Actinomyces israelii		23.35	0.55			
Actinomyces naeslundii		2.30	0.47			
67773153	**Low discrimination**					**67773153**
Actinomyces naeslundii		77.98	0.50	ESL 12		
Bifidobacterium breve		16.68	0.24	MLZ 5		
Actinomyces israelii		5.34	0.21			
67773160	**Low discrimination**					**67773160**
Actinomyces naeslundii		66.74	0.69	ESL 12		
Bifidobacterium breve		19.61	0.46			
Actinomyces israelii		10.17	0.46			
67773161	**Very good identification**					**67773161**
Actinomyces naeslundii		99.80	0.72	ESL 12		

Code / Taxon		ID score	T index	tests against	Differentiation tests	Code
Bifidobacterium breve		0.20	0.12			
67773162	**Low discrimination**					**67773162**
Bifidobacterium breve		68.26	0.63	NIT 1		
Actinomyces naeslundii		22.97	0.69			
Actinomyces israelii		5.25	0.49			
67773163	**Very good identification**					**67773163**
Actinomyces naeslundii		97.89	0.72	ESL 12		
Bifidobacterium breve		1.96	0.29			
Actinomyces israelii		0.15	0.15			
67773170	**Low discrimination**					**67773170**
Bifidobacterium breve		78.44	0.46	NIT 1		
Actinomyces naeslundii		14.05	0.47	BGL 91		
Actinomyces israelii		6.08	0.32			
67773171	**Good identification**					**67773171**
Actinomyces naeslundii		96.38	0.50	ESL 12		
Bifidobacterium breve		3.62	0.12	MLZ 5		
67773172	**Very good identification**					**67773172**
Bifidobacterium breve		97.08	0.63	NIT 1		
Actinomyces naeslundii		1.72	0.47			
Actinomyces israelii		1.12	0.35			
67773173	**Low discrimination**					**67773173**
Actinomyces naeslundii		72.45	0.50	ESL 12		
Bifidobacterium breve		27.55	0.29	MLZ 5		
67773200	**Low discrimination**					**67773200**
Actinomyces israelii		85.18	0.44	XYL 92		
Actinomyces naeslundii		14.82	0.40	RHA 7		
67773201	**Good identification**					**67773201**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67773202	**Low discrimination**					**67773202**
					SPO	
Actinomyces israelii		88.72	0.47	XYL 92	1	
Actinomyces naeslundii		10.29	0.40	RHA 7	1	
Clostridium ramosum		0.99	-.03		92	
67773203	**Good identification**					**67773203**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67773221	**Good identification**					**67773221**
Actinomyces naeslundii		100.0	0.43	ESL 12		
67773223	**Acceptable identification**					**67773223**
					SPO	
Actinomyces naeslundii		94.00	0.43	ESL 12	1	
Clostridium ramosum		6.01	-.03	RHA 1	92	
67773240	**Acceptable identification**					**67773240**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		92.68	0.37	XYL 92			
Actinomyces naeslundii		7.32	0.27	RHA 7			
67773242	**Acceptable identification**						**67773242**
Actinomyces israelii		95.00	0.40	XYL 92			
Actinomyces naeslundii		5.00	0.27	RHA 7			
67773300	**Low discrimination**						**67773300**
					GRA		
Actinomyces israelii		67.27	0.54	XYL 92	99		
Mitsuokella multacida		17.46	0.32	RHA 7	1		
Actinomyces naeslundii		8.33	0.48		99		
67773301	**Low discrimination**						**67773301**
					GRA		
Actinomyces naeslundii		82.67	0.51	ESL 12	99		
Mitsuokella multacida		12.84	0.15	RHA 1	1		
Actinomyces israelii		4.50	0.20		99		
67773302	**Low discrimination**						**67773302**
					SPO		
Actinomyces israelii		70.97	0.57	XYL 92	1		
Clostridium ramosum		19.53	0.32	RHA 7	92		
Actinomyces naeslundii		5.86	0.48		1		
67773303	**Acceptable identification**						**67773303**
					SPO		
Actinomyces naeslundii		90.58	0.51	ESL 12	1		
Actinomyces israelii		7.39	0.23	RHA 1	1		
Clostridium ramosum		2.03	-.03		92		
67773310	**Low discrimination**						**67773310**
					GRA		
Actinomyces israelii		61.63	0.40	XYL 92	99		
Mitsuokella multacida		35.68	0.24	RHA 7	1		
Actinomyces naeslundii		2.69	0.26	MLZ 13	99		
67773312	**Low discrimination**						**67773312**
					GRA		
Actinomyces israelii		67.57	0.43	XYL 92	99		
Leptotrichia buccalis		15.30	0.07	RHA 7	1		
Bifidobacterium breve		11.01	0.24	MLZ 13	99		
67773320	**Good identification**						**67773320**
					SPO	GRA	
Clostridium ramosum		97.53	0.56	NIT 1	92	50	
Actinomyces naeslundii		1.18	0.48		1	99	
Mitsuokella multacida		0.62	0.21		1	1	
67773321	**Low discrimination**						**67773321**
					SPO	GRA	
Actinomyces naeslundii		62.72	0.51	ESL 12	1	99	
Clostridium ramosum		34.84	0.21	RHA 1	92	50	
Mitsuokella multacida		2.43	0.05		1	1	
67773322	**Very good identification**						**67773322**
					SPO		
Clostridium ramosum		99.22	0.66	NIT 1	92		
Actinomyces naeslundii		0.30	0.48		1		
Actinomyces israelii		0.23	0.37		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
67773340	**Low discrimination**					**67773340**
					GRA	
Actinomyces israelii		58.02	0.47	XYL 92	99	
Mitsuokella multacida		38.72	0.32	RHA 7	1	
Actinomyces naeslundii		3.26	0.35		99	
67773341	**Low discrimination**					**67773341**
					GRA	
Actinomyces naeslundii		50.01	0.38	ESL 12	99	
Mitsuokella multacida		44.00	0.15	RHA 1	1	
Actinomyces israelii		5.99	0.13		99	
67773342	**Low discrimination**					**67773342**
					GRA	
Actinomyces israelii		86.46	0.50	XYL 92	99	
Bifidobacterium breve		5.41	0.24	RHA 7	99	
Mitsuokella multacida		4.27	0.15		1	
67773343	**Low discrimination**					**67773343**
Actinomyces naeslundii		84.76	0.38	ESL 12		
Actinomyces israelii		15.24	0.16	RHA 1		
67773352	**Low discrimination**					**67773352**
					GRA	
Actinomyces israelii		65.39	0.36	XYL 92	99	
Bifidobacterium breve		27.40	0.24	RHA 7	99	
Mitsuokella multacida		7.21	0.07	MLZ 13	1	
67773361	**Low discrimination**					**67773361**
					GRA	
Actinomyces naeslundii		81.97	0.38	ESL 12	99	
Mitsuokella multacida		18.03	0.05	RHA 1	1	
67773363	**Low discrimination**					**67773363**
					SPO	
Actinomyces naeslundii		88.72	0.38	ESL 12	1	
Clostridium ramosum		11.28	-.03	RHA 1	92	
67773400	**Good identification**					**67773400**
					GRA	
Actinomyces israelii		93.32	0.62	XYL 92	99	
Mitsuokella multacida		4.80	0.27		1	
Actinomyces naeslundii		1.49	0.40		99	
67773401	**Low discrimination**					**67773401**
					GRA	
Actinomyces naeslundii		60.29	0.43	ESL 12	99	
Actinomyces israelii		25.36	0.28	ARA 1	99	
Mitsuokella multacida		14.35	0.11		1	
67773402	**Very good identification**					**67773402**
					GRA	
Actinomyces israelii		98.29	0.65	XYL 92	99	
Actinomyces naeslundii		1.05	0.40		99	
Mitsuokella multacida		0.37	0.11		1	
67773403	**Low discrimination**					**67773403**
Actinomyces naeslundii		61.31	0.43	ESL 12		
Actinomyces israelii		38.69	0.31	ARA 1		
67773410	**Low discrimination**					**67773410**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		89.71	0.48	XYL 92	99		
Mitsuokella multacida		10.29	0.19	MLZ 13	1		
67773412	**Good identification**						**67773412**
Actinomyces israelii		96.63	0.51	XYL 92			
Bifidobacterium adolescentis		1.43	0.13	MLZ 13			
Bifidobacterium dentium		1.12	-.07				
67773420	**Low discrimination**						**67773420**
					GRA		
Actinomyces israelii		68.86	0.41	XYL 92	99		
Actinomyces naeslundii		17.27	0.40	NAG 6	99		
Mitsuokella multacida		13.87	0.17		1		
67773421	**Good identification**						**67773421**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67773422	**Low discrimination**						**67773422**
Actinomyces israelii		85.67	0.44	XYL 92			
Actinomyces naeslundii		14.33	0.40	NAG 6			
67773423	**Good identification**						**67773423**
Actinomyces naeslundii		100.0	0.43	ESL 12			
67773440	**Low discrimination**						**67773440**
					GRA		
Actinomyces israelii		87.76	0.55	XYL 92	99		
Mitsuokella multacida		11.60	0.27		1		
Actinomyces naeslundii		0.64	0.27		99		
67773442	**Very good identification**						**67773442**
					GRA		
Actinomyces israelii		98.56	0.58	XYL 92	99		
Mitsuokella multacida		0.97	0.11		1		
Actinomyces naeslundii		0.48	0.27		99		
67773450	**Low discrimination**						**67773450**
					COC	GRA	
Actinomyces israelii		74.67	0.41	XYL 92	1	99	
Mitsuokella multacida		22.03	0.19	MLZ 13	1	1	
Blautia producta		3.30	-.03		99	99	
67773452	**Good identification**						**67773452**
					GRA		
Actinomyces israelii		97.86	0.44	XYL 92	99		
Mitsuokella multacida		2.14	0.03	MLZ 13	1		
67773462	**Acceptable identification**						**67773462**
Actinomyces israelii		92.95	0.37	XYL 92			
Actinomyces naeslundii		7.05	0.27	NAG 6			
67773500	**Low discrimination**						**67773500**
					GRA	SPO	
Mitsuokella multacida		55.10	0.61	XYL 99	1	1	
Actinomyces israelii		43.29	0.72		99	1	
Clostridium butyricum		1.09	0.24		99	99	
67773501	**Low discrimination**						**67773501**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Mitsuokella multacida		83.87	0.45	XYL 99	1		
Actinomyces naeslundii		10.14	0.51	URE 10	99		
Actinomyces israelii		5.99	0.38		99		
67773502	**Low discrimination**						**67773502**
					GRA		
Actinomyces israelii		86.52	0.75	XYL 92	99		
Mitsuokella multacida		8.16	0.45		1		
Bifidobacterium adolescentis		3.87	0.46		99		
67773503	**Low discrimination**						**67773503**
					GRA		
Actinomyces naeslundii		35.14	0.51	ESL 12	99		
Mitsuokella multacida		32.31	0.29	ARA 1	1		
Actinomyces israelii		31.16	0.41		99		
67773510	**Low discrimination**						**67773510**
					GRA		
Mitsuokella multacida		72.71	0.53	XYL 99	1		
Actinomyces israelii		25.61	0.58		99		
Bifidobacterium dentium		1.09	0.10		99		
67773511	**Good identification**						**67773511**
					GRA		
Mitsuokella multacida		95.14	0.37	XYL 99	1		
Actinomyces israelii		3.05	0.24	URE 10	99		
Actinomyces naeslundii		1.82	0.29		99		
67773512	**Low discrimination**						**67773512**
Actinomyces israelii		52.49	0.61	XYL 92			
Bifidobacterium adolescentis		19.22	0.47	MLZ 13			
Bifidobacterium dentium		15.12	0.27				
67773520	**Low discrimination**						**67773520**
					GRA		
Mitsuokella multacida		78.89	0.51	XYL 99	1		
Actinomyces israelii		15.83	0.52		99		
Actinomyces naeslundii		2.82	0.48		99		
67773522	**Low discrimination**						**67773522**
					SPO	GRA	
Actinomyces israelii		53.25	0.55	XYL 92	1	99	
Mitsuokella multacida		19.66	0.35	NAG 6	1	1	
Clostridium ramosum		15.93	0.30		92	50	
67773523	**Low discrimination**						**67773523**
					GRA		
Actinomyces naeslundii		77.73	0.51	ESL 12	99		
Mitsuokella multacida		17.87	0.18	ARA 1	1		
Actinomyces israelii		4.40	0.21		99		
67773530	**Low discrimination**						**67773530**
					GRA		
Mitsuokella multacida		82.77	0.43	XYL 99	1		
Bifidobacterium dentium		8.86	0.15		99		
Actinomyces israelii		7.44	0.38		99		
67773540	**Low discrimination**						**67773540**
					COC	GRA	
Mitsuokella multacida		75.44	0.61	XYL 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		23.05	0.65		1	99	
Blautia producta		1.26	0.24		99	99	
67773541	**Good identification**						**67773541**
					GRA		
Mitsuokella multacida		95.32	0.45	XYL 99	1		
Actinomyces israelii		2.65	0.31	URE 10	99		
Actinomyces naeslundii		2.03	0.38		99		
67773542	**Low discrimination**						**67773542**
					GRA		
Actinomyces israelii		78.66	0.68	XYL 92	99		
Mitsuokella multacida		19.07	0.45		1		
Bifidobacterium adolescentis		1.12	0.30		99		
67773550	**Low discrimination**						**67773550**
					COC	GRA	
Mitsuokella multacida		77.70	0.53	XYL 99	1	1	
Blautia producta		11.66	0.32		99	99	
Actinomyces israelii		10.64	0.51		1	99	
67773551	**Good identification**						**67773551**
					COC	GRA	
Mitsuokella multacida		97.46	0.37	XYL 99	1	1	
Blautia producta		1.33	-.03	URE 10	99	99	
Actinomyces israelii		1.21	0.17		1	99	
67773552	**Low discrimination**						**67773552**
					GRA		
Actinomyces israelii		56.04	0.54	XYL 92	99		
Mitsuokella multacida		30.31	0.37	MLZ 13	1		
Bifidobacterium adolescentis		6.52	0.32		99		
67773560	**Acceptable identification**						**67773560**
					COC	GRA	
Mitsuokella multacida		91.44	0.51	XYL 99	1	1	
Actinomyces israelii		7.13	0.45		1	99	
Blautia producta		0.68	0.07		99	99	
67773562	**Low discrimination**						**67773562**
					GRA		
Actinomyces israelii		46.21	0.48	XYL 92	99		
Mitsuokella multacida		43.88	0.35	NAG 6	1		
Bifidobacterium breve		7.41	0.29		99		
67773570	**Acceptable identification**						**67773570**
					COC	GRA	
Mitsuokella multacida		90.32	0.43	XYL 99	1	1	
Blautia producta		6.02	0.15		99	99	
Actinomyces israelii		3.16	0.31		1	99	
67773602	**Acceptable identification**						**67773602**
					GRA		
Actinomyces israelii		91.41	0.46	XYL 92	99		
Mitsuokella multacida		8.59	0.15	RHA 7	1		
67773642	**Low discrimination**						**67773642**
					GRA		
Actinomyces israelii		80.54	0.39	XYL 92	99		
Mitsuokella multacida		19.46	0.15	RHA 7	1		
67773700	**Very good identification**						**67773700**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		96.88	0.66	XYL 99	1	1	
Actinomyces israelii		3.09	0.53		99	1	
Clostridium butyricum		0.03	-.02		99	99	
67773701	**Good identification**						**67773701**
					GRA		
Mitsuokella multacida		99.71	0.50	XYL 99	1		
Actinomyces israelii		0.29	0.19	URE 10	99		
67773702	**Low discrimination**						**67773702**
					GRA		
Mitsuokella multacida		69.74	0.50	XYL 99	1		
Actinomyces israelii		29.98	0.56	BGL 10	99		
Bifidobacterium adolescentis		0.18	0.12		99		
67773710	**Very good identification**						**67773710**
					COC	GRA	
Mitsuokella multacida		98.51	0.58	XYL 99	1	1	
Actinomyces israelii		1.41	0.39		1	99	
Blautia producta		0.08	-.03		99	99	
67773711	**Good identification**						**67773711**
Mitsuokella multacida		100.0	0.41	XYL 99			
67773712	**Low discrimination**						**67773712**
					GRA		
Mitsuokella multacida		81.49	0.41	XYL 99	1		
Actinomyces israelii		15.70	0.42	BGL 10	99		
Clostridium clostridioforme		1.43	0.15		50		
67773720	**Good identification**						**67773720**
					SPO	GRA	
Mitsuokella multacida		97.64	0.56	XYL 99	1	1	
Clostridium ramosum		1.57	0.21		92	50	
Actinomyces israelii		0.79	0.32		1	99	
67773721	**Good identification**						**67773721**
Mitsuokella multacida		100.0	0.39	XYL 99			
67773722	**Low discrimination**						**67773722**
					SPO	GRA	
Mitsuokella multacida		58.90	0.39	XYL 99	1	1	
Clostridium ramosum		34.06	0.32	BGL 10	92	50	
Actinomyces israelii		6.47	0.35		1	99	
67773730	**Good identification**						**67773730**
					GRA		
Mitsuokella multacida		99.64	0.47	XYL 99	1		
Actinomyces israelii		0.36	0.18		99		
67773740	**Very good identification**						**67773740**
					COC	GRA	
Mitsuokella multacida		97.91	0.66	XYL 99	1	1	
Actinomyces israelii		1.21	0.46		1	99	
Blautia producta		0.88	0.24		99	99	
67773741	**Good identification**						**67773741**
Mitsuokella multacida		100.0	0.50	XYL 99			
67773742	**Low discrimination**						**67773742**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		85.02	0.50	XYL 99	1	1	
Actinomyces israelii		14.22	0.49	BGL 10	1	99	
Blautia producta		0.76	0.07		99	99	
67773750	**Good identification**						**67773750**
					COC	GRA	
Mitsuokella multacida		92.05	0.58	XYL 99	1	1	
Blautia producta		7.44	0.32		99	99	
Actinomyces israelii		0.51	0.32		1	99	
67773751	**Good identification**						**67773751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	XYL 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67773752	**Low discrimination**						**67773752**
					COC	GRA	
Mitsuokella multacida		86.53	0.41	XYL 99	1	1	
Blautia producta		6.99	0.15	BGL 10	99	99	
Actinomyces israelii		6.49	0.35		1	99	
67773760	**Good identification**						**67773760**
					COC	GRA	
Mitsuokella multacida		99.29	0.56	XYL 99	1	1	
Blautia producta		0.40	0.07		99	99	
Actinomyces israelii		0.31	0.25		1	99	
67773761	**Good identification**						**67773761**
Mitsuokella multacida		100.0	0.39	XYL 99			
67773762	**Good identification**						**67773762**
					SPO	GRA	
Mitsuokella multacida		95.37	0.39	XYL 99	1	1	
Actinomyces israelii		4.07	0.28	BGL 10	1	99	
Clostridium ramosum		0.56	-.03		92	50	
67773770	**Good identification**						**67773770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	XYL 99	1	1	
Blautia producta		3.47	0.15		99	99	
67774002	**Good identification**						**67774002**
Actinomyces israelii		100.0	0.37	GAL 99			
67774100	**Low discrimination**						**67774100**
					GRA	SPO	
Actinomyces israelii		63.98	0.45	GAL 99	99	1	
Clostridium butyricum		21.01	0.15		99	99	
Mitsuokella multacida		14.07	0.21		1	1	
67774102	**Low discrimination**						**67774102**
					SPO		
Actinomyces israelii		86.99	0.48	GAL 99	1		
Clostridium butyricum		7.41	0.08		99		
Clostridium clostridioforme		2.27	0.11		66		
67774140	**Low discrimination**						**67774140**
					GRA	SPO	
Actinomyces israelii		61.02	0.38	GAL 99	99	1	
Mitsuokella multacida		34.50	0.21		1	1	
Clostridium butyricum		4.48	-.03		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67774142	**Good identification**						**67774142**
					GRA		
Actinomyces israelii		95.98	0.41	GAL 99	99		
Mitsuokella multacida		4.02	0.04		1		
67774312	**Good identification**						**67774312**
					GRA		
Clostridium clostridioforme		97.61	0.37	NIT 1	50		
Actinomyces israelii		1.26	0.14	ARA 94	99		
Mitsuokella multacida		1.13	0.01		1		
67774332	**Acceptable identification**						**67774332**
Clostridium clostridioforme		93.96	0.37	NIT 1			
Parabacteroides distasonis		6.04	0.23	ARA 94			
67774402	**Low discrimination**						**67774402**
					SPO		
Actinomyces israelii		54.60	0.36	GAL 99	1		
Clostridium clostridioforme		32.53	0.23		66		
Clostridium butyricum		8.52	0.01		99		
67774500	**Low discrimination**						**67774500**
					GRA	SPO	
Mitsuokella multacida		85.23	0.55	GAL 99	1	1	
Clostridium butyricum		11.57	0.32		99	99	
Actinomyces israelii		3.20	0.43		99	1	
67774501	**Good identification**						**67774501**
					GRA	SPO	
Mitsuokella multacida		98.78	0.39	GAL 99	1	1	
Clostridium butyricum		1.22	-.03	URE 10	99	99	
67774502	**Low discrimination**						**67774502**
					GRA	SPO	
Mitsuokella multacida		42.97	0.39	GAL 99	1	1	
Actinomyces israelii		21.80	0.46	BGL 10	99	1	
Clostridium butyricum		20.42	0.25		99	99	
67774510	**Good identification**						**67774510**
					GRA	SPO	
Mitsuokella multacida		95.95	0.47	GAL 99	1	1	
Clostridium butyricum		2.06	0.10		99	99	
Actinomyces israelii		1.62	0.29		99	1	
67774512	**Low discrimination**						**67774512**
					GRA		
Clostridium clostridioforme		67.43	0.43	NIT 1	50		
Mitsuokella multacida		19.50	0.30	RHA 88	1		
Bifidobacterium adolescentis		6.55	0.29		99		
67774520	**Acceptable identification**						**67774520**
					GRA	SPO	
Mitsuokella multacida		94.62	0.45	GAL 99	1	1	
Clostridium butyricum		4.47	0.14		99	99	
Actinomyces israelii		0.91	0.23		99	1	
67774530	**Good identification**						**67774530**
Mitsuokella multacida		98.47	0.37	GAL 99			
Clostridium clostridioforme		1.53	0.09				
67774532	**Acceptable identification**						**67774532**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Clostridium clostridioforme		91.32	0.43	NIT 1	50		
Mitsuokella multacida		6.60	0.20	RHA 88	1		
Bifidobacterium dentium		1.52	-.03		99		
67774540	**Good identification**						**67774540**
					GRA	SPO	
Mitsuokella multacida		97.43	0.55	GAL 99	1	1	
Actinomyces israelii		1.42	0.36		99	1	
Clostridium butyricum		1.15	0.14		99	99	
67774541	**Good identification**						**67774541**
Mitsuokella multacida		100.0	0.39	GAL 99			
67774542	**Low discrimination**						**67774542**
					GRA	SPO	
Mitsuokella multacida		80.00	0.39	GAL 99	1	1	
Actinomyces israelii		15.79	0.39	BGL 10	99	1	
Clostridium butyricum		3.31	0.07		99	99	
67774550	**Good identification**						**67774550**
					COC	GRA	
Mitsuokella multacida		99.00	0.47	GAL 99	1	1	
Actinomyces israelii		0.65	0.22		1	99	
Blautia producta		0.35	-.03		99	99	
67774560	**Good identification**						**67774560**
					GRA	SPO	
Mitsuokella multacida		99.22	0.45	GAL 99	1	1	
Clostridium butyricum		0.41	-.05		99	99	
Actinomyces israelii		0.37	0.16		99	1	
67774570	**Good identification**						**67774570**
Mitsuokella multacida		100.0	0.37	GAL 99			
67774602	**Good identification**						**67774602**
					GRA		
Clostridium clostridioforme		95.14	0.37	NIT 1	50		
Mitsuokella multacida		3.22	0.09		1		
Actinomyces israelii		1.64	0.17		99		
67774612	**Good identification**						**67774612**
Clostridium clostridioforme		99.75	0.49	NIT 1			
Mitsuokella multacida		0.25	0.01				
67774622	**Good identification**						**67774622**
Clostridium clostridioforme		100.0	0.37	NIT 1			
67774632	**Good identification**						**67774632**
Clostridium clostridioforme		100.0	0.49	NIT 1			
67774700	**Very good identification**						**67774700**
					GRA	SPO	
Mitsuokella multacida		99.51	0.60	GAL 99	1	1	
Clostridium butyricum		0.23	0.06		99	99	
Actinomyces israelii		0.15	0.24		99	1	
67774701	**Good identification**						**67774701**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.43	GAL 99			
67774702	**Low discrimination**						**67774702**
					GRA		
Mitsuokella multacida		49.32	0.43	GAL 99	1		
Clostridium clostridioforme		49.28	0.46	BGL 10	50		
Actinomyces israelii		1.01	0.27		99		
67774710	**Good identification**						**67774710**
Mitsuokella multacida		98.49	0.51	GAL 99			
Clostridium clostridioforme		1.51	0.24				
67774711	**Good identification**						**67774711**
Mitsuokella multacida		100.0	0.35	GAL 99			
67774712	**Good identification**						**67774712**
Clostridium clostridioforme		93.18	0.58	NIT 1			
Mitsuokella multacida		6.82	0.35				
67774720	**Good identification**						**67774720**
Mitsuokella multacida		99.55	0.49	GAL 99			
Clostridium clostridioforme		0.45	0.12				
67774722	**Low discrimination**						**67774722**
Clostridium clostridioforme		77.61	0.46	NIT 1			
Mitsuokella multacida		19.42	0.33				
Parabacteroides distasonis		1.65	0.23				
67774730	**Acceptable identification**						**67774730**
Mitsuokella multacida		94.22	0.41	GAL 99			
Clostridium clostridioforme		5.78	0.24				
67774732	**Very good identification**						**67774732**
Clostridium clostridioforme		97.72	0.58	NIT 1			
Mitsuokella multacida		1.79	0.25				
Parabacteroides distasonis		0.25	0.19				
67774740	**Very good identification**						**67774740**
					GRA		
Mitsuokella multacida		99.94	0.60	GAL 99	1		
Actinomyces israelii		0.06	0.17		99		
67774741	**Good identification**						**67774741**
Mitsuokella multacida		100.0	0.43	GAL 99			
67774742	**Good identification**						**67774742**
					GRA		
Mitsuokella multacida		98.22	0.43	GAL 99	1		
Clostridium clostridioforme		0.99	0.12	BGL 10	50		
Actinomyces israelii		0.79	0.20		99		
67774750	**Good identification**						**67774750**
					COC	GRA	
Mitsuokella multacida		99.81	0.51	GAL 99	1	1	
Blautia producta		0.19	-.03		99	99	
67774751	**Good identification**						**67774751**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.35	GAL 99			
67774752	**Low discrimination**						**67774752**
Mitsuokella multacida		87.88	0.35	GAL 99			
Clostridium clostridioforme		12.12	0.24	BGL 10			
67774760	**Good identification**						**67774760**
Mitsuokella multacida		100.0	0.49	GAL 99			
67774770	**Good identification**						**67774770**
Mitsuokella multacida		100.0	0.41	GAL 99			
67775000	**Good identification**						**67775000**
					GRA	SPO	
Actinomyces israelii		93.73	0.69		99	1	
Clostridium butyricum		5.13	0.26		99	99	
Mitsuokella multacida		0.83	0.21		1	1	
67775002	**Very good identification**						**67775002**
					SPO		
Actinomyces israelii		98.15	0.72		1		
Clostridium butyricum		1.39	0.19		99		
Actinomyces naeslundii		0.22	0.35		1		
67775003	**Low discrimination**						**67775003**
Actinomyces israelii		75.24	0.37	URE 1			
Actinomyces naeslundii		24.77	0.38				
67775010	**Good identification**						**67775010**
					GRA	SPO	
Actinomyces israelii		96.24	0.54	MLZ 13	99	1	
Mitsuokella multacida		1.91	0.13		1	1	
Clostridium butyricum		1.86	0.04		99	99	
67775012	**Good identification**						**67775012**
					GRA		
Actinomyces israelii		93.06	0.57	MLZ 13	99		
Leptotrichia buccalis		3.13	0.07		1		
Clostridium clostridioforme		2.68	0.22		50		
67775020	**Low discrimination**						**67775020**
					SPO		
Actinomyces israelii		86.10	0.48	NAG 6	1		
Clostridium butyricum		6.42	0.08		99		
Actinomyces naeslundii		4.49	0.35		1		
67775021	**Low discrimination**						**67775021**
Actinomyces naeslundii		88.55	0.38	ESL 12			
Actinomyces israelii		11.45	0.14	XYL 1			
67775022	**Low discrimination**						**67775022**
					GRA		
Actinomyces israelii		88.93	0.51	NAG 6	99		
Actinomyces naeslundii		3.09	0.35		99		
Prevotella oralis		3.00	0.09		1		
67775023	**Low discrimination**						**67775023**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		83.76	0.38	ESL 12			
Actinomyces israelii		16.24	0.17	XYL 1			
67775032	**Low discrimination**						**67775032**
					GRA		
Actinomyces israelii		62.98	0.37	MLZ 13	99		
Clostridium clostridioforme		28.37	0.22	NAG 6	50		
Parabacteroides distasonis		4.98	0.14		1		
67775040	**Very good identification**						**67775040**
					GRA	SPO	
Actinomyces israelii		96.47	0.62		99	1	
Mitsuokella multacida		2.20	0.21		1	1	
Clostridium butyricum		1.18	0.08		99	99	
67775042	**Very good identification**						**67775042**
					GRA	SPO	
Actinomyces israelii		99.42	0.65		99	1	
Clostridium butyricum		0.32	0.01		99	99	
Mitsuokella multacida		0.17	0.04		1	1	
67775050	**Good identification**						**67775050**
					GRA		
Actinomyces israelii		95.15	0.47	MLZ 13	99		
Mitsuokella multacida		4.85	0.13		1		
67775052	**Good identification**						**67775052**
Actinomyces israelii		100.0	0.50	MLZ 13			
67775060	**Low discrimination**						**67775060**
					GRA		
Actinomyces israelii		89.84	0.41	NAG 6	99		
Mitsuokella multacida		8.04	0.11		1		
Actinomyces naeslundii		2.12	0.22		99		
67775062	**Good identification**						**67775062**
Actinomyces israelii		98.45	0.44	NAG 6			
Actinomyces naeslundii		1.55	0.22				
67775100	**Low discrimination**						**67775100**
					GRA	SPO	
Actinomyces israelii		64.00	0.79		99	1	
Clostridium butyricum		21.02	0.50		99	99	
Mitsuokella multacida		14.07	0.55		1	1	
67775101	**Low discrimination**						**67775101**
					GRA		
Mitsuokella multacida		59.02	0.39	ARA 99	1		
Actinomyces israelii		24.40	0.45	URE 10	99		
Actinomyces naeslundii		8.57	0.46		99		
67775102	**Low discrimination**						**67775102**
					SPO		
Actinomyces israelii		88.43	0.82		1		
Clostridium butyricum		7.53	0.43		99		
Bifidobacterium adolescentis		1.95	0.47		1		
67775103	**Low discrimination**						**67775103**
					GRA		
Actinomyces israelii		65.76	0.48	URE 1	99		
Actinomyces naeslundii		15.40	0.46		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		11.78	0.22		1		
67775110	**Low discrimination**						**67775110**
					GRA	SPO	
Actinomyces israelii		61.67	0.65	MLZ 13	99	1	
Mitsuokella multacida		30.25	0.47		1	1	
Clostridium butyricum		7.13	0.28		99	99	
67775112	**Low discrimination**						**67775112**
					GRA		
Actinomyces israelii		47.39	0.68	MLZ 13	99		
Leptotrichia buccalis		39.38	0.41		1		
Bifidobacterium adolescentis		8.55	0.49		99		
67775120	**Low discrimination**						**67775120**
					GRA	SPO	
Actinomyces israelii		41.30	0.58	NAG 6	99	1	
Mitsuokella multacida		35.57	0.45		1	1	
Clostridium butyricum		18.48	0.31		99	99	
67775122	**Low discrimination**						**67775122**
					GRA		
Actinomyces israelii		50.45	0.61	NAG 6	99		
Parabacteroides distasonis		27.38	0.53		1		
Prevotella oralis		6.65	0.30		1		
67775123	**Low discrimination**						**67775123**
					GRA		
Actinomyces naeslundii		62.05	0.46	ESL 12	99		
Actinomyces israelii		16.91	0.27	XYL 1	99		
Mitsuokella multacida		11.87	0.12		1		
67775130	**Low discrimination**						**67775130**
					GRA	SPO	
Mitsuokella multacida		60.89	0.37	ARA 99	1	1	
Actinomyces israelii		31.69	0.44		99	1	
Clostridium butyricum		4.99	0.10		99	99	
67775132	**Low discrimination**						**67775132**
					GRA		
Parabacteroides distasonis		37.67	0.49	NIT 1	1		
Leptotrichia buccalis		27.86	0.25		1		
Actinomyces israelii		19.26	0.47		99		
67775140	**Low discrimination**						**67775140**
					GRA	SPO	
Actinomyces israelii		60.98	0.72		99	1	
Mitsuokella multacida		34.48	0.55		1	1	
Clostridium butyricum		4.48	0.31		99	99	
67775141	**Low discrimination**						**67775141**
					GRA		
Mitsuokella multacida		83.46	0.39	ARA 99	1		
Actinomyces israelii		13.42	0.38	URE 10	99		
Actinomyces naeslundii		2.14	0.33		99		
67775142	**Good identification**						**67775142**
					GRA	SPO	
Actinomyces israelii		93.37	0.75		99	1	
Mitsuokella multacida		3.91	0.39		1	1	
Clostridium butyricum		1.78	0.24		99	99	
67775143	**Low discrimination**						**67775143**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Actinomyces israelii		63.82	0.41	URE 1	99		
Mitsuokella multacida		29.40	0.22		1		
Actinomyces naeslundii		6.78	0.33		99		
67775150	**Low discrimination**						**67775150**
					GRA	SPO	
Mitsuokella multacida		55.04	0.47	ARA 99	1	1	
Actinomyces israelii		43.64	0.58		99	1	
Clostridium butyricum		1.13	0.10		99	99	
67775152	**Low discrimination**						**67775152**
					GRA		
Actinomyces israelii		83.71	0.61	MLZ 13	99		
Mitsuokella multacida		7.82	0.30		1		
Bifidobacterium adolescentis		4.80	0.33		99		
67775160	**Low discrimination**						**67775160**
					GRA	SPO	
Mitsuokella multacida		66.29	0.45	ARA 99	1	1	
Actinomyces israelii		29.93	0.51		99	1	
Clostridium butyricum		2.99	0.13		99	99	
67775162	**Low discrimination**						**67775162**
					GRA		
Actinomyces israelii		78.13	0.54	NAG 6	99		
Mitsuokella multacida		12.82	0.28		1		
Bifidobacterium breve		5.05	0.29		99		
67775170	**Low discrimination**						**67775170**
					GRA		
Mitsuokella multacida		82.28	0.37	ARA 99	1		
Actinomyces israelii		16.65	0.37		99		
Bifidobacterium breve		1.07	0.12		99		
67775172	**Low discrimination**						**67775172**
					GRA		
Actinomyces israelii		52.99	0.40	MLZ 13	99		
Bifidobacterium breve		22.94	0.29	NAG 6	99		
Mitsuokella multacida		19.39	0.20		1		
67775200	**Low discrimination**						**67775200**
					GRA	SPO	
Actinomyces israelii		80.53	0.49	RHA 7	99	1	
Mitsuokella multacida		17.66	0.25		1	1	
Clostridium butyricum		1.81	0.00		99	99	
67775202	**Low discrimination**						**67775202**
					GRA		
Actinomyces israelii		89.77	0.52	RHA 7	99		
Clostridium clostridioforme		8.25	0.25		50		
Mitsuokella multacida		1.46	0.09		1		
67775210	**Low discrimination**						**67775210**
					GRA		
Actinomyces israelii		67.16	0.35	RHA 7	99		
Mitsuokella multacida		32.84	0.17	MLZ 13	1		
67775212	**Low discrimination**						**67775212**
					GRA		
Clostridium clostridioforme		73.00	0.36	NIT 1	50		
Actinomyces israelii		26.06	0.38	ARA 94	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		0.94	0.01		1		
67775232	**Acceptable identification**						**67775232**
					GRA		
Clostridium clostridioforme		91.36	0.36	NIT 1	50		
Parabacteroides distasonis		6.56	0.23	ARA 94	1		
Actinomyces israelii		2.08	0.18		99		
67775240	**Low discrimination**						**67775240**
					GRA		
Actinomyces israelii		63.95	0.42	RHA 7	99		
Mitsuokella multacida		36.05	0.25		1		
67775242	**Good identification**						**67775242**
					GRA		
Actinomyces israelii		95.99	0.45	RHA 7	99		
Mitsuokella multacida		4.01	0.09		1		
67775300	**Low discrimination**						**67775300**
					GRA	SPO	
Mitsuokella multacida		82.70	0.60	ARA 99	1	1	
Actinomyces israelii		15.24	0.60		99	1	
Clostridium butyricum		2.06	0.24		99	99	
67775301	**Good identification**						**67775301**
					GRA		
Mitsuokella multacida		98.35	0.43	ARA 99	1		
Actinomyces israelii		1.65	0.25	URE 10	99		
67775302	**Low discrimination**						**67775302**
					GRA		
Actinomyces israelii		65.22	0.63	RHA 7	99		
Mitsuokella multacida		26.21	0.43		1		
Clostridium clostridioforme		5.02	0.34		50		
67775310	**Acceptable identification**						**67775310**
					GRA	SPO	
Mitsuokella multacida		91.79	0.51	ARA 99	1	1	
Actinomyces israelii		7.58	0.46		99	1	
Clostridium butyricum		0.36	0.02		99	99	
67775311	**Good identification**						**67775311**
Mitsuokella multacida		100.0	0.35	ARA 99			
67775312	**Low discrimination**						**67775312**
					GRA		
Clostridium clostridioforme		52.81	0.45	NIT 1	50		
Actinomyces israelii		22.53	0.49	ARA 94	99		
Mitsuokella multacida		20.20	0.35		1		
67775320	**Acceptable identification**						**67775320**
					GRA		
Mitsuokella multacida		92.55	0.49	ARA 99	1		
Actinomyces israelii		4.36	0.39		99		
Parabacteroides distasonis		1.43	0.27		1		
67775322	**Low discrimination**						**67775322**
Parabacteroides distasonis		80.38	0.61	NIT 1			
Mitsuokella multacida		5.85	0.33				
Clostridium clostridioforme		4.48	0.34				
67775330	**Acceptable identification**						**67775330**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Mitsuokella multacida		94.54	0.41	ARA 99	1		
Parabacteroides distasonis		2.36	0.23		1		
Actinomyces israelii		1.99	0.25		99		
67775332	**Low discrimination**						**67775332**
Parabacteroides distasonis		64.50	0.57	NIT 1			
Clostridium clostridioforme		30.39	0.45				
Mitsuokella multacida		2.91	0.25				
67775340	**Good identification**						**67775340**
					GRA	SPO	
Mitsuokella multacida		93.12	0.60	ARA 99	1	1	
Actinomyces israelii		6.68	0.53		99	1	
Clostridium butyricum		0.20	0.06		99	99	
67775341	**Good identification**						**67775341**
					GRA		
Mitsuokella multacida		99.35	0.43	ARA 99	1		
Actinomyces israelii		0.65	0.18	URE 10	99		
67775342	**Low discrimination**						**67775342**
					GRA	SPO	
Mitsuokella multacida		50.63	0.43	ARA 99	1	1	
Actinomyces israelii		48.99	0.56	BGL 10	99	1	
Clostridium butyricum		0.38	-.02		99	99	
67775350	**Good identification**						**67775350**
					COC	GRA	
Mitsuokella multacida		96.71	0.51	ARA 99	1	1	
Actinomyces israelii		3.11	0.38		1	99	
Blautia producta		0.18	-.03		99	99	
67775351	**Good identification**						**67775351**
Mitsuokella multacida		100.0	0.35	ARA 99			
67775352	**Low discrimination**						**67775352**
					GRA		
Mitsuokella multacida		68.49	0.35	ARA 99	1		
Actinomyces israelii		29.71	0.42	BGL 10	99		
Clostridium clostridioforme		1.81	0.11		50		
67775360	**Good identification**						**67775360**
					GRA		
Mitsuokella multacida		98.20	0.49	ARA 99	1		
Actinomyces israelii		1.80	0.32		99		
67775370	**Good identification**						**67775370**
					GRA		
Mitsuokella multacida		99.19	0.41	ARA 99	1		
Actinomyces israelii		0.81	0.18		99		
67775400	**Low discrimination**						**67775400**
					GRA	SPO	
Mitsuokella multacida		40.16	0.55	LAC 99	1	1	
Actinomyces israelii		37.35	0.67		99	1	
Clostridium butyricum		22.49	0.42		99	99	
67775401	**Low discrimination**						**67775401**
					GRA	SPO	
Mitsuokella multacida		88.07	0.39	LAC 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		7.45	0.33	URE 10	99	1	
Clostridium butyricum		4.48	0.08		99	99	
67775402	**Low discrimination**						**67775402**
					GRA	SPO	
Actinomyces israelii		79.30	0.70		99	1	
Clostridium butyricum		12.38	0.35		99	99	
Mitsuokella multacida		6.32	0.39		1	1	
67775403	**Low discrimination**						**67775403**
					GRA	SPO	
Actinomyces israelii		49.21	0.36	URE 1	99	1	
Mitsuokella multacida		43.11	0.22		1	1	
Clostridium butyricum		7.68	0.01		99	99	
67775410	**Low discrimination**						**67775410**
					GRA	SPO	
Mitsuokella multacida		66.27	0.47	LAC 99	1	1	
Actinomyces israelii		27.64	0.53		99	1	
Clostridium butyricum		5.86	0.20		99	99	
67775412	**Low discrimination**						**67775412**
					GRA		
Actinomyces israelii		54.10	0.56	MLZ 13	99		
Clostridium clostridioforme		29.78	0.42		50		
Mitsuokella multacida		9.61	0.30		1		
67775420	**Low discrimination**						**67775420**
					GRA	SPO	
Mitsuokella multacida		69.82	0.45	LAC 99	1	1	
Actinomyces israelii		16.58	0.47		99	1	
Clostridium butyricum		13.60	0.24		99	99	
67775422	**Low discrimination**						**67775422**
					GRA		
Actinomyces israelii		54.83	0.50	NAG 6	99		
Mitsuokella multacida		17.10	0.28		1		
Clostridium clostridioforme		15.51	0.31		50		
67775430	**Low discrimination**						**67775430**
					GRA	SPO	
Mitsuokella multacida		86.87	0.37	LAC 99	1	1	
Actinomyces israelii		9.25	0.32		99	1	
Clostridium butyricum		2.67	0.02		99	99	
67775432	**Low discrimination**						**67775432**
					GRA		
Clostridium clostridioforme		81.26	0.42	NIT 1	50		
Actinomyces israelii		9.42	0.36	RHA 88	99		
Mitsuokella multacida		6.56	0.20		1		
67775440	**Low discrimination**						**67775440**
					GRA	SPO	
Mitsuokella multacida		70.90	0.55	LAC 99	1	1	
Actinomyces israelii		25.65	0.60		99	1	
Clostridium butyricum		3.45	0.24		99	99	
67775441	**Good identification**						**67775441**
					GRA		
Mitsuokella multacida		96.82	0.39	LAC 99	1		
Actinomyces israelii		3.18	0.26	URE 10	99		
67775442	**Low discrimination**						**67775442**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA	SPO	
Actinomyces israelii		80.52	0.63		99	1	
Mitsuokella multacida		16.49	0.39		1	1	
Clostridium butyricum		2.81	0.17		99	99	
67775450	**Low discrimination**						**67775450**
					GRA	SPO	
Mitsuokella multacida		85.22	0.47	LAC 99	1	1	
Actinomyces israelii		13.82	0.46		99	1	
Clostridium butyricum		0.66	0.02		99	99	
67775452	**Low discrimination**						**67775452**
					GRA		
Actinomyces israelii		66.56	0.49	MLZ 13	99		
Mitsuokella multacida		30.41	0.30		1		
Bifidobacterium adolescentis		1.26	0.13		99		
67775460	**Acceptable identification**						**67775460**
					GRA	SPO	
Mitsuokella multacida		90.15	0.45	LAC 99	1	1	
Actinomyces israelii		8.33	0.40		99	1	
Clostridium butyricum		1.53	0.06		99	99	
67775462	**Low discrimination**						**67775462**
					GRA	SPO	
Actinomyces israelii		54.07	0.43	NAG 6	99	1	
Mitsuokella multacida		43.36	0.28		1	1	
Clostridium butyricum		2.57	-.01		99	99	
67775470	**Good identification**						**67775470**
					GRA		
Mitsuokella multacida		96.02	0.37	LAC 99	1		
Actinomyces israelii		3.98	0.25		99		
67775500	**Low discrimination**						**67775500**
					GRA	SPO	
Mitsuokella multacida		85.22	0.89		1	1	
Clostridium butyricum		11.57	0.66		99	99	
Actinomyces israelii		3.20	0.77		99	1	
67775501	**Very good identification**						**67775501**
					GRA	SPO	
Mitsuokella multacida		98.45	0.73	URE 10	1	1	
Clostridium butyricum		1.22	0.32		99	99	
Actinomyces israelii		0.34	0.43		99	1	
67775502	**Low discrimination**						**67775502**
					GRA	SPO	
Mitsuokella multacida		48.03	0.73	BGL 10	1	1	
Actinomyces israelii		24.37	0.80		99	1	
Clostridium butyricum		22.82	0.59		99	99	
67775503	**Good identification**						**67775503**
					GRA	SPO	
Mitsuokella multacida		91.05	0.56	URE 10	1	1	
Actinomyces israelii		4.20	0.46	BGL 10	99	1	
Clostridium butyricum		3.93	0.25		99	99	
67775510	**Very good identification**						**67775510**
					GRA	SPO	
Mitsuokella multacida		96.23	0.81		1	1	
Clostridium butyricum		2.06	0.44		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		1.62	0.63		99	1	
67775511	**Very good identification**						**67775511**
					GRA	SPO	
Mitsuokella multacida		99.65	0.65	URE 10	1	1	
Clostridium butyricum		0.19	0.10		99	99	
Actinomyces israelii		0.15	0.29		99	1	
67775512	**Low discrimination**						**67775512**
					GRA		
Mitsuokella multacida		56.25	0.65	BGL 10	1		
Bifidobacterium adolescentis		18.88	0.63		99		
Actinomyces israelii		12.79	0.66		99		
67775513	**Acceptable identification**						**67775513**
					GRA		
Mitsuokella multacida		93.42	0.48	URE 10	1		
Bifidobacterium adolescentis		2.85	0.29	BGL 10	99		
Actinomyces israelii		1.93	0.32		99		
67775520	**Good identification**						**67775520**
					GRA	SPO	
Mitsuokella multacida		94.61	0.79		1	1	
Clostridium butyricum		4.47	0.48		99	99	
Actinomyces israelii		0.91	0.57		99	1	
67775521	**Very good identification**						**67775521**
					GRA	SPO	
Mitsuokella multacida		99.49	0.63	URE 10	1	1	
Clostridium butyricum		0.43	0.14		99	99	
Actinomyces israelii		0.09	0.23		99	1	
67775522	**Low discrimination**						**67775522**
					GRA	SPO	
Mitsuokella multacida		71.89	0.63	BGL 10	1	1	
Clostridium butyricum		11.88	0.41		99	99	
Actinomyces israelii		9.31	0.60		99	1	
67775523	**Good identification**						**67775523**
					GRA	SPO	
Mitsuokella multacida		96.68	0.46	URE 10	1	1	
Clostridium butyricum		1.45	0.07	BGL 10	99	99	
Actinomyces israelii		1.14	0.26		99	1	
67775530	**Very good identification**						**67775530**
					GRA	SPO	
Mitsuokella multacida		98.54	0.71		1	1	
Clostridium butyricum		0.74	0.26		99	99	
Actinomyces israelii		0.42	0.43		99	1	
67775531	**Good identification**						**67775531**
Mitsuokella multacida		100.0	0.54	URE 10			
67775532	**Low discrimination**						**67775532**
					GRA		
Mitsuokella multacida		54.17	0.54	BGL 10	1		
Clostridium clostridioforme		22.71	0.51		50		
Bifidobacterium dentium		12.44	0.32		99		
67775533	**Acceptable identification**						**67775533**
					GRA		
Mitsuokella multacida		93.75	0.38	URE 10	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		3.57	0.17	BGL 10	50		
Bifidobacterium dentium		1.96	-.03		99		
67775540	**Very good identification**						**67775540**
					GRA	SPO	
Mitsuokella multacida		97.39	0.89		1	1	
Actinomyces israelii		1.42	0.70		99	1	
Clostridium butyricum		1.15	0.48		99	99	
67775541	**Very good identification**						**67775541**
					GRA	SPO	
Mitsuokella multacida		99.76	0.73	URE 10	1	1	
Actinomyces israelii		0.13	0.36		99	1	
Clostridium butyricum		0.11	0.14		99	99	
67775542	**Low discrimination**						**67775542**
					GRA	SPO	
Mitsuokella multacida		79.97	0.73	BGL 10	1	1	
Actinomyces israelii		15.78	0.73		99	1	
Clostridium butyricum		3.31	0.41		99	99	
67775543	**Very good identification**						**67775543**
					GRA	SPO	
Mitsuokella multacida		97.78	0.56	URE 10	1	1	
Actinomyces israelii		1.75	0.39	BGL 10	99	1	
Clostridium butyricum		0.37	0.07		99	99	
67775550	**Very good identification**						**67775550**
					COC	GRA	
Mitsuokella multacida		98.81	0.81		1	1	
Actinomyces israelii		0.65	0.56		1	99	
Blautia producta		0.35	0.32		99	99	
67775551	**Very good identification**						**67775551**
					COC	GRA	
Mitsuokella multacida		99.91	0.65	URE 10	1	1	
Actinomyces israelii		0.06	0.22		1	99	
Blautia producta		0.03	-.03		99	99	
67775552	**Low discrimination**						**67775552**
					GRA		
Mitsuokella multacida		87.61	0.65	BGL 10	1		
Actinomyces israelii		7.75	0.59		99		
Bifidobacterium adolescentis		3.64	0.47		99		
67775553	**Good identification**						**67775553**
					GRA		
Mitsuokella multacida		98.83	0.48	URE 10	1		
Actinomyces israelii		0.80	0.25	BGL 10	99		
Bifidobacterium adolescentis		0.37	0.13		99		
67775560	**Very good identification**						**67775560**
					GRA	SPO	
Mitsuokella multacida		99.21	0.79		1	1	
Clostridium butyricum		0.41	0.30		99	99	
Actinomyces israelii		0.37	0.50		99	1	
67775561	**Very good identification**						**67775561**
					GRA	SPO	
Mitsuokella multacida		99.93	0.63	URE 10	1	1	
Clostridium butyricum		0.04	-.05		99	99	
Actinomyces israelii		0.03	0.16		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67775562	**Good identification**						**67775562**
					GRA	SPO	
Mitsuokella multacida		93.78	0.63	BGL 10	1	1	
Actinomyces israelii		4.72	0.53		99	1	
Clostridium butyricum		1.35	0.23		99	99	
67775563	**Good identification**						**67775563**
					GRA		
Mitsuokella multacida		99.54	0.46	URE 10	1		
Actinomyces israelii		0.46	0.19	BGL 10	99		
67775570	**Very good identification**						**67775570**
					COC	GRA	
Mitsuokella multacida		99.61	0.71		1	1	
Actinomyces israelii		0.17	0.36		1	99	
Blautia producta		0.16	0.15		99	99	
67775571	**Good identification**						**67775571**
Mitsuokella multacida		100.0	0.54	URE 10			
67775572	**Good identification**						**67775572**
					GRA		
Mitsuokella multacida		96.57	0.54	BGL 10	1		
Actinomyces israelii		2.18	0.39		99		
Clostridium clostridioforme		0.41	0.17		50		
67775573	**Good identification**						**67775573**
Mitsuokella multacida		100.0	0.38	URE 10			
67775600	**Very good identification**						**67775600**
					GRA	SPO	
Mitsuokella multacida		95.42	0.60	LAC 99	1	1	
Actinomyces israelii		3.60	0.48		99	1	
Clostridium butyricum		0.89	0.16		99	99	
67775601	**Good identification**						**67775601**
					GRA		
Mitsuokella multacida		99.66	0.43	LAC 99	1		
Actinomyces israelii		0.34	0.14	URE 10	99		
67775602	**Low discrimination**						**67775602**
					GRA		
Mitsuokella multacida		41.03	0.43	LAC 99	1		
Clostridium clostridioforme		36.74	0.46	BGL 10	50		
Actinomyces israelii		20.88	0.51		99		
67775610	**Good identification**						**67775610**
					GRA		
Mitsuokella multacida		96.89	0.51	LAC 99	1		
Actinomyces israelii		1.64	0.34		99		
Clostridium clostridioforme		1.33	0.23		50		
67775611	**Good identification**						**67775611**
Mitsuokella multacida		100.0	0.35	LAC 99			
67775612	**Good identification**						**67775612**
					GRA		
Clostridium clostridioforme		90.88	0.57	NIT 1	50		
Mitsuokella multacida		7.43	0.35		1		
Actinomyces israelii		1.69	0.37		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67775620	**Good identification**						**67775620**
					GRA		
Mitsuokella multacida		98.34	0.49	LAC 99	1		
Actinomyces israelii		0.95	0.27		99		
Clostridium clostridioforme		0.40	0.11		50		
67775622	**Low discrimination**						**67775622**
					GRA		
Clostridium clostridioforme		73.62	0.46	NIT 1	50		
Mitsuokella multacida		20.56	0.33		1		
Actinomyces israelii		2.67	0.30		99		
67775630	**Acceptable identification**						**67775630**
					GRA		
Mitsuokella multacida		94.41	0.41	LAC 99	1		
Clostridium clostridioforme		5.19	0.23		50		
Actinomyces israelii		0.41	0.13		99		
67775632	**Very good identification**						**67775632**
Clostridium clostridioforme		97.35	0.57	NIT 1			
Mitsuokella multacida		1.99	0.25				
Parabacteroides distasonis		0.27	0.19				
67775640	**Very good identification**						**67775640**
					GRA	SPO	
Mitsuokella multacida		98.48	0.60	LAC 99	1	1	
Actinomyces israelii		1.44	0.41		99	1	
Clostridium butyricum		0.08	-.02		99	99	
67775641	**Good identification**						**67775641**
Mitsuokella multacida		100.0	0.43	LAC 99			
67775642	**Low discrimination**						**67775642**
					GRA		
Mitsuokella multacida		82.85	0.43	LAC 99	1		
Actinomyces israelii		16.40	0.44	BGL 10	99		
Clostridium clostridioforme		0.75	0.11		50		
67775650	**Good identification**						**67775650**
					COC	GRA	
Mitsuokella multacida		99.16	0.51	LAC 99	1	1	
Actinomyces israelii		0.65	0.27		1	99	
Blautia producta		0.19	-.03		99	99	
67775651	**Good identification**						**67775651**
Mitsuokella multacida		100.0	0.35	LAC 99			
67775652	**Low discrimination**						**67775652**
					GRA		
Mitsuokella multacida		82.49	0.35	LAC 99	1		
Clostridium clostridioforme		10.20	0.23	BGL 10	50		
Actinomyces israelii		7.32	0.30		99		
67775660	**Good identification**						**67775660**
					GRA		
Mitsuokella multacida		99.63	0.49	LAC 99	1		
Actinomyces israelii		0.37	0.20		99		
67775670	**Good identification**						**67775670**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		100.0	0.41	LAC 99			
67775700	**Excellent identification**						**67775700**
					GRA	SPO	
Mitsuokella multacida		99.62	0.94		1	1	
Clostridium butyricum		0.23	0.40		99	99	
Actinomyces israelii		0.15	0.58		99	1	
67775701	**Very good identification**						**67775701**
					GRA	SPO	
Mitsuokella multacida		99.97	0.77	URE 10	1	1	
Clostridium butyricum		0.02	0.06		99	99	
Actinomyces israelii		0.01	0.24		99	1	
67775702	**Good identification**						**67775702**
					GRA		
Mitsuokella multacida		94.37	0.77	BGL 10	1		
Clostridium clostridioforme		2.86	0.55		50		
Actinomyces israelii		1.94	0.61		99		
67775703	**Very good identification**						**67775703**
					GRA		
Mitsuokella multacida		99.47	0.61	URE 10	1		
Clostridium clostridioforme		0.27	0.20	BGL 10	50		
Actinomyces israelii		0.19	0.27		99		
67775710	**Very good identification**						**67775710**
					GRA		
Mitsuokella multacida		99.85	0.86		1		
Actinomyces israelii		0.07	0.44		99		
Clostridium clostridioforme		0.05	0.32		50		
67775711	**Very good identification**						**67775711**
Mitsuokella multacida		100.0	0.69	URE 10			
67775712	**Low discrimination**						**67775712**
					GRA		
Mitsuokella multacida		70.08	0.69	BGL 10	1		
Clostridium clostridioforme		29.00	0.66		50		
Actinomyces israelii		0.65	0.47		99		
67775713	**Good identification**						**67775713**
Mitsuokella multacida		96.37	0.53	URE 10			
Clostridium clostridioforme		3.63	0.32	BGL 10			
67775720	**Very good identification**						**67775720**
					GRA	SPO	
Mitsuokella multacida		99.83	0.83		1	1	
Clostridium butyricum		0.08	0.22		99	99	
Actinomyces israelii		0.04	0.38		99	1	
67775721	**Very good identification**						**67775721**
Mitsuokella multacida		100.0	0.67	URE 10			
67775722	**Low discrimination**						**67775722**
Mitsuokella multacida		77.70	0.67	BGL 10			
Clostridium clostridioforme		9.41	0.55				
Parabacteroides distasonis		6.61	0.58				
67775723	**Good identification**						**67775723**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		97.57	0.51	URE 10			
Clostridium clostridioforme		1.07	0.20	BGL 10			
Parabacteroides distasonis		0.76	0.23				
67775730	**Very good identification**						**67775730**
					GRA		
Mitsuokella multacida		99.75	0.75		1		
Clostridium clostridioforme		0.19	0.32		50		
Actinomyces israelii		0.02	0.24		99		
67775731	**Very good identification**						**67775731**
Mitsuokella multacida		100.0	0.59	URE 10			
67775732	**Low discrimination**						**67775732**
Clostridium clostridioforme		56.35	0.66	NIT 1			
Mitsuokella multacida		34.04	0.59				
Parabacteroides distasonis		4.68	0.53				
67775733	**Low discrimination**						**67775733**
Mitsuokella multacida		85.08	0.43	URE 10			
Clostridium clostridioforme		12.80	0.32	BGL 10			
Parabacteroides distasonis		1.06	0.19				
67775740	**Excellent identification**						**67775740**
					COC	GRA	
Mitsuokella multacida		99.90	0.94		1	1	
Actinomyces israelii		0.06	0.51		1	99	
Blautia producta		0.02	0.24		99	99	
67775741	**Very good identification**						**67775741**
					GRA		
Mitsuokella multacida		100.0	0.77	URE 10	1		
Actinomyces israelii		0.01	0.17		99		
67775742	**Very good identification**						**67775742**
					GRA	SPO	
Mitsuokella multacida		99.08	0.77	BGL 10	1	1	
Actinomyces israelii		0.79	0.54		99	1	
Clostridium butyricum		0.07	0.15		99	99	
67775743	**Very good identification**						**67775743**
					GRA		
Mitsuokella multacida		99.93	0.61	URE 10	1		
Actinomyces israelii		0.07	0.20	BGL 10	99		
67775750	**Very good identification**						**67775750**
					COC	GRA	
Mitsuokella multacida		99.78	0.86		1	1	
Blautia producta		0.19	0.32		99	99	
Actinomyces israelii		0.03	0.37		1	99	
67775751	**Very good identification**						**67775751**
					COC	GRA	
Mitsuokella multacida		99.98	0.69	URE 10	1	1	
Blautia producta		0.02	-.03		99	99	
67775752	**Very good identification**						**67775752**
					GRA		
Mitsuokella multacida		99.01	0.69	BGL 10	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium clostridioforme		0.41	0.32		50		
Actinomyces israelii		0.36	0.40		99		
67775753	**Good identification**						**67775753**
Mitsuokella multacida		100.0	0.53	URE 10			
67775760	**Very good identification**						**67775760**
					COC	GRA	
Mitsuokella multacida		99.97	0.83		1	1	
Actinomyces israelii		0.02	0.31		1	99	
Blautia producta		0.01	0.07		99	99	
67775761	**Very good identification**						**67775761**
Mitsuokella multacida		100.0	0.67	URE 10			
67775762	**Very good identification**						**67775762**
					GRA		
Mitsuokella multacida		99.29	0.67	BGL 10	1		
Bacteroides ovatus		0.28	0.16		1		
Actinomyces israelii		0.20	0.34		99		
67775763	**Good identification**						**67775763**
Mitsuokella multacida		100.0	0.51	URE 10			
67775770	**Very good identification**						**67775770**
					COC	GRA	
Mitsuokella multacida		99.91	0.75		1	1	
Blautia producta		0.09	0.15		99	99	
Actinomyces israelii		0.01	0.17		1	99	
67775771	**Very good identification**						**67775771**
Mitsuokella multacida		100.0	0.59	URE 10			
67775772	**Very good identification**						**67775772**
Mitsuokella multacida		97.51	0.59	BGL 10			
Clostridium clostridioforme		1.63	0.32				
Bacteroides ovatus		0.55	0.13				
67775773	**Good identification**						**67775773**
Mitsuokella multacida		100.0	0.43	URE 10			
67776000	**Good identification**						**67776000**
Actinomyces israelii		100.0	0.47	GAL 99			
67776002	**Good identification**						**67776002**
Actinomyces israelii		100.0	0.50	GAL 99			
67776012	**Good identification**						**67776012**
Actinomyces israelii		100.0	0.36	GAL 99			
67776040	**Good identification**						**67776040**
Actinomyces israelii		100.0	0.40	GAL 99			
67776042	**Good identification**						**67776042**
Actinomyces israelii		100.0	0.43	GAL 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67776100	**Good identification**						**67776100**
					GRA	SPO	
Actinomyces israelii		90.88	0.58	GAL 99	99	1	
Mitsuokella multacida		8.23	0.27		1	1	
Clostridium butyricum		0.59	-.01		99	99	
67776102	**Very good identification**						**67776102**
					GRA		
Actinomyces israelii		98.80	0.61	GAL 99	99		
Mitsuokella multacida		0.66	0.11		1		
Bifidobacterium adolescentis		0.32	0.12		99		
67776110	**Low discrimination**						**67776110**
					GRA		
Actinomyces israelii		83.20	0.43	GAL 99	99		
Mitsuokella multacida		16.80	0.19	MLZ 13	1		
67776112	**Acceptable identification**						**67776112**
Actinomyces israelii		94.21	0.46	GAL 99			
Bifidobacterium adolescentis		2.46	0.13	MLZ 13			
Bifidobacterium dentium		1.93	-.07				
67776120	**Low discrimination**						**67776120**
					GRA		
Actinomyces israelii		71.07	0.37	GAL 99	99		
Mitsuokella multacida		25.21	0.17	NAG 6	1		
Actinomyces naeslundii		3.72	0.24		99		
67776122	**Good identification**						**67776122**
Actinomyces israelii		96.63	0.40	GAL 99			
Actinomyces naeslundii		3.38	0.24	NAG 6			
67776140	**Low discrimination**						**67776140**
					GRA		
Actinomyces israelii		81.11	0.51	GAL 99	99		
Mitsuokella multacida		18.89	0.27		1		
67776142	**Good identification**						**67776142**
					GRA		
Actinomyces israelii		98.30	0.54	GAL 99	99		
Mitsuokella multacida		1.70	0.11		1		
67776150	**Low discrimination**						**67776150**
					COC	GRA	
Actinomyces israelii		62.61	0.36	GAL 99	1	99	
Mitsuokella multacida		32.52	0.19	MLZ 13	1	1	
Blautia producta		4.88	-.03		99	99	
67776152	**Good identification**						**67776152**
					GRA		
Actinomyces israelii		96.29	0.39	GAL 99	99		
Mitsuokella multacida		3.71	0.03	MLZ 13	1		
67776302	**Low discrimination**						**67776302**
					GRA		
Actinomyces israelii		85.80	0.41	GAL 99	99		
Mitsuokella multacida		14.20	0.15	RHA 7	1		
67776400	**Low discrimination**						**67776400**
					GRA		
Actinomyces israelii		69.32	0.46	GAL 99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		30.68	0.27		1		
67776402	**Good identification**						**67776402**
					GRA		
Actinomyces israelii		96.83	0.49	GAL 99	99		
Mitsuokella multacida		3.18	0.11		1		
67776442	**Acceptable identification**						**67776442**
					GRA		
Actinomyces israelii		92.22	0.42	GAL 99	99		
Mitsuokella multacida		7.78	0.11		1		
67776500	**Good identification**						**67776500**
					GRA	SPO	
Mitsuokella multacida		91.10	0.61	GAL 99	1	1	
Actinomyces israelii		8.32	0.56		99	1	
Clostridium butyricum		0.59	0.15		99	99	
67776501	**Good identification**						**67776501**
					GRA		
Mitsuokella multacida		99.18	0.45	GAL 99	1		
Actinomyces israelii		0.82	0.22	URE 10	99		
67776502	**Low discrimination**						**67776502**
					GRA		
Actinomyces israelii		53.77	0.59	GAL 99	99		
Mitsuokella multacida		43.64	0.45		1		
Bifidobacterium adolescentis		1.40	0.26		99		
67776510	**Acceptable identification**						**67776510**
					GRA		
Mitsuokella multacida		94.47	0.53	GAL 99	1		
Actinomyces israelii		3.87	0.42		99		
Bifidobacterium dentium		1.42	0.10		99		
67776511	**Good identification**						**67776511**
Mitsuokella multacida		100.0	0.37	GAL 99			
67776520	**Good identification**						**67776520**
					GRA	SPO	
Mitsuokella multacida		97.51	0.51	GAL 99	1	1	
Actinomyces israelii		2.27	0.36		99	1	
Clostridium butyricum		0.22	-.03		99	99	
67776530	**Low discrimination**						**67776530**
					GRA		
Mitsuokella multacida		89.49	0.43	GAL 99	1		
Bifidobacterium dentium		9.58	0.15		99		
Actinomyces israelii		0.94	0.22		99		
67776540	**Good identification**						**67776540**
					COC	GRA	
Mitsuokella multacida		94.99	0.61	GAL 99	1	1	
Actinomyces israelii		3.37	0.49		1	99	
Blautia producta		1.58	0.24		99	99	
67776541	**Good identification**						**67776541**
					GRA		
Mitsuokella multacida		99.68	0.45	GAL 99	1		
Actinomyces israelii		0.32	0.15	URE 10	99		
67776542	**Low discrimination**						**67776542**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		66.68	0.45	GAL 99	1	1	
Actinomyces israelii		31.95	0.52	BGL 10	1	99	
Blautia producta		1.11	0.07		99	99	
67776550	**Low discrimination**						**67776550**
					COC	GRA	
Mitsuokella multacida		85.77	0.53	GAL 99	1	1	
Blautia producta		12.87	0.32		99	99	
Actinomyces israelii		1.37	0.35		1	99	
67776551	**Good identification**						**67776551**
					COC	GRA	
Mitsuokella multacida		98.66	0.37	GAL 99	1	1	
Blautia producta		1.35	-.03	URE 10	99	99	
67776552	**Low discrimination**						**67776552**
					COC	GRA	
Mitsuokella multacida		71.78	0.37	GAL 99	1	1	
Actinomyces israelii		15.42	0.38	BGL 10	1	99	
Blautia producta		10.77	0.15		99	99	
67776560	**Good identification**						**67776560**
					COC	GRA	
Mitsuokella multacida		98.38	0.51	GAL 99	1	1	
Actinomyces israelii		0.89	0.29		1	99	
Blautia producta		0.73	0.07		99	99	
67776570	**Acceptable identification**						**67776570**
					COC	GRA	
Mitsuokella multacida		93.39	0.43	GAL 99	1	1	
Blautia producta		6.23	0.15		99	99	
Actinomyces israelii		0.38	0.15		1	99	
67776700	**Very good identification**						**67776700**
					GRA		
Mitsuokella multacida		99.63	0.66	GAL 99	1		
Actinomyces israelii		0.37	0.37		99		
67776701	**Good identification**						**67776701**
Mitsuokella multacida		100.0	0.50	GAL 99			
67776702	**Acceptable identification**						**67776702**
					GRA		
Mitsuokella multacida		94.85	0.50	GAL 99	1		
Actinomyces israelii		4.74	0.40	BGL 10	99		
Clostridium clostridioforme		0.41	0.12		50		
67776710	**Very good identification**						**67776710**
					COC	GRA	
Mitsuokella multacida		99.75	0.58	GAL 99	1	1	
Actinomyces israelii		0.17	0.23		1	99	
Blautia producta		0.08	-.03		99	99	
67776711	**Good identification**						**67776711**
Mitsuokella multacida		100.0	0.41	GAL 99			
67776712	**Acceptable identification**						**67776712**
					GRA		
Mitsuokella multacida		91.83	0.41	GAL 99	1		
Clostridium clostridioforme		5.43	0.24	BGL 10	50		
Actinomyces israelii		2.06	0.26		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67776720	**Good identification**						**67776720**
					GRA		
Mitsuokella multacida		99.91	0.56	GAL 99	1		
Actinomyces israelii		0.09	0.16		99		
67776721	**Good identification**						**67776721**
Mitsuokella multacida		100.0	0.39	GAL 99			
67776722	**Good identification**						**67776722**
					GRA		
Mitsuokella multacida		96.17	0.39	GAL 99	1		
Clostridium clostridioforme		1.66	0.12	BGL 10	50		
Actinomyces israelii		1.23	0.19		99		
67776730	**Good identification**						**67776730**
Mitsuokella multacida		100.0	0.47	GAL 99			
67776740	**Very good identification**						**67776740**
					COC	GRA	
Mitsuokella multacida		98.97	0.66	GAL 99	1	1	
Blautia producta		0.89	0.24		99	99	
Actinomyces israelii		0.14	0.30		1	99	
67776741	**Good identification**						**67776741**
Mitsuokella multacida		100.0	0.50	GAL 99			
67776742	**Good identification**						**67776742**
					COC	GRA	
Mitsuokella multacida		97.24	0.50	GAL 99	1	1	
Actinomyces israelii		1.89	0.33	BGL 10	1	99	
Blautia producta		0.87	0.07		99	99	
67776750	**Good identification**						**67776750**
					COC	GRA	
Mitsuokella multacida		92.47	0.58	GAL 99	1	1	
Blautia producta		7.47	0.32		99	99	
Actinomyces israelii		0.06	0.16		1	99	
67776751	**Good identification**						**67776751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	GAL 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67776752	**Acceptable identification**						**67776752**
					COC	GRA	
Mitsuokella multacida		91.79	0.41	GAL 99	1	1	
Blautia producta		7.41	0.15	BGL 10	99	99	
Actinomyces israelii		0.80	0.19		1	99	
67776760	**Good identification**						**67776760**
					COC	GRA	
Mitsuokella multacida		99.60	0.56	GAL 99	1	1	
Blautia producta		0.40	0.07		99	99	
67776761	**Good identification**						**67776761**
Mitsuokella multacida		100.0	0.39	GAL 99			
67776762	**Good identification**						**67776762**
Mitsuokella multacida		100.0	0.39	GAL 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67776770	**Good identification**						**67776770**
					COC	GRA	
Mitsuokella multacida		96.53	0.47	GAL 99	1	1	
Blautia producta		3.47	0.15		99	99	
67777000	**Very good identification**						**67777000**
					GRA		
Actinomyces israelii		99.42	0.81		99		
Mitsuokella multacida		0.36	0.27		1		
Actinomyces naeslundii		0.11	0.40		99		
67777001	**Low discrimination**						**67777001**
					GRA		
Actinomyces israelii		82.69	0.47	URE 1	99		
Actinomyces naeslundii		13.99	0.43		99		
Mitsuokella multacida		3.33	0.11		1		
67777002	**Very good identification**						**67777002**
					GRA		
Actinomyces israelii		99.86	0.84		99		
Actinomyces naeslundii		0.08	0.40		99		
Mitsuokella multacida		0.03	0.11		1		
67777003	**Low discrimination**						**67777003**
Actinomyces israelii		89.87	0.50	URE 1			
Actinomyces naeslundii		10.13	0.43				
67777010	**Very good identification**						**67777010**
					GRA		
Actinomyces israelii		99.19	0.67	MLZ 13	99		
Mitsuokella multacida		0.81	0.19		1		
67777012	**Very good identification**						**67777012**
Actinomyces israelii		99.75	0.70	MLZ 13			
Bifidobacterium adolescentis		0.11	0.13				
Bifidobacterium dentium		0.08	-.07				
67777013	**Good identification**						**67777013**
Actinomyces israelii		100.0	0.36	MLZ 13			
67777020	**Very good identification**						**67777020**
					GRA		
Actinomyces israelii		96.88	0.61	NAG 6	99		
Actinomyces naeslundii		1.73	0.40		99		
Mitsuokella multacida		1.39	0.17		1		
67777021	**Low discrimination**						**67777021**
Actinomyces naeslundii		72.60	0.43	ESL 12			
Actinomyces israelii		27.40	0.27	XYL 1			
67777022	**Very good identification**						**67777022**
Actinomyces israelii		98.82	0.64	NAG 6			
Actinomyces naeslundii		1.18	0.40				
67777023	**Low discrimination**						**67777023**
Actinomyces naeslundii		63.85	0.43	ESL 12			
Actinomyces israelii		36.15	0.30	XYL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777030	**Good identification**						**67777030**
					GRA		
Actinomyces israelii		96.90	0.47	MLZ 13	99		
Mitsuokella multacida		3.10	0.09	NAG 6	1		
67777032	**Good identification**						**67777032**
Actinomyces israelii		97.75	0.50	MLZ 13			
Bifidobacterium dentium		2.25	-.03	NAG 6			
67777040	**Very good identification**						**67777040**
					GRA		
Actinomyces israelii		99.02	0.74		99		
Mitsuokella multacida		0.93	0.27		1		
Actinomyces naeslundii		0.05	0.27		99		
67777041	**Low discrimination**						**67777041**
					GRA		
Actinomyces israelii		84.73	0.40	URE 1	99		
Mitsuokella multacida		8.77	0.11		1		
Actinomyces naeslundii		6.50	0.30		99		
67777042	**Very good identification**						**67777042**
					GRA		
Actinomyces israelii		99.90	0.77		99		
Mitsuokella multacida		0.07	0.11		1		
Actinomyces naeslundii		0.03	0.27		99		
67777043	**Good identification**						**67777043**
Actinomyces israelii		95.13	0.43	URE 1			
Actinomyces naeslundii		4.87	0.30				
67777050	**Very good identification**						**67777050**
					COC	GRA	
Actinomyces israelii		97.64	0.60	MLZ 13	1	99	
Mitsuokella multacida		2.05	0.19		1	1	
Blautia producta		0.31	-.03		99	99	
67777052	**Very good identification**						**67777052**
					GRA		
Actinomyces israelii		99.85	0.63	MLZ 13	99		
Mitsuokella multacida		0.16	0.03		1		
67777060	**Good identification**						**67777060**
					GRA		
Actinomyces israelii		95.70	0.54	NAG 6	99		
Mitsuokella multacida		3.53	0.17		1		
Actinomyces naeslundii		0.78	0.27		99		
67777062	**Very good identification**						**67777062**
Actinomyces israelii		99.46	0.57	NAG 6			
Actinomyces naeslundii		0.54	0.27				
67777070	**Acceptable identification**						**67777070**
					GRA		
Actinomyces israelii		92.41	0.40	MLZ 13	99		
Mitsuokella multacida		7.60	0.09	NAG 6	1		
67777072	**Good identification**						**67777072**
Actinomyces israelii		100.0	0.43	MLZ 13			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777100	**Good identification**						**67777100**
					GRA	SPO	
Actinomyces israelii		91.08	0.92		99	1	
Mitsuokella multacida		8.25	0.61		1	1	
Clostridium butyricum		0.59	0.33		99	99	
67777101	**Low discrimination**						**67777101**
					GRA		
Actinomyces israelii		47.11	0.58	URE 1	99		
Mitsuokella multacida		46.92	0.45		1		
Actinomyces naeslundii		5.67	0.51		99		
67777102	**Very good identification**						**67777102**
					GRA		
Actinomyces israelii		98.76	0.95		99		
Mitsuokella multacida		0.66	0.45		1		
Bifidobacterium adolescentis		0.32	0.46		99		
67777103	**Low discrimination**						**67777103**
					GRA		
Actinomyces israelii		86.42	0.61	URE 1	99		
Actinomyces naeslundii		6.93	0.51		99		
Mitsuokella multacida		6.38	0.29		1		
67777110	**Low discrimination**						**67777110**
					GRA		
Actinomyces israelii		82.65	0.78	MLZ 13	99		
Mitsuokella multacida		16.70	0.53		1		
Bifidobacterium dentium		0.25	0.10		99		
67777111	**Low discrimination**						**67777111**
					GRA		
Mitsuokella multacida		68.07	0.37	ARA 99	1		
Actinomyces israelii		30.64	0.43	URE 10	99		
Actinomyces naeslundii		1.30	0.29		99		
67777112	**Good identification**						**67777112**
Actinomyces israelii		93.63	0.81	MLZ 13			
Bifidobacterium adolescentis		2.44	0.47				
Bifidobacterium dentium		1.92	0.27				
67777113	**Low discrimination**						**67777113**
					GRA		
Actinomyces israelii		80.69	0.46	MLZ 13	99		
Mitsuokella multacida		13.28	0.20	URE 1	1		
Actinomyces naeslundii		2.28	0.29		99		
67777120	**Low discrimination**						**67777120**
					GRA		
Actinomyces israelii		71.89	0.71	NAG 6	99		
Mitsuokella multacida		25.49	0.51		1		
Actinomyces naeslundii		0.91	0.48		99		
67777122	**Good identification**						**67777122**
					SPO	GRA	
Actinomyces israelii		92.79	0.74	NAG 6	1	99	
Mitsuokella multacida		2.44	0.35		1	1	
Clostridium ramosum		1.98	0.30		92	50	
67777123	**Low discrimination**						**67777123**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		49.37	0.51	ESL 12	99		
Actinomyces israelii		39.28	0.40	XYL 1	99		
Mitsuokella multacida		11.35	0.18		1		
67777130	**Low discrimination**						**67777130**
					GRA		
Actinomyces israelii		50.95	0.57	MLZ 13	99		
Mitsuokella multacida		40.31	0.43	NAG 6	1		
Bifidobacterium dentium		4.31	0.15		99		
67777132	**Low discrimination**						**67777132**
					SPO		
Actinomyces israelii		54.88	0.60	MLZ 13	1		
Bifidobacterium dentium		31.32	0.32	NAG 6	1		
Clostridium tertium		8.22	0.32		99		
67777140	**Low discrimination**						**67777140**
					COC	GRA	
Actinomyces israelii		80.74	0.85		1	99	
Mitsuokella multacida		18.80	0.61		1	1	
Blautia producta		0.31	0.24		99	99	
67777141	**Low discrimination**						**67777141**
					GRA		
Mitsuokella multacida		70.83	0.45	ARA 99	1		
Actinomyces israelii		27.66	0.51	URE 10	99		
Actinomyces naeslundii		1.51	0.38		99		
67777142	**Very good identification**						**67777142**
					GRA		
Actinomyces israelii		98.08	0.88		99		
Mitsuokella multacida		1.69	0.45		1		
Bifidobacterium adolescentis		0.10	0.30		99		
67777143	**Low discrimination**						**67777143**
					GRA		
Actinomyces israelii		81.56	0.54	URE 1	99		
Mitsuokella multacida		15.47	0.29		1		
Actinomyces naeslundii		2.97	0.38		99		
67777150	**Low discrimination**						**67777150**
					COC	GRA	
Actinomyces israelii		62.59	0.71	MLZ 13	1	99	
Mitsuokella multacida		32.51	0.53		1	1	
Blautia producta		4.88	0.32		99	99	
67777151	**Low discrimination**						**67777151**
					COC	GRA	
Mitsuokella multacida		84.13	0.37	ARA 99	1	1	
Actinomyces israelii		14.73	0.36	URE 10	1	99	
Blautia producta		1.15	-.03		99	99	
67777152	**Good identification**						**67777152**
					GRA		
Actinomyces israelii		94.72	0.74	MLZ 13	99		
Mitsuokella multacida		3.64	0.37		1		
Bifidobacterium adolescentis		0.78	0.32		99		
67777153	**Low discrimination**						**67777153**
					GRA		
Actinomyces israelii		70.26	0.39	MLZ 13	99		
Mitsuokella multacida		29.74	0.20	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777160	**Low discrimination**						**67777160**
					COC	GRA	
Actinomyces israelii		51.86	0.64	NAG 6	1	99	
Mitsuokella multacida		47.30	0.51		1	1	
Blautia producta		0.35	0.07		99	99	
67777162	**Good identification**						**67777162**
					GRA		
Actinomyces israelii		92.35	0.67	NAG 6	99		
Mitsuokella multacida		6.24	0.35		1		
Bifidobacterium breve		1.05	0.29		99		
67777170	**Low discrimination**						**67777170**
					COC	GRA	
Mitsuokella multacida		63.95	0.43	ARA 99	1	1	
Actinomyces israelii		31.43	0.50		1	99	
Blautia producta		4.26	0.15		99	99	
67777172	**Low discrimination**						**67777172**
					GRA		
Actinomyces israelii		79.88	0.53	MLZ 13	99		
Mitsuokella multacida		12.04	0.26	NAG 6	1		
Bifidobacterium breve		6.10	0.29		99		
67777200	**Good identification**						**67777200**
					GRA		
Actinomyces israelii		91.72	0.62	RHA 7	99		
Mitsuokella multacida		8.28	0.32		1		
67777202	**Very good identification**						**67777202**
					GRA		
Actinomyces israelii		99.34	0.65	RHA 7	99		
Mitsuokella multacida		0.66	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777210	**Low discrimination**						**67777210**
					GRA		
Actinomyces israelii		83.24	0.48	RHA 7	99		
Mitsuokella multacida		16.76	0.24	MLZ 13	1		
67777212	**Good identification**						**67777212**
					GRA		
Actinomyces israelii		98.53	0.51	RHA 7	99		
Mitsuokella multacida		1.47	0.07	MLZ 13	1		
67777220	**Low discrimination**						**67777220**
					GRA		
Actinomyces israelii		73.88	0.42	RHA 7	99		
Mitsuokella multacida		26.12	0.21	NAG 6	1		
67777222	**Good identification**						**67777222**
					SPO	GRA	
Actinomyces israelii		96.03	0.45	RHA 7	1	99	
Mitsuokella multacida		2.52	0.05	NAG 6	1	1	
Clostridium ramosum		1.45	-.03		92	50	
67777240	**Low discrimination**						**67777240**
					GRA		
Actinomyces israelii		81.16	0.55	RHA 7	99		
Mitsuokella multacida		18.84	0.32		1		
67777242	**Very good identification**						**67777242**
					GRA		
Actinomyces israelii		98.31	0.58	RHA 7	99		
Mitsuokella multacida		1.69	0.15		1		
67777250	**Low discrimination**						**67777250**
					COC	GRA	
Actinomyces israelii		64.12	0.41	RHA 7	1	99	
Mitsuokella multacida		33.20	0.24	MLZ 13	1	1	
Blautia producta		2.68	-.03		99	99	
67777252	**Good identification**						**67777252**
					GRA		
Actinomyces israelii		96.31	0.44	RHA 7	99		
Mitsuokella multacida		3.69	0.07	MLZ 13	1		
67777262	**Acceptable identification**						**67777262**
					GRA		
Actinomyces israelii		93.69	0.38	RHA 7	99		
Mitsuokella multacida		6.31	0.05	NAG 6	1		
67777300	**Low discrimination**						**67777300**
					GRA	SPO	
Mitsuokella multacida		69.02	0.66	ARA 99	1	1	
Actinomyces israelii		30.90	0.73		99	1	
Clostridium butyricum		0.08	0.07		99	99	
67777301	**Good identification**						**67777301**
					GRA		
Mitsuokella multacida		96.09	0.50	ARA 99	1		
Actinomyces israelii		3.91	0.38	URE 10	99		
67777302	**Low discrimination**						**67777302**
					GRA	SPO	
Actinomyces israelii		85.70	0.76	RHA 7	99	1	
Mitsuokella multacida		14.18	0.50		1	1	
Clostridium butyricum		0.06	0.00		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777310	**Low discrimination**						**67777310**
					COC	GRA	
Mitsuokella multacida		83.23	0.58	ARA 99	1	1	
Actinomyces israelii		16.70	0.58		1	99	
Blautia producta		0.07	-.03		99	99	
67777311	**Good identification**						**67777311**
					GRA		
Mitsuokella multacida		98.21	0.41	ARA 99	1		
Actinomyces israelii		1.79	0.24	URE 10	99		
67777312	**Low discrimination**						**67777312**
					GRA		
Actinomyces israelii		72.49	0.61	RHA 7	99		
Mitsuokella multacida		26.76	0.41	MLZ 13	1		
Clostridium clostridioforme		0.30	0.11		50		
67777320	**Low discrimination**						**67777320**
					SPO	GRA	
Mitsuokella multacida		88.47	0.56	ARA 99	1	1	
Actinomyces israelii		10.11	0.52		1	99	
Clostridium ramosum		1.42	0.21		92	50	
67777321	**Good identification**						**67777321**
					GRA		
Mitsuokella multacida		98.97	0.39	ARA 99	1		
Actinomyces israelii		1.03	0.18	URE 10	99		
67777322	**Low discrimination**						**67777322**
					SPO	GRA	
Actinomyces israelii		48.06	0.55	RHA 7	1	99	
Mitsuokella multacida		31.15	0.39	NAG 6	1	1	
Clostridium ramosum		18.01	0.32		92	50	
67777330	**Good identification**						**67777330**
					GRA		
Mitsuokella multacida		95.13	0.47	ARA 99	1		
Actinomyces israelii		4.87	0.38		99		
67777340	**Low discrimination**						**67777340**
					COC	GRA	
Mitsuokella multacida		84.53	0.66	ARA 99	1	1	
Actinomyces israelii		14.71	0.66		1	99	
Blautia producta		0.76	0.24		99	99	
67777341	**Good identification**						**67777341**
					GRA		
Mitsuokella multacida		98.44	0.50	ARA 99	1		
Actinomyces israelii		1.56	0.31	URE 10	99		
67777342	**Low discrimination**						**67777342**
					COC	GRA	
Actinomyces israelii		69.96	0.69	RHA 7	1	99	
Mitsuokella multacida		29.77	0.50		1	1	
Blautia producta		0.27	0.07		99	99	
67777350	**Low discrimination**						**67777350**
					COC	GRA	
Mitsuokella multacida		86.30	0.58	ARA 99	1	1	
Blautia producta		6.97	0.32		99	99	
Actinomyces israelii		6.73	0.51		1	99	
67777351	**Good identification**						**67777351**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Mitsuokella multacida		98.58	0.41	ARA 99	1	1	
Blautia producta		0.72	-.03	URE 10	99	99	
Actinomyces israelii		0.70	0.17		1	99	
67777352	**Low discrimination**						**67777352**
					COC	GRA	
Actinomyces israelii		49.36	0.54	RHA 7	1	99	
Mitsuokella multacida		46.86	0.41	MLZ 13	1	1	
Blautia producta		3.79	0.15		99	99	
67777360	**Good identification**						**67777360**
					COC	GRA	
Mitsuokella multacida		95.38	0.56	ARA 99	1	1	
Actinomyces israelii		4.24	0.45		1	99	
Blautia producta		0.38	0.07		99	99	
67777361	**Good identification**						**67777361**
Mitsuokella multacida		100.0	0.39	ARA 99			
67777362	**Low discrimination**						**67777362**
					SPO	GRA	
Mitsuokella multacida		62.27	0.39	ARA 99	1	1	
Actinomyces israelii		37.36	0.48	BGL 10	1	99	
Clostridium ramosum		0.36	-.03		92	50	
67777370	**Acceptable identification**						**67777370**
					COC	GRA	
Mitsuokella multacida		94.71	0.47	ARA 99	1	1	
Blautia producta		3.40	0.15		99	99	
Actinomyces israelii		1.89	0.31		1	99	
67777400	**Low discrimination**						**67777400**
					GRA	SPO	
Actinomyces israelii		68.75	0.80		99	1	
Mitsuokella multacida		30.44	0.61		1	1	
Clostridium butyricum		0.81	0.26		99	99	
67777401	**Low discrimination**						**67777401**
					GRA		
Mitsuokella multacida		82.96	0.45	LAC 99	1		
Actinomyces israelii		17.04	0.46	URE 10	99		
67777402	**Very good identification**						**67777402**
					GRA	SPO	
Actinomyces israelii		96.43	0.83		99	1	
Mitsuokella multacida		3.16	0.45		1	1	
Clostridium butyricum		0.30	0.19		99	99	
67777403	**Low discrimination**						**67777403**
					GRA		
Actinomyces israelii		73.49	0.49	URE 1	99		
Mitsuokella multacida		26.51	0.29		1		
67777410	**Low discrimination**						**67777410**
					GRA		
Actinomyces israelii		49.80	0.66	MLZ 13	99		
Mitsuokella multacida		49.18	0.53		1		
Bifidobacterium dentium		0.74	0.10		99		
67777411	**Acceptable identification**						**67777411**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		91.57	0.37	LAC 99	1		
Actinomyces israelii		8.43	0.32	URE 10	99		
67777412	**Low discrimination**						**67777412**
					GRA		
Actinomyces israelii		84.46	0.69	MLZ 13	99		
Bifidobacterium dentium		8.46	0.27		99		
Mitsuokella multacida		6.18	0.37		1		
67777420	**Low discrimination**						**67777420**
					GRA	SPO	
Mitsuokella multacida		63.05	0.51	LAC 99	1	1	
Actinomyces israelii		36.37	0.60		99	1	
Clostridium butyricum		0.59	0.08		99	99	
67777422	**Low discrimination**						**67777422**
					GRA	SPO	
Actinomyces israelii		87.97	0.63	NAG 6	99	1	
Mitsuokella multacida		11.30	0.35		1	1	
Clostridium butyricum		0.37	0.01		99	99	
67777430	**Low discrimination**						**67777430**
					GRA		
Mitsuokella multacida		73.23	0.43	LAC 99	1		
Actinomyces israelii		18.93	0.45		99		
Bifidobacterium dentium		7.84	0.15		99		
67777440	**Low discrimination**						**67777440**
					COC	GRA	
Mitsuokella multacida		52.70	0.61	LAC 99	1	1	
Actinomyces israelii		46.30	0.73		1	99	
Blautia producta		0.88	0.24		99	99	
67777441	**Acceptable identification**						**67777441**
					GRA		
Mitsuokella multacida		92.60	0.45	LAC 99	1		
Actinomyces israelii		7.40	0.39	URE 10	99		
67777442	**Good identification**						**67777442**
					COC	GRA	
Actinomyces israelii		92.02	0.76		1	99	
Mitsuokella multacida		7.76	0.45		1	1	
Blautia producta		0.13	0.07		99	99	
67777443	**Low discrimination**						**67777443**
					GRA		
Actinomyces israelii		51.88	0.42	URE 1	99		
Mitsuokella multacida		48.12	0.29		1		
67777450	**Low discrimination**						**67777450**
					COC	GRA	
Mitsuokella multacida		64.77	0.53	LAC 99	1	1	
Actinomyces israelii		25.51	0.59		1	99	
Blautia producta		9.72	0.32		99	99	
67777451	**Good identification**						**67777451**
					COC	GRA	
Mitsuokella multacida		95.29	0.37	LAC 99	1	1	
Actinomyces israelii		3.41	0.25	URE 10	1	99	
Blautia producta		1.30	-.03		99	99	
67777452	**Low discrimination**						**67777452**
					COC	GRA	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		81.86	0.62	MLZ 13	1	99	
Mitsuokella multacida		15.40	0.37		1	1	
Blautia producta		2.31	0.15		99	99	
67777460	**Low discrimination**						**67777460**
					COC	GRA	
Mitsuokella multacida		81.19	0.51	LAC 99	1	1	
Actinomyces israelii		18.21	0.52		1	99	
Blautia producta		0.60	0.07		99	99	
67777462	**Low discrimination**						**67777462**
					GRA		
Actinomyces israelii		75.17	0.56	NAG 6	99		
Mitsuokella multacida		24.83	0.35		1		
67777470	**Low discrimination**						**67777470**
					COC	GRA	
Mitsuokella multacida		85.67	0.43	LAC 99	1	1	
Actinomyces israelii		8.61	0.38		1	99	
Blautia producta		5.71	0.15		99	99	
67777472	**Low discrimination**						**67777472**
					GRA		
Actinomyces israelii		53.81	0.41	MLZ 13	99		
Mitsuokella multacida		39.64	0.26	NAG 6	1		
Bifidobacterium dentium		3.90	-.03		99		
67777500	**Good identification**						**67777500**
					GRA	SPO	
Mitsuokella multacida		91.08	0.95		1	1	
Actinomyces israelii		8.31	0.90		99	1	
Clostridium butyricum		0.59	0.50		99	99	
67777501	**Very good identification**						**67777501**
					GRA	SPO	
Mitsuokella multacida		99.12	0.79	URE 10	1	1	
Actinomyces israelii		0.82	0.56		99	1	
Clostridium butyricum		0.06	0.15		99	99	
67777502	**Low discrimination**						**67777502**
					GRA		
Actinomyces israelii		53.77	0.93		99		
Mitsuokella multacida		43.64	0.79		1		
Bifidobacterium adolescentis		1.40	0.60		99		
67777503	**Low discrimination**						**67777503**
					GRA		
Mitsuokella multacida		89.49	0.63	URE 10	1		
Actinomyces israelii		10.03	0.59	BGL 10	99		
Bifidobacterium adolescentis		0.26	0.26		99		
67777510	**Good identification**						**67777510**
					GRA		
Mitsuokella multacida		94.46	0.87		1		
Actinomyces israelii		3.87	0.76		99		
Bifidobacterium dentium		1.42	0.44		99		
67777511	**Very good identification**						**67777511**
					GRA		
Mitsuokella multacida		99.47	0.71	URE 10	1		
Actinomyces israelii		0.37	0.42		99		
Bifidobacterium dentium		0.14	0.10		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777512	**Low discrimination**						**67777512**
					GRA		
Bifidobacterium dentium		44.97	0.62	NIT 1	99		
Mitsuokella multacida		32.84	0.71		1		
Actinomyces israelii		18.14	0.79		99		
67777513	**Low discrimination**						**67777513**
					GRA		
Mitsuokella multacida		84.36	0.55	URE 10	1		
Bifidobacterium dentium		10.50	0.27	BGL 10	99		
Actinomyces israelii		4.24	0.45		99		
67777520	**Very good identification**						**67777520**
					GRA	SPO	
Mitsuokella multacida		97.47	0.85		1	1	
Actinomyces israelii		2.27	0.70		99	1	
Clostridium butyricum		0.22	0.31		99	99	
67777521	**Very good identification**						**67777521**
					GRA	SPO	
Mitsuokella multacida		99.77	0.69	URE 10	1	1	
Actinomyces israelii		0.21	0.36		99	1	
Clostridium butyricum		0.02	-.03		99	99	
67777522	**Low discrimination**						**67777522**
					GRA		
Mitsuokella multacida		73.65	0.69	BGL 10	1		
Actinomyces israelii		23.17	0.73		99		
Bifidobacterium dentium		2.42	0.32		99		
67777523	**Good identification**						**67777523**
					GRA		
Mitsuokella multacida		96.94	0.52	URE 10	1		
Actinomyces israelii		2.77	0.39	BGL 10	99		
Bifidobacterium dentium		0.29	-.03		99		
67777530	**Low discrimination**						**67777530**
					GRA		
Mitsuokella multacida		89.41	0.77		1		
Bifidobacterium dentium		9.57	0.49		99		
Actinomyces israelii		0.93	0.56		99		
67777531	**Very good identification**						**67777531**
					GRA		
Mitsuokella multacida		98.94	0.61	URE 10	1		
Bifidobacterium dentium		0.96	0.15		99		
Actinomyces israelii		0.09	0.22		99		
67777532	**Low discrimination**						**67777532**
					GRA		
Bifidobacterium dentium		89.42	0.66	NIT 1	99		
Mitsuokella multacida		9.18	0.61		1		
Actinomyces israelii		1.30	0.59		99		
67777533	**Low discrimination**						**67777533**
					GRA		
Mitsuokella multacida		52.68	0.44	URE 10	1		
Bifidobacterium dentium		46.64	0.32	BGL 10	99		
Actinomyces israelii		0.68	0.25		99		
67777540	**Good identification**						**67777540**
					COC	GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		94.99	0.95		1	1	
Actinomyces israelii		3.37	0.83		1	99	
Blautia producta		1.58	0.58		99	99	
67777541	**Very good identification**						**67777541**
					COC	GRA	
Mitsuokella multacida		99.52	0.79	URE 10	1	1	
Actinomyces israelii		0.32	0.49		1	99	
Blautia producta		0.15	0.24		99	99	
67777542	**Low discrimination**						**67777542**
					COC	GRA	
Mitsuokella multacida		66.59	0.79	BGL 10	1	1	
Actinomyces israelii		31.91	0.86		1	99	
Blautia producta		1.11	0.41		99	99	
67777543	**Very good identification**						**67777543**
					COC	GRA	
Mitsuokella multacida		95.65	0.63	URE 10	1	1	
Actinomyces israelii		4.17	0.52	BGL 10	1	99	
Blautia producta		0.15	0.07		99	99	
67777550	**Low discrimination**						**67777550**
					COC	GRA	
Mitsuokella multacida		85.75	0.87		1	1	
Blautia producta		12.86	0.66		99	99	
Actinomyces israelii		1.36	0.69		1	99	
67777551	**Very good identification**						**67777551**
					COC	GRA	
Mitsuokella multacida		98.51	0.71	URE 10	1	1	
Blautia producta		1.34	0.32		99	99	
Actinomyces israelii		0.14	0.35		1	99	
67777552	**Low discrimination**						**67777552**
					COC	GRA	
Mitsuokella multacida		71.76	0.71	BGL 10	1	1	
Actinomyces israelii		15.42	0.72		1	99	
Blautia producta		10.77	0.50		99	99	
67777553	**Good identification**						**67777553**
					COC	GRA	
Mitsuokella multacida		96.55	0.55	URE 10	1	1	
Actinomyces israelii		1.89	0.38	BGL 10	1	99	
Blautia producta		1.32	0.15		99	99	
67777560	**Very good identification**						**67777560**
					COC	GRA	
Mitsuokella multacida		98.36	0.85		1	1	
Actinomyces israelii		0.89	0.63		1	99	
Blautia producta		0.73	0.41		99	99	
67777561	**Very good identification**						**67777561**
					COC	GRA	
Mitsuokella multacida		99.85	0.69	URE 10	1	1	
Actinomyces israelii		0.08	0.29		1	99	
Blautia producta		0.07	0.07		99	99	
67777562	**Low discrimination**						**67777562**
					COC	GRA	
Mitsuokella multacida		88.42	0.69	BGL 10	1	1	
Actinomyces israelii		10.82	0.66		1	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		0.66	0.25		99	99	
67777563	**Good identification**						**67777563**
					GRA		
Mitsuokella multacida		98.90	0.52	URE 10	1		
Actinomyces israelii		1.10	0.32	BGL 10	99		
67777570	**Good identification**						**67777570**
					COC	GRA	
Mitsuokella multacida		93.30	0.77		1	1	
Blautia producta		6.22	0.50		99	99	
Actinomyces israelii		0.38	0.49		1	99	
67777571	**Very good identification**						**67777571**
					COC	GRA	
Mitsuokella multacida		99.36	0.61	URE 10	1	1	
Blautia producta		0.60	0.15		99	99	
Actinomyces israelii		0.04	0.15		1	99	
67777572	**Low discrimination**						**67777572**
					COC	GRA	
Mitsuokella multacida		81.94	0.61	BGL 10	1	1	
Bifidobacterium dentium		8.06	0.32		1	99	
Blautia producta		5.46	0.33		99	99	
67777573	**Good identification**						**67777573**
					COC	GRA	
Mitsuokella multacida		98.04	0.44	URE 10	1	1	
Bifidobacterium dentium		0.88	-.03	BGL 10	1	99	
Blautia producta		0.59	-.01		99	99	
67777600	**Good identification**						**67777600**
					GRA	SPO	
Mitsuokella multacida		91.58	0.66	LAC 99	1	1	
Actinomyces israelii		8.38	0.61		99	1	
Clostridium butyricum		0.04	0.00		99	99	
67777601	**Good identification**						**67777601**
					GRA		
Mitsuokella multacida		99.18	0.50	LAC 99	1		
Actinomyces israelii		0.83	0.27	URE 10	99		
67777602	**Low discrimination**						**67777602**
					GRA		
Actinomyces israelii		55.14	0.64	RHA 7	99		
Mitsuokella multacida		44.61	0.50		1		
Clostridium clostridioforme		0.17	0.11		50		
67777610	**Very good identification**						**67777610**
					COC	GRA	
Mitsuokella multacida		95.98	0.58	LAC 99	1	1	
Actinomyces israelii		3.94	0.47		1	99	
Blautia producta		0.08	-.03		99	99	
67777611	**Good identification**						**67777611**
Mitsuokella multacida		100.0	0.41	LAC 99			
67777612	**Low discrimination**						**67777612**
					GRA		
Mitsuokella multacida		61.94	0.41	LAC 99	1		
Actinomyces israelii		34.32	0.50	BGL 10	99		
Clostridium clostridioforme		3.28	0.23		50		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
67777620	**Good identification**						**67777620**
					GRA		
Mitsuokella multacida		97.72	0.56	LAC 99	1		
Actinomyces israelii		2.28	0.40		99		
67777621	**Good identification**						**67777621**
Mitsuokella multacida		100.0	0.39	LAC 99			
67777622	**Low discrimination**						**67777622**
					GRA		
Mitsuokella multacida		74.58	0.39	LAC 99	1		
Actinomyces israelii		23.54	0.43	BGL 10	99		
Clostridium clostridioforme		1.16	0.11		50		
67777630	**Good identification**						**67777630**
					GRA		
Mitsuokella multacida		98.96	0.47	LAC 99	1		
Actinomyces israelii		1.04	0.26		99		
67777640	**Very good identification**						**67777640**
					COC	GRA	
Mitsuokella multacida		95.73	0.66	LAC 99	1	1	
Actinomyces israelii		3.41	0.54		1	99	
Blautia producta		0.86	0.24		99	99	
67777641	**Good identification**						**67777641**
					GRA		
Mitsuokella multacida		99.68	0.50	LAC 99	1		
Actinomyces israelii		0.32	0.20	URE 10	99		
67777642	**Low discrimination**						**67777642**
					COC	GRA	
Mitsuokella multacida		67.13	0.50	LAC 99	1	1	
Actinomyces israelii		32.27	0.57	BGL 10	1	99	
Blautia producta		0.60	0.07		99	99	
67777650	**Good identification**						**67777650**
					COC	GRA	
Mitsuokella multacida		91.18	0.58	LAC 99	1	1	
Blautia producta		7.37	0.32		99	99	
Actinomyces israelii		1.46	0.40		1	99	
67777651	**Good identification**						**67777651**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	LAC 99	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
67777652	**Low discrimination**						**67777652**
					COC	GRA	
Mitsuokella multacida		77.15	0.41	LAC 99	1	1	
Actinomyces israelii		16.62	0.43	BGL 10	1	99	
Blautia producta		6.23	0.15		99	99	
67777660	**Good identification**						**67777660**
					COC	GRA	
Mitsuokella multacida		98.71	0.56	LAC 99	1	1	
Actinomyces israelii		0.90	0.33		1	99	
Blautia producta		0.39	0.07		99	99	
67777661	**Good identification**						**67777661**
Mitsuokella multacida		100.0	0.39	LAC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777662	**Low discrimination**						**67777662**
					GRA		
Mitsuokella multacida		89.07	0.39	LAC 99	1		
Actinomyces israelii		10.93	0.36	BGL 10	99		
67777670	**Good identification**						**67777670**
					COC	GRA	
Mitsuokella multacida		96.16	0.47	LAC 99	1	1	
Blautia producta		3.45	0.15		99	99	
Actinomyces israelii		0.39	0.19		1	99	
67777700	**Excellent identification**						**67777700**
					GRA	SPO	
Mitsuokella multacida		99.61	1.00		1	1	
Actinomyces israelii		0.37	0.71		99	1	
Clostridium butyricum		0.01	0.24		99	99	
67777701	**Very good identification**						**67777701**
					GRA		
Mitsuokella multacida		99.97	0.84	URE 10	1		
Actinomyces israelii		0.03	0.37		99		
67777702	**Very good identification**						**67777702**
					GRA	SPO	
Mitsuokella multacida		95.17	0.84	BGL 10	1	1	
Actinomyces israelii		4.75	0.74		99	1	
Clostridium butyricum		0.04	0.17		99	99	
67777703	**Very good identification**						**67777703**
					GRA		
Mitsuokella multacida		99.55	0.67	URE 10	1		
Actinomyces israelii		0.45	0.40	BGL 10	99		
67777710	**Excellent identification**						**67777710**
					COC	GRA	
Mitsuokella multacida		99.74	0.92		1	1	
Actinomyces israelii		0.17	0.57		1	99	
Blautia producta		0.08	0.32		99	99	
67777711	**Very good identification**						**67777711**
					COC	GRA	
Mitsuokella multacida		99.98	0.76	URE 10	1	1	
Actinomyces israelii		0.02	0.23		1	99	
Blautia producta		0.01	-.03		99	99	
67777712	**Very good identification**						**67777712**
					GRA		
Mitsuokella multacida		96.79	0.76	BGL 10	1		
Actinomyces israelii		2.17	0.60		99		
Bifidobacterium dentium		0.72	0.27		99		
67777713	**Very good identification**						**67777713**
					GRA		
Mitsuokella multacida		99.73	0.59	URE 10	1		
Actinomyces israelii		0.20	0.26	BGL 10	99		
Bifidobacterium dentium		0.07	-.07		99		
67777720	**Very good identification**						**67777720**
					COC	GRA	
Mitsuokella multacida		99.90	0.90		1	1	
Actinomyces israelii		0.09	0.51		1	99	
Blautia producta		0.00	0.07		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777721	**Very good identification**						**67777721**
					GRA		
Mitsuokella multacida		99.99	0.73	URE 10	1		
Actinomyces israelii		0.01	0.16		99		
67777722	**Very good identification**						**67777722**
					GRA		
Mitsuokella multacida		97.68	0.73	BGL 10	1		
Actinomyces israelii		1.25	0.54		99		
Bacteroides ovatus		0.95	0.32		1		
67777723	**Very good identification**						**67777723**
					GRA		
Mitsuokella multacida		99.80	0.57	URE 10	1		
Actinomyces israelii		0.12	0.19	BGL 10	99		
Bacteroides ovatus		0.09	-.02		1		
67777730	**Very good identification**						**67777730**
					GRA		
Mitsuokella multacida		99.86	0.82		1		
Bifidobacterium dentium		0.06	0.15		99		
Actinomyces israelii		0.04	0.36		99		
67777731	**Very good identification**						**67777731**
Mitsuokella multacida		100.0	0.65	URE 10			
67777732	**Good identification**						**67777732**
					GRA		
Mitsuokella multacida		92.06	0.65	BGL 10	1		
Bifidobacterium dentium		4.88	0.32		99		
Bacteroides ovatus		1.79	0.29		1		
67777733	**Good identification**						**67777733**
					GRA		
Mitsuokella multacida		99.35	0.49	URE 10	1		
Bifidobacterium dentium		0.48	-.03	BGL 10	99		
Bacteroides ovatus		0.18	-.05		1		
67777740	**Very good identification**						**67777740**
					COC	GRA	
Mitsuokella multacida		98.97	1.00		1	1	
Blautia producta		0.89	0.58		99	99	
Actinomyces israelii		0.14	0.64		1	99	
67777741	**Very good identification**						**67777741**
					COC	GRA	
Mitsuokella multacida		99.91	0.84	URE 10	1	1	
Blautia producta		0.08	0.24		99	99	
Actinomyces israelii		0.01	0.30		1	99	
67777742	**Very good identification**						**67777742**
					COC	GRA	
Mitsuokella multacida		97.23	0.84	BGL 10	1	1	
Actinomyces israelii		1.89	0.67		1	99	
Blautia producta		0.87	0.41		99	99	
67777743	**Very good identification**						**67777743**
					COC	GRA	
Mitsuokella multacida		99.74	0.67	URE 10	1	1	
Actinomyces israelii		0.18	0.33	BGL 10	1	99	
Blautia producta		0.08	0.07		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
67777750	**Good identification**						**67777750**
					COC	GRA	
Mitsuokella multacida		92.47	0.92		1	1	
Blautia producta		7.47	0.66		99	99	
Actinomyces israelii		0.06	0.50		1	99	
67777751	**Very good identification**						**67777751**
					COC	GRA	
Mitsuokella multacida		99.27	0.76	URE 10	1	1	
Blautia producta		0.73	0.32		99	99	
Actinomyces israelii		0.01	0.16		1	99	
67777752	**Good identification**						**67777752**
					COC	GRA	
Mitsuokella multacida		91.77	0.76	BGL 10	1	1	
Blautia producta		7.41	0.50		99	99	
Actinomyces israelii		0.80	0.53		1	99	
67777753	**Very good identification**						**67777753**
					COC	GRA	
Mitsuokella multacida		99.19	0.59	URE 10	1	1	
Blautia producta		0.73	0.15	BGL 10	99	99	
Actinomyces israelii		0.08	0.19		1	99	
67777760	**Very good identification**						**67777760**
					COC	GRA	
Mitsuokella multacida		99.57	0.90		1	1	
Blautia producta		0.40	0.41		99	99	
Actinomyces israelii		0.04	0.44		1	99	
67777761	**Very good identification**						**67777761**
					COC	GRA	
Mitsuokella multacida		99.96	0.73	URE 10	1	1	
Blautia producta		0.04	0.07		99	99	
67777762	**Very good identification**						**67777762**
					COC	GRA	
Mitsuokella multacida		99.07	0.73	BGL 10	1	1	
Actinomyces israelii		0.49	0.47		1	99	
Blautia producta		0.40	0.25		99	99	
67777763	**Very good identification**						**67777763**
Mitsuokella multacida		100.0	0.57	URE 10			
67777770	**Very good identification**						**67777770**
					COC	GRA	
Mitsuokella multacida		96.52	0.82		1	1	
Blautia producta		3.47	0.50		99	99	
Actinomyces israelii		0.02	0.29		1	99	
67777771	**Very good identification**						**67777771**
					COC	GRA	
Mitsuokella multacida		99.68	0.65	URE 10	1	1	
Blautia producta		0.33	0.15		99	99	
67777772	**Very good identification**						**67777772**
					COC	GRA	
Mitsuokella multacida		96.20	0.65	BGL 10	1	1	
Blautia producta		3.45	0.33		99	99	
Actinomyces israelii		0.21	0.32		1	99	
67777773	**Good identification**						**67777773**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COC	GRA		
Mitsuokella multacida		99.68	0.49	URE 10	1	1		
Blautia producta		0.33	-.01	BGL 10	99	99		
70000000	**Low discrimination**							**70000000**
					COC	GRA		
Peptoniphilus asaccharolyticus		35.21	0.40	NIT 3	99	99		
Veillonella parvula		11.74	0.32	ESL 1	99	1		
Fusobacterium nucleatum subsp. nucleatum		11.50	0.32		1	1		
70100000	**Acceptable identification**							**70100000**
					COC	SPO		
Clostridium bifermentans		90.62	0.46	NIT 1	1	99		
Clostridium sporogenes		2.87	0.19	GLU 92	1	99		
Peptoniphilus asaccharolyticus		2.07	0.06		99	1		
71000000	**Good identification**							**71000000**
Clostridium bifermentans		93.19	0.58	NIT 1				
Clostridium botulinum biovar A		2.49	0.32					
Clostridium sporogenes		1.83	0.27					
71000020	**Low discrimination**							**71000020**
					SPO	CAT		
Clostridium bifermentans		56.43	0.38	NIT 1	99	1		
Clostridium cadaveris		33.98	0.32	NAG 6	99	1		
Propionibacterium acnes		8.14	0.24		1	95		
71001520	**Good identification**							**71001520**
Odoribacter splanchnicus		100.0	0.51	MNS 99				
71010000	**Low discrimination**							**71010000**
					GRA	SPO	CAT	
Clostridium bifermentans		51.66	0.36	NIT 1	99	99	1	
Fusobacterium varium		23.60	0.25	FRU 5	1	1	1	
Propionibacterium acnes		9.04	0.24		99	1	95	
71010020	**Very good identification**							**71010020**
					SPO	CAT		
Propionibacterium acnes		95.49	0.58	ESL 1	1	95		
Clostridium cadaveris		4.03	0.32		99	1		
Clostridium bifermentans		0.35	0.16		99	1		
71010060	**Good identification**							**71010060**
					SPO	CAT		
Propionibacterium acnes		99.81	0.47	ESL 1	1	95		
Clostridium cadaveris		0.19	-.03		99	1		
71011020	**Very good identification**							**71011020**
					SPO	CAT		
Propionibacterium acnes		99.91	0.58	ESL 1	1	95		
Atopobium minutum		0.05	0.09		1	1		
Clostridium cadaveris		0.04	-.03		99	1		
71011060	**Good identification**							**71011060**
Propionibacterium acnes		100.0	0.47	ESL 1				
71011520	**Good identification**							**71011520**
Odoribacter splanchnicus		100.0	0.56	MNS 99				
71011560	**Good identification**							**71011560**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Odoribacter splanchnicus		100.0	0.39	MNS 99			
71012020	**Good identification**						**71012020**
					SPO	CAT	
Propionibacterium acnes		99.86	0.54	ESL 1	1	95	
Clostridium cadaveris		0.08	-.03		99	1	
Eubacterium limosum		0.06	-.06		1	1	
71012060	**Good identification**						**71012060**
Propionibacterium acnes		100.0	0.43	ESL 1			
71013020	**Good identification**						**71013020**
Propionibacterium acnes		100.0	0.54	ESL 1			
71013060	**Good identification**						**71013060**
Propionibacterium acnes		100.0	0.43	ESL 1			
71100000	**Very good identification**						**71100000**
Clostridium bifermentans		97.40	0.64	NIT 1			
Clostridium sporogenes		1.52	0.32				
Clostridium botulinum biovar A		0.91	0.30				
71100001	**Acceptable identification**						**71100001**
Clostridium sordellii		91.61	0.44	NIT 1			
Clostridium bifermentans		7.94	0.30	ESL 5			
Clostridium novyi biovar A		0.33	0.19				
71100020	**Good identification**						**71100020**
Clostridium bifermentans		98.42	0.44	NIT 1			
Clostridium cadaveris		1.34	0.10	NAG 6			
Clostridium sporogenes		0.24	-.03				
71100040	**Acceptable identification**						**71100040**
Clostridium bifermentans		92.81	0.41	NIT 1			
Clostridium botulinum biovar A		6.23	0.21	SOR 4			
Clostridium novyi biovar A		0.61	0.16				
71110000	**Acceptable identification**						**71110000**
Clostridium bifermentans		90.44	0.43	NIT 1			
Clostridium sporogenes		4.74	0.19	FRU 5			
Clostridium sordellii		2.02	0.10				
71110001	**Good identification**						**71110001**
Clostridium sordellii		99.55	0.44	NIT 1			
Clostridium bifermentans		0.45	0.08	ESL 5			
71200000	**Good identification**						**71200000**
					SPO		
Clostridium bifermentans		99.68	0.42	NIT 1	99		
Eubacterium tortuosum		0.32	-.04	SAL 10	1		
71300000	**Good identification**						**71300000**
Clostridium bifermentans		99.86	0.48	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Clostridium sporogenes		0.14	-.03	SAL 10		
73001010	**Good identification**					**73001010**
Eubacterium saburreum		97.89	0.40	NIT 1		
Eubacterium tortuosum		2.11	-.04			
73001100	**Good identification**					**73001100**
Eubacterium saburreum		99.25	0.47	NIT 1		
Eubacterium tortuosum		0.75	-.04			
73001110	**Good identification**					**73001110**
Eubacterium saburreum		100.0	0.46	NIT 1		
73005000	**Good identification**					**73005000**
Eubacterium saburreum		96.83	0.37	NIT 1		
Eubacterium tortuosum		3.17	-.04			
73005010	**Good identification**					**73005010**
Eubacterium saburreum		100.0	0.35	NIT 1		
73005100	**Good identification**					**73005100**
Eubacterium saburreum		100.0	0.43	NIT 1		
73005110	**Good identification**					**73005110**
Eubacterium saburreum		100.0	0.41	NIT 1		
73011000	**Good identification**					**73011000**
Eubacterium saburreum		99.42	0.49	NIT 1		
Eubacterium tortuosum		0.58	-.04			
73011002	**Good identification**					**73011002**
Eubacterium saburreum		100.0	0.36	NIT 1		
73011010	**Good identification**					**73011010**
Eubacterium saburreum		100.0	0.48	NIT 1		
73011020	**Low discrimination**					**73011020**
					CAT	
Eubacterium saburreum		53.77	0.36	NIT 1	1	
Propionibacterium acnes		46.23	0.24		95	
73011100	**Good identification**					**73011100**
Eubacterium saburreum		99.56	0.56	NIT 1		
Fusobacterium mortiferum		0.44	0.17			
73011102	**Good identification**					**73011102**
Eubacterium saburreum		100.0	0.43	NIT 1		
73011110	**Good identification**					**73011110**
Eubacterium saburreum		100.0	0.54	NIT 1		
73011112	**Good identification**					**73011112**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.41	NIT 1		
73011120	**Good identification**					**73011120**
Eubacterium saburreum		100.0	0.43	NIT 1		
73011130	**Good identification**					**73011130**
Eubacterium saburreum		100.0	0.41	NIT 1		
73015000	**Good identification**					**73015000**
Eubacterium saburreum		100.0	0.45	NIT 1		
73015010	**Good identification**					**73015010**
Eubacterium saburreum		100.0	0.43	NIT 1		
73015100	**Good identification**					**73015100**
Eubacterium saburreum		100.0	0.51	NIT 1		
73015102	**Good identification**					**73015102**
Eubacterium saburreum		100.0	0.38	NIT 1		
73015110	**Good identification**					**73015110**
Eubacterium saburreum		100.0	0.49	NIT 1		
73015112	**Good identification**					**73015112**
Eubacterium saburreum		100.0	0.37	NIT 1		
73015120	**Good identification**					**73015120**
Eubacterium saburreum		100.0	0.38	NIT 1		
73015130	**Good identification**					**73015130**
Eubacterium saburreum		100.0	0.37	NIT 1		
73021000	**Good identification**					**73021000**
Eubacterium saburreum		97.43	0.38	NIT 1		
Eubacterium tortuosum		2.57	-.04			
73021010	**Good identification**					**73021010**
Eubacterium saburreum		100.0	0.37	NIT 1		
73021100	**Good identification**					**73021100**
Eubacterium saburreum		100.0	0.44	NIT 1		
73021110	**Good identification**					**73021110**
Eubacterium saburreum		100.0	0.43	NIT 1		
73025100	**Good identification**					**73025100**
Eubacterium saburreum		100.0	0.40	NIT 1		
73025110	**Good identification**					**73025110**
Eubacterium saburreum		100.0	0.38	NIT 1		
73031000	**Good identification**					**73031000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.46	NIT 1		
73031010	**Good identification**					**73031010**
Eubacterium saburreum		100.0	0.45	NIT 1		
73031100	**Good identification**					**73031100**
Eubacterium saburreum		100.0	0.53	NIT 1		
73031102	**Good identification**					**73031102**
Eubacterium saburreum		100.0	0.40	NIT 1		
73031110	**Good identification**					**73031110**
Eubacterium saburreum		100.0	0.51	NIT 1		
73031112	**Good identification**					**73031112**
Eubacterium saburreum		100.0	0.38	NIT 1		
73031120	**Good identification**					**73031120**
Eubacterium saburreum		100.0	0.40	NIT 1		
73031130	**Good identification**					**73031130**
Eubacterium saburreum		100.0	0.38	NIT 1		
73035000	**Good identification**					**73035000**
Eubacterium saburreum		100.0	0.42	NIT 1		
73035010	**Good identification**					**73035010**
Eubacterium saburreum		100.0	0.40	NIT 1		
73035100	**Good identification**					**73035100**
Eubacterium saburreum		100.0	0.48	NIT 1		
73035102	**Good identification**					**73035102**
Eubacterium saburreum		100.0	0.35	NIT 1		
73035110	**Good identification**					**73035110**
Eubacterium saburreum		100.0	0.46	NIT 1		
73035120	**Good identification**					**73035120**
Eubacterium saburreum		100.0	0.35	NIT 1		
73101000	**Good identification**					**73101000**
Eubacterium saburreum		99.56	0.51	NIT 1		
Eubacterium tortuosum		0.44	-.04			
73101002	**Good identification**					**73101002**
Eubacterium saburreum		100.0	0.39	NIT 1		
73101010	**Good identification**					**73101010**
Eubacterium saburreum		100.0	0.50	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
73101012	**Good identification**					**73101012**
Eubacterium saburreum		100.0	0.37	NIT 1		
73101020	**Good identification**					**73101020**
Eubacterium saburreum		100.0	0.39	NIT 1		
73101030	**Good identification**					**73101030**
Eubacterium saburreum		100.0	0.37	NIT 1		
73101100	**Very good identification**					**73101100**
					SPO	
Eubacterium saburreum		99.47	0.58	NIT 1	1	
Clostridium chauvoei		0.54	0.09		99	
73101101	**Good identification**					**73101101**
					SPO	
Clostridium chauvoei		98.14	0.43	IND 1	99	
Eubacterium saburreum		1.86	0.24	MNS 90	1	
73101102	**Good identification**					**73101102**
Eubacterium saburreum		100.0	0.45	NIT 1		
73101110	**Very good identification**					**73101110**
Eubacterium saburreum		100.0	0.56	NIT 1		
73101112	**Good identification**					**73101112**
Eubacterium saburreum		100.0	0.43	NIT 1		
73101120	**Good identification**					**73101120**
					SPO	
Eubacterium saburreum		98.99	0.45	NIT 1	1	
Clostridium chauvoei		1.01	0.01		99	
73101121	**Good identification**					**73101121**
					SPO	
Clostridium chauvoei		99.01	0.35	IND 1	99	
Eubacterium saburreum		0.99	0.11	MNS 90	1	
73101130	**Good identification**					**73101130**
Eubacterium saburreum		100.0	0.43	NIT 1		
73105000	**Good identification**					**73105000**
Eubacterium saburreum		100.0	0.47	NIT 1		
73105010	**Good identification**					**73105010**
Eubacterium saburreum		100.0	0.45	NIT 1		
73105100	**Good identification**					**73105100**
Eubacterium saburreum		100.0	0.53	NIT 1		
73105102	**Good identification**					**73105102**
Eubacterium saburreum		100.0	0.40	NIT 1		
73105110	**Good identification**					**73105110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.52	NIT 1			
73105112	**Good identification**						**73105112**
Eubacterium saburreum		100.0	0.39	NIT 1			
73105120	**Good identification**						**73105120**
Eubacterium saburreum		100.0	0.40	NIT 1			
73105130	**Good identification**						**73105130**
Eubacterium saburreum		100.0	0.39	NIT 1			
73111000	**Very good identification**						**73111000**
					GRA		
Eubacterium saburreum		98.82	0.60	NIT 1	85		
Actinomyces odontolyticus		0.70	0.19		99		
Prevotella intermedia		0.48	0.11		1		
73111002	**Good identification**						**73111002**
Eubacterium saburreum		97.59	0.47	NIT 1			
Actinomyces odontolyticus		2.41	0.15				
73111010	**Very good identification**						**73111010**
Eubacterium saburreum		100.0	0.58	NIT 1			
73111012	**Good identification**						**73111012**
Eubacterium saburreum		100.0	0.45	NIT 1			
73111020	**Good identification**						**73111020**
Eubacterium saburreum		99.35	0.47	NIT 1			
Actinomyces odontolyticus		0.65	0.05				
73111030	**Good identification**						**73111030**
Eubacterium saburreum		100.0	0.45	NIT 1			
73111100	**Very good identification**						**73111100**
					SPO		
Eubacterium saburreum		99.41	0.66	NIT 1	1		
Clostridium chauvoei		0.42	0.15		99		
Actinomyces odontolyticus		0.10	0.10		1		
73111101	**Good identification**						**73111101**
					SPO		
Clostridium chauvoei		97.62	0.50	IND 1	99		
Eubacterium saburreum		2.38	0.32	MNS 90	1		
73111102	**Good identification**						**73111102**
					SPO	CAT	
Eubacterium saburreum		95.48	0.53	NIT 1	1	1	
Capnocytophaga ochracea		3.80	0.17		1	90	
Clostridium chauvoei		0.40	0.03		99	1	
73111103	**Good identification**						**73111103**
					SPO		
Clostridium chauvoei		97.62	0.37	IND 1	99		
Eubacterium saburreum		2.38	0.19	MNS 90	1		
73111110	**Very good identification**						**73111110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.64	NIT 1			
73111112	**Good identification**						**73111112**
					CAT		
Eubacterium saburreum		99.46	0.52	NIT 1	1		
Capnocytophaga ochracea		0.54	0.01		90		
73111120	**Acceptable identification**						**73111120**
					CAT	GRA	
Eubacterium saburreum		94.34	0.53	NIT 1	1	85	
Capnocytophaga ochracea		3.76	0.17		90	1	
Clostridium perfringens		0.78	0.08		1	99	
73111121	**Good identification**						**73111121**
					SPO		
Clostridium chauvoei		98.73	0.41	IND 1	99		
Eubacterium saburreum		1.27	0.19	MNS 90	1		
73111130	**Good identification**						**73111130**
					CAT		
Eubacterium saburreum		99.46	0.52	NIT 1	1		
Capnocytophaga ochracea		0.54	0.01		90		
73111132	**Low discrimination**						**73111132**
					CAT		
Eubacterium saburreum		78.38	0.39	NIT 1	1		
Capnocytophaga ochracea		21.62	0.17		90		
73115000	**Good identification**						**73115000**
Eubacterium saburreum		99.36	0.55	NIT 1			
Actinomyces odontolyticus		0.65	0.14				
73115002	**Good identification**						**73115002**
Eubacterium saburreum		97.79	0.42	NIT 1			
Actinomyces odontolyticus		2.21	0.10				
73115010	**Good identification**						**73115010**
Eubacterium saburreum		100.0	0.53	NIT 1			
73115012	**Good identification**						**73115012**
Eubacterium saburreum		100.0	0.41	NIT 1			
73115020	**Good identification**						**73115020**
Eubacterium saburreum		100.0	0.42	NIT 1			
73115030	**Good identification**						**73115030**
Eubacterium saburreum		100.0	0.41	NIT 1			
73115100	**Very good identification**						**73115100**
Eubacterium saburreum		100.0	0.61	NIT 1			
73115102	**Good identification**						**73115102**
					CAT		
Eubacterium saburreum		98.18	0.48	NIT 1	1		
Capnocytophaga ochracea		1.82	0.07		90		
73115110	**Very good identification**						**73115110**
Eubacterium saburreum		100.0	0.60	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
73115112	**Good identification**					**73115112**
Eubacterium saburreum		100.0	0.47	NIT 1		
73115120	**Good identification**					**73115120**
					CAT	
Eubacterium saburreum		98.18	0.48	NIT 1	1	
Capnocytophaga ochracea		1.82	0.07		90	
73115122	**Low discrimination**					**73115122**
					CAT	
Eubacterium saburreum		51.46	0.35	NIT 1	1	
Capnocytophaga ochracea		48.54	0.23		90	
73115130	**Good identification**					**73115130**
Eubacterium saburreum		100.0	0.47	NIT 1		
73121000	**Good identification**					**73121000**
Eubacterium saburreum		100.0	0.48	NIT 1		
73121002	**Good identification**					**73121002**
Eubacterium saburreum		100.0	0.36	NIT 1		
73121010	**Good identification**					**73121010**
Eubacterium saburreum		100.0	0.47	NIT 1		
73121020	**Good identification**					**73121020**
Eubacterium saburreum		100.0	0.36	NIT 1		
73121100	**Good identification**					**73121100**
Eubacterium saburreum		100.0	0.55	NIT 1		
73121102	**Good identification**					**73121102**
Eubacterium saburreum		100.0	0.42	NIT 1		
73121110	**Good identification**					**73121110**
Eubacterium saburreum		100.0	0.53	NIT 1		
73121112	**Good identification**					**73121112**
Eubacterium saburreum		100.0	0.40	NIT 1		
73121120	**Good identification**					**73121120**
Eubacterium saburreum		100.0	0.42	NIT 1		
73121130	**Good identification**					**73121130**
Eubacterium saburreum		100.0	0.40	NIT 1		
73125000	**Good identification**					**73125000**
Eubacterium saburreum		100.0	0.44	NIT 1		
73125010	**Good identification**					**73125010**
Eubacterium saburreum		100.0	0.42	NIT 1		
73125100	**Good identification**					**73125100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.50	NIT 1		
73125102	**Good identification**					**73125102**
Eubacterium saburreum		100.0	0.37	NIT 1		
73125110	**Good identification**					**73125110**
Eubacterium saburreum		100.0	0.49	NIT 1		
73125112	**Good identification**					**73125112**
Eubacterium saburreum		100.0	0.36	NIT 1		
73125120	**Good identification**					**73125120**
Eubacterium saburreum		100.0	0.37	NIT 1		
73125130	**Good identification**					**73125130**
Eubacterium saburreum		100.0	0.36	NIT 1		
73131000	**Very good identification**					**73131000**
Eubacterium saburreum		100.0	0.57	NIT 1		
73131002	**Good identification**					**73131002**
Eubacterium saburreum		100.0	0.44	NIT 1		
73131010	**Good identification**					**73131010**
Eubacterium saburreum		100.0	0.55	NIT 1		
73131012	**Good identification**					**73131012**
Eubacterium saburreum		100.0	0.42	NIT 1		
73131020	**Good identification**					**73131020**
Eubacterium saburreum		100.0	0.44	NIT 1		
73131030	**Good identification**					**73131030**
Eubacterium saburreum		100.0	0.42	NIT 1		
73131100	**Very good identification**					**73131100**
Eubacterium saburreum		100.0	0.63	NIT 1		
73131102	**Good identification**					**73131102**
Eubacterium saburreum		100.0	0.50	NIT 1		
73131110	**Very good identification**					**73131110**
Eubacterium saburreum		100.0	0.61	NIT 1		
73131112	**Good identification**					**73131112**
Eubacterium saburreum		100.0	0.48	NIT 1		
73131120	**Good identification**					**73131120**
Eubacterium saburreum		97.76	0.50	NIT 1		
Clostridium perfringens		2.24	0.13			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
73131122	**Acceptable identification**						**73131122**
					CAT	GRA	
Eubacterium saburreum		91.59	0.37	NIT 1	1	85	
Clostridium perfringens		5.59	0.07		1	99	
Capnocytophaga ochracea		2.82	-.01		90	1	
73131130	**Good identification**						**73131130**
Eubacterium saburreum		100.0	0.48	NIT 1			
73131132	**Good identification**						**73131132**
Eubacterium saburreum		100.0	0.36	NIT 1			
73135000	**Good identification**						**73135000**
Eubacterium saburreum		100.0	0.52	NIT 1			
73135002	**Good identification**						**73135002**
Eubacterium saburreum		100.0	0.39	NIT 1			
73135010	**Good identification**						**73135010**
Eubacterium saburreum		100.0	0.50	NIT 1			
73135012	**Good identification**						**73135012**
Eubacterium saburreum		100.0	0.38	NIT 1			
73135020	**Good identification**						**73135020**
Eubacterium saburreum		100.0	0.39	NIT 1			
73135030	**Good identification**						**73135030**
Eubacterium saburreum		100.0	0.38	NIT 1			
73135100	**Very good identification**						**73135100**
Eubacterium saburreum		100.0	0.58	NIT 1			
73135102	**Good identification**						**73135102**
Eubacterium saburreum		100.0	0.45	NIT 1			
73135110	**Very good identification**						**73135110**
Eubacterium saburreum		100.0	0.57	NIT 1			
73135112	**Good identification**						**73135112**
Eubacterium saburreum		100.0	0.44	NIT 1			
73135120	**Good identification**						**73135120**
Eubacterium saburreum		100.0	0.45	NIT 1			
73135130	**Good identification**						**73135130**
Eubacterium saburreum		100.0	0.44	NIT 1			
73201000	**Very good identification**						**73201000**
Eubacterium tortuosum		100.0	0.64	IND 1			
73201002	**Good identification**						**73201002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium tortuosum		100.0	0.52	IND 1		
73201020	**Good identification**					**73201020**
Eubacterium tortuosum		100.0	0.52	IND 1		
73201022	**Good identification**					**73201022**
Eubacterium tortuosum		100.0	0.39	IND 1		
73241000	**Good identification**					**73241000**
Eubacterium tortuosum		100.0	0.48	IND 1		
73241002	**Good identification**					**73241002**
Eubacterium tortuosum		100.0	0.35	IND 1		
73241020	**Good identification**					**73241020**
Eubacterium tortuosum		100.0	0.35	IND 1		
73377100	**Good identification**					**73377100**
Actinomyces israelii		100.0	0.41	IND 1		
73377102	**Good identification**					**73377102**
Actinomyces israelii		100.0	0.44	IND 1		
73377142	**Good identification**					**73377142**
Actinomyces israelii		100.0	0.37	IND 1		
73377500	**Good identification**					**73377500**
Actinomyces israelii		100.0	0.40	IND 1		
73377502	**Good identification**					**73377502**
Actinomyces israelii		100.0	0.43	IND 1		
73377542	**Good identification**					**73377542**
Actinomyces israelii		100.0	0.36	IND 1		
73401010	**Good identification**					**73401010**
Eubacterium saburreum		100.0	0.35	NIT 1		
73401100	**Good identification**					**73401100**
Eubacterium saburreum		100.0	0.43	NIT 1		
73401110	**Good identification**					**73401110**
Eubacterium saburreum		100.0	0.41	NIT 1		
73405100	**Good identification**					**73405100**
Eubacterium saburreum		100.0	0.38	NIT 1		
73405110	**Good identification**					**73405110**
Eubacterium saburreum		100.0	0.37	NIT 1		
73411000	**Good identification**					**73411000**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		98.44	0.45	NIT 1		
Fusobacterium mortiferum		1.57	0.16			
73411010	**Good identification**					**73411010**
Eubacterium saburreum		100.0	0.43	NIT 1		
73411100	**Good identification**					**73411100**
Eubacterium saburreum		98.44	0.51	NIT 1		
Fusobacterium mortiferum		1.57	0.22			
73411102	**Good identification**					**73411102**
					CAT	
Eubacterium saburreum		97.38	0.38	NIT 1	1	
Capnocytophaga ochracea		2.62	-.01		90	
73411110	**Good identification**					**73411110**
Eubacterium saburreum		100.0	0.49	NIT 1		
73411112	**Good identification**					**73411112**
Eubacterium saburreum		100.0	0.37	NIT 1		
73411120	**Good identification**					**73411120**
					CAT	
Eubacterium saburreum		97.38	0.38	NIT 1	1	
Capnocytophaga ochracea		2.62	-.01		90	
73411130	**Good identification**					**73411130**
Eubacterium saburreum		100.0	0.37	NIT 1		
73415000	**Good identification**					**73415000**
Eubacterium saburreum		100.0	0.40	NIT 1		
73415010	**Good identification**					**73415010**
Eubacterium saburreum		100.0	0.39	NIT 1		
73415100	**Good identification**					**73415100**
Eubacterium saburreum		100.0	0.46	NIT 1		
73415110	**Good identification**					**73415110**
Eubacterium saburreum		100.0	0.45	NIT 1		
73421100	**Good identification**					**73421100**
Eubacterium saburreum		100.0	0.40	NIT 1		
73421110	**Good identification**					**73421110**
Eubacterium saburreum		100.0	0.38	NIT 1		
73425100	**Good identification**					**73425100**
Eubacterium saburreum		100.0	0.35	NIT 1		
73431000	**Good identification**					**73431000**
Eubacterium saburreum		100.0	0.42	NIT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
73431010	**Good identification**					**73431010**
Eubacterium saburreum		100.0	0.40	NIT 1		
73431100	**Good identification**					**73431100**
Eubacterium saburreum		99.29	0.48	NIT 1		
Fusobacterium mortiferum		0.71	0.13			
73431102	**Good identification**					**73431102**
Eubacterium saburreum		100.0	0.35	NIT 1		
73431110	**Good identification**					**73431110**
Eubacterium saburreum		100.0	0.46	NIT 1		
73431120	**Good identification**					**73431120**
Eubacterium saburreum		100.0	0.35	NIT 1		
73435000	**Good identification**					**73435000**
Eubacterium saburreum		100.0	0.37	NIT 1		
73435010	**Good identification**					**73435010**
Eubacterium saburreum		100.0	0.36	NIT 1		
73435100	**Good identification**					**73435100**
Eubacterium saburreum		100.0	0.43	NIT 1		
73435110	**Good identification**					**73435110**
Eubacterium saburreum		100.0	0.42	NIT 1		
73501000	**Good identification**					**73501000**
Eubacterium saburreum		100.0	0.47	NIT 1		
73501010	**Good identification**					**73501010**
Eubacterium saburreum		100.0	0.45	NIT 1		
73501100	**Good identification**					**73501100**
					COC	
Eubacterium saburreum		99.54	0.53	NIT 1	1	
Blautia hansenii		0.47	-.03		99	
73501102	**Good identification**					**73501102**
					CAT	
Eubacterium saburreum		98.02	0.40	NIT 1	1	
Capnocytophaga ochracea		1.98	-.01		90	
73501110	**Good identification**					**73501110**
Eubacterium saburreum		100.0	0.52	NIT 1		
73501112	**Good identification**					**73501112**
Eubacterium saburreum		100.0	0.39	NIT 1		
73501120	**Good identification**					**73501120**
					CAT	
Eubacterium saburreum		95.92	0.40	NIT 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Prevotella melaninogenica		2.15	-.04		1		
Capnocytophaga ochracea		1.94	-.01		90		
73501130	**Good identification**						**73501130**
Eubacterium saburreum		100.0	0.39	NIT 1			
73505000	**Good identification**						**73505000**
Eubacterium saburreum		100.0	0.42	NIT 1			
73505010	**Good identification**						**73505010**
Eubacterium saburreum		100.0	0.41	NIT 1			
73505100	**Good identification**						**73505100**
Eubacterium saburreum		100.0	0.49	NIT 1			
73505102	**Good identification**						**73505102**
Eubacterium saburreum		100.0	0.36	NIT 1			
73505110	**Good identification**						**73505110**
Eubacterium saburreum		100.0	0.47	NIT 1			
73505120	**Good identification**						**73505120**
Eubacterium saburreum		100.0	0.36	NIT 1			
73511000	**Good identification**						**73511000**
Eubacterium saburreum		98.66	0.55	NIT 1			
Prevotella intermedia		1.34	0.14				
73511002	**Good identification**						**73511002**
					GRA	CAT	
Eubacterium saburreum		96.98	0.42	NIT 1	85	1	
Capnocytophaga ochracea		1.52	-.01		1	90	
Actinomyces viscosus		1.50	0.00		99	76	
73511010	**Good identification**						**73511010**
Eubacterium saburreum		100.0	0.53	NIT 1			
73511012	**Good identification**						**73511012**
Eubacterium saburreum		100.0	0.41	NIT 1			
73511020	**Good identification**						**73511020**
					CAT		
Eubacterium saburreum		98.46	0.42	NIT 1	1		
Capnocytophaga ochracea		1.55	-.01		90		
73511030	**Good identification**						**73511030**
Eubacterium saburreum		100.0	0.41	NIT 1			
73511100	**Very good identification**						**73511100**
					CAT		
Eubacterium saburreum		98.71	0.61	NIT 1	1		
Capnocytophaga ochracea		1.29	0.17		90		
73511102	**Low discrimination**						**73511102**
					GRA	CAT	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		59.61	0.48	NIT 1	85	1	
Capnocytophaga ochracea		39.69	0.33		1	90	
Actinomyces viscosus		0.70	0.04		99	76	
73511110	**Very good identification**						**73511110**
					CAT		
Eubacterium saburreum		99.82	0.60	NIT 1	1		
Capnocytophaga ochracea		0.18	0.01		90		
73511112	**Acceptable identification**						**73511112**
					CAT		
Eubacterium saburreum		91.71	0.47	NIT 1	1		
Capnocytophaga ochracea		8.29	0.17		90		
73511120	**Low discrimination**						**73511120**
					CAT		
Eubacterium saburreum		57.70	0.48	NIT 1	1		
Capnocytophaga ochracea		38.42	0.33		90		
Prevotella melaninogenica		3.88	0.12		1		
73511122	**Good identification**						**73511122**
					CAT		
Capnocytophaga ochracea		96.94	0.50	IND 1	90		
Eubacterium saburreum		2.86	0.35	MNS 90	1		
Prevotella melaninogenica		0.21	0.00		1		
73511130	**Acceptable identification**						**73511130**
					CAT		
Eubacterium saburreum		91.71	0.47	NIT 1	1		
Capnocytophaga ochracea		8.29	0.17		90		
73511522	**Good identification**						**73511522**
					GRA	CAT	
Capnocytophaga ochracea		99.51	0.37	IND 1	1	90	
Bifidobacterium longum subsp. longum		0.49	0.05	MNS 90	99	1	
73515000	**Good identification**						**73515000**
Eubacterium saburreum		100.0	0.50	NIT 1			
73515002	**Good identification**						**73515002**
Eubacterium saburreum		100.0	0.37	NIT 1			
73515010	**Good identification**						**73515010**
Eubacterium saburreum		100.0	0.49	NIT 1			
73515012	**Good identification**						**73515012**
Eubacterium saburreum		100.0	0.36	NIT 1			
73515020	**Good identification**						**73515020**
Eubacterium saburreum		100.0	0.37	NIT 1			
73515030	**Good identification**						**73515030**
Eubacterium saburreum		100.0	0.36	NIT 1			
73515100	**Very good identification**						**73515100**
					CAT		
Eubacterium saburreum		99.40	0.57	NIT 1	1		
Capnocytophaga ochracea		0.60	0.07		90		
73515102	**Low discrimination**						**73515102**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Eubacterium saburreum		76.39	0.44	NIT 1	1	
Capnocytophaga ochracea		23.61	0.23		90	
73515110	**Good identification**					**73515110**
Eubacterium saburreum		100.0	0.55	NIT 1		
73515112	**Good identification**					**73515112**
					CAT	
Eubacterium saburreum		95.97	0.42	NIT 1	1	
Capnocytophaga ochracea		4.03	0.07		90	
73515120	**Low discrimination**					**73515120**
					CAT	
Eubacterium saburreum		76.39	0.44	NIT 1	1	
Capnocytophaga ochracea		23.61	0.23		90	
73515122	**Acceptable identification**					**73515122**
					CAT	
Capnocytophaga ochracea		90.71	0.39	IND 1	90	
Eubacterium saburreum		5.75	0.31	MNS 90	1	
Bacteroides fragilis		3.54	0.06		99	
73515130	**Good identification**					**73515130**
					CAT	
Eubacterium saburreum		95.97	0.42	NIT 1	1	
Capnocytophaga ochracea		4.03	0.07		90	
73521000	**Good identification**					**73521000**
Eubacterium saburreum		100.0	0.44	NIT 1		
73521010	**Good identification**					**73521010**
Eubacterium saburreum		100.0	0.42	NIT 1		
73521100	**Good identification**					**73521100**
Eubacterium saburreum		100.0	0.50	NIT 1		
73521102	**Good identification**					**73521102**
Eubacterium saburreum		100.0	0.37	NIT 1		
73521110	**Good identification**					**73521110**
Eubacterium saburreum		100.0	0.49	NIT 1		
73521112	**Good identification**					**73521112**
Eubacterium saburreum		100.0	0.36	NIT 1		
73521120	**Good identification**					**73521120**
Eubacterium saburreum		100.0	0.37	NIT 1		
73521130	**Good identification**					**73521130**
Eubacterium saburreum		100.0	0.36	NIT 1		
73525000	**Good identification**					**73525000**
Eubacterium saburreum		100.0	0.39	NIT 1		
73525010	**Good identification**					**73525010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.38	NIT 1			
73525100	**Good identification**						**73525100**
Eubacterium saburreum		100.0	0.46	NIT 1			
73525110	**Good identification**						**73525110**
Eubacterium saburreum		100.0	0.44	NIT 1			
73531000	**Good identification**						**73531000**
Eubacterium saburreum		100.0	0.52	NIT 1			
73531002	**Good identification**						**73531002**
Eubacterium saburreum		100.0	0.39	NIT 1			
73531010	**Good identification**						**73531010**
Eubacterium saburreum		100.0	0.50	NIT 1			
73531012	**Good identification**						**73531012**
Eubacterium saburreum		100.0	0.38	NIT 1			
73531020	**Good identification**						**73531020**
Eubacterium saburreum		100.0	0.39	NIT 1			
73531030	**Good identification**						**73531030**
Eubacterium saburreum		100.0	0.38	NIT 1			
73531100	**Very good identification**						**73531100**
Eubacterium saburreum		100.0	0.58	NIT 1			
73531102	**Good identification**						**73531102**
					CAT		
Eubacterium saburreum		99.00	0.45	NIT 1	1		
Capnocytophaga ochracea		1.00	-.01		90		
73531110	**Very good identification**						**73531110**
Eubacterium saburreum		100.0	0.57	NIT 1			
73531112	**Good identification**						**73531112**
Eubacterium saburreum		100.0	0.44	NIT 1			
73531120	**Good identification**						**73531120**
					CAT	GRA	
Eubacterium saburreum		98.22	0.45	NIT 1	1	85	
Capnocytophaga ochracea		0.99	-.01		90	1	
Clostridium perfringens		0.80	0.01		1	99	
73531130	**Good identification**						**73531130**
Eubacterium saburreum		100.0	0.44	NIT 1			
73535000	**Good identification**						**73535000**
Eubacterium saburreum		100.0	0.47	NIT 1			
73535010	**Good identification**						**73535010**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.46	NIT 1		
73535100	**Good identification**					**73535100**
Eubacterium saburreum		100.0	0.54	NIT 1		
73535102	**Good identification**					**73535102**
Eubacterium saburreum		100.0	0.41	NIT 1		
73535110	**Good identification**					**73535110**
Eubacterium saburreum		100.0	0.52	NIT 1		
73535112	**Good identification**					**73535112**
Eubacterium saburreum		100.0	0.39	NIT 1		
73535120	**Good identification**					**73535120**
Eubacterium saburreum		100.0	0.41	NIT 1		
73535130	**Good identification**					**73535130**
Eubacterium saburreum		100.0	0.39	NIT 1		
73551122	**Good identification**					**73551122**
Capnocytophaga ochracea		100.0	0.39	IND 1		
73577102	**Good identification**					**73577102**
Actinomyces israelii		100.0	0.36	IND 1		
73601000	**Very good identification**					**73601000**
Eubacterium tortuosum		100.0	0.56	IND 1		
73601002	**Good identification**					**73601002**
Eubacterium tortuosum		100.0	0.43	IND 1		
73601020	**Good identification**					**73601020**
Eubacterium tortuosum		100.0	0.43	IND 1		
73641000	**Good identification**					**73641000**
Eubacterium tortuosum		100.0	0.40	IND 1		
73757100	**Good identification**					**73757100**
Actinomyces israelii		100.0	0.41	IND 1		
73757102	**Good identification**					**73757102**
Actinomyces israelii		100.0	0.44	IND 1		
73757142	**Good identification**					**73757142**
Actinomyces israelii		100.0	0.37	IND 1		
73757500	**Good identification**					**73757500**
Actinomyces israelii		100.0	0.40	IND 1		
73757502	**Acceptable identification**					**73757502**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		94.90	0.43	IND 1			
Bifidobacterium adolescentis		5.10	0.15				
73757542	**Good identification**						**73757542**
Actinomyces israelii		100.0	0.36	IND 1			
73773101	**Good identification**						**73773101**
Actinomyces naeslundii		100.0	0.36	IND 1			
73773102	**Low discrimination**						**73773102**
Actinomyces israelii		82.96	0.36	IND 1			
Actinomyces naeslundii		17.04	0.33	XYL 92			
73773103	**Good identification**						**73773103**
Actinomyces naeslundii		100.0	0.36	IND 1			
73773121	**Good identification**						**73773121**
Actinomyces naeslundii		100.0	0.36	IND 1			
73773123	**Good identification**						**73773123**
Actinomyces naeslundii		100.0	0.36	IND 1			
73775100	**Good identification**						**73775100**
Actinomyces israelii		100.0	0.38	IND 1			
73775102	**Good identification**						**73775102**
Actinomyces israelii		100.0	0.41	IND 1			
73775500	**Low discrimination**						**73775500**
					GRA	SPO	
Actinomyces israelii		51.77	0.37	IND 1	99	1	
Mitsuokella multacida		34.07	0.21		1	1	
Clostridium butyricum		14.16	0.06		99	99	
73775502	**Low discrimination**						**73775502**
					SPO		
Actinomyces israelii		84.34	0.40	IND 1	1		
Clostridium butyricum		5.98	-.01		99		
Bifidobacterium adolescentis		5.57	0.13		1		
73777000	**Good identification**						**73777000**
Actinomyces israelii		100.0	0.41	IND 1			
73777002	**Good identification**						**73777002**
Actinomyces israelii		100.0	0.44	IND 1			
73777042	**Good identification**						**73777042**
Actinomyces israelii		100.0	0.37	IND 1			
73777100	**Good identification**						**73777100**
Actinomyces israelii		100.0	0.51	IND 1			
73777102	**Good identification**						**73777102**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Actinomyces israelii		100.0	0.54	IND 1				
73777110	**Good identification**							**73777110**
Actinomyces israelii		100.0	0.37	IND 1				
73777112	**Good identification**							**73777112**
Actinomyces israelii		100.0	0.40	IND 1				
73777140	**Good identification**							**73777140**
Actinomyces israelii		100.0	0.44	IND 1				
73777142	**Good identification**							**73777142**
Actinomyces israelii		100.0	0.47	IND 1				
73777400	**Good identification**							**73777400**
Actinomyces israelii		100.0	0.39	IND 1				
73777402	**Good identification**							**73777402**
Actinomyces israelii		100.0	0.42	IND 1				
73777442	**Good identification**							**73777442**
Actinomyces israelii		100.0	0.35	IND 1				
73777500	**Low discrimination**							**73777500**
					GRA			
Actinomyces israelii		78.68	0.49	IND 1	99			
Mitsuokella multacida		21.32	0.27		1			
73777502	**Good identification**							**73777502**
					GRA			
Actinomyces israelii		97.13	0.52	IND 1	99			
Mitsuokella multacida		1.95	0.11		1			
Bifidobacterium adolescentis		0.93	0.12		99			
73777510	**Low discrimination**							**73777510**
					GRA			
Actinomyces israelii		62.33	0.35	IND 1	99			
Mitsuokella multacida		37.67	0.19	MLZ 13	1			
73777512	**Low discrimination**							**73777512**
Actinomyces israelii		84.46	0.38	IND 1				
Bifidobacterium adolescentis		6.59	0.13	MLZ 13				
Bifidobacterium dentium		5.18	-.07					
73777540	**Low discrimination**							**73777540**
					GRA			
Actinomyces israelii		58.94	0.42	IND 1	99			
Mitsuokella multacida		41.06	0.27		1			
73777542	**Good identification**							**73777542**
					GRA			
Actinomyces israelii		95.09	0.45	IND 1	99			
Mitsuokella multacida		4.91	0.11		1			
74000000	**Low discrimination**							**74000000**
					COC	GRA	SPO	
Clostridium bifermentans		88.96	0.41	NIT 1	1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Peptoniphilus asaccharolyticus		3.88	0.06	GLU 92	99	99	1	
Veillonella parvula		1.29	-.03		99	1	1	
74001520	**Good identification**							**74001520**
Odoribacter splanchnicus		100.0	0.51	GLU 99				
74011520	**Good identification**							**74011520**
Odoribacter splanchnicus		100.0	0.56	GLU 99				
74011560	**Good identification**							**74011560**
Odoribacter splanchnicus		100.0	0.39	GLU 99				
74100000	**Good identification**							**74100000**
Clostridium bifermentans		100.0	0.48	NIT 1				
75000000	**Very good identification**							**75000000**
					SPO			
Clostridium bifermentans		99.69	0.60	NIT 1	99			
Clostridium botulinum biovar A		0.16	0.13		99			
Atopobium minutum		0.09	0.16		1			
75000020	**Low discrimination**							**75000020**
					SPO	CAT		
Clostridium bifermentans		60.52	0.39	NIT 1	99	1		
Clostridium cadaveris		19.88	0.29	NAG 6	99	1		
Propionibacterium acnes		19.31	0.32		1	95		
75000040	**Good identification**							**75000040**
Clostridium bifermentans		98.84	0.36	NIT 1				
Clostridium botulinum biovar A		1.17	0.04	SOR 4				
75000520	**Good identification**							**75000520**
Odoribacter splanchnicus		100.0	0.51	GAL 99				
75001120	**Good identification**							**75001120**
Odoribacter splanchnicus		100.0	0.51	ARA 99				
75001420	**Good identification**							**75001420**
Odoribacter splanchnicus		100.0	0.51	LAC 99				
75001500	**Good identification**							**75001500**
Odoribacter splanchnicus		100.0	0.51	NAG 99				
75001520	**Very good identification**							**75001520**
Odoribacter splanchnicus		100.0	0.85					
75001521	**Good identification**							**75001521**
Odoribacter splanchnicus		100.0	0.51	URE 1				
75001522	**Good identification**							**75001522**
Odoribacter splanchnicus		100.0	0.51	BGL 1				
75001530	**Good identification**							**75001530**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Odoribacter splanchnicus		100.0	0.51	MLZ 1				
75001560	**Very good identification**							**75001560**
Odoribacter splanchnicus		100.0	0.69	SOR 10				
75001720	**Good identification**							**75001720**
Odoribacter splanchnicus		100.0	0.51	RHA 1				
75003520	**Good identification**							**75003520**
Odoribacter splanchnicus		100.0	0.51	MAN 1				
75005520	**Good identification**							**75005520**
Odoribacter splanchnicus		100.0	0.51	XYL 1				
75010000	**Low discrimination**							**75010000**
					GRA	SPO	CAT	
Clostridium bifermentans		39.98	0.38	NIT 1	99	99	1	
Fusobacterium varium		38.42	0.32	FRU 5	1	1	1	
Propionibacterium acnes		15.47	0.32		99	1	95	
75010020	**Very good identification**							**75010020**
					SPO	CAT		
Propionibacterium acnes		98.77	0.66	ESL 1	1	95		
Clostridium cadaveris		1.03	0.29		99	1		
Clostridium bifermentans		0.17	0.17		99	1		
75010060	**Good identification**							**75010060**
Propionibacterium acnes		100.0	0.55	ESL 1				
75010520	**Good identification**							**75010520**
Odoribacter splanchnicus		100.0	0.56	GAL 99				
75010560	**Good identification**							**75010560**
Odoribacter splanchnicus		100.0	0.39	GAL 99				
75011020	**Very good identification**							**75011020**
					GRA	CAT		
Propionibacterium acnes		99.26	0.66	ESL 1	99	95		
Odoribacter splanchnicus		0.74	0.21		1	1		
75011060	**Good identification**							**75011060**
					GRA	CAT		
Propionibacterium acnes		99.62	0.55	ESL 1	99	95		
Odoribacter splanchnicus		0.38	0.05		1	1		
75011120	**Good identification**							**75011120**
					GRA	CAT		
Odoribacter splanchnicus		98.65	0.56	ARA 99	1	1		
Propionibacterium acnes		1.35	0.32		99	95		
75011160	**Good identification**							**75011160**
					GRA	CAT		
Odoribacter splanchnicus		97.38	0.39	ARA 99	1	1		
Propionibacterium acnes		2.62	0.20	SOR 10	99	95		
75011420	**Good identification**							**75011420**
					GRA	CAT		
Odoribacter splanchnicus		98.65	0.56	LAC 99	1	1		
Propionibacterium acnes		1.35	0.32		99	95		

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
75011460	**Good identification**						**75011460**
					GRA	CAT	
Odoribacter splanchnicus		97.38	0.39	LAC 99	1	1	
Propionibacterium acnes		2.62	0.20	SOR 10	99	95	
75011500	**Good identification**						**75011500**
Odoribacter splanchnicus		100.0	0.56	NAG 99			
75011520	**Very good identification**						**75011520**
Odoribacter splanchnicus		100.0	0.90				
75011521	**Good identification**						**75011521**
Odoribacter splanchnicus		100.0	0.56	URE 1			
75011522	**Good identification**						**75011522**
Odoribacter splanchnicus		100.0	0.56	BGL 1			
75011530	**Good identification**						**75011530**
Odoribacter splanchnicus		100.0	0.56	MLZ 1			
75011540	**Good identification**						**75011540**
Odoribacter splanchnicus		100.0	0.39	NAG 99			
75011560	**Very good identification**						**75011560**
Odoribacter splanchnicus		100.0	0.73	SOR 10			
75011561	**Good identification**						**75011561**
Odoribacter splanchnicus		100.0	0.39	SOR 10			
75011562	**Good identification**						**75011562**
Odoribacter splanchnicus		100.0	0.39	SOR 10			
75011570	**Good identification**						**75011570**
Odoribacter splanchnicus		100.0	0.39	MLZ 1			
75011720	**Good identification**						**75011720**
Odoribacter splanchnicus		100.0	0.56	RHA 1			
75011760	**Good identification**						**75011760**
Odoribacter splanchnicus		100.0	0.39	RHA 1			
75012020	**Very good identification**						**75012020**
					CAT		
Propionibacterium acnes		99.96	0.62	ESL 1	95		
Eubacterium limosum		0.04	-.03		1		
75012060	**Good identification**						**75012060**
Propionibacterium acnes		100.0	0.50	ESL 1			
75013020	**Very good identification**						**75013020**
Propionibacterium acnes		100.0	0.62	ESL 1			
75013060	**Good identification**						**75013060**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Propionibacterium acnes		100.0	0.50	ESL 1			
75013520	**Good identification**						**75013520**
Odoribacter splanchnicus		100.0	0.56	MAN 1			
75013560	**Good identification**						**75013560**
Odoribacter splanchnicus		100.0	0.39	MAN 1			
75015520	**Good identification**						**75015520**
Odoribacter splanchnicus		100.0	0.56	XYL 1			
75015560	**Good identification**						**75015560**
Odoribacter splanchnicus		100.0	0.39	XYL 1			
75021520	**Good identification**						**75021520**
Odoribacter splanchnicus		100.0	0.51	TRE 1			
75031520	**Good identification**						**75031520**
Odoribacter splanchnicus		100.0	0.56	TRE 1			
75031560	**Good identification**						**75031560**
Odoribacter splanchnicus		100.0	0.39	TRE 1			
75041520	**Good identification**						**75041520**
Odoribacter splanchnicus		100.0	0.51	CEL 1			
75051520	**Good identification**						**75051520**
Odoribacter splanchnicus		100.0	0.56	CEL 1			
75051560	**Good identification**						**75051560**
Odoribacter splanchnicus		100.0	0.39	CEL 1			
75100000	**Very good identification**						**75100000**
					GRA	SPO	
Clostridium bifermentans		99.91	0.66	NIT 1	99	99	
Clostridium botulinum biovar A		0.06	0.11		99	99	
Prevotella disiens		0.02	-.03		1	1	
75100020	**Good identification**						**75100020**
Clostridium bifermentans		99.26	0.45	NIT 1			
Clostridium cadaveris		0.74	0.07	NAG 6			
75100040	**Good identification**						**75100040**
Clostridium bifermentans		99.59	0.42	NIT 1			
Clostridium botulinum biovar A		0.41	0.02	SOR 4			
75101520	**Good identification**						**75101520**
Odoribacter splanchnicus		100.0	0.51	MLT 1			
75101722	**Good identification**						**75101722**
Bacteroides eggerthii		100.0	0.36	NIT 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
75105722	**Good identification**						**75105722**
Bacteroides eggerthii		100.0	0.56	NIT 1			
75110000	**Acceptable identification**						**75110000**
					GRA	SPO	
Clostridium bifermentans		90.99	0.44	NIT 1	99	99	
Prevotella intermedia		8.27	0.29	FRU 5	1	1	
Fusobacterium varium		0.38	-.03		1	1	
75111520	**Good identification**						**75111520**
Odoribacter splanchnicus		100.0	0.56	MLT 1			
75111560	**Good identification**						**75111560**
Odoribacter splanchnicus		100.0	0.39	MLT 1			
75141722	**Good identification**						**75141722**
Bacteroides eggerthii		100.0	0.47	NIT 1			
75145722	**Very good identification**						**75145722**
Bacteroides eggerthii		100.0	0.66	NIT 1			
75200000	**Good identification**						**75200000**
Clostridium bifermentans		100.0	0.43	NIT 1			
75201520	**Good identification**						**75201520**
Odoribacter splanchnicus		100.0	0.51	SAL 1			
75211520	**Good identification**						**75211520**
Odoribacter splanchnicus		100.0	0.56	SAL 1			
75211560	**Good identification**						**75211560**
Odoribacter splanchnicus		100.0	0.39	SAL 1			
75300000	**Good identification**						**75300000**
Clostridium bifermentans		100.0	0.50	NIT 1			
75401520	**Good identification**						**75401520**
Odoribacter splanchnicus		100.0	0.51	RAF 1			
75411520	**Good identification**						**75411520**
Odoribacter splanchnicus		100.0	0.56	RAF 1			
75411560	**Good identification**						**75411560**
Odoribacter splanchnicus		100.0	0.39	RAF 1			
75575722	**Acceptable identification**						**75575722**
Bacteroides thetaiotaomicron		93.35	0.37	NIT 1			
Bacteroides ovatus		6.24	0.17	SUC 98			
Parabacteroides distasonis		0.41	0.13				
76575722	**Good identification**						**76575722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides thetaiotaomicron		95.77	0.40	NIT 1		
Bacteroides ovatus		4.23	0.17	GLU 97		
77001100	**Good identification**					**77001100**
Eubacterium saburreum		100.0	0.37	NIT 1		
77001110	**Good identification**					**77001110**
Eubacterium saburreum		100.0	0.36	NIT 1		
77001520	**Good identification**					**77001520**
Odoribacter splanchnicus		100.0	0.51	SUC 1		
77011000	**Good identification**					**77011000**
Eubacterium saburreum		96.69	0.39	NIT 1		
Fusobacterium mortiferum		3.31	0.16			
77011010	**Good identification**					**77011010**
Eubacterium saburreum		100.0	0.37	NIT 1		
77011100	**Good identification**					**77011100**
Eubacterium saburreum		96.69	0.45	NIT 1		
Fusobacterium mortiferum		3.31	0.22			
77011110	**Good identification**					**77011110**
Eubacterium saburreum		100.0	0.44	NIT 1		
77011520	**Good identification**					**77011520**
Odoribacter splanchnicus		100.0	0.56	SUC 1		
77011560	**Good identification**					**77011560**
Odoribacter splanchnicus		100.0	0.39	SUC 1		
77015100	**Good identification**					**77015100**
Eubacterium saburreum		100.0	0.41	NIT 1		
77015110	**Good identification**					**77015110**
Eubacterium saburreum		100.0	0.39	NIT 1		
77031000	**Good identification**					**77031000**
Eubacterium saburreum		100.0	0.36	NIT 1		
77031100	**Good identification**					**77031100**
Eubacterium saburreum		98.49	0.42	NIT 1		
Fusobacterium mortiferum		1.51	0.13			
77031110	**Good identification**					**77031110**
Eubacterium saburreum		100.0	0.41	NIT 1		
77035100	**Good identification**					**77035100**
Eubacterium saburreum		100.0	0.38	NIT 1		
77035110	**Good identification**					**77035110**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Eubacterium saburreum		100.0	0.36	NIT 1			
77101000	**Good identification**						**77101000**
					SPO		
Eubacterium saburreum		98.58	0.41	NIT 1	1		
Clostridium bifermentans		1.42	-.03		99		
77101010	**Good identification**						**77101010**
Eubacterium saburreum		100.0	0.40	NIT 1			
77101100	**Low discrimination**						**77101100**
					SPO		
Eubacterium saburreum		83.77	0.47	NIT 1	1		
Clostridium chauvoei		16.23	0.25		99		
77101101	**Very good identification**						**77101101**
					SPO		
Clostridium chauvoei		99.95	0.60	IND 1	99		
Eubacterium saburreum		0.05	0.13		1		
77101103	**Good identification**						**77101103**
Clostridium chauvoei		100.0	0.47	IND 1			
77101110	**Good identification**						**77101110**
Eubacterium saburreum		100.0	0.46	NIT 1			
77101121	**Good identification**						**77101121**
Clostridium chauvoei		100.0	0.51	IND 1			
77101123	**Good identification**						**77101123**
Clostridium chauvoei		100.0	0.39	IND 1			
77105000	**Good identification**						**77105000**
Eubacterium saburreum		100.0	0.37	NIT 1			
77105010	**Good identification**						**77105010**
Eubacterium saburreum		100.0	0.35	NIT 1			
77105100	**Good identification**						**77105100**
Eubacterium saburreum		100.0	0.43	NIT 1			
77105110	**Good identification**						**77105110**
Eubacterium saburreum		100.0	0.41	NIT 1			
77111000	**Low discrimination**						**77111000**
					GRA		
Eubacterium saburreum		89.57	0.49	NIT 1	85		
Prevotella intermedia		9.92	0.23		1		
Clostridium perfringens		0.51	0.02		99		
77111002	**Acceptable identification**						**77111002**
					GRA	CAT	
Eubacterium saburreum		92.19	0.36	NIT 1	85	1	
Actinomyces viscosus		4.69	0.03		99	76	
Capnocytophaga ochracea		3.12	-.01		1	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
77111010	**Good identification**						**77111010**
Eubacterium saburreum		100.0	0.48	NIT 1			
77111100	**Low discrimination**						**77111100**
					SPO	CAT	
Eubacterium saburreum		83.57	0.56	NIT 1	1	1	
Clostridium chauvoei		12.60	0.32		99	1	
Capnocytophaga ochracea		2.35	0.17		1	90	
77111101	**Very good identification**						**77111101**
					SPO		
Clostridium chauvoei		99.93	0.66	IND 1	99		
Eubacterium saburreum		0.07	0.21		1		
77111103	**Good identification**						**77111103**
					GRA	SPO	
Clostridium chauvoei		99.82	0.53	IND 1	99	99	
Capnocytophaga ochracea		0.10	-.01		1	1	
Actinomyces viscosus		0.09	-.01		99	1	
77111110	**Good identification**						**77111110**
					CAT		
Eubacterium saburreum		99.62	0.54	NIT 1	1		
Capnocytophaga ochracea		0.38	0.01		90		
77111112	**Low discrimination**						**77111112**
					CAT		
Eubacterium saburreum		83.70	0.41	NIT 1	1		
Capnocytophaga ochracea		16.30	0.17		90		
77111120	**Low discrimination**						**77111120**
					CAT	GRA	
Clostridium perfringens		53.18	0.42	IND 1	1	99	
Capnocytophaga ochracea		23.45	0.33		90	1	
Eubacterium saburreum		16.35	0.43		1	85	
77111121	**Very good identification**						**77111121**
					GRA	CAT	
Clostridium chauvoei		99.83	0.58	IND 1	99	1	
Clostridium perfringens		0.12	0.08		99	1	
Capnocytophaga ochracea		0.05	-.01		1	90	
77111122	**Low discrimination**						**77111122**
					CAT	GRA	
Capnocytophaga ochracea		87.75	0.50	IND 1	90	1	
Clostridium perfringens		10.41	0.37	RAF 90	1	99	
Eubacterium saburreum		1.20	0.30		1	85	
77111123	**Good identification**						**77111123**
					GRA	CAT	
Clostridium chauvoei		97.15	0.45	IND 1	99	1	
Capnocytophaga ochracea		2.55	0.15		1	90	
Clostridium perfringens		0.30	0.03		99	1	
77111130	**Low discrimination**						**77111130**
					CAT	GRA	
Eubacterium saburreum		80.98	0.41	NIT 1	1	85	
Capnocytophaga ochracea		15.77	0.17		90	1	
Clostridium perfringens		3.25	0.08		1	99	
77111522	**Good identification**						**77111522**
					GRA	CAT	

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
Capnocytophaga ochracea	99.33	0.37	IND 1	1	90	
Clostridium perfringens	0.67	0.03	RAF 90	99	1	
77115000 Good identification						**77115000**
Eubacterium saburreum	100.0	0.45	NIT 1			
77115010 Good identification						**77115010**
Eubacterium saburreum	100.0	0.43	NIT 1			
77115100 Good identification						**77115100**
				CAT		
Eubacterium saburreum	98.71	0.51	NIT 1	1		
Capnocytophaga ochracea	1.29	0.07		90		
77115102 Low discrimination						**77115102**
				CAT		
Eubacterium saburreum	60.03	0.38	NIT 1	1		
Capnocytophaga ochracea	39.97	0.23		90		
77115110 Good identification						**77115110**
Eubacterium saburreum	100.0	0.49	NIT 1			
77115112 Acceptable identification						**77115112**
				CAT		
Eubacterium saburreum	91.71	0.37	NIT 1	1		
Capnocytophaga ochracea	8.29	0.07		90		
77115120 Low discrimination						**77115120**
				CAT	GRA	
Eubacterium saburreum	57.91	0.38	NIT 1	1	85	
Capnocytophaga ochracea	38.56	0.23		90	1	
Clostridium perfringens	3.53	0.08		1	99	
77115122 Acceptable identification						**77115122**
				CAT		
Capnocytophaga ochracea	93.18	0.39	IND 1	90		
Bacteroides fragilis	3.63	0.06	RAF 90	99		
Eubacterium saburreum	2.74	0.25		1		
77115130 Acceptable identification						**77115130**
				CAT		
Eubacterium saburreum	91.71	0.37	NIT 1	1		
Capnocytophaga ochracea	8.29	0.07		90		
77121000 Good identification						**77121000**
Eubacterium saburreum	100.0	0.38	NIT 1			
77121010 Good identification						**77121010**
Eubacterium saburreum	100.0	0.37	NIT 1			
77121100 Good identification						**77121100**
				SPO		
Eubacterium saburreum	96.87	0.44	NIT 1	1		
Clostridium chauvoei	3.13	0.09		99		
77121101 Good identification						**77121101**
				SPO		
Clostridium chauvoei	99.69	0.43	IND 1	99		
Eubacterium saburreum	0.32	0.10	TRE 10	1		
77121110 Good identification						**77121110**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Eubacterium saburreum		100.0	0.43	NIT 1		
77121121	**Good identification**					**77121121**
Clostridium chauvoei		100.0	0.35	IND 1		
77125100	**Good identification**					**77125100**
Eubacterium saburreum		100.0	0.40	NIT 1		
77125110	**Good identification**					**77125110**
Eubacterium saburreum		100.0	0.38	NIT 1		
77131000	**Good identification**					**77131000**
Eubacterium saburreum		98.45	0.46	NIT 1		
Clostridium perfringens		1.55	0.06			
77131010	**Good identification**					**77131010**
Eubacterium saburreum		100.0	0.45	NIT 1		
77131100	**Low discrimination**					**77131100**
					COC	
Eubacterium saburreum		89.79	0.53	NIT 1	1	
Clostridium perfringens		4.44	0.21		1	
Atopobium parvulum		3.52	0.19		99	
77131101	**Good identification**					**77131101**
					SPO	
Clostridium chauvoei		99.60	0.50	IND 1	99	
Eubacterium saburreum		0.41	0.18	TRE 10	1	
77131102	**Low discrimination**					**77131102**
					COC	
Eubacterium saburreum		82.11	0.40	NIT 1	1	
Clostridium perfringens		10.83	0.16		1	
Atopobium parvulum		3.22	0.06		99	
77131103	**Good identification**					**77131103**
Clostridium chauvoei		100.0	0.37	IND 1		
77131110	**Good identification**					**77131110**
Eubacterium saburreum		100.0	0.51	NIT 1		
77131112	**Good identification**					**77131112**
Eubacterium saburreum		100.0	0.38	NIT 1		
77131120	**Low discrimination**					**77131120**
					SPO	
Clostridium perfringens		89.10	0.47	IND 1	84	
Eubacterium saburreum		9.83	0.40		1	
Clostridium chauvoei		0.47	0.07		99	
77131121	**Good identification**					**77131121**
Clostridium chauvoei		98.09	0.41	IND 1		
Clostridium perfringens		1.91	0.13	TRE 10		
77131122	**Acceptable identification**					**77131122**
					CAT GRA	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		91.98	0.41	IND 1	1	99	
Capnocytophaga ochracea		4.22	0.15		90	1	
Eubacterium saburreum		3.81	0.27		1	85	
77131130	**Low discrimination**						**77131130**
Eubacterium saburreum		89.94	0.38	NIT 1			
Clostridium perfringens		10.06	0.13				
77135000	**Good identification**						**77135000**
Eubacterium saburreum		100.0	0.42	NIT 1			
77135010	**Good identification**						**77135010**
Eubacterium saburreum		100.0	0.40	NIT 1			
77135100	**Low discrimination**						**77135100**
					COC		
Eubacterium saburreum		88.10	0.48	NIT 1	1		
Atopobium parvulum		11.90	0.23		99		
77135102	**Low discrimination**						**77135102**
					COC		
Eubacterium saburreum		88.10	0.35	NIT 1	1		
Atopobium parvulum		11.90	0.10		99		
77135110	**Good identification**						**77135110**
Eubacterium saburreum		100.0	0.46	NIT 1			
77135120	**Low discrimination**						**77135120**
					COC		
Eubacterium saburreum		76.62	0.35	NIT 1	1		
Clostridium perfringens		13.03	0.13		1		
Atopobium parvulum		10.35	0.10		99		
77151122	**Good identification**						**77151122**
					GRA	CAT	
Capnocytophaga ochracea		95.56	0.39	IND 1	1	90	
Clostridium perfringens		1.89	0.13	RAF 90	99	1	
Prevotella oralis		1.77	0.09		1	1	
77331120	**Good identification**						**77331120**
					COC		
Clostridium perfringens		97.02	0.37	IND 1	1		
Atopobium parvulum		2.22	0.08		99		
Actinomyces naeslundii		0.75	0.23		1		
77355522	**Good identification**						**77355522**
Bacteroides uniformis		100.0	0.38	NIT 1			
77355722	**Good identification**						**77355722**
					CAT		
Bacteroides uniformis		97.56	0.40	NIT 1	5		
Bacteroides ovatus		2.15	0.15	RAF 97	68		
Bacteroides thetaiotaomicron		0.29	0.00		89		
77357100	**Good identification**						**77357100**
Actinomyces israelii		100.0	0.38	IND 1			
77357102	**Good identification**						**77357102**
					SPO		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		96.00	0.41	IND 1	1		
Clostridium tertium		4.00	0.03		99		
77357500	**Good identification**						**77357500**
Actinomyces israelii		100.0	0.36	IND 1			
77357502	**Good identification**						**77357502**
Actinomyces israelii		95.41	0.39	IND 1			
Bifidobacterium adolescentis		4.59	0.11				
77371112	**Good identification**						**77371112**
Leptotrichia buccalis		100.0	0.50	IND 1			
77375102	**Low discrimination**						**77375102**
					COC	SPO	
Actinomyces israelii		64.70	0.38	IND 1	1	1	
Atopobium parvulum		31.96	0.19		99	1	
Clostridium butyricum		3.34	-.05		1	99	
77375502	**Low discrimination**						**77375502**
					GRA	SPO	
Actinomyces israelii		58.77	0.36	IND 1	99	1	
Clostridium butyricum		33.37	0.11		99	99	
Mitsuokella multacida		4.40	0.04		1	1	
77377000	**Good identification**						**77377000**
Actinomyces israelii		100.0	0.37	IND 1			
77377002	**Good identification**						**77377002**
Actinomyces israelii		100.0	0.40	IND 1			
77377100	**Good identification**						**77377100**
					SPO		
Actinomyces israelii		95.25	0.48	IND 1	1		
Clostridium tertium		4.75	0.11		99		
77377102	**Acceptable identification**						**77377102**
					SPO		
Actinomyces israelii		90.93	0.51	IND 1	1		
Clostridium tertium		9.07	0.19		99		
77377140	**Good identification**						**77377140**
Actinomyces israelii		100.0	0.41	IND 1			
77377142	**Good identification**						**77377142**
Actinomyces israelii		100.0	0.44	IND 1			
77377400	**Good identification**						**77377400**
Actinomyces israelii		100.0	0.36	IND 1			
77377402	**Good identification**						**77377402**
Actinomyces israelii		100.0	0.39	IND 1			
77377500	**Low discrimination**						**77377500**
					GRA	SPO	
Actinomyces israelii		68.53	0.46	IND 1	99	1	
Mitsuokella multacida		28.53	0.27		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		2.94	0.02		99	99	
77377502	**Good identification**						**77377502**
					GRA	SPO	
Actinomyces israelii		95.19	0.49	IND 1	99	1	
Mitsuokella multacida		2.94	0.11		1	1	
Clostridium butyricum		1.06	-.05		99	99	
77377512	**Low discrimination**						**77377512**
Actinomyces israelii		61.37	0.35	IND 1			
Bifidobacterium dentium		30.13	0.06	MLZ 13			
Bifidobacterium adolescentis		4.28	0.09				
77377542	**Acceptable identification**						**77377542**
					GRA		
Actinomyces israelii		92.65	0.42	IND 1	99		
Mitsuokella multacida		7.35	0.11		1		
77411100	**Low discrimination**						**77411100**
					CAT		
Eubacterium saburreum		87.53	0.41	NIT 1	1		
Fusobacterium mortiferum		10.81	0.27		5		
Capnocytophaga ochracea		1.66	-.01		90		
77411110	**Good identification**						**77411110**
Eubacterium saburreum		100.0	0.39	NIT 1			
77415100	**Good identification**						**77415100**
Eubacterium saburreum		100.0	0.36	NIT 1			
77431100	**Acceptable identification**						**77431100**
Eubacterium saburreum		94.76	0.38	NIT 1			
Fusobacterium mortiferum		5.24	0.18				
77431110	**Good identification**						**77431110**
Eubacterium saburreum		100.0	0.36	NIT 1			
77435722	**Good identification**						**77435722**
Bacteroides thetaiotaomicron		99.38	0.40	NIT 1			
Parabacteroides distasonis		0.62	0.18	MLT 94			
77475722	**Good identification**						**77475722**
					CAT		
Bacteroides thetaiotaomicron		97.59	0.45	NIT 1	89		
Bacteroides ovatus		2.09	0.17	MLT 94	68		
Bacteroides uniformis		0.17	-.05		5		
77475732	**Good identification**						**77475732**
Bacteroides thetaiotaomicron		96.47	0.38	NIT 1			
Bacteroides ovatus		3.54	0.14	MLT 94			
77501000	**Good identification**						**77501000**
Eubacterium saburreum		100.0	0.37	NIT 1			
77501010	**Good identification**						**77501010**
Eubacterium saburreum		100.0	0.35	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
77501100	**Good identification**						**77501100**
					CAT		
Eubacterium saburreum		98.60	0.43	NIT 1	1		
Capnocytophaga ochracea		1.40	-.01		90		
77501110	**Good identification**						**77501110**
Eubacterium saburreum		100.0	0.41	NIT 1			
77505100	**Good identification**						**77505100**
Eubacterium saburreum		100.0	0.38	NIT 1			
77505110	**Good identification**						**77505110**
Eubacterium saburreum		100.0	0.37	NIT 1			
77511000	**Low discrimination**						**77511000**
					CAT		
Eubacterium saburreum		75.78	0.45	NIT 1	1		
Prevotella intermedia		23.38	0.26		1		
Capnocytophaga ochracea		0.84	-.01		90		
77511010	**Good identification**						**77511010**
Eubacterium saburreum		100.0	0.43	NIT 1			
77511100	**Low discrimination**						**77511100**
					GRA	CAT	
Eubacterium saburreum		67.08	0.51	NIT 1	85	1	
Capnocytophaga ochracea		31.53	0.33		1	90	
Actinomyces naeslundii		0.43	0.27		99	25	
77511102	**Acceptable identification**						**77511102**
					GRA	CAT	
Capnocytophaga ochracea		91.64	0.50	IND 1	1	90	
Actinomyces viscosus		4.32	0.28	NAG 90	99	76	
Eubacterium saburreum		3.82	0.38		85	1	
77511110	**Acceptable identification**						**77511110**
					CAT		
Eubacterium saburreum		94.00	0.49	NIT 1	1		
Capnocytophaga ochracea		6.00	0.17		90		
77511120	**Low discrimination**						**77511120**
					GRA	CAT	
Capnocytophaga ochracea		83.88	0.50	IND 1	1	90	
Prevotella melaninogenica		8.46	0.29	BGL 90	1	1	
Clostridium perfringens		4.03	0.30		99	1	
77511122	**Very good identification**						**77511122**
					GRA	CAT	
Capnocytophaga ochracea		99.15	0.66	IND 1	1	90	
Clostridium perfringens		0.25	0.24		99	1	
Prevotella melaninogenica		0.21	0.16		1	1	
77511132	**Good identification**						**77511132**
					CAT		
Capnocytophaga ochracea		99.40	0.50	IND 1	90		
Eubacterium saburreum		0.60	0.24	MLZ 10	1		
77511162	**Good identification**						**77511162**
					GRA	CAT	
Capnocytophaga ochracea		99.57	0.50	IND 1	1	90	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium perfringens		0.43	0.12	SOR 10	99	1	
77511502	**Good identification**						**77511502**
					GRA	CAT	
Capnocytophaga ochracea		99.18	0.37	IND 1	1	90	
Bifidobacterium longum subsp. longum		0.82	0.09	NAG 90	99	1	
77511520	**Good identification**						**77511520**
					GRA	CAT	
Capnocytophaga ochracea		96.26	0.37	IND 1	1	90	
Bifidobacterium longum subsp. longum		3.18	0.20	BGL 90	99	1	
Prevotella melaninogenica		0.56	-.06		1	1	
77511522	**Good identification**						**77511522**
					GRA	CAT	
Capnocytophaga ochracea		99.53	0.53	IND 1	1	90	
Bifidobacterium longum subsp. longum		0.37	0.20		99	1	
Bacteroides thetaiotaomicron		0.10	-.02		1	89	
77511532	**Acceptable identification**						**77511532**
					GRA	CAT	
Capnocytophaga ochracea		94.67	0.37	IND 1	1	90	
Bifidobacterium longum subsp. longum		5.33	0.24	MLZ 10	99	1	
77511562	**Good identification**						**77511562**
Capnocytophaga ochracea		100.0	0.37	IND 1			
77515000	**Good identification**						**77515000**
Eubacterium saburreum		100.0	0.40	NIT 1			
77515010	**Good identification**						**77515010**
Eubacterium saburreum		100.0	0.39	NIT 1			
77515100	**Low discrimination**						**77515100**
					CAT		
Eubacterium saburreum		82.09	0.46	NIT 1	1		
Capnocytophaga ochracea		17.91	0.23		90		
77515102	**Low discrimination**						**77515102**
					CAT		
Capnocytophaga ochracea		88.59	0.39	IND 1	90		
Eubacterium saburreum		7.96	0.33	NAG 90	1		
Bacteroides fragilis		3.45	0.06		99		
77515110	**Good identification**						**77515110**
					CAT		
Eubacterium saburreum		97.12	0.45	NIT 1	1		
Capnocytophaga ochracea		2.88	0.07		90		
77515120	**Low discrimination**						**77515120**
					CAT		
Capnocytophaga ochracea		76.70	0.39	IND 1	90		
Bacteroides fragilis		15.58	0.18	BGL 90	99		
Eubacterium saburreum		6.89	0.33		1		
77515122	**Low discrimination**						**77515122**
					CAT		
Capnocytophaga ochracea		69.79	0.56	IND 1	90		
Bacteroides fragilis		29.92	0.40		99		
Eubacterium saburreum		0.12	0.21		1		
77515132	**Good identification**						**77515132**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					CAT		
Capnocytophaga ochracea		95.06	0.39	IND 1	90		
Bacteroides fragilis		3.71	0.06	MLZ 10	99		
Eubacterium saburreum		1.23	0.19		1		
77515162	**Good identification**						**77515162**
Capnocytophaga ochracea		96.25	0.39	IND 1			
Bacteroides fragilis		3.75	0.06	SOR 10			
77515522	**Low discrimination**						**77515522**
					GRA	CAT	
Capnocytophaga ochracea		83.30	0.43	IND 1	1	90	
Bacteroides thetaiotaomicron		8.16	0.22		1	89	
Bifidobacterium longum subsp. longum		3.67	0.28		99	1	
77515722	**Acceptable identification**						**77515722**
					CAT		
Bacteroides thetaiotaomicron		94.73	0.41	NIT 1	89		
Bacteroides uniformis		3.41	0.14	TRE 93	5		
Capnocytophaga ochracea		0.74	0.09		90		
77521100	**Good identification**						**77521100**
Eubacterium saburreum		100.0	0.40	NIT 1			
77521110	**Good identification**						**77521110**
Eubacterium saburreum		100.0	0.38	NIT 1			
77525100	**Good identification**						**77525100**
Eubacterium saburreum		100.0	0.35	NIT 1			
77531000	**Good identification**						**77531000**
Eubacterium saburreum		98.49	0.42	NIT 1			
Actinomyces naeslundii		1.51	0.24				
77531010	**Good identification**						**77531010**
Eubacterium saburreum		100.0	0.40	NIT 1			
77531100	**Good identification**						**77531100**
Eubacterium saburreum		95.85	0.48	NIT 1			
Actinomyces naeslundii		1.79	0.32				
Clostridium perfringens		1.68	0.09				
77531102	**Low discrimination**						**77531102**
					GRA	CAT	
Eubacterium saburreum		60.86	0.35	NIT 1	85	1	
Capnocytophaga ochracea		22.11	0.15		1	90	
Actinomyces viscosus		7.76	0.09		99	76	
77531110	**Good identification**						**77531110**
Eubacterium saburreum		100.0	0.46	NIT 1			
77531722	**Good identification**						**77531722**
Bacteroides thetaiotaomicron		99.63	0.37	NIT 1			
Parabacteroides distasonis		0.37	0.11	XYL 96			
77533100	**Low discrimination**						**77533100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		56.17	0.37	IND 1			
Propionibacterium propionicum		28.17	0.21	ESL 12			
Eubacterium saburreum		15.67	0.14				
77533101	**Good identification**						**77533101**
Actinomyces naeslundii		100.0	0.40	IND 1			
77533102	**Low discrimination**						**77533102**
Actinomyces naeslundii		57.07	0.37	IND 1			
Propionibacterium propionicum		42.93	0.24	ESL 12			
77533103	**Good identification**						**77533103**
Actinomyces naeslundii		100.0	0.40	IND 1			
77533120	**Low discrimination**						**77533120**
Actinomyces naeslundii		86.38	0.37	IND 1			
Clostridium perfringens		13.62	0.01	ESL 12			
77533121	**Good identification**						**77533121**
Actinomyces naeslundii		100.0	0.40	IND 1			
77533122	**Low discrimination**						**77533122**
					GRA		
Actinomyces naeslundii		86.14	0.37	IND 1	99		
Capnocytophaga ochracea		13.86	-.03	ESL 12	1		
77533123	**Good identification**						**77533123**
Actinomyces naeslundii		100.0	0.40	IND 1			
77535000	**Good identification**						**77535000**
Eubacterium saburreum		100.0	0.37	NIT 1			
77535010	**Good identification**						**77535010**
Eubacterium saburreum		100.0	0.36	NIT 1			
77535100	**Good identification**						**77535100**
Eubacterium saburreum		100.0	0.43	NIT 1			
77535110	**Good identification**						**77535110**
Eubacterium saburreum		100.0	0.42	NIT 1			
77535522	**Good identification**						**77535522**
					GRA	CAT	
Bacteroides thetaiotaomicron		97.61	0.41	NIT 1	1	89	
Bifidobacterium longum subsp. longum		0.83	0.17	RHA 93	99	1	
Capnocytophaga ochracea		0.76	0.09		1	90	
77535722	**Very good identification**						**77535722**
Bacteroides thetaiotaomicron		99.57	0.60	NIT 1			
Bacteroides ovatus		0.29	0.17				
Parabacteroides distasonis		0.14	0.27				
77535732	**Good identification**						**77535732**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides thetaiotaomicron		99.26	0.53	NIT 1			
Bacteroides ovatus		0.49	0.14				
Parabacteroides distasonis		0.19	0.23				
77551102	**Acceptable identification**						**77551102**
					GRA	CAT	
Capnocytophaga ochracea		93.52	0.39	IND 1	1	90	
Actinomyces viscosus		4.68	0.18	NAG 90	99	76	
Prevotella oralis		1.05	0.06		1	1	
77551120	**Acceptable identification**						**77551120**
					CAT		
Capnocytophaga ochracea		93.28	0.39	IND 1	90		
Prevotella melaninogenica		4.18	0.12	BGL 90	1		
Prevotella oralis		1.04	0.06		1		
77551122	**Good identification**						**77551122**
					CAT		
Capnocytophaga ochracea		97.25	0.56	IND 1	90		
Prevotella oralis		2.30	0.27		1		
Bacteroides fragilis		0.18	-.01		99		
77551132	**Good identification**						**77551132**
Capnocytophaga ochracea		100.0	0.39	IND 1			
77551162	**Good identification**						**77551162**
Capnocytophaga ochracea		100.0	0.39	IND 1			
77551522	**Good identification**						**77551522**
					CAT		
Capnocytophaga ochracea		95.91	0.43	IND 1	90		
Bacteroides uniformis		3.25	0.11		5		
Bacteroides thetaiotaomicron		0.83	0.04		89		
77553101	**Good identification**						**77553101**
Actinomyces naeslundii		100.0	0.37	IND 1			
77553103	**Good identification**						**77553103**
Actinomyces naeslundii		100.0	0.37	IND 1			
77553121	**Good identification**						**77553121**
Actinomyces naeslundii		100.0	0.37	IND 1			
77553123	**Good identification**						**77553123**
Actinomyces naeslundii		100.0	0.37	IND 1			
77555122	**Low discrimination**						**77555122**
					CAT		
Capnocytophaga ochracea		51.21	0.45	IND 1	90		
Bacteroides fragilis		37.64	0.34		99		
Bacteroides uniformis		10.56	0.27		5		
77555522	**Low discrimination**						**77555522**
					CAT		
Bacteroides uniformis		87.17	0.45	NIT 1	5		
Capnocytophaga ochracea		6.49	0.32	SAL 92	90		
Bacteroides thetaiotaomicron		5.40	0.27		89		
77555722	**Low discrimination**						**77555722**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					CAT	
Bacteroides uniformis		58.71	0.48	NIT 1	5	
Bacteroides thetaiotaomicron		35.01	0.47	SAL 92	89	
Bacteroides ovatus		6.21	0.34		68	
77555732	**Low discrimination**					**77555732**
					CAT	
Bacteroides thetaiotaomicron		74.19	0.40	NIT 1	89	
Bacteroides ovatus		22.58	0.31	TRE 93	68	
Bacteroides uniformis		3.23	0.14		5	
77571101	**Good identification**					**77571101**
Actinomyces naeslundii		100.0	0.37	IND 1		
77571103	**Good identification**					**77571103**
Actinomyces naeslundii		100.0	0.37	IND 1		
77571121	**Good identification**					**77571121**
Actinomyces naeslundii		100.0	0.37	IND 1		
77571123	**Good identification**					**77571123**
Actinomyces naeslundii		100.0	0.37	IND 1		
77571722	**Good identification**					**77571722**
Bacteroides thetaiotaomicron		96.83	0.42	NIT 1		
Bacteroides ovatus		3.17	0.17	XYL 96		
77571732	**Acceptable identification**					**77571732**
Bacteroides thetaiotaomicron		94.68	0.35	NIT 1		
Bacteroides ovatus		5.32	0.14	XYL 96		
77573100	**Low discrimination**					**77573100**
Actinomyces naeslundii		74.96	0.39	IND 1		
Actinomyces israelii		25.05	0.21	ESL 12		
77573101	**Good identification**					**77573101**
Actinomyces naeslundii		100.0	0.42	IND 1		
77573102	**Low discrimination**					**77573102**
Actinomyces naeslundii		66.61	0.39	IND 1		
Actinomyces israelii		33.39	0.24	ESL 12		
77573103	**Good identification**					**77573103**
Actinomyces naeslundii		100.0	0.42	IND 1		
77573120	**Good identification**					**77573120**
Actinomyces naeslundii		100.0	0.39	IND 1		
77573121	**Good identification**					**77573121**
Actinomyces naeslundii		100.0	0.42	IND 1		
77573122	**Good identification**					**77573122**
Actinomyces naeslundii		100.0	0.39	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
77573123	**Good identification**					**77573123**
Actinomyces naeslundii		100.0	0.42	IND 1		
77575522	**Acceptable identification**					**77575522**
					CAT	
Bacteroides thetaiotaomicron		90.15	0.47	NIT 1	89	
Bacteroides ovatus		8.52	0.29	RHA 93	68	
Bacteroides uniformis		1.11	0.11		5	
77575532	**Low discrimination**					**77575532**
Bacteroides thetaiotaomicron		85.90	0.40	NIT 1		
Bacteroides ovatus		13.91	0.26	RHA 93		
Parabacteroides distasonis		0.19	0.10			
77575622	**Acceptable identification**					**77575622**
Bacteroides thetaiotaomicron		93.73	0.37	NIT 1		
Bacteroides ovatus		6.27	0.17	LAC 98		
77575722	**Low discrimination**					**77575722**
					CAT	
Bacteroides thetaiotaomicron		87.97	0.66	NIT 1	89	
Bacteroides ovatus		11.88	0.51		68	
Bacteroides uniformis		0.11	0.14		5	
77575732	**Low discrimination**					**77575732**
Bacteroides thetaiotaomicron		81.11	0.59	NIT 1		
Bacteroides ovatus		18.78	0.48			
Clostridium clostridioforme		0.07	0.17			
77577100	**Good identification**					**77577100**
Actinomyces israelii		100.0	0.39	IND 1		
77577102	**Good identification**					**77577102**
Actinomyces israelii		100.0	0.42	IND 1		
77577142	**Good identification**					**77577142**
Actinomyces israelii		100.0	0.35	IND 1		
77577502	**Low discrimination**					**77577502**
					GRA	
Actinomyces israelii		87.17	0.41	IND 1	99	
Mitsuokella multacida		8.22	0.11	SAL 92	1	
Bifidobacterium adolescentis		4.61	0.13		99	
77577722	**Low discrimination**					**77577722**
					CAT	
Bacteroides ovatus		81.52	0.43	NIT 1	68	
Bacteroides thetaiotaomicron		18.29	0.32	SAL 88	89	
Mitsuokella multacida		0.19	0.05		1	
77577732	**Low discrimination**					**77577732**
Bacteroides ovatus		88.43	0.40	NIT 1		
Bacteroides thetaiotaomicron		11.57	0.25	SAL 88		
77655522	**Good identification**					**77655522**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides uniformis		99.93	0.45	NIT 1	5		
Bacteroides thetaiotaomicron		0.07	-.06	MLT 92	89		
77655722	**Good identification**						**77655722**
					CAT		
Bacteroides uniformis		98.57	0.48	NIT 1	5		
Bacteroides ovatus		0.77	0.15	MLT 92	68		
Bacteroides thetaiotaomicron		0.66	0.13		89		
77711101	**Good identification**						**77711101**
Actinomyces naeslundii		100.0	0.39	IND 1			
77711103	**Low discrimination**						**77711103**
Actinomyces naeslundii		61.59	0.39	IND 1			
Actinomyces viscosus		31.90	0.10	ESL 12			
Bifidobacterium longum subsp. infantis		6.52	-.07				
77711121	**Good identification**						**77711121**
Actinomyces naeslundii		100.0	0.39	IND 1			
77711123	**Low discrimination**						**77711123**
					GRA	CAT	
Actinomyces naeslundii		76.79	0.39	IND 1	99	25	
Bifidobacterium longum subsp. infantis		14.46	-.03	ESL 12	99	1	
Capnocytophaga ochracea		8.75	-.03		1	90	
77713001	**Good identification**						**77713001**
Actinomyces naeslundii		100.0	0.36	IND 1			
77713003	**Good identification**						**77713003**
Actinomyces naeslundii		100.0	0.36	IND 1			
77713021	**Good identification**						**77713021**
Actinomyces naeslundii		100.0	0.36	IND 1			
77713023	**Good identification**						**77713023**
Actinomyces naeslundii		100.0	0.36	IND 1			
77713100	**Good identification**						**77713100**
Actinomyces naeslundii		100.0	0.41	IND 1			
77713101	**Good identification**						**77713101**
Actinomyces naeslundii		100.0	0.44	IND 1			
77713102	**Low discrimination**						**77713102**
Actinomyces naeslundii		85.80	0.41	IND 1			
Actinomyces israelii		7.18	0.13	ESL 12			
Bifidobacterium longum subsp. infantis		7.02	-.07				
77713103	**Good identification**						**77713103**
Actinomyces naeslundii		100.0	0.44	IND 1			
77713120	**Good identification**						**77713120**
Actinomyces naeslundii		100.0	0.41	IND 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
77713121	**Good identification**						**77713121**
Actinomyces naeslundii		100.0	0.44	IND 1			
77713122	**Low discrimination**						**77713122**
					GRA	CAT	
Actinomyces naeslundii		81.07	0.41	IND 1	99	25	
Bifidobacterium longum subsp. infantis		11.79	-.03	ESL 12	99	1	
Capnocytophaga ochracea		7.14	-.03		1	90	
77713123	**Good identification**						**77713123**
Actinomyces naeslundii		100.0	0.44	IND 1			
77731001	**Good identification**						**77731001**
Actinomyces naeslundii		100.0	0.36	IND 1			
77731003	**Good identification**						**77731003**
Actinomyces naeslundii		100.0	0.36	IND 1			
77731021	**Good identification**						**77731021**
Actinomyces naeslundii		100.0	0.36	IND 1			
77731023	**Good identification**						**77731023**
Actinomyces naeslundii		100.0	0.36	IND 1			
77731100	**Low discrimination**						**77731100**
Actinomyces naeslundii		86.75	0.41	IND 1			
Eubacterium saburreum		13.25	0.14	ESL 12			
77731101	**Good identification**						**77731101**
Actinomyces naeslundii		100.0	0.44	IND 1			
77731102	**Low discrimination**						**77731102**
Actinomyces naeslundii		85.31	0.41	IND 1			
Actinomyces viscosus		7.71	-.01	ESL 12			
Bifidobacterium longum subsp. infantis		6.98	-.07				
77731103	**Good identification**						**77731103**
Actinomyces naeslundii		100.0	0.44	IND 1			
77731121	**Good identification**						**77731121**
Actinomyces naeslundii		100.0	0.44	IND 1			
77731123	**Good identification**						**77731123**
Actinomyces naeslundii		100.0	0.44	IND 1			
77733000	**Good identification**						**77733000**
Actinomyces naeslundii		100.0	0.38	IND 1			
77733001	**Good identification**						**77733001**
Actinomyces naeslundii		100.0	0.41	IND 1			
77733002	**Good identification**						**77733002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.38	IND 1		
77733003	**Good identification**					**77733003**
Actinomyces naeslundii		100.0	0.41	IND 1		
77733020	**Good identification**					**77733020**
Actinomyces naeslundii		100.0	0.38	IND 1		
77733021	**Good identification**					**77733021**
Actinomyces naeslundii		100.0	0.41	IND 1		
77733022	**Good identification**					**77733022**
Actinomyces naeslundii		100.0	0.38	IND 1		
77733023	**Good identification**					**77733023**
Actinomyces naeslundii		100.0	0.41	IND 1		
77733100	**Acceptable identification**					**77733100**
Actinomyces naeslundii		90.24	0.46	IND 1		
Actinomyces israelii		9.76	0.20	ESL 12		
77733101	**Good identification**					**77733101**
Actinomyces naeslundii		100.0	0.49	IND 1		
77733102	**Low discrimination**					**77733102**
Actinomyces naeslundii		86.04	0.46	IND 1		
Actinomyces israelii		13.96	0.23	ESL 12		
77733103	**Good identification**					**77733103**
Actinomyces naeslundii		100.0	0.49	IND 1		
77733120	**Good identification**					**77733120**
Actinomyces naeslundii		100.0	0.46	IND 1		
77733121	**Good identification**					**77733121**
Actinomyces naeslundii		100.0	0.49	IND 1		
77733122	**Good identification**					**77733122**
Actinomyces naeslundii		100.0	0.46	IND 1		
77733123	**Good identification**					**77733123**
Actinomyces naeslundii		100.0	0.49	IND 1		
77733141	**Good identification**					**77733141**
Actinomyces naeslundii		100.0	0.36	IND 1		
77733143	**Good identification**					**77733143**
Actinomyces naeslundii		100.0	0.36	IND 1		
77733161	**Good identification**					**77733161**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		100.0	0.36	IND 1		
77733163	**Good identification**					**77733163**
Actinomyces naeslundii		100.0	0.36	IND 1		
77735722	**Low discrimination**					**77735722**
Bacteroides thetaiotaomicron		88.27	0.47	NIT 1		
Bacteroides ovatus		10.64	0.32			
Parabacteroides distasonis		0.76	0.28			
77735732	**Low discrimination**					**77735732**
Bacteroides thetaiotaomicron		80.53	0.40	NIT 1		
Bacteroides ovatus		16.64	0.29			
Clostridium clostridioforme		1.87	0.23			
77737100	**Good identification**					**77737100**
Actinomyces israelii		100.0	0.38	IND 1		
77737102	**Good identification**					**77737102**
Actinomyces israelii		100.0	0.41	IND 1		
77737502	**Low discrimination**					**77737502**
					GRA	
Actinomyces israelii		78.63	0.40	IND 1	99	
Bifidobacterium adolescentis		12.81	0.20	CEL 93	99	
Mitsuokella multacida		8.56	0.11		1	
77751100	**Low discrimination**					**77751100**
Actinomyces naeslundii		52.29	0.38	IND 1		
Collinsella aerofaciens		29.07	-.03	ESL 12		
Actinomyces israelii		10.02	0.17			
77751101	**Good identification**					**77751101**
Actinomyces naeslundii		100.0	0.41	IND 1		
77751103	**Acceptable identification**					**77751103**
Actinomyces naeslundii		90.59	0.41	IND 1		
Actinomyces viscosus		9.41	0.00	ESL 12		
77751121	**Good identification**					**77751121**
Actinomyces naeslundii		100.0	0.41	IND 1		
77751123	**Acceptable identification**					**77751123**
					GRA	
Actinomyces naeslundii		91.80	0.41	IND 1	99	
Prevotella oralis		8.20	-.03	ESL 12	1	
77753000	**Low discrimination**					**77753000**
Actinomyces naeslundii		71.53	0.35	IND 1		
Actinomyces israelii		28.47	0.19	ESL 12		
77753001	**Good identification**					**77753001**
Actinomyces naeslundii		100.0	0.38	IND 1		
77753002	**Low discrimination**					**77753002**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Actinomyces naeslundii		62.62	0.35	IND 1		
Actinomyces israelii		37.38	0.22	ESL 12		
77753003	**Good identification**					**77753003**
Actinomyces naeslundii		100.0	0.38	IND 1		
77753020	**Good identification**					**77753020**
Actinomyces naeslundii		100.0	0.35	IND 1		
77753021	**Good identification**					**77753021**
Actinomyces naeslundii		100.0	0.38	IND 1		
77753022	**Good identification**					**77753022**
Actinomyces naeslundii		100.0	0.35	IND 1		
77753023	**Good identification**					**77753023**
Actinomyces naeslundii		100.0	0.38	IND 1		
77753100	**Low discrimination**					**77753100**
Actinomyces naeslundii		60.82	0.43	IND 1		
Actinomyces israelii		34.01	0.30	ESL 12		
Bifidobacterium breve		5.17	0.10			
77753101	**Good identification**					**77753101**
Actinomyces naeslundii		100.0	0.46	IND 1		
77753102	**Low discrimination**					**77753102**
Actinomyces naeslundii		37.07	0.43	IND 1		
Bifidobacterium breve		31.84	0.27	ESL 12		
Actinomyces israelii		31.09	0.33			
77753103	**Good identification**					**77753103**
Actinomyces naeslundii		100.0	0.46	IND 1		
77753120	**Low discrimination**					**77753120**
					SPO	
Actinomyces naeslundii		81.38	0.43	IND 1	1	
Bifidobacterium breve		12.29	0.15	ESL 12	1	
Clostridium ramosum		6.34	-.01		92	
77753121	**Good identification**					**77753121**
Actinomyces naeslundii		100.0	0.46	IND 1		
77753123	**Good identification**					**77753123**
Actinomyces naeslundii		100.0	0.46	IND 1		
77755102	**Low discrimination**					**77755102**
					GRA	
Actinomyces israelii		56.83	0.38	IND 1	99	
Bacteroides uniformis		32.54	0.11		1	
Bifidobacterium adolescentis		7.07	0.16		99	
77755122	**Good identification**					**77755122**
					CAT	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Bacteroides uniformis		98.04	0.45	NIT 1	5	
Prevotella oralis		0.86	0.16	ARA 92	1	
Capnocytophaga ochracea		0.42	0.11		90	
77755322	**Good identification**					**77755322**
Bacteroides uniformis		97.19	0.48	NIT 1		
Bacteroides ovatus		2.33	0.23	ARA 92		
Prevotella oralis		0.26	0.10			
77755522	**Very good identification**					**77755522**
					CAT	
Bacteroides uniformis		99.40	0.63	NIT 1	5	
Bacteroides ovatus		0.49	0.27		68	
Bacteroides thetaiotaomicron		0.10	0.15		89	
77755622	**Good identification**					**77755622**
					CAT	
Bacteroides uniformis		95.92	0.37	NIT 1	5	
Bacteroides ovatus		3.20	0.15	LAC 98	68	
Bacteroides thetaiotaomicron		0.88	0.05		89	
77755722	**Good identification**					**77755722**
					CAT	
Bacteroides uniformis		92.87	0.66	NIT 1	5	
Bacteroides ovatus		6.27	0.49		68	
Bacteroides thetaiotaomicron		0.85	0.34		89	
77755732	**Low discrimination**					**77755732**
					CAT	
Bacteroides ovatus		76.41	0.46	NIT 1	68	
Bacteroides uniformis		17.15	0.32	TRE 91	5	
Bacteroides thetaiotaomicron		6.04	0.27		89	
77757000	**Good identification**					**77757000**
Actinomyces israelii		100.0	0.37	IND 1		
77757002	**Good identification**					**77757002**
Actinomyces israelii		100.0	0.40	IND 1		
77757100	**Good identification**					**77757100**
Actinomyces israelii		100.0	0.48	IND 1		
77757102	**Good identification**					**77757102**
Actinomyces israelii		98.23	0.51	IND 1		
Bifidobacterium adolescentis		1.77	0.15			
77757112	**Low discrimination**					**77757112**
Actinomyces israelii		87.18	0.37	IND 1		
Bifidobacterium adolescentis		12.82	0.16	MLZ 13		
77757140	**Good identification**					**77757140**
Actinomyces israelii		100.0	0.41	IND 1		
77757142	**Good identification**					**77757142**
Actinomyces israelii		100.0	0.44	IND 1		
77757400	**Good identification**					**77757400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		100.0	0.36	IND 1			
77757402	**Good identification**						**77757402**
Actinomyces israelii		100.0	0.39	IND 1			
77757500	**Low discrimination**						**77757500**
					GRA	SPO	
Actinomyces israelii		68.10	0.46	IND 1	99	1	
Mitsuokella multacida		28.35	0.27		1	1	
Clostridium butyricum		3.55	0.03		99	99	
77757502	**Low discrimination**						**77757502**
					GRA		
Actinomyces israelii		82.94	0.49	IND 1	99		
Bifidobacterium adolescentis		12.19	0.29		99		
Mitsuokella multacida		2.56	0.11		1		
77757542	**Low discrimination**						**77757542**
					GRA		
Actinomyces israelii		88.81	0.42	IND 1	99		
Mitsuokella multacida		7.04	0.11		1		
Bifidobacterium adolescentis		4.15	0.13		99		
77757722	**Low discrimination**						**77757722**
					CAT		
Bacteroides ovatus		68.66	0.41	NIT 1	68		
Bacteroides uniformis		30.83	0.32	TRE 91	5		
Bacteroides thetaiotaomicron		0.28	0.00		89		
77757732	**Good identification**						**77757732**
Bacteroides ovatus		99.32	0.38	NIT 1			
Bacteroides uniformis		0.68	-.03	TRE 91			
77771000	**Low discrimination**						**77771000**
Actinomyces naeslundii		79.11	0.35	IND 1			
Actinomyces israelii		20.90	0.16	ESL 12			
77771001	**Good identification**						**77771001**
Actinomyces naeslundii		100.0	0.38	IND 1			
77771002	**Low discrimination**						**77771002**
Actinomyces naeslundii		71.62	0.35	IND 1			
Actinomyces israelii		28.38	0.19	ESL 12			
77771003	**Good identification**						**77771003**
Actinomyces naeslundii		100.0	0.38	IND 1			
77771020	**Good identification**						**77771020**
Actinomyces naeslundii		100.0	0.35	IND 1			
77771021	**Good identification**						**77771021**
Actinomyces naeslundii		100.0	0.38	IND 1			
77771022	**Good identification**						**77771022**
Actinomyces naeslundii		100.0	0.35	IND 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
77771023	**Good identification**					**77771023**
Actinomyces naeslundii		100.0	0.38	IND 1		
77771100	**Low discrimination**					**77771100**
					SPO	
Actinomyces naeslundii		66.49	0.43	IND 1	1	
Actinomyces israelii		24.68	0.27	ESL 12	1	
Clostridium butyricum		8.83	-.02		99	
77771101	**Good identification**					**77771101**
Actinomyces naeslundii		100.0	0.46	IND 1		
77771102	**Low discrimination**					**77771102**
					GRA	
Actinomyces naeslundii		29.44	0.43	IND 1	99	
Leptotrichia buccalis		23.42	0.07	ESL 12	1	
Bifidobacterium breve		16.86	0.24		99	
77771103	**Good identification**					**77771103**
Actinomyces naeslundii		100.0	0.46	IND 1		
77771112	**Good identification**					**77771112**
					GRA	
Leptotrichia buccalis		99.17	0.41	IND 1	1	
Bifidobacterium breve		0.72	0.24	NIT 10	99	
Actinomyces israelii		0.11	0.15		99	
77771120	**Low discrimination**					**77771120**
					SPO	
Actinomyces naeslundii		58.80	0.43	IND 1	1	
Clostridium paraputrificum		18.67	0.03	ESL 12	99	
Clostridium ramosum		12.66	0.06		92	
77771121	**Good identification**					**77771121**
Actinomyces naeslundii		100.0	0.46	IND 1		
77771123	**Good identification**					**77771123**
Actinomyces naeslundii		100.0	0.46	IND 1		
77773000	**Low discrimination**					**77773000**
Actinomyces naeslundii		56.46	0.40	IND 1		
Actinomyces israelii		43.54	0.29	ESL 12		
77773001	**Good identification**					**77773001**
Actinomyces naeslundii		100.0	0.43	IND 1		
77773003	**Good identification**					**77773003**
Actinomyces naeslundii		100.0	0.43	IND 1		
77773020	**Good identification**					**77773020**
Actinomyces naeslundii		100.0	0.40	IND 1		
77773021	**Good identification**					**77773021**
Actinomyces naeslundii		100.0	0.43	IND 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
77773022	**Good identification**					**77773022**
Actinomyces naeslundii		100.0	0.40	IND 1		
77773023	**Good identification**					**77773023**
Actinomyces naeslundii		100.0	0.43	IND 1		
77773100	**Low discrimination**					**77773100**
Actinomyces israelii		52.01	0.39	IND 1		
Actinomyces naeslundii		47.99	0.48	XYL 92		
77773101	**Good identification**					**77773101**
Actinomyces naeslundii		100.0	0.51	IND 1		
77773102	**Low discrimination**					**77773102**
Actinomyces israelii		55.66	0.42	IND 1		
Actinomyces naeslundii		34.24	0.48	XYL 92		
Bifidobacterium breve		10.10	0.24			
77773103	**Good identification**					**77773103**
Actinomyces naeslundii		100.0	0.51	IND 1		
77773120	**Low discrimination**					**77773120**
					SPO	
Actinomyces naeslundii		57.15	0.48	IND 1	1	
Clostridium ramosum		35.93	0.19	ESL 12	92	
Actinomyces israelii		3.95	0.19		1	
77773121	**Good identification**					**77773121**
Actinomyces naeslundii		100.0	0.51	IND 1		
77773123	**Good identification**					**77773123**
Actinomyces naeslundii		100.0	0.51	IND 1		
77773141	**Good identification**					**77773141**
Actinomyces naeslundii		100.0	0.38	IND 1		
77773142	**Low discrimination**					**77773142**
Actinomyces israelii		57.28	0.35	IND 1		
Bifidobacterium breve		26.73	0.24	XYL 92		
Actinomyces naeslundii		15.99	0.35			
77773143	**Good identification**					**77773143**
Actinomyces naeslundii		100.0	0.38	IND 1		
77773160	**Low discrimination**					**77773160**
Actinomyces naeslundii		77.29	0.35	IND 1		
Bifidobacterium breve		22.71	0.12	ESL 12		
77773161	**Good identification**					**77773161**
Actinomyces naeslundii		100.0	0.38	IND 1		
77773163	**Good identification**					**77773163**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces naeslundii		100.0	0.38	IND 1			
77773502	**Low discrimination**						**77773502**
					GRA		
Actinomyces israelii		87.79	0.41	IND 1	99		
Mitsuokella multacida		8.28	0.11	XYL 92	1		
Bifidobacterium adolescentis		3.93	0.12		99		
77775002	**Good identification**						**77775002**
					SPO		
Actinomyces israelii		95.84	0.37	IND 1	1		
Clostridium butyricum		4.16	-.07		99		
77775102	**Low discrimination**						**77775102**
					SPO		
Actinomyces israelii		76.98	0.48	IND 1	1		
Clostridium butyricum		20.07	0.17		99		
Bifidobacterium adolescentis		1.70	0.13		1		
77775140	**Low discrimination**						**77775140**
					GRA	SPO	
Actinomyces israelii		55.86	0.38	IND 1	99	1	
Mitsuokella multacida		31.58	0.21		1	1	
Clostridium butyricum		12.56	0.06		99	99	
77775142	**Acceptable identification**						**77775142**
					GRA	SPO	
Actinomyces israelii		90.89	0.41	IND 1	99	1	
Clostridium butyricum		5.30	-.02		99	99	
Mitsuokella multacida		3.81	0.04		1	1	
77775322	**Low discrimination**						**77775322**
					CAT		
Bacteroides ovatus		89.26	0.40	NIT 1	68		
Bacteroides thetaiotaomicron		5.19	0.19	ARA 97	89		
Bacteroides uniformis		3.72	0.14		5		
77775332	**Acceptable identification**						**77775332**
Bacteroides ovatus		94.48	0.37	NIT 1			
Bacteroides thetaiotaomicron		3.21	0.12	ARA 97			
Parabacteroides distasonis		1.57	0.23				
77775402	**Low discrimination**						**77775402**
					GRA	SPO	
Actinomyces israelii		64.20	0.36	IND 1	99	1	
Clostridium butyricum		30.68	0.09		99	99	
Mitsuokella multacida		5.11	0.04		1	1	
77775500	**Low discrimination**						**77775500**
					GRA	SPO	
Mitsuokella multacida		68.81	0.55	IND 1	1	1	
Clostridium butyricum		28.60	0.40		99	99	
Actinomyces israelii		2.59	0.43		99	1	
77775501	**Good identification**						**77775501**
					GRA	SPO	
Mitsuokella multacida		96.36	0.39	IND 1	1	1	
Clostridium butyricum		3.64	0.06	URE 10	99	99	
77775510	**Acceptable identification**						**77775510**
					GRA	SPO	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Mitsuokella multacida		92.38	0.47	IND 1	1	1	
Clostridium butyricum		6.06	0.18		99	99	
Actinomyces israelii		1.56	0.29		99	1	
77775520	**Low discrimination**						**77775520**
					GRA	SPO	
Mitsuokella multacida		84.10	0.45	IND 1	1	1	
Clostridium butyricum		12.16	0.22		99	99	
Bacteroides ovatus		2.01	0.10		1	1	
77775522	**Low discrimination**						**77775522**
					CAT		
Bacteroides ovatus		64.98	0.44	NIT 1	68		
Bacteroides thetaiotaomicron		16.55	0.34	RHA 95	89		
Bacteroides uniformis		13.23	0.29		5		
77775530	**Acceptable identification**						**77775530**
					GRA	SPO	
Mitsuokella multacida		93.40	0.37	IND 1	1	1	
Bacteroides ovatus		4.47	0.07		1	1	
Clostridium butyricum		2.13	0.00		99	99	
77775532	**Low discrimination**						**77775532**
					CAT		
Bacteroides ovatus		83.74	0.41	NIT 1	68		
Bacteroides thetaiotaomicron		12.44	0.27	RHA 95	89		
Mitsuokella multacida		1.96	0.20		1		
77775540	**Good identification**						**77775540**
					GRA	SPO	
Mitsuokella multacida		95.17	0.55	IND 1	1	1	
Clostridium butyricum		3.44	0.22		99	99	
Actinomyces israelii		1.39	0.36		99	1	
77775541	**Good identification**						**77775541**
Mitsuokella multacida		100.0	0.39	IND 1			
77775542	**Low discrimination**						**77775542**
					GRA	SPO	
Mitsuokella multacida		74.90	0.39	IND 1	1	1	
Actinomyces israelii		14.78	0.39	BGL 10	99	1	
Clostridium butyricum		9.48	0.15		99	99	
77775550	**Good identification**						**77775550**
					GRA	SPO	
Mitsuokella multacida		98.45	0.47	IND 1	1	1	
Actinomyces israelii		0.65	0.22		99	1	
Clostridium butyricum		0.56	0.00		99	99	
77775560	**Good identification**						**77775560**
					GRA	SPO	
Mitsuokella multacida		98.40	0.45	IND 1	1	1	
Clostridium butyricum		1.24	0.04		99	99	
Actinomyces israelii		0.37	0.16		99	1	
77775570	**Good identification**						**77775570**
Mitsuokella multacida		100.0	0.37	IND 1			
77775700	**Very good identification**						**77775700**
					GRA	SPO	
Mitsuokella multacida		99.10	0.60	IND 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.69	0.14		99	99	
Actinomyces israelii		0.15	0.24		99	1	
77775701	**Good identification**						**77775701**
Mitsuokella multacida		100.0	0.43	IND 1			
77775702	**Low discrimination**						**77775702**
					CAT		
Mitsuokella multacida		57.73	0.43	IND 1	1		
Bacteroides ovatus		31.80	0.32	BGL 10	68		
Bacteroides thetaiotaomicron		5.66	0.19		89		
77775710	**Good identification**						**77775710**
					GRA	SPO	
Mitsuokella multacida		99.77	0.51	IND 1	1	1	
Bacteroides ovatus		0.12	-.06		1	1	
Clostridium butyricum		0.11	-.08		99	99	
77775711	**Good identification**						**77775711**
Mitsuokella multacida		100.0	0.35	IND 1			
77775720	**Low discrimination**						**77775720**
					CAT		
Mitsuokella multacida		77.25	0.49	IND 1	1		
Bacteroides ovatus		18.91	0.32		68		
Bacteroides thetaiotaomicron		3.37	0.19		89		
77775722	**Low discrimination**						**77775722**
					CAT		
Bacteroides ovatus		83.44	0.66	NIT 1	68		
Bacteroides thetaiotaomicron		14.86	0.53		89		
Bacteroides uniformis		1.24	0.32		5		
77775730	**Low discrimination**						**77775730**
					CAT		
Mitsuokella multacida		64.91	0.41	IND 1	1		
Bacteroides ovatus		31.78	0.29		68		
Bacteroides thetaiotaomicron		3.30	0.12		89		
77775732	**Good identification**						**77775732**
Bacteroides ovatus		90.05	0.63	NIT 1			
Bacteroides thetaiotaomicron		9.36	0.46				
Clostridium clostridioforme		0.34	0.32				
77775740	**Very good identification**						**77775740**
					GRA	SPO	
Mitsuokella multacida		99.88	0.60	IND 1	1	1	
Clostridium butyricum		0.06	-.04		99	99	
Actinomyces israelii		0.06	0.17		99	1	
77775741	**Good identification**						**77775741**
Mitsuokella multacida		100.0	0.43	IND 1			
77775742	**Good identification**						**77775742**
					GRA		
Mitsuokella multacida		97.00	0.43	IND 1	1		
Bacteroides ovatus		2.23	0.08	BGL 10	1		
Actinomyces israelii		0.78	0.20		99		
77775750	**Good identification**						**77775750**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COC	GRA	
Mitsuokella multacida		99.81	0.51	IND 1	1	1	
Blautia producta		0.19	-.03		99	99	
77775751	**Good identification**						**77775751**
Mitsuokella multacida		100.0	0.35	IND 1			
77775752	**Good identification**						**77775752**
Mitsuokella multacida		95.61	0.35	IND 1			
Bacteroides ovatus		4.39	0.05	BGL 10			
77775760	**Good identification**						**77775760**
Mitsuokella multacida		98.99	0.49	IND 1			
Bacteroides ovatus		1.01	0.08				
77775762	**Low discrimination**						**77775762**
					CAT		
Bacteroides ovatus		86.45	0.42	NIT 1	68		
Mitsuokella multacida		9.51	0.33	SOR 4	1		
Bacteroides thetaiotaomicron		3.73	0.19		89		
77775770	**Good identification**						**77775770**
Mitsuokella multacida		98.00	0.41	IND 1			
Bacteroides ovatus		2.00	0.05				
77775772	**Acceptable identification**						**77775772**
					CAT		
Bacteroides ovatus		92.58	0.39	NIT 1	68		
Mitsuokella multacida		5.09	0.25	SOR 4	1		
Bacteroides thetaiotaomicron		2.33	0.12		89		
77777000	**Good identification**						**77777000**
Actinomyces israelii		100.0	0.47	IND 1			
77777002	**Good identification**						**77777002**
Actinomyces israelii		100.0	0.50	IND 1			
77777012	**Good identification**						**77777012**
Actinomyces israelii		100.0	0.36	IND 1			
77777040	**Good identification**						**77777040**
Actinomyces israelii		100.0	0.40	IND 1			
77777042	**Good identification**						**77777042**
Actinomyces israelii		100.0	0.43	IND 1			
77777100	**Good identification**						**77777100**
					GRA	SPO	
Actinomyces israelii		90.07	0.58	IND 1	99	1	
Mitsuokella multacida		8.16	0.27		1	1	
Clostridium butyricum		1.78	0.07		99	99	
77777102	**Very good identification**						**77777102**
					GRA	SPO	
Actinomyces israelii		98.52	0.61	IND 1	99	1	
Mitsuokella multacida		0.66	0.11		1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Clostridium butyricum		0.50	0.00		99	99	
77777110	**Low discrimination**						**77777110**
					GRA		
Actinomyces israelii		83.20	0.43	IND 1	99		
Mitsuokella multacida		16.80	0.19	MLZ 13	1		
77777112	**Acceptable identification**						**77777112**
Actinomyces israelii		94.21	0.46	IND 1			
Bifidobacterium adolescentis		2.46	0.13	MLZ 13			
Bifidobacterium dentium		1.93	-.07				
77777120	**Low discrimination**						**77777120**
					GRA		
Actinomyces israelii		73.82	0.37	IND 1	99		
Mitsuokella multacida		26.18	0.17	NAG 6	1		
77777122	**Low discrimination**						**77777122**
					GRA		
Actinomyces israelii		80.32	0.40	IND 1	99		
Bacteroides ovatus		19.68	0.10	NAG 6	1		
77777140	**Low discrimination**						**77777140**
					GRA		
Actinomyces israelii		81.11	0.51	IND 1	99		
Mitsuokella multacida		18.89	0.27		1		
77777142	**Good identification**						**77777142**
					GRA		
Actinomyces israelii		98.30	0.54	IND 1	99		
Mitsuokella multacida		1.70	0.11		1		
77777150	**Low discrimination**						**77777150**
					COC	GRA	
Actinomyces israelii		62.61	0.36	IND 1	1	99	
Mitsuokella multacida		32.52	0.19	MLZ 13	1	1	
Blautia producta		4.88	-.03		99	99	
77777152	**Good identification**						**77777152**
					GRA		
Actinomyces israelii		96.29	0.39	IND 1	99		
Mitsuokella multacida		3.71	0.03	MLZ 13	1		
77777302	**Low discrimination**						**77777302**
					GRA		
Actinomyces israelii		82.96	0.41	IND 1	99		
Mitsuokella multacida		13.73	0.15	RHA 7	1		
Bacteroides ovatus		3.31	-.02		1		
77777400	**Low discrimination**						**77777400**
					GRA	SPO	
Actinomyces israelii		67.62	0.46	IND 1	99	1	
Mitsuokella multacida		29.93	0.27		1	1	
Clostridium butyricum		2.44	0.00		99	99	
77777402	**Good identification**						**77777402**
					GRA	SPO	
Actinomyces israelii		95.95	0.49	IND 1	99	1	
Mitsuokella multacida		3.15	0.11		1	1	
Clostridium butyricum		0.90	-.07		99	99	
77777442	**Acceptable identification**						**77777442**
					GRA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Actinomyces israelii		92.22	0.42	IND 1	99		
Mitsuokella multacida		7.78	0.11		1		
77777500	**Good identification**						**77777500**
					GRA	SPO	
Mitsuokella multacida		90.00	0.61	IND 1	1	1	
Actinomyces israelii		8.22	0.56		99	1	
Clostridium butyricum		1.78	0.24		99	99	
77777501	**Good identification**						**77777501**
					GRA		
Mitsuokella multacida		99.18	0.45	IND 1	1		
Actinomyces israelii		0.82	0.22	URE 10	99		
77777502	**Low discrimination**						**77777502**
					GRA	SPO	
Actinomyces israelii		52.53	0.59	IND 1	99	1	
Mitsuokella multacida		42.63	0.45		1	1	
Clostridium butyricum		2.95	0.17		99	99	
77777510	**Acceptable identification**						**77777510**
					GRA		
Mitsuokella multacida		94.29	0.53	IND 1	1		
Actinomyces israelii		3.86	0.42		99		
Bifidobacterium dentium		1.42	0.10		99		
77777511	**Good identification**						**77777511**
Mitsuokella multacida		100.0	0.37	IND 1			
77777520	**Good identification**						**77777520**
					GRA	SPO	
Mitsuokella multacida		96.75	0.51	IND 1	1	1	
Actinomyces israelii		2.26	0.36		99	1	
Clostridium butyricum		0.67	0.06		99	99	
77777522	**Low discrimination**						**77777522**
					GRA		
Bacteroides ovatus		68.30	0.36	NIT 1	1		
Mitsuokella multacida		22.42	0.35	RHA 95	1		
Actinomyces israelii		7.06	0.39		99		
77777530	**Low discrimination**						**77777530**
					GRA		
Mitsuokella multacida		88.94	0.43	IND 1	1		
Bifidobacterium dentium		9.52	0.15		99		
Actinomyces israelii		0.93	0.22		99		
77777540	**Good identification**						**77777540**
					COC	GRA	
Mitsuokella multacida		94.89	0.61	IND 1	1	1	
Actinomyces israelii		3.37	0.49		1	99	
Blautia producta		1.58	0.24		99	99	
77777541	**Good identification**						**77777541**
					GRA		
Mitsuokella multacida		99.68	0.45	IND 1	1		
Actinomyces israelii		0.32	0.15	URE 10	99		
77777542	**Low discrimination**						**77777542**
					COC	GRA	
Mitsuokella multacida		66.41	0.45	IND 1	1	1	
Actinomyces israelii		31.82	0.52	BGL 10	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Blautia producta		1.11	0.07		99	99	
77777550	**Low discrimination**						**77777550**
					COC	GRA	
Mitsuokella multacida		85.77	0.53	IND 1	1	1	
Blautia producta		12.87	0.32		99	99	
Actinomyces israelii		1.37	0.35		1	99	
77777551	**Good identification**						**77777551**
					COC	GRA	
Mitsuokella multacida		98.66	0.37	IND 1	1	1	
Blautia producta		1.35	-.03	URE 10	99	99	
77777552	**Low discrimination**						**77777552**
					COC	GRA	
Mitsuokella multacida		71.78	0.37	IND 1	1	1	
Actinomyces israelii		15.42	0.38	BGL 10	1	99	
Blautia producta		10.77	0.15		99	99	
77777560	**Good identification**						**77777560**
					COC	GRA	
Mitsuokella multacida		98.38	0.51	IND 1	1	1	
Actinomyces israelii		0.89	0.29		1	99	
Blautia producta		0.73	0.07		99	99	
77777570	**Acceptable identification**						**77777570**
					COC	GRA	
Mitsuokella multacida		93.39	0.43	IND 1	1	1	
Blautia producta		6.23	0.15		99	99	
Actinomyces israelii		0.38	0.15		1	99	
77777700	**Very good identification**						**77777700**
					GRA	SPO	
Mitsuokella multacida		99.60	0.66	IND 1	1	1	
Actinomyces israelii		0.37	0.37		99	1	
Clostridium butyricum		0.03	-.02		99	99	
77777701	**Good identification**						**77777701**
Mitsuokella multacida		100.0	0.50	IND 1			
77777702	**Low discrimination**						**77777702**
					GRA		
Mitsuokella multacida		88.60	0.50	IND 1	1		
Bacteroides ovatus		6.97	0.24	BGL 10	1		
Actinomyces israelii		4.43	0.40		99		
77777710	**Very good identification**						**77777710**
					COC	GRA	
Mitsuokella multacida		99.75	0.58	IND 1	1	1	
Actinomyces israelii		0.17	0.23		1	99	
Blautia producta		0.08	-.03		99	99	
77777711	**Good identification**						**77777711**
Mitsuokella multacida		100.0	0.41	IND 1			
77777712	**Low discrimination**						**77777712**
					GRA		
Mitsuokella multacida		84.23	0.41	IND 1	1		
Bacteroides ovatus		13.26	0.21	BGL 10	1		
Actinomyces israelii		1.89	0.26		99		
77777720	**Good identification**						**77777720**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					GRA		
Mitsuokella multacida		96.53	0.56	IND 1	1		
Bacteroides ovatus		3.38	0.24		1		
Actinomyces israelii		0.09	0.16		99		
77777721	**Good identification**						**77777721**
Mitsuokella multacida		100.0	0.39	IND 1			
77777722	**Very good identification**						**77777722**
					CAT		
Bacteroides ovatus		96.31	0.58	NIT 1	68		
Mitsuokella multacida		3.09	0.39		1		
Bacteroides thetaiotaomicron		0.52	0.19		89		
77777730	**Acceptable identification**						**77777730**
Mitsuokella multacida		93.46	0.47	IND 1			
Bacteroides ovatus		6.54	0.21				
77777732	**Good identification**						**77777732**
					CAT		
Bacteroides ovatus		98.04	0.55	NIT 1	68		
Mitsuokella multacida		1.57	0.31		1		
Bacteroides thetaiotaomicron		0.31	0.12		89		
77777740	**Very good identification**						**77777740**
					COC	GRA	
Mitsuokella multacida		98.97	0.66	IND 1	1	1	
Blautia producta		0.89	0.24		99	99	
Actinomyces israelii		0.14	0.30		1	99	
77777741	**Good identification**						**77777741**
Mitsuokella multacida		100.0	0.50	IND 1			
77777742	**Good identification**						**77777742**
					COC	GRA	
Mitsuokella multacida		96.93	0.50	IND 1	1	1	
Actinomyces israelii		1.88	0.33	BGL 10	1	99	
Blautia producta		0.87	0.07		99	99	
77777750	**Good identification**						**77777750**
					COC	GRA	
Mitsuokella multacida		92.47	0.58	IND 1	1	1	
Blautia producta		7.47	0.32		99	99	
Actinomyces israelii		0.06	0.16		1	99	
77777751	**Good identification**						**77777751**
					COC	GRA	
Mitsuokella multacida		99.27	0.41	IND 1	1	1	
Blautia producta		0.73	-.03	URE 10	99	99	
77777752	**Acceptable identification**						**77777752**
					COC	GRA	
Mitsuokella multacida		91.24	0.41	IND 1	1	1	
Blautia producta		7.37	0.15	BGL 10	99	99	
Actinomyces israelii		0.79	0.19		1	99	
77777760	**Good identification**						**77777760**
					COC	GRA	
Mitsuokella multacida		99.46	0.56	IND 1	1	1	
Blautia producta		0.40	0.07		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Bacteroides ovatus		0.15	0.00		1	1	
77777761	**Good identification**						**77777761**
Mitsuokella multacida		100.0	0.39	IND 1			
77777770	**Good identification**						**77777770**
					COC	GRA	
Mitsuokella multacida		96.26	0.47	IND 1	1	1	
Blautia producta		3.46	0.15		99	99	
Bacteroides ovatus		0.28	-.03		1	1	

Erba Lachema s.r.o.
Karásek 2219/1d, 621 00 Brno, Czech Republic
Tel.: +420 517 077 111
E-mail: diagnostics@erbamannheim.com
www.erbalachema.com

Contact info / map